



Full wwPDB EM Validation Report ⓘ

Mar 7, 2026 – 01:18 AM UTC

PDB ID : 8ZET / pdb_00008zet
EMDB ID : EMD-60044
Title : Tp-PSI-FCPI-S in Thalassiosira pseudonana
Authors : Feng, Y.; Li, Z.; Shen, J.R.; Wang, W.
Deposited on : 2024-05-06
Resolution : 3.20 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.dev132
Mogul : 2022.3.0, CSD as543be (2022)
MolProbity : 4-5-2 with Phenix2.0
Buster-report : wwPDB partial adaption of 1.1.7 (2018)
Percentile statistics : 20250101.v01 (using entries in the PDB archive January 1st 2025)
EM percentile statistics : 202505.v01 (Using data in the EMDB archive up until May 2025)
MapQ : 1.9.13
Ideal geometry (proteins) : Engh & Huber (2001)
Ideal geometry (DNA, RNA) : Parkinson et al. (1996)
Validation Pipeline (wwPDB-VP) : 2.49

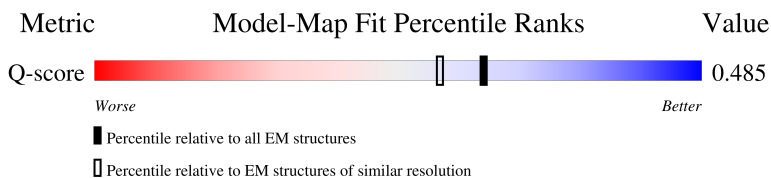
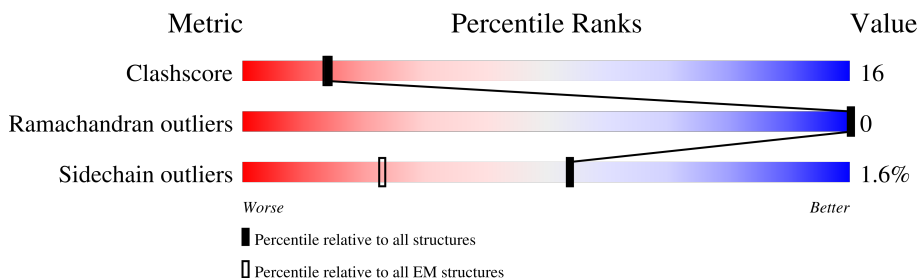
1 Overall quality at a glance

The following experimental techniques were used to determine the structure:

ELECTRON MICROSCOPY

The reported resolution of this entry is 3.20 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.



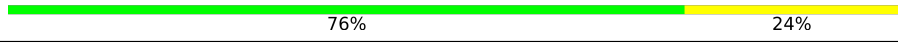
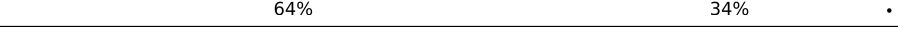
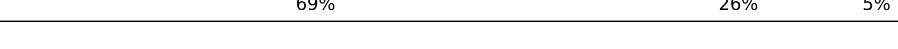
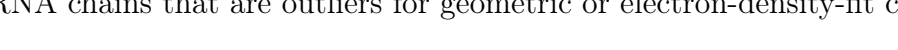
Metric	Whole archive (#Entries)	EM structures (#Entries)	Similar EM resolution (#Entries, resolution range(Å))
Clashscore	229148	23984	-
Ramachandran outliers	224038	23583	-
Sidechain outliers	223484	23102	-
Q-score	-	25397	15020 (2.70 - 3.70)

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 $\leq 5\%$. 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	743	
2	b	732	
3	c	80	
4	d	132	

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Mol	Chain	Length	Quality of chain
5	e	62	
6	f	160	
7	g	131	
8	i	33	
9	j	40	
10	l	146	
11	m	29	
12	r	89	
13	B	165	
14	C	170	
15	D	164	
16	E	185	
17	H	168	

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
18	CLA	B	306	X	-	-	-
18	CLA	B	307	X	-	-	-
18	CLA	B	308	X	-	-	-
18	CLA	B	309	X	-	-	-
18	CLA	B	310	X	-	-	-
18	CLA	B	311	X	-	-	-
18	CLA	B	312	X	-	-	-
18	CLA	B	314	X	-	-	-
18	CLA	C	306	X	-	-	-
18	CLA	C	307	X	-	-	-
18	CLA	C	309	X	-	-	-
18	CLA	C	310	X	-	-	-
18	CLA	C	311	X	-	-	-
18	CLA	C	312	X	-	-	-
18	CLA	C	314	X	-	-	-

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Mol	Type	Chain	Res	Chirality	Geometry	Clashes	Electron density
18	CLA	C	315	X	-	-	-
18	CLA	C	316	X	-	-	-
18	CLA	C	317	X	-	-	-
18	CLA	C	318	X	-	-	-
18	CLA	D	207	X	-	-	-
18	CLA	D	208	X	-	-	-
18	CLA	D	209	X	-	-	-
18	CLA	D	210	X	-	-	-
18	CLA	D	211	X	-	-	-
18	CLA	D	212	X	-	-	-
18	CLA	D	213	X	-	-	-
18	CLA	D	214	X	-	-	-
18	CLA	D	215	X	-	-	-
18	CLA	D	216	X	-	-	-
18	CLA	D	217	X	-	-	-
18	CLA	E	309	X	-	-	-
18	CLA	E	310	X	-	-	-
18	CLA	E	311	X	-	-	-
18	CLA	E	312	X	-	-	-
18	CLA	E	313	X	-	-	-
18	CLA	E	314	X	-	-	-
18	CLA	E	315	X	-	-	-
18	CLA	E	316	X	-	-	-
18	CLA	H	305	X	-	-	-
18	CLA	H	306	X	-	-	-
18	CLA	H	307	X	-	-	-
18	CLA	H	308	X	-	-	-
18	CLA	H	309	X	-	-	-
18	CLA	H	310	X	-	-	-
18	CLA	H	311	X	-	-	-
18	CLA	H	313	X	-	-	-
18	CLA	H	314	X	-	-	-
18	CLA	H	315	X	-	-	-
18	CLA	a	801	X	-	-	-
18	CLA	a	802	X	-	-	-
18	CLA	a	803	X	-	-	-
18	CLA	a	804	X	-	-	-
18	CLA	a	805	X	-	-	-
18	CLA	a	806	X	-	-	-
18	CLA	a	807	X	-	-	-
18	CLA	a	808	X	-	-	-
18	CLA	a	809	X	-	-	-

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Mol	Type	Chain	Res	Chirality	Geometry	Clashes	Electron density
18	CLA	a	810	X	-	-	-
18	CLA	a	811	X	-	-	-
18	CLA	a	812	X	-	-	-
18	CLA	a	813	X	-	-	-
18	CLA	a	814	X	-	-	-
18	CLA	a	815	X	-	-	-
18	CLA	a	816	X	-	-	-
18	CLA	a	817	X	-	-	-
18	CLA	a	818	X	-	-	-
18	CLA	a	819	X	-	-	-
18	CLA	a	820	X	-	-	-
18	CLA	a	821	X	-	-	-
18	CLA	a	822	X	-	-	-
18	CLA	a	823	X	-	-	-
18	CLA	a	824	X	-	-	-
18	CLA	a	825	X	-	-	-
18	CLA	a	826	X	-	-	-
18	CLA	a	827	X	-	-	-
18	CLA	a	828	X	-	-	-
18	CLA	a	830	X	-	-	-
18	CLA	a	836	X	-	-	-
18	CLA	a	837	X	-	-	-
18	CLA	a	838	X	-	-	-
18	CLA	a	839	X	-	-	-
18	CLA	a	840	X	-	-	-
18	CLA	a	841	X	-	-	-
18	CLA	a	842	X	-	-	-
18	CLA	a	843	X	-	-	-
18	CLA	a	844	X	-	-	-
18	CLA	a	846	X	-	-	-
18	CLA	a	847	X	-	-	-
18	CLA	a	848	X	-	-	-
18	CLA	a	853	X	-	-	-
18	CLA	a	854	X	-	-	-
18	CLA	a	855	X	-	-	-
18	CLA	b	801	X	-	-	-
18	CLA	b	803	X	-	-	-
18	CLA	b	804	X	-	-	-
18	CLA	b	805	X	-	-	-
18	CLA	b	806	X	-	-	-
18	CLA	b	807	X	-	-	-
18	CLA	b	808	X	-	-	-

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Mol	Type	Chain	Res	Chirality	Geometry	Clashes	Electron density
18	CLA	b	809	X	-	-	-
18	CLA	b	810	X	-	-	-
18	CLA	b	811	X	-	-	-
18	CLA	b	812	X	-	-	-
18	CLA	b	813	X	-	-	-
18	CLA	b	814	X	-	-	-
18	CLA	b	815	X	-	-	-
18	CLA	b	816	X	-	-	-
18	CLA	b	817	X	-	-	-
18	CLA	b	818	X	-	-	-
18	CLA	b	819	X	-	-	-
18	CLA	b	820	X	-	-	-
18	CLA	b	821	X	-	-	-
18	CLA	b	822	X	-	-	-
18	CLA	b	823	X	-	-	-
18	CLA	b	824	X	-	-	-
18	CLA	b	825	X	-	-	-
18	CLA	b	826	X	-	-	-
18	CLA	b	827	X	-	-	-
18	CLA	b	828	X	-	-	-
18	CLA	b	829	X	-	-	-
18	CLA	b	830	X	-	-	-
18	CLA	b	837	X	-	-	-
18	CLA	b	838	X	-	-	-
18	CLA	b	840	X	-	-	-
18	CLA	b	841	X	-	-	-
18	CLA	b	842	X	-	-	-
18	CLA	b	843	X	-	-	-
18	CLA	b	844	X	-	-	-
18	CLA	b	845	X	-	-	-
18	CLA	b	846	X	-	-	-
18	CLA	b	848	X	-	-	-
18	CLA	b	849	X	-	-	-
18	CLA	b	850	X	-	-	-
18	CLA	f	202	X	-	-	-
18	CLA	f	203	X	-	-	-
18	CLA	f	204	X	-	-	-
18	CLA	i	101	X	-	-	-
18	CLA	j	104	X	-	-	-
18	CLA	l	202	X	-	-	-
18	CLA	l	203	X	-	-	-
18	CLA	l	205	X	-	-	-

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Mol	Type	Chain	Res	Chirality	Geometry	Clashes	Electron density
18	CLA	r	202	X	-	-	-
24	SF4	b	802	-	-	X	-
24	SF4	c	101	-	-	X	-
24	SF4	c	102	-	-	X	-

2 Entry composition

There are 29 unique types of molecules in this entry. The entry contains 36577 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.

Mol	Chain	Residues	Atoms					AltConf	Trace
1	a	743	Total	C	N	O	S	0	0
			5852	3822	992	1009	29		

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

Mol	Chain	Residues	Atoms					AltConf	Trace
2	b	732	Total	C	N	O	S	0	0
			5824	3827	982	996	19		

- Molecule 3 is a protein called Photosystem I iron-sulfur center.

Mol	Chain	Residues	Atoms					AltConf	Trace
3	c	80	Total	C	N	O	S	0	0
			599	368	103	118	10		

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

Mol	Chain	Residues	Atoms					AltConf	Trace
4	d	132	Total	C	N	O	S	0	0
			1040	665	177	195	3		

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

Mol	Chain	Residues	Atoms					AltConf	Trace
5	e	62	Total	C	N	O	S	0	0
			503	317	89	96	1		

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

Mol	Chain	Residues	Atoms					AltConf	Trace
6	f	160	Total	C	N	O	S	0	0
			1242	795	211	233	3		

- Molecule 7 is a protein called Photosystem I reaction center subunit Psa29.

Mol	Chain	Residues	Atoms					AltConf	Trace
7	g	131	Total	C	N	O	S	0	0
			981	619	154	204	4		

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

Mol	Chain	Residues	Atoms					AltConf	Trace
8	i	33	Total	C	N	O	S	0	0
			256	177	34	44	1		

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

Mol	Chain	Residues	Atoms					AltConf	Trace
9	j	40	Total	C	N	O	S	0	0
			332	224	48	57	3		

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

Mol	Chain	Residues	Atoms					AltConf	Trace
10	l	146	Total	C	N	O	S	0	0
			1095	722	178	193	2		

- Molecule 11 is a protein called Photosystem I reaction center subunit XII.

Mol	Chain	Residues	Atoms					AltConf	Trace
11	m	29	Total	C	N	O	S	0	0
			220	147	33	38	2		

- Molecule 12 is a protein called Tp-PsaR.

Mol	Chain	Residues	Atoms					AltConf	Trace
12	r	89	Total	C	N	O	S	0	0
			683	443	112	121	7		

- Molecule 13 is a protein called Fucoxanthin chlorophyll a/c-binding protein Lhcq8.

Mol	Chain	Residues	Atoms					AltConf	Trace
13	B	165	Total	C	N	O	S	0	0
			1285	833	204	240	8		

- Molecule 14 is a protein called Fucoxanthin chl a/c light-harvesting protein, major type.

Mol	Chain	Residues	Atoms					AltConf	Trace
14	C	170	Total	C	N	O	S	0	0
			1302	841	213	240	8		

- Molecule 15 is a protein called Pt17531-like protein.

Mol	Chain	Residues	Atoms					AltConf	Trace
15	D	164	Total	C	N	O	S	0	0
			1271	819	208	234	10		

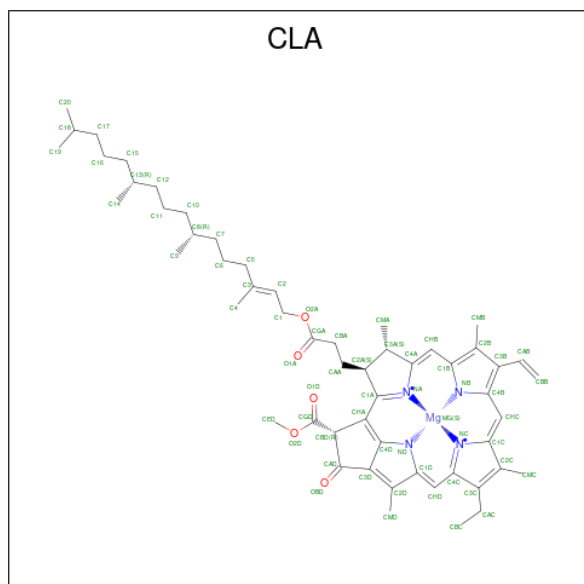
- Molecule 16 is a protein called Tp-RedCAP.

Mol	Chain	Residues	Atoms					AltConf	Trace
16	E	185	Total	C	N	O	S	0	0
			1413	902	235	262	14		

- Molecule 17 is a protein called Fucoxanthin chl a/c light-harvesting protein.

Mol	Chain	Residues	Atoms					AltConf	Trace
17	H	168	Total	C	N	O	S	0	0
			1300	834	214	242	10		

- Molecule 18 is CHLOROPHYLL A (CCD ID: CLA) (formula: $C_{55}H_{72}MgN_4O_5$) (labeled as "Ligand of Interest" by depositor).



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

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Mol	Chain	Residues	Atoms					AltConf
18	a	1	Total	C	Mg	N	O	0
			65	55	1	4	5	
18	a	1	Total	C	Mg	N	O	0
			65	55	1	4	5	
18	a	1	Total	C	Mg	N	O	0
			55	45	1	4	5	
18	a	1	Total	C	Mg	N	O	0
			65	55	1	4	5	
18	a	1	Total	C	Mg	N	O	0
			65	55	1	4	5	
18	a	1	Total	C	Mg	N	O	0
			65	55	1	4	5	
18	a	1	Total	C	Mg	N	O	0
			62	52	1	4	5	
18	a	1	Total	C	Mg	N	O	0
			65	55	1	4	5	
18	a	1	Total	C	Mg	N	O	0
			50	40	1	4	5	
18	a	1	Total	C	Mg	N	O	0
			45	35	1	4	5	
18	a	1	Total	C	Mg	N	O	0
			65	55	1	4	5	
18	a	1	Total	C	Mg	N	O	0
			65	55	1	4	5	
18	a	1	Total	C	Mg	N	O	0
			65	55	1	4	5	
18	a	1	Total	C	Mg	N	O	0
			45	35	1	4	5	
18	a	1	Total	C	Mg	N	O	0
			49	39	1	4	5	
18	a	1	Total	C	Mg	N	O	0
			51	41	1	4	5	
18	a	1	Total	C	Mg	N	O	0
			65	55	1	4	5	
18	a	1	Total	C	Mg	N	O	0
			65	55	1	4	5	
18	a	1	Total	C	Mg	N	O	0
			65	55	1	4	5	
18	a	1	Total	C	Mg	N	O	0
			50	40	1	4	5	

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Mol	Chain	Residues	Atoms					AltConf
18	a	1	Total	C	Mg	N	O	0
			45	35	1	4	5	
18	a	1	Total	C	Mg	N	O	0
			51	41	1	4	5	
18	a	1	Total	C	Mg	N	O	0
			60	50	1	4	5	
18	a	1	Total	C	Mg	N	O	0
			65	55	1	4	5	
18	a	1	Total	C	Mg	N	O	0
			65	55	1	4	5	
18	a	1	Total	C	Mg	N	O	0
			65	55	1	4	5	
18	a	1	Total	C	Mg	N	O	0
			52	42	1	4	5	
18	a	1	Total	C	Mg	N	O	0
			65	55	1	4	5	
18	a	1	Total	C	Mg	N	O	0
			60	50	1	4	5	
18	a	1	Total	C	Mg	N	O	0
			65	55	1	4	5	
18	a	1	Total	C	Mg	N	O	0
			65	55	1	4	5	
18	a	1	Total	C	Mg	N	O	0
			65	55	1	4	5	
18	a	1	Total	C	Mg	N	O	0
			65	55	1	4	5	
18	a	1	Total	C	Mg	N	O	0
			65	55	1	4	5	
18	a	1	Total	C	Mg	N	O	0
			65	55	1	4	5	
18	a	1	Total	C	Mg	N	O	0
			65	55	1	4	5	
18	a	1	Total	C	Mg	N	O	0
			56	46	1	4	5	
18	a	1	Total	C	Mg	N	O	0
			55	45	1	4	5	
18	a	1	Total	C	Mg	N	O	0
			65	55	1	4	5	

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Mol	Chain	Residues	Atoms					AltConf
18	a	1	Total 65	C 55	Mg 1	N 4	O 5	0
18	b	1	Total 65	C 55	Mg 1	N 4	O 5	0
18	b	1	Total 65	C 55	Mg 1	N 4	O 5	0
18	b	1	Total 45	C 35	Mg 1	N 4	O 5	0
18	b	1	Total 65	C 55	Mg 1	N 4	O 5	0
18	b	1	Total 65	C 55	Mg 1	N 4	O 5	0
18	b	1	Total 65	C 55	Mg 1	N 4	O 5	0
18	b	1	Total 65	C 55	Mg 1	N 4	O 5	0
18	b	1	Total 55	C 45	Mg 1	N 4	O 5	0
18	b	1	Total 65	C 55	Mg 1	N 4	O 5	0
18	b	1	Total 60	C 50	Mg 1	N 4	O 5	0
18	b	1	Total 59	C 49	Mg 1	N 4	O 5	0
18	b	1	Total 55	C 45	Mg 1	N 4	O 5	0
18	b	1	Total 65	C 55	Mg 1	N 4	O 5	0
18	b	1	Total 46	C 36	Mg 1	N 4	O 5	0
18	b	1	Total 55	C 45	Mg 1	N 4	O 5	0
18	b	1	Total 60	C 50	Mg 1	N 4	O 5	0
18	b	1	Total 65	C 55	Mg 1	N 4	O 5	0
18	b	1	Total 65	C 55	Mg 1	N 4	O 5	0
18	b	1	Total 65	C 55	Mg 1	N 4	O 5	0
18	b	1	Total 50	C 40	Mg 1	N 4	O 5	0

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Mol	Chain	Residues	Atoms					AltConf
18	b	1	Total 65	C 55	Mg 1	N 4	O 5	0
18	b	1	Total 58	C 48	Mg 1	N 4	O 5	0
18	b	1	Total 65	C 55	Mg 1	N 4	O 5	0
18	b	1	Total 45	C 35	Mg 1	N 4	O 5	0
18	b	1	Total 60	C 50	Mg 1	N 4	O 5	0
18	b	1	Total 47	C 37	Mg 1	N 4	O 5	0
18	b	1	Total 65	C 55	Mg 1	N 4	O 5	0
18	b	1	Total 65	C 55	Mg 1	N 4	O 5	0
18	b	1	Total 65	C 55	Mg 1	N 4	O 5	0
18	b	1	Total 60	C 50	Mg 1	N 4	O 5	0
18	b	1	Total 65	C 55	Mg 1	N 4	O 5	0
18	b	1	Total 61	C 51	Mg 1	N 4	O 5	0
18	b	1	Total 65	C 55	Mg 1	N 4	O 5	0
18	b	1	Total 65	C 55	Mg 1	N 4	O 5	0
18	b	1	Total 65	C 55	Mg 1	N 4	O 5	0
18	b	1	Total 62	C 52	Mg 1	N 4	O 5	0
18	b	1	Total 65	C 55	Mg 1	N 4	O 5	0
18	b	1	Total 65	C 55	Mg 1	N 4	O 5	0
18	b	1	Total 54	C 44	Mg 1	N 4	O 5	0
18	b	1	Total 65	C 55	Mg 1	N 4	O 5	0
18	b	1	Total 65	C 55	Mg 1	N 4	O 5	0

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Mol	Chain	Residues	Atoms					AltConf
18	f	1	Total 65	C 55	Mg 1	N 4	O 5	0
18	f	1	Total 65	C 55	Mg 1	N 4	O 5	0
18	f	1	Total 45	C 35	Mg 1	N 4	O 5	0
18	i	1	Total 60	C 50	Mg 1	N 4	O 5	0
18	j	1	Total 41	C 33	Mg 1	N 4	O 3	0
18	l	1	Total 49	C 39	Mg 1	N 4	O 5	0
18	l	1	Total 65	C 55	Mg 1	N 4	O 5	0
18	l	1	Total 45	C 35	Mg 1	N 4	O 5	0
18	r	1	Total 45	C 35	Mg 1	N 4	O 5	0
18	B	1	Total 57	C 49	Mg 1	N 4	O 3	0
18	B	1	Total 65	C 55	Mg 1	N 4	O 5	0
18	B	1	Total 65	C 55	Mg 1	N 4	O 5	0
18	B	1	Total 65	C 55	Mg 1	N 4	O 5	0
18	B	1	Total 46	C 36	Mg 1	N 4	O 5	0
18	B	1	Total 65	C 55	Mg 1	N 4	O 5	0
18	B	1	Total 65	C 55	Mg 1	N 4	O 5	0
18	B	1	Total 41	C 33	Mg 1	N 4	O 3	0
18	C	1	Total 45	C 35	Mg 1	N 4	O 5	0
18	C	1	Total 65	C 55	Mg 1	N 4	O 5	0
18	C	1	Total 62	C 52	Mg 1	N 4	O 5	0
18	C	1	Total 46	C 36	Mg 1	N 4	O 5	0

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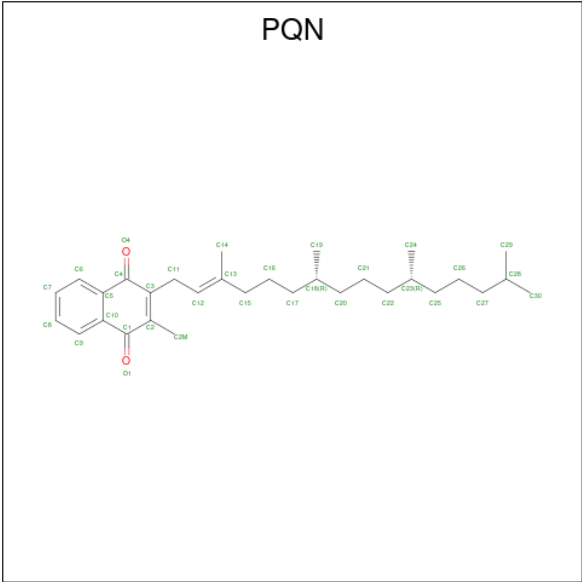
Mol	Chain	Residues	Atoms					AltConf
18	C	1	Total 60	C 50	Mg 1	N 4	O 5	0
18	C	1	Total 61	C 51	Mg 1	N 4	O 5	0
18	C	1	Total 41	C 33	Mg 1	N 4	O 3	0
18	C	1	Total 42	C 34	Mg 1	N 4	O 3	0
18	C	1	Total 41	C 33	Mg 1	N 4	O 3	0
18	C	1	Total 42	C 34	Mg 1	N 4	O 3	0
18	C	1	Total 42	C 34	Mg 1	N 4	O 3	0
18	D	1	Total 61	C 51	Mg 1	N 4	O 5	0
18	D	1	Total 65	C 55	Mg 1	N 4	O 5	0
18	D	1	Total 49	C 39	Mg 1	N 4	O 5	0
18	D	1	Total 56	C 46	Mg 1	N 4	O 5	0
18	D	1	Total 46	C 36	Mg 1	N 4	O 5	0
18	D	1	Total 42	C 34	Mg 1	N 4	O 3	0
18	D	1	Total 65	C 55	Mg 1	N 4	O 5	0
18	D	1	Total 56	C 46	Mg 1	N 4	O 5	0
18	D	1	Total 41	C 33	Mg 1	N 4	O 3	0
18	D	1	Total 46	C 36	Mg 1	N 4	O 5	0
18	D	1	Total 41	C 33	Mg 1	N 4	O 3	0
18	E	1	Total 65	C 55	Mg 1	N 4	O 5	0
18	E	1	Total 65	C 55	Mg 1	N 4	O 5	0
18	E	1	Total 46	C 36	Mg 1	N 4	O 5	0

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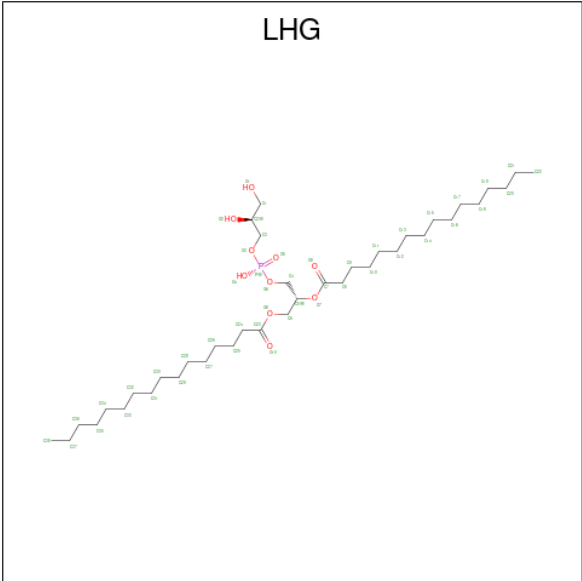
Mol	Chain	Residues	Atoms					AltConf
18	E	1	Total	C	Mg	N	O	0
			65	55	1	4	5	
18	E	1	Total	C	Mg	N	O	0
			55	45	1	4	5	
18	E	1	Total	C	Mg	N	O	0
			45	35	1	4	5	
18	E	1	Total	C	Mg	N	O	0
			65	55	1	4	5	
18	E	1	Total	C	Mg	N	O	0
			56	46	1	4	5	
18	H	1	Total	C	Mg	N	O	0
			54	44	1	4	5	
18	H	1	Total	C	Mg	N	O	0
			65	55	1	4	5	
18	H	1	Total	C	Mg	N	O	0
			65	55	1	4	5	
18	H	1	Total	C	Mg	N	O	0
			65	55	1	4	5	
18	H	1	Total	C	Mg	N	O	0
			46	36	1	4	5	
18	H	1	Total	C	Mg	N	O	0
			60	50	1	4	5	
18	H	1	Total	C	Mg	N	O	0
			57	47	1	4	5	
18	H	1	Total	C	Mg	N	O	0
			65	55	1	4	5	
18	H	1	Total	C	Mg	N	O	0
			65	55	1	4	5	
18	H	1	Total	C	Mg	N	O	0
			45	35	1	4	5	

- Molecule 19 is PHYLLOQUINONE (CCD ID: PQN) (formula: $C_{31}H_{46}O_2$) (labeled as "Ligand of Interest" by depositor).



Mol	Chain	Residues	Atoms			AltConf
19	a	1	Total	C	O	0
			33	31	2	
19	b	1	Total	C	O	0
			33	31	2	

- Molecule 20 is 1,2-DIPALMITOYL-PHOSPHATIDYL-GLYCEROLE (CCD ID: LHG) (formula: C₃₈H₇₅O₁₀P) (labeled as "Ligand of Interest" by depositor).



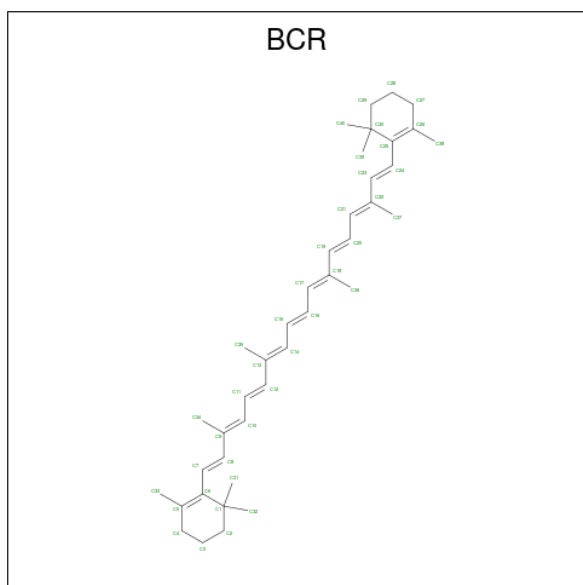
Mol	Chain	Residues	Atoms				AltConf
20	a	1	Total	C	O	P	0
			49	38	10	1	

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Mol	Chain	Residues	Atoms				AltConf
20	a	1	Total	C	O	P	0
			27	16	10	1	
20	a	1	Total	C	O	P	0
			33	22	10	1	
20	b	1	Total	C	O	P	0
			49	38	10	1	
20	j	1	Total	C	O	P	0
			49	38	10	1	
20	B	1	Total	C	O	P	0
			42	31	10	1	
20	D	1	Total	C	O	P	0
			49	38	10	1	
20	E	1	Total	C	O	P	0
			49	38	10	1	
20	E	1	Total	C	O	P	0
			42	31	10	1	
20	H	1	Total	C	O	P	0
			35	24	10	1	

- Molecule 21 is BETA-CAROTENE (CCD ID: BCR) (formula: $C_{40}H_{56}$) (labeled as "Ligand of Interest" by depositor).



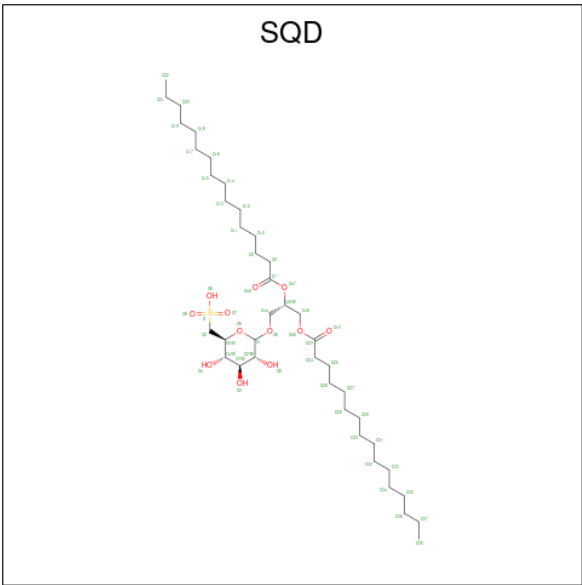
Mol	Chain	Residues	Atoms		AltConf
21	a	1	Total	C	0
			40	40	
21	a	1	Total	C	0
			40	40	

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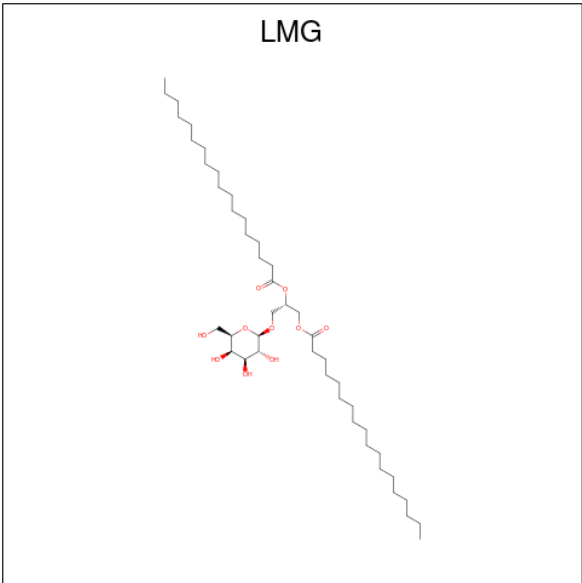
Mol	Chain	Residues	Atoms	AltConf
21	a	1	Total C 40 40	0
21	a	1	Total C 40 40	0
21	a	1	Total C 40 40	0
21	b	1	Total C 40 40	0
21	b	1	Total C 40 40	0
21	b	1	Total C 40 40	0
21	b	1	Total C 40 40	0
21	f	1	Total C 40 40	0
21	f	1	Total C 40 40	0
21	i	1	Total C 40 40	0
21	i	1	Total C 40 40	0
21	j	1	Total C 40 40	0
21	j	1	Total C 40 40	0
21	l	1	Total C 40 40	0
21	l	1	Total C 40 40	0
21	m	1	Total C 40 40	0
21	r	1	Total C 40 40	0
21	E	1	Total C 40 40	0
21	E	1	Total C 40 40	0

- Molecule 22 is 1,2-DI-O-ACYL-3-O-[6-DEOXY-6-SULFO-ALPHA-D-GLUCOPYRANOSYL]-SN-GLYCEROL (CCD ID: SQD) (formula: C₄₁H₇₈O₁₂S) (labeled as "Ligand of Interest" by depositor).



Mol	Chain	Residues	Atoms				AltConf
22	a	1	Total	C	O	S	0
			33	23	9	1	

- Molecule 23 is 1,2-DISTEAROYL-MONOGALACTOSYL-DIGLYCERIDE (CCD ID: LMG) (formula: C₄₅H₈₆O₁₀) (labeled as "Ligand of Interest" by depositor).



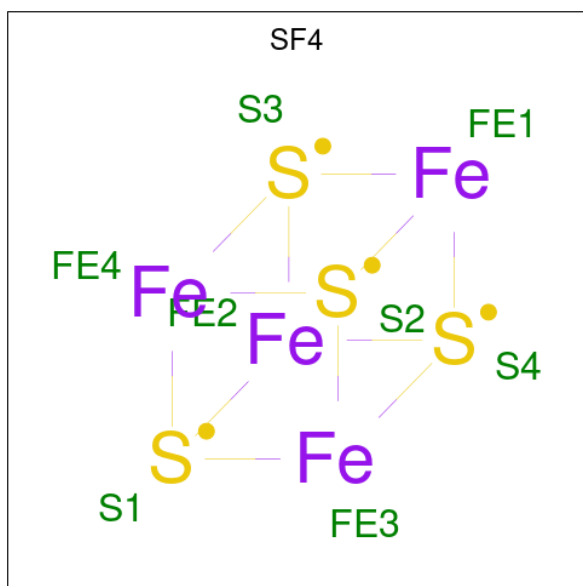
Mol	Chain	Residues	Atoms			AltConf
23	a	1	Total	C	O	0
			33	23	10	
23	j	1	Total	C	O	0
			37	27	10	

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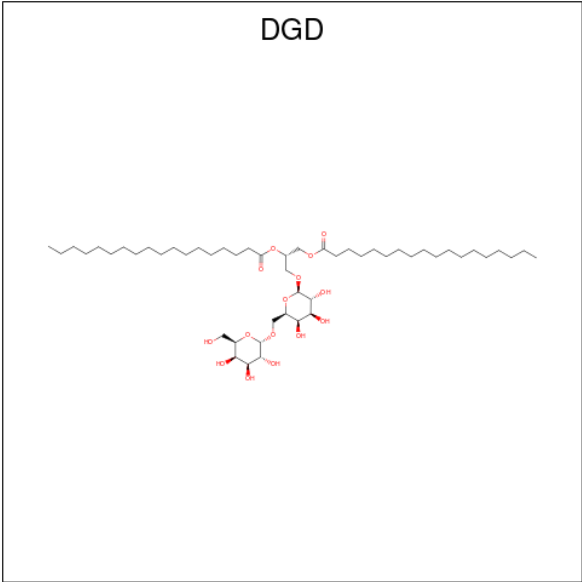
Mol	Chain	Residues	Atoms			AltConf
23	l	1	Total	C	O	0
			40	30	10	
23	C	1	Total	C	O	0
			31	21	10	
23	C	1	Total	C	O	0
			44	34	10	
23	D	1	Total	C	O	0
			46	36	10	
23	E	1	Total	C	O	0
			43	33	10	

- Molecule 24 is IRON/SULFUR CLUSTER (CCD ID: SF4) (formula: Fe_4S_4) (labeled as "Ligand of Interest" by depositor).



