Full wwPDB EM Validation Report ⓘ

Feb 28, 2024 – 05:21 am GMT

PDB ID	:	8CMO
EMDB ID	:	EMD-16732
Title	:	Cryo-EM structure of the Photosystem I - LHCI supercomplex from Coelastrella sp.
Authors	:	Fadeeva, M.; Klaiman, D.; Nelson, N.
Deposited on	:	2023-02-20
Resolution	:	2.81 Å(reported)

This is a Full wwPDB EM Validation Report for a publicly released PDB entry.

We welcome your comments at *validation@mail.wwpdb.org*
A user guide is available at
https://www.wwpdb.org/validation/2017/EMValidationReportHelp
with specific help available everywhere you see the ⓘ symbol.

The types of validation reports are described at
http://www.wwpdb.org/validation/2017/FAQs#types.

The following versions of software and data (see references ⓘ) were used in the production of this report:

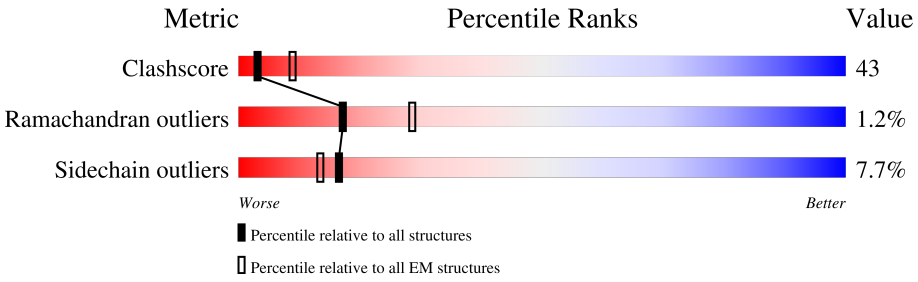
EMDB validation analysis	:	0.0.1.dev70
Mogul	:	1.8.4, CSD as541be (2020)
MolProbity	:	4.02b-467
buster-report	:	1.1.7 (2018)
Percentile statistics	:	20191225.v01 (using entries in the PDB archive December 25th 2019)
MapQ	:	1.9.13
Ideal geometry (proteins)	:	Engh & Huber (2001)
Ideal geometry (DNA, RNA)	:	Parkinson et al. (1996)
Validation Pipeline (wwPDB-VP)	:	2.36

1 Overall quality at a glance (i)

The following experimental techniques were used to determine the structure:
ELECTRON MICROSCOPY

The reported resolution of this entry is 2.81 Å.

Percentile scores (ranging between 0-100) for global validation metrics of the entry are shown in the following graphic. The table shows the number of entries on which the scores are based.



Metric	Whole archive (#Entries)	EM structures (#Entries)
Clashscore	158937	4297
Ramachandran outliers	154571	4023
Sidechain outliers	154315	3826

The table below summarises the geometric issues observed across the polymeric chains and their fit to the map. The red, orange, yellow and green segments of the bar indicate the fraction of residues that contain outliers for >=3, 2, 1 and 0 types of geometric quality criteria respectively. A grey segment represents the fraction of residues that are not modelled. The numeric value for each fraction is indicated below the corresponding segment, with a dot representing fractions <=5% The upper red bar (where present) indicates the fraction of residues that have poor fit to the EM map (all-atom inclusion < 40%). The numeric value is given above the bar.

Mol	Chain	Length	Quality of chain
1	A	740	61% 38%
2	B	732	55% 42%
3	C	80	49% 49%
4	D	142	8% 56% 41%
5	E	61	10% 59% 38%
6	F	165	52% 44%
7	G	93	35% 41% 55%
8	I	41	7% 41% 59%

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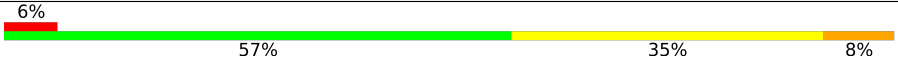
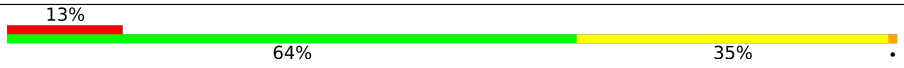
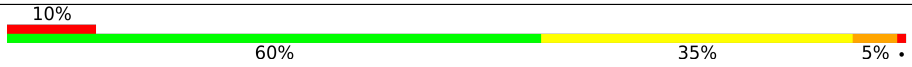
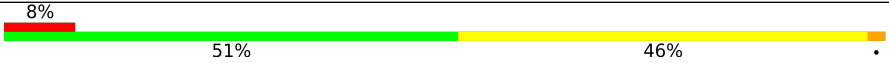
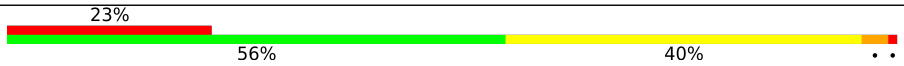
Mol	Chain	Length	Quality of chain
9	J	40	
10	K	89	
11	L	123	
12	1	194	
12	Z	194	
13	3	218	
14	7	220	
15	8	214	
16	4	207	
17	5	226	
18	6	226	
19	9	184	
20	2	70	

The following table lists non-polymeric compounds, carbohydrate monomers and non-standard residues in protein, DNA, RNA chains that are outliers for geometric or electron-density-fit criteria:

Mol	Type	Chain	Res	Chirality	Geometry	Clashes	Electron density
21	CL0	A	801	X	-	-	-
22	CLA	1	304	X	-	-	-
22	CLA	1	305	X	-	-	-
22	CLA	1	306	X	-	-	-
22	CLA	1	307	X	-	X	-
22	CLA	1	308	X	-	X	-
22	CLA	1	309	X	-	-	-
22	CLA	1	310	X	-	-	-
22	CLA	1	311	X	-	-	-
22	CLA	1	313	X	-	-	-
22	CLA	1	314	X	-	-	-
22	CLA	1	315	X	-	-	-
22	CLA	1	316	X	-	-	-
22	CLA	2	301	X	-	-	-
22	CLA	2	302	X	-	-	-

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Mol	Type	Chain	Res	Chirality	Geometry	Clashes	Electron density
22	CLA	2	304	X	-	-	-
22	CLA	3	308	X	-	-	-
22	CLA	3	309	X	-	-	-
22	CLA	3	310	X	-	-	-
22	CLA	3	311	X	-	-	-
22	CLA	3	312	X	-	X	-
22	CLA	3	313	X	-	X	-
22	CLA	3	314	X	-	-	-
22	CLA	3	315	X	-	-	-
22	CLA	3	316	X	-	-	-
22	CLA	3	318	X	-	-	-
22	CLA	3	319	X	-	-	-
22	CLA	3	320	X	-	-	-
22	CLA	3	322	X	-	-	-
22	CLA	4	805	X	-	-	-
22	CLA	4	806	X	-	-	-
22	CLA	4	807	X	-	-	-
22	CLA	4	808	X	-	-	-
22	CLA	4	809	X	-	-	-
22	CLA	4	810	X	-	-	-
22	CLA	4	811	X	-	-	-
22	CLA	4	812	X	-	-	-
22	CLA	4	815	X	-	-	-
22	CLA	4	817	X	-	-	-
22	CLA	4	818	X	-	-	-
22	CLA	5	301	X	-	-	-
22	CLA	5	307	X	-	X	-
22	CLA	5	308	X	-	X	-
22	CLA	5	309	X	-	-	-
22	CLA	5	310	X	-	-	-
22	CLA	5	311	X	-	-	-
22	CLA	5	312	X	-	-	-
22	CLA	5	313	X	-	-	-
22	CLA	5	314	X	-	-	-
22	CLA	5	315	X	-	-	-
22	CLA	5	318	X	-	X	-
22	CLA	5	319	X	-	-	-
22	CLA	5	320	X	-	-	-
22	CLA	5	322	X	-	-	-
22	CLA	5	323	X	-	-	-
22	CLA	5	326	X	-	-	-
22	CLA	6	301	X	-	-	-

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Mol	Type	Chain	Res	Chirality	Geometry	Clashes	Electron density
22	CLA	6	302	X	-	-	-
22	CLA	6	307	X	-	X	-
22	CLA	6	308	X	-	-	-
22	CLA	6	309	X	-	X	-
22	CLA	6	310	X	-	-	-
22	CLA	6	311	X	-	-	-
22	CLA	6	312	X	-	-	-
22	CLA	6	313	X	-	-	-
22	CLA	6	314	X	-	-	-
22	CLA	6	317	X	-	-	-
22	CLA	6	319	X	-	-	-
22	CLA	6	321	X	-	-	-
22	CLA	6	322	X	-	X	-
22	CLA	7	304	X	-	-	-
22	CLA	7	305	X	-	-	-
22	CLA	7	306	X	-	-	-
22	CLA	7	307	X	-	X	-
22	CLA	7	308	X	-	-	-
22	CLA	7	309	X	-	-	-
22	CLA	7	310	X	-	-	-
22	CLA	7	311	X	-	-	-
22	CLA	7	312	X	-	-	-
22	CLA	7	314	X	-	-	-
22	CLA	7	315	X	-	-	-
22	CLA	7	316	X	-	-	-
22	CLA	7	317	X	-	-	-
22	CLA	7	322	X	-	-	-
22	CLA	7	323	X	-	-	-
22	CLA	7	324	X	-	-	-
22	CLA	8	305	X	-	X	-
22	CLA	8	306	X	-	-	-
22	CLA	8	307	X	-	-	-
22	CLA	8	308	X	-	-	-
22	CLA	8	309	X	-	-	-
22	CLA	8	310	X	-	-	-
22	CLA	8	311	X	-	-	-
22	CLA	8	313	X	-	-	-
22	CLA	8	314	X	-	-	-
22	CLA	8	316	X	-	-	-
22	CLA	9	303	X	-	-	-
22	CLA	9	304	X	-	-	-
22	CLA	9	305	X	-	-	-

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Mol	Type	Chain	Res	Chirality	Geometry	Clashes	Electron density
22	CLA	9	306	X	-	-	-
22	CLA	9	307	X	-	-	-
22	CLA	9	308	X	-	-	-
22	CLA	9	309	X	-	-	-
22	CLA	9	310	X	-	-	-
22	CLA	9	311	X	-	-	-
22	CLA	9	313	X	-	-	-
22	CLA	A	802	X	-	-	-
22	CLA	A	803	X	-	-	-
22	CLA	A	804	X	-	-	-
22	CLA	A	805	X	-	X	-
22	CLA	A	806	X	-	-	-
22	CLA	A	807	X	-	-	-
22	CLA	A	808	X	-	-	-
22	CLA	A	809	X	-	-	-
22	CLA	A	810	X	-	X	-
22	CLA	A	811	X	-	-	-
22	CLA	A	812	X	-	X	-
22	CLA	A	813	X	-	-	-
22	CLA	A	814	X	-	X	-
22	CLA	A	815	X	-	-	-
22	CLA	A	816	X	-	X	-
22	CLA	A	817	X	-	-	-
22	CLA	A	818	X	-	-	-
22	CLA	A	819	X	-	-	-
22	CLA	A	820	X	-	-	-
22	CLA	A	821	X	-	X	-
22	CLA	A	822	X	-	X	-
22	CLA	A	823	X	-	X	-
22	CLA	A	824	X	-	-	-
22	CLA	A	825	X	-	-	-
22	CLA	A	826	X	-	-	-
22	CLA	A	827	X	-	-	-
22	CLA	A	828	X	-	-	-
22	CLA	A	829	X	-	-	-
22	CLA	A	830	X	-	-	-
22	CLA	A	831	X	-	-	-
22	CLA	A	832	X	-	-	-
22	CLA	A	833	X	-	-	-
22	CLA	A	834	X	-	-	-
22	CLA	A	835	X	-	-	-
22	CLA	A	836	X	-	-	-

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Mol	Type	Chain	Res	Chirality	Geometry	Clashes	Electron density
22	CLA	A	837	X	-	-	-
22	CLA	A	838	X	-	-	-
22	CLA	A	839	X	-	-	-
22	CLA	A	840	X	-	-	-
22	CLA	A	841	X	-	-	-
22	CLA	A	842	X	-	-	-
22	CLA	A	854	X	-	-	-
22	CLA	A	855	X	-	-	-
22	CLA	B	801	X	-	-	-
22	CLA	B	805	X	-	-	-
22	CLA	B	806	X	-	X	-
22	CLA	B	807	X	-	-	-
22	CLA	B	808	X	-	-	-
22	CLA	B	809	X	-	-	-
22	CLA	B	810	X	-	-	-
22	CLA	B	811	X	-	X	-
22	CLA	B	812	X	-	-	-
22	CLA	B	813	X	-	-	-
22	CLA	B	814	X	-	-	-
22	CLA	B	815	X	-	X	-
22	CLA	B	816	X	-	-	-
22	CLA	B	817	X	-	-	-
22	CLA	B	818	X	-	-	-
22	CLA	B	819	X	-	-	-
22	CLA	B	820	X	-	-	-
22	CLA	B	821	X	-	-	-
22	CLA	B	822	X	-	-	-
22	CLA	B	823	X	-	-	-
22	CLA	B	824	X	-	X	-
22	CLA	B	825	X	-	X	-
22	CLA	B	826	X	-	-	-
22	CLA	B	827	X	-	X	-
22	CLA	B	828	X	-	X	-
22	CLA	B	829	X	-	-	-
22	CLA	B	830	X	-	-	-
22	CLA	B	831	X	-	-	-
22	CLA	B	832	X	-	-	-
22	CLA	B	833	X	-	-	-
22	CLA	B	834	X	-	-	-
22	CLA	B	835	X	-	-	-
22	CLA	B	836	X	-	-	-
22	CLA	B	837	X	-	-	-

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Mol	Type	Chain	Res	Chirality	Geometry	Clashes	Electron density
22	CLA	B	838	X	-	-	-
22	CLA	B	839	X	-	-	-
22	CLA	B	840	X	-	-	-
22	CLA	B	841	X	-	-	-
22	CLA	B	842	X	-	X	-
22	CLA	F	301	X	-	-	-
22	CLA	F	302	X	-	-	-
22	CLA	F	304	X	-	X	-
22	CLA	F	305	X	-	-	-
22	CLA	G	201	X	-	-	-
22	CLA	G	202	X	-	-	-
22	CLA	J	103	X	-	-	-
22	CLA	K	202	X	-	-	-
22	CLA	K	203	X	-	-	-
22	CLA	K	204	X	-	-	-
22	CLA	K	205	X	-	-	-
22	CLA	L	201	X	-	-	-
22	CLA	L	202	X	-	-	-
22	CLA	L	203	X	-	-	-
22	CLA	Z	303	X	-	X	-
22	CLA	Z	304	X	-	-	-
22	CLA	Z	305	X	-	-	-
22	CLA	Z	306	X	-	-	-
22	CLA	Z	307	X	-	-	-
22	CLA	Z	308	X	-	-	-
22	CLA	Z	309	X	-	X	-
22	CLA	Z	310	X	-	-	-
22	CLA	Z	313	X	-	X	-
22	CLA	Z	314	X	-	-	-
22	CLA	Z	315	X	-	-	-
22	CLA	Z	316	X	-	-	-
24	BCR	3	304	-	X	-	-
24	BCR	3	307	-	X	-	-
24	BCR	4	804	-	-	X	-
24	BCR	5	304	-	-	X	-
24	BCR	6	306	-	X	-	-
24	BCR	A	844	-	-	X	-
24	BCR	A	845	-	X	-	-
24	BCR	A	856	-	X	X	-
24	BCR	B	846	-	X	-	-
24	BCR	B	847	-	-	X	-
24	BCR	B	848	-	X	-	-

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Mol	Type	Chain	Res	Chirality	Geometry	Clashes	Electron density
24	BCR	B	853	-	X	X	-
29	SF4	B	802	-	-	X	-
29	SF4	C	101	-	-	X	-
29	SF4	C	102	-	-	X	-
31	DGD	3	301	-	-	X	-
37	DAO	K	201	-	-	X	-
38	LUT	Z	301	-	-	X	-
39	CHL	1	312	X	-	-	-
39	CHL	2	303	X	-	-	-
39	CHL	3	317	X	-	-	-
39	CHL	4	813	X	-	-	-
39	CHL	4	814	X	-	-	-
39	CHL	4	816	X	-	-	-
39	CHL	4	819	X	-	-	-
39	CHL	5	316	X	-	-	-
39	CHL	5	317	X	-	-	-
39	CHL	5	321	X	-	-	-
39	CHL	6	315	X	-	X	-
39	CHL	6	316	X	-	-	-
39	CHL	6	318	X	-	-	-
39	CHL	6	320	X	-	-	-
39	CHL	7	313	X	-	-	-
39	CHL	8	301	X	-	-	-
39	CHL	8	312	X	-	-	-
39	CHL	8	315	X	-	X	-
39	CHL	9	312	X	-	X	-
39	CHL	9	314	X	-	-	-
39	CHL	Z	311	X	-	-	-
39	CHL	Z	312	X	-	-	-

2 Entry composition

There are 44 unique types of molecules in this entry. The entry contains 50057 atoms, of which 0 are hydrogens and 0 are deuteriums.

In the tables below, the AltConf column contains the number of residues with at least one atom in alternate conformation and the Trace column contains the number of residues modelled with at most 2 atoms.

- Molecule 1 is a protein called Photosystem I P700 chlorophyll a apoprotein A1 PsaA.

Mol	Chain	Residues	Atoms					AltConf	Trace
			Total	C	N	O	S		
1	A	740	5819	3806	991	1000	22	0	0

- Molecule 2 is a protein called Photosystem I P700 chlorophyll a apoprotein A2 PsaB.

Mol	Chain	Residues	Atoms					AltConf	Trace
			Total	C	N	O	S		
2	B	732	5800	3804	973	1005	18	0	0

- Molecule 3 is a protein called Photosystem I subunit VII PsaC.

Mol	Chain	Residues	Atoms					AltConf	Trace
			Total	C	N	O	S		
3	C	80	603	370	105	116	12	0	0

- Molecule 4 is a protein called Photosystem I reaction center subunit II PsaD.

Mol	Chain	Residues	Atoms					AltConf	Trace
			Total	C	N	O	S		
4	D	142	1129	724	196	204	5	0	0

- Molecule 5 is a protein called Photosystem I reaction center subunit IV PsaE.

Mol	Chain	Residues	Atoms				AltConf	Trace
			Total	C	N	O		
5	E	61	484	309	83	92	0	0

- Molecule 6 is a protein called Photosystem I reaction center subunit III PsaF.

Mol	Chain	Residues	Atoms					AltConf	Trace
			Total	C	N	O	S		
6	F	165	1265	813	219	230	3	0	0

- Molecule 7 is a protein called Photosystem I reaction center subunit V PsaG.

Mol	Chain	Residues	Atoms					AltConf	Trace
7	G	93	Total	C	N	O	S	0	0
			690	440	116	131	3		

- Molecule 8 is a protein called Photosystem I reaction center subunit VIII PsaI.

Mol	Chain	Residues	Atoms					AltConf	Trace
8	I	41	Total	C	N	O	S	0	0
			308	210	44	53	1		

- Molecule 9 is a protein called Photosystem I subunit IX PsaJ.

Mol	Chain	Residues	Atoms					AltConf	Trace
9	J	40	Total	C	N	O	S	0	0
			319	217	47	54	1		

- Molecule 10 is a protein called Photosystem I reaction center subunit X psaK.

Mol	Chain	Residues	Atoms					AltConf	Trace
10	K	89	Total	C	N	O	S	0	0
			617	390	110	115	2		

- Molecule 11 is a protein called Photosystem I reaction centre subunit XI PsaL.

Mol	Chain	Residues	Atoms					AltConf	Trace
11	L	123	Total	C	N	O	S	0	0
			897	580	148	165	4		

- Molecule 12 is a protein called Light-harvesting protein of photosystem I Lhca1.

Mol	Chain	Residues	Atoms					AltConf	Trace
12	1	194	Total	C	N	O	S	0	0
			1457	936	248	265	8		
12	Z	194	Total	C	N	O	S	0	0
			1457	936	248	265	8		

- Molecule 13 is a protein called Light-harvesting protein of photosystem I Lhca3.

Mol	Chain	Residues	Atoms					AltConf	Trace
13	3	218	Total	C	N	O	S	0	0
			1695	1107	272	309	7		

- Molecule 14 is a protein called Light-harvesting protein of photosystem I Lhca7.

Mol	Chain	Residues	Atoms					AltConf	Trace
14	7	220	Total	C	N	O	S	0	0
			1705	1102	283	313	7		

- Molecule 15 is a protein called Light-harvesting protein of photosystem I Lhca8.

Mol	Chain	Residues	Atoms					AltConf	Trace
15	8	214	Total	C	N	O	S	0	0
			1602	1032	269	291	10		

- Molecule 16 is a protein called Light-harvesting protein of photosystem I Lhca4.

Mol	Chain	Residues	Atoms					AltConf	Trace
16	4	207	Total	C	N	O	S	0	0
			1598	1036	272	284	6		

- Molecule 17 is a protein called Light-harvesting protein of photosystem I Lhca5.

Mol	Chain	Residues	Atoms					AltConf	Trace
17	5	226	Total	C	N	O	S	0	0
			1764	1136	302	318	8		

- Molecule 18 is a protein called Light-harvesting protein of photosystem I Lhca6.

Mol	Chain	Residues	Atoms					AltConf	Trace
18	6	226	Total	C	N	O	S	0	0
			1762	1162	289	303	8		

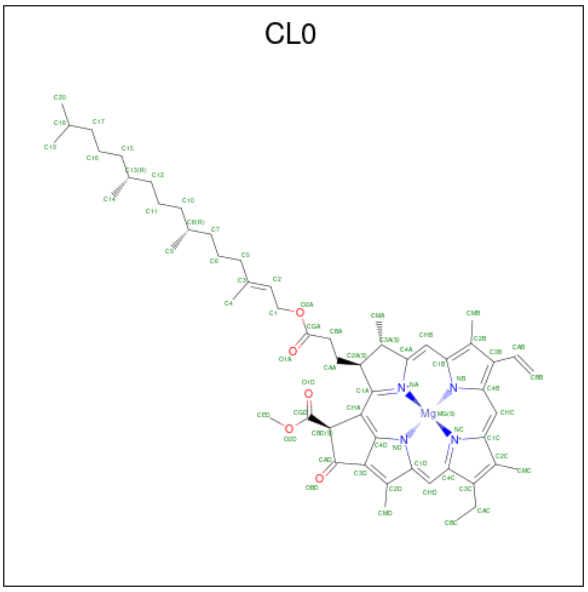
- Molecule 19 is a protein called Light-harvesting protein of photosystem I Lhca9.

Mol	Chain	Residues	Atoms					AltConf	Trace
19	9	184	Total	C	N	O	S	0	0
			1416	914	239	254	9		

- Molecule 20 is a protein called Light-harvesting protein of photosystem I Lhca2 partial.

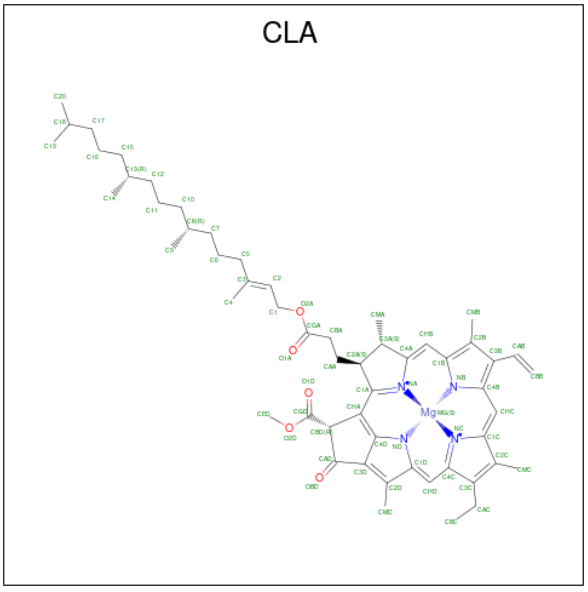
Mol	Chain	Residues	Atoms					AltConf	Trace
20	2	70	Total	C	N	O	S	0	0
			525	336	90	94	5		

- Molecule 21 is CHLOROPHYLL A ISOMER (three-letter code: CL0) (formula: $C_{55}H_{72}MgN_4O_5$).



Mol	Chain	Residues	Atoms					AltConf
21	A	1	Total	C	Mg	N	O	0
			65	55	1	4	5	

- Molecule 22 is CHLOROPHYLL A (three-letter code: CLA) (formula: $C_{55}H_{72}MgN_4O_5$).



Mol	Chain	Residues	Atoms					AltConf
22	A	1	Total	C	Mg	N	O	0
			65	55	1	4	5	
22	A	1	Total	C	Mg	N	O	0
			65	55	1	4	5	
22	A	1	Total	C	Mg	N	O	0
			45	35	1	4	5	

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Mol	Chain	Residues	Atoms					AltConf
22	A	1	Total	C	Mg	N	O	0
			65	55	1	4	5	
22	A	1	Total	C	Mg	N	O	0
			65	55	1	4	5	
22	A	1	Total	C	Mg	N	O	0
			60	50	1	4	5	
22	A	1	Total	C	Mg	N	O	0
			65	55	1	4	5	
22	A	1	Total	C	Mg	N	O	0
			55	45	1	4	5	
22	A	1	Total	C	Mg	N	O	0
			65	55	1	4	5	
22	A	1	Total	C	Mg	N	O	0
			65	55	1	4	5	
22	A	1	Total	C	Mg	N	O	0
			65	55	1	4	5	
22	A	1	Total	C	Mg	N	O	0
			65	55	1	4	5	
22	A	1	Total	C	Mg	N	O	0
			60	50	1	4	5	
22	A	1	Total	C	Mg	N	O	0
			65	55	1	4	5	
22	A	1	Total	C	Mg	N	O	0
			56	46	1	4	5	
22	A	1	Total	C	Mg	N	O	0
			60	50	1	4	5	
22	A	1	Total	C	Mg	N	O	0
			60	50	1	4	5	
22	A	1	Total	C	Mg	N	O	0
			65	55	1	4	5	
22	A	1	Total	C	Mg	N	O	0
			60	50	1	4	5	
22	A	1	Total	C	Mg	N	O	0
			65	55	1	4	5	
22	A	1	Total	C	Mg	N	O	0
			55	45	1	4	5	
22	A	1	Total	C	Mg	N	O	0
			65	55	1	4	5	
22	A	1	Total	C	Mg	N	O	0
			65	55	1	4	5	
22	A	1	Total	C	Mg	N	O	0
			65	55	1	4	5	

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Mol	Chain	Residues	Atoms					AltConf
22	A	1	Total	C	Mg	N	O	0
			50	40	1	4	5	
22	A	1	Total	C	Mg	N	O	0
			65	55	1	4	5	
22	A	1	Total	C	Mg	N	O	0
			65	55	1	4	5	
22	A	1	Total	C	Mg	N	O	0
			65	55	1	4	5	
22	A	1	Total	C	Mg	N	O	0
			65	55	1	4	5	
22	A	1	Total	C	Mg	N	O	0
			50	40	1	4	5	
22	A	1	Total	C	Mg	N	O	0
			65	55	1	4	5	
22	A	1	Total	C	Mg	N	O	0
			65	55	1	4	5	
22	A	1	Total	C	Mg	N	O	0
			65	55	1	4	5	
22	A	1	Total	C	Mg	N	O	0
			65	55	1	4	5	
22	A	1	Total	C	Mg	N	O	0
			55	45	1	4	5	
22	A	1	Total	C	Mg	N	O	0
			51	41	1	4	5	
22	A	1	Total	C	Mg	N	O	0
			65	55	1	4	5	
22	A	1	Total	C	Mg	N	O	0
			51	41	1	4	5	
22	A	1	Total	C	Mg	N	O	0
			65	55	1	4	5	
22	A	1	Total	C	Mg	N	O	0
			65	55	1	4	5	
22	A	1	Total	C	Mg	N	O	0
			52	42	1	4	5	
22	A	1	Total	C	Mg	N	O	0
			65	55	1	4	5	
22	A	1	Total	C	Mg	N	O	0
			65	55	1	4	5	
22	B	1	Total	C	Mg	N	O	0
			65	55	1	4	5	
22	B	1	Total	C	Mg	N	O	0
			65	55	1	4	5	

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Mol	Chain	Residues	Atoms					AltConf
22	B	1	Total	C	Mg	N	O	0
			65	55	1	4	5	
22	B	1	Total	C	Mg	N	O	0
			45	35	1	4	5	
22	B	1	Total	C	Mg	N	O	0
			65	55	1	4	5	
22	B	1	Total	C	Mg	N	O	0
			65	55	1	4	5	
22	B	1	Total	C	Mg	N	O	0
			56	46	1	4	5	
22	B	1	Total	C	Mg	N	O	0
			65	55	1	4	5	
22	B	1	Total	C	Mg	N	O	0
			60	50	1	4	5	
22	B	1	Total	C	Mg	N	O	0
			56	46	1	4	5	
22	B	1	Total	C	Mg	N	O	0
			60	50	1	4	5	
22	B	1	Total	C	Mg	N	O	0
			65	55	1	4	5	
22	B	1	Total	C	Mg	N	O	0
			60	50	1	4	5	
22	B	1	Total	C	Mg	N	O	0
			57	47	1	4	5	
22	B	1	Total	C	Mg	N	O	0
			65	55	1	4	5	
22	B	1	Total	C	Mg	N	O	0
			59	49	1	4	5	
22	B	1	Total	C	Mg	N	O	0
			60	50	1	4	5	
22	B	1	Total	C	Mg	N	O	0
			65	55	1	4	5	
22	B	1	Total	C	Mg	N	O	0
			56	46	1	4	5	
22	B	1	Total	C	Mg	N	O	0
			55	45	1	4	5	
22	B	1	Total	C	Mg	N	O	0
			59	49	1	4	5	
22	B	1	Total	C	Mg	N	O	0
			65	55	1	4	5	
22	B	1	Total	C	Mg	N	O	0
			65	55	1	4	5	

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Mol	Chain	Residues	Atoms					AltConf
22	B	1	Total	C	Mg	N	O	0
			65	55	1	4	5	
22	B	1	Total	C	Mg	N	O	0
			65	55	1	4	5	
22	B	1	Total	C	Mg	N	O	0
			57	47	1	4	5	
22	B	1	Total	C	Mg	N	O	0
			65	55	1	4	5	
22	B	1	Total	C	Mg	N	O	0
			58	48	1	4	5	
22	B	1	Total	C	Mg	N	O	0
			50	40	1	4	5	
22	B	1	Total	C	Mg	N	O	0
			65	55	1	4	5	
22	B	1	Total	C	Mg	N	O	0
			58	48	1	4	5	
22	B	1	Total	C	Mg	N	O	0
			51	41	1	4	5	
22	B	1	Total	C	Mg	N	O	0
			45	35	1	4	5	
22	B	1	Total	C	Mg	N	O	0
			51	41	1	4	5	
22	B	1	Total	C	Mg	N	O	0
			65	55	1	4	5	
22	B	1	Total	C	Mg	N	O	0
			52	42	1	4	5	
22	B	1	Total	C	Mg	N	O	0
			65	55	1	4	5	
22	B	1	Total	C	Mg	N	O	0
			65	55	1	4	5	
22	B	1	Total	C	Mg	N	O	0
			65	55	1	4	5	
22	F	1	Total	C	Mg	N	O	0
			65	55	1	4	5	
22	F	1	Total	C	Mg	N	O	0
			65	55	1	4	5	
22	F	1	Total	C	Mg	N	O	0
			65	55	1	4	5	
22	F	1	Total	C	Mg	N	O	0
			45	35	1	4	5	
22	G	1	Total	C	Mg	N	O	0
			50	40	1	4	5	

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Mol	Chain	Residues	Atoms					AltConf
22	G	1	Total	C	Mg	N	O	0
			46	36	1	4	5	
22	J	1	Total	C	Mg	N	O	0
			42	34	1	4	3	
22	K	1	Total	C	Mg	N	O	0
			46	36	1	4	5	
22	K	1	Total	C	Mg	N	O	0
			55	45	1	4	5	
22	K	1	Total	C	Mg	N	O	0
			49	39	1	4	5	
22	K	1	Total	C	Mg	N	O	0
			55	45	1	4	5	
22	L	1	Total	C	Mg	N	O	0
			65	55	1	4	5	
22	L	1	Total	C	Mg	N	O	0
			65	55	1	4	5	
22	L	1	Total	C	Mg	N	O	0
			50	40	1	4	5	
22	1	1	Total	C	Mg	N	O	0
			65	55	1	4	5	
22	1	1	Total	C	Mg	N	O	0
			45	35	1	4	5	
22	1	1	Total	C	Mg	N	O	0
			65	55	1	4	5	
22	1	1	Total	C	Mg	N	O	0
			60	50	1	4	5	
22	1	1	Total	C	Mg	N	O	0
			55	45	1	4	5	
22	1	1	Total	C	Mg	N	O	0
			61	51	1	4	5	
22	1	1	Total	C	Mg	N	O	0
			60	50	1	4	5	
22	1	1	Total	C	Mg	N	O	0
			46	36	1	4	5	
22	1	1	Total	C	Mg	N	O	0
			65	55	1	4	5	
22	1	1	Total	C	Mg	N	O	0
			65	55	1	4	5	
22	1	1	Total	C	Mg	N	O	0
			46	36	1	4	5	
22	1	1	Total	C	Mg	N	O	0
			65	55	1	4	5	

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Mol	Chain	Residues	Atoms					AltConf
22	Z	1	Total	C	Mg	N	O	0
			65	55	1	4	5	
22	Z	1	Total	C	Mg	N	O	0
			45	35	1	4	5	
22	Z	1	Total	C	Mg	N	O	0
			56	46	1	4	5	
22	Z	1	Total	C	Mg	N	O	0
			60	50	1	4	5	
22	Z	1	Total	C	Mg	N	O	0
			55	45	1	4	5	
22	Z	1	Total	C	Mg	N	O	0
			61	51	1	4	5	
22	Z	1	Total	C	Mg	N	O	0
			60	50	1	4	5	
22	Z	1	Total	C	Mg	N	O	0
			50	40	1	4	5	
22	Z	1	Total	C	Mg	N	O	0
			65	55	1	4	5	
22	Z	1	Total	C	Mg	N	O	0
			51	41	1	4	5	
22	Z	1	Total	C	Mg	N	O	0
			46	36	1	4	5	
22	Z	1	Total	C	Mg	N	O	0
			65	55	1	4	5	
22	3	1	Total	C	Mg	N	O	0
			65	55	1	4	5	
22	3	1	Total	C	Mg	N	O	0
			46	36	1	4	5	
22	3	1	Total	C	Mg	N	O	0
			65	55	1	4	5	
22	3	1	Total	C	Mg	N	O	0
			60	50	1	4	5	
22	3	1	Total	C	Mg	N	O	0
			65	55	1	4	5	
22	3	1	Total	C	Mg	N	O	0
			65	55	1	4	5	
22	3	1	Total	C	Mg	N	O	0
			60	50	1	4	5	
22	3	1	Total	C	Mg	N	O	0
			45	35	1	4	5	
22	3	1	Total	C	Mg	N	O	0
			52	42	1	4	5	

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Mol	Chain	Residues	Atoms					AltConf
22	3	1	Total	C	Mg	N	O	0
			60	50	1	4	5	
22	3	1	Total	C	Mg	N	O	0
			55	45	1	4	5	
22	3	1	Total	C	Mg	N	O	0
			46	36	1	4	5	
22	3	1	Total	C	Mg	N	O	0
			65	55	1	4	5	
22	7	1	Total	C	Mg	N	O	0
			60	50	1	4	5	
22	7	1	Total	C	Mg	N	O	0
			50	40	1	4	5	
22	7	1	Total	C	Mg	N	O	0
			65	55	1	4	5	
22	7	1	Total	C	Mg	N	O	0
			65	55	1	4	5	
22	7	1	Total	C	Mg	N	O	0
			61	51	1	4	5	
22	7	1	Total	C	Mg	N	O	0
			56	46	1	4	5	
22	7	1	Total	C	Mg	N	O	0
			65	55	1	4	5	
22	7	1	Total	C	Mg	N	O	0
			43	35	1	4	3	
22	7	1	Total	C	Mg	N	O	0
			60	50	1	4	5	
22	7	1	Total	C	Mg	N	O	0
			50	40	1	4	5	
22	7	1	Total	C	Mg	N	O	0
			50	40	1	4	5	
22	7	1	Total	C	Mg	N	O	0
			42	34	1	4	3	
22	7	1	Total	C	Mg	N	O	0
			58	48	1	4	5	
22	7	1	Total	C	Mg	N	O	0
			55	45	1	4	5	
22	7	1	Total	C	Mg	N	O	0
			65	55	1	4	5	
22	7	1	Total	C	Mg	N	O	0
			60	50	1	4	5	
22	8	1	Total	C	Mg	N	O	0
			60	50	1	4	5	

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Mol	Chain	Residues	Atoms					AltConf
22	8	1	Total	C	Mg	N	O	0
			51	41	1	4	5	
22	8	1	Total	C	Mg	N	O	0
			65	55	1	4	5	
22	8	1	Total	C	Mg	N	O	0
			62	52	1	4	5	
22	8	1	Total	C	Mg	N	O	0
			65	55	1	4	5	
22	8	1	Total	C	Mg	N	O	0
			60	50	1	4	5	
22	8	1	Total	C	Mg	N	O	0
			50	40	1	4	5	
22	8	1	Total	C	Mg	N	O	0
			50	40	1	4	5	
22	8	1	Total	C	Mg	N	O	0
			46	36	1	4	5	
22	8	1	Total	C	Mg	N	O	0
			46	36	1	4	5	
22	4	1	Total	C	Mg	N	O	0
			60	50	1	4	5	
22	4	1	Total	C	Mg	N	O	0
			52	42	1	4	5	
22	4	1	Total	C	Mg	N	O	0
			65	55	1	4	5	
22	4	1	Total	C	Mg	N	O	0
			60	50	1	4	5	
22	4	1	Total	C	Mg	N	O	0
			51	41	1	4	5	
22	4	1	Total	C	Mg	N	O	0
			50	40	1	4	5	
22	4	1	Total	C	Mg	N	O	0
			55	45	1	4	5	
22	4	1	Total	C	Mg	N	O	0
			55	45	1	4	5	
22	4	1	Total	C	Mg	N	O	0
			50	40	1	4	5	
22	4	1	Total	C	Mg	N	O	0
			41	33	1	4	3	
22	4	1	Total	C	Mg	N	O	0
			51	41	1	4	5	
22	5	1	Total	C	Mg	N	O	0
			56	46	1	4	5	

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Mol	Chain	Residues	Atoms					AltConf
22	5	1	Total	C	Mg	N	O	0
			60	50	1	4	5	
22	5	1	Total	C	Mg	N	O	0
			61	51	1	4	5	
22	5	1	Total	C	Mg	N	O	0
			56	46	1	4	5	
22	5	1	Total	C	Mg	N	O	0
			65	55	1	4	5	
22	5	1	Total	C	Mg	N	O	0
			55	45	1	4	5	
22	5	1	Total	C	Mg	N	O	0
			50	40	1	4	5	
22	5	1	Total	C	Mg	N	O	0
			61	51	1	4	5	
22	5	1	Total	C	Mg	N	O	0
			45	35	1	4	5	
22	5	1	Total	C	Mg	N	O	0
			65	55	1	4	5	
22	5	1	Total	C	Mg	N	O	0
			65	55	1	4	5	
22	5	1	Total	C	Mg	N	O	0
			55	45	1	4	5	
22	5	1	Total	C	Mg	N	O	0
			50	40	1	4	5	
22	5	1	Total	C	Mg	N	O	0
			65	55	1	4	5	
22	5	1	Total	C	Mg	N	O	0
			46	36	1	4	5	
22	5	1	Total	C	Mg	N	O	0
			55	45	1	4	5	
22	6	1	Total	C	Mg	N	O	0
			60	50	1	4	5	
22	6	1	Total	C	Mg	N	O	0
			60	50	1	4	5	
22	6	1	Total	C	Mg	N	O	0
			60	50	1	4	5	
22	6	1	Total	C	Mg	N	O	0
			52	42	1	4	5	
22	6	1	Total	C	Mg	N	O	0
			65	55	1	4	5	
22	6	1	Total	C	Mg	N	O	0
			65	55	1	4	5	

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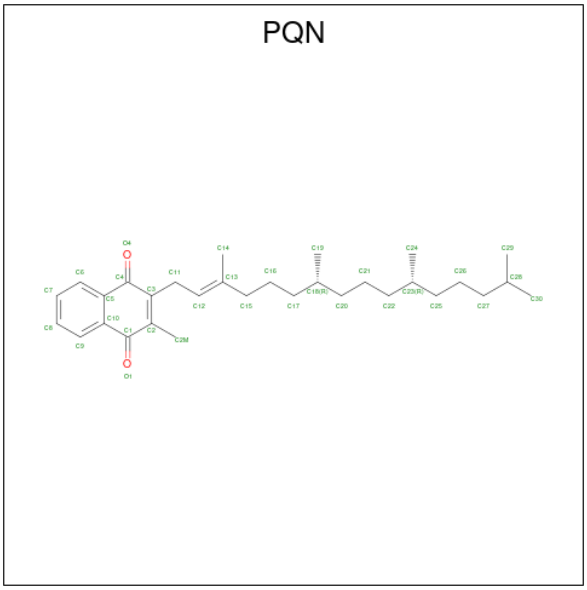
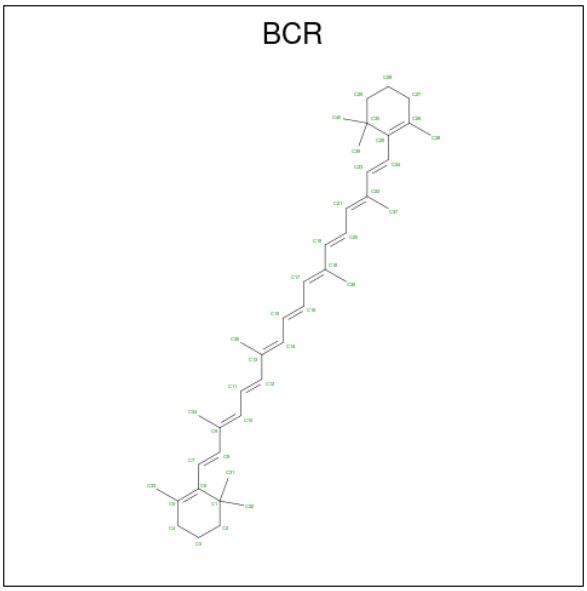
Mol	Chain	Residues	Atoms					AltConf
22	6	1	Total	C	Mg	N	O	0
			55	45	1	4	5	
22	6	1	Total	C	Mg	N	O	0
			65	55	1	4	5	
22	6	1	Total	C	Mg	N	O	0
			55	45	1	4	5	
22	6	1	Total	C	Mg	N	O	0
			65	55	1	4	5	
22	6	1	Total	C	Mg	N	O	0
			50	40	1	4	5	
22	6	1	Total	C	Mg	N	O	0
			61	51	1	4	5	
22	6	1	Total	C	Mg	N	O	0
			46	36	1	4	5	
22	6	1	Total	C	Mg	N	O	0
			65	55	1	4	5	
22	9	1	Total	C	Mg	N	O	0
			60	50	1	4	5	
22	9	1	Total	C	Mg	N	O	0
			45	35	1	4	5	
22	9	1	Total	C	Mg	N	O	0
			60	50	1	4	5	
22	9	1	Total	C	Mg	N	O	0
			62	52	1	4	5	
22	9	1	Total	C	Mg	N	O	0
			55	45	1	4	5	
22	9	1	Total	C	Mg	N	O	0
			50	40	1	4	5	
22	9	1	Total	C	Mg	N	O	0
			55	45	1	4	5	
22	9	1	Total	C	Mg	N	O	0
			45	35	1	4	5	
22	9	1	Total	C	Mg	N	O	0
			46	36	1	4	5	
22	9	1	Total	C	Mg	N	O	0
			50	40	1	4	5	
22	2	1	Total	C	Mg	N	O	0
			55	45	1	4	5	
22	2	1	Total	C	Mg	N	O	0
			45	35	1	4	5	
22	2	1	Total	C	Mg	N	O	0
			46	36	1	4	5	

- Molecule 23 is PHYLLOQUINONE (three-letter code: PQN) (formula: $C_{31}H_{46}O_2$).



Mol	Chain	Residues	Atoms	AltConf
23	A	1	Total C O 33 31 2	0
23	B	1	Total C O 33 31 2	0

- Molecule 24 is BETA-CAROTENE (three-letter code: BCR) (formula: $C_{40}H_{56}$).



Mol	Chain	Residues	Atoms	AltConf
24	A	1	Total C 40 40	0

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Mol	Chain	Residues	Atoms	AltConf
24	A	1	Total C 40 40	0
24	A	1	Total C 40 40	0
24	A	1	Total C 40 40	0
24	A	1	Total C 40 40	0
24	A	1	Total C 40 40	0
24	B	1	Total C 40 40	0
24	B	1	Total C 40 40	0
24	B	1	Total C 40 40	0
24	B	1	Total C 40 40	0
24	B	1	Total C 40 40	0
24	B	1	Total C 40 40	0
24	B	1	Total C 40 40	0
24	F	1	Total C 40 40	0
24	G	1	Total C 40 40	0
24	I	1	Total C 40 40	0
24	J	1	Total C 40 40	0
24	K	1	Total C 40 40	0
24	L	1	Total C 40 40	0
24	3	1	Total C 40 40	0
24	3	1	Total C 40 40	0
24	3	1	Total C 40 40	0

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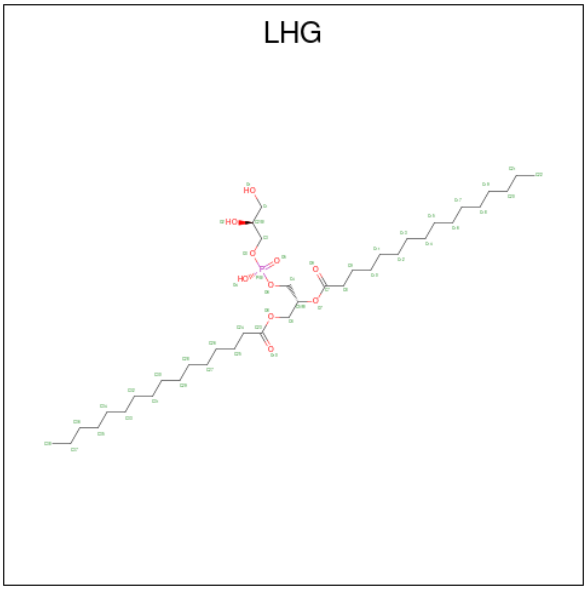
Mol	Chain	Residues	Atoms	AltConf
24	3	1	Total C 40 40	0
24	7	1	Total C 40 40	0
24	8	1	Total C 40 40	0
24	4	1	Total C 40 40	0
24	5	1	Total C 40 40	0
24	5	1	Total C 40 40	0
24	6	1	Total C 40 40	0
24	6	1	Total C 40 40	0

- Molecule 25 is 1,2-DIPALMITOYL-PHOSPHATIDYL-GLYCEROLE (three-letter code: LHG) (formula: $C_{38}H_{75}O_{10}P$).



Mol	Chain	Residues	Atoms	AltConf
25	A	1	Total C O P 35 24 10 1	0
25	A	1	Total C O P 49 38 10 1	0
25	B	1	Total C O P 33 22 10 1	0

Continued on next page...

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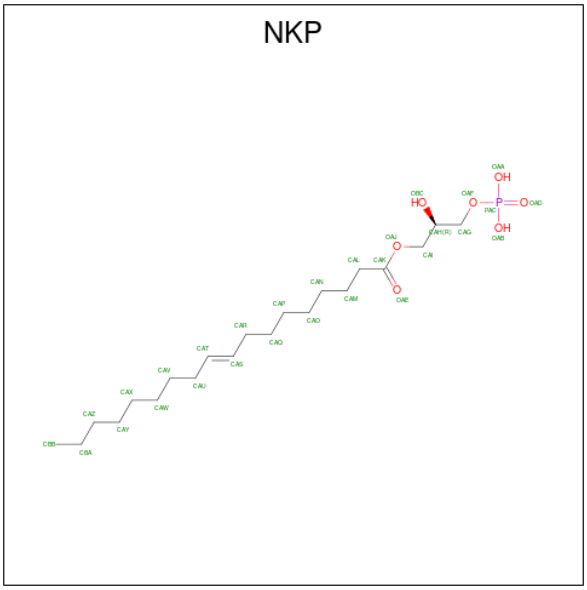
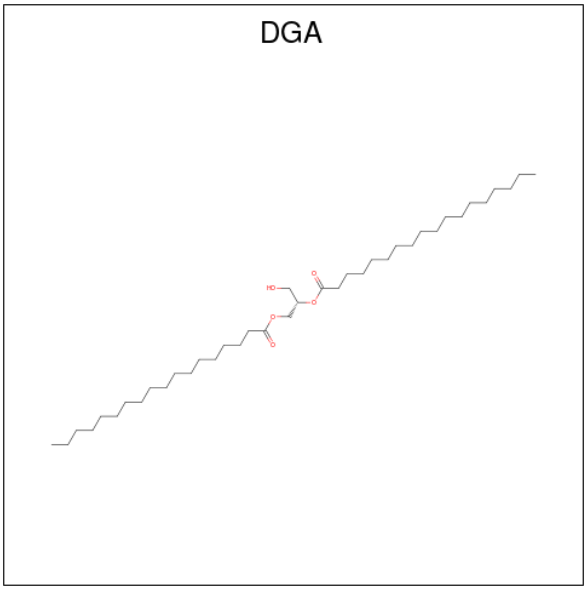
Mol	Chain	Residues	Atoms				AltConf
25	B	1	Total	C	O	P	0
			23	12	10	1	
25	B	1	Total	C	O	P	0
			20	9	10	1	
25	1	1	Total	C	O	P	0
			43	32	10	1	
25	Z	1	Total	C	O	P	0
			43	32	10	1	
25	3	1	Total	C	O	P	0
			20	9	10	1	
25	7	1	Total	C	O	P	0
			37	26	10	1	
25	8	1	Total	C	O	P	0
			38	27	10	1	
25	4	1	Total	C	O	P	0
			49	38	10	1	
25	4	1	Total	C	O	P	0
			32	21	10	1	
25	5	1	Total	C	O	P	0
			37	26	10	1	
25	6	1	Total	C	O	P	0
			49	38	10	1	
25	9	1	Total	C	O	P	0
			33	22	10	1	

- Molecule 26 is (2R)-2-hydroxy-3-(phosphonooxy)propyl (9E)-octadec-9-enoate (three-letter code: NKP) (formula: $C_{21}H_{41}O_7P$).



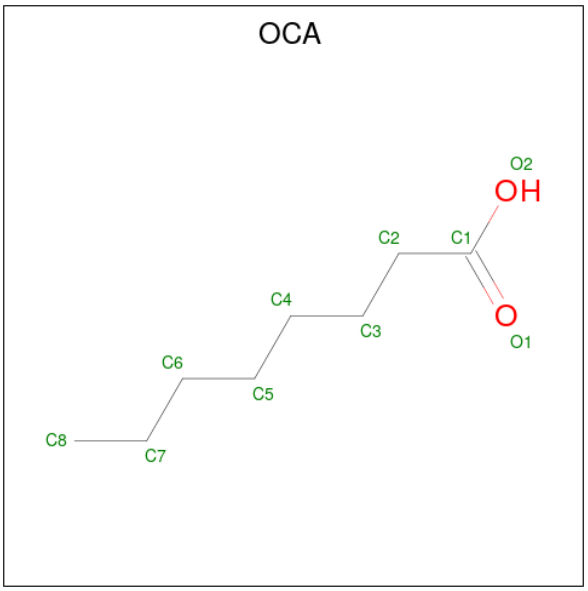
Mol	Chain	Residues	Atoms				AltConf
26	A	1	Total	C	O	P	0
			23	15	7	1	
26	8	1	Total	C	O	P	0
			29	21	7	1	

- Molecule 27 is DIACYL GLYCEROL (three-letter code: DGA) (formula: $C_{39}H_{76}O_5$).



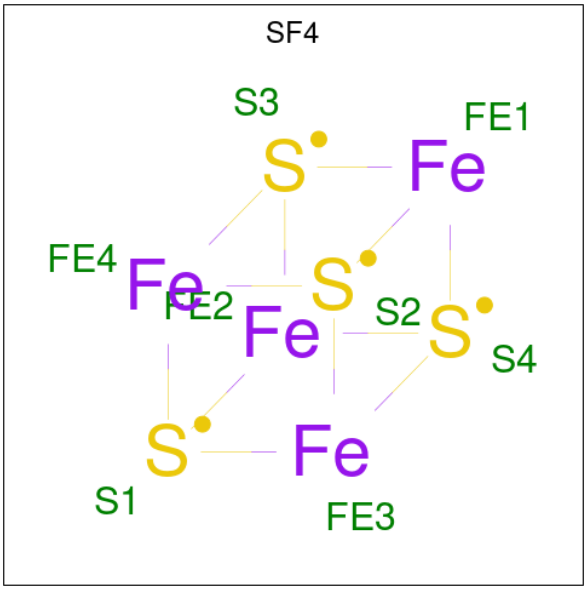
Mol	Chain	Residues	Atoms			AltConf
27	A	1	Total	C	O	0
			40	35	5	

- Molecule 28 is OCTANOIC ACID (CAPRYLIC ACID) (three-letter code: OCA) (formula: $C_8H_{16}O_2$).



Mol	Chain	Residues	Atoms			AltConf
28	A	1	Total	C	O	0
			10	8	2	

- Molecule 29 is IRON/SULFUR CLUSTER (three-letter code: SF4) (formula: Fe_4S_4).

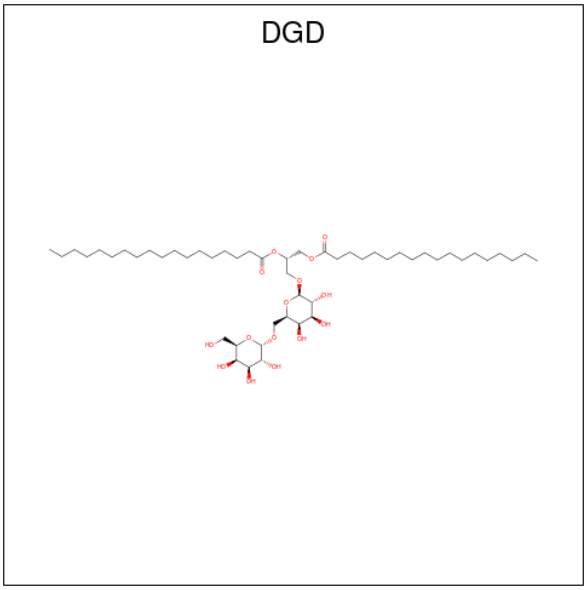


Mol	Chain	Residues	Atoms			AltConf
29	B	1	Total	Fe	S	0
			8	4	4	
29	C	1	Total	Fe	S	0
			8	4	4	
29	C	1	Total	Fe	S	0
			8	4	4	

- Molecule 30 is CALCIUM ION (three-letter code: CA) (formula: Ca).

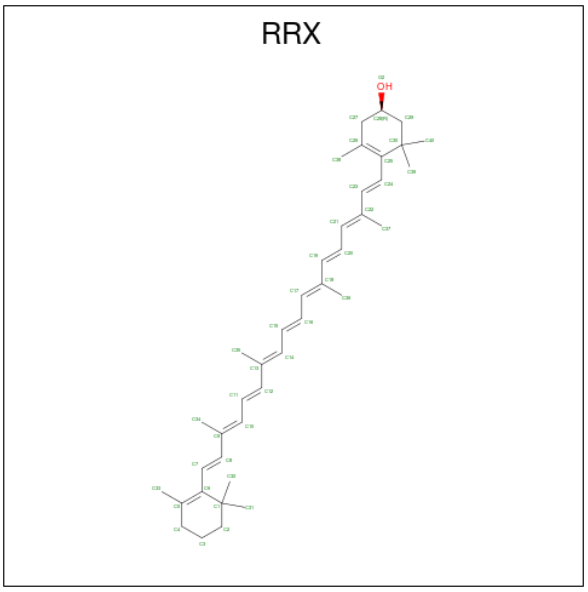
Mol	Chain	Residues	Atoms	AltConf
30	B	1	Total Ca 1 1	0

- Molecule 31 is DIGALACTOSYL DIACYL GLYCEROL (DGDG) (three-letter code: DGD) (formula: $C_{51}H_{96}O_{15}$).



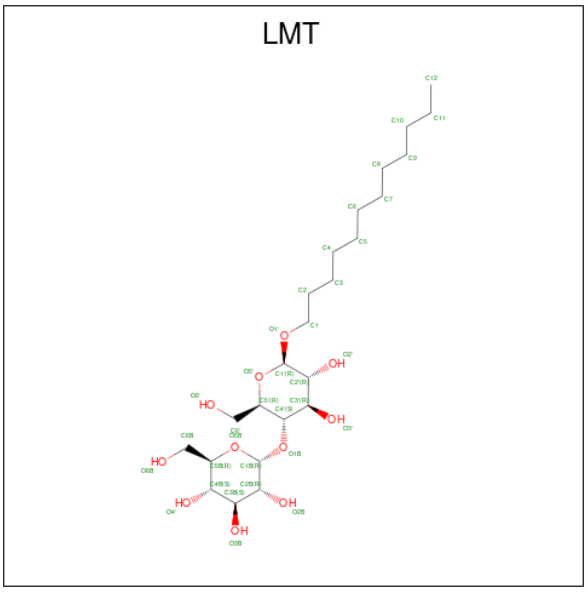
Mol	Chain	Residues	Atoms	AltConf
31	B	1	Total C O 61 46 15	0
31	1	1	Total C O 51 36 15	0
31	3	1	Total C O 51 36 15	0

- Molecule 32 is (3R)-beta,beta-caroten-3-ol (three-letter code: RRX) (formula: $C_{40}H_{56}O$).



Mol	Chain	Residues	Atoms			AltConf
32	F	1	Total	C	O	0
			41	40	1	

- Molecule 33 is DODECYL-BETA-D-MALTOSIDE (three-letter code: LMT) (formula: $C_{24}H_{46}O_{11}$).



Mol	Chain	Residues	Atoms			AltConf
33	F	1	Total	C	O	0
			35	24	11	
33	G	1	Total	C	O	0
			35	24	11	

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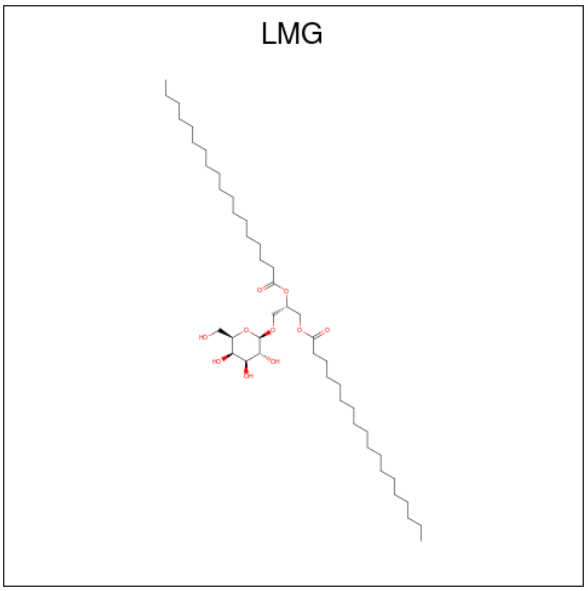
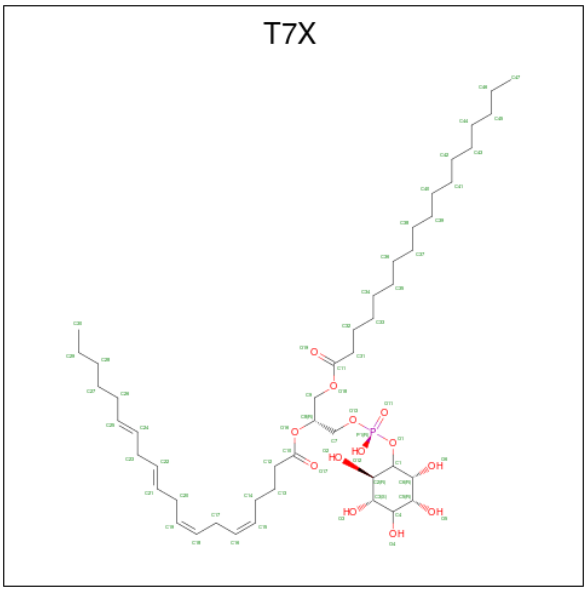
Mol	Chain	Residues	Atoms			AltConf
33	1	1	Total 35	C 24	O 11	0
33	4	1	Total 35	C 24	O 11	0
33	4	1	Total 35	C 24	O 11	0

- Molecule 34 is 1,2-DISTEAROYL-MONOGALACTOSYL-DIGLYCERIDE (three-letter code: LMG) (formula: $C_{45}H_{86}O_{10}$).



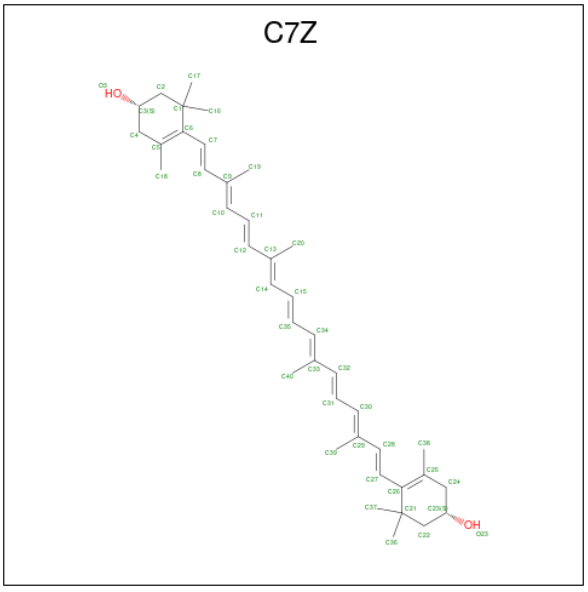
Mol	Chain	Residues	Atoms			AltConf
34	F	1	Total 35	C 25	O 10	0
34	J	1	Total 29	C 19	O 10	0

- Molecule 35 is Phosphatidylinositol (three-letter code: T7X) (formula: $C_{47}H_{83}O_{13}P$).



Mol	Chain	Residues	Atoms				AltConf
35	J	1	Total	C	O	P	0
			49	35	13	1	

- Molecule 36 is (1 {S})-3,5,5-trimethyl-4-[(1 {E},3 {E},5 {E},7 {E},9 {E},11 {E},13 {E},15 {E},17 {E})-3,7,12,16-tetramethyl-18-[(4 {S})-2,6,6-trimethyl-4-oxidanyl-cyclohexen-1-yl]octadeca-1,3,5,7,9,11,13,15,17-nonaenyl]cyclohex-3-en-1-ol (three-letter code: C7Z) (formula: $C_{40}H_{56}O_2$).



Mol	Chain	Residues	Atoms			AltConf
36	J	1	Total	C	O	0
			42	40	2	

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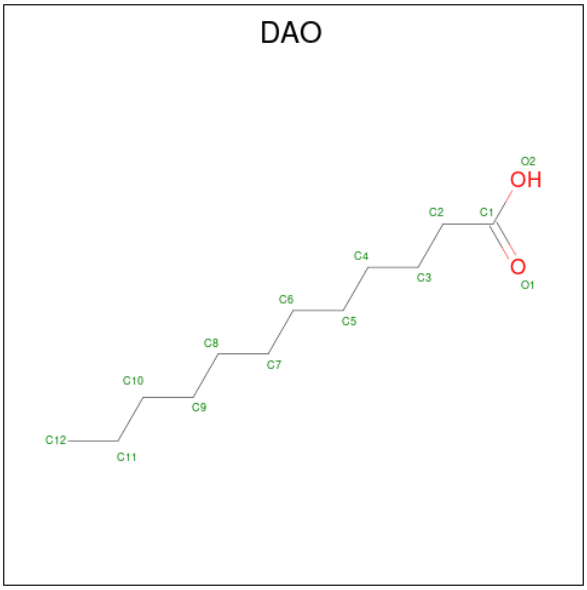
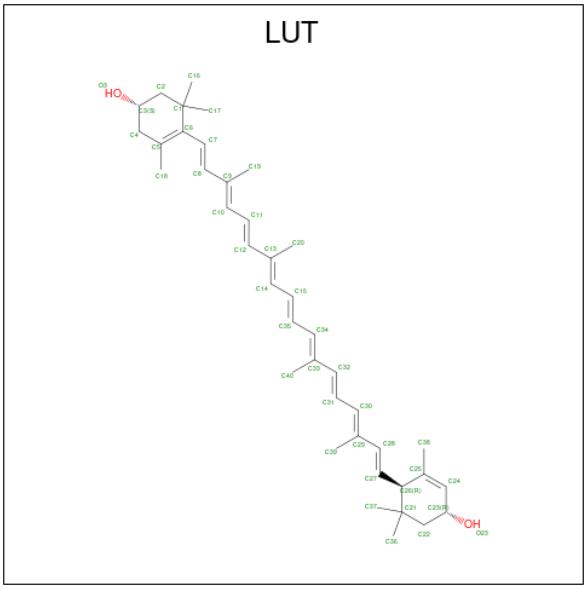
Mol	Chain	Residues	Atoms			AltConf
36	5	1	Total	C	O	0
			42	40	2	

- Molecule 37 is LAURIC ACID (three-letter code: DAO) (formula: $C_{12}H_{24}O_2$).



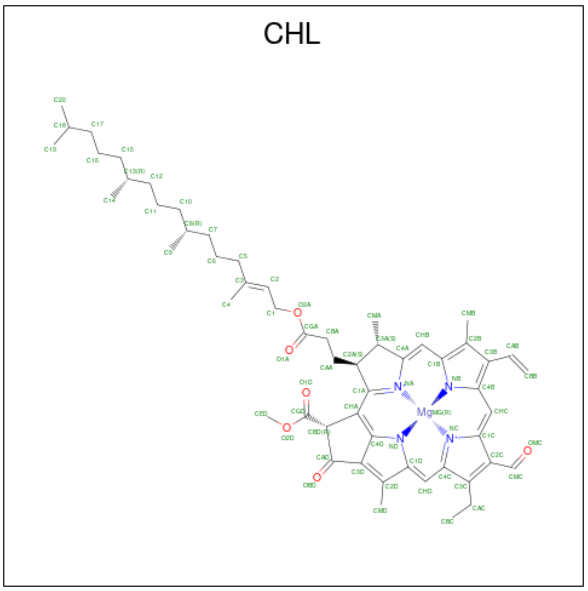
Mol	Chain	Residues	Atoms			AltConf
37	K	1	Total	C	O	0
			14	12	2	

- Molecule 38 is (3R,3'R,6S)-4,5-DIDEHYDRO-5,6-DIHYDRO-BETA,BETA-CAROTENE-3 ,3'-DIOL (three-letter code: LUT) (formula: $C_{40}H_{56}O_2$).



Mol	Chain	Residues	Atoms			AltConf
38	1	1	Total	C	O	0
			42	40	2	
38	1	1	Total	C	O	0
			42	40	2	
38	Z	1	Total	C	O	0
			42	40	2	
38	Z	1	Total	C	O	0
			42	40	2	
38	3	1	Total	C	O	0
			42	40	2	
38	3	1	Total	C	O	0
			42	40	2	
38	7	1	Total	C	O	0
			42	40	2	
38	7	1	Total	C	O	0
			42	40	2	
38	8	1	Total	C	O	0
			42	40	2	
38	8	1	Total	C	O	0
			42	40	2	
38	4	1	Total	C	O	0
			42	40	2	
38	4	1	Total	C	O	0
			42	40	2	
38	5	1	Total	C	O	0
			42	40	2	
38	5	1	Total	C	O	0
			42	40	2	
38	6	1	Total	C	O	0
			42	40	2	
38	6	1	Total	C	O	0
			42	40	2	
38	9	1	Total	C	O	0
			42	40	2	
38	9	1	Total	C	O	0
			42	40	2	

- Molecule 39 is CHLOROPHYLL B (three-letter code: CHL) (formula: $C_{55}H_{70}MgN_4O_6$).



Mol	Chain	Residues	Atoms					AltConf
39	1	1	Total	C	Mg	N	O	0
			48	37	1	4	6	
39	Z	1	Total	C	Mg	N	O	0
			51	40	1	4	6	
39	Z	1	Total	C	Mg	N	O	0
			48	37	1	4	6	
39	3	1	Total	C	Mg	N	O	0
			66	55	1	4	6	
39	7	1	Total	C	Mg	N	O	0
			54	43	1	4	6	
39	8	1	Total	C	Mg	N	O	0
			58	47	1	4	6	
39	8	1	Total	C	Mg	N	O	0
			56	45	1	4	6	
39	8	1	Total	C	Mg	N	O	0
			66	55	1	4	6	
39	4	1	Total	C	Mg	N	O	0
			51	40	1	4	6	
39	4	1	Total	C	Mg	N	O	0
			51	40	1	4	6	
39	4	1	Total	C	Mg	N	O	0
			47	36	1	4	6	
39	4	1	Total	C	Mg	N	O	0
			43	34	1	4	4	
39	5	1	Total	C	Mg	N	O	0
			66	55	1	4	6	
39	5	1	Total	C	Mg	N	O	0
			51	40	1	4	6	

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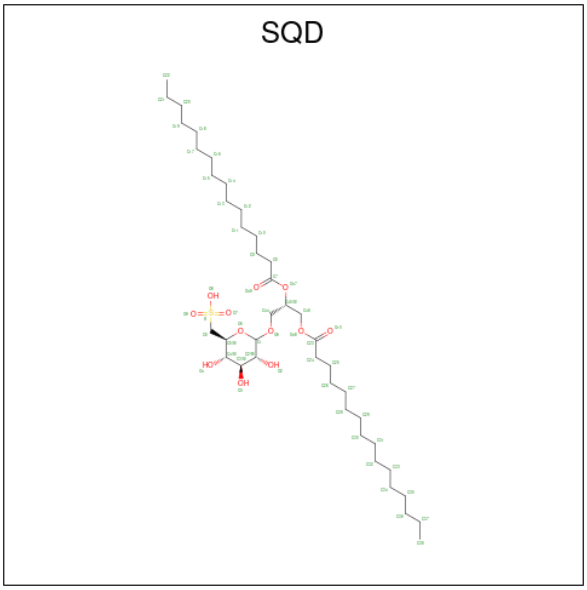
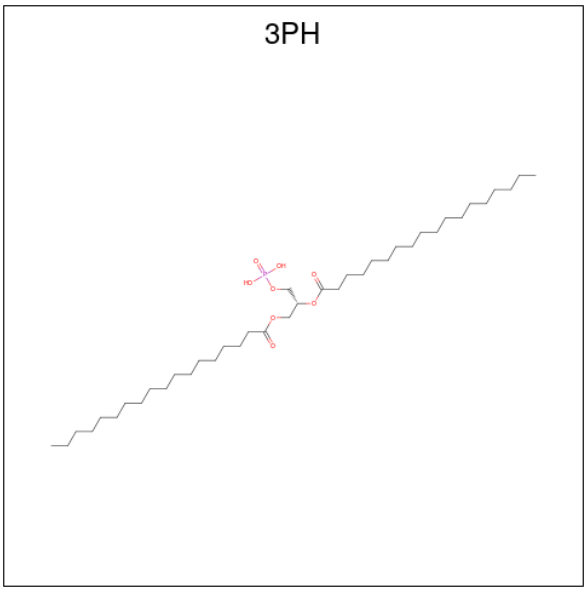
Mol	Chain	Residues	Atoms					AltConf
39	5	1	Total	C	Mg	N	O	0
			43	34	1	4	4	
39	6	1	Total	C	Mg	N	O	0
			56	45	1	4	6	
39	6	1	Total	C	Mg	N	O	0
			51	40	1	4	6	
39	6	1	Total	C	Mg	N	O	0
			56	45	1	4	6	
39	6	1	Total	C	Mg	N	O	0
			43	34	1	4	4	
39	9	1	Total	C	Mg	N	O	0
			66	55	1	4	6	
39	9	1	Total	C	Mg	N	O	0
			42	33	1	4	4	
39	2	1	Total	C	Mg	N	O	0
			51	40	1	4	6	

- Molecule 40 is 1,2-DI-O-ACYL-3-O-[6-DEOXY-6-SULFO-ALPHA-D-GLUCOPYRANOSYL]-SN-GLYCEROL (three-letter code: SQD) (formula: $C_{41}H_{78}O_{12}S$).



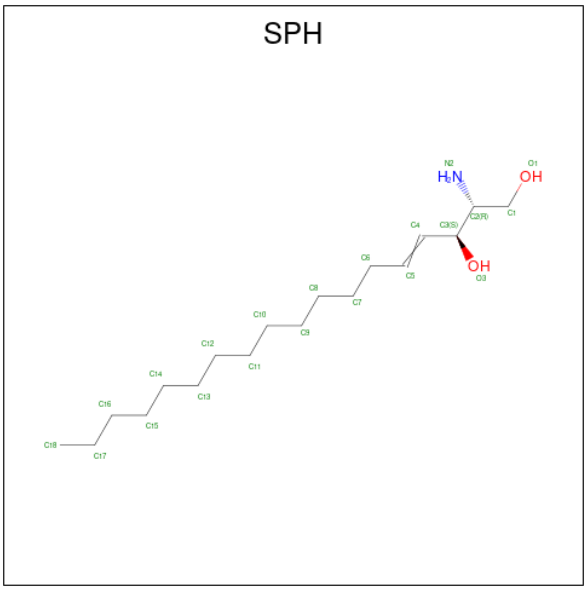
Mol	Chain	Residues	Atoms				AltConf
40	1	1	Total	C	O	S	0
			48	35	12	1	

- Molecule 41 is 1,2-DIACYL-GLYCEROL-3-SN-PHOSPHATE (three-letter code: 3PH) (formula: $C_{39}H_{77}O_8P$).



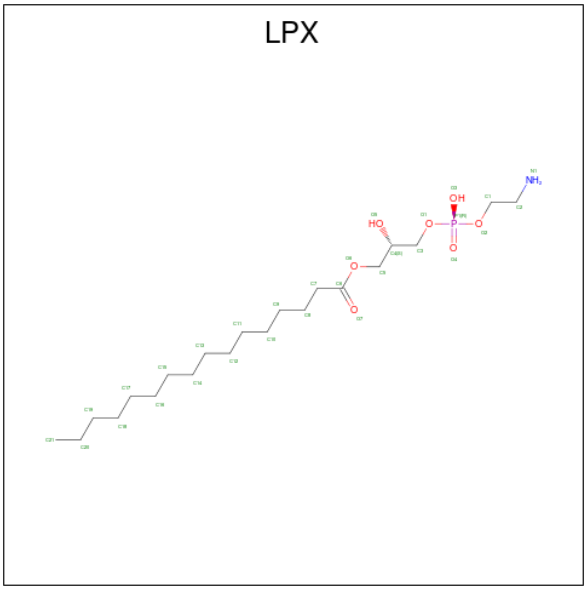
Mol	Chain	Residues	Atoms				AltConf
41	7	1	Total	C	O	P	0
			39	30	8	1	
41	8	1	Total	C	O	P	0
			30	21	8	1	
41	5	1	Total	C	O	P	0
			23	14	8	1	
41	6	1	Total	C	O	P	0
			29	20	8	1	

- Molecule 42 is SPHINGOSINE (three-letter code: SPH) (formula: $C_{18}H_{37}NO_2$).



Mol	Chain	Residues	Atoms	AltConf
42	7	1	Total C N O 21 18 1 2	0
42	7	1	Total C N O 21 18 1 2	0

- Molecule 43 is (2S)-3-{[(R)-(2-aminoethoxy)(hydroxy)phosphoryl]oxy}-2-hydroxypropyl hexadecanoate (three-letter code: LPX) (formula: $C_{21}H_{44}NO_7P$).



Mol	Chain	Residues	Atoms	AltConf
43	8	1	Total C N O P 30 21 1 7 1	0

- Molecule 44 is water.

Mol	Chain	Residues	Atoms	AltConf
44	A	19	Total O 19 19	0
44	B	17	Total O 17 17	0
44	F	3	Total O 3 3	0
44	I	1	Total O 1 1	0
44	J	1	Total O 1 1	0
44	K	1	Total O 1 1	0

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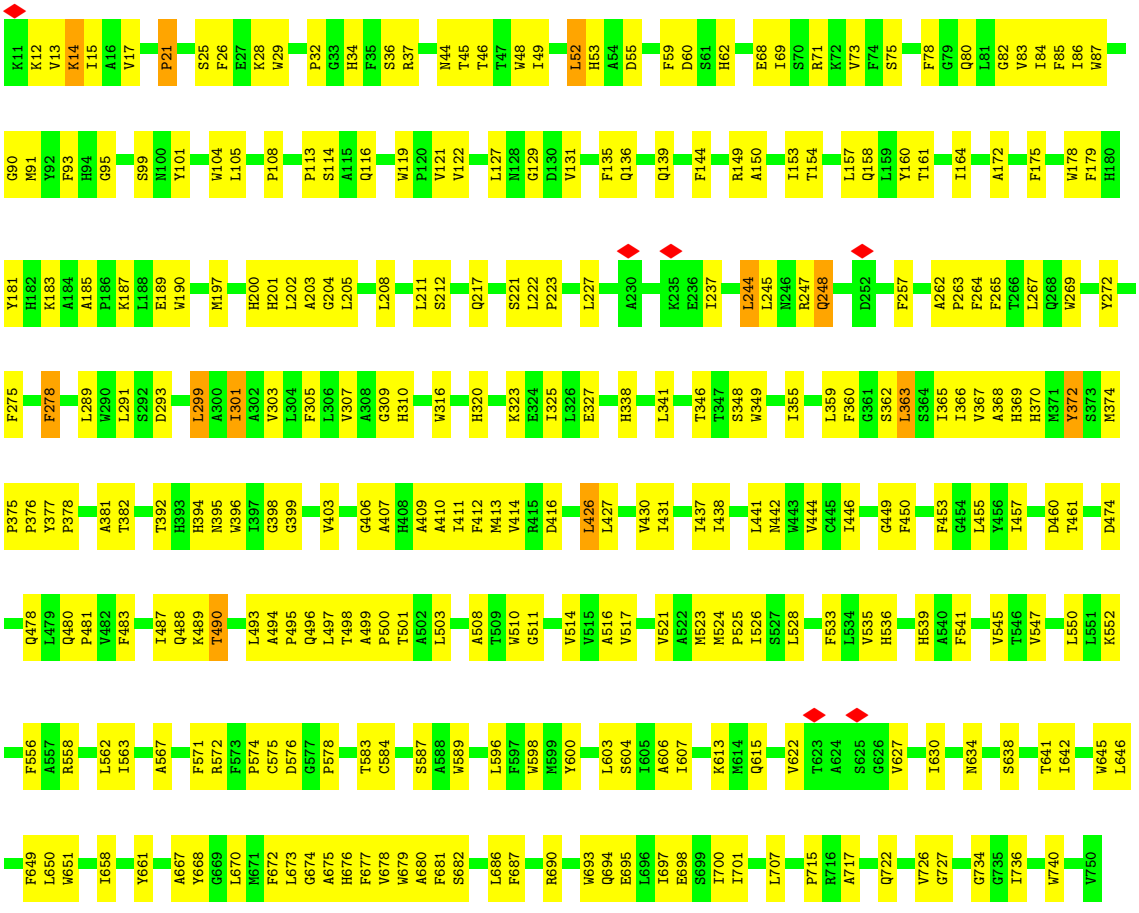
Mol	Chain	Residues	Atoms	AltConf
44	1	1	Total O 1 1	0
44	Z	1	Total O 1 1	0
44	3	2	Total O 2 2	0
44	7	4	Total O 4 4	0
44	8	4	Total O 4 4	0
44	4	5	Total O 5 5	0
44	5	5	Total O 5 5	0
44	6	3	Total O 3 3	0
44	2	1	Total O 1 1	0

3 Residue-property plots (i)

These plots are drawn for all protein, RNA, DNA and oligosaccharide chains in the entry. The first graphic for a chain summarises the proportions of the various outlier classes displayed in the second graphic. The second graphic shows the sequence view annotated by issues in geometry and atom inclusion in map density. Residues are color-coded according to the number of geometric quality criteria for which they contain at least one outlier: green = 0, yellow = 1, orange = 2 and red = 3 or more. A red diamond above a residue indicates a poor fit to the EM map for this residue (all-atom inclusion < 40%). Stretches of 2 or more consecutive residues without any outlier are shown as a green connector. Residues present in the sample, but not in the model, are shown in grey.

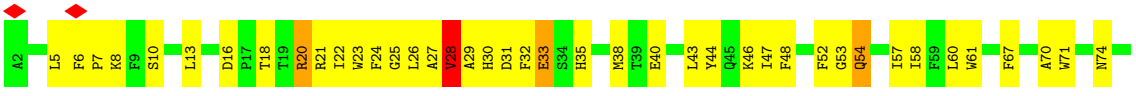
• Molecule 1: Photosystem I P700 chlorophyll a apoprotein A1 PsaA

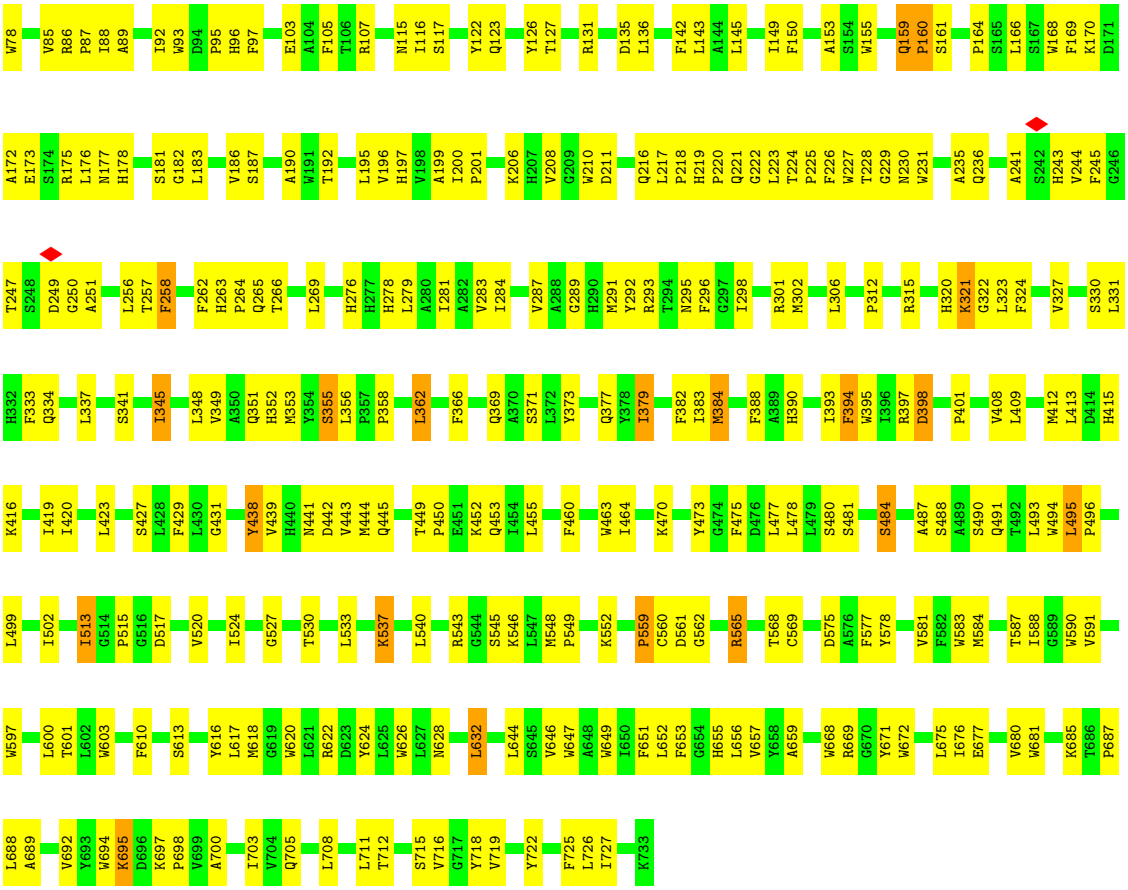
Chain A:



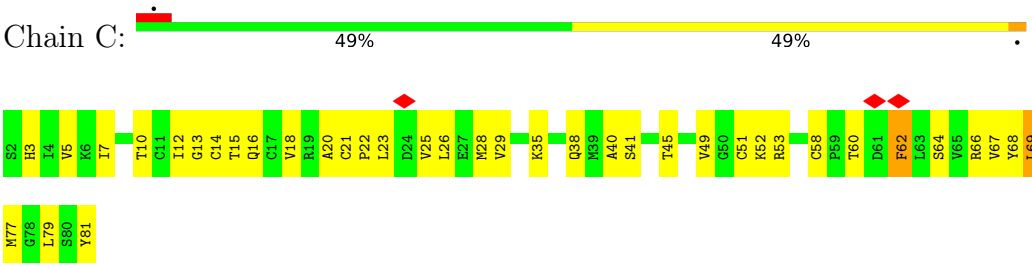
• Molecule 2: Photosystem I P700 chlorophyll a apoprotein A2 PsaB

Chain B:

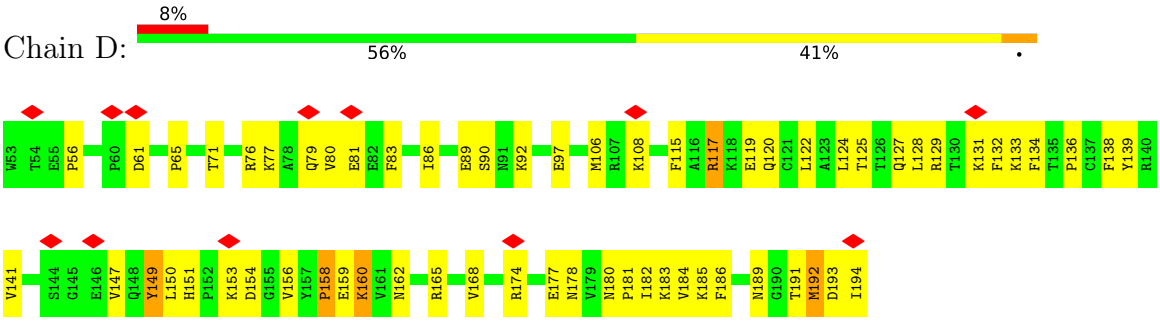




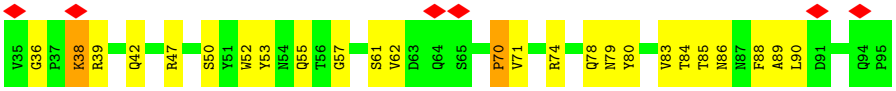
• Molecule 3: Photosystem I subunit VII PsaC



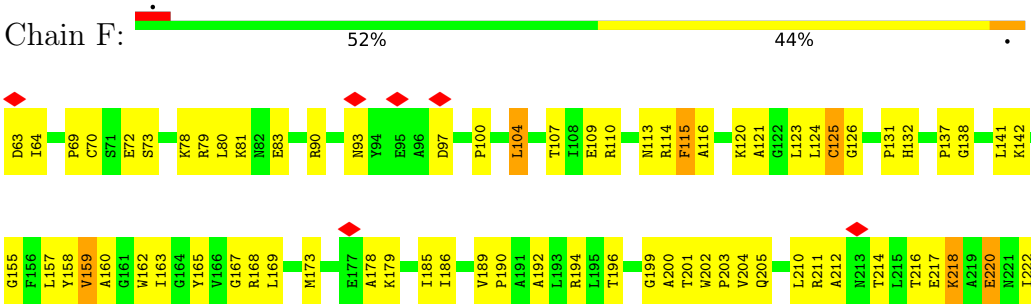
• Molecule 4: Photosystem I reaction center subunit II PsaD



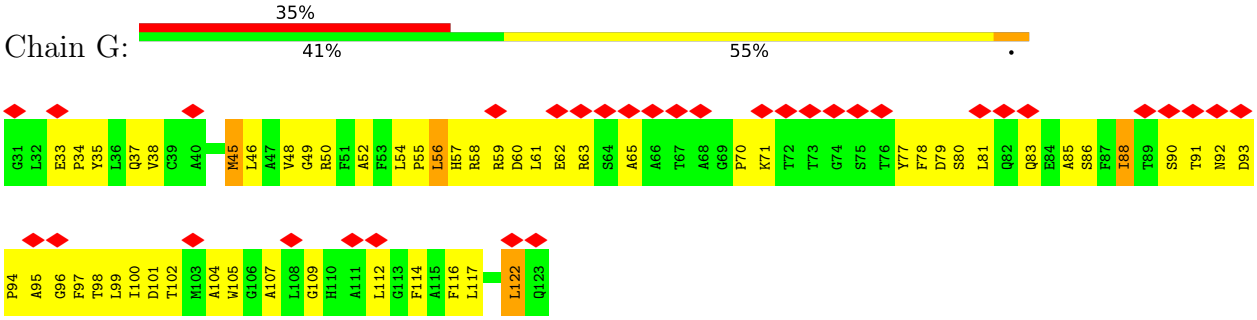
• Molecule 5: Photosystem I reaction center subunit IV PsaE



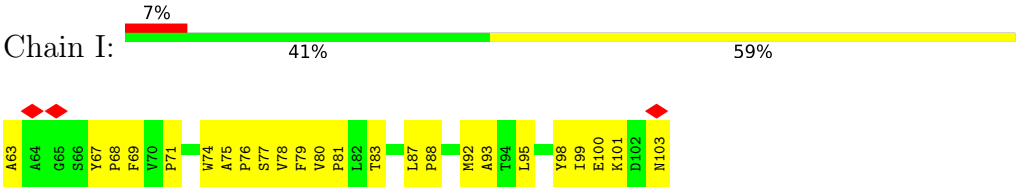
• Molecule 6: Photosystem I reaction center subunit III PsaF



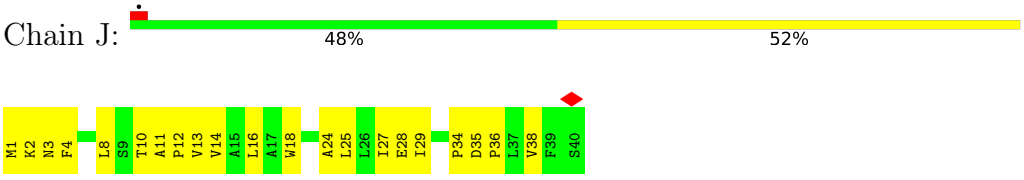
• Molecule 7: Photosystem I reaction center subunit V PsaG



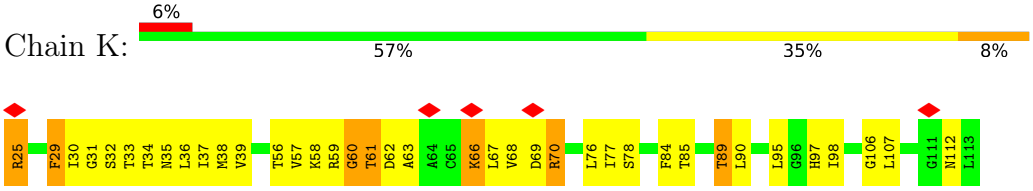
• Molecule 8: Photosystem I reaction center subunit VIII PsaI



• Molecule 9: Photosystem I subunit IX PsaJ

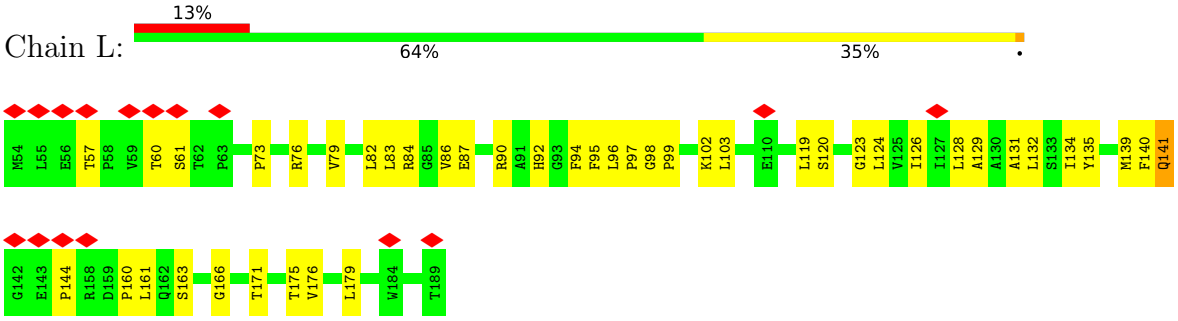


• Molecule 10: Photosystem I reaction center subunit X psaK



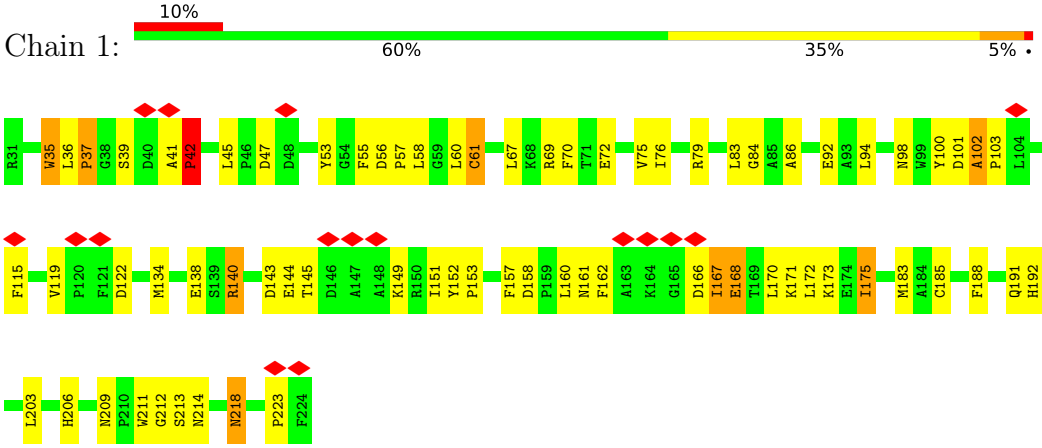
• Molecule 11: Photosystem I reaction centre subunit XI PsaL

Chain L:



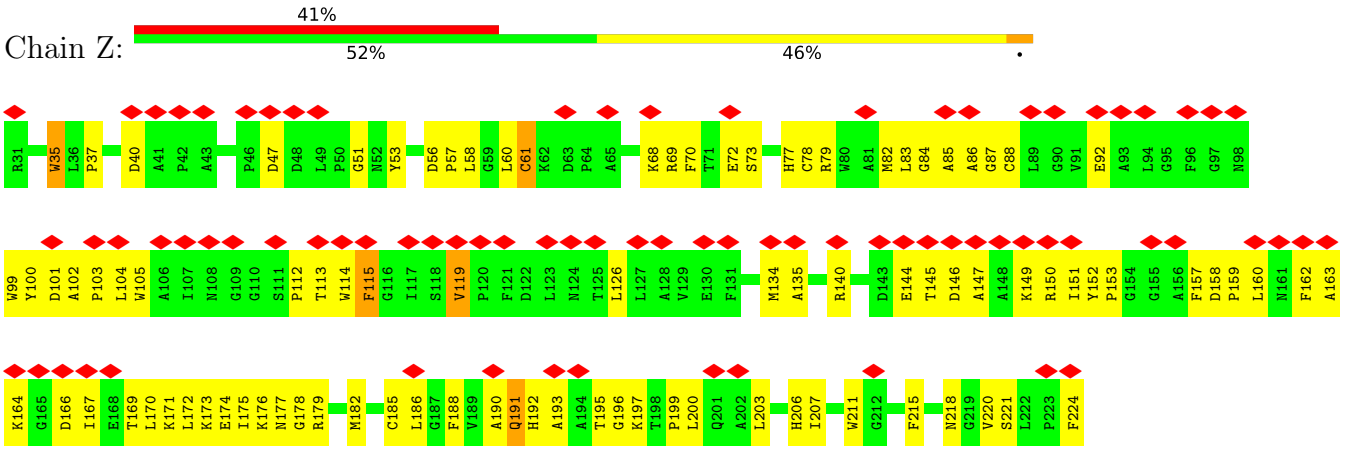
• Molecule 12: Light-harvesting protein of photosystem I Lhca1

Chain 1:



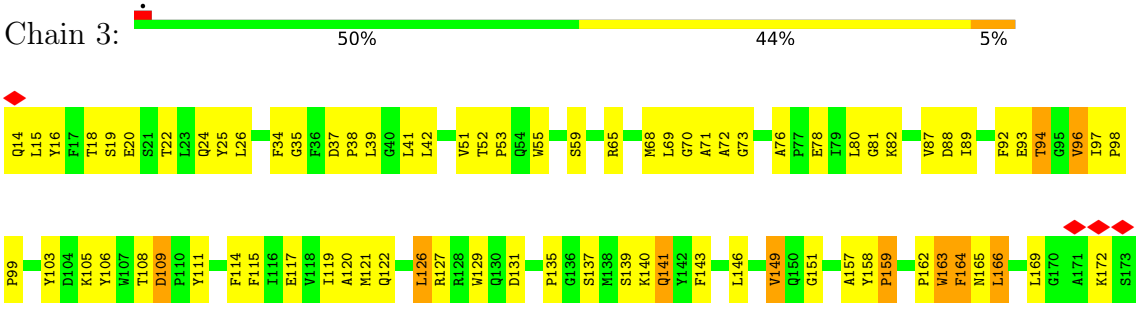
• Molecule 12: Light-harvesting protein of photosystem I Lhca1

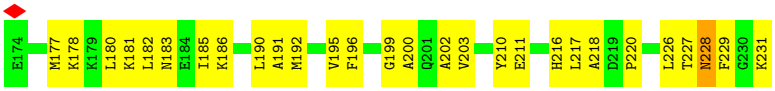
Chain Z:



• Molecule 13: Light-harvesting protein of photosystem I Lhca3

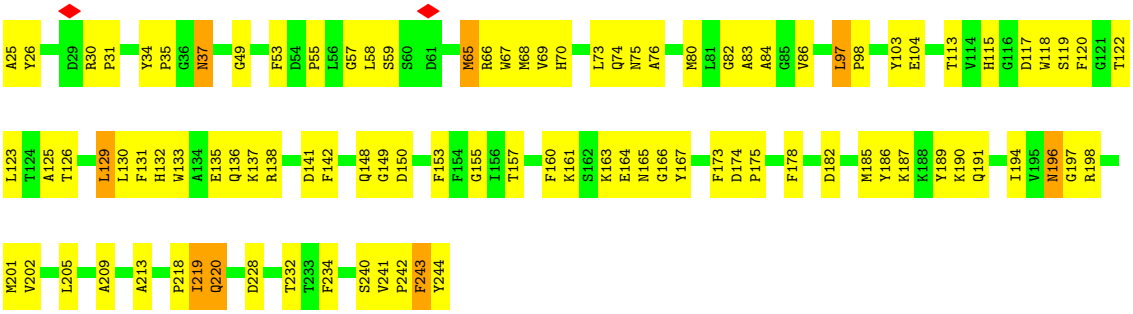
Chain 3:





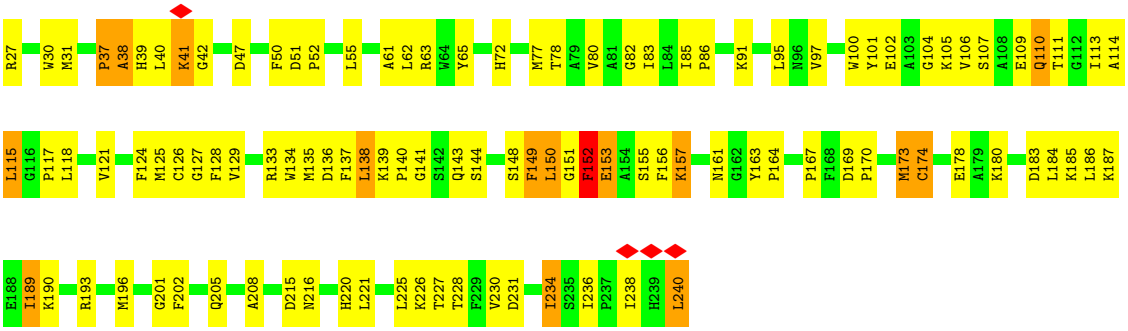
• Molecule 14: Light-harvesting protein of photosystem I Lhca7

Chain 7:



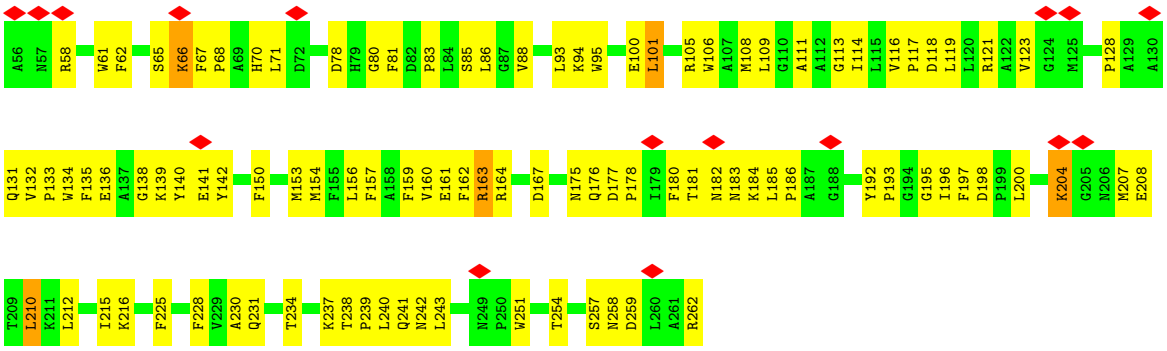
• Molecule 15: Light-harvesting protein of photosystem I Lhca8

Chain 8:



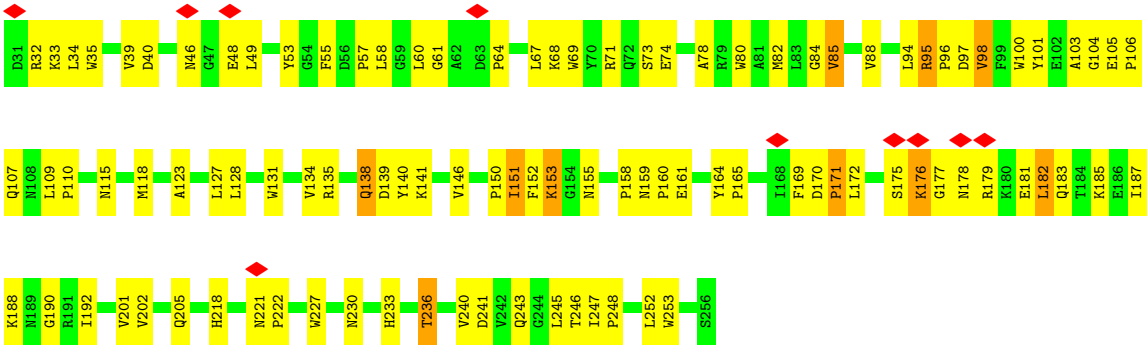
• Molecule 16: Light-harvesting protein of photosystem I Lhca4

Chain 4:



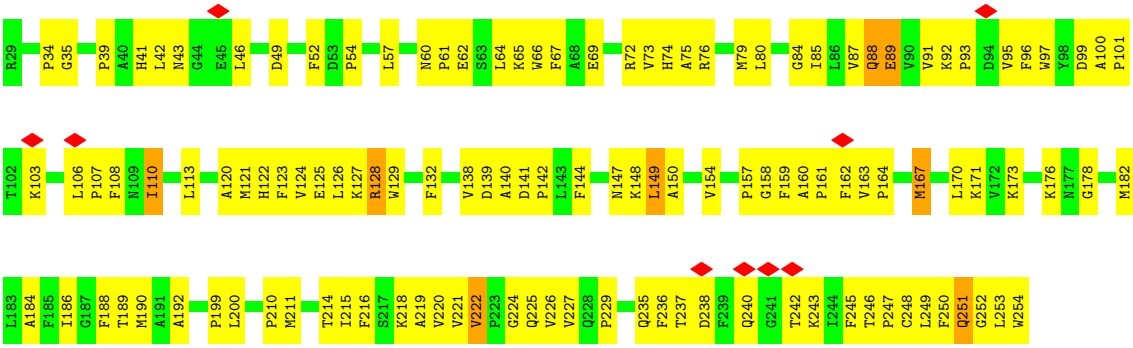
• Molecule 17: Light-harvesting protein of photosystem I Lhca5

Chain 5:



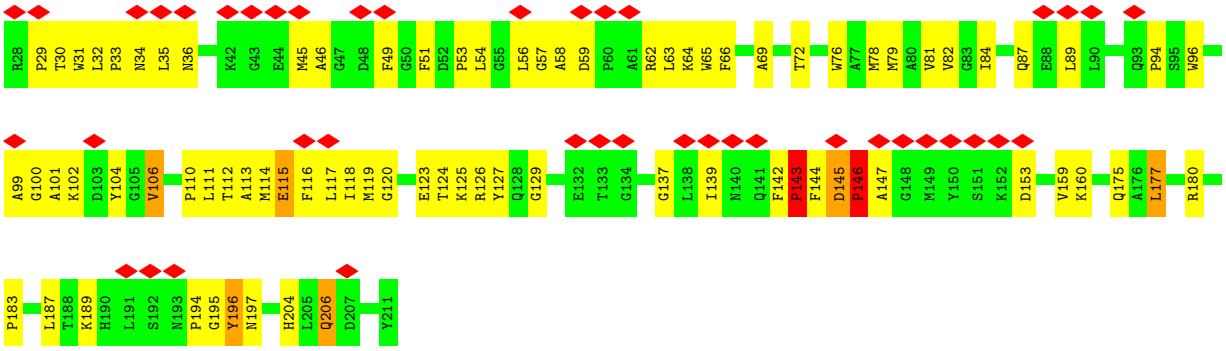
• Molecule 18: Light-harvesting protein of photosystem I Lhca6

Chain 6:



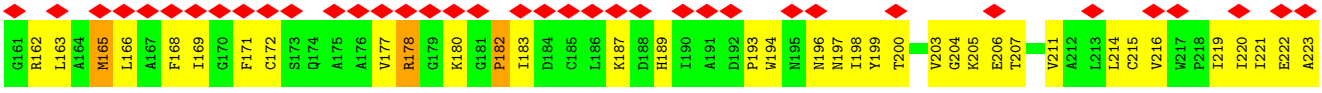
• Molecule 19: Light-harvesting protein of photosystem I Lhca9

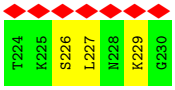
Chain 9:



• Molecule 20: Light-harvesting protein of photosystem I Lhca2 partial

Chain 2:




T224
K225
S226
L227
N228
K229
G230

4 Experimental information (i)

Property	Value	Source
EM reconstruction method	SINGLE PARTICLE	Depositor
Imposed symmetry	POINT, C1	Depositor
Number of particles used	167160	Depositor
Resolution determination method	FSC 0.143 CUT-OFF	Depositor
CTF correction method	NONE	Depositor
Microscope	FEI TITAN KRIOS	Depositor
Voltage (kV)	300	Depositor
Electron dose ($e^-/Å^2$)	50	Depositor
Minimum defocus (nm)	150	Depositor
Maximum defocus (nm)	1750	Depositor
Magnification	130000	Depositor
Image detector	GATAN K3 BIOQUANTUM (6k x 4k)	Depositor
Maximum map value	0.032	Depositor
Minimum map value	-0.012	Depositor
Average map value	0.000	Depositor
Map value standard deviation	0.001	Depositor
Recommended contour level	0.007	Depositor
Map size (Å)	325.5, 325.5, 325.5	wwPDB
Map dimensions	500, 500, 500	wwPDB
Map angles (°)	90.0, 90.0, 90.0	wwPDB
Pixel spacing (Å)	0.651, 0.651, 0.651	Depositor

5 Model quality

5.1 Standard geometry

Bond lengths and bond angles in the following residue types are not validated in this section: 3PH, OCA, RRX, LHG, PQN, NKP, C7Z, DGD, BCR, LMT, LMG, LPX, SQD, CA, SPH, CL0, CLA, LUT, SF4, CHL, T7X, DGA, DAO

The Z score for a bond length (or angle) is the number of standard deviations the observed value is removed from the expected value. A bond length (or angle) with $|Z| > 5$ is considered an outlier worth inspection. RMSZ is the root-mean-square of all Z scores of the bond lengths (or angles).

Mol	Chain	Bond lengths		Bond angles	
		RMSZ	#$\|Z\|$ >5	RMSZ	#$\|Z\|$ >5
1	A	0.61	0/6016	0.61	0/8200
2	B	0.62	0/6010	0.60	0/8207
3	C	0.65	0/613	0.63	0/830
4	D	0.59	0/1159	0.69	0/1569
5	E	0.64	0/495	0.60	0/675
6	F	0.61	0/1292	0.68	0/1755
7	G	0.63	0/706	0.64	0/960
8	I	0.62	0/320	0.57	0/443
9	J	0.63	0/329	0.60	0/452
10	K	0.66	0/625	0.70	0/848
11	L	0.64	0/919	0.70	0/1253
12	1	0.61	0/1501	0.72	1/2045 (0.0%)
12	Z	0.54	0/1501	0.56	0/2045
13	3	0.61	0/1747	0.67	1/2376 (0.0%)
14	7	0.61	0/1765	0.67	0/2407
15	8	0.61	0/1643	0.66	0/2226
16	4	0.61	0/1648	0.62	0/2240
17	5	0.62	0/1819	0.66	0/2480
18	6	0.60	0/1824	0.60	0/2488
19	9	0.57	0/1456	0.77	3/1975 (0.2%)
20	2	0.64	0/535	0.64	0/726
All	All	0.61	0/33923	0.64	5/46200 (0.0%)

There are no bond length outliers.

All (5) bond angle outliers are listed below:

Mol	Chain	Res	Type	Atoms	Z	Observed(^{o})	Ideal(^{o})
19	9	143	PRO	CA-N-CD	-11.42	95.51	111.50
12	1	42	PRO	CA-N-CD	-9.60	98.06	111.50

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Mol	Chain	Res	Type	Atoms	Z	Observed(^{o})	Ideal(^{o})
19	9	146	PRO	CA-N-CD	-7.25	101.35	111.50
13	3	164	PHE	CB-CA-C	5.17	120.74	110.40
19	9	146	PRO	N-CA-CB	-5.16	96.92	102.60

There are no chirality outliers.

There are no planarity outliers.

5.2 Too-close contacts (i)

In the following table, the Non-H and H(model) columns list the number of non-hydrogen atoms and hydrogen atoms in the chain respectively. The H(added) column lists the number of hydrogen atoms added and optimized by MolProbity. The Clashes column lists the number of clashes within the asymmetric unit, whereas Symm-Clashes lists symmetry-related clashes.

Mol	Chain	Non-H	H(model)	H(added)	Clashes	Symm-Clashes
1	A	5819	0	5666	308	0
2	B	5800	0	5556	367	0
3	C	603	0	587	41	0
4	D	1129	0	1127	46	0
5	E	484	0	477	24	0
6	F	1265	0	1292	74	0
7	G	690	0	671	67	0
8	I	308	0	310	33	0
9	J	319	0	327	25	0
10	K	617	0	635	53	0
11	L	897	0	891	44	0
12	1	1457	0	1406	82	0
12	Z	1457	0	1406	126	0
13	3	1695	0	1641	136	0
14	7	1705	0	1601	107	0
15	8	1602	0	1600	132	0
16	4	1598	0	1569	120	0
17	5	1764	0	1709	125	0
18	6	1762	0	1740	182	0
19	9	1416	0	1392	100	0
20	2	525	0	539	44	0
21	A	65	0	72	9	0
22	1	698	0	678	128	0
22	2	146	0	112	34	0
22	3	749	0	722	166	0
22	4	590	0	511	114	0

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Mol	Chain	Non-H	H(model)	H(added)	Clashes	Symm-Clashes
22	5	910	0	851	218	0
22	6	824	0	800	213	0
22	7	905	0	848	154	0
22	8	555	0	502	123	0
22	9	528	0	451	72	0
22	A	2640	0	2709	558	0
22	B	2340	0	2333	499	0
22	F	240	0	245	69	0
22	G	96	0	70	28	0
22	J	42	0	30	4	0
22	K	205	0	166	28	0
22	L	180	0	180	40	0
22	Z	679	0	631	166	0
23	A	33	0	46	9	0
23	B	33	0	46	5	0
24	3	160	0	224	61	0
24	4	40	0	56	27	0
24	5	80	0	112	33	0
24	6	80	0	112	35	0
24	7	40	0	56	15	0
24	8	40	0	56	15	0
24	A	240	0	336	99	0
24	B	280	0	392	105	0
24	F	40	0	56	16	0
24	G	40	0	56	13	0
24	I	40	0	56	12	0
24	J	40	0	56	19	0
24	K	40	0	55	11	0
24	L	40	0	56	17	0
25	1	43	0	56	8	0
25	3	20	0	12	1	0
25	4	81	0	108	20	0
25	5	37	0	44	7	0
25	6	49	0	74	12	0
25	7	37	0	44	10	0
25	8	38	0	46	7	0
25	9	33	0	36	2	0
25	A	84	0	114	14	0
25	B	76	0	64	14	0
25	Z	43	0	56	9	0
26	8	29	0	39	6	0
26	A	23	0	24	3	0

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Mol	Chain	Non-H	H(model)	H(added)	Clashes	Symm-Clashes
27	A	40	0	65	20	0
28	A	10	0	15	0	0
29	B	8	0	0	9	0
29	C	16	0	0	9	0
30	B	1	0	0	0	0
31	1	51	0	60	10	0
31	3	51	0	60	21	0
31	B	61	0	83	17	0
32	F	41	0	56	19	0
33	1	35	0	46	9	0
33	4	70	0	92	10	0
33	F	35	0	46	5	0
33	G	35	0	46	9	0
34	F	35	0	40	1	0
34	J	29	0	28	11	0
35	J	49	0	0	2	0
36	5	42	0	0	2	0
36	J	42	0	0	2	0
37	K	14	0	23	8	0
38	1	84	0	112	31	0
38	3	84	0	112	32	0
38	4	84	0	112	35	0
38	5	84	0	112	32	0
38	6	84	0	112	29	0
38	7	84	0	112	33	0
38	8	84	0	112	31	0
38	9	84	0	112	32	0
38	Z	84	0	112	39	0
39	1	48	0	32	9	0
39	2	51	0	36	6	0
39	3	66	0	69	13	0
39	4	192	0	130	47	0
39	5	160	0	134	32	0
39	6	206	0	156	53	0
39	7	54	0	42	12	0
39	8	180	0	165	51	0
39	9	108	0	95	35	0
39	Z	99	0	68	27	0
40	1	48	0	62	13	0
41	5	23	0	19	4	0
41	6	29	0	31	5	0
41	7	39	0	51	12	0

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Mol	Chain	Non-H	H(model)	H(added)	Clashes	Symm-Clashes
41	8	30	0	33	4	0
42	7	42	0	74	12	0
43	8	30	0	43	7	0
44	1	1	0	0	0	0
44	2	1	0	0	0	0
44	3	2	0	0	1	0
44	4	5	0	0	0	0
44	5	5	0	0	0	0
44	6	3	0	0	0	0
44	7	4	0	0	0	0
44	8	4	0	0	0	0
44	A	19	0	0	0	0
44	B	17	0	0	0	0
44	F	3	0	0	0	0
44	I	1	0	0	0	0
44	J	1	0	0	0	0
44	K	1	0	0	0	0
44	Z	1	0	0	0	0
All	All	50057	0	49449	4232	0

The all-atom clashscore is defined as the number of clashes found per 1000 atoms (including hydrogen atoms). The all-atom clashscore for this structure is 43.

All (4232) close contacts within the same asymmetric unit are listed below, sorted by their clash magnitude.

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
18:6:238:ASP:HA	18:6:243:LYS:HA	1.26	1.13
22:B:820:CLA:HBB1	22:B:820:CLA:HMB1	1.35	1.08
24:3:306:BCR:H21C	22:3:318:CLA:H91	1.25	1.06
22:B:813:CLA:HMB1	22:B:813:CLA:HBB1	1.35	1.04
24:5:304:BCR:H19C	22:5:319:CLA:HBA2	1.40	1.04
22:3:320:CLA:HMD2	41:7:319:3PH:H342	1.40	1.03
1:A:80:GLN:HG2	22:A:805:CLA:HMA1	1.41	1.03
12:Z:153:PRO:HD3	22:Z:313:CLA:HMD2	1.40	1.03
19:9:53:PRO:HD2	38:9:302:LUT:H23	1.42	1.02
38:6:304:LUT:H28	22:6:310:CLA:H61	1.39	1.02
17:5:175:SER:HB3	22:5:307:CLA:HAA2	1.37	1.00
12:Z:72:GLU:HG2	12:Z:151:ILE:HG12	1.45	0.99
22:Z:306:CLA:H71	22:Z:307:CLA:HMA1	1.43	0.99
38:7:302:LUT:H32	22:7:307:CLA:HAB	1.44	0.98
18:6:127:LYS:HG3	22:6:302:CLA:HED3	1.43	0.98

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
22:A:822:CLA:HMB2	22:A:823:CLA:H2	1.44	0.98
22:5:309:CLA:HBD	22:5:309:CLA:HBA1	1.46	0.97
22:A:817:CLA:HAB	24:K:206:BCR:HC32	1.42	0.97
22:B:810:CLA:H61	24:I:4001:BCR:HC31	1.45	0.97
15:8:52:PRO:HD2	38:8:303:LUT:H23	1.46	0.97
22:B:811:CLA:HHC	22:B:811:CLA:HBB1	1.47	0.96
16:4:198:ASP:HA	38:4:802:LUT:H24	1.47	0.95
22:B:840:CLA:H161	24:I:4001:BCR:H19C	1.49	0.95
1:A:307:VAL:HA	24:A:856:BCR:H17C	1.45	0.95
22:6:309:CLA:H71	25:6:323:LHG:H341	1.48	0.95
22:B:811:CLA:HBA2	22:B:829:CLA:HBD	1.45	0.95
22:3:310:CLA:HBD	22:3:310:CLA:HBA1	1.43	0.94
22:7:306:CLA:H91	22:7:306:CLA:H142	1.45	0.94
22:A:831:CLA:HMA2	11:L:57:THR:HG21	1.47	0.94
38:9:301:LUT:H30	22:9:303:CLA:H52	1.47	0.94
24:6:305:BCR:H16C	39:6:316:CHL:HMB3	1.49	0.93
17:5:58:LEU:HD13	38:5:303:LUT:H222	1.49	0.93
22:7:306:CLA:HBD	22:7:306:CLA:HBA1	1.48	0.93
3:C:73:SER:H	3:C:76:SER:HB3	1.32	0.93
20:2:221:ILE:HG23	20:2:227:LEU:HD21	1.50	0.93
22:A:808:CLA:HHC	22:A:808:CLA:HBB1	1.51	0.93
12:Z:182:MET:HE3	22:Z:306:CLA:HMC3	1.48	0.93
22:5:311:CLA:H71	22:5:323:CLA:HAB	1.51	0.92
22:B:838:CLA:H202	32:F:306:RRX:H34	1.49	0.92
6:F:204:VAL:HG13	22:8:309:CLA:H2	1.49	0.92
12:1:158:ASP:HA	38:1:302:LUT:H24	1.50	0.92
20:2:177:VAL:HG21	22:2:304:CLA:HMD1	1.51	0.92
1:A:575:CYS:HG	29:B:802:SF4:FE3	0.69	0.92
22:A:837:CLA:HHC	22:A:837:CLA:HBB1	1.50	0.92
22:1:310:CLA:HBB1	22:1:310:CLA:HHC	1.52	0.92
22:B:814:CLA:HAA1	7:G:81:LEU:HB3	1.52	0.92
39:4:816:CHL:HMB1	39:4:816:CHL:HBB1	1.52	0.92
1:A:212:SER:HB3	24:A:844:BCR:H363	1.52	0.91
22:B:823:CLA:HHC	22:B:823:CLA:HBB1	1.52	0.91
22:A:816:CLA:HBB1	22:A:816:CLA:HMB1	1.51	0.91
2:B:348:LEU:HD11	22:B:828:CLA:HAA2	1.50	0.91
22:F:302:CLA:HHC	22:F:302:CLA:HBB1	1.50	0.91
22:B:814:CLA:HBA1	19:9:54:LEU:HB3	1.51	0.91
22:4:809:CLA:HHC	22:4:809:CLA:HBB1	1.51	0.91
17:5:182:LEU:HB3	22:5:307:CLA:HMA1	1.52	0.90
22:6:314:CLA:HHC	22:6:314:CLA:HBB1	1.53	0.90

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
19:9:115:GLU:HA	39:9:314:CHL:HMA3	1.53	0.90
22:8:306:CLA:HBB1	22:8:306:CLA:HHC	1.52	0.90
2:B:210:TRP:HB2	25:B:851:LHG:HC5	1.54	0.90
22:B:825:CLA:H92	22:B:825:CLA:H2	1.54	0.90
22:B:836:CLA:HMB1	22:B:836:CLA:HBB1	1.54	0.90
22:9:306:CLA:HMB1	22:9:306:CLA:HBB1	1.54	0.90
22:3:313:CLA:HMB1	22:3:313:CLA:HBB1	1.54	0.90
22:B:805:CLA:HBB1	22:B:805:CLA:HHC	1.55	0.89
22:B:835:CLA:HMB1	22:B:835:CLA:HBB1	1.54	0.89
5:E:78:GLN:HB3	5:E:84:THR:HG22	1.54	0.89
22:A:824:CLA:H2	22:A:835:CLA:H171	1.54	0.89
22:A:823:CLA:HBB1	22:A:823:CLA:HHC	1.54	0.89
22:7:322:CLA:HMB1	22:7:322:CLA:HBB1	1.55	0.89
22:B:827:CLA:H151	22:B:842:CLA:H161	1.55	0.89
2:B:382:PHE:HB3	22:B:809:CLA:H112	1.54	0.89
15:8:78:THR:HG23	22:8:305:CLA:HMC1	1.55	0.89
22:Z:303:CLA:HBB1	22:Z:303:CLA:HMB1	1.54	0.89
22:A:802:CLA:H13	22:A:828:CLA:H143	1.53	0.88
22:A:830:CLA:HMB1	22:A:830:CLA:HBB1	1.52	0.88
22:6:308:CLA:HMB1	22:6:308:CLA:HBB1	1.55	0.88
38:5:302:LUT:H11	22:5:308:CLA:HMC2	1.54	0.88
24:B:844:BCR:H362	24:G:203:BCR:H312	1.56	0.88
22:1:305:CLA:HHC	22:1:305:CLA:HBB1	1.53	0.88
22:5:322:CLA:HMB1	22:5:322:CLA:HBB1	1.56	0.88
13:3:195:VAL:HG11	38:3:303:LUT:H12	1.56	0.88
22:B:833:CLA:H8	32:F:306:RRX:H27	1.56	0.88
22:A:815:CLA:HBB1	22:A:815:CLA:HMB1	1.55	0.88
2:B:560:CYS:HG	29:B:802:SF4:FE4	0.83	0.88
22:6:317:CLA:HED2	22:6:317:CLA:H2A	1.52	0.88
22:A:807:CLA:HMB3	22:A:808:CLA:HHB	1.55	0.88
22:K:202:CLA:HMB1	22:K:202:CLA:HBB1	1.55	0.88
22:B:831:CLA:HBB1	22:B:831:CLA:HMB1	1.55	0.87
12:Z:145:THR:HA	12:Z:150:ARG:HH21	1.39	0.87
24:3:304:BCR:H281	22:5:301:CLA:HAB	1.57	0.87
22:B:821:CLA:H151	22:B:826:CLA:H172	1.57	0.87
24:5:305:BCR:H321	24:5:305:BCR:HC8	1.56	0.87
22:A:826:CLA:HBB1	22:A:826:CLA:HMB1	1.56	0.87
22:B:808:CLA:HBB1	22:B:808:CLA:HMB1	1.54	0.87
14:7:205:LEU:HD13	22:7:308:CLA:HBB2	1.55	0.87
22:7:308:CLA:HMB1	22:7:308:CLA:HBB1	1.57	0.87
16:4:95:TRP:HZ2	22:4:815:CLA:HAA2	1.38	0.87

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
22:A:814:CLA:HBB1	22:A:814:CLA:HMB1	1.56	0.87
22:Z:316:CLA:HHC	22:Z:316:CLA:HBB1	1.55	0.87
22:7:312:CLA:HHC	22:7:312:CLA:HBB1	1.57	0.87
6:F:79:ARG:HG2	6:F:115:PHE:HZ	1.40	0.86
38:3:302:LUT:H361	22:3:308:CLA:H143	1.57	0.86
22:4:815:CLA:HBB1	22:4:815:CLA:HMB1	1.54	0.86
22:B:812:CLA:HMB1	22:B:812:CLA:HBB1	1.57	0.86
12:1:35:TRP:HH2	40:1:318:SQD:H2	1.40	0.86
13:3:55:TRP:HZ2	22:3:318:CLA:HAA2	1.40	0.86
18:6:247:PRO:HB2	22:6:301:CLA:H11	1.56	0.86
19:9:81:VAL:HG11	38:9:301:LUT:H12	1.57	0.86
13:3:203:VAL:HG22	31:3:301:DGD:HB62	1.58	0.86
22:A:820:CLA:HHC	22:A:820:CLA:HBB1	1.56	0.86
22:A:832:CLA:HMB1	22:A:832:CLA:HBB1	1.57	0.86
24:5:304:BCR:H16C	39:5:317:CHL:HMB3	1.56	0.86
22:A:836:CLA:H61	22:K:203:CLA:HBA2	1.58	0.86
3:C:14:CYS:SG	29:C:102:SF4:FE3	1.68	0.85
12:Z:149:LYS:HB3	22:Z:313:CLA:HBC1	1.57	0.85
20:2:197:ASN:HA	22:2:301:CLA:HBA2	1.58	0.85
10:K:59:ARG:HA	10:K:69:ASP:HA	1.58	0.85
15:8:55:LEU:HD12	38:8:303:LUT:H221	1.57	0.85
22:4:807:CLA:HBD	22:4:807:CLA:HBA1	1.59	0.85
22:6:309:CLA:HBD	22:6:309:CLA:HBA1	1.56	0.85
22:3:309:CLA:HBB1	22:3:309:CLA:HMB1	1.57	0.85
22:A:835:CLA:H102	24:A:847:BCR:H20C	1.59	0.85
11:L:102:LYS:HG2	22:L:203:CLA:HMA1	1.58	0.85
22:B:840:CLA:H2	22:B:841:CLA:H121	1.59	0.85
22:A:803:CLA:HHC	22:A:803:CLA:HBB1	1.58	0.84
22:B:808:CLA:HBA1	22:B:815:CLA:HBA1	1.57	0.84
15:8:169:ASP:HB2	22:8:305:CLA:HBD	1.57	0.84
24:A:856:BCR:HC41	31:3:301:DGD:HAT1	1.58	0.84
2:B:345:ILE:HD11	22:B:820:CLA:H122	1.59	0.84
24:B:853:BCR:HC21	22:L:202:CLA:H62	1.60	0.84
22:7:309:CLA:HMB1	22:7:309:CLA:HBB1	1.57	0.84
22:8:307:CLA:HBD	22:8:307:CLA:HBA1	1.59	0.84
22:A:822:CLA:HMB1	22:A:822:CLA:HBB1	1.59	0.84
24:B:853:BCR:H11C	22:L:202:CLA:HAB	1.58	0.84
11:L:131:ALA:HB1	24:L:204:BCR:H19C	1.57	0.84
22:L:201:CLA:HHC	22:L:201:CLA:HBB1	1.57	0.84
17:5:227:TRP:HB3	22:5:309:CLA:H2	1.58	0.84
22:L:203:CLA:HBB1	22:L:203:CLA:HMB1	1.59	0.84

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
17:5:109:LEU:HD21	22:5:319:CLA:HAC1	1.57	0.84
22:G:201:CLA:HBB1	22:G:201:CLA:HHC	1.57	0.83
38:9:302:LUT:H28	22:9:306:CLA:H52	1.58	0.83
39:9:312:CHL:HBC2	39:9:312:CHL:HMC	1.60	0.83
1:A:496:GLN:HG3	1:A:497:LEU:HG	1.59	0.83
38:8:302:LUT:H11	22:8:306:CLA:HMC2	1.60	0.83
6:F:204:VAL:HG22	22:8:309:CLA:H72	1.60	0.83
12:Z:73:SER:HB2	22:Z:314:CLA:HED3	1.61	0.83
22:A:825:CLA:H193	22:A:825:CLA:H13	1.61	0.83
39:4:813:CHL:HMB1	39:4:813:CHL:HBB1	1.60	0.83
18:6:110:ILE:HD13	24:6:306:BCR:HC8	1.59	0.83
22:6:311:CLA:HBB1	22:6:311:CLA:HMB1	1.60	0.83
22:1:304:CLA:HBB1	22:1:304:CLA:HMB1	1.59	0.83
22:7:306:CLA:HBB1	22:7:306:CLA:HMB1	1.61	0.83
22:5:308:CLA:HMB3	22:5:308:CLA:H151	1.59	0.83
18:6:54:PRO:HD2	38:6:304:LUT:H23	1.59	0.83
2:B:54:GLN:HG2	22:B:808:CLA:HMA1	1.60	0.82
22:B:833:CLA:HMB1	22:B:833:CLA:HBB1	1.61	0.82
18:6:249:LEU:HG	24:6:306:BCR:H291	1.61	0.82
39:6:318:CHL:HBB1	39:6:318:CHL:HMB1	1.59	0.82
17:5:170:ASP:HA	38:5:302:LUT:H24	1.59	0.82
22:B:823:CLA:HMD2	24:B:844:BCR:HC7	1.59	0.82
38:5:303:LUT:H28	22:5:310:CLA:H52	1.61	0.82
18:6:170:LEU:HB3	22:6:307:CLA:HMA1	1.60	0.82
13:3:82:LYS:HZ2	13:3:210:TYR:HB3	1.43	0.82
22:6:301:CLA:H122	39:6:315:CHL:H43	1.62	0.82
22:9:306:CLA:H71	22:9:307:CLA:HMA1	1.61	0.82
1:A:29:TRP:HE1	22:A:811:CLA:CHB	1.93	0.82
22:B:814:CLA:HHC	22:B:814:CLA:HBB1	1.59	0.82
22:B:827:CLA:HMB1	22:B:827:CLA:HBB1	1.62	0.82
22:8:305:CLA:HMB1	22:8:305:CLA:HBB1	1.61	0.82
22:A:829:CLA:H2	24:A:845:BCR:HC7	1.59	0.81
39:Z:311:CHL:HMA1	25:Z:317:LHG:H192	1.60	0.81
18:6:222:VAL:HG12	18:6:225:GLN:H	1.45	0.81
22:6:317:CLA:HMB1	22:6:317:CLA:HBB1	1.60	0.81
1:A:338:HIS:HE1	25:A:849:LHG:HC12	1.45	0.81
22:A:812:CLA:HBB1	22:A:812:CLA:HHC	1.62	0.81
2:B:657:VAL:HG22	22:B:841:CLA:HMB3	1.62	0.81
22:B:841:CLA:HAC2	24:B:849:BCR:H24C	1.62	0.81
12:Z:215:PHE:HB3	22:Z:305:CLA:H11	1.60	0.81
22:3:311:CLA:HBB1	22:3:311:CLA:HMB1	1.60	0.81

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
22:1:314:CLA:H172	22:1:314:CLA:H91	1.62	0.81
22:B:825:CLA:HMB1	22:B:825:CLA:HBB1	1.62	0.81
39:8:312:CHL:HBB1	39:8:312:CHL:HMB1	1.63	0.81
3:C:14:CYS:HG	29:C:102:SF4:FE3	0.96	0.81
22:A:821:CLA:H101	24:A:847:BCR:H10C	1.63	0.81
22:F:304:CLA:HMD3	33:F:307:LMT:H32	1.61	0.81
24:A:844:BCR:H362	24:A:845:BCR:H21C	1.62	0.80
15:8:163:TYR:HB3	22:8:305:CLA:HED3	1.63	0.80
19:9:126:ARG:HH22	22:9:313:CLA:HMA3	1.44	0.80
22:4:807:CLA:HBB1	22:4:807:CLA:HMB1	1.62	0.80
39:Z:312:CHL:HMB1	39:Z:312:CHL:HBB1	1.62	0.80
22:3:316:CLA:HHC	22:3:316:CLA:HBB1	1.61	0.80
22:A:836:CLA:HMB1	24:A:847:BCR:H281	1.64	0.80
39:8:315:CHL:HBB1	39:8:315:CHL:HMB1	1.63	0.80
11:L:119:LEU:O	22:L:201:CLA:H112	1.80	0.80
39:6:315:CHL:HBB1	39:6:315:CHL:HMB1	1.62	0.80
22:B:819:CLA:H92	22:B:828:CLA:HAB	1.63	0.80
17:5:57:PRO:HD2	38:5:303:LUT:H23	1.63	0.80
22:A:810:CLA:HMB1	22:A:810:CLA:HBB1	1.63	0.80
2:B:560:CYS:SG	29:B:802:SF4:FE4	1.74	0.80
22:B:830:CLA:HHC	22:B:830:CLA:HBB1	1.62	0.79
22:5:307:CLA:H72	22:5:308:CLA:HMA1	1.64	0.79
15:8:127:GLY:HA3	39:8:301:CHL:H12	1.63	0.79
16:4:83:PRO:HD2	38:4:803:LUT:H23	1.63	0.79
39:4:814:CHL:HMB1	39:4:814:CHL:HBB1	1.64	0.79
19:9:56:LEU:HD11	38:9:302:LUT:H221	1.64	0.79
14:7:104:GLU:HA	39:7:313:CHL:HED3	1.64	0.79
18:6:142:PRO:HB2	22:6:302:CLA:HMD1	1.64	0.79
22:A:840:CLA:HBC2	24:F:303:BCR:H21C	1.63	0.79
1:A:338:HIS:HA	1:A:426:LEU:HD11	1.62	0.79
22:A:842:CLA:HMB1	24:A:846:BCR:H12C	1.63	0.79
22:A:821:CLA:HMB1	22:A:821:CLA:HBB1	1.65	0.79
22:A:834:CLA:HMA2	11:L:103:LEU:HB3	1.65	0.79
22:8:307:CLA:HBB1	22:8:307:CLA:HMB1	1.65	0.79
22:6:309:CLA:HMB1	22:6:309:CLA:HBB1	1.64	0.79
12:1:36:LEU:HD11	40:1:318:SQD:H172	1.65	0.79
39:1:312:CHL:HMB1	39:1:312:CHL:HBB1	1.63	0.79
22:B:840:CLA:H172	22:B:841:CLA:H91	1.65	0.79
12:1:86:ALA:HB1	22:1:304:CLA:H172	1.64	0.78
24:3:305:BCR:H21C	22:3:313:CLA:HAC1	1.63	0.78
14:7:115:HIS:HB2	15:8:225:LEU:HD11	1.66	0.78

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
22:B:840:CLA:HBB1	22:B:840:CLA:HMB1	1.64	0.78
22:Z:308:CLA:H61	22:Z:315:CLA:HAA2	1.65	0.78
15:8:128:PHE:CE1	24:8:304:BCR:H10C	2.18	0.78
22:5:318:CLA:HBB1	22:5:318:CLA:HMB1	1.65	0.78
24:A:844:BCR:H23C	24:3:306:BCR:H333	1.63	0.78
24:6:306:BCR:H382	22:6:319:CLA:H112	1.65	0.78
22:G:202:CLA:HBB1	22:G:202:CLA:HMB1	1.65	0.78
22:A:814:CLA:H121	22:A:816:CLA:CAB	2.13	0.78
19:9:142:PHE:HB3	19:9:143:PRO:HD2	1.65	0.78
38:4:802:LUT:H30	22:4:805:CLA:H52	1.65	0.78
39:5:317:CHL:HMB1	39:5:317:CHL:HBB1	1.66	0.78
22:A:832:CLA:H18	22:L:202:CLA:H18	1.65	0.78
22:A:841:CLA:HBA2	25:A:850:LHG:H161	1.64	0.78
14:7:189:TYR:HB3	22:7:304:CLA:HMA1	1.66	0.78
12:1:140:ARG:HG3	22:1:313:CLA:HMC3	1.66	0.77
25:1:317:LHG:H101	24:8:304:BCR:H313	1.65	0.77
22:A:823:CLA:H111	24:A:856:BCR:H401	1.66	0.77
22:B:828:CLA:HBB1	22:B:828:CLA:HHC	1.66	0.77
7:G:94:PRO:HG2	22:G:202:CLA:HBC2	1.65	0.77
12:Z:140:ARG:HG3	22:Z:313:CLA:HMC3	1.66	0.77
22:A:814:CLA:H121	22:A:816:CLA:HAB	1.65	0.77
2:B:543:ARG:HB2	2:B:552:LYS:HB3	1.65	0.77
22:1:315:CLA:HBB1	22:1:315:CLA:HMB1	1.65	0.77
24:3:307:BCR:H311	22:3:315:CLA:HHB	1.66	0.77
38:7:302:LUT:H28	22:7:307:CLA:H61	1.65	0.77
6:F:142:LYS:HG3	33:F:307:LMT:H2'	1.66	0.77
14:7:133:TRP:CD1	24:7:303:BCR:H12C	2.18	0.77
22:A:804:CLA:HBB1	22:A:804:CLA:HMB1	1.63	0.77
22:B:813:CLA:H11	22:9:307:CLA:H61	1.67	0.77
12:Z:206:HIS:HB2	22:Z:305:CLA:HAA2	1.65	0.77
22:7:309:CLA:H52	22:7:316:CLA:HBD	1.65	0.77
12:Z:173:LYS:HD3	22:Z:304:CLA:HAA2	1.67	0.77
2:B:284:ILE:HA	22:B:818:CLA:H172	1.67	0.77
16:4:133:PRO:HG2	16:4:136:GLU:HB2	1.66	0.77
2:B:227:TRP:HH2	22:B:817:CLA:H71	1.50	0.77
17:5:218:HIS:CG	22:5:309:CLA:HAA1	2.19	0.77
24:6:306:BCR:H16C	39:6:315:CHL:CHD	2.14	0.77
22:6:307:CLA:HBB1	22:6:308:CLA:HAA1	1.65	0.77
22:B:819:CLA:H91	22:B:828:CLA:HMB1	1.67	0.77
38:Z:301:LUT:H28	22:Z:303:CLA:H2	1.65	0.77
15:8:164:PRO:HD3	22:8:313:CLA:HMD2	1.66	0.77

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
24:B:844:BCR:H272	24:B:846:BCR:HC31	1.67	0.76
22:A:840:CLA:H111	22:A:840:CLA:HAB	1.66	0.76
7:G:59:ARG:HG2	7:G:63:ARG:HE	1.48	0.76
25:4:820:LHG:H191	22:6:321:CLA:HBA2	1.65	0.76
38:6:303:LUT:H403	22:6:307:CLA:H122	1.66	0.76
22:B:810:CLA:HMB3	22:B:811:CLA:HMA1	1.68	0.76
22:B:814:CLA:C4D	22:B:814:CLA:H2	2.15	0.76
22:5:322:CLA:H61	22:6:310:CLA:H193	1.66	0.76
39:9:312:CHL:HHC	39:9:312:CHL:HBB1	1.67	0.76
22:A:854:CLA:H152	22:L:201:CLA:HBC1	1.65	0.76
22:B:836:CLA:HBC2	31:1:319:DGD:HA32	1.68	0.76
23:B:843:PQN:H221	24:B:849:BCR:H17C	1.66	0.76
12:1:35:TRP:CH2	40:1:318:SQD:H2	2.20	0.76
22:4:815:CLA:HMB2	33:4:822:LMT:H51	1.68	0.76
39:6:316:CHL:CHB	39:6:318:CHL:H71	2.16	0.76
24:A:848:BCR:H372	2:B:439:VAL:HG21	1.65	0.76
22:B:841:CLA:HBD	23:B:843:PQN:H302	1.68	0.76
19:9:79:MET:HG2	22:9:308:CLA:HAB	1.67	0.76
8:I:83:THR:HG21	24:I:4001:BCR:HC42	1.68	0.76
12:1:152:TYR:HB3	22:1:304:CLA:HED2	1.66	0.76
18:6:75:ALA:HB1	18:6:178:GLY:HA3	1.67	0.76
20:2:177:VAL:HG21	22:2:304:CLA:CMD	2.16	0.76
22:B:834:CLA:HBA2	24:J:104:BCR:H272	1.68	0.76
12:Z:203:LEU:HB2	38:Z:301:LUT:H22	1.68	0.76
31:3:301:DGD:HAE2	24:3:306:BCR:H10C	1.66	0.76
6:F:204:VAL:HA	22:8:309:CLA:H51	1.68	0.76
12:Z:103:PRO:HB3	22:Z:315:CLA:HMC1	1.67	0.76
13:3:55:TRP:CZ2	22:3:318:CLA:HAA2	2.19	0.76
18:6:46:LEU:H	18:6:46:LEU:HD12	1.50	0.76
22:A:804:CLA:HBA2	22:A:811:CLA:H92	1.66	0.75
11:L:132:LEU:HB3	11:L:171:THR:HG22	1.68	0.75
15:8:80:VAL:HG11	38:8:302:LUT:H12	1.68	0.75
11:L:132:LEU:HD21	24:L:204:BCR:H383	1.68	0.75
16:4:136:GLU:HA	39:4:813:CHL:HED2	1.67	0.75
17:5:241:ASP:HA	17:5:246:THR:HA	1.69	0.75
19:9:32:LEU:HD22	19:9:33:PRO:HD2	1.69	0.75
12:Z:190:ALA:HB2	22:Z:316:CLA:H61	1.68	0.75
38:Z:301:LUT:H30	22:Z:303:CLA:H52	1.67	0.75
22:3:313:CLA:H43	22:5:320:CLA:HED2	1.68	0.75
22:B:827:CLA:H143	22:B:842:CLA:H18	1.67	0.75
39:5:316:CHL:HMC	22:5:319:CLA:HAB	1.66	0.75

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:B:86:ARG:NH2	2:B:107:ARG:HB2	2.02	0.75
22:B:820:CLA:HMB1	22:B:820:CLA:CBB	2.17	0.75
24:B:847:BCR:H323	33:1:301:LMT:H102	1.67	0.75
17:5:159:ASN:HA	17:5:165:PRO:HA	1.69	0.75
2:B:351:GLN:HG3	22:B:827:CLA:HED2	1.68	0.75
12:1:94:LEU:HD12	22:1:309:CLA:HMD3	1.69	0.75
22:K:205:CLA:HMD2	24:K:206:BCR:H21C	1.69	0.74
11:L:120:SER:HA	22:L:201:CLA:H71	1.69	0.74
22:1:306:CLA:H152	25:1:317:LHG:HC92	1.69	0.74
13:3:37:ASP:HA	38:3:303:LUT:H24	1.70	0.74
22:B:825:CLA:HBB2	22:B:842:CLA:H61	1.70	0.74
39:1:312:CHL:HAA2	31:1:319:DGD:HG31	1.67	0.74
22:A:820:CLA:H102	22:A:820:CLA:H142	1.69	0.74
22:7:322:CLA:HMC3	15:8:221:LEU:HD21	1.68	0.74
12:Z:51:GLY:HA3	12:Z:175:ILE:HD13	1.69	0.74
39:Z:311:CHL:HBC3	39:Z:311:CHL:HHD	1.68	0.74
22:3:320:CLA:HMA2	22:3:322:CLA:H112	1.68	0.74
22:7:307:CLA:HBB1	22:7:307:CLA:HMB1	1.68	0.74
1:A:575:CYS:SG	29:B:802:SF4:FE3	1.79	0.74
2:B:86:ARG:HH21	2:B:107:ARG:HB2	1.52	0.74
13:3:165:ASN:HA	38:3:302:LUT:H24	1.69	0.74
14:7:205:LEU:HD11	22:7:307:CLA:H202	1.69	0.74
22:A:854:CLA:HBB2	2:B:659:ALA:HB3	1.70	0.74
22:3:310:CLA:HBA1	22:3:310:CLA:CBD	2.17	0.74
22:A:802:CLA:HAB	2:B:583:TRP:CH2	2.23	0.74
22:B:824:CLA:HMB1	22:B:824:CLA:HBB1	1.68	0.74
22:B:838:CLA:H171	32:F:306:RRX:H38	1.68	0.74
12:Z:68:LYS:HD2	12:Z:147:ALA:HB1	1.69	0.74
22:9:308:CLA:HMB1	22:9:308:CLA:HBB1	1.70	0.74
1:A:363:LEU:HD11	22:A:819:CLA:H111	1.68	0.74
34:J:101:LMG:HC91	34:J:101:LMG:H121	1.70	0.74
22:3:308:CLA:HMB1	22:3:308:CLA:HBB1	1.69	0.74
22:6:322:CLA:HBB1	22:6:322:CLA:HMB1	1.70	0.74
1:A:453:PHE:HE1	22:B:806:CLA:HMA1	1.53	0.73
1:A:676:HIS:HB2	22:A:802:CLA:HBA1	1.70	0.73
22:A:822:CLA:HBA2	22:A:823:CLA:H72	1.70	0.73
11:L:140:PHE:HA	11:L:144:PRO:HD2	1.70	0.73
12:Z:162:PHE:HB2	22:Z:303:CLA:CGA	2.18	0.73
22:Z:304:CLA:HMD1	22:Z:309:CLA:HBA2	1.71	0.73
22:A:840:CLA:H121	22:F:305:CLA:HAC1	1.70	0.73
22:Z:304:CLA:NA	22:Z:309:CLA:H71	2.03	0.73

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
22:5:320:CLA:HAC2	41:5:325:3PH:H232	1.70	0.73
22:4:807:CLA:H102	25:4:820:LHG:H162	1.68	0.73
22:6:314:CLA:H92	25:6:323:LHG:H152	1.71	0.73
22:5:322:CLA:H143	22:6:310:CLA:H101	1.71	0.73
19:9:119:MET:HE3	39:9:314:CHL:HHB	1.69	0.73
22:4:818:CLA:H2	22:4:818:CLA:HBA1	1.71	0.73
22:5:307:CLA:HBB1	22:5:307:CLA:HMB1	1.69	0.73
22:B:825:CLA:HBA2	24:B:847:BCR:H14C	1.71	0.73
2:B:26:LEU:HD21	24:L:204:BCR:H332	1.71	0.73
22:B:813:CLA:HMB1	22:B:813:CLA:CBB	2.15	0.73
10:K:59:ARG:CA	10:K:69:ASP:HA	2.19	0.73
31:3:301:DGD:HB21	31:3:301:DGD:HG31	1.71	0.73
14:7:174:ASP:HA	38:7:301:LUT:H24	1.70	0.73
24:5:304:BCR:H353	39:5:317:CHL:HBB1	1.69	0.73
39:6:320:CHL:HHC	39:6:320:CHL:HBB1	1.70	0.73
19:9:100:GLY:HA2	39:9:314:CHL:HAC2	1.70	0.73
22:A:840:CLA:H91	22:F:305:CLA:HAC2	1.71	0.72
24:3:305:BCR:H321	24:3:305:BCR:HC8	1.69	0.72
22:7:322:CLA:H61	22:7:323:CLA:H72	1.70	0.72
24:4:804:BCR:H272	22:4:810:CLA:H2	1.69	0.72
22:A:823:CLA:H191	10:K:98:ILE:HD12	1.71	0.72
2:B:569:CYS:SG	29:B:802:SF4:FE1	1.81	0.72
22:A:815:CLA:H161	13:3:195:VAL:HG12	1.70	0.72
22:A:818:CLA:H62	22:A:818:CLA:C4C	2.19	0.72
24:3:307:BCR:H311	22:3:315:CLA:HMA1	1.70	0.72
14:7:167:TYR:HB3	22:7:304:CLA:HED3	1.70	0.72
16:4:121:ARG:HH22	16:4:133:PRO:HA	1.55	0.72
22:A:822:CLA:HED1	10:K:70:ARG:HE	1.54	0.72
22:5:308:CLA:HBB1	22:5:308:CLA:HMB1	1.70	0.72
2:B:296:PHE:O	7:G:88:ILE:HG13	1.89	0.72
22:B:834:CLA:H51	24:F:303:BCR:HC22	1.72	0.72
22:B:816:CLA:HMA2	24:B:846:BCR:H282	1.71	0.72
22:B:827:CLA:H192	22:B:842:CLA:H91	1.70	0.72
1:A:615:GLN:HG2	1:A:630:ILE:HD12	1.70	0.72
22:A:806:CLA:HHC	22:A:806:CLA:HBB1	1.70	0.72
22:A:822:CLA:HMD2	24:A:856:BCR:H23C	1.72	0.72
22:B:840:CLA:H8	22:B:841:CLA:H142	1.70	0.72
13:3:70:GLY:HA2	38:3:303:LUT:H181	1.71	0.72
22:3:308:CLA:HMD2	39:3:317:CHL:HBA2	1.72	0.72
15:8:187:LYS:HG3	22:8:311:CLA:HED2	1.69	0.72
22:B:823:CLA:H71	22:B:824:CLA:H112	1.71	0.71

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
22:4:810:CLA:HMB1	22:4:810:CLA:HBB1	1.71	0.71
14:7:133:TRP:CZ2	24:7:303:BCR:HC8	2.24	0.71
22:B:828:CLA:H112	24:B:847:BCR:H373	1.71	0.71
6:F:202:TRP:CG	6:F:203:PRO:HD3	2.25	0.71
22:7:323:CLA:H18	22:7:324:CLA:HMC2	1.72	0.71
22:A:810:CLA:HMD2	22:3:312:CLA:H102	1.71	0.71
22:A:827:CLA:H192	22:A:835:CLA:H91	1.70	0.71
24:F:303:BCR:H321	24:F:303:BCR:HC8	1.72	0.71
22:7:306:CLA:NC	22:7:306:CLA:H52	2.05	0.71
39:4:819:CHL:HBB1	39:4:819:CHL:HHC	1.73	0.71
22:A:822:CLA:H2	22:A:823:CLA:H92	1.73	0.71
18:6:250:PHE:H	22:6:301:CLA:H43	1.55	0.71
7:G:38:VAL:HG11	22:G:201:CLA:C4D	2.21	0.71
13:3:120:ALA:HB1	24:3:304:BCR:H363	1.72	0.71
22:B:827:CLA:CAB	24:B:848:BCR:H19C	2.21	0.71
10:K:31:GLY:H	10:K:35:ASN:HD22	1.37	0.71
22:A:825:CLA:HBB1	22:A:825:CLA:HMB1	1.72	0.71
6:F:202:TRP:CD1	6:F:203:PRO:HD3	2.25	0.71
39:4:813:CHL:HAB	39:4:816:CHL:HBB2	1.72	0.71
18:6:126:LEU:HD12	22:6:302:CLA:HBA1	1.72	0.71
12:1:45:LEU:HD21	22:1:307:CLA:HAA2	1.71	0.71
15:8:85:ILE:HB	15:8:86:PRO:HD3	1.72	0.71
22:F:302:CLA:H193	32:F:306:RRX:H21	1.73	0.71
22:J:103:CLA:HBB1	22:J:103:CLA:HMB1	1.73	0.71
22:1:308:CLA:HHC	22:1:308:CLA:HBB1	1.73	0.71
22:7:316:CLA:HED2	22:7:324:CLA:HMD2	1.72	0.71
6:F:201:THR:HA	22:8:309:CLA:H152	1.72	0.70
38:1:302:LUT:H30	22:1:304:CLA:H52	1.73	0.70
12:1:92:GLU:HG2	12:1:98:ASN:HA	1.72	0.70
12:Z:78:CYS:HB3	12:Z:178:GLY:HA3	1.73	0.70
19:9:129:GLY:HA3	19:9:142:PHE:CZ	2.26	0.70
1:A:316:TRP:HD1	10:K:76:LEU:HD21	1.55	0.70
22:B:824:CLA:HAB	24:B:847:BCR:C10	2.20	0.70
22:7:306:CLA:HBA1	22:7:306:CLA:CBD	2.21	0.70
25:4:820:LHG:HC82	24:6:305:BCR:HC42	1.73	0.70
39:9:312:CHL:H2	39:9:312:CHL:HMA3	1.73	0.70
22:A:820:CLA:HBB2	24:A:844:BCR:H402	1.73	0.70
2:B:549:PRO:HB3	6:F:226:PRO:HG2	1.72	0.70
2:B:719:VAL:HG21	31:B:852:DGD:HBV1	1.73	0.70
13:3:159:PRO:HG3	39:3:317:CHL:HMD2	1.73	0.70
16:4:200:LEU:HD12	38:4:802:LUT:H222	1.74	0.70

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
22:Z:308:CLA:H151	22:Z:313:CLA:HAA2	1.73	0.70
2:B:620:TRP:O	2:B:624:TYR:HB3	1.91	0.70
22:Z:304:CLA:HHC	22:Z:304:CLA:HBB1	1.72	0.70
22:Z:309:CLA:HMC3	24:4:804:BCR:HC21	1.74	0.70
22:Z:316:CLA:HMB2	22:8:310:CLA:H122	1.73	0.70
38:6:303:LUT:H371	22:6:307:CLA:H92	1.73	0.70
2:B:618:MET:HG3	2:B:622:ARG:HH12	1.55	0.70
22:B:815:CLA:H51	22:B:820:CLA:HBC3	1.70	0.70
8:I:88:PRO:HG3	24:I:4001:BCR:H12C	1.72	0.70
22:4:815:CLA:HED2	22:4:815:CLA:H2A	1.72	0.70
17:5:106:PRO:HB2	17:5:115:ASN:HA	1.74	0.70
39:5:316:CHL:H122	39:5:316:CHL:H91	1.74	0.70
1:A:45:THR:HG23	5:E:78:GLN:HE21	1.55	0.70
13:3:96:VAL:HG11	38:3:303:LUT:H172	1.71	0.70
13:3:180:LEU:HB3	22:3:308:CLA:HMA1	1.74	0.70
22:3:308:CLA:H152	39:3:317:CHL:H72	1.73	0.70
22:3:312:CLA:HHC	22:3:312:CLA:HBB1	1.74	0.70
22:5:311:CLA:C7	22:5:323:CLA:HAB	2.22	0.70
22:A:813:CLA:HBB1	22:A:813:CLA:HMB1	1.72	0.70
22:A:819:CLA:HMB1	22:A:819:CLA:HBB1	1.73	0.70
24:B:853:BCR:H271	22:L:201:CLA:HED1	1.73	0.70
18:6:247:PRO:HG3	22:6:301:CLA:H111	1.73	0.70
22:6:312:CLA:HBC3	22:6:312:CLA:H122	1.74	0.70
13:3:203:VAL:HG13	31:3:301:DGD:HG32	1.74	0.69
1:A:205:LEU:HA	24:A:845:BCR:H363	1.74	0.69
22:F:304:CLA:HBB1	32:F:306:RRX:H51	1.74	0.69
24:3:304:BCR:H281	22:5:301:CLA:CAB	2.22	0.69
39:3:317:CHL:HHC	39:3:317:CHL:HBB1	1.74	0.69
22:F:305:CLA:HMB1	22:F:305:CLA:HBB1	1.72	0.69
38:Z:301:LUT:H403	22:Z:303:CLA:H141	1.73	0.69
39:6:316:CHL:HHC	39:6:316:CHL:HBB1	1.73	0.69
13:3:103:TYR:HB3	13:3:106:TYR:CE2	2.26	0.69
14:7:53:PHE:HB3	22:7:307:CLA:CAD	2.23	0.69
14:7:178:PHE:HE2	38:7:301:LUT:H221	1.56	0.69
15:8:104:GLY:HA2	39:8:315:CHL:HAC2	1.75	0.69
22:A:809:CLA:HHC	22:A:809:CLA:HBB1	1.74	0.69
22:B:837:CLA:HMB1	22:B:837:CLA:HBB1	1.74	0.69
12:Z:84:GLY:HA2	38:Z:302:LUT:H181	1.75	0.69
22:A:813:CLA:HBA1	22:A:825:CLA:H41	1.74	0.69
22:B:809:CLA:H41	31:B:852:DGD:HB61	1.75	0.69
22:B:816:CLA:HBB1	22:B:816:CLA:HMB1	1.75	0.69

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
18:6:123:PHE:CZ	39:6:320:CHL:HBB2	2.28	0.69
39:2:303:CHL:HHC	39:2:303:CHL:HBB1	1.73	0.69
22:A:822:CLA:HAA2	10:K:61:THR:HG21	1.75	0.69
13:3:203:VAL:HA	31:3:301:DGD:HB42	1.73	0.69
39:5:321:CHL:HBB1	39:5:321:CHL:HHC	1.74	0.69
22:1:305:CLA:HBC3	22:1:310:CLA:H143	1.74	0.69
17:5:185:LYS:HD3	22:5:308:CLA:HAA2	1.75	0.69
22:A:823:CLA:HAA2	10:K:63:ALA:O	1.93	0.68
22:A:838:CLA:HBB1	22:A:838:CLA:HMB1	1.74	0.68
22:B:833:CLA:HBC2	22:B:839:CLA:HMC2	1.75	0.68
6:F:153:THR:HA	22:F:302:CLA:HAC2	1.75	0.68
12:Z:170:LEU:HB3	22:Z:303:CLA:HMA1	1.75	0.68
38:7:302:LUT:C32	22:7:308:CLA:HMB2	2.22	0.68
1:A:157:LEU:HD11	27:A:852:DGA:HA81	1.75	0.68
22:A:812:CLA:HBA1	22:A:812:CLA:CHA	2.23	0.68
22:A:841:CLA:H12	22:A:841:CLA:NA	2.08	0.68
22:B:824:CLA:HAB	24:B:847:BCR:C9	2.24	0.68
24:A:848:BCR:H292	22:F:302:CLA:HMA1	1.75	0.68
22:A:855:CLA:H122	22:B:841:CLA:H141	1.76	0.68
13:3:73:GLY:HA3	38:3:303:LUT:H182	1.76	0.68
22:A:809:CLA:HMA1	9:J:27:ILE:HD13	1.75	0.68
22:A:825:CLA:H143	22:A:825:CLA:HMD2	1.75	0.68
2:B:283:VAL:HG13	24:B:844:BCR:H363	1.74	0.68
12:Z:134:MET:HG3	22:Z:314:CLA:HMC3	1.74	0.68
12:Z:152:TYR:HB3	22:Z:303:CLA:HED3	1.74	0.68
39:9:314:CHL:HBB1	39:9:314:CHL:HHC	1.74	0.68
22:A:808:CLA:HBB2	22:A:828:CLA:H171	1.74	0.68
22:F:301:CLA:H152	9:J:18:TRP:HE3	1.58	0.68
24:4:804:BCR:H373	39:4:816:CHL:HHB	1.76	0.68
22:5:312:CLA:HMB1	22:5:312:CLA:HBB1	1.75	0.68
22:A:807:CLA:H8	22:A:809:CLA:H61	1.73	0.68
22:A:823:CLA:H152	24:A:856:BCR:H401	1.73	0.68
22:B:815:CLA:HMB1	22:B:815:CLA:HBB1	1.75	0.68
11:L:141:GLN:HE21	11:L:141:GLN:H	1.41	0.68
22:B:817:CLA:H51	24:B:844:BCR:H282	1.75	0.68
22:A:805:CLA:HBB1	22:A:805:CLA:HMB1	1.74	0.68
2:B:687:PRO:HG2	22:L:202:CLA:H52	1.76	0.68
16:4:210:LEU:HB3	22:4:805:CLA:HMA1	1.76	0.68
18:6:99:ASP:HA	39:6:315:CHL:O1D	1.94	0.68
22:6:310:CLA:H71	22:6:311:CLA:HMA1	1.75	0.68
21:A:801:CL0:H13	22:A:802:CLA:CAD	2.23	0.67

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:B:499:LEU:HA	2:B:502:ILE:HG22	1.76	0.67
22:B:801:CLA:HMB1	22:B:801:CLA:HBB1	1.75	0.67
6:F:79:ARG:HG2	6:F:115:PHE:CZ	2.26	0.67
22:5:322:CLA:H152	22:6:319:CLA:H71	1.77	0.67
24:A:856:BCR:H24C	10:K:95:LEU:HD21	1.76	0.67
2:B:197:HIS:O	2:B:201:PRO:HG2	1.95	0.67
22:G:201:CLA:CAB	24:G:203:BCR:H352	2.24	0.67
10:K:39:VAL:HG22	22:K:202:CLA:HBA1	1.75	0.67
39:6:315:CHL:H51	39:6:315:CHL:H92	1.77	0.67
22:Z:313:CLA:H101	22:Z:313:CLA:HBA1	1.76	0.67
22:A:830:CLA:HMD2	25:A:850:LHG:H291	1.77	0.67
22:A:840:CLA:H111	22:A:840:CLA:CAB	2.25	0.67
22:B:833:CLA:C8	32:F:306:RRX:H27	2.24	0.67
13:3:89:ILE:HD13	13:3:94:THR:HG22	1.76	0.67
18:6:149:LEU:HD11	18:6:158:GLY:HA3	1.75	0.67
20:2:198:ILE:HG13	22:2:301:CLA:HMD2	1.77	0.67
16:4:67:PHE:HD1	16:4:80:GLY:HA3	1.60	0.67
16:4:121:ARG:HG3	16:4:131:GLN:HA	1.75	0.67
23:B:843:PQN:H201	24:B:849:BCR:H21C	1.76	0.67
9:J:1:MET:H2	34:J:101:LMG:HC2	1.60	0.67
14:7:26:TYR:HE2	14:7:191:GLN:HB2	1.59	0.67
22:9:313:CLA:HHC	22:9:313:CLA:HBB1	1.76	0.67
1:A:446:ILE:HA	22:A:854:CLA:CGA	2.25	0.67
22:A:815:CLA:H43	24:3:306:BCR:HC21	1.77	0.67
22:A:840:CLA:HMB1	22:A:840:CLA:HBB1	1.77	0.67
2:B:453:GLN:HE21	2:B:455:LEU:HD11	1.58	0.67
39:5:316:CHL:HHC	39:5:316:CHL:HBB1	1.76	0.67
18:6:108:PHE:HD2	18:6:113:LEU:HD21	1.60	0.67
22:A:804:CLA:HAB	22:A:806:CLA:CAD	2.24	0.67
7:G:91:THR:HG21	7:G:98:THR:HA	1.77	0.67
7:G:99:LEU:HD12	7:G:99:LEU:H	1.60	0.67
13:3:146:LEU:HD22	24:3:305:BCR:HC32	1.76	0.67
22:5:309:CLA:H12	22:5:309:CLA:NA	2.09	0.67
1:A:453:PHE:CE1	22:B:806:CLA:HMA1	2.30	0.67
22:B:827:CLA:H2	22:B:839:CLA:HBD	1.76	0.67
7:G:77:TYR:HD2	7:G:79:ASP:HB3	1.59	0.67
40:1:318:SQD:C7	40:1:318:SQD:H121	2.25	0.67
17:5:85:VAL:HG11	38:5:302:LUT:H12	1.77	0.67
19:9:145:ASP:H	19:9:146:PRO:HD2	1.59	0.67
22:B:811:CLA:CBA	22:B:829:CLA:HBD	2.24	0.66
4:D:178:ASN:HB3	4:D:191:THR:HG22	1.77	0.66

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
22:1:308:CLA:HMD2	22:1:314:CLA:C1D	2.25	0.66
38:3:302:LUT:C30	22:3:308:CLA:H72	2.24	0.66
22:B:817:CLA:HBB1	22:B:817:CLA:HMB1	1.77	0.66
38:1:303:LUT:H8	22:1:309:CLA:HBB1	1.78	0.66
22:1:308:CLA:HMD2	22:1:314:CLA:ND	2.10	0.66
38:Z:301:LUT:H26	22:Z:303:CLA:H2	1.76	0.66
14:7:209:ALA:HB2	22:7:317:CLA:H8	1.77	0.66
39:8:301:CHL:HHC	39:8:301:CHL:HBB1	1.76	0.66
18:6:64:LEU:HD12	22:6:310:CLA:HMA2	1.75	0.66
22:A:810:CLA:H161	24:A:845:BCR:H372	1.77	0.66
4:D:189:ASN:HD22	4:D:194:ILE:HD13	1.59	0.66
38:7:302:LUT:H28	22:7:307:CLA:C6	2.25	0.66
15:8:156:PHE:CE1	22:8:313:CLA:HAB	2.30	0.66
24:5:304:BCR:HC8	22:5:326:CLA:H72	1.77	0.66
22:3:310:CLA:H52	22:3:310:CLA:NC	2.10	0.66
18:6:62:GLU:HA	18:6:65:LYS:HD3	1.76	0.66
2:B:475:PHE:HB3	22:F:304:CLA:H191	1.77	0.66
22:B:831:CLA:H52	31:B:852:DGD:HB52	1.77	0.66
22:8:306:CLA:HMD3	33:4:801:LMT:H82	1.77	0.66
16:4:95:TRP:CZ2	22:4:815:CLA:HAA2	2.27	0.66
1:A:114:SER:HB2	1:A:131:VAL:HG11	1.77	0.66
22:5:326:CLA:H71	22:6:314:CLA:H72	1.78	0.66
1:A:722:GLN:HA	25:A:850:LHG:HC5	1.78	0.66
33:G:204:LMT:H51	22:1:314:CLA:H2	1.78	0.66
23:A:843:PQN:H193	24:F:303:BCR:H23C	1.76	0.66
2:B:89:ALA:HB2	2:B:117:SER:HB3	1.77	0.66
12:1:60:LEU:HD13	22:1:307:CLA:H42	1.76	0.66
18:6:99:ASP:HB3	39:6:315:CHL:HED3	1.77	0.66
12:1:57:PRO:HD2	38:1:303:LUT:H23	1.77	0.66
15:8:63:ARG:HG2	15:8:137:PHE:CE2	2.31	0.66
17:5:205:GLN:HG3	17:5:253:TRP:HZ3	1.61	0.66
12:1:39:SER:HA	15:8:135:MET:HE1	1.78	0.66
38:9:301:LUT:H26	22:9:303:CLA:O1A	1.95	0.66
22:5:309:CLA:HBA1	22:5:309:CLA:CBD	2.24	0.65
1:A:680:ALA:HB3	22:B:801:CLA:HBB2	1.77	0.65
22:A:818:CLA:H41	22:A:836:CLA:HMA2	1.79	0.65
22:A:824:CLA:HBB1	22:A:824:CLA:HHC	1.78	0.65
13:3:89:ILE:HG12	22:3:313:CLA:HED2	1.78	0.65
14:7:49:GLY:H	14:7:191:GLN:HE22	1.44	0.65
17:5:230:ASN:HA	17:5:233:HIS:NE2	2.12	0.65
39:9:312:CHL:HBC3	39:9:312:CHL:H151	1.77	0.65

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
38:3:302:LUT:H383	39:3:317:CHL:H11	1.79	0.65
22:3:312:CLA:O1D	22:3:318:CLA:H62	1.96	0.65
22:7:304:CLA:H101	22:7:310:CLA:H18	1.78	0.65
1:A:316:TRP:CD1	10:K:76:LEU:HD21	2.31	0.65
22:B:824:CLA:HMA2	33:1:301:LMT:H3'	1.78	0.65
22:8:311:CLA:HHC	22:8:311:CLA:HBB1	1.79	0.65
38:5:302:LUT:C11	22:5:308:CLA:HMC2	2.25	0.65
19:9:117:LEU:HD21	39:2:303:CHL:H2	1.77	0.65
1:A:449:GLY:HA3	22:A:854:CLA:O1A	1.95	0.65
22:A:814:CLA:HBA2	22:A:816:CLA:HHB	1.78	0.65
24:F:303:BCR:H12C	22:F:305:CLA:CAB	2.27	0.65
22:7:324:CLA:HMB1	22:7:324:CLA:HBB1	1.77	0.65
15:8:41:LYS:H	15:8:41:LYS:HD2	1.61	0.65
6:F:160:ALA:HB1	22:F:302:CLA:H8	1.79	0.65
33:G:204:LMT:H91	22:1:308:CLA:HMD3	1.77	0.65
24:3:307:BCR:H392	22:3:314:CLA:HAC2	1.78	0.65
16:4:67:PHE:CD1	16:4:80:GLY:HA3	2.31	0.65
22:A:822:CLA:H71	22:K:204:CLA:HAB	1.79	0.65
5:E:36:GLY:HA3	5:E:62:VAL:HG11	1.78	0.65
12:Z:158:ASP:HA	38:Z:301:LUT:H24	1.79	0.65
13:3:82:LYS:NZ	13:3:211:GLU:HG2	2.11	0.65
2:B:463:TRP:CD1	22:F:304:CLA:H193	2.32	0.65
2:B:487:ALA:O	2:B:491:GLN:HG3	1.97	0.65
22:B:814:CLA:H12	19:9:56:LEU:HD22	1.79	0.65
22:B:827:CLA:H8	22:B:839:CLA:HMD2	1.79	0.65
13:3:146:LEU:HD11	22:5:315:CLA:H171	1.77	0.65
22:3:322:CLA:HBB2	22:7:307:CLA:HHD	1.79	0.65
39:7:313:CHL:HHC	39:7:313:CHL:HBB1	1.77	0.65
39:5:316:CHL:H162	22:6:322:CLA:H193	1.78	0.65
22:B:833:CLA:HBC2	22:B:839:CLA:CMC	2.27	0.65
22:B:838:CLA:H171	32:F:306:RRX:C35	2.26	0.65
15:8:55:LEU:CD1	38:8:303:LUT:H221	2.25	0.65
15:8:125:MET:HB3	24:8:304:BCR:H15C	1.78	0.65
22:5:326:CLA:HBD	22:5:326:CLA:HBA1	1.79	0.65
22:A:814:CLA:H112	24:A:844:BCR:H353	1.79	0.64
22:A:833:CLA:HAB	22:A:834:CLA:HHB	1.77	0.64
22:A:833:CLA:NC	22:A:834:CLA:H101	2.12	0.64
39:Z:311:CHL:HHC	39:Z:311:CHL:HBB1	1.78	0.64
22:5:326:CLA:HMD1	22:6:309:CLA:H2	1.77	0.64
38:4:802:LUT:H383	39:4:814:CHL:H12	1.78	0.64
22:A:832:CLA:HMA2	2:B:685:LYS:HB2	1.79	0.64

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
22:B:838:CLA:HMB1	22:B:838:CLA:HBB1	1.79	0.64
17:5:131:TRP:CE3	22:6:314:CLA:HBA2	2.32	0.64
17:5:182:LEU:HB3	22:5:307:CLA:CMA	2.26	0.64
18:6:125:GLU:HG2	22:6:317:CLA:C1B	2.26	0.64
20:2:189:HIS:CG	22:2:301:CLA:HAA2	2.33	0.64
24:A:856:BCR:H323	24:3:306:BCR:HC41	1.78	0.64
22:B:810:CLA:H62	22:B:810:CLA:NC	2.11	0.64
22:3:320:CLA:CMA	22:3:322:CLA:H112	2.28	0.64
22:7:306:CLA:HMB1	22:7:306:CLA:CBB	2.27	0.64
15:8:62:LEU:HD12	22:8:308:CLA:HMA2	1.80	0.64
24:8:304:BCR:H292	22:8:310:CLA:HBA1	1.80	0.64
7:G:57:HIS:CE1	7:G:61:LEU:HD11	2.33	0.64
12:Z:61:CYS:HB2	22:Z:306:CLA:HBA1	1.80	0.64
13:3:122:GLN:O	13:3:126:LEU:HB2	1.97	0.64
22:4:805:CLA:H71	22:4:806:CLA:HMA1	1.77	0.64
19:9:102:LYS:HB3	19:9:104:TYR:CE2	2.32	0.64
22:A:836:CLA:HMB1	22:A:836:CLA:HBB1	1.80	0.64
22:B:830:CLA:H151	24:B:845:BCR:H15C	1.78	0.64
6:F:224:ILE:HB	6:F:227:ARG:HH22	1.61	0.64
7:G:35:TYR:HB3	22:G:201:CLA:HED2	1.80	0.64
38:Z:301:LUT:H382	22:Z:303:CLA:C4D	2.27	0.64
15:8:135:MET:HB3	15:8:143:GLN:NE2	2.13	0.64
22:8:309:CLA:HED2	22:8:309:CLA:H2A	1.80	0.64
38:4:802:LUT:H28	22:4:805:CLA:H2	1.78	0.64
24:4:804:BCR:H19C	39:4:816:CHL:O1A	1.98	0.64
1:A:584:CYS:SG	29:B:802:SF4:FE2	1.90	0.64
22:A:821:CLA:H72	24:A:846:BCR:H361	1.80	0.64
22:B:816:CLA:H61	24:B:846:BCR:H362	1.79	0.64
10:K:57:VAL:HG11	10:K:76:LEU:HD12	1.78	0.64
38:7:301:LUT:C30	22:7:304:CLA:H72	2.27	0.64
22:A:855:CLA:HAB	2:B:692:VAL:HG11	1.79	0.64
3:C:51:CYS:SG	29:C:101:SF4:FE1	1.89	0.64
13:3:98:PRO:HD2	22:3:316:CLA:C1B	2.28	0.64
38:5:303:LUT:H28	22:5:310:CLA:H72	1.78	0.64
24:3:306:BCR:H19C	22:3:312:CLA:C2C	2.28	0.64
17:5:57:PRO:HG2	22:5:315:CLA:H12	1.78	0.64
18:6:123:PHE:CD1	24:6:305:BCR:H10C	2.33	0.64
18:6:247:PRO:HG2	24:6:306:BCR:H272	1.79	0.64
22:A:832:CLA:H112	22:L:202:CLA:H171	1.80	0.64
2:B:394:PHE:CD2	24:B:848:BCR:H281	2.32	0.64
22:B:810:CLA:HMB2	24:I:4001:BCR:H322	1.80	0.64

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
14:7:178:PHE:CE2	38:7:301:LUT:H221	2.32	0.64
38:6:304:LUT:H393	22:6:310:CLA:H141	1.80	0.64
22:A:806:CLA:H43	25:A:850:LHG:H262	1.79	0.63
2:B:283:VAL:HG13	24:B:844:BCR:C36	2.28	0.63
22:1:313:CLA:HBA1	22:1:313:CLA:H101	1.78	0.63
22:Z:306:CLA:HMD2	39:Z:311:CHL:CBB	2.28	0.63
39:7:313:CHL:C4C	22:7:316:CLA:HMC3	2.28	0.63
22:8:306:CLA:HMD2	22:8:311:CLA:C4D	2.28	0.63
7:G:78:PHE:CE1	7:G:81:LEU:HD21	2.32	0.63
7:G:78:PHE:CZ	7:G:81:LEU:HD21	2.33	0.63
12:1:223:PRO:HG3	22:1:316:CLA:HAB	1.79	0.63
13:3:22:THR:HA	13:3:25:TYR:CE2	2.34	0.63
16:4:228:PHE:CD1	16:4:239:PRO:HB3	2.33	0.63
18:6:88:GLN:HE22	38:6:304:LUT:H42	1.62	0.63
22:A:810:CLA:H71	22:3:312:CLA:H51	1.80	0.63
22:B:821:CLA:HMB2	22:B:826:CLA:HMA3	1.79	0.63
24:B:847:BCR:H331	24:B:847:BCR:HC8	1.80	0.63
24:B:853:BCR:H403	24:I:4001:BCR:H372	1.80	0.63
22:F:302:CLA:C2B	24:F:303:BCR:H10C	2.28	0.63
13:3:195:VAL:HG11	38:3:303:LUT:C12	2.28	0.63
24:3:307:BCR:H403	24:3:307:BCR:H23C	1.78	0.63
22:7:317:CLA:H11	22:7:317:CLA:HMA2	1.79	0.63
22:A:838:CLA:H52	22:A:838:CLA:C1C	2.29	0.63
10:K:57:VAL:O	10:K:58:LYS:HB3	1.97	0.63
12:1:194:ALA:HB1	12:1:218:ASN:HD21	1.64	0.63
13:3:42:LEU:HB2	22:3:311:CLA:HBA1	1.78	0.63
1:A:514:VAL:HG22	1:A:524:MET:HG3	1.79	0.63
2:B:420:ILE:HD13	2:B:537:LYS:HG2	1.81	0.63
22:B:809:CLA:H43	22:B:831:CLA:H2	1.78	0.63
22:B:834:CLA:HMB1	22:B:834:CLA:HBB1	1.80	0.63
22:F:301:CLA:HMC2	24:F:303:BCR:H381	1.79	0.63
11:L:87:GLU:HA	11:L:90:ARG:NH2	2.13	0.63
1:A:690:ARG:HG3	1:A:717:ALA:HB3	1.80	0.63
21:A:801:CL0:H15	21:A:801:CL0:H11	1.80	0.63
22:A:816:CLA:CBB	24:A:844:BCR:H12C	2.28	0.63
22:A:827:CLA:H193	22:A:827:CLA:H152	1.79	0.63
2:B:27:ALA:HB1	31:B:852:DGD:HG2	1.79	0.63
31:1:319:DGD:HG11	31:1:319:DGD:HB21	1.79	0.63
18:6:88:GLN:HG2	22:6:312:CLA:C4D	2.29	0.63
22:A:822:CLA:HBB1	37:K:201:DAO:H112	1.79	0.63
22:B:811:CLA:HBA2	22:B:829:CLA:CBD	2.27	0.63

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
38:1:303:LUT:H373	22:1:307:CLA:H91	1.78	0.63
13:3:71:ALA:HB1	38:3:302:LUT:H12	1.81	0.63
1:A:598:TRP:CH2	22:B:806:CLA:HAB	2.34	0.63
22:B:827:CLA:HBA2	22:B:839:CLA:HAA1	1.81	0.63
6:F:69:PRO:HG2	6:F:72:GLU:HB2	1.81	0.63
14:7:70:HIS:O	14:7:74:GLN:HG2	1.99	0.63
22:7:322:CLA:H2	22:7:323:CLA:H101	1.80	0.63
1:A:244:LEU:HD12	22:A:815:CLA:HMA3	1.81	0.63
22:A:822:CLA:H62	22:A:823:CLA:H92	1.81	0.63
22:A:855:CLA:H152	24:B:853:BCR:H17C	1.81	0.63
2:B:258:PHE:HZ	22:B:819:CLA:H111	1.63	0.63
12:1:168:GLU:CD	12:1:168:GLU:H	2.02	0.63
17:5:155:ASN:HB3	39:5:321:CHL:C3D	2.27	0.63
22:A:810:CLA:H43	22:A:812:CLA:H42	1.81	0.62
22:A:833:CLA:HBA1	22:A:855:CLA:H2	1.81	0.62
12:1:153:PRO:HB3	22:1:313:CLA:HBC2	1.80	0.62
22:8:314:CLA:C4B	26:8:318:NKP:HAY	2.29	0.62
16:4:66:LYS:HD3	25:4:821:LHG:HC31	1.81	0.62
18:6:96:PHE:HB3	18:6:99:ASP:OD1	1.99	0.62
22:A:835:CLA:HBB1	22:A:835:CLA:HMB1	1.81	0.62
14:7:228:ASP:HB3	14:7:232:THR:HG23	1.81	0.62
18:6:238:ASP:CA	18:6:243:LYS:HA	2.17	0.62
23:A:843:PQN:H212	22:F:301:CLA:C3B	2.29	0.62
4:D:115:PHE:HB3	4:D:120:GLN:HE21	1.63	0.62
14:7:133:TRP:CH2	24:7:303:BCR:HC8	2.34	0.62
15:8:101:TYR:HA	39:8:312:CHL:HMA3	1.80	0.62
22:8:305:CLA:HMD2	22:8:313:CLA:CGA	2.30	0.62
22:9:307:CLA:HMB1	22:9:307:CLA:HBB1	1.81	0.62
39:9:312:CHL:HED1	39:9:312:CHL:H93	1.80	0.62
7:G:93:ASP:HB2	22:G:202:CLA:CHA	2.29	0.62
14:7:175:PRO:HD2	38:7:301:LUT:H23	1.80	0.62
15:8:135:MET:HB3	15:8:143:GLN:HE22	1.63	0.62
38:8:303:LUT:H371	22:8:308:CLA:H142	1.81	0.62
22:2:304:CLA:HBB1	22:2:304:CLA:HHC	1.82	0.62
2:B:491:GLN:HA	2:B:495:LEU:HD12	1.82	0.62
22:1:307:CLA:HMB1	22:1:307:CLA:HBB1	1.81	0.62
22:1:310:CLA:HBA1	22:1:310:CLA:CHA	2.28	0.62
12:Z:103:PRO:HG2	39:Z:312:CHL:HED1	1.80	0.62
16:4:86:LEU:HD23	33:4:801:LMT:H2'	1.81	0.62
17:5:171:PRO:HD2	38:5:302:LUT:H23	1.82	0.62
19:9:101:ALA:HB2	39:9:312:CHL:HBA2	1.81	0.62

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
19:9:194:PRO:HB3	22:9:310:CLA:HMB3	1.81	0.62
22:A:815:CLA:H41	24:3:306:BCR:HC8	1.81	0.62
22:A:818:CLA:H8	22:A:836:CLA:CMA	2.30	0.62
22:B:820:CLA:HMB2	22:B:820:CLA:H51	1.80	0.62
22:B:834:CLA:NC	24:J:104:BCR:H292	2.15	0.62
22:3:309:CLA:HMD3	22:3:314:CLA:H43	1.82	0.62
24:8:304:BCR:H19C	39:8:315:CHL:HMA2	1.82	0.62
1:A:59:PHE:HA	1:A:62:HIS:ND1	2.15	0.62
22:A:817:CLA:HBB1	22:A:817:CLA:HMB1	1.80	0.62
22:A:823:CLA:HBC2	22:A:824:CLA:HAA1	1.81	0.62
38:3:303:LUT:H26	22:3:311:CLA:H2	1.82	0.62
22:3:318:CLA:HMB1	22:3:318:CLA:HBB1	1.80	0.62
22:7:308:CLA:HMD2	22:7:315:CLA:C1D	2.30	0.62
22:A:806:CLA:HMD1	22:A:811:CLA:H162	1.81	0.62
11:L:135:TYR:HB2	24:L:204:BCR:H361	1.81	0.62
12:Z:70:PHE:HB3	22:Z:306:CLA:HMA1	1.82	0.62
13:3:26:LEU:HD13	13:3:35:GLY:HA2	1.81	0.62
22:5:309:CLA:HHC	22:5:309:CLA:HBB1	1.82	0.62
22:4:811:CLA:HHC	22:4:811:CLA:HBB1	1.82	0.62
2:B:175:ARG:HE	22:B:826:CLA:HMD1	1.64	0.62
22:Z:309:CLA:H102	22:Z:309:CLA:H142	1.80	0.62
14:7:97:LEU:HD12	22:6:301:CLA:HMA2	1.82	0.62
19:9:124:THR:HG21	39:2:303:CHL:HED3	1.80	0.62
1:A:153:ILE:HA	1:A:158:GLN:OE1	2.00	0.61
1:A:541:PHE:HZ	22:B:806:CLA:HBB2	1.65	0.61
24:A:844:BCR:H17C	24:A:845:BCR:H382	1.80	0.61
2:B:6:PHE:HB3	2:B:7:PRO:HD3	1.82	0.61
23:B:843:PQN:H142	24:B:849:BCR:H281	1.81	0.61
13:3:82:LYS:HD2	13:3:210:TYR:HD2	1.64	0.61
22:3:313:CLA:H202	22:5:320:CLA:HHB	1.82	0.61
24:7:303:BCR:H332	25:8:317:LHG:H252	1.82	0.61
13:3:163:TRP:HH2	24:3:305:BCR:H10C	1.65	0.61
22:7:322:CLA:ND	22:7:323:CLA:H122	2.16	0.61
15:8:170:PRO:HD2	38:8:302:LUT:H24	1.82	0.61
16:4:113:GLY:HA3	38:4:803:LUT:H182	1.83	0.61
19:9:56:LEU:HB2	22:9:306:CLA:H11	1.81	0.61
20:2:166:LEU:HA	20:2:169:ILE:HG12	1.81	0.61
2:B:597:TRP:CH2	2:B:613:SER:HB2	2.35	0.61
22:B:818:CLA:HBB1	22:B:818:CLA:HHC	1.82	0.61
12:1:140:ARG:HG2	22:1:313:CLA:HHC	1.81	0.61
22:1:306:CLA:HBC1	22:1:316:CLA:H101	1.81	0.61

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
24:6:306:BCR:H16C	39:6:315:CHL:C4C	2.29	0.61
38:9:302:LUT:H11	22:9:307:CLA:HAC2	1.82	0.61
1:A:508:ALA:HB3	1:A:523:MET:HG3	1.81	0.61
2:B:95:PRO:HG3	22:L:201:CLA:HMB2	1.80	0.61
22:B:810:CLA:H11	8:I:79:PHE:HB3	1.81	0.61
22:F:301:CLA:H121	9:J:14:VAL:HG13	1.81	0.61
7:G:107:ALA:HA	24:G:203:BCR:H341	1.82	0.61
38:1:303:LUT:H161	22:1:309:CLA:HMB3	1.82	0.61
12:Z:157:PHE:HA	22:Z:313:CLA:H72	1.81	0.61
15:8:80:VAL:HG13	15:8:202:PHE:HE2	1.64	0.61
39:4:816:CHL:HMB1	39:4:816:CHL:CBB	2.28	0.61
18:6:226:VAL:HG21	41:6:324:3PH:H221	1.80	0.61
22:A:829:CLA:H11	24:A:845:BCR:H311	1.82	0.61
8:I:67:TYR:CE2	8:I:69:PHE:HB2	2.36	0.61
22:5:307:CLA:H93	22:5:308:CLA:H62	1.81	0.61
18:6:188:PHE:CE1	18:6:199:PRO:HB3	2.36	0.61
22:A:829:CLA:H42	22:A:829:CLA:HBD	1.82	0.61
23:A:843:PQN:H212	22:F:301:CLA:C4B	2.31	0.61
12:1:98:ASN:HD21	12:1:101:ASP:HB2	1.63	0.61
38:Z:301:LUT:H28	22:Z:303:CLA:H52	1.83	0.61
13:3:120:ALA:HB2	22:3:319:CLA:H2	1.82	0.61
16:4:243:LEU:HB2	38:4:802:LUT:H22	1.81	0.61
22:6:312:CLA:H52	22:6:312:CLA:C3D	2.30	0.61
19:9:120:GLY:HA2	22:9:313:CLA:HAB	1.81	0.61
1:A:197:MET:HG3	22:A:813:CLA:HBC2	1.83	0.61
1:A:545:VAL:HG11	1:A:598:TRP:CE2	2.35	0.61
2:B:54:GLN:HG2	22:B:808:CLA:CMA	2.30	0.61
22:L:201:CLA:H162	22:L:201:CLA:H91	1.81	0.61
18:6:147:ASN:HB3	39:6:320:CHL:C4D	2.30	0.61
20:2:193:PRO:HG3	22:2:302:CLA:HMB3	1.82	0.61
1:A:409:ALA:HB1	24:A:846:BCR:H271	1.83	0.61
1:A:694:GLN:HE21	5:E:80:TYR:HE1	1.49	0.61
22:A:833:CLA:HAB	22:A:834:CLA:CHB	2.30	0.61
22:B:817:CLA:HHB	24:B:846:BCR:HC8	1.83	0.61
22:B:824:CLA:HMC1	22:B:825:CLA:H43	1.82	0.61
13:3:190:LEU:HB2	24:3:307:BCR:H393	1.82	0.61
22:3:312:CLA:H112	22:3:318:CLA:H42	1.81	0.61
22:A:836:CLA:NB	22:K:203:CLA:H52	2.15	0.61
22:A:854:CLA:H142	24:I:4001:BCR:H271	1.83	0.61
2:B:413:LEU:HB3	6:F:227:ARG:HG3	1.82	0.61
22:B:812:CLA:H71	24:B:849:BCR:H343	1.82	0.61

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
13:3:38:PRO:HD2	38:3:303:LUT:H23	1.82	0.61
24:3:304:BCR:H382	24:3:305:BCR:H351	1.82	0.61
38:7:301:LUT:H383	22:7:314:CLA:H11	1.82	0.61
22:5:312:CLA:HMA2	22:5:319:CLA:HAC2	1.81	0.61
22:9:307:CLA:H2	22:9:307:CLA:HED2	1.81	0.61
2:B:694:TRP:CD1	2:B:698:PRO:HD3	2.36	0.61
25:B:803:LHG:HC5	22:B:807:CLA:HBA2	1.83	0.61
12:Z:188:PHE:HA	12:Z:191:GLN:NE2	2.15	0.61
22:4:815:CLA:H43	33:4:822:LMT:H91	1.83	0.61
17:5:131:TRP:CD1	24:5:304:BCR:H12C	2.36	0.61
38:5:303:LUT:C28	22:5:310:CLA:H72	2.31	0.61
24:5:304:BCR:H342	22:5:326:CLA:H61	1.83	0.61
2:B:366:PHE:HB3	2:B:603:TRP:CZ3	2.36	0.60
13:3:42:LEU:HD13	22:3:311:CLA:HMA2	1.81	0.60
16:4:159:PHE:CD2	24:4:804:BCR:H12C	2.36	0.60
18:6:95:VAL:HG13	18:6:103:LYS:HB2	1.82	0.60
38:6:303:LUT:H11	22:6:308:CLA:HMC2	1.83	0.60
19:9:84:ILE:HG23	19:9:96:TRP:CG	2.36	0.60
19:9:110:PRO:HG3	20:2:199:TYR:HB2	1.82	0.60
22:A:818:CLA:HMD2	22:A:827:CLA:HMB2	1.83	0.60
14:7:26:TYR:CE2	14:7:191:GLN:HB2	2.37	0.60
24:8:304:BCR:H282	22:8:310:CLA:NB	2.17	0.60
17:5:205:GLN:HG3	17:5:253:TRP:CZ3	2.35	0.60
22:9:311:CLA:HED3	22:9:311:CLA:H2A	1.83	0.60
1:A:211:LEU:HD23	24:A:844:BCR:H373	1.84	0.60
1:A:363:LEU:HD21	22:A:819:CLA:H72	1.83	0.60
22:A:807:CLA:HHB	22:A:808:CLA:HMB3	1.82	0.60
22:B:828:CLA:H71	24:B:848:BCR:H16C	1.82	0.60
14:7:49:GLY:HA3	14:7:194:ILE:HG21	1.83	0.60
15:8:27:ARG:HD3	15:8:47:ASP:O	2.01	0.60
16:4:86:LEU:HD11	38:4:803:LUT:H221	1.83	0.60
38:4:803:LUT:H26	22:4:808:CLA:H52	1.83	0.60
38:9:301:LUT:H14	22:9:304:CLA:HBB1	1.83	0.60
1:A:157:LEU:HD23	22:A:816:CLA:HED1	1.82	0.60
22:Z:308:CLA:H61	22:Z:315:CLA:CAA	2.30	0.60
17:5:222:PRO:HG2	22:5:314:CLA:HMB3	1.83	0.60
22:6:312:CLA:HBB1	22:6:312:CLA:HMB1	1.83	0.60
22:A:821:CLA:HMB2	22:A:825:CLA:HMA3	1.83	0.60
4:D:131:LYS:HE3	4:D:132:PHE:CE2	2.35	0.60
22:1:305:CLA:ND	22:1:310:CLA:H8	2.17	0.60
40:1:318:SQD:H192	39:8:301:CHL:HMA2	1.83	0.60

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
22:8:310:CLA:H121	43:8:319:LPX:H8A	1.83	0.60
38:5:303:LUT:H372	22:5:315:CLA:H72	1.82	0.60
18:6:52:PHE:HB3	22:6:310:CLA:CAD	2.32	0.60
18:6:247:PRO:HB2	22:6:301:CLA:C1	2.30	0.60
22:A:828:CLA:H191	24:J:104:BCR:H19C	1.83	0.60
15:8:63:ARG:HG2	15:8:137:PHE:HE2	1.66	0.60
15:8:102:GLU:HA	39:8:312:CHL:HED3	1.84	0.60
22:A:823:CLA:HBA1	37:K:201:DAO:H51	1.82	0.60
22:B:828:CLA:HBA2	24:B:848:BCR:H10C	1.83	0.60
10:K:38:MET:HG2	22:K:202:CLA:C3D	2.31	0.60
10:K:97:HIS:ND1	24:K:206:BCR:H14C	2.15	0.60
22:1:306:CLA:C15	25:1:317:LHG:HC92	2.32	0.60
22:1:306:CLA:H172	25:1:317:LHG:H121	1.82	0.60
12:Z:144:GLU:HB3	12:Z:150:ARG:HG2	1.82	0.60
22:Z:304:CLA:HBC3	22:Z:309:CLA:H143	1.84	0.60
13:3:137:SER:HA	13:3:140:LYS:NZ	2.16	0.60
22:A:804:CLA:HBD	22:A:811:CLA:H2	1.83	0.60
22:A:811:CLA:HBB1	22:A:811:CLA:HHC	1.83	0.60
2:B:262:PHE:CE1	2:B:356:LEU:HA	2.37	0.60
15:8:174:CYS:HB2	22:8:305:CLA:HAA2	1.83	0.60
16:4:159:PHE:CD1	24:4:804:BCR:H10C	2.37	0.60
18:6:176:LYS:HD3	22:6:313:CLA:O1D	2.01	0.60
22:6:307:CLA:HBB1	22:6:307:CLA:HMB1	1.82	0.60
1:A:680:ALA:CB	22:B:801:CLA:HBB2	2.32	0.60
2:B:155:TRP:CZ2	2:B:159:GLN:HG2	2.37	0.60
2:B:183:LEU:HD13	22:B:815:CLA:HHB	1.83	0.60
7:G:46:LEU:HD12	22:G:201:CLA:H11	1.82	0.60
7:G:49:GLY:HA3	7:G:105:TRP:CD2	2.37	0.60
10:K:97:HIS:CE1	24:K:206:BCR:H14C	2.37	0.60
22:Z:304:CLA:C1C	22:Z:309:CLA:H121	2.31	0.60
22:5:307:CLA:H72	22:5:308:CLA:CMA	2.31	0.60
18:6:190:MET:HE1	22:6:319:CLA:H12	1.84	0.60
1:A:211:LEU:HD11	24:A:856:BCR:HC7	1.84	0.60
22:A:817:CLA:CAB	24:K:206:BCR:HC32	2.25	0.60
2:B:493:LEU:HD12	22:B:836:CLA:HED1	1.84	0.60
22:F:301:CLA:HHC	22:F:301:CLA:HBB1	1.84	0.60
16:4:228:PHE:HD1	16:4:239:PRO:HB3	1.67	0.60
17:5:98:VAL:HG11	22:5:312:CLA:HED2	1.84	0.60
22:5:318:CLA:HBC2	22:5:319:CLA:HMB1	1.83	0.60
18:6:173:LYS:HD3	22:6:308:CLA:O1A	2.02	0.60
1:A:136:GLN:HE22	6:F:104:LEU:HD11	1.67	0.59

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:B:70:ALA:HB2	2:B:136:LEU:HB2	1.84	0.59
22:B:811:CLA:HBA1	22:B:811:CLA:CHA	2.31	0.59
22:G:201:CLA:HHC	22:G:201:CLA:CBB	2.31	0.59
17:5:181:GLU:OE2	17:5:185:LYS:HE2	2.02	0.59
22:A:854:CLA:HMB1	22:A:854:CLA:HBB1	1.84	0.59
22:Z:310:CLA:HBA2	22:Z:310:CLA:CHA	2.29	0.59
39:4:813:CHL:HAB	39:4:816:CHL:CBB	2.32	0.59
24:5:305:BCR:H333	18:6:227:VAL:HG21	1.83	0.59
19:9:115:GLU:HB2	39:9:312:CHL:HBB2	1.83	0.59
22:A:805:CLA:HBD	22:A:825:CLA:H52	1.83	0.59
2:B:155:TRP:CH2	2:B:159:GLN:HG2	2.37	0.59
24:B:853:BCR:HC42	22:L:202:CLA:NC	2.17	0.59
5:E:39:ARG:HH21	5:E:62:VAL:H	1.49	0.59
13:3:82:LYS:NZ	13:3:210:TYR:HB3	2.16	0.59
22:8:308:CLA:HBC3	41:8:320:3PH:H3C1	1.83	0.59
16:4:177:ASP:HB3	16:4:180:PHE:O	2.02	0.59
22:4:818:CLA:HBA1	22:4:818:CLA:CHA	2.32	0.59
22:6:309:CLA:HBC1	22:6:319:CLA:HAA1	1.83	0.59
39:9:312:CHL:HBD	39:9:312:CHL:HAA2	1.83	0.59
22:A:821:CLA:C4C	22:A:827:CLA:H161	2.32	0.59
2:B:195:LEU:HA	2:B:199:ALA:HB3	1.84	0.59
22:B:814:CLA:HHC	22:B:814:CLA:CBB	2.32	0.59
22:B:819:CLA:HMD2	22:B:828:CLA:HMB2	1.82	0.59
12:Z:157:PHE:CE2	22:Z:313:CLA:H161	2.37	0.59
13:3:190:LEU:HB2	24:3:307:BCR:C39	2.33	0.59
22:5:310:CLA:H101	22:5:315:CLA:H121	1.85	0.59
22:5:326:CLA:HHC	22:5:326:CLA:HBB1	1.83	0.59
1:A:320:HIS:HB3	1:A:325:ILE:HD11	1.83	0.59
22:A:834:CLA:HBB1	22:A:834:CLA:HMB1	1.85	0.59
14:7:126:THR:HG21	22:7:322:CLA:HMA2	1.84	0.59
24:7:303:BCR:H11C	22:7:322:CLA:H41	1.84	0.59
15:8:78:THR:HG23	22:8:305:CLA:CMC	2.30	0.59
16:4:138:GLY:HA2	39:4:816:CHL:HAC2	1.83	0.59
22:B:811:CLA:HBB2	22:B:812:CLA:HAA1	1.84	0.59
22:8:309:CLA:H43	22:8:309:CLA:HED3	1.84	0.59
22:4:806:CLA:HMD2	22:4:811:CLA:C1D	2.32	0.59
1:A:598:TRP:HE1	22:A:854:CLA:C1D	2.15	0.59
22:A:814:CLA:H141	31:3:301:DGD:HA41	1.84	0.59
2:B:349:VAL:HA	22:B:820:CLA:H42	1.85	0.59
14:7:49:GLY:H	14:7:191:GLN:NE2	2.01	0.59
39:8:301:CHL:HAA2	39:8:301:CHL:HBD	1.84	0.59

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
38:4:803:LUT:C32	22:4:809:CLA:HMB2	2.32	0.59
24:5:305:BCR:H343	18:6:220:VAL:HG11	1.84	0.59
20:2:207:THR:HG23	22:2:301:CLA:H8	1.82	0.59
22:A:855:CLA:H102	22:B:840:CLA:H43	1.85	0.59
2:B:78:TRP:HA	2:B:85:VAL:HB	1.85	0.59
2:B:229:GLY:HA3	7:G:117:LEU:HD13	1.84	0.59
22:B:814:CLA:O1A	7:G:81:LEU:HD22	2.02	0.59
22:Z:303:CLA:H102	22:Z:304:CLA:HMB3	1.84	0.59
22:Z:304:CLA:HMD2	22:Z:309:CLA:C4D	2.33	0.59
15:8:133:ARG:HH12	22:8:314:CLA:HED3	1.67	0.59
22:5:322:CLA:CAD	22:6:314:CLA:HMA1	2.33	0.59
18:6:39:PRO:HB2	18:6:41:HIS:CE1	2.37	0.59
38:Z:302:LUT:H393	22:Z:306:CLA:H141	1.85	0.59
24:5:305:BCR:C12	24:5:305:BCR:H341	2.32	0.59
18:6:41:HIS:CD2	18:6:42:LEU:HG	2.38	0.59
18:6:245:PHE:CZ	18:6:247:PRO:HB3	2.36	0.59
22:6:314:CLA:HHC	22:6:314:CLA:CBB	2.31	0.59
2:B:182:GLY:HA3	22:B:815:CLA:HBB1	1.84	0.59
22:B:822:CLA:HBA1	7:G:100:ILE:HG22	1.85	0.59
12:Z:135:ALA:HA	22:Z:314:CLA:HAB	1.84	0.59
22:Z:309:CLA:HHC	22:Z:309:CLA:HBB1	1.85	0.59
13:3:217:LEU:HD21	22:3:315:CLA:HMC3	1.85	0.59
22:3:322:CLA:H52	25:7:318:LHG:H321	1.84	0.59
22:7:312:CLA:HHC	22:7:312:CLA:CBB	2.32	0.59
38:9:301:LUT:H181	22:9:305:CLA:HBB1	1.85	0.59
1:A:217:GLN:HA	1:A:221:SER:HB2	1.84	0.58
22:A:814:CLA:H11	22:A:816:CLA:CHB	2.33	0.58
22:B:811:CLA:HBA1	22:B:811:CLA:HBD	1.85	0.58
6:F:141:LEU:HD21	6:F:150:PHE:CE2	2.37	0.58
22:F:304:CLA:H12	22:F:304:CLA:NA	2.17	0.58
12:Z:153:PRO:HB3	22:Z:313:CLA:HBC2	1.85	0.58
15:8:40:LEU:HD11	15:8:51:ASP:HB2	1.84	0.58
22:B:811:CLA:HHC	22:B:811:CLA:CBB	2.28	0.58
4:D:184:VAL:HG11	4:D:194:ILE:HG21	1.85	0.58
15:8:39:HIS:HA	15:8:41:LYS:NZ	2.18	0.58
15:8:205:GLN:HB3	15:8:216:ASN:ND2	2.18	0.58
22:B:806:CLA:H171	22:L:201:CLA:HMC2	1.85	0.58
22:B:819:CLA:HHC	22:B:819:CLA:HBB1	1.85	0.58
22:B:827:CLA:H121	22:B:842:CLA:H202	1.86	0.58
6:F:165:TYR:O	6:F:169:LEU:HG	2.03	0.58
10:K:62:ASP:HB2	10:K:68:VAL:CG2	2.33	0.58

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
10:K:66:LYS:HB2	10:K:66:LYS:NZ	2.18	0.58
11:L:73:PRO:HB3	11:L:87:GLU:OE1	2.02	0.58
15:8:184:LEU:HB3	22:8:305:CLA:HMA1	1.84	0.58
16:4:116:VAL:HB	16:4:117:PRO:HD3	1.86	0.58
22:4:809:CLA:OBD	22:4:815:CLA:HAA1	2.04	0.58
17:5:160:PRO:HG2	17:5:179:ARG:NH1	2.18	0.58
17:5:176:LYS:HD2	17:5:177:GLY:N	2.19	0.58
22:A:824:CLA:CMD	22:A:825:CLA:HAB	2.34	0.58
24:A:856:BCR:C24	10:K:95:LEU:HD21	2.34	0.58
2:B:10:SER:HB3	31:B:852:DGD:HE2	1.84	0.58
2:B:168:TRP:CE2	22:B:813:CLA:HMA1	2.38	0.58
22:B:814:CLA:H43	19:9:56:LEU:HD22	1.85	0.58
22:B:842:CLA:CHC	22:1:308:CLA:H12	2.34	0.58
38:1:302:LUT:H28	22:1:304:CLA:H2	1.83	0.58
13:3:129:TRP:CZ3	41:7:319:3PH:H322	2.39	0.58
22:7:308:CLA:HMD2	22:7:315:CLA:CHD	2.32	0.58
39:7:313:CHL:CBB	22:7:316:CLA:HBB2	2.33	0.58
38:5:302:LUT:H30	22:5:307:CLA:C5	2.33	0.58
1:A:59:PHE:CD2	22:A:805:CLA:HMC2	2.38	0.58
22:A:811:CLA:HBA1	22:A:811:CLA:CHA	2.33	0.58
11:L:141:GLN:H	11:L:141:GLN:NE2	2.00	0.58
12:Z:179:ARG:HA	12:Z:182:MET:HE2	1.85	0.58
38:3:303:LUT:H162	22:3:313:CLA:HMB3	1.85	0.58
38:4:802:LUT:H28	22:4:805:CLA:C2	2.34	0.58
25:4:821:LHG:H142	22:6:321:CLA:C4C	2.33	0.58
17:5:49:LEU:HD13	17:5:71:ARG:HD3	1.84	0.58
1:A:375:PRO:HG2	1:A:381:ALA:HB2	1.85	0.58
27:A:852:DGA:HG32	22:3:316:CLA:HMB2	1.84	0.58
22:A:855:CLA:HHC	22:A:855:CLA:HBB1	1.84	0.58
2:B:628:ASN:HB2	2:B:725:PHE:CE1	2.38	0.58
22:B:828:CLA:C7	24:B:848:BCR:H16C	2.34	0.58
22:B:841:CLA:CAC	24:B:849:BCR:H24C	2.34	0.58
12:Z:101:ASP:HA	12:Z:104:LEU:HD12	1.85	0.58
12:Z:115:PHE:CD2	22:Z:308:CLA:HMD2	2.38	0.58
38:Z:301:LUT:H8	22:Z:305:CLA:HBB1	1.86	0.58
22:Z:307:CLA:H52	15:8:152:PHE:HB2	1.84	0.58
13:3:203:VAL:HG11	31:3:301:DGD:HA72	1.86	0.58
22:5:313:CLA:HBC1	25:5:324:LHG:H302	1.86	0.58
22:A:814:CLA:HMC2	24:A:844:BCR:H352	1.85	0.58
22:A:818:CLA:H8	22:A:836:CLA:HMA2	1.86	0.58
6:F:203:PRO:HB2	22:8:309:CLA:H101	1.86	0.58

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
11:L:96:LEU:HD11	11:L:124:LEU:HD23	1.84	0.58
12:Z:203:LEU:HA	22:Z:305:CLA:CMA	2.34	0.58
17:5:164:TYR:HB3	22:5:307:CLA:HED3	1.85	0.58
38:9:301:LUT:H161	22:9:305:CLA:HMB3	1.84	0.58
2:B:243:HIS:HA	2:B:249:ASP:O	2.04	0.58
2:B:284:ILE:HG23	22:B:818:CLA:H192	1.86	0.58
2:B:464:ILE:HG12	22:B:835:CLA:HMC3	1.86	0.58
4:D:178:ASN:HB3	4:D:191:THR:CG2	2.33	0.58
5:E:74:ARG:HD2	5:E:85:THR:HB	1.85	0.58
24:7:303:BCR:H292	42:7:321:SPH:H122	1.85	0.58
39:8:315:CHL:HAA2	39:8:315:CHL:HBD	1.86	0.58
22:4:809:CLA:HMD2	22:4:815:CLA:ND	2.19	0.58
39:4:813:CHL:HAA1	39:4:813:CHL:HBD	1.85	0.58
17:5:247:ILE:HG23	39:5:316:CHL:H42	1.86	0.58
22:5:313:CLA:HBC2	25:5:324:LHG:HC91	1.86	0.58
22:A:807:CLA:H41	22:J:103:CLA:HMC1	1.84	0.58
4:D:182:ILE:HG23	4:D:183:LYS:HD2	1.85	0.58
22:F:304:CLA:C8	22:F:304:CLA:H2	2.33	0.58
7:G:45:MET:HG3	7:G:109:GLY:HA2	1.85	0.58
10:K:25:ARG:HA	10:K:29:PHE:CE1	2.39	0.58
40:1:318:SQD:H132	15:8:134:TRP:CE2	2.39	0.58
1:A:190:TRP:CZ2	22:A:810:CLA:HMA1	2.39	0.58
2:B:341:SER:HA	22:B:828:CLA:H51	1.84	0.58
12:1:37:PRO:O	39:8:301:CHL:HED1	2.03	0.58
17:5:84:GLY:HA2	38:5:303:LUT:H181	1.86	0.58
18:6:108:PHE:HB2	18:6:113:LEU:HG	1.85	0.58
22:2:301:CLA:HBA1	22:2:301:CLA:CHA	2.31	0.58
22:B:821:CLA:HMB1	22:B:826:CLA:H2	1.86	0.57
22:A:816:CLA:HBA2	22:A:816:CLA:CHA	2.33	0.57
2:B:296:PHE:CD2	22:B:822:CLA:HAA2	2.39	0.57
2:B:296:PHE:CE2	22:B:822:CLA:HBD	2.39	0.57
22:B:832:CLA:C4D	22:B:842:CLA:H52	2.33	0.57
22:B:832:CLA:H42	22:B:842:CLA:H62	1.86	0.57
8:I:88:PRO:CG	24:I:4001:BCR:H12C	2.34	0.57
12:1:76:ILE:HG13	22:1:314:CLA:O1D	2.04	0.57
22:4:810:CLA:CGA	22:4:818:CLA:HAB	2.34	0.57
22:6:307:CLA:HMD2	39:6:316:CHL:CGA	2.34	0.57
39:6:320:CHL:HAA2	39:6:320:CHL:HBD	1.87	0.57
19:9:101:ALA:HB2	39:9:312:CHL:CBA	2.35	0.57
1:A:362:SER:O	1:A:366:ILE:HG13	2.04	0.57
1:A:670:LEU:HB3	22:A:802:CLA:H92	1.85	0.57

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
22:A:812:CLA:O1D	22:A:813:CLA:HMC1	2.03	0.57
22:A:835:CLA:ND	24:A:847:BCR:H282	2.19	0.57
22:A:841:CLA:HBA1	22:A:841:CLA:CHA	2.34	0.57
22:B:821:CLA:HMC2	22:B:826:CLA:H202	1.85	0.57
22:B:825:CLA:HMA1	22:B:842:CLA:HED2	1.86	0.57
6:F:153:THR:O	6:F:157:LEU:HG	2.04	0.57
8:I:87:LEU:HB3	8:I:88:PRO:HD3	1.84	0.57
12:Z:157:PHE:HB3	22:Z:303:CLA:HMD1	1.86	0.57
12:Z:157:PHE:HZ	22:Z:313:CLA:NB	2.01	0.57
14:7:153:PHE:HB3	14:7:160:PHE:HE2	1.69	0.57
15:8:129:VAL:HG11	24:8:304:BCR:H16C	1.86	0.57
22:4:809:CLA:HMA2	22:4:809:CLA:H11	1.85	0.57
39:9:312:CHL:HED1	39:9:312:CHL:C9	2.34	0.57
22:A:807:CLA:H102	36:J:105:C7Z:C11	2.34	0.57
22:A:810:CLA:HAB	22:A:813:CLA:H101	1.85	0.57
22:A:827:CLA:HMB3	22:A:835:CLA:H2	1.86	0.57
2:B:646:VAL:HA	22:B:812:CLA:HAC1	1.86	0.57
24:B:844:BCR:H343	7:G:104:ALA:HA	1.86	0.57
12:1:170:LEU:HB2	22:1:304:CLA:HMA1	1.86	0.57
24:3:304:BCR:H392	22:3:319:CLA:HAA1	1.86	0.57
22:7:306:CLA:H2	22:7:311:CLA:HMD1	1.86	0.57
25:4:820:LHG:H151	22:6:302:CLA:H72	1.84	0.57
22:5:311:CLA:HMD2	22:5:318:CLA:C1D	2.35	0.57
18:6:157:PRO:HG3	39:6:316:CHL:C1D	2.34	0.57
18:6:188:PHE:CD1	18:6:199:PRO:HB3	2.39	0.57
22:A:807:CLA:HMA1	22:A:808:CLA:HMB3	1.86	0.57
22:A:807:CLA:HAB	24:J:104:BCR:C6	2.34	0.57
22:A:832:CLA:H152	22:A:832:CLA:H8	1.87	0.57
22:B:824:CLA:O1A	22:B:824:CLA:HBD	2.05	0.57
22:F:304:CLA:H2	22:F:304:CLA:H92	1.86	0.57
22:1:310:CLA:HHC	22:1:310:CLA:CBB	2.32	0.57
22:Z:308:CLA:HMA2	22:Z:315:CLA:C3C	2.34	0.57
24:3:304:BCR:H292	22:5:301:CLA:HMB1	1.87	0.57
24:3:304:BCR:H313	25:7:318:LHG:H111	1.86	0.57
22:7:316:CLA:HBB1	22:7:316:CLA:HMB1	1.85	0.57
15:8:169:ASP:OD2	15:8:174:CYS:HB2	2.04	0.57
18:6:120:ALA:HB1	24:6:305:BCR:H15C	1.85	0.57
38:9:302:LUT:H371	22:9:306:CLA:H112	1.86	0.57
22:A:816:CLA:HBB2	24:A:844:BCR:H12C	1.87	0.57
24:A:844:BCR:H20C	24:A:845:BCR:H23C	1.87	0.57
2:B:681:TRP:O	2:B:685:LYS:HG2	2.05	0.57

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
4:D:174:ARG:HB2	4:D:177:GLU:HG3	1.85	0.57
8:I:75:ALA:HB3	8:I:76:PRO:HD3	1.87	0.57
8:I:98:TYR:HD2	8:I:99:ILE:HD12	1.68	0.57
22:3:314:CLA:HHC	22:3:314:CLA:HBB1	1.85	0.57
22:8:308:CLA:HBB1	22:8:308:CLA:HMB1	1.86	0.57
22:8:310:CLA:HBB1	22:8:310:CLA:HMB1	1.87	0.57
22:A:807:CLA:HMB1	24:J:104:BCR:HC7	1.85	0.57
22:B:809:CLA:HBB1	22:B:809:CLA:HMB1	1.87	0.57
22:B:827:CLA:HAA2	22:B:828:CLA:OBD	2.05	0.57
11:L:87:GLU:HA	11:L:90:ARG:HH21	1.68	0.57
22:L:202:CLA:HBC1	24:L:204:BCR:H23C	1.87	0.57
22:1:308:CLA:C4B	22:1:314:CLA:H161	2.34	0.57
13:3:159:PRO:HD2	22:3:308:CLA:OBD	2.04	0.57
22:3:312:CLA:C4B	22:3:318:CLA:H122	2.35	0.57
22:7:309:CLA:CHD	42:7:321:SPH:H151	2.35	0.57
15:8:115:LEU:HD21	39:8:312:CHL:HHD	1.85	0.57
39:4:813:CHL:HMB1	39:4:813:CHL:CBB	2.34	0.57
1:A:396:TRP:CD1	22:A:828:CLA:HAB	2.39	0.57
1:A:604:SER:O	1:A:607:ILE:HG12	2.04	0.57
21:A:801:CL0:H21	2:B:626:TRP:HD1	1.69	0.57
22:A:819:CLA:H172	22:A:827:CLA:O2A	2.05	0.57
3:C:51:CYS:HG	29:C:101:SF4:FE1	1.22	0.57
10:K:77:ILE:HD12	13:3:18:THR:HG23	1.86	0.57
38:1:303:LUT:H24	22:1:307:CLA:O1A	2.03	0.57
22:Z:316:CLA:NA	22:Z:316:CLA:H162	2.19	0.57
13:3:121:MET:HB2	22:3:318:CLA:HAB	1.86	0.57
13:3:149:VAL:CG2	13:3:163:TRP:HB2	2.35	0.57
14:7:234:PHE:CD1	22:7:306:CLA:H11	2.40	0.57
38:7:301:LUT:H403	22:7:304:CLA:H13	1.86	0.57
15:8:208:ALA:HA	15:8:234:ILE:HD13	1.87	0.57
24:8:304:BCR:H371	39:8:315:CHL:HMB2	1.87	0.57
16:4:71:LEU:HD21	22:4:808:CLA:HAA2	1.86	0.57
17:5:185:LYS:NZ	22:5:313:CLA:HBD	2.19	0.57
17:5:245:LEU:HD11	18:6:222:VAL:HB	1.86	0.57
19:9:118:ILE:HB	39:9:314:CHL:HMA2	1.86	0.57
1:A:160:TYR:O	1:A:164:ILE:HG12	2.03	0.57
12:1:102:ALA:H	12:1:103:PRO:HD2	1.69	0.57
12:Z:177:ASN:ND2	22:Z:309:CLA:HMD1	2.19	0.57
15:8:173:MET:HB2	22:8:305:CLA:HBA1	1.86	0.57
22:5:301:CLA:H51	22:5:301:CLA:ND	2.20	0.57
1:A:204:GLY:HA2	1:A:208:LEU:HD12	1.87	0.57

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
22:A:808:CLA:HHC	22:A:808:CLA:CBB	2.32	0.57
22:A:819:CLA:ND	22:A:829:CLA:H72	2.19	0.57
22:A:823:CLA:C1C	37:K:201:DAO:H111	2.34	0.57
2:B:142:PHE:HZ	22:B:816:CLA:H122	1.68	0.57
22:F:302:CLA:HHC	22:F:302:CLA:CBB	2.31	0.57
11:L:171:THR:O	11:L:175:THR:HG23	2.04	0.57
39:1:312:CHL:HBD	39:1:312:CHL:HAA1	1.87	0.57
22:3:313:CLA:H121	22:5:320:CLA:NC	2.20	0.57
14:7:25:ALA:HA	14:7:30:ARG:HB2	1.87	0.57
16:4:116:VAL:HG11	22:4:810:CLA:HAC2	1.87	0.57
39:5:316:CHL:H141	39:5:316:CHL:H172	1.86	0.57
18:6:182:MET:HE3	22:6:310:CLA:HMC3	1.87	0.57
39:6:318:CHL:HMB1	39:6:318:CHL:CBB	2.34	0.57
2:B:460:PHE:CG	22:F:304:CLA:HMB3	2.39	0.56
2:B:478:LEU:HD13	22:B:835:CLA:HMD3	1.86	0.56
22:G:202:CLA:NC	24:G:203:BCR:H282	2.20	0.56
39:4:814:CHL:HAA1	39:4:814:CHL:HBD	1.86	0.56
22:6:312:CLA:HMA2	39:6:318:CHL:HAC2	1.87	0.56
38:9:302:LUT:H162	22:9:308:CLA:HHB	1.87	0.56
20:2:178:ARG:NH1	20:2:203:VAL:HG22	2.19	0.56
1:A:355:ILE:HD13	24:A:846:BCR:H371	1.86	0.56
1:A:563:ILE:HD13	1:A:576:ASP:HB2	1.87	0.56
22:A:821:CLA:H102	22:A:824:CLA:H93	1.87	0.56
7:G:56:LEU:HD11	33:G:204:LMT:H3'	1.87	0.56
14:7:196:ASN:ND2	22:7:310:CLA:HMD1	2.19	0.56
16:4:159:PHE:CE1	24:4:804:BCR:HC8	2.40	0.56
25:4:820:LHG:H101	24:6:305:BCR:HC42	1.86	0.56
22:5:315:CLA:H2A	25:5:324:LHG:H151	1.85	0.56
18:6:128:ARG:HG3	39:6:316:CHL:C1D	2.35	0.56
1:A:28:LYS:HG3	22:A:811:CLA:HMA2	1.87	0.56
22:A:803:CLA:HHC	22:A:803:CLA:CBB	2.35	0.56
22:A:824:CLA:C2	22:A:835:CLA:H171	2.30	0.56
22:A:841:CLA:NC	22:A:841:CLA:H52	2.19	0.56
22:B:815:CLA:H102	24:B:845:BCR:H381	1.88	0.56
22:B:834:CLA:NA	24:J:104:BCR:H282	2.21	0.56
10:K:95:LEU:HD13	22:K:204:CLA:HBB2	1.88	0.56
38:1:303:LUT:H31	22:1:307:CLA:HMC2	1.86	0.56
14:7:163:LYS:HD3	14:7:164:GLU:HG3	1.86	0.56
15:8:193:ARG:HA	15:8:196:MET:HE2	1.87	0.56
38:8:302:LUT:C28	22:8:305:CLA:H61	2.35	0.56
22:8:307:CLA:H101	25:8:317:LHG:H312	1.86	0.56

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
22:8:310:CLA:HBD	43:8:319:LPX:H3	1.86	0.56
17:5:253:TRP:HD1	22:5:301:CLA:O1A	1.88	0.56
24:5:305:BCR:C14	22:6:322:CLA:HMA2	2.35	0.56
22:5:308:CLA:C4B	22:5:313:CLA:H121	2.35	0.56
38:6:304:LUT:C28	22:6:310:CLA:H61	2.26	0.56
19:9:62:ARG:HA	19:9:62:ARG:HE	1.69	0.56
19:9:112:THR:HG22	19:9:116:PHE:CZ	2.40	0.56
39:9:312:CHL:H141	39:9:312:CHL:CHD	2.35	0.56
1:A:26:PHE:HD2	34:J:101:LMG:HC1	1.69	0.56
1:A:622:VAL:HA	1:A:627:VAL:HG22	1.86	0.56
22:B:810:CLA:HAA2	8:I:80:VAL:HG22	1.87	0.56
39:Z:312:CHL:HAB	22:Z:315:CLA:HBB2	1.86	0.56
24:3:307:BCR:H352	22:3:315:CLA:C4D	2.34	0.56
15:8:118:LEU:HD21	39:8:315:CHL:CHD	2.35	0.56
16:4:81:PHE:HB3	22:4:808:CLA:C3D	2.35	0.56
18:6:245:PHE:CE2	22:6:301:CLA:H112	2.41	0.56
20:2:196:ASN:O	22:2:301:CLA:HAA1	2.06	0.56
1:A:303:VAL:HG12	22:A:817:CLA:H142	1.87	0.56
22:A:805:CLA:H171	24:A:844:BCR:H323	1.87	0.56
2:B:224:THR:HB	2:B:225:PRO:HD3	1.88	0.56
2:B:495:LEU:HB3	2:B:496:PRO:HD3	1.88	0.56
7:G:50:ARG:HG2	22:G:202:CLA:CAB	2.35	0.56
7:G:90:SER:HB2	19:9:34:ASN:O	2.05	0.56
15:8:170:PRO:CD	38:8:302:LUT:H24	2.35	0.56
25:8:317:LHG:H122	25:8:317:LHG:O7	2.05	0.56
38:5:303:LUT:H162	22:5:312:CLA:HMB3	1.87	0.56
18:6:184:ALA:HB1	38:6:303:LUT:H7	1.86	0.56
19:9:72:THR:HG21	22:9:313:CLA:O1D	2.05	0.56
38:9:301:LUT:C14	22:9:304:CLA:HBB1	2.35	0.56
22:2:301:CLA:C1C	22:2:301:CLA:H52	2.34	0.56
22:A:803:CLA:H8	9:J:16:LEU:HD23	1.87	0.56
22:A:831:CLA:HBA2	25:A:849:LHG:HC92	1.87	0.56
2:B:57:ILE:HG21	22:B:809:CLA:HMD2	1.86	0.56
11:L:140:PHE:HD1	11:L:144:PRO:HD2	1.71	0.56
12:1:170:LEU:CB	22:1:304:CLA:HMA1	2.35	0.56
12:Z:176:LYS:HD3	22:Z:309:CLA:O1D	2.05	0.56
22:Z:306:CLA:HBB1	22:Z:306:CLA:HMB1	1.88	0.56
22:Z:309:CLA:H142	22:Z:309:CLA:H61	1.86	0.56
22:7:308:CLA:OBD	22:7:315:CLA:HBA2	2.04	0.56
39:6:318:CHL:HAA2	39:6:318:CHL:HBD	1.88	0.56
22:6:321:CLA:HBB1	22:6:321:CLA:HMB1	1.85	0.56

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
22:9:307:CLA:HMD2	22:9:313:CLA:C1D	2.35	0.56
1:A:461:THR:HG21	22:B:812:CLA:HAB	1.87	0.56
22:A:812:CLA:HBA1	22:A:812:CLA:HBD	1.87	0.56
22:A:815:CLA:H121	24:3:306:BCR:H16C	1.88	0.56
22:A:818:CLA:HHD	22:A:835:CLA:H42	1.87	0.56
22:A:820:CLA:HHC	22:A:820:CLA:CBB	2.33	0.56
22:A:828:CLA:H43	22:A:828:CLA:HBA1	1.86	0.56
2:B:296:PHE:HB3	7:G:100:ILE:HG21	1.86	0.56
22:B:806:CLA:H13	24:B:849:BCR:H10C	1.88	0.56
8:I:74:TRP:HZ2	22:2:304:CLA:HMA1	1.71	0.56
39:Z:311:CHL:NC	25:Z:317:LHG:H102	2.21	0.56
38:3:303:LUT:H28	22:3:311:CLA:H2	1.87	0.56
22:3:320:CLA:CAD	22:3:322:CLA:HMA1	2.36	0.56
15:8:110:GLN:HB3	43:8:319:LPX:H2	1.87	0.56
22:8:313:CLA:HMA2	39:8:315:CHL:H122	1.88	0.56
16:4:185:LEU:HD23	39:4:819:CHL:HMD3	1.86	0.56
22:4:810:CLA:H12	39:4:816:CHL:C3D	2.35	0.56
17:5:67:LEU:HD12	22:5:310:CLA:HMA2	1.88	0.56
19:9:51:PHE:HB3	22:9:306:CLA:CAD	2.36	0.56
1:A:201:HIS:O	1:A:205:LEU:HB3	2.05	0.56
22:A:812:CLA:O1A	13:3:39:LEU:HB3	2.06	0.56
22:A:838:CLA:H91	22:A:838:CLA:H143	1.86	0.56
22:A:855:CLA:H111	22:B:840:CLA:H62	1.88	0.56
2:B:481:SER:O	2:B:484:SER:HB2	2.05	0.56
22:B:825:CLA:H141	22:B:832:CLA:HBC3	1.88	0.56
4:D:139:TYR:CE2	4:D:149:TYR:HD1	2.23	0.56
12:Z:159:PRO:HD2	38:Z:301:LUT:H23	1.88	0.56
31:3:301:DGD:HA21	31:3:301:DGD:HA61	1.86	0.56
39:8:312:CHL:HMC	26:8:318:NKP:HARA	1.86	0.56
18:6:123:PHE:CE1	22:6:302:CLA:HBD	2.41	0.56
18:6:245:PHE:CE2	22:6:301:CLA:H91	2.40	0.56
2:B:93:TRP:H	22:B:812:CLA:HED3	1.70	0.56
2:B:190:ALA:HB1	22:B:816:CLA:HAB	1.88	0.56
2:B:315:ARG:HB3	22:B:842:CLA:H43	1.88	0.56
2:B:349:VAL:HA	22:B:820:CLA:H11	1.88	0.56
2:B:463:TRP:CG	22:F:304:CLA:HMA2	2.41	0.56
2:B:470:LYS:HG2	2:B:473:TYR:CE2	2.41	0.56
22:B:814:CLA:HAB	22:B:822:CLA:H72	1.87	0.56
22:B:827:CLA:H172	22:B:833:CLA:HBC1	1.87	0.56
3:C:22:PRO:HB3	3:C:53:ARG:NH1	2.21	0.56
12:Z:100:TYR:O	12:Z:103:PRO:HD2	2.05	0.56

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
22:3:320:CLA:C4D	41:7:319:3PH:H392	2.35	0.56
14:7:164:GLU:HB2	14:7:167:TYR:HB2	1.87	0.56
14:7:186:TYR:O	14:7:190:LYS:HG3	2.06	0.56
22:7:322:CLA:HBA2	22:7:323:CLA:H102	1.88	0.56
1:A:677:PHE:HB2	22:A:802:CLA:HAA1	1.89	0.56
22:B:813:CLA:HBB1	22:B:813:CLA:CMB	2.24	0.56
7:G:54:LEU:N	7:G:55:PRO:HD2	2.21	0.56
12:1:134:MET:HG3	22:1:314:CLA:HMC3	1.88	0.56
12:Z:206:HIS:CB	22:Z:305:CLA:HAA2	2.33	0.56
15:8:128:PHE:CZ	24:8:304:BCR:H10C	2.41	0.56
38:8:302:LUT:H392	22:8:313:CLA:H41	1.89	0.56
22:4:807:CLA:HBA1	22:4:807:CLA:CBD	2.33	0.56
22:4:809:CLA:HHC	22:4:809:CLA:CBB	2.31	0.56
19:9:51:PHE:HB3	22:9:306:CLA:C3D	2.36	0.56
2:B:569:CYS:HG	29:B:802:SF4:FE1	1.22	0.55
2:B:708:LEU:HD21	31:B:852:DGD:HA21	1.87	0.55
22:B:805:CLA:HHC	22:B:805:CLA:CBB	2.32	0.55
12:1:75:VAL:HG22	12:1:175:ILE:HD11	1.88	0.55
12:Z:57:PRO:HG2	38:Z:302:LUT:H23	1.87	0.55
22:Z:309:CLA:C2B	25:Z:317:LHG:HC2	2.35	0.55
22:3:313:CLA:H2	22:3:319:CLA:CAD	2.36	0.55
22:5:315:CLA:C1A	25:5:324:LHG:H142	2.36	0.55
18:6:238:ASP:HA	18:6:243:LYS:CA	2.19	0.55
39:2:303:CHL:HAA1	39:2:303:CHL:HBD	1.88	0.55
1:A:93:PHE:CZ	22:A:807:CLA:HMD3	2.40	0.55
22:A:834:CLA:H172	22:L:202:CLA:HMB2	1.86	0.55
22:A:841:CLA:H111	22:B:801:CLA:H121	1.89	0.55
27:A:852:DGA:HBW1	31:3:301:DGD:HA31	1.87	0.55
2:B:427:SER:HB3	22:B:801:CLA:HBA1	1.89	0.55
22:B:810:CLA:HHB	22:B:811:CLA:HMB3	1.88	0.55
22:F:304:CLA:H2	22:F:304:CLA:H8	1.88	0.55
38:8:303:LUT:H28	22:8:308:CLA:H72	1.87	0.55
24:8:304:BCR:H372	39:8:315:CHL:H162	1.87	0.55
38:4:803:LUT:H193	39:4:816:CHL:CBB	2.36	0.55
38:4:803:LUT:H30	22:4:808:CLA:H72	1.89	0.55
22:4:810:CLA:O1A	22:4:818:CLA:HHC	2.06	0.55
17:5:106:PRO:O	17:5:109:LEU:HB2	2.06	0.55
17:5:127:LEU:HD13	22:6:314:CLA:H8	1.87	0.55
1:A:349:TRP:CD2	22:A:825:CLA:H192	2.42	0.55
1:A:535:VAL:HG22	1:A:539:HIS:NE2	2.22	0.55
22:A:814:CLA:H11	22:A:816:CLA:HHB	1.88	0.55

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
22:A:816:CLA:H43	22:A:816:CLA:HBA1	1.88	0.55
2:B:6:PHE:CE2	2:B:25:GLY:HA3	2.41	0.55
2:B:168:TRP:CE2	22:B:815:CLA:HAC2	2.41	0.55
2:B:291:MET:HB2	22:B:823:CLA:HAC2	1.88	0.55
11:L:131:ALA:HB1	24:L:204:BCR:C19	2.35	0.55
12:Z:102:ALA:HB3	12:Z:103:PRO:HD3	1.88	0.55
15:8:190:LYS:HD3	22:8:311:CLA:O1D	2.06	0.55
16:4:135:PHE:HA	39:4:813:CHL:HMA3	1.88	0.55
17:5:202:VAL:HG22	17:5:252:LEU:HD12	1.89	0.55
18:6:89:GLU:HB3	18:6:200:LEU:HD12	1.88	0.55
38:6:304:LUT:H28	22:6:310:CLA:C6	2.26	0.55
6:F:126:GLY:H	6:F:132:HIS:CE1	2.24	0.55
22:1:304:CLA:H122	22:1:305:CLA:CMB	2.36	0.55
16:4:68:PRO:HG2	16:4:71:LEU:HD12	1.87	0.55
1:A:438:ILE:HG13	1:A:556:PHE:HE2	1.71	0.55
22:A:830:CLA:H151	22:A:841:CLA:HMA2	1.88	0.55
2:B:221:GLN:NE2	2:B:225:PRO:HG3	2.22	0.55
22:B:817:CLA:H111	24:B:844:BCR:H383	1.88	0.55
22:B:825:CLA:HMD2	22:B:826:CLA:HAB	1.86	0.55
10:K:77:ILE:HG13	13:3:19:SER:HA	1.87	0.55
12:Z:206:HIS:CG	22:Z:305:CLA:HAA2	2.42	0.55
22:3:309:CLA:HMD2	22:3:314:CLA:C1D	2.35	0.55
15:8:196:MET:HE3	22:8:308:CLA:HMC3	1.87	0.55
22:A:818:CLA:O1A	22:A:827:CLA:HBB2	2.05	0.55
2:B:54:GLN:HE22	22:B:809:CLA:C1D	2.19	0.55
2:B:87:PRO:HB3	2:B:122:TYR:CD1	2.42	0.55
2:B:183:LEU:HD11	22:B:815:CLA:H43	1.87	0.55
2:B:722:TYR:HB2	22:B:805:CLA:HED3	1.89	0.55
8:I:80:VAL:HB	8:I:81:PRO:HD3	1.89	0.55
12:1:115:PHE:CZ	22:1:309:CLA:HMD2	2.41	0.55
13:3:68:MET:SD	22:3:308:CLA:HAB	2.46	0.55
38:3:302:LUT:C11	22:3:309:CLA:HMC2	2.37	0.55
22:3:312:CLA:NB	22:3:318:CLA:H122	2.22	0.55
14:7:189:TYR:CB	22:7:304:CLA:HMA1	2.36	0.55
22:4:810:CLA:HMA2	39:4:816:CHL:C3C	2.37	0.55
22:6:308:CLA:OBD	22:6:313:CLA:HBD	2.06	0.55
1:A:426:LEU:O	1:A:430:VAL:HG23	2.07	0.55
22:A:840:CLA:HMC3	22:F:301:CLA:C4D	2.37	0.55
2:B:408:VAL:O	2:B:412:MET:HG2	2.05	0.55
2:B:546:LYS:HG3	6:F:225:SER:OG	2.07	0.55
22:B:823:CLA:HHC	22:B:823:CLA:CBB	2.32	0.55

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
22:B:842:CLA:HBB2	22:1:307:CLA:H61	1.88	0.55
6:F:116:ALA:HB1	6:F:120:LYS:NZ	2.21	0.55
6:F:201:THR:HA	22:8:309:CLA:C15	2.36	0.55
8:I:63:ALA:HB2	20:2:206:GLU:OE2	2.07	0.55
22:Z:308:CLA:HMB1	22:Z:308:CLA:HBB1	1.89	0.55
13:3:186:LYS:HA	24:3:307:BCR:H282	1.87	0.55
22:3:320:CLA:C2D	41:7:319:3PH:H371	2.37	0.55
15:8:157:LYS:HB2	15:8:157:LYS:NZ	2.22	0.55
39:5:316:CHL:CBB	22:5:319:CLA:HBB2	2.37	0.55
19:9:123:GLU:HB3	19:9:126:ARG:HH21	1.71	0.55
20:2:211:VAL:HG13	39:2:303:CHL:H2	1.87	0.55
1:A:414:VAL:HG13	1:A:567:ALA:HA	1.88	0.55
1:A:575:CYS:HB3	1:A:584:CYS:HA	1.89	0.55
22:A:804:CLA:C4B	22:A:811:CLA:H122	2.37	0.55
22:A:806:CLA:HHC	22:A:806:CLA:CBB	2.37	0.55
22:B:816:CLA:H13	24:B:846:BCR:H19C	1.87	0.55
4:D:128:LEU:HA	4:D:132:PHE:HD2	1.72	0.55
13:3:97:ILE:HG22	13:3:99:PRO:HG2	1.89	0.55
22:3:311:CLA:C6	22:3:312:CLA:HMA1	2.36	0.55
14:7:186:TYR:CE2	14:7:190:LYS:HD2	2.42	0.55
15:8:77:MET:HB2	22:8:305:CLA:HMC3	1.89	0.55
18:6:220:VAL:HG22	22:6:322:CLA:HMB3	1.88	0.55
22:A:818:CLA:HMD2	22:A:835:CLA:H43	1.87	0.55
2:B:493:LEU:O	2:B:496:PRO:HD2	2.07	0.55
22:B:801:CLA:HMB1	22:B:801:CLA:CBB	2.37	0.55
22:B:822:CLA:HMB1	22:B:822:CLA:HBB1	1.89	0.55
22:F:304:CLA:H12	22:F:304:CLA:C4A	2.36	0.55
22:F:304:CLA:HBD	22:F:304:CLA:H151	1.88	0.55
7:G:99:LEU:HG	22:G:202:CLA:CGA	2.37	0.55
12:1:55:PHE:CD2	38:1:303:LUT:H383	2.41	0.55
22:Z:303:CLA:HMB1	22:Z:303:CLA:CBB	2.34	0.55
18:6:162:PHE:CZ	18:6:164:PRO:HB3	2.42	0.55
20:2:171:PHE:O	20:2:172:CYS:C	2.45	0.55
2:B:40:GLU:O	2:B:44:TYR:HD1	1.90	0.55
22:Z:304:CLA:HED2	22:Z:304:CLA:H2A	1.88	0.55
16:4:85:SER:HB2	16:4:88:VAL:CG2	2.37	0.55
16:4:157:PHE:CG	22:4:815:CLA:HMC3	2.42	0.55
16:4:237:LYS:HB2	16:4:242:ASN:HD21	1.72	0.55
39:4:819:CHL:HAA2	39:4:819:CHL:HBD	1.89	0.55
17:5:88:VAL:HG22	17:5:100:TRP:CD1	2.42	0.55
18:6:237:THR:HG21	39:6:315:CHL:HED2	1.89	0.55

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
18:6:252:GLY:HA3	22:6:301:CLA:HED1	1.89	0.55
1:A:348:SER:O	22:A:825:CLA:H191	2.07	0.54
22:A:822:CLA:C3D	24:A:856:BCR:H292	2.37	0.54
22:B:823:CLA:H62	22:B:824:CLA:H72	1.89	0.54
7:G:48:VAL:O	7:G:52:ALA:HB3	2.06	0.54
22:L:201:CLA:HHC	22:L:201:CLA:CBB	2.33	0.54
39:1:312:CHL:HBA1	31:1:319:DGD:HB31	1.89	0.54
40:1:318:SQD:H131	15:8:138:LEU:HG	1.89	0.54
22:3:311:CLA:H62	22:3:312:CLA:HMA1	1.88	0.54
22:8:307:CLA:HBA1	22:8:307:CLA:CBD	2.33	0.54
22:5:301:CLA:H61	22:5:320:CLA:HED1	1.88	0.54
19:9:126:ARG:HH22	22:9:313:CLA:CMA	2.17	0.54
19:9:206:GLN:NE2	19:9:206:GLN:H	2.05	0.54
1:A:21:PRO:HD3	1:A:183:LYS:O	2.08	0.54
1:A:672:PHE:CE2	1:A:676:HIS:HE1	2.26	0.54
27:A:852:DGA:HA52	14:7:243:PHE:HB3	1.89	0.54
6:F:196:THR:HB	22:F:305:CLA:HED2	1.89	0.54
13:3:114:PHE:CE2	22:3:316:CLA:H2	2.42	0.54
13:3:226:LEU:HA	13:3:229:PHE:CD2	2.42	0.54
14:7:68:MET:HB3	22:7:307:CLA:HMA1	1.90	0.54
14:7:103:TYR:HA	39:7:313:CHL:HMA3	1.90	0.54
25:7:318:LHG:H122	25:7:318:LHG:H322	1.89	0.54
22:5:307:CLA:H71	22:5:307:CLA:H41	1.87	0.54
1:A:13:VAL:HG12	22:A:812:CLA:H2A	1.88	0.54
22:A:805:CLA:HAA2	22:A:825:CLA:H41	1.90	0.54
2:B:23:TRP:HB3	31:B:852:DGD:HA41	1.89	0.54
12:Z:195:THR:CG2	12:Z:197:LYS:HE2	2.37	0.54
13:3:126:LEU:HD13	22:3:318:CLA:HMB3	1.88	0.54
22:7:322:CLA:HMB1	22:7:322:CLA:CBB	2.34	0.54
17:5:103:ALA:HB1	22:5:312:CLA:HED3	1.89	0.54
1:A:75:SER:OG	1:A:181:TYR:HB2	2.07	0.54
1:A:247:ARG:NH1	31:3:301:DGD:HE4	2.22	0.54
22:A:803:CLA:HMA1	22:A:808:CLA:H201	1.88	0.54
22:B:807:CLA:HBA1	22:B:807:CLA:CHA	2.38	0.54
22:B:827:CLA:HMA2	22:B:828:CLA:CED	2.38	0.54
12:Z:86:ALA:HB1	22:Z:303:CLA:H192	1.89	0.54
12:Z:185:CYS:HA	12:Z:188:PHE:CD2	2.42	0.54
22:Z:303:CLA:HBC2	22:Z:313:CLA:H12	1.88	0.54
14:7:66:ARG:HG2	14:7:142:PHE:CZ	2.43	0.54
15:8:220:HIS:HA	15:8:227:THR:OG1	2.08	0.54
17:5:58:LEU:HD22	22:5:315:CLA:H142	1.90	0.54

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
18:6:123:PHE:HB2	22:6:302:CLA:C1	2.37	0.54
18:6:139:ASP:O	18:6:148:LYS:HG3	2.06	0.54
22:6:308:CLA:HMD2	22:6:313:CLA:C4D	2.37	0.54
20:2:177:VAL:C	20:2:178:ARG:HG3	2.27	0.54
22:A:816:CLA:H42	27:A:852:DGA:HAF1	1.89	0.54
2:B:160:PRO:O	2:B:161:SER:HB3	2.07	0.54
2:B:298:ILE:HG22	7:G:57:HIS:CE1	2.43	0.54
2:B:549:PRO:HB2	3:C:62:PHE:CE2	2.42	0.54
38:1:302:LUT:C11	22:1:305:CLA:HMC2	2.38	0.54
38:4:803:LUT:H28	22:4:808:CLA:C5	2.38	0.54
22:6:309:CLA:C4B	22:6:309:CLA:H62	2.37	0.54
22:6:322:CLA:HMC1	41:6:324:3PH:H2A1	1.88	0.54
2:B:92:ILE:HD11	2:B:115:ASN:ND2	2.23	0.54
2:B:617:LEU:HD21	22:B:805:CLA:H202	1.90	0.54
2:B:695:LYS:HD2	8:I:100:GLU:HB3	1.90	0.54
2:B:697:LYS:HD3	3:C:81:TYR:HD2	1.71	0.54
22:B:827:CLA:HMA2	22:B:828:CLA:HED2	1.89	0.54
11:L:79:VAL:HB	11:L:84:ARG:HG3	1.90	0.54
12:1:39:SER:HB2	40:1:318:SQD:H62	1.88	0.54
13:3:80:LEU:HD11	24:3:305:BCR:H401	1.89	0.54
13:3:216:HIS:CG	22:3:310:CLA:HAA1	2.42	0.54
22:3:316:CLA:HHC	22:3:316:CLA:CBB	2.34	0.54
17:5:192:ILE:HD11	22:5:313:CLA:HHD	1.89	0.54
18:6:66:TRP:HE1	22:6:311:CLA:H11	1.72	0.54
18:6:254:TRP:CG	22:6:319:CLA:HED1	2.43	0.54
38:6:303:LUT:C30	22:6:307:CLA:H8	2.38	0.54
24:A:848:BCR:H393	22:B:801:CLA:H102	1.90	0.54
2:B:263:HIS:HE1	2:B:265:GLN:HB3	1.73	0.54
2:B:680:VAL:HG13	2:B:694:TRP:CZ2	2.43	0.54
6:F:165:TYR:O	6:F:168:ARG:HG2	2.07	0.54
7:G:56:LEU:HG	7:G:59:ARG:HH22	1.72	0.54
22:3:312:CLA:HMD2	22:3:318:CLA:C1D	2.37	0.54
22:8:306:CLA:HMD2	22:8:311:CLA:ND	2.23	0.54
22:4:808:CLA:HBB2	22:4:809:CLA:CHB	2.37	0.54
39:5:316:CHL:H122	39:5:316:CHL:C9	2.36	0.54
18:6:238:ASP:HB2	18:6:243:LYS:HG2	1.89	0.54
1:A:584:CYS:HG	29:B:802:SF4:FE2	1.25	0.54
22:A:812:CLA:HBA1	22:A:812:CLA:CBD	2.37	0.54
22:A:815:CLA:H92	13:3:203:VAL:HG21	1.90	0.54
2:B:235:ALA:HA	2:B:257:THR:HG22	1.90	0.54
2:B:515:PRO:HG2	6:F:132:HIS:CD2	2.42	0.54

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
22:B:819:CLA:O1A	22:B:828:CLA:HBB2	2.07	0.54
22:B:834:CLA:HMB2	22:F:302:CLA:CBB	2.38	0.54
7:G:46:LEU:HB3	22:G:202:CLA:HMB1	1.89	0.54
22:1:305:CLA:HHC	22:1:305:CLA:CBB	2.32	0.54
38:7:302:LUT:C32	22:7:307:CLA:HAB	2.27	0.54
2:B:438:TYR:CD1	2:B:617:LEU:HG	2.43	0.54
22:Z:307:CLA:HMD2	22:Z:314:CLA:C1D	2.38	0.54
39:7:313:CHL:HAA1	39:7:313:CHL:HBD	1.90	0.54
22:8:310:CLA:HBA2	43:8:319:LPX:H3	1.90	0.54
39:8:312:CHL:HMB1	39:8:312:CHL:CBB	2.37	0.54
22:5:311:CLA:HMD1	22:6:322:CLA:H92	1.90	0.54
22:6:314:CLA:HMC2	25:6:323:LHG:HC61	1.88	0.54
1:A:578:PRO:HG3	3:C:52:LYS:HD2	1.89	0.54
1:A:682:SER:HB2	1:A:727:GLY:O	2.08	0.54
22:A:804:CLA:C3B	22:A:811:CLA:H122	2.38	0.54
22:A:814:CLA:H43	22:A:816:CLA:H41	1.90	0.54
22:B:823:CLA:C1B	22:B:824:CLA:H8	2.38	0.54
4:D:77:LYS:HG3	4:D:81:GLU:HB3	1.90	0.54
7:G:33:GLU:HB2	7:G:35:TYR:CD2	2.43	0.54
22:1:308:CLA:HHC	22:1:308:CLA:CBB	2.37	0.54
12:Z:157:PHE:HA	22:Z:313:CLA:C7	2.38	0.54
22:8:307:CLA:H161	22:8:311:CLA:HMC1	1.90	0.54
39:8:312:CHL:CMC	26:8:318:NKP:HARA	2.38	0.54
22:4:810:CLA:H43	39:4:816:CHL:HBD	1.90	0.54
38:5:302:LUT:H373	22:5:307:CLA:H42	1.89	0.54
25:5:324:LHG:H102	25:5:324:LHG:H281	1.89	0.54
18:6:251:GLN:HB2	22:6:301:CLA:HBA1	1.90	0.54
19:9:137:GLY:HA2	19:9:144:PHE:HB2	1.90	0.54
1:A:49:ILE:H	1:A:49:ILE:HD12	1.74	0.53
1:A:646:LEU:HD12	1:A:650:LEU:HD12	1.90	0.53
1:A:707:LEU:HD23	6:F:216:THR:HG22	1.90	0.53
22:A:805:CLA:H101	24:A:845:BCR:HC8	1.89	0.53
6:F:123:LEU:HD11	9:J:38:VAL:HG11	1.90	0.53
22:G:202:CLA:HMB1	22:G:202:CLA:CBB	2.38	0.53
12:Z:218:ASN:HD22	22:Z:305:CLA:HMD1	1.72	0.53
38:4:802:LUT:C14	22:4:806:CLA:HAB	2.37	0.53
17:5:128:LEU:HB3	24:5:304:BCR:H15C	1.89	0.53
18:6:219:ALA:HB2	18:6:229:PRO:HD3	1.90	0.53
19:9:29:PRO:HB3	22:9:311:CLA:HED2	1.90	0.53
1:A:80:GLN:HG2	22:A:805:CLA:CMA	2.28	0.53
1:A:80:GLN:O	1:A:83:VAL:HG22	2.08	0.53

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
22:A:805:CLA:H143	24:A:844:BCR:C5	2.38	0.53
22:A:812:CLA:H172	22:A:812:CLA:H143	1.89	0.53
2:B:200:ILE:HB	2:B:201:PRO:HD3	1.90	0.53
3:C:62:PHE:HE1	3:C:66:ARG:HH21	1.55	0.53
6:F:190:PRO:O	6:F:194:ARG:HG3	2.08	0.53
7:G:97:PHE:CD2	7:G:102:THR:HG22	2.43	0.53
8:I:67:TYR:CZ	8:I:69:PHE:HB2	2.43	0.53
11:L:140:PHE:HA	11:L:144:PRO:CD	2.38	0.53
13:3:82:LYS:HZ3	13:3:211:GLU:HG2	1.70	0.53
38:7:301:LUT:H30	22:7:304:CLA:H72	1.89	0.53
15:8:202:PHE:CZ	38:8:302:LUT:H8	2.42	0.53
17:5:33:LYS:HB3	22:5:315:CLA:CBC	2.37	0.53
22:5:301:CLA:HHC	22:5:301:CLA:HBB1	1.89	0.53
19:9:63:LEU:HD12	22:9:306:CLA:HMA2	1.88	0.53
22:A:812:CLA:HHC	22:A:812:CLA:CBB	2.36	0.53
22:A:821:CLA:C2C	22:A:827:CLA:H18	2.38	0.53
2:B:442:ASP:OD1	2:B:616:TYR:HB2	2.08	0.53
22:B:820:CLA:HBB1	22:B:820:CLA:CMB	2.25	0.53
13:3:78:GLU:HB3	13:3:82:LYS:HE3	1.91	0.53
24:3:305:BCR:H21C	22:3:313:CLA:CAC	2.36	0.53
22:3:312:CLA:HHC	22:3:312:CLA:CBB	2.38	0.53
22:7:307:CLA:H92	22:7:308:CLA:H72	1.90	0.53
16:4:153:MET:HB2	39:4:816:CHL:HMA3	1.89	0.53
22:A:837:CLA:HHC	22:A:837:CLA:CBB	2.31	0.53
2:B:241:ALA:HA	2:B:264:PRO:HG3	1.90	0.53
2:B:590:TRP:CD1	22:B:805:CLA:H152	2.42	0.53
6:F:163:ILE:HG12	22:F:305:CLA:C4D	2.39	0.53
11:L:160:PRO:O	11:L:166:GLY:HA3	2.07	0.53
39:Z:312:CHL:CAB	22:Z:315:CLA:HBB2	2.38	0.53
15:8:136:ASP:O	15:8:140:PRO:HA	2.09	0.53
16:4:106:TRP:CH2	24:4:804:BCR:H372	2.44	0.53
22:4:806:CLA:C4D	22:4:811:CLA:H61	2.38	0.53
22:5:326:CLA:CMD	22:6:309:CLA:H2	2.37	0.53
18:6:221:VAL:HB	22:6:322:CLA:C1C	2.39	0.53
22:A:806:CLA:CBB	24:A:845:BCR:HC31	2.39	0.53
2:B:48:PHE:HB2	2:B:169:PHE:HE1	1.74	0.53
2:B:355:SER:HB3	22:B:828:CLA:HAC2	1.89	0.53
22:B:827:CLA:C3B	24:B:848:BCR:H19C	2.38	0.53
22:B:828:CLA:C11	24:B:847:BCR:H373	2.38	0.53
39:Z:312:CHL:HMB1	39:Z:312:CHL:CBB	2.36	0.53
24:3:305:BCR:H19C	22:3:313:CLA:HMC1	1.90	0.53

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
22:3:312:CLA:C1D	22:3:318:CLA:H93	2.38	0.53
22:3:313:CLA:H202	22:5:320:CLA:CHB	2.37	0.53
24:7:303:BCR:H333	22:7:322:CLA:H8	1.90	0.53
22:4:807:CLA:H71	25:4:820:LHG:H171	1.90	0.53
17:5:140:TYR:OH	22:5:323:CLA:HBA2	2.09	0.53
24:6:305:BCR:H291	22:6:312:CLA:H2	1.91	0.53
22:A:819:CLA:HHD	24:A:845:BCR:H352	1.90	0.53
2:B:244:VAL:HB	2:B:247:THR:OG1	2.09	0.53
2:B:320:HIS:O	2:B:323:LEU:HB2	2.08	0.53
2:B:379:ILE:O	2:B:383:ILE:HG13	2.09	0.53
22:B:811:CLA:H142	22:B:811:CLA:HMB2	1.89	0.53
10:K:61:THR:HA	10:K:67:LEU:HA	1.91	0.53
22:Z:304:CLA:HHC	22:Z:304:CLA:CBB	2.37	0.53
22:Z:316:CLA:HBD	22:Z:316:CLA:H72	1.91	0.53
15:8:110:GLN:HG2	43:8:319:LPX:H1	1.89	0.53
22:8:311:CLA:HMA1	33:4:801:LMT:H5'	1.89	0.53
17:5:53:TYR:HB2	22:5:310:CLA:HMD1	1.90	0.53
17:5:131:TRP:HH2	24:5:304:BCR:H323	1.74	0.53
38:5:302:LUT:H30	22:5:307:CLA:H51	1.91	0.53
1:A:44:ASN:HB3	5:E:78:GLN:NE2	2.23	0.53
1:A:95:GLY:HA2	1:A:99:SER:HB3	1.89	0.53
2:B:312:PRO:HB2	25:B:850:LHG:HC32	1.90	0.53
38:1:303:LUT:C28	22:1:307:CLA:H111	2.39	0.53
22:1:307:CLA:H102	22:1:308:CLA:HMB2	1.90	0.53
22:Z:304:CLA:ND	22:Z:309:CLA:H62	2.23	0.53
22:8:308:CLA:H12	22:8:308:CLA:CHB	2.39	0.53
22:8:309:CLA:HMD2	22:8:314:CLA:C4C	2.38	0.53
22:2:302:CLA:HBA1	22:2:302:CLA:CHA	2.38	0.53
1:A:190:TRP:CD1	22:A:812:CLA:HED2	2.43	0.53
22:A:822:CLA:HMB1	22:A:822:CLA:CBB	2.36	0.53
2:B:588:ILE:O	2:B:591:VAL:HG22	2.09	0.53
2:B:655:HIS:HB2	22:B:806:CLA:HBA1	1.89	0.53
22:B:825:CLA:H111	24:B:847:BCR:C15	2.38	0.53
22:B:830:CLA:HHC	22:B:830:CLA:CBB	2.35	0.53
6:F:107:THR:HA	6:F:110:ARG:HE	1.73	0.53
24:G:203:BCR:H341	24:G:203:BCR:C12	2.39	0.53
12:1:194:ALA:HB1	12:1:218:ASN:ND2	2.23	0.53
22:Z:313:CLA:H172	22:Z:313:CLA:CHC	2.39	0.53
38:7:302:LUT:H26	22:7:307:CLA:H61	1.89	0.53
22:4:806:CLA:HMD2	22:4:811:CLA:ND	2.24	0.53
22:6:302:CLA:C3D	24:6:305:BCR:HC41	2.38	0.53

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
19:9:111:LEU:HD22	39:9:312:CHL:HAB	1.89	0.53
22:A:809:CLA:O1A	22:A:809:CLA:HBD	2.08	0.53
22:A:812:CLA:H92	22:3:311:CLA:H112	1.91	0.53
22:A:812:CLA:H142	13:3:39:LEU:HD11	1.89	0.53
22:A:815:CLA:H2	13:3:203:VAL:CG1	2.38	0.53
22:A:816:CLA:HMB1	22:A:816:CLA:CBB	2.31	0.53
22:A:826:CLA:HMB1	22:A:826:CLA:CBB	2.34	0.53
27:A:852:DGA:HB91	22:3:316:CLA:HMC2	1.90	0.53
24:A:856:BCR:C13	24:K:206:BCR:HC8	2.39	0.53
22:B:819:CLA:C9	22:B:828:CLA:HMB1	2.38	0.53
10:K:78:SER:HB2	10:K:85:THR:HG22	1.91	0.53
22:Z:303:CLA:H62	22:Z:304:CLA:HMA1	1.91	0.53
22:Z:316:CLA:HHC	22:Z:316:CLA:CBB	2.32	0.53
38:3:303:LUT:H183	22:3:313:CLA:C4B	2.38	0.53
14:7:76:ALA:HB1	14:7:197:GLY:HA3	1.91	0.53
22:7:307:CLA:H71	22:7:308:CLA:HMA1	1.91	0.53
22:7:323:CLA:CAB	25:8:317:LHG:H151	2.38	0.53
15:8:236:ILE:HD11	22:8:316:CLA:HBC3	1.89	0.53
22:8:307:CLA:CGA	22:8:307:CLA:C1A	2.86	0.53
24:5:304:BCR:H271	22:5:312:CLA:CHB	2.39	0.53
22:5:311:CLA:H52	22:5:323:CLA:HAB	1.91	0.53
18:6:66:TRP:CZ2	22:6:317:CLA:HAA2	2.44	0.53
38:6:303:LUT:H26	22:6:307:CLA:O1A	2.08	0.53
1:A:204:GLY:HA3	22:A:813:CLA:HBB1	1.90	0.53
1:A:488:GLN:HA	1:A:510:TRP:HE3	1.74	0.53
1:A:740:TRP:HD1	22:A:828:CLA:HMB2	1.74	0.53
22:A:821:CLA:H43	22:A:835:CLA:H143	1.91	0.53
22:B:836:CLA:HMB1	22:B:836:CLA:CBB	2.34	0.53
22:F:305:CLA:HMB1	22:F:305:CLA:CBB	2.39	0.53
12:Z:144:GLU:HB2	22:Z:313:CLA:HAC1	1.90	0.53
13:3:227:THR:O	13:3:231:LYS:HG2	2.09	0.53
38:3:302:LUT:H11	22:3:309:CLA:HMC2	1.90	0.53
16:4:133:PRO:HG2	16:4:136:GLU:CB	2.38	0.53
17:5:185:LYS:HZ3	22:5:313:CLA:HBD	1.73	0.53
17:5:227:TRP:NE1	22:5:314:CLA:HED2	2.23	0.53
18:6:141:ASP:HB3	18:6:144:PHE:O	2.08	0.53
22:6:308:CLA:HMB1	22:6:308:CLA:CBB	2.36	0.53
19:9:81:VAL:CG1	38:9:301:LUT:H12	2.36	0.53
22:9:305:CLA:HBA1	22:9:305:CLA:HBD	1.90	0.53
1:A:264:PHE:HA	1:A:269:TRP:CD1	2.44	0.52
1:A:681:PHE:HZ	22:A:841:CLA:HBC2	1.74	0.52

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
22:A:802:CLA:H12	2:B:617:LEU:HD13	1.90	0.52
22:A:816:CLA:NC	27:A:852:DGA:HB52	2.24	0.52
22:B:810:CLA:H62	22:B:810:CLA:C1C	2.38	0.52
22:B:828:CLA:HMA3	24:B:848:BCR:C6	2.39	0.52
4:D:156:VAL:HG12	4:D:160:LYS:HB2	1.90	0.52
12:1:203:LEU:HB2	38:1:302:LUT:H22	1.91	0.52
22:Z:304:CLA:CAD	22:Z:309:CLA:HBD	2.39	0.52
22:Z:307:CLA:HMD2	22:Z:314:CLA:ND	2.24	0.52
14:7:119:SER:HB3	15:8:226:LYS:HG3	1.91	0.52
16:4:185:LEU:HD11	16:4:197:PHE:CD2	2.44	0.52
22:4:809:CLA:HMD2	22:4:815:CLA:C1D	2.38	0.52
18:6:147:ASN:HD21	39:6:320:CHL:HAA1	1.72	0.52
1:A:45:THR:HG23	5:E:78:GLN:NE2	2.21	0.52
1:A:247:ARG:HB2	31:3:301:DGD:O3D	2.09	0.52
22:A:855:CLA:H202	24:L:204:BCR:H391	1.91	0.52
2:B:394:PHE:HA	2:B:398:ASP:OD1	2.09	0.52
12:Z:88:CYS:HA	12:Z:99:TRP:HB3	1.90	0.52
12:Z:144:GLU:HG3	22:Z:313:CLA:HBC1	1.91	0.52
39:Z:312:CHL:HHC	22:Z:315:CLA:HAB	1.90	0.52
13:3:169:LEU:HB2	22:3:308:CLA:O1A	2.09	0.52
14:7:132:HIS:CD2	22:7:315:CLA:HMB1	2.44	0.52
18:6:141:ASP:CB	18:6:147:ASN:HB2	2.40	0.52
20:2:180:LYS:HE2	22:2:301:CLA:HED3	1.91	0.52
22:1:307:CLA:H102	22:1:308:CLA:CMB	2.40	0.52
39:1:312:CHL:HAA2	31:1:319:DGD:O2G	2.09	0.52
17:5:33:LYS:HB3	22:5:315:CLA:HBC3	1.91	0.52
22:5:301:CLA:H62	22:5:301:CLA:NB	2.25	0.52
22:9:313:CLA:HHC	22:9:313:CLA:CBB	2.40	0.52
20:2:197:ASN:CA	22:2:301:CLA:HBA2	2.35	0.52
1:A:366:ILE:HG22	1:A:370:HIS:CE1	2.45	0.52
22:A:830:CLA:HMB1	22:A:830:CLA:CBB	2.33	0.52
6:F:205:GLN:HG2	15:8:55:LEU:HD23	1.91	0.52
22:F:302:CLA:HMC3	35:J:102:T7X:C19	2.39	0.52
7:G:38:VAL:HG11	22:G:201:CLA:C3D	2.40	0.52
13:3:199:GLY:O	13:3:203:VAL:HG23	2.09	0.52
24:3:304:BCR:H402	22:3:313:CLA:C1B	2.39	0.52
17:5:171:PRO:HD2	38:5:302:LUT:C23	2.38	0.52
1:A:32:PRO:HB2	1:A:48:TRP:HH2	1.74	0.52
1:A:114:SER:HG	1:A:135:PHE:HE2	1.56	0.52
22:A:826:CLA:HAA2	22:A:827:CLA:OBD	2.10	0.52
26:A:851:NKP:HAGA	26:A:851:NKP:OAE	2.09	0.52

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:B:628:ASN:HB2	2:B:725:PHE:CZ	2.44	0.52
2:B:708:LEU:HD21	31:B:852:DGD:HA51	1.91	0.52
22:B:818:CLA:NC	22:B:818:CLA:H51	2.24	0.52
3:C:12:ILE:HG12	29:C:102:SF4:S2	2.50	0.52
8:I:92:MET:HB3	24:L:204:BCR:H12C	1.91	0.52
22:L:203:CLA:HMB1	22:L:203:CLA:CBB	2.37	0.52
12:1:94:LEU:HD12	22:1:309:CLA:CMD	2.38	0.52
38:1:303:LUT:H28	22:1:307:CLA:H8	1.90	0.52
22:1:313:CLA:HBB1	22:1:313:CLA:HMB1	1.91	0.52
22:Z:303:CLA:HMD2	22:Z:313:CLA:O2A	2.09	0.52
39:9:312:CHL:H2A	39:9:312:CHL:O2A	2.09	0.52
1:A:278:PHE:HZ	22:A:818:CLA:H43	1.75	0.52
22:A:823:CLA:NB	37:K:201:DAO:H91	2.24	0.52
22:B:828:CLA:HHC	22:B:828:CLA:CBB	2.38	0.52
10:K:107:LEU:HD21	22:K:203:CLA:HBB1	1.91	0.52
24:3:304:BCR:C5	22:3:322:CLA:HMD2	2.40	0.52
24:3:307:BCR:H402	25:3:321:LHG:O6	2.09	0.52
18:6:107:PRO:HD2	39:6:318:CHL:CMD	2.40	0.52
18:6:157:PRO:HG2	22:6:307:CLA:HMD1	1.90	0.52
18:6:186:ILE:HD11	22:6:311:CLA:HAB	1.91	0.52
1:A:299:LEU:HG	22:A:815:CLA:HMC1	1.91	0.52
1:A:349:TRP:HE3	22:A:805:CLA:HMD2	1.75	0.52
22:A:804:CLA:HED1	22:A:811:CLA:HBD	1.90	0.52
2:B:390:HIS:HA	2:B:393:ILE:HD12	1.91	0.52
22:B:825:CLA:HAB	22:B:832:CLA:HMD2	1.92	0.52
6:F:116:ALA:HB1	6:F:120:LYS:HZ2	1.73	0.52
9:J:1:MET:H2	34:J:101:LMG:C2	2.21	0.52
22:K:202:CLA:HMB1	22:K:202:CLA:CBB	2.36	0.52
22:Z:304:CLA:OBD	22:Z:309:CLA:HBD	2.08	0.52
13:3:38:PRO:HD2	38:3:303:LUT:C23	2.40	0.52
24:3:305:BCR:H24C	22:3:313:CLA:CAC	2.39	0.52
24:3:305:BCR:H342	39:3:317:CHL:HHB	1.92	0.52
22:7:309:CLA:HMB1	22:7:309:CLA:CBB	2.36	0.52
38:4:802:LUT:H382	22:4:805:CLA:C4D	2.39	0.52
17:5:95:ARG:NH1	17:5:110:PRO:HB3	2.25	0.52
17:5:138:GLN:NE2	18:6:35:GLY:HA3	2.25	0.52
22:5:311:CLA:OBD	22:5:318:CLA:HAA1	2.09	0.52
18:6:34:PRO:HD2	22:6:314:CLA:O1D	2.09	0.52
18:6:108:PHE:CD2	18:6:113:LEU:HD21	2.44	0.52
22:6:319:CLA:HHC	22:6:319:CLA:HBB1	1.91	0.52
1:A:197:MET:CG	22:A:813:CLA:HBC2	2.38	0.52

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:B:477:LEU:H	2:B:480:SER:HB2	1.75	0.52
2:B:688:LEU:HB2	24:B:853:BCR:HC31	1.92	0.52
22:B:831:CLA:HMB1	22:B:831:CLA:CBB	2.35	0.52
12:Z:87:GLY:HA3	38:Z:302:LUT:H182	1.91	0.52
12:Z:203:LEU:HD22	38:Z:301:LUT:H172	1.92	0.52
14:7:55:PRO:HD2	38:7:302:LUT:H23	1.90	0.52
24:7:303:BCR:H292	22:7:309:CLA:H71	1.90	0.52
38:8:302:LUT:H373	22:8:305:CLA:H42	1.91	0.52
17:5:106:PRO:HG3	17:5:243:GLN:HG3	1.91	0.52
22:A:819:CLA:HMB1	22:A:819:CLA:CBB	2.40	0.52
22:A:829:CLA:C1C	24:A:845:BCR:HC42	2.40	0.52
22:A:832:CLA:H121	22:A:834:CLA:H201	1.91	0.52
22:A:833:CLA:H202	24:L:204:BCR:C10	2.40	0.52
2:B:552:LYS:HD2	4:D:181:PRO:HD3	1.91	0.52
22:B:814:CLA:H3A	19:9:54:LEU:HD22	1.92	0.52
22:B:827:CLA:H2	22:B:839:CLA:HBA2	1.91	0.52
22:B:828:CLA:HBA2	24:B:848:BCR:C10	2.40	0.52
22:B:833:CLA:H112	22:F:302:CLA:H151	1.92	0.52
22:B:835:CLA:HMB1	22:B:835:CLA:CBB	2.35	0.52
11:L:92:HIS:HA	11:L:95:PHE:HE2	1.75	0.52
12:1:60:LEU:CD1	22:1:307:CLA:H42	2.40	0.52
12:Z:105:TRP:CH2	12:Z:112:PRO:HG3	2.45	0.52
13:3:65:ARG:HA	13:3:68:MET:HE3	1.92	0.52
13:3:180:LEU:HB3	22:3:308:CLA:CMA	2.39	0.52
24:3:307:BCR:HC7	22:3:315:CLA:H3A	1.90	0.52
22:3:313:CLA:HAA1	22:5:301:CLA:HMB3	1.90	0.52
14:7:82:GLY:O	14:7:86:VAL:HG23	2.10	0.52
22:8:307:CLA:HMB1	22:8:307:CLA:CBB	2.37	0.52
22:8:313:CLA:CMA	39:8:315:CHL:H52	2.40	0.52
16:4:86:LEU:CD1	38:4:803:LUT:H221	2.40	0.52
16:4:139:LYS:HG3	39:4:813:CHL:O2D	2.10	0.52
17:5:131:TRP:CH2	24:5:304:BCR:H323	2.44	0.52
17:5:243:GLN:HB3	24:5:305:BCR:H312	1.91	0.52
22:5:318:CLA:HMB1	22:5:318:CLA:CBB	2.38	0.52
18:6:88:GLN:NE2	38:6:304:LUT:H42	2.24	0.52
19:9:82:VAL:HG12	22:9:308:CLA:HBC2	1.91	0.52
19:9:101:ALA:N	39:9:312:CHL:HAA1	2.25	0.52
22:A:810:CLA:HMB1	22:A:810:CLA:CBB	2.37	0.52
2:B:127:THR:HA	2:B:245:PHE:CZ	2.45	0.52
2:B:394:PHE:HZ	2:B:409:LEU:HD21	1.75	0.52
12:1:194:ALA:HB2	22:1:316:CLA:HED3	1.91	0.52

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
12:Z:53:TYR:CE1	39:Z:311:CHL:HAC2	2.45	0.52
14:7:175:PRO:HD2	38:7:301:LUT:C23	2.40	0.52
39:5:317:CHL:HAA1	39:5:317:CHL:HBD	1.90	0.52
22:6:312:CLA:H193	22:6:312:CLA:H152	1.92	0.52
22:A:824:CLA:HHC	22:A:824:CLA:CBB	2.41	0.51
2:B:327:VAL:HG21	22:B:826:CLA:HMC2	1.92	0.51
2:B:394:PHE:CZ	2:B:409:LEU:HD21	2.45	0.51
22:B:824:CLA:C1C	33:1:301:LMT:H101	2.39	0.51
22:1:313:CLA:CHC	22:1:313:CLA:H172	2.41	0.51
22:3:313:CLA:H11	22:5:301:CLA:HMB2	1.91	0.51
14:7:149:GLY:HA3	14:7:161:LYS:HE3	1.93	0.51
1:A:360:PHE:HD1	22:A:829:CLA:H192	1.76	0.51
2:B:362:LEU:HD22	2:B:369:GLN:HE21	1.75	0.51
22:B:835:CLA:HMB2	24:B:848:BCR:H332	1.91	0.51
6:F:78:LYS:HZ2	6:F:81:LYS:HD3	1.75	0.51
12:Z:56:ASP:HA	38:Z:302:LUT:H24	1.91	0.51
13:3:129:TRP:HZ3	41:7:319:3PH:H322	1.74	0.51
24:7:303:BCR:H383	22:7:316:CLA:NA	2.25	0.51
22:7:312:CLA:H72	22:7:312:CLA:C4B	2.40	0.51
22:7:323:CLA:H92	25:8:317:LHG:H321	1.93	0.51
16:4:108:MET:SD	22:4:805:CLA:HAB	2.50	0.51
16:4:210:LEU:HB3	22:4:805:CLA:CMA	2.39	0.51
16:4:254:THR:HA	22:4:807:CLA:HBA2	1.91	0.51
22:6:319:CLA:C3D	25:6:323:LHG:H372	2.40	0.51
19:9:65:TRP:CZ2	22:9:313:CLA:HAA1	2.45	0.51
1:A:460:ASP:OD1	1:A:641:THR:HB	2.10	0.51
22:A:810:CLA:HBA2	22:A:810:CLA:H42	1.92	0.51
2:B:463:TRP:NE1	2:B:477:LEU:HG	2.24	0.51
22:B:810:CLA:H2	22:B:810:CLA:C4D	2.41	0.51
22:B:816:CLA:HMB1	22:B:816:CLA:CBB	2.40	0.51
22:B:824:CLA:HBC2	22:B:825:CLA:HBA1	1.91	0.51
22:G:202:CLA:C4C	24:G:203:BCR:H282	2.41	0.51
10:K:58:LYS:O	10:K:59:ARG:HB2	2.09	0.51
22:1:310:CLA:C1C	25:1:317:LHG:HC41	2.41	0.51
13:3:139:SER:HB3	13:3:151:GLY:HA2	1.92	0.51
16:4:163:ARG:NH2	16:4:178:PRO:HG3	2.26	0.51
22:4:810:CLA:HMB1	22:4:810:CLA:CBB	2.40	0.51
18:6:132:PHE:HD1	18:6:154:VAL:HG11	1.75	0.51
18:6:147:ASN:HB3	39:6:320:CHL:C3D	2.40	0.51
1:A:267:LEU:HD12	10:K:34:THR:HG23	1.92	0.51
1:A:407:ALA:O	1:A:411:ILE:HG13	2.10	0.51

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
22:A:804:CLA:HMB1	22:A:804:CLA:CBB	2.37	0.51
22:A:842:CLA:H42	22:A:842:CLA:HED2	1.92	0.51
2:B:530:THR:HG21	2:B:583:TRP:CE2	2.46	0.51
2:B:694:TRP:HA	22:B:840:CLA:O1D	2.10	0.51
22:B:811:CLA:HBA1	22:B:811:CLA:CBD	2.40	0.51
22:B:814:CLA:H112	22:B:814:CLA:C3B	2.41	0.51
22:B:840:CLA:H111	22:B:841:CLA:H142	1.92	0.51
10:K:59:ARG:N	10:K:69:ASP:HA	2.25	0.51
22:K:205:CLA:CMD	24:K:206:BCR:H21C	2.39	0.51
38:1:303:LUT:C28	22:1:307:CLA:H8	2.40	0.51
22:7:312:CLA:H143	22:7:312:CLA:HMC2	1.93	0.51
15:8:118:LEU:HD21	39:8:315:CHL:C1D	2.40	0.51
22:8:311:CLA:HHC	22:8:311:CLA:CBB	2.40	0.51
22:4:805:CLA:HMD2	39:4:814:CHL:O1A	2.09	0.51
17:5:227:TRP:HH2	22:5:301:CLA:HBC2	1.75	0.51
22:5:326:CLA:H51	22:6:314:CLA:H61	1.93	0.51
18:6:79:MET:SD	22:6:307:CLA:HAB	2.50	0.51
22:6:309:CLA:HMB1	22:6:309:CLA:CBB	2.38	0.51
1:A:15:ILE:HG23	22:A:810:CLA:HMA2	1.92	0.51
1:A:413:MET:HG3	1:A:558:ARG:HG3	1.93	0.51
22:A:820:CLA:H143	10:K:90:LEU:HD21	1.91	0.51
22:A:825:CLA:HMB1	22:A:825:CLA:CBB	2.40	0.51
22:B:812:CLA:HMB1	22:B:812:CLA:CBB	2.35	0.51
10:K:32:SER:O	10:K:36:LEU:HG	2.10	0.51
17:5:82:MET:SD	22:5:307:CLA:HAB	2.51	0.51
18:6:235:GLN:HG2	18:6:245:PHE:HE1	1.76	0.51
1:A:75:SER:HA	1:A:78:PHE:HD2	1.76	0.51
1:A:104:TRP:CH2	1:A:149:ARG:HD2	2.45	0.51
22:A:816:CLA:CMB	24:A:844:BCR:H341	2.40	0.51
22:A:824:CLA:H11	24:A:846:BCR:H363	1.92	0.51
2:B:355:SER:CB	22:B:828:CLA:HAC2	2.41	0.51
22:B:824:CLA:HMB1	22:B:824:CLA:CBB	2.37	0.51
22:B:827:CLA:HMB1	22:B:827:CLA:CBB	2.36	0.51
22:B:833:CLA:HMB1	22:B:833:CLA:CBB	2.36	0.51
22:B:842:CLA:HAB	12:1:70:PHE:CZ	2.45	0.51
3:C:29:VAL:HG12	4:D:165:ARG:HB3	1.92	0.51
6:F:162:TRP:CD1	22:F:305:CLA:HBD	2.46	0.51
11:L:123:GLY:HA2	22:L:201:CLA:H161	1.91	0.51
12:Z:100:TYR:O	39:Z:312:CHL:HED1	2.10	0.51
14:7:196:ASN:HD22	14:7:196:ASN:N	2.08	0.51
22:7:310:CLA:HBA2	22:7:310:CLA:CHA	2.40	0.51

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
38:8:303:LUT:H163	39:8:315:CHL:HMC	1.92	0.51
39:8:312:CHL:HBD	39:8:312:CHL:HAA1	1.93	0.51
22:4:811:CLA:HHC	22:4:811:CLA:CBB	2.40	0.51
25:4:820:LHG:H312	22:6:321:CLA:HED1	1.92	0.51
22:5:322:CLA:HMB1	22:5:322:CLA:CBB	2.34	0.51
18:6:120:ALA:HB1	24:6:305:BCR:C15	2.40	0.51
24:6:306:BCR:H352	39:6:315:CHL:HAC2	1.91	0.51
1:A:349:TRP:HZ3	22:A:825:CLA:H91	1.76	0.51
22:A:815:CLA:C4	24:3:306:BCR:HC8	2.40	0.51
22:A:818:CLA:HMB1	22:A:818:CLA:HBB1	1.91	0.51
2:B:93:TRP:CZ2	8:I:77:SER:HA	2.46	0.51
2:B:463:TRP:HD1	22:F:304:CLA:H193	1.75	0.51
22:B:842:CLA:HBB2	22:1:307:CLA:H41	1.93	0.51
22:F:302:CLA:C1B	24:F:303:BCR:H10C	2.41	0.51
12:Z:79:ARG:HA	12:Z:82:MET:HE2	1.92	0.51
15:8:149:PHE:O	15:8:150:LEU:C	2.47	0.51
22:8:306:CLA:HHC	22:8:306:CLA:CBB	2.32	0.51
16:4:234:THR:HG22	22:4:817:CLA:O1D	2.10	0.51
17:5:165:PRO:HD2	22:5:307:CLA:OBD	2.10	0.51
22:5:308:CLA:HMD2	22:5:313:CLA:C1D	2.41	0.51
39:5:317:CHL:HMB1	39:5:317:CHL:CBB	2.40	0.51
18:6:106:LEU:CD1	18:6:110:ILE:HG23	2.41	0.51
22:9:308:CLA:HMB1	22:9:308:CLA:CBB	2.39	0.51
1:A:396:TRP:HA	1:A:603:LEU:HD23	1.93	0.51
22:A:822:CLA:HBC3	22:A:823:CLA:H171	1.93	0.51
2:B:145:LEU:O	2:B:149:ILE:HG13	2.11	0.51
22:B:822:CLA:C1B	24:B:844:BCR:H351	2.40	0.51
10:K:62:ASP:HB2	10:K:68:VAL:HG23	1.92	0.51
13:3:14:GLN:N	13:3:16:TYR:HH	2.09	0.51
22:3:309:CLA:HMB1	22:3:309:CLA:CBB	2.36	0.51
16:4:157:PHE:CZ	22:4:815:CLA:HBC2	2.46	0.51
22:5:313:CLA:HHC	22:5:313:CLA:HBB1	1.93	0.51
22:5:326:CLA:HHC	22:5:326:CLA:CBB	2.41	0.51
22:A:810:CLA:C4C	22:A:813:CLA:H203	2.40	0.51
22:A:833:CLA:H43	22:A:855:CLA:H43	1.92	0.51
23:A:843:PQN:C19	24:F:303:BCR:H23C	2.41	0.51
22:F:304:CLA:H2	22:F:304:CLA:C9	2.41	0.51
18:6:142:PRO:HD2	39:6:320:CHL:CHC	2.41	0.51
25:6:323:LHG:C12	25:6:323:LHG:H292	2.40	0.51
19:9:114:MET:SD	22:2:302:CLA:HAA1	2.51	0.51
19:9:204:HIS:HA	19:9:206:GLN:OE1	2.11	0.51

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
20:2:197:ASN:HA	22:2:301:CLA:CBA	2.34	0.51
1:A:493:LEU:O	1:A:496:GLN:HG2	2.09	0.51
22:A:833:CLA:CAD	24:B:853:BCR:H10C	2.41	0.51
2:B:46:LYS:HG2	25:B:803:LHG:HC12	1.92	0.51
22:B:841:CLA:HBA2	23:B:843:PQN:H241	1.92	0.51
6:F:186:ILE:HG12	9:J:10:THR:HG22	1.92	0.51
11:L:94:PHE:O	11:L:97:PRO:HD2	2.11	0.51
12:1:58:LEU:HD12	38:1:303:LUT:H222	1.92	0.51
12:1:173:LYS:HD3	22:1:305:CLA:HAA2	1.92	0.51
15:8:164:PRO:HB3	22:8:313:CLA:HBC2	1.94	0.51
16:4:71:LEU:HB3	16:4:78:ASP:OD1	2.11	0.51
22:4:805:CLA:HMD2	39:4:814:CHL:H11	1.92	0.51
25:4:820:LHG:H131	22:6:302:CLA:H2	1.92	0.51
18:6:64:LEU:CD1	22:6:310:CLA:HMA2	2.41	0.51
18:6:245:PHE:HE2	22:6:301:CLA:H91	1.76	0.51
22:2:304:CLA:HHC	22:2:304:CLA:CBB	2.41	0.51
1:A:526:ILE:HG23	1:A:613:LYS:HE3	1.93	0.50
22:A:814:CLA:HMB1	22:A:814:CLA:CBB	2.36	0.50
22:A:829:CLA:HHC	22:A:829:CLA:HBB1	1.92	0.50
22:A:855:CLA:HHC	22:A:855:CLA:CBB	2.41	0.50
2:B:263:HIS:HB3	2:B:266:THR:OG1	2.11	0.50
2:B:276:HIS:ND1	22:B:819:CLA:HAB	2.26	0.50
22:F:304:CLA:C2D	22:F:304:CLA:H72	2.40	0.50
12:1:84:GLY:HA2	38:1:303:LUT:H181	1.93	0.50
24:3:305:BCR:H16C	39:3:317:CHL:H112	1.94	0.50
22:7:308:CLA:HMB1	22:7:308:CLA:CBB	2.35	0.50
15:8:125:MET:HB3	24:8:304:BCR:C15	2.41	0.50
36:5:306:C7Z:C36	22:5:322:CLA:HAC1	2.42	0.50
22:A:809:CLA:H11	24:J:104:BCR:C12	2.41	0.50
22:A:840:CLA:H92	24:F:303:BCR:H17C	1.93	0.50
2:B:463:TRP:HE1	2:B:477:LEU:HG	1.75	0.50
2:B:597:TRP:HH2	2:B:613:SER:HB2	1.77	0.50
12:Z:203:LEU:HA	22:Z:305:CLA:HMA3	1.92	0.50
14:7:234:PHE:HD1	22:7:306:CLA:H11	1.75	0.50
22:7:310:CLA:H171	22:7:310:CLA:H122	1.94	0.50
38:5:303:LUT:H373	22:5:310:CLA:H8	1.94	0.50
22:6:309:CLA:HBA1	22:6:309:CLA:CBD	2.34	0.50
1:A:49:ILE:O	1:A:52:LEU:HD23	2.11	0.50
1:A:204:GLY:O	1:A:208:LEU:HB2	2.12	0.50
1:A:552:LYS:HD3	2:B:671:TYR:CE1	2.46	0.50
22:A:809:CLA:HHC	22:A:809:CLA:CBB	2.39	0.50

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
24:A:844:BCR:C40	24:A:856:BCR:H12C	2.42	0.50
25:B:803:LHG:HC32	25:B:803:LHG:H242	1.91	0.50
12:Z:163:ALA:HB2	22:Z:303:CLA:HAA2	1.92	0.50
22:7:311:CLA:HBA2	22:7:311:CLA:CHA	2.41	0.50
15:8:52:PRO:HD2	38:8:303:LUT:C23	2.31	0.50
15:8:152:PHE:HB3	15:8:156:PHE:CE2	2.47	0.50
15:8:169:ASP:HA	38:8:302:LUT:C24	2.42	0.50
39:4:814:CHL:HMB1	39:4:814:CHL:CBB	2.38	0.50
22:4:815:CLA:HMB1	22:4:815:CLA:CBB	2.34	0.50
17:5:127:LEU:HD12	22:5:326:CLA:CBA	2.42	0.50
1:A:75:SER:O	1:A:78:PHE:HB2	2.11	0.50
22:A:833:CLA:CHA	24:B:853:BCR:H12C	2.41	0.50
22:B:808:CLA:HMB1	22:B:808:CLA:CBB	2.34	0.50
3:C:22:PRO:HG2	3:C:23:LEU:HD12	1.94	0.50
6:F:155:GLY:O	6:F:159:VAL:HG12	2.11	0.50
22:F:302:CLA:H162	32:F:306:RRX:C40	2.42	0.50
11:L:92:HIS:HA	11:L:95:PHE:CE2	2.47	0.50
24:3:307:BCR:C31	22:3:315:CLA:HHB	2.38	0.50
18:6:107:PRO:HD2	39:6:318:CHL:HMD1	1.94	0.50
18:6:214:THR:HA	22:6:309:CLA:CBA	2.40	0.50
25:6:323:LHG:HC92	25:6:323:LHG:H272	1.93	0.50
1:A:28:LYS:HE2	1:A:34:HIS:CG	2.47	0.50
22:A:806:CLA:C4	25:A:850:LHG:H262	2.42	0.50
2:B:444:MET:HB3	2:B:449:THR:O	2.11	0.50
24:B:847:BCR:H311	22:1:308:CLA:H72	1.93	0.50
12:Z:218:ASN:ND2	22:Z:305:CLA:HMD1	2.26	0.50
15:8:169:ASP:N	15:8:170:PRO:HD3	2.27	0.50
18:6:216:PHE:HA	22:6:322:CLA:HMB1	1.93	0.50
22:A:839:CLA:H11	24:A:847:BCR:C15	2.41	0.50
24:A:844:BCR:H23C	24:3:306:BCR:C33	2.40	0.50
22:B:806:CLA:H171	22:L:201:CLA:CMC	2.41	0.50
22:B:811:CLA:CBB	22:B:812:CLA:HAA1	2.42	0.50
22:B:825:CLA:NA	24:B:847:BCR:H362	2.27	0.50
11:L:99:PRO:HG3	22:L:203:CLA:CAB	2.41	0.50
12:1:160:LEU:H	12:1:160:LEU:HD12	1.77	0.50
38:Z:301:LUT:H30	22:Z:303:CLA:C5	2.39	0.50
22:3:308:CLA:HMB1	22:3:308:CLA:CBB	2.39	0.50
14:7:129:LEU:HD21	25:8:317:LHG:H322	1.94	0.50
15:8:202:PHE:CD1	38:8:302:LUT:H41	2.47	0.50
16:4:183:ASN:HB3	39:4:819:CHL:C3D	2.41	0.50
38:4:803:LUT:H32	22:4:808:CLA:CAB	2.41	0.50

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
18:6:127:LYS:HB3	39:6:316:CHL:CMC	2.40	0.50
1:A:305:PHE:HE1	22:A:821:CLA:HAB	1.76	0.50
1:A:359:LEU:HD21	22:A:821:CLA:H93	1.94	0.50
22:A:814:CLA:H143	27:A:852:DGA:HBT1	1.94	0.50
22:A:815:CLA:HMB1	22:A:815:CLA:CBB	2.36	0.50
22:A:821:CLA:HMB1	22:A:821:CLA:CBB	2.39	0.50
2:B:16:ASP:OD2	2:B:20:ARG:HG3	2.12	0.50
22:B:811:CLA:CGA	22:B:811:CLA:C1A	2.89	0.50
22:B:817:CLA:H112	24:B:844:BCR:H21C	1.93	0.50
22:B:818:CLA:HHC	22:B:818:CLA:CBB	2.42	0.50
4:D:125:THR:HG22	4:D:129:ARG:HD2	1.93	0.50
9:J:35:ASP:N	9:J:36:PRO:HD3	2.27	0.50
22:1:305:CLA:C4D	22:1:310:CLA:H8	2.42	0.50
12:Z:221:SER:HA	22:Z:316:CLA:C4D	2.41	0.50
22:Z:306:CLA:H72	22:Z:307:CLA:HHB	1.92	0.50
22:8:313:CLA:HMB1	22:8:313:CLA:HBB1	1.92	0.50
38:6:304:LUT:H31	22:6:310:CLA:HMC2	1.94	0.50
38:9:301:LUT:H403	22:9:303:CLA:H13	1.93	0.50
38:9:302:LUT:C32	22:9:306:CLA:HAB	2.42	0.50
1:A:536:HIS:HA	1:A:539:HIS:HD2	1.76	0.50
1:A:572:ARG:HH21	25:A:850:LHG:C3	2.24	0.50
1:A:679:TRP:CD2	21:A:801:CL0:H5	2.47	0.50
22:A:813:CLA:H201	22:3:312:CLA:H72	1.93	0.50
22:A:829:CLA:CHA	22:A:829:CLA:H12	2.42	0.50
22:A:838:CLA:HMB1	22:A:838:CLA:CBB	2.41	0.50
2:B:5:LEU:HD12	8:I:101:LYS:HG3	1.94	0.50
2:B:222:GLY:O	2:B:225:PRO:HD2	2.12	0.50
22:B:830:CLA:H8	24:B:845:BCR:H19C	1.94	0.50
22:B:836:CLA:NC	31:1:319:DGD:HA82	2.26	0.50
22:F:304:CLA:HMC1	33:F:307:LMT:H121	1.94	0.50
11:L:86:VAL:HG22	11:L:90:ARG:HD3	1.93	0.50
11:L:120:SER:OG	22:L:201:CLA:H2	2.12	0.50
12:Z:58:LEU:HD21	15:8:170:PRO:HB3	1.93	0.50
22:Z:303:CLA:H102	22:Z:304:CLA:CMB	2.41	0.50
14:7:130:LEU:HB3	24:7:303:BCR:C15	2.42	0.50
25:4:820:LHG:H381	22:6:321:CLA:HBD	1.94	0.50
17:5:64:PRO:O	17:5:68:LYS:HG3	2.11	0.50
18:6:147:ASN:ND2	39:6:320:CHL:HAA1	2.27	0.50
18:6:157:PRO:HG3	39:6:316:CHL:C2D	2.42	0.50
22:9:307:CLA:HED2	22:9:307:CLA:H2A	1.94	0.50
20:2:166:LEU:HA	20:2:169:ILE:CG1	2.41	0.50

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:71:ARG:HH21	1:A:187:LYS:HA	1.77	0.50
1:A:245:LEU:HD23	31:3:301:DGD:HG12	1.93	0.50
1:A:697:ILE:O	1:A:701:ILE:HG13	2.12	0.50
22:A:815:CLA:H51	13:3:200:ALA:HA	1.93	0.50
2:B:13:LEU:HD21	2:B:20:ARG:HH11	1.76	0.50
12:1:211:TRP:CZ3	15:8:118:LEU:HA	2.47	0.50
22:1:308:CLA:HMA2	22:1:308:CLA:H11	1.94	0.50
12:Z:103:PRO:HG2	39:Z:312:CHL:CED	2.41	0.50
12:Z:140:ARG:HG2	22:Z:313:CLA:CAB	2.42	0.50
22:Z:309:CLA:CBB	24:4:804:BCR:H323	2.42	0.50
22:7:307:CLA:HMB1	22:7:307:CLA:CBB	2.39	0.50
22:7:324:CLA:HMB1	22:7:324:CLA:CBB	2.42	0.50
16:4:185:LEU:HD11	16:4:197:PHE:CE2	2.47	0.50
24:4:804:BCR:H402	22:4:810:CLA:C5	2.42	0.50
22:5:312:CLA:HMB1	22:5:312:CLA:CBB	2.41	0.50
1:A:60:ASP:HA	1:A:69:ILE:HD13	1.93	0.49
2:B:143:LEU:HD21	24:B:846:BCR:H24C	1.94	0.49
2:B:178:HIS:HB3	22:B:815:CLA:CAB	2.42	0.49
2:B:478:LEU:HD13	22:B:835:CLA:HHD	1.94	0.49
22:B:838:CLA:H152	32:F:306:RRX:C35	2.42	0.49
7:G:78:PHE:CG	7:G:78:PHE:O	2.65	0.49
7:G:114:PHE:CE2	22:G:201:CLA:HBC2	2.47	0.49
22:1:307:CLA:HHD	39:8:301:CHL:HBB2	1.93	0.49
38:Z:302:LUT:H28	22:Z:306:CLA:H61	1.94	0.49
39:Z:311:CHL:HAA1	39:Z:311:CHL:HBD	1.93	0.49
38:8:302:LUT:C30	22:8:305:CLA:H61	2.41	0.49
17:5:188:LYS:HE3	25:5:324:LHG:HC61	1.94	0.49
22:5:311:CLA:C2	22:5:311:CLA:HMA2	2.42	0.49
19:9:115:GLU:CB	39:9:312:CHL:HBB2	2.41	0.49
1:A:498:THR:HB	22:A:835:CLA:HMD1	1.93	0.49
22:A:823:CLA:HHC	22:A:823:CLA:CBB	2.33	0.49
24:A:856:BCR:H321	24:A:856:BCR:HC8	1.94	0.49
2:B:393:ILE:CG2	2:B:397:ARG:HD2	2.42	0.49
2:B:646:VAL:HG21	22:B:811:CLA:HAC1	1.93	0.49
22:B:807:CLA:HBB1	22:B:807:CLA:HMB1	1.94	0.49
22:B:815:CLA:H141	22:B:820:CLA:NB	2.27	0.49
22:B:827:CLA:H111	22:B:839:CLA:HMD2	1.93	0.49
6:F:138:GLY:HA3	22:F:304:CLA:C2D	2.41	0.49
22:F:302:CLA:HMA2	24:F:303:BCR:H362	1.93	0.49
12:1:160:LEU:HD12	12:1:160:LEU:N	2.28	0.49
38:1:302:LUT:H11	22:1:305:CLA:HMC2	1.93	0.49

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
12:Z:144:GLU:HG3	22:Z:313:CLA:HAC1	1.93	0.49
13:3:220:PRO:HG2	22:3:315:CLA:HMB3	1.93	0.49
17:5:128:LEU:HB3	24:5:304:BCR:C15	2.42	0.49
38:9:301:LUT:H34	22:9:303:CLA:HBB2	1.94	0.49
38:9:301:LUT:H28	22:9:303:CLA:H2	1.92	0.49
22:A:826:CLA:HHB	22:A:839:CLA:O2A	2.12	0.49
2:B:71:TRP:CD1	8:I:71:PRO:HD2	2.47	0.49
2:B:278:HIS:HA	2:B:281:ILE:HG12	1.94	0.49
2:B:349:VAL:O	2:B:353:MET:HB2	2.13	0.49
2:B:689:ALA:HB2	24:B:853:BCR:HC22	1.94	0.49
2:B:705:GLN:O	2:B:708:LEU:HG	2.11	0.49
22:B:820:CLA:O1A	22:B:830:CLA:HMD1	2.12	0.49
22:F:301:CLA:HHC	22:F:301:CLA:CBB	2.42	0.49
12:Z:144:GLU:HG3	22:Z:313:CLA:CAC	2.43	0.49
22:3:311:CLA:H62	22:3:312:CLA:HHB	1.95	0.49
22:4:812:CLA:C4D	22:6:302:CLA:H111	2.42	0.49
22:5:309:CLA:HHC	22:5:309:CLA:CBB	2.41	0.49
22:5:310:CLA:HMB1	22:5:310:CLA:HBB1	1.94	0.49
18:6:140:ALA:HA	18:6:148:LYS:HA	1.93	0.49
19:9:84:ILE:HG22	19:9:183:PRO:HG2	1.94	0.49
2:B:6:PHE:CE2	2:B:22:ILE:HA	2.47	0.49
2:B:478:LEU:CD1	22:B:835:CLA:HHD	2.42	0.49
4:D:184:VAL:CG1	4:D:194:ILE:HG21	2.42	0.49
22:G:201:CLA:HMC1	24:G:203:BCR:H332	1.94	0.49
22:1:305:CLA:HED2	22:1:305:CLA:H2A	1.93	0.49
22:1:316:CLA:HED1	22:1:316:CLA:H2	1.95	0.49
22:3:314:CLA:HHC	22:3:314:CLA:CBB	2.42	0.49
16:4:118:ASP:HA	16:4:121:ARG:HE	1.78	0.49
17:5:106:PRO:HA	17:5:109:LEU:HD12	1.93	0.49
1:A:178:TRP:HB2	22:A:811:CLA:HMC3	1.94	0.49
1:A:272:TYR:HB3	1:A:275:PHE:CZ	2.48	0.49
22:A:805:CLA:HAA2	22:A:825:CLA:C4	2.41	0.49
22:A:806:CLA:H8	22:A:830:CLA:HBC3	1.94	0.49
22:A:817:CLA:HMB1	22:A:817:CLA:CBB	2.43	0.49
22:A:854:CLA:HMB1	22:A:854:CLA:CBB	2.42	0.49
2:B:18:THR:HG22	3:C:79:LEU:HD11	1.94	0.49
2:B:107:ARG:HH22	2:B:115:ASN:HD22	1.60	0.49
22:B:817:CLA:HMB1	22:B:817:CLA:CBB	2.43	0.49
22:B:840:CLA:HMB1	22:B:840:CLA:CBB	2.38	0.49
22:Z:307:CLA:HBB1	22:Z:307:CLA:HMB1	1.93	0.49
25:Z:317:LHG:HC92	24:4:804:BCR:HC22	1.94	0.49

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
13:3:103:TYR:HE2	13:3:105:LYS:O	1.95	0.49
22:3:318:CLA:HMB1	22:3:318:CLA:CBB	2.42	0.49
14:7:58:LEU:HD11	38:7:302:LUT:H221	1.94	0.49
17:5:127:LEU:HD12	22:5:326:CLA:HBA1	1.93	0.49
18:6:87:VAL:C	18:6:89:GLU:H	2.16	0.49
18:6:235:GLN:HE21	22:6:301:CLA:HHB	1.78	0.49
39:6:315:CHL:HMB1	39:6:315:CHL:CBB	2.36	0.49
19:9:62:ARG:HB3	19:9:66:PHE:CE2	2.47	0.49
22:A:815:CLA:H122	13:3:196:PHE:CD2	2.47	0.49
22:A:841:CLA:H51	22:B:801:CLA:H203	1.94	0.49
22:A:855:CLA:H193	24:B:853:BCR:C35	2.43	0.49
2:B:651:PHE:CE2	2:B:655:HIS:HE1	2.31	0.49
2:B:715:SER:HB2	31:B:852:DGD:HBH2	1.95	0.49
22:B:805:CLA:HMB3	22:B:806:CLA:OBD	2.13	0.49
22:B:838:CLA:HMB1	22:B:838:CLA:CBB	2.43	0.49
39:1:312:CHL:HAB	22:1:315:CLA:HBB2	1.93	0.49
14:7:202:VAL:HG21	25:7:318:LHG:H271	1.93	0.49
39:8:315:CHL:HMB1	39:8:315:CHL:CBB	2.37	0.49
22:5:307:CLA:H101	22:5:308:CLA:H122	1.94	0.49
22:5:308:CLA:HMB1	22:5:308:CLA:CBB	2.39	0.49
1:A:25:SER:O	22:A:811:CLA:HMA1	2.13	0.49
22:A:815:CLA:HMA2	31:3:301:DGD:O1A	2.12	0.49
22:A:824:CLA:H141	25:A:849:LHG:H171	1.94	0.49
22:A:837:CLA:C3A	22:A:838:CLA:HAA2	2.42	0.49
22:A:841:CLA:HHC	22:A:841:CLA:HBB1	1.94	0.49
2:B:164:PRO:HB2	2:B:169:PHE:CE2	2.48	0.49
2:B:227:TRP:CZ2	22:B:817:CLA:H52	2.48	0.49
22:B:838:CLA:CGA	22:B:838:CLA:C1A	2.90	0.49
22:B:838:CLA:NC	22:B:838:CLA:H42	2.28	0.49
4:D:76:ARG:O	4:D:80:VAL:HG12	2.12	0.49
24:L:204:BCR:C12	24:L:204:BCR:H341	2.43	0.49
22:1:304:CLA:HMB1	22:1:304:CLA:CBB	2.36	0.49
22:1:310:CLA:HBC3	25:1:317:LHG:O8	2.13	0.49
12:Z:191:GLN:HE22	12:Z:199:PRO:HG3	1.77	0.49
22:Z:307:CLA:H12	15:8:151:GLY:O	2.13	0.49
22:3:313:CLA:HMB1	22:3:313:CLA:CBB	2.35	0.49
22:3:314:CLA:H41	22:3:314:CLA:H101	1.93	0.49
14:7:58:LEU:HB2	22:7:307:CLA:H11	1.94	0.49
14:7:83:ALA:HB1	38:7:301:LUT:H12	1.93	0.49
25:4:820:LHG:H122	24:6:305:BCR:H333	1.94	0.49
17:5:35:TRP:CZ3	22:5:315:CLA:HAA1	2.48	0.49

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
17:5:80:TRP:CE2	22:5:318:CLA:HBC3	2.47	0.49
22:5:309:CLA:H11	22:5:314:CLA:HMD1	1.94	0.49
18:6:73:VAL:HG21	18:6:128:ARG:CZ	2.41	0.49
22:6:310:CLA:H151	22:6:311:CLA:HBB1	1.94	0.49
22:A:807:CLA:HHC	22:A:807:CLA:HBB1	1.94	0.49
2:B:306:LEU:HD23	22:B:825:CLA:O1D	2.12	0.49
22:B:810:CLA:C1B	24:I:4001:BCR:HC22	2.42	0.49
22:B:814:CLA:O1D	22:B:815:CLA:HMC1	2.12	0.49
22:1:309:CLA:HMA2	22:1:315:CLA:C3C	2.43	0.49
38:Z:302:LUT:H161	22:Z:308:CLA:HMB3	1.95	0.49
13:3:121:MET:CB	22:3:318:CLA:HAB	2.42	0.49
38:8:303:LUT:H163	39:8:315:CHL:CMC	2.43	0.49
22:8:309:CLA:HMD2	22:8:314:CLA:NC	2.28	0.49
16:4:259:ASP:HB3	22:4:817:CLA:CHB	2.42	0.49
22:5:311:CLA:HMD3	22:5:318:CLA:H93	1.94	0.49
22:5:322:CLA:C14	22:6:310:CLA:H101	2.42	0.49
1:A:21:PRO:HG2	1:A:185:ALA:HB3	1.95	0.49
1:A:53:HIS:HD2	22:A:804:CLA:HAC1	1.77	0.49
1:A:545:VAL:HG11	1:A:598:TRP:CZ2	2.48	0.49
2:B:71:TRP:CZ2	8:I:71:PRO:HB2	2.47	0.49
6:F:138:GLY:O	6:F:142:LYS:HB2	2.13	0.49
7:G:71:LYS:HA	7:G:71:LYS:HE2	1.94	0.49
10:K:30:ILE:HA	10:K:35:ASN:ND2	2.28	0.49
22:1:306:CLA:H171	24:8:304:BCR:H321	1.93	0.49
22:Z:308:CLA:HBA2	22:Z:308:CLA:CHA	2.43	0.49
13:3:114:PHE:HB2	22:3:316:CLA:HAA1	1.95	0.49
16:4:210:LEU:C	22:4:805:CLA:HMA1	2.33	0.49
38:9:301:LUT:H34	22:9:303:CLA:CBB	2.42	0.49
22:A:810:CLA:H142	22:A:813:CLA:H151	1.95	0.49
22:A:835:CLA:H201	22:A:842:CLA:HBB2	1.95	0.49
22:B:815:CLA:HMB1	22:B:815:CLA:CBB	2.42	0.49
22:B:837:CLA:HMB1	22:B:837:CLA:CBB	2.41	0.49
8:I:74:TRP:HZ2	22:2:304:CLA:HHB	1.78	0.49
10:K:107:LEU:HD22	10:K:112:ASN:HD22	1.78	0.49
22:K:205:CLA:CHA	24:K:206:BCR:H393	2.43	0.49
22:1:315:CLA:HMB1	22:1:315:CLA:CBB	2.39	0.49
13:3:117:GLU:O	13:3:121:MET:HG2	2.13	0.49
13:3:203:VAL:O	31:3:301:DGD:HG31	2.12	0.49
22:3:311:CLA:HMB1	22:3:311:CLA:CBB	2.36	0.49
39:7:313:CHL:C1C	39:7:313:CHL:H71	2.43	0.49
15:8:187:LYS:HD3	22:8:306:CLA:HBA1	1.94	0.49

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
16:4:180:PHE:HB3	16:4:182:ASN:OD1	2.12	0.49
39:4:816:CHL:HBA2	39:4:816:CHL:O2D	2.12	0.49
22:6:322:CLA:H112	22:6:322:CLA:H72	1.60	0.49
22:9:307:CLA:C3D	22:9:313:CLA:H43	2.43	0.49
20:2:178:ARG:HB3	22:2:304:CLA:HED2	1.94	0.49
1:A:202:LEU:O	1:A:305:PHE:HB3	2.12	0.48
1:A:203:ALA:HB2	1:A:309:GLY:HA3	1.93	0.48
1:A:668:TYR:CE1	22:A:808:CLA:HAC1	2.48	0.48
22:A:840:CLA:HMB1	22:A:840:CLA:CBB	2.41	0.48
2:B:142:PHE:CZ	22:B:816:CLA:H122	2.48	0.48
22:B:825:CLA:HMB1	22:B:825:CLA:CBB	2.38	0.48
3:C:5:VAL:HG11	3:C:26:LEU:HD21	1.95	0.48
6:F:107:THR:HA	6:F:110:ARG:NE	2.27	0.48
22:F:301:CLA:H122	22:F:301:CLA:H8	1.61	0.48
10:K:59:ARG:H	10:K:69:ASP:HA	1.78	0.48
12:Z:192:HIS:O	12:Z:196:GLY:HA2	2.12	0.48
14:7:137:LYS:HE3	22:7:314:CLA:HMC3	1.95	0.48
38:7:302:LUT:C31	22:7:307:CLA:HMC2	2.43	0.48
15:8:153:GLU:HA	15:8:156:PHE:HD2	1.78	0.48
22:4:808:CLA:H43	22:4:808:CLA:HMB2	1.94	0.48
22:4:808:CLA:HBC1	25:4:820:LHG:H282	1.95	0.48
17:5:109:LEU:HD23	22:5:312:CLA:HAA2	1.95	0.48
18:6:110:ILE:CD1	24:6:306:BCR:HC8	2.38	0.48
18:6:141:ASP:HB2	18:6:147:ASN:HB2	1.95	0.48
22:6:309:CLA:NC	22:6:309:CLA:H51	2.28	0.48
39:9:312:CHL:CHB	39:9:314:CHL:HBC2	2.43	0.48
1:A:396:TRP:CZ3	1:A:600:TYR:HA	2.48	0.48
21:A:801:CL0:H13	22:A:802:CLA:OBD	2.13	0.48
2:B:223:LEU:HB3	2:B:227:TRP:NE1	2.28	0.48
2:B:412:MET:HA	2:B:415:HIS:CE1	2.48	0.48
8:I:78:VAL:C	8:I:81:PRO:HD2	2.33	0.48
12:Z:69:ARG:NH2	15:8:150:LEU:HD23	2.27	0.48
22:Z:309:CLA:HHC	22:Z:309:CLA:CBB	2.43	0.48
13:3:143:PHE:CE2	39:3:317:CHL:HBB2	2.48	0.48
16:4:258:ASN:O	16:4:262:ARG:HG3	2.13	0.48
17:5:105:GLU:HB3	17:5:107:GLN:NE2	2.27	0.48
24:5:305:BCR:H21C	24:5:305:BCR:H361	1.94	0.48
22:5:307:CLA:HMB1	22:5:307:CLA:CBB	2.39	0.48
22:6:311:CLA:HMB1	22:6:311:CLA:CBB	2.36	0.48
19:9:31:TRP:CD1	19:9:51:PHE:HA	2.48	0.48
1:A:501:THR:HG21	22:K:203:CLA:HED1	1.94	0.48

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:B:54:GLN:O	2:B:58:ILE:HG13	2.14	0.48
2:B:584:MET:O	2:B:588:ILE:HG12	2.13	0.48
22:B:814:CLA:HBB2	22:B:822:CLA:H93	1.95	0.48
22:B:825:CLA:HMC3	22:B:832:CLA:HMD3	1.94	0.48
13:3:149:VAL:HG22	13:3:163:TRP:HB2	1.93	0.48
14:7:82:GLY:HA2	38:7:302:LUT:H181	1.95	0.48
14:7:125:ALA:HB2	15:8:230:VAL:HG21	1.94	0.48
22:8:309:CLA:HAC1	39:8:312:CHL:HBB2	1.95	0.48
16:4:161:GLU:HG3	22:4:815:CLA:C4B	2.43	0.48
24:4:804:BCR:C16	39:4:814:CHL:HMB3	2.42	0.48
22:5:301:CLA:HHC	22:5:301:CLA:CBB	2.43	0.48
19:9:76:TRP:CD2	22:9:313:CLA:HAC2	2.47	0.48
1:A:678:VAL:HG21	24:A:848:BCR:H353	1.95	0.48
22:A:822:CLA:C4D	24:A:856:BCR:H292	2.44	0.48
23:A:843:PQN:H201	24:F:303:BCR:H382	1.95	0.48
24:A:848:BCR:H24C	22:B:801:CLA:H101	1.95	0.48
2:B:23:TRP:HB3	31:B:852:DGD:C4A	2.44	0.48
2:B:32:PHE:HB2	2:B:43:LEU:HD13	1.95	0.48
2:B:177:ASN:OD1	2:B:292:TYR:HD2	1.97	0.48
22:B:824:CLA:H11	33:G:204:LMT:C6'	2.43	0.48
22:B:834:CLA:HMB1	22:B:834:CLA:CBB	2.42	0.48
22:B:841:CLA:H203	24:L:204:BCR:H15C	1.95	0.48
24:B:844:BCR:H272	24:B:846:BCR:H323	1.96	0.48
3:C:3:HIS:ND1	3:C:69:LEU:HA	2.28	0.48
6:F:121:ALA:HB3	6:F:123:LEU:HD13	1.95	0.48
22:F:304:CLA:C1D	22:F:304:CLA:H52	2.44	0.48
7:G:33:GLU:HB3	7:G:34:PRO:HD2	1.95	0.48
7:G:54:LEU:O	7:G:58:ARG:HG2	2.13	0.48
9:J:34:PRO:HD2	35:J:102:T7X:C5	2.43	0.48
12:1:191:GLN:HG3	12:1:192:HIS:N	2.27	0.48
22:1:305:CLA:HMD2	22:1:310:CLA:C1D	2.43	0.48
13:3:141:GLN:NE2	22:3:322:CLA:HMD3	2.27	0.48
22:7:310:CLA:H2	22:7:310:CLA:H61	1.64	0.48
22:7:316:CLA:HMB1	22:7:316:CLA:CBB	2.43	0.48
39:8:312:CHL:CHD	39:8:315:CHL:HBC2	2.43	0.48
16:4:114:ILE:HG23	16:4:134:TRP:CD1	2.48	0.48
16:4:177:ASP:CB	16:4:183:ASN:HB2	2.42	0.48
18:6:159:PHE:O	38:6:303:LUT:H24	2.12	0.48
1:A:69:ILE:O	1:A:73:VAL:HG23	2.13	0.48
1:A:382:THR:HG21	1:A:517:VAL:HB	1.94	0.48
1:A:487:ILE:HD11	22:A:837:CLA:H43	1.95	0.48

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:563:ILE:HD11	1:A:583:THR:HB	1.95	0.48
22:A:810:CLA:O1A	22:A:813:CLA:H193	2.13	0.48
22:A:813:CLA:H191	22:3:312:CLA:H41	1.95	0.48
22:A:824:CLA:HMD1	22:A:825:CLA:HAB	1.94	0.48
22:A:835:CLA:NB	24:A:847:BCR:H292	2.28	0.48
22:A:836:CLA:C4D	22:K:203:CLA:H12	2.44	0.48
22:A:841:CLA:H62	22:A:841:CLA:H102	1.48	0.48
22:B:819:CLA:HHC	22:B:819:CLA:CBB	2.43	0.48
22:B:821:CLA:H42	22:B:825:CLA:O1A	2.14	0.48
22:B:842:CLA:HBA1	25:B:850:LHG:HC42	1.95	0.48
33:1:301:LMT:H31	22:1:308:CLA:H42	1.95	0.48
22:1:314:CLA:HBB1	22:1:314:CLA:HMB1	1.94	0.48
13:3:163:TRP:O	13:3:166:LEU:HB3	2.14	0.48
38:3:302:LUT:H382	22:3:308:CLA:C4D	2.43	0.48
22:7:308:CLA:HMA2	22:7:308:CLA:H12	1.96	0.48
25:4:820:LHG:HC91	25:4:820:LHG:H292	1.94	0.48
25:6:323:LHG:H292	25:6:323:LHG:H122	1.96	0.48
38:9:302:LUT:H161	39:9:314:CHL:HBB1	1.96	0.48
39:9:312:CHL:HBC2	39:9:312:CHL:CMC	2.39	0.48
1:A:25:SER:O	22:A:811:CLA:HHB	2.13	0.48
1:A:427:LEU:O	1:A:431:ILE:HG13	2.14	0.48
1:A:441:LEU:O	1:A:444:VAL:HG12	2.13	0.48
1:A:541:PHE:CZ	22:B:806:CLA:HBB2	2.46	0.48
22:A:811:CLA:HHC	22:A:811:CLA:CBB	2.42	0.48
22:A:823:CLA:NC	37:K:201:DAO:H111	2.29	0.48
2:B:490:SER:CB	2:B:495:LEU:HB2	2.44	0.48
22:B:810:CLA:HMB3	22:B:811:CLA:HHB	1.96	0.48
22:B:832:CLA:H2	22:B:842:CLA:C3	2.43	0.48
22:Z:305:CLA:H43	22:Z:310:CLA:OBD	2.14	0.48
13:3:202:ALA:O	31:3:301:DGD:HB31	2.13	0.48
24:3:306:BCR:C12	24:3:306:BCR:H341	2.42	0.48
22:3:310:CLA:H62	22:3:315:CLA:HMD1	1.95	0.48
15:8:208:ALA:HA	15:8:234:ILE:HG21	1.95	0.48
22:4:807:CLA:C1A	22:4:807:CLA:CGA	2.91	0.48
22:6:312:CLA:HMB1	22:6:312:CLA:CBB	2.43	0.48
20:2:199:TYR:O	20:2:204:GLY:HA3	2.14	0.48
1:A:82:GLY:O	1:A:86:ILE:HG13	2.13	0.48
1:A:367:VAL:HG11	22:A:829:CLA:HMD3	1.95	0.48
1:A:495:PRO:O	1:A:500:PRO:HA	2.13	0.48
1:A:673:LEU:HB3	22:A:802:CLA:O2A	2.13	0.48
22:A:805:CLA:HMB1	22:A:805:CLA:CBB	2.41	0.48

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
22:A:814:CLA:HBA1	24:A:844:BCR:C34	2.44	0.48
22:A:821:CLA:H2	22:A:821:CLA:H61	1.56	0.48
22:A:821:CLA:C10	24:A:847:BCR:H10C	2.41	0.48
22:A:834:CLA:HMB1	22:A:834:CLA:CBB	2.44	0.48
2:B:295:ASN:HB2	7:G:83:GLN:O	2.13	0.48
6:F:123:LEU:HB3	6:F:131:PRO:HB3	1.96	0.48
9:J:16:LEU:HD21	36:J:105:C7Z:C12	2.44	0.48
22:Z:303:CLA:H41	22:Z:304:CLA:HBA1	1.95	0.48
22:Z:305:CLA:HBC1	22:Z:316:CLA:C10	2.42	0.48
14:7:243:PHE:HE2	22:7:317:CLA:CAB	2.26	0.48
16:4:128:PRO:HD2	22:4:818:CLA:HMA3	1.96	0.48
16:4:183:ASN:OD1	39:4:819:CHL:HAA1	2.14	0.48
17:5:160:PRO:HG2	17:5:179:ARG:HH12	1.77	0.48
39:5:316:CHL:HAA1	39:5:316:CHL:HBD	1.95	0.48
22:5:322:CLA:O2D	22:6:314:CLA:HMB2	2.13	0.48
18:6:237:THR:HG23	18:6:246:THR:OG1	2.13	0.48
22:6:311:CLA:CBB	22:6:319:CLA:H42	2.44	0.48
19:9:100:GLY:O	39:9:314:CHL:HBC1	2.14	0.48
39:9:312:CHL:HBC1	22:9:313:CLA:HBC3	1.94	0.48
1:A:248:GLN:CD	1:A:248:GLN:H	2.17	0.48
22:A:822:CLA:H8	22:A:823:CLA:C9	2.44	0.48
2:B:613:SER:HA	2:B:616:TYR:CE1	2.49	0.48
22:B:834:CLA:O2A	9:J:36:PRO:HG2	2.14	0.48
11:L:126:ILE:HD12	22:L:201:CLA:H152	1.96	0.48
15:8:107:SER:OG	39:8:315:CHL:HAC1	2.14	0.48
15:8:184:LEU:C	22:8:305:CLA:HMA1	2.34	0.48
22:5:312:CLA:HBA1	22:5:312:CLA:H3A	1.72	0.48
22:5:318:CLA:H92	22:6:322:CLA:H62	1.95	0.48
18:6:122:HIS:CE1	18:6:126:LEU:HD11	2.48	0.48
20:2:199:TYR:OH	20:2:211:VAL:HG21	2.13	0.48
1:A:26:PHE:CD2	34:J:101:LMG:HC1	2.49	0.48
22:A:805:CLA:C17	24:A:844:BCR:H323	2.44	0.48
2:B:168:TRP:NE1	22:B:813:CLA:HMA1	2.29	0.48
2:B:217:LEU:HD23	22:B:817:CLA:O1D	2.14	0.48
2:B:688:LEU:CB	24:B:853:BCR:HC31	2.43	0.48
22:B:815:CLA:H71	22:B:815:CLA:H111	1.56	0.48
22:B:815:CLA:H172	22:B:830:CLA:CAD	2.44	0.48
22:G:202:CLA:HED2	22:G:202:CLA:H2A	1.95	0.48
9:J:3:ASN:HB2	34:J:101:LMG:HC3	1.95	0.48
22:J:103:CLA:HMB1	22:J:103:CLA:CBB	2.40	0.48
11:L:135:TYR:CZ	11:L:139:MET:HG3	2.49	0.48

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
39:Z:312:CHL:HAA1	39:Z:312:CHL:HBD	1.95	0.48
22:3:322:CLA:H141	22:7:317:CLA:HBC2	1.96	0.48
39:8:312:CHL:C1C	26:8:318:NKP:HAPA	2.44	0.48
16:4:237:LYS:HB2	16:4:242:ASN:ND2	2.28	0.48
22:5:322:CLA:H2A	22:5:322:CLA:HED3	1.95	0.48
18:6:52:PHE:HB3	22:6:310:CLA:C3D	2.43	0.48
19:9:69:ALA:HA	19:9:72:THR:HG22	1.96	0.48
22:A:810:CLA:H51	22:3:312:CLA:H51	1.95	0.48
2:B:32:PHE:HB2	2:B:43:LEU:CD1	2.43	0.48
2:B:312:PRO:HD2	25:B:850:LHG:P	2.54	0.48
2:B:655:HIS:HB3	22:B:806:CLA:HBD	1.96	0.48
5:E:47:ARG:HB3	5:E:50:SER:OG	2.13	0.48
11:L:96:LEU:HB3	11:L:97:PRO:HD3	1.95	0.48
12:Z:60:LEU:CD1	38:Z:302:LUT:H221	2.43	0.48
13:3:20:GLU:O	13:3:24:GLN:HG3	2.14	0.48
38:7:301:LUT:H26	22:7:304:CLA:O1A	2.14	0.48
22:4:812:CLA:H72	22:6:302:CLA:H121	1.96	0.48
22:6:309:CLA:HBD	22:6:309:CLA:CBA	2.38	0.48
1:A:307:VAL:CA	24:A:856:BCR:H17C	2.31	0.47
22:A:823:CLA:H12	37:K:201:DAO:H82	1.95	0.47
22:A:830:CLA:H112	25:A:850:LHG:H202	1.95	0.47
22:A:833:CLA:CBD	24:B:853:BCR:H12C	2.43	0.47
22:A:855:CLA:H141	24:B:853:BCR:C17	2.44	0.47
22:B:811:CLA:H111	22:B:811:CLA:H2	1.95	0.47
22:1:307:CLA:HMB1	22:1:307:CLA:CBB	2.43	0.47
14:7:58:LEU:CD1	38:7:302:LUT:H221	2.44	0.47
22:8:313:CLA:CHA	22:8:313:CLA:HBA2	2.44	0.47
16:4:192:TYR:HB3	22:4:805:CLA:O1D	2.14	0.47
18:6:148:LYS:C	39:6:320:CHL:HMD2	2.34	0.47
39:9:312:CHL:HHC	39:9:312:CHL:CBB	2.41	0.47
1:A:409:ALA:CB	24:A:846:BCR:H271	2.43	0.47
22:A:822:CLA:C1D	24:A:856:BCR:H402	2.43	0.47
22:A:833:CLA:H42	22:B:840:CLA:H41	1.96	0.47
22:A:835:CLA:O1A	22:A:836:CLA:HHB	2.13	0.47
22:A:842:CLA:CBB	24:A:846:BCR:H12C	2.44	0.47
2:B:561:ASP:CG	3:C:66:ARG:HH12	2.17	0.47
22:B:810:CLA:C2B	24:I:4001:BCR:HC22	2.45	0.47
22:B:818:CLA:H143	22:B:818:CLA:H111	1.64	0.47
6:F:153:THR:HG21	32:F:306:RRX:H57	1.95	0.47
7:G:35:TYR:HA	22:G:201:CLA:OBD	2.15	0.47
33:1:301:LMT:H52	22:1:314:CLA:H42	1.95	0.47

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
12:Z:203:LEU:O	12:Z:207:ILE:HG13	2.14	0.47
38:Z:302:LUT:C16	22:Z:308:CLA:HMB3	2.44	0.47
22:Z:308:CLA:HMB1	22:Z:308:CLA:CBB	2.44	0.47
13:3:59:SER:OG	22:3:312:CLA:HAA1	2.14	0.47
13:3:70:GLY:HA2	38:3:303:LUT:C18	2.40	0.47
15:8:30:TRP:CH2	41:8:320:3PH:H332	2.49	0.47
16:4:183:ASN:HB3	39:4:819:CHL:C4D	2.43	0.47
17:5:53:TYR:CG	22:5:315:CLA:HMD2	2.49	0.47
17:5:134:VAL:HG12	22:6:314:CLA:HED1	1.95	0.47
22:5:307:CLA:H52	22:5:307:CLA:H8	1.66	0.47
22:A:805:CLA:OBD	22:A:825:CLA:H101	2.14	0.47
3:C:66:ARG:HB3	3:C:68:TYR:CE1	2.48	0.47
5:E:50:SER:OG	5:E:88:PHE:HE1	1.97	0.47
7:G:112:LEU:O	7:G:116:PHE:HD1	1.97	0.47
12:Z:170:LEU:HA	12:Z:173:LYS:HD2	1.95	0.47
13:3:82:LYS:HD2	13:3:210:TYR:CD2	2.47	0.47
24:3:306:BCR:C21	22:3:318:CLA:H91	2.19	0.47
22:7:322:CLA:HMD1	22:8:307:CLA:H61	1.96	0.47
22:8:307:CLA:H62	22:8:307:CLA:CHC	2.44	0.47
18:6:216:PHE:CD1	22:6:322:CLA:HBB1	2.48	0.47
19:9:123:GLU:HG3	22:9:313:CLA:C4B	2.43	0.47
22:A:805:CLA:H151	24:A:844:BCR:HC31	1.96	0.47
2:B:31:ASP:HB2	22:B:831:CLA:CAA	2.45	0.47
2:B:153:ALA:HB2	22:B:813:CLA:HBC2	1.95	0.47
2:B:680:VAL:HG13	2:B:694:TRP:HZ2	1.79	0.47
2:B:705:GLN:HG3	31:B:852:DGD:HA52	1.95	0.47
22:B:806:CLA:H72	22:B:806:CLA:H112	1.77	0.47
24:B:853:BCR:H272	22:L:201:CLA:H92	1.95	0.47
7:G:102:THR:HG21	22:G:202:CLA:HBA1	1.96	0.47
9:J:8:LEU:HA	9:J:13:VAL:HG11	1.96	0.47
22:K:203:CLA:HBB1	22:K:203:CLA:HMB1	1.97	0.47
12:Z:112:PRO:HB2	12:Z:119:VAL:HG23	1.97	0.47
13:3:127:ARG:HG3	22:3:322:CLA:O1D	2.14	0.47
22:8:305:CLA:HMB1	22:8:305:CLA:CBB	2.38	0.47
16:4:176:GLN:NE2	16:4:181:THR:HA	2.29	0.47
17:5:55:PHE:CE1	22:5:315:CLA:HAA2	2.49	0.47
22:5:310:CLA:H172	22:5:311:CLA:HBB2	1.96	0.47
18:6:123:PHE:HZ	39:6:320:CHL:HBB2	1.76	0.47
22:A:821:CLA:H18	24:A:846:BCR:H272	1.96	0.47
22:A:855:CLA:HMC2	22:B:840:CLA:H11	1.96	0.47
24:A:856:BCR:H281	22:K:204:CLA:HMB3	1.95	0.47

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:B:330:SER:O	2:B:334:GLN:HG3	2.13	0.47
22:B:808:CLA:CBA	22:B:815:CLA:HBA1	2.38	0.47
4:D:192:MET:SD	4:D:192:MET:N	2.83	0.47
12:1:185:CYS:HA	12:1:188:PHE:CD2	2.49	0.47
38:Z:301:LUT:H162	22:Z:305:CLA:HMB3	1.96	0.47
38:Z:301:LUT:C28	22:Z:303:CLA:H52	2.45	0.47
13:3:131:ASP:O	13:3:135:PRO:HA	2.15	0.47
24:3:304:BCR:HC31	22:7:310:CLA:HAB	1.96	0.47
22:3:322:CLA:C4B	25:7:318:LHG:H101	2.44	0.47
22:7:306:CLA:H8	22:7:306:CLA:H51	1.48	0.47
25:4:820:LHG:H121	25:4:820:LHG:HC92	1.60	0.47
17:5:123:ALA:HB1	22:5:326:CLA:HAA1	1.95	0.47
24:5:304:BCR:C37	22:5:319:CLA:HMB2	2.45	0.47
22:5:315:CLA:H152	22:5:315:CLA:H18	1.76	0.47
18:6:73:VAL:HG21	18:6:128:ARG:NH1	2.30	0.47
18:6:123:PHE:HE1	24:6:305:BCR:H332	1.80	0.47
18:6:161:PRO:HD2	38:6:303:LUT:C23	2.44	0.47
18:6:251:GLN:HG2	18:6:254:TRP:CZ3	2.50	0.47
22:6:308:CLA:C1D	22:6:313:CLA:H72	2.44	0.47
22:6:308:CLA:HMD1	22:6:313:CLA:HBA2	1.95	0.47
22:6:321:CLA:HMB1	22:6:321:CLA:CBB	2.45	0.47
19:9:111:LEU:HD11	39:9:314:CHL:C2D	2.45	0.47
22:2:302:CLA:HED2	22:2:302:CLA:H2A	1.95	0.47
1:A:87:TRP:O	1:A:91:MET:HG2	2.15	0.47
1:A:264:PHE:HA	1:A:269:TRP:HD1	1.80	0.47
1:A:437:ILE:HG23	22:A:839:CLA:HBB2	1.95	0.47
21:A:801:CL0:H34	21:A:801:CL0:H41	1.52	0.47
22:A:807:CLA:CHC	26:A:851:NKP:HAUA	2.44	0.47
22:A:813:CLA:HMB1	22:A:813:CLA:CBB	2.40	0.47
2:B:18:THR:HG21	3:C:77:MET:HG3	1.96	0.47
2:B:30:HIS:CD2	22:B:808:CLA:HBB2	2.50	0.47
2:B:269:LEU:HD21	2:B:356:LEU:HB3	1.95	0.47
22:B:823:CLA:HBD	7:G:57:HIS:HD2	1.79	0.47
22:B:842:CLA:HAB	12:1:70:PHE:HZ	1.79	0.47
6:F:157:LEU:HB3	22:F:302:CLA:H142	1.97	0.47
7:G:59:ARG:HG2	7:G:63:ARG:NE	2.24	0.47
9:J:11:ALA:N	9:J:12:PRO:HD2	2.29	0.47
22:L:202:CLA:H2	22:L:202:CLA:H92	1.97	0.47
12:Z:144:GLU:HG3	22:Z:313:CLA:CBC	2.45	0.47
13:3:203:VAL:CG1	31:3:301:DGD:HG32	2.42	0.47
14:7:201:MET:HE3	22:7:307:CLA:HMC3	1.96	0.47

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
15:8:37:PRO:HG2	15:8:40:LEU:HD12	1.96	0.47
15:8:51:ASP:HA	38:8:303:LUT:H24	1.96	0.47
15:8:105:LYS:HB2	39:8:312:CHL:OBD	2.14	0.47
16:4:85:SER:HB2	16:4:88:VAL:HG23	1.95	0.47
16:4:192:TYR:HD1	22:4:805:CLA:O1D	1.98	0.47
22:5:311:CLA:O2D	22:5:311:CLA:HBA2	2.15	0.47
1:A:200:HIS:HB3	22:A:813:CLA:CAB	2.44	0.47
1:A:204:GLY:HA3	22:A:813:CLA:CBB	2.45	0.47
22:A:809:CLA:HAB	24:J:104:BCR:H383	1.97	0.47
22:A:829:CLA:HHC	22:A:829:CLA:CBB	2.45	0.47
22:A:832:CLA:HMB1	22:A:832:CLA:CBB	2.36	0.47
22:A:835:CLA:HMB1	22:A:835:CLA:CBB	2.43	0.47
22:A:836:CLA:HMB1	22:A:836:CLA:CBB	2.44	0.47
2:B:31:ASP:HB2	22:B:831:CLA:HAA1	1.97	0.47
2:B:92:ILE:HG23	22:B:812:CLA:CAD	2.45	0.47
2:B:218:PRO:HD2	22:B:817:CLA:CAD	2.45	0.47
2:B:262:PHE:HZ	2:B:356:LEU:HD23	1.79	0.47
2:B:429:PHE:HA	22:F:302:CLA:O1D	2.15	0.47
22:B:811:CLA:H11	22:B:829:CLA:O1A	2.13	0.47
22:B:813:CLA:HBA1	22:B:814:CLA:H11	1.97	0.47
22:B:821:CLA:HMB2	22:B:826:CLA:H92	1.96	0.47
22:B:824:CLA:HMC2	24:B:847:BCR:H10C	1.97	0.47
22:B:841:CLA:H91	22:B:841:CLA:H111	1.68	0.47
3:C:14:CYS:HB2	3:C:16:GLN:HG2	1.97	0.47
5:E:71:VAL:CG2	5:E:90:LEU:HD23	2.44	0.47
6:F:79:ARG:O	6:F:83:GLU:HB2	2.14	0.47
7:G:78:PHE:CD1	19:9:58:ALA:HB3	2.49	0.47
8:I:75:ALA:O	8:I:79:PHE:HD1	1.98	0.47
11:L:86:VAL:O	11:L:90:ARG:HG3	2.14	0.47
12:1:53:TYR:HB2	22:1:307:CLA:HMD1	1.97	0.47
12:Z:83:LEU:HD11	22:Z:303:CLA:HAC1	1.94	0.47
12:Z:166:ASP:O	12:Z:170:LEU:HD13	2.14	0.47
22:3:322:CLA:CBB	22:7:307:CLA:HMD2	2.44	0.47
14:7:189:TYR:HB3	22:7:304:CLA:CMA	2.40	0.47
38:7:301:LUT:H161	22:7:306:CLA:HMB3	1.96	0.47
24:7:303:BCR:C19	22:7:316:CLA:HMA2	2.45	0.47
39:7:313:CHL:HMC	22:7:316:CLA:HAB	1.95	0.47
15:8:72:HIS:HB3	15:8:196:MET:SD	2.55	0.47
22:5:307:CLA:HMD2	39:5:317:CHL:H11	1.97	0.47
22:5:311:CLA:C1B	22:5:318:CLA:H191	2.45	0.47
39:5:317:CHL:CHB	22:5:319:CLA:H71	2.44	0.47

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
18:6:125:GLU:HG2	22:6:317:CLA:CHB	2.45	0.47
22:9:311:CLA:NC	25:9:315:LHG:H271	2.30	0.47
22:A:809:CLA:CAB	24:J:104:BCR:H24C	2.45	0.47
2:B:301:ARG:HG3	7:G:65:ALA:HA	1.97	0.47
2:B:395:TRP:NE1	24:B:848:BCR:H272	2.29	0.47
7:G:58:ARG:NH2	7:G:96:GLY:O	2.46	0.47
22:1:309:CLA:HBB1	22:1:309:CLA:HMB1	1.97	0.47
13:3:186:LYS:HD3	22:3:314:CLA:O1D	2.15	0.47
14:7:131:PHE:HB3	22:7:315:CLA:HAB	1.96	0.47
15:8:80:VAL:HG13	15:8:202:PHE:CE2	2.46	0.47
38:8:303:LUT:C28	22:8:308:CLA:H72	2.45	0.47
22:8:305:CLA:H71	22:8:306:CLA:CMA	2.44	0.47
16:4:81:PHE:HB3	22:4:808:CLA:CAD	2.44	0.47
16:4:140:TYR:CD2	22:4:810:CLA:HED1	2.49	0.47
22:4:807:CLA:HMB1	22:4:807:CLA:CBB	2.38	0.47
22:5:301:CLA:H61	22:5:301:CLA:H2	1.68	0.47
38:5:302:LUT:H183	22:5:309:CLA:C3B	2.45	0.47
24:5:304:BCR:H17C	22:5:319:CLA:H2	1.97	0.47
18:6:61:PRO:O	18:6:65:LYS:HG3	2.15	0.47
18:6:84:GLY:O	18:6:88:GLN:HG3	2.15	0.47
18:6:160:ALA:HA	22:6:307:CLA:HBA2	1.97	0.47
1:A:59:PHE:CE2	22:A:805:CLA:HMC2	2.50	0.47
1:A:262:ALA:HB3	1:A:263:PRO:HD3	1.97	0.47
22:A:804:CLA:HAA1	22:A:811:CLA:H2	1.97	0.47
22:A:821:CLA:H72	24:A:846:BCR:C36	2.44	0.47
22:A:833:CLA:C1D	22:A:834:CLA:H121	2.45	0.47
2:B:219:HIS:CG	2:B:220:PRO:HD2	2.50	0.47
2:B:705:GLN:HG3	31:B:852:DGD:C5A	2.45	0.47
22:B:813:CLA:H42	22:9:307:CLA:H72	1.96	0.47
22:B:823:CLA:HED2	22:B:823:CLA:H2A	1.97	0.47
22:B:828:CLA:H143	22:B:828:CLA:H111	1.75	0.47
3:C:5:VAL:HG22	3:C:67:VAL:HG22	1.96	0.47
22:F:302:CLA:H162	32:F:306:RRX:H6	1.95	0.47
22:G:201:CLA:HHB	24:G:203:BCR:H363	1.96	0.47
12:Z:70:PHE:CB	22:Z:306:CLA:HMA1	2.45	0.47
15:8:77:MET:HE3	22:8:305:CLA:HMC3	1.97	0.47
15:8:97:VAL:HG13	22:8:310:CLA:O1D	2.14	0.47
38:4:803:LUT:H24	22:4:808:CLA:O1A	2.15	0.47
18:6:69:GLU:HG3	18:6:132:PHE:CD1	2.50	0.47
22:6:301:CLA:C1	22:6:301:CLA:H92	2.45	0.47
22:6:309:CLA:H121	25:6:323:LHG:H122	1.97	0.47

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
22:6:317:CLA:HMB1	22:6:317:CLA:CBB	2.37	0.47
19:9:49:PHE:HZ	19:9:160:LYS:HE2	1.80	0.47
19:9:87:GLN:HG2	22:9:308:CLA:C1D	2.45	0.47
22:2:301:CLA:HMA1	22:2:302:CLA:HAC2	1.97	0.47
1:A:410:ALA:O	1:A:414:VAL:HG23	2.15	0.47
22:A:817:CLA:H62	22:A:817:CLA:H41	1.51	0.47
2:B:208:VAL:HG22	2:B:216:GLN:HG2	1.97	0.47
22:B:830:CLA:H72	24:B:846:BCR:H372	1.96	0.47
12:1:157:PHE:HZ	22:1:313:CLA:NB	2.13	0.47
12:Z:195:THR:HG22	12:Z:197:LYS:H	1.80	0.47
38:Z:301:LUT:H26	22:Z:303:CLA:C2	2.44	0.47
14:7:55:PRO:HG2	38:7:302:LUT:H23	1.97	0.47
14:7:120:PHE:CE1	39:7:313:CHL:HMD2	2.49	0.47
42:7:321:SPH:H4	42:7:321:SPH:H71	1.72	0.47
15:8:50:PHE:HB3	22:8:308:CLA:CAD	2.45	0.47
15:8:124:PHE:CD1	39:8:301:CHL:H51	2.50	0.47
22:8:307:CLA:H141	22:8:311:CLA:HMC1	1.97	0.47
22:8:309:CLA:HBD	22:8:314:CLA:HBD	1.97	0.47
16:4:93:LEU:HD12	22:4:808:CLA:HMA2	1.96	0.47
17:5:188:LYS:HD3	22:5:313:CLA:O1D	2.15	0.47
22:5:320:CLA:HMC3	41:5:325:3PH:H341	1.97	0.47
39:6:315:CHL:HAB	39:6:318:CHL:HBB2	1.96	0.47
19:9:62:ARG:HB3	19:9:66:PHE:CZ	2.50	0.47
19:9:117:LEU:HD23	20:2:215:CYS:SG	2.55	0.47
20:2:214:LEU:HB3	39:2:303:CHL:HBA1	1.97	0.47
22:2:302:CLA:HHC	22:2:302:CLA:HBB1	1.96	0.47
1:A:301:ILE:HD12	1:A:305:PHE:CE2	2.50	0.46
22:A:814:CLA:HBA1	24:A:844:BCR:H341	1.97	0.46
22:A:825:CLA:HMB3	22:A:827:CLA:H93	1.97	0.46
22:A:827:CLA:HMA1	22:A:835:CLA:H62	1.96	0.46
2:B:187:SER:O	22:B:816:CLA:HBB2	2.15	0.46
2:B:495:LEU:O	2:B:495:LEU:HD22	2.15	0.46
22:B:823:CLA:CMB	33:1:301:LMT:H71	2.45	0.46
7:G:86:SER:HB3	19:9:36:ASN:HB2	1.97	0.46
10:K:35:ASN:HA	22:K:202:CLA:OBD	2.15	0.46
39:1:312:CHL:HMB1	39:1:312:CHL:CBB	2.38	0.46
22:Z:307:CLA:CBC	22:Z:314:CLA:HAC1	2.45	0.46
13:3:146:LEU:HD11	22:5:315:CLA:C17	2.42	0.46
13:3:149:VAL:HG21	17:5:57:PRO:O	2.15	0.46
13:3:149:VAL:HG23	13:3:162:PRO:HD2	1.96	0.46
24:3:304:BCR:H312	22:3:322:CLA:C1D	2.44	0.46

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
22:3:316:CLA:CHA	22:3:316:CLA:HBA1	2.45	0.46
42:7:321:SPH:H5	42:7:321:SPH:H2	1.58	0.46
16:4:198:ASP:HA	38:4:802:LUT:C24	2.31	0.46
38:4:802:LUT:H181	38:4:802:LUT:H8	1.97	0.46
24:4:804:BCR:H271	22:4:810:CLA:C4A	2.45	0.46
24:4:804:BCR:H271	22:4:810:CLA:NA	2.30	0.46
22:4:818:CLA:C2D	22:4:818:CLA:H42	2.45	0.46
22:5:326:CLA:HMD3	22:6:309:CLA:CHB	2.45	0.46
18:6:164:PRO:O	18:6:170:LEU:HD21	2.15	0.46
22:6:307:CLA:HMB1	22:6:307:CLA:CBB	2.45	0.46
22:6:314:CLA:NC	25:6:323:LHG:HC91	2.29	0.46
1:A:114:SER:CB	1:A:131:VAL:HG21	2.46	0.46
1:A:310:HIS:HB2	24:A:856:BCR:H19C	1.96	0.46
1:A:316:TRP:HH2	22:A:812:CLA:HHB	1.80	0.46
1:A:396:TRP:HB3	22:A:828:CLA:HMC3	1.97	0.46
1:A:528:LEU:HD11	22:A:837:CLA:CAD	2.46	0.46
22:A:807:CLA:HHC	22:A:807:CLA:CBB	2.46	0.46
2:B:96:HIS:CE1	22:B:812:CLA:HMB3	2.50	0.46
2:B:687:PRO:C	2:B:689:ALA:H	2.19	0.46
22:B:826:CLA:HMB1	22:B:826:CLA:HBB1	1.97	0.46
3:C:7:ILE:HG13	3:C:40:ALA:O	2.15	0.46
3:C:25:VAL:HG13	3:C:45:THR:HA	1.97	0.46
8:I:74:TRP:CZ2	22:2:304:CLA:HMA1	2.48	0.46
11:L:129:ALA:CB	11:L:175:THR:HG22	2.45	0.46
12:Z:85:ALA:HB1	38:Z:301:LUT:H12	1.96	0.46
12:Z:195:THR:HG22	12:Z:197:LYS:HE2	1.96	0.46
13:3:69:LEU:O	22:3:313:CLA:HMC3	2.15	0.46
13:3:228:ASN:HA	13:3:231:LYS:HG2	1.96	0.46
14:7:117:ASP:HB3	15:8:226:LYS:HD2	1.97	0.46
38:7:301:LUT:H11	22:7:305:CLA:HMC2	1.97	0.46
24:7:303:BCR:H271	22:7:309:CLA:C4A	2.44	0.46
22:8:307:CLA:H111	22:8:307:CLA:H152	1.66	0.46
22:8:308:CLA:HMB1	22:8:308:CLA:CBB	2.45	0.46
22:5:313:CLA:HHC	22:5:313:CLA:CBB	2.46	0.46
18:6:167:MET:SD	18:6:171:LYS:HE3	2.55	0.46
22:9:306:CLA:HMB1	22:9:306:CLA:CBB	2.35	0.46
1:A:366:ILE:HA	22:A:826:CLA:HED2	1.96	0.46
22:A:808:CLA:HAB	24:J:104:BCR:H352	1.98	0.46
22:A:826:CLA:HMB2	22:A:839:CLA:HBA1	1.97	0.46
2:B:196:VAL:HA	2:B:200:ILE:HD12	1.98	0.46
2:B:632:LEU:HD11	2:B:651:PHE:HB2	1.97	0.46

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:C:58:CYS:HA	29:C:102:SF4:S4	2.56	0.46
4:D:150:LEU:O	4:D:153:LYS:HE3	2.15	0.46
22:F:302:CLA:H61	32:F:306:RRX:H7	1.96	0.46
22:F:304:CLA:H72	22:F:304:CLA:C3D	2.44	0.46
10:K:38:MET:HB3	22:K:202:CLA:CAD	2.46	0.46
12:1:98:ASN:ND2	12:1:101:ASP:HB2	2.29	0.46
38:7:301:LUT:H382	22:7:304:CLA:C4D	2.44	0.46
25:7:318:LHG:H111	25:7:318:LHG:HC81	1.41	0.46
22:7:322:CLA:C2	22:7:323:CLA:H101	2.45	0.46
38:8:303:LUT:H162	22:8:310:CLA:HHB	1.96	0.46
16:4:237:LYS:HB3	16:4:241:GLN:OE1	2.15	0.46
17:5:131:TRP:HZ2	17:5:150:PRO:HG3	1.80	0.46
17:5:171:PRO:HD2	38:5:302:LUT:C24	2.45	0.46
17:5:248:PRO:O	39:5:316:CHL:H2	2.14	0.46
22:5:310:CLA:H92	22:5:310:CLA:H61	1.64	0.46
18:6:149:LEU:HD13	18:6:150:ALA:N	2.31	0.46
18:6:215:ILE:HG13	22:6:322:CLA:HBB2	1.97	0.46
22:6:322:CLA:HMB1	22:6:322:CLA:CBB	2.41	0.46
39:9:312:CHL:HAA2	39:9:312:CHL:CBD	2.46	0.46
22:2:301:CLA:HHC	22:2:301:CLA:HBB1	1.97	0.46
1:A:80:GLN:O	1:A:84:ILE:HG13	2.16	0.46
1:A:122:VAL:HB	22:B:834:CLA:HMD1	1.97	0.46
1:A:175:PHE:CD2	22:A:810:CLA:HAC2	2.50	0.46
1:A:289:LEU:HD21	1:A:374:MET:HB3	1.97	0.46
2:B:450:PRO:O	2:B:453:GLN:HG2	2.16	0.46
2:B:543:ARG:CB	2:B:552:LYS:HB3	2.39	0.46
2:B:601:THR:HG21	2:B:610:PHE:HB2	1.97	0.46
22:B:824:CLA:HHC	24:B:847:BCR:C10	2.45	0.46
22:B:842:CLA:HED2	24:B:847:BCR:H11C	1.97	0.46
7:G:98:THR:OG1	7:G:100:ILE:HG12	2.15	0.46
12:1:79:ARG:O	12:1:83:LEU:HG	2.15	0.46
12:Z:220:VAL:HG12	22:Z:316:CLA:H2A	1.97	0.46
22:Z:306:CLA:HMB1	22:Z:306:CLA:CBB	2.45	0.46
22:3:313:CLA:H121	22:5:320:CLA:C1C	2.45	0.46
24:7:303:BCR:H291	42:7:321:SPH:H142	1.98	0.46
39:8:312:CHL:HHD	39:8:315:CHL:HBC2	1.96	0.46
18:6:141:ASP:HA	39:6:320:CHL:C1C	2.46	0.46
19:9:106:VAL:HG13	19:9:111:LEU:HD12	1.98	0.46
1:A:150:ALA:HB2	1:A:378:PRO:CD	2.46	0.46
1:A:450:PHE:CD2	22:A:854:CLA:H12	2.51	0.46
1:A:493:LEU:HB3	1:A:496:GLN:NE2	2.30	0.46

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
22:A:803:CLA:H193	24:J:104:BCR:H17C	1.98	0.46
22:A:818:CLA:CMD	22:A:827:CLA:HMB2	2.44	0.46
2:B:256:LEU:HD11	22:B:817:CLA:HBC1	1.98	0.46
2:B:337:LEU:HD13	22:B:826:CLA:C3B	2.46	0.46
24:B:844:BCR:C36	24:G:203:BCR:H312	2.36	0.46
4:D:125:THR:HA	4:D:136:PRO:HG2	1.97	0.46
7:G:49:GLY:HA3	7:G:105:TRP:CE2	2.50	0.46
7:G:98:THR:O	7:G:102:THR:HG23	2.15	0.46
22:Z:307:CLA:OBD	22:Z:314:CLA:HAA1	2.15	0.46
39:Z:311:CHL:CMA	25:Z:317:LHG:H192	2.40	0.46
39:Z:312:CHL:HMC	22:Z:315:CLA:HAB	1.97	0.46
22:3:309:CLA:HMC3	22:3:314:CLA:H143	1.97	0.46
22:8:307:CLA:H18	22:8:311:CLA:HAC2	1.98	0.46
39:8:312:CHL:ND	26:8:318:NKP:HAL	2.30	0.46
16:4:230:ALA:CB	22:4:807:CLA:HAC2	2.46	0.46
22:5:307:CLA:CHB	22:5:307:CLA:H2	2.46	0.46
22:6:301:CLA:H41	22:6:301:CLA:H61	1.75	0.46
38:6:303:LUT:H35	38:6:303:LUT:H401	1.73	0.46
38:6:303:LUT:H30	22:6:307:CLA:H52	1.96	0.46
39:6:315:CHL:HBC3	39:6:318:CHL:CBC	2.45	0.46
22:6:322:CLA:HBC2	41:6:324:3PH:H281	1.98	0.46
22:2:301:CLA:HHC	22:2:301:CLA:CBB	2.46	0.46
1:A:119:TRP:HE3	22:A:809:CLA:H2A	1.81	0.46
1:A:526:ILE:CG2	1:A:613:LYS:HE3	2.46	0.46
1:A:694:GLN:O	1:A:698:GLU:HG3	2.14	0.46
2:B:263:HIS:CE1	2:B:265:GLN:HB3	2.48	0.46
2:B:692:VAL:HG13	11:L:135:TYR:CE1	2.50	0.46
22:B:841:CLA:H161	24:L:204:BCR:H15C	1.97	0.46
4:D:125:THR:O	4:D:129:ARG:HG3	2.16	0.46
10:K:66:LYS:O	10:K:68:VAL:HG23	2.16	0.46
12:1:153:PRO:HD2	22:1:304:CLA:OBD	2.16	0.46
40:1:318:SQD:H291	39:8:301:CHL:HMB2	1.98	0.46
38:3:303:LUT:H42	44:3:402:HOH:O	2.16	0.46
38:7:302:LUT:C16	22:7:309:CLA:HMB3	2.46	0.46
15:8:151:GLY:O	15:8:152:PHE:HB2	2.16	0.46
16:4:231:GLN:HB3	16:4:242:ASN:OD1	2.16	0.46
22:4:806:CLA:H12	22:4:806:CLA:H52	1.66	0.46
17:5:61:GLY:HA3	22:5:310:CLA:HBA1	1.96	0.46
17:5:164:TYR:CE1	17:5:183:GLN:HG2	2.50	0.46
22:5:315:CLA:H2	22:5:315:CLA:H61	1.55	0.46
18:6:66:TRP:NE1	22:6:311:CLA:H11	2.30	0.46

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
18:6:164:PRO:CG	22:6:308:CLA:H12	2.45	0.46
22:6:308:CLA:ND	22:6:313:CLA:H72	2.30	0.46
22:9:307:CLA:HMB1	22:9:307:CLA:CBB	2.43	0.46
1:A:589:TRP:NE1	22:A:830:CLA:HMD1	2.30	0.46
22:A:806:CLA:HBB2	24:A:845:BCR:HC31	1.96	0.46
22:A:821:CLA:H43	22:A:824:CLA:H12	1.98	0.46
22:A:825:CLA:H122	22:A:825:CLA:H8	1.65	0.46
22:A:828:CLA:H191	24:J:104:BCR:C19	2.45	0.46
2:B:190:ALA:CB	22:B:816:CLA:HAB	2.44	0.46
22:B:822:CLA:HBA2	22:B:822:CLA:H12	1.73	0.46
9:J:4:PHE:O	9:J:8:LEU:HG	2.16	0.46
14:7:53:PHE:HB3	22:7:307:CLA:C3D	2.45	0.46
39:7:313:CHL:HMC	22:7:316:CLA:CAB	2.46	0.46
22:8:310:CLA:HMB1	22:8:310:CLA:CBB	2.45	0.46
39:4:813:CHL:HHC	39:4:816:CHL:HAB	1.98	0.46
22:4:815:CLA:H2A	22:4:815:CLA:CED	2.43	0.46
22:5:307:CLA:HMB1	22:5:308:CLA:HBA2	1.96	0.46
18:6:173:LYS:HG3	22:6:313:CLA:HED2	1.97	0.46
22:6:314:CLA:H11	22:6:314:CLA:H51	1.75	0.46
22:2:302:CLA:HHC	22:2:302:CLA:CBB	2.46	0.46
1:A:87:TRP:HA	22:A:807:CLA:HBB2	1.97	0.46
1:A:687:PHE:HB2	22:B:801:CLA:HBC2	1.97	0.46
22:A:805:CLA:H202	22:A:805:CLA:H161	1.71	0.46
22:A:838:CLA:HMB2	22:A:839:CLA:C3D	2.45	0.46
22:A:839:CLA:H11	24:A:847:BCR:H15C	1.98	0.46
27:A:852:DGA:HAW1	27:A:852:DGA:HAN2	1.57	0.46
2:B:87:PRO:HB3	2:B:122:TYR:CG	2.51	0.46
2:B:296:PHE:HE2	22:B:822:CLA:HBD	1.79	0.46
2:B:358:PRO:HG3	22:B:820:CLA:HBA1	1.98	0.46
2:B:401:PRO:HD2	4:D:180:ASN:ND2	2.30	0.46
2:B:540:LEU:HD21	24:B:847:BCR:H282	1.97	0.46
2:B:577:PHE:CE2	22:B:831:CLA:HMD2	2.51	0.46
22:B:812:CLA:H8	22:B:812:CLA:H51	1.27	0.46
22:B:838:CLA:HMB2	22:B:839:CLA:C2D	2.45	0.46
4:D:131:LYS:HE3	4:D:132:PHE:CZ	2.51	0.46
7:G:99:LEU:HD21	19:9:32:LEU:HD21	1.98	0.46
10:K:107:LEU:CD2	22:K:203:CLA:HBB1	2.45	0.46
22:Z:304:CLA:C4D	22:Z:309:CLA:H62	2.45	0.46
31:3:301:DGD:O2G	31:3:301:DGD:HD1	2.16	0.46
16:4:154:MET:HA	16:4:154:MET:CE	2.46	0.46
17:5:73:SER:OG	22:5:311:CLA:HAA2	2.15	0.46

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
17:5:105:GLU:OE1	17:5:240:VAL:HG21	2.16	0.46
17:5:153:LYS:HA	17:5:153:LYS:HD3	1.69	0.46
36:5:306:C7Z:C28	22:5:323:CLA:HMD3	2.46	0.46
18:6:250:PHE:H	22:6:301:CLA:C4	2.27	0.46
18:6:250:PHE:N	22:6:301:CLA:H43	2.28	0.46
1:A:200:HIS:CD2	22:A:813:CLA:HAB	2.50	0.46
1:A:263:PRO:HB2	1:A:272:TYR:CZ	2.50	0.46
24:A:844:BCR:H15C	24:A:844:BCR:H351	1.39	0.46
22:B:808:CLA:H43	24:B:845:BCR:H24C	1.97	0.46
22:B:810:CLA:CAA	8:I:80:VAL:HG22	2.46	0.46
22:B:816:CLA:C6	24:B:846:BCR:H362	2.46	0.46
22:B:817:CLA:C4B	24:B:844:BCR:H271	2.46	0.46
22:B:831:CLA:HMD2	31:B:852:DGD:HB62	1.97	0.46
24:B:847:BCR:H15C	24:B:847:BCR:H351	1.75	0.46
6:F:218:LYS:HE2	6:F:218:LYS:HB2	1.63	0.46
7:G:92:ASN:HB3	22:G:202:CLA:OBD	2.16	0.46
12:1:61:CYS:SG	12:1:67:LEU:HA	2.56	0.46
25:Z:317:LHG:H121	25:Z:317:LHG:HC91	1.78	0.46
42:7:320:SPH:H11	42:7:320:SPH:H4	1.55	0.46
16:4:163:ARG:NE	39:4:819:CHL:OMC	2.49	0.46
17:5:57:PRO:HB2	17:5:58:LEU:HD12	1.97	0.46
22:5:326:CLA:HBA1	22:5:326:CLA:CBD	2.45	0.46
24:6:306:BCR:H19C	39:6:315:CHL:H2	1.98	0.46
22:6:311:CLA:HED3	22:6:317:CLA:CGA	2.46	0.46
19:9:51:PHE:HE1	22:9:311:CLA:HBC3	1.80	0.46
1:A:12:LYS:HE3	1:A:12:LYS:HB3	1.70	0.46
22:A:803:CLA:HMB2	25:A:850:LHG:H172	1.97	0.46
22:A:814:CLA:H121	22:A:816:CLA:CBB	2.44	0.46
22:A:823:CLA:H172	24:A:856:BCR:H403	1.98	0.46
2:B:32:PHE:CD2	22:B:808:CLA:HMC2	2.51	0.46
2:B:488:SER:HA	2:B:491:GLN:CD	2.35	0.46
2:B:653:PHE:HD1	22:B:806:CLA:H71	1.81	0.46
22:B:809:CLA:H71	22:B:809:CLA:H111	1.52	0.46
22:B:822:CLA:HMB1	22:B:822:CLA:CBB	2.46	0.46
24:B:853:BCR:H372	24:I:4001:BCR:H403	1.98	0.46
10:K:98:ILE:HG12	24:K:206:BCR:H311	1.98	0.46
22:1:306:CLA:C4B	22:1:306:CLA:H72	2.46	0.46
22:Z:307:CLA:HBC1	22:Z:314:CLA:HAC1	1.98	0.46
39:Z:311:CHL:C1C	25:Z:317:LHG:H102	2.46	0.46
13:3:182:LEU:HA	13:3:185:ILE:HG22	1.98	0.46
24:3:307:BCR:C19	22:3:310:CLA:H91	2.46	0.46

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
16:4:119:LEU:O	16:4:123:VAL:HG23	2.16	0.46
24:5:305:BCR:H343	18:6:220:VAL:HG21	1.97	0.46
22:5:315:CLA:HHC	22:5:315:CLA:HBB1	1.97	0.46
39:5:316:CHL:H152	22:6:322:CLA:HAA1	1.97	0.46
18:6:96:PHE:HZ	18:6:192:ALA:HB1	1.81	0.46
18:6:170:LEU:CB	22:6:307:CLA:HMA1	2.40	0.46
22:6:309:CLA:HBC2	22:6:319:CLA:HBD	1.98	0.46
39:6:315:CHL:HBC3	39:6:318:CHL:HBC3	1.97	0.46
19:9:142:PHE:HB3	19:9:143:PRO:CD	2.40	0.46
1:A:572:ARG:HH21	25:A:850:LHG:HC31	1.82	0.45
1:A:676:HIS:HB3	22:A:802:CLA:HBD	1.98	0.45
22:A:802:CLA:H202	22:A:830:CLA:H192	1.97	0.45
22:A:817:CLA:H122	22:A:817:CLA:H8	1.64	0.45
2:B:470:LYS:HG2	2:B:473:TYR:HE2	1.80	0.45
25:B:803:LHG:HC32	25:B:803:LHG:H262	1.98	0.45
22:B:821:CLA:H93	22:B:821:CLA:H111	1.63	0.45
6:F:199:GLY:O	6:F:202:TRP:HD1	1.99	0.45
22:F:304:CLA:H172	22:F:304:CLA:HAA1	1.98	0.45
8:I:95:LEU:CD1	24:L:204:BCR:HC8	2.46	0.45
12:Z:190:ALA:CB	22:Z:316:CLA:H61	2.41	0.45
22:7:324:CLA:H102	22:4:817:CLA:C4B	2.45	0.45
15:8:202:PHE:HD1	38:8:302:LUT:H41	1.81	0.45
16:4:66:LYS:CD	25:4:821:LHG:HC31	2.44	0.45
17:5:151:ILE:HD12	39:5:321:CHL:CMB	2.46	0.45
24:5:305:BCR:H351	39:5:316:CHL:HAC1	1.97	0.45
22:5:308:CLA:HMA2	22:5:308:CLA:HBA1	1.97	0.45
38:6:304:LUT:C16	22:6:312:CLA:HMB3	2.46	0.45
22:6:319:CLA:HHC	22:6:319:CLA:CBB	2.45	0.45
19:9:59:ASP:OD1	19:9:62:ARG:HB2	2.16	0.45
20:2:163:LEU:HB3	20:2:165:MET:HG2	1.99	0.45
1:A:578:PRO:HD3	2:B:562:GLY:HA2	1.97	0.45
1:A:740:TRP:CD1	22:A:828:CLA:HMB2	2.52	0.45
24:A:848:BCR:H23C	22:B:834:CLA:HMC2	1.98	0.45
27:A:852:DGA:HA61	14:7:242:PRO:HB2	1.97	0.45
22:B:809:CLA:HMB1	22:B:809:CLA:CBB	2.46	0.45
22:B:810:CLA:HMA1	22:B:811:CLA:HMB3	1.99	0.45
22:B:829:CLA:ND	22:B:829:CLA:H8	2.31	0.45
22:B:836:CLA:O2A	31:1:319:DGD:HA72	2.16	0.45
22:B:842:CLA:HHC	12:1:70:PHE:HZ	1.81	0.45
6:F:114:ARG:C	6:F:116:ALA:H	2.20	0.45
9:J:2:LYS:HE3	9:J:2:LYS:HB3	1.59	0.45

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
12:1:37:PRO:HG2	39:8:301:CHL:CMD	2.46	0.45
24:3:307:BCR:H11C	24:3:307:BCR:H341	1.67	0.45
22:3:308:CLA:CMD	39:3:317:CHL:HBA2	2.45	0.45
22:3:322:CLA:H92	22:7:306:CLA:H72	1.99	0.45
15:8:201:GLY:O	15:8:205:GLN:HG2	2.16	0.45
16:4:184:LYS:C	39:4:819:CHL:HMD2	2.37	0.45
18:6:126:LEU:CD1	22:6:302:CLA:HBA1	2.42	0.45
18:6:219:ALA:CB	18:6:229:PRO:HD3	2.47	0.45
22:6:307:CLA:H71	22:6:308:CLA:HMA1	1.97	0.45
20:2:223:ALA:HB3	20:2:226:SER:OG	2.16	0.45
1:A:516:ALA:HB2	1:A:622:VAL:HG21	1.98	0.45
22:A:812:CLA:C1C	22:A:812:CLA:H51	2.46	0.45
2:B:226:PHE:HA	2:B:231:TRP:CD1	2.51	0.45
2:B:348:LEU:HG	2:B:352:HIS:CE1	2.52	0.45
4:D:127:GLN:HG3	4:D:131:LYS:HE2	1.98	0.45
4:D:129:ARG:O	4:D:133:LYS:HA	2.16	0.45
4:D:141:VAL:HG22	4:D:147:VAL:HG22	1.98	0.45
13:3:192:MET:SD	22:3:311:CLA:HAB	2.57	0.45
24:3:307:BCR:H401	22:3:314:CLA:CHD	2.47	0.45
22:7:305:CLA:HMB3	22:7:310:CLA:H193	1.97	0.45
22:8:310:CLA:H62	22:8:310:CLA:H2	1.72	0.45
22:4:808:CLA:HBB1	22:4:808:CLA:HMB1	1.98	0.45
17:5:118:MET:HB3	24:5:305:BCR:H12C	1.98	0.45
17:5:131:TRP:HZ3	24:5:304:BCR:HC41	1.81	0.45
39:5:316:CHL:H13	22:6:322:CLA:HBD	1.98	0.45
39:6:315:CHL:HBD	39:6:315:CHL:HAA1	1.97	0.45
22:9:311:CLA:CGA	25:9:315:LHG:HC62	2.47	0.45
1:A:265:PHE:HA	22:K:202:CLA:HAC2	1.99	0.45
1:A:700:ILE:HG23	22:A:840:CLA:O1D	2.16	0.45
22:A:802:CLA:H51	2:B:439:VAL:HG13	1.98	0.45
22:A:833:CLA:H13	22:B:840:CLA:HBA1	1.98	0.45
22:A:834:CLA:HBC3	22:A:854:CLA:H13	1.97	0.45
2:B:293:ARG:HD2	2:B:301:ARG:HD3	1.98	0.45
2:B:321:LYS:HD2	2:B:322:GLY:N	2.31	0.45
25:B:803:LHG:H111	22:B:807:CLA:C1D	2.46	0.45
22:B:816:CLA:H61	22:B:816:CLA:H2	1.52	0.45
22:B:825:CLA:H13	24:B:847:BCR:H17C	1.99	0.45
22:B:827:CLA:CHB	22:B:839:CLA:HAA2	2.47	0.45
22:B:834:CLA:HBA2	22:B:834:CLA:CHA	2.43	0.45
3:C:10:THR:O	3:C:60:THR:HG21	2.17	0.45
6:F:125:CYS:HB3	6:F:131:PRO:HA	1.99	0.45

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
10:K:60:GLY:O	10:K:67:LEU:HG	2.16	0.45
12:Z:126:LEU:HD11	22:Z:315:CLA:HMD3	1.98	0.45
22:Z:306:CLA:HBC1	25:Z:317:LHG:H292	1.98	0.45
22:7:322:CLA:H72	22:7:323:CLA:H171	1.99	0.45
22:8:305:CLA:HBB1	22:8:306:CLA:HAA1	1.98	0.45
22:8:311:CLA:C2B	25:8:317:LHG:HC2	2.46	0.45
1:A:101:TYR:O	1:A:105:LEU:HG	2.16	0.45
1:A:113:PRO:O	1:A:139:GLN:HA	2.17	0.45
1:A:248:GLN:H	1:A:248:GLN:NE2	2.14	0.45
1:A:442:ASN:HD22	2:B:675:LEU:HD11	1.82	0.45
22:A:810:CLA:H121	22:A:812:CLA:HBC1	1.98	0.45
22:A:814:CLA:H92	27:A:852:DGA:HB81	1.99	0.45
22:A:815:CLA:C1B	24:A:856:BCR:HC42	2.47	0.45
22:A:829:CLA:H11	24:A:845:BCR:H321	1.98	0.45
2:B:228:THR:O	7:G:34:PRO:HG3	2.17	0.45
2:B:616:TYR:HD2	2:B:618:MET:HB3	1.81	0.45
22:B:806:CLA:C19	22:B:812:CLA:H52	2.46	0.45
6:F:151:ILE:HB	6:F:152:PRO:HD3	1.97	0.45
11:L:134:ILE:HG13	24:L:204:BCR:H363	1.97	0.45
12:1:72:GLU:O	12:1:76:ILE:HG12	2.17	0.45
12:1:138:GLU:HG3	22:1:314:CLA:C4B	2.47	0.45
12:1:153:PRO:HD3	22:1:313:CLA:HMD2	1.98	0.45
38:Z:302:LUT:C28	22:Z:306:CLA:H61	2.47	0.45
13:3:191:ALA:O	13:3:195:VAL:HG23	2.17	0.45
14:7:198:ARG:HA	14:7:201:MET:CE	2.47	0.45
24:4:804:BCR:H393	22:4:810:CLA:CHC	2.47	0.45
25:4:820:LHG:HC61	22:6:302:CLA:HAC2	1.99	0.45
18:6:66:TRP:HZ2	22:6:317:CLA:HAA2	1.80	0.45
18:6:101:PRO:HG3	39:6:318:CHL:OMC	2.16	0.45
19:9:57:GLY:HA3	19:9:63:LEU:HD13	1.98	0.45
38:9:302:LUT:C37	22:9:306:CLA:H112	2.47	0.45
20:2:197:ASN:CB	22:2:301:CLA:HBA2	2.47	0.45
20:2:216:VAL:O	20:2:220:ILE:HG13	2.16	0.45
1:A:49:ILE:HD12	1:A:49:ILE:N	2.32	0.45
1:A:403:VAL:HG11	1:A:596:LEU:CD2	2.46	0.45
22:A:822:CLA:H71	22:K:204:CLA:CAB	2.45	0.45
23:A:843:PQN:H262	23:A:843:PQN:H302	1.63	0.45
2:B:647:TRP:CZ2	2:B:727:ILE:HG21	2.52	0.45
22:B:818:CLA:H112	22:B:818:CLA:H93	1.69	0.45
22:B:819:CLA:CMD	22:B:828:CLA:HMB2	2.46	0.45
4:D:117:ARG:HB3	4:D:120:GLN:HG3	1.99	0.45

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
6:F:163:ILE:HG23	22:F:305:CLA:C1D	2.47	0.45
7:G:78:PHE:CE1	19:9:58:ALA:HB3	2.52	0.45
11:L:129:ALA:HB2	11:L:175:THR:HG22	1.97	0.45
12:Z:147:ALA:HA	12:Z:150:ARG:CD	2.47	0.45
22:3:313:CLA:H111	22:3:313:CLA:H152	1.68	0.45
14:7:118:TRP:CZ3	14:7:123:LEU:HA	2.52	0.45
22:8:307:CLA:NC	22:8:307:CLA:H51	2.32	0.45
17:5:60:LEU:HD12	38:5:303:LUT:H221	1.99	0.45
17:5:104:GLY:HA2	22:5:319:CLA:HAC2	1.98	0.45
18:6:127:LYS:O	18:6:138:VAL:HG21	2.16	0.45
22:6:302:CLA:C4D	24:6:305:BCR:HC41	2.47	0.45
22:6:311:CLA:HMA2	22:6:311:CLA:O1A	2.16	0.45
19:9:94:PRO:HB2	19:9:99:ALA:HA	1.98	0.45
1:A:175:PHE:CE2	22:A:810:CLA:HAC2	2.52	0.45
21:A:801:CL0:H72	21:A:801:CL0:H10	1.57	0.45
2:B:107:ARG:NH2	2:B:115:ASN:HA	2.32	0.45
2:B:227:TRP:CH2	22:B:817:CLA:H52	2.50	0.45
2:B:243:HIS:CD2	2:B:251:ALA:HA	2.52	0.45
2:B:423:LEU:HD13	2:B:533:LEU:HA	1.97	0.45
22:B:810:CLA:C1D	22:B:810:CLA:H51	2.46	0.45
22:B:824:CLA:NC	33:1:301:LMT:H101	2.32	0.45
22:B:825:CLA:H2	22:B:825:CLA:H61	1.76	0.45
22:B:825:CLA:O1A	24:B:847:BCR:H16C	2.16	0.45
22:B:835:CLA:CMB	24:B:848:BCR:H332	2.47	0.45
22:F:301:CLA:H152	9:J:18:TRP:CE3	2.46	0.45
13:3:164:PHE:HB3	22:3:308:CLA:OBD	2.16	0.45
41:7:319:3PH:H362	41:7:319:3PH:H231	1.98	0.45
15:8:101:TYR:CD1	15:8:102:GLU:HG3	2.52	0.45
16:4:156:LEU:O	24:4:804:BCR:H14C	2.17	0.45
22:4:805:CLA:H61	22:4:805:CLA:H92	1.74	0.45
17:5:60:LEU:HD11	22:5:315:CLA:C14	2.47	0.45
22:5:301:CLA:HBA1	22:5:320:CLA:HED3	1.97	0.45
38:5:303:LUT:H183	22:5:312:CLA:C3B	2.47	0.45
22:5:307:CLA:HMD2	39:5:317:CHL:O1A	2.17	0.45
18:6:97:TRP:HH2	18:6:189:THR:HG22	1.80	0.45
38:6:304:LUT:H162	22:6:312:CLA:HMB3	1.97	0.45
24:6:305:BCR:C27	22:6:312:CLA:H101	2.46	0.45
24:6:306:BCR:H11C	24:6:306:BCR:H341	1.80	0.45
19:9:65:TRP:HZ2	22:9:313:CLA:HAA1	1.81	0.45
19:9:106:VAL:HG22	19:9:111:LEU:HB2	1.98	0.45
19:9:120:GLY:CA	22:9:313:CLA:HAB	2.44	0.45

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:114:SER:HB3	1:A:131:VAL:HG21	1.99	0.45
1:A:222:LEU:N	1:A:223:PRO:HD2	2.32	0.45
1:A:571:PHE:HD2	22:A:830:CLA:HED1	1.82	0.45
22:A:805:CLA:HMC3	22:A:830:CLA:HMA1	1.98	0.45
22:A:829:CLA:H61	24:A:845:BCR:H342	1.97	0.45
22:A:841:CLA:HHC	22:A:841:CLA:CBB	2.47	0.45
22:A:855:CLA:HMA2	24:B:853:BCR:H322	1.99	0.45
2:B:192:THR:HG21	2:B:279:LEU:HB2	1.99	0.45
2:B:262:PHE:HE1	2:B:356:LEU:HA	1.82	0.45
22:B:821:CLA:H172	22:B:826:CLA:H203	1.99	0.45
22:B:825:CLA:H141	22:B:832:CLA:CBC	2.47	0.45
22:B:829:CLA:O1D	22:B:830:CLA:HHB	2.16	0.45
3:C:18:VAL:C	3:C:20:ALA:H	2.19	0.45
3:C:74:THR:HG23	3:C:79:LEU:HB2	1.98	0.45
5:E:39:ARG:HE	5:E:62:VAL:N	2.15	0.45
6:F:158:TYR:HA	6:F:202:TRP:HZ2	1.82	0.45
9:J:24:ALA:O	9:J:28:GLU:HG2	2.17	0.45
22:Z:306:CLA:H3A	22:Z:306:CLA:HBA2	1.15	0.45
13:3:157:ALA:O	13:3:159:PRO:HD3	2.17	0.45
14:7:122:THR:O	14:7:126:THR:HG23	2.17	0.45
15:8:50:PHE:CZ	41:8:320:3PH:H391	2.52	0.45
15:8:83:ILE:HG23	15:8:100:TRP:HB3	1.99	0.45
16:4:204:LYS:H	16:4:204:LYS:HG2	1.66	0.45
24:4:804:BCR:H282	22:4:818:CLA:HMC1	1.97	0.45
17:5:46:ASN:HB2	17:5:48:GLU:OE1	2.17	0.45
17:5:74:GLU:CD	17:5:187:ILE:HD11	2.38	0.45
24:5:304:BCR:H371	22:5:319:CLA:HMB2	1.99	0.45
22:5:307:CLA:H3A	22:5:307:CLA:HBA2	1.35	0.45
22:5:311:CLA:OBD	22:5:318:CLA:H12	2.17	0.45
22:9:313:CLA:CHA	22:9:313:CLA:HBA2	2.45	0.45
1:A:80:GLN:HE21	1:A:84:ILE:HG13	1.82	0.45
22:A:821:CLA:H101	24:A:847:BCR:C10	2.40	0.45
2:B:475:PHE:HB3	22:F:304:CLA:C19	2.44	0.45
2:B:548:MET:HG2	2:B:565:ARG:NH2	2.32	0.45
2:B:561:ASP:HB3	2:B:568:THR:HG21	1.99	0.45
2:B:681:TRP:CZ2	4:D:71:THR:HB	2.52	0.45
5:E:42:GLN:HA	5:E:57:GLY:O	2.16	0.45
14:7:53:PHE:HZ	41:7:319:3PH:H282	1.82	0.45
15:8:115:LEU:HD21	39:8:312:CHL:CHD	2.47	0.45
15:8:139:LYS:HB3	15:8:139:LYS:HE2	1.71	0.45
16:4:157:PHE:CD2	22:4:815:CLA:HMC3	2.52	0.45

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
17:5:169:PHE:HD1	39:5:317:CHL:O1A	1.98	0.45
22:5:308:CLA:C1C	22:5:313:CLA:H143	2.47	0.45
22:6:322:CLA:H2	22:6:322:CLA:H121	1.99	0.45
1:A:85:PHE:HE1	24:A:844:BCR:HC21	1.82	0.45
22:A:803:CLA:HMA2	22:A:808:CLA:H191	1.99	0.45
2:B:293:ARG:HH22	7:G:61:LEU:HB3	1.81	0.45
22:B:838:CLA:H152	32:F:306:RRX:H37	1.98	0.45
7:G:80:SER:O	7:G:83:GLN:HB3	2.17	0.45
22:G:201:CLA:HAB	24:G:203:BCR:H352	1.98	0.45
10:K:68:VAL:O	10:K:68:VAL:HG12	2.16	0.45
12:Z:147:ALA:HA	12:Z:150:ARG:HD2	1.99	0.45
38:Z:301:LUT:H15	38:Z:301:LUT:H201	1.79	0.45
13:3:164:PHE:HB3	22:3:308:CLA:HMD1	1.99	0.45
24:3:304:BCR:H341	24:3:304:BCR:H11C	1.34	0.45
14:7:58:LEU:HD11	22:7:308:CLA:H92	1.99	0.45
15:8:113:ILE:HG22	15:8:118:LEU:HB2	1.99	0.45
16:4:140:TYR:HD2	16:4:142:TYR:CE1	2.35	0.45
24:4:804:BCR:C37	39:4:816:CHL:HHB	2.44	0.45
17:5:69:TRP:CG	22:5:323:CLA:HAA1	2.52	0.45
22:5:326:CLA:C1	22:6:314:CLA:H152	2.46	0.45
39:9:312:CHL:H2A	39:9:312:CHL:C2	2.47	0.45
1:A:172:ALA:HA	22:3:312:CLA:H18	1.99	0.44
1:A:442:ASN:ND2	2:B:675:LEU:HD11	2.33	0.44
1:A:481:PRO:HG3	1:A:533:PHE:HB2	1.99	0.44
1:A:536:HIS:HA	1:A:539:HIS:CD2	2.52	0.44
22:A:807:CLA:HAB	24:J:104:BCR:C7	2.47	0.44
22:A:820:CLA:H62	22:A:820:CLA:H2	1.41	0.44
22:A:821:CLA:HMA2	22:A:825:CLA:NB	2.31	0.44
22:A:840:CLA:HBB2	6:F:167:GLY:HA3	1.99	0.44
27:A:852:DGA:HA22	27:A:852:DGA:HA51	1.65	0.44
2:B:181:SER:HB3	2:B:289:GLY:HA3	2.00	0.44
22:B:801:CLA:H122	22:B:801:CLA:H161	1.67	0.44
22:B:810:CLA:CMB	22:B:811:CLA:HMA1	2.41	0.44
22:B:811:CLA:C3D	22:B:829:CLA:HBA1	2.47	0.44
22:B:842:CLA:C4B	22:1:308:CLA:H51	2.47	0.44
22:B:842:CLA:HBC1	33:1:301:LMT:O6'	2.16	0.44
6:F:224:ILE:HB	6:F:227:ARG:NH2	2.30	0.44
7:G:94:PRO:O	7:G:95:ALA:HB3	2.17	0.44
11:L:128:LEU:HB3	24:L:204:BCR:H401	1.99	0.44
12:1:144:GLU:CB	22:1:313:CLA:HAC1	2.47	0.44
12:Z:145:THR:CA	12:Z:150:ARG:HH21	2.22	0.44

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
12:Z:150:ARG:O	22:Z:313:CLA:HMD3	2.17	0.44
13:3:126:LEU:HD23	22:3:322:CLA:HMA2	1.99	0.44
38:3:303:LUT:H32	22:3:311:CLA:CAB	2.47	0.44
22:8:313:CLA:HMA1	39:8:315:CHL:H52	1.98	0.44
38:4:803:LUT:H32	22:4:808:CLA:HAB	1.99	0.44
17:5:131:TRP:CZ3	22:6:314:CLA:HBA2	2.52	0.44
17:5:151:ILE:HD12	39:5:321:CHL:HMB1	2.00	0.44
41:5:325:3PH:H221	41:5:325:3PH:H2	1.51	0.44
19:9:145:ASP:N	19:9:146:PRO:HD2	2.27	0.44
19:9:187:LEU:HD13	38:9:301:LUT:H163	1.99	0.44
20:2:177:VAL:HG22	22:2:301:CLA:HMD3	1.99	0.44
1:A:53:HIS:CD2	22:A:804:CLA:HAC1	2.52	0.44
1:A:658:ILE:HD12	2:B:622:ARG:HG3	2.00	0.44
22:A:804:CLA:C4A	22:A:811:CLA:H71	2.48	0.44
22:A:816:CLA:O1A	27:A:852:DGA:HG12	2.16	0.44
22:A:817:CLA:C4D	22:A:817:CLA:H12	2.47	0.44
22:A:830:CLA:H151	22:A:841:CLA:CMA	2.48	0.44
22:A:832:CLA:H101	22:L:202:CLA:H11	2.00	0.44
22:A:836:CLA:H61	22:K:203:CLA:CBA	2.39	0.44
2:B:32:PHE:HA	2:B:35:HIS:CG	2.52	0.44
2:B:622:ARG:HA	2:B:626:TRP:HB3	1.98	0.44
22:B:811:CLA:H202	22:B:811:CLA:H162	1.64	0.44
22:B:819:CLA:H111	22:B:819:CLA:H71	1.25	0.44
22:B:824:CLA:H91	22:B:824:CLA:H111	1.75	0.44
6:F:186:ILE:HA	9:J:10:THR:HA	1.97	0.44
12:1:57:PRO:CD	38:1:303:LUT:H23	2.45	0.44
12:1:211:TRP:HZ3	15:8:118:LEU:HA	1.81	0.44
12:Z:68:LYS:HD2	12:Z:147:ALA:CB	2.44	0.44
22:3:312:CLA:HBD	22:3:318:CLA:HBD	1.99	0.44
14:7:157:THR:O	14:7:161:LYS:HD3	2.17	0.44
22:7:310:CLA:H122	22:7:310:CLA:C17	2.46	0.44
15:8:50:PHE:HB3	22:8:308:CLA:C3D	2.48	0.44
15:8:205:GLN:HG3	38:8:302:LUT:H42	1.99	0.44
22:4:809:CLA:H11	22:4:809:CLA:HBA1	1.62	0.44
25:4:821:LHG:H142	22:6:321:CLA:CHD	2.47	0.44
25:5:324:LHG:H111	25:5:324:LHG:HC82	1.50	0.44
22:A:809:CLA:HBA1	24:J:104:BCR:H363	1.99	0.44
22:A:827:CLA:C3B	22:A:835:CLA:HMA2	2.48	0.44
22:A:833:CLA:H41	22:A:855:CLA:HBC2	1.99	0.44
24:A:844:BCR:C20	24:A:845:BCR:H23C	2.47	0.44
24:A:844:BCR:H21C	24:A:845:BCR:H271	1.99	0.44

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
27:A:852:DGA:HG11	14:7:243:PHE:HB3	1.99	0.44
2:B:135:ASP:HB3	22:B:816:CLA:HED1	1.99	0.44
2:B:292:TYR:CD1	2:B:302:MET:HG2	2.52	0.44
2:B:649:TRP:CZ2	22:B:806:CLA:H72	2.52	0.44
22:B:807:CLA:HMB1	22:B:807:CLA:CBB	2.47	0.44
22:B:823:CLA:NA	22:B:824:CLA:H92	2.32	0.44
22:B:829:CLA:O1D	22:B:830:CLA:HMA1	2.17	0.44
22:B:839:CLA:H61	22:B:839:CLA:H2	1.67	0.44
33:G:204:LMT:H91	33:G:204:LMT:H62	1.79	0.44
12:Z:193:ALA:HB3	22:Z:316:CLA:HED1	1.99	0.44
22:Z:303:CLA:H62	22:Z:304:CLA:CMA	2.47	0.44
22:Z:309:CLA:HBB2	24:4:804:BCR:H323	1.99	0.44
13:3:71:ALA:CB	38:3:302:LUT:H12	2.47	0.44
13:3:146:LEU:O	13:3:149:VAL:N	2.46	0.44
24:3:307:BCR:H342	22:3:315:CLA:HAA1	1.98	0.44
39:3:317:CHL:H61	39:3:317:CHL:H101	1.65	0.44
14:7:141:ASP:HB2	14:7:148:GLN:HG2	1.98	0.44
22:7:310:CLA:HBC1	25:7:318:LHG:H281	1.99	0.44
15:8:51:ASP:OD2	22:8:308:CLA:HBA1	2.18	0.44
22:8:310:CLA:H72	22:8:310:CLA:H112	1.31	0.44
16:4:83:PRO:HD2	38:4:803:LUT:C23	2.42	0.44
16:4:193:PRO:HB2	16:4:197:PHE:HB2	2.00	0.44
22:4:805:CLA:HBA2	22:4:805:CLA:H3A	1.29	0.44
39:4:813:CHL:CAB	39:4:816:CHL:HBB2	2.41	0.44
17:5:230:ASN:HA	17:5:233:HIS:HE2	1.82	0.44
38:5:303:LUT:H15	38:5:303:LUT:H201	1.85	0.44
19:9:31:TRP:NE1	19:9:51:PHE:HA	2.32	0.44
1:A:212:SER:HA	22:A:815:CLA:HBB2	1.98	0.44
1:A:269:TRP:HB3	22:A:817:CLA:HHB	1.98	0.44
1:A:369:HIS:HA	1:A:372:TYR:CD2	2.53	0.44
1:A:412:PHE:CD1	1:A:416:ASP:HB2	2.53	0.44
22:A:821:CLA:NC	22:A:827:CLA:H161	2.32	0.44
2:B:52:PHE:HD1	2:B:150:PHE:HD1	1.64	0.44
2:B:287:VAL:HB	22:B:818:CLA:H202	1.98	0.44
25:B:803:LHG:H262	22:B:807:CLA:HAA2	2.00	0.44
22:B:835:CLA:HBA2	22:B:835:CLA:H3A	1.25	0.44
24:B:848:BCR:C8	24:B:848:BCR:H321	2.47	0.44
5:E:70:PRO:HD2	5:E:89:ALA:HA	1.97	0.44
10:K:62:ASP:HB2	10:K:68:VAL:HG21	1.99	0.44
12:Z:160:LEU:HB2	38:Z:301:LUT:H222	1.99	0.44
13:3:127:ARG:HD3	39:3:317:CHL:CBB	2.47	0.44

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
15:8:105:LYS:NZ	15:8:105:LYS:HB3	2.32	0.44
22:8:308:CLA:H161	22:8:309:CLA:HMB1	1.99	0.44
22:4:818:CLA:HBA1	22:4:818:CLA:C2	2.45	0.44
17:5:100:TRP:HD1	38:5:303:LUT:H41	1.82	0.44
17:5:245:LEU:HD22	18:6:220:VAL:HG11	1.99	0.44
22:5:307:CLA:H43	22:5:308:CLA:H2	1.99	0.44
18:6:57:LEU:CD1	38:6:304:LUT:H221	2.47	0.44
22:A:810:CLA:O1A	22:A:812:CLA:HMD2	2.17	0.44
22:A:815:CLA:H143	13:3:196:PHE:HA	1.98	0.44
22:A:840:CLA:C4B	22:F:301:CLA:HMD2	2.48	0.44
27:A:852:DGA:HA22	27:A:852:DGA:HG12	1.72	0.44
2:B:373:TYR:CE2	2:B:377:GLN:HG3	2.52	0.44
22:B:816:CLA:HMA1	24:B:846:BCR:H402	1.99	0.44
7:G:97:PHE:CE2	22:G:202:CLA:HMB2	2.52	0.44
22:1:308:CLA:C1C	22:1:314:CLA:H18	2.48	0.44
39:1:312:CHL:CAA	31:1:319:DGD:HG31	2.42	0.44
25:1:317:LHG:C10	24:8:304:BCR:H313	2.43	0.44
12:Z:79:ARG:O	12:Z:83:LEU:HG	2.17	0.44
12:Z:190:ALA:CB	22:Z:305:CLA:HAC2	2.47	0.44
14:7:75:ASN:ND2	22:7:315:CLA:HMD1	2.32	0.44
15:8:114:ALA:HB3	15:8:117:PRO:HD2	2.00	0.44
15:8:126:CYS:HB2	22:8:314:CLA:HAB	1.99	0.44
38:5:302:LUT:H401	38:5:302:LUT:H35	1.77	0.44
18:6:253:LEU:HA	41:6:324:3PH:H292	1.98	0.44
22:6:311:CLA:HMD2	22:6:317:CLA:C1D	2.47	0.44
38:9:302:LUT:H28	22:9:306:CLA:C5	2.40	0.44
22:9:306:CLA:H111	22:9:306:CLA:H72	1.27	0.44
20:2:182:PRO:HG2	20:2:183:ILE:H	1.81	0.44
1:A:71:ARG:HD3	1:A:185:ALA:HB1	1.99	0.44
22:A:812:CLA:H91	22:A:812:CLA:H111	1.71	0.44
22:A:814:CLA:H41	22:A:814:CLA:H61	1.65	0.44
22:A:832:CLA:H13	22:A:832:CLA:H91	2.00	0.44
24:A:844:BCR:C21	24:A:845:BCR:H291	2.47	0.44
2:B:423:LEU:HB3	2:B:533:LEU:HD13	1.98	0.44
2:B:429:PHE:CE2	22:B:838:CLA:HAB	2.52	0.44
2:B:473:TYR:HB3	2:B:475:PHE:CE2	2.53	0.44
22:B:818:CLA:H72	22:B:818:CLA:H122	1.99	0.44
22:B:818:CLA:HBA2	22:B:818:CLA:H3A	1.53	0.44
3:C:49:VAL:HG11	4:D:117:ARG:HH21	1.82	0.44
4:D:115:PHE:HB3	4:D:120:GLN:NE2	2.29	0.44
22:F:304:CLA:H92	22:F:304:CLA:H61	1.57	0.44

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
22:G:201:CLA:C1C	24:G:203:BCR:H10C	2.48	0.44
10:K:107:LEU:HD22	10:K:112:ASN:CB	2.47	0.44
22:Z:307:CLA:H12	22:Z:307:CLA:H51	1.70	0.44
24:3:307:BCR:C39	22:3:314:CLA:HAC2	2.46	0.44
22:7:309:CLA:C2	22:7:309:CLA:H72	2.44	0.44
22:7:310:CLA:CBC	25:7:318:LHG:H281	2.48	0.44
22:8:311:CLA:C4A	33:4:801:LMT:H62	2.47	0.44
16:4:113:GLY:O	38:4:803:LUT:H42	2.17	0.44
16:4:121:ARG:NH2	16:4:133:PRO:HA	2.28	0.44
16:4:164:ARG:HH12	22:4:815:CLA:CED	2.31	0.44
25:4:820:LHG:H222	22:6:321:CLA:HMA2	1.99	0.44
22:5:310:CLA:C2B	22:5:310:CLA:H51	2.48	0.44
22:6:309:CLA:HBC1	22:6:319:CLA:CAA	2.46	0.44
22:6:312:CLA:H52	22:6:312:CLA:C2D	2.48	0.44
19:9:102:LYS:HD2	22:9:308:CLA:HED1	2.00	0.44
1:A:511:GLY:HA2	1:A:525:PRO:HB3	1.99	0.44
1:A:606:ALA:HB2	22:A:837:CLA:HBB2	1.99	0.44
21:A:801:CL0:CAC	2:B:626:TRP:HD1	2.31	0.44
22:A:812:CLA:C4	13:3:41:LEU:HD21	2.48	0.44
22:A:830:CLA:H111	22:A:841:CLA:HMA2	1.99	0.44
22:B:840:CLA:C2	22:B:841:CLA:H143	2.47	0.44
24:B:844:BCR:H341	24:B:844:BCR:H11C	1.52	0.44
3:C:28:MET:CE	3:C:38:GLN:HB3	2.48	0.44
22:F:302:CLA:H62	22:F:302:CLA:H2	1.48	0.44
22:F:304:CLA:H111	22:F:304:CLA:H152	1.71	0.44
13:3:203:VAL:HG11	31:3:301:DGD:C7A	2.48	0.44
14:7:187:LYS:HD3	14:7:190:LYS:HD3	1.98	0.44
22:7:309:CLA:C3C	42:7:321:SPH:H162	2.48	0.44
41:7:319:3PH:H372	41:7:319:3PH:H3A1	1.41	0.44
22:8:309:CLA:H93	22:8:309:CLA:H111	1.71	0.44
22:5:326:CLA:HMB3	18:6:210:PRO:CB	2.47	0.44
18:6:235:GLN:NE2	22:6:301:CLA:HHB	2.32	0.44
41:6:324:3PH:H221	41:6:324:3PH:H251	1.52	0.44
1:A:399:GLY:HA3	1:A:603:LEU:HD21	2.00	0.44
22:A:807:CLA:H2	22:A:807:CLA:H61	1.62	0.44
2:B:183:LEU:CD1	22:B:815:CLA:H43	2.47	0.44
2:B:695:LYS:HE2	11:L:141:GLN:NE2	2.32	0.44
22:B:821:CLA:O1A	22:B:826:CLA:H142	2.17	0.44
22:B:830:CLA:H193	24:B:846:BCR:H17C	1.99	0.44
22:B:842:CLA:HBB1	22:B:842:CLA:HMB1	2.00	0.44
22:L:201:CLA:HBA2	22:L:201:CLA:HBD	1.99	0.44

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
22:L:202:CLA:CBC	24:L:204:BCR:H23C	2.48	0.44
12:1:209:ASN:ND2	12:1:212:GLY:HA3	2.33	0.44
38:1:303:LUT:H11	38:1:303:LUT:H191	1.88	0.44
38:1:303:LUT:H382	22:1:307:CLA:O2A	2.17	0.44
22:1:308:CLA:CMD	22:1:314:CLA:ND	2.79	0.44
12:Z:149:LYS:CB	22:Z:313:CLA:HBC1	2.40	0.44
12:Z:177:ASN:HD22	12:Z:177:ASN:H	1.65	0.44
13:3:149:VAL:HG21	13:3:163:TRP:HB2	2.00	0.44
14:7:135:GLU:HG3	22:7:315:CLA:C4B	2.48	0.44
38:7:301:LUT:C14	22:7:305:CLA:HAB	2.48	0.44
15:8:115:LEU:HD22	15:8:115:LEU:HA	1.74	0.44
15:8:141:GLY:HA2	15:8:157:LYS:O	2.18	0.44
39:8:315:CHL:H61	39:8:315:CHL:H111	1.98	0.44
38:4:803:LUT:H28	22:4:808:CLA:H72	2.00	0.44
22:4:807:CLA:H152	22:4:807:CLA:H111	1.67	0.44
17:5:155:ASN:HB3	39:5:321:CHL:C2D	2.47	0.44
17:5:179:ARG:O	17:5:183:GLN:HG3	2.17	0.44
39:6:320:CHL:HHC	39:6:320:CHL:CBB	2.46	0.44
22:A:826:CLA:HMA1	24:A:847:BCR:H15C	1.99	0.44
2:B:258:PHE:CD1	2:B:494:TRP:HB3	2.53	0.44
22:B:808:CLA:H122	22:B:808:CLA:HBD	1.99	0.44
22:B:817:CLA:H3A	22:B:817:CLA:HBA2	1.21	0.44
6:F:222:ILE:O	6:F:222:ILE:HD12	2.18	0.44
9:J:1:MET:H1	34:J:101:LMG:HC71	1.83	0.44
38:1:302:LUT:C30	22:1:304:CLA:H8	2.47	0.44
22:1:313:CLA:HMB1	22:1:313:CLA:CBB	2.48	0.44
12:Z:114:TRP:HB2	22:Z:308:CLA:CGA	2.47	0.44
38:Z:302:LUT:H8	22:Z:308:CLA:HBB1	2.00	0.44
13:3:42:LEU:HD12	13:3:53:PRO:HA	2.00	0.44
13:3:81:GLY:HA3	13:3:88:ASP:HA	2.00	0.44
38:3:303:LUT:C16	22:3:313:CLA:HMB3	2.47	0.44
22:3:313:CLA:H112	22:5:320:CLA:ND	2.33	0.44
22:7:305:CLA:HAC2	22:7:306:CLA:H191	2.00	0.44
22:8:313:CLA:HMB1	22:8:313:CLA:CBB	2.47	0.44
39:4:816:CHL:HAC2	39:4:816:CHL:OMC	2.18	0.44
17:5:106:PRO:C	17:5:115:ASN:H	2.21	0.44
17:5:179:ARG:HG2	17:5:183:GLN:HG3	2.00	0.44
24:5:304:BCR:HC32	22:6:314:CLA:C1D	2.48	0.44
22:5:326:CLA:HAC2	22:6:309:CLA:HMA1	2.00	0.44
19:9:57:GLY:N	22:9:306:CLA:O1A	2.51	0.44
1:A:316:TRP:CZ3	22:A:820:CLA:HBD	2.52	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:693:TRP:HZ3	22:B:801:CLA:O1D	2.01	0.43
22:A:818:CLA:H62	22:A:818:CLA:NC	2.32	0.43
22:A:823:CLA:HBA2	37:K:201:DAO:H81	2.00	0.43
2:B:231:TRP:HB2	22:B:818:CLA:O1A	2.18	0.43
2:B:320:HIS:HA	2:B:323:LEU:HD12	2.00	0.43
2:B:351:GLN:HG3	22:B:827:CLA:CED	2.44	0.43
2:B:388:PHE:CZ	24:B:848:BCR:H20C	2.53	0.43
2:B:449:THR:HB	2:B:452:LYS:HG3	1.99	0.43
2:B:616:TYR:CD2	2:B:618:MET:HB3	2.52	0.43
22:B:817:CLA:NC	24:B:844:BCR:H291	2.33	0.43
22:B:829:CLA:H92	22:B:829:CLA:H62	1.55	0.43
22:B:834:CLA:C2	24:F:303:BCR:HC42	2.47	0.43
7:G:50:ARG:HG2	22:G:202:CLA:C3B	2.48	0.43
11:L:96:LEU:O	11:L:99:PRO:HD2	2.18	0.43
22:1:309:CLA:HMB1	22:1:309:CLA:CBB	2.48	0.43
12:Z:179:ARG:HA	12:Z:182:MET:CE	2.48	0.43
39:Z:311:CHL:HBC3	39:Z:311:CHL:CHD	2.44	0.43
13:3:115:PHE:O	13:3:119:ILE:HG12	2.18	0.43
38:8:302:LUT:C33	22:8:306:CLA:HMB2	2.48	0.43
22:8:306:CLA:CMD	22:8:311:CLA:C4D	2.96	0.43
16:4:259:ASP:HA	16:4:262:ARG:HD3	1.99	0.43
22:5:311:CLA:CMD	22:6:322:CLA:H92	2.47	0.43
22:5:315:CLA:HHC	22:5:315:CLA:CBB	2.48	0.43
22:6:302:CLA:HHC	22:6:302:CLA:CBB	2.48	0.43
1:A:32:PRO:HB2	1:A:48:TRP:CH2	2.53	0.43
1:A:127:LEU:HD22	1:A:667:ALA:HB2	2.00	0.43
1:A:227:LEU:HD12	1:A:237:ILE:HG23	2.00	0.43
1:A:338:HIS:CE1	25:A:849:LHG:HC12	2.36	0.43
1:A:392:THR:HG23	1:A:607:ILE:HG21	2.00	0.43
22:A:817:CLA:H61	22:A:817:CLA:C4C	2.48	0.43
22:A:821:CLA:H11	22:A:827:CLA:H122	2.00	0.43
23:A:843:PQN:H142	24:F:303:BCR:H291	2.00	0.43
2:B:5:LEU:HB3	2:B:8:LYS:HE3	2.00	0.43
2:B:520:VAL:O	2:B:524:ILE:HG13	2.18	0.43
22:B:821:CLA:CMB	22:B:826:CLA:HMA3	2.47	0.43
22:B:833:CLA:H18	22:B:833:CLA:H152	1.74	0.43
33:G:204:LMT:H52	22:1:308:CLA:HMD1	2.00	0.43
9:J:1:MET:N	34:J:101:LMG:HC71	2.32	0.43
12:1:57:PRO:HD2	38:1:303:LUT:C23	2.46	0.43
12:1:70:PHE:HB3	22:1:307:CLA:HMA1	1.99	0.43
22:Z:316:CLA:C2	22:Z:316:CLA:H71	2.42	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
13:3:103:TYR:CE2	13:3:105:LYS:HB2	2.53	0.43
14:7:70:HIS:CE1	14:7:138:ARG:HH11	2.35	0.43
22:8:311:CLA:HMC2	33:4:801:LMT:H121	2.00	0.43
38:4:803:LUT:H28	22:4:808:CLA:H52	2.00	0.43
22:4:808:CLA:HMB1	22:4:808:CLA:CBB	2.49	0.43
24:5:304:BCR:HC42	25:6:323:LHG:HC81	2.00	0.43
22:5:310:CLA:HMD3	22:5:315:CLA:HMD1	2.00	0.43
22:5:311:CLA:HMB1	22:5:311:CLA:HBB1	1.99	0.43
22:5:318:CLA:H12	22:5:318:CLA:H51	1.80	0.43
22:5:322:CLA:HED1	25:6:323:LHG:H301	2.00	0.43
39:6:316:CHL:HBD	39:6:316:CHL:HAA1	2.00	0.43
19:9:35:LEU:HG	19:9:36:ASN:N	2.32	0.43
19:9:129:GLY:HA3	19:9:142:PHE:CE1	2.53	0.43
22:9:307:CLA:HMD2	22:9:313:CLA:CHD	2.48	0.43
1:A:521:VAL:HG21	1:A:627:VAL:HG11	1.99	0.43
1:A:550:LEU:CB	24:A:847:BCR:H322	2.48	0.43
22:A:835:CLA:H51	22:A:835:CLA:H12	1.76	0.43
23:A:843:PQN:H111	23:A:843:PQN:H2M1	1.77	0.43
2:B:16:ASP:HB3	2:B:21:ARG:HB3	1.98	0.43
2:B:57:ILE:HD13	22:B:809:CLA:HMD2	2.00	0.43
22:B:812:CLA:NB	24:B:849:BCR:H322	2.34	0.43
22:B:817:CLA:H51	22:B:817:CLA:H11	1.76	0.43
22:B:821:CLA:HAA2	22:B:825:CLA:OBD	2.17	0.43
22:B:825:CLA:H111	24:B:847:BCR:C16	2.49	0.43
22:B:838:CLA:HHC	22:B:838:CLA:H102	2.01	0.43
3:C:66:ARG:HD3	3:C:66:ARG:HA	1.84	0.43
12:1:55:PHE:HZ	40:1:318:SQD:H271	1.83	0.43
12:1:69:ARG:NH2	22:1:308:CLA:O1D	2.51	0.43
12:Z:82:MET:SD	22:Z:303:CLA:HAB	2.58	0.43
12:Z:211:TRP:CE3	22:Z:310:CLA:HMA1	2.54	0.43
38:Z:301:LUT:H362	22:Z:313:CLA:H42	1.99	0.43
22:7:324:CLA:H143	22:4:817:CLA:ND	2.32	0.43
15:8:121:VAL:O	15:8:125:MET:HG3	2.18	0.43
24:8:304:BCR:H19C	39:8:315:CHL:CGA	2.49	0.43
22:8:316:CLA:CBB	22:8:316:CLA:HHC	2.48	0.43
22:8:316:CLA:HHC	22:8:316:CLA:HBB1	2.00	0.43
17:5:230:ASN:HA	17:5:233:HIS:CE1	2.53	0.43
18:6:215:ILE:O	18:6:218:LYS:HG2	2.18	0.43
18:6:240:GLN:HB2	24:6:306:BCR:H313	1.99	0.43
24:6:305:BCR:H20C	22:6:312:CLA:H203	1.99	0.43
19:9:30:THR:HB	19:9:35:LEU:HB3	2.00	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:412:PHE:HD1	1:A:416:ASP:HB2	1.83	0.43
1:A:675:ALA:HB1	1:A:734:GLY:O	2.18	0.43
22:A:823:CLA:H191	10:K:98:ILE:CD1	2.46	0.43
22:A:827:CLA:CBB	22:A:835:CLA:HMA2	2.48	0.43
22:A:828:CLA:C9	24:J:104:BCR:H341	2.48	0.43
2:B:331:LEU:HD22	22:B:808:CLA:HMD2	2.01	0.43
22:B:823:CLA:NB	22:B:824:CLA:H8	2.33	0.43
3:C:14:CYS:SG	29:C:102:SF4:S4	3.17	0.43
4:D:160:LYS:HB3	4:D:160:LYS:HE2	1.78	0.43
5:E:78:GLN:CB	5:E:84:THR:HG22	2.38	0.43
6:F:137:PRO:HG2	22:F:304:CLA:C2C	2.49	0.43
12:1:143:ASP:O	12:1:145:THR:HG23	2.18	0.43
38:1:302:LUT:H35	38:1:302:LUT:H401	1.84	0.43
38:1:302:LUT:C33	22:1:305:CLA:HMB2	2.48	0.43
22:Z:307:CLA:HMB1	22:Z:307:CLA:CBB	2.48	0.43
24:3:307:BCR:H312	22:3:315:CLA:HMB3	2.00	0.43
39:3:317:CHL:HAA1	39:3:317:CHL:HBD	1.99	0.43
38:7:301:LUT:H35	38:7:301:LUT:H401	1.83	0.43
15:8:228:THR:HA	22:8:307:CLA:CBA	2.48	0.43
22:8:308:CLA:C17	22:8:308:CLA:H122	2.48	0.43
16:4:101:LEU:HD12	16:4:101:LEU:HA	1.82	0.43
17:5:135:ARG:HB3	39:5:317:CHL:CMC	2.48	0.43
17:5:178:ASN:ND2	17:5:181:GLU:HB2	2.32	0.43
38:5:303:LUT:H31	38:5:303:LUT:H391	1.91	0.43
22:5:318:CLA:H2	22:5:323:CLA:HAC1	1.99	0.43
18:6:235:GLN:HG2	18:6:245:PHE:CE1	2.53	0.43
22:6:309:CLA:H203	22:6:309:CLA:H152	2.01	0.43
22:9:305:CLA:H92	22:9:305:CLA:H61	1.68	0.43
1:A:26:PHE:H	34:J:101:LMG:HC5	1.83	0.43
22:A:804:CLA:HED1	22:A:811:CLA:HED2	2.00	0.43
22:A:807:CLA:ND	26:A:851:NKP:HARA	2.33	0.43
22:A:810:CLA:HAA1	13:3:51:VAL:HG21	2.01	0.43
22:A:814:CLA:H61	27:A:852:DGA:HB62	1.99	0.43
22:A:841:CLA:CAB	22:B:801:CLA:H162	2.48	0.43
22:A:842:CLA:H62	22:A:842:CLA:H2	1.65	0.43
2:B:258:PHE:HE1	22:B:819:CLA:H52	1.83	0.43
25:B:803:LHG:H122	22:B:807:CLA:ND	2.32	0.43
22:B:807:CLA:HAB	22:B:809:CLA:CAD	2.48	0.43
22:B:817:CLA:H111	24:B:844:BCR:H23C	2.01	0.43
4:D:191:THR:O	4:D:194:ILE:HG22	2.19	0.43
12:Z:164:LYS:HB2	12:Z:164:LYS:NZ	2.34	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
38:Z:301:LUT:C16	22:Z:305:CLA:HMB3	2.49	0.43
14:7:80:MET:SD	22:7:304:CLA:HAB	2.58	0.43
15:8:82:GLY:O	15:8:86:PRO:HG2	2.19	0.43
16:4:193:PRO:HD2	22:4:805:CLA:OBD	2.18	0.43
17:5:131:TRP:HB2	22:6:314:CLA:H2	2.00	0.43
22:5:309:CLA:H12	22:5:309:CLA:C1A	2.49	0.43
22:5:318:CLA:HMA1	22:5:322:CLA:HBC2	2.00	0.43
22:6:312:CLA:C4D	22:6:312:CLA:H52	2.48	0.43
39:6:315:CHL:HBB2	39:6:318:CHL:HBB2	2.00	0.43
39:6:315:CHL:HBB2	22:6:317:CLA:HBC1	2.00	0.43
1:A:161:THR:HG21	22:A:816:CLA:HAA2	2.00	0.43
1:A:487:ILE:HG21	22:A:837:CLA:HED1	2.01	0.43
1:A:674:GLY:HA2	24:A:848:BCR:H17C	2.00	0.43
22:A:833:CLA:H102	22:B:840:CLA:C5	2.48	0.43
2:B:48:PHE:HB2	2:B:169:PHE:CE1	2.54	0.43
2:B:262:PHE:CZ	2:B:356:LEU:HA	2.54	0.43
2:B:687:PRO:HG2	22:L:202:CLA:C5	2.45	0.43
22:B:825:CLA:CAB	22:B:832:CLA:HMD2	2.49	0.43
22:B:831:CLA:CHA	22:B:831:CLA:HBA1	2.42	0.43
22:B:832:CLA:HAA1	22:B:842:CLA:H41	2.00	0.43
22:B:838:CLA:H122	32:F:306:RRX:H38	2.00	0.43
4:D:56:PRO:HD2	4:D:149:TYR:CZ	2.54	0.43
5:E:39:ARG:HG3	5:E:61:SER:HA	1.98	0.43
5:E:86:ASN:HB3	5:E:88:PHE:CE2	2.53	0.43
22:F:304:CLA:OBD	22:F:304:CLA:H122	2.19	0.43
34:J:101:LMG:HC8	14:7:57:GLY:HA3	2.00	0.43
12:1:100:TYR:O	12:1:103:PRO:HD2	2.17	0.43
38:1:303:LUT:C16	22:1:309:CLA:HMB3	2.46	0.43
22:1:307:CLA:C1B	22:1:307:CLA:H2	2.49	0.43
22:1:314:CLA:HMB1	22:1:314:CLA:CBB	2.48	0.43
14:7:205:LEU:HD21	22:7:307:CLA:H171	2.00	0.43
22:7:317:CLA:HMA2	22:7:317:CLA:C1	2.49	0.43
22:7:323:CLA:HED2	15:8:31:MET:HG2	2.00	0.43
41:8:320:3PH:H3A1	41:8:320:3PH:H3D1	1.45	0.43
16:4:100:GLU:CD	16:4:215:ILE:HD11	2.38	0.43
16:4:196:ILE:HD12	16:4:196:ILE:HA	1.80	0.43
24:4:804:BCR:H281	22:4:810:CLA:NB	2.33	0.43
22:5:308:CLA:H72	22:5:308:CLA:H112	1.68	0.43
22:5:323:CLA:O1A	22:5:323:CLA:H2A	2.19	0.43
18:6:88:GLN:HG2	22:6:312:CLA:C3D	2.48	0.43
1:A:119:TRP:CE3	22:A:809:CLA:H2A	2.53	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:376:PRO:HB3	22:A:819:CLA:HMA2	2.00	0.43
1:A:651:TRP:HD1	22:B:805:CLA:CAC	2.31	0.43
1:A:673:LEU:HD12	22:A:802:CLA:H42	1.99	0.43
22:A:834:CLA:H92	22:A:834:CLA:H61	1.71	0.43
2:B:38:MET:O	4:D:186:PHE:HD1	2.02	0.43
2:B:116:ILE:O	22:B:811:CLA:HMD3	2.18	0.43
2:B:590:TRP:CG	22:B:805:CLA:H152	2.53	0.43
22:B:833:CLA:C1B	32:F:306:RRX:H26	2.48	0.43
33:G:204:LMT:H111	22:1:314:CLA:CMC	2.47	0.43
9:J:1:MET:N	34:J:101:LMG:HC2	2.29	0.43
12:1:173:LYS:HG3	22:1:310:CLA:HED2	2.00	0.43
40:1:318:SQD:H192	39:8:301:CHL:CMA	2.48	0.43
12:Z:144:GLU:CG	22:Z:313:CLA:HAC1	2.49	0.43
12:Z:195:THR:HG23	12:Z:197:LYS:HE2	2.01	0.43
38:Z:301:LUT:H35	38:Z:301:LUT:H401	1.81	0.43
22:Z:306:CLA:C1B	22:Z:306:CLA:H2	2.49	0.43
13:3:52:THR:HG23	13:3:55:TRP:H	1.84	0.43
38:7:302:LUT:C28	22:7:307:CLA:H61	2.44	0.43
24:4:804:BCR:H343	39:4:819:CHL:HAB	2.01	0.43
24:5:304:BCR:HC7	39:5:321:CHL:HBB2	2.01	0.43
22:5:318:CLA:H2	22:5:323:CLA:CAC	2.49	0.43
38:6:304:LUT:C31	22:6:310:CLA:HMC2	2.49	0.43
38:9:302:LUT:H35	38:9:302:LUT:H401	1.81	0.43
1:A:349:TRP:CE3	22:A:805:CLA:HMD2	2.54	0.43
21:A:801:CL0:H22	2:B:626:TRP:CD1	2.54	0.43
22:A:816:CLA:H92	27:A:852:DGA:HB61	2.01	0.43
22:A:821:CLA:H111	22:A:821:CLA:H152	1.75	0.43
22:A:826:CLA:O1A	22:A:839:CLA:HAA2	2.19	0.43
22:A:832:CLA:C11	22:L:202:CLA:H171	2.47	0.43
2:B:126:TYR:O	2:B:131:ARG:NH1	2.51	0.43
2:B:431:GLY:HA3	22:B:801:CLA:H11	2.00	0.43
22:B:806:CLA:HBB1	22:B:806:CLA:HMB1	2.01	0.43
22:B:815:CLA:H142	22:B:815:CLA:H112	1.80	0.43
22:B:815:CLA:H151	22:B:820:CLA:C4D	2.48	0.43
22:B:824:CLA:HBC2	22:B:825:CLA:CBA	2.48	0.43
22:F:304:CLA:CBB	32:F:306:RRX:H51	2.45	0.43
7:G:45:MET:HG3	7:G:109:GLY:CA	2.47	0.43
22:L:202:CLA:HBB1	22:L:202:CLA:HMB1	2.01	0.43
12:Z:35:TRP:HD1	39:Z:311:CHL:C1B	2.32	0.43
12:Z:160:LEU:HB3	12:Z:162:PHE:CZ	2.54	0.43
12:Z:166:ASP:OD1	12:Z:169:THR:OG1	2.28	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
31:3:301:DGD:HE5	31:3:301:DGD:O4D	2.18	0.43
24:3:304:BCR:H23C	24:3:305:BCR:H353	2.01	0.43
24:3:305:BCR:C16	39:3:317:CHL:H112	2.48	0.43
14:7:59:SER:HB2	14:7:65:MET:HG2	2.01	0.43
14:7:135:GLU:HG3	22:7:315:CLA:C1B	2.48	0.43
22:7:308:CLA:H92	22:7:308:CLA:H62	1.71	0.43
22:7:322:CLA:HMD1	22:8:307:CLA:H2	2.01	0.43
15:8:37:PRO:O	15:8:38:ALA:HB2	2.19	0.43
15:8:39:HIS:HA	15:8:41:LYS:HZ1	1.82	0.43
22:8:307:CLA:O1D	22:8:307:CLA:H2A	2.19	0.43
16:4:70:HIS:CD2	16:4:71:LEU:HG	2.54	0.43
16:4:105:ARG:HB3	39:4:814:CHL:HED2	2.00	0.43
17:5:78:ALA:HB1	17:5:190:GLY:HA3	2.01	0.43
24:5:304:BCR:H381	22:5:319:CLA:NC	2.34	0.43
22:5:310:CLA:H152	22:5:311:CLA:HBB1	2.00	0.43
18:6:167:MET:O	18:6:171:LYS:HG3	2.19	0.43
18:6:173:LYS:NZ	22:6:308:CLA:H43	2.34	0.43
19:9:32:LEU:O	19:9:35:LEU:HB2	2.18	0.43
19:9:78:MET:HB2	22:9:303:CLA:HMC3	2.00	0.43
20:2:177:VAL:O	20:2:178:ARG:HG3	2.19	0.43
1:A:521:VAL:HG11	1:A:524:MET:HB2	2.00	0.43
22:A:819:CLA:H151	22:A:819:CLA:H18	1.74	0.43
22:A:821:CLA:O2A	22:A:824:CLA:HBA2	2.19	0.43
2:B:669:ARG:NE	2:B:700:ALA:HB3	2.34	0.43
22:B:805:CLA:H51	22:B:805:CLA:H8	1.64	0.43
22:B:814:CLA:H2	22:B:814:CLA:ND	2.33	0.43
22:B:827:CLA:HBA1	22:B:827:CLA:H12	1.73	0.43
6:F:201:THR:O	6:F:205:GLN:HG3	2.19	0.43
33:G:204:LMT:C9	22:1:308:CLA:HMD3	2.48	0.43
12:1:56:ASP:HA	38:1:303:LUT:H24	2.00	0.43
12:1:115:PHE:CE1	22:1:309:CLA:HMD2	2.53	0.43
22:1:316:CLA:H141	39:8:301:CHL:H111	2.00	0.43
38:3:303:LUT:H15	38:3:303:LUT:H201	1.86	0.43
14:7:198:ARG:HA	14:7:201:MET:HE2	2.01	0.43
22:4:808:CLA:H62	22:4:808:CLA:H41	1.58	0.43
17:5:58:LEU:HD12	17:5:58:LEU:N	2.34	0.43
22:5:311:CLA:HMD2	22:5:318:CLA:ND	2.33	0.43
38:6:303:LUT:C28	22:6:307:CLA:H52	2.49	0.43
19:9:195:GLY:O	19:9:196:TYR:C	2.57	0.43
22:2:301:CLA:HBA1	22:2:301:CLA:H11	1.61	0.43
1:A:494:ALA:HB1	1:A:498:THR:OG1	2.18	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:726:VAL:HG22	22:A:841:CLA:O1D	2.19	0.43
22:A:822:CLA:HAA2	10:K:61:THR:CG2	2.47	0.43
22:A:832:CLA:H112	22:L:202:CLA:H152	2.00	0.43
2:B:652:LEU:O	22:B:806:CLA:HBA1	2.19	0.43
22:B:809:CLA:CBC	22:B:811:CLA:H193	2.49	0.43
22:B:819:CLA:H93	22:B:835:CLA:HBA1	2.00	0.43
24:B:848:BCR:H321	24:B:848:BCR:HC8	2.00	0.43
3:C:79:LEU:HD22	3:C:81:TYR:CE2	2.54	0.43
6:F:70:CYS:HA	6:F:73:SER:HB3	2.01	0.43
7:G:77:TYR:CG	7:G:78:PHE:N	2.87	0.43
22:G:202:CLA:CHC	24:G:203:BCR:H402	2.49	0.43
8:I:74:TRP:CZ2	22:2:304:CLA:HHB	2.54	0.43
12:1:157:PHE:HB3	22:1:304:CLA:HMD1	2.01	0.43
38:Z:301:LUT:H371	22:Z:303:CLA:H92	2.01	0.43
13:3:109:ASP:N	13:3:109:ASP:OD1	2.52	0.43
38:3:302:LUT:H30	22:3:308:CLA:H72	1.98	0.43
22:3:308:CLA:H3A	22:3:308:CLA:HBA2	1.17	0.43
22:3:312:CLA:H61	22:3:312:CLA:H2	1.73	0.43
22:3:312:CLA:H61	22:3:312:CLA:H92	1.76	0.43
22:3:322:CLA:HHC	22:3:322:CLA:HBB1	2.01	0.43
42:7:320:SPH:H91	42:7:320:SPH:H61	1.63	0.43
15:8:105:LYS:O	15:8:109:GLU:HG2	2.19	0.43
16:4:225:PHE:HA	16:4:228:PHE:HD2	1.83	0.43
22:4:806:CLA:HBA1	22:4:806:CLA:H61	2.01	0.43
17:5:94:LEU:O	17:5:96:PRO:HD3	2.18	0.43
17:5:107:GLN:H	17:5:107:GLN:CD	2.21	0.43
17:5:227:TRP:N	22:5:309:CLA:O1A	2.52	0.43
18:6:222:VAL:CG1	18:6:224:GLY:H	2.32	0.43
24:6:306:BCR:H19C	39:6:315:CHL:C2	2.49	0.43
24:6:306:BCR:C8	24:6:306:BCR:H311	2.48	0.43
1:A:694:GLN:NE2	5:E:80:TYR:HE1	2.16	0.42
22:A:819:CLA:H203	22:A:827:CLA:HAA1	2.00	0.42
22:A:840:CLA:H11	22:A:840:CLA:C1A	2.49	0.42
2:B:600:LEU:HD23	2:B:600:LEU:HA	1.94	0.42
22:B:809:CLA:HBA1	22:B:809:CLA:H3A	1.55	0.42
24:B:844:BCR:C22	24:G:203:BCR:HC42	2.49	0.42
3:C:74:THR:HG21	4:D:81:GLU:OE1	2.18	0.42
5:E:47:ARG:HD2	5:E:50:SER:OG	2.18	0.42
12:Z:149:LYS:HB3	22:Z:313:CLA:CBC	2.40	0.42
14:7:113:THR:HG22	14:7:115:HIS:H	1.84	0.42
14:7:137:LYS:HB3	22:7:314:CLA:HMC3	2.00	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
38:8:303:LUT:H31	22:8:308:CLA:HMC2	2.01	0.42
16:4:106:TRP:CZ3	24:4:804:BCR:H372	2.54	0.42
16:4:216:LYS:HD2	22:4:811:CLA:O1D	2.19	0.42
17:5:69:TRP:CZ2	22:5:318:CLA:HAA2	2.54	0.42
22:5:309:CLA:H11	22:5:314:CLA:CMD	2.49	0.42
18:6:242:THR:HG23	24:6:306:BCR:H312	2.01	0.42
38:6:304:LUT:H15	38:6:304:LUT:H201	1.84	0.42
20:2:198:ILE:HD12	22:2:301:CLA:CHD	2.48	0.42
1:A:365:ILE:CD1	1:A:398:GLY:HA3	2.49	0.42
1:A:589:TRP:HE1	22:A:830:CLA:HMD1	1.84	0.42
22:A:806:CLA:H12	22:A:806:CLA:HBA2	1.53	0.42
22:A:818:CLA:HMB1	22:A:818:CLA:CBB	2.48	0.42
22:A:821:CLA:H93	22:A:821:CLA:H62	1.85	0.42
22:A:831:CLA:HAB	22:A:839:CLA:CBB	2.49	0.42
2:B:107:ARG:HE	2:B:116:ILE:HG12	1.84	0.42
2:B:371:SER:HA	2:B:726:LEU:HD21	2.00	0.42
2:B:513:ILE:HG22	2:B:517:ASP:OD2	2.19	0.42
24:B:847:BCR:HC31	22:1:314:CLA:H41	2.00	0.42
5:E:38:LYS:HB3	5:E:38:LYS:HE3	1.65	0.42
22:1:308:CLA:C1D	22:1:314:CLA:H93	2.49	0.42
12:Z:115:PHE:H	22:Z:308:CLA:CAD	2.32	0.42
39:Z:312:CHL:HAB	22:Z:315:CLA:CBB	2.49	0.42
39:Z:312:CHL:C1C	22:Z:315:CLA:HMC3	2.49	0.42
13:3:141:GLN:HE22	22:3:322:CLA:HMD3	1.84	0.42
14:7:65:MET:O	14:7:69:VAL:HG23	2.20	0.42
24:7:303:BCR:H373	22:7:316:CLA:HHB	2.00	0.42
22:7:310:CLA:H93	22:7:310:CLA:H62	1.80	0.42
16:4:85:SER:HB2	16:4:88:VAL:HG21	2.00	0.42
17:5:227:TRP:HE1	22:5:314:CLA:HED2	1.83	0.42
38:5:303:LUT:H31	22:5:310:CLA:HMC2	2.01	0.42
38:5:303:LUT:C16	22:5:312:CLA:HMB3	2.48	0.42
22:5:308:CLA:HMD2	22:5:313:CLA:CHD	2.48	0.42
22:5:310:CLA:HMB1	22:5:310:CLA:CBB	2.49	0.42
22:5:314:CLA:CBB	22:5:314:CLA:HHC	2.49	0.42
19:9:59:ASP:OD2	19:9:62:ARG:HG3	2.19	0.42
38:9:301:LUT:H162	22:9:305:CLA:HHB	2.01	0.42
1:A:365:ILE:HG13	1:A:395:ASN:HA	2.01	0.42
1:A:366:ILE:HD12	22:A:827:CLA:H42	2.01	0.42
22:A:810:CLA:CAB	22:A:813:CLA:H121	2.48	0.42
22:A:814:CLA:C2C	24:A:844:BCR:H352	2.49	0.42
2:B:5:LEU:HG	8:I:103:ASN:HB2	2.01	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:B:176:LEU:HD12	2:B:176:LEU:HA	1.87	0.42
2:B:211:ASP:HB3	25:B:851:LHG:HC42	2.01	0.42
2:B:527:GLY:HA2	2:B:583:TRP:CZ3	2.53	0.42
24:B:853:BCR:HC32	22:L:202:CLA:C4C	2.49	0.42
3:C:13:GLY:HA2	3:C:28:MET:HE1	2.00	0.42
22:F:304:CLA:C2	22:F:304:CLA:HBA1	2.49	0.42
8:I:98:TYR:CD2	8:I:99:ILE:HD12	2.51	0.42
10:K:33:THR:O	10:K:37:ILE:HG13	2.19	0.42
38:Z:301:LUT:C30	22:Z:303:CLA:H8	2.49	0.42
22:3:309:CLA:O1D	22:3:314:CLA:H51	2.19	0.42
22:7:307:CLA:H18	22:7:307:CLA:H151	1.80	0.42
22:7:312:CLA:CGD	22:7:315:CLA:HMA1	2.50	0.42
38:8:302:LUT:H30	22:8:305:CLA:H61	2.00	0.42
18:6:46:LEU:H	18:6:46:LEU:CD1	2.27	0.42
24:6:305:BCR:H402	22:6:312:CLA:C1B	2.50	0.42
1:A:483:PHE:HB3	22:A:837:CLA:C3	2.48	0.42
22:A:812:CLA:C9	38:3:303:LUT:H372	2.49	0.42
22:A:824:CLA:H52	24:A:847:BCR:H363	2.01	0.42
22:A:832:CLA:H18	22:A:832:CLA:H151	1.69	0.42
22:A:835:CLA:H11	22:A:835:CLA:HBA1	1.72	0.42
2:B:43:LEU:O	2:B:47:ILE:HG13	2.20	0.42
2:B:243:HIS:HD2	2:B:251:ALA:HA	1.84	0.42
22:B:815:CLA:H192	22:B:830:CLA:H93	2.01	0.42
22:B:817:CLA:HBA1	19:9:177:LEU:HD22	2.00	0.42
22:B:826:CLA:HMB1	22:B:826:CLA:CBB	2.50	0.42
4:D:138:PHE:CZ	4:D:151:HIS:HB2	2.54	0.42
6:F:162:TRP:CZ3	6:F:200:ALA:HA	2.54	0.42
12:1:158:ASP:O	12:1:161:ASN:ND2	2.50	0.42
38:Z:302:LUT:H35	38:Z:302:LUT:H401	1.83	0.42
13:3:182:LEU:HD23	22:3:314:CLA:HED2	2.01	0.42
22:3:318:CLA:H62	22:3:318:CLA:H2	1.64	0.42
14:7:84:ALA:HB3	22:7:309:CLA:HMC3	2.01	0.42
38:7:302:LUT:H35	38:7:302:LUT:H401	1.87	0.42
42:7:320:SPH:H121	42:7:320:SPH:H152	1.45	0.42
15:8:155:SER:HB3	15:8:167:PRO:HD3	2.02	0.42
22:8:310:CLA:H62	43:8:319:LPX:H5A	2.01	0.42
39:8:312:CHL:C1D	26:8:318:NKP:HAL	2.50	0.42
16:4:62:PHE:HD2	16:4:65:SER:HB3	1.85	0.42
24:4:804:BCR:H371	39:4:816:CHL:HMB2	2.00	0.42
18:6:72:ARG:O	18:6:76:ARG:HG3	2.20	0.42
18:6:91:VAL:O	18:6:93:PRO:HD3	2.20	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
18:6:95:VAL:HA	18:6:103:LYS:HE2	2.02	0.42
18:6:123:PHE:CD2	24:6:305:BCR:H12C	2.54	0.42
22:6:302:CLA:HHC	22:6:302:CLA:HBB1	2.01	0.42
22:6:310:CLA:H13	22:6:310:CLA:H102	1.82	0.42
19:9:110:PRO:HB2	20:2:194:TRP:CZ3	2.54	0.42
19:9:125:LYS:NZ	19:9:125:LYS:HB3	2.35	0.42
20:2:189:HIS:ND1	22:2:301:CLA:HAA2	2.34	0.42
1:A:101:TYR:CZ	1:A:105:LEU:HD21	2.54	0.42
1:A:150:ALA:HB2	1:A:378:PRO:HD2	2.02	0.42
1:A:650:LEU:HD13	22:B:805:CLA:CBB	2.50	0.42
1:A:686:LEU:HG	1:A:727:GLY:HA3	2.00	0.42
22:A:805:CLA:HMA2	22:A:805:CLA:H12	2.01	0.42
22:A:808:CLA:H93	22:A:808:CLA:H62	1.87	0.42
22:A:810:CLA:HBD	13:3:51:VAL:HG23	2.00	0.42
22:A:829:CLA:H2	22:A:829:CLA:H62	1.73	0.42
22:A:838:CLA:HHB	22:A:839:CLA:O1D	2.19	0.42
24:A:846:BCR:H15C	24:A:846:BCR:H351	1.66	0.42
2:B:27:ALA:C	2:B:29:ALA:H	2.22	0.42
2:B:231:TRP:CH2	22:B:818:CLA:H92	2.54	0.42
22:B:841:CLA:NB	24:B:849:BCR:H15C	2.35	0.42
24:B:844:BCR:H20C	24:B:844:BCR:H361	1.42	0.42
25:B:850:LHG:HC81	25:B:850:LHG:HC5	1.83	0.42
4:D:86:ILE:HG21	4:D:124:LEU:HD23	2.01	0.42
8:I:68:PRO:HG3	20:2:205:LYS:HZ2	1.84	0.42
8:I:95:LEU:HD13	24:L:204:BCR:HC8	2.00	0.42
12:Z:160:LEU:N	12:Z:160:LEU:HD22	2.34	0.42
12:Z:188:PHE:HA	12:Z:191:GLN:HE22	1.82	0.42
38:Z:302:LUT:H15	38:Z:302:LUT:H201	1.83	0.42
39:Z:312:CHL:HED2	39:Z:312:CHL:H2A	2.00	0.42
13:3:41:LEU:HD12	13:3:41:LEU:N	2.34	0.42
24:3:307:BCR:H351	24:3:307:BCR:H15C	1.46	0.42
14:7:149:GLY:HA3	14:7:161:LYS:HD2	2.02	0.42
14:7:167:TYR:CZ	14:7:190:LYS:HE3	2.54	0.42
38:7:302:LUT:H15	38:7:302:LUT:H201	1.83	0.42
16:4:118:ASP:HB2	16:4:240:LEU:HD23	2.01	0.42
22:4:812:CLA:H61	22:4:812:CLA:H2	1.60	0.42
22:5:318:CLA:H11	22:6:322:CLA:H91	2.00	0.42
18:6:80:LEU:HD11	22:6:307:CLA:CBC	2.49	0.42
18:6:142:PRO:HB2	22:6:302:CLA:CMD	2.44	0.42
22:6:311:CLA:OBD	22:6:317:CLA:HAA1	2.19	0.42
1:A:278:PHE:HB2	22:A:817:CLA:O1D	2.20	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:490:THR:O	1:A:494:ALA:HB2	2.20	0.42
22:A:805:CLA:H93	22:A:805:CLA:H111	1.71	0.42
22:A:806:CLA:HMA1	22:A:830:CLA:C3B	2.49	0.42
22:A:812:CLA:H2	22:A:812:CLA:ND	2.35	0.42
22:A:812:CLA:H71	22:A:812:CLA:CHC	2.48	0.42
22:A:841:CLA:HBC1	24:F:303:BCR:H392	2.00	0.42
2:B:166:LEU:HD11	2:B:170:LYS:HE3	2.01	0.42
2:B:517:ASP:HA	2:B:520:VAL:HG12	2.01	0.42
2:B:712:THR:O	2:B:716:VAL:HG23	2.19	0.42
22:B:821:CLA:O1A	22:B:826:CLA:H112	2.20	0.42
6:F:192:ALA:O	6:F:196:THR:HG23	2.18	0.42
7:G:122:LEU:HD22	7:G:122:LEU:HA	1.83	0.42
12:1:192:HIS:O	12:1:196:GLY:HA2	2.19	0.42
13:3:15:LEU:HA	13:3:34:PHE:O	2.19	0.42
22:7:324:CLA:H11	22:7:324:CLA:H51	1.83	0.42
15:8:186:LEU:HD23	22:8:311:CLA:HED3	2.00	0.42
16:4:225:PHE:CD2	38:4:803:LUT:H14	2.54	0.42
38:4:802:LUT:H31	38:4:802:LUT:H391	1.90	0.42
22:4:812:CLA:H93	24:6:305:BCR:H311	2.01	0.42
17:5:55:PHE:HB3	22:5:310:CLA:C3D	2.50	0.42
19:9:76:TRP:CE3	22:9:313:CLA:HAC2	2.55	0.42
1:A:161:THR:HA	27:A:852:DGA:HAW1	2.01	0.42
1:A:370:HIS:HD1	22:A:819:CLA:H41	1.84	0.42
1:A:736:ILE:HD13	22:A:828:CLA:HMC2	2.02	0.42
22:A:817:CLA:H61	22:A:817:CLA:C3C	2.50	0.42
22:A:829:CLA:C2	24:A:845:BCR:HC7	2.41	0.42
23:A:843:PQN:H18	23:A:843:PQN:H222	1.42	0.42
2:B:33:GLU:H	2:B:33:GLU:HG3	1.48	0.42
2:B:61:TRP:CG	22:B:809:CLA:HBC1	2.54	0.42
2:B:443:VAL:HG21	22:B:834:CLA:HAC2	2.01	0.42
2:B:453:GLN:NE2	2:B:455:LEU:HD11	2.32	0.42
22:B:806:CLA:HMB1	22:B:806:CLA:CBB	2.50	0.42
22:B:822:CLA:NB	24:B:844:BCR:H351	2.35	0.42
22:B:829:CLA:H111	22:B:829:CLA:C2D	2.50	0.42
6:F:202:TRP:N	6:F:203:PRO:CD	2.83	0.42
22:L:201:CLA:H62	22:L:201:CLA:H102	1.30	0.42
13:3:72:ALA:HB1	24:3:305:BCR:H383	2.01	0.42
22:3:320:CLA:NA	41:7:319:3PH:H3D2	2.34	0.42
22:3:322:CLA:CMB	41:7:319:3PH:H2A1	2.49	0.42
16:4:111:ALA:HB1	38:4:802:LUT:H12	2.00	0.42
17:5:252:LEU:HD23	41:5:325:3PH:H342	2.00	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
18:6:248:CYS:O	18:6:249:LEU:C	2.57	0.42
22:6:301:CLA:H11	22:6:301:CLA:H52	1.71	0.42
22:6:314:CLA:H93	22:6:314:CLA:H112	1.63	0.42
19:9:113:ALA:HA	19:9:116:PHE:HD2	1.84	0.42
20:2:165:MET:O	20:2:169:ILE:HG12	2.20	0.42
1:A:478:GLN:HB3	1:A:480:GLN:HE21	1.83	0.42
22:A:805:CLA:H62	22:A:805:CLA:H2	1.55	0.42
22:A:822:CLA:CBB	37:K:201:DAO:H112	2.46	0.42
22:A:833:CLA:HBB1	22:L:203:CLA:HBB2	2.02	0.42
2:B:67:PHE:HZ	8:I:71:PRO:HG2	1.85	0.42
2:B:89:ALA:HB1	22:B:811:CLA:O2D	2.19	0.42
25:B:803:LHG:HC91	22:B:807:CLA:C3D	2.50	0.42
22:B:829:CLA:H62	22:B:829:CLA:H2	1.63	0.42
5:E:79:ASN:OD1	5:E:83:VAL:N	2.53	0.42
6:F:220:GLU:H	6:F:220:GLU:HG3	1.50	0.42
22:L:202:CLA:HMB1	22:L:202:CLA:CBB	2.50	0.42
22:1:308:CLA:HMD2	22:1:314:CLA:CHD	2.50	0.42
22:Z:316:CLA:H111	22:Z:316:CLA:H152	1.49	0.42
13:3:82:LYS:HZ1	13:3:211:GLU:HG2	1.82	0.42
22:3:311:CLA:H62	22:3:311:CLA:H93	1.80	0.42
22:3:322:CLA:CBB	22:3:322:CLA:HHC	2.48	0.42
14:7:67:TRP:CH2	22:7:315:CLA:HAA1	2.54	0.42
14:7:135:GLU:HG3	22:7:315:CLA:NB	2.35	0.42
22:7:323:CLA:H93	22:7:323:CLA:H111	1.68	0.42
15:8:205:GLN:NE2	38:8:302:LUT:H42	2.35	0.42
22:8:307:CLA:H51	22:8:307:CLA:C4C	2.50	0.42
39:8:315:CHL:HAA2	39:8:315:CHL:CBD	2.50	0.42
24:4:804:BCR:H24C	39:4:816:CHL:HAA1	2.01	0.42
17:5:100:TRP:CH2	17:5:201:VAL:HA	2.55	0.42
17:5:123:ALA:CB	22:5:326:CLA:HAA1	2.50	0.42
39:5:316:CHL:HMC	22:5:319:CLA:CAB	2.45	0.42
18:6:214:THR:HA	22:6:309:CLA:HBA2	2.01	0.42
18:6:248:CYS:HB3	18:6:254:TRP:CH2	2.55	0.42
39:9:312:CHL:H162	39:9:312:CHL:H121	1.85	0.42
1:A:341:LEU:HD11	1:A:426:LEU:CD1	2.50	0.42
1:A:461:THR:CG2	22:B:812:CLA:HAB	2.48	0.42
22:A:815:CLA:C1C	24:A:856:BCR:H312	2.49	0.42
2:B:13:LEU:HD11	31:B:852:DGD:HE1	2.02	0.42
2:B:653:PHE:CD1	22:B:806:CLA:H71	2.54	0.42
22:B:801:CLA:H152	22:B:801:CLA:H18	1.83	0.42
22:B:819:CLA:H43	22:B:835:CLA:HED1	2.02	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
22:B:823:CLA:HMB1	33:1:301:LMT:H71	2.01	0.42
6:F:114:ARG:HB2	9:J:35:ASP:OD2	2.20	0.42
22:1:307:CLA:HMD2	39:8:301:CHL:CBB	2.50	0.42
13:3:216:HIS:CD2	22:3:310:CLA:HAA1	2.54	0.42
14:7:209:ALA:CB	22:7:317:CLA:H8	2.48	0.42
22:7:305:CLA:C4B	22:7:310:CLA:H121	2.50	0.42
15:8:187:LYS:HG2	22:8:306:CLA:HBD	2.01	0.42
38:8:302:LUT:H161	22:8:307:CLA:HMB3	2.01	0.42
17:5:158:PRO:HB2	17:5:160:PRO:HD3	2.02	0.42
24:5:304:BCR:H19C	22:5:319:CLA:CBA	2.28	0.42
39:9:312:CHL:H18	39:9:312:CHL:C1C	2.50	0.42
39:9:314:CHL:HED2	20:2:194:TRP:CH2	2.54	0.42
1:A:129:GLY:O	1:A:136:GLN:HA	2.20	0.42
1:A:205:LEU:HD12	24:A:845:BCR:H363	2.02	0.42
1:A:278:PHE:CZ	22:A:818:CLA:H43	2.54	0.42
1:A:369:HIS:HA	1:A:372:TYR:CE2	2.55	0.42
1:A:651:TRP:CD1	22:B:805:CLA:HBC1	2.55	0.42
22:A:806:CLA:H62	22:A:806:CLA:H2	1.64	0.42
22:A:815:CLA:O1A	13:3:203:VAL:HG12	2.20	0.42
22:A:840:CLA:HHD	24:F:303:BCR:H383	2.01	0.42
2:B:229:GLY:O	2:B:231:TRP:N	2.50	0.42
2:B:419:ILE:HG23	22:B:839:CLA:HBB2	2.02	0.42
2:B:490:SER:HB3	2:B:495:LEU:HB2	2.01	0.42
2:B:712:THR:HG22	31:B:852:DGD:HBW1	2.02	0.42
8:I:93:ALA:HB1	11:L:134:ILE:HG21	2.00	0.42
11:L:140:PHE:CD1	11:L:144:PRO:HD2	2.53	0.42
12:1:206:HIS:HA	12:1:213:SER:OG	2.20	0.42
12:Z:186:LEU:HG	22:Z:305:CLA:HAC1	2.02	0.42
22:Z:309:CLA:HHD	25:Z:317:LHG:H242	2.02	0.42
22:3:320:CLA:C4A	41:7:319:3PH:H3D2	2.50	0.42
22:7:307:CLA:H93	22:7:307:CLA:H111	1.77	0.42
16:4:193:PRO:HD2	22:4:805:CLA:CAD	2.50	0.42
25:4:820:LHG:H352	22:6:321:CLA:C1	2.50	0.42
18:6:79:MET:HE3	22:6:307:CLA:HMC3	2.01	0.42
18:6:85:ILE:HG12	38:6:304:LUT:H41	2.01	0.42
18:6:100:ALA:HB3	18:6:101:PRO:HD3	2.02	0.42
39:6:315:CHL:H61	39:6:315:CHL:H41	1.88	0.42
39:6:318:CHL:H62	39:6:318:CHL:H41	1.74	0.42
19:9:102:LYS:CD	22:9:308:CLA:HED1	2.50	0.42
19:9:175:GLN:OE1	38:9:301:LUT:H42	2.19	0.42
22:9:305:CLA:H142	22:9:305:CLA:H112	1.89	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:634:ASN:O	1:A:638:SER:HB2	2.19	0.41
22:A:820:CLA:H71	22:A:820:CLA:H111	1.20	0.41
22:A:832:CLA:H112	22:A:832:CLA:H152	1.66	0.41
22:A:855:CLA:H161	24:B:853:BCR:H351	2.01	0.41
2:B:687:PRO:O	2:B:688:LEU:HB2	2.20	0.41
22:B:826:CLA:H92	22:B:826:CLA:H61	1.65	0.41
3:C:15:THR:HG22	3:C:28:MET:HG3	2.02	0.41
38:Z:301:LUT:H403	22:Z:303:CLA:C14	2.47	0.41
22:3:314:CLA:H41	22:3:314:CLA:H62	1.32	0.41
14:7:241:VAL:CG1	14:7:244:TYR:HB3	2.50	0.41
22:7:312:CLA:HED2	22:7:312:CLA:HBD	1.85	0.41
39:7:313:CHL:CMC	22:7:316:CLA:HAB	2.50	0.41
38:8:302:LUT:H35	38:8:302:LUT:H401	1.79	0.41
38:5:303:LUT:H35	38:5:303:LUT:H401	1.85	0.41
22:5:326:CLA:HMD1	22:6:309:CLA:H61	2.01	0.41
22:6:307:CLA:H41	22:6:308:CLA:HBA1	2.02	0.41
22:6:308:CLA:H51	22:6:308:CLA:H11	1.63	0.41
39:6:320:CHL:OMC	39:6:320:CHL:HAC2	2.20	0.41
1:A:179:PHE:O	1:A:183:LYS:HB2	2.20	0.41
1:A:483:PHE:HB3	22:A:837:CLA:H42	2.01	0.41
22:A:823:CLA:H111	22:A:823:CLA:H152	1.46	0.41
24:A:844:BCR:H393	24:3:306:BCR:HC42	2.02	0.41
2:B:105:PHE:HD2	2:B:107:ARG:HH12	1.67	0.41
2:B:178:HIS:ND1	22:B:815:CLA:HMC2	2.35	0.41
2:B:384:MET:HA	2:B:384:MET:HE3	2.02	0.41
2:B:388:PHE:HZ	22:B:827:CLA:HAB	1.85	0.41
2:B:587:THR:O	2:B:591:VAL:HG13	2.21	0.41
2:B:656:LEU:HA	22:B:806:CLA:HAA1	2.01	0.41
2:B:727:ILE:CD1	22:B:829:CLA:H12	2.50	0.41
22:B:806:CLA:H92	22:B:806:CLA:H62	1.71	0.41
22:B:810:CLA:C1	8:I:79:PHE:HB3	2.50	0.41
22:B:830:CLA:H52	24:B:845:BCR:C23	2.50	0.41
6:F:222:ILE:H	6:F:222:ILE:HG13	1.79	0.41
33:G:204:LMT:H41	33:G:204:LMT:H12	1.79	0.41
12:Z:167:ILE:O	12:Z:171:LYS:HG3	2.20	0.41
13:3:183:ASN:HB3	22:3:309:CLA:CAA	2.51	0.41
14:7:155:GLY:N	42:7:320:SPH:H62	2.35	0.41
38:7:302:LUT:H24	22:7:307:CLA:O1A	2.20	0.41
15:8:111:THR:HG21	39:8:315:CHL:CMD	2.50	0.41
38:8:303:LUT:H15	38:8:303:LUT:H201	1.95	0.41
24:8:304:BCR:H19C	39:8:315:CHL:CMA	2.49	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
16:4:156:LEU:HD22	24:4:804:BCR:H15C	2.02	0.41
17:5:172:LEU:HD12	38:5:302:LUT:H222	2.01	0.41
22:5:311:CLA:CHB	22:5:318:CLA:H191	2.51	0.41
22:5:318:CLA:C9	22:6:322:CLA:H62	2.49	0.41
18:6:87:VAL:HG12	22:6:312:CLA:HMD3	2.02	0.41
18:6:121:MET:HG2	22:6:317:CLA:HMC3	2.02	0.41
19:9:31:TRP:CZ3	19:9:32:LEU:HD23	2.55	0.41
1:A:46:THR:CG2	1:A:715:PRO:HA	2.51	0.41
1:A:376:PRO:HG3	22:A:819:CLA:HBA1	2.03	0.41
22:A:828:CLA:H72	22:A:828:CLA:H111	1.19	0.41
22:A:828:CLA:H93	24:J:104:BCR:H353	2.01	0.41
22:A:841:CLA:H141	22:A:841:CLA:H162	1.94	0.41
2:B:495:LEU:HD22	2:B:499:LEU:HG	2.02	0.41
22:B:811:CLA:H51	22:B:811:CLA:H8	1.22	0.41
22:B:818:CLA:H121	22:B:818:CLA:HMC2	2.01	0.41
22:B:821:CLA:H41	22:B:825:CLA:H91	2.02	0.41
22:B:823:CLA:HMA1	22:B:824:CLA:O1A	2.20	0.41
22:B:823:CLA:HAA2	7:G:60:ASP:OD2	2.21	0.41
22:B:830:CLA:C1D	24:B:845:BCR:H272	2.50	0.41
6:F:204:VAL:CG2	22:8:309:CLA:H72	2.42	0.41
22:1:308:CLA:HBC1	22:1:314:CLA:HAC1	2.02	0.41
22:1:313:CLA:H102	22:1:313:CLA:H13	1.88	0.41
12:Z:83:LEU:HB2	22:Z:308:CLA:HBB2	2.02	0.41
13:3:114:PHE:CD1	22:3:316:CLA:HBD	2.55	0.41
13:3:122:GLN:HE21	13:3:122:GLN:HB3	1.73	0.41
13:3:158:TYR:CE1	13:3:181:LYS:HG2	2.55	0.41
14:7:150:ASP:OD1	14:7:150:ASP:N	2.53	0.41
39:4:813:CHL:HAB	39:4:816:CHL:CAB	2.50	0.41
24:5:305:BCR:H363	39:5:316:CHL:ND	2.34	0.41
22:5:308:CLA:CMA	22:5:308:CLA:HBA1	2.49	0.41
22:5:314:CLA:HHC	22:5:314:CLA:HBB1	2.02	0.41
22:6:309:CLA:H62	22:6:309:CLA:CHC	2.51	0.41
1:A:346:THR:HG23	22:A:825:CLA:HBC1	2.00	0.41
1:A:547:VAL:CG2	22:A:826:CLA:HMC2	2.50	0.41
1:A:690:ARG:HG3	1:A:717:ALA:CB	2.48	0.41
1:A:740:TRP:CG	24:A:848:BCR:HC41	2.55	0.41
22:A:808:CLA:CAD	22:A:828:CLA:HAA2	2.51	0.41
22:A:823:CLA:C11	24:A:856:BCR:H401	2.41	0.41
22:A:840:CLA:CBB	6:F:167:GLY:HA3	2.51	0.41
2:B:356:LEU:HD13	22:B:819:CLA:HAA2	2.03	0.41
2:B:439:VAL:HG12	22:B:834:CLA:HAC1	2.03	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
22:B:827:CLA:C2	22:B:839:CLA:HBD	2.46	0.41
3:C:5:VAL:HA	3:C:66:ARG:O	2.21	0.41
22:F:304:CLA:NB	22:F:304:CLA:H43	2.36	0.41
32:F:306:RRX:H40	32:F:306:RRX:H36	1.34	0.41
22:K:203:CLA:HMB1	22:K:203:CLA:CBB	2.51	0.41
22:L:202:CLA:H2	22:L:202:CLA:H61	1.61	0.41
12:1:140:ARG:HG2	22:1:313:CLA:CHC	2.50	0.41
38:1:302:LUT:H15	38:1:302:LUT:H201	1.86	0.41
22:1:313:CLA:O1A	22:1:313:CLA:H93	2.21	0.41
12:Z:85:ALA:CB	38:Z:301:LUT:H12	2.51	0.41
39:Z:311:CHL:HAA2	16:4:162:PHE:HB2	2.01	0.41
13:3:38:PRO:HD2	38:3:303:LUT:C24	2.50	0.41
13:3:111:TYR:CE2	14:7:244:TYR:HB2	2.55	0.41
38:3:303:LUT:H31	38:3:303:LUT:H391	1.92	0.41
24:3:306:BCR:H393	22:3:318:CLA:H92	2.02	0.41
22:3:322:CLA:C2C	25:7:318:LHG:HC82	2.51	0.41
22:7:305:CLA:O1D	22:7:310:CLA:H92	2.19	0.41
16:4:85:SER:OG	33:4:801:LMT:H1B	2.20	0.41
17:5:240:VAL:HG13	17:5:247:ILE:HB	2.03	0.41
24:5:304:BCR:H382	22:5:319:CLA:NB	2.36	0.41
22:5:309:CLA:CHB	22:5:314:CLA:HMD3	2.50	0.41
22:5:315:CLA:H93	22:5:315:CLA:H62	1.75	0.41
22:5:318:CLA:H42	22:5:323:CLA:HAC2	2.02	0.41
18:6:245:PHE:HE2	22:6:301:CLA:H112	1.85	0.41
39:6:316:CHL:H12	39:6:316:CHL:HBA1	1.52	0.41
38:9:302:LUT:H31	38:9:302:LUT:H391	1.90	0.41
1:A:28:LYS:CG	22:A:811:CLA:HMA2	2.49	0.41
1:A:495:PRO:HA	1:A:499:ALA:O	2.20	0.41
22:A:814:CLA:C9	27:A:852:DGA:HB81	2.50	0.41
22:A:817:CLA:C1A	22:A:817:CLA:CGA	2.98	0.41
22:A:822:CLA:CBC	22:A:823:CLA:H171	2.50	0.41
22:A:834:CLA:CHC	24:I:4001:BCR:H291	2.51	0.41
2:B:298:ILE:HG22	7:G:57:HIS:NE2	2.35	0.41
2:B:398:ASP:OD2	4:D:185:LYS:HE3	2.21	0.41
2:B:559:PRO:HB3	2:B:703:ILE:HB	2.02	0.41
22:B:815:CLA:H102	24:B:845:BCR:C38	2.50	0.41
22:B:829:CLA:HBC3	31:B:852:DGD:HBG3	2.02	0.41
22:B:838:CLA:HAC2	32:F:306:RRX:H2	2.02	0.41
22:B:838:CLA:C19	22:F:302:CLA:H101	2.51	0.41
22:B:842:CLA:CED	24:B:847:BCR:H11C	2.51	0.41
3:C:22:PRO:O	4:D:122:LEU:HD23	2.20	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
12:1:162:PHE:HB2	22:1:304:CLA:O2A	2.20	0.41
12:1:167:ILE:HG22	22:1:304:CLA:HMA2	2.02	0.41
22:1:310:CLA:H91	22:1:310:CLA:H111	1.77	0.41
12:Z:174:GLU:HG3	22:Z:303:CLA:C4B	2.50	0.41
14:7:136:GLN:HG3	22:7:323:CLA:HBA1	2.02	0.41
14:7:218:PRO:C	14:7:220:GLN:H	2.24	0.41
14:7:241:VAL:HG12	14:7:244:TYR:H	1.85	0.41
22:7:309:CLA:C4C	42:7:321:SPH:H151	2.49	0.41
15:8:61:ALA:HB1	15:8:65:TYR:CE2	2.56	0.41
15:8:106:VAL:O	15:8:110:GLN:HB2	2.21	0.41
39:8:315:CHL:H111	39:8:315:CHL:C6	2.46	0.41
22:5:322:CLA:H152	22:6:319:CLA:H52	2.03	0.41
38:6:304:LUT:H35	38:6:304:LUT:H401	1.86	0.41
24:6:305:BCR:H271	22:6:312:CLA:H101	2.03	0.41
24:6:306:BCR:H363	39:6:315:CHL:C4D	2.50	0.41
22:6:322:CLA:H2A	22:6:322:CLA:CED	2.49	0.41
19:9:110:PRO:HG2	20:2:200:THR:HG23	2.02	0.41
1:A:14:LYS:HA	22:A:812:CLA:HAA1	2.03	0.41
1:A:15:ILE:HG12	22:A:812:CLA:OBD	2.21	0.41
1:A:211:LEU:CD2	24:A:844:BCR:H373	2.50	0.41
1:A:584:CYS:HB3	2:B:668:TRP:HE3	1.86	0.41
22:A:809:CLA:NC	22:A:828:CLA:H172	2.35	0.41
22:A:814:CLA:H43	22:A:816:CLA:C4	2.50	0.41
2:B:78:TRP:HH2	2:B:123:GLN:HA	1.85	0.41
22:B:808:CLA:H2	22:B:808:CLA:H61	1.61	0.41
22:B:808:CLA:H143	22:B:815:CLA:C3D	2.50	0.41
22:B:824:CLA:HMB3	22:B:842:CLA:ND	2.36	0.41
22:B:830:CLA:H52	24:B:845:BCR:H23C	2.02	0.41
7:G:33:GLU:CD	7:G:33:GLU:H	2.23	0.41
12:1:206:HIS:CG	22:1:306:CLA:HAA2	2.56	0.41
22:Z:306:CLA:HMD2	39:Z:311:CHL:HBB1	2.01	0.41
22:Z:316:CLA:H141	22:Z:316:CLA:H172	2.01	0.41
22:Z:316:CLA:ND	22:Z:316:CLA:H13	2.35	0.41
22:3:313:CLA:H13	22:5:320:CLA:NB	2.35	0.41
22:3:313:CLA:H3A	22:3:313:CLA:HBA2	1.46	0.41
22:3:322:CLA:HBC1	14:7:31:PRO:HB2	2.03	0.41
22:7:322:CLA:H61	22:7:322:CLA:H2	1.66	0.41
22:7:322:CLA:C3	22:7:323:CLA:H193	2.50	0.41
22:7:323:CLA:H161	22:7:323:CLA:H121	1.79	0.41
22:4:808:CLA:HMD2	22:6:302:CLA:HBB1	2.03	0.41
19:9:56:LEU:CD1	38:9:302:LUT:H221	2.45	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
19:9:113:ALA:HA	19:9:116:PHE:CD2	2.56	0.41
22:9:313:CLA:HMA1	20:2:219:ILE:HD12	2.02	0.41
20:2:163:LEU:C	20:2:165:MET:H	2.24	0.41
1:A:368:ALA:HB2	1:A:394:HIS:HB2	2.03	0.41
1:A:377:TYR:CE2	22:A:829:CLA:HED2	2.56	0.41
1:A:396:TRP:HZ3	1:A:600:TYR:HA	1.85	0.41
1:A:406:GLY:CA	24:A:847:BCR:H323	2.51	0.41
1:A:499:ALA:HB1	22:A:836:CLA:HED3	2.02	0.41
1:A:695:GLU:HA	1:A:698:GLU:CD	2.41	0.41
22:A:803:CLA:HBA2	22:A:803:CLA:H12	1.72	0.41
22:A:807:CLA:O2D	22:A:809:CLA:HED1	2.19	0.41
22:A:822:CLA:CHD	24:A:856:BCR:H402	2.50	0.41
22:A:836:CLA:HBA2	22:A:836:CLA:CHA	2.49	0.41
2:B:24:PHE:O	2:B:28:VAL:HG23	2.21	0.41
2:B:413:LEU:HA	2:B:416:LYS:HB3	2.03	0.41
22:B:819:CLA:C9	22:B:828:CLA:HAB	2.41	0.41
22:B:832:CLA:CHA	22:B:842:CLA:H52	2.51	0.41
4:D:106:MET:SD	4:D:134:PHE:HE2	2.43	0.41
6:F:142:LYS:HA	33:F:307:LMT:H2'	2.03	0.41
12:1:157:PHE:CE2	22:1:313:CLA:H161	2.55	0.41
12:Z:215:PHE:CG	22:Z:305:CLA:H42	2.55	0.41
22:3:312:CLA:ND	22:3:318:CLA:H101	2.35	0.41
15:8:39:HIS:HA	15:8:41:LYS:HZ3	1.83	0.41
15:8:124:PHE:HD1	39:8:301:CHL:H51	1.85	0.41
22:8:305:CLA:HBB1	22:8:306:CLA:CAA	2.51	0.41
22:8:305:CLA:H71	22:8:306:CLA:HMA1	2.01	0.41
16:4:78:ASP:HA	22:4:808:CLA:O1D	2.21	0.41
16:4:83:PRO:CD	38:4:803:LUT:H23	2.42	0.41
16:4:150:PHE:HB2	39:4:813:CHL:HAC2	2.01	0.41
16:4:238:THR:HB	16:4:239:PRO:HD2	2.03	0.41
17:5:161:GLU:HG3	17:5:179:ARG:HH22	1.86	0.41
18:6:95:VAL:CG1	18:6:100:ALA:HA	2.50	0.41
39:9:312:CHL:C3C	39:9:314:CHL:CBB	2.98	0.41
1:A:575:CYS:SG	29:B:802:SF4:S1	3.19	0.41
1:A:661:TYR:OH	1:A:670:LEU:HD23	2.21	0.41
22:A:817:CLA:H12	22:A:817:CLA:ND	2.35	0.41
2:B:32:PHE:HD1	2:B:35:HIS:CE1	2.38	0.41
2:B:143:LEU:HD21	24:B:846:BCR:C24	2.50	0.41
2:B:172:ALA:O	2:B:173:GLU:C	2.59	0.41
2:B:243:HIS:HB2	2:B:250:GLY:O	2.21	0.41
2:B:315:ARG:HB2	22:B:842:CLA:O1A	2.21	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:B:331:LEU:HD23	2:B:331:LEU:HA	1.90	0.41
22:B:830:CLA:ND	24:B:845:BCR:H282	2.35	0.41
22:B:838:CLA:H2	22:B:839:CLA:CMD	2.51	0.41
22:B:842:CLA:HMB1	22:B:842:CLA:CBB	2.51	0.41
6:F:153:THR:HG21	32:F:306:RRX:C4	2.51	0.41
12:Z:176:LYS:HE2	22:Z:309:CLA:C2D	2.50	0.41
38:Z:302:LUT:H371	22:Z:306:CLA:H142	2.02	0.41
22:3:312:CLA:H162	22:3:312:CLA:H141	1.85	0.41
14:7:73:LEU:HD13	14:7:166:GLY:HA3	2.03	0.41
14:7:129:LEU:O	22:7:323:CLA:H11	2.21	0.41
14:7:173:PHE:HZ	22:7:314:CLA:CHC	2.34	0.41
22:7:304:CLA:H101	22:7:305:CLA:HMA1	2.03	0.41
22:7:322:CLA:C3D	22:7:323:CLA:H91	2.51	0.41
16:4:109:LEU:HB3	22:4:810:CLA:HAB	2.03	0.41
18:6:164:PRO:HG2	22:6:308:CLA:H12	2.03	0.41
18:6:249:LEU:O	18:6:250:PHE:C	2.58	0.41
19:9:82:VAL:HG21	38:9:301:LUT:H401	2.03	0.41
19:9:111:LEU:HD21	39:9:314:CHL:ND	2.35	0.41
1:A:59:PHE:HB2	1:A:69:ILE:HG23	2.03	0.41
1:A:205:LEU:HD13	22:A:813:CLA:HHB	2.03	0.41
22:A:807:CLA:H91	22:A:807:CLA:H111	1.69	0.41
22:A:816:CLA:HBC2	22:3:316:CLA:CBB	2.51	0.41
22:A:821:CLA:O1A	22:A:824:CLA:HBD	2.21	0.41
22:A:828:CLA:H91	24:J:104:BCR:H11C	2.02	0.41
22:A:829:CLA:NC	24:A:845:BCR:HC42	2.35	0.41
22:A:838:CLA:H52	22:A:838:CLA:NC	2.36	0.41
24:A:847:BCR:H311	24:A:847:BCR:HC7	1.81	0.41
2:B:74:ASN:ND2	2:B:88:ILE:HG13	2.35	0.41
2:B:127:THR:HA	2:B:245:PHE:HZ	1.84	0.41
2:B:168:TRP:NE1	22:B:815:CLA:HAC2	2.36	0.41
2:B:302:MET:HB2	2:B:324:PHE:CE2	2.55	0.41
2:B:362:LEU:HD22	2:B:369:GLN:NE2	2.35	0.41
2:B:575:ASP:HA	2:B:578:TYR:HB3	2.02	0.41
22:B:812:CLA:H41	22:B:812:CLA:H62	1.88	0.41
22:B:821:CLA:H51	22:B:828:CLA:H101	2.03	0.41
22:B:826:CLA:H161	22:B:826:CLA:H122	1.67	0.41
22:B:827:CLA:HBA1	22:B:828:CLA:HED2	2.03	0.41
24:B:853:BCR:H312	22:L:202:CLA:C2C	2.50	0.41
5:E:53:TYR:C	5:E:55:GLN:H	2.25	0.41
34:F:308:LMG:H161	34:F:308:LMG:H132	1.83	0.41
22:J:103:CLA:HED1	15:8:240:LEU:HB2	2.03	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
22:K:205:CLA:NA	24:K:206:BCR:H281	2.36	0.41
38:1:303:LUT:H35	38:1:303:LUT:H401	1.84	0.41
39:1:312:CHL:H11	31:1:319:DGD:HD3	2.02	0.41
12:Z:92:GLU:OE1	12:Z:199:PRO:HD2	2.20	0.41
12:Z:92:GLU:OE1	12:Z:200:LEU:HD23	2.20	0.41
12:Z:103:PRO:HB2	39:Z:312:CHL:C1D	2.51	0.41
13:3:98:PRO:N	13:3:99:PRO:HD2	2.36	0.41
13:3:158:TYR:HB3	22:3:308:CLA:HED3	2.03	0.41
24:3:304:BCR:H383	22:3:313:CLA:CAB	2.51	0.41
22:3:309:CLA:CMD	22:3:314:CLA:H43	2.50	0.41
22:3:313:CLA:H13	22:5:320:CLA:C4B	2.51	0.41
14:7:58:LEU:CD1	22:7:308:CLA:H92	2.51	0.41
24:7:303:BCR:C29	42:7:321:SPH:H122	2.50	0.41
15:8:77:MET:SD	22:8:305:CLA:HAB	2.61	0.41
15:8:113:ILE:CG2	15:8:118:LEU:HB2	2.51	0.41
15:8:144:SER:OG	15:8:157:LYS:HD3	2.21	0.41
15:8:184:LEU:HB3	22:8:305:CLA:CMA	2.50	0.41
15:8:189:ILE:HD12	15:8:189:ILE:HA	1.62	0.41
22:8:305:CLA:H111	43:8:319:LPX:H19A	2.03	0.41
16:4:132:VAL:HG21	16:4:140:TYR:CD1	2.56	0.41
16:4:231:GLN:NE2	38:4:802:LUT:H42	2.35	0.41
16:4:251:TRP:HB3	18:6:108:PHE:HB3	2.02	0.41
38:4:803:LUT:H31	38:4:803:LUT:H391	1.91	0.41
22:4:807:CLA:HBB1	22:4:807:CLA:H171	2.02	0.41
22:4:811:CLA:HMC3	24:6:305:BCR:H313	2.03	0.41
18:6:61:PRO:HG2	18:6:62:GLU:OE1	2.21	0.41
18:6:215:ILE:HA	18:6:218:LYS:HG2	2.01	0.41
22:6:310:CLA:HBC1	25:6:323:LHG:H282	2.03	0.41
39:6:315:CHL:CBB	39:6:318:CHL:HBB2	2.51	0.41
38:9:302:LUT:H32	22:9:306:CLA:CAB	2.50	0.41
20:2:178:ARG:O	20:2:178:ARG:NE	2.52	0.41
20:2:194:TRP:O	20:2:200:THR:HG21	2.21	0.41
1:A:90:GLY:CA	22:A:807:CLA:HMC3	2.51	0.41
1:A:562:LEU:HB2	2:B:677:GLU:OE2	2.21	0.41
22:A:836:CLA:H42	22:K:203:CLA:C2D	2.51	0.41
22:A:840:CLA:H61	22:A:840:CLA:H2	1.75	0.41
2:B:53:GLY:O	2:B:57:ILE:HG13	2.21	0.41
24:B:847:BCR:C8	22:1:308:CLA:H92	2.51	0.41
3:C:21:CYS:HA	29:C:101:SF4:S4	2.61	0.41
6:F:79:ARG:HB2	6:F:79:ARG:NH1	2.36	0.41
10:K:59:ARG:NH2	10:K:67:LEU:HD22	2.36	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
24:K:206:BCR:H15C	24:K:206:BCR:H351	1.96	0.41
12:1:36:LEU:HD21	40:1:318:SQD:H161	2.03	0.41
12:1:167:ILE:O	12:1:171:LYS:HG3	2.20	0.41
22:1:306:CLA:H112	25:1:317:LHG:H312	2.02	0.41
22:1:307:CLA:H61	22:1:307:CLA:H92	1.76	0.41
12:Z:60:LEU:CD1	22:Z:306:CLA:H42	2.51	0.41
12:Z:77:HIS:HB3	12:Z:182:MET:SD	2.61	0.41
38:3:302:LUT:H31	38:3:302:LUT:H391	1.88	0.41
24:3:307:BCR:H10C	22:3:315:CLA:C1B	2.50	0.41
22:7:304:CLA:HMD2	22:7:314:CLA:CGA	2.50	0.41
39:7:313:CHL:OMC	39:7:313:CHL:HAC1	2.20	0.41
22:8:306:CLA:CMD	33:4:801:LMT:H61	2.51	0.41
39:8:315:CHL:HAC2	39:8:315:CHL:OMC	2.20	0.41
16:4:225:PHE:CE1	38:4:803:LUT:H10	2.56	0.41
18:6:99:ASP:CB	39:6:315:CHL:HED3	2.48	0.41
18:6:123:PHE:CG	24:6:305:BCR:H10C	2.55	0.41
18:6:157:PRO:HG2	22:6:307:CLA:CMD	2.51	0.41
18:6:199:PRO:HB2	38:6:303:LUT:H183	2.03	0.41
22:6:317:CLA:HBA2	22:6:317:CLA:H12	1.69	0.41
39:9:314:CHL:HHC	39:9:314:CHL:CBB	2.48	0.41
20:2:177:VAL:HG22	22:2:301:CLA:HHD	2.03	0.41
1:A:578:PRO:HB3	2:B:560:CYS:SG	2.60	0.40
22:A:807:CLA:HAA2	22:A:809:CLA:O2D	2.22	0.40
22:A:834:CLA:HED2	22:A:834:CLA:H2A	2.02	0.40
22:A:838:CLA:HMC2	22:A:838:CLA:H102	2.02	0.40
2:B:460:PHE:CD1	22:F:304:CLA:HMB3	2.56	0.40
31:B:852:DGD:HE5	31:B:852:DGD:HD4	2.03	0.40
4:D:117:ARG:HD2	4:D:119:GLU:HB3	2.03	0.40
4:D:192:MET:HG2	4:D:193:ASP:N	2.36	0.40
5:E:52:TRP:O	5:E:55:GLN:HB2	2.21	0.40
10:K:106:GLY:HA3	22:K:203:CLA:C1B	2.52	0.40
22:1:304:CLA:HMD3	22:1:313:CLA:CAD	2.50	0.40
22:Z:306:CLA:H102	22:Z:307:CLA:HMB3	2.02	0.40
22:7:310:CLA:HBC3	25:7:318:LHG:O8	2.21	0.40
22:7:323:CLA:HMB1	22:7:323:CLA:HBB1	2.03	0.40
22:8:311:CLA:CMA	33:4:801:LMT:H5'	2.51	0.40
16:4:150:PHE:O	16:4:153:MET:HG2	2.22	0.40
16:4:258:ASN:HD22	16:4:258:ASN:H	1.69	0.40
17:5:101:TYR:HD1	39:5:316:CHL:HED3	1.86	0.40
18:6:57:LEU:HD11	38:6:304:LUT:H221	2.01	0.40
18:6:60:ASN:OD1	18:6:61:PRO:HD2	2.21	0.40

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
18:6:163:VAL:HG22	22:6:307:CLA:HAA2	2.02	0.40
1:A:293:ASP:CB	22:A:818:CLA:HMA1	2.52	0.40
1:A:323:LYS:O	1:A:327:GLU:HG3	2.21	0.40
22:A:802:CLA:H12	22:A:802:CLA:HMA2	2.03	0.40
22:A:814:CLA:CMC	24:A:844:BCR:H352	2.51	0.40
22:A:821:CLA:H71	22:A:824:CLA:H71	2.01	0.40
22:A:827:CLA:H2	22:A:827:CLA:H61	1.69	0.40
22:A:828:CLA:H91	24:J:104:BCR:H341	2.03	0.40
22:A:833:CLA:H192	22:B:840:CLA:HHB	2.04	0.40
2:B:672:TRP:O	2:B:676:ILE:HG13	2.20	0.40
22:B:821:CLA:H193	22:B:826:CLA:H203	2.03	0.40
22:B:836:CLA:C1B	31:1:319:DGD:HAT2	2.51	0.40
5:E:39:ARG:NH2	5:E:62:VAL:H	2.17	0.40
6:F:69:PRO:HA	6:F:124:LEU:HD23	2.03	0.40
22:F:304:CLA:HAC1	33:F:307:LMT:H121	2.02	0.40
12:Z:82:MET:HE3	22:Z:303:CLA:HMC3	2.02	0.40
22:Z:303:CLA:H12	22:Z:303:CLA:HBA2	1.59	0.40
22:3:312:CLA:H111	22:3:312:CLA:H93	1.62	0.40
16:4:178:PRO:HD2	39:4:819:CHL:CHC	2.51	0.40
16:4:193:PRO:HG2	22:4:805:CLA:HMD1	2.03	0.40
18:6:74:HIS:ND1	22:6:317:CLA:HMD1	2.36	0.40
22:6:301:CLA:HHC	22:6:301:CLA:CBB	2.52	0.40
22:6:307:CLA:HMB2	22:6:308:CLA:HBA2	2.03	0.40
1:A:104:TRP:CD1	1:A:108:PRO:HA	2.56	0.40
1:A:651:TRP:HB2	22:B:805:CLA:CBC	2.51	0.40
22:A:815:CLA:H42	31:3:301:DGD:HAT2	2.03	0.40
22:A:831:CLA:NA	25:A:849:LHG:H111	2.37	0.40
24:A:844:BCR:H403	24:A:856:BCR:H10C	2.02	0.40
24:A:856:BCR:H392	10:K:95:LEU:CD2	2.52	0.40
2:B:258:PHE:CE1	22:B:819:CLA:H52	2.57	0.40
22:B:814:CLA:H111	19:9:54:LEU:HD11	2.04	0.40
22:B:833:CLA:H2	22:B:833:CLA:H61	1.77	0.40
10:K:38:MET:HG2	22:K:202:CLA:C2D	2.52	0.40
10:K:107:LEU:CD2	10:K:112:ASN:HD22	2.34	0.40
11:L:144:PRO:O	11:L:163:SER:HA	2.21	0.40
12:1:191:GLN:OE1	12:1:199:PRO:HA	2.20	0.40
12:Z:146:ASP:O	12:Z:147:ALA:HB3	2.20	0.40
14:7:129:LEU:HD13	22:7:322:CLA:O1A	2.22	0.40
15:8:228:THR:HA	22:8:307:CLA:HBA2	2.02	0.40
22:5:308:CLA:H101	22:5:308:CLA:H13	1.80	0.40
22:5:311:CLA:CHD	22:5:318:CLA:H101	2.50	0.40

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
18:6:110:ILE:HD13	24:6:306:BCR:C8	2.41	0.40
19:9:127:TYR:HB2	22:9:313:CLA:HMA3	2.03	0.40
1:A:121:VAL:HG13	1:A:122:VAL:HG22	2.03	0.40
1:A:310:HIS:O	22:A:822:CLA:HAC2	2.22	0.40
1:A:370:HIS:NE2	22:A:827:CLA:NB	2.70	0.40
1:A:571:PHE:O	1:A:587:SER:HB2	2.22	0.40
22:A:810:CLA:H62	22:A:810:CLA:H2	1.77	0.40
22:A:833:CLA:HBD	24:B:853:BCR:H12C	2.03	0.40
2:B:441:ASN:O	2:B:445:GLN:HG2	2.21	0.40
3:C:22:PRO:HD2	29:C:101:SF4:S3	2.61	0.40
3:C:41:SER:HB2	4:D:168:VAL:H	1.87	0.40
4:D:83:PHE:CZ	4:D:141:VAL:HB	2.56	0.40
6:F:107:THR:O	6:F:110:ARG:HG2	2.21	0.40
6:F:185:ILE:HG13	6:F:186:ILE:H	1.87	0.40
9:J:25:LEU:O	9:J:29:ILE:HG13	2.22	0.40
12:1:144:GLU:HB2	22:1:313:CLA:HAC1	2.03	0.40
12:1:170:LEU:HB3	22:1:304:CLA:HMA1	2.02	0.40
38:1:302:LUT:H371	22:1:304:CLA:H92	2.04	0.40
38:1:303:LUT:H15	38:1:303:LUT:H201	1.86	0.40
22:1:308:CLA:ND	22:1:314:CLA:H93	2.37	0.40
12:Z:77:HIS:CE1	22:Z:314:CLA:HMD1	2.57	0.40
22:Z:305:CLA:CGA	22:Z:305:CLA:C1A	2.99	0.40
22:3:320:CLA:HBA1	41:7:319:3PH:C3D	2.51	0.40
14:7:37:ASN:HD22	14:7:37:ASN:HA	1.57	0.40
16:4:138:GLY:HA3	39:4:813:CHL:C3D	2.51	0.40
16:4:186:PRO:HD2	16:4:195:GLY:HA3	2.04	0.40
38:4:802:LUT:H7	38:4:802:LUT:H171	1.85	0.40
24:4:804:BCR:C27	22:4:810:CLA:H2	2.44	0.40
22:4:807:CLA:H93	22:4:807:CLA:H61	1.90	0.40
22:4:809:CLA:HMC1	39:4:813:CHL:CBB	2.52	0.40
17:5:69:TRP:HZ2	22:5:318:CLA:HAA2	1.86	0.40
38:5:303:LUT:H193	22:5:319:CLA:HBB2	2.03	0.40
24:5:304:BCR:H271	22:5:312:CLA:C1B	2.51	0.40
22:5:311:CLA:HMB1	22:5:311:CLA:CBB	2.52	0.40
22:5:326:CLA:C1D	22:6:314:CLA:H13	2.52	0.40
18:6:49:ASP:HA	22:6:310:CLA:O1D	2.21	0.40
18:6:54:PRO:HD2	38:6:304:LUT:C23	2.40	0.40
38:9:301:LUT:H162	22:9:305:CLA:CHB	2.51	0.40
38:9:302:LUT:H371	22:9:306:CLA:H142	2.03	0.40
1:A:161:THR:HG23	27:A:852:DGA:HAW2	2.03	0.40
1:A:247:ARG:HD3	1:A:257:PHE:O	2.21	0.40

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:457:ILE:HG21	22:B:806:CLA:H92	2.03	0.40
1:A:494:ALA:N	1:A:495:PRO:CD	2.84	0.40
1:A:645:TRP:O	1:A:649:PHE:HB3	2.22	0.40
22:A:804:CLA:HBA1	22:A:811:CLA:C1D	2.52	0.40
22:A:809:CLA:H42	24:J:104:BCR:C10	2.52	0.40
22:A:813:CLA:HBA1	22:A:825:CLA:C4	2.49	0.40
22:A:822:CLA:H8	22:A:823:CLA:H93	2.02	0.40
22:A:822:CLA:C9	22:K:204:CLA:HAB	2.52	0.40
22:A:836:CLA:HED3	22:A:836:CLA:H2A	2.03	0.40
2:B:263:HIS:CG	2:B:264:PRO:HD2	2.56	0.40
2:B:333:PHE:CD2	24:B:848:BCR:H292	2.56	0.40
2:B:581:VAL:HG21	2:B:711:LEU:HD21	2.03	0.40
2:B:695:LYS:HE2	11:L:141:GLN:HE22	1.86	0.40
22:B:819:CLA:ND	22:B:819:CLA:H8	2.36	0.40
22:B:823:CLA:HMD2	24:B:844:BCR:C7	2.40	0.40
22:B:825:CLA:H162	22:B:825:CLA:HMB2	2.02	0.40
3:C:10:THR:O	3:C:64:SER:HB2	2.21	0.40
4:D:76:ARG:HA	4:D:79:GLN:OE1	2.21	0.40
4:D:128:LEU:HB3	4:D:134:PHE:HB2	2.03	0.40
10:K:84:PHE:CD2	10:K:89:THR:HG22	2.57	0.40
40:1:318:SQD:H171	40:1:318:SQD:H141	1.92	0.40
13:3:22:THR:HA	13:3:25:TYR:CZ	2.56	0.40
13:3:76:ALA:CB	24:3:305:BCR:H292	2.52	0.40
13:3:169:LEU:HD12	38:3:302:LUT:H221	2.03	0.40
24:3:304:BCR:H19C	22:3:319:CLA:HBA1	2.03	0.40
22:3:318:CLA:H13	22:3:318:CLA:H102	1.89	0.40
14:7:155:GLY:O	14:7:157:THR:HG23	2.22	0.40
22:7:305:CLA:ND	22:7:310:CLA:H8	2.36	0.40
15:8:164:PRO:HG3	22:8:313:CLA:C1D	2.52	0.40
24:4:804:BCR:H282	22:4:818:CLA:CMC	2.51	0.40
38:5:302:LUT:H30	22:5:307:CLA:H52	2.01	0.40
22:5:308:CLA:OBD	22:5:313:CLA:H12	2.22	0.40
18:6:67:PHE:HB2	22:6:310:CLA:HMA1	2.04	0.40
18:6:236:PHE:CE2	18:6:245:PHE:HB2	2.56	0.40
22:6:309:CLA:H51	22:6:309:CLA:C1C	2.52	0.40

There are no symmetry-related clashes.

5.3 Torsion angles (i)

5.3.1 Protein backbone (i)

In the following table, the Percentiles column shows the percent Ramachandran outliers of the chain as a percentile score with respect to all PDB entries followed by that with respect to all EM entries.

The Analysed column shows the number of residues for which the backbone conformation was analysed, and the total number of residues.

Mol	Chain	Analysed	Favoured	Allowed	Outliers	Percentiles
1	A	738/740 (100%)	685 (93%)	51 (7%)	2 (0%)	41 70
2	B	730/732 (100%)	683 (94%)	42 (6%)	5 (1%)	22 51
3	C	78/80 (98%)	72 (92%)	6 (8%)	0	100 100
4	D	140/142 (99%)	124 (89%)	13 (9%)	3 (2%)	7 22
5	E	59/61 (97%)	50 (85%)	8 (14%)	1 (2%)	9 27
6	F	163/165 (99%)	145 (89%)	14 (9%)	4 (2%)	5 18
7	G	91/93 (98%)	73 (80%)	16 (18%)	2 (2%)	6 21
8	I	39/41 (95%)	34 (87%)	5 (13%)	0	100 100
9	J	38/40 (95%)	38 (100%)	0	0	100 100
10	K	87/89 (98%)	78 (90%)	8 (9%)	1 (1%)	14 39
11	L	119/123 (97%)	106 (89%)	11 (9%)	2 (2%)	9 27
12	1	192/194 (99%)	165 (86%)	23 (12%)	4 (2%)	7 22
12	Z	192/194 (99%)	181 (94%)	10 (5%)	1 (0%)	29 59
13	3	216/218 (99%)	195 (90%)	18 (8%)	3 (1%)	11 32
14	7	218/220 (99%)	202 (93%)	12 (6%)	4 (2%)	8 26
15	8	212/214 (99%)	181 (85%)	24 (11%)	7 (3%)	4 12
16	4	205/207 (99%)	190 (93%)	15 (7%)	0	100 100
17	5	224/226 (99%)	207 (92%)	13 (6%)	4 (2%)	8 26
18	6	224/226 (99%)	212 (95%)	11 (5%)	1 (0%)	34 64
19	9	182/184 (99%)	152 (84%)	23 (13%)	7 (4%)	3 10
20	2	68/70 (97%)	53 (78%)	14 (21%)	1 (2%)	10 31
All	All	4215/4259 (99%)	3826 (91%)	337 (8%)	52 (1%)	17 37

All (52) Ramachandran outliers are listed below:

Mol	Chain	Res	Type
4	D	154	ASP
12	1	41	ALA
12	1	42	PRO
14	7	219	ILE
15	8	38	ALA
15	8	148	SER
15	8	152	PHE
15	8	161	ASN
19	9	46	ALA
19	9	143	PRO
19	9	146	PRO
19	9	147	ALA
20	2	182	PRO
1	A	21	PRO
2	B	28	VAL
6	F	64	ILE
6	F	150	PHE
12	1	166	ASP
15	8	42	GLY
19	9	139	ILE
6	F	178	ALA
6	F	212	ALA
11	L	61	SER
13	3	87	VAL
13	3	159	PRO
14	7	213	ALA
15	8	37	PRO
19	9	196	TYR
2	B	230	ASN
7	G	70	PRO
7	G	85	ALA
12	1	102	ALA
12	Z	40	ASP
14	7	98	PRO
14	7	240	SER
15	8	150	LEU
18	6	88	GLN
19	9	145	ASP
1	A	574	PRO
4	D	149	TYR
13	3	218	ALA
17	5	152	PHE
2	B	160	PRO

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Mol	Chain	Res	Type
2	B	545	SER
2	B	559	PRO
17	5	236	THR
4	D	158	PRO
5	E	70	PRO
11	L	98	GLY
17	5	171	PRO
10	K	60	GLY
17	5	151	ILE

5.3.2 Protein sidechains (i)

In the following table, the Percentiles column shows the percent sidechain outliers of the chain as a percentile score with respect to all PDB entries followed by that with respect to all EM entries.

The Analysed column shows the number of residues for which the sidechain conformation was analysed, and the total number of residues.

Mol	Chain	Analysed	Rotameric	Outliers	Percentiles
1	A	602/602 (100%)	576 (96%)	26 (4%)	29 60
2	B	596/596 (100%)	566 (95%)	30 (5%)	24 55
3	C	70/70 (100%)	67 (96%)	3 (4%)	29 60
4	D	122/122 (100%)	109 (89%)	13 (11%)	6 19
5	E	54/54 (100%)	53 (98%)	1 (2%)	57 84
6	F	128/128 (100%)	107 (84%)	21 (16%)	2 6
7	G	69/69 (100%)	62 (90%)	7 (10%)	7 21
8	I	33/33 (100%)	33 (100%)	0	100 100
9	J	35/35 (100%)	35 (100%)	0	100 100
10	K	60/60 (100%)	53 (88%)	7 (12%)	5 16
11	L	90/90 (100%)	82 (91%)	8 (9%)	9 27
12	1	144/144 (100%)	125 (87%)	19 (13%)	4 11
12	Z	144/144 (100%)	134 (93%)	10 (7%)	15 40
13	3	173/173 (100%)	158 (91%)	15 (9%)	10 28
14	7	169/169 (100%)	156 (92%)	13 (8%)	13 34
15	8	162/162 (100%)	140 (86%)	22 (14%)	3 11
16	4	162/162 (100%)	146 (90%)	16 (10%)	8 22

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Mol	Chain	Analysed	Rotameric	Outliers	Percentiles
17	5	185/185 (100%)	168 (91%)	17 (9%)	9 26
18	6	184/184 (100%)	172 (94%)	12 (6%)	17 43
19	9	143/143 (100%)	129 (90%)	14 (10%)	8 23
20	2	57/57 (100%)	50 (88%)	7 (12%)	4 14
All	All	3382/3382 (100%)	3121 (92%)	261 (8%)	16 34

All (261) residues with a non-rotameric sidechain are listed below:

Mol	Chain	Res	Type
1	A	14	LYS
1	A	17	VAL
1	A	36	SER
1	A	37	ARG
1	A	52	LEU
1	A	55	ASP
1	A	68	GLU
1	A	116	GLN
1	A	144	PHE
1	A	154	THR
1	A	189	GLU
1	A	244	LEU
1	A	248	GLN
1	A	278	PHE
1	A	291	LEU
1	A	299	LEU
1	A	301	ILE
1	A	363	LEU
1	A	372	TYR
1	A	426	LEU
1	A	455	LEU
1	A	474	ASP
1	A	489	LYS
1	A	490	THR
1	A	503	LEU
1	A	642	ILE
2	B	20	ARG
2	B	28	VAL
2	B	33	GLU
2	B	54	GLN
2	B	60	LEU
2	B	97	PHE

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Mol	Chain	Res	Type
2	B	103	GLU
2	B	159	GLN
2	B	186	VAL
2	B	206	LYS
2	B	236	GLN
2	B	258	PHE
2	B	321	LYS
2	B	345	ILE
2	B	355	SER
2	B	362	LEU
2	B	379	ILE
2	B	384	MET
2	B	394	PHE
2	B	398	ASP
2	B	438	TYR
2	B	484	SER
2	B	495	LEU
2	B	513	ILE
2	B	537	LYS
2	B	565	ARG
2	B	632	LEU
2	B	644	LEU
2	B	695	LYS
2	B	718	TYR
3	C	35	LYS
3	C	62	PHE
3	C	69	LEU
4	D	61	ASP
4	D	65	PRO
4	D	89	GLU
4	D	90	SER
4	D	92	LYS
4	D	97	GLU
4	D	108	LYS
4	D	117	ARG
4	D	158	PRO
4	D	159	GLU
4	D	160	LYS
4	D	162	ASN
4	D	192	MET
5	E	38	LYS
6	F	63	ASP

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Mol	Chain	Res	Type
6	F	80	LEU
6	F	90	ARG
6	F	93	ASN
6	F	97	ASP
6	F	100	PRO
6	F	104	LEU
6	F	109	GLU
6	F	113	ASN
6	F	115	PHE
6	F	125	CYS
6	F	159	VAL
6	F	173	MET
6	F	179	LYS
6	F	189	VAL
6	F	210	LEU
6	F	211	ARG
6	F	214	THR
6	F	217	GLU
6	F	218	LYS
6	F	220	GLU
7	G	37	GLN
7	G	45	MET
7	G	56	LEU
7	G	62	GLU
7	G	88	ILE
7	G	101	ASP
7	G	122	LEU
10	K	25	ARG
10	K	29	PHE
10	K	56	THR
10	K	61	THR
10	K	66	LYS
10	K	70	ARG
10	K	89	THR
11	L	60	THR
11	L	76	ARG
11	L	82	LEU
11	L	83	LEU
11	L	141	GLN
11	L	161	LEU
11	L	176	VAL
11	L	179	LEU

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Mol	Chain	Res	Type
12	1	35	TRP
12	1	37	PRO
12	1	42	PRO
12	1	47	ASP
12	1	61	CYS
12	1	113	THR
12	1	119	VAL
12	1	122	ASP
12	1	140	ARG
12	1	149	LYS
12	1	151	ILE
12	1	167	ILE
12	1	168	GLU
12	1	172	LEU
12	1	175	ILE
12	1	183	MET
12	1	198	THR
12	1	214	ASN
12	1	218	ASN
12	Z	35	TRP
12	Z	37	PRO
12	Z	47	ASP
12	Z	61	CYS
12	Z	113	THR
12	Z	115	PHE
12	Z	119	VAL
12	Z	172	LEU
12	Z	191	GLN
12	Z	224	PHE
13	3	92	PHE
13	3	93	GLU
13	3	94	THR
13	3	96	VAL
13	3	108	THR
13	3	109	ASP
13	3	126	LEU
13	3	141	GLN
13	3	149	VAL
13	3	163	TRP
13	3	166	LEU
13	3	172	LYS
13	3	177	MET

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Mol	Chain	Res	Type
13	3	178	LYS
13	3	228	ASN
14	7	34	TYR
14	7	35	PRO
14	7	37	ASN
14	7	65	MET
14	7	97	LEU
14	7	129	LEU
14	7	165	ASN
14	7	182	ASP
14	7	185	MET
14	7	196	ASN
14	7	219	ILE
14	7	220	GLN
14	7	243	PHE
15	8	41	LYS
15	8	91	LYS
15	8	95	LEU
15	8	110	GLN
15	8	115	LEU
15	8	138	LEU
15	8	149	PHE
15	8	152	PHE
15	8	153	GLU
15	8	157	LYS
15	8	173	MET
15	8	174	CYS
15	8	178	GLU
15	8	180	LYS
15	8	183	ASP
15	8	185	LYS
15	8	189	ILE
15	8	215	ASP
15	8	231	ASP
15	8	234	ILE
15	8	238	ILE
15	8	240	LEU
16	4	58	ARG
16	4	61	TRP
16	4	66	LYS
16	4	94	LYS
16	4	101	LEU

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Mol	Chain	Res	Type
16	4	141	GLU
16	4	160	VAL
16	4	163	ARG
16	4	167	ASP
16	4	175	ASN
16	4	204	LYS
16	4	207	MET
16	4	208	GLU
16	4	210	LEU
16	4	212	LEU
16	4	257	SER
17	5	32	ARG
17	5	34	LEU
17	5	39	VAL
17	5	40	ASP
17	5	85	VAL
17	5	95	ARG
17	5	97	ASP
17	5	98	VAL
17	5	138	GLN
17	5	139	ASP
17	5	141	LYS
17	5	146	VAL
17	5	153	LYS
17	5	176	LYS
17	5	182	LEU
17	5	221	ASN
17	5	236	THR
18	6	43	ASN
18	6	89	GLU
18	6	92	LYS
18	6	110	ILE
18	6	124	VAL
18	6	128	ARG
18	6	129	TRP
18	6	149	LEU
18	6	167	MET
18	6	211	MET
18	6	222	VAL
18	6	251	GLN
19	9	45	MET
19	9	64	LYS

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Mol	Chain	Res	Type
19	9	89	LEU
19	9	106	VAL
19	9	115	GLU
19	9	143	PRO
19	9	146	PRO
19	9	153	ASP
19	9	159	VAL
19	9	177	LEU
19	9	180	ARG
19	9	189	LYS
19	9	197	ASN
19	9	206	GLN
20	2	162	ARG
20	2	165	MET
20	2	168	PHE
20	2	178	ARG
20	2	187	LYS
20	2	222	GLU
20	2	229	LYS

Sometimes sidechains can be flipped to improve hydrogen bonding and reduce clashes. All (71) such sidechains are listed below:

Mol	Chain	Res	Type
1	A	80	GLN
1	A	136	GLN
1	A	219	HIS
1	A	248	GLN
1	A	279	ASN
1	A	296	HIS
1	A	338	HIS
1	A	442	ASN
1	A	637	GLN
1	A	659	GLN
1	A	676	HIS
1	A	722	GLN
2	B	11	GLN
2	B	30	HIS
2	B	35	HIS
2	B	42	ASN
2	B	72	GLN
2	B	90	HIS
2	B	115	ASN

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Mol	Chain	Res	Type
2	B	216	GLN
2	B	221	GLN
2	B	300	HIS
2	B	364	ASN
2	B	369	GLN
2	B	445	GLN
2	B	491	GLN
2	B	655	HIS
4	D	120	GLN
4	D	162	ASN
4	D	180	ASN
4	D	189	ASN
5	E	78	GLN
5	E	94	GLN
6	F	82	ASN
6	F	132	HIS
6	F	145	HIS
6	F	205	GLN
7	G	57	HIS
7	G	92	ASN
8	I	103	ASN
9	J	30	ASN
10	K	35	ASN
10	K	112	ASN
11	L	141	GLN
12	1	209	ASN
12	Z	191	GLN
13	3	141	GLN
13	3	201	GLN
13	3	222	ASN
13	3	223	ASN
14	7	37	ASN
14	7	132	HIS
14	7	196	ASN
15	8	110	GLN
15	8	205	GLN
16	4	231	GLN
16	4	233	GLN
16	4	242	ASN
16	4	258	ASN
17	5	108	ASN
17	5	114	GLN

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Mol	Chain	Res	Type
17	5	230	ASN
18	6	122	HIS
18	6	134	ASN
18	6	147	ASN
18	6	235	GLN
19	9	40	HIS
19	9	87	GLN
19	9	128	GLN
19	9	201	ASN
19	9	206	GLN

5.3.3 RNA

There are no RNA molecules in this entry.

5.4 Non-standard residues in protein, DNA, RNA chains

There are no non-standard protein/DNA/RNA residues in this entry.

5.5 Carbohydrates

There are no monosaccharides in this entry.

5.6 Ligand geometry

Of 332 ligands modelled in this entry, 1 is monoatomic - leaving 331 for Mogul analysis.

In the following table, the Counts columns list the number of bonds (or angles) for which Mogul statistics could be retrieved, the number of bonds (or angles) that are observed in the model and the number of bonds (or angles) that are defined in the Chemical Component Dictionary. The Link column lists molecule types, if any, to which the group is linked. The Z score for a bond length (or angle) is the number of standard deviations the observed value is removed from the expected value. A bond length (or angle) with $|Z| > 2$ is considered an outlier worth inspection. RMSZ is the root-mean-square of all Z scores of the bond lengths (or angles).

Mol	Type	Chain	Res	Link	Bond lengths			Bond angles		
					Counts	RMSZ	$\#\|Z\| > 2$	Counts	RMSZ	$\#\|Z\| > 2$
22	CLA	5	315	17	65,73,73	1.34	7 (10%)	76,113,113	1.97	17 (22%)
22	CLA	3	319	-	55,63,73	1.45	7 (12%)	64,101,113	2.12	16 (25%)
22	CLA	A	812	-	65,73,73	1.33	8 (12%)	76,113,113	1.97	17 (22%)

Mol	Type	Chain	Res	Link	Bond lengths			Bond angles		
					Counts	RMSZ	$\#\|Z\|>2$	Counts	RMSZ	$\#\|Z\|>2$
22	CLA	9	308	-	50,58,73	1.53	7 (14%)	58,95,113	2.21	18 (31%)
22	CLA	G	201	-	50,58,73	1.53	6 (12%)	58,95,113	2.25	18 (31%)
42	SPH	7	321	-	19,20,20	0.65	1 (5%)	18,21,21	1.15	0
22	CLA	7	314	44	50,58,73	1.51	6 (12%)	58,95,113	2.25	17 (29%)
22	CLA	6	312	-	65,73,73	1.34	8 (12%)	76,113,113	2.00	18 (23%)
22	CLA	Z	304	12	45,53,73	1.61	7 (15%)	52,89,113	2.07	12 (23%)
26	NKP	A	851	-	22,22,28	0.37	0	24,26,32	0.39	0
39	CHL	8	315	44	66,74,74	0.81	2 (3%)	73,114,114	1.29	11 (15%)
39	CHL	5	316	44	66,74,74	0.85	3 (4%)	73,114,114	1.23	12 (16%)
22	CLA	A	831	-	50,58,73	1.51	7 (14%)	58,95,113	2.22	17 (29%)
22	CLA	5	311	-	55,63,73	1.45	8 (14%)	64,101,113	2.11	18 (28%)
22	CLA	A	854	-	65,73,73	1.33	7 (10%)	76,113,113	1.94	17 (22%)
22	CLA	B	816	-	60,68,73	1.38	7 (11%)	70,107,113	2.04	17 (24%)
22	CLA	J	103	9	42,50,73	1.65	7 (16%)	48,85,113	2.22	15 (31%)
22	CLA	3	308	13	65,73,73	1.34	8 (12%)	76,113,113	2.02	18 (23%)
24	BCR	B	853	-	41,41,41	4.83	27 (65%)	56,56,56	2.40	22 (39%)
39	CHL	6	318	-	56,64,74	0.91	2 (3%)	61,102,114	1.40	12 (19%)
24	BCR	A	844	-	41,41,41	4.78	27 (65%)	56,56,56	3.12	23 (41%)
25	LHG	8	317	22	37,37,48	0.43	0	40,43,54	1.03	2 (5%)
22	CLA	A	817	-	60,68,73	1.39	9 (15%)	70,107,113	2.08	16 (22%)
22	CLA	B	806	44	65,73,73	1.35	7 (10%)	76,113,113	1.99	16 (21%)
38	LUT	3	303	-	42,43,43	6.06	28 (66%)	51,60,60	2.00	13 (25%)
39	CHL	9	314	-	42,50,74	1.13	4 (9%)	44,85,114	1.53	11 (25%)
22	CLA	A	855	44	65,73,73	1.34	7 (10%)	76,113,113	1.95	16 (21%)
22	CLA	B	808	-	65,73,73	1.33	6 (9%)	76,113,113	1.98	18 (23%)
22	CLA	A	814	-	60,68,73	1.38	6 (10%)	70,107,113	2.10	19 (27%)
39	CHL	4	814	44	51,59,74	0.90	2 (3%)	55,96,114	1.47	13 (23%)
22	CLA	A	818	-	60,68,73	1.38	7 (11%)	70,107,113	2.12	17 (24%)
41	3PH	5	325	-	22,22,47	1.22	4 (18%)	26,27,52	1.28	2 (7%)
25	LHG	4	820	22	48,48,48	0.42	0	51,54,54	1.04	4 (7%)
22	CLA	5	318	17	65,73,73	1.32	7 (10%)	76,113,113	1.95	17 (22%)
22	CLA	Z	315	-	46,54,73	1.59	7 (15%)	53,90,113	2.15	13 (24%)
41	3PH	7	319	-	38,38,47	0.94	4 (10%)	42,43,52	1.13	2 (4%)
24	BCR	3	307	-	41,41,41	4.78	27 (65%)	56,56,56	3.25	27 (48%)
22	CLA	A	815	-	65,73,73	1.32	6 (9%)	76,113,113	2.00	17 (22%)

Mol	Type	Chain	Res	Link	Bond lengths			Bond angles		
					Counts	RMSZ	$\#\|Z\|>2$	Counts	RMSZ	$\#\|Z\|>2$
22	CLA	4	805	16	60,68,73	1.42	8 (13%)	70,107,113	1.93	16 (22%)
28	OCA	A	853	-	9,9,9	0.70	0	9,9,9	0.70	0
22	CLA	A	804	22	45,53,73	1.59	7 (15%)	52,89,113	2.16	15 (28%)
22	CLA	3	318	13	60,68,73	1.38	7 (11%)	70,107,113	2.06	18 (25%)
22	CLA	4	807	16	65,73,73	1.33	8 (12%)	76,113,113	2.03	16 (21%)
35	T7X	J	102	-	49,49,61	0.91	4 (8%)	59,61,73	0.94	3 (5%)
22	CLA	A	807	-	60,68,73	1.39	7 (11%)	70,107,113	2.04	15 (21%)
34	LMG	F	308	-	35,35,55	0.46	0	43,43,63	1.38	5 (11%)
22	CLA	7	306	-	65,73,73	1.34	7 (10%)	76,113,113	2.03	17 (22%)
24	BCR	A	856	-	41,41,41	4.77	27 (65%)	56,56,56	2.50	23 (41%)
22	CLA	2	302	-	45,53,73	1.60	7 (15%)	52,89,113	2.19	15 (28%)
22	CLA	7	311	-	43,51,73	1.63	7 (16%)	49,86,113	2.20	14 (28%)
22	CLA	1	311	-	46,54,73	1.58	8 (17%)	53,90,113	2.22	12 (22%)
22	CLA	1	314	12	65,73,73	1.32	7 (10%)	76,113,113	1.95	16 (21%)
37	DAO	K	201	-	13,13,13	0.59	0	13,13,13	0.56	0
24	BCR	7	303	-	41,41,41	4.78	27 (65%)	56,56,56	2.39	21 (37%)
22	CLA	6	308	18	52,60,73	1.47	6 (11%)	60,97,113	2.27	20 (33%)
22	CLA	B	821	44	65,73,73	1.34	7 (10%)	76,113,113	1.96	16 (21%)
22	CLA	A	805	-	65,73,73	1.33	9 (13%)	76,113,113	1.98	17 (22%)
22	CLA	B	810	-	56,64,73	1.43	6 (10%)	65,102,113	2.18	20 (30%)
22	CLA	A	819	-	65,73,73	1.33	7 (10%)	76,113,113	1.95	17 (22%)
22	CLA	8	311	25	50,58,73	1.53	7 (14%)	58,95,113	2.18	15 (25%)
24	BCR	I	4001	-	41,41,41	4.78	27 (65%)	56,56,56	2.58	19 (33%)
33	LMT	4	822	-	36,36,36	0.45	0	47,47,47	1.17	3 (6%)
38	LUT	4	802	-	42,43,43	6.08	28 (66%)	51,60,60	2.02	12 (23%)
22	CLA	7	317	-	58,66,73	1.41	7 (12%)	67,104,113	2.06	16 (23%)
39	CHL	7	313	44	54,62,74	0.92	3 (5%)	58,99,114	1.36	12 (20%)
22	CLA	A	829	-	65,73,73	1.34	7 (10%)	76,113,113	1.92	15 (19%)
22	CLA	5	319	44	55,63,73	1.45	7 (12%)	64,101,113	2.12	17 (26%)
22	CLA	A	803	-	65,73,73	1.34	7 (10%)	76,113,113	1.99	17 (22%)
22	CLA	B	801	-	65,73,73	1.33	7 (10%)	76,113,113	1.91	17 (22%)
22	CLA	B	824	-	59,67,73	1.40	7 (11%)	68,105,113	2.12	18 (26%)
22	CLA	6	301	44	60,68,73	1.40	8 (13%)	70,107,113	2.06	18 (25%)
22	CLA	5	310	17	65,73,73	1.33	7 (10%)	76,113,113	2.00	17 (22%)
39	CHL	6	316	-	51,59,74	0.96	3 (5%)	55,96,114	1.49	12 (21%)

Mol	Type	Chain	Res	Link	Bond lengths			Bond angles		
					Counts	RMSZ	$\#\|Z\|>2$	Counts	RMSZ	$\#\|Z\|>2$
22	CLA	3	310	13	65,73,73	1.34	8 (12%)	76,113,113	2.06	20 (26%)
40	SQD	1	318	-	47,48,54	0.83	0	56,59,65	0.96	3 (5%)
39	CHL	5	317	44	51,59,74	0.90	2 (3%)	55,96,114	1.42	13 (23%)
22	CLA	A	810	-	65,73,73	1.33	8 (12%)	76,113,113	1.98	17 (22%)
38	LUT	1	302	-	42,43,43	6.10	28 (66%)	51,60,60	2.08	12 (23%)
24	BCR	B	847	-	41,41,41	4.78	27 (65%)	56,56,56	2.46	21 (37%)
22	CLA	4	806	16	52,60,73	1.51	8 (15%)	60,97,113	2.23	18 (30%)
39	CHL	3	317	-	66,74,74	0.84	3 (4%)	73,114,114	1.29	9 (12%)
39	CHL	Z	311	12	51,59,74	0.98	3 (5%)	55,96,114	1.53	13 (23%)
38	LUT	3	302	-	42,43,43	6.05	28 (66%)	51,60,60	2.02	13 (25%)
22	CLA	6	317	18	50,58,73	1.52	6 (12%)	58,95,113	2.24	16 (27%)
24	BCR	K	206	-	41,41,41	4.79	27 (65%)	56,56,56	2.50	24 (42%)
22	CLA	4	818	44	51,59,73	1.52	7 (13%)	59,96,113	2.26	19 (32%)
25	LHG	5	324	22	36,36,48	0.44	0	39,42,54	1.12	3 (7%)
22	CLA	1	310	25	60,68,73	1.38	8 (13%)	70,107,113	2.04	16 (22%)
25	LHG	B	850	22	22,22,48	0.56	0	25,28,54	1.26	2 (8%)
22	CLA	B	841	-	65,73,73	1.34	7 (10%)	76,113,113	2.02	17 (22%)
22	CLA	5	309	-	56,64,73	1.45	8 (14%)	65,102,113	2.23	18 (27%)
22	CLA	5	312	44	50,58,73	1.52	7 (14%)	58,95,113	2.19	16 (27%)
25	LHG	A	850	-	48,48,48	0.39	0	51,54,54	1.04	3 (5%)
24	BCR	4	804	-	41,41,41	4.82	26 (63%)	56,56,56	2.25	20 (35%)
22	CLA	A	841	-	65,73,73	1.34	7 (10%)	76,113,113	1.99	17 (22%)
25	LHG	1	317	22	42,42,48	0.45	0	45,48,54	1.18	3 (6%)
22	CLA	B	809	2	65,73,73	1.33	8 (12%)	76,113,113	1.97	16 (21%)
22	CLA	8	313	44	50,58,73	1.52	7 (14%)	58,95,113	2.22	16 (27%)
22	CLA	4	812	-	55,63,73	1.46	7 (12%)	64,101,113	2.12	17 (26%)
27	DGA	A	852	-	39,39,43	1.14	3 (7%)	41,41,45	1.32	3 (7%)
22	CLA	A	824	-	65,73,73	1.34	7 (10%)	76,113,113	1.94	17 (22%)
22	CLA	B	805	-	65,73,73	1.33	7 (10%)	76,113,113	1.97	18 (23%)
33	LMT	1	301	-	36,36,36	0.37	0	47,47,47	0.84	0
36	C7Z	5	306	-	43,43,43	5.32	27 (62%)	58,60,60	2.37	22 (37%)
22	CLA	4	815	16	50,58,73	1.51	6 (12%)	58,95,113	2.24	15 (25%)
24	BCR	B	848	-	41,41,41	4.79	27 (65%)	56,56,56	6.79	24 (42%)
22	CLA	4	808	16	60,68,73	1.39	9 (15%)	70,107,113	2.04	17 (24%)
24	BCR	J	104	-	41,41,41	4.76	27 (65%)	56,56,56	2.38	20 (35%)

Mol	Type	Chain	Res	Link	Bond lengths			Bond angles		
					Counts	RMSZ	$\#\|Z\|>2$	Counts	RMSZ	$\#\|Z\|>2$
22	CLA	A	809	1	55,63,73	1.45	7 (12%)	64,101,113	2.12	19 (29%)
22	CLA	Z	310	-	50,58,73	1.53	8 (16%)	58,95,113	2.17	17 (29%)
22	CLA	6	314	18	65,73,73	1.34	7 (10%)	76,113,113	1.95	16 (21%)
22	CLA	B	829	-	57,65,73	1.43	7 (12%)	66,103,113	2.11	18 (27%)
22	CLA	B	815	-	65,73,73	1.34	6 (9%)	76,113,113	2.05	18 (23%)
22	CLA	A	820	-	60,68,73	1.40	7 (11%)	70,107,113	2.00	17 (24%)
29	SF4	B	802	-	0,12,12	-	-	-		
22	CLA	5	301	44	56,64,73	1.43	7 (12%)	65,102,113	2.11	17 (26%)
22	CLA	7	322	-	55,63,73	1.44	6 (10%)	64,101,113	2.11	18 (28%)
38	LUT	Z	301	-	42,43,43	6.07	27 (64%)	51,60,60	1.99	14 (27%)
42	SPH	7	320	-	19,20,20	0.63	0	18,21,21	1.10	0
22	CLA	A	806	1	65,73,73	1.34	7 (10%)	76,113,113	1.95	16 (21%)
22	CLA	B	831	-	58,66,73	1.41	8 (13%)	67,104,113	2.12	17 (25%)
22	CLA	Z	305	-	56,64,73	1.43	7 (12%)	65,102,113	2.09	15 (23%)
25	LHG	B	803	-	32,32,48	0.46	0	35,38,54	1.18	3 (8%)
22	CLA	A	842	25	52,60,73	1.49	8 (15%)	60,97,113	2.22	17 (28%)
22	CLA	3	315	-	45,53,73	1.66	8 (17%)	52,89,113	2.16	12 (23%)
24	BCR	B	844	-	41,41,41	4.79	26 (63%)	56,56,56	2.77	24 (42%)
22	CLA	1	305	12	45,53,73	1.61	8 (17%)	52,89,113	2.15	16 (30%)
24	BCR	A	847	-	41,41,41	4.80	27 (65%)	56,56,56	2.38	20 (35%)
22	CLA	Z	307	-	55,63,73	1.46	7 (12%)	64,101,113	2.18	17 (26%)
38	LUT	7	301	-	42,43,43	6.06	28 (66%)	51,60,60	1.95	13 (25%)
22	CLA	4	817	16	41,49,73	1.67	7 (17%)	47,84,113	2.26	15 (31%)
31	DGD	3	301	-	52,52,67	0.89	2 (3%)	66,66,81	1.06	4 (6%)
22	CLA	B	838	-	65,73,73	1.32	7 (10%)	76,113,113	1.93	15 (19%)
22	CLA	F	304	6	65,73,73	1.33	7 (10%)	76,113,113	1.99	17 (22%)
39	CHL	2	303	-	51,59,74	0.95	2 (3%)	55,96,114	1.48	13 (23%)
22	CLA	1	304	-	65,73,73	1.34	6 (9%)	76,113,113	1.99	18 (23%)
22	CLA	9	310	-	45,53,73	1.59	8 (17%)	52,89,113	2.21	14 (26%)
22	CLA	1	313	44	65,73,73	1.33	7 (10%)	76,113,113	2.03	18 (23%)
22	CLA	A	835	-	65,73,73	1.33	7 (10%)	76,113,113	1.94	18 (23%)
22	CLA	7	310	25	65,73,73	1.34	7 (10%)	76,113,113	1.96	15 (19%)
22	CLA	8	305	15	60,68,73	1.39	6 (10%)	70,107,113	2.09	17 (24%)
22	CLA	Z	316	12	65,73,73	1.34	7 (10%)	76,113,113	1.95	16 (21%)
22	CLA	L	202	-	65,73,73	1.34	7 (10%)	76,113,113	1.96	17 (22%)

Mol	Type	Chain	Res	Link	Bond lengths			Bond angles		
					Counts	RMSZ	$\#\|Z\|>2$	Counts	RMSZ	$\#\|Z\|>2$
22	CLA	5	322	-	65,73,73	1.33	7 (10%)	76,113,113	1.97	17 (22%)
22	CLA	5	326	-	55,63,73	1.45	7 (12%)	64,101,113	2.11	18 (28%)
24	BCR	L	204	-	41,41,41	4.78	27 (65%)	56,56,56	2.34	18 (32%)
22	CLA	A	839	-	51,59,73	1.51	8 (15%)	59,96,113	2.22	17 (28%)
24	BCR	A	848	-	41,41,41	4.78	27 (65%)	56,56,56	2.42	22 (39%)
22	CLA	5	314	-	45,53,73	1.60	7 (15%)	52,89,113	2.13	13 (25%)
22	CLA	B	837	-	51,59,73	1.51	7 (13%)	59,96,113	2.24	18 (30%)
24	BCR	6	306	-	41,41,41	4.77	27 (65%)	56,56,56	2.79	23 (41%)
22	CLA	L	201	-	65,73,73	1.33	7 (10%)	76,113,113	2.00	17 (22%)
39	CHL	6	320	18	43,51,74	1.02	3 (6%)	45,86,114	1.53	10 (22%)
22	CLA	B	842	25	65,73,73	1.33	7 (10%)	76,113,113	1.98	16 (21%)
22	CLA	L	203	-	50,58,73	1.52	6 (12%)	58,95,113	2.24	17 (29%)
22	CLA	8	310	44	60,68,73	1.39	7 (11%)	70,107,113	2.09	17 (24%)
22	CLA	A	822	-	55,63,73	1.46	8 (14%)	64,101,113	2.20	17 (26%)
22	CLA	B	817	-	57,65,73	1.42	7 (12%)	66,103,113	2.09	16 (24%)
22	CLA	3	312	-	65,73,73	1.34	8 (12%)	76,113,113	1.95	17 (22%)
24	BCR	F	303	-	41,41,41	4.81	27 (65%)	56,56,56	2.21	18 (32%)
33	LMT	4	801	-	36,36,36	0.36	0	47,47,47	0.72	1 (2%)
22	CLA	5	308	17	61,69,73	1.37	8 (13%)	71,108,113	2.05	18 (25%)
22	CLA	9	306	19	62,70,73	1.37	7 (11%)	72,109,113	2.10	19 (26%)
39	CHL	6	315	44	56,64,74	0.87	2 (3%)	61,102,114	1.36	12 (19%)
22	CLA	B	836	-	45,53,73	1.58	7 (15%)	52,89,113	2.16	16 (30%)
22	CLA	A	823	-	65,73,73	1.33	8 (12%)	76,113,113	1.98	17 (22%)
22	CLA	6	311	-	55,63,73	1.45	7 (12%)	64,101,113	2.08	17 (26%)
25	LHG	7	318	22	36,36,48	0.43	0	39,42,54	1.14	3 (7%)
22	CLA	A	828	-	65,73,73	1.33	7 (10%)	76,113,113	1.99	15 (19%)
43	LPX	8	319	-	29,29,29	0.29	0	31,33,33	0.35	0
22	CLA	3	311	13	60,68,73	1.39	7 (11%)	70,107,113	2.05	19 (27%)
22	CLA	4	811	25	55,63,73	1.46	7 (12%)	64,101,113	2.14	17 (26%)
22	CLA	K	203	44	55,63,73	1.45	7 (12%)	64,101,113	2.15	17 (26%)
22	CLA	G	202	7	46,54,73	1.58	7 (15%)	53,90,113	2.16	15 (28%)
22	CLA	6	321	-	46,54,73	1.58	7 (15%)	53,90,113	2.15	13 (24%)
22	CLA	A	808	1	65,73,73	1.33	7 (10%)	76,113,113	2.00	17 (22%)
24	BCR	A	846	-	41,41,41	4.80	27 (65%)	56,56,56	2.47	22 (39%)
38	LUT	4	803	-	42,43,43	6.06	28 (66%)	51,60,60	2.15	13 (25%)

Mol	Type	Chain	Res	Link	Bond lengths			Bond angles		
					Counts	RMSZ	$\#\|Z\|>2$	Counts	RMSZ	$\#\|Z\|>2$
22	CLA	F	305	-	45,53,73	1.59	8 (17%)	52,89,113	2.12	13 (25%)
39	CHL	9	312	-	66,74,74	0.84	3 (4%)	73,114,114	1.41	10 (13%)
38	LUT	5	302	-	42,43,43	6.07	28 (66%)	51,60,60	1.96	14 (27%)
22	CLA	1	315	-	46,54,73	1.58	6 (13%)	53,90,113	2.16	14 (26%)
22	CLA	B	823	-	55,63,73	1.46	9 (16%)	64,101,113	2.09	18 (28%)
22	CLA	A	840	-	65,73,73	1.33	8 (12%)	76,113,113	1.99	17 (22%)
22	CLA	8	316	15	46,54,73	1.59	8 (17%)	53,90,113	2.11	13 (24%)
22	CLA	A	825	44	65,73,73	1.34	7 (10%)	76,113,113	2.02	18 (23%)
22	CLA	F	302	-	65,73,73	1.33	8 (12%)	76,113,113	1.97	18 (23%)
22	CLA	1	309	-	61,69,73	1.37	8 (13%)	71,108,113	2.07	16 (22%)
22	CLA	9	303	19	60,68,73	1.42	8 (13%)	70,107,113	2.15	22 (31%)
22	CLA	4	809	16	51,59,73	1.52	8 (15%)	59,96,113	2.15	17 (28%)
22	CLA	F	301	44	65,73,73	1.34	7 (10%)	76,113,113	2.00	19 (25%)
22	CLA	1	307	12	60,68,73	1.39	8 (13%)	70,107,113	2.09	16 (22%)
22	CLA	B	832	-	50,58,73	1.52	7 (14%)	58,95,113	2.21	17 (29%)
22	CLA	B	828	-	65,73,73	1.34	7 (10%)	76,113,113	1.97	17 (22%)
22	CLA	A	827	-	65,73,73	1.34	7 (10%)	76,113,113	1.99	19 (25%)
22	CLA	B	811	-	65,73,73	1.34	7 (10%)	76,113,113	2.01	18 (23%)
22	CLA	7	316	44	42,50,73	1.64	7 (16%)	48,85,113	2.24	16 (33%)
24	BCR	B	845	-	41,41,41	4.78	27 (65%)	56,56,56	2.39	20 (35%)
22	CLA	7	304	14	60,68,73	1.39	7 (11%)	70,107,113	2.13	19 (27%)
31	DGD	B	852	-	62,62,67	1.12	6 (9%)	76,76,81	0.96	3 (3%)
22	CLA	1	316	12	65,73,73	1.38	9 (13%)	76,113,113	1.92	17 (22%)
39	CHL	1	312	-	48,56,74	0.95	2 (4%)	51,92,114	1.45	10 (19%)
34	LMG	J	101	-	29,29,55	0.53	0	37,37,63	1.12	3 (8%)
22	CLA	8	307	-	65,73,73	1.33	8 (12%)	76,113,113	2.06	16 (21%)
22	CLA	A	802	44	65,73,73	1.34	7 (10%)	76,113,113	2.03	17 (22%)
22	CLA	A	813	-	65,73,73	1.34	6 (9%)	76,113,113	2.01	18 (23%)
22	CLA	9	309	25	55,63,73	1.45	7 (12%)	64,101,113	2.17	13 (20%)
22	CLA	6	319	44	61,69,73	1.38	7 (11%)	71,108,113	2.06	17 (23%)
22	CLA	B	827	44	65,73,73	1.33	7 (10%)	76,113,113	2.02	17 (22%)
24	BCR	5	305	-	41,41,41	4.80	27 (65%)	56,56,56	2.67	22 (39%)
22	CLA	A	816	44	56,64,73	1.44	8 (14%)	65,102,113	2.18	19 (29%)
22	CLA	B	840	44	65,73,73	1.33	7 (10%)	76,113,113	1.97	17 (22%)
39	CHL	4	813	44	51,59,74	0.95	2 (3%)	55,96,114	1.44	12 (21%)

Mol	Type	Chain	Res	Link	Bond lengths			Bond angles		
					Counts	RMSZ	$\#\|Z\|>2$	Counts	RMSZ	$\#\|Z\|>2$
22	CLA	B	835	-	50,58,73	1.51	7 (14%)	58,95,113	2.31	18 (31%)
39	CHL	4	816	44	47,55,74	0.99	3 (6%)	50,91,114	1.44	8 (16%)
22	CLA	Z	306	12	60,68,73	1.39	8 (13%)	70,107,113	2.04	17 (24%)
22	CLA	1	306	-	65,73,73	1.34	7 (10%)	76,113,113	2.01	17 (22%)
22	CLA	A	836	-	55,63,73	1.44	8 (14%)	64,101,113	2.11	19 (29%)
25	LHG	6	323	22	48,48,48	0.38	0	51,54,54	1.09	3 (5%)
25	LHG	3	321	22	19,19,48	1.07	1 (5%)	20,24,54	1.53	1 (5%)
32	RRX	F	306	-	42,42,42	5.06	25 (59%)	57,58,58	2.59	23 (40%)
22	CLA	7	324	-	60,68,73	1.39	7 (11%)	70,107,113	2.07	19 (27%)
25	LHG	Z	317	22	42,42,48	0.41	0	45,48,54	1.08	3 (6%)
22	CLA	B	813	-	56,64,73	1.44	7 (12%)	65,102,113	2.16	17 (26%)
22	CLA	5	313	25	61,69,73	1.39	8 (13%)	71,108,113	2.06	17 (23%)
22	CLA	6	307	18	60,68,73	1.39	8 (13%)	70,107,113	2.06	17 (24%)
22	CLA	3	316	13	52,60,73	1.49	7 (13%)	60,97,113	2.16	17 (28%)
22	CLA	B	822	-	56,64,73	1.44	7 (12%)	65,102,113	2.06	16 (24%)
38	LUT	6	304	-	42,43,43	6.08	28 (66%)	51,60,60	2.05	13 (25%)
22	CLA	6	309	44	65,73,73	1.34	8 (12%)	76,113,113	2.07	19 (25%)
22	CLA	4	810	44	50,58,73	1.51	6 (12%)	58,95,113	2.24	18 (31%)
22	CLA	Z	309	25	60,68,73	1.39	7 (11%)	70,107,113	2.03	16 (22%)
22	CLA	3	313	44	65,73,73	1.31	6 (9%)	76,113,113	2.00	17 (22%)
38	LUT	6	303	-	42,43,43	6.11	27 (64%)	51,60,60	1.92	16 (31%)
22	CLA	7	308	14	61,69,73	1.39	8 (13%)	71,108,113	2.04	18 (25%)
22	CLA	8	314	15	46,54,73	1.57	7 (15%)	53,90,113	2.09	13 (24%)
22	CLA	3	309	13	46,54,73	1.58	8 (17%)	53,90,113	2.13	15 (28%)
22	CLA	7	305	-	50,58,73	1.53	7 (14%)	58,95,113	2.23	17 (29%)
25	LHG	4	821	-	31,31,48	0.48	0	34,37,54	1.19	3 (8%)
36	C7Z	J	105	-	43,43,43	5.34	27 (62%)	58,60,60	2.21	21 (36%)
22	CLA	9	313	19	50,58,73	1.52	7 (14%)	58,95,113	2.23	17 (29%)
22	CLA	5	323	44	46,54,73	1.57	7 (15%)	53,90,113	2.20	14 (26%)
39	CHL	4	819	16	43,51,74	1.01	2 (4%)	45,86,114	1.54	10 (22%)
22	CLA	B	833	-	65,73,73	1.32	6 (9%)	76,113,113	1.97	17 (22%)
38	LUT	1	303	-	42,43,43	6.07	28 (66%)	51,60,60	2.06	12 (23%)
22	CLA	6	313	25	55,63,73	1.47	7 (12%)	64,101,113	2.08	16 (25%)
22	CLA	B	819	-	59,67,73	1.40	7 (11%)	68,105,113	2.09	19 (27%)
22	CLA	8	308	15	62,70,73	1.36	7 (11%)	72,109,113	2.03	17 (23%)

Mol	Type	Chain	Res	Link	Bond lengths			Bond angles		
					Counts	RMSZ	$\#\|Z\|>2$	Counts	RMSZ	$\#\|Z\|>2$
22	CLA	Z	313	44	65,73,73	1.34	7 (10%)	76,113,113	1.98	18 (23%)
22	CLA	Z	308	-	61,69,73	1.37	7 (11%)	71,108,113	2.04	17 (23%)
22	CLA	B	825	-	65,73,73	1.33	6 (9%)	76,113,113	1.98	18 (23%)
22	CLA	B	818	-	65,73,73	1.33	8 (12%)	76,113,113	1.96	17 (22%)
22	CLA	B	820	-	60,68,73	1.40	7 (11%)	70,107,113	2.00	17 (24%)
22	CLA	B	839	-	52,60,73	1.49	8 (15%)	60,97,113	2.22	18 (30%)
38	LUT	8	303	-	42,43,43	6.07	28 (66%)	51,60,60	1.97	14 (27%)
24	BCR	5	304	-	41,41,41	4.78	25 (60%)	56,56,56	2.54	22 (39%)
29	SF4	C	101	3	0,12,12	-	-	-		
22	CLA	A	833	-	65,73,73	1.34	7 (10%)	76,113,113	1.99	18 (23%)
24	BCR	3	306	-	41,41,41	4.80	27 (65%)	56,56,56	2.28	17 (30%)
22	CLA	9	304	-	45,53,73	1.60	7 (15%)	52,89,113	2.14	13 (25%)
24	BCR	G	203	-	41,41,41	4.79	27 (65%)	56,56,56	2.41	22 (39%)
22	CLA	Z	303	12	65,73,73	1.33	6 (9%)	76,113,113	2.03	19 (25%)
24	BCR	3	304	-	41,41,41	4.82	27 (65%)	56,56,56	3.07	28 (50%)
22	CLA	B	807	-	45,53,73	1.59	7 (15%)	52,89,113	2.13	14 (26%)
33	LMT	G	204	-	36,36,36	0.39	0	47,47,47	0.74	2 (4%)
41	3PH	6	324	-	28,28,47	1.09	4 (14%)	32,33,52	1.20	2 (6%)
22	CLA	9	307	19	55,63,73	1.46	9 (16%)	64,101,113	2.07	16 (25%)
24	BCR	A	845	-	41,41,41	4.77	27 (65%)	56,56,56	2.50	24 (42%)
22	CLA	6	322	18	65,73,73	1.33	7 (10%)	76,113,113	2.02	18 (23%)
41	3PH	8	320	-	29,29,47	1.06	4 (13%)	33,34,52	1.16	2 (6%)
22	CLA	K	204	10	49,57,73	1.55	8 (16%)	55,93,113	2.25	17 (30%)
22	CLA	A	838	-	65,73,73	1.33	7 (10%)	76,113,113	2.02	17 (22%)
22	CLA	Z	314	12	51,59,73	1.50	7 (13%)	59,96,113	2.17	16 (27%)
22	CLA	7	312	-	60,68,73	1.39	7 (11%)	70,107,113	2.03	17 (24%)
22	CLA	B	826	44	65,73,73	1.34	7 (10%)	76,113,113	2.00	18 (23%)
22	CLA	3	314	25	60,68,73	1.39	7 (11%)	70,107,113	2.03	16 (22%)
22	CLA	6	302	16	60,68,73	1.39	7 (11%)	70,107,113	2.00	16 (22%)
33	LMT	F	307	-	36,36,36	0.38	0	47,47,47	0.64	0
22	CLA	3	322	14	65,73,73	1.34	8 (12%)	76,113,113	1.94	16 (21%)
22	CLA	K	202	-	46,54,73	1.58	6 (13%)	53,90,113	2.20	16 (30%)
24	BCR	8	304	-	41,41,41	4.80	27 (65%)	56,56,56	2.43	21 (37%)
23	PQN	A	843	-	34,34,34	0.37	0	42,45,45	1.20	6 (14%)
39	CHL	5	321	17	43,51,74	1.02	2 (4%)	45,86,114	1.52	10 (22%)

Mol	Type	Chain	Res	Link	Bond lengths			Bond angles		
					Counts	RMSZ	$\#\|Z\|>2$	Counts	RMSZ	$\#\|Z\|>2$
22	CLA	B	812	2	59,67,73	1.39	7 (11%)	68,105,113	2.01	16 (23%)
24	BCR	B	849	-	41,41,41	4.81	27 (65%)	56,56,56	2.35	21 (37%)
31	DGD	1	319	-	52,52,67	0.89	2 (3%)	66,66,81	0.97	4 (6%)
22	CLA	7	307	14	65,73,73	1.34	8 (12%)	76,113,113	2.02	16 (21%)
22	CLA	9	311	19	46,54,73	1.59	7 (15%)	53,90,113	2.12	14 (26%)
22	CLA	2	304	44	46,54,73	1.58	7 (15%)	53,90,113	2.12	13 (24%)
22	CLA	A	830	-	65,73,73	1.34	7 (10%)	76,113,113	1.98	16 (21%)
22	CLA	7	323	15	65,73,73	1.33	7 (10%)	76,113,113	2.01	18 (23%)
22	CLA	A	834	-	65,73,73	1.34	7 (10%)	76,113,113	1.96	17 (22%)
26	NKP	8	318	-	28,28,28	0.33	0	31,32,32	0.34	0
38	LUT	Z	302	-	42,43,43	6.09	27 (64%)	51,60,60	2.05	13 (25%)
39	CHL	8	301	12	58,66,74	0.92	3 (5%)	63,104,114	1.42	12 (19%)
22	CLA	2	301	-	55,63,73	1.45	8 (14%)	64,101,113	2.14	15 (23%)
22	CLA	8	306	15	51,59,73	1.52	7 (13%)	59,96,113	2.13	17 (28%)
23	PQN	B	843	-	34,34,34	0.37	0	42,45,45	1.13	3 (7%)
24	BCR	B	846	-	41,41,41	4.79	27 (65%)	56,56,56	2.40	22 (39%)
22	CLA	7	315	14	50,58,73	1.53	8 (16%)	58,95,113	2.25	18 (31%)
22	CLA	B	834	-	58,66,73	1.41	7 (12%)	67,104,113	2.10	17 (25%)
38	LUT	7	302	-	42,43,43	6.08	28 (66%)	51,60,60	2.06	15 (29%)
22	CLA	8	309	-	65,73,73	1.34	8 (12%)	76,113,113	1.97	17 (22%)
22	CLA	K	205	10	55,63,73	1.54	10 (18%)	64,101,113	2.21	15 (23%)
22	CLA	A	832	-	65,73,73	1.32	6 (9%)	76,113,113	1.96	19 (25%)
24	BCR	3	305	-	41,41,41	4.78	27 (65%)	56,56,56	2.19	19 (33%)
25	LHG	9	315	22	32,32,48	0.48	0	35,38,54	1.05	3 (8%)
22	CLA	6	310	18	65,73,73	1.34	9 (13%)	76,113,113	1.94	18 (23%)
22	CLA	1	308	-	55,63,73	1.46	8 (14%)	64,101,113	2.20	16 (25%)
22	CLA	7	309	-	56,64,73	1.43	7 (12%)	65,102,113	2.17	19 (29%)
22	CLA	A	811	22	65,73,73	1.33	8 (12%)	76,113,113	1.93	17 (22%)
24	BCR	6	305	-	41,41,41	4.79	26 (63%)	56,56,56	2.49	21 (37%)
22	CLA	B	814	-	60,68,73	1.39	7 (11%)	70,107,113	2.04	15 (21%)
22	CLA	9	305	19	60,68,73	1.44	9 (15%)	70,107,113	2.44	22 (31%)
25	LHG	A	849	22	34,34,48	0.44	0	37,40,54	1.14	3 (8%)
22	CLA	3	320	13	46,54,73	1.60	8 (17%)	53,90,113	2.21	13 (24%)
38	LUT	5	303	-	42,43,43	6.11	28 (66%)	51,60,60	2.00	14 (27%)
38	LUT	8	302	-	42,43,43	6.02	28 (66%)	51,60,60	1.96	13 (25%)

Mol	Type	Chain	Res	Link	Bond lengths			Bond angles		
					Counts	RMSZ	$\#\|Z\|>2$	Counts	RMSZ	$\#\|Z\|>2$
22	CLA	5	307	17	60,68,73	1.38	8 (13%)	70,107,113	2.07	17 (24%)
39	CHL	8	312	-	56,64,74	0.89	2 (3%)	61,102,114	1.38	12 (19%)
39	CHL	Z	312	-	48,56,74	0.96	2 (4%)	51,92,114	1.43	11 (21%)
25	LHG	B	851	-	19,19,48	0.87	1 (5%)	20,24,54	1.36	1 (5%)
29	SF4	C	102	3	0,12,12	-	-	-		
22	CLA	A	826	44	50,58,73	1.52	7 (14%)	58,95,113	2.24	16 (27%)
22	CLA	5	320	-	50,58,73	1.59	9 (18%)	58,95,113	2.18	15 (25%)
22	CLA	A	821	44	65,73,73	1.33	7 (10%)	76,113,113	1.96	17 (22%)
22	CLA	A	837	-	51,59,73	1.52	7 (13%)	59,96,113	2.18	16 (27%)
38	LUT	9	301	-	42,43,43	6.08	28 (66%)	51,60,60	1.96	12 (23%)
38	LUT	9	302	-	42,43,43	6.07	28 (66%)	51,60,60	1.87	12 (23%)
22	CLA	B	830	-	65,73,73	1.34	7 (10%)	76,113,113	1.94	16 (21%)
21	CL0	A	801	-	65,73,73	2.37	19 (29%)	76,113,113	2.56	20 (26%)

In the following table, the Chirals column lists the number of chiral outliers, the number of chiral centers analysed, the number of these observed in the model and the number defined in the Chemical Component Dictionary. Similar counts are reported in the Torsion and Rings columns. '-' means no outliers of that kind were identified.