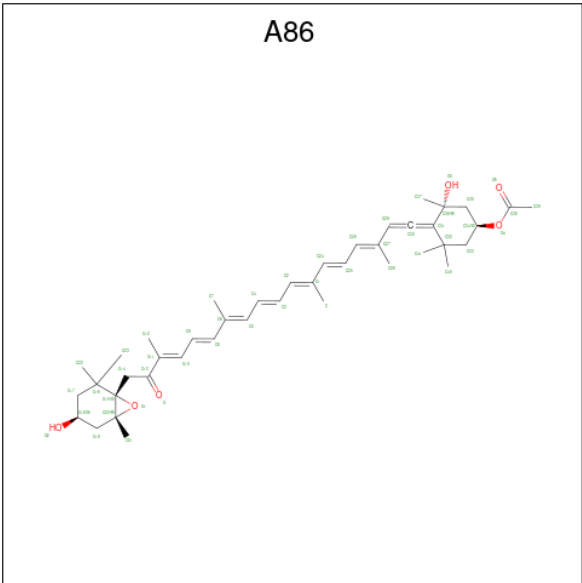
Mol	Chain	Residues	Atoms			AltConf
24	b	1	Total	Fe	S	0
			8	4	4	
24	c	1	Total	Fe	S	0
			8	4	4	
24	c	1	Total	Fe	S	0
			8	4	4	

- Molecule 25 is DIGALACTOSYL DIACYL GLYCEROL (DGDG) (CCD ID: DGD) (formula: $\text{C}_{51}\text{H}_{96}\text{O}_{15}$) (labeled as "Ligand of Interest" by depositor).



Mol	Chain	Residues	Atoms			AltConf
25	b	1	Total	C	O	0
			60	45	15	

- Molecule 26 is (3S,3'S,5R,5'R,6S,6'R,8'R)-3,5'-dihydroxy-8-oxo-6',7'-didehydro-5,5',6,6',7,8-hexahydro-5,6-epoxy-beta,beta-caroten-3'- yl acetate (CCD ID: A86) (formula: C₄₂H₅₈O₆) (labeled as "Ligand of Interest" by depositor).



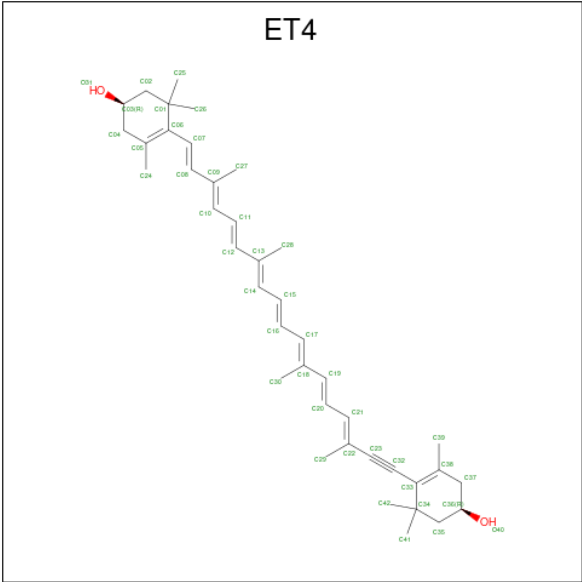
Mol	Chain	Residues	Atoms			AltConf
26	b	1	Total	C	O	0
			48	42	6	
26	m	1	Total	C	O	0
			48	42	6	

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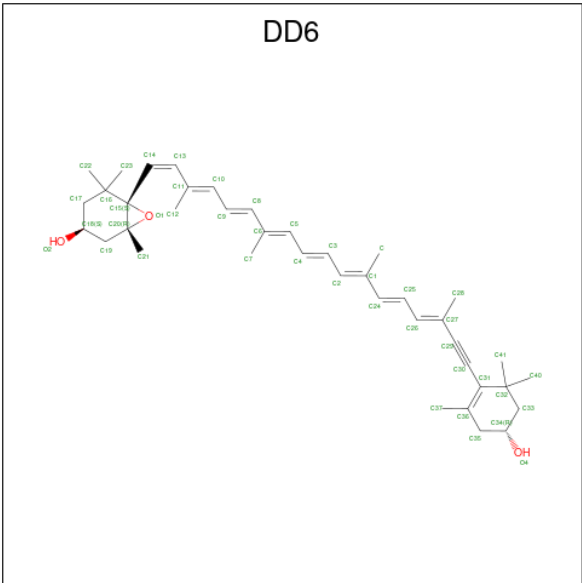
Mol	Chain	Residues	Atoms			AltConf
26	r	1	Total	C	O	0
			48	42	6	
26	B	1	Total	C	O	0
			48	42	6	
26	B	1	Total	C	O	0
			48	42	6	
26	B	1	Total	C	O	0
			48	42	6	
26	B	1	Total	C	O	0
			48	42	6	
26	C	1	Total	C	O	0
			48	42	6	
26	C	1	Total	C	O	0
			48	42	6	
26	C	1	Total	C	O	0
			48	42	6	
26	D	1	Total	C	O	0
			48	42	6	
26	D	1	Total	C	O	0
			48	42	6	
26	D	1	Total	C	O	0
			48	42	6	
26	E	1	Total	C	O	0
			48	42	6	
26	H	1	Total	C	O	0
			48	42	6	
26	H	1	Total	C	O	0
			48	42	6	
26	H	1	Total	C	O	0
			48	42	6	

- Molecule 27 is (1 {R})-3,5,5-trimethyl-4-[(1 {E},3 {E},5 {E},7 {E},9 {E},11 {E},13 {E},15 {E})-3,7,12,16-tetramethyl-18-[(4 {R})-2,6,6-trimethyl-4-oxidanyl-cyclohexen-1-yl]octadeca-1,3,5,7,9,11,13,15-octaen-17-ynyl]cyclohex-3-en-1-ol (CCD ID: ET4) (formula: C₄₀H₅₄O₂) (labeled as "Ligand of Interest" by depositor).



Mol	Chain	Residues	Atoms			AltConf
27	1	1	Total	C	O	0
			42	40	2	

- Molecule 28 is (3S,3'R,5R,6S,7cis)-7',8'-didehydro-5,6-dihydro-5,6-epoxy-beta,beta-carotene-3,3'-diol (CCD ID: DD6) (formula: C₄₀H₅₄O₃) (labeled as "Ligand of Interest" by depositor).



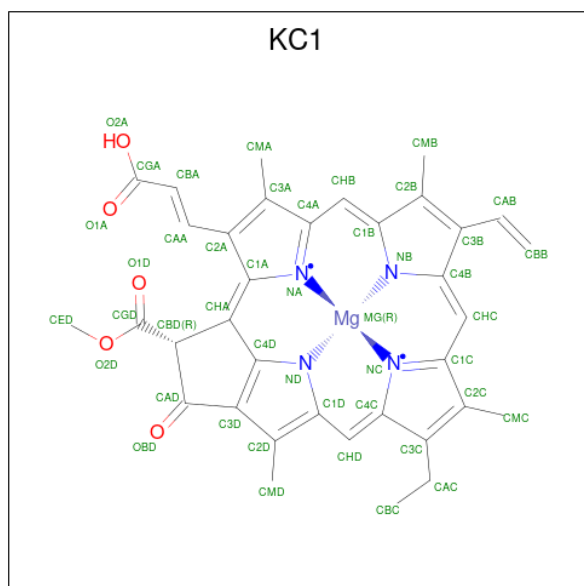
Mol	Chain	Residues	Atoms			AltConf
28	B	1	Total	C	O	0
			43	40	3	
28	C	1	Total	C	O	0
			43	40	3	

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Mol	Chain	Residues	Atoms			AltConf
28	D	1	Total	C	O	0
			43	40	3	
28	E	1	Total	C	O	0
			43	40	3	
28	E	1	Total	C	O	0
			43	40	3	
28	E	1	Total	C	O	0
			43	40	3	
28	E	1	Total	C	O	0
			43	40	3	
28	H	1	Total	C	O	0
			43	40	3	

- Molecule 29 is Chlorophyll c1 (CCD ID: KC1) (formula: $C_{35}H_{30}MgN_4O_5$) (labeled as "Ligand of Interest" by depositor).

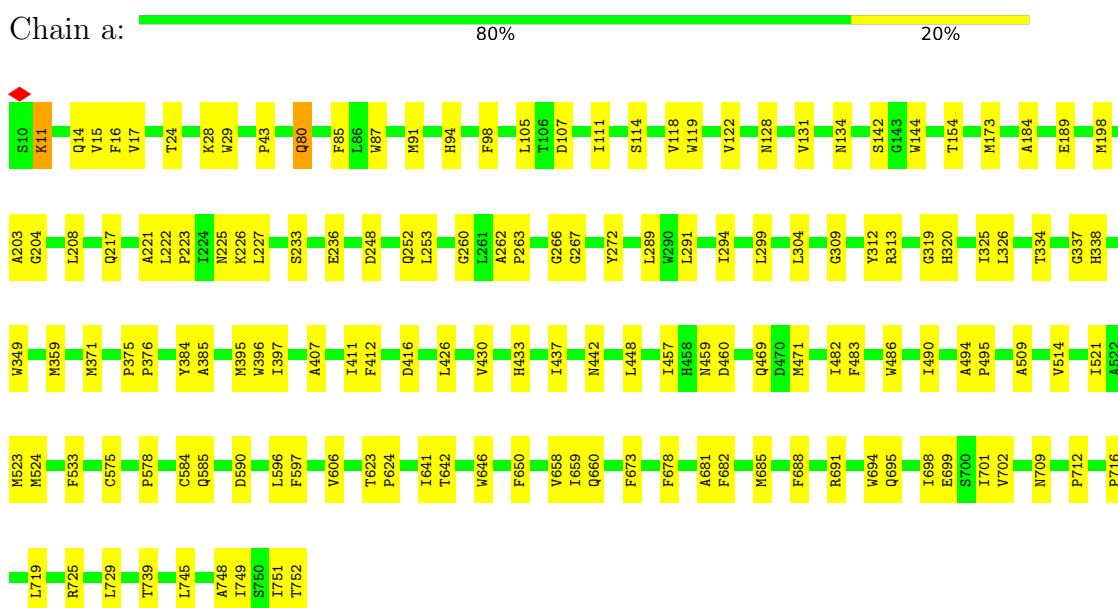


Mol	Chain	Residues	Atoms					AltConf
29	B	1	Total	C	Mg	N	O	0
			45	35	1	4	5	
29	C	1	Total	C	Mg	N	O	0
			45	35	1	4	5	
29	C	1	Total	C	Mg	N	O	0
			45	35	1	4	5	
29	H	1	Total	C	Mg	N	O	0
			45	35	1	4	5	

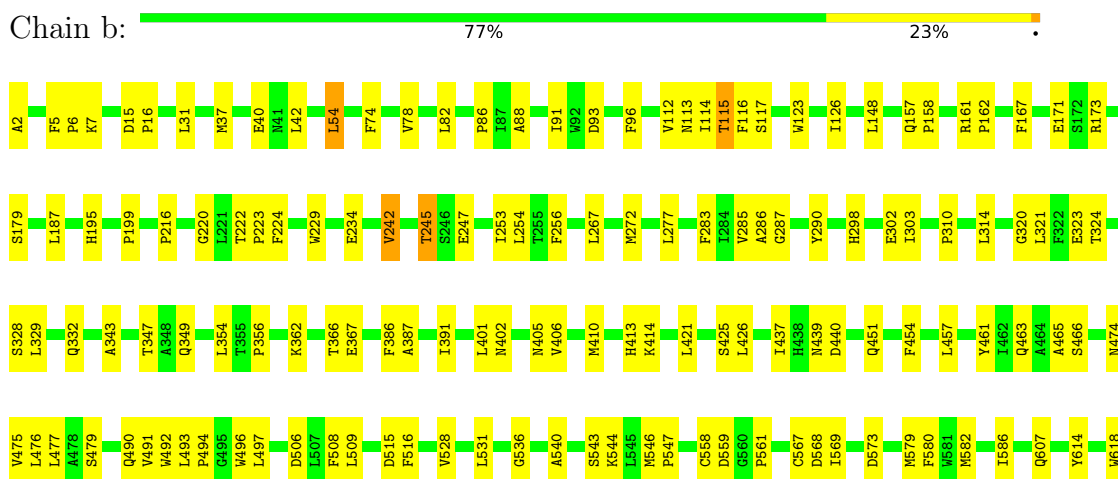
3 Residue-property plots

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

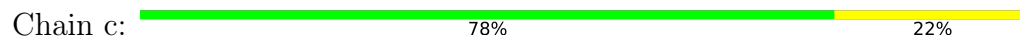


• Molecule 2: Photosystem I P700 chlorophyll a apoprotein A2





- Molecule 3: Photosystem I iron-sulfur center



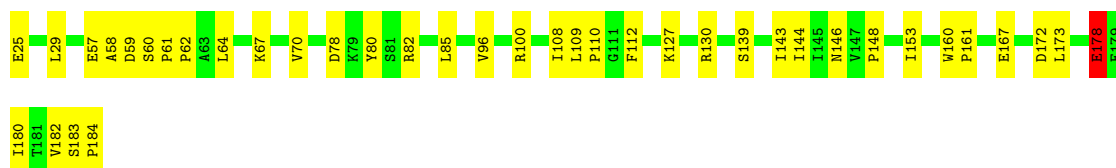
- Molecule 4: Photosystem I reaction center subunit II



- Molecule 5: Photosystem I reaction center subunit IV



- Molecule 6: Photosystem I reaction center subunit III



- Molecule 7: Photosystem I reaction center subunit Psa29



- Molecule 8: Photosystem I reaction center subunit VIII





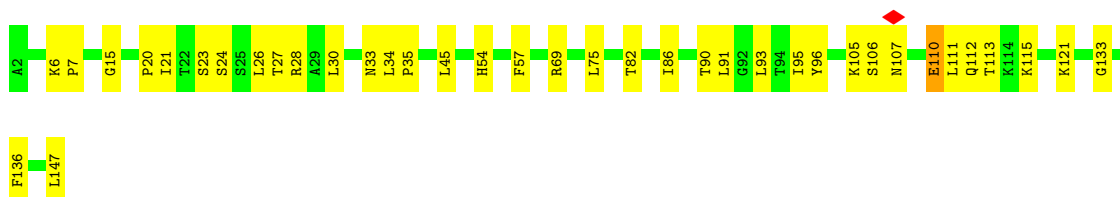
- Molecule 9: Photosystem I reaction center subunit IX

Chain j: 70% 22% 8%



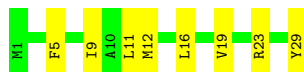
- Molecule 10: Photosystem I reaction center subunit XI

Chain l: 74% 25%



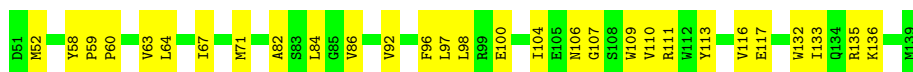
- Molecule 11: Photosystem I reaction center subunit XII

Chain m: 72% 28%



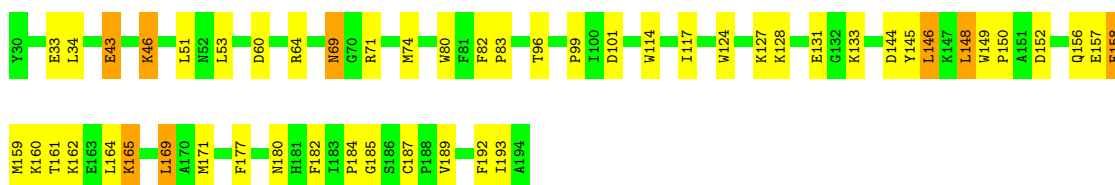
- Molecule 12: Tp-PsaR

Chain r: 67% 33%



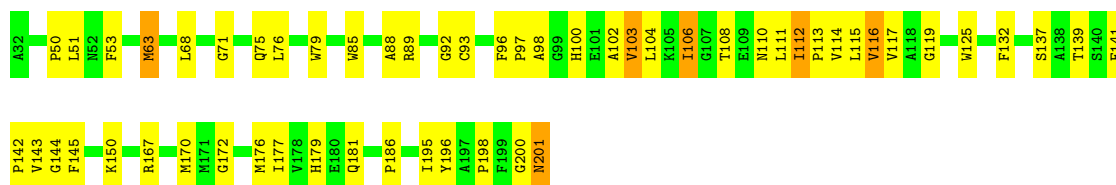
- Molecule 13: Fucoxanthin chlorophyll a/c-binding protein Lhcq8

Chain B: 69% 26% 5%



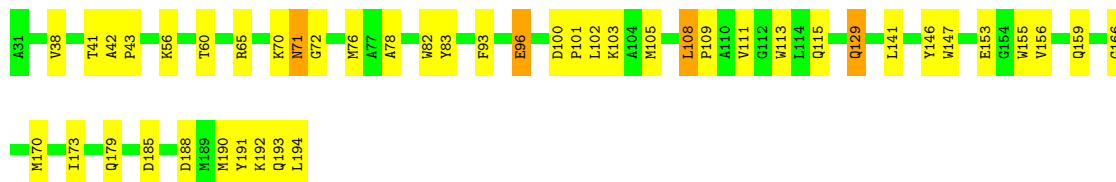
- Molecule 14: Fucoxanthin chl a/c light-harvesting protein, major type

Chain C: 68% 29%



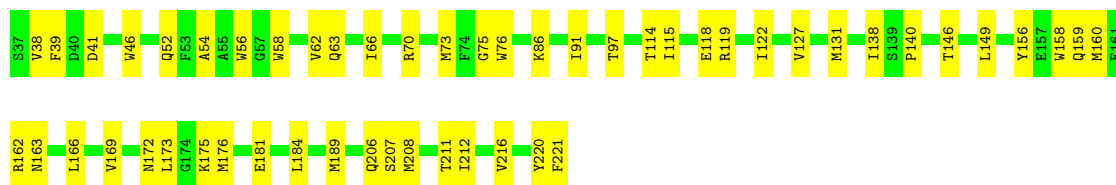
- Molecule 15: Pt17531-like protein

Chain D: 73% 25%



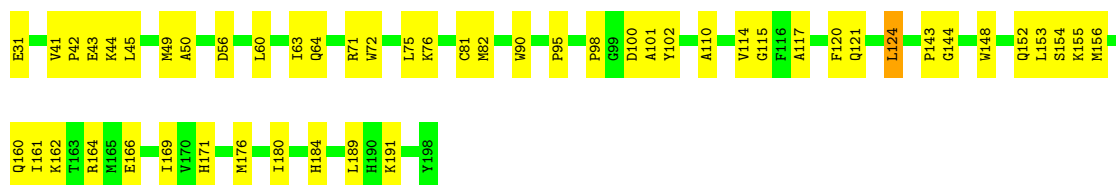
- Molecule 16: Tp-RedCAP

Chain E: 72% 28%



- Molecule 17: Fucoxanthin chl a/c light-harvesting protein

Chain H: 70% 30%



4 Experimental information

Property	Value	Source
EM reconstruction method	SINGLE PARTICLE	Depositor
Imposed symmetry	POINT, Not provided	
Number of particles used	52772	Depositor
Resolution determination method	FSC 0.143 CUT-OFF	Depositor
CTF correction method	PHASE FLIPPING AND AMPLITUDE CORRECTION	Depositor
Microscope	FEI TITAN KRIOS	Depositor
Voltage (kV)	300	Depositor
Electron dose ($e^-/\text{\AA}^2$)	60	Depositor
Minimum defocus (nm)	1000	Depositor
Maximum defocus (nm)	2000	Depositor
Magnification	Not provided	
Image detector	GATAN K3 BIOCONTINUUM (6k x 4k)	Depositor
Maximum map value	1.370	Depositor
Minimum map value	-0.381	Depositor
Average map value	-0.000	Depositor
Map value standard deviation	0.019	Depositor
Recommended contour level	0.08	Depositor
Map size (Å)	563.2, 563.2, 563.2	wwPDB
Map dimensions	512, 512, 512	wwPDB
Map angles (°)	90.0, 90.0, 90.0	wwPDB
Pixel spacing (Å)	1.1, 1.1, 1.1	Depositor

5 Model quality [i](#)

5.1 Standard geometry [i](#)

Bond lengths and bond angles in the following residue types are not validated in this section: CLA, LHG, LMG, BCR, A86, SQD, PQN, ET4, KC1, SF4, DD6, DGD

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	0.53	0/6049	0.67	2/8234 (0.0%)
2	b	0.53	0/6034	0.66	1/8236 (0.0%)
3	c	0.52	0/609	0.80	1/826 (0.1%)
4	d	0.47	0/1067	0.72	0/1441
5	e	0.43	0/511	0.72	0/690
6	f	0.53	0/1271	0.73	3/1727 (0.2%)
7	g	0.43	0/1003	0.68	1/1354 (0.1%)
8	i	0.51	0/264	0.73	0/360
9	j	0.52	0/342	0.78	2/463 (0.4%)
10	l	0.54	0/1123	0.67	1/1523 (0.1%)
11	m	0.56	0/222	0.56	0/300
12	r	0.48	0/704	0.60	0/957
13	B	0.53	0/1324	0.66	0/1804
14	C	0.50	0/1339	0.68	1/1813 (0.1%)
15	D	0.51	0/1304	0.64	0/1771
16	E	0.52	0/1450	0.67	0/1974
17	H	0.51	0/1334	0.65	1/1809 (0.1%)
All	All	0.52	0/25950	0.67	13/35282 (0.0%)

Chiral center outliers are detected by calculating the chiral volume of a chiral center and verifying if the center is modelled as a planar moiety or with the opposite hand. A planarity outlier is detected by checking planarity of atoms in a peptide group, atoms in a mainchain group or atoms of a sidechain that are expected to be planar.

Mol	Chain	#Chirality outliers	#Planarity outliers
4	d	0	1

There are no bond length outliers.

All (13) bond angle outliers are listed below:

Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
2	b	667	ARG	N-CA-C	8.65	120.33	111.07
7	g	156	GLY	N-CA-C	-8.09	103.12	112.50
6	f	58	ALA	CB-CA-C	-6.27	109.33	116.54
1	a	266	GLY	N-CA-C	-6.25	102.80	112.85
1	a	122	VAL	N-CA-C	6.09	122.02	109.34
10	l	110	GLU	N-CA-C	-5.79	97.53	107.61
14	C	112	ILE	CB-CA-C	-5.59	108.43	114.35
6	f	57	GLU	N-CA-C	5.56	118.51	109.06
6	f	178	GLU	N-CA-C	-5.53	104.07	111.87
3	c	64	SER	N-CA-C	-5.37	106.73	113.28
9	j	36	MET	N-CA-C	5.31	117.38	109.25
9	j	35	ASP	N-CA-C	5.22	119.60	113.23
17	H	115	GLY	N-CA-C	5.22	116.91	111.95

There are no chirality outliers.

All (1) planarity outliers are listed below:

Mol	Chain	Res	Type	Group
4	d	96	HIS	Peptide

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	5852	0	5682	113	0
2	b	5824	0	5643	144	0
3	c	599	0	582	19	0
4	d	1040	0	1027	26	0
5	e	503	0	500	18	0
6	f	1242	0	1239	34	0
7	g	981	0	922	40	0
8	i	256	0	259	13	0
9	j	332	0	331	19	0
10	l	1095	0	1116	37	0
11	m	220	0	241	8	0
12	r	683	0	675	26	0
13	B	1285	0	1239	61	0
14	C	1302	0	1243	69	0

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Mol	Chain	Non-H	H(model)	H(added)	Clashes	Symm-Clashes
15	D	1271	0	1227	41	0
16	E	1413	0	1365	47	0
17	H	1300	0	1262	44	0
18	B	469	0	479	33	0
18	C	547	0	472	60	0
18	D	568	0	501	34	0
18	E	462	0	454	21	0
18	H	587	0	583	54	0
18	a	2671	0	2759	134	0
18	b	2492	0	2569	138	0
18	f	175	0	177	3	0
18	i	60	0	59	1	0
18	j	41	0	29	1	0
18	l	159	0	144	10	0
18	r	45	0	33	2	0
19	a	33	0	46	1	0
19	b	33	0	46	4	0
20	B	42	0	54	3	0
20	D	49	0	74	6	0
20	E	91	0	131	5	0
20	H	35	0	43	0	0
20	a	109	0	134	6	0
20	b	49	0	74	5	0
20	j	49	0	74	5	0
21	E	80	0	112	9	0
21	a	200	0	280	31	0
21	b	160	0	224	19	0
21	f	80	0	112	6	0
21	i	80	0	112	14	0
21	j	80	0	112	17	0
21	l	80	0	112	11	0
21	m	40	0	56	5	0
21	r	40	0	56	2	0
22	a	33	0	39	7	0
23	C	75	0	88	1	0
23	D	46	0	61	1	0
23	E	43	0	55	5	0
23	a	33	0	33	2	0
23	j	37	0	42	0	0
23	l	40	0	49	1	0
24	b	8	0	0	2	0
24	c	16	0	0	7	0

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Mol	Chain	Non-H	H(model)	H(added)	Clashes	Symm-Clashes
25	b	60	0	81	0	0
26	B	192	0	0	7	0
26	C	144	0	0	5	0
26	D	144	0	0	3	0
26	E	48	0	0	0	0
26	H	144	0	0	17	0
26	b	48	0	0	4	0
26	m	48	0	0	5	0
26	r	48	0	0	1	0
27	l	42	0	0	0	0
28	B	43	0	0	0	0
28	C	43	0	0	1	0
28	D	43	0	0	3	0
28	E	172	0	0	2	0
28	H	43	0	0	3	0
29	B	45	0	0	5	0
29	C	90	0	0	4	0
29	H	45	0	0	1	0
All	All	36577	0	35112	1120	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 16.

All (1120) 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)
26:H:302:A86:C1	18:H:309:CLA:CMD	2.27	1.12
26:H:302:A86:C1	18:H:309:CLA:HMD1	1.82	1.10
18:a:807:CLA:H202	21:j:105:BCR:H392	1.33	1.05
13:B:165:LYS:CB	18:B:306:CLA:HMD2	1.86	1.05
13:B:165:LYS:HB3	18:B:306:CLA:CMD	1.88	1.02
26:H:302:A86:C	18:H:309:CLA:C1D	2.40	0.98
3:c:62:PHE:HD2	4:d:120:ILE:HG21	1.27	0.98
26:H:302:A86:C	18:H:309:CLA:C2D	2.43	0.95
26:H:302:A86:C2	18:H:309:CLA:HMD1	1.97	0.95
26:H:302:A86:C	18:H:309:CLA:HMD1	1.97	0.95
13:B:165:LYS:HB3	18:B:306:CLA:HMD2	0.96	0.94
1:a:312:TYR:HE1	18:a:842:CLA:HED1	1.29	0.94
18:a:807:CLA:H202	21:j:105:BCR:C39	1.97	0.94
22:a:849:SQD:H272	22:a:849:SQD:H441	1.49	0.94
6:f:182:VAL:HG22	6:f:184:PRO:HD2	1.50	0.93

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:c:14:CYS:SG	24:c:102:SF4:FE3	1.59	0.93
14:C:144:GLY:H	18:C:316:CLA:HBC2	1.32	0.92
26:H:302:A86:C	18:H:309:CLA:CHD	2.49	0.90
18:l:202:CLA:HHC	18:l:202:CLA:HBB1	1.51	0.89
26:B:301:A86:C22	18:B:306:CLA:HAC2	2.04	0.88
26:H:302:A86:C	18:H:309:CLA:CMD	2.51	0.87
18:a:807:CLA:H2	18:a:808:CLA:H2	1.57	0.87
13:B:171:MET:HE1	18:B:307:CLA:HAB	1.56	0.85
10:l:105:LYS:HD3	10:l:115:LYS:HB3	1.58	0.84
18:a:825:CLA:HHC	18:a:825:CLA:HBB1	1.60	0.83
21:a:833:BCR:H271	21:a:852:BCR:H372	1.59	0.83
18:a:803:CLA:H51	18:a:803:CLA:H92	1.60	0.83
19:b:831:PQN:H161	21:b:839:BCR:H401	1.61	0.83
3:c:14:CYS:HG	24:c:102:SF4:FE3	0.92	0.83
18:b:801:CLA:H171	18:b:850:CLA:HMC2	1.61	0.82
18:a:807:CLA:C20	21:j:105:BCR:H392	2.09	0.82
3:c:61:ASP:HB2	5:e:51:ASN:ND2	1.94	0.82
16:E:75:GLY:HA3	28:E:304:DD6:C22	2.10	0.81
18:a:803:CLA:H8	18:a:803:CLA:H2	1.62	0.81
18:b:812:CLA:H51	18:H:314:CLA:HED1	1.62	0.81
5:e:21:ILE:HD12	5:e:61:VAL:HG11	1.63	0.81
18:a:855:CLA:H143	21:l:204:BCR:H24C	1.60	0.81
18:C:317:CLA:HBC2	18:C:317:CLA:HHD	1.61	0.81
14:C:97:PRO:HG2	14:C:102:ALA:HA	1.62	0.79
18:b:829:CLA:HHC	18:b:829:CLA:HBB1	1.63	0.79
20:D:201:LHG:H321	18:D:214:CLA:HMA3	1.63	0.79
2:b:116:PHE:HA	2:b:366:THR:HG22	1.65	0.78
14:C:142:PRO:HG2	18:C:316:CLA:CAA	2.14	0.78
7:g:144:PHE:HA	7:g:175:TRP:HH2	1.49	0.78
10:l:95:ILE:HD11	20:E:317:LHG:H241	1.65	0.78
6:f:127:LYS:HD2	6:f:153:ILE:HG23	1.66	0.77
18:a:807:CLA:C20	21:j:105:BCR:C39	2.62	0.77
14:C:144:GLY:N	18:C:316:CLA:HBC2	1.99	0.77
1:a:312:TYR:CE1	18:a:842:CLA:HED1	2.17	0.77
3:c:62:PHE:CD2	4:d:120:ILE:HG21	2.18	0.77
17:H:42:PRO:HG3	17:H:56:ASP:HB3	1.67	0.77
18:b:848:CLA:HAA2	17:H:63:ILE:HD13	1.67	0.76
18:a:848:CLA:H2	22:a:849:SQD:H281	1.67	0.76
17:H:180:ILE:HD12	18:H:306:CLA:H203	1.68	0.75
21:a:833:BCR:H291	21:a:852:BCR:C37	2.16	0.75
18:b:813:CLA:HBD	12:r:110:VAL:HG21	1.70	0.74

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
13:B:169:LEU:HD11	18:B:306:CLA:CBC	2.17	0.74
17:H:82:MET:HE2	17:H:171:HIS:HB3	1.69	0.74
4:d:33:TRP:HE1	4:d:51:MET:HE3	1.50	0.74
26:H:302:A86:C1	18:H:309:CLA:HMD3	2.15	0.74
18:a:813:CLA:HBC3	18:a:813:CLA:HHD	1.70	0.73
26:C:305:A86:C41	26:C:305:A86:C37	2.66	0.73
26:C:305:A86:C41	18:C:312:CLA:O1A	2.36	0.73
1:a:395:MET:HE2	1:a:606:VAL:HG11	1.70	0.73
18:a:855:CLA:H161	21:l:204:BCR:H402	1.68	0.73
16:E:208:MET:HE3	18:E:316:CLA:H3A	1.71	0.72
18:b:819:CLA:H161	18:H:314:CLA:H91	1.70	0.72
18:a:804:CLA:C10	18:a:807:CLA:H201	2.19	0.72
18:b:840:CLA:H2	18:b:840:CLA:H102	1.71	0.71
1:a:514:VAL:HG22	1:a:524:MET:HG3	1.72	0.71
21:f:201:BCR:HC31	20:j:102:LHG:H241	1.71	0.71
11:m:12:MET:HG3	21:m:101:BCR:H382	1.73	0.71
14:C:198:PRO:HG2	18:C:318:CLA:HAA1	1.72	0.71
21:a:833:BCR:HC7	18:a:836:CLA:H42	1.70	0.71
1:a:659:ILE:HG13	1:a:660:GLN:HG3	1.70	0.71
2:b:37:MET:HE3	2:b:42:LEU:HA	1.73	0.71
18:b:814:CLA:HBB1	18:b:838:CLA:H61	1.73	0.70
12:r:64:LEU:HD21	12:r:133:ILE:HD12	1.70	0.70
26:B:301:A86:C22	18:B:306:CLA:CAC	2.70	0.70
18:B:314:CLA:HMB1	18:B:314:CLA:HBB1	1.73	0.70
26:H:302:A86:C2	18:H:309:CLA:CMD	2.63	0.70
1:a:338:HIS:HE1	20:a:832:LHG:HC12	1.56	0.69
2:b:491:VAL:CG1	18:b:813:CLA:HED2	2.22	0.69
21:b:839:BCR:H403	21:b:839:BCR:H23C	1.74	0.69
18:a:810:CLA:H112	22:a:849:SQD:H383	1.73	0.69
7:g:141:PRO:HB2	7:g:144:PHE:HB2	1.75	0.69
2:b:283:PHE:HE1	18:b:814:CLA:HAB	1.57	0.69
18:a:837:CLA:H91	18:D:213:CLA:H143	1.74	0.68
18:a:802:CLA:HBC3	2:b:580:PHE:HB2	1.74	0.68
26:m:102:A86:C39	21:E:306:BCR:C39	2.72	0.68
18:a:802:CLA:H172	21:f:201:BCR:H14C	1.75	0.68
10:l:6:LYS:HE3	15:D:147:TRP:CD1	2.29	0.67
21:E:306:BCR:H10C	18:E:309:CLA:HBC2	1.75	0.67
18:a:804:CLA:H91	18:a:807:CLA:C20	2.25	0.67
26:C:305:A86:C37	26:C:305:A86:O4	2.42	0.67
14:C:172:GLY:O	14:C:176:MET:HG3	1.95	0.67
18:a:816:CLA:HBB1	21:a:850:BCR:H14C	1.77	0.67