Mol	Type	Chain	Res	Link	Chirals	Torsions	Rings
22	CLA	5	315	17	1/1/15/20	14/37/115/115	-
22	CLA	3	319	-	1/1/13/20	9/25/103/115	-
22	CLA	A	812	-	1/1/15/20	16/37/115/115	-
22	CLA	9	308	-	1/1/12/20	8/19/97/115	-
22	CLA	G	201	-	1/1/12/20	10/19/97/115	-
42	SPH	7	321	-	-	11/21/21/21	-
22	CLA	7	314	44	1/1/12/20	7/19/97/115	-
22	CLA	6	312	-	1/1/15/20	15/37/115/115	-
22	CLA	Z	304	12	1/1/11/20	6/13/91/115	-
26	NKP	A	851	-	-	9/22/22/28	-
39	CHL	8	315	44	3/3/20/26	12/39/137/137	-
39	CHL	5	316	44	3/3/20/26	12/39/137/137	-
22	CLA	A	831	-	1/1/12/20	10/19/97/115	-
22	CLA	5	311	-	1/1/13/20	11/25/103/115	-
22	CLA	A	854	-	1/1/15/20	13/37/115/115	-
22	CLA	B	816	-	1/1/14/20	13/31/109/115	-

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Mol	Type	Chain	Res	Link	Chirals	Torsions	Rings
22	CLA	J	103	9	1/1/10/20	5/10/88/115	-
22	CLA	3	308	13	1/1/15/20	18/37/115/115	-
24	BCR	B	853	-	-	17/29/63/63	0/2/2/2
39	CHL	6	318	-	3/3/18/26	8/27/125/137	-
24	BCR	A	844	-	-	13/29/63/63	0/2/2/2
25	LHG	8	317	22	-	27/42/42/53	-
22	CLA	A	817	-	1/1/14/20	13/31/109/115	-
22	CLA	B	806	44	1/1/15/20	12/37/115/115	-
38	LUT	3	303	-	-	7/29/67/67	0/2/2/2
39	CHL	9	314	-	3/3/15/26	0/10/108/137	-
22	CLA	A	855	44	1/1/15/20	18/37/115/115	-
22	CLA	B	808	-	1/1/15/20	15/37/115/115	-
22	CLA	A	814	-	1/1/14/20	19/31/109/115	-
39	CHL	4	814	44	3/3/17/26	10/21/119/137	-
22	CLA	A	818	-	1/1/14/20	15/31/109/115	-
41	3PH	5	325	-	-	5/24/24/49	-
25	LHG	4	820	22	-	30/53/53/53	-
22	CLA	5	318	17	1/1/15/20	14/37/115/115	-
22	CLA	Z	315	-	1/1/11/20	8/15/93/115	-
41	3PH	7	319	-	-	17/40/40/49	-
24	BCR	3	307	-	-	13/29/63/63	0/2/2/2
22	CLA	A	815	-	1/1/15/20	16/37/115/115	-
22	CLA	4	805	16	1/1/14/20	13/31/109/115	-
28	OCA	A	853	-	-	3/7/7/7	-
22	CLA	A	804	22	1/1/11/20	6/13/91/115	-
22	CLA	3	318	13	1/1/14/20	11/31/109/115	-
22	CLA	4	807	16	1/1/15/20	12/37/115/115	-
35	T7X	J	102	-	-	17/44/68/80	0/1/1/1
22	CLA	A	807	-	1/1/14/20	17/31/109/115	-
34	LMG	F	308	-	-	16/30/50/70	0/1/1/1
22	CLA	7	306	-	1/1/15/20	16/37/115/115	-
24	BCR	A	856	-	-	16/29/63/63	0/2/2/2
22	CLA	2	302	-	1/1/11/20	8/13/91/115	-
22	CLA	7	311	-	1/1/10/20	5/11/89/115	-

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Mol	Type	Chain	Res	Link	Chirals	Torsions	Rings
22	CLA	1	311	-	1/1/11/20	6/15/93/115	-
22	CLA	1	314	12	1/1/15/20	13/37/115/115	-
37	DAO	K	201	-	-	1/11/11/11	-
24	BCR	7	303	-	-	16/29/63/63	0/2/2/2
22	CLA	6	308	18	1/1/12/20	7/22/100/115	-
22	CLA	B	821	44	1/1/15/20	18/37/115/115	-
22	CLA	A	805	-	1/1/15/20	19/37/115/115	-
22	CLA	B	810	-	1/1/13/20	11/27/105/115	-
22	CLA	A	819	-	1/1/15/20	12/37/115/115	-
22	CLA	8	311	25	1/1/12/20	10/19/97/115	-
24	BCR	I	4001	-	-	14/29/63/63	0/2/2/2
33	LMT	4	822	-	-	6/21/61/61	0/2/2/2
38	LUT	4	802	-	-	10/29/67/67	0/2/2/2
22	CLA	7	317	-	1/1/13/20	15/29/107/115	-
39	CHL	7	313	44	3/3/17/26	7/25/123/137	-
22	CLA	A	829	-	1/1/15/20	11/37/115/115	-
22	CLA	5	319	44	1/1/13/20	11/25/103/115	-
22	CLA	A	803	-	1/1/15/20	12/37/115/115	-
22	CLA	B	801	-	1/1/15/20	15/37/115/115	-
22	CLA	B	824	-	1/1/13/20	19/30/108/115	-
22	CLA	6	301	44	1/1/14/20	18/31/109/115	-
22	CLA	5	310	17	1/1/15/20	13/37/115/115	-
39	CHL	6	316	-	3/3/17/26	9/21/119/137	-
22	CLA	3	310	13	1/1/15/20	14/37/115/115	-
40	SQD	1	318	-	-	12/43/63/69	0/1/1/1
39	CHL	5	317	44	3/3/17/26	6/21/119/137	-
22	CLA	A	810	-	1/1/15/20	17/37/115/115	-
38	LUT	1	302	-	-	9/29/67/67	0/2/2/2
24	BCR	B	847	-	-	11/29/63/63	0/2/2/2
22	CLA	4	806	16	1/1/12/20	10/22/100/115	-
39	CHL	3	317	-	3/3/20/26	12/39/137/137	-
39	CHL	Z	311	12	3/3/17/26	10/21/119/137	-
38	LUT	3	302	-	-	4/29/67/67	0/2/2/2
22	CLA	6	317	18	1/1/12/20	10/19/97/115	-

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Mol	Type	Chain	Res	Link	Chirals	Torsions	Rings
24	BCR	K	206	-	-	13/29/63/63	0/2/2/2
22	CLA	4	818	44	1/1/12/20	10/21/99/115	-
25	LHG	5	324	22	-	24/41/41/53	-
22	CLA	1	310	25	1/1/14/20	17/31/109/115	-
25	LHG	B	850	22	-	16/26/26/53	-
22	CLA	B	841	-	1/1/15/20	11/37/115/115	-
22	CLA	5	309	-	1/1/13/20	17/27/105/115	-
22	CLA	5	312	44	1/1/12/20	6/19/97/115	-
25	LHG	A	850	-	-	29/53/53/53	-
24	BCR	4	804	-	-	14/29/63/63	0/2/2/2
22	CLA	A	841	-	1/1/15/20	10/37/115/115	-
25	LHG	1	317	22	-	22/47/47/53	-
22	CLA	B	809	2	1/1/15/20	19/37/115/115	-
22	CLA	8	313	44	1/1/12/20	7/19/97/115	-
22	CLA	4	812	-	1/1/13/20	8/25/103/115	-
27	DGA	A	852	-	-	23/41/41/45	-
22	CLA	A	824	-	1/1/15/20	17/37/115/115	-
22	CLA	B	805	-	1/1/15/20	16/37/115/115	-
33	LMT	1	301	-	-	4/21/61/61	0/2/2/2
36	C7Z	5	306	-	-	13/29/67/67	0/2/2/2
22	CLA	4	815	16	1/1/12/20	8/19/97/115	-
24	BCR	B	848	-	-	16/29/63/63	0/2/2/2
22	CLA	4	808	16	1/1/14/20	16/31/109/115	-
24	BCR	J	104	-	-	14/29/63/63	0/2/2/2
22	CLA	A	809	1	1/1/13/20	7/25/103/115	-
22	CLA	Z	310	-	1/1/12/20	10/19/97/115	-
22	CLA	6	314	18	1/1/15/20	15/37/115/115	-
22	CLA	B	829	-	1/1/13/20	15/28/106/115	-
22	CLA	B	815	-	1/1/15/20	13/37/115/115	-
22	CLA	A	820	-	1/1/14/20	21/31/109/115	-
29	SF4	B	802	-	-	-	0/6/5/5
22	CLA	5	301	44	1/1/13/20	13/27/105/115	-
22	CLA	7	322	-	1/1/13/20	9/25/103/115	-
38	LUT	Z	301	-	-	7/29/67/67	0/2/2/2

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Mol	Type	Chain	Res	Link	Chirals	Torsions	Rings
42	SPH	7	320	-	-	11/21/21/21	-
22	CLA	A	806	1	1/1/15/20	16/37/115/115	-
22	CLA	B	831	-	1/1/13/20	17/29/107/115	-
22	CLA	Z	305	-	1/1/13/20	12/27/105/115	-
25	LHG	B	803	-	-	22/37/37/53	-
22	CLA	A	842	25	1/1/12/20	12/22/100/115	-
22	CLA	3	315	-	1/1/11/20	6/13/91/115	-
24	BCR	B	844	-	-	12/29/63/63	0/2/2/2
22	CLA	1	305	12	1/1/11/20	6/13/91/115	-
24	BCR	A	847	-	-	16/29/63/63	0/2/2/2
22	CLA	Z	307	-	1/1/13/20	10/25/103/115	-
38	LUT	7	301	-	-	9/29/67/67	0/2/2/2
22	CLA	4	817	16	1/1/10/20	2/8/86/115	-
31	DGD	3	301	-	-	12/40/80/95	0/2/2/2
22	CLA	B	838	-	1/1/15/20	15/37/115/115	-
22	CLA	F	304	6	1/1/15/20	23/37/115/115	-
39	CHL	2	303	-	3/3/17/26	7/21/119/137	-
22	CLA	1	304	-	1/1/15/20	13/37/115/115	-
22	CLA	9	310	-	1/1/11/20	4/13/91/115	-
22	CLA	1	313	44	1/1/15/20	12/37/115/115	-
22	CLA	A	835	-	1/1/15/20	13/37/115/115	-
22	CLA	7	310	25	1/1/15/20	15/37/115/115	-
22	CLA	8	305	15	1/1/14/20	14/31/109/115	-
22	CLA	Z	316	12	1/1/15/20	21/37/115/115	-
22	CLA	L	202	-	1/1/15/20	21/37/115/115	-
22	CLA	5	322	-	1/1/15/20	11/37/115/115	-
22	CLA	5	326	-	1/1/13/20	8/25/103/115	-
24	BCR	L	204	-	-	13/29/63/63	0/2/2/2
22	CLA	A	839	-	1/1/12/20	10/21/99/115	-
24	BCR	A	848	-	-	15/29/63/63	0/2/2/2
22	CLA	5	314	-	1/1/11/20	6/13/91/115	-
22	CLA	B	837	-	1/1/12/20	9/21/99/115	-
24	BCR	6	306	-	-	16/29/63/63	0/2/2/2
22	CLA	L	201	-	1/1/15/20	17/37/115/115	-

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Mol	Type	Chain	Res	Link	Chirals	Torsions	Rings
39	CHL	6	320	18	3/3/15/26	0/12/110/137	-
22	CLA	B	842	25	1/1/15/20	17/37/115/115	-
22	CLA	L	203	-	1/1/12/20	7/19/97/115	-
22	CLA	8	310	44	1/1/14/20	15/31/109/115	-
22	CLA	A	822	-	1/1/13/20	12/25/103/115	-
22	CLA	B	817	-	1/1/13/20	18/28/106/115	-
22	CLA	3	312	-	1/1/15/20	12/37/115/115	-
24	BCR	F	303	-	-	15/29/63/63	0/2/2/2
39	CHL	6	315	44	3/3/18/26	11/27/125/137	-
22	CLA	5	308	17	1/1/14/20	17/33/111/115	-
22	CLA	9	306	19	1/1/14/20	14/34/112/115	-
33	LMT	4	801	-	-	1/21/61/61	0/2/2/2
22	CLA	B	836	-	1/1/11/20	6/13/91/115	-
22	CLA	A	823	-	1/1/15/20	15/37/115/115	-
22	CLA	6	311	-	1/1/13/20	12/25/103/115	-
25	LHG	7	318	22	-	25/41/41/53	-
22	CLA	A	828	-	1/1/15/20	21/37/115/115	-
43	LPX	8	319	-	-	6/31/31/31	-
22	CLA	3	311	13	1/1/14/20	13/31/109/115	-
22	CLA	4	811	25	1/1/13/20	9/25/103/115	-
22	CLA	K	203	44	1/1/13/20	9/25/103/115	-
22	CLA	G	202	7	1/1/11/20	8/15/93/115	-
22	CLA	6	321	-	1/1/11/20	6/15/93/115	-
22	CLA	A	808	1	1/1/15/20	19/37/115/115	-
24	BCR	A	846	-	-	13/29/63/63	0/2/2/2
38	LUT	4	803	-	-	8/29/67/67	0/2/2/2
22	CLA	F	305	-	1/1/11/20	6/13/91/115	-
39	CHL	9	312	-	3/3/20/26	16/39/137/137	-
38	LUT	5	302	-	-	9/29/67/67	0/2/2/2
22	CLA	1	315	-	1/1/11/20	7/15/93/115	-
22	CLA	B	823	-	1/1/13/20	13/25/103/115	-
22	CLA	A	840	-	1/1/15/20	18/37/115/115	-
22	CLA	8	316	15	1/1/11/20	11/15/93/115	-
22	CLA	A	825	44	1/1/15/20	16/37/115/115	-

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Mol	Type	Chain	Res	Link	Chirals	Torsions	Rings
22	CLA	F	302	-	1/1/15/20	17/37/115/115	-
22	CLA	1	309	-	1/1/14/20	13/33/111/115	-
22	CLA	9	303	19	1/1/14/20	11/31/109/115	-
22	CLA	4	809	16	1/1/12/20	5/21/99/115	-
22	CLA	F	301	44	1/1/15/20	13/37/115/115	-
22	CLA	1	307	12	1/1/14/20	18/31/109/115	-
22	CLA	B	832	-	1/1/12/20	6/19/97/115	-
22	CLA	B	828	-	1/1/15/20	18/37/115/115	-
22	CLA	A	827	-	1/1/15/20	17/37/115/115	-
22	CLA	B	811	-	1/1/15/20	15/37/115/115	-
22	CLA	7	316	44	1/1/10/20	6/10/88/115	-
24	BCR	B	845	-	-	16/29/63/63	0/2/2/2
22	CLA	7	304	14	1/1/14/20	13/31/109/115	-
31	DGD	B	852	-	-	11/50/90/95	0/2/2/2
22	CLA	1	316	12	1/1/15/20	20/37/115/115	-
39	CHL	1	312	-	3/3/16/26	3/18/116/137	-
34	LMG	J	101	-	-	7/24/44/70	0/1/1/1
22	CLA	8	307	-	1/1/15/20	12/37/115/115	-
22	CLA	A	802	44	1/1/15/20	17/37/115/115	-
22	CLA	A	813	-	1/1/15/20	14/37/115/115	-
22	CLA	9	309	25	1/1/13/20	7/25/103/115	-
22	CLA	6	319	44	1/1/14/20	17/33/111/115	-
22	CLA	B	827	44	1/1/15/20	14/37/115/115	-
24	BCR	5	305	-	-	13/29/63/63	0/2/2/2
22	CLA	A	816	44	1/1/13/20	14/27/105/115	-
22	CLA	B	840	44	1/1/15/20	9/37/115/115	-
39	CHL	4	813	44	3/3/17/26	6/21/119/137	-
22	CLA	B	835	-	1/1/12/20	8/19/97/115	-
39	CHL	4	816	44	3/3/16/26	6/17/115/137	-
22	CLA	Z	306	12	1/1/14/20	20/31/109/115	-
22	CLA	1	306	-	1/1/15/20	18/37/115/115	-
22	CLA	A	836	-	1/1/13/20	12/25/103/115	-
25	LHG	6	323	22	-	38/53/53/53	-
25	LHG	3	321	22	-	11/22/22/53	-

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Mol	Type	Chain	Res	Link	Chirals	Torsions	Rings
32	RRX	F	306	-	-	15/29/65/65	0/2/2/2
22	CLA	7	324	-	1/1/14/20	23/31/109/115	-
25	LHG	Z	317	22	-	27/47/47/53	-
22	CLA	B	813	-	1/1/13/20	14/27/105/115	-
22	CLA	5	313	25	1/1/14/20	12/33/111/115	-
22	CLA	6	307	18	1/1/14/20	12/31/109/115	-
22	CLA	3	316	13	1/1/12/20	8/22/100/115	-
22	CLA	B	822	-	1/1/13/20	17/27/105/115	-
38	LUT	6	304	-	-	6/29/67/67	0/2/2/2
22	CLA	6	309	44	1/1/15/20	17/37/115/115	-
22	CLA	4	810	44	1/1/12/20	8/19/97/115	-
22	CLA	Z	309	25	1/1/14/20	16/31/109/115	-
22	CLA	3	313	44	1/1/15/20	17/37/115/115	-
38	LUT	6	303	-	-	9/29/67/67	0/2/2/2
22	CLA	7	308	14	1/1/14/20	16/33/111/115	-
22	CLA	8	314	15	1/1/11/20	10/15/93/115	-
22	CLA	3	309	13	1/1/11/20	6/15/93/115	-
22	CLA	7	305	-	1/1/12/20	4/19/97/115	-
25	LHG	4	821	-	-	16/36/36/53	-
36	C7Z	J	105	-	-	13/29/67/67	0/2/2/2
22	CLA	9	313	19	1/1/12/20	6/19/97/115	-
22	CLA	5	323	44	1/1/11/20	7/15/93/115	-
39	CHL	4	819	16	3/3/15/26	0/12/110/137	-
22	CLA	B	833	-	1/1/15/20	19/37/115/115	-
38	LUT	1	303	-	-	8/29/67/67	0/2/2/2
22	CLA	6	313	25	1/1/13/20	8/25/103/115	-
22	CLA	B	819	-	1/1/13/20	15/30/108/115	-
22	CLA	8	308	15	1/1/14/20	15/34/112/115	-
22	CLA	Z	313	44	1/1/15/20	13/37/115/115	-
22	CLA	Z	308	-	1/1/14/20	14/33/111/115	-
22	CLA	B	825	-	1/1/15/20	19/37/115/115	-
22	CLA	B	818	-	1/1/15/20	15/37/115/115	-
22	CLA	B	820	-	1/1/14/20	18/31/109/115	-
22	CLA	B	839	-	1/1/12/20	11/22/100/115	-

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Mol	Type	Chain	Res	Link	Chirals	Torsions	Rings
38	LUT	8	303	-	-	3/29/67/67	0/2/2/2
24	BCR	5	304	-	-	14/29/63/63	0/2/2/2
29	SF4	C	101	3	-	-	0/6/5/5
22	CLA	A	833	-	1/1/15/20	14/37/115/115	-
24	BCR	3	306	-	-	13/29/63/63	0/2/2/2
22	CLA	9	304	-	1/1/11/20	4/13/91/115	-
24	BCR	G	203	-	-	15/29/63/63	0/2/2/2
22	CLA	Z	303	12	1/1/15/20	16/37/115/115	-
24	BCR	3	304	-	-	10/29/63/63	0/2/2/2
22	CLA	B	807	-	1/1/11/20	8/13/91/115	-
33	LMT	G	204	-	-	6/21/61/61	0/2/2/2
41	3PH	6	324	-	-	11/30/30/49	-
22	CLA	9	307	19	1/1/13/20	10/25/103/115	-
24	BCR	A	845	-	-	14/29/63/63	0/2/2/2
22	CLA	6	322	18	1/1/15/20	15/37/115/115	-
41	3PH	8	320	-	-	13/31/31/49	-
22	CLA	K	204	10	1/1/11/20	10/18/96/115	-
22	CLA	A	838	-	1/1/15/20	15/37/115/115	-
22	CLA	Z	314	12	1/1/12/20	9/21/99/115	-
22	CLA	7	312	-	1/1/14/20	18/31/109/115	-
22	CLA	B	826	44	1/1/15/20	17/37/115/115	-
22	CLA	3	314	25	1/1/14/20	19/31/109/115	-
22	CLA	6	302	16	1/1/14/20	17/31/109/115	-
33	LMT	F	307	-	-	8/21/61/61	0/2/2/2
22	CLA	3	322	14	1/1/15/20	14/37/115/115	-
22	CLA	K	202	-	1/1/11/20	10/15/93/115	-
24	BCR	8	304	-	-	12/29/63/63	0/2/2/2
23	PQN	A	843	-	-	8/23/43/43	0/2/2/2
39	CHL	5	321	17	3/3/15/26	3/12/110/137	-
22	CLA	B	812	2	2/2/14/20	15/27/105/115	-
24	BCR	B	849	-	-	13/29/63/63	0/2/2/2
31	DGD	1	319	-	-	14/40/80/95	0/2/2/2
22	CLA	7	307	14	1/1/15/20	14/37/115/115	-
22	CLA	9	311	19	1/1/11/20	5/15/93/115	-

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Mol	Type	Chain	Res	Link	Chirals	Torsions	Rings
22	CLA	2	304	44	1/1/11/20	9/15/93/115	-
22	CLA	A	830	-	1/1/15/20	14/37/115/115	-
22	CLA	7	323	15	1/1/15/20	19/37/115/115	-
22	CLA	A	834	-	1/1/15/20	15/37/115/115	-
26	NKP	8	318	-	-	7/28/28/28	-
38	LUT	Z	302	-	-	7/29/67/67	0/2/2/2
39	CHL	8	301	12	3/3/18/26	5/30/128/137	-
22	CLA	2	301	-	1/1/13/20	11/25/103/115	-
22	CLA	8	306	15	1/1/12/20	9/21/99/115	-
23	PQN	B	843	-	-	9/23/43/43	0/2/2/2
24	BCR	B	846	-	-	17/29/63/63	0/2/2/2
22	CLA	7	315	14	1/1/12/20	8/19/97/115	-
22	CLA	B	834	-	1/1/13/20	15/29/107/115	-
38	LUT	7	302	-	-	5/29/67/67	0/2/2/2
22	CLA	8	309	-	1/1/15/20	17/37/115/115	-
22	CLA	K	205	10	1/1/13/20	15/25/103/115	-
22	CLA	A	832	-	1/1/15/20	17/37/115/115	-
24	BCR	3	305	-	-	14/29/63/63	0/2/2/2
25	LHG	9	315	22	-	20/37/37/53	-
22	CLA	6	310	18	1/1/15/20	16/37/115/115	-
22	CLA	1	308	-	1/1/13/20	13/25/103/115	-
22	CLA	7	309	-	1/1/13/20	14/27/105/115	-
22	CLA	A	811	22	1/1/15/20	20/37/115/115	-
24	BCR	6	305	-	-	13/29/63/63	0/2/2/2
22	CLA	B	814	-	1/1/14/20	19/31/109/115	-
22	CLA	9	305	19	1/1/14/20	18/31/109/115	-
25	LHG	A	849	22	-	27/39/39/53	-
22	CLA	3	320	13	1/1/11/20	5/15/93/115	-
38	LUT	5	303	-	-	6/29/67/67	0/2/2/2
38	LUT	8	302	-	-	8/29/67/67	0/2/2/2
22	CLA	5	307	17	1/1/14/20	14/31/109/115	-
39	CHL	8	312	-	3/3/18/26	9/27/125/137	-
39	CHL	Z	312	-	3/3/16/26	9/18/116/137	-
25	LHG	B	851	-	-	15/22/22/53	-

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Mol	Type	Chain	Res	Link	Chirals	Torsions	Rings
29	SF4	C	102	3	-	-	0/6/5/5
22	CLA	A	826	44	1/1/12/20	6/19/97/115	-
22	CLA	5	320	-	1/1/12/20	9/19/97/115	-
22	CLA	A	821	44	1/1/15/20	12/37/115/115	-
22	CLA	A	837	-	1/1/12/20	10/21/99/115	-
38	LUT	9	301	-	-	5/29/67/67	0/2/2/2
38	LUT	9	302	-	-	4/29/67/67	0/2/2/2
22	CLA	B	830	-	1/1/15/20	13/37/115/115	-
21	CL0	A	801	-	3/3/20/25	13/37/135/135	-

All (3038) bond length outliers are listed below:

Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
38	6	303	LUT	C24-C25	23.39	1.62	1.33
38	5	303	LUT	C24-C25	23.37	1.62	1.33
38	1	302	LUT	C24-C25	23.34	1.62	1.33
38	7	302	LUT	C24-C25	23.27	1.62	1.33
38	Z	302	LUT	C24-C25	23.26	1.62	1.33
38	6	304	LUT	C24-C25	23.22	1.62	1.33
38	5	302	LUT	C24-C25	23.21	1.62	1.33
38	4	803	LUT	C24-C25	23.19	1.62	1.33
38	1	303	LUT	C24-C25	23.18	1.62	1.33
38	9	302	LUT	C24-C25	23.17	1.62	1.33
38	9	301	LUT	C24-C25	23.16	1.62	1.33
38	8	303	LUT	C24-C25	23.12	1.62	1.33
38	3	303	LUT	C24-C25	23.10	1.61	1.33
38	4	802	LUT	C24-C25	23.09	1.61	1.33
38	3	302	LUT	C24-C25	23.07	1.61	1.33
38	Z	301	LUT	C24-C25	23.04	1.61	1.33
38	7	301	LUT	C24-C25	22.90	1.61	1.33
38	8	302	LUT	C24-C25	22.74	1.61	1.33
36	J	105	C7Z	C25-C26	15.85	1.61	1.34
32	F	306	RRX	C26-C25	15.80	1.61	1.34
36	5	306	C7Z	C25-C26	15.76	1.61	1.34
24	5	305	BCR	C26-C25	15.60	1.61	1.34
24	B	848	BCR	C26-C25	15.57	1.61	1.34
24	6	306	BCR	C26-C25	15.55	1.61	1.34
24	F	303	BCR	C26-C25	15.52	1.61	1.34
24	7	303	BCR	C26-C25	15.49	1.61	1.34
24	A	846	BCR	C26-C25	15.46	1.61	1.34

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
24	3	304	BCR	C26-C25	15.45	1.61	1.34
24	4	804	BCR	C26-C25	15.45	1.61	1.34
24	6	305	BCR	C26-C25	15.42	1.61	1.34
24	8	304	BCR	C26-C25	15.42	1.61	1.34
24	B	844	BCR	C26-C25	15.41	1.61	1.34
24	B	846	BCR	C26-C25	15.40	1.61	1.34
24	A	847	BCR	C26-C25	15.39	1.61	1.34
24	G	203	BCR	C26-C25	15.36	1.61	1.34
24	B	853	BCR	C26-C25	15.36	1.61	1.34
24	A	848	BCR	C26-C25	15.31	1.61	1.34
24	3	307	BCR	C26-C25	15.31	1.61	1.34
24	A	845	BCR	C26-C25	15.30	1.60	1.34
38	4	802	LUT	C5-C6	15.30	1.60	1.34
24	L	204	BCR	C26-C25	15.29	1.60	1.34
38	6	303	LUT	C5-C6	15.28	1.60	1.34
24	B	849	BCR	C26-C25	15.27	1.60	1.34
38	5	303	LUT	C5-C6	15.26	1.60	1.34
24	3	306	BCR	C26-C25	15.26	1.60	1.34
38	1	302	LUT	C5-C6	15.22	1.60	1.34
38	9	302	LUT	C5-C6	15.21	1.60	1.34
38	7	301	LUT	C5-C6	15.19	1.60	1.34
38	Z	301	LUT	C5-C6	15.19	1.60	1.34
38	9	301	LUT	C5-C6	15.17	1.60	1.34
24	K	206	BCR	C26-C25	15.17	1.60	1.34
24	3	305	BCR	C26-C25	15.16	1.60	1.34
38	7	302	LUT	C5-C6	15.15	1.60	1.34
38	8	302	LUT	C5-C6	15.14	1.60	1.34
24	B	847	BCR	C26-C25	15.13	1.60	1.34
38	6	304	LUT	C5-C6	15.11	1.60	1.34
38	Z	302	LUT	C5-C6	15.10	1.60	1.34
24	I	4001	BCR	C26-C25	15.10	1.60	1.34
24	B	845	BCR	C26-C25	15.10	1.60	1.34
38	8	303	LUT	C5-C6	15.09	1.60	1.34
24	A	844	BCR	C26-C25	15.09	1.60	1.34
36	J	105	C7Z	C5-C6	15.09	1.60	1.34
24	5	304	BCR	C26-C25	15.08	1.60	1.34
38	5	302	LUT	C5-C6	15.07	1.60	1.34
38	3	302	LUT	C5-C6	15.07	1.60	1.34
38	3	303	LUT	C5-C6	15.06	1.60	1.34
38	1	303	LUT	C5-C6	15.05	1.60	1.34
38	4	803	LUT	C5-C6	15.01	1.60	1.34
24	A	856	BCR	C26-C25	15.01	1.60	1.34

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
24	J	104	BCR	C26-C25	15.00	1.60	1.34
36	5	306	C7Z	C5-C6	14.91	1.60	1.34
32	F	306	RRX	C5-C6	14.38	1.59	1.34
24	3	304	BCR	C5-C6	14.21	1.59	1.34
24	B	849	BCR	C5-C6	14.20	1.59	1.34
24	4	804	BCR	C5-C6	14.20	1.59	1.34
24	F	303	BCR	C5-C6	14.11	1.58	1.34
24	A	844	BCR	C5-C6	14.08	1.58	1.34
24	B	853	BCR	C5-C6	14.08	1.58	1.34
24	A	847	BCR	C5-C6	14.08	1.58	1.34
24	B	847	BCR	C5-C6	14.06	1.58	1.34
24	3	305	BCR	C5-C6	14.04	1.58	1.34
24	A	846	BCR	C5-C6	14.02	1.58	1.34
24	B	844	BCR	C5-C6	14.01	1.58	1.34
24	G	203	BCR	C5-C6	14.00	1.58	1.34
24	K	206	BCR	C5-C6	13.99	1.58	1.34
24	B	846	BCR	C5-C6	13.99	1.58	1.34
24	5	304	BCR	C5-C6	13.97	1.58	1.34
24	6	305	BCR	C5-C6	13.96	1.58	1.34
24	5	305	BCR	C5-C6	13.94	1.58	1.34
24	A	856	BCR	C5-C6	13.93	1.58	1.34
24	6	306	BCR	C5-C6	13.91	1.58	1.34
24	B	845	BCR	C5-C6	13.88	1.58	1.34
24	B	848	BCR	C5-C6	13.87	1.58	1.34
24	J	104	BCR	C5-C6	13.82	1.58	1.34
24	I	4001	BCR	C5-C6	13.81	1.58	1.34
24	3	306	BCR	C5-C6	13.81	1.58	1.34
24	A	845	BCR	C5-C6	13.79	1.58	1.34
24	8	304	BCR	C5-C6	13.73	1.58	1.34
24	L	204	BCR	C5-C6	13.72	1.58	1.34
24	A	848	BCR	C5-C6	13.68	1.58	1.34
24	3	307	BCR	C5-C6	13.67	1.58	1.34
24	7	303	BCR	C5-C6	13.64	1.58	1.34
38	4	802	LUT	C22-C21	-13.09	1.38	1.54
38	5	302	LUT	C22-C21	-13.07	1.38	1.54
38	5	303	LUT	C22-C21	-13.06	1.38	1.54
38	6	303	LUT	C22-C21	-13.04	1.38	1.54
38	4	803	LUT	C22-C21	-12.98	1.38	1.54
38	8	303	LUT	C22-C21	-12.98	1.38	1.54
38	6	304	LUT	C22-C21	-12.93	1.38	1.54
38	Z	302	LUT	C22-C21	-12.93	1.38	1.54
38	7	301	LUT	C22-C21	-12.88	1.38	1.54

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
38	9	302	LUT	C22-C21	-12.85	1.38	1.54
38	7	302	LUT	C22-C21	-12.84	1.38	1.54
38	1	303	LUT	C22-C21	-12.84	1.38	1.54
38	9	301	LUT	C22-C21	-12.83	1.38	1.54
38	3	303	LUT	C22-C21	-12.82	1.38	1.54
38	1	302	LUT	C22-C21	-12.80	1.38	1.54
38	3	302	LUT	C22-C21	-12.72	1.38	1.54
38	Z	301	LUT	C22-C21	-12.65	1.38	1.54
38	8	302	LUT	C22-C21	-12.53	1.39	1.54
36	J	105	C7Z	C24-C23	11.66	1.72	1.52
36	5	306	C7Z	C24-C23	11.62	1.72	1.52
38	4	802	LUT	C2-C3	-11.26	1.36	1.52
38	7	301	LUT	C2-C3	-11.09	1.36	1.52
38	9	301	LUT	C2-C3	-11.06	1.36	1.52
38	5	302	LUT	C2-C3	-11.04	1.36	1.52
38	9	302	LUT	C2-C3	-11.03	1.36	1.52
38	3	302	LUT	C2-C3	-11.03	1.36	1.52
38	8	302	LUT	C2-C3	-11.02	1.36	1.52
36	J	105	C7Z	C22-C23	-11.02	1.36	1.52
38	5	303	LUT	C2-C3	-11.01	1.36	1.52
38	3	303	LUT	C2-C3	-10.97	1.36	1.52
36	5	306	C7Z	C22-C23	-10.95	1.36	1.52
38	1	302	LUT	C2-C3	-10.95	1.36	1.52
38	7	302	LUT	C2-C3	-10.92	1.36	1.52
38	4	803	LUT	C2-C3	-10.88	1.36	1.52
38	6	304	LUT	C2-C3	-10.87	1.36	1.52
38	1	303	LUT	C2-C3	-10.85	1.36	1.52
38	Z	302	LUT	C2-C3	-10.85	1.36	1.52
38	8	303	LUT	C2-C3	-10.84	1.36	1.52
38	Z	301	LUT	C2-C3	-10.81	1.36	1.52
38	6	303	LUT	C2-C3	-10.79	1.36	1.52
32	F	306	RRX	C29-C28	-10.64	1.37	1.52
36	J	105	C7Z	C2-C3	-9.85	1.38	1.52
36	5	306	C7Z	C2-C3	-9.83	1.38	1.52
21	A	801	CL0	MG-NA	9.22	2.28	2.06
36	5	306	C7Z	C4-C3	8.54	1.67	1.52
36	J	105	C7Z	C4-C3	8.47	1.67	1.52
38	1	303	LUT	C22-C23	8.33	1.66	1.53
38	3	302	LUT	C22-C23	8.26	1.66	1.53
38	6	303	LUT	C22-C23	8.18	1.66	1.53
38	8	302	LUT	C22-C23	8.17	1.66	1.53
38	Z	301	LUT	C22-C23	8.15	1.66	1.53

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
38	7	301	LUT	C22-C23	8.13	1.66	1.53
38	9	302	LUT	C22-C23	8.12	1.66	1.53
38	5	303	LUT	C22-C23	8.10	1.66	1.53
38	8	303	LUT	C22-C23	8.08	1.66	1.53
38	7	302	LUT	C22-C23	8.08	1.66	1.53
38	4	803	LUT	C22-C23	8.08	1.66	1.53
32	F	306	RRX	C27-C28	8.05	1.66	1.52
38	Z	302	LUT	C22-C23	8.05	1.66	1.53
38	6	304	LUT	C22-C23	8.04	1.66	1.53
38	3	302	LUT	C4-C3	8.02	1.66	1.52
38	1	302	LUT	C4-C3	8.02	1.66	1.52
38	1	302	LUT	C22-C23	7.99	1.66	1.53
38	5	302	LUT	C22-C23	7.99	1.66	1.53
38	9	301	LUT	C22-C23	7.98	1.66	1.53
38	3	303	LUT	C22-C23	7.97	1.66	1.53
38	Z	301	LUT	C4-C3	7.96	1.66	1.52
38	4	802	LUT	C22-C23	7.96	1.66	1.53
38	1	303	LUT	C4-C3	7.95	1.66	1.52
38	6	303	LUT	C4-C3	7.92	1.66	1.52
38	Z	302	LUT	C4-C3	7.90	1.66	1.52
38	8	302	LUT	C4-C3	7.88	1.65	1.52
38	5	303	LUT	C4-C3	7.87	1.65	1.52
38	5	302	LUT	C4-C3	7.87	1.65	1.52
38	9	301	LUT	C4-C3	7.86	1.65	1.52
38	6	304	LUT	C4-C3	7.85	1.65	1.52
38	3	303	LUT	C4-C3	7.83	1.65	1.52
38	8	303	LUT	C4-C3	7.81	1.65	1.52
38	7	301	LUT	C4-C3	7.80	1.65	1.52
38	7	302	LUT	C4-C3	7.79	1.65	1.52
38	4	803	LUT	C4-C3	7.78	1.65	1.52
38	4	802	LUT	C4-C3	7.73	1.65	1.52
38	9	302	LUT	C4-C3	7.62	1.65	1.52
38	Z	301	LUT	C32-C33	7.61	1.62	1.45
38	7	302	LUT	C32-C33	7.50	1.62	1.45
38	1	303	LUT	C32-C33	7.49	1.62	1.45
38	1	302	LUT	C32-C33	7.49	1.62	1.45
38	5	303	LUT	C32-C33	7.47	1.62	1.45
38	Z	302	LUT	C32-C33	7.46	1.62	1.45
38	7	301	LUT	C32-C33	7.43	1.61	1.45
38	6	304	LUT	C32-C33	7.42	1.61	1.45
38	3	303	LUT	C32-C33	7.42	1.61	1.45
38	6	303	LUT	C32-C33	7.40	1.61	1.45

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
38	3	302	LUT	C32-C33	7.40	1.61	1.45
38	9	301	LUT	C32-C33	7.39	1.61	1.45
38	4	802	LUT	C32-C33	7.36	1.61	1.45
38	8	303	LUT	C32-C33	7.36	1.61	1.45
38	9	302	LUT	C32-C33	7.35	1.61	1.45
38	4	803	LUT	C32-C33	7.35	1.61	1.45
38	5	302	LUT	C32-C33	7.32	1.61	1.45
38	8	302	LUT	C32-C33	7.28	1.61	1.45
24	3	306	BCR	C30-C25	-7.11	1.44	1.53
24	5	304	BCR	C30-C25	-7.10	1.44	1.53
24	B	845	BCR	C30-C25	-7.03	1.44	1.53
24	L	204	BCR	C30-C25	-7.03	1.44	1.53
24	B	847	BCR	C30-C25	-7.02	1.44	1.53
24	A	844	BCR	C30-C25	-7.02	1.44	1.53
24	I	4001	BCR	C30-C25	-7.02	1.44	1.53
24	A	847	BCR	C30-C25	-6.99	1.44	1.53
24	K	206	BCR	C30-C25	-6.95	1.44	1.53
24	8	304	BCR	C1-C6	-6.89	1.44	1.53
24	3	305	BCR	C30-C25	-6.89	1.44	1.53
24	I	4001	BCR	C1-C6	-6.88	1.44	1.53
24	B	853	BCR	C30-C25	-6.86	1.44	1.53
24	B	849	BCR	C30-C25	-6.86	1.44	1.53
24	A	856	BCR	C30-C25	-6.85	1.44	1.53
24	J	104	BCR	C30-C25	-6.84	1.44	1.53
24	B	848	BCR	C30-C25	-6.80	1.44	1.53
24	A	848	BCR	C30-C25	-6.80	1.44	1.53
24	3	307	BCR	C30-C25	-6.79	1.44	1.53
24	4	804	BCR	C30-C25	-6.79	1.44	1.53
24	B	844	BCR	C30-C25	-6.78	1.44	1.53
24	3	304	BCR	C2-C3	-6.78	1.35	1.52
24	A	845	BCR	C30-C25	-6.77	1.44	1.53
24	A	846	BCR	C30-C25	-6.77	1.44	1.53
24	5	305	BCR	C30-C25	-6.77	1.44	1.53
24	6	305	BCR	C30-C25	-6.74	1.44	1.53
24	6	306	BCR	C30-C25	-6.73	1.44	1.53
24	7	303	BCR	C30-C25	-6.72	1.44	1.53
24	3	304	BCR	C30-C25	-6.71	1.44	1.53
24	B	847	BCR	C1-C6	-6.70	1.44	1.53
24	L	204	BCR	C1-C6	-6.70	1.44	1.53
24	F	303	BCR	C30-C25	-6.69	1.44	1.53
24	8	304	BCR	C30-C25	-6.69	1.44	1.53
24	G	203	BCR	C2-C3	-6.69	1.36	1.52

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
24	B	845	BCR	C1-C6	-6.68	1.44	1.53
24	6	305	BCR	C2-C3	-6.68	1.36	1.52
24	A	844	BCR	C2-C3	-6.67	1.36	1.52
24	5	304	BCR	C2-C3	-6.67	1.36	1.52
24	B	844	BCR	C2-C3	-6.67	1.36	1.52
24	K	206	BCR	C2-C3	-6.66	1.36	1.52
24	3	305	BCR	C2-C3	-6.66	1.36	1.52
24	3	307	BCR	C1-C6	-6.66	1.44	1.53
24	B	849	BCR	C2-C3	-6.66	1.36	1.52
24	B	848	BCR	C2-C3	-6.65	1.36	1.52
24	4	804	BCR	C2-C3	-6.65	1.36	1.52
24	F	303	BCR	C2-C3	-6.65	1.36	1.52
24	B	846	BCR	C2-C3	-6.64	1.36	1.52
24	5	304	BCR	C1-C6	-6.63	1.44	1.53
24	A	846	BCR	C2-C3	-6.63	1.36	1.52
24	B	846	BCR	C30-C25	-6.62	1.44	1.53
24	K	206	BCR	C1-C6	-6.62	1.44	1.53
24	5	305	BCR	C2-C3	-6.61	1.36	1.52
24	A	847	BCR	C2-C3	-6.61	1.36	1.52
24	B	847	BCR	C2-C3	-6.61	1.36	1.52
24	A	848	BCR	C1-C6	-6.59	1.44	1.53
24	A	856	BCR	C2-C3	-6.57	1.36	1.52
24	B	853	BCR	C2-C3	-6.57	1.36	1.52
24	G	203	BCR	C30-C25	-6.56	1.44	1.53
24	A	847	BCR	C1-C6	-6.54	1.44	1.53
24	3	306	BCR	C2-C3	-6.54	1.36	1.52
24	6	306	BCR	C2-C3	-6.53	1.36	1.52
24	I	4001	BCR	C2-C3	-6.53	1.36	1.52
32	F	306	RRX	C2-C3	-6.51	1.36	1.52
24	6	306	BCR	C1-C6	-6.51	1.44	1.53
24	A	845	BCR	C2-C3	-6.50	1.36	1.52
24	B	846	BCR	C1-C6	-6.50	1.44	1.53
22	4	805	CLA	MG-NA	6.49	2.21	2.06
24	G	203	BCR	C1-C6	-6.48	1.44	1.53
32	F	306	RRX	C30-C25	-6.48	1.44	1.53
24	3	305	BCR	C1-C6	-6.48	1.44	1.53
24	J	104	BCR	C1-C6	-6.47	1.44	1.53
24	B	845	BCR	C2-C3	-6.46	1.36	1.52
24	J	104	BCR	C2-C3	-6.45	1.36	1.52
24	A	846	BCR	C1-C6	-6.45	1.44	1.53
24	A	856	BCR	C1-C6	-6.45	1.44	1.53
24	L	204	BCR	C2-C3	-6.45	1.36	1.52

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
24	8	304	BCR	C2-C3	-6.44	1.36	1.52
24	A	848	BCR	C2-C3	-6.43	1.36	1.52
22	8	307	CLA	MG-NA	6.42	2.21	2.06
24	5	305	BCR	C29-C28	-6.42	1.36	1.52
24	3	307	BCR	C2-C3	-6.41	1.36	1.52
24	6	305	BCR	C29-C28	-6.41	1.36	1.52
24	A	847	BCR	C29-C28	-6.41	1.36	1.52
24	B	844	BCR	C1-C6	-6.40	1.45	1.53
24	B	849	BCR	C1-C6	-6.40	1.45	1.53
24	7	303	BCR	C29-C28	-6.40	1.36	1.52
24	A	845	BCR	C1-C6	-6.40	1.45	1.53
22	4	807	CLA	MG-NA	6.40	2.21	2.06
24	B	848	BCR	C29-C28	-6.39	1.36	1.52
22	4	806	CLA	MG-NA	6.38	2.21	2.06
22	7	308	CLA	MG-NA	6.38	2.21	2.06
22	4	809	CLA	MG-NA	6.38	2.21	2.06
24	4	804	BCR	C29-C28	-6.38	1.36	1.52
38	6	303	LUT	C1-C6	-6.38	1.45	1.53
24	F	303	BCR	C29-C28	-6.38	1.36	1.52
22	6	309	CLA	MG-NA	6.38	2.21	2.06
22	A	816	CLA	MG-NA	6.37	2.21	2.06
22	5	311	CLA	MG-NA	6.37	2.21	2.06
24	8	304	BCR	C29-C28	-6.37	1.36	1.52
22	B	820	CLA	MG-NA	6.37	2.21	2.06
22	7	306	CLA	MG-NA	6.36	2.21	2.06
22	3	310	CLA	MG-NA	6.36	2.21	2.06
22	A	817	CLA	MG-NA	6.36	2.21	2.06
22	3	312	CLA	MG-NA	6.36	2.21	2.06
24	7	303	BCR	C2-C3	-6.36	1.36	1.52
24	6	306	BCR	C29-C28	-6.36	1.36	1.52
22	A	825	CLA	MG-NA	6.36	2.21	2.06
22	5	315	CLA	MG-NA	6.35	2.21	2.06
22	B	828	CLA	MG-NA	6.35	2.21	2.06
22	Z	313	CLA	MG-NA	6.34	2.21	2.06
22	7	324	CLA	MG-NA	6.34	2.21	2.06
24	B	853	BCR	C1-C6	-6.34	1.45	1.53
22	6	301	CLA	MG-NA	6.34	2.21	2.06
22	A	837	CLA	MG-NA	6.34	2.21	2.06
24	7	303	BCR	C1-C6	-6.33	1.45	1.53
22	1	315	CLA	MG-NA	6.33	2.21	2.06
22	5	313	CLA	MG-NA	6.33	2.21	2.06
22	G	201	CLA	MG-NA	6.33	2.21	2.06

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
22	B	826	CLA	MG-NA	6.33	2.21	2.06
24	3	304	BCR	C29-C28	-6.33	1.36	1.52
22	A	820	CLA	MG-NA	6.32	2.21	2.06
24	B	844	BCR	C29-C28	-6.32	1.37	1.52
22	A	822	CLA	MG-NA	6.32	2.21	2.06
24	B	846	BCR	C29-C28	-6.32	1.37	1.52
22	5	322	CLA	MG-NA	6.32	2.21	2.06
24	A	846	BCR	C29-C28	-6.32	1.37	1.52
24	4	804	BCR	C1-C6	-6.32	1.45	1.53
22	A	813	CLA	MG-NA	6.32	2.21	2.06
22	A	829	CLA	MG-NA	6.32	2.21	2.06
22	F	301	CLA	MG-NA	6.31	2.21	2.06
22	K	203	CLA	MG-NA	6.31	2.21	2.06
22	9	308	CLA	MG-NA	6.31	2.21	2.06
22	1	310	CLA	MG-NA	6.31	2.21	2.06
22	8	306	CLA	MG-NA	6.31	2.21	2.06
24	G	203	BCR	C29-C28	-6.31	1.37	1.52
22	5	326	CLA	MG-NA	6.31	2.21	2.06
22	3	322	CLA	MG-NA	6.31	2.21	2.06
22	5	309	CLA	MG-NA	6.31	2.21	2.06
22	B	810	CLA	MG-NA	6.31	2.21	2.06
22	L	203	CLA	MG-NA	6.31	2.21	2.06
22	B	824	CLA	MG-NA	6.30	2.21	2.06
22	B	840	CLA	MG-NA	6.30	2.21	2.06
22	7	323	CLA	MG-NA	6.30	2.21	2.06
22	A	840	CLA	MG-NA	6.30	2.21	2.06
22	L	202	CLA	MG-NA	6.30	2.21	2.06
22	4	811	CLA	MG-NA	6.30	2.21	2.06
22	1	305	CLA	MG-NA	6.30	2.21	2.06
22	A	834	CLA	MG-NA	6.30	2.21	2.06
22	J	103	CLA	MG-NA	6.30	2.21	2.06
22	B	829	CLA	MG-NA	6.30	2.21	2.06
22	Z	315	CLA	MG-NA	6.30	2.21	2.06
22	8	309	CLA	MG-NA	6.30	2.21	2.06
22	9	307	CLA	MG-NA	6.30	2.21	2.06
22	6	319	CLA	MG-NA	6.30	2.21	2.06
22	B	834	CLA	MG-NA	6.30	2.21	2.06
22	B	806	CLA	MG-NA	6.30	2.21	2.06
22	1	304	CLA	MG-NA	6.30	2.21	2.06
22	A	807	CLA	MG-NA	6.30	2.21	2.06
22	7	310	CLA	MG-NA	6.30	2.21	2.06
22	A	839	CLA	MG-NA	6.29	2.21	2.06

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
22	A	855	CLA	MG-NA	6.29	2.21	2.06
22	Z	304	CLA	MG-NA	6.29	2.21	2.06
22	K	202	CLA	MG-NA	6.29	2.21	2.06
22	2	302	CLA	MG-NA	6.29	2.21	2.06
22	F	302	CLA	MG-NA	6.29	2.21	2.06
22	Z	310	CLA	MG-NA	6.29	2.21	2.06
22	7	305	CLA	MG-NA	6.29	2.21	2.06
22	2	304	CLA	MG-NA	6.29	2.21	2.06
22	A	814	CLA	MG-NA	6.29	2.21	2.06
22	7	312	CLA	MG-NA	6.29	2.21	2.06
22	A	824	CLA	MG-NA	6.29	2.21	2.06
22	B	823	CLA	MG-NA	6.29	2.21	2.06
22	B	822	CLA	MG-NA	6.29	2.21	2.06
22	7	317	CLA	MG-NA	6.29	2.21	2.06
22	6	311	CLA	MG-NA	6.29	2.21	2.06
22	B	842	CLA	MG-NA	6.29	2.21	2.06
22	A	812	CLA	MG-NA	6.29	2.21	2.06
22	F	305	CLA	MG-NA	6.29	2.21	2.06
22	A	841	CLA	MG-NA	6.29	2.21	2.06
22	A	808	CLA	MG-NA	6.28	2.21	2.06
22	A	809	CLA	MG-NA	6.28	2.21	2.06
22	B	833	CLA	MG-NA	6.28	2.21	2.06
22	B	815	CLA	MG-NA	6.28	2.21	2.06
24	F	303	BCR	C1-C6	-6.28	1.45	1.53
22	6	322	CLA	MG-NA	6.28	2.21	2.06
22	4	812	CLA	MG-NA	6.28	2.21	2.06
22	B	837	CLA	MG-NA	6.28	2.21	2.06
22	9	313	CLA	MG-NA	6.28	2.21	2.06
22	K	204	CLA	MG-NA	6.28	2.21	2.06
22	A	826	CLA	MG-NA	6.28	2.21	2.06
22	4	808	CLA	MG-NA	6.28	2.21	2.06
22	A	827	CLA	MG-NA	6.28	2.21	2.06
22	Z	307	CLA	MG-NA	6.28	2.21	2.06
24	3	306	BCR	C29-C28	-6.28	1.37	1.52
22	7	315	CLA	MG-NA	6.27	2.21	2.06
22	B	805	CLA	MG-NA	6.27	2.21	2.06
22	6	314	CLA	MG-NA	6.27	2.21	2.06
22	7	314	CLA	MG-NA	6.27	2.21	2.06
22	A	810	CLA	MG-NA	6.27	2.21	2.06
22	A	842	CLA	MG-NA	6.27	2.21	2.06
22	L	201	CLA	MG-NA	6.27	2.21	2.06
22	3	319	CLA	MG-NA	6.27	2.21	2.06

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
22	B	827	CLA	MG-NA	6.27	2.21	2.06
22	3	309	CLA	MG-NA	6.26	2.21	2.06
22	3	316	CLA	MG-NA	6.26	2.21	2.06
22	6	313	CLA	MG-NA	6.26	2.21	2.06
22	B	839	CLA	MG-NA	6.26	2.21	2.06
22	7	311	CLA	MG-NA	6.26	2.21	2.06
22	6	321	CLA	MG-NA	6.26	2.21	2.06
22	B	821	CLA	MG-NA	6.26	2.21	2.06
22	9	311	CLA	MG-NA	6.26	2.21	2.06
38	8	303	LUT	C1-C6	-6.26	1.45	1.53
22	B	819	CLA	MG-NA	6.26	2.21	2.06
24	3	307	BCR	C29-C28	-6.26	1.37	1.52
22	1	307	CLA	MG-NA	6.26	2.21	2.06
22	1	306	CLA	MG-NA	6.26	2.21	2.06
22	B	811	CLA	MG-NA	6.25	2.21	2.06
22	6	312	CLA	MG-NA	6.25	2.21	2.06
22	Z	314	CLA	MG-NA	6.25	2.21	2.06
22	Z	316	CLA	MG-NA	6.25	2.21	2.06
22	A	830	CLA	MG-NA	6.25	2.21	2.06
22	A	802	CLA	MG-NA	6.25	2.21	2.06
22	7	316	CLA	MG-NA	6.25	2.21	2.06
22	4	818	CLA	MG-NA	6.25	2.21	2.06
22	6	302	CLA	MG-NA	6.25	2.21	2.06
38	Z	301	LUT	C1-C6	-6.25	1.45	1.53
24	K	206	BCR	C29-C28	-6.25	1.37	1.52
22	A	803	CLA	MG-NA	6.25	2.21	2.06
22	A	806	CLA	MG-NA	6.25	2.21	2.06
22	7	309	CLA	MG-NA	6.25	2.21	2.06
22	A	833	CLA	MG-NA	6.25	2.21	2.06
22	B	831	CLA	MG-NA	6.25	2.21	2.06
22	1	314	CLA	MG-NA	6.25	2.21	2.06
38	4	803	LUT	C1-C6	-6.25	1.45	1.53
22	8	314	CLA	MG-NA	6.25	2.21	2.06
22	A	815	CLA	MG-NA	6.25	2.21	2.06
24	A	845	BCR	C29-C28	-6.25	1.37	1.52
22	5	310	CLA	MG-NA	6.25	2.21	2.06
22	6	310	CLA	MG-NA	6.25	2.21	2.06
22	A	832	CLA	MG-NA	6.25	2.21	2.06
22	B	816	CLA	MG-NA	6.24	2.21	2.06
22	B	825	CLA	MG-NA	6.24	2.21	2.06
22	F	304	CLA	MG-NA	6.24	2.21	2.06
22	B	841	CLA	MG-NA	6.24	2.21	2.06

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
22	8	316	CLA	MG-NA	6.24	2.21	2.06
22	B	809	CLA	MG-NA	6.24	2.21	2.06
22	B	813	CLA	MG-NA	6.24	2.21	2.06
22	1	313	CLA	MG-NA	6.24	2.21	2.06
22	A	854	CLA	MG-NA	6.24	2.21	2.06
22	4	817	CLA	MG-NA	6.24	2.21	2.06
22	6	317	CLA	MG-NA	6.24	2.21	2.06
22	3	314	CLA	MG-NA	6.24	2.21	2.06
22	5	308	CLA	MG-NA	6.24	2.21	2.06
22	5	319	CLA	MG-NA	6.24	2.21	2.06
38	1	303	LUT	C1-C6	-6.23	1.45	1.53
22	5	314	CLA	MG-NA	6.23	2.21	2.06
22	A	811	CLA	MG-NA	6.23	2.21	2.06
22	Z	309	CLA	MG-NA	6.23	2.21	2.06
22	A	818	CLA	MG-NA	6.23	2.21	2.06
22	B	807	CLA	MG-NA	6.23	2.21	2.06
22	B	808	CLA	MG-NA	6.23	2.21	2.06
22	Z	308	CLA	MG-NA	6.23	2.21	2.06
22	8	311	CLA	MG-NA	6.23	2.21	2.06
22	5	307	CLA	MG-NA	6.23	2.21	2.06
24	A	848	BCR	C29-C28	-6.23	1.37	1.52
22	8	310	CLA	MG-NA	6.23	2.21	2.06
38	3	302	LUT	C1-C6	-6.23	1.45	1.53
22	A	831	CLA	MG-NA	6.23	2.21	2.06
22	B	812	CLA	MG-NA	6.23	2.21	2.06
22	B	830	CLA	MG-NA	6.22	2.21	2.06
22	4	815	CLA	MG-NA	6.22	2.21	2.06
22	G	202	CLA	MG-NA	6.22	2.21	2.06
22	5	312	CLA	MG-NA	6.22	2.21	2.06
22	B	836	CLA	MG-NA	6.22	2.21	2.06
22	A	804	CLA	MG-NA	6.22	2.21	2.06
22	A	819	CLA	MG-NA	6.22	2.21	2.06
22	4	810	CLA	MG-NA	6.22	2.21	2.06
24	A	844	BCR	C1-C6	-6.22	1.45	1.53
22	A	821	CLA	MG-NA	6.22	2.21	2.06
22	5	301	CLA	MG-NA	6.22	2.21	2.06
22	B	801	CLA	MG-NA	6.22	2.21	2.06
22	A	838	CLA	MG-NA	6.21	2.21	2.06
22	6	308	CLA	MG-NA	6.21	2.21	2.06
22	9	306	CLA	MG-NA	6.21	2.21	2.06
22	B	835	CLA	MG-NA	6.21	2.21	2.06
24	B	848	BCR	C1-C6	-6.21	1.45	1.53

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
22	3	308	CLA	MG-NA	6.21	2.21	2.06
24	5	305	BCR	C1-C6	-6.21	1.45	1.53
22	8	313	CLA	MG-NA	6.20	2.21	2.06
22	3	311	CLA	MG-NA	6.20	2.21	2.06
22	8	308	CLA	MG-NA	6.20	2.21	2.06
22	A	805	CLA	MG-NA	6.20	2.21	2.06
22	B	832	CLA	MG-NA	6.20	2.21	2.06
22	6	307	CLA	MG-NA	6.20	2.21	2.06
22	5	318	CLA	MG-NA	6.20	2.21	2.06
24	B	849	BCR	C29-C28	-6.20	1.37	1.52
22	7	322	CLA	MG-NA	6.20	2.21	2.06
22	B	817	CLA	MG-NA	6.19	2.21	2.06
24	3	306	BCR	C1-C6	-6.19	1.45	1.53
22	B	838	CLA	MG-NA	6.19	2.21	2.06
22	2	301	CLA	MG-NA	6.19	2.21	2.06
22	A	823	CLA	MG-NA	6.19	2.21	2.06
22	7	307	CLA	MG-NA	6.19	2.21	2.06
38	8	302	LUT	C1-C6	-6.18	1.45	1.53
22	A	828	CLA	MG-NA	6.18	2.21	2.06
22	Z	306	CLA	MG-NA	6.18	2.21	2.06
22	1	308	CLA	MG-NA	6.18	2.21	2.06
38	1	302	LUT	C1-C6	-6.18	1.45	1.53
22	3	318	CLA	MG-NA	6.18	2.20	2.06
38	Z	302	LUT	C1-C6	-6.18	1.45	1.53
24	3	305	BCR	C29-C28	-6.17	1.37	1.52
24	5	304	BCR	C29-C28	-6.17	1.37	1.52
24	B	847	BCR	C29-C28	-6.17	1.37	1.52
38	3	303	LUT	C1-C6	-6.16	1.45	1.53
24	6	305	BCR	C1-C6	-6.16	1.45	1.53
24	B	845	BCR	C29-C28	-6.15	1.37	1.52
38	5	302	LUT	C1-C6	-6.15	1.45	1.53
22	5	323	CLA	MG-NA	6.15	2.20	2.06
22	Z	303	CLA	MG-NA	6.15	2.20	2.06
38	6	304	LUT	C1-C6	-6.15	1.45	1.53
22	B	818	CLA	MG-NA	6.15	2.20	2.06
22	Z	305	CLA	MG-NA	6.15	2.20	2.06
22	A	835	CLA	MG-NA	6.15	2.20	2.06
24	A	844	BCR	C29-C28	-6.15	1.37	1.52
22	1	309	CLA	MG-NA	6.14	2.20	2.06
22	A	836	CLA	MG-NA	6.14	2.20	2.06
22	B	814	CLA	MG-NA	6.14	2.20	2.06
24	L	204	BCR	C29-C28	-6.13	1.37	1.52

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
32	F	306	RRX	C19-C18	6.12	1.59	1.45
22	3	313	CLA	MG-NA	6.12	2.20	2.06
24	3	304	BCR	C1-C6	-6.11	1.45	1.53
24	I	4001	BCR	C29-C28	-6.11	1.37	1.52
24	B	853	BCR	C29-C28	-6.10	1.37	1.52
22	7	304	CLA	MG-NA	6.10	2.20	2.06
22	8	305	CLA	MG-NA	6.10	2.20	2.06
22	1	316	CLA	MG-NA	6.09	2.20	2.06
24	J	104	BCR	C29-C28	-6.09	1.37	1.52
22	9	304	CLA	MG-NA	6.09	2.20	2.06
38	7	302	LUT	C1-C6	-6.09	1.45	1.53
38	9	301	LUT	C1-C6	-6.05	1.45	1.53
22	9	310	CLA	MG-NA	6.05	2.20	2.06
38	5	303	LUT	C1-C6	-6.04	1.45	1.53
24	A	856	BCR	C29-C28	-6.02	1.37	1.52
38	9	302	LUT	C1-C6	-6.02	1.45	1.53
36	J	105	C7Z	C12-C13	6.02	1.58	1.45
38	7	301	LUT	C1-C6	-6.01	1.45	1.53
24	B	853	BCR	C12-C13	6.00	1.58	1.45
22	K	205	CLA	MG-NA	5.99	2.20	2.06
22	9	309	CLA	MG-NA	5.96	2.20	2.06
22	3	315	CLA	MG-NA	5.95	2.20	2.06
22	5	320	CLA	MG-NA	5.93	2.20	2.06
36	5	306	C7Z	C12-C13	5.92	1.58	1.45
22	3	320	CLA	MG-NA	5.92	2.20	2.06
22	1	311	CLA	MG-NA	5.91	2.20	2.06
22	9	305	CLA	MG-NA	5.89	2.20	2.06
24	3	306	BCR	C12-C13	5.89	1.58	1.45
38	4	802	LUT	C1-C6	-5.87	1.45	1.53
24	3	304	BCR	C12-C13	5.85	1.58	1.45
24	A	844	BCR	C12-C13	5.82	1.58	1.45
24	J	104	BCR	C12-C13	5.80	1.58	1.45
24	A	848	BCR	C12-C13	5.79	1.58	1.45
24	B	846	BCR	C12-C13	5.75	1.58	1.45
24	A	856	BCR	C12-C13	5.74	1.58	1.45
22	9	303	CLA	MG-NA	5.73	2.19	2.06
24	A	845	BCR	C12-C13	5.73	1.58	1.45
24	B	849	BCR	C12-C13	5.73	1.58	1.45
24	B	845	BCR	C12-C13	5.72	1.58	1.45
24	B	844	BCR	C12-C13	5.72	1.58	1.45
24	G	203	BCR	C12-C13	5.72	1.58	1.45
24	3	307	BCR	C12-C13	5.71	1.58	1.45

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
24	I	4001	BCR	C12-C13	5.71	1.58	1.45
24	F	303	BCR	C12-C13	5.71	1.58	1.45
24	7	303	BCR	C12-C13	5.71	1.58	1.45
24	3	305	BCR	C12-C13	5.68	1.58	1.45
24	6	305	BCR	C12-C13	5.68	1.58	1.45
24	4	804	BCR	C12-C13	5.68	1.58	1.45
24	8	304	BCR	C12-C13	5.68	1.58	1.45
24	B	853	BCR	C8-C9	5.67	1.58	1.45
24	5	304	BCR	C12-C13	5.67	1.58	1.45
24	B	847	BCR	C12-C13	5.65	1.58	1.45
24	5	305	BCR	C12-C13	5.64	1.58	1.45
24	L	204	BCR	C12-C13	5.62	1.58	1.45
24	A	846	BCR	C12-C13	5.62	1.58	1.45
24	K	206	BCR	C12-C13	5.62	1.58	1.45
24	A	847	BCR	C12-C13	5.60	1.58	1.45
24	B	848	BCR	C12-C13	5.60	1.58	1.45
24	6	306	BCR	C12-C13	5.59	1.58	1.45
24	B	844	BCR	C8-C9	5.56	1.57	1.45
24	4	804	BCR	C8-C9	5.52	1.57	1.45
24	I	4001	BCR	C8-C9	5.51	1.57	1.45
24	A	856	BCR	C8-C9	5.49	1.57	1.45
24	3	307	BCR	C8-C9	5.48	1.57	1.45
24	B	845	BCR	C8-C9	5.46	1.57	1.45
24	F	303	BCR	C8-C9	5.45	1.57	1.45
24	3	305	BCR	C8-C9	5.44	1.57	1.45
24	B	849	BCR	C8-C9	5.44	1.57	1.45
24	J	104	BCR	C8-C9	5.44	1.57	1.45
24	A	845	BCR	C8-C9	5.43	1.57	1.45
24	B	846	BCR	C8-C9	5.43	1.57	1.45
24	A	848	BCR	C8-C9	5.42	1.57	1.45
32	F	306	RRX	C1-C6	-5.41	1.46	1.53
24	L	204	BCR	C8-C9	5.41	1.57	1.45
24	3	306	BCR	C8-C9	5.41	1.57	1.45
36	5	306	C7Z	C1-C6	-5.41	1.46	1.53
24	5	305	BCR	C8-C9	5.40	1.57	1.45
24	B	853	BCR	C23-C22	5.40	1.57	1.45
24	B	848	BCR	C8-C9	5.40	1.57	1.45
24	7	303	BCR	C8-C9	5.39	1.57	1.45
24	6	305	BCR	C8-C9	5.39	1.57	1.45
24	8	304	BCR	C8-C9	5.38	1.57	1.45
24	A	846	BCR	C8-C9	5.37	1.57	1.45
24	A	856	BCR	C23-C22	5.37	1.57	1.45

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
24	8	304	BCR	C29-C30	5.35	1.66	1.54
24	3	304	BCR	C8-C9	5.35	1.57	1.45
24	3	305	BCR	C23-C22	5.35	1.57	1.45
24	5	304	BCR	C8-C9	5.34	1.57	1.45
24	A	847	BCR	C8-C9	5.34	1.57	1.45
24	K	206	BCR	C8-C9	5.34	1.57	1.45
24	A	856	BCR	C29-C30	5.33	1.66	1.54
24	B	847	BCR	C8-C9	5.32	1.57	1.45
24	5	304	BCR	C29-C30	5.32	1.66	1.54
24	G	203	BCR	C8-C9	5.32	1.57	1.45
24	8	304	BCR	C23-C22	5.31	1.57	1.45
24	B	846	BCR	C29-C30	5.31	1.66	1.54
24	G	203	BCR	C29-C30	5.31	1.66	1.54
24	F	303	BCR	C23-C22	5.31	1.57	1.45
24	B	849	BCR	C29-C30	5.30	1.66	1.54
24	J	104	BCR	C29-C30	5.30	1.66	1.54
24	A	848	BCR	C23-C22	5.30	1.57	1.45
24	B	849	BCR	C23-C22	5.29	1.57	1.45
24	A	844	BCR	C8-C9	5.29	1.57	1.45
24	6	306	BCR	C8-C9	5.29	1.57	1.45
24	K	206	BCR	C23-C22	5.28	1.57	1.45
24	3	304	BCR	C29-C30	5.28	1.66	1.54
32	F	306	RRX	C8-C9	5.28	1.57	1.45
24	3	305	BCR	C29-C30	5.28	1.66	1.54
24	A	845	BCR	C23-C22	5.28	1.57	1.45
24	7	303	BCR	C23-C22	5.27	1.57	1.45
24	I	4001	BCR	C29-C30	5.27	1.66	1.54
24	5	304	BCR	C23-C22	5.27	1.57	1.45
24	B	844	BCR	C23-C22	5.27	1.57	1.45
24	A	846	BCR	C29-C30	5.27	1.66	1.54
24	3	306	BCR	C23-C22	5.27	1.57	1.45
24	A	844	BCR	C29-C30	5.27	1.66	1.54
24	3	304	BCR	C23-C22	5.26	1.57	1.45
24	J	104	BCR	C23-C22	5.26	1.57	1.45
24	K	206	BCR	C29-C30	5.25	1.66	1.54
24	A	845	BCR	C29-C30	5.25	1.66	1.54
24	B	847	BCR	C29-C30	5.25	1.66	1.54
24	B	845	BCR	C23-C22	5.25	1.57	1.45
24	4	804	BCR	C23-C22	5.24	1.57	1.45
36	J	105	C7Z	C1-C6	-5.24	1.46	1.53
24	6	305	BCR	C29-C30	5.24	1.66	1.54
24	B	846	BCR	C23-C22	5.23	1.57	1.45

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
24	A	848	BCR	C29-C30	5.22	1.66	1.54
24	L	204	BCR	C29-C30	5.22	1.66	1.54
24	3	306	BCR	C29-C30	5.21	1.66	1.54
24	B	845	BCR	C29-C30	5.21	1.66	1.54
24	G	203	BCR	C23-C22	5.20	1.57	1.45
21	A	801	CL0	CHC-C1C	5.20	1.48	1.35
24	3	307	BCR	C29-C30	5.20	1.66	1.54
24	6	305	BCR	C23-C22	5.20	1.57	1.45
24	B	844	BCR	C29-C30	5.19	1.66	1.54
24	A	847	BCR	C23-C22	5.19	1.57	1.45
24	B	847	BCR	C23-C22	5.19	1.57	1.45
24	L	204	BCR	C23-C22	5.18	1.57	1.45
24	A	846	BCR	C23-C22	5.18	1.57	1.45
24	A	844	BCR	C23-C22	5.18	1.57	1.45
24	F	303	BCR	C29-C30	5.18	1.66	1.54
24	3	307	BCR	C23-C22	5.18	1.57	1.45
24	B	848	BCR	C23-C22	5.16	1.57	1.45
24	5	305	BCR	C23-C22	5.16	1.57	1.45
24	6	306	BCR	C29-C30	5.16	1.66	1.54
24	I	4001	BCR	C23-C22	5.15	1.57	1.45
21	A	801	CL0	O2D-CGD	5.15	1.45	1.33
24	4	804	BCR	C29-C30	5.14	1.66	1.54
24	7	303	BCR	C29-C30	5.11	1.65	1.54
24	B	848	BCR	C29-C30	5.10	1.65	1.54
21	A	801	CL0	O2A-C1	5.10	1.60	1.46
24	5	305	BCR	C29-C30	5.09	1.65	1.54
38	1	302	LUT	C8-C9	5.07	1.56	1.45
38	Z	301	LUT	C8-C9	5.03	1.56	1.45
38	6	303	LUT	C8-C9	5.02	1.56	1.45
32	F	306	RRX	C2-C1	5.02	1.65	1.54
36	5	306	C7Z	C24-C25	-5.01	1.43	1.51
24	B	853	BCR	C15-C14	5.01	1.59	1.43
38	7	302	LUT	C8-C9	4.99	1.56	1.45
38	Z	302	LUT	C8-C9	4.99	1.56	1.45
38	8	303	LUT	C8-C9	4.98	1.56	1.45
38	1	303	LUT	C8-C9	4.98	1.56	1.45
38	6	304	LUT	C8-C9	4.96	1.56	1.45
24	6	306	BCR	C23-C22	4.96	1.56	1.45
24	A	847	BCR	C29-C30	4.96	1.65	1.54
38	5	303	LUT	C8-C9	4.95	1.56	1.45
38	9	302	LUT	C8-C9	4.94	1.56	1.45
38	7	301	LUT	C8-C9	4.94	1.56	1.45

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
38	9	301	LUT	C8-C9	4.93	1.56	1.45
38	8	302	LUT	C8-C9	4.93	1.56	1.45
38	3	303	LUT	C8-C9	4.93	1.56	1.45
32	F	306	RRX	C23-C22	4.92	1.56	1.45
24	3	306	BCR	C15-C14	4.91	1.58	1.43
24	A	844	BCR	C15-C14	4.87	1.58	1.43
38	4	802	LUT	C8-C9	4.87	1.56	1.45
38	5	302	LUT	C8-C9	4.87	1.56	1.45
38	4	803	LUT	C8-C9	4.83	1.56	1.45
24	3	304	BCR	C15-C14	4.83	1.58	1.43
24	B	853	BCR	C19-C18	4.83	1.56	1.45
24	7	303	BCR	C15-C14	4.81	1.58	1.43
36	J	105	C7Z	C24-C25	-4.81	1.43	1.51
24	J	104	BCR	C15-C14	4.80	1.58	1.43
21	A	801	CL0	C3B-C2B	4.80	1.47	1.40
24	3	305	BCR	C15-C14	4.80	1.58	1.43
24	B	853	BCR	C29-C30	4.79	1.65	1.54
38	3	302	LUT	C8-C9	4.79	1.56	1.45
24	A	848	BCR	C15-C14	4.78	1.58	1.43
24	5	304	BCR	C15-C14	4.77	1.58	1.43
24	6	305	BCR	C15-C14	4.77	1.58	1.43
24	A	856	BCR	C15-C14	4.76	1.58	1.43
24	I	4001	BCR	C15-C14	4.76	1.58	1.43
24	8	304	BCR	C15-C14	4.76	1.58	1.43
24	B	849	BCR	C15-C14	4.75	1.58	1.43
24	3	307	BCR	C15-C14	4.74	1.58	1.43
24	F	303	BCR	C15-C14	4.74	1.58	1.43
24	B	848	BCR	C15-C14	4.74	1.58	1.43
24	4	804	BCR	C15-C14	4.73	1.58	1.43
24	G	203	BCR	C15-C14	4.73	1.58	1.43
24	7	303	BCR	C19-C18	4.73	1.56	1.45
24	B	845	BCR	C15-C14	4.73	1.58	1.43
24	B	846	BCR	C15-C14	4.72	1.58	1.43
24	5	305	BCR	C15-C14	4.72	1.58	1.43
24	5	305	BCR	C19-C18	4.72	1.56	1.45
24	A	845	BCR	C15-C14	4.72	1.58	1.43
24	A	847	BCR	C15-C14	4.72	1.58	1.43
21	A	801	CL0	CHD-C1D	4.71	1.47	1.38
38	Z	301	LUT	C12-C13	4.71	1.56	1.45
24	3	307	BCR	C19-C18	4.71	1.56	1.45
24	B	844	BCR	C15-C14	4.71	1.58	1.43
24	7	303	BCR	C2-C1	4.69	1.64	1.54

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
38	1	302	LUT	C12-C13	4.69	1.56	1.45
24	A	848	BCR	C19-C18	4.69	1.56	1.45
24	4	804	BCR	C2-C1	4.69	1.64	1.54
24	A	856	BCR	C19-C18	4.68	1.56	1.45
24	J	104	BCR	C2-C1	4.68	1.64	1.54
38	9	302	LUT	C4-C5	-4.68	1.43	1.51
24	K	206	BCR	C15-C14	4.68	1.57	1.43
24	L	204	BCR	C15-C14	4.67	1.57	1.43
24	B	849	BCR	C19-C18	4.67	1.56	1.45
38	Z	302	LUT	C12-C13	4.67	1.56	1.45
24	3	306	BCR	C2-C1	4.67	1.64	1.54
24	B	848	BCR	C2-C1	4.67	1.64	1.54
24	8	304	BCR	C2-C1	4.66	1.64	1.54
24	G	203	BCR	C19-C18	4.65	1.55	1.45
24	A	846	BCR	C15-C14	4.65	1.57	1.43
36	5	306	C7Z	C28-C29	4.65	1.55	1.45
38	4	802	LUT	C4-C5	-4.65	1.43	1.51
24	B	845	BCR	C2-C1	4.65	1.64	1.54
24	B	847	BCR	C15-C14	4.65	1.57	1.43
24	4	804	BCR	C19-C18	4.64	1.55	1.45
24	A	847	BCR	C2-C1	4.64	1.64	1.54
24	8	304	BCR	C19-C18	4.64	1.55	1.45
32	F	306	RRX	C24-C25	4.64	1.61	1.45
24	J	104	BCR	C19-C18	4.64	1.55	1.45
36	J	105	C7Z	C28-C29	4.64	1.55	1.45
24	A	846	BCR	C2-C1	4.63	1.64	1.54
24	F	303	BCR	C19-C18	4.63	1.55	1.45
24	6	305	BCR	C2-C1	4.63	1.64	1.54
24	L	204	BCR	C2-C1	4.63	1.64	1.54
38	7	302	LUT	C12-C13	4.63	1.55	1.45
24	3	305	BCR	C19-C18	4.63	1.55	1.45
24	3	307	BCR	C2-C1	4.62	1.64	1.54
24	6	306	BCR	C15-C14	4.62	1.57	1.43
38	6	303	LUT	C12-C13	4.62	1.55	1.45
38	1	303	LUT	C12-C13	4.62	1.55	1.45
24	3	306	BCR	C19-C18	4.61	1.55	1.45
24	3	304	BCR	C19-C18	4.61	1.55	1.45
24	3	304	BCR	C2-C1	4.61	1.64	1.54
24	A	844	BCR	C2-C1	4.61	1.64	1.54
24	A	848	BCR	C2-C1	4.60	1.64	1.54
24	6	306	BCR	C2-C1	4.59	1.64	1.54
24	K	206	BCR	C19-C18	4.59	1.55	1.45

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
24	A	844	BCR	C19-C18	4.59	1.55	1.45
38	5	303	LUT	C12-C13	4.59	1.55	1.45
24	B	848	BCR	C19-C18	4.59	1.55	1.45
24	B	845	BCR	C19-C18	4.59	1.55	1.45
38	3	303	LUT	C12-C13	4.59	1.55	1.45
38	8	303	LUT	C12-C13	4.58	1.55	1.45
24	B	847	BCR	C19-C18	4.58	1.55	1.45
24	B	853	BCR	C28-C27	4.58	1.66	1.52
38	7	301	LUT	C12-C13	4.58	1.55	1.45
24	B	853	BCR	C2-C1	4.58	1.64	1.54
21	A	801	CL0	C3C-C2C	4.57	1.46	1.36
38	7	302	LUT	C4-C5	-4.57	1.44	1.51
38	4	803	LUT	C12-C13	4.57	1.55	1.45
24	A	845	BCR	C19-C18	4.56	1.55	1.45
38	6	304	LUT	C12-C13	4.56	1.55	1.45
24	B	844	BCR	C19-C18	4.56	1.55	1.45
24	I	4001	BCR	C19-C18	4.55	1.55	1.45
24	I	4001	BCR	C2-C1	4.54	1.64	1.54
24	B	849	BCR	C2-C1	4.54	1.64	1.54
24	L	204	BCR	C19-C18	4.54	1.55	1.45
24	5	304	BCR	C19-C18	4.54	1.55	1.45
24	B	846	BCR	C19-C18	4.54	1.55	1.45
24	6	305	BCR	C19-C18	4.54	1.55	1.45
36	5	306	C7Z	C31-C30	4.54	1.57	1.43
24	5	304	BCR	C2-C1	4.53	1.64	1.54
38	5	302	LUT	C12-C13	4.52	1.55	1.45
24	A	845	BCR	C2-C1	4.52	1.64	1.54
38	4	803	LUT	C4-C5	-4.52	1.44	1.51
24	5	305	BCR	C2-C1	4.52	1.64	1.54
24	B	846	BCR	C2-C1	4.52	1.64	1.54
38	7	301	LUT	C4-C5	-4.52	1.44	1.51
38	9	302	LUT	C12-C13	4.51	1.55	1.45
24	K	206	BCR	C2-C1	4.51	1.64	1.54
38	5	303	LUT	C4-C5	-4.51	1.44	1.51
38	8	302	LUT	C12-C13	4.51	1.55	1.45
38	9	301	LUT	C12-C13	4.51	1.55	1.45
38	3	303	LUT	C4-C5	-4.51	1.44	1.51
38	4	802	LUT	C12-C13	4.50	1.55	1.45
38	6	304	LUT	C4-C5	-4.50	1.44	1.51
38	9	301	LUT	C4-C5	-4.50	1.44	1.51
38	8	303	LUT	C4-C5	-4.50	1.44	1.51
24	A	856	BCR	C2-C1	4.49	1.64	1.54

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
24	A	846	BCR	C19-C18	4.49	1.55	1.45
24	B	844	BCR	C2-C1	4.48	1.64	1.54
24	A	847	BCR	C19-C18	4.48	1.55	1.45
38	Z	302	LUT	C4-C5	-4.48	1.44	1.51
24	F	303	BCR	C2-C1	4.47	1.64	1.54
38	5	302	LUT	C4-C5	-4.46	1.44	1.51
32	F	306	RRX	C12-C13	4.46	1.55	1.45
24	B	847	BCR	C2-C1	4.45	1.64	1.54
38	8	302	LUT	C4-C5	-4.45	1.44	1.51
36	J	105	C7Z	C31-C30	4.44	1.57	1.43
38	Z	301	LUT	C31-C30	4.41	1.57	1.43
38	3	302	LUT	C4-C5	-4.40	1.44	1.51
38	Z	301	LUT	C15-C14	4.40	1.57	1.43
38	1	302	LUT	C15-C14	4.40	1.57	1.43
24	G	203	BCR	C2-C1	4.40	1.64	1.54
24	3	305	BCR	C2-C1	4.38	1.64	1.54
38	1	303	LUT	C4-C5	-4.38	1.44	1.51
38	Z	302	LUT	C31-C30	4.37	1.57	1.43
38	Z	302	LUT	C15-C14	4.37	1.57	1.43
38	1	303	LUT	C15-C14	4.37	1.57	1.43
24	8	304	BCR	C20-C21	4.36	1.57	1.43
24	A	856	BCR	C20-C21	4.36	1.57	1.43
38	Z	301	LUT	C4-C5	-4.36	1.44	1.51
38	7	302	LUT	C15-C14	4.36	1.56	1.43
22	5	320	CLA	MG-ND	-4.36	1.97	2.05
38	6	303	LUT	C4-C5	-4.35	1.44	1.51
38	5	303	LUT	C31-C30	4.35	1.56	1.43
38	3	302	LUT	C31-C30	4.35	1.56	1.43
24	7	303	BCR	C20-C21	4.34	1.56	1.43
38	5	303	LUT	C15-C14	4.34	1.56	1.43
38	1	302	LUT	C35-C34	4.34	1.56	1.43
24	A	848	BCR	C20-C21	4.33	1.56	1.43
38	3	303	LUT	C15-C14	4.33	1.56	1.43
38	3	303	LUT	C31-C30	4.33	1.56	1.43
38	7	301	LUT	C15-C14	4.32	1.56	1.43
38	6	304	LUT	C15-C14	4.32	1.56	1.43
38	9	302	LUT	C31-C30	4.32	1.56	1.43
32	F	306	RRX	C3-C4	4.31	1.66	1.52
38	1	302	LUT	C31-C30	4.31	1.56	1.43
31	B	852	DGD	O1G-C1A	4.31	1.45	1.33
38	8	303	LUT	C31-C30	4.31	1.56	1.43
22	K	205	CLA	MG-ND	-4.31	1.97	2.05

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
24	B	847	BCR	C28-C27	4.30	1.66	1.52
24	3	307	BCR	C20-C21	4.30	1.56	1.43
24	3	304	BCR	C20-C21	4.30	1.56	1.43
38	1	303	LUT	C31-C30	4.30	1.56	1.43
38	8	303	LUT	C15-C14	4.30	1.56	1.43
24	B	849	BCR	C20-C21	4.30	1.56	1.43
38	7	301	LUT	C31-C30	4.30	1.56	1.43
24	A	856	BCR	C28-C27	4.30	1.66	1.52
24	4	804	BCR	C20-C21	4.30	1.56	1.43
24	A	844	BCR	C28-C27	4.29	1.66	1.52
38	Z	301	LUT	C35-C34	4.29	1.56	1.43
24	J	104	BCR	C28-C27	4.29	1.66	1.52
38	7	302	LUT	C31-C30	4.29	1.56	1.43
38	6	303	LUT	C15-C14	4.29	1.56	1.43
24	3	305	BCR	C20-C21	4.29	1.56	1.43
36	5	306	C7Z	C32-C33	4.29	1.55	1.45
38	6	304	LUT	C31-C30	4.29	1.56	1.43
24	B	853	BCR	C20-C21	4.29	1.56	1.43
24	F	303	BCR	C20-C21	4.29	1.56	1.43
24	5	304	BCR	C20-C21	4.28	1.56	1.43
31	3	301	DGD	O1G-C1A	4.28	1.45	1.33
36	J	105	C7Z	C32-C33	4.28	1.55	1.45
38	3	302	LUT	C12-C13	4.28	1.55	1.45
24	3	306	BCR	C20-C21	4.28	1.56	1.43
38	9	301	LUT	C15-C14	4.27	1.56	1.43
38	8	302	LUT	C15-C14	4.27	1.56	1.43
24	B	845	BCR	C28-C27	4.27	1.65	1.52
38	6	303	LUT	C31-C30	4.27	1.56	1.43
24	L	204	BCR	C28-C27	4.27	1.65	1.52
24	I	4001	BCR	C28-C27	4.27	1.65	1.52
24	5	304	BCR	C28-C27	4.26	1.65	1.52
38	9	301	LUT	C31-C30	4.26	1.56	1.43
24	K	206	BCR	C20-C21	4.26	1.56	1.43
24	J	104	BCR	C20-C21	4.26	1.56	1.43
38	4	803	LUT	C15-C14	4.26	1.56	1.43
38	9	302	LUT	C15-C14	4.26	1.56	1.43
24	A	848	BCR	C28-C27	4.26	1.65	1.52
38	1	303	LUT	C35-C34	4.26	1.56	1.43
24	B	849	BCR	C28-C27	4.25	1.65	1.52
24	B	847	BCR	C20-C21	4.25	1.56	1.43
38	4	802	LUT	C31-C30	4.25	1.56	1.43
24	B	844	BCR	C20-C21	4.25	1.56	1.43

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
24	I	4001	BCR	C20-C21	4.25	1.56	1.43
24	3	305	BCR	C28-C27	4.25	1.65	1.52
38	5	302	LUT	C15-C14	4.25	1.56	1.43
38	Z	302	LUT	C35-C34	4.24	1.56	1.43
38	5	303	LUT	C35-C34	4.24	1.56	1.43
21	A	801	CL0	C3D-C4D	-4.24	1.34	1.44
24	6	306	BCR	C19-C18	4.24	1.55	1.45
32	F	306	RRX	C11-C10	4.24	1.56	1.43
38	4	802	LUT	C15-C14	4.24	1.56	1.43
38	8	302	LUT	C31-C30	4.24	1.56	1.43
38	7	301	LUT	C35-C34	4.23	1.56	1.43
38	7	302	LUT	C35-C34	4.23	1.56	1.43
24	A	845	BCR	C20-C21	4.23	1.56	1.43
24	6	305	BCR	C20-C21	4.23	1.56	1.43
24	3	306	BCR	C28-C27	4.23	1.65	1.52
24	L	204	BCR	C20-C21	4.23	1.56	1.43
38	6	304	LUT	C35-C34	4.23	1.56	1.43
38	6	303	LUT	C35-C34	4.23	1.56	1.43
24	A	845	BCR	C28-C27	4.23	1.65	1.52
24	K	206	BCR	C28-C27	4.22	1.65	1.52
24	B	845	BCR	C20-C21	4.22	1.56	1.43
24	A	844	BCR	C20-C21	4.22	1.56	1.43
24	G	203	BCR	C20-C21	4.22	1.56	1.43
38	4	803	LUT	C31-C30	4.21	1.56	1.43
24	B	846	BCR	C20-C21	4.21	1.56	1.43
24	A	847	BCR	C20-C21	4.21	1.56	1.43
24	B	848	BCR	C20-C21	4.21	1.56	1.43
24	A	847	BCR	C28-C27	4.20	1.65	1.52
24	5	305	BCR	C20-C21	4.20	1.56	1.43
38	8	302	LUT	C35-C34	4.19	1.56	1.43
38	8	303	LUT	C35-C34	4.19	1.56	1.43
38	5	302	LUT	C31-C30	4.19	1.56	1.43
38	Z	301	LUT	C28-C29	4.19	1.54	1.45
22	K	205	CLA	C1C-NC	-4.19	1.31	1.37
24	B	844	BCR	C28-C27	4.19	1.65	1.52
24	A	846	BCR	C20-C21	4.19	1.56	1.43
38	4	803	LUT	C35-C34	4.19	1.56	1.43
38	9	301	LUT	C35-C34	4.19	1.56	1.43
38	3	303	LUT	C35-C34	4.18	1.56	1.43
38	9	302	LUT	C35-C34	4.18	1.56	1.43
24	5	305	BCR	C28-C27	4.18	1.65	1.52
24	7	303	BCR	C28-C27	4.16	1.65	1.52

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
24	A	846	BCR	C28-C27	4.16	1.65	1.52
24	4	804	BCR	C28-C27	4.16	1.65	1.52
38	4	802	LUT	C35-C34	4.15	1.56	1.43
31	1	319	DGD	O1G-C1A	4.15	1.45	1.33
22	1	316	CLA	MG-ND	-4.14	1.97	2.05
24	3	307	BCR	C28-C27	4.14	1.65	1.52
24	3	304	BCR	C28-C27	4.14	1.65	1.52
32	F	306	RRX	C29-C30	4.14	1.67	1.54
24	B	846	BCR	C28-C27	4.13	1.65	1.52
38	3	302	LUT	C15-C14	4.13	1.56	1.43
38	3	302	LUT	C28-C29	4.13	1.54	1.45
24	F	303	BCR	C28-C27	4.13	1.65	1.52
38	1	302	LUT	C28-C29	4.13	1.54	1.45
24	B	848	BCR	C28-C27	4.11	1.65	1.52
24	6	306	BCR	C28-C27	4.10	1.65	1.52
38	5	302	LUT	C35-C34	4.10	1.56	1.43
32	F	306	RRX	C15-C14	4.10	1.56	1.43
24	G	203	BCR	C28-C27	4.09	1.65	1.52
24	8	304	BCR	C28-C27	4.08	1.65	1.52
38	3	302	LUT	C35-C34	4.08	1.56	1.43
24	6	305	BCR	C28-C27	4.07	1.65	1.52
25	3	321	LHG	O7-C5	-4.07	1.42	1.46
24	6	306	BCR	C20-C21	4.07	1.56	1.43
38	Z	301	LUT	C7-C6	4.07	1.59	1.45
38	Z	302	LUT	C28-C29	4.06	1.54	1.45
38	6	303	LUT	C7-C6	4.06	1.59	1.45
38	5	303	LUT	C28-C29	4.04	1.54	1.45
38	1	302	LUT	C7-C6	4.04	1.59	1.45
38	8	302	LUT	C28-C29	4.04	1.54	1.45
32	F	306	RRX	C16-C17	4.03	1.55	1.43
38	1	303	LUT	C28-C29	4.03	1.54	1.45
21	A	801	CL0	CHD-C4C	4.03	1.48	1.39
38	3	303	LUT	C28-C29	4.03	1.54	1.45
38	9	302	LUT	C28-C29	4.02	1.54	1.45
38	6	303	LUT	C28-C29	4.02	1.54	1.45
38	7	302	LUT	C7-C6	4.02	1.59	1.45
38	8	303	LUT	C28-C29	4.02	1.54	1.45
38	Z	302	LUT	C7-C6	4.02	1.59	1.45
22	9	309	CLA	MG-ND	-4.02	1.97	2.05
38	6	304	LUT	C28-C29	4.02	1.54	1.45
38	7	301	LUT	C28-C29	4.01	1.54	1.45
38	5	303	LUT	C7-C6	4.01	1.59	1.45

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
38	8	302	LUT	C21-C26	4.01	1.67	1.56
38	6	304	LUT	C7-C6	4.01	1.59	1.45
38	9	301	LUT	C28-C29	4.00	1.54	1.45
38	1	302	LUT	C4-C5	-4.00	1.44	1.51
38	9	301	LUT	C7-C6	4.00	1.59	1.45
22	3	315	CLA	MG-ND	-3.99	1.97	2.05
38	5	302	LUT	C7-C6	3.99	1.59	1.45
38	9	302	LUT	C7-C6	3.99	1.59	1.45
38	8	302	LUT	C7-C6	3.99	1.59	1.45
38	3	303	LUT	C7-C6	3.99	1.59	1.45
38	1	303	LUT	C7-C6	3.98	1.59	1.45
38	5	302	LUT	C28-C29	3.98	1.54	1.45
24	8	304	BCR	C3-C4	3.98	1.65	1.52
38	8	303	LUT	C7-C6	3.98	1.59	1.45
24	3	307	BCR	C3-C4	3.98	1.64	1.52
38	7	301	LUT	C7-C6	3.97	1.59	1.45
24	A	845	BCR	C3-C4	3.97	1.64	1.52
24	L	204	BCR	C3-C4	3.96	1.64	1.52
38	4	803	LUT	C7-C6	3.96	1.59	1.45
24	J	104	BCR	C3-C4	3.96	1.64	1.52
24	7	303	BCR	C3-C4	3.96	1.64	1.52
38	4	803	LUT	C28-C29	3.96	1.54	1.45
38	4	802	LUT	C7-C6	3.95	1.59	1.45
32	F	306	RRX	C20-C21	3.95	1.55	1.43
38	7	302	LUT	C28-C29	3.95	1.54	1.45
22	9	303	CLA	MG-ND	-3.95	1.98	2.05
24	A	848	BCR	C3-C4	3.95	1.64	1.52
38	4	802	LUT	C28-C29	3.94	1.54	1.45
22	9	304	CLA	MG-ND	-3.94	1.98	2.05
24	3	306	BCR	C11-C10	3.93	1.55	1.43
38	3	302	LUT	C7-C6	3.93	1.59	1.45
36	5	306	C7Z	C22-C21	3.93	1.67	1.54
24	I	4001	BCR	C3-C4	3.93	1.64	1.52
24	B	845	BCR	C3-C4	3.93	1.64	1.52
38	Z	301	LUT	C21-C26	3.93	1.67	1.56
38	9	302	LUT	C21-C26	3.92	1.67	1.56
38	9	301	LUT	C21-C26	3.92	1.67	1.56
24	A	856	BCR	C3-C4	3.92	1.64	1.52
38	Z	302	LUT	C21-C26	3.91	1.67	1.56
24	5	305	BCR	C11-C10	3.90	1.55	1.43
24	6	306	BCR	C3-C4	3.90	1.64	1.52
38	8	303	LUT	C21-C26	3.89	1.67	1.56

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
24	B	853	BCR	C3-C4	3.89	1.64	1.52
24	B	846	BCR	C3-C4	3.89	1.64	1.52
38	3	303	LUT	C21-C26	3.89	1.67	1.56
24	3	305	BCR	C3-C4	3.88	1.64	1.52
22	9	305	CLA	MG-ND	-3.88	1.98	2.05
24	G	203	BCR	C3-C4	3.88	1.64	1.52
22	1	316	CLA	C1C-NC	-3.88	1.32	1.37
22	9	310	CLA	MG-ND	-3.88	1.98	2.05
36	5	306	C7Z	C4-C5	-3.88	1.45	1.51
24	B	853	BCR	C16-C17	3.88	1.55	1.43
38	6	304	LUT	C21-C26	3.87	1.67	1.56
24	B	847	BCR	C3-C4	3.87	1.64	1.52
24	F	303	BCR	C3-C4	3.87	1.64	1.52
38	4	803	LUT	C21-C26	3.87	1.67	1.56
24	3	306	BCR	C3-C4	3.86	1.64	1.52
38	1	302	LUT	C21-C26	3.86	1.67	1.56
36	J	105	C7Z	C11-C10	3.86	1.55	1.43
36	5	306	C7Z	C15-C14	3.86	1.55	1.43
38	7	302	LUT	C21-C26	3.86	1.67	1.56
24	3	306	BCR	C16-C17	3.86	1.55	1.43
24	5	305	BCR	C3-C4	3.85	1.64	1.52
22	3	320	CLA	C1C-NC	-3.85	1.32	1.37
22	5	320	CLA	C1C-NC	-3.85	1.32	1.37
22	3	320	CLA	MG-ND	-3.85	1.98	2.05
36	J	105	C7Z	C15-C14	3.85	1.55	1.43
24	A	844	BCR	C16-C17	3.84	1.55	1.43
38	5	302	LUT	C21-C26	3.84	1.67	1.56
24	A	847	BCR	C3-C4	3.84	1.64	1.52
24	5	304	BCR	C3-C4	3.84	1.64	1.52
24	B	853	BCR	C11-C10	3.83	1.55	1.43
24	B	844	BCR	C3-C4	3.83	1.64	1.52
24	K	206	BCR	C3-C4	3.83	1.64	1.52
22	9	303	CLA	C1C-NC	-3.82	1.32	1.37
38	7	301	LUT	C21-C26	3.82	1.67	1.56
24	L	204	BCR	C11-C10	3.82	1.55	1.43
36	J	105	C7Z	C22-C21	3.81	1.66	1.54
24	3	304	BCR	C16-C17	3.81	1.55	1.43
24	B	848	BCR	C3-C4	3.81	1.64	1.52
24	B	849	BCR	C3-C4	3.80	1.64	1.52
38	Z	301	LUT	C11-C10	3.80	1.55	1.43
24	G	203	BCR	C11-C10	3.80	1.55	1.43
24	A	844	BCR	C11-C10	3.80	1.55	1.43

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
24	7	303	BCR	C16-C17	3.79	1.55	1.43
36	5	306	C7Z	C11-C10	3.79	1.55	1.43
38	3	302	LUT	C21-C26	3.79	1.66	1.56
38	1	303	LUT	C21-C26	3.79	1.66	1.56
24	A	846	BCR	C3-C4	3.79	1.64	1.52
24	A	848	BCR	C16-C17	3.79	1.55	1.43
32	F	306	RRX	C27-C26	-3.78	1.45	1.51
38	6	303	LUT	C11-C10	3.78	1.55	1.43
24	6	305	BCR	C3-C4	3.78	1.64	1.52
38	4	802	LUT	C21-C26	3.78	1.66	1.56
38	5	303	LUT	C21-C26	3.77	1.66	1.56
24	A	856	BCR	C16-C17	3.77	1.55	1.43
24	3	305	BCR	C16-C17	3.77	1.55	1.43
24	B	849	BCR	C11-C10	3.77	1.55	1.43
24	B	849	BCR	C16-C17	3.77	1.55	1.43
38	9	302	LUT	C38-C25	3.76	1.57	1.50
38	Z	302	LUT	C11-C10	3.76	1.55	1.43
24	J	104	BCR	C16-C17	3.76	1.55	1.43
24	I	4001	BCR	C16-C17	3.76	1.55	1.43
24	4	804	BCR	C3-C4	3.75	1.64	1.52
36	5	306	C7Z	C27-C26	3.75	1.58	1.45
24	3	307	BCR	C16-C17	3.75	1.55	1.43
24	F	303	BCR	C16-C17	3.75	1.55	1.43
24	5	305	BCR	C16-C17	3.74	1.55	1.43
38	1	303	LUT	C38-C25	3.74	1.57	1.50
38	1	302	LUT	C11-C10	3.74	1.55	1.43
38	6	303	LUT	C21-C26	3.74	1.66	1.56
22	3	315	CLA	C1C-NC	-3.74	1.32	1.37
38	7	301	LUT	C11-C10	3.74	1.55	1.43
24	A	845	BCR	C16-C17	3.73	1.55	1.43
24	A	856	BCR	C11-C10	3.73	1.55	1.43
24	G	203	BCR	C16-C17	3.73	1.55	1.43
24	4	804	BCR	C16-C17	3.73	1.55	1.43
24	B	845	BCR	C16-C17	3.73	1.55	1.43
38	7	302	LUT	C11-C10	3.73	1.55	1.43
24	B	846	BCR	C16-C17	3.73	1.55	1.43
24	A	844	BCR	C3-C4	3.73	1.64	1.52
38	1	303	LUT	C11-C10	3.73	1.55	1.43
24	F	303	BCR	C11-C10	3.72	1.55	1.43
38	5	303	LUT	C11-C10	3.72	1.55	1.43
38	8	303	LUT	C11-C10	3.72	1.55	1.43
36	J	105	C7Z	C4-C5	-3.72	1.45	1.51

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
24	B	844	BCR	C11-C10	3.72	1.55	1.43
24	A	848	BCR	C11-C10	3.72	1.55	1.43
21	A	801	CL0	OBD-CAD	3.72	1.28	1.22
38	9	301	LUT	C38-C25	3.72	1.57	1.50
24	K	206	BCR	C16-C17	3.72	1.55	1.43
24	6	305	BCR	C16-C17	3.71	1.55	1.43
24	J	104	BCR	C11-C10	3.71	1.55	1.43
36	J	105	C7Z	C27-C26	3.71	1.58	1.45
38	8	303	LUT	C38-C25	3.71	1.57	1.50
38	3	303	LUT	C11-C10	3.71	1.54	1.43
38	6	304	LUT	C11-C10	3.71	1.54	1.43
24	3	304	BCR	C3-C4	3.71	1.64	1.52
24	7	303	BCR	C11-C10	3.71	1.54	1.43
24	B	845	BCR	C11-C10	3.70	1.54	1.43
38	9	301	LUT	C11-C10	3.70	1.54	1.43
38	6	304	LUT	C38-C25	3.70	1.57	1.50
24	5	304	BCR	C16-C17	3.70	1.54	1.43
24	8	304	BCR	C16-C17	3.70	1.54	1.43
24	4	804	BCR	C11-C10	3.70	1.54	1.43
24	B	846	BCR	C11-C10	3.70	1.54	1.43
24	3	305	BCR	C11-C10	3.69	1.54	1.43
24	B	844	BCR	C16-C17	3.69	1.54	1.43
22	5	313	CLA	MG-ND	-3.68	1.98	2.05
38	1	302	LUT	C19-C9	3.68	1.58	1.50
24	I	4001	BCR	C11-C10	3.68	1.54	1.43
24	L	204	BCR	C16-C17	3.68	1.54	1.43
24	A	847	BCR	C16-C17	3.68	1.54	1.43
22	4	806	CLA	MG-ND	-3.68	1.98	2.05
38	7	302	LUT	C38-C25	3.68	1.57	1.50
38	4	803	LUT	C38-C25	3.68	1.57	1.50
22	G	201	CLA	MG-ND	-3.68	1.98	2.05
24	3	304	BCR	C11-C10	3.68	1.54	1.43
22	Z	305	CLA	MG-ND	-3.68	1.98	2.05
32	F	306	RRX	C7-C6	3.67	1.58	1.45
22	7	306	CLA	MG-ND	-3.67	1.98	2.05
38	5	302	LUT	C11-C10	3.67	1.54	1.43
22	A	820	CLA	MG-ND	-3.67	1.98	2.05
22	B	839	CLA	MG-ND	-3.67	1.98	2.05
38	4	802	LUT	C11-C10	3.67	1.54	1.43
24	B	848	BCR	C16-C17	3.67	1.54	1.43
22	A	816	CLA	MG-ND	-3.67	1.98	2.05
22	Z	313	CLA	MG-ND	-3.67	1.98	2.05

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
24	B	847	BCR	C16-C17	3.67	1.54	1.43
38	8	302	LUT	C11-C10	3.67	1.54	1.43
36	J	105	C7Z	C35-C34	3.67	1.54	1.43
22	6	312	CLA	MG-ND	-3.66	1.98	2.05
22	K	204	CLA	MG-ND	-3.66	1.98	2.05
22	1	305	CLA	MG-ND	-3.66	1.98	2.05
24	5	304	BCR	C11-C10	3.66	1.54	1.43
22	1	311	CLA	MG-ND	-3.66	1.98	2.05
22	5	308	CLA	MG-ND	-3.66	1.98	2.05
36	5	306	C7Z	C35-C34	3.66	1.54	1.43
24	A	845	BCR	C11-C10	3.66	1.54	1.43
38	4	803	LUT	C11-C10	3.65	1.54	1.43
22	6	309	CLA	MG-ND	-3.65	1.98	2.05
22	8	309	CLA	MG-ND	-3.65	1.98	2.05
38	8	302	LUT	C38-C25	3.65	1.57	1.50
38	3	302	LUT	C19-C9	3.65	1.58	1.50
22	A	839	CLA	MG-ND	-3.65	1.98	2.05
22	6	311	CLA	MG-ND	-3.65	1.98	2.05
24	A	846	BCR	C11-C10	3.64	1.54	1.43
22	1	311	CLA	C1C-NC	-3.64	1.32	1.37
22	9	308	CLA	MG-ND	-3.64	1.98	2.05
38	Z	302	LUT	C19-C9	3.64	1.58	1.50
24	B	847	BCR	C11-C10	3.64	1.54	1.43
22	4	818	CLA	MG-ND	-3.64	1.98	2.05
22	B	815	CLA	MG-ND	-3.63	1.98	2.05
22	5	309	CLA	MG-ND	-3.63	1.98	2.05
22	4	807	CLA	MG-ND	-3.63	1.98	2.05
22	B	817	CLA	MG-ND	-3.63	1.98	2.05
24	6	305	BCR	C11-C10	3.63	1.54	1.43
22	A	836	CLA	MG-ND	-3.63	1.98	2.05
22	5	311	CLA	MG-ND	-3.63	1.98	2.05
22	K	203	CLA	MG-ND	-3.63	1.98	2.05
22	7	308	CLA	MG-ND	-3.63	1.98	2.05
24	K	206	BCR	C11-C10	3.63	1.54	1.43
38	5	303	LUT	C38-C25	3.63	1.57	1.50
38	9	302	LUT	C11-C10	3.63	1.54	1.43
38	Z	302	LUT	C38-C25	3.63	1.57	1.50
24	B	848	BCR	C11-C10	3.63	1.54	1.43
22	6	302	CLA	MG-ND	-3.63	1.98	2.05
22	6	322	CLA	MG-ND	-3.63	1.98	2.05
24	A	856	BCR	C24-C25	3.63	1.58	1.45
22	1	304	CLA	MG-ND	-3.62	1.98	2.05

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
24	3	307	BCR	C11-C10	3.62	1.54	1.43
22	A	802	CLA	MG-ND	-3.62	1.98	2.05
22	A	842	CLA	MG-ND	-3.62	1.98	2.05
22	4	817	CLA	MG-ND	-3.62	1.98	2.05
22	B	834	CLA	MG-ND	-3.62	1.98	2.05
22	1	313	CLA	MG-ND	-3.62	1.98	2.05
22	4	808	CLA	MG-ND	-3.62	1.98	2.05
22	A	835	CLA	MG-ND	-3.62	1.98	2.05
22	8	306	CLA	MG-ND	-3.62	1.98	2.05
22	9	311	CLA	MG-ND	-3.62	1.98	2.05
38	5	302	LUT	C38-C25	3.62	1.57	1.50
38	6	303	LUT	C19-C9	3.62	1.58	1.50
22	4	809	CLA	MG-ND	-3.61	1.98	2.05
38	5	303	LUT	C19-C9	3.61	1.58	1.50
24	6	306	BCR	C11-C10	3.61	1.54	1.43
38	1	303	LUT	C19-C9	3.61	1.58	1.50
22	A	829	CLA	MG-ND	-3.61	1.98	2.05
22	K	202	CLA	MG-ND	-3.61	1.98	2.05
22	3	309	CLA	MG-ND	-3.61	1.98	2.05
38	9	301	LUT	C19-C9	3.61	1.58	1.50
24	8	304	BCR	C11-C10	3.61	1.54	1.43
22	A	804	CLA	MG-ND	-3.61	1.98	2.05
22	7	309	CLA	MG-ND	-3.61	1.98	2.05
22	6	319	CLA	MG-ND	-3.61	1.98	2.05
22	B	821	CLA	MG-ND	-3.61	1.98	2.05
22	B	822	CLA	MG-ND	-3.61	1.98	2.05
38	6	303	LUT	C38-C25	3.61	1.57	1.50
24	7	303	BCR	C24-C25	3.61	1.57	1.45
22	7	305	CLA	MG-ND	-3.61	1.98	2.05
24	A	847	BCR	C11-C10	3.61	1.54	1.43
22	5	315	CLA	MG-ND	-3.60	1.98	2.05
22	A	832	CLA	MG-ND	-3.60	1.98	2.05
22	B	806	CLA	MG-ND	-3.60	1.98	2.05
22	F	304	CLA	MG-ND	-3.60	1.98	2.05
22	1	315	CLA	MG-ND	-3.60	1.98	2.05
22	5	314	CLA	MG-ND	-3.60	1.98	2.05
24	A	846	BCR	C16-C17	3.60	1.54	1.43
24	B	844	BCR	C24-C25	3.60	1.57	1.45
22	6	313	CLA	MG-ND	-3.60	1.98	2.05
22	Z	304	CLA	MG-ND	-3.60	1.98	2.05
22	B	841	CLA	MG-ND	-3.60	1.98	2.05
22	9	305	CLA	C1C-NC	-3.60	1.32	1.37

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
22	3	310	CLA	MG-ND	-3.60	1.98	2.05
22	B	836	CLA	MG-ND	-3.60	1.98	2.05
24	A	845	BCR	C24-C25	3.60	1.57	1.45
38	4	802	LUT	C19-C9	3.60	1.58	1.50
22	B	807	CLA	MG-ND	-3.60	1.98	2.05
22	4	812	CLA	MG-ND	-3.60	1.98	2.05
22	L	203	CLA	MG-ND	-3.60	1.98	2.05
22	5	326	CLA	MG-ND	-3.60	1.98	2.05
22	8	313	CLA	MG-ND	-3.59	1.98	2.05
22	B	801	CLA	MG-ND	-3.59	1.98	2.05
24	A	848	BCR	C24-C25	3.59	1.57	1.45
22	A	831	CLA	MG-ND	-3.59	1.98	2.05
22	6	310	CLA	MG-ND	-3.59	1.98	2.05
22	2	302	CLA	MG-ND	-3.59	1.98	2.05
38	3	302	LUT	C38-C25	3.59	1.57	1.50
22	A	838	CLA	MG-ND	-3.59	1.98	2.05
38	6	304	LUT	C19-C9	3.59	1.58	1.50
22	A	814	CLA	MG-ND	-3.59	1.98	2.05
22	A	854	CLA	MG-ND	-3.59	1.98	2.05
22	5	310	CLA	MG-ND	-3.59	1.98	2.05
22	B	813	CLA	MG-ND	-3.59	1.98	2.05
22	2	301	CLA	MG-ND	-3.59	1.98	2.05
24	3	307	BCR	C24-C25	3.59	1.57	1.45
24	6	306	BCR	C16-C17	3.59	1.54	1.43
24	3	306	BCR	C24-C25	3.59	1.57	1.45
24	3	305	BCR	C24-C25	3.59	1.57	1.45
22	B	808	CLA	MG-ND	-3.59	1.98	2.05
22	3	308	CLA	MG-ND	-3.59	1.98	2.05
22	7	322	CLA	MG-ND	-3.59	1.98	2.05
22	8	305	CLA	MG-ND	-3.59	1.98	2.05
22	B	809	CLA	MG-ND	-3.59	1.98	2.05
22	3	322	CLA	MG-ND	-3.59	1.98	2.05
22	Z	307	CLA	MG-ND	-3.59	1.98	2.05
22	6	307	CLA	MG-ND	-3.59	1.98	2.05
38	4	803	LUT	C19-C9	3.58	1.58	1.50
24	J	104	BCR	C24-C25	3.58	1.57	1.45
24	8	304	BCR	C24-C25	3.58	1.57	1.45
22	J	103	CLA	MG-ND	-3.58	1.98	2.05
22	8	314	CLA	MG-ND	-3.58	1.98	2.05
22	5	312	CLA	MG-ND	-3.58	1.98	2.05
22	B	837	CLA	MG-ND	-3.58	1.98	2.05
22	3	312	CLA	MG-ND	-3.58	1.98	2.05

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
22	9	313	CLA	MG-ND	-3.58	1.98	2.05
22	3	314	CLA	MG-ND	-3.58	1.98	2.05
38	3	303	LUT	C38-C25	3.58	1.57	1.50
38	7	302	LUT	C19-C9	3.58	1.58	1.50
22	7	314	CLA	MG-ND	-3.58	1.98	2.05
22	Z	315	CLA	MG-ND	-3.58	1.98	2.05
22	8	310	CLA	MG-ND	-3.58	1.98	2.05
22	6	317	CLA	MG-ND	-3.58	1.98	2.05
22	3	319	CLA	MG-ND	-3.58	1.98	2.05
38	5	302	LUT	C19-C9	3.58	1.58	1.50
22	B	826	CLA	MG-ND	-3.58	1.98	2.05
22	A	817	CLA	MG-ND	-3.57	1.98	2.05
22	B	832	CLA	MG-ND	-3.57	1.98	2.05
38	3	303	LUT	C19-C9	3.57	1.58	1.50
22	A	813	CLA	MG-ND	-3.57	1.98	2.05
22	A	821	CLA	MG-ND	-3.57	1.98	2.05
22	A	827	CLA	MG-ND	-3.57	1.98	2.05
22	7	323	CLA	MG-ND	-3.57	1.98	2.05
22	B	833	CLA	MG-ND	-3.57	1.98	2.05
22	1	306	CLA	MG-ND	-3.57	1.98	2.05
22	Z	310	CLA	MG-ND	-3.57	1.98	2.05
38	7	301	LUT	C19-C9	3.57	1.58	1.50
22	7	324	CLA	MG-ND	-3.57	1.98	2.05
38	3	302	LUT	C11-C10	3.57	1.54	1.43
38	9	302	LUT	C19-C9	3.57	1.58	1.50
22	6	308	CLA	MG-ND	-3.57	1.98	2.05
22	4	805	CLA	MG-ND	-3.57	1.98	2.05
24	G	203	BCR	C24-C25	3.57	1.57	1.45
22	1	307	CLA	MG-ND	-3.57	1.98	2.05
38	Z	301	LUT	C19-C9	3.57	1.58	1.50
22	B	828	CLA	MG-ND	-3.57	1.98	2.05
22	9	306	CLA	MG-ND	-3.57	1.98	2.05
22	6	321	CLA	MG-ND	-3.57	1.98	2.05
22	B	823	CLA	MG-ND	-3.57	1.98	2.05
22	7	307	CLA	MG-ND	-3.56	1.98	2.05
22	A	809	CLA	MG-ND	-3.56	1.98	2.05
22	L	202	CLA	MG-ND	-3.56	1.98	2.05
22	3	316	CLA	MG-ND	-3.56	1.98	2.05
22	8	307	CLA	MG-ND	-3.56	1.98	2.05
22	B	824	CLA	MG-ND	-3.56	1.98	2.05
22	Z	303	CLA	MG-ND	-3.56	1.98	2.05
22	A	828	CLA	MG-ND	-3.56	1.98	2.05

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
22	A	840	CLA	MG-ND	-3.56	1.98	2.05
22	4	811	CLA	MG-ND	-3.56	1.98	2.05
22	B	829	CLA	MG-ND	-3.56	1.98	2.05
22	3	311	CLA	MG-ND	-3.56	1.98	2.05
22	B	831	CLA	MG-ND	-3.56	1.98	2.05
22	7	317	CLA	MG-ND	-3.56	1.98	2.05
22	5	307	CLA	MG-ND	-3.56	1.98	2.05
22	B	825	CLA	MG-ND	-3.56	1.98	2.05
24	5	304	BCR	C24-C25	3.56	1.57	1.45
22	B	835	CLA	MG-ND	-3.56	1.98	2.05
22	B	816	CLA	MG-ND	-3.56	1.98	2.05
22	Z	306	CLA	MG-ND	-3.56	1.98	2.05
22	B	830	CLA	MG-ND	-3.56	1.98	2.05
38	8	302	LUT	C19-C9	3.56	1.58	1.50
22	B	811	CLA	MG-ND	-3.56	1.98	2.05
22	7	310	CLA	MG-ND	-3.56	1.98	2.05
22	9	307	CLA	MG-ND	-3.56	1.98	2.05
22	F	305	CLA	MG-ND	-3.55	1.98	2.05
22	7	304	CLA	MG-ND	-3.55	1.98	2.05
22	7	316	CLA	MG-ND	-3.55	1.98	2.05
22	2	304	CLA	MG-ND	-3.55	1.98	2.05
24	3	304	BCR	C24-C25	3.55	1.57	1.45
22	A	826	CLA	MG-ND	-3.55	1.98	2.05
22	B	840	CLA	MG-ND	-3.55	1.98	2.05
22	A	815	CLA	MG-ND	-3.55	1.98	2.05
22	5	301	CLA	MG-ND	-3.55	1.98	2.05
22	A	822	CLA	MG-ND	-3.55	1.98	2.05
22	A	825	CLA	MG-ND	-3.55	1.98	2.05
22	1	310	CLA	MG-ND	-3.55	1.98	2.05
22	A	812	CLA	MG-ND	-3.55	1.98	2.05
22	8	311	CLA	MG-ND	-3.55	1.98	2.05
24	5	305	BCR	C24-C25	3.55	1.57	1.45
24	B	849	BCR	C24-C25	3.55	1.57	1.45
24	4	804	BCR	C24-C25	3.55	1.57	1.45
22	5	322	CLA	MG-ND	-3.54	1.98	2.05
24	L	204	BCR	C24-C25	3.54	1.57	1.45
22	A	830	CLA	MG-ND	-3.54	1.98	2.05
22	B	810	CLA	MG-ND	-3.54	1.98	2.05
24	A	846	BCR	C24-C25	3.54	1.57	1.45
38	1	302	LUT	C38-C25	3.54	1.57	1.50
22	A	818	CLA	MG-ND	-3.54	1.98	2.05
22	5	323	CLA	MG-ND	-3.54	1.98	2.05

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
22	7	312	CLA	MG-ND	-3.54	1.98	2.05
22	7	315	CLA	MG-ND	-3.54	1.98	2.05
24	B	845	BCR	C24-C25	3.54	1.57	1.45
22	A	803	CLA	MG-ND	-3.54	1.98	2.05
22	6	301	CLA	MG-ND	-3.54	1.98	2.05
24	K	206	BCR	C24-C25	3.54	1.57	1.45
22	B	820	CLA	MG-ND	-3.54	1.98	2.05
22	A	806	CLA	MG-ND	-3.54	1.98	2.05
22	Z	309	CLA	MG-ND	-3.54	1.98	2.05
22	8	308	CLA	MG-ND	-3.53	1.98	2.05
22	Z	308	CLA	MG-ND	-3.53	1.98	2.05
22	9	310	CLA	C1C-NC	-3.53	1.32	1.37
36	J	105	C7Z	C2-C1	3.53	1.65	1.54
24	6	305	BCR	C4-C5	-3.53	1.44	1.51
22	4	815	CLA	MG-ND	-3.53	1.98	2.05
22	B	812	CLA	MG-ND	-3.53	1.98	2.05
22	G	202	CLA	MG-ND	-3.53	1.98	2.05
24	A	844	BCR	C24-C25	3.53	1.57	1.45
22	B	819	CLA	MG-ND	-3.53	1.98	2.05
22	F	302	CLA	MG-ND	-3.53	1.98	2.05
24	F	303	BCR	C24-C25	3.53	1.57	1.45
38	8	303	LUT	C19-C9	3.53	1.58	1.50
22	7	311	CLA	MG-ND	-3.53	1.98	2.05
22	6	314	CLA	MG-ND	-3.52	1.98	2.05
22	B	842	CLA	MG-ND	-3.52	1.98	2.05
22	A	819	CLA	MG-ND	-3.52	1.98	2.05
22	1	309	CLA	MG-ND	-3.52	1.98	2.05
22	Z	316	CLA	MG-ND	-3.52	1.98	2.05
21	A	801	CL0	MG-NC	3.52	2.14	2.06
22	A	823	CLA	MG-ND	-3.52	1.98	2.05
22	3	313	CLA	MG-ND	-3.52	1.98	2.05
22	A	824	CLA	MG-ND	-3.52	1.98	2.05
22	A	807	CLA	MG-ND	-3.52	1.98	2.05
22	A	805	CLA	MG-ND	-3.52	1.98	2.05
22	B	805	CLA	MG-ND	-3.52	1.98	2.05
22	A	841	CLA	MG-ND	-3.51	1.98	2.05
22	5	319	CLA	MG-ND	-3.51	1.98	2.05
24	6	305	BCR	C24-C25	3.51	1.57	1.45
22	A	810	CLA	MG-ND	-3.51	1.98	2.05
24	A	847	BCR	C24-C25	3.51	1.57	1.45
22	4	810	CLA	MG-ND	-3.51	1.98	2.05
22	A	834	CLA	MG-ND	-3.51	1.98	2.05

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
22	B	818	CLA	MG-ND	-3.51	1.98	2.05
24	B	847	BCR	C24-C25	3.50	1.57	1.45
22	A	811	CLA	MG-ND	-3.50	1.98	2.05
22	B	814	CLA	MG-ND	-3.50	1.98	2.05
22	A	833	CLA	MG-ND	-3.50	1.98	2.05
24	B	848	BCR	C24-C25	3.50	1.57	1.45
22	B	827	CLA	MG-ND	-3.50	1.98	2.05
22	8	316	CLA	MG-ND	-3.50	1.98	2.05
38	9	302	LUT	C2-C1	3.49	1.65	1.54
22	Z	314	CLA	MG-ND	-3.49	1.98	2.05
36	5	306	C7Z	C2-C1	3.49	1.65	1.54
22	5	318	CLA	MG-ND	-3.49	1.98	2.05
22	1	308	CLA	MG-ND	-3.49	1.98	2.05
22	3	318	CLA	MG-ND	-3.49	1.98	2.05
24	B	846	BCR	C24-C25	3.48	1.57	1.45
22	A	837	CLA	MG-ND	-3.48	1.98	2.05
22	B	838	CLA	MG-ND	-3.47	1.98	2.05
38	Z	302	LUT	C2-C1	3.47	1.65	1.54
22	9	309	CLA	C1C-NC	-3.47	1.32	1.37
24	B	846	BCR	C27-C26	-3.47	1.44	1.51
38	8	303	LUT	C2-C1	3.47	1.65	1.54
22	F	301	CLA	MG-ND	-3.46	1.98	2.05
38	7	301	LUT	C38-C25	3.46	1.57	1.50
22	A	855	CLA	MG-ND	-3.46	1.98	2.05
38	4	803	LUT	C2-C1	3.45	1.65	1.54
38	6	304	LUT	C2-C1	3.45	1.65	1.54
24	J	104	BCR	C27-C26	-3.45	1.44	1.51
38	3	303	LUT	C2-C1	3.45	1.65	1.54
24	B	845	BCR	C4-C5	-3.45	1.44	1.51
24	I	4001	BCR	C24-C25	3.45	1.57	1.45
22	1	314	CLA	MG-ND	-3.45	1.99	2.05
24	G	203	BCR	C27-C26	-3.44	1.44	1.51
22	4	805	CLA	C1C-NC	-3.44	1.32	1.37
38	7	301	LUT	C2-C1	3.44	1.65	1.54
24	B	845	BCR	C27-C26	-3.44	1.44	1.51
24	7	303	BCR	C4-C5	-3.43	1.44	1.51
24	L	204	BCR	C4-C5	-3.43	1.44	1.51
22	9	304	CLA	C1C-NC	-3.43	1.32	1.37
22	8	305	CLA	C1C-NC	-3.43	1.32	1.37
38	5	303	LUT	C2-C1	3.42	1.65	1.54
24	3	306	BCR	C4-C5	-3.42	1.44	1.51
38	1	303	LUT	C2-C1	3.42	1.65	1.54

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
24	3	304	BCR	C4-C5	-3.42	1.44	1.51
38	7	302	LUT	C2-C1	3.42	1.65	1.54
22	A	808	CLA	MG-ND	-3.42	1.99	2.05
36	J	105	C7Z	C18-C5	3.41	1.56	1.50
24	5	304	BCR	C27-C26	-3.41	1.44	1.51
22	L	201	CLA	MG-ND	-3.41	1.99	2.05
38	8	302	LUT	C2-C1	3.41	1.65	1.54
38	4	802	LUT	C2-C1	3.40	1.65	1.54
38	1	302	LUT	C2-C1	3.40	1.65	1.54
38	Z	301	LUT	C2-C1	3.40	1.65	1.54
38	8	303	LUT	C40-C33	3.40	1.57	1.50
27	A	852	DGA	OG2-CB1	3.40	1.43	1.34
38	6	303	LUT	C2-C1	3.40	1.65	1.54
24	8	304	BCR	C4-C5	-3.40	1.44	1.51
38	5	302	LUT	C2-C1	3.39	1.65	1.54
38	9	301	LUT	C2-C1	3.39	1.65	1.54
24	A	848	BCR	C4-C5	-3.39	1.44	1.51
24	6	306	BCR	C24-C25	3.39	1.57	1.45
24	A	844	BCR	C27-C26	-3.38	1.44	1.51
24	4	804	BCR	C4-C5	-3.38	1.44	1.51
24	8	304	BCR	C27-C26	-3.38	1.44	1.51
24	3	307	BCR	C4-C5	-3.38	1.44	1.51
38	Z	301	LUT	C38-C25	3.38	1.56	1.50
24	3	307	BCR	C27-C26	-3.38	1.44	1.51
38	6	304	LUT	C40-C33	3.38	1.57	1.50
38	7	302	LUT	C40-C33	3.37	1.57	1.50
24	A	856	BCR	C27-C26	-3.37	1.44	1.51
39	Z	312	CHL	CBB-CAB	3.37	1.51	1.29
36	5	306	C7Z	C18-C5	3.37	1.56	1.50
22	A	817	CLA	CBB-CAB	3.37	1.51	1.29
39	4	814	CHL	CBB-CAB	3.37	1.51	1.29
38	8	302	LUT	C40-C33	3.37	1.57	1.50
22	A	830	CLA	CBB-CAB	3.37	1.51	1.29
24	B	848	BCR	C4-C5	-3.37	1.44	1.51
22	8	307	CLA	CBB-CAB	3.37	1.51	1.29
22	5	322	CLA	CBB-CAB	3.36	1.51	1.29
22	6	308	CLA	CBB-CAB	3.36	1.51	1.29
24	A	846	BCR	C27-C26	-3.36	1.44	1.51
24	I	4001	BCR	C27-C26	-3.36	1.44	1.51
38	Z	302	LUT	C40-C33	3.36	1.57	1.50
38	4	802	LUT	C38-C25	3.36	1.56	1.50
22	B	831	CLA	CBB-CAB	3.36	1.51	1.29

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
22	Z	306	CLA	CBB-CAB	3.36	1.51	1.29
39	4	816	CHL	CBB-CAB	3.36	1.51	1.29
38	3	303	LUT	C40-C33	3.36	1.57	1.50
39	6	318	CHL	CBB-CAB	3.36	1.51	1.29
39	6	316	CHL	CBB-CAB	3.36	1.51	1.29
38	1	303	LUT	C40-C33	3.36	1.57	1.50
24	3	304	BCR	C27-C26	-3.36	1.44	1.51
24	5	304	BCR	C4-C5	-3.36	1.44	1.51
22	L	201	CLA	CBB-CAB	3.36	1.51	1.29
22	1	308	CLA	CBB-CAB	3.36	1.51	1.29
22	A	808	CLA	CBB-CAB	3.36	1.51	1.29
22	8	313	CLA	CBB-CAB	3.36	1.51	1.29
22	5	309	CLA	CBB-CAB	3.36	1.51	1.29
22	B	834	CLA	CBB-CAB	3.36	1.51	1.29
22	B	808	CLA	CBB-CAB	3.36	1.51	1.29
22	Z	310	CLA	CBB-CAB	3.36	1.51	1.29
22	A	829	CLA	CBB-CAB	3.35	1.51	1.29
22	Z	304	CLA	CBB-CAB	3.35	1.51	1.29
22	B	822	CLA	CBB-CAB	3.35	1.51	1.29
22	6	321	CLA	CBB-CAB	3.35	1.51	1.29
22	4	809	CLA	CBB-CAB	3.35	1.51	1.29
22	5	318	CLA	CBB-CAB	3.35	1.51	1.29
22	2	301	CLA	CBB-CAB	3.35	1.51	1.29
22	7	304	CLA	CBB-CAB	3.35	1.51	1.29
22	1	308	CLA	C1C-NC	-3.35	1.32	1.37
38	9	302	LUT	C40-C33	3.35	1.57	1.50
22	Z	305	CLA	CBB-CAB	3.35	1.51	1.29
22	7	310	CLA	CBB-CAB	3.35	1.51	1.29
22	1	305	CLA	CBB-CAB	3.35	1.51	1.29
22	1	307	CLA	CBB-CAB	3.35	1.51	1.29
22	Z	308	CLA	CBB-CAB	3.35	1.51	1.29
22	7	311	CLA	CBB-CAB	3.35	1.51	1.29
22	8	310	CLA	CBB-CAB	3.35	1.51	1.29
22	3	314	CLA	CBB-CAB	3.35	1.51	1.29
39	1	312	CHL	CBB-CAB	3.35	1.51	1.29
22	A	820	CLA	CBB-CAB	3.35	1.51	1.29
38	7	301	LUT	C40-C33	3.35	1.57	1.50
22	1	309	CLA	CBB-CAB	3.35	1.51	1.29
22	8	306	CLA	CBB-CAB	3.35	1.51	1.29
22	5	307	CLA	CBB-CAB	3.35	1.51	1.29
22	B	826	CLA	CBB-CAB	3.35	1.51	1.29
22	A	821	CLA	CBB-CAB	3.35	1.51	1.29

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
22	A	807	CLA	CBB-CAB	3.35	1.51	1.29
22	3	313	CLA	CBB-CAB	3.35	1.51	1.29
22	4	810	CLA	CBB-CAB	3.35	1.51	1.29
22	5	313	CLA	CBB-CAB	3.35	1.51	1.29
22	9	311	CLA	CBB-CAB	3.35	1.51	1.29
22	B	838	CLA	CBB-CAB	3.35	1.51	1.29
22	1	304	CLA	CBB-CAB	3.35	1.51	1.29
22	A	826	CLA	CBB-CAB	3.35	1.51	1.29
22	7	314	CLA	CBB-CAB	3.35	1.51	1.29
22	B	817	CLA	CBB-CAB	3.35	1.51	1.29
22	9	313	CLA	CBB-CAB	3.35	1.51	1.29
22	6	302	CLA	CBB-CAB	3.35	1.51	1.29
22	3	309	CLA	CBB-CAB	3.35	1.51	1.29
22	7	315	CLA	CBB-CAB	3.35	1.51	1.29
22	5	326	CLA	CBB-CAB	3.35	1.51	1.29
22	J	103	CLA	CBB-CAB	3.35	1.51	1.29
22	B	807	CLA	CBB-CAB	3.35	1.51	1.29
22	F	305	CLA	CBB-CAB	3.35	1.51	1.29
22	5	301	CLA	CBB-CAB	3.35	1.51	1.29
22	7	316	CLA	CBB-CAB	3.35	1.51	1.29
22	5	323	CLA	CBB-CAB	3.35	1.51	1.29
22	A	819	CLA	CBB-CAB	3.35	1.51	1.29
22	B	823	CLA	CBB-CAB	3.35	1.51	1.29
22	B	833	CLA	CBB-CAB	3.35	1.51	1.29
22	3	322	CLA	CBB-CAB	3.35	1.51	1.29
22	8	308	CLA	CBB-CAB	3.35	1.51	1.29
22	6	307	CLA	CBB-CAB	3.35	1.51	1.29
38	5	302	LUT	C40-C33	3.35	1.57	1.50
22	F	302	CLA	CBB-CAB	3.35	1.51	1.29
22	A	835	CLA	CBB-CAB	3.35	1.51	1.29
22	G	201	CLA	CBB-CAB	3.35	1.51	1.29
22	7	317	CLA	CBB-CAB	3.35	1.51	1.29
22	B	825	CLA	CBB-CAB	3.35	1.51	1.29
22	5	315	CLA	CBB-CAB	3.35	1.51	1.29
22	B	820	CLA	CBB-CAB	3.35	1.51	1.29
22	7	322	CLA	CBB-CAB	3.35	1.51	1.29
22	5	314	CLA	CBB-CAB	3.35	1.51	1.29
22	B	830	CLA	CBB-CAB	3.35	1.51	1.29
22	2	304	CLA	CBB-CAB	3.35	1.51	1.29
24	K	206	BCR	C27-C26	-3.35	1.44	1.51
22	3	310	CLA	CBB-CAB	3.35	1.51	1.29
22	8	311	CLA	CBB-CAB	3.35	1.51	1.29

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
22	4	817	CLA	CBB-CAB	3.35	1.51	1.29
22	9	308	CLA	CBB-CAB	3.35	1.51	1.29
24	3	305	BCR	C27-C26	-3.35	1.44	1.51
22	A	854	CLA	CBB-CAB	3.35	1.51	1.29
22	B	815	CLA	C1C-NC	-3.35	1.32	1.37
22	A	834	CLA	CBB-CAB	3.35	1.51	1.29
22	6	317	CLA	CBB-CAB	3.35	1.51	1.29
22	B	801	CLA	CBB-CAB	3.35	1.51	1.29
22	A	810	CLA	CBB-CAB	3.34	1.51	1.29
22	5	310	CLA	CBB-CAB	3.34	1.51	1.29
39	2	303	CHL	CBB-CAB	3.34	1.51	1.29
22	A	828	CLA	CBB-CAB	3.34	1.51	1.29
22	A	825	CLA	CBB-CAB	3.34	1.51	1.29
22	B	819	CLA	CBB-CAB	3.34	1.51	1.29
22	B	835	CLA	CBB-CAB	3.34	1.51	1.29
22	2	302	CLA	CBB-CAB	3.34	1.51	1.29
39	8	312	CHL	CBB-CAB	3.34	1.51	1.29
22	B	841	CLA	CBB-CAB	3.34	1.51	1.29
22	K	202	CLA	CBB-CAB	3.34	1.51	1.29
22	1	315	CLA	CBB-CAB	3.34	1.51	1.29
22	3	312	CLA	CBB-CAB	3.34	1.51	1.29
22	9	307	CLA	CBB-CAB	3.34	1.51	1.29
22	A	804	CLA	CBB-CAB	3.34	1.51	1.29
38	4	802	LUT	C40-C33	3.34	1.57	1.50
38	5	303	LUT	C40-C33	3.34	1.57	1.50
22	B	821	CLA	CBB-CAB	3.34	1.51	1.29
22	5	308	CLA	CBB-CAB	3.34	1.51	1.29
22	4	815	CLA	CBB-CAB	3.34	1.51	1.29
22	B	829	CLA	CBB-CAB	3.34	1.51	1.29
36	J	105	C7Z	C8-C9	3.34	1.53	1.45
24	A	846	BCR	C4-C5	-3.34	1.44	1.51
22	B	840	CLA	CBB-CAB	3.34	1.51	1.29
22	1	310	CLA	CBB-CAB	3.34	1.51	1.29
22	3	311	CLA	CBB-CAB	3.34	1.51	1.29
22	A	823	CLA	CBB-CAB	3.34	1.51	1.29
22	4	812	CLA	CBB-CAB	3.34	1.51	1.29
22	3	319	CLA	CBB-CAB	3.34	1.51	1.29
22	A	831	CLA	CBB-CAB	3.34	1.51	1.29
22	A	855	CLA	CBB-CAB	3.34	1.51	1.29
22	F	301	CLA	CBB-CAB	3.34	1.51	1.29
22	Z	313	CLA	CBB-CAB	3.34	1.51	1.29
22	B	836	CLA	CBB-CAB	3.34	1.51	1.29

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
22	8	314	CLA	CBB-CAB	3.34	1.51	1.29
22	7	323	CLA	CBB-CAB	3.34	1.51	1.29
22	B	809	CLA	CBB-CAB	3.34	1.51	1.29
22	Z	303	CLA	CBB-CAB	3.34	1.51	1.29
22	A	832	CLA	CBB-CAB	3.34	1.51	1.29
22	7	309	CLA	CBB-CAB	3.34	1.51	1.29
22	8	316	CLA	CBB-CAB	3.34	1.51	1.29
24	K	206	BCR	C4-C5	-3.34	1.44	1.51
22	4	818	CLA	CBB-CAB	3.34	1.51	1.29
22	A	824	CLA	CBB-CAB	3.34	1.51	1.29
22	G	202	CLA	CBB-CAB	3.34	1.51	1.29
22	7	307	CLA	CBB-CAB	3.34	1.51	1.29
22	1	314	CLA	CBB-CAB	3.34	1.51	1.29
22	7	308	CLA	CBB-CAB	3.34	1.51	1.29
22	B	827	CLA	CBB-CAB	3.34	1.51	1.29
22	7	312	CLA	CBB-CAB	3.34	1.51	1.29
22	5	312	CLA	CBB-CAB	3.34	1.51	1.29
24	6	305	BCR	C27-C26	-3.34	1.44	1.51
22	6	319	CLA	CBB-CAB	3.34	1.51	1.29
22	A	818	CLA	CBB-CAB	3.34	1.51	1.29
39	4	813	CHL	CBB-CAB	3.34	1.51	1.29
22	7	324	CLA	CBB-CAB	3.34	1.51	1.29
39	8	315	CHL	CBB-CAB	3.34	1.51	1.29
24	J	104	BCR	C4-C5	-3.34	1.44	1.51
22	B	813	CLA	CBB-CAB	3.34	1.51	1.29
22	3	318	CLA	CBB-CAB	3.34	1.51	1.29
22	B	812	CLA	CBB-CAB	3.34	1.51	1.29
22	A	833	CLA	CBB-CAB	3.34	1.51	1.29
22	B	832	CLA	CBB-CAB	3.34	1.51	1.29
22	6	301	CLA	CBB-CAB	3.34	1.51	1.29
22	A	815	CLA	CBB-CAB	3.34	1.51	1.29
22	1	313	CLA	CBB-CAB	3.34	1.51	1.29
24	B	853	BCR	C4-C5	-3.34	1.44	1.51
22	3	308	CLA	CBB-CAB	3.34	1.51	1.29
22	A	805	CLA	CBB-CAB	3.34	1.51	1.29
22	B	842	CLA	CBB-CAB	3.34	1.51	1.29
22	A	814	CLA	CBB-CAB	3.34	1.51	1.29
22	A	822	CLA	CBB-CAB	3.34	1.51	1.29
22	L	202	CLA	CBB-CAB	3.34	1.51	1.29
22	A	802	CLA	CBB-CAB	3.34	1.51	1.29
22	6	311	CLA	CBB-CAB	3.34	1.51	1.29
22	3	316	CLA	CBB-CAB	3.34	1.51	1.29

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
38	9	301	LUT	C40-C33	3.34	1.57	1.50
22	A	841	CLA	CBB-CAB	3.34	1.51	1.29
22	1	306	CLA	CBB-CAB	3.34	1.51	1.29
22	Z	314	CLA	CBB-CAB	3.34	1.51	1.29
22	B	818	CLA	CBB-CAB	3.34	1.51	1.29
22	6	313	CLA	CBB-CAB	3.34	1.51	1.29
22	B	816	CLA	CBB-CAB	3.34	1.51	1.29
22	B	839	CLA	CBB-CAB	3.34	1.51	1.29
22	B	811	CLA	CBB-CAB	3.34	1.51	1.29
22	A	803	CLA	CBB-CAB	3.34	1.51	1.29
22	L	203	CLA	CBB-CAB	3.34	1.51	1.29
22	4	806	CLA	CBB-CAB	3.34	1.51	1.29
24	A	845	BCR	C4-C5	-3.34	1.44	1.51
22	B	837	CLA	CBB-CAB	3.33	1.51	1.29
22	7	305	CLA	CBB-CAB	3.33	1.51	1.29
22	B	814	CLA	CBB-CAB	3.33	1.51	1.29
22	1	311	CLA	CBB-CAB	3.33	1.51	1.29
22	6	322	CLA	CBB-CAB	3.33	1.51	1.29
22	A	839	CLA	CBB-CAB	3.33	1.51	1.29
39	9	312	CHL	CBB-CAB	3.33	1.51	1.29
38	Z	301	LUT	C40-C33	3.33	1.57	1.50
24	A	847	BCR	C4-C5	-3.33	1.44	1.51
22	B	805	CLA	CBB-CAB	3.33	1.51	1.29
22	6	312	CLA	CBB-CAB	3.33	1.51	1.29
22	4	807	CLA	CBB-CAB	3.33	1.51	1.29
22	B	824	CLA	CBB-CAB	3.33	1.51	1.29
22	Z	307	CLA	CBB-CAB	3.33	1.51	1.29
22	A	816	CLA	CBB-CAB	3.33	1.51	1.29
22	4	811	CLA	CBB-CAB	3.33	1.51	1.29
22	9	306	CLA	CBB-CAB	3.33	1.51	1.29
22	3	315	CLA	CBB-CAB	3.33	1.51	1.29
22	A	812	CLA	CBB-CAB	3.33	1.51	1.29
22	6	314	CLA	CBB-CAB	3.33	1.51	1.29
22	A	809	CLA	CBB-CAB	3.33	1.51	1.29
22	Z	316	CLA	CBB-CAB	3.33	1.51	1.29
24	B	848	BCR	C27-C26	-3.33	1.44	1.51
22	6	309	CLA	CBB-CAB	3.33	1.51	1.29
39	5	317	CHL	CBB-CAB	3.33	1.51	1.29
24	B	853	BCR	C24-C25	3.33	1.56	1.45
24	A	844	BCR	C4-C5	-3.33	1.44	1.51
22	A	840	CLA	CBB-CAB	3.33	1.51	1.29
22	8	309	CLA	CBB-CAB	3.33	1.51	1.29

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
22	8	305	CLA	CBB-CAB	3.33	1.51	1.29
24	6	306	BCR	C4-C5	-3.33	1.44	1.51
22	B	815	CLA	CBB-CAB	3.33	1.51	1.29
22	B	828	CLA	CBB-CAB	3.33	1.51	1.29
22	Z	309	CLA	CBB-CAB	3.33	1.51	1.29
22	F	304	CLA	CBB-CAB	3.33	1.51	1.29
38	3	302	LUT	C40-C33	3.33	1.57	1.50
22	4	805	CLA	CBB-CAB	3.33	1.51	1.29
36	J	105	C7Z	C7-C6	3.33	1.56	1.45
22	5	311	CLA	CBB-CAB	3.32	1.51	1.29
22	A	811	CLA	CBB-CAB	3.32	1.51	1.29
22	5	319	CLA	CBB-CAB	3.32	1.51	1.29
24	6	306	BCR	C27-C26	-3.32	1.44	1.51
22	A	842	CLA	CBB-CAB	3.32	1.51	1.29
22	A	806	CLA	CBB-CAB	3.32	1.51	1.29
22	Z	315	CLA	CBB-CAB	3.32	1.51	1.29
22	A	837	CLA	CBB-CAB	3.32	1.51	1.29
39	9	314	CHL	CBB-CAB	3.32	1.51	1.29
22	A	813	CLA	CBB-CAB	3.32	1.51	1.29
38	6	303	LUT	C40-C33	3.32	1.57	1.50
22	K	203	CLA	CBB-CAB	3.32	1.51	1.29
22	A	836	CLA	CBB-CAB	3.32	1.51	1.29
22	7	306	CLA	CBB-CAB	3.32	1.51	1.29
22	A	838	CLA	CBB-CAB	3.32	1.51	1.29
22	G	201	CLA	C1C-NC	-3.32	1.32	1.37
39	3	317	CHL	CBB-CAB	3.32	1.51	1.29
22	B	806	CLA	CBB-CAB	3.32	1.51	1.29
22	4	808	CLA	CBB-CAB	3.31	1.51	1.29
22	B	810	CLA	CBB-CAB	3.31	1.51	1.29
22	K	204	CLA	CBB-CAB	3.31	1.51	1.29
22	K	205	CLA	CBB-CAB	3.31	1.51	1.29
22	9	304	CLA	CBB-CAB	3.31	1.51	1.29
39	4	819	CHL	CBB-CAB	3.31	1.51	1.29
24	B	847	BCR	C4-C5	-3.31	1.44	1.51
22	6	310	CLA	CBB-CAB	3.31	1.51	1.29
22	A	827	CLA	CBB-CAB	3.31	1.51	1.29
24	B	849	BCR	C4-C5	-3.30	1.44	1.51
22	A	813	CLA	C1C-NC	-3.30	1.32	1.37
38	3	302	LUT	C2-C1	3.30	1.65	1.54
39	7	313	CHL	CBB-CAB	3.30	1.51	1.29
38	4	803	LUT	C40-C33	3.30	1.57	1.50
39	8	301	CHL	CBB-CAB	3.30	1.51	1.29

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
22	9	305	CLA	CBB-CAB	3.30	1.51	1.29
24	5	305	BCR	C4-C5	-3.30	1.44	1.51
24	L	204	BCR	C27-C26	-3.30	1.44	1.51
38	1	302	LUT	C40-C33	3.30	1.57	1.50
22	1	316	CLA	CBB-CAB	3.29	1.51	1.29
39	Z	311	CHL	CBB-CAB	3.29	1.51	1.29
39	6	315	CHL	CBB-CAB	3.29	1.51	1.29
22	Z	304	CLA	C1C-NC	-3.29	1.32	1.37
39	5	321	CHL	CBB-CAB	3.29	1.51	1.29
24	B	844	BCR	C27-C26	-3.29	1.44	1.51
22	9	310	CLA	CBB-CAB	3.28	1.51	1.29
22	4	809	CLA	C1C-NC	-3.28	1.32	1.37
22	B	814	CLA	C1C-NC	-3.28	1.32	1.37
22	9	309	CLA	CBB-CAB	3.28	1.51	1.29
22	3	320	CLA	CBB-CAB	3.28	1.51	1.29
24	B	849	BCR	C27-C26	-3.28	1.44	1.51
24	B	844	BCR	C4-C5	-3.28	1.44	1.51
24	B	847	BCR	C27-C26	-3.28	1.44	1.51
36	J	105	C7Z	C21-C26	-3.28	1.49	1.53
31	B	852	DGD	CAA-C9A	-3.27	1.33	1.51
24	4	804	BCR	C27-C26	-3.27	1.44	1.51
31	B	852	DGD	CDB-CCB	-3.27	1.33	1.51
24	A	848	BCR	C27-C26	-3.27	1.44	1.51
39	6	320	CHL	CBB-CAB	3.27	1.50	1.29
24	I	4001	BCR	C4-C5	-3.26	1.44	1.51
22	5	320	CLA	CBB-CAB	3.26	1.50	1.29
22	6	302	CLA	C1C-NC	-3.26	1.32	1.37
22	B	818	CLA	C1C-NC	-3.26	1.32	1.37
22	A	827	CLA	C1C-NC	-3.26	1.32	1.37
27	A	852	DGA	OG1-CA1	3.26	1.42	1.33
31	B	852	DGD	CAB-C9B	-3.26	1.33	1.51
24	3	306	BCR	C27-C26	-3.26	1.44	1.51
36	5	306	C7Z	C7-C6	3.26	1.56	1.45
39	8	301	CHL	C4B-NB	3.25	1.38	1.35
22	1	305	CLA	C1C-NC	-3.25	1.33	1.37
22	Z	303	CLA	C1C-NC	-3.25	1.33	1.37
22	8	306	CLA	C1C-NC	-3.25	1.33	1.37
24	A	845	BCR	C27-C26	-3.25	1.44	1.51
22	1	313	CLA	C1C-NC	-3.25	1.33	1.37
39	5	316	CHL	CBB-CAB	3.25	1.50	1.29
22	Z	305	CLA	C1C-NC	-3.24	1.33	1.37
22	3	312	CLA	C1C-NC	-3.24	1.33	1.37

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
22	Z	316	CLA	C1C-NC	-3.24	1.33	1.37
39	3	317	CHL	C4B-NB	3.24	1.38	1.35
31	B	852	DGD	CGB-CFB	-3.23	1.33	1.51
22	A	805	CLA	C1C-NC	-3.23	1.33	1.37
31	1	319	DGD	CAA-C9A	-3.23	1.33	1.51
22	A	835	CLA	C1C-NC	-3.23	1.33	1.37
22	4	818	CLA	C1C-NC	-3.23	1.33	1.37
24	F	303	BCR	C4-C5	-3.23	1.44	1.51
22	A	823	CLA	C1C-NC	-3.23	1.33	1.37
31	3	301	DGD	CAA-C9A	-3.23	1.33	1.51
22	B	817	CLA	C1C-NC	-3.22	1.33	1.37
22	3	316	CLA	C1C-NC	-3.22	1.33	1.37
22	B	830	CLA	C1C-NC	-3.22	1.33	1.37
22	6	301	CLA	C1C-NC	-3.22	1.33	1.37
24	B	846	BCR	C4-C5	-3.22	1.44	1.51
22	2	304	CLA	C1C-NC	-3.21	1.33	1.37
22	A	812	CLA	C1C-NC	-3.21	1.33	1.37
22	3	311	CLA	C1C-NC	-3.21	1.33	1.37
22	6	307	CLA	C1C-NC	-3.21	1.33	1.37
22	1	304	CLA	C1C-NC	-3.21	1.33	1.37
22	Z	313	CLA	C1C-NC	-3.21	1.33	1.37
36	5	306	C7Z	C8-C9	3.21	1.52	1.45
22	5	314	CLA	C1C-NC	-3.21	1.33	1.37
22	A	855	CLA	C1C-NC	-3.20	1.33	1.37
32	F	306	RRX	C4-C5	-3.20	1.44	1.51
22	7	312	CLA	C1C-NC	-3.20	1.33	1.37
22	9	311	CLA	C1C-NC	-3.20	1.33	1.37
22	5	309	CLA	C1C-NC	-3.20	1.33	1.37
22	9	303	CLA	CBB-CAB	3.20	1.50	1.29
39	Z	311	CHL	C4B-NB	3.20	1.38	1.35
38	9	301	LUT	C23-C24	-3.20	1.45	1.50
22	A	828	CLA	C1C-NC	-3.20	1.33	1.37
22	B	827	CLA	C1C-NC	-3.20	1.33	1.37
22	B	811	CLA	C1C-NC	-3.20	1.33	1.37
22	8	311	CLA	C1C-NC	-3.20	1.33	1.37
22	9	308	CLA	C1C-NC	-3.19	1.33	1.37
22	7	307	CLA	C1C-NC	-3.19	1.33	1.37
22	A	840	CLA	C1C-NC	-3.19	1.33	1.37
39	2	303	CHL	C4B-NB	3.19	1.38	1.35
22	A	806	CLA	C1C-NC	-3.19	1.33	1.37
22	Z	309	CLA	C1C-NC	-3.19	1.33	1.37
22	F	301	CLA	C1C-NC	-3.19	1.33	1.37

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
22	K	203	CLA	C1C-NC	-3.19	1.33	1.37
22	8	316	CLA	C1C-NC	-3.19	1.33	1.37
24	A	847	BCR	C27-C26	-3.19	1.44	1.51
38	Z	301	LUT	C23-C24	-3.19	1.45	1.50
22	B	821	CLA	C1C-NC	-3.19	1.33	1.37
24	A	856	BCR	C4-C5	-3.19	1.44	1.51
24	7	303	BCR	C27-C26	-3.18	1.44	1.51
39	9	314	CHL	C3A-C2A	-3.18	1.51	1.54
22	6	311	CLA	C1C-NC	-3.18	1.33	1.37
22	1	307	CLA	C1C-NC	-3.18	1.33	1.37
22	B	823	CLA	C1C-NC	-3.18	1.33	1.37
22	5	301	CLA	C1C-NC	-3.18	1.33	1.37
22	G	202	CLA	C1C-NC	-3.18	1.33	1.37
22	7	305	CLA	C1C-NC	-3.18	1.33	1.37
22	B	834	CLA	C1C-NC	-3.18	1.33	1.37
22	2	302	CLA	C1C-NC	-3.18	1.33	1.37
24	F	303	BCR	C27-C26	-3.18	1.44	1.51
22	B	819	CLA	C1C-NC	-3.18	1.33	1.37
22	5	308	CLA	C1C-NC	-3.18	1.33	1.37
22	A	811	CLA	C1C-NC	-3.17	1.33	1.37
22	A	803	CLA	C1C-NC	-3.17	1.33	1.37
22	A	833	CLA	C1C-NC	-3.17	1.33	1.37
22	3	309	CLA	C1C-NC	-3.17	1.33	1.37
22	J	103	CLA	C1C-NC	-3.17	1.33	1.37
22	Z	315	CLA	C1C-NC	-3.17	1.33	1.37
22	A	821	CLA	C1C-NC	-3.17	1.33	1.37
22	9	306	CLA	C1C-NC	-3.17	1.33	1.37
22	9	313	CLA	C1C-NC	-3.17	1.33	1.37
22	B	828	CLA	C1C-NC	-3.17	1.33	1.37
22	5	307	CLA	C1C-NC	-3.17	1.33	1.37
22	9	307	CLA	C1C-NC	-3.17	1.33	1.37
22	A	809	CLA	C1C-NC	-3.17	1.33	1.37
22	2	301	CLA	C1C-NC	-3.17	1.33	1.37
22	A	824	CLA	C1C-NC	-3.17	1.33	1.37
22	3	308	CLA	C1C-NC	-3.17	1.33	1.37
22	A	817	CLA	C1C-NC	-3.16	1.33	1.37
22	A	818	CLA	C1C-NC	-3.16	1.33	1.37
22	A	820	CLA	C1C-NC	-3.16	1.33	1.37
22	B	822	CLA	C1C-NC	-3.16	1.33	1.37
22	F	304	CLA	C1C-NC	-3.16	1.33	1.37
22	3	314	CLA	C1C-NC	-3.16	1.33	1.37
22	B	832	CLA	C1C-NC	-3.16	1.33	1.37

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
22	6	313	CLA	C1C-NC	-3.16	1.33	1.37
22	6	314	CLA	C1C-NC	-3.16	1.33	1.37
25	B	851	LHG	O7-C5	-3.16	1.43	1.46
22	5	326	CLA	C1C-NC	-3.16	1.33	1.37
24	G	203	BCR	C4-C5	-3.16	1.44	1.51
22	L	203	CLA	C1C-NC	-3.16	1.33	1.37
22	B	809	CLA	C1C-NC	-3.16	1.33	1.37
22	6	317	CLA	C1C-NC	-3.16	1.33	1.37
22	7	308	CLA	C1C-NC	-3.16	1.33	1.37
22	A	839	CLA	C1C-NC	-3.16	1.33	1.37
22	A	829	CLA	C1C-NC	-3.16	1.33	1.37
22	4	807	CLA	C1C-NC	-3.16	1.33	1.37
22	7	322	CLA	C1C-NC	-3.16	1.33	1.37
22	B	825	CLA	C1C-NC	-3.15	1.33	1.37
22	A	808	CLA	C1C-NC	-3.15	1.33	1.37
22	8	313	CLA	C1C-NC	-3.15	1.33	1.37
22	7	304	CLA	C1C-NC	-3.15	1.33	1.37
22	5	312	CLA	C1C-NC	-3.15	1.33	1.37
39	6	316	CHL	C4B-NB	3.15	1.38	1.35
22	A	838	CLA	C1C-NC	-3.15	1.33	1.37
22	K	202	CLA	C1C-NC	-3.15	1.33	1.37
22	8	309	CLA	C1C-NC	-3.15	1.33	1.37
22	A	841	CLA	C1C-NC	-3.15	1.33	1.37
22	7	310	CLA	C1C-NC	-3.15	1.33	1.37
22	Z	310	CLA	C1C-NC	-3.15	1.33	1.37
22	A	836	CLA	C1C-NC	-3.15	1.33	1.37
22	F	302	CLA	C1C-NC	-3.15	1.33	1.37
22	L	201	CLA	C1C-NC	-3.15	1.33	1.37
22	4	811	CLA	C1C-NC	-3.15	1.33	1.37
24	5	305	BCR	C27-C26	-3.15	1.44	1.51
39	7	313	CHL	C4B-NB	3.14	1.38	1.35
22	7	314	CLA	C1C-NC	-3.14	1.33	1.37
22	B	839	CLA	C1C-NC	-3.14	1.33	1.37
22	4	817	CLA	C1C-NC	-3.14	1.33	1.37
22	5	315	CLA	C1C-NC	-3.14	1.33	1.37
22	Z	314	CLA	C1C-NC	-3.14	1.33	1.37
22	B	836	CLA	C1C-NC	-3.14	1.33	1.37
22	A	804	CLA	C1C-NC	-3.14	1.33	1.37
22	A	842	CLA	C1C-NC	-3.14	1.33	1.37
22	5	319	CLA	C1C-NC	-3.14	1.33	1.37
24	B	853	BCR	C27-C26	-3.14	1.44	1.51
22	Z	307	CLA	C1C-NC	-3.14	1.33	1.37

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
22	B	829	CLA	C1C-NC	-3.14	1.33	1.37
22	1	310	CLA	C1C-NC	-3.14	1.33	1.37
22	3	313	CLA	C1C-NC	-3.14	1.33	1.37
22	8	310	CLA	C1C-NC	-3.14	1.33	1.37
22	A	830	CLA	C1C-NC	-3.14	1.33	1.37
22	Z	306	CLA	C1C-NC	-3.14	1.33	1.37
22	B	837	CLA	C1C-NC	-3.13	1.33	1.37
22	4	815	CLA	C1C-NC	-3.13	1.33	1.37
22	B	835	CLA	C1C-NC	-3.13	1.33	1.37
22	7	309	CLA	C1C-NC	-3.13	1.33	1.37
36	5	306	C7Z	C21-C26	-3.13	1.49	1.53
22	B	805	CLA	C1C-NC	-3.13	1.33	1.37
39	4	813	CHL	C4B-NB	3.13	1.38	1.35
22	L	202	CLA	C1C-NC	-3.13	1.33	1.37
22	1	314	CLA	C1C-NC	-3.13	1.33	1.37
22	4	812	CLA	C1C-NC	-3.13	1.33	1.37
22	4	808	CLA	C1C-NC	-3.13	1.33	1.37
22	A	825	CLA	C1C-NC	-3.13	1.33	1.37
22	3	319	CLA	C1C-NC	-3.13	1.33	1.37
22	B	831	CLA	C1C-NC	-3.13	1.33	1.37
22	8	314	CLA	C1C-NC	-3.13	1.33	1.37
22	B	841	CLA	C1C-NC	-3.12	1.33	1.37
22	A	810	CLA	C1C-NC	-3.12	1.33	1.37
22	8	308	CLA	C1C-NC	-3.12	1.33	1.37
22	5	323	CLA	C1C-NC	-3.12	1.33	1.37
22	5	318	CLA	C1C-NC	-3.12	1.33	1.37
22	A	822	CLA	C1C-NC	-3.12	1.33	1.37
22	B	801	CLA	C1C-NC	-3.12	1.33	1.37
22	5	313	CLA	C1C-NC	-3.12	1.33	1.37
39	4	816	CHL	C4B-NB	3.12	1.38	1.35
39	5	317	CHL	C4B-NB	3.12	1.38	1.35
22	6	322	CLA	C1C-NC	-3.12	1.33	1.37
22	1	306	CLA	C1C-NC	-3.12	1.33	1.37
22	7	315	CLA	C1C-NC	-3.12	1.33	1.37
38	8	302	LUT	C23-C24	-3.12	1.45	1.50
22	A	832	CLA	C1C-NC	-3.12	1.33	1.37
22	B	810	CLA	C1C-NC	-3.12	1.33	1.37
22	B	808	CLA	C1C-NC	-3.11	1.33	1.37
22	3	310	CLA	C1C-NC	-3.11	1.33	1.37
22	B	807	CLA	C1C-NC	-3.11	1.33	1.37
22	1	309	CLA	C1C-NC	-3.11	1.33	1.37
22	A	814	CLA	C1C-NC	-3.11	1.33	1.37

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
22	6	312	CLA	C1C-NC	-3.11	1.33	1.37
22	A	815	CLA	C1C-NC	-3.11	1.33	1.37
22	7	306	CLA	C1C-NC	-3.11	1.33	1.37
22	4	806	CLA	C1C-NC	-3.11	1.33	1.37
22	7	317	CLA	C1C-NC	-3.11	1.33	1.37
22	7	324	CLA	C1C-NC	-3.11	1.33	1.37
22	B	840	CLA	C1C-NC	-3.10	1.33	1.37
22	K	204	CLA	C1C-NC	-3.10	1.33	1.37
22	3	318	CLA	C1C-NC	-3.10	1.33	1.37
22	A	831	CLA	C1C-NC	-3.10	1.33	1.37
22	Z	308	CLA	C1C-NC	-3.10	1.33	1.37
22	B	833	CLA	C1C-NC	-3.10	1.33	1.37
22	B	812	CLA	C1C-NC	-3.10	1.33	1.37
22	7	311	CLA	C1C-NC	-3.10	1.33	1.37
22	4	810	CLA	C1C-NC	-3.10	1.33	1.37
39	9	314	CHL	C4B-NB	3.10	1.38	1.35
22	A	819	CLA	C1C-NC	-3.10	1.33	1.37
22	A	802	CLA	C1C-NC	-3.10	1.33	1.37
22	B	806	CLA	C1C-NC	-3.10	1.33	1.37
22	6	309	CLA	C1C-NC	-3.09	1.33	1.37
22	F	305	CLA	C1C-NC	-3.09	1.33	1.37
22	3	322	CLA	C1C-NC	-3.09	1.33	1.37
22	B	838	CLA	C1C-NC	-3.09	1.33	1.37
22	A	854	CLA	C1C-NC	-3.09	1.33	1.37
22	A	816	CLA	C1C-NC	-3.09	1.33	1.37
39	6	318	CHL	C4B-NB	3.09	1.38	1.35
22	7	323	CLA	C1C-NC	-3.09	1.33	1.37
22	6	308	CLA	C1C-NC	-3.08	1.33	1.37
22	6	321	CLA	C1C-NC	-3.08	1.33	1.37
39	5	316	CHL	C4B-NB	3.08	1.38	1.35
22	A	807	CLA	C1C-NC	-3.08	1.33	1.37
22	6	310	CLA	C1C-NC	-3.08	1.33	1.37
22	5	322	CLA	C1C-NC	-3.08	1.33	1.37
22	7	316	CLA	C1C-NC	-3.08	1.33	1.37
22	B	842	CLA	C1C-NC	-3.08	1.33	1.37
39	6	320	CHL	C4B-NB	3.07	1.38	1.35
22	5	311	CLA	C1C-NC	-3.07	1.33	1.37
22	5	310	CLA	C1C-NC	-3.07	1.33	1.37
22	A	837	CLA	C1C-NC	-3.07	1.33	1.37
39	Z	312	CHL	C4B-NB	3.07	1.37	1.35
38	4	802	LUT	C23-C24	-3.07	1.45	1.50
24	3	305	BCR	C4-C5	-3.06	1.45	1.51

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
22	B	824	CLA	C1C-NC	-3.06	1.33	1.37
38	1	302	LUT	C23-C24	-3.06	1.45	1.50
22	B	816	CLA	C1C-NC	-3.06	1.33	1.37
22	A	834	CLA	C1C-NC	-3.06	1.33	1.37
22	B	813	CLA	C1C-NC	-3.05	1.33	1.37
39	5	321	CHL	C4B-NB	3.05	1.37	1.35
22	6	319	CLA	C1C-NC	-3.05	1.33	1.37
22	B	820	CLA	C1C-NC	-3.04	1.33	1.37
22	1	315	CLA	C1C-NC	-3.04	1.33	1.37
39	4	814	CHL	C4B-NB	3.04	1.37	1.35
22	1	311	CLA	C3B-C2B	-3.04	1.36	1.40
22	B	826	CLA	C1C-NC	-3.03	1.33	1.37
39	8	315	CHL	C4B-NB	3.03	1.37	1.35
39	9	312	CHL	C4B-NB	3.03	1.37	1.35
39	4	819	CHL	C4B-NB	3.02	1.37	1.35
21	A	801	CL0	C1D-ND	-3.02	1.34	1.37
22	5	320	CLA	C3B-C2B	-3.02	1.36	1.40
38	7	301	LUT	C23-C24	-3.02	1.45	1.50
38	3	303	LUT	C23-C24	-3.01	1.45	1.50
39	6	315	CHL	C4B-NB	3.01	1.37	1.35
39	8	312	CHL	C4B-NB	3.00	1.37	1.35
22	8	307	CLA	C1C-NC	-3.00	1.33	1.37
21	A	801	CL0	C3D-C2D	2.99	1.47	1.39
22	A	826	CLA	C1C-NC	-2.99	1.33	1.37
22	3	320	CLA	C3B-C2B	-2.98	1.36	1.40
38	3	302	LUT	C23-C24	-2.97	1.45	1.50
38	Z	302	LUT	C23-C24	-2.94	1.46	1.50
39	1	312	CHL	C4B-NB	2.93	1.37	1.35
38	6	304	LUT	C23-C24	-2.93	1.46	1.50
38	8	303	LUT	C23-C24	-2.93	1.46	1.50
24	L	204	BCR	C34-C9	2.88	1.56	1.50
24	5	305	BCR	C34-C9	2.87	1.56	1.50
24	B	853	BCR	C34-C9	2.87	1.56	1.50
38	7	302	LUT	C23-C24	-2.85	1.46	1.50
24	A	845	BCR	C34-C9	2.85	1.56	1.50
24	B	849	BCR	C34-C9	2.85	1.56	1.50
24	B	847	BCR	C34-C9	2.84	1.56	1.50
24	A	846	BCR	C34-C9	2.84	1.56	1.50
24	F	303	BCR	C34-C9	2.84	1.56	1.50
38	5	302	LUT	C23-C24	-2.84	1.46	1.50
22	K	205	CLA	C3B-C2B	-2.82	1.36	1.40
22	B	824	CLA	CHC-C1C	2.82	1.42	1.35

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
24	A	847	BCR	C34-C9	2.82	1.56	1.50
24	5	304	BCR	C34-C9	2.82	1.56	1.50
24	6	306	BCR	C34-C9	2.82	1.56	1.50
24	B	844	BCR	C34-C9	2.82	1.56	1.50
22	9	305	CLA	C3B-C2B	-2.82	1.36	1.40
24	B	853	BCR	C7-C6	2.81	1.55	1.45
24	B	845	BCR	C34-C9	2.81	1.56	1.50
22	9	310	CLA	C3B-C2B	-2.81	1.36	1.40
24	3	307	BCR	C34-C9	2.81	1.56	1.50
24	6	305	BCR	C34-C9	2.81	1.56	1.50
21	A	801	CL0	C4D-CHA	2.81	1.48	1.38
38	4	803	LUT	C23-C24	-2.80	1.46	1.50
22	A	834	CLA	CHC-C1C	2.80	1.42	1.35
24	A	856	BCR	C34-C9	2.80	1.56	1.50
24	I	4001	BCR	C34-C9	2.80	1.56	1.50
24	3	305	BCR	C34-C9	2.79	1.56	1.50
22	3	318	CLA	CHC-C1C	2.79	1.42	1.35
22	F	305	CLA	CHC-C1C	2.79	1.42	1.35
22	A	823	CLA	CHC-C1C	2.79	1.42	1.35
24	5	305	BCR	C7-C6	2.79	1.55	1.45
22	A	825	CLA	CHC-C1C	2.78	1.42	1.35
22	7	322	CLA	CHC-C1C	2.78	1.42	1.35
38	9	302	LUT	C23-C24	-2.78	1.46	1.50
24	K	206	BCR	C34-C9	2.78	1.56	1.50
22	9	307	CLA	CHC-C1C	2.78	1.42	1.35
24	8	304	BCR	C34-C9	2.78	1.56	1.50
24	J	104	BCR	C34-C9	2.78	1.56	1.50
22	B	801	CLA	CHC-C1C	2.78	1.42	1.35
24	B	848	BCR	C34-C9	2.78	1.56	1.50
24	3	304	BCR	C34-C9	2.78	1.56	1.50
22	B	808	CLA	CHC-C1C	2.77	1.42	1.35
22	7	315	CLA	CHC-C1C	2.77	1.42	1.35
22	B	835	CLA	CHC-C1C	2.77	1.42	1.35
22	7	311	CLA	CHC-C1C	2.77	1.42	1.35
24	4	804	BCR	C34-C9	2.77	1.56	1.50
22	B	807	CLA	CHC-C1C	2.77	1.42	1.35
24	A	848	BCR	C34-C9	2.77	1.56	1.50
22	A	811	CLA	CHC-C1C	2.77	1.42	1.35
22	9	309	CLA	C3B-C2B	-2.77	1.36	1.40
22	A	819	CLA	CHC-C1C	2.77	1.42	1.35
24	7	303	BCR	C34-C9	2.77	1.56	1.50
22	A	809	CLA	CHC-C1C	2.76	1.42	1.35

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
24	G	203	BCR	C34-C9	2.76	1.56	1.50
22	7	324	CLA	CHC-C1C	2.76	1.42	1.35
22	A	824	CLA	CHC-C1C	2.76	1.42	1.35
22	6	310	CLA	CHC-C1C	2.76	1.42	1.35
22	A	841	CLA	CHC-C1C	2.76	1.42	1.35
22	4	806	CLA	CHC-C1C	2.76	1.42	1.35
22	B	831	CLA	CHC-C1C	2.76	1.42	1.35
22	Z	309	CLA	CHC-C1C	2.76	1.42	1.35
22	A	810	CLA	CHC-C1C	2.76	1.42	1.35
22	4	817	CLA	CHC-C1C	2.76	1.42	1.35
22	A	836	CLA	CHC-C1C	2.76	1.42	1.35
22	5	322	CLA	CHC-C1C	2.76	1.42	1.35
22	L	203	CLA	CHC-C1C	2.76	1.42	1.35
22	B	838	CLA	CHC-C1C	2.75	1.42	1.35
22	B	818	CLA	CHC-C1C	2.75	1.42	1.35
22	B	836	CLA	CHC-C1C	2.75	1.42	1.35
24	3	306	BCR	C7-C6	2.75	1.54	1.45
22	J	103	CLA	CHC-C1C	2.75	1.42	1.35
22	4	810	CLA	CHC-C1C	2.75	1.42	1.35
22	B	823	CLA	CHC-C1C	2.75	1.42	1.35
22	Z	306	CLA	CHC-C1C	2.75	1.42	1.35
22	7	316	CLA	CHC-C1C	2.75	1.42	1.35
22	Z	308	CLA	CHC-C1C	2.74	1.42	1.35
24	F	303	BCR	C7-C6	2.74	1.54	1.45
22	A	812	CLA	CHC-C1C	2.74	1.42	1.35
22	L	202	CLA	CHC-C1C	2.74	1.42	1.35
22	A	805	CLA	CHC-C1C	2.74	1.42	1.35
22	A	828	CLA	CHC-C1C	2.74	1.42	1.35
38	5	303	LUT	C23-C24	-2.74	1.46	1.50
22	6	317	CLA	CHC-C1C	2.74	1.42	1.35
22	9	304	CLA	C3B-C2B	-2.74	1.36	1.40
22	F	301	CLA	CHC-C1C	2.74	1.42	1.35
22	B	840	CLA	CHC-C1C	2.74	1.42	1.35
22	A	818	CLA	CHC-C1C	2.74	1.42	1.35
22	5	318	CLA	CHC-C1C	2.74	1.42	1.35
22	B	816	CLA	CHC-C1C	2.74	1.42	1.35
22	1	309	CLA	CHC-C1C	2.73	1.42	1.35
22	A	826	CLA	CHC-C1C	2.73	1.42	1.35
22	Z	314	CLA	CHC-C1C	2.73	1.42	1.35
22	6	301	CLA	CHC-C1C	2.73	1.42	1.35
22	6	313	CLA	CHC-C1C	2.73	1.42	1.35
22	A	803	CLA	CHC-C1C	2.73	1.42	1.35

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
22	A	854	CLA	CHC-C1C	2.73	1.42	1.35
22	B	822	CLA	CHC-C1C	2.73	1.42	1.35
24	B	848	BCR	C7-C6	2.73	1.54	1.45
22	1	314	CLA	CHC-C1C	2.73	1.42	1.35
22	B	828	CLA	CHC-C1C	2.73	1.42	1.35
22	Z	316	CLA	CHC-C1C	2.73	1.42	1.35
22	7	305	CLA	CHC-C1C	2.73	1.42	1.35
22	1	307	CLA	CHC-C1C	2.73	1.42	1.35
22	1	308	CLA	CHC-C1C	2.73	1.42	1.35
22	9	306	CLA	CHC-C1C	2.73	1.42	1.35
24	B	844	BCR	C7-C6	2.73	1.54	1.45
22	B	826	CLA	CHC-C1C	2.73	1.42	1.35
22	3	313	CLA	CHC-C1C	2.73	1.42	1.35
22	5	323	CLA	CHC-C1C	2.73	1.42	1.35
22	4	815	CLA	CHC-C1C	2.73	1.42	1.35
24	B	846	BCR	C34-C9	2.73	1.56	1.50
22	8	310	CLA	CHC-C1C	2.73	1.42	1.35
22	7	307	CLA	CHC-C1C	2.73	1.42	1.35
22	5	312	CLA	CHC-C1C	2.73	1.42	1.35
22	A	833	CLA	CHC-C1C	2.73	1.42	1.35
22	B	820	CLA	CHC-C1C	2.72	1.42	1.35
22	B	809	CLA	CHC-C1C	2.72	1.41	1.35
22	8	306	CLA	CHC-C1C	2.72	1.41	1.35
22	A	837	CLA	CHC-C1C	2.72	1.41	1.35
22	A	835	CLA	CHC-C1C	2.72	1.41	1.35
22	8	311	CLA	CHC-C1C	2.72	1.41	1.35
24	3	306	BCR	C34-C9	2.72	1.56	1.50
22	3	308	CLA	CHC-C1C	2.72	1.41	1.35
22	9	313	CLA	CHC-C1C	2.72	1.41	1.35
22	B	832	CLA	CHC-C1C	2.72	1.41	1.35
22	5	314	CLA	CHC-C1C	2.72	1.41	1.35
22	A	807	CLA	CHC-C1C	2.72	1.41	1.35
22	3	316	CLA	CHC-C1C	2.72	1.41	1.35
24	3	305	BCR	C7-C6	2.72	1.54	1.45
22	A	802	CLA	CHC-C1C	2.71	1.41	1.35
22	3	311	CLA	CHC-C1C	2.71	1.41	1.35
22	G	202	CLA	CHC-C1C	2.71	1.41	1.35
22	B	837	CLA	CHC-C1C	2.71	1.41	1.35
22	Z	307	CLA	CHC-C1C	2.71	1.41	1.35
22	B	817	CLA	CHC-C1C	2.71	1.41	1.35
22	6	321	CLA	CHC-C1C	2.71	1.41	1.35
22	B	825	CLA	CHC-C1C	2.71	1.41	1.35

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
22	A	831	CLA	CHC-C1C	2.71	1.41	1.35
22	K	202	CLA	CHC-C1C	2.71	1.41	1.35
22	5	315	CLA	CHC-C1C	2.71	1.41	1.35
22	Z	310	CLA	CHC-C1C	2.71	1.41	1.35
22	1	304	CLA	CHC-C1C	2.71	1.41	1.35
22	3	309	CLA	CHC-C1C	2.71	1.41	1.35
22	1	315	CLA	CHC-C1C	2.71	1.41	1.35
22	A	806	CLA	CHC-C1C	2.71	1.41	1.35
22	7	323	CLA	CHC-C1C	2.71	1.41	1.35
22	A	804	CLA	CHC-C1C	2.70	1.41	1.35
22	A	817	CLA	CHC-C1C	2.70	1.41	1.35
22	7	314	CLA	CHC-C1C	2.70	1.41	1.35
22	8	309	CLA	CHC-C1C	2.70	1.41	1.35
22	2	304	CLA	CHC-C1C	2.70	1.41	1.35
22	B	839	CLA	CHC-C1C	2.70	1.41	1.35
22	3	310	CLA	CHC-C1C	2.70	1.41	1.35
22	8	313	CLA	CHC-C1C	2.70	1.41	1.35
22	4	811	CLA	CHC-C1C	2.70	1.41	1.35
22	2	301	CLA	CHC-C1C	2.70	1.41	1.35
22	3	319	CLA	CHC-C1C	2.70	1.41	1.35
24	A	856	BCR	C7-C6	2.70	1.54	1.45
22	8	308	CLA	CHC-C1C	2.70	1.41	1.35
22	2	302	CLA	CHC-C1C	2.70	1.41	1.35
22	B	819	CLA	CHC-C1C	2.70	1.41	1.35
24	4	804	BCR	C7-C6	2.70	1.54	1.45
22	B	833	CLA	CHC-C1C	2.70	1.41	1.35
22	3	322	CLA	CHC-C1C	2.70	1.41	1.35
22	5	308	CLA	CHC-C1C	2.70	1.41	1.35
38	1	302	LUT	C18-C5	2.70	1.55	1.50
22	Z	303	CLA	CHC-C1C	2.70	1.41	1.35
22	A	815	CLA	CHC-C1C	2.70	1.41	1.35
22	F	304	CLA	CHC-C1C	2.70	1.41	1.35
22	6	319	CLA	CHC-C1C	2.70	1.41	1.35
22	1	306	CLA	CHC-C1C	2.69	1.41	1.35
22	B	821	CLA	CHC-C1C	2.69	1.41	1.35
22	5	307	CLA	CHC-C1C	2.69	1.41	1.35
24	7	303	BCR	C7-C6	2.69	1.54	1.45
24	J	104	BCR	C7-C6	2.69	1.54	1.45
38	9	301	LUT	C18-C5	2.69	1.55	1.50
22	A	855	CLA	CHC-C1C	2.69	1.41	1.35
38	6	303	LUT	C23-C24	-2.69	1.46	1.50
22	A	814	CLA	CHC-C1C	2.69	1.41	1.35

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
22	1	310	CLA	CHC-C1C	2.69	1.41	1.35
38	6	303	LUT	C18-C5	2.69	1.55	1.50
22	7	310	CLA	CHC-C1C	2.69	1.41	1.35
22	4	818	CLA	CHC-C1C	2.69	1.41	1.35
22	B	814	CLA	CHC-C1C	2.69	1.41	1.35
38	Z	301	LUT	C18-C5	2.69	1.55	1.50
24	A	845	BCR	C7-C6	2.69	1.54	1.45
21	A	801	CL0	C1B-CHB	2.69	1.48	1.41
22	B	830	CLA	CHC-C1C	2.69	1.41	1.35
22	4	808	CLA	CHC-C1C	2.69	1.41	1.35
22	6	307	CLA	CHC-C1C	2.69	1.41	1.35
22	B	841	CLA	CHC-C1C	2.69	1.41	1.35
22	5	311	CLA	CHC-C1C	2.69	1.41	1.35
22	A	829	CLA	CHC-C1C	2.69	1.41	1.35
22	8	316	CLA	CHC-C1C	2.69	1.41	1.35
38	4	802	LUT	C18-C5	2.69	1.55	1.50
22	A	839	CLA	CHC-C1C	2.69	1.41	1.35
22	B	813	CLA	CHC-C1C	2.68	1.41	1.35
24	B	845	BCR	C7-C6	2.68	1.54	1.45
22	B	812	CLA	CHC-C1C	2.68	1.41	1.35
22	5	309	CLA	CHC-C1C	2.68	1.41	1.35
24	B	853	BCR	C36-C18	2.68	1.56	1.50
22	7	308	CLA	CHC-C1C	2.68	1.41	1.35
22	B	829	CLA	CHC-C1C	2.68	1.41	1.35
22	5	310	CLA	CHC-C1C	2.68	1.41	1.35
22	A	820	CLA	CHC-C1C	2.68	1.41	1.35
22	A	838	CLA	CHC-C1C	2.68	1.41	1.35
22	B	827	CLA	CHC-C1C	2.68	1.41	1.35
22	7	304	CLA	CHC-C1C	2.68	1.41	1.35
22	7	317	CLA	CHC-C1C	2.68	1.41	1.35
22	6	322	CLA	CHC-C1C	2.68	1.41	1.35
22	L	201	CLA	CHC-C1C	2.68	1.41	1.35
22	6	302	CLA	CHC-C1C	2.68	1.41	1.35
22	B	805	CLA	CHC-C1C	2.68	1.41	1.35
22	A	830	CLA	CHC-C1C	2.68	1.41	1.35
22	8	314	CLA	CHC-C1C	2.68	1.41	1.35
22	6	311	CLA	CHC-C1C	2.68	1.41	1.35
22	F	302	CLA	CHC-C1C	2.68	1.41	1.35
22	9	311	CLA	CHC-C1C	2.68	1.41	1.35
22	A	821	CLA	CHC-C1C	2.68	1.41	1.35
22	7	309	CLA	CHC-C1C	2.68	1.41	1.35
22	3	312	CLA	CHC-C1C	2.67	1.41	1.35

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
24	B	849	BCR	C7-C6	2.67	1.54	1.45
38	7	301	LUT	C18-C5	2.67	1.55	1.50
22	4	809	CLA	CHC-C1C	2.67	1.41	1.35
22	6	312	CLA	CHC-C1C	2.67	1.41	1.35
22	K	204	CLA	CHC-C1C	2.67	1.41	1.35
22	6	314	CLA	CHC-C1C	2.66	1.41	1.35
22	A	822	CLA	CHC-C1C	2.66	1.41	1.35
22	Z	304	CLA	CHC-C1C	2.66	1.41	1.35
22	B	810	CLA	CHC-C1C	2.66	1.41	1.35
24	A	844	BCR	C34-C9	2.66	1.56	1.50
22	A	816	CLA	CHC-C1C	2.66	1.41	1.35
22	6	308	CLA	CHC-C1C	2.66	1.41	1.35
22	5	313	CLA	CHC-C1C	2.66	1.41	1.35
22	1	316	CLA	C3B-C2B	-2.66	1.36	1.40
22	A	827	CLA	CHC-C1C	2.66	1.41	1.35
22	A	840	CLA	CHC-C1C	2.66	1.41	1.35
22	K	203	CLA	CHC-C1C	2.66	1.41	1.35
24	B	846	BCR	C7-C6	2.66	1.54	1.45
38	5	302	LUT	C18-C5	2.66	1.55	1.50
22	5	326	CLA	CHC-C1C	2.65	1.41	1.35
22	A	808	CLA	CHC-C1C	2.65	1.41	1.35
22	A	832	CLA	CHC-C1C	2.65	1.41	1.35
22	Z	315	CLA	CHC-C1C	2.65	1.41	1.35
22	3	314	CLA	CHC-C1C	2.65	1.41	1.35
22	7	312	CLA	CHC-C1C	2.65	1.41	1.35
22	4	812	CLA	CHC-C1C	2.65	1.41	1.35
22	B	806	CLA	CHC-C1C	2.65	1.41	1.35
24	B	847	BCR	C7-C6	2.65	1.54	1.45
38	3	302	LUT	C18-C5	2.64	1.55	1.50
22	5	319	CLA	CHC-C1C	2.64	1.41	1.35
22	A	842	CLA	CHC-C1C	2.64	1.41	1.35
22	1	305	CLA	CHC-C1C	2.64	1.41	1.35
24	A	848	BCR	C7-C6	2.64	1.54	1.45
24	6	305	BCR	C7-C6	2.64	1.54	1.45
38	8	302	LUT	O23-C23	-2.64	1.38	1.43
22	B	834	CLA	CHC-C1C	2.64	1.41	1.35
22	9	303	CLA	C3B-C2B	-2.64	1.36	1.40
22	8	305	CLA	CHC-C1C	2.64	1.41	1.35
22	B	842	CLA	CHC-C1C	2.64	1.41	1.35
21	A	801	CL0	C4B-CHC	2.64	1.48	1.41
22	9	308	CLA	CHC-C1C	2.64	1.41	1.35
22	Z	313	CLA	CHC-C1C	2.64	1.41	1.35

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
22	8	307	CLA	CHC-C1C	2.63	1.41	1.35
22	6	309	CLA	CHC-C1C	2.63	1.41	1.35
24	B	853	BCR	C38-C26	2.63	1.55	1.50
22	B	811	CLA	CHC-C1C	2.63	1.41	1.35
24	G	203	BCR	C7-C6	2.63	1.54	1.45
24	I	4001	BCR	C7-C6	2.63	1.54	1.45
22	5	301	CLA	CHC-C1C	2.63	1.41	1.35
38	6	304	LUT	C18-C5	2.63	1.55	1.50
38	8	302	LUT	C18-C5	2.63	1.55	1.50
24	3	307	BCR	C7-C6	2.62	1.54	1.45
22	3	315	CLA	C3B-C2B	-2.62	1.36	1.40
38	4	802	LUT	O23-C23	-2.62	1.38	1.43
24	A	847	BCR	C7-C6	2.61	1.54	1.45
24	L	204	BCR	C7-C6	2.61	1.54	1.45
24	6	306	BCR	C7-C6	2.61	1.54	1.45
22	3	315	CLA	CHC-C1C	2.61	1.41	1.35
24	8	304	BCR	C7-C6	2.61	1.54	1.45
24	A	846	BCR	C7-C6	2.61	1.54	1.45
22	9	303	CLA	CHC-C1C	2.61	1.41	1.35
22	1	313	CLA	CHC-C1C	2.60	1.41	1.35
22	4	807	CLA	CHC-C1C	2.60	1.41	1.35
22	G	201	CLA	CHC-C1C	2.60	1.41	1.35
22	7	306	CLA	CHC-C1C	2.60	1.41	1.35
22	A	813	CLA	CHC-C1C	2.60	1.41	1.35
22	9	304	CLA	CHC-C1C	2.59	1.41	1.35
22	K	204	CLA	C3B-C2B	-2.59	1.36	1.40
24	A	844	BCR	C7-C6	2.59	1.54	1.45
24	3	304	BCR	C7-C6	2.59	1.54	1.45
38	1	303	LUT	C23-C24	-2.59	1.46	1.50
38	9	301	LUT	O23-C23	-2.59	1.38	1.43
24	5	304	BCR	C7-C6	2.58	1.54	1.45
41	5	325	3PH	O21-C2	-2.58	1.40	1.46
38	7	302	LUT	C18-C5	2.58	1.55	1.50
24	K	206	BCR	C7-C6	2.58	1.54	1.45
38	3	303	LUT	C18-C5	2.58	1.55	1.50
22	Z	305	CLA	CHC-C1C	2.57	1.41	1.35
38	5	303	LUT	C18-C5	2.57	1.55	1.50
35	J	102	T7X	O16-C8	-2.56	1.40	1.46
38	4	803	LUT	C18-C5	2.56	1.55	1.50
38	5	303	LUT	O23-C23	-2.55	1.39	1.43
38	1	303	LUT	C18-C5	2.55	1.55	1.50
38	9	302	LUT	C18-C5	2.55	1.55	1.50

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
38	Z	301	LUT	O23-C23	-2.54	1.39	1.43
22	A	812	CLA	C3B-C2B	-2.54	1.36	1.40
38	3	303	LUT	O23-C23	-2.53	1.39	1.43
22	B	815	CLA	CHC-C1C	2.52	1.41	1.35
22	5	313	CLA	C3B-C2B	-2.52	1.36	1.40
38	7	301	LUT	O23-C23	-2.52	1.39	1.43
38	8	303	LUT	C18-C5	2.51	1.55	1.50
22	F	304	CLA	C3B-C2B	-2.51	1.36	1.40
22	A	833	CLA	C3B-C2B	-2.50	1.36	1.40
38	Z	302	LUT	C18-C5	2.50	1.55	1.50
22	6	302	CLA	C3B-C2B	-2.50	1.36	1.40
22	4	805	CLA	CHC-C1C	2.49	1.41	1.35
22	6	301	CLA	C3B-C2B	-2.49	1.36	1.40
38	3	302	LUT	O23-C23	-2.49	1.39	1.43
22	5	301	CLA	C3B-C2B	-2.49	1.36	1.40
22	5	315	CLA	C3B-C2B	-2.49	1.36	1.40
41	7	319	3PH	O21-C2	-2.48	1.40	1.46
38	Z	302	LUT	O23-C23	-2.48	1.39	1.43
21	A	801	CL0	C1D-C2D	2.48	1.50	1.45
38	6	303	LUT	O23-C23	-2.48	1.39	1.43
38	4	803	LUT	O23-C23	-2.48	1.39	1.43
22	1	305	CLA	C3B-C2B	-2.47	1.36	1.40
22	6	310	CLA	C3B-C2B	-2.47	1.36	1.40
41	8	320	3PH	O21-C2	-2.47	1.40	1.46
38	1	302	LUT	O23-C23	-2.47	1.39	1.43
22	6	314	CLA	C3B-C2B	-2.47	1.36	1.40
38	6	304	LUT	O23-C23	-2.47	1.39	1.43
22	A	803	CLA	C3B-C2B	-2.47	1.36	1.40
38	1	303	LUT	O23-C23	-2.46	1.39	1.43
22	B	821	CLA	C3B-C2B	-2.46	1.37	1.40
22	4	811	CLA	C3B-C2B	-2.46	1.37	1.40
22	9	311	CLA	C3B-C2B	-2.46	1.37	1.40
22	B	805	CLA	C3B-C2B	-2.45	1.37	1.40
22	5	319	CLA	C3B-C2B	-2.45	1.37	1.40
22	6	313	CLA	C3B-C2B	-2.45	1.37	1.40
22	1	308	CLA	C1C-C2C	2.45	1.49	1.44
41	6	324	3PH	O21-C2	-2.44	1.40	1.46
22	9	310	CLA	CHC-C1C	2.44	1.41	1.35
22	8	311	CLA	C3B-C2B	-2.44	1.37	1.40
24	7	303	BCR	C36-C18	2.44	1.55	1.50
22	Z	304	CLA	C3B-C2B	-2.44	1.37	1.40
38	5	302	LUT	O23-C23	-2.43	1.39	1.43

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
22	5	320	CLA	C4B-NB	-2.43	1.33	1.35
22	A	837	CLA	C3B-C2B	-2.43	1.37	1.40
41	6	324	3PH	O31-C31	2.43	1.40	1.33
38	8	303	LUT	O23-C23	-2.43	1.39	1.43
38	9	302	LUT	O23-C23	-2.43	1.39	1.43
38	Z	301	LUT	C26-C27	2.43	1.53	1.50
22	A	806	CLA	C3B-C2B	-2.43	1.37	1.40
24	F	303	BCR	C36-C18	2.43	1.55	1.50
22	3	316	CLA	C3B-C2B	-2.43	1.37	1.40
24	5	305	BCR	C36-C18	2.43	1.55	1.50
22	B	811	CLA	C3B-C2B	-2.43	1.37	1.40
22	5	314	CLA	C3B-C2B	-2.43	1.37	1.40
24	J	104	BCR	C36-C18	2.42	1.55	1.50
22	A	808	CLA	C3B-C2B	-2.42	1.37	1.40
22	A	841	CLA	C3B-C2B	-2.42	1.37	1.40
22	Z	316	CLA	C3B-C2B	-2.42	1.37	1.40
38	8	302	LUT	C30-C29	-2.42	1.32	1.35
21	A	801	CL0	C1C-NC	-2.42	1.34	1.37
22	B	819	CLA	C3B-C2B	-2.42	1.37	1.40
22	5	323	CLA	C3B-C2B	-2.42	1.37	1.40
32	F	306	RRX	C34-C9	2.42	1.55	1.50
22	3	314	CLA	C3B-C2B	-2.41	1.37	1.40
22	L	201	CLA	C3B-C2B	-2.41	1.37	1.40
36	J	105	C7Z	C20-C13	2.41	1.55	1.50
22	B	814	CLA	C3B-C2B	-2.41	1.37	1.40
22	9	303	CLA	C4B-NB	-2.41	1.33	1.35
22	7	317	CLA	C3B-C2B	-2.41	1.37	1.40
22	Z	309	CLA	C3B-C2B	-2.40	1.37	1.40
24	3	307	BCR	C36-C18	2.40	1.55	1.50
22	B	828	CLA	C3B-C2B	-2.40	1.37	1.40
22	B	830	CLA	C3B-C2B	-2.40	1.37	1.40
22	2	302	CLA	C3B-C2B	-2.40	1.37	1.40
24	B	853	BCR	C39-C30	-2.40	1.49	1.53
22	Z	310	CLA	C3B-C2B	-2.40	1.37	1.40
22	8	316	CLA	C3B-C2B	-2.40	1.37	1.40
38	7	302	LUT	O23-C23	-2.39	1.39	1.43
24	B	849	BCR	C36-C18	2.39	1.55	1.50
24	G	203	BCR	C36-C18	2.39	1.55	1.50
22	8	306	CLA	C3B-C2B	-2.39	1.37	1.40
41	8	320	3PH	O31-C31	2.39	1.40	1.33
22	B	823	CLA	C3B-C2B	-2.39	1.37	1.40
36	5	306	C7Z	C20-C13	2.39	1.55	1.50

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
22	A	809	CLA	C3B-C2B	-2.39	1.37	1.40
24	A	848	BCR	C36-C18	2.39	1.55	1.50
22	1	310	CLA	C3B-C2B	-2.39	1.37	1.40
22	3	312	CLA	C3B-C2B	-2.39	1.37	1.40
22	A	829	CLA	C3B-C2B	-2.39	1.37	1.40
22	7	306	CLA	C3B-C2B	-2.39	1.37	1.40
22	6	319	CLA	C3B-C2B	-2.39	1.37	1.40
22	4	818	CLA	C3B-C2B	-2.38	1.37	1.40
22	A	824	CLA	C3B-C2B	-2.38	1.37	1.40
22	5	326	CLA	C3B-C2B	-2.38	1.37	1.40
22	1	311	CLA	CHC-C1C	2.38	1.41	1.35
36	5	306	C7Z	C38-C25	2.38	1.54	1.50
24	A	856	BCR	C36-C18	2.38	1.55	1.50
38	1	302	LUT	C26-C27	2.38	1.53	1.50
24	I	4001	BCR	C36-C18	2.38	1.55	1.50
32	F	306	RRX	C17-C18	-2.38	1.32	1.35
22	B	813	CLA	C3B-C2B	-2.38	1.37	1.40
41	5	325	3PH	O31-C31	2.38	1.40	1.33
24	A	847	BCR	C36-C18	2.38	1.55	1.50
22	A	807	CLA	C3B-C2B	-2.38	1.37	1.40
24	5	304	BCR	C36-C18	2.38	1.55	1.50
22	4	806	CLA	C3B-C2B	-2.38	1.37	1.40
22	B	820	CLA	C3B-C2B	-2.38	1.37	1.40
22	A	820	CLA	C3B-C2B	-2.37	1.37	1.40
24	3	306	BCR	C36-C18	2.37	1.55	1.50
22	9	305	CLA	C1A-CHA	2.37	1.52	1.43
24	3	305	BCR	C36-C18	2.37	1.55	1.50
24	8	304	BCR	C36-C18	2.37	1.55	1.50
22	7	304	CLA	C3B-C2B	-2.37	1.37	1.40
22	4	809	CLA	C3B-C2B	-2.37	1.37	1.40
24	6	305	BCR	C36-C18	2.37	1.55	1.50
22	A	823	CLA	C3B-C2B	-2.37	1.37	1.40
35	J	102	T7X	O18-C11	2.37	1.40	1.33
22	3	320	CLA	CHC-C1C	2.37	1.41	1.35
24	3	307	BCR	C39-C30	-2.36	1.49	1.53
24	B	847	BCR	C36-C18	2.36	1.55	1.50
24	B	845	BCR	C36-C18	2.36	1.55	1.50
22	7	312	CLA	C3B-C2B	-2.36	1.37	1.40
22	B	818	CLA	C3B-C2B	-2.36	1.37	1.40
22	Z	314	CLA	C3B-C2B	-2.36	1.37	1.40
24	K	206	BCR	C36-C18	2.36	1.55	1.50
24	4	804	BCR	C36-C18	2.35	1.55	1.50

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
36	J	105	C7Z	C40-C33	2.35	1.55	1.50
22	1	316	CLA	C4B-NB	-2.35	1.33	1.35
24	B	848	BCR	C36-C18	2.35	1.55	1.50
22	2	304	CLA	C3B-C2B	-2.35	1.37	1.40
24	B	846	BCR	C36-C18	2.35	1.55	1.50
22	K	205	CLA	C4B-NB	-2.35	1.33	1.35
22	2	301	CLA	C3B-C2B	-2.34	1.37	1.40
22	7	311	CLA	C3B-C2B	-2.34	1.37	1.40
36	J	105	C7Z	C38-C25	2.34	1.54	1.50
22	5	309	CLA	C3B-C2B	-2.34	1.37	1.40
24	6	306	BCR	C36-C18	2.33	1.55	1.50
24	L	204	BCR	C36-C18	2.33	1.55	1.50
38	9	301	LUT	C30-C29	-2.33	1.32	1.35
24	8	304	BCR	C39-C30	-2.33	1.49	1.53
22	G	201	CLA	C3B-C2B	-2.33	1.37	1.40
22	3	322	CLA	C3B-C2B	-2.32	1.37	1.40
22	B	841	CLA	C3B-C2B	-2.32	1.37	1.40
38	7	301	LUT	C30-C29	-2.32	1.32	1.35
22	9	313	CLA	C3B-C2B	-2.32	1.37	1.40
24	A	844	BCR	C36-C18	2.32	1.55	1.50
24	A	846	BCR	C36-C18	2.32	1.55	1.50
22	1	308	CLA	C3B-C2B	-2.32	1.37	1.40
22	A	811	CLA	C3B-C2B	-2.32	1.37	1.40
38	6	303	LUT	C30-C29	-2.31	1.32	1.35
22	F	301	CLA	C3B-C2B	-2.31	1.37	1.40
22	F	302	CLA	C3B-C2B	-2.31	1.37	1.40
22	7	310	CLA	C3B-C2B	-2.31	1.37	1.40
22	A	816	CLA	C3B-C2B	-2.31	1.37	1.40
22	Z	315	CLA	C3B-C2B	-2.31	1.37	1.40
38	3	302	LUT	C34-C33	-2.30	1.32	1.35
24	3	306	BCR	C39-C30	-2.30	1.49	1.53
36	5	306	C7Z	C40-C33	2.30	1.55	1.50
39	5	316	CHL	C3B-C2B	-2.30	1.37	1.40
41	7	319	3PH	O31-C31	2.29	1.40	1.33
22	7	305	CLA	C3B-C2B	-2.29	1.37	1.40
39	8	301	CHL	C3B-C2B	-2.29	1.37	1.40
22	1	316	CLA	CHC-C1C	2.29	1.40	1.35
22	A	855	CLA	C3B-C2B	-2.29	1.37	1.40
38	Z	302	LUT	C26-C27	2.29	1.53	1.50
24	3	304	BCR	C36-C18	2.29	1.55	1.50
22	Z	305	CLA	C3B-C2B	-2.28	1.37	1.40
22	A	802	CLA	C3B-C2B	-2.28	1.37	1.40

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
22	9	309	CLA	CHC-C1C	2.28	1.40	1.35
24	A	845	BCR	C36-C18	2.28	1.55	1.50
38	3	303	LUT	C26-C27	2.27	1.53	1.50
22	3	315	CLA	C3D-C4D	-2.27	1.39	1.44
38	5	302	LUT	C30-C29	-2.27	1.32	1.35
36	J	105	C7Z	C19-C9	2.26	1.55	1.50
22	9	305	CLA	CHC-C1C	2.26	1.40	1.35
24	A	856	BCR	C39-C30	-2.25	1.49	1.53
22	4	812	CLA	C3B-C2B	-2.25	1.37	1.40
38	4	802	LUT	C30-C29	-2.25	1.32	1.35
22	9	303	CLA	C3D-C4D	-2.25	1.39	1.44
22	Z	306	CLA	C1C-C2C	2.25	1.48	1.44
22	B	824	CLA	C1C-C2C	2.25	1.48	1.44
22	B	806	CLA	C3B-C2B	-2.25	1.37	1.40
38	8	303	LUT	C26-C27	2.25	1.53	1.50
38	6	304	LUT	C26-C27	2.24	1.53	1.50
22	K	205	CLA	C1A-CHA	2.24	1.52	1.43
36	5	306	C7Z	C19-C9	2.24	1.55	1.50
22	A	827	CLA	C3B-C2B	-2.23	1.37	1.40
39	9	312	CHL	C3B-C2B	-2.23	1.37	1.40
24	B	844	BCR	C39-C30	-2.23	1.49	1.53
22	B	829	CLA	C3B-C2B	-2.23	1.37	1.40
22	4	817	CLA	C3B-C2B	-2.23	1.37	1.40
38	9	302	LUT	C26-C27	2.23	1.53	1.50
38	1	302	LUT	C30-C29	-2.22	1.32	1.35
22	A	826	CLA	C1C-C2C	2.22	1.48	1.44
38	4	803	LUT	C30-C29	-2.22	1.32	1.35
38	7	302	LUT	C26-C27	2.22	1.53	1.50
24	3	304	BCR	C39-C30	-2.22	1.49	1.53
24	K	206	BCR	C39-C30	-2.22	1.49	1.53
24	6	305	BCR	C39-C30	-2.21	1.49	1.53
38	5	303	LUT	C26-C27	2.21	1.53	1.50
22	3	320	CLA	C3D-C4D	-2.21	1.39	1.44
22	4	806	CLA	C1C-C2C	2.21	1.48	1.44
38	8	302	LUT	C26-C27	2.21	1.53	1.50
22	7	305	CLA	C1C-C2C	2.21	1.48	1.44
22	7	307	CLA	C1C-C2C	2.21	1.48	1.44
22	6	310	CLA	C1C-C2C	2.21	1.48	1.44
22	6	313	CLA	C1C-C2C	2.21	1.48	1.44
24	A	847	BCR	C39-C30	-2.21	1.49	1.53
24	I	4001	BCR	C38-C26	2.21	1.54	1.50
38	7	301	LUT	C26-C27	2.21	1.53	1.50

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
24	F	303	BCR	C39-C30	-2.20	1.49	1.53
38	9	302	LUT	C30-C29	-2.20	1.32	1.35
22	B	807	CLA	C1C-C2C	2.20	1.48	1.44
38	3	302	LUT	C30-C29	-2.20	1.32	1.35
22	K	205	CLA	CHC-C1C	2.20	1.40	1.35
22	4	815	CLA	C1C-C2C	2.20	1.48	1.44
22	A	834	CLA	C3B-C2B	-2.20	1.37	1.40
22	F	302	CLA	C1C-C2C	2.20	1.48	1.44
22	1	316	CLA	C3D-C4D	-2.20	1.39	1.44
27	A	852	DGA	OG2-CG2	-2.19	1.41	1.46
24	5	305	BCR	C39-C30	-2.19	1.49	1.53
22	B	828	CLA	C1C-C2C	2.19	1.48	1.44
24	7	303	BCR	C39-C30	-2.19	1.49	1.53
24	L	204	BCR	C39-C30	-2.19	1.49	1.53
22	3	319	CLA	C3B-C2B	-2.19	1.37	1.40
22	K	205	CLA	C3D-C4D	-2.19	1.39	1.44
22	1	313	CLA	C1C-C2C	2.19	1.48	1.44
24	5	305	BCR	C38-C26	2.19	1.54	1.50
22	B	808	CLA	C1C-C2C	2.19	1.48	1.44
22	F	301	CLA	C1C-C2C	2.18	1.48	1.44
22	B	810	CLA	C3B-C2B	-2.18	1.37	1.40
24	3	305	BCR	C39-C30	-2.18	1.49	1.53
22	A	828	CLA	C3B-C2B	-2.18	1.37	1.40
41	6	324	3PH	O21-C21	2.18	1.40	1.34
22	4	811	CLA	C1C-C2C	2.18	1.48	1.44
22	5	322	CLA	C1C-C2C	2.18	1.48	1.44
22	A	825	CLA	C3B-C2B	-2.18	1.37	1.40
39	6	320	CHL	C3B-C2B	-2.18	1.37	1.40
38	4	802	LUT	C26-C27	2.18	1.53	1.50
22	6	312	CLA	C3B-C2B	-2.18	1.37	1.40
38	3	303	LUT	C30-C29	-2.18	1.32	1.35
22	B	801	CLA	C3B-C2B	-2.18	1.37	1.40
41	7	319	3PH	O31-C3	-2.18	1.40	1.45
22	8	313	CLA	C3B-C2B	-2.18	1.37	1.40
22	J	103	CLA	C1C-C2C	2.18	1.48	1.44
22	9	307	CLA	C1C-C2C	2.18	1.48	1.44
24	4	804	BCR	C39-C30	-2.18	1.49	1.53
22	3	310	CLA	C3B-C2B	-2.18	1.37	1.40
22	8	307	CLA	C3B-C2B	-2.17	1.37	1.40
22	6	311	CLA	C3B-C2B	-2.17	1.37	1.40
22	5	310	CLA	C1C-C2C	2.17	1.48	1.44
35	J	102	T7X	O18-C9	-2.17	1.40	1.45

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
24	B	845	BCR	C39-C30	-2.17	1.49	1.53
38	4	803	LUT	C26-C27	2.17	1.53	1.50
38	8	303	LUT	C30-C29	-2.17	1.32	1.35
22	7	322	CLA	C1C-C2C	2.17	1.48	1.44
22	7	323	CLA	C3B-C2B	-2.17	1.37	1.40
22	1	315	CLA	C1C-C2C	2.17	1.48	1.44
22	Z	308	CLA	C1C-C2C	2.17	1.48	1.44
24	A	846	BCR	C39-C30	-2.17	1.49	1.53
22	6	319	CLA	C1C-C2C	2.17	1.48	1.44
22	B	813	CLA	C1C-C2C	2.16	1.48	1.44
22	5	320	CLA	CHC-C1C	2.16	1.40	1.35
22	7	316	CLA	C3B-C2B	-2.16	1.37	1.40
22	A	823	CLA	C1C-C2C	2.16	1.48	1.44
22	7	311	CLA	C1C-C2C	2.16	1.48	1.44
22	3	318	CLA	C1C-C2C	2.16	1.48	1.44
41	6	324	3PH	O31-C3	-2.16	1.40	1.45
22	A	854	CLA	C3B-C2B	-2.16	1.37	1.40
22	Z	314	CLA	C1C-C2C	2.16	1.48	1.44
22	3	318	CLA	C3B-C2B	-2.16	1.37	1.40
22	A	833	CLA	C1C-C2C	2.16	1.48	1.44
22	A	841	CLA	C1C-C2C	2.16	1.48	1.44
22	7	315	CLA	C1C-C2C	2.16	1.48	1.44
22	B	825	CLA	C1C-C2C	2.16	1.48	1.44
22	L	202	CLA	C3B-C2B	-2.16	1.37	1.40
22	Z	306	CLA	C3B-C2B	-2.16	1.37	1.40
22	B	834	CLA	C3B-C2B	-2.16	1.37	1.40
22	B	842	CLA	C3B-C2B	-2.16	1.37	1.40
22	B	805	CLA	C1C-C2C	2.16	1.48	1.44
22	B	831	CLA	C1C-C2C	2.16	1.48	1.44
22	8	311	CLA	C1C-C2C	2.16	1.48	1.44
22	9	308	CLA	C3B-C2B	-2.16	1.37	1.40
38	7	302	LUT	C30-C29	-2.16	1.32	1.35
22	Z	313	CLA	C1C-C2C	2.16	1.48	1.44
22	A	834	CLA	C1C-C2C	2.16	1.48	1.44
22	4	808	CLA	C3B-C2B	-2.16	1.37	1.40
24	B	846	BCR	C39-C30	-2.15	1.49	1.53
22	1	307	CLA	C1C-C2C	2.15	1.48	1.44
22	A	831	CLA	C3B-C2B	-2.15	1.37	1.40
24	6	306	BCR	C39-C30	-2.15	1.49	1.53
22	L	203	CLA	C1C-C2C	2.15	1.48	1.44
22	8	309	CLA	C3B-C2B	-2.15	1.37	1.40
22	7	314	CLA	C3B-C2B	-2.15	1.37	1.40

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
24	8	304	BCR	C38-C26	2.15	1.54	1.50
22	7	324	CLA	C1C-C2C	2.15	1.48	1.44
22	A	839	CLA	C3B-C2B	-2.15	1.37	1.40
24	A	846	BCR	C38-C26	2.15	1.54	1.50
22	B	832	CLA	C3B-C2B	-2.15	1.37	1.40
39	9	314	CHL	C3B-C2B	-2.15	1.37	1.40
22	A	812	CLA	C1C-C2C	2.15	1.48	1.44
24	K	206	BCR	C38-C26	2.15	1.54	1.50
41	5	325	3PH	O31-C3	-2.15	1.40	1.45
22	A	824	CLA	C1C-C2C	2.15	1.48	1.44
22	A	842	CLA	C3B-C2B	-2.15	1.37	1.40
39	Z	311	CHL	C3B-C2B	-2.15	1.37	1.40
22	A	818	CLA	C1C-C2C	2.15	1.48	1.44
24	B	847	BCR	C39-C30	-2.15	1.49	1.53
32	F	306	RRX	C33-C5	2.14	1.54	1.50
22	A	825	CLA	C1C-C2C	2.14	1.48	1.44
22	B	829	CLA	C1C-C2C	2.14	1.48	1.44
22	B	824	CLA	C3B-C2B	-2.14	1.37	1.40
24	A	848	BCR	C38-C26	2.14	1.54	1.50
41	8	320	3PH	O21-C21	2.14	1.40	1.34
22	7	310	CLA	C1C-C2C	2.14	1.48	1.44
22	6	317	CLA	C1C-C2C	2.14	1.48	1.44
41	7	319	3PH	O21-C21	2.14	1.40	1.34
22	8	306	CLA	C1C-C2C	2.14	1.48	1.44
22	5	314	CLA	C1C-C2C	2.14	1.48	1.44
22	1	314	CLA	C3B-C2B	-2.14	1.37	1.40
22	B	837	CLA	C1C-C2C	2.14	1.48	1.44
38	6	304	LUT	C30-C29	-2.14	1.32	1.35
24	B	848	BCR	C39-C30	-2.14	1.49	1.53
22	B	826	CLA	C1C-C2C	2.14	1.48	1.44
22	Z	313	CLA	C3B-C2B	-2.14	1.37	1.40
22	B	816	CLA	C3B-C2B	-2.14	1.37	1.40
22	B	801	CLA	C1C-C2C	2.14	1.48	1.44
24	B	853	BCR	C37-C22	2.14	1.55	1.50
22	A	835	CLA	C1C-C2C	2.14	1.48	1.44
22	B	822	CLA	C1C-C2C	2.14	1.48	1.44
22	B	823	CLA	C1C-C2C	2.14	1.48	1.44
22	3	319	CLA	C1C-C2C	2.14	1.48	1.44
22	8	316	CLA	C1C-C2C	2.14	1.48	1.44
24	J	104	BCR	C39-C30	-2.14	1.49	1.53
22	B	838	CLA	C1C-C2C	2.14	1.48	1.44
22	4	817	CLA	C1C-C2C	2.14	1.48	1.44

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
22	3	315	CLA	C4B-NB	-2.14	1.33	1.35
24	3	306	BCR	C38-C26	2.14	1.54	1.50
22	7	315	CLA	C3B-C2B	-2.14	1.37	1.40
22	9	311	CLA	C1C-C2C	2.14	1.48	1.44
22	B	833	CLA	C1C-C2C	2.14	1.48	1.44
22	B	807	CLA	C3B-C2B	-2.14	1.37	1.40
22	A	829	CLA	C1C-C2C	2.14	1.48	1.44
22	B	820	CLA	C1C-C2C	2.14	1.48	1.44
22	4	808	CLA	C1C-C2C	2.14	1.48	1.44
22	A	819	CLA	C1C-C2C	2.13	1.48	1.44
24	B	847	BCR	C38-C26	2.13	1.54	1.50
38	1	303	LUT	C26-C27	2.13	1.53	1.50
22	A	840	CLA	C3B-C2B	-2.13	1.37	1.40
24	A	848	BCR	C39-C30	-2.13	1.49	1.53
22	3	311	CLA	C1C-C2C	2.13	1.48	1.44
22	1	304	CLA	C1C-C2C	2.13	1.48	1.44
38	9	301	LUT	C26-C27	2.13	1.53	1.50
24	B	849	BCR	C39-C30	-2.13	1.49	1.53
22	Z	309	CLA	C1C-C2C	2.13	1.48	1.44
24	A	844	BCR	C39-C30	-2.13	1.49	1.53
22	7	308	CLA	C1C-C2C	2.13	1.48	1.44
24	G	203	BCR	C39-C30	-2.13	1.49	1.53
41	8	320	3PH	O31-C3	-2.13	1.40	1.45
38	5	302	LUT	C26-C27	2.13	1.53	1.50
22	B	821	CLA	C1C-C2C	2.13	1.48	1.44
22	5	311	CLA	C1C-C2C	2.13	1.48	1.44
24	B	845	BCR	C38-C26	2.13	1.54	1.50
22	3	314	CLA	C1C-C2C	2.13	1.48	1.44
22	B	816	CLA	C1C-C2C	2.13	1.48	1.44
22	B	836	CLA	C1C-C2C	2.13	1.48	1.44
22	8	309	CLA	C1C-C2C	2.13	1.48	1.44
24	B	849	BCR	C37-C22	2.13	1.55	1.50
22	A	830	CLA	C1C-C2C	2.13	1.48	1.44
22	8	307	CLA	C1C-C2C	2.12	1.48	1.44
24	B	844	BCR	C38-C26	2.12	1.54	1.50
22	A	819	CLA	C3B-C2B	-2.12	1.37	1.40
24	A	845	BCR	C39-C30	-2.12	1.49	1.53
38	4	802	LUT	C34-C33	-2.12	1.33	1.35
22	A	809	CLA	C1C-C2C	2.12	1.48	1.44
24	B	848	BCR	C38-C26	2.12	1.54	1.50
22	3	316	CLA	C1C-C2C	2.12	1.48	1.44
22	Z	304	CLA	C1C-C2C	2.12	1.48	1.44

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
22	K	202	CLA	C1C-C2C	2.12	1.48	1.44
22	8	308	CLA	C1C-C2C	2.12	1.48	1.44
22	1	305	CLA	C3D-C4D	-2.12	1.39	1.44
22	Z	303	CLA	C1C-C2C	2.12	1.48	1.44
24	I	4001	BCR	C39-C30	-2.12	1.49	1.53
22	A	831	CLA	C1C-C2C	2.12	1.48	1.44
38	6	303	LUT	C26-C27	2.12	1.53	1.50
22	5	307	CLA	C3B-C2B	-2.12	1.37	1.40
24	6	305	BCR	C38-C26	2.12	1.54	1.50
22	A	806	CLA	C1C-C2C	2.12	1.48	1.44
38	4	803	LUT	C34-C33	-2.12	1.33	1.35
22	B	811	CLA	C1C-C2C	2.12	1.48	1.44
22	A	810	CLA	C1C-C2C	2.12	1.48	1.44
22	B	835	CLA	C1C-C2C	2.12	1.48	1.44
22	3	322	CLA	C1C-C2C	2.12	1.48	1.44
22	A	820	CLA	C1C-C2C	2.12	1.48	1.44
22	1	309	CLA	C1C-C2C	2.12	1.48	1.44
22	2	304	CLA	C1C-C2C	2.12	1.48	1.44
22	6	321	CLA	C3B-C2B	-2.12	1.37	1.40
22	B	815	CLA	C3B-C2B	-2.12	1.37	1.40
22	B	806	CLA	C1C-C2C	2.12	1.48	1.44
22	A	811	CLA	C1C-C2C	2.11	1.48	1.44
22	F	305	CLA	C1C-C2C	2.11	1.48	1.44
22	5	315	CLA	C1C-C2C	2.11	1.48	1.44
24	A	844	BCR	C38-C26	2.11	1.54	1.50
38	5	302	LUT	C34-C33	-2.11	1.33	1.35
38	9	302	LUT	C34-C33	-2.11	1.33	1.35
22	A	836	CLA	C1C-C2C	2.11	1.48	1.44
22	9	305	CLA	CAA-C2A	-2.11	1.50	1.54
22	3	313	CLA	C1C-C2C	2.11	1.48	1.44
38	9	301	LUT	C34-C33	-2.11	1.33	1.35
22	A	813	CLA	C3B-C2B	-2.11	1.37	1.40
22	G	202	CLA	C3B-C2B	-2.11	1.37	1.40
41	5	325	3PH	O21-C21	2.11	1.40	1.34
24	B	846	BCR	C38-C26	2.11	1.54	1.50
22	4	805	CLA	C1C-C2C	2.11	1.48	1.44
22	1	311	CLA	C3D-C4D	-2.11	1.39	1.44
24	5	304	BCR	C39-C30	-2.11	1.49	1.53
22	Z	316	CLA	C1C-C2C	2.11	1.48	1.44
24	B	844	BCR	C37-C22	2.11	1.55	1.50
22	A	808	CLA	C1C-C2C	2.11	1.48	1.44
22	A	855	CLA	C1C-C2C	2.11	1.48	1.44

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
22	6	322	CLA	C1C-C2C	2.11	1.48	1.44
22	A	807	CLA	C1C-C2C	2.11	1.48	1.44
22	Z	307	CLA	C1C-C2C	2.11	1.48	1.44
22	8	310	CLA	C3B-C2B	-2.11	1.37	1.40
24	3	306	BCR	C37-C22	2.10	1.55	1.50
22	Z	307	CLA	C3B-C2B	-2.10	1.37	1.40
38	3	302	LUT	C26-C27	2.10	1.53	1.50
22	8	314	CLA	C3B-C2B	-2.10	1.37	1.40
22	5	312	CLA	C3B-C2B	-2.10	1.37	1.40
22	6	311	CLA	C1C-C2C	2.10	1.48	1.44
22	5	309	CLA	C1C-C2C	2.10	1.48	1.44
22	K	203	CLA	C3B-C2B	-2.10	1.37	1.40
22	K	204	CLA	C1B-NB	2.10	1.37	1.35
22	B	832	CLA	C1C-C2C	2.10	1.48	1.44
24	A	845	BCR	C37-C22	2.10	1.55	1.50
22	8	308	CLA	C3B-C2B	-2.10	1.37	1.40
22	A	822	CLA	C3B-C2B	-2.10	1.37	1.40
22	Z	308	CLA	C3B-C2B	-2.10	1.37	1.40
24	A	847	BCR	C37-C22	2.10	1.55	1.50
22	A	803	CLA	C1C-C2C	2.10	1.48	1.44
22	6	321	CLA	C1C-C2C	2.10	1.48	1.44
24	A	847	BCR	C38-C26	2.10	1.54	1.50
22	Z	310	CLA	C1C-C2C	2.10	1.48	1.44
24	6	306	BCR	C38-C26	2.10	1.54	1.50
22	5	308	CLA	C3B-C2B	-2.10	1.37	1.40
38	7	301	LUT	C34-C33	-2.10	1.33	1.35
22	8	310	CLA	C1C-C2C	2.10	1.48	1.44
24	K	206	BCR	C37-C22	2.10	1.55	1.50
22	B	817	CLA	C1C-C2C	2.10	1.48	1.44
22	L	201	CLA	C1C-C2C	2.10	1.48	1.44
22	7	316	CLA	C1C-C2C	2.10	1.48	1.44
22	B	841	CLA	C1C-C2C	2.09	1.48	1.44
22	3	308	CLA	C1C-C2C	2.09	1.48	1.44
22	A	817	CLA	C3B-C2B	-2.09	1.37	1.40
22	B	822	CLA	C3B-C2B	-2.09	1.37	1.40
22	8	314	CLA	C1C-C2C	2.09	1.48	1.44
22	7	317	CLA	C1C-C2C	2.09	1.48	1.44
22	5	313	CLA	C1A-CHA	2.09	1.51	1.43
24	G	203	BCR	C37-C22	2.09	1.55	1.50
22	5	326	CLA	C1C-C2C	2.09	1.48	1.44
22	B	812	CLA	C3B-C2B	-2.09	1.37	1.40
22	5	313	CLA	C1C-C2C	2.09	1.48	1.44

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
22	Z	315	CLA	C1C-C2C	2.09	1.48	1.44
22	9	306	CLA	C1C-C2C	2.09	1.48	1.44
24	L	204	BCR	C38-C26	2.09	1.54	1.50
22	7	304	CLA	C1C-C2C	2.09	1.48	1.44
22	A	814	CLA	C1C-C2C	2.09	1.48	1.44
22	1	306	CLA	C1C-C2C	2.09	1.48	1.44
22	6	312	CLA	C1C-C2C	2.09	1.48	1.44
22	G	202	CLA	C1C-C2C	2.09	1.48	1.44
38	7	302	LUT	C34-C33	-2.09	1.33	1.35
24	F	303	BCR	C37-C22	2.09	1.55	1.50
24	3	305	BCR	C37-C22	2.09	1.55	1.50
22	1	306	CLA	C3B-C2B	-2.08	1.37	1.40
22	3	308	CLA	C3B-C2B	-2.08	1.37	1.40
22	6	322	CLA	C3B-C2B	-2.08	1.37	1.40
22	B	827	CLA	C1C-C2C	2.08	1.48	1.44
22	1	307	CLA	C3B-C2B	-2.08	1.37	1.40
22	B	830	CLA	C1C-C2C	2.08	1.48	1.44
22	A	830	CLA	C3B-C2B	-2.08	1.37	1.40
22	B	809	CLA	C1C-C2C	2.08	1.48	1.44
22	L	202	CLA	C1C-C2C	2.08	1.48	1.44
24	3	307	BCR	C38-C26	2.08	1.54	1.50
22	4	805	CLA	MG-NC	2.08	2.11	2.06
24	3	304	BCR	C37-C22	2.08	1.55	1.50
24	B	845	BCR	C37-C22	2.08	1.55	1.50
22	9	309	CLA	C1A-CHA	2.08	1.51	1.43
22	A	815	CLA	C1C-C2C	2.08	1.48	1.44
22	7	307	CLA	C3D-C4D	-2.08	1.39	1.44
22	B	839	CLA	C3B-C2B	-2.08	1.37	1.40
22	5	320	CLA	C1A-CHA	2.08	1.51	1.43
22	B	818	CLA	C1C-C2C	2.08	1.48	1.44
24	3	304	BCR	C38-C26	2.08	1.54	1.50
22	A	839	CLA	C1C-C2C	2.08	1.48	1.44
22	B	819	CLA	C1C-C2C	2.08	1.48	1.44
22	7	309	CLA	C1C-C2C	2.08	1.48	1.44
24	G	203	BCR	C38-C26	2.08	1.54	1.50
22	6	309	CLA	C1A-CHA	2.08	1.51	1.43
22	4	807	CLA	C1C-C2C	2.08	1.48	1.44
22	A	827	CLA	C1C-C2C	2.08	1.48	1.44
22	3	310	CLA	C1C-C2C	2.08	1.48	1.44
22	B	840	CLA	C1C-C2C	2.08	1.48	1.44
38	1	303	LUT	C30-C29	-2.08	1.33	1.35
24	J	104	BCR	C38-C26	2.07	1.54	1.50

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
22	7	324	CLA	C3B-C2B	-2.07	1.37	1.40
24	A	848	BCR	C37-C22	2.07	1.55	1.50
22	6	307	CLA	C1C-C2C	2.07	1.48	1.44
22	F	304	CLA	C1C-C2C	2.07	1.48	1.44
22	9	306	CLA	C3B-C2B	-2.07	1.37	1.40
22	A	837	CLA	C1C-C2C	2.07	1.48	1.44
22	1	314	CLA	C1C-C2C	2.07	1.48	1.44
22	B	826	CLA	C3B-C2B	-2.07	1.37	1.40
22	4	807	CLA	C3B-C2B	-2.07	1.37	1.40
24	A	846	BCR	C37-C22	2.07	1.55	1.50
22	1	310	CLA	C1C-C2C	2.07	1.48	1.44
22	4	812	CLA	C1C-C2C	2.07	1.48	1.44
22	6	302	CLA	C1C-C2C	2.07	1.48	1.44
22	9	313	CLA	C1C-C2C	2.07	1.48	1.44
22	B	836	CLA	C3B-C2B	-2.07	1.37	1.40
39	6	316	CHL	C3B-C2B	-2.07	1.37	1.40
22	5	312	CLA	C1C-C2C	2.07	1.48	1.44
22	9	304	CLA	C3D-C4D	-2.07	1.39	1.44
22	9	310	CLA	C1A-CHA	2.07	1.51	1.43
22	A	832	CLA	C1C-C2C	2.07	1.48	1.44
22	3	312	CLA	C1C-C2C	2.07	1.48	1.44
22	5	307	CLA	C1C-C2C	2.07	1.48	1.44
24	J	104	BCR	C37-C22	2.07	1.55	1.50
22	6	307	CLA	C3B-C2B	-2.07	1.37	1.40
22	K	205	CLA	C1D-ND	-2.07	1.35	1.37
22	3	310	CLA	C1A-CHA	2.07	1.51	1.43
22	A	838	CLA	C1C-C2C	2.06	1.48	1.44
22	K	203	CLA	C1C-C2C	2.06	1.48	1.44
22	6	308	CLA	C1C-C2C	2.06	1.48	1.44
22	7	308	CLA	C3D-C4D	-2.06	1.39	1.44
22	B	812	CLA	C1C-C2C	2.06	1.48	1.44
22	4	809	CLA	C1C-C2C	2.06	1.48	1.44
22	5	319	CLA	C1C-C2C	2.06	1.48	1.44
22	6	301	CLA	C1C-C2C	2.06	1.48	1.44
24	3	307	BCR	C37-C22	2.06	1.55	1.50
22	A	816	CLA	C1C-C2C	2.06	1.48	1.44
22	7	323	CLA	C1C-C2C	2.06	1.48	1.44
22	8	309	CLA	C3D-C4D	-2.06	1.39	1.44
22	A	854	CLA	C1C-C2C	2.06	1.48	1.44
22	5	323	CLA	C1C-C2C	2.06	1.48	1.44
22	9	307	CLA	C1B-NB	2.06	1.37	1.35
22	A	802	CLA	C1C-C2C	2.06	1.48	1.44

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
38	5	303	LUT	C34-C33	-2.06	1.33	1.35
22	8	307	CLA	C1A-CHA	2.06	1.51	1.43
35	J	102	T7X	O16-C10	2.06	1.40	1.34
24	7	303	BCR	C37-C22	2.06	1.55	1.50
22	A	804	CLA	C1C-C2C	2.06	1.48	1.44
22	5	301	CLA	C1C-C2C	2.06	1.48	1.44
22	7	312	CLA	C1C-C2C	2.06	1.48	1.44
24	B	849	BCR	C38-C26	2.06	1.54	1.50
22	A	810	CLA	C3B-C2B	-2.06	1.37	1.40
22	A	836	CLA	C3B-C2B	-2.06	1.37	1.40
22	F	305	CLA	C3B-C2B	-2.06	1.37	1.40
24	B	846	BCR	C37-C22	2.06	1.55	1.50
22	2	302	CLA	C1C-C2C	2.06	1.48	1.44
22	5	308	CLA	C1C-C2C	2.06	1.48	1.44
22	3	320	CLA	C4B-NB	-2.05	1.33	1.35
24	A	845	BCR	C38-C26	2.05	1.54	1.50
22	9	310	CLA	C3D-C4D	-2.05	1.39	1.44
22	A	804	CLA	C3B-C2B	-2.05	1.37	1.40
22	A	828	CLA	C1C-C2C	2.05	1.48	1.44
22	B	837	CLA	C3B-C2B	-2.05	1.37	1.40
24	6	306	BCR	C37-C22	2.05	1.55	1.50
22	9	305	CLA	C3D-C4D	-2.05	1.39	1.44
22	8	305	CLA	C3B-C2B	-2.05	1.37	1.40
22	5	311	CLA	C3B-C2B	-2.05	1.37	1.40
24	B	848	BCR	C37-C22	2.05	1.55	1.50
24	I	4001	BCR	C37-C22	2.05	1.55	1.50
24	B	847	BCR	C37-C22	2.05	1.55	1.50
22	A	835	CLA	C3B-C2B	-2.05	1.37	1.40
22	7	315	CLA	C1A-CHA	2.05	1.51	1.43
22	B	842	CLA	C1C-C2C	2.05	1.48	1.44
22	A	822	CLA	C1C-C2C	2.05	1.48	1.44
22	A	840	CLA	C1C-C2C	2.05	1.48	1.44
22	5	308	CLA	C3D-C4D	-2.05	1.39	1.44
22	B	823	CLA	C1B-NB	2.05	1.37	1.35
22	8	313	CLA	C1C-C2C	2.05	1.48	1.44
22	3	309	CLA	C1C-C2C	2.05	1.48	1.44
22	7	307	CLA	C3B-C2B	-2.05	1.37	1.40
38	1	303	LUT	C34-C33	-2.05	1.33	1.35
22	A	842	CLA	C1C-C2C	2.05	1.48	1.44
24	5	305	BCR	C37-C22	2.05	1.55	1.50
22	7	308	CLA	C3B-C2B	-2.05	1.37	1.40
22	3	311	CLA	C3B-C2B	-2.05	1.37	1.40

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
22	B	831	CLA	C3D-C4D	-2.04	1.39	1.44
22	A	838	CLA	C3B-C2B	-2.04	1.37	1.40
24	A	856	BCR	C38-C26	2.04	1.54	1.50
22	4	807	CLA	C1A-CHA	2.04	1.51	1.43
24	L	204	BCR	C37-C22	2.04	1.55	1.50
22	K	204	CLA	C1C-C2C	2.04	1.48	1.44
39	3	317	CHL	C3B-C2B	-2.04	1.37	1.40
38	5	303	LUT	C30-C29	-2.04	1.33	1.35
22	4	805	CLA	CHD-C1D	2.04	1.42	1.38
22	1	309	CLA	C3B-C2B	-2.04	1.37	1.40
22	5	309	CLA	C1A-CHA	2.04	1.51	1.43
22	4	818	CLA	C1C-C2C	2.04	1.48	1.44
22	A	805	CLA	C3B-C2B	-2.04	1.37	1.40
38	Z	301	LUT	C28-C27	2.04	1.37	1.32
22	9	307	CLA	C3B-C2B	-2.04	1.37	1.40
22	Z	310	CLA	C1B-NB	2.04	1.37	1.35
22	B	834	CLA	C1C-C2C	2.04	1.48	1.44
22	1	308	CLA	C3D-C4D	-2.04	1.39	1.44
22	9	307	CLA	C3D-C4D	-2.04	1.39	1.44
22	2	301	CLA	C3D-C4D	-2.04	1.39	1.44
22	4	810	CLA	C1C-C2C	2.04	1.48	1.44
22	5	318	CLA	C1C-C2C	2.04	1.48	1.44
22	J	103	CLA	C3B-C2B	-2.04	1.37	1.40
39	7	313	CHL	C3B-C2B	-2.04	1.37	1.40
22	A	823	CLA	C3D-C4D	-2.03	1.39	1.44
22	B	827	CLA	C3B-C2B	-2.03	1.37	1.40
22	A	836	CLA	C3D-C4D	-2.03	1.39	1.44
22	1	311	CLA	C1A-CHA	2.03	1.51	1.43
22	5	310	CLA	C3B-C2B	-2.03	1.37	1.40
22	A	817	CLA	C1B-NB	2.03	1.37	1.35
22	Z	306	CLA	C3D-C4D	-2.03	1.39	1.44
22	3	308	CLA	C3D-C4D	-2.03	1.39	1.44
22	5	322	CLA	C3B-C2B	-2.03	1.37	1.40
22	6	314	CLA	C1C-C2C	2.03	1.48	1.44
24	F	303	BCR	C38-C26	2.03	1.54	1.50
38	8	303	LUT	C34-C33	-2.03	1.33	1.35
22	1	309	CLA	C3D-C4D	-2.03	1.39	1.44
22	B	809	CLA	C3B-C2B	-2.03	1.37	1.40
22	A	816	CLA	C1A-CHA	2.03	1.51	1.43
38	6	304	LUT	C34-C33	-2.03	1.33	1.35
22	A	821	CLA	C1C-C2C	2.03	1.48	1.44
21	A	801	CL0	C4C-C3C	2.03	1.48	1.45

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
22	4	809	CLA	C3D-C4D	-2.03	1.39	1.44
22	A	826	CLA	C3B-C2B	-2.03	1.37	1.40
22	B	814	CLA	C3D-C4D	-2.03	1.39	1.44
22	B	840	CLA	C3B-C2B	-2.03	1.37	1.40
22	B	839	CLA	C3D-C4D	-2.03	1.39	1.44
22	3	309	CLA	C3B-C2B	-2.03	1.37	1.40
22	6	309	CLA	C1C-C2C	2.03	1.48	1.44
22	4	806	CLA	C3D-C4D	-2.03	1.39	1.44
22	6	309	CLA	C3B-C2B	-2.03	1.37	1.40
22	1	316	CLA	C1A-CHA	2.02	1.51	1.43
22	B	809	CLA	C3D-C4D	-2.02	1.39	1.44
22	Z	305	CLA	C1C-C2C	2.02	1.48	1.44
38	Z	302	LUT	C30-C29	-2.02	1.33	1.35
22	A	811	CLA	C3D-C4D	-2.02	1.39	1.44
22	7	309	CLA	C3B-C2B	-2.02	1.37	1.40
22	3	309	CLA	C3D-C4D	-2.02	1.39	1.44
22	B	818	CLA	C3D-C4D	-2.02	1.39	1.44
22	A	817	CLA	C1C-C2C	2.02	1.48	1.44
24	7	303	BCR	C38-C26	2.02	1.54	1.50
22	2	301	CLA	C1B-NB	2.02	1.37	1.35
38	1	302	LUT	C28-C27	2.02	1.37	1.32
22	B	835	CLA	C3D-C4D	-2.02	1.39	1.44
39	4	816	CHL	CHC-C1C	2.02	1.40	1.35
22	A	818	CLA	C3D-C4D	-2.02	1.39	1.44
22	A	842	CLA	C3D-C4D	-2.02	1.39	1.44
22	1	310	CLA	C1A-CHA	2.02	1.51	1.43
22	F	305	CLA	C3D-C4D	-2.01	1.39	1.44
22	1	307	CLA	C3D-C4D	-2.01	1.39	1.44
22	A	822	CLA	C1A-CHA	2.01	1.51	1.43
24	8	304	BCR	C37-C22	2.01	1.55	1.50
38	3	303	LUT	C34-C33	-2.01	1.33	1.35
38	8	302	LUT	C34-C33	-2.01	1.33	1.35
22	6	312	CLA	C1A-CHA	2.01	1.51	1.43
22	B	838	CLA	C3B-C2B	-2.01	1.37	1.40
22	3	312	CLA	C3D-C4D	-2.01	1.39	1.44
22	4	808	CLA	C1B-NB	2.01	1.37	1.35
22	B	839	CLA	C1C-C2C	2.01	1.48	1.44
22	5	311	CLA	C3D-C4D	-2.01	1.39	1.44
22	5	318	CLA	C3B-C2B	-2.01	1.37	1.40
22	A	839	CLA	C3D-C4D	-2.01	1.39	1.44
22	5	307	CLA	C3D-C4D	-2.01	1.39	1.44
22	7	306	CLA	C1A-CHA	2.01	1.51	1.43

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
22	A	805	CLA	C1B-NB	2.01	1.37	1.35
22	6	310	CLA	C3D-C4D	-2.01	1.39	1.44
22	A	805	CLA	C3D-C4D	-2.01	1.39	1.44
22	B	831	CLA	C3B-C2B	-2.01	1.37	1.40
22	1	313	CLA	C3B-C2B	-2.01	1.37	1.40
22	3	322	CLA	C3D-C4D	-2.01	1.39	1.44
24	3	305	BCR	C38-C26	2.01	1.54	1.50
31	B	852	DGD	CDA-CCA	-2.01	1.33	1.49
22	A	840	CLA	C3D-C4D	-2.01	1.39	1.44
22	8	316	CLA	C3D-C4D	-2.01	1.39	1.44
22	5	320	CLA	C3D-C4D	-2.01	1.39	1.44
22	9	308	CLA	C1C-C2C	2.01	1.48	1.44
22	A	812	CLA	C3D-C4D	-2.01	1.39	1.44
22	1	305	CLA	C1C-C2C	2.01	1.48	1.44
22	6	307	CLA	C3D-C4D	-2.01	1.39	1.44
22	B	817	CLA	C3B-C2B	-2.01	1.37	1.40
22	A	817	CLA	C3D-C4D	-2.01	1.39	1.44
22	B	823	CLA	C3D-C4D	-2.01	1.39	1.44
22	F	302	CLA	C3D-C4D	-2.01	1.39	1.44
22	A	821	CLA	C3B-C2B	-2.00	1.37	1.40
42	7	321	SPH	C3-C4	2.00	1.53	1.50
24	A	856	BCR	C37-C22	2.00	1.55	1.50
22	4	808	CLA	C3D-C4D	-2.00	1.39	1.44
22	6	310	CLA	C1B-NB	2.00	1.37	1.35
24	4	804	BCR	C38-C26	2.00	1.54	1.50
22	6	301	CLA	C3D-C4D	-2.00	1.39	1.44
24	A	844	BCR	C37-C22	2.00	1.55	1.50
22	A	805	CLA	C1C-C2C	2.00	1.48	1.44
22	A	810	CLA	C3D-C4D	-2.00	1.39	1.44

All (4873) bond angle outliers are listed below:

Mol	Chain	Res	Type	Atoms	Z	Observed(^{o})	Ideal(^{o})
24	B	848	BCR	C37-C22-C21	-29.76	81.23	122.92
24	B	848	BCR	C23-C22-C21	28.06	162.01	118.94
24	B	848	BCR	C37-C22-C23	-23.62	80.87	118.08
24	3	304	BCR	C11-C10-C9	-11.56	110.82	127.31
22	K	205	CLA	C4A-NA-C1A	11.24	111.76	106.71
22	9	309	CLA	C4A-NA-C1A	10.90	111.61	106.71
22	9	305	CLA	C4A-NA-C1A	10.78	111.55	106.71
24	A	844	BCR	C15-C14-C13	-10.68	112.06	127.31
22	1	311	CLA	C4A-NA-C1A	10.16	111.27	106.71

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Mol	Chain	Res	Type	Atoms	Z	Observed(^{o})	Ideal(^{o})
22	3	320	CLA	C4A-NA-C1A	9.77	111.10	106.71
22	9	310	CLA	C4A-NA-C1A	9.65	111.05	106.71
24	3	307	BCR	C20-C21-C22	-9.52	113.73	127.31
22	8	307	CLA	C4A-NA-C1A	9.46	110.96	106.71
22	5	320	CLA	C4A-NA-C1A	9.44	110.95	106.71
22	A	822	CLA	C4A-NA-C1A	9.39	110.93	106.71
22	1	316	CLA	C4A-NA-C1A	9.38	110.92	106.71
22	3	315	CLA	C4A-NA-C1A	9.35	110.91	106.71
22	5	323	CLA	C4A-NA-C1A	9.31	110.89	106.71
24	3	307	BCR	C15-C14-C13	-9.30	114.04	127.31
22	6	307	CLA	C4A-NA-C1A	9.29	110.88	106.71
22	7	306	CLA	C4A-NA-C1A	9.18	110.83	106.71
22	4	807	CLA	C4A-NA-C1A	9.17	110.83	106.71
22	B	815	CLA	C4A-NA-C1A	9.16	110.82	106.71
22	9	304	CLA	C4A-NA-C1A	9.16	110.82	106.71
22	A	813	CLA	C4A-NA-C1A	9.12	110.81	106.71
22	3	313	CLA	C4A-NA-C1A	9.12	110.81	106.71
22	A	818	CLA	C4A-NA-C1A	9.07	110.78	106.71
22	A	816	CLA	C4A-NA-C1A	9.07	110.78	106.71
24	A	844	BCR	C7-C8-C9	-9.06	112.55	126.23
22	A	838	CLA	C4A-NA-C1A	9.05	110.77	106.71
22	1	309	CLA	C4A-NA-C1A	9.04	110.77	106.71
22	L	201	CLA	C4A-NA-C1A	9.02	110.76	106.71
22	7	310	CLA	C4A-NA-C1A	9.01	110.75	106.71
22	6	317	CLA	C4A-NA-C1A	8.99	110.75	106.71
22	A	808	CLA	C4A-NA-C1A	8.98	110.75	106.71
22	B	810	CLA	C4A-NA-C1A	8.98	110.74	106.71
22	B	827	CLA	C4A-NA-C1A	8.97	110.74	106.71
22	A	815	CLA	C4A-NA-C1A	8.96	110.73	106.71
22	A	826	CLA	C4A-NA-C1A	8.95	110.73	106.71
22	K	202	CLA	C4A-NA-C1A	8.93	110.72	106.71
22	4	815	CLA	C4A-NA-C1A	8.93	110.72	106.71
22	Z	305	CLA	C4A-NA-C1A	8.93	110.72	106.71
22	B	805	CLA	C4A-NA-C1A	8.90	110.71	106.71
22	A	832	CLA	C4A-NA-C1A	8.89	110.70	106.71
22	8	305	CLA	C4A-NA-C1A	8.89	110.70	106.71
22	A	827	CLA	C4A-NA-C1A	8.88	110.70	106.71
22	7	314	CLA	C4A-NA-C1A	8.88	110.70	106.71
22	5	326	CLA	C4A-NA-C1A	8.87	110.70	106.71
22	5	309	CLA	C4A-NA-C1A	8.87	110.69	106.71
22	B	817	CLA	C4A-NA-C1A	8.86	110.69	106.71
22	7	309	CLA	C4A-NA-C1A	8.85	110.69	106.71

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Mol	Chain	Res	Type	Atoms	Z	Observed(^{o})	Ideal(^{o})
22	5	313	CLA	C4A-NA-C1A	8.85	110.68	106.71
22	B	841	CLA	C4A-NA-C1A	8.84	110.68	106.71
22	5	318	CLA	C4A-NA-C1A	8.82	110.67	106.71
22	A	817	CLA	C4A-NA-C1A	8.80	110.66	106.71
22	B	833	CLA	C4A-NA-C1A	8.80	110.66	106.71
22	B	834	CLA	C4A-NA-C1A	8.80	110.66	106.71
22	A	839	CLA	C4A-NA-C1A	8.80	110.66	106.71
22	Z	314	CLA	C4A-NA-C1A	8.80	110.66	106.71
22	7	304	CLA	C4A-NA-C1A	8.80	110.66	106.71
22	7	315	CLA	C4A-NA-C1A	8.79	110.66	106.71
22	4	810	CLA	C4A-NA-C1A	8.79	110.66	106.71
22	A	842	CLA	C4A-NA-C1A	8.78	110.65	106.71
22	B	837	CLA	C4A-NA-C1A	8.77	110.65	106.71
22	B	811	CLA	C4A-NA-C1A	8.77	110.65	106.71
22	7	311	CLA	C4A-NA-C1A	8.76	110.65	106.71
22	Z	313	CLA	C4A-NA-C1A	8.76	110.64	106.71
22	6	308	CLA	C4A-NA-C1A	8.76	110.64	106.71
22	8	313	CLA	C4A-NA-C1A	8.75	110.64	106.71
22	6	313	CLA	C4A-NA-C1A	8.75	110.64	106.71
22	A	807	CLA	C4A-NA-C1A	8.75	110.64	106.71
22	A	812	CLA	C4A-NA-C1A	8.75	110.64	106.71
22	3	310	CLA	C4A-NA-C1A	8.75	110.64	106.71
22	3	314	CLA	C4A-NA-C1A	8.74	110.64	106.71
22	1	304	CLA	C4A-NA-C1A	8.74	110.64	106.71
22	L	203	CLA	C4A-NA-C1A	8.73	110.63	106.71
22	Z	303	CLA	C4A-NA-C1A	8.73	110.63	106.71
22	6	322	CLA	C4A-NA-C1A	8.73	110.63	106.71
22	6	309	CLA	C4A-NA-C1A	8.73	110.63	106.71
24	B	844	BCR	C11-C10-C9	-8.72	114.86	127.31
22	B	813	CLA	C4A-NA-C1A	8.71	110.62	106.71
22	1	313	CLA	C4A-NA-C1A	8.71	110.62	106.71
22	5	314	CLA	C4A-NA-C1A	8.71	110.62	106.71
22	A	833	CLA	C4A-NA-C1A	8.71	110.62	106.71
22	2	304	CLA	C4A-NA-C1A	8.71	110.62	106.71
22	8	308	CLA	C4A-NA-C1A	8.71	110.62	106.71
22	5	312	CLA	C4A-NA-C1A	8.71	110.62	106.71
22	A	836	CLA	C4A-NA-C1A	8.70	110.62	106.71
22	4	811	CLA	C4A-NA-C1A	8.70	110.62	106.71
22	A	831	CLA	C4A-NA-C1A	8.70	110.62	106.71
22	A	840	CLA	C4A-NA-C1A	8.70	110.62	106.71
22	B	819	CLA	C4A-NA-C1A	8.70	110.62	106.71
22	K	203	CLA	C4A-NA-C1A	8.70	110.62	106.71

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Mol	Chain	Res	Type	Atoms	Z	Observed(^{o})	Ideal(^{o})
22	Z	308	CLA	C4A-NA-C1A	8.69	110.61	106.71
22	1	310	CLA	C4A-NA-C1A	8.69	110.61	106.71
22	A	825	CLA	C4A-NA-C1A	8.69	110.61	106.71
22	G	201	CLA	C4A-NA-C1A	8.68	110.61	106.71
22	A	828	CLA	C4A-NA-C1A	8.68	110.61	106.71
22	8	311	CLA	C4A-NA-C1A	8.68	110.61	106.71
22	F	304	CLA	C4A-NA-C1A	8.67	110.61	106.71
22	1	314	CLA	C4A-NA-C1A	8.67	110.61	106.71
22	G	202	CLA	C4A-NA-C1A	8.67	110.60	106.71
22	7	324	CLA	C4A-NA-C1A	8.67	110.60	106.71
22	3	318	CLA	C4A-NA-C1A	8.67	110.60	106.71
22	B	830	CLA	C4A-NA-C1A	8.66	110.60	106.71
22	6	321	CLA	C4A-NA-C1A	8.66	110.60	106.71
22	A	805	CLA	C4A-NA-C1A	8.65	110.60	106.71
22	3	308	CLA	C4A-NA-C1A	8.65	110.60	106.71
22	B	826	CLA	C4A-NA-C1A	8.65	110.60	106.71
22	3	311	CLA	C4A-NA-C1A	8.65	110.60	106.71
22	F	302	CLA	C4A-NA-C1A	8.65	110.59	106.71
22	A	804	CLA	C4A-NA-C1A	8.64	110.59	106.71
22	B	842	CLA	C4A-NA-C1A	8.64	110.59	106.71
22	8	316	CLA	C4A-NA-C1A	8.64	110.59	106.71
22	B	838	CLA	C4A-NA-C1A	8.64	110.59	106.71
22	5	310	CLA	C4A-NA-C1A	8.64	110.59	106.71
22	5	315	CLA	C4A-NA-C1A	8.64	110.59	106.71
22	A	837	CLA	C4A-NA-C1A	8.64	110.59	106.71
22	1	307	CLA	C4A-NA-C1A	8.64	110.59	106.71
22	6	312	CLA	C4A-NA-C1A	8.64	110.59	106.71
22	B	818	CLA	C4A-NA-C1A	8.63	110.59	106.71
22	1	315	CLA	C4A-NA-C1A	8.63	110.59	106.71
22	3	319	CLA	C4A-NA-C1A	8.63	110.58	106.71
22	A	814	CLA	C4A-NA-C1A	8.62	110.58	106.71
22	6	301	CLA	C4A-NA-C1A	8.62	110.58	106.71
22	B	839	CLA	C4A-NA-C1A	8.62	110.58	106.71
22	7	316	CLA	C4A-NA-C1A	8.62	110.58	106.71
22	5	322	CLA	C4A-NA-C1A	8.62	110.58	106.71
22	5	311	CLA	C4A-NA-C1A	8.61	110.58	106.71
22	4	812	CLA	C4A-NA-C1A	8.61	110.58	106.71
22	B	820	CLA	C4A-NA-C1A	8.61	110.58	106.71
22	B	832	CLA	C4A-NA-C1A	8.60	110.57	106.71
22	A	830	CLA	C4A-NA-C1A	8.59	110.57	106.71
22	4	818	CLA	C4A-NA-C1A	8.59	110.57	106.71
22	5	307	CLA	C4A-NA-C1A	8.59	110.57	106.71

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Mol	Chain	Res	Type	Atoms	Z	Observed(^{o})	Ideal(^{o})
22	B	808	CLA	C4A-NA-C1A	8.59	110.57	106.71
22	4	808	CLA	C4A-NA-C1A	8.58	110.56	106.71
22	A	841	CLA	C4A-NA-C1A	8.58	110.56	106.71
22	2	302	CLA	C4A-NA-C1A	8.57	110.56	106.71
22	6	319	CLA	C4A-NA-C1A	8.57	110.56	106.71
22	Z	306	CLA	C4A-NA-C1A	8.57	110.56	106.71
22	Z	315	CLA	C4A-NA-C1A	8.57	110.56	106.71
22	A	802	CLA	C4A-NA-C1A	8.57	110.56	106.71
22	A	810	CLA	C4A-NA-C1A	8.57	110.56	106.71
22	B	806	CLA	C4A-NA-C1A	8.57	110.56	106.71
22	L	202	CLA	C4A-NA-C1A	8.57	110.56	106.71
22	B	829	CLA	C4A-NA-C1A	8.56	110.55	106.71
22	A	821	CLA	C4A-NA-C1A	8.56	110.55	106.71
22	Z	309	CLA	C4A-NA-C1A	8.56	110.55	106.71
22	5	319	CLA	C4A-NA-C1A	8.55	110.55	106.71
22	9	313	CLA	C4A-NA-C1A	8.55	110.55	106.71
22	9	306	CLA	C4A-NA-C1A	8.55	110.55	106.71
22	1	305	CLA	C4A-NA-C1A	8.55	110.55	106.71
22	F	301	CLA	C4A-NA-C1A	8.54	110.54	106.71
22	9	311	CLA	C4A-NA-C1A	8.54	110.54	106.71
22	B	807	CLA	C4A-NA-C1A	8.53	110.54	106.71
22	A	824	CLA	C4A-NA-C1A	8.53	110.54	106.71
22	B	840	CLA	C4A-NA-C1A	8.52	110.54	106.71
22	K	204	CLA	C4A-NA-C1A	8.51	110.53	106.71
22	Z	304	CLA	C4A-NA-C1A	8.51	110.53	106.71
22	A	803	CLA	C4A-NA-C1A	8.51	110.53	106.71
22	A	854	CLA	C4A-NA-C1A	8.51	110.53	106.71
22	B	831	CLA	C4A-NA-C1A	8.51	110.53	106.71
22	7	323	CLA	C4A-NA-C1A	8.50	110.53	106.71
22	5	308	CLA	C4A-NA-C1A	8.50	110.53	106.71
22	A	823	CLA	C4A-NA-C1A	8.49	110.52	106.71
22	4	817	CLA	C4A-NA-C1A	8.49	110.52	106.71
22	B	835	CLA	C4A-NA-C1A	8.49	110.52	106.71
22	B	836	CLA	C4A-NA-C1A	8.48	110.52	106.71
22	5	301	CLA	C4A-NA-C1A	8.48	110.52	106.71
22	2	301	CLA	C4A-NA-C1A	8.48	110.52	106.71
22	B	814	CLA	C4A-NA-C1A	8.48	110.52	106.71
22	3	309	CLA	C4A-NA-C1A	8.48	110.52	106.71
22	6	302	CLA	C4A-NA-C1A	8.47	110.51	106.71
22	B	824	CLA	C4A-NA-C1A	8.47	110.51	106.71
22	B	823	CLA	C4A-NA-C1A	8.46	110.51	106.71
22	7	322	CLA	C4A-NA-C1A	8.46	110.51	106.71

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Mol	Chain	Res	Type	Atoms	Z	Observed(^{o})	Ideal(^{o})
22	8	309	CLA	C4A-NA-C1A	8.44	110.50	106.71
22	Z	310	CLA	C4A-NA-C1A	8.43	110.50	106.71
22	J	103	CLA	C4A-NA-C1A	8.43	110.50	106.71
22	A	819	CLA	C4A-NA-C1A	8.43	110.50	106.71
22	4	806	CLA	C4A-NA-C1A	8.43	110.50	106.71
22	8	314	CLA	C4A-NA-C1A	8.43	110.50	106.71
22	9	308	CLA	C4A-NA-C1A	8.43	110.50	106.71
22	7	312	CLA	C4A-NA-C1A	8.43	110.49	106.71
22	A	820	CLA	C4A-NA-C1A	8.41	110.49	106.71
22	A	834	CLA	C4A-NA-C1A	8.41	110.49	106.71
22	F	305	CLA	C4A-NA-C1A	8.41	110.49	106.71
22	B	801	CLA	C4A-NA-C1A	8.40	110.48	106.71
22	8	310	CLA	C4A-NA-C1A	8.39	110.48	106.71
22	A	829	CLA	C4A-NA-C1A	8.38	110.47	106.71
22	B	816	CLA	C4A-NA-C1A	8.38	110.47	106.71
22	Z	316	CLA	C4A-NA-C1A	8.38	110.47	106.71
22	7	305	CLA	C4A-NA-C1A	8.38	110.47	106.71
22	B	821	CLA	C4A-NA-C1A	8.37	110.47	106.71
22	7	308	CLA	C4A-NA-C1A	8.37	110.47	106.71
22	3	316	CLA	C4A-NA-C1A	8.37	110.47	106.71
22	B	812	CLA	C4A-NA-C1A	8.36	110.46	106.71
22	A	811	CLA	C4A-NA-C1A	8.35	110.46	106.71
22	A	835	CLA	C4A-NA-C1A	8.33	110.45	106.71
22	B	822	CLA	C4A-NA-C1A	8.33	110.45	106.71
22	8	306	CLA	C4A-NA-C1A	8.30	110.44	106.71
22	1	306	CLA	C4A-NA-C1A	8.29	110.43	106.71
22	A	855	CLA	C4A-NA-C1A	8.29	110.43	106.71
24	5	305	BCR	C20-C21-C22	-8.28	115.49	127.31
22	6	310	CLA	C4A-NA-C1A	8.27	110.42	106.71
22	7	307	CLA	C4A-NA-C1A	8.27	110.42	106.71
22	7	317	CLA	C4A-NA-C1A	8.27	110.42	106.71
22	B	825	CLA	C4A-NA-C1A	8.25	110.42	106.71
22	6	314	CLA	C4A-NA-C1A	8.25	110.42	106.71
22	3	322	CLA	C4A-NA-C1A	8.24	110.41	106.71
22	B	828	CLA	C4A-NA-C1A	8.23	110.41	106.71
22	4	809	CLA	C4A-NA-C1A	8.22	110.40	106.71
22	Z	307	CLA	C4A-NA-C1A	8.20	110.39	106.71
22	A	806	CLA	C4A-NA-C1A	8.19	110.39	106.71
22	6	311	CLA	C4A-NA-C1A	8.19	110.39	106.71
22	9	307	CLA	C4A-NA-C1A	8.19	110.39	106.71
24	6	306	BCR	C24-C23-C22	-8.19	113.87	126.23
24	B	848	BCR	C20-C21-C22	-8.17	115.65	127.31

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Mol	Chain	Res	Type	Atoms	Z	Observed(^{o})	Ideal(^{o})
22	3	312	CLA	C4A-NA-C1A	8.17	110.38	106.71
22	B	809	CLA	C4A-NA-C1A	8.12	110.36	106.71
22	1	308	CLA	C4A-NA-C1A	8.10	110.35	106.71
24	3	304	BCR	C7-C8-C9	-8.03	114.10	126.23
22	9	303	CLA	C4A-NA-C1A	8.02	110.31	106.71
22	A	809	CLA	C4A-NA-C1A	8.01	110.31	106.71
21	A	801	CL0	CMD-C2D-C1D	7.95	138.73	124.71
22	4	805	CLA	C4A-NA-C1A	7.80	110.21	106.71
24	6	306	BCR	C16-C17-C18	-7.58	116.49	127.31
24	3	307	BCR	C24-C23-C22	-7.48	114.93	126.23
24	6	306	BCR	C11-C10-C9	-7.35	116.82	127.31
22	9	305	CLA	CAA-C2A-C3A	-7.25	92.94	112.78
24	5	305	BCR	C15-C14-C13	-7.24	116.98	127.31
24	B	847	BCR	C15-C14-C13	-7.16	117.09	127.31
24	I	4001	BCR	C16-C17-C18	-7.15	117.11	127.31
24	A	846	BCR	C15-C14-C13	-7.14	117.12	127.31
32	F	306	RRX	C11-C10-C9	-6.88	117.49	127.31
24	3	304	BCR	C15-C16-C17	-6.66	109.82	123.47
21	A	801	CL0	C4A-NA-C1A	6.60	109.67	106.71
24	5	305	BCR	C11-C10-C9	-6.40	118.18	127.31
36	5	306	C7Z	C11-C10-C9	-6.35	118.25	127.31
24	3	307	BCR	C11-C10-C9	-6.31	118.30	127.31
24	G	203	BCR	C20-C21-C22	-6.30	118.31	127.31
24	5	304	BCR	C20-C21-C22	-6.28	118.34	127.31
24	B	844	BCR	C20-C21-C22	-6.26	118.38	127.31
24	K	206	BCR	C11-C10-C9	-6.22	118.43	127.31
21	A	801	CL0	C2D-C1D-ND	6.19	114.67	110.10
22	1	308	CLA	CMD-C2D-C1D	6.07	135.42	124.71
21	A	801	CL0	C2C-C1C-NC	6.06	115.65	109.97
24	A	848	BCR	C20-C21-C22	-6.06	118.66	127.31
24	6	305	BCR	C20-C21-C22	-6.06	118.67	127.31
24	B	848	BCR	C33-C5-C6	-6.03	117.76	124.53
24	A	856	BCR	C20-C21-C22	-6.03	118.71	127.31
24	3	307	BCR	C38-C26-C25	-6.01	117.78	124.53
24	B	849	BCR	C20-C21-C22	-5.99	118.77	127.31
38	5	302	LUT	C35-C34-C33	-5.98	118.78	127.31
22	7	304	CLA	O2A-C1-C2	5.97	124.34	108.64
22	9	305	CLA	O2D-CGD-CBD	5.96	121.86	111.27
24	A	845	BCR	C16-C17-C18	-5.95	118.81	127.31
24	5	304	BCR	C16-C17-C18	-5.94	118.83	127.31
24	B	846	BCR	C15-C14-C13	-5.93	118.85	127.31
22	B	831	CLA	CMD-C2D-C1D	5.93	135.16	124.71

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Mol	Chain	Res	Type	Atoms	Z	Observed(^{o})	Ideal(^{o})
38	4	803	LUT	C35-C34-C33	-5.92	118.86	127.31
22	7	307	CLA	CMD-C2D-C1D	5.86	135.03	124.71
25	3	321	LHG	O7-C7-O9	-5.85	118.12	125.57
22	B	835	CLA	CMD-C2D-C1D	5.85	135.02	124.71
24	I	4001	BCR	C24-C23-C22	-5.84	117.41	126.23
24	3	305	BCR	C33-C5-C6	-5.84	117.97	124.53
38	6	303	LUT	C35-C34-C33	-5.83	118.99	127.31
24	B	845	BCR	C1-C6-C5	-5.83	114.41	122.61
24	F	303	BCR	C16-C17-C18	-5.80	119.04	127.31
22	9	303	CLA	O2A-C1-C2	5.78	123.81	108.64
22	5	307	CLA	CMD-C2D-C1D	5.77	134.89	124.71
22	A	818	CLA	CMD-C2D-C1D	5.77	134.88	124.71
22	6	322	CLA	O2D-CGD-CBD	5.77	121.51	111.27
22	2	301	CLA	CMD-C2D-C1D	5.76	134.86	124.71
22	B	838	CLA	CMD-C2D-C1D	5.73	134.82	124.71
36	5	306	C7Z	C15-C14-C13	-5.73	119.13	127.31
22	1	309	CLA	CMD-C2D-C1D	5.72	134.79	124.71
24	A	847	BCR	C38-C26-C25	-5.71	118.11	124.53
22	B	816	CLA	CMD-C2D-C1D	5.70	134.76	124.71
32	F	306	RRX	C12-C13-C14	5.70	127.69	118.94
22	7	315	CLA	O2D-CGD-CBD	5.69	121.39	111.27
22	B	824	CLA	CMD-C2D-C1D	5.69	134.74	124.71
22	5	323	CLA	CMD-C2D-C1D	5.68	134.73	124.71
22	A	825	CLA	CMD-C2D-C1D	5.68	134.73	124.71
22	A	811	CLA	CMD-C2D-C1D	5.68	134.72	124.71
22	F	304	CLA	CMD-C2D-C1D	5.67	134.70	124.71
22	Z	306	CLA	CMD-C2D-C1D	5.66	134.69	124.71
22	A	807	CLA	CMD-C2D-C1D	5.66	134.68	124.71
22	B	826	CLA	CMD-C2D-C1D	5.66	134.68	124.71
22	A	823	CLA	CMD-C2D-C1D	5.65	134.67	124.71
22	6	309	CLA	CMD-C2D-C1D	5.65	134.67	124.71
22	Z	307	CLA	CMD-C2D-C1D	5.65	134.67	124.71
22	K	204	CLA	O2A-C1-C2	5.64	122.16	108.97
22	A	805	CLA	CMD-C2D-C1D	5.64	134.65	124.71
24	J	104	BCR	C27-C26-C25	-5.64	114.55	122.73
38	Z	301	LUT	C11-C10-C9	-5.63	119.27	127.31
22	8	305	CLA	CMD-C2D-C1D	5.63	134.64	124.71
22	Z	303	CLA	CMD-C2D-C1D	5.63	134.64	124.71
22	1	307	CLA	CMD-C2D-C1D	5.63	134.63	124.71
38	1	302	LUT	C11-C10-C9	-5.62	119.29	127.31
22	A	810	CLA	CMD-C2D-C1D	5.62	134.61	124.71
22	4	818	CLA	O2A-C1-C2	5.61	123.38	108.64

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Mol	Chain	Res	Type	Atoms	Z	Observed(^{o})	Ideal(^{o})
22	9	306	CLA	CMD-C2D-C1D	5.61	134.59	124.71
22	B	832	CLA	CMD-C2D-C1D	5.61	134.59	124.71
22	1	315	CLA	CMD-C2D-C1D	5.61	134.59	124.71
22	5	310	CLA	CMD-C2D-C1D	5.61	134.59	124.71
38	4	802	LUT	C7-C8-C9	-5.61	117.77	126.23
32	F	306	RRX	C20-C21-C22	-5.60	119.31	127.31
22	1	305	CLA	CMD-C2D-C1D	5.60	134.59	124.71
24	5	304	BCR	C11-C10-C9	-5.60	119.32	127.31
22	B	840	CLA	CMD-C2D-C1D	5.59	134.57	124.71
22	1	313	CLA	CMD-C2D-C1D	5.59	134.57	124.71
22	A	827	CLA	CMD-C2D-C1D	5.59	134.57	124.71
22	3	322	CLA	CMD-C2D-C1D	5.59	134.56	124.71
22	B	814	CLA	CMD-C2D-C1D	5.59	134.56	124.71
22	B	828	CLA	CMD-C2D-C1D	5.59	134.56	124.71
22	8	309	CLA	CMD-C2D-C1D	5.58	134.55	124.71
22	A	812	CLA	CMD-C2D-C1D	5.58	134.55	124.71
22	A	832	CLA	CMD-C2D-C1D	5.58	134.55	124.71
24	A	847	BCR	C16-C17-C18	-5.58	119.35	127.31
22	4	812	CLA	CMD-C2D-C1D	5.58	134.54	124.71
22	4	809	CLA	CMD-C2D-C1D	5.58	134.54	124.71
22	B	818	CLA	CMD-C2D-C1D	5.57	134.54	124.71
22	A	809	CLA	CMD-C2D-C1D	5.57	134.54	124.71
22	A	839	CLA	CMD-C2D-C1D	5.57	134.53	124.71
22	5	301	CLA	CMD-C2D-C1D	5.57	134.53	124.71
22	3	313	CLA	CMD-C2D-C1D	5.57	134.53	124.71
22	F	305	CLA	CMD-C2D-C1D	5.57	134.53	124.71
22	3	308	CLA	CMD-C2D-C1D	5.57	134.53	124.71
22	A	819	CLA	CMD-C2D-C1D	5.57	134.53	124.71
22	5	309	CLA	CMD-C2D-C1D	5.57	134.52	124.71
22	5	308	CLA	CMD-C2D-C1D	5.56	134.52	124.71
22	6	322	CLA	CMD-C2D-C1D	5.56	134.52	124.71
22	6	310	CLA	CMD-C2D-C1D	5.56	134.51	124.71
22	3	309	CLA	CMD-C2D-C1D	5.56	134.51	124.71
22	Z	308	CLA	CMD-C2D-C1D	5.56	134.51	124.71
24	A	848	BCR	C7-C8-C9	-5.56	117.84	126.23
22	A	833	CLA	CMD-C2D-C1D	5.55	134.50	124.71
22	7	304	CLA	CMD-C2D-C1D	5.55	134.50	124.71
22	3	318	CLA	CMD-C2D-C1D	5.55	134.50	124.71
22	A	840	CLA	CMD-C2D-C1D	5.55	134.50	124.71
22	7	323	CLA	CMD-C2D-C1D	5.55	134.49	124.71
22	A	802	CLA	O2A-C1-C2	5.55	123.22	108.64
22	Z	315	CLA	CMD-C2D-C1D	5.55	134.49	124.71

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Mol	Chain	Res	Type	Atoms	Z	Observed(^{o})	Ideal(^{o})
22	B	815	CLA	CMD-C2D-C1D	5.55	134.49	124.71
22	G	201	CLA	CMD-C2D-C1D	5.54	134.49	124.71
22	7	324	CLA	CMD-C2D-C1D	5.54	134.48	124.71
22	A	822	CLA	CMD-C2D-C1D	5.54	134.48	124.71
22	A	837	CLA	CMD-C2D-C1D	5.54	134.48	124.71
22	B	842	CLA	CMD-C2D-C1D	5.54	134.48	124.71
22	B	813	CLA	CMD-C2D-C1D	5.54	134.48	124.71
22	Z	316	CLA	CMD-C2D-C1D	5.54	134.47	124.71
22	K	203	CLA	CMD-C2D-C1D	5.54	134.47	124.71
22	G	202	CLA	CMD-C2D-C1D	5.53	134.47	124.71
22	L	201	CLA	CMD-C2D-C1D	5.53	134.47	124.71
22	B	836	CLA	CMD-C2D-C1D	5.53	134.47	124.71
24	I	4001	BCR	C20-C21-C22	-5.53	119.41	127.31
22	5	309	CLA	O2A-C1-C2	5.53	123.17	108.64
22	7	309	CLA	CMD-C2D-C1D	5.53	134.46	124.71
22	7	305	CLA	CMD-C2D-C1D	5.53	134.46	124.71
24	A	846	BCR	C11-C10-C9	-5.53	119.42	127.31
22	B	829	CLA	CMD-C2D-C1D	5.53	134.46	124.71
22	B	810	CLA	CMD-C2D-C1D	5.53	134.45	124.71
22	B	819	CLA	CMD-C2D-C1D	5.53	134.45	124.71
22	B	835	CLA	O2D-CGD-CBD	5.53	121.09	111.27
22	5	326	CLA	CMD-C2D-C1D	5.52	134.44	124.71
22	B	808	CLA	CMD-C2D-C1D	5.52	134.44	124.71
24	B	849	BCR	C16-C17-C18	-5.52	119.44	127.31
22	4	810	CLA	CMD-C2D-C1D	5.52	134.44	124.71
22	8	308	CLA	CMD-C2D-C1D	5.52	134.44	124.71
22	A	831	CLA	CMD-C2D-C1D	5.52	134.43	124.71
22	2	302	CLA	CMD-C2D-C1D	5.51	134.43	124.71
22	3	316	CLA	CMD-C2D-C1D	5.51	134.43	124.71
22	7	308	CLA	CMD-C2D-C1D	5.51	134.43	124.71
22	A	838	CLA	CMD-C2D-C1D	5.51	134.42	124.71
22	6	312	CLA	CMD-C2D-C1D	5.51	134.42	124.71
22	B	825	CLA	CMD-C2D-C1D	5.51	134.42	124.71
22	B	817	CLA	O2D-CGD-CBD	5.51	121.05	111.27
22	B	833	CLA	CMD-C2D-C1D	5.50	134.41	124.71
22	7	314	CLA	CMD-C2D-C1D	5.50	134.41	124.71
22	A	841	CLA	CMD-C2D-C1D	5.50	134.41	124.71
22	A	834	CLA	CMD-C2D-C1D	5.50	134.41	124.71
22	6	307	CLA	CMD-C2D-C1D	5.50	134.41	124.71
22	A	802	CLA	CMD-C2D-C1D	5.50	134.41	124.71
22	K	204	CLA	CMD-C2D-C1D	5.50	134.41	124.71
22	B	823	CLA	CMD-C2D-C1D	5.50	134.41	124.71

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Mol	Chain	Res	Type	Atoms	Z	Observed(^{o})	Ideal(^{o})
22	F	301	CLA	CMD-C2D-C1D	5.50	134.41	124.71
22	4	817	CLA	CMD-C2D-C1D	5.50	134.41	124.71
22	B	827	CLA	CMD-C2D-C1D	5.50	134.41	124.71
22	A	836	CLA	CMD-C2D-C1D	5.50	134.40	124.71
22	3	319	CLA	CMD-C2D-C1D	5.50	134.40	124.71
22	5	311	CLA	CMD-C2D-C1D	5.50	134.40	124.71
22	3	312	CLA	CMD-C2D-C1D	5.50	134.40	124.71
22	B	834	CLA	CMD-C2D-C1D	5.50	134.40	124.71
22	8	307	CLA	CMD-C2D-C1D	5.49	134.40	124.71
22	F	302	CLA	CMD-C2D-C1D	5.49	134.40	124.71
24	A	845	BCR	C15-C14-C13	-5.49	119.47	127.31
22	4	818	CLA	CMD-C2D-C1D	5.49	134.40	124.71
22	7	312	CLA	CMD-C2D-C1D	5.49	134.39	124.71
22	A	815	CLA	CMD-C2D-C1D	5.49	134.39	124.71
22	A	817	CLA	CMD-C2D-C1D	5.49	134.38	124.71
22	B	811	CLA	CMD-C2D-C1D	5.49	134.38	124.71
22	A	821	CLA	CMD-C2D-C1D	5.49	134.38	124.71
22	A	813	CLA	CMD-C2D-C1D	5.48	134.38	124.71
22	A	855	CLA	CMD-C2D-C1D	5.48	134.38	124.71
22	2	302	CLA	O2D-CGD-CBD	5.48	121.01	111.27
22	9	308	CLA	CMD-C2D-C1D	5.48	134.37	124.71
22	B	809	CLA	CMD-C2D-C1D	5.48	134.37	124.71
22	8	310	CLA	CMD-C2D-C1D	5.48	134.37	124.71
22	B	821	CLA	CMD-C2D-C1D	5.47	134.36	124.71
22	7	323	CLA	O2D-CGD-CBD	5.47	121.00	111.27
38	7	301	LUT	C35-C34-C33	-5.47	119.50	127.31
22	L	203	CLA	CMD-C2D-C1D	5.47	134.35	124.71
22	6	301	CLA	CMD-C2D-C1D	5.47	134.35	124.71
38	7	302	LUT	C11-C10-C9	-5.47	119.51	127.31
24	I	4001	BCR	C30-C25-C26	-5.47	114.91	122.61
21	A	801	CL0	C1C-C2C-C3C	-5.47	101.21	106.96
22	6	311	CLA	CMD-C2D-C1D	5.47	134.35	124.71
22	B	806	CLA	CMD-C2D-C1D	5.46	134.34	124.71
22	4	808	CLA	CMD-C2D-C1D	5.46	134.34	124.71
22	A	816	CLA	CMD-C2D-C1D	5.46	134.34	124.71
22	5	312	CLA	CMD-C2D-C1D	5.46	134.33	124.71
22	6	319	CLA	CMD-C2D-C1D	5.46	134.33	124.71
22	8	310	CLA	O2A-C1-C2	5.46	122.97	108.64
22	7	311	CLA	CMD-C2D-C1D	5.46	134.33	124.71
22	J	103	CLA	CMD-C2D-C1D	5.46	134.33	124.71
22	9	311	CLA	CMD-C2D-C1D	5.45	134.32	124.71
22	7	316	CLA	CMD-C2D-C1D	5.45	134.32	124.71

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Mol	Chain	Res	Type	Atoms	Z	Observed(^{o})	Ideal(^{o})
22	K	202	CLA	CMD-C2D-C1D	5.45	134.32	124.71
22	6	319	CLA	O2D-CGD-CBD	5.45	120.95	111.27
24	6	305	BCR	C11-C10-C9	-5.44	119.54	127.31
22	A	803	CLA	CMD-C2D-C1D	5.44	134.30	124.71
22	1	306	CLA	CMD-C2D-C1D	5.44	134.30	124.71
22	7	322	CLA	CMD-C2D-C1D	5.44	134.30	124.71
22	4	806	CLA	CMD-C2D-C1D	5.43	134.29	124.71
22	B	826	CLA	O2D-CGD-CBD	5.43	120.92	111.27
22	B	820	CLA	CMD-C2D-C1D	5.43	134.28	124.71
22	A	828	CLA	CMD-C2D-C1D	5.43	134.28	124.71
22	3	314	CLA	O2A-C1-C2	5.43	122.90	108.64
22	5	319	CLA	CMD-C2D-C1D	5.42	134.27	124.71
22	3	310	CLA	CMD-C2D-C1D	5.42	134.27	124.71
22	B	839	CLA	CMD-C2D-C1D	5.42	134.27	124.71
22	4	807	CLA	CMD-C2D-C1D	5.42	134.27	124.71
22	7	317	CLA	CMD-C2D-C1D	5.42	134.27	124.71
32	F	306	RRX	C11-C12-C13	-5.41	111.21	126.42
22	8	316	CLA	CMD-C2D-C1D	5.41	134.25	124.71
22	A	826	CLA	CMD-C2D-C1D	5.41	134.25	124.71
22	A	814	CLA	CMD-C2D-C1D	5.41	134.25	124.71
22	6	314	CLA	CMD-C2D-C1D	5.41	134.25	124.71
24	B	847	BCR	C11-C10-C9	-5.41	119.59	127.31
22	4	805	CLA	CMD-C2D-C1D	5.41	134.25	124.71
22	A	842	CLA	CMD-C2D-C1D	5.41	134.25	124.71
24	G	203	BCR	C16-C17-C18	-5.41	119.59	127.31
22	A	830	CLA	CMD-C2D-C1D	5.41	134.25	124.71
24	A	856	BCR	C27-C26-C25	-5.41	114.88	122.73
22	6	308	CLA	CMD-C2D-C1D	5.41	134.24	124.71
22	1	304	CLA	CMD-C2D-C1D	5.40	134.24	124.71
22	A	820	CLA	CMD-C2D-C1D	5.40	134.24	124.71
22	B	812	CLA	CMD-C2D-C1D	5.40	134.23	124.71
22	A	808	CLA	CMD-C2D-C1D	5.40	134.23	124.71
22	6	321	CLA	CMD-C2D-C1D	5.39	134.22	124.71
22	8	313	CLA	CMD-C2D-C1D	5.39	134.22	124.71
24	3	306	BCR	C1-C6-C5	-5.39	115.02	122.61
22	A	824	CLA	CMD-C2D-C1D	5.39	134.21	124.71
22	1	310	CLA	CMD-C2D-C1D	5.39	134.20	124.71
22	L	202	CLA	CMD-C2D-C1D	5.38	134.20	124.71
24	3	307	BCR	C15-C16-C17	-5.38	112.45	123.47
22	5	318	CLA	CMD-C2D-C1D	5.38	134.20	124.71
24	A	847	BCR	C20-C21-C22	-5.38	119.63	127.31
22	Z	303	CLA	O2D-CGD-CBD	5.38	120.82	111.27

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Mol	Chain	Res	Type	Atoms	Z	Observed(^{o})	Ideal(^{o})
22	B	837	CLA	CMD-C2D-C1D	5.38	134.19	124.71
22	3	311	CLA	CMD-C2D-C1D	5.38	134.19	124.71
22	B	807	CLA	CMD-C2D-C1D	5.38	134.19	124.71
22	5	308	CLA	O2D-CGD-CBD	5.37	120.82	111.27
24	B	853	BCR	C1-C6-C5	-5.37	115.05	122.61
22	9	307	CLA	CMD-C2D-C1D	5.37	134.18	124.71
22	5	322	CLA	CMD-C2D-C1D	5.37	134.17	124.71
22	8	306	CLA	CMD-C2D-C1D	5.36	134.17	124.71
22	A	804	CLA	CMD-C2D-C1D	5.36	134.16	124.71
22	4	815	CLA	O2D-CGD-CBD	5.36	120.79	111.27
22	1	314	CLA	CMD-C2D-C1D	5.36	134.16	124.71
22	9	305	CLA	CMD-C2D-C1D	5.35	134.15	124.71
22	A	806	CLA	CMD-C2D-C1D	5.35	134.14	124.71
22	A	803	CLA	O2D-CGD-CBD	5.35	120.77	111.27
22	B	841	CLA	CMD-C2D-C1D	5.35	134.13	124.71
24	G	203	BCR	C33-C5-C6	-5.34	118.53	124.53
22	7	306	CLA	CMD-C2D-C1D	5.33	134.12	124.71
22	Z	307	CLA	O2D-CGD-CBD	5.33	120.75	111.27
22	A	854	CLA	CMD-C2D-C1D	5.33	134.11	124.71
24	K	206	BCR	C15-C14-C13	-5.33	119.70	127.31
38	9	302	LUT	C35-C34-C33	-5.33	119.71	127.31
22	4	811	CLA	CMD-C2D-C1D	5.32	134.09	124.71
22	7	323	CLA	O2A-C1-C2	5.32	122.61	108.64
22	7	324	CLA	O2A-C1-C2	5.32	122.61	108.64
22	B	805	CLA	CMD-C2D-C1D	5.32	134.08	124.71
24	B	846	BCR	C24-C23-C22	-5.31	118.20	126.23
22	B	801	CLA	CMD-C2D-C1D	5.31	134.07	124.71
22	A	804	CLA	O2D-CGD-CBD	5.31	120.69	111.27
24	5	304	BCR	C38-C26-C25	-5.30	118.57	124.53
22	B	822	CLA	CMD-C2D-C1D	5.30	134.05	124.71
22	5	313	CLA	O2A-C1-C2	5.30	122.56	108.64
22	A	816	CLA	O2D-CGD-CBD	5.29	120.67	111.27
22	Z	310	CLA	CMD-C2D-C1D	5.29	134.04	124.71
22	1	308	CLA	O2D-CGD-CBD	5.29	120.67	111.27
38	3	302	LUT	C35-C34-C33	-5.29	119.76	127.31
22	8	306	CLA	O2D-CGD-CBD	5.29	120.66	111.27
22	A	809	CLA	O2D-CGD-CBD	5.28	120.65	111.27
22	A	834	CLA	O2D-CGD-CBD	5.28	120.64	111.27
22	4	811	CLA	O2A-C1-C2	5.27	122.50	108.64
22	1	306	CLA	O2A-C1-C2	5.27	122.49	108.64
22	6	309	CLA	O2A-C1-C2	5.27	122.49	108.64
22	9	313	CLA	CMD-C2D-C1D	5.27	134.00	124.71

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Mol	Chain	Res	Type	Atoms	Z	Observed(^{o})	Ideal(^{o})
36	J	105	C7Z	C35-C34-C33	-5.27	119.79	127.31
38	8	303	LUT	C35-C34-C33	-5.27	119.79	127.31
22	Z	313	CLA	O2A-C1-C2	5.27	122.48	108.64
38	8	302	LUT	C35-C34-C33	-5.27	119.79	127.31
36	J	105	C7Z	C27-C28-C29	-5.26	118.28	126.23
22	9	306	CLA	O2A-C1-C2	5.25	122.44	108.64
22	A	818	CLA	O2A-C1-C2	5.25	122.44	108.64
22	A	833	CLA	O2D-CGD-CBD	5.25	120.60	111.27
22	6	308	CLA	O2D-CGD-CBD	5.25	120.59	111.27
22	B	831	CLA	O2D-CGD-CBD	5.24	120.58	111.27
22	5	313	CLA	CMD-C2D-C1D	5.24	133.95	124.71
22	B	820	CLA	O2D-CGD-CBD	5.24	120.58	111.27
24	A	856	BCR	C16-C17-C18	-5.24	119.83	127.31
22	9	308	CLA	O2D-CGD-CBD	5.23	120.56	111.27
22	A	835	CLA	CMD-C2D-C1D	5.23	133.93	124.71
22	9	303	CLA	CMD-C2D-C1D	5.23	133.93	124.71
24	6	305	BCR	C7-C8-C9	-5.23	118.34	126.23
22	B	810	CLA	O2D-CGD-CBD	5.22	120.55	111.27
22	1	311	CLA	O2D-CGD-CBD	5.22	120.54	111.27
22	A	829	CLA	CMD-C2D-C1D	5.22	133.91	124.71
22	A	814	CLA	O2A-C1-C2	5.22	122.35	108.64
22	1	313	CLA	O2A-C1-C2	5.22	122.35	108.64
22	Z	313	CLA	O2D-CGD-CBD	5.22	120.54	111.27
22	8	314	CLA	CMD-C2D-C1D	5.20	133.89	124.71
22	9	313	CLA	O2D-CGD-CBD	5.20	120.51	111.27
22	7	307	CLA	O2D-CGD-CBD	5.20	120.51	111.27
38	7	302	LUT	C15-C14-C13	-5.20	119.89	127.31
24	A	847	BCR	C24-C23-C22	-5.20	118.38	126.23
22	5	314	CLA	CMD-C2D-C1D	5.19	133.87	124.71
22	7	305	CLA	O2D-CGD-CBD	5.19	120.50	111.27
22	B	834	CLA	O2D-CGD-CBD	5.19	120.49	111.27
22	B	811	CLA	O2D-CGD-CBD	5.19	120.49	111.27
24	3	306	BCR	C15-C14-C13	-5.19	119.91	127.31
22	5	319	CLA	O2A-C1-C2	5.18	122.26	108.64
22	B	837	CLA	O2D-CGD-CBD	5.18	120.48	111.27
24	7	303	BCR	C11-C10-C9	-5.18	119.91	127.31
22	B	814	CLA	O2D-CGD-CBD	5.18	120.47	111.27
22	7	310	CLA	CMD-C2D-C1D	5.18	133.84	124.71
22	6	314	CLA	O2D-CGD-CBD	5.18	120.47	111.27
21	A	801	CL0	O2D-CGD-CBD	5.18	120.47	111.27
22	5	311	CLA	O2D-CGD-CBD	5.17	120.46	111.27
22	5	301	CLA	O2A-C1-C2	5.17	122.23	108.64

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Mol	Chain	Res	Type	Atoms	Z	Observed(^{o})	Ideal(^{o})
24	B	853	BCR	C7-C8-C9	-5.17	118.42	126.23
24	7	303	BCR	C38-C26-C25	-5.17	118.72	124.53
22	A	841	CLA	O2A-C1-C2	5.17	122.22	108.64
22	3	308	CLA	O2D-CGD-CBD	5.17	120.45	111.27
38	3	303	LUT	C7-C8-C9	-5.16	118.43	126.23
22	3	314	CLA	CMD-C2D-C1D	5.16	133.81	124.71
22	B	839	CLA	O2D-CGD-CBD	5.16	120.44	111.27
22	B	813	CLA	O2A-C1-C2	5.16	122.20	108.64
22	8	305	CLA	O2D-CGD-CBD	5.16	120.44	111.27
22	B	817	CLA	CMD-C2D-C1D	5.16	133.81	124.71
22	B	830	CLA	CMD-C2D-C1D	5.16	133.81	124.71
22	3	320	CLA	CMD-C2D-C1D	5.16	133.80	124.71
22	4	818	CLA	O2D-CGD-CBD	5.15	120.43	111.27
22	A	838	CLA	O2D-CGD-CBD	5.15	120.42	111.27
25	B	851	LHG	O7-C7-O9	-5.15	119.01	125.57
22	6	302	CLA	CMD-C2D-C1D	5.15	133.79	124.71
22	Z	314	CLA	CMD-C2D-C1D	5.15	133.79	124.71
22	3	316	CLA	O2D-CGD-CBD	5.15	120.42	111.27
22	A	813	CLA	O2D-CGD-CBD	5.15	120.41	111.27
22	B	827	CLA	O2D-CGD-CBD	5.15	120.41	111.27
22	B	819	CLA	O2D-CGD-CBD	5.14	120.41	111.27
22	2	304	CLA	CMD-C2D-C1D	5.14	133.78	124.71
22	5	326	CLA	O2D-CGD-CBD	5.14	120.41	111.27
22	6	301	CLA	O2D-CGD-CBD	5.14	120.40	111.27
38	1	302	LUT	C7-C8-C9	-5.14	118.47	126.23
24	7	303	BCR	C1-C6-C5	-5.14	115.38	122.61
38	1	302	LUT	C1-C6-C5	-5.14	115.38	122.61
22	K	202	CLA	O2D-CGD-CBD	5.13	120.39	111.27
21	A	801	CL0	CHD-C1D-ND	-5.13	119.74	124.45
22	B	822	CLA	O2D-CGD-CBD	5.13	120.38	111.27
22	5	315	CLA	CMD-C2D-C1D	5.13	133.75	124.71
22	Z	305	CLA	O2D-CGD-CBD	5.13	120.38	111.27
22	B	829	CLA	O2A-C1-C2	5.12	122.10	108.64
22	8	311	CLA	CMD-C2D-C1D	5.12	133.74	124.71
22	A	810	CLA	O2A-C1-C2	5.12	122.10	108.64
22	F	301	CLA	O2D-CGD-CBD	5.12	120.36	111.27
22	5	310	CLA	O2D-CGD-CBD	5.12	120.36	111.27
22	1	313	CLA	O2D-CGD-CBD	5.12	120.36	111.27
22	B	815	CLA	O2D-CGD-CBD	5.11	120.36	111.27
24	B	845	BCR	C27-C26-C25	-5.11	115.31	122.73
22	G	201	CLA	O2D-CGD-CBD	5.11	120.35	111.27
36	J	105	C7Z	C31-C30-C29	-5.11	120.01	127.31

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Mol	Chain	Res	Type	Atoms	Z	Observed(^{o})	Ideal(^{o})
22	A	839	CLA	O2D-CGD-CBD	5.11	120.35	111.27
22	A	817	CLA	O2A-C1-C2	5.11	122.06	108.64
22	6	317	CLA	CMD-C2D-C1D	5.11	133.71	124.71
36	5	306	C7Z	C35-C34-C33	-5.11	120.02	127.31
32	F	306	RRX	C33-C5-C6	-5.10	118.80	124.53
22	6	312	CLA	O2D-CGD-CBD	5.10	120.33	111.27
22	A	837	CLA	O2D-CGD-CBD	5.10	120.33	111.27
22	8	313	CLA	O2D-CGD-CBD	5.10	120.33	111.27
24	B	848	BCR	C7-C8-C9	-5.10	118.54	126.23
22	9	306	CLA	O2D-CGD-CBD	5.09	120.32	111.27
22	B	840	CLA	O2D-CGD-CBD	5.09	120.31	111.27
22	B	841	CLA	O2D-CGD-CBD	5.09	120.31	111.27
24	7	303	BCR	C4-C5-C6	-5.09	115.34	122.73
22	7	307	CLA	O2A-C1-C2	5.09	122.00	108.64
22	B	824	CLA	O2D-CGD-CBD	5.09	120.30	111.27
22	B	828	CLA	O2D-CGD-CBD	5.09	120.30	111.27
22	B	832	CLA	O2D-CGD-CBD	5.08	120.30	111.27
24	F	303	BCR	C20-C21-C22	-5.08	120.06	127.31
22	A	808	CLA	O2D-CGD-CBD	5.08	120.30	111.27
22	4	812	CLA	O2D-CGD-CBD	5.08	120.30	111.27
22	7	315	CLA	CMD-C2D-C1D	5.08	133.67	124.71
22	A	841	CLA	O2D-CGD-CBD	5.08	120.29	111.27
22	3	318	CLA	O2D-CGD-CBD	5.08	120.29	111.27
22	4	817	CLA	O2D-CGD-CBD	5.08	120.29	111.27
38	3	303	LUT	C15-C14-C13	-5.07	120.07	127.31
22	Z	309	CLA	CMD-C2D-C1D	5.07	133.65	124.71
22	A	818	CLA	O2D-CGD-CBD	5.07	120.28	111.27
22	4	808	CLA	O2D-CGD-CBD	5.07	120.28	111.27
22	4	815	CLA	CMD-C2D-C1D	5.07	133.65	124.71
22	7	316	CLA	O2D-CGD-CBD	5.07	120.27	111.27
22	A	819	CLA	O2D-CGD-CBD	5.07	120.27	111.27
22	A	831	CLA	O2D-CGD-CBD	5.07	120.27	111.27
22	A	802	CLA	O2D-CGD-CBD	5.06	120.27	111.27
22	A	814	CLA	O2D-CGD-CBD	5.06	120.27	111.27
22	8	307	CLA	O2D-CGD-CBD	5.06	120.27	111.27
22	5	322	CLA	O2A-C1-C2	5.06	121.94	108.64
22	6	319	CLA	O2A-C1-C2	5.06	121.94	108.64
22	7	304	CLA	O2D-CGD-CBD	5.06	120.26	111.27
22	B	805	CLA	O2D-CGD-CBD	5.06	120.25	111.27
22	G	202	CLA	O2D-CGD-CBD	5.06	120.25	111.27
22	1	305	CLA	O2D-CGD-CBD	5.05	120.25	111.27
22	4	806	CLA	O2D-CGD-CBD	5.05	120.25	111.27

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Mol	Chain	Res	Type	Atoms	Z	Observed(^{o})	Ideal(^{o})
22	6	317	CLA	O2D-CGD-CBD	5.05	120.25	111.27
22	B	818	CLA	O2D-CGD-CBD	5.05	120.25	111.27
22	B	808	CLA	O2D-CGD-CBD	5.05	120.25	111.27
21	A	801	CL0	C3D-C2D-C1D	-5.05	98.94	105.83
22	7	314	CLA	O2D-CGD-CBD	5.05	120.24	111.27
22	A	838	CLA	O2A-C1-C2	5.05	121.90	108.64
22	3	319	CLA	O2D-CGD-CBD	5.05	120.24	111.27
22	B	806	CLA	O2D-CGD-CBD	5.05	120.24	111.27
24	5	305	BCR	C1-C6-C5	-5.05	115.50	122.61
24	6	305	BCR	C16-C17-C18	-5.05	120.11	127.31
21	A	801	CL0	O2A-CGA-O1A	-5.04	110.86	123.59
34	F	308	LMG	O7-C10-C11	5.04	122.36	111.50
22	B	841	CLA	O2A-C1-C2	5.04	121.88	108.64
22	6	310	CLA	O2D-CGD-CBD	5.04	120.22	111.27
24	6	306	BCR	C38-C26-C25	-5.04	118.87	124.53
22	4	811	CLA	O2D-CGD-CBD	5.04	120.22	111.27
24	8	304	BCR	C15-C14-C13	-5.04	120.12	127.31
22	5	319	CLA	O2D-CGD-CBD	5.03	120.21	111.27
22	3	311	CLA	O2D-CGD-CBD	5.03	120.21	111.27
24	A	845	BCR	C11-C10-C9	-5.03	120.13	127.31
22	6	313	CLA	CMD-C2D-C1D	5.03	133.57	124.71
24	7	303	BCR	C20-C21-C22	-5.03	120.14	127.31
22	A	822	CLA	O2A-C1-C2	5.03	121.85	108.64
22	A	825	CLA	O2D-CGD-CBD	5.03	120.20	111.27
24	K	206	BCR	C20-C21-C22	-5.02	120.14	127.31
22	3	322	CLA	O2D-CGD-CBD	5.02	120.19	111.27
24	A	848	BCR	C15-C14-C13	-5.02	120.14	127.31
24	3	307	BCR	C4-C5-C6	-5.02	115.44	122.73
22	A	829	CLA	O2A-C1-C2	5.02	121.83	108.64
38	1	303	LUT	C35-C34-C33	-5.02	120.14	127.31
22	B	809	CLA	O2A-C1-C2	5.02	121.83	108.64
22	8	308	CLA	O2D-CGD-CBD	5.02	120.19	111.27
22	1	310	CLA	O2A-C1-C2	5.02	121.83	108.64
22	6	301	CLA	O2A-C1-C2	5.02	121.82	108.64
22	A	855	CLA	O2D-CGD-CBD	5.01	120.18	111.27
22	A	807	CLA	O2D-CGD-CBD	5.01	120.17	111.27
22	5	301	CLA	O2D-CGD-CBD	5.01	120.17	111.27
24	B	844	BCR	C20-C19-C18	-5.01	112.34	126.42
22	B	836	CLA	O2D-CGD-CBD	5.01	120.17	111.27
22	8	307	CLA	O2A-C1-C2	5.01	121.80	108.64
22	B	816	CLA	O2D-CGD-CBD	5.00	120.16	111.27
38	Z	302	LUT	C11-C10-C9	-5.00	120.17	127.31

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Mol	Chain	Res	Type	Atoms	Z	Observed(^{o})	Ideal(^{o})
22	3	315	CLA	O2D-CGD-CBD	5.00	120.16	111.27
22	A	835	CLA	O2D-CGD-CBD	5.00	120.16	111.27
22	Z	308	CLA	O2D-CGD-CBD	5.00	120.16	111.27
24	B	849	BCR	C7-C8-C9	-5.00	118.68	126.23
22	K	205	CLA	O2A-C1-C2	5.00	121.78	108.64
22	A	812	CLA	O2D-CGD-CBD	5.00	120.15	111.27
22	1	309	CLA	O2D-CGD-CBD	5.00	120.15	111.27
24	4	804	BCR	C16-C17-C18	-5.00	120.18	127.31
22	A	805	CLA	O2D-CGD-CBD	5.00	120.15	111.27
22	5	315	CLA	O2D-CGD-CBD	5.00	120.15	111.27
22	1	308	CLA	O2A-C1-C2	5.00	121.76	108.64
24	K	206	BCR	C16-C17-C18	-5.00	120.18	127.31
22	F	304	CLA	O2A-C1-C2	4.99	121.75	108.64
22	F	305	CLA	O2D-CGD-CBD	4.99	120.14	111.27
22	Z	309	CLA	O2D-CGD-CBD	4.99	120.13	111.27
22	A	830	CLA	O2A-C1-C2	4.99	121.74	108.64
22	4	809	CLA	O2D-CGD-CBD	4.99	120.13	111.27
22	A	822	CLA	O2D-CGD-CBD	4.99	120.13	111.27
22	B	807	CLA	O2D-CGD-CBD	4.99	120.13	111.27
22	4	806	CLA	O2A-C1-C2	4.99	121.74	108.64
22	5	310	CLA	O2A-C1-C2	4.99	121.74	108.64
24	A	845	BCR	C20-C21-C22	-4.98	120.20	127.31
22	A	827	CLA	O2D-CGD-CBD	4.98	120.12	111.27
22	6	321	CLA	O2D-CGD-CBD	4.98	120.12	111.27
22	G	201	CLA	O2A-C1-C2	4.98	121.72	108.64
22	5	307	CLA	O2D-CGD-CBD	4.98	120.11	111.27
22	B	815	CLA	O2A-C1-C2	4.97	121.71	108.64
24	4	804	BCR	C24-C23-C22	-4.97	118.72	126.23
22	J	103	CLA	O2D-CGD-CBD	4.97	120.11	111.27
22	L	202	CLA	O2D-CGD-CBD	4.97	120.11	111.27
22	Z	313	CLA	CMD-C2D-C1D	4.97	133.48	124.71
38	9	301	LUT	C35-C34-C33	-4.97	120.22	127.31
24	A	845	BCR	C33-C5-C6	-4.97	118.95	124.53
38	4	803	LUT	C7-C8-C9	-4.97	118.73	126.23
22	1	306	CLA	O2D-CGD-CBD	4.97	120.09	111.27
22	2	304	CLA	O2D-CGD-CBD	4.97	120.09	111.27
22	7	324	CLA	O2D-CGD-CBD	4.97	120.09	111.27
22	A	806	CLA	O2D-CGD-CBD	4.96	120.09	111.27
22	4	810	CLA	O2D-CGD-CBD	4.96	120.09	111.27
24	L	204	BCR	C1-C6-C5	-4.96	115.63	122.61
22	F	302	CLA	O2D-CGD-CBD	4.96	120.08	111.27
39	9	312	CHL	C4A-NA-C1A	4.96	108.94	106.71

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Mol	Chain	Res	Type	Atoms	Z	Observed(^{o})	Ideal(^{o})
22	5	314	CLA	O2D-CGD-CBD	4.96	120.08	111.27
22	1	304	CLA	O2D-CGD-CBD	4.96	120.07	111.27
22	1	315	CLA	O2D-CGD-CBD	4.96	120.07	111.27
22	Z	304	CLA	CMD-C2D-C1D	4.95	133.44	124.71
22	A	815	CLA	O2D-CGD-CBD	4.95	120.07	111.27
22	3	314	CLA	O2D-CGD-CBD	4.95	120.07	111.27
22	A	832	CLA	O2D-CGD-CBD	4.95	120.07	111.27
24	B	846	BCR	C33-C5-C6	-4.95	118.97	124.53
22	8	309	CLA	O2A-C1-C2	4.95	121.64	108.64
22	A	821	CLA	O2D-CGD-CBD	4.95	120.06	111.27
38	8	302	LUT	C18-C5-C6	-4.95	118.97	124.53
22	F	304	CLA	O2D-CGD-CBD	4.94	120.05	111.27
22	7	322	CLA	O2D-CGD-CBD	4.94	120.05	111.27
38	4	802	LUT	C35-C34-C33	-4.94	120.26	127.31
22	B	828	CLA	O2A-C1-C2	4.94	121.61	108.64
22	Z	315	CLA	O2D-CGD-CBD	4.94	120.04	111.27
22	L	201	CLA	O2D-CGD-CBD	4.93	120.03	111.27
22	1	307	CLA	O2D-CGD-CBD	4.93	120.03	111.27
22	B	842	CLA	O2D-CGD-CBD	4.93	120.03	111.27
22	8	314	CLA	O2D-CGD-CBD	4.93	120.03	111.27
22	B	823	CLA	O2D-CGD-CBD	4.93	120.02	111.27
22	3	308	CLA	O2A-C1-C2	4.93	121.58	108.64
22	B	806	CLA	O2A-C1-C2	4.93	121.58	108.64
22	Z	314	CLA	O2A-C1-C2	4.93	121.58	108.64
38	1	303	LUT	C11-C10-C9	-4.92	120.28	127.31
22	B	814	CLA	O2A-C1-C2	4.92	121.58	108.64
22	K	205	CLA	O2D-CGD-CBD	4.92	120.02	111.27
22	F	301	CLA	O2A-C1-C2	4.92	121.57	108.64
22	7	308	CLA	O2D-CGD-CBD	4.92	120.01	111.27
38	1	303	LUT	C7-C8-C9	-4.92	118.80	126.23
22	Z	308	CLA	O2A-C1-C2	4.92	121.56	108.64
22	1	310	CLA	O2D-CGD-CBD	4.92	120.01	111.27
22	B	811	CLA	O2A-C1-C2	4.92	121.56	108.64
24	B	847	BCR	C7-C8-C9	-4.92	118.81	126.23
22	A	836	CLA	O2D-CGD-CBD	4.91	120.00	111.27
22	5	309	CLA	O2D-CGD-CBD	4.91	120.00	111.27
22	B	833	CLA	O2D-CGD-CBD	4.91	120.00	111.27
22	A	820	CLA	O2D-CGD-CBD	4.91	120.00	111.27
22	9	307	CLA	O2A-C1-C2	4.91	121.54	108.64
22	B	830	CLA	O2A-C1-C2	4.91	121.54	108.64
22	6	313	CLA	O2D-CGD-CBD	4.91	119.99	111.27
22	B	831	CLA	O2A-C1-C2	4.91	121.53	108.64

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Mol	Chain	Res	Type	Atoms	Z	Observed(^{o})	Ideal(^{o})
22	B	809	CLA	O2D-CGD-CBD	4.90	119.98	111.27
22	B	837	CLA	O2A-C1-C2	4.90	121.53	108.64
22	B	835	CLA	O2A-C1-C2	4.90	121.52	108.64
22	Z	314	CLA	O2D-CGD-CBD	4.90	119.98	111.27
22	A	840	CLA	O2A-C1-C2	4.90	121.52	108.64
22	8	310	CLA	O2D-CGD-CBD	4.90	119.98	111.27
22	7	312	CLA	O2D-CGD-CBD	4.90	119.97	111.27
22	A	817	CLA	O2D-CGD-CBD	4.89	119.97	111.27
24	B	848	BCR	C11-C10-C9	-4.89	120.33	127.31
38	6	304	LUT	C15-C14-C13	-4.89	120.33	127.31
22	B	839	CLA	O2A-C1-C2	4.89	121.49	108.64
24	3	306	BCR	C30-C25-C26	-4.89	115.72	122.61
22	A	830	CLA	O2D-CGD-CBD	4.89	119.96	111.27
22	B	813	CLA	O2D-CGD-CBD	4.89	119.95	111.27
22	5	313	CLA	O2D-CGD-CBD	4.89	119.95	111.27
22	A	828	CLA	O2D-CGD-CBD	4.89	119.95	111.27
22	A	823	CLA	O2D-CGD-CBD	4.89	119.95	111.27
22	7	311	CLA	O2D-CGD-CBD	4.89	119.95	111.27
22	6	309	CLA	O2D-CGD-CBD	4.89	119.95	111.27
22	3	310	CLA	O2A-C1-C2	4.88	121.47	108.64
22	B	838	CLA	O2D-CGD-CBD	4.88	119.94	111.27
22	A	824	CLA	O2D-CGD-CBD	4.88	119.94	111.27
22	A	810	CLA	O2D-CGD-CBD	4.87	119.93	111.27
22	2	301	CLA	O2A-C1-C2	4.87	121.44	108.64
22	7	309	CLA	O2A-C1-C2	4.87	121.43	108.64
38	6	304	LUT	C35-C34-C33	-4.87	120.36	127.31
22	B	801	CLA	O2D-CGD-CBD	4.87	119.92	111.27
24	6	306	BCR	C7-C8-C9	-4.87	118.88	126.23
22	3	310	CLA	O2D-CGD-CBD	4.87	119.92	111.27
22	7	308	CLA	O2A-C1-C2	4.87	121.43	108.64
24	J	104	BCR	C20-C21-C22	-4.87	120.37	127.31
22	6	302	CLA	O2D-CGD-CBD	4.86	119.91	111.27
22	B	824	CLA	O2A-C1-C2	4.86	121.41	108.64
22	6	313	CLA	O2A-C1-C2	4.86	121.41	108.64
22	B	816	CLA	O2A-C1-C2	4.86	121.41	108.64
22	3	318	CLA	O2A-C1-C2	4.86	121.40	108.64
22	B	812	CLA	O2A-C1-C2	4.86	121.40	108.64
22	L	203	CLA	O2D-CGD-CBD	4.86	119.90	111.27
22	6	311	CLA	O2A-C1-C2	4.86	121.40	108.64
24	A	844	BCR	C33-C5-C6	-4.86	119.08	124.53
24	8	304	BCR	C11-C10-C9	-4.85	120.38	127.31
22	7	305	CLA	O2A-C1-C2	4.85	121.39	108.64

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Mol	Chain	Res	Type	Atoms	Z	Observed(^{o})	Ideal(^{o})
22	A	826	CLA	O2A-C1-C2	4.85	121.39	108.64
22	9	311	CLA	O2D-CGD-CBD	4.85	119.89	111.27
22	B	810	CLA	O2A-C1-C2	4.85	121.39	108.64
24	A	844	BCR	C12-C13-C14	4.85	126.39	118.94
22	7	317	CLA	O2A-C1-C2	4.85	121.38	108.64
22	5	308	CLA	O2A-C1-C2	4.85	121.38	108.64
22	K	203	CLA	O2D-CGD-CBD	4.85	119.88	111.27
22	2	301	CLA	O2D-CGD-CBD	4.85	119.88	111.27
22	L	203	CLA	O2A-C1-C2	4.85	121.37	108.64
32	F	306	RRX	C1-C6-C5	-4.84	115.79	122.61
22	B	821	CLA	O2D-CGD-CBD	4.84	119.88	111.27
21	A	801	CL0	O2A-C1-C2	4.84	121.37	108.64
38	5	303	LUT	C11-C10-C9	-4.84	120.40	127.31
24	I	4001	BCR	C1-C6-C5	-4.84	115.80	122.61
24	G	203	BCR	C38-C26-C25	-4.84	119.10	124.53
24	K	206	BCR	C7-C8-C9	-4.84	118.93	126.23
24	A	844	BCR	C38-C26-C25	-4.83	119.10	124.53
22	3	322	CLA	O2A-C1-C2	4.83	121.34	108.64
22	K	204	CLA	O2D-CGD-CBD	4.83	119.86	111.27
39	9	312	CHL	CMA-C3A-C4A	4.83	124.75	111.77
36	5	306	C7Z	C38-C25-C26	-4.83	119.11	124.53
22	7	309	CLA	O2D-CGD-CBD	4.83	119.84	111.27
22	A	842	CLA	O2D-CGD-CBD	4.82	119.84	111.27
22	A	842	CLA	O2A-C1-C2	4.82	121.31	108.64
22	B	829	CLA	O2D-CGD-CBD	4.82	119.84	111.27
22	5	323	CLA	O2D-CGD-CBD	4.82	119.83	111.27
24	A	844	BCR	C8-C9-C10	4.82	126.33	118.94
22	A	854	CLA	O2D-CGD-CBD	4.82	119.83	111.27
24	A	844	BCR	C15-C16-C17	-4.81	113.61	123.47
22	6	322	CLA	O2A-C1-C2	4.81	121.28	108.64
38	Z	301	LUT	C15-C14-C13	-4.81	120.45	127.31
22	B	812	CLA	O2D-CGD-CBD	4.81	119.81	111.27
24	B	845	BCR	C16-C17-C18	-4.81	120.45	127.31
22	4	815	CLA	O2A-C1-C2	4.81	121.27	108.64
22	1	316	CLA	CMD-C2D-C1D	4.81	133.18	124.71
24	B	846	BCR	C7-C8-C9	-4.80	118.98	126.23
22	Z	310	CLA	O2D-CGD-CBD	4.80	119.80	111.27
22	B	826	CLA	O2A-C1-C2	4.80	121.25	108.64
24	A	845	BCR	C1-C6-C5	-4.80	115.86	122.61
24	B	844	BCR	C19-C18-C17	4.80	126.30	118.94
22	1	314	CLA	O2D-CGD-CBD	4.79	119.78	111.27
22	7	306	CLA	O2D-CGD-CBD	4.79	119.78	111.27

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Mol	Chain	Res	Type	Atoms	Z	Observed(^{o})	Ideal(^{o})
22	6	311	CLA	O2D-CGD-CBD	4.79	119.78	111.27
22	Z	304	CLA	O2D-CGD-CBD	4.79	119.78	111.27
24	J	104	BCR	C38-C26-C25	-4.79	119.15	124.53
22	6	317	CLA	O2A-C1-C2	4.79	121.22	108.64
22	8	311	CLA	O2D-CGD-CBD	4.79	119.77	111.27
22	9	309	CLA	O2A-C1-C2	4.78	121.21	108.64
38	Z	302	LUT	C7-C8-C9	-4.78	119.01	126.23
38	6	304	LUT	C7-C8-C9	-4.78	119.01	126.23
24	A	856	BCR	C38-C26-C25	-4.78	119.16	124.53
22	B	825	CLA	O2D-CGD-CBD	4.78	119.76	111.27
22	A	815	CLA	O2A-C1-C2	4.78	121.20	108.64
22	A	855	CLA	O2A-C1-C2	4.78	121.19	108.64
24	B	853	BCR	C20-C21-C22	-4.78	120.49	127.31
22	B	818	CLA	O2A-C1-C2	4.78	121.19	108.64
24	J	104	BCR	C4-C5-C6	-4.78	115.80	122.73
24	8	304	BCR	C7-C8-C9	-4.77	119.02	126.23
22	9	313	CLA	O2A-C1-C2	4.77	121.18	108.64
24	3	305	BCR	C27-C26-C25	-4.77	115.80	122.73
22	4	807	CLA	O2A-C1-C2	4.77	121.18	108.64
22	7	317	CLA	O2D-CGD-CBD	4.77	119.75	111.27
22	3	309	CLA	O2D-CGD-CBD	4.77	119.75	111.27
22	8	308	CLA	O2A-C1-C2	4.77	121.16	108.64
22	9	307	CLA	O2D-CGD-CBD	4.76	119.73	111.27
22	6	312	CLA	O2A-C1-C2	4.76	121.16	108.64
22	A	839	CLA	O2A-C1-C2	4.76	121.14	108.64
22	A	840	CLA	O2D-CGD-CBD	4.76	119.72	111.27
24	7	303	BCR	C15-C14-C13	-4.76	120.52	127.31
22	B	819	CLA	O2A-C1-C2	4.75	121.13	108.64
22	A	816	CLA	O2A-C1-C2	4.75	121.12	108.64
24	8	304	BCR	C4-C5-C6	-4.75	115.83	122.73
22	Z	316	CLA	O2A-C1-C2	4.75	121.12	108.64
36	J	105	C7Z	C11-C10-C9	-4.75	120.53	127.31
22	6	307	CLA	O2D-CGD-CBD	4.75	119.71	111.27
22	A	835	CLA	O2A-C1-C2	4.75	121.11	108.64
24	5	305	BCR	C38-C26-C25	-4.75	119.20	124.53
22	9	304	CLA	CMD-C2D-C1D	4.75	133.08	124.71
24	6	305	BCR	C24-C23-C22	-4.74	119.07	126.23
38	5	303	LUT	C15-C14-C13	-4.74	120.54	127.31
22	3	319	CLA	O2A-C1-C2	4.74	121.09	108.64
22	A	826	CLA	O2D-CGD-CBD	4.73	119.67	111.27
38	7	302	LUT	C7-C8-C9	-4.73	119.09	126.23
22	7	309	CLA	C1-C2-C3	-4.73	117.86	126.04

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Mol	Chain	Res	Type	Atoms	Z	Observed(^{o})	Ideal(^{o})
22	1	314	CLA	O2A-C1-C2	4.73	121.06	108.64
22	9	310	CLA	O2D-CGD-CBD	4.73	119.67	111.27
22	8	316	CLA	O2D-CGD-CBD	4.72	119.66	111.27
24	3	305	BCR	C30-C25-C26	-4.72	115.96	122.61
38	3	302	LUT	C11-C10-C9	-4.72	120.57	127.31
24	4	804	BCR	C15-C14-C13	-4.72	120.58	127.31
22	A	809	CLA	O2A-C1-C2	4.71	121.02	108.64
22	8	309	CLA	O2D-CGD-CBD	4.71	119.64	111.27
22	5	312	CLA	O2D-CGD-CBD	4.71	119.64	111.27
22	Z	316	CLA	O2D-CGD-CBD	4.71	119.63	111.27
22	A	817	CLA	C1-C2-C3	-4.70	117.91	126.04
22	B	808	CLA	O2A-C1-C2	4.70	120.99	108.64
24	3	307	BCR	C7-C8-C9	-4.70	119.13	126.23
38	5	302	LUT	C18-C5-C6	-4.70	119.25	124.53
22	B	840	CLA	O2A-C1-C2	4.70	120.98	108.64
24	4	804	BCR	C38-C26-C25	-4.69	119.26	124.53
22	B	805	CLA	O2A-C1-C2	4.69	120.97	108.64
22	Z	306	CLA	O2D-CGD-CBD	4.69	119.61	111.27
22	A	803	CLA	O2A-C1-C2	4.69	120.96	108.64
22	7	314	CLA	O2A-C1-C2	4.69	120.96	108.64
22	3	312	CLA	O2A-C1-C2	4.69	120.96	108.64
38	4	802	LUT	C15-C14-C13	-4.69	120.62	127.31
24	L	204	BCR	C38-C26-C25	-4.69	119.26	124.53
22	9	304	CLA	O2D-CGD-CBD	4.69	119.60	111.27
31	1	319	DGD	O2G-C1B-C2B	4.69	121.60	111.50
38	5	303	LUT	C7-C8-C9	-4.69	119.15	126.23
22	Z	307	CLA	O2A-C1-C2	4.69	120.95	108.64
22	8	310	CLA	C1-C2-C3	-4.68	117.94	126.04
22	L	202	CLA	O2A-C1-C2	4.68	120.94	108.64
22	6	314	CLA	O2A-C1-C2	4.68	120.94	108.64
22	Z	305	CLA	CMD-C2D-C1D	4.68	132.96	124.71
22	6	302	CLA	O2A-C1-C2	4.68	120.93	108.64
22	A	829	CLA	O2D-CGD-CBD	4.68	119.58	111.27
38	Z	302	LUT	C15-C14-C13	-4.67	120.64	127.31
22	B	827	CLA	O2A-C1-C2	4.67	120.90	108.64
22	1	307	CLA	O2A-C1-C2	4.67	120.90	108.64
24	B	844	BCR	C33-C5-C6	-4.66	119.29	124.53
22	A	811	CLA	O2D-CGD-CBD	4.66	119.55	111.27
24	L	204	BCR	C27-C26-C25	-4.66	115.97	122.73
22	5	322	CLA	O2D-CGD-CBD	4.66	119.54	111.27
22	3	313	CLA	O2D-CGD-CBD	4.65	119.54	111.27
22	3	312	CLA	O2D-CGD-CBD	4.65	119.53	111.27

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Mol	Chain	Res	Type	Atoms	Z	Observed(^{o})	Ideal(^{o})
24	8	304	BCR	C16-C17-C18	-4.65	120.67	127.31
38	7	302	LUT	C35-C34-C33	-4.65	120.68	127.31
24	B	847	BCR	C27-C26-C25	-4.64	115.99	122.73
24	6	305	BCR	C38-C26-C25	-4.64	119.32	124.53
22	7	306	CLA	O2A-C1-C2	4.64	120.82	108.64
22	Z	309	CLA	O2A-C1-C2	4.64	120.82	108.64
24	3	305	BCR	C38-C26-C25	-4.63	119.33	124.53
22	6	308	CLA	O2A-C1-C2	4.63	120.80	108.64
38	3	302	LUT	C7-C8-C9	-4.63	119.24	126.23
22	1	311	CLA	CMD-C2D-C1D	4.62	132.86	124.71
22	L	201	CLA	O2A-C1-C2	4.62	120.78	108.64
24	A	856	BCR	C33-C5-C6	-4.62	119.34	124.53
36	5	306	C7Z	C7-C8-C9	-4.62	119.26	126.23
22	B	821	CLA	O2A-C1-C2	4.62	120.77	108.64
22	A	805	CLA	O2A-C1-C2	4.62	120.77	108.64
22	K	203	CLA	O2A-C1-C2	4.62	120.76	108.64
22	A	825	CLA	O2A-C1-C2	4.61	120.76	108.64
22	B	825	CLA	O2A-C1-C2	4.61	120.75	108.64
24	3	305	BCR	C24-C23-C22	-4.61	119.27	126.23
22	B	822	CLA	O2A-C1-C2	4.61	120.74	108.64
24	B	844	BCR	C7-C8-C9	-4.60	119.28	126.23
24	F	303	BCR	C33-C5-C6	-4.60	119.37	124.53
22	B	830	CLA	O2D-CGD-CBD	4.59	119.43	111.27
38	3	303	LUT	C11-C10-C9	-4.59	120.76	127.31
22	6	307	CLA	O2A-C1-C2	4.59	120.69	108.64
22	4	810	CLA	O2A-C1-C2	4.59	120.69	108.64
38	4	803	LUT	C11-C10-C9	-4.59	120.77	127.31
22	9	308	CLA	O2A-C1-C2	4.58	120.69	108.64
22	A	834	CLA	O2A-C1-C2	4.58	120.68	108.64
31	3	301	DGD	O2G-C1B-C2B	4.58	121.37	111.50
24	5	305	BCR	C15-C16-C17	-4.58	114.09	123.47
22	7	310	CLA	O2A-C1-C2	4.58	120.67	108.64
24	B	844	BCR	C11-C12-C13	-4.58	113.56	126.42
22	B	842	CLA	O2A-C1-C2	4.57	120.65	108.64
38	1	303	LUT	C15-C14-C13	-4.57	120.79	127.31
22	Z	305	CLA	O2A-C1-C2	4.57	120.64	108.64
24	L	204	BCR	C4-C5-C6	-4.57	116.10	122.73
24	B	849	BCR	C38-C26-C25	-4.57	119.40	124.53
24	A	844	BCR	C27-C26-C25	-4.56	116.11	122.73
38	9	301	LUT	C31-C30-C29	-4.56	120.80	127.31
22	1	304	CLA	O2A-C1-C2	4.56	120.62	108.64
24	J	104	BCR	C33-C5-C6	-4.56	119.41	124.53

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Mol	Chain	Res	Type	Atoms	Z	Observed(^{o})	Ideal(^{o})
22	A	819	CLA	O2A-C1-C2	4.56	120.61	108.64
22	A	854	CLA	O2A-C1-C2	4.55	120.60	108.64
22	B	833	CLA	O2A-C1-C2	4.55	120.60	108.64
24	3	306	BCR	C16-C17-C18	-4.55	120.81	127.31
22	A	833	CLA	O2A-C1-C2	4.55	120.59	108.64
24	A	848	BCR	C1-C6-C5	-4.55	116.21	122.61
24	5	304	BCR	C24-C23-C22	-4.54	119.37	126.23
22	5	318	CLA	O2A-C1-C2	4.54	120.57	108.64
22	7	315	CLA	O2A-C1-C2	4.54	120.57	108.64
24	A	856	BCR	C15-C14-C13	-4.54	120.83	127.31
22	6	310	CLA	O2A-C1-C2	4.54	120.57	108.64
38	9	301	LUT	C7-C8-C9	-4.54	119.38	126.23
22	7	312	CLA	O2A-C1-C2	4.54	120.56	108.64
22	B	834	CLA	O2A-C1-C2	4.54	120.56	108.64
24	A	844	BCR	C20-C21-C22	-4.54	120.83	127.31
22	5	318	CLA	O2D-CGD-CBD	4.54	119.33	111.27
22	7	322	CLA	O2A-C1-C2	4.53	120.55	108.64
22	A	806	CLA	O2A-C1-C2	4.53	120.54	108.64
38	Z	302	LUT	C35-C34-C33	-4.53	120.85	127.31
38	5	303	LUT	C35-C34-C33	-4.53	120.85	127.31
39	3	317	CHL	CHD-C1D-ND	-4.52	120.30	124.45
24	A	856	BCR	C24-C23-C22	-4.52	119.41	126.23
22	A	823	CLA	O2A-C1-C2	4.52	120.50	108.64
22	8	313	CLA	O2A-C1-C2	4.52	120.50	108.64
24	B	848	BCR	C38-C26-C25	-4.51	119.46	124.53
22	8	306	CLA	O2A-C1-C2	4.51	120.49	108.64
24	G	203	BCR	C15-C14-C13	-4.51	120.87	127.31
38	Z	301	LUT	C35-C34-C33	-4.51	120.87	127.31
22	8	311	CLA	O2A-C1-C2	4.51	120.48	108.64
24	K	206	BCR	C30-C25-C26	-4.50	116.27	122.61
22	4	807	CLA	O2D-CGD-CBD	4.50	119.27	111.27
22	Z	303	CLA	O2A-C1-C2	4.50	120.46	108.64
22	5	312	CLA	O2A-C1-C2	4.50	120.46	108.64
24	5	305	BCR	C33-C5-C6	-4.50	119.48	124.53
24	B	845	BCR	C38-C26-C25	-4.50	119.48	124.53
24	A	846	BCR	C24-C23-C22	-4.50	119.44	126.23
22	A	837	CLA	O2A-C1-C2	4.49	120.44	108.64
24	B	848	BCR	C24-C23-C22	-4.49	119.45	126.23
22	A	821	CLA	O2A-C1-C2	4.49	120.43	108.64
24	3	304	BCR	C33-C5-C6	-4.49	119.49	124.53
22	B	801	CLA	O2A-C1-C2	4.49	120.43	108.64
22	5	311	CLA	O2A-C1-C2	4.48	120.42	108.64

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Mol	Chain	Res	Type	Atoms	Z	Observed(^{o})	Ideal(^{o})
22	9	305	CLA	O2A-C1-C2	4.48	120.42	108.64
24	A	856	BCR	C1-C6-C5	-4.48	116.30	122.61
22	5	315	CLA	O2A-C1-C2	4.48	120.41	108.64
22	A	808	CLA	O2A-C1-C2	4.48	120.40	108.64
22	4	808	CLA	O2A-C1-C2	4.48	120.40	108.64
22	9	310	CLA	CMD-C2D-C1D	4.47	132.60	124.71
22	A	811	CLA	O2A-C1-C2	4.47	120.38	108.64
24	B	847	BCR	C33-C5-C6	-4.47	119.51	124.53
22	A	831	CLA	O2A-C1-C2	4.46	120.36	108.64
22	B	835	CLA	C1-C2-C3	-4.46	119.53	126.75
24	J	104	BCR	C1-C6-C5	-4.46	116.33	122.61
32	F	306	RRX	C7-C8-C9	-4.46	119.50	126.23
22	A	813	CLA	O2A-C1-C2	4.46	120.35	108.64
24	8	304	BCR	C33-C5-C6	-4.46	119.52	124.53
22	3	316	CLA	O2A-C1-C2	4.45	120.34	108.64
22	A	828	CLA	O2A-C1-C2	4.45	120.34	108.64
22	B	832	CLA	O2A-C1-C2	4.45	120.33	108.64
22	3	311	CLA	O2A-C1-C2	4.45	120.33	108.64
22	5	320	CLA	O2A-C1-C2	4.45	120.33	108.64
22	4	809	CLA	O2A-C1-C2	4.45	120.33	108.64
25	B	803	LHG	O7-C7-C8	4.44	121.08	111.50
22	3	313	CLA	O2A-C1-C2	4.44	120.31	108.64
39	8	315	CHL	CHD-C1D-ND	-4.44	120.37	124.45
24	B	853	BCR	C16-C17-C18	-4.44	120.97	127.31
22	A	820	CLA	O2A-C1-C2	4.44	120.31	108.64
22	1	306	CLA	C1-C2-C3	-4.44	118.37	126.04
24	B	853	BCR	C27-C26-C25	-4.44	116.29	122.73
24	3	304	BCR	C12-C13-C14	-4.43	112.14	118.94
24	A	846	BCR	C7-C8-C9	-4.43	119.54	126.23
22	A	827	CLA	O2A-C1-C2	4.43	120.28	108.64
39	6	318	CHL	CHD-C1D-ND	-4.42	120.39	124.45
22	3	320	CLA	O2D-CGD-CBD	4.42	119.13	111.27
24	B	847	BCR	C38-C26-C25	-4.42	119.56	124.53
22	Z	310	CLA	O2A-C1-C2	4.42	120.25	108.64
22	8	305	CLA	O2A-C1-C2	4.42	120.24	108.64
38	4	802	LUT	C11-C10-C9	-4.41	121.02	127.31
24	7	303	BCR	C16-C17-C18	-4.41	121.02	127.31
25	6	323	LHG	O7-C7-C8	4.41	121.00	111.50
24	B	846	BCR	C16-C17-C18	-4.40	121.03	127.31
22	5	320	CLA	CMD-C2D-C1D	4.40	132.47	124.71
24	B	844	BCR	C38-C26-C25	-4.40	119.59	124.53
22	1	316	CLA	O2D-CGD-CBD	4.40	119.08	111.27

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Mol	Chain	Res	Type	Atoms	Z	Observed(^{o})	Ideal(^{o})
22	1	309	CLA	O2A-C1-C2	4.40	120.19	108.64
24	A	847	BCR	C7-C8-C9	-4.40	119.59	126.23
24	4	804	BCR	C20-C21-C22	-4.39	121.04	127.31
22	A	812	CLA	O2A-C1-C2	4.39	120.18	108.64
22	F	302	CLA	O2A-C1-C2	4.39	120.17	108.64
22	A	824	CLA	O2A-C1-C2	4.39	120.17	108.64
22	A	836	CLA	O2A-C1-C2	4.38	120.16	108.64
38	1	302	LUT	C35-C34-C33	-4.37	121.07	127.31
24	8	304	BCR	C24-C23-C22	-4.37	119.64	126.23
38	7	301	LUT	C15-C14-C13	-4.36	121.08	127.31
22	A	807	CLA	O2A-C1-C2	4.36	120.09	108.64
24	F	303	BCR	C38-C26-C25	-4.35	119.64	124.53
24	B	845	BCR	C11-C10-C9	-4.35	121.10	127.31
24	5	304	BCR	C30-C25-C26	-4.35	116.49	122.61
24	L	204	BCR	C33-C5-C6	-4.35	119.65	124.53
24	L	204	BCR	C16-C17-C18	-4.35	121.11	127.31
39	5	321	CHL	CHD-C1D-ND	-4.34	120.46	124.45
22	B	813	CLA	C1-C2-C3	-4.34	118.54	126.04
24	A	844	BCR	C35-C13-C14	-4.34	116.84	122.92
24	6	305	BCR	C15-C14-C13	-4.34	121.12	127.31
22	4	805	CLA	O2D-CGD-CBD	4.33	118.97	111.27
39	6	320	CHL	CHD-C1D-ND	-4.33	120.47	124.45
24	3	306	BCR	C27-C26-C25	-4.33	116.45	122.73
24	B	853	BCR	C38-C26-C25	-4.32	119.68	124.53
24	A	846	BCR	C33-C5-C6	-4.32	119.68	124.53
39	4	819	CHL	CHD-C1D-ND	-4.32	120.49	124.45
24	B	853	BCR	C38-C26-C27	4.31	121.90	113.62
22	7	310	CLA	O2D-CGD-CBD	4.31	118.93	111.27
24	B	845	BCR	C15-C14-C13	-4.31	121.16	127.31
24	I	4001	BCR	C7-C8-C9	-4.30	119.73	126.23
24	I	4001	BCR	C11-C10-C9	-4.30	121.17	127.31
39	Z	312	CHL	CHD-C1D-ND	-4.30	120.50	124.45
24	8	304	BCR	C1-C6-C5	-4.30	116.56	122.61
22	B	837	CLA	C1-C2-C3	-4.29	118.62	126.04
39	4	816	CHL	CHD-C1D-ND	-4.29	120.51	124.45
39	4	813	CHL	CHD-C1D-ND	-4.29	120.51	124.45
22	9	303	CLA	O2D-CGD-CBD	4.29	118.89	111.27
22	5	307	CLA	C1-C2-C3	-4.28	118.63	126.04
24	L	204	BCR	C15-C14-C13	-4.28	121.20	127.31
39	2	303	CHL	CHD-C1D-ND	-4.27	120.53	124.45
24	B	848	BCR	C16-C17-C18	-4.27	121.22	127.31
24	A	846	BCR	C20-C21-C22	-4.27	121.22	127.31

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Mol	Chain	Res	Type	Atoms	Z	Observed(^{o})	Ideal(^{o})
24	3	307	BCR	C33-C5-C6	-4.26	119.74	124.53
24	3	304	BCR	C27-C26-C25	-4.26	116.54	122.73
22	5	307	CLA	O2A-C1-C2	4.26	119.84	108.64
36	5	306	C7Z	C1-C6-C5	-4.26	116.61	122.61
24	L	204	BCR	C30-C25-C26	-4.26	116.62	122.61
25	1	317	LHG	O7-C7-C8	4.26	120.67	111.50
39	8	301	CHL	CHD-C1D-ND	-4.25	120.55	124.45
22	B	838	CLA	O2A-C1-C2	4.25	119.81	108.64
25	Z	317	LHG	O7-C7-C8	4.25	120.67	111.50
22	4	806	CLA	C1-C2-C3	-4.25	118.69	126.04
22	K	203	CLA	C1-C2-C3	-4.25	118.69	126.04
22	L	203	CLA	C1-C2-C3	-4.25	119.88	126.75
22	7	305	CLA	C1-C2-C3	-4.25	119.88	126.75
24	3	307	BCR	C1-C6-C5	-4.25	116.63	122.61
22	4	812	CLA	O2A-C1-C2	4.24	119.79	108.64
24	F	303	BCR	C24-C23-C22	-4.24	119.82	126.23
24	I	4001	BCR	C38-C26-C27	4.24	121.76	113.62
24	A	856	BCR	C11-C10-C9	-4.24	121.26	127.31
38	7	301	LUT	C7-C8-C9	-4.24	119.83	126.23
22	4	810	CLA	C1-C2-C3	-4.24	119.90	126.75
22	B	817	CLA	O2A-C1-C2	4.23	119.75	108.64
22	B	820	CLA	O2A-C1-C2	4.23	119.75	108.64
36	J	105	C7Z	C18-C5-C6	-4.23	119.78	124.53
24	G	203	BCR	C27-C26-C25	-4.23	116.60	122.73
22	7	310	CLA	C1-C2-C3	-4.22	118.75	126.04
38	4	803	LUT	C18-C5-C6	-4.21	119.80	124.53
22	A	826	CLA	C1-C2-C3	-4.21	119.94	126.75
22	B	823	CLA	O2A-C1-C2	4.21	119.71	108.64
24	A	848	BCR	C4-C5-C6	-4.21	116.62	122.73
24	K	206	BCR	C24-C23-C22	-4.21	119.88	126.23
39	9	312	CHL	CHD-C1D-ND	-4.20	120.59	124.45
22	7	306	CLA	C1-C2-C3	-4.20	118.77	126.04
24	B	845	BCR	C4-C5-C6	-4.20	116.63	122.73
22	9	309	CLA	CMD-C2D-C1D	4.20	132.11	124.71
24	A	848	BCR	C30-C25-C26	-4.20	116.70	122.61
39	1	312	CHL	CHD-C1D-ND	-4.19	120.60	124.45
22	3	315	CLA	CMD-C2D-C1D	4.19	132.10	124.71
24	A	846	BCR	C38-C26-C25	-4.19	119.82	124.53
39	6	316	CHL	CHD-C1D-ND	-4.19	120.60	124.45
22	5	326	CLA	O2A-C1-C2	4.19	119.65	108.64
24	5	304	BCR	C33-C5-C6	-4.19	119.83	124.53
39	Z	311	CHL	CHD-C1D-ND	-4.18	120.61	124.45

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Mol	Chain	Res	Type	Atoms	Z	Observed(^{o})	Ideal(^{o})
22	9	306	CLA	C1-C2-C3	-4.18	118.82	126.04
38	7	301	LUT	C11-C10-C9	-4.17	121.35	127.31
24	A	844	BCR	C30-C25-C26	-4.17	116.73	122.61
38	Z	302	LUT	C18-C5-C6	-4.17	119.84	124.53
22	A	806	CLA	C1-C2-C3	-4.17	118.83	126.04
24	3	304	BCR	C8-C9-C10	4.17	125.34	118.94
36	5	306	C7Z	C31-C30-C29	-4.17	121.36	127.31
22	A	814	CLA	C1-C2-C3	-4.17	118.84	126.04
24	B	849	BCR	C24-C23-C22	-4.16	119.95	126.23
38	1	303	LUT	C18-C5-C6	-4.16	119.86	124.53
38	6	304	LUT	C11-C10-C9	-4.16	121.38	127.31
22	Z	306	CLA	O2A-C1-C2	4.16	119.56	108.64
24	I	4001	BCR	C4-C5-C6	-4.15	116.70	122.73
39	9	314	CHL	CHD-C1D-ND	-4.15	120.64	124.45
24	I	4001	BCR	C27-C26-C25	-4.15	116.71	122.73
22	6	319	CLA	C1-C2-C3	-4.15	118.87	126.04
36	J	105	C7Z	C38-C25-C26	-4.15	119.87	124.53
22	4	815	CLA	C1-C2-C3	-4.15	120.05	126.75
24	8	304	BCR	C20-C21-C22	-4.14	121.40	127.31
36	J	105	C7Z	C1-C6-C5	-4.14	116.78	122.61
22	7	315	CLA	C1-C2-C3	-4.14	120.05	126.75
22	Z	307	CLA	C1-C2-C3	-4.14	118.89	126.04
39	5	316	CHL	CHD-C1D-ND	-4.14	120.65	124.45
38	3	302	LUT	C18-C5-C6	-4.14	119.88	124.53
38	4	803	LUT	C15-C14-C13	-4.14	121.41	127.31
21	A	801	CL0	O2A-CGA-CBA	4.13	124.88	111.91
24	B	845	BCR	C7-C8-C9	-4.13	119.99	126.23
24	3	307	BCR	C27-C26-C25	-4.12	116.75	122.73
22	7	314	CLA	C1-C2-C3	-4.12	120.09	126.75
24	A	847	BCR	C15-C14-C13	-4.12	121.43	127.31
39	8	312	CHL	CHD-C1D-ND	-4.12	120.67	124.45
22	4	805	CLA	O2A-C1-C2	4.11	119.44	108.64
24	6	306	BCR	C33-C5-C6	-4.11	119.91	124.53
33	4	822	LMT	C1B-O1B-C4'	-4.10	107.81	117.96
22	5	320	CLA	C2C-C1C-NC	4.10	113.82	109.97
22	6	317	CLA	C1-C2-C3	-4.10	120.11	126.75
22	7	317	CLA	C1-C2-C3	-4.10	118.95	126.04
38	8	303	LUT	C18-C5-C6	-4.10	119.92	124.53
24	A	847	BCR	C11-C10-C9	-4.10	121.46	127.31
22	A	825	CLA	C1-C2-C3	-4.09	118.97	126.04
38	Z	301	LUT	C1-C6-C5	-4.09	116.86	122.61
25	A	850	LHG	O7-C7-C8	4.08	120.30	111.50

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Mol	Chain	Res	Type	Atoms	Z	Observed(^{o})	Ideal(^{o})
38	4	803	LUT	C1-C6-C5	-4.08	116.87	122.61
24	L	204	BCR	C24-C23-C22	-4.08	120.07	126.23
41	5	325	3PH	O21-C21-C22	4.07	120.27	111.50
25	7	318	LHG	O7-C7-C8	4.06	120.26	111.50
21	A	801	CL0	C1D-ND-C4D	-4.06	103.45	106.33
24	3	306	BCR	C4-C5-C6	-4.06	116.83	122.73
22	1	308	CLA	C1-C2-C3	-4.06	119.02	126.04
24	6	306	BCR	C15-C14-C13	-4.06	121.52	127.31
22	A	828	CLA	C1-C2-C3	-4.06	119.02	126.04
36	5	306	C7Z	C27-C28-C29	-4.06	120.10	126.23
24	3	306	BCR	C38-C26-C25	-4.06	119.97	124.53
24	B	845	BCR	C30-C25-C26	-4.05	116.90	122.61
22	1	316	CLA	O2A-C1-C2	4.05	119.28	108.64
24	A	856	BCR	C30-C25-C26	-4.05	116.91	122.61
24	3	306	BCR	C7-C8-C9	-4.05	120.12	126.23
39	4	814	CHL	CHD-C1D-ND	-4.03	120.75	124.45
22	B	810	CLA	C1-C2-C3	-4.02	119.08	126.04
22	9	308	CLA	C1-C2-C3	-4.02	120.25	126.75
24	G	203	BCR	C1-C6-C5	-4.02	116.95	122.61
22	9	313	CLA	C1-C2-C3	-4.01	120.26	126.75
22	9	304	CLA	C2D-C1D-ND	4.01	113.06	110.10
38	9	301	LUT	C11-C10-C9	-4.00	121.60	127.31
22	1	307	CLA	C1-C2-C3	-4.00	119.12	126.04
38	3	303	LUT	C18-C5-C6	-4.00	120.04	124.53
25	B	850	LHG	O7-C7-C8	4.00	120.12	111.50
22	7	307	CLA	C1-C2-C3	-3.99	119.13	126.04
32	F	306	RRX	C16-C15-C14	-3.99	115.30	123.47
22	4	807	CLA	C1-C2-C3	-3.99	119.14	126.04
22	8	311	CLA	C1-C2-C3	-3.99	120.30	126.75
24	6	306	BCR	C4-C5-C6	-3.99	116.94	122.73
39	8	301	CHL	C1-C2-C3	-3.99	119.15	126.04
24	8	304	BCR	C29-C30-C25	3.98	116.61	110.48
22	B	824	CLA	C1-C2-C3	-3.98	119.16	126.04
22	B	825	CLA	C1-C2-C3	-3.98	119.16	126.04
24	3	304	BCR	C38-C26-C25	-3.97	120.07	124.53
24	B	849	BCR	C27-C26-C25	-3.97	116.97	122.73
34	J	101	LMG	O7-C10-C11	3.97	120.05	111.50
22	A	831	CLA	C1-C2-C3	-3.97	120.33	126.75
25	A	849	LHG	O7-C7-C8	3.96	120.04	111.50
24	6	306	BCR	C11-C12-C13	-3.96	115.29	126.42
24	7	303	BCR	C24-C23-C22	-3.96	120.26	126.23
39	7	313	CHL	CHD-C1D-ND	-3.96	120.82	124.45

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Mol	Chain	Res	Type	Atoms	Z	Observed(^{o})	Ideal(^{o})
24	B	847	BCR	C30-C25-C26	-3.95	117.05	122.61
24	B	853	BCR	C33-C5-C6	-3.95	120.10	124.53
24	J	104	BCR	C30-C25-C26	-3.94	117.06	122.61
22	1	316	CLA	C2C-C1C-NC	3.94	113.66	109.97
38	8	303	LUT	C1-C6-C5	-3.94	117.07	122.61
24	B	847	BCR	C24-C23-C22	-3.94	120.29	126.23
22	G	201	CLA	C1-C2-C3	-3.94	120.39	126.75
22	B	841	CLA	C1-C2-C3	-3.93	119.24	126.04
22	3	312	CLA	C1-C2-C3	-3.93	119.25	126.04
22	9	310	CLA	C2D-C1D-ND	3.93	113.00	110.10
32	F	306	RRX	C38-C26-C25	-3.93	120.12	124.53
24	I	4001	BCR	C33-C5-C6	-3.92	120.12	124.53
24	A	844	BCR	C24-C23-C22	-3.92	120.31	126.23
38	9	302	LUT	C15-C14-C13	-3.92	121.71	127.31
22	5	313	CLA	C1-C2-C3	-3.92	119.26	126.04
41	6	324	3PH	O21-C21-C22	3.92	119.95	111.50
38	5	302	LUT	C7-C8-C9	-3.92	120.32	126.23
22	2	301	CLA	C1-C2-C3	-3.91	119.27	126.04
36	5	306	C7Z	C18-C5-C6	-3.91	120.13	124.53
24	B	844	BCR	C12-C13-C14	3.91	124.94	118.94
24	A	845	BCR	C4-C5-C6	-3.91	117.06	122.73
22	9	309	CLA	O2D-CGD-CBD	3.90	118.21	111.27
24	3	304	BCR	C34-C9-C10	-3.90	117.45	122.92
24	3	304	BCR	C16-C15-C14	3.90	131.46	123.47
39	5	317	CHL	CHD-C1D-ND	-3.89	120.88	124.45
22	Z	316	CLA	C1-C2-C3	-3.89	119.31	126.04
24	A	845	BCR	C38-C26-C25	-3.89	120.16	124.53
24	6	305	BCR	C33-C5-C6	-3.89	120.16	124.53
22	5	312	CLA	C1-C2-C3	-3.89	120.46	126.75
24	A	844	BCR	C10-C11-C12	-3.88	111.09	123.22
24	6	306	BCR	C27-C26-C25	-3.88	117.10	122.73
24	3	307	BCR	C11-C12-C13	-3.88	115.53	126.42
24	A	847	BCR	C4-C5-C6	-3.87	117.11	122.73
38	5	303	LUT	C18-C5-C6	-3.87	120.18	124.53
24	6	305	BCR	C4-C5-C6	-3.87	117.11	122.73
24	A	844	BCR	C1-C6-C5	-3.87	117.17	122.61
22	9	303	CLA	C2D-C1D-ND	3.86	112.95	110.10
38	6	304	LUT	C18-C5-C6	-3.86	120.20	124.53
24	5	304	BCR	C4-C5-C6	-3.86	117.13	122.73
24	K	206	BCR	C33-C5-C6	-3.86	120.20	124.53
38	1	303	LUT	C31-C30-C29	-3.86	121.81	127.31
24	F	303	BCR	C11-C12-C13	-3.86	115.59	126.42

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Mol	Chain	Res	Type	Atoms	Z	Observed(^{o})	Ideal(^{o})
24	7	303	BCR	C33-C5-C6	-3.85	120.20	124.53
24	J	104	BCR	C24-C23-C22	-3.85	120.41	126.23
24	A	848	BCR	C33-C5-C4	3.85	121.02	113.62
25	4	820	LHG	O7-C7-C8	3.85	119.79	111.50
24	3	307	BCR	C30-C25-C26	-3.84	117.20	122.61
22	B	839	CLA	C1-C2-C3	-3.84	119.40	126.04
24	B	846	BCR	C20-C21-C22	-3.84	121.83	127.31
22	8	305	CLA	C1-C2-C3	-3.84	119.40	126.04
22	A	832	CLA	O2A-C1-C2	3.84	118.73	108.64
24	B	853	BCR	C24-C23-C22	-3.84	120.44	126.23
24	B	846	BCR	C15-C16-C17	-3.84	115.62	123.47
22	B	832	CLA	C1-C2-C3	-3.82	120.58	126.75
22	Z	308	CLA	C1-C2-C3	-3.81	119.45	126.04
22	6	309	CLA	C1-C2-C3	-3.81	119.46	126.04
38	8	303	LUT	C21-C26-C27	-3.81	107.89	112.70
22	5	309	CLA	C1-C2-C3	-3.80	119.46	126.04
22	F	301	CLA	C1-C2-C3	-3.80	119.46	126.04
38	6	304	LUT	C31-C30-C29	-3.80	121.88	127.31
24	5	304	BCR	C27-C26-C25	-3.80	117.21	122.73
24	A	847	BCR	C33-C5-C6	-3.80	120.26	124.53
24	A	845	BCR	C30-C25-C26	-3.79	117.28	122.61
38	8	303	LUT	C11-C10-C9	-3.79	121.90	127.31
22	Z	310	CLA	C1-C2-C3	-3.79	120.62	126.75
35	J	102	T7X	O16-C10-C12	3.78	119.66	111.50
24	B	846	BCR	C38-C26-C27	3.78	120.89	113.62
22	B	821	CLA	C1-C2-C3	-3.78	119.50	126.04
24	A	846	BCR	C16-C17-C18	-3.78	121.92	127.31
25	5	324	LHG	O7-C7-C8	3.78	119.64	111.50
38	1	302	LUT	C15-C14-C13	-3.76	121.94	127.31
27	A	852	DGA	CDB-CCB-CBB	-3.76	80.39	115.30
24	3	304	BCR	C2-C1-C6	3.76	116.27	110.48
24	L	204	BCR	C7-C8-C9	-3.75	120.56	126.23
38	Z	301	LUT	C7-C8-C9	-3.75	120.57	126.23
24	A	845	BCR	C7-C8-C9	-3.75	120.57	126.23
24	A	844	BCR	C34-C9-C10	-3.75	117.68	122.92
38	6	303	LUT	C18-C5-C6	-3.74	120.33	124.53
22	A	841	CLA	C1-C2-C3	-3.74	119.58	126.04
38	6	304	LUT	C21-C26-C27	-3.74	107.98	112.70
22	5	319	CLA	C1-C2-C3	-3.73	119.58	126.04
39	6	315	CHL	CHD-C1D-ND	-3.73	121.02	124.45
24	B	849	BCR	C15-C14-C13	-3.73	121.98	127.31
22	3	315	CLA	C2D-C1D-ND	3.73	112.85	110.10

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Mol	Chain	Res	Type	Atoms	Z	Observed(^{o})	Ideal(^{o})
38	3	303	LUT	C35-C34-C33	-3.73	121.99	127.31
32	F	306	RRX	C35-C13-C14	-3.73	117.70	122.92
24	K	206	BCR	C4-C5-C6	-3.73	117.32	122.73
22	3	308	CLA	C1-C2-C3	-3.73	119.60	126.04
39	5	321	CHL	CMA-C3A-C4A	3.73	121.79	111.77
22	A	816	CLA	C1-C2-C3	-3.73	119.60	126.04
27	A	852	DGA	OG2-CB1-CB2	3.72	119.52	111.50
24	3	304	BCR	C15-C14-C13	3.72	132.62	127.31
22	8	313	CLA	C1-C2-C3	-3.72	120.74	126.75
22	A	830	CLA	C1-C2-C3	-3.71	119.63	126.04
22	9	309	CLA	C2D-C1D-ND	3.71	112.84	110.10
38	Z	302	LUT	C31-C30-C29	-3.70	122.03	127.31
38	3	302	LUT	C31-C30-C29	-3.70	122.03	127.31
24	J	104	BCR	C11-C10-C9	-3.70	122.03	127.31
24	B	848	BCR	C15-C16-C17	-3.69	115.91	123.47
25	4	821	LHG	O7-C7-C8	3.69	119.46	111.50
24	A	848	BCR	C38-C26-C27	3.69	120.70	113.62
22	K	205	CLA	C2C-C1C-NC	3.68	113.42	109.97
22	A	808	CLA	C1-C2-C3	-3.68	119.67	126.04
22	7	322	CLA	C1-C2-C3	-3.68	119.67	126.04
24	A	846	BCR	C15-C16-C17	-3.68	115.93	123.47
22	5	310	CLA	C1-C2-C3	-3.68	119.68	126.04
22	B	809	CLA	C1-C2-C3	-3.68	119.68	126.04
38	4	802	LUT	C31-C30-C29	-3.68	122.06	127.31
24	K	206	BCR	C38-C26-C25	-3.68	120.40	124.53
22	6	302	CLA	C1-C2-C3	-3.67	119.69	126.04
39	Z	311	CHL	C1-C2-C3	-3.67	120.81	126.75
24	4	804	BCR	C11-C10-C9	-3.67	122.07	127.31
22	5	308	CLA	C1-C2-C3	-3.67	119.70	126.04
22	A	842	CLA	C1-C2-C3	-3.67	119.70	126.04
39	9	312	CHL	C1-O2A-CGA	3.67	126.06	116.44
21	A	801	CL0	C1-C2-C3	-3.67	119.70	126.04
22	B	828	CLA	C1-C2-C3	-3.66	119.70	126.04
24	J	104	BCR	C15-C14-C13	-3.66	122.09	127.31
22	7	312	CLA	C1-C2-C3	-3.65	119.72	126.04
22	B	815	CLA	C1-C2-C3	-3.65	119.73	126.04
39	6	318	CHL	C1-C2-C3	-3.64	119.75	126.04
22	A	809	CLA	C1-C2-C3	-3.64	119.75	126.04
24	L	204	BCR	C20-C21-C22	-3.64	122.12	127.31
24	3	304	BCR	C35-C13-C12	3.64	123.81	118.08
22	7	308	CLA	C1-C2-C3	-3.63	119.76	126.04
22	B	830	CLA	C1-C2-C3	-3.63	119.76	126.04

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Mol	Chain	Res	Type	Atoms	Z	Observed(^{o})	Ideal(^{o})
24	B	845	BCR	C33-C5-C4	3.63	120.59	113.62
24	B	847	BCR	C20-C21-C22	-3.63	122.13	127.31
22	6	309	CLA	CAA-C2A-C3A	-3.63	102.84	112.78
25	8	317	LHG	O7-C7-C8	3.63	119.32	111.50
24	A	848	BCR	C27-C26-C25	-3.63	117.47	122.73
22	5	320	CLA	O2D-CGD-CBD	3.62	117.71	111.27
22	8	307	CLA	C1-C2-C3	-3.62	119.78	126.04
41	7	319	3PH	O21-C21-C22	3.62	119.30	111.50
24	8	304	BCR	C38-C26-C27	3.62	120.57	113.62
22	A	840	CLA	C1-C2-C3	-3.62	119.78	126.04
22	7	304	CLA	C1-C2-C3	-3.61	119.80	126.04
38	7	302	LUT	C18-C5-C6	-3.61	120.47	124.53
38	9	302	LUT	C31-C30-C29	-3.61	122.16	127.31
24	B	849	BCR	C15-C16-C17	-3.61	116.09	123.47
38	9	301	LUT	C18-C5-C6	-3.60	120.48	124.53
24	J	104	BCR	C16-C17-C18	-3.60	122.17	127.31
22	9	310	CLA	O2D-CGD-O1D	-3.60	116.80	123.84
22	3	310	CLA	CAA-C2A-C3A	-3.60	102.92	112.78
22	3	320	CLA	C2D-C1D-ND	3.60	112.75	110.10
24	5	304	BCR	C1-C6-C5	-3.60	117.55	122.61
22	1	310	CLA	C1-C2-C3	-3.60	119.83	126.04
38	5	302	LUT	C11-C10-C9	-3.59	122.18	127.31
24	J	104	BCR	C7-C8-C9	-3.59	120.81	126.23
22	6	308	CLA	C1-C2-C3	-3.59	119.84	126.04
22	8	308	CLA	C1-C2-C3	-3.59	119.84	126.04
38	3	302	LUT	C15-C14-C13	-3.58	122.20	127.31
24	K	206	BCR	C38-C26-C27	3.58	120.49	113.62
40	1	318	SQD	O7-S-C6	-3.58	102.69	106.94
22	A	803	CLA	C1-C2-C3	-3.58	119.86	126.04
24	B	846	BCR	C29-C30-C25	3.57	115.98	110.48
22	Z	309	CLA	C1-C2-C3	-3.57	119.86	126.04
22	B	818	CLA	C1-C2-C3	-3.57	119.87	126.04
32	F	306	RRX	C4-C5-C6	-3.57	117.55	122.73
24	5	305	BCR	C24-C23-C22	-3.57	120.84	126.23
24	5	304	BCR	C8-C7-C6	-3.56	117.19	127.20
22	K	205	CLA	C2D-C1D-ND	3.56	112.73	110.10
38	4	803	LUT	C21-C26-C27	-3.56	108.20	112.70
24	6	306	BCR	C20-C21-C22	-3.56	122.23	127.31
22	4	818	CLA	C1-C2-C3	-3.56	119.89	126.04
24	B	847	BCR	C15-C16-C17	-3.56	116.19	123.47
24	B	845	BCR	C24-C23-C22	-3.56	120.86	126.23
22	B	827	CLA	C1-C2-C3	-3.55	119.89	126.04

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Mol	Chain	Res	Type	Atoms	Z	Observed(^{o})	Ideal(^{o})
24	3	305	BCR	C11-C12-C13	-3.55	116.43	126.42
22	6	311	CLA	C1-C2-C3	-3.55	119.90	126.04
22	3	315	CLA	O2D-CGD-O1D	-3.55	116.89	123.84
31	B	852	DGD	O2G-C1B-C2B	3.55	119.16	111.50
22	A	838	CLA	C1-C2-C3	-3.55	119.91	126.04
24	F	303	BCR	C1-C6-C5	-3.55	117.62	122.61
39	Z	311	CHL	C3C-C4C-NC	-3.54	106.60	110.57
24	F	303	BCR	C15-C14-C13	-3.54	122.26	127.31
22	A	807	CLA	C1-C2-C3	-3.54	119.92	126.04
22	4	812	CLA	C1-C2-C3	-3.54	119.92	126.04
24	A	845	BCR	C27-C26-C25	-3.54	117.59	122.73
39	4	814	CHL	C1-C2-C3	-3.53	121.03	126.75
22	1	309	CLA	C1-C2-C3	-3.53	119.93	126.04
24	A	846	BCR	C4-C5-C6	-3.53	117.61	122.73
24	5	304	BCR	C11-C12-C13	-3.53	116.51	126.42
38	8	302	LUT	C21-C26-C27	-3.52	108.25	112.70
24	B	846	BCR	C4-C5-C6	-3.52	117.62	122.73
24	A	845	BCR	C24-C23-C22	-3.52	120.92	126.23
22	1	308	CLA	CHD-C1D-ND	-3.51	121.22	124.45
22	9	305	CLA	C2D-C1D-ND	3.51	112.69	110.10
24	B	844	BCR	C4-C5-C6	-3.51	117.63	122.73
24	B	849	BCR	C1-C6-C5	-3.51	117.67	122.61
22	3	313	CLA	C2D-C1D-ND	3.51	112.69	110.10
24	K	206	BCR	C1-C6-C5	-3.51	117.67	122.61
22	5	322	CLA	C1-C2-C3	-3.51	119.97	126.04
22	A	822	CLA	C1-C2-C3	-3.51	119.98	126.04
24	B	848	BCR	C15-C14-C13	-3.49	122.32	127.31
38	9	302	LUT	C18-C5-C6	-3.49	120.61	124.53
32	F	306	RRX	C16-C17-C18	-3.49	122.33	127.31
24	4	804	BCR	C2-C1-C6	3.49	115.85	110.48
22	B	834	CLA	C1-C2-C3	-3.48	120.02	126.04
22	9	305	CLA	C2C-C1C-NC	3.48	113.23	109.97
22	A	818	CLA	C1-C2-C3	-3.48	120.03	126.04
24	B	846	BCR	C38-C26-C25	-3.47	120.63	124.53
22	8	309	CLA	C1-C2-C3	-3.47	120.04	126.04
24	3	306	BCR	C33-C5-C6	-3.47	120.63	124.53
24	A	845	BCR	C38-C26-C27	3.46	120.27	113.62
22	1	311	CLA	C2C-C1C-NC	3.46	113.22	109.97
22	5	309	CLA	CAA-C2A-C3A	-3.46	103.30	112.78
38	1	302	LUT	C18-C5-C4	3.46	120.77	114.36
22	B	835	CLA	CHD-C1D-ND	-3.45	121.28	124.45
24	7	303	BCR	C33-C5-C4	3.45	120.24	113.62

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Mol	Chain	Res	Type	Atoms	Z	Observed(^{o})	Ideal(^{o})
24	B	844	BCR	C34-C9-C10	-3.45	118.09	122.92
38	8	303	LUT	C7-C8-C9	-3.44	121.03	126.23
38	Z	302	LUT	C21-C26-C27	-3.44	108.35	112.70
22	4	805	CLA	C2C-C1C-NC	3.44	113.19	109.97
38	3	303	LUT	C31-C30-C29	-3.44	122.41	127.31
22	3	310	CLA	C1-C2-C3	-3.44	120.10	126.04
24	3	304	BCR	C24-C23-C22	-3.44	121.04	126.23
39	6	320	CHL	C4D-CHA-C1A	3.43	125.43	121.25
39	3	317	CHL	C2C-C3C-C4C	3.43	108.94	106.49
22	B	822	CLA	C1-C2-C3	-3.43	120.11	126.04
22	A	813	CLA	C2C-C1C-NC	3.43	113.19	109.97
22	B	805	CLA	C1-C2-C3	-3.43	120.11	126.04
22	B	806	CLA	C1-C2-C3	-3.43	120.11	126.04
39	8	315	CHL	C4D-CHA-C1A	3.43	125.42	121.25
38	9	302	LUT	C11-C10-C9	-3.43	122.42	127.31
22	B	816	CLA	C1-C2-C3	-3.42	120.12	126.04
24	G	203	BCR	C29-C30-C25	3.42	115.75	110.48
22	1	308	CLA	C1D-ND-C4D	-3.42	103.90	106.33
22	6	308	CLA	C6-C5-C3	-3.42	109.02	114.62
24	6	305	BCR	C29-C30-C25	3.42	115.75	110.48
24	G	203	BCR	C11-C12-C13	-3.42	116.82	126.42
24	L	204	BCR	C33-C5-C4	3.42	120.18	113.62
38	4	803	LUT	C31-C30-C29	-3.42	122.44	127.31
24	K	206	BCR	C27-C26-C25	-3.41	117.78	122.73
24	B	844	BCR	C1-C6-C5	-3.41	117.81	122.61
22	B	829	CLA	C1-C2-C3	-3.41	120.15	126.04
22	A	823	CLA	C1-C2-C3	-3.40	120.16	126.04
38	5	303	LUT	C31-C30-C29	-3.40	122.45	127.31
22	9	303	CLA	CAA-C2A-C3A	-3.40	103.47	112.78
38	6	303	LUT	C1-C6-C5	-3.40	117.83	122.61
22	A	815	CLA	C1-C2-C3	-3.40	120.17	126.04
24	4	804	BCR	C33-C5-C4	3.39	120.14	113.62
22	7	304	CLA	CHD-C1D-ND	-3.39	121.33	124.45
38	4	802	LUT	C18-C5-C6	-3.39	120.72	124.53
24	4	804	BCR	C7-C8-C9	-3.39	121.11	126.23
22	7	306	CLA	C2C-C1C-NC	3.39	113.15	109.97
24	B	848	BCR	C11-C12-C13	-3.39	116.90	126.42
24	B	844	BCR	C27-C26-C25	-3.38	117.82	122.73
38	Z	302	LUT	C1-C6-C5	-3.38	117.85	122.61
22	5	315	CLA	C1-C2-C3	-3.38	120.20	126.04
22	3	311	CLA	C1-C2-C3	-3.38	120.20	126.04
24	B	849	BCR	C30-C25-C26	-3.38	117.86	122.61

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Mol	Chain	Res	Type	Atoms	Z	Observed(^{o})	Ideal(^{o})
22	B	842	CLA	C1-C2-C3	-3.38	120.20	126.04
22	9	303	CLA	C1D-ND-C4D	-3.37	103.94	106.33
24	3	305	BCR	C20-C21-C22	-3.37	122.50	127.31
24	A	848	BCR	C38-C26-C25	-3.37	120.74	124.53
24	A	846	BCR	C29-C30-C25	3.37	115.67	110.48
22	B	831	CLA	C1-C2-C3	-3.37	120.21	126.04
22	6	313	CLA	C1-C2-C3	-3.37	120.22	126.04
38	5	303	LUT	C1-C6-C5	-3.36	117.87	122.61
24	B	844	BCR	C36-C18-C17	-3.36	118.21	122.92
22	B	815	CLA	C2C-C1C-NC	3.36	113.12	109.97
39	4	813	CHL	C1-C2-C3	-3.36	121.32	126.75
24	6	305	BCR	C15-C16-C17	-3.36	116.60	123.47
22	Z	303	CLA	C1-C2-C3	-3.35	120.24	126.04
38	Z	301	LUT	C18-C5-C4	3.35	120.57	114.36
39	3	317	CHL	C3C-C4C-NC	-3.35	106.81	110.57
22	G	201	CLA	C2C-C1C-NC	3.35	113.11	109.97
22	9	305	CLA	C1-C2-C3	-3.35	120.25	126.04
22	9	309	CLA	C2C-C1C-NC	3.35	113.11	109.97
22	A	805	CLA	CHD-C1D-ND	-3.34	121.38	124.45
24	B	853	BCR	C4-C5-C6	-3.34	117.88	122.73
22	5	320	CLA	C2D-C1D-ND	3.34	112.57	110.10
22	4	806	CLA	C6-C5-C3	-3.34	109.16	114.62
24	G	203	BCR	C24-C23-C22	-3.34	121.19	126.23
24	A	848	BCR	C16-C17-C18	-3.34	122.55	127.31
22	F	302	CLA	C1-C2-C3	-3.34	120.27	126.04
22	3	318	CLA	C1-C2-C3	-3.33	120.28	126.04
39	4	819	CHL	C4D-CHA-C1A	3.33	125.31	121.25
22	7	311	CLA	C2D-C1D-ND	3.33	112.56	110.10
41	8	320	3PH	O21-C21-C22	3.33	120.08	110.80
22	6	308	CLA	C2D-C1D-ND	3.33	112.56	110.10
22	3	313	CLA	CHD-C1D-ND	-3.32	121.40	124.45
22	B	824	CLA	C2D-C1D-ND	3.32	112.55	110.10
22	6	301	CLA	C1-C2-C3	-3.32	120.30	126.04
24	3	307	BCR	C35-C13-C14	-3.32	118.27	122.92
32	F	306	RRX	C24-C23-C22	-3.32	121.22	126.23
22	Z	315	CLA	C2C-C1C-NC	3.32	113.08	109.97
22	4	807	CLA	C2C-C1C-NC	3.32	113.08	109.97
22	7	307	CLA	CHD-C1D-ND	-3.32	121.41	124.45
22	L	201	CLA	C2D-C1D-ND	3.32	112.55	110.10
22	4	808	CLA	C1-C2-C3	-3.31	120.31	126.04
22	5	313	CLA	C2C-C1C-NC	3.31	113.08	109.97
24	3	305	BCR	C15-C14-C13	-3.31	122.59	127.31

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Mol	Chain	Res	Type	Atoms	Z	Observed(^{o})	Ideal(^{o})
39	8	301	CHL	C3C-C4C-NC	-3.30	106.87	110.57
38	5	302	LUT	C35-C15-C14	-3.30	116.71	123.47
22	7	309	CLA	C2D-C1D-ND	3.30	112.53	110.10
22	G	202	CLA	CHD-C1D-ND	-3.30	121.42	124.45
38	8	302	LUT	C15-C14-C13	-3.30	122.60	127.31
22	L	202	CLA	C1-C2-C3	-3.29	120.35	126.04
22	6	312	CLA	C1-C2-C3	-3.29	120.35	126.04
22	Z	309	CLA	C2D-C1D-ND	3.29	112.53	110.10
24	B	845	BCR	C20-C21-C22	-3.29	122.62	127.31
22	1	304	CLA	C1-C2-C3	-3.29	120.36	126.04
22	9	307	CLA	C1-C2-C3	-3.29	120.36	126.04
24	6	305	BCR	C11-C12-C13	-3.29	117.19	126.42
24	A	856	BCR	C4-C5-C6	-3.28	117.96	122.73
22	A	816	CLA	C2C-C1C-NC	3.28	113.05	109.97
22	Z	307	CLA	CHD-C1D-ND	-3.28	121.44	124.45
22	L	201	CLA	C1-C2-C3	-3.28	120.37	126.04
38	8	303	LUT	C15-C14-C13	-3.28	122.63	127.31
22	1	308	CLA	C2D-C1D-ND	3.28	112.52	110.10
22	A	833	CLA	C1-C2-C3	-3.28	120.38	126.04
22	4	805	CLA	CMA-C3A-C4A	3.27	120.57	111.77
22	3	318	CLA	CHD-C1D-ND	-3.27	121.45	124.45
38	8	302	LUT	C11-C10-C9	-3.27	122.64	127.31
22	5	307	CLA	CHD-C1D-ND	-3.27	121.45	124.45
22	7	305	CLA	C2D-C1D-ND	3.27	112.51	110.10
38	1	303	LUT	C1-C6-C5	-3.27	118.01	122.61
22	A	805	CLA	C2D-C1D-ND	3.26	112.51	110.10
22	B	805	CLA	C2D-C1D-ND	3.26	112.51	110.10
24	A	847	BCR	C1-C6-C5	-3.26	118.02	122.61
22	K	204	CLA	CHD-C1D-ND	-3.26	121.45	124.45
22	7	304	CLA	C2D-C1D-ND	3.26	112.51	110.10
22	1	313	CLA	C2D-C1D-ND	3.26	112.51	110.10
36	J	105	C7Z	C31-C32-C33	-3.26	117.26	126.42
22	7	322	CLA	C2D-C1D-ND	3.26	112.51	110.10
22	A	839	CLA	C1-C2-C3	-3.26	120.41	126.04
22	A	842	CLA	C2C-C1C-NC	3.25	113.02	109.97
22	A	842	CLA	C6-C5-C3	-3.25	109.30	114.62
38	4	802	LUT	C21-C26-C27	-3.25	108.59	112.70
22	Z	308	CLA	CHD-C1D-ND	-3.25	121.47	124.45
22	B	834	CLA	C2C-C1C-NC	3.25	113.01	109.97
24	B	853	BCR	C30-C25-C26	-3.25	118.04	122.61
39	Z	311	CHL	C2C-C3C-C4C	3.24	108.80	106.49
22	B	838	CLA	CHD-C1D-ND	-3.24	121.47	124.45

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Mol	Chain	Res	Type	Atoms	Z	Observed(^{o})	Ideal(^{o})
22	1	316	CLA	C2D-C1D-ND	3.24	112.49	110.10
39	2	303	CHL	C4D-CHA-C1A	3.24	125.19	121.25
24	6	306	BCR	C33-C5-C4	3.24	119.84	113.62
39	2	303	CHL	C1-C2-C3	-3.24	121.51	126.75
22	B	839	CLA	C2C-C1C-NC	3.24	113.00	109.97
22	8	306	CLA	C1-C2-C3	-3.24	120.44	126.04
24	F	303	BCR	C38-C26-C27	3.24	119.83	113.62
24	B	846	BCR	C1-C6-C5	-3.23	118.06	122.61
22	7	310	CLA	C2D-C1D-ND	3.23	112.49	110.10
24	L	204	BCR	C11-C10-C9	-3.23	122.70	127.31
22	A	814	CLA	C2C-C1C-NC	3.23	113.00	109.97
22	A	817	CLA	CMA-C3A-C4A	3.23	120.46	111.77
22	K	203	CLA	C2C-C1C-NC	3.23	113.00	109.97
39	5	317	CHL	C1-C2-C3	-3.23	121.53	126.75
22	9	308	CLA	C2C-C1C-NC	3.23	113.00	109.97
22	1	314	CLA	CHD-C1D-ND	-3.23	121.49	124.45
22	A	812	CLA	C2D-C1D-ND	3.23	112.48	110.10
22	6	310	CLA	C1-C2-C3	-3.23	120.47	126.04
24	3	306	BCR	C24-C23-C22	-3.22	121.36	126.23
22	A	818	CLA	C2D-C1D-ND	3.22	112.48	110.10
22	5	322	CLA	C2D-C1D-ND	3.22	112.48	110.10
22	F	302	CLA	C2D-C1D-ND	3.22	112.48	110.10
22	B	833	CLA	C1-C2-C3	-3.22	120.47	126.04
22	5	323	CLA	CHD-C1D-ND	-3.22	121.50	124.45
22	6	310	CLA	CHD-C1D-ND	-3.22	121.50	124.45
22	B	810	CLA	C2C-C1C-NC	3.22	112.99	109.97
22	5	326	CLA	C2D-C1D-ND	3.22	112.47	110.10
24	4	804	BCR	C30-C25-C26	-3.21	118.08	122.61
22	Z	305	CLA	C2C-C1C-NC	3.21	112.98	109.97
22	B	824	CLA	CHD-C1D-ND	-3.21	121.50	124.45
22	4	805	CLA	CHD-C1D-ND	-3.21	121.50	124.45
22	6	314	CLA	C2C-C1C-NC	3.21	112.98	109.97
24	A	848	BCR	C24-C23-C22	-3.21	121.39	126.23
38	9	302	LUT	C7-C8-C9	-3.21	121.39	126.23
22	A	827	CLA	C2D-C1D-ND	3.21	112.47	110.10
24	I	4001	BCR	C33-C5-C4	3.21	119.78	113.62
22	F	301	CLA	C2D-C1D-ND	3.21	112.47	110.10
24	5	305	BCR	C4-C5-C6	-3.21	118.08	122.73
22	A	833	CLA	C2D-C1D-ND	3.21	112.47	110.10
22	Z	313	CLA	C2D-C1D-ND	3.21	112.47	110.10
22	3	308	CLA	CHD-C1D-ND	-3.20	121.51	124.45
22	B	823	CLA	C1-C2-C3	-3.20	120.50	126.04

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Mol	Chain	Res	Type	Atoms	Z	Observed(^{o})	Ideal(^{o})
22	3	313	CLA	C1D-ND-C4D	-3.20	104.06	106.33
22	5	301	CLA	CHD-C1D-ND	-3.20	121.51	124.45
22	6	312	CLA	C2C-C1C-NC	3.20	112.97	109.97
22	2	304	CLA	C2D-C1D-ND	3.20	112.46	110.10
22	B	806	CLA	C2C-C1C-NC	3.20	112.97	109.97
38	8	302	LUT	C7-C8-C9	-3.20	121.40	126.23
24	5	305	BCR	C33-C5-C4	3.20	119.76	113.62
24	6	306	BCR	C1-C6-C5	-3.20	118.11	122.61
22	2	301	CLA	CHD-C1D-ND	-3.20	121.52	124.45
22	Z	303	CLA	C2D-C1D-ND	3.20	112.46	110.10
38	9	301	LUT	C15-C14-C13	-3.20	122.75	127.31
22	B	809	CLA	CHD-C1D-ND	-3.19	121.52	124.45
22	A	855	CLA	C1-C2-C3	-3.19	120.52	126.04
22	9	304	CLA	C2C-C1C-NC	3.19	112.96	109.97
22	7	305	CLA	CHD-C1D-ND	-3.19	121.52	124.45
24	G	203	BCR	C7-C8-C9	-3.19	121.41	126.23
22	A	839	CLA	C2C-C1C-NC	3.19	112.96	109.97
22	5	309	CLA	C2C-C1C-NC	3.19	112.96	109.97
38	8	302	LUT	C31-C30-C29	-3.19	122.76	127.31
22	A	810	CLA	C1-C2-C3	-3.19	120.53	126.04
22	5	315	CLA	C2D-C1D-ND	3.19	112.45	110.10
22	A	827	CLA	C1-C2-C3	-3.19	120.53	126.04
22	A	823	CLA	C1D-ND-C4D	-3.18	104.07	106.33
22	B	818	CLA	C2D-C1D-ND	3.18	112.45	110.10
24	3	307	BCR	C33-C5-C4	3.18	119.73	113.62
22	5	320	CLA	C1-C2-C3	-3.18	121.60	126.75
24	B	847	BCR	C38-C26-C27	3.18	119.73	113.62
22	Z	308	CLA	C2D-C1D-ND	3.18	112.45	110.10
22	A	822	CLA	C2C-C1C-NC	3.18	112.95	109.97
22	B	806	CLA	CHD-C1D-ND	-3.18	121.53	124.45
24	5	304	BCR	C38-C26-C27	3.18	119.73	113.62
24	B	844	BCR	C30-C25-C26	-3.18	118.13	122.61
22	A	812	CLA	C1-C2-C3	-3.18	120.54	126.04
38	8	303	LUT	C31-C30-C29	-3.18	122.77	127.31
22	1	307	CLA	CHD-C1D-ND	-3.18	121.53	124.45
22	B	815	CLA	CMA-C3A-C4A	3.18	120.31	111.77
22	Z	305	CLA	C2D-C1D-ND	3.18	112.45	110.10
22	6	309	CLA	C2C-C1C-NC	3.18	112.95	109.97
22	9	305	CLA	CMA-C3A-C4A	3.18	120.31	111.77
22	A	824	CLA	C2D-C1D-ND	3.18	112.44	110.10
22	4	812	CLA	C2C-C1C-NC	3.18	112.95	109.97
22	Z	314	CLA	C2D-C1D-ND	3.18	112.44	110.10

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Mol	Chain	Res	Type	Atoms	Z	Observed(^{o})	Ideal(^{o})
24	5	304	BCR	C15-C16-C17	-3.18	116.97	123.47
22	A	817	CLA	C2C-C1C-NC	3.17	112.95	109.97
22	B	811	CLA	C2C-C1C-NC	3.17	112.95	109.97
22	6	311	CLA	C2C-C1C-NC	3.17	112.95	109.97
22	B	811	CLA	C1-C2-C3	-3.17	120.55	126.04
22	A	802	CLA	CHD-C1D-ND	-3.17	121.54	124.45
22	2	301	CLA	CMA-C3A-C4A	3.17	120.30	111.77
22	Z	304	CLA	C2D-C1D-ND	3.17	112.44	110.10
22	Z	305	CLA	C1-C2-C3	-3.17	120.56	126.04
22	A	810	CLA	CHD-C1D-ND	-3.17	121.54	124.45
22	A	819	CLA	C1-C2-C3	-3.17	120.56	126.04
22	A	823	CLA	C2D-C1D-ND	3.17	112.44	110.10
24	8	304	BCR	C33-C5-C4	3.17	119.71	113.62
22	K	204	CLA	C2C-C1C-NC	3.17	112.94	109.97
22	Z	310	CLA	C2D-C1D-ND	3.17	112.44	110.10
22	5	319	CLA	C2C-C1C-NC	3.17	112.94	109.97
22	7	312	CLA	C2D-C1D-ND	3.17	112.44	110.10
22	3	310	CLA	C2C-C1C-NC	3.17	112.94	109.97
22	B	813	CLA	C2D-C1D-ND	3.17	112.44	110.10
22	6	311	CLA	CHD-C1D-ND	-3.17	121.54	124.45
22	1	310	CLA	C2C-C1C-NC	3.17	112.94	109.97
22	4	807	CLA	C2D-C1D-ND	3.17	112.44	110.10
24	8	304	BCR	C38-C26-C25	-3.16	120.97	124.53
24	B	849	BCR	C33-C5-C4	3.16	119.69	113.62
22	6	314	CLA	C1-C2-C3	-3.16	120.57	126.04
22	Z	303	CLA	CHD-C1D-ND	-3.16	121.55	124.45
22	4	811	CLA	C2D-C1D-ND	3.16	112.43	110.10
22	3	322	CLA	CHD-C1D-ND	-3.16	121.55	124.45
24	B	849	BCR	C33-C5-C6	-3.16	120.98	124.53
22	A	839	CLA	CHD-C1D-ND	-3.16	121.55	124.45
22	B	832	CLA	CHD-C1D-ND	-3.16	121.55	124.45
39	5	321	CHL	C3C-C4C-NC	-3.16	107.03	110.57
36	5	306	C7Z	C21-C26-C25	-3.16	118.16	122.61
22	A	837	CLA	C1-C2-C3	-3.16	120.58	126.04
22	8	307	CLA	C2C-C1C-NC	3.16	112.93	109.97
22	B	821	CLA	CHD-C1D-ND	-3.15	121.56	124.45
22	A	823	CLA	CHD-C1D-ND	-3.15	121.56	124.45
22	3	319	CLA	CHD-C1D-ND	-3.15	121.56	124.45
22	5	309	CLA	C2D-C1D-ND	3.15	112.43	110.10
22	3	319	CLA	C2C-C1C-NC	3.15	112.92	109.97
22	B	819	CLA	C1-C2-C3	-3.15	120.59	126.04
22	3	318	CLA	C2D-C1D-ND	3.15	112.43	110.10

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Mol	Chain	Res	Type	Atoms	Z	Observed(^{o})	Ideal(^{o})
22	3	312	CLA	C2C-C1C-NC	3.15	112.92	109.97
22	A	803	CLA	C2D-C1D-ND	3.15	112.42	110.10
24	B	844	BCR	C24-C23-C22	-3.15	121.48	126.23
22	B	842	CLA	C2C-C1C-NC	3.15	112.92	109.97
22	B	841	CLA	C2D-C1D-ND	3.15	112.42	110.10
22	B	801	CLA	CHD-C1D-ND	-3.15	121.56	124.45
22	K	204	CLA	C2D-C1D-ND	3.15	112.42	110.10
22	J	103	CLA	C2D-C1D-ND	3.14	112.42	110.10
22	5	323	CLA	C2D-C1D-ND	3.14	112.42	110.10
22	Z	316	CLA	CHD-C1D-ND	-3.14	121.56	124.45
24	G	203	BCR	C8-C7-C6	-3.14	118.37	127.20
22	B	830	CLA	C2D-C1D-ND	3.14	112.42	110.10
22	2	302	CLA	C2D-C1D-ND	3.14	112.42	110.10
22	B	836	CLA	CHD-C1D-ND	-3.14	121.57	124.45
25	9	315	LHG	O7-C7-C8	3.14	118.27	111.50
22	7	311	CLA	CHD-C1D-ND	-3.14	121.57	124.45
22	A	827	CLA	C2C-C1C-NC	3.14	112.92	109.97
22	B	811	CLA	C2D-C1D-ND	3.14	112.42	110.10
22	6	317	CLA	C2D-C1D-ND	3.14	112.42	110.10
39	6	316	CHL	C4D-CHA-C1A	3.14	125.07	121.25
22	5	301	CLA	C1-C2-C3	-3.14	120.61	126.04
22	9	303	CLA	O2A-CGA-CBA	3.14	121.76	111.91
38	7	301	LUT	C31-C30-C29	-3.14	122.83	127.31
22	B	808	CLA	CHD-C1D-ND	-3.14	121.57	124.45
22	7	314	CLA	CHD-C1D-ND	-3.14	121.57	124.45
22	8	313	CLA	C2D-C1D-ND	3.14	112.42	110.10
22	4	809	CLA	C2C-C1C-NC	3.14	112.91	109.97
22	6	302	CLA	C2C-C1C-NC	3.14	112.91	109.97
24	A	848	BCR	C33-C5-C6	-3.14	121.01	124.53
22	Z	307	CLA	CMA-C3A-C4A	3.14	120.20	111.77
24	B	849	BCR	C11-C10-C9	-3.13	122.84	127.31
22	B	817	CLA	C2D-C1D-ND	3.13	112.41	110.10
22	8	310	CLA	CHD-C1D-ND	-3.13	121.58	124.45
38	7	302	LUT	C21-C26-C27	-3.13	108.74	112.70
22	B	812	CLA	C2C-C1C-NC	3.13	112.91	109.97
22	8	305	CLA	C2C-C1C-NC	3.13	112.91	109.97
24	A	848	BCR	C20-C19-C18	-3.13	117.62	126.42
22	A	854	CLA	CHD-C1D-ND	-3.13	121.58	124.45
22	1	305	CLA	C2C-C1C-NC	3.13	112.90	109.97
22	7	312	CLA	C2C-C1C-NC	3.13	112.90	109.97
22	4	808	CLA	C2C-C1C-NC	3.13	112.90	109.97
22	B	825	CLA	CHD-C1D-ND	-3.13	121.58	124.45

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Mol	Chain	Res	Type	Atoms	Z	Observed(^{o})	Ideal(^{o})
39	6	316	CHL	CMA-C3A-C4A	3.13	120.18	111.77
22	8	307	CLA	C2D-C1D-ND	3.13	112.41	110.10
22	4	815	CLA	C2D-C1D-ND	3.13	112.41	110.10
22	Z	307	CLA	C2C-C1C-NC	3.13	112.90	109.97
38	7	301	LUT	C18-C5-C4	3.13	120.15	114.36
23	A	843	PQN	C11-C12-C13	-3.13	121.59	126.79
22	8	308	CLA	C2D-C1D-ND	3.13	112.41	110.10
24	6	305	BCR	C33-C5-C4	3.13	119.62	113.62
24	3	307	BCR	C23-C22-C21	3.13	123.74	118.94
22	A	834	CLA	C1-C2-C3	-3.13	120.64	126.04
22	A	838	CLA	C2C-C1C-NC	3.13	112.90	109.97
22	3	320	CLA	C2C-C1C-NC	3.13	112.90	109.97
22	A	825	CLA	CMA-C3A-C4A	3.12	120.17	111.77
22	A	805	CLA	C1-C2-C3	-3.12	120.64	126.04
22	A	808	CLA	C2C-C1C-NC	3.12	112.90	109.97
24	3	307	BCR	C37-C22-C21	-3.12	118.55	122.92
22	A	808	CLA	C2D-C1D-ND	3.12	112.41	110.10
22	2	304	CLA	C2C-C1C-NC	3.12	112.90	109.97
39	1	312	CHL	CMA-C3A-C4A	3.12	120.16	111.77
22	1	307	CLA	CMA-C3A-C4A	3.12	120.16	111.77
22	6	322	CLA	C2C-C1C-NC	3.12	112.90	109.97
22	B	828	CLA	C2D-C1D-ND	3.12	112.40	110.10
22	6	302	CLA	C2D-C1D-ND	3.12	112.40	110.10
22	8	305	CLA	CMA-C3A-C4A	3.12	120.16	111.77
22	B	811	CLA	CMA-C3A-C4A	3.12	120.16	111.77
22	6	314	CLA	CMA-C3A-C4A	3.12	120.16	111.77
22	Z	310	CLA	CHD-C1D-ND	-3.12	121.59	124.45
22	3	316	CLA	C2C-C1C-NC	3.12	112.89	109.97
22	3	310	CLA	CMA-C3A-C4A	3.12	120.15	111.77
39	6	316	CHL	C1-C2-C3	-3.12	121.71	126.75
22	A	809	CLA	C2D-C1D-ND	3.12	112.40	110.10
22	B	819	CLA	C2D-C1D-ND	3.12	112.40	110.10
22	B	823	CLA	C2D-C1D-ND	3.12	112.40	110.10
22	9	311	CLA	C2D-C1D-ND	3.12	112.40	110.10
22	Z	313	CLA	C2C-C1C-NC	3.12	112.89	109.97
22	B	819	CLA	CHD-C1D-ND	-3.12	121.59	124.45
22	8	305	CLA	CHD-C1D-ND	-3.12	121.59	124.45
22	B	836	CLA	C2D-C1D-ND	3.12	112.40	110.10
22	5	311	CLA	C2D-C1D-ND	3.12	112.40	110.10
22	A	837	CLA	CHD-C1D-ND	-3.11	121.59	124.45
22	4	812	CLA	C2D-C1D-ND	3.11	112.40	110.10
22	A	816	CLA	CMA-C3A-C4A	3.11	120.14	111.77

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Mol	Chain	Res	Type	Atoms	Z	Observed(^{o})	Ideal(^{o})
22	A	829	CLA	C2C-C1C-NC	3.11	112.89	109.97
22	A	809	CLA	CHD-C1D-ND	-3.11	121.59	124.45
22	9	308	CLA	CMA-C3A-C4A	3.11	120.14	111.77
22	A	811	CLA	C2D-C1D-ND	3.11	112.40	110.10
22	A	841	CLA	C2D-C1D-ND	3.11	112.40	110.10
22	B	827	CLA	C2C-C1C-NC	3.11	112.89	109.97
22	B	807	CLA	C2D-C1D-ND	3.11	112.40	110.10
22	B	808	CLA	C2D-C1D-ND	3.11	112.40	110.10
22	B	813	CLA	C2C-C1C-NC	3.11	112.89	109.97
22	7	310	CLA	C2C-C1C-NC	3.11	112.89	109.97
24	5	305	BCR	C36-C18-C19	3.11	122.97	118.08
22	B	820	CLA	C2C-C1C-NC	3.11	112.88	109.97
22	2	302	CLA	C2C-C1C-NC	3.11	112.88	109.97
22	A	819	CLA	CHD-C1D-ND	-3.11	121.60	124.45
39	4	819	CHL	C3C-C4C-NC	-3.11	107.09	110.57
22	A	829	CLA	C2D-C1D-ND	3.11	112.39	110.10
22	3	309	CLA	C2D-C1D-ND	3.11	112.39	110.10
22	B	829	CLA	C2C-C1C-NC	3.11	112.88	109.97
39	8	301	CHL	C2C-C3C-C4C	3.11	108.70	106.49
22	1	309	CLA	CHD-C1D-ND	-3.11	121.60	124.45
22	8	313	CLA	CHD-C1D-ND	-3.11	121.60	124.45
38	3	303	LUT	C1-C6-C5	-3.11	118.24	122.61
23	B	843	PQN	C14-C13-C15	3.11	120.50	115.27
22	B	840	CLA	C1-C2-C3	-3.10	120.67	126.04
39	4	816	CHL	C3C-C4C-NC	-3.10	107.09	110.57
22	A	838	CLA	C2D-C1D-ND	3.10	112.39	110.10
22	A	832	CLA	CHD-C1D-ND	-3.10	121.60	124.45
22	B	840	CLA	CHD-C1D-ND	-3.10	121.60	124.45
22	8	308	CLA	CHD-C1D-ND	-3.10	121.60	124.45
22	9	307	CLA	CMA-C3A-C4A	3.10	120.11	111.77
22	A	826	CLA	C2D-C1D-ND	3.10	112.39	110.10
22	B	817	CLA	C1-C2-C3	-3.10	120.68	126.04
24	3	306	BCR	C33-C5-C4	3.10	119.58	113.62
22	A	836	CLA	CHD-C1D-ND	-3.10	121.60	124.45
22	7	316	CLA	C2D-C1D-ND	3.10	112.39	110.10
22	4	809	CLA	C2D-C1D-ND	3.10	112.39	110.10
22	3	316	CLA	C6-C5-C3	-3.10	109.55	114.62
22	7	309	CLA	CHD-C1D-ND	-3.10	121.60	124.45
22	3	312	CLA	C2D-C1D-ND	3.10	112.39	110.10
22	A	811	CLA	CHD-C1D-ND	-3.10	121.61	124.45
22	B	829	CLA	CHD-C1D-ND	-3.10	121.61	124.45
22	A	840	CLA	C2C-C1C-NC	3.10	112.88	109.97

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Mol	Chain	Res	Type	Atoms	Z	Observed(^{o})	Ideal(^{o})
22	7	323	CLA	C2C-C1C-NC	3.10	112.88	109.97
24	B	847	BCR	C11-C12-C13	-3.10	117.71	126.42
24	B	848	BCR	C38-C26-C27	3.10	119.57	113.62
22	A	802	CLA	C2C-C1C-NC	3.10	112.87	109.97
38	6	304	LUT	C1-C6-C5	-3.10	118.25	122.61
38	5	302	LUT	C21-C26-C27	-3.10	108.79	112.70
24	B	853	BCR	C33-C5-C4	3.10	119.56	113.62
22	A	820	CLA	C2C-C1C-NC	3.10	112.87	109.97
22	5	314	CLA	C2C-C1C-NC	3.10	112.87	109.97
22	3	311	CLA	C2D-C1D-ND	3.10	112.39	110.10
22	7	304	CLA	CMA-C3A-C4A	3.10	120.09	111.77
22	4	808	CLA	CHD-C1D-ND	-3.09	121.61	124.45
22	5	314	CLA	C2D-C1D-ND	3.09	112.38	110.10
22	A	818	CLA	CHD-C1D-ND	-3.09	121.61	124.45
22	A	837	CLA	C2C-C1C-NC	3.09	112.87	109.97
22	B	839	CLA	CHD-C1D-ND	-3.09	121.61	124.45
22	6	307	CLA	C2C-C1C-NC	3.09	112.87	109.97
22	1	315	CLA	C2C-C1C-NC	3.09	112.87	109.97
22	5	318	CLA	C2D-C1D-ND	3.09	112.38	110.10
22	9	313	CLA	C2D-C1D-ND	3.09	112.38	110.10
22	9	310	CLA	C2C-C1C-NC	3.09	112.87	109.97
22	A	806	CLA	C2C-C1C-NC	3.09	112.87	109.97
22	B	833	CLA	C2C-C1C-NC	3.09	112.87	109.97
22	7	305	CLA	C2C-C1C-NC	3.09	112.87	109.97
22	1	311	CLA	C2D-C1D-ND	3.09	112.38	110.10
22	B	833	CLA	CHD-C1D-ND	-3.09	121.62	124.45
38	6	303	LUT	C11-C10-C9	-3.09	122.90	127.31
22	B	825	CLA	C2D-C1D-ND	3.09	112.38	110.10
22	4	817	CLA	C2D-C1D-ND	3.09	112.38	110.10
39	6	315	CHL	C4D-CHA-C1A	3.09	125.00	121.25
22	1	313	CLA	CHD-C1D-ND	-3.09	121.62	124.45
22	3	314	CLA	C2C-C1C-NC	3.09	112.86	109.97
22	6	301	CLA	C2D-C1D-ND	3.09	112.38	110.10
22	5	301	CLA	C2D-C1D-ND	3.08	112.38	110.10
22	A	804	CLA	C2C-C1C-NC	3.08	112.86	109.97
22	Z	316	CLA	C2D-C1D-ND	3.08	112.38	110.10
22	B	807	CLA	CHD-C1D-ND	-3.08	121.62	124.45
22	4	810	CLA	CHD-C1D-ND	-3.08	121.62	124.45
22	5	318	CLA	CHD-C1D-ND	-3.08	121.62	124.45
22	7	323	CLA	C2D-C1D-ND	3.08	112.38	110.10
22	B	812	CLA	C1-C2-C3	-3.08	120.71	126.04
24	B	847	BCR	C16-C17-C18	-3.08	122.91	127.31

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Mol	Chain	Res	Type	Atoms	Z	Observed(^{o})	Ideal(^{o})
38	6	303	LUT	C22-C23-C24	3.08	115.25	111.74
22	9	305	CLA	C2A-C1A-CHA	3.08	129.25	123.86
22	6	322	CLA	C1-C2-C3	-3.08	120.72	126.04
22	Z	310	CLA	C2C-C1C-NC	3.08	112.86	109.97
22	5	308	CLA	CHD-C1D-ND	-3.08	121.62	124.45
22	B	840	CLA	C2C-C1C-NC	3.08	112.86	109.97
22	A	831	CLA	CHD-C1D-ND	-3.08	121.63	124.45
22	7	312	CLA	CHD-C1D-ND	-3.08	121.63	124.45
22	B	841	CLA	C2C-C1C-NC	3.08	112.85	109.97
22	5	311	CLA	C2C-C1C-NC	3.08	112.85	109.97
22	8	311	CLA	C2D-C1D-ND	3.08	112.37	110.10
22	B	818	CLA	CHD-C1D-ND	-3.08	121.63	124.45
22	L	203	CLA	CHD-C1D-ND	-3.08	121.63	124.45
22	8	311	CLA	C2C-C1C-NC	3.08	112.85	109.97
22	B	823	CLA	CHD-C1D-ND	-3.08	121.63	124.45
24	A	846	BCR	C33-C5-C4	3.07	119.52	113.62
22	6	310	CLA	C2D-C1D-ND	3.07	112.37	110.10
38	7	301	LUT	C18-C5-C6	-3.07	121.08	124.53
22	5	308	CLA	C2D-C1D-ND	3.07	112.37	110.10
22	8	308	CLA	C2C-C1C-NC	3.07	112.85	109.97
22	5	301	CLA	C2C-C1C-NC	3.07	112.85	109.97
36	5	306	C7Z	C11-C12-C13	-3.07	117.78	126.42
22	A	836	CLA	C1-C2-C3	-3.07	120.73	126.04
22	4	812	CLA	CHD-C1D-ND	-3.07	121.63	124.45
22	2	301	CLA	C2C-C1C-NC	3.07	112.85	109.97
22	7	314	CLA	C2D-C1D-ND	3.07	112.37	110.10
22	4	818	CLA	C2D-C1D-ND	3.07	112.37	110.10
22	A	828	CLA	CHD-C1D-ND	-3.07	121.63	124.45
22	5	320	CLA	O2D-CGD-O1D	-3.07	117.84	123.84
22	B	825	CLA	CMA-C3A-C4A	3.07	120.02	111.77
22	B	835	CLA	C2D-C1D-ND	3.07	112.37	110.10
22	4	808	CLA	C2D-C1D-ND	3.07	112.37	110.10
22	5	319	CLA	CHD-C1D-ND	-3.07	121.63	124.45
22	F	304	CLA	C2C-C1C-NC	3.07	112.85	109.97
22	9	306	CLA	CHD-C1D-ND	-3.07	121.64	124.45
22	A	807	CLA	C2C-C1C-NC	3.07	112.84	109.97
22	A	837	CLA	C2D-C1D-ND	3.07	112.36	110.10
22	B	833	CLA	C2D-C1D-ND	3.07	112.36	110.10
22	A	821	CLA	CHD-C1D-ND	-3.07	121.64	124.45
22	3	322	CLA	C2C-C1C-NC	3.06	112.84	109.97
22	9	311	CLA	C2C-C1C-NC	3.06	112.84	109.97
36	5	306	C7Z	C18-C5-C4	3.06	120.03	114.36

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Mol	Chain	Res	Type	Atoms	Z	Observed(^{o})	Ideal(^{o})
22	6	321	CLA	C2C-C1C-NC	3.06	112.84	109.97
22	L	201	CLA	CHD-C1D-ND	-3.06	121.64	124.45
38	1	302	LUT	C18-C5-C6	-3.06	121.09	124.53
22	1	306	CLA	C2C-C1C-NC	3.06	112.84	109.97
22	6	313	CLA	C2C-C1C-NC	3.06	112.84	109.97
22	7	322	CLA	CHD-C1D-ND	-3.06	121.64	124.45
22	K	205	CLA	CMD-C2D-C1D	3.06	130.11	124.71
22	B	829	CLA	C2D-C1D-ND	3.06	112.36	110.10
22	A	825	CLA	CHD-C1D-ND	-3.06	121.64	124.45
39	4	814	CHL	C3C-C4C-NC	-3.06	107.14	110.57
22	1	304	CLA	C2C-C1C-NC	3.06	112.84	109.97
22	3	319	CLA	C1-C2-C3	-3.06	120.75	126.04
22	L	203	CLA	C2C-C1C-NC	3.06	112.84	109.97
22	A	834	CLA	C2D-C1D-ND	3.06	112.36	110.10
22	A	836	CLA	C2D-C1D-ND	3.06	112.36	110.10
22	7	315	CLA	C2D-C1D-ND	3.06	112.36	110.10
22	7	305	CLA	CMA-C3A-C4A	3.06	119.99	111.77
22	4	806	CLA	C2D-C1D-ND	3.06	112.36	110.10
22	A	835	CLA	CHD-C1D-ND	-3.06	121.65	124.45
39	1	312	CHL	C4D-CHA-C1A	3.06	124.97	121.25
24	B	845	BCR	C38-C26-C27	3.05	119.48	113.62
22	B	821	CLA	C2C-C1C-NC	3.05	112.83	109.97
22	G	202	CLA	C2D-C1D-ND	3.05	112.35	110.10
22	7	309	CLA	C2C-C1C-NC	3.05	112.83	109.97
24	A	844	BCR	C38-C26-C27	3.05	119.48	113.62
39	6	320	CHL	C3C-C4C-NC	-3.05	107.15	110.57
22	B	809	CLA	C2C-C1C-NC	3.05	112.83	109.97
22	7	317	CLA	CHD-C1D-ND	-3.05	121.65	124.45
22	Z	309	CLA	C2C-C1C-NC	3.05	112.83	109.97
22	F	305	CLA	CHD-C1D-ND	-3.05	121.65	124.45
22	Z	304	CLA	C2C-C1C-NC	3.05	112.83	109.97
24	A	846	BCR	C1-C6-C5	-3.05	118.32	122.61
22	K	204	CLA	CMA-C3A-C4A	3.05	119.97	111.77
22	B	835	CLA	C1D-ND-C4D	-3.05	104.17	106.33
22	B	827	CLA	CHD-C1D-ND	-3.05	121.65	124.45
22	1	314	CLA	C1-C2-C3	-3.05	120.77	126.04
24	3	306	BCR	C38-C26-C27	3.05	119.47	113.62
22	6	319	CLA	C2D-C1D-ND	3.05	112.35	110.10
22	K	202	CLA	C2C-C1C-NC	3.05	112.83	109.97
22	5	312	CLA	C2C-C1C-NC	3.05	112.83	109.97
22	6	317	CLA	C2C-C1C-NC	3.05	112.83	109.97
22	A	835	CLA	C2D-C1D-ND	3.05	112.35	110.10

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Mol	Chain	Res	Type	Atoms	Z	Observed(^{o})	Ideal(^{o})
22	K	202	CLA	C2D-C1D-ND	3.05	112.35	110.10
22	A	834	CLA	CHD-C1D-ND	-3.04	121.66	124.45
22	B	837	CLA	C2C-C1C-NC	3.04	112.82	109.97
22	1	307	CLA	C2C-C1C-NC	3.04	112.82	109.97
24	A	844	BCR	C4-C5-C6	-3.04	118.31	122.73
24	6	305	BCR	C38-C26-C27	3.04	119.46	113.62
22	6	312	CLA	C2D-C1D-ND	3.04	112.35	110.10
22	6	314	CLA	CHD-C1D-ND	-3.04	121.66	124.45
22	A	832	CLA	C2D-C1D-ND	3.04	112.35	110.10
22	1	304	CLA	C2D-C1D-ND	3.04	112.35	110.10
22	8	314	CLA	C2C-C1C-NC	3.04	112.82	109.97
22	4	811	CLA	C2C-C1C-NC	3.04	112.82	109.97
22	Z	314	CLA	C1-C2-C3	-3.04	120.78	126.04
24	3	305	BCR	C1-C6-C5	-3.04	118.33	122.61
22	B	816	CLA	CHD-C1D-ND	-3.04	121.66	124.45
39	6	318	CHL	C4D-CHA-C1A	3.04	124.95	121.25
22	8	310	CLA	C2C-C1C-NC	3.04	112.82	109.97
22	4	810	CLA	C2C-C1C-NC	3.04	112.82	109.97
22	A	831	CLA	C2D-C1D-ND	3.04	112.34	110.10
22	B	827	CLA	C2D-C1D-ND	3.04	112.34	110.10
22	J	103	CLA	CHD-C1D-ND	-3.04	121.66	124.45
22	5	310	CLA	CHD-C1D-ND	-3.04	121.66	124.45
22	9	308	CLA	CHD-C1D-ND	-3.04	121.66	124.45
22	1	313	CLA	C2C-C1C-NC	3.04	112.82	109.97
22	3	313	CLA	C1-C2-C3	-3.04	120.79	126.04
22	K	203	CLA	C2D-C1D-ND	3.04	112.34	110.10
22	7	324	CLA	C2C-C1C-NC	3.04	112.82	109.97
22	A	805	CLA	CMA-C3A-C4A	3.04	119.94	111.77
22	A	832	CLA	C2C-C1C-NC	3.04	112.82	109.97
22	6	301	CLA	C2C-C1C-NC	3.04	112.82	109.97
24	B	853	BCR	C36-C18-C17	-3.04	118.67	122.92
39	Z	311	CHL	CMA-C3A-C4A	3.04	119.93	111.77
24	3	307	BCR	C36-C18-C19	3.04	122.86	118.08
22	3	309	CLA	CHD-C1D-ND	-3.04	121.66	124.45
22	7	324	CLA	CHD-C1D-ND	-3.04	121.66	124.45
22	B	801	CLA	C2D-C1D-ND	3.04	112.34	110.10
22	B	810	CLA	C2D-C1D-ND	3.04	112.34	110.10
22	B	832	CLA	C2D-C1D-ND	3.04	112.34	110.10
22	A	842	CLA	CHD-C1D-ND	-3.03	121.67	124.45
22	Z	315	CLA	CHD-C1D-ND	-3.03	121.67	124.45
22	A	825	CLA	C2C-C1C-NC	3.03	112.81	109.97
22	7	304	CLA	C2C-C1C-NC	3.03	112.81	109.97

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Mol	Chain	Res	Type	Atoms	Z	Observed(^{o})	Ideal(^{o})
22	A	807	CLA	C2D-C1D-ND	3.03	112.34	110.10
22	1	306	CLA	CHD-C1D-ND	-3.03	121.67	124.45
24	4	804	BCR	C33-C5-C6	-3.03	121.12	124.53
22	L	201	CLA	C1D-ND-C4D	-3.03	104.18	106.33
24	B	853	BCR	C11-C10-C9	-3.03	122.98	127.31
22	B	834	CLA	C2D-C1D-ND	3.03	112.34	110.10
22	K	202	CLA	CMA-C3A-C4A	3.03	119.92	111.77
39	2	303	CHL	C3C-C4C-NC	-3.03	107.17	110.57
22	6	321	CLA	CHD-C1D-ND	-3.03	121.67	124.45
22	9	306	CLA	C2C-C1C-NC	3.03	112.81	109.97
24	7	303	BCR	C7-C8-C9	-3.03	121.66	126.23
22	A	802	CLA	C2D-C1D-ND	3.03	112.34	110.10
22	G	201	CLA	C2D-C1D-ND	3.03	112.34	110.10
22	A	808	CLA	CHD-C1D-ND	-3.03	121.67	124.45
22	8	309	CLA	C2C-C1C-NC	3.03	112.81	109.97
22	8	313	CLA	C2C-C1C-NC	3.03	112.81	109.97
22	5	326	CLA	C2C-C1C-NC	3.03	112.81	109.97
22	A	855	CLA	CHD-C1D-ND	-3.03	121.67	124.45
36	J	105	C7Z	C38-C25-C24	3.03	119.96	114.36
22	L	201	CLA	C2C-C1C-NC	3.03	112.81	109.97
22	5	307	CLA	C2C-C1C-NC	3.03	112.81	109.97
22	5	322	CLA	C2C-C1C-NC	3.03	112.81	109.97
22	8	309	CLA	C2D-C1D-ND	3.03	112.33	110.10
22	B	831	CLA	CHD-C1D-ND	-3.03	121.67	124.45
22	A	803	CLA	CHD-C1D-ND	-3.02	121.67	124.45
22	7	308	CLA	C2C-C1C-NC	3.02	112.80	109.97
22	9	313	CLA	CHD-C1D-ND	-3.02	121.68	124.45
22	3	316	CLA	C1-C2-C3	-3.02	120.81	126.04
39	Z	312	CHL	C4D-CHA-C1A	3.02	124.93	121.25
22	L	202	CLA	C2D-C1D-ND	3.02	112.33	110.10
22	7	324	CLA	C2D-C1D-ND	3.02	112.33	110.10
22	A	831	CLA	C2C-C1C-NC	3.02	112.80	109.97
22	A	814	CLA	CHD-C1D-ND	-3.02	121.68	124.45
22	2	304	CLA	CMA-C3A-C4A	3.02	119.89	111.77
22	A	821	CLA	C2C-C1C-NC	3.02	112.80	109.97
22	F	304	CLA	C2D-C1D-ND	3.02	112.33	110.10
22	9	313	CLA	C2C-C1C-NC	3.02	112.80	109.97
22	5	315	CLA	CMA-C3A-C4A	3.02	119.89	111.77
22	B	822	CLA	C2D-C1D-ND	3.02	112.33	110.10
22	3	319	CLA	C2D-C1D-ND	3.02	112.33	110.10
22	5	312	CLA	CHD-C1D-ND	-3.02	121.68	124.45
21	A	801	CL0	CMC-C2C-C1C	3.02	129.63	125.04

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Mol	Chain	Res	Type	Atoms	Z	Observed(^{o})	Ideal(^{o})
22	6	308	CLA	CHD-C1D-ND	-3.02	121.68	124.45
24	3	305	BCR	C16-C17-C18	-3.02	123.00	127.31
22	A	822	CLA	CMA-C3A-C4A	3.02	119.88	111.77
22	1	309	CLA	C2D-C1D-ND	3.02	112.33	110.10
22	7	308	CLA	C2D-C1D-ND	3.02	112.33	110.10
22	B	832	CLA	C2C-C1C-NC	3.02	112.80	109.97
22	J	103	CLA	C2C-C1C-NC	3.02	112.80	109.97
22	9	306	CLA	C2D-C1D-ND	3.02	112.33	110.10
22	B	828	CLA	CHD-C1D-ND	-3.01	121.68	124.45
22	7	304	CLA	C1D-ND-C4D	-3.01	104.19	106.33
22	B	830	CLA	CMA-C3A-C4A	3.01	119.87	111.77
24	3	306	BCR	C20-C21-C22	-3.01	123.01	127.31
22	6	308	CLA	C2C-C1C-NC	3.01	112.80	109.97
22	A	838	CLA	CHD-C1D-ND	-3.01	121.69	124.45
22	A	815	CLA	C2D-C1D-ND	3.01	112.32	110.10
22	B	830	CLA	C2C-C1C-NC	3.01	112.79	109.97
22	B	812	CLA	CHD-C1D-ND	-3.01	121.69	124.45
22	B	813	CLA	CHD-C1D-ND	-3.01	121.69	124.45
22	7	323	CLA	CHD-C1D-ND	-3.01	121.69	124.45
22	9	307	CLA	C2C-C1C-NC	3.01	112.79	109.97
24	A	846	BCR	C27-C26-C25	-3.01	118.36	122.73
24	8	304	BCR	C27-C26-C25	-3.01	118.36	122.73
22	A	806	CLA	C2D-C1D-ND	3.01	112.32	110.10
22	B	823	CLA	CMA-C3A-C4A	3.01	119.86	111.77
38	4	803	LUT	C18-C5-C4	3.01	119.93	114.36
22	A	823	CLA	CMA-C3A-C4A	3.01	119.86	111.77
39	5	316	CHL	C4D-CHA-C1A	3.01	124.91	121.25
22	F	302	CLA	CMA-C3A-C4A	3.01	119.86	111.77
22	A	824	CLA	C2C-C1C-NC	3.01	112.79	109.97
38	7	302	LUT	C31-C30-C29	-3.01	123.02	127.31
22	1	306	CLA	CMA-C3A-C4A	3.01	119.85	111.77
22	F	301	CLA	C2C-C1C-NC	3.01	112.79	109.97
22	7	306	CLA	C2D-C1D-ND	3.01	112.32	110.10
22	A	841	CLA	CHD-C1D-ND	-3.01	121.69	124.45
22	4	817	CLA	CHD-C1D-ND	-3.01	121.69	124.45
22	4	818	CLA	CHD-C1D-ND	-3.01	121.69	124.45
22	A	841	CLA	C1D-ND-C4D	-3.01	104.20	106.33
22	5	309	CLA	CMA-C3A-C4A	3.00	119.85	111.77
24	B	848	BCR	C33-C5-C4	3.00	119.39	113.62
22	7	316	CLA	CHD-C1D-ND	-3.00	121.69	124.45
22	B	831	CLA	C2C-C1C-NC	3.00	112.79	109.97
22	B	814	CLA	CHD-C1D-ND	-3.00	121.69	124.45

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Mol	Chain	Res	Type	Atoms	Z	Observed(^{o})	Ideal(^{o})
22	5	319	CLA	C2D-C1D-ND	3.00	112.32	110.10
22	B	842	CLA	CHD-C1D-ND	-3.00	121.69	124.45
22	6	313	CLA	C2D-C1D-ND	3.00	112.32	110.10
22	A	818	CLA	C1D-ND-C4D	-3.00	104.20	106.33
22	B	818	CLA	C1D-ND-C4D	-3.00	104.20	106.33
22	A	820	CLA	C2D-C1D-ND	3.00	112.31	110.10
22	B	826	CLA	CHD-C1D-ND	-3.00	121.70	124.45
22	1	304	CLA	CHD-C1D-ND	-3.00	121.70	124.45
22	B	826	CLA	C2C-C1C-NC	3.00	112.78	109.97
22	4	806	CLA	C2C-C1C-NC	3.00	112.78	109.97
22	B	807	CLA	CMA-C3A-C4A	3.00	119.83	111.77
38	5	302	LUT	C1-C6-C5	-3.00	118.39	122.61
22	4	811	CLA	C1-C2-C3	-3.00	120.86	126.04
22	A	825	CLA	C2D-C1D-ND	3.00	112.31	110.10
22	3	314	CLA	C2D-C1D-ND	3.00	112.31	110.10
22	A	838	CLA	CMA-C3A-C4A	3.00	119.83	111.77
22	F	302	CLA	C2C-C1C-NC	3.00	112.78	109.97
22	6	309	CLA	CMA-C3A-C4A	3.00	119.83	111.77
22	B	824	CLA	C1D-ND-C4D	-3.00	104.21	106.33
22	F	301	CLA	CHD-C1D-ND	-3.00	121.70	124.45
22	6	312	CLA	CHD-C1D-ND	-3.00	121.70	124.45
22	A	814	CLA	C2D-C1D-ND	3.00	112.31	110.10
22	L	203	CLA	C2D-C1D-ND	2.99	112.31	110.10
22	9	303	CLA	CHD-C1D-ND	-2.99	121.70	124.45
22	Z	314	CLA	C2C-C1C-NC	2.99	112.78	109.97
22	8	316	CLA	C2C-C1C-NC	2.99	112.78	109.97
22	B	823	CLA	C1D-ND-C4D	-2.99	104.21	106.33
22	7	311	CLA	C1D-ND-C4D	-2.99	104.21	106.33
22	A	830	CLA	CMA-C3A-C4A	2.99	119.82	111.77
22	B	815	CLA	CHD-C1D-ND	-2.99	121.70	124.45
22	A	818	CLA	CMA-C3A-C4A	2.99	119.82	111.77
22	B	811	CLA	C1D-ND-C4D	-2.99	104.21	106.33
22	A	818	CLA	C2C-C1C-NC	2.99	112.78	109.97
22	7	311	CLA	C2C-C1C-NC	2.99	112.78	109.97
22	1	308	CLA	C1C-C2C-C3C	-2.99	103.81	106.96
22	A	829	CLA	CMA-C3A-C4A	2.99	119.81	111.77
22	A	807	CLA	CHD-C1D-ND	-2.99	121.70	124.45
22	L	202	CLA	CHD-C1D-ND	-2.99	121.70	124.45
22	6	321	CLA	CMA-C3A-C4A	2.99	119.81	111.77
22	A	830	CLA	C2C-C1C-NC	2.99	112.77	109.97
22	6	322	CLA	CHD-C1D-ND	-2.99	121.71	124.45
22	9	310	CLA	CMA-C3A-C4A	2.99	119.81	111.77

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Mol	Chain	Res	Type	Atoms	Z	Observed(^{o})	Ideal(^{o})
22	A	828	CLA	C2C-C1C-NC	2.99	112.77	109.97
22	B	810	CLA	CMA-C3A-C4A	2.99	119.81	111.77
22	A	833	CLA	CHD-C1D-ND	-2.99	121.71	124.45
22	B	805	CLA	C2C-C1C-NC	2.99	112.77	109.97
22	L	202	CLA	C2C-C1C-NC	2.99	112.77	109.97
22	7	314	CLA	C2C-C1C-NC	2.99	112.77	109.97
39	8	315	CHL	C3C-C4C-NC	-2.99	107.22	110.57
22	A	813	CLA	C1-C2-C3	-2.99	120.88	126.04
22	7	314	CLA	CMA-C3A-C4A	2.99	119.80	111.77
22	A	836	CLA	C2C-C1C-NC	2.99	112.77	109.97
22	A	828	CLA	C2D-C1D-ND	2.99	112.31	110.10
22	3	310	CLA	C2D-C1D-ND	2.99	112.31	110.10
22	Z	308	CLA	C1D-ND-C4D	-2.99	104.21	106.33
22	4	815	CLA	CMA-C3A-C4A	2.99	119.80	111.77
22	Z	306	CLA	CHD-C1D-ND	-2.98	121.71	124.45
39	8	312	CHL	C4D-CHA-C1A	2.98	124.88	121.25
22	A	805	CLA	C1D-ND-C4D	-2.98	104.22	106.33
24	B	846	BCR	C34-C9-C10	-2.98	118.74	122.92
22	4	810	CLA	C2D-C1D-ND	2.98	112.30	110.10
22	5	308	CLA	C2C-C1C-NC	2.98	112.77	109.97
22	B	836	CLA	CMA-C3A-C4A	2.98	119.79	111.77
22	B	838	CLA	C2C-C1C-NC	2.98	112.77	109.97
22	A	811	CLA	C1-C2-C3	-2.98	120.89	126.04
22	7	306	CLA	CMA-C3A-C4A	2.98	119.79	111.77
22	8	309	CLA	CHD-C1D-ND	-2.98	121.71	124.45
22	A	826	CLA	C2C-C1C-NC	2.98	112.77	109.97
22	3	308	CLA	C2C-C1C-NC	2.98	112.77	109.97
22	5	323	CLA	C2C-C1C-NC	2.98	112.77	109.97
24	A	844	BCR	C33-C5-C4	2.98	119.34	113.62
22	A	830	CLA	CHD-C1D-ND	-2.98	121.72	124.45
22	K	203	CLA	CHD-C1D-ND	-2.98	121.72	124.45
22	A	815	CLA	C2C-C1C-NC	2.98	112.76	109.97
22	7	317	CLA	C2C-C1C-NC	2.98	112.76	109.97
22	A	854	CLA	C2D-C1D-ND	2.98	112.30	110.10
22	9	308	CLA	C2D-C1D-ND	2.98	112.30	110.10
22	B	808	CLA	C1-C2-C3	-2.98	120.89	126.04
22	B	819	CLA	C2C-C1C-NC	2.98	112.76	109.97
24	K	206	BCR	C33-C5-C4	2.98	119.34	113.62
22	A	813	CLA	CHD-C1D-ND	-2.98	121.72	124.45
22	A	840	CLA	CHD-C1D-ND	-2.98	121.72	124.45
22	3	316	CLA	CHD-C1D-ND	-2.98	121.72	124.45
22	Z	316	CLA	C2C-C1C-NC	2.98	112.76	109.97

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Mol	Chain	Res	Type	Atoms	Z	Observed(^{o})	Ideal(^{o})
22	7	307	CLA	C2C-C1C-NC	2.98	112.76	109.97
38	1	302	LUT	C31-C30-C29	-2.98	123.06	127.31
22	6	301	CLA	CHD-C1D-ND	-2.98	121.72	124.45
22	A	834	CLA	C2C-C1C-NC	2.98	112.76	109.97
39	9	314	CHL	C4D-CHA-C1A	2.98	124.87	121.25
22	B	814	CLA	C1-C2-C3	-2.98	120.89	126.04
22	A	810	CLA	CMA-C3A-C4A	2.98	119.77	111.77
22	A	819	CLA	CMA-C3A-C4A	2.98	119.77	111.77
38	7	301	LUT	C21-C26-C27	-2.98	108.94	112.70
22	3	322	CLA	CMA-C3A-C4A	2.97	119.77	111.77
39	6	315	CHL	C3C-C4C-NC	-2.97	107.24	110.57
22	1	305	CLA	C2D-C1D-ND	2.97	112.30	110.10
39	7	313	CHL	C3C-C4C-NC	-2.97	107.24	110.57
24	A	856	BCR	C38-C26-C27	2.97	119.33	113.62
22	6	321	CLA	C2D-C1D-ND	2.97	112.29	110.10
22	A	855	CLA	CMA-C3A-C4A	2.97	119.76	111.77
22	3	311	CLA	CHD-C1D-ND	-2.97	121.72	124.45
22	5	309	CLA	CHD-C1D-ND	-2.97	121.72	124.45
38	9	302	LUT	C21-C26-C27	-2.97	108.95	112.70
22	K	205	CLA	C1-C2-C3	-2.97	120.91	126.04
39	4	813	CHL	C4D-CHA-C1A	2.97	124.86	121.25
22	1	305	CLA	CHD-C1D-ND	-2.97	121.72	124.45
23	B	843	PQN	C11-C12-C13	-2.97	121.85	126.79
39	8	312	CHL	CMA-C3A-C4A	2.97	119.75	111.77
22	1	304	CLA	CMA-C3A-C4A	2.97	119.75	111.77
22	A	827	CLA	CHD-C1D-ND	-2.97	121.73	124.45
22	A	819	CLA	C2C-C1C-NC	2.97	112.75	109.97
22	1	309	CLA	C2C-C1C-NC	2.97	112.75	109.97
22	5	326	CLA	C1-C2-C3	-2.97	120.91	126.04
24	B	847	BCR	C4-C5-C6	-2.97	118.42	122.73
22	B	817	CLA	C2C-C1C-NC	2.97	112.75	109.97
22	1	314	CLA	C2C-C1C-NC	2.97	112.75	109.97
22	7	316	CLA	C2C-C1C-NC	2.97	112.75	109.97
22	B	837	CLA	CMA-C3A-C4A	2.97	119.75	111.77
22	7	310	CLA	CMA-C3A-C4A	2.97	119.75	111.77
36	J	105	C7Z	C35-C15-C14	-2.97	117.40	123.47
39	6	318	CHL	CMA-C3A-C4A	2.96	119.74	111.77
24	B	846	BCR	C27-C26-C25	-2.96	118.43	122.73
22	A	854	CLA	C1-C2-C3	-2.96	120.92	126.04
22	A	809	CLA	C2C-C1C-NC	2.96	112.75	109.97
22	3	311	CLA	C2C-C1C-NC	2.96	112.75	109.97
22	8	316	CLA	CHD-C1D-ND	-2.96	121.73	124.45

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Mol	Chain	Res	Type	Atoms	Z	Observed(^{o})	Ideal(^{o})
24	6	305	BCR	C27-C26-C25	-2.96	118.43	122.73
22	Z	306	CLA	C2D-C1D-ND	2.96	112.29	110.10
22	3	316	CLA	C2D-C1D-ND	2.96	112.29	110.10
22	A	854	CLA	C2C-C1C-NC	2.96	112.75	109.97
22	9	305	CLA	O2D-CGD-O1D	-2.96	118.05	123.84
38	9	301	LUT	C1-C6-C5	-2.96	118.44	122.61
22	4	818	CLA	C2C-C1C-NC	2.96	112.75	109.97
22	5	310	CLA	C2C-C1C-NC	2.96	112.75	109.97
22	B	820	CLA	CHD-C1D-ND	-2.96	121.73	124.45
24	G	203	BCR	C33-C5-C4	2.96	119.31	113.62
22	6	319	CLA	C2C-C1C-NC	2.96	112.75	109.97
22	B	840	CLA	C2D-C1D-ND	2.96	112.29	110.10
22	B	820	CLA	CMA-C3A-C4A	2.96	119.73	111.77
22	B	810	CLA	CHD-C1D-ND	-2.96	121.73	124.45
22	B	841	CLA	CHD-C1D-ND	-2.96	121.73	124.45
24	B	853	BCR	C23-C24-C25	-2.96	118.89	127.20
24	3	307	BCR	C12-C13-C14	2.96	123.48	118.94
22	9	313	CLA	CMA-C3A-C4A	2.96	119.73	111.77
22	Z	315	CLA	CMA-C3A-C4A	2.96	119.72	111.77
22	3	308	CLA	CMA-C3A-C4A	2.96	119.72	111.77
22	A	806	CLA	CHD-C1D-ND	-2.96	121.74	124.45
22	6	319	CLA	CHD-C1D-ND	-2.96	121.74	124.45
22	1	306	CLA	C2D-C1D-ND	2.96	112.28	110.10
22	4	817	CLA	C2C-C1C-NC	2.96	112.74	109.97
22	1	315	CLA	CHD-C1D-ND	-2.96	121.74	124.45
39	9	314	CHL	C3C-C4C-NC	-2.95	107.26	110.57
22	3	309	CLA	C2C-C1C-NC	2.95	112.74	109.97
22	B	813	CLA	CMA-C3A-C4A	2.95	119.71	111.77
22	B	840	CLA	CMA-C3A-C4A	2.95	119.71	111.77
22	4	812	CLA	CMA-C3A-C4A	2.95	119.71	111.77
22	7	315	CLA	C2C-C1C-NC	2.95	112.74	109.97
22	4	806	CLA	CMA-C3A-C4A	2.95	119.71	111.77
22	K	202	CLA	CHD-C1D-ND	-2.95	121.74	124.45
22	1	310	CLA	C2D-C1D-ND	2.95	112.28	110.10
22	1	309	CLA	C1D-ND-C4D	-2.95	104.24	106.33
22	A	833	CLA	CMA-C3A-C4A	2.95	119.70	111.77
22	A	824	CLA	C1D-ND-C4D	-2.95	104.24	106.33
22	A	826	CLA	CHD-C1D-ND	-2.95	121.74	124.45
24	B	849	BCR	C4-C5-C6	-2.95	118.45	122.73
22	A	815	CLA	CHD-C1D-ND	-2.95	121.75	124.45
22	3	318	CLA	C1D-ND-C4D	-2.95	104.24	106.33
22	1	315	CLA	CMA-C3A-C4A	2.95	119.69	111.77

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Mol	Chain	Res	Type	Atoms	Z	Observed(^{o})	Ideal(^{o})
22	8	314	CLA	C2D-C1D-ND	2.95	112.28	110.10
39	8	301	CHL	CMA-C3A-C4A	2.95	119.69	111.77
22	3	312	CLA	C1D-ND-C4D	-2.95	104.24	106.33
22	F	305	CLA	C2C-C1C-NC	2.95	112.73	109.97
22	A	810	CLA	C2D-C1D-ND	2.95	112.28	110.10
22	A	809	CLA	CMA-C3A-C4A	2.95	119.69	111.77
22	G	202	CLA	C2C-C1C-NC	2.95	112.73	109.97
22	J	103	CLA	CMA-C3A-C4A	2.94	119.69	111.77
25	4	821	LHG	C5-O7-C7	-2.94	110.54	117.79
22	A	824	CLA	CHD-C1D-ND	-2.94	121.75	124.45
22	8	311	CLA	CMA-C3A-C4A	2.94	119.69	111.77
22	8	306	CLA	CMA-C3A-C4A	2.94	119.68	111.77
22	B	833	CLA	CMA-C3A-C4A	2.94	119.68	111.77
36	J	105	C7Z	C8-C7-C6	-2.94	118.94	127.20
22	A	855	CLA	C2D-C1D-ND	2.94	112.27	110.10
22	A	824	CLA	CMA-C3A-C4A	2.94	119.67	111.77
39	3	317	CHL	C4D-CHA-C1A	2.94	124.83	121.25
22	9	313	CLA	C1D-ND-C4D	-2.94	104.25	106.33
22	B	837	CLA	C2D-C1D-ND	2.94	112.27	110.10
22	A	802	CLA	C1-C2-C3	-2.94	120.96	126.04
22	5	314	CLA	CMA-C3A-C4A	2.94	119.67	111.77
24	A	844	BCR	C2-C1-C6	2.94	115.00	110.48
22	7	315	CLA	CMA-C3A-C4A	2.94	119.67	111.77
22	5	312	CLA	C2D-C1D-ND	2.94	112.27	110.10
22	1	311	CLA	O2D-CGD-O1D	-2.94	118.10	123.84
22	A	802	CLA	CMA-C3A-C4A	2.94	119.67	111.77
22	Z	310	CLA	CMA-C3A-C4A	2.94	119.67	111.77
22	5	318	CLA	CMA-C3A-C4A	2.94	119.66	111.77
24	B	853	BCR	C34-C9-C10	-2.94	118.81	122.92
23	A	843	PQN	C14-C13-C15	2.94	120.21	115.27
38	8	302	LUT	C8-C7-C6	-2.93	118.96	127.20
22	8	310	CLA	C2D-C1D-ND	2.93	112.27	110.10
22	Z	304	CLA	CMA-C3A-C4A	2.93	119.66	111.77
22	8	314	CLA	CHD-C1D-ND	-2.93	121.76	124.45
22	2	302	CLA	CHD-C1D-ND	-2.93	121.76	124.45
22	Z	309	CLA	C1D-ND-C4D	-2.93	104.25	106.33
22	Z	316	CLA	C1D-ND-C4D	-2.93	104.25	106.33
22	B	824	CLA	CMA-C3A-C4A	2.93	119.66	111.77
24	3	306	BCR	C11-C12-C13	-2.93	118.18	126.42
22	B	806	CLA	CMA-C3A-C4A	2.93	119.65	111.77
36	J	105	C7Z	C15-C14-C13	-2.93	123.13	127.31
22	2	302	CLA	CMA-C3A-C4A	2.93	119.65	111.77

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Mol	Chain	Res	Type	Atoms	Z	Observed(^{o})	Ideal(^{o})
22	A	830	CLA	C2D-C1D-ND	2.93	112.26	110.10
22	1	307	CLA	C2D-C1D-ND	2.93	112.26	110.10
22	Z	314	CLA	CHD-C1D-ND	-2.93	121.76	124.45
22	B	842	CLA	CMA-C3A-C4A	2.93	119.65	111.77
22	B	823	CLA	C2C-C1C-NC	2.93	112.72	109.97
22	8	306	CLA	C2C-C1C-NC	2.93	112.72	109.97
22	9	309	CLA	C1C-C2C-C3C	-2.93	103.88	106.96
22	5	307	CLA	CMA-C3A-C4A	2.93	119.64	111.77
22	A	839	CLA	C2D-C1D-ND	2.93	112.26	110.10
22	B	834	CLA	CHD-C1D-ND	-2.93	121.76	124.45
22	L	202	CLA	CMA-C3A-C4A	2.93	119.64	111.77
22	B	814	CLA	C2C-C1C-NC	2.93	112.72	109.97
22	B	805	CLA	C1D-ND-C4D	-2.93	104.25	106.33
22	Z	313	CLA	CMA-C3A-C4A	2.93	119.64	111.77
22	4	811	CLA	CMA-C3A-C4A	2.93	119.64	111.77
22	B	816	CLA	C2C-C1C-NC	2.93	112.71	109.97
22	B	822	CLA	C2C-C1C-NC	2.93	112.71	109.97
22	A	837	CLA	CMA-C3A-C4A	2.93	119.64	111.77
24	5	304	BCR	C33-C5-C4	2.92	119.23	113.62
22	B	825	CLA	C2C-C1C-NC	2.92	112.71	109.97
36	J	105	C7Z	C18-C5-C4	2.92	119.77	114.36
22	B	808	CLA	C1D-ND-C4D	-2.92	104.26	106.33
24	A	856	BCR	C33-C5-C4	2.92	119.23	113.62
22	B	805	CLA	CHD-C1D-ND	-2.92	121.77	124.45
22	5	322	CLA	CHD-C1D-ND	-2.92	121.77	124.45
22	Z	303	CLA	C2C-C1C-NC	2.92	112.71	109.97
22	A	803	CLA	C1D-ND-C4D	-2.92	104.26	106.33
22	3	312	CLA	CHD-C1D-ND	-2.92	121.77	124.45
22	2	304	CLA	CHD-C1D-ND	-2.92	121.77	124.45
22	A	855	CLA	C2C-C1C-NC	2.92	112.71	109.97
22	A	821	CLA	C2D-C1D-ND	2.92	112.26	110.10
22	5	323	CLA	CMA-C3A-C4A	2.92	119.62	111.77
22	9	311	CLA	CMA-C3A-C4A	2.92	119.62	111.77
22	B	837	CLA	CHD-C1D-ND	-2.92	121.77	124.45
22	A	812	CLA	CMA-C3A-C4A	2.92	119.62	111.77
22	Z	309	CLA	CMA-C3A-C4A	2.92	119.62	111.77
22	1	313	CLA	CMA-C3A-C4A	2.92	119.61	111.77
22	A	809	CLA	C1D-ND-C4D	-2.92	104.26	106.33
22	A	812	CLA	C1D-ND-C4D	-2.92	104.26	106.33
24	B	849	BCR	C38-C26-C27	2.92	119.22	113.62
39	4	816	CHL	C4D-CHA-C1A	2.92	124.80	121.25
22	A	814	CLA	CMA-C3A-C4A	2.92	119.61	111.77

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Mol	Chain	Res	Type	Atoms	Z	Observed(^{o})	Ideal(^{o})
22	7	309	CLA	C1D-ND-C4D	-2.92	104.26	106.33
39	1	312	CHL	C3C-C4C-NC	-2.91	107.30	110.57
22	Z	307	CLA	C2D-C1D-ND	2.91	112.25	110.10
22	7	317	CLA	C2D-C1D-ND	2.91	112.25	110.10
22	A	810	CLA	C2C-C1C-NC	2.91	112.70	109.97
22	5	315	CLA	C2C-C1C-NC	2.91	112.70	109.97
22	B	811	CLA	CHD-C1D-ND	-2.91	121.78	124.45
22	7	310	CLA	CHD-C1D-ND	-2.91	121.78	124.45
22	B	839	CLA	C2D-C1D-ND	2.91	112.25	110.10
39	1	312	CHL	C1B-CHB-C4A	-2.91	124.35	130.12
22	A	816	CLA	C2D-C1D-ND	2.91	112.25	110.10
22	1	311	CLA	C1C-C2C-C3C	-2.91	103.89	106.96
22	A	803	CLA	CMA-C3A-C4A	2.91	119.60	111.77
22	4	808	CLA	CMA-C3A-C4A	2.91	119.60	111.77
22	3	313	CLA	C1C-C2C-C3C	-2.91	103.90	106.96
22	4	807	CLA	CMA-C3A-C4A	2.91	119.59	111.77
24	5	305	BCR	C23-C24-C25	-2.91	119.03	127.20
22	A	813	CLA	C2D-C1D-ND	2.91	112.25	110.10
22	3	322	CLA	C2D-C1D-ND	2.91	112.25	110.10
39	3	317	CHL	C1-C2-C3	-2.91	121.01	126.04
22	B	838	CLA	CMA-C3A-C4A	2.91	119.59	111.77
22	1	310	CLA	CMA-C3A-C4A	2.91	119.59	111.77
22	5	310	CLA	C2D-C1D-ND	2.91	112.25	110.10
22	A	835	CLA	C2C-C1C-NC	2.91	112.70	109.97
22	B	814	CLA	C1D-ND-C4D	-2.91	104.27	106.33
22	A	803	CLA	C2C-C1C-NC	2.91	112.69	109.97
22	Z	308	CLA	C2C-C1C-NC	2.91	112.69	109.97
22	8	316	CLA	C2D-C1D-ND	2.91	112.25	110.10
22	3	312	CLA	CMA-C3A-C4A	2.91	119.58	111.77
22	B	835	CLA	C2C-C1C-NC	2.91	112.69	109.97
22	F	302	CLA	CHD-C1D-ND	-2.91	121.78	124.45
22	B	821	CLA	C2D-C1D-ND	2.91	112.25	110.10
22	B	826	CLA	C2D-C1D-ND	2.91	112.25	110.10
22	B	815	CLA	C2D-C1D-ND	2.90	112.24	110.10
22	1	315	CLA	C2D-C1D-ND	2.90	112.24	110.10
22	8	306	CLA	C2D-C1D-ND	2.90	112.24	110.10
22	B	829	CLA	CMA-C3A-C4A	2.90	119.58	111.77
22	A	805	CLA	C2C-C1C-NC	2.90	112.69	109.97
22	6	313	CLA	CMA-C3A-C4A	2.90	119.58	111.77
22	Z	314	CLA	C1D-ND-C4D	-2.90	104.27	106.33
22	B	814	CLA	C2D-C1D-ND	2.90	112.24	110.10
22	2	301	CLA	C1D-ND-C4D	-2.90	104.27	106.33

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Mol	Chain	Res	Type	Atoms	Z	Observed(^{o})	Ideal(^{o})
22	7	307	CLA	CMA-C3A-C4A	2.90	119.57	111.77
22	A	813	CLA	CMA-C3A-C4A	2.90	119.57	111.77
24	A	848	BCR	C34-C9-C10	-2.90	118.86	122.92
22	3	308	CLA	C2D-C1D-ND	2.90	112.24	110.10
24	J	104	BCR	C33-C5-C4	2.90	119.18	113.62
22	1	314	CLA	C2D-C1D-ND	2.90	112.24	110.10
22	A	832	CLA	CMA-C3A-C4A	2.90	119.56	111.77
22	3	314	CLA	CMA-C3A-C4A	2.90	119.56	111.77
22	8	316	CLA	CMA-C3A-C4A	2.90	119.56	111.77
22	9	311	CLA	CHD-C1D-ND	-2.90	121.79	124.45
39	5	317	CHL	C3C-C4C-NC	-2.90	107.32	110.57
22	A	833	CLA	C2C-C1C-NC	2.89	112.68	109.97
22	3	315	CLA	CMA-C3A-C4A	2.89	119.55	111.77
22	7	307	CLA	C2D-C1D-ND	2.89	112.24	110.10
22	Z	306	CLA	C1D-ND-C4D	-2.89	104.28	106.33
22	7	309	CLA	CMA-C3A-C4A	2.89	119.55	111.77
22	A	837	CLA	C1D-ND-C4D	-2.89	104.28	106.33
22	Z	307	CLA	C1D-ND-C4D	-2.89	104.28	106.33
22	A	842	CLA	C2D-C1D-ND	2.89	112.23	110.10
22	B	838	CLA	C2D-C1D-ND	2.89	112.23	110.10
22	6	314	CLA	C2D-C1D-ND	2.89	112.23	110.10
22	B	832	CLA	C1D-ND-C4D	-2.89	104.28	106.33
22	7	307	CLA	C1D-ND-C4D	-2.89	104.28	106.33
22	A	812	CLA	C2C-C1C-NC	2.89	112.68	109.97
22	B	806	CLA	C2D-C1D-ND	2.89	112.23	110.10
22	B	842	CLA	C2D-C1D-ND	2.89	112.23	110.10
22	B	828	CLA	CMA-C3A-C4A	2.89	119.54	111.77
22	B	807	CLA	C1D-ND-C4D	-2.89	104.28	106.33
22	6	301	CLA	C1D-ND-C4D	-2.89	104.28	106.33
22	A	841	CLA	C2C-C1C-NC	2.89	112.68	109.97
22	5	314	CLA	CHD-C1D-ND	-2.89	121.80	124.45
22	B	831	CLA	CMA-C3A-C4A	2.89	119.53	111.77
24	F	303	BCR	C8-C7-C6	-2.89	119.09	127.20
22	6	311	CLA	C2D-C1D-ND	2.89	112.23	110.10
22	Z	306	CLA	CMA-C3A-C4A	2.89	119.53	111.77
24	J	104	BCR	C38-C26-C27	2.89	119.16	113.62
39	Z	312	CHL	C3C-C4C-NC	-2.89	107.33	110.57
22	8	311	CLA	CHD-C1D-ND	-2.89	121.80	124.45
24	3	304	BCR	C29-C30-C25	2.89	114.92	110.48
22	Z	303	CLA	CMA-C3A-C4A	2.89	119.53	111.77
22	A	811	CLA	C1D-ND-C4D	-2.89	104.28	106.33
22	7	310	CLA	C1D-ND-C4D	-2.89	104.28	106.33

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Mol	Chain	Res	Type	Atoms	Z	Observed(^{o})	Ideal(^{o})
24	B	845	BCR	C23-C24-C25	-2.89	119.10	127.20
22	B	828	CLA	C2C-C1C-NC	2.89	112.67	109.97
39	6	315	CHL	C1-O2A-CGA	2.88	124.01	116.44
22	A	808	CLA	CMA-C3A-C4A	2.88	119.53	111.77
22	7	314	CLA	C1D-ND-C4D	-2.88	104.29	106.33
22	B	838	CLA	C1D-ND-C4D	-2.88	104.29	106.33
22	6	310	CLA	C1D-ND-C4D	-2.88	104.29	106.33
22	A	834	CLA	CMA-C3A-C4A	2.88	119.52	111.77
22	B	819	CLA	CMA-C3A-C4A	2.88	119.52	111.77
22	5	318	CLA	C2C-C1C-NC	2.88	112.67	109.97
22	5	323	CLA	C1D-ND-C4D	-2.88	104.29	106.33
22	Z	305	CLA	CMA-C3A-C4A	2.88	119.52	111.77
22	B	812	CLA	C2D-C1D-ND	2.88	112.23	110.10
22	F	305	CLA	C2D-C1D-ND	2.88	112.23	110.10
22	A	804	CLA	CHD-C1D-ND	-2.88	121.81	124.45
22	J	103	CLA	C1D-ND-C4D	-2.88	104.29	106.33
22	L	202	CLA	C1D-ND-C4D	-2.88	104.29	106.33
22	B	830	CLA	C1D-ND-C4D	-2.88	104.29	106.33
22	A	842	CLA	CMA-C3A-C4A	2.88	119.51	111.77
22	Z	314	CLA	CMA-C3A-C4A	2.88	119.51	111.77
38	Z	301	LUT	C11-C12-C13	-2.88	118.33	126.42
22	A	822	CLA	CHD-C1D-ND	-2.88	121.81	124.45
24	F	303	BCR	C33-C5-C4	2.88	119.14	113.62
39	8	312	CHL	C3C-C4C-NC	-2.88	107.34	110.57
22	L	203	CLA	CMA-C3A-C4A	2.88	119.50	111.77
22	Z	303	CLA	C1D-ND-C4D	-2.88	104.29	106.33
38	3	302	LUT	C18-C5-C4	2.88	119.68	114.36
22	B	808	CLA	C2C-C1C-NC	2.88	112.67	109.97
22	5	319	CLA	C1C-C2C-C3C	-2.88	103.93	106.96
22	A	811	CLA	C2C-C1C-NC	2.87	112.67	109.97
38	1	303	LUT	C18-C5-C4	2.87	119.68	114.36
24	A	856	BCR	C7-C8-C9	-2.87	121.89	126.23
22	5	310	CLA	CMA-C3A-C4A	2.87	119.50	111.77
22	1	313	CLA	C1-C2-C3	-2.87	121.07	126.04
22	5	313	CLA	CMA-C3A-C4A	2.87	119.49	111.77
39	Z	312	CHL	CMA-C3A-C4A	2.87	119.49	111.77
22	4	809	CLA	C1-C2-C3	-2.87	121.08	126.04
22	A	807	CLA	CMA-C3A-C4A	2.87	119.49	111.77
22	4	818	CLA	CMA-C3A-C4A	2.87	119.49	111.77
22	9	303	CLA	CBA-CAA-C2A	2.87	122.34	113.86
22	9	307	CLA	CHD-C1D-ND	-2.87	121.81	124.45
39	6	316	CHL	C1B-CHB-C4A	-2.87	124.43	130.12

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Mol	Chain	Res	Type	Atoms	Z	Observed(^{o})	Ideal(^{o})
22	5	313	CLA	C2D-C1D-ND	2.87	112.22	110.10
22	7	308	CLA	CMA-C3A-C4A	2.87	119.49	111.77
22	Z	310	CLA	C1D-ND-C4D	-2.87	104.30	106.33
22	L	201	CLA	CMA-C3A-C4A	2.87	119.49	111.77
22	A	840	CLA	C2D-C1D-ND	2.87	112.22	110.10
22	Z	315	CLA	C2D-C1D-ND	2.87	112.22	110.10
22	4	817	CLA	CMA-C3A-C4A	2.87	119.49	111.77
22	K	203	CLA	CMA-C3A-C4A	2.87	119.48	111.77
22	8	313	CLA	C1D-ND-C4D	-2.87	104.30	106.33
22	A	839	CLA	CMA-C3A-C4A	2.87	119.48	111.77
24	6	306	BCR	C21-C20-C19	-2.87	114.27	123.22
22	6	310	CLA	C2C-C1C-NC	2.87	112.66	109.97
22	5	318	CLA	C1D-ND-C4D	-2.87	104.30	106.33
22	6	301	CLA	CMA-C3A-C4A	2.87	119.48	111.77
22	5	301	CLA	CMA-C3A-C4A	2.87	119.48	111.77
22	B	809	CLA	C2D-C1D-ND	2.87	112.22	110.10
22	1	308	CLA	CMA-C3A-C4A	2.86	119.47	111.77
22	5	322	CLA	CMA-C3A-C4A	2.86	119.47	111.77
39	8	315	CHL	C1B-CHB-C4A	-2.86	124.44	130.12
34	F	308	LMG	C9-C8-C7	-2.86	105.01	111.79
22	B	822	CLA	CHD-C1D-ND	-2.86	121.82	124.45
24	B	847	BCR	C23-C24-C25	-2.86	119.16	127.20
24	K	206	BCR	C11-C12-C13	-2.86	118.37	126.42
24	B	844	BCR	C35-C13-C14	-2.86	118.91	122.92
22	B	820	CLA	C2D-C1D-ND	2.86	112.21	110.10
22	6	322	CLA	C2D-C1D-ND	2.86	112.21	110.10
38	9	302	LUT	C18-C5-C4	2.86	119.66	114.36
22	4	809	CLA	CHD-C1D-ND	-2.86	121.83	124.45
22	B	836	CLA	C2C-C1C-NC	2.86	112.65	109.97
22	A	804	CLA	C2D-C1D-ND	2.86	112.21	110.10
22	A	820	CLA	CMA-C3A-C4A	2.86	119.46	111.77
22	A	833	CLA	C1D-ND-C4D	-2.86	104.30	106.33
22	Z	304	CLA	C1D-ND-C4D	-2.86	104.30	106.33
22	A	835	CLA	C1-C2-C3	-2.86	121.10	126.04
22	5	320	CLA	C1C-C2C-C3C	-2.86	103.95	106.96
22	7	323	CLA	CMA-C3A-C4A	2.86	119.45	111.77
22	1	314	CLA	CMA-C3A-C4A	2.86	119.45	111.77
22	6	312	CLA	CMA-C3A-C4A	2.86	119.45	111.77
22	2	301	CLA	C2D-C1D-ND	2.86	112.21	110.10
22	B	841	CLA	CMA-C3A-C4A	2.86	119.45	111.77
24	4	804	BCR	C29-C30-C25	2.86	114.88	110.48
22	4	815	CLA	C2C-C1C-NC	2.86	112.65	109.97

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Mol	Chain	Res	Type	Atoms	Z	Observed(^{o})	Ideal(^{o})
22	7	308	CLA	CHD-C1D-ND	-2.86	121.83	124.45
24	A	845	BCR	C33-C5-C4	2.86	119.10	113.62
22	9	305	CLA	C1D-ND-C4D	-2.85	104.31	106.33
24	G	203	BCR	C38-C26-C27	2.85	119.10	113.62
22	A	840	CLA	CMA-C3A-C4A	2.85	119.44	111.77
22	8	306	CLA	C1D-ND-C4D	-2.85	104.31	106.33
38	6	303	LUT	C35-C15-C14	-2.85	117.63	123.47
22	Z	305	CLA	CHD-C1D-ND	-2.85	121.83	124.45
22	6	308	CLA	C1D-ND-C4D	-2.85	104.31	106.33
22	6	307	CLA	C2D-C1D-ND	2.85	112.20	110.10
22	4	809	CLA	CMA-C3A-C4A	2.85	119.43	111.77
22	3	318	CLA	C2C-C1C-NC	2.85	112.64	109.97
22	9	303	CLA	C1-C2-C3	-2.85	121.11	126.04
24	I	4001	BCR	C20-C19-C18	-2.85	118.41	126.42
22	B	818	CLA	CMA-C3A-C4A	2.85	119.43	111.77
39	Z	311	CHL	C4D-CHA-C1A	2.85	124.72	121.25
39	5	321	CHL	C4D-CHA-C1A	2.85	124.71	121.25
24	A	844	BCR	C23-C24-C25	-2.85	119.21	127.20
22	5	307	CLA	C2D-C1D-ND	2.85	112.20	110.10
25	4	820	LHG	O8-C23-C24	2.85	120.84	111.91
22	7	305	CLA	C1D-ND-C4D	-2.85	104.31	106.33
22	6	313	CLA	C1D-ND-C4D	-2.85	104.31	106.33
22	1	313	CLA	C1C-C2C-C3C	-2.85	103.96	106.96
22	3	318	CLA	CMA-C3A-C4A	2.85	119.42	111.77
22	Z	309	CLA	CHD-C1D-ND	-2.85	121.84	124.45
39	6	316	CHL	C3C-C4C-NC	-2.85	107.38	110.57
22	1	313	CLA	C1D-ND-C4D	-2.84	104.31	106.33
22	F	301	CLA	CMA-C3A-C4A	2.84	119.41	111.77
22	4	811	CLA	CHD-C1D-ND	-2.84	121.84	124.45
22	7	311	CLA	CMA-C3A-C4A	2.84	119.41	111.77
22	A	838	CLA	C1D-ND-C4D	-2.84	104.32	106.33
22	7	324	CLA	CMA-C3A-C4A	2.84	119.41	111.77
22	A	806	CLA	CMA-C3A-C4A	2.84	119.41	111.77
22	7	312	CLA	C1D-ND-C4D	-2.84	104.32	106.33
22	B	808	CLA	CMA-C3A-C4A	2.84	119.40	111.77
24	B	844	BCR	C38-C26-C27	2.84	119.07	113.62
22	F	302	CLA	C1D-ND-C4D	-2.84	104.32	106.33
24	L	204	BCR	C15-C16-C17	-2.84	117.66	123.47
22	3	313	CLA	C2C-C1C-NC	2.84	112.63	109.97
38	3	303	LUT	C18-C5-C4	2.84	119.61	114.36
22	B	816	CLA	CMA-C3A-C4A	2.84	119.39	111.77
22	A	836	CLA	C1D-ND-C4D	-2.84	104.32	106.33

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Mol	Chain	Res	Type	Atoms	Z	Observed(^{o})	Ideal(^{o})
22	4	807	CLA	CHD-C1D-ND	-2.83	121.85	124.45
24	3	304	BCR	C38-C26-C27	2.83	119.06	113.62
22	6	319	CLA	CMA-C3A-C4A	2.83	119.39	111.77
39	6	320	CHL	C1B-CHB-C4A	-2.83	124.50	130.12
39	4	816	CHL	CMA-C3A-C4A	2.83	119.39	111.77
22	A	827	CLA	CMA-C3A-C4A	2.83	119.39	111.77
22	B	801	CLA	C2C-C1C-NC	2.83	112.62	109.97
38	9	301	LUT	C18-C5-C4	2.83	119.60	114.36
22	8	309	CLA	CMA-C3A-C4A	2.83	119.38	111.77
22	A	819	CLA	C2D-C1D-ND	2.83	112.19	110.10
38	6	303	LUT	C7-C8-C9	-2.83	121.96	126.23
22	B	826	CLA	C1-C2-C3	-2.83	121.15	126.04
39	6	316	CHL	C1-O2A-CGA	2.83	123.87	116.44
22	3	316	CLA	CMA-C3A-C4A	2.83	119.38	111.77
24	3	306	BCR	C23-C24-C25	-2.83	119.26	127.20
22	6	307	CLA	CHD-C1D-ND	-2.83	121.86	124.45
22	A	807	CLA	C1D-ND-C4D	-2.83	104.33	106.33
22	A	828	CLA	C1D-ND-C4D	-2.83	104.33	106.33
22	7	322	CLA	C2C-C1C-NC	2.83	112.62	109.97
24	G	203	BCR	C23-C24-C25	-2.83	119.26	127.20
24	B	844	BCR	C15-C16-C17	-2.83	117.68	123.47
38	6	304	LUT	C18-C5-C4	2.83	119.59	114.36
22	8	310	CLA	CMA-C3A-C4A	2.83	119.37	111.77
22	F	304	CLA	C1-C2-C3	-2.82	121.16	126.04
22	A	822	CLA	C2D-C1D-ND	2.82	112.19	110.10
22	2	302	CLA	C1D-ND-C4D	-2.82	104.33	106.33
39	6	318	CHL	C1B-CHB-C4A	-2.82	124.53	130.12
22	B	822	CLA	CMA-C3A-C4A	2.82	119.36	111.77
22	F	302	CLA	C1C-C2C-C3C	-2.82	103.99	106.96
22	A	829	CLA	CHD-C1D-ND	-2.82	121.86	124.45
22	Z	304	CLA	CHD-C1D-ND	-2.82	121.86	124.45
22	5	326	CLA	CHD-C1D-ND	-2.82	121.86	124.45
22	B	819	CLA	C1D-ND-C4D	-2.82	104.33	106.33
22	3	308	CLA	C1D-ND-C4D	-2.82	104.33	106.33
22	B	830	CLA	CHD-C1D-ND	-2.82	121.86	124.45
22	6	302	CLA	CHD-C1D-ND	-2.82	121.86	124.45
39	8	312	CHL	C1-C2-C3	-2.82	121.17	126.04
22	B	842	CLA	C1C-C2C-C3C	-2.82	103.99	106.96
22	F	304	CLA	CHD-C1D-ND	-2.82	121.86	124.45
22	9	303	CLA	CMA-C3A-C4A	2.82	119.35	111.77
24	3	304	BCR	C33-C5-C4	2.82	119.03	113.62
22	1	307	CLA	C1D-ND-C4D	-2.82	104.33	106.33

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Mol	Chain	Res	Type	Atoms	Z	Observed(^{o})	Ideal(^{o})
22	5	310	CLA	C1D-ND-C4D	-2.82	104.33	106.33
22	7	312	CLA	CMA-C3A-C4A	2.82	119.34	111.77
22	A	820	CLA	CHD-C1D-ND	-2.82	121.87	124.45
22	6	317	CLA	CMA-C3A-C4A	2.82	119.34	111.77
22	A	810	CLA	C1D-ND-C4D	-2.82	104.33	106.33
22	Z	306	CLA	C1-C2-C3	-2.81	121.17	126.04
22	4	809	CLA	C1D-ND-C4D	-2.81	104.34	106.33
22	4	810	CLA	C1D-ND-C4D	-2.81	104.34	106.33
31	B	852	DGD	C2G-O2G-C1B	-2.81	110.86	117.79
25	4	821	LHG	O8-C23-C24	2.81	120.73	111.91
22	Z	306	CLA	C2C-C1C-NC	2.81	112.61	109.97
22	F	301	CLA	C1D-ND-C4D	-2.81	104.34	106.33
24	I	4001	BCR	C38-C26-C25	-2.81	121.37	124.53
22	A	812	CLA	CHD-C1D-ND	-2.81	121.87	124.45
22	8	308	CLA	C1D-ND-C4D	-2.81	104.34	106.33
24	6	305	BCR	C2-C1-C6	2.81	114.81	110.48
22	F	305	CLA	CMA-C3A-C4A	2.81	119.32	111.77
22	B	816	CLA	C2D-C1D-ND	2.81	112.17	110.10
22	A	811	CLA	CMA-C3A-C4A	2.81	119.32	111.77
39	9	314	CHL	CMA-C3A-C2A	2.81	122.65	116.10
22	A	840	CLA	C1C-C2C-C3C	-2.81	104.00	106.96
24	A	847	BCR	C33-C5-C4	2.81	119.01	113.62
22	5	301	CLA	C1D-ND-C4D	-2.81	104.34	106.33
22	5	307	CLA	C1D-ND-C4D	-2.81	104.34	106.33
38	1	302	LUT	C11-C12-C13	-2.81	118.53	126.42
22	6	302	CLA	CMA-C3A-C4A	2.80	119.31	111.77
22	5	326	CLA	CMA-C3A-C4A	2.80	119.31	111.77
39	Z	311	CHL	CHD-C4C-C3C	2.80	128.96	124.84
38	8	303	LUT	C8-C7-C6	-2.80	119.33	127.20
22	A	823	CLA	C2C-C1C-NC	2.80	112.60	109.97
22	B	813	CLA	C1D-ND-C4D	-2.80	104.34	106.33
22	7	322	CLA	C1D-ND-C4D	-2.80	104.34	106.33
22	1	310	CLA	CHD-C1D-ND	-2.80	121.88	124.45
39	4	813	CHL	C3C-C4C-NC	-2.80	107.43	110.57
22	8	311	CLA	C1D-ND-C4D	-2.80	104.34	106.33
22	3	319	CLA	C1C-C2C-C3C	-2.80	104.01	106.96
22	A	826	CLA	C1D-ND-C4D	-2.80	104.35	106.33
22	4	808	CLA	C1D-ND-C4D	-2.80	104.35	106.33
39	5	316	CHL	C3C-C4C-NC	-2.80	107.43	110.57
22	4	818	CLA	C1D-ND-C4D	-2.80	104.35	106.33
38	9	302	LUT	C8-C7-C6	-2.80	119.35	127.20
22	5	311	CLA	CHD-C1D-ND	-2.79	121.89	124.45

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Mol	Chain	Res	Type	Atoms	Z	Observed(^{o})	Ideal(^{o})
22	2	304	CLA	C1D-ND-C4D	-2.79	104.35	106.33
22	8	305	CLA	C2D-C1D-ND	2.79	112.16	110.10
22	6	311	CLA	C1C-C2C-C3C	-2.79	104.02	106.96
22	6	314	CLA	C1C-C2C-C3C	-2.79	104.02	106.96
39	5	317	CHL	C4D-CHA-C1A	2.79	124.65	121.25
22	A	832	CLA	C1D-ND-C4D	-2.79	104.35	106.33
22	A	806	CLA	C1D-ND-C4D	-2.79	104.35	106.33
22	F	304	CLA	C1D-ND-C4D	-2.79	104.35	106.33
22	9	307	CLA	C2D-C1D-ND	2.79	112.16	110.10
24	3	305	BCR	C38-C26-C27	2.79	118.98	113.62
24	B	844	BCR	C33-C5-C4	2.79	118.98	113.62
22	8	314	CLA	CMA-C3A-C4A	2.79	119.27	111.77
22	A	815	CLA	C1D-ND-C4D	-2.79	104.35	106.33
22	K	204	CLA	C1D-ND-C4D	-2.79	104.35	106.33
39	4	814	CHL	C4D-CHA-C1A	2.79	124.64	121.25
22	8	313	CLA	CMA-C3A-C4A	2.79	119.27	111.77
24	B	848	BCR	C30-C25-C26	-2.79	118.69	122.61
39	6	318	CHL	C3C-C4C-NC	-2.79	107.44	110.57
22	L	203	CLA	C1D-ND-C4D	-2.79	104.36	106.33
22	A	826	CLA	CMA-C3A-C4A	2.79	119.26	111.77
22	6	309	CLA	CHD-C1D-ND	-2.79	121.89	124.45
22	1	309	CLA	O2D-CGD-O1D	-2.79	118.39	123.84
22	4	807	CLA	C1C-C2C-C3C	-2.79	104.03	106.96
22	7	306	CLA	C1C-C2C-C3C	-2.79	104.03	106.96
35	J	102	T7X	C12-C13-C14	-2.79	108.26	113.23
22	7	324	CLA	C1-C2-C3	-2.78	121.23	126.04
22	7	308	CLA	C1D-ND-C4D	-2.78	104.36	106.33
22	3	320	CLA	CHD-C1D-ND	-2.78	121.90	124.45
22	6	309	CLA	C2D-C1D-ND	2.78	112.16	110.10
22	A	808	CLA	C1D-ND-C4D	-2.78	104.36	106.33
22	A	827	CLA	C1D-ND-C4D	-2.78	104.36	106.33
22	A	829	CLA	C1D-ND-C4D	-2.78	104.36	106.33
22	3	319	CLA	C1D-ND-C4D	-2.78	104.36	106.33
22	5	314	CLA	C1D-ND-C4D	-2.78	104.36	106.33
22	3	310	CLA	CHD-C1D-ND	-2.78	121.90	124.45
24	4	804	BCR	C4-C5-C6	-2.78	118.69	122.73
22	9	310	CLA	C1D-ND-C4D	-2.78	104.36	106.33
22	B	832	CLA	CMA-C3A-C4A	2.78	119.25	111.77
39	8	301	CHL	C4D-CHA-C1A	2.78	124.63	121.25
22	B	817	CLA	C1D-ND-C4D	-2.78	104.36	106.33
22	A	808	CLA	C1C-C2C-C3C	-2.78	104.04	106.96
22	A	802	CLA	C1D-ND-C4D	-2.78	104.36	106.33

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Mol	Chain	Res	Type	Atoms	Z	Observed(^{o})	Ideal(^{o})
24	L	204	BCR	C8-C7-C6	-2.78	119.40	127.20
39	7	313	CHL	C4D-CHA-C1A	2.78	124.63	121.25
22	B	817	CLA	CMA-C3A-C4A	2.78	119.23	111.77
22	B	833	CLA	C1D-ND-C4D	-2.78	104.36	106.33
22	A	842	CLA	C1C-C2C-C3C	-2.77	104.04	106.96
21	A	801	CL0	C4-C3-C5	2.77	119.94	115.27
39	Z	311	CHL	C1B-CHB-C4A	-2.77	124.62	130.12
22	9	305	CLA	C1C-C2C-C3C	-2.77	104.04	106.96
25	5	324	LHG	O8-C23-C24	2.77	120.61	111.91
22	1	304	CLA	C1D-ND-C4D	-2.77	104.36	106.33
22	B	805	CLA	CMA-C3A-C4A	2.77	119.23	111.77
22	7	304	CLA	C1C-C2C-C3C	-2.77	104.04	106.96
22	3	314	CLA	CHD-C1D-ND	-2.77	121.91	124.45
22	4	818	CLA	C1-O2A-CGA	2.77	123.72	116.44
22	Z	305	CLA	C1C-C2C-C3C	-2.77	104.04	106.96
22	B	825	CLA	C1D-ND-C4D	-2.77	104.37	106.33
22	7	323	CLA	O2D-CGD-O1D	-2.77	118.42	123.84
22	Z	315	CLA	C1C-C2C-C3C	-2.77	104.04	106.96
22	A	841	CLA	CMA-C3A-C4A	2.77	119.22	111.77
22	B	807	CLA	C2C-C1C-NC	2.77	112.57	109.97
22	A	839	CLA	C1D-ND-C4D	-2.77	104.37	106.33
22	4	815	CLA	C1D-ND-C4D	-2.77	104.37	106.33
22	3	322	CLA	C1-C2-C3	-2.77	121.25	126.04
22	A	828	CLA	CMA-C3A-C4A	2.77	119.21	111.77
22	A	832	CLA	C1-C2-C3	-2.77	121.26	126.04
39	8	312	CHL	C1B-CHB-C4A	-2.77	124.64	130.12
39	2	303	CHL	C1B-CHB-C4A	-2.77	124.64	130.12
22	B	810	CLA	C1C-C2C-C3C	-2.77	104.05	106.96
22	6	308	CLA	CMA-C3A-C4A	2.77	119.21	111.77
22	8	308	CLA	CMA-C3A-C4A	2.77	119.21	111.77
22	7	310	CLA	C1C-C2C-C3C	-2.77	104.05	106.96
38	6	303	LUT	C15-C14-C13	-2.76	123.36	127.31
22	A	820	CLA	C1D-ND-C4D	-2.76	104.37	106.33
22	B	831	CLA	C1D-ND-C4D	-2.76	104.37	106.33
22	6	317	CLA	C1D-ND-C4D	-2.76	104.37	106.33
22	A	816	CLA	CHD-C1D-ND	-2.76	121.91	124.45
22	A	817	CLA	CHD-C1D-ND	-2.76	121.91	124.45
22	B	817	CLA	CHD-C1D-ND	-2.76	121.91	124.45
22	G	201	CLA	CHD-C1D-ND	-2.76	121.91	124.45
38	6	304	LUT	C11-C12-C13	-2.76	118.66	126.42
22	3	316	CLA	C1D-ND-C4D	-2.76	104.37	106.33
39	4	813	CHL	CMA-C3A-C4A	2.76	119.20	111.77

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Mol	Chain	Res	Type	Atoms	Z	Observed(^{o})	Ideal(^{o})
22	6	309	CLA	O2A-CGA-CBA	2.76	120.57	111.91
22	9	306	CLA	CMA-C3A-C4A	2.76	119.19	111.77
22	B	818	CLA	C2C-C1C-NC	2.76	112.56	109.97
24	A	848	BCR	C11-C10-C9	-2.76	123.37	127.31
38	7	302	LUT	C1-C6-C5	-2.76	118.73	122.61
22	B	836	CLA	C1D-ND-C4D	-2.76	104.38	106.33
22	Z	305	CLA	C1D-ND-C4D	-2.76	104.38	106.33
24	3	304	BCR	C23-C24-C25	-2.76	119.45	127.20
22	9	311	CLA	C1D-ND-C4D	-2.76	104.38	106.33
22	4	815	CLA	CHD-C1D-ND	-2.76	121.92	124.45
24	5	304	BCR	C15-C14-C13	-2.76	123.38	127.31
22	A	854	CLA	O2A-CGA-CBA	2.76	120.56	111.91
22	5	323	CLA	C1C-C2C-C3C	-2.76	104.06	106.96
22	5	322	CLA	C1D-ND-C4D	-2.76	104.38	106.33
39	Z	312	CHL	C1B-CHB-C4A	-2.75	124.66	130.12
22	A	829	CLA	C1-C2-C3	-2.75	121.28	126.04
22	6	307	CLA	CMA-C3A-C4A	2.75	119.17	111.77
22	Z	307	CLA	C1C-C2C-C3C	-2.75	104.06	106.96
22	3	311	CLA	C1D-ND-C4D	-2.75	104.38	106.33
38	8	302	LUT	C35-C15-C14	-2.75	117.84	123.47
22	A	814	CLA	C1C-C2C-C3C	-2.75	104.06	106.96
24	5	305	BCR	C8-C9-C10	2.75	123.16	118.94
22	8	307	CLA	C1C-C2C-C3C	-2.75	104.07	106.96
24	A	856	BCR	C37-C22-C21	-2.75	119.07	122.92
22	9	304	CLA	C1D-ND-C4D	-2.75	104.38	106.33
22	6	308	CLA	C1C-C2C-C3C	-2.75	104.07	106.96
24	I	4001	BCR	C15-C14-C13	-2.75	123.39	127.31
24	5	305	BCR	C37-C22-C21	-2.75	119.08	122.92
22	A	835	CLA	C1D-ND-C4D	-2.74	104.39	106.33
22	B	841	CLA	C1D-ND-C4D	-2.74	104.39	106.33
22	4	812	CLA	C1D-ND-C4D	-2.74	104.39	106.33
24	4	804	BCR	C27-C26-C25	-2.74	118.75	122.73
22	A	831	CLA	C1D-ND-C4D	-2.74	104.39	106.33
22	A	855	CLA	C1D-ND-C4D	-2.74	104.39	106.33
24	A	847	BCR	C15-C16-C17	-2.74	117.86	123.47
22	B	834	CLA	CMA-C3A-C4A	2.74	119.14	111.77
22	8	307	CLA	CHD-C1D-ND	-2.74	121.94	124.45
24	A	848	BCR	C23-C24-C25	-2.74	119.51	127.20
24	B	844	BCR	C23-C24-C25	-2.74	119.51	127.20
39	6	315	CHL	C2C-C3C-C4C	2.74	108.44	106.49
22	B	834	CLA	C1C-C2C-C3C	-2.74	104.08	106.96
22	4	811	CLA	C1D-ND-C4D	-2.74	104.39	106.33

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Mol	Chain	Res	Type	Atoms	Z	Observed(^{o})	Ideal(^{o})
24	3	307	BCR	C19-C18-C17	-2.74	114.74	118.94
22	L	202	CLA	C1C-C2C-C3C	-2.74	104.08	106.96
22	5	313	CLA	C1C-C2C-C3C	-2.74	104.08	106.96
22	6	322	CLA	C1C-C2C-C3C	-2.74	104.08	106.96
22	7	316	CLA	C1D-ND-C4D	-2.74	104.39	106.33
22	6	317	CLA	CHD-C1D-ND	-2.74	121.94	124.45
22	8	309	CLA	C1D-ND-C4D	-2.74	104.39	106.33
24	A	856	BCR	C8-C7-C6	-2.73	119.52	127.20
22	1	314	CLA	C1C-C2C-C3C	-2.73	104.08	106.96
22	1	308	CLA	CMD-C2D-C3D	-2.73	121.33	127.61
24	F	303	BCR	C29-C30-C25	2.73	114.69	110.48
22	7	315	CLA	CHD-C1D-ND	-2.73	121.94	124.45
24	8	304	BCR	C11-C12-C13	-2.73	118.74	126.42
24	A	846	BCR	C38-C26-C27	2.73	118.86	113.62
39	4	819	CHL	C1B-CHB-C4A	-2.73	124.71	130.12
23	A	843	PQN	C2M-C2-C3	-2.73	119.95	124.40
22	8	307	CLA	O2D-CGD-O1D	-2.73	118.50	123.84
22	Z	314	CLA	C1C-C2C-C3C	-2.73	104.09	106.96
24	A	844	BCR	C16-C17-C18	-2.73	123.42	127.31
22	B	839	CLA	C1D-ND-C4D	-2.73	104.40	106.33
22	6	302	CLA	C1D-ND-C4D	-2.73	104.40	106.33
39	8	301	CHL	C1B-CHB-C4A	-2.73	124.71	130.12
22	A	838	CLA	C1C-C2C-C3C	-2.73	104.09	106.96
22	A	830	CLA	O2D-CGD-O1D	-2.73	118.51	123.84
22	3	309	CLA	C1D-ND-C4D	-2.73	104.40	106.33
38	3	303	LUT	C15-C35-C34	-2.73	117.89	123.47
22	B	809	CLA	C1C-C2C-C3C	-2.73	104.09	106.96
22	1	305	CLA	CMA-C3A-C4A	2.73	119.10	111.77
22	9	308	CLA	C1C-C2C-C3C	-2.73	104.09	106.96
22	A	854	CLA	C1D-ND-C4D	-2.73	104.40	106.33
22	7	305	CLA	C1C-C2C-C3C	-2.72	104.09	106.96
22	B	809	CLA	C1D-ND-C4D	-2.72	104.40	106.33
22	6	321	CLA	C1D-ND-C4D	-2.72	104.40	106.33
22	A	802	CLA	C1C-C2C-C3C	-2.72	104.09	106.96
22	A	816	CLA	C1C-C2C-C3C	-2.72	104.09	106.96
22	B	811	CLA	C1C-C2C-C3C	-2.72	104.09	106.96
22	B	821	CLA	C1D-ND-C4D	-2.72	104.40	106.33
22	8	316	CLA	C1D-ND-C4D	-2.72	104.40	106.33
32	F	306	RRX	C38-C26-C27	2.72	119.40	114.36
22	7	323	CLA	C1C-C2C-C3C	-2.72	104.09	106.96
22	B	829	CLA	C1D-ND-C4D	-2.72	104.40	106.33
22	8	307	CLA	C1D-ND-C4D	-2.72	104.40	106.33

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Mol	Chain	Res	Type	Atoms	Z	Observed(^{o})	Ideal(^{o})
22	B	839	CLA	C1C-C2C-C3C	-2.72	104.10	106.96
39	6	315	CHL	C1B-CHB-C4A	-2.72	124.73	130.12
22	3	312	CLA	C1C-C2C-C3C	-2.72	104.10	106.96
22	7	312	CLA	C1C-C2C-C3C	-2.72	104.10	106.96
22	B	828	CLA	C1D-ND-C4D	-2.72	104.40	106.33
22	5	315	CLA	CHD-C1D-ND	-2.72	121.95	124.45
22	K	203	CLA	C1C-C2C-C3C	-2.72	104.10	106.96
22	5	315	CLA	C1D-ND-C4D	-2.72	104.40	106.33
22	B	812	CLA	C1C-C2C-C3C	-2.72	104.10	106.96
22	4	812	CLA	C1C-C2C-C3C	-2.72	104.10	106.96
22	9	306	CLA	C1D-ND-C4D	-2.72	104.41	106.33
38	7	301	LUT	C1-C6-C5	-2.72	118.79	122.61
22	4	806	CLA	CHD-C1D-ND	-2.72	121.96	124.45
39	7	313	CHL	C1-O2A-CGA	2.72	123.57	116.44
39	5	316	CHL	C1B-CHB-C4A	-2.72	124.74	130.12
22	B	812	CLA	CMA-C3A-C4A	2.72	119.07	111.77
22	8	305	CLA	C1D-ND-C4D	-2.72	104.41	106.33
22	3	311	CLA	CMA-C3A-C4A	2.71	119.07	111.77
22	F	305	CLA	C1D-ND-C4D	-2.71	104.41	106.33
22	1	315	CLA	C1D-ND-C4D	-2.71	104.41	106.33
22	5	309	CLA	C1D-ND-C4D	-2.71	104.41	106.33
22	A	839	CLA	C1C-C2C-C3C	-2.71	104.10	106.96
22	Z	313	CLA	C1C-C2C-C3C	-2.71	104.10	106.96
22	B	827	CLA	CMA-C3A-C4A	2.71	119.07	111.77
22	5	311	CLA	CMA-C3A-C4A	2.71	119.06	111.77
22	B	840	CLA	C1D-ND-C4D	-2.71	104.41	106.33
25	6	323	LHG	C5-O7-C7	-2.71	111.11	117.79
22	8	313	CLA	C1C-C2C-C3C	-2.71	104.11	106.96
22	4	818	CLA	O2A-CGA-CBA	2.71	120.42	111.91
22	B	806	CLA	C1C-C2C-C3C	-2.71	104.11	106.96
24	F	303	BCR	C4-C5-C6	-2.71	118.80	122.73
22	A	830	CLA	C1D-ND-C4D	-2.71	104.41	106.33
22	B	816	CLA	C1D-ND-C4D	-2.71	104.41	106.33
22	3	310	CLA	C1C-C2C-C3C	-2.71	104.11	106.96
22	B	813	CLA	C1C-C2C-C3C	-2.71	104.11	106.96
22	A	825	CLA	C1D-ND-C4D	-2.71	104.41	106.33
22	3	314	CLA	C1D-ND-C4D	-2.71	104.41	106.33
22	1	304	CLA	C1C-C2C-C3C	-2.71	104.11	106.96
22	9	305	CLA	O2A-CGA-CBA	2.71	120.40	111.91
24	3	305	BCR	C20-C19-C18	-2.71	118.81	126.42
22	3	315	CLA	C1D-ND-C4D	-2.71	104.41	106.33
22	4	808	CLA	C1C-C2C-C3C	-2.71	104.11	106.96

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Mol	Chain	Res	Type	Atoms	Z	Observed(^{o})	Ideal(^{o})
22	6	309	CLA	C1C-C2C-C3C	-2.70	104.11	106.96
24	5	305	BCR	C35-C13-C14	-2.70	119.14	122.92
22	B	831	CLA	C2D-C1D-ND	2.70	112.10	110.10
39	5	316	CHL	CMA-C3A-C4A	2.70	119.04	111.77
22	B	826	CLA	O2A-CGA-CBA	2.70	120.39	111.91
22	6	307	CLA	C1D-ND-C4D	-2.70	104.42	106.33
22	B	821	CLA	CMA-C3A-C4A	2.70	119.04	111.77
22	K	204	CLA	C1C-C2C-C3C	-2.70	104.11	106.96
22	3	320	CLA	C1C-C2C-C3C	-2.70	104.11	106.96
22	6	302	CLA	C1C-C2C-C3C	-2.70	104.11	106.96
24	6	306	BCR	C30-C25-C26	-2.70	118.81	122.61
25	6	323	LHG	O8-C23-C24	2.70	120.38	111.91
22	A	831	CLA	CMA-C3A-C4A	2.70	119.03	111.77
22	7	316	CLA	C1C-C2C-C3C	-2.70	104.12	106.96
22	1	309	CLA	C1C-C2C-C3C	-2.70	104.12	106.96
22	8	314	CLA	C1C-C2C-C3C	-2.70	104.12	106.96
22	B	827	CLA	C1C-C2C-C3C	-2.70	104.12	106.96
38	6	303	LUT	C8-C7-C6	-2.70	119.63	127.20
38	7	302	LUT	C18-C5-C4	2.70	119.35	114.36
22	8	307	CLA	CMA-C3A-C4A	2.70	119.02	111.77
22	8	310	CLA	C1C-C2C-C3C	-2.70	104.12	106.96
22	A	834	CLA	C1D-ND-C4D	-2.70	104.42	106.33
22	4	817	CLA	C1D-ND-C4D	-2.70	104.42	106.33
22	6	319	CLA	C1D-ND-C4D	-2.70	104.42	106.33
24	3	304	BCR	C1-C6-C5	-2.69	118.82	122.61
22	5	309	CLA	C1C-C2C-C3C	-2.69	104.13	106.96
22	8	310	CLA	C1D-ND-C4D	-2.69	104.42	106.33
22	A	836	CLA	C1C-C2C-C3C	-2.69	104.13	106.96
22	5	307	CLA	C1C-C2C-C3C	-2.69	104.13	106.96
22	6	313	CLA	CHD-C1D-ND	-2.69	121.98	124.45
24	5	305	BCR	C27-C26-C25	-2.69	118.82	122.73
24	B	844	BCR	C8-C9-C10	2.69	123.07	118.94
24	3	304	BCR	C21-C20-C19	-2.69	114.82	123.22
22	7	317	CLA	C1D-ND-C4D	-2.69	104.42	106.33
22	7	306	CLA	CHD-C1D-ND	-2.69	121.98	124.45
22	B	805	CLA	C1C-C2C-C3C	-2.69	104.13	106.96
22	5	315	CLA	C1C-C2C-C3C	-2.69	104.13	106.96
22	3	322	CLA	C1D-ND-C4D	-2.69	104.43	106.33
22	7	323	CLA	C1D-ND-C4D	-2.69	104.43	106.33
22	B	821	CLA	C1C-C2C-C3C	-2.69	104.13	106.96
22	6	310	CLA	CMA-C3A-C4A	2.69	118.99	111.77
22	Z	313	CLA	C1D-ND-C4D	-2.69	104.43	106.33

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Mol	Chain	Res	Type	Atoms	Z	Observed(^{o})	Ideal(^{o})
22	9	308	CLA	C1D-ND-C4D	-2.69	104.43	106.33
22	5	311	CLA	C1-C2-C3	-2.69	121.40	126.04
22	B	841	CLA	C1C-C2C-C3C	-2.69	104.13	106.96
24	B	847	BCR	C1-C6-C5	-2.68	118.83	122.61
22	B	826	CLA	C1D-ND-C4D	-2.68	104.43	106.33
22	6	314	CLA	C1D-ND-C4D	-2.68	104.43	106.33
22	F	304	CLA	C1C-C2C-C3C	-2.68	104.14	106.96
39	4	813	CHL	C1B-CHB-C4A	-2.68	124.81	130.12
22	B	810	CLA	C1D-ND-C4D	-2.68	104.43	106.33
22	B	827	CLA	C1D-ND-C4D	-2.68	104.43	106.33
22	A	807	CLA	C1C-C2C-C3C	-2.68	104.14	106.96
22	8	311	CLA	C1C-C2C-C3C	-2.68	104.14	106.96
22	5	311	CLA	C1C-C2C-C3C	-2.68	104.14	106.96
22	5	326	CLA	C1D-ND-C4D	-2.68	104.43	106.33
22	A	832	CLA	C1C-C2C-C3C	-2.68	104.14	106.96
22	9	311	CLA	C1C-C2C-C3C	-2.68	104.14	106.96
22	G	202	CLA	C1D-ND-C4D	-2.68	104.43	106.33
22	B	838	CLA	C1C-C2C-C3C	-2.68	104.14	106.96
22	6	307	CLA	C1C-C2C-C3C	-2.68	104.14	106.96
24	A	845	BCR	C23-C24-C25	-2.68	119.68	127.20
22	1	306	CLA	C1D-ND-C4D	-2.68	104.43	106.33
22	7	324	CLA	C1D-ND-C4D	-2.68	104.43	106.33
24	7	303	BCR	C15-C16-C17	-2.68	117.99	123.47
22	A	817	CLA	C2D-C1D-ND	2.68	112.08	110.10
22	B	840	CLA	C1C-C2C-C3C	-2.68	104.14	106.96
24	B	846	BCR	C11-C10-C9	-2.68	123.49	127.31
38	8	303	LUT	C18-C5-C4	2.67	119.31	114.36
22	8	308	CLA	C1C-C2C-C3C	-2.67	104.14	106.96
22	1	314	CLA	C1D-ND-C4D	-2.67	104.44	106.33
22	5	301	CLA	C1C-C2C-C3C	-2.67	104.15	106.96
36	J	105	C7Z	C21-C26-C25	-2.67	118.85	122.61
22	B	801	CLA	C1D-ND-C4D	-2.67	104.44	106.33
22	4	817	CLA	CAA-C2A-C3A	-2.67	109.87	116.10
22	A	806	CLA	C1C-C2C-C3C	-2.67	104.15	106.96
22	3	316	CLA	C1C-C2C-C3C	-2.67	104.15	106.96
24	6	305	BCR	C1-C6-C5	-2.67	118.86	122.61
22	A	840	CLA	C1D-ND-C4D	-2.67	104.44	106.33
22	A	833	CLA	C1C-C2C-C3C	-2.67	104.15	106.96
22	7	307	CLA	C1C-C2C-C3C	-2.67	104.15	106.96
22	4	811	CLA	O2A-CGA-CBA	2.67	120.27	111.91
22	A	819	CLA	C1D-ND-C4D	-2.66	104.44	106.33
22	5	312	CLA	C1C-C2C-C3C	-2.66	104.16	106.96

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Mol	Chain	Res	Type	Atoms	Z	Observed(^{o})	Ideal(^{o})
25	1	317	LHG	O8-C23-C24	2.66	120.27	111.91
22	9	304	CLA	C1C-C2C-C3C	-2.66	104.16	106.96
22	8	314	CLA	C1D-ND-C4D	-2.66	104.44	106.33
22	B	826	CLA	CMA-C3A-C4A	2.66	118.93	111.77
22	1	316	CLA	C1C-C2C-C3C	-2.66	104.16	106.96
22	5	320	CLA	CMA-C3A-C4A	2.66	118.93	111.77
24	6	306	BCR	C16-C15-C14	-2.66	118.02	123.47
22	B	822	CLA	C1D-ND-C4D	-2.66	104.44	106.33
22	Z	316	CLA	C1C-C2C-C3C	-2.66	104.16	106.96
22	3	309	CLA	CMA-C3A-C4A	2.66	118.92	111.77
22	B	831	CLA	CMD-C2D-C3D	-2.66	121.49	127.61
22	B	820	CLA	C1C-C2C-C3C	-2.66	104.16	106.96
22	1	315	CLA	C1C-C2C-C3C	-2.66	104.16	106.96
22	Z	308	CLA	C1C-C2C-C3C	-2.66	104.16	106.96
38	8	302	LUT	C1-C6-C5	-2.66	118.87	122.61
22	A	818	CLA	C1C-C2C-C3C	-2.66	104.16	106.96
22	4	810	CLA	C1C-C2C-C3C	-2.66	104.16	106.96
22	B	809	CLA	CMA-C3A-C4A	2.66	118.92	111.77
22	G	201	CLA	C1C-C2C-C3C	-2.66	104.16	106.96
22	5	308	CLA	C1D-ND-C4D	-2.66	104.45	106.33
22	6	311	CLA	CMA-C3A-C4A	2.66	118.92	111.77
22	5	309	CLA	O2A-CGA-CBA	2.66	120.25	111.91
24	5	304	BCR	C37-C22-C21	-2.66	119.20	122.92
22	K	202	CLA	C1D-ND-C4D	-2.66	104.45	106.33
22	A	815	CLA	CMA-C3A-C4A	2.66	118.91	111.77
24	4	804	BCR	C38-C26-C27	2.66	118.72	113.62
22	L	201	CLA	C1C-C2C-C3C	-2.65	104.17	106.96
22	4	817	CLA	C1C-C2C-C3C	-2.65	104.17	106.96
22	5	310	CLA	C1C-C2C-C3C	-2.65	104.17	106.96
22	A	821	CLA	C1D-ND-C4D	-2.65	104.45	106.33
39	8	315	CHL	CMA-C3A-C4A	2.65	118.90	111.77
22	J	103	CLA	C1C-C2C-C3C	-2.65	104.17	106.96
22	5	314	CLA	C1C-C2C-C3C	-2.65	104.17	106.96
22	L	203	CLA	C1C-C2C-C3C	-2.65	104.17	106.96
22	6	313	CLA	C1C-C2C-C3C	-2.65	104.17	106.96
22	2	301	CLA	C1C-C2C-C3C	-2.65	104.17	106.96
22	B	835	CLA	CMA-C3A-C4A	2.65	118.90	111.77
39	9	312	CHL	C1B-CHB-C4A	-2.65	124.87	130.12
22	3	322	CLA	C1C-C2C-C3C	-2.65	104.17	106.96
22	7	309	CLA	C1C-C2C-C3C	-2.65	104.17	106.96
22	7	324	CLA	C1C-C2C-C3C	-2.65	104.17	106.96
22	6	312	CLA	C1C-C2C-C3C	-2.65	104.17	106.96

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Mol	Chain	Res	Type	Atoms	Z	Observed(^{o})	Ideal(^{o})
22	A	821	CLA	C1-C2-C3	-2.65	121.46	126.04
22	Z	309	CLA	C1C-C2C-C3C	-2.65	104.17	106.96
22	3	308	CLA	C1C-C2C-C3C	-2.65	104.17	106.96
22	F	304	CLA	O2A-CGA-CBA	2.65	120.22	111.91
24	B	848	BCR	C2-C1-C6	2.65	114.56	110.48
22	A	826	CLA	C1C-C2C-C3C	-2.65	104.17	106.96
22	B	801	CLA	C1C-C2C-C3C	-2.65	104.17	106.96
22	B	831	CLA	C1C-C2C-C3C	-2.65	104.17	106.96
22	4	809	CLA	C1C-C2C-C3C	-2.65	104.17	106.96
39	4	814	CHL	C1-O2A-CGA	2.65	123.39	116.44
24	B	846	BCR	C33-C5-C4	2.65	118.70	113.62
24	A	848	BCR	C36-C18-C17	-2.65	119.22	122.92
22	A	842	CLA	C1D-ND-C4D	-2.65	104.46	106.33
22	B	833	CLA	C1C-C2C-C3C	-2.64	104.18	106.96
22	G	202	CLA	CMA-C3A-C4A	2.64	118.88	111.77
38	4	802	LUT	C18-C5-C4	2.64	119.25	114.36
22	3	314	CLA	C1C-C2C-C3C	-2.64	104.18	106.96
24	3	304	BCR	C30-C25-C26	-2.64	118.89	122.61
39	9	312	CHL	CHB-C4A-NA	2.64	128.16	124.51
22	F	301	CLA	C1C-C2C-C3C	-2.64	104.18	106.96
22	7	308	CLA	C1C-C2C-C3C	-2.64	104.18	106.96
22	5	322	CLA	C1C-C2C-C3C	-2.64	104.18	106.96
22	1	308	CLA	C2C-C1C-NC	2.64	112.45	109.97
22	A	827	CLA	C1C-C2C-C3C	-2.64	104.18	106.96
22	3	318	CLA	C1C-C2C-C3C	-2.64	104.18	106.96
22	B	839	CLA	C6-C5-C3	-2.64	110.30	114.62
22	Z	307	CLA	O2D-CGD-O1D	-2.64	118.68	123.84
38	6	303	LUT	C18-C5-C4	2.64	119.24	114.36
22	B	819	CLA	C1C-C2C-C3C	-2.64	104.18	106.96
22	K	203	CLA	C1D-ND-C4D	-2.64	104.46	106.33
22	Z	304	CLA	C1C-C2C-C3C	-2.64	104.18	106.96
22	B	817	CLA	O2D-CGD-O1D	-2.64	118.68	123.84
39	3	317	CHL	C1B-CHB-C4A	-2.64	124.89	130.12
22	4	806	CLA	C1C-C2C-C3C	-2.64	104.18	106.96
36	J	105	C7Z	C7-C8-C9	-2.64	122.25	126.23
22	1	305	CLA	C1D-ND-C4D	-2.64	104.46	106.33
22	2	304	CLA	C1C-C2C-C3C	-2.64	104.19	106.96
22	6	321	CLA	C1C-C2C-C3C	-2.64	104.19	106.96
22	7	322	CLA	CMA-C3A-C4A	2.63	118.85	111.77
24	A	846	BCR	C8-C7-C6	-2.63	119.81	127.20
22	Z	310	CLA	C1C-C2C-C3C	-2.63	104.19	106.96
38	7	302	LUT	C11-C12-C13	-2.63	119.02	126.42

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Mol	Chain	Res	Type	Atoms	Z	Observed(^{o})	Ideal(^{o})
22	B	824	CLA	C2C-C1C-NC	2.63	112.44	109.97
22	3	315	CLA	CAA-C2A-C3A	-2.63	105.57	112.78
22	1	307	CLA	C1C-C2C-C3C	-2.63	104.19	106.96
22	5	308	CLA	C1C-C2C-C3C	-2.63	104.19	106.96
24	7	303	BCR	C8-C7-C6	-2.63	119.81	127.20
22	1	311	CLA	CHD-C1D-ND	-2.63	122.04	124.45
22	3	311	CLA	C1C-C2C-C3C	-2.63	104.19	106.96
22	9	306	CLA	C1C-C2C-C3C	-2.63	104.19	106.96
22	7	315	CLA	C1C-C2C-C3C	-2.63	104.19	106.96
24	B	848	BCR	C27-C26-C25	-2.63	118.92	122.73
22	7	315	CLA	O2D-CGD-O1D	-2.63	118.70	123.84
22	5	313	CLA	CHD-C1D-ND	-2.63	122.04	124.45
22	6	312	CLA	C1D-ND-C4D	-2.63	104.47	106.33
22	A	835	CLA	C1C-C2C-C3C	-2.63	104.20	106.96
22	1	305	CLA	C1C-C2C-C3C	-2.63	104.20	106.96
22	A	822	CLA	O2D-CGD-O1D	-2.63	118.70	123.84
22	7	317	CLA	CMA-C3A-C4A	2.63	118.83	111.77
38	Z	301	LUT	C18-C5-C6	-2.63	121.58	124.53
22	4	815	CLA	O2D-CGD-O1D	-2.62	118.71	123.84
22	8	306	CLA	CHD-C1D-ND	-2.62	122.04	124.45
24	J	104	BCR	C20-C19-C18	-2.62	119.04	126.42
22	7	311	CLA	C1C-C2C-C3C	-2.62	104.20	106.96
22	6	301	CLA	C1C-C2C-C3C	-2.62	104.20	106.96
22	5	319	CLA	CMA-C3A-C4A	2.62	118.83	111.77
22	B	801	CLA	C1-C2-C3	-2.62	121.50	126.04
39	3	317	CHL	CMA-C3A-C4A	2.62	118.83	111.77
22	1	306	CLA	C1C-C2C-C3C	-2.62	104.20	106.96
38	1	302	LUT	C19-C9-C10	-2.62	119.25	122.92
22	3	319	CLA	CMA-C3A-C4A	2.62	118.82	111.77
22	B	817	CLA	C1C-C2C-C3C	-2.62	104.20	106.96
41	7	319	3PH	O31-C31-C32	2.62	120.13	111.91
22	1	310	CLA	C1D-ND-C4D	-2.62	104.47	106.33
22	B	828	CLA	C1C-C2C-C3C	-2.62	104.20	106.96
25	B	850	LHG	C5-O7-C7	-2.62	111.35	117.79
22	3	315	CLA	C2C-C1C-NC	2.62	112.42	109.97
22	B	813	CLA	O2D-CGD-O1D	-2.62	118.72	123.84
22	5	311	CLA	C1D-ND-C4D	-2.62	104.48	106.33
39	2	303	CHL	CMA-C3A-C4A	2.62	118.81	111.77
22	F	304	CLA	CMA-C3A-C4A	2.62	118.80	111.77
22	9	310	CLA	C1C-C2C-C3C	-2.62	104.21	106.96
22	K	205	CLA	O2D-CGD-O1D	-2.61	118.73	123.84
36	5	306	C7Z	C28-C27-C26	-2.61	119.86	127.20

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Mol	Chain	Res	Type	Atoms	Z	Observed(^{o})	Ideal(^{o})
22	B	835	CLA	C1C-C2C-C3C	-2.61	104.21	106.96
22	9	307	CLA	C1C-C2C-C3C	-2.61	104.21	106.96
22	4	805	CLA	O2A-CGA-CBA	2.61	120.11	111.91
22	Z	303	CLA	C1C-C2C-C3C	-2.61	104.21	106.96
38	5	302	LUT	C30-C31-C32	-2.61	115.07	123.22
22	G	201	CLA	CMA-C3A-C4A	2.61	118.79	111.77
22	A	821	CLA	C1C-C2C-C3C	-2.61	104.21	106.96
22	A	837	CLA	C1C-C2C-C3C	-2.61	104.21	106.96
22	A	841	CLA	C1C-C2C-C3C	-2.61	104.21	106.96
22	B	806	CLA	C1D-ND-C4D	-2.61	104.48	106.33
22	A	834	CLA	C1C-C2C-C3C	-2.61	104.21	106.96
22	1	310	CLA	C1C-C2C-C3C	-2.61	104.21	106.96
22	3	309	CLA	C1C-C2C-C3C	-2.61	104.21	106.96
22	8	309	CLA	C1C-C2C-C3C	-2.61	104.22	106.96
22	6	317	CLA	C1C-C2C-C3C	-2.61	104.22	106.96
32	F	306	RRX	C23-C24-C25	-2.61	119.88	127.20
22	B	820	CLA	C1D-ND-C4D	-2.61	104.48	106.33
22	B	837	CLA	C1C-C2C-C3C	-2.61	104.22	106.96
22	5	320	CLA	C2A-C1A-CHA	2.61	128.42	123.86
22	5	312	CLA	CMA-C3A-C4A	2.61	118.78	111.77
22	B	834	CLA	C1D-ND-C4D	-2.61	104.48	106.33
22	Z	316	CLA	CMA-C3A-C4A	2.61	118.78	111.77
24	B	853	BCR	C30-C25-C24	2.60	123.15	115.78
24	L	204	BCR	C38-C26-C27	2.60	118.62	113.62
22	4	806	CLA	C1D-ND-C4D	-2.60	104.48	106.33
22	A	814	CLA	C1D-ND-C4D	-2.60	104.49	106.33
22	4	807	CLA	C1D-ND-C4D	-2.60	104.49	106.33
24	6	306	BCR	C38-C26-C27	2.60	118.62	113.62
22	4	811	CLA	C1C-C2C-C3C	-2.60	104.22	106.96
38	Z	301	LUT	C19-C9-C10	-2.60	119.28	122.92
22	B	829	CLA	C1C-C2C-C3C	-2.60	104.22	106.96
34	F	308	LMG	O8-C28-C29	2.60	120.07	111.91
22	B	820	CLA	C1-C2-C3	-2.60	121.55	126.04
24	A	856	BCR	C20-C19-C18	-2.60	119.11	126.42
24	3	305	BCR	C7-C8-C9	-2.60	122.31	126.23
22	A	819	CLA	C1C-C2C-C3C	-2.60	104.22	106.96
32	F	306	RRX	C33-C5-C4	2.60	118.61	113.62
25	8	317	LHG	O8-C23-C24	2.60	120.06	111.91
38	1	303	LUT	C22-C23-C24	2.60	114.70	111.74
22	A	822	CLA	C1C-C2C-C3C	-2.60	104.22	106.96
22	B	808	CLA	C1C-C2C-C3C	-2.60	104.22	106.96
22	Z	306	CLA	C1C-C2C-C3C	-2.60	104.22	106.96

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Mol	Chain	Res	Type	Atoms	Z	Observed(^{o})	Ideal(^{o})
22	A	820	CLA	C1C-C2C-C3C	-2.60	104.22	106.96
22	B	815	CLA	C1D-ND-C4D	-2.60	104.49	106.33
22	9	309	CLA	C1D-ND-C4D	-2.60	104.49	106.33
22	A	815	CLA	C1C-C2C-C3C	-2.60	104.23	106.96
22	A	817	CLA	C1C-C2C-C3C	-2.60	104.23	106.96
22	8	316	CLA	C1C-C2C-C3C	-2.60	104.23	106.96
22	G	202	CLA	C1C-C2C-C3C	-2.60	104.23	106.96
25	Z	317	LHG	O8-C23-C24	2.59	120.05	111.91
22	A	804	CLA	C1C-C2C-C3C	-2.59	104.23	106.96
22	7	316	CLA	CMA-C3A-C4A	2.59	118.74	111.77
22	4	815	CLA	C1C-C2C-C3C	-2.59	104.23	106.96
22	K	205	CLA	C2A-C1A-CHA	2.59	128.39	123.86
22	A	804	CLA	C1D-ND-C4D	-2.59	104.49	106.33
22	F	305	CLA	C1C-C2C-C3C	-2.59	104.23	106.96
39	4	814	CHL	CMA-C3A-C4A	2.59	118.74	111.77
22	6	322	CLA	O2D-CGD-O1D	-2.59	118.78	123.84
22	A	810	CLA	C1C-C2C-C3C	-2.59	104.23	106.96
38	3	302	LUT	C8-C7-C6	-2.59	119.93	127.20
22	A	813	CLA	C1C-C2C-C3C	-2.59	104.23	106.96
22	B	814	CLA	CMA-C3A-C4A	2.59	118.73	111.77
25	9	315	LHG	O8-C23-C24	2.59	120.03	111.91
22	5	326	CLA	C1C-C2C-C3C	-2.59	104.24	106.96
24	3	305	BCR	C8-C7-C6	-2.59	119.94	127.20
24	4	804	BCR	C15-C16-C17	-2.59	118.18	123.47
22	A	803	CLA	C1C-C2C-C3C	-2.58	104.24	106.96
22	A	831	CLA	C1C-C2C-C3C	-2.58	104.24	106.96
22	2	302	CLA	C1C-C2C-C3C	-2.58	104.24	106.96
25	A	850	LHG	O8-C23-C24	2.58	120.01	111.91
22	1	308	CLA	O2D-CGD-O1D	-2.58	118.79	123.84
22	5	318	CLA	C1-C2-C3	-2.58	121.58	126.04
22	B	814	CLA	O2D-CGD-O1D	-2.58	118.79	123.84
22	B	830	CLA	C1C-C2C-C3C	-2.58	104.24	106.96
24	7	303	BCR	C29-C30-C25	2.58	114.45	110.48
22	G	201	CLA	CHA-C4D-ND	2.58	137.90	132.50
22	1	316	CLA	CHA-C4D-ND	2.58	137.89	132.50
38	5	302	LUT	C8-C7-C6	-2.58	119.96	127.20
36	5	306	C7Z	C38-C25-C24	2.58	119.13	114.36
22	5	312	CLA	C1D-ND-C4D	-2.58	104.50	106.33
22	A	829	CLA	C1C-C2C-C3C	-2.58	104.25	106.96
31	B	852	DGD	O1G-C1A-C2A	2.58	119.99	111.91
22	A	825	CLA	C1C-C2C-C3C	-2.58	104.25	106.96
22	9	304	CLA	CHA-C4D-ND	2.57	137.88	132.50

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Mol	Chain	Res	Type	Atoms	Z	Observed(^{o})	Ideal(^{o})
22	A	854	CLA	CMA-C3A-C4A	2.57	118.69	111.77
22	7	306	CLA	C1D-ND-C4D	-2.57	104.51	106.33
22	6	307	CLA	C1-C2-C3	-2.57	121.59	126.04
22	9	313	CLA	C1C-C2C-C3C	-2.57	104.25	106.96
22	A	809	CLA	C1C-C2C-C3C	-2.57	104.25	106.96
24	B	853	BCR	C2-C1-C6	2.57	114.44	110.48
22	B	837	CLA	C1D-ND-C4D	-2.57	104.51	106.33
22	B	825	CLA	C1C-C2C-C3C	-2.57	104.25	106.96
22	B	826	CLA	C1C-C2C-C3C	-2.57	104.25	106.96
24	B	848	BCR	C1-C6-C5	-2.57	118.99	122.61
22	A	824	CLA	C1C-C2C-C3C	-2.57	104.25	106.96
22	9	303	CLA	CMB-C2B-C3B	2.57	129.48	124.68
22	7	314	CLA	C1C-C2C-C3C	-2.57	104.26	106.96
22	4	806	CLA	CHA-C4D-ND	2.57	137.87	132.50
24	5	304	BCR	C7-C8-C9	-2.57	122.36	126.23
24	3	304	BCR	C4-C5-C6	-2.57	119.00	122.73
22	B	816	CLA	C1C-C2C-C3C	-2.57	104.26	106.96
39	5	321	CHL	C1B-CHB-C4A	-2.57	125.03	130.12
22	5	319	CLA	C1D-ND-C4D	-2.57	104.51	106.33
24	3	305	BCR	C33-C5-C4	2.57	118.54	113.62
32	F	306	RRX	C36-C18-C17	-2.56	119.33	122.92
22	A	813	CLA	C1D-ND-C4D	-2.56	104.51	106.33
24	J	104	BCR	C23-C24-C25	-2.56	120.00	127.20
24	I	4001	BCR	C36-C18-C17	-2.56	119.33	122.92
22	A	854	CLA	C1C-C2C-C3C	-2.56	104.26	106.96
25	B	803	LHG	O8-C23-C24	2.56	119.95	111.91
22	6	310	CLA	C1C-C2C-C3C	-2.56	104.26	106.96
22	B	839	CLA	CMA-C3A-C4A	2.56	118.66	111.77
22	7	315	CLA	C1D-ND-C4D	-2.56	104.52	106.33
22	3	308	CLA	O2A-CGA-CBA	2.56	119.94	111.91
22	B	836	CLA	C1C-C2C-C3C	-2.56	104.26	106.96
24	B	848	BCR	C23-C24-C25	-2.56	120.01	127.20
21	A	801	CL0	CMB-C2B-C3B	2.56	129.47	124.68
22	3	314	CLA	C1-C2-C3	-2.56	121.62	126.04
24	B	844	BCR	C15-C14-C13	-2.56	123.66	127.31
24	5	305	BCR	C8-C7-C6	-2.56	120.02	127.20
24	B	847	BCR	C33-C5-C4	2.56	118.53	113.62
22	6	308	CLA	O2D-CGD-O1D	-2.56	118.84	123.84
22	7	324	CLA	O2A-CGA-CBA	2.55	119.92	111.91
22	7	322	CLA	C1C-C2C-C3C	-2.55	104.27	106.96
38	Z	302	LUT	C18-C5-C4	2.55	119.08	114.36
39	8	315	CHL	C2C-C3C-C4C	2.55	108.31	106.49

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Mol	Chain	Res	Type	Atoms	Z	Observed(^{o})	Ideal(^{o})
39	8	315	CHL	C1-O2A-CGA	2.55	123.14	116.44
22	A	812	CLA	O2D-CGD-O1D	-2.55	118.85	123.84
39	6	315	CHL	CMA-C3A-C4A	2.55	118.63	111.77
39	6	320	CHL	C2C-C3C-C4C	2.55	108.31	106.49
24	A	846	BCR	C2-C1-C6	2.55	114.41	110.48
24	G	203	BCR	C20-C19-C18	-2.55	119.25	126.42
22	3	310	CLA	C1D-ND-C4D	-2.55	104.52	106.33
22	6	322	CLA	CMA-C3A-C4A	2.55	118.62	111.77
22	5	318	CLA	C1C-C2C-C3C	-2.55	104.28	106.96
38	1	303	LUT	C11-C12-C13	-2.55	119.26	126.42
38	3	302	LUT	C35-C15-C14	-2.55	118.26	123.47
22	9	303	CLA	O2D-CGD-O1D	-2.55	118.86	123.84
22	K	202	CLA	C1C-C2C-C3C	-2.55	104.28	106.96
22	B	831	CLA	CHA-C4D-ND	2.54	137.82	132.50
22	5	313	CLA	CHA-C4D-ND	2.54	137.82	132.50
22	A	809	CLA	C6-C5-C3	-2.54	106.78	113.45
22	A	835	CLA	O2A-CGA-CBA	2.54	119.89	111.91
39	6	320	CHL	CMA-C3A-C4A	2.54	118.61	111.77
22	A	855	CLA	C1C-C2C-C3C	-2.54	104.28	106.96
22	6	314	CLA	O2D-CGD-O1D	-2.54	118.87	123.84
38	5	303	LUT	C18-C5-C4	2.54	119.06	114.36
22	9	307	CLA	C1D-ND-C4D	-2.54	104.53	106.33
22	B	842	CLA	C1D-ND-C4D	-2.54	104.53	106.33
22	6	309	CLA	CHA-C4D-ND	2.54	137.81	132.50
22	B	807	CLA	C1C-C2C-C3C	-2.54	104.29	106.96
22	5	308	CLA	CHA-C4D-ND	2.54	137.81	132.50
38	1	302	LUT	C21-C26-C27	-2.54	109.49	112.70
39	4	819	CHL	CHC-C1C-NC	2.54	128.05	124.20
22	1	316	CLA	CHA-C1A-NA	-2.54	120.59	126.40
24	G	203	BCR	C30-C25-C26	-2.54	119.04	122.61
22	A	811	CLA	C1C-C2C-C3C	-2.54	104.29	106.96
22	A	841	CLA	O2A-CGA-CBA	2.54	119.87	111.91
22	A	828	CLA	O2A-CGA-CBA	2.54	119.86	111.91
22	1	309	CLA	CMA-C3A-C4A	2.53	118.59	111.77
36	5	306	C7Z	C8-C7-C6	-2.53	120.08	127.20
22	4	809	CLA	CHA-C4D-ND	2.53	137.80	132.50
22	B	832	CLA	C1C-C2C-C3C	-2.53	104.29	106.96
22	7	305	CLA	O2D-CGD-O1D	-2.53	118.89	123.84
38	6	303	LUT	C38-C25-C24	-2.53	118.14	123.56
22	6	311	CLA	C1D-ND-C4D	-2.53	104.54	106.33
22	B	812	CLA	C1D-ND-C4D	-2.53	104.54	106.33
24	K	206	BCR	C15-C16-C17	-2.53	118.29	123.47

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Mol	Chain	Res	Type	Atoms	Z	Observed(^{o})	Ideal(^{o})
38	3	303	LUT	C21-C26-C27	-2.53	109.50	112.70
22	F	304	CLA	CHA-C4D-ND	2.53	137.79	132.50
24	B	849	BCR	C2-C1-C6	2.53	114.37	110.48
22	5	326	CLA	CHA-C4D-ND	2.53	137.79	132.50
22	9	304	CLA	CHA-C1A-NA	-2.53	120.61	126.40
22	6	322	CLA	CHA-C4D-ND	2.53	137.78	132.50
22	3	320	CLA	C1D-ND-C4D	-2.53	104.54	106.33
38	5	303	LUT	C21-C26-C27	-2.53	109.51	112.70
39	4	814	CHL	C1B-CHB-C4A	-2.53	125.11	130.12
22	8	306	CLA	C1C-C2C-C3C	-2.53	104.30	106.96
22	9	309	CLA	CHA-C4D-ND	2.53	137.78	132.50
22	5	308	CLA	O2D-CGD-O1D	-2.53	118.90	123.84
22	Z	315	CLA	C1D-ND-C4D	-2.52	104.54	106.33
22	A	828	CLA	C1C-C2C-C3C	-2.52	104.30	106.96
22	4	805	CLA	C1C-C2C-C3C	-2.52	104.30	106.96
22	1	305	CLA	CHA-C4D-ND	2.52	137.78	132.50
39	4	819	CHL	CMA-C3A-C4A	2.52	118.55	111.77
39	2	303	CHL	C2C-C3C-C4C	2.52	108.29	106.49
22	4	810	CLA	CMA-C3A-C4A	2.52	118.55	111.77
22	1	307	CLA	O2A-CGA-CBA	2.52	119.82	111.91
22	Z	303	CLA	O1D-CGD-CBD	-2.52	119.33	124.48
39	6	316	CHL	CHC-C1C-NC	2.52	128.03	124.20
22	K	205	CLA	CMA-C3A-C2A	2.52	124.00	113.83
24	8	304	BCR	C8-C7-C6	-2.52	120.12	127.20
22	9	306	CLA	O2A-CGA-CBA	2.52	119.82	111.91
22	7	308	CLA	CHA-C4D-ND	2.52	137.77	132.50
39	7	313	CHL	C1B-CHB-C4A	-2.52	125.13	130.12
39	4	814	CHL	C2C-C3C-C4C	2.52	108.28	106.49
22	B	826	CLA	CHA-C4D-ND	2.52	137.77	132.50
22	A	809	CLA	O2A-CGA-CBA	2.52	119.81	111.91
39	7	313	CHL	C2C-C3C-C4C	2.52	108.28	106.49
22	A	834	CLA	O2D-CGD-O1D	-2.52	118.92	123.84
22	A	830	CLA	C1C-C2C-C3C	-2.52	104.31	106.96
22	3	315	CLA	CHA-C4D-ND	2.52	137.76	132.50
24	6	305	BCR	C23-C24-C25	-2.51	120.14	127.20
24	3	307	BCR	C8-C7-C6	-2.51	120.14	127.20
22	7	317	CLA	C1C-C2C-C3C	-2.51	104.32	106.96
38	Z	302	LUT	C11-C12-C13	-2.51	119.36	126.42
38	9	301	LUT	C21-C26-C25	2.51	115.91	111.42
22	7	312	CLA	O2A-CGA-CBA	2.51	119.78	111.91
22	A	812	CLA	C1C-C2C-C3C	-2.51	104.32	106.96
38	3	302	LUT	C1-C6-C5	-2.51	119.08	122.61

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Mol	Chain	Res	Type	Atoms	Z	Observed(^{o})	Ideal(^{o})
22	B	822	CLA	C1C-C2C-C3C	-2.51	104.32	106.96
24	A	845	BCR	C8-C7-C6	-2.51	120.16	127.20
39	9	314	CHL	CMA-C3A-C4A	2.51	118.51	111.77
22	7	307	CLA	CMD-C2D-C3D	-2.51	121.85	127.61
22	5	311	CLA	CHA-C4D-ND	2.51	137.74	132.50
22	8	305	CLA	C1C-C2C-C3C	-2.50	104.32	106.96
22	2	301	CLA	CMD-C2D-C3D	-2.50	121.86	127.61
22	B	813	CLA	CHA-C4D-ND	2.50	137.74	132.50
22	9	309	CLA	CMA-C3A-C4A	2.50	118.50	111.77
22	Z	309	CLA	O2A-CGA-CBA	2.50	119.76	111.91
24	J	104	BCR	C8-C7-C6	-2.50	120.17	127.20
22	9	310	CLA	CHA-C4D-ND	2.50	137.73	132.50
24	A	847	BCR	C8-C7-C6	-2.50	120.18	127.20
22	8	306	CLA	CHA-C4D-ND	2.50	137.73	132.50
24	3	304	BCR	C10-C11-C12	2.50	131.02	123.22
22	A	817	CLA	CHA-C4D-ND	2.50	137.73	132.50
22	7	315	CLA	CHA-C4D-ND	2.50	137.73	132.50
22	B	834	CLA	CHA-C4D-ND	2.50	137.73	132.50
24	B	847	BCR	C35-C13-C14	-2.50	119.42	122.92
22	9	304	CLA	O2D-CGD-O1D	-2.50	118.95	123.84
24	3	307	BCR	C34-C9-C10	-2.50	119.42	122.92
39	7	313	CHL	CMA-C3A-C4A	2.50	118.49	111.77
22	1	310	CLA	CHA-C4D-ND	2.50	137.72	132.50
22	5	320	CLA	CHA-C1A-NA	-2.50	120.68	126.40
22	A	807	CLA	CHA-C4D-ND	2.50	137.72	132.50
22	A	816	CLA	CHA-C4D-ND	2.49	137.72	132.50
39	5	317	CHL	C1B-CHB-C4A	-2.49	125.18	130.12
22	3	320	CLA	CHA-C4D-ND	2.49	137.72	132.50
22	B	835	CLA	CMD-C2D-C3D	-2.49	121.88	127.61
22	9	307	CLA	O2D-CGD-O1D	-2.49	118.97	123.84
22	8	309	CLA	CHA-C4D-ND	2.49	137.71	132.50
22	6	312	CLA	CHA-C4D-ND	2.49	137.71	132.50
22	A	805	CLA	C1C-C2C-C3C	-2.49	104.34	106.96
22	4	811	CLA	CHA-C4D-ND	2.49	137.71	132.50
24	L	204	BCR	C23-C24-C25	-2.49	120.21	127.20
22	9	304	CLA	C3D-C2D-C1D	-2.49	102.44	105.83
22	6	307	CLA	O2A-CGA-CBA	2.49	119.72	111.91
22	A	823	CLA	C1C-C2C-C3C	-2.49	104.34	106.96
22	6	319	CLA	C1C-C2C-C3C	-2.49	104.34	106.96
22	6	322	CLA	C1D-ND-C4D	-2.49	104.57	106.33
22	B	823	CLA	C1C-C2C-C3C	-2.49	104.34	106.96
22	A	811	CLA	O2A-CGA-CBA	2.49	119.71	111.91

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Mol	Chain	Res	Type	Atoms	Z	Observed(^{o})	Ideal(^{o})
24	F	303	BCR	C23-C24-C25	-2.49	120.22	127.20
22	B	828	CLA	CHA-C4D-ND	2.49	137.70	132.50
24	B	848	BCR	C4-C5-C6	-2.49	119.12	122.73
22	3	313	CLA	CMA-C3A-C4A	2.49	118.45	111.77
22	A	812	CLA	CHA-C4D-ND	2.48	137.69	132.50
22	K	202	CLA	CHA-C4D-ND	2.48	137.69	132.50
24	A	856	BCR	C15-C16-C17	-2.48	118.39	123.47
22	2	302	CLA	O2D-CGD-O1D	-2.48	118.98	123.84
22	Z	306	CLA	CHA-C4D-ND	2.48	137.69	132.50
22	6	307	CLA	CHA-C4D-ND	2.48	137.69	132.50
32	F	306	RRX	C2-C1-C6	2.48	114.30	110.48
22	A	804	CLA	CMA-C3A-C4A	2.48	118.44	111.77
22	7	323	CLA	CHA-C4D-ND	2.48	137.69	132.50
22	B	810	CLA	CHA-C4D-ND	2.48	137.69	132.50
22	1	309	CLA	CHA-C4D-ND	2.48	137.69	132.50
39	9	314	CHL	C1B-CHB-C4A	-2.48	125.20	130.12
22	K	205	CLA	CHA-C4D-ND	2.48	137.69	132.50
22	1	313	CLA	CHA-C4D-ND	2.48	137.68	132.50
22	B	838	CLA	CMD-C2D-C3D	-2.48	121.91	127.61
39	4	819	CHL	C2C-C3C-C4C	2.48	108.25	106.49
22	B	835	CLA	O2D-CGD-O1D	-2.48	119.00	123.84
22	A	804	CLA	CHA-C4D-ND	2.48	137.68	132.50
24	7	303	BCR	C30-C25-C26	-2.48	119.13	122.61
22	1	315	CLA	CHA-C4D-ND	2.48	137.68	132.50
22	Z	313	CLA	CHA-C4D-ND	2.48	137.68	132.50
22	K	203	CLA	CHA-C4D-ND	2.48	137.68	132.50
22	7	324	CLA	CHA-C4D-ND	2.48	137.68	132.50
22	B	814	CLA	C1C-C2C-C3C	-2.48	104.35	106.96
24	3	306	BCR	C35-C13-C14	-2.48	119.46	122.92
22	B	831	CLA	O2D-CGD-O1D	-2.47	119.00	123.84
22	5	312	CLA	CHA-C4D-ND	2.47	137.67	132.50
22	B	840	CLA	O2D-CGD-O1D	-2.47	119.00	123.84
22	B	805	CLA	CHA-C4D-ND	2.47	137.67	132.50
22	B	815	CLA	C1C-C2C-C3C	-2.47	104.36	106.96
38	9	301	LUT	C35-C15-C14	-2.47	118.41	123.47
24	4	804	BCR	C8-C7-C6	-2.47	120.26	127.20
22	A	816	CLA	O2A-CGA-CBA	2.47	119.66	111.91
22	5	309	CLA	CHA-C4D-ND	2.47	137.66	132.50
21	A	801	CL0	O2D-CGD-O1D	-2.47	119.01	123.84
39	8	312	CHL	CHC-C1C-NC	2.47	127.95	124.20
24	F	303	BCR	C20-C19-C18	-2.47	119.48	126.42
22	3	309	CLA	CHA-C4D-ND	2.47	137.66	132.50

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Mol	Chain	Res	Type	Atoms	Z	Observed(^{o})	Ideal(^{o})
24	A	845	BCR	C36-C18-C17	-2.47	119.47	122.92
24	B	853	BCR	C20-C19-C18	-2.47	119.49	126.42
22	B	806	CLA	CHA-C4D-ND	2.47	137.66	132.50
22	4	805	CLA	CHA-C4D-ND	2.46	137.65	132.50
22	B	825	CLA	O2A-CGA-CBA	2.46	119.64	111.91
22	B	834	CLA	O2D-CGD-O1D	-2.46	119.02	123.84
39	4	816	CHL	C1B-CHB-C4A	-2.46	125.24	130.12
22	4	818	CLA	C1C-C2C-C3C	-2.46	104.37	106.96
22	G	201	CLA	C1D-ND-C4D	-2.46	104.58	106.33
22	4	805	CLA	O2D-CGD-O1D	-2.46	119.02	123.84
22	9	307	CLA	CHA-C4D-ND	2.46	137.65	132.50
22	A	820	CLA	CHA-C4D-ND	2.46	137.65	132.50
22	5	311	CLA	O2D-CGD-O1D	-2.46	119.03	123.84
22	9	305	CLA	CAA-CBA-CGA	-2.46	106.06	113.25
22	A	826	CLA	CHA-C4D-ND	2.46	137.65	132.50
22	5	307	CLA	CMD-C2D-C3D	-2.46	121.95	127.61
22	A	827	CLA	CHA-C4D-ND	2.46	137.65	132.50
22	3	312	CLA	CHA-C4D-ND	2.46	137.65	132.50
22	5	307	CLA	CHA-C4D-ND	2.46	137.65	132.50
22	B	826	CLA	O2D-CGD-O1D	-2.46	119.03	123.84
22	A	802	CLA	CHA-C4D-ND	2.46	137.64	132.50
22	A	824	CLA	C1-C2-C3	-2.46	121.79	126.04
22	A	833	CLA	O2D-CGD-O1D	-2.46	119.03	123.84
22	A	806	CLA	CHA-C4D-ND	2.46	137.64	132.50
22	6	309	CLA	CHA-C1A-NA	-2.46	120.77	126.40
22	B	816	CLA	CHA-C4D-ND	2.46	137.64	132.50
22	A	826	CLA	O2D-CGD-O1D	-2.46	119.03	123.84
22	A	818	CLA	CHA-C4D-ND	2.46	137.64	132.50
22	B	837	CLA	CHA-C4D-ND	2.46	137.64	132.50
22	4	806	CLA	O2D-CGD-O1D	-2.46	119.03	123.84
22	3	322	CLA	CHA-C4D-ND	2.46	137.64	132.50
22	F	302	CLA	CHA-C4D-ND	2.46	137.64	132.50
22	5	310	CLA	CHA-C4D-ND	2.46	137.64	132.50
22	6	312	CLA	O2A-CGA-CBA	2.46	119.61	111.91
22	A	825	CLA	CHA-C4D-ND	2.46	137.64	132.50
22	3	310	CLA	O2A-CGA-CBA	2.46	119.61	111.91
22	9	305	CLA	CHA-C4D-ND	2.46	137.63	132.50
22	3	312	CLA	O2D-CGD-O1D	-2.46	119.04	123.84
22	Z	313	CLA	C1-C2-C3	-2.45	121.80	126.04
22	B	842	CLA	CHA-C4D-ND	2.45	137.63	132.50
22	7	312	CLA	CHA-C4D-ND	2.45	137.63	132.50
22	A	840	CLA	CHA-C4D-ND	2.45	137.63	132.50

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Mol	Chain	Res	Type	Atoms	Z	Observed(^{o})	Ideal(^{o})
22	Z	303	CLA	CHA-C4D-ND	2.45	137.63	132.50
22	3	316	CLA	CHA-C4D-ND	2.45	137.63	132.50
22	A	813	CLA	CHA-C4D-ND	2.45	137.63	132.50
31	3	301	DGD	O1G-C1A-C2A	2.45	119.60	111.91
22	A	837	CLA	CHA-C4D-ND	2.45	137.63	132.50
22	6	309	CLA	C1D-ND-C4D	-2.45	104.59	106.33
24	A	847	BCR	C11-C12-C13	-2.45	119.53	126.42
22	A	804	CLA	O2D-CGD-O1D	-2.45	119.05	123.84
24	K	206	BCR	C8-C7-C6	-2.45	120.32	127.20
22	A	833	CLA	CHA-C4D-ND	2.45	137.63	132.50
22	4	810	CLA	CHA-C4D-ND	2.45	137.63	132.50
22	3	320	CLA	CMA-C3A-C4A	2.45	118.36	111.77
21	A	801	CL0	C3D-C4D-ND	2.45	114.20	110.24
22	3	320	CLA	CHA-C1A-NA	-2.45	120.79	126.40
22	4	807	CLA	CHA-C4D-ND	2.45	137.62	132.50
22	7	306	CLA	CHA-C1A-NA	-2.45	120.79	126.40
22	5	315	CLA	CHA-C4D-ND	2.45	137.62	132.50
22	B	827	CLA	O2D-CGD-O1D	-2.45	119.05	123.84
22	B	816	CLA	CMD-C2D-C3D	-2.45	121.98	127.61
22	9	306	CLA	CHA-C4D-ND	2.45	137.62	132.50
22	A	830	CLA	CHA-C4D-ND	2.45	137.62	132.50
24	A	846	BCR	C23-C24-C25	-2.45	120.33	127.20
41	5	325	3PH	O31-C31-C32	2.45	119.58	111.91
38	5	303	LUT	C11-C12-C13	-2.45	119.55	126.42
25	5	324	LHG	C5-O7-C7	-2.44	111.77	117.79
22	A	839	CLA	CHA-C4D-ND	2.44	137.61	132.50
22	L	201	CLA	CHA-C4D-ND	2.44	137.61	132.50
22	5	308	CLA	CMA-C3A-C4A	2.44	118.33	111.77
22	8	307	CLA	CHA-C4D-ND	2.44	137.60	132.50
22	B	827	CLA	CHA-C4D-ND	2.44	137.60	132.50
22	F	305	CLA	CHA-C4D-ND	2.44	137.60	132.50
24	5	305	BCR	C38-C26-C27	2.44	118.30	113.62
22	B	808	CLA	O2A-CGA-CBA	2.44	119.56	111.91
22	A	836	CLA	CHA-C4D-ND	2.44	137.60	132.50
22	3	308	CLA	CHA-C4D-ND	2.44	137.60	132.50
22	A	803	CLA	O2A-CGA-CBA	2.44	119.56	111.91
24	I	4001	BCR	C8-C7-C6	-2.44	120.35	127.20
22	5	322	CLA	CHA-C4D-ND	2.44	137.60	132.50
22	9	311	CLA	CHA-C4D-ND	2.44	137.60	132.50
22	A	819	CLA	CHA-C4D-ND	2.44	137.60	132.50
22	A	815	CLA	CHA-C4D-ND	2.44	137.60	132.50
22	A	836	CLA	CMA-C3A-C4A	2.44	118.32	111.77

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Mol	Chain	Res	Type	Atoms	Z	Observed(^{o})	Ideal(^{o})
22	1	307	CLA	CHA-C4D-ND	2.44	137.60	132.50
22	3	310	CLA	CHA-C4D-ND	2.44	137.60	132.50
25	A	849	LHG	O8-C23-C24	2.44	119.56	111.91
22	B	819	CLA	CHA-C4D-ND	2.44	137.60	132.50
22	B	820	CLA	CHA-C4D-ND	2.44	137.60	132.50
22	6	301	CLA	CHA-C4D-ND	2.44	137.60	132.50
22	6	314	CLA	CHA-C4D-ND	2.44	137.60	132.50
22	A	854	CLA	O2D-CGD-O1D	-2.44	119.08	123.84
22	7	306	CLA	CHA-C4D-ND	2.44	137.59	132.50
22	A	842	CLA	O2D-CGD-O1D	-2.44	119.08	123.84
22	B	811	CLA	CHA-C4D-ND	2.44	137.59	132.50
22	8	310	CLA	CHA-C4D-ND	2.44	137.59	132.50
22	3	310	CLA	CHA-C1A-NA	-2.44	120.82	126.40
22	B	814	CLA	CHA-C4D-ND	2.44	137.59	132.50
22	F	301	CLA	CHA-C4D-ND	2.43	137.59	132.50
22	9	309	CLA	O2A-CGA-CBA	2.43	119.55	111.91
22	A	814	CLA	CHA-C4D-ND	2.43	137.59	132.50
22	6	310	CLA	O2A-CGA-CBA	2.43	119.55	111.91
24	A	847	BCR	C2-C1-C6	2.43	114.23	110.48
22	A	821	CLA	CHA-C4D-ND	2.43	137.59	132.50
22	B	817	CLA	CHA-C4D-ND	2.43	137.59	132.50
22	7	323	CLA	C1-C2-C3	-2.43	121.84	126.04
22	B	836	CLA	CHA-C4D-ND	2.43	137.59	132.50
22	3	314	CLA	CHA-C4D-ND	2.43	137.59	132.50
22	2	301	CLA	CHA-C4D-ND	2.43	137.59	132.50
22	A	855	CLA	CHA-C4D-ND	2.43	137.59	132.50
22	6	311	CLA	CHA-C4D-ND	2.43	137.58	132.50
22	A	819	CLA	O2D-CGD-O1D	-2.43	119.09	123.84
22	G	202	CLA	CHA-C4D-ND	2.43	137.58	132.50
22	B	833	CLA	CHA-C4D-ND	2.43	137.58	132.50
22	B	840	CLA	CHA-C4D-ND	2.43	137.58	132.50
22	7	311	CLA	CHA-C4D-ND	2.43	137.58	132.50
24	8	304	BCR	C23-C24-C25	-2.43	120.38	127.20
22	B	815	CLA	CHA-C4D-ND	2.43	137.58	132.50
22	2	302	CLA	CHA-C4D-ND	2.43	137.58	132.50
22	L	203	CLA	CHA-C4D-ND	2.43	137.58	132.50
22	1	308	CLA	CHA-C4D-ND	2.43	137.58	132.50
24	K	206	BCR	C23-C24-C25	-2.43	120.39	127.20
22	A	854	CLA	CHA-C4D-ND	2.43	137.57	132.50
22	4	815	CLA	CHA-C4D-ND	2.43	137.57	132.50
22	6	319	CLA	CHA-C4D-ND	2.43	137.57	132.50
22	A	811	CLA	CHA-C4D-ND	2.43	137.57	132.50

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Mol	Chain	Res	Type	Atoms	Z	Observed(^{o})	Ideal(^{o})
22	8	313	CLA	O2D-CGD-O1D	-2.43	119.10	123.84
38	6	304	LUT	C15-C35-C34	-2.43	118.51	123.47
22	5	320	CLA	CHA-C4D-ND	2.43	137.57	132.50
22	B	834	CLA	O2A-CGA-CBA	2.43	119.52	111.91
22	A	803	CLA	O2D-CGD-O1D	-2.42	119.10	123.84
38	5	302	LUT	C18-C5-C4	2.42	118.84	114.36
22	B	838	CLA	CHA-C4D-ND	2.42	137.57	132.50
22	Z	315	CLA	CHA-C4D-ND	2.42	137.57	132.50
24	6	306	BCR	C37-C22-C23	2.42	121.89	118.08
22	6	302	CLA	O2D-CGD-O1D	-2.42	119.10	123.84
22	A	822	CLA	CHA-C4D-ND	2.42	137.57	132.50
22	A	831	CLA	CHA-C4D-ND	2.42	137.57	132.50
22	A	828	CLA	CHA-C4D-ND	2.42	137.56	132.50
22	A	832	CLA	CHA-C4D-ND	2.42	137.56	132.50
22	3	319	CLA	CHA-C4D-ND	2.42	137.56	132.50
22	7	314	CLA	O2A-CGA-CBA	2.42	119.51	111.91
22	B	818	CLA	C1C-C2C-C3C	-2.42	104.41	106.96
22	A	842	CLA	CHA-C4D-ND	2.42	137.56	132.50
22	1	309	CLA	CMD-C2D-C3D	-2.42	122.05	127.61
22	B	841	CLA	O2D-CGD-O1D	-2.42	119.11	123.84
38	7	302	LUT	C19-C9-C10	-2.42	119.53	122.92
22	A	808	CLA	CHA-C4D-ND	2.42	137.56	132.50
22	7	307	CLA	CHA-C4D-ND	2.42	137.56	132.50
22	B	841	CLA	CHA-C4D-ND	2.42	137.56	132.50
22	8	316	CLA	CHA-C4D-ND	2.42	137.56	132.50
22	6	319	CLA	O2D-CGD-O1D	-2.42	119.11	123.84
22	7	316	CLA	CHA-C4D-ND	2.42	137.56	132.50
22	B	801	CLA	CHA-C4D-ND	2.42	137.56	132.50
22	A	822	CLA	O2A-CGA-CBA	2.42	119.49	111.91
39	5	317	CHL	CMA-C3A-C4A	2.42	118.27	111.77
22	7	305	CLA	CHA-C4D-ND	2.42	137.56	132.50
22	8	311	CLA	CHA-C4D-ND	2.42	137.55	132.50
22	8	313	CLA	CHA-C4D-ND	2.42	137.55	132.50
22	B	809	CLA	CHA-C4D-ND	2.42	137.55	132.50
22	Z	307	CLA	CHA-C4D-ND	2.42	137.55	132.50
22	1	311	CLA	CHA-C4D-ND	2.42	137.55	132.50
22	6	302	CLA	CHA-C4D-ND	2.42	137.55	132.50
36	5	306	C7Z	C24-C25-C26	-2.42	115.47	120.85
22	B	820	CLA	O2D-CGD-O1D	-2.42	119.12	123.84
24	6	306	BCR	C34-C9-C10	-2.42	119.54	122.92
22	A	829	CLA	CHA-C4D-ND	2.42	137.55	132.50
22	4	817	CLA	O2D-CGD-O1D	-2.41	119.12	123.84

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Mol	Chain	Res	Type	Atoms	Z	Observed(^{o})	Ideal(^{o})
22	4	817	CLA	CHA-C4D-ND	2.41	137.55	132.50
24	A	856	BCR	C29-C30-C25	2.41	114.20	110.48
22	B	818	CLA	CHA-C4D-ND	2.41	137.55	132.50
22	B	821	CLA	CHA-C4D-ND	2.41	137.55	132.50
22	B	835	CLA	CHA-C4D-ND	2.41	137.55	132.50
22	4	809	CLA	O2A-CGA-CBA	2.41	119.48	111.91
22	A	835	CLA	CMA-C3A-C4A	2.41	118.26	111.77
22	4	805	CLA	C1-C2-C3	-2.41	121.87	126.04
22	Z	308	CLA	CHA-C4D-ND	2.41	137.54	132.50
24	6	306	BCR	C8-C7-C6	-2.41	120.43	127.20
38	1	302	LUT	C38-C25-C24	-2.41	118.40	123.56
32	F	306	RRX	C30-C25-C26	-2.41	119.22	122.61
22	8	305	CLA	O1D-CGD-CBD	-2.41	119.55	124.48
22	A	816	CLA	C1D-ND-C4D	-2.41	104.62	106.33
22	1	306	CLA	CHA-C4D-ND	2.41	137.54	132.50
22	3	313	CLA	CHA-C4D-ND	2.41	137.54	132.50
22	9	313	CLA	CHA-C4D-ND	2.41	137.54	132.50
22	6	317	CLA	CHA-C4D-ND	2.41	137.54	132.50
22	6	321	CLA	CHA-C4D-ND	2.41	137.54	132.50
22	6	307	CLA	O2D-CGD-O1D	-2.41	119.12	123.84
22	1	306	CLA	O2A-CGA-CBA	2.41	119.47	111.91
22	1	304	CLA	CHA-C4D-ND	2.41	137.54	132.50
22	5	323	CLA	CHA-C4D-ND	2.41	137.54	132.50
22	3	316	CLA	O2D-CGD-O1D	-2.41	119.13	123.84
22	B	839	CLA	CHA-C4D-ND	2.41	137.54	132.50
22	B	826	CLA	CMD-C2D-C3D	-2.41	122.07	127.61
22	4	812	CLA	O2D-CGD-O1D	-2.41	119.13	123.84
22	A	838	CLA	CHA-C4D-ND	2.41	137.54	132.50
22	B	808	CLA	CHA-C4D-ND	2.41	137.54	132.50
22	9	308	CLA	CHA-C4D-ND	2.41	137.53	132.50
22	3	312	CLA	O2A-CGA-CBA	2.41	119.46	111.91
22	Z	309	CLA	O2D-CGD-O1D	-2.41	119.13	123.84
22	A	810	CLA	CHA-C4D-ND	2.41	137.53	132.50
22	3	311	CLA	O2A-CGA-CBA	2.41	119.46	111.91
33	4	801	LMT	C1B-O1B-C4'	-2.41	112.01	117.96
24	A	847	BCR	C27-C26-C25	-2.41	119.24	122.73
22	3	311	CLA	CHA-C4D-ND	2.41	137.53	132.50
24	7	303	BCR	C1-C6-C7	2.41	122.58	115.78
22	3	314	CLA	C1-O2A-CGA	2.40	122.75	116.44
22	B	822	CLA	CHA-C4D-ND	2.40	137.53	132.50
39	5	317	CHL	C2C-C3C-C4C	2.40	108.20	106.49
22	A	820	CLA	C6-C5-C3	-2.40	107.15	113.45

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Mol	Chain	Res	Type	Atoms	Z	Observed(^{o})	Ideal(^{o})
22	1	304	CLA	O2A-CGA-CBA	2.40	119.45	111.91
22	4	818	CLA	CHA-C4D-ND	2.40	137.53	132.50
22	7	324	CLA	O2D-CGD-O1D	-2.40	119.14	123.84
25	A	849	LHG	C5-O7-C7	-2.40	111.87	117.79
22	8	308	CLA	CHA-C4D-ND	2.40	137.53	132.50
22	4	812	CLA	CHA-C4D-ND	2.40	137.53	132.50
38	6	303	LUT	C11-C12-C13	-2.40	119.67	126.42
22	5	318	CLA	CHA-C4D-ND	2.40	137.53	132.50
22	B	812	CLA	CHA-C4D-ND	2.40	137.53	132.50
22	7	322	CLA	CHA-C4D-ND	2.40	137.53	132.50
22	5	311	CLA	O2A-CGA-CBA	2.40	119.45	111.91
24	B	848	BCR	C1-C6-C7	2.40	122.57	115.78
38	4	803	LUT	C11-C12-C13	-2.40	119.67	126.42
39	9	312	CHL	C4D-CHA-C1A	2.40	124.17	121.25
22	A	816	CLA	O2D-CGD-O1D	-2.40	119.14	123.84
22	B	837	CLA	O2D-CGD-O1D	-2.40	119.14	123.84
22	5	314	CLA	CHA-C4D-ND	2.40	137.52	132.50
22	Z	306	CLA	O2A-CGA-CBA	2.40	119.44	111.91
22	8	314	CLA	CHA-C4D-ND	2.40	137.52	132.50
24	B	846	BCR	C10-C11-C12	-2.40	115.73	123.22
22	7	317	CLA	CHA-C4D-ND	2.40	137.52	132.50
22	8	311	CLA	O2D-CGD-O1D	-2.40	119.15	123.84
22	7	304	CLA	C1-O2A-CGA	2.40	122.74	116.44
22	B	801	CLA	CMA-C3A-C4A	2.40	118.22	111.77
22	A	823	CLA	CHA-C4D-ND	2.40	137.52	132.50
22	B	824	CLA	CHA-C4D-ND	2.40	137.52	132.50
22	7	316	CLA	O2D-CGD-O1D	-2.40	119.15	123.84
24	3	306	BCR	C8-C7-C6	-2.40	120.47	127.20
39	1	312	CHL	CHC-C1C-NC	2.40	127.84	124.20
22	A	834	CLA	CHA-C4D-ND	2.40	137.51	132.50
22	4	807	CLA	CHA-C1A-NA	-2.40	120.91	126.40
22	B	832	CLA	CHA-C4D-ND	2.40	137.51	132.50
22	B	824	CLA	C1C-C2C-C3C	-2.40	104.44	106.96
22	B	828	CLA	O2D-CGD-O1D	-2.40	119.15	123.84
22	J	103	CLA	CHA-C4D-ND	2.40	137.51	132.50
22	7	314	CLA	CHA-C4D-ND	2.40	137.51	132.50
22	6	310	CLA	CHA-C4D-ND	2.40	137.51	132.50
41	6	324	3PH	O31-C31-C32	2.40	119.43	111.91
22	6	313	CLA	CHA-C4D-ND	2.40	137.51	132.50
22	A	803	CLA	CHA-C4D-ND	2.40	137.51	132.50
22	B	827	CLA	O2A-CGA-CBA	2.40	119.42	111.91
22	B	841	CLA	O2A-CGA-CBA	2.40	119.42	111.91

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Mol	Chain	Res	Type	Atoms	Z	Observed(^{o})	Ideal(^{o})
25	7	318	LHG	C5-O7-C7	-2.39	111.89	117.79
24	3	304	BCR	C7-C6-C5	-2.39	115.66	121.46
22	Z	306	CLA	CMD-C2D-C3D	-2.39	122.11	127.61
23	B	843	PQN	C2M-C2-C3	-2.39	120.50	124.40
22	8	305	CLA	CHA-C4D-ND	2.39	137.50	132.50
22	Z	316	CLA	CHA-C4D-ND	2.39	137.50	132.50
22	B	807	CLA	CHA-C4D-ND	2.39	137.50	132.50
22	5	319	CLA	CHA-C4D-ND	2.39	137.50	132.50
22	9	303	CLA	CMB-C2B-C1B	-2.39	124.79	128.46
35	J	102	T7X	O18-C11-C31	2.39	119.41	111.91
22	A	835	CLA	CHA-C4D-ND	2.39	137.50	132.50
22	5	310	CLA	CMD-C2D-C3D	-2.39	122.11	127.61
22	B	823	CLA	CHA-C4D-ND	2.39	137.50	132.50
22	3	318	CLA	CHA-C4D-ND	2.39	137.50	132.50
22	A	818	CLA	O2A-CGA-CBA	2.39	119.41	111.91
22	Z	307	CLA	O2A-CGA-CBA	2.39	119.41	111.91
24	3	304	BCR	C36-C18-C19	2.39	121.84	118.08
22	Z	313	CLA	CHD-C1D-ND	-2.39	122.26	124.45
22	B	825	CLA	CHA-C4D-ND	2.39	137.49	132.50
22	6	308	CLA	CHA-C4D-ND	2.39	137.49	132.50
34	J	101	LMG	C8-O7-C10	-2.39	111.91	117.79
22	K	204	CLA	CHA-C4D-ND	2.39	137.49	132.50
22	1	311	CLA	CMA-C3A-C4A	2.39	118.19	111.77
22	Z	307	CLA	CMD-C2D-C3D	-2.39	122.12	127.61
22	4	811	CLA	O2D-CGD-O1D	-2.39	119.17	123.84
22	A	841	CLA	CHA-C4D-ND	2.39	137.49	132.50
22	5	301	CLA	CHA-C4D-ND	2.39	137.49	132.50
22	6	301	CLA	O2A-CGA-CBA	2.39	119.40	111.91
22	A	839	CLA	O2D-CGD-O1D	-2.39	119.17	123.84
22	L	202	CLA	CHA-C4D-ND	2.39	137.49	132.50
22	6	309	CLA	CMD-C2D-C3D	-2.39	122.13	127.61
24	B	846	BCR	C8-C9-C10	2.39	122.60	118.94
22	A	832	CLA	C6-C5-C3	-2.38	107.20	113.45
22	8	306	CLA	O2D-CGD-O1D	-2.38	119.18	123.84
22	8	305	CLA	CMD-C2D-C3D	-2.38	122.13	127.61
24	3	307	BCR	C38-C26-C27	2.38	118.19	113.62
22	6	308	CLA	O2A-CGA-CBA	2.38	119.39	111.91
38	Z	301	LUT	C21-C26-C27	-2.38	109.69	112.70
22	9	313	CLA	O2A-CGA-CBA	2.38	119.38	111.91
22	B	830	CLA	CHA-C4D-ND	2.38	137.48	132.50
22	Z	314	CLA	CHA-C4D-ND	2.38	137.48	132.50
22	A	822	CLA	C1D-ND-C4D	-2.38	104.64	106.33

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Mol	Chain	Res	Type	Atoms	Z	Observed(^{o})	Ideal(^{o})
22	Z	313	CLA	O2D-CGD-O1D	-2.38	119.19	123.84
22	A	820	CLA	C1-C2-C3	-2.38	121.93	126.04
24	7	303	BCR	C23-C24-C25	-2.38	120.52	127.20
22	A	818	CLA	CMD-C2D-C3D	-2.38	122.14	127.61
24	B	849	BCR	C37-C22-C21	-2.38	119.59	122.92
22	9	303	CLA	C3D-C2D-C1D	-2.38	102.58	105.83
22	B	810	CLA	O2D-CGD-O1D	-2.38	119.19	123.84
22	4	808	CLA	CHA-C4D-ND	2.38	137.47	132.50
22	Z	303	CLA	O2A-CGA-CBA	2.38	119.37	111.91
22	2	304	CLA	CHA-C4D-ND	2.38	137.47	132.50
24	B	846	BCR	C37-C22-C21	-2.38	119.59	122.92
22	A	806	CLA	O2D-CGD-O1D	-2.38	119.19	123.84
39	9	312	CHL	C3C-C4C-NC	-2.37	107.91	110.57
22	B	818	CLA	O2A-CGA-CBA	2.37	119.36	111.91
22	B	814	CLA	CMD-C2D-C3D	-2.37	122.15	127.61
25	Z	317	LHG	C5-O7-C7	-2.37	111.95	117.79
22	9	308	CLA	O2D-CGD-O1D	-2.37	119.20	123.84
22	1	316	CLA	C2A-C1A-CHA	2.37	128.00	123.86
39	9	314	CHL	C2C-C3C-C4C	2.37	108.18	106.49
24	5	304	BCR	C23-C24-C25	-2.37	120.54	127.20
22	7	304	CLA	O2A-CGA-CBA	2.37	119.35	111.91
22	A	823	CLA	C11-C12-C13	-2.37	108.26	115.92
38	Z	301	LUT	C31-C30-C29	-2.37	123.93	127.31
22	B	825	CLA	CAA-CBA-CGA	-2.37	106.33	113.25
22	A	838	CLA	O2D-CGD-O1D	-2.37	119.21	123.84
22	A	805	CLA	CHA-C4D-ND	2.37	137.45	132.50
39	8	312	CHL	C1-O2A-CGA	2.37	122.65	116.44
22	A	824	CLA	CHA-C4D-ND	2.37	137.45	132.50
22	3	314	CLA	O2D-CGD-O1D	-2.37	119.21	123.84
22	7	309	CLA	O2A-CGA-CBA	2.37	119.33	111.91
22	B	824	CLA	O2A-CGA-CBA	2.37	119.33	111.91
32	F	306	RRX	C37-C22-C21	-2.37	119.61	122.92
22	A	807	CLA	CMD-C2D-C3D	-2.36	122.17	127.61
22	A	821	CLA	O2A-CGA-CBA	2.36	119.33	111.91
22	B	839	CLA	O2A-CGA-CBA	2.36	119.33	111.91
22	A	823	CLA	CMD-C2D-C3D	-2.36	122.17	127.61
25	7	318	LHG	O8-C23-C24	2.36	119.33	111.91
22	F	302	CLA	O2A-CGA-CBA	2.36	119.33	111.91
22	F	304	CLA	O2D-CGD-O1D	-2.36	119.22	123.84
22	B	829	CLA	CHA-C4D-ND	2.36	137.44	132.50
22	Z	304	CLA	CHA-C4D-ND	2.36	137.44	132.50
22	1	313	CLA	O2D-CGD-O1D	-2.36	119.22	123.84

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Mol	Chain	Res	Type	Atoms	Z	Observed(^{o})	Ideal(^{o})
22	1	316	CLA	O2D-CGD-O1D	-2.36	119.22	123.84
24	3	307	BCR	C29-C30-C25	2.36	114.12	110.48
22	5	320	CLA	C1-O2A-CGA	2.36	122.64	116.44
22	9	313	CLA	O2D-CGD-O1D	-2.36	119.22	123.84
24	A	848	BCR	C37-C22-C21	-2.36	119.62	122.92
22	1	315	CLA	CMD-C2D-C3D	-2.36	122.19	127.61
39	5	321	CHL	CHC-C1C-NC	2.36	127.78	124.20
24	B	845	BCR	C30-C25-C24	2.36	122.45	115.78
22	4	805	CLA	CMD-C2D-C3D	-2.36	122.19	127.61
39	4	813	CHL	C1-O2A-CGA	2.36	122.63	116.44
24	B	847	BCR	C2-C1-C6	2.36	114.11	110.48
22	G	202	CLA	O2D-CGD-O1D	-2.36	119.23	123.84
22	A	839	CLA	O2A-CGA-CBA	2.36	119.31	111.91
22	A	808	CLA	O2D-CGD-O1D	-2.36	119.23	123.84
22	5	313	CLA	CHA-C1A-NA	-2.36	121.00	126.40
27	A	852	DGA	OG1-CA1-CA2	2.36	119.30	111.91
22	A	825	CLA	CMD-C2D-C3D	-2.36	122.19	127.61
22	5	319	CLA	O2D-CGD-O1D	-2.36	119.23	123.84
22	F	304	CLA	CMD-C2D-C3D	-2.36	122.20	127.61
38	8	303	LUT	C35-C15-C14	-2.35	118.65	123.47
22	A	837	CLA	O2D-CGD-O1D	-2.35	119.24	123.84
22	Z	310	CLA	CHA-C4D-ND	2.35	137.42	132.50
22	5	326	CLA	O2D-CGD-O1D	-2.35	119.24	123.84
22	B	809	CLA	O2A-CGA-CBA	2.35	119.29	111.91
22	5	322	CLA	O2A-CGA-CBA	2.35	119.29	111.91
22	G	201	CLA	O2D-CGD-O1D	-2.35	119.24	123.84
39	5	321	CHL	C2C-C3C-C4C	2.35	108.17	106.49
39	5	317	CHL	CHC-C1C-NC	2.35	127.77	124.20
22	Z	309	CLA	CHA-C4D-ND	2.35	137.42	132.50
22	7	309	CLA	CHA-C4D-ND	2.35	137.42	132.50
36	J	105	C7Z	C11-C12-C13	-2.35	119.81	126.42
22	B	824	CLA	O2D-CGD-O1D	-2.35	119.24	123.84
22	1	315	CLA	O2D-CGD-O1D	-2.35	119.24	123.84
22	3	320	CLA	C3D-C2D-C1D	-2.35	102.62	105.83
22	8	309	CLA	O2A-CGA-CBA	2.35	119.28	111.91
22	9	303	CLA	CHA-C1A-NA	-2.35	121.02	126.40
22	1	311	CLA	C1D-ND-C4D	-2.35	104.67	106.33
24	B	849	BCR	C23-C24-C25	-2.35	120.60	127.20
22	A	802	CLA	O2D-CGD-O1D	-2.35	119.25	123.84
22	A	833	CLA	O2A-CGA-CBA	2.35	119.28	111.91
22	B	806	CLA	O2A-CGA-CBA	2.35	119.28	111.91
22	B	801	CLA	O2D-CGD-O1D	-2.35	119.25	123.84

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Mol	Chain	Res	Type	Atoms	Z	Observed(^{o})	Ideal(^{o})
22	J	103	CLA	O2D-CGD-O1D	-2.35	119.25	123.84
22	4	808	CLA	O2D-CGD-O1D	-2.35	119.25	123.84
39	4	814	CHL	CHC-C1C-NC	2.35	127.76	124.20
22	A	809	CLA	CHA-C4D-ND	2.35	137.41	132.50
22	B	811	CLA	O2D-CGD-O1D	-2.35	119.25	123.84
39	5	316	CHL	C1-C2-C3	-2.35	121.99	126.04
24	A	844	BCR	C7-C6-C5	-2.35	115.78	121.46
22	A	841	CLA	O2D-CGD-O1D	-2.35	119.25	123.84
22	7	311	CLA	O2D-CGD-O1D	-2.35	119.25	123.84
24	A	846	BCR	C35-C13-C14	-2.35	119.64	122.92
22	B	832	CLA	CMD-C2D-C3D	-2.34	122.22	127.61
22	A	811	CLA	O2D-CGD-O1D	-2.34	119.25	123.84
22	7	307	CLA	O2D-CGD-O1D	-2.34	119.25	123.84
24	F	303	BCR	C30-C25-C26	-2.34	119.31	122.61
22	A	813	CLA	O2A-CGA-CBA	2.34	119.26	111.91
22	1	307	CLA	O2D-CGD-O1D	-2.34	119.26	123.84
22	1	307	CLA	CMD-C2D-C3D	-2.34	122.23	127.61
22	B	808	CLA	O2D-CGD-O1D	-2.34	119.26	123.84
22	9	305	CLA	CHD-C1D-ND	-2.34	122.30	124.45
22	B	816	CLA	O2A-CGA-CBA	2.34	119.24	111.91
22	B	806	CLA	O2D-CGD-O1D	-2.34	119.27	123.84
22	Z	305	CLA	O2D-CGD-O1D	-2.34	119.27	123.84
24	A	845	BCR	C11-C12-C13	-2.34	119.85	126.42
22	B	822	CLA	O2D-CGD-O1D	-2.34	119.27	123.84
31	1	319	DGD	O6D-C5D-C6D	2.34	111.38	106.67
39	6	320	CHL	CHC-C1C-NC	2.34	127.75	124.20
22	B	812	CLA	O2D-CGD-O1D	-2.34	119.27	123.84
38	3	303	LUT	C30-C31-C32	-2.34	115.93	123.22
22	3	315	CLA	C1C-C2C-C3C	-2.34	104.50	106.96
24	A	856	BCR	C29-C28-C27	2.33	116.59	111.38
22	Z	305	CLA	O2A-CGA-CBA	2.33	119.23	111.91
22	7	312	CLA	O2D-CGD-O1D	-2.33	119.27	123.84
22	A	842	CLA	O2A-CGA-CBA	2.33	119.23	111.91
39	8	315	CHL	C1-C2-C3	-2.33	122.01	126.04
22	A	819	CLA	CMD-C2D-C3D	-2.33	122.25	127.61
22	6	322	CLA	O1D-CGD-CBD	-2.33	119.71	124.48
38	9	301	LUT	C21-C26-C27	-2.33	109.75	112.70
22	7	304	CLA	CHA-C4D-ND	2.33	137.38	132.50
24	A	847	BCR	C38-C26-C27	2.33	118.10	113.62
24	B	853	BCR	C35-C13-C14	-2.33	119.66	122.92
22	B	829	CLA	O2A-CGA-CBA	2.33	119.22	111.91
39	4	816	CHL	CHC-C1C-NC	2.33	127.74	124.20

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Mol	Chain	Res	Type	Atoms	Z	Observed(^{o})	Ideal(^{o})
22	Z	315	CLA	O2D-CGD-O1D	-2.33	119.28	123.84
22	K	205	CLA	C1D-ND-C4D	-2.33	104.68	106.33
22	A	810	CLA	CMD-C2D-C3D	-2.33	122.25	127.61
22	7	308	CLA	O2D-CGD-O1D	-2.33	119.28	123.84
22	A	838	CLA	O2A-CGA-CBA	2.33	119.22	111.91
22	7	310	CLA	CHA-C4D-ND	2.33	137.37	132.50
22	3	322	CLA	O2D-CGD-O1D	-2.33	119.28	123.84
22	3	319	CLA	O2D-CGD-O1D	-2.33	119.29	123.84
22	B	819	CLA	O2D-CGD-O1D	-2.33	119.29	123.84
22	5	314	CLA	O2D-CGD-O1D	-2.33	119.29	123.84
22	3	308	CLA	CMD-C2D-C3D	-2.33	122.26	127.61
22	6	317	CLA	O2A-CGA-CBA	2.33	119.21	111.91
22	1	309	CLA	CAA-CBA-CGA	-2.33	106.46	113.25
22	A	820	CLA	O2A-CGA-CBA	2.32	119.20	111.91
22	A	805	CLA	O2A-CGA-CBA	2.32	119.20	111.91
22	5	323	CLA	CMD-C2D-C3D	-2.32	122.27	127.61
22	5	309	CLA	CHA-C1A-NA	-2.32	121.08	126.40
22	Z	314	CLA	O2D-CGD-O1D	-2.32	119.30	123.84
22	B	840	CLA	O2A-CGA-CBA	2.32	119.20	111.91
40	1	318	SQD	O3-C3-C2	-2.32	104.98	110.35
22	A	836	CLA	O2D-CGD-O1D	-2.32	119.30	123.84
22	9	305	CLA	CHA-C1A-NA	-2.32	121.08	126.40
22	1	306	CLA	O2D-CGD-O1D	-2.32	119.30	123.84
22	5	326	CLA	CHA-C1A-NA	-2.32	121.08	126.40
22	8	316	CLA	O2D-CGD-O1D	-2.32	119.30	123.84
22	3	318	CLA	O2D-CGD-O1D	-2.32	119.30	123.84
38	7	301	LUT	C8-C7-C6	-2.32	120.69	127.20
22	B	801	CLA	O2A-CGA-CBA	2.32	119.19	111.91
22	Z	310	CLA	O2A-CGA-CBA	2.32	119.19	111.91
24	G	203	BCR	C37-C22-C21	-2.32	119.67	122.92
22	5	313	CLA	O2D-CGD-O1D	-2.32	119.31	123.84
24	6	306	BCR	C29-C30-C25	2.32	114.05	110.48
22	A	821	CLA	O2D-CGD-O1D	-2.32	119.31	123.84
22	6	301	CLA	O2D-CGD-O1D	-2.32	119.31	123.84
36	5	306	C7Z	C2-C3-C4	2.32	113.48	110.30
22	9	303	CLA	C6-C5-C3	-2.32	107.38	113.45
22	A	807	CLA	O2D-CGD-O1D	-2.32	119.31	123.84
22	6	317	CLA	O2D-CGD-O1D	-2.32	119.31	123.84
22	4	810	CLA	O2A-CGA-CBA	2.32	119.17	111.91
22	A	828	CLA	O2D-CGD-O1D	-2.32	119.31	123.84
22	F	305	CLA	CMD-C2D-C3D	-2.32	122.29	127.61
22	B	819	CLA	O2A-CGA-CBA	2.32	119.17	111.91

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Mol	Chain	Res	Type	Atoms	Z	Observed(^{o})	Ideal(^{o})
24	B	844	BCR	C37-C22-C21	-2.32	119.68	122.92
24	A	847	BCR	C30-C25-C26	-2.31	119.35	122.61
22	1	314	CLA	CHA-C4D-ND	2.31	137.34	132.50
22	A	839	CLA	CMD-C2D-C3D	-2.31	122.29	127.61
22	A	840	CLA	CMD-C2D-C3D	-2.31	122.29	127.61
39	2	303	CHL	CHC-C1C-NC	2.31	127.71	124.20
22	5	315	CLA	O2D-CGD-O1D	-2.31	119.32	123.84
39	5	316	CHL	CMA-C3A-C2A	2.31	123.16	113.83
22	8	307	CLA	CHA-C1A-NA	-2.31	121.10	126.40
22	6	314	CLA	O2A-CGA-CBA	2.31	119.16	111.91
22	5	307	CLA	O1D-CGD-CBD	-2.31	119.75	124.48
22	6	313	CLA	O2A-CGA-CBA	2.31	119.16	111.91
22	K	202	CLA	O2D-CGD-O1D	-2.31	119.32	123.84
22	A	816	CLA	CHA-C1A-NA	-2.31	121.11	126.40
38	6	303	LUT	C30-C31-C32	-2.31	116.01	123.22
22	6	301	CLA	C6-C5-C3	-2.31	107.40	113.45
39	Z	312	CHL	CHC-C1C-NC	2.31	127.71	124.20
22	B	815	CLA	O2D-CGD-O1D	-2.31	119.32	123.84
22	F	305	CLA	O2D-CGD-O1D	-2.31	119.32	123.84
22	6	313	CLA	O2D-CGD-O1D	-2.31	119.32	123.84
22	6	311	CLA	O2D-CGD-O1D	-2.31	119.32	123.84
22	A	811	CLA	CMD-C2D-C3D	-2.31	122.30	127.61
22	Z	308	CLA	CMA-C3A-C4A	2.31	117.98	111.77
22	1	309	CLA	O2A-CGA-CBA	2.31	119.15	111.91
22	B	816	CLA	O2D-CGD-O1D	-2.31	119.33	123.84
24	B	845	BCR	C8-C7-C6	-2.31	120.72	127.20
22	8	311	CLA	O2A-CGA-CBA	2.31	119.15	111.91
41	8	320	3PH	O31-C31-C32	2.31	119.15	111.91
22	6	307	CLA	CMD-C2D-C3D	-2.31	122.31	127.61
38	8	302	LUT	C18-C5-C4	2.31	118.63	114.36
24	B	845	BCR	C11-C12-C13	-2.31	119.94	126.42
22	B	815	CLA	O2A-CGA-CBA	2.31	119.14	111.91
24	6	305	BCR	C37-C22-C21	-2.31	119.69	122.92
22	3	322	CLA	CMD-C2D-C3D	-2.31	122.31	127.61
22	7	308	CLA	O2A-CGA-CBA	2.31	119.14	111.91
22	7	323	CLA	O2A-CGA-CBA	2.31	119.14	111.91
24	G	203	BCR	C16-C15-C14	-2.31	118.75	123.47
24	B	847	BCR	C8-C7-C6	-2.31	120.73	127.20
22	6	302	CLA	O2A-CGA-CBA	2.30	119.14	111.91
22	A	821	CLA	CMA-C3A-C4A	2.30	117.97	111.77
22	K	205	CLA	CHA-C1A-NA	-2.30	121.12	126.40
22	8	310	CLA	O2D-CGD-O1D	-2.30	119.34	123.84

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Mol	Chain	Res	Type	Atoms	Z	Observed(^{o})	Ideal(^{o})
22	B	840	CLA	CMD-C2D-C3D	-2.30	122.32	127.61
22	9	306	CLA	O2D-CGD-O1D	-2.30	119.34	123.84
24	6	306	BCR	C35-C13-C12	2.30	121.70	118.08
22	A	855	CLA	O2D-CGD-O1D	-2.30	119.34	123.84
22	B	807	CLA	O2D-CGD-O1D	-2.30	119.34	123.84
22	3	310	CLA	O2D-CGD-O1D	-2.30	119.34	123.84
24	3	305	BCR	C1-C6-C7	2.30	122.28	115.78
24	7	303	BCR	C11-C12-C13	-2.30	119.96	126.42
22	9	306	CLA	CMD-C2D-C3D	-2.30	122.33	127.61
22	3	319	CLA	O2A-CGA-CBA	2.30	119.11	111.91
22	A	809	CLA	O2D-CGD-O1D	-2.30	119.35	123.84
21	A	801	CL0	CMD-C2D-C3D	-2.30	122.33	127.61
38	4	802	LUT	C15-C35-C34	-2.30	118.77	123.47
22	7	304	CLA	CAA-C2A-C3A	-2.30	106.49	112.78
22	A	823	CLA	O2A-CGA-CBA	2.30	119.11	111.91
22	6	322	CLA	CMD-C2D-C3D	-2.30	122.33	127.61
22	5	301	CLA	O2D-CGD-O1D	-2.29	119.35	123.84
38	3	303	LUT	C11-C12-C13	-2.29	119.97	126.42
22	4	806	CLA	O2A-CGA-CBA	2.29	119.11	111.91
22	L	201	CLA	O2D-CGD-O1D	-2.29	119.35	123.84
39	4	819	CHL	CHD-C4C-C3C	2.29	128.21	124.84
38	4	802	LUT	C11-C12-C13	-2.29	119.97	126.42
22	A	813	CLA	O2D-CGD-O1D	-2.29	119.35	123.84
24	B	853	BCR	C37-C22-C21	-2.29	119.71	122.92
22	A	817	CLA	O2D-CGD-O1D	-2.29	119.36	123.84
22	A	834	CLA	O2A-CGA-CBA	2.29	119.09	111.91
22	6	312	CLA	O2D-CGD-O1D	-2.29	119.36	123.84
22	1	311	CLA	CHA-C1A-NA	-2.29	121.16	126.40
22	B	824	CLA	CMD-C2D-C3D	-2.29	122.35	127.61
39	2	303	CHL	C1-O2A-CGA	2.29	122.44	116.44
22	1	305	CLA	O2D-CGD-O1D	-2.29	119.37	123.84
22	3	310	CLA	C6-C7-C8	-2.29	108.53	115.92
24	B	853	BCR	C15-C14-C13	-2.29	124.05	127.31
22	A	820	CLA	O2D-CGD-O1D	-2.29	119.37	123.84
22	7	315	CLA	CAA-C2A-C3A	-2.29	106.52	112.78
22	A	814	CLA	O2D-CGD-O1D	-2.28	119.37	123.84
22	A	831	CLA	O2D-CGD-O1D	-2.28	119.37	123.84
36	5	306	C7Z	C31-C32-C33	-2.28	120.00	126.42
22	3	309	CLA	O2D-CGD-O1D	-2.28	119.37	123.84
22	A	837	CLA	CMD-C2D-C3D	-2.28	122.36	127.61
22	7	324	CLA	C1-O2A-CGA	2.28	122.44	116.44
22	A	832	CLA	O2D-CGD-O1D	-2.28	119.38	123.84

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Mol	Chain	Res	Type	Atoms	Z	Observed(^{o})	Ideal(^{o})
22	A	814	CLA	O2A-CGA-CBA	2.28	119.07	111.91
38	Z	302	LUT	C19-C9-C10	-2.28	119.73	122.92
38	7	301	LUT	C11-C12-C13	-2.28	120.01	126.42
22	B	842	CLA	CMD-C2D-C3D	-2.28	122.37	127.61
22	4	810	CLA	CMD-C2D-C3D	-2.28	122.37	127.61
39	7	313	CHL	CHC-C1C-NC	2.28	127.66	124.20
22	B	828	CLA	CHA-C1A-NA	-2.28	121.17	126.40
22	1	316	CLA	C1-C2-C3	-2.28	122.10	126.04
22	6	321	CLA	O2D-CGD-O1D	-2.28	119.38	123.84
22	4	805	CLA	CHA-C1A-NA	-2.28	121.18	126.40
38	Z	301	LUT	C40-C33-C34	-2.28	119.73	122.92
24	5	305	BCR	C19-C18-C17	-2.28	115.45	118.94
22	A	825	CLA	O2D-CGD-O1D	-2.28	119.39	123.84
22	8	305	CLA	CMB-C2B-C3B	2.28	128.94	124.68
22	B	823	CLA	O2A-CGA-CBA	2.28	119.05	111.91
32	F	306	RRX	C34-C9-C10	-2.28	119.73	122.92
22	B	815	CLA	CMD-C2D-C3D	-2.28	122.38	127.61
22	A	831	CLA	O2A-CGA-CBA	2.28	119.05	111.91
22	L	203	CLA	O2D-CGD-O1D	-2.28	119.39	123.84
22	5	311	CLA	CHA-C1A-NA	-2.28	121.19	126.40
22	4	809	CLA	O2D-CGD-O1D	-2.28	119.39	123.84
22	B	818	CLA	CMD-C2D-C3D	-2.27	122.38	127.61
22	8	306	CLA	CMD-C2D-C3D	-2.27	122.38	127.61
22	2	301	CLA	O2D-CGD-O1D	-2.27	119.39	123.84
22	A	840	CLA	O2A-CGA-CBA	2.27	119.04	111.91
22	B	813	CLA	O2A-CGA-CBA	2.27	119.04	111.91
22	1	310	CLA	O2D-CGD-O1D	-2.27	119.39	123.84
22	7	322	CLA	O2D-CGD-O1D	-2.27	119.39	123.84
22	3	316	CLA	CMD-C2D-C3D	-2.27	122.39	127.61
24	A	848	BCR	C8-C9-C10	2.27	122.43	118.94
24	G	203	BCR	C11-C10-C9	-2.27	124.07	127.31
22	A	832	CLA	CMD-C2D-C3D	-2.27	122.39	127.61
22	Z	316	CLA	CMD-C2D-C3D	-2.27	122.39	127.61
22	A	818	CLA	O2D-CGD-O1D	-2.27	119.40	123.84
22	Z	303	CLA	CMD-C2D-C3D	-2.27	122.39	127.61
22	1	316	CLA	C3D-C2D-C1D	-2.27	102.73	105.83
22	Z	306	CLA	O2D-CGD-O1D	-2.27	119.40	123.84
24	7	303	BCR	C38-C26-C27	2.27	117.97	113.62
22	Z	308	CLA	O2D-CGD-O1D	-2.27	119.40	123.84
24	A	846	BCR	C36-C18-C19	2.27	121.65	118.08
39	8	315	CHL	C4A-NA-C1A	2.27	107.72	106.71
22	A	810	CLA	O2D-CGD-O1D	-2.27	119.41	123.84

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Mol	Chain	Res	Type	Atoms	Z	Observed(^{o})	Ideal(^{o})
36	J	105	C7Z	C28-C27-C26	-2.27	120.84	127.20
22	A	812	CLA	O2A-CGA-CBA	2.27	119.02	111.91
22	A	814	CLA	CHA-C1A-NA	-2.27	121.21	126.40
22	8	308	CLA	O2D-CGD-O1D	-2.27	119.41	123.84
22	7	322	CLA	CMB-C2B-C3B	2.27	128.92	124.68
22	A	823	CLA	O2D-CGD-O1D	-2.26	119.41	123.84
22	Z	305	CLA	CHA-C4D-ND	2.26	137.24	132.50
22	Z	316	CLA	O2A-CGA-CBA	2.26	119.01	111.91
22	A	815	CLA	O2D-CGD-O1D	-2.26	119.41	123.84
22	Z	315	CLA	CMD-C2D-C3D	-2.26	122.41	127.61
22	A	826	CLA	O2A-CGA-CBA	2.26	119.01	111.91
22	9	310	CLA	CHA-C1A-NA	-2.26	121.21	126.40
22	K	204	CLA	O2A-CGA-CBA	2.26	119.01	111.91
22	6	310	CLA	CMD-C2D-C3D	-2.26	122.41	127.61
22	A	809	CLA	CMD-C2D-C3D	-2.26	122.41	127.61
22	B	809	CLA	CMD-C2D-C3D	-2.26	122.41	127.61
22	9	303	CLA	CHA-C4D-ND	2.26	137.23	132.50
39	6	316	CHL	CMA-C3A-C2A	2.26	122.95	113.83
22	A	824	CLA	O2D-CGD-O1D	-2.26	119.42	123.84
22	3	308	CLA	O1D-CGD-CBD	-2.26	119.86	124.48
22	B	811	CLA	CMD-C2D-C3D	-2.26	122.41	127.61
24	6	305	BCR	C8-C7-C6	-2.26	120.85	127.20
22	A	817	CLA	O2A-CGA-CBA	2.26	119.00	111.91
22	5	301	CLA	CMD-C2D-C3D	-2.26	122.41	127.61
22	5	310	CLA	O2D-CGD-O1D	-2.26	119.42	123.84
24	A	847	BCR	C23-C24-C25	-2.26	120.86	127.20
24	3	305	BCR	C11-C10-C9	-2.26	124.08	127.31
22	Z	306	CLA	C6-C5-C3	-2.26	107.53	113.45
22	A	839	CLA	CAA-CBA-CGA	-2.26	106.65	113.25
22	G	201	CLA	CHA-C1A-NA	-2.26	121.22	126.40
22	A	821	CLA	CMD-C2D-C3D	-2.26	122.42	127.61
22	Z	308	CLA	CMD-C2D-C3D	-2.26	122.42	127.61
22	B	818	CLA	O2D-CGD-O1D	-2.26	119.42	123.84
22	B	801	CLA	CHA-C1A-NA	-2.26	121.23	126.40
38	9	301	LUT	C8-C7-C6	-2.26	120.86	127.20
22	A	812	CLA	CMD-C2D-C3D	-2.26	122.42	127.61
22	A	806	CLA	O2A-CGA-CBA	2.26	118.99	111.91
22	8	309	CLA	CMD-C2D-C3D	-2.26	122.42	127.61
22	6	310	CLA	O2D-CGD-O1D	-2.26	119.43	123.84
22	B	812	CLA	O2A-CGA-CBA	2.26	118.99	111.91
22	A	855	CLA	CMD-C2D-C3D	-2.25	122.43	127.61
22	7	323	CLA	CMD-C2D-C3D	-2.25	122.43	127.61

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Mol	Chain	Res	Type	Atoms	Z	Observed(^{o})	Ideal(^{o})
22	6	309	CLA	O2D-CGD-O1D	-2.25	119.43	123.84
39	Z	311	CHL	C1-O2A-CGA	2.25	122.35	116.44
22	A	802	CLA	CMD-C2D-C3D	-2.25	122.43	127.61
22	3	318	CLA	CMD-C2D-C3D	-2.25	122.43	127.61
22	1	305	CLA	CMD-C2D-C3D	-2.25	122.44	127.61
22	A	841	CLA	CMD-C2D-C3D	-2.25	122.44	127.61
22	F	301	CLA	C6-C5-C3	-2.25	107.56	113.45
22	7	306	CLA	O2D-CGD-O1D	-2.25	119.44	123.84
22	1	310	CLA	CHA-C1A-NA	-2.25	121.25	126.40
22	B	836	CLA	O2D-CGD-O1D	-2.25	119.44	123.84
22	B	809	CLA	O2D-CGD-O1D	-2.25	119.44	123.84
25	A	850	LHG	C5-O7-C7	-2.25	112.26	117.79
22	7	324	CLA	CMD-C2D-C3D	-2.25	122.44	127.61
22	B	825	CLA	O2D-CGD-O1D	-2.25	119.44	123.84
22	B	836	CLA	CMB-C2B-C3B	2.25	128.88	124.68
22	B	805	CLA	O2A-CGA-CBA	2.25	118.95	111.91
22	A	822	CLA	CMD-C2D-C3D	-2.25	122.45	127.61
36	5	306	C7Z	C19-C9-C10	-2.25	119.78	122.92
22	5	323	CLA	O2D-CGD-O1D	-2.24	119.45	123.84
22	K	202	CLA	CMB-C2B-C3B	2.24	128.88	124.68
22	5	313	CLA	C1D-ND-C4D	-2.24	104.74	106.33
22	B	821	CLA	CMD-C2D-C3D	-2.24	122.45	127.61
22	L	203	CLA	O2A-CGA-CBA	2.24	118.95	111.91
22	F	302	CLA	O2D-CGD-O1D	-2.24	119.45	123.84
22	9	303	CLA	C2C-C1C-NC	2.24	112.07	109.97
22	3	319	CLA	CMD-C2D-C3D	-2.24	122.45	127.61
22	7	314	CLA	CMD-C2D-C3D	-2.24	122.45	127.61
22	B	821	CLA	O2D-CGD-O1D	-2.24	119.45	123.84
22	A	831	CLA	CMD-C2D-C3D	-2.24	122.46	127.61
38	5	302	LUT	C20-C13-C12	2.24	121.61	118.08
22	B	833	CLA	O2D-CGD-O1D	-2.24	119.46	123.84
38	4	802	LUT	C1-C6-C5	-2.24	119.46	122.61
22	5	315	CLA	CHA-C1A-NA	-2.24	121.27	126.40
22	4	805	CLA	C2D-C1D-ND	2.24	111.75	110.10
22	B	814	CLA	O2A-CGA-CBA	2.24	118.94	111.91
25	4	820	LHG	C5-O7-C7	-2.24	112.28	117.79
22	B	806	CLA	CMD-C2D-C3D	-2.24	122.46	127.61
22	A	838	CLA	CMD-C2D-C3D	-2.24	122.46	127.61
22	B	842	CLA	O2D-CGD-O1D	-2.24	119.46	123.84
22	B	808	CLA	CMD-C2D-C3D	-2.24	122.47	127.61
39	6	320	CHL	CHD-C4C-C3C	2.24	128.13	124.84
39	6	315	CHL	CHB-C4A-NA	2.24	127.61	124.51

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Mol	Chain	Res	Type	Atoms	Z	Observed(^{o})	Ideal(^{o})
22	A	805	CLA	CMD-C2D-C3D	-2.24	122.47	127.61
22	7	308	CLA	CMD-C2D-C3D	-2.24	122.47	127.61
22	8	310	CLA	CMD-C2D-C3D	-2.24	122.47	127.61
22	7	315	CLA	O1D-CGD-CBD	-2.24	119.91	124.48
22	1	313	CLA	CMD-C2D-C3D	-2.24	122.47	127.61
22	5	309	CLA	O2D-CGD-O1D	-2.24	119.47	123.84
24	8	304	BCR	C30-C25-C26	-2.24	119.47	122.61
39	5	316	CHL	CHB-C4A-NA	2.24	127.60	124.51
22	G	202	CLA	CMD-C2D-C3D	-2.24	122.47	127.61
22	9	306	CLA	CMB-C2B-C3B	2.24	128.86	124.68
36	5	306	C7Z	C4-C5-C6	-2.23	115.87	120.85
22	4	809	CLA	CMD-C2D-C3D	-2.23	122.47	127.61
24	3	307	BCR	C1-C6-C7	2.23	122.10	115.78
22	B	829	CLA	CMD-C2D-C3D	-2.23	122.47	127.61
38	6	303	LUT	C40-C33-C34	-2.23	119.79	122.92
22	6	311	CLA	CMB-C2B-C3B	2.23	128.86	124.68
22	A	805	CLA	O2D-CGD-O1D	-2.23	119.47	123.84
22	8	308	CLA	CMD-C2D-C3D	-2.23	122.48	127.61
22	5	301	CLA	O2A-CGA-CBA	2.23	118.91	111.91
22	B	823	CLA	CMD-C2D-C3D	-2.23	122.48	127.61
24	B	848	BCR	C29-C30-C25	2.23	113.92	110.48
39	8	301	CHL	CMA-C3A-C2A	2.23	122.83	113.83
22	B	835	CLA	O1D-CGD-CBD	-2.23	119.92	124.48
22	B	810	CLA	CMD-C2D-C3D	-2.23	122.48	127.61
22	4	812	CLA	CMD-C2D-C3D	-2.23	122.48	127.61
22	B	839	CLA	CMD-C2D-C3D	-2.23	122.48	127.61
22	7	304	CLA	CMD-C2D-C3D	-2.23	122.48	127.61
22	B	839	CLA	O1D-CGD-CBD	-2.23	119.92	124.48
22	K	203	CLA	CMD-C2D-C3D	-2.23	122.49	127.61
22	L	201	CLA	O2A-CGA-CBA	2.23	118.90	111.91
22	B	838	CLA	C1-C2-C3	-2.23	122.19	126.04
31	3	301	DGD	C2G-O2G-C1B	-2.23	112.31	117.79
22	7	305	CLA	O2A-CGA-CBA	2.23	118.90	111.91
24	B	848	BCR	C7-C6-C5	-2.23	116.06	121.46
22	3	316	CLA	O2A-CGA-CBA	2.23	118.90	111.91
22	Z	316	CLA	O2D-CGD-O1D	-2.23	119.48	123.84
22	A	817	CLA	CMD-C2D-C3D	-2.23	122.49	127.61
22	B	805	CLA	O2D-CGD-O1D	-2.23	119.48	123.84
22	L	203	CLA	CMD-C2D-C3D	-2.23	122.49	127.61
22	A	815	CLA	CMD-C2D-C3D	-2.23	122.49	127.61
22	A	833	CLA	CMD-C2D-C3D	-2.23	122.49	127.61
22	5	309	CLA	CMD-C2D-C3D	-2.23	122.49	127.61

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Mol	Chain	Res	Type	Atoms	Z	Observed(^{o})	Ideal(^{o})
22	A	813	CLA	CMD-C2D-C3D	-2.23	122.49	127.61
22	L	201	CLA	CMD-C2D-C3D	-2.23	122.49	127.61
22	B	832	CLA	O2D-CGD-O1D	-2.23	119.49	123.84
24	7	303	BCR	C27-C26-C25	-2.22	119.50	122.73
39	5	317	CHL	CHD-C4C-C3C	2.22	128.11	124.84
22	6	312	CLA	CMD-C2D-C3D	-2.22	122.50	127.61
22	B	827	CLA	CMD-C2D-C3D	-2.22	122.50	127.61
22	3	309	CLA	CMD-C2D-C3D	-2.22	122.50	127.61
36	J	105	C7Z	C2-C3-C4	2.22	113.34	110.30
38	4	803	LUT	C8-C7-C6	-2.22	120.96	127.20
22	B	813	CLA	CMD-C2D-C3D	-2.22	122.50	127.61
22	6	319	CLA	O1D-CGD-CBD	-2.22	119.94	124.48
24	B	849	BCR	C29-C30-C25	2.22	113.90	110.48
22	A	816	CLA	CAA-CBA-CGA	-2.22	106.76	113.25
22	B	825	CLA	CMD-C2D-C3D	-2.22	122.51	127.61
24	3	304	BCR	C19-C18-C17	-2.22	115.53	118.94
22	4	811	CLA	CHA-C1A-NA	-2.22	121.31	126.40
24	F	303	BCR	C27-C26-C25	-2.22	119.51	122.73
22	8	316	CLA	CMD-C2D-C3D	-2.22	122.51	127.61
24	I	4001	BCR	C23-C24-C25	-2.22	120.97	127.20
36	5	306	C7Z	C40-C33-C34	-2.22	119.81	122.92
22	B	828	CLA	CMD-C2D-C3D	-2.22	122.51	127.61
22	K	203	CLA	CHA-C1A-NA	-2.22	121.32	126.40
22	4	806	CLA	CHA-C1A-NA	-2.22	121.32	126.40
24	B	846	BCR	C35-C13-C14	-2.22	119.82	122.92
24	B	846	BCR	C23-C24-C25	-2.22	120.97	127.20
22	6	311	CLA	O2A-CGA-CBA	2.22	118.86	111.91
22	3	322	CLA	O2A-CGA-CBA	2.22	118.86	111.91
22	A	835	CLA	O2D-CGD-O1D	-2.22	119.51	123.84
22	9	310	CLA	C3D-C2D-C1D	-2.22	102.81	105.83
22	4	818	CLA	O1D-CGD-CBD	-2.21	119.95	124.48
22	A	836	CLA	CMD-C2D-C3D	-2.21	122.52	127.61
22	8	307	CLA	CMD-C2D-C3D	-2.21	122.52	127.61
22	5	308	CLA	O2A-CGA-CBA	2.21	118.85	111.91
22	3	313	CLA	CMD-C2D-C3D	-2.21	122.52	127.61
39	Z	311	CHL	CHC-C1C-NC	2.21	127.56	124.20
24	B	845	BCR	C15-C16-C17	-2.21	118.94	123.47
22	9	305	CLA	C3A-C2A-C1A	2.21	104.65	101.34
22	B	820	CLA	CMD-C2D-C3D	-2.21	122.52	127.61
22	A	817	CLA	C1D-ND-C4D	-2.21	104.76	106.33
22	4	818	CLA	CMD-C2D-C3D	-2.21	122.53	127.61
22	A	812	CLA	CHA-C1A-NA	-2.21	121.33	126.40

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Mol	Chain	Res	Type	Atoms	Z	Observed(^{o})	Ideal(^{o})
22	9	311	CLA	CHA-C1A-NA	-2.21	121.33	126.40
22	B	833	CLA	CMD-C2D-C3D	-2.21	122.53	127.61
22	6	311	CLA	CMD-C2D-C3D	-2.21	122.53	127.61
22	2	302	CLA	CMD-C2D-C3D	-2.21	122.53	127.61
22	6	308	CLA	CAA-CBA-CGA	-2.21	106.79	113.25
39	8	315	CHL	CHC-C1C-NC	2.21	127.56	124.20
22	3	311	CLA	O2D-CGD-O1D	-2.21	119.52	123.84
22	A	828	CLA	CMD-C2D-C3D	-2.21	122.53	127.61
22	6	301	CLA	CMD-C2D-C3D	-2.21	122.53	127.61
22	6	308	CLA	CMB-C2B-C3B	2.21	128.81	124.68
22	A	830	CLA	CMD-C2D-C3D	-2.21	122.53	127.61
22	3	312	CLA	CMD-C2D-C3D	-2.21	122.53	127.61
22	8	306	CLA	O2A-CGA-CBA	2.21	118.84	111.91
38	5	302	LUT	C39-C29-C28	2.21	121.56	118.08
22	L	202	CLA	O2D-CGD-O1D	-2.21	119.52	123.84
22	7	312	CLA	CHA-C1A-NA	-2.21	121.34	126.40
24	K	206	BCR	C34-C9-C10	-2.21	119.83	122.92
22	B	842	CLA	CHA-C1A-NA	-2.21	121.35	126.40
24	5	304	BCR	C36-C18-C17	-2.21	119.83	122.92
22	8	314	CLA	O2D-CGD-O1D	-2.20	119.53	123.84
22	B	819	CLA	CMD-C2D-C3D	-2.20	122.54	127.61
22	B	836	CLA	CMD-C2D-C3D	-2.20	122.54	127.61
22	9	308	CLA	CMD-C2D-C3D	-2.20	122.54	127.61
24	4	804	BCR	C23-C24-C25	-2.20	121.01	127.20
22	F	304	CLA	CHA-C1A-NA	-2.20	121.35	126.40
39	6	318	CHL	CHC-C1C-NC	2.20	127.55	124.20
22	A	834	CLA	CMD-C2D-C3D	-2.20	122.55	127.61
22	L	202	CLA	CMD-C2D-C3D	-2.20	122.55	127.61
22	B	833	CLA	O2A-CGA-CBA	2.20	118.82	111.91
33	4	822	LMT	C1B-O5B-C5B	2.20	118.01	113.69
24	B	849	BCR	C36-C18-C17	-2.20	119.84	122.92
22	F	301	CLA	O2D-CGD-O1D	-2.20	119.53	123.84
22	4	810	CLA	O2D-CGD-O1D	-2.20	119.53	123.84
22	5	312	CLA	CMD-C2D-C3D	-2.20	122.55	127.61
24	5	305	BCR	C1-C6-C7	2.20	122.00	115.78
22	4	805	CLA	C1D-ND-C4D	-2.20	104.77	106.33
31	1	319	DGD	O2G-C1B-O1B	-2.20	118.39	123.70
22	A	802	CLA	C1-O2A-CGA	2.20	122.22	116.44
32	F	306	RRX	C35-C13-C12	-2.20	114.61	118.08
22	4	817	CLA	CMD-C2D-C3D	-2.20	122.55	127.61
24	7	303	BCR	C37-C22-C21	-2.20	119.84	122.92
39	4	816	CHL	C2C-C3C-C4C	2.20	108.06	106.49

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Mol	Chain	Res	Type	Atoms	Z	Observed(^{o})	Ideal(^{o})
22	Z	310	CLA	CHA-C1A-NA	-2.20	121.37	126.40
22	A	827	CLA	CMD-C2D-C3D	-2.20	122.56	127.61
22	B	834	CLA	CMD-C2D-C3D	-2.20	122.56	127.61
22	A	824	CLA	C6-C5-C3	-2.20	107.70	113.45
22	K	204	CLA	CMD-C2D-C3D	-2.19	122.57	127.61
22	B	831	CLA	O2A-CGA-CBA	2.19	118.79	111.91
22	A	820	CLA	CMD-C2D-C3D	-2.19	122.57	127.61
22	7	317	CLA	CMD-C2D-C3D	-2.19	122.57	127.61
39	4	816	CHL	CMB-C2B-C1B	-2.19	125.09	128.46
22	A	842	CLA	CMD-C2D-C3D	-2.19	122.57	127.61
39	6	318	CHL	CMB-C2B-C1B	-2.19	125.09	128.46
38	5	303	LUT	C22-C23-C24	2.19	114.24	111.74
38	6	304	LUT	C8-C7-C6	-2.19	121.05	127.20
22	F	301	CLA	CMD-C2D-C3D	-2.19	122.57	127.61
22	5	308	CLA	CMD-C2D-C3D	-2.19	122.57	127.61
39	5	316	CHL	C2C-C3C-C4C	2.19	108.05	106.49
22	A	804	CLA	CMD-C2D-C3D	-2.19	122.57	127.61
22	G	201	CLA	CMD-C2D-C3D	-2.19	122.57	127.61
24	K	206	BCR	C37-C22-C21	-2.19	119.85	122.92
39	6	315	CHL	CMB-C2B-C1B	-2.19	125.10	128.46
22	7	309	CLA	CMD-C2D-C3D	-2.19	122.58	127.61
22	6	319	CLA	CMD-C2D-C3D	-2.19	122.58	127.61
24	B	846	BCR	C8-C7-C6	-2.19	121.05	127.20
39	9	314	CHL	CHB-C4A-NA	2.19	127.54	124.51
24	A	846	BCR	C34-C9-C10	-2.19	119.86	122.92
22	A	809	CLA	O1D-CGD-CBD	-2.19	120.01	124.48
31	3	301	DGD	O2G-C1B-O1B	-2.19	118.41	123.70
22	7	322	CLA	CHA-C1A-NA	-2.19	121.39	126.40
22	1	308	CLA	O2A-CGA-CBA	2.19	118.77	111.91
22	5	326	CLA	CAA-C2A-C3A	-2.19	106.79	112.78
22	B	821	CLA	O2A-CGA-CBA	2.19	118.77	111.91
22	3	318	CLA	O2A-CGA-CBA	2.19	118.77	111.91
22	A	840	CLA	O2D-CGD-O1D	-2.19	119.56	123.84
22	K	205	CLA	CMA-C3A-C4A	2.19	117.65	111.77
22	A	820	CLA	CHA-C1A-NA	-2.19	121.39	126.40
39	4	814	CHL	CMB-C2B-C1B	-2.19	125.10	128.46
22	6	321	CLA	CMD-C2D-C3D	-2.19	122.58	127.61
22	A	816	CLA	CMD-C2D-C3D	-2.19	122.58	127.61
22	2	302	CLA	O1D-CGD-CBD	-2.19	120.01	124.48
39	1	312	CHL	CMB-C2B-C1B	-2.19	125.11	128.46
39	5	316	CHL	CHC-C1C-NC	2.19	127.52	124.20
22	7	314	CLA	O2D-CGD-O1D	-2.18	119.57	123.84

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Mol	Chain	Res	Type	Atoms	Z	Observed(^{o})	Ideal(^{o})
22	5	319	CLA	O2A-CGA-CBA	2.18	118.76	111.91
22	6	322	CLA	O2A-CGA-CBA	2.18	118.76	111.91
22	6	314	CLA	CMD-C2D-C3D	-2.18	122.59	127.61
39	4	813	CHL	CMB-C2B-C1B	-2.18	125.11	128.46
22	B	835	CLA	CMB-C2B-C3B	2.18	128.76	124.68
22	J	103	CLA	CMD-C2D-C3D	-2.18	122.59	127.61
22	2	304	CLA	O2D-CGD-O1D	-2.18	119.57	123.84
22	A	803	CLA	CMD-C2D-C3D	-2.18	122.59	127.61
22	K	203	CLA	O2D-CGD-O1D	-2.18	119.57	123.84
22	6	311	CLA	CHA-C1A-NA	-2.18	121.40	126.40
22	7	312	CLA	CMD-C2D-C3D	-2.18	122.59	127.61
22	A	836	CLA	CMB-C2B-C3B	2.18	128.76	124.68
22	6	322	CLA	CMB-C2B-C3B	2.18	128.76	124.68
25	9	315	LHG	C9-C8-C7	-2.18	105.69	113.62
22	Z	304	CLA	O2D-CGD-O1D	-2.18	119.58	123.84
22	K	202	CLA	CMD-C2D-C3D	-2.18	122.60	127.61
22	B	830	CLA	O2A-CGA-CBA	2.18	118.74	111.91
22	A	826	CLA	CMD-C2D-C3D	-2.18	122.60	127.61
22	A	804	CLA	CHA-C1A-NA	-2.18	121.41	126.40
22	A	811	CLA	CHA-C1A-NA	-2.18	121.41	126.40
22	1	304	CLA	O1D-CGD-CBD	-2.18	120.03	124.48
22	4	809	CLA	CHA-C1A-NA	-2.18	121.41	126.40
22	7	304	CLA	O2D-CGD-O1D	-2.18	119.58	123.84
22	7	316	CLA	CMD-C2D-C3D	-2.18	122.61	127.61
22	9	307	CLA	CMD-C2D-C3D	-2.18	122.61	127.61
21	A	801	CL0	C4D-C3D-CAD	2.17	110.66	108.10
22	B	838	CLA	O2A-CGA-CBA	2.17	118.73	111.91
22	1	306	CLA	CMD-C2D-C3D	-2.17	122.61	127.61
39	Z	312	CHL	CMB-C2B-C1B	-2.17	125.12	128.46
39	8	301	CHL	CMB-C2B-C1B	-2.17	125.12	128.46
22	3	320	CLA	C2A-C1A-CHA	2.17	127.66	123.86
22	4	808	CLA	CMD-C2D-C3D	-2.17	122.62	127.61
22	8	308	CLA	O2A-CGA-CBA	2.17	118.72	111.91
22	7	309	CLA	O2D-CGD-O1D	-2.17	119.59	123.84
22	6	308	CLA	C3D-C2D-C1D	-2.17	102.87	105.83
22	3	310	CLA	CMD-C2D-C3D	-2.17	122.62	127.61
22	A	827	CLA	C3D-C2D-C1D	-2.17	102.87	105.83
22	9	304	CLA	C2A-C1A-CHA	2.17	127.65	123.86
22	Z	313	CLA	CHA-C1A-NA	-2.17	121.43	126.40
39	1	312	CHL	CHD-C4C-C3C	2.17	128.03	124.84
22	1	310	CLA	CMD-C2D-C3D	-2.17	122.62	127.61
22	2	302	CLA	CHA-C1A-NA	-2.17	121.43	126.40

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Mol	Chain	Res	Type	Atoms	Z	Observed(^{o})	Ideal(^{o})
22	B	826	CLA	O1D-CGD-CBD	-2.17	120.05	124.48
24	G	203	BCR	C4-C5-C6	-2.17	119.58	122.73
22	B	825	CLA	CMB-C2B-C3B	2.17	128.73	124.68
22	B	829	CLA	O2D-CGD-O1D	-2.17	119.60	123.84
39	2	303	CHL	CMA-C3A-C2A	2.17	122.57	113.83
22	A	827	CLA	O2D-CGD-O1D	-2.17	119.60	123.84
22	7	305	CLA	C3D-C2D-C1D	-2.17	102.87	105.83
22	1	304	CLA	CMD-C2D-C3D	-2.17	122.63	127.61
39	5	317	CHL	C1-O2A-CGA	2.17	122.13	116.44
22	B	832	CLA	O2A-CGA-CBA	2.17	118.70	111.91
22	5	310	CLA	O2A-CGA-CBA	2.17	118.70	111.91
39	8	301	CHL	CHB-C4A-NA	2.17	127.51	124.51
22	F	302	CLA	CMD-C2D-C3D	-2.17	122.63	127.61
22	A	802	CLA	O2A-CGA-CBA	2.17	118.70	111.91
22	A	831	CLA	CHA-C1A-NA	-2.17	121.44	126.40
22	B	829	CLA	CHA-C1A-NA	-2.17	121.44	126.40
22	7	316	CLA	CAA-C2A-C3A	-2.17	108.85	114.26
39	8	315	CHL	CMB-C2B-C1B	-2.17	125.14	128.46
22	4	808	CLA	O2A-CGA-CBA	2.17	118.70	111.91
22	8	309	CLA	O2D-CGD-O1D	-2.17	119.61	123.84
22	5	318	CLA	O2A-CGA-CBA	2.16	118.70	111.91
22	Z	308	CLA	CHA-C1A-NA	-2.16	121.44	126.40
22	B	838	CLA	O2D-CGD-O1D	-2.16	119.61	123.84
22	9	311	CLA	CMD-C2D-C3D	-2.16	122.64	127.61
22	A	805	CLA	C3D-C2D-C1D	-2.16	102.88	105.83
22	4	812	CLA	O2A-CGA-CBA	2.16	118.70	111.91
22	J	103	CLA	CHA-C1A-NA	-2.16	121.44	126.40
38	3	302	LUT	C11-C12-C13	-2.16	120.34	126.42
22	5	326	CLA	CMD-C2D-C3D	-2.16	122.64	127.61
22	B	812	CLA	CMD-C2D-C3D	-2.16	122.64	127.61
22	5	320	CLA	C3D-C2D-C1D	-2.16	102.88	105.83
22	A	808	CLA	CMD-C2D-C3D	-2.16	122.64	127.61
39	7	313	CHL	C1-C2-C3	-2.16	122.31	126.04
22	F	301	CLA	O2A-CGA-CBA	2.16	118.69	111.91
22	K	204	CLA	CHA-C1A-NA	-2.16	121.45	126.40
22	9	307	CLA	O2A-CGA-CBA	2.16	118.69	111.91
22	1	314	CLA	CMD-C2D-C3D	-2.16	122.64	127.61
22	5	311	CLA	CMD-C2D-C3D	-2.16	122.64	127.61
22	4	818	CLA	O2D-CGD-O1D	-2.16	119.61	123.84
22	1	304	CLA	CMB-C2B-C3B	2.16	128.72	124.68
39	5	321	CHL	C4A-NA-C1A	2.16	107.68	106.71
22	7	323	CLA	CAA-CBA-CGA	-2.16	106.94	113.25

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Mol	Chain	Res	Type	Atoms	Z	Observed(^{o})	Ideal(^{o})
22	8	313	CLA	CMD-C2D-C3D	-2.16	122.65	127.61
22	A	830	CLA	O2A-CGA-CBA	2.16	118.68	111.91
39	6	316	CHL	CHD-C4C-C3C	2.16	128.01	124.84
22	A	809	CLA	CHA-C1A-NA	-2.16	121.45	126.40
22	7	316	CLA	CHA-C1A-NA	-2.16	121.45	126.40
39	4	813	CHL	C2C-C3C-C4C	2.16	108.03	106.49
22	2	301	CLA	O2A-CGA-CBA	2.16	118.68	111.91
22	7	310	CLA	CHA-C1A-NA	-2.16	121.46	126.40
39	8	312	CHL	CMB-C2B-C1B	-2.16	125.15	128.46
25	1	317	LHG	O7-C7-O9	-2.16	118.49	123.70
22	B	823	CLA	CAA-CBA-CGA	-2.16	106.95	113.25
22	A	815	CLA	CMB-C2B-C3B	2.16	128.71	124.68
38	9	302	LUT	C11-C12-C13	-2.15	120.36	126.42
22	4	818	CLA	CHA-C1A-NA	-2.15	121.47	126.40
22	A	813	CLA	CHA-C1A-NA	-2.15	121.47	126.40
22	9	303	CLA	C1-O2A-CGA	2.15	122.09	116.44
22	B	810	CLA	CHA-C1A-NA	-2.15	121.47	126.40
22	A	854	CLA	CMD-C2D-C3D	-2.15	122.66	127.61
22	B	835	CLA	O2A-CGA-CBA	2.15	118.66	111.91
22	A	814	CLA	CMD-C2D-C3D	-2.15	122.66	127.61
22	B	837	CLA	CMD-C2D-C3D	-2.15	122.66	127.61
24	3	307	BCR	C21-C20-C19	2.15	129.93	123.22
22	B	820	CLA	O2A-CGA-CBA	2.15	118.66	111.91
22	3	322	CLA	CHA-C1A-NA	-2.15	121.47	126.40
22	5	322	CLA	CHA-C1A-NA	-2.15	121.47	126.40
22	5	318	CLA	CMD-C2D-C3D	-2.15	122.67	127.61
22	B	828	CLA	O2A-CGA-CBA	2.15	118.66	111.91
22	Z	315	CLA	CHA-C1A-NA	-2.15	121.47	126.40
22	7	305	CLA	CMD-C2D-C3D	-2.15	122.67	127.61
22	1	310	CLA	O2A-CGA-CBA	2.15	118.66	111.91
22	B	839	CLA	O2D-CGD-O1D	-2.15	119.63	123.84
34	F	308	LMG	O6-C1-C2	-2.15	105.80	110.35
22	7	317	CLA	O2A-CGA-CBA	2.15	118.65	111.91
39	5	317	CHL	CMB-C2B-C1B	-2.15	125.16	128.46
39	4	813	CHL	CHC-C1C-NC	2.15	127.46	124.20
22	B	834	CLA	CHA-C1A-NA	-2.15	121.48	126.40
22	5	322	CLA	O2D-CGD-O1D	-2.15	119.64	123.84
22	B	808	CLA	CMB-C2B-C3B	2.15	128.70	124.68
22	6	312	CLA	CHA-C1A-NA	-2.15	121.48	126.40
38	5	303	LUT	C19-C9-C10	-2.15	119.91	122.92
22	A	827	CLA	O2A-CGA-CBA	2.15	118.64	111.91
22	9	311	CLA	O2D-CGD-O1D	-2.15	119.64	123.84

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Mol	Chain	Res	Type	Atoms	Z	Observed(^{o})	Ideal(^{o})
24	6	305	BCR	C34-C9-C10	-2.15	119.92	122.92
24	B	847	BCR	C21-C20-C19	-2.15	116.52	123.22
22	A	836	CLA	CHA-C1A-NA	-2.15	121.48	126.40
38	7	301	LUT	C40-C33-C34	-2.15	119.92	122.92
22	A	825	CLA	CHA-C1A-NA	-2.15	121.48	126.40
22	3	312	CLA	CHA-C1A-NA	-2.15	121.48	126.40
22	9	305	CLA	O1D-CGD-CBD	-2.15	120.09	124.48
24	K	206	BCR	C35-C13-C14	-2.15	119.92	122.92
22	A	806	CLA	CMD-C2D-C3D	-2.14	122.68	127.61
22	7	311	CLA	CMD-C2D-C3D	-2.14	122.68	127.61
22	3	319	CLA	CHA-C1A-NA	-2.14	121.49	126.40
24	B	845	BCR	C33-C5-C6	-2.14	122.12	124.53
39	6	320	CHL	CMB-C2B-C1B	-2.14	125.17	128.46
22	5	313	CLA	OBD-CAD-C3D	-2.14	123.36	128.52
22	F	301	CLA	O1D-CGD-CBD	-2.14	120.10	124.48
22	B	824	CLA	CHA-C1A-NA	-2.14	121.49	126.40
22	B	823	CLA	O2D-CGD-O1D	-2.14	119.65	123.84
39	4	814	CHL	CHD-C4C-C3C	2.14	127.99	124.84
22	4	812	CLA	CHA-C1A-NA	-2.14	121.49	126.40
22	B	806	CLA	CHA-C1A-NA	-2.14	121.49	126.40
22	8	314	CLA	CHA-C1A-NA	-2.14	121.49	126.40
39	9	314	CHL	C4A-NA-C1A	2.14	107.67	106.71
22	A	824	CLA	CMD-C2D-C3D	-2.14	122.69	127.61
39	8	301	CHL	CHD-C4C-C3C	2.14	127.99	124.84
22	B	824	CLA	C3D-C2D-C1D	-2.14	102.91	105.83
22	5	308	CLA	C3D-C2D-C1D	-2.14	102.91	105.83
22	B	839	CLA	CHA-C1A-NA	-2.14	121.50	126.40
34	J	101	LMG	O8-C28-C29	2.14	118.62	111.91
39	2	303	CHL	CHB-C4A-NA	2.14	127.47	124.51
22	A	827	CLA	CHA-C1A-NA	-2.14	121.50	126.40
39	6	318	CHL	CMA-C3A-C2A	2.14	122.45	113.83
22	4	806	CLA	CMD-C2D-C3D	-2.14	122.70	127.61
22	Z	307	CLA	CAA-CBA-CGA	-2.14	107.01	113.25
22	A	810	CLA	O2A-CGA-CBA	2.14	118.61	111.91
38	1	303	LUT	C19-C9-C10	-2.14	119.93	122.92
22	A	840	CLA	CHA-C1A-NA	-2.14	121.51	126.40
22	4	815	CLA	CMB-C2B-C3B	2.13	128.67	124.68
22	5	326	CLA	C3D-C2D-C1D	-2.13	102.92	105.83
24	K	206	BCR	C36-C18-C17	-2.13	119.93	122.92
22	5	319	CLA	CMD-C2D-C3D	-2.13	122.70	127.61
22	7	306	CLA	O2A-CGA-CBA	2.13	118.61	111.91
22	B	840	CLA	CHA-C1A-NA	-2.13	121.51	126.40

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Mol	Chain	Res	Type	Atoms	Z	Observed(^{o})	Ideal(^{o})
39	9	314	CHL	CHC-C1C-NC	2.13	127.44	124.20
38	1	303	LUT	C8-C7-C6	-2.13	121.21	127.20
22	A	835	CLA	CHA-C1A-NA	-2.13	121.51	126.40
22	A	833	CLA	CHA-C1A-NA	-2.13	121.52	126.40
22	1	305	CLA	OBD-CAD-C3D	-2.13	123.39	128.52
39	2	303	CHL	CHD-C4C-C3C	2.13	127.97	124.84
22	3	311	CLA	CMB-C2B-C3B	2.13	128.66	124.68
22	7	317	CLA	CHA-C1A-NA	-2.13	121.52	126.40
39	9	314	CHL	CMB-C2B-C1B	-2.13	125.19	128.46
22	6	309	CLA	CMB-C2B-C3B	2.13	128.66	124.68
22	3	311	CLA	CHA-C1A-NA	-2.13	121.52	126.40
22	7	315	CLA	O2A-CGA-CBA	2.13	118.59	111.91
38	3	302	LUT	C21-C26-C27	-2.13	110.01	112.70
39	5	321	CHL	CMB-C2B-C1B	-2.13	125.19	128.46
22	A	803	CLA	O1D-CGD-CBD	-2.13	120.13	124.48
22	A	821	CLA	CHA-C1A-NA	-2.13	121.53	126.40
22	B	836	CLA	CHA-C1A-NA	-2.13	121.53	126.40
39	9	312	CHL	CMA-C3A-C2A	2.13	122.41	113.83
22	B	828	CLA	C3D-C2D-C1D	-2.13	102.93	105.83
22	B	807	CLA	CMD-C2D-C3D	-2.12	122.73	127.61
22	7	322	CLA	CMD-C2D-C3D	-2.12	122.73	127.61
24	4	804	BCR	C1-C6-C5	-2.12	119.62	122.61
22	6	317	CLA	CMB-C2B-C3B	2.12	128.65	124.68
22	3	311	CLA	CMD-C2D-C3D	-2.12	122.73	127.61
22	A	829	CLA	CHA-C1A-NA	-2.12	121.53	126.40
22	A	854	CLA	CHA-C1A-NA	-2.12	121.53	126.40
22	B	820	CLA	CHA-C1A-NA	-2.12	121.53	126.40
39	4	819	CHL	CMB-C2B-C1B	-2.12	125.20	128.46
22	9	304	CLA	CHD-C1D-ND	-2.12	122.50	124.45
24	5	304	BCR	C21-C20-C19	-2.12	116.59	123.22
24	A	845	BCR	C34-C9-C10	-2.12	119.95	122.92
22	A	825	CLA	O2A-CGA-CBA	2.12	118.57	111.91
39	Z	312	CHL	CMA-C3A-C2A	2.12	122.39	113.83
36	5	306	C7Z	C20-C13-C14	-2.12	119.95	122.92
38	Z	301	LUT	C8-C7-C6	-2.12	121.24	127.20
22	F	302	CLA	CHA-C1A-NA	-2.12	121.54	126.40
22	3	309	CLA	CHA-C1A-NA	-2.12	121.54	126.40
24	J	104	BCR	C30-C25-C24	2.12	121.78	115.78
39	8	312	CHL	CHB-C4A-NA	2.12	127.45	124.51
22	6	319	CLA	CHA-C1A-NA	-2.12	121.54	126.40
24	3	305	BCR	C23-C24-C25	-2.12	121.24	127.20
38	3	302	LUT	C22-C23-C24	2.12	114.16	111.74

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Mol	Chain	Res	Type	Atoms	Z	Observed(^{o})	Ideal(^{o})
22	7	308	CLA	CHA-C1A-NA	-2.12	121.54	126.40
39	Z	311	CHL	CMB-C2B-C1B	-2.12	125.20	128.46
22	1	315	CLA	CMB-C2B-C3B	2.12	128.65	124.68
22	3	308	CLA	O2D-CGD-O1D	-2.12	119.69	123.84
22	9	310	CLA	CHD-C1D-ND	-2.12	122.51	124.45
38	7	302	LUT	C15-C35-C34	-2.12	119.13	123.47
22	3	316	CLA	CHA-C1A-NA	-2.12	121.55	126.40
22	1	315	CLA	CHA-C1A-NA	-2.12	121.55	126.40
22	A	855	CLA	CHA-C1A-NA	-2.12	121.55	126.40
22	J	103	CLA	CAA-C2A-C3A	-2.12	108.97	114.26
22	3	310	CLA	CAA-CBA-CGA	-2.12	107.06	113.25
25	B	803	LHG	O7-C7-O9	-2.12	118.58	123.70
39	5	316	CHL	CMB-C2B-C1B	-2.12	125.21	128.46
22	4	807	CLA	CMD-C2D-C3D	-2.12	122.74	127.61
22	G	201	CLA	C3D-C2D-C1D	-2.12	102.94	105.83
39	7	313	CHL	CHB-C4A-NA	2.12	127.44	124.51
22	6	301	CLA	CHA-C1A-NA	-2.12	121.55	126.40
22	3	309	CLA	CMB-C2B-C3B	2.12	128.64	124.68
22	B	816	CLA	CHA-C1A-NA	-2.12	121.55	126.40
22	A	815	CLA	O2A-CGA-CBA	2.11	118.54	111.91
22	7	304	CLA	O1D-CGD-CBD	-2.11	120.16	124.48
39	Z	312	CHL	C2C-C3C-C4C	2.11	108.00	106.49
22	3	313	CLA	C3D-C2D-C1D	-2.11	102.95	105.83
22	Z	303	CLA	CMB-C2B-C3B	2.11	128.63	124.68
22	7	317	CLA	O2D-CGD-O1D	-2.11	119.71	123.84
22	A	842	CLA	CHA-C1A-NA	-2.11	121.56	126.40
22	7	322	CLA	O2A-CGA-CBA	2.11	118.54	111.91
22	K	202	CLA	CHA-C1A-NA	-2.11	121.56	126.40
22	9	306	CLA	CHA-C1A-NA	-2.11	121.56	126.40
22	B	801	CLA	CMB-C2B-C3B	2.11	128.63	124.68
24	K	206	BCR	C29-C30-C25	2.11	113.73	110.48
22	8	306	CLA	O1D-CGD-CBD	-2.11	120.16	124.48
39	6	315	CHL	CHD-C4C-C3C	2.11	127.94	124.84
24	G	203	BCR	C36-C18-C17	-2.11	119.97	122.92
39	5	317	CHL	C4A-NA-C1A	2.11	107.66	106.71
22	6	321	CLA	CHA-C1A-NA	-2.11	121.56	126.40
22	8	310	CLA	O2A-CGA-CBA	2.11	118.53	111.91
22	7	324	CLA	CHA-C1A-NA	-2.11	121.57	126.40
22	1	304	CLA	CHA-C1A-NA	-2.11	121.57	126.40
38	5	303	LUT	C8-C7-C6	-2.11	121.28	127.20
22	5	311	CLA	C3D-C2D-C1D	-2.11	102.95	105.83
39	3	317	CHL	CMB-C2B-C1B	-2.11	125.22	128.46

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Mol	Chain	Res	Type	Atoms	Z	Observed(^{o})	Ideal(^{o})
22	B	812	CLA	CHA-C1A-NA	-2.11	121.57	126.40
22	4	810	CLA	CHA-C1A-NA	-2.11	121.57	126.40
22	A	832	CLA	O2A-CGA-CBA	2.11	118.52	111.91
22	B	810	CLA	CAA-CBA-CGA	-2.11	107.09	113.25
22	7	306	CLA	CMD-C2D-C3D	-2.11	122.76	127.61
22	F	304	CLA	C1-O2A-CGA	2.11	121.97	116.44
22	B	811	CLA	CHA-C1A-NA	-2.11	121.57	126.40
22	9	308	CLA	CHA-C1A-NA	-2.11	121.57	126.40
22	5	315	CLA	O2A-CGA-CBA	2.11	118.52	111.91
24	3	304	BCR	C20-C21-C22	-2.11	124.31	127.31
22	4	807	CLA	CAA-C2A-C3A	-2.11	107.01	112.78
22	A	816	CLA	O1D-CGD-CBD	-2.11	120.18	124.48
22	1	313	CLA	C3D-C2D-C1D	-2.11	102.96	105.83
22	K	205	CLA	C1C-C2C-C3C	-2.11	104.74	106.96
22	4	815	CLA	O2A-CGA-CBA	2.10	118.51	111.91
22	6	302	CLA	CHA-C1A-NA	-2.10	121.58	126.40
22	Z	310	CLA	O2D-CGD-O1D	-2.10	119.72	123.84
38	5	303	LUT	C15-C35-C34	-2.10	119.16	123.47
22	9	307	CLA	CHA-C1A-NA	-2.10	121.58	126.40
22	1	316	CLA	CMA-C3A-C4A	2.10	117.43	111.77
22	1	314	CLA	O2A-CGA-CBA	2.10	118.51	111.91
24	A	856	BCR	C10-C11-C12	-2.10	116.65	123.22
22	B	833	CLA	CHA-C1A-NA	-2.10	121.58	126.40
24	5	304	BCR	C34-C9-C10	-2.10	119.98	122.92
22	7	309	CLA	C3D-C2D-C1D	-2.10	102.96	105.83
22	3	318	CLA	C6-C7-C8	-2.10	109.12	115.92
39	Z	312	CHL	CHD-C4C-C3C	2.10	127.93	124.84
22	B	819	CLA	CHA-C1A-NA	-2.10	121.58	126.40
39	8	301	CHL	CHC-C1C-NC	2.10	127.39	124.20
22	7	311	CLA	CHA-C1A-NA	-2.10	121.58	126.40
22	5	308	CLA	CHA-C1A-NA	-2.10	121.58	126.40
22	7	309	CLA	CAA-CBA-CGA	-2.10	107.11	113.25
31	1	319	DGD	O1G-C1A-C2A	2.10	118.50	111.91
22	B	801	CLA	CMD-C2D-C3D	-2.10	122.78	127.61
22	B	822	CLA	CMD-C2D-C3D	-2.10	122.78	127.61
22	B	807	CLA	CHA-C1A-NA	-2.10	121.59	126.40
22	9	313	CLA	CHA-C1A-NA	-2.10	121.59	126.40
38	6	303	LUT	C31-C30-C29	-2.10	124.31	127.31
39	2	303	CHL	CMB-C2B-C1B	-2.10	125.24	128.46
38	Z	301	LUT	C20-C13-C14	-2.10	119.98	122.92
22	7	314	CLA	O1D-CGD-CBD	-2.10	120.19	124.48
22	5	319	CLA	CHA-C1A-NA	-2.10	121.59	126.40