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
21:a:833:BCR:H291	21:a:852:BCR:H371	1.76	0.67
2:b:654:LEU:HB2	18:b:801:CLA:O1A	1.95	0.66
7:g:154:LEU:HD21	7:g:175:TRP:HB2	1.78	0.66
1:a:233:SER:HB3	1:a:236:GLU:HG2	1.76	0.66
1:a:678:PHE:HB2	18:a:841:CLA:O1A	1.95	0.66
6:f:67:LYS:O	6:f:70:VAL:HG22	1.95	0.66
2:b:579:MET:HG3	2:b:709:LEU:HD21	1.76	0.66
21:b:833:BCR:H14C	18:b:845:CLA:HMA1	1.78	0.66
26:D:203:A86:C2	18:D:215:CLA:HBC2	2.26	0.66
10:l:35:PRO:HG3	18:l:203:CLA:HED2	1.78	0.66
18:H:314:CLA:O1D	18:H:314:CLA:HBA2	1.96	0.65
7:g:158:LEU:O	7:g:162:LYS:HB3	1.95	0.65
12:r:86:VAL:HG12	12:r:117:GLU:HB3	1.76	0.65
2:b:567:CYS:SG	24:b:802:SF4:FE1	1.88	0.65
7:g:144:PHE:HA	7:g:175:TRP:CH2	2.30	0.65
7:g:158:LEU:HD21	7:g:174:THR:HG21	1.79	0.65
12:r:132:TRP:CD1	18:r:202:CLA:HAA2	2.31	0.65
21:a:833:BCR:H21C	21:a:852:BCR:H362	1.79	0.65
10:l:105:LYS:HD3	10:l:115:LYS:CB	2.27	0.65
14:C:145:PHE:HA	18:C:312:CLA:HBA1	1.78	0.65
16:E:115:ILE:HD11	16:E:119:ARG:HG2	1.78	0.65
2:b:277:LEU:HG	18:b:813:CLA:HMC3	1.76	0.65
18:a:803:CLA:H193	18:a:803:CLA:H12	1.79	0.65
10:l:133:GLY:HA3	23:l:207:LMG:H161	1.77	0.65
13:B:71:ARG:HA	13:B:74:MET:HE3	1.78	0.65
3:c:51:CYS:SG	24:c:101:SF4:FE1	1.87	0.65
1:a:43:PRO:HG3	6:f:143:ILE:HD13	1.79	0.64
18:l:202:CLA:H2A	18:l:202:CLA:O2D	1.97	0.64
20:B:315:LHG:HC11	20:B:315:LHG:HC41	1.77	0.64
18:a:815:CLA:HAB	18:a:815:CLA:H8	1.79	0.64
18:B:306:CLA:CAD	29:B:313:KC1:OBD	2.45	0.64
13:B:169:LEU:HD11	18:B:306:CLA:HBC2	1.79	0.64
10:l:21:ILE:O	10:l:27:THR:HG21	1.99	0.63
15:D:193:GLN:O	15:D:194:LEU:C	2.40	0.63
17:H:153:LEU:HA	17:H:156:MET:SD	2.37	0.63
18:a:840:CLA:H13	19:b:831:PQN:C19	2.28	0.63
2:b:491:VAL:HG11	18:b:813:CLA:HED2	1.80	0.63
14:C:113:PRO:O	14:C:116:VAL:HG22	1.98	0.63
18:b:817:CLA:H51	18:b:838:CLA:H152	1.81	0.63
17:H:60:LEU:HD23	18:H:306:CLA:H42	1.81	0.63
21:i:102:BCR:H383	21:l:201:BCR:C19	2.29	0.63

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
14:C:195:ILE:HG22	18:C:318:CLA:HAC1	1.80	0.63
6:f:59:ASP:HA	6:f:64:LEU:HD21	1.80	0.63
2:b:715:GLY:O	2:b:719:THR:HG22	1.97	0.63
19:b:831:PQN:H161	21:b:839:BCR:C40	2.27	0.63
7:g:57:TYR:HB3	7:g:67:GLN:HE22	1.63	0.62
26:C:305:A86:C29	18:C:312:CLA:H72	2.30	0.62
17:H:98:PRO:HB2	26:H:302:A86:C41	2.29	0.62
26:B:302:A86:C2	18:B:309:CLA:H12	2.30	0.62
4:d:72:GLN:HG2	4:d:76:PHE:HE2	1.64	0.62
14:C:200:GLY:O	14:C:201:ASN:C	2.42	0.62
18:b:830:CLA:H11	20:b:835:LHG:H242	1.82	0.62
2:b:267:LEU:HB2	2:b:272:MET:HE3	1.82	0.61
12:r:97:LEU:O	12:r:100:GLU:HG3	2.00	0.61
16:E:175:LYS:HD3	16:E:181:GLU:HG2	1.81	0.61
13:B:169:LEU:HD11	18:B:306:CLA:HBC1	1.81	0.61
5:e:1:MET:N	5:e:60:GLU:HG3	2.16	0.61
9:j:31:ARG:HD3	21:j:105:BCR:H332	1.83	0.61
2:b:310:PRO:HB2	20:b:835:LHG:HC32	1.83	0.61
16:E:189:MET:HE2	21:E:305:BCR:H19C	1.83	0.61
2:b:650:LEU:HB3	18:b:801:CLA:H11	1.82	0.61
15:D:82:TRP:CE2	15:D:101:PRO:HG2	2.36	0.61
15:D:102:LEU:O	15:D:105:MET:HG2	2.00	0.61
18:a:806:CLA:H161	18:a:836:CLA:HBB2	1.83	0.60
18:C:312:CLA:HBD	18:C:316:CLA:CBC	2.31	0.60
15:D:83:TYR:O	18:D:216:CLA:HBA1	2.01	0.60
1:a:533:PHE:HA	18:a:825:CLA:HED1	1.82	0.60
21:a:833:BCR:C27	21:a:852:BCR:H372	2.31	0.60
18:D:208:CLA:H51	18:D:209:CLA:H12	1.83	0.60
18:b:823:CLA:HMA3	20:j:102:LHG:H252	1.82	0.60
15:D:108:LEU:HD11	18:D:210:CLA:HAA2	1.83	0.60
18:B:311:CLA:H13	18:E:312:CLA:H93	1.82	0.60
6:f:85:LEU:HD11	9:j:37:LEU:HD11	1.82	0.60
14:C:63:MET:HE2	14:C:63:MET:HA	1.82	0.60
18:a:803:CLA:H8	18:a:803:CLA:C2	2.29	0.60
4:d:40:ILE:HG12	4:d:50:ILE:HG12	1.83	0.60
18:C:312:CLA:HBD	18:C:316:CLA:HBC1	1.82	0.60
21:a:845:BCR:H403	21:a:845:BCR:H23C	1.84	0.60
2:b:546:MET:HE3	2:b:569:ILE:HD13	1.84	0.60
21:a:833:BCR:H291	21:a:852:BCR:H372	1.84	0.59
18:B:307:CLA:H41	18:B:308:CLA:H43	1.82	0.59
21:a:834:BCR:H10C	18:a:842:CLA:H102	1.84	0.59

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
13:B:149:TRP:HZ3	18:B:312:CLA:HMA2	1.68	0.59
18:B:306:CLA:C3D	29:B:313:KC1:OBD	2.51	0.59
14:C:85:TRP:CE2	14:C:186:PRO:HG3	2.38	0.59
18:b:829:CLA:HHC	18:b:829:CLA:CBB	2.32	0.59
1:a:584:CYS:HB3	2:b:666:TRP:HE3	1.68	0.59
7:g:80:VAL:HG21	7:g:91:TYR:HB3	1.84	0.59
26:D:203:A86:C1	18:D:215:CLA:HBC2	2.33	0.59
18:a:804:CLA:H91	18:a:807:CLA:H203	1.85	0.58
2:b:302:GLU:HB3	12:r:52:MET:HG3	1.85	0.58
17:H:110:ALA:O	17:H:114:VAL:HG23	2.03	0.58
1:a:712:PRO:HG2	1:a:716:PRO:HD3	1.85	0.58
1:a:326:LEU:HA	18:a:844:CLA:HED1	1.85	0.58
8:i:28:LEU:HD21	10:l:95:ILE:HG23	1.84	0.58
18:b:807:CLA:H102	21:i:102:BCR:HC21	1.86	0.58
18:b:813:CLA:CAA	12:r:107:GLY:HA3	2.34	0.58
5:e:1:MET:H3	5:e:60:GLU:HG3	1.68	0.58
14:C:51:LEU:HB2	14:C:53:PHE:CE2	2.38	0.58
15:D:76:MET:HE2	15:D:166:GLY:CA	2.32	0.58
14:C:53:PHE:CD2	18:C:307:CLA:H11	2.38	0.58
1:a:144:TRP:HE1	18:a:838:CLA:HED1	1.69	0.58
20:D:201:LHG:H102	18:D:214:CLA:H42	1.85	0.58
16:E:146:THR:H	16:E:159:GLN:HE22	1.51	0.58
1:a:313:ARG:HG3	1:a:319:GLY:C	2.29	0.57
16:E:173:LEU:HD12	16:E:176:MET:SD	2.43	0.57
18:a:818:CLA:HBB1	18:a:818:CLA:HMB3	1.86	0.57
2:b:457:LEU:HD11	6:f:96:VAL:HG13	1.86	0.57
18:b:848:CLA:H2	18:H:307:CLA:H41	1.86	0.57
11:m:23:ARG:HD3	26:m:102:A86:C41	2.35	0.57
14:C:195:ILE:HG23	18:C:317:CLA:HBC3	1.86	0.57
18:C:309:CLA:C1A	18:C:309:CLA:CGA	2.81	0.57
6:f:25:GLU:HB2	6:f:29:LEU:O	2.04	0.57
1:a:119:TRP:CD2	18:a:808:CLA:HED3	2.40	0.57
10:l:45:LEU:HD21	20:B:315:LHG:HC82	1.86	0.57
1:a:701:ILE:HD13	18:a:827:CLA:HMD1	1.87	0.57
3:c:14:CYS:SG	24:c:102:SF4:S4	3.02	0.57
18:f:204:CLA:H3A	21:f:205:BCR:H382	1.87	0.57
2:b:15:ASP:OD1	2:b:16:PRO:HD2	2.04	0.57
2:b:410:MET:HA	2:b:413:HIS:CE1	2.39	0.57
17:H:72:TRP:CE2	17:H:143:PRO:HA	2.39	0.57
4:d:39:GLN:HG3	4:d:51:MET:SD	2.45	0.57
15:D:70:LYS:HD2	18:D:212:CLA:C3D	2.35	0.57

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
18:b:824:CLA:H11	18:b:842:CLA:C9	2.35	0.56
4:d:98:LYS:HE2	7:g:69:GLN:NE2	2.20	0.56
26:H:302:A86:C24	18:H:309:CLA:HMD3	2.34	0.56
1:a:745:LEU:O	1:a:749:ILE:HG12	2.05	0.56
3:c:21:CYS:HA	24:c:101:SF4:S4	2.45	0.56
13:B:146:LEU:HB3	13:B:148:LEU:HG	1.87	0.56
18:C:307:CLA:HMB3	18:C:307:CLA:HBB1	1.86	0.56
15:D:194:LEU:HD13	18:D:217:CLA:CHB	2.35	0.56
26:H:302:A86:C1	18:H:309:CLA:C2D	2.80	0.56
3:c:58:CYS:HA	24:c:102:SF4:S4	2.46	0.56
6:f:167:GLU:HG2	6:f:173:LEU:HB2	1.88	0.56
18:b:819:CLA:HMD2	18:b:837:CLA:H143	1.88	0.56
2:b:414:LYS:HD3	6:f:182:VAL:HG11	1.88	0.56
2:b:651:PHE:HD1	18:b:801:CLA:H62	1.71	0.56
2:b:546:MET:HE2	2:b:559:ASP:HB2	1.88	0.56
18:b:801:CLA:HBB1	18:b:801:CLA:HMB3	1.88	0.56
1:a:430:VAL:HA	1:a:433:HIS:CE1	2.41	0.56
1:a:681:ALA:HB3	18:a:802:CLA:HBB2	1.88	0.56
18:a:803:CLA:H42	18:a:803:CLA:O1A	2.06	0.55
21:a:833:BCR:H23C	21:a:852:BCR:H362	1.87	0.55
18:D:208:CLA:H41	18:D:209:CLA:H12	1.87	0.55
2:b:618:TRP:O	2:b:622:TYR:HB3	2.05	0.55
18:b:813:CLA:HED3	12:r:109:TRP:HZ2	1.71	0.55
21:l:204:BCR:H393	18:D:211:CLA:CMC	2.36	0.55
18:H:305:CLA:HBC1	18:H:313:CLA:H203	1.87	0.55
1:a:575:CYS:SG	24:b:802:SF4:S4	3.04	0.55
12:r:135:ARG:HD2	12:r:136:LYS:HZ2	1.71	0.55
15:D:111:VAL:O	15:D:115:GLN:HG3	2.06	0.55
5:e:1:MET:HE3	5:e:3:ASP:HB2	1.87	0.55
26:H:302:A86:C7	18:H:309:CLA:HBC1	2.36	0.55
2:b:195:HIS:O	2:b:199:PRO:HG2	2.06	0.55
26:b:847:A86:C24	12:r:116:VAL:HG22	2.37	0.55
4:d:9:PHE:O	4:d:52:ARG:HD2	2.06	0.55
14:C:177:ILE:O	14:C:181:GLN:HG2	2.07	0.55
1:a:442:ASN:HD22	2:b:677:LEU:HD21	1.72	0.54
18:a:808:CLA:H11	21:j:101:BCR:H19C	1.88	0.54
18:a:830:CLA:NB	20:a:832:LHG:O4	2.40	0.54
2:b:324:THR:HG21	2:b:402:ASN:OD1	2.07	0.54
14:C:143:VAL:HG12	18:C:316:CLA:C1C	2.36	0.54
1:a:134:ASN:HD21	6:f:60:SER:HB2	1.72	0.54
5:e:32:ILE:HD13	5:e:32:ILE:H	1.71	0.54

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
20:D:201:LHG:H102	18:D:214:CLA:C4	2.38	0.54
21:b:839:BCR:H393	18:b:849:CLA:HMD2	1.90	0.54
18:l:203:CLA:H51	18:l:203:CLA:H92	1.89	0.54
14:C:141:PHE:CZ	18:C:316:CLA:HBB1	2.42	0.54
2:b:286:ALA:HB2	18:b:814:CLA:HBC2	1.88	0.54
2:b:343:ALA:O	2:b:347:THR:HG23	2.06	0.54
18:b:813:CLA:HBA2	12:r:110:VAL:HB	1.90	0.54
13:B:82:PHE:HZ	14:C:198:PRO:HG3	1.72	0.54
16:E:73:MET:HE3	18:E:313:CLA:HMC2	1.89	0.54
2:b:222:THR:HB	2:b:223:PRO:HD3	1.90	0.54
6:f:167:GLU:HG3	6:f:172:ASP:HB2	1.89	0.54
2:b:224:PHE:HE1	18:b:813:CLA:HBB2	1.73	0.54
26:m:102:A86:C12	18:H:307:CLA:HBC1	2.37	0.54
15:D:173:ILE:HG21	28:D:205:DD6:C24	2.37	0.54
20:D:201:LHG:H321	18:D:214:CLA:CMA	2.36	0.54
2:b:173:ARG:NH1	18:b:805:CLA:H203	2.23	0.54
17:H:184:HIS:HB2	18:H:313:CLA:HBC3	1.90	0.54
18:a:848:CLA:C2	22:a:849:SQD:H311	2.38	0.54
18:b:813:CLA:HBD	12:r:110:VAL:CG2	2.37	0.54
21:i:103:BCR:H24C	10:l:93:LEU:CD2	2.38	0.54
13:B:148:LEU:HD12	13:B:148:LEU:H	1.72	0.54
18:a:804:CLA:C9	18:a:807:CLA:H201	2.38	0.54
4:d:35:SER:HB2	4:d:51:MET:HE2	1.88	0.54
13:B:51:LEU:HB2	13:B:53:LEU:HD13	1.88	0.54
13:B:114:TRP:O	13:B:117:ILE:HG22	2.08	0.54
13:B:165:LYS:CB	18:B:306:CLA:CMD	2.65	0.54
1:a:114:SER:HB2	1:a:131:VAL:HG11	1.90	0.54
6:f:80:TYR:OH	9:j:35:ASP:O	2.24	0.53
9:j:22:THR:HA	9:j:25:PHE:CD2	2.43	0.53
14:C:50:PRO:HB2	14:C:51:LEU:HD22	1.89	0.53
14:C:198:PRO:CG	18:C:318:CLA:H3A	2.39	0.53
29:C:308:KC1:OBD	18:C:311:CLA:HBD	2.09	0.53
18:b:823:CLA:H72	9:j:25:PHE:HE1	1.73	0.53
7:g:112:LEU:HD12	7:g:116:LYS:HE3	1.91	0.53
18:a:848:CLA:H2	22:a:849:SQD:H311	1.90	0.53
2:b:298:HIS:HB3	2:b:303:ILE:HD11	1.90	0.53
7:g:162:LYS:O	7:g:166:MET:HG3	2.08	0.53
13:B:128:LYS:HD2	13:B:133:LYS:HD2	1.90	0.53
15:D:108:LEU:HD23	15:D:109:PRO:HD2	1.88	0.53
18:H:311:CLA:CGA	18:H:311:CLA:C3A	2.87	0.53
2:b:285:VAL:HG11	26:b:847:A86:C41	2.38	0.53

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
4:d:72:GLN:HG2	4:d:76:PHE:CE2	2.42	0.53
8:i:29:TYR:O	8:i:32:THR:HG22	2.08	0.53
14:C:88:ALA:HB1	18:C:317:CLA:HMD2	1.90	0.53
1:a:11:LYS:N	1:a:14:GLN:HE21	2.06	0.53
21:b:832:BCR:HC8	26:b:847:A86:O3	2.09	0.53
15:D:82:TRP:CD2	15:D:101:PRO:HG2	2.43	0.53
2:b:187:LEU:HD21	21:b:832:BCR:H383	1.91	0.53
18:b:843:CLA:HMB1	18:b:843:CLA:HBB1	1.91	0.53
7:g:80:VAL:HG22	7:g:81:ALA:H	1.74	0.53
13:B:80:TRP:CE2	13:B:99:PRO:HG2	2.44	0.53
16:E:97:THR:HB	16:E:114:THR:OG1	2.08	0.53
18:a:804:CLA:H91	18:a:807:CLA:H201	1.89	0.53
18:a:815:CLA:H203	18:a:819:CLA:H3A	1.91	0.53
2:b:314:LEU:HD22	18:b:821:CLA:HMD2	1.90	0.53
2:b:391:ILE:HD13	18:b:820:CLA:HED3	1.91	0.53
7:g:137:ALA:O	7:g:141:PRO:HD3	2.08	0.53
16:E:207:SER:HA	23:E:318:LMG:HC3	1.91	0.53
1:a:15:VAL:HG11	18:a:848:CLA:HAA2	1.90	0.53
18:a:802:CLA:H102	19:a:829:PQN:H302	1.90	0.53
18:b:813:CLA:HAA1	12:r:107:GLY:HA3	1.91	0.53
6:f:160:TRP:CG	6:f:161:PRO:HD3	2.43	0.53
14:C:71:GLY:O	14:C:75:GLN:HG3	2.09	0.53
20:a:835:LHG:HC11	10:l:69:ARG:HE	1.74	0.53
5:e:20:GLN:HE21	6:f:180:ILE:HB	1.74	0.53
14:C:106:ILE:HG21	14:C:111:LEU:HD21	1.91	0.53
15:D:188:ASP:OD2	18:D:216:CLA:ND	2.42	0.53
2:b:454:PHE:HZ	20:j:102:LHG:H292	1.74	0.52
18:b:817:CLA:H2A	18:b:817:CLA:HED3	1.91	0.52
18:a:841:CLA:HBB1	18:a:841:CLA:HMB3	1.90	0.52
1:a:14:GLN:HB2	1:a:16:PHE:CZ	2.45	0.52
1:a:575:CYS:SG	2:b:561:PRO:HB3	2.50	0.52
2:b:516:PHE:HA	18:b:844:CLA:HED1	1.91	0.52
26:m:102:A86:C39	21:E:306:BCR:H392	2.39	0.52
1:a:304:LEU:HD11	18:a:814:CLA:H18	1.89	0.52
2:b:493:LEU:O	2:b:494:PRO:C	2.51	0.52
10:l:33:ASN:HD21	15:D:129:GLN:HG2	1.73	0.52
1:a:442:ASN:ND2	2:b:677:LEU:HD21	2.24	0.52
1:a:748:ALA:HA	1:a:751:ILE:HG12	1.91	0.52
18:a:842:CLA:HMB3	18:a:842:CLA:HBB1	1.91	0.52
7:g:158:LEU:CD2	7:g:174:THR:HG21	2.40	0.52
18:B:311:CLA:H171	18:E:312:CLA:H71	1.92	0.52

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
18:b:850:CLA:H193	8:i:17:ILE:HD13	1.91	0.52
13:B:33:GLU:H	13:B:33:GLU:CD	2.18	0.52
7:g:160:TYR:CE1	7:g:164:LEU:HD11	2.45	0.52
13:B:145:TYR:CD2	13:B:146:LEU:HD13	2.45	0.52
18:B:306:CLA:H43	29:B:313:KC1:O1D	2.10	0.52
18:b:801:CLA:H151	18:b:850:CLA:HBC3	1.92	0.52
7:g:51:TYR:CE2	7:g:75:PRO:HG2	2.45	0.52
21:i:103:BCR:H24C	10:l:93:LEU:HD21	1.91	0.52
2:b:2:ALA:HB3	2:b:7:LYS:HA	1.93	0.52
2:b:86:PRO:HB2	2:b:115:THR:CG2	2.40	0.52
13:B:157:GLU:HA	13:B:160:LYS:HE2	1.92	0.52
18:D:208:CLA:H201	18:D:215:CLA:HBC3	1.92	0.52
16:E:66:ILE:HD13	16:E:184:LEU:HD13	1.92	0.52
17:H:153:LEU:HD11	17:H:161:ILE:HG23	1.92	0.52
2:b:229:TRP:HB3	18:b:813:CLA:CMB	2.40	0.51
14:C:198:PRO:CG	18:C:318:CLA:HAA1	2.40	0.51
17:H:180:ILE:HG22	18:H:313:CLA:HBC2	1.92	0.51
14:C:79:TRP:CD2	18:C:309:CLA:HBC2	2.45	0.51
15:D:190:MET:HE3	15:D:191:TYR:CE2	2.45	0.51
21:l:204:BCR:H393	18:D:211:CLA:HMC1	1.93	0.51
16:E:172:ASN:HD21	16:E:181:GLU:HG2	1.74	0.51
18:b:850:CLA:HBB2	8:i:15:GLY:HA3	1.93	0.51
5:e:62:LYS:HD2	5:e:62:LYS:C	2.36	0.51
14:C:79:TRP:HE1	18:C:317:CLA:C4B	2.24	0.51
1:a:338:HIS:CE1	20:a:832:LHG:HC12	2.42	0.51
1:a:385:ALA:H	1:a:752:THR:HG21	1.76	0.51
18:a:802:CLA:CGA	18:a:802:CLA:H3A	2.40	0.51
14:C:150:LYS:NZ	14:C:150:LYS:HB2	2.25	0.51
17:H:81:CYS:SG	28:H:303:DD6:C5	2.99	0.51
2:b:157:GLN:O	2:b:161:ARG:HG3	2.10	0.51
2:b:543:SER:HA	6:f:182:VAL:HG23	1.92	0.51
6:f:182:VAL:HG22	6:f:183:SER:N	2.25	0.51
12:r:67:ILE:HG22	12:r:71:MET:HE3	1.93	0.51
13:B:182:PHE:O	13:B:184:PRO:HD3	2.11	0.51
14:C:89:ARG:NH2	14:C:98:ALA:HB2	2.25	0.51
16:E:63:GLN:HG2	16:E:163:ASN:HD21	1.76	0.51
5:e:32:ILE:HD13	5:e:32:ILE:N	2.26	0.51
18:a:803:CLA:H191	20:a:831:LHG:H182	1.92	0.51
10:l:20:PRO:O	10:l:24:SER:HB3	2.10	0.51
18:H:311:CLA:CGA	18:H:311:CLA:H3A	2.40	0.51
6:f:61:PRO:HB2	6:f:62:PRO:HD3	1.93	0.50

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
13:B:46:LYS:HD2	13:B:46:LYS:C	2.36	0.50
13:B:69:ASN:HD22	13:B:69:ASN:N	2.09	0.50
16:E:56:TRP:HB2	16:E:58:TRP:CD1	2.46	0.50
1:a:260:GLY:O	1:a:263:PRO:HD2	2.12	0.50
2:b:91:ILE:HD11	2:b:113:ASN:ND2	2.26	0.50
21:m:101:BCR:H23C	23:E:318:LMG:H151	1.94	0.50
14:C:102:ALA:O	14:C:106:ILE:HG22	2.11	0.50
14:C:119:GLY:HA3	18:C:311:CLA:HAB	1.94	0.50
1:a:385:ALA:HA	1:a:748:ALA:HB1	1.92	0.50
14:C:114:VAL:O	14:C:117:VAL:HG12	2.11	0.50
14:C:170:MET:HE3	18:C:307:CLA:HMC2	1.92	0.50
15:D:179:GLN:HG2	15:D:185:ASP:O	2.11	0.50
16:E:63:GLN:HG2	16:E:163:ASN:ND2	2.26	0.50
16:E:118:GLU:O	16:E:122:ILE:HG12	2.11	0.50
17:H:100:ASP:O	17:H:101:ALA:HB3	2.11	0.50
21:b:839:BCR:H393	18:b:849:CLA:CMD	2.41	0.50
9:j:37:LEU:HD12	9:j:37:LEU:H	1.76	0.50
18:B:311:CLA:HBA1	20:B:315:LHG:H261	1.91	0.50
21:a:833:BCR:C29	21:a:852:BCR:H372	2.42	0.50
18:b:810:CLA:H13	18:b:837:CLA:HAB	1.93	0.50
7:g:113:ASP:HB3	7:g:116:LYS:HB3	1.93	0.50
13:B:158:GLU:OE2	13:B:162:LYS:HD2	2.12	0.50
16:E:189:MET:HA	16:E:189:MET:HE3	1.94	0.50
18:a:842:CLA:HBC2	18:a:842:CLA:HHD	1.94	0.50
2:b:224:PHE:CE1	18:b:813:CLA:HBB2	2.46	0.50
18:D:208:CLA:CGA	18:D:208:CLA:H3A	2.42	0.50
16:E:38:VAL:HG23	16:E:41:ASP:H	1.76	0.50
16:E:156:TYR:O	16:E:160:MET:HG2	2.12	0.50
1:a:107:ASP:HB3	1:a:111:ILE:HD13	1.93	0.50
2:b:681:HIS:O	2:b:688:ASN:ND2	2.45	0.50
18:b:821:CLA:HAB	18:b:827:CLA:HBB2	1.92	0.50
6:f:143:ILE:HG13	6:f:144:ILE:HG13	1.93	0.50
1:a:289:LEU:HG	1:a:375:PRO:O	2.13	0.49
13:B:114:TRP:HE1	26:B:302:A86:C9	2.25	0.49
18:D:217:CLA:HBC2	18:D:217:CLA:HHD	1.93	0.49
20:a:831:LHG:O10	20:a:831:LHG:H272	2.12	0.49
2:b:40:GLU:CD	2:b:40:GLU:H	2.19	0.49
18:a:808:CLA:H42	21:j:101:BCR:H21C	1.94	0.49
18:b:849:CLA:H122	21:i:103:BCR:H19C	1.95	0.49
11:m:5:PHE:CZ	11:m:9:ILE:HD11	2.47	0.49
14:C:88:ALA:CB	18:C:317:CLA:HMD2	2.42	0.49

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
26:D:203:A86:C24	18:D:215:CLA:HBC2	2.42	0.49
2:b:474:ASN:HD21	2:b:479:SER:HA	1.78	0.49
4:d:39:GLN:O	4:d:51:MET:HG2	2.11	0.49
17:H:152:GLN:HB3	17:H:164:ARG:HH12	1.76	0.49
1:a:189:GLU:H	1:a:189:GLU:CD	2.19	0.49
21:a:833:BCR:H21C	21:a:852:BCR:C36	2.42	0.49
26:m:102:A86:C39	21:E:306:BCR:H393	2.40	0.49
1:a:349:TRP:HB3	18:a:805:CLA:HAC1	1.93	0.49
2:b:229:TRP:HB3	18:b:813:CLA:HMB1	1.94	0.49
5:e:10:ILE:HG22	5:e:12:ARG:H	1.77	0.49
13:B:46:LYS:NZ	13:B:46:LYS:H	2.11	0.49
14:C:139:THR:HG21	14:C:141:PHE:CZ	2.47	0.49
1:a:457:ILE:HG23	18:b:801:CLA:H52	1.94	0.49
15:D:38:VAL:HG23	15:D:65:ARG:HH12	1.77	0.49
1:a:459:ASN:HB3	1:a:471:MET:HE3	1.94	0.49
9:j:2:ASN:HD22	9:j:2:ASN:C	2.20	0.49
13:B:148:LEU:HD11	26:B:304:A86:C17	2.43	0.49
15:D:170:MET:HE1	18:D:208:CLA:HHC	1.94	0.49
16:E:169:VAL:HG22	18:E:309:CLA:OBD	2.12	0.49
1:a:267:GLY:O	18:a:813:CLA:H52	2.12	0.49
1:a:596:LEU:HD12	1:a:729:LEU:HD21	1.95	0.49
21:a:833:BCR:H393	21:a:852:BCR:C37	2.42	0.49
26:B:302:A86:C27	18:C:318:CLA:HBB2	2.43	0.49
17:H:184:HIS:CB	18:H:313:CLA:HBC3	2.42	0.49
28:H:303:DD6:C24	18:H:306:CLA:HAB	2.43	0.49
18:H:314:CLA:HBB1	18:H:314:CLA:HMB1	1.95	0.49
13:B:157:GLU:O	13:B:161:THR:HG23	2.12	0.49
1:a:585:GLN:HA	1:a:590:ASP:OD2	2.13	0.48
6:f:160:TRP:CD1	6:f:161:PRO:HD3	2.48	0.48
14:C:137:SER:HA	14:C:143:VAL:HG23	1.95	0.48
15:D:41:THR:HG23	15:D:43:PRO:HD2	1.95	0.48
18:a:802:CLA:H2	18:a:802:CLA:CMA	2.42	0.48
18:a:807:CLA:HMB2	21:j:101:BCR:H23C	1.95	0.48
2:b:461:TYR:CE1	2:b:475:VAL:HG11	2.48	0.48
4:d:98:LYS:HE2	7:g:69:GLN:CD	2.38	0.48
13:B:158:GLU:O	13:B:162:LYS:HG3	2.13	0.48
15:D:93:PHE:O	15:D:96:GLU:HB2	2.13	0.48
26:H:302:A86:C24	18:H:309:CLA:CMD	2.91	0.48
18:a:802:CLA:HBC2	18:a:802:CLA:HHD	1.95	0.48
2:b:54:LEU:HB3	2:b:148:LEU:HD13	1.94	0.48
2:b:117:SER:HA	18:b:841:CLA:HMA2	1.95	0.48

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:b:437:ILE:HG22	18:b:823:CLA:HAC1	1.96	0.48
2:b:568:ASP:HA	2:b:573:ASP:OD1	2.13	0.48
18:b:805:CLA:HBA1	18:b:805:CLA:H3A	1.52	0.48
4:d:116:ARG:NH2	4:d:118:PHE:HZ	2.12	0.48
1:a:371:MET:HG3	1:a:376:PRO:HG3	1.95	0.48
2:b:685:PRO:O	2:b:686:LEU:HB2	2.13	0.48
9:j:11:ALA:N	9:j:12:PRO:HD2	2.28	0.48
14:C:195:ILE:HG22	18:C:318:CLA:CAC	2.44	0.48
18:a:855:CLA:H43	18:l:203:CLA:H43	1.95	0.48
2:b:320:GLY:O	2:b:324:THR:HG23	2.12	0.48
2:b:466:SER:HA	2:b:477:LEU:HD12	1.95	0.48
7:g:114:LEU:H	7:g:114:LEU:HD22	1.78	0.48
2:b:267:LEU:HB2	2:b:272:MET:CE	2.42	0.48
2:b:490:GLN:HB2	12:r:104:ILE:HD11	1.96	0.48
18:b:818:CLA:H112	21:b:833:BCR:H17C	1.96	0.48
18:b:823:CLA:H72	9:j:25:PHE:CE1	2.49	0.48
13:B:171:MET:CE	18:B:307:CLA:HAB	2.38	0.48
18:C:309:CLA:H111	18:C:309:CLA:H72	1.63	0.48
1:a:80:GLN:HA	1:a:80:GLN:HE21	1.79	0.48
2:b:719:THR:HG23	18:b:803:CLA:O1D	2.14	0.48
18:b:822:CLA:HBB2	21:f:201:BCR:H312	1.95	0.48
14:C:93:CYS:SG	18:C:310:CLA:HBB1	2.53	0.48
1:a:263:PRO:HB2	1:a:272:TYR:CZ	2.48	0.48
3:c:60:THR:O	3:c:61:ASP:C	2.57	0.48
8:i:29:TYR:HB2	20:E:317:LHG:H161	1.96	0.48
13:B:180:ASN:OD1	13:B:189:VAL:HG12	2.13	0.48
1:a:685:MET:HB3	18:a:802:CLA:C1C	2.44	0.48
18:a:838:CLA:H152	21:j:101:BCR:H372	1.95	0.48
2:b:547:PRO:HD2	3:c:62:PHE:CE1	2.49	0.48
18:b:819:CLA:H62	21:b:836:BCR:H331	1.96	0.48
13:B:82:PHE:HB3	13:B:83:PRO:HD3	1.96	0.48
14:C:167:ARG:HA	14:C:170:MET:HE3	1.96	0.48
14:C:201:ASN:C	14:C:201:ASN:HD22	2.22	0.48
18:a:807:CLA:HBB2	21:j:101:BCR:H272	1.95	0.48
18:C:317:CLA:HBB1	18:C:318:CLA:HBC3	1.96	0.48
18:H:306:CLA:HBB1	18:H:306:CLA:HMB3	1.94	0.48
1:a:521:ILE:HG21	1:a:524:MET:HE3	1.95	0.47
18:a:847:CLA:H41	18:a:847:CLA:H61	1.71	0.47
2:b:492:TRP:CZ3	18:b:813:CLA:HED1	2.49	0.47
18:b:829:CLA:H61	18:b:829:CLA:H41	1.70	0.47
9:j:25:PHE:O	9:j:29:VAL:HG23	2.14	0.47

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
16:E:38:VAL:O	16:E:39:PHE:HB2	2.14	0.47
18:a:815:CLA:H151	18:a:815:CLA:HBB2	1.95	0.47
2:b:386:PHE:HZ	18:b:845:CLA:HAB	1.79	0.47
2:b:640:ASN:HD21	2:b:642:LEU:HB2	1.78	0.47
18:r:202:CLA:HBA2	18:r:202:CLA:H3A	1.49	0.47
1:a:682:PHE:HZ	18:a:828:CLA:HBC2	1.78	0.47
18:a:803:CLA:H193	18:a:803:CLA:C1	2.44	0.47
2:b:463:GLN:HG2	2:b:508:PHE:CD1	2.49	0.47
18:b:827:CLA:HHB	18:b:845:CLA:HBB1	1.96	0.47
7:g:117:ALA:O	7:g:120:THR:HG22	2.15	0.47
13:B:156:GLN:O	13:B:160:LYS:HG3	2.15	0.47
14:C:108:THR:O	14:C:112:ILE:HG12	2.14	0.47
17:H:114:VAL:HG21	18:H:308:CLA:HAA2	1.96	0.47
18:a:803:CLA:CBB	21:j:105:BCR:H292	2.44	0.47
18:a:809:CLA:H61	18:a:809:CLA:H41	1.71	0.47
2:b:123:TRP:HA	2:b:126:ILE:HG12	1.94	0.47
2:b:635:ASN:HB2	2:b:636:PRO:HD2	1.97	0.47
18:b:841:CLA:H3A	18:b:841:CLA:HBA2	1.35	0.47
11:m:9:ILE:HA	11:m:12:MET:HE3	1.96	0.47
12:r:60:PRO:O	12:r:63:VAL:HG22	2.14	0.47
14:C:141:PHE:CE2	18:C:316:CLA:HBB1	2.50	0.47
16:E:216:VAL:HG21	16:E:220:TYR:CE2	2.49	0.47
18:a:815:CLA:H3A	18:a:815:CLA:HBA2	1.52	0.47
21:a:850:BCR:H15C	21:a:850:BCR:H351	1.77	0.47
18:b:804:CLA:HHC	18:b:806:CLA:OBD	2.14	0.47
18:b:821:CLA:HBA2	18:b:821:CLA:H3A	1.47	0.47
12:r:113:TYR:O	12:r:117:GLU:HG3	2.14	0.47
28:H:303:DD6:C13	28:H:303:DD6:C21	2.92	0.47
2:b:543:SER:HA	6:f:182:VAL:CG2	2.44	0.47
2:b:707:VAL:HA	18:b:829:CLA:HED1	1.96	0.47
5:e:28:ASP:HB3	5:e:32:ILE:HD11	1.96	0.47
13:B:124:TRP:CZ2	13:B:128:LYS:HE2	2.49	0.47
16:E:206:GLN:NE2	16:E:211:THR:HG22	2.29	0.47
1:a:460:ASP:OD2	1:a:642:THR:HB	2.15	0.47
21:a:850:BCR:H322	21:a:852:BCR:H21C	1.96	0.47
2:b:242:VAL:HG12	2:b:247:GLU:HB2	1.96	0.47
2:b:349:GLN:HG3	18:b:845:CLA:HED1	1.95	0.47
2:b:663:LEU:HD22	2:b:708:GLY:HA3	1.97	0.47
4:d:85:ARG:HB2	4:d:95:LEU:HD11	1.96	0.47
10:l:23:SER:HB2	10:l:28:ARG:NH1	2.30	0.47
11:m:29:TYR:CD1	20:E:301:LHG:HC61	2.50	0.47