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Mol	Chain	Res	Type	Atoms	Z	Observed(^{o})	Ideal(^{o})
39	6	316	CHL	CMB-C2B-C1B	-2.10	125.24	128.46
22	1	313	CLA	CHA-C1A-NA	-2.10	121.59	126.40
36	J	105	C7Z	C40-C33-C34	-2.10	119.98	122.92
22	A	802	CLA	CHA-C1A-NA	-2.10	121.59	126.40
22	A	819	CLA	CMB-C2B-C3B	2.10	128.60	124.68
22	3	315	CLA	OBD-CAD-C3D	-2.10	123.47	128.52
22	B	815	CLA	CHA-C1A-NA	-2.10	121.59	126.40
22	A	807	CLA	CHA-C1A-NA	-2.10	121.60	126.40
22	8	305	CLA	O2A-CGA-CBA	2.10	118.49	111.91
22	L	203	CLA	CMB-C2B-C3B	2.10	128.60	124.68
22	F	302	CLA	C3D-C2D-C1D	-2.10	102.97	105.83
22	5	322	CLA	CMD-C2D-C3D	-2.10	122.79	127.61
22	A	824	CLA	CHA-C1A-NA	-2.10	121.60	126.40
22	Z	303	CLA	C3D-C2D-C1D	-2.09	102.97	105.83
22	7	322	CLA	C3D-C2D-C1D	-2.09	102.97	105.83
24	J	104	BCR	C1-C6-C7	2.09	121.70	115.78
22	4	812	CLA	C3D-C2D-C1D	-2.09	102.97	105.83
22	B	805	CLA	CHA-C1A-NA	-2.09	121.60	126.40
22	7	307	CLA	CMB-C2B-C3B	2.09	128.59	124.68
22	B	841	CLA	CMD-C2D-C3D	-2.09	122.80	127.61
22	1	313	CLA	O2A-CGA-CBA	2.09	118.48	111.91
24	5	305	BCR	C12-C13-C14	2.09	122.15	118.94
22	1	305	CLA	C3D-C2D-C1D	-2.09	102.97	105.83
39	8	312	CHL	CMA-C3A-C2A	2.09	122.27	113.83
22	A	814	CLA	CMB-C2B-C3B	2.09	128.59	124.68
22	B	813	CLA	CHA-C1A-NA	-2.09	121.61	126.40
38	7	302	LUT	C20-C13-C14	-2.09	119.99	122.92
22	5	312	CLA	O2A-CGA-CBA	2.09	118.47	111.91
22	A	818	CLA	C3D-C2D-C1D	-2.09	102.98	105.83
22	9	309	CLA	CHA-C1A-NA	-2.09	121.61	126.40
39	1	312	CHL	CMA-C3A-C2A	2.09	122.26	113.83
22	B	822	CLA	CHA-C1A-NA	-2.09	121.61	126.40
22	B	821	CLA	CHA-C1A-NA	-2.09	121.61	126.40
22	B	832	CLA	O1D-CGD-CBD	-2.09	120.21	124.48
39	7	313	CHL	CMB-C2B-C1B	-2.09	125.25	128.46
22	A	810	CLA	CHA-C1A-NA	-2.09	121.61	126.40
22	A	811	CLA	C3D-C2D-C1D	-2.09	102.98	105.83
22	A	837	CLA	CHA-C1A-NA	-2.09	121.62	126.40
22	9	305	CLA	C3D-C2D-C1D	-2.09	102.98	105.83
23	A	843	PQN	C16-C17-C18	-2.09	109.17	115.92
22	7	315	CLA	CHA-C1A-NA	-2.09	121.62	126.40
22	4	809	CLA	C3D-C2D-C1D	-2.09	102.98	105.83

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Mol	Chain	Res	Type	Atoms	Z	Observed(^{o})	Ideal(^{o})
22	5	309	CLA	C3D-C2D-C1D	-2.09	102.98	105.83
22	6	307	CLA	CHA-C1A-NA	-2.09	121.62	126.40
39	5	321	CHL	CHD-C4C-C3C	2.09	127.91	124.84
24	8	304	BCR	C2-C1-C6	2.09	113.69	110.48
22	F	301	CLA	CHA-C1A-NA	-2.09	121.62	126.40
22	5	310	CLA	O1D-CGD-CBD	-2.09	120.22	124.48
39	8	312	CHL	CHD-C4C-C3C	2.09	127.91	124.84
39	6	318	CHL	CHB-C4A-NA	2.09	127.40	124.51
39	6	315	CHL	CHC-C1C-NC	2.09	127.37	124.20
22	5	307	CLA	O2A-CGA-CBA	2.09	118.45	111.91
22	A	804	CLA	CMB-C2B-C3B	2.09	128.58	124.68
22	9	313	CLA	CMD-C2D-C3D	-2.08	122.82	127.61
22	4	807	CLA	C3D-C2D-C1D	-2.08	102.99	105.83
22	B	842	CLA	O2A-CGA-CBA	2.08	118.45	111.91
39	4	814	CHL	C4A-NA-C1A	2.08	107.64	106.71
38	8	303	LUT	C30-C31-C32	-2.08	116.71	123.22
22	7	309	CLA	CMB-C2B-C3B	2.08	128.58	124.68
22	Z	313	CLA	C6-C5-C3	-2.08	107.99	113.45
22	3	309	CLA	C3D-C2D-C1D	-2.08	102.99	105.83
39	6	315	CHL	CMA-C3A-C2A	2.08	122.23	113.83
22	A	803	CLA	CHA-C1A-NA	-2.08	121.63	126.40
24	8	304	BCR	C34-C9-C10	-2.08	120.01	122.92
22	7	311	CLA	C3D-C2D-C1D	-2.08	102.99	105.83
38	7	302	LUT	C8-C7-C6	-2.08	121.36	127.20
22	A	834	CLA	CHA-C1A-NA	-2.08	121.63	126.40
22	B	835	CLA	CHA-C1A-NA	-2.08	121.63	126.40
22	A	832	CLA	CMB-C2B-C3B	2.08	128.57	124.68
22	A	815	CLA	CHA-C1A-NA	-2.08	121.63	126.40
25	4	820	LHG	O7-C7-O9	-2.08	118.67	123.70
22	6	322	CLA	CHA-C1A-NA	-2.08	121.63	126.40
22	B	826	CLA	CHA-C1A-NA	-2.08	121.64	126.40
22	A	821	CLA	CMB-C2B-C3B	2.08	128.57	124.68
22	B	836	CLA	C3D-C2D-C1D	-2.08	102.99	105.83
22	Z	316	CLA	CHA-C1A-NA	-2.08	121.64	126.40
38	8	302	LUT	C38-C25-C24	-2.08	119.11	123.56
39	4	813	CHL	CMA-C3A-C2A	2.08	122.21	113.83
22	A	841	CLA	CHA-C1A-NA	-2.08	121.64	126.40
22	L	201	CLA	CHA-C1A-NA	-2.08	121.64	126.40
22	4	817	CLA	CHA-C1A-NA	-2.08	121.64	126.40
22	K	204	CLA	O2D-CGD-O1D	-2.08	119.77	123.84
22	7	307	CLA	O1D-CGD-CBD	-2.08	120.23	124.48
22	Z	314	CLA	O2A-CGA-CBA	2.08	118.43	111.91

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Mol	Chain	Res	Type	Atoms	Z	Observed(^{o})	Ideal(^{o})
22	A	813	CLA	O1D-CGD-CBD	-2.08	120.23	124.48
22	A	825	CLA	CMB-C2B-C3B	2.08	128.56	124.68
22	8	313	CLA	O2A-CGA-CBA	2.08	118.42	111.91
38	5	302	LUT	C38-C25-C24	-2.08	119.11	123.56
22	9	308	CLA	O1D-CGD-CBD	-2.08	120.24	124.48
22	A	808	CLA	O2A-CGA-CBA	2.08	118.42	111.91
22	B	815	CLA	CAC-C3C-C4C	2.08	127.50	124.81
24	A	856	BCR	C36-C18-C17	-2.08	120.02	122.92
22	5	307	CLA	CHA-C1A-NA	-2.07	121.65	126.40
22	A	855	CLA	O2A-CGA-CBA	2.07	118.42	111.91
22	B	837	CLA	O2A-CGA-CBA	2.07	118.42	111.91
22	Z	313	CLA	O2A-CGA-CBA	2.07	118.41	111.91
22	L	202	CLA	CHA-C1A-NA	-2.07	121.65	126.40
22	5	323	CLA	C3D-C2D-C1D	-2.07	103.00	105.83
22	B	810	CLA	C6-C7-C8	-2.07	109.22	115.92
24	5	305	BCR	C7-C8-C9	-2.07	123.10	126.23
22	5	301	CLA	CHA-C1A-NA	-2.07	121.65	126.40
22	7	309	CLA	CHA-C1A-NA	-2.07	121.65	126.40
22	B	805	CLA	CMD-C2D-C3D	-2.07	122.85	127.61
22	Z	314	CLA	CHA-C1A-NA	-2.07	121.66	126.40
22	A	833	CLA	C3D-C2D-C1D	-2.07	103.00	105.83
22	1	305	CLA	CHA-C1A-NA	-2.07	121.66	126.40
22	5	312	CLA	O2D-CGD-O1D	-2.07	119.79	123.84
22	B	819	CLA	C3D-C2D-C1D	-2.07	103.01	105.83
22	4	811	CLA	CMD-C2D-C3D	-2.07	122.85	127.61
22	A	830	CLA	CHA-C1A-NA	-2.07	121.66	126.40
22	B	841	CLA	CHA-C1A-NA	-2.07	121.66	126.40
39	5	317	CHL	CHB-C4A-NA	2.07	127.37	124.51
39	9	312	CHL	CMB-C2B-C1B	-2.07	125.28	128.46
22	B	829	CLA	C6-C5-C3	-2.07	108.03	113.45
38	6	303	LUT	C39-C29-C28	2.07	121.34	118.08
22	Z	309	CLA	CHA-C1A-NA	-2.07	121.66	126.40
39	3	317	CHL	CHD-C4C-C3C	2.07	127.88	124.84
22	8	309	CLA	CHA-C1A-NA	-2.07	121.66	126.40
22	A	810	CLA	CMB-C2B-C3B	2.07	128.55	124.68
39	4	819	CHL	CHB-C4A-NA	2.07	127.37	124.51
22	4	806	CLA	C3D-C2D-C1D	-2.07	103.01	105.83
22	B	831	CLA	CHA-C1A-NA	-2.07	121.66	126.40
22	3	308	CLA	CMB-C2B-C3B	2.07	128.54	124.68
22	B	810	CLA	O1D-CGD-CBD	-2.07	120.26	124.48
22	5	310	CLA	CHA-C1A-NA	-2.07	121.67	126.40
24	F	303	BCR	C36-C18-C17	-2.07	120.03	122.92

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Mol	Chain	Res	Type	Atoms	Z	Observed(^{o})	Ideal(^{o})
22	6	310	CLA	CHA-C1A-NA	-2.06	121.67	126.40
22	A	804	CLA	O1D-CGD-CBD	-2.06	120.26	124.48
22	8	310	CLA	CHA-C1A-NA	-2.06	121.67	126.40
33	G	204	LMT	C1'-O5'-C5'	-2.06	109.64	113.69
22	7	312	CLA	C3D-C2D-C1D	-2.06	103.01	105.83
22	B	805	CLA	O1D-CGD-CBD	-2.06	120.26	124.48
22	B	811	CLA	O1D-CGD-CBD	-2.06	120.26	124.48
22	9	310	CLA	OBD-CAD-C3D	-2.06	123.56	128.52
22	8	313	CLA	CHA-C1A-NA	-2.06	121.67	126.40
22	9	313	CLA	O1D-CGD-CBD	-2.06	120.26	124.48
22	7	305	CLA	CHA-C1A-NA	-2.06	121.67	126.40
22	7	304	CLA	C3D-C2D-C1D	-2.06	103.02	105.83
23	A	843	PQN	C2M-C2-C1	2.06	119.69	116.27
22	9	305	CLA	CMD-C2D-C3D	-2.06	122.87	127.61
22	B	817	CLA	O1D-CGD-CBD	-2.06	120.27	124.48
22	8	311	CLA	CMD-C2D-C3D	-2.06	122.87	127.61
22	A	835	CLA	CMB-C2B-C3B	2.06	128.53	124.68
38	6	304	LUT	C20-C13-C14	-2.06	120.04	122.92
22	1	307	CLA	CHA-C1A-NA	-2.06	121.68	126.40
22	B	813	CLA	C3D-C2D-C1D	-2.06	103.02	105.83
22	4	810	CLA	CAA-CBA-CGA	-2.06	107.23	113.25
22	B	830	CLA	O2D-CGD-O1D	-2.06	119.81	123.84
22	F	301	CLA	C3D-C2D-C1D	-2.06	103.02	105.83
22	K	204	CLA	C3D-C2D-C1D	-2.06	103.02	105.83
22	8	309	CLA	C3D-C2D-C1D	-2.06	103.02	105.83
22	A	826	CLA	CHA-C1A-NA	-2.06	121.68	126.40
22	5	319	CLA	C3D-C2D-C1D	-2.06	103.02	105.83
22	A	827	CLA	O1D-CGD-CBD	-2.06	120.27	124.48
22	B	823	CLA	CHA-C1A-NA	-2.06	121.69	126.40
22	B	837	CLA	CHA-C1A-NA	-2.06	121.69	126.40
22	1	306	CLA	CHA-C1A-NA	-2.06	121.69	126.40
22	1	314	CLA	CHA-C1A-NA	-2.06	121.69	126.40
24	A	846	BCR	C11-C12-C13	-2.06	120.64	126.42
22	7	324	CLA	CMB-C2B-C3B	2.06	128.53	124.68
22	6	307	CLA	CMB-C2B-C3B	2.06	128.53	124.68
22	3	313	CLA	O2A-CGA-CBA	2.06	118.36	111.91
22	Z	313	CLA	O1D-CGD-CBD	-2.06	120.27	124.48
22	5	312	CLA	CHA-C1A-NA	-2.06	121.69	126.40
22	5	318	CLA	CHA-C1A-NA	-2.06	121.69	126.40
22	3	311	CLA	O1D-CGD-CBD	-2.06	120.28	124.48
22	5	308	CLA	O1D-CGD-CBD	-2.06	120.28	124.48
22	4	808	CLA	CHA-C1A-NA	-2.06	121.69	126.40

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Mol	Chain	Res	Type	Atoms	Z	Observed(^{o})	Ideal(^{o})
22	B	810	CLA	C6-C5-C3	-2.06	108.06	113.45
22	G	201	CLA	O2A-CGA-CBA	2.06	118.36	111.91
22	Z	306	CLA	CHA-C1A-NA	-2.05	121.69	126.40
22	1	314	CLA	O2D-CGD-O1D	-2.05	119.82	123.84
22	A	839	CLA	CHA-C1A-NA	-2.05	121.69	126.40
22	A	829	CLA	CMD-C2D-C3D	-2.05	122.89	127.61
22	B	808	CLA	CHA-C1A-NA	-2.05	121.69	126.40
22	A	832	CLA	CHA-C1A-NA	-2.05	121.70	126.40
22	5	307	CLA	CMB-C2B-C3B	2.05	128.52	124.68
22	8	306	CLA	CHA-C1A-NA	-2.05	121.70	126.40
22	6	308	CLA	CMD-C2D-C3D	-2.05	122.89	127.61
22	A	812	CLA	C3D-C2D-C1D	-2.05	103.03	105.83
22	K	202	CLA	O1D-CGD-CBD	-2.05	120.29	124.48
22	B	826	CLA	C16-C15-C13	-2.05	109.29	115.92
22	A	838	CLA	CHA-C1A-NA	-2.05	121.70	126.40
22	A	829	CLA	O2D-CGD-O1D	-2.05	119.83	123.84
22	3	318	CLA	C11-C10-C8	-2.05	109.29	115.92
38	3	303	LUT	C40-C33-C32	2.05	121.31	118.08
39	4	813	CHL	CHB-C4A-NA	2.05	127.35	124.51
22	6	301	CLA	O1D-CGD-CBD	-2.05	120.29	124.48
22	B	816	CLA	CMB-C2B-C3B	2.05	128.51	124.68
22	7	314	CLA	CHA-C1A-NA	-2.05	121.71	126.40
22	8	316	CLA	CHA-C1A-NA	-2.05	121.71	126.40
22	B	831	CLA	CMB-C2B-C3B	2.05	128.51	124.68
22	A	817	CLA	CHA-C1A-NA	-2.05	121.71	126.40
22	B	827	CLA	CHA-C1A-NA	-2.05	121.71	126.40
33	G	204	LMT	C1B-O1B-C4'	-2.05	112.89	117.96
24	B	845	BCR	C2-C1-C6	2.05	113.63	110.48
22	A	836	CLA	C6-C5-C3	-2.05	108.09	113.45
22	B	834	CLA	C3D-C2D-C1D	-2.05	103.04	105.83
22	5	322	CLA	C3D-C2D-C1D	-2.05	103.04	105.83
22	Z	308	CLA	O2A-CGA-CBA	2.05	118.33	111.91
22	A	834	CLA	C3D-C2D-C1D	-2.05	103.04	105.83
22	L	201	CLA	C3D-C2D-C1D	-2.05	103.04	105.83
22	9	311	CLA	C3D-C2D-C1D	-2.05	103.04	105.83
22	6	317	CLA	CHA-C1A-NA	-2.05	121.71	126.40
22	7	323	CLA	C1-O2A-CGA	2.05	121.81	116.44
22	4	817	CLA	C3D-C2D-C1D	-2.05	103.04	105.83
22	K	203	CLA	C3D-C2D-C1D	-2.04	103.04	105.83
22	A	819	CLA	O2A-CGA-CBA	2.04	118.32	111.91
22	A	835	CLA	CMD-C2D-C3D	-2.04	122.91	127.61
22	B	820	CLA	O1D-CGD-CBD	-2.04	120.30	124.48

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Mol	Chain	Res	Type	Atoms	Z	Observed(^{o})	Ideal(^{o})
24	A	844	BCR	C21-C20-C19	-2.04	116.84	123.22
22	B	819	CLA	O1D-CGD-CBD	-2.04	120.30	124.48
22	Z	307	CLA	OBD-CAD-C3D	-2.04	123.60	128.52
22	9	309	CLA	C2A-C1A-CHA	2.04	127.43	123.86
24	A	856	BCR	C30-C25-C24	2.04	121.56	115.78
22	Z	310	CLA	C3D-C2D-C1D	-2.04	103.04	105.83
39	Z	311	CHL	CMA-C3A-C2A	2.04	122.07	113.83
22	3	310	CLA	C11-C10-C8	-2.04	109.32	115.92
22	5	314	CLA	CHA-C1A-NA	-2.04	121.72	126.40
39	6	318	CHL	CHD-C4C-C3C	2.04	127.84	124.84
24	A	848	BCR	C8-C7-C6	-2.04	121.47	127.20
22	Z	310	CLA	CMD-C2D-C3D	-2.04	122.92	127.61
22	8	314	CLA	CMD-C2D-C3D	-2.04	122.92	127.61
22	6	312	CLA	O1D-CGD-CBD	-2.04	120.31	124.48
22	4	808	CLA	C3D-C2D-C1D	-2.04	103.05	105.83
22	2	302	CLA	C3D-C2D-C1D	-2.04	103.05	105.83
22	A	824	CLA	O2A-CGA-CBA	2.04	118.31	111.91
22	F	305	CLA	CHA-C1A-NA	-2.04	121.73	126.40
22	Z	303	CLA	O2D-CGD-O1D	-2.04	119.85	123.84
22	G	202	CLA	CHA-C1A-NA	-2.04	121.73	126.40
22	2	304	CLA	CHA-C1A-NA	-2.04	121.73	126.40
24	J	104	BCR	C16-C15-C14	-2.04	119.30	123.47
38	8	303	LUT	C11-C12-C13	-2.04	120.69	126.42
39	Z	312	CHL	CHB-C4A-NA	2.04	127.33	124.51
22	A	823	CLA	C6-C5-C3	-2.04	108.11	113.45
22	6	313	CLA	CHA-C1A-NA	-2.04	121.73	126.40
22	6	314	CLA	CHA-C1A-NA	-2.04	121.73	126.40
22	3	314	CLA	CMD-C2D-C3D	-2.04	122.93	127.61
22	F	302	CLA	CAA-CBA-CGA	-2.04	107.30	113.25
22	7	310	CLA	O2A-CGA-CBA	2.04	118.30	111.91
39	Z	311	CHL	CHB-C4A-NA	2.04	127.33	124.51
39	6	316	CHL	CHB-C4A-NA	2.04	127.33	124.51
22	A	806	CLA	CHA-C1A-NA	-2.04	121.74	126.40
38	Z	302	LUT	C8-C7-C6	-2.04	121.49	127.20
24	3	304	BCR	C11-C12-C13	2.04	132.13	126.42
22	5	301	CLA	C3D-C2D-C1D	-2.03	103.05	105.83
38	5	302	LUT	C15-C14-C13	-2.03	124.41	127.31
22	B	815	CLA	O1D-CGD-CBD	-2.03	120.32	124.48
22	L	202	CLA	O2A-CGA-CBA	2.03	118.29	111.91
22	B	837	CLA	CMB-C2B-C3B	2.03	128.48	124.68
22	5	313	CLA	C3D-C2D-C1D	-2.03	103.06	105.83
22	A	809	CLA	C3D-C2D-C1D	-2.03	103.06	105.83

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Mol	Chain	Res	Type	Atoms	Z	Observed(^{o})	Ideal(^{o})
22	G	201	CLA	OBD-CAD-C3D	-2.03	123.63	128.52
22	B	833	CLA	C3D-C2D-C1D	-2.03	103.06	105.83
22	A	818	CLA	O1D-CGD-CBD	-2.03	120.33	124.48
22	8	308	CLA	CHA-C1A-NA	-2.03	121.74	126.40
22	B	823	CLA	O1D-CGD-CBD	-2.03	120.33	124.48
22	4	811	CLA	C3D-C2D-C1D	-2.03	103.06	105.83
39	6	318	CHL	C1-O2A-CGA	2.03	121.78	116.44
22	L	203	CLA	CHA-C1A-NA	-2.03	121.75	126.40
24	A	845	BCR	C29-C30-C25	2.03	113.61	110.48
22	G	202	CLA	C3D-C2D-C1D	-2.03	103.06	105.83
22	A	835	CLA	O1D-CGD-CBD	-2.03	120.33	124.48
22	B	818	CLA	O1D-CGD-CBD	-2.03	120.33	124.48
22	A	827	CLA	C1-O2A-CGA	2.03	121.77	116.44
22	1	316	CLA	CHD-C1D-ND	-2.03	122.59	124.45
24	B	849	BCR	C8-C7-C6	-2.03	121.50	127.20
22	Z	303	CLA	CHA-C1A-NA	-2.03	121.75	126.40
22	3	312	CLA	C3D-C2D-C1D	-2.03	103.06	105.83
22	2	304	CLA	O1D-CGD-CBD	-2.03	120.33	124.48
36	J	105	C7Z	C19-C9-C10	-2.03	120.08	122.92
22	A	837	CLA	O2A-CGA-CBA	2.03	118.27	111.91
22	B	810	CLA	C3D-C2D-C1D	-2.03	103.06	105.83
40	1	318	SQD	O8-S-C6	-2.03	102.51	105.74
38	Z	302	LUT	C20-C13-C14	-2.03	120.08	122.92
22	A	832	CLA	C3D-C2D-C1D	-2.03	103.06	105.83
22	B	817	CLA	CMB-C2B-C3B	2.03	128.47	124.68
22	G	202	CLA	CMB-C2B-C3B	2.03	128.47	124.68
22	B	830	CLA	CHA-C1A-NA	-2.03	121.76	126.40
22	A	808	CLA	C11-C12-C13	-2.03	109.37	115.92
22	9	306	CLA	O1D-CGD-CBD	-2.02	120.34	124.48
22	B	829	CLA	C3D-C2D-C1D	-2.02	103.07	105.83
22	B	841	CLA	C3D-C2D-C1D	-2.02	103.07	105.83
22	8	310	CLA	CAA-C2A-C3A	-2.02	107.23	112.78
22	A	841	CLA	C1-O2A-CGA	2.02	121.75	116.44
22	7	308	CLA	CMB-C2B-C3B	2.02	128.47	124.68
38	8	303	LUT	C20-C13-C12	2.02	121.27	118.08
38	7	302	LUT	C38-C25-C24	-2.02	119.23	123.56
22	B	805	CLA	C3D-C2D-C1D	-2.02	103.07	105.83
22	3	318	CLA	C3D-C2D-C1D	-2.02	103.07	105.83
22	7	316	CLA	C3D-C2D-C1D	-2.02	103.07	105.83
39	5	316	CHL	C1-O2A-CGA	2.02	121.75	116.44
22	1	316	CLA	C6-C7-C8	-2.02	109.38	115.92
22	Z	309	CLA	C6-C5-C3	-2.02	108.15	113.45

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Mol	Chain	Res	Type	Atoms	Z	Observed(^{o})	Ideal(^{o})
22	K	203	CLA	O2A-CGA-CBA	2.02	118.26	111.91
22	Z	305	CLA	O1D-CGD-CBD	-2.02	120.34	124.48
22	Z	304	CLA	CHA-C1A-NA	-2.02	121.77	126.40
22	A	840	CLA	CMB-C2B-C3B	2.02	128.46	124.68
22	B	811	CLA	O2A-CGA-CBA	2.02	118.25	111.91
22	6	308	CLA	CHA-C1A-NA	-2.02	121.77	126.40
32	F	306	RRX	C19-C18-C17	2.02	122.04	118.94
22	A	854	CLA	CMB-C2B-C3B	2.02	128.46	124.68
22	B	822	CLA	O1D-CGD-CBD	-2.02	120.35	124.48
22	K	204	CLA	O1D-CGD-CBD	-2.02	120.35	124.48
22	A	822	CLA	C3D-C2D-C1D	-2.02	103.07	105.83
22	6	310	CLA	O1D-CGD-CBD	-2.02	120.35	124.48
22	9	303	CLA	CMC-C2C-C1C	2.02	128.12	125.04
22	5	326	CLA	O1D-CGD-CBD	-2.02	120.35	124.48
22	7	310	CLA	C3D-C2D-C1D	-2.02	103.07	105.83
24	3	305	BCR	C29-C30-C25	2.02	113.59	110.48
22	5	318	CLA	O2D-CGD-O1D	-2.02	119.89	123.84
38	4	803	LUT	C40-C33-C34	-2.02	120.09	122.92
22	Z	308	CLA	C3D-C2D-C1D	-2.02	103.08	105.83
22	7	324	CLA	C3D-C2D-C1D	-2.02	103.08	105.83
22	B	825	CLA	C3D-C2D-C1D	-2.02	103.08	105.83
39	1	312	CHL	C2C-C3C-C4C	2.02	107.93	106.49
38	9	302	LUT	C35-C15-C14	-2.02	119.34	123.47
22	A	836	CLA	C3D-C2D-C1D	-2.02	103.08	105.83
22	7	323	CLA	C3D-C2D-C1D	-2.02	103.08	105.83
22	9	306	CLA	C3D-C2D-C1D	-2.02	103.08	105.83
22	4	818	CLA	C3D-C2D-C1D	-2.02	103.08	105.83
38	7	301	LUT	C15-C35-C34	-2.02	119.34	123.47
22	A	825	CLA	C3D-C2D-C1D	-2.02	103.08	105.83
22	6	310	CLA	C3D-C2D-C1D	-2.02	103.08	105.83
22	1	308	CLA	CMC-C2C-C1C	2.02	128.11	125.04
22	A	814	CLA	O1D-CGD-CBD	-2.02	120.36	124.48
22	A	819	CLA	CHA-C1A-NA	-2.02	121.78	126.40
34	F	308	LMG	O7-C8-C9	2.02	115.70	108.40
22	1	304	CLA	O2D-CGD-O1D	-2.02	119.90	123.84
22	A	831	CLA	O1D-CGD-CBD	-2.02	120.36	124.48
22	A	816	CLA	C3D-C2D-C1D	-2.02	103.08	105.83
22	5	314	CLA	CMD-C2D-C3D	-2.02	122.98	127.61
24	A	845	BCR	C20-C19-C18	-2.02	120.75	126.42
24	A	848	BCR	C35-C13-C14	-2.02	120.10	122.92
22	B	818	CLA	C3D-C2D-C1D	-2.02	103.08	105.83
22	J	103	CLA	C3D-C2D-C1D	-2.02	103.08	105.83

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Mol	Chain	Res	Type	Atoms	Z	Observed(^{o})	Ideal(^{o})
24	K	206	BCR	C2-C1-C6	2.02	113.58	110.48
22	7	316	CLA	CMB-C2B-C3B	2.01	128.45	124.68
22	K	202	CLA	C3D-C2D-C1D	-2.01	103.08	105.83
22	6	312	CLA	C3D-C2D-C1D	-2.01	103.08	105.83
22	8	307	CLA	C3D-C2D-C1D	-2.01	103.08	105.83
22	A	833	CLA	O1D-CGD-CBD	-2.01	120.36	124.48
22	3	308	CLA	CHA-C1A-NA	-2.01	121.79	126.40
22	1	306	CLA	C3D-C2D-C1D	-2.01	103.08	105.83
22	3	311	CLA	C3D-C2D-C1D	-2.01	103.08	105.83
22	9	308	CLA	C3D-C2D-C1D	-2.01	103.08	105.83
22	3	313	CLA	O2D-CGD-O1D	-2.01	119.90	123.84
24	A	845	BCR	C16-C15-C14	-2.01	119.35	123.47
22	7	306	CLA	C6-C5-C3	-2.01	108.18	113.45
22	1	316	CLA	C1D-ND-C4D	-2.01	104.91	106.33
22	B	830	CLA	CMD-C2D-C3D	-2.01	122.99	127.61
22	B	824	CLA	CMB-C2B-C3B	2.01	128.44	124.68
22	A	838	CLA	O1D-CGD-CBD	-2.01	120.37	124.48
22	8	305	CLA	C6-C5-C3	-2.01	108.18	113.45
38	4	803	LUT	C15-C35-C34	-2.01	119.36	123.47
22	3	314	CLA	CHA-C1A-NA	-2.01	121.79	126.40
22	A	814	CLA	C3D-C2D-C1D	-2.01	103.09	105.83
22	Z	314	CLA	CMD-C2D-C3D	-2.01	122.99	127.61
24	8	304	BCR	C15-C16-C17	-2.01	119.36	123.47
22	B	807	CLA	C3D-C2D-C1D	-2.01	103.09	105.83
22	Z	313	CLA	C3D-C2D-C1D	-2.01	103.09	105.83
22	8	308	CLA	C3D-C2D-C1D	-2.01	103.09	105.83
38	8	302	LUT	C30-C31-C32	-2.01	116.94	123.22
22	6	319	CLA	C3D-C2D-C1D	-2.01	103.09	105.83
22	L	202	CLA	O1D-CGD-CBD	-2.01	120.37	124.48
38	4	802	LUT	C38-C25-C24	-2.01	119.26	123.56
24	A	845	BCR	C35-C13-C14	-2.01	120.11	122.92
24	A	845	BCR	C15-C16-C17	-2.01	119.36	123.47
33	4	822	LMT	O5B-C5B-C4B	-2.01	106.05	109.69
22	5	313	CLA	CMD-C2D-C3D	-2.01	122.99	127.61
22	6	309	CLA	C2A-C1A-CHA	2.01	127.37	123.86
22	B	837	CLA	O1D-CGD-CBD	-2.01	120.38	124.48
24	4	804	BCR	C36-C18-C17	-2.01	120.11	122.92
22	3	313	CLA	CMB-C2B-C3B	2.01	128.44	124.68
22	6	313	CLA	CMD-C2D-C3D	-2.01	123.00	127.61
22	B	840	CLA	CMB-C2B-C3B	2.01	128.43	124.68
22	A	822	CLA	CMB-C2B-C3B	2.01	128.43	124.68
39	7	313	CHL	CHD-C4C-C3C	2.01	127.79	124.84

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Mol	Chain	Res	Type	Atoms	Z	Observed(^{o})	Ideal(^{o})
24	3	307	BCR	C30-C25-C24	2.01	121.45	115.78
22	4	810	CLA	O1D-CGD-CBD	-2.01	120.38	124.48
22	A	805	CLA	O1D-CGD-CBD	-2.01	120.38	124.48
22	B	808	CLA	C3D-C2D-C1D	-2.01	103.09	105.83
22	5	315	CLA	C6-C5-C3	-2.01	108.20	113.45
22	B	817	CLA	CMD-C2D-C3D	-2.01	123.00	127.61
22	5	315	CLA	C3D-C2D-C1D	-2.00	103.09	105.83
22	1	305	CLA	O1D-CGD-CBD	-2.00	120.38	124.48
24	L	204	BCR	C29-C30-C25	2.00	113.57	110.48
22	B	827	CLA	C3D-C2D-C1D	-2.00	103.10	105.83
22	6	302	CLA	C3D-C2D-C1D	-2.00	103.10	105.83
39	4	814	CHL	CHB-C4A-NA	2.00	127.28	124.51
22	7	315	CLA	CMB-C2B-C3B	2.00	128.43	124.68
22	A	813	CLA	C6-C5-C3	-2.00	108.20	113.45
22	1	313	CLA	OBD-CAD-C3D	-2.00	123.70	128.52
22	5	318	CLA	CMB-C2B-C3B	2.00	128.43	124.68
38	9	302	LUT	C30-C31-C32	-2.00	116.97	123.22
22	5	311	CLA	CMB-C2B-C3B	2.00	128.43	124.68
22	B	809	CLA	CHA-C1A-NA	-2.00	121.81	126.40
22	5	323	CLA	CHA-C1A-NA	-2.00	121.81	126.40
36	J	105	C7Z	C4-C5-C6	-2.00	116.39	120.85
22	B	836	CLA	O1D-CGD-CBD	-2.00	120.39	124.48
22	9	308	CLA	O2A-CGA-CBA	2.00	118.19	111.91
22	A	836	CLA	O2A-CGA-CBA	2.00	118.19	111.91
22	A	808	CLA	CHA-C1A-NA	-2.00	121.81	126.40
23	A	843	PQN	C21-C22-C23	-2.00	109.45	115.92
39	6	320	CHL	CMA-C3A-C2A	2.00	121.90	113.83
22	B	811	CLA	CAA-C2A-C3A	-2.00	107.30	112.78
24	I	4001	BCR	C16-C15-C14	-2.00	119.38	123.47
24	6	306	BCR	C36-C18-C17	-2.00	120.12	122.92
22	B	819	CLA	C6-C5-C3	-2.00	108.21	113.45
22	7	308	CLA	C3D-C2D-C1D	-2.00	103.10	105.83
24	A	847	BCR	C36-C18-C17	-2.00	120.12	122.92
22	B	832	CLA	CHA-C1A-NA	-2.00	121.82	126.40

All (283) chirality outliers are listed below:

Mol	Chain	Res	Type	Atom
21	A	801	CL0	ND
21	A	801	CL0	NC
21	A	801	CL0	NA
22	A	802	CLA	ND

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Mol	Chain	Res	Type	Atom
22	A	803	CLA	ND
22	A	804	CLA	ND
22	A	805	CLA	ND
22	A	806	CLA	ND
22	A	807	CLA	ND
22	A	808	CLA	ND
22	A	809	CLA	ND
22	A	810	CLA	ND
22	A	811	CLA	ND
22	A	812	CLA	ND
22	A	813	CLA	ND
22	A	814	CLA	ND
22	A	815	CLA	ND
22	A	816	CLA	ND
22	A	817	CLA	ND
22	A	818	CLA	ND
22	A	819	CLA	ND
22	A	820	CLA	ND
22	A	821	CLA	ND
22	A	822	CLA	ND
22	A	823	CLA	ND
22	A	824	CLA	ND
22	A	825	CLA	ND
22	A	826	CLA	ND
22	A	827	CLA	ND
22	A	828	CLA	ND
22	A	829	CLA	ND
22	A	830	CLA	ND
22	A	831	CLA	ND
22	A	832	CLA	ND
22	A	833	CLA	ND
22	A	834	CLA	ND
22	A	835	CLA	ND
22	A	836	CLA	ND
22	A	837	CLA	ND
22	A	838	CLA	ND
22	A	839	CLA	ND
22	A	840	CLA	ND
22	A	841	CLA	ND
22	A	842	CLA	ND
22	A	854	CLA	ND
22	A	855	CLA	ND

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Mol	Chain	Res	Type	Atom
22	B	801	CLA	ND
22	B	805	CLA	ND
22	B	806	CLA	ND
22	B	807	CLA	ND
22	B	808	CLA	ND
22	B	809	CLA	ND
22	B	810	CLA	ND
22	B	811	CLA	ND
22	B	812	CLA	ND
22	B	812	CLA	C13
22	B	813	CLA	ND
22	B	814	CLA	ND
22	B	815	CLA	ND
22	B	816	CLA	ND
22	B	817	CLA	ND
22	B	818	CLA	ND
22	B	819	CLA	ND
22	B	820	CLA	ND
22	B	821	CLA	ND
22	B	822	CLA	ND
22	B	823	CLA	ND
22	B	824	CLA	ND
22	B	825	CLA	ND
22	B	826	CLA	ND
22	B	827	CLA	ND
22	B	828	CLA	ND
22	B	829	CLA	ND
22	B	830	CLA	ND
22	B	831	CLA	ND
22	B	832	CLA	ND
22	B	833	CLA	ND
22	B	834	CLA	ND
22	B	835	CLA	ND
22	B	836	CLA	ND
22	B	837	CLA	ND
22	B	838	CLA	ND
22	B	839	CLA	ND
22	B	840	CLA	ND
22	B	841	CLA	ND
22	B	842	CLA	ND
22	F	301	CLA	ND
22	F	302	CLA	ND

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Mol	Chain	Res	Type	Atom
22	F	304	CLA	ND
22	F	305	CLA	ND
22	G	201	CLA	ND
22	G	202	CLA	ND
22	J	103	CLA	ND
22	K	202	CLA	ND
22	K	203	CLA	ND
22	K	204	CLA	ND
22	K	205	CLA	ND
22	L	201	CLA	ND
22	L	202	CLA	ND
22	L	203	CLA	ND
22	1	304	CLA	ND
22	1	305	CLA	ND
22	1	306	CLA	ND
22	1	307	CLA	ND
22	1	308	CLA	ND
22	1	309	CLA	ND
22	1	310	CLA	ND
22	1	311	CLA	ND
22	1	313	CLA	ND
22	1	314	CLA	ND
22	1	315	CLA	ND
22	1	316	CLA	ND
22	Z	303	CLA	ND
22	Z	304	CLA	ND
22	Z	305	CLA	ND
22	Z	306	CLA	ND
22	Z	307	CLA	ND
22	Z	308	CLA	ND
22	Z	309	CLA	ND
22	Z	310	CLA	ND
22	Z	313	CLA	ND
22	Z	314	CLA	ND
22	Z	315	CLA	ND
22	Z	316	CLA	ND
22	3	308	CLA	ND
22	3	309	CLA	ND
22	3	310	CLA	ND
22	3	311	CLA	ND
22	3	312	CLA	ND
22	3	313	CLA	ND

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Mol	Chain	Res	Type	Atom
22	3	314	CLA	ND
22	3	315	CLA	ND
22	3	316	CLA	ND
22	3	318	CLA	ND
22	3	319	CLA	ND
22	3	320	CLA	ND
22	3	322	CLA	ND
22	7	304	CLA	ND
22	7	305	CLA	ND
22	7	306	CLA	ND
22	7	307	CLA	ND
22	7	308	CLA	ND
22	7	309	CLA	ND
22	7	310	CLA	ND
22	7	311	CLA	ND
22	7	312	CLA	ND
22	7	314	CLA	ND
22	7	315	CLA	ND
22	7	316	CLA	ND
22	7	317	CLA	ND
22	7	322	CLA	ND
22	7	323	CLA	ND
22	7	324	CLA	ND
22	8	305	CLA	ND
22	8	306	CLA	ND
22	8	307	CLA	ND
22	8	308	CLA	ND
22	8	309	CLA	ND
22	8	310	CLA	ND
22	8	311	CLA	ND
22	8	313	CLA	ND
22	8	314	CLA	ND
22	8	316	CLA	ND
22	4	805	CLA	ND
22	4	806	CLA	ND
22	4	807	CLA	ND
22	4	808	CLA	ND
22	4	809	CLA	ND
22	4	810	CLA	ND
22	4	811	CLA	ND
22	4	812	CLA	ND
22	4	815	CLA	ND

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Mol	Chain	Res	Type	Atom
22	4	817	CLA	ND
22	4	818	CLA	ND
22	5	301	CLA	ND
22	5	307	CLA	ND
22	5	308	CLA	ND
22	5	309	CLA	ND
22	5	310	CLA	ND
22	5	311	CLA	ND
22	5	312	CLA	ND
22	5	313	CLA	ND
22	5	314	CLA	ND
22	5	315	CLA	ND
22	5	318	CLA	ND
22	5	319	CLA	ND
22	5	320	CLA	ND
22	5	322	CLA	ND
22	5	323	CLA	ND
22	5	326	CLA	ND
22	6	301	CLA	ND
22	6	302	CLA	ND
22	6	307	CLA	ND
22	6	308	CLA	ND
22	6	309	CLA	ND
22	6	310	CLA	ND
22	6	311	CLA	ND
22	6	312	CLA	ND
22	6	313	CLA	ND
22	6	314	CLA	ND
22	6	317	CLA	ND
22	6	319	CLA	ND
22	6	321	CLA	ND
22	6	322	CLA	ND
22	9	303	CLA	ND
22	9	304	CLA	ND
22	9	305	CLA	ND
22	9	306	CLA	ND
22	9	307	CLA	ND
22	9	308	CLA	ND
22	9	309	CLA	ND
22	9	310	CLA	ND
22	9	311	CLA	ND
22	9	313	CLA	ND

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Mol	Chain	Res	Type	Atom
22	2	301	CLA	ND
22	2	302	CLA	ND
22	2	304	CLA	ND
39	1	312	CHL	ND
39	1	312	CHL	NC
39	1	312	CHL	NA
39	Z	311	CHL	ND
39	Z	311	CHL	NC
39	Z	311	CHL	NA
39	Z	312	CHL	ND
39	Z	312	CHL	NC
39	Z	312	CHL	NA
39	3	317	CHL	ND
39	3	317	CHL	NC
39	3	317	CHL	NA
39	7	313	CHL	ND
39	7	313	CHL	NC
39	7	313	CHL	NA
39	8	301	CHL	ND
39	8	301	CHL	NC
39	8	301	CHL	NA
39	8	312	CHL	ND
39	8	312	CHL	NC
39	8	312	CHL	NA
39	8	315	CHL	ND
39	8	315	CHL	NC
39	8	315	CHL	NA
39	4	813	CHL	ND
39	4	813	CHL	NC
39	4	813	CHL	NA
39	4	814	CHL	ND
39	4	814	CHL	NC
39	4	814	CHL	NA
39	4	816	CHL	ND
39	4	816	CHL	NC
39	4	816	CHL	NA
39	4	819	CHL	ND
39	4	819	CHL	NC
39	4	819	CHL	NA
39	5	316	CHL	ND
39	5	316	CHL	NC
39	5	316	CHL	NA

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Mol	Chain	Res	Type	Atom
39	5	317	CHL	ND
39	5	317	CHL	NC
39	5	317	CHL	NA
39	5	321	CHL	ND
39	5	321	CHL	NC
39	5	321	CHL	NA
39	6	315	CHL	ND
39	6	315	CHL	NC
39	6	315	CHL	NA
39	6	316	CHL	ND
39	6	316	CHL	NC
39	6	316	CHL	NA
39	6	318	CHL	ND
39	6	318	CHL	NC
39	6	318	CHL	NA
39	6	320	CHL	ND
39	6	320	CHL	NC
39	6	320	CHL	NA
39	9	312	CHL	ND
39	9	312	CHL	NC
39	9	312	CHL	NA
39	9	314	CHL	ND
39	9	314	CHL	NC
39	9	314	CHL	NA
39	2	303	CHL	ND
39	2	303	CHL	NC
39	2	303	CHL	NA

All (4072) torsion outliers are listed below:

Mol	Chain	Res	Type	Atoms
22	A	802	CLA	C1A-C2A-CAA-CBA
22	A	802	CLA	C2-C1-O2A-CGA
22	A	803	CLA	C1A-C2A-CAA-CBA
22	A	803	CLA	CBA-CGA-O2A-C1
22	A	803	CLA	O1A-CGA-O2A-C1
22	A	804	CLA	C3A-C2A-CAA-CBA
22	A	805	CLA	C3A-C2A-CAA-CBA
22	A	806	CLA	C1A-C2A-CAA-CBA
22	A	806	CLA	C3A-C2A-CAA-CBA
22	A	806	CLA	CBA-CGA-O2A-C1
22	A	806	CLA	O1A-CGA-O2A-C1

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Mol	Chain	Res	Type	Atoms
22	A	806	CLA	C6-C7-C8-C9
22	A	807	CLA	C1A-C2A-CAA-CBA
22	A	807	CLA	C3A-C2A-CAA-CBA
22	A	807	CLA	CBD-CGD-O2D-CED
22	A	807	CLA	C11-C10-C8-C9
22	A	808	CLA	C3A-C2A-CAA-CBA
22	A	808	CLA	CHA-CBD-CGD-O1D
22	A	808	CLA	CHA-CBD-CGD-O2D
22	A	810	CLA	C2-C1-O2A-CGA
22	A	811	CLA	C1A-C2A-CAA-CBA
22	A	811	CLA	CBD-CGD-O2D-CED
22	A	812	CLA	C1A-C2A-CAA-CBA
22	A	812	CLA	C3A-C2A-CAA-CBA
22	A	814	CLA	C2-C1-O2A-CGA
22	A	815	CLA	C3A-C2A-CAA-CBA
22	A	815	CLA	C2-C1-O2A-CGA
22	A	816	CLA	C1A-C2A-CAA-CBA
22	A	816	CLA	CBD-CGD-O2D-CED
22	A	817	CLA	CBD-CGD-O2D-CED
22	A	817	CLA	C2-C3-C5-C6
22	A	817	CLA	C4-C3-C5-C6
22	A	818	CLA	C1A-C2A-CAA-CBA
22	A	818	CLA	C3A-C2A-CAA-CBA
22	A	818	CLA	C2-C1-O2A-CGA
22	A	818	CLA	CBD-CGD-O2D-CED
22	A	819	CLA	C1A-C2A-CAA-CBA
22	A	819	CLA	C3A-C2A-CAA-CBA
22	A	820	CLA	C1A-C2A-CAA-CBA
22	A	820	CLA	CBD-CGD-O2D-CED
22	A	821	CLA	C6-C7-C8-C9
22	A	822	CLA	C1A-C2A-CAA-CBA
22	A	822	CLA	C3A-C2A-CAA-CBA
22	A	822	CLA	CAD-CBD-CGD-O1D
22	A	822	CLA	CAD-CBD-CGD-O2D
22	A	823	CLA	CBD-CGD-O2D-CED
22	A	824	CLA	C1A-C2A-CAA-CBA
22	A	825	CLA	C1A-C2A-CAA-CBA
22	A	825	CLA	CBA-CGA-O2A-C1
22	A	825	CLA	O1A-CGA-O2A-C1
22	A	826	CLA	C3A-C2A-CAA-CBA
22	A	826	CLA	CBA-CGA-O2A-C1
22	A	827	CLA	C1A-C2A-CAA-CBA

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Mol	Chain	Res	Type	Atoms
22	A	827	CLA	C3A-C2A-CAA-CBA
22	A	828	CLA	C1A-C2A-CAA-CBA
22	A	828	CLA	C3A-C2A-CAA-CBA
22	A	828	CLA	C2-C1-O2A-CGA
22	A	828	CLA	CBD-CGD-O2D-CED
22	A	828	CLA	C11-C10-C8-C7
22	A	829	CLA	CBA-CGA-O2A-C1
22	A	829	CLA	O1A-CGA-O2A-C1
22	A	829	CLA	CHA-CBD-CGD-O1D
22	A	830	CLA	CAD-CBD-CGD-O1D
22	A	831	CLA	C2-C1-O2A-CGA
22	A	832	CLA	C1A-C2A-CAA-CBA
22	A	832	CLA	C3A-C2A-CAA-CBA
22	A	833	CLA	C1A-C2A-CAA-CBA
22	A	833	CLA	C3A-C2A-CAA-CBA
22	A	833	CLA	C2-C1-O2A-CGA
22	A	833	CLA	CBD-CGD-O2D-CED
22	A	834	CLA	CHA-CBD-CGD-O1D
22	A	834	CLA	CHA-CBD-CGD-O2D
22	A	835	CLA	CBA-CGA-O2A-C1
22	A	835	CLA	O1A-CGA-O2A-C1
22	A	835	CLA	CBD-CGD-O2D-CED
22	A	836	CLA	C1A-C2A-CAA-CBA
22	A	836	CLA	CHA-CBD-CGD-O1D
22	A	836	CLA	CHA-CBD-CGD-O2D
22	A	838	CLA	C1A-C2A-CAA-CBA
22	A	838	CLA	C3A-C2A-CAA-CBA
22	A	839	CLA	C1A-C2A-CAA-CBA
22	A	839	CLA	C3A-C2A-CAA-CBA
22	A	839	CLA	CHA-CBD-CGD-O1D
22	A	840	CLA	C1A-C2A-CAA-CBA
22	A	840	CLA	C3A-C2A-CAA-CBA
22	A	841	CLA	C1A-C2A-CAA-CBA
22	A	842	CLA	C1A-C2A-CAA-CBA
22	A	842	CLA	C3A-C2A-CAA-CBA
22	A	842	CLA	CHA-CBD-CGD-O1D
22	A	842	CLA	CHA-CBD-CGD-O2D
22	A	842	CLA	CBD-CGD-O2D-CED
22	A	854	CLA	CBD-CGD-O2D-CED
22	A	854	CLA	O1D-CGD-O2D-CED
22	A	855	CLA	C2-C1-O2A-CGA
22	A	855	CLA	CBD-CGD-O2D-CED

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Mol	Chain	Res	Type	Atoms
22	B	801	CLA	CBD-CGD-O2D-CED
22	B	805	CLA	CHA-CBD-CGD-O1D
22	B	805	CLA	CHA-CBD-CGD-O2D
22	B	805	CLA	CBD-CGD-O2D-CED
22	B	806	CLA	CBD-CGD-O2D-CED
22	B	807	CLA	C1A-C2A-CAA-CBA
22	B	808	CLA	CBD-CGD-O2D-CED
22	B	810	CLA	C1A-C2A-CAA-CBA
22	B	810	CLA	C3A-C2A-CAA-CBA
22	B	811	CLA	C1A-C2A-CAA-CBA
22	B	811	CLA	C3A-C2A-CAA-CBA
22	B	811	CLA	CHA-CBD-CGD-O1D
22	B	811	CLA	CHA-CBD-CGD-O2D
22	B	812	CLA	C1A-C2A-CAA-CBA
22	B	812	CLA	C3A-C2A-CAA-CBA
22	B	812	CLA	O1A-CGA-O2A-C1
22	B	812	CLA	C2-C1-O2A-CGA
22	B	812	CLA	CBD-CGD-O2D-CED
22	B	813	CLA	O1A-CGA-O2A-C1
22	B	813	CLA	CAD-CBD-CGD-O1D
22	B	814	CLA	CBD-CGD-O2D-CED
22	B	815	CLA	C1A-C2A-CAA-CBA
22	B	815	CLA	CHA-CBD-CGD-O1D
22	B	815	CLA	CHA-CBD-CGD-O2D
22	B	816	CLA	CBD-CGD-O2D-CED
22	B	817	CLA	C3A-C2A-CAA-CBA
22	B	817	CLA	CHA-CBD-CGD-O1D
22	B	817	CLA	CHA-CBD-CGD-O2D
22	B	818	CLA	C1A-C2A-CAA-CBA
22	B	818	CLA	C3A-C2A-CAA-CBA
22	B	819	CLA	C1A-C2A-CAA-CBA
22	B	819	CLA	C3A-C2A-CAA-CBA
22	B	819	CLA	C2-C1-O2A-CGA
22	B	820	CLA	C3A-C2A-CAA-CBA
22	B	821	CLA	C1A-C2A-CAA-CBA
22	B	821	CLA	C3A-C2A-CAA-CBA
22	B	822	CLA	C1A-C2A-CAA-CBA
22	B	822	CLA	C3A-C2A-CAA-CBA
22	B	822	CLA	CBA-CGA-O2A-C1
22	B	822	CLA	O1A-CGA-O2A-C1
22	B	824	CLA	C1A-C2A-CAA-CBA
22	B	824	CLA	CBD-CGD-O2D-CED

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Mol	Chain	Res	Type	Atoms
22	B	825	CLA	C3A-C2A-CAA-CBA
22	B	826	CLA	CHA-CBD-CGD-O1D
22	B	826	CLA	CHA-CBD-CGD-O2D
22	B	827	CLA	C3A-C2A-CAA-CBA
22	B	827	CLA	CBA-CGA-O2A-C1
22	B	827	CLA	O1A-CGA-O2A-C1
22	B	828	CLA	CHA-CBD-CGD-O1D
22	B	829	CLA	C3A-C2A-CAA-CBA
22	B	829	CLA	CHA-CBD-CGD-O1D
22	B	829	CLA	CHA-CBD-CGD-O2D
22	B	830	CLA	C1A-C2A-CAA-CBA
22	B	830	CLA	C3A-C2A-CAA-CBA
22	B	831	CLA	C1A-C2A-CAA-CBA
22	B	831	CLA	CBD-CGD-O2D-CED
22	B	832	CLA	C3A-C2A-CAA-CBA
22	B	833	CLA	C1A-C2A-CAA-CBA
22	B	833	CLA	C3A-C2A-CAA-CBA
22	B	834	CLA	C2-C1-O2A-CGA
22	B	834	CLA	CHA-CBD-CGD-O1D
22	B	834	CLA	CHA-CBD-CGD-O2D
22	B	835	CLA	C1A-C2A-CAA-CBA
22	B	835	CLA	C3A-C2A-CAA-CBA
22	B	835	CLA	C2-C1-O2A-CGA
22	B	836	CLA	C1A-C2A-CAA-CBA
22	B	837	CLA	C3A-C2A-CAA-CBA
22	B	841	CLA	CBD-CGD-O2D-CED
22	B	841	CLA	C11-C10-C8-C9
22	F	301	CLA	C1A-C2A-CAA-CBA
22	F	301	CLA	C3A-C2A-CAA-CBA
22	F	301	CLA	CHA-CBD-CGD-O1D
22	F	301	CLA	CHA-CBD-CGD-O2D
22	F	302	CLA	C2-C1-O2A-CGA
22	F	302	CLA	CBD-CGD-O2D-CED
22	F	304	CLA	C1A-C2A-CAA-CBA
22	F	304	CLA	CBA-CGA-O2A-C1
22	F	304	CLA	O1A-CGA-O2A-C1
22	F	304	CLA	CHA-CBD-CGD-O1D
22	F	304	CLA	CHA-CBD-CGD-O2D
22	F	304	CLA	CAD-CBD-CGD-O1D
22	F	304	CLA	CAD-CBD-CGD-O2D
22	F	305	CLA	C1A-C2A-CAA-CBA
22	F	305	CLA	C3A-C2A-CAA-CBA

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Mol	Chain	Res	Type	Atoms
22	G	202	CLA	C1A-C2A-CAA-CBA
22	G	202	CLA	C3A-C2A-CAA-CBA
22	G	202	CLA	CHA-CBD-CGD-O2D
22	J	103	CLA	C1A-C2A-CAA-CBA
22	K	202	CLA	CHA-CBD-CGD-O1D
22	K	204	CLA	CAD-CBD-CGD-O2D
22	K	205	CLA	CBD-CGD-O2D-CED
22	K	205	CLA	C2-C3-C5-C6
22	K	205	CLA	C4-C3-C5-C6
22	L	201	CLA	C1A-C2A-CAA-CBA
22	L	202	CLA	C1A-C2A-CAA-CBA
22	L	202	CLA	C3A-C2A-CAA-CBA
22	L	203	CLA	C3A-C2A-CAA-CBA
22	1	304	CLA	C1A-C2A-CAA-CBA
22	1	304	CLA	C3A-C2A-CAA-CBA
22	1	306	CLA	CBD-CGD-O2D-CED
22	1	307	CLA	C1A-C2A-CAA-CBA
22	1	307	CLA	C3A-C2A-CAA-CBA
22	1	308	CLA	C2-C1-O2A-CGA
22	1	308	CLA	CHA-CBD-CGD-O1D
22	1	308	CLA	CHA-CBD-CGD-O2D
22	1	309	CLA	C1A-C2A-CAA-CBA
22	1	309	CLA	C3A-C2A-CAA-CBA
22	1	310	CLA	C1A-C2A-CAA-CBA
22	1	313	CLA	CBD-CGD-O2D-CED
22	1	313	CLA	O2A-C1-C2-C3
22	1	314	CLA	C3A-C2A-CAA-CBA
22	1	315	CLA	C1A-C2A-CAA-CBA
22	Z	303	CLA	C3A-C2A-CAA-CBA
22	Z	303	CLA	CBA-CGA-O2A-C1
22	Z	303	CLA	O1A-CGA-O2A-C1
22	Z	303	CLA	CHA-CBD-CGD-O1D
22	Z	303	CLA	CHA-CBD-CGD-O2D
22	Z	303	CLA	CBD-CGD-O2D-CED
22	Z	305	CLA	CBD-CGD-O2D-CED
22	Z	306	CLA	C3A-C2A-CAA-CBA
22	Z	307	CLA	CHA-CBD-CGD-O1D
22	Z	307	CLA	CHA-CBD-CGD-O2D
22	Z	308	CLA	C1A-C2A-CAA-CBA
22	Z	308	CLA	CBD-CGD-O2D-CED
22	Z	310	CLA	C1A-C2A-CAA-CBA
22	Z	310	CLA	C2-C1-O2A-CGA

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Mol	Chain	Res	Type	Atoms
22	Z	313	CLA	O2A-C1-C2-C3
22	Z	314	CLA	C1A-C2A-CAA-CBA
22	Z	314	CLA	C3A-C2A-CAA-CBA
22	Z	314	CLA	CBD-CGD-O2D-CED
22	Z	315	CLA	CBD-CGD-O2D-CED
22	Z	316	CLA	O1A-CGA-O2A-C1
22	Z	316	CLA	CBD-CGD-O2D-CED
22	Z	316	CLA	C11-C10-C8-C9
22	3	308	CLA	C1A-C2A-CAA-CBA
22	3	308	CLA	C3A-C2A-CAA-CBA
22	3	309	CLA	CBD-CGD-O2D-CED
22	3	310	CLA	CHA-CBD-CGD-O1D
22	3	310	CLA	CHA-CBD-CGD-O2D
22	3	310	CLA	CBD-CGD-O2D-CED
22	3	311	CLA	C3A-C2A-CAA-CBA
22	3	311	CLA	C6-C7-C8-C9
22	3	311	CLA	C11-C10-C8-C7
22	3	312	CLA	CBD-CGD-O2D-CED
22	3	313	CLA	C3A-C2A-CAA-CBA
22	3	314	CLA	C1A-C2A-CAA-CBA
22	3	314	CLA	C3A-C2A-CAA-CBA
22	3	315	CLA	C1A-C2A-CAA-CBA
22	3	315	CLA	CBD-CGD-O2D-CED
22	3	316	CLA	C1A-C2A-CAA-CBA
22	3	316	CLA	CBD-CGD-O2D-CED
22	3	318	CLA	C3A-C2A-CAA-CBA
22	3	319	CLA	C4-C3-C5-C6
22	7	304	CLA	C2-C1-O2A-CGA
22	7	305	CLA	CBD-CGD-O2D-CED
22	7	306	CLA	C1A-C2A-CAA-CBA
22	7	306	CLA	CBD-CGD-O2D-CED
22	7	307	CLA	C3A-C2A-CAA-CBA
22	7	308	CLA	CBA-CGA-O2A-C1
22	7	308	CLA	O1A-CGA-O2A-C1
22	7	309	CLA	CBD-CGD-O2D-CED
22	7	310	CLA	C1A-C2A-CAA-CBA
22	7	310	CLA	C6-C7-C8-C9
22	7	311	CLA	C1A-C2A-CAA-CBA
22	7	312	CLA	C3A-C2A-CAA-CBA
22	7	312	CLA	CHA-CBD-CGD-O1D
22	7	312	CLA	CBD-CGD-O2D-CED
22	7	314	CLA	C1A-C2A-CAA-CBA

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Mol	Chain	Res	Type	Atoms
22	7	314	CLA	C3A-C2A-CAA-CBA
22	7	314	CLA	CBA-CGA-O2A-C1
22	7	314	CLA	O1A-CGA-O2A-C1
22	7	315	CLA	CHA-CBD-CGD-O1D
22	7	315	CLA	CHA-CBD-CGD-O2D
22	7	316	CLA	C1A-C2A-CAA-CBA
22	7	316	CLA	CBD-CGD-O2D-CED
22	7	317	CLA	C1A-C2A-CAA-CBA
22	7	317	CLA	C3A-C2A-CAA-CBA
22	7	317	CLA	C2-C1-O2A-CGA
22	7	323	CLA	C1A-C2A-CAA-CBA
22	7	323	CLA	C3A-C2A-CAA-CBA
22	7	323	CLA	C2-C1-O2A-CGA
22	7	323	CLA	CHA-CBD-CGD-O1D
22	7	323	CLA	CHA-CBD-CGD-O2D
22	7	323	CLA	CAD-CBD-CGD-O1D
22	7	323	CLA	CAD-CBD-CGD-O2D
22	7	324	CLA	C3A-C2A-CAA-CBA
22	7	324	CLA	C2-C1-O2A-CGA
22	8	305	CLA	C1A-C2A-CAA-CBA
22	8	305	CLA	C3A-C2A-CAA-CBA
22	8	305	CLA	CBD-CGD-O2D-CED
22	8	307	CLA	C1A-C2A-CAA-CBA
22	8	308	CLA	C3A-C2A-CAA-CBA
22	8	309	CLA	C1A-C2A-CAA-CBA
22	8	309	CLA	C3A-C2A-CAA-CBA
22	8	310	CLA	CBD-CGD-O2D-CED
22	8	311	CLA	C1A-C2A-CAA-CBA
22	8	311	CLA	C3A-C2A-CAA-CBA
22	8	311	CLA	CHA-CBD-CGD-O1D
22	8	313	CLA	CBD-CGD-O2D-CED
22	8	314	CLA	C1A-C2A-CAA-CBA
22	8	316	CLA	C1A-C2A-CAA-CBA
22	8	316	CLA	C3A-C2A-CAA-CBA
22	4	805	CLA	C1A-C2A-CAA-CBA
22	4	805	CLA	C3A-C2A-CAA-CBA
22	4	806	CLA	C2A-CAA-CBA-CGA
22	4	806	CLA	C2-C1-O2A-CGA
22	4	807	CLA	C1A-C2A-CAA-CBA
22	4	808	CLA	C3A-C2A-CAA-CBA
22	4	808	CLA	C2-C3-C5-C6
22	4	808	CLA	C4-C3-C5-C6

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Mol	Chain	Res	Type	Atoms
22	4	809	CLA	CBA-CGA-O2A-C1
22	4	809	CLA	O1A-CGA-O2A-C1
22	4	809	CLA	CBD-CGD-O2D-CED
22	4	811	CLA	CBA-CGA-O2A-C1
22	4	811	CLA	O1A-CGA-O2A-C1
22	4	815	CLA	C3A-C2A-CAA-CBA
22	4	815	CLA	CHA-CBD-CGD-O1D
22	4	815	CLA	CHA-CBD-CGD-O2D
22	4	815	CLA	CBD-CGD-O2D-CED
22	4	818	CLA	C1A-C2A-CAA-CBA
22	4	818	CLA	C2-C1-O2A-CGA
22	5	301	CLA	CBD-CGD-O2D-CED
22	5	307	CLA	C1A-C2A-CAA-CBA
22	5	307	CLA	C3A-C2A-CAA-CBA
22	5	308	CLA	C1A-C2A-CAA-CBA
22	5	308	CLA	CBA-CGA-O2A-C1
22	5	308	CLA	O1A-CGA-O2A-C1
22	5	308	CLA	CBD-CGD-O2D-CED
22	5	309	CLA	C2-C1-O2A-CGA
22	5	310	CLA	C3A-C2A-CAA-CBA
22	5	310	CLA	C6-C7-C8-C9
22	5	311	CLA	C3A-C2A-CAA-CBA
22	5	313	CLA	CBA-CGA-O2A-C1
22	5	313	CLA	CHA-CBD-CGD-O1D
22	5	313	CLA	CHA-CBD-CGD-O2D
22	5	314	CLA	C1A-C2A-CAA-CBA
22	5	314	CLA	CHA-CBD-CGD-O1D
22	5	315	CLA	C1A-C2A-CAA-CBA
22	5	315	CLA	C3A-C2A-CAA-CBA
22	5	315	CLA	CHA-CBD-CGD-O1D
22	5	315	CLA	CBD-CGD-O2D-CED
22	5	315	CLA	C6-C7-C8-C9
22	5	318	CLA	C3A-C2A-CAA-CBA
22	5	319	CLA	C1A-C2A-CAA-CBA
22	5	319	CLA	C3A-C2A-CAA-CBA
22	5	320	CLA	C1A-C2A-CAA-CBA
22	5	320	CLA	C3A-C2A-CAA-CBA
22	5	320	CLA	CBD-CGD-O2D-CED
22	5	322	CLA	C1A-C2A-CAA-CBA
22	5	322	CLA	C3A-C2A-CAA-CBA
22	5	323	CLA	CBD-CGD-O2D-CED
22	5	326	CLA	C1A-C2A-CAA-CBA

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Mol	Chain	Res	Type	Atoms
22	5	326	CLA	CBD-CGD-O2D-CED
22	6	301	CLA	C1A-C2A-CAA-CBA
22	6	301	CLA	C3A-C2A-CAA-CBA
22	6	302	CLA	C1A-C2A-CAA-CBA
22	6	302	CLA	C3A-C2A-CAA-CBA
22	6	302	CLA	CHA-CBD-CGD-O1D
22	6	302	CLA	CHA-CBD-CGD-O2D
22	6	302	CLA	CBD-CGD-O2D-CED
22	6	307	CLA	C1A-C2A-CAA-CBA
22	6	307	CLA	CBD-CGD-O2D-CED
22	6	308	CLA	C2-C1-O2A-CGA
22	6	309	CLA	CHA-CBD-CGD-O1D
22	6	311	CLA	C1A-C2A-CAA-CBA
22	6	311	CLA	C3A-C2A-CAA-CBA
22	6	311	CLA	CBD-CGD-O2D-CED
22	6	312	CLA	C2-C1-O2A-CGA
22	6	313	CLA	CHA-CBD-CGD-O1D
22	6	313	CLA	CHA-CBD-CGD-O2D
22	6	314	CLA	CHA-CBD-CGD-O1D
22	6	317	CLA	C3A-C2A-CAA-CBA
22	6	317	CLA	CBA-CGA-O2A-C1
22	6	317	CLA	O1A-CGA-O2A-C1
22	6	317	CLA	CHA-CBD-CGD-O2D
22	6	319	CLA	C3A-C2A-CAA-CBA
22	6	321	CLA	C1A-C2A-CAA-CBA
22	6	321	CLA	C3A-C2A-CAA-CBA
22	6	321	CLA	CBA-CGA-O2A-C1
22	6	322	CLA	C1A-C2A-CAA-CBA
22	6	322	CLA	C3A-C2A-CAA-CBA
22	6	322	CLA	CHA-CBD-CGD-O2D
22	9	303	CLA	C1A-C2A-CAA-CBA
22	9	308	CLA	C2-C1-O2A-CGA
22	9	308	CLA	CBD-CGD-O2D-CED
22	9	309	CLA	CHA-CBD-CGD-O1D
22	9	311	CLA	C1A-C2A-CAA-CBA
22	9	311	CLA	CBD-CGD-O2D-CED
22	9	313	CLA	CBA-CGA-O2A-C1
22	2	301	CLA	CBA-CGA-O2A-C1
22	2	301	CLA	O1A-CGA-O2A-C1
22	2	301	CLA	CHA-CBD-CGD-O1D
22	2	301	CLA	CHA-CBD-CGD-O2D
22	2	302	CLA	C1A-C2A-CAA-CBA

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Mol	Chain	Res	Type	Atoms
22	2	302	CLA	CHA-CBD-CGD-O2D
22	2	304	CLA	C1A-C2A-CAA-CBA
22	2	304	CLA	C3A-C2A-CAA-CBA
22	2	304	CLA	CBA-CGA-O2A-C1
24	A	844	BCR	C5-C6-C7-C8
24	A	844	BCR	C21-C22-C23-C24
24	A	844	BCR	C37-C22-C23-C24
24	A	845	BCR	C7-C8-C9-C34
24	A	845	BCR	C21-C22-C23-C24
24	A	845	BCR	C37-C22-C23-C24
24	A	846	BCR	C7-C8-C9-C10
24	A	846	BCR	C7-C8-C9-C34
24	A	847	BCR	C1-C6-C7-C8
24	A	847	BCR	C11-C12-C13-C14
24	A	847	BCR	C11-C12-C13-C35
24	A	847	BCR	C13-C14-C15-C16
24	A	847	BCR	C19-C20-C21-C22
24	A	847	BCR	C23-C24-C25-C26
24	A	848	BCR	C15-C16-C17-C18
24	A	848	BCR	C17-C18-C19-C20
24	A	848	BCR	C36-C18-C19-C20
24	A	848	BCR	C21-C22-C23-C24
24	A	848	BCR	C37-C22-C23-C24
24	A	856	BCR	C1-C6-C7-C8
24	A	856	BCR	C9-C10-C11-C12
24	A	856	BCR	C11-C12-C13-C14
24	A	856	BCR	C11-C12-C13-C35
24	A	856	BCR	C17-C18-C19-C20
24	A	856	BCR	C36-C18-C19-C20
24	A	856	BCR	C23-C24-C25-C30
24	B	844	BCR	C1-C6-C7-C8
24	B	844	BCR	C7-C8-C9-C34
24	B	845	BCR	C7-C8-C9-C10
24	B	845	BCR	C7-C8-C9-C34
24	B	845	BCR	C9-C10-C11-C12
24	B	845	BCR	C11-C12-C13-C14
24	B	846	BCR	C11-C12-C13-C14
24	B	846	BCR	C11-C12-C13-C35
24	B	846	BCR	C15-C16-C17-C18
24	B	846	BCR	C17-C18-C19-C20
24	B	846	BCR	C36-C18-C19-C20
24	B	846	BCR	C21-C22-C23-C24

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Mol	Chain	Res	Type	Atoms
24	B	846	BCR	C37-C22-C23-C24
24	B	847	BCR	C11-C12-C13-C14
24	B	847	BCR	C17-C18-C19-C20
24	B	847	BCR	C36-C18-C19-C20
24	B	848	BCR	C1-C6-C7-C8
24	B	848	BCR	C5-C6-C7-C8
24	B	848	BCR	C11-C12-C13-C14
24	B	848	BCR	C11-C12-C13-C35
24	B	848	BCR	C13-C14-C15-C16
24	B	848	BCR	C21-C22-C23-C24
24	B	848	BCR	C37-C22-C23-C24
24	B	849	BCR	C9-C10-C11-C12
24	B	849	BCR	C11-C12-C13-C14
24	B	849	BCR	C11-C12-C13-C35
24	B	849	BCR	C36-C18-C19-C20
24	B	849	BCR	C21-C22-C23-C24
24	B	849	BCR	C37-C22-C23-C24
24	B	849	BCR	C23-C24-C25-C30
24	B	853	BCR	C7-C8-C9-C10
24	B	853	BCR	C7-C8-C9-C34
24	B	853	BCR	C36-C18-C19-C20
24	B	853	BCR	C19-C20-C21-C22
24	F	303	BCR	C1-C6-C7-C8
24	F	303	BCR	C9-C10-C11-C12
24	F	303	BCR	C11-C12-C13-C14
24	F	303	BCR	C11-C12-C13-C35
24	F	303	BCR	C13-C14-C15-C16
24	F	303	BCR	C17-C18-C19-C20
24	F	303	BCR	C36-C18-C19-C20
24	F	303	BCR	C19-C20-C21-C22
24	G	203	BCR	C7-C8-C9-C10
24	G	203	BCR	C7-C8-C9-C34
24	I	4001	BCR	C5-C6-C7-C8
24	I	4001	BCR	C17-C18-C19-C20
24	I	4001	BCR	C36-C18-C19-C20
24	I	4001	BCR	C19-C20-C21-C22
24	I	4001	BCR	C21-C22-C23-C24
24	I	4001	BCR	C37-C22-C23-C24
24	J	104	BCR	C11-C12-C13-C14
24	J	104	BCR	C11-C12-C13-C35
24	J	104	BCR	C17-C18-C19-C20
24	J	104	BCR	C36-C18-C19-C20

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Mol	Chain	Res	Type	Atoms
24	K	206	BCR	C7-C8-C9-C10
24	K	206	BCR	C7-C8-C9-C34
24	K	206	BCR	C19-C20-C21-C22
24	K	206	BCR	C23-C24-C25-C26
24	K	206	BCR	C23-C24-C25-C30
24	L	204	BCR	C7-C8-C9-C10
24	L	204	BCR	C13-C14-C15-C16
24	L	204	BCR	C15-C16-C17-C18
24	L	204	BCR	C36-C18-C19-C20
24	L	204	BCR	C19-C20-C21-C22
24	L	204	BCR	C21-C22-C23-C24
24	L	204	BCR	C37-C22-C23-C24
24	3	304	BCR	C7-C8-C9-C10
24	3	304	BCR	C7-C8-C9-C34
24	3	304	BCR	C13-C14-C15-C16
24	3	304	BCR	C17-C18-C19-C20
24	3	304	BCR	C36-C18-C19-C20
24	3	305	BCR	C1-C6-C7-C8
24	3	305	BCR	C5-C6-C7-C8
24	3	305	BCR	C9-C10-C11-C12
24	3	305	BCR	C11-C12-C13-C14
24	3	305	BCR	C11-C12-C13-C35
24	3	305	BCR	C19-C20-C21-C22
24	3	305	BCR	C21-C22-C23-C24
24	3	305	BCR	C37-C22-C23-C24
24	3	306	BCR	C11-C12-C13-C14
24	3	306	BCR	C11-C12-C13-C35
24	3	306	BCR	C17-C18-C19-C20
24	3	306	BCR	C19-C20-C21-C22
24	3	306	BCR	C21-C22-C23-C24
24	3	306	BCR	C37-C22-C23-C24
24	3	307	BCR	C11-C12-C13-C14
24	3	307	BCR	C11-C12-C13-C35
24	3	307	BCR	C17-C18-C19-C20
24	3	307	BCR	C36-C18-C19-C20
24	3	307	BCR	C23-C24-C25-C26
24	7	303	BCR	C7-C8-C9-C10
24	7	303	BCR	C7-C8-C9-C34
24	7	303	BCR	C11-C12-C13-C14
24	7	303	BCR	C11-C12-C13-C35
24	7	303	BCR	C13-C14-C15-C16
24	7	303	BCR	C23-C24-C25-C26

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Mol	Chain	Res	Type	Atoms
24	8	304	BCR	C11-C12-C13-C14
24	8	304	BCR	C11-C12-C13-C35
24	8	304	BCR	C13-C14-C15-C16
24	4	804	BCR	C7-C8-C9-C10
24	4	804	BCR	C7-C8-C9-C34
24	4	804	BCR	C9-C10-C11-C12
24	4	804	BCR	C23-C24-C25-C26
24	5	304	BCR	C7-C8-C9-C10
24	5	304	BCR	C7-C8-C9-C34
24	5	304	BCR	C11-C12-C13-C14
24	5	304	BCR	C11-C12-C13-C35
24	5	304	BCR	C13-C14-C15-C16
24	5	304	BCR	C23-C24-C25-C26
24	5	305	BCR	C5-C6-C7-C8
24	5	305	BCR	C9-C10-C11-C12
24	5	305	BCR	C17-C18-C19-C20
24	5	305	BCR	C36-C18-C19-C20
24	6	305	BCR	C1-C6-C7-C8
24	6	305	BCR	C11-C12-C13-C14
24	6	305	BCR	C11-C12-C13-C35
24	6	306	BCR	C1-C6-C7-C8
24	6	306	BCR	C5-C6-C7-C8
24	6	306	BCR	C7-C8-C9-C34
24	6	306	BCR	C23-C24-C25-C26
25	A	849	LHG	C4-O6-P-O4
25	A	849	LHG	C4-O6-P-O5
25	A	850	LHG	O1-C1-C2-C3
25	A	850	LHG	C1-C2-C3-O3
25	A	850	LHG	C4-O6-P-O5
25	B	803	LHG	O9-C7-O7-C5
25	B	803	LHG	C8-C7-O7-C5
25	B	850	LHG	C3-O3-P-O6
25	B	850	LHG	O9-C7-O7-C5
25	B	851	LHG	O1-C1-C2-O2
25	B	851	LHG	C3-O3-P-O5
25	B	851	LHG	O9-C7-O7-C5
25	1	317	LHG	C8-C7-O7-C5
25	Z	317	LHG	O1-C1-C2-O2
25	3	321	LHG	O9-C7-O7-C5
25	7	318	LHG	C1-C2-C3-O3
25	7	318	LHG	C4-O6-P-O4
25	8	317	LHG	O1-C1-C2-C3

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Mol	Chain	Res	Type	Atoms
25	8	317	LHG	C4-O6-P-O4
25	4	820	LHG	C3-O3-P-O4
25	4	820	LHG	C8-C7-O7-C5
25	5	324	LHG	O1-C1-C2-O2
25	6	323	LHG	O1-C1-C2-O2
25	6	323	LHG	C3-O3-P-O5
25	6	323	LHG	C4-O6-P-O4
25	6	323	LHG	C4-O6-P-O5
25	6	323	LHG	C24-C23-O8-C6
25	9	315	LHG	C3-O3-P-O4
27	A	852	DGA	CA2-CA1-OG1-CG1
27	A	852	DGA	OA1-CA1-OG1-CG1
31	B	852	DGD	O1B-C1B-O2G-C2G
31	1	319	DGD	C2A-C1A-O1G-C1G
31	1	319	DGD	O1A-C1A-O1G-C1G
31	1	319	DGD	C2B-C1B-O2G-C2G
31	1	319	DGD	O1B-C1B-O2G-C2G
31	1	319	DGD	O6E-C1E-O5D-C6D
31	3	301	DGD	C2B-C1B-O2G-C2G
31	3	301	DGD	O1B-C1B-O2G-C2G
32	F	306	RRX	C7-C8-C9-C10
32	F	306	RRX	C7-C8-C9-C34
32	F	306	RRX	C1-C6-C7-C8
32	F	306	RRX	C5-C6-C7-C8
33	F	307	LMT	C2'-C1'-O1'-C1
33	F	307	LMT	O5'-C1'-O1'-C1
33	G	204	LMT	C2'-C1'-O1'-C1
33	G	204	LMT	O5'-C1'-O1'-C1
33	1	301	LMT	C2'-C1'-O1'-C1
33	1	301	LMT	O5'-C1'-O1'-C1
33	1	301	LMT	C2-C1-O1'-C1'
34	F	308	LMG	C11-C10-O7-C8
36	J	105	C7Z	C1-C6-C7-C8
36	J	105	C7Z	C5-C6-C7-C8
36	J	105	C7Z	C7-C8-C9-C10
36	J	105	C7Z	C31-C32-C33-C34
36	J	105	C7Z	C31-C32-C33-C40
36	5	306	C7Z	C11-C12-C13-C20
36	5	306	C7Z	C11-C12-C13-C14
36	5	306	C7Z	C31-C32-C33-C40
38	1	302	LUT	C11-C12-C13-C14
38	1	302	LUT	C11-C12-C13-C20

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Mol	Chain	Res	Type	Atoms
38	1	302	LUT	C27-C28-C29-C30
38	1	302	LUT	C27-C28-C29-C39
38	1	302	LUT	C31-C32-C33-C34
38	1	302	LUT	C31-C32-C33-C40
38	1	303	LUT	C11-C12-C13-C14
38	1	303	LUT	C11-C12-C13-C20
38	1	303	LUT	C31-C32-C33-C34
38	1	303	LUT	C31-C32-C33-C40
38	Z	301	LUT	C11-C12-C13-C14
38	Z	301	LUT	C11-C12-C13-C20
38	Z	302	LUT	C7-C8-C9-C10
38	Z	302	LUT	C11-C12-C13-C14
38	Z	302	LUT	C11-C12-C13-C20
38	3	302	LUT	C11-C12-C13-C14
38	3	302	LUT	C11-C12-C13-C20
38	3	302	LUT	C31-C32-C33-C34
38	3	302	LUT	C31-C32-C33-C40
38	3	303	LUT	C11-C12-C13-C14
38	3	303	LUT	C11-C12-C13-C20
38	7	301	LUT	C11-C12-C13-C14
38	7	301	LUT	C11-C12-C13-C20
38	7	301	LUT	C31-C32-C33-C34
38	7	301	LUT	C31-C32-C33-C40
38	7	302	LUT	C7-C8-C9-C10
38	8	302	LUT	C11-C12-C13-C14
38	8	302	LUT	C11-C12-C13-C20
38	8	302	LUT	C27-C28-C29-C30
38	8	302	LUT	C27-C28-C29-C39
38	8	303	LUT	C11-C12-C13-C14
38	8	303	LUT	C11-C12-C13-C20
38	4	802	LUT	C5-C6-C7-C8
38	4	802	LUT	C7-C8-C9-C10
38	4	802	LUT	C7-C8-C9-C19
38	4	802	LUT	C31-C32-C33-C34
38	4	802	LUT	C31-C32-C33-C40
38	4	803	LUT	C1-C6-C7-C8
38	4	803	LUT	C11-C12-C13-C14
38	4	803	LUT	C11-C12-C13-C20
38	5	302	LUT	C7-C8-C9-C19
38	5	302	LUT	C11-C12-C13-C14
38	5	302	LUT	C11-C12-C13-C20
38	5	302	LUT	C31-C32-C33-C34

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Mol	Chain	Res	Type	Atoms
38	5	302	LUT	C31-C32-C33-C40
38	5	303	LUT	C1-C6-C7-C8
38	5	303	LUT	C11-C12-C13-C14
38	5	303	LUT	C11-C12-C13-C20
38	6	303	LUT	C7-C8-C9-C10
38	6	303	LUT	C7-C8-C9-C19
38	6	303	LUT	C11-C12-C13-C14
38	6	303	LUT	C11-C12-C13-C20
38	6	303	LUT	C31-C32-C33-C40
38	6	304	LUT	C11-C12-C13-C14
38	6	304	LUT	C11-C12-C13-C20
38	9	301	LUT	C11-C12-C13-C14
38	9	301	LUT	C11-C12-C13-C20
38	9	301	LUT	C31-C32-C33-C34
38	9	301	LUT	C31-C32-C33-C40
38	9	302	LUT	C7-C8-C9-C10
38	9	302	LUT	C7-C8-C9-C19
38	9	302	LUT	C11-C12-C13-C20
39	Z	312	CHL	CBD-CGD-O2D-CED
39	7	313	CHL	CHA-CBD-CGD-O1D
39	7	313	CHL	CHA-CBD-CGD-O2D
39	8	312	CHL	CHA-CBD-CGD-O1D
39	8	312	CHL	CHA-CBD-CGD-O2D
39	8	312	CHL	CBD-CGD-O2D-CED
39	4	814	CHL	CHA-CBD-CGD-O1D
39	4	814	CHL	CHA-CBD-CGD-O2D
39	4	814	CHL	CBD-CGD-O2D-CED
39	4	816	CHL	CHA-CBD-CGD-O1D
39	4	816	CHL	CHA-CBD-CGD-O2D
39	5	317	CHL	C1A-C2A-CAA-CBA
39	5	317	CHL	C3A-C2A-CAA-CBA
39	5	321	CHL	C3A-C2A-CAA-CBA
39	6	315	CHL	CHA-CBD-CGD-O1D
39	6	315	CHL	CHA-CBD-CGD-O2D
39	6	316	CHL	C1A-C2A-CAA-CBA
39	6	316	CHL	C3A-C2A-CAA-CBA
39	6	316	CHL	CBA-CGA-O2A-C1
39	6	316	CHL	O1A-CGA-O2A-C1
39	6	318	CHL	CBD-CGD-O2D-CED
39	9	312	CHL	CBD-CGD-O2D-CED
39	2	303	CHL	CBD-CGD-O2D-CED
40	1	318	SQD	O5-C5-C6-S

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Mol	Chain	Res	Type	Atoms
41	7	319	3PH	C22-C21-O21-C2
41	8	320	3PH	C1-O11-P-O13
41	8	320	3PH	C1-O11-P-O14
41	8	320	3PH	C2-C1-O11-P
41	5	325	3PH	O22-C21-O21-C2
41	5	325	3PH	C22-C21-O21-C2
41	6	324	3PH	C22-C21-O21-C2
42	7	320	SPH	O1-C1-C2-N2
42	7	321	SPH	C1-C2-C3-O3
42	7	321	SPH	C1-C2-C3-C4
42	7	321	SPH	N2-C2-C3-O3
42	7	321	SPH	N2-C2-C3-C4
43	8	319	LPX	C4-C3-O1-P1
43	8	319	LPX	C2-C1-O2-P1
22	B	809	CLA	O1D-CGD-O2D-CED
22	F	302	CLA	O1D-CGD-O2D-CED
22	1	315	CLA	O1D-CGD-O2D-CED
22	Z	315	CLA	O1D-CGD-O2D-CED
22	3	308	CLA	O1D-CGD-O2D-CED
22	7	306	CLA	O1D-CGD-O2D-CED
22	7	309	CLA	O1D-CGD-O2D-CED
22	8	316	CLA	O1D-CGD-O2D-CED
22	4	818	CLA	O1D-CGD-O2D-CED
22	5	319	CLA	O1D-CGD-O2D-CED
22	6	302	CLA	O1D-CGD-O2D-CED
22	6	308	CLA	O1D-CGD-O2D-CED
22	6	311	CLA	O1D-CGD-O2D-CED
22	9	310	CLA	O1D-CGD-O2D-CED
33	F	307	LMT	O5B-C1B-O1B-C4'
22	A	811	CLA	O1D-CGD-O2D-CED
22	A	823	CLA	O1D-CGD-O2D-CED
22	A	825	CLA	O1D-CGD-O2D-CED
22	A	842	CLA	O1D-CGD-O2D-CED
22	B	805	CLA	O1D-CGD-O2D-CED
22	B	829	CLA	O1D-CGD-O2D-CED
22	J	103	CLA	O1D-CGD-O2D-CED
22	K	204	CLA	O1D-CGD-O2D-CED
22	L	203	CLA	O1D-CGD-O2D-CED
22	3	309	CLA	O1D-CGD-O2D-CED
22	3	314	CLA	O1D-CGD-O2D-CED
22	7	308	CLA	O1D-CGD-O2D-CED
22	7	312	CLA	O1D-CGD-O2D-CED

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Mol	Chain	Res	Type	Atoms
22	8	310	CLA	O1D-CGD-O2D-CED
22	4	809	CLA	O1D-CGD-O2D-CED
22	4	817	CLA	O1D-CGD-O2D-CED
22	5	309	CLA	O1D-CGD-O2D-CED
22	5	315	CLA	O1D-CGD-O2D-CED
22	9	303	CLA	O1D-CGD-O2D-CED
22	A	802	CLA	CBD-CGD-O2D-CED
22	A	804	CLA	CBD-CGD-O2D-CED
22	A	808	CLA	CBD-CGD-O2D-CED
22	A	809	CLA	CBD-CGD-O2D-CED
22	A	812	CLA	CBD-CGD-O2D-CED
22	A	814	CLA	CBD-CGD-O2D-CED
22	A	819	CLA	CBD-CGD-O2D-CED
22	A	821	CLA	CBD-CGD-O2D-CED
22	A	825	CLA	CBD-CGD-O2D-CED
22	A	832	CLA	CBD-CGD-O2D-CED
22	A	834	CLA	CBD-CGD-O2D-CED
22	A	836	CLA	CBD-CGD-O2D-CED
22	A	838	CLA	CBD-CGD-O2D-CED
22	B	807	CLA	CBD-CGD-O2D-CED
22	B	809	CLA	CBD-CGD-O2D-CED
22	B	811	CLA	CBD-CGD-O2D-CED
22	B	818	CLA	CBD-CGD-O2D-CED
22	B	819	CLA	CBD-CGD-O2D-CED
22	B	820	CLA	CBD-CGD-O2D-CED
22	B	821	CLA	CBD-CGD-O2D-CED
22	B	823	CLA	CBD-CGD-O2D-CED
22	B	827	CLA	CBD-CGD-O2D-CED
22	B	828	CLA	CBD-CGD-O2D-CED
22	B	829	CLA	CBD-CGD-O2D-CED
22	B	836	CLA	CBD-CGD-O2D-CED
22	B	840	CLA	CBD-CGD-O2D-CED
22	F	301	CLA	CBD-CGD-O2D-CED
22	F	304	CLA	CBD-CGD-O2D-CED
22	F	305	CLA	CBD-CGD-O2D-CED
22	G	201	CLA	CBD-CGD-O2D-CED
22	G	202	CLA	CBD-CGD-O2D-CED
22	J	103	CLA	CBD-CGD-O2D-CED
22	K	204	CLA	CBD-CGD-O2D-CED
22	L	203	CLA	CBD-CGD-O2D-CED
22	1	305	CLA	CBD-CGD-O2D-CED
22	1	307	CLA	CBD-CGD-O2D-CED

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Mol	Chain	Res	Type	Atoms
22	1	309	CLA	CBD-CGD-O2D-CED
22	1	311	CLA	CBD-CGD-O2D-CED
22	1	315	CLA	CBD-CGD-O2D-CED
22	1	316	CLA	CBD-CGD-O2D-CED
22	Z	304	CLA	CBD-CGD-O2D-CED
22	Z	306	CLA	CBD-CGD-O2D-CED
22	Z	307	CLA	CBD-CGD-O2D-CED
22	Z	313	CLA	CBD-CGD-O2D-CED
22	3	308	CLA	CBD-CGD-O2D-CED
22	3	314	CLA	CBD-CGD-O2D-CED
22	3	318	CLA	CBD-CGD-O2D-CED
22	3	319	CLA	CBD-CGD-O2D-CED
22	3	320	CLA	CBD-CGD-O2D-CED
22	7	304	CLA	CBD-CGD-O2D-CED
22	7	308	CLA	CBD-CGD-O2D-CED
22	7	315	CLA	CBD-CGD-O2D-CED
22	7	324	CLA	CBD-CGD-O2D-CED
22	8	306	CLA	CBD-CGD-O2D-CED
22	8	307	CLA	CBD-CGD-O2D-CED
22	8	309	CLA	CBD-CGD-O2D-CED
22	8	316	CLA	CBD-CGD-O2D-CED
22	4	806	CLA	CBD-CGD-O2D-CED
22	4	811	CLA	CBD-CGD-O2D-CED
22	4	812	CLA	CBD-CGD-O2D-CED
22	4	817	CLA	CBD-CGD-O2D-CED
22	4	818	CLA	CBD-CGD-O2D-CED
22	5	309	CLA	CBD-CGD-O2D-CED
22	5	319	CLA	CBD-CGD-O2D-CED
22	5	322	CLA	CBD-CGD-O2D-CED
22	6	301	CLA	CBD-CGD-O2D-CED
22	6	308	CLA	CBD-CGD-O2D-CED
22	6	309	CLA	CBD-CGD-O2D-CED
22	6	317	CLA	CBD-CGD-O2D-CED
22	6	322	CLA	CBD-CGD-O2D-CED
22	9	303	CLA	CBD-CGD-O2D-CED
22	9	304	CLA	CBD-CGD-O2D-CED
22	9	305	CLA	CBD-CGD-O2D-CED
22	9	306	CLA	CBD-CGD-O2D-CED
22	9	307	CLA	CBD-CGD-O2D-CED
22	9	310	CLA	CBD-CGD-O2D-CED
22	9	313	CLA	CBD-CGD-O2D-CED
22	2	301	CLA	CBD-CGD-O2D-CED

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Mol	Chain	Res	Type	Atoms
22	2	302	CLA	CBD-CGD-O2D-CED
39	7	313	CHL	CBD-CGD-O2D-CED
39	4	813	CHL	CBD-CGD-O2D-CED
22	A	816	CLA	O1A-CGA-O2A-C1
22	A	826	CLA	O1A-CGA-O2A-C1
22	A	831	CLA	O1A-CGA-O2A-C1
22	A	841	CLA	O1A-CGA-O2A-C1
22	A	855	CLA	O1A-CGA-O2A-C1
22	B	805	CLA	O1A-CGA-O2A-C1
22	B	808	CLA	O1A-CGA-O2A-C1
22	B	814	CLA	O1A-CGA-O2A-C1
22	B	823	CLA	O1A-CGA-O2A-C1
22	B	832	CLA	O1A-CGA-O2A-C1
22	B	839	CLA	O1A-CGA-O2A-C1
22	K	204	CLA	O1A-CGA-O2A-C1
22	Z	308	CLA	O1A-CGA-O2A-C1
22	Z	309	CLA	O1A-CGA-O2A-C1
22	7	315	CLA	O1A-CGA-O2A-C1
22	7	322	CLA	O1A-CGA-O2A-C1
22	8	309	CLA	O1A-CGA-O2A-C1
22	5	307	CLA	O1A-CGA-O2A-C1
22	5	313	CLA	O1A-CGA-O2A-C1
22	6	314	CLA	O1A-CGA-O2A-C1
22	9	313	CLA	O1A-CGA-O2A-C1
25	6	323	LHG	O10-C23-O8-C6
34	J	101	LMG	O10-C28-O8-C9
39	5	316	CHL	O1A-CGA-O2A-C1
39	9	312	CHL	O1A-CGA-O2A-C1
22	K	202	CLA	O1A-CGA-O2A-C1
22	8	316	CLA	O1A-CGA-O2A-C1
22	2	304	CLA	O1A-CGA-O2A-C1
33	F	307	LMT	C2B-C1B-O1B-C4'
22	A	802	CLA	O1D-CGD-O2D-CED
22	A	821	CLA	O1D-CGD-O2D-CED
22	A	832	CLA	O1D-CGD-O2D-CED
22	B	827	CLA	O1D-CGD-O2D-CED
22	B	828	CLA	O1D-CGD-O2D-CED
22	B	836	CLA	O1D-CGD-O2D-CED
22	F	301	CLA	O1D-CGD-O2D-CED
22	Z	305	CLA	O1D-CGD-O2D-CED
22	Z	314	CLA	O1D-CGD-O2D-CED
22	Z	316	CLA	O1D-CGD-O2D-CED

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Mol	Chain	Res	Type	Atoms
22	3	310	CLA	O1D-CGD-O2D-CED
22	3	319	CLA	O1D-CGD-O2D-CED
22	5	320	CLA	O1D-CGD-O2D-CED
22	2	301	CLA	O1D-CGD-O2D-CED
22	7	308	CLA	C8-C10-C11-C12
22	K	202	CLA	CBA-CGA-O2A-C1
22	1	311	CLA	CBA-CGA-O2A-C1
22	1	315	CLA	CBA-CGA-O2A-C1
22	Z	315	CLA	CBA-CGA-O2A-C1
22	3	309	CLA	CBA-CGA-O2A-C1
22	8	316	CLA	CBA-CGA-O2A-C1
39	9	312	CHL	C4C-C3C-CAC-CBC
22	A	818	CLA	O1D-CGD-O2D-CED
22	A	820	CLA	O1D-CGD-O2D-CED
22	B	801	CLA	O1D-CGD-O2D-CED
22	B	806	CLA	O1D-CGD-O2D-CED
22	B	808	CLA	O1D-CGD-O2D-CED
22	B	831	CLA	O1D-CGD-O2D-CED
22	B	841	CLA	O1D-CGD-O2D-CED
22	F	304	CLA	O1D-CGD-O2D-CED
22	K	205	CLA	O1D-CGD-O2D-CED
22	1	306	CLA	O1D-CGD-O2D-CED
22	1	313	CLA	O1D-CGD-O2D-CED
22	1	316	CLA	O1D-CGD-O2D-CED
22	Z	303	CLA	O1D-CGD-O2D-CED
22	3	312	CLA	O1D-CGD-O2D-CED
22	3	315	CLA	O1D-CGD-O2D-CED
22	3	316	CLA	O1D-CGD-O2D-CED
22	7	324	CLA	O1D-CGD-O2D-CED
22	4	815	CLA	O1D-CGD-O2D-CED
22	5	301	CLA	O1D-CGD-O2D-CED
22	5	323	CLA	O1D-CGD-O2D-CED
22	6	317	CLA	O1D-CGD-O2D-CED
22	9	313	CLA	O1D-CGD-O2D-CED
39	Z	312	CHL	O1D-CGD-O2D-CED
39	6	318	CHL	O1D-CGD-O2D-CED
22	A	816	CLA	CBA-CGA-O2A-C1
22	A	841	CLA	CBA-CGA-O2A-C1
22	B	805	CLA	CBA-CGA-O2A-C1
22	B	812	CLA	CBA-CGA-O2A-C1
22	B	839	CLA	CBA-CGA-O2A-C1
22	K	204	CLA	CBA-CGA-O2A-C1

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Mol	Chain	Res	Type	Atoms
22	Z	308	CLA	CBA-CGA-O2A-C1
22	Z	309	CLA	CBA-CGA-O2A-C1
22	7	315	CLA	CBA-CGA-O2A-C1
22	7	322	CLA	CBA-CGA-O2A-C1
22	4	812	CLA	CBA-CGA-O2A-C1
22	6	314	CLA	CBA-CGA-O2A-C1
34	J	101	LMG	C29-C28-O8-C9
39	9	312	CHL	CBA-CGA-O2A-C1
25	3	321	LHG	C24-C23-O8-C6
22	A	803	CLA	CBD-CGD-O2D-CED
22	A	830	CLA	CBD-CGD-O2D-CED
22	A	831	CLA	CBD-CGD-O2D-CED
22	A	841	CLA	CBD-CGD-O2D-CED
22	B	813	CLA	CBD-CGD-O2D-CED
22	B	817	CLA	CBD-CGD-O2D-CED
22	B	833	CLA	CBD-CGD-O2D-CED
22	B	834	CLA	CBD-CGD-O2D-CED
22	B	837	CLA	CBD-CGD-O2D-CED
22	B	842	CLA	CBD-CGD-O2D-CED
22	K	202	CLA	CBD-CGD-O2D-CED
22	1	308	CLA	CBD-CGD-O2D-CED
22	7	311	CLA	CBD-CGD-O2D-CED
22	8	311	CLA	CBD-CGD-O2D-CED
22	4	805	CLA	CBD-CGD-O2D-CED
22	4	808	CLA	CBD-CGD-O2D-CED
22	5	307	CLA	CBD-CGD-O2D-CED
22	5	311	CLA	CBD-CGD-O2D-CED
39	5	316	CHL	CBD-CGD-O2D-CED
39	6	315	CHL	CBD-CGD-O2D-CED
39	Z	311	CHL	C2C-C3C-CAC-CBC
39	9	312	CHL	C2C-C3C-CAC-CBC
22	A	807	CLA	O1A-CGA-O2A-C1
22	A	812	CLA	O1A-CGA-O2A-C1
22	A	814	CLA	O1A-CGA-O2A-C1
22	A	815	CLA	O1A-CGA-O2A-C1
22	A	819	CLA	O1A-CGA-O2A-C1
22	A	821	CLA	O1A-CGA-O2A-C1
22	A	822	CLA	O1A-CGA-O2A-C1
22	A	823	CLA	O1A-CGA-O2A-C1
22	A	827	CLA	O1A-CGA-O2A-C1
22	A	837	CLA	O1A-CGA-O2A-C1
22	B	819	CLA	O1A-CGA-O2A-C1

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Mol	Chain	Res	Type	Atoms
22	B	821	CLA	O1A-CGA-O2A-C1
22	B	834	CLA	O1A-CGA-O2A-C1
22	1	307	CLA	O1A-CGA-O2A-C1
22	1	309	CLA	O1A-CGA-O2A-C1
22	Z	307	CLA	O1A-CGA-O2A-C1
22	3	308	CLA	O1A-CGA-O2A-C1
22	7	310	CLA	O1A-CGA-O2A-C1
22	7	324	CLA	O1A-CGA-O2A-C1
22	8	305	CLA	O1A-CGA-O2A-C1
22	4	805	CLA	O1A-CGA-O2A-C1
22	4	810	CLA	O1A-CGA-O2A-C1
22	4	812	CLA	O1A-CGA-O2A-C1
22	4	818	CLA	O1A-CGA-O2A-C1
22	5	301	CLA	O1A-CGA-O2A-C1
22	6	312	CLA	O1A-CGA-O2A-C1
22	6	313	CLA	O1A-CGA-O2A-C1
22	9	309	CLA	O1A-CGA-O2A-C1
40	1	318	SQD	O10-C23-O48-C46
41	7	319	3PH	O32-C31-O31-C3
22	1	311	CLA	O1A-CGA-O2A-C1
22	1	315	CLA	O1A-CGA-O2A-C1
22	Z	315	CLA	O1A-CGA-O2A-C1
22	3	309	CLA	O1A-CGA-O2A-C1
22	6	321	CLA	O1A-CGA-O2A-C1
22	9	311	CLA	O1A-CGA-O2A-C1
22	A	817	CLA	O1D-CGD-O2D-CED
22	A	828	CLA	O1D-CGD-O2D-CED
22	B	812	CLA	O1D-CGD-O2D-CED
22	B	814	CLA	O1D-CGD-O2D-CED
22	B	824	CLA	O1D-CGD-O2D-CED
22	7	305	CLA	O1D-CGD-O2D-CED
22	7	316	CLA	O1D-CGD-O2D-CED
22	8	305	CLA	O1D-CGD-O2D-CED
22	8	313	CLA	O1D-CGD-O2D-CED
22	5	326	CLA	O1D-CGD-O2D-CED
22	6	307	CLA	O1D-CGD-O2D-CED
22	9	308	CLA	O1D-CGD-O2D-CED
22	9	311	CLA	O1D-CGD-O2D-CED
39	8	312	CHL	O1D-CGD-O2D-CED
39	9	312	CHL	O1D-CGD-O2D-CED
22	B	811	CLA	C5-C6-C7-C8
22	B	812	CLA	C5-C6-C7-C8

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Mol	Chain	Res	Type	Atoms
22	7	306	CLA	C5-C6-C7-C8
22	A	807	CLA	O1D-CGD-O2D-CED
22	A	833	CLA	O1D-CGD-O2D-CED
22	A	855	CLA	O1D-CGD-O2D-CED
22	B	816	CLA	O1D-CGD-O2D-CED
22	B	819	CLA	O1D-CGD-O2D-CED
22	Z	308	CLA	O1D-CGD-O2D-CED
22	8	309	CLA	O1D-CGD-O2D-CED
22	5	308	CLA	O1D-CGD-O2D-CED
22	B	826	CLA	CBD-CGD-O2D-CED
22	A	810	CLA	C8-C10-C11-C12
22	B	838	CLA	C8-C10-C11-C12
23	A	843	PQN	C18-C20-C21-C22
22	A	816	CLA	O1D-CGD-O2D-CED
22	A	835	CLA	O1D-CGD-O2D-CED
22	B	821	CLA	O1D-CGD-O2D-CED
22	F	305	CLA	O1D-CGD-O2D-CED
22	G	201	CLA	O1D-CGD-O2D-CED
22	Z	313	CLA	O1D-CGD-O2D-CED
22	9	304	CLA	O1D-CGD-O2D-CED
39	4	814	CHL	O1D-CGD-O2D-CED
39	2	303	CHL	O1D-CGD-O2D-CED
25	1	317	LHG	O9-C7-O7-C5
34	F	308	LMG	O9-C10-O7-C8
34	J	101	LMG	O9-C10-O7-C8
41	7	319	3PH	O22-C21-O21-C2
41	6	324	3PH	O22-C21-O21-C2
22	1	306	CLA	O1A-CGA-O2A-C1
22	8	314	CLA	CBA-CGA-O2A-C1
22	9	311	CLA	CBA-CGA-O2A-C1
39	Z	311	CHL	C4C-C3C-CAC-CBC
31	3	301	DGD	C2G-C1G-O1G-C1A
22	A	802	CLA	C3-C5-C6-C7
22	A	805	CLA	C3-C5-C6-C7
22	A	818	CLA	C3-C5-C6-C7
22	A	820	CLA	C3-C5-C6-C7
22	A	855	CLA	C3-C5-C6-C7
22	B	801	CLA	C3-C5-C6-C7
22	B	816	CLA	C3-C5-C6-C7
22	B	819	CLA	C3-C5-C6-C7
22	B	827	CLA	C3-C5-C6-C7
22	B	828	CLA	C3-C5-C6-C7

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Mol	Chain	Res	Type	Atoms
22	B	829	CLA	C3-C5-C6-C7
22	F	302	CLA	C3-C5-C6-C7
22	L	202	CLA	C3-C5-C6-C7
22	1	314	CLA	C3-C5-C6-C7
22	Z	306	CLA	C3-C5-C6-C7
22	Z	307	CLA	C3-C5-C6-C7
22	3	314	CLA	C3-C5-C6-C7
22	8	307	CLA	C3-C5-C6-C7
22	8	310	CLA	C3-C5-C6-C7
22	5	315	CLA	C3-C5-C6-C7
22	5	322	CLA	C3-C5-C6-C7
22	6	311	CLA	C3-C5-C6-C7
22	6	313	CLA	C3-C5-C6-C7
22	9	305	CLA	C3-C5-C6-C7
22	A	807	CLA	CBA-CGA-O2A-C1
22	A	808	CLA	CBA-CGA-O2A-C1
22	A	814	CLA	CBA-CGA-O2A-C1
22	A	821	CLA	CBA-CGA-O2A-C1
22	A	823	CLA	CBA-CGA-O2A-C1
22	A	831	CLA	CBA-CGA-O2A-C1
22	A	837	CLA	CBA-CGA-O2A-C1
22	A	855	CLA	CBA-CGA-O2A-C1
22	B	808	CLA	CBA-CGA-O2A-C1
22	B	813	CLA	CBA-CGA-O2A-C1
22	B	814	CLA	CBA-CGA-O2A-C1
22	B	823	CLA	CBA-CGA-O2A-C1
22	B	831	CLA	CBA-CGA-O2A-C1
22	B	832	CLA	CBA-CGA-O2A-C1
22	1	307	CLA	CBA-CGA-O2A-C1
22	1	309	CLA	CBA-CGA-O2A-C1
22	Z	316	CLA	CBA-CGA-O2A-C1
22	3	316	CLA	CBA-CGA-O2A-C1
22	7	324	CLA	CBA-CGA-O2A-C1
22	8	305	CLA	CBA-CGA-O2A-C1
22	8	309	CLA	CBA-CGA-O2A-C1
22	4	810	CLA	CBA-CGA-O2A-C1
22	4	818	CLA	CBA-CGA-O2A-C1
22	5	301	CLA	CBA-CGA-O2A-C1
22	5	307	CLA	CBA-CGA-O2A-C1
22	5	326	CLA	CBA-CGA-O2A-C1
22	6	312	CLA	CBA-CGA-O2A-C1
26	A	851	NKP	CAL-CAK-OAJ-CAI

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Mol	Chain	Res	Type	Atoms
39	5	316	CHL	CBA-CGA-O2A-C1
40	1	318	SQD	C24-C23-O48-C46
41	7	319	3PH	C32-C31-O31-C3
25	B	850	LHG	C24-C23-O8-C6
25	B	851	LHG	C24-C23-O8-C6
22	A	814	CLA	C8-C10-C11-C12
22	B	831	CLA	C5-C6-C7-C8
25	B	850	LHG	C8-C7-O7-C5
31	B	852	DGD	C2B-C1B-O2G-C2G
34	J	101	LMG	C11-C10-O7-C8
22	B	840	CLA	O1D-CGD-O2D-CED
22	3	318	CLA	O1D-CGD-O2D-CED
22	5	322	CLA	O1D-CGD-O2D-CED
22	6	309	CLA	O1D-CGD-O2D-CED
22	6	322	CLA	O1D-CGD-O2D-CED
22	A	815	CLA	CBD-CGD-O2D-CED
22	B	825	CLA	CBD-CGD-O2D-CED
22	1	304	CLA	CBD-CGD-O2D-CED
22	Z	310	CLA	CBD-CGD-O2D-CED
22	7	310	CLA	CBD-CGD-O2D-CED
22	5	314	CLA	CBD-CGD-O2D-CED
39	6	316	CHL	CBD-CGD-O2D-CED
22	8	314	CLA	O1A-CGA-O2A-C1
39	4	813	CHL	O1D-CGD-O2D-CED
22	3	320	CLA	CBA-CGA-O2A-C1
22	6	308	CLA	C3-C5-C6-C7
22	3	319	CLA	C2-C3-C5-C6
22	L	201	CLA	CBD-CGD-O2D-CED
22	3	313	CLA	CBD-CGD-O2D-CED
22	2	304	CLA	CBD-CGD-O2D-CED
22	B	839	CLA	C2A-CAA-CBA-CGA
22	K	204	CLA	C2A-CAA-CBA-CGA
22	K	205	CLA	C2A-CAA-CBA-CGA
22	3	310	CLA	C2A-CAA-CBA-CGA
22	3	315	CLA	C2A-CAA-CBA-CGA
22	3	320	CLA	C2A-CAA-CBA-CGA
22	8	309	CLA	C2A-CAA-CBA-CGA
22	4	810	CLA	C2A-CAA-CBA-CGA
22	5	313	CLA	C2A-CAA-CBA-CGA
22	5	318	CLA	C2A-CAA-CBA-CGA
22	9	305	CLA	C2A-CAA-CBA-CGA
22	A	836	CLA	O1A-CGA-O2A-C1

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Mol	Chain	Res	Type	Atoms
22	B	840	CLA	O1A-CGA-O2A-C1
22	5	312	CLA	O1A-CGA-O2A-C1
22	G	202	CLA	O1D-CGD-O2D-CED
22	Z	306	CLA	O1D-CGD-O2D-CED
22	3	320	CLA	O1D-CGD-O2D-CED
25	Z	317	LHG	C11-C12-C13-C14
31	B	852	DGD	C8A-C9A-CAA-CBA
31	3	301	DGD	C8A-C9A-CAA-CBA
22	A	803	CLA	C3-C5-C6-C7
22	A	812	CLA	C3-C5-C6-C7
22	A	828	CLA	C3-C5-C6-C7
22	A	836	CLA	C3-C5-C6-C7
22	A	841	CLA	C3-C5-C6-C7
22	B	815	CLA	C3-C5-C6-C7
22	1	313	CLA	C3-C5-C6-C7
22	Z	313	CLA	C3-C5-C6-C7
22	3	308	CLA	C3-C5-C6-C7
22	3	310	CLA	C3-C5-C6-C7
22	3	312	CLA	C3-C5-C6-C7
22	7	312	CLA	C3-C5-C6-C7
22	7	324	CLA	C3-C5-C6-C7
22	8	308	CLA	C3-C5-C6-C7
22	9	307	CLA	C3-C5-C6-C7
22	A	812	CLA	CBA-CGA-O2A-C1
22	A	815	CLA	CBA-CGA-O2A-C1
22	A	819	CLA	CBA-CGA-O2A-C1
22	A	822	CLA	CBA-CGA-O2A-C1
22	A	827	CLA	CBA-CGA-O2A-C1
22	A	828	CLA	CBA-CGA-O2A-C1
22	A	832	CLA	CBA-CGA-O2A-C1
22	B	819	CLA	CBA-CGA-O2A-C1
22	B	820	CLA	CBA-CGA-O2A-C1
22	B	821	CLA	CBA-CGA-O2A-C1
22	B	834	CLA	CBA-CGA-O2A-C1
22	B	840	CLA	CBA-CGA-O2A-C1
22	1	304	CLA	CBA-CGA-O2A-C1
22	1	306	CLA	CBA-CGA-O2A-C1
22	1	310	CLA	CBA-CGA-O2A-C1
22	Z	307	CLA	CBA-CGA-O2A-C1
22	Z	310	CLA	CBA-CGA-O2A-C1
22	3	308	CLA	CBA-CGA-O2A-C1
22	7	310	CLA	CBA-CGA-O2A-C1

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Mol	Chain	Res	Type	Atoms
22	7	312	CLA	CBA-CGA-O2A-C1
22	8	311	CLA	CBA-CGA-O2A-C1
22	4	805	CLA	CBA-CGA-O2A-C1
22	4	806	CLA	CBA-CGA-O2A-C1
22	6	301	CLA	CBA-CGA-O2A-C1
22	6	311	CLA	CBA-CGA-O2A-C1
22	6	313	CLA	CBA-CGA-O2A-C1
22	9	306	CLA	CBA-CGA-O2A-C1
22	9	309	CLA	CBA-CGA-O2A-C1
39	2	303	CHL	CBA-CGA-O2A-C1
22	A	804	CLA	O1D-CGD-O2D-CED
22	A	809	CLA	O1D-CGD-O2D-CED
22	4	812	CLA	O1D-CGD-O2D-CED
22	4	807	CLA	CBD-CGD-O2D-CED
22	A	814	CLA	O1D-CGD-O2D-CED
22	A	838	CLA	O1D-CGD-O2D-CED
22	B	818	CLA	O1D-CGD-O2D-CED
22	1	311	CLA	O1D-CGD-O2D-CED
22	Z	307	CLA	O1D-CGD-O2D-CED
22	7	304	CLA	O1D-CGD-O2D-CED
22	7	315	CLA	O1D-CGD-O2D-CED
22	4	806	CLA	O1D-CGD-O2D-CED
22	9	307	CLA	O1D-CGD-O2D-CED
25	4	820	LHG	O9-C7-O7-C5
33	G	204	LMT	C4'-C5'-C6'-O6'
22	A	828	CLA	O1A-CGA-O2A-C1
22	A	832	CLA	O1A-CGA-O2A-C1
22	A	842	CLA	O1A-CGA-O2A-C1
22	B	820	CLA	O1A-CGA-O2A-C1
22	B	831	CLA	O1A-CGA-O2A-C1
22	1	304	CLA	O1A-CGA-O2A-C1
22	1	310	CLA	O1A-CGA-O2A-C1
22	Z	306	CLA	O1A-CGA-O2A-C1
22	Z	310	CLA	O1A-CGA-O2A-C1
22	3	316	CLA	O1A-CGA-O2A-C1
22	7	312	CLA	O1A-CGA-O2A-C1
22	4	806	CLA	O1A-CGA-O2A-C1
22	5	322	CLA	O1A-CGA-O2A-C1
22	6	301	CLA	O1A-CGA-O2A-C1
22	9	306	CLA	O1A-CGA-O2A-C1
26	A	851	NKP	OAE-CAK-OAJ-CAI
39	7	313	CHL	O1A-CGA-O2A-C1

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Mol	Chain	Res	Type	Atoms
39	2	303	CHL	O1A-CGA-O2A-C1
22	3	320	CLA	O1A-CGA-O2A-C1
24	A	845	BCR	C9-C10-C11-C12
24	A	845	BCR	C19-C20-C21-C22
24	A	846	BCR	C9-C10-C11-C12
24	A	846	BCR	C15-C16-C17-C18
24	B	848	BCR	C15-C16-C17-C18
24	B	849	BCR	C19-C20-C21-C22
24	B	853	BCR	C15-C16-C17-C18
24	G	203	BCR	C19-C20-C21-C22
24	I	4001	BCR	C13-C14-C15-C16
24	J	104	BCR	C19-C20-C21-C22
24	K	206	BCR	C15-C16-C17-C18
24	L	204	BCR	C9-C10-C11-C12
24	7	303	BCR	C9-C10-C11-C12
24	7	303	BCR	C19-C20-C21-C22
24	4	804	BCR	C13-C14-C15-C16
24	5	304	BCR	C9-C10-C11-C12
24	6	305	BCR	C9-C10-C11-C12
24	6	305	BCR	C13-C14-C15-C16
24	6	306	BCR	C13-C14-C15-C16
24	6	306	BCR	C19-C20-C21-C22
36	5	306	C7Z	C29-C30-C31-C32
22	B	822	CLA	CBD-CGD-O2D-CED
22	B	838	CLA	CBD-CGD-O2D-CED
22	3	322	CLA	CBD-CGD-O2D-CED
22	6	310	CLA	CBD-CGD-O2D-CED
22	B	820	CLA	O1D-CGD-O2D-CED
22	1	309	CLA	O1D-CGD-O2D-CED
39	7	313	CHL	O1D-CGD-O2D-CED
25	A	849	LHG	O2-C2-C3-O3
25	A	850	LHG	O2-C2-C3-O3
25	B	803	LHG	O2-C2-C3-O3
25	B	850	LHG	O2-C2-C3-O3
25	3	321	LHG	O2-C2-C3-O3
25	9	315	LHG	O2-C2-C3-O3
21	A	801	CL0	C3-C5-C6-C7
22	A	809	CLA	C3-C5-C6-C7
22	A	816	CLA	C3-C5-C6-C7
22	A	821	CLA	C3-C5-C6-C7
22	B	810	CLA	C3-C5-C6-C7
22	B	822	CLA	C3-C5-C6-C7

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Mol	Chain	Res	Type	Atoms
22	B	823	CLA	C3-C5-C6-C7
22	B	825	CLA	C3-C5-C6-C7
22	1	308	CLA	C3-C5-C6-C7
22	3	319	CLA	C3-C5-C6-C7
22	5	318	CLA	C3-C5-C6-C7
22	6	302	CLA	C3-C5-C6-C7
22	6	309	CLA	C3-C5-C6-C7
22	6	319	CLA	C3-C5-C6-C7
22	6	322	CLA	C3-C5-C6-C7
22	2	301	CLA	C3-C5-C6-C7
22	A	833	CLA	CBA-CGA-O2A-C1
22	A	836	CLA	CBA-CGA-O2A-C1
22	A	839	CLA	CBA-CGA-O2A-C1
22	B	816	CLA	CBA-CGA-O2A-C1
22	F	302	CLA	CBA-CGA-O2A-C1
22	Z	306	CLA	CBA-CGA-O2A-C1
22	3	311	CLA	CBA-CGA-O2A-C1
22	4	808	CLA	CBA-CGA-O2A-C1
22	5	312	CLA	CBA-CGA-O2A-C1
39	7	313	CHL	CBA-CGA-O2A-C1
22	A	808	CLA	O1A-CGA-O2A-C1
22	B	833	CLA	O1A-CGA-O2A-C1
22	3	311	CLA	O1A-CGA-O2A-C1
22	5	326	CLA	O1A-CGA-O2A-C1
22	6	311	CLA	O1A-CGA-O2A-C1
22	A	808	CLA	O1D-CGD-O2D-CED
22	A	834	CLA	O1D-CGD-O2D-CED
22	A	836	CLA	O1D-CGD-O2D-CED
22	B	807	CLA	O1D-CGD-O2D-CED
22	B	823	CLA	O1D-CGD-O2D-CED
22	Z	304	CLA	O1D-CGD-O2D-CED
22	8	307	CLA	O1D-CGD-O2D-CED
22	9	305	CLA	O1D-CGD-O2D-CED
22	2	302	CLA	O1D-CGD-O2D-CED
22	A	825	CLA	C8-C10-C11-C12
25	7	318	LHG	C8-C7-O7-C5
35	J	102	T7X	C12-C10-O16-C8
25	Z	317	LHG	C23-C24-C25-C26
22	A	819	CLA	O1D-CGD-O2D-CED
22	A	810	CLA	CBD-CGD-O2D-CED
22	1	310	CLA	CBD-CGD-O2D-CED
22	7	323	CLA	CBD-CGD-O2D-CED

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Mol	Chain	Res	Type	Atoms
22	8	314	CLA	CBD-CGD-O2D-CED
22	5	313	CLA	CBD-CGD-O2D-CED
22	B	816	CLA	O1A-CGA-O2A-C1
22	1	307	CLA	O1D-CGD-O2D-CED
22	A	837	CLA	CBD-CGD-O2D-CED
22	A	807	CLA	C3-C5-C6-C7
22	A	814	CLA	C3-C5-C6-C7
22	B	821	CLA	C3-C5-C6-C7
22	B	838	CLA	C3-C5-C6-C7
22	F	301	CLA	C3-C5-C6-C7
22	7	322	CLA	C3-C5-C6-C7
22	4	812	CLA	C3-C5-C6-C7
22	5	301	CLA	C3-C5-C6-C7
22	5	310	CLA	C3-C5-C6-C7
22	A	842	CLA	CBA-CGA-O2A-C1
22	B	833	CLA	CBA-CGA-O2A-C1
22	5	322	CLA	CBA-CGA-O2A-C1
41	6	324	3PH	C22-C23-C24-C25
26	A	851	NKP	CAH-CAG-OAF-PAC
22	A	839	CLA	O1A-CGA-O2A-C1
22	F	302	CLA	O1A-CGA-O2A-C1
22	8	311	CLA	O1A-CGA-O2A-C1
22	4	808	CLA	O1A-CGA-O2A-C1
22	A	842	CLA	C3-C5-C6-C7
22	A	817	CLA	C8-C10-C11-C12
22	3	314	CLA	C4-C3-C5-C6
22	3	314	CLA	C2-C3-C5-C6
25	3	321	LHG	O10-C23-O8-C6
22	A	824	CLA	C2A-CAA-CBA-CGA
22	F	304	CLA	C2A-CAA-CBA-CGA
22	1	306	CLA	C2A-CAA-CBA-CGA
22	Z	303	CLA	C2A-CAA-CBA-CGA
22	9	308	CLA	C2A-CAA-CBA-CGA
22	B	811	CLA	O1D-CGD-O2D-CED
22	1	305	CLA	O1D-CGD-O2D-CED
22	8	306	CLA	O1D-CGD-O2D-CED
22	6	301	CLA	O1D-CGD-O2D-CED
22	9	306	CLA	O1D-CGD-O2D-CED
22	A	833	CLA	O1A-CGA-O2A-C1
22	L	203	CLA	O1A-CGA-O2A-C1
22	8	310	CLA	O1A-CGA-O2A-C1
22	A	840	CLA	CBA-CGA-O2A-C1

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Mol	Chain	Res	Type	Atoms
22	L	203	CLA	CBA-CGA-O2A-C1
22	7	317	CLA	CBA-CGA-O2A-C1
22	8	310	CLA	CBA-CGA-O2A-C1
22	9	308	CLA	CBA-CGA-O2A-C1
39	6	315	CHL	CBA-CGA-O2A-C1
22	A	840	CLA	CBD-CGD-O2D-CED
22	G	202	CLA	CBA-CGA-O2A-C1
22	A	812	CLA	O1D-CGD-O2D-CED
22	B	834	CLA	O1D-CGD-O2D-CED
22	4	805	CLA	O1D-CGD-O2D-CED
22	4	808	CLA	O1D-CGD-O2D-CED
22	4	811	CLA	O1D-CGD-O2D-CED
25	8	317	LHG	C25-C26-C27-C28
22	B	817	CLA	O1D-CGD-O2D-CED
22	B	833	CLA	O1D-CGD-O2D-CED
22	B	842	CLA	O1D-CGD-O2D-CED
22	7	311	CLA	O1D-CGD-O2D-CED
39	6	315	CHL	O1D-CGD-O2D-CED
22	A	806	CLA	CBD-CGD-O2D-CED
22	A	841	CLA	O1D-CGD-O2D-CED
22	5	311	CLA	O1D-CGD-O2D-CED
25	B	803	LHG	C1-C2-C3-O3
25	4	820	LHG	C1-C2-C3-O3
25	5	324	LHG	C1-C2-C3-O3
25	6	323	LHG	C1-C2-C3-O3
35	J	102	T7X	O17-C10-O16-C8
22	9	308	CLA	O1A-CGA-O2A-C1
22	A	817	CLA	C3-C5-C6-C7
22	A	830	CLA	C3-C5-C6-C7
22	B	811	CLA	C3-C5-C6-C7
22	7	310	CLA	C3-C5-C6-C7
22	9	306	CLA	C3-C5-C6-C7
22	B	837	CLA	O1D-CGD-O2D-CED
22	A	810	CLA	CBA-CGA-O2A-C1
22	A	824	CLA	CBA-CGA-O2A-C1
22	A	834	CLA	CBA-CGA-O2A-C1
22	B	809	CLA	CBA-CGA-O2A-C1
22	B	817	CLA	CBA-CGA-O2A-C1
22	K	205	CLA	CBA-CGA-O2A-C1
22	1	308	CLA	CBA-CGA-O2A-C1
22	7	309	CLA	CBA-CGA-O2A-C1
22	8	308	CLA	CBA-CGA-O2A-C1

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Mol	Chain	Res	Type	Atoms
22	5	319	CLA	CBA-CGA-O2A-C1
22	6	308	CLA	CBA-CGA-O2A-C1
22	9	307	CLA	CBA-CGA-O2A-C1
25	5	324	LHG	C24-C23-O8-C6
39	8	315	CHL	CBA-CGA-O2A-C1
22	B	839	CLA	CBD-CGD-O2D-CED
24	A	847	BCR	C15-C16-C17-C18
24	A	856	BCR	C13-C14-C15-C16
24	A	856	BCR	C19-C20-C21-C22
24	B	845	BCR	C15-C16-C17-C18
24	B	845	BCR	C19-C20-C21-C22
24	I	4001	BCR	C15-C16-C17-C18
24	7	303	BCR	C15-C16-C17-C18
24	8	304	BCR	C15-C16-C17-C18
24	8	304	BCR	C19-C20-C21-C22
24	4	804	BCR	C15-C16-C17-C18
24	4	804	BCR	C19-C20-C21-C22
24	6	306	BCR	C15-C16-C17-C18
22	9	307	CLA	O1A-CGA-O2A-C1
22	1	306	CLA	C5-C6-C7-C8
22	6	314	CLA	C8-C10-C11-C12
25	4	820	LHG	O2-C2-C3-O3
43	8	319	LPX	O1-C3-C4-O5
39	9	312	CHL	C3-C5-C6-C7
25	6	323	LHG	C7-C8-C9-C10
22	B	839	CLA	C3-C5-C6-C7
22	3	316	CLA	C3-C5-C6-C7
25	B	850	LHG	O10-C23-O8-C6
25	B	851	LHG	O10-C23-O8-C6
22	A	834	CLA	O1A-CGA-O2A-C1
22	A	840	CLA	O1A-CGA-O2A-C1
22	B	817	CLA	O1A-CGA-O2A-C1
22	K	205	CLA	O1A-CGA-O2A-C1
22	1	308	CLA	O1A-CGA-O2A-C1
33	G	204	LMT	O5'-C5'-C6'-O6'
22	A	802	CLA	C6-C7-C8-C9
22	A	803	CLA	C6-C7-C8-C9
22	A	807	CLA	C6-C7-C8-C9
22	A	808	CLA	C6-C7-C8-C9
22	A	812	CLA	C6-C7-C8-C9
22	A	812	CLA	C11-C10-C8-C9
22	A	814	CLA	C6-C7-C8-C9

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Mol	Chain	Res	Type	Atoms
22	A	815	CLA	C6-C7-C8-C9
22	A	816	CLA	C6-C7-C8-C9
22	A	818	CLA	C6-C7-C8-C9
22	A	818	CLA	C11-C10-C8-C9
22	A	820	CLA	C6-C7-C8-C9
22	A	823	CLA	C6-C7-C8-C9
22	A	823	CLA	C11-C10-C8-C9
22	A	830	CLA	C6-C7-C8-C9
22	A	832	CLA	C6-C7-C8-C9
22	A	834	CLA	C6-C7-C8-C9
22	A	835	CLA	C6-C7-C8-C9
22	A	838	CLA	C11-C10-C8-C9
22	A	854	CLA	C6-C7-C8-C9
22	A	855	CLA	C11-C12-C13-C14
22	B	801	CLA	C6-C7-C8-C9
22	B	809	CLA	C6-C7-C8-C9
22	B	813	CLA	C6-C7-C8-C9
22	B	815	CLA	C6-C7-C8-C9
22	B	818	CLA	C11-C12-C13-C14
22	B	819	CLA	C6-C7-C8-C9
22	B	820	CLA	C6-C7-C8-C9
22	B	824	CLA	C11-C10-C8-C9
22	B	825	CLA	C6-C7-C8-C9
22	B	828	CLA	C11-C10-C8-C9
22	B	829	CLA	C6-C7-C8-C9
22	B	830	CLA	C6-C7-C8-C9
22	B	833	CLA	C6-C7-C8-C9
22	B	842	CLA	C11-C10-C8-C9
22	F	304	CLA	C6-C7-C8-C9
22	L	202	CLA	C11-C12-C13-C14
22	1	304	CLA	C6-C7-C8-C9
22	1	306	CLA	C11-C12-C13-C14
22	1	307	CLA	C6-C7-C8-C9
22	1	309	CLA	C6-C7-C8-C9
22	1	310	CLA	C11-C10-C8-C9
22	1	313	CLA	C6-C7-C8-C9
22	1	316	CLA	C11-C10-C8-C9
22	Z	303	CLA	C6-C7-C8-C9
22	Z	313	CLA	C6-C7-C8-C9
22	3	312	CLA	C11-C10-C8-C9
22	3	314	CLA	C11-C10-C8-C9
22	3	318	CLA	C6-C7-C8-C9

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Mol	Chain	Res	Type	Atoms
22	3	322	CLA	C6-C7-C8-C9
22	3	322	CLA	C11-C10-C8-C9
22	3	322	CLA	C11-C12-C13-C14
22	7	304	CLA	C6-C7-C8-C9
22	7	307	CLA	C11-C10-C8-C9
22	7	312	CLA	C11-C10-C8-C9
22	7	324	CLA	C6-C7-C8-C9
22	7	324	CLA	C11-C10-C8-C9
22	8	308	CLA	C6-C7-C8-C9
22	4	807	CLA	C6-C7-C8-C9
22	5	313	CLA	C11-C10-C8-C9
22	5	318	CLA	C6-C7-C8-C9
22	5	322	CLA	C11-C10-C8-C9
22	6	307	CLA	C6-C7-C8-C9
22	6	312	CLA	C6-C7-C8-C9
22	9	305	CLA	C6-C7-C8-C9
39	3	317	CHL	C11-C10-C8-C9
39	5	316	CHL	C14-C13-C15-C16
22	A	830	CLA	O1D-CGD-O2D-CED
22	A	831	CLA	O1D-CGD-O2D-CED
22	7	307	CLA	CBD-CGD-O2D-CED
24	A	844	BCR	C11-C12-C13-C35
24	A	844	BCR	C36-C18-C19-C20
24	A	846	BCR	C11-C12-C13-C35
24	A	847	BCR	C7-C8-C9-C34
24	A	847	BCR	C36-C18-C19-C20
24	A	848	BCR	C11-C12-C13-C35
24	A	856	BCR	C7-C8-C9-C34
24	A	856	BCR	C37-C22-C23-C24
24	B	845	BCR	C11-C12-C13-C35
24	B	845	BCR	C36-C18-C19-C20
24	B	846	BCR	C7-C8-C9-C34
24	B	847	BCR	C11-C12-C13-C35
24	B	848	BCR	C7-C8-C9-C34
24	B	853	BCR	C11-C12-C13-C35
24	B	853	BCR	C37-C22-C23-C24
24	F	303	BCR	C37-C22-C23-C24
24	G	203	BCR	C37-C22-C23-C24
24	I	4001	BCR	C7-C8-C9-C34
24	I	4001	BCR	C11-C12-C13-C35
24	J	104	BCR	C7-C8-C9-C34
24	K	206	BCR	C11-C12-C13-C35

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Mol	Chain	Res	Type	Atoms
24	K	206	BCR	C37-C22-C23-C24
24	L	204	BCR	C7-C8-C9-C34
24	3	304	BCR	C37-C22-C23-C24
24	3	305	BCR	C7-C8-C9-C34
24	3	306	BCR	C36-C18-C19-C20
24	3	307	BCR	C7-C8-C9-C34
24	3	307	BCR	C37-C22-C23-C24
24	7	303	BCR	C36-C18-C19-C20
24	4	804	BCR	C36-C18-C19-C20
24	5	304	BCR	C36-C18-C19-C20
24	5	305	BCR	C37-C22-C23-C24
24	6	305	BCR	C7-C8-C9-C34
24	6	305	BCR	C36-C18-C19-C20
24	6	306	BCR	C11-C12-C13-C35
24	6	306	BCR	C36-C18-C19-C20
36	J	105	C7Z	C7-C8-C9-C19
36	J	105	C7Z	C11-C12-C13-C20
36	5	306	C7Z	C7-C8-C9-C19
36	5	306	C7Z	C27-C28-C29-C39
38	3	303	LUT	C7-C8-C9-C19
38	7	302	LUT	C11-C12-C13-C20
38	5	302	LUT	C27-C28-C29-C39
38	5	303	LUT	C7-C8-C9-C19
38	6	304	LUT	C7-C8-C9-C19
24	A	844	BCR	C11-C12-C13-C14
24	A	844	BCR	C17-C18-C19-C20
24	A	845	BCR	C7-C8-C9-C10
24	A	846	BCR	C11-C12-C13-C14
24	A	847	BCR	C17-C18-C19-C20
24	A	856	BCR	C7-C8-C9-C10
24	A	856	BCR	C21-C22-C23-C24
24	B	844	BCR	C7-C8-C9-C10
24	B	845	BCR	C21-C22-C23-C24
24	B	848	BCR	C7-C8-C9-C10
24	B	849	BCR	C17-C18-C19-C20
24	B	853	BCR	C11-C12-C13-C14
24	B	853	BCR	C21-C22-C23-C24
24	F	303	BCR	C21-C22-C23-C24
24	G	203	BCR	C21-C22-C23-C24
24	I	4001	BCR	C7-C8-C9-C10
24	I	4001	BCR	C11-C12-C13-C14
24	J	104	BCR	C7-C8-C9-C10

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Mol	Chain	Res	Type	Atoms
24	K	206	BCR	C21-C22-C23-C24
24	L	204	BCR	C11-C12-C13-C14
24	L	204	BCR	C17-C18-C19-C20
24	3	304	BCR	C21-C22-C23-C24
24	3	305	BCR	C7-C8-C9-C10
24	3	305	BCR	C17-C18-C19-C20
24	3	307	BCR	C7-C8-C9-C10
24	3	307	BCR	C21-C22-C23-C24
24	7	303	BCR	C17-C18-C19-C20
24	4	804	BCR	C11-C12-C13-C14
24	4	804	BCR	C17-C18-C19-C20
24	5	304	BCR	C17-C18-C19-C20
24	6	305	BCR	C7-C8-C9-C10
24	6	305	BCR	C17-C18-C19-C20
24	6	306	BCR	C7-C8-C9-C10
24	6	306	BCR	C11-C12-C13-C14
24	6	306	BCR	C17-C18-C19-C20
36	J	105	C7Z	C11-C12-C13-C14
36	5	306	C7Z	C7-C8-C9-C10
36	5	306	C7Z	C31-C32-C33-C34
36	5	306	C7Z	C27-C28-C29-C30
38	1	302	LUT	C7-C8-C9-C10
38	3	303	LUT	C7-C8-C9-C10
38	5	302	LUT	C7-C8-C9-C10
38	5	303	LUT	C7-C8-C9-C10
38	6	304	LUT	C7-C8-C9-C10
38	9	302	LUT	C11-C12-C13-C14
25	7	318	LHG	O9-C7-O7-C5
22	B	809	CLA	O1A-CGA-O2A-C1
22	7	309	CLA	O1A-CGA-O2A-C1
22	6	308	CLA	O1A-CGA-O2A-C1
39	8	315	CHL	O1A-CGA-O2A-C1
22	B	805	CLA	C5-C6-C7-C8
22	B	818	CLA	C10-C11-C12-C13
22	5	310	CLA	C5-C6-C7-C8
27	A	852	DGA	CA7-CA8-CA9-CAA
22	A	834	CLA	C3-C5-C6-C7
22	K	203	CLA	C3-C5-C6-C7
22	7	307	CLA	C3-C5-C6-C7
22	6	310	CLA	C3-C5-C6-C7
22	3	310	CLA	CBA-CGA-O2A-C1
22	9	303	CLA	CBA-CGA-O2A-C1

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Mol	Chain	Res	Type	Atoms
22	A	818	CLA	C5-C6-C7-C8
22	B	829	CLA	C5-C6-C7-C8
22	9	303	CLA	C10-C11-C12-C13
22	5	307	CLA	O1D-CGD-O2D-CED
39	5	316	CHL	O1D-CGD-O2D-CED
22	A	839	CLA	CBD-CGD-O2D-CED
22	A	819	CLA	C5-C6-C7-C8
22	B	825	CLA	C5-C6-C7-C8
22	B	828	CLA	C10-C11-C12-C13
22	F	304	CLA	C5-C6-C7-C8
22	1	316	CLA	C10-C11-C12-C13
22	3	313	CLA	C10-C11-C12-C13
22	7	306	CLA	C8-C10-C11-C12
22	7	306	CLA	C10-C11-C12-C13
22	6	312	CLA	C5-C6-C7-C8
22	9	303	CLA	C5-C6-C7-C8
22	9	305	CLA	C5-C6-C7-C8
41	8	320	3PH	C3A-C3B-C3C-C3D
22	B	813	CLA	O1D-CGD-O2D-CED
22	K	202	CLA	O1D-CGD-O2D-CED
25	7	318	LHG	O1-C1-C2-O2
22	A	810	CLA	O1A-CGA-O2A-C1
25	A	849	LHG	C23-C24-C25-C26
25	A	850	LHG	C7-C8-C9-C10
25	A	850	LHG	C23-C24-C25-C26
25	B	803	LHG	C7-C8-C9-C10
25	Z	317	LHG	C7-C8-C9-C10
25	8	317	LHG	C23-C24-C25-C26
25	4	820	LHG	C23-C24-C25-C26
25	6	323	LHG	C23-C24-C25-C26
34	F	308	LMG	C10-C11-C12-C13
25	5	324	LHG	C11-C10-C9-C8
22	A	819	CLA	C8-C10-C11-C12
22	5	309	CLA	CBA-CGA-O2A-C1
25	7	318	LHG	C11-C10-C9-C8
22	A	803	CLA	O1D-CGD-O2D-CED
22	1	308	CLA	O1D-CGD-O2D-CED
22	A	832	CLA	C2-C1-O2A-CGA
22	A	838	CLA	C2-C1-O2A-CGA
22	A	839	CLA	C2-C1-O2A-CGA
22	B	806	CLA	C2-C1-O2A-CGA
22	B	825	CLA	C2-C1-O2A-CGA

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Mol	Chain	Res	Type	Atoms
22	B	831	CLA	C2-C1-O2A-CGA
22	1	314	CLA	C2-C1-O2A-CGA
22	3	310	CLA	C2-C1-O2A-CGA
22	3	316	CLA	C2-C1-O2A-CGA
22	8	310	CLA	C2-C1-O2A-CGA
22	8	311	CLA	C2-C1-O2A-CGA
22	4	810	CLA	C2-C1-O2A-CGA
22	6	302	CLA	C2-C1-O2A-CGA
22	6	307	CLA	C2-C1-O2A-CGA
22	3	312	CLA	C8-C10-C11-C12
25	B	803	LHG	C23-C24-C25-C26
22	4	806	CLA	C3-C5-C6-C7
22	A	806	CLA	C11-C10-C8-C7
22	A	810	CLA	C6-C7-C8-C10
22	A	820	CLA	C11-C10-C8-C7
22	A	824	CLA	C6-C7-C8-C10
22	A	827	CLA	C11-C12-C13-C15
22	A	841	CLA	C6-C7-C8-C10
22	B	810	CLA	C6-C7-C8-C10
22	B	815	CLA	C11-C10-C8-C7
22	B	819	CLA	C6-C7-C8-C10
22	B	819	CLA	C11-C10-C8-C7
22	B	828	CLA	C11-C10-C8-C7
22	B	834	CLA	C6-C7-C8-C10
22	B	838	CLA	C6-C7-C8-C10
22	L	201	CLA	C6-C7-C8-C10
22	Z	309	CLA	C6-C7-C8-C10
22	3	314	CLA	C6-C7-C8-C10
22	7	312	CLA	C6-C7-C8-C10
22	7	317	CLA	C6-C7-C8-C10
22	8	310	CLA	C11-C10-C8-C7
22	6	302	CLA	C6-C7-C8-C10
22	6	319	CLA	C6-C7-C8-C10
22	6	322	CLA	C6-C7-C8-C10
22	9	306	CLA	C11-C10-C8-C7
39	8	315	CHL	C11-C10-C8-C7
39	9	312	CHL	C12-C13-C15-C16
22	A	815	CLA	C3-C5-C6-C7
22	A	854	CLA	C3-C5-C6-C7
22	B	808	CLA	C3-C5-C6-C7
22	B	814	CLA	C3-C5-C6-C7
22	1	309	CLA	C3-C5-C6-C7

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Mol	Chain	Res	Type	Atoms
22	7	309	CLA	C3-C5-C6-C7
22	5	319	CLA	O1A-CGA-O2A-C1
25	5	324	LHG	O10-C23-O8-C6
24	A	844	BCR	C15-C16-C17-C18
24	A	848	BCR	C13-C14-C15-C16
24	B	845	BCR	C13-C14-C15-C16
24	B	847	BCR	C9-C10-C11-C12
24	B	849	BCR	C13-C14-C15-C16
24	B	853	BCR	C13-C14-C15-C16
24	I	4001	BCR	C9-C10-C11-C12
24	J	104	BCR	C13-C14-C15-C16
24	J	104	BCR	C15-C16-C17-C18
24	3	306	BCR	C13-C14-C15-C16
24	5	304	BCR	C15-C16-C17-C18
24	6	305	BCR	C15-C16-C17-C18
38	6	303	LUT	C9-C10-C11-C12
38	6	304	LUT	C9-C10-C11-C12
22	A	840	CLA	C2A-CAA-CBA-CGA
22	A	855	CLA	C2A-CAA-CBA-CGA
22	9	306	CLA	C2A-CAA-CBA-CGA
22	A	815	CLA	O1D-CGD-O2D-CED
22	B	826	CLA	O1D-CGD-O2D-CED
22	7	310	CLA	O1D-CGD-O2D-CED
22	8	311	CLA	O1D-CGD-O2D-CED
22	1	306	CLA	C8-C10-C11-C12
22	9	305	CLA	C8-C10-C11-C12
33	4	822	LMT	O1'-C1-C2-C3
22	A	824	CLA	O1A-CGA-O2A-C1
22	L	202	CLA	C8-C10-C11-C12
25	1	317	LHG	C23-C24-C25-C26
25	B	851	LHG	O2-C2-C3-O3
25	Z	317	LHG	O2-C2-C3-O3
25	7	318	LHG	O2-C2-C3-O3
25	8	317	LHG	O2-C2-C3-O3
25	5	324	LHG	O2-C2-C3-O3
25	6	323	LHG	O2-C2-C3-O3
22	A	808	CLA	C3-C5-C6-C7
22	1	307	CLA	C3-C5-C6-C7
22	A	825	CLA	C5-C6-C7-C8
22	Z	309	CLA	C5-C6-C7-C8
22	A	830	CLA	CBA-CGA-O2A-C1
22	6	310	CLA	CBA-CGA-O2A-C1

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Mol	Chain	Res	Type	Atoms
22	7	317	CLA	O1A-CGA-O2A-C1
22	8	308	CLA	O1A-CGA-O2A-C1
22	5	309	CLA	O1A-CGA-O2A-C1
39	6	315	CHL	O1A-CGA-O2A-C1
25	5	324	LHG	C7-C8-C9-C10
25	9	315	LHG	C23-C24-C25-C26
25	4	820	LHG	C9-C10-C11-C12
42	7	320	SPH	C12-C13-C14-C15
22	A	827	CLA	C8-C10-C11-C12
22	A	840	CLA	C8-C10-C11-C12
22	A	855	CLA	C8-C10-C11-C12
22	B	801	CLA	C13-C15-C16-C17
22	B	806	CLA	C8-C10-C11-C12
22	Z	303	CLA	C5-C6-C7-C8
22	3	313	CLA	C8-C10-C11-C12
22	1	304	CLA	O1D-CGD-O2D-CED
22	5	314	CLA	O1D-CGD-O2D-CED
22	3	310	CLA	O1A-CGA-O2A-C1
22	9	303	CLA	O1A-CGA-O2A-C1
25	4	821	LHG	C8-C7-O7-C5
22	A	806	CLA	C13-C15-C16-C17
22	A	820	CLA	C5-C6-C7-C8
22	A	825	CLA	C15-C16-C17-C18
22	A	828	CLA	C8-C10-C11-C12
22	B	817	CLA	C5-C6-C7-C8
22	B	830	CLA	C8-C10-C11-C12
22	B	841	CLA	C5-C6-C7-C8
22	7	323	CLA	C13-C15-C16-C17
22	8	307	CLA	C8-C10-C11-C12
22	5	315	CLA	C8-C10-C11-C12
22	6	307	CLA	C5-C6-C7-C8
22	6	322	CLA	C8-C10-C11-C12
25	A	849	LHG	C3-O3-P-O6
25	A	849	LHG	C4-O6-P-O3
25	A	850	LHG	C4-O6-P-O3
25	B	803	LHG	C3-O3-P-O6
25	7	318	LHG	C4-O6-P-O3
25	4	820	LHG	C4-O6-P-O3
25	5	324	LHG	C3-O3-P-O6
25	6	323	LHG	C4-O6-P-O3
25	9	315	LHG	C3-O3-P-O6
25	9	315	LHG	C4-O6-P-O3

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Mol	Chain	Res	Type	Atoms
25	5	324	LHG	C23-C24-C25-C26
22	8	309	CLA	C3-C5-C6-C7
22	5	309	CLA	C3-C5-C6-C7
27	A	852	DGA	CCA-CDA-CEA-CFA
22	B	824	CLA	CBA-CGA-O2A-C1
22	B	841	CLA	CBA-CGA-O2A-C1
31	3	301	DGD	C2A-C1A-O1G-C1G
22	2	304	CLA	O1D-CGD-O2D-CED
22	L	201	CLA	O1D-CGD-O2D-CED
22	Z	310	CLA	O1D-CGD-O2D-CED
25	7	318	LHG	C7-C8-C9-C10
22	B	825	CLA	O1D-CGD-O2D-CED
39	6	316	CHL	O1D-CGD-O2D-CED
25	A	849	LHG	C1-C2-C3-O3
25	B	850	LHG	C1-C2-C3-O3
25	B	851	LHG	C1-C2-C3-O3
25	Z	317	LHG	C1-C2-C3-O3
25	8	317	LHG	C1-C2-C3-O3
25	4	821	LHG	O9-C7-O7-C5
22	B	813	CLA	C2A-CAA-CBA-CGA
22	Z	306	CLA	C2A-CAA-CBA-CGA
22	3	313	CLA	C2A-CAA-CBA-CGA
22	5	320	CLA	C2A-CAA-CBA-CGA
39	1	312	CHL	C2A-CAA-CBA-CGA
39	Z	312	CHL	C2A-CAA-CBA-CGA
39	4	813	CHL	C2A-CAA-CBA-CGA
39	4	814	CHL	C2A-CAA-CBA-CGA
39	5	317	CHL	C2A-CAA-CBA-CGA
22	B	823	CLA	C6-C7-C8-C9
22	5	311	CLA	C6-C7-C8-C9
22	6	313	CLA	C6-C7-C8-C10
22	2	301	CLA	C6-C7-C8-C9
22	K	205	CLA	C3-C5-C6-C7
22	5	323	CLA	CBA-CGA-O2A-C1
22	A	805	CLA	CBA-CGA-O2A-C1
22	G	201	CLA	CBA-CGA-O2A-C1
25	4	820	LHG	C24-C23-O8-C6
25	4	821	LHG	C24-C23-O8-C6
22	G	202	CLA	O1A-CGA-O2A-C1
23	A	843	PQN	C25-C26-C27-C28
41	7	319	3PH	C37-C38-C39-C3A
24	A	844	BCR	C19-C20-C21-C22

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Mol	Chain	Res	Type	Atoms
24	A	847	BCR	C9-C10-C11-C12
24	A	848	BCR	C19-C20-C21-C22
24	A	856	BCR	C15-C16-C17-C18
24	B	848	BCR	C9-C10-C11-C12
24	B	849	BCR	C15-C16-C17-C18
24	F	303	BCR	C15-C16-C17-C18
24	5	304	BCR	C19-C20-C21-C22
24	6	305	BCR	C19-C20-C21-C22
38	4	802	LUT	C9-C10-C11-C12
27	A	852	DGA	CB1-CB2-CB3-CB4
33	4	822	LMT	C6-C7-C8-C9
22	1	316	CLA	C8-C10-C11-C12
22	7	323	CLA	C3-C5-C6-C7
25	A	849	LHG	C11-C12-C13-C14
25	4	820	LHG	C11-C12-C13-C14
27	A	852	DGA	CB9-CAB-CBB-CCB
40	1	318	SQD	C11-C10-C9-C8
22	3	313	CLA	O1D-CGD-O2D-CED
22	3	319	CLA	C6-C7-C8-C9
25	A	850	LHG	C13-C14-C15-C16
25	1	317	LHG	C11-C12-C13-C14
25	8	317	LHG	C11-C12-C13-C14
25	4	821	LHG	C11-C12-C13-C14
22	4	807	CLA	O1D-CGD-O2D-CED
22	A	807	CLA	C8-C10-C11-C12
22	3	322	CLA	C13-C15-C16-C17
25	A	849	LHG	C13-C14-C15-C16
25	1	317	LHG	C27-C28-C29-C30
41	8	320	3PH	C36-C37-C38-C39
25	3	321	LHG	C2-C3-O3-P
22	A	830	CLA	O1A-CGA-O2A-C1
22	6	310	CLA	O1A-CGA-O2A-C1
25	1	317	LHG	C12-C13-C14-C15
22	B	822	CLA	O1D-CGD-O2D-CED
25	1	317	LHG	O2-C2-C3-O3
25	1	317	LHG	C10-C11-C12-C13
25	1	317	LHG	C16-C17-C18-C19
22	7	317	CLA	C3-C5-C6-C7
22	8	305	CLA	C3-C5-C6-C7
22	A	811	CLA	CBA-CGA-O2A-C1
25	B	803	LHG	C24-C23-O8-C6
25	9	315	LHG	C24-C23-O8-C6

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Mol	Chain	Res	Type	Atoms
25	Z	317	LHG	C13-C14-C15-C16
25	8	317	LHG	C11-C10-C9-C8
27	A	852	DGA	CA6-CA7-CA8-CA9
22	A	832	CLA	C8-C10-C11-C12
22	B	830	CLA	C5-C6-C7-C8
22	B	812	CLA	C6-C7-C8-C9
22	K	205	CLA	C6-C7-C8-C10
22	7	322	CLA	C6-C7-C8-C9
22	4	812	CLA	C6-C7-C8-C10
22	6	311	CLA	C6-C7-C8-C10
22	B	838	CLA	O1D-CGD-O2D-CED
22	6	310	CLA	O1D-CGD-O2D-CED
25	A	850	LHG	C11-C12-C13-C14
22	A	827	CLA	C11-C12-C13-C14
22	A	828	CLA	C11-C10-C8-C9
22	A	840	CLA	C6-C7-C8-C9
22	B	814	CLA	C6-C7-C8-C9
22	B	817	CLA	C6-C7-C8-C9
22	B	818	CLA	C6-C7-C8-C9
22	1	314	CLA	C14-C13-C15-C16
22	1	316	CLA	C14-C13-C15-C16
22	3	311	CLA	C11-C10-C8-C9
22	4	805	CLA	C6-C7-C8-C9
22	8	308	CLA	C13-C15-C16-C17
25	9	315	LHG	C7-C8-C9-C10
25	4	820	LHG	C28-C29-C30-C31
25	9	315	LHG	C11-C10-C9-C8
22	A	809	CLA	C2A-CAA-CBA-CGA
22	3	311	CLA	C2A-CAA-CBA-CGA
22	7	323	CLA	C2A-CAA-CBA-CGA
22	8	311	CLA	C2A-CAA-CBA-CGA
22	5	308	CLA	C2A-CAA-CBA-CGA
22	6	319	CLA	C2A-CAA-CBA-CGA
39	6	316	CHL	C2A-CAA-CBA-CGA
24	A	845	BCR	C36-C18-C19-C20
24	A	846	BCR	C36-C18-C19-C20
24	A	846	BCR	C37-C22-C23-C24
24	B	845	BCR	C37-C22-C23-C24
24	B	848	BCR	C36-C18-C19-C20
24	B	849	BCR	C7-C8-C9-C34
24	F	303	BCR	C7-C8-C9-C34
24	G	203	BCR	C36-C18-C19-C20

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Mol	Chain	Res	Type	Atoms
24	J	104	BCR	C37-C22-C23-C24
24	K	206	BCR	C36-C18-C19-C20
24	L	204	BCR	C11-C12-C13-C35
24	3	305	BCR	C36-C18-C19-C20
24	8	304	BCR	C7-C8-C9-C34
24	8	304	BCR	C36-C18-C19-C20
24	8	304	BCR	C37-C22-C23-C24
24	4	804	BCR	C11-C12-C13-C35
24	5	305	BCR	C11-C12-C13-C35
24	6	306	BCR	C37-C22-C23-C24
32	F	306	RRX	C37-C22-C23-C24
32	F	306	RRX	C36-C18-C19-C20
38	1	302	LUT	C7-C8-C9-C19
38	1	303	LUT	C7-C8-C9-C19
38	Z	302	LUT	C7-C8-C9-C19
38	7	301	LUT	C7-C8-C9-C19
38	7	302	LUT	C7-C8-C9-C19
38	8	302	LUT	C31-C32-C33-C40
38	4	803	LUT	C7-C8-C9-C19
25	A	849	LHG	O1-C1-C2-C3
25	B	803	LHG	O1-C1-C2-C3
25	B	850	LHG	O1-C1-C2-C3
25	B	851	LHG	O1-C1-C2-C3
25	Z	317	LHG	O1-C1-C2-C3
25	7	318	LHG	O1-C1-C2-C3
25	4	820	LHG	O1-C1-C2-C3
25	4	821	LHG	O1-C1-C2-C3
25	5	324	LHG	O1-C1-C2-C3
25	6	323	LHG	O1-C1-C2-C3
24	A	845	BCR	C17-C18-C19-C20
24	A	846	BCR	C17-C18-C19-C20
24	A	846	BCR	C21-C22-C23-C24
24	A	847	BCR	C7-C8-C9-C10
24	A	848	BCR	C11-C12-C13-C14
24	B	845	BCR	C17-C18-C19-C20
24	B	846	BCR	C7-C8-C9-C10
24	B	848	BCR	C17-C18-C19-C20
24	B	849	BCR	C7-C8-C9-C10
24	F	303	BCR	C7-C8-C9-C10
24	G	203	BCR	C17-C18-C19-C20
24	J	104	BCR	C21-C22-C23-C24
24	K	206	BCR	C11-C12-C13-C14

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Mol	Chain	Res	Type	Atoms
24	K	206	BCR	C17-C18-C19-C20
24	8	304	BCR	C7-C8-C9-C10
24	8	304	BCR	C17-C18-C19-C20
24	8	304	BCR	C21-C22-C23-C24
24	5	305	BCR	C11-C12-C13-C14
24	5	305	BCR	C21-C22-C23-C24
24	6	306	BCR	C21-C22-C23-C24
32	F	306	RRX	C21-C22-C23-C24
32	F	306	RRX	C17-C18-C19-C20
36	J	105	C7Z	C27-C28-C29-C30
38	1	303	LUT	C7-C8-C9-C10
38	7	301	LUT	C7-C8-C9-C10
38	4	803	LUT	C7-C8-C9-C10
22	A	827	CLA	C3-C5-C6-C7
22	A	833	CLA	C3-C5-C6-C7
22	B	833	CLA	C3-C5-C6-C7
22	B	834	CLA	C3-C5-C6-C7
22	3	318	CLA	C5-C6-C7-C8
25	4	821	LHG	C9-C10-C11-C12
22	3	322	CLA	O1D-CGD-O2D-CED
25	5	324	LHG	C28-C29-C30-C31
22	B	824	CLA	O1A-CGA-O2A-C1
22	A	836	CLA	C6-C7-C8-C9
22	A	836	CLA	C6-C7-C8-C10
22	6	311	CLA	C6-C7-C8-C9
22	9	307	CLA	C6-C7-C8-C9
22	9	307	CLA	C6-C7-C8-C10
22	B	809	CLA	C5-C6-C7-C8
39	8	315	CHL	C10-C11-C12-C13
25	Z	317	LHG	C26-C27-C28-C29
25	4	820	LHG	C29-C30-C31-C32
34	F	308	LMG	C29-C30-C31-C32
22	6	319	CLA	CBD-CGD-O2D-CED
25	B	803	LHG	C11-C10-C9-C8
25	8	317	LHG	C28-C29-C30-C31
25	4	820	LHG	C33-C34-C35-C36
27	A	852	DGA	CB7-CB8-CB9-CAB
25	8	317	LHG	C7-C8-C9-C10
22	A	805	CLA	C5-C6-C7-C8
22	1	304	CLA	C5-C6-C7-C8
23	B	843	PQN	C23-C25-C26-C27
31	3	301	DGD	O1A-C1A-O1G-C1G

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Mol	Chain	Res	Type	Atoms
25	9	315	LHG	C26-C27-C28-C29
27	A	852	DGA	CA4-CA5-CA6-CA7
22	A	818	CLA	CBA-CGA-O2A-C1
22	L	201	CLA	CBA-CGA-O2A-C1
22	L	202	CLA	CBA-CGA-O2A-C1
25	A	850	LHG	C24-C23-O8-C6
25	4	820	LHG	C34-C35-C36-C37
41	7	319	3PH	C22-C23-C24-C25
22	A	810	CLA	O1D-CGD-O2D-CED
22	A	802	CLA	C3A-C2A-CAA-CBA
22	A	803	CLA	C3A-C2A-CAA-CBA
22	A	811	CLA	C3A-C2A-CAA-CBA
22	A	813	CLA	C3A-C2A-CAA-CBA
22	A	816	CLA	C3A-C2A-CAA-CBA
22	A	825	CLA	C3A-C2A-CAA-CBA
22	A	837	CLA	C3A-C2A-CAA-CBA
22	B	808	CLA	C3A-C2A-CAA-CBA
22	B	815	CLA	C3A-C2A-CAA-CBA
22	B	828	CLA	C3A-C2A-CAA-CBA
22	B	836	CLA	C3A-C2A-CAA-CBA
22	K	202	CLA	C3A-C2A-CAA-CBA
22	K	205	CLA	C3A-C2A-CAA-CBA
22	L	201	CLA	C3A-C2A-CAA-CBA
22	1	310	CLA	C3A-C2A-CAA-CBA
22	1	315	CLA	C3A-C2A-CAA-CBA
22	Z	315	CLA	C3A-C2A-CAA-CBA
22	3	315	CLA	C3A-C2A-CAA-CBA
22	7	306	CLA	C3A-C2A-CAA-CBA
22	8	306	CLA	C3A-C2A-CAA-CBA
22	8	307	CLA	C3A-C2A-CAA-CBA
22	8	314	CLA	C3A-C2A-CAA-CBA
22	4	807	CLA	C3A-C2A-CAA-CBA
22	5	308	CLA	C3A-C2A-CAA-CBA
22	5	314	CLA	C3A-C2A-CAA-CBA
22	5	326	CLA	C3A-C2A-CAA-CBA
22	6	307	CLA	C3A-C2A-CAA-CBA
22	9	303	CLA	C3A-C2A-CAA-CBA
39	4	816	CHL	C3A-C2A-CAA-CBA
22	A	838	CLA	C10-C11-C12-C13
22	1	307	CLA	C5-C6-C7-C8
22	4	808	CLA	C5-C6-C7-C8
22	6	311	CLA	C5-C6-C7-C8

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Mol	Chain	Res	Type	Atoms
24	G	203	BCR	C9-C10-C11-C12
25	7	318	LHG	C11-C12-C13-C14
25	7	318	LHG	C28-C29-C30-C31
25	6	323	LHG	C9-C10-C11-C12
22	A	822	CLA	C6-C7-C8-C9
22	8	314	CLA	O1D-CGD-O2D-CED
25	4	821	LHG	C4-C5-C6-O8
22	6	313	CLA	O2A-C1-C2-C3
22	A	811	CLA	C3-C5-C6-C7
22	A	828	CLA	C4-C3-C5-C6
22	F	304	CLA	C4-C3-C5-C6
39	8	315	CHL	C4-C3-C5-C6
22	5	318	CLA	CBA-CGA-O2A-C1
22	A	828	CLA	C2-C3-C5-C6
22	F	304	CLA	C2-C3-C5-C6
25	B	803	LHG	O1-C1-C2-O2
25	B	850	LHG	O1-C1-C2-O2
25	8	317	LHG	O1-C1-C2-O2
25	4	821	LHG	O1-C1-C2-O2
25	6	323	LHG	C25-C26-C27-C28
25	6	323	LHG	C28-C29-C30-C31
28	A	853	OCA	C4-C5-C6-C7
25	B	850	LHG	C7-C8-C9-C10
26	A	851	NKP	CAP-CAQ-CAR-CAS
22	A	805	CLA	O1A-CGA-O2A-C1
22	Z	307	CLA	C6-C7-C8-C9
22	3	319	CLA	C6-C7-C8-C10
22	5	326	CLA	C6-C7-C8-C9
25	4	821	LHG	C11-C10-C9-C8
22	A	802	CLA	C5-C6-C7-C8
22	B	826	CLA	C5-C6-C7-C8
41	7	319	3PH	C27-C28-C29-C2A
22	A	813	CLA	C3-C5-C6-C7
22	A	840	CLA	C3-C5-C6-C7
22	5	307	CLA	C3-C5-C6-C7
25	5	324	LHG	C11-C12-C13-C14
27	A	852	DGA	CB5-CB6-CB7-CB8
33	4	822	LMT	C7-C8-C9-C10
22	B	841	CLA	O1A-CGA-O2A-C1
22	G	201	CLA	O1A-CGA-O2A-C1
25	4	820	LHG	O10-C23-O8-C6
25	4	821	LHG	O10-C23-O8-C6

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Mol	Chain	Res	Type	Atoms
22	B	801	CLA	C8-C10-C11-C12
22	B	814	CLA	C8-C10-C11-C12
22	B	827	CLA	C5-C6-C7-C8
22	6	302	CLA	C5-C6-C7-C8
22	A	805	CLA	C2-C1-O2A-CGA
22	A	820	CLA	C2-C1-O2A-CGA
22	A	822	CLA	C2-C1-O2A-CGA
22	B	808	CLA	C2-C1-O2A-CGA
22	B	820	CLA	C2-C1-O2A-CGA
22	B	823	CLA	C2-C1-O2A-CGA
22	B	829	CLA	C2-C1-O2A-CGA
22	F	301	CLA	C2-C1-O2A-CGA
22	F	304	CLA	C2-C1-O2A-CGA
22	G	201	CLA	C2-C1-O2A-CGA
22	K	205	CLA	C2-C1-O2A-CGA
22	1	313	CLA	C2-C1-O2A-CGA
22	Z	313	CLA	C2-C1-O2A-CGA
22	Z	314	CLA	C2-C1-O2A-CGA
22	3	314	CLA	C2-C1-O2A-CGA
22	7	307	CLA	C2-C1-O2A-CGA
22	6	319	CLA	C2-C1-O2A-CGA
22	6	322	CLA	C2-C1-O2A-CGA
22	9	303	CLA	C2-C1-O2A-CGA
22	A	815	CLA	C5-C6-C7-C8
22	A	820	CLA	C8-C10-C11-C12
22	5	301	CLA	C5-C6-C7-C8
22	A	818	CLA	O1A-CGA-O2A-C1
25	B	803	LHG	O10-C23-O8-C6
25	9	315	LHG	O10-C23-O8-C6
25	1	317	LHG	C9-C10-C11-C12
25	Z	317	LHG	C28-C29-C30-C31
22	B	831	CLA	C3-C5-C6-C7
22	3	311	CLA	C3-C5-C6-C7
24	A	844	BCR	C1-C6-C7-C8
24	A	844	BCR	C23-C24-C25-C26
24	A	845	BCR	C1-C6-C7-C8
24	A	845	BCR	C23-C24-C25-C26
24	A	846	BCR	C5-C6-C7-C8
24	A	846	BCR	C23-C24-C25-C26
24	A	847	BCR	C5-C6-C7-C8
24	A	847	BCR	C23-C24-C25-C30
24	A	848	BCR	C23-C24-C25-C30

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Mol	Chain	Res	Type	Atoms
24	A	856	BCR	C5-C6-C7-C8
24	A	856	BCR	C23-C24-C25-C26
24	B	845	BCR	C1-C6-C7-C8
24	B	845	BCR	C5-C6-C7-C8
24	B	845	BCR	C23-C24-C25-C30
24	B	846	BCR	C1-C6-C7-C8
24	B	846	BCR	C23-C24-C25-C26
24	B	847	BCR	C23-C24-C25-C26
24	B	848	BCR	C23-C24-C25-C26
24	B	853	BCR	C23-C24-C25-C26
24	B	853	BCR	C23-C24-C25-C30
24	F	303	BCR	C5-C6-C7-C8
24	F	303	BCR	C23-C24-C25-C26
24	G	203	BCR	C1-C6-C7-C8
24	G	203	BCR	C23-C24-C25-C30
24	J	104	BCR	C1-C6-C7-C8
24	J	104	BCR	C23-C24-C25-C26
24	J	104	BCR	C23-C24-C25-C30
24	L	204	BCR	C5-C6-C7-C8
24	3	304	BCR	C5-C6-C7-C8
24	3	305	BCR	C23-C24-C25-C26
24	3	306	BCR	C1-C6-C7-C8
24	7	303	BCR	C1-C6-C7-C8
24	7	303	BCR	C23-C24-C25-C30
24	4	804	BCR	C23-C24-C25-C30
24	5	304	BCR	C23-C24-C25-C30
24	5	305	BCR	C1-C6-C7-C8
24	6	306	BCR	C23-C24-C25-C30
36	5	306	C7Z	C1-C6-C7-C8
36	5	306	C7Z	C21-C26-C27-C28
38	7	301	LUT	C5-C6-C7-C8
38	8	303	LUT	C5-C6-C7-C8
38	4	802	LUT	C1-C6-C7-C8
22	B	829	CLA	C8-C10-C11-C12
31	B	852	DGD	C3A-C4A-C5A-C6A
41	7	319	3PH	C24-C25-C26-C27
41	7	319	3PH	C25-C26-C27-C28
22	B	825	CLA	CBA-CGA-O2A-C1
22	B	835	CLA	CBA-CGA-O2A-C1
22	7	305	CLA	CBA-CGA-O2A-C1
39	3	317	CHL	CBA-CGA-O2A-C1
33	4	822	LMT	C1-C2-C3-C4

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Mol	Chain	Res	Type	Atoms
22	L	202	CLA	O1A-CGA-O2A-C1
41	6	324	3PH	C27-C28-C29-C2A
22	1	313	CLA	C8-C10-C11-C12
22	7	317	CLA	C5-C6-C7-C8
22	5	307	CLA	C8-C10-C11-C12
33	4	822	LMT	C3-C4-C5-C6
22	9	305	CLA	C4-C3-C5-C6
39	9	312	CHL	C4-C3-C5-C6
22	A	805	CLA	C11-C12-C13-C15
22	A	810	CLA	C11-C12-C13-C15
22	A	811	CLA	C11-C10-C8-C7
22	A	819	CLA	C6-C7-C8-C10
22	A	823	CLA	C11-C10-C8-C7
22	A	823	CLA	C11-C12-C13-C15
22	A	829	CLA	C11-C12-C13-C15
22	A	833	CLA	C6-C7-C8-C10
22	A	838	CLA	C6-C7-C8-C10
22	A	838	CLA	C11-C10-C8-C7
22	A	840	CLA	C6-C7-C8-C10
22	B	801	CLA	C12-C13-C15-C16
22	B	809	CLA	C11-C10-C8-C7
22	B	811	CLA	C6-C7-C8-C10
22	B	814	CLA	C6-C7-C8-C10
22	B	815	CLA	C2-C3-C5-C6
22	B	817	CLA	C6-C7-C8-C10
22	B	818	CLA	C6-C7-C8-C10
22	B	818	CLA	C11-C12-C13-C15
22	B	822	CLA	C6-C7-C8-C10
22	B	827	CLA	C6-C7-C8-C10
22	B	831	CLA	C6-C7-C8-C10
22	L	202	CLA	C11-C12-C13-C15
22	L	202	CLA	C12-C13-C15-C16
22	1	306	CLA	C11-C12-C13-C15
22	1	316	CLA	C12-C13-C15-C16
22	Z	305	CLA	C6-C7-C8-C10
22	Z	316	CLA	C11-C10-C8-C7
22	3	312	CLA	C11-C10-C8-C7
22	3	322	CLA	C11-C10-C8-C7
22	3	322	CLA	C11-C12-C13-C15
22	7	307	CLA	C11-C10-C8-C7
22	7	312	CLA	C11-C10-C8-C7
22	7	323	CLA	C6-C7-C8-C10

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Mol	Chain	Res	Type	Atoms
22	8	307	CLA	C11-C12-C13-C15
22	5	308	CLA	C6-C7-C8-C10
22	5	313	CLA	C11-C10-C8-C7
22	6	314	CLA	C6-C7-C8-C10
22	6	322	CLA	C11-C10-C8-C7
23	B	843	PQN	C17-C18-C20-C21
39	3	317	CHL	C2-C3-C5-C6
39	8	315	CHL	C2-C3-C5-C6
39	9	312	CHL	C11-C12-C13-C15
42	7	320	SPH	C4-C5-C6-C7
22	B	809	CLA	C3-C5-C6-C7
22	B	820	CLA	C3-C5-C6-C7
22	A	811	CLA	O1A-CGA-O2A-C1
22	B	825	CLA	O1A-CGA-O2A-C1
22	L	201	CLA	O1A-CGA-O2A-C1
22	5	318	CLA	O1A-CGA-O2A-C1
25	A	850	LHG	O10-C23-O8-C6
22	Z	313	CLA	C8-C10-C11-C12
24	B	847	BCR	C15-C16-C17-C18
24	G	203	BCR	C15-C16-C17-C18
24	3	306	BCR	C15-C16-C17-C18
24	5	305	BCR	C15-C16-C17-C18
32	F	306	RRX	C19-C20-C21-C22
22	B	842	CLA	C16-C17-C18-C20
22	K	205	CLA	C6-C7-C8-C9
22	7	322	CLA	C6-C7-C8-C10
22	5	319	CLA	C6-C7-C8-C9
42	7	320	SPH	C5-C6-C7-C8
31	1	319	DGD	C1B-C2B-C3B-C4B
22	8	306	CLA	CBA-CGA-O2A-C1
25	Z	317	LHG	C24-C23-O8-C6
35	J	102	T7X	C31-C11-O18-C9
22	A	826	CLA	C2A-CAA-CBA-CGA
22	B	815	CLA	C2A-CAA-CBA-CGA
22	B	831	CLA	C2A-CAA-CBA-CGA
22	Z	307	CLA	C2A-CAA-CBA-CGA
22	3	318	CLA	C2A-CAA-CBA-CGA
22	4	815	CLA	C2A-CAA-CBA-CGA
22	1	310	CLA	O1D-CGD-O2D-CED
22	7	323	CLA	O1D-CGD-O2D-CED
25	7	318	LHG	C9-C10-C11-C12
22	5	313	CLA	O1D-CGD-O2D-CED

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Mol	Chain	Res	Type	Atoms
25	A	850	LHG	C34-C35-C36-C37
25	Z	317	LHG	C9-C10-C11-C12
34	F	308	LMG	C12-C13-C14-C15
22	A	840	CLA	O1D-CGD-O2D-CED
22	1	307	CLA	C8-C10-C11-C12
22	Z	305	CLA	C3-C5-C6-C7
22	A	806	CLA	O1D-CGD-O2D-CED
22	B	826	CLA	CBA-CGA-O2A-C1
22	6	302	CLA	CBA-CGA-O2A-C1
22	Z	309	CLA	C11-C12-C13-C15
22	A	834	CLA	C5-C6-C7-C8
22	6	309	CLA	C5-C6-C7-C8
25	A	850	LHG	C28-C29-C30-C31
27	A	852	DGA	CB3-CB4-CB5-CB6
33	F	307	LMT	C2-C3-C4-C5
42	7	321	SPH	C10-C11-C12-C13
25	A	849	LHG	C7-C8-C9-C10
25	6	323	LHG	C8-C7-O7-C5
41	8	320	3PH	C22-C21-O21-C2
25	6	323	LHG	C29-C30-C31-C32
25	4	820	LHG	C30-C31-C32-C33
41	8	320	3PH	O22-C21-O21-C2
22	A	824	CLA	C3-C5-C6-C7
22	Z	309	CLA	C3-C5-C6-C7
22	A	806	CLA	C5-C6-C7-C8
25	A	850	LHG	O7-C5-C6-O8
25	4	821	LHG	O7-C5-C6-O8
25	5	324	LHG	O7-C5-C6-O8
25	6	323	LHG	C12-C13-C14-C15
25	9	315	LHG	C9-C10-C11-C12
22	B	823	CLA	C6-C7-C8-C10
22	5	311	CLA	C6-C7-C8-C10
35	J	102	T7X	C25-C26-C27-C28
22	B	815	CLA	C4-C3-C5-C6
39	3	317	CHL	C4-C3-C5-C6
39	6	318	CHL	C4-C3-C5-C6
22	A	805	CLA	C6-C7-C8-C9
22	A	810	CLA	C6-C7-C8-C9
22	A	819	CLA	C6-C7-C8-C9
22	A	824	CLA	C6-C7-C8-C9
22	A	833	CLA	C6-C7-C8-C9
22	A	855	CLA	C6-C7-C8-C9

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Mol	Chain	Res	Type	Atoms
22	B	810	CLA	C6-C7-C8-C9
22	B	811	CLA	C6-C7-C8-C9
22	B	822	CLA	C6-C7-C8-C9
22	B	824	CLA	C6-C7-C8-C9
22	B	827	CLA	C6-C7-C8-C9
22	B	831	CLA	C6-C7-C8-C9
22	B	834	CLA	C6-C7-C8-C9
22	B	838	CLA	C6-C7-C8-C9
22	F	302	CLA	C6-C7-C8-C9
22	L	202	CLA	C14-C13-C15-C16
22	1	314	CLA	C6-C7-C8-C9
22	Z	305	CLA	C6-C7-C8-C9
22	Z	306	CLA	C6-C7-C8-C9
22	7	323	CLA	C11-C12-C13-C14
22	8	309	CLA	C6-C7-C8-C9
22	6	314	CLA	C6-C7-C8-C9
22	6	314	CLA	C11-C12-C13-C14
22	6	322	CLA	C6-C7-C8-C9
22	9	306	CLA	C11-C10-C8-C9
23	B	843	PQN	C21-C22-C23-C24
39	9	312	CHL	C14-C13-C15-C16
41	6	324	3PH	C24-C25-C26-C27
22	A	819	CLA	C3-C5-C6-C7
22	B	817	CLA	C3-C5-C6-C7
22	F	304	CLA	C3-C5-C6-C7
22	Z	308	CLA	C3-C5-C6-C7
22	4	807	CLA	C3-C5-C6-C7
22	A	808	CLA	C2A-CAA-CBA-CGA
22	A	842	CLA	C2A-CAA-CBA-CGA
22	B	830	CLA	C2A-CAA-CBA-CGA
22	B	840	CLA	C2A-CAA-CBA-CGA
22	7	312	CLA	C2A-CAA-CBA-CGA
22	6	314	CLA	C2A-CAA-CBA-CGA
33	G	204	LMT	O5B-C5B-C6B-O6B
24	B	844	BCR	C37-C22-C23-C24
24	B	847	BCR	C7-C8-C9-C34
36	J	105	C7Z	C27-C28-C29-C39
38	Z	301	LUT	C7-C8-C9-C19
22	A	837	CLA	O1D-CGD-O2D-CED
22	A	808	CLA	C8-C10-C11-C12
22	F	301	CLA	C8-C10-C11-C12
22	5	307	CLA	C5-C6-C7-C8

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Mol	Chain	Res	Type	Atoms
25	7	318	LHG	C29-C30-C31-C32
24	B	853	BCR	C17-C18-C19-C20
38	Z	301	LUT	C7-C8-C9-C10
38	6	303	LUT	C31-C32-C33-C34
22	B	835	CLA	O1A-CGA-O2A-C1
22	7	305	CLA	O1A-CGA-O2A-C1
35	J	102	T7X	O19-C11-O18-C9
39	3	317	CHL	O1A-CGA-O2A-C1
22	B	839	CLA	O1D-CGD-O2D-CED
21	A	801	CL0	C1A-C2A-CAA-CBA
22	A	804	CLA	C1A-C2A-CAA-CBA
22	A	805	CLA	C1A-C2A-CAA-CBA
22	A	808	CLA	C1A-C2A-CAA-CBA
22	A	809	CLA	C1A-C2A-CAA-CBA
22	A	810	CLA	C1A-C2A-CAA-CBA
22	A	813	CLA	C1A-C2A-CAA-CBA
22	A	814	CLA	C1A-C2A-CAA-CBA
22	A	815	CLA	C1A-C2A-CAA-CBA
22	A	826	CLA	C1A-C2A-CAA-CBA
22	A	830	CLA	C1A-C2A-CAA-CBA
22	A	837	CLA	C1A-C2A-CAA-CBA
22	B	809	CLA	C1A-C2A-CAA-CBA
22	B	813	CLA	C1A-C2A-CAA-CBA
22	B	814	CLA	C1A-C2A-CAA-CBA
22	B	816	CLA	C1A-C2A-CAA-CBA
22	B	817	CLA	C1A-C2A-CAA-CBA
22	B	820	CLA	C1A-C2A-CAA-CBA
22	B	823	CLA	C1A-C2A-CAA-CBA
22	B	825	CLA	C1A-C2A-CAA-CBA
22	B	826	CLA	C1A-C2A-CAA-CBA
22	B	827	CLA	C1A-C2A-CAA-CBA
22	B	828	CLA	C1A-C2A-CAA-CBA
22	B	829	CLA	C1A-C2A-CAA-CBA
22	B	832	CLA	C1A-C2A-CAA-CBA
22	B	834	CLA	C1A-C2A-CAA-CBA
22	B	837	CLA	C1A-C2A-CAA-CBA
22	B	839	CLA	C1A-C2A-CAA-CBA
22	B	840	CLA	C1A-C2A-CAA-CBA
22	B	841	CLA	C1A-C2A-CAA-CBA
22	B	842	CLA	C1A-C2A-CAA-CBA
22	F	302	CLA	C1A-C2A-CAA-CBA
22	K	202	CLA	C1A-C2A-CAA-CBA

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Mol	Chain	Res	Type	Atoms
22	K	203	CLA	C1A-C2A-CAA-CBA
22	K	205	CLA	C1A-C2A-CAA-CBA
22	L	203	CLA	C1A-C2A-CAA-CBA
22	1	314	CLA	C1A-C2A-CAA-CBA
22	Z	303	CLA	C1A-C2A-CAA-CBA
22	Z	306	CLA	C1A-C2A-CAA-CBA
22	Z	313	CLA	C1A-C2A-CAA-CBA
22	Z	315	CLA	C1A-C2A-CAA-CBA
22	3	311	CLA	C1A-C2A-CAA-CBA
22	3	313	CLA	C1A-C2A-CAA-CBA
22	3	318	CLA	C1A-C2A-CAA-CBA
22	3	322	CLA	C1A-C2A-CAA-CBA
22	7	304	CLA	C1A-C2A-CAA-CBA
22	7	307	CLA	C1A-C2A-CAA-CBA
22	7	312	CLA	C1A-C2A-CAA-CBA
22	7	322	CLA	C1A-C2A-CAA-CBA
22	7	324	CLA	C1A-C2A-CAA-CBA
22	8	306	CLA	C1A-C2A-CAA-CBA
22	8	308	CLA	C1A-C2A-CAA-CBA
22	8	310	CLA	C1A-C2A-CAA-CBA
22	8	313	CLA	C1A-C2A-CAA-CBA
22	4	808	CLA	C1A-C2A-CAA-CBA
22	4	811	CLA	C1A-C2A-CAA-CBA
22	4	815	CLA	C1A-C2A-CAA-CBA
22	5	310	CLA	C1A-C2A-CAA-CBA
22	5	311	CLA	C1A-C2A-CAA-CBA
22	5	313	CLA	C1A-C2A-CAA-CBA
22	5	318	CLA	C1A-C2A-CAA-CBA
22	6	312	CLA	C1A-C2A-CAA-CBA
22	6	314	CLA	C1A-C2A-CAA-CBA
22	6	317	CLA	C1A-C2A-CAA-CBA
22	6	319	CLA	C1A-C2A-CAA-CBA
22	9	308	CLA	C1A-C2A-CAA-CBA
22	9	313	CLA	C1A-C2A-CAA-CBA
22	2	301	CLA	C1A-C2A-CAA-CBA
39	8	315	CHL	C1A-C2A-CAA-CBA
39	4	816	CHL	C1A-C2A-CAA-CBA
22	A	822	CLA	C6-C7-C8-C10
22	B	812	CLA	C6-C7-C8-C10
22	5	319	CLA	C6-C7-C8-C10
22	5	326	CLA	C6-C7-C8-C10
22	2	301	CLA	C6-C7-C8-C10

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Mol	Chain	Res	Type	Atoms
25	6	323	LHG	O9-C7-O7-C5
25	A	850	LHG	C33-C34-C35-C36
25	4	821	LHG	C10-C11-C12-C13
24	A	846	BCR	C19-C20-C21-C22
24	B	846	BCR	C13-C14-C15-C16
36	5	306	C7Z	C13-C14-C15-C35
38	7	301	LUT	C9-C10-C11-C12
22	3	310	CLA	C5-C6-C7-C8
22	8	310	CLA	C8-C10-C11-C12
25	Z	317	LHG	C4-O6-P-O3
22	3	318	CLA	C3-C5-C6-C7
25	B	851	LHG	C2-C3-O3-P
22	A	835	CLA	C5-C6-C7-C8
22	1	316	CLA	C15-C16-C17-C18
22	B	811	CLA	CBA-CGA-O2A-C1
25	A	850	LHG	O6-C4-C5-C6
25	8	317	LHG	O6-C4-C5-C6
25	5	324	LHG	O6-C4-C5-C6
31	1	319	DGD	O6E-C5E-C6E-O5E
22	A	805	CLA	C15-C16-C17-C18
22	1	308	CLA	C6-C7-C8-C9
22	Z	309	CLA	C11-C12-C13-C14
22	5	308	CLA	C3-C5-C6-C7
22	B	828	CLA	C13-C15-C16-C17
22	B	834	CLA	C8-C10-C11-C12
41	8	320	3PH	C35-C36-C37-C38
22	6	307	CLA	CBA-CGA-O2A-C1
25	1	317	LHG	C1-C2-C3-O3
33	F	307	LMT	O5'-C5'-C6'-O6'
22	A	814	CLA	C4-C3-C5-C6
39	6	315	CHL	C4-C3-C5-C6
22	7	316	CLA	C3A-C2A-CAA-CBA
22	A	812	CLA	C8-C10-C11-C12
22	A	829	CLA	C8-C10-C11-C12
22	B	842	CLA	C5-C6-C7-C8
22	Z	309	CLA	C8-C10-C11-C12
22	8	306	CLA	O1A-CGA-O2A-C1
22	1	316	CLA	C3-C5-C6-C7
22	5	311	CLA	C3-C5-C6-C7
25	3	321	LHG	C4-C5-C6-O8
25	7	318	LHG	C4-C5-C6-O8
25	5	324	LHG	C13-C14-C15-C16

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Mol	Chain	Res	Type	Atoms
25	6	323	LHG	C4-C5-C6-O8
31	B	852	DGD	O1G-C1G-C2G-C3G
41	7	319	3PH	C3C-C3D-C3E-C3F
41	7	319	3PH	C3A-C3B-C3C-C3D
22	B	811	CLA	O1A-CGA-O2A-C1
31	B	852	DGD	C5D-C6D-O5D-C1E
31	3	301	DGD	C2G-C3G-O3G-C1D
22	B	817	CLA	C8-C10-C11-C12
22	Z	306	CLA	C8-C10-C11-C12
39	3	317	CHL	C8-C10-C11-C12
34	F	308	LMG	C15-C16-C17-C18
41	6	324	3PH	C21-C22-C23-C24
25	Z	317	LHG	O10-C23-O8-C6
25	7	318	LHG	C26-C27-C28-C29
21	A	801	CL0	CBA-CGA-O2A-C1
25	8	317	LHG	C24-C23-O8-C6
33	1	301	LMT	O5'-C5'-C6'-O6'
33	4	801	LMT	O5'-C5'-C6'-O6'
25	A	850	LHG	O1-C1-C2-O2
25	4	821	LHG	C23-C24-C25-C26
33	F	307	LMT	O5B-C5B-C6B-O6B
22	B	822	CLA	C5-C6-C7-C8
22	7	310	CLA	C8-C10-C11-C12
22	8	309	CLA	C10-C11-C12-C13
22	A	822	CLA	C4-C3-C5-C6
22	B	842	CLA	C4-C3-C5-C6
25	B	803	LHG	C9-C10-C11-C12
41	7	319	3PH	C39-C3A-C3B-C3C
22	6	302	CLA	O1A-CGA-O2A-C1
22	B	842	CLA	C16-C17-C18-C19
22	A	820	CLA	CBA-CGA-O2A-C1
22	9	305	CLA	CBA-CGA-O2A-C1
41	5	325	3PH	C22-C23-C24-C25
22	B	805	CLA	C8-C10-C11-C12
22	B	823	CLA	C5-C6-C7-C8
22	7	310	CLA	C13-C15-C16-C17
42	7	321	SPH	C2-C3-C4-C5
25	4	820	LHG	C31-C32-C33-C34
22	A	821	CLA	C2A-CAA-CBA-CGA
22	B	814	CLA	C2A-CAA-CBA-CGA
22	B	824	CLA	C2A-CAA-CBA-CGA
22	3	314	CLA	C2A-CAA-CBA-CGA

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Mol	Chain	Res	Type	Atoms
22	A	838	CLA	C5-C6-C7-C8
22	Z	303	CLA	C8-C10-C11-C12
22	A	811	CLA	C2-C1-O2A-CGA
22	Z	316	CLA	C2-C1-O2A-CGA
22	7	315	CLA	C2-C1-O2A-CGA
22	4	808	CLA	C2-C1-O2A-CGA
22	5	301	CLA	C2-C1-O2A-CGA
22	9	307	CLA	C2-C1-O2A-CGA
22	9	313	CLA	C2-C1-O2A-CGA
33	4	822	LMT	O5B-C5B-C6B-O6B
22	7	307	CLA	O1D-CGD-O2D-CED
42	7	321	SPH	C4-C5-C6-C7
22	5	323	CLA	O1A-CGA-O2A-C1
22	A	854	CLA	C5-C6-C7-C8
22	Z	316	CLA	C5-C6-C7-C8
22	4	805	CLA	C5-C6-C7-C8
22	5	313	CLA	C10-C11-C12-C13
25	B	803	LHG	C10-C11-C12-C13
25	8	317	LHG	C13-C14-C15-C16
28	A	853	OCA	C5-C6-C7-C8
22	A	817	CLA	CBA-CGA-O2A-C1
22	1	313	CLA	CBA-CGA-O2A-C1
22	3	313	CLA	CBA-CGA-O2A-C1
35	J	102	T7X	O13-C7-C8-O16
27	A	852	DGA	CA1-CA2-CA3-CA4
25	B	803	LHG	C11-C12-C13-C14
22	B	813	CLA	C5-C6-C7-C8
22	B	818	CLA	C8-C10-C11-C12
22	K	205	CLA	C5-C6-C7-C8
22	7	317	CLA	C8-C10-C11-C12
22	A	839	CLA	O1D-CGD-O2D-CED
22	B	826	CLA	O1A-CGA-O2A-C1
34	J	101	LMG	C2-C1-O1-C7
25	4	820	LHG	O8-C23-C24-C25
25	7	318	LHG	O7-C5-C6-O8
25	8	317	LHG	O7-C5-C6-O8
25	1	317	LHG	C31-C32-C33-C34
22	A	823	CLA	C8-C10-C11-C12
21	A	801	CL0	O1A-CGA-O2A-C1
22	1	313	CLA	O1A-CGA-O2A-C1
21	A	801	CL0	C6-C7-C8-C10
22	A	805	CLA	C6-C7-C8-C10

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Mol	Chain	Res	Type	Atoms
22	A	807	CLA	C11-C10-C8-C7
22	A	813	CLA	C6-C7-C8-C10
22	A	815	CLA	C11-C12-C13-C15
22	A	817	CLA	C11-C10-C8-C7
22	A	818	CLA	C11-C10-C8-C7
22	A	821	CLA	C6-C7-C8-C10
22	A	828	CLA	C11-C12-C13-C15
22	A	830	CLA	C11-C12-C13-C15
22	A	832	CLA	C11-C10-C8-C7
22	A	855	CLA	C6-C7-C8-C10
22	A	855	CLA	C11-C12-C13-C15
22	B	801	CLA	C11-C10-C8-C7
22	B	805	CLA	C6-C7-C8-C10
22	B	808	CLA	C6-C7-C8-C10
22	B	811	CLA	C11-C10-C8-C7
22	B	820	CLA	C11-C10-C8-C7
22	B	821	CLA	C6-C7-C8-C10
22	B	821	CLA	C11-C12-C13-C15
22	B	824	CLA	C6-C7-C8-C10
22	B	824	CLA	C11-C10-C8-C7
22	B	825	CLA	C6-C7-C8-C10
22	B	838	CLA	C12-C13-C15-C16
22	B	842	CLA	C6-C7-C8-C10
22	F	301	CLA	C6-C7-C8-C10
22	F	302	CLA	C6-C7-C8-C10
22	F	304	CLA	C11-C12-C13-C15
22	L	201	CLA	C11-C10-C8-C7
22	1	306	CLA	C6-C7-C8-C10
22	1	310	CLA	C11-C10-C8-C7
22	1	314	CLA	C6-C7-C8-C10
22	1	314	CLA	C11-C10-C8-C7
22	1	314	CLA	C12-C13-C15-C16
22	1	316	CLA	C11-C10-C8-C7
22	Z	306	CLA	C6-C7-C8-C10
22	Z	308	CLA	C6-C7-C8-C10
22	Z	316	CLA	C6-C7-C8-C10
22	Z	316	CLA	C11-C12-C13-C15
22	3	308	CLA	C6-C7-C8-C10
22	3	313	CLA	C11-C12-C13-C15
22	3	314	CLA	C11-C10-C8-C7
22	7	304	CLA	C11-C10-C8-C7
22	7	307	CLA	C6-C7-C8-C10

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Mol	Chain	Res	Type	Atoms
22	7	307	CLA	C11-C12-C13-C15
22	7	324	CLA	C11-C10-C8-C7
22	8	305	CLA	C6-C7-C8-C10
22	8	307	CLA	C6-C7-C8-C10
22	8	309	CLA	C6-C7-C8-C10
22	8	309	CLA	C11-C12-C13-C15
22	4	805	CLA	C11-C10-C8-C7
22	4	807	CLA	C11-C10-C8-C7
22	5	301	CLA	C6-C7-C8-C10
22	5	309	CLA	C6-C7-C8-C10
22	5	310	CLA	C11-C10-C8-C7
22	5	315	CLA	C6-C7-C8-C10
22	5	318	CLA	C6-C7-C8-C10
22	5	322	CLA	C11-C10-C8-C7
22	6	314	CLA	C11-C12-C13-C15
22	9	306	CLA	C6-C7-C8-C10
23	A	843	PQN	C17-C18-C20-C21
23	B	843	PQN	C21-C22-C23-C25
39	3	317	CHL	C11-C10-C8-C7
22	A	806	CLA	C3-C5-C6-C7
21	A	801	CL0	C6-C7-C8-C9
22	A	805	CLA	C11-C10-C8-C9
22	A	811	CLA	C6-C7-C8-C9
22	A	813	CLA	C6-C7-C8-C9
22	A	820	CLA	C11-C10-C8-C9
22	A	827	CLA	C6-C7-C8-C9
22	A	828	CLA	C6-C7-C8-C9
22	A	832	CLA	C11-C10-C8-C9
22	A	838	CLA	C6-C7-C8-C9
22	A	841	CLA	C6-C7-C8-C9
22	A	855	CLA	C11-C10-C8-C9
22	B	801	CLA	C11-C10-C8-C9
22	B	805	CLA	C6-C7-C8-C9
22	B	808	CLA	C6-C7-C8-C9
22	B	821	CLA	C6-C7-C8-C9
22	B	828	CLA	C6-C7-C8-C9
22	B	828	CLA	C11-C12-C13-C14
22	B	838	CLA	C14-C13-C15-C16
22	B	841	CLA	C6-C7-C8-C9
22	B	842	CLA	C6-C7-C8-C9
22	F	301	CLA	C6-C7-C8-C9
22	F	302	CLA	C11-C12-C13-C14

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Mol	Chain	Res	Type	Atoms
22	L	201	CLA	C11-C10-C8-C9
22	1	314	CLA	C11-C10-C8-C9
22	Z	308	CLA	C6-C7-C8-C9
22	Z	316	CLA	C6-C7-C8-C9
22	3	308	CLA	C6-C7-C8-C9
22	3	314	CLA	C6-C7-C8-C9
22	7	306	CLA	C6-C7-C8-C9
22	7	307	CLA	C6-C7-C8-C9
22	7	308	CLA	C6-C7-C8-C9
22	7	312	CLA	C6-C7-C8-C9
22	7	323	CLA	C6-C7-C8-C9
22	8	305	CLA	C6-C7-C8-C9
22	8	307	CLA	C6-C7-C8-C9
22	8	309	CLA	C11-C12-C13-C14
22	8	310	CLA	C11-C10-C8-C9
22	4	805	CLA	C11-C10-C8-C9
22	4	807	CLA	C11-C10-C8-C9
22	5	301	CLA	C6-C7-C8-C9
22	5	307	CLA	C6-C7-C8-C9
22	5	308	CLA	C6-C7-C8-C9
22	5	309	CLA	C6-C7-C8-C9
22	6	301	CLA	C6-C7-C8-C9
22	6	302	CLA	C6-C7-C8-C9
22	6	310	CLA	C6-C7-C8-C9
22	6	319	CLA	C6-C7-C8-C9
22	9	306	CLA	C6-C7-C8-C9
23	A	843	PQN	C19-C18-C20-C21
39	8	315	CHL	C11-C10-C8-C9
39	9	312	CHL	C11-C12-C13-C14
22	1	314	CLA	C2A-CAA-CBA-CGA
22	3	308	CLA	C2A-CAA-CBA-CGA
22	7	324	CLA	C2A-CAA-CBA-CGA
39	8	312	CHL	C2A-CAA-CBA-CGA
27	A	852	DGA	CA8-CA9-CAA-CBA
25	8	317	LHG	O10-C23-O8-C6
24	G	203	BCR	C11-C12-C13-C35
24	7	303	BCR	C37-C22-C23-C24
24	4	804	BCR	C37-C22-C23-C24
24	5	304	BCR	C37-C22-C23-C24
24	5	305	BCR	C7-C8-C9-C34
38	Z	301	LUT	C27-C28-C29-C39
22	L	202	CLA	C5-C6-C7-C8

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Mol	Chain	Res	Type	Atoms
22	K	203	CLA	C6-C7-C8-C9
24	B	847	BCR	C7-C8-C9-C10
24	7	303	BCR	C21-C22-C23-C24
24	4	804	BCR	C21-C22-C23-C24
24	5	305	BCR	C7-C8-C9-C10
22	B	826	CLA	C8-C10-C11-C12
22	9	305	CLA	O1A-CGA-O2A-C1
22	A	813	CLA	CBA-CGA-O2A-C1
22	3	314	CLA	CBA-CGA-O2A-C1
22	7	310	CLA	C15-C16-C17-C18
25	6	323	LHG	C19-C20-C21-C22
22	A	817	CLA	O1A-CGA-O2A-C1
22	6	307	CLA	O1A-CGA-O2A-C1
22	L	202	CLA	CBD-CGD-O2D-CED
25	4	820	LHG	C15-C16-C17-C18
25	Z	317	LHG	O6-C4-C5-C6
25	7	318	LHG	O6-C4-C5-C6
41	6	324	3PH	O11-C1-C2-C3
22	7	308	CLA	C3-C5-C6-C7
22	B	842	CLA	CBA-CGA-O2A-C1
21	A	801	CL0	C5-C6-C7-C8
22	A	830	CLA	C5-C6-C7-C8
22	5	308	CLA	C8-C10-C11-C12
22	5	301	CLA	C11-C10-C8-C9
25	6	323	LHG	C35-C36-C37-C38
22	L	201	CLA	C3-C5-C6-C7
41	6	324	3PH	C25-C26-C27-C28
25	1	317	LHG	C24-C23-O8-C6
25	8	317	LHG	C9-C10-C11-C12
26	8	318	NKP	CAH-CAG-OAF-PAC
21	A	801	CL0	C3A-C2A-CAA-CBA
22	B	809	CLA	C3A-C2A-CAA-CBA
22	B	824	CLA	C3A-C2A-CAA-CBA
22	B	826	CLA	C3A-C2A-CAA-CBA
22	Z	310	CLA	C3A-C2A-CAA-CBA
39	Z	311	CHL	C3A-C2A-CAA-CBA
39	8	315	CHL	C3A-C2A-CAA-CBA
31	3	301	DGD	C3A-C4A-C5A-C6A
24	K	206	BCR	C13-C14-C15-C16
24	8	304	BCR	C9-C10-C11-C12
22	4	807	CLA	C5-C6-C7-C8
22	1	308	CLA	C6-C7-C8-C10

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Mol	Chain	Res	Type	Atoms
22	A	854	CLA	C8-C10-C11-C12
25	A	850	LHG	C4-C5-C6-O8
25	B	851	LHG	C4-C5-C6-O8
25	1	317	LHG	C4-C5-C6-O8
25	Z	317	LHG	C4-C5-C6-O8
25	5	324	LHG	C4-C5-C6-O8
31	3	301	DGD	O1G-C1G-C2G-C3G
34	F	308	LMG	C7-C8-C9-O8
40	1	318	SQD	C44-C45-C46-O48
22	A	820	CLA	O1A-CGA-O2A-C1
25	A	849	LHG	C16-C17-C18-C19
25	Z	317	LHG	C31-C32-C33-C34
22	A	829	CLA	O2A-C1-C2-C3
22	L	202	CLA	O2A-C1-C2-C3
23	A	843	PQN	C13-C15-C16-C17
25	4	821	LHG	C7-C8-C9-C10
22	3	308	CLA	C8-C10-C11-C12
22	8	308	CLA	C8-C10-C11-C12
26	A	851	NKP	OBC-CAH-CAI-OAJ
22	K	203	CLA	C6-C7-C8-C10
22	4	812	CLA	C6-C7-C8-C9
22	9	305	CLA	C2-C3-C5-C6
22	B	824	CLA	C11-C12-C13-C14
39	3	317	CHL	CBD-CGD-O2D-CED
35	J	102	T7X	C7-O13-P1-O1
35	J	102	T7X	C18-C19-C20-C21
35	J	102	T7X	C22-C23-C24-C25
22	6	309	CLA	CAA-CBA-CGA-O2A
22	A	816	CLA	C2A-CAA-CBA-CGA
22	B	835	CLA	C2A-CAA-CBA-CGA
39	7	313	CHL	C2A-CAA-CBA-CGA
22	A	830	CLA	C8-C10-C11-C12
22	2	301	CLA	C5-C6-C7-C8
39	8	315	CHL	C8-C10-C11-C12
25	A	850	LHG	O6-C4-C5-O7
25	B	850	LHG	O6-C4-C5-O7
25	Z	317	LHG	O6-C4-C5-O7
25	8	317	LHG	O6-C4-C5-O7
39	Z	312	CHL	CBA-CGA-O2A-C1
25	6	323	LHG	C11-C12-C13-C14
31	1	319	DGD	C1A-C2A-C3A-C4A
22	4	811	CLA	C6-C7-C8-C9

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Mol	Chain	Res	Type	Atoms
23	A	843	PQN	C26-C27-C28-C30
22	1	314	CLA	C8-C10-C11-C12
22	3	314	CLA	O1A-CGA-O2A-C1
22	A	822	CLA	C3-C5-C6-C7
25	A	849	LHG	O7-C5-C6-O8
25	6	323	LHG	O7-C5-C6-O8
31	1	319	DGD	O2G-C2G-C3G-O3G
40	1	318	SQD	O6-C44-C45-O47
40	1	318	SQD	O47-C45-C46-O48
24	3	306	BCR	C9-C10-C11-C12
22	A	809	CLA	C6-C7-C8-C9
22	1	310	CLA	C11-C12-C13-C15
22	Z	308	CLA	C8-C10-C11-C12
25	3	321	LHG	C1-C2-C3-O3
25	9	315	LHG	C1-C2-C3-O3
25	9	315	LHG	C24-C25-C26-C27
22	B	801	CLA	C2-C1-O2A-CGA
22	5	315	CLA	C2-C1-O2A-CGA
22	6	301	CLA	C2-C1-O2A-CGA
22	A	813	CLA	O1A-CGA-O2A-C1
22	B	819	CLA	C8-C10-C11-C12
22	B	821	CLA	C8-C10-C11-C12
22	B	805	CLA	C11-C12-C13-C14
22	B	816	CLA	C6-C7-C8-C9
22	B	840	CLA	C6-C7-C8-C9
22	L	201	CLA	C6-C7-C8-C9
22	L	202	CLA	C11-C10-C8-C9
22	1	316	CLA	C11-C12-C13-C14
22	Z	309	CLA	C6-C7-C8-C9
22	4	808	CLA	C6-C7-C8-C9
22	6	312	CLA	C11-C12-C13-C14
25	Z	317	LHG	C25-C26-C27-C28
25	7	318	LHG	C5-C4-O6-P
25	5	324	LHG	C5-C4-O6-P
35	J	102	T7X	C8-C7-O13-P1
22	3	313	CLA	O1A-CGA-O2A-C1
25	4	820	LHG	C13-C14-C15-C16
22	A	820	CLA	C11-C12-C13-C15
22	Z	307	CLA	C6-C7-C8-C10
22	6	313	CLA	C6-C7-C8-C9
22	K	204	CLA	O2A-C1-C2-C3
39	5	321	CHL	CBD-CGD-O2D-CED

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Mol	Chain	Res	Type	Atoms
24	G	203	BCR	C23-C24-C25-C26
36	5	306	C7Z	C25-C26-C27-C28
38	3	303	LUT	C1-C6-C7-C8
38	8	302	LUT	C5-C6-C7-C8
22	3	318	CLA	C8-C10-C11-C12
38	3	303	LUT	C31-C32-C33-C40
38	4	802	LUT	C11-C12-C13-C20
24	A	847	BCR	C21-C22-C23-C24
24	5	304	BCR	C21-C22-C23-C24
38	7	302	LUT	C11-C12-C13-C14
38	8	302	LUT	C31-C32-C33-C34
38	5	302	LUT	C27-C28-C29-C30
22	B	824	CLA	C5-C6-C7-C8
22	K	203	CLA	C5-C6-C7-C8
22	7	310	CLA	C5-C6-C7-C8
25	B	803	LHG	C25-C26-C27-C28
27	A	852	DGA	CA2-CA3-CA4-CA5
22	B	810	CLA	C11-C10-C8-C9
22	5	309	CLA	C5-C6-C7-C8
22	6	319	CLA	O1D-CGD-O2D-CED
25	B	803	LHG	O6-C4-C5-C6
25	B	850	LHG	O6-C4-C5-C6
25	6	323	LHG	O6-C4-C5-C6
25	6	323	LHG	C31-C32-C33-C34
22	A	802	CLA	C6-C7-C8-C10
22	A	803	CLA	C6-C7-C8-C10
22	A	806	CLA	C6-C7-C8-C10
22	A	808	CLA	C6-C7-C8-C10
22	A	808	CLA	C12-C13-C15-C16
22	A	811	CLA	C6-C7-C8-C10
22	A	812	CLA	C6-C7-C8-C10
22	A	812	CLA	C11-C10-C8-C7
22	A	815	CLA	C6-C7-C8-C10
22	A	816	CLA	C6-C7-C8-C10
22	A	820	CLA	C6-C7-C8-C10
22	A	823	CLA	C6-C7-C8-C10
22	A	827	CLA	C6-C7-C8-C10
22	A	828	CLA	C6-C7-C8-C10
22	A	830	CLA	C11-C10-C8-C7
22	A	833	CLA	C11-C10-C8-C7
22	A	834	CLA	C6-C7-C8-C10
22	A	854	CLA	C6-C7-C8-C10

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Mol	Chain	Res	Type	Atoms
22	A	855	CLA	C11-C10-C8-C7
22	B	806	CLA	C11-C12-C13-C15
22	B	814	CLA	C11-C10-C8-C7
22	B	815	CLA	C6-C7-C8-C10
22	B	816	CLA	C6-C7-C8-C10
22	B	826	CLA	C11-C12-C13-C15
22	B	827	CLA	C11-C12-C13-C15
22	B	828	CLA	C6-C7-C8-C10
22	B	833	CLA	C11-C10-C8-C7
22	B	840	CLA	C6-C7-C8-C10
22	B	841	CLA	C6-C7-C8-C10
22	B	841	CLA	C11-C10-C8-C7
22	B	842	CLA	C11-C10-C8-C7
22	L	202	CLA	C11-C10-C8-C7
22	1	306	CLA	C11-C10-C8-C7
22	1	309	CLA	C6-C7-C8-C10
22	1	313	CLA	C6-C7-C8-C10
22	1	316	CLA	C11-C12-C13-C15
22	3	311	CLA	C6-C7-C8-C10
22	7	304	CLA	C6-C7-C8-C10
22	7	306	CLA	C6-C7-C8-C10
22	7	308	CLA	C11-C12-C13-C15
22	7	310	CLA	C6-C7-C8-C10
22	7	317	CLA	C11-C10-C8-C7
22	7	324	CLA	C6-C7-C8-C10
22	8	308	CLA	C11-C10-C8-C7
22	4	807	CLA	C6-C7-C8-C10
22	5	307	CLA	C6-C7-C8-C10
22	5	322	CLA	C6-C7-C8-C10
22	6	301	CLA	C6-C7-C8-C10
22	6	310	CLA	C6-C7-C8-C10
22	6	312	CLA	C6-C7-C8-C10
39	5	316	CHL	C12-C13-C15-C16
35	J	102	T7X	C1-O1-P1-O13
35	J	102	T7X	C24-C25-C26-C27
25	6	323	LHG	C30-C31-C32-C33
22	A	812	CLA	C13-C15-C16-C17
22	1	316	CLA	C5-C6-C7-C8
24	3	304	BCR	C19-C20-C21-C22
24	6	306	BCR	C9-C10-C11-C12
38	Z	301	LUT	C9-C10-C11-C12
22	A	842	CLA	CAA-CBA-CGA-O2A

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Mol	Chain	Res	Type	Atoms
22	K	202	CLA	C2A-CAA-CBA-CGA
39	Z	311	CHL	C2A-CAA-CBA-CGA
39	8	301	CHL	C2A-CAA-CBA-CGA
39	5	316	CHL	C2A-CAA-CBA-CGA
39	6	318	CHL	C2A-CAA-CBA-CGA
39	2	303	CHL	C2A-CAA-CBA-CGA
25	1	317	LHG	C15-C16-C17-C18
22	5	319	CLA	C5-C6-C7-C8
22	6	314	CLA	C10-C11-C12-C13
22	B	812	CLA	C3-C5-C6-C7
22	4	811	CLA	C6-C7-C8-C10
22	A	805	CLA	C8-C10-C11-C12
22	A	824	CLA	C8-C10-C11-C12
21	A	801	CL0	CAA-CBA-CGA-O2A
42	7	321	SPH	C12-C13-C14-C15
22	B	810	CLA	C11-C10-C8-C7
22	7	309	CLA	C11-C10-C8-C7
22	5	301	CLA	C11-C10-C8-C7
22	A	806	CLA	C8-C10-C11-C12
22	A	828	CLA	C5-C6-C7-C8
22	B	809	CLA	C8-C10-C11-C12
22	4	808	CLA	C8-C10-C11-C12
26	8	318	NKP	CAS-CAT-CAU-CAV
25	6	323	LHG	C10-C11-C12-C13
22	B	823	CLA	CAD-CBD-CGD-O2D
22	1	311	CLA	CAD-CBD-CGD-O2D
22	3	322	CLA	CAD-CBD-CGD-O2D
22	8	309	CLA	CAD-CBD-CGD-O2D
22	Z	316	CLA	C3-C5-C6-C7
22	A	825	CLA	C13-C15-C16-C17
22	B	824	CLA	C8-C10-C11-C12
22	B	825	CLA	C8-C10-C11-C12
25	7	318	LHG	C12-C13-C14-C15
22	1	316	CLA	CBA-CGA-O2A-C1
22	8	313	CLA	CBA-CGA-O2A-C1
26	8	318	NKP	CAL-CAK-OAJ-CAI
39	8	312	CHL	CBA-CGA-O2A-C1
22	B	820	CLA	C11-C12-C13-C14
25	Z	317	LHG	C30-C31-C32-C33
25	6	323	LHG	C2-C3-O3-P
25	9	315	LHG	C4-C5-C6-O8
40	1	318	SQD	O6-C44-C45-C46

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Mol	Chain	Res	Type	Atoms
22	4	810	CLA	CBD-CGD-O2D-CED
22	B	842	CLA	O1A-CGA-O2A-C1
25	7	318	LHG	O6-C4-C5-O7
25	5	324	LHG	O6-C4-C5-O7
22	B	801	CLA	C5-C6-C7-C8
22	1	306	CLA	C3-C5-C6-C7
22	5	309	CLA	CAA-CBA-CGA-O2A
25	A	849	LHG	C24-C23-O8-C6
22	L	202	CLA	C2A-CAA-CBA-CGA
39	Z	312	CHL	O1A-CGA-O2A-C1
22	A	820	CLA	C11-C12-C13-C14
22	1	310	CLA	C11-C12-C13-C14
22	7	304	CLA	C11-C12-C13-C15
22	A	806	CLA	CHA-CBD-CGD-O1D
22	A	806	CLA	CHA-CBD-CGD-O2D
22	A	814	CLA	CHA-CBD-CGD-O1D
22	A	814	CLA	CHA-CBD-CGD-O2D
22	A	815	CLA	CHA-CBD-CGD-O1D
22	A	815	CLA	CHA-CBD-CGD-O2D
22	A	829	CLA	CHA-CBD-CGD-O2D
22	A	837	CLA	CHA-CBD-CGD-O1D
22	A	839	CLA	CHA-CBD-CGD-O2D
22	B	828	CLA	CHA-CBD-CGD-O2D
22	B	837	CLA	CHA-CBD-CGD-O1D
22	B	837	CLA	CHA-CBD-CGD-O2D
22	G	202	CLA	CHA-CBD-CGD-O1D
22	K	202	CLA	CHA-CBD-CGD-O2D
22	L	201	CLA	CHA-CBD-CGD-O1D
22	L	201	CLA	CHA-CBD-CGD-O2D
22	1	305	CLA	CHA-CBD-CGD-O1D
22	1	305	CLA	CHA-CBD-CGD-O2D
22	Z	304	CLA	CHA-CBD-CGD-O1D
22	Z	305	CLA	CHA-CBD-CGD-O1D
22	Z	305	CLA	CHA-CBD-CGD-O2D
22	7	312	CLA	CHA-CBD-CGD-O2D
22	8	311	CLA	CHA-CBD-CGD-O2D
22	5	314	CLA	CHA-CBD-CGD-O2D
22	5	315	CLA	CHA-CBD-CGD-O2D
22	6	309	CLA	CHA-CBD-CGD-O2D
22	6	312	CLA	CHA-CBD-CGD-O1D
22	6	312	CLA	CHA-CBD-CGD-O2D
22	6	314	CLA	CHA-CBD-CGD-O2D

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Mol	Chain	Res	Type	Atoms
22	6	317	CLA	CHA-CBD-CGD-O1D
22	6	319	CLA	CHA-CBD-CGD-O1D
22	6	319	CLA	CHA-CBD-CGD-O2D
22	6	322	CLA	CHA-CBD-CGD-O1D
22	9	305	CLA	CHA-CBD-CGD-O1D
22	9	307	CLA	CHA-CBD-CGD-O1D
22	9	309	CLA	CHA-CBD-CGD-O2D
22	2	302	CLA	CHA-CBD-CGD-O1D
39	Z	311	CHL	CHA-CBD-CGD-O1D
39	8	301	CHL	CHA-CBD-CGD-O1D
41	8	320	3PH	C3B-C3C-C3D-C3E
22	1	316	CLA	O1A-CGA-O2A-C1
33	F	307	LMT	C1-C2-C3-C4
22	A	813	CLA	C13-C15-C16-C17
25	B	850	LHG	O7-C5-C6-O8
25	B	851	LHG	O7-C5-C6-O8
25	Z	317	LHG	O7-C5-C6-O8
25	3	321	LHG	O7-C5-C6-O8
31	B	852	DGD	O1G-C1G-C2G-O2G
31	3	301	DGD	O1G-C1G-C2G-O2G
34	F	308	LMG	O1-C7-C8-O7
34	F	308	LMG	O7-C8-C9-O8
22	1	316	CLA	C13-C15-C16-C17
25	A	849	LHG	O1-C1-C2-O2
27	A	852	DGA	CA9-CAA-CBA-CCA
40	1	318	SQD	C10-C11-C12-C13
25	1	317	LHG	O10-C23-O8-C6
22	B	842	CLA	C2-C3-C5-C6
22	B	826	CLA	C6-C7-C8-C9
22	F	302	CLA	C11-C10-C8-C9
22	1	306	CLA	C11-C10-C8-C9
22	3	310	CLA	C6-C7-C8-C9
22	7	309	CLA	C6-C7-C8-C9
22	7	310	CLA	C11-C10-C8-C9
22	7	323	CLA	C11-C10-C8-C9
22	6	319	CLA	C11-C12-C13-C14
22	6	301	CLA	C5-C6-C7-C8
40	1	318	SQD	C4-C5-C6-S
22	B	821	CLA	C2A-CAA-CBA-CGA
22	A	824	CLA	CBD-CGD-O2D-CED
22	7	309	CLA	C11-C10-C8-C9
22	B	841	CLA	C8-C10-C11-C12

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Mol	Chain	Res	Type	Atoms
22	Z	313	CLA	CBA-CGA-O2A-C1
24	A	847	BCR	C37-C22-C23-C24
24	6	305	BCR	C37-C22-C23-C24
24	G	203	BCR	C11-C12-C13-C14
24	6	305	BCR	C21-C22-C23-C24
38	4	802	LUT	C11-C12-C13-C14
22	6	301	CLA	C3-C5-C6-C7
22	B	808	CLA	C1A-C2A-CAA-CBA
22	1	308	CLA	C1A-C2A-CAA-CBA
22	1	313	CLA	C1A-C2A-CAA-CBA
22	Z	309	CLA	C1A-C2A-CAA-CBA
22	7	315	CLA	C1A-C2A-CAA-CBA
39	4	814	CHL	C1A-C2A-CAA-CBA
22	B	828	CLA	C8-C10-C11-C12
22	Z	309	CLA	C10-C11-C12-C13
22	1	316	CLA	C2-C1-O2A-CGA
22	3	312	CLA	C2-C1-O2A-CGA
22	4	812	CLA	C2-C1-O2A-CGA
22	4	815	CLA	C2-C1-O2A-CGA
24	3	307	BCR	C19-C20-C21-C22
25	8	317	LHG	C4-O6-P-O3
25	6	323	LHG	C34-C35-C36-C37
22	A	838	CLA	C3-C5-C6-C7
22	A	822	CLA	C2-C3-C5-C6
39	6	315	CHL	C2-C3-C5-C6
41	7	319	3PH	C33-C34-C35-C36
22	Z	313	CLA	O1A-CGA-O2A-C1
26	8	318	NKP	OAE-CAK-OAJ-CAI
25	A	849	LHG	C3-O3-P-O5
25	A	850	LHG	C4-O6-P-O4
25	B	803	LHG	C3-O3-P-O5
25	B	850	LHG	C3-O3-P-O4
25	4	820	LHG	C4-O6-P-O5
25	5	324	LHG	C3-O3-P-O4
25	5	324	LHG	C3-O3-P-O5
25	9	315	LHG	C4-O6-P-O5
25	A	849	LHG	O6-C4-C5-C6
35	J	102	T7X	O13-C7-C8-C9
41	7	319	3PH	O11-C1-C2-C3
22	3	310	CLA	C8-C10-C11-C12
22	1	310	CLA	C3-C5-C6-C7
22	3	313	CLA	C3-C5-C6-C7

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Mol	Chain	Res	Type	Atoms
25	A	849	LHG	C11-C10-C9-C8
25	6	323	LHG	C11-C10-C9-C8
26	A	851	NKP	CAG-CAH-CAI-OAJ
23	A	843	PQN	C26-C27-C28-C29
22	A	826	CLA	CAD-CBD-CGD-O1D
22	K	205	CLA	CAD-CBD-CGD-O1D
22	1	309	CLA	CAD-CBD-CGD-O1D
22	8	307	CLA	CAD-CBD-CGD-O1D
22	9	305	CLA	CAD-CBD-CGD-O1D
39	Z	311	CHL	CAD-CBD-CGD-O1D
39	8	301	CHL	CAD-CBD-CGD-O1D
22	A	840	CLA	C5-C6-C7-C8
22	F	302	CLA	C13-C15-C16-C17
22	7	308	CLA	C5-C6-C7-C8
22	1	308	CLA	C5-C6-C7-C8
22	A	807	CLA	C11-C12-C13-C15
22	7	324	CLA	C4-C3-C5-C6
22	5	311	CLA	C4-C3-C5-C6
22	A	802	CLA	C11-C12-C13-C15
22	A	807	CLA	C6-C7-C8-C10
22	A	808	CLA	C11-C12-C13-C15
22	A	813	CLA	C11-C12-C13-C15
22	A	814	CLA	C6-C7-C8-C10
22	A	814	CLA	C11-C10-C8-C7
22	A	818	CLA	C6-C7-C8-C10
22	A	835	CLA	C6-C7-C8-C10
22	A	835	CLA	C11-C10-C8-C7
22	A	840	CLA	C11-C10-C8-C7
22	A	854	CLA	C11-C12-C13-C15
22	B	801	CLA	C6-C7-C8-C10
22	B	809	CLA	C6-C7-C8-C10
22	B	813	CLA	C6-C7-C8-C10
22	B	816	CLA	C11-C10-C8-C7
22	B	820	CLA	C6-C7-C8-C10
22	B	825	CLA	C11-C10-C8-C7
22	B	826	CLA	C11-C10-C8-C7
22	B	829	CLA	C6-C7-C8-C10
22	B	830	CLA	C11-C12-C13-C15
22	B	833	CLA	C6-C7-C8-C10
22	B	834	CLA	C11-C10-C8-C7
22	B	838	CLA	C11-C12-C13-C15
22	B	840	CLA	C11-C12-C13-C15

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Mol	Chain	Res	Type	Atoms
22	F	301	CLA	C11-C10-C8-C7
22	F	302	CLA	C11-C10-C8-C7
22	1	304	CLA	C6-C7-C8-C10
22	1	307	CLA	C6-C7-C8-C10
22	1	307	CLA	C11-C10-C8-C7
22	1	308	CLA	C3A-C2A-CAA-CBA
22	1	309	CLA	C11-C10-C8-C7
22	Z	303	CLA	C6-C7-C8-C10
22	Z	306	CLA	C11-C10-C8-C7
22	Z	309	CLA	C11-C10-C8-C7
22	Z	316	CLA	C12-C13-C15-C16
22	3	308	CLA	C11-C12-C13-C15
22	3	318	CLA	C6-C7-C8-C10
22	3	322	CLA	C6-C7-C8-C10
22	7	306	CLA	C11-C10-C8-C7
22	7	306	CLA	C11-C12-C13-C15
22	7	308	CLA	C11-C10-C8-C7
22	7	309	CLA	C6-C7-C8-C10
22	7	310	CLA	C11-C10-C8-C7
22	8	305	CLA	C11-C10-C8-C7
22	8	310	CLA	C6-C7-C8-C10
22	5	308	CLA	C11-C12-C13-C15
22	5	312	CLA	C3A-C2A-CAA-CBA
22	5	315	CLA	C11-C12-C13-C15
22	6	301	CLA	C11-C10-C8-C7
22	6	302	CLA	C11-C10-C8-C7
22	6	309	CLA	C6-C7-C8-C10
22	6	310	CLA	C11-C10-C8-C7
22	6	310	CLA	C11-C12-C13-C15
22	6	319	CLA	C11-C10-C8-C7
22	6	319	CLA	C11-C12-C13-C15
22	9	303	CLA	C11-C10-C8-C7
23	B	843	PQN	C16-C17-C18-C20
25	A	849	LHG	O6-C4-C5-O7
25	B	803	LHG	O6-C4-C5-O7
25	B	851	LHG	O6-C4-C5-O7
25	6	323	LHG	O6-C4-C5-O7
39	Z	312	CHL	C3A-C2A-CAA-CBA
41	7	319	3PH	O11-C1-C2-O21
41	6	324	3PH	O11-C1-C2-O21
22	4	810	CLA	O1D-CGD-O2D-CED
25	5	324	LHG	C26-C27-C28-C29

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Mol	Chain	Res	Type	Atoms
24	G	203	BCR	C13-C14-C15-C16
22	A	838	CLA	CAA-CBA-CGA-O2A
22	A	827	CLA	C15-C16-C17-C18
22	7	309	CLA	C5-C6-C7-C8
22	8	313	CLA	O1A-CGA-O2A-C1
25	A	849	LHG	O10-C23-O8-C6
39	8	312	CHL	O1A-CGA-O2A-C1
25	A	850	LHG	C30-C31-C32-C33
22	B	835	CLA	CBD-CGD-O2D-CED
22	A	813	CLA	C2A-CAA-CBA-CGA
39	3	317	CHL	C2A-CAA-CBA-CGA
22	Z	306	CLA	C11-C12-C13-C15
25	B	850	LHG	C4-C5-C6-O8
25	4	820	LHG	C4-C5-C6-O8
31	1	319	DGD	C1G-C2G-C3G-O3G
34	F	308	LMG	O1-C7-C8-C9
25	1	317	LHG	O7-C5-C6-O8
25	4	820	LHG	O7-C5-C6-O8
25	7	318	LHG	C30-C31-C32-C33
31	1	319	DGD	C5D-C6D-O5D-C1E
22	3	311	CLA	C8-C10-C11-C12
22	5	313	CLA	C5-C6-C7-C8
22	6	307	CLA	C8-C10-C11-C12
22	L	202	CLA	O1D-CGD-O2D-CED
22	A	816	CLA	C11-C10-C8-C9
22	A	805	CLA	C14-C13-C15-C16
22	A	806	CLA	C11-C10-C8-C9
22	A	817	CLA	C11-C10-C8-C9
22	A	828	CLA	C11-C12-C13-C14
22	A	830	CLA	C11-C10-C8-C9
22	A	833	CLA	C11-C10-C8-C9
22	B	806	CLA	C11-C12-C13-C14
22	B	814	CLA	C11-C10-C8-C9
22	B	818	CLA	C11-C10-C8-C9
22	B	819	CLA	C11-C10-C8-C9
22	B	821	CLA	C11-C12-C13-C14
22	B	826	CLA	C11-C12-C13-C14
22	B	827	CLA	C11-C12-C13-C14
22	B	833	CLA	C11-C10-C8-C9
22	Z	316	CLA	C14-C13-C15-C16
22	7	308	CLA	C11-C12-C13-C14
22	7	317	CLA	C6-C7-C8-C9

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Mol	Chain	Res	Type	Atoms
22	7	317	CLA	C11-C10-C8-C9
22	8	308	CLA	C11-C10-C8-C9
22	5	322	CLA	C6-C7-C8-C9
22	6	302	CLA	C11-C10-C8-C9
39	5	316	CHL	C6-C7-C8-C9
22	B	813	CLA	C3-C5-C6-C7
22	F	304	CLA	C16-C17-C18-C20
22	8	310	CLA	C11-C12-C13-C15
22	1	311	CLA	C2A-CAA-CBA-CGA
39	6	315	CHL	C2A-CAA-CBA-CGA
22	A	805	CLA	CAA-CBA-CGA-O2A
22	B	842	CLA	CAA-CBA-CGA-O2A
22	6	302	CLA	CAA-CBA-CGA-O2A
25	4	820	LHG	O1-C1-C2-O2
24	B	848	BCR	C18-C19-C20-C21
24	3	304	BCR	C10-C11-C12-C13
24	3	307	BCR	C18-C19-C20-C21
24	5	305	BCR	C18-C19-C20-C21
38	6	303	LUT	C27-C28-C29-C39
25	6	323	LHG	C13-C14-C15-C16
22	A	809	CLA	C6-C7-C8-C10
22	B	805	CLA	C16-C17-C18-C20
22	B	820	CLA	C11-C12-C13-C15
27	A	852	DGA	CBB-CCB-CDB-CEB
39	3	317	CHL	O1D-CGD-O2D-CED
22	A	834	CLA	C4-C3-C5-C6
22	A	802	CLA	O1A-CGA-O2A-C1
22	6	312	CLA	C15-C16-C17-C18
26	8	318	NKP	CAQ-CAR-CAS-CAT
22	A	811	CLA	C8-C10-C11-C12
22	7	312	CLA	C8-C10-C11-C12
34	F	308	LMG	C9-C8-O7-C10
25	B	851	LHG	O6-C4-C5-C6
22	A	854	CLA	C2A-CAA-CBA-CGA
22	B	805	CLA	C2A-CAA-CBA-CGA
22	B	810	CLA	C2A-CAA-CBA-CGA
22	B	812	CLA	C2A-CAA-CBA-CGA
22	Z	314	CLA	C2A-CAA-CBA-CGA
22	8	316	CLA	C2A-CAA-CBA-CGA
22	6	312	CLA	O1D-CGD-O2D-CED
21	A	801	CL0	C2-C1-O2A-CGA
22	A	816	CLA	C2-C1-O2A-CGA

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Mol	Chain	Res	Type	Atoms
22	7	308	CLA	C2-C1-O2A-CGA
25	A	850	LHG	C27-C28-C29-C30
25	8	317	LHG	C29-C30-C31-C32
25	B	803	LHG	C26-C27-C28-C29
25	3	321	LHG	C5-C4-O6-P
41	5	325	3PH	O11-C1-C2-O21
22	6	309	CLA	C16-C17-C18-C19
24	B	847	BCR	C23-C24-C25-C30
24	B	853	BCR	C5-C6-C7-C8
24	7	303	BCR	C5-C6-C7-C8
38	1	302	LUT	C1-C6-C7-C8
38	1	303	LUT	C5-C6-C7-C8
38	Z	302	LUT	C1-C6-C7-C8
38	6	304	LUT	C1-C6-C7-C8
39	9	312	CHL	C2-C3-C5-C6
22	B	835	CLA	O1D-CGD-O2D-CED
39	5	321	CHL	O1D-CGD-O2D-CED
25	7	318	LHG	C10-C11-C12-C13
42	7	321	SPH	C14-C15-C16-C17
22	7	324	CLA	C11-C12-C13-C15
39	4	816	CHL	CBD-CGD-O2D-CED
22	B	825	CLA	C2A-CAA-CBA-CGA
25	9	315	LHG	O7-C5-C6-O8
22	6	312	CLA	CBD-CGD-O2D-CED
25	A	850	LHG	C3-O3-P-O6
25	B	851	LHG	C3-O3-P-O6
25	3	321	LHG	C3-O3-P-O6
25	7	318	LHG	C3-O3-P-O6
25	4	820	LHG	C3-O3-P-O6
25	5	324	LHG	C4-O6-P-O3
43	8	319	LPX	C3-O1-P1-O2
31	1	319	DGD	C6A-C7A-C8A-C9A
25	A	849	LHG	C4-C5-C6-O8
25	8	317	LHG	C4-C5-C6-O8
22	A	814	CLA	C2-C3-C5-C6
22	A	824	CLA	C11-C10-C8-C7
22	A	825	CLA	C6-C7-C8-C10
22	B	817	CLA	C11-C10-C8-C7
22	F	302	CLA	C11-C12-C13-C15
22	1	304	CLA	C11-C12-C13-C15
22	Z	308	CLA	C11-C10-C8-C7
22	7	323	CLA	C11-C12-C13-C15

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Mol	Chain	Res	Type	Atoms
22	A	835	CLA	C11-C10-C8-C9
22	A	840	CLA	C11-C10-C8-C9
22	B	811	CLA	C11-C10-C8-C9
22	B	826	CLA	C11-C10-C8-C9
22	B	830	CLA	C11-C12-C13-C14
22	B	834	CLA	C11-C10-C8-C9
22	F	301	CLA	C11-C10-C8-C9
22	Z	306	CLA	C11-C10-C8-C9
22	Z	309	CLA	C11-C10-C8-C9
22	7	304	CLA	C11-C10-C8-C9
22	7	307	CLA	C11-C12-C13-C14
22	7	308	CLA	C11-C10-C8-C9
22	8	310	CLA	C6-C7-C8-C9
22	5	308	CLA	C11-C12-C13-C14
22	6	301	CLA	C11-C10-C8-C9
22	6	309	CLA	C6-C7-C8-C9
22	6	319	CLA	C11-C10-C8-C9
22	B	821	CLA	C13-C15-C16-C17
24	3	305	BCR	C15-C16-C17-C18
38	4	803	LUT	C9-C10-C11-C12
38	4	803	LUT	C33-C34-C35-C15
22	4	807	CLA	C16-C17-C18-C20
42	7	320	SPH	C11-C10-C9-C8
41	8	320	3PH	O22-C21-C22-C23
22	Z	313	CLA	C5-C6-C7-C8
22	8	307	CLA	O1A-CGA-O2A-C1
24	3	306	BCR	C7-C8-C9-C34
22	A	807	CLA	C11-C12-C13-C14
22	A	816	CLA	C11-C10-C8-C7
22	B	813	CLA	C11-C10-C8-C7
22	Z	305	CLA	C11-C10-C8-C7
22	5	309	CLA	C11-C10-C8-C7
26	A	851	NKP	OAF-CAG-CAH-CAI
22	A	816	CLA	C5-C6-C7-C8
22	5	311	CLA	C2-C3-C5-C6
39	6	318	CHL	C2-C3-C5-C6
22	A	802	CLA	CBA-CGA-O2A-C1
22	8	307	CLA	CBA-CGA-O2A-C1
22	F	302	CLA	C8-C10-C11-C12
22	6	302	CLA	C8-C10-C11-C12
42	7	320	SPH	C10-C11-C12-C13
22	A	824	CLA	O1D-CGD-O2D-CED

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Mol	Chain	Res	Type	Atoms
25	8	317	LHG	C26-C27-C28-C29
22	B	817	CLA	C2A-CAA-CBA-CGA
22	8	305	CLA	C2A-CAA-CBA-CGA
22	5	310	CLA	C2A-CAA-CBA-CGA
38	8	302	LUT	C9-C10-C11-C12
38	5	302	LUT	C9-C10-C11-C12
25	5	324	LHG	C25-C26-C27-C28
22	B	824	CLA	C3-C5-C6-C7
22	7	304	CLA	C3-C5-C6-C7
22	A	802	CLA	C8-C10-C11-C12
22	B	820	CLA	C8-C10-C11-C12
40	1	318	SQD	C7-C8-C9-C10
22	6	307	CLA	C11-C12-C13-C15
22	L	202	CLA	C4-C3-C5-C6
22	3	312	CLA	C4-C3-C5-C6
22	A	834	CLA	C2-C3-C5-C6
22	L	202	CLA	C2-C3-C5-C6
22	3	312	CLA	C2-C3-C5-C6
41	7	319	3PH	C34-C35-C36-C37
22	5	318	CLA	C5-C6-C7-C8
25	6	323	LHG	C33-C34-C35-C36
22	A	817	CLA	C2-C1-O2A-CGA
22	B	816	CLA	C2-C1-O2A-CGA
22	Z	303	CLA	C2-C1-O2A-CGA
22	4	811	CLA	C2-C1-O2A-CGA
22	9	309	CLA	C2-C1-O2A-CGA
22	Z	306	CLA	C11-C12-C13-C14
22	4	808	CLA	C11-C12-C13-C15
26	8	318	NKP	CAR-CAS-CAT-CAU
22	6	310	CLA	C2A-CAA-CBA-CGA
31	1	319	DGD	O6D-C5D-C6D-O5D
22	Z	316	CLA	C3A-C2A-CAA-CBA
22	5	309	CLA	C3A-C2A-CAA-CBA
22	6	309	CLA	C3A-C2A-CAA-CBA
39	4	814	CHL	C3A-C2A-CAA-CBA
22	B	826	CLA	CAA-CBA-CGA-O2A
36	J	105	C7Z	C29-C30-C31-C32
27	A	852	DGA	CBB-CAB-CB9-CB8
22	B	829	CLA	C4-C3-C5-C6
22	7	309	CLA	C4-C3-C5-C6
42	7	320	SPH	C7-C8-C9-C10
22	B	821	CLA	C11-C10-C8-C9

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Mol	Chain	Res	Type	Atoms
22	B	838	CLA	C11-C12-C13-C14
22	1	304	CLA	C14-C13-C15-C16
22	1	307	CLA	C11-C10-C8-C9
22	3	313	CLA	C6-C7-C8-C9
22	7	306	CLA	C11-C10-C8-C9
22	7	306	CLA	C11-C12-C13-C14
22	8	309	CLA	C11-C10-C8-C9
22	6	310	CLA	C11-C10-C8-C9
22	6	314	CLA	C11-C10-C8-C9
39	5	316	CHL	C11-C12-C13-C14
22	7	312	CLA	C11-C12-C13-C14
22	A	804	CLA	CAA-CBA-CGA-O1A
22	3	313	CLA	C13-C15-C16-C17
28	A	853	OCA	C3-C4-C5-C6
22	5	315	CLA	C15-C16-C17-C18
24	A	844	BCR	C11-C10-C9-C34
24	A	845	BCR	C35-C13-C14-C15
24	A	848	BCR	C11-C10-C9-C34
24	A	848	BCR	C16-C17-C18-C36
24	B	844	BCR	C35-C13-C14-C15
24	B	844	BCR	C16-C17-C18-C36
24	B	846	BCR	C11-C10-C9-C34
24	B	846	BCR	C20-C21-C22-C37
24	B	848	BCR	C20-C21-C22-C37
24	B	853	BCR	C11-C10-C9-C34
32	F	306	RRX	C16-C17-C18-C36
32	F	306	RRX	C35-C13-C14-C15
32	F	306	RRX	C11-C10-C9-C34
22	A	802	CLA	C2A-CAA-CBA-CGA
22	7	324	CLA	C11-C12-C13-C14
22	A	810	CLA	O2A-C1-C2-C3
22	A	839	CLA	O2A-C1-C2-C3
22	1	305	CLA	CAA-CBA-CGA-O1A
24	B	844	BCR	C11-C12-C13-C35
38	Z	302	LUT	C31-C32-C33-C40
22	5	308	CLA	C5-C6-C7-C8
25	4	820	LHG	C25-C26-C27-C28
34	J	101	LMG	C9-C8-O7-C10
22	7	322	CLA	C5-C6-C7-C8
22	B	833	CLA	C4-C3-C5-C6
22	A	834	CLA	C1A-C2A-CAA-CBA
22	Z	316	CLA	C1A-C2A-CAA-CBA

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Mol	Chain	Res	Type	Atoms
22	5	309	CLA	C1A-C2A-CAA-CBA
22	6	310	CLA	C1A-C2A-CAA-CBA
39	Z	311	CHL	C1A-C2A-CAA-CBA
39	Z	312	CHL	C1A-C2A-CAA-CBA
22	A	805	CLA	C11-C10-C8-C7
22	A	821	CLA	C11-C12-C13-C15
22	A	825	CLA	C11-C10-C8-C7
22	A	830	CLA	C6-C7-C8-C10
22	A	832	CLA	C6-C7-C8-C10
22	A	840	CLA	C11-C12-C13-C15
22	B	805	CLA	C11-C10-C8-C7
22	B	809	CLA	C11-C12-C13-C15
22	B	831	CLA	C11-C10-C8-C7
22	1	310	CLA	C6-C7-C8-C10
22	Z	313	CLA	C6-C7-C8-C10
22	5	307	CLA	C11-C10-C8-C7
22	5	308	CLA	C11-C10-C8-C7
22	6	309	CLA	C11-C10-C8-C7
26	A	851	NKP	CAQ-CAR-CAS-CAT
41	8	320	3PH	O21-C21-C22-C23
24	B	847	BCR	C13-C14-C15-C16
38	5	303	LUT	C9-C10-C11-C12
25	1	317	LHG	C17-C18-C19-C20
35	J	102	T7X	C16-C17-C18-C19
33	G	204	LMT	C5-C6-C7-C8
22	8	314	CLA	CAA-CBA-CGA-O2A
31	B	852	DGD	O1A-C1A-O1G-C1G
25	6	323	LHG	C14-C15-C16-C17
39	4	816	CHL	O1D-CGD-O2D-CED
22	A	805	CLA	C2A-CAA-CBA-CGA
22	B	809	CLA	C2A-CAA-CBA-CGA
22	B	822	CLA	C2A-CAA-CBA-CGA
22	A	808	CLA	C5-C6-C7-C8
22	B	811	CLA	C15-C16-C17-C18
22	3	308	CLA	C15-C16-C17-C18
22	3	312	CLA	C5-C6-C7-C8
22	A	835	CLA	C8-C10-C11-C12
22	A	838	CLA	C8-C10-C11-C12
22	B	806	CLA	CBA-CGA-O2A-C1
27	A	852	DGA	CB6-CB7-CB8-CB9
22	A	811	CLA	C10-C11-C12-C13
22	B	831	CLA	C8-C10-C11-C12

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Mol	Chain	Res	Type	Atoms
22	3	314	CLA	C5-C6-C7-C8
22	6	319	CLA	C8-C10-C11-C12
22	B	807	CLA	CAA-CBA-CGA-O2A
25	A	849	LHG	C10-C11-C12-C13
40	1	318	SQD	C24-C25-C26-C27
22	8	308	CLA	C5-C6-C7-C8
22	B	829	CLA	C2-C3-C5-C6
22	6	301	CLA	C8-C10-C11-C12
22	7	323	CLA	C16-C17-C18-C20
24	A	844	BCR	C11-C10-C9-C8
24	A	845	BCR	C12-C13-C14-C15
24	A	848	BCR	C11-C10-C9-C8
24	A	848	BCR	C16-C17-C18-C19
24	B	844	BCR	C12-C13-C14-C15
24	B	844	BCR	C16-C17-C18-C19
24	B	846	BCR	C11-C10-C9-C8
24	B	846	BCR	C20-C21-C22-C23
24	B	853	BCR	C11-C10-C9-C8
32	F	306	RRX	C16-C17-C18-C19
32	F	306	RRX	C12-C13-C14-C15
32	F	306	RRX	C11-C10-C9-C8
22	B	833	CLA	C8-C10-C11-C12
22	A	820	CLA	C2A-CAA-CBA-CGA
24	A	845	BCR	C15-C16-C17-C18
24	3	307	BCR	C15-C16-C17-C18
38	7	302	LUT	C9-C10-C11-C12
22	Z	304	CLA	CAA-CBA-CGA-O1A
22	6	309	CLA	C16-C17-C18-C20
22	1	310	CLA	C2C-C3C-CAC-CBC
41	6	324	3PH	C29-C2A-C2B-C2C
22	Z	316	CLA	C13-C15-C16-C17
25	4	820	LHG	O10-C23-C24-C25
22	A	836	CLA	C2-C1-O2A-CGA
22	7	309	CLA	C2-C3-C5-C6
22	B	807	CLA	CAA-CBA-CGA-O1A
22	1	305	CLA	CAA-CBA-CGA-O2A
27	A	852	DGA	CAA-CBA-CCA-CDA
22	A	828	CLA	CAA-CBA-CGA-O2A
22	5	311	CLA	CAA-CBA-CGA-O2A
22	A	814	CLA	C11-C10-C8-C9
22	B	815	CLA	C11-C10-C8-C9
22	6	310	CLA	C11-C12-C13-C14

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Mol	Chain	Res	Type	Atoms
42	7	320	SPH	O1-C1-C2-C3
22	2	304	CLA	CAA-CBA-CGA-O2A
26	A	851	NKP	CAN-CAO-CAP-CAQ
42	7	320	SPH	C13-C14-C15-C16
22	7	308	CLA	C2A-CAA-CBA-CGA
22	4	805	CLA	C2A-CAA-CBA-CGA
22	B	814	CLA	C11-C12-C13-C15
22	1	307	CLA	C11-C12-C13-C15
22	3	314	CLA	C11-C12-C13-C15
22	A	804	CLA	CAA-CBA-CGA-O2A
22	9	310	CLA	CAA-CBA-CGA-O1A
24	A	845	BCR	C23-C24-C25-C30
24	B	844	BCR	C23-C24-C25-C30
24	3	307	BCR	C23-C24-C25-C30
36	J	105	C7Z	C21-C26-C27-C28
38	1	303	LUT	C1-C6-C7-C8
22	B	834	CLA	CAA-CBA-CGA-O2A
41	6	324	3PH	O21-C21-C22-C23
25	3	321	LHG	O1-C1-C2-C3
25	6	323	LHG	C16-C17-C18-C19
42	7	321	SPH	C13-C14-C15-C16
24	A	848	BCR	C9-C10-C11-C12
24	B	844	BCR	C21-C22-C23-C24
38	Z	301	LUT	C27-C28-C29-C30
38	3	303	LUT	C31-C32-C33-C34
22	L	201	CLA	C5-C6-C7-C8
22	B	833	CLA	C2-C3-C5-C6
22	Z	304	CLA	CAA-CBA-CGA-O2A
22	9	310	CLA	CAA-CBA-CGA-O2A
22	6	322	CLA	O1A-CGA-O2A-C1
39	6	318	CHL	O1A-CGA-O2A-C1
22	Z	310	CLA	CAA-CBA-CGA-O2A
22	6	301	CLA	C11-C12-C13-C15
22	Z	305	CLA	C11-C10-C8-C9
42	7	320	SPH	C11-C12-C13-C14
22	B	809	CLA	CAA-CBA-CGA-O2A
22	B	815	CLA	CAA-CBA-CGA-O2A
22	B	806	CLA	O1A-CGA-O2A-C1
22	B	820	CLA	C5-C6-C7-C8
25	A	850	LHG	C9-C10-C11-C12
25	8	317	LHG	C24-C25-C26-C27
22	7	306	CLA	C3-C5-C6-C7

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Mol	Chain	Res	Type	Atoms
22	Z	316	CLA	C15-C16-C17-C18
22	Z	309	CLA	O1D-CGD-O2D-CED
25	A	850	LHG	C25-C26-C27-C28
21	A	801	CL0	C11-C12-C13-C15
22	A	832	CLA	C11-C12-C13-C15
22	B	828	CLA	C11-C12-C13-C15
22	B	830	CLA	C2-C3-C5-C6
22	3	313	CLA	C6-C7-C8-C10
22	7	323	CLA	C12-C13-C15-C16
22	6	309	CLA	C11-C12-C13-C15
39	3	317	CHL	C6-C7-C8-C10
22	9	304	CLA	CAA-CBA-CGA-O2A
22	4	807	CLA	C13-C15-C16-C17
25	B	803	LHG	C2-C3-O3-P
34	F	308	LMG	O10-C28-O8-C9
31	B	852	DGD	C2A-C1A-O1G-C1G
34	F	308	LMG	C29-C28-O8-C9
22	A	823	CLA	C3-C5-C6-C7
22	A	807	CLA	CAA-CBA-CGA-O2A
22	B	810	CLA	CAA-CBA-CGA-O2A
22	L	203	CLA	CAA-CBA-CGA-O2A
22	3	322	CLA	CAA-CBA-CGA-O2A
22	1	313	CLA	C5-C6-C7-C8
22	6	307	CLA	C11-C12-C13-C14
39	6	318	CHL	CBA-CGA-O2A-C1
22	A	808	CLA	C13-C15-C16-C17
25	A	850	LHG	C11-C10-C9-C8
22	A	841	CLA	C4-C3-C5-C6
22	5	310	CLA	C4-C3-C5-C6
22	B	808	CLA	C8-C10-C11-C12
22	F	304	CLA	C8-C10-C11-C12
22	7	324	CLA	C8-C10-C11-C12
43	8	319	LPX	C1-O2-P1-O1
22	A	825	CLA	C2-C3-C5-C6
22	4	808	CLA	C11-C12-C13-C14
22	6	309	CLA	CAA-CBA-CGA-O1A
22	B	833	CLA	C13-C15-C16-C17
22	B	822	CLA	C11-C10-C8-C7
22	A	802	CLA	C11-C12-C13-C14
22	A	808	CLA	C11-C12-C13-C14
22	A	808	CLA	C14-C13-C15-C16
22	A	811	CLA	C14-C13-C15-C16

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Mol	Chain	Res	Type	Atoms
22	A	854	CLA	C11-C12-C13-C14
22	B	805	CLA	C11-C10-C8-C9
22	B	809	CLA	C11-C12-C13-C14
22	B	816	CLA	C11-C10-C8-C9
22	B	817	CLA	C11-C10-C8-C9
22	B	825	CLA	C11-C10-C8-C9
22	B	831	CLA	C11-C10-C8-C9
22	1	304	CLA	C11-C12-C13-C14
22	1	309	CLA	C11-C10-C8-C9
22	3	308	CLA	C11-C12-C13-C14
22	3	313	CLA	C11-C10-C8-C9
22	8	305	CLA	C11-C10-C8-C9
22	5	307	CLA	C11-C10-C8-C9
22	5	315	CLA	C11-C12-C13-C14
22	6	322	CLA	C11-C10-C8-C9
22	9	303	CLA	C11-C10-C8-C9
23	B	843	PQN	C16-C17-C18-C19
23	B	843	PQN	C19-C18-C20-C21
39	8	315	CHL	C6-C7-C8-C9
31	B	852	DGD	C9B-CAB-CBB-CCB
25	8	317	LHG	C30-C31-C32-C33
22	B	807	CLA	C3A-C2A-CAA-CBA
22	B	838	CLA	C3A-C2A-CAA-CBA
22	F	304	CLA	C3A-C2A-CAA-CBA
22	Z	308	CLA	C3A-C2A-CAA-CBA
22	2	302	CLA	C3A-C2A-CAA-CBA
22	A	812	CLA	CAA-CBA-CGA-O2A
22	3	308	CLA	CAA-CBA-CGA-O2A
25	A	849	LHG	O7-C7-C8-C9
25	1	317	LHG	O8-C23-C24-C25
31	3	301	DGD	O1G-C1A-C2A-C3A
41	8	320	3PH	C38-C39-C3A-C3B
22	2	302	CLA	CAA-CBA-CGA-O2A
22	B	813	CLA	CAD-CBD-CGD-O2D
22	3	315	CLA	CAD-CBD-CGD-O2D
22	5	323	CLA	CAD-CBD-CGD-O2D
39	1	312	CHL	CAD-CBD-CGD-O2D
39	Z	312	CHL	CAD-CBD-CGD-O2D
39	4	813	CHL	CAD-CBD-CGD-O2D
39	6	318	CHL	CAD-CBD-CGD-O2D
39	9	312	CHL	CAD-CBD-CGD-O2D
22	7	317	CLA	C2A-CAA-CBA-CGA

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Mol	Chain	Res	Type	Atoms
25	Z	317	LHG	C24-C25-C26-C27
22	A	821	CLA	C2-C1-O2A-CGA
22	B	838	CLA	C2-C1-O2A-CGA
22	A	837	CLA	CAA-CBA-CGA-O2A
22	4	806	CLA	CAA-CBA-CGA-O2A
22	5	323	CLA	CAA-CBA-CGA-O2A
22	9	306	CLA	CAA-CBA-CGA-O2A
22	9	304	CLA	CAA-CBA-CGA-O1A
22	B	824	CLA	CAA-CBA-CGA-O2A
22	1	306	CLA	CAA-CBA-CGA-O2A
22	3	314	CLA	CAA-CBA-CGA-O2A
24	B	844	BCR	C11-C12-C13-C14
24	3	306	BCR	C7-C8-C9-C10
38	Z	302	LUT	C31-C32-C33-C34
38	6	303	LUT	C27-C28-C29-C30
22	B	813	CLA	C11-C10-C8-C9
22	B	822	CLA	C11-C10-C8-C9
22	5	309	CLA	C11-C10-C8-C9
42	7	321	SPH	C5-C6-C7-C8
25	A	850	LHG	C35-C36-C37-C38
22	7	324	CLA	C10-C11-C12-C13
25	A	850	LHG	C2-C3-O3-P
41	8	320	3PH	C1-O11-P-O12
22	1	304	CLA	C8-C10-C11-C12
22	Z	305	CLA	C5-C6-C7-C8
22	A	815	CLA	CAA-CBA-CGA-O2A
22	8	309	CLA	CAA-CBA-CGA-O2A
22	A	831	CLA	O2A-C1-C2-C3
22	1	309	CLA	O2A-C1-C2-C3
22	5	301	CLA	O2A-C1-C2-C3
22	5	320	CLA	O2A-C1-C2-C3
22	9	309	CLA	O2A-C1-C2-C3
39	3	317	CHL	O2A-C1-C2-C3
39	4	814	CHL	O2A-C1-C2-C3
39	5	317	CHL	O2A-C1-C2-C3
39	9	312	CHL	O2A-C1-C2-C3
22	B	812	CLA	CAA-CBA-CGA-O2A
22	7	324	CLA	CAA-CBA-CGA-O2A
22	9	305	CLA	CAA-CBA-CGA-O2A
22	2	302	CLA	CAA-CBA-CGA-O1A
25	1	317	LHG	C28-C29-C30-C31
22	A	802	CLA	CHA-CBD-CGD-O1D

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Mol	Chain	Res	Type	Atoms
22	A	802	CLA	CHA-CBD-CGD-O2D
22	A	805	CLA	CHA-CBD-CGD-O1D
22	A	805	CLA	CHA-CBD-CGD-O2D
22	A	810	CLA	CHA-CBD-CGD-O1D
22	A	810	CLA	CHA-CBD-CGD-O2D
22	A	811	CLA	CHA-CBD-CGD-O2D
22	A	813	CLA	CHA-CBD-CGD-O1D
22	A	813	CLA	CHA-CBD-CGD-O2D
22	A	818	CLA	CHA-CBD-CGD-O1D
22	A	818	CLA	CHA-CBD-CGD-O2D
22	A	819	CLA	CHA-CBD-CGD-O2D
22	A	820	CLA	CHA-CBD-CGD-O1D
22	A	820	CLA	CHA-CBD-CGD-O2D
22	A	821	CLA	CHA-CBD-CGD-O2D
22	A	823	CLA	CHA-CBD-CGD-O1D
22	A	823	CLA	CHA-CBD-CGD-O2D
22	A	824	CLA	CHA-CBD-CGD-O1D
22	A	824	CLA	CHA-CBD-CGD-O2D
22	A	827	CLA	CHA-CBD-CGD-O1D
22	A	827	CLA	CHA-CBD-CGD-O2D
22	A	828	CLA	CHA-CBD-CGD-O1D
22	A	828	CLA	CHA-CBD-CGD-O2D
22	A	831	CLA	CHA-CBD-CGD-O1D
22	A	831	CLA	CHA-CBD-CGD-O2D
22	A	832	CLA	CHA-CBD-CGD-O2D
22	A	835	CLA	CHA-CBD-CGD-O1D
22	A	835	CLA	CHA-CBD-CGD-O2D
22	A	837	CLA	CHA-CBD-CGD-O2D
22	A	838	CLA	CHA-CBD-CGD-O2D
22	A	840	CLA	CHA-CBD-CGD-O1D
22	A	840	CLA	CHA-CBD-CGD-O2D
22	A	854	CLA	CHA-CBD-CGD-O1D
22	A	854	CLA	CHA-CBD-CGD-O2D
22	A	855	CLA	CHA-CBD-CGD-O1D
22	A	855	CLA	CHA-CBD-CGD-O2D
22	B	801	CLA	CHA-CBD-CGD-O1D
22	B	801	CLA	CHA-CBD-CGD-O2D
22	B	806	CLA	CHA-CBD-CGD-O1D
22	B	806	CLA	CHA-CBD-CGD-O2D
22	B	807	CLA	CHA-CBD-CGD-O1D
22	B	807	CLA	CHA-CBD-CGD-O2D
22	B	808	CLA	CHA-CBD-CGD-O2D

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Mol	Chain	Res	Type	Atoms
22	B	809	CLA	CHA-CBD-CGD-O1D
22	B	809	CLA	CHA-CBD-CGD-O2D
22	B	810	CLA	CHA-CBD-CGD-O1D
22	B	810	CLA	CHA-CBD-CGD-O2D
22	B	812	CLA	CHA-CBD-CGD-O2D
22	B	814	CLA	CHA-CBD-CGD-O1D
22	B	814	CLA	CHA-CBD-CGD-O2D
22	B	816	CLA	CHA-CBD-CGD-O1D
22	B	816	CLA	CHA-CBD-CGD-O2D
22	B	818	CLA	CHA-CBD-CGD-O1D
22	B	818	CLA	CHA-CBD-CGD-O2D
22	B	819	CLA	CHA-CBD-CGD-O1D
22	B	819	CLA	CHA-CBD-CGD-O2D
22	B	820	CLA	CHA-CBD-CGD-O1D
22	B	820	CLA	CHA-CBD-CGD-O2D
22	B	821	CLA	CHA-CBD-CGD-O1D
22	B	821	CLA	CHA-CBD-CGD-O2D
22	B	822	CLA	CHA-CBD-CGD-O1D
22	B	822	CLA	CHA-CBD-CGD-O2D
22	B	824	CLA	CHA-CBD-CGD-O2D
22	B	830	CLA	CHA-CBD-CGD-O1D
22	B	830	CLA	CHA-CBD-CGD-O2D
22	B	832	CLA	CHA-CBD-CGD-O1D
22	B	832	CLA	CHA-CBD-CGD-O2D
22	B	833	CLA	CHA-CBD-CGD-O1D
22	B	833	CLA	CHA-CBD-CGD-O2D
22	B	836	CLA	CHA-CBD-CGD-O2D
22	B	838	CLA	CHA-CBD-CGD-O1D
22	B	838	CLA	CHA-CBD-CGD-O2D
22	B	839	CLA	CHA-CBD-CGD-O1D
22	B	839	CLA	CHA-CBD-CGD-O2D
22	B	842	CLA	CHA-CBD-CGD-O1D
22	B	842	CLA	CHA-CBD-CGD-O2D
22	F	305	CLA	CHA-CBD-CGD-O1D
22	F	305	CLA	CHA-CBD-CGD-O2D
22	G	201	CLA	CHA-CBD-CGD-O1D
22	G	201	CLA	CHA-CBD-CGD-O2D
22	J	103	CLA	CHA-CBD-CGD-O1D
22	J	103	CLA	CHA-CBD-CGD-O2D
22	K	203	CLA	CHA-CBD-CGD-O1D
22	K	203	CLA	CHA-CBD-CGD-O2D
22	L	202	CLA	CHA-CBD-CGD-O1D

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Mol	Chain	Res	Type	Atoms
22	L	202	CLA	CHA-CBD-CGD-O2D
22	1	306	CLA	CHA-CBD-CGD-O1D
22	1	306	CLA	CHA-CBD-CGD-O2D
22	1	307	CLA	CHA-CBD-CGD-O1D
22	1	307	CLA	CHA-CBD-CGD-O2D
22	1	310	CLA	CHA-CBD-CGD-O2D
22	1	316	CLA	CHA-CBD-CGD-O2D
22	Z	304	CLA	CHA-CBD-CGD-O2D
22	Z	306	CLA	CHA-CBD-CGD-O1D
22	Z	306	CLA	CHA-CBD-CGD-O2D
22	Z	308	CLA	CHA-CBD-CGD-O1D
22	Z	308	CLA	CHA-CBD-CGD-O2D
22	Z	310	CLA	CHA-CBD-CGD-O1D
22	Z	310	CLA	CHA-CBD-CGD-O2D
22	Z	314	CLA	CHA-CBD-CGD-O1D
22	Z	314	CLA	CHA-CBD-CGD-O2D
22	Z	316	CLA	CHA-CBD-CGD-O1D
22	Z	316	CLA	CHA-CBD-CGD-O2D
22	3	308	CLA	CHA-CBD-CGD-O1D
22	3	308	CLA	CHA-CBD-CGD-O2D
22	3	309	CLA	CHA-CBD-CGD-O2D
22	3	311	CLA	CHA-CBD-CGD-O1D
22	3	311	CLA	CHA-CBD-CGD-O2D
22	3	312	CLA	CHA-CBD-CGD-O2D
22	3	318	CLA	CHA-CBD-CGD-O2D
22	3	319	CLA	CHA-CBD-CGD-O1D
22	3	319	CLA	CHA-CBD-CGD-O2D
22	7	304	CLA	CHA-CBD-CGD-O2D
22	7	306	CLA	CHA-CBD-CGD-O1D
22	7	306	CLA	CHA-CBD-CGD-O2D
22	7	307	CLA	CHA-CBD-CGD-O1D
22	7	307	CLA	CHA-CBD-CGD-O2D
22	7	308	CLA	CHA-CBD-CGD-O1D
22	7	308	CLA	CHA-CBD-CGD-O2D
22	7	309	CLA	CHA-CBD-CGD-O1D
22	7	309	CLA	CHA-CBD-CGD-O2D
22	7	311	CLA	CHA-CBD-CGD-O1D
22	7	311	CLA	CHA-CBD-CGD-O2D
22	7	314	CLA	CHA-CBD-CGD-O2D
22	7	316	CLA	CHA-CBD-CGD-O2D
22	7	317	CLA	CHA-CBD-CGD-O1D
22	7	317	CLA	CHA-CBD-CGD-O2D

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Mol	Chain	Res	Type	Atoms
22	7	322	CLA	CHA-CBD-CGD-O1D
22	7	322	CLA	CHA-CBD-CGD-O2D
22	7	324	CLA	CHA-CBD-CGD-O1D
22	7	324	CLA	CHA-CBD-CGD-O2D
22	8	305	CLA	CHA-CBD-CGD-O1D
22	8	305	CLA	CHA-CBD-CGD-O2D
22	8	310	CLA	CHA-CBD-CGD-O1D
22	8	310	CLA	CHA-CBD-CGD-O2D
22	8	313	CLA	CHA-CBD-CGD-O1D
22	8	313	CLA	CHA-CBD-CGD-O2D
22	8	314	CLA	CHA-CBD-CGD-O1D
22	8	314	CLA	CHA-CBD-CGD-O2D
22	8	316	CLA	CHA-CBD-CGD-O1D
22	8	316	CLA	CHA-CBD-CGD-O2D
22	4	808	CLA	CHA-CBD-CGD-O1D
22	4	810	CLA	CHA-CBD-CGD-O1D
22	4	810	CLA	CHA-CBD-CGD-O2D
22	4	818	CLA	CHA-CBD-CGD-O1D
22	4	818	CLA	CHA-CBD-CGD-O2D
22	5	309	CLA	CHA-CBD-CGD-O1D
22	5	309	CLA	CHA-CBD-CGD-O2D
22	5	310	CLA	CHA-CBD-CGD-O1D
22	5	310	CLA	CHA-CBD-CGD-O2D
22	5	311	CLA	CHA-CBD-CGD-O2D
22	5	312	CLA	CHA-CBD-CGD-O1D
22	5	312	CLA	CHA-CBD-CGD-O2D
22	5	318	CLA	CHA-CBD-CGD-O1D
22	5	318	CLA	CHA-CBD-CGD-O2D
22	5	319	CLA	CHA-CBD-CGD-O1D
22	5	319	CLA	CHA-CBD-CGD-O2D
22	5	320	CLA	CHA-CBD-CGD-O1D
22	5	320	CLA	CHA-CBD-CGD-O2D
22	6	301	CLA	CHA-CBD-CGD-O1D
22	6	301	CLA	CHA-CBD-CGD-O2D
22	6	310	CLA	CHA-CBD-CGD-O1D
22	6	310	CLA	CHA-CBD-CGD-O2D
22	6	311	CLA	CHA-CBD-CGD-O1D
22	6	311	CLA	CHA-CBD-CGD-O2D
22	6	321	CLA	CHA-CBD-CGD-O1D
22	6	321	CLA	CHA-CBD-CGD-O2D
22	9	305	CLA	CHA-CBD-CGD-O2D
22	9	306	CLA	CHA-CBD-CGD-O1D

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Mol	Chain	Res	Type	Atoms
22	9	306	CLA	CHA-CBD-CGD-O2D
22	9	307	CLA	CHA-CBD-CGD-O2D
22	2	304	CLA	CHA-CBD-CGD-O1D
22	2	304	CLA	CHA-CBD-CGD-O2D
39	Z	311	CHL	CHA-CBD-CGD-O2D
39	8	301	CHL	CHA-CBD-CGD-O2D
22	4	805	CLA	CAA-CBA-CGA-O2A
25	7	318	LHG	C25-C26-C27-C28
22	A	835	CLA	C3-C5-C6-C7
41	5	325	3PH	O11-C1-C2-C3
25	A	850	LHG	C12-C13-C14-C15
22	A	825	CLA	C16-C17-C18-C20
22	5	307	CLA	C11-C12-C13-C15
22	A	833	CLA	C8-C10-C11-C12
22	A	811	CLA	CAA-CBA-CGA-O2A
22	A	820	CLA	CAA-CBA-CGA-O2A
22	B	825	CLA	CAA-CBA-CGA-O2A
22	K	203	CLA	CAA-CBA-CGA-O2A
22	7	312	CLA	CAA-CBA-CGA-O2A
22	8	316	CLA	CAA-CBA-CGA-O2A
22	4	818	CLA	CAA-CBA-CGA-O2A
25	Z	317	LHG	O8-C23-C24-C25
25	9	315	LHG	O7-C7-C8-C9
25	9	315	LHG	O8-C23-C24-C25
34	F	308	LMG	O7-C10-C11-C12
35	J	102	T7X	O18-C11-C31-C32
39	4	814	CHL	CAA-CBA-CGA-O2A
35	J	102	T7X	O16-C8-C9-O18
39	8	301	CHL	O1A-CGA-O2A-C1
22	7	304	CLA	C8-C10-C11-C12
22	5	310	CLA	C8-C10-C11-C12
22	A	813	CLA	CAA-CBA-CGA-O2A
22	A	855	CLA	CAA-CBA-CGA-O2A
22	B	814	CLA	CAA-CBA-CGA-O2A
22	B	817	CLA	CAA-CBA-CGA-O2A
22	B	823	CLA	CAA-CBA-CGA-O2A
22	Z	305	CLA	CAA-CBA-CGA-O2A
22	6	317	CLA	CAA-CBA-CGA-O2A
22	7	314	CLA	O1D-CGD-O2D-CED
22	B	820	CLA	C2A-CAA-CBA-CGA
22	A	821	CLA	C16-C17-C18-C20
23	B	843	PQN	C26-C27-C28-C29

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Mol	Chain	Res	Type	Atoms
22	A	810	CLA	CAA-CBA-CGA-O2A
22	B	818	CLA	CAA-CBA-CGA-O2A
34	F	308	LMG	C13-C14-C15-C16
22	A	803	CLA	C11-C10-C8-C7
22	A	827	CLA	C11-C10-C8-C7
22	B	805	CLA	C11-C12-C13-C15
22	B	827	CLA	C11-C10-C8-C7
22	B	833	CLA	C11-C12-C13-C15
22	3	310	CLA	C11-C10-C8-C7
22	3	313	CLA	C11-C10-C8-C7
22	5	310	CLA	C6-C7-C8-C10
22	5	318	CLA	C11-C12-C13-C15
25	A	849	LHG	O9-C7-O7-C5
39	5	316	CHL	C8-C10-C11-C12
22	6	314	CLA	CAA-CBA-CGA-O2A
22	A	803	CLA	C11-C10-C8-C9
22	A	824	CLA	C11-C10-C8-C9
22	A	825	CLA	C11-C10-C8-C9
22	A	829	CLA	C6-C7-C8-C9
22	A	829	CLA	C11-C12-C13-C14
22	A	840	CLA	C11-C12-C13-C14
22	Z	308	CLA	C11-C10-C8-C9
22	6	309	CLA	C11-C10-C8-C9
36	J	105	C7Z	C33-C34-C35-C15
42	7	320	SPH	C6-C7-C8-C9
22	6	322	CLA	CBA-CGA-O2A-C1
22	A	827	CLA	CAA-CBA-CGA-O2A
25	8	317	LHG	O7-C7-C8-C9
22	8	308	CLA	O1D-CGD-O2D-CED
22	Z	309	CLA	CBD-CGD-O2D-CED
22	8	308	CLA	CBD-CGD-O2D-CED
22	9	309	CLA	O1D-CGD-O2D-CED
22	1	310	CLA	C4C-C3C-CAC-CBC
22	B	830	CLA	C4-C3-C5-C6
25	1	317	LHG	C24-C25-C26-C27
22	A	807	CLA	CAA-CBA-CGA-O1A
22	A	812	CLA	CAA-CBA-CGA-O1A
38	7	301	LUT	C27-C28-C29-C30
27	A	852	DGA	CDA-CEA-CFA-CGA
22	B	806	CLA	C1A-C2A-CAA-CBA
22	G	201	CLA	C1A-C2A-CAA-CBA
22	4	806	CLA	C1A-C2A-CAA-CBA

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Mol	Chain	Res	Type	Atoms
22	5	312	CLA	C1A-C2A-CAA-CBA
39	9	312	CHL	C1A-C2A-CAA-CBA
22	A	815	CLA	CAA-CBA-CGA-O1A
22	A	837	CLA	CAA-CBA-CGA-O1A
22	Z	306	CLA	C5-C6-C7-C8
25	Z	317	LHG	C29-C30-C31-C32
22	1	306	CLA	CAA-CBA-CGA-O1A
22	3	322	CLA	CAA-CBA-CGA-O1A
22	9	305	CLA	CAA-CBA-CGA-O1A
27	A	852	DGA	OG1-CG1-CG2-CG3
22	B	828	CLA	CAA-CBA-CGA-O2A
22	G	201	CLA	CAA-CBA-CGA-O2A
22	5	308	CLA	CAA-CBA-CGA-O2A
39	Z	311	CHL	CAA-CBA-CGA-O2A
27	A	852	DGA	CB4-CB5-CB6-CB7
22	A	834	CLA	C2A-CAA-CBA-CGA
22	6	301	CLA	C2A-CAA-CBA-CGA
25	6	323	LHG	C3-O3-P-O6
22	6	310	CLA	C16-C17-C18-C19
39	6	315	CHL	C6-C7-C8-C10
22	8	309	CLA	CAA-CBA-CGA-O1A
25	A	849	LHG	O9-C7-C8-C9
25	1	317	LHG	O10-C23-C24-C25
25	9	315	LHG	O10-C23-C24-C25
22	A	811	CLA	C13-C15-C16-C17
22	4	808	CLA	C10-C11-C12-C13
22	Z	306	CLA	CAA-CBA-CGA-O2A
39	8	312	CHL	CAA-CBA-CGA-O2A
39	4	813	CHL	CAA-CBA-CGA-O2A
39	5	317	CHL	CAA-CBA-CGA-O2A
39	6	316	CHL	CAA-CBA-CGA-O2A
39	2	303	CHL	CAA-CBA-CGA-O2A
22	Z	303	CLA	C10-C11-C12-C13
25	4	820	LHG	C5-C4-O6-P
22	B	824	CLA	CAA-CBA-CGA-O1A
22	3	308	CLA	CAA-CBA-CGA-O1A
22	7	324	CLA	CAA-CBA-CGA-O1A
22	4	806	CLA	CAA-CBA-CGA-O1A
22	5	323	CLA	CAA-CBA-CGA-O1A
31	1	319	DGD	C2E-C1E-O5D-C6D
22	A	814	CLA	C5-C6-C7-C8
22	A	824	CLA	C5-C6-C7-C8

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Mol	Chain	Res	Type	Atoms
25	B	850	LHG	C4-O6-P-O5
25	B	851	LHG	C3-O3-P-O4
25	Z	317	LHG	C4-O6-P-O5
25	8	317	LHG	C4-O6-P-O5
25	4	820	LHG	C3-O3-P-O5
25	5	324	LHG	C4-O6-P-O5
43	8	319	LPX	C1-O2-P1-O4
22	B	814	CLA	C11-C12-C13-C14
22	A	811	CLA	CAA-CBA-CGA-O1A
22	K	203	CLA	CAA-CBA-CGA-O1A
22	8	316	CLA	CAA-CBA-CGA-O1A
22	4	805	CLA	CAA-CBA-CGA-O1A
31	3	301	DGD	O1A-C1A-C2A-C3A
22	B	822	CLA	CAA-CBA-CGA-O2A
22	B	837	CLA	CAA-CBA-CGA-O2A
22	1	310	CLA	C10-C11-C12-C13
24	B	845	BCR	C23-C24-C25-C26
24	I	4001	BCR	C1-C6-C7-C8
22	A	820	CLA	CAA-CBA-CGA-O1A
22	B	812	CLA	CAA-CBA-CGA-O1A
22	B	814	CLA	CAA-CBA-CGA-O1A
22	3	314	CLA	CAA-CBA-CGA-O1A
25	9	315	LHG	O9-C7-C8-C9
39	4	814	CHL	CAA-CBA-CGA-O1A
25	Z	317	LHG	C11-C10-C9-C8
22	Z	303	CLA	CAA-CBA-CGA-O2A
22	9	308	CLA	CAA-CBA-CGA-O2A
26	8	318	NKP	OAJ-CAK-CAL-CAM
22	1	307	CLA	C11-C12-C13-C14
22	7	312	CLA	C11-C12-C13-C15
22	A	810	CLA	CAA-CBA-CGA-O1A
22	A	855	CLA	CAA-CBA-CGA-O1A
25	Z	317	LHG	O10-C23-C24-C25
34	F	308	LMG	O9-C10-C11-C12
35	J	102	T7X	O19-C11-C31-C32
22	A	824	CLA	C15-C16-C17-C18
22	A	834	CLA	C13-C15-C16-C17
21	A	801	CL0	CAA-CBA-CGA-O1A
22	A	842	CLA	CAA-CBA-CGA-O1A
22	Z	305	CLA	CAA-CBA-CGA-O1A
22	5	309	CLA	CAA-CBA-CGA-O1A
24	B	846	BCR	C9-C10-C11-C12

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Mol	Chain	Res	Type	Atoms
24	B	853	BCR	C9-C10-C11-C12
22	7	324	CLA	C2-C3-C5-C6
23	B	843	PQN	C20-C21-C22-C23
22	A	807	CLA	CAD-CBD-CGD-O1D
22	A	811	CLA	CAD-CBD-CGD-O1D
22	A	817	CLA	CAD-CBD-CGD-O1D
22	A	832	CLA	CAD-CBD-CGD-O1D
22	B	824	CLA	CAD-CBD-CGD-O1D
22	B	836	CLA	CAD-CBD-CGD-O1D
22	B	837	CLA	C2-C3-C5-C6
22	K	204	CLA	CAD-CBD-CGD-O1D
22	1	310	CLA	CAD-CBD-CGD-O1D
22	1	316	CLA	CAD-CBD-CGD-O1D
22	3	309	CLA	CAD-CBD-CGD-O1D
22	3	312	CLA	CAD-CBD-CGD-O1D
22	7	304	CLA	CAD-CBD-CGD-O1D
22	7	314	CLA	CAD-CBD-CGD-O1D
22	7	316	CLA	CAD-CBD-CGD-O1D
22	8	306	CLA	CAD-CBD-CGD-O1D
22	8	308	CLA	CAD-CBD-CGD-O1D
22	4	809	CLA	CAD-CBD-CGD-O1D
22	4	811	CLA	CAD-CBD-CGD-O1D
22	5	301	CLA	CAD-CBD-CGD-O1D
22	5	320	CLA	CAD-CBD-CGD-O1D
41	7	319	3PH	C21-C22-C23-C24
22	A	827	CLA	CAA-CBA-CGA-O1A
22	9	306	CLA	CAA-CBA-CGA-O1A
22	A	814	CLA	CAA-CBA-CGA-O2A
22	K	204	CLA	CAA-CBA-CGA-O2A
22	3	316	CLA	CAA-CBA-CGA-O2A
22	A	810	CLA	C11-C12-C13-C14
22	A	811	CLA	C11-C10-C8-C9
22	A	827	CLA	C11-C10-C8-C9
22	B	801	CLA	C14-C13-C15-C16
22	B	808	CLA	C11-C10-C8-C9
22	B	809	CLA	C11-C10-C8-C9
22	B	827	CLA	C11-C10-C8-C9
22	B	833	CLA	C11-C12-C13-C14
22	1	306	CLA	C6-C7-C8-C9
22	3	310	CLA	C11-C10-C8-C9
22	5	318	CLA	C11-C12-C13-C14
23	A	843	PQN	C16-C17-C18-C19

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Mol	Chain	Res	Type	Atoms
22	A	832	CLA	C15-C16-C17-C18
22	B	814	CLA	C5-C6-C7-C8
22	B	817	CLA	CAA-CBA-CGA-O1A
22	4	818	CLA	CAA-CBA-CGA-O1A
22	6	317	CLA	CAA-CBA-CGA-O1A
22	A	823	CLA	CAA-CBA-CGA-O2A
22	A	854	CLA	CAA-CBA-CGA-O2A
22	B	831	CLA	CAA-CBA-CGA-O2A
22	F	304	CLA	CAA-CBA-CGA-O2A
22	Z	315	CLA	CAA-CBA-CGA-O2A
22	8	306	CLA	CAA-CBA-CGA-O2A
22	5	310	CLA	CAA-CBA-CGA-O2A
25	6	323	LHG	O7-C7-C8-C9
27	A	852	DGA	OG1-CA1-CA2-CA3
37	K	201	DAO	C7-C8-C9-C10
22	8	314	CLA	C2A-CAA-CBA-CGA
22	A	829	CLA	CAA-CBA-CGA-O2A
22	A	831	CLA	CAA-CBA-CGA-O2A
22	B	808	CLA	CAA-CBA-CGA-O2A
22	B	839	CLA	CAA-CBA-CGA-O2A
22	1	315	CLA	CAA-CBA-CGA-O2A
22	Z	314	CLA	CAA-CBA-CGA-O2A
22	5	318	CLA	CAA-CBA-CGA-O2A
25	4	821	LHG	O7-C7-C8-C9
39	5	316	CHL	CAA-CBA-CGA-O2A
22	L	201	CLA	C8-C10-C11-C12
22	A	813	CLA	CAA-CBA-CGA-O1A
22	6	314	CLA	CAA-CBA-CGA-O1A
22	A	825	CLA	C4-C3-C5-C6
22	3	322	CLA	C4-C3-C5-C6
21	A	801	CL0	C11-C10-C8-C7
22	A	803	CLA	C11-C12-C13-C15
22	A	810	CLA	C11-C10-C8-C7
22	A	812	CLA	C11-C12-C13-C15
22	A	820	CLA	C3A-C2A-CAA-CBA
22	A	824	CLA	C3A-C2A-CAA-CBA
22	A	829	CLA	C6-C7-C8-C10
22	A	836	CLA	C3A-C2A-CAA-CBA
22	A	841	CLA	C3A-C2A-CAA-CBA
22	B	808	CLA	C11-C10-C8-C7
22	B	818	CLA	C11-C10-C8-C7
22	B	825	CLA	C2-C3-C5-C6

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Mol	Chain	Res	Type	Atoms
22	B	826	CLA	C12-C13-C15-C16
22	B	829	CLA	C11-C10-C8-C7
22	B	830	CLA	C6-C7-C8-C10
22	B	831	CLA	C3A-C2A-CAA-CBA
22	F	304	CLA	C6-C7-C8-C10
22	L	201	CLA	C11-C12-C13-C15
22	7	310	CLA	C3A-C2A-CAA-CBA
22	8	308	CLA	C11-C12-C13-C15
22	6	312	CLA	C11-C12-C13-C15
22	9	305	CLA	C3A-C2A-CAA-CBA
22	9	305	CLA	C6-C7-C8-C10
39	8	315	CHL	C6-C7-C8-C10
22	B	823	CLA	CAA-CBA-CGA-O1A
22	F	304	CLA	CAA-CBA-CGA-O1A
25	8	317	LHG	O9-C7-C8-C9
39	Z	311	CHL	CAA-CBA-CGA-O1A
39	6	316	CHL	CAA-CBA-CGA-O1A
39	2	303	CHL	CAA-CBA-CGA-O1A
22	K	202	CLA	CAA-CBA-CGA-O2A
22	Z	309	CLA	CAA-CBA-CGA-O2A
22	3	313	CLA	CAA-CBA-CGA-O2A
22	6	312	CLA	CAA-CBA-CGA-O2A
22	6	319	CLA	CAA-CBA-CGA-O2A
25	A	849	LHG	O8-C23-C24-C25
39	1	312	CHL	CAA-CBA-CGA-O2A
39	Z	312	CHL	CAA-CBA-CGA-O2A
31	B	852	DGD	C5B-C6B-C7B-C8B
38	4	802	LUT	C27-C28-C29-C30
38	9	301	LUT	C7-C8-C9-C10
22	A	831	CLA	CAA-CBA-CGA-O1A
22	G	201	CLA	CAA-CBA-CGA-O1A
25	A	849	LHG	O10-C23-C24-C25
25	4	821	LHG	O9-C7-C8-C9
39	8	312	CHL	CAA-CBA-CGA-O1A
38	4	803	LUT	C13-C14-C15-C35
39	6	315	CHL	C6-C7-C8-C9
22	A	833	CLA	CAA-CBA-CGA-O2A
22	B	838	CLA	CAA-CBA-CGA-O2A
22	1	307	CLA	CAA-CBA-CGA-O2A
22	Z	313	CLA	CAA-CBA-CGA-O2A
22	Z	316	CLA	CAA-CBA-CGA-O2A
22	B	806	CLA	C5-C6-C7-C8

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Mol	Chain	Res	Type	Atoms
22	B	822	CLA	CAA-CBA-CGA-O1A
22	B	825	CLA	CAA-CBA-CGA-O1A
22	B	831	CLA	CAA-CBA-CGA-O1A
22	B	837	CLA	CAA-CBA-CGA-O1A
22	B	839	CLA	CAA-CBA-CGA-O1A
22	K	204	CLA	CAA-CBA-CGA-O1A
22	Z	306	CLA	CAA-CBA-CGA-O1A
22	Z	315	CLA	CAA-CBA-CGA-O1A
22	5	308	CLA	CAA-CBA-CGA-O1A
22	A	817	CLA	CAA-CBA-CGA-O2A
22	B	821	CLA	CAA-CBA-CGA-O2A
22	L	201	CLA	CAA-CBA-CGA-O2A
22	6	308	CLA	CAA-CBA-CGA-O2A
25	B	803	LHG	O8-C23-C24-C25
34	J	101	LMG	O7-C10-C11-C12
22	F	302	CLA	C10-C11-C12-C13
22	A	823	CLA	CAA-CBA-CGA-O1A
22	B	828	CLA	CAA-CBA-CGA-O1A
39	4	813	CHL	CAA-CBA-CGA-O1A
39	5	317	CHL	CAA-CBA-CGA-O1A
22	A	832	CLA	C10-C11-C12-C13
22	1	314	CLA	C13-C15-C16-C17
22	6	309	CLA	C15-C16-C17-C18
22	A	814	CLA	CAA-CBA-CGA-O1A
22	8	306	CLA	CAA-CBA-CGA-O1A
39	5	316	CHL	CAA-CBA-CGA-O1A
22	B	825	CLA	C4-C3-C5-C6
22	1	316	CLA	C4C-C3C-CAC-CBC
22	F	302	CLA	CAA-CBA-CGA-O2A

There are no ring outliers.

328 monomers are involved in 3221 short contacts:

Mol	Chain	Res	Type	Clashes	Symm-Clashes
22	5	315	CLA	20	0
22	3	319	CLA	4	0
22	A	812	CLA	24	0
22	9	308	CLA	8	0
22	G	201	CLA	13	0
42	7	321	SPH	8	0
22	7	314	CLA	5	0
22	6	312	CLA	18	0

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Mol	Chain	Res	Type	Clashes	Symm-Clashes
22	Z	304	CLA	18	0
26	A	851	NKP	3	0
39	8	315	CHL	24	0
39	5	316	CHL	16	0
22	A	831	CLA	4	0
22	5	311	CLA	20	0
22	A	854	CLA	10	0
22	B	816	CLA	14	0
22	J	103	CLA	4	0
22	3	308	CLA	18	0
24	B	853	BCR	21	0
39	6	318	CHL	14	0
24	A	844	BCR	30	0
25	8	317	LHG	7	0
22	A	817	CLA	14	0
22	B	806	CLA	22	0
38	3	303	LUT	20	0
39	9	314	CHL	13	0
22	A	855	CLA	16	0
22	B	808	CLA	13	0
22	A	814	CLA	22	0
39	4	814	CHL	8	0
22	A	818	CLA	15	0
41	5	325	3PH	4	0
25	4	820	LHG	16	0
22	5	318	CLA	22	0
22	Z	315	CLA	11	0
41	7	319	3PH	12	0
24	3	307	BCR	19	0
22	A	815	CLA	20	0
22	4	805	CLA	18	0
22	A	804	CLA	14	0
22	3	318	CLA	20	0
22	4	807	CLA	12	0
35	J	102	T7X	2	0
22	A	807	CLA	20	0
34	F	308	LMG	1	0
22	7	306	CLA	13	0
24	A	856	BCR	25	0
22	2	302	CLA	7	0
22	7	311	CLA	2	0
22	1	314	CLA	19	0

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Mol	Chain	Res	Type	Clashes	Symm-Clashes
37	K	201	DAO	8	0
24	7	303	BCR	15	0
22	6	308	CLA	17	0
22	B	821	CLA	15	0
22	A	805	CLA	22	0
22	B	810	CLA	17	0
22	A	819	CLA	12	0
22	8	311	CLA	16	0
24	I	4001	BCR	12	0
33	4	822	LMT	2	0
38	4	802	LUT	15	0
22	7	317	CLA	6	0
39	7	313	CHL	12	0
22	A	829	CLA	16	0
22	5	319	CLA	16	0
22	A	803	CLA	8	0
22	B	801	CLA	15	0
22	B	824	CLA	22	0
22	6	301	CLA	20	0
22	5	310	CLA	17	0
39	6	316	CHL	10	0
22	3	310	CLA	7	0
40	1	318	SQD	13	0
39	5	317	CHL	10	0
22	A	810	CLA	22	0
38	1	302	LUT	11	0
24	B	847	BCR	21	0
22	4	806	CLA	7	0
39	3	317	CHL	13	0
39	Z	311	CHL	13	0
38	3	302	LUT	12	0
22	6	317	CLA	14	0
24	K	206	BCR	11	0
22	4	818	CLA	9	0
25	5	324	LHG	7	0
22	1	310	CLA	11	0
25	B	850	LHG	4	0
22	B	841	CLA	15	0
22	5	309	CLA	13	0
22	5	312	CLA	12	0
25	A	850	LHG	9	0
24	4	804	BCR	27	0

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Mol	Chain	Res	Type	Clashes	Symm-Clashes
22	A	841	CLA	17	0
25	1	317	LHG	8	0
22	B	809	CLA	13	0
22	8	313	CLA	12	0
22	4	812	CLA	4	0
27	A	852	DGA	20	0
22	A	824	CLA	15	0
22	B	805	CLA	12	0
33	1	301	LMT	9	0
36	5	306	C7Z	2	0
22	4	815	CLA	16	0
24	B	848	BCR	15	0
22	4	808	CLA	20	0
24	J	104	BCR	19	0
22	A	809	CLA	15	0
22	Z	310	CLA	3	0
22	6	314	CLA	20	0
22	B	829	CLA	13	0
22	B	815	CLA	23	0
22	A	820	CLA	8	0
29	B	802	SF4	9	0
22	5	301	CLA	14	0
22	7	322	CLA	18	0
38	Z	301	LUT	25	0
42	7	320	SPH	4	0
22	A	806	CLA	12	0
22	B	831	CLA	9	0
22	Z	305	CLA	17	0
25	B	803	LHG	8	0
22	A	842	CLA	5	0
22	3	315	CLA	11	0
24	B	844	BCR	20	0
22	1	305	CLA	12	0
24	A	847	BCR	14	0
22	Z	307	CLA	13	0
38	7	301	LUT	16	0
22	4	817	CLA	4	0
31	3	301	DGD	21	0
22	B	838	CLA	16	0
22	F	304	CLA	30	0
39	2	303	CHL	6	0
22	1	304	CLA	17	0

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Mol	Chain	Res	Type	Clashes	Symm-Clashes
22	9	310	CLA	1	0
22	1	313	CLA	16	0
22	A	835	CLA	20	0
22	7	310	CLA	15	0
22	8	305	CLA	24	0
22	Z	316	CLA	15	0
22	L	202	CLA	19	0
22	5	322	CLA	13	0
22	5	326	CLA	20	0
24	L	204	BCR	17	0
22	A	839	CLA	9	0
24	A	848	BCR	8	0
22	5	314	CLA	8	0
22	B	837	CLA	2	0
24	6	306	BCR	16	0
22	L	201	CLA	16	0
39	6	320	CHL	13	0
22	B	842	CLA	27	0
22	L	203	CLA	5	0
22	8	310	CLA	13	0
22	A	822	CLA	24	0
22	B	817	CLA	18	0
22	3	312	CLA	25	0
24	F	303	BCR	16	0
33	4	801	LMT	8	0
22	5	308	CLA	21	0
22	9	306	CLA	16	0
39	6	315	CHL	22	0
22	B	836	CLA	7	0
22	A	823	CLA	25	0
22	6	311	CLA	12	0
25	7	318	LHG	10	0
22	A	828	CLA	17	0
43	8	319	LPX	7	0
22	3	311	CLA	13	0
22	4	811	CLA	7	0
22	K	203	CLA	11	0
22	G	202	CLA	15	0
22	6	321	CLA	9	0
22	A	808	CLA	12	0
24	A	846	BCR	10	0
38	4	803	LUT	20	0

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Mol	Chain	Res	Type	Clashes	Symm-Clashes
22	F	305	CLA	9	0
39	9	312	CHL	24	0
38	5	302	LUT	14	0
22	1	315	CLA	4	0
22	B	823	CLA	16	0
22	A	840	CLA	16	0
22	8	316	CLA	3	0
22	A	825	CLA	20	0
22	F	302	CLA	19	0
22	1	309	CLA	10	0
22	9	303	CLA	7	0
22	4	809	CLA	10	0
22	F	301	CLA	11	0
22	1	307	CLA	22	0
22	B	832	CLA	10	0
22	B	828	CLA	26	0
22	A	827	CLA	20	0
22	B	811	CLA	26	0
22	7	316	CLA	12	0
24	B	845	BCR	9	0
22	7	304	CLA	13	0
31	B	852	DGD	17	0
22	1	316	CLA	5	0
39	1	312	CHL	9	0
34	J	101	LMG	11	0
22	8	307	CLA	19	0
22	A	802	CLA	14	0
22	A	813	CLA	19	0
22	6	319	CLA	12	0
22	B	827	CLA	24	0
24	5	305	BCR	11	0
22	A	816	CLA	22	0
22	B	840	CLA	16	0
39	4	813	CHL	14	0
22	B	835	CLA	11	0
39	4	816	CHL	20	0
22	Z	306	CLA	19	0
22	1	306	CLA	8	0
22	A	836	CLA	15	0
25	6	323	LHG	12	0
25	3	321	LHG	1	0
32	F	306	RRX	19	0

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Mol	Chain	Res	Type	Clashes	Symm-Clashes
22	7	324	CLA	7	0
25	Z	317	LHG	9	0
22	B	813	CLA	9	0
22	5	313	CLA	13	0
22	6	307	CLA	22	0
22	3	316	CLA	10	0
22	B	822	CLA	11	0
38	6	304	LUT	17	0
22	6	309	CLA	22	0
22	4	810	CLA	17	0
22	Z	309	CLA	21	0
22	3	313	CLA	26	0
38	6	303	LUT	12	0
22	7	308	CLA	13	0
22	8	314	CLA	6	0
22	3	309	CLA	10	0
22	7	305	CLA	8	0
25	4	821	LHG	4	0
36	J	105	C7Z	2	0
22	9	313	CLA	19	0
22	5	323	CLA	10	0
39	4	819	CHL	10	0
22	B	833	CLA	11	0
38	1	303	LUT	20	0
22	6	313	CLA	7	0
22	B	819	CLA	18	0
22	8	308	CLA	15	0
22	Z	313	CLA	24	0
22	Z	308	CLA	14	0
22	B	825	CLA	26	0
22	B	818	CLA	13	0
22	B	820	CLA	12	0
22	B	839	CLA	13	0
38	8	303	LUT	13	0
24	5	304	BCR	22	0
29	C	101	SF4	4	0
22	A	833	CLA	17	0
24	3	306	BCR	14	0
22	9	304	CLA	2	0
24	G	203	BCR	13	0
22	Z	303	CLA	32	0
24	3	304	BCR	15	0

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Mol	Chain	Res	Type	Clashes	Symm-Clashes
22	B	807	CLA	9	0
33	G	204	LMT	9	0
41	6	324	3PH	5	0
22	9	307	CLA	11	0
24	A	845	BCR	20	0
22	6	322	CLA	21	0
41	8	320	3PH	4	0
22	K	204	CLA	5	0
22	A	838	CLA	9	0
22	Z	314	CLA	9	0
22	7	312	CLA	6	0
22	B	826	CLA	18	0
22	3	314	CLA	14	0
22	6	302	CLA	17	0
33	F	307	LMT	5	0
22	3	322	CLA	20	0
22	K	202	CLA	8	0
24	8	304	BCR	15	0
23	A	843	PQN	9	0
39	5	321	CHL	6	0
22	B	812	CLA	15	0
24	B	849	BCR	9	0
31	1	319	DGD	10	0
22	7	307	CLA	23	0
22	9	311	CLA	5	0
22	2	304	CLA	9	0
22	A	830	CLA	14	0
22	7	323	CLA	17	0
22	A	834	CLA	13	0
26	8	318	NKP	6	0
38	Z	302	LUT	14	0
39	8	301	CHL	13	0
22	2	301	CLA	19	0
22	8	306	CLA	15	0
23	B	843	PQN	5	0
24	B	846	BCR	12	0
22	7	315	CLA	11	0
22	B	834	CLA	14	0
38	7	302	LUT	17	0
22	8	309	CLA	15	0
22	K	205	CLA	4	0
22	A	832	CLA	13	0

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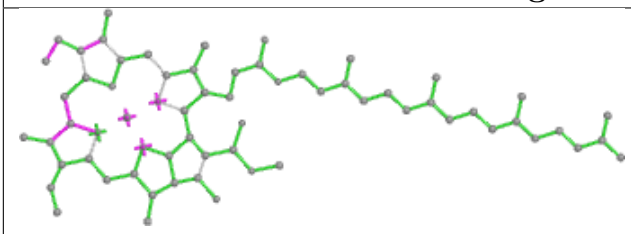
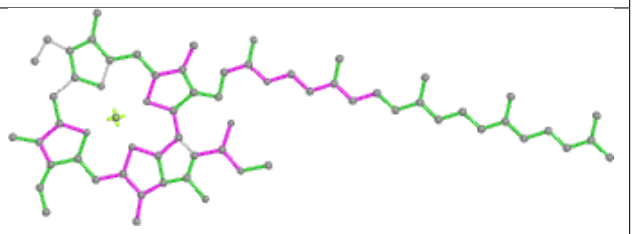
Mol	Chain	Res	Type	Clashes	Symm-Clashes
24	3	305	BCR	15	0
25	9	315	LHG	2	0
22	6	310	CLA	20	0
22	1	308	CLA	23	0
22	7	309	CLA	11	0
22	A	811	CLA	19	0
24	6	305	BCR	19	0
22	B	814	CLA	16	0
22	9	305	CLA	7	0
25	A	849	LHG	5	0
22	3	320	CLA	9	0
38	5	303	LUT	18	0
38	8	302	LUT	18	0
22	5	307	CLA	24	0
39	8	312	CHL	16	0
39	Z	312	CHL	14	0
25	B	851	LHG	2	0
29	C	102	SF4	5	0
22	A	826	CLA	9	0
22	5	320	CLA	12	0
22	A	821	CLA	25	0
22	A	837	CLA	9	0
38	9	301	LUT	17	0
38	9	302	LUT	15	0
22	B	830	CLA	15	0
21	A	801	CL0	9	0

The following is a two-dimensional graphical depiction of Mogul quality analysis of bond lengths, bond angles, torsion angles, and ring geometry for all instances of the Ligand of Interest. In addition, ligands with molecular weight > 250 and outliers as shown on the validation Tables will also be included. For torsion angles, if less then 5% of the Mogul distribution of torsion angles is within 10 degrees of the torsion angle in question, then that torsion angle is considered an outlier. Any bond that is central to one or more torsion angles identified as an outlier by Mogul will be highlighted in the graph. For rings, the root-mean-square deviation (RMSD) between the ring in question and similar rings identified by Mogul is calculated over all ring torsion angles. If the average RMSD is greater than 60 degrees and the minimal RMSD between the ring in question and any Mogul-identified rings is also greater than 60 degrees, then that ring is considered an outlier. The outliers are highlighted in purple. The color gray indicates Mogul did not find sufficient equivalents in the CSD to analyse the geometry.

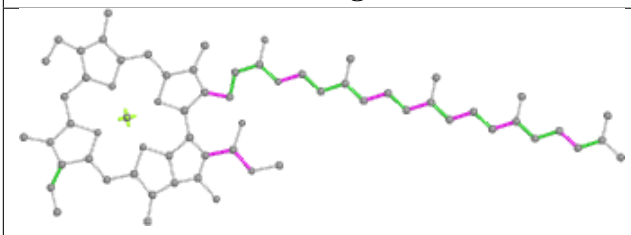
Ligand CLA 5 315



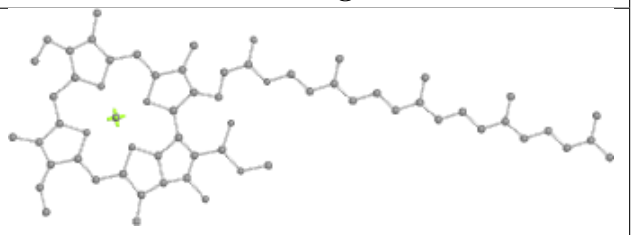
Bond lengths



Bond angles

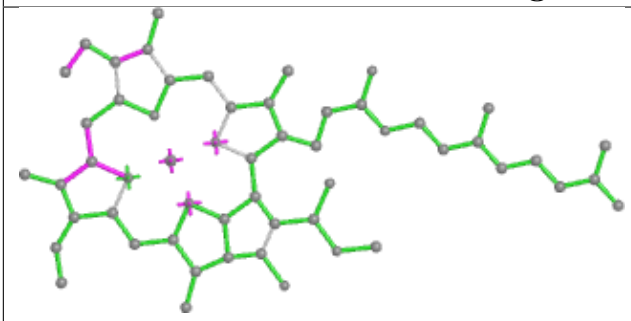


Torsions

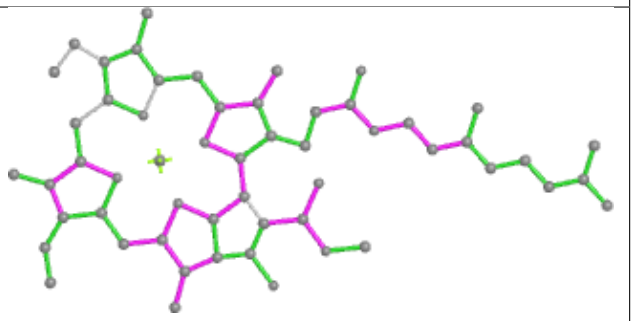


Rings

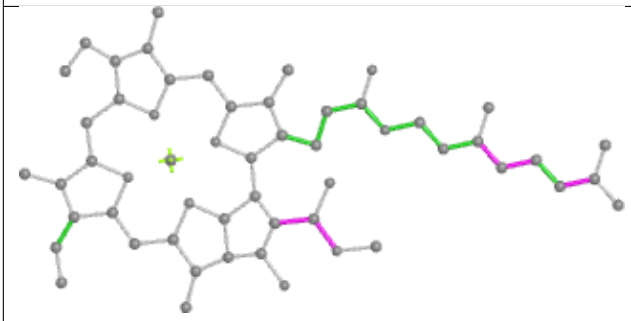
Ligand CLA 3 319



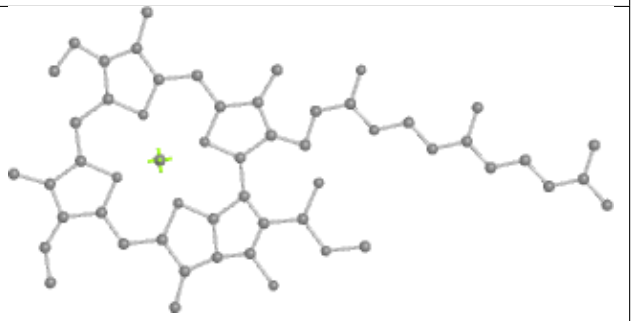
Bond lengths



Bond angles

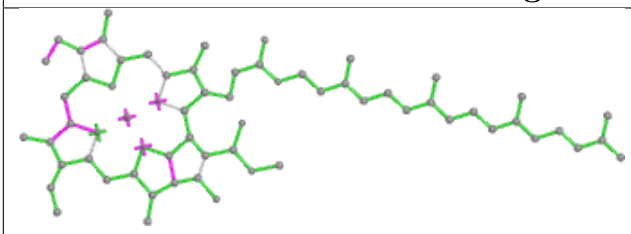


Torsions

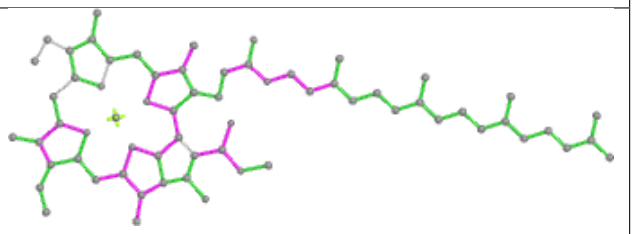


Rings

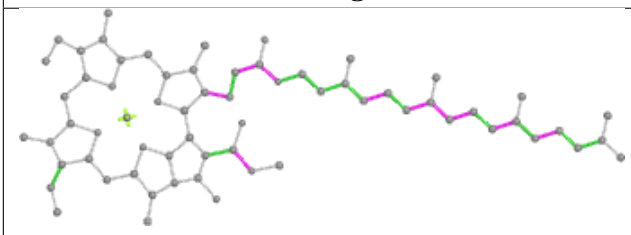
Ligand CLA A 812



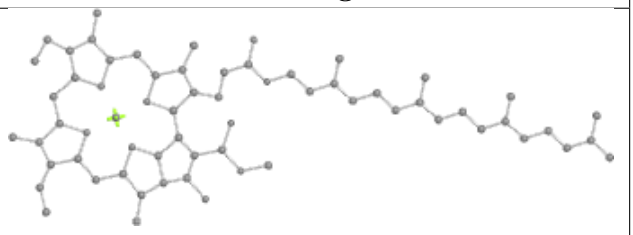
Bond lengths



Bond angles

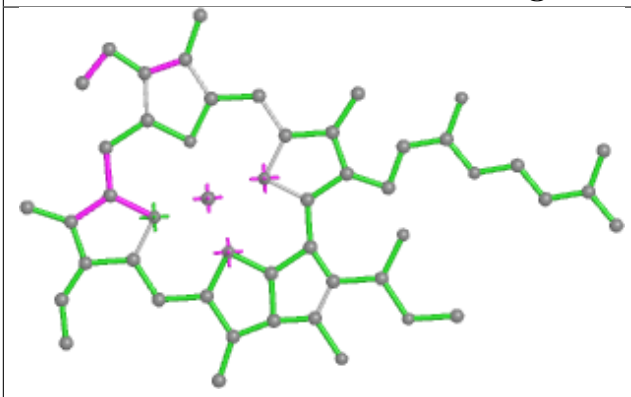


Torsions

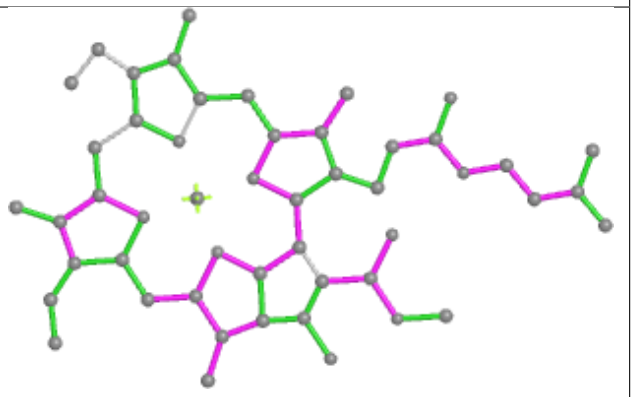


Rings

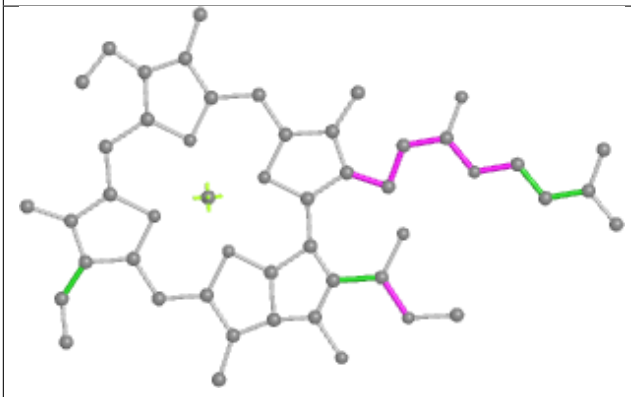
Ligand CLA 9 308



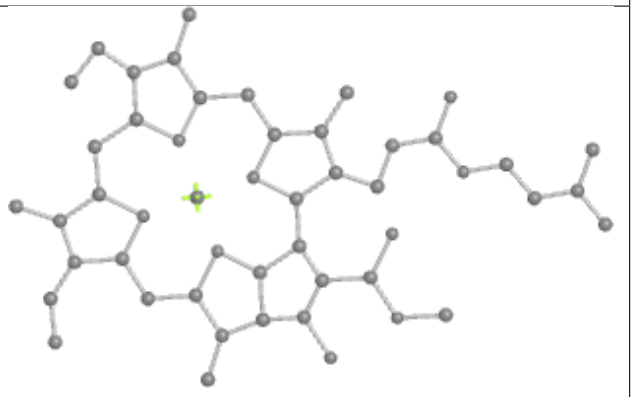
Bond lengths



Bond angles

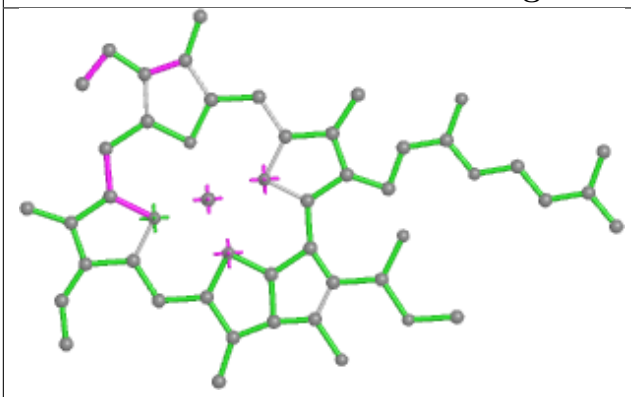


Torsions



Rings

Ligand CLA G 201



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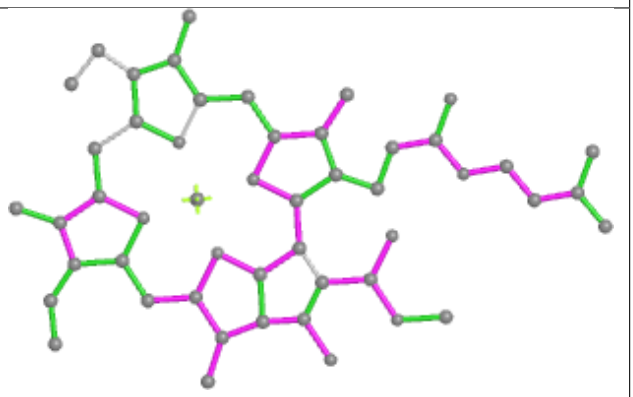

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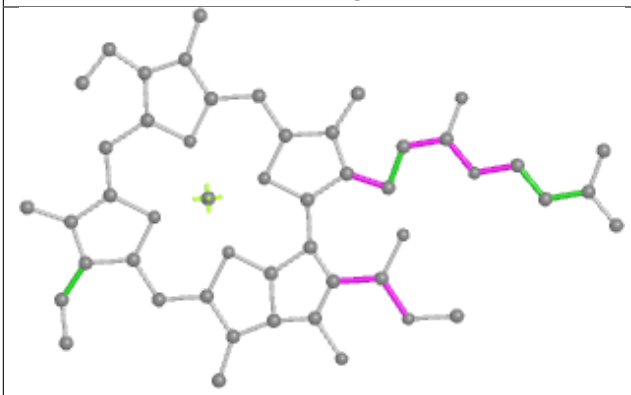

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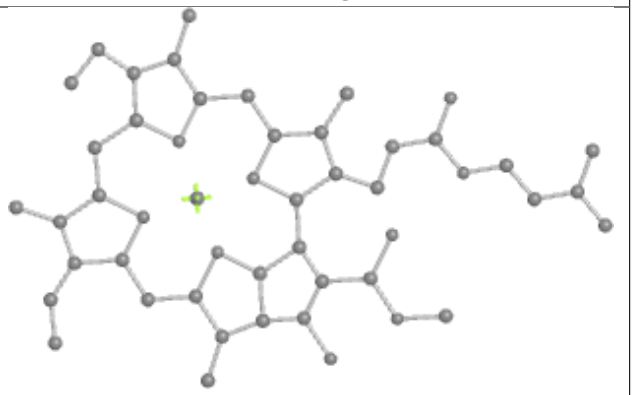
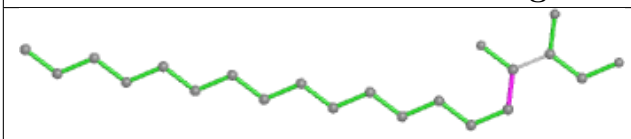
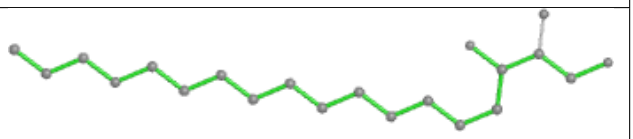


Rings

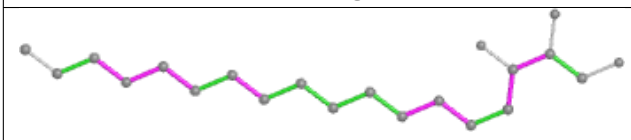
Ligand SPH 7 321



Bond lengths



Bond angles



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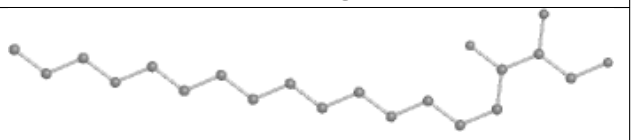

Rings

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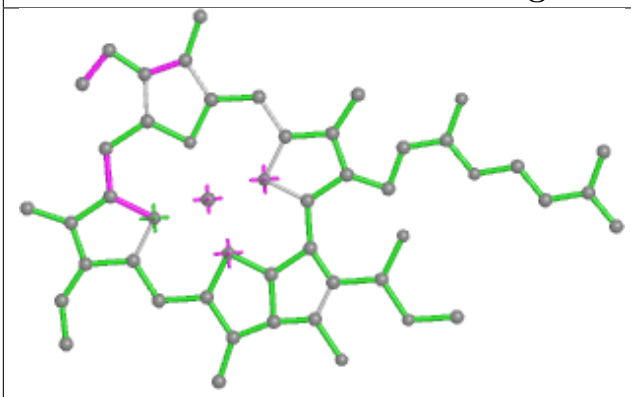
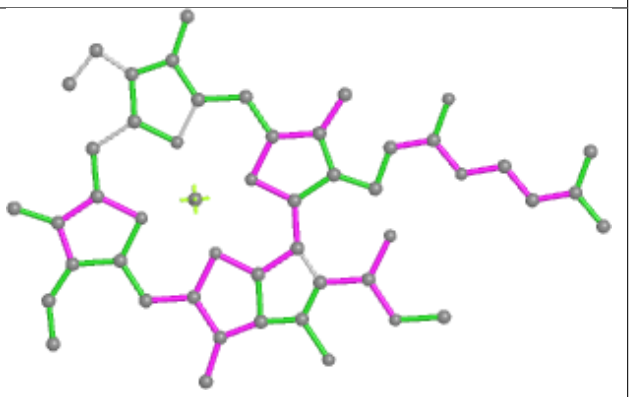
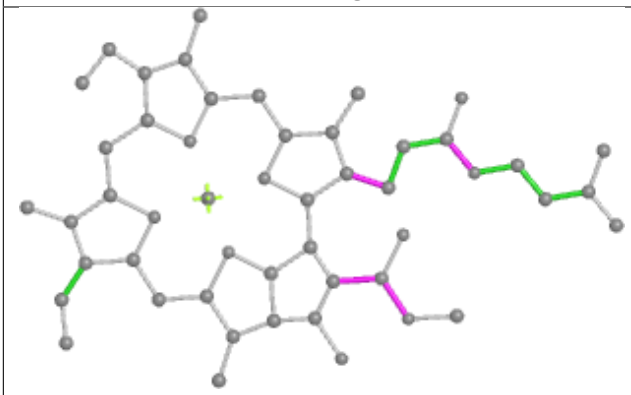
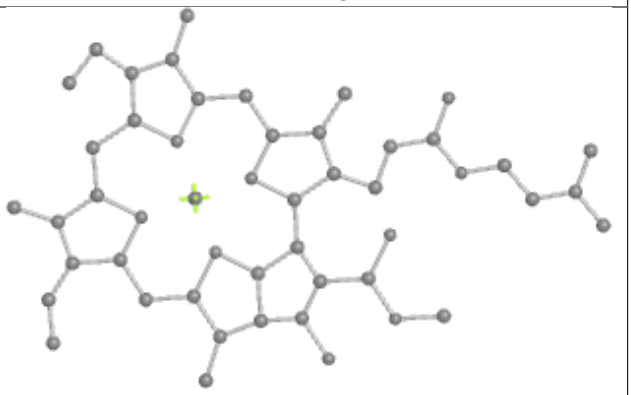
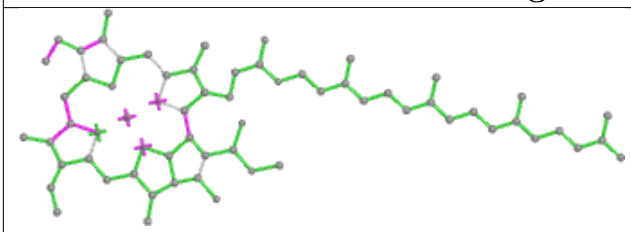
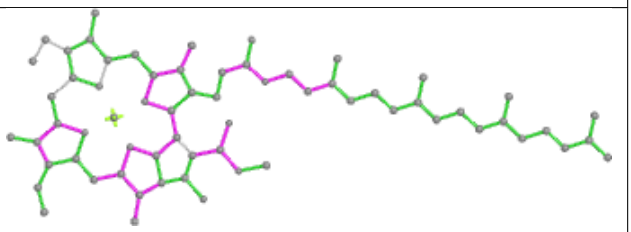
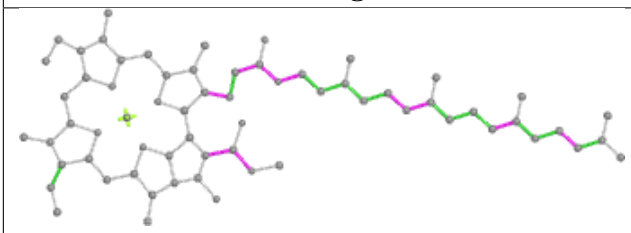
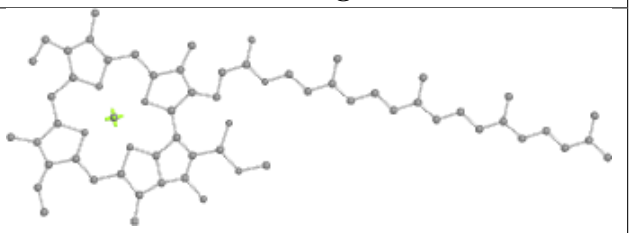
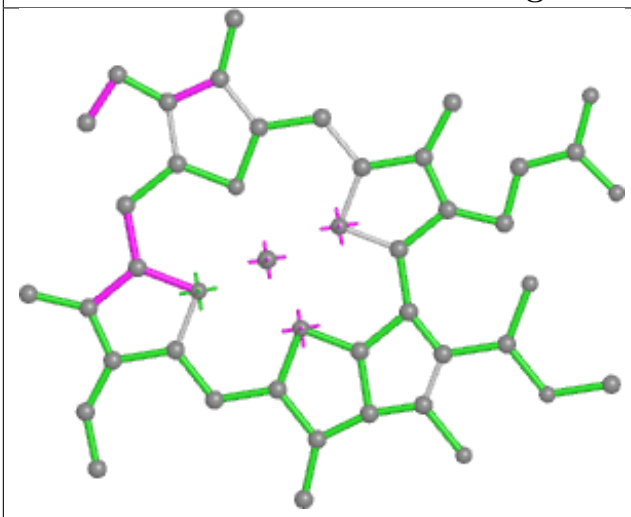
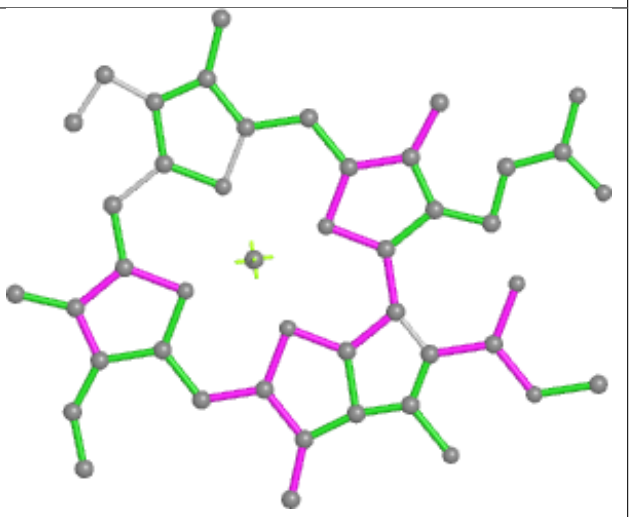
	
Bond lengths	Bond angles
	
Torsions	Rings

Ligand CLA 6 312	
	
Bond lengths	Bond angles
	
Torsions	Rings

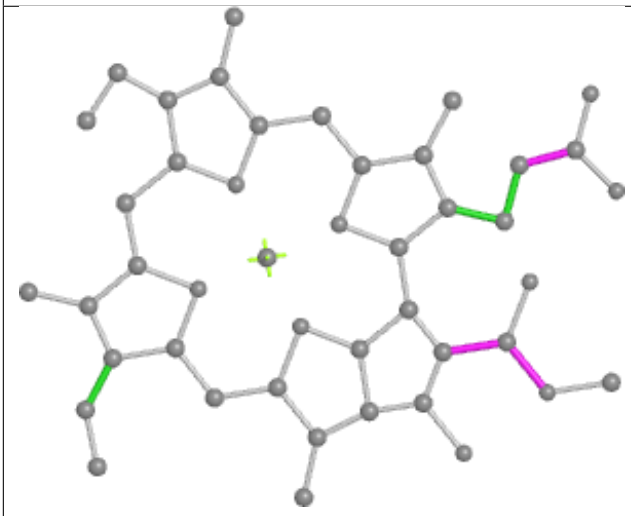
Ligand CLA Z 304



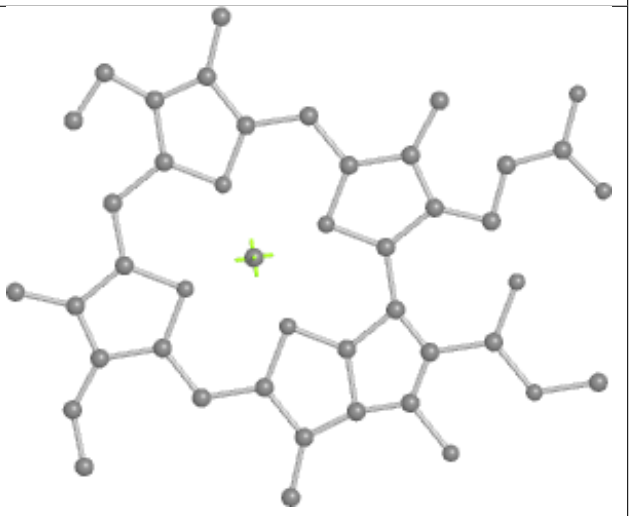
Bond lengths



Bond angles

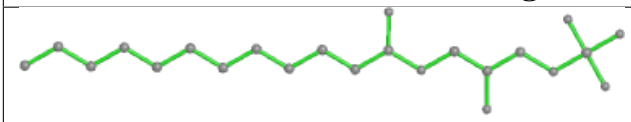


Torsions

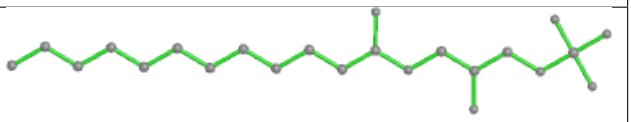


Rings

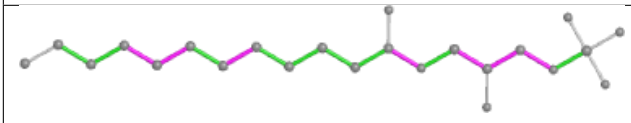
Ligand NKP A 851



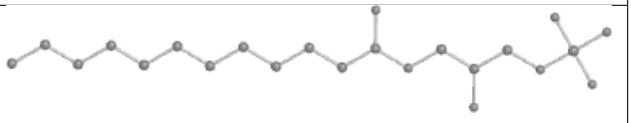
Bond lengths



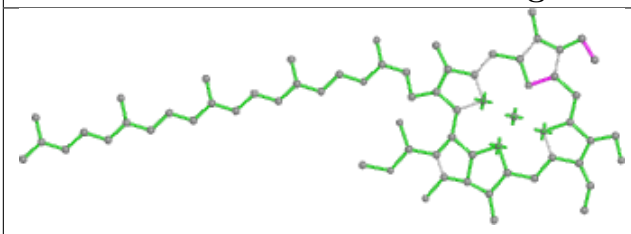
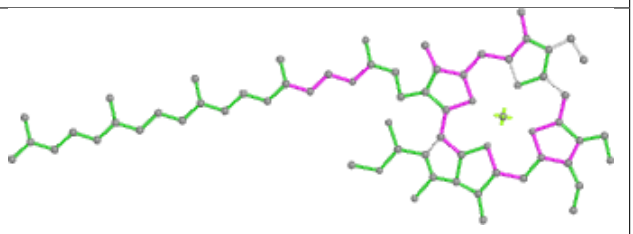
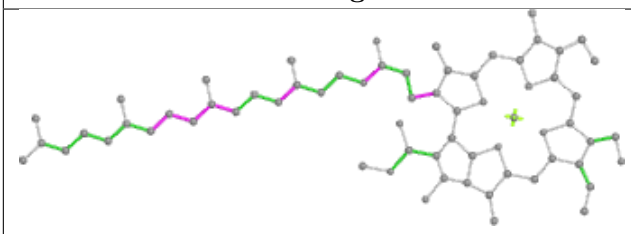
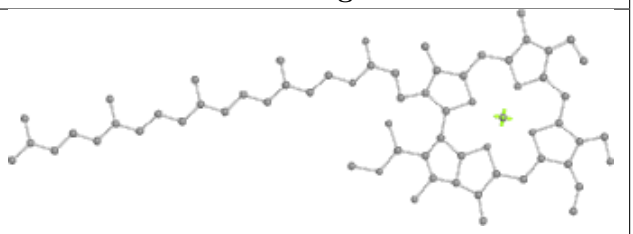
Bond angles

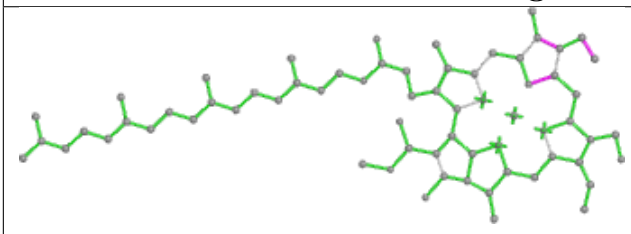
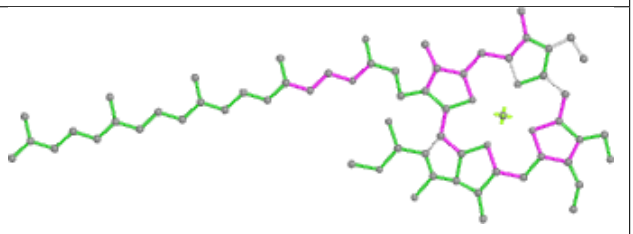
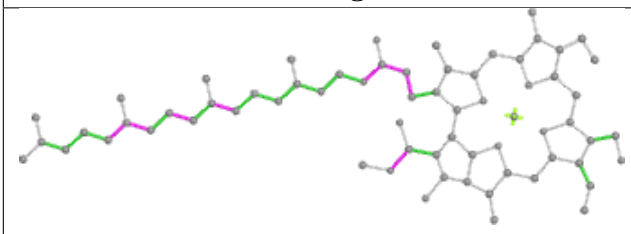
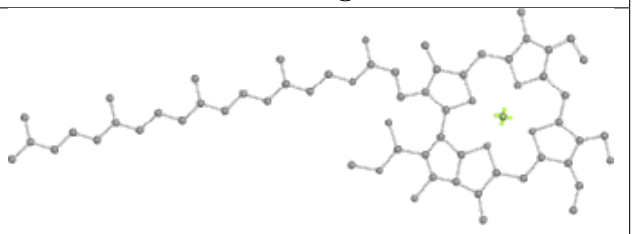


Torsions

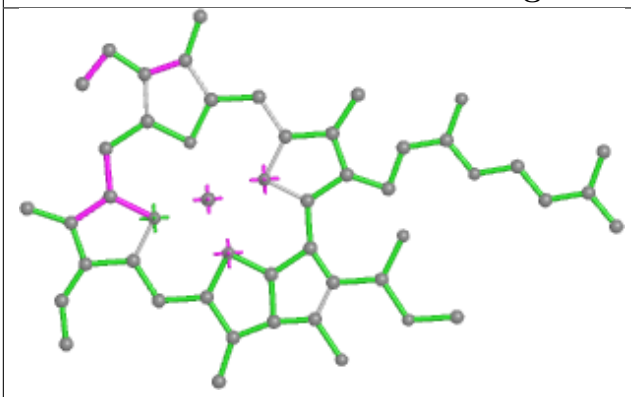


Rings

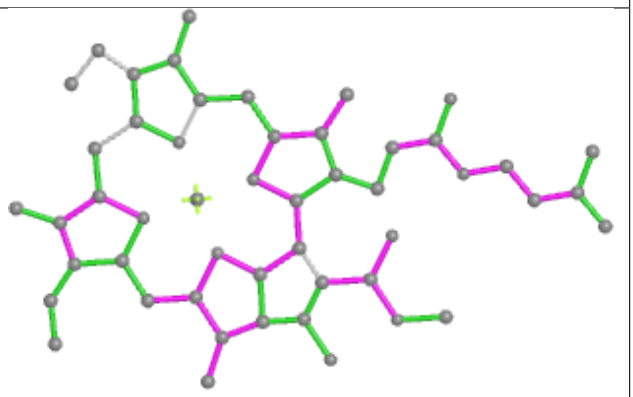
Ligand CHL 8 315	
 Bond lengths	 Bond angles
 Torsions	 Rings

Ligand CHL 5 316	
 Bond lengths	 Bond angles
 Torsions	 Rings

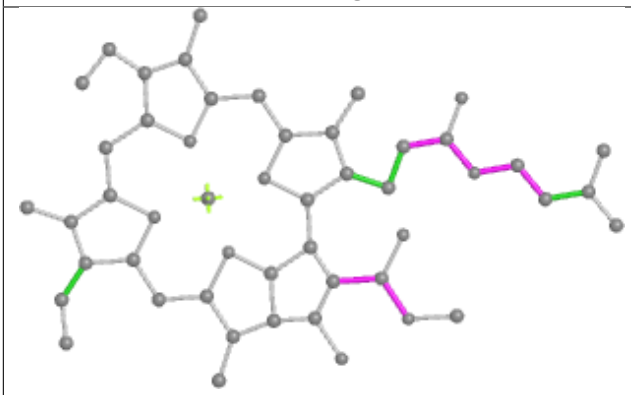
Ligand CLA A 831



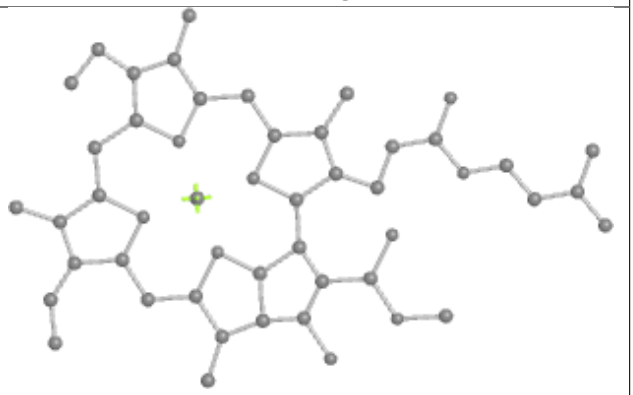
Bond lengths



Bond angles

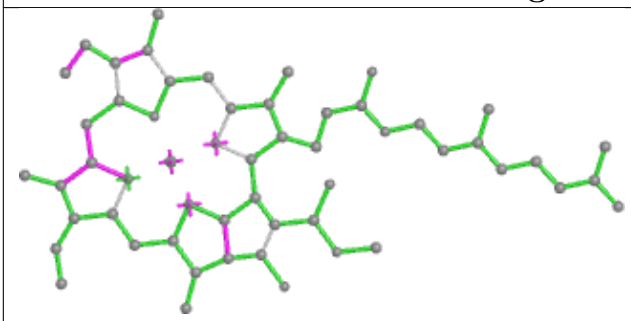


Torsions

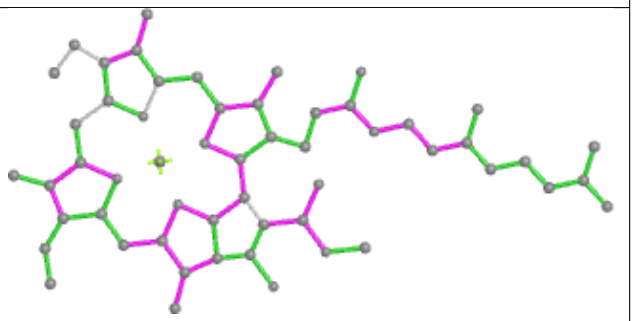


Rings

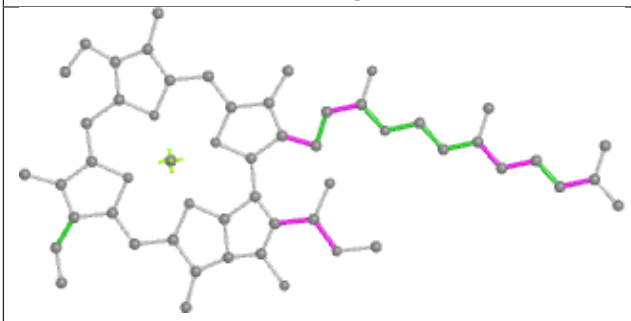
Ligand CLA 5 311



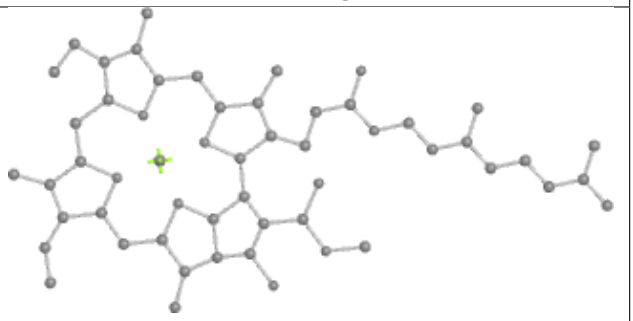
Bond lengths



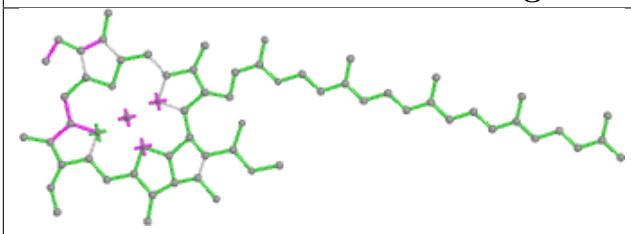
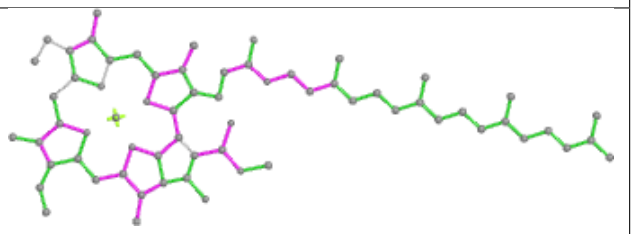
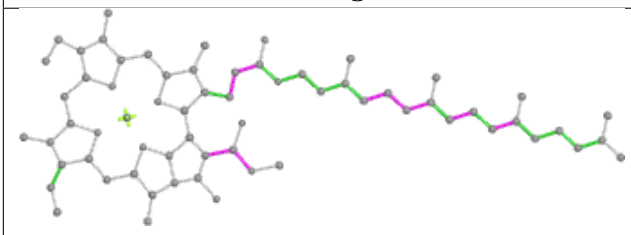
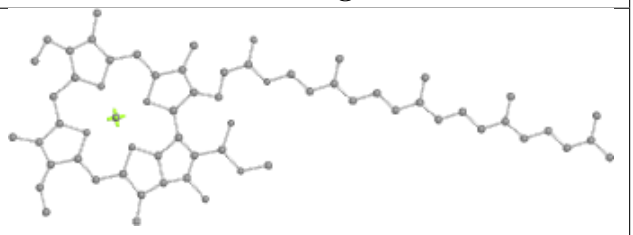
Bond angles



Torsions



Rings

Ligand CLA A 854	
 Bond lengths	 Bond angles
 Torsions	 Rings

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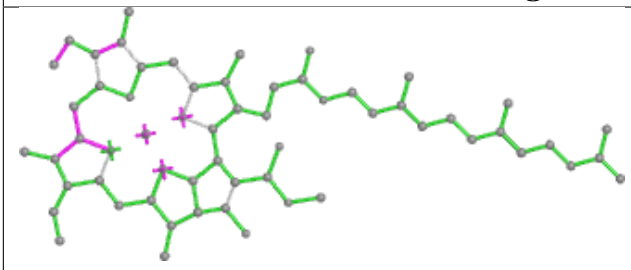
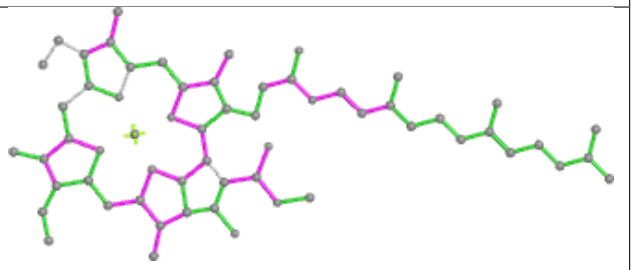
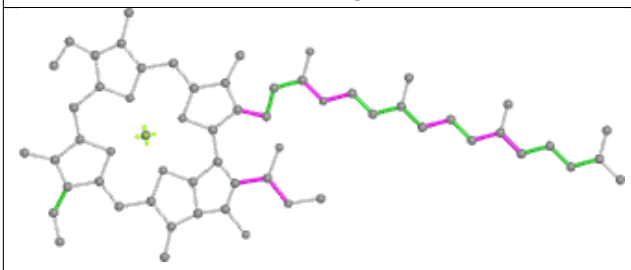
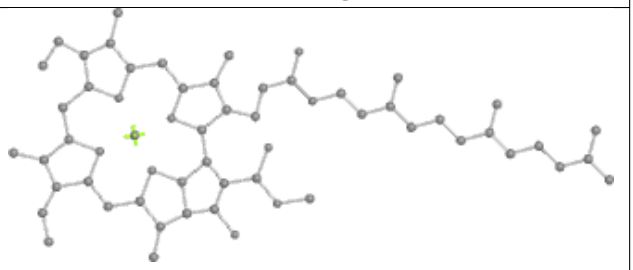
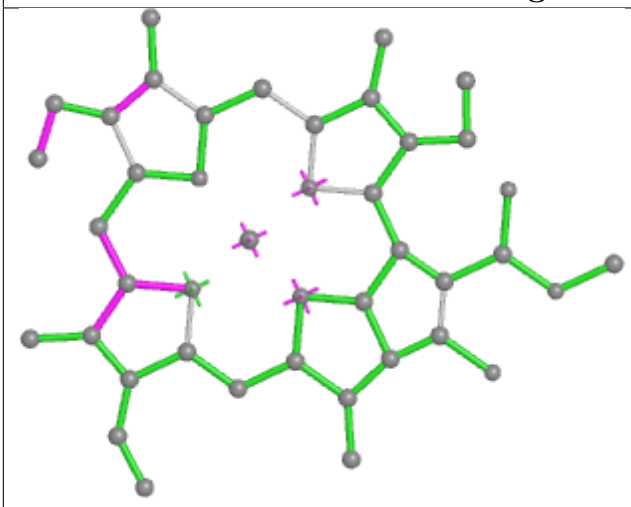
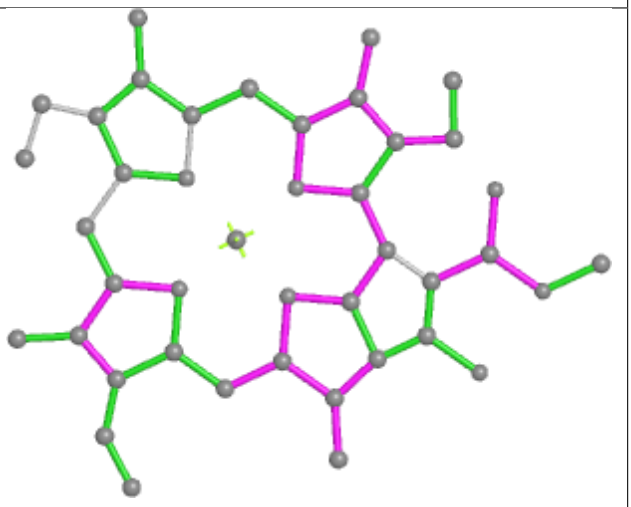
 Bond lengths	 Bond angles
 Torsions	 Rings

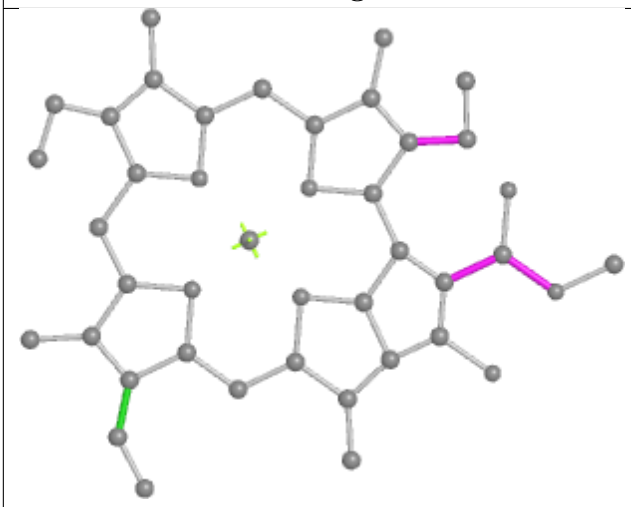
Ligand CLA J 103



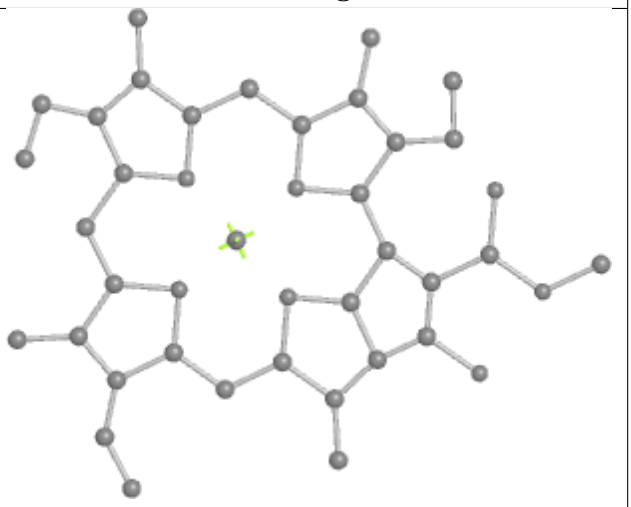
Bond lengths



Bond angles

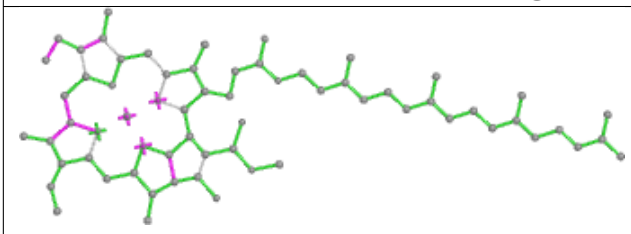


Torsions

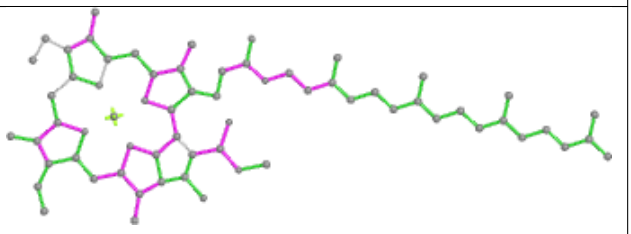


Rings

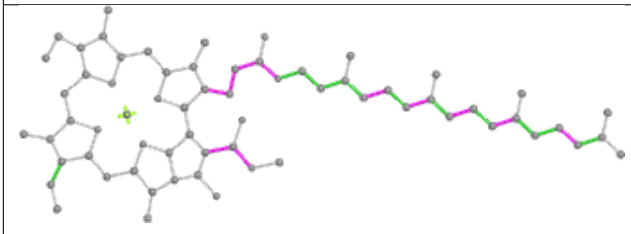
Ligand CLA 3 308



Bond lengths



Bond angles



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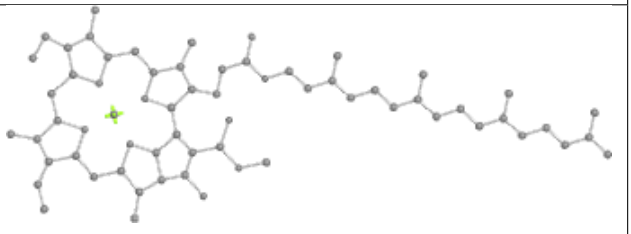
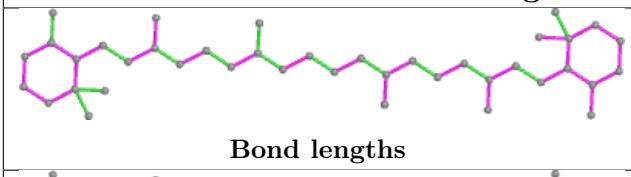


Rings

Ligand BCR B 853



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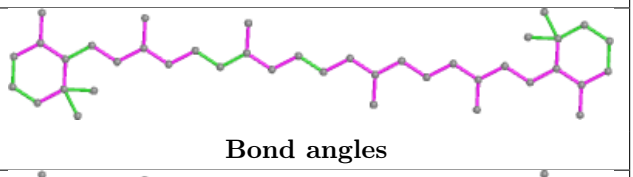
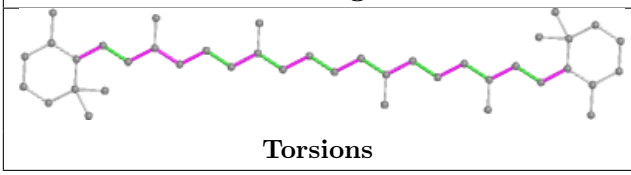
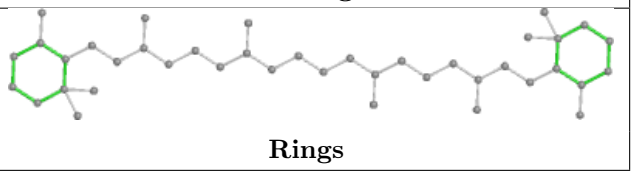


Bond angles

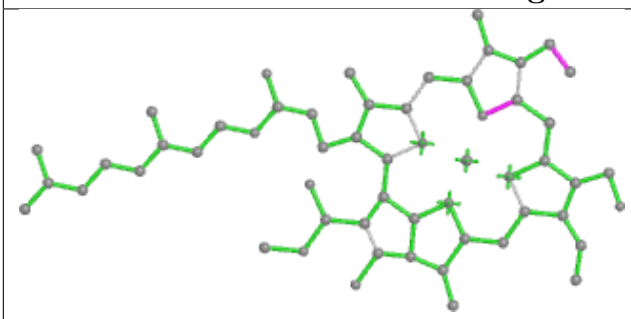


Torsions

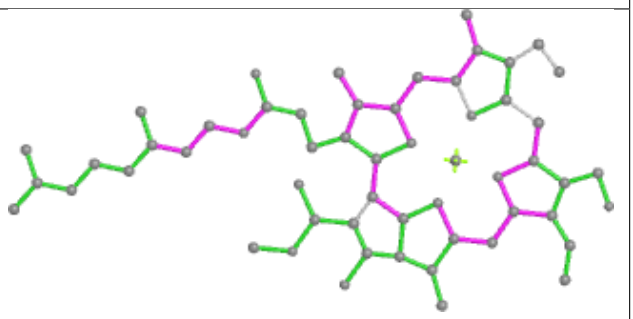


Rings

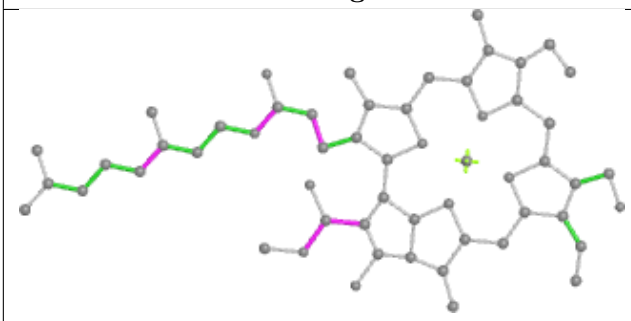
Ligand CHL 6 318



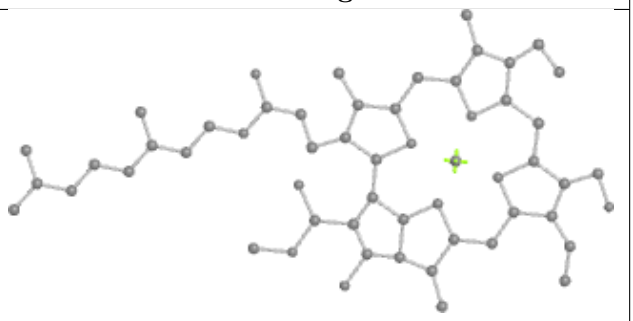
Bond lengths



Bond angles

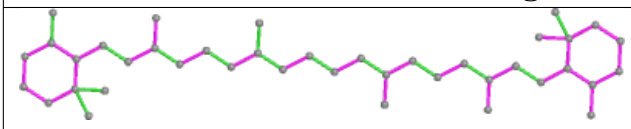


Torsions

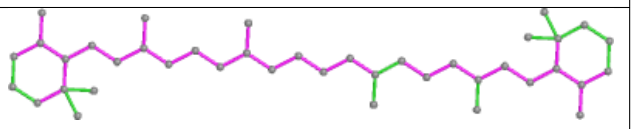


Rings

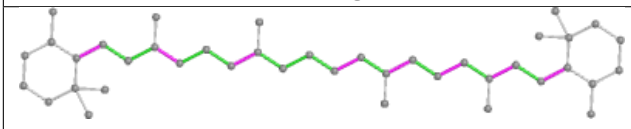
Ligand BCR A 844



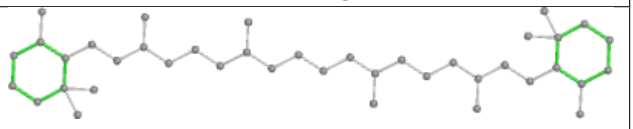
Bond lengths



Bond angles

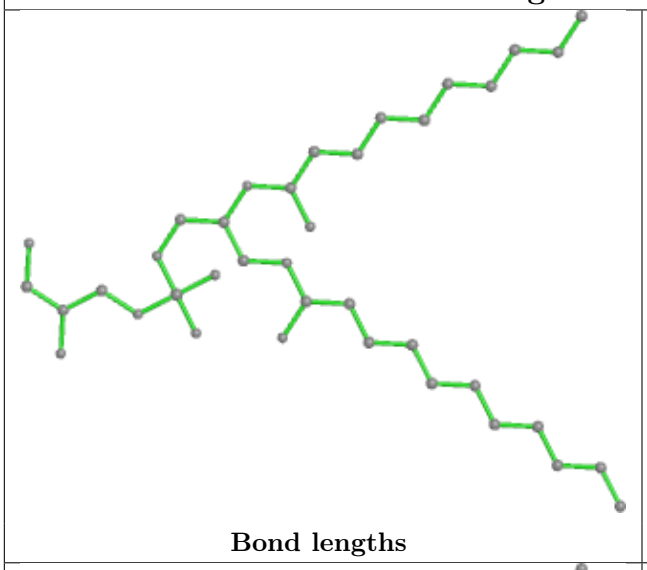


Torsions

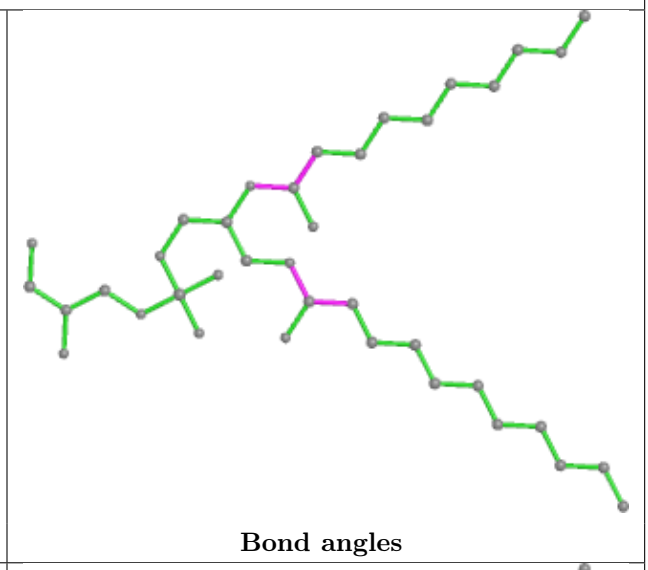


Rings

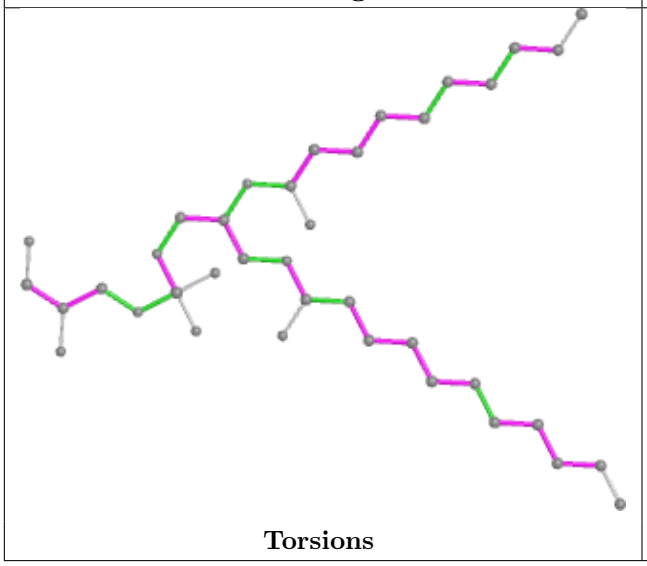
Ligand LHG 8 317



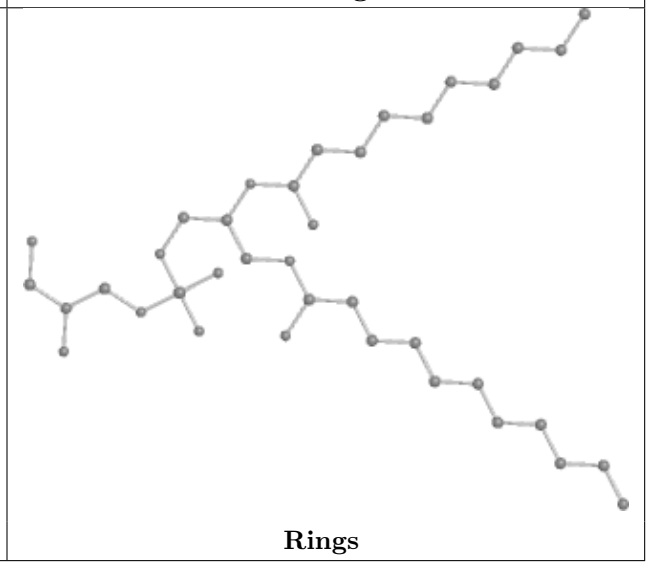
Bond lengths



Bond angles

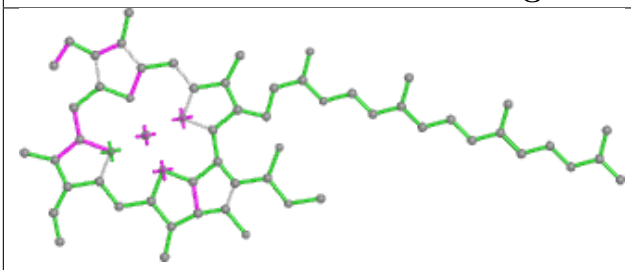


Torsions

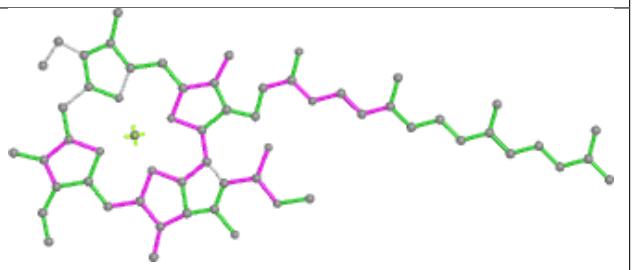


Rings

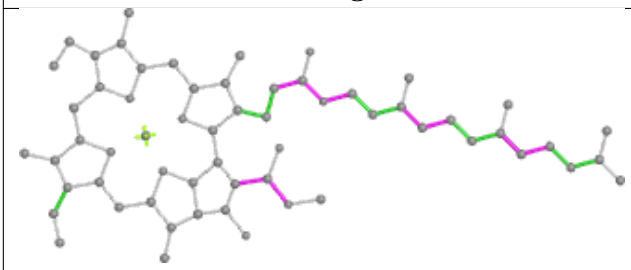
Ligand CLA A 817



Bond lengths



Bond angles



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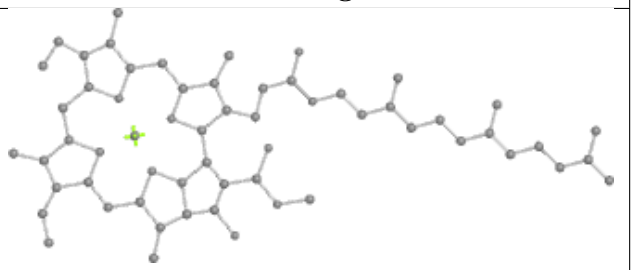
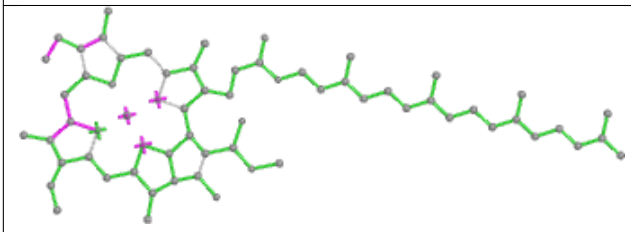
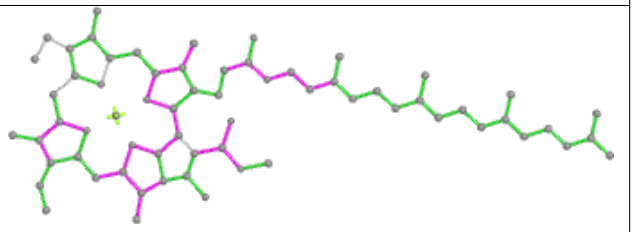


Rings

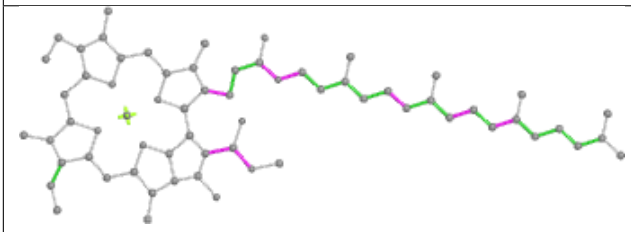
Ligand CLA B 806



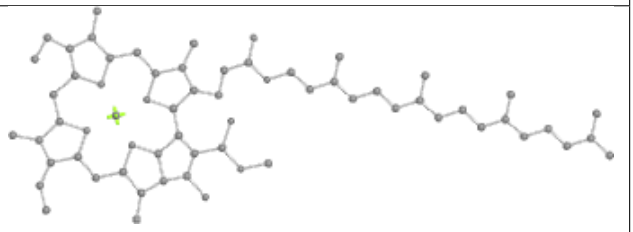
Bond lengths



Bond angles

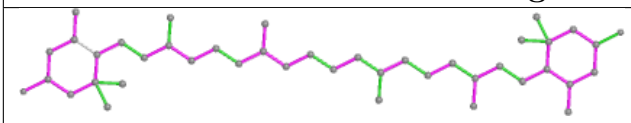


Torsions



Rings

Ligand LUT 3 303



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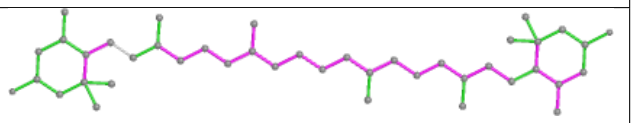
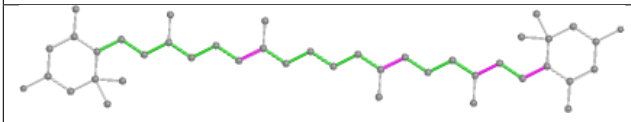


Bond angles



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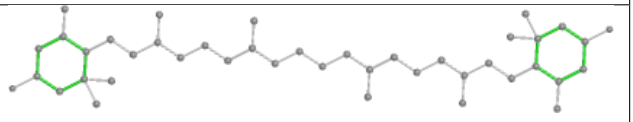
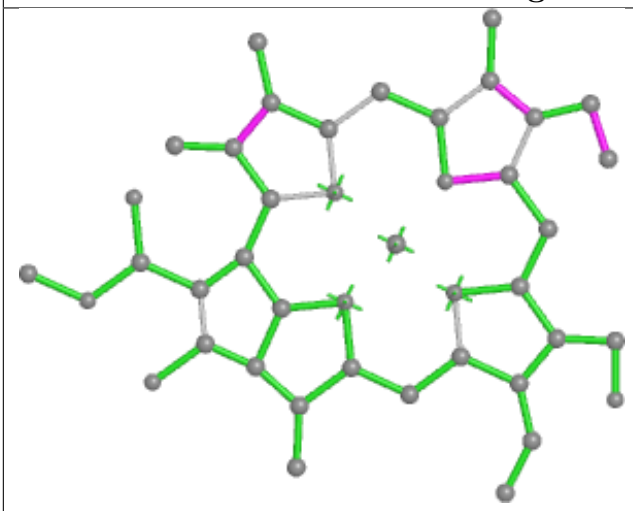


Rings

Ligand CHL 9 314



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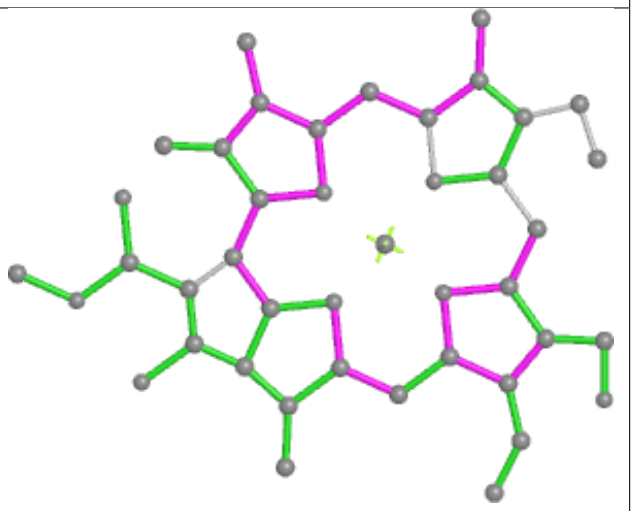
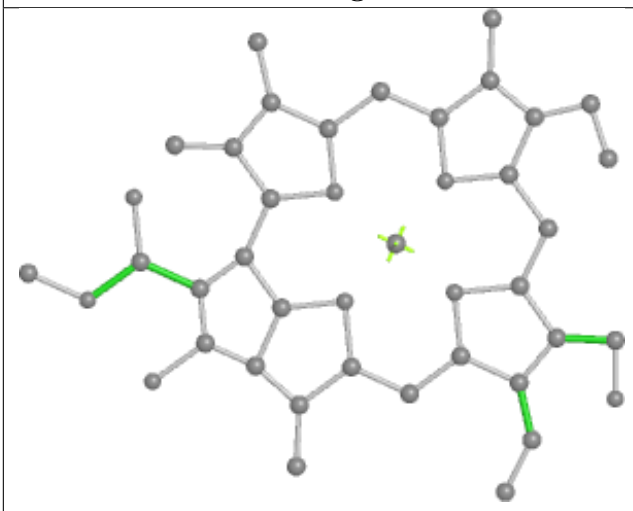


Bond angles



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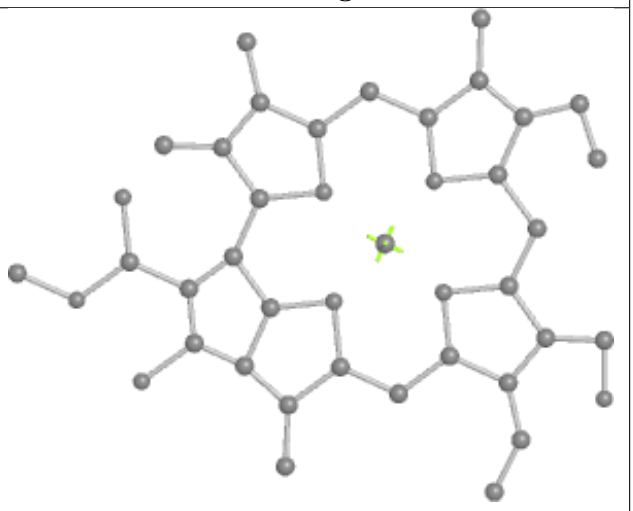
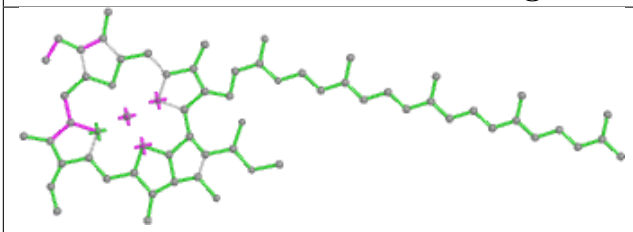


Rings

Ligand CLA A 855



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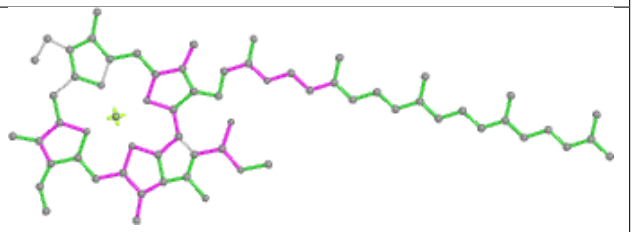
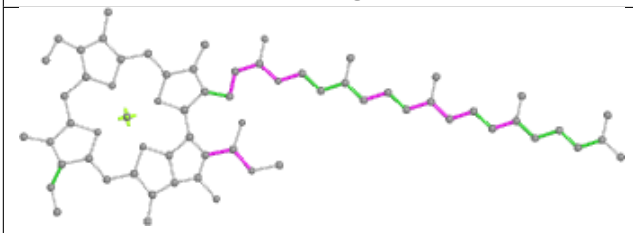
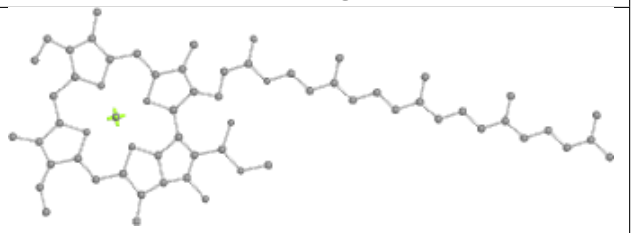


Bond angles

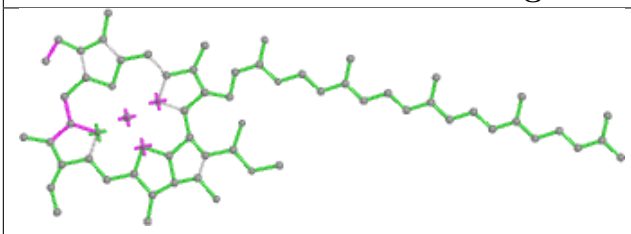


Torsions

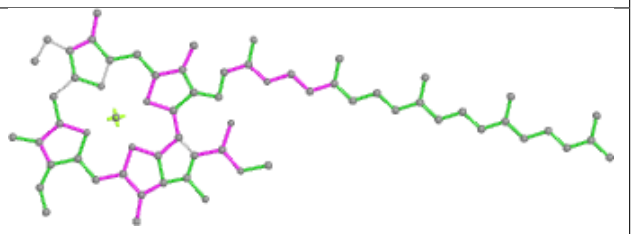


Rings

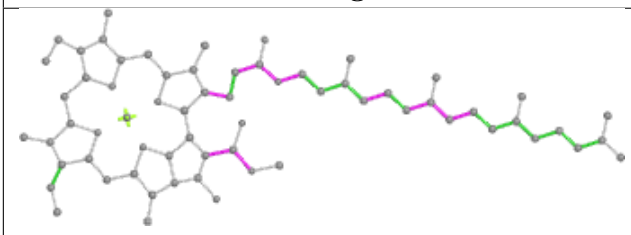
Ligand CLA B 808



Bond lengths



Bond angles



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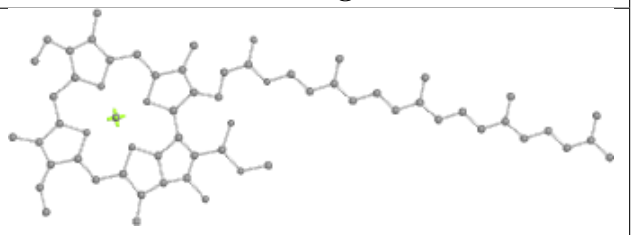
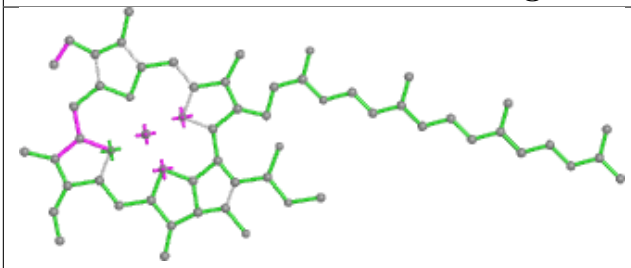


Rings

Ligand CLA A 814



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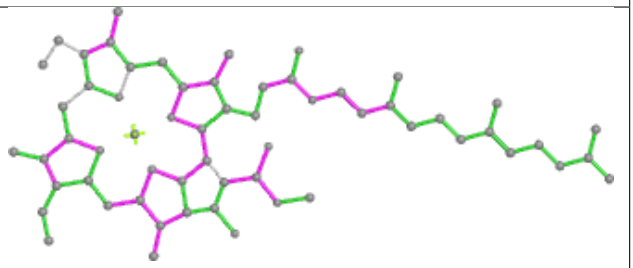
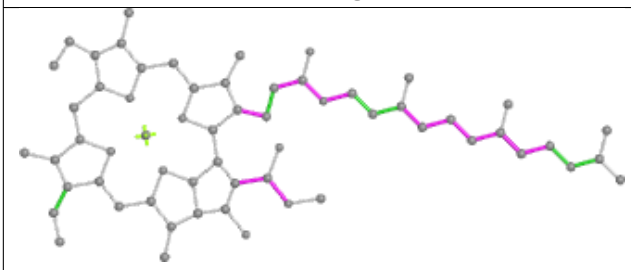


Bond angles



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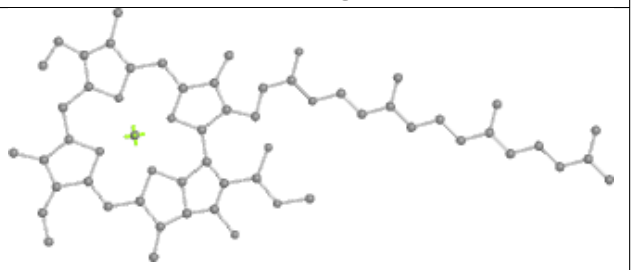
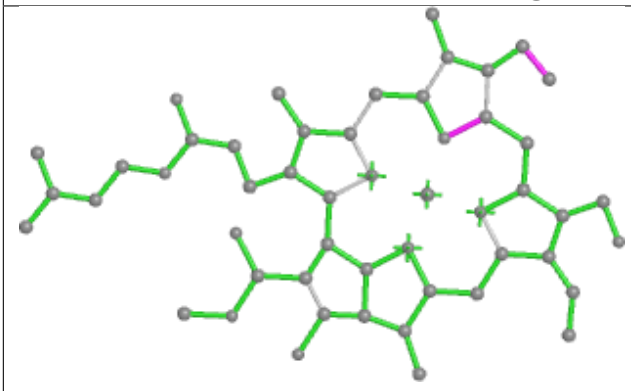
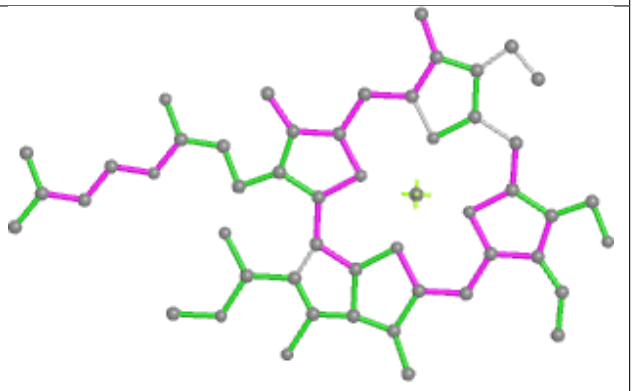


Rings

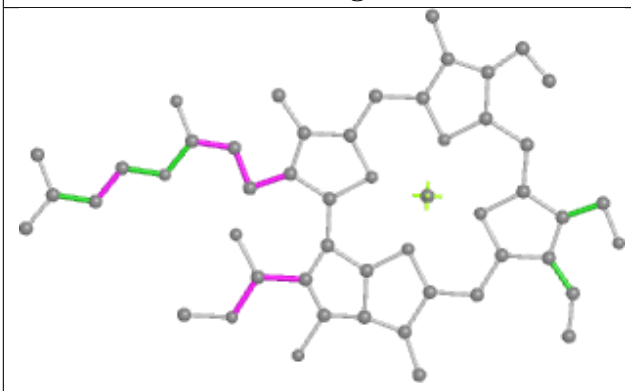
Ligand CHL 4 814



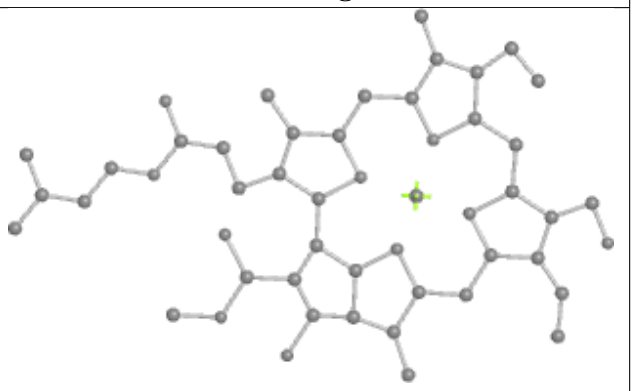
Bond lengths



Bond angles

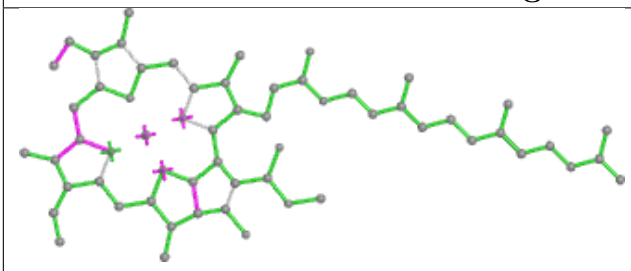


Torsions

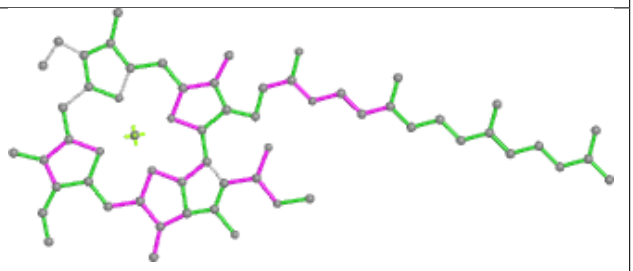


Rings

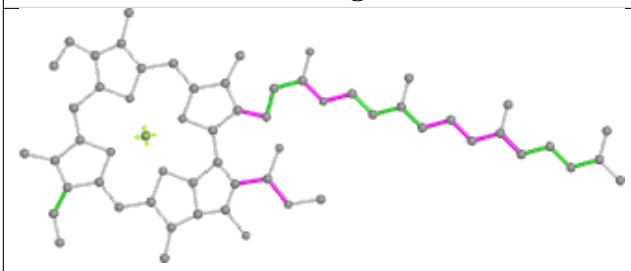
Ligand CLA A 818



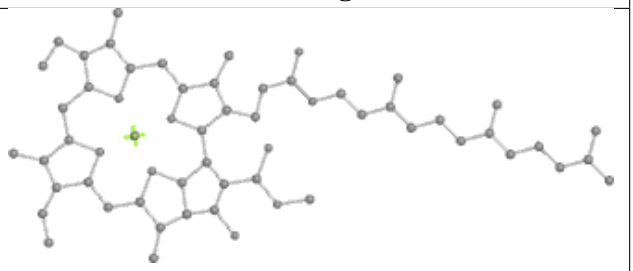
Bond lengths



Bond angles

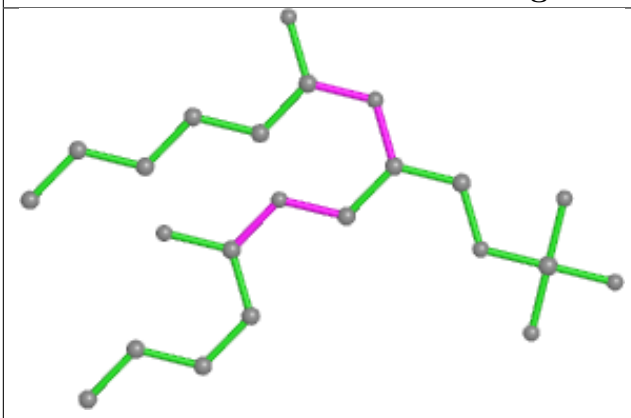


Torsions



Rings

Ligand 3PH 5 325



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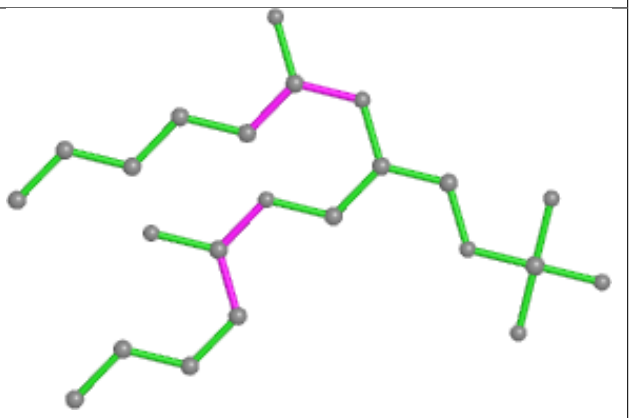
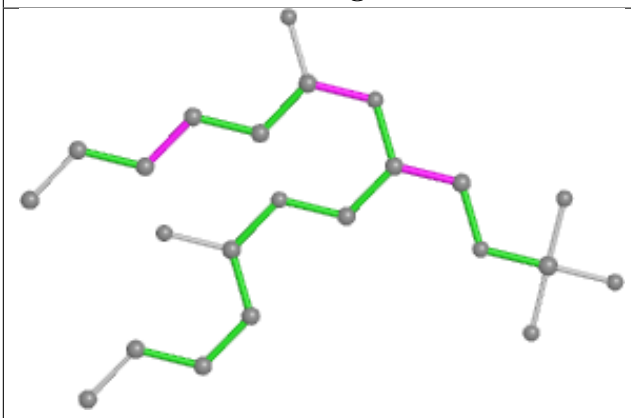
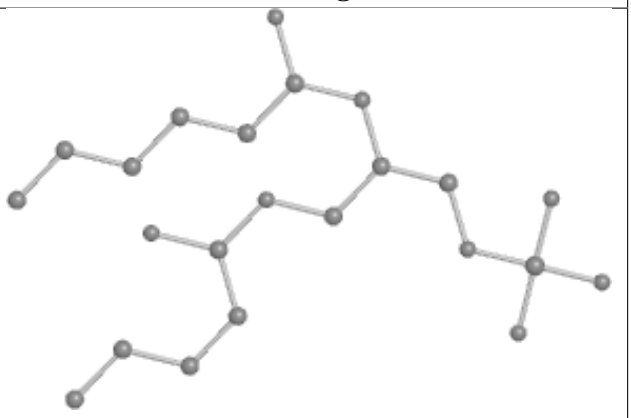


Bond angles

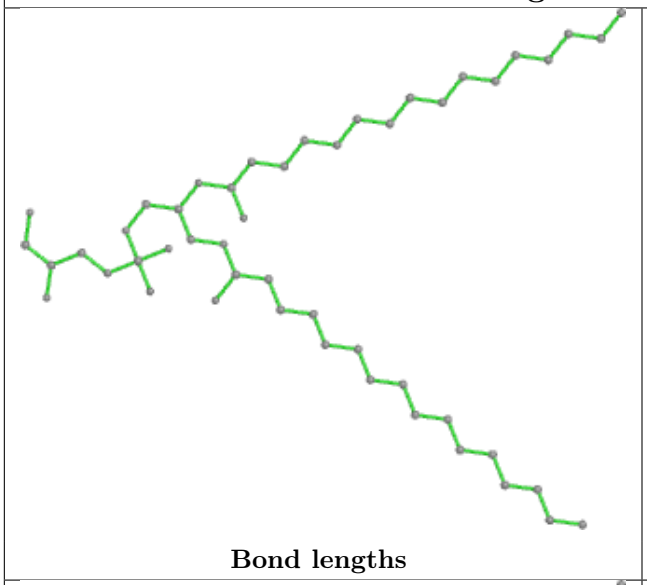


Torsions



Rings

Ligand LHG 4 820



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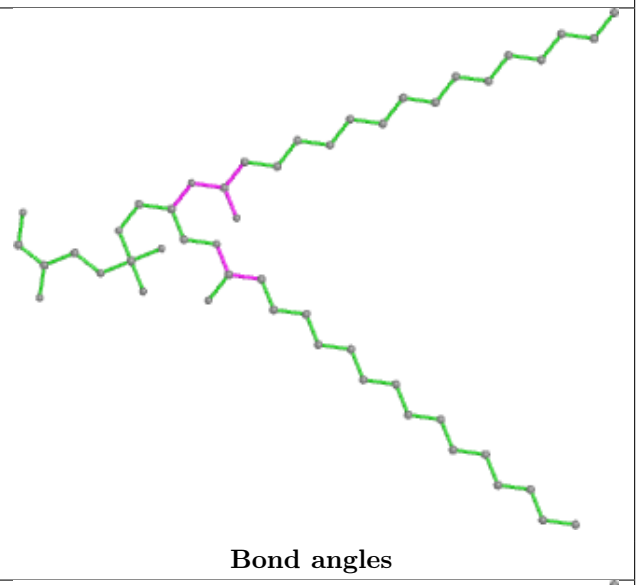
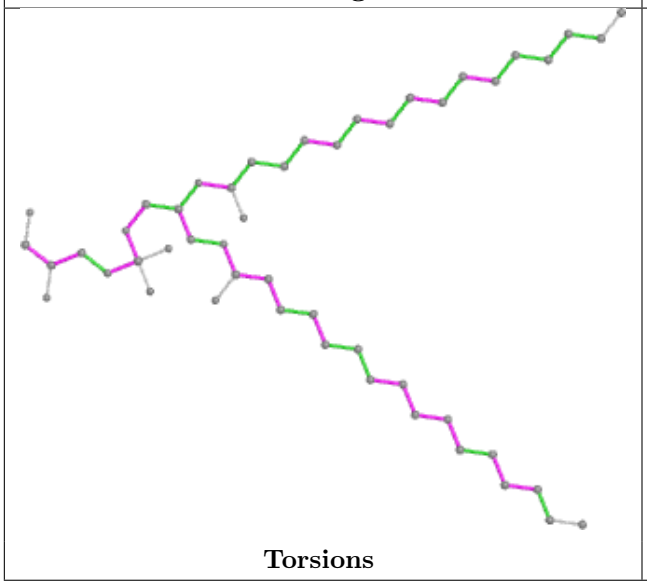


Bond angles



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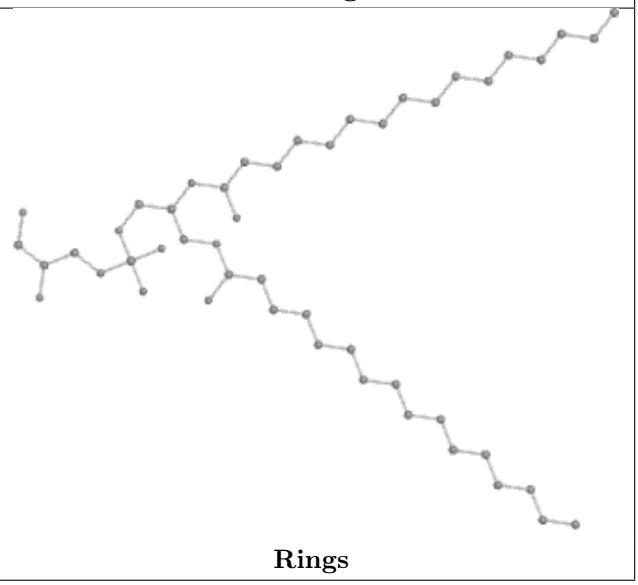
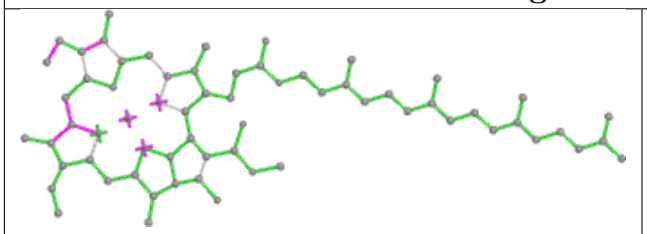


Rings

Ligand CLA 5 318



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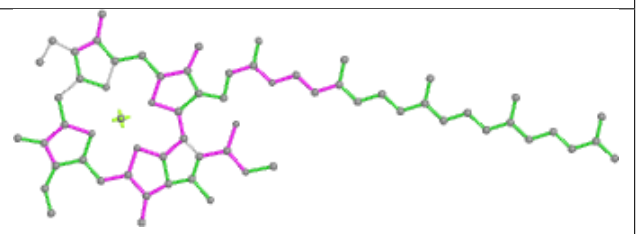
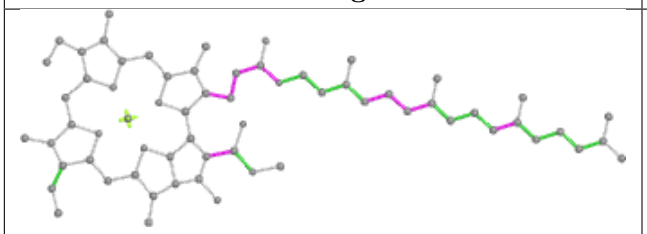
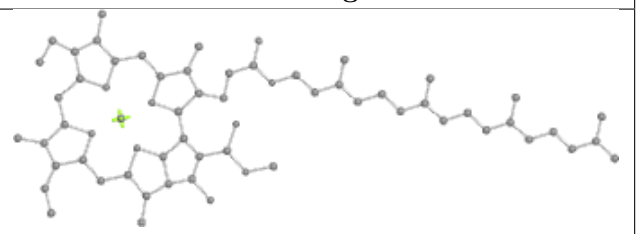


Bond angles

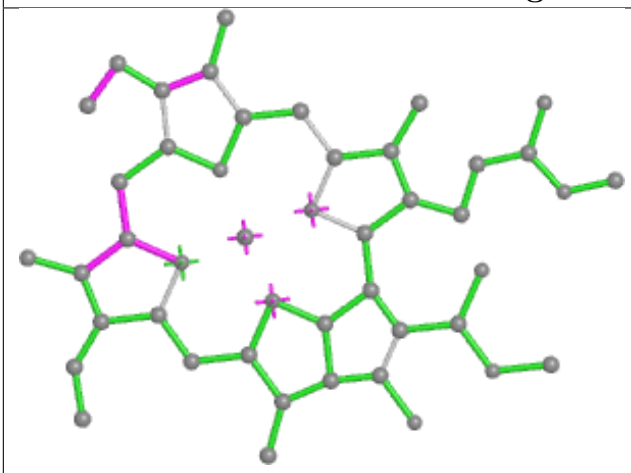


Torsions

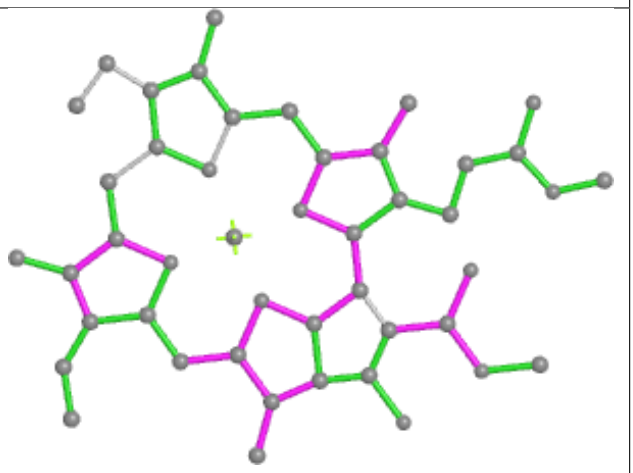


Rings

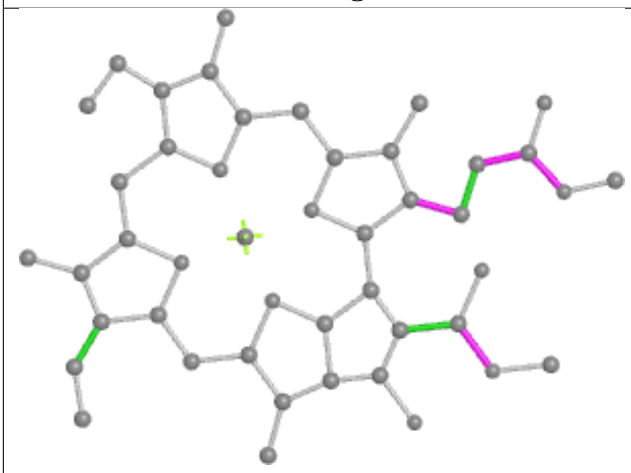
Ligand CLA Z 315



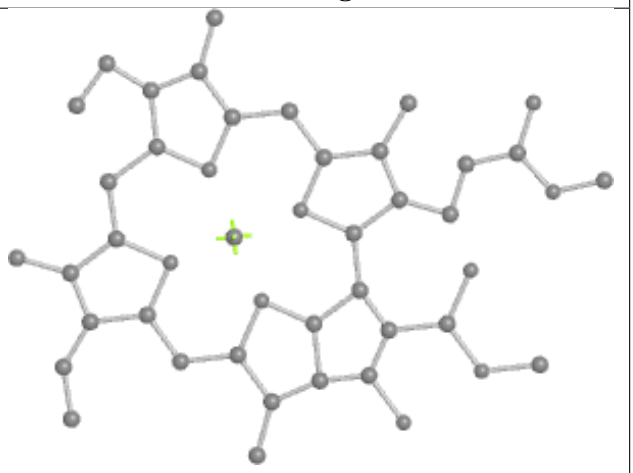
Bond lengths



Bond angles

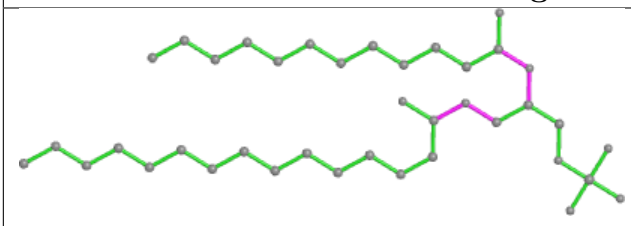


Torsions

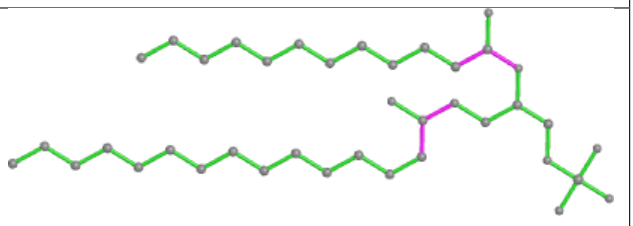


Rings

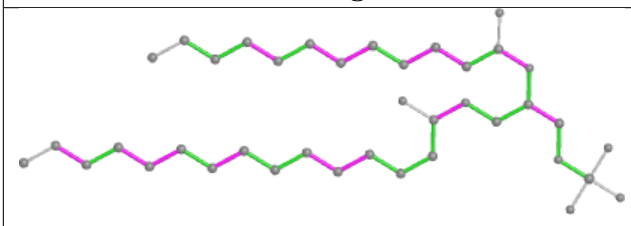
Ligand 3PH 7 319



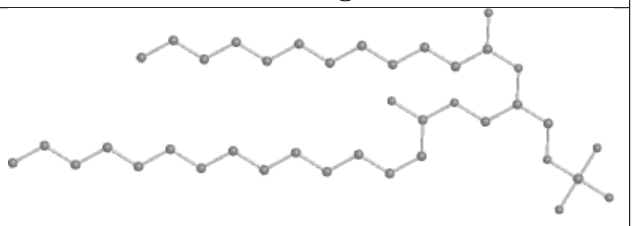
Bond lengths



Bond angles

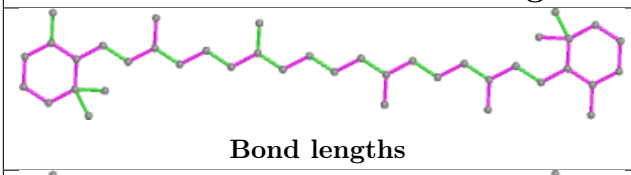


Torsions

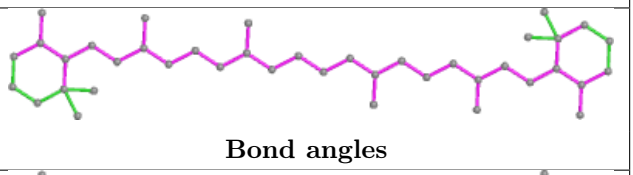


Rings

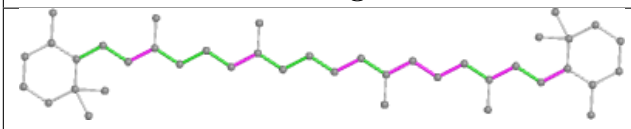
Ligand BCR 3 307



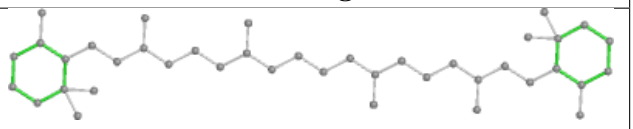
Bond lengths



Bond angles

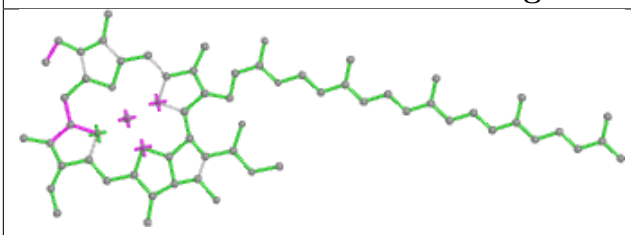


Torsions

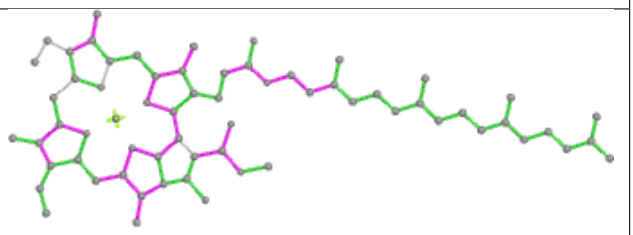


Rings

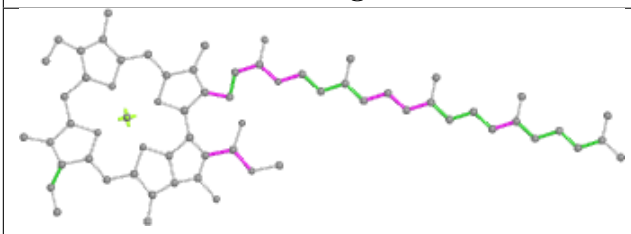
Ligand CLA A 815



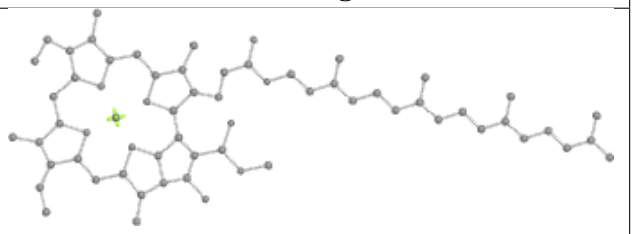
Bond lengths



Bond angles

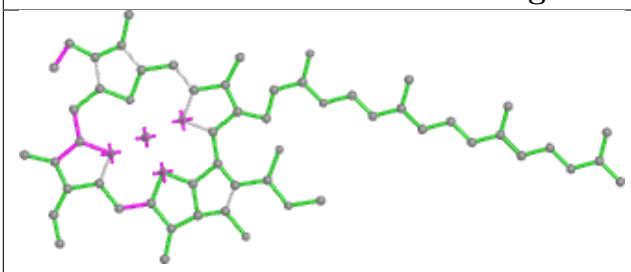


Torsions

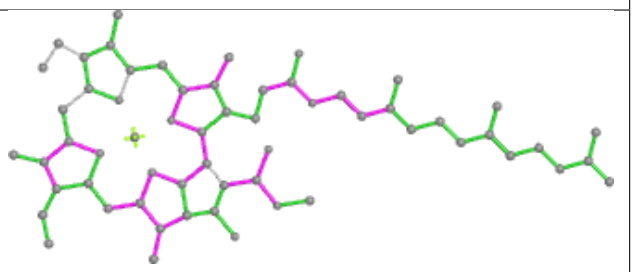


Rings

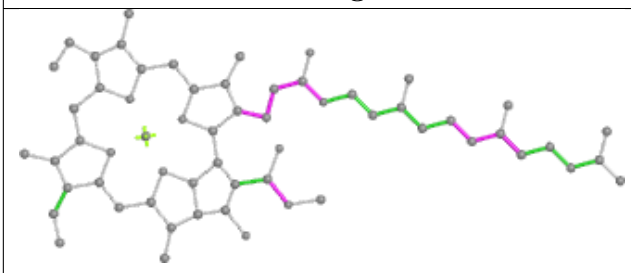
Ligand CLA 4 805



Bond lengths



Bond angles



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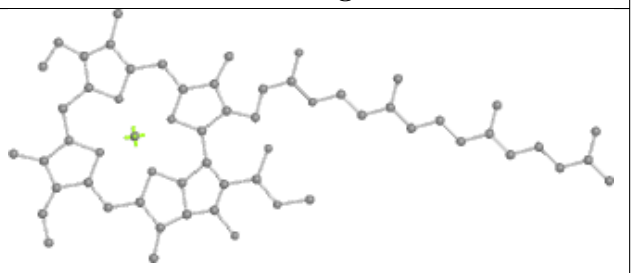
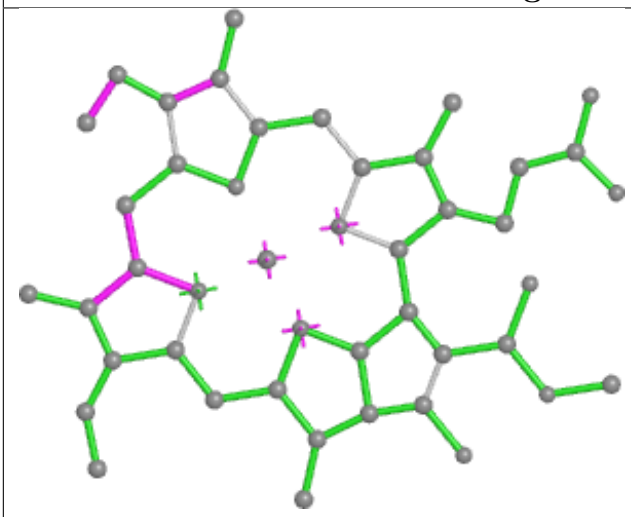
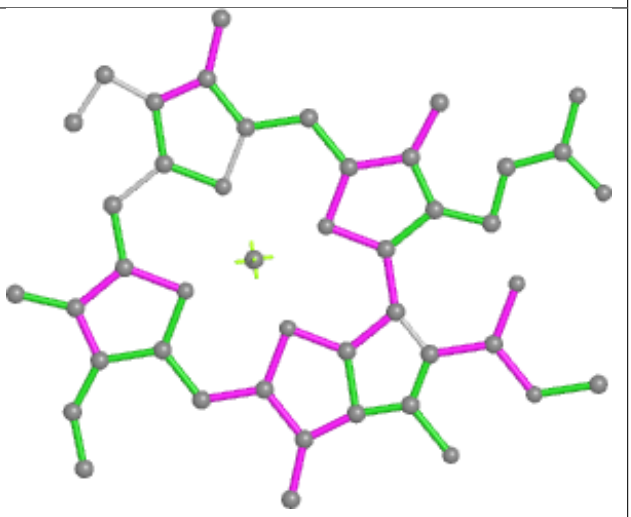


Rings

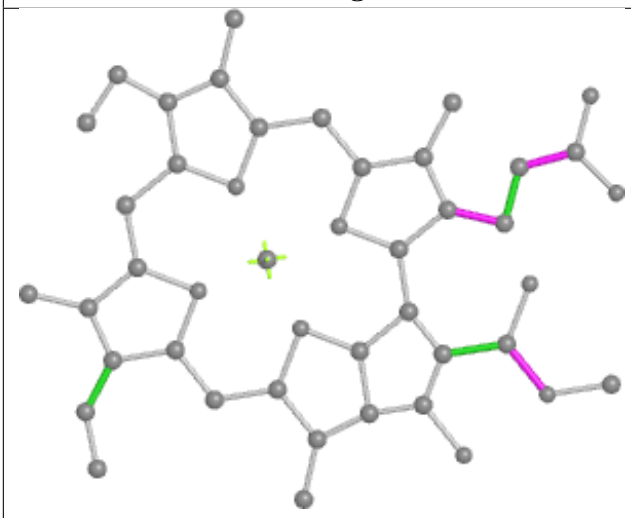
Ligand CLA A 804



Bond lengths



Bond angles



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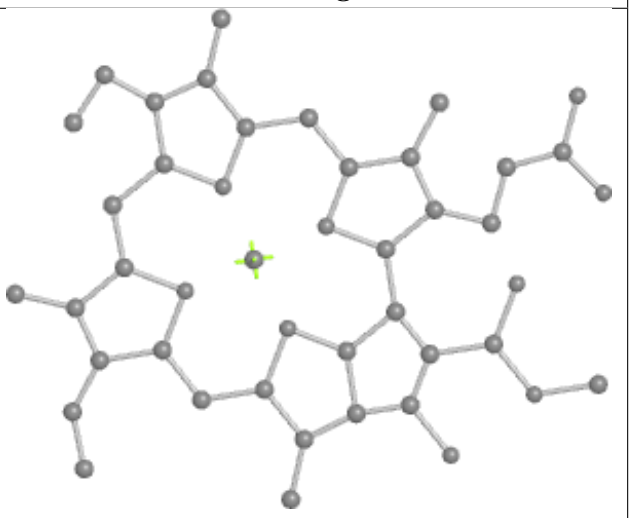
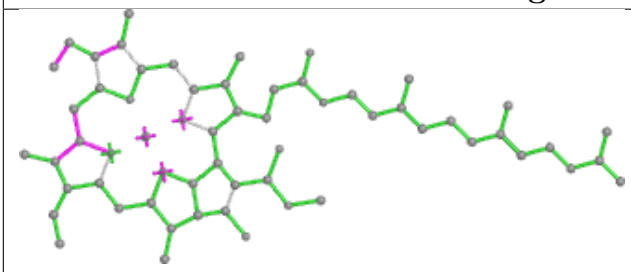
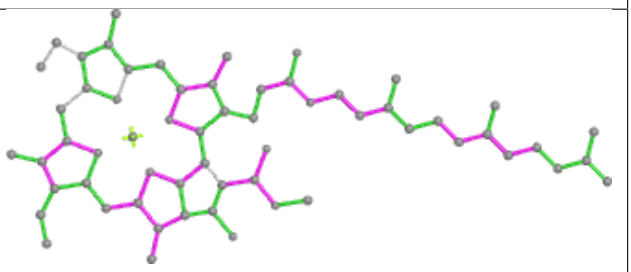


Rings

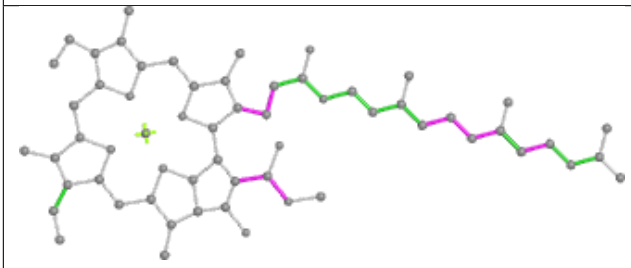
Ligand CLA 3 318



Bond lengths



Bond angles



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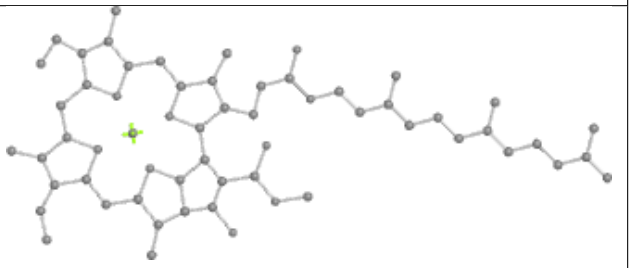
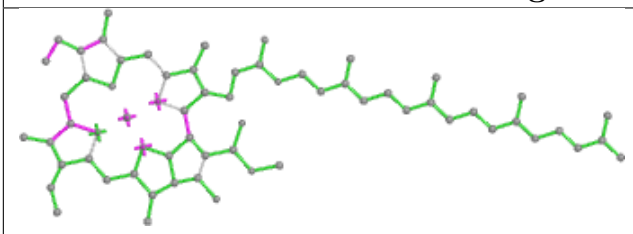


Rings

Ligand CLA 4 807



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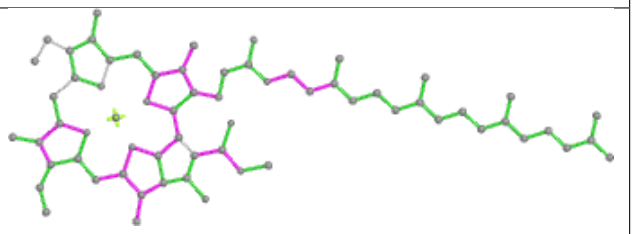
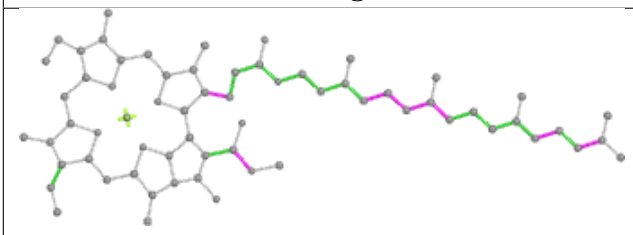
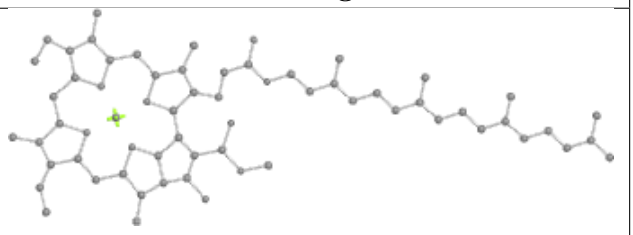


Bond angles

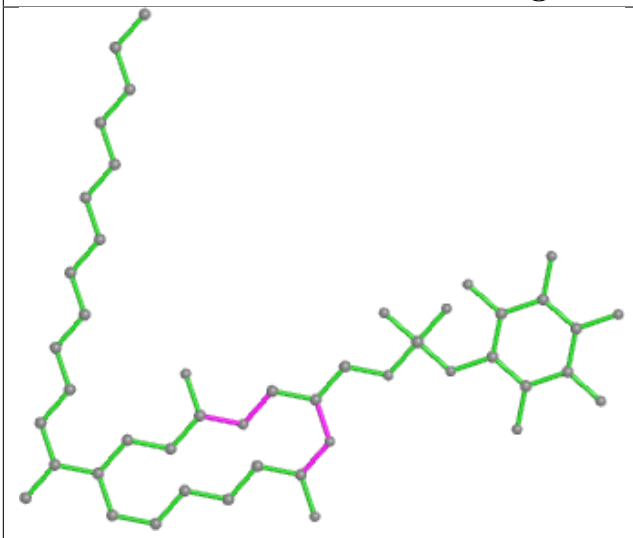


Torsions

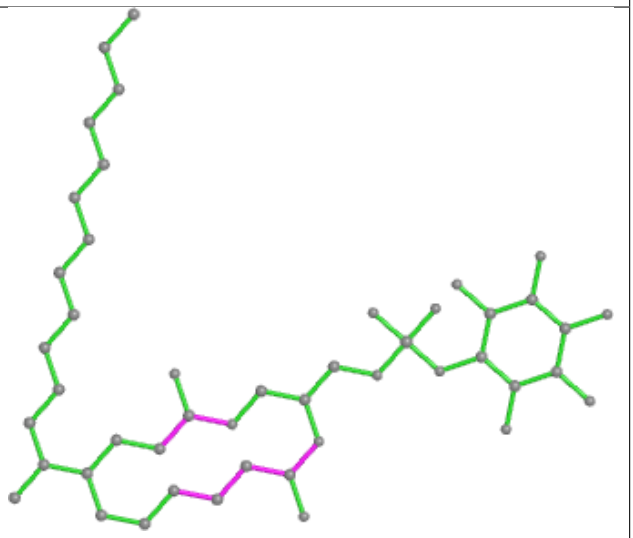


Rings

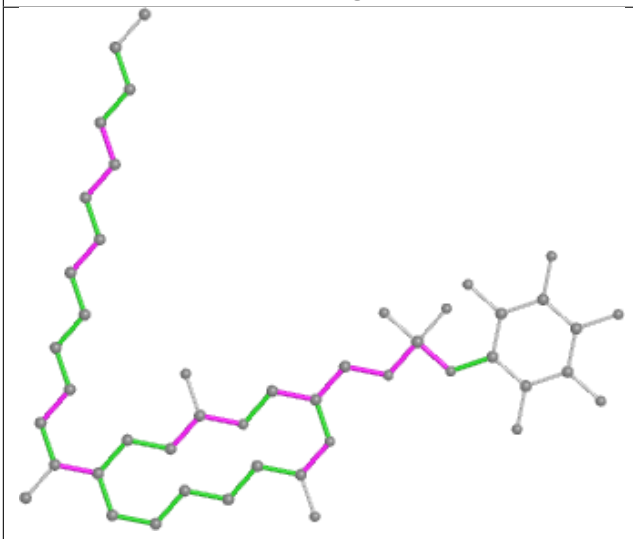
Ligand T7X J 102



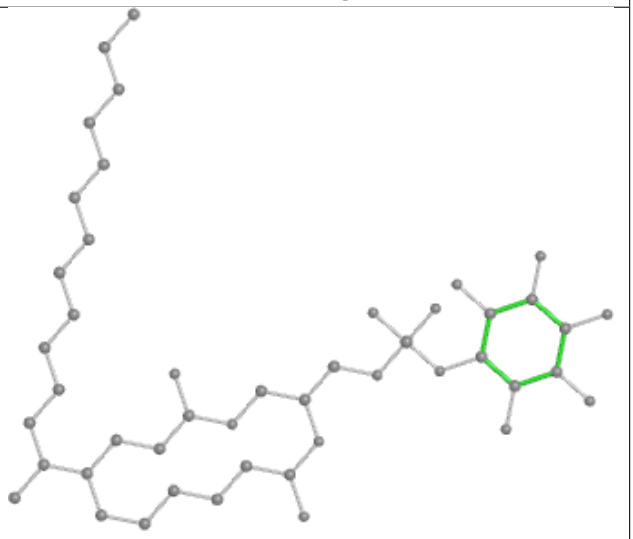
Bond lengths



Bond angles

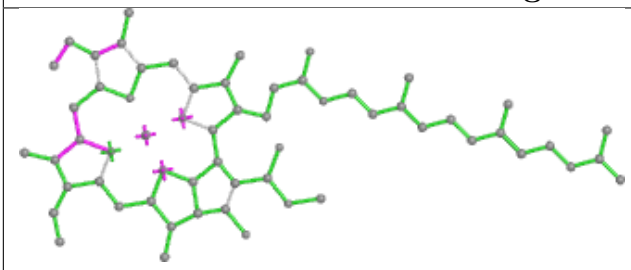


Torsions



Rings

Ligand CLA A 807



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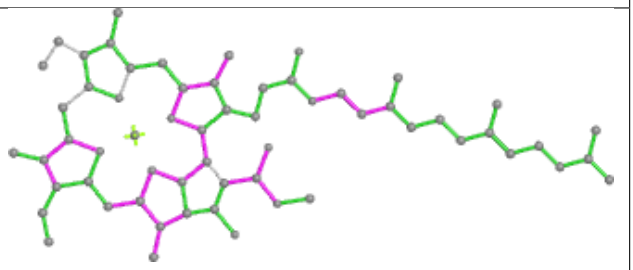
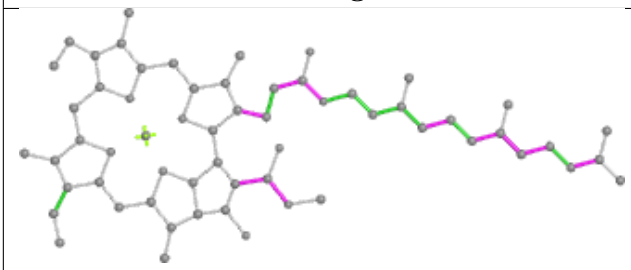
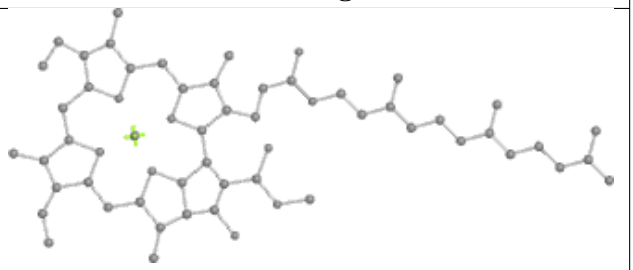


Bond angles

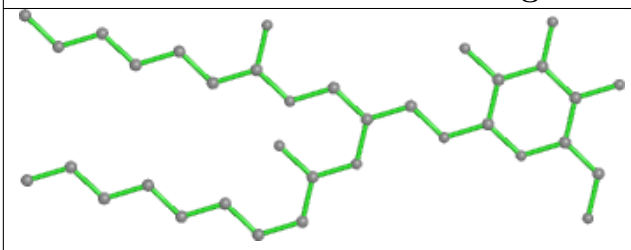


Torsions

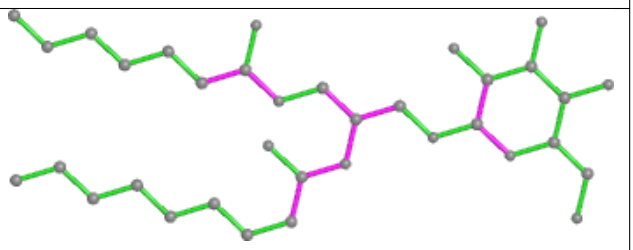


Rings

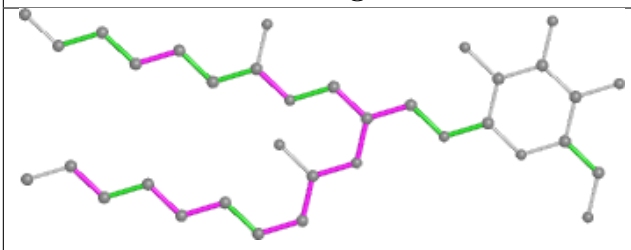
Ligand LMG F 308



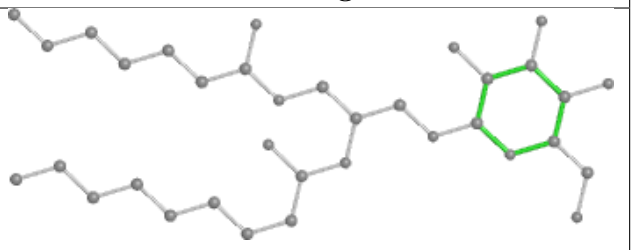
Bond lengths



Bond angles

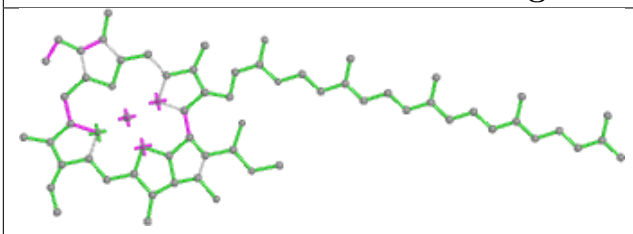


Torsions

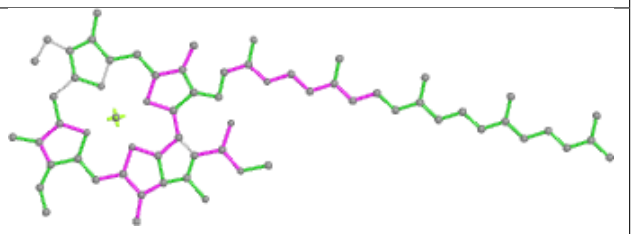


Rings

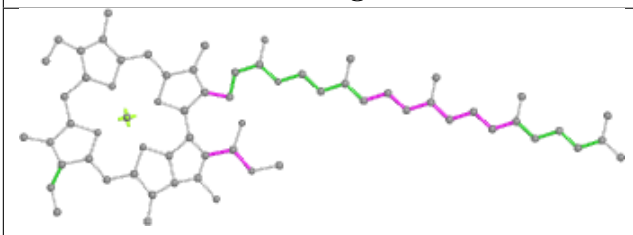
Ligand CLA 7 306



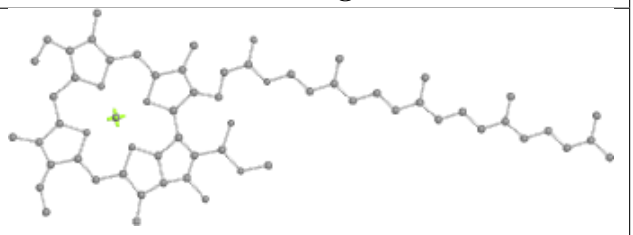
Bond lengths



Bond angles

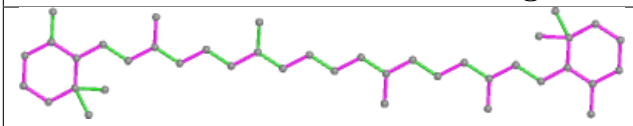


Torsions

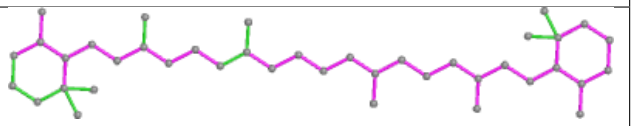


Rings

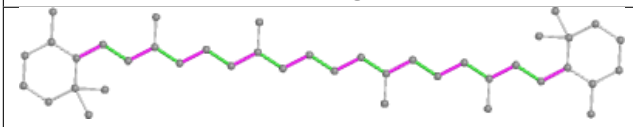
Ligand BCR A 856



Bond lengths



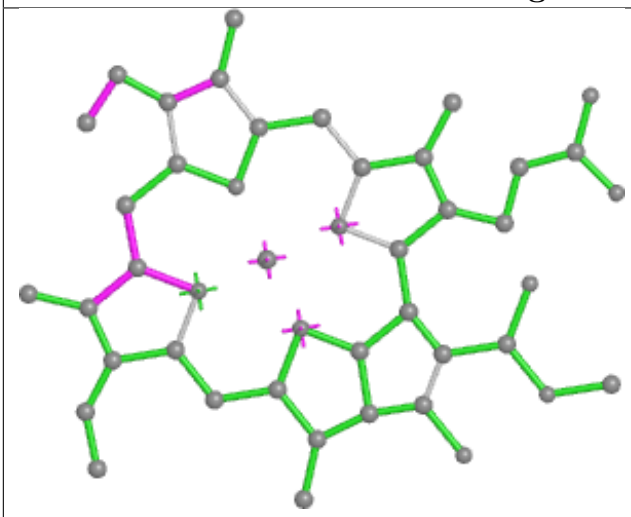
Bond angles



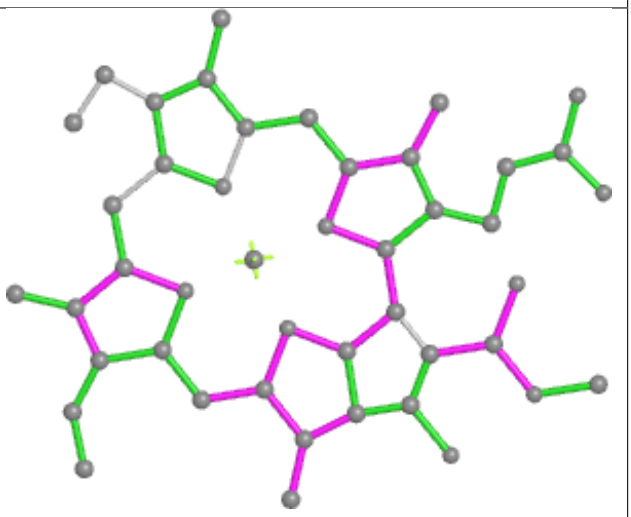
Torsions

Rings

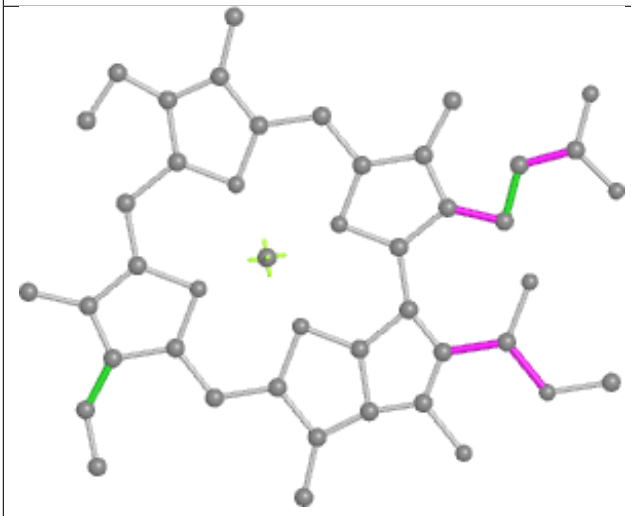
Ligand CLA 2 302



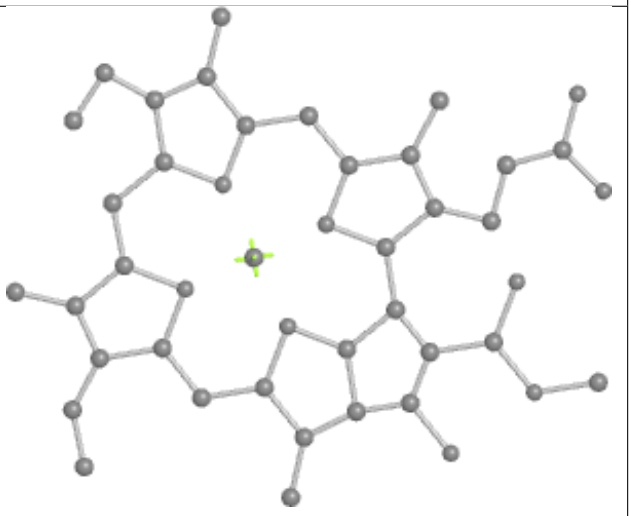
Bond lengths



Bond angles

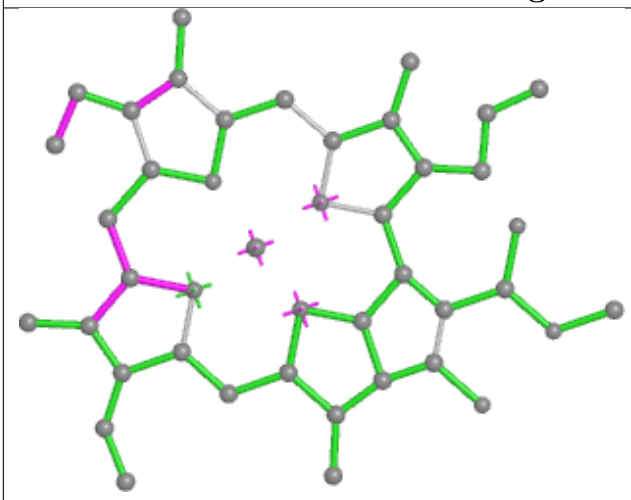


Torsions

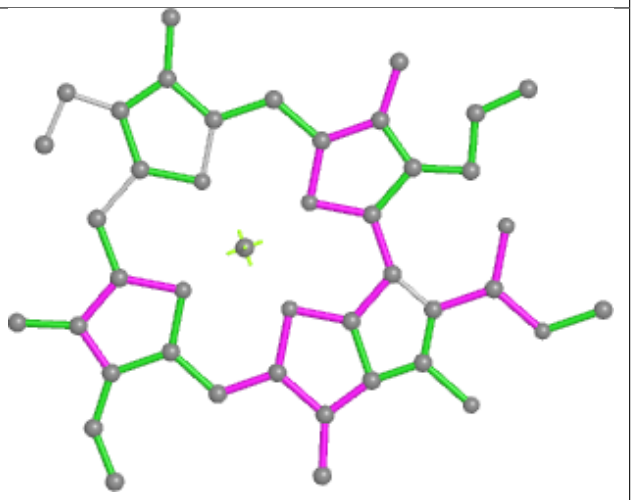


Rings

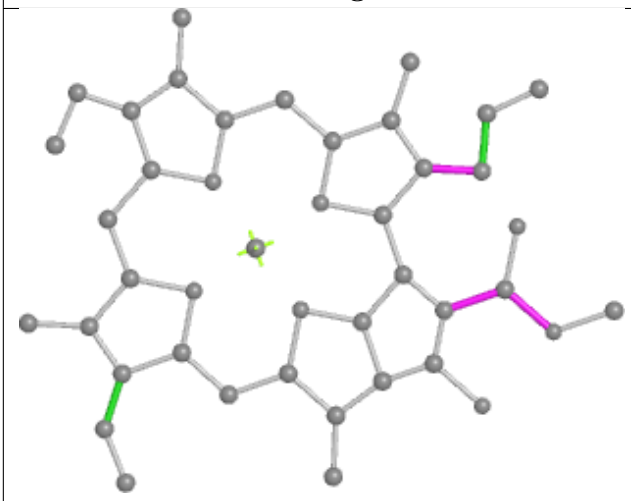
Ligand CLA 7 311



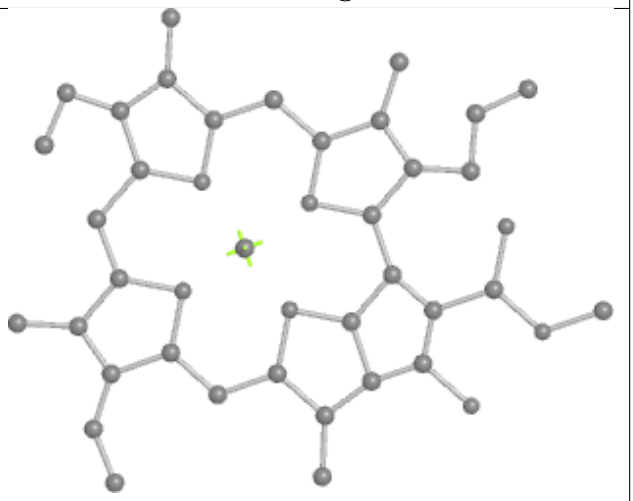
Bond lengths



Bond angles

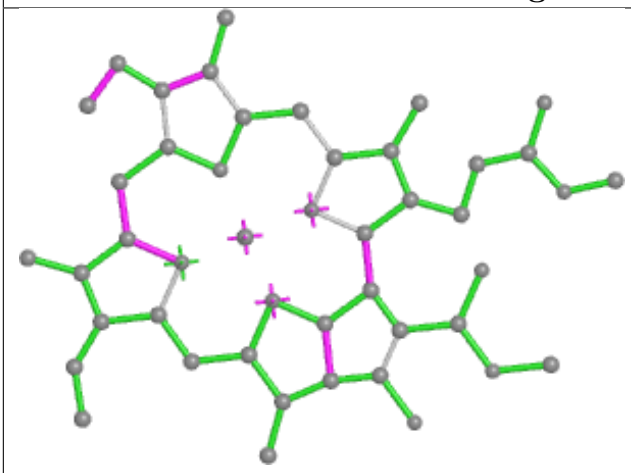


Torsions



Rings

Ligand CLA 1 311



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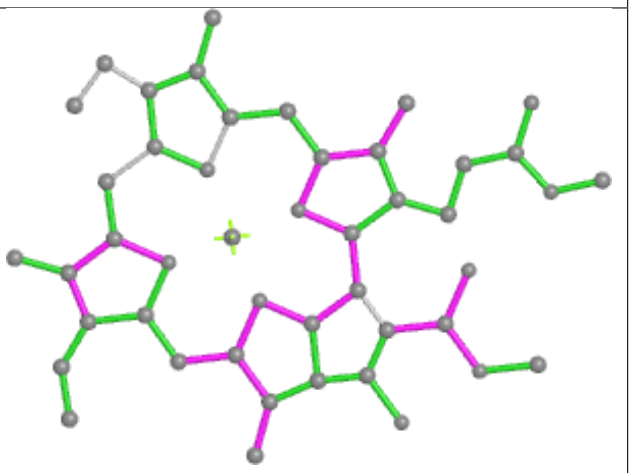
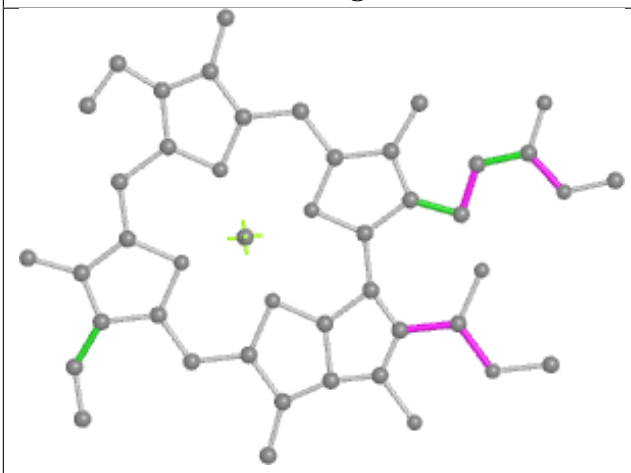


Bond angles



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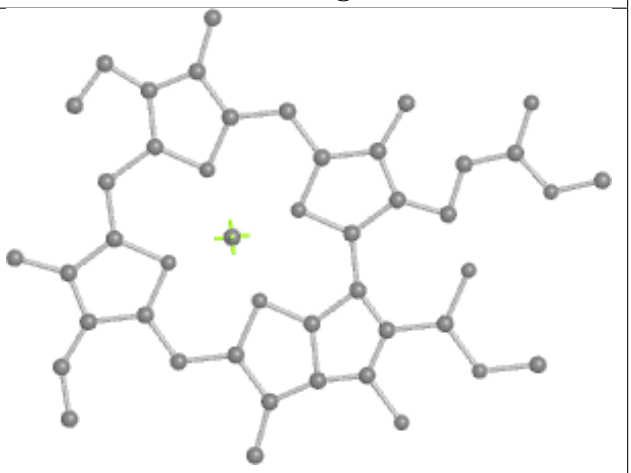
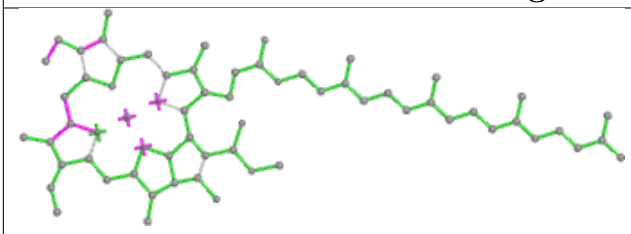


Rings

Ligand CLA 1 314



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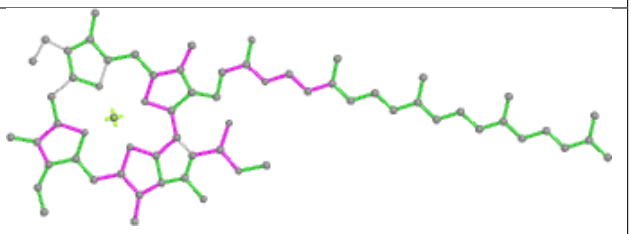
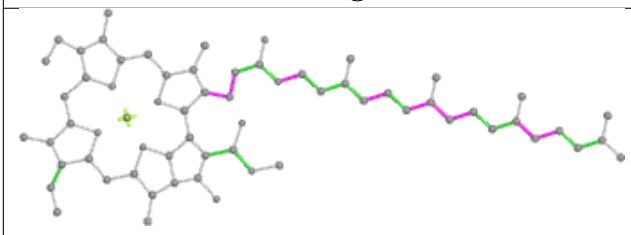


Bond angles



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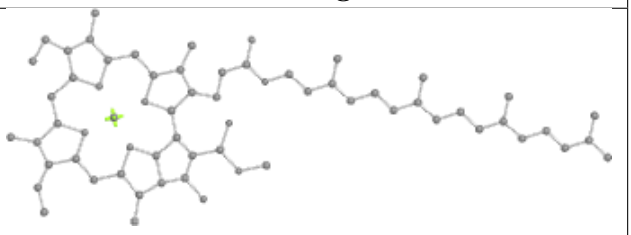
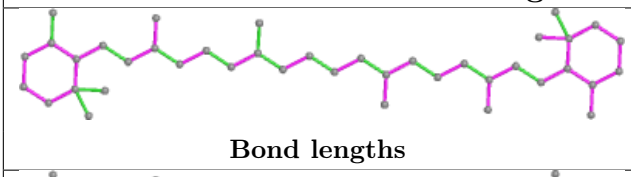
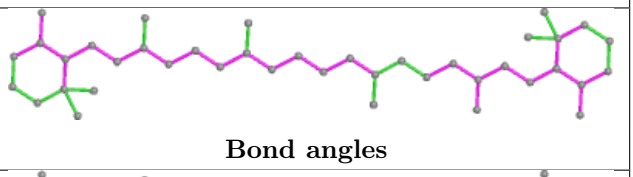


Rings

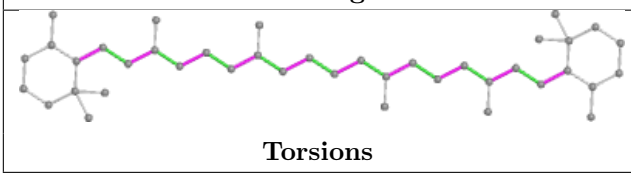
Ligand BCR 7 303



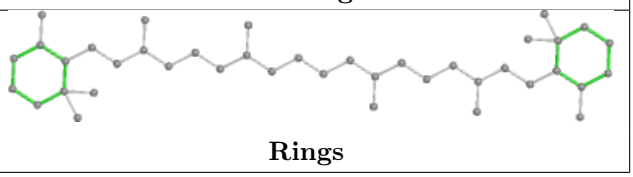
Bond lengths



Bond angles

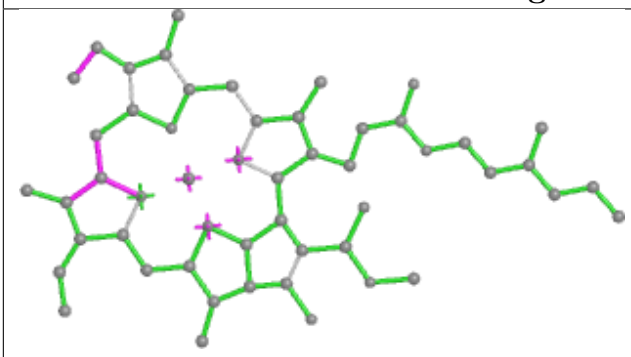


Torsions

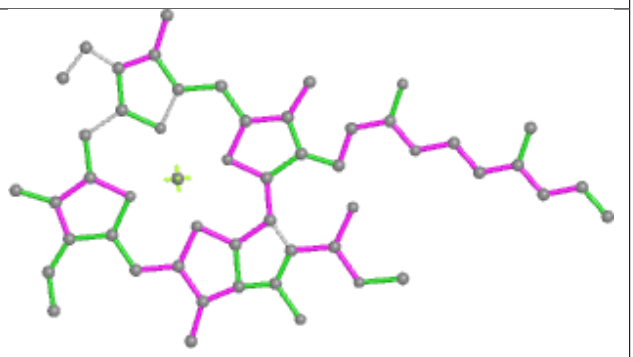


Rings

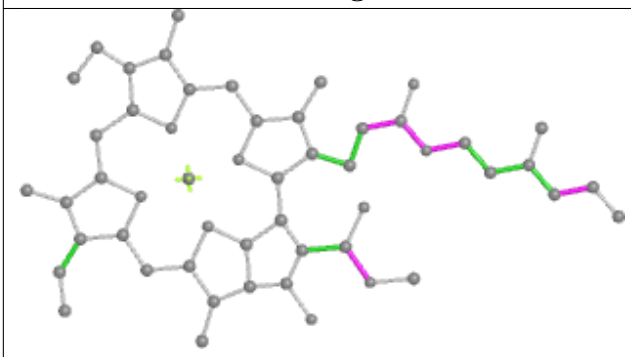
Ligand CLA 6 308



Bond lengths



Bond angles



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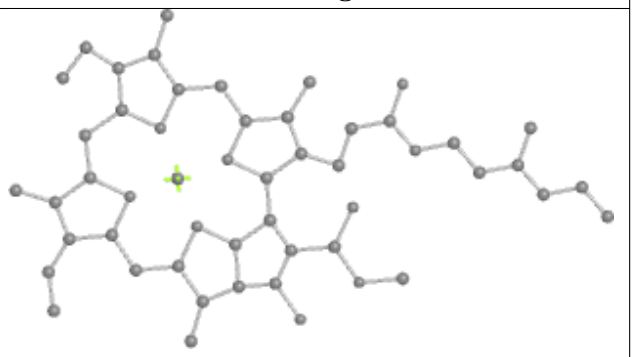
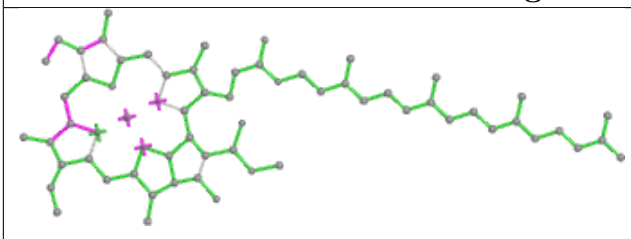
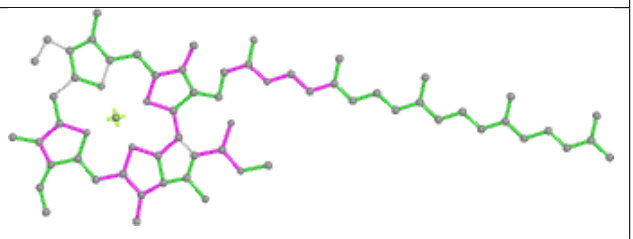


Rings

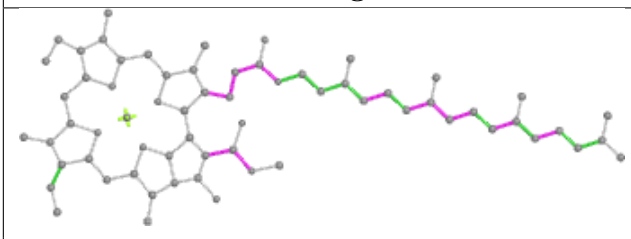
Ligand CLA B 821



Bond lengths



Bond angles



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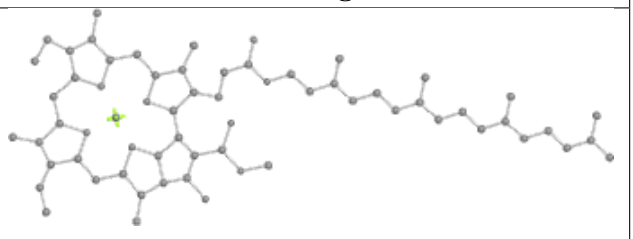
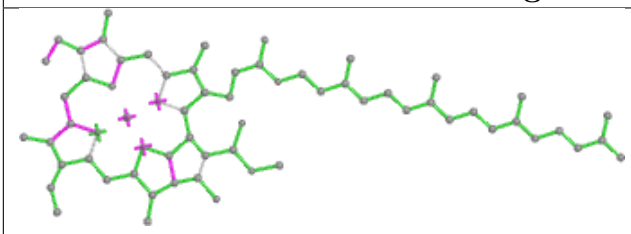
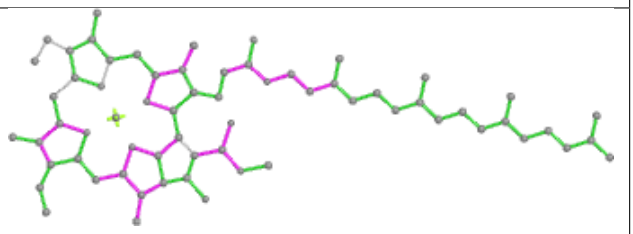


Rings

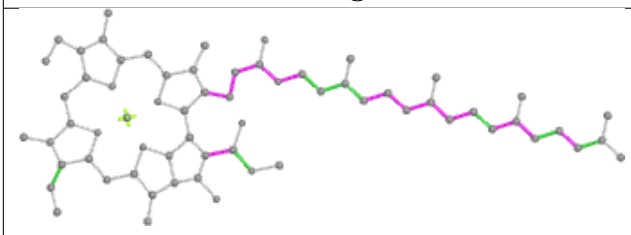
Ligand CLA A 805



Bond lengths



Bond angles



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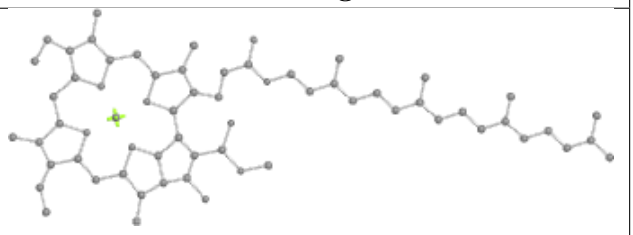
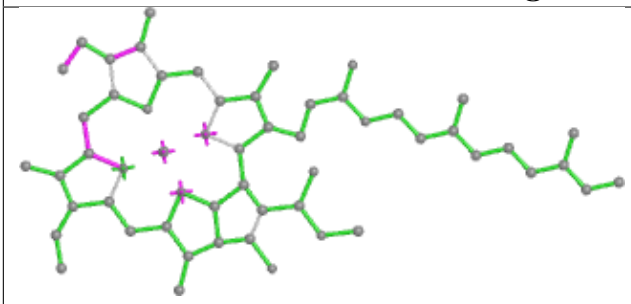
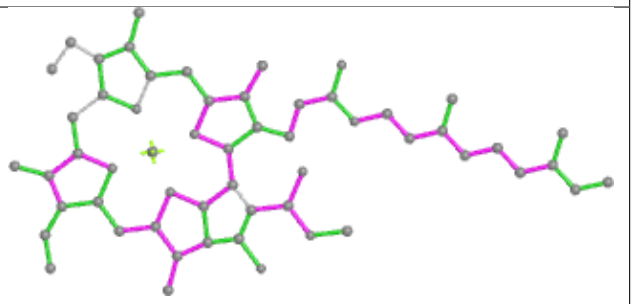


Rings

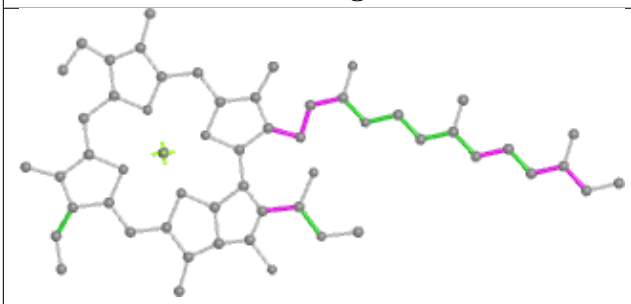
Ligand CLA B 810



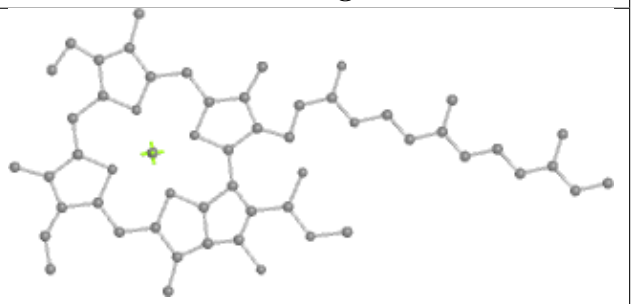
Bond lengths



Bond angles

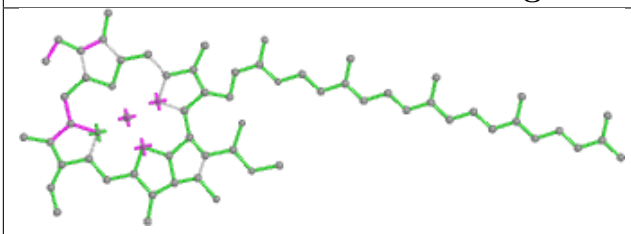


Torsions

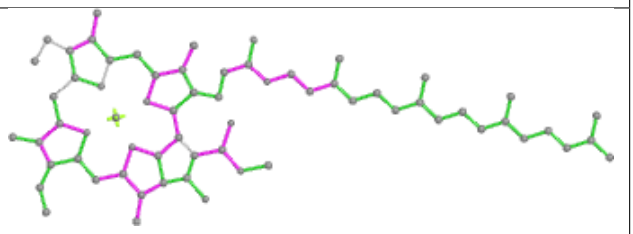


Rings

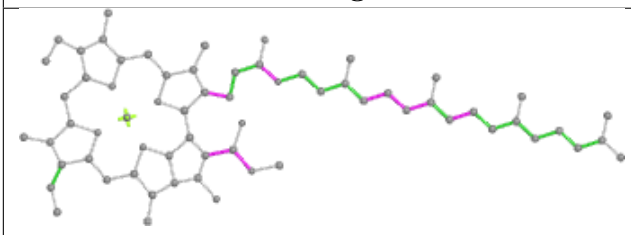
Ligand CLA A 819



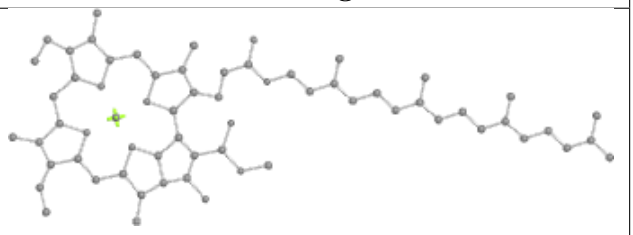
Bond lengths



Bond angles

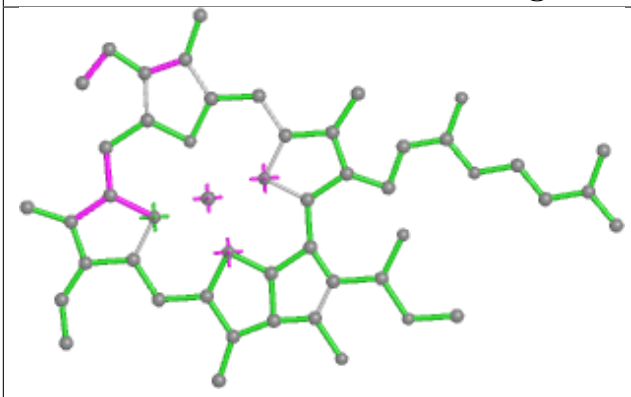


Torsions

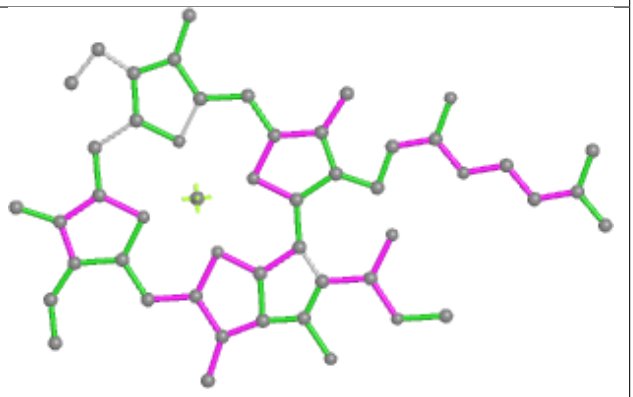


Rings

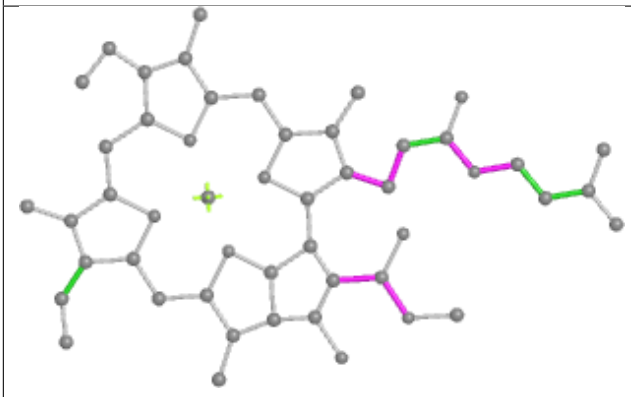
Ligand CLA 8 311



Bond lengths



Bond angles



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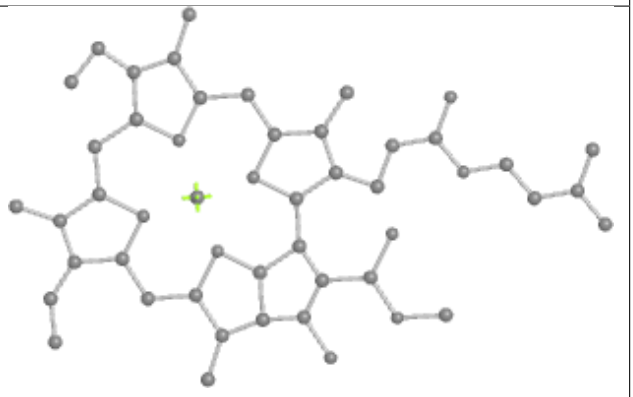
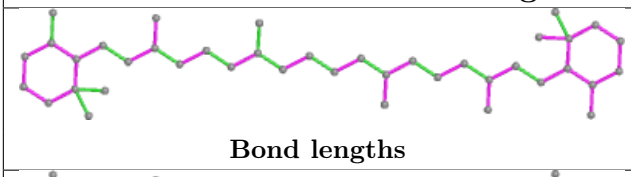
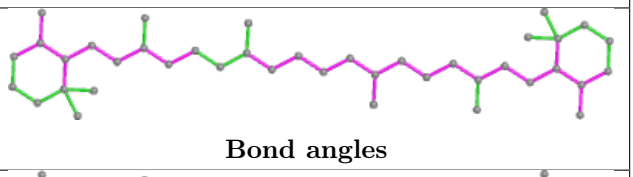


Rings

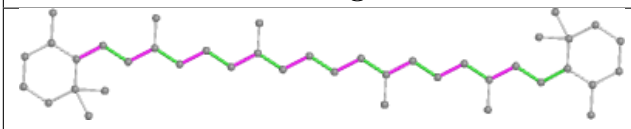
Ligand BCR I 4001



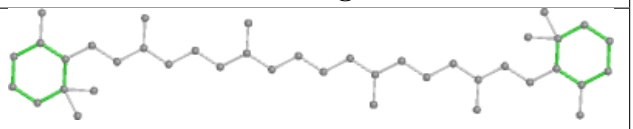
Bond lengths



Bond angles

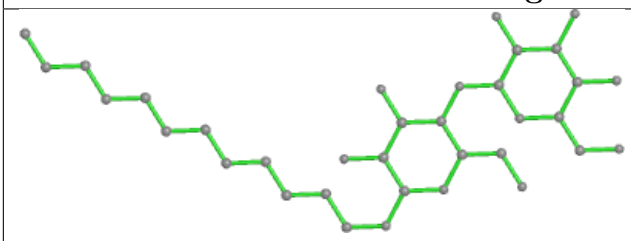


Torsions

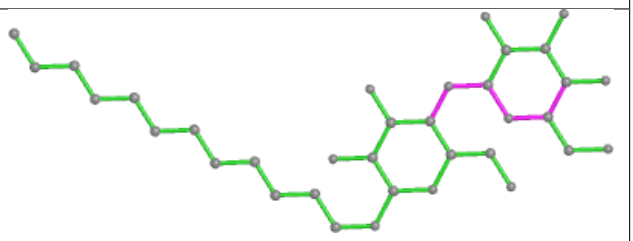


Rings

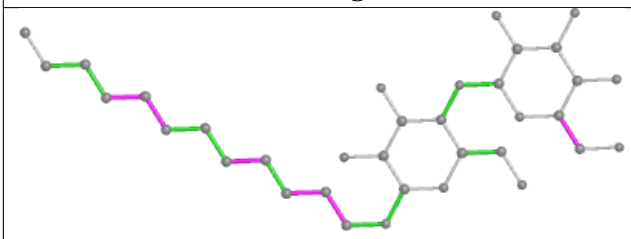
Ligand LMT 4 822



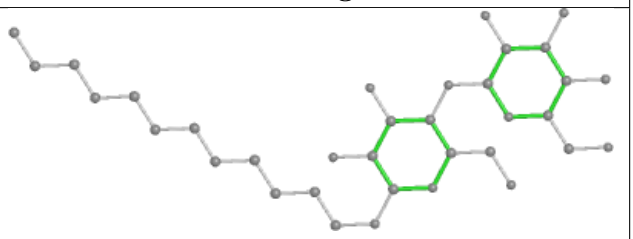
Bond lengths



Bond angles

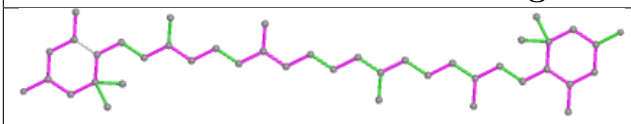


Torsions

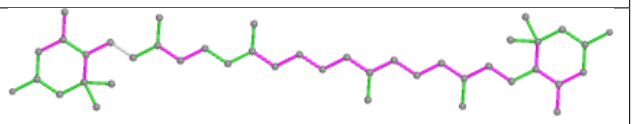


Rings

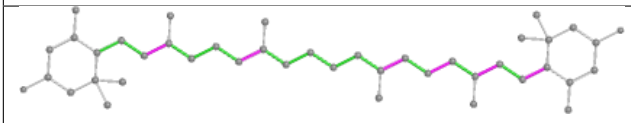
Ligand LUT 4 802



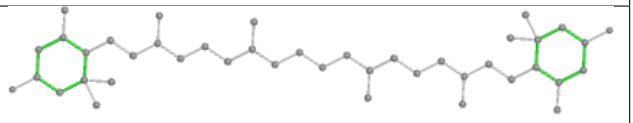
Bond lengths



Bond angles

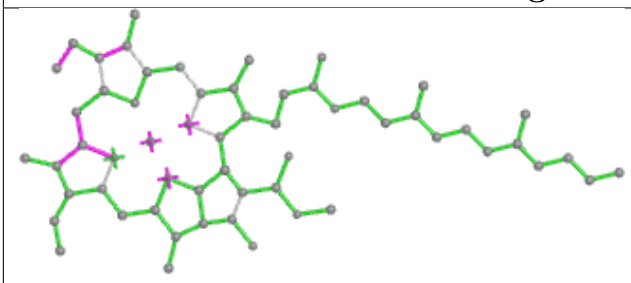


Torsions

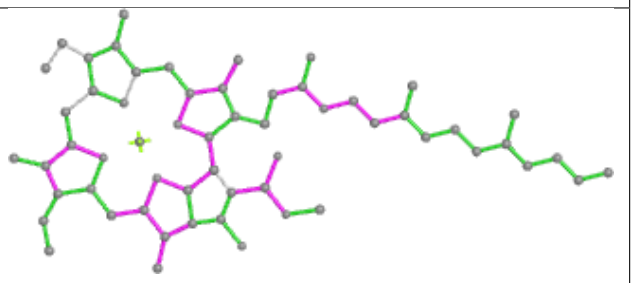


Rings

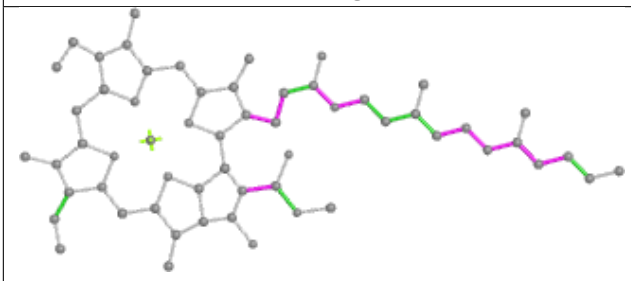
Ligand CLA 7 317



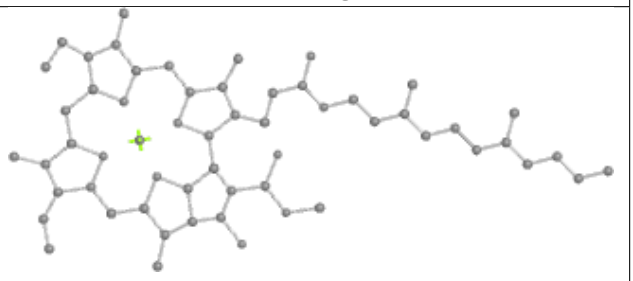
Bond lengths



Bond angles

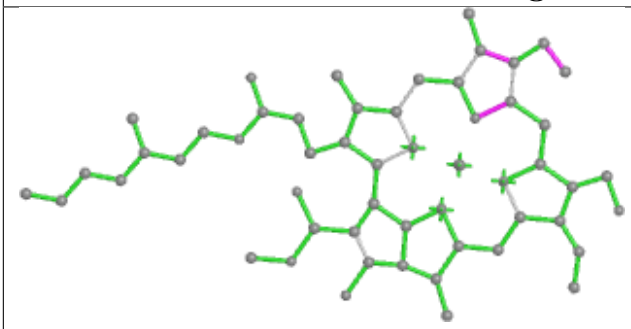


Torsions



Rings

Ligand CHL 7 313



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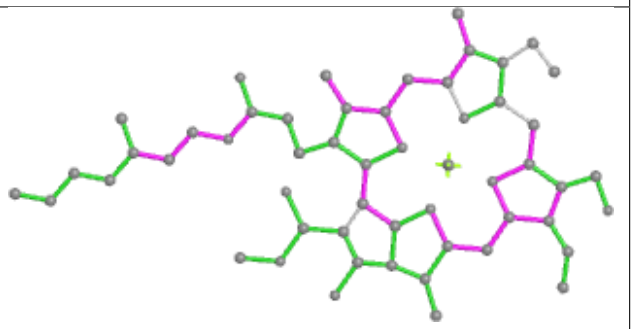
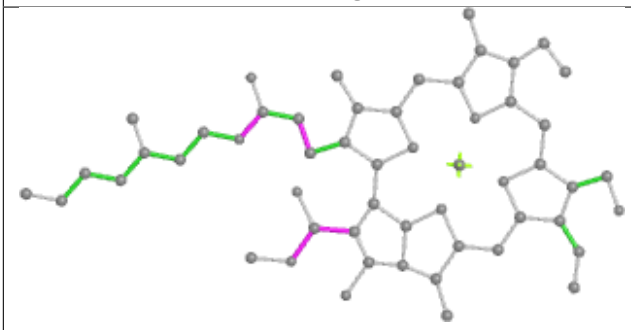


Bond angles



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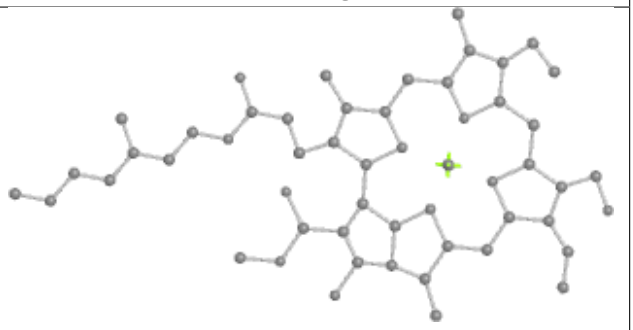
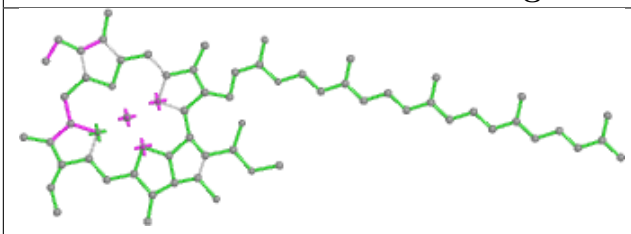
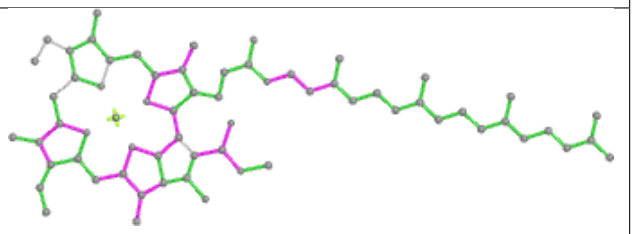


Rings

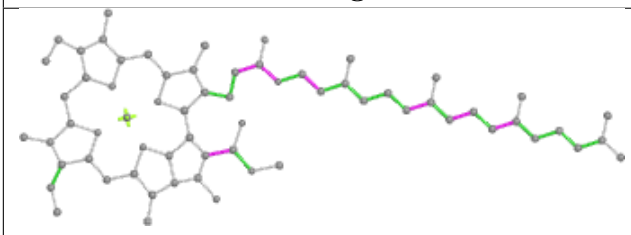
Ligand CLA A 829



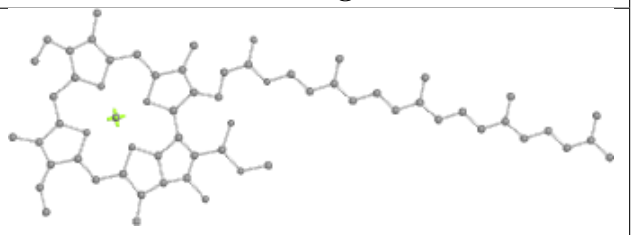
Bond lengths



Bond angles

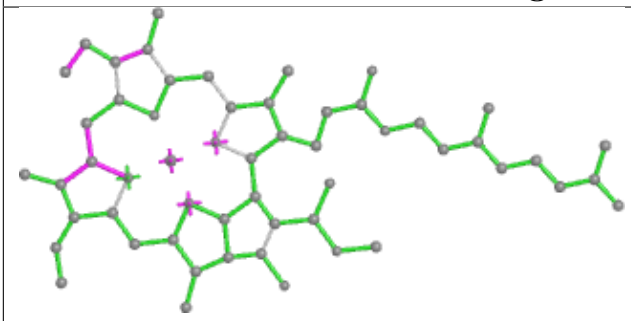


Torsions

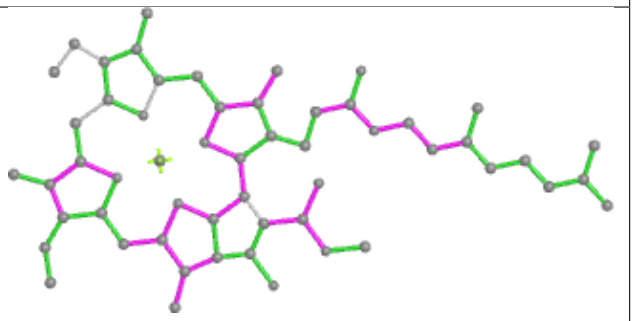


Rings

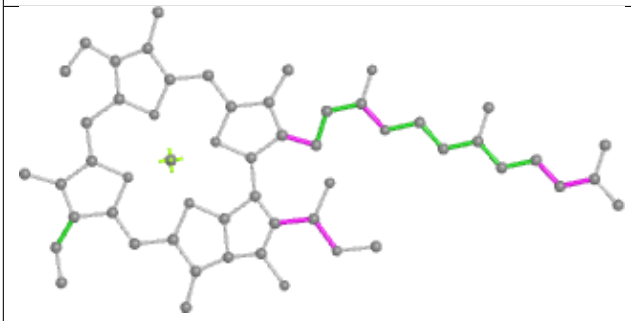
Ligand CLA 5 319



Bond lengths



Bond angles



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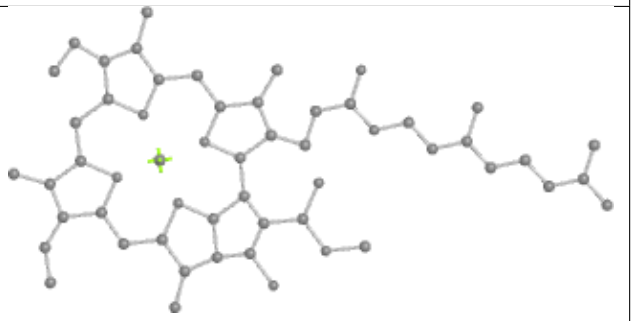
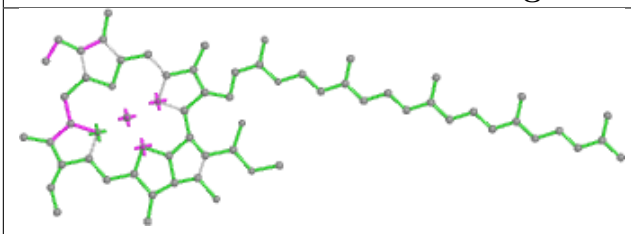
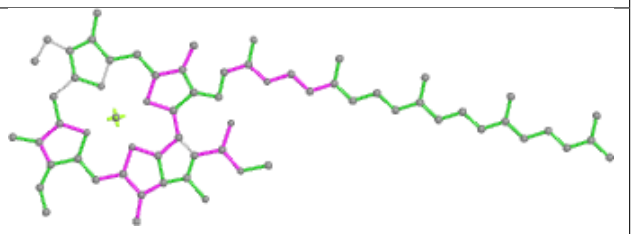


Rings

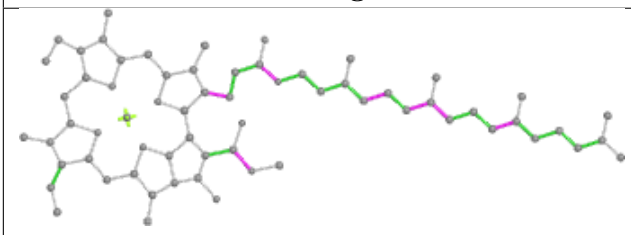
Ligand CLA A 803



Bond lengths



Bond angles



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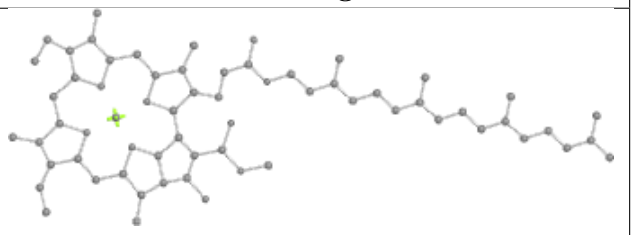
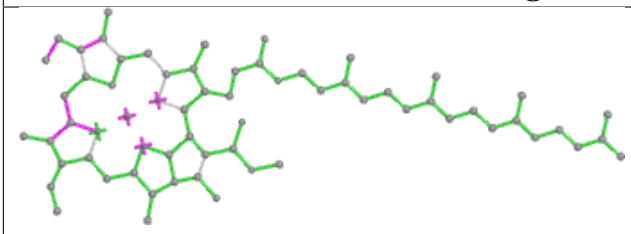
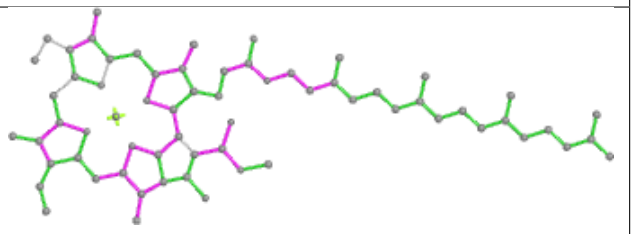


Rings

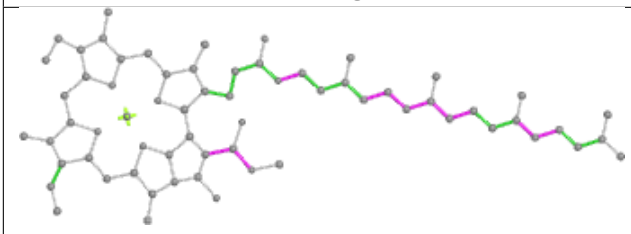
Ligand CLA B 801



Bond lengths



Bond angles



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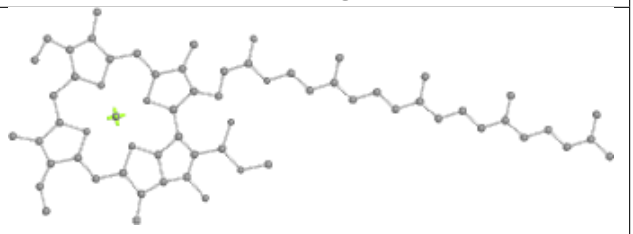

Rings

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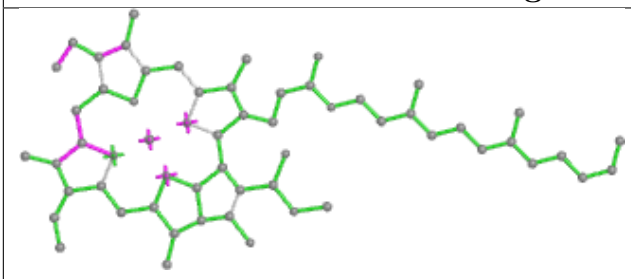
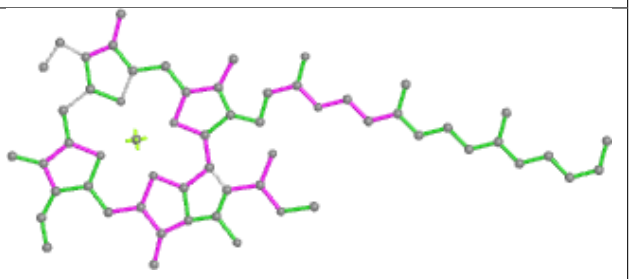
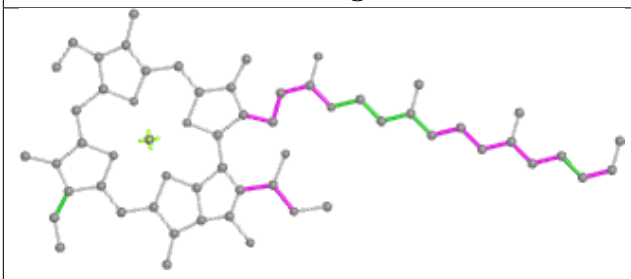


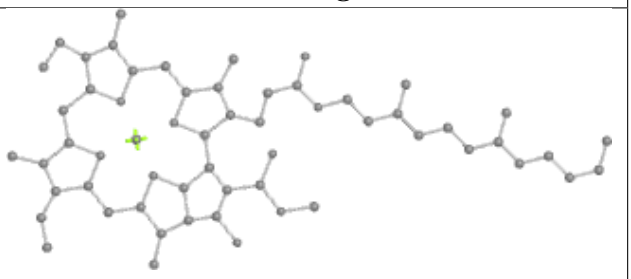
Bond lengths



Bond angles

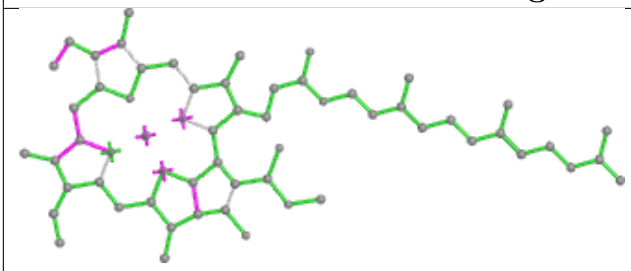


Torsions

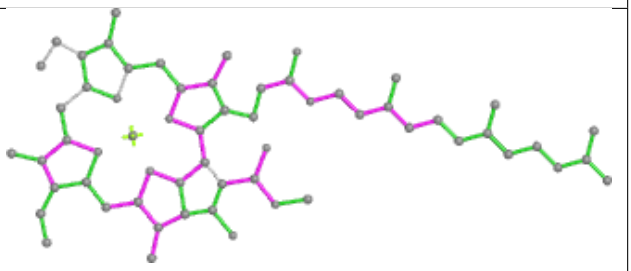


Rings

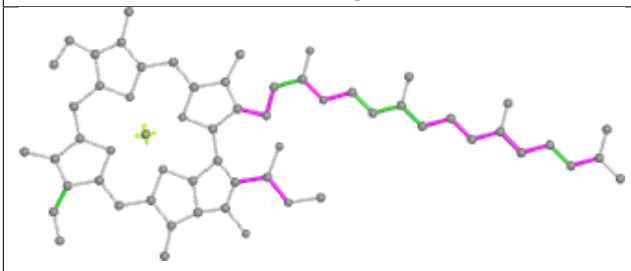
Ligand CLA 6 301



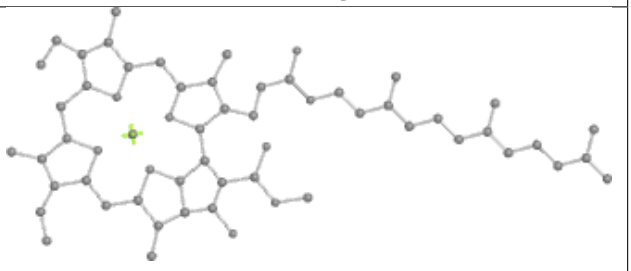
Bond lengths



Bond angles

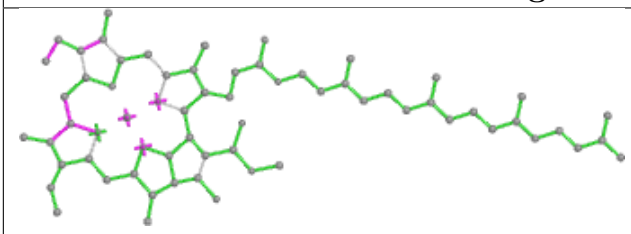


Torsions



Rings

Ligand CLA 5 310



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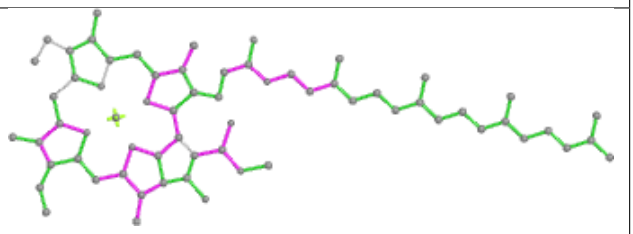
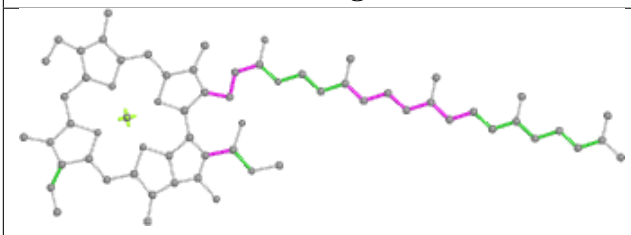
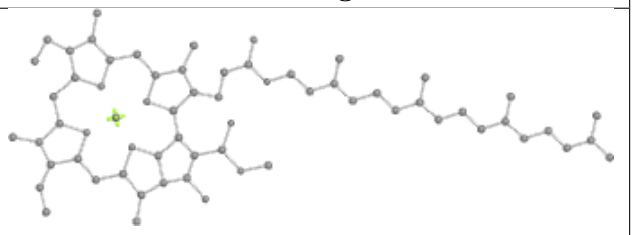


Bond angles

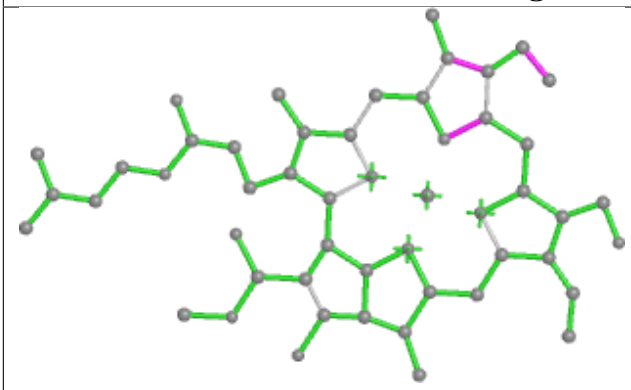


Torsions

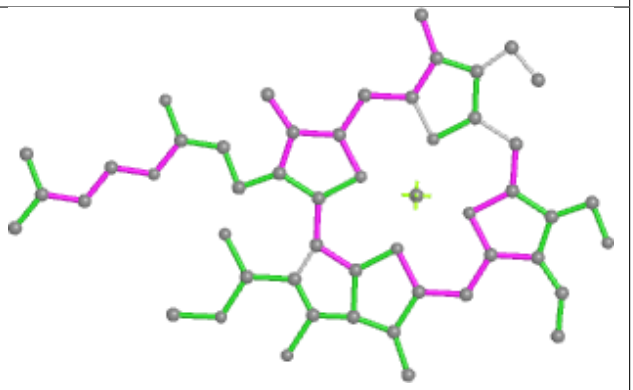


Rings

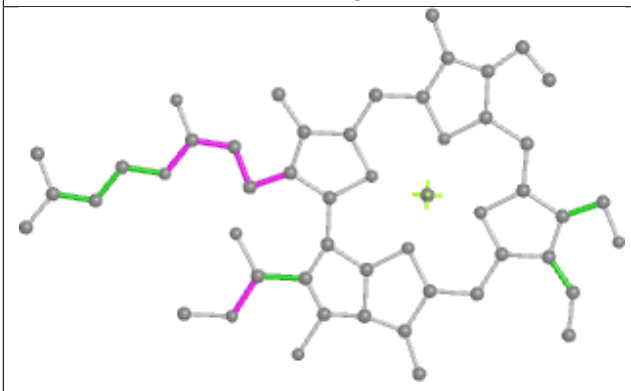
Ligand CHL 6 316



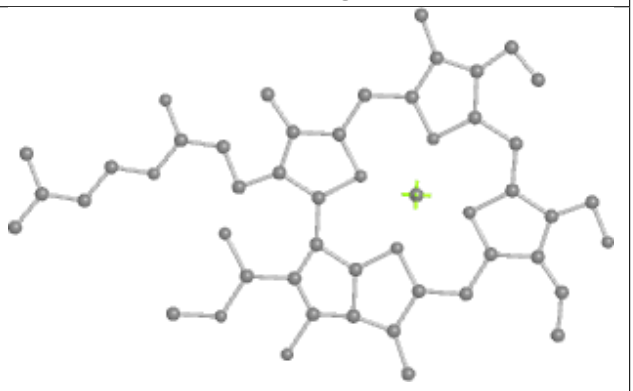
Bond lengths



Bond angles

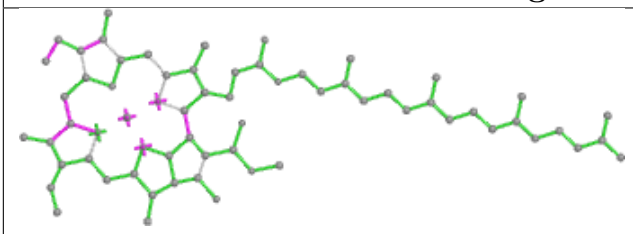


Torsions

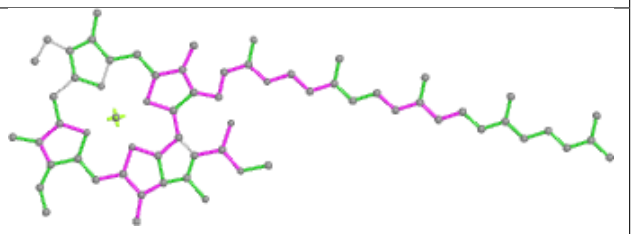


Rings

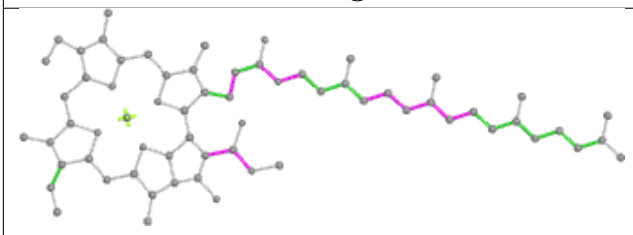
Ligand CLA 3 310



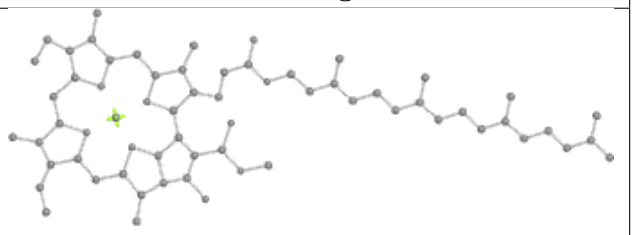
Bond lengths



Bond angles

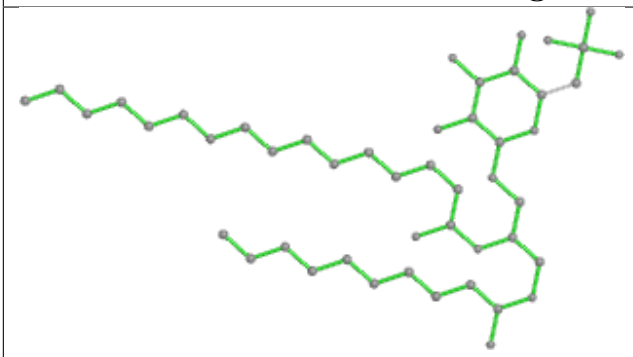


Torsions

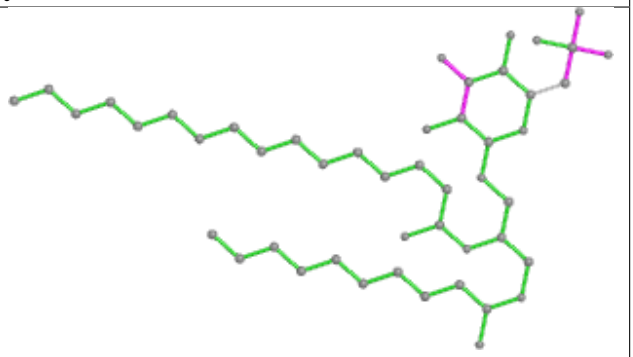


Rings

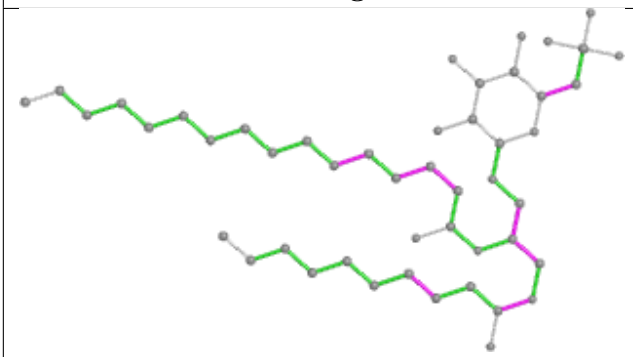
Ligand SQD 1 318



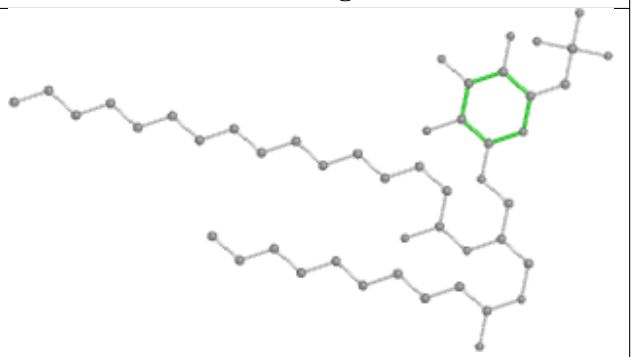
Bond lengths



Bond angles

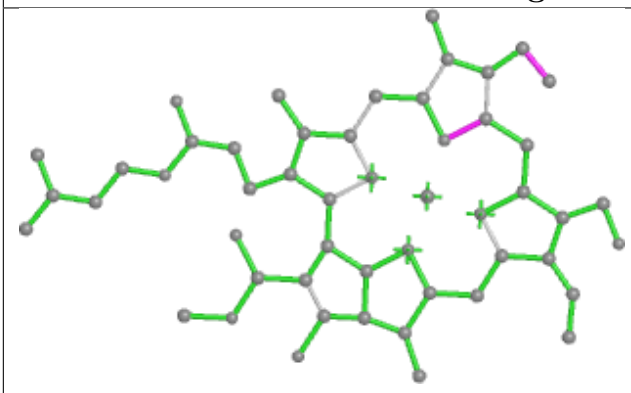


Torsions

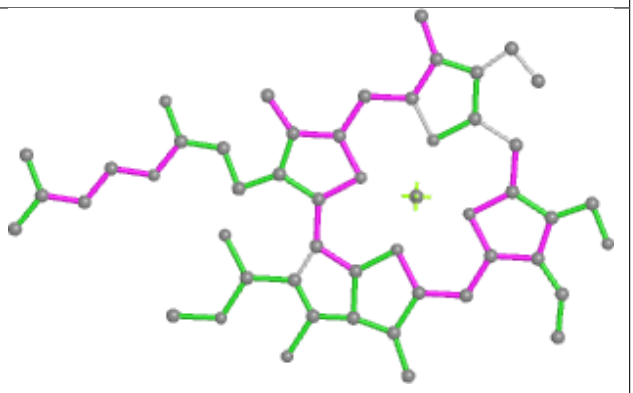


Rings

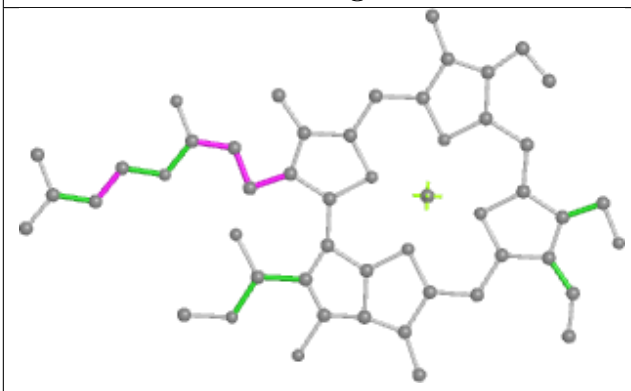
Ligand CHL 5 317



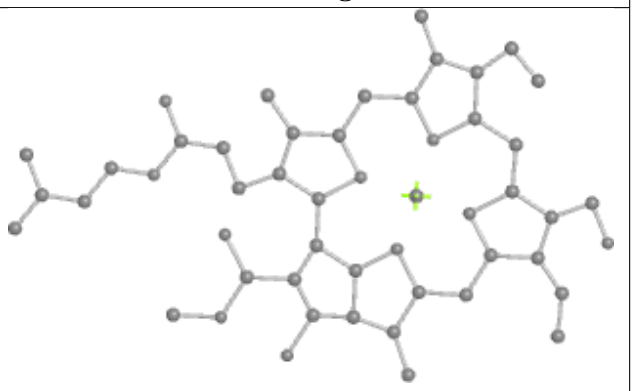
Bond lengths



Bond angles

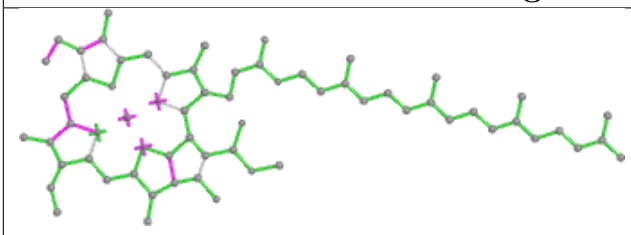


Torsions

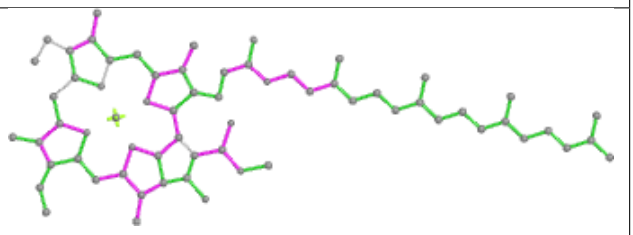


Rings

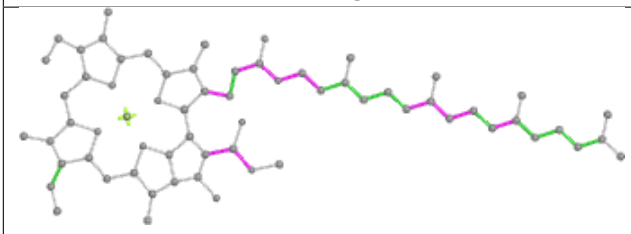
Ligand CLA A 810



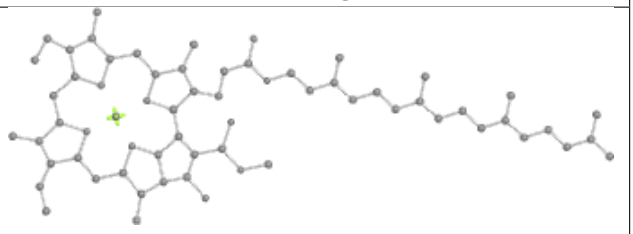
Bond lengths



Bond angles

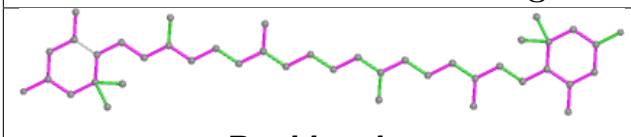


Torsions

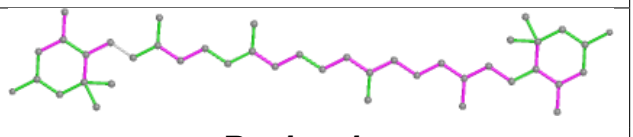


Rings

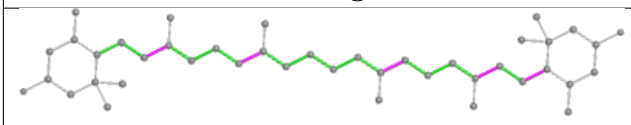
Ligand LUT 1 302



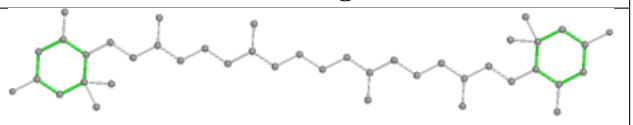
Bond lengths



Bond angles

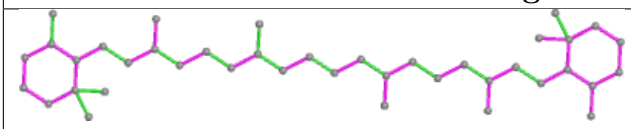


Torsions

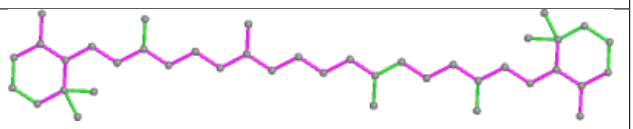


Rings

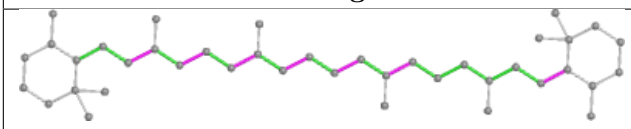
Ligand BCR B 847



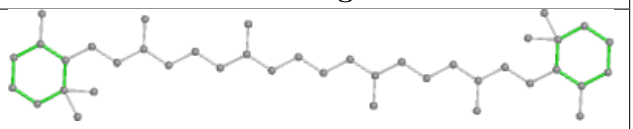
Bond lengths



Bond angles

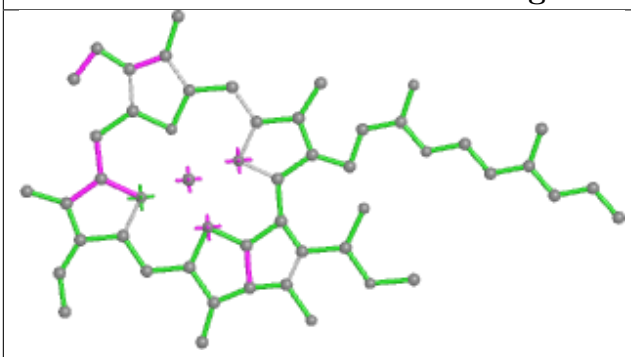


Torsions

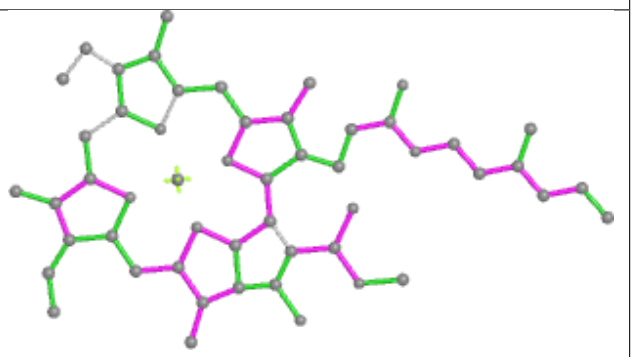


Rings

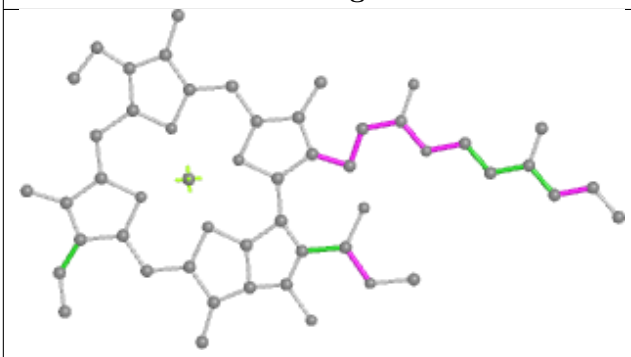
Ligand CLA 4 806



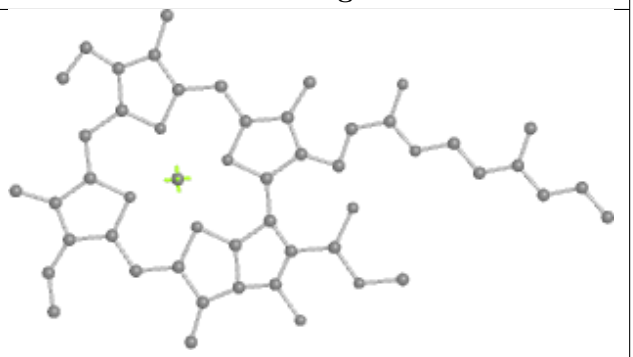
Bond lengths



Bond angles

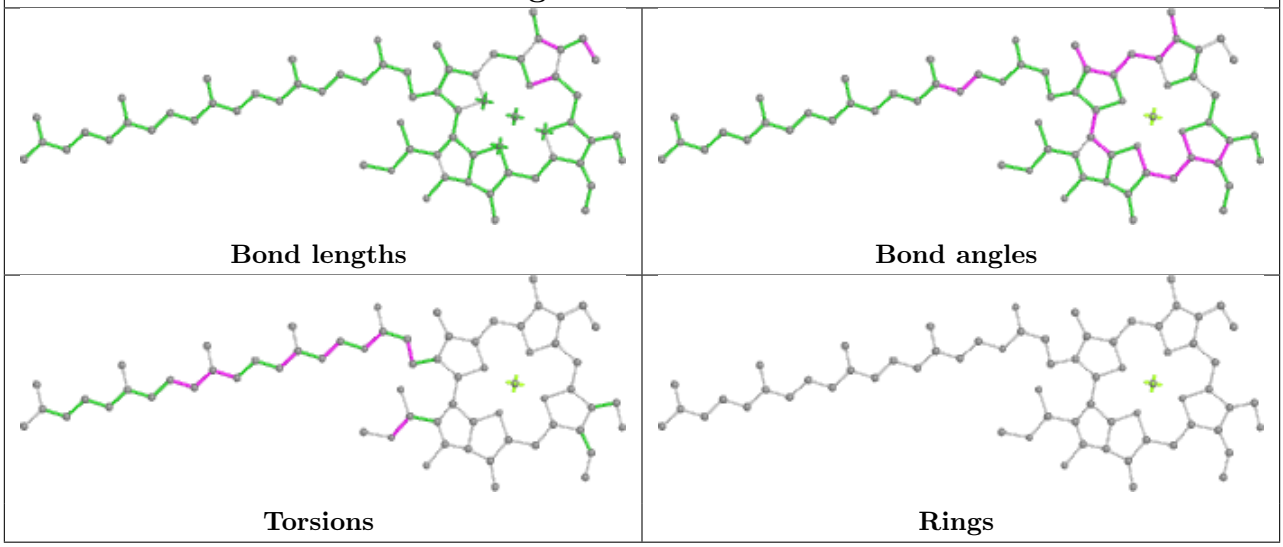


Torsions

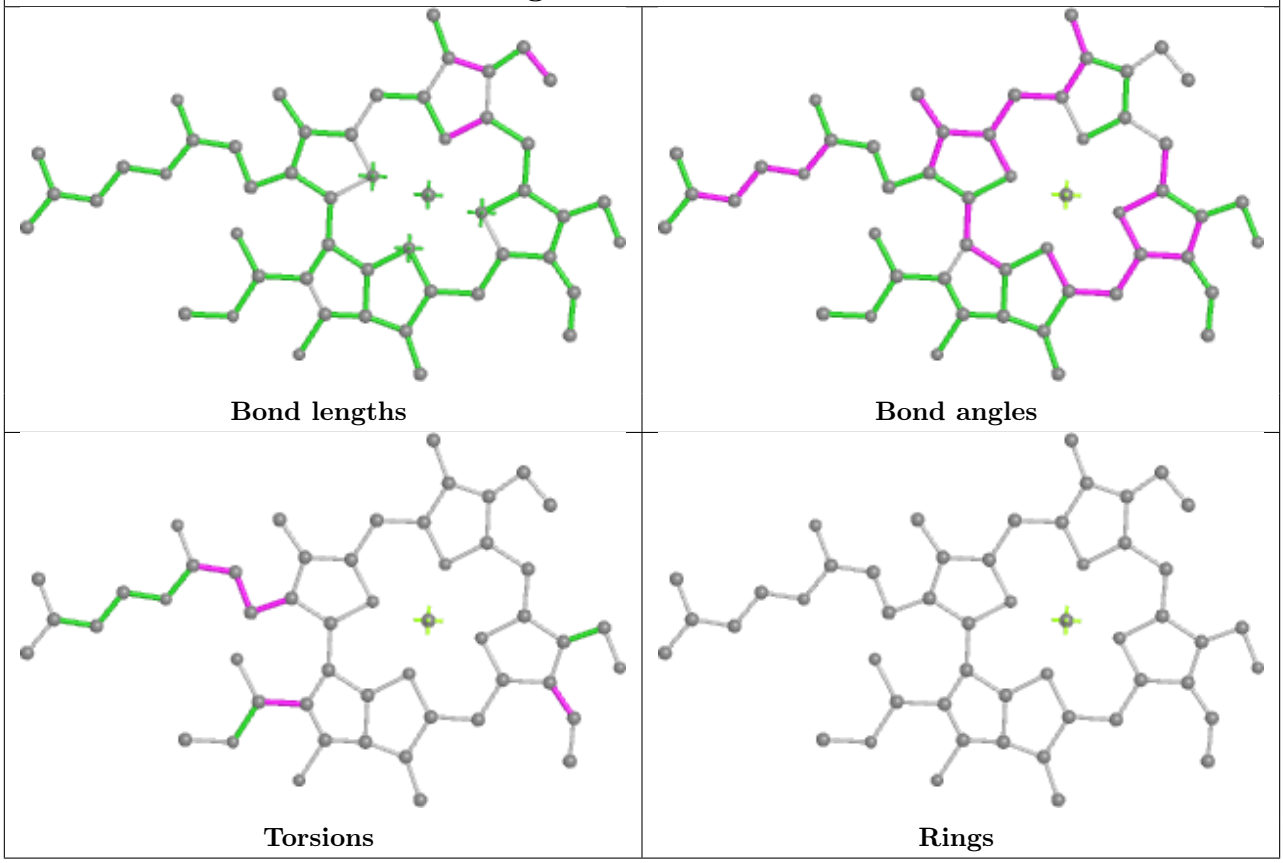


Rings

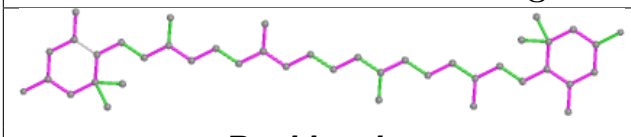
Ligand CHL 3 317



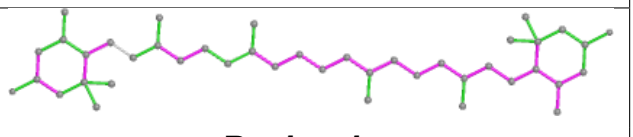
Ligand CHL Z 311



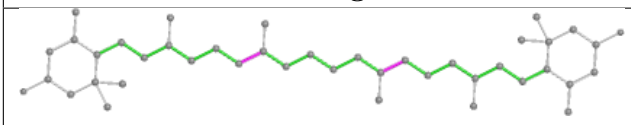
Ligand LUT 3 302



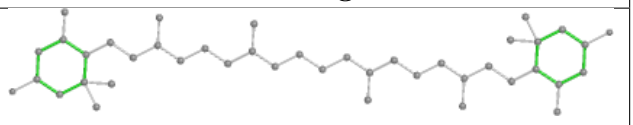
Bond lengths



Bond angles

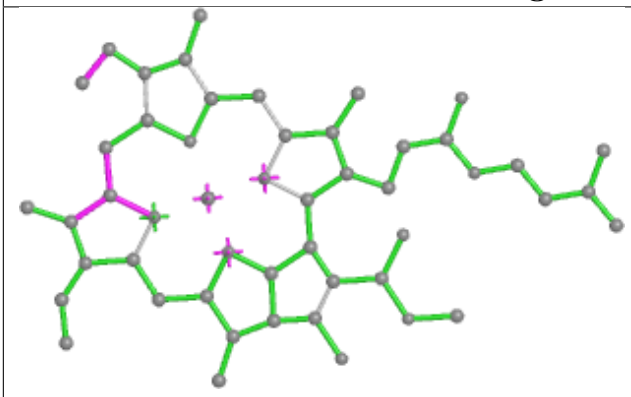


Torsions

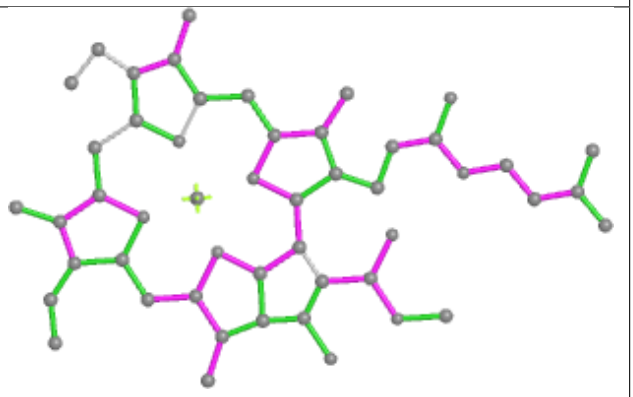


Rings

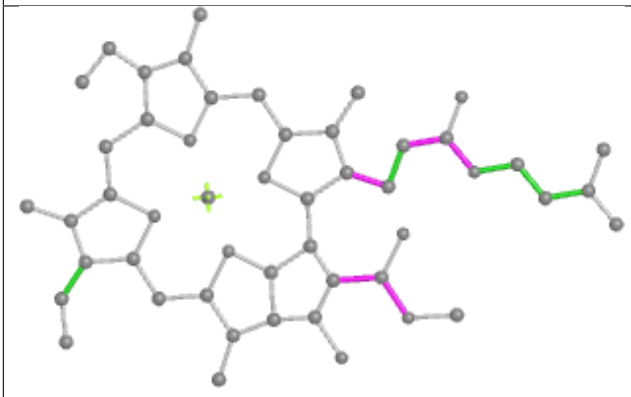
Ligand CLA 6 317



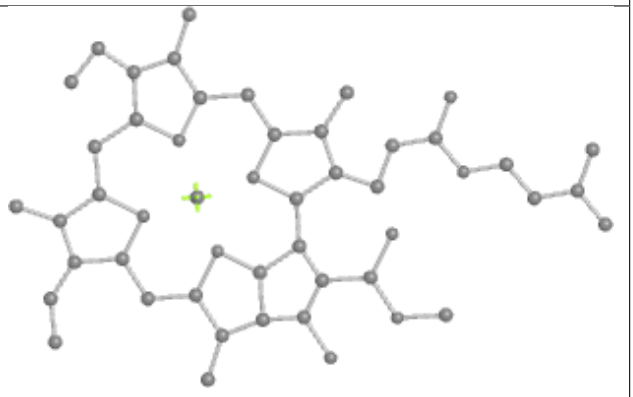
Bond lengths



Bond angles

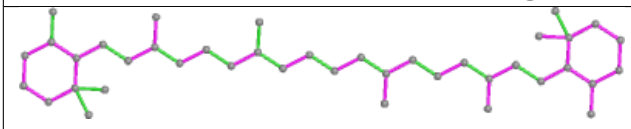


Torsions

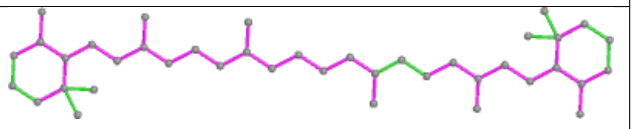


Rings

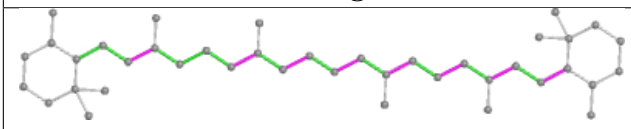
Ligand BCR K 206



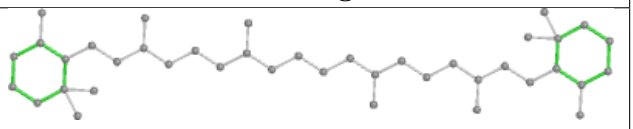
Bond lengths



Bond angles

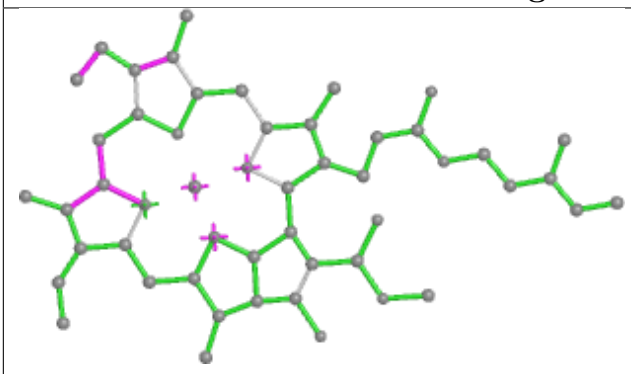


Torsions

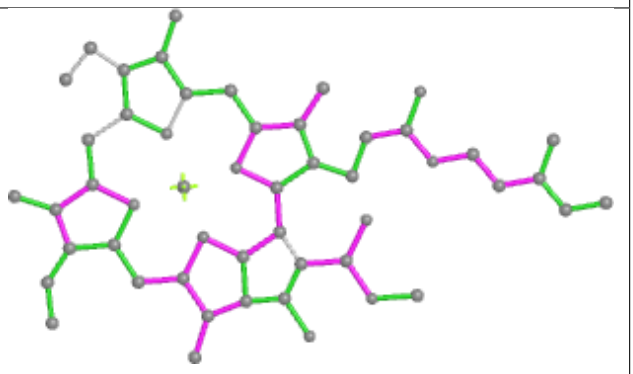


Rings

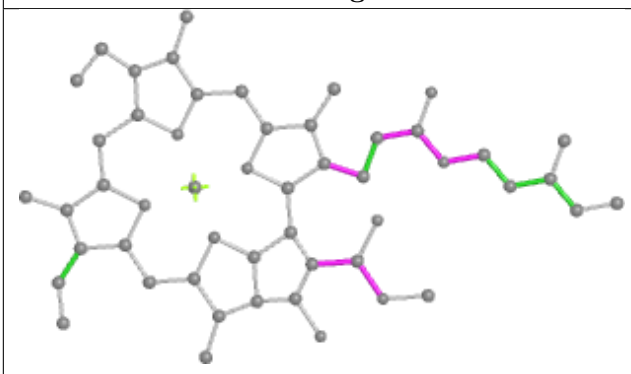
Ligand CLA 4 818



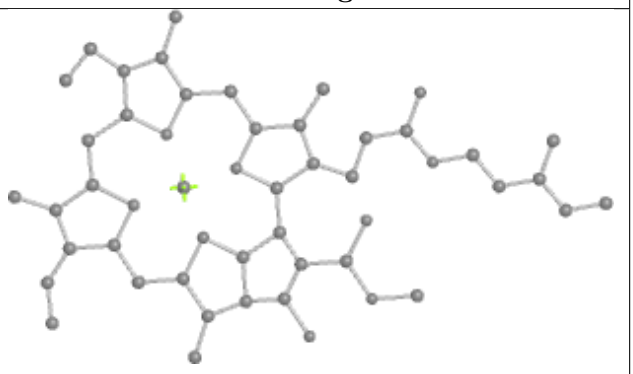
Bond lengths



Bond angles

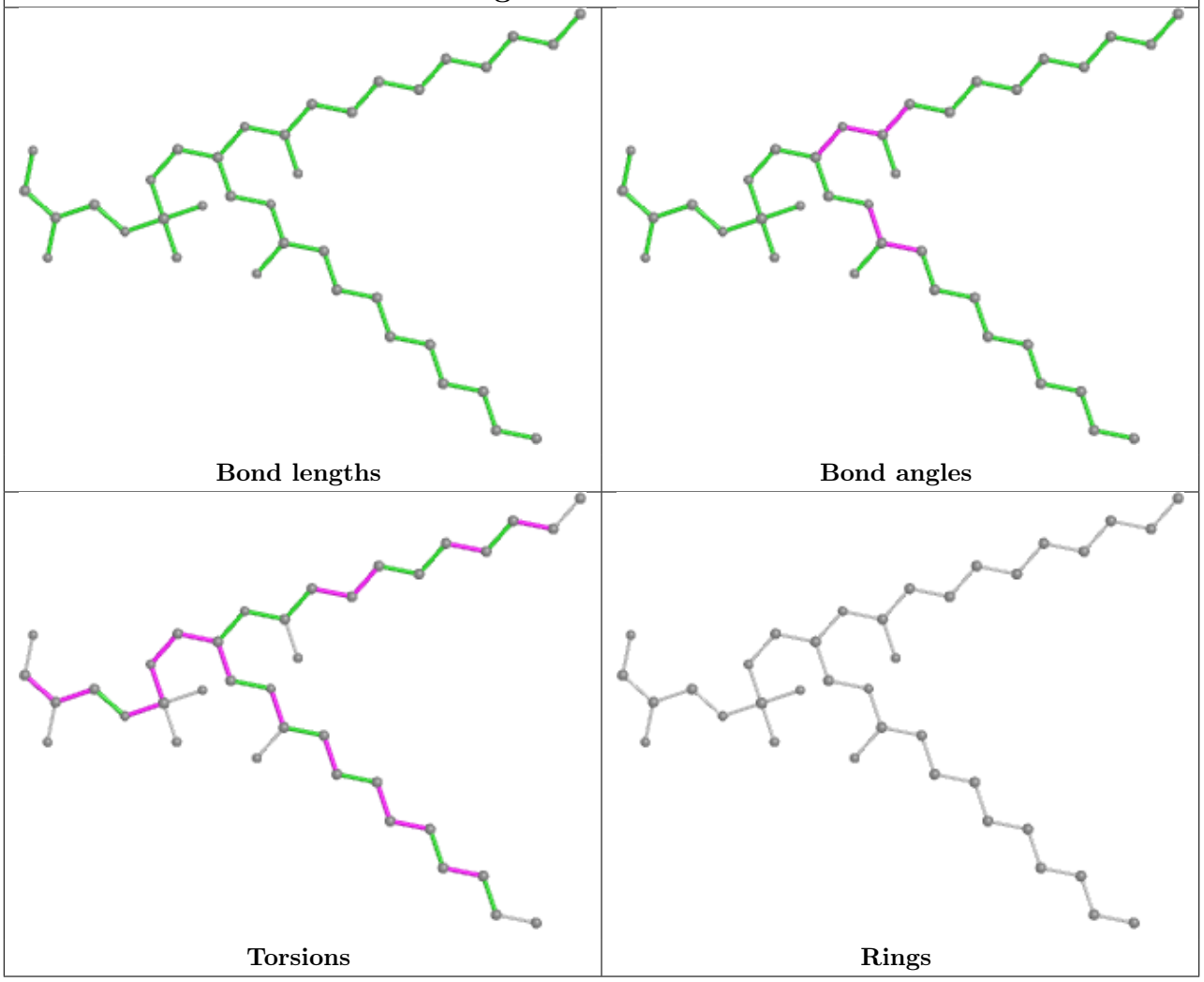


Torsions

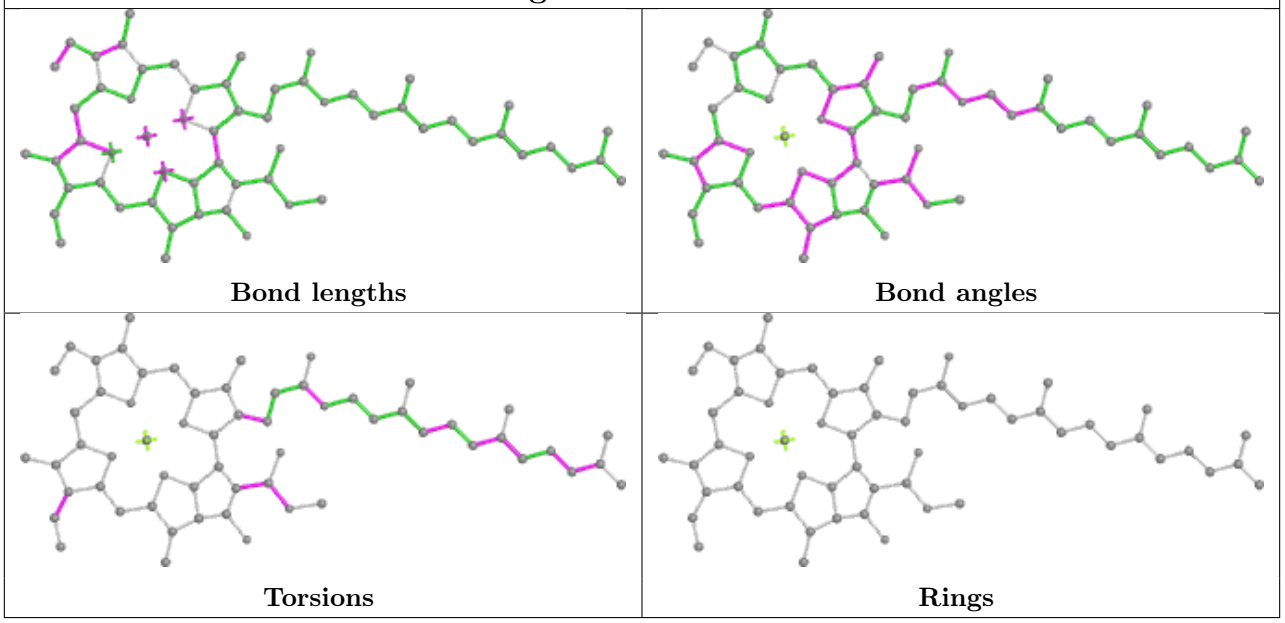


Rings

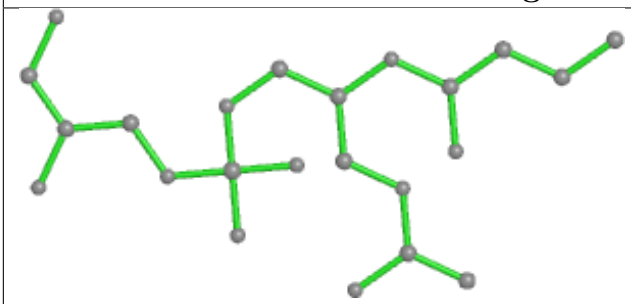
Ligand LHG 5 324



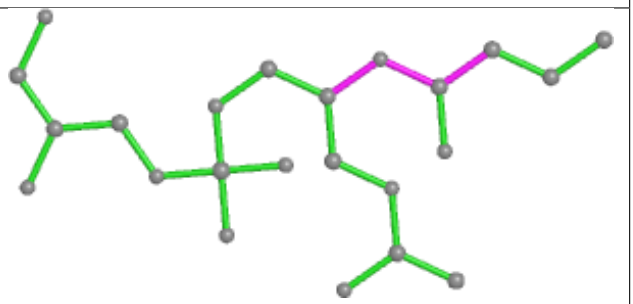
Ligand CLA 1 310



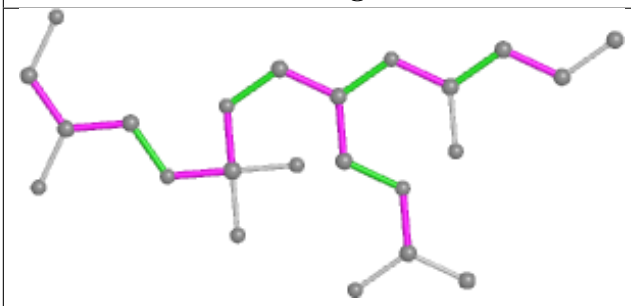
Ligand LHG B 850



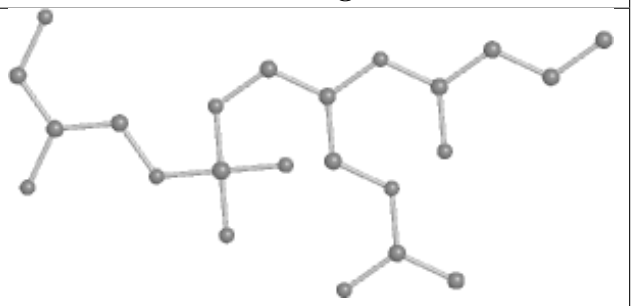
Bond lengths



Bond angles

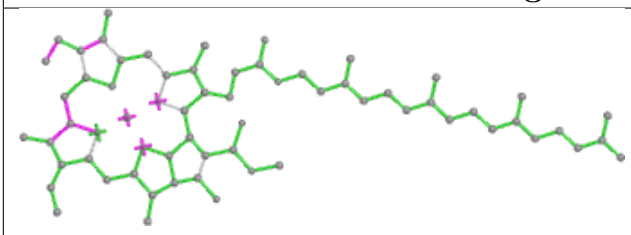


Torsions

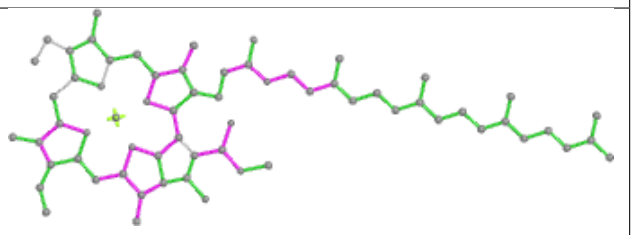


Rings

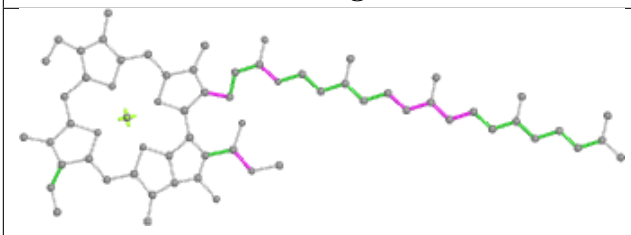
Ligand CLA B 841



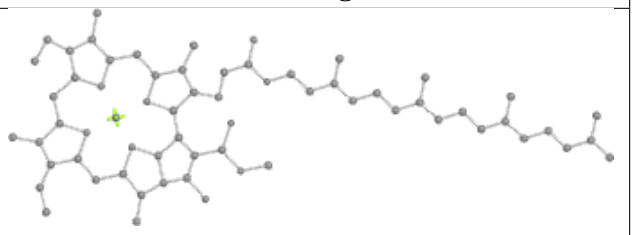
Bond lengths



Bond angles

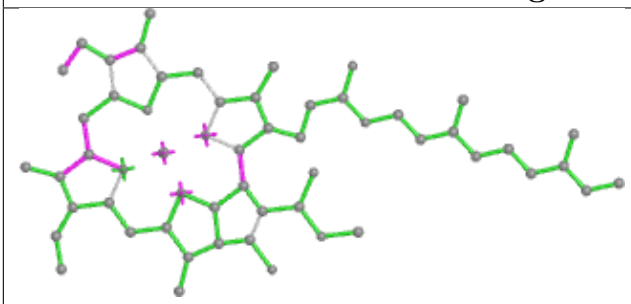


Torsions

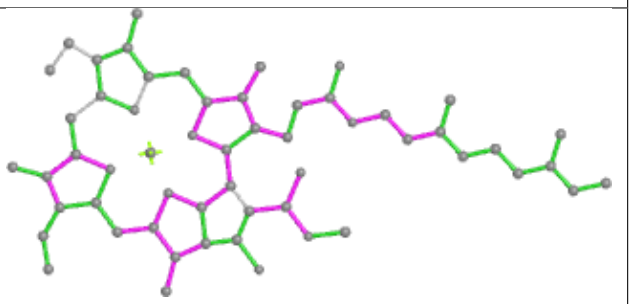


Rings

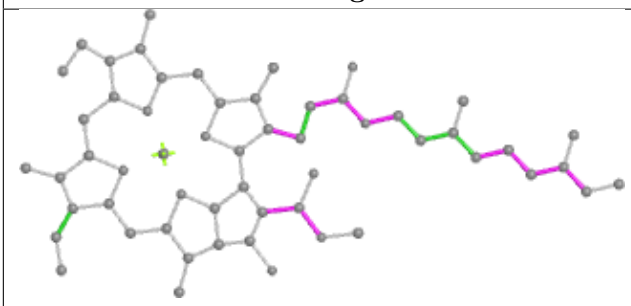
Ligand CLA 5 309



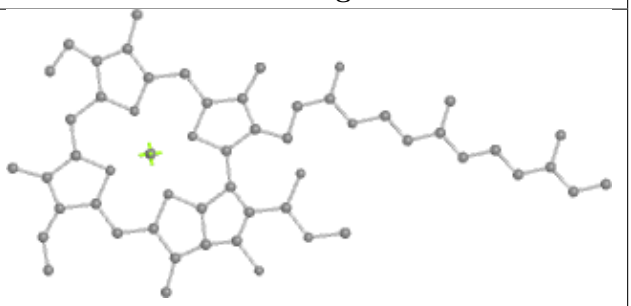
Bond lengths



Bond angles

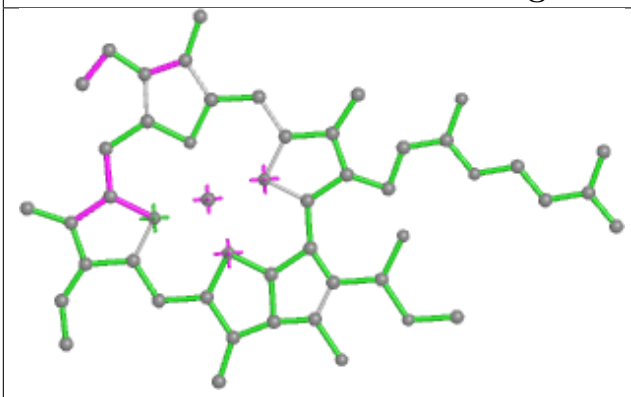


Torsions

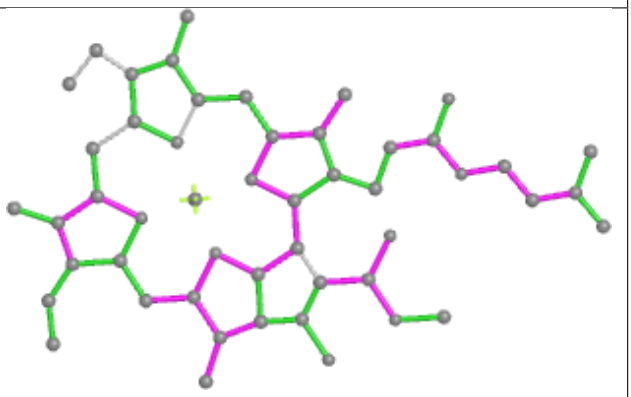


Rings

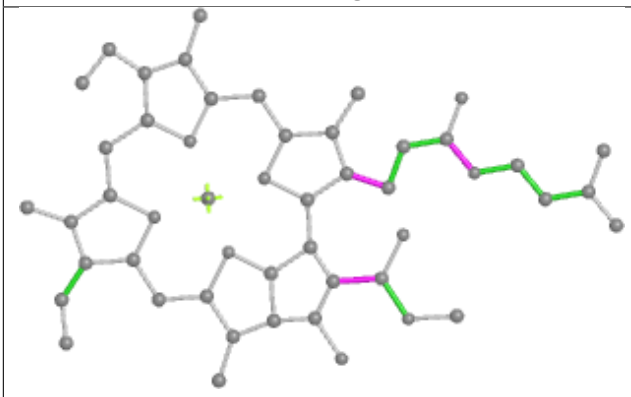
Ligand CLA 5 312



Bond lengths



Bond angles



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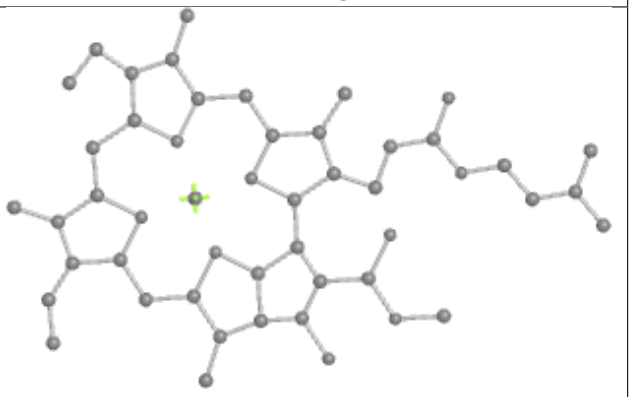
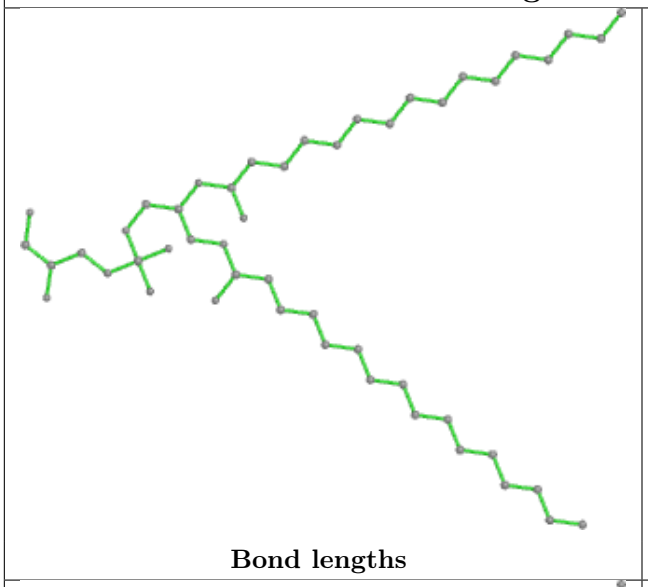
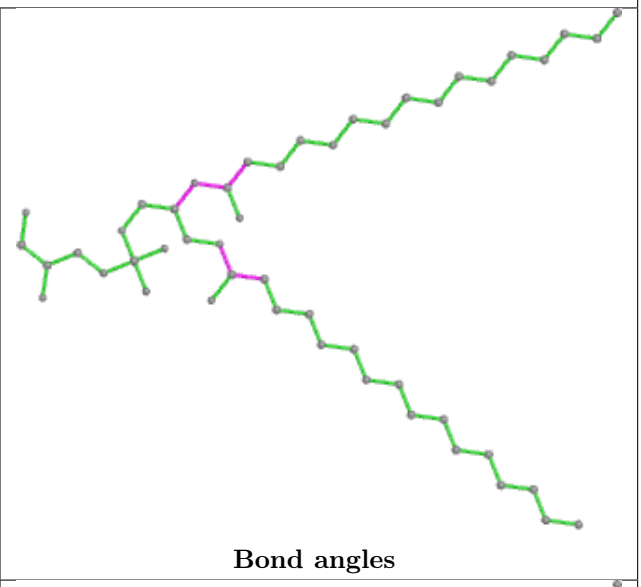


Rings

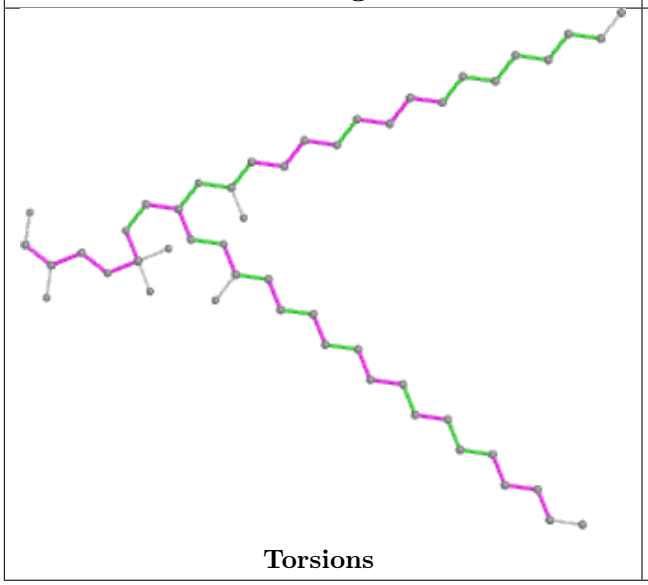
Ligand LHG A 850



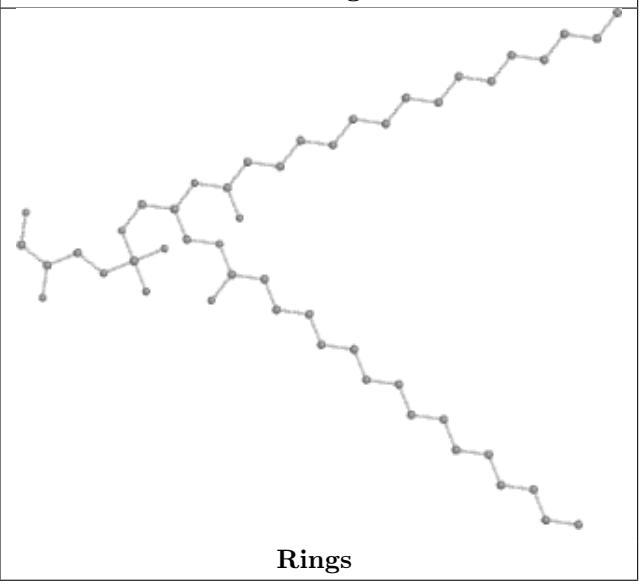
Bond lengths



Bond angles

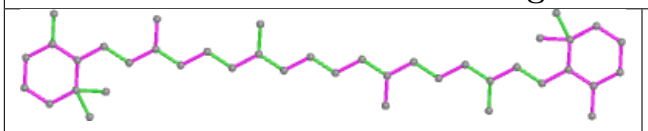


Torsions

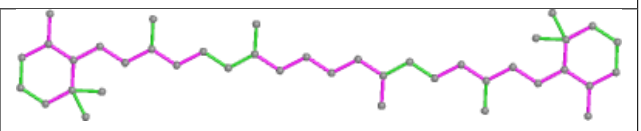


Rings

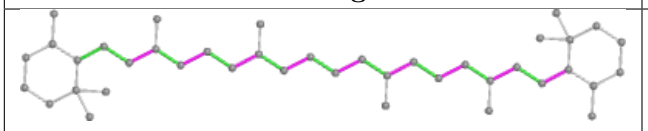
Ligand BCR 4 804



Bond lengths



Bond angles



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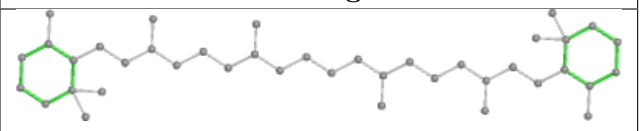
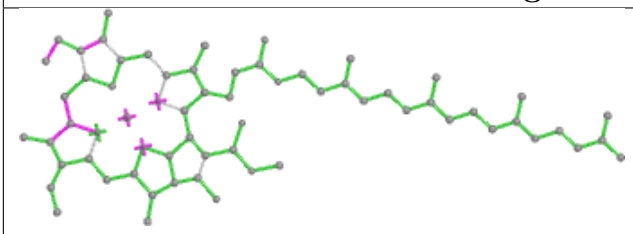


Rings

Ligand CLA A 841



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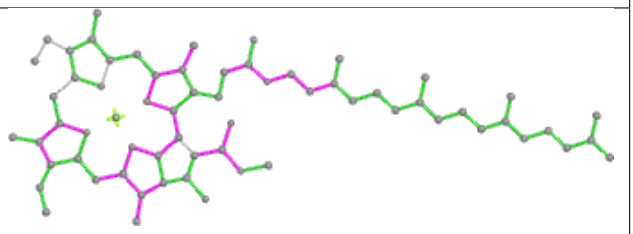
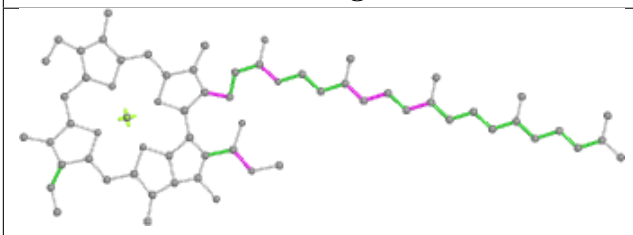


Bond angles



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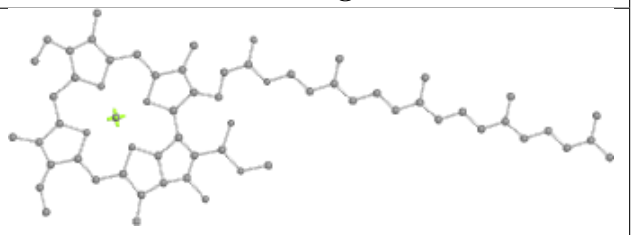
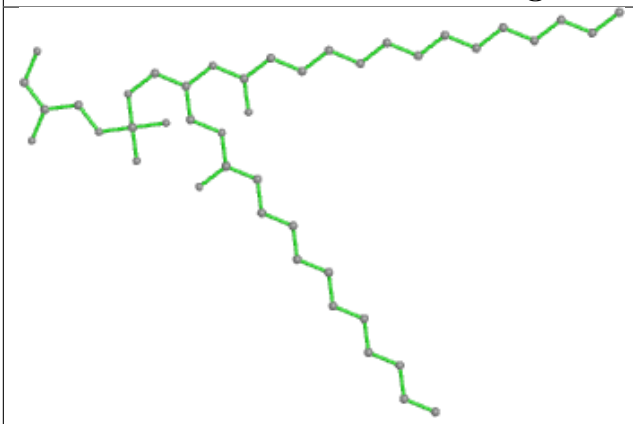


Rings

Ligand LHG 1 317



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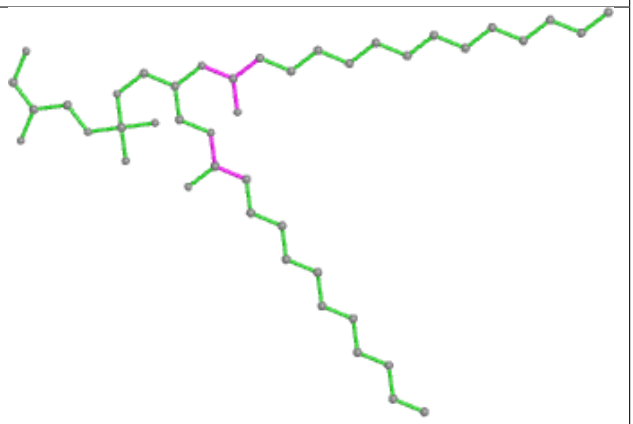
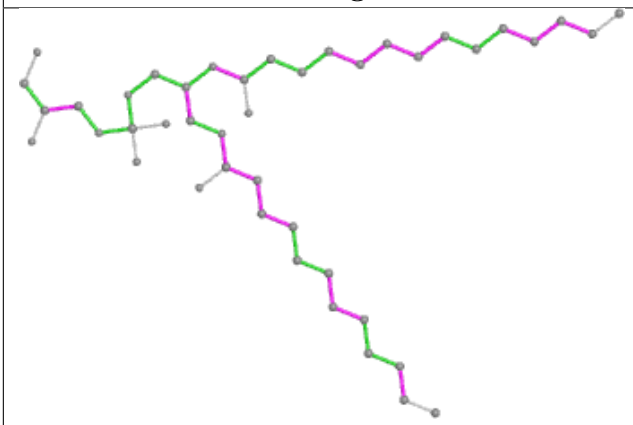


Bond angles



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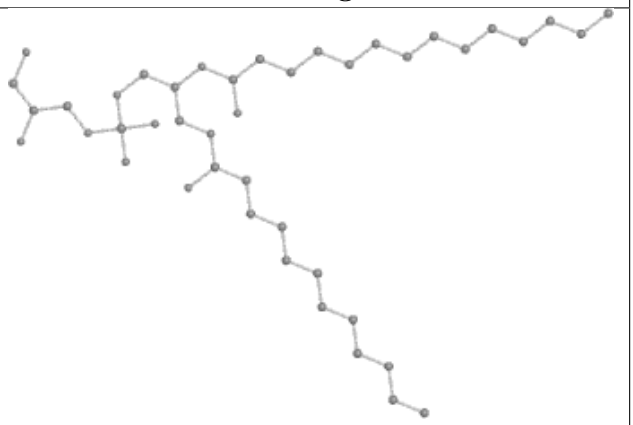
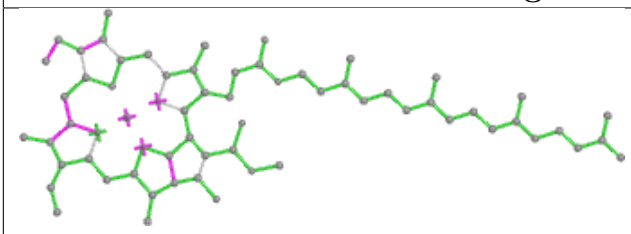


Rings

Ligand CLA B 809



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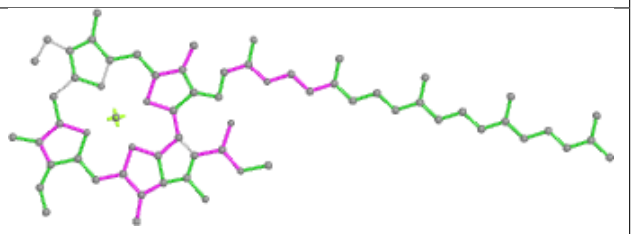
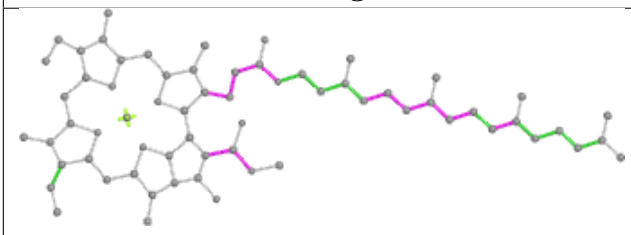
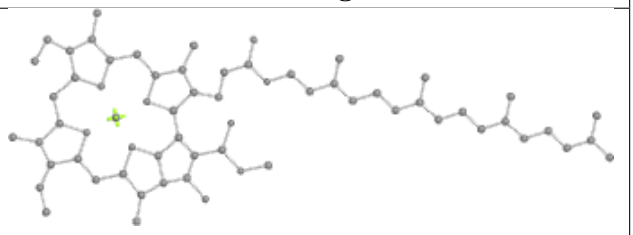


Bond angles

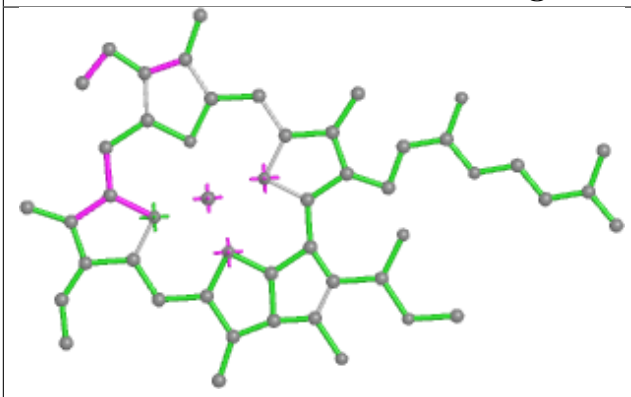


Torsions

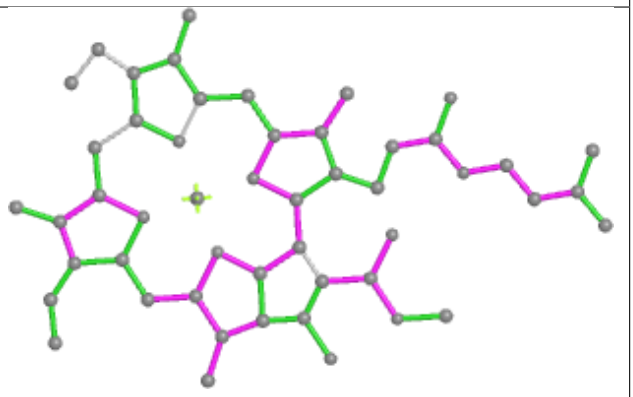


Rings

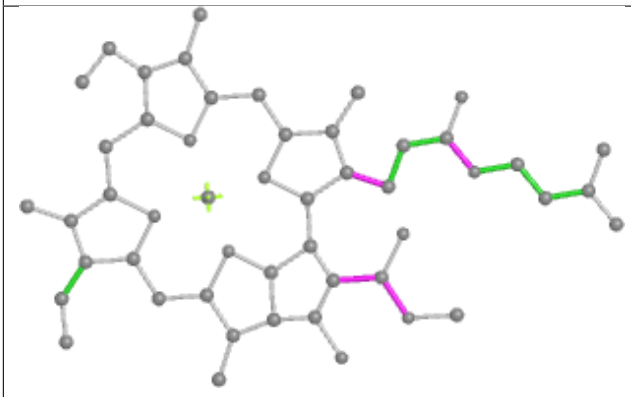
Ligand CLA 8 313



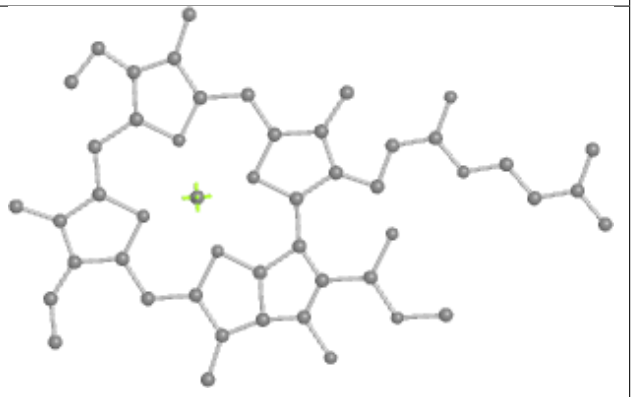
Bond lengths



Bond angles

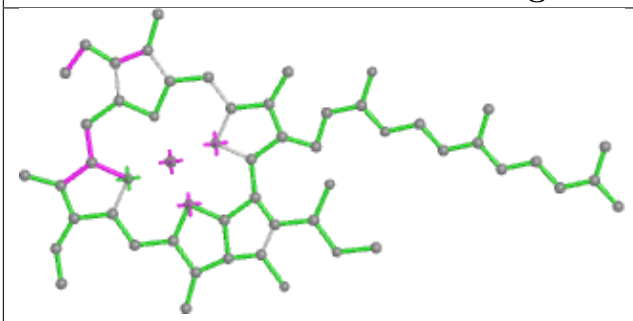


Torsions

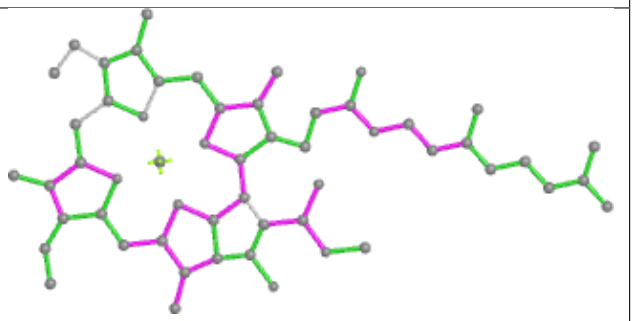


Rings

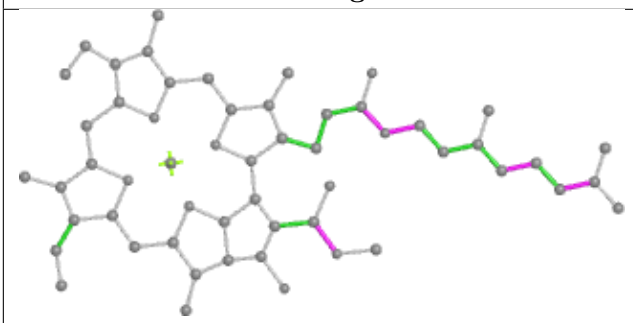
Ligand CLA 4 812



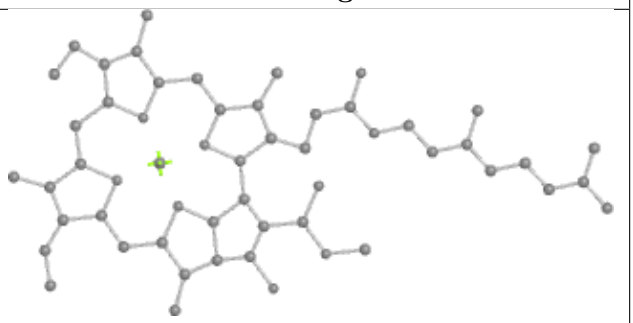
Bond lengths



Bond angles

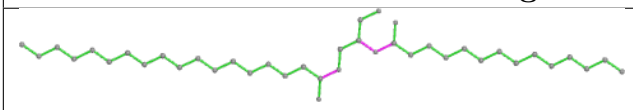


Torsions

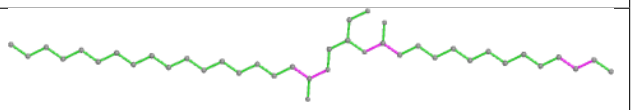


Rings

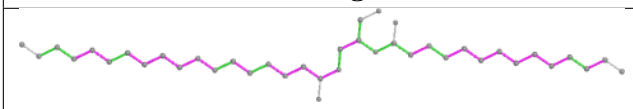
Ligand DGA A 852



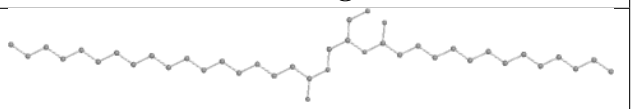
Bond lengths



Bond angles

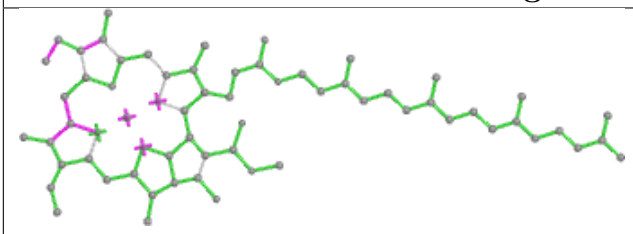


Torsions

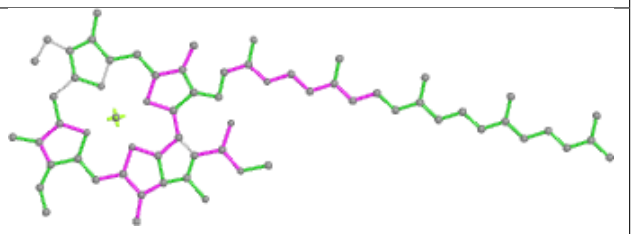


Rings

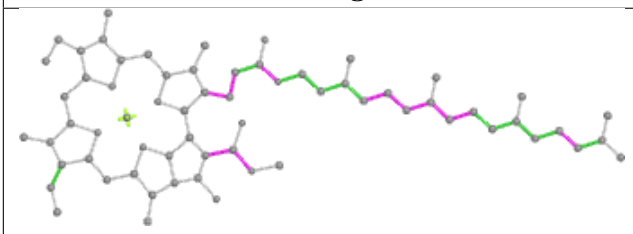
Ligand CLA A 824



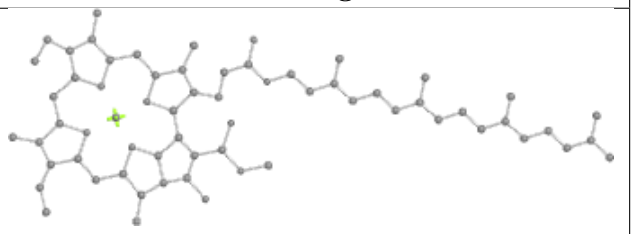
Bond lengths



Bond angles

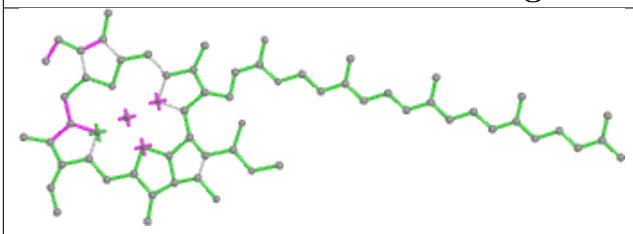


Torsions

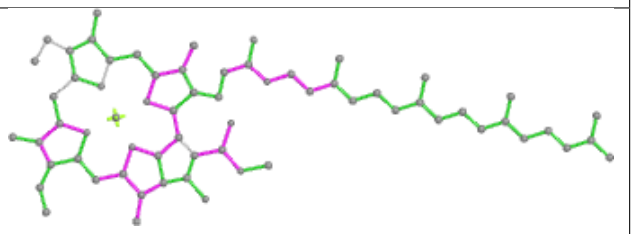


Rings

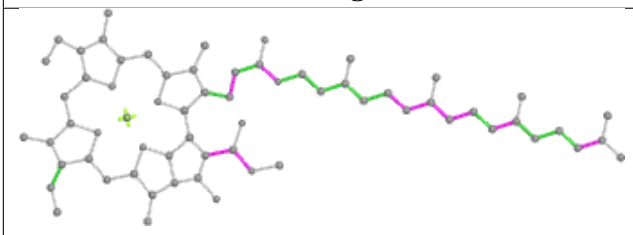
Ligand CLA B 805



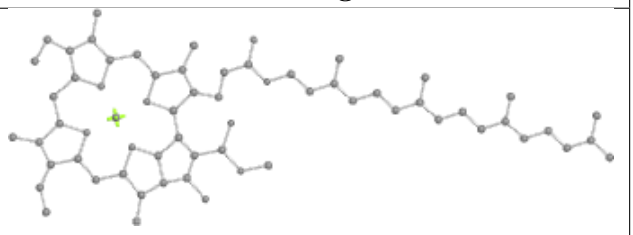
Bond lengths



Bond angles

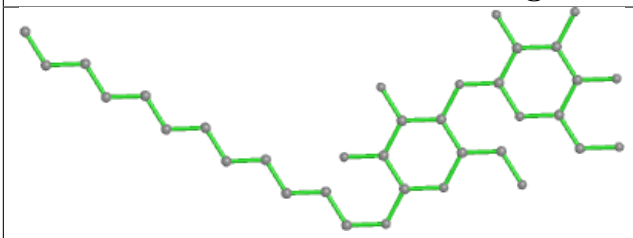


Torsions

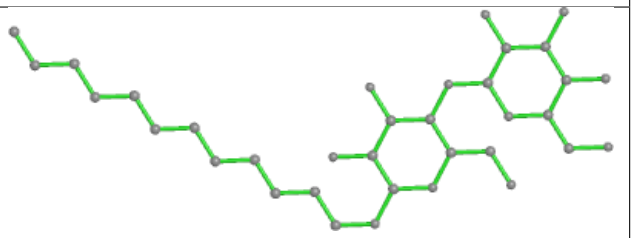


Rings

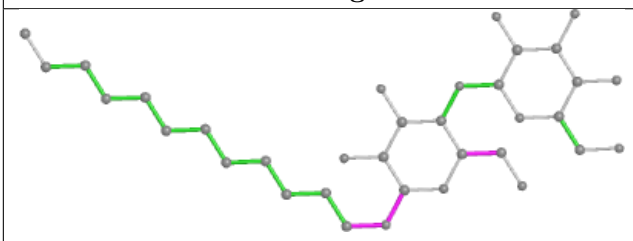
Ligand LMT 1 301



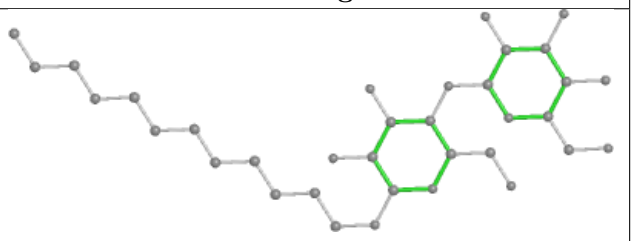
Bond lengths



Bond angles

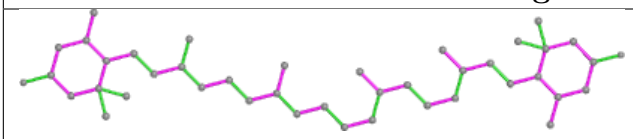


Torsions

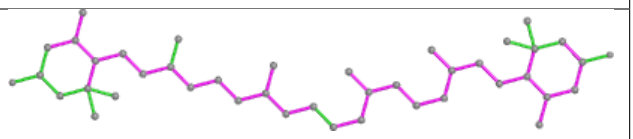


Rings

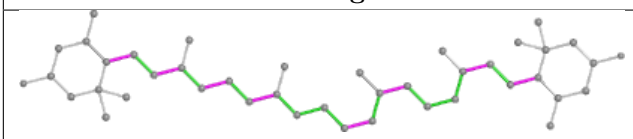
Ligand C7Z 5 306



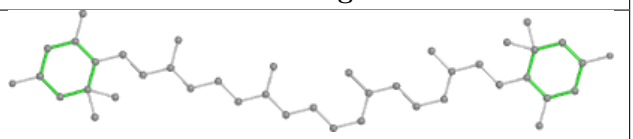
Bond lengths



Bond angles

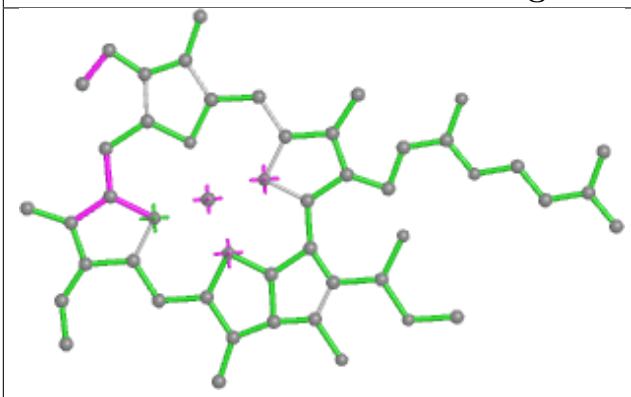


Torsions



Rings

Ligand CLA 4 815



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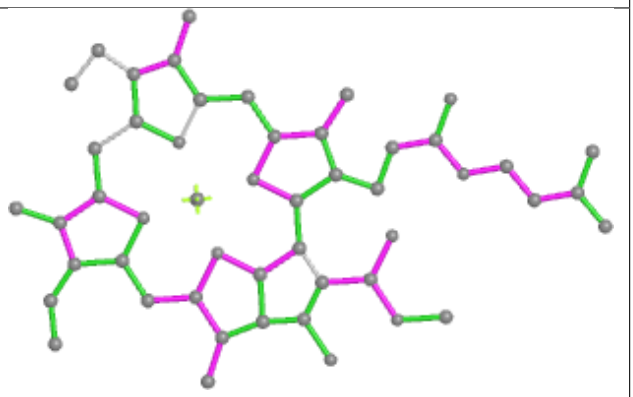
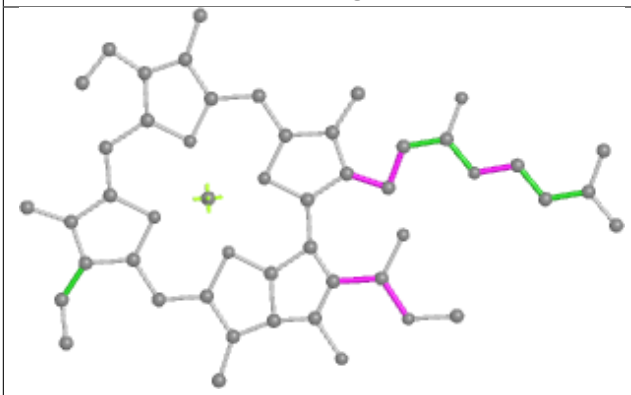


Bond angles



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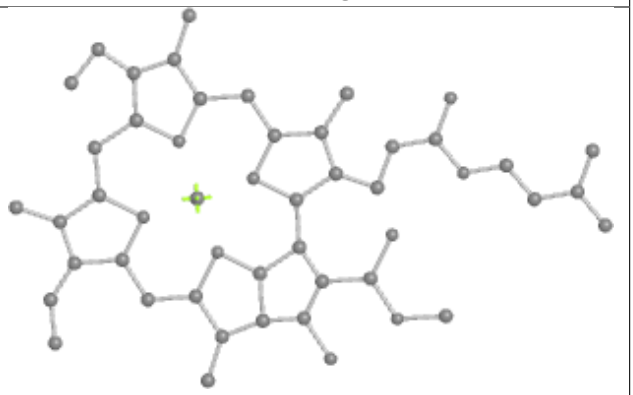
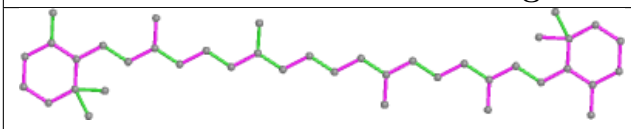
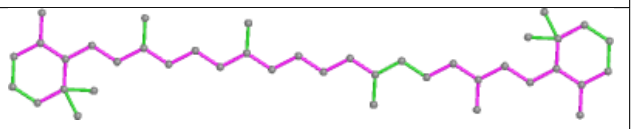


Rings

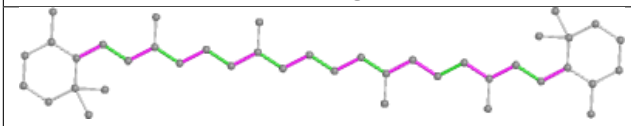
Ligand BCR B 848



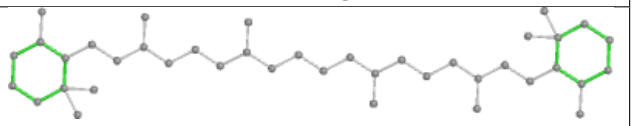
Bond lengths



Bond angles

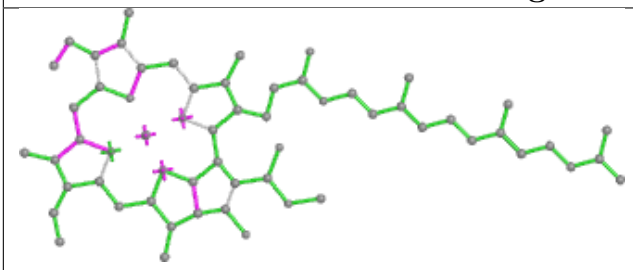


Torsions

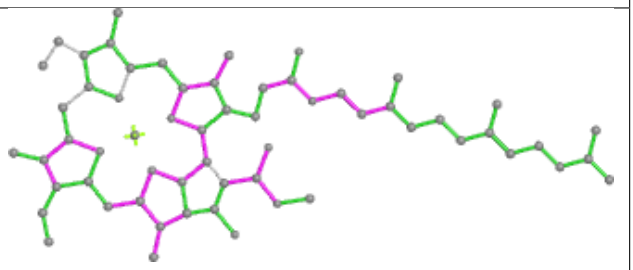


Rings

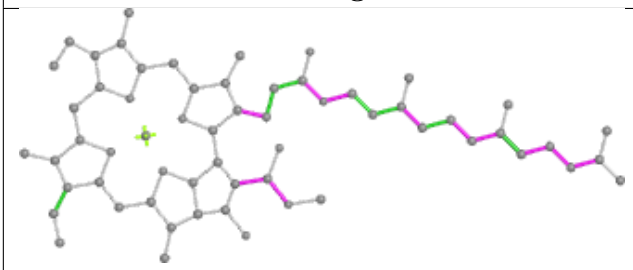
Ligand CLA 4 808



Bond lengths



Bond angles



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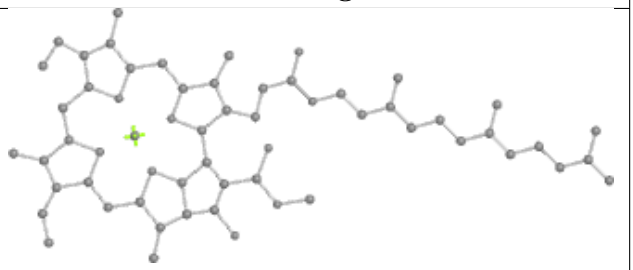
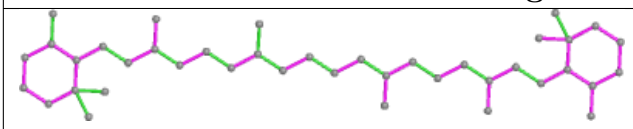
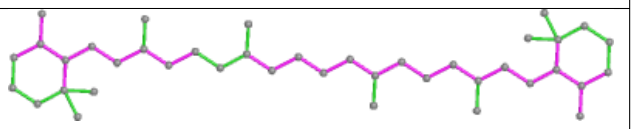


Rings

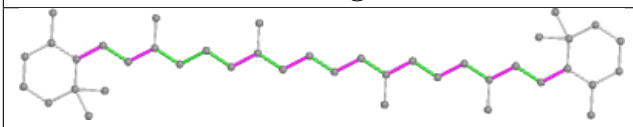
Ligand BCR J 104



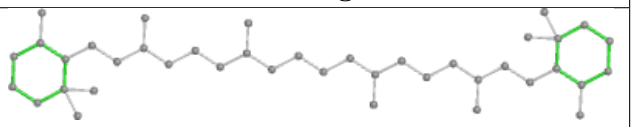
Bond lengths



Bond angles

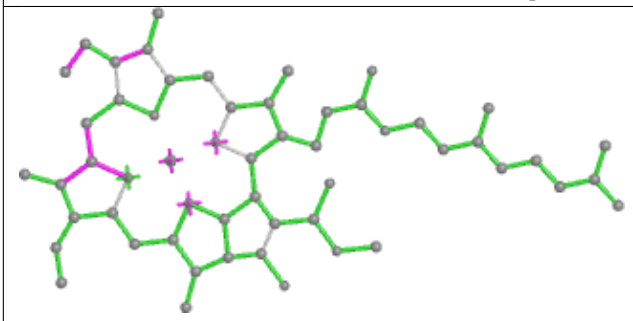


Torsions

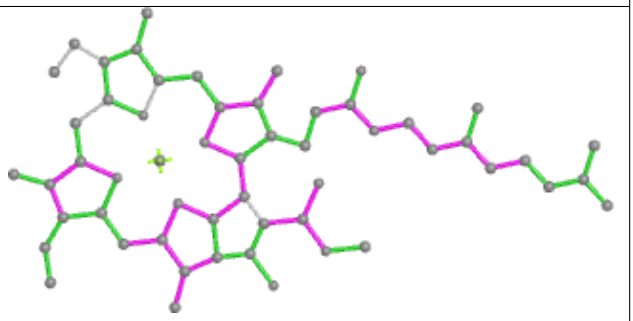


Rings

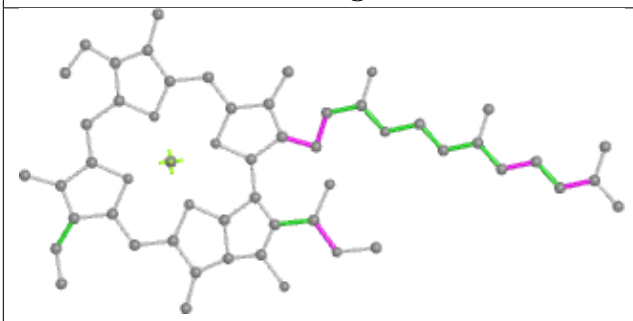
Ligand CLA A 809



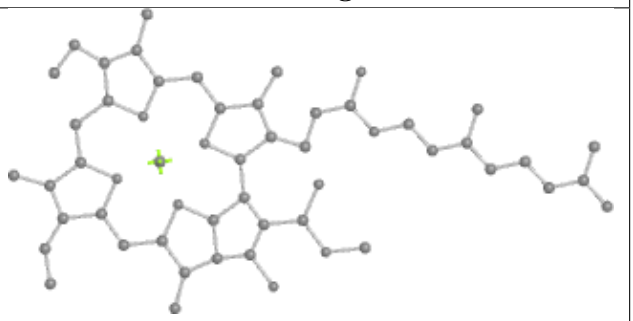
Bond lengths



Bond angles

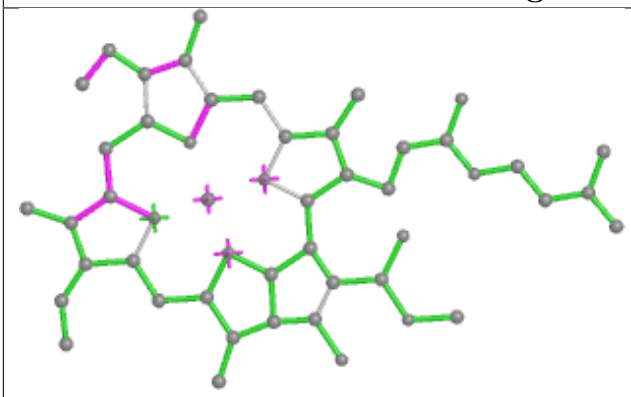


Torsions

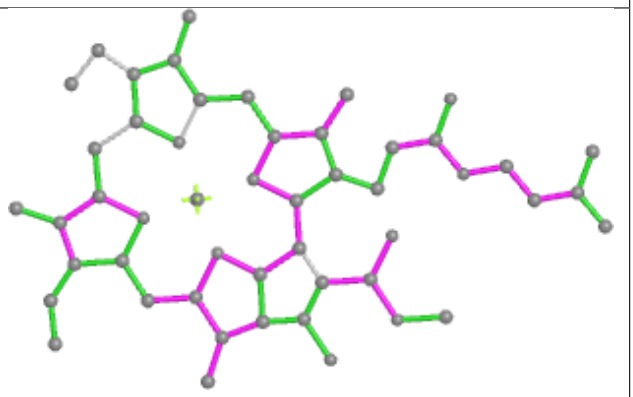


Rings

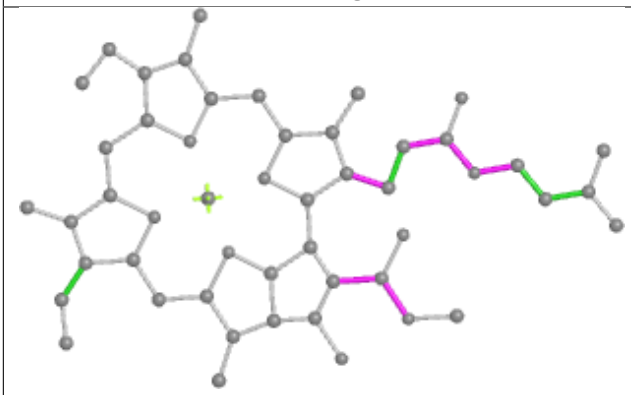
Ligand CLA Z 310



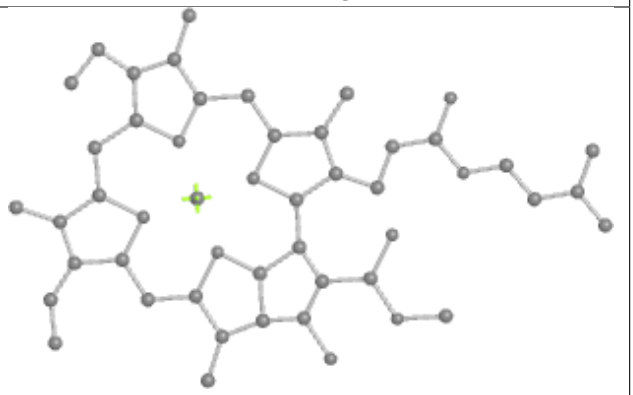
Bond lengths



Bond angles

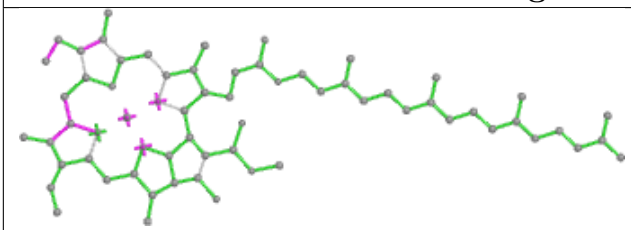


Torsions

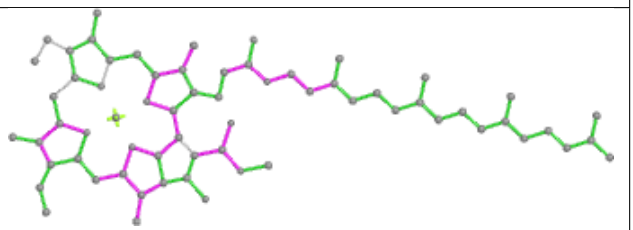


Rings

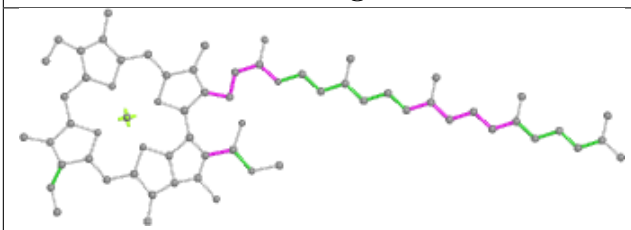
Ligand CLA 6 314



Bond lengths



Bond angles



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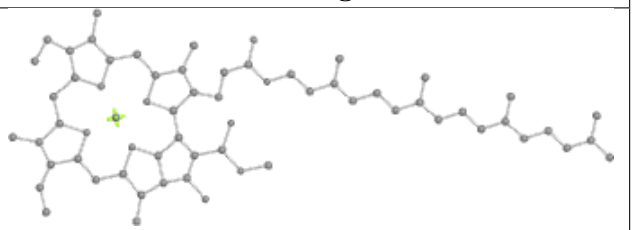
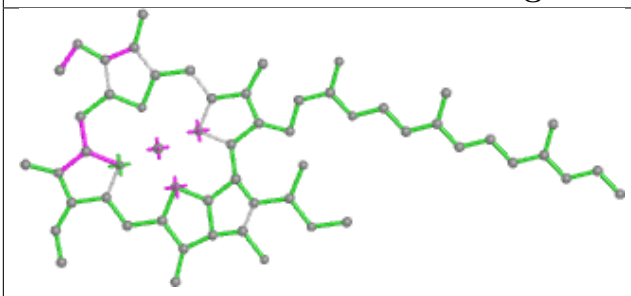
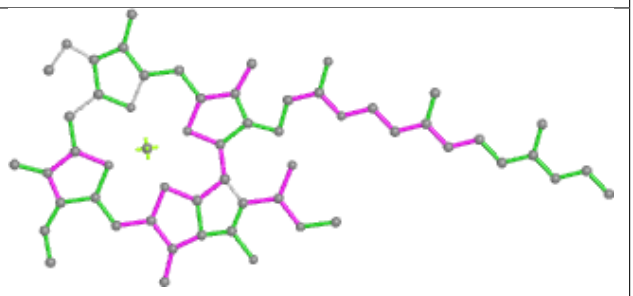