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
21:m:101:BCR:H272	23:E:318:LMG:H121	1.97	0.47
13:B:192:PHE:CE1	13:B:193:ILE:HG13	2.49	0.47
14:C:96:PHE:CD1	14:C:96:PHE:N	2.81	0.47
16:E:52:GLN:CD	16:E:52:GLN:H	2.22	0.47
16:E:149:LEU:HD13	16:E:160:MET:SD	2.54	0.47
26:H:302:A86:C	18:H:309:CLA:HHD	2.39	0.47
4:d:62:ARG:HB2	4:d:65:GLN:HG3	1.96	0.47
18:D:208:CLA:H91	18:D:208:CLA:H112	1.51	0.47
1:a:412:PHE:CD1	1:a:416:ASP:HB2	2.50	0.47
18:a:813:CLA:H102	23:a:851:LMG:H311	1.97	0.47
18:a:841:CLA:O1A	18:a:841:CLA:H2	2.15	0.47
2:b:74:PHE:O	2:b:78:VAL:HG23	2.15	0.47
2:b:323:GLU:HB2	2:b:401:LEU:HD21	1.97	0.47
20:b:835:LHG:H332	20:b:835:LHG:H361	1.69	0.47
5:e:18:PHE:HB3	6:f:180:ILE:HD12	1.97	0.47
6:f:96:VAL:HG11	6:f:100:ARG:HD2	1.96	0.47
7:g:132:LYS:HD2	7:g:171:LEU:HD11	1.97	0.47
13:B:60:ASP:O	13:B:64:ARG:HG3	2.14	0.47
29:C:308:KC1:CED	18:C:311:CLA:H51	2.45	0.47
18:D:207:CLA:H61	18:D:207:CLA:H41	1.51	0.47
18:H:310:CLA:HBB1	18:H:310:CLA:HMB3	1.97	0.47
2:b:690:ILE:HG23	10:l:96:TYR:CE1	2.49	0.47
18:b:813:CLA:H41	26:b:847:A86:C22	2.45	0.47
13:B:144:ASP:OD1	13:B:148:LEU:HD12	2.15	0.47
18:B:311:CLA:H62	18:B:311:CLA:H41	1.54	0.47
14:C:179:HIS:HE1	18:C:314:CLA:NA	2.12	0.47
18:a:803:CLA:H202	18:a:803:CLA:H161	1.65	0.46
2:b:157:GLN:HB3	2:b:158:PRO:HD2	1.98	0.46
2:b:328:SER:O	2:b:332:GLN:HG3	2.14	0.46
18:b:845:CLA:H142	18:b:845:CLA:H111	1.81	0.46
6:f:109:LEU:N	6:f:110:PRO:HD2	2.30	0.46
15:D:71:ASN:HD22	15:D:71:ASN:N	2.12	0.46
17:H:176:MET:HE3	18:H:306:CLA:HMC3	1.96	0.46
18:H:311:CLA:C2B	18:H:311:CLA:H43	2.44	0.46
1:a:698:ILE:O	1:a:702:VAL:HG23	2.15	0.46
2:b:496:TRP:HD1	2:b:497:LEU:HD12	1.80	0.46
2:b:651:PHE:CE2	2:b:655:ILE:HD11	2.50	0.46
18:b:809:CLA:H12	17:H:60:LEU:HD11	1.97	0.46
18:b:824:CLA:HBC2	18:b:824:CLA:HHD	1.98	0.46
16:E:172:ASN:HD21	16:E:181:GLU:CG	2.28	0.46
17:H:75:LEU:HD13	17:H:144:GLY:HA3	1.96	0.46

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:a:227:LEU:HD21	1:a:253:LEU:HD21	1.97	0.46
6:f:146:ASN:OD1	6:f:148:PRO:HD2	2.14	0.46
13:B:177:PHE:CE1	13:B:193:ILE:HG21	2.50	0.46
18:D:208:CLA:H52	18:D:208:CLA:CMB	2.46	0.46
18:a:844:CLA:HMB2	21:a:845:BCR:H362	1.98	0.46
4:d:32:THR:HG22	4:d:57:LEU:HD13	1.97	0.46
14:C:119:GLY:CA	18:C:311:CLA:HAB	2.45	0.46
17:H:180:ILE:HD12	18:H:306:CLA:C20	2.43	0.46
1:a:29:TRP:HE1	18:a:809:CLA:CHB	2.28	0.46
1:a:248:ASP:O	1:a:252:GLN:HG3	2.15	0.46
18:a:827:CLA:HAA2	18:b:843:CLA:HMB1	1.98	0.46
18:a:854:CLA:CBA	18:a:854:CLA:HED3	2.45	0.46
2:b:242:VAL:HG13	2:b:245:THR:OG1	2.15	0.46
18:b:817:CLA:H3A	18:b:817:CLA:HBA2	1.61	0.46
3:c:13:GLY:HA3	3:c:38:GLN:HG3	1.98	0.46
13:B:43:GLU:H	13:B:43:GLU:HG2	1.38	0.46
13:B:159:MET:HG3	18:B:312:CLA:HMA1	1.96	0.46
14:C:106:ILE:HG13	14:C:110:ASN:OD1	2.15	0.46
18:D:216:CLA:H3A	18:D:216:CLA:HBA2	1.58	0.46
18:a:839:CLA:H191	18:a:841:CLA:H141	1.98	0.46
22:a:849:SQD:H262	22:a:849:SQD:H291	1.54	0.46
2:b:256:PHE:CE2	18:b:842:CLA:HBB1	2.51	0.46
18:b:823:CLA:HBA1	18:b:823:CLA:H3A	1.65	0.46
2:b:256:PHE:HZ	18:b:842:CLA:H62	1.80	0.46
18:b:822:CLA:CBB	21:f:201:BCR:H312	2.46	0.46
18:b:823:CLA:H61	18:b:823:CLA:H41	1.79	0.46
18:b:824:CLA:H93	18:b:825:CLA:HBB1	1.98	0.46
21:b:839:BCR:H11C	21:b:839:BCR:H341	1.61	0.46
7:g:84:SER:OG	7:g:90:SER:HB3	2.16	0.46
10:l:91:LEU:O	10:l:95:ILE:HG12	2.16	0.46
13:B:152:ASP:O	13:B:156:GLN:HG3	2.16	0.46
14:C:167:ARG:HA	14:C:170:MET:CE	2.45	0.46
18:C:314:CLA:HHD	18:C:314:CLA:HBC2	1.96	0.46
18:D:215:CLA:HED3	18:D:215:CLA:H2A	1.97	0.46
17:H:72:TRP:NE1	17:H:143:PRO:HA	2.31	0.46
2:b:173:ARG:HH21	18:b:838:CLA:HMD2	1.81	0.46
18:b:801:CLA:H92	18:b:801:CLA:H61	1.76	0.46
18:b:843:CLA:C4A	18:b:843:CLA:HBA2	2.46	0.46
3:c:38:GLN:HB3	4:d:106:VAL:HG21	1.97	0.46
20:D:201:LHG:H331	20:D:201:LHG:H301	1.58	0.46
18:a:810:CLA:H202	21:a:833:BCR:H382	1.97	0.46

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
18:a:837:CLA:C9	18:D:213:CLA:H143	2.44	0.46
18:b:825:CLA:HBA2	18:b:842:CLA:H141	1.98	0.46
21:j:101:BCR:H15C	21:j:101:BCR:H351	1.77	0.46
10:l:110:GLU:O	10:l:111:LEU:C	2.59	0.46
18:D:207:CLA:H111	18:D:207:CLA:H91	1.42	0.46
16:E:39:PHE:HD1	18:E:309:CLA:HBA2	1.80	0.46
21:E:306:BCR:H24C	21:E:306:BCR:H371	1.76	0.46
18:E:312:CLA:CBB	18:E:316:CLA:HBC2	2.46	0.46
18:E:316:CLA:HBA1	23:E:318:LMG:O2	2.16	0.46
18:a:818:CLA:HMA1	18:a:830:CLA:CBC	2.46	0.46
18:f:203:CLA:HBB1	9:j:26:ILE:HD11	1.97	0.46
18:H:311:CLA:H3A	18:H:311:CLA:HBA2	1.75	0.46
1:a:623:THR:HB	1:a:624:PRO:HD2	1.98	0.45
1:a:695:GLN:O	1:a:699:GLU:HG3	2.17	0.45
18:a:821:CLA:H2A	18:a:821:CLA:HED2	1.97	0.45
18:b:801:CLA:OBD	18:b:803:CLA:HMB3	2.17	0.45
18:b:810:CLA:H193	18:b:814:CLA:H171	1.97	0.45
18:b:840:CLA:H43	18:b:841:CLA:H11	1.98	0.45
4:d:130:LYS:HG2	4:d:131:PHE:CD2	2.51	0.45
4:d:137:TYR:CD1	4:d:138:GLU:HG3	2.51	0.45
7:g:151:MET:O	7:g:155:VAL:HG22	2.16	0.45
18:a:803:CLA:H93	18:a:827:CLA:C9	2.46	0.45
13:B:46:LYS:H	13:B:46:LYS:HZ2	1.64	0.45
14:C:125:TRP:CZ3	14:C:141:PHE:HE2	2.34	0.45
1:a:320:HIS:HB3	1:a:325:ILE:HD11	1.98	0.45
21:a:833:BCR:H393	21:a:852:BCR:H373	1.98	0.45
18:a:837:CLA:H101	18:a:855:CLA:HBC1	1.98	0.45
18:b:814:CLA:H72	18:b:838:CLA:H122	1.97	0.45
18:b:817:CLA:H71	18:b:817:CLA:H111	1.67	0.45
13:B:127:LYS:O	13:B:131:GLU:HG3	2.17	0.45
16:E:62:VAL:O	16:E:66:ILE:HG12	2.16	0.45
17:H:102:TYR:HA	18:H:308:CLA:HED2	1.98	0.45
1:a:396:TRP:O	1:a:397:ILE:C	2.59	0.45
1:a:688:PHE:CD1	2:b:663:LEU:HB3	2.51	0.45
12:r:100:GLU:OE1	12:r:104:ILE:HG23	2.16	0.45
18:b:817:CLA:H62	18:b:817:CLA:H2	1.65	0.45
18:b:849:CLA:H62	18:b:849:CLA:H41	1.62	0.45
13:B:124:TRP:CH2	13:B:128:LYS:HE2	2.51	0.45
13:B:152:ASP:OD1	13:B:152:ASP:N	2.50	0.45
14:C:132:PHE:HZ	18:C:312:CLA:HED2	1.81	0.45
14:C:198:PRO:HD2	18:C:318:CLA:HAA1	1.97	0.45

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
18:C:317:CLA:NA	18:C:318:CLA:HMD1	2.32	0.45
8:i:18:PHE:N	8:i:19:PRO:HD2	2.32	0.45
16:E:169:VAL:HG13	18:E:309:CLA:HMD3	1.97	0.45
18:H:311:CLA:H102	18:H:311:CLA:H62	1.50	0.45
1:a:226:LYS:HD2	1:a:253:LEU:HD22	1.99	0.45
2:b:5:PHE:HB3	2:b:6:PRO:HD3	1.99	0.45
2:b:439:ASN:HB3	2:b:451:GLN:HE21	1.80	0.45
18:b:803:CLA:H111	18:b:803:CLA:H143	1.48	0.45
18:b:850:CLA:H3A	18:b:850:CLA:HBA1	1.74	0.45
7:g:84:SER:HB3	7:g:85:PRO:HD2	1.99	0.45
18:C:312:CLA:H111	18:C:312:CLA:H142	1.68	0.45
17:H:42:PRO:HG2	17:H:45:LEU:HG	1.98	0.45
1:a:17:VAL:HG11	1:a:184:ALA:HB1	1.98	0.45
1:a:509:ALA:HB2	1:a:523:MET:CE	2.47	0.45
18:a:806:CLA:HED3	18:a:820:CLA:H72	1.98	0.45
21:b:839:BCR:H24C	21:b:839:BCR:H371	1.79	0.45
3:c:62:PHE:O	3:c:63:LEU:C	2.59	0.45
1:a:299:LEU:HD21	18:a:813:CLA:HAB	1.98	0.45
4:d:124:PRO:HG3	4:d:139:SER:HB3	1.97	0.45
29:B:313:KC1:CHC	29:B:313:KC1:CBB	2.94	0.45
23:C:319:LMG:H111	23:C:319:LMG:H142	1.67	0.45
20:E:301:LHG:H371	20:E:301:LHG:H342	1.48	0.45
17:H:148:TRP:N	17:H:148:TRP:CD1	2.84	0.45
17:H:189:LEU:HB2	17:H:191:LYS:HG2	1.98	0.45
1:a:448:LEU:HD12	1:a:448:LEU:HA	1.82	0.45
1:a:641:ILE:HG13	1:a:642:THR:HG23	1.99	0.45
1:a:694:TRP:O	1:a:698:ILE:HG13	2.17	0.45
18:b:806:CLA:H2	18:b:806:CLA:H62	1.60	0.45
20:b:835:LHG:H301	20:b:835:LHG:H271	1.37	0.45
21:b:839:BCR:H15C	21:b:839:BCR:H351	1.82	0.45
7:g:149:ALA:O	7:g:153:GLU:HG3	2.17	0.45
8:i:26:PHE:CD2	21:i:103:BCR:HC8	2.52	0.45
10:l:26:LEU:HG	15:D:141:LEU:O	2.17	0.45
14:C:198:PRO:CD	18:C:318:CLA:HAA1	2.46	0.45
18:C:309:CLA:H102	18:C:309:CLA:H13	1.75	0.45
16:E:46:TRP:HZ3	16:E:54:ALA:HB2	1.82	0.45
16:E:70:ARG:HA	16:E:73:MET:CE	2.46	0.45
17:H:162:LYS:O	17:H:166:GLU:HG3	2.16	0.45
1:a:337:GLY:O	1:a:426:LEU:HG	2.17	0.44
18:a:814:CLA:HBA2	18:a:814:CLA:H3A	1.55	0.44
2:b:93:ASP:HB3	2:b:96:PHE:CD1	2.51	0.44

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
18:b:813:CLA:HAA2	12:r:107:GLY:HA3	1.99	0.44
7:g:173:LYS:HD2	7:g:173:LYS:C	2.42	0.44
21:l:204:BCR:H341	21:l:204:BCR:H11C	1.60	0.44
1:a:521:ILE:HD13	1:a:524:MET:CE	2.47	0.44
6:f:85:LEU:HD11	9:j:37:LEU:CD1	2.47	0.44
18:i:101:CLA:H101	18:i:101:CLA:H13	1.84	0.44
18:H:308:CLA:H91	18:H:308:CLA:H111	1.60	0.44
18:a:815:CLA:H122	18:a:815:CLA:H162	1.74	0.44
21:a:852:BCR:H15C	21:a:852:BCR:H351	1.63	0.44
18:b:801:CLA:H171	18:b:850:CLA:CMC	2.40	0.44
21:b:833:BCR:H15C	21:b:833:BCR:H351	1.68	0.44
21:i:103:BCR:H15C	21:i:103:BCR:H351	1.74	0.44
20:j:102:LHG:H182	20:j:102:LHG:H212	1.78	0.44
21:l:201:BCR:H15C	21:l:201:BCR:H351	1.79	0.44
13:B:149:TRP:CE3	18:B:312:CLA:HAA2	2.52	0.44
17:H:120:PHE:CZ	17:H:124:LEU:HD11	2.51	0.44
18:a:802:CLA:CBC	2:b:580:PHE:HB2	2.43	0.44
21:a:833:BCR:H24C	21:a:833:BCR:H371	1.78	0.44
2:b:354:LEU:HD11	18:b:818:CLA:HAB	1.99	0.44
2:b:465:ALA:HB1	2:b:476:LEU:HD12	1.99	0.44
18:b:846:CLA:H141	18:b:846:CLA:H161	1.82	0.44
14:C:117:VAL:HG11	28:C:303:DD6:C5	2.46	0.44
16:E:172:ASN:ND2	16:E:181:GLU:HG2	2.32	0.44
18:E:312:CLA:H112	18:E:312:CLA:H142	1.71	0.44
1:a:85:PHE:HE2	1:a:173:MET:HE3	1.83	0.44
18:a:813:CLA:H3A	18:a:813:CLA:HBA2	1.49	0.44
18:a:825:CLA:H12	18:a:825:CLA:NA	2.32	0.44
21:a:845:BCR:H24C	21:a:845:BCR:H371	1.79	0.44
18:b:845:CLA:H121	18:b:845:CLA:H161	1.71	0.44
15:D:78:ALA:HB1	28:D:205:DD6:C37	2.46	0.44
16:E:75:GLY:O	16:E:76:TRP:C	2.60	0.44
1:a:359:MET:HE2	1:a:359:MET:HA	1.99	0.44
8:i:33:ASP:OD1	8:i:33:ASP:N	2.51	0.44
10:l:6:LYS:HE3	15:D:147:TRP:HD1	1.80	0.44
13:B:158:GLU:HG3	13:B:159:MET:N	2.32	0.44
15:D:146:TYR:HD2	15:D:147:TRP:CE2	2.36	0.44
2:b:283:PHE:CE1	18:b:814:CLA:HAB	2.46	0.44
18:b:846:CLA:H142	18:b:846:CLA:H112	1.85	0.44
7:g:113:ASP:H	7:g:116:LYS:HE2	1.83	0.44
14:C:112:ILE:N	14:C:113:PRO:HD2	2.32	0.44
14:C:179:HIS:CE1	18:C:314:CLA:NA	2.86	0.44

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
15:D:72:GLY:O	15:D:76:MET:HG3	2.18	0.44
15:D:170:MET:HE2	28:D:205:DD6:C8	2.47	0.44
18:D:208:CLA:H92	18:D:208:CLA:H61	1.54	0.44
1:a:222:LEU:N	1:a:223:PRO:HD2	2.33	0.44
1:a:262:ALA:HB3	1:a:263:PRO:HD3	1.98	0.44
1:a:469:GLN:CD	1:a:469:GLN:H	2.25	0.44
1:a:702:VAL:HG22	18:f:202:CLA:HBC2	1.99	0.44
18:a:826:CLA:H111	18:a:844:CLA:H12	1.98	0.44
2:b:720:TYR:HB2	18:b:803:CLA:HED3	2.00	0.44
7:g:89:THR:HG23	7:g:94:ALA:HB3	1.99	0.44
18:l:202:CLA:H3A	18:l:202:CLA:HBA1	1.80	0.44
13:B:71:ARG:HA	13:B:74:MET:CE	2.47	0.44
17:H:166:GLU:O	17:H:169:ILE:HG22	2.17	0.44
1:a:326:LEU:O	1:a:338:HIS:HB2	2.18	0.44
1:a:673:PHE:CE1	1:a:739:THR:HG23	2.53	0.44
18:a:806:CLA:H12	18:a:820:CLA:H2	1.99	0.44
3:c:12:ILE:HG12	24:c:102:SF4:S2	2.58	0.44
18:D:217:CLA:H2A	18:D:217:CLA:HED2	2.00	0.44
18:a:807:CLA:H201	21:j:105:BCR:C39	2.47	0.43
2:b:607:GLN:OE1	2:b:621:ASP:HB3	2.18	0.43
10:l:54:HIS:HA	10:l:57:PHE:CE2	2.53	0.43
13:B:187:CYS:SG	18:B:314:CLA:C4B	3.06	0.43
18:E:315:CLA:H62	18:E:315:CLA:H92	1.84	0.43
1:a:334:THR:HB	1:a:426:LEU:HD21	2.01	0.43
1:a:596:LEU:CD1	1:a:729:LEU:HD21	2.47	0.43
18:a:801:CLA:H203	18:a:801:CLA:H162	1.58	0.43
21:a:833:BCR:H402	18:a:847:CLA:H143	2.00	0.43
18:a:841:CLA:H62	18:a:841:CLA:H41	1.52	0.43
18:b:840:CLA:H142	18:b:840:CLA:H112	1.79	0.43
6:f:182:VAL:HG22	6:f:183:SER:H	1.81	0.43
18:E:316:CLA:H3A	18:E:316:CLA:HBA2	1.79	0.43
1:a:11:LYS:H	1:a:11:LYS:HG2	1.63	0.43
18:a:808:CLA:H151	18:a:809:CLA:H142	2.00	0.43
18:a:823:CLA:HBA2	18:a:823:CLA:H3A	1.57	0.43
2:b:88:ALA:HB2	2:b:115:THR:HB	1.99	0.43
2:b:234:GLU:HG2	12:r:106:ASN:ND2	2.33	0.43
2:b:253:ILE:HG13	2:b:254:LEU:HG	1.99	0.43
13:B:34:LEU:HB3	18:B:307:CLA:HED3	2.00	0.43
18:B:312:CLA:HBA2	18:B:312:CLA:H3A	1.45	0.43
18:C:309:CLA:H2A	18:C:309:CLA:O2D	2.18	0.43
18:D:207:CLA:H71	18:D:214:CLA:C1D	2.48	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
16:E:138:ILE:C	16:E:140:PRO:HD3	2.42	0.43
21:b:836:BCR:H361	21:b:836:BCR:H20C	1.78	0.43
18:b:843:CLA:H41	18:b:843:CLA:H62	1.41	0.43
13:B:128:LYS:HD3	13:B:128:LYS:HA	1.74	0.43
14:C:112:ILE:O	14:C:113:PRO:C	2.62	0.43
29:C:308:KC1:CAD	18:C:311:CLA:HBD	2.48	0.43
15:D:100:ASP:HB3	15:D:103:LYS:HD3	1.99	0.43
18:a:854:CLA:H61	18:a:854:CLA:H41	1.61	0.43
5:e:32:ILE:H	5:e:32:ILE:CD1	2.32	0.43
8:i:28:LEU:CD2	10:l:95:ILE:HG23	2.48	0.43
20:j:102:LHG:H161	20:j:102:LHG:H191	1.79	0.43
10:l:30:LEU:O	10:l:34:LEU:HG	2.18	0.43
12:r:82:ALA:O	12:r:86:VAL:HG23	2.18	0.43
13:B:149:TRP:CE3	13:B:150:PRO:HD2	2.53	0.43
13:B:159:MET:HG3	18:B:312:CLA:CMA	2.48	0.43
18:B:306:CLA:HBA2	18:B:306:CLA:H3A	1.16	0.43
18:E:311:CLA:O1A	18:E:312:CLA:H203	2.19	0.43
21:a:833:BCR:C7	18:a:836:CLA:H42	2.45	0.43
2:b:321:LEU:HG	2:b:405:ASN:HD21	1.83	0.43
18:b:829:CLA:H18	21:i:102:BCR:H362	1.99	0.43
21:b:836:BCR:H15C	21:b:836:BCR:H351	1.91	0.43
10:l:105:LYS:HD2	10:l:113:THR:HB	2.00	0.43
21:l:204:BCR:C8	21:l:204:BCR:H331	2.48	0.43
11:m:16:LEU:O	11:m:19:VAL:HG22	2.18	0.43
18:B:314:CLA:HBB1	18:B:314:CLA:CMB	2.45	0.43
21:E:306:BCR:H20C	21:E:306:BCR:H361	1.77	0.43
1:a:585:GLN:HG2	1:a:725:ARG:CZ	2.48	0.43
18:a:802:CLA:HMB1	18:a:802:CLA:HBB1	2.01	0.43
18:a:837:CLA:C15	18:a:844:CLA:H192	2.49	0.43
2:b:321:LEU:HD11	2:b:406:VAL:HG12	1.99	0.43
2:b:440:ASP:OD1	2:b:614:TYR:HB2	2.18	0.43
20:b:835:LHG:H161	20:b:835:LHG:H132	1.35	0.43
7:g:85:PRO:HG2	7:g:88:GLU:CD	2.43	0.43
7:g:136:PHE:HE1	7:g:172:MET:HA	1.84	0.43
10:l:106:SER:O	10:l:107:ASN:C	2.60	0.43
21:l:204:BCR:H17C	23:D:202:LMG:H211	2.01	0.43
14:C:106:ILE:HG13	14:C:110:ASN:HD21	1.83	0.43
21:E:305:BCR:H11C	21:E:305:BCR:H341	1.83	0.43
17:H:100:ASP:C	17:H:102:TYR:H	2.25	0.43
1:a:94:HIS:HA	1:a:98:PHE:HD2	1.82	0.43
1:a:597:PHE:CE2	1:a:729:LEU:HD13	2.54	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:c:51:CYS:O	3:c:52:LYS:HB2	2.18	0.43
14:C:115:LEU:HD22	14:C:115:LEU:H	1.84	0.43
17:H:76:LYS:HD2	18:H:310:CLA:C3D	2.49	0.43
17:H:102:TYR:CE1	17:H:114:VAL:HG13	2.54	0.43
1:a:691:ARG:HG3	1:a:719:LEU:O	2.19	0.43
21:a:845:BCR:H15C	21:a:845:BCR:H351	1.87	0.43
18:a:854:CLA:H93	18:a:854:CLA:H62	1.52	0.43
2:b:82:LEU:CD2	2:b:362:LYS:HD3	2.49	0.43
7:g:136:PHE:HA	7:g:168:THR:HG23	2.01	0.43
8:i:4:SER:O	8:i:7:PRO:HD2	2.18	0.43
16:E:91:ILE:HD13	16:E:118:GLU:HG3	2.00	0.43
16:E:166:LEU:HD11	16:E:181:GLU:HA	2.01	0.43
1:a:291:LEU:HD23	1:a:294:ILE:HD12	2.00	0.43
18:a:825:CLA:H62	18:a:825:CLA:H41	1.64	0.43
2:b:667:ARG:O	2:b:668:GLY:C	2.61	0.43
18:b:820:CLA:H143	18:b:820:CLA:H111	1.82	0.43
18:b:842:CLA:H62	18:b:842:CLA:H2	1.73	0.43
5:e:4:ARG:O	5:e:5:ASN:HB2	2.19	0.43
9:j:22:THR:HA	9:j:25:PHE:HD2	1.82	0.43
12:r:92:VAL:HG12	12:r:96:PHE:CE2	2.54	0.43
12:r:111:ARG:HB3	12:r:113:TYR:CZ	2.54	0.43
15:D:153:GLU:O	15:D:156:VAL:HG22	2.19	0.43
15:D:170:MET:O	15:D:173:ILE:HG22	2.19	0.43
17:H:31:GLU:O	17:H:41:VAL:HG12	2.18	0.43
18:H:313:CLA:H93	18:H:313:CLA:H61	1.83	0.43
1:a:87:TRP:O	1:a:91:MET:HG2	2.19	0.42
18:a:802:CLA:H201	18:a:828:CLA:H191	1.99	0.42
2:b:173:ARG:HH12	18:b:805:CLA:H203	1.82	0.42
2:b:425:SER:HB3	2:b:528:VAL:HG22	2.01	0.42
2:b:536:GLY:O	2:b:540:ALA:HB2	2.19	0.42
4:d:28:LYS:HE2	4:d:59:TYR:HB3	2.01	0.42
10:l:121:LYS:HE2	10:l:121:LYS:HB3	1.88	0.42
18:C:312:CLA:H3A	18:C:312:CLA:HBA2	1.54	0.42
1:a:509:ALA:HB2	1:a:523:MET:HE3	2.02	0.42
1:a:578:PRO:HB3	2:b:558:CYS:SG	2.59	0.42
18:a:813:CLA:H142	18:a:813:CLA:H111	1.75	0.42
18:b:807:CLA:H91	18:b:807:CLA:H112	1.78	0.42
8:i:27:PHE:CE2	10:l:95:ILE:HG22	2.54	0.42
18:C:315:CLA:HED2	18:C:315:CLA:H2A	2.01	0.42
18:E:310:CLA:HBB2	18:E:316:CLA:HAB	2.02	0.42
18:a:804:CLA:H52	18:a:804:CLA:H12	1.76	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
21:a:834:BCR:H11C	21:a:834:BCR:H341	1.93	0.42
2:b:272:MET:HE1	2:b:356:PRO:O	2.20	0.42
4:d:58:LEU:HD23	4:d:59:TYR:N	2.35	0.42
6:f:108:ILE:HG22	6:f:112:PHE:CE2	2.54	0.42
21:i:102:BCR:H11C	21:i:102:BCR:H341	1.84	0.42
21:i:102:BCR:H371	21:i:102:BCR:H24C	1.78	0.42
18:j:104:CLA:HBC3	21:j:105:BCR:H311	2.00	0.42
1:a:433:HIS:NE2	1:a:437:ILE:HD11	2.34	0.42
18:a:848:CLA:H41	18:a:848:CLA:H62	1.66	0.42
2:b:406:VAL:HG21	18:b:817:CLA:CHC	2.50	0.42
18:b:811:CLA:HBB1	18:b:811:CLA:HMB3	2.01	0.42
18:b:820:CLA:HBB1	18:b:820:CLA:HMB3	2.00	0.42
5:e:62:LYS:HD2	5:e:62:LYS:OXT	2.19	0.42
10:l:6:LYS:HD2	15:D:146:TYR:CE2	2.53	0.42
12:r:58:TYR:HA	12:r:59:PRO:HD3	1.89	0.42
21:r:201:BCR:H371	21:r:201:BCR:H24C	1.82	0.42
15:D:193:GLN:CD	15:D:193:GLN:H	2.28	0.42
17:H:43:GLU:H	17:H:43:GLU:CD	2.26	0.42
1:a:294:ILE:HG23	18:a:815:CLA:HMA3	2.01	0.42
1:a:426:LEU:HD13	18:a:844:CLA:C1C	2.48	0.42
2:b:91:ILE:HD13	18:b:808:CLA:HMD1	2.01	0.42
14:C:112:ILE:O	14:C:116:VAL:HG13	2.20	0.42
18:C:312:CLA:HAA1	18:C:316:CLA:HBC1	2.00	0.42
15:D:192:LYS:NZ	18:D:216:CLA:OBD	2.51	0.42
18:E:312:CLA:HBA1	18:E:312:CLA:H3A	1.67	0.42
18:H:307:CLA:H41	18:H:307:CLA:H62	1.39	0.42
18:a:837:CLA:O1A	10:l:20:PRO:HD2	2.19	0.42
2:b:162:PRO:HB2	2:b:167:PHE:CE2	2.55	0.42
7:g:116:LYS:HB3	7:g:116:LYS:HE2	1.82	0.42
9:j:25:PHE:CD1	9:j:25:PHE:C	2.96	0.42
29:C:308:KC1:C2D	18:C:311:CLA:H62	2.49	0.42
15:D:155:TRP:O	15:D:159:GLN:HG3	2.19	0.42
18:a:801:CLA:HED2	18:a:801:CLA:HBD	1.48	0.42
18:a:805:CLA:HBA1	18:a:805:CLA:H3A	1.54	0.42
21:a:834:BCR:H15C	21:a:834:BCR:H351	1.78	0.42
2:b:86:PRO:O	2:b:114:ILE:HG23	2.20	0.42
18:b:814:CLA:H112	18:b:814:CLA:H91	1.83	0.42
18:b:845:CLA:HBA1	18:b:845:CLA:H3A	1.43	0.42
18:b:849:CLA:H141	18:b:849:CLA:H161	1.77	0.42
5:e:9:ARG:CZ	6:f:178:GLU:HG3	2.49	0.42
5:e:28:ASP:HB3	5:e:32:ILE:CD1	2.49	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
10:l:110:GLU:C	10:l:112:GLN:N	2.76	0.42
21:m:101:BCR:H361	21:m:101:BCR:H20C	1.82	0.42
12:r:84:LEU:HD13	26:r:203:A86:C7	2.49	0.42
13:B:180:ASN:HD22	13:B:184:PRO:HA	1.85	0.42
14:C:196:TYR:CD2	14:C:198:PRO:HD3	2.54	0.42
26:C:304:A86:C24	18:C:307:CLA:HAB	2.49	0.42
15:D:153:GLU:HA	15:D:156:VAL:HG22	2.00	0.42
28:E:307:DD6:C10	28:E:307:DD6:C7	2.98	0.42
17:H:154:SER:CB	17:H:155:LYS:HD3	2.49	0.42
18:H:308:CLA:CGA	18:H:308:CLA:C1A	2.97	0.42
1:a:111:ILE:H	1:a:111:ILE:HD12	1.85	0.42
1:a:118:VAL:HG22	1:a:128:ASN:CG	2.45	0.42
1:a:142:SER:HB2	1:a:144:TRP:CD1	2.55	0.42
1:a:225:ASN:N	1:a:225:ASN:HD22	2.16	0.42
18:a:811:CLA:HBB1	21:a:852:BCR:H19C	2.01	0.42
18:a:855:CLA:HBC2	18:l:203:CLA:H143	2.02	0.42
2:b:391:ILE:HD13	18:b:820:CLA:CED	2.50	0.42
21:i:102:BCR:H383	21:l:201:BCR:C18	2.49	0.42
13:B:96:THR:HG21	13:B:101:ASP:HB2	2.00	0.42
15:D:41:THR:OG1	15:D:42:ALA:N	2.52	0.42
15:D:56:LYS:HD2	15:D:60:THR:HG21	2.01	0.42
1:a:384:TYR:HB2	1:a:752:THR:HG22	2.02	0.42
1:a:646:TRP:O	1:a:650:PHE:HB3	2.19	0.42
4:d:51:MET:HE1	4:d:78:ILE:CD1	2.50	0.42
4:d:73:LEU:HA	4:d:76:PHE:HD2	1.85	0.42
21:i:102:BCR:H15C	21:i:102:BCR:H351	1.85	0.42
9:j:22:THR:HG23	9:j:25:PHE:CE2	2.54	0.42
15:D:105:MET:HA	15:D:113:TRP:HZ2	1.85	0.42
16:E:70:ARG:HA	16:E:73:MET:HE3	2.01	0.42
1:a:105:LEU:HD21	1:a:154:THR:HA	2.02	0.42
18:a:801:CLA:H102	18:a:801:CLA:H62	1.76	0.42
18:a:825:CLA:HHC	18:a:825:CLA:CBB	2.42	0.42
2:b:544:LYS:NZ	6:f:180:ILE:HD11	2.35	0.42
8:i:27:PHE:CD2	10:l:95:ILE:CG2	3.03	0.42
9:j:22:THR:HG23	9:j:25:PHE:HE2	1.84	0.42
18:B:309:CLA:H41	18:B:309:CLA:H61	1.78	0.42
14:C:142:PRO:HB2	18:C:316:CLA:CHA	2.49	0.42
2:b:31:LEU:HB3	2:b:37:MET:HE2	2.01	0.41
2:b:544:LYS:HE3	2:b:544:LYS:HB3	1.83	0.41
3:c:46:GLU:HG2	3:c:47:ASP:OD2	2.20	0.41
7:g:113:ASP:HB3	7:g:116:LYS:HE2	2.01	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
7:g:137:ALA:HA	7:g:141:PRO:HG3	2.02	0.41
9:j:20:THR:HG23	21:j:105:BCR:H351	2.02	0.41
14:C:76:LEU:HD11	18:C:312:CLA:HBC1	2.01	0.41
18:C:310:CLA:O1D	18:C:310:CLA:H2A	2.20	0.41
16:E:158:TRP:HZ3	16:E:162:ARG:HH21	1.66	0.41
21:E:306:BCR:H15C	21:E:306:BCR:H351	1.83	0.41
17:H:49:MET:HE1	17:H:71:ARG:HB2	2.02	0.41
18:H:307:CLA:HMB3	18:H:307:CLA:HBB1	2.01	0.41
18:H:313:CLA:H143	18:H:313:CLA:H161	1.65	0.41
2:b:476:LEU:HD13	18:b:824:CLA:CMD	2.49	0.41
2:b:476:LEU:HD13	18:b:824:CLA:HMD1	2.01	0.41
13:B:69:ASN:HD22	13:B:69:ASN:H	1.67	0.41
14:C:100:HIS:HE1	14:C:104:LEU:HD21	1.85	0.41
14:C:103:VAL:CG2	18:C:315:CLA:HAB	2.50	0.41
18:C:307:CLA:H193	18:C:307:CLA:H161	1.73	0.41
18:C:309:CLA:H161	18:C:309:CLA:H141	1.75	0.41
18:D:209:CLA:O1A	18:D:209:CLA:H2A	2.19	0.41
16:E:39:PHE:CD1	18:E:309:CLA:HBA2	2.56	0.41
2:b:387:ALA:O	2:b:391:ILE:HG13	2.20	0.41
5:e:18:PHE:HB3	6:f:180:ILE:CD1	2.50	0.41
6:f:78:ASP:O	6:f:82:ARG:HG3	2.21	0.41
7:g:116:LYS:HA	7:g:119:GLU:OE1	2.20	0.41
21:j:105:BCR:H351	21:j:105:BCR:H15C	1.80	0.41
10:l:6:LYS:HB2	10:l:7:PRO:HD2	2.02	0.41
18:E:315:CLA:H3A	18:E:315:CLA:H12	2.02	0.41
17:H:50:ALA:HB1	17:H:166:GLU:HG2	2.02	0.41
1:a:385:ALA:H	1:a:752:THR:CG2	2.34	0.41
22:a:849:SQD:C5	22:a:849:SQD:H271	2.51	0.41
23:a:851:LMG:H312	23:a:851:LMG:H341	1.31	0.41
2:b:220:GLY:O	2:b:223:PRO:HD2	2.20	0.41
2:b:329:LEU:HD13	18:b:805:CLA:HAC1	2.03	0.41
4:d:76:PHE:O	4:d:77:LYS:HB2	2.21	0.41
21:r:201:BCR:H11C	21:r:201:BCR:H341	1.93	0.41
20:D:201:LHG:H142	20:D:201:LHG:H111	1.47	0.41
1:a:203:ALA:HB2	1:a:309:GLY:HA3	2.02	0.41
1:a:217:GLN:O	1:a:222:LEU:HD13	2.21	0.41
1:a:486:TRP:CE2	1:a:490:ILE:HD11	2.56	0.41
2:b:37:MET:HE3	2:b:42:LEU:CA	2.47	0.41
2:b:492:TRP:HZ3	18:b:813:CLA:HED1	1.85	0.41
18:b:823:CLA:O1D	9:j:35:ASP:HA	2.21	0.41
10:l:147:LEU:HD12	10:l:147:LEU:C	2.46	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
14:C:195:ILE:CG2	18:C:317:CLA:HBC3	2.51	0.41
18:E:315:CLA:H91	18:E:315:CLA:H112	1.84	0.41
17:H:117:ALA:O	17:H:121:GLN:HG3	2.21	0.41
17:H:156:MET:HE3	17:H:160:GLN:HB3	2.01	0.41
18:H:306:CLA:CGA	18:H:306:CLA:H3A	2.50	0.41
1:a:111:ILE:HD12	1:a:111:ILE:N	2.35	0.41
1:a:204:GLY:O	1:a:208:LEU:HB2	2.21	0.41
1:a:217:GLN:HA	1:a:221:ALA:HB3	2.03	0.41
1:a:225:ASN:OD1	1:a:291:LEU:HD11	2.20	0.41
18:a:806:CLA:H122	18:a:806:CLA:H162	1.87	0.41
18:a:813:CLA:H141	18:a:813:CLA:H161	1.53	0.41
12:r:98:LEU:HD21	12:r:109:TRP:CD2	2.55	0.41
20:E:301:LHG:H191	20:E:301:LHG:H222	1.81	0.41
17:H:90:TRP:CZ2	17:H:95:PRO:HD2	2.54	0.41
18:a:810:CLA:H61	18:a:810:CLA:H41	1.90	0.41
18:a:813:CLA:H2	18:a:813:CLA:H62	1.74	0.41
21:a:834:BCR:H20C	21:a:834:BCR:H361	1.81	0.41
13:B:180:ASN:ND2	13:B:185:GLY:H	2.18	0.41
16:E:86:LYS:HE3	16:E:221:PHE:HE1	1.85	0.41
17:H:44:LYS:HE2	17:H:64:GLN:O	2.21	0.41
17:H:120:PHE:O	17:H:124:LEU:HD13	2.20	0.41
18:a:804:CLA:C10	18:a:807:CLA:H191	2.51	0.41
2:b:635:ASN:HB2	2:b:636:PRO:CD	2.50	0.41
18:b:818:CLA:H141	18:b:818:CLA:H161	1.77	0.41
18:b:820:CLA:H41	18:b:820:CLA:H62	1.63	0.41
18:b:822:CLA:H92	18:b:822:CLA:H61	1.90	0.41
3:c:5:VAL:HG22	3:c:67:VAL:HG22	2.02	0.41
21:j:105:BCR:H11C	21:j:105:BCR:H341	1.80	0.41
10:l:82:THR:O	10:l:86:ILE:HG13	2.21	0.41
13:B:146:LEU:HD23	13:B:148:LEU:HD21	2.02	0.41
14:C:112:ILE:HA	14:C:115:LEU:HD23	2.02	0.41
16:E:208:MET:HE1	18:E:316:CLA:HHB	2.01	0.41
16:E:208:MET:HE1	23:E:318:LMG:HC91	2.02	0.41
1:a:482:ILE:HG13	1:a:483:PHE:N	2.36	0.41
21:a:833:BCR:H403	18:a:847:CLA:H121	2.03	0.41
18:a:841:CLA:H142	18:a:841:CLA:H111	1.61	0.41
18:a:846:CLA:H152	18:a:846:CLA:H112	1.90	0.41
2:b:86:PRO:HB2	2:b:115:THR:HG23	2.03	0.41
2:b:179:SER:HB3	2:b:287:GLY:HA3	2.02	0.41
2:b:421:LEU:HB3	2:b:531:LEU:HD13	2.03	0.41
2:b:491:VAL:HG11	18:b:813:CLA:CED	2.49	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
18:b:828:CLA:H2A	8:i:27:PHE:HE1	1.86	0.41
21:b:832:BCR:H11C	21:b:832:BCR:H341	1.85	0.41
18:b:843:CLA:H193	18:b:843:CLA:H161	1.69	0.41
9:j:2:ASN:C	9:j:2:ASN:ND2	2.79	0.41
10:l:90:THR:HG21	16:E:138:ILE:HG21	2.03	0.41
13:B:145:TYR:HD2	13:B:146:LEU:HD13	1.85	0.41
13:B:149:TRP:CZ3	18:B:312:CLA:HMA2	2.53	0.41
13:B:169:LEU:HD22	26:B:304:A86:C28	2.51	0.41
18:C:318:CLA:CBD	18:C:318:CLA:HAA2	2.51	0.41
18:D:208:CLA:H161	18:D:208:CLA:H141	1.62	0.41
16:E:181:GLU:H	16:E:181:GLU:CD	2.28	0.41
17:H:155:LYS:HE2	17:H:155:LYS:HB2	1.70	0.41
26:H:302:A86:C3	18:H:309:CLA:HMD1	2.48	0.41
18:H:306:CLA:H61	18:H:306:CLA:H92	1.77	0.41
18:H:310:CLA:H62	18:H:310:CLA:H41	1.48	0.41
2:b:492:TRP:HZ2	18:b:825:CLA:HMA3	1.86	0.41
2:b:582:MET:O	2:b:586:ILE:HG12	2.21	0.41
2:b:663:LEU:CD2	2:b:708:GLY:HA3	2.50	0.41
21:f:205:BCR:H11C	21:f:205:BCR:H341	1.76	0.41
21:l:204:BCR:H15C	21:l:204:BCR:H351	1.80	0.41
14:C:125:TRP:CZ3	14:C:141:PHE:CE2	3.08	0.41
15:D:42:ALA:N	15:D:43:PRO:HD2	2.37	0.41
18:E:312:CLA:HED3	18:E:313:CLA:HBC2	2.03	0.41
1:a:494:ALA:HB3	1:a:495:PRO:HD3	2.04	0.40
18:a:830:CLA:HBA2	18:a:830:CLA:H3A	1.78	0.40
18:a:840:CLA:H13	19:b:831:PQN:H192	2.01	0.40
2:b:171:GLU:HG3	2:b:290:TYR:HB3	2.03	0.40
2:b:493:LEU:N	2:b:494:PRO:CD	2.84	0.40
2:b:506:ASP:HA	2:b:509:LEU:HD21	2.03	0.40
2:b:647:TRP:HZ3	21:b:839:BCR:H312	1.86	0.40
18:b:809:CLA:HBA2	18:b:809:CLA:H3A	1.60	0.40
18:b:829:CLA:HBB1	18:b:829:CLA:CHC	2.41	0.40
21:b:836:BCR:H11C	21:b:836:BCR:H341	1.85	0.40
18:b:848:CLA:H61	18:b:848:CLA:H41	1.88	0.40
4:d:18:GLY:H	10:l:15:GLY:HA2	1.86	0.40
21:i:102:BCR:H20C	21:i:102:BCR:H361	1.87	0.40
14:C:112:ILE:N	14:C:113:PRO:CD	2.84	0.40
16:E:208:MET:O	16:E:212:ILE:HG13	2.21	0.40
17:H:152:GLN:HG3	18:H:311:CLA:H12	2.01	0.40
18:H:305:CLA:H62	18:H:305:CLA:C1B	2.51	0.40
18:H:305:CLA:HBA2	18:H:305:CLA:H3A	1.18	0.40