Rings

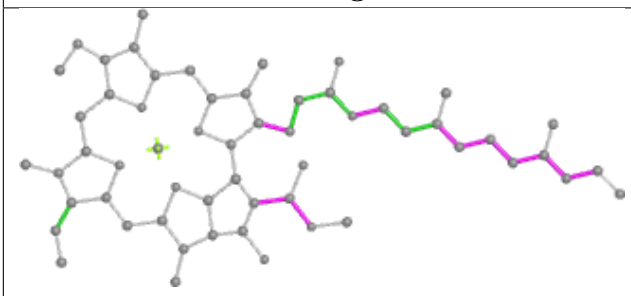
Ligand CLA B 829



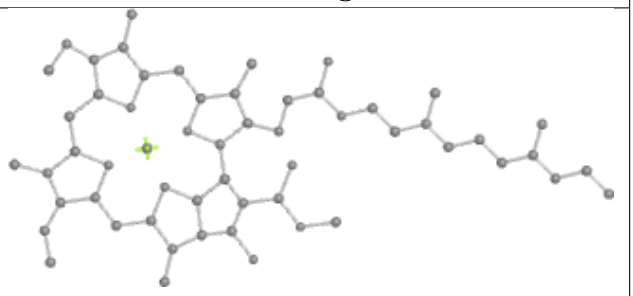
Bond lengths



Bond angles

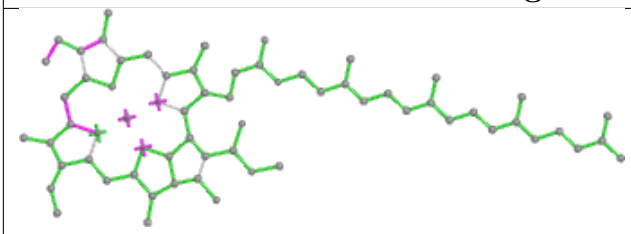


Torsions

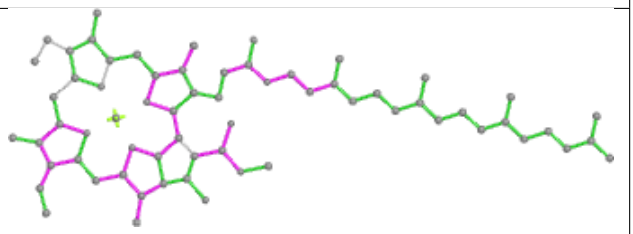


Rings

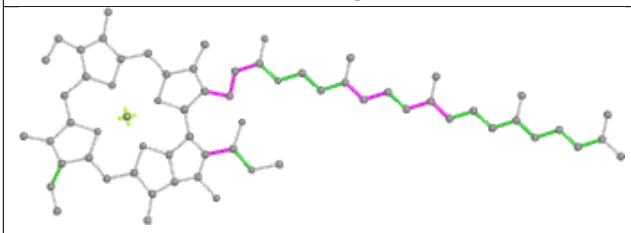
Ligand CLA B 815



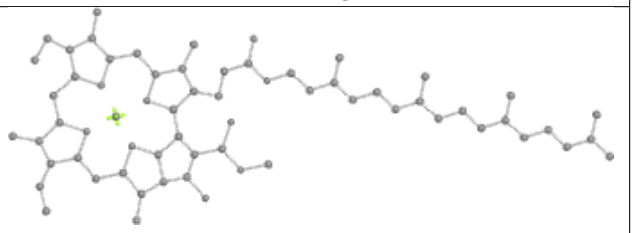
Bond lengths



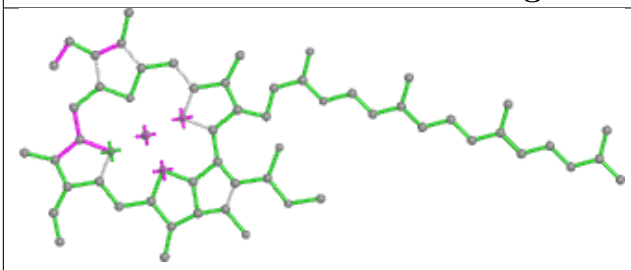
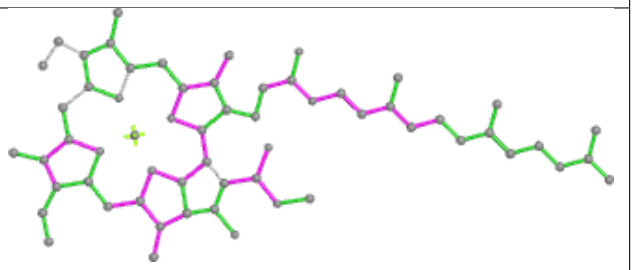
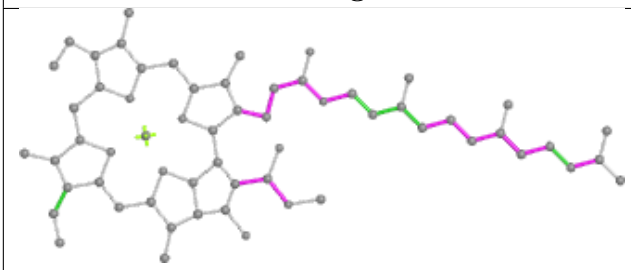
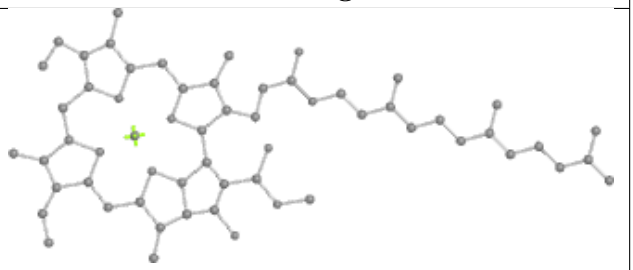
Bond angles

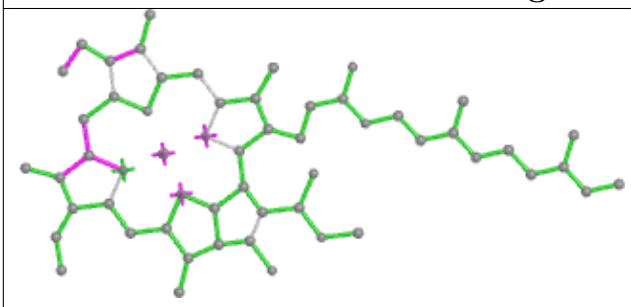
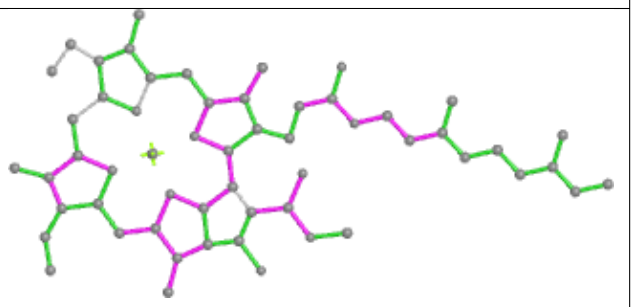
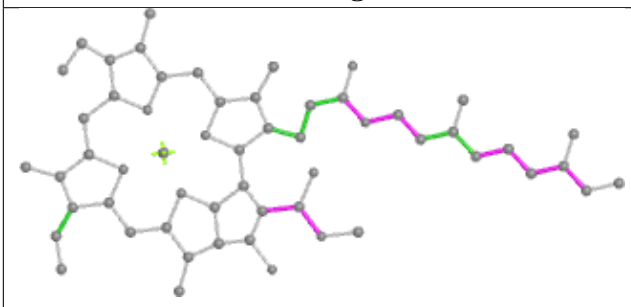
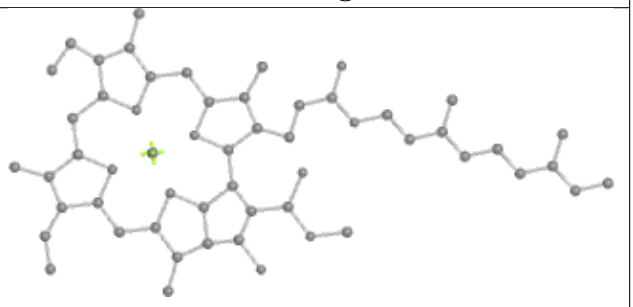


Torsions

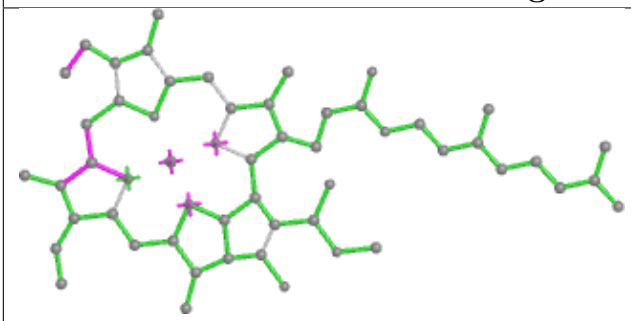


Rings

Ligand CLA A 820	
 Bond lengths	 Bond angles
 Torsions	 Rings

Ligand CLA 5 301	
 Bond lengths	 Bond angles
 Torsions	 Rings

Ligand CLA 7 322



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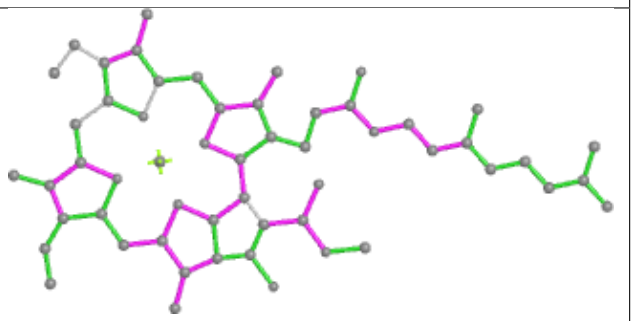

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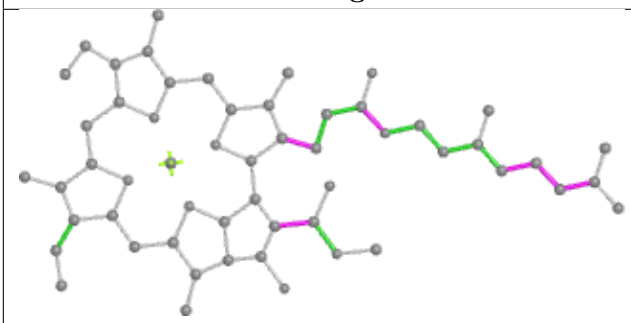

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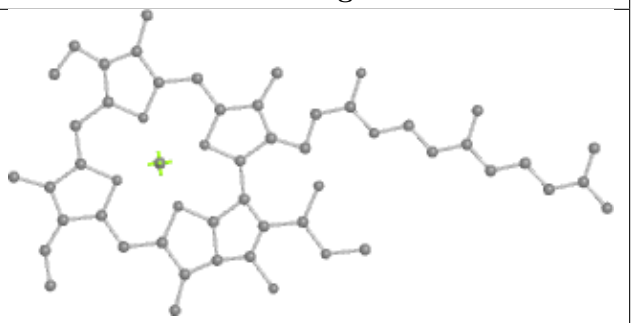
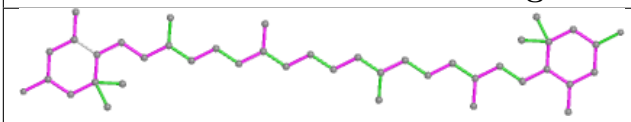


Rings

Ligand LUT Z 301



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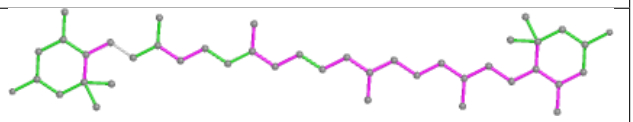
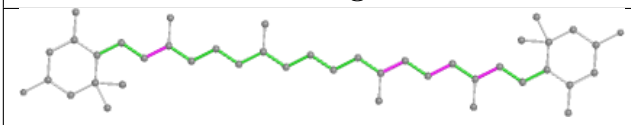


Bond angles



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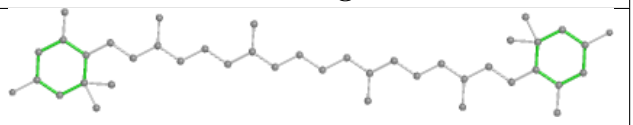
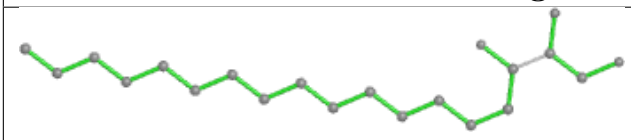


Rings

Ligand SPH 7 320



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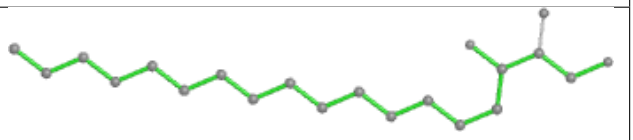

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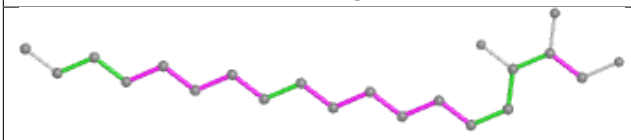
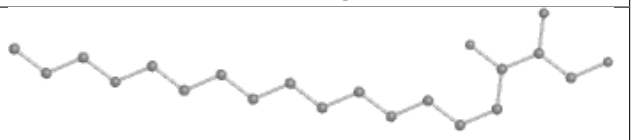
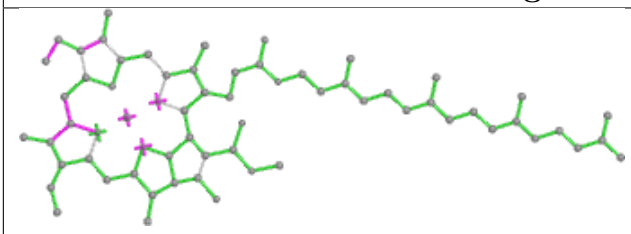


Torsions



Rings

Ligand CLA A 806



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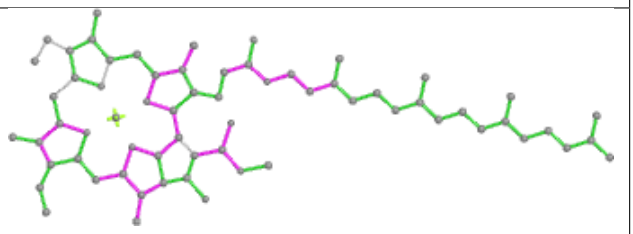
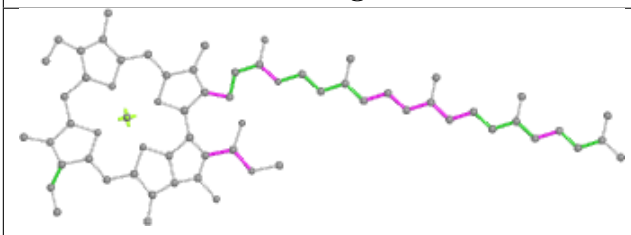


Bond angles



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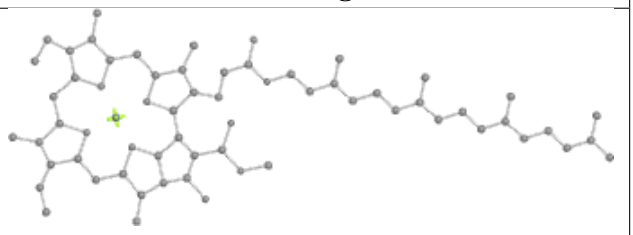
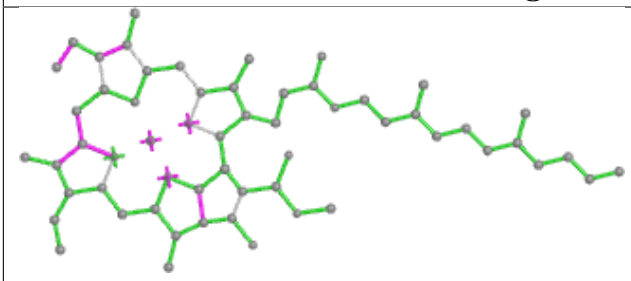


Rings

Ligand CLA B 831



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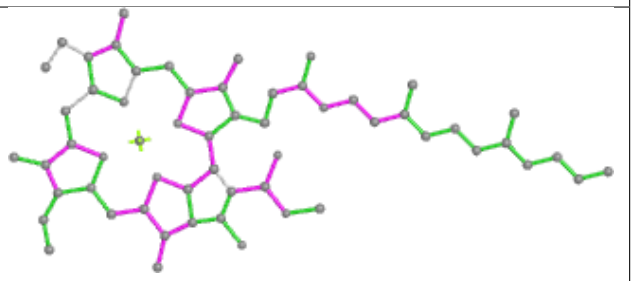
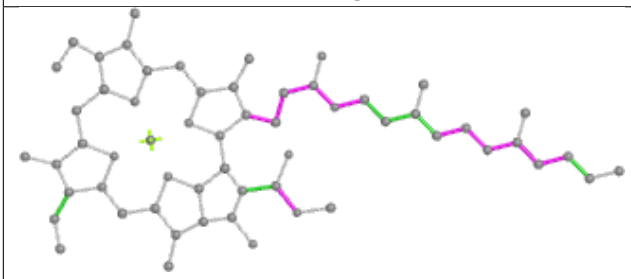


Bond angles



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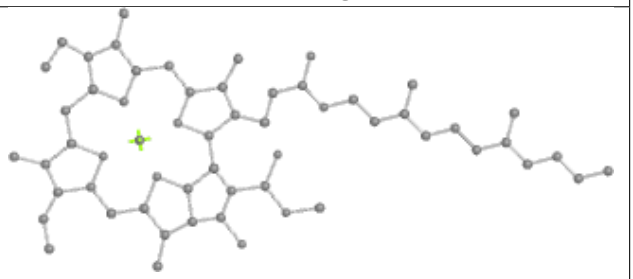
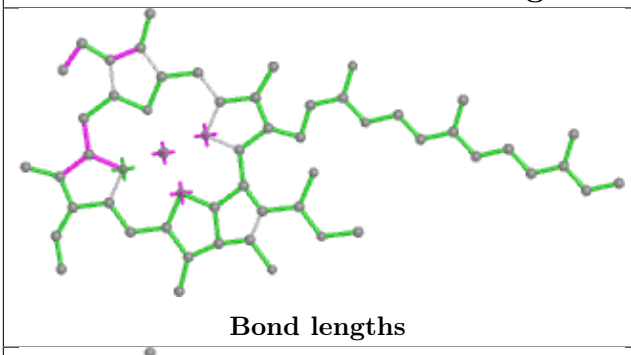
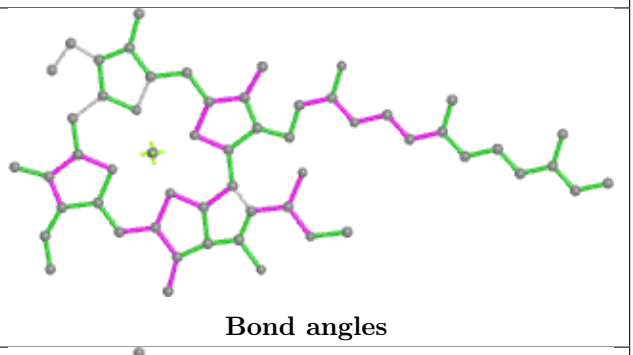


Rings

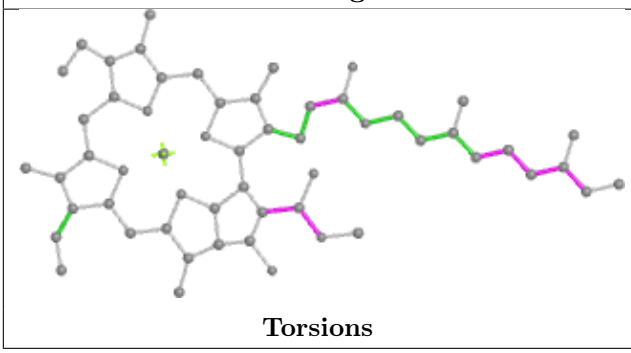
Ligand CLA Z 305



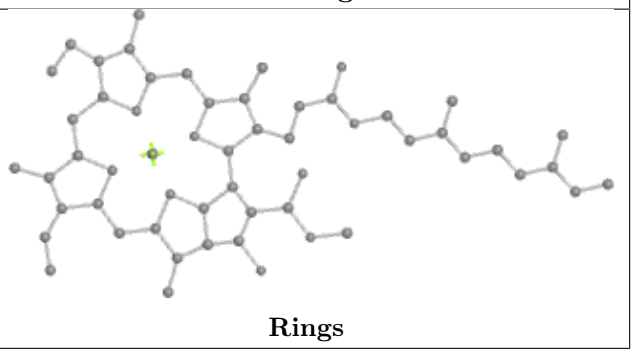
Bond lengths



Bond angles

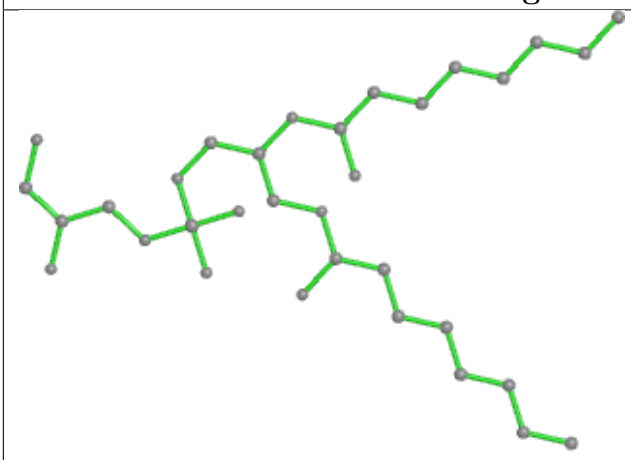


Torsions



Rings

Ligand LHG B 803



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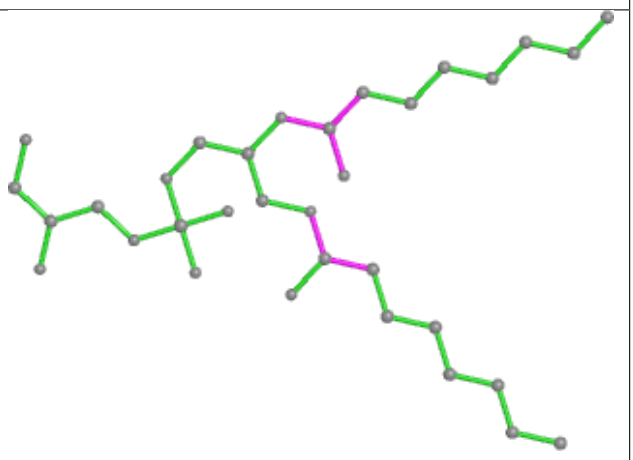
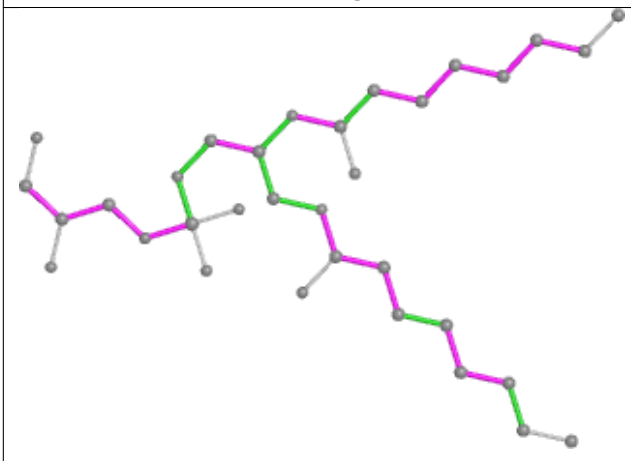


Bond angles



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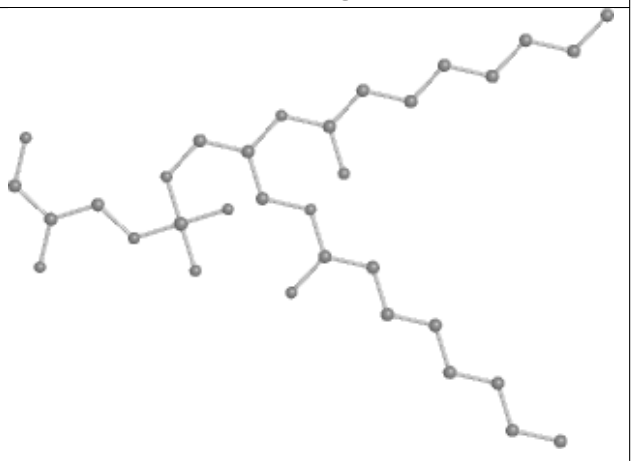
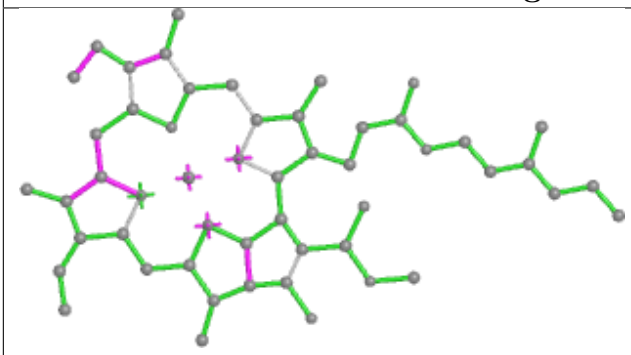


Rings

Ligand CLA A 842



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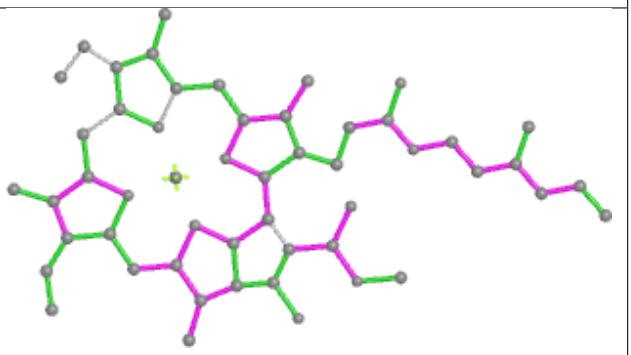
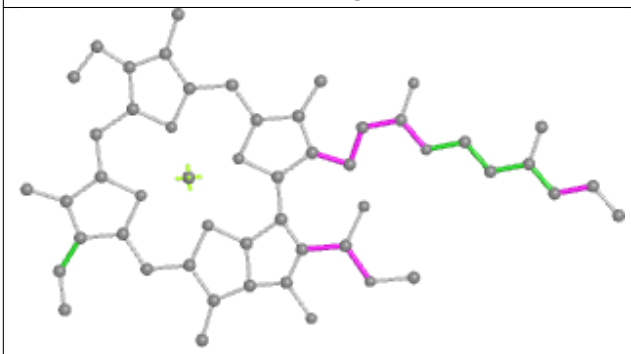
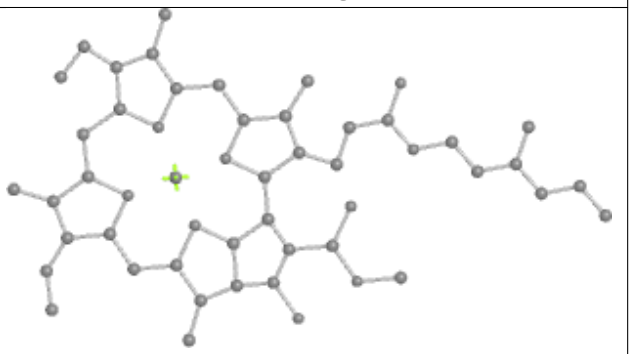


Bond angles

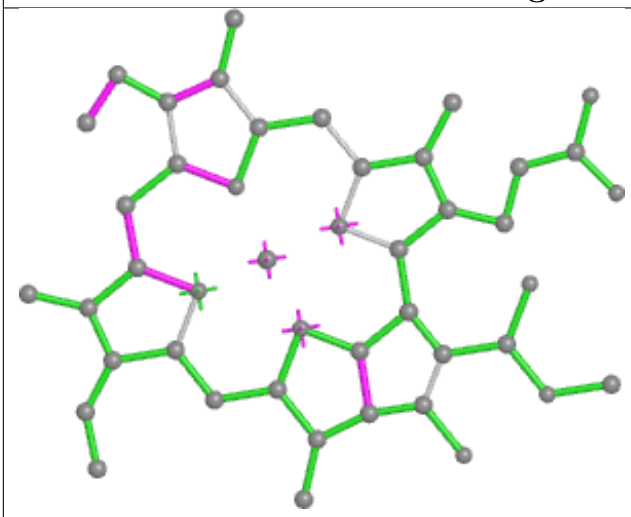


Torsions

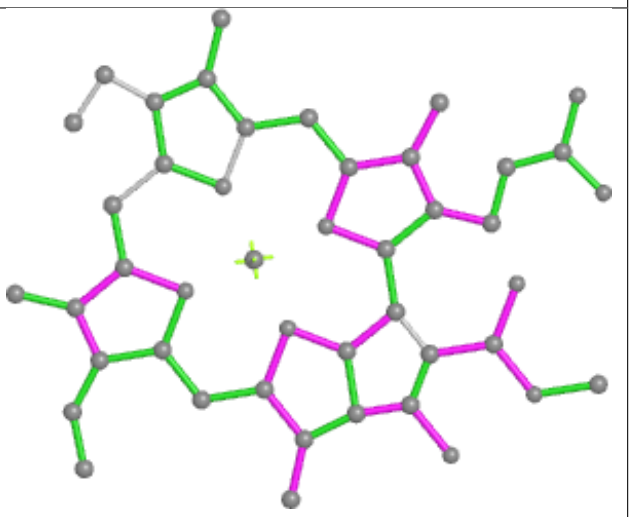


Rings

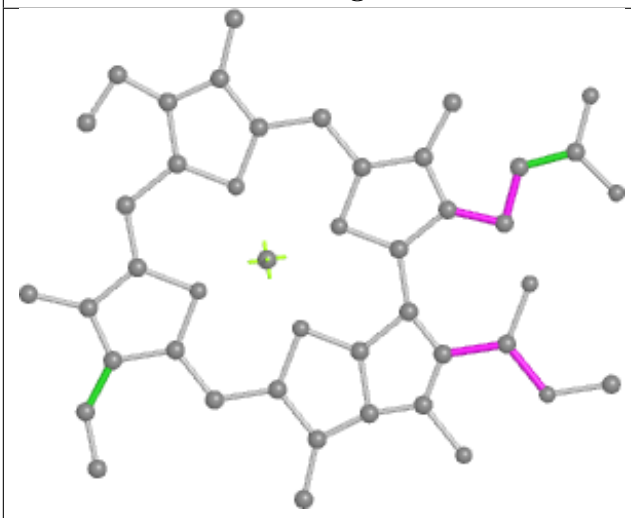
Ligand CLA 3 315



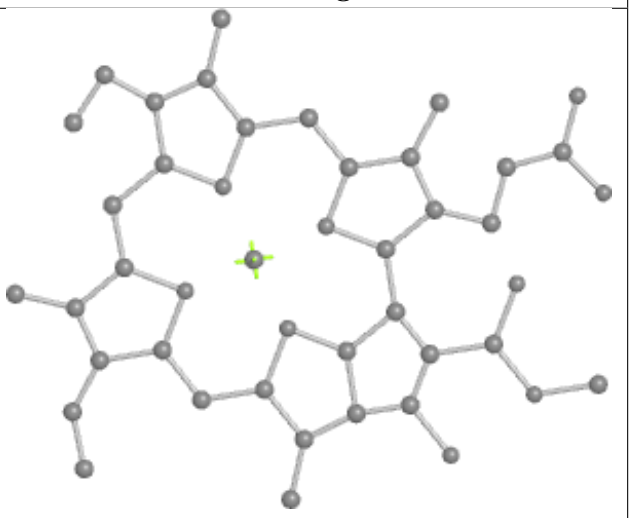
Bond lengths



Bond angles

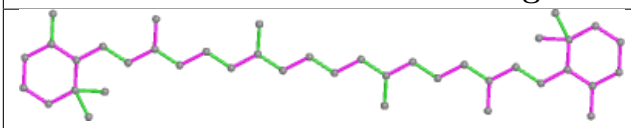


Torsions

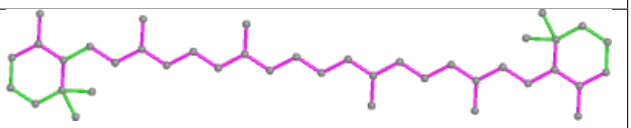


Rings

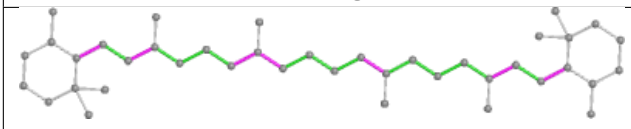
Ligand BCR B 844



Bond lengths



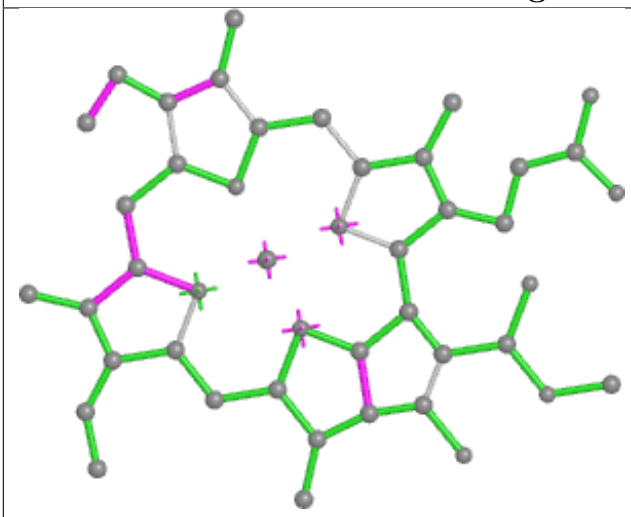
Bond angles



Torsions

Rings

Ligand CLA 1 305



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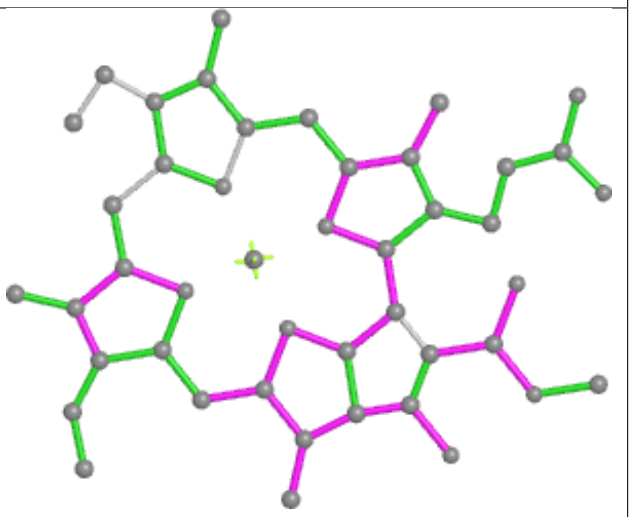
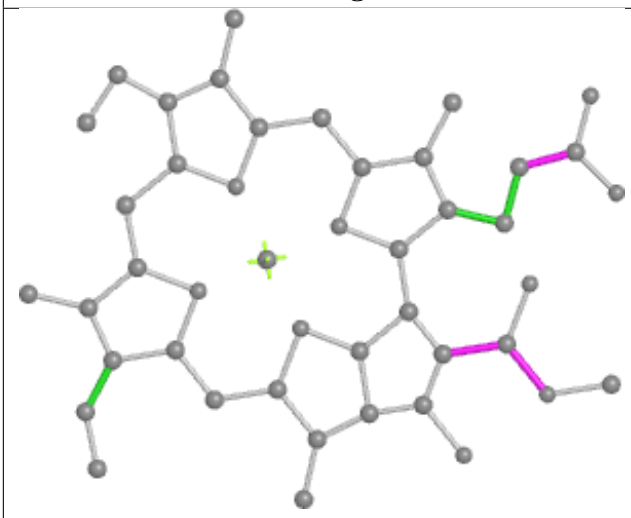
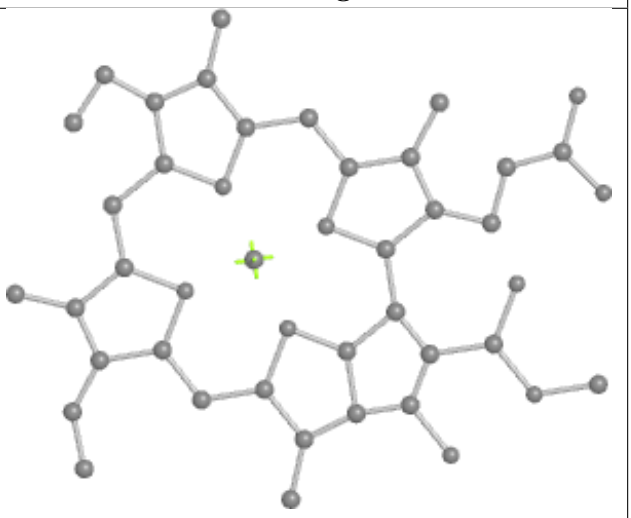


Bond angles

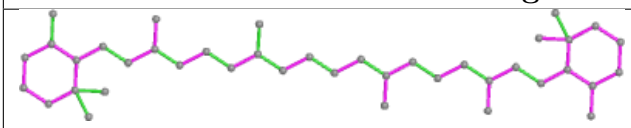


Torsions

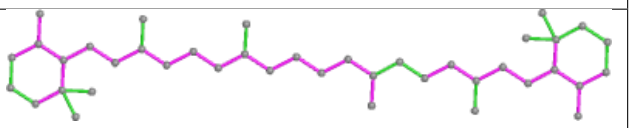


Rings

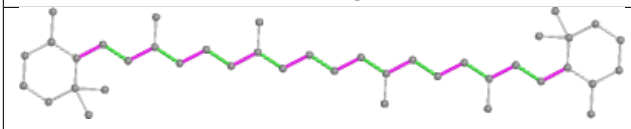
Ligand BCR A 847



Bond lengths



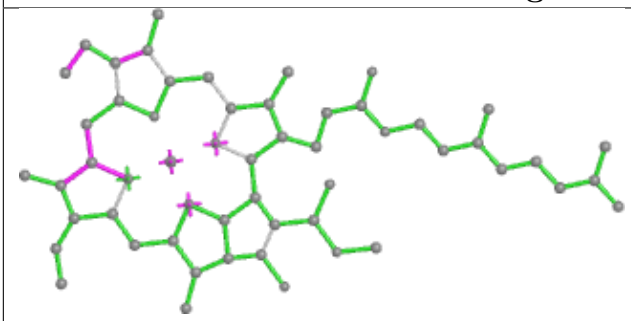
Bond angles



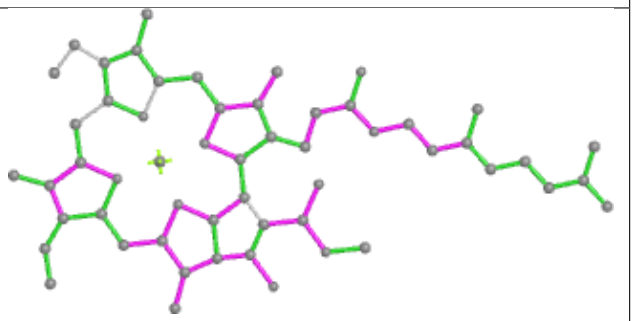
Torsions

Rings

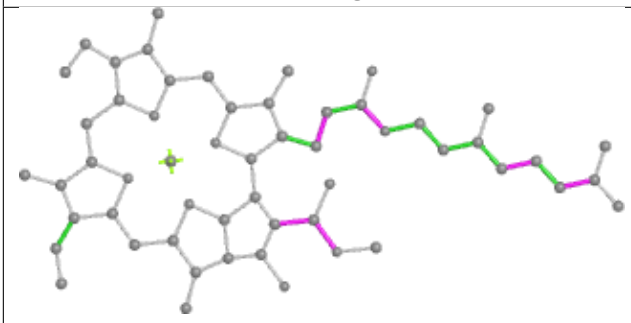
Ligand CLA Z 307



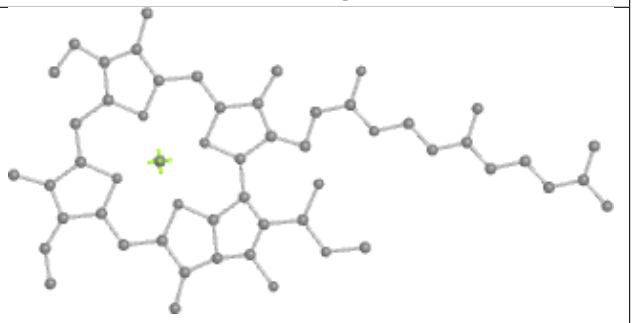
Bond lengths



Bond angles

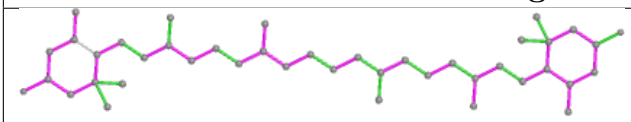


Torsions



Rings

Ligand LUT 7 301



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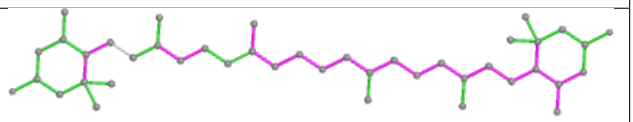
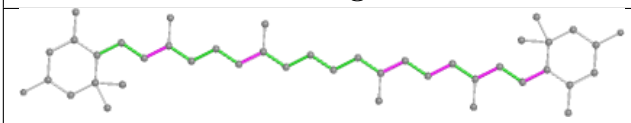


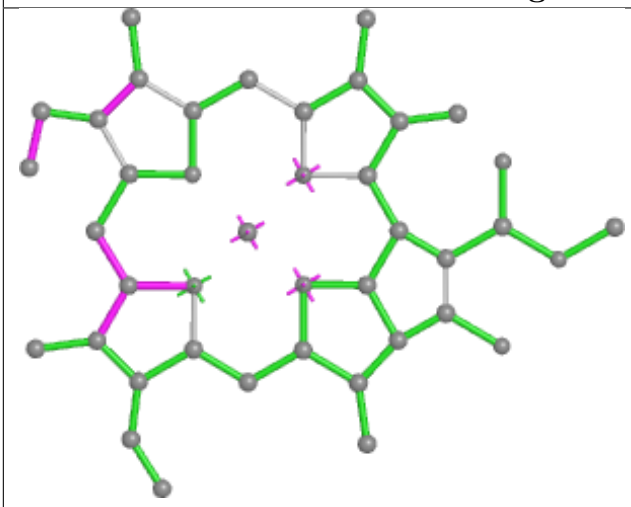
Bond angles



Torsions

Rings

Ligand CLA 4 817



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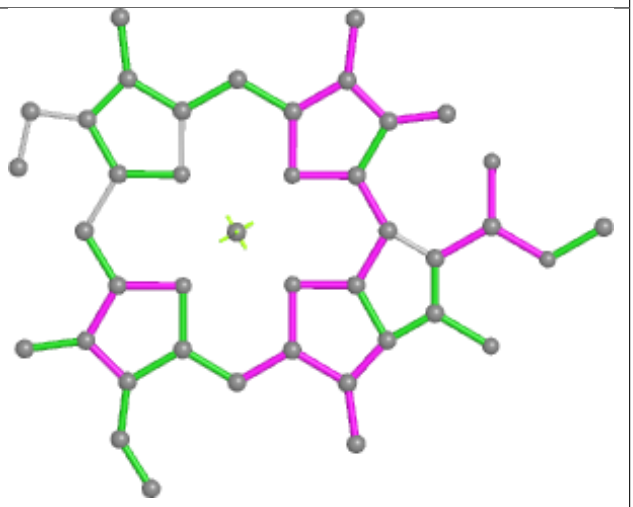
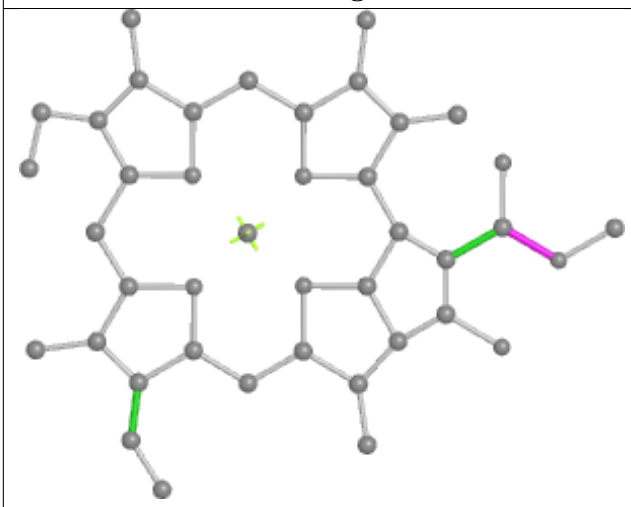
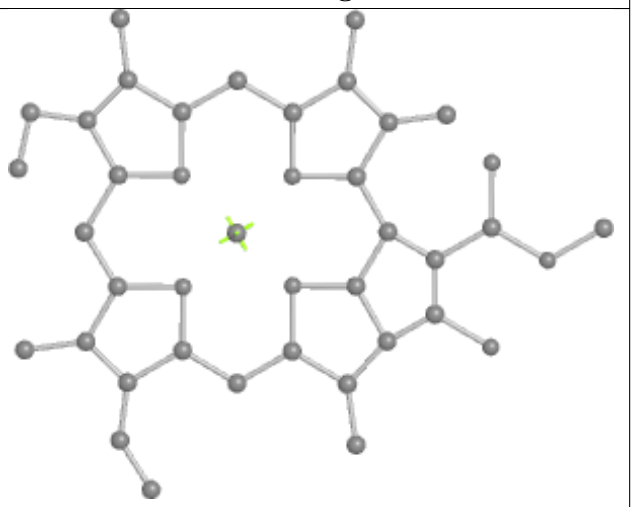


Bond angles

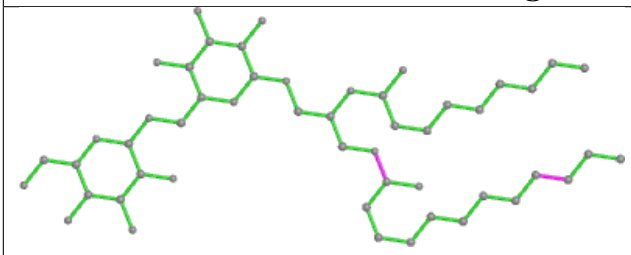


Torsions



Rings

Ligand DGD 3 301



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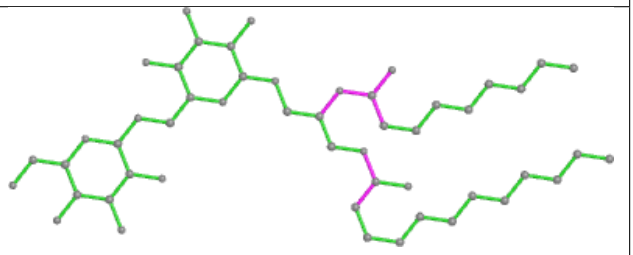
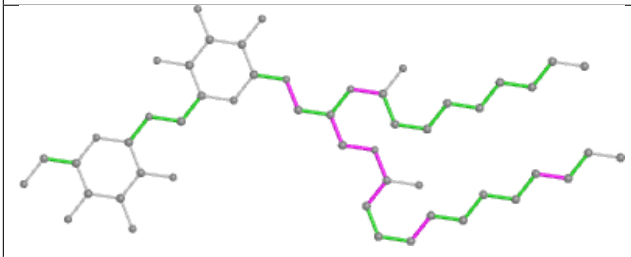


Bond angles



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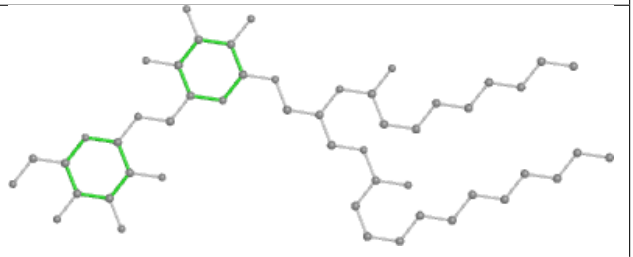
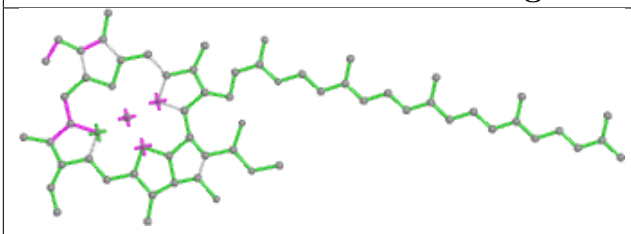


Rings

Ligand CLA B 838



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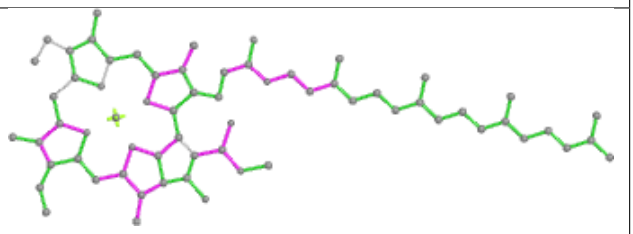
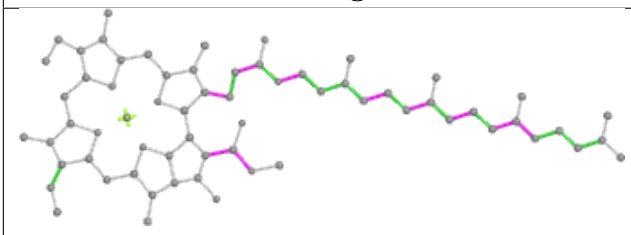


Bond angles



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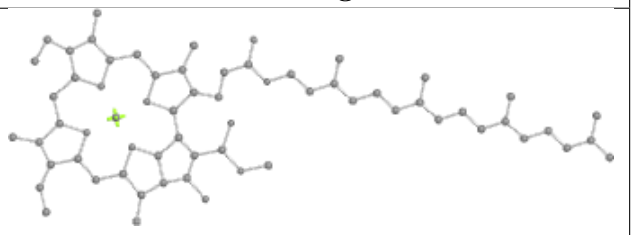
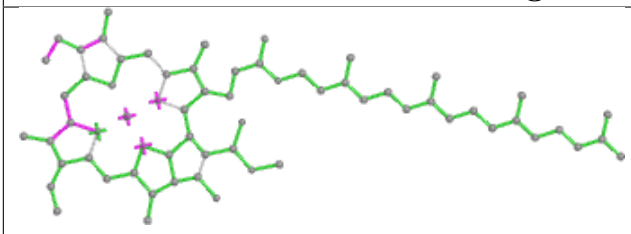


Rings

Ligand CLA F 304



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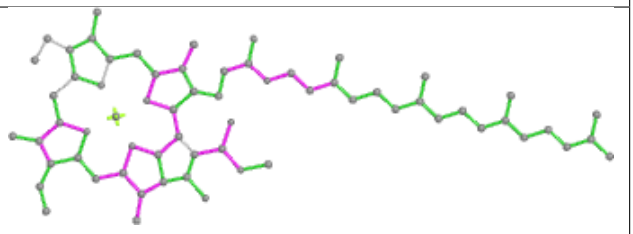
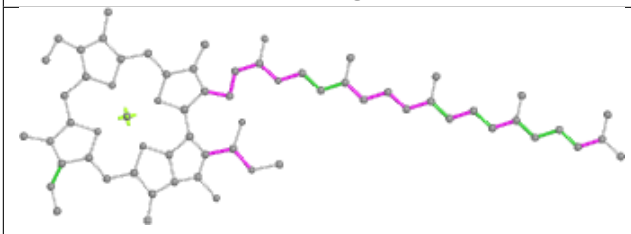


Bond angles



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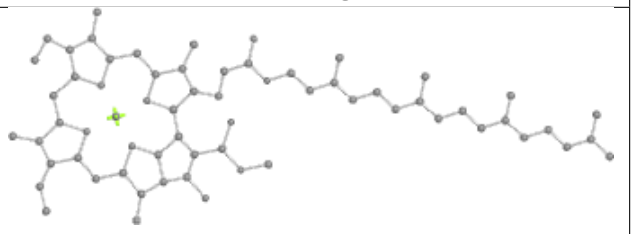
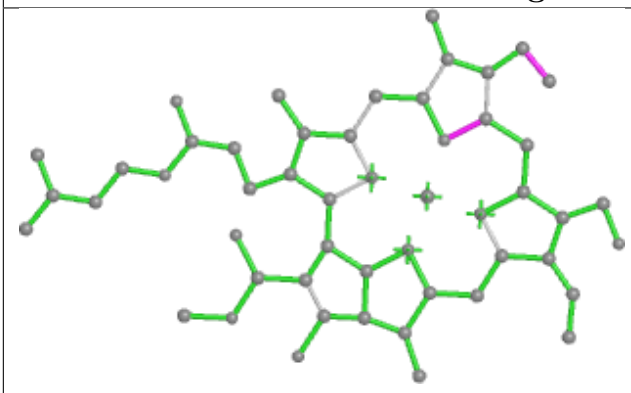
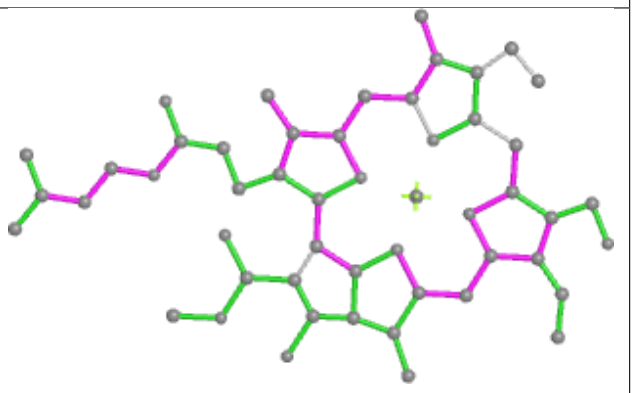


Rings

Ligand CHL 2 303



Bond lengths



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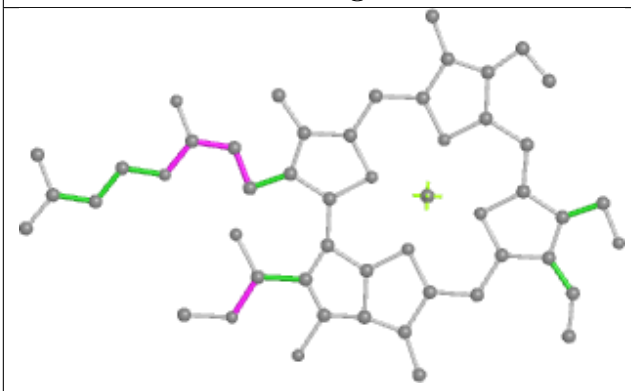
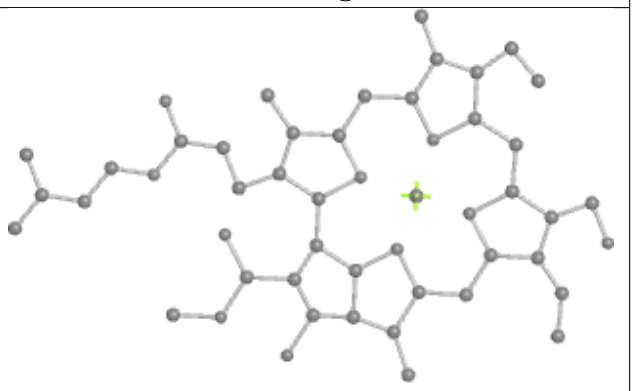
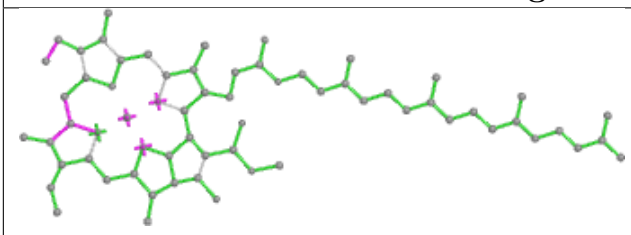


Torsions



Rings

Ligand CLA 1 304



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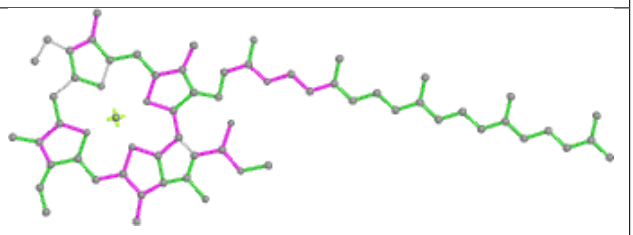
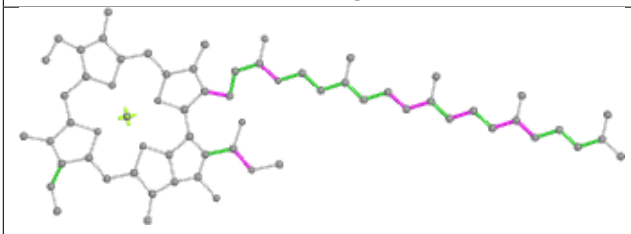
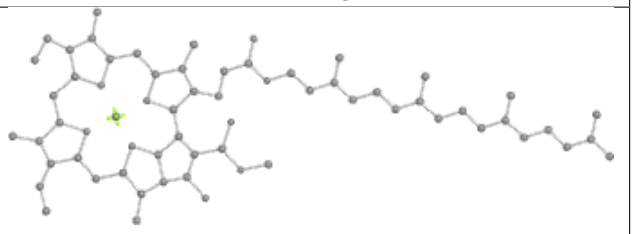


Bond angles

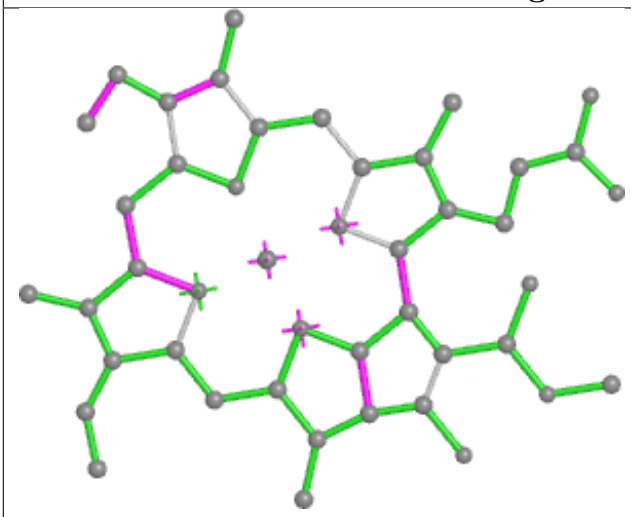


Torsions

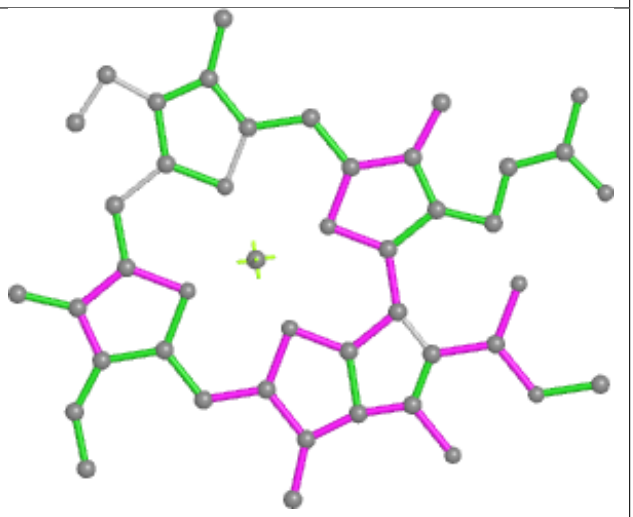


Rings

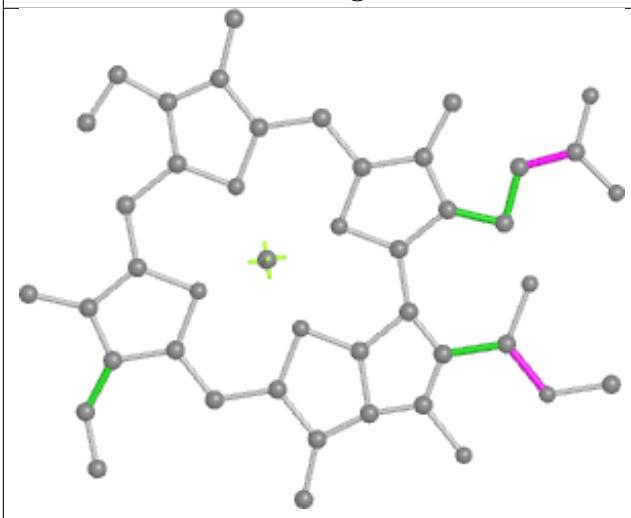
Ligand CLA 9 310



Bond lengths



Bond angles



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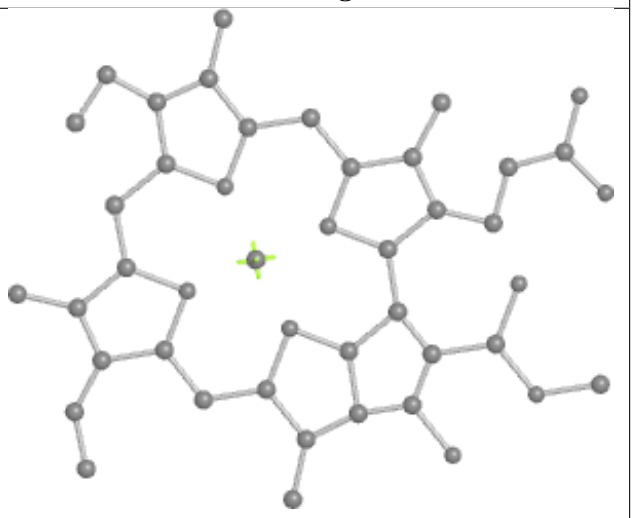
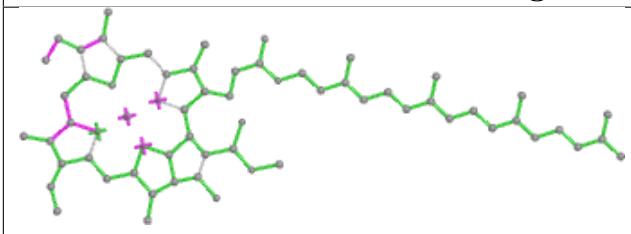
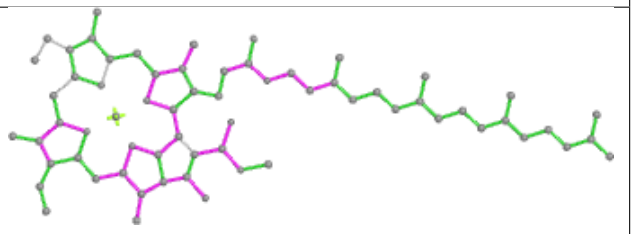


Rings

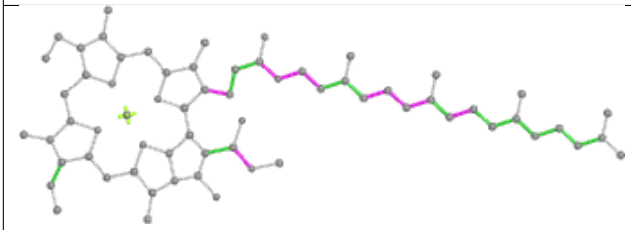
Ligand CLA 1 313



Bond lengths



Bond angles



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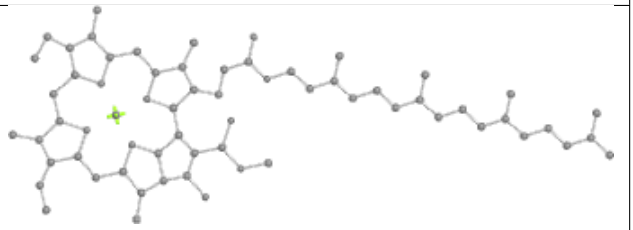
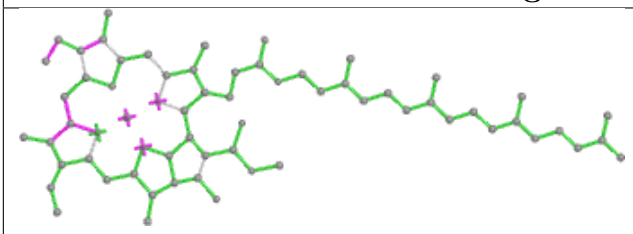
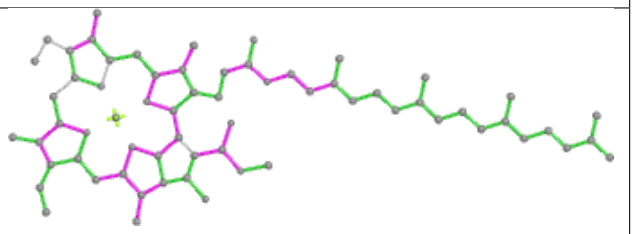


Rings

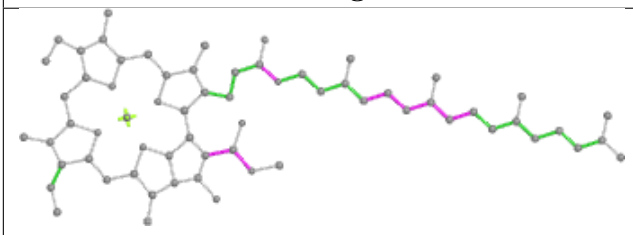
Ligand CLA A 835



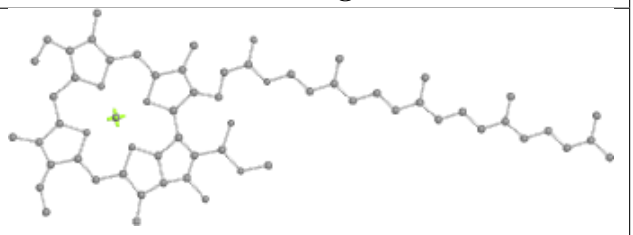
Bond lengths



Bond angles

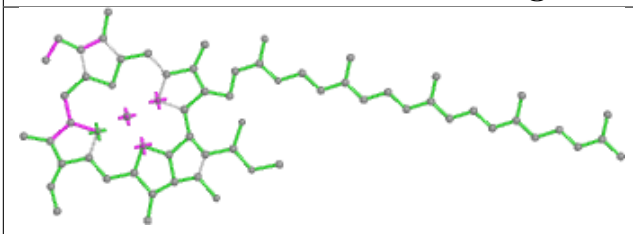


Torsions

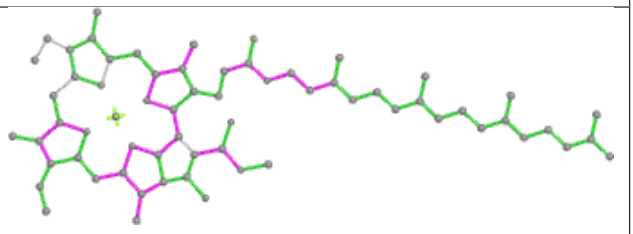


Rings

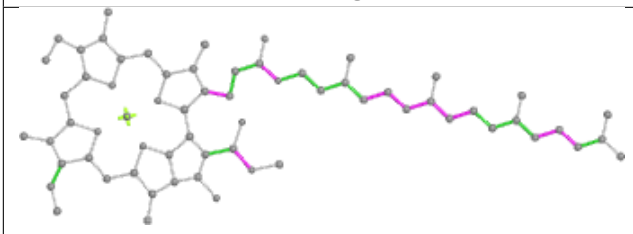
Ligand CLA 7 310



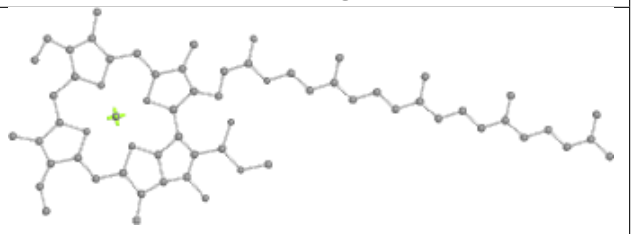
Bond lengths



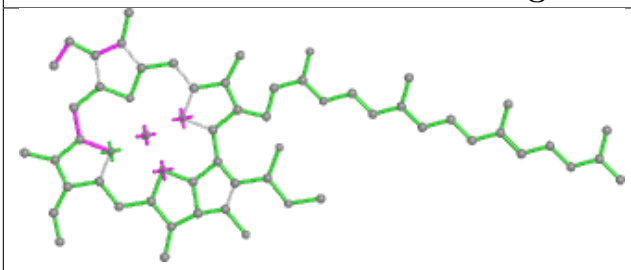
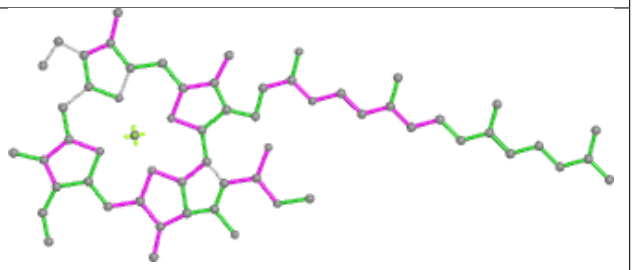
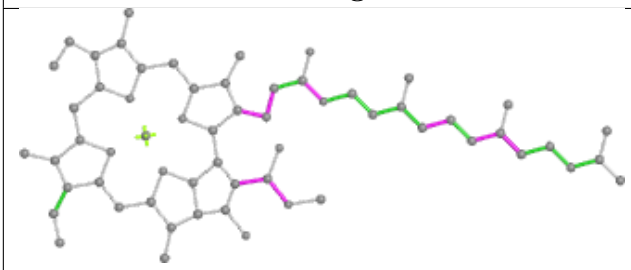
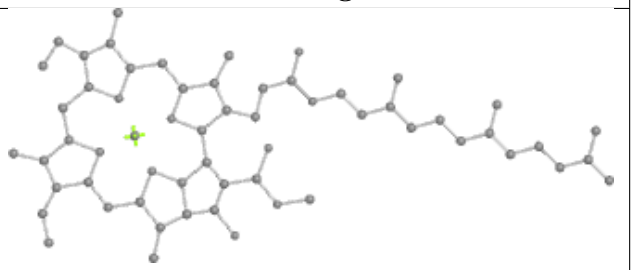
Bond angles

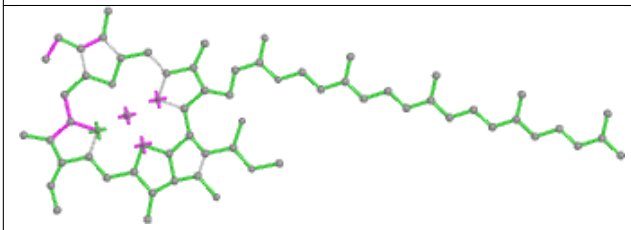
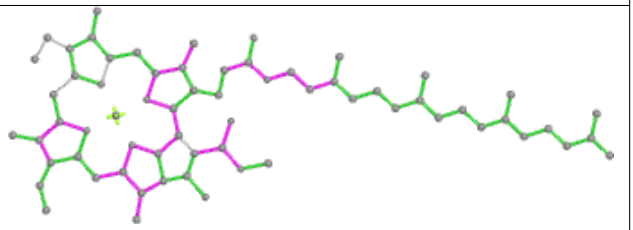
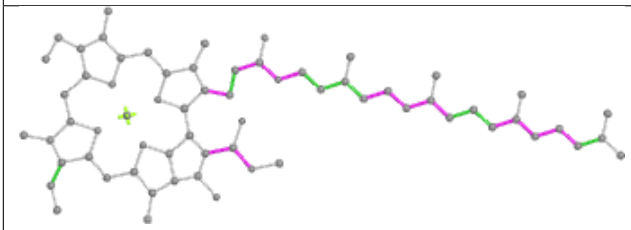
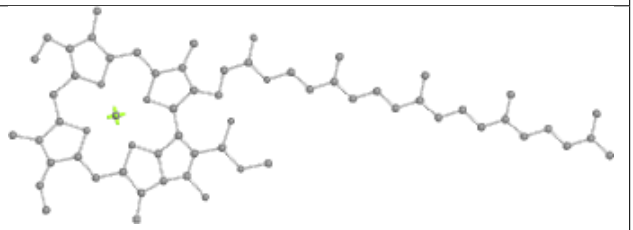


Torsions

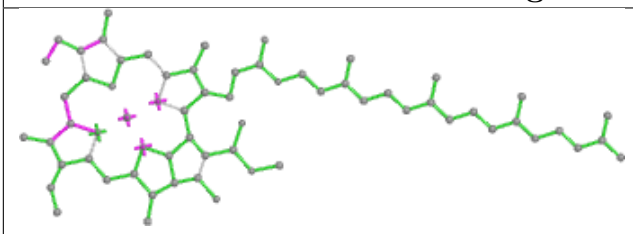


Rings

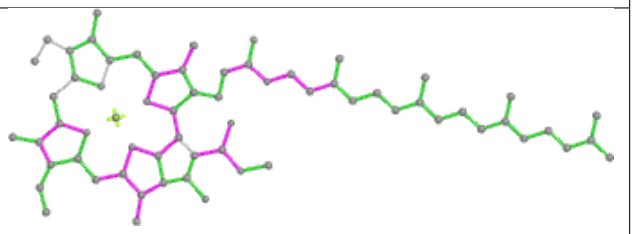
Ligand CLA 8 305	
 Bond lengths	 Bond angles
 Torsions	 Rings

Ligand CLA Z 316	
 Bond lengths	 Bond angles
 Torsions	 Rings

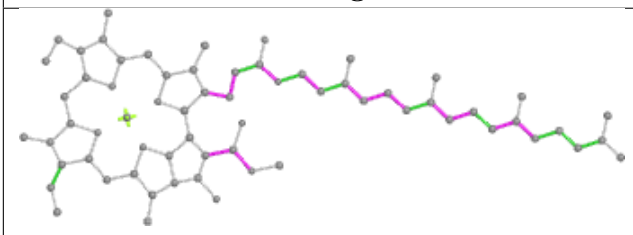
Ligand CLA L 202



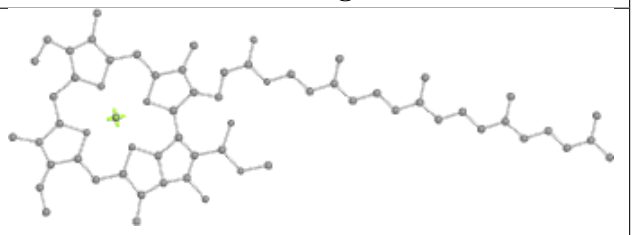
Bond lengths



Bond angles

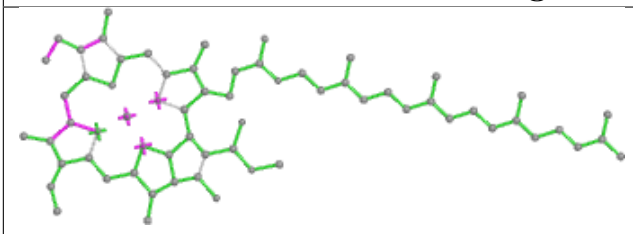


Torsions



Rings

Ligand CLA 5 322



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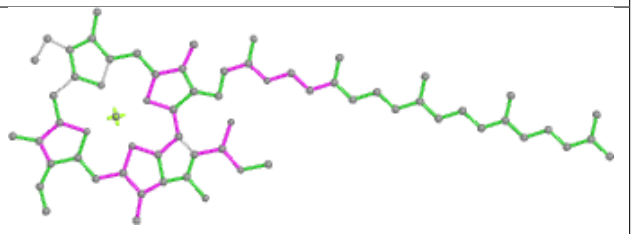
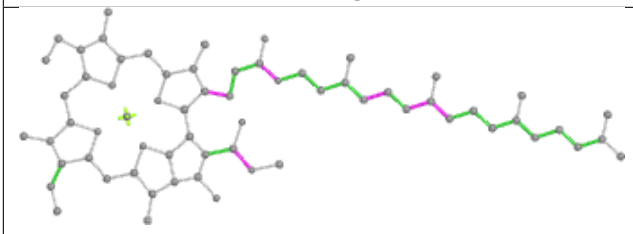
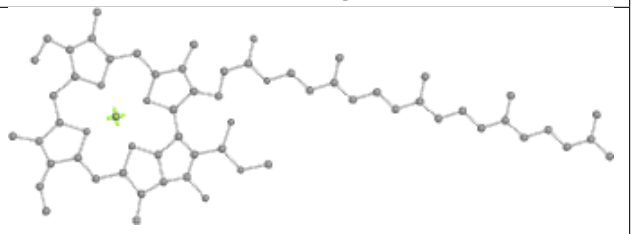


Bond angles

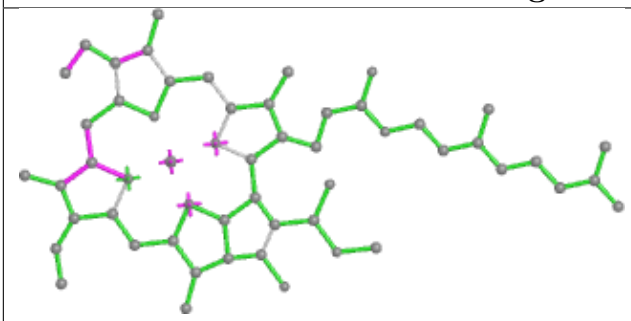


Torsions



Rings

Ligand CLA 5 326



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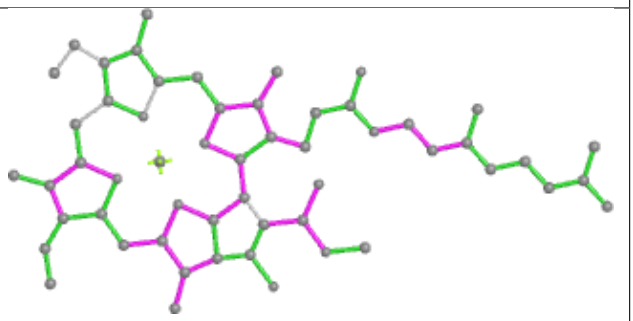
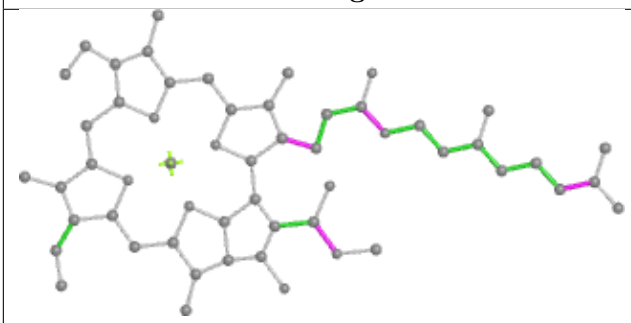


Bond angles



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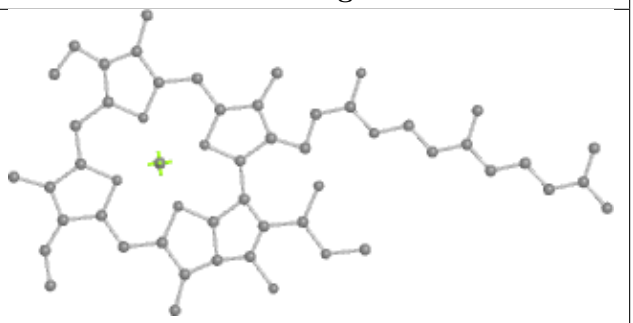
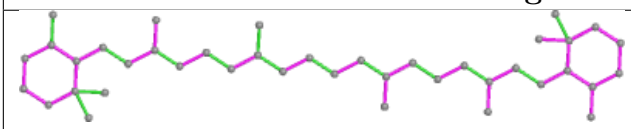


Rings

Ligand BCR L 204



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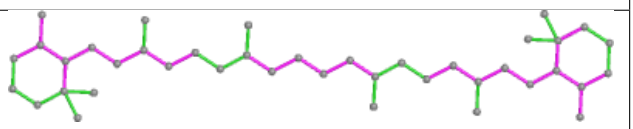
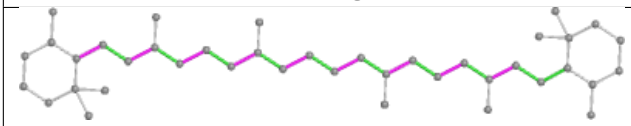


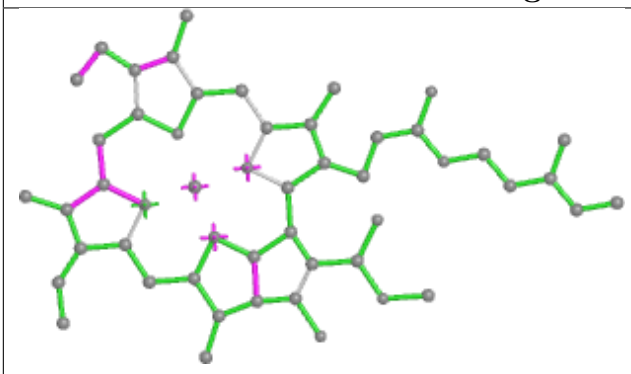
Bond angles



Torsions

Rings

Ligand CLA A 839



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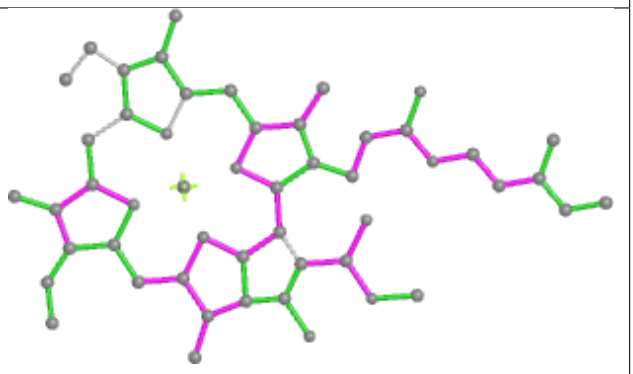
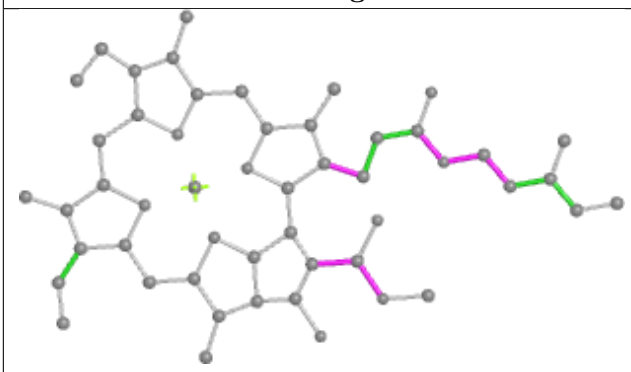
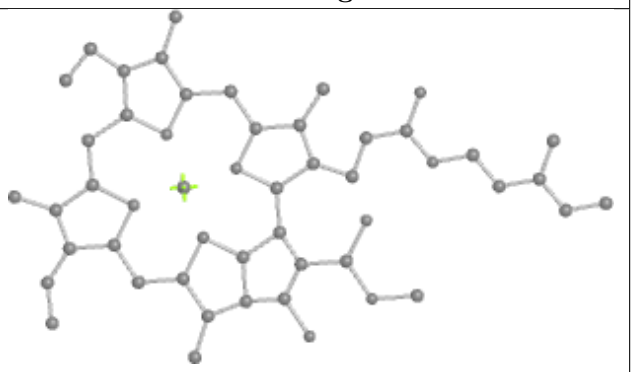


Bond angles

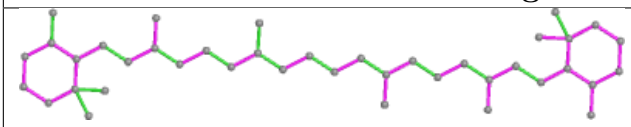


Torsions



Rings

Ligand BCR A 848



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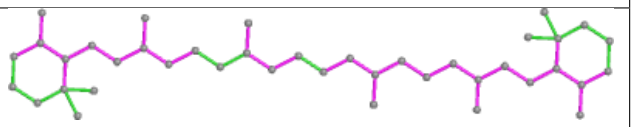
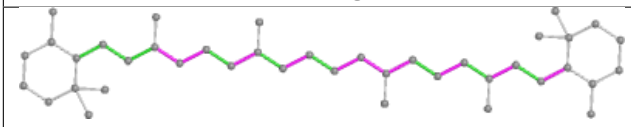


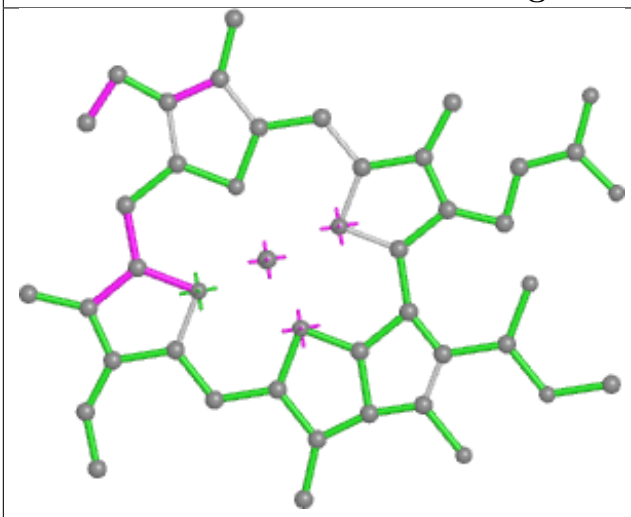
Bond angles



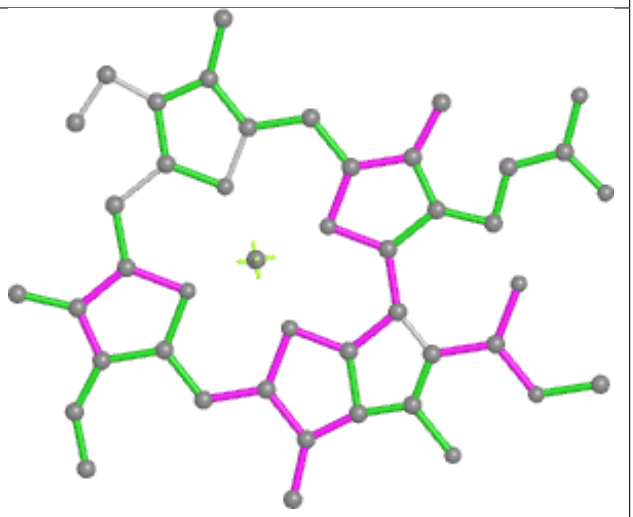
Torsions

Rings

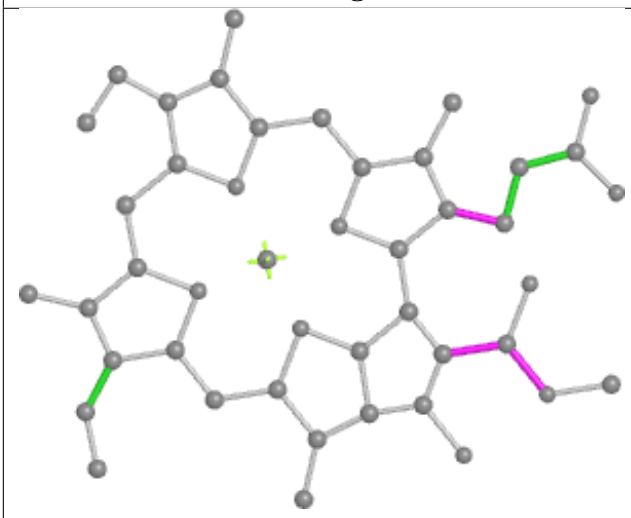
Ligand CLA 5 314



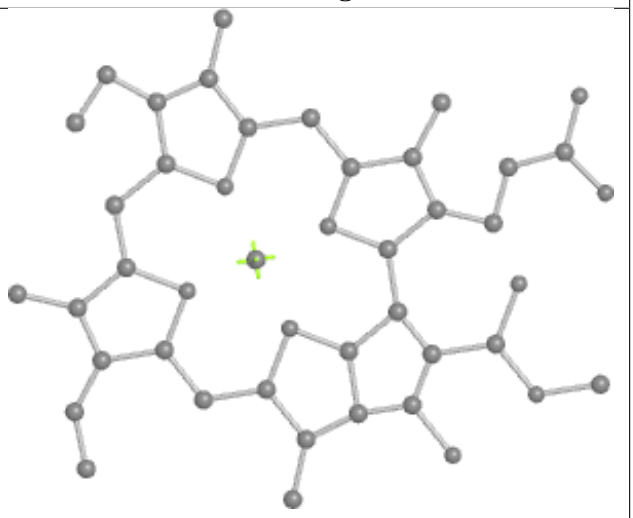
Bond lengths



Bond angles

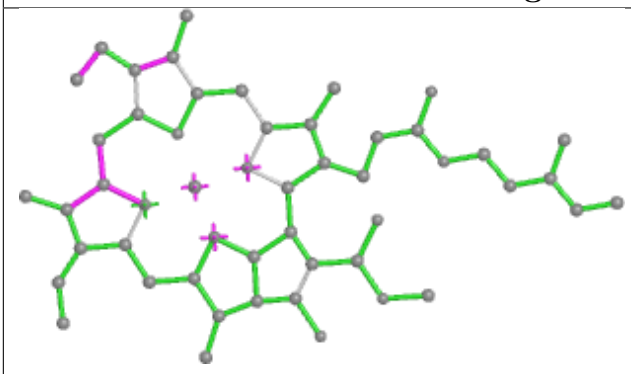


Torsions

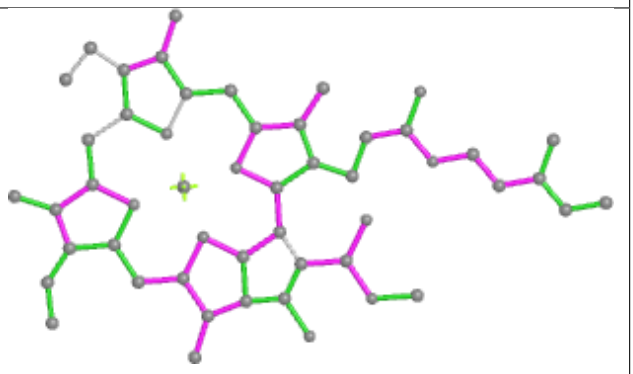


Rings

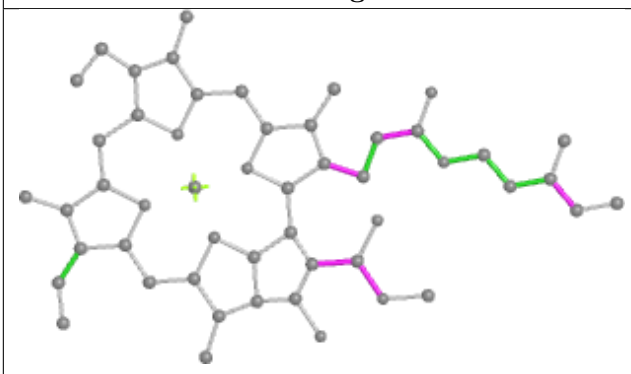
Ligand CLA B 837



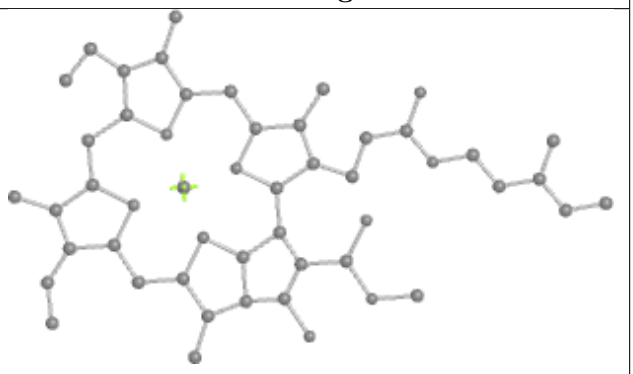
Bond lengths



Bond angles

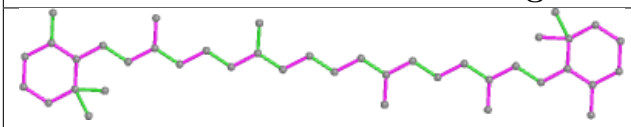


Torsions

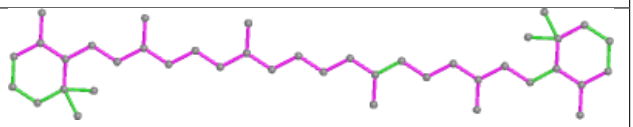


Rings

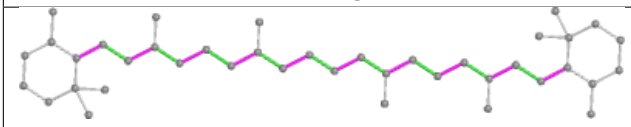
Ligand BCR 6 306



Bond lengths



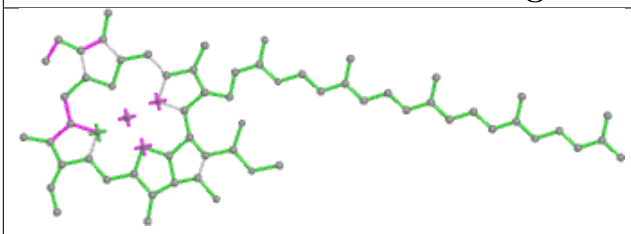
Bond angles



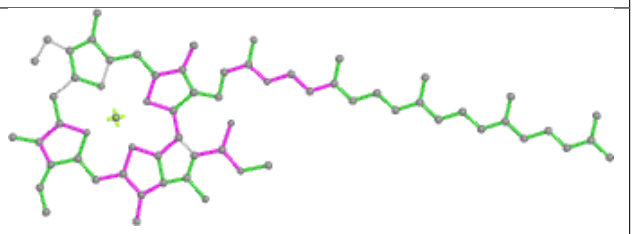
Torsions

Rings

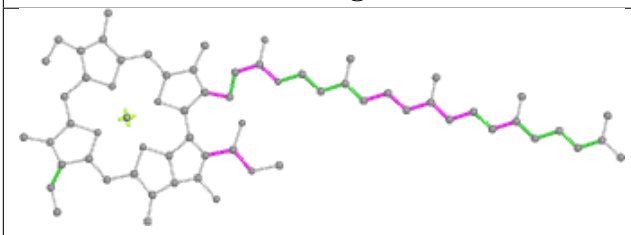
Ligand CLA L 201



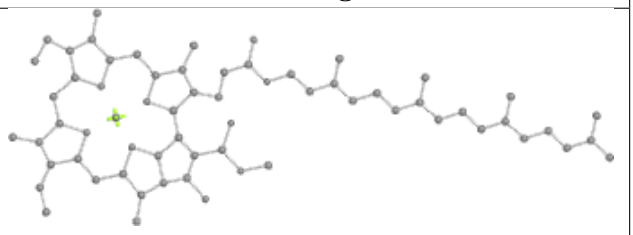
Bond lengths



Bond angles

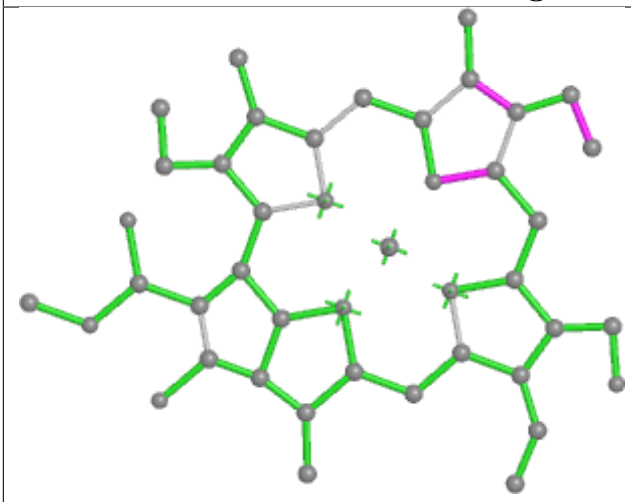


Torsions



Rings

Ligand CHL 6 320



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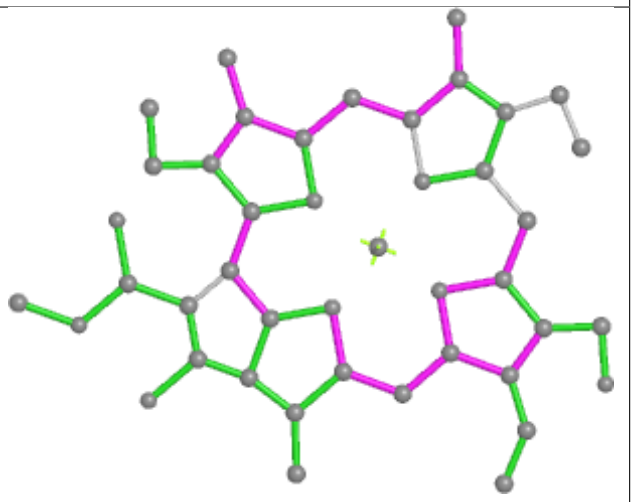
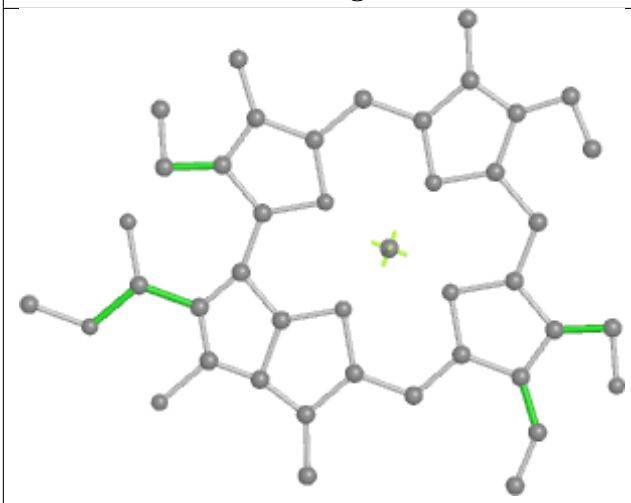
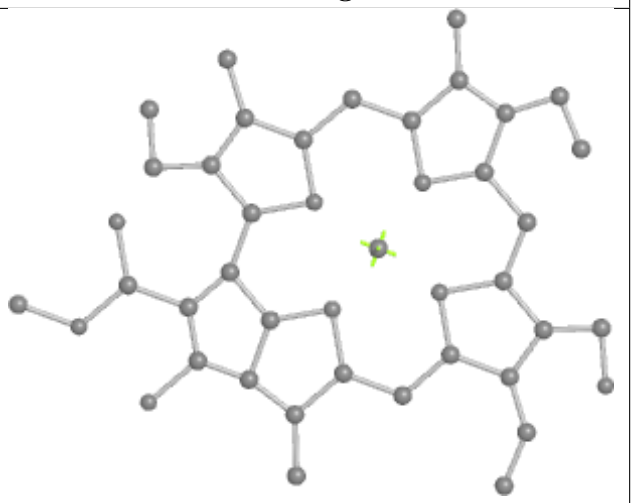


Bond angles

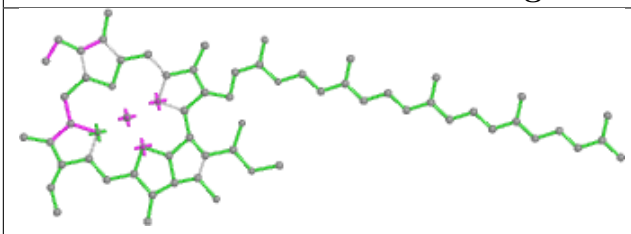


Torsions

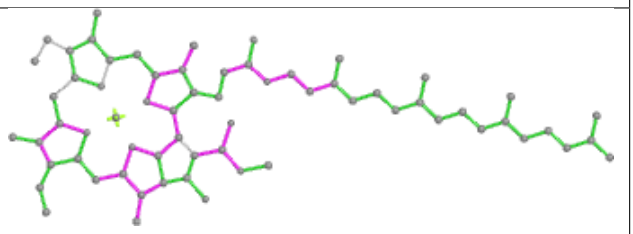


Rings

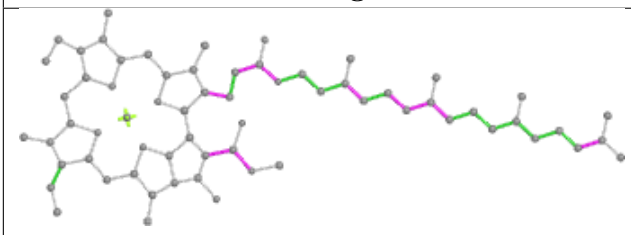
Ligand CLA B 842



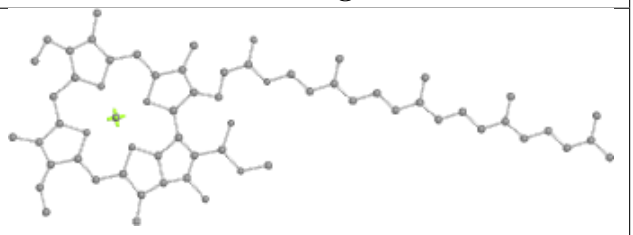
Bond lengths



Bond angles

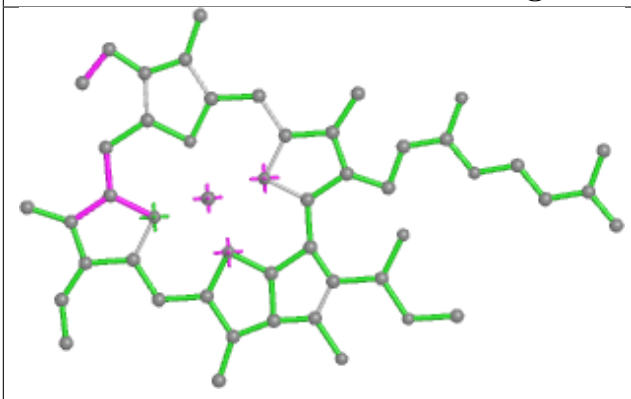


Torsions

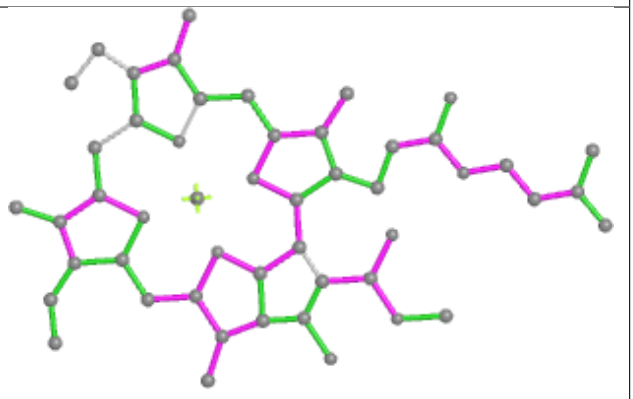


Rings

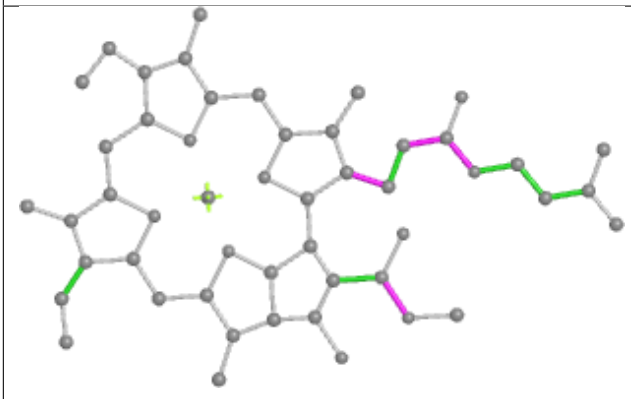
Ligand CLA L 203



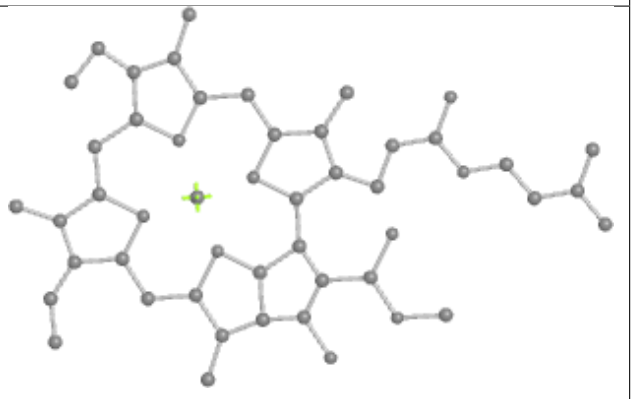
Bond lengths



Bond angles

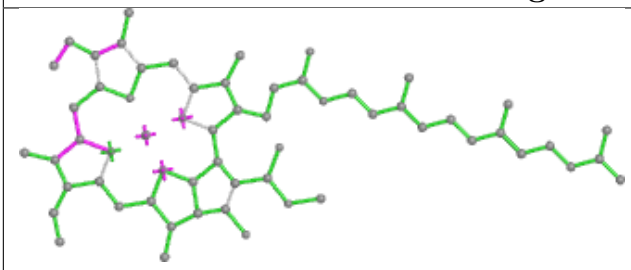


Torsions



Rings

Ligand CLA 8 310



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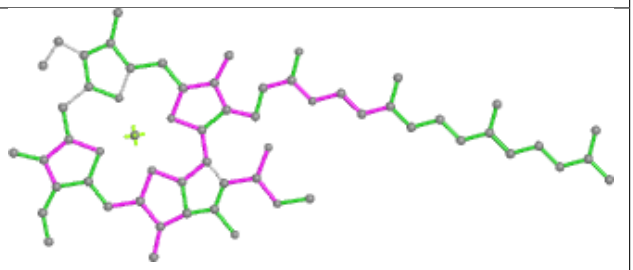
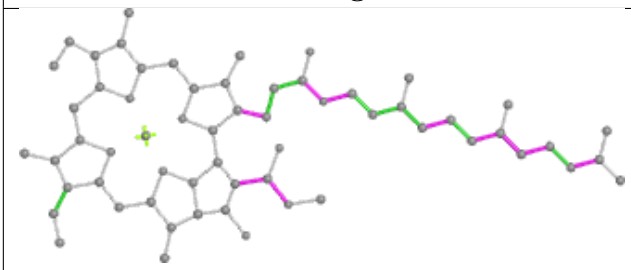
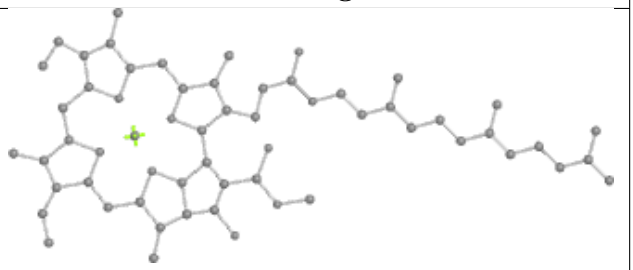


Bond angles

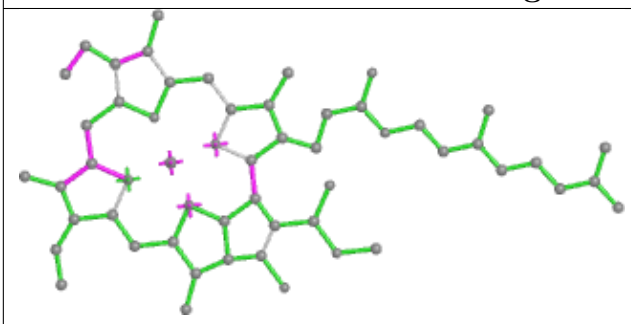


Torsions

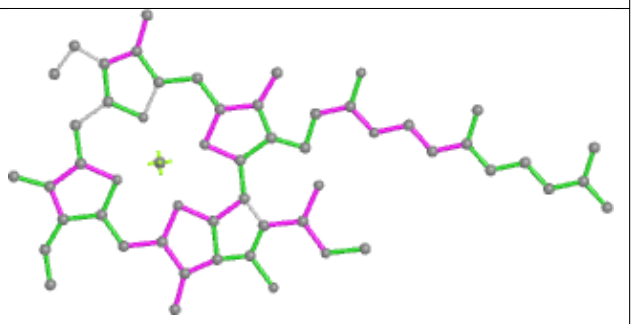


Rings

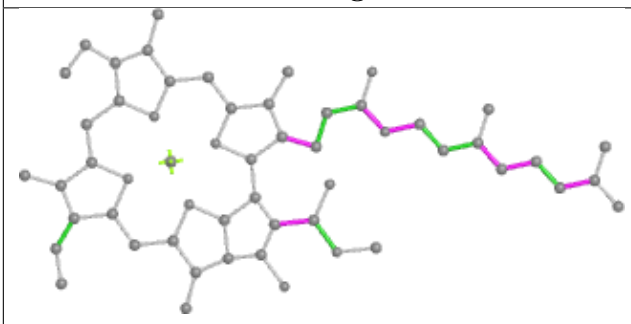
Ligand CLA A 822



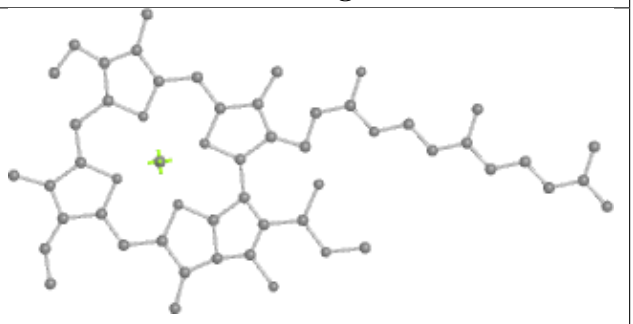
Bond lengths



Bond angles

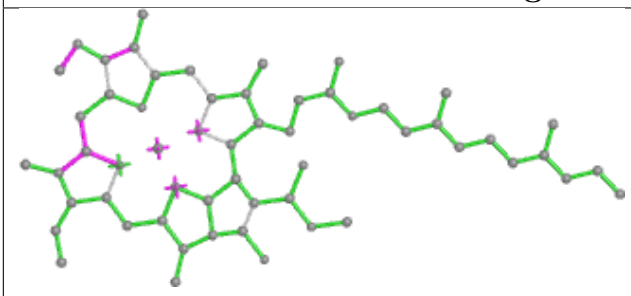


Torsions



Rings

Ligand CLA B 817



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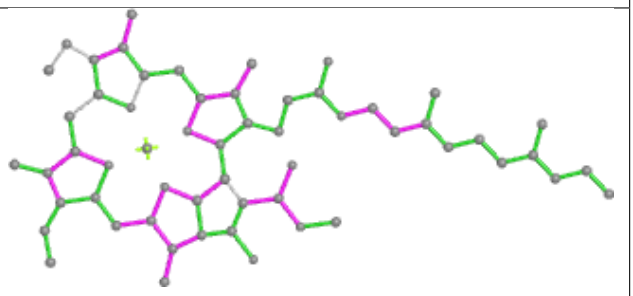
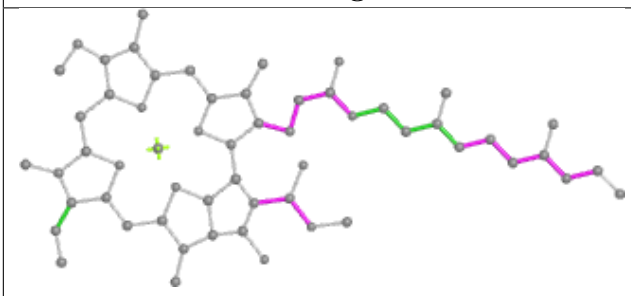


Bond angles



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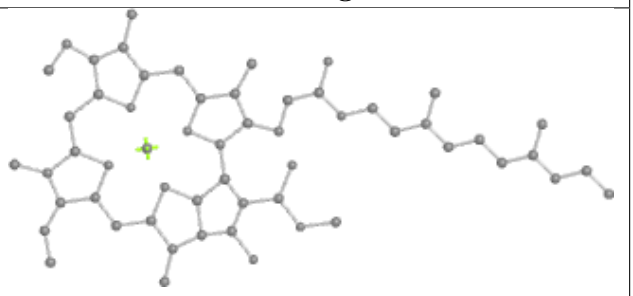
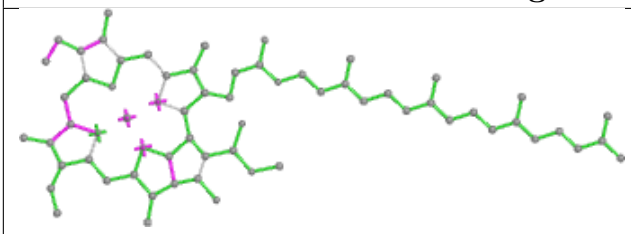


Rings

Ligand CLA 3 312



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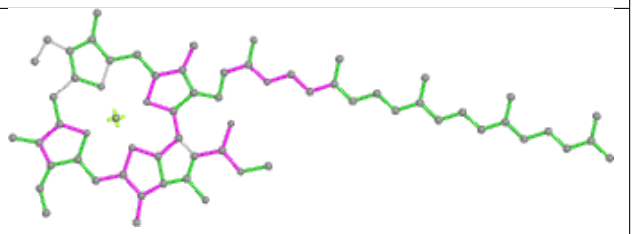
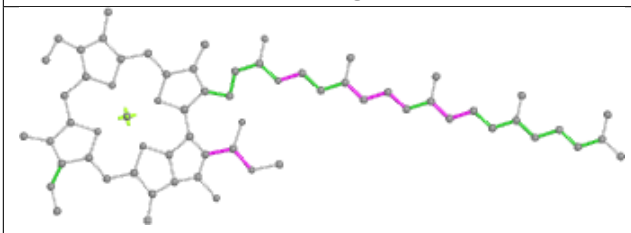
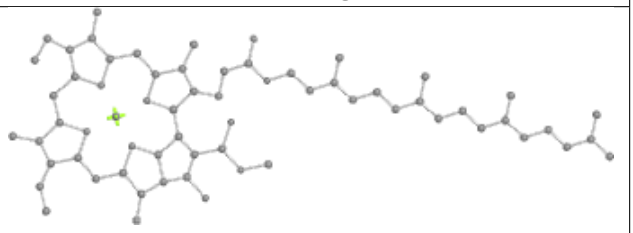


Bond angles

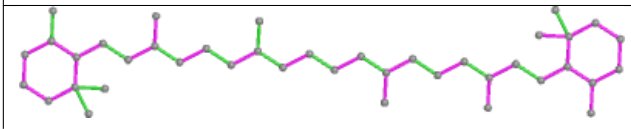


Torsions

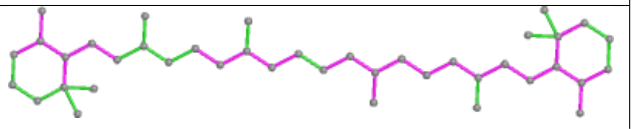


Rings

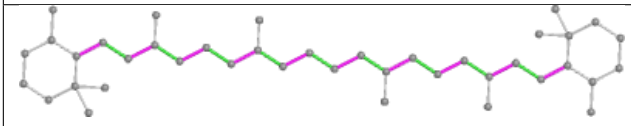
Ligand BCR F 303



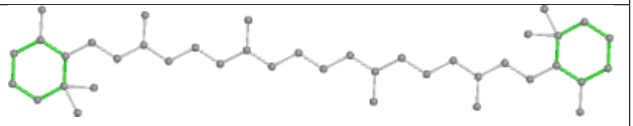
Bond lengths



Bond angles

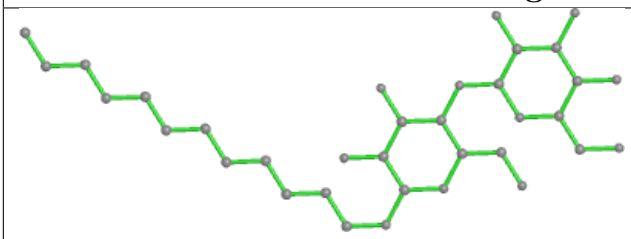


Torsions

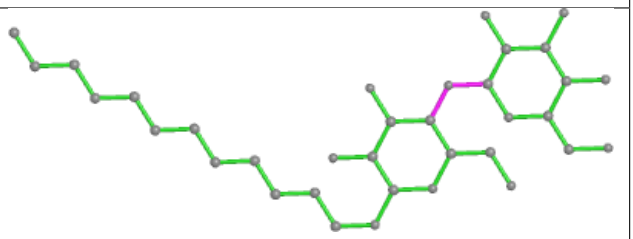


Rings

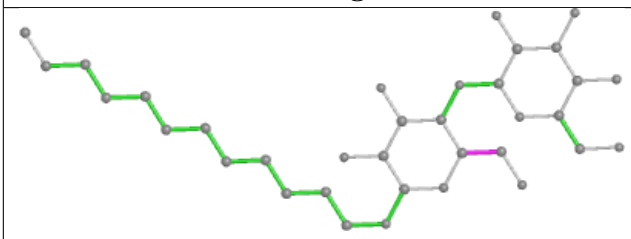
Ligand LMT 4 801



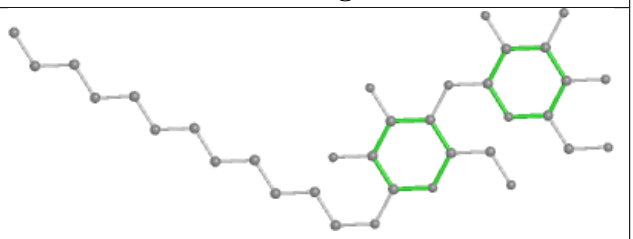
Bond lengths



Bond angles

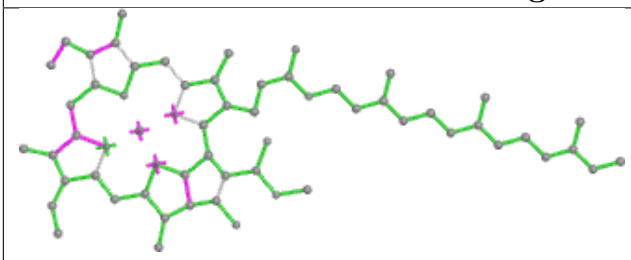


Torsions

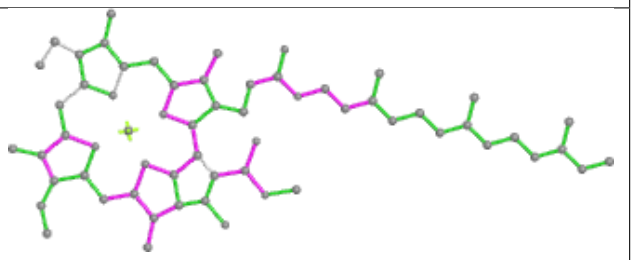


Rings

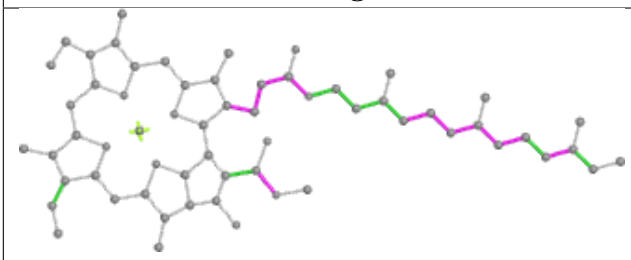
Ligand CLA 5 308



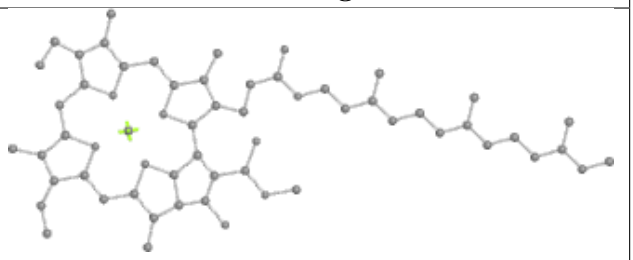
Bond lengths



Bond angles

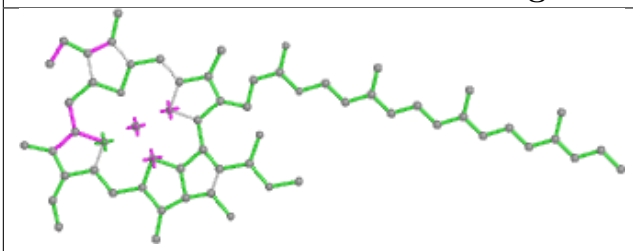


Torsions

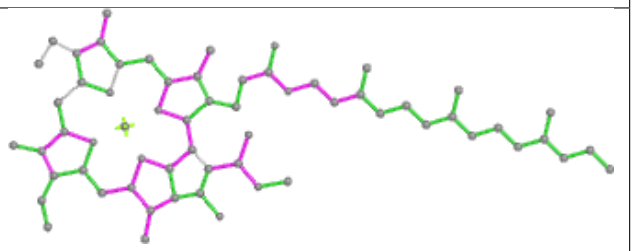


Rings

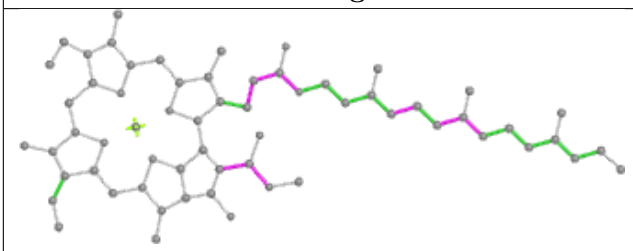
Ligand CLA 9 306



Bond lengths



Bond angles



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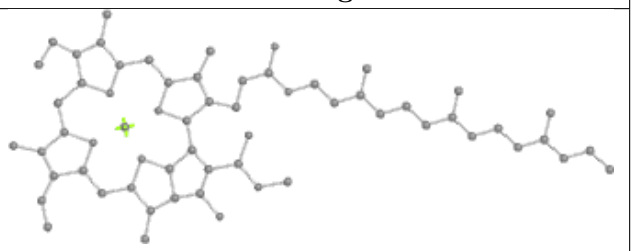
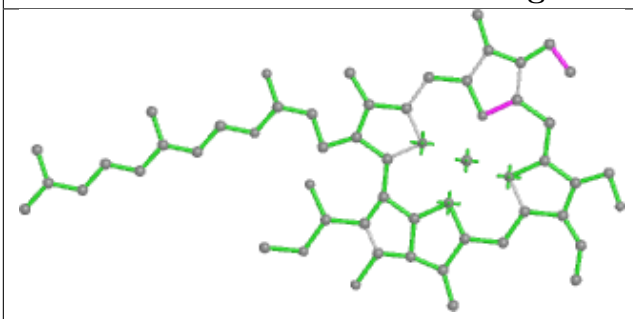
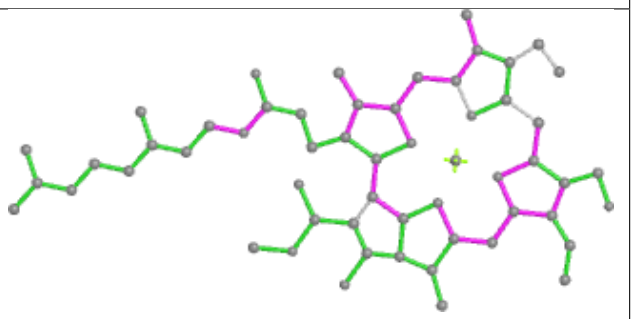


Rings

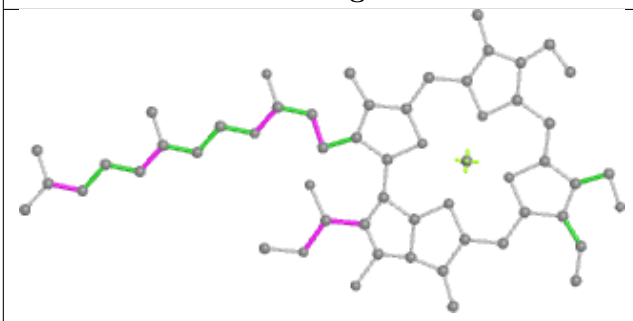
Ligand CHL 6 315



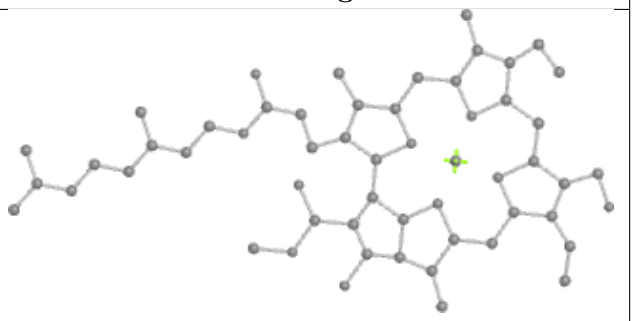
Bond lengths



Bond angles

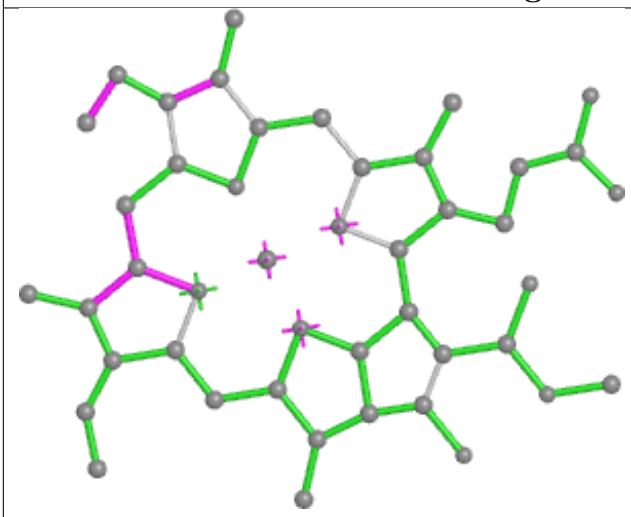


Torsions

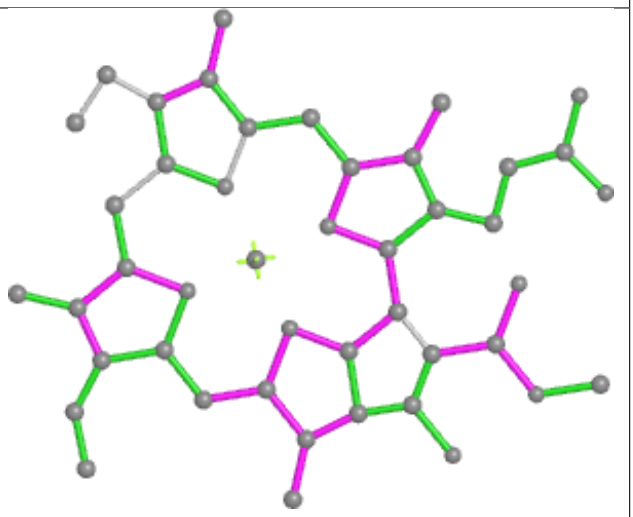


Rings

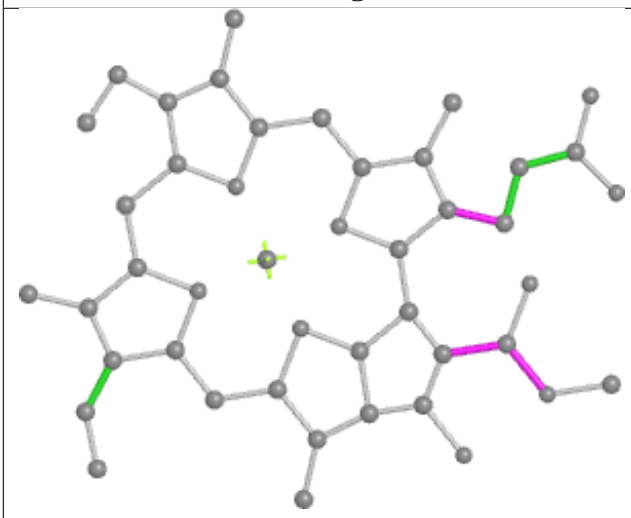
Ligand CLA B 836



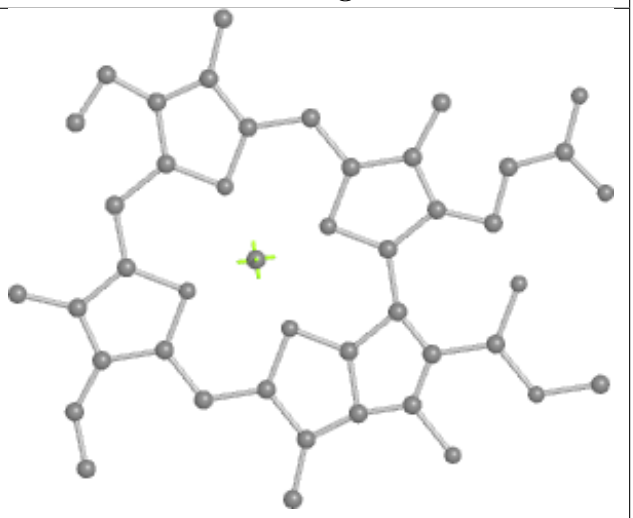
Bond lengths



Bond angles

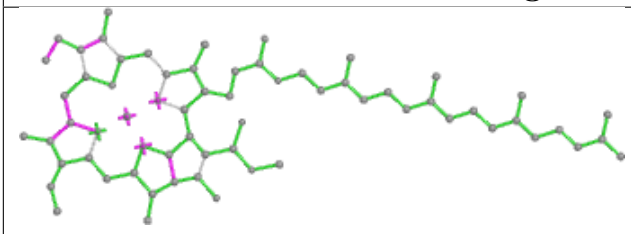


Torsions

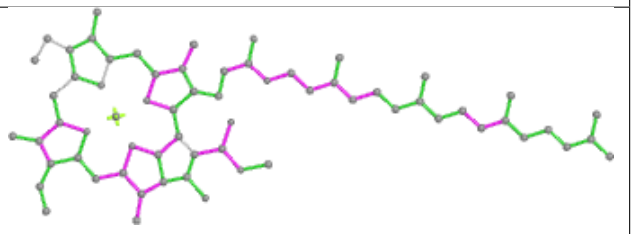


Rings

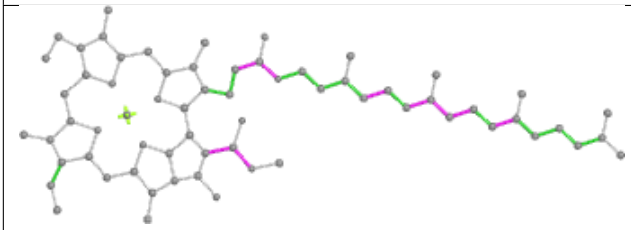
Ligand CLA A 823



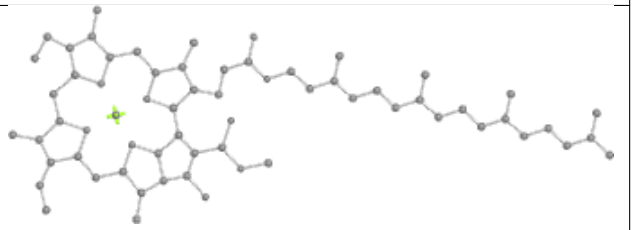
Bond lengths



Bond angles

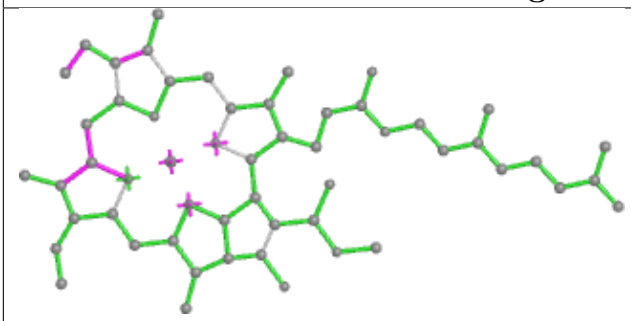


Torsions

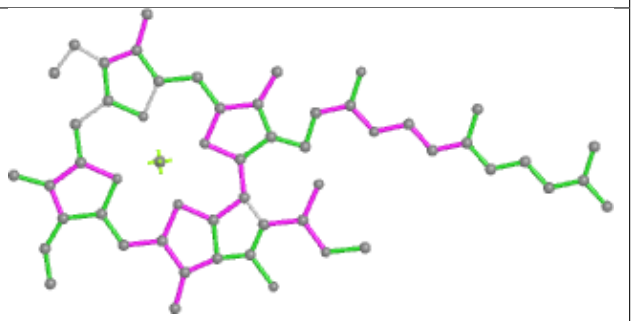


Rings

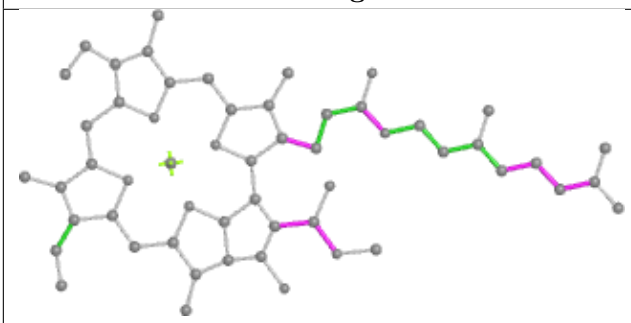
Ligand CLA 6 311



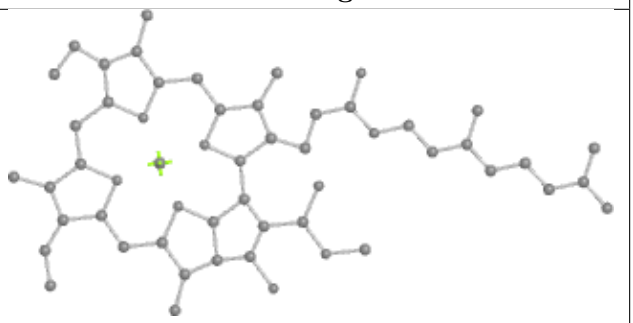
Bond lengths



Bond angles

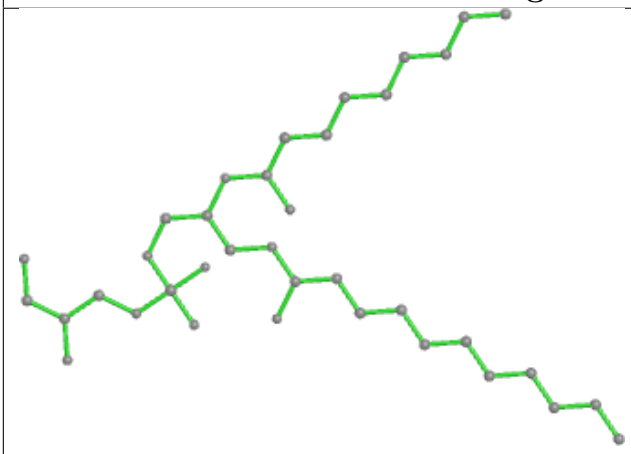


Torsions

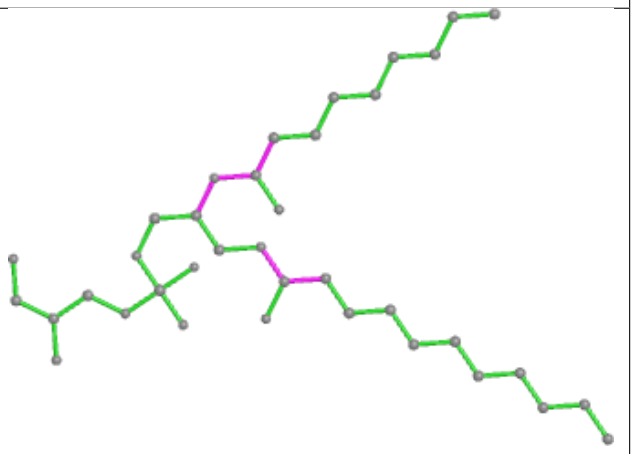


Rings

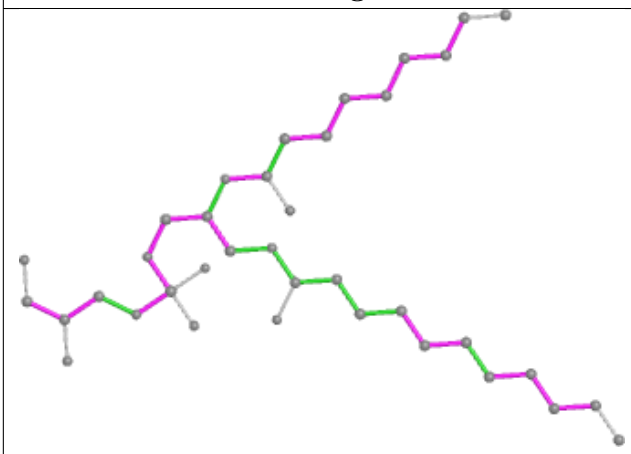
Ligand LHG 7 318



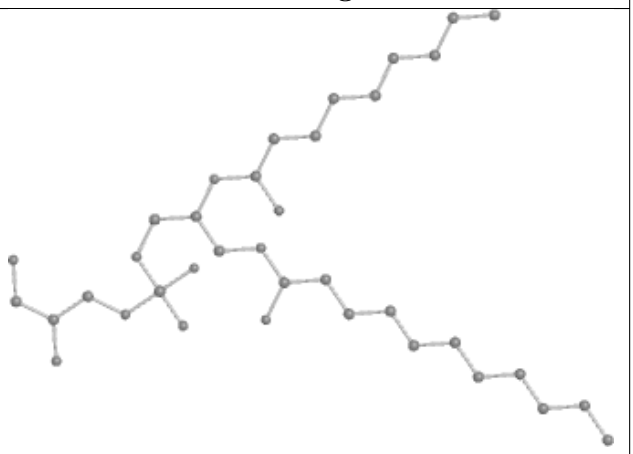
Bond lengths



Bond angles

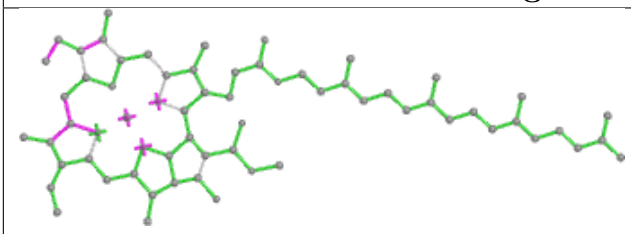


Torsions

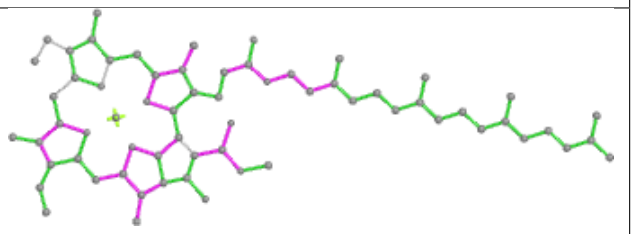


Rings

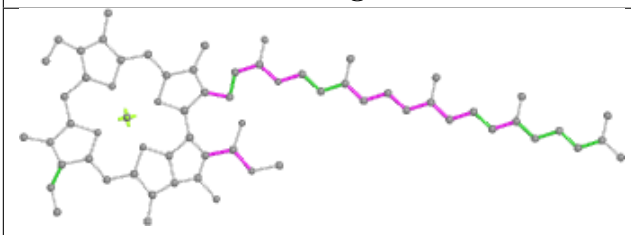
Ligand CLA A 828



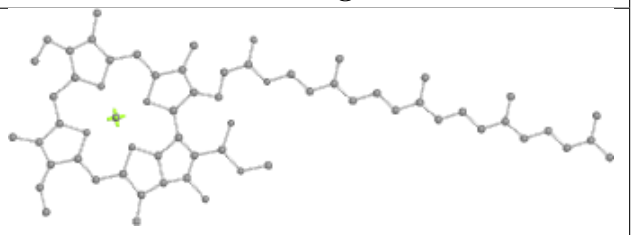
Bond lengths



Bond angles

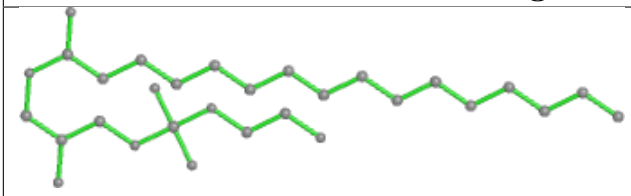


Torsions

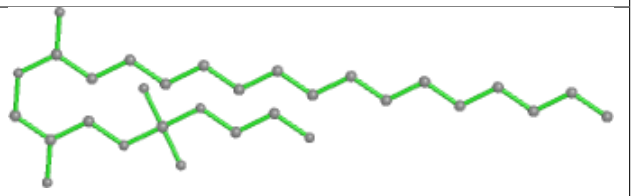


Rings

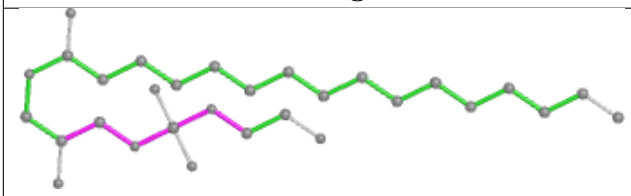
Ligand LPX 8 319



Bond lengths



Bond angles



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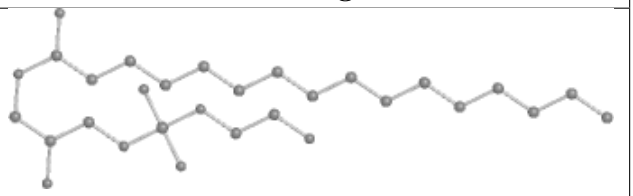
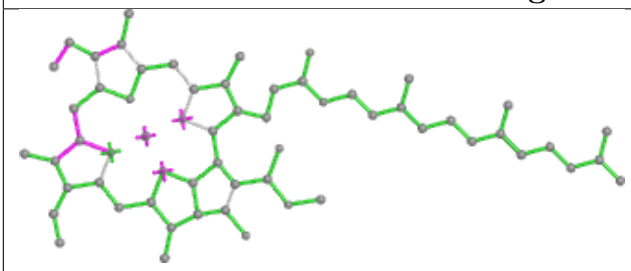
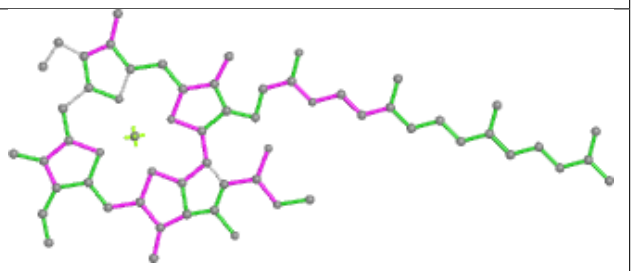


Rings

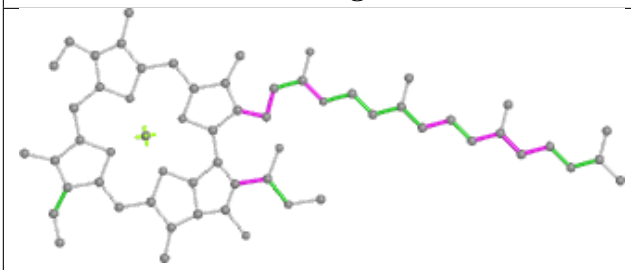
Ligand CLA 3 311



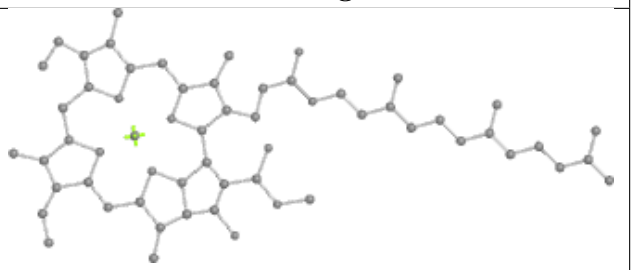
Bond lengths



Bond angles

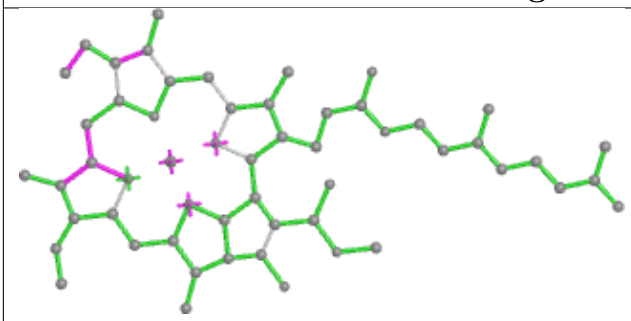


Torsions

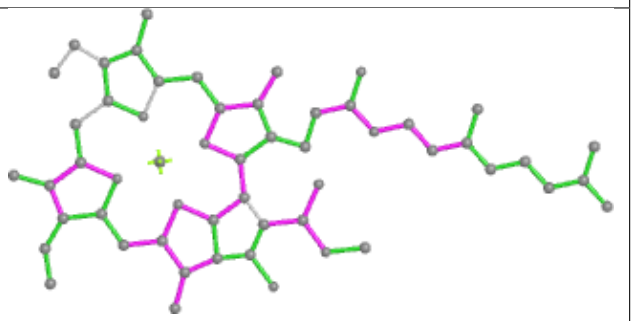


Rings

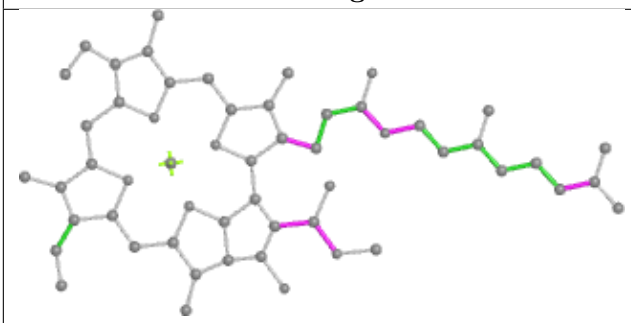
Ligand CLA 4 811



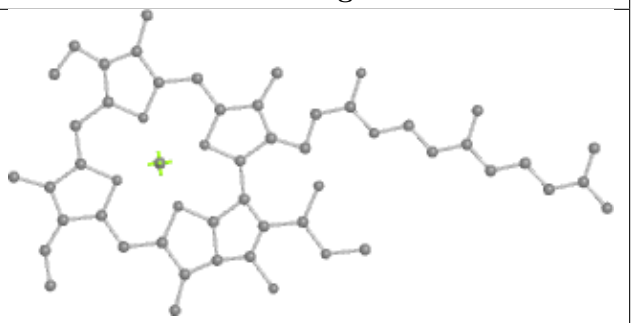
Bond lengths



Bond angles

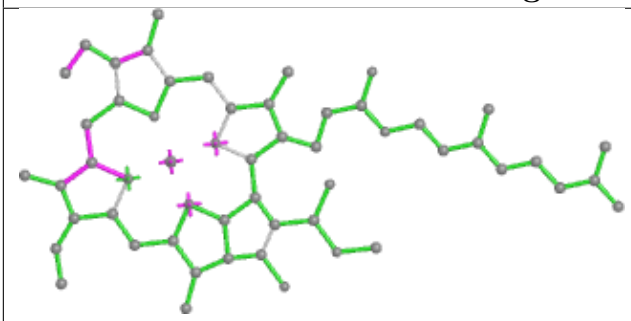


Torsions

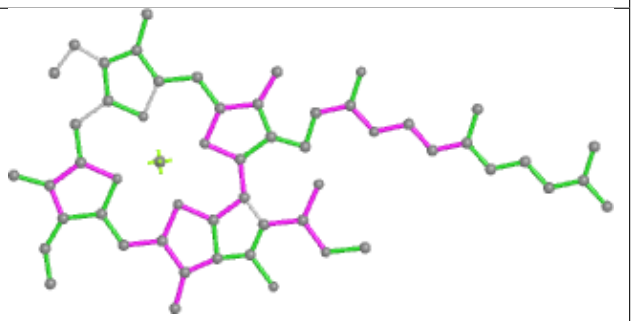


Rings

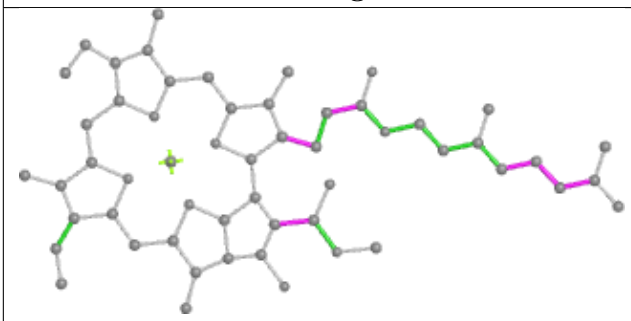
Ligand CLA K 203



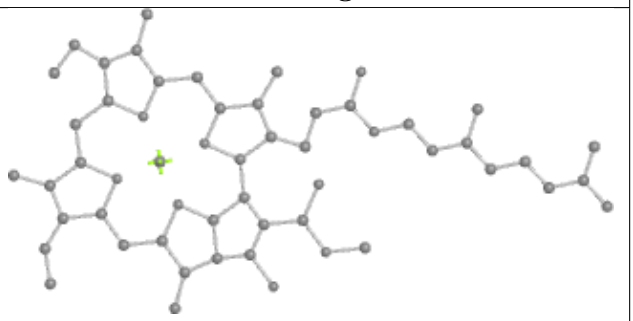
Bond lengths



Bond angles

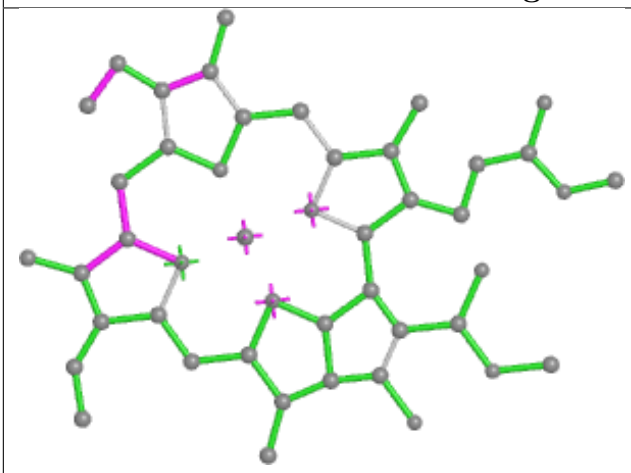


Torsions

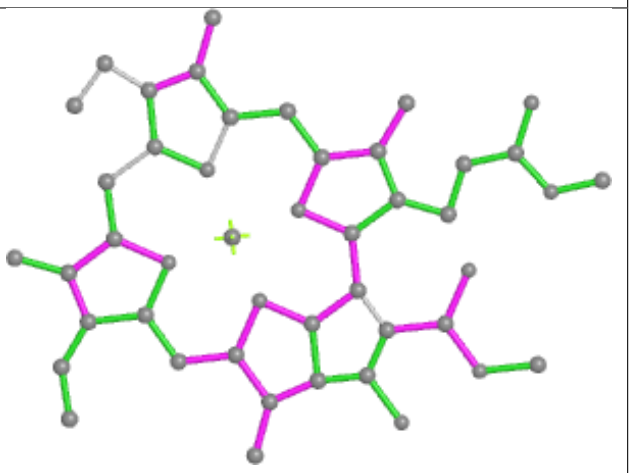


Rings

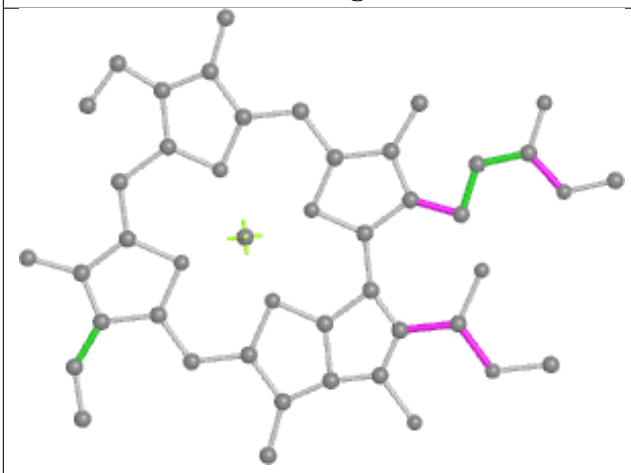
Ligand CLA G 202



Bond lengths



Bond angles



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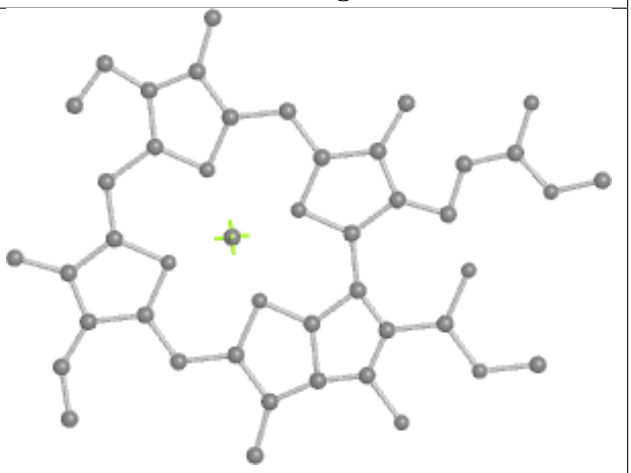
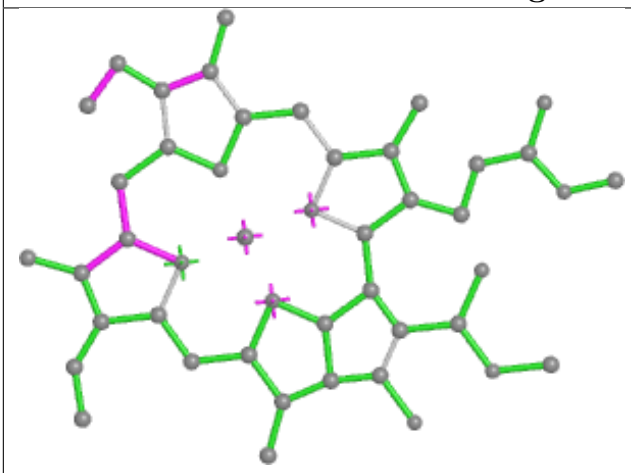
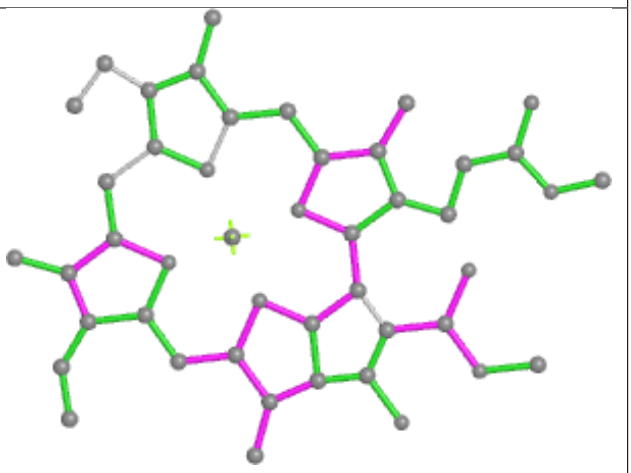


Rings

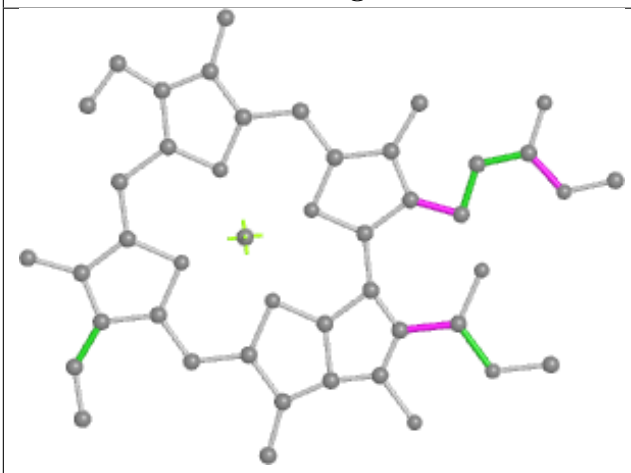
Ligand CLA 6 321



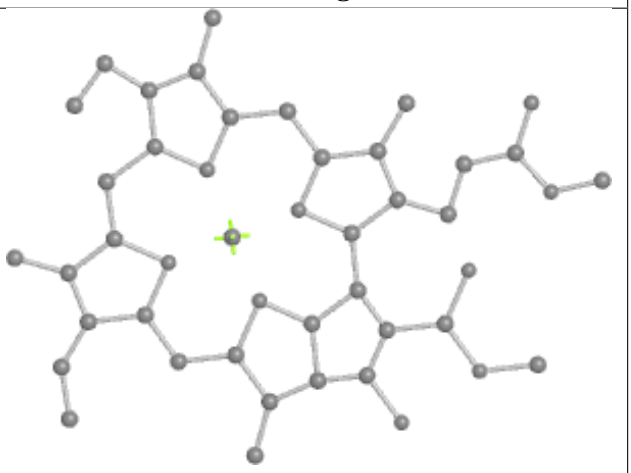
Bond lengths



Bond angles

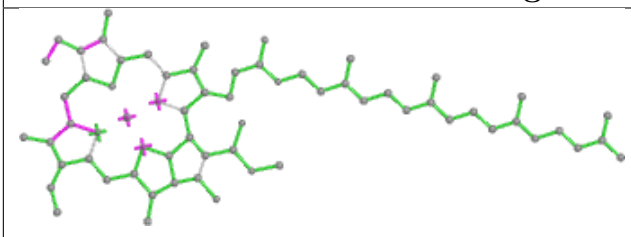


Torsions

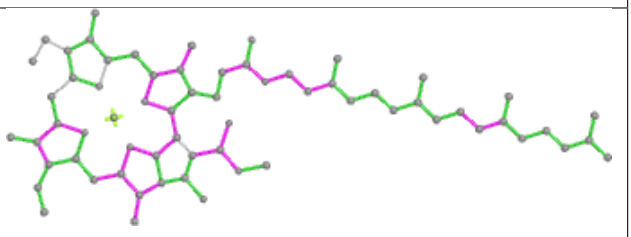


Rings

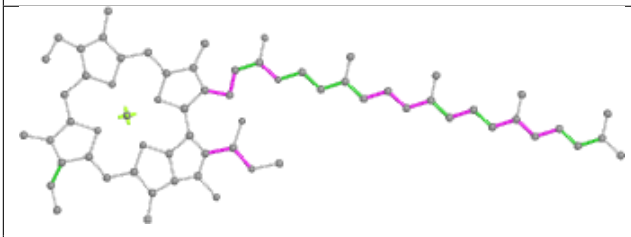
Ligand CLA A 808



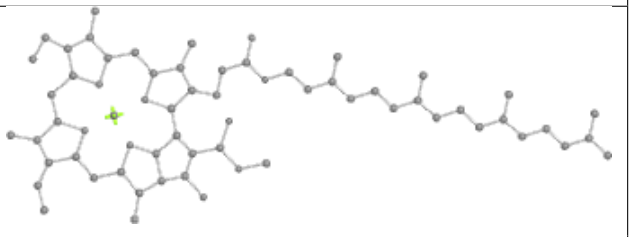
Bond lengths



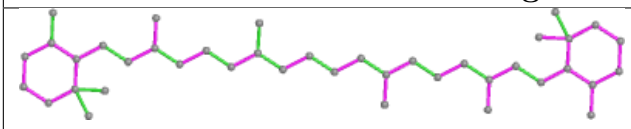
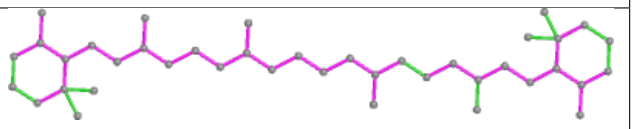
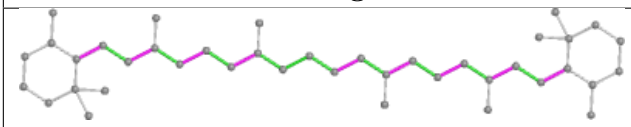
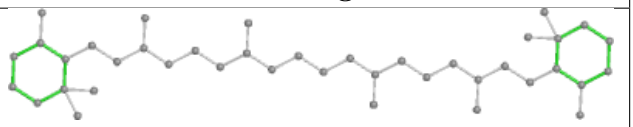
Bond angles

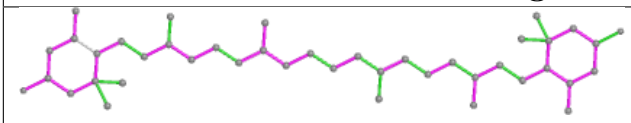
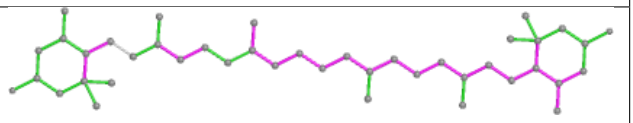
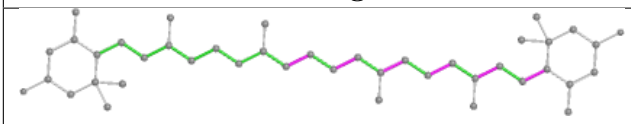
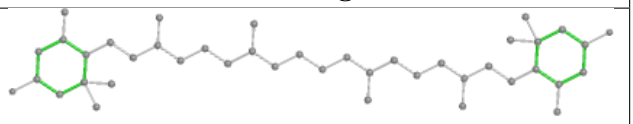


Torsions

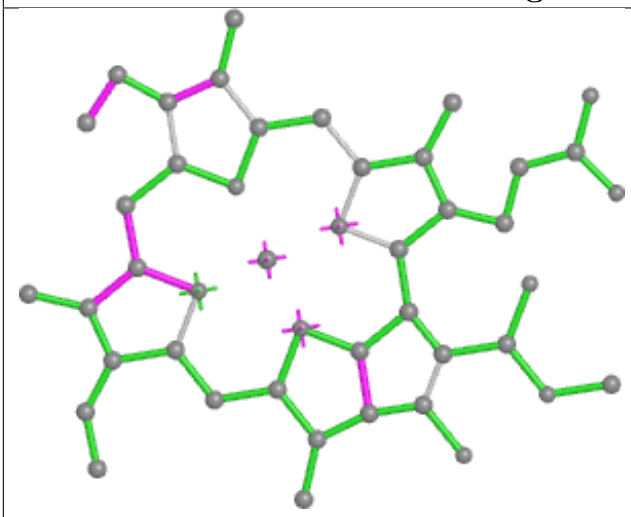


Rings

Ligand BCR A 846	
	
Bond lengths	Bond angles
	
Torsions	Rings

Ligand LUT 4 803	
	
Bond lengths	Bond angles
	
Torsions	Rings

Ligand CLA F 305



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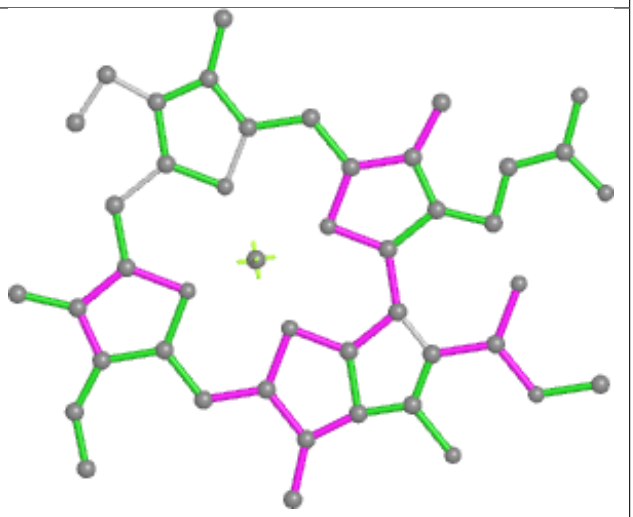

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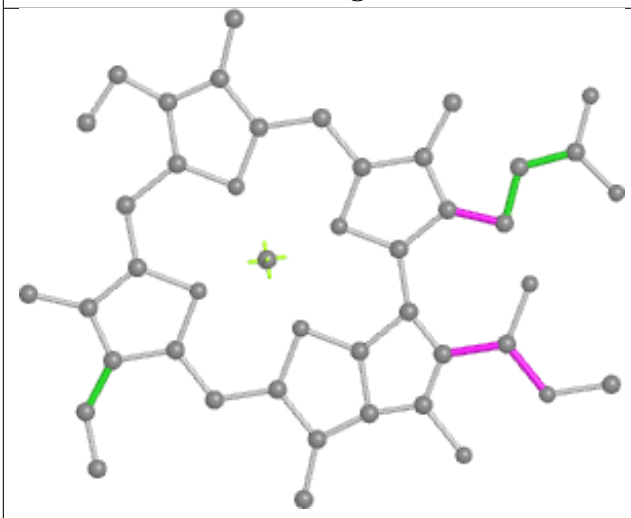

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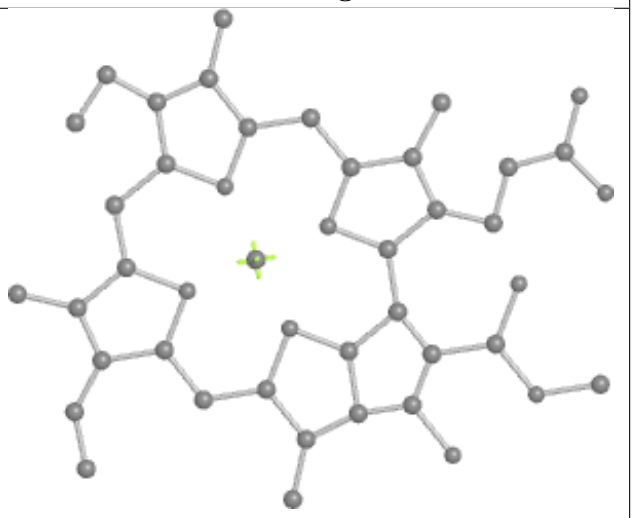
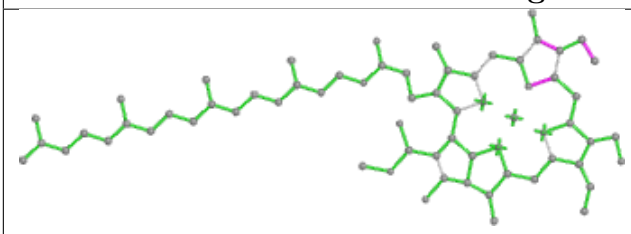


Rings

Ligand CHL 9 312



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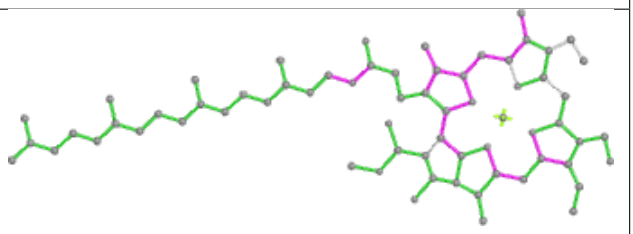

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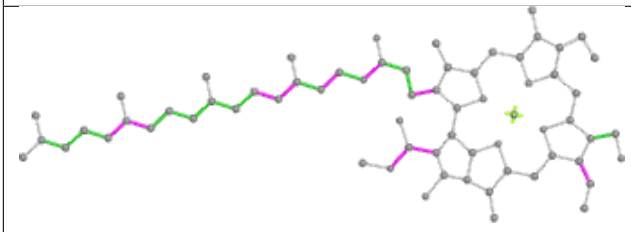

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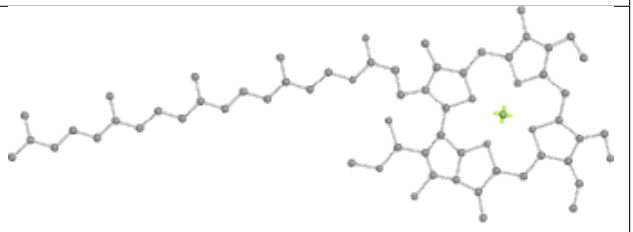
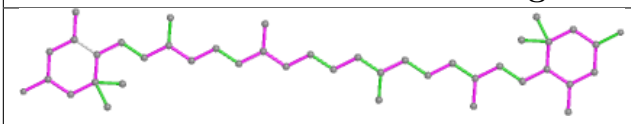


Rings

Ligand LUT 5 302



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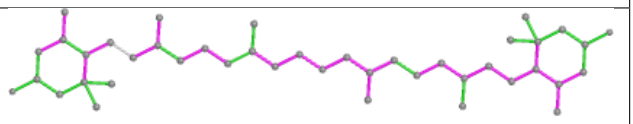
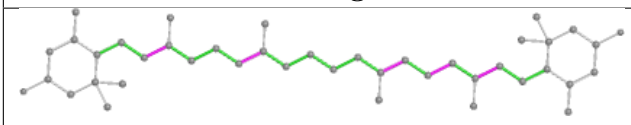


Bond angles



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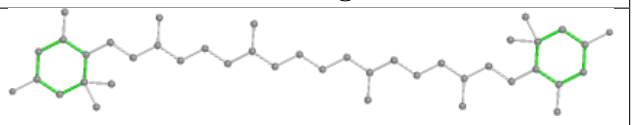
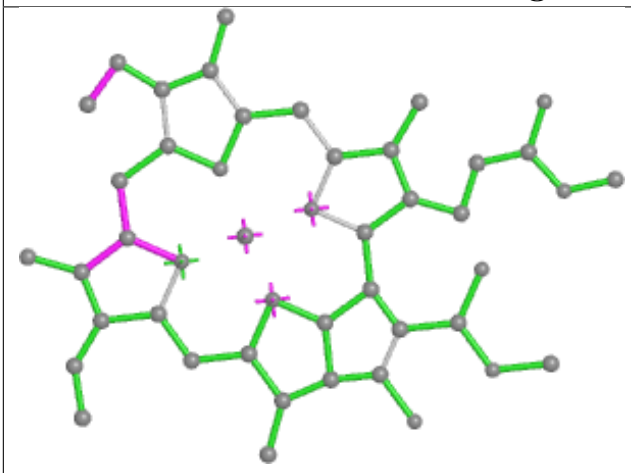


Rings

Ligand CLA 1 315



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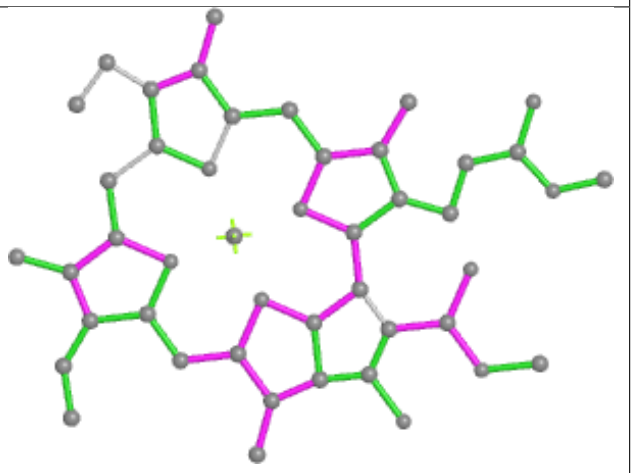
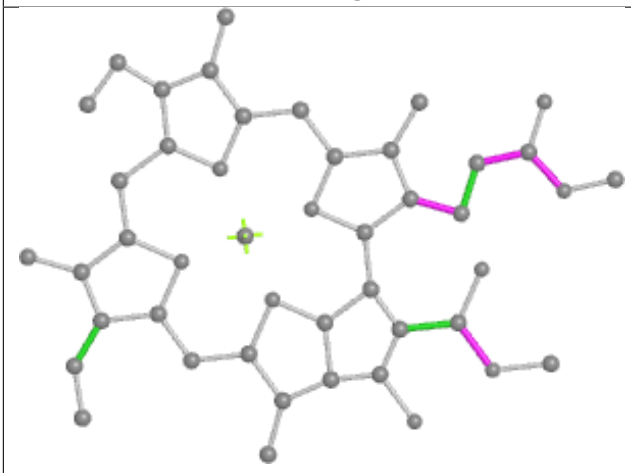


Bond angles



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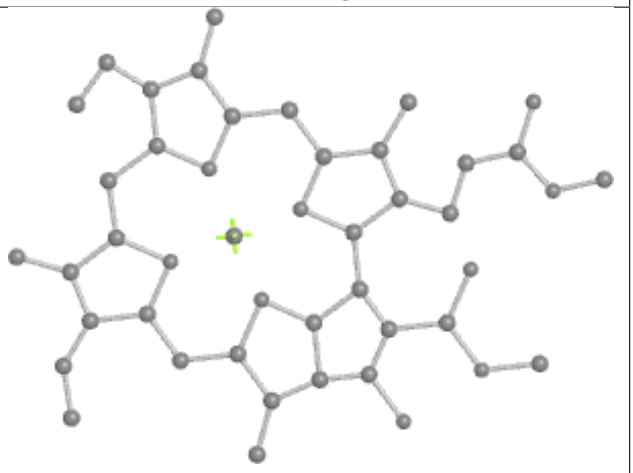
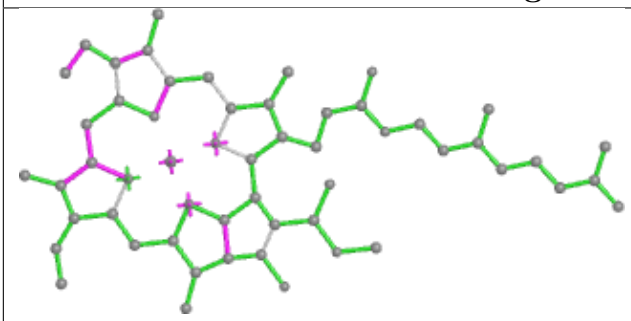


Rings

Ligand CLA B 823



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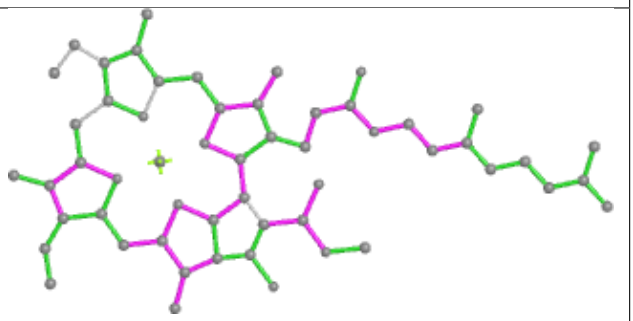
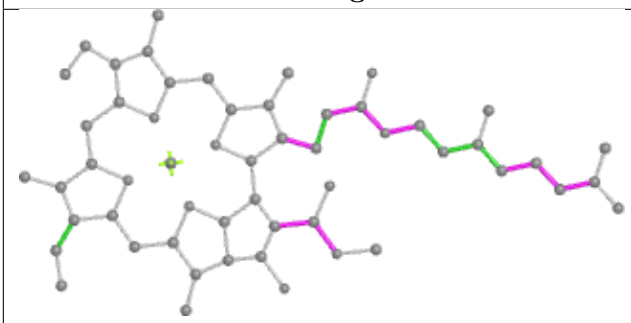


Bond angles



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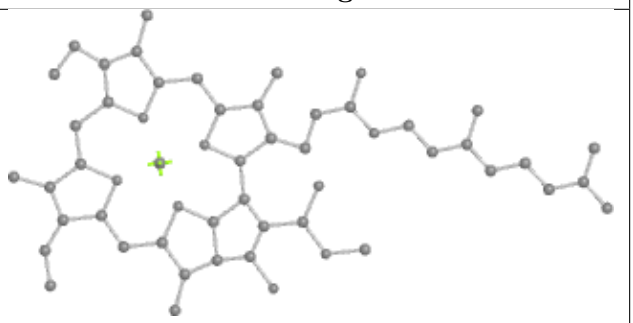
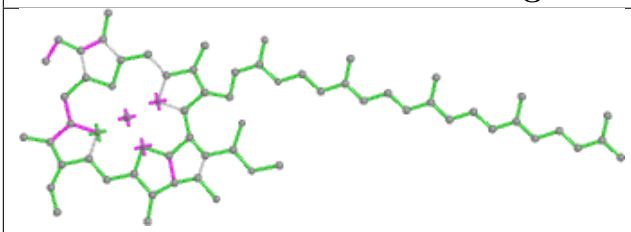


Rings

Ligand CLA A 840



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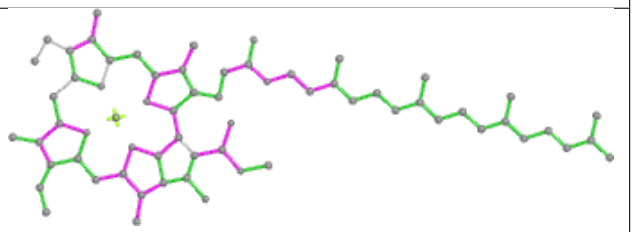
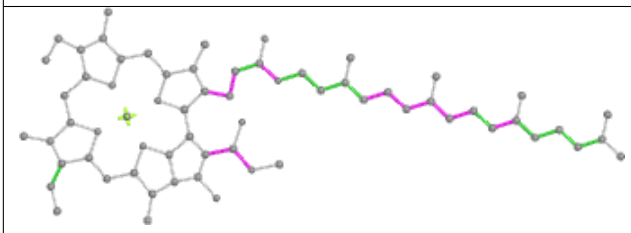


Bond angles



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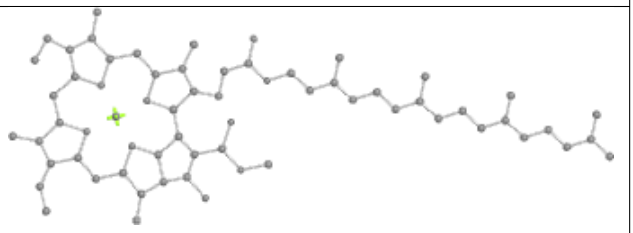
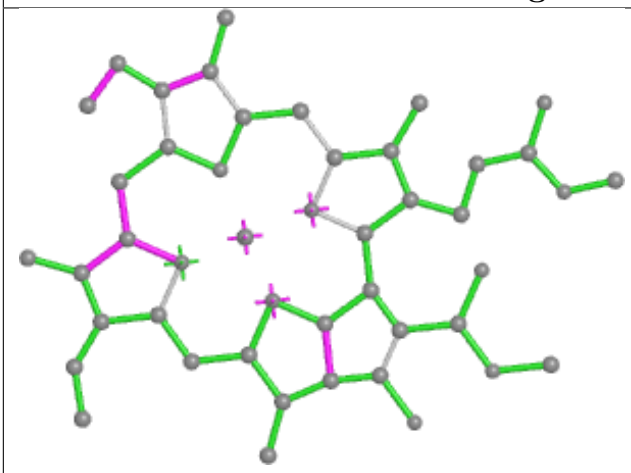


Rings

Ligand CLA 8 316



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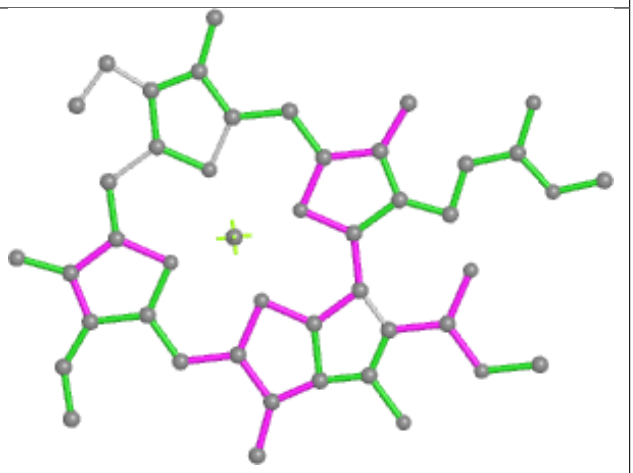
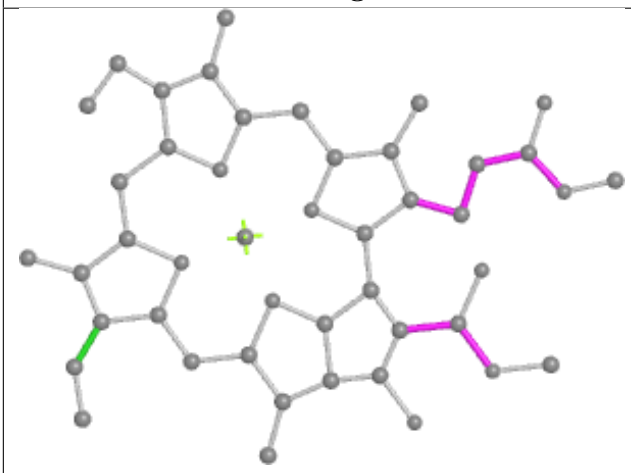


Bond angles



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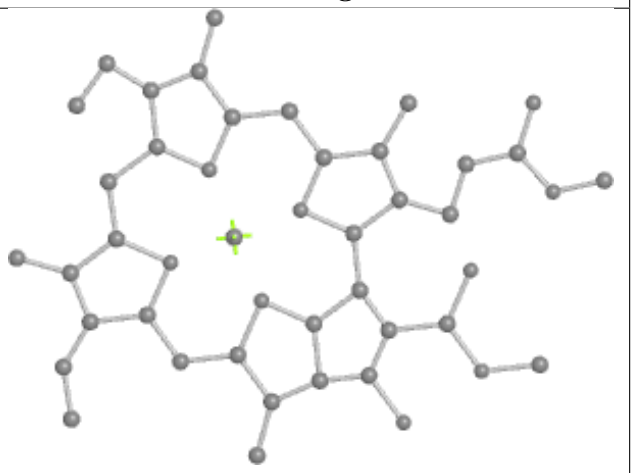
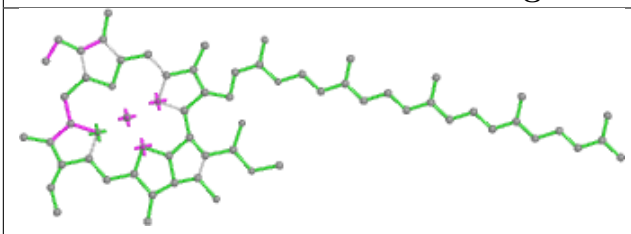


Rings

Ligand CLA A 825



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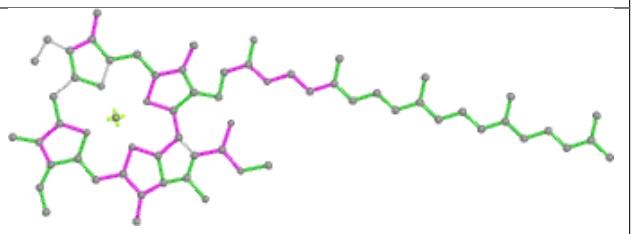
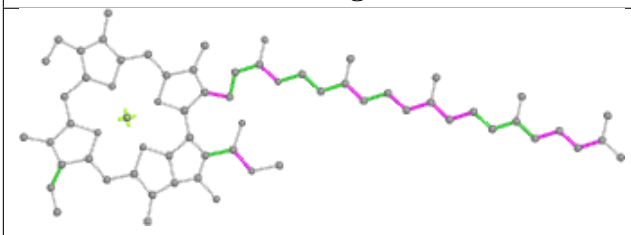
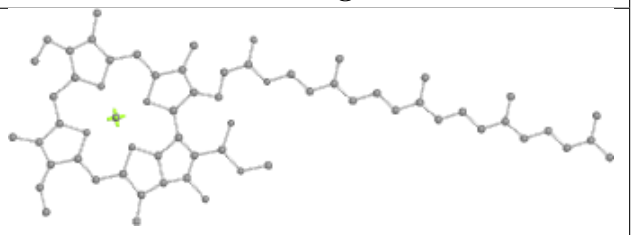


Bond angles

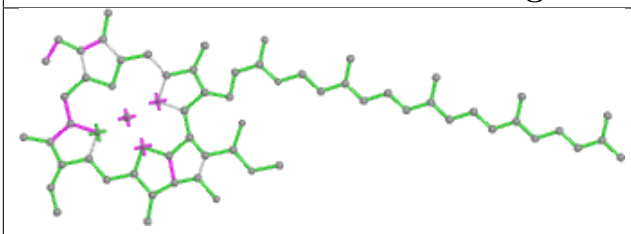


Torsions

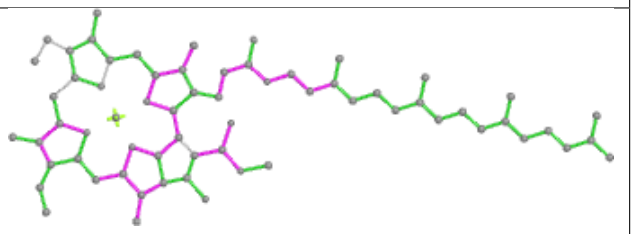


Rings

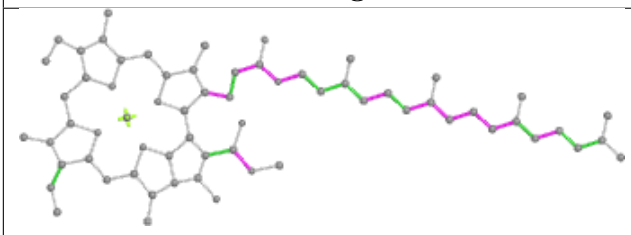
Ligand CLA F 302



Bond lengths



Bond angles



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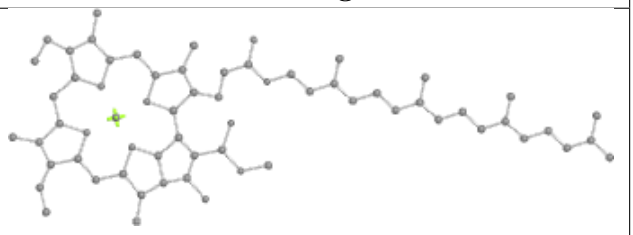
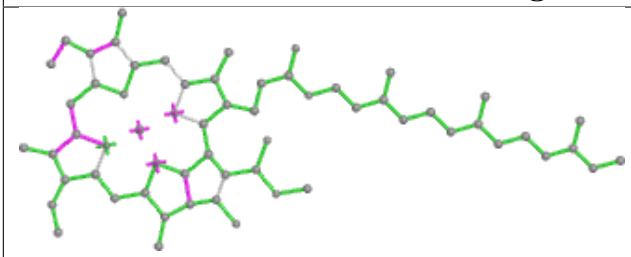


Rings

Ligand CLA 1 309



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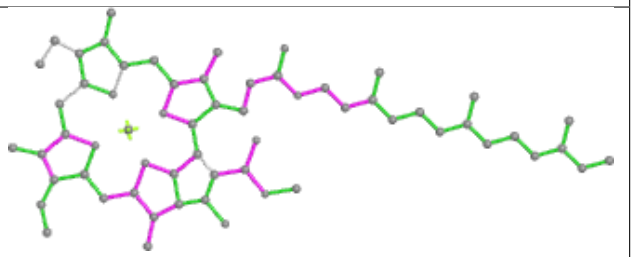
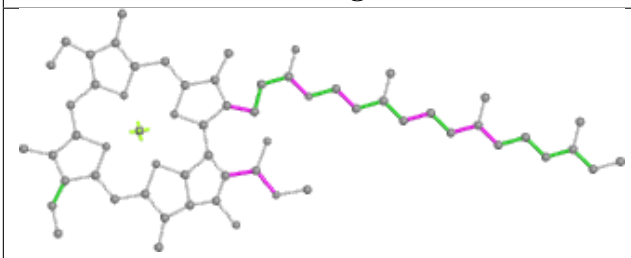
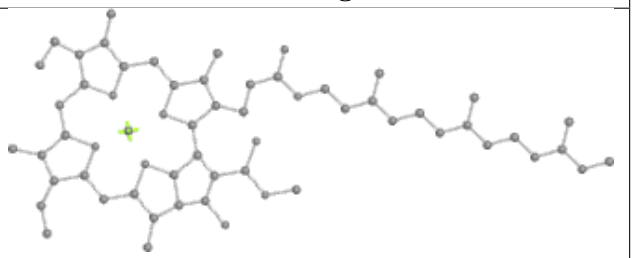


Bond angles

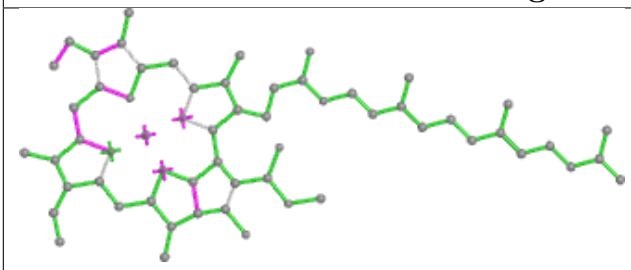


Torsions



Rings

Ligand CLA 9 303



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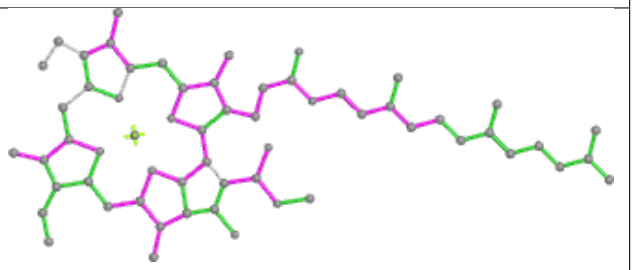
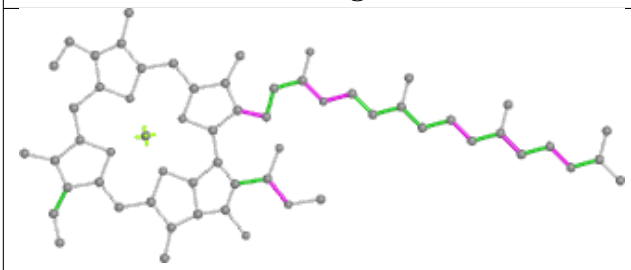
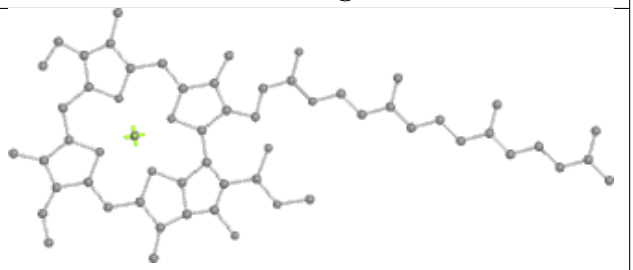


Bond angles

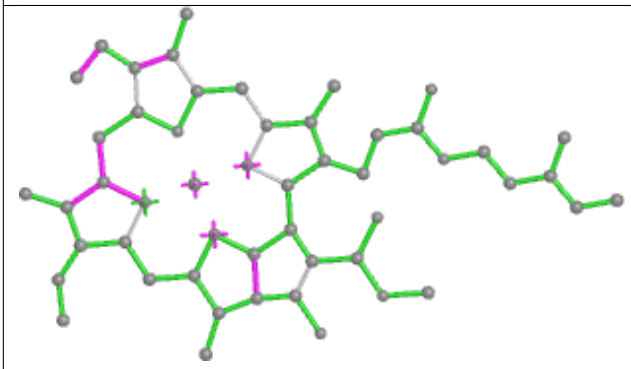


Torsions

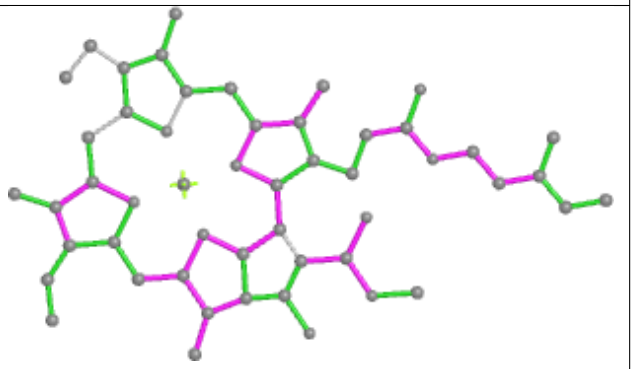


Rings

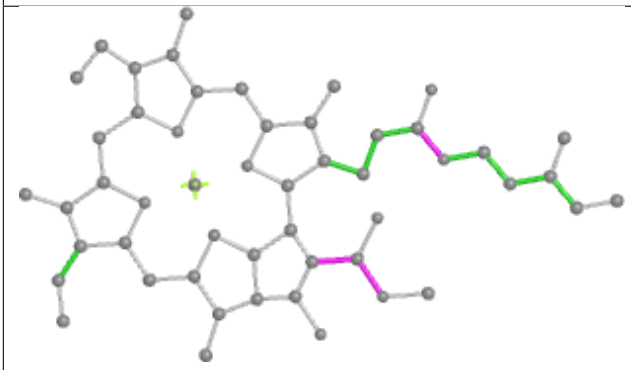
Ligand CLA 4 809



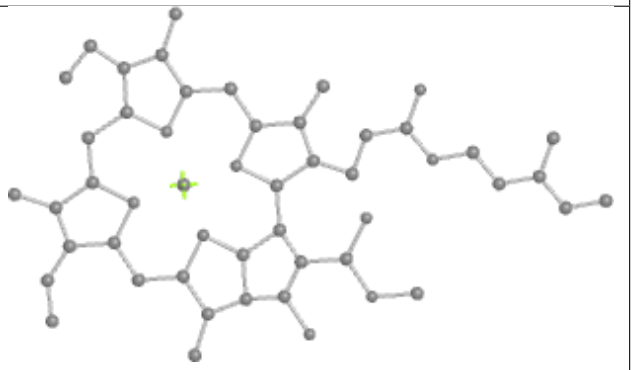
Bond lengths



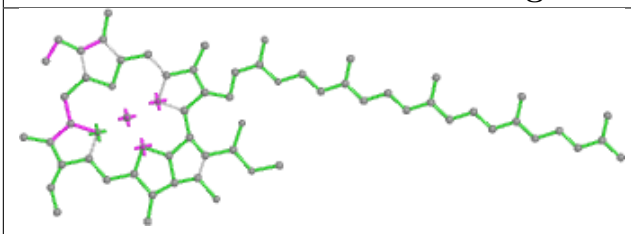
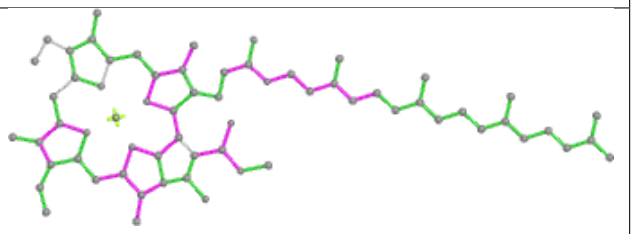
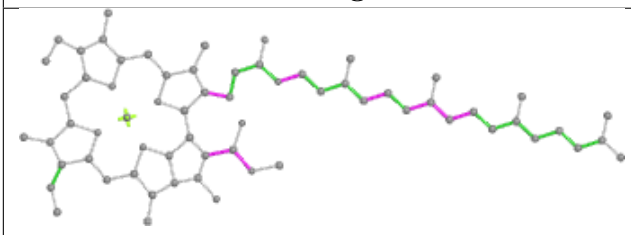
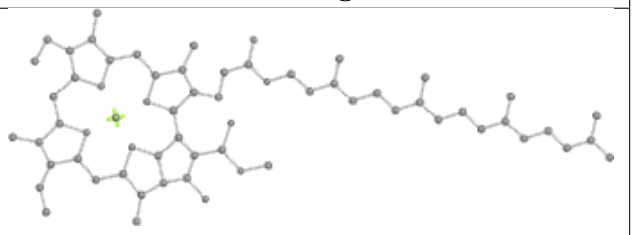
Bond angles

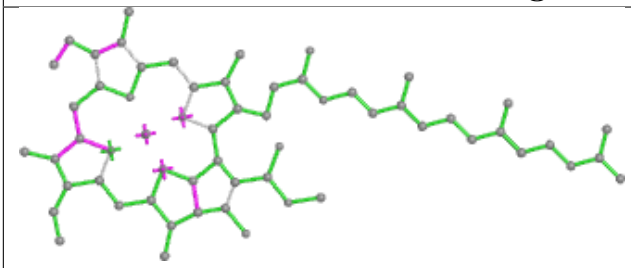
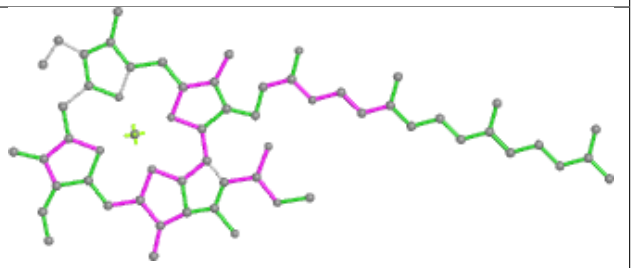
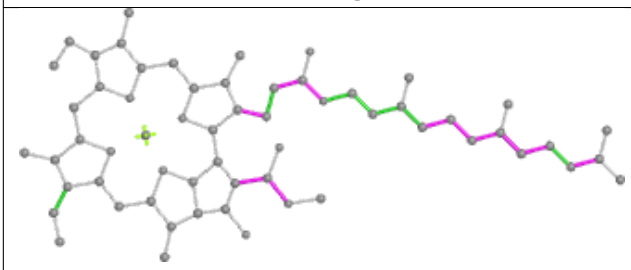
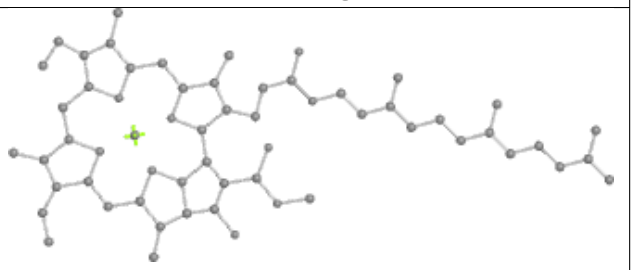


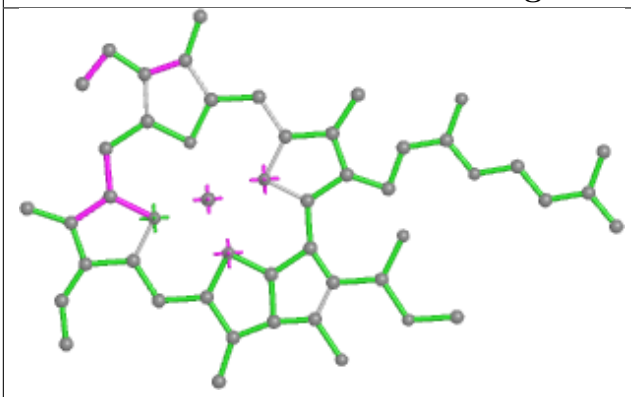
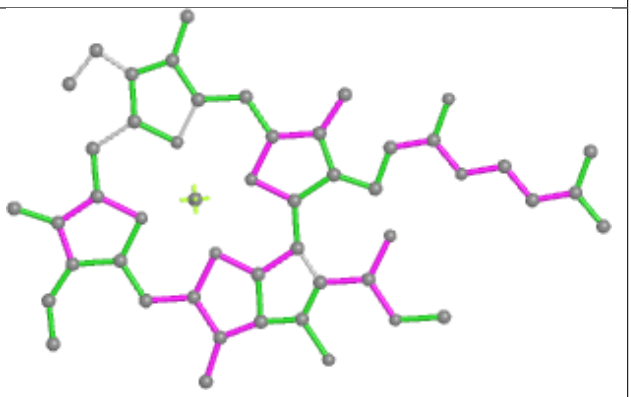
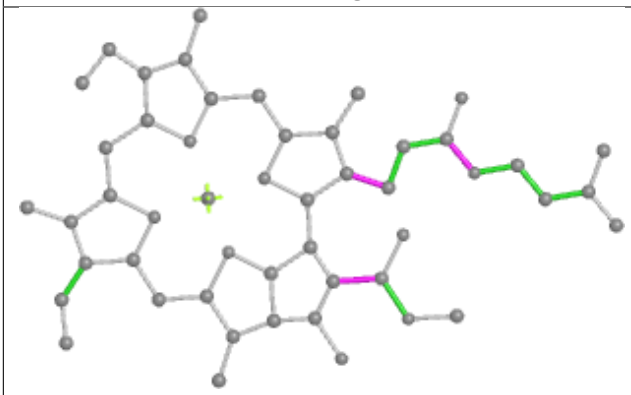
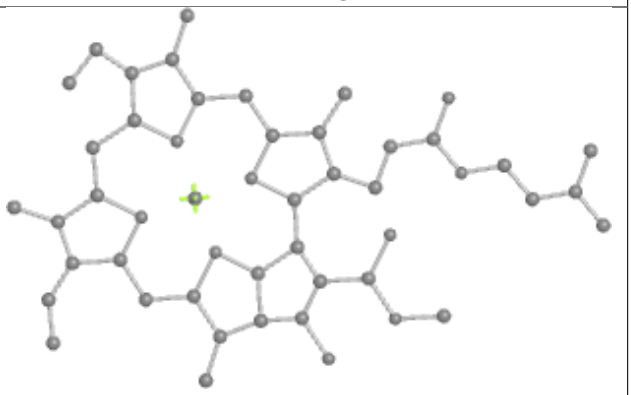
Torsions



Rings

Ligand CLA F 301	
 Bond lengths	 Bond angles
 Torsions	 Rings

Ligand CLA 1 307	
 Bond lengths	 Bond angles
 Torsions	 Rings

Ligand CLA B 832	
 Bond lengths	 Bond angles
 Torsions	 Rings

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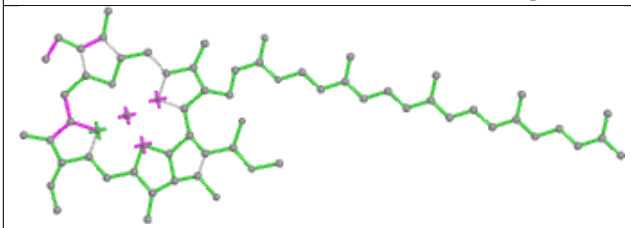
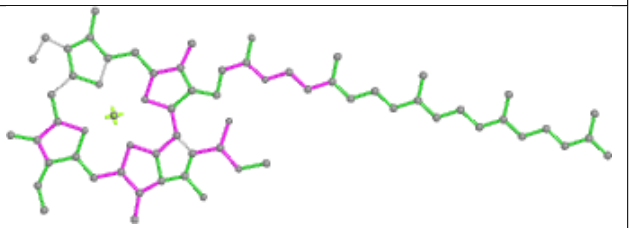
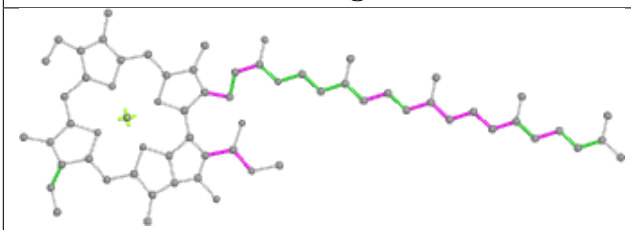
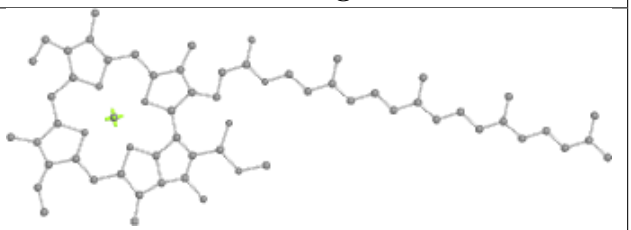
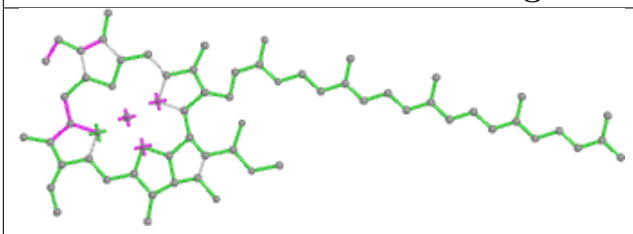
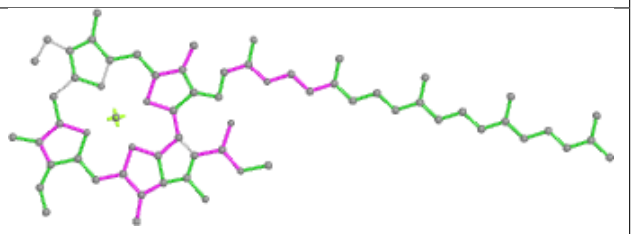
 Bond lengths	 Bond angles
 Torsions	 Rings

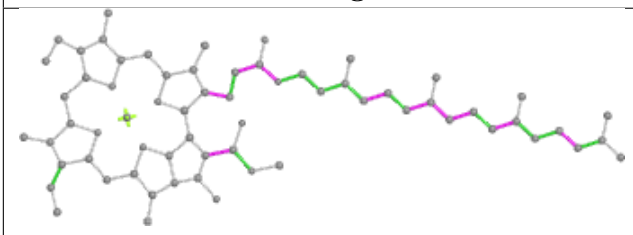
Ligand CLA A 827



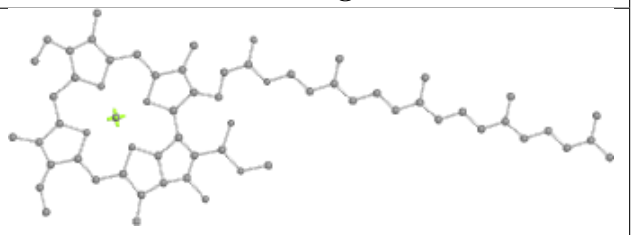
Bond lengths



Bond angles

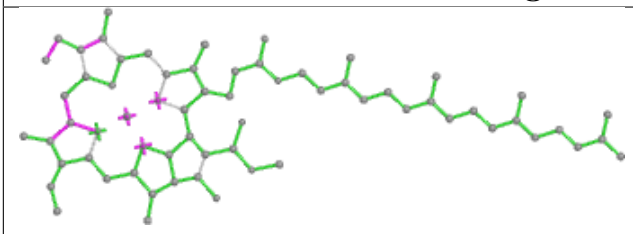


Torsions

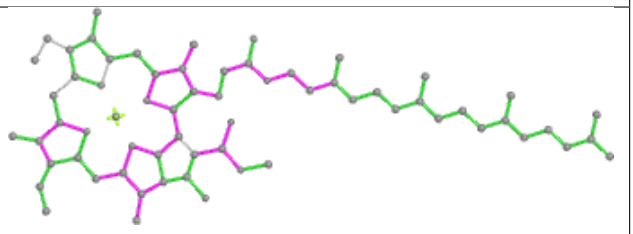


Rings

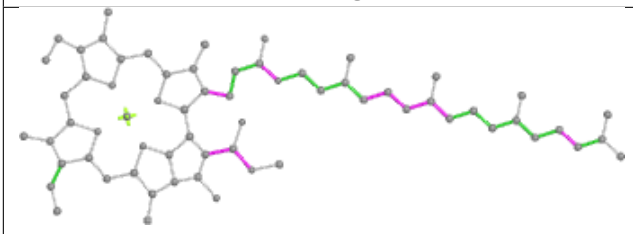
Ligand CLA B 811



Bond lengths



Bond angles



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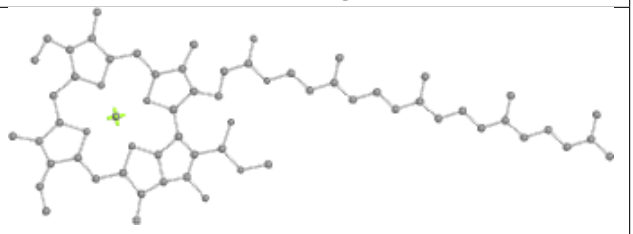
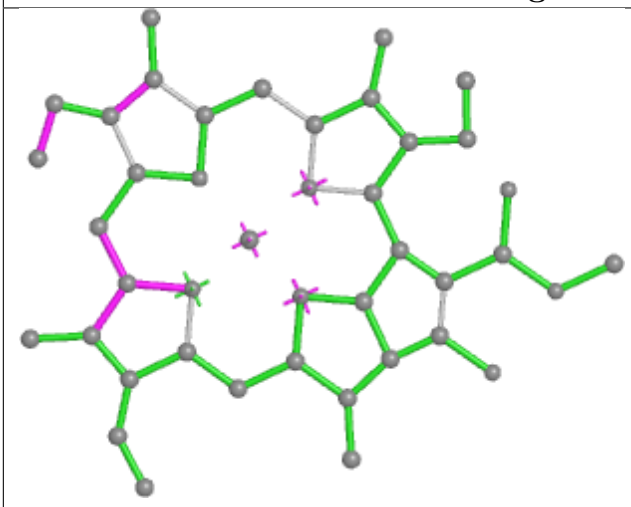
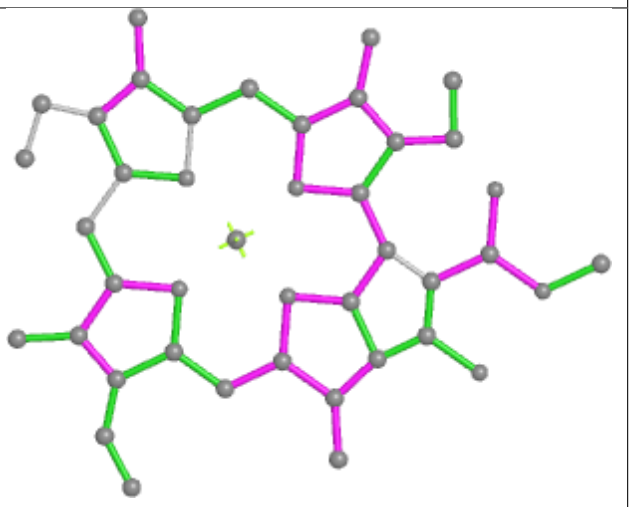


Rings

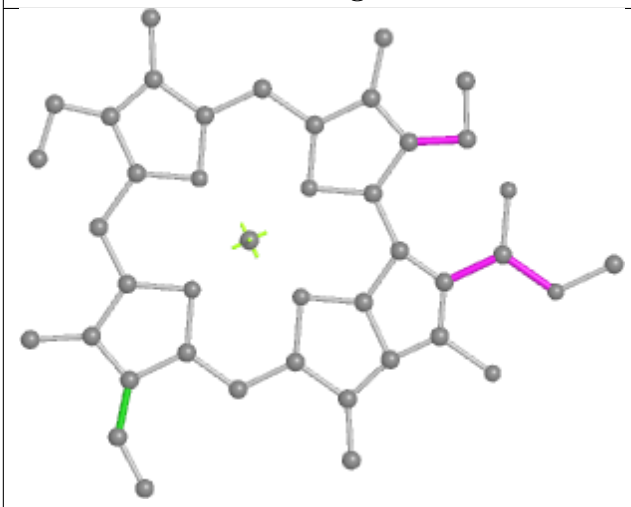
Ligand CLA 7 316



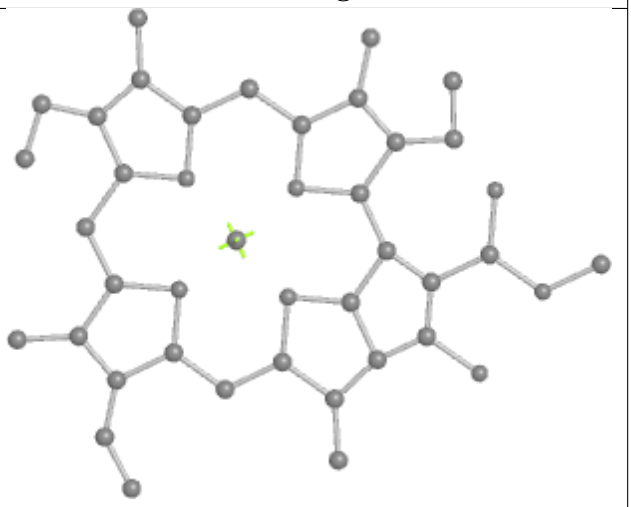
Bond lengths



Bond angles

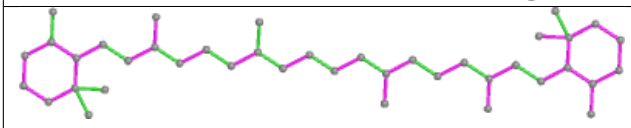


Torsions

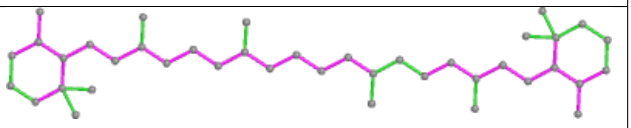


Rings

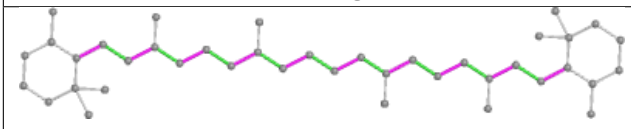
Ligand BCR B 845



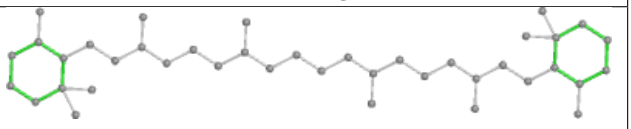
Bond lengths



Bond angles

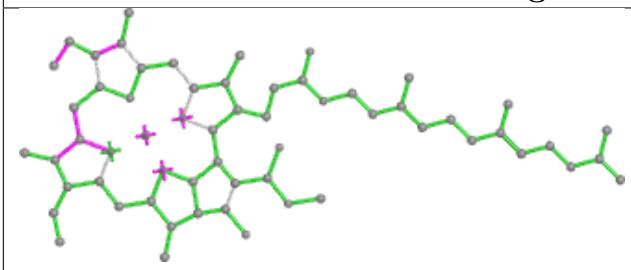


Torsions

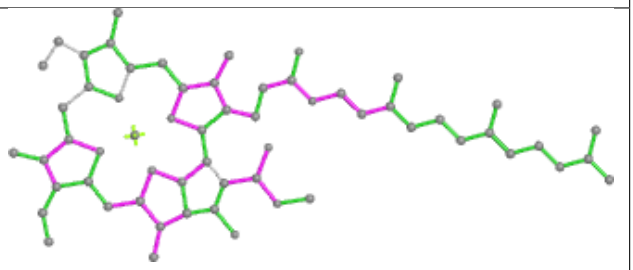


Rings

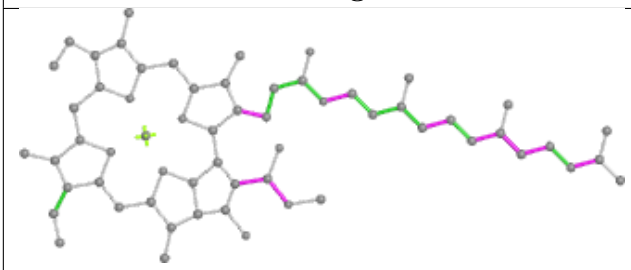
Ligand CLA 7 304



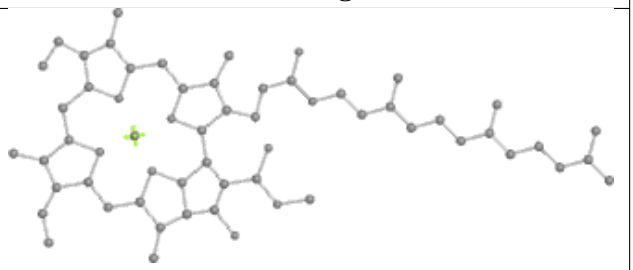
Bond lengths



Bond angles

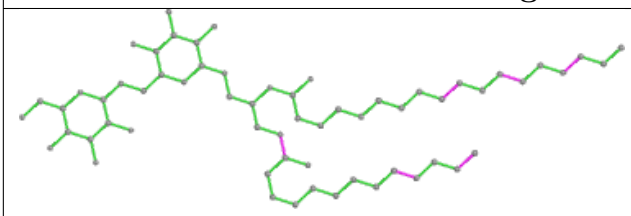


Torsions

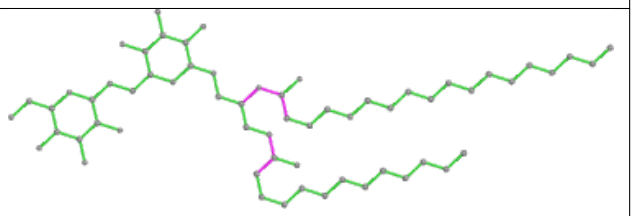


Rings

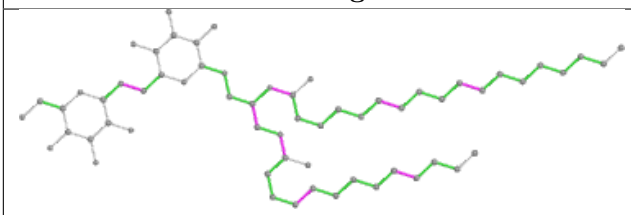
Ligand DGD B 852



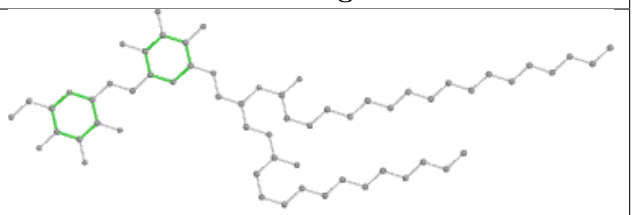
Bond lengths



Bond angles

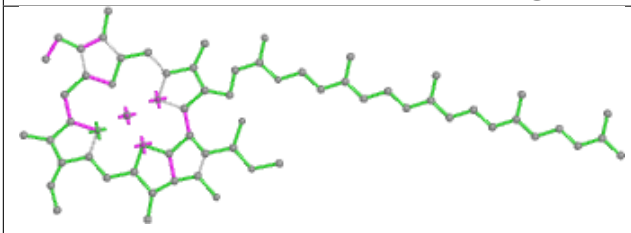


Torsions

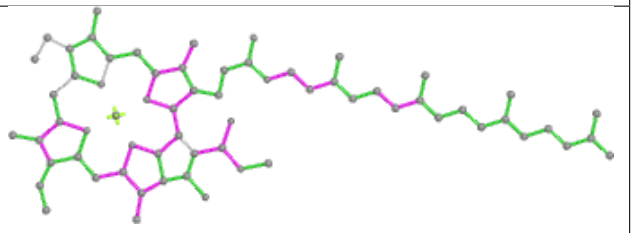


Rings

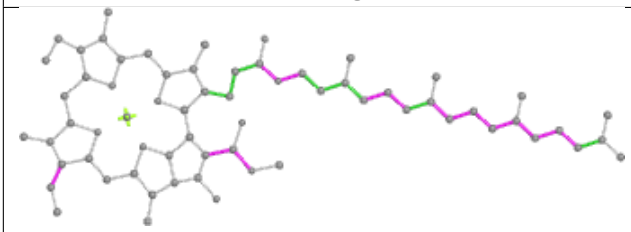
Ligand CLA 1 316



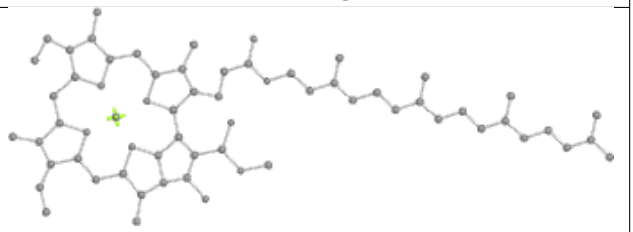
Bond lengths



Bond angles

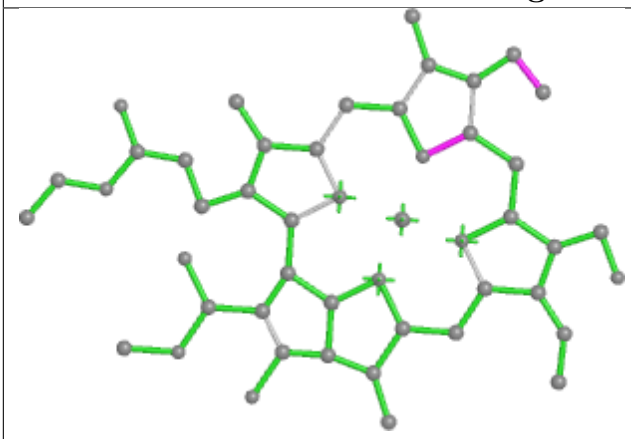


Torsions

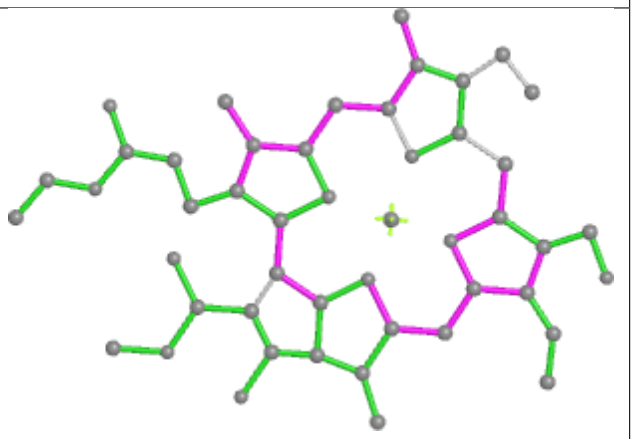


Rings

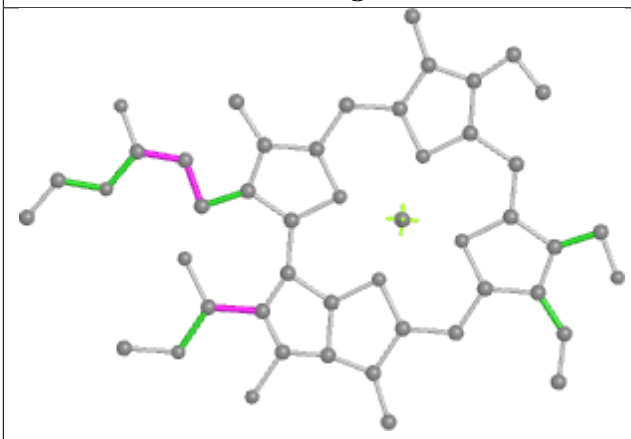
Ligand CHL 1 312



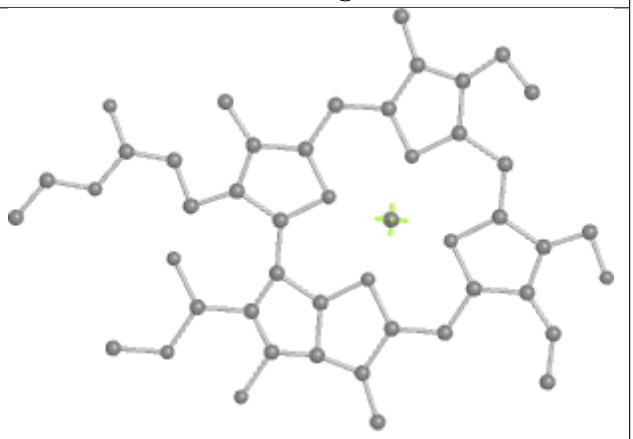
Bond lengths



Bond angles

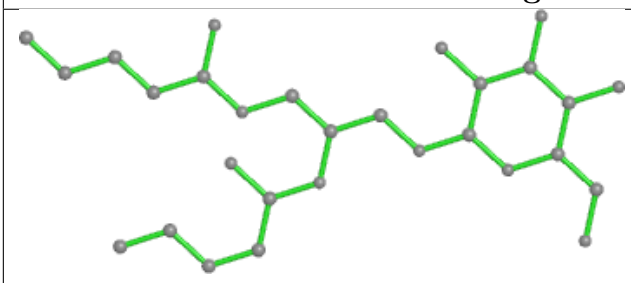


Torsions

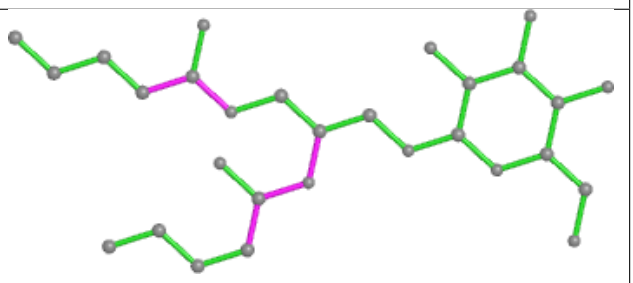


Rings

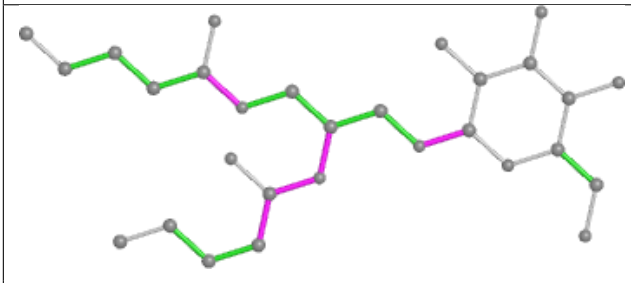
Ligand LMG J 101



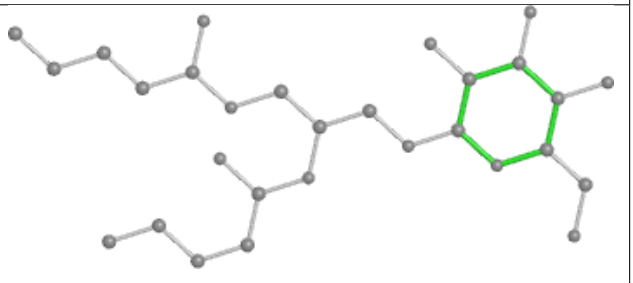
Bond lengths



Bond angles

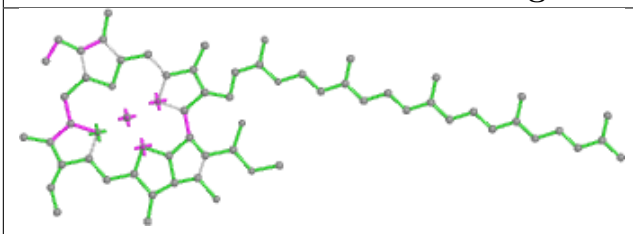


Torsions



Rings

Ligand CLA 8 307



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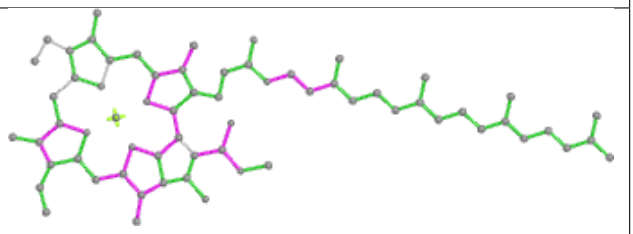
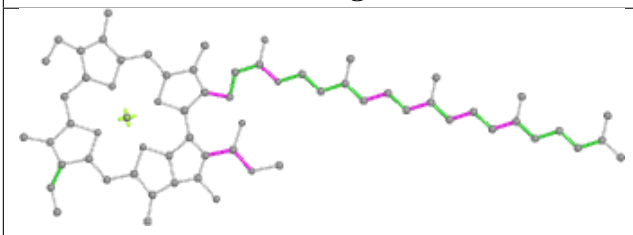


Bond angles



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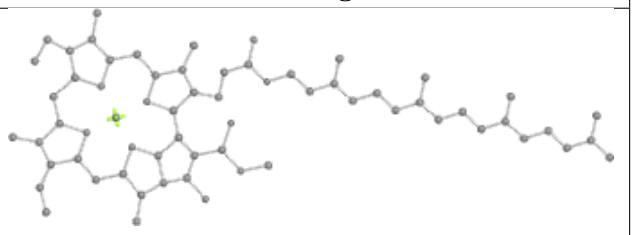
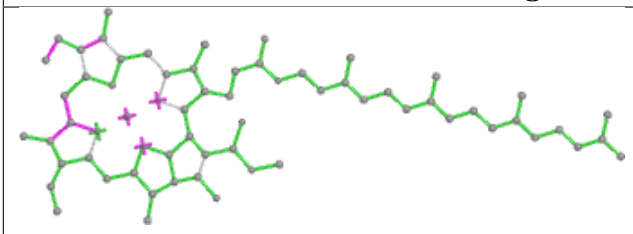


Rings

Ligand CLA A 802



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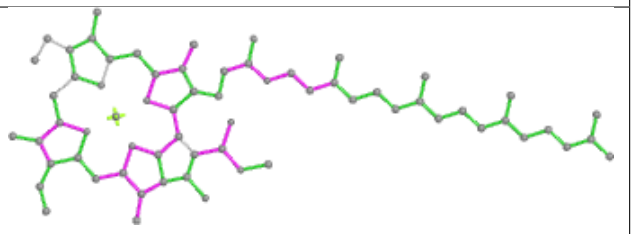
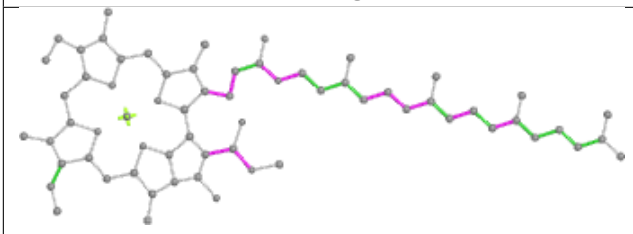


Bond angles



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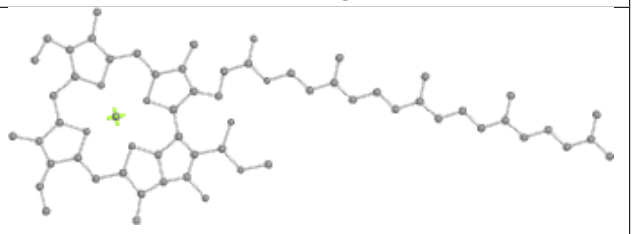
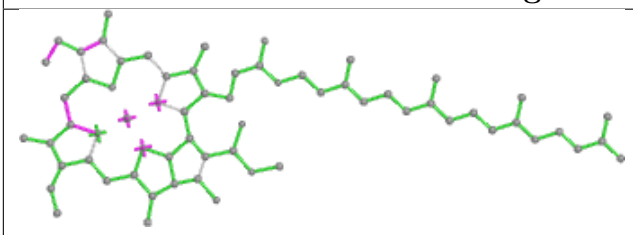
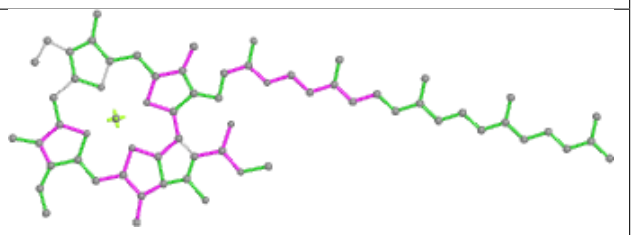


Rings

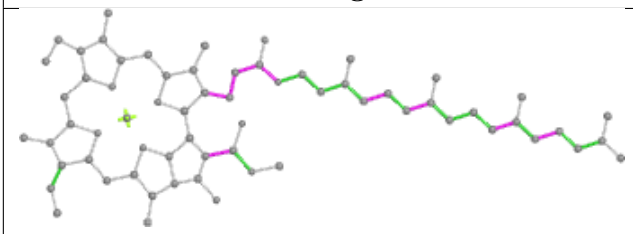
Ligand CLA A 813



Bond lengths



Bond angles



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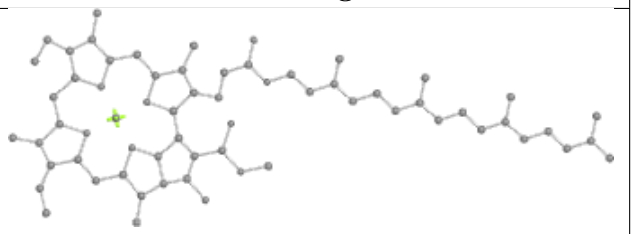
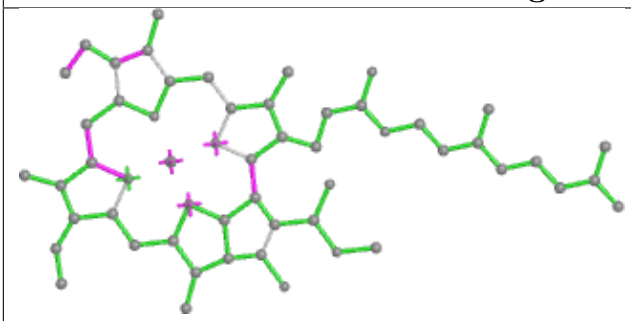
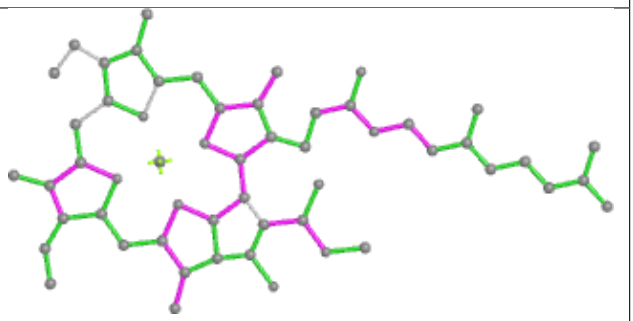


Rings

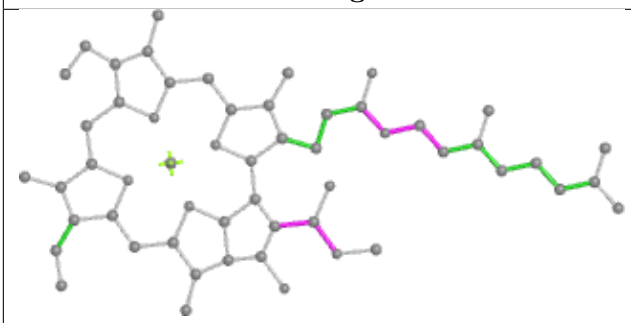
Ligand CLA 9 309



Bond lengths



Bond angles



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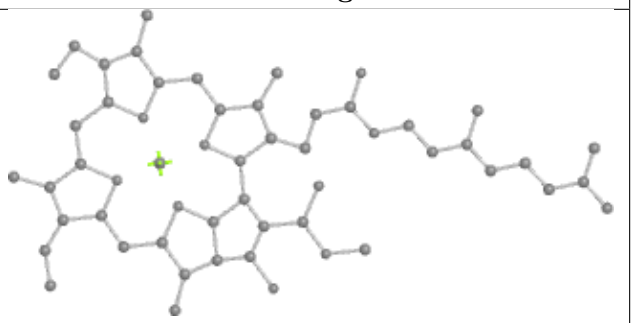
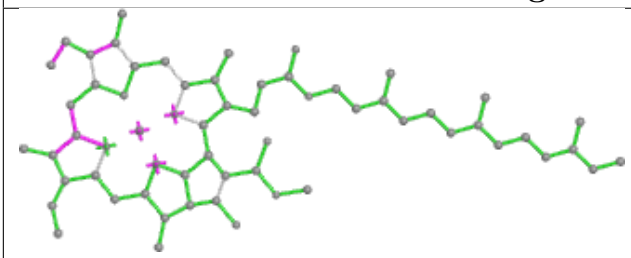
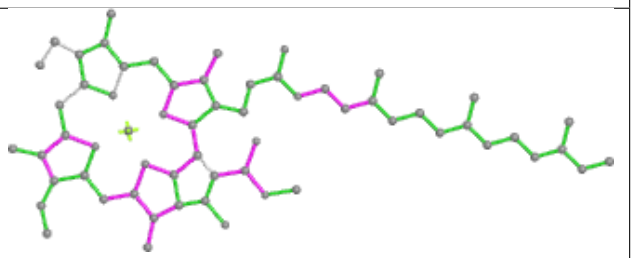


Rings

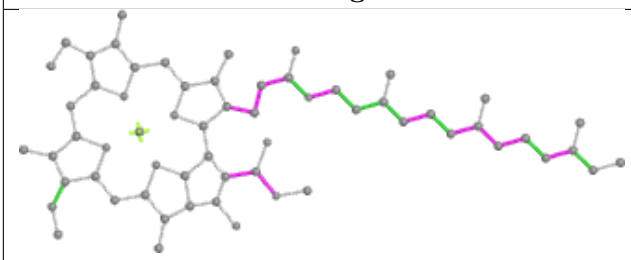
Ligand CLA 6 319



Bond lengths



Bond angles



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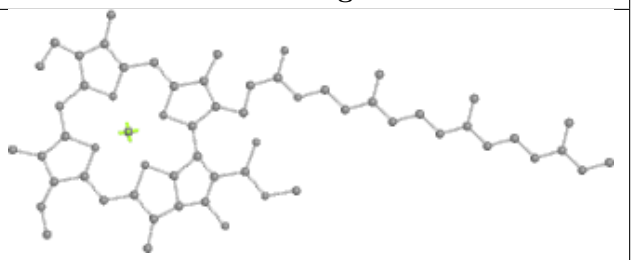
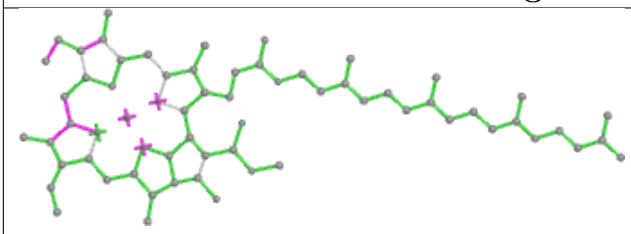


Rings

Ligand CLA B 827



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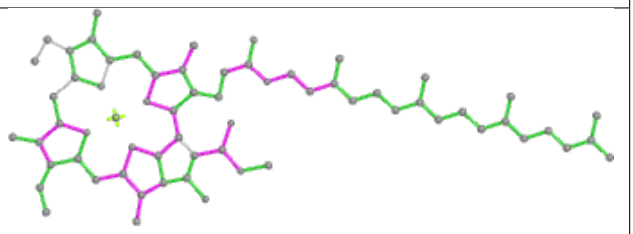
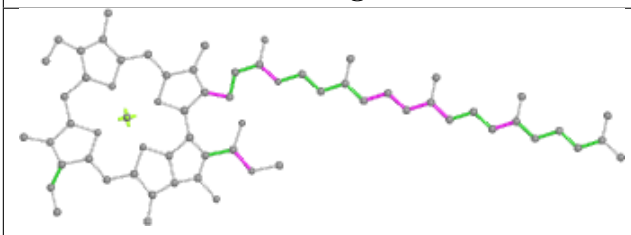
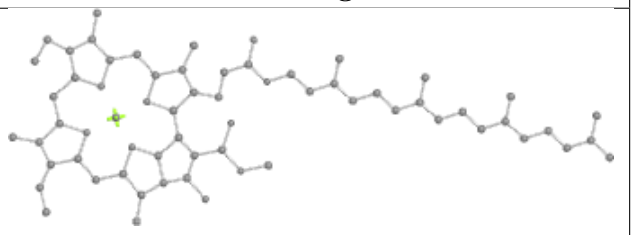


Bond angles

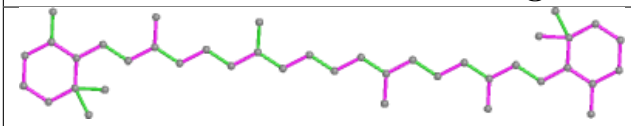


Torsions

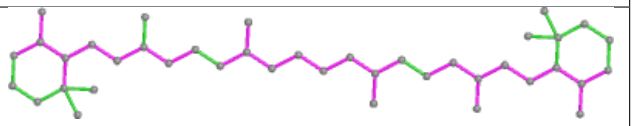


Rings

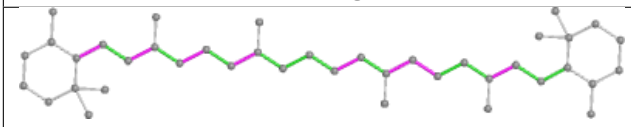
Ligand BCR 5 305



Bond lengths



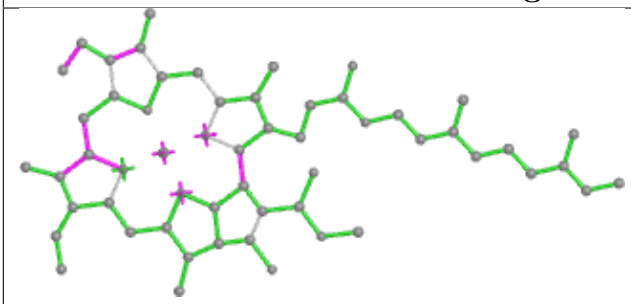
Bond angles



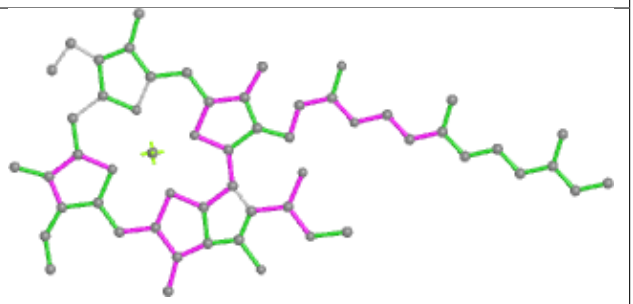
Torsions

Rings

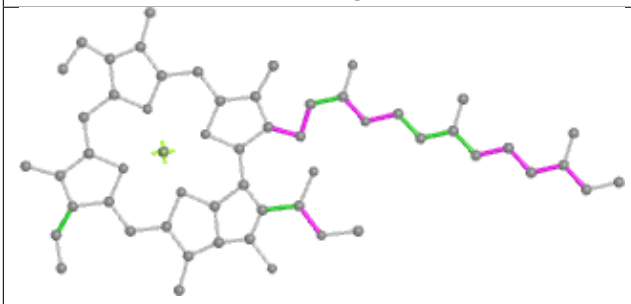
Ligand CLA A 816



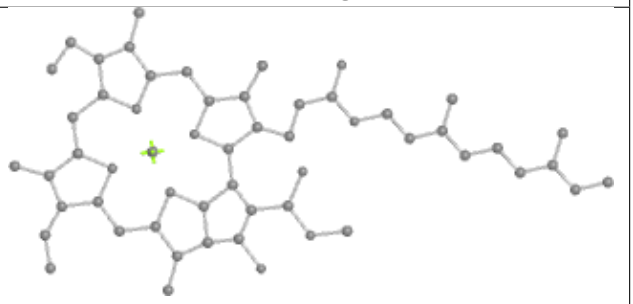
Bond lengths



Bond angles

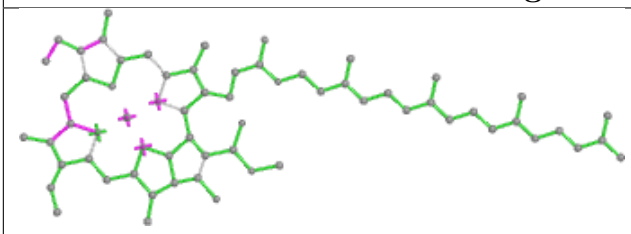


Torsions



Rings

Ligand CLA B 840



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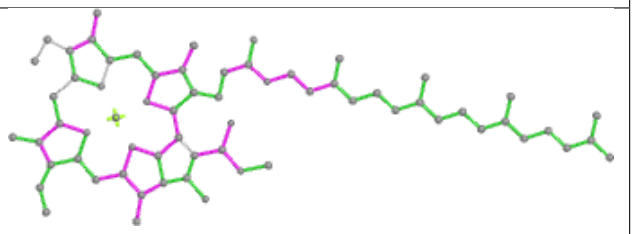
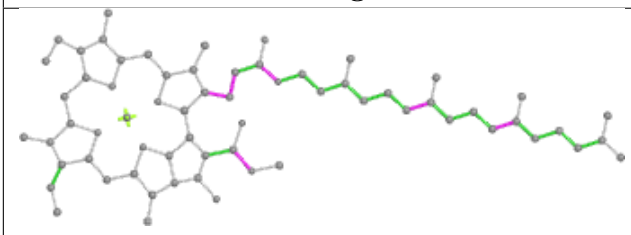
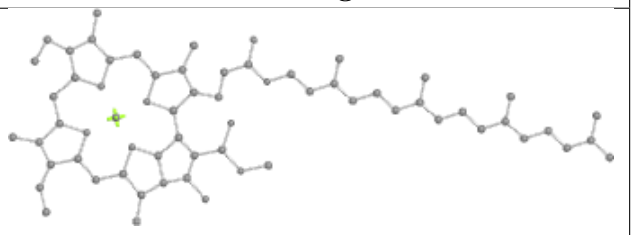


Bond angles

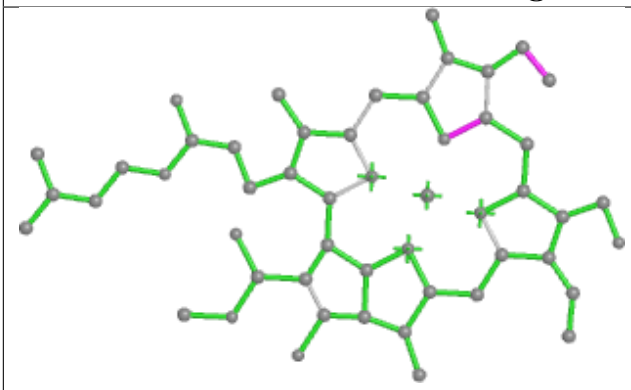


Torsions

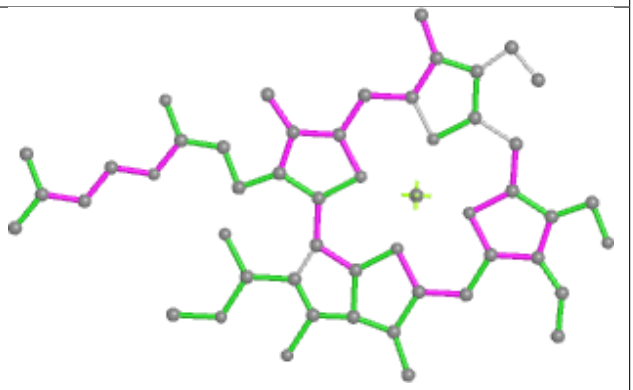


Rings

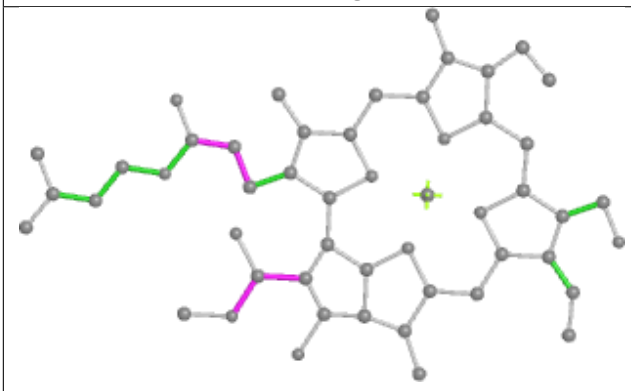
Ligand CHL 4 813



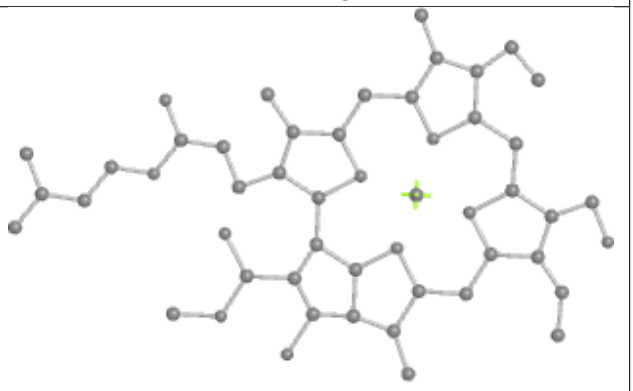
Bond lengths



Bond angles

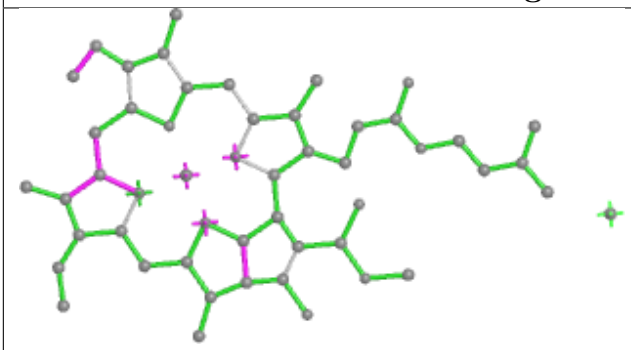


Torsions

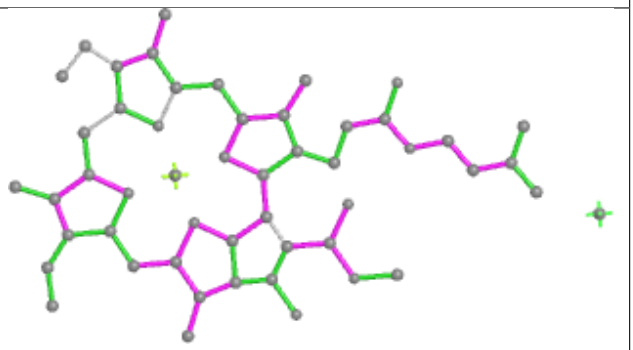


Rings

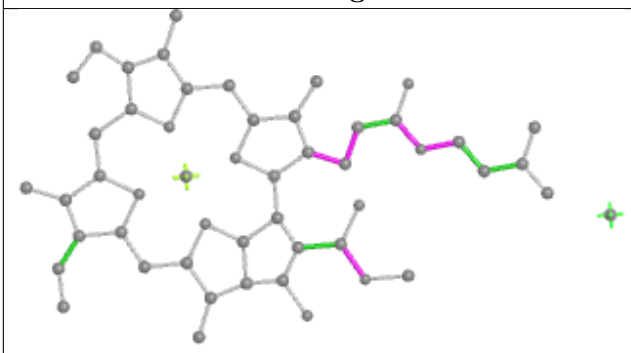
Ligand CLA B 835



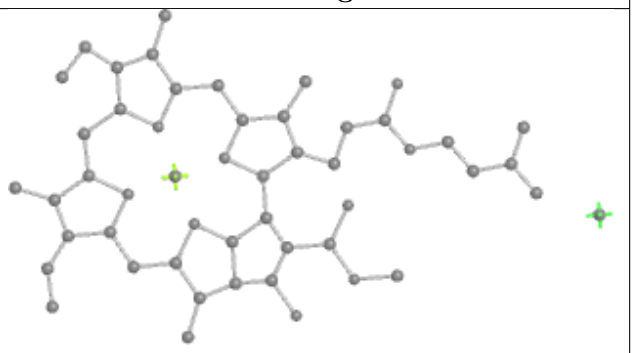
Bond lengths



Bond angles

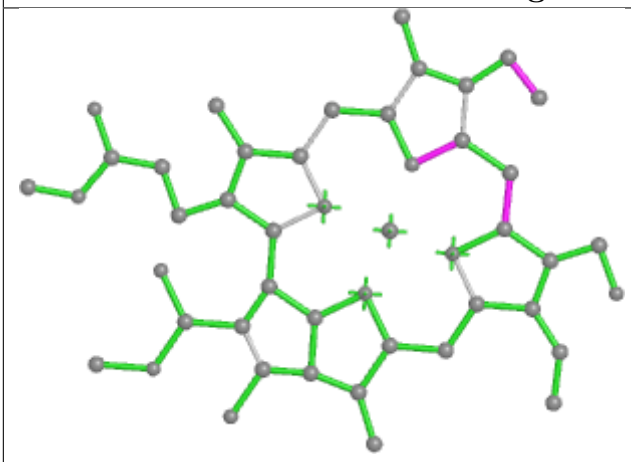


Torsions

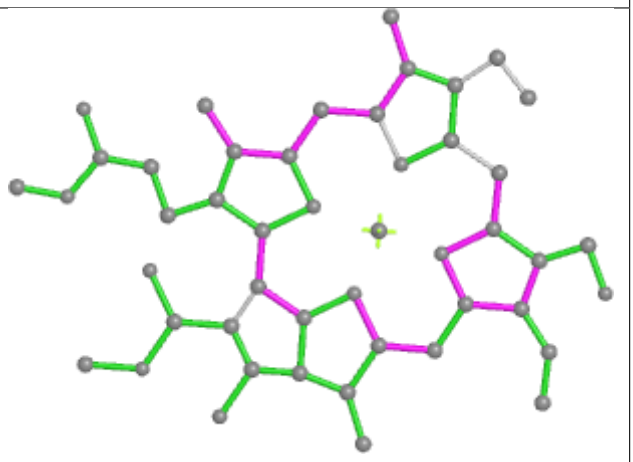


Rings

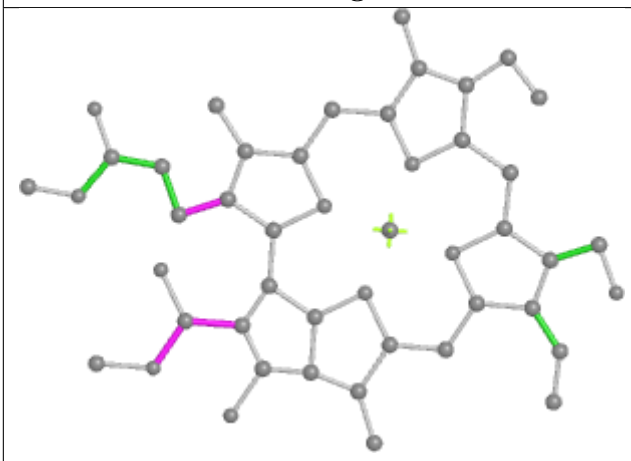
Ligand CHL 4 816



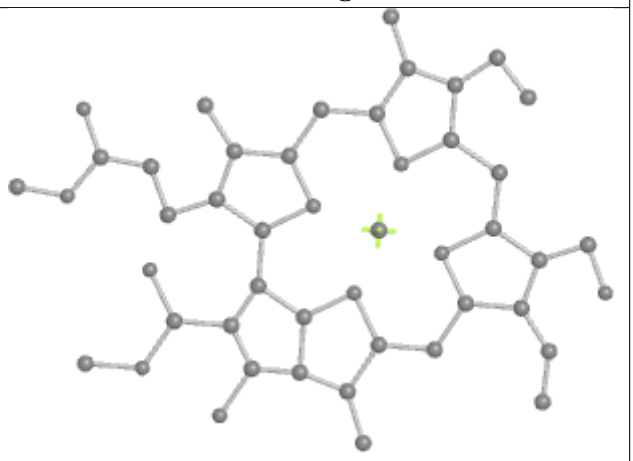
Bond lengths



Bond angles

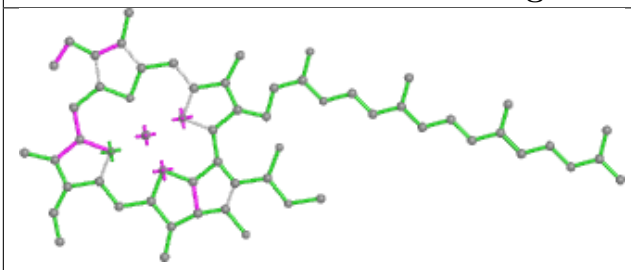


Torsions

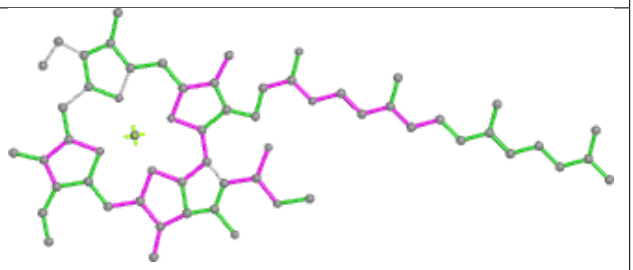


Rings

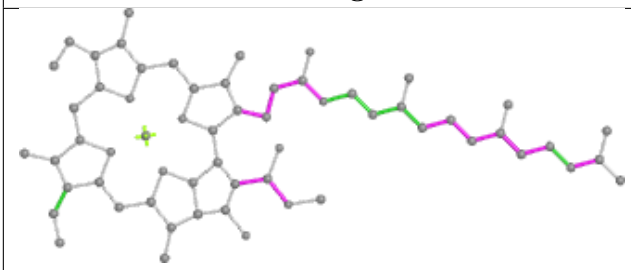
Ligand CLA Z 306



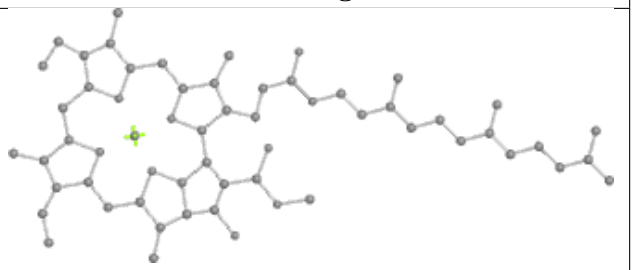
Bond lengths



Bond angles

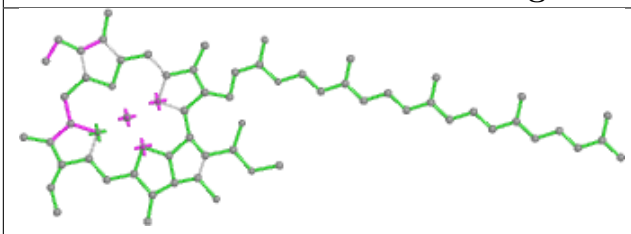


Torsions



Rings

Ligand CLA 1 306



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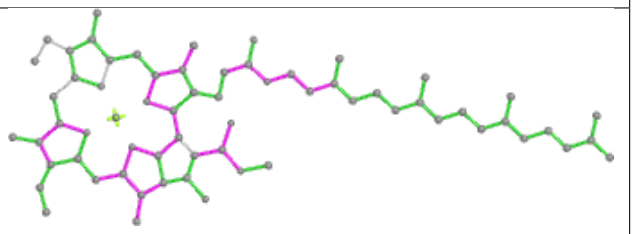
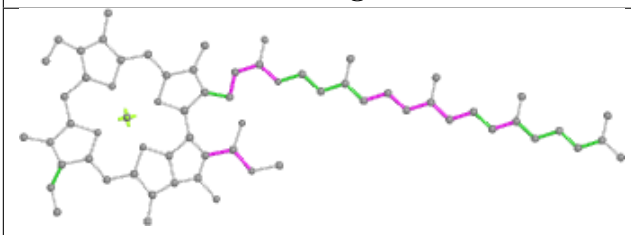


Bond angles



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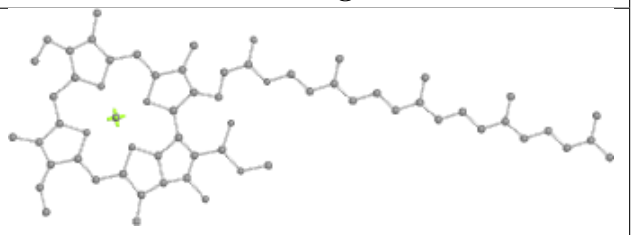
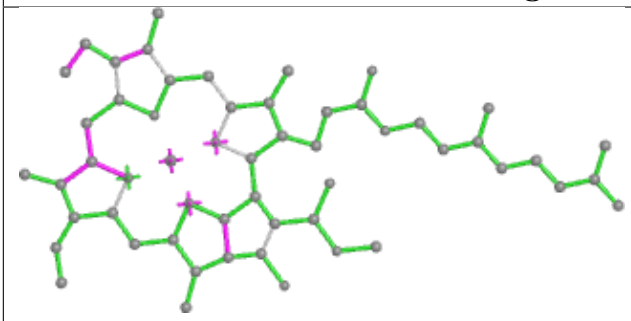
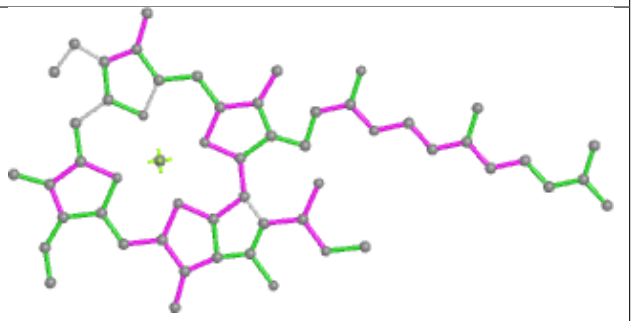


Rings

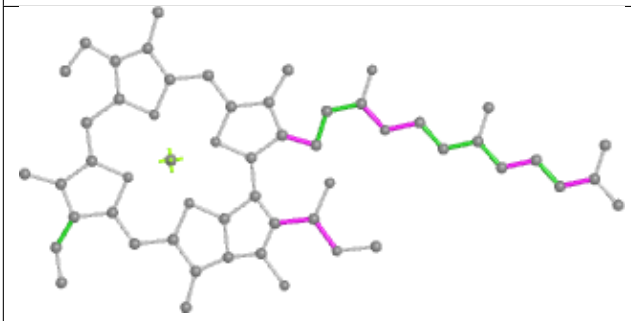
Ligand CLA A 836



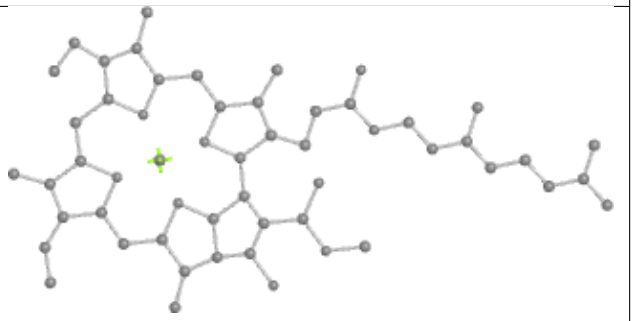
Bond lengths



Bond angles

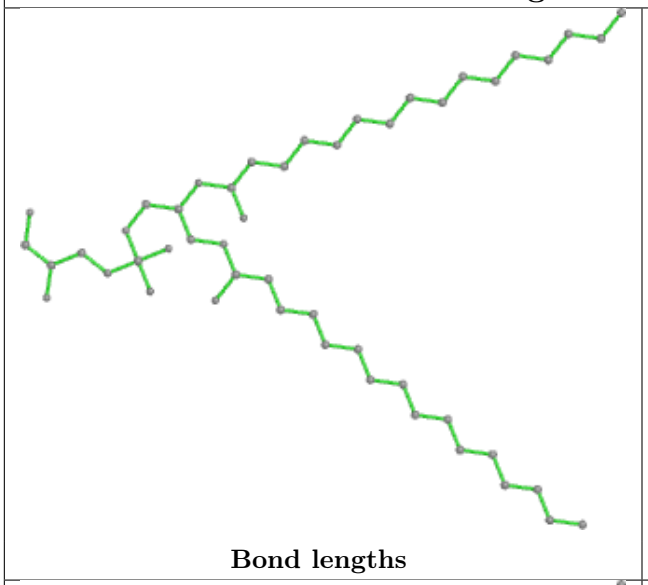


Torsions

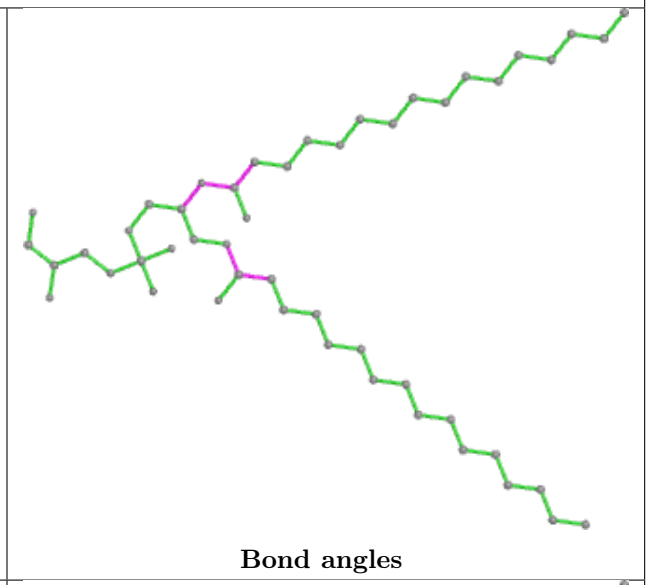


Rings

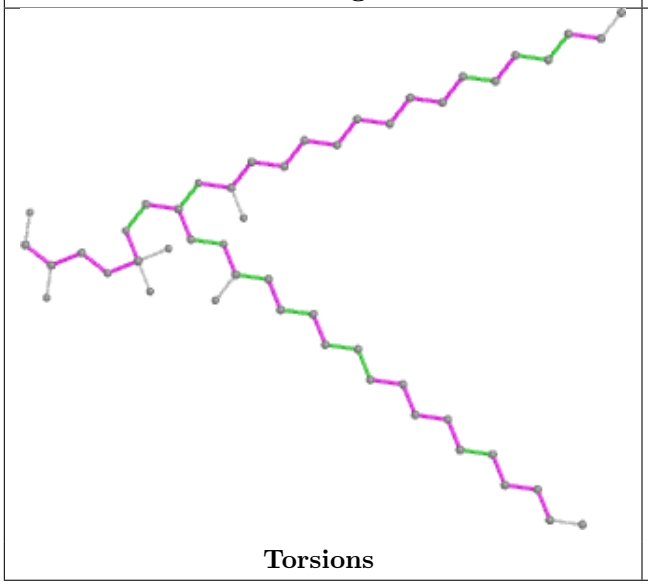
Ligand LHG 6 323



Bond lengths



Bond angles



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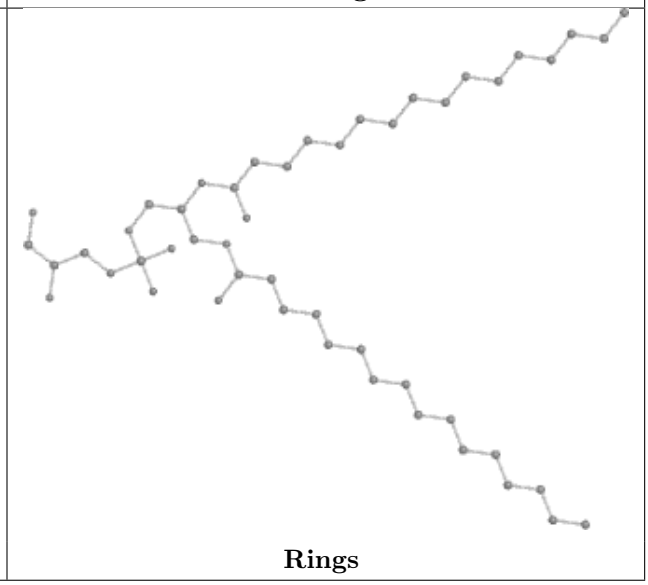
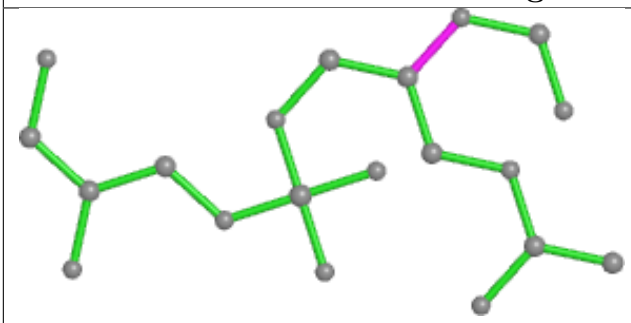
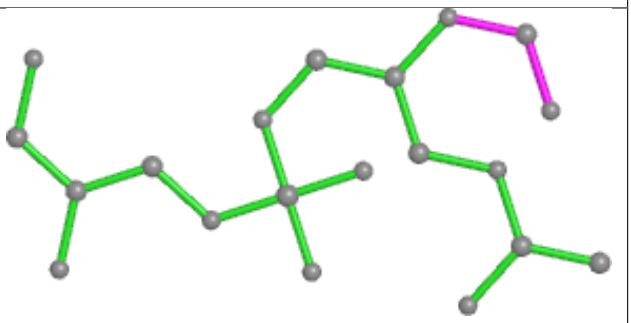


Rings

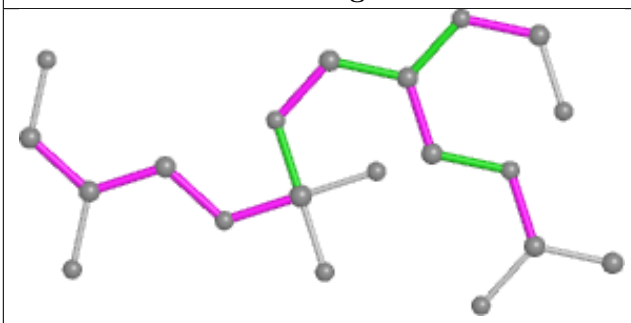
Ligand LHG 3 321



Bond lengths



Bond angles



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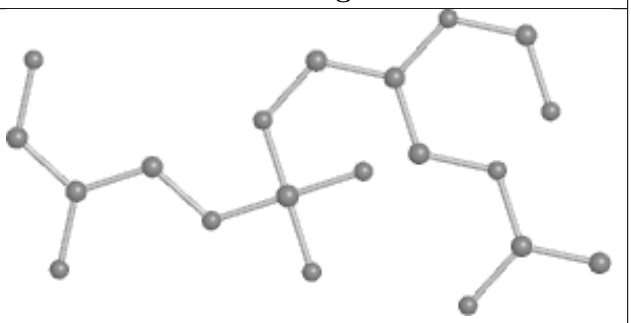
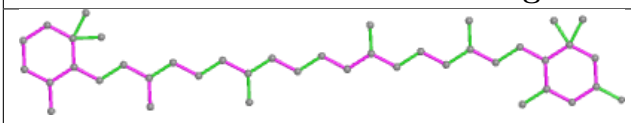
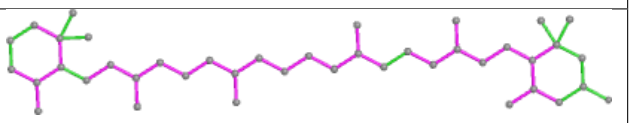


Rings

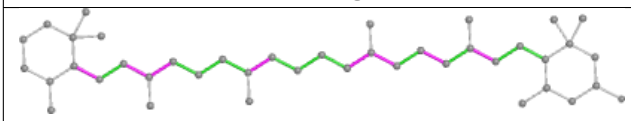
Ligand RRX F 306



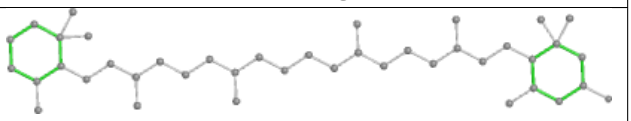
Bond lengths



Bond angles

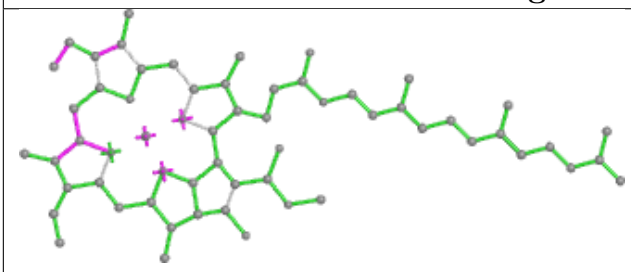


Torsions

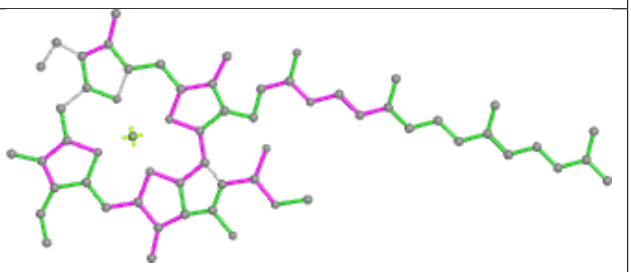


Rings

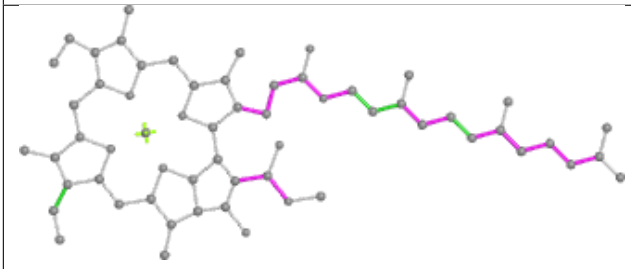
Ligand CLA 7 324



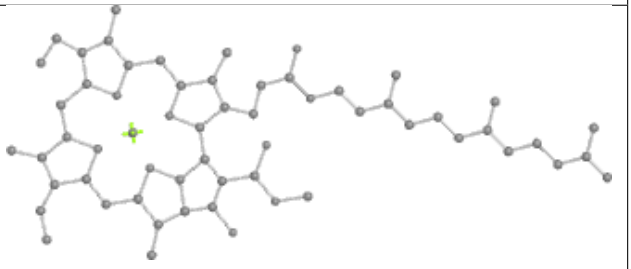
Bond lengths



Bond angles

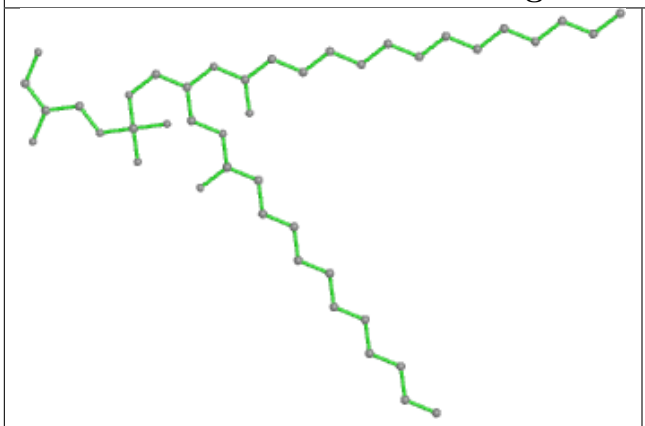


Torsions

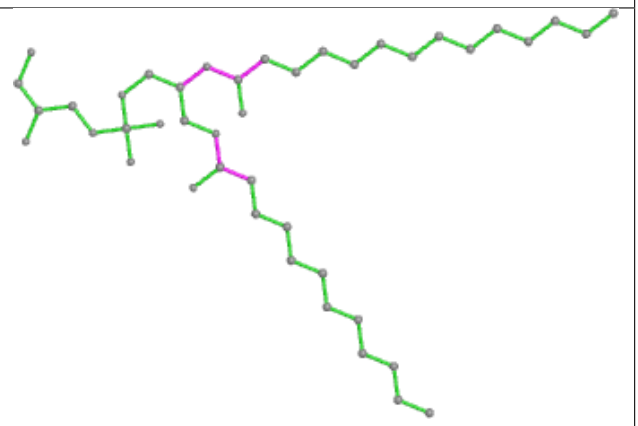


Rings

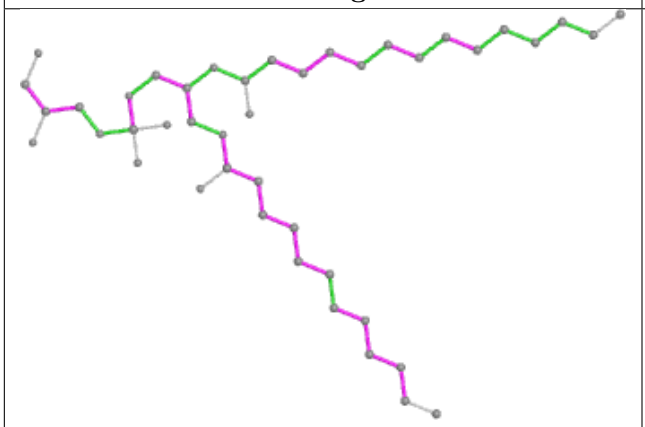
Ligand LHG Z 317



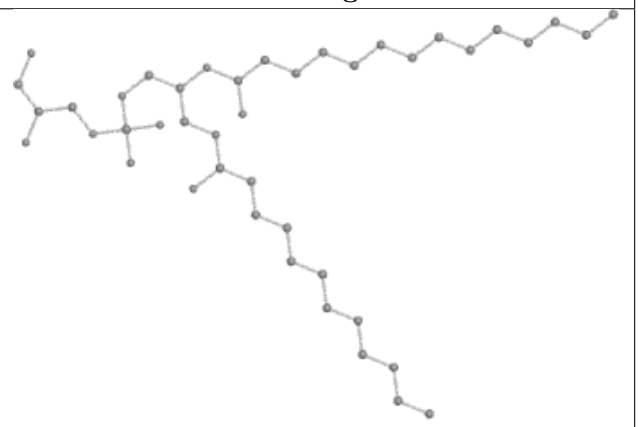
Bond lengths



Bond angles

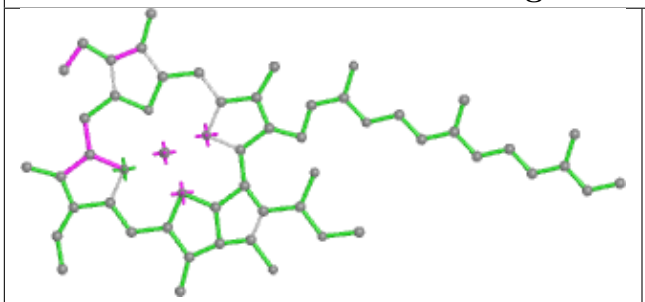


Torsions

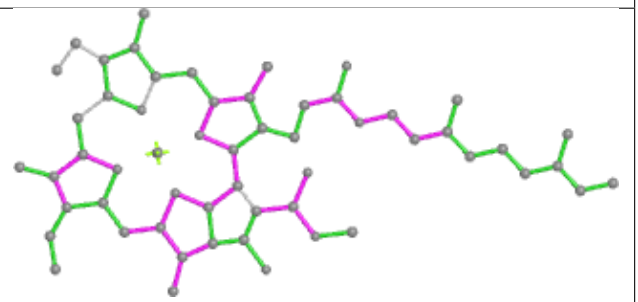


Rings

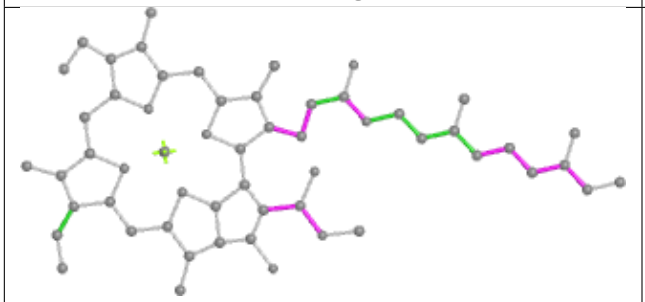
Ligand CLA B 813



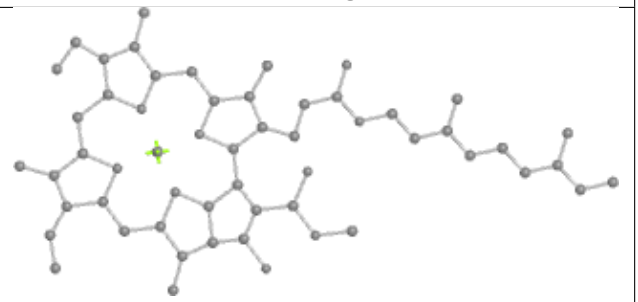
Bond lengths



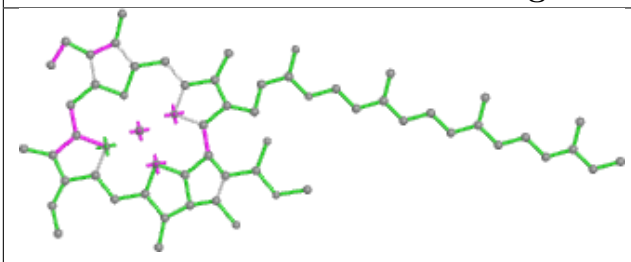
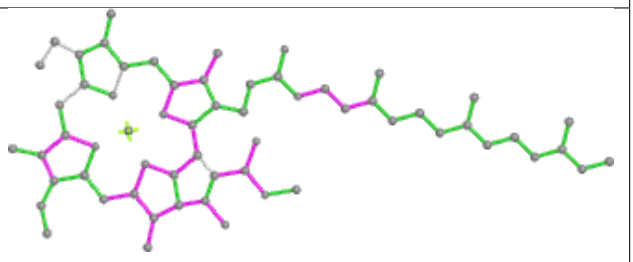
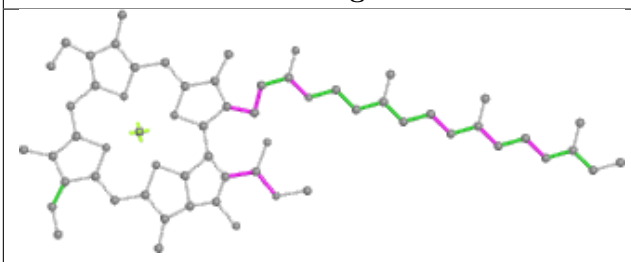
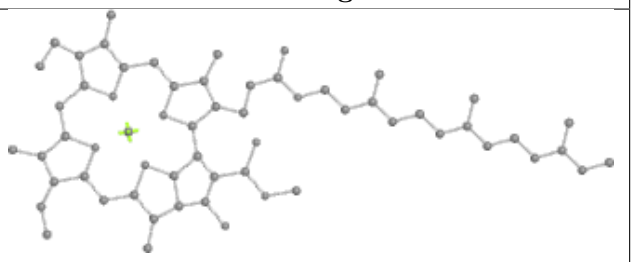
Bond angles

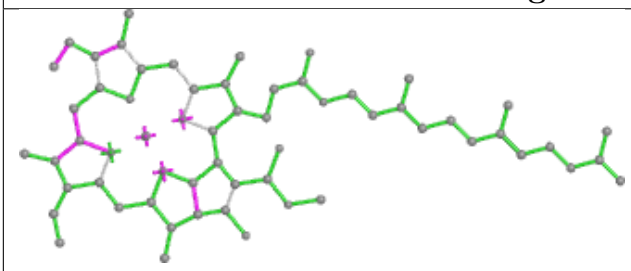
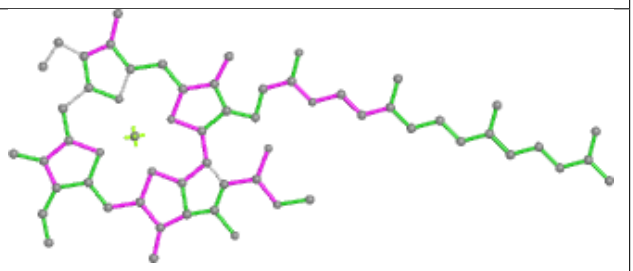
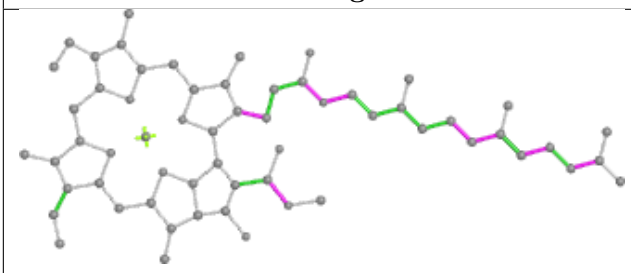
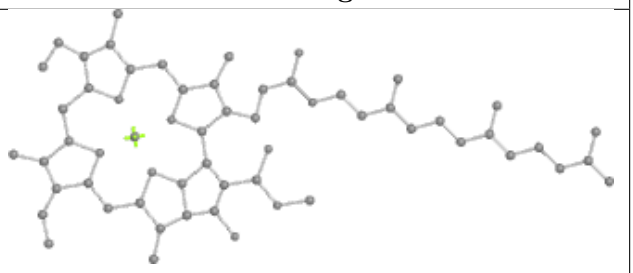


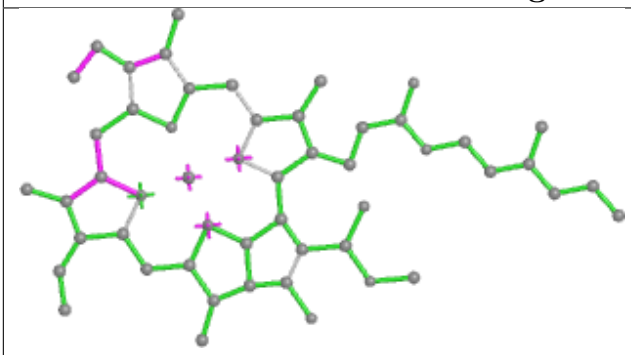
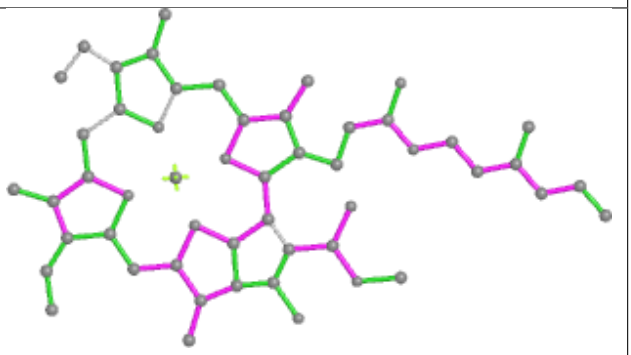
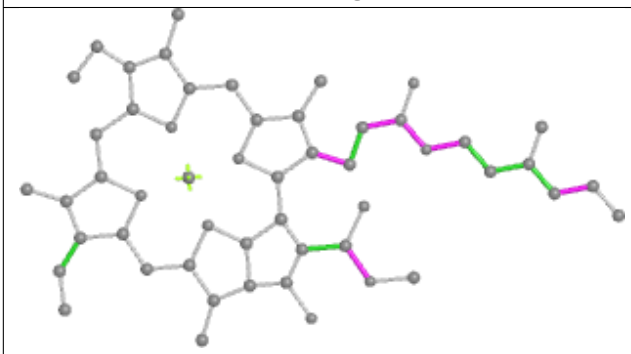
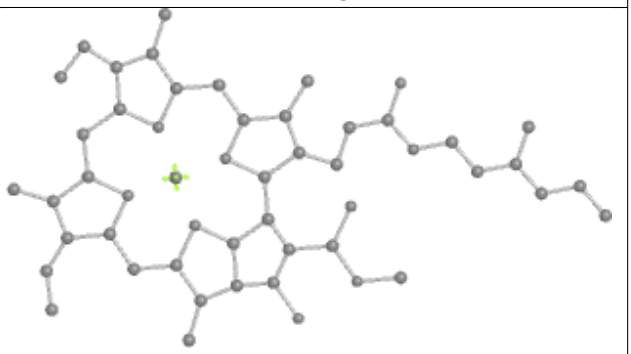
Torsions

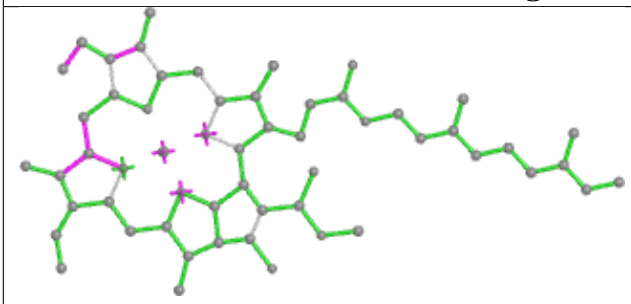
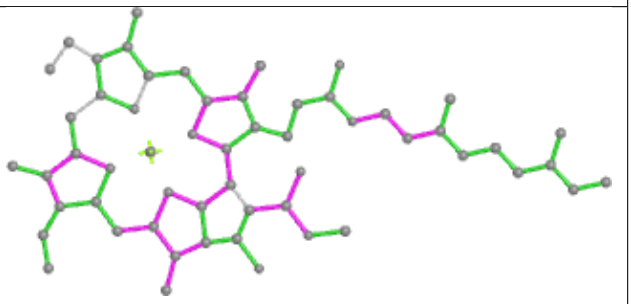
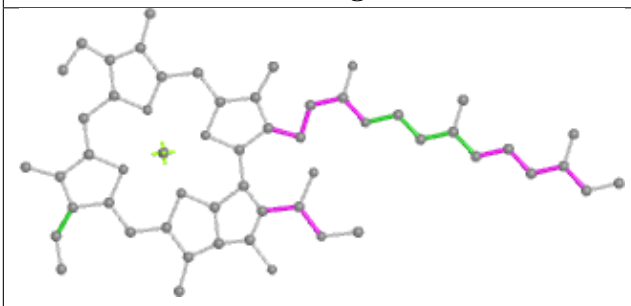
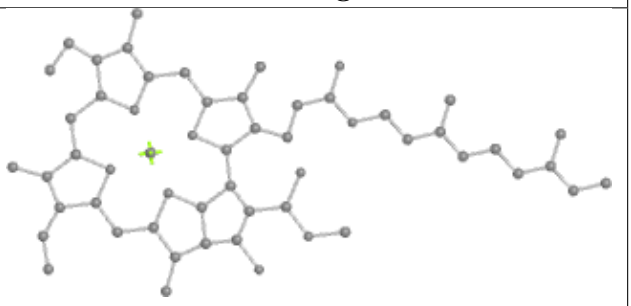


Rings

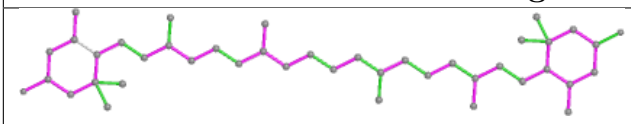
Ligand CLA 5 313	
 Bond lengths	 Bond angles
 Torsions	 Rings

Ligand CLA 6 307	
 Bond lengths	 Bond angles
 Torsions	 Rings

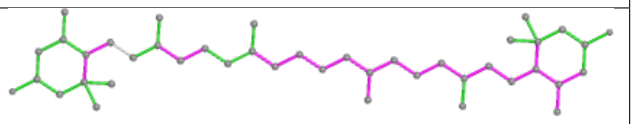
Ligand CLA 3 316	
 Bond lengths	 Bond angles
 Torsions	 Rings

Ligand CLA B 822	
 Bond lengths	 Bond angles
 Torsions	 Rings

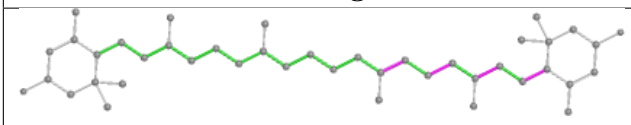
Ligand LUT 6 304



Bond lengths



Bond angles



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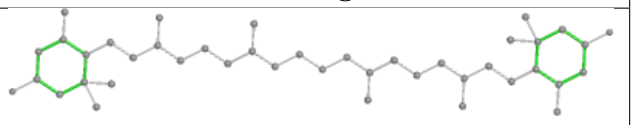
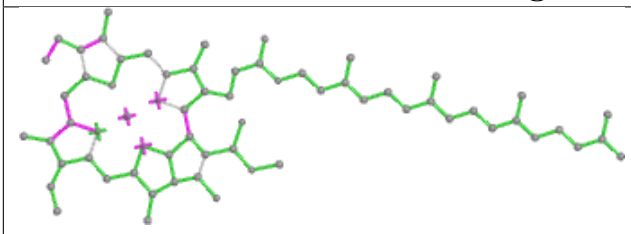


Rings

Ligand CLA 6 309



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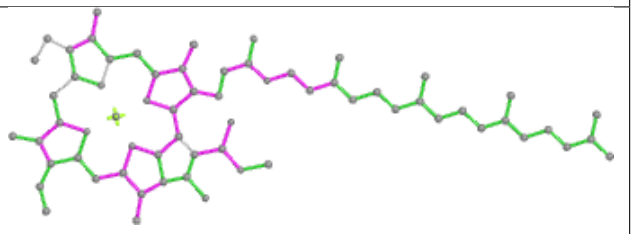
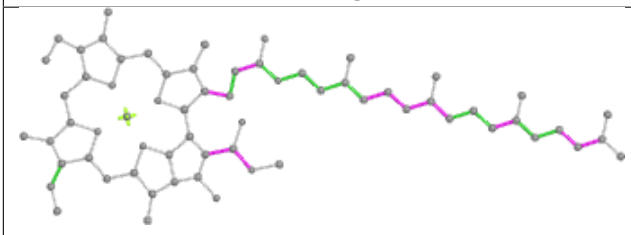


Bond angles



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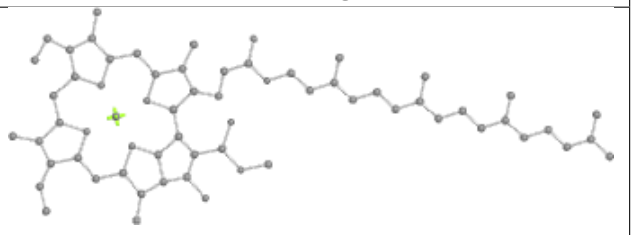
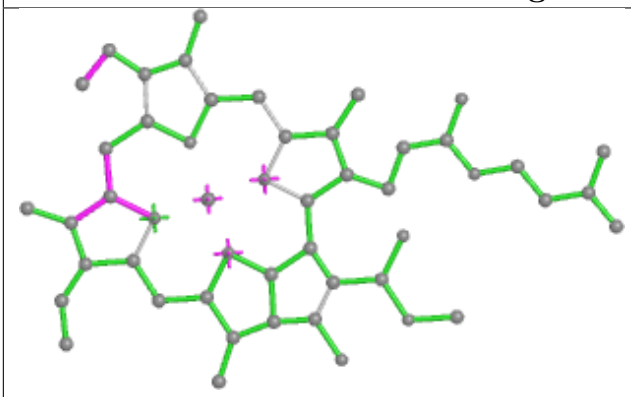


Rings

Ligand CLA 4 810



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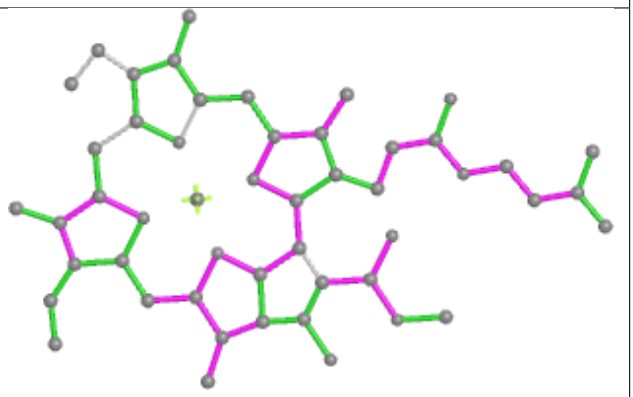
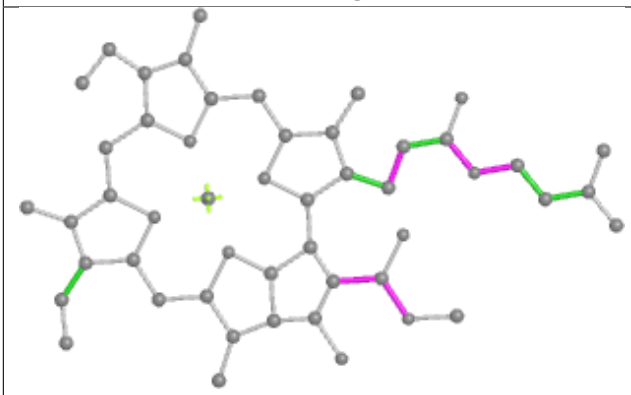


Bond angles



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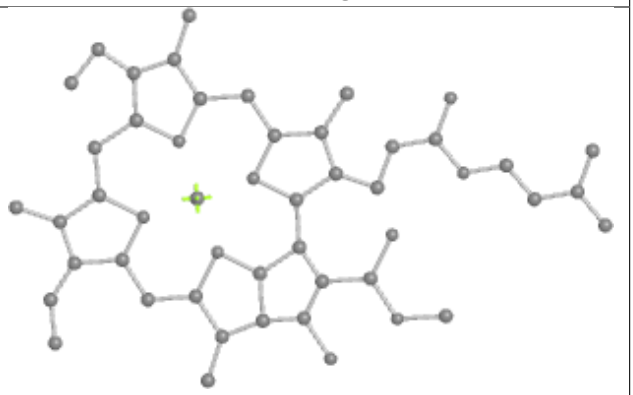
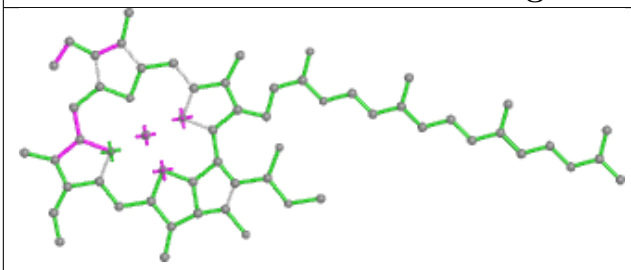
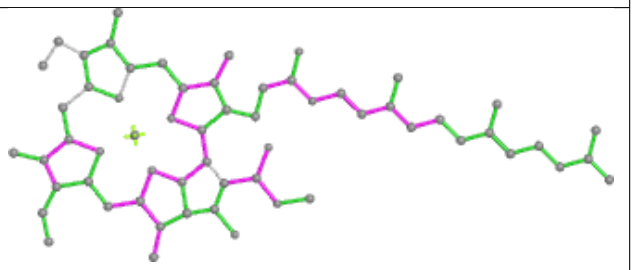


Rings

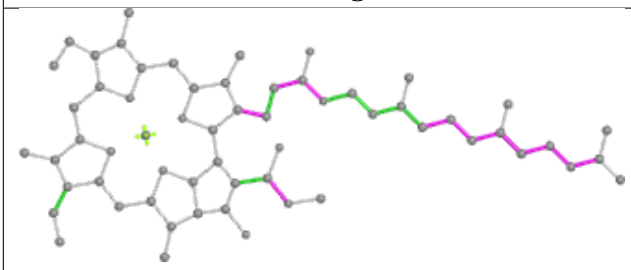
Ligand CLA Z 309



Bond lengths



Bond angles



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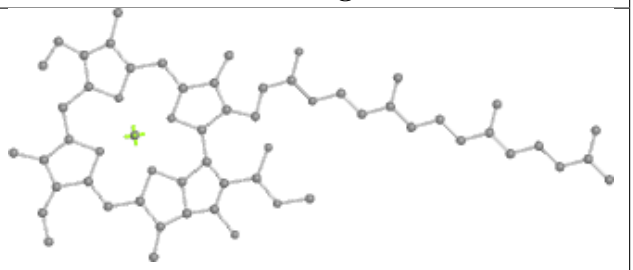
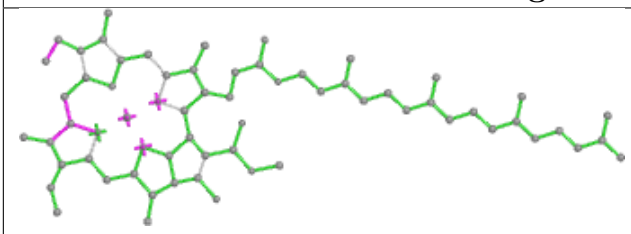
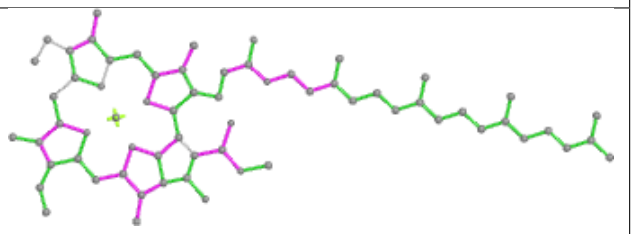


Rings

Ligand CLA 3 313



Bond lengths



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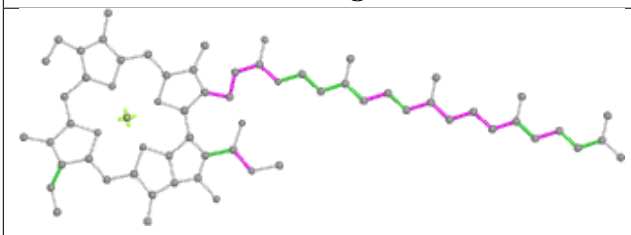

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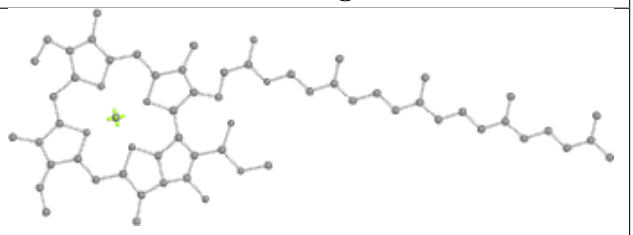
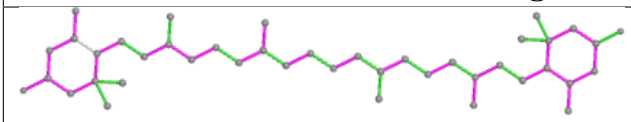


Rings

Ligand LUT 6 303



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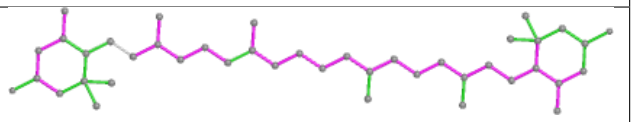
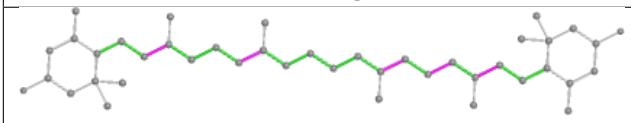
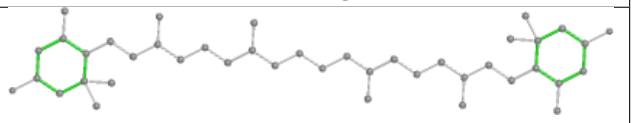


Bond angles

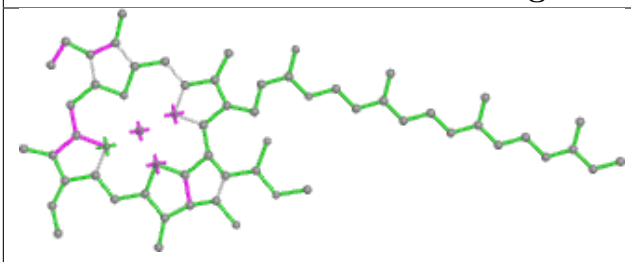


Torsions



Rings

Ligand CLA 7 308



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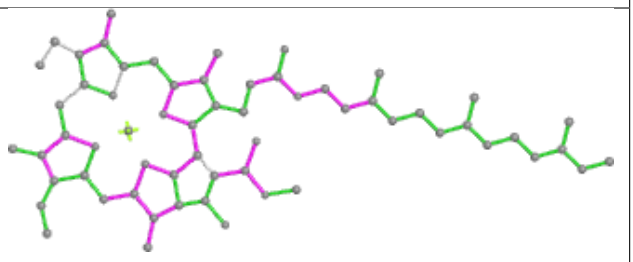
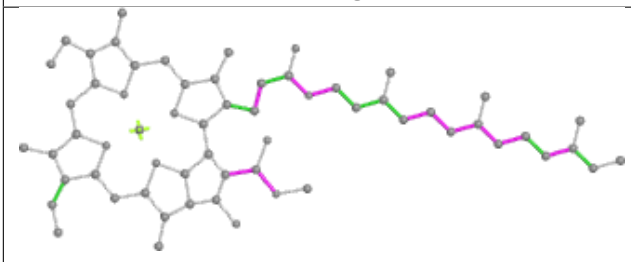
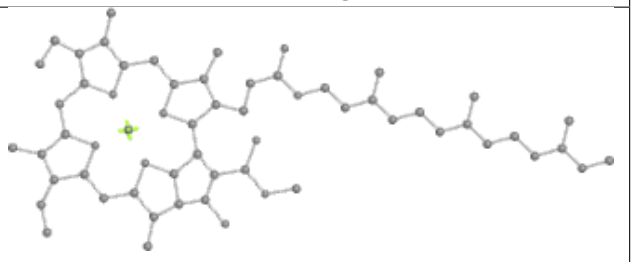


Bond angles

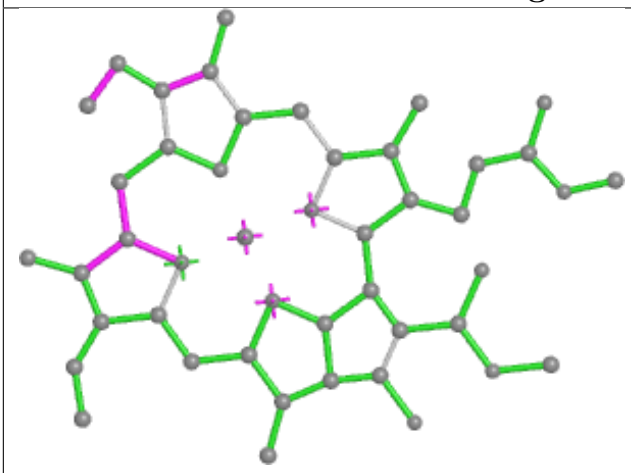


Torsions

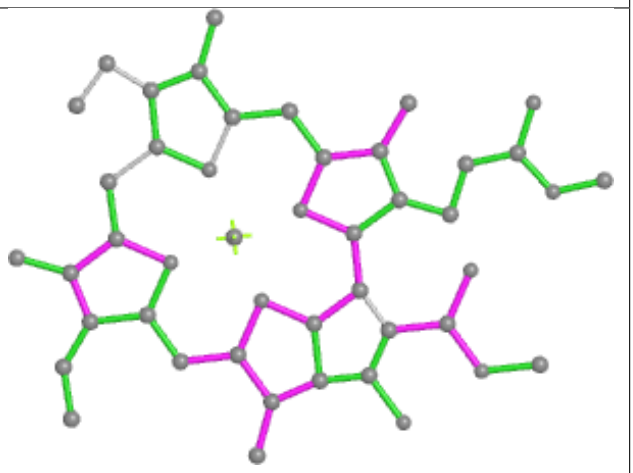


Rings

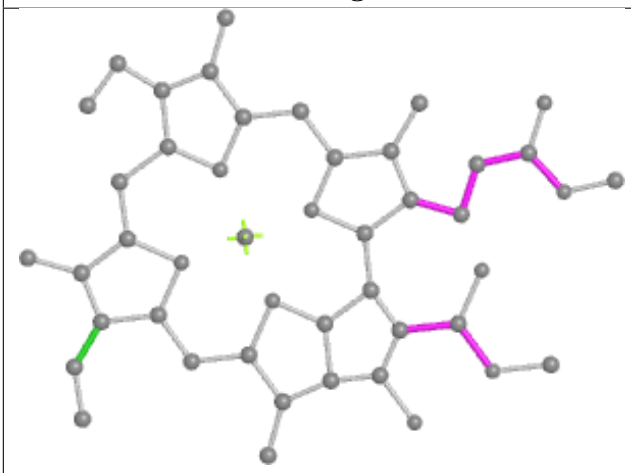
Ligand CLA 8 314



Bond lengths



Bond angles



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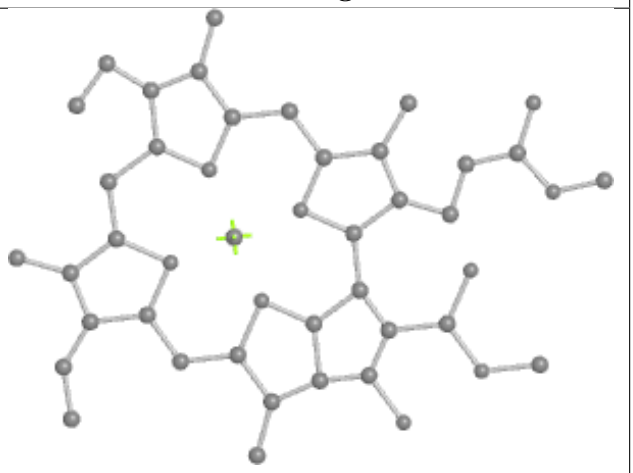
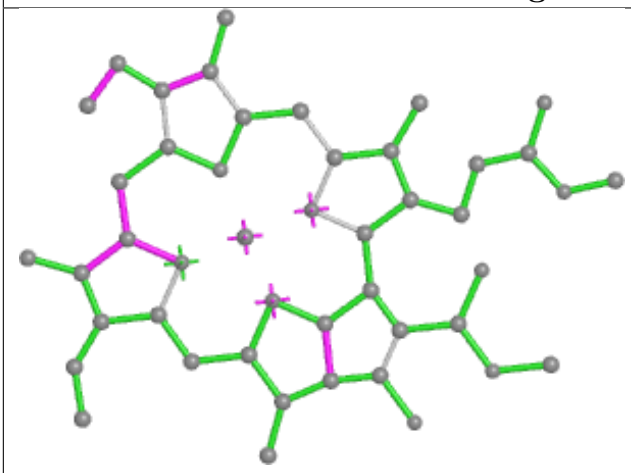


Rings

Ligand CLA 3 309



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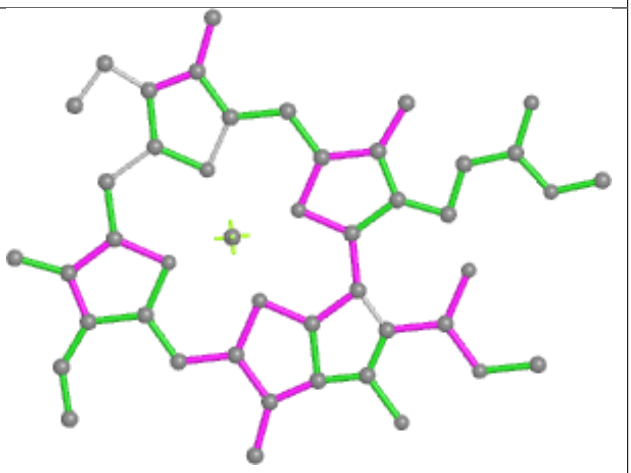
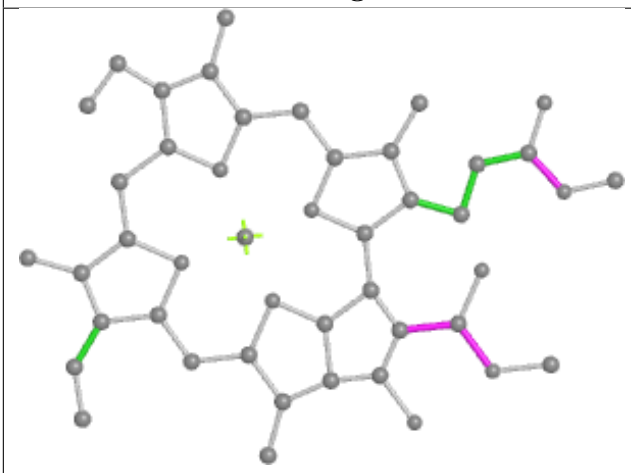


Bond angles



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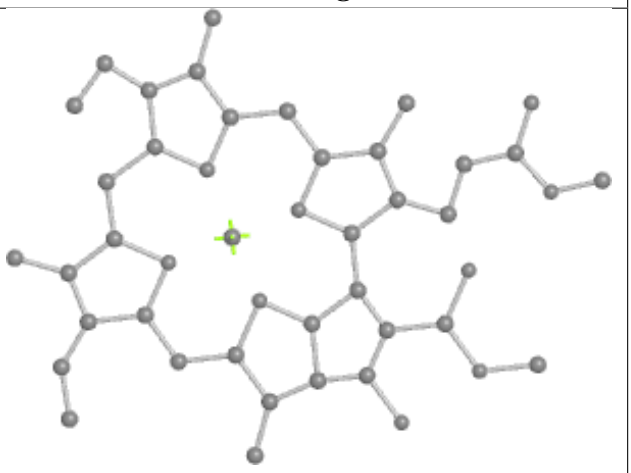
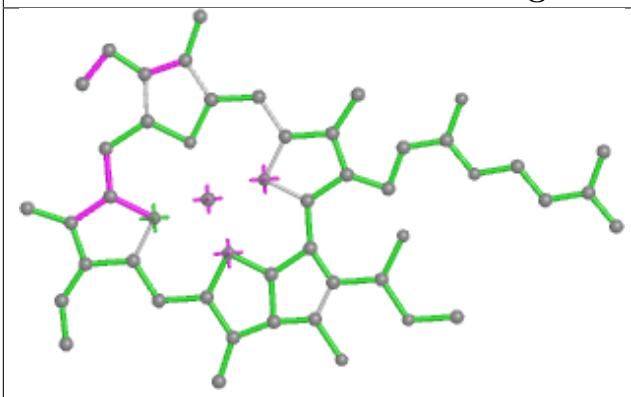
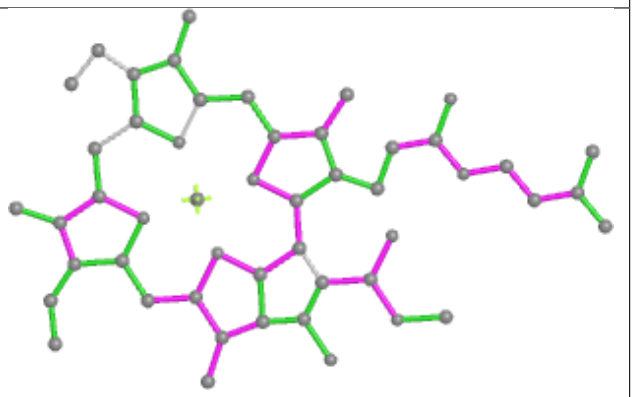


Rings

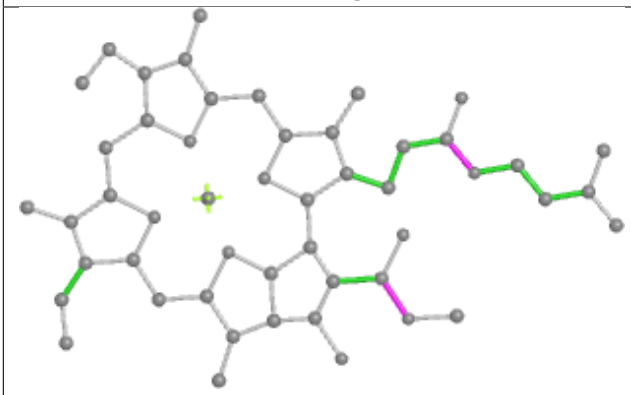
Ligand CLA 7 305



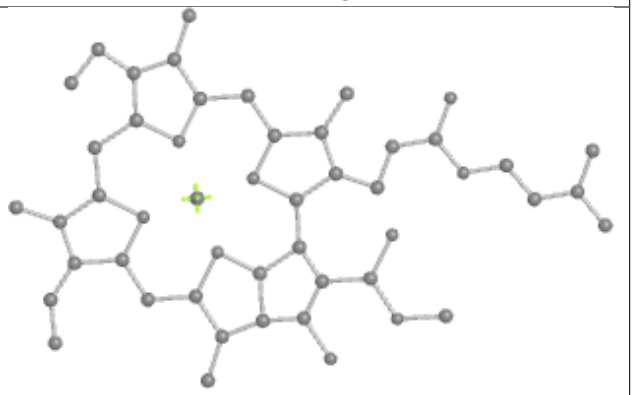
Bond lengths



Bond angles

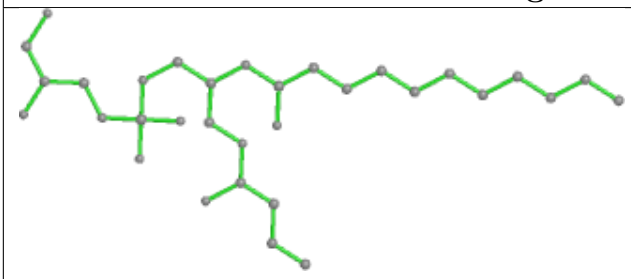


Torsions

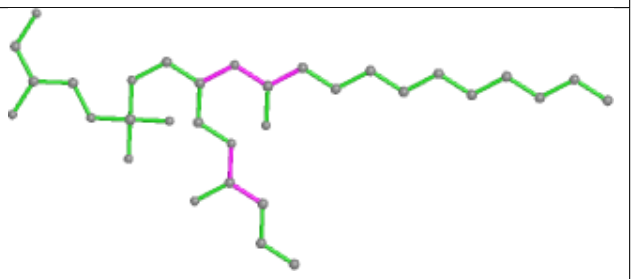


Rings

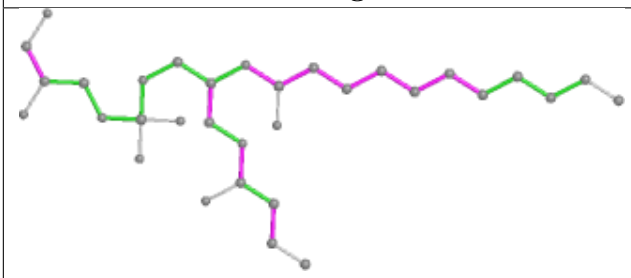
Ligand LHG 4 821



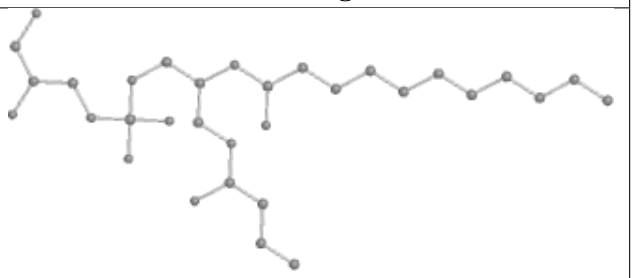
Bond lengths



Bond angles

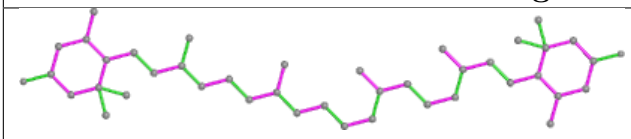


Torsions

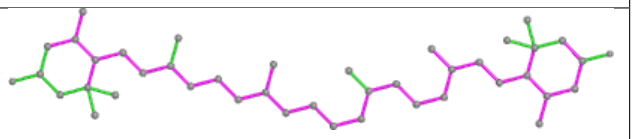


Rings

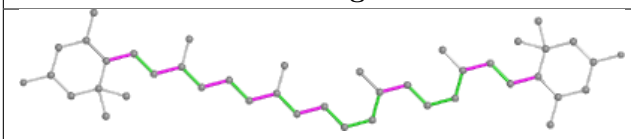
Ligand C7Z J 105



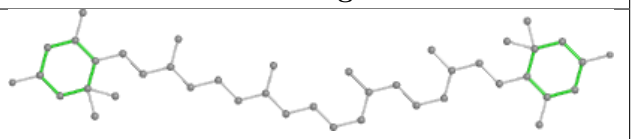
Bond lengths



Bond angles

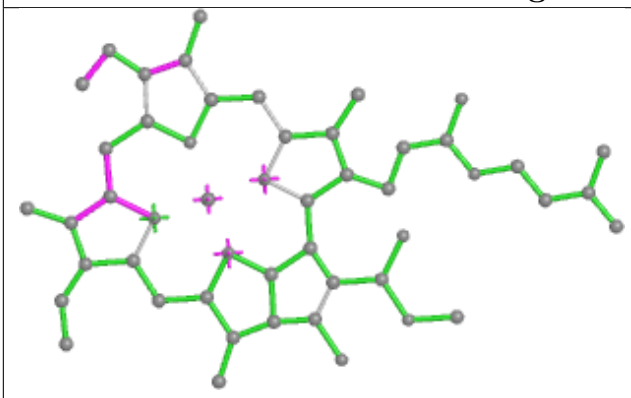


Torsions



Rings

Ligand CLA 9 313



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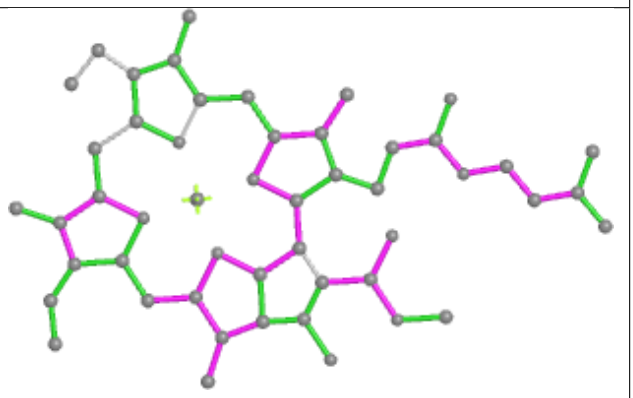
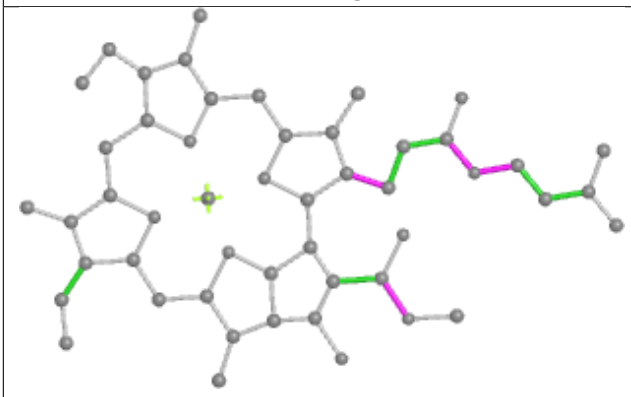


Bond angles



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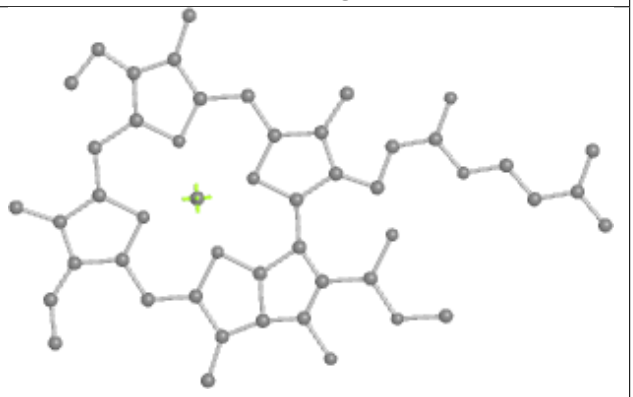
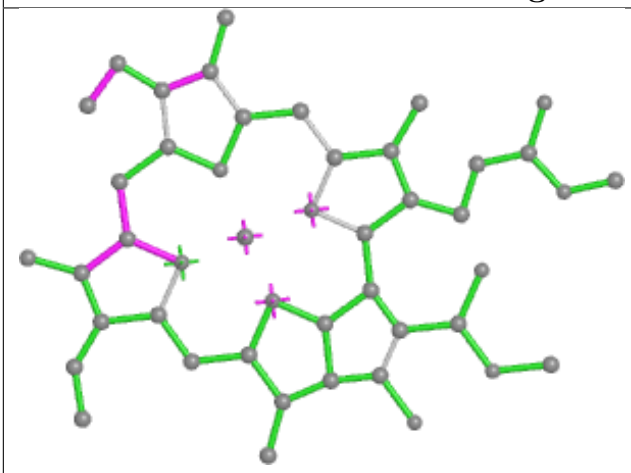
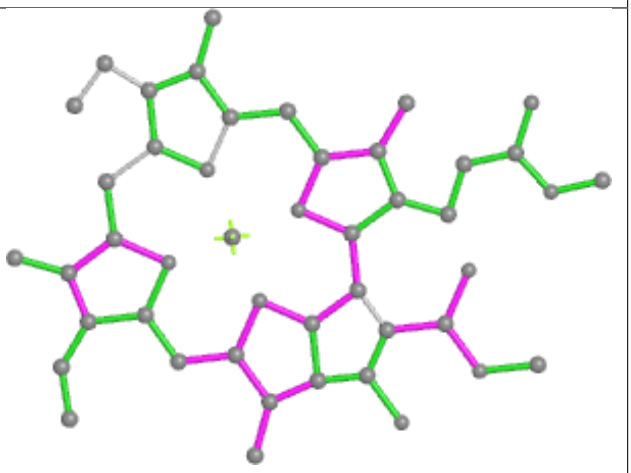


Rings

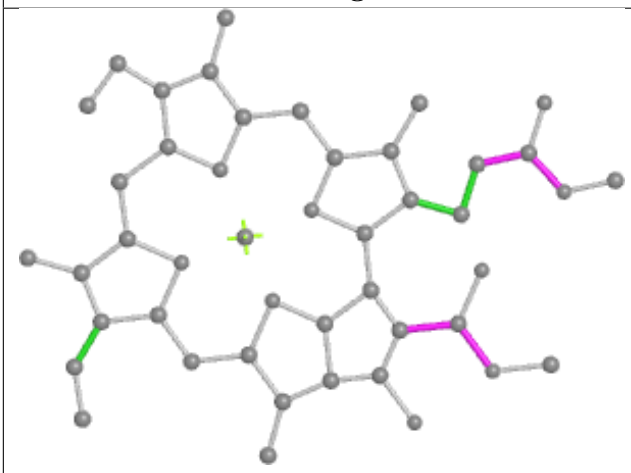
Ligand CLA 5 323



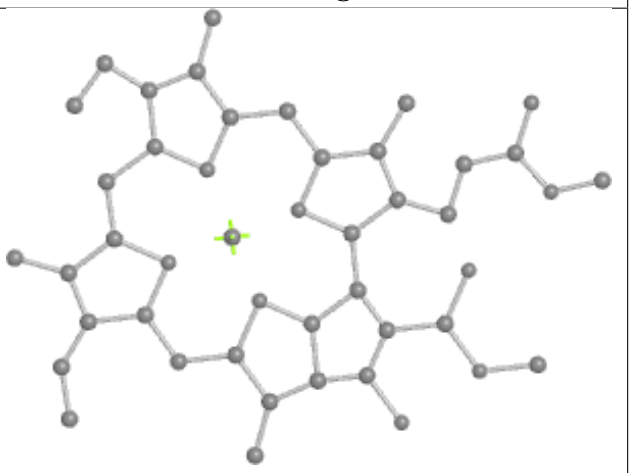
Bond lengths



Bond angles

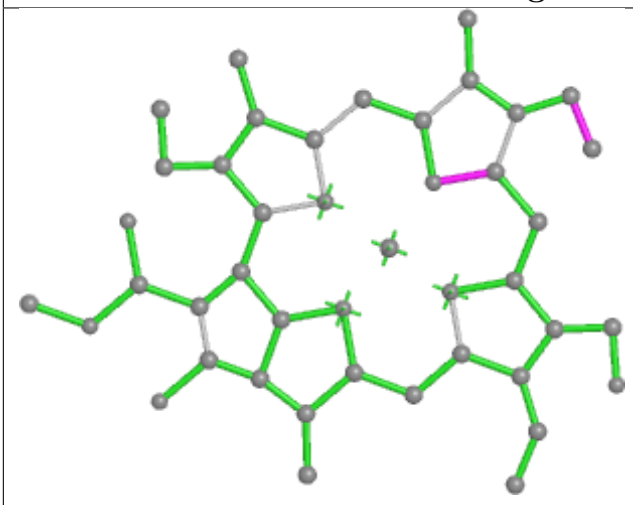


Torsions



Rings

Ligand CHL 4 819



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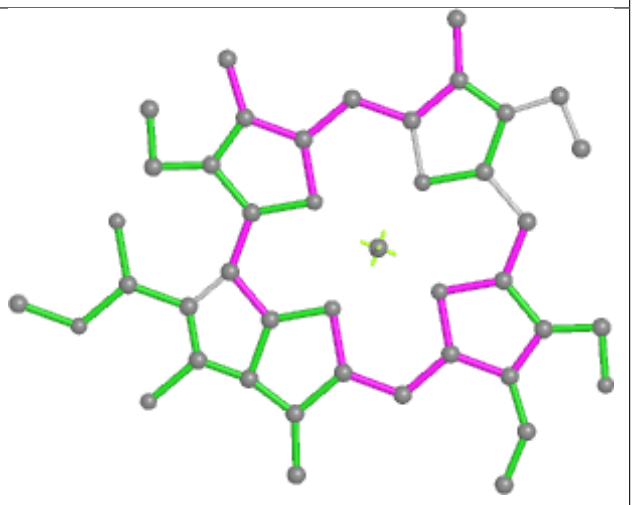
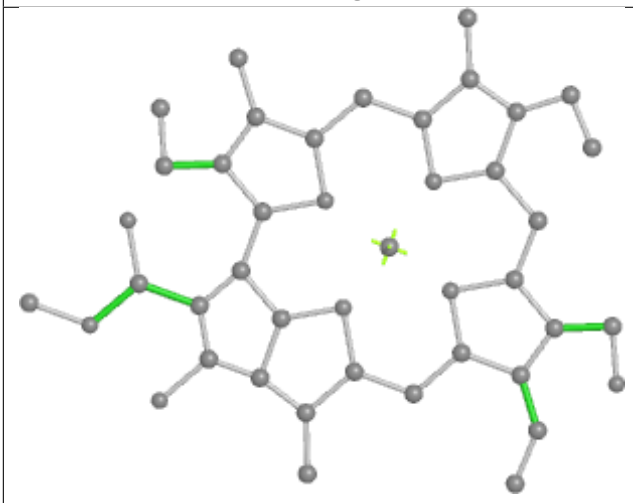


Bond angles



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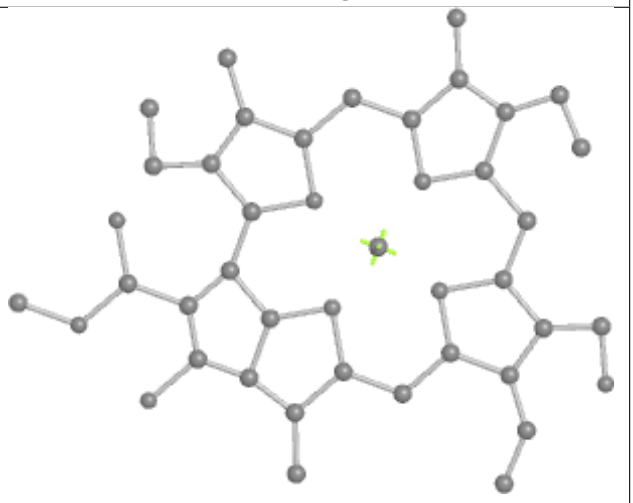
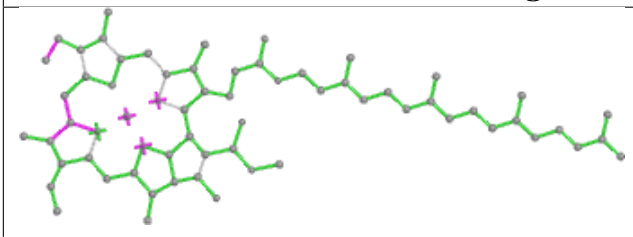


Rings

Ligand CLA B 833



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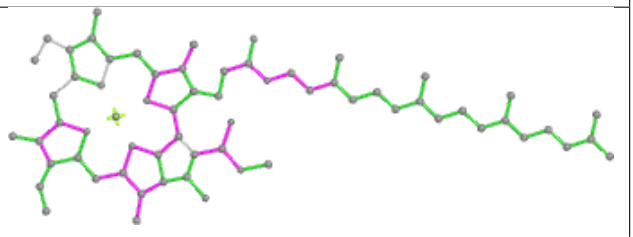
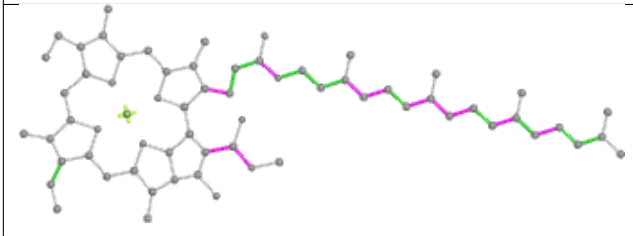
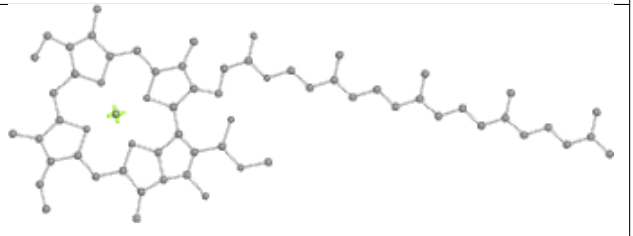


Bond angles

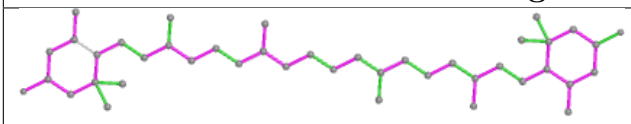


Torsions

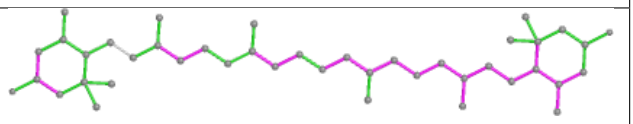


Rings

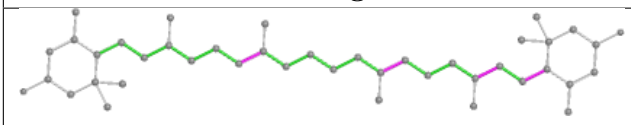
Ligand LUT 1 303



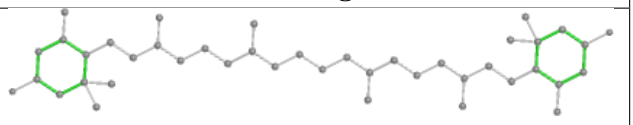
Bond lengths



Bond angles

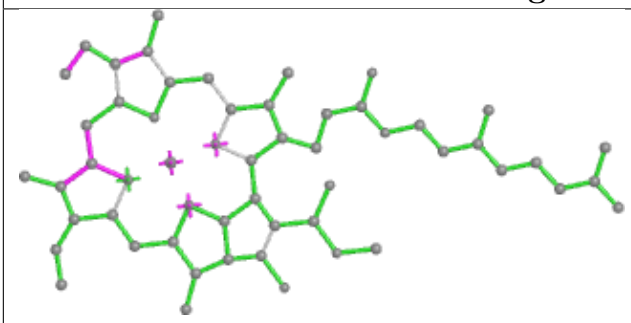


Torsions

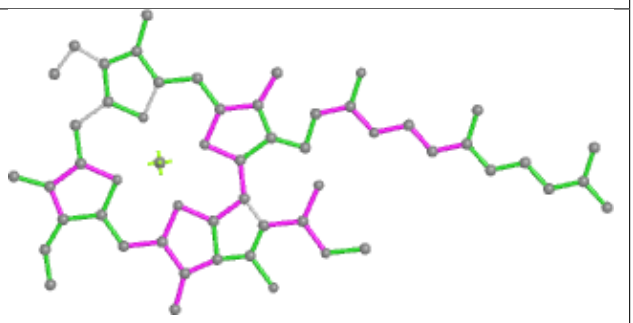


Rings

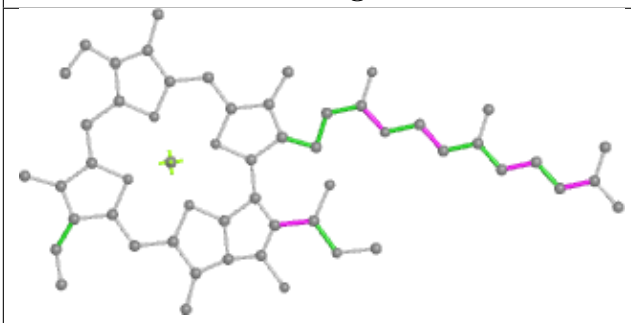
Ligand CLA 6 313



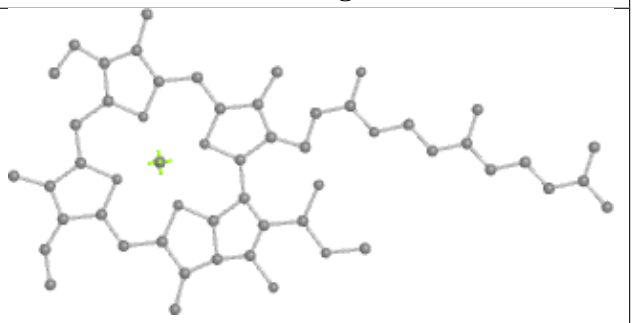
Bond lengths



Bond angles

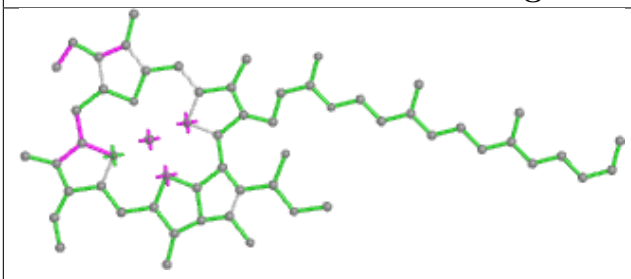


Torsions

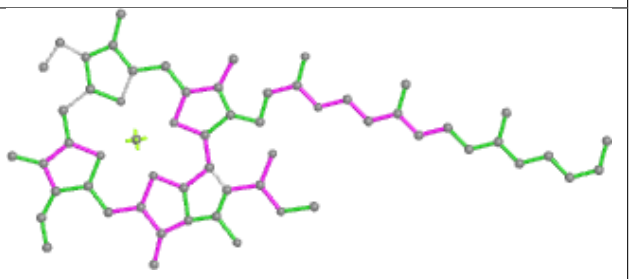


Rings

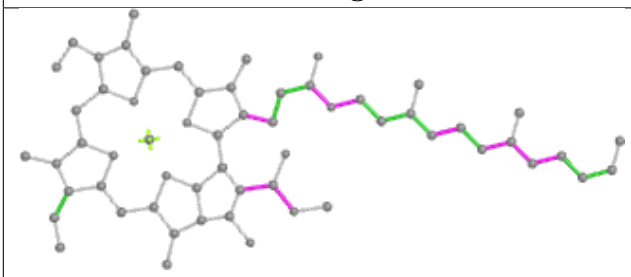
Ligand CLA B 819



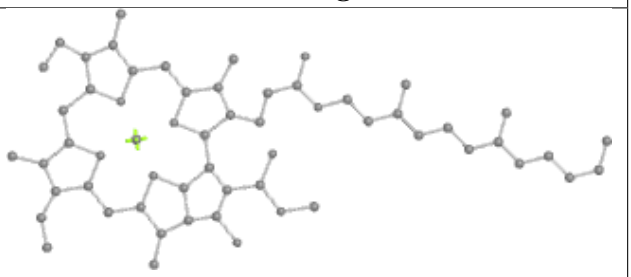
Bond lengths



Bond angles

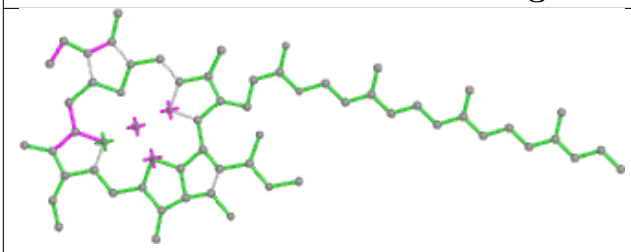


Torsions



Rings

Ligand CLA 8 308



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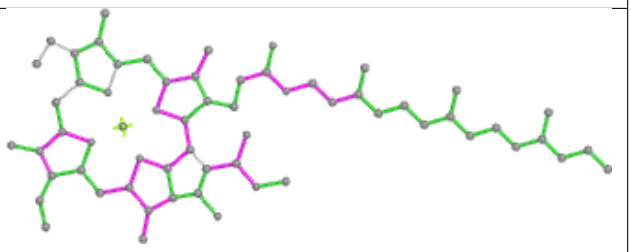
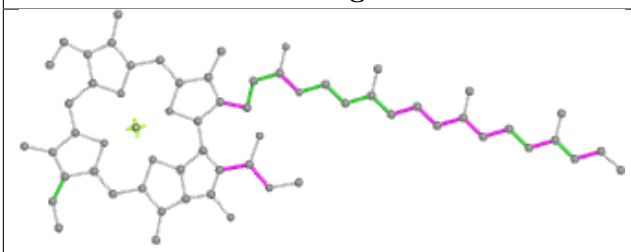
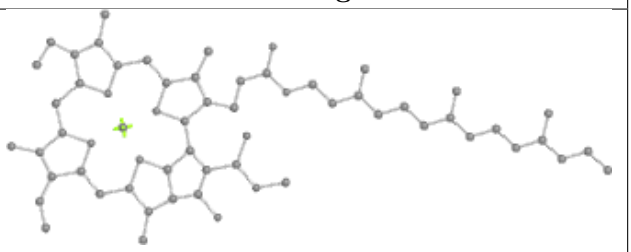


Bond angles

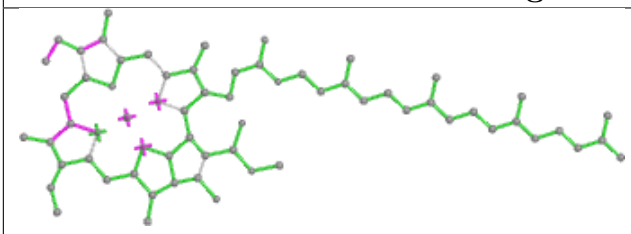


Torsions

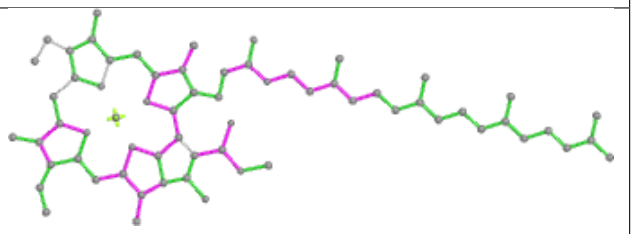


Rings

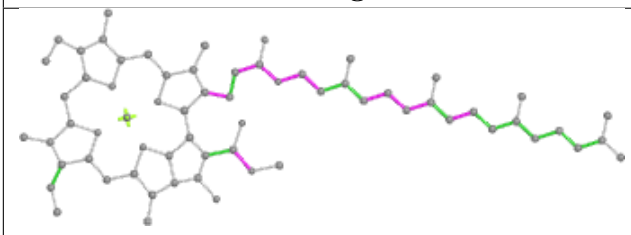
Ligand CLA Z 313



Bond lengths



Bond angles



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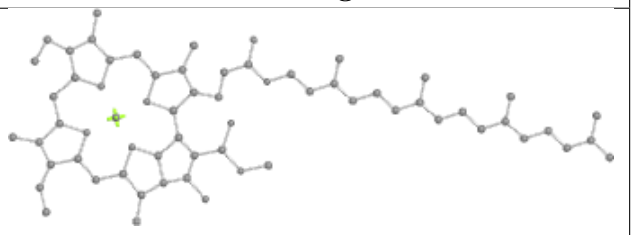
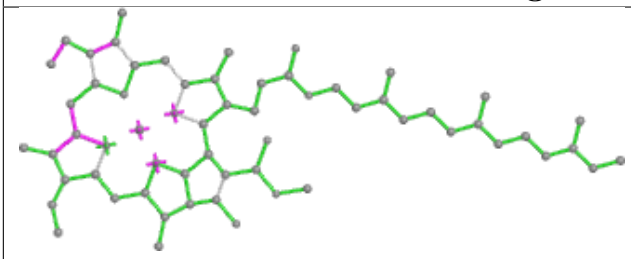
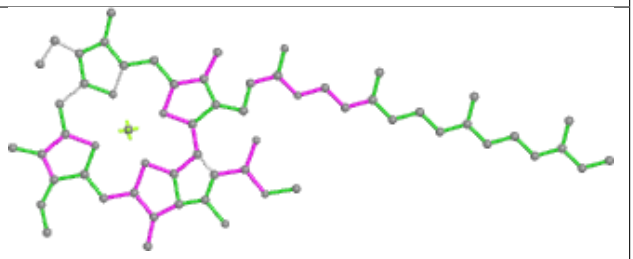


Rings

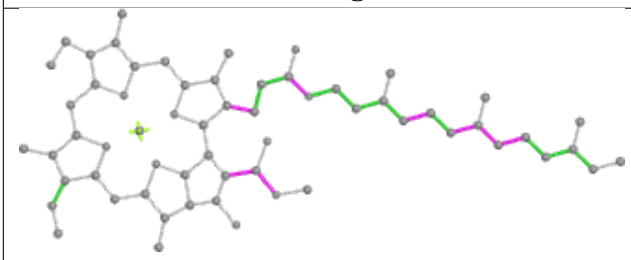
Ligand CLA Z 308



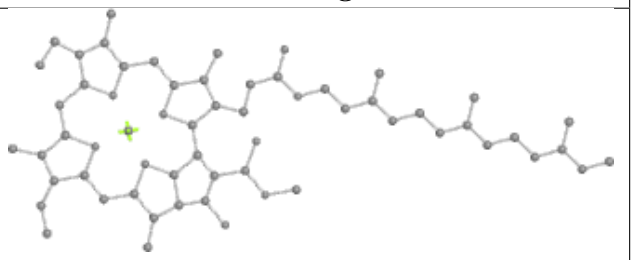
Bond lengths



Bond angles

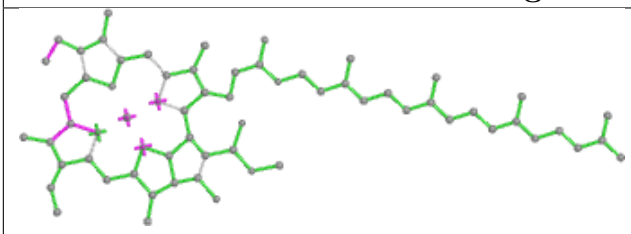


Torsions

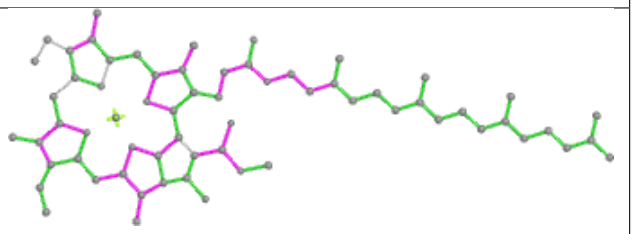


Rings

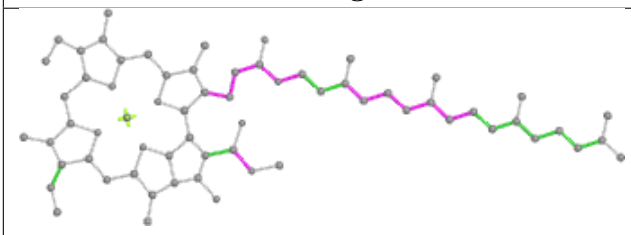
Ligand CLA B 825



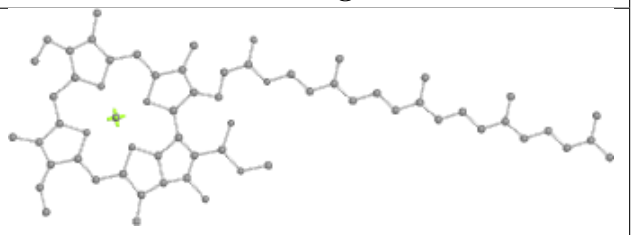
Bond lengths



Bond angles

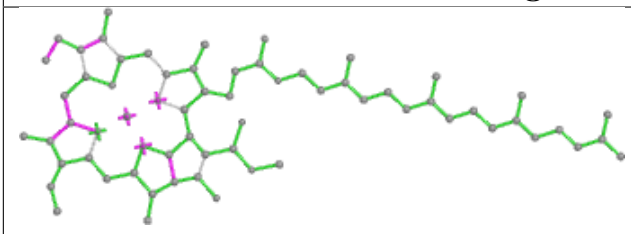


Torsions

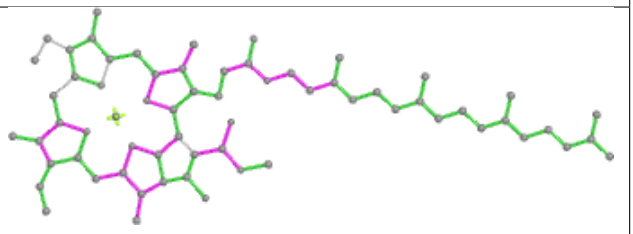


Rings

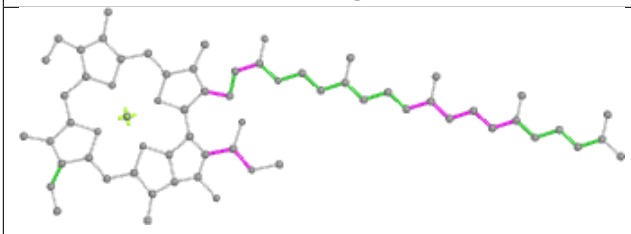
Ligand CLA B 818



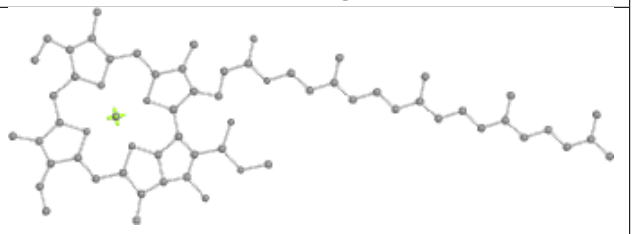
Bond lengths



Bond angles

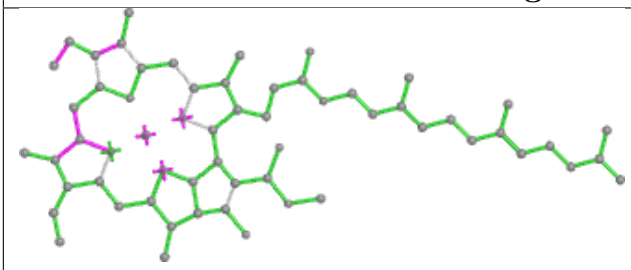


Torsions

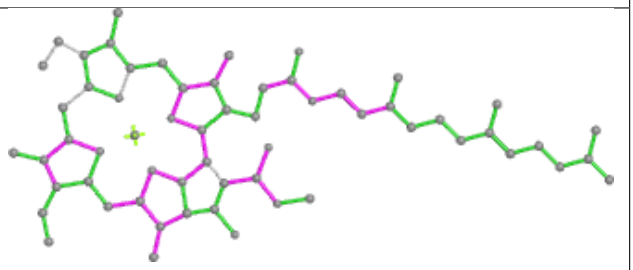


Rings

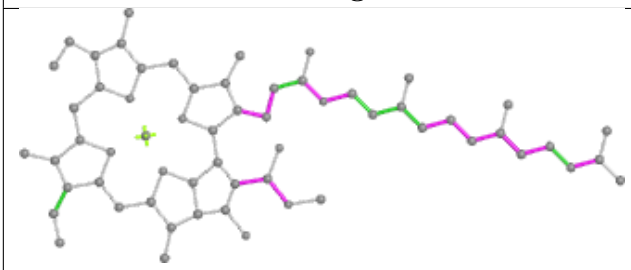
Ligand CLA B 820



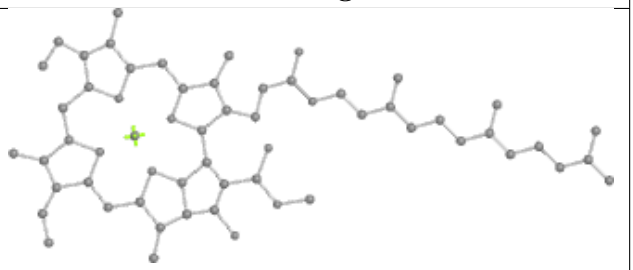
Bond lengths



Bond angles

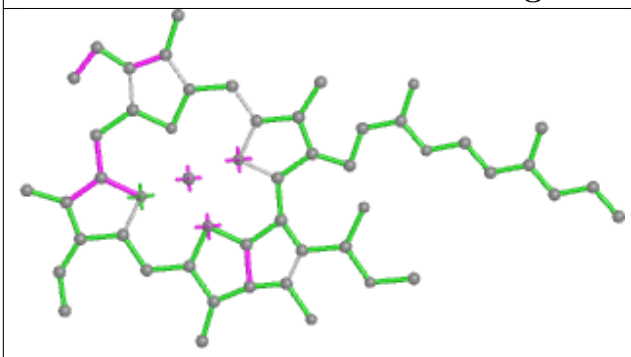


Torsions

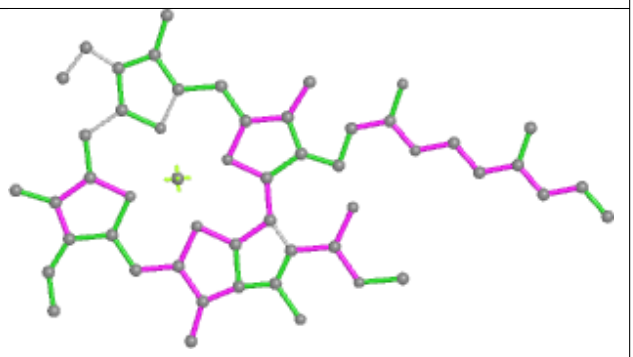


Rings

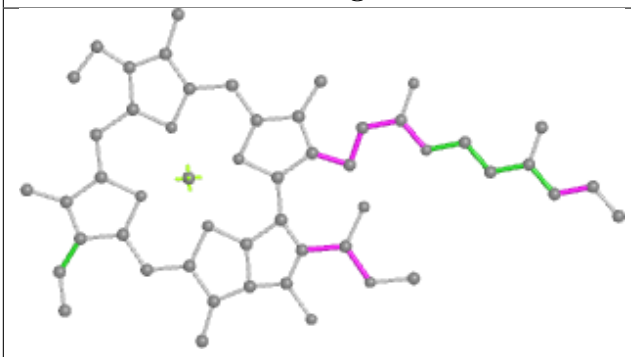
Ligand CLA B 839



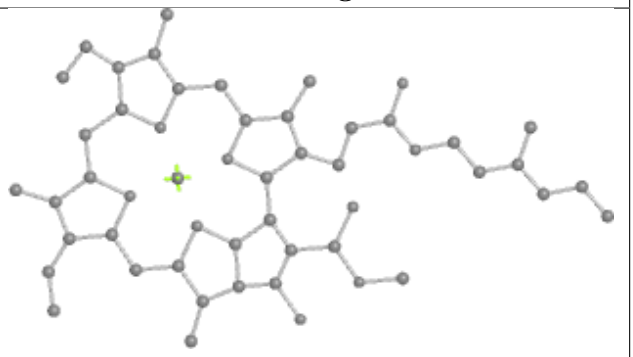
Bond lengths



Bond angles

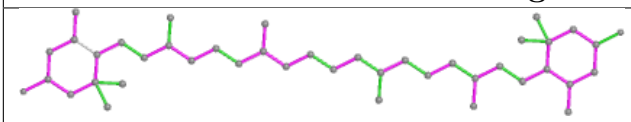


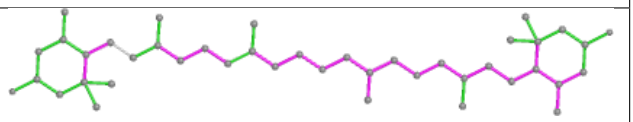
Torsions

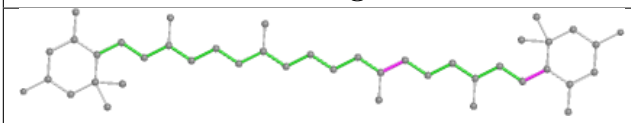


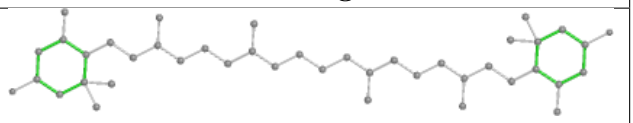
Rings

Ligand LUT 8 303

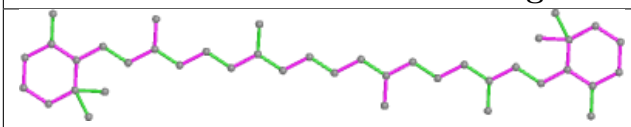

Bond lengths


Bond angles


Torsions


Rings

Ligand BCR 5 304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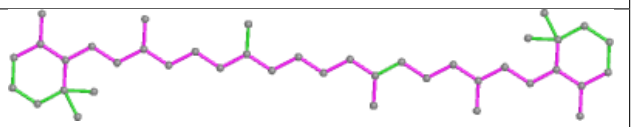
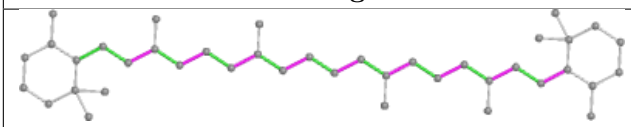
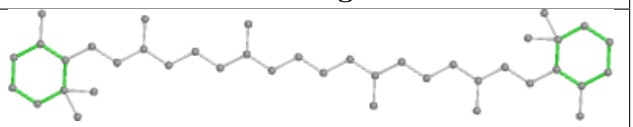
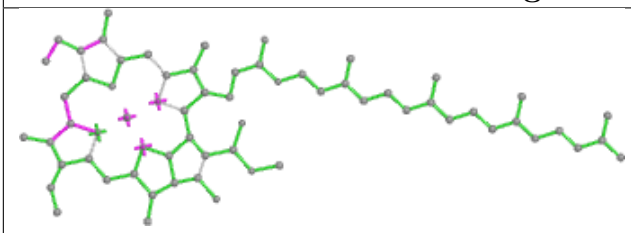
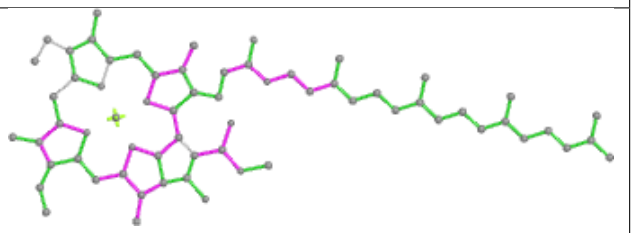

Bond angles

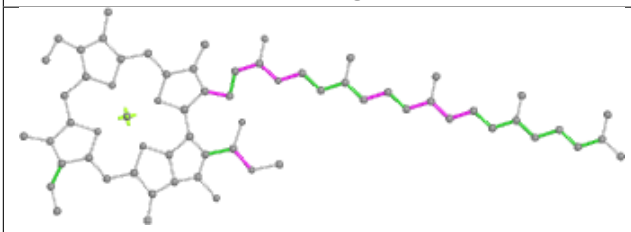

Torsions

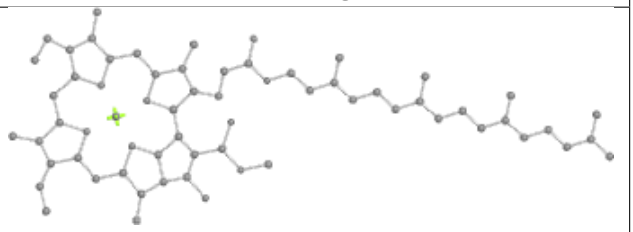

Rings

Ligand CLA A 833

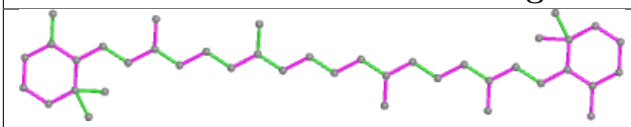

Bond lengths

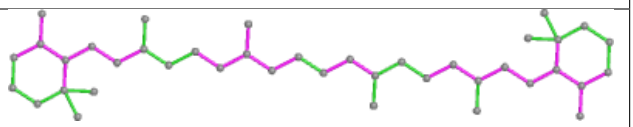

Bond angles

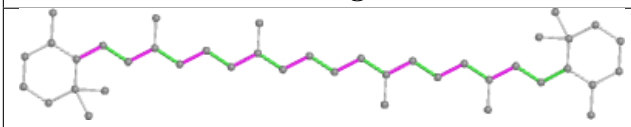

Torsions


Rings

Ligand BCR 3 306

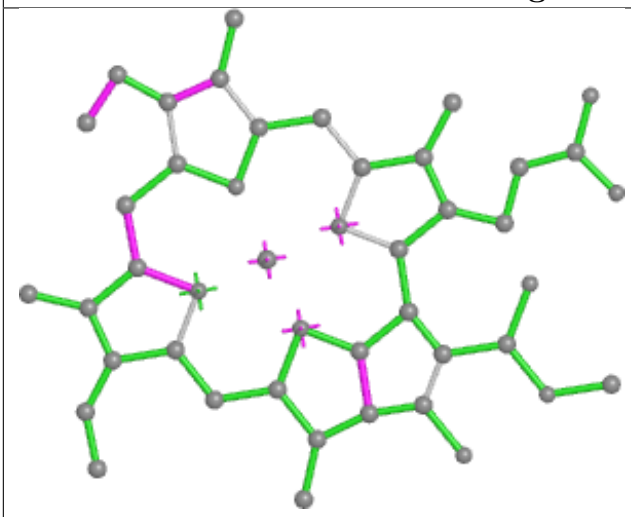

Bond lengths


Bond angles

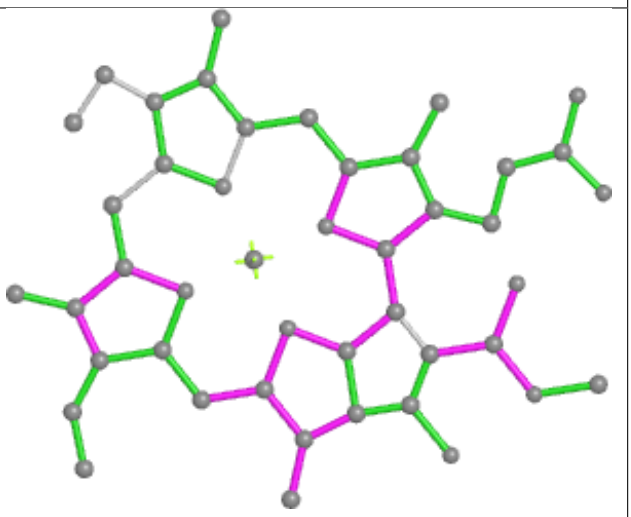

Torsions


Rings

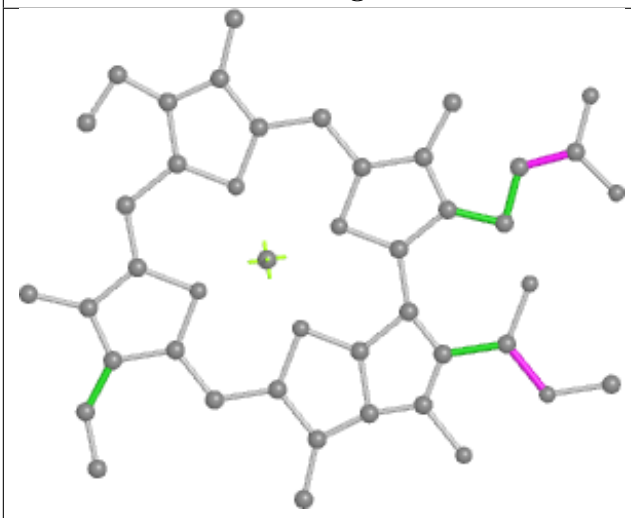
Ligand CLA 9 304



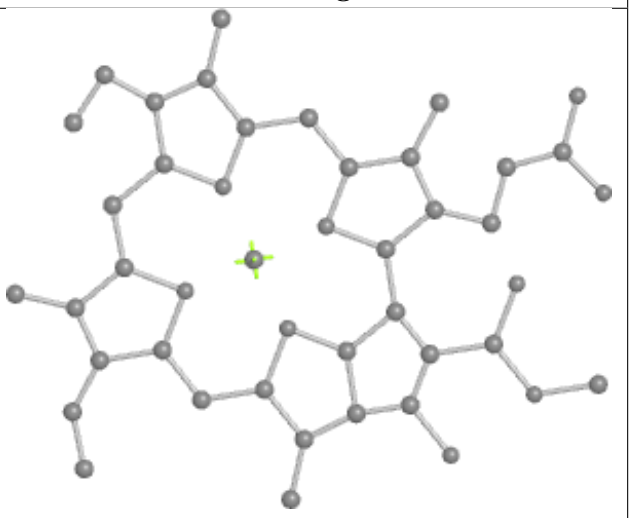
Bond lengths



Bond angles

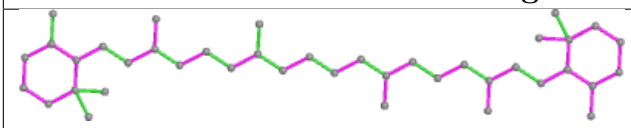


Torsions



Rings

Ligand BCR G 203



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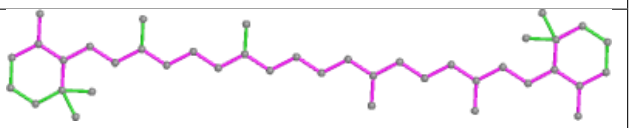
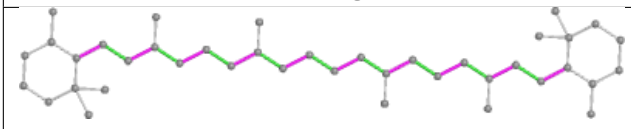


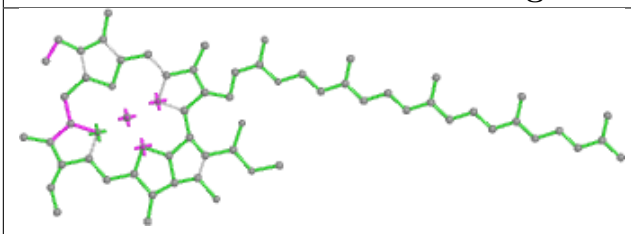
Bond angles



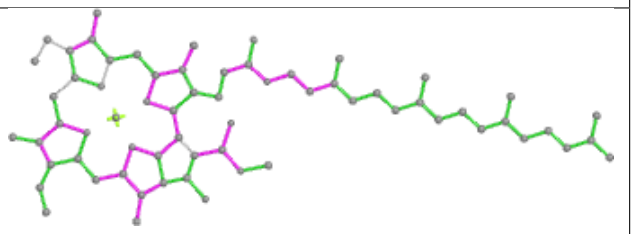
Torsions

Rings

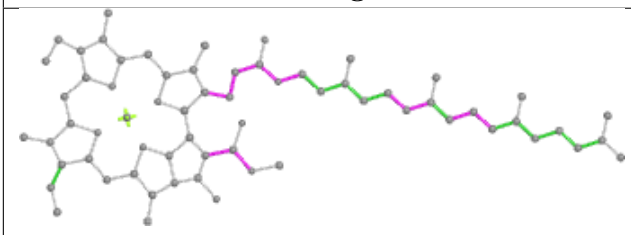
Ligand CLA Z 303



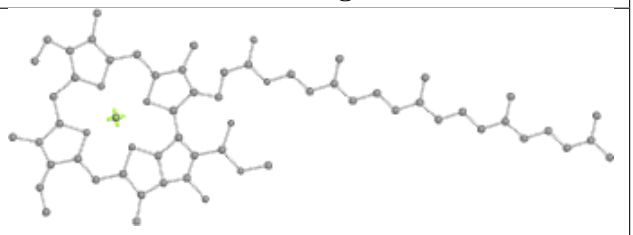
Bond lengths



Bond angles

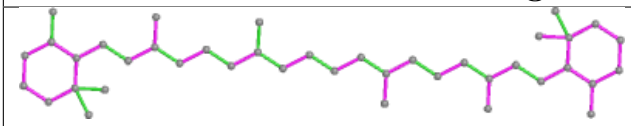


Torsions

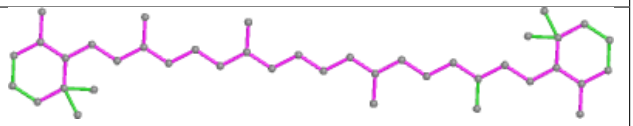


Rings

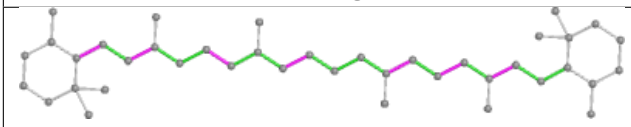
Ligand BCR 3 304



Bond lengths



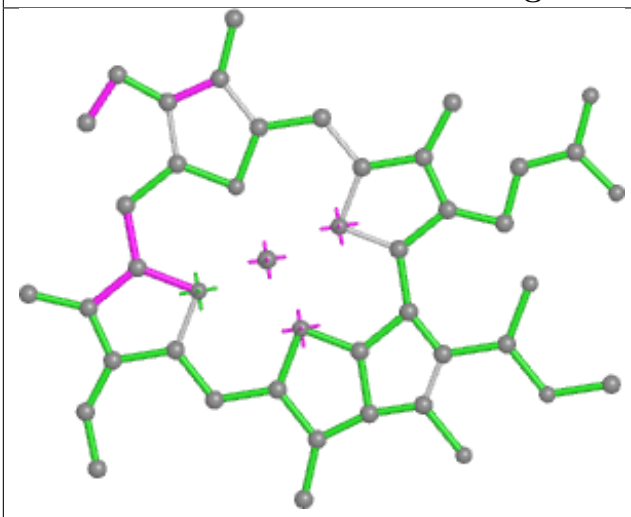
Bond angles



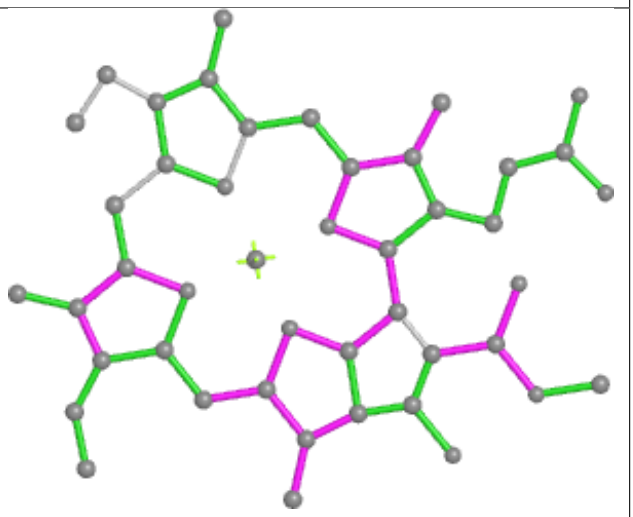
Torsions

Rings

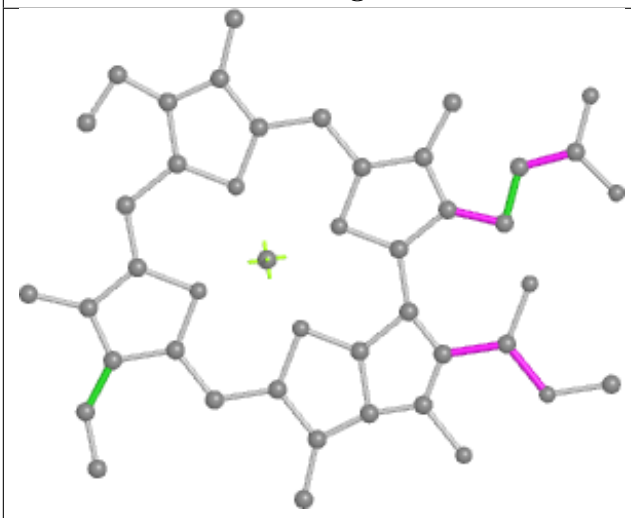
Ligand CLA B 807



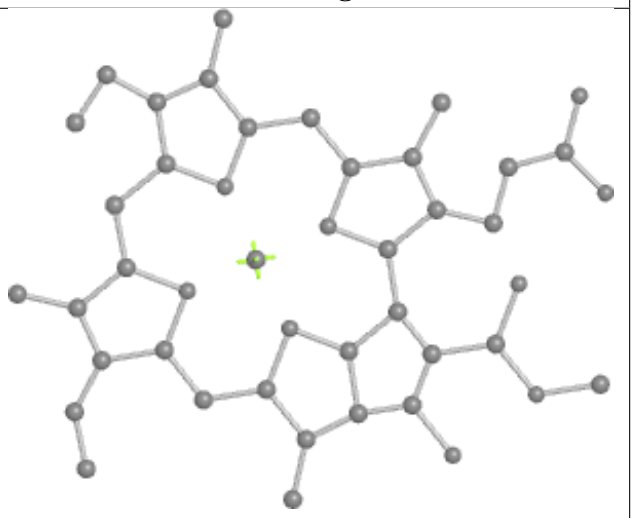
Bond lengths



Bond angles

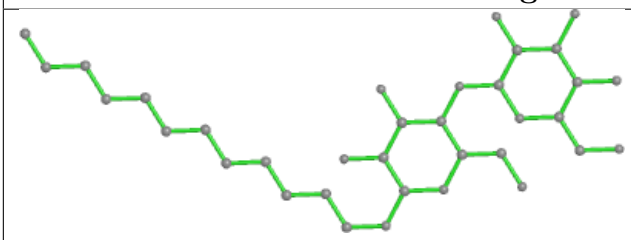


Torsions

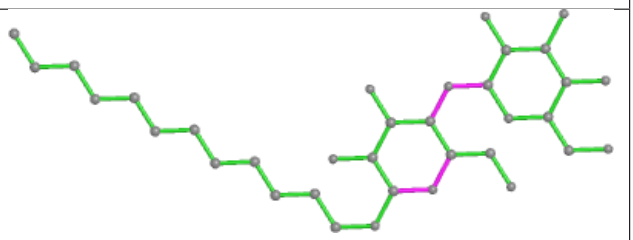


Rings

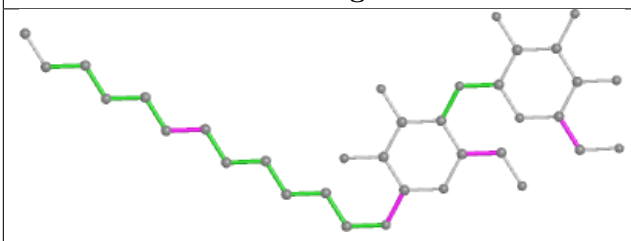
Ligand LMT G 204



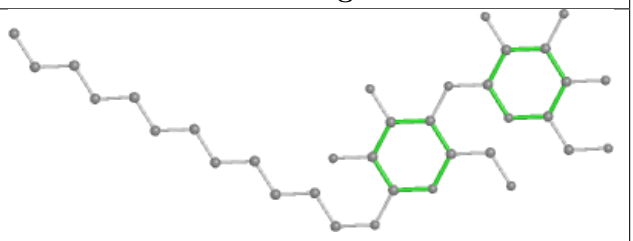
Bond lengths



Bond angles

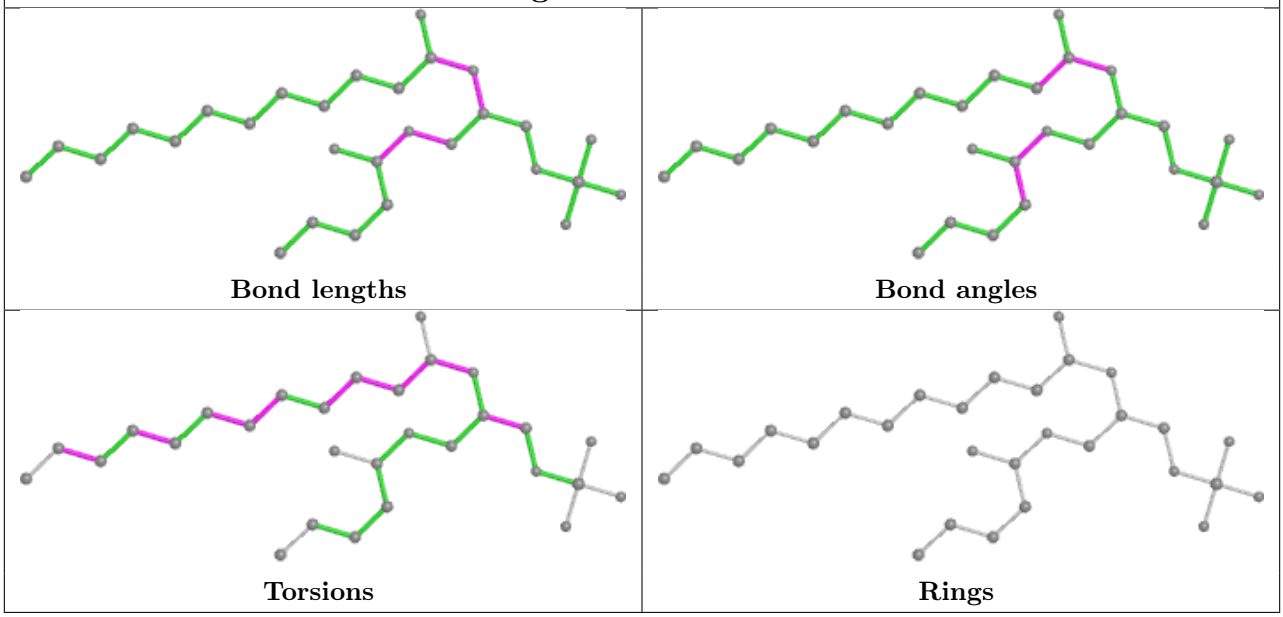


Torsions

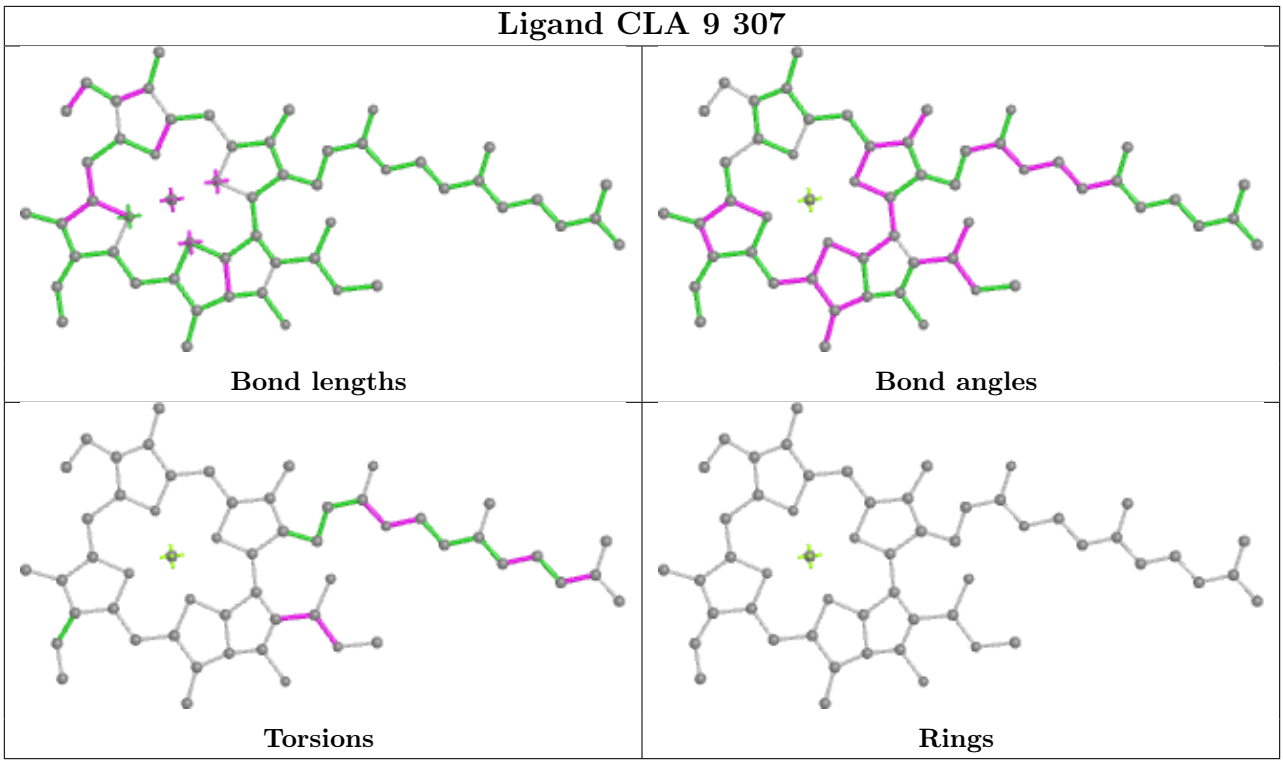


Rings

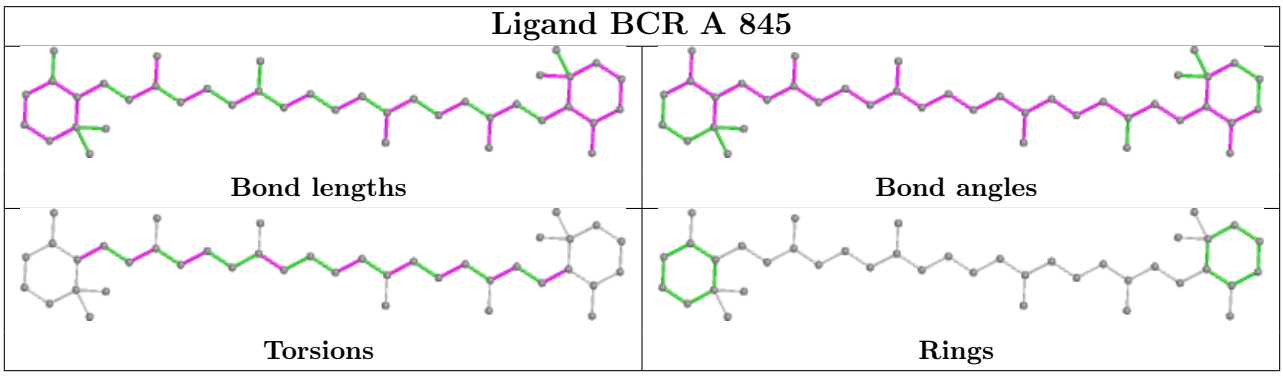
Ligand 3PH 6 324



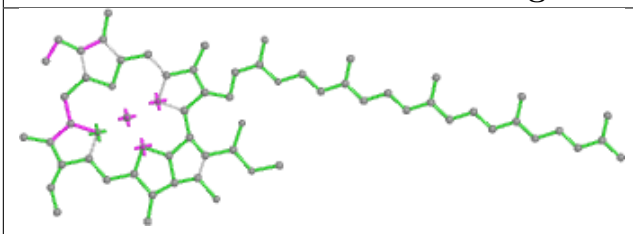
Ligand CLA 9 307



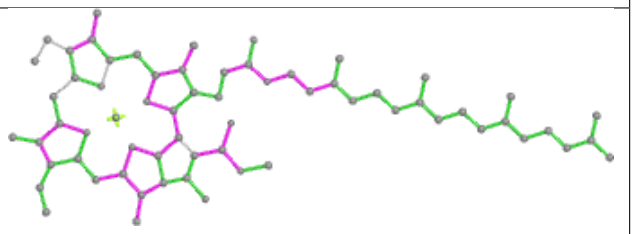
Ligand BCR A 845



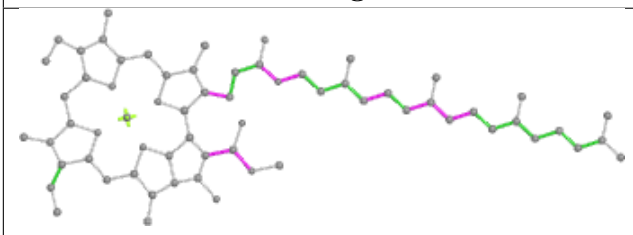
Ligand CLA 6 322



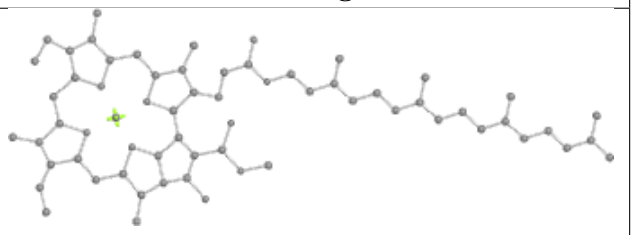
Bond lengths



Bond angles

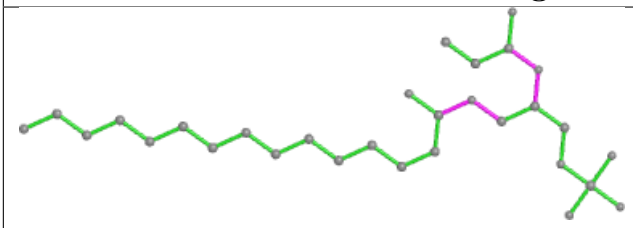


Torsions

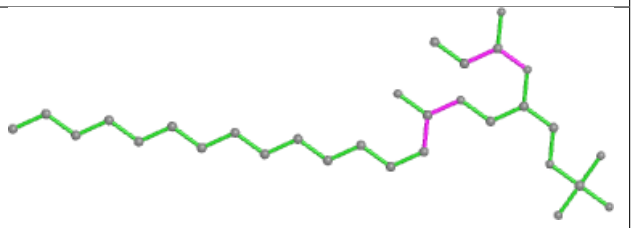


Rings

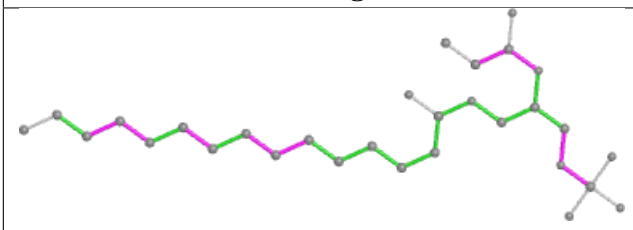
Ligand 3PH 8 320



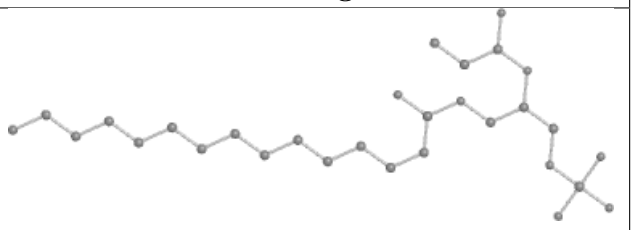
Bond lengths



Bond angles

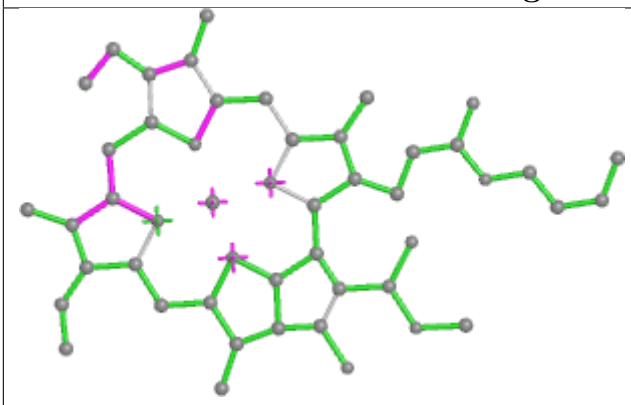


Torsions

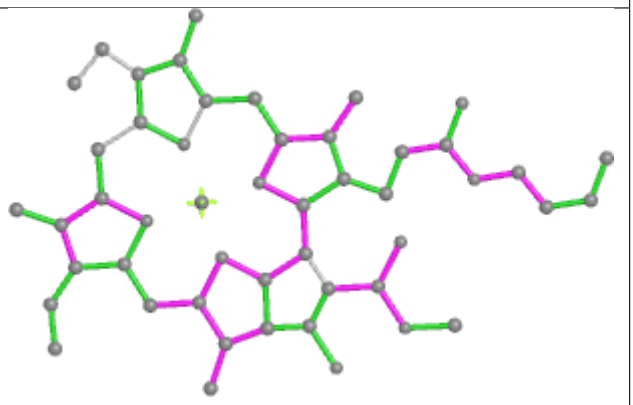


Rings

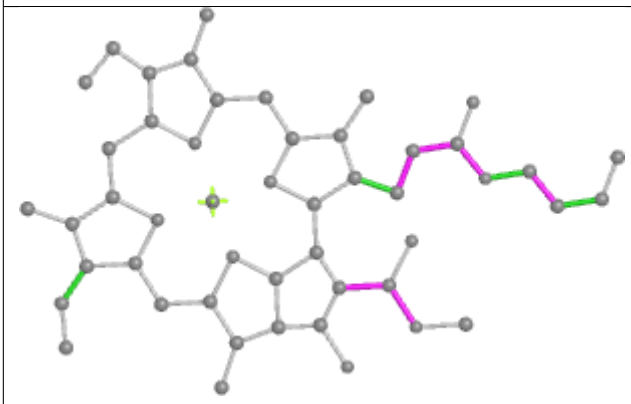
Ligand CLA K 204



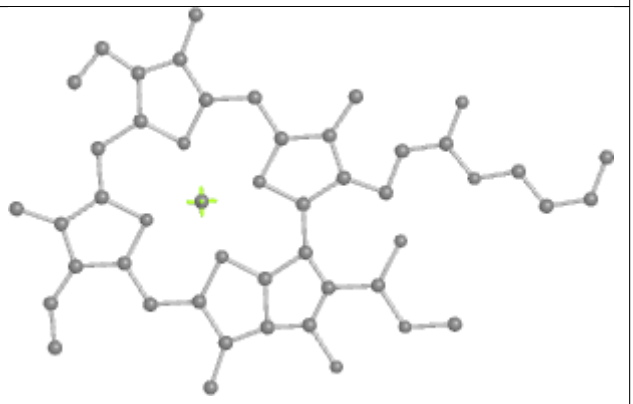
Bond lengths



Bond angles

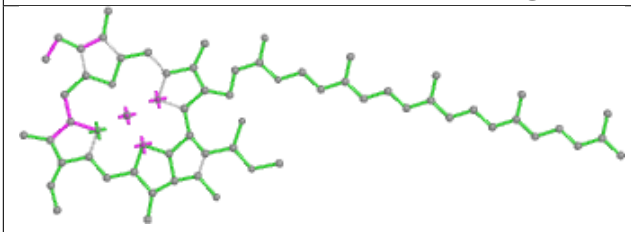


Torsions

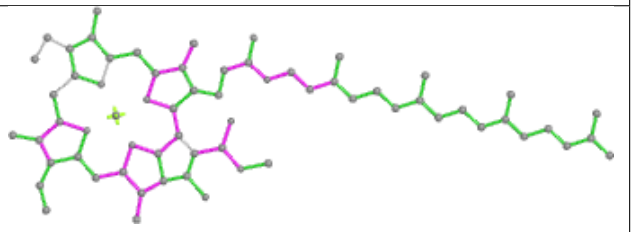


Rings

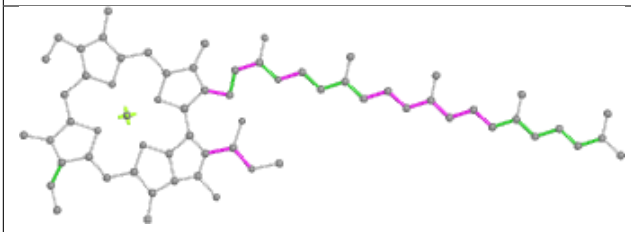
Ligand CLA A 838



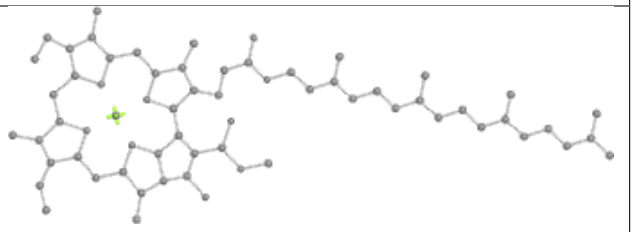
Bond lengths



Bond angles

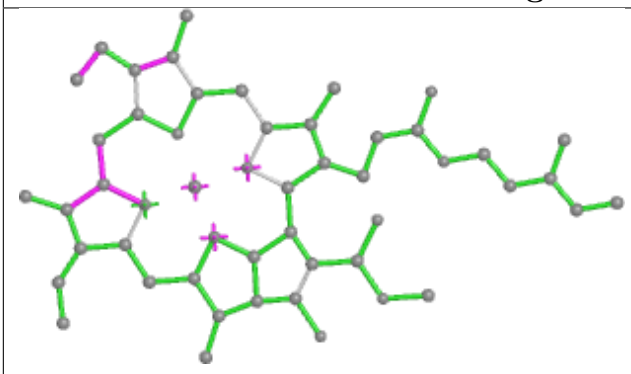


Torsions

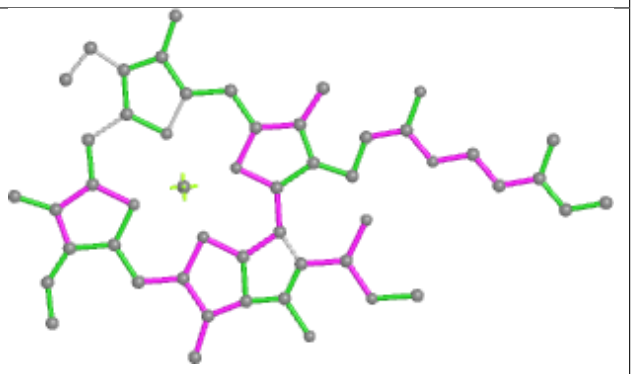


Rings

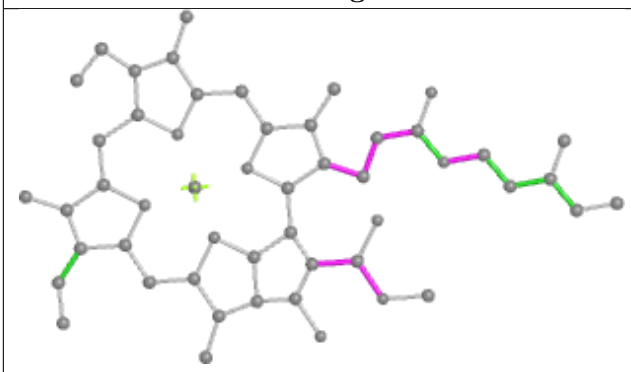
Ligand CLA Z 314



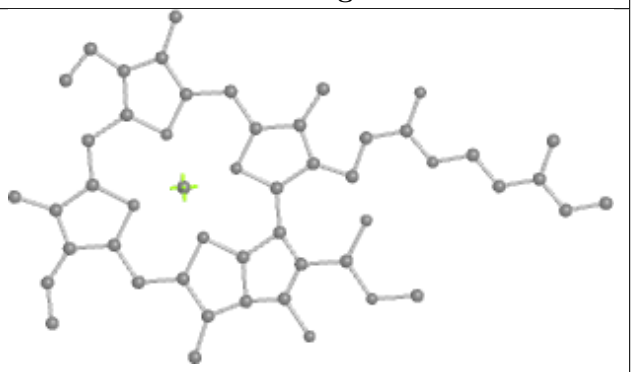
Bond lengths



Bond angles

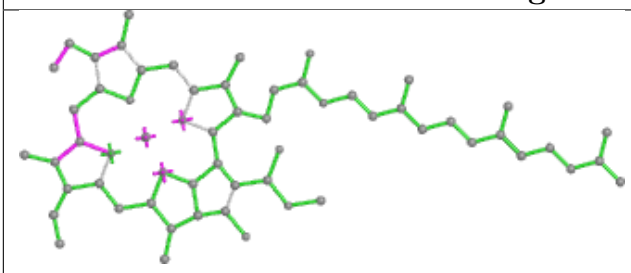


Torsions

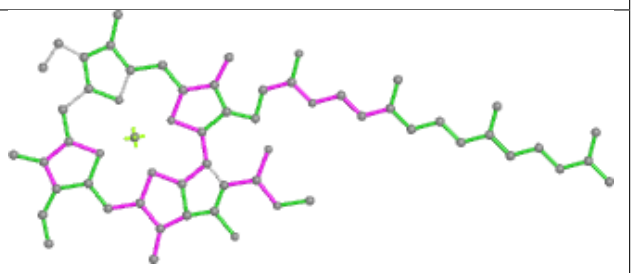


Rings

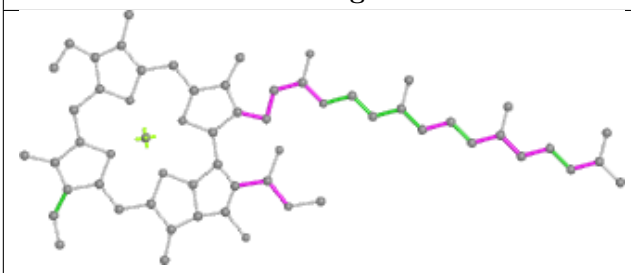
Ligand CLA 7 312



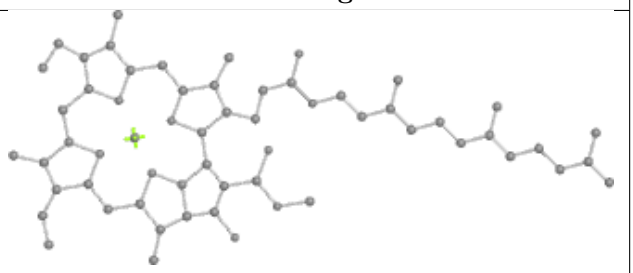
Bond lengths



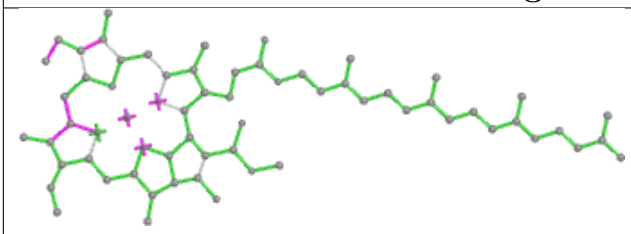
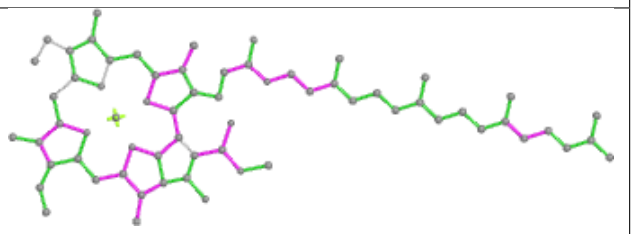
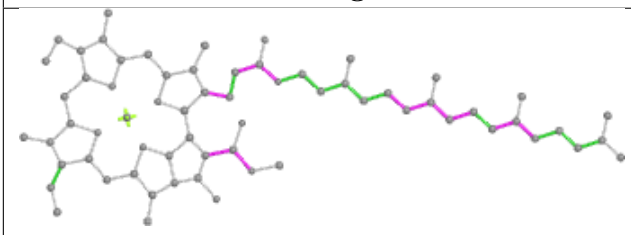
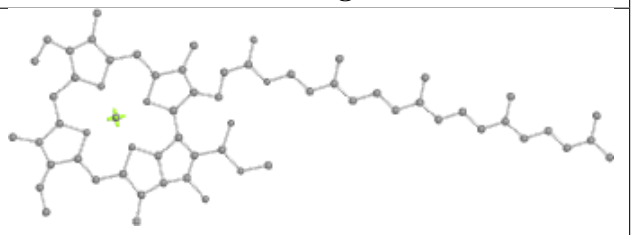
Bond angles

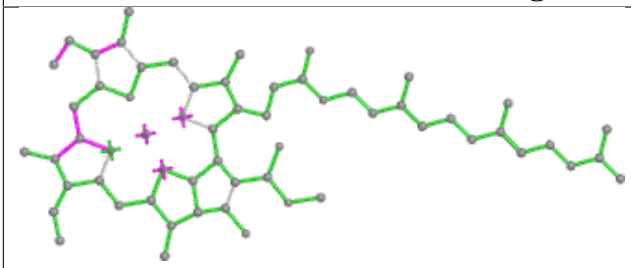
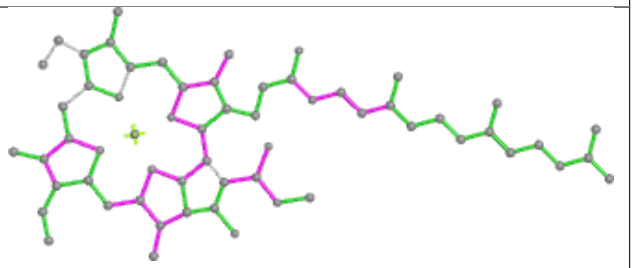
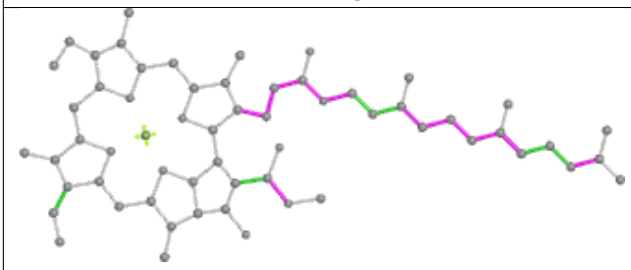
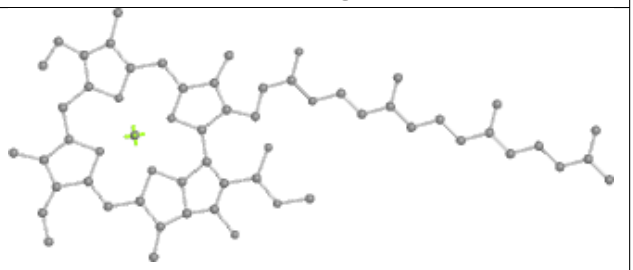


Torsions

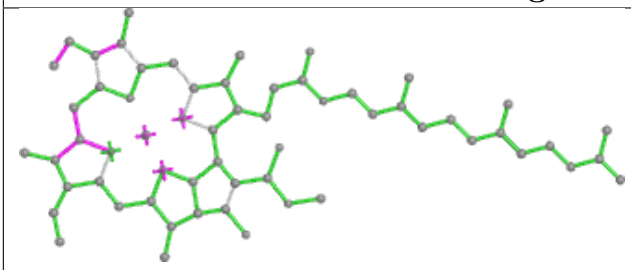


Rings

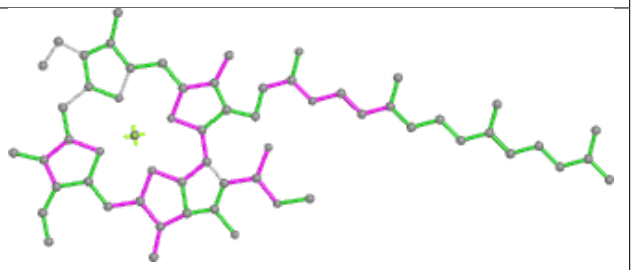
Ligand CLA B 826	
 Bond lengths	 Bond angles
 Torsions	 Rings

Ligand CLA 3 314	
 Bond lengths	 Bond angles
 Torsions	 Rings

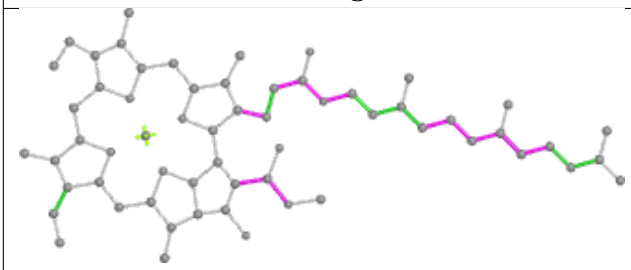
Ligand CLA 6 302



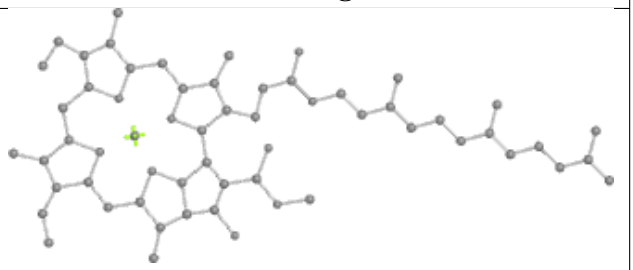
Bond lengths



Bond angles

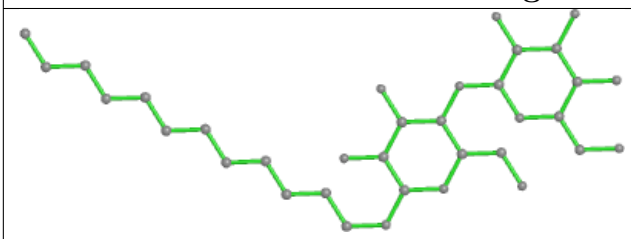


Torsions

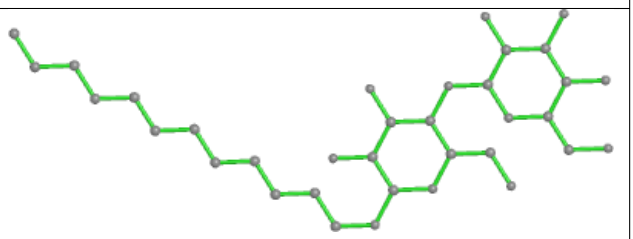


Rings

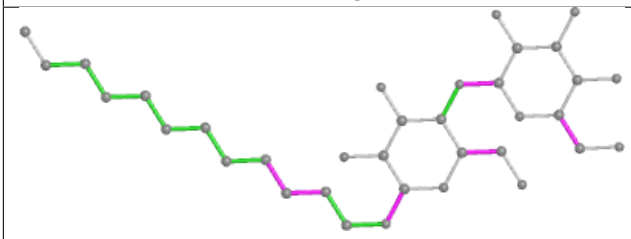
Ligand LMT F 307



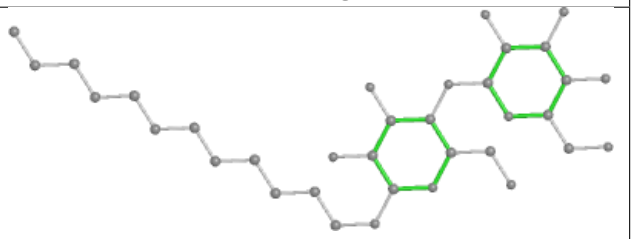
Bond lengths



Bond angles

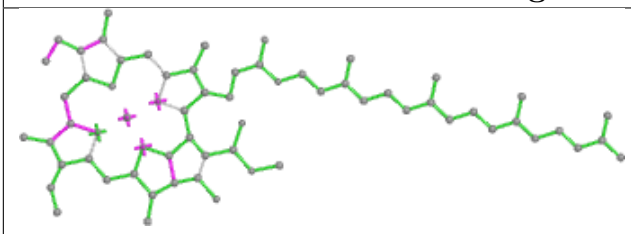


Torsions

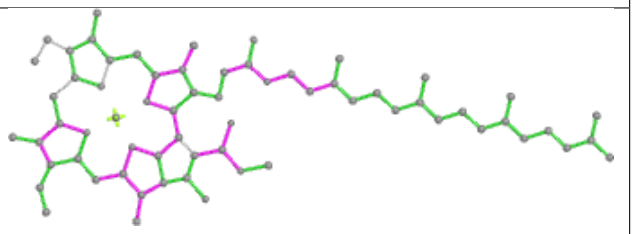


Rings

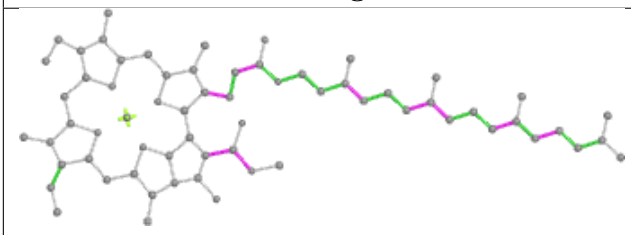
Ligand CLA 3 322



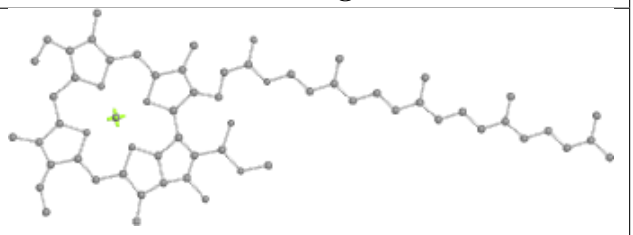
Bond lengths



Bond angles

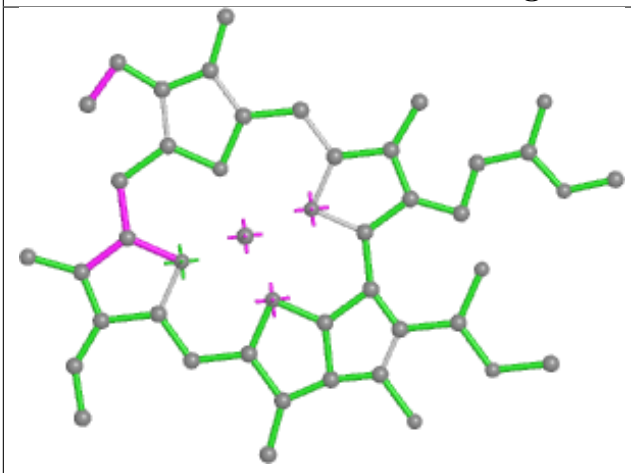


Torsions

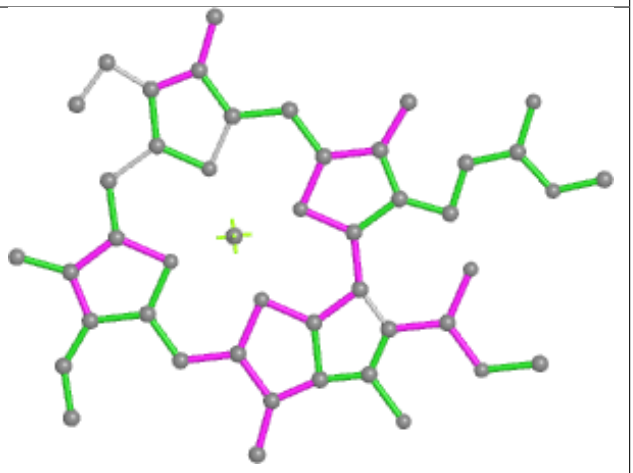


Rings

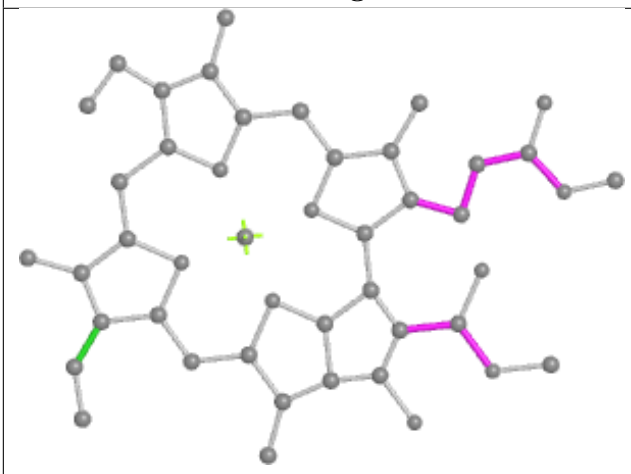
Ligand CLA K 202



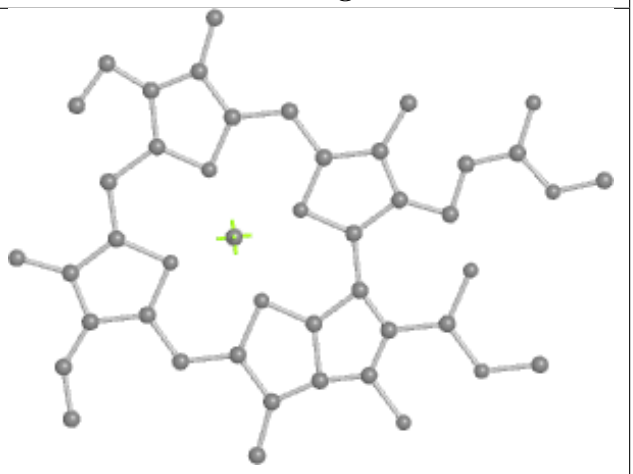
Bond lengths



Bond angles

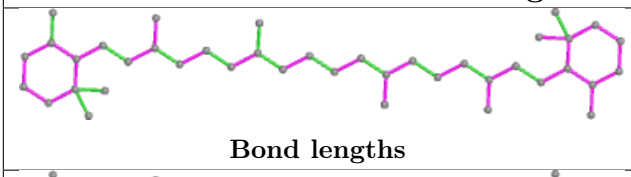


Torsions



Rings

Ligand BCR 8 304



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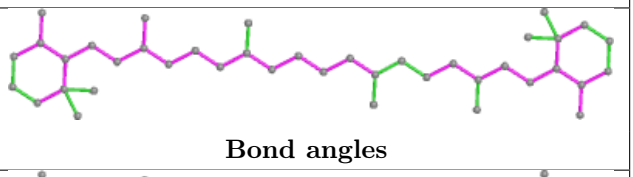
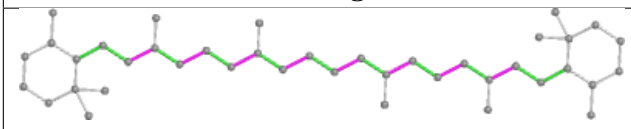
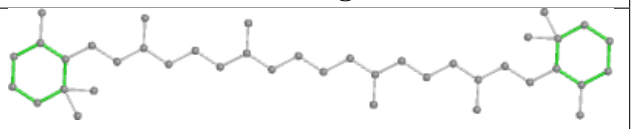


Bond angles

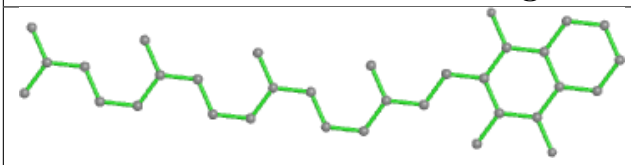


Torsions

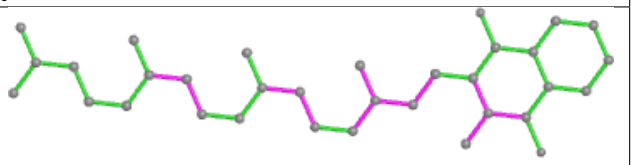


Rings

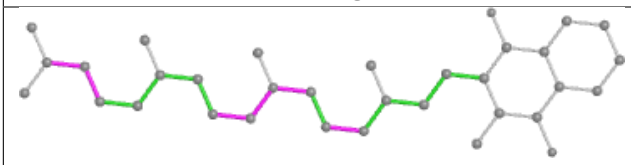
Ligand PQN A 843



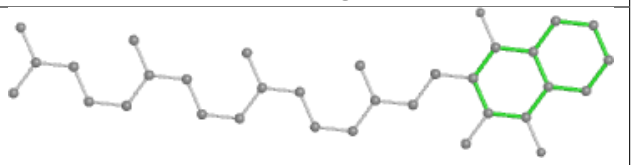
Bond lengths



Bond angles

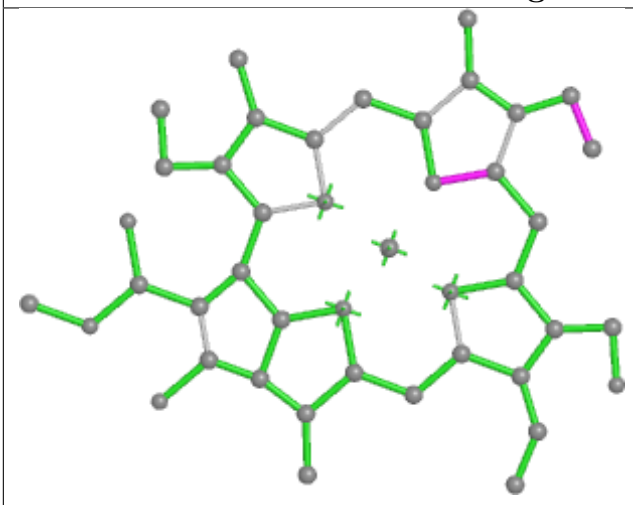


Torsions



Rings

Ligand CHL 5 321



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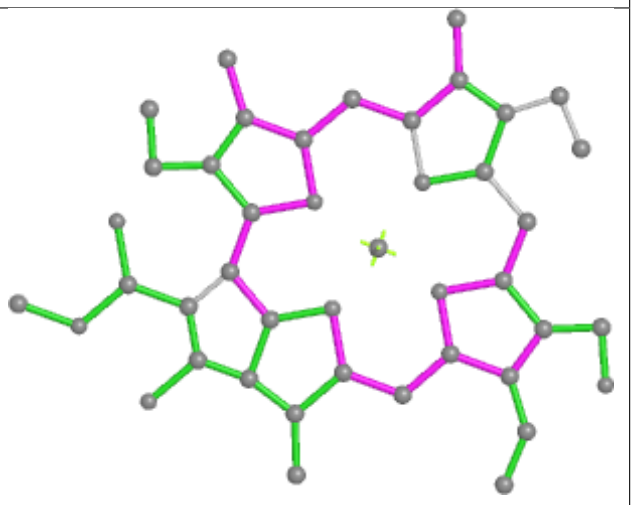
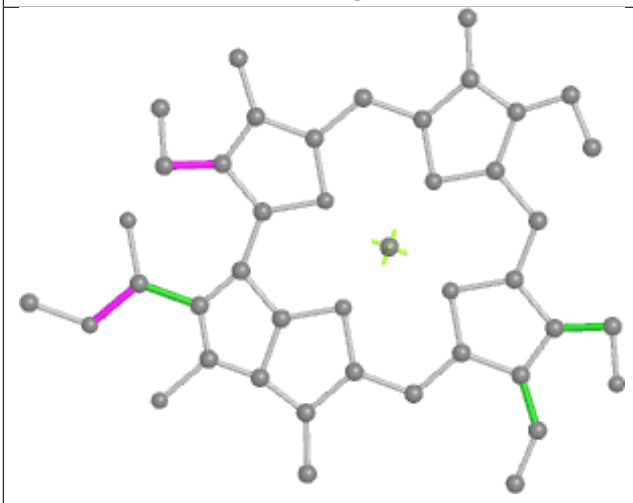
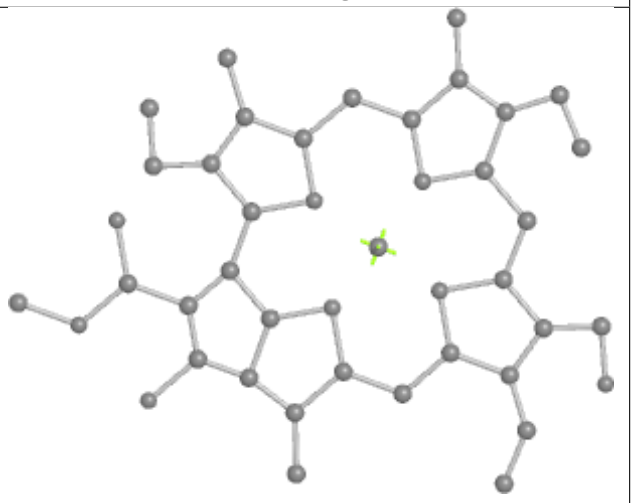


Bond angles

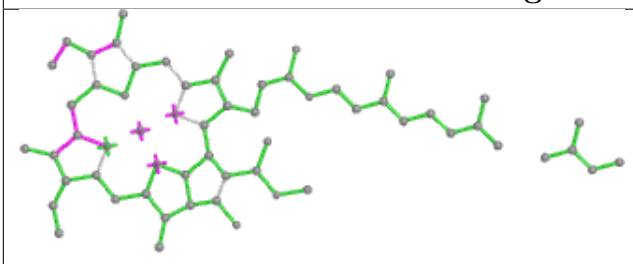


Torsions

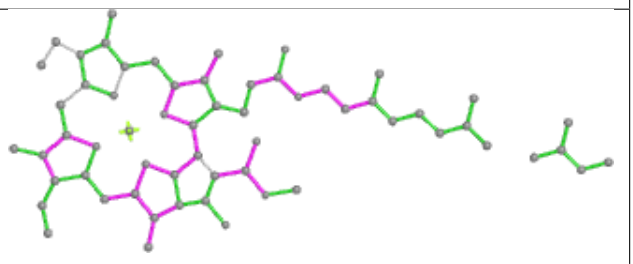


Rings

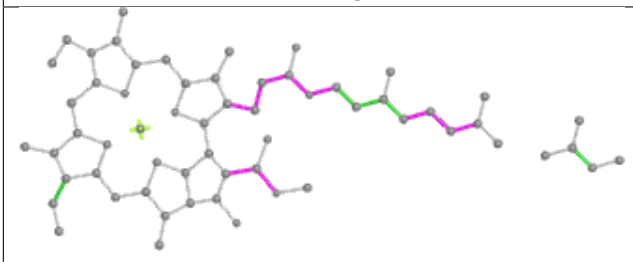
Ligand CLA B 812



Bond lengths



Bond angles



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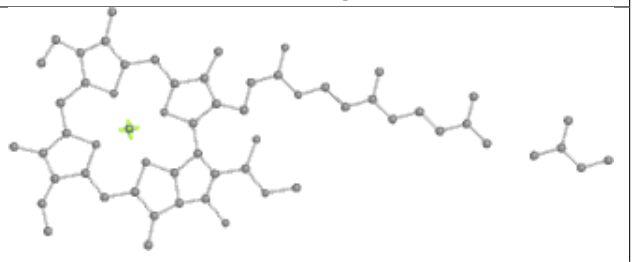
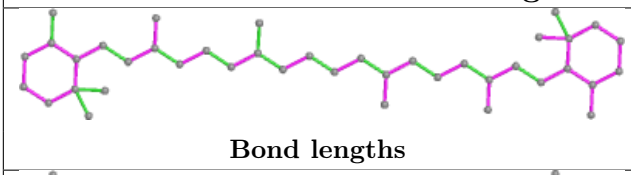
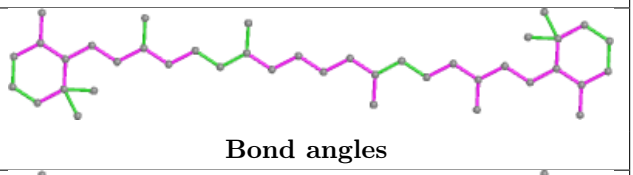


Rings

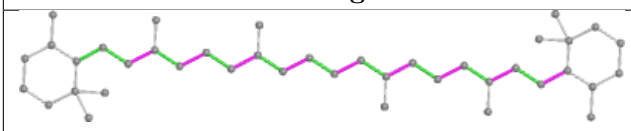
Ligand BCR B 849



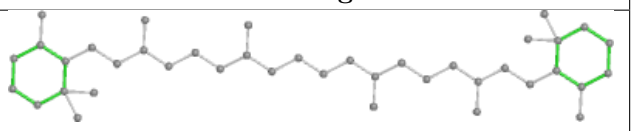
Bond lengths



Bond angles

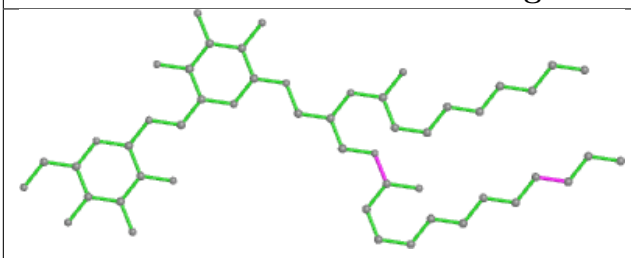


Torsions

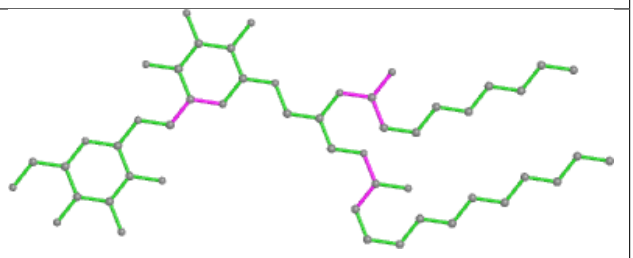


Rings

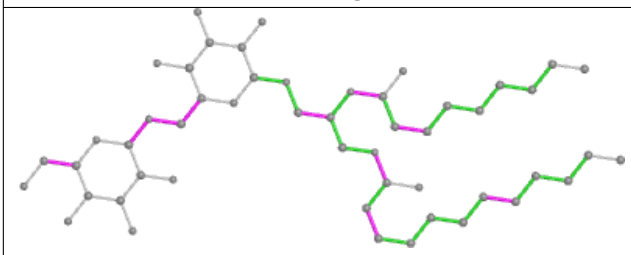
Ligand DGD 1 319



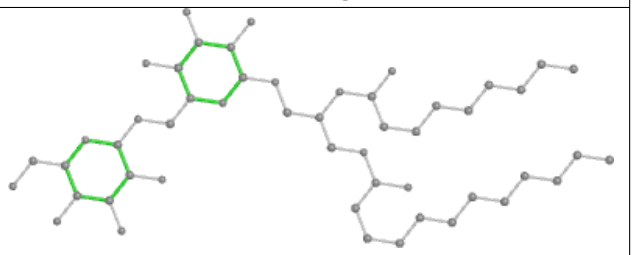
Bond lengths



Bond angles

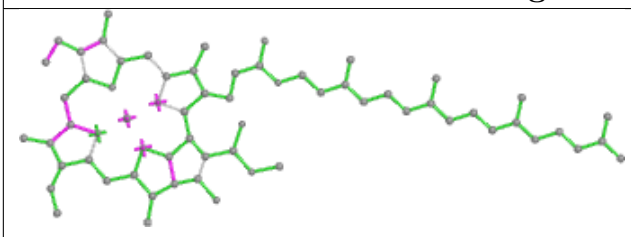


Torsions

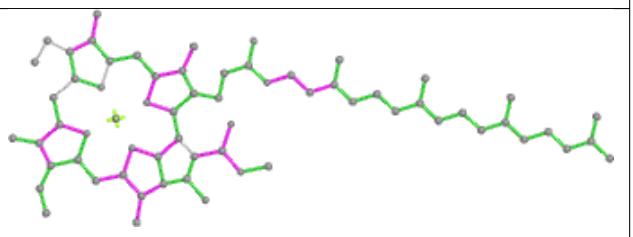


Rings

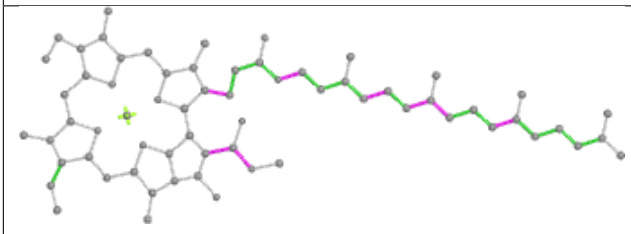
Ligand CLA 7 307



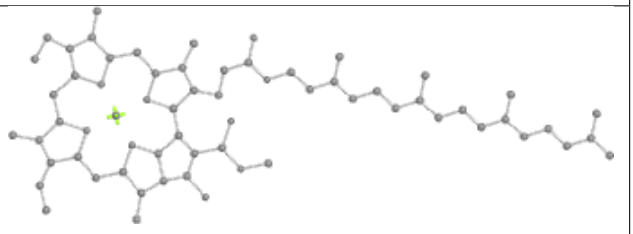
Bond lengths



Bond angles

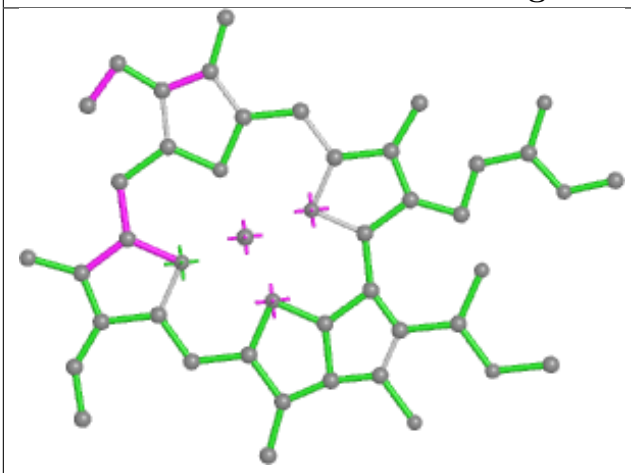


Torsions

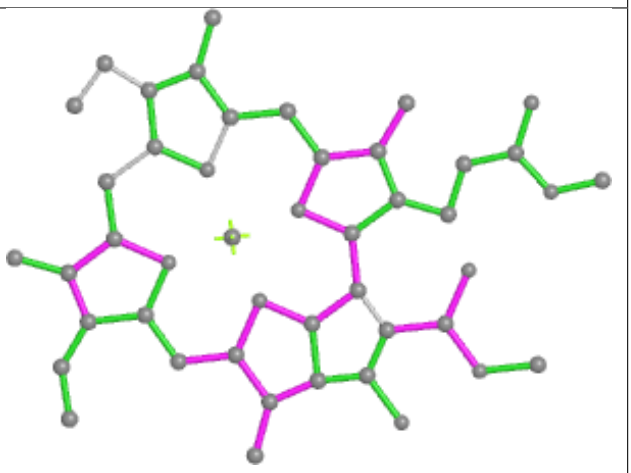


Rings

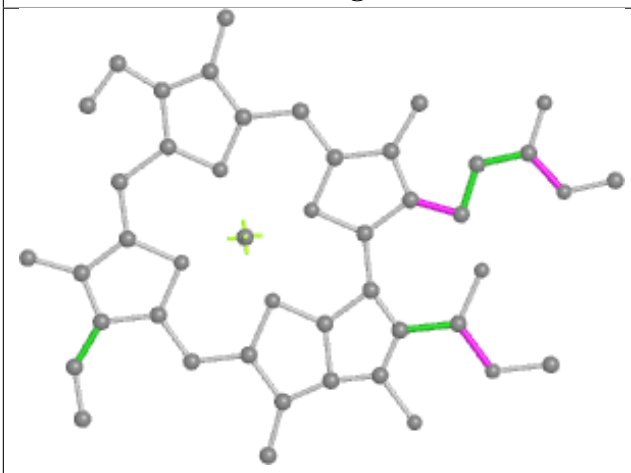
Ligand CLA 9 311



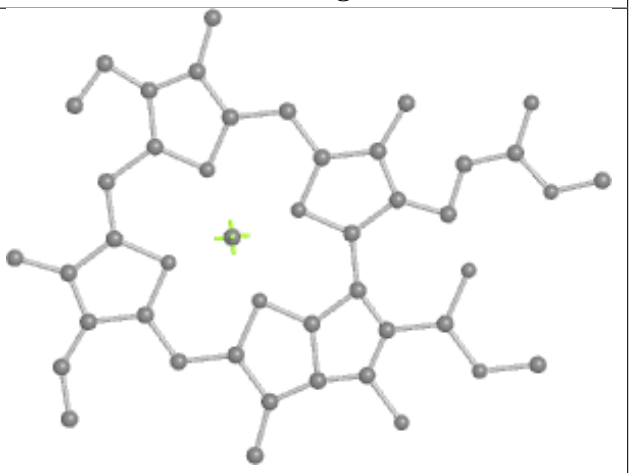
Bond lengths



Bond angles

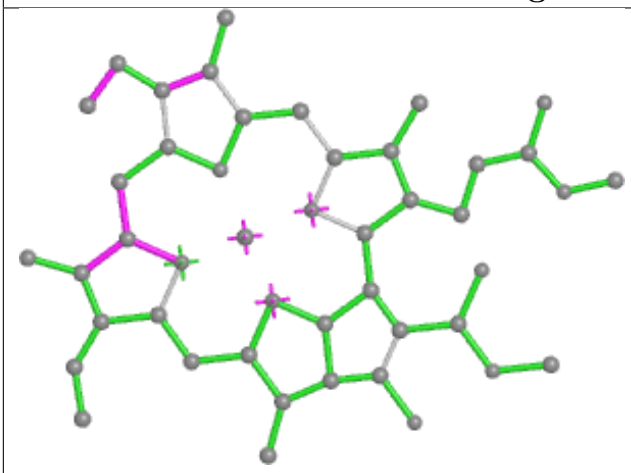


Torsions

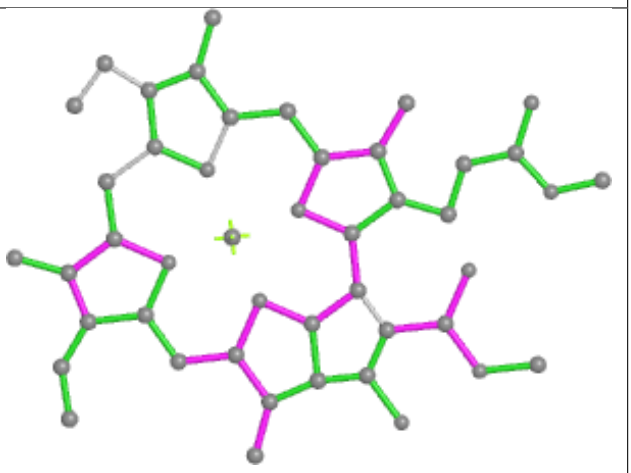


Rings

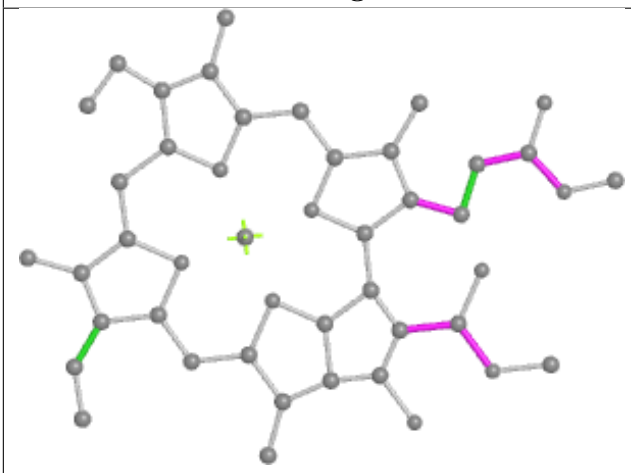
Ligand CLA 2 304



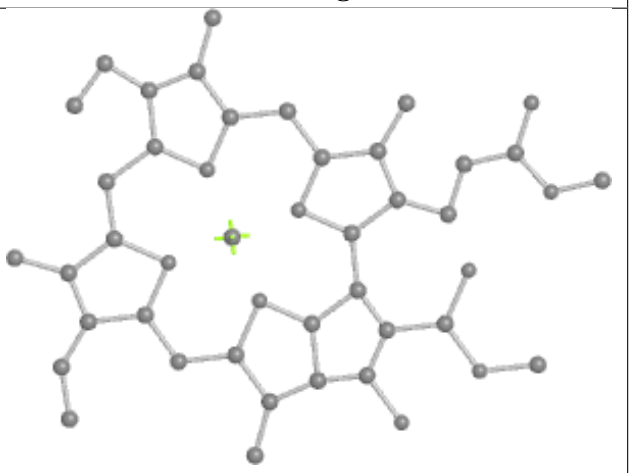
Bond lengths



Bond angles

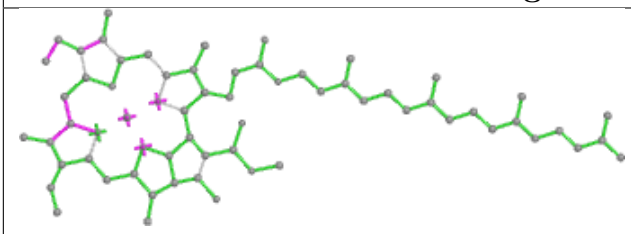


Torsions

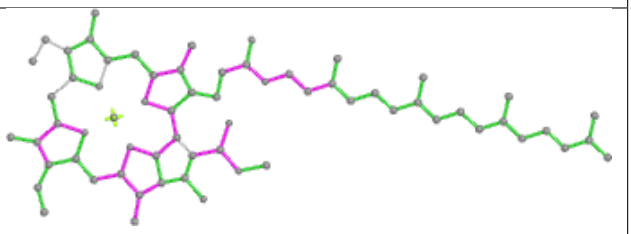


Rings

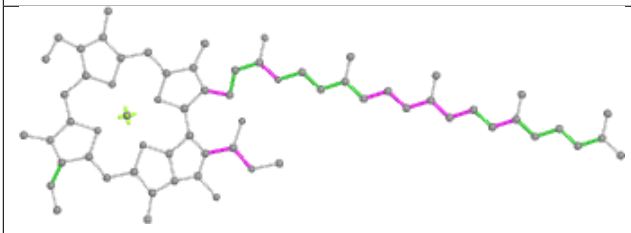
Ligand CLA A 830



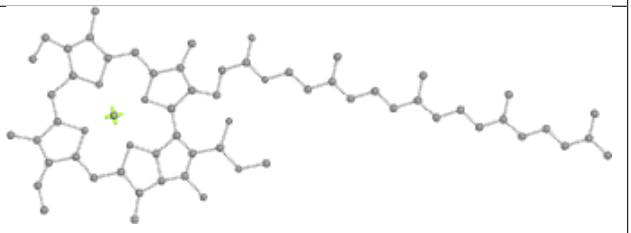
Bond lengths



Bond angles

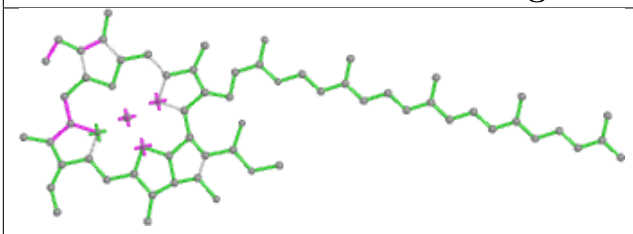


Torsions

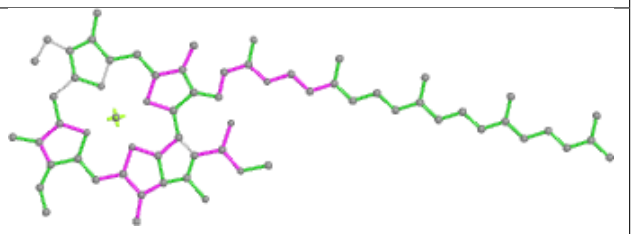


Rings

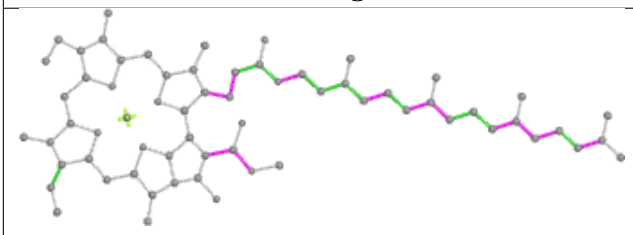
Ligand CLA 7 323



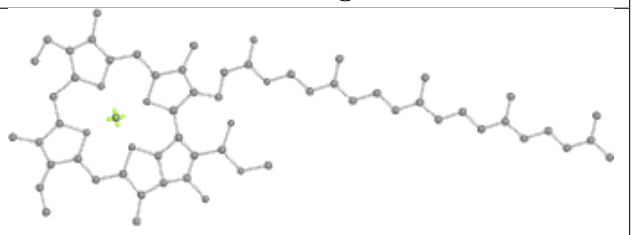
Bond lengths



Bond angles

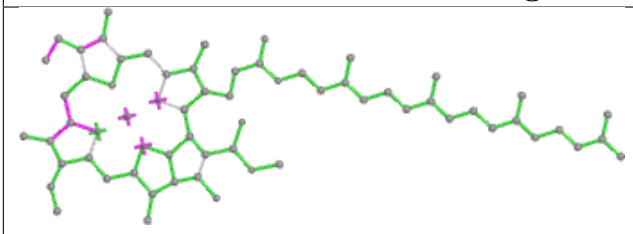


Torsions

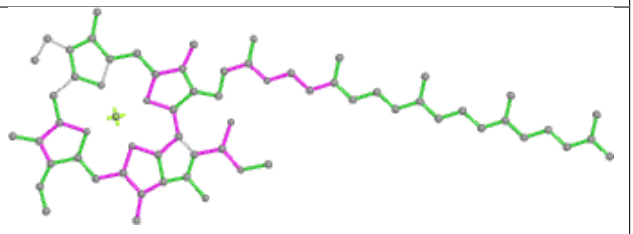


Rings

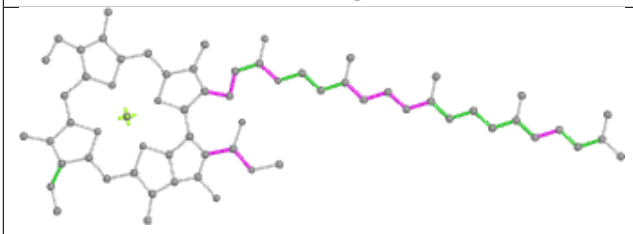
Ligand CLA A 834



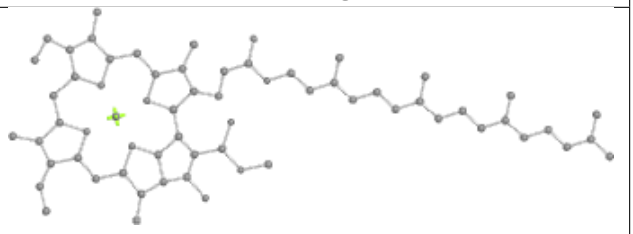
Bond lengths



Bond angles

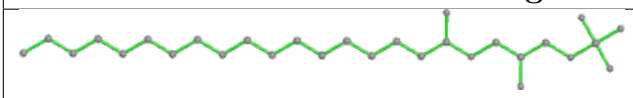


Torsions

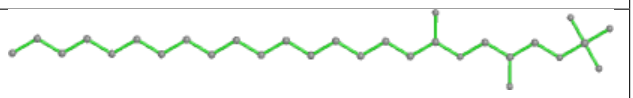


Rings

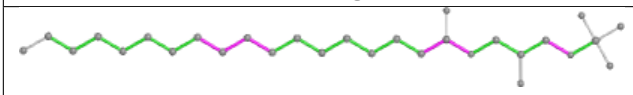
Ligand NKP 8 318



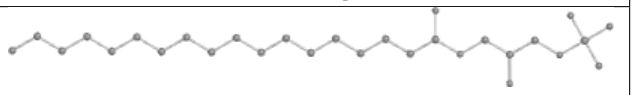
Bond lengths



Bond angles

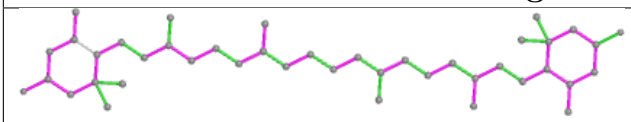


Torsions



Rings

Ligand LUT Z 302



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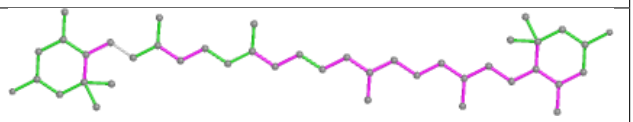
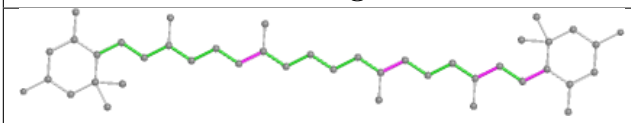
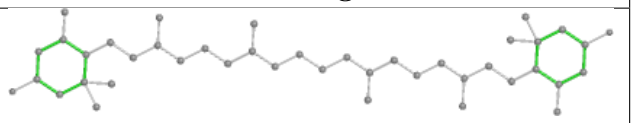


Bond angles

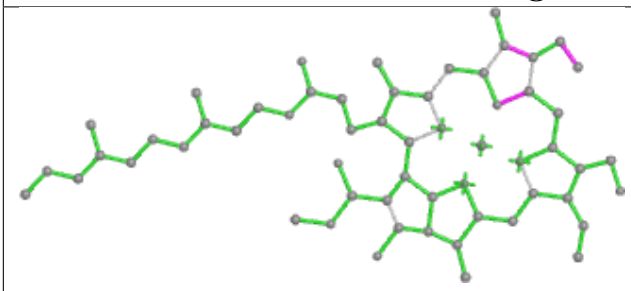


Torsions

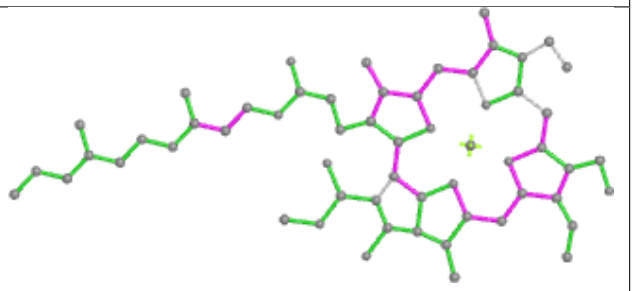


Rings

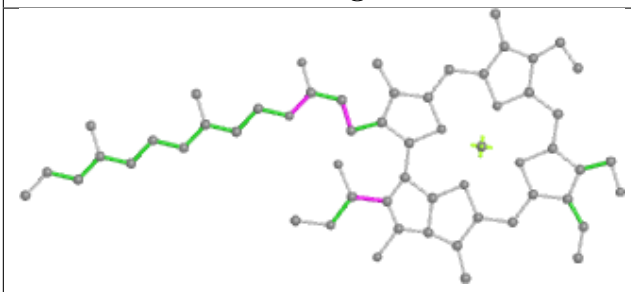
Ligand CHL 8 301



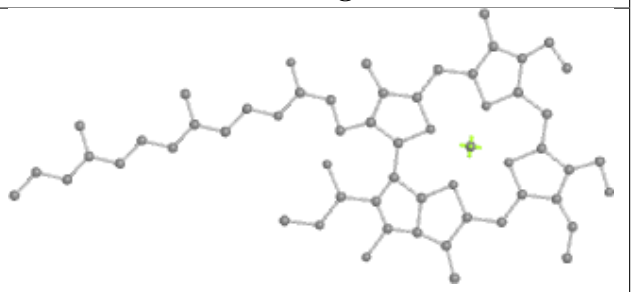
Bond lengths



Bond angles

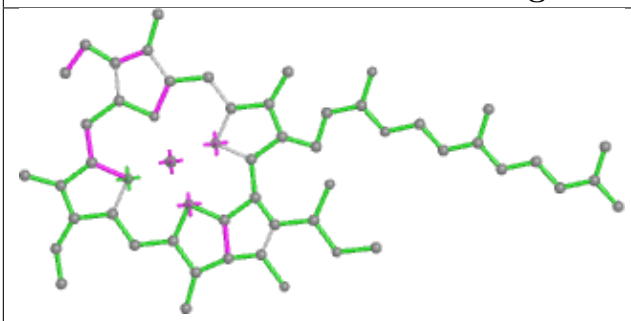


Torsions

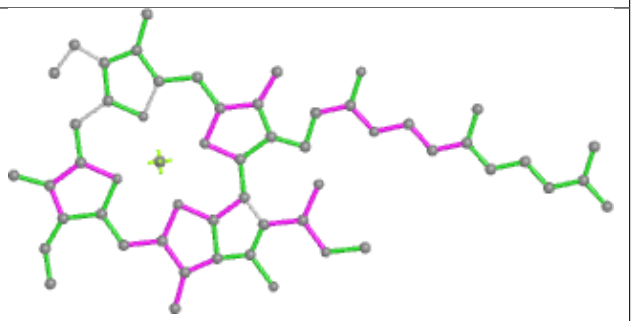


Rings

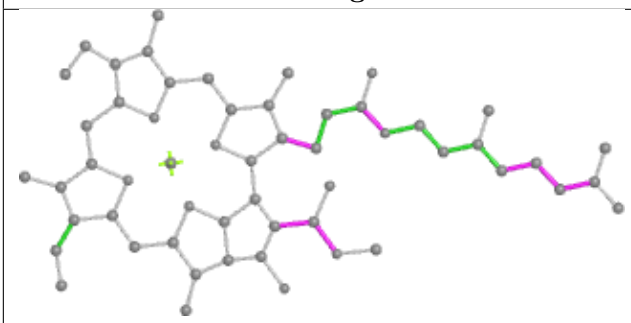
Ligand CLA 2 301



Bond lengths



Bond angles



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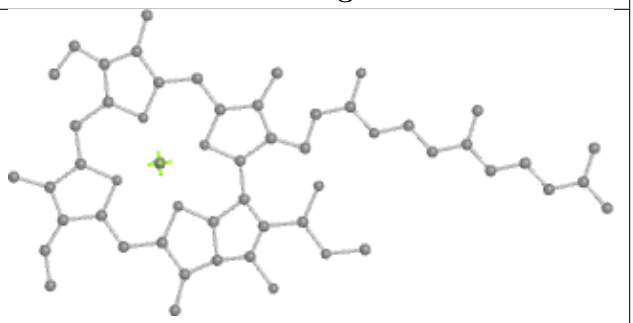
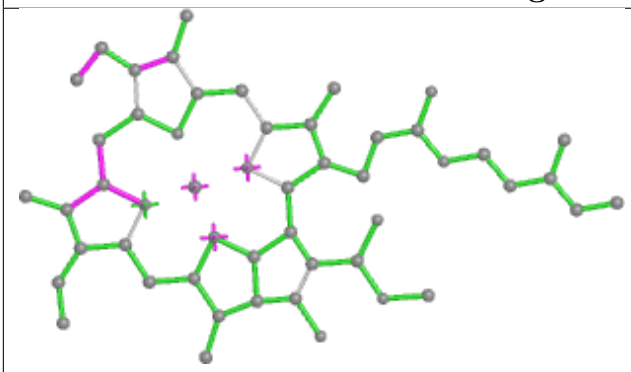
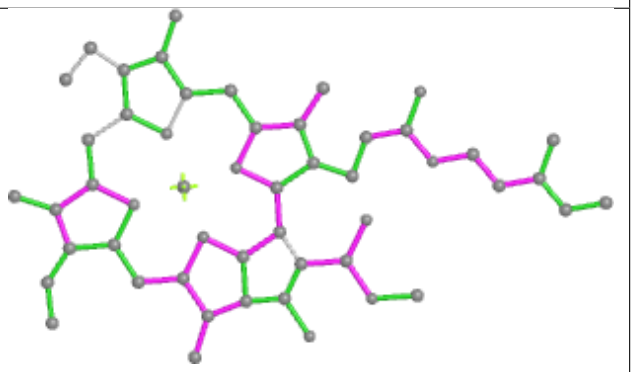


Rings

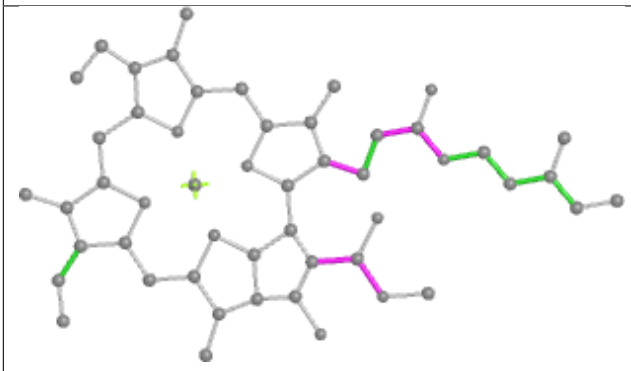
Ligand CLA 8 306



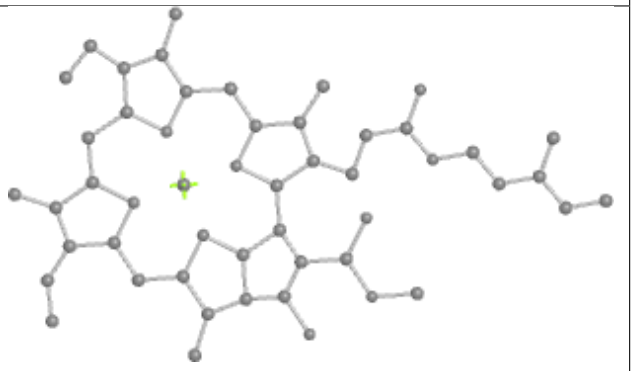
Bond lengths



Bond angles

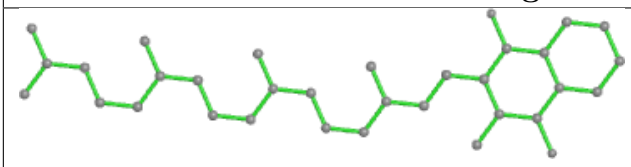


Torsions

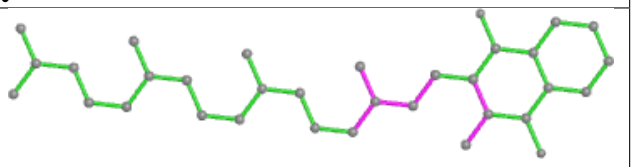


Rings

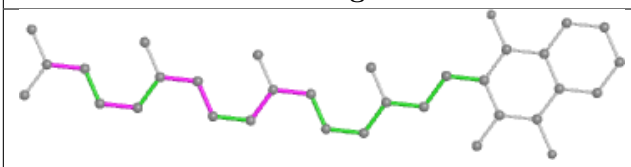
Ligand PQN B 843



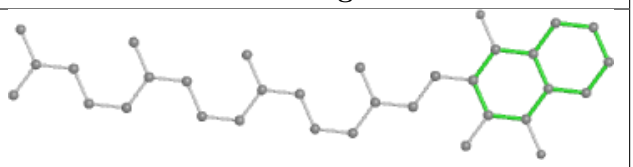
Bond lengths



Bond angles

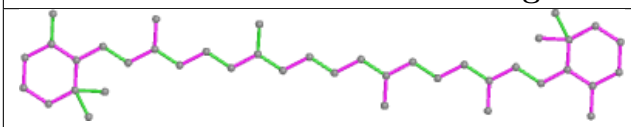


Torsions

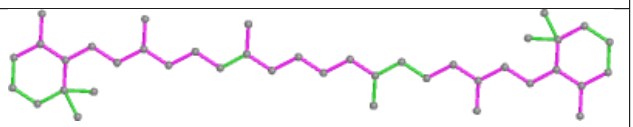


Rings

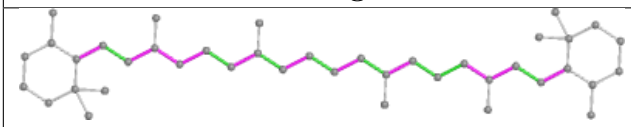
Ligand BCR B 846



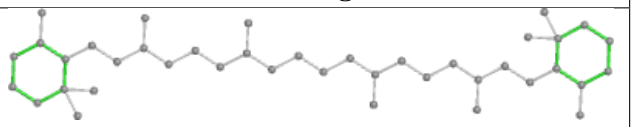
Bond lengths



Bond angles

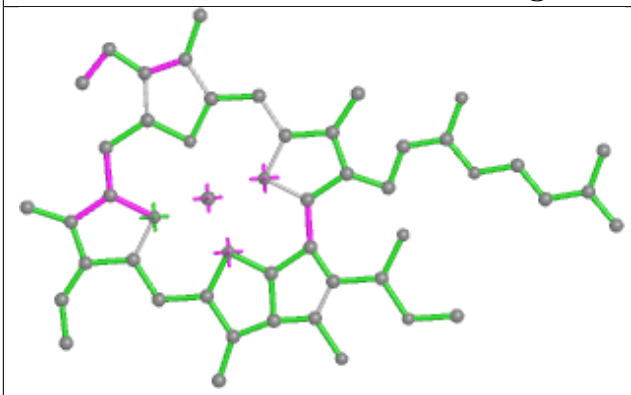


Torsions

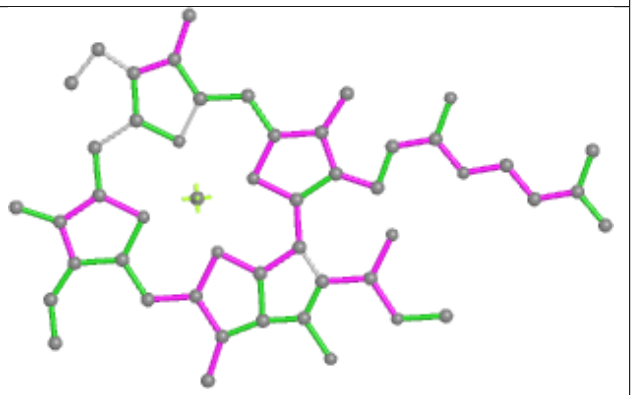


Rings

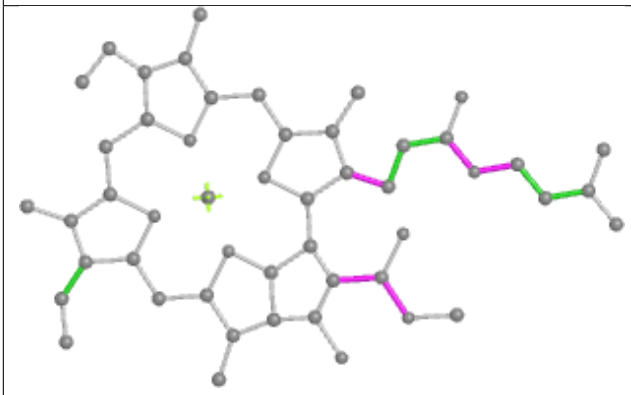
Ligand CLA 7 315



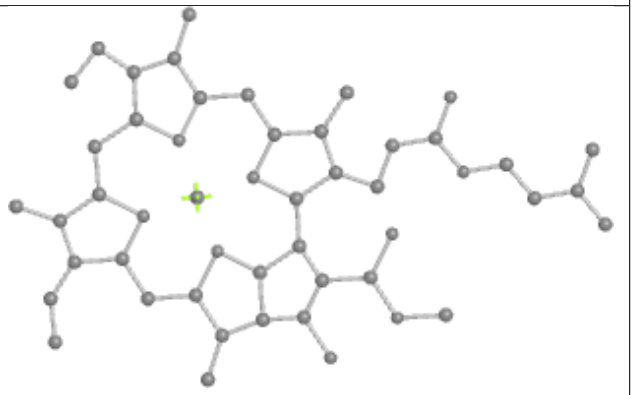
Bond lengths



Bond angles

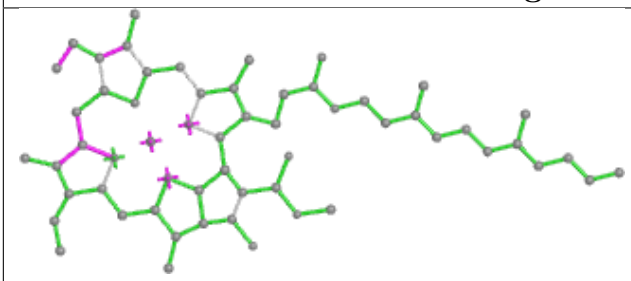


Torsions

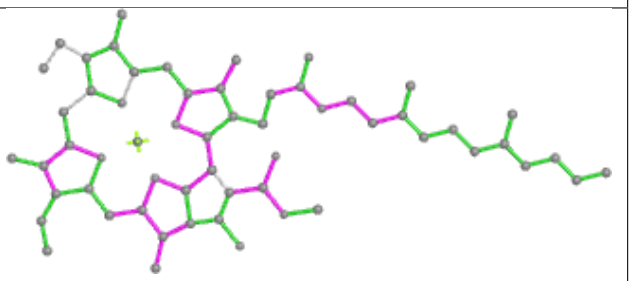


Rings

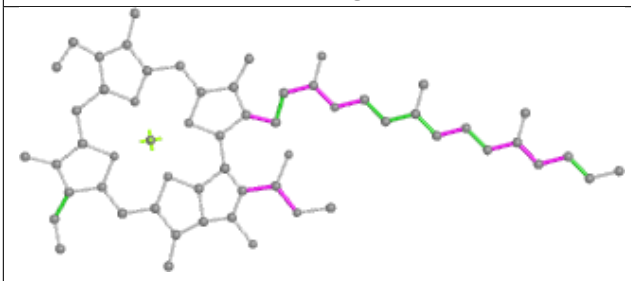
Ligand CLA B 834



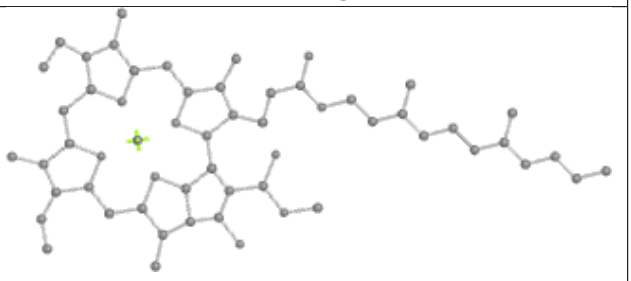
Bond lengths



Bond angles

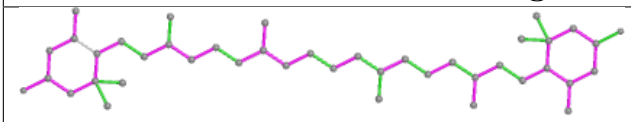


Torsions

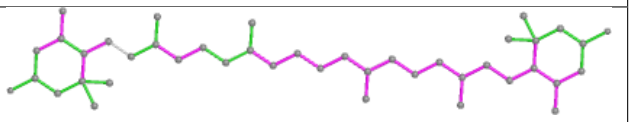


Rings

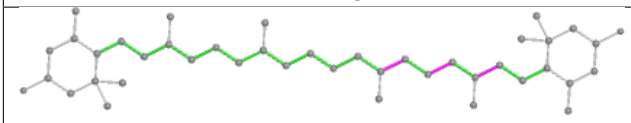
Ligand LUT 7 302



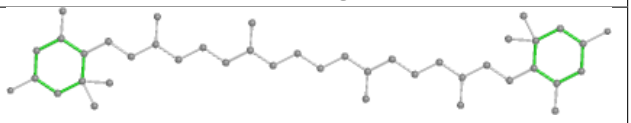
Bond lengths



Bond angles

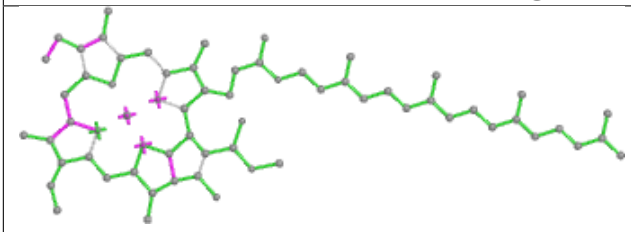


Torsions

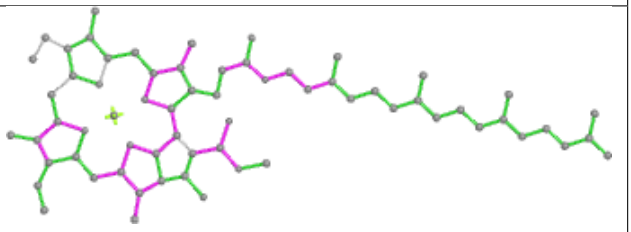


Rings

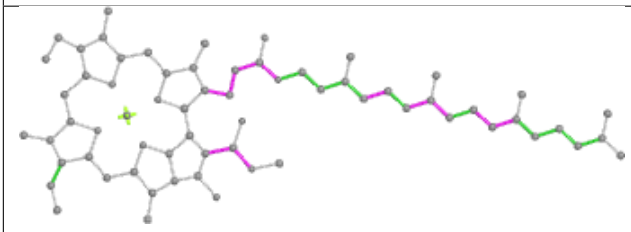
Ligand CLA 8 309



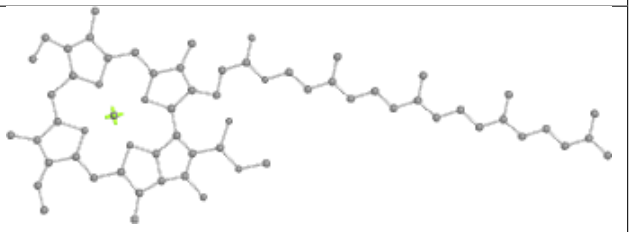
Bond lengths



Bond angles

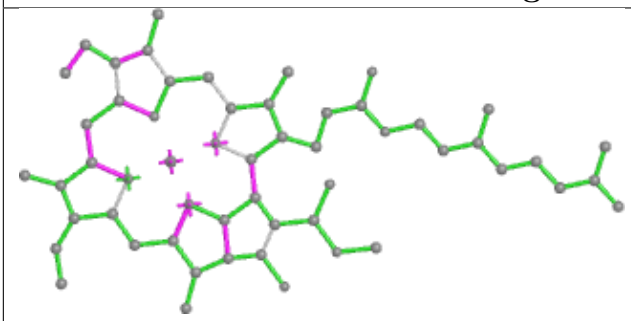


Torsions

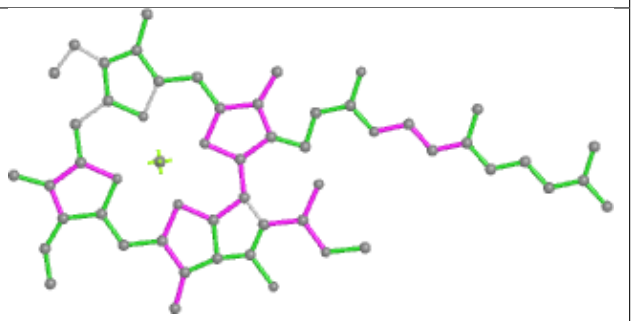


Rings

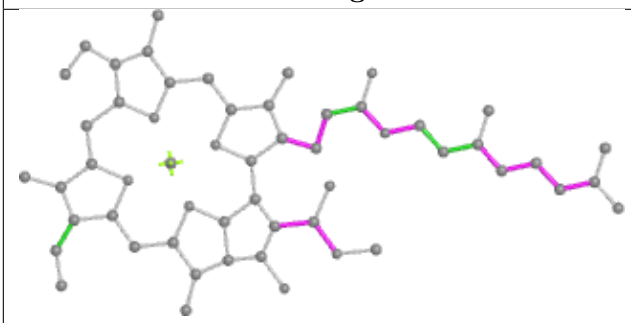
Ligand CLA K 205



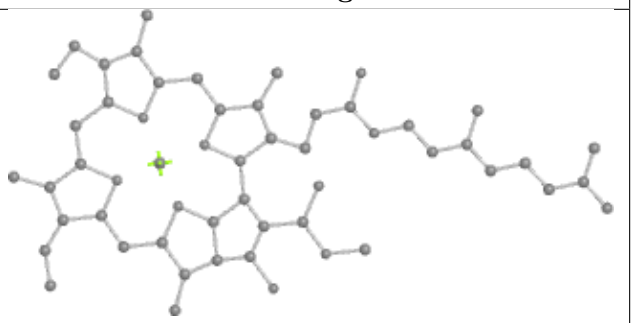
Bond lengths



Bond angles

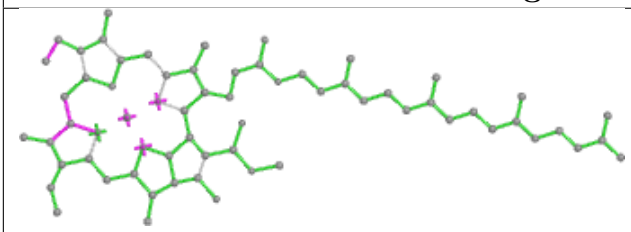


Torsions

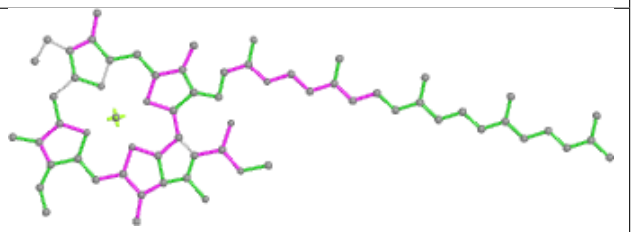


Rings

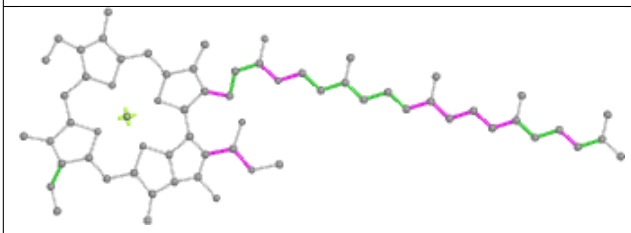
Ligand CLA A 832



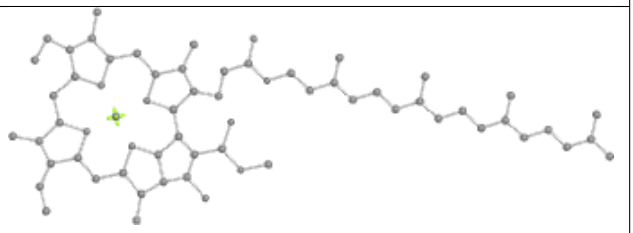
Bond lengths



Bond angles

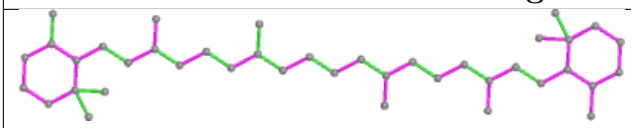


Torsions

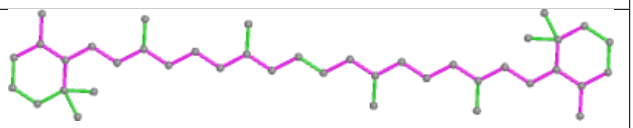


Rings

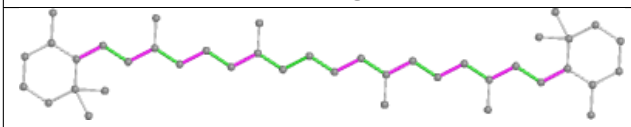
Ligand BCR 3 305



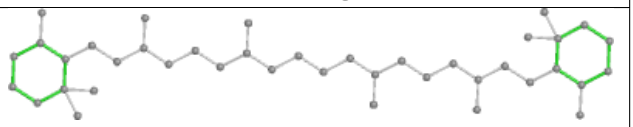
Bond lengths



Bond angles

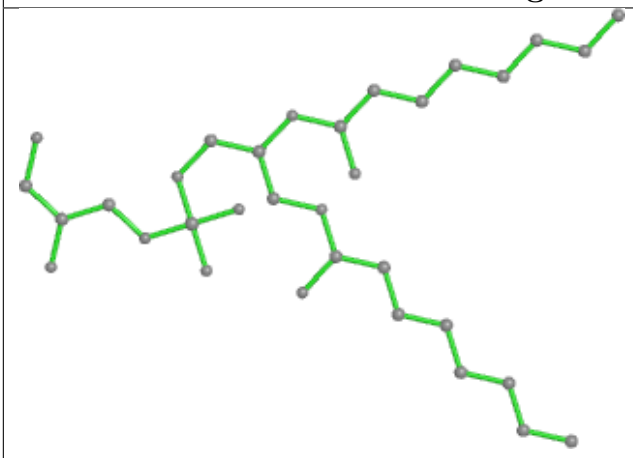


Torsions

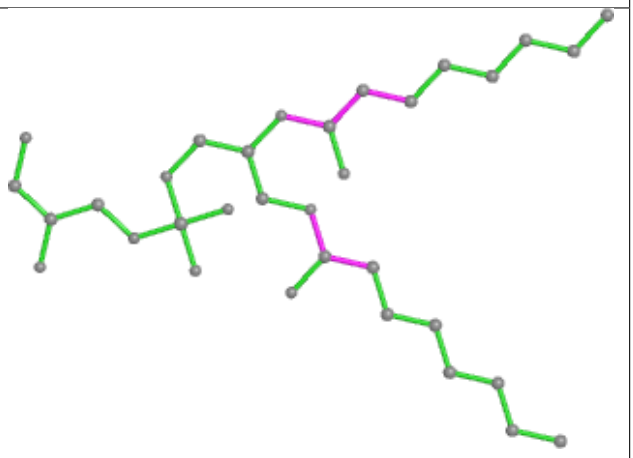


Rings

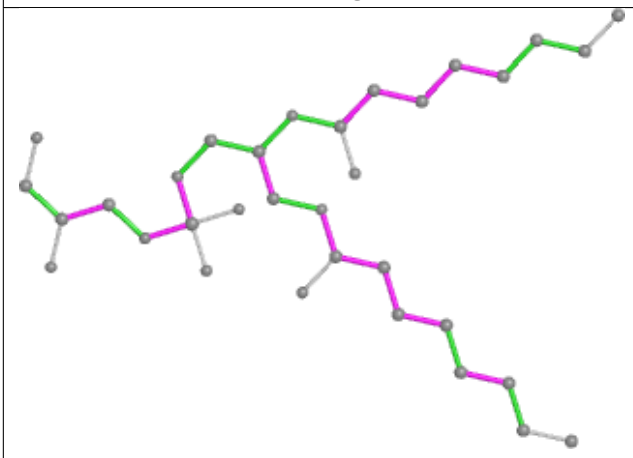
Ligand LHG 9 315



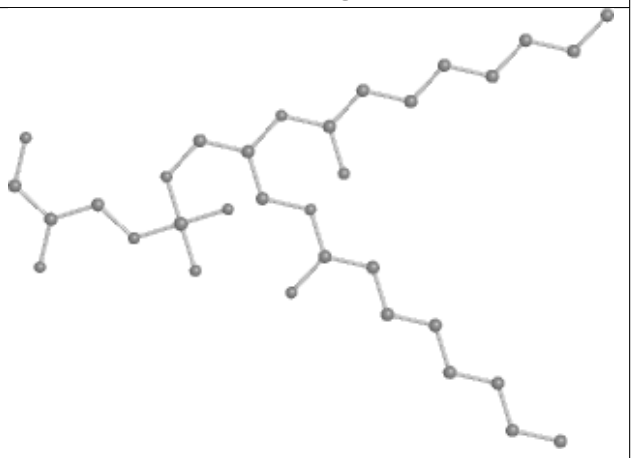
Bond lengths



Bond angles

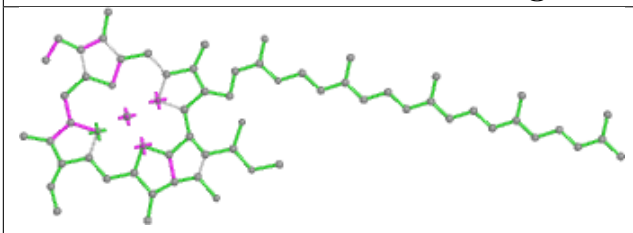


Torsions

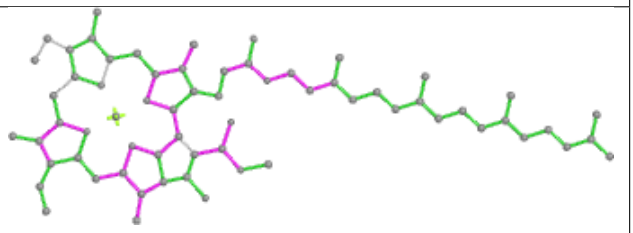


Rings

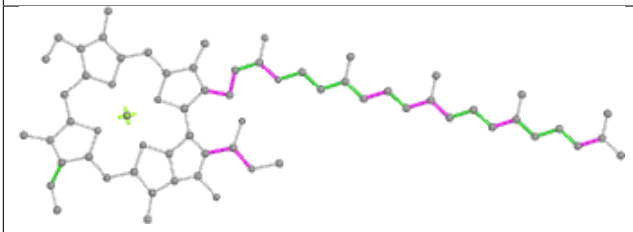
Ligand CLA 6 310



Bond lengths



Bond angles



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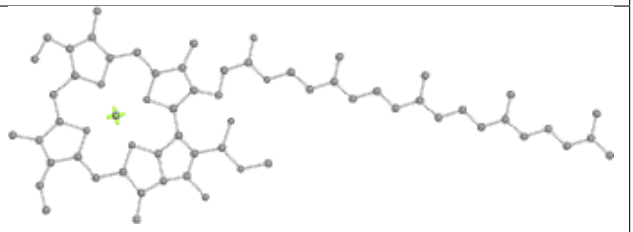
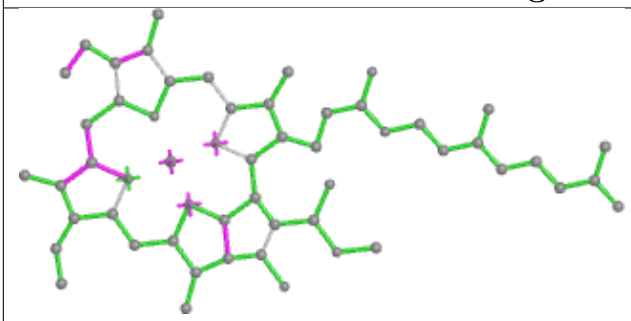


Rings

Ligand CLA 1 308



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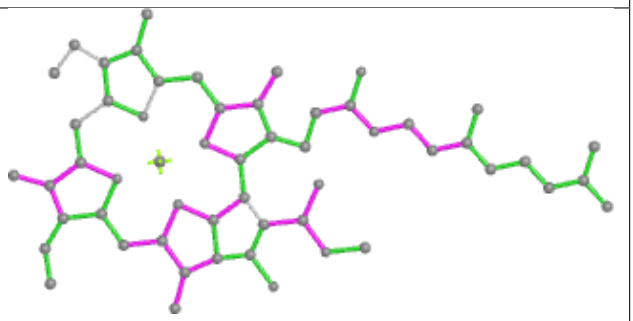
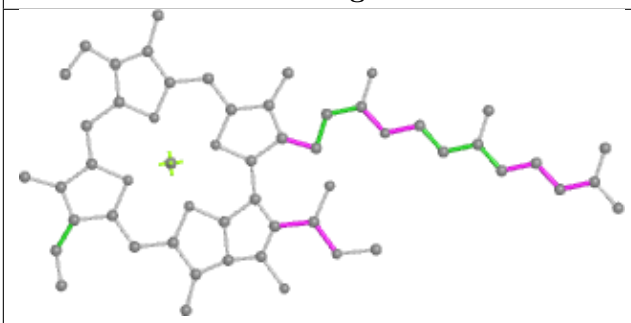


Bond angles



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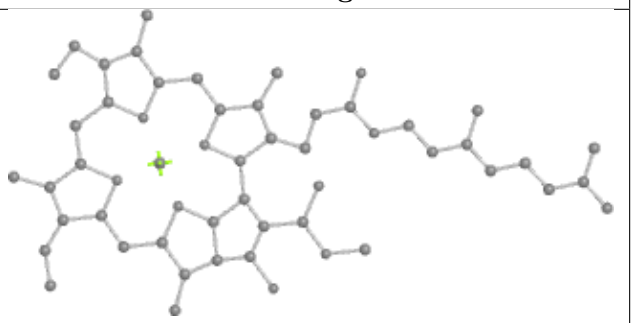
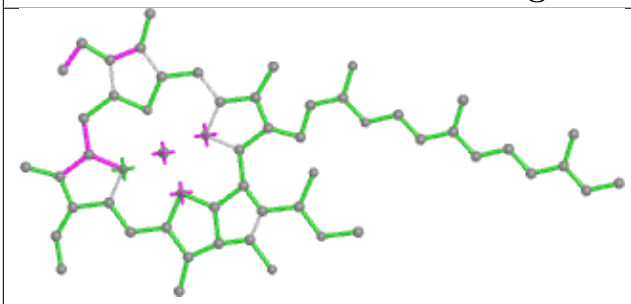
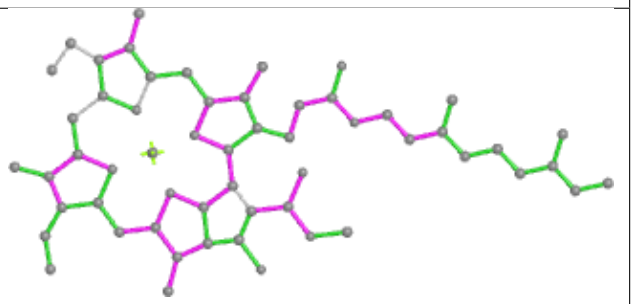


Rings

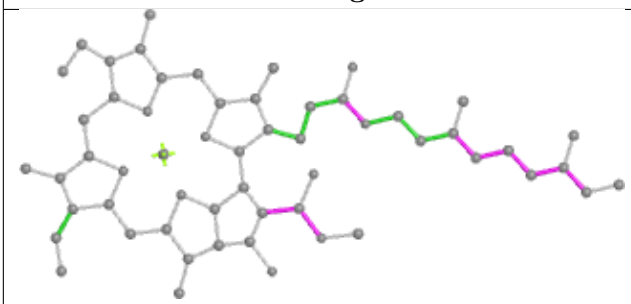
Ligand CLA 7 309



Bond lengths



Bond angles



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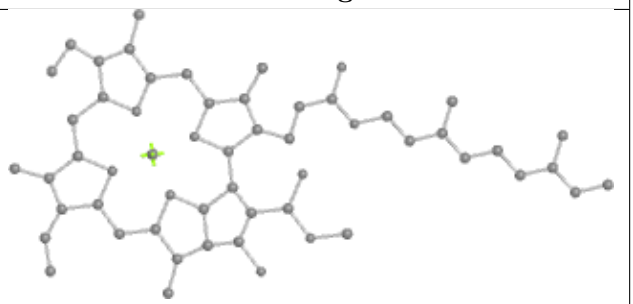
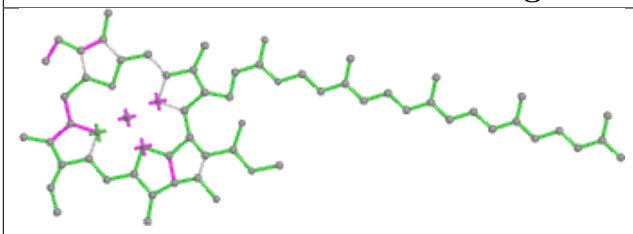
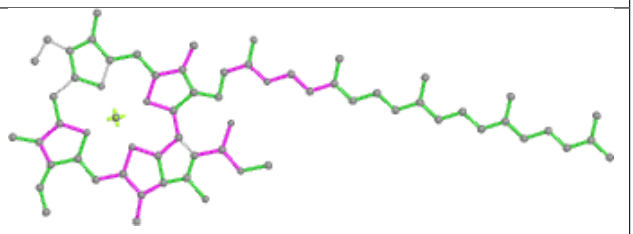


Rings

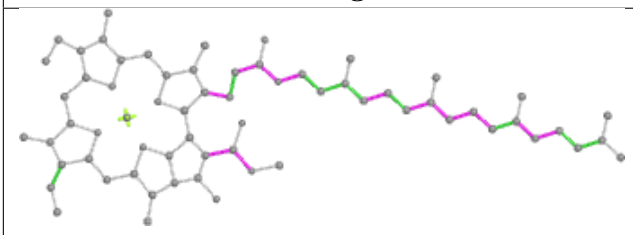
Ligand CLA A 811



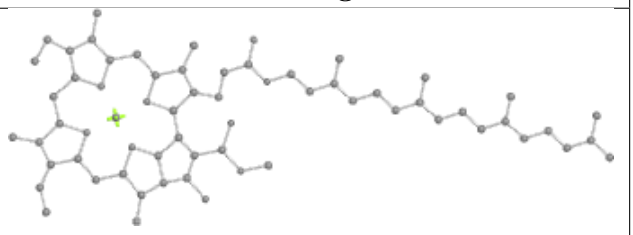
Bond lengths



Bond angles

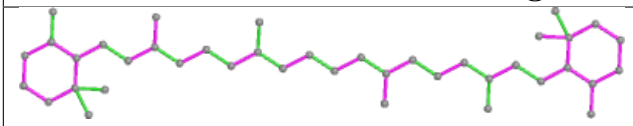


Torsions

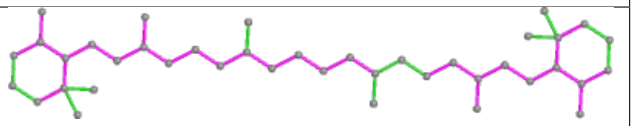


Rings

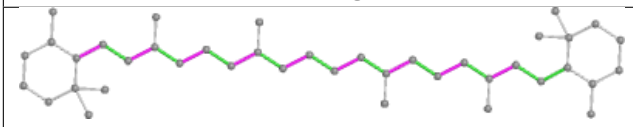
Ligand BCR 6 305



Bond lengths



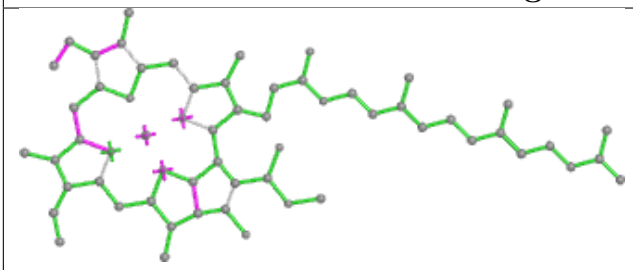
Bond angles



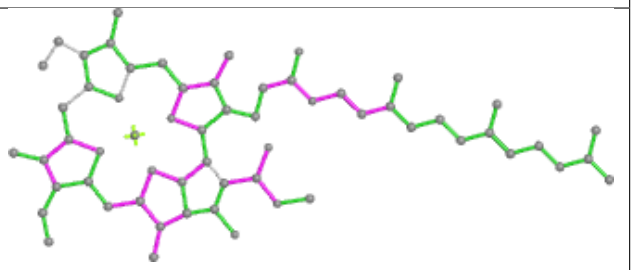
Torsions

Rings

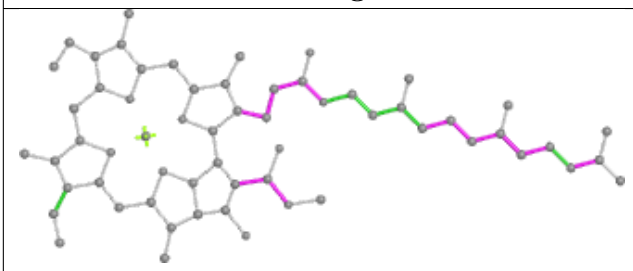
Ligand CLA B 814



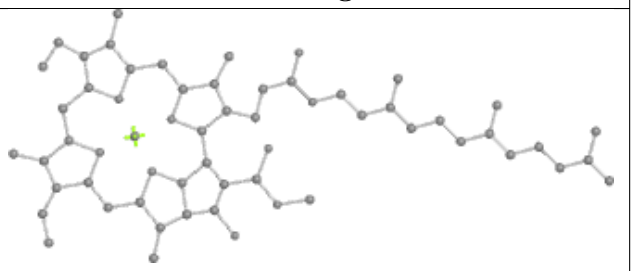
Bond lengths



Bond angles

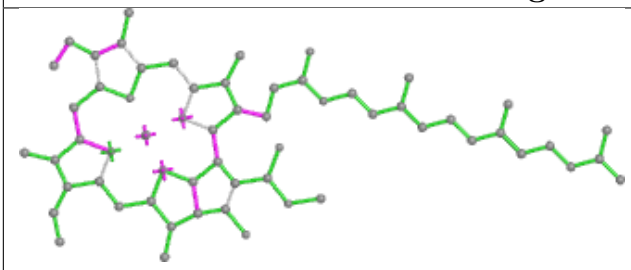


Torsions

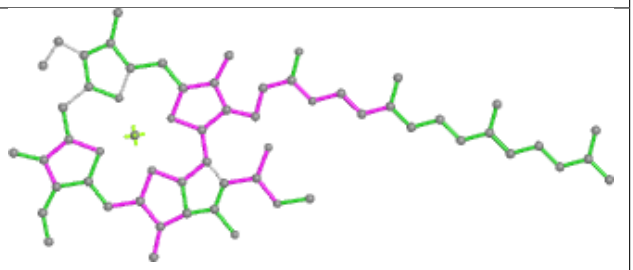


Rings

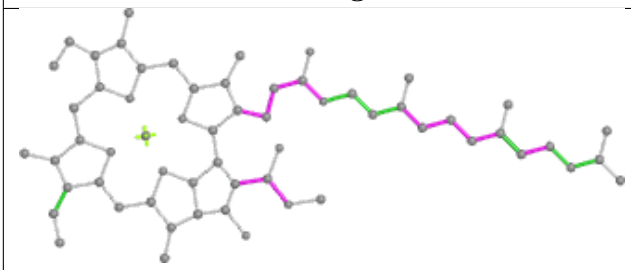
Ligand CLA 9 305



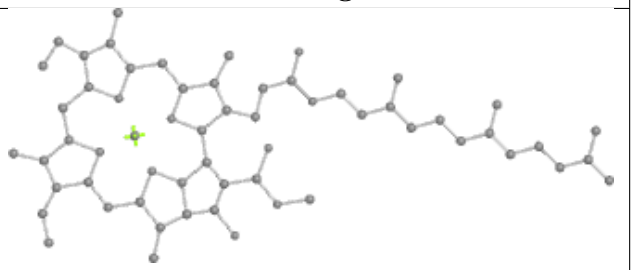
Bond lengths



Bond angles

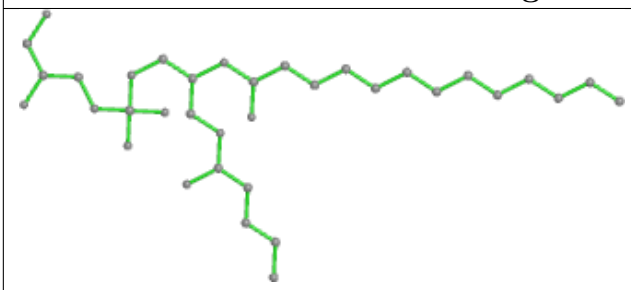


Torsions

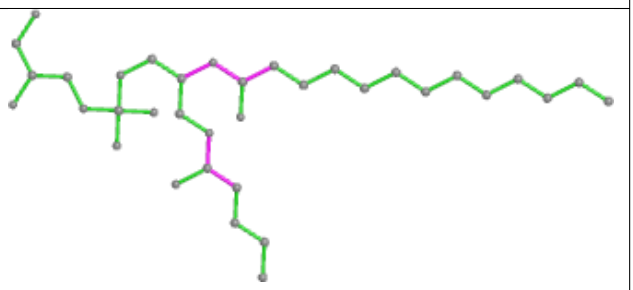


Rings

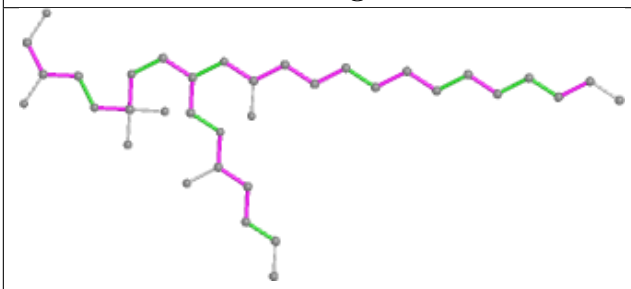
Ligand LHG A 849



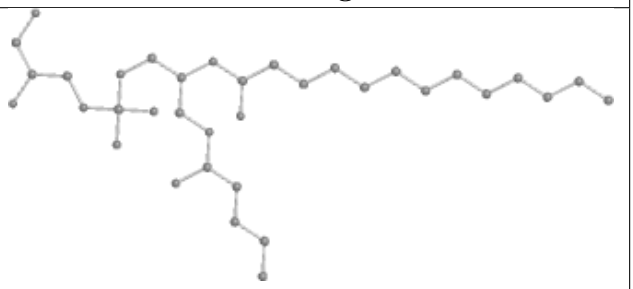
Bond lengths



Bond angles

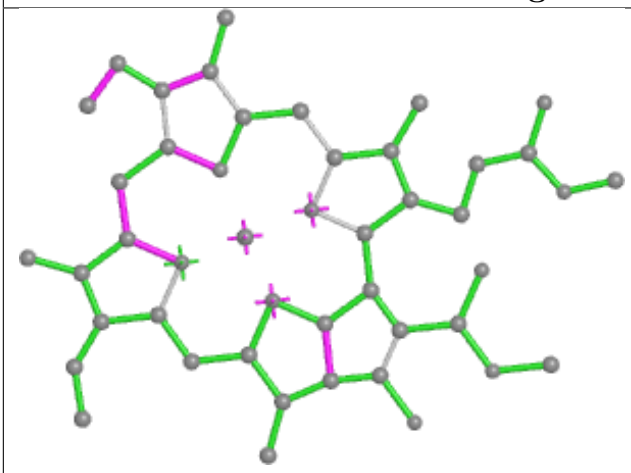


Torsions

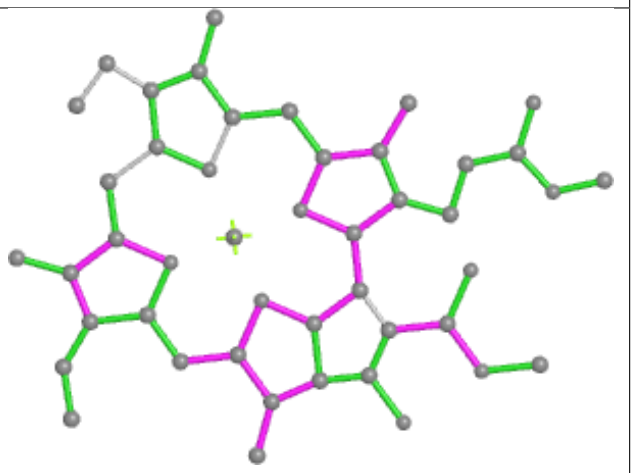


Rings

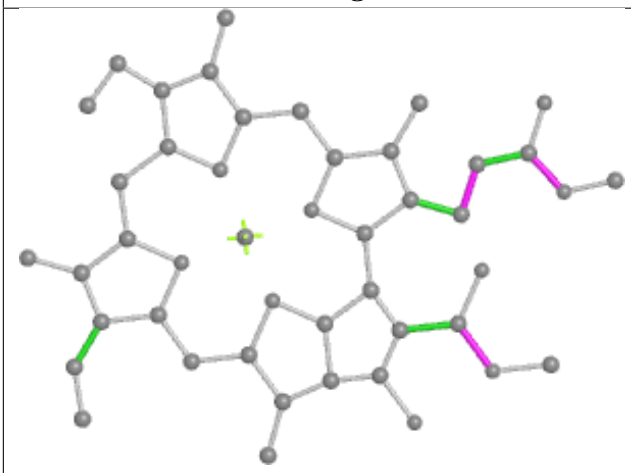
Ligand CLA 3 320



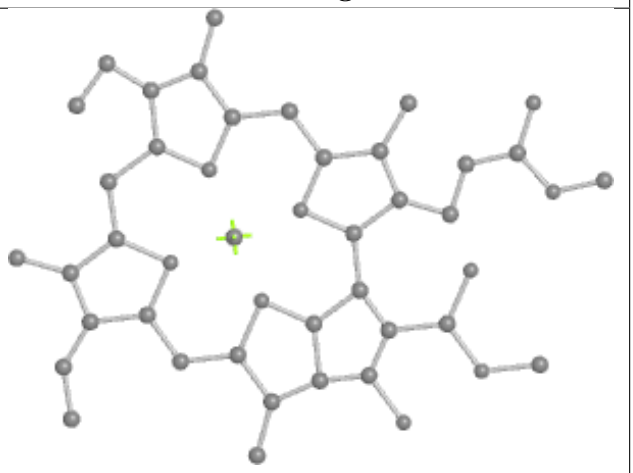
Bond lengths



Bond angles

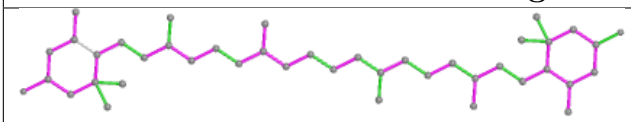


Torsions

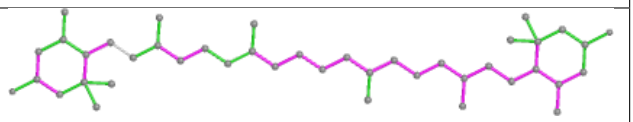


Rings

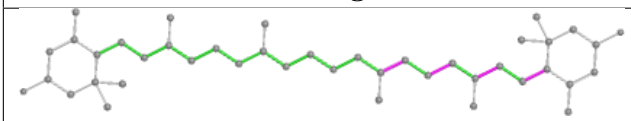
Ligand LUT 5 303



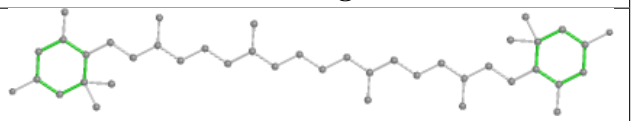
Bond lengths



Bond angles

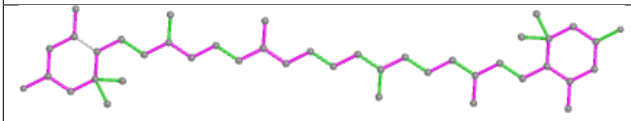


Torsions

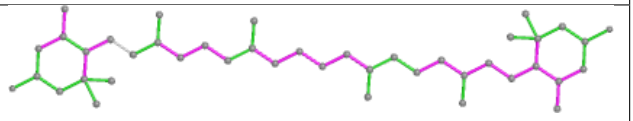


Rings

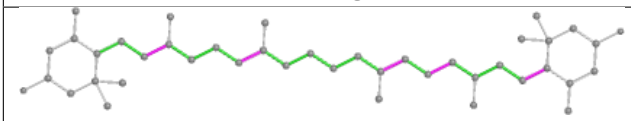
Ligand LUT 8 302



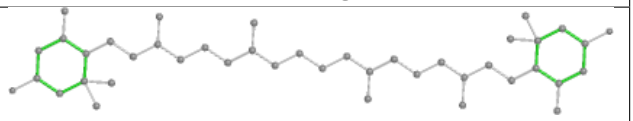
Bond lengths



Bond angles

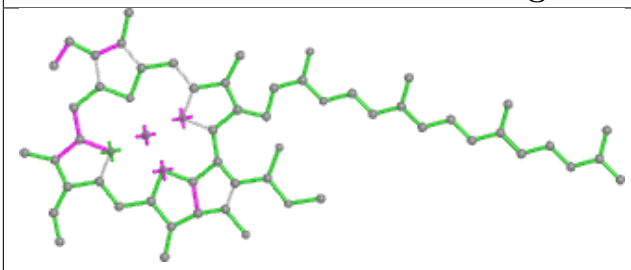


Torsions

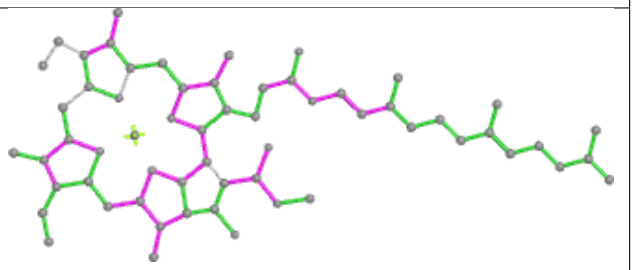


Rings

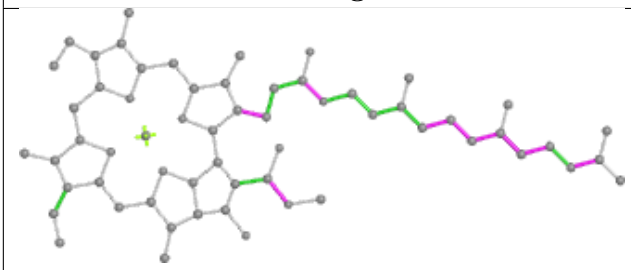
Ligand CLA 5 307



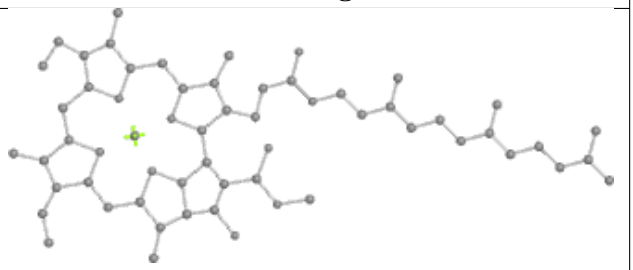
Bond lengths



Bond angles

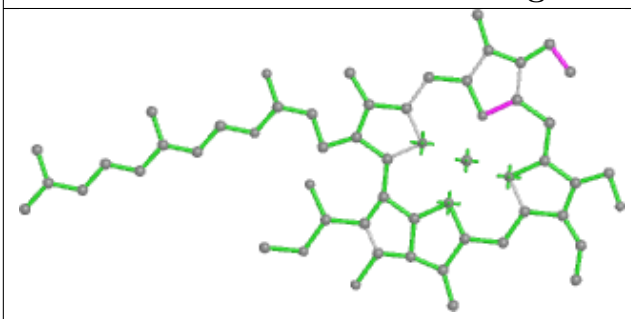


Torsions

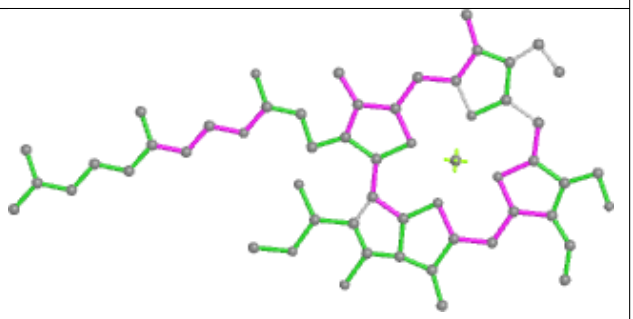


Rings

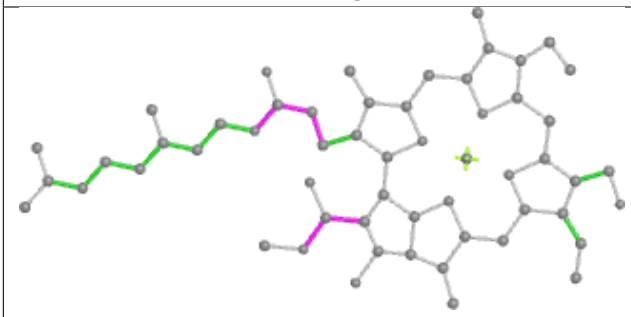
Ligand CHL 8 312



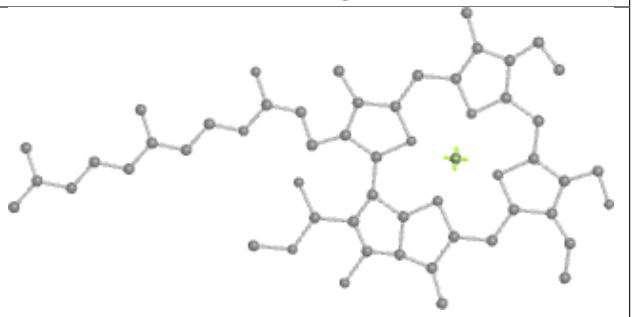
Bond lengths



Bond angles

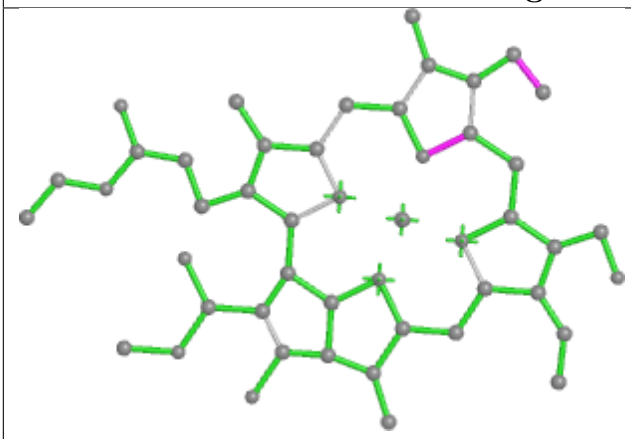


Torsions



Rings

Ligand CHL Z 312



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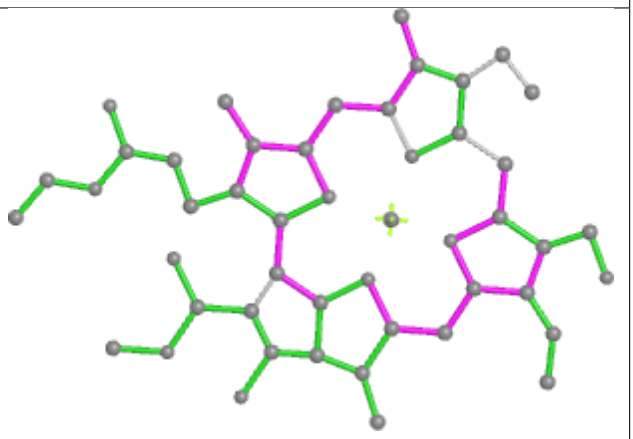
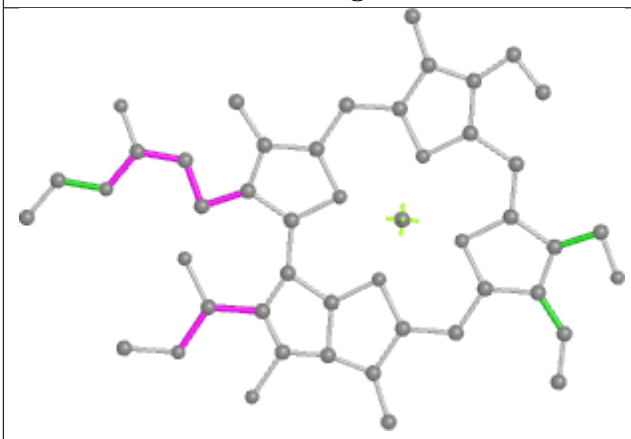
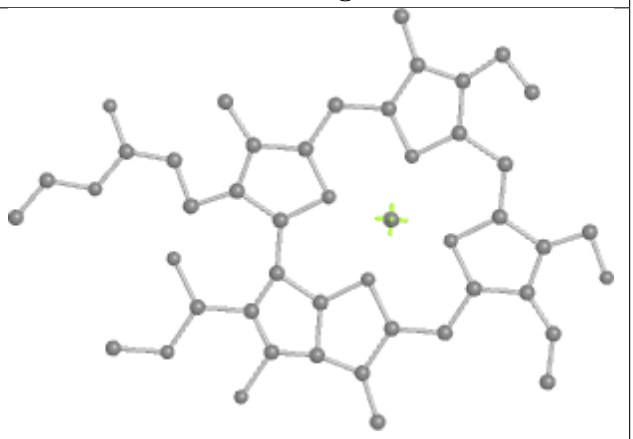


Bond angles

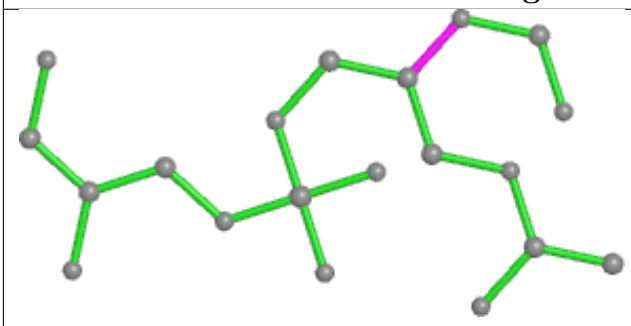


Torsions

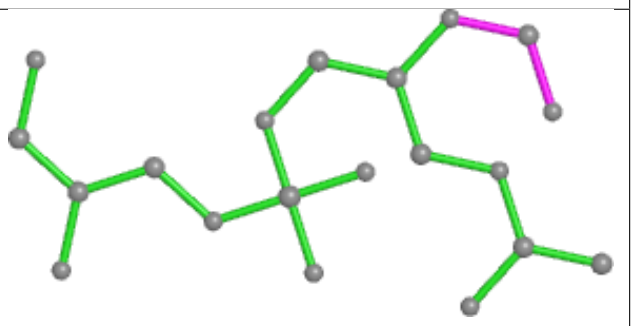


Rings

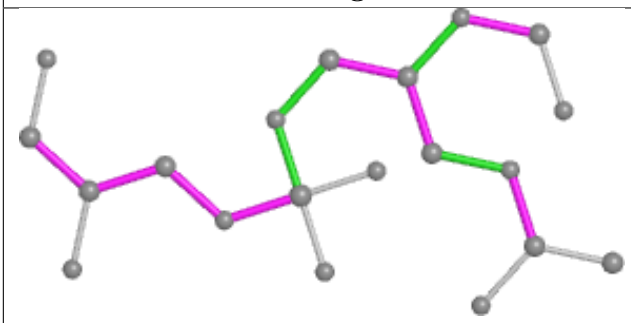
Ligand LHG B 851



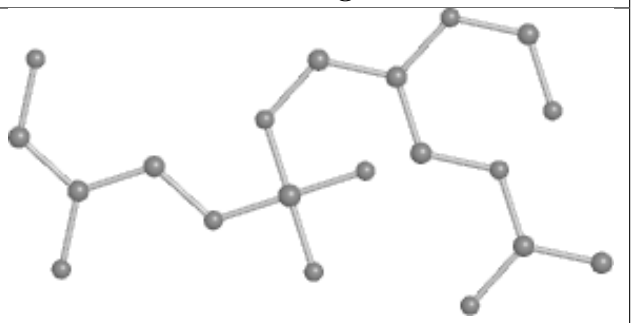
Bond lengths



Bond angles

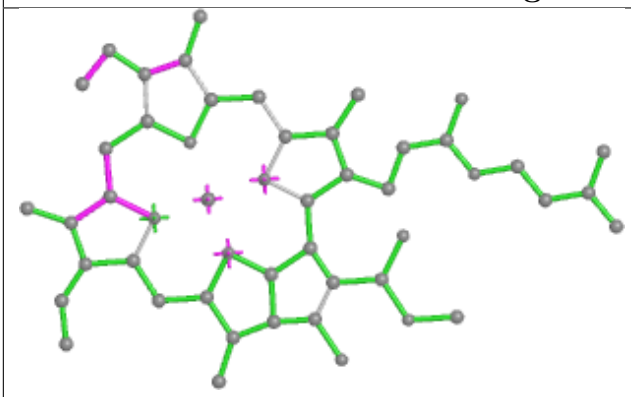


Torsions

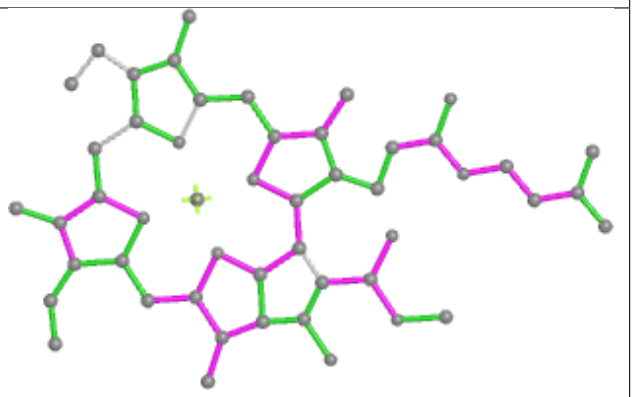


Rings

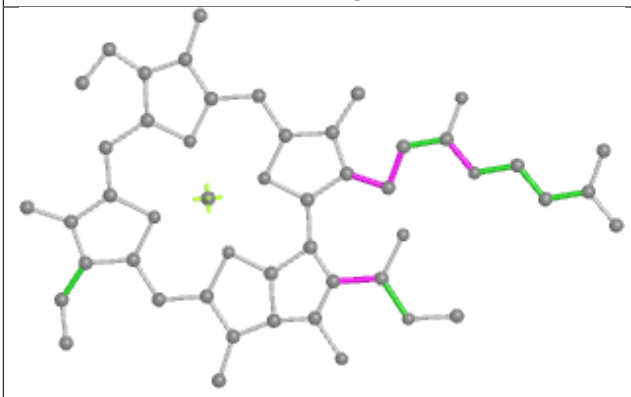
Ligand CLA A 826



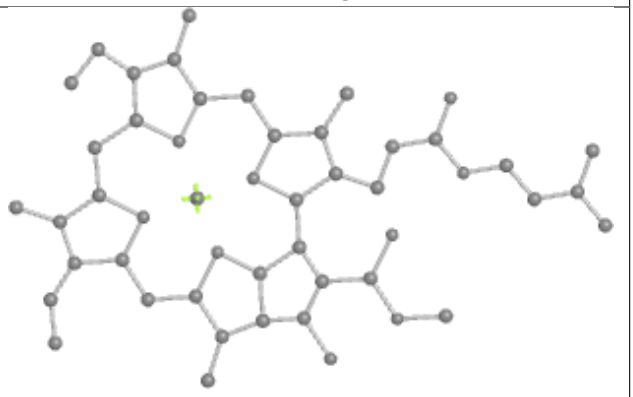
Bond lengths



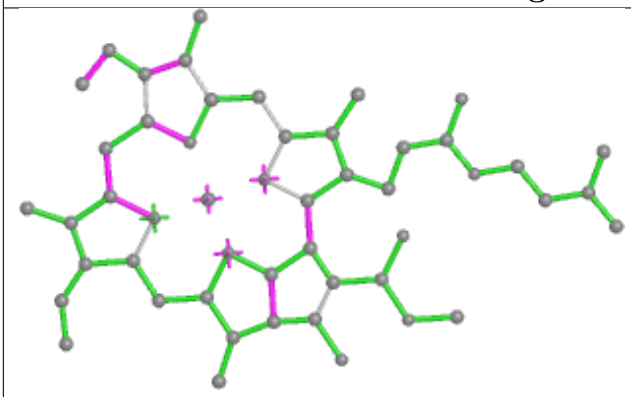
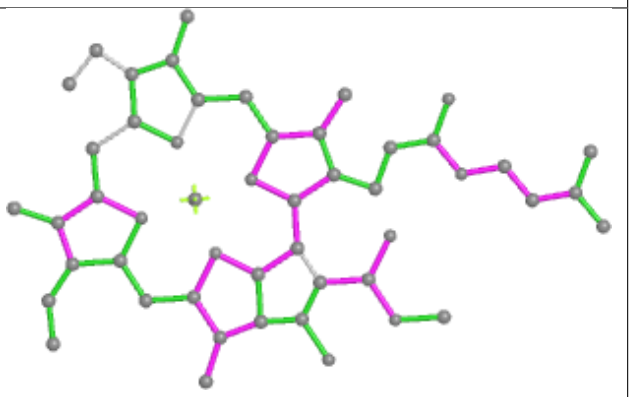
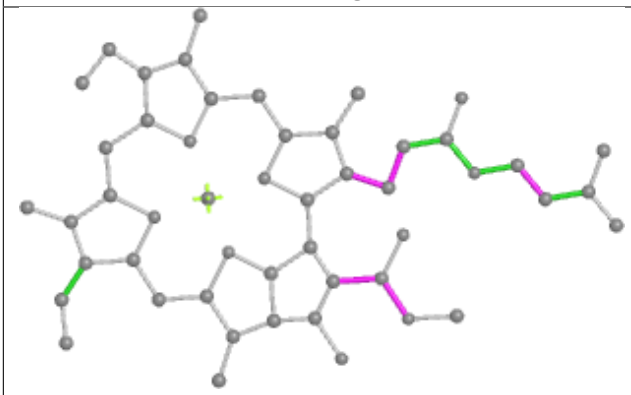
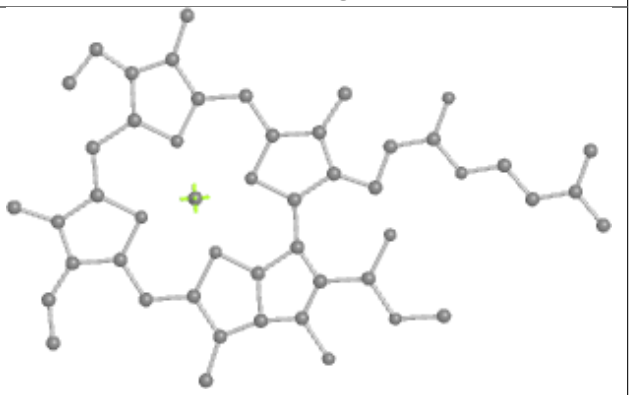
Bond angles

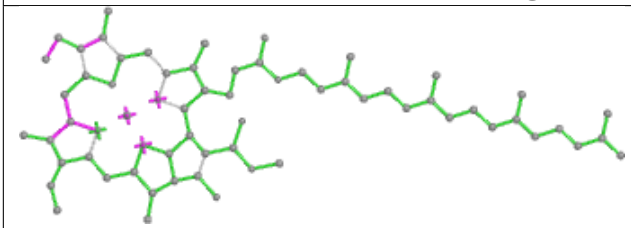
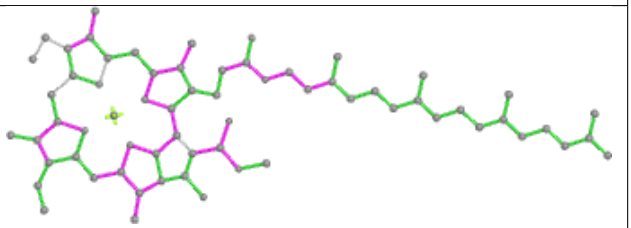
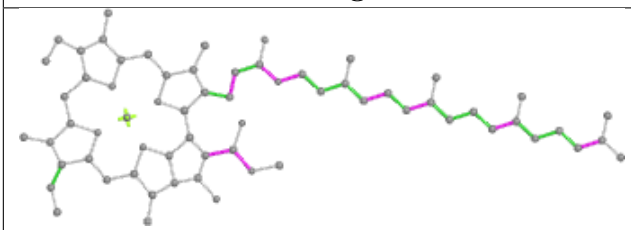
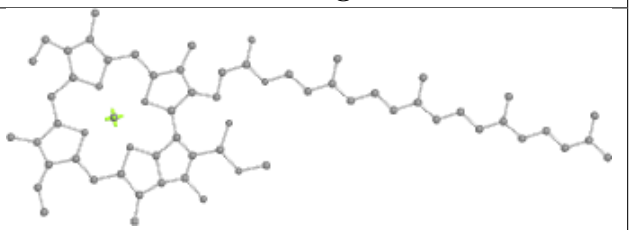


Torsions

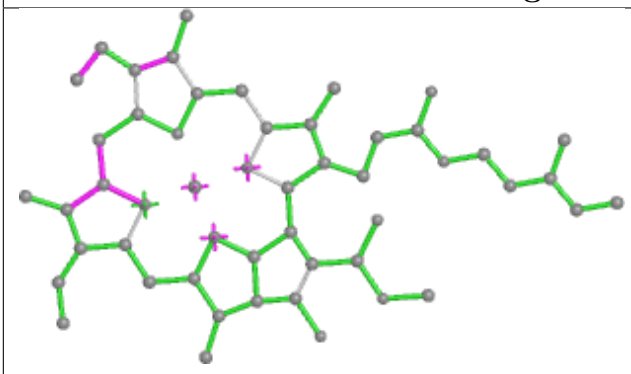


Rings

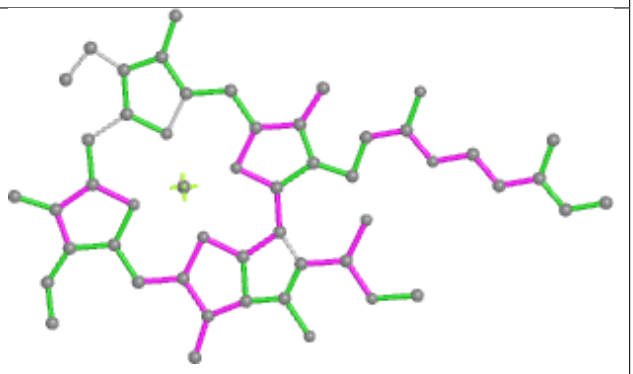
Ligand CLA 5 320	
 Bond lengths	 Bond angles
 Torsions	 Rings

Ligand CLA A 821	
 Bond lengths	 Bond angles
 Torsions	 Rings

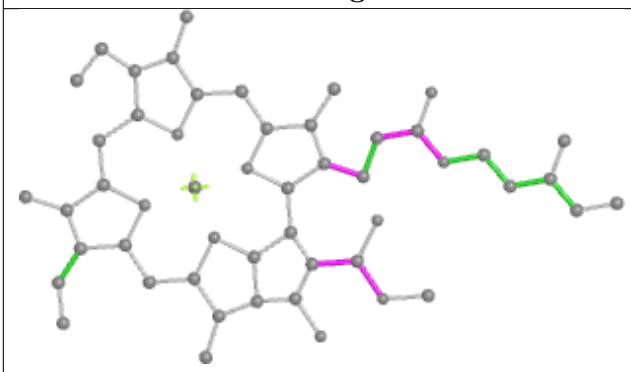
Ligand CLA A 837



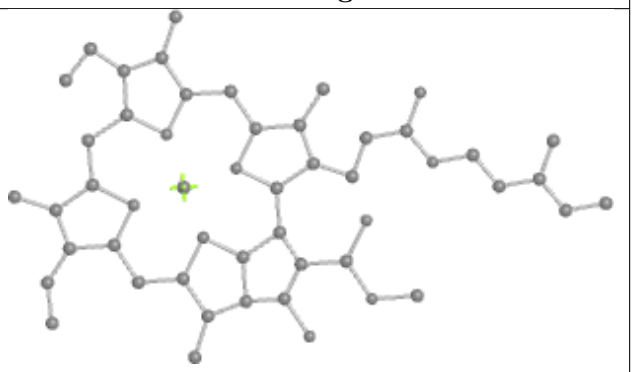
Bond lengths



Bond angles

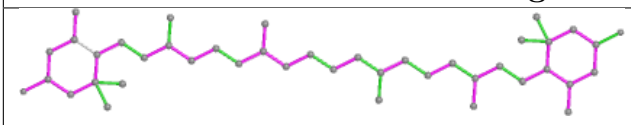


Torsions

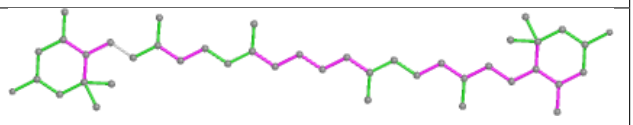


Rings

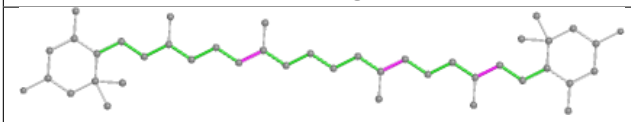
Ligand LUT 9 301



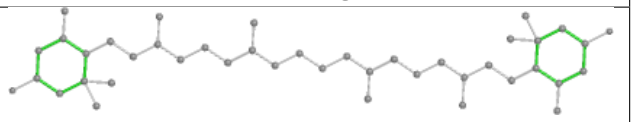
Bond lengths



Bond angles

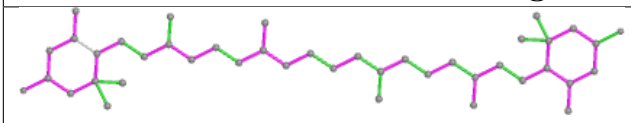


Torsions

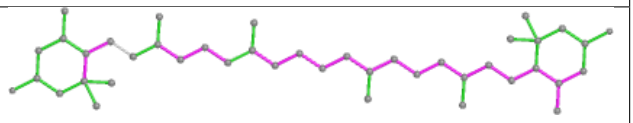


Rings

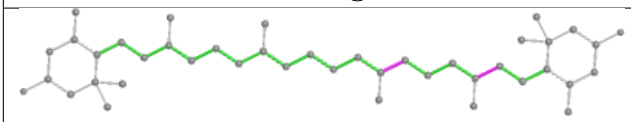
Ligand LUT 9 302



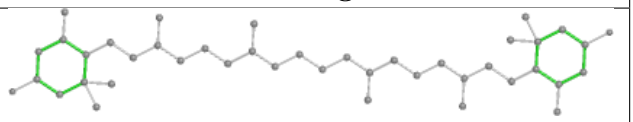
Bond lengths



Bond angles

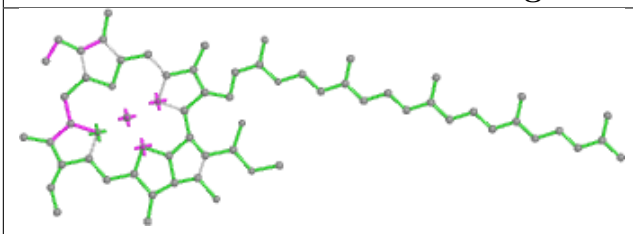


Torsions

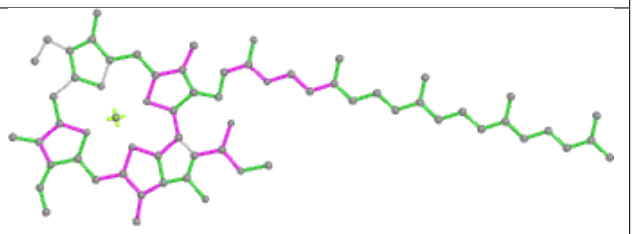


Rings

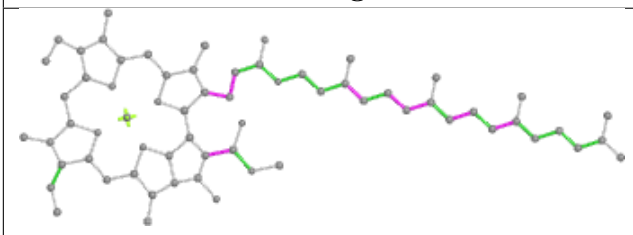
Ligand CLA B 830



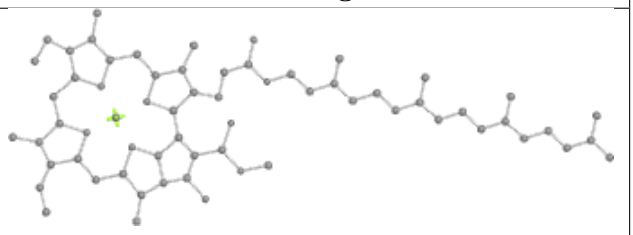
Bond lengths



Bond angles

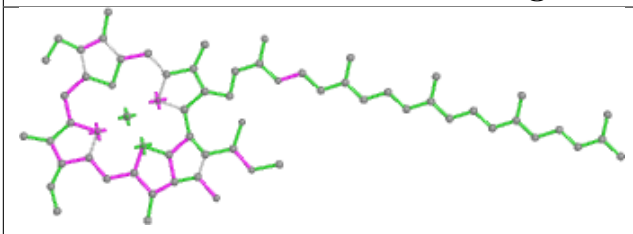


Torsions

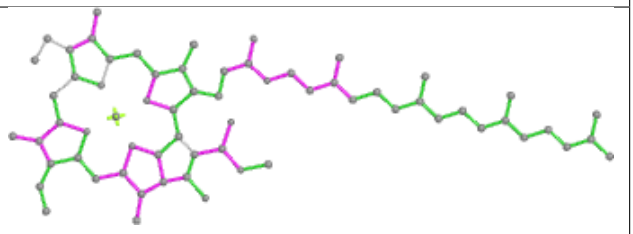


Rings

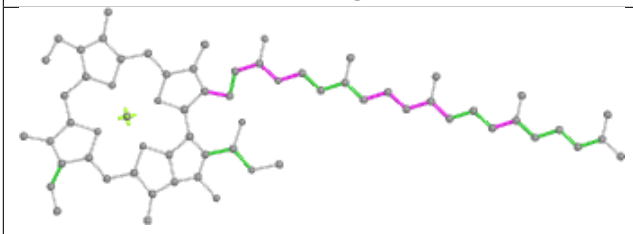
Ligand CL0 A 801



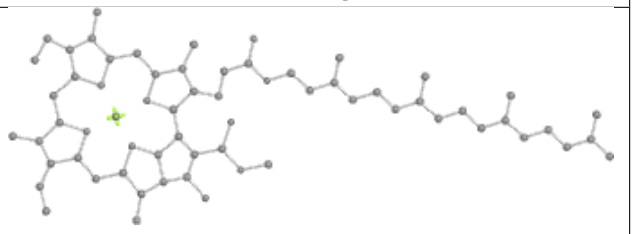
Bond lengths



Bond angles



Torsions



Rings

5.7 Other polymers

There are no such residues in this entry.

5.8 Polymer linkage issues

The following chains have linkage breaks:

Mol	Chain	Number of breaks
11	L	1

All chain breaks are listed below:

Model	Chain	Residue-1	Atom-1	Residue-2	Atom-2	Distance (Å)
1	L	144:PRO	C	158:ARG	N	6.44

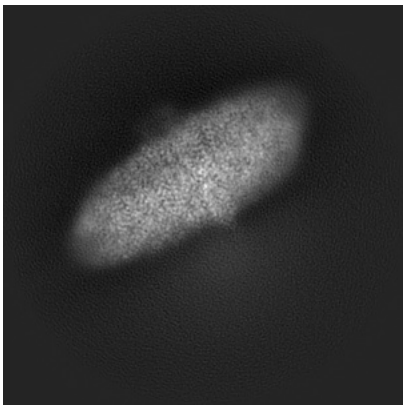
6 Map visualisation

This section contains visualisations of the EMDB entry EMD-16732. These allow visual inspection of the internal detail of the map and identification of artifacts.

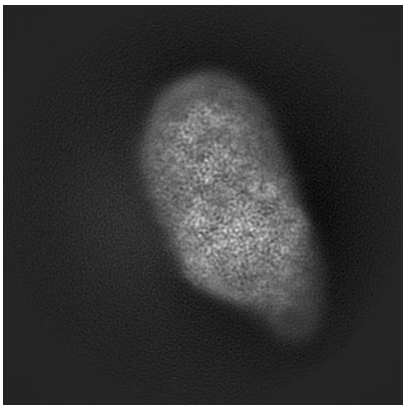
Images derived from a raw map, generated by summing the deposited half-maps, are presented below the corresponding image components of the primary map to allow further visual inspection and comparison with those of the primary map.

6.1 Orthogonal projections

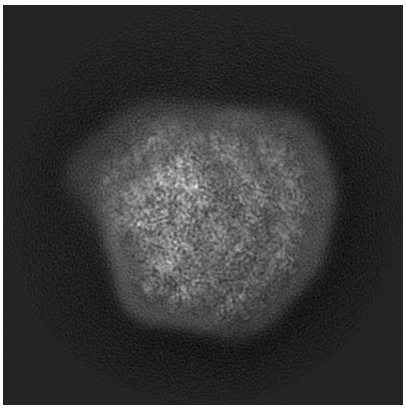
6.1.1 Primary map



X

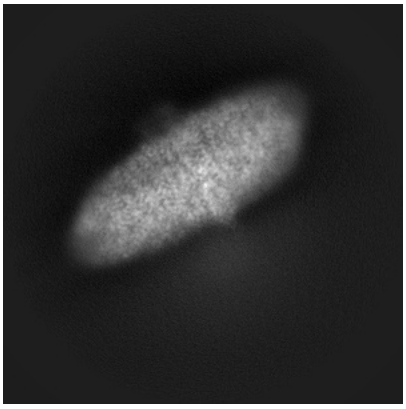


Y

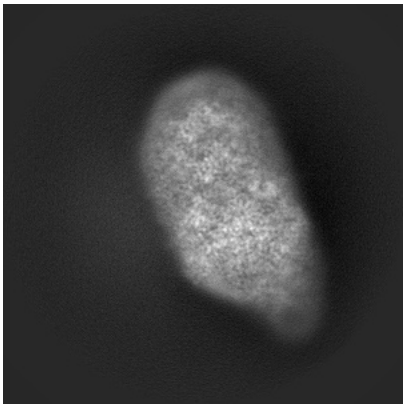


Z

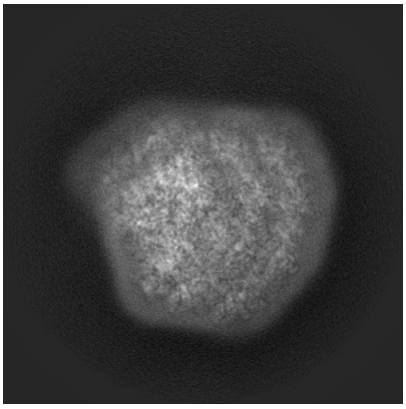
6.1.2 Raw map



X



Y

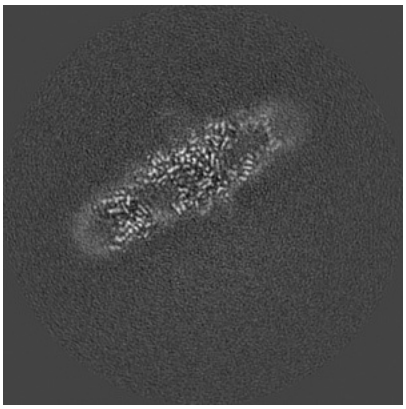


Z

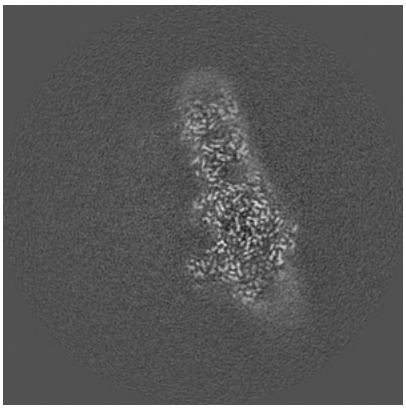
The images above show the map projected in three orthogonal directions.

6.2 Central slices (i)

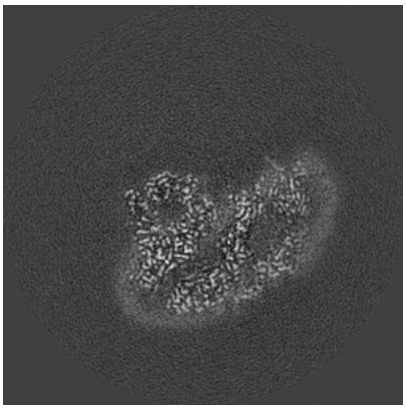
6.2.1 Primary map



X Index: 250

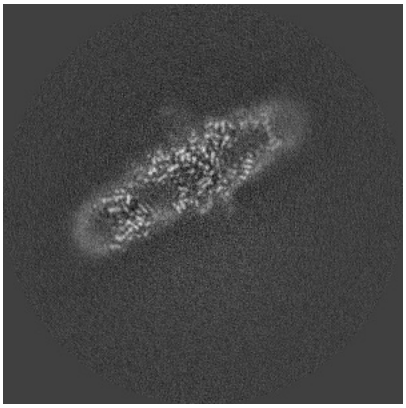


Y Index: 250

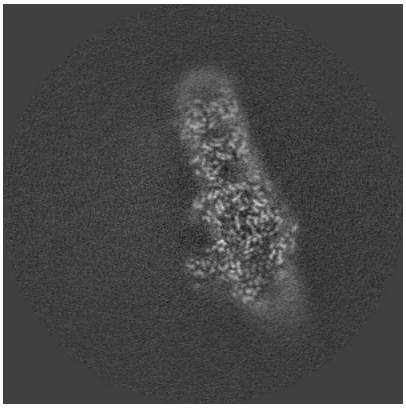


Z Index: 250

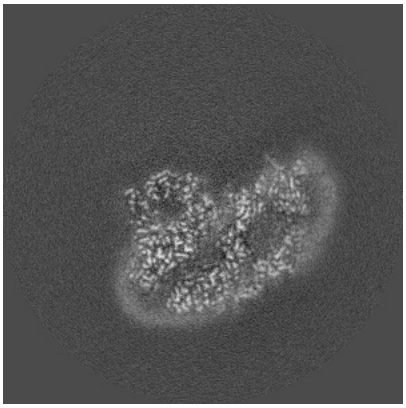
6.2.2 Raw map



X Index: 250



Y Index: 250

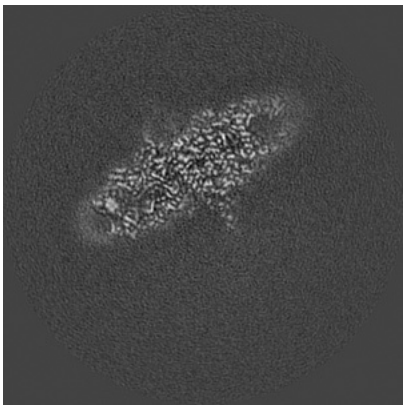


Z Index: 250

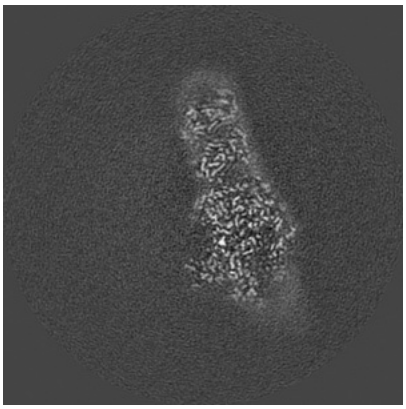
The images above show central slices of the map in three orthogonal directions.

6.3 Largest variance slices (i)

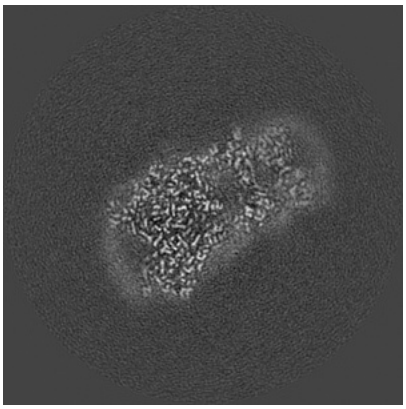
6.3.1 Primary map



X Index: 218

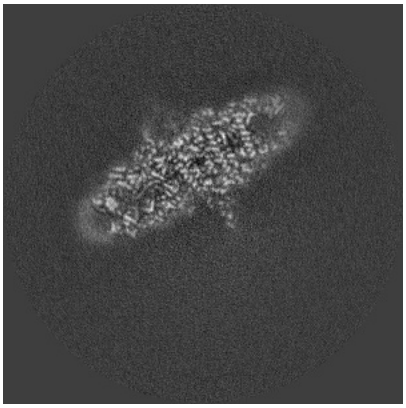


Y Index: 252

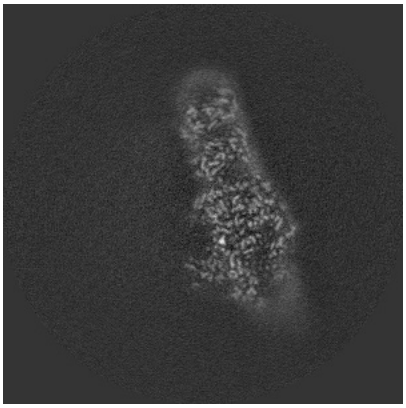


Z Index: 289

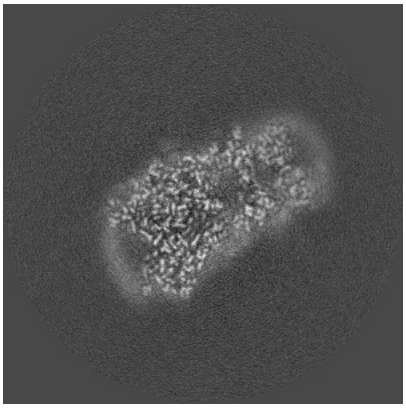
6.3.2 Raw map



X Index: 218



Y Index: 252

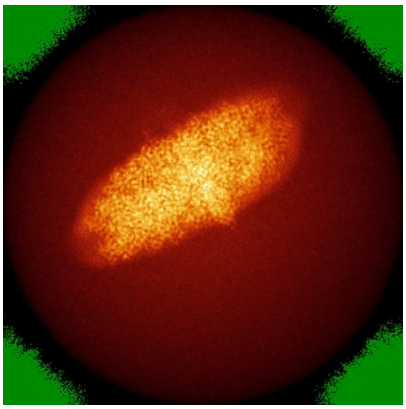


Z Index: 289

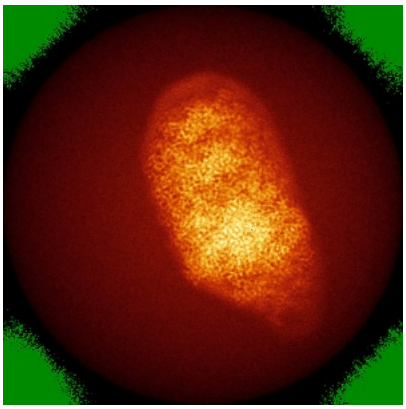
The images above show the largest variance slices of the map in three orthogonal directions.

6.4 Orthogonal standard-deviation projections (False-color) (i)

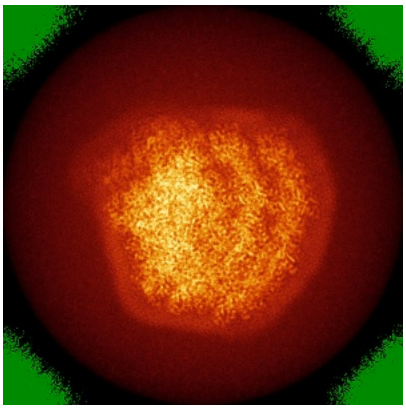
6.4.1 Primary map



X

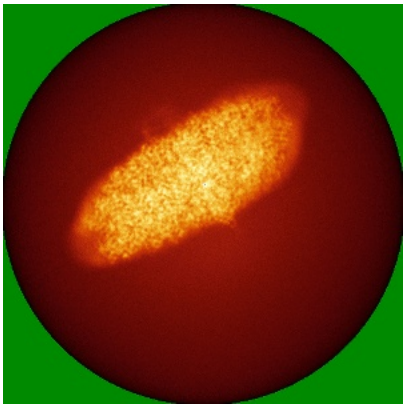


Y

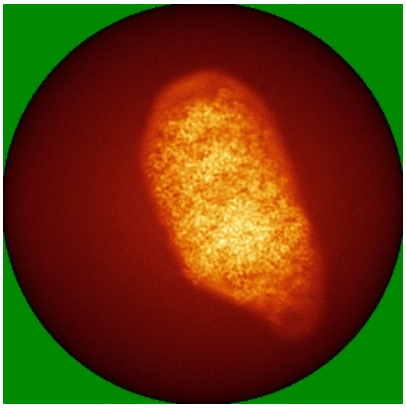


Z

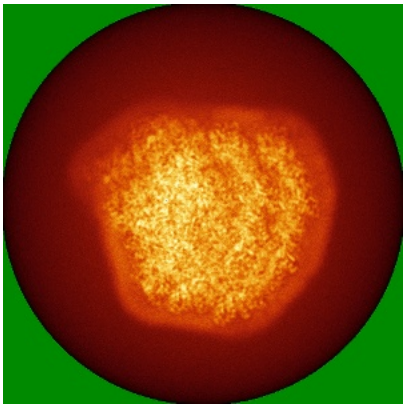
6.4.2 Raw map



X



Y



Z

The images above show the map standard deviation projections with false color in three orthogonal directions. Minimum values are shown in green, max in blue, and dark to light orange shades represent small to large values respectively.

6.5 Orthogonal surface views (i)

6.5.1 Primary map

X Y Z

The images above show the 3D surface view of the map at the recommended contour level 0.007. These images, in conjunction with the slice images, may facilitate assessment of whether an appropriate contour level has been provided.

6.5.2 Raw map

X Y Z

These images show the 3D surface of the raw map. The raw map's contour level was selected so that its surface encloses the same volume as the primary map does at its recommended contour level.

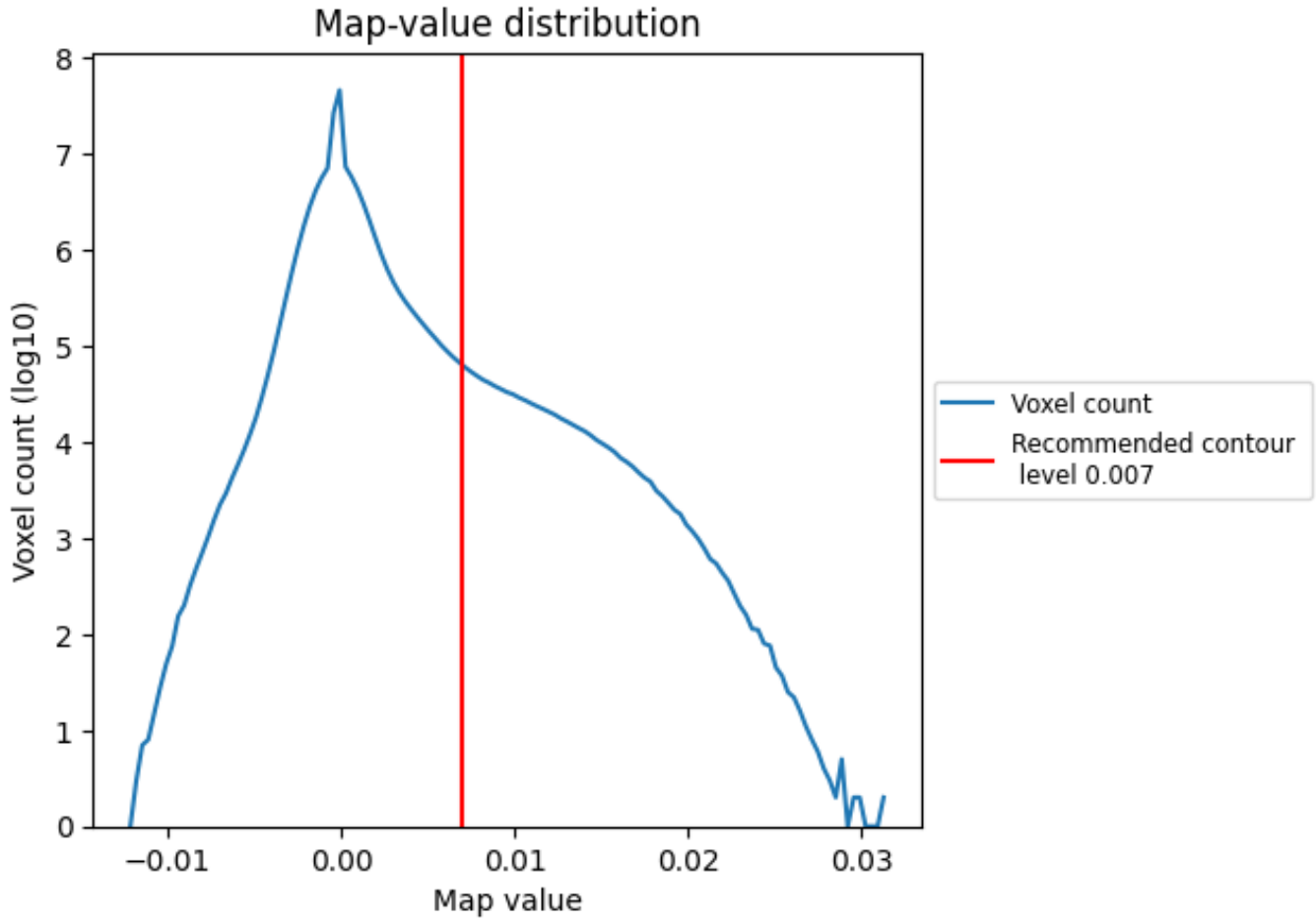
6.6 Mask visualisation (i)

This section was not generated. No masks/segmentation were deposited.

7 Map analysis

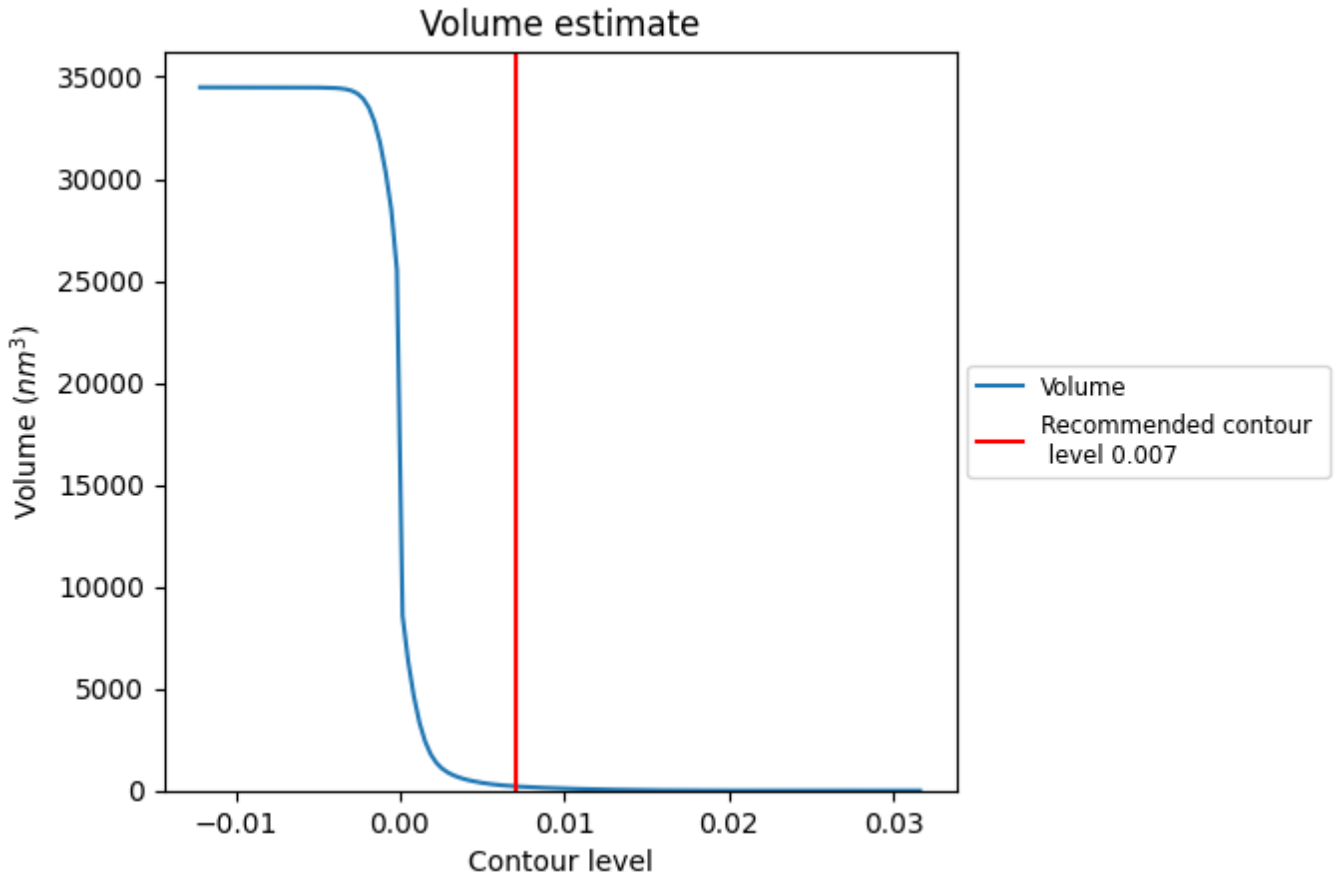
This section contains the results of statistical analysis of the map.

7.1 Map-value distribution



The map-value distribution is plotted in 128 intervals along the x-axis. The y-axis is logarithmic. A spike in this graph at zero usually indicates that the volume has been masked.

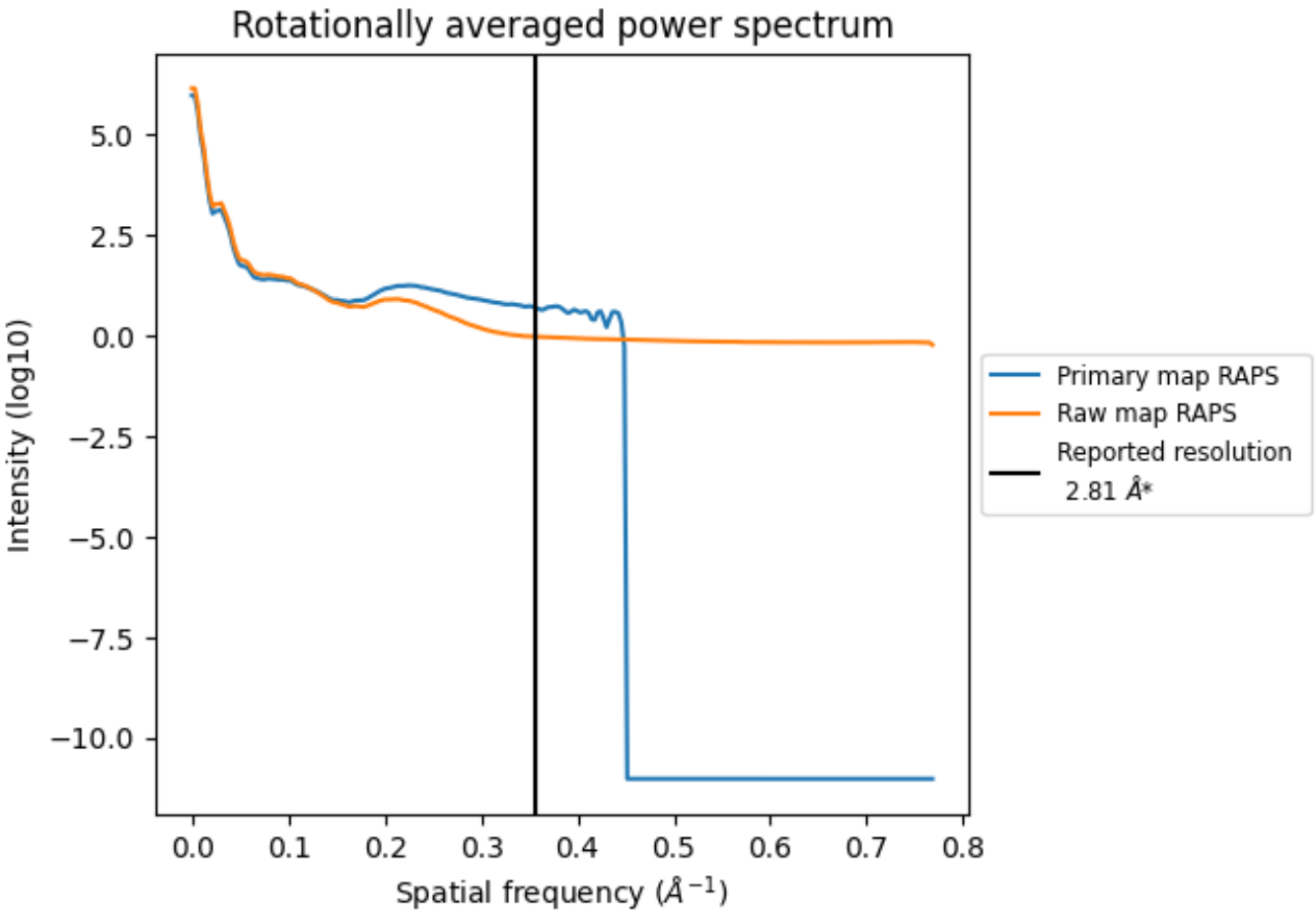
7.2 Volume estimate



The volume at the recommended contour level is 210 nm^3; this corresponds to an approximate mass of 190 kDa.

The volume estimate graph shows how the enclosed volume varies with the contour level. The recommended contour level is shown as a vertical line and the intersection between the line and the curve gives the volume of the enclosed surface at the given level.

7.3 Rotationally averaged power spectrum (i)

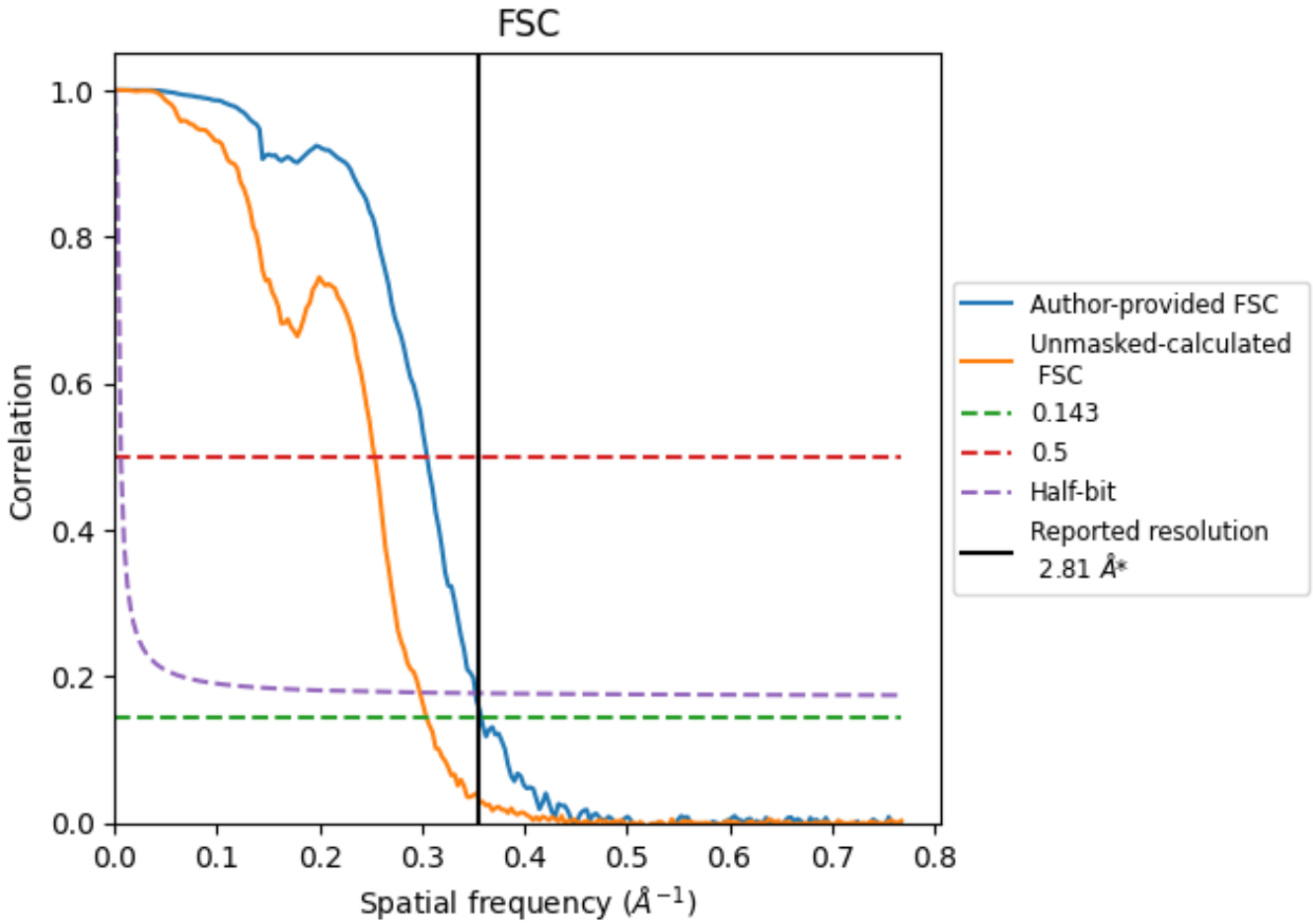


*Reported resolution corresponds to spatial frequency of 0.356 $Å^{-1}$

8 Fourier-Shell correlation

Fourier-Shell Correlation (FSC) is the most commonly used method to estimate the resolution of single-particle and subtomogram-averaged maps. The shape of the curve depends on the imposed symmetry, mask and whether or not the two 3D reconstructions used were processed from a common reference. The reported resolution is shown as a black line. A curve is displayed for the half-bit criterion in addition to lines showing the 0.143 gold standard cut-off and 0.5 cut-off.

8.1 FSC



*Reported resolution corresponds to spatial frequency of 0.356 Å^{-1}

8.2 Resolution estimates (i)

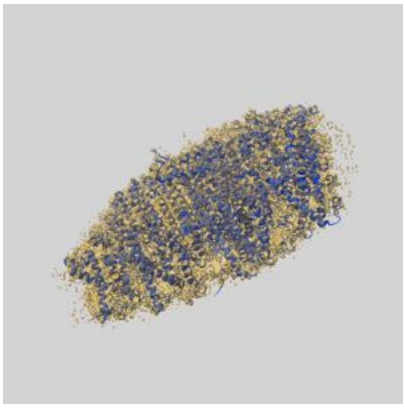
Resolution estimate (Å)	Estimation criterion (FSC cut-off)		
	0.143	0.5	Half-bit
Reported by author	2.81	-	-
Author-provided FSC curve	2.79	3.28	2.84
Unmasked-calculated*	3.28	3.94	3.36

*Resolution estimate based on FSC curve calculated by comparison of deposited half-maps. The value from deposited half-maps intersecting FSC 0.143 CUT-OFF 3.28 differs from the reported value 2.81 by more than 10 %

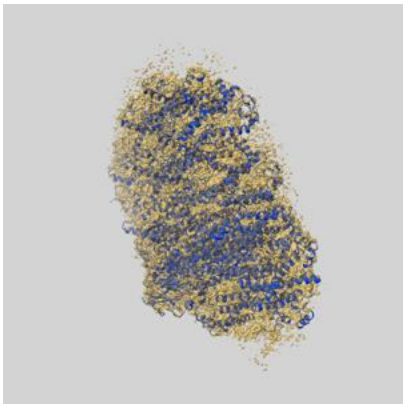
9 Map-model fit

This section contains information regarding the fit between EMDB map EMD-16732 and PDB model 8CMO. Per-residue inclusion information can be found in section 3 on page 41.

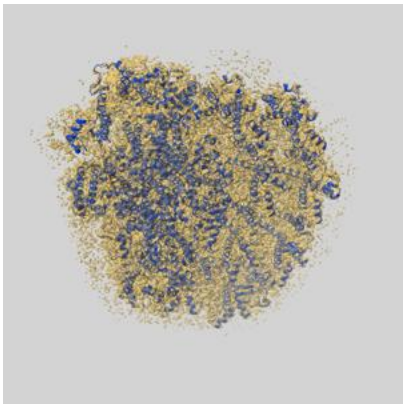
9.1 Map-model overlay



X



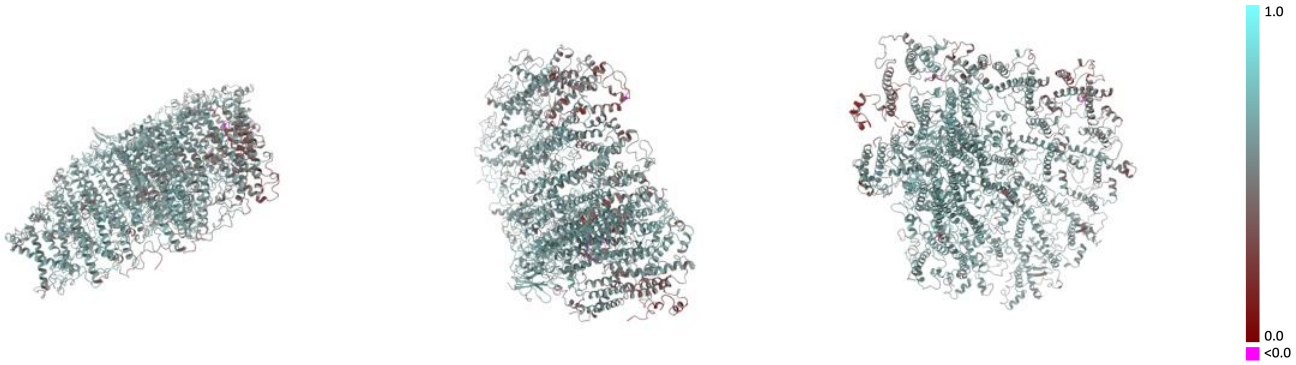
Y



Z

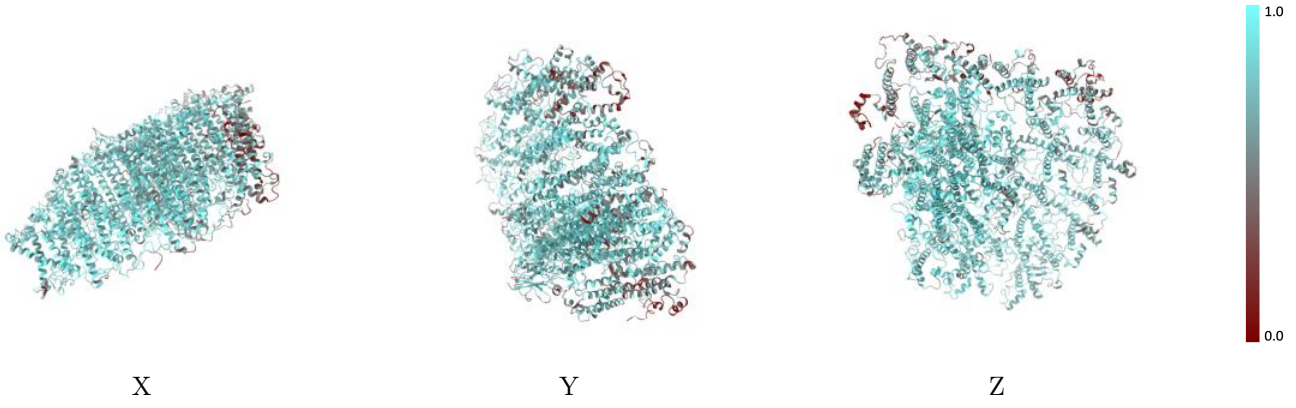
The images above show the 3D surface view of the map at the recommended contour level 0.007 at 50% transparency in yellow overlaid with a ribbon representation of the model coloured in blue. These images allow for the visual assessment of the quality of fit between the atomic model and the map.

9.2 Q-score mapped to coordinate model (i)



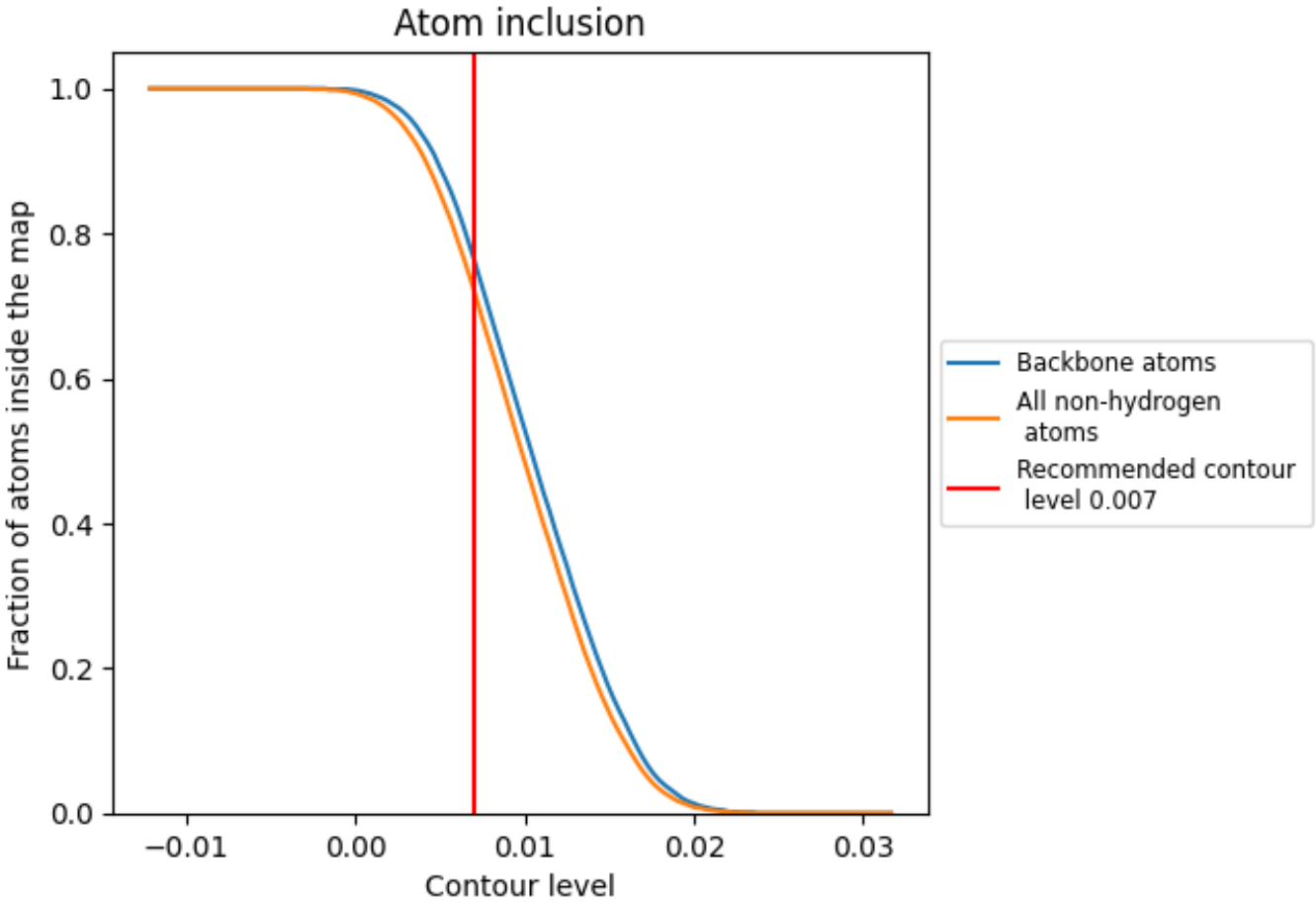
The images above show the model with each residue coloured according its Q-score. This shows their resolvability in the map with higher Q-score values reflecting better resolvability. Please note: Q-score is calculating the resolvability of atoms, and thus high values are only expected at resolutions at which atoms can be resolved. Low Q-score values may therefore be expected for many entries.

9.3 Atom inclusion mapped to coordinate model (i)



The images above show the model with each residue coloured according to its atom inclusion. This shows to what extent they are inside the map at the recommended contour level (0.007).

9.4 Atom inclusion (i)



At the recommended contour level, 76% of all backbone atoms, 72% of all non-hydrogen atoms, are inside the map.

9.5 Map-model fit summary

The table lists the average atom inclusion at the recommended contour level (0.007) and Q-score for the entire model and for each chain.

Chain	Atom inclusion	Q-score
All	0.7200	0.5370
1	0.6350	0.4980
2	0.3310	0.3380
3	0.7710	0.5610
4	0.6470	0.5050
5	0.6780	0.5150
6	0.6810	0.5140
7	0.7840	0.5510
8	0.7410	0.5310
9	0.5380	0.4770
A	0.8270	0.5920
B	0.8210	0.5830
C	0.7720	0.6020
D	0.6520	0.5660
E	0.6830	0.5660
F	0.6980	0.5310
G	0.4660	0.4150
I	0.6720	0.5270
J	0.7000	0.5310
K	0.7110	0.5230
L	0.6550	0.5030
Z	0.4730	0.4100