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
29:H:312:KC1:CGD	29:H:312:KC1:CAA	2.98	0.40
18:a:802:CLA:HBA2	2:b:426:LEU:HD23	2.03	0.40
18:a:817:CLA:H11	18:a:817:CLA:HBA1	1.88	0.40
2:b:216:PRO:HD2	18:b:812:CLA:C3D	2.51	0.40
2:b:440:ASP:CG	2:b:614:TYR:HB2	2.46	0.40
3:c:46:GLU:OE1	7:g:68:GLY:HA3	2.22	0.40
7:g:146:SER:O	7:g:147:SER:C	2.64	0.40
18:l:203:CLA:H122	18:l:203:CLA:H161	1.93	0.40
11:m:12:MET:CG	21:m:101:BCR:H382	2.48	0.40
1:a:407:ALA:O	1:a:411:ILE:HG13	2.21	0.40
18:a:803:CLA:H111	18:a:803:CLA:H151	1.42	0.40
18:a:841:CLA:H202	18:a:841:CLA:H161	1.60	0.40
2:b:515:ASP:OD1	2:b:515:ASP:N	2.53	0.40
18:b:844:CLA:H162	18:b:844:CLA:H141	1.90	0.40
7:g:151:MET:HE3	7:g:151:MET:HB3	1.96	0.40
18:C:307:CLA:HBB1	18:C:307:CLA:CMB	2.49	0.40
15:D:65:ARG:HD3	15:D:65:ARG:O	2.21	0.40
16:E:91:ILE:CD1	16:E:118:GLU:HG3	2.52	0.40
16:E:127:VAL:O	16:E:131:MET:HG2	2.21	0.40
18:H:314:CLA:H61	18:H:314:CLA:H41	1.79	0.40
1:a:118:VAL:HG22	1:a:128:ASN:OD1	2.21	0.40
18:b:808:CLA:H92	21:i:102:BCR:H352	2.03	0.40
7:g:114:LEU:HD22	7:g:114:LEU:N	2.36	0.40
10:l:75:LEU:HG	10:l:136:PHE:CD1	2.57	0.40
18:l:202:CLA:HHC	18:l:202:CLA:CBB	2.35	0.40
18:l:203:CLA:H91	18:l:203:CLA:H111	1.33	0.40
14:C:92:GLY:HA3	18:C:310:CLA:C4A	2.51	0.40
1:a:28:LYS:HB2	18:a:809:CLA:HAA2	2.04	0.40
1:a:709:ASN:HB3	6:f:130:ARG:NH1	2.37	0.40
18:a:803:CLA:H12	18:a:803:CLA:C19	2.49	0.40
21:a:850:BCR:H312	21:a:850:BCR:HC32	1.88	0.40
2:b:354:LEU:HD13	18:b:842:CLA:HBD	2.04	0.40
2:b:457:LEU:CD1	6:f:96:VAL:HG13	2.49	0.40
2:b:684:THR:HG23	2:b:687:ALA:HB3	2.03	0.40
18:b:806:CLA:H152	18:b:819:CLA:HBB2	2.03	0.40
18:b:807:CLA:HBC2	11:m:11:LEU:HD23	2.03	0.40
7:g:137:ALA:HA	7:g:141:PRO:CG	2.52	0.40
18:B:312:CLA:H43	29:B:313:KC1:O1A	2.22	0.40
14:C:106:ILE:HG13	14:C:110:ASN:ND2	2.36	0.40

There are no symmetry-related clashes.

5.3 Torsion angles

5.3.1 Protein backbone

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	741/743 (100%)	723 (98%)	18 (2%)	0	100	100
2	b	730/732 (100%)	713 (98%)	17 (2%)	0	100	100
3	c	78/80 (98%)	74 (95%)	4 (5%)	0	100	100
4	d	130/132 (98%)	126 (97%)	4 (3%)	0	100	100
5	e	60/62 (97%)	58 (97%)	2 (3%)	0	100	100
6	f	158/160 (99%)	152 (96%)	6 (4%)	0	100	100
7	g	129/131 (98%)	121 (94%)	8 (6%)	0	100	100
8	i	31/33 (94%)	30 (97%)	1 (3%)	0	100	100
9	j	38/40 (95%)	36 (95%)	2 (5%)	0	100	100
10	l	144/146 (99%)	138 (96%)	6 (4%)	0	100	100
11	m	27/29 (93%)	26 (96%)	1 (4%)	0	100	100
12	r	87/89 (98%)	86 (99%)	1 (1%)	0	100	100
13	B	163/165 (99%)	157 (96%)	6 (4%)	0	100	100
14	C	168/170 (99%)	165 (98%)	3 (2%)	0	100	100
15	D	162/164 (99%)	156 (96%)	6 (4%)	0	100	100
16	E	183/185 (99%)	180 (98%)	3 (2%)	0	100	100
17	H	166/168 (99%)	162 (98%)	4 (2%)	0	100	100
All	All	3195/3229 (99%)	3103 (97%)	92 (3%)	0	100	100

There are no Ramachandran outliers to report.

5.3.2 Protein sidechains

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	603/603 (100%)	598 (99%)	5 (1%)	73	82
2	b	597/597 (100%)	591 (99%)	6 (1%)	68	80
3	c	69/69 (100%)	68 (99%)	1 (1%)	59	77
4	d	111/111 (100%)	111 (100%)	0	100	100
5	e	55/55 (100%)	52 (94%)	3 (6%)	19	52
6	f	132/132 (100%)	130 (98%)	2 (2%)	57	76
7	g	98/98 (100%)	95 (97%)	3 (3%)	35	66
8	i	28/28 (100%)	28 (100%)	0	100	100
9	j	36/36 (100%)	34 (94%)	2 (6%)	19	52
10	l	113/113 (100%)	113 (100%)	0	100	100
11	m	22/22 (100%)	22 (100%)	0	100	100
12	r	72/72 (100%)	72 (100%)	0	100	100
13	B	133/133 (100%)	124 (93%)	9 (7%)	14	46
14	C	129/130 (99%)	123 (95%)	6 (5%)	23	57
15	D	128/128 (100%)	124 (97%)	4 (3%)	35	66
16	E	144/144 (100%)	144 (100%)	0	100	100
17	H	136/136 (100%)	135 (99%)	1 (1%)	76	83
All	All	2606/2607 (100%)	2564 (98%)	42 (2%)	54	75

All (42) residues with a non-rotameric sidechain are listed below:

Mol	Chain	Res	Type
1	a	11	LYS
1	a	24	THR
1	a	80	GLN
1	a	198	MET
1	a	658	VAL
2	b	54	LEU
2	b	112	VAL
2	b	115	THR
2	b	242	VAL
2	b	245	THR
2	b	367	GLU
3	c	65	VAL

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Mol	Chain	Res	Type
5	e	14	GLU
5	e	32	ILE
5	e	59	VAL
6	f	139	SER
6	f	178	GLU
7	g	91	TYR
7	g	152	GLU
7	g	158	LEU
9	j	2	ASN
9	j	37	LEU
13	B	43	GLU
13	B	46	LYS
13	B	69	ASN
13	B	146	LEU
13	B	148	LEU
13	B	158	GLU
13	B	164	LEU
13	B	165	LYS
13	B	169	LEU
14	C	63	MET
14	C	68	LEU
14	C	103	VAL
14	C	106	ILE
14	C	116	VAL
14	C	201	ASN
15	D	71	ASN
15	D	96	GLU
15	D	108	LEU
15	D	129	GLN
17	H	124	LEU

Sometimes sidechains can be flipped to improve hydrogen bonding and reduce clashes. All (44) such sidechains are listed below:

Mol	Chain	Res	Type
1	a	12	ASN
1	a	14	GLN
1	a	116	GLN
1	a	134	ASN
1	a	217	GLN
1	a	321	ASN
1	a	421	ASN
1	a	478	GLN

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Mol	Chain	Res	Type
1	a	489	ASN
1	a	501	ASN
1	a	723	GLN
2	b	10	GLN
2	b	34	HIS
2	b	89	HIS
2	b	113	ASN
2	b	157	GLN
2	b	293	ASN
2	b	451	GLN
2	b	472	GLN
2	b	474	ASN
2	b	626	ASN
2	b	688	ASN
2	b	703	GLN
4	d	53	ASN
4	d	114	ASN
5	e	20	GLN
6	f	55	GLN
6	f	171	ASN
9	j	2	ASN
10	l	9	ASN
10	l	33	ASN
13	B	52	ASN
13	B	69	ASN
13	B	180	ASN
13	B	181	HIS
14	C	52	ASN
14	C	201	ASN
15	D	71	ASN
15	D	85	ASN
16	E	159	GLN
16	E	213	ASN
17	H	91	GLN
17	H	185	GLN
17	H	190	HIS

5.3.3 RNA ⓘ

There are no RNA molecules in this entry.

5.4 Non-standard residues in protein, DNA, RNA chains [i](#)

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

5.5 Carbohydrates [i](#)

There are no oligosaccharides in this entry.

5.6 Ligand geometry [i](#)

217 ligands are modelled in this entry.

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	$\# Z > 2$
18	CLA	a	809	18,1	66,70,73	1.18	7 (10%)	78,109,113	1.27	4 (5%)
26	A86	D	206	-	47,50,50	1.70	9 (19%)	51,76,76	3.31	21 (41%)
21	BCR	i	103	-	41,41,41	0.67	0	56,56,56	2.05	13 (23%)
21	BCR	f	201	-	41,41,41	0.94	2 (4%)	56,56,56	1.94	17 (30%)
26	A86	b	847	-	47,50,50	1.63	9 (19%)	51,76,76	2.69	18 (35%)
18	CLA	b	849	-	69,73,73	1.17	9 (13%)	82,113,113	1.30	6 (7%)
18	CLA	C	312	14	65,69,73	1.20	9 (13%)	77,108,113	1.38	7 (9%)
18	CLA	D	217	-	45,49,73	1.39	8 (17%)	54,84,113	1.47	6 (11%)
18	CLA	b	814	-	69,73,73	1.18	9 (13%)	82,113,113	1.30	7 (8%)
18	CLA	a	815	1	69,73,73	1.16	8 (11%)	82,113,113	1.26	7 (8%)
18	CLA	a	816	1	49,53,73	1.39	8 (16%)	58,89,113	1.42	4 (6%)
20	LHG	b	835	18	48,48,48	1.14	3 (6%)	51,54,54	1.04	3 (5%)
18	CLA	E	311	16	50,54,73	1.36	9 (18%)	59,90,113	1.38	5 (8%)
18	CLA	a	825	1	64,68,73	1.24	11 (17%)	76,107,113	1.47	10 (13%)
28	DD6	B	303	-	40,45,45	2.06	7 (17%)	51,67,67	2.92	18 (35%)
18	CLA	a	828	1	69,73,73	1.16	8 (11%)	82,113,113	1.26	6 (7%)
21	BCR	m	101	-	41,41,41	0.66	0	56,56,56	2.14	15 (26%)
18	CLA	b	817	2	64,68,73	1.21	9 (14%)	76,107,113	1.29	5 (6%)

Mol	Type	Chain	Res	Link	Bond lengths			Bond angles		
					Counts	RMSZ	# Z > 2	Counts	RMSZ	# Z > 2
18	CLA	B	311	13	69,73,73	1.16	8 (11%)	82,113,113	1.27	6 (7%)
26	A86	m	102	-	47,50,50	1.56	9 (19%)	51,76,76	3.32	25 (49%)
18	CLA	a	814	1	69,73,73	1.16	9 (13%)	82,113,113	1.27	6 (7%)
18	CLA	E	310	16	69,73,73	1.17	8 (11%)	82,113,113	1.25	5 (6%)
18	CLA	a	802	-	69,73,73	1.16	9 (13%)	82,113,113	1.27	5 (6%)
23	LMG	l	207	-	40,40,55	1.23	4 (10%)	48,48,63	1.79	4 (8%)
18	CLA	b	846	2	69,73,73	1.16	8 (11%)	82,113,113	1.24	5 (6%)
18	CLA	H	314	17	69,73,73	1.17	9 (13%)	82,113,113	1.38	8 (9%)
18	CLA	a	838	1	69,73,73	1.16	7 (10%)	82,113,113	1.27	6 (7%)
18	CLA	b	830	20	69,73,73	1.16	9 (13%)	82,113,113	1.27	5 (6%)
18	CLA	H	310	-	64,68,73	1.21	8 (12%)	76,107,113	1.45	7 (9%)
18	CLA	a	824	1	55,59,73	1.29	8 (14%)	64,96,113	1.43	8 (12%)
21	BCR	r	201	-	41,41,41	0.67	0	56,56,56	1.77	15 (26%)
18	CLA	a	844	1	69,73,73	1.17	7 (10%)	82,113,113	1.28	5 (6%)
23	LMG	E	318	-	43,43,55	2.00	7 (16%)	51,51,63	1.99	7 (13%)
20	LHG	B	315	-	41,41,48	0.99	2 (4%)	44,47,54	1.10	3 (6%)
18	CLA	B	308	13	69,73,73	1.17	9 (13%)	82,113,113	1.27	5 (6%)
18	CLA	a	823	1	49,53,73	1.37	8 (16%)	58,89,113	1.42	4 (6%)
18	CLA	a	817	1	53,57,73	1.32	8 (15%)	61,93,113	1.42	5 (8%)
29	KC1	B	313	-	49,53,53	1.55	8 (16%)	61,89,89	1.89	13 (21%)
18	CLA	a	839	1	69,73,73	1.14	9 (13%)	82,113,113	1.30	5 (6%)
18	CLA	D	207	-	65,69,73	1.20	10 (15%)	77,108,113	1.39	7 (9%)
18	CLA	b	818	2	69,73,73	1.17	8 (11%)	82,113,113	1.27	6 (7%)
18	CLA	B	312	13	69,73,73	1.15	8 (11%)	82,113,113	1.27	4 (4%)
18	CLA	B	307	13	69,73,73	1.16	9 (13%)	82,113,113	1.26	5 (6%)
18	CLA	b	811	2	64,68,73	1.20	8 (12%)	76,107,113	1.40	6 (7%)
18	CLA	l	202	10	53,57,73	1.33	9 (16%)	61,93,113	1.49	8 (13%)
18	CLA	b	828	-	69,73,73	1.17	9 (13%)	82,113,113	1.28	6 (7%)
18	CLA	a	808	1	69,73,73	1.17	9 (13%)	82,113,113	1.28	5 (6%)
18	CLA	b	805	2	69,73,73	1.16	8 (11%)	82,113,113	1.34	6 (7%)
21	BCR	E	306	-	41,41,41	0.69	0	56,56,56	2.02	15 (26%)
23	LMG	C	319	18	44,44,55	1.32	7 (15%)	52,52,63	1.85	4 (7%)
18	CLA	H	306	17	69,73,73	1.17	9 (13%)	82,113,113	1.29	6 (7%)
18	CLA	a	807	1	69,73,73	1.19	9 (13%)	82,113,113	1.46	9 (10%)
20	LHG	D	201	-	48,48,48	1.12	4 (8%)	51,54,54	1.05	2 (3%)

Mol	Type	Chain	Res	Link	Bond lengths			Bond angles		
					Counts	RMSZ	# Z > 2	Counts	RMSZ	# Z > 2
29	KC1	C	313	-	49,53,53	1.36	6 (12%)	61,89,89	1.76	13 (21%)
25	DGD	b	834	-	61,61,67	0.86	2 (3%)	75,75,81	1.11	4 (5%)
18	CLA	b	816	2	59,63,73	1.25	8 (13%)	70,101,113	1.31	4 (5%)
18	CLA	a	822	1	54,58,73	1.30	7 (12%)	64,95,113	1.41	6 (9%)
21	BCR	a	852	-	41,41,41	0.71	0	56,56,56	2.15	17 (30%)
18	CLA	a	806	1	69,73,73	1.16	9 (13%)	82,113,113	1.27	5 (6%)
18	CLA	b	838	-	69,73,73	1.17	9 (13%)	82,113,113	1.30	5 (6%)
18	CLA	E	314	16	49,53,73	1.38	11 (22%)	58,89,113	1.47	6 (10%)
18	CLA	a	801	1	69,73,73	1.22	9 (13%)	82,113,113	1.34	8 (9%)
18	CLA	D	214	15	60,64,73	1.25	9 (15%)	71,102,113	1.38	7 (9%)
21	BCR	j	105	-	41,41,41	0.73	0	56,56,56	2.09	11 (19%)
18	CLA	B	309	-	69,73,73	1.16	9 (13%)	82,113,113	1.29	5 (6%)
18	CLA	a	805	1	69,73,73	1.15	8 (11%)	82,113,113	1.31	6 (7%)
26	A86	H	302	-	47,50,50	1.80	10 (21%)	51,76,76	3.56	24 (47%)
21	BCR	b	832	-	41,41,41	0.68	0	56,56,56	1.92	14 (25%)
18	CLA	C	316	14	45,49,73	1.40	10 (22%)	54,84,113	1.52	5 (9%)
18	CLA	a	846	-	69,73,73	1.15	8 (11%)	82,113,113	1.26	5 (6%)
18	CLA	b	825	-	49,53,73	1.40	8 (16%)	58,89,113	1.42	5 (8%)
18	CLA	b	824	-	69,73,73	1.20	8 (11%)	82,113,113	1.48	7 (8%)
21	BCR	j	101	-	41,41,41	0.69	0	56,56,56	2.06	13 (23%)
24	SF4	c	101	3	0,12,12	-	-	-		
18	CLA	b	844	2	66,70,73	1.18	8 (12%)	78,109,113	1.32	5 (6%)
18	CLA	a	843	-	69,73,73	1.16	8 (11%)	82,113,113	1.32	7 (8%)
18	CLA	B	314	13	45,49,73	1.41	9 (20%)	54,84,113	1.49	7 (12%)
18	CLA	E	315	16	69,73,73	1.18	9 (13%)	82,113,113	1.27	5 (6%)
18	CLA	D	215	-	45,49,73	1.41	9 (20%)	54,84,113	1.46	5 (9%)
26	A86	H	301	-	47,50,50	1.74	10 (21%)	51,76,76	3.37	25 (49%)
18	CLA	H	307	17	69,73,73	1.18	8 (11%)	82,113,113	1.48	12 (14%)
18	CLA	a	830	20	56,60,73	1.30	9 (16%)	65,97,113	1.37	5 (7%)
18	CLA	b	829	2	69,73,73	1.20	9 (13%)	82,113,113	1.44	8 (9%)
18	CLA	a	812	-	49,53,73	1.39	9 (18%)	58,89,113	1.45	4 (6%)
21	BCR	a	833	-	41,41,41	0.89	1 (2%)	56,56,56	1.93	21 (37%)
18	CLA	a	841	-	69,73,73	1.20	9 (13%)	82,113,113	1.34	8 (9%)
18	CLA	b	812	2	63,67,73	1.22	9 (14%)	74,105,113	1.31	6 (8%)
26	A86	C	304	-	47,50,50	1.67	9 (19%)	51,76,76	2.83	16 (31%)

Mol	Type	Chain	Res	Link	Bond lengths			Bond angles		
					Counts	RMSZ	# Z > 2	Counts	RMSZ	# Z > 2
18	CLA	C	310	14	50,54,73	1.31	8 (16%)	59,90,113	1.41	5 (8%)
18	CLA	a	854	-	69,73,73	1.20	7 (10%)	82,113,113	1.43	6 (7%)
26	A86	D	203	-	47,50,50	1.58	7 (14%)	51,76,76	3.56	20 (39%)
18	CLA	a	847	1	69,73,73	1.17	9 (13%)	82,113,113	1.28	5 (6%)
20	LHG	a	831	-	48,48,48	1.16	3 (6%)	51,54,54	1.02	3 (5%)
21	BCR	b	839	-	41,41,41	1.01	2 (4%)	56,56,56	2.04	20 (35%)
18	CLA	b	823	2	62,66,73	1.22	8 (12%)	73,104,113	1.36	6 (8%)
18	CLA	f	202	-	69,73,73	1.16	7 (10%)	82,113,113	1.30	7 (8%)
28	DD6	E	308	-	40,45,45	1.54	2 (5%)	51,67,67	1.77	12 (23%)
24	SF4	c	102	3	0,12,12	-	-	-	-	-
18	CLA	a	811	1	54,58,73	1.29	8 (14%)	64,95,113	1.43	7 (10%)
18	CLA	H	309	17	50,54,73	1.35	8 (16%)	59,90,113	1.40	4 (6%)
18	CLA	a	804	18,1	59,63,73	1.27	9 (15%)	70,101,113	1.34	6 (8%)
18	CLA	b	806	2	69,73,73	1.17	8 (11%)	82,113,113	1.26	5 (6%)
18	CLA	b	821	2	54,58,73	1.31	9 (16%)	64,95,113	1.36	6 (9%)
18	CLA	b	826	2	64,68,73	1.21	8 (12%)	76,107,113	1.36	7 (9%)
21	BCR	E	305	-	41,41,41	0.67	0	56,56,56	1.88	17 (30%)
18	CLA	a	820	1	69,73,73	1.17	8 (11%)	82,113,113	1.31	7 (8%)
18	CLA	b	815	2	50,54,73	1.36	9 (18%)	59,90,113	1.45	4 (6%)
24	SF4	b	802	2	0,12,12	-	-	-	-	-
18	CLA	a	848	1	60,64,73	1.26	8 (13%)	71,102,113	1.31	5 (7%)
18	CLA	C	306	23	49,53,73	1.39	9 (18%)	58,89,113	1.44	4 (6%)
18	CLA	a	837	1	64,68,73	1.25	8 (12%)	76,107,113	1.52	8 (10%)
26	A86	B	304	-	47,50,50	1.70	9 (19%)	51,76,76	3.02	19 (37%)
18	CLA	E	309	16	69,73,73	1.16	8 (11%)	82,113,113	1.28	7 (8%)
21	BCR	f	205	-	41,41,41	0.71	0	56,56,56	1.96	15 (26%)
28	DD6	D	205	-	40,45,45	1.98	7 (17%)	51,67,67	3.21	24 (47%)
19	PQN	a	829	-	34,34,34	1.63	2 (5%)	43,45,45	1.10	3 (6%)
23	LMG	D	202	-	46,46,55	1.80	7 (15%)	54,54,63	1.37	5 (9%)
18	CLA	H	313	17	69,73,73	1.17	9 (13%)	82,113,113	1.44	7 (8%)
27	ET4	l	206	-	41,43,43	1.45	8 (19%)	52,60,60	2.23	18 (34%)
18	CLA	b	841	2	69,73,73	1.17	8 (11%)	82,113,113	1.26	6 (7%)
18	CLA	a	819	1	69,73,73	1.16	8 (11%)	82,113,113	1.30	6 (7%)
28	DD6	H	303	-	40,45,45	2.16	11 (27%)	51,67,67	2.44	19 (37%)
29	KC1	H	312	17	49,53,53	1.63	8 (16%)	61,89,89	1.80	13 (21%)

Mol	Type	Chain	Res	Link	Bond lengths			Bond angles		
					Counts	RMSZ	# Z > 2	Counts	RMSZ	# Z > 2
26	A86	B	305	-	47,50,50	1.72	12 (25%)	51,76,76	3.43	27 (52%)
21	BCR	l	201	-	41,41,41	0.96	2 (4%)	56,56,56	2.05	16 (28%)
18	CLA	C	317	-	46,50,73	1.44	8 (17%)	53,85,113	1.60	5 (9%)
18	CLA	b	827	2	51,55,73	1.35	9 (17%)	60,91,113	1.40	5 (8%)
18	CLA	D	211	15	50,54,73	1.37	9 (18%)	59,90,113	1.41	4 (6%)
18	CLA	b	843	2	69,73,73	1.17	8 (11%)	82,113,113	1.35	6 (7%)
18	CLA	b	804	2	49,53,73	1.38	9 (18%)	58,89,113	1.52	6 (10%)
20	LHG	j	102	-	48,48,48	1.15	6 (12%)	51,54,54	0.93	2 (3%)
26	A86	C	305	-	47,50,50	1.76	10 (21%)	51,76,76	3.40	26 (50%)
18	CLA	H	311	17	61,65,73	1.24	7 (11%)	72,103,113	1.40	6 (8%)
18	CLA	l	203	10	69,73,73	1.18	8 (11%)	82,113,113	1.45	7 (8%)
18	CLA	b	813	2	59,63,73	1.26	8 (13%)	70,101,113	1.35	5 (7%)
18	CLA	C	307	14	69,73,73	1.17	7 (10%)	82,113,113	1.35	8 (9%)
18	CLA	b	837	2	64,68,73	1.20	8 (12%)	76,107,113	1.44	6 (7%)
19	PQN	b	831	-	34,34,34	1.61	2 (5%)	43,45,45	1.21	4 (9%)
26	A86	r	203	-	47,50,50	1.78	10 (21%)	51,76,76	3.90	27 (52%)
29	KC1	C	308	14	49,53,53	1.35	6 (12%)	61,89,89	1.78	13 (21%)
20	LHG	H	316	-	34,34,48	1.15	4 (11%)	37,40,54	0.98	2 (5%)
28	DD6	E	303	-	40,45,45	1.53	2 (5%)	51,67,67	1.88	8 (15%)
18	CLA	b	820	2	69,73,73	1.16	9 (13%)	82,113,113	1.48	7 (8%)
18	CLA	a	853	1	59,63,73	1.27	8 (13%)	70,101,113	1.37	6 (8%)
18	CLA	B	306	13	61,65,73	1.23	9 (14%)	73,102,113	1.25	3 (4%)
18	CLA	H	305	-	58,62,73	1.27	8 (13%)	68,99,113	1.48	6 (8%)
18	CLA	b	803	2	69,73,73	1.22	9 (13%)	82,113,113	1.33	8 (9%)
18	CLA	b	840	2	65,69,73	1.19	9 (13%)	77,108,113	1.34	6 (7%)
18	CLA	b	848	2	58,62,73	1.27	8 (13%)	68,99,113	1.31	4 (5%)
28	DD6	C	303	-	40,45,45	1.70	3 (7%)	51,67,67	2.28	17 (33%)
18	CLA	b	850	2	69,73,73	1.16	7 (10%)	82,113,113	1.28	5 (6%)
21	BCR	b	836	-	41,41,41	0.74	0	56,56,56	2.00	17 (30%)
18	CLA	b	808	2	69,73,73	1.16	9 (13%)	82,113,113	1.28	6 (7%)
18	CLA	D	208	15	69,73,73	1.20	11 (15%)	82,113,113	1.46	9 (10%)
18	CLA	a	810	1	69,73,73	1.16	9 (13%)	82,113,113	1.27	5 (6%)
18	CLA	H	308	17	69,73,73	1.17	9 (13%)	82,113,113	1.43	7 (8%)
21	BCR	b	833	-	41,41,41	0.69	0	56,56,56	3.34	20 (35%)
18	CLA	b	809	2	59,63,73	1.28	9 (15%)	70,101,113	1.36	5 (7%)

Mol	Type	Chain	Res	Link	Bond lengths			Bond angles		
					Counts	RMSZ	# Z > 2	Counts	RMSZ	# Z > 2
18	CLA	f	204	6	49,53,73	1.39	8 (16%)	58,89,113	1.43	4 (6%)
18	CLA	E	316	-	60,64,73	1.25	8 (13%)	71,102,113	1.34	5 (7%)
18	CLA	j	104	9	45,49,73	1.41	8 (17%)	54,84,113	1.51	5 (9%)
18	CLA	a	818	1	55,59,73	1.31	8 (14%)	64,96,113	1.55	7 (10%)
18	CLA	a	840	-	69,73,73	1.21	8 (11%)	82,113,113	1.36	6 (7%)
18	CLA	D	209	15	53,57,73	1.33	9 (16%)	61,93,113	1.40	5 (8%)
20	LHG	E	301	-	48,48,48	1.15	6 (12%)	51,54,54	1.05	3 (5%)
18	CLA	D	212	15	46,50,73	1.39	8 (17%)	53,85,113	1.41	4 (7%)
18	CLA	C	315	14	46,50,73	1.41	7 (15%)	53,85,113	1.41	4 (7%)
21	BCR	a	834	-	41,41,41	0.70	0	56,56,56	1.98	15 (26%)
23	LMG	C	301	-	31,31,55	1.60	6 (19%)	39,39,63	3.05	7 (17%)
18	CLA	b	842	2	69,73,73	1.16	8 (11%)	82,113,113	1.28	5 (6%)
26	A86	E	302	-	47,50,50	1.68	9 (19%)	51,76,76	3.04	24 (47%)
18	CLA	C	314	14	45,49,73	1.41	11 (24%)	54,84,113	1.49	6 (11%)
22	SQD	a	849	-	31,32,54	3.13	8 (25%)	33,36,65	2.78	8 (24%)
18	CLA	b	819	2	69,73,73	1.16	9 (13%)	82,113,113	1.25	5 (6%)
18	CLA	i	101	-	64,68,73	1.21	8 (12%)	76,107,113	1.32	7 (9%)
18	CLA	b	822	2	69,73,73	1.16	8 (11%)	82,113,113	1.30	6 (7%)
18	CLA	f	203	-	69,73,73	1.16	8 (11%)	82,113,113	1.27	5 (6%)
18	CLA	a	842	-	69,73,73	1.17	7 (10%)	82,113,113	1.39	6 (7%)
21	BCR	a	850	-	41,41,41	1.05	2 (4%)	56,56,56	1.80	11 (19%)
18	CLA	l	205	-	49,53,73	1.38	8 (16%)	58,89,113	1.44	6 (10%)
28	DD6	E	307	-	40,45,45	1.71	3 (7%)	51,67,67	2.90	20 (39%)
18	CLA	a	836	1	69,73,73	1.15	7 (10%)	82,113,113	1.24	5 (6%)
18	CLA	b	810	2	69,73,73	1.17	9 (13%)	82,113,113	1.29	5 (6%)
18	CLA	a	821	1	69,73,73	1.17	8 (11%)	82,113,113	1.27	5 (6%)
21	BCR	l	204	-	41,41,41	0.87	0	56,56,56	2.06	15 (26%)
18	CLA	C	318	14	46,50,73	1.38	8 (17%)	53,85,113	1.48	3 (5%)
20	LHG	a	832	18	26,26,48	1.31	5 (19%)	29,32,54	1.17	2 (6%)
26	A86	D	204	-	47,50,50	1.73	10 (21%)	51,76,76	3.96	24 (47%)
21	BCR	i	102	-	41,41,41	0.73	1 (2%)	56,56,56	1.96	17 (30%)
26	A86	B	302	-	47,50,50	1.81	10 (21%)	51,76,76	3.58	24 (47%)
18	CLA	b	845	-	69,73,73	1.16	8 (11%)	82,113,113	1.26	5 (6%)
18	CLA	a	827	1	69,73,73	1.15	8 (11%)	82,113,113	1.31	6 (7%)
26	A86	H	304	-	47,50,50	1.71	9 (19%)	51,76,76	3.22	23 (45%)

Mol	Type	Chain	Res	Link	Bond lengths			Bond angles		
					Counts	RMSZ	# Z > 2	Counts	RMSZ	# Z > 2
18	CLA	D	210	-	60,64,73	1.25	8 (13%)	71,102,113	1.43	6 (8%)
20	LHG	a	835	-	32,32,48	1.33	5 (15%)	35,38,54	1.20	2 (5%)
23	LMG	a	851	-	33,33,55	1.59	5 (15%)	41,41,63	3.67	7 (17%)
18	CLA	b	807	2	69,73,73	1.16	8 (11%)	82,113,113	1.23	5 (6%)
18	CLA	a	813	1	69,73,73	1.16	9 (13%)	82,113,113	1.47	10 (12%)
18	CLA	a	826	1	69,73,73	1.18	8 (11%)	82,113,113	1.30	6 (7%)
26	A86	C	302	-	47,50,50	1.74	11 (23%)	51,76,76	3.28	24 (47%)
18	CLA	C	309	-	66,70,73	1.19	9 (13%)	78,109,113	1.48	8 (10%)
23	LMG	j	103	-	37,37,55	1.57	5 (13%)	45,45,63	3.67	7 (15%)
18	CLA	C	311	14	64,68,73	1.22	9 (14%)	76,107,113	1.35	6 (7%)
18	CLA	D	216	15	50,54,73	1.34	8 (16%)	59,90,113	1.41	5 (8%)
18	CLA	E	313	16	59,63,73	1.26	9 (15%)	70,101,113	1.39	6 (8%)
18	CLA	b	801	-	69,73,73	1.20	10 (14%)	82,113,113	1.45	7 (8%)
18	CLA	r	202	12	49,53,73	1.37	8 (16%)	58,89,113	1.40	5 (8%)
18	CLA	a	855	1	69,73,73	1.16	8 (11%)	82,113,113	1.31	7 (8%)
18	CLA	B	310	13	50,54,73	1.36	9 (18%)	59,90,113	1.39	4 (6%)
18	CLA	E	312	16	69,73,73	1.22	9 (13%)	82,113,113	1.34	7 (8%)
18	CLA	H	315	17	49,53,73	1.38	9 (18%)	58,89,113	1.45	4 (6%)
21	BCR	a	845	-	41,41,41	0.93	2 (4%)	56,56,56	1.96	20 (35%)
18	CLA	D	213	15	69,73,73	1.16	9 (13%)	82,113,113	1.27	5 (6%)
20	LHG	E	317	-	41,41,48	1.00	2 (4%)	44,47,54	1.22	5 (11%)
28	DD6	E	304	-	40,45,45	2.17	8 (20%)	51,67,67	3.34	15 (29%)
18	CLA	a	803	1	69,73,73	1.18	9 (13%)	82,113,113	1.49	7 (8%)
26	A86	B	301	-	47,50,50	1.50	7 (14%)	51,76,76	2.88	17 (33%)

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
18	CLA	a	809	18,1	1/1/14/20	13/36/112/115	-
26	A86	D	206	-	-	7/34/90/90	0/3/3/3
21	BCR	i	103	-	-	4/29/63/63	0/2/2/2
21	BCR	f	201	-	-	8/29/63/63	0/2/2/2
26	A86	b	847	-	-	15/34/90/90	0/3/3/3

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Mol	Type	Chain	Res	Link	Chirals	Torsions	Rings
18	CLA	b	849	-	1/1/15/20	16/39/115/115	-
18	CLA	C	312	14	1/1/14/20	12/35/111/115	-
18	CLA	D	217	-	1/1/10/20	4/10/86/115	-
18	CLA	b	814	-	1/1/15/20	11/39/115/115	-
18	CLA	a	815	1	1/1/15/20	15/39/115/115	-
18	CLA	a	816	1	1/1/11/20	10/15/91/115	-
20	LHG	b	835	18	-	32/53/53/53	-
18	CLA	E	311	16	1/1/11/20	6/17/93/115	-
18	CLA	a	825	1	1/1/14/20	13/33/109/115	-
28	DD6	B	303	-	-	0/26/80/80	0/3/3/3
18	CLA	a	828	1	1/1/15/20	10/39/115/115	-
21	BCR	m	101	-	-	7/29/63/63	0/2/2/2
18	CLA	b	817	2	1/1/14/20	17/33/109/115	-
18	CLA	B	311	13	1/1/15/20	12/39/115/115	-
26	A86	m	102	-	-	7/34/90/90	0/3/3/3
18	CLA	a	814	1	1/1/15/20	8/39/115/115	-
18	CLA	E	310	16	1/1/15/20	13/39/115/115	-
18	CLA	a	802	-	1/1/15/20	8/39/115/115	-
23	LMG	l	207	-	-	16/35/55/70	0/1/1/1
18	CLA	b	846	2	1/1/15/20	14/39/115/115	-
18	CLA	H	314	17	1/1/15/20	12/39/115/115	-
18	CLA	a	838	1	1/1/15/20	12/39/115/115	-
18	CLA	b	830	20	1/1/15/20	9/39/115/115	-
18	CLA	H	310	-	1/1/14/20	9/33/109/115	-
18	CLA	a	824	1	1/1/12/20	9/23/99/115	-
21	BCR	r	201	-	-	4/29/63/63	0/2/2/2
18	CLA	a	844	1	1/1/15/20	16/39/115/115	-
23	LMG	E	318	-	-	21/38/58/70	0/1/1/1
20	LHG	B	315	-	-	25/46/46/53	-
18	CLA	B	308	13	1/1/15/20	11/39/115/115	-
18	CLA	a	823	1	1/1/11/20	9/15/91/115	-
18	CLA	a	817	1	1/1/11/20	11/20/96/115	-
29	KC1	B	313	-	-	6/15/71/71	-
18	CLA	a	839	1	1/1/15/20	11/39/115/115	-

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Mol	Type	Chain	Res	Link	Chirals	Torsions	Rings
18	CLA	D	207	-	1/1/14/20	11/35/111/115	-
18	CLA	b	818	2	1/1/15/20	7/39/115/115	-
18	CLA	B	312	13	1/1/15/20	12/39/115/115	-
18	CLA	B	307	13	1/1/15/20	8/39/115/115	-
18	CLA	b	811	2	1/1/14/20	8/33/109/115	-
18	CLA	l	202	10	1/1/11/20	7/20/96/115	-
18	CLA	b	828	-	1/1/15/20	10/39/115/115	-
18	CLA	a	808	1	1/1/15/20	12/39/115/115	-
18	CLA	b	805	2	1/1/15/20	13/39/115/115	-
21	BCR	E	306	-	-	0/29/63/63	0/2/2/2
23	LMG	C	319	18	-	20/39/59/70	0/1/1/1
18	CLA	H	306	17	1/1/15/20	12/39/115/115	-
18	CLA	a	807	1	1/1/15/20	9/39/115/115	-
20	LHG	D	201	-	-	32/53/53/53	-
29	KC1	C	313	-	-	5/15/71/71	-
25	DGD	b	834	-	-	28/49/89/95	0/2/2/2
18	CLA	b	816	2	1/1/13/20	6/27/103/115	-
18	CLA	a	822	1	1/1/12/20	7/21/97/115	-
21	BCR	a	852	-	-	8/29/63/63	0/2/2/2
18	CLA	a	806	1	1/1/15/20	13/39/115/115	-
18	CLA	b	838	-	1/1/15/20	8/39/115/115	-
18	CLA	E	314	16	1/1/11/20	4/15/91/115	-
18	CLA	a	801	1	1/1/15/20	15/39/115/115	-
18	CLA	D	214	15	1/1/13/20	9/29/105/115	-
21	BCR	j	105	-	-	5/29/63/63	0/2/2/2
18	CLA	B	309	-	1/1/15/20	9/39/115/115	-
18	CLA	a	805	1	1/1/15/20	12/39/115/115	-
26	A86	H	302	-	-	5/34/90/90	0/3/3/3
21	BCR	b	832	-	-	0/29/63/63	0/2/2/2
18	CLA	C	316	14	1/1/10/20	4/10/86/115	-
18	CLA	a	846	-	1/1/15/20	8/39/115/115	-
18	CLA	b	825	-	1/1/11/20	4/15/91/115	-
18	CLA	b	824	-	1/1/15/20	17/39/115/115	-
21	BCR	j	101	-	-	2/29/63/63	0/2/2/2
28	DD6	E	308	-	-	1/26/80/80	0/3/3/3

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Mol	Type	Chain	Res	Link	Chirals	Torsions	Rings
18	CLA	b	844	2	1/1/14/20	10/36/112/115	-
18	CLA	a	843	-	1/1/15/20	10/39/115/115	-
18	CLA	B	314	13	1/1/10/20	2/10/86/115	-
18	CLA	E	315	16	1/1/15/20	11/39/115/115	-
18	CLA	D	215	-	1/1/10/20	4/10/86/115	-
26	A86	H	301	-	-	5/34/90/90	0/3/3/3
18	CLA	H	307	17	1/1/15/20	15/39/115/115	-
18	CLA	a	830	20	1/1/12/20	6/24/100/115	-
18	CLA	b	829	2	1/1/15/20	13/39/115/115	-
18	CLA	a	812	-	1/1/11/20	6/15/91/115	-
21	BCR	a	833	-	-	0/29/63/63	0/2/2/2
18	CLA	a	841	-	1/1/15/20	20/39/115/115	-
18	CLA	b	812	2	1/1/13/20	9/32/108/115	-
26	A86	C	304	-	-	4/34/90/90	0/3/3/3
18	CLA	C	310	14	1/1/11/20	8/17/93/115	-
18	CLA	a	854	-	1/1/15/20	16/39/115/115	-
26	A86	D	203	-	-	10/34/90/90	0/3/3/3
18	CLA	a	847	1	1/1/15/20	14/39/115/115	-
20	LHG	a	831	-	-	25/53/53/53	-
21	BCR	b	839	-	-	6/29/63/63	0/2/2/2
18	CLA	b	823	2	1/1/13/20	13/31/107/115	-
18	CLA	f	202	-	1/1/15/20	7/39/115/115	-
24	SF4	c	102	3	-	-	0/6/5/5
18	CLA	a	811	1	1/1/12/20	6/21/97/115	-
18	CLA	H	309	17	1/1/11/20	9/17/93/115	-
18	CLA	a	804	18,1	1/1/13/20	5/27/103/115	-
18	CLA	b	806	2	1/1/15/20	15/39/115/115	-
18	CLA	b	821	2	1/1/12/20	9/21/97/115	-
18	CLA	b	826	2	1/1/14/20	9/33/109/115	-
21	BCR	E	305	-	-	2/29/63/63	0/2/2/2
18	CLA	a	820	1	1/1/15/20	8/39/115/115	-
18	CLA	b	815	2	1/1/11/20	6/17/93/115	-
24	SF4	b	802	2	-	-	0/6/5/5
18	CLA	a	848	1	1/1/13/20	11/29/105/115	-
18	CLA	C	306	23	1/1/11/20	4/15/91/115	-

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Mol	Type	Chain	Res	Link	Chirals	Torsions	Rings
18	CLA	a	837	1	1/1/14/20	14/33/109/115	-
26	A86	B	304	-	-	3/34/90/90	0/3/3/3
18	CLA	E	309	16	1/1/15/20	13/39/115/115	-
21	BCR	f	205	-	-	2/29/63/63	0/2/2/2
28	DD6	D	205	-	-	3/26/80/80	0/3/3/3
19	PQN	a	829	-	-	2/23/43/43	0/2/2/2
23	LMG	D	202	-	-	17/41/61/70	0/1/1/1
18	CLA	H	313	17	1/1/15/20	18/39/115/115	-
27	ET4	l	206	-	-	3/25/67/67	0/2/2/2
18	CLA	b	841	2	1/1/15/20	7/39/115/115	-
18	CLA	a	819	1	1/1/15/20	4/39/115/115	-
28	DD6	H	303	-	-	3/26/80/80	0/3/3/3
29	KC1	H	312	17	-	9/15/71/71	-
26	A86	B	305	-	-	4/34/90/90	0/3/3/3
21	BCR	l	201	-	-	4/29/63/63	0/2/2/2
18	CLA	C	317	-	1/1/10/20	8/12/88/115	-
18	CLA	b	827	2	1/1/11/20	5/18/94/115	-
18	CLA	D	211	15	1/1/11/20	10/17/93/115	-
18	CLA	b	843	2	1/1/15/20	17/39/115/115	-
18	CLA	b	804	2	1/1/11/20	3/15/91/115	-
20	LHG	j	102	-	-	18/53/53/53	-
26	A86	C	305	-	-	4/34/90/90	0/3/3/3
18	CLA	H	311	17	1/1/13/20	8/30/106/115	-
18	CLA	l	203	10	1/1/15/20	16/39/115/115	-
18	CLA	b	813	2	1/1/13/20	12/27/103/115	-
18	CLA	C	307	14	1/1/15/20	5/39/115/115	-
18	CLA	b	837	2	1/1/14/20	13/33/109/115	-
19	PQN	b	831	-	-	3/23/43/43	0/2/2/2
26	A86	r	203	-	-	6/34/90/90	0/3/3/3
29	KC1	C	308	14	-	7/15/71/71	-
20	LHG	H	316	-	-	19/38/38/53	-
28	DD6	E	303	-	-	3/26/80/80	0/3/3/3
18	CLA	b	820	2	1/1/15/20	15/39/115/115	-
18	CLA	a	853	1	1/1/13/20	10/27/103/115	-

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Mol	Type	Chain	Res	Link	Chirals	Torsions	Rings
18	CLA	B	306	13	1/1/12/20	18/29/101/115	-
18	CLA	H	305	-	1/1/12/20	10/26/102/115	-
24	SF4	c	101	3	-	-	0/6/5/5
18	CLA	b	803	2	1/1/15/20	17/39/115/115	-
18	CLA	b	840	2	1/1/14/20	12/35/111/115	-
18	CLA	b	848	2	1/1/12/20	10/26/102/115	-
28	DD6	C	303	-	-	5/26/80/80	0/3/3/3
18	CLA	b	850	2	1/1/15/20	11/39/115/115	-
21	BCR	b	836	-	-	5/29/63/63	0/2/2/2
18	CLA	b	808	2	1/1/15/20	10/39/115/115	-
18	CLA	D	208	15	1/1/15/20	15/39/115/115	-
18	CLA	a	810	1	1/1/15/20	8/39/115/115	-
18	CLA	H	308	17	1/1/15/20	10/39/115/115	-
21	BCR	b	833	-	-	6/29/63/63	0/2/2/2
18	CLA	b	809	2	1/1/13/20	4/27/103/115	-
18	CLA	f	204	6	1/1/11/20	5/15/91/115	-
18	CLA	E	316	-	1/1/13/20	4/29/105/115	-
18	CLA	j	104	9	1/1/10/20	4/10/86/115	-
18	CLA	a	818	1	1/1/12/20	4/23/99/115	-
18	CLA	a	840	-	1/1/15/20	13/39/115/115	-
18	CLA	D	209	15	1/1/11/20	11/20/96/115	-
20	LHG	E	301	-	-	33/53/53/53	-
18	CLA	D	212	15	1/1/10/20	2/12/88/115	-
18	CLA	C	315	14	1/1/10/20	6/12/88/115	-
21	BCR	a	834	-	-	4/29/63/63	0/2/2/2
23	LMG	C	301	-	-	14/26/46/70	0/1/1/1
18	CLA	b	842	2	1/1/15/20	15/39/115/115	-
26	A86	E	302	-	-	5/34/90/90	0/3/3/3
18	CLA	C	314	14	1/1/10/20	4/10/86/115	-
22	SQD	a	849	-	-	19/33/33/69	-
18	CLA	b	819	2	1/1/15/20	14/39/115/115	-
18	CLA	i	101	-	1/1/14/20	10/33/109/115	-
18	CLA	b	822	2	1/1/15/20	15/39/115/115	-
18	CLA	f	203	-	1/1/15/20	11/39/115/115	-

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Mol	Type	Chain	Res	Link	Chirals	Torsions	Rings
18	CLA	a	842	-	1/1/15/20	9/39/115/115	-
21	BCR	a	850	-	-	4/29/63/63	0/2/2/2
18	CLA	l	205	-	1/1/11/20	3/15/91/115	-
28	DD6	E	307	-	-	11/26/80/80	0/3/3/3
18	CLA	a	836	1	1/1/15/20	12/39/115/115	-
18	CLA	b	810	2	1/1/15/20	7/39/115/115	-
18	CLA	a	821	1	1/1/15/20	13/39/115/115	-
21	BCR	l	204	-	-	8/29/63/63	0/2/2/2
18	CLA	C	318	14	1/1/10/20	6/12/88/115	-
20	LHG	a	832	18	-	13/31/31/53	-
26	A86	D	204	-	-	4/34/90/90	0/3/3/3
21	BCR	i	102	-	-	0/29/63/63	0/2/2/2
26	A86	B	302	-	-	5/34/90/90	0/3/3/3
18	CLA	b	845	-	1/1/15/20	13/39/115/115	-
18	CLA	a	827	1	1/1/15/20	7/39/115/115	-
26	A86	H	304	-	-	4/34/90/90	0/3/3/3
18	CLA	D	210	-	1/1/13/20	6/29/105/115	-
20	LHG	a	835	-	-	19/36/36/53	-
23	LMG	a	851	-	-	16/28/48/70	0/1/1/1
18	CLA	b	807	2	1/1/15/20	9/39/115/115	-
18	CLA	a	813	1	1/1/15/20	24/39/115/115	-
18	CLA	a	826	1	1/1/15/20	11/39/115/115	-
26	A86	C	302	-	-	5/34/90/90	0/3/3/3
18	CLA	C	309	-	1/1/14/20	14/36/112/115	-
23	LMG	j	103	-	-	16/32/52/70	0/1/1/1
18	CLA	C	311	14	1/1/14/20	13/33/109/115	-
18	CLA	D	216	15	1/1/11/20	5/17/93/115	-
18	CLA	E	313	16	1/1/13/20	9/27/103/115	-
18	CLA	b	801	-	1/1/15/20	8/39/115/115	-
18	CLA	r	202	12	1/1/11/20	11/15/91/115	-
18	CLA	a	855	1	1/1/15/20	12/39/115/115	-
18	CLA	B	310	13	1/1/11/20	6/17/93/115	-
18	CLA	E	312	16	1/1/15/20	10/39/115/115	-
18	CLA	H	315	17	1/1/11/20	6/15/91/115	-

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Mol	Type	Chain	Res	Link	Chirals	Torsions	Rings
21	BCR	a	845	-	-	2/29/63/63	0/2/2/2
18	CLA	D	213	15	1/1/15/20	10/39/115/115	-
20	LHG	E	317	-	-	5/46/46/53	-
28	DD6	E	304	-	-	6/26/80/80	0/3/3/3
18	CLA	a	803	1	1/1/15/20	13/39/115/115	-
26	A86	B	301	-	-	10/34/90/90	0/3/3/3

All (1547) bond length outliers are listed below:

Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
22	a	849	SQD	C6-S	-12.69	1.59	1.77
23	E	318	LMG	O2-C2	-10.20	1.17	1.43
28	E	307	DD6	C30-C31	-9.01	1.25	1.42
28	E	304	DD6	C30-C31	-8.98	1.25	1.42
28	C	303	DD6	C30-C31	-8.95	1.25	1.42
23	D	202	LMG	O2-C2	-8.88	1.21	1.43
28	B	303	DD6	C30-C31	-8.49	1.26	1.42
28	H	303	DD6	C30-C31	-8.47	1.26	1.42
28	E	303	DD6	C30-C31	-8.35	1.26	1.42
28	E	308	DD6	C30-C31	-8.33	1.26	1.42
19	a	829	PQN	C3-C2	7.87	1.49	1.35
19	b	831	PQN	C3-C2	7.72	1.49	1.35
28	D	205	DD6	C30-C31	-7.56	1.27	1.42
22	a	849	SQD	O9-S	7.03	1.64	1.45
23	j	103	LMG	O2-C2	-5.66	1.28	1.43
23	a	851	LMG	O2-C2	-5.64	1.29	1.43
29	B	313	KC1	MG-ND	-5.53	1.94	2.05
29	H	312	KC1	MG-ND	-5.52	1.94	2.05
22	a	849	SQD	O47-C45	-5.27	1.39	1.46
29	H	312	KC1	MG-NB	-5.15	1.95	2.05
26	b	847	A86	O4-C38	5.10	1.46	1.35
29	B	313	KC1	MG-NB	-5.04	1.95	2.05
29	C	313	KC1	MG-ND	-4.99	1.95	2.05
29	C	308	KC1	MG-ND	-4.95	1.96	2.05
26	B	302	A86	C30-C31	-4.93	1.24	1.30
26	B	301	A86	O4-C38	4.92	1.46	1.35
26	H	302	A86	C30-C31	-4.89	1.24	1.30
26	r	203	A86	C30-C31	-4.88	1.24	1.30
26	m	102	A86	O4-C38	4.81	1.45	1.35
19	a	829	PQN	C10-C5	4.79	1.48	1.40
26	E	302	A86	O4-C38	4.78	1.45	1.35

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
19	b	831	PQN	C10-C5	4.77	1.48	1.40
22	a	849	SQD	O8-S	4.76	1.65	1.47
29	C	308	KC1	MG-NB	-4.72	1.96	2.05
26	H	301	A86	C30-C31	-4.69	1.25	1.30
26	E	302	A86	C30-C31	-4.69	1.25	1.30
26	C	305	A86	C30-C31	-4.66	1.25	1.30
26	r	203	A86	C20-C15	4.66	1.54	1.48
26	C	302	A86	C30-C31	-4.65	1.25	1.30
26	B	305	A86	C30-C31	-4.64	1.25	1.30
26	C	305	A86	O4-C38	4.62	1.45	1.35
29	C	313	KC1	MG-NB	-4.62	1.96	2.05
26	D	203	A86	O4-C38	4.61	1.45	1.35
23	C	301	LMG	O2-C2	4.60	1.54	1.43
26	H	301	A86	O4-C38	4.60	1.45	1.35
26	H	304	A86	O4-C38	4.57	1.45	1.35
26	D	206	A86	C30-C31	-4.55	1.25	1.30
26	C	304	A86	C30-C31	-4.55	1.25	1.30
26	D	204	A86	C30-C31	-4.51	1.25	1.30
26	D	204	A86	O4-C38	4.51	1.45	1.35
26	B	304	A86	O4-C38	4.42	1.45	1.35
26	H	304	A86	C30-C31	-4.41	1.25	1.30
22	a	849	SQD	O49-C7	4.38	1.40	1.22
26	D	206	A86	O4-C38	4.33	1.44	1.35
26	C	302	A86	O4-C38	4.32	1.44	1.35
26	C	302	A86	C30-C29	-4.31	1.24	1.31
26	H	304	A86	C20-C15	4.25	1.53	1.48
26	B	304	A86	C30-C31	-4.23	1.25	1.30
26	H	302	A86	O4-C38	4.22	1.44	1.35
20	B	315	LHG	O8-C23	4.21	1.45	1.33
20	E	317	LHG	O8-C23	4.20	1.45	1.33
26	B	302	A86	O4-C38	4.20	1.44	1.35
26	C	305	A86	C20-C15	4.19	1.53	1.48
26	B	302	A86	C30-C29	-4.19	1.25	1.31
26	C	304	A86	C30-C29	-4.15	1.25	1.31
27	l	206	ET4	C08-C07	4.15	1.45	1.33
26	r	203	A86	C30-C29	-4.15	1.25	1.31
25	b	834	DGD	O1G-C1A	4.14	1.45	1.33
26	C	304	A86	O4-C38	4.14	1.44	1.35
26	H	302	A86	C30-C29	-4.13	1.25	1.31
26	E	302	A86	C30-C29	-4.12	1.25	1.31
26	r	203	A86	O4-C38	4.11	1.44	1.35
26	B	305	A86	O4-C38	4.10	1.44	1.35

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
26	D	204	A86	C30-C29	-4.09	1.25	1.31
28	H	303	DD6	C28-C27	-4.09	1.48	1.50
20	E	317	LHG	O7-C7	4.07	1.45	1.34
26	H	301	A86	C30-C29	-4.05	1.25	1.31
26	b	847	A86	C30-C31	-4.05	1.25	1.30
28	E	304	DD6	C21-C20	-4.01	1.45	1.51
26	D	206	A86	C30-C29	-4.01	1.25	1.31
26	C	305	A86	C30-C29	-4.00	1.25	1.31
26	B	301	A86	C30-C31	-3.99	1.25	1.30
26	D	203	A86	O1-C20	-3.97	1.41	1.46
26	B	305	A86	C30-C29	-3.97	1.25	1.31
25	b	834	DGD	O2G-C1B	3.95	1.45	1.34
26	D	203	A86	C20-C15	3.95	1.53	1.48
26	H	304	A86	C30-C29	-3.93	1.25	1.31
26	b	847	A86	C20-C15	3.92	1.53	1.48
26	B	304	A86	C30-C29	-3.92	1.25	1.31
28	B	303	DD6	C28-C27	-3.90	1.48	1.50
20	B	315	LHG	O7-C7	3.84	1.45	1.34
28	B	303	DD6	C-C1	-3.83	1.43	1.50
26	H	301	A86	C20-C15	3.82	1.53	1.48
26	C	304	A86	C20-C15	3.81	1.53	1.48
26	C	302	A86	C20-C15	3.79	1.53	1.48
26	B	302	A86	O1-C20	-3.78	1.41	1.46
26	D	203	A86	C30-C31	-3.75	1.26	1.30
26	B	305	A86	C20-C15	3.75	1.53	1.48
26	b	847	A86	C30-C29	-3.74	1.25	1.31
28	H	303	DD6	C-C1	-3.71	1.43	1.50
18	b	842	CLA	C1D-ND	3.71	1.42	1.37
26	D	206	A86	O1-C20	-3.69	1.41	1.46
18	a	826	CLA	C1D-ND	3.69	1.42	1.37
26	D	206	A86	C20-C15	3.68	1.52	1.48
28	D	205	DD6	C28-C27	-3.66	1.48	1.50
18	a	814	CLA	C1D-ND	3.65	1.42	1.37
18	a	844	CLA	C1D-ND	3.65	1.42	1.37
26	C	302	A86	O1-C20	-3.64	1.41	1.46
18	a	838	CLA	C1D-ND	3.63	1.42	1.37
26	E	302	A86	C20-C15	3.63	1.52	1.48
29	H	312	KC1	C4B-NB	-3.62	1.33	1.37
26	D	204	A86	C20-C15	3.62	1.52	1.48
18	b	818	CLA	C1D-ND	3.62	1.42	1.37
18	D	211	CLA	C1D-ND	3.62	1.42	1.37
18	b	841	CLA	C1D-ND	3.61	1.42	1.37

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
18	a	821	CLA	C1D-ND	3.60	1.42	1.37
18	b	809	CLA	C1D-ND	3.58	1.42	1.37
26	B	302	A86	C20-C15	3.57	1.52	1.48
26	H	302	A86	C20-C15	3.57	1.52	1.48
26	m	102	A86	C20-C15	3.56	1.52	1.48
18	E	313	CLA	C1D-ND	3.55	1.42	1.37
18	b	813	CLA	C1D-ND	3.55	1.42	1.37
18	b	825	CLA	C1D-ND	3.55	1.42	1.37
28	E	304	DD6	C19-C18	-3.55	1.47	1.52
18	a	817	CLA	C1D-ND	3.55	1.42	1.37
18	b	815	CLA	C1D-ND	3.55	1.42	1.37
26	H	302	A86	O1-C20	-3.54	1.41	1.46
18	a	812	CLA	C1D-ND	3.54	1.42	1.37
26	B	301	A86	C30-C29	-3.54	1.26	1.31
18	C	315	CLA	C1D-ND	3.53	1.42	1.37
18	f	204	CLA	C1D-ND	3.53	1.42	1.37
18	a	848	CLA	C1D-ND	3.53	1.42	1.37
18	D	209	CLA	C1D-ND	3.52	1.42	1.37
20	D	201	LHG	C26-C25	-3.52	1.34	1.51
18	b	805	CLA	C1D-ND	3.52	1.42	1.37
26	D	203	A86	C30-C29	-3.51	1.26	1.31
18	a	853	CLA	C1D-ND	3.50	1.42	1.37
18	D	215	CLA	C1D-ND	3.49	1.42	1.37
18	a	830	CLA	C1D-ND	3.49	1.42	1.37
18	B	308	CLA	C1D-ND	3.49	1.42	1.37
26	B	304	A86	O1-C20	-3.49	1.41	1.46
18	b	817	CLA	C1D-ND	3.49	1.42	1.37
18	b	838	CLA	C1D-ND	3.49	1.42	1.37
18	C	306	CLA	C1D-ND	3.49	1.42	1.37
18	D	214	CLA	C1D-ND	3.49	1.42	1.37
18	b	812	CLA	C1D-ND	3.48	1.42	1.37
18	B	312	CLA	C1D-ND	3.48	1.42	1.37
18	a	819	CLA	C1D-ND	3.48	1.42	1.37
18	a	824	CLA	C1D-ND	3.48	1.42	1.37
23	C	319	LMG	O2-C2	-3.48	1.34	1.43
23	D	202	LMG	O6-C1	3.47	1.50	1.41
18	a	816	CLA	C1D-ND	3.47	1.42	1.37
18	a	806	CLA	C1D-ND	3.46	1.42	1.37
18	E	311	CLA	C1D-ND	3.46	1.42	1.37
18	H	309	CLA	C1D-ND	3.46	1.42	1.37
18	a	811	CLA	C1D-ND	3.46	1.42	1.37
18	B	310	CLA	C1D-ND	3.46	1.42	1.37

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
18	b	807	CLA	C1D-ND	3.45	1.42	1.37
18	b	806	CLA	C1D-ND	3.45	1.42	1.37
18	b	827	CLA	C1D-ND	3.45	1.42	1.37
18	a	823	CLA	C1D-ND	3.45	1.42	1.37
18	b	810	CLA	C1D-ND	3.44	1.42	1.37
18	E	310	CLA	C1D-ND	3.44	1.42	1.37
18	b	850	CLA	C1D-ND	3.44	1.42	1.37
18	b	846	CLA	C1D-ND	3.44	1.42	1.37
18	E	309	CLA	C1D-ND	3.44	1.42	1.37
18	a	827	CLA	C1D-ND	3.44	1.42	1.37
18	r	202	CLA	C1D-ND	3.43	1.42	1.37
18	b	814	CLA	C1D-ND	3.43	1.42	1.37
18	b	823	CLA	C1D-ND	3.43	1.42	1.37
18	a	822	CLA	C1D-ND	3.43	1.42	1.37
18	E	315	CLA	C1D-ND	3.42	1.42	1.37
18	a	828	CLA	C1D-ND	3.42	1.42	1.37
18	b	826	CLA	C1D-ND	3.42	1.42	1.37
18	H	315	CLA	C1D-ND	3.42	1.42	1.37
18	b	819	CLA	C1D-ND	3.42	1.42	1.37
18	a	809	CLA	C1D-ND	3.42	1.42	1.37
18	l	205	CLA	C1D-ND	3.41	1.42	1.37
18	B	307	CLA	C1D-ND	3.41	1.42	1.37
18	j	104	CLA	C1D-ND	3.41	1.42	1.37
18	D	213	CLA	C1D-ND	3.40	1.42	1.37
28	E	304	DD6	C36-C31	-3.40	1.31	1.35
23	j	103	LMG	O6-C1	3.40	1.50	1.41
18	i	101	CLA	C1D-ND	3.39	1.42	1.37
18	a	843	CLA	C1D-ND	3.39	1.42	1.37
26	H	304	A86	O1-C20	-3.39	1.41	1.46
18	a	847	CLA	C1D-ND	3.38	1.42	1.37
26	m	102	A86	O1-C20	-3.38	1.41	1.46
18	b	849	CLA	C1D-ND	3.38	1.42	1.37
18	a	810	CLA	C1D-ND	3.38	1.42	1.37
18	B	309	CLA	C1D-ND	3.38	1.42	1.37
18	a	808	CLA	C1D-ND	3.37	1.42	1.37
18	b	828	CLA	C1D-ND	3.37	1.42	1.37
18	b	845	CLA	C1D-ND	3.37	1.42	1.37
18	b	816	CLA	C1D-ND	3.37	1.42	1.37
23	l	207	LMG	O6-C1	3.37	1.50	1.41
18	b	840	CLA	C1D-ND	3.37	1.42	1.37
18	a	855	CLA	C1D-ND	3.37	1.42	1.37
18	D	212	CLA	C1D-ND	3.37	1.42	1.37

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
26	B	305	A86	O1-C20	-3.36	1.41	1.46
18	a	804	CLA	C1D-ND	3.36	1.42	1.37
18	a	815	CLA	C1D-ND	3.36	1.42	1.37
18	a	805	CLA	C1D-ND	3.35	1.42	1.37
18	b	844	CLA	C1D-ND	3.35	1.42	1.37
18	a	846	CLA	C1D-ND	3.35	1.42	1.37
26	C	304	A86	O1-C20	-3.35	1.41	1.46
18	b	808	CLA	C1D-ND	3.35	1.42	1.37
18	f	203	CLA	C1D-ND	3.34	1.42	1.37
18	b	830	CLA	C1D-ND	3.34	1.42	1.37
20	E	301	LHG	C26-C25	-3.33	1.35	1.51
26	m	102	A86	C30-C29	-3.33	1.26	1.31
18	a	820	CLA	C1D-ND	3.32	1.42	1.37
18	a	826	CLA	C4B-NB	3.32	1.42	1.37
18	b	822	CLA	C1D-ND	3.31	1.42	1.37
20	a	831	LHG	C26-C25	-3.31	1.35	1.51
28	B	303	DD6	O1-C20	-3.31	1.41	1.46
18	b	848	CLA	C1D-ND	3.31	1.42	1.37
26	H	301	A86	O1-C20	-3.30	1.41	1.46
18	f	202	CLA	C1D-ND	3.30	1.42	1.37
26	C	305	A86	O1-C20	-3.30	1.41	1.46
26	m	102	A86	C30-C31	-3.30	1.26	1.30
18	a	836	CLA	C1D-ND	3.29	1.42	1.37
20	b	835	LHG	C26-C25	-3.29	1.35	1.51
28	H	303	DD6	O1-C20	-3.29	1.42	1.46
18	a	839	CLA	C1D-ND	3.28	1.42	1.37
28	E	304	DD6	O1-C20	-3.28	1.42	1.46
28	D	205	DD6	C36-C31	-3.28	1.31	1.35
20	a	835	LHG	C26-C25	-3.27	1.35	1.51
18	B	311	CLA	C1D-ND	3.26	1.42	1.37
26	B	301	A86	C20-C15	3.26	1.52	1.48
26	H	302	A86	C32-C31	-3.25	1.49	1.54
18	b	821	CLA	C1D-ND	3.25	1.42	1.37
18	E	316	CLA	C1D-ND	3.24	1.42	1.37
18	C	317	CLA	MG-NB	-3.24	1.99	2.05
23	E	318	LMG	O6-C1	3.23	1.50	1.41
18	a	802	CLA	C1D-ND	3.22	1.42	1.37
20	j	102	LHG	C26-C25	-3.22	1.35	1.51
18	a	804	CLA	C4B-NB	3.21	1.42	1.37
29	B	313	KC1	C4B-NB	-3.21	1.33	1.37
26	B	304	A86	C20-C15	3.21	1.52	1.48
26	B	302	A86	C32-C31	-3.19	1.49	1.54

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
18	b	827	CLA	C4B-NB	3.18	1.42	1.37
26	r	203	A86	O1-C20	-3.18	1.42	1.46
26	E	302	A86	O1-C20	-3.18	1.42	1.46
18	a	816	CLA	C4B-NB	3.17	1.42	1.37
28	D	205	DD6	O1-C20	-3.17	1.42	1.46
26	D	204	A86	O1-C20	-3.17	1.42	1.46
18	b	825	CLA	C4B-NB	3.15	1.42	1.37
26	C	305	A86	C32-C31	-3.15	1.49	1.54
23	E	318	LMG	O1-C1	-3.14	1.35	1.40
18	D	217	CLA	C1D-ND	3.14	1.42	1.37
26	B	305	A86	C32-C31	-3.13	1.49	1.54
29	H	312	KC1	CBA-CGA	-3.12	1.40	1.48
18	b	841	CLA	C4B-NB	3.12	1.42	1.37
23	C	301	LMG	O6-C1	3.11	1.49	1.41
18	B	311	CLA	C4B-NB	3.11	1.41	1.37
18	a	838	CLA	C4B-NB	3.10	1.41	1.37
18	b	809	CLA	C4B-NB	3.08	1.41	1.37
18	b	821	CLA	C4B-NB	3.08	1.41	1.37
18	j	104	CLA	C4B-NB	3.07	1.41	1.37
18	b	822	CLA	C4B-NB	3.07	1.41	1.37
29	H	312	KC1	C1B-NB	-3.07	1.33	1.37
18	B	314	CLA	C1D-ND	3.06	1.41	1.37
23	C	301	LMG	O8-C28	3.06	1.42	1.33
18	D	209	CLA	C4B-NB	3.06	1.41	1.37
20	b	835	LHG	O8-C6	-3.06	1.38	1.45
18	b	805	CLA	C4B-NB	3.06	1.41	1.37
18	B	310	CLA	C4B-NB	3.06	1.41	1.37
18	C	306	CLA	C4B-NB	3.05	1.41	1.37
18	C	315	CLA	C4B-NB	3.05	1.41	1.37
18	b	816	CLA	C4B-NB	3.05	1.41	1.37
18	D	215	CLA	C4B-NB	3.04	1.41	1.37
18	a	812	CLA	C4B-NB	3.04	1.41	1.37
18	a	821	CLA	C4B-NB	3.04	1.41	1.37
18	b	806	CLA	C4B-NB	3.04	1.41	1.37
18	a	820	CLA	C4B-NB	3.03	1.41	1.37
18	a	823	CLA	C4B-NB	3.03	1.41	1.37
22	a	849	SQD	O47-C7	3.02	1.39	1.33
18	a	830	CLA	C4B-NB	3.02	1.41	1.37
18	H	309	CLA	C4B-NB	3.02	1.41	1.37
18	D	214	CLA	C4B-NB	3.01	1.41	1.37
18	a	828	CLA	C4B-NB	3.01	1.41	1.37
18	E	312	CLA	MG-NB	-3.01	1.99	2.05

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
18	a	814	CLA	C4B-NB	3.00	1.41	1.37
18	E	310	CLA	C4B-NB	3.00	1.41	1.37
18	a	827	CLA	C4B-NB	3.00	1.41	1.37
18	a	854	CLA	MG-NB	-3.00	1.99	2.05
18	b	814	CLA	C4B-NB	2.99	1.41	1.37
18	a	808	CLA	C4B-NB	2.99	1.41	1.37
18	b	828	CLA	C4B-NB	2.99	1.41	1.37
18	a	805	CLA	C4B-NB	2.99	1.41	1.37
18	a	853	CLA	C4B-NB	2.99	1.41	1.37
18	b	844	CLA	C4B-NB	2.99	1.41	1.37
18	b	803	CLA	MG-NB	-2.99	1.99	2.05
18	b	850	CLA	C4B-NB	2.99	1.41	1.37
18	b	823	CLA	C4B-NB	2.99	1.41	1.37
18	a	847	CLA	C4B-NB	2.98	1.41	1.37
18	E	311	CLA	C4B-NB	2.98	1.41	1.37
18	b	840	CLA	C4B-NB	2.98	1.41	1.37
18	b	817	CLA	C4B-NB	2.97	1.41	1.37
18	b	810	CLA	C4B-NB	2.97	1.41	1.37
18	a	815	CLA	C4B-NB	2.97	1.41	1.37
18	C	310	CLA	C1D-ND	2.96	1.41	1.37
29	H	312	KC1	C4A-C3A	-2.96	1.38	1.44
28	B	303	DD6	C17-C16	-2.96	1.50	1.54
18	b	849	CLA	C4B-NB	2.96	1.41	1.37
18	i	101	CLA	C4B-NB	2.96	1.41	1.37
18	b	807	CLA	C4B-NB	2.96	1.41	1.37
23	D	202	LMG	O8-C28	2.96	1.42	1.33
18	b	845	CLA	C4B-NB	2.95	1.41	1.37
21	a	850	BCR	C30-C25	-2.95	1.50	1.53
18	f	203	CLA	C4B-NB	2.95	1.41	1.37
18	f	204	CLA	C4B-NB	2.95	1.41	1.37
18	D	211	CLA	C4B-NB	2.95	1.41	1.37
18	E	315	CLA	C4B-NB	2.94	1.41	1.37
18	a	810	CLA	C4B-NB	2.94	1.41	1.37
18	a	822	CLA	C4B-NB	2.94	1.41	1.37
18	D	213	CLA	C4B-NB	2.94	1.41	1.37
29	B	313	KC1	C1B-NB	-2.94	1.34	1.37
18	a	844	CLA	C4B-NB	2.93	1.41	1.37
23	C	319	LMG	O1-C1	-2.93	1.35	1.40
23	D	202	LMG	O7-C8	-2.93	1.39	1.46
18	D	210	CLA	C1D-ND	2.92	1.41	1.37
23	D	202	LMG	O1-C1	-2.92	1.35	1.40
18	E	309	CLA	C4B-NB	2.92	1.41	1.37

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
26	H	301	A86	C32-C31	-2.92	1.49	1.54
18	b	830	CLA	C4B-NB	2.92	1.41	1.37
18	f	202	CLA	C4B-NB	2.92	1.41	1.37
18	r	202	CLA	C4B-NB	2.92	1.41	1.37
18	b	801	CLA	CMB-C2B	-2.91	1.44	1.50
18	H	315	CLA	C4B-NB	2.91	1.41	1.37
18	C	316	CLA	C1D-ND	2.91	1.41	1.37
18	B	307	CLA	C4B-NB	2.91	1.41	1.37
18	E	316	CLA	C4B-NB	2.91	1.41	1.37
29	B	313	KC1	CBA-CGA	-2.91	1.41	1.48
28	D	205	DD6	C17-C16	-2.91	1.50	1.54
18	a	848	CLA	C4B-NB	2.91	1.41	1.37
18	b	848	CLA	C4B-NB	2.91	1.41	1.37
18	b	819	CLA	C4B-NB	2.90	1.41	1.37
26	H	304	A86	C32-C31	-2.90	1.49	1.54
18	a	809	CLA	C4B-NB	2.90	1.41	1.37
18	D	212	CLA	C4B-NB	2.89	1.41	1.37
18	H	305	CLA	C1D-ND	2.89	1.41	1.37
23	l	207	LMG	O8-C28	2.89	1.41	1.33
18	b	808	CLA	C4B-NB	2.89	1.41	1.37
18	b	818	CLA	C4B-NB	2.89	1.41	1.37
18	a	843	CLA	C4B-NB	2.88	1.41	1.37
18	l	205	CLA	C4B-NB	2.88	1.41	1.37
18	b	815	CLA	C4B-NB	2.87	1.41	1.37
18	a	802	CLA	C4B-NB	2.87	1.41	1.37
18	a	837	CLA	MG-NB	-2.87	2.00	2.05
18	b	812	CLA	C4B-NB	2.86	1.41	1.37
18	b	846	CLA	C4B-NB	2.86	1.41	1.37
26	H	302	A86	C-C1	-2.86	1.45	1.50
18	a	817	CLA	C4B-NB	2.85	1.41	1.37
18	E	313	CLA	C4B-NB	2.85	1.41	1.37
28	C	303	DD6	C36-C31	-2.85	1.31	1.35
18	b	813	CLA	C4B-NB	2.85	1.41	1.37
18	b	801	CLA	MG-NB	-2.85	2.00	2.05
18	b	824	CLA	CMD-C2D	-2.85	1.44	1.50
18	a	855	CLA	C4B-NB	2.85	1.41	1.37
21	a	850	BCR	C1-C6	-2.84	1.50	1.53
18	a	836	CLA	C4B-NB	2.84	1.41	1.37
18	H	311	CLA	MG-NB	-2.84	2.00	2.05
18	H	314	CLA	MG-NB	-2.84	2.00	2.05
29	H	312	KC1	C1D-ND	-2.84	1.34	1.37
18	B	308	CLA	C4B-NB	2.84	1.41	1.37

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
18	C	307	CLA	C1D-ND	2.84	1.41	1.37
18	b	842	CLA	C4B-NB	2.83	1.41	1.37
18	a	819	CLA	C4B-NB	2.83	1.41	1.37
18	B	312	CLA	C4B-NB	2.83	1.41	1.37
18	C	314	CLA	MG-NB	-2.83	2.00	2.05
26	B	302	A86	C-C1	-2.83	1.45	1.50
20	a	831	LHG	O8-C6	-2.83	1.38	1.45
18	H	308	CLA	C1D-ND	2.82	1.41	1.37
18	b	838	CLA	C4B-NB	2.82	1.41	1.37
18	b	820	CLA	CMD-C2D	-2.81	1.45	1.50
23	a	851	LMG	O6-C1	2.81	1.49	1.41
18	b	826	CLA	C4B-NB	2.81	1.41	1.37
18	l	203	CLA	MG-NB	-2.81	2.00	2.05
18	E	316	CLA	C1B-C2B	2.81	1.49	1.43
28	D	205	DD6	C-C1	-2.80	1.45	1.50
18	C	314	CLA	C1D-ND	2.80	1.41	1.37
18	C	318	CLA	MG-NB	-2.80	2.00	2.05
26	D	204	A86	C32-C31	-2.80	1.49	1.54
21	b	839	BCR	C30-C25	-2.80	1.50	1.53
18	C	309	CLA	C1D-ND	2.80	1.41	1.37
18	H	307	CLA	MG-NB	-2.79	2.00	2.05
18	B	309	CLA	C4B-NB	2.79	1.41	1.37
18	a	803	CLA	MG-NB	-2.79	2.00	2.05
18	B	306	CLA	C1D-ND	2.79	1.41	1.37
18	D	211	CLA	C1B-C2B	2.79	1.49	1.43
18	C	311	CLA	C1D-ND	2.79	1.41	1.37
18	b	824	CLA	MG-NB	-2.79	2.00	2.05
18	a	806	CLA	C4B-NB	2.79	1.41	1.37
18	l	202	CLA	C1D-ND	2.78	1.41	1.37
28	H	303	DD6	C36-C31	-2.78	1.31	1.35
18	a	824	CLA	C4B-NB	2.77	1.41	1.37
23	j	103	LMG	O7-C8	-2.77	1.40	1.46
23	j	103	LMG	O8-C28	2.77	1.41	1.33
18	a	839	CLA	C4B-NB	2.77	1.41	1.37
18	b	843	CLA	MG-NB	-2.76	2.00	2.05
26	b	847	A86	O1-C20	-2.76	1.42	1.46
18	l	202	CLA	CMB-C2B	-2.76	1.45	1.50
18	D	216	CLA	C1D-ND	2.76	1.41	1.37
18	a	846	CLA	C4B-NB	2.76	1.41	1.37
18	a	825	CLA	CMB-C2B	-2.75	1.45	1.50
18	b	850	CLA	C1B-C2B	2.75	1.49	1.43
18	a	848	CLA	C1B-C2B	2.75	1.49	1.43

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
18	a	840	CLA	MG-NB	-2.75	2.00	2.05
18	a	841	CLA	MG-NB	-2.75	2.00	2.05
18	b	848	CLA	C1B-C2B	2.74	1.49	1.43
18	b	815	CLA	C1B-C2B	2.74	1.49	1.43
18	H	306	CLA	MG-NB	-2.74	2.00	2.05
18	H	307	CLA	C1D-ND	2.74	1.41	1.37
26	B	304	A86	C15-C16	-2.74	1.51	1.55
18	B	306	CLA	C4B-NB	2.74	1.41	1.37
18	C	315	CLA	C1B-C2B	2.74	1.49	1.43
18	a	854	CLA	CMB-C2B	-2.74	1.45	1.50
18	a	801	CLA	C1D-ND	2.74	1.41	1.37
18	a	805	CLA	C1B-C2B	2.74	1.49	1.43
18	a	827	CLA	C1B-C2B	2.73	1.49	1.43
27	l	206	ET4	C37-C38	-2.73	1.46	1.51
18	B	314	CLA	MG-NB	-2.73	2.00	2.05
18	a	811	CLA	C4B-NB	2.73	1.41	1.37
18	a	817	CLA	C1B-C2B	2.72	1.49	1.43
18	b	821	CLA	C1B-C2B	2.72	1.49	1.43
18	H	315	CLA	C1B-C2B	2.72	1.49	1.43
18	C	318	CLA	CMD-C2D	-2.72	1.45	1.50
18	b	829	CLA	MG-NB	-2.72	2.00	2.05
18	b	825	CLA	C1B-C2B	2.71	1.49	1.43
18	H	310	CLA	MG-NB	-2.71	2.00	2.05
18	b	820	CLA	MG-NB	-2.71	2.00	2.05
18	b	803	CLA	CMD-C2D	-2.71	1.45	1.50
18	a	828	CLA	C1B-C2B	2.71	1.49	1.43
18	b	823	CLA	C1B-C2B	2.71	1.49	1.43
18	f	204	CLA	C1B-C2B	2.71	1.49	1.43
18	C	317	CLA	CMD-C2D	-2.71	1.45	1.50
18	a	844	CLA	C1B-C2B	2.70	1.49	1.43
18	b	820	CLA	CMB-C2B	-2.70	1.45	1.50
18	b	806	CLA	C1B-C2B	2.70	1.49	1.43
20	D	201	LHG	O8-C6	-2.70	1.39	1.45
18	b	807	CLA	C1B-C2B	2.70	1.49	1.43
18	B	308	CLA	C1B-C2B	2.70	1.49	1.43
18	a	842	CLA	C1D-ND	2.70	1.41	1.37
18	D	208	CLA	MG-NB	-2.70	2.00	2.05
18	C	312	CLA	C1D-ND	2.69	1.41	1.37
21	b	839	BCR	C1-C6	-2.69	1.50	1.53
18	f	202	CLA	C1B-C2B	2.69	1.49	1.43
18	b	809	CLA	C1B-C2B	2.69	1.49	1.43
18	E	312	CLA	C1D-ND	2.69	1.41	1.37

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
26	B	301	A86	O1-C20	-2.69	1.42	1.46
18	b	837	CLA	CMB-C2B	-2.68	1.45	1.50
18	a	818	CLA	MG-NB	-2.68	2.00	2.05
18	D	215	CLA	C1B-C2B	2.68	1.49	1.43
18	D	214	CLA	C1B-C2B	2.68	1.49	1.43
18	a	840	CLA	CMB-C2B	-2.68	1.45	1.50
29	B	313	KC1	C4A-C3A	-2.68	1.39	1.44
18	a	842	CLA	MG-NB	-2.68	2.00	2.05
18	b	812	CLA	C1B-C2B	2.68	1.49	1.43
18	a	830	CLA	C1B-C2B	2.68	1.49	1.43
18	b	838	CLA	C1B-C2B	2.68	1.49	1.43
18	f	203	CLA	C1B-C2B	2.68	1.49	1.43
18	a	812	CLA	C1B-C2B	2.68	1.49	1.43
18	r	202	CLA	C1B-C2B	2.68	1.49	1.43
29	C	313	KC1	CBA-CGA	-2.67	1.41	1.48
18	C	306	CLA	C1B-C2B	2.67	1.49	1.43
18	a	801	CLA	CMD-C2D	-2.67	1.45	1.50
18	b	811	CLA	CMC-C2C	-2.67	1.45	1.50
18	a	806	CLA	C1B-C2B	2.67	1.49	1.43
18	b	840	CLA	C1B-C2B	2.67	1.49	1.43
20	a	831	LHG	O7-C5	-2.67	1.40	1.46
18	a	813	CLA	C1D-ND	2.67	1.41	1.37
18	B	309	CLA	C1B-C2B	2.67	1.49	1.43
18	b	803	CLA	MG-ND	-2.67	2.00	2.05
18	b	805	CLA	C1B-C2B	2.67	1.49	1.43
18	b	846	CLA	C1B-C2B	2.67	1.49	1.43
18	E	313	CLA	C1B-C2B	2.67	1.49	1.43
20	H	316	LHG	O8-C6	-2.66	1.39	1.45
18	a	824	CLA	C1B-C2B	2.66	1.49	1.43
18	b	829	CLA	CMB-C2B	-2.66	1.45	1.50
18	a	811	CLA	C1B-C2B	2.66	1.49	1.43
18	b	827	CLA	C1B-C2B	2.66	1.49	1.43
26	C	302	A86	C32-C31	-2.66	1.50	1.54
18	C	316	CLA	MG-NB	-2.66	2.00	2.05
29	C	313	KC1	C4B-NB	-2.66	1.34	1.37
18	a	837	CLA	CMB-C2B	-2.66	1.45	1.50
18	l	202	CLA	CMD-C2D	-2.66	1.45	1.50
18	b	816	CLA	C1B-C2B	2.66	1.49	1.43
18	a	822	CLA	C1B-C2B	2.66	1.49	1.43
18	E	312	CLA	CMD-C2D	-2.66	1.45	1.50
18	b	818	CLA	C1B-C2B	2.66	1.49	1.43
18	E	310	CLA	C1B-C2B	2.65	1.49	1.43

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
18	a	821	CLA	C1B-C2B	2.65	1.49	1.43
18	l	202	CLA	MG-NB	-2.65	2.00	2.05
18	b	843	CLA	CMB-C2B	-2.65	1.45	1.50
18	a	841	CLA	CMD-C2D	-2.65	1.45	1.50
18	D	217	CLA	C4B-NB	2.65	1.41	1.37
18	b	829	CLA	CMD-C2D	-2.65	1.45	1.50
18	a	836	CLA	C1B-C2B	2.65	1.49	1.43
18	b	817	CLA	C1B-C2B	2.65	1.49	1.43
23	E	318	LMG	O7-C8	-2.65	1.40	1.46
18	E	309	CLA	C1B-C2B	2.64	1.49	1.43
18	B	310	CLA	C1B-C2B	2.64	1.49	1.43
18	a	840	CLA	C1D-ND	2.64	1.41	1.37
18	a	837	CLA	CMD-C2D	-2.64	1.45	1.50
18	b	844	CLA	C1B-C2B	2.64	1.49	1.43
23	C	319	LMG	O8-C28	2.64	1.41	1.33
18	D	216	CLA	MG-NB	-2.64	2.00	2.05
18	a	801	CLA	MG-NB	-2.64	2.00	2.05
18	b	811	CLA	MG-NB	-2.63	2.00	2.05
18	a	837	CLA	CMC-C2C	-2.63	1.45	1.50
18	l	205	CLA	C1B-C2B	2.63	1.49	1.43
18	a	823	CLA	C1B-C2B	2.63	1.49	1.43
18	a	807	CLA	C1D-ND	2.63	1.41	1.37
18	B	314	CLA	C4B-NB	2.63	1.41	1.37
18	a	825	CLA	MG-NB	-2.63	2.00	2.05
18	j	104	CLA	C1B-C2B	2.63	1.49	1.43
23	C	319	LMG	O6-C1	2.63	1.48	1.41
18	a	847	CLA	C1B-C2B	2.63	1.49	1.43
18	b	845	CLA	C1B-C2B	2.63	1.49	1.43
18	H	308	CLA	MG-NB	-2.63	2.00	2.05
18	b	842	CLA	C1B-C2B	2.63	1.49	1.43
18	l	203	CLA	CMB-C2B	-2.63	1.45	1.50
18	b	822	CLA	C1B-C2B	2.62	1.49	1.43
18	H	309	CLA	C1B-C2B	2.62	1.49	1.43
18	a	810	CLA	C1B-C2B	2.62	1.49	1.43
26	C	302	A86	C13-C11	-2.62	1.44	1.49
18	b	837	CLA	C1D-ND	2.62	1.41	1.37
23	l	207	LMG	O7-C8	-2.61	1.40	1.46
18	a	816	CLA	C1B-C2B	2.61	1.49	1.43
18	D	208	CLA	CMC-C2C	-2.61	1.45	1.50
18	B	312	CLA	C1B-C2B	2.61	1.49	1.43
26	D	203	A86	C13-C11	-2.61	1.44	1.49
18	a	843	CLA	C1B-C2B	2.61	1.49	1.43

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
18	b	829	CLA	CMC-C2C	-2.61	1.45	1.50
18	b	830	CLA	C1B-C2B	2.61	1.49	1.43
18	a	802	CLA	C1B-C2B	2.61	1.49	1.43
18	a	819	CLA	C1B-C2B	2.61	1.49	1.43
18	b	849	CLA	C1B-C2B	2.61	1.49	1.43
18	b	801	CLA	CMC-C2C	-2.61	1.45	1.50
18	a	809	CLA	C1B-C2B	2.60	1.49	1.43
18	b	810	CLA	C1B-C2B	2.60	1.49	1.43
18	a	853	CLA	C1B-C2B	2.60	1.49	1.43
18	a	818	CLA	CMD-C2D	-2.60	1.45	1.50
18	b	819	CLA	C1B-C2B	2.60	1.49	1.43
18	b	804	CLA	CMB-C2B	-2.60	1.45	1.50
18	a	814	CLA	C1B-C2B	2.60	1.49	1.43
18	D	212	CLA	C1B-C2B	2.60	1.49	1.43
18	B	306	CLA	MG-NB	-2.60	2.00	2.05
18	D	209	CLA	C1B-C2B	2.60	1.49	1.43
18	a	842	CLA	CMB-C2B	-2.60	1.45	1.50
18	b	829	CLA	C1D-ND	2.60	1.41	1.37
20	E	301	LHG	O7-C5	-2.60	1.40	1.46
18	a	807	CLA	MG-NB	-2.60	2.00	2.05
18	b	824	CLA	CMC-C2C	-2.59	1.45	1.50
18	b	804	CLA	MG-NB	-2.59	2.00	2.05
26	D	204	A86	C13-C11	-2.59	1.44	1.49
18	a	837	CLA	C1B-NB	-2.59	1.34	1.37
18	D	210	CLA	CMB-C2B	-2.59	1.45	1.50
18	i	101	CLA	C1B-C2B	2.59	1.49	1.43
21	a	833	BCR	C1-C6	-2.59	1.50	1.53
18	C	310	CLA	C1B-C2B	2.59	1.49	1.43
18	H	311	CLA	CMC-C2C	-2.59	1.45	1.50
18	E	311	CLA	C1B-C2B	2.59	1.49	1.43
23	E	318	LMG	O8-C28	2.59	1.40	1.33
18	a	840	CLA	CMD-C2D	-2.58	1.45	1.50
18	a	842	CLA	CMD-C2D	-2.58	1.45	1.50
18	b	804	CLA	C1D-ND	2.58	1.41	1.37
21	l	201	BCR	C30-C25	-2.58	1.50	1.53
18	a	807	CLA	CMB-C2B	-2.58	1.45	1.50
18	H	313	CLA	MG-NB	-2.58	2.00	2.05
18	H	310	CLA	CMD-C2D	-2.58	1.45	1.50
20	E	301	LHG	O8-C6	-2.58	1.39	1.45
18	a	808	CLA	C1B-C2B	2.58	1.49	1.43
18	a	841	CLA	C1D-ND	2.58	1.41	1.37
20	a	835	LHG	O8-C6	-2.58	1.39	1.45

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
18	a	826	CLA	C1B-C2B	2.58	1.49	1.43
26	C	304	A86	C13-C11	-2.58	1.44	1.49
20	b	835	LHG	O7-C5	-2.58	1.40	1.46
18	a	801	CLA	CMB-C2B	-2.58	1.45	1.50
18	a	841	CLA	CMB-C2B	-2.57	1.45	1.50
18	E	315	CLA	C1B-C2B	2.57	1.49	1.43
18	b	808	CLA	C1B-C2B	2.57	1.49	1.43
20	a	835	LHG	O7-C5	-2.57	1.40	1.46
18	D	217	CLA	C1B-C2B	2.57	1.49	1.43
18	a	803	CLA	C1D-ND	2.57	1.41	1.37
18	b	813	CLA	C1B-C2B	2.57	1.49	1.43
21	a	845	BCR	C30-C25	-2.57	1.50	1.53
18	B	311	CLA	C1B-C2B	2.56	1.49	1.43
18	C	309	CLA	MG-NB	-2.56	2.00	2.05
18	C	312	CLA	MG-NB	-2.56	2.00	2.05
18	a	846	CLA	C1B-C2B	2.56	1.49	1.43
18	C	310	CLA	MG-NB	-2.56	2.00	2.05
18	a	840	CLA	CMC-C2C	-2.56	1.45	1.50
18	b	843	CLA	CMD-C2D	-2.56	1.45	1.50
18	B	306	CLA	CBD-CAD	2.56	1.57	1.51
23	C	319	LMG	O7-C8	-2.56	1.40	1.46
18	b	811	CLA	CMD-C2D	-2.56	1.45	1.50
18	C	314	CLA	CMD-C2D	-2.55	1.45	1.50
18	C	312	CLA	CMB-C2B	-2.55	1.45	1.50
18	a	854	CLA	CMC-C2C	-2.55	1.45	1.50
18	H	311	CLA	CMD-C2D	-2.54	1.45	1.50
18	b	837	CLA	MG-NB	-2.54	2.00	2.05
18	b	824	CLA	CMB-C2B	-2.54	1.45	1.50
18	a	820	CLA	C1B-C2B	2.54	1.49	1.43
18	E	314	CLA	C1D-ND	2.54	1.41	1.37
26	E	302	A86	C32-C31	-2.54	1.50	1.54
18	H	305	CLA	MG-NB	-2.54	2.00	2.05
18	b	820	CLA	C1D-ND	2.54	1.41	1.37
18	a	813	CLA	MG-NB	-2.54	2.00	2.05
18	b	841	CLA	C1B-C2B	2.54	1.49	1.43
18	B	306	CLA	C1B-C2B	2.54	1.49	1.43
18	D	208	CLA	CMB-C2B	-2.53	1.45	1.50
18	a	813	CLA	CMD-C2D	-2.53	1.45	1.50
18	a	841	CLA	CMC-C2C	-2.53	1.45	1.50
18	H	306	CLA	CMB-C2B	-2.53	1.45	1.50
18	H	308	CLA	CMD-C2D	-2.53	1.45	1.50
29	C	308	KC1	C4B-NB	-2.53	1.34	1.37

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
18	a	825	CLA	CMD-C2D	-2.53	1.45	1.50
18	b	811	CLA	CMB-C2B	-2.53	1.45	1.50
18	a	838	CLA	C1B-C2B	2.53	1.49	1.43
28	E	304	DD6	C-C1	-2.53	1.45	1.50
18	a	804	CLA	C1B-C2B	2.53	1.49	1.43
18	D	207	CLA	MG-NB	-2.52	2.00	2.05
18	C	311	CLA	C4B-NB	2.52	1.41	1.37
18	D	216	CLA	CMD-C2D	-2.52	1.45	1.50
18	H	307	CLA	CMC-C2C	-2.52	1.45	1.50
18	a	855	CLA	C1B-C2B	2.52	1.49	1.43
18	a	807	CLA	CAA-C2A	-2.52	1.49	1.54
26	B	304	A86	C32-C31	-2.52	1.50	1.54
18	a	804	CLA	CHC-C1C	2.52	1.43	1.38
18	C	317	CLA	CMB-C2B	-2.52	1.45	1.50
18	C	311	CLA	CMB-C2B	-2.52	1.45	1.50
18	a	807	CLA	CMD-C2D	-2.52	1.45	1.50
18	D	208	CLA	CMD-C2D	-2.52	1.45	1.50
18	H	314	CLA	CMB-C2B	-2.51	1.45	1.50
18	b	837	CLA	CMD-C2D	-2.51	1.45	1.50
18	C	317	CLA	MG-ND	-2.51	2.00	2.05
18	a	803	CLA	CMB-C2B	-2.51	1.45	1.50
18	b	803	CLA	CMC-C2C	-2.51	1.45	1.50
26	b	847	A86	C32-C31	-2.51	1.50	1.54
18	a	803	CLA	CMD-C2D	-2.51	1.45	1.50
18	C	311	CLA	MG-NB	-2.51	2.00	2.05
18	E	312	CLA	CMB-C2B	-2.51	1.45	1.50
18	a	839	CLA	C1B-C2B	2.51	1.49	1.43
18	a	837	CLA	MG-ND	-2.51	2.00	2.05
29	C	308	KC1	CBA-CGA	-2.51	1.42	1.48
18	D	216	CLA	CMC-C2C	-2.50	1.45	1.50
18	D	217	CLA	MG-NB	-2.50	2.00	2.05
18	E	312	CLA	MG-ND	-2.50	2.00	2.05
18	a	803	CLA	CMC-C2C	-2.50	1.45	1.50
18	I	203	CLA	CMD-C2D	-2.50	1.45	1.50
18	D	207	CLA	CMB-C2B	-2.50	1.45	1.50
26	b	847	A86	C17-C16	-2.50	1.50	1.54
18	C	316	CLA	CMD-C2D	-2.50	1.45	1.50
18	H	313	CLA	C1D-ND	2.49	1.41	1.37
18	C	309	CLA	CMD-C2D	-2.49	1.45	1.50
18	b	803	CLA	CMB-C2B	-2.49	1.45	1.50
18	a	818	CLA	CMB-C2B	-2.49	1.45	1.50
18	b	801	CLA	CMD-C2D	-2.49	1.45	1.50

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
18	b	824	CLA	C1D-ND	2.49	1.41	1.37
18	b	826	CLA	C1B-C2B	2.49	1.49	1.43
26	r	203	A86	C32-C31	-2.49	1.50	1.54
18	H	311	CLA	C1D-ND	2.49	1.41	1.37
18	b	843	CLA	CMC-C2C	-2.49	1.45	1.50
18	a	813	CLA	CMC-C2C	-2.49	1.45	1.50
18	H	307	CLA	CMD-C2D	-2.48	1.45	1.50
21	a	845	BCR	C1-C6	-2.48	1.50	1.53
18	E	314	CLA	CMD-C2D	-2.48	1.45	1.50
18	H	305	CLA	C4B-NB	2.48	1.41	1.37
18	H	314	CLA	C1D-ND	2.48	1.41	1.37
18	H	307	CLA	CMB-C2B	-2.48	1.45	1.50
28	E	304	DD6	C17-C16	-2.48	1.50	1.54
18	D	207	CLA	C1D-ND	2.48	1.41	1.37
18	l	203	CLA	CMC-C2C	-2.48	1.45	1.50
18	D	207	CLA	CMD-C2D	-2.48	1.45	1.50
18	D	208	CLA	C1D-ND	2.48	1.41	1.37
28	H	303	DD6	C21-C20	-2.48	1.48	1.51
28	H	303	DD6	C17-C16	-2.47	1.50	1.54
18	a	854	CLA	C1B-NB	-2.47	1.34	1.37
18	D	213	CLA	C1B-C2B	2.47	1.48	1.43
18	b	814	CLA	C1B-C2B	2.47	1.48	1.43
26	C	304	A86	C32-C31	-2.47	1.50	1.54
18	B	307	CLA	C1B-C2B	2.47	1.48	1.43
23	C	301	LMG	O7-C8	-2.47	1.40	1.46
18	a	825	CLA	CMC-C2C	-2.47	1.45	1.50
18	l	202	CLA	CMC-C2C	-2.47	1.45	1.50
18	H	305	CLA	CMD-C2D	-2.46	1.45	1.50
18	E	314	CLA	MG-NB	-2.46	2.00	2.05
18	H	306	CLA	CMD-C2D	-2.46	1.45	1.50
18	b	804	CLA	CMC-C2C	-2.46	1.45	1.50
18	b	843	CLA	C1D-ND	2.46	1.41	1.37
18	C	309	CLA	CMB-C2B	-2.46	1.45	1.50
18	D	210	CLA	CMD-C2D	-2.46	1.45	1.50
18	a	815	CLA	C1B-C2B	2.46	1.48	1.43
18	H	306	CLA	C1D-ND	2.46	1.41	1.37
18	b	813	CLA	CMC-C2C	-2.46	1.45	1.50
18	a	807	CLA	CMC-C2C	-2.46	1.45	1.50
26	D	204	A86	C17-C16	-2.46	1.50	1.54
18	H	310	CLA	CMC-C2C	-2.45	1.45	1.50
18	a	854	CLA	CMD-C2D	-2.45	1.45	1.50
18	a	837	CLA	C1D-ND	2.45	1.41	1.37

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
23	C	301	LMG	O7-C10	2.45	1.41	1.34
18	C	307	CLA	CMC-C2C	-2.44	1.45	1.50
18	b	804	CLA	CMD-C2D	-2.44	1.45	1.50
20	D	201	LHG	O7-C5	-2.44	1.40	1.46
18	b	837	CLA	CMC-C2C	-2.44	1.45	1.50
18	H	310	CLA	C1D-ND	2.44	1.41	1.37
18	H	314	CLA	CMC-C2C	-2.44	1.45	1.50
18	a	818	CLA	C1D-ND	2.44	1.41	1.37
18	H	314	CLA	CMD-C2D	-2.44	1.45	1.50
18	a	801	CLA	MG-ND	-2.44	2.01	2.05
23	a	851	LMG	O8-C28	2.44	1.40	1.33
18	C	312	CLA	CMD-C2D	-2.43	1.45	1.50
27	l	206	ET4	C19-C18	2.43	1.51	1.46
26	H	302	A86	C17-C16	-2.43	1.50	1.54
21	l	201	BCR	C1-C6	-2.43	1.50	1.53
18	a	840	CLA	MG-ND	-2.43	2.01	2.05
20	a	832	LHG	O8-C6	-2.43	1.39	1.45
21	f	201	BCR	C1-C6	-2.43	1.50	1.53
23	C	319	LMG	O7-C10	2.43	1.41	1.34
18	a	818	CLA	CMC-C2C	-2.43	1.45	1.50
18	C	318	CLA	CMB-C2B	-2.42	1.45	1.50
18	D	216	CLA	CMB-C2B	-2.42	1.45	1.50
26	D	206	A86	C32-C31	-2.42	1.50	1.54
18	C	307	CLA	C1B-NB	-2.42	1.34	1.37
18	D	213	CLA	C3B-C4B	2.42	1.49	1.42
18	E	312	CLA	CMC-C2C	-2.41	1.45	1.50
18	C	316	CLA	C4B-NB	2.41	1.41	1.37
18	b	803	CLA	C1D-ND	2.41	1.41	1.37
18	b	801	CLA	C1D-ND	2.41	1.41	1.37
18	H	308	CLA	CMC-C2C	-2.41	1.45	1.50
18	C	309	CLA	CMC-C2C	-2.41	1.45	1.50
18	b	828	CLA	C3B-C4B	2.40	1.49	1.42
18	D	210	CLA	MG-NB	-2.40	2.01	2.05
28	H	303	DD6	C35-C36	-2.40	1.47	1.51
18	D	207	CLA	CMC-C2C	-2.40	1.45	1.50
18	a	825	CLA	C1B-NB	-2.40	1.34	1.37
29	C	308	KC1	C1B-NB	-2.40	1.34	1.37
18	B	314	CLA	C1B-C2B	2.40	1.48	1.43
18	H	308	CLA	CMB-C2B	-2.40	1.45	1.50
26	C	305	A86	C17-C16	-2.40	1.50	1.54
20	a	832	LHG	O7-C5	-2.40	1.41	1.46
18	B	311	CLA	C3B-C4B	2.40	1.49	1.42

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
26	B	304	A86	C-C1	-2.40	1.46	1.50
20	a	835	LHG	O7-C7	2.40	1.40	1.35
18	D	213	CLA	CHC-C1C	2.40	1.43	1.38
18	a	815	CLA	C3B-C4B	2.40	1.49	1.42
18	a	825	CLA	C4B-NB	2.40	1.41	1.37
28	H	303	DD6	C19-C18	-2.39	1.48	1.52
26	b	847	A86	C13-C11	-2.39	1.45	1.49
18	D	216	CLA	C1B-C2B	2.39	1.48	1.43
18	b	824	CLA	MG-ND	-2.39	2.01	2.05
18	l	202	CLA	MG-ND	-2.39	2.01	2.05
18	b	814	CLA	C3B-C4B	2.39	1.49	1.42
18	C	314	CLA	CMB-C2B	-2.39	1.45	1.50
18	E	314	CLA	CMB-C2B	-2.39	1.45	1.50
18	a	842	CLA	MG-ND	-2.39	2.01	2.05
18	H	305	CLA	CMB-C2B	-2.39	1.45	1.50
18	b	801	CLA	MG-ND	-2.39	2.01	2.05
18	a	841	CLA	C1B-NB	-2.38	1.34	1.37
26	B	302	A86	C17-C16	-2.38	1.50	1.54
18	a	801	CLA	CMC-C2C	-2.38	1.45	1.50
18	a	838	CLA	C3B-C4B	2.38	1.49	1.42
18	C	307	CLA	CMB-C2B	-2.38	1.45	1.50
28	B	303	DD6	C36-C31	-2.38	1.32	1.35
18	a	841	CLA	MG-ND	-2.37	2.01	2.05
23	a	851	LMG	O7-C10	2.37	1.41	1.34
18	C	307	CLA	MG-NB	-2.37	2.01	2.05
18	b	811	CLA	C1D-ND	2.37	1.41	1.37
18	B	310	CLA	CMD-C2D	-2.37	1.45	1.50
18	a	820	CLA	CMB-C2B	-2.37	1.45	1.50
22	a	849	SQD	O48-C23	2.37	1.40	1.33
18	a	804	CLA	C3B-C4B	2.37	1.49	1.42
20	j	102	LHG	O8-C6	-2.37	1.39	1.45
18	E	313	CLA	C3B-C4B	2.37	1.49	1.42
18	b	841	CLA	C3B-C4B	2.37	1.49	1.42
18	b	804	CLA	C1B-NB	-2.36	1.34	1.37
18	b	828	CLA	C1B-C2B	2.36	1.48	1.43
26	r	203	A86	C-C1	-2.36	1.46	1.50
18	H	313	CLA	CMD-C2D	-2.36	1.45	1.50
18	D	210	CLA	CMC-C2C	-2.36	1.45	1.50
18	b	843	CLA	MG-ND	-2.36	2.01	2.05
18	a	815	CLA	CHC-C1C	2.36	1.43	1.38
26	B	302	A86	O1-C15	-2.36	1.41	1.45
18	C	314	CLA	CMC-C2C	-2.35	1.45	1.50

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
18	H	313	CLA	CMB-C2B	-2.35	1.45	1.50
18	l	203	CLA	C1B-NB	-2.35	1.34	1.37
18	a	853	CLA	CHC-C1C	2.35	1.43	1.38
18	C	318	CLA	C1D-ND	2.35	1.41	1.37
18	C	309	CLA	C4B-NB	2.35	1.40	1.37
26	C	305	A86	C13-C11	-2.35	1.45	1.49
18	H	310	CLA	MG-ND	-2.35	2.01	2.05
18	a	825	CLA	C1D-ND	2.35	1.40	1.37
18	b	841	CLA	CHC-C1C	2.35	1.43	1.38
18	E	309	CLA	CHC-C1C	2.35	1.43	1.38
29	B	313	KC1	C1D-ND	-2.35	1.34	1.37
18	a	813	CLA	CMB-C2B	-2.35	1.46	1.50
18	C	312	CLA	CMC-C2C	-2.35	1.46	1.50
18	B	312	CLA	C3B-C4B	2.34	1.49	1.42
18	B	314	CLA	MG-ND	-2.34	2.01	2.05
18	C	315	CLA	C3B-C4B	2.34	1.49	1.42
18	a	819	CLA	C3B-C4B	2.34	1.49	1.42
26	C	302	A86	O1-C15	-2.34	1.41	1.45
18	H	310	CLA	C1B-C2B	2.34	1.48	1.43
18	H	313	CLA	C1B-C2B	2.34	1.48	1.43
18	a	842	CLA	CMC-C2C	-2.34	1.46	1.50
28	E	307	DD6	C36-C31	-2.34	1.32	1.35
20	j	102	LHG	O8-C23	2.34	1.40	1.33
18	E	314	CLA	C4B-NB	2.34	1.40	1.37
18	E	315	CLA	C3B-C4B	2.34	1.49	1.42
18	D	211	CLA	C3B-C4B	2.34	1.49	1.42
18	b	814	CLA	CHC-C1C	2.33	1.43	1.38
18	b	826	CLA	C3B-C4B	2.33	1.49	1.42
18	E	311	CLA	C3B-C4B	2.33	1.49	1.42
20	H	316	LHG	O7-C7	2.33	1.40	1.34
20	a	835	LHG	O8-C23	2.33	1.40	1.33
18	a	818	CLA	MG-ND	-2.33	2.01	2.05
18	a	854	CLA	MG-ND	-2.33	2.01	2.05
18	C	316	CLA	CMB-C2B	-2.33	1.46	1.50
18	D	207	CLA	C4B-NB	2.33	1.40	1.37
18	l	205	CLA	C3B-C4B	2.33	1.49	1.42
18	H	308	CLA	C1B-C2B	2.33	1.48	1.43
18	i	101	CLA	C3B-C4B	2.33	1.49	1.42
18	b	824	CLA	C1B-NB	-2.33	1.34	1.37
18	C	314	CLA	C4B-NB	2.33	1.40	1.37
18	b	849	CLA	CMD-C2D	-2.33	1.46	1.50
18	H	311	CLA	CMB-C2B	-2.33	1.46	1.50

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
20	a	832	LHG	O7-C7	2.33	1.40	1.34
20	j	102	LHG	O7-C7	2.33	1.40	1.34
18	a	846	CLA	C3B-C4B	2.33	1.49	1.42
18	C	306	CLA	C3B-C4B	2.32	1.49	1.42
18	D	212	CLA	C3B-C4B	2.32	1.49	1.42
18	H	308	CLA	C4B-NB	2.32	1.40	1.37
18	a	853	CLA	C3B-C4B	2.32	1.49	1.42
18	a	839	CLA	C3B-C4B	2.32	1.49	1.42
23	C	301	LMG	O1-C1	-2.32	1.36	1.40
18	C	317	CLA	C1D-ND	2.32	1.40	1.37
18	E	310	CLA	C3B-C4B	2.32	1.49	1.42
18	a	847	CLA	C3B-C4B	2.32	1.49	1.42
18	b	803	CLA	C1B-C2B	2.32	1.48	1.43
18	E	314	CLA	CMC-C2C	-2.32	1.46	1.50
28	E	308	DD6	O1-C20	-2.32	1.43	1.46
26	H	301	A86	C15-C16	-2.31	1.51	1.55
18	C	311	CLA	C1B-C2B	2.31	1.48	1.43
18	H	310	CLA	CMB-C2B	-2.31	1.46	1.50
18	b	822	CLA	C3B-C4B	2.31	1.49	1.42
18	C	316	CLA	CMC-C2C	-2.31	1.46	1.50
18	H	313	CLA	CMC-C2C	-2.31	1.46	1.50
18	B	314	CLA	CMB-C2B	-2.31	1.46	1.50
18	b	809	CLA	C3B-C4B	2.31	1.49	1.42
26	D	206	A86	O1-C15	-2.31	1.42	1.45
18	a	808	CLA	CMD-C2D	-2.31	1.46	1.50
18	b	843	CLA	C1B-NB	-2.31	1.34	1.37
18	H	306	CLA	CMC-C2C	-2.30	1.46	1.50
18	H	315	CLA	C3B-C4B	2.30	1.49	1.42
18	b	828	CLA	CHC-C1C	2.30	1.43	1.38
18	E	309	CLA	C3B-C4B	2.30	1.49	1.42
18	b	808	CLA	CMB-C2B	-2.30	1.46	1.50
18	b	811	CLA	C1B-NB	-2.30	1.34	1.37
18	i	101	CLA	CHC-C1C	2.30	1.43	1.38
18	a	843	CLA	C3B-C4B	2.30	1.49	1.42
18	j	104	CLA	C3B-C4B	2.30	1.49	1.42
18	D	214	CLA	C3B-C4B	2.30	1.49	1.42
18	D	210	CLA	C1B-C2B	2.30	1.48	1.43
26	C	302	A86	C15-C16	-2.30	1.51	1.55
18	a	838	CLA	CHC-C1C	2.30	1.43	1.38
18	a	826	CLA	C3B-C4B	2.30	1.49	1.42
18	b	846	CLA	C3B-C4B	2.30	1.49	1.42
18	D	215	CLA	C3B-C4B	2.30	1.49	1.42

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
18	E	312	CLA	CAA-C2A	-2.30	1.49	1.54
18	b	844	CLA	C3B-C4B	2.30	1.49	1.42
18	H	305	CLA	C1B-C2B	2.30	1.48	1.43
18	f	202	CLA	C3B-C4B	2.29	1.49	1.42
20	j	102	LHG	O7-C5	-2.29	1.41	1.46
18	b	829	CLA	C1B-C2B	2.29	1.48	1.43
18	E	316	CLA	CMD-C2D	-2.29	1.46	1.50
18	a	840	CLA	C1B-NB	-2.29	1.34	1.37
18	b	821	CLA	CMD-C2D	-2.29	1.46	1.50
29	C	313	KC1	C4A-C3A	-2.29	1.39	1.44
18	a	810	CLA	C3B-C4B	2.29	1.49	1.42
18	b	848	CLA	CMD-C2D	-2.29	1.46	1.50
18	f	203	CLA	C3B-C4B	2.29	1.49	1.42
18	C	315	CLA	CHC-C1C	2.29	1.43	1.38
18	H	313	CLA	C4B-NB	2.29	1.40	1.37
18	C	307	CLA	MG-ND	-2.29	2.01	2.05
18	C	311	CLA	CMC-C2C	-2.28	1.46	1.50
18	b	812	CLA	C3B-C4B	2.28	1.49	1.42
18	B	307	CLA	C3B-C4B	2.28	1.49	1.42
18	b	826	CLA	CHC-C1C	2.28	1.43	1.38
20	a	832	LHG	O8-C23	2.28	1.40	1.33
29	H	312	KC1	C1B-C2B	-2.28	1.40	1.45
18	a	827	CLA	C3B-C4B	2.28	1.49	1.42
18	b	830	CLA	C3B-C4B	2.28	1.49	1.42
18	b	827	CLA	C3B-C4B	2.28	1.49	1.42
18	H	309	CLA	C3B-C4B	2.28	1.49	1.42
18	D	209	CLA	C3B-C4B	2.28	1.49	1.42
18	b	813	CLA	MG-NB	-2.28	2.01	2.05
18	D	207	CLA	MG-ND	-2.28	2.01	2.05
18	a	814	CLA	C3B-C4B	2.28	1.49	1.42
18	B	307	CLA	CHC-C1C	2.28	1.43	1.38
18	l	205	CLA	CHC-C1C	2.27	1.43	1.38
18	C	318	CLA	MG-ND	-2.27	2.01	2.05
18	a	806	CLA	C3B-C4B	2.27	1.49	1.42
18	a	816	CLA	C3B-C4B	2.27	1.49	1.42
18	a	812	CLA	C3B-C4B	2.27	1.49	1.42
18	C	310	CLA	C4B-NB	2.27	1.40	1.37
18	H	311	CLA	C1B-C2B	2.27	1.48	1.43
18	H	311	CLA	MG-ND	-2.27	2.01	2.05
23	a	851	LMG	O7-C8	-2.27	1.41	1.46
18	b	811	CLA	MG-ND	-2.27	2.01	2.05
18	b	849	CLA	C3B-C4B	2.27	1.49	1.42

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
29	B	313	KC1	C1B-C2B	-2.27	1.40	1.45
18	E	309	CLA	CMC-C2C	-2.27	1.46	1.50
18	a	855	CLA	CHC-C1C	2.27	1.43	1.38
26	B	301	A86	C13-C11	-2.26	1.45	1.49
18	a	836	CLA	CMD-C2D	-2.26	1.46	1.50
18	D	212	CLA	CHC-C1C	2.26	1.43	1.38
26	D	206	A86	O4-C34	-2.26	1.41	1.46
18	b	845	CLA	C3B-C4B	2.26	1.49	1.42
18	j	104	CLA	CHC-C1C	2.26	1.43	1.38
18	a	830	CLA	CMC-C2C	-2.26	1.46	1.50
26	r	203	A86	O4-C34	-2.26	1.41	1.46
18	b	810	CLA	C3B-C4B	2.26	1.49	1.42
18	C	307	CLA	CMD-C2D	-2.26	1.46	1.50
18	b	837	CLA	MG-ND	-2.26	2.01	2.05
18	E	315	CLA	CHC-C1C	2.26	1.43	1.38
18	b	818	CLA	C3B-C4B	2.26	1.49	1.42
18	H	306	CLA	MG-ND	-2.26	2.01	2.05
18	b	848	CLA	C3B-C4B	2.25	1.49	1.42
18	f	204	CLA	C3B-C4B	2.25	1.49	1.42
18	b	849	CLA	CMB-C2B	-2.25	1.46	1.50
29	C	313	KC1	C1B-NB	-2.25	1.34	1.37
18	b	840	CLA	C3B-C4B	2.25	1.49	1.42
18	a	821	CLA	C3B-C4B	2.25	1.49	1.42
18	a	820	CLA	CMD-C2D	-2.25	1.46	1.50
18	B	310	CLA	C3B-C4B	2.25	1.49	1.42
18	b	822	CLA	CHC-C1C	2.25	1.43	1.38
18	a	817	CLA	C3B-C4B	2.25	1.49	1.42
18	a	855	CLA	C3B-C4B	2.25	1.49	1.42
18	a	846	CLA	CHC-C1C	2.25	1.43	1.38
18	r	202	CLA	C3B-C4B	2.25	1.49	1.42
18	C	311	CLA	CMD-C2D	-2.25	1.46	1.50
18	a	811	CLA	C3B-C4B	2.25	1.49	1.42
18	H	314	CLA	MG-ND	-2.25	2.01	2.05
18	B	310	CLA	CHC-C1C	2.25	1.43	1.38
18	C	317	CLA	CMC-C2C	-2.24	1.46	1.50
18	E	314	CLA	C1B-C2B	2.24	1.48	1.43
18	B	308	CLA	C3B-C4B	2.24	1.49	1.42
18	b	820	CLA	CMC-C2C	-2.24	1.46	1.50
18	a	809	CLA	C3B-C4B	2.24	1.49	1.42
18	l	203	CLA	C1D-ND	2.24	1.40	1.37
18	b	804	CLA	MG-ND	-2.24	2.01	2.05
18	b	830	CLA	MG-NB	-2.24	2.01	2.05

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
18	C	306	CLA	CHC-C1C	2.24	1.43	1.38
18	D	208	CLA	C1B-C2B	2.24	1.48	1.43
18	b	828	CLA	CMB-C2B	-2.24	1.46	1.50
18	a	822	CLA	CHC-C1C	2.23	1.43	1.38
18	a	802	CLA	C3B-C4B	2.23	1.49	1.42
18	D	210	CLA	C4B-NB	2.23	1.40	1.37
18	a	825	CLA	C3B-C4B	2.23	1.49	1.42
27	l	206	ET4	C04-C05	-2.23	1.47	1.51
18	b	812	CLA	CHC-C1C	2.23	1.43	1.38
18	b	801	CLA	C1B-NB	-2.23	1.34	1.37
18	l	202	CLA	C1B-NB	-2.23	1.34	1.37
18	E	316	CLA	MG-NB	-2.23	2.01	2.05
18	a	823	CLA	C3B-C4B	2.23	1.49	1.42
18	H	308	CLA	MG-ND	-2.23	2.01	2.05
26	H	301	A86	O1-C15	-2.23	1.42	1.45
18	a	842	CLA	C1B-C2B	2.23	1.48	1.43
18	b	808	CLA	C3B-C4B	2.23	1.49	1.42
18	a	805	CLA	CMD-C2D	-2.23	1.46	1.50
18	H	305	CLA	CMC-C2C	-2.23	1.46	1.50
18	H	307	CLA	MG-ND	-2.23	2.01	2.05
18	b	810	CLA	CHC-C1C	2.23	1.42	1.38
18	b	809	CLA	CHC-C1C	2.23	1.42	1.38
18	b	825	CLA	C3B-C4B	2.23	1.49	1.42
18	C	309	CLA	C1B-C2B	2.22	1.48	1.43
18	b	817	CLA	C3B-C4B	2.22	1.49	1.42
18	a	813	CLA	C1B-C2B	2.22	1.48	1.43
18	b	842	CLA	C3B-C4B	2.22	1.49	1.42
18	b	819	CLA	C3B-C4B	2.22	1.49	1.42
18	a	819	CLA	CHC-C1C	2.22	1.42	1.38
26	r	203	A86	O1-C15	-2.22	1.42	1.45
28	E	304	DD6	C28-C27	-2.22	1.49	1.50
18	a	822	CLA	C3B-C4B	2.22	1.49	1.42
18	b	817	CLA	CHC-C1C	2.22	1.42	1.38
18	E	313	CLA	CHC-C1C	2.22	1.42	1.38
18	B	314	CLA	CMD-C2D	-2.22	1.46	1.50
18	b	804	CLA	C1B-C2B	2.22	1.48	1.43
18	r	202	CLA	CHC-C1C	2.22	1.42	1.38
18	a	848	CLA	C3B-C4B	2.22	1.49	1.42
18	D	211	CLA	CHC-C1C	2.22	1.42	1.38
18	a	836	CLA	C3B-C4B	2.22	1.49	1.42
18	b	806	CLA	C3B-C4B	2.22	1.49	1.42
18	a	810	CLA	CHC-C1C	2.22	1.42	1.38

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
26	H	302	A86	C15-C16	-2.22	1.52	1.55
18	a	803	CLA	C1B-C2B	2.22	1.48	1.43
18	b	810	CLA	CMD-C2D	-2.21	1.46	1.50
18	a	806	CLA	CHC-C1C	2.21	1.42	1.38
23	E	318	LMG	O1-C7	-2.21	1.39	1.43
18	a	802	CLA	MG-NB	-2.21	2.01	2.05
18	l	205	CLA	CMD-C2D	-2.21	1.46	1.50
18	b	826	CLA	MG-NB	-2.21	2.01	2.05
18	b	807	CLA	CHC-C1C	2.21	1.42	1.38
18	b	805	CLA	C3B-C4B	2.21	1.49	1.42
18	a	823	CLA	CHC-C1C	2.21	1.42	1.38
18	b	805	CLA	CHC-C1C	2.21	1.42	1.38
18	b	818	CLA	CHC-C1C	2.21	1.42	1.38
18	a	808	CLA	C3B-C4B	2.21	1.49	1.42
18	a	854	CLA	C1D-ND	2.21	1.40	1.37
18	a	824	CLA	CHC-C1C	2.20	1.42	1.38
18	b	830	CLA	CHC-C1C	2.20	1.42	1.38
18	D	217	CLA	CMD-C2D	-2.20	1.46	1.50
18	a	802	CLA	CHC-C1C	2.20	1.42	1.38
18	a	830	CLA	CHC-C1C	2.20	1.42	1.38
18	a	839	CLA	CHC-C1C	2.20	1.42	1.38
18	b	815	CLA	C3B-C4B	2.20	1.49	1.42
18	B	306	CLA	CMB-C2B	-2.20	1.46	1.50
18	C	311	CLA	C3B-C4B	2.20	1.49	1.42
28	H	303	DD6	C40-C32	-2.20	1.49	1.53
18	C	310	CLA	CMD-C2D	-2.20	1.46	1.50
18	H	309	CLA	CHC-C1C	2.20	1.42	1.38
18	D	208	CLA	C4B-NB	2.20	1.40	1.37
18	a	815	CLA	CMB-C2B	-2.20	1.46	1.50
18	b	840	CLA	CMC-C2C	-2.20	1.46	1.50
18	b	838	CLA	C3B-C4B	2.20	1.49	1.42
18	B	309	CLA	CMD-C2D	-2.20	1.46	1.50
18	B	314	CLA	CMC-C2C	-2.20	1.46	1.50
18	b	850	CLA	CHC-C1C	2.20	1.42	1.38
18	f	202	CLA	CHC-C1C	2.20	1.42	1.38
18	B	308	CLA	CHC-C1C	2.20	1.42	1.38
18	a	805	CLA	C3B-C4B	2.20	1.49	1.42
18	b	807	CLA	C3B-C4B	2.20	1.49	1.42
18	b	850	CLA	C3B-C4B	2.20	1.49	1.42
18	b	813	CLA	CMB-C2B	-2.20	1.46	1.50
18	a	843	CLA	CHC-C1C	2.20	1.42	1.38
18	a	847	CLA	CHC-C1C	2.19	1.42	1.38

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
18	a	841	CLA	C4B-NB	2.19	1.40	1.37
18	b	837	CLA	C1B-NB	-2.19	1.35	1.37
26	C	305	A86	O1-C15	-2.19	1.42	1.45
18	E	310	CLA	CHC-C1C	2.19	1.42	1.38
18	D	216	CLA	C4B-NB	2.19	1.40	1.37
23	j	103	LMG	O7-C10	2.19	1.40	1.34
18	a	828	CLA	CHC-C1C	2.19	1.42	1.38
18	a	816	CLA	CHC-C1C	2.19	1.42	1.38
18	a	826	CLA	CHC-C1C	2.19	1.42	1.38
26	H	301	A86	C-C1	-2.19	1.46	1.50
18	C	312	CLA	C4B-NB	2.19	1.40	1.37
18	H	314	CLA	C1B-NB	-2.19	1.35	1.37
18	C	318	CLA	CMC-C2C	-2.19	1.46	1.50
18	b	815	CLA	CHC-C1C	2.19	1.42	1.38
18	a	844	CLA	C3B-C4B	2.19	1.49	1.42
18	a	828	CLA	C3B-C4B	2.18	1.49	1.42
18	a	822	CLA	CMD-C2D	-2.18	1.46	1.50
18	H	315	CLA	CHC-C1C	2.18	1.42	1.38
26	m	102	A86	O1-C15	-2.18	1.42	1.45
18	D	214	CLA	CHC-C1C	2.18	1.42	1.38
18	E	311	CLA	CMD-C2D	-2.18	1.46	1.50
18	a	825	CLA	MG-ND	-2.18	2.01	2.05
18	a	814	CLA	CHC-C1C	2.18	1.42	1.38
18	B	308	CLA	CMD-C2D	-2.18	1.46	1.50
18	a	812	CLA	CHC-C1C	2.18	1.42	1.38
18	b	848	CLA	CHC-C1C	2.18	1.42	1.38
18	b	849	CLA	CHC-C1C	2.18	1.42	1.38
18	E	310	CLA	CMD-C2D	-2.18	1.46	1.50
18	C	310	CLA	CMC-C2C	-2.18	1.46	1.50
18	a	837	CLA	CAA-C2A	-2.18	1.50	1.54
18	b	801	CLA	C4B-NB	2.18	1.40	1.37
18	b	838	CLA	CHC-C1C	2.18	1.42	1.38
18	b	816	CLA	C3B-C4B	2.18	1.49	1.42
18	a	836	CLA	CHC-C1C	2.18	1.42	1.38
18	H	313	CLA	MG-ND	-2.18	2.01	2.05
26	B	302	A86	C15-C16	-2.18	1.52	1.55
26	H	302	A86	O1-C15	-2.18	1.42	1.45
18	b	842	CLA	CHC-C1C	2.18	1.42	1.38
18	f	204	CLA	CHC-C1C	2.18	1.42	1.38
18	E	316	CLA	C3B-C4B	2.18	1.49	1.42
18	b	823	CLA	CHC-C1C	2.18	1.42	1.38
26	B	305	A86	O1-C15	-2.18	1.42	1.45

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
18	D	212	CLA	CMD-C2D	-2.18	1.46	1.50
26	C	304	A86	C15-C16	-2.18	1.52	1.55
18	B	311	CLA	CHC-C1C	2.18	1.42	1.38
20	E	301	LHG	P-O6	2.17	1.67	1.59
18	b	845	CLA	CMD-C2D	-2.17	1.46	1.50
18	a	827	CLA	CHC-C1C	2.17	1.42	1.38
18	b	844	CLA	CHC-C1C	2.17	1.42	1.38
18	a	809	CLA	CMD-C2D	-2.17	1.46	1.50
18	b	816	CLA	CMB-C2B	-2.17	1.46	1.50
18	a	820	CLA	C3B-C4B	2.17	1.49	1.42
18	b	809	CLA	MG-NB	-2.17	2.01	2.05
18	b	825	CLA	CHC-C1C	2.17	1.42	1.38
18	C	317	CLA	C1B-C2B	2.17	1.48	1.43
18	b	840	CLA	CMD-C2D	-2.17	1.46	1.50
18	D	213	CLA	CMC-C2C	-2.17	1.46	1.50
18	D	212	CLA	CMB-C2B	-2.17	1.46	1.50
18	b	846	CLA	CHC-C1C	2.17	1.42	1.38
18	b	803	CLA	CAA-C2A	-2.17	1.50	1.54
18	a	812	CLA	CMB-C2B	-2.17	1.46	1.50
18	E	311	CLA	CHC-C1C	2.17	1.42	1.38
18	E	314	CLA	MG-ND	-2.17	2.01	2.05
18	a	819	CLA	CMC-C2C	-2.17	1.46	1.50
28	C	303	DD6	C17-C16	-2.16	1.51	1.54
18	b	819	CLA	CHC-C1C	2.16	1.42	1.38
18	a	824	CLA	C3B-C4B	2.16	1.49	1.42
18	a	801	CLA	C1B-C2B	2.16	1.48	1.43
18	B	310	CLA	MG-NB	-2.16	2.01	2.05
18	a	803	CLA	CAA-C2A	-2.16	1.50	1.54
18	a	803	CLA	C1B-NB	-2.16	1.35	1.37
18	E	315	CLA	CMB-C2B	-2.16	1.46	1.50
18	b	819	CLA	CMD-C2D	-2.16	1.46	1.50
18	B	312	CLA	CHC-C1C	2.16	1.42	1.38
22	a	849	SQD	O48-C46	-2.16	1.40	1.45
18	B	306	CLA	CMC-C2C	-2.16	1.46	1.50
18	D	215	CLA	CHC-C1C	2.16	1.42	1.38
18	a	855	CLA	CMD-C2D	-2.16	1.46	1.50
18	B	311	CLA	CMD-C2D	-2.16	1.46	1.50
18	a	855	CLA	MG-NB	-2.16	2.01	2.05
18	b	827	CLA	CHC-C1C	2.16	1.42	1.38
27	l	206	ET4	C35-C36	-2.16	1.49	1.52
28	E	303	DD6	O1-C20	-2.16	1.43	1.46
18	a	807	CLA	MG-ND	-2.16	2.01	2.05

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
18	a	840	CLA	C4B-NB	2.16	1.40	1.37
18	b	823	CLA	C3B-C4B	2.16	1.49	1.42
26	H	304	A86	O1-C15	-2.15	1.42	1.45
18	i	101	CLA	MG-NB	-2.15	2.01	2.05
26	m	102	A86	C15-C16	-2.15	1.52	1.55
18	D	209	CLA	CHC-C1C	2.15	1.42	1.38
18	l	202	CLA	C3B-C4B	2.15	1.49	1.42
18	a	844	CLA	CMD-C2D	-2.15	1.46	1.50
18	a	848	CLA	CHC-C1C	2.15	1.42	1.38
20	E	301	LHG	O7-C7	2.15	1.40	1.34
18	a	802	CLA	CMD-C2D	-2.15	1.46	1.50
18	C	316	CLA	C1B-C2B	2.15	1.48	1.43
18	a	820	CLA	MG-NB	-2.15	2.01	2.05
18	B	306	CLA	MG-ND	-2.15	2.01	2.05
18	a	809	CLA	CHC-C1C	2.15	1.42	1.38
18	b	808	CLA	MG-NB	-2.15	2.01	2.05
28	H	303	DD6	C41-C32	-2.15	1.49	1.53
18	b	805	CLA	CMD-C2D	-2.15	1.46	1.50
18	D	217	CLA	CMB-C2B	-2.15	1.46	1.50
18	b	845	CLA	CHC-C1C	2.15	1.42	1.38
18	a	843	CLA	CMD-C2D	-2.15	1.46	1.50
18	b	830	CLA	CMD-C2D	-2.15	1.46	1.50
18	D	214	CLA	CMC-C2C	-2.15	1.46	1.50
18	D	209	CLA	CMD-C2D	-2.15	1.46	1.50
18	a	818	CLA	C1B-C2B	2.15	1.48	1.43
18	a	839	CLA	CMB-C2B	-2.15	1.46	1.50
18	b	816	CLA	CMD-C2D	-2.15	1.46	1.50
18	b	838	CLA	CMD-C2D	-2.15	1.46	1.50
18	C	314	CLA	C1B-C2B	2.14	1.48	1.43
18	a	830	CLA	C3B-C4B	2.14	1.49	1.42
18	a	815	CLA	CMD-C2D	-2.14	1.46	1.50
18	b	828	CLA	CMD-C2D	-2.14	1.46	1.50
18	H	313	CLA	CAA-C2A	-2.14	1.50	1.54
18	C	310	CLA	CMB-C2B	-2.14	1.46	1.50
18	b	810	CLA	CMC-C2C	-2.14	1.46	1.50
18	a	802	CLA	CMB-C2B	-2.14	1.46	1.50
18	b	804	CLA	CAA-C2A	-2.14	1.50	1.54
18	H	306	CLA	C1B-C2B	2.14	1.48	1.43
18	f	202	CLA	CMD-C2D	-2.14	1.46	1.50
20	j	102	LHG	P-O6	2.14	1.67	1.59
18	D	216	CLA	MG-ND	-2.14	2.01	2.05
18	D	208	CLA	MG-ND	-2.14	2.01	2.05

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
18	a	817	CLA	CHC-C1C	2.14	1.42	1.38
18	a	819	CLA	CMB-C2B	-2.14	1.46	1.50
18	l	203	CLA	MG-ND	-2.14	2.01	2.05
18	C	318	CLA	C1B-C2B	2.14	1.48	1.43
18	b	814	CLA	CMB-C2B	-2.14	1.46	1.50
18	D	212	CLA	MG-NB	-2.14	2.01	2.05
26	D	204	A86	O1-C15	-2.14	1.42	1.45
23	D	202	LMG	O1-C7	-2.14	1.40	1.43
18	r	202	CLA	MG-NB	-2.14	2.01	2.05
18	a	817	CLA	CMD-C2D	-2.13	1.46	1.50
18	C	309	CLA	MG-ND	-2.13	2.01	2.05
18	B	306	CLA	CMD-C2D	-2.13	1.46	1.50
18	b	841	CLA	CMD-C2D	-2.13	1.46	1.50
18	E	310	CLA	CMB-C2B	-2.13	1.46	1.50
26	B	305	A86	O4-C34	-2.13	1.41	1.46
18	b	820	CLA	MG-ND	-2.13	2.01	2.05
26	B	305	A86	C17-C16	-2.13	1.51	1.54
18	a	826	CLA	MG-NB	-2.13	2.01	2.05
18	b	813	CLA	C3B-C4B	2.13	1.49	1.42
18	a	818	CLA	C1B-NB	-2.13	1.35	1.37
18	a	853	CLA	MG-NB	-2.13	2.01	2.05
18	E	315	CLA	MG-NB	-2.13	2.01	2.05
18	a	804	CLA	CMD-C2D	-2.13	1.46	1.50
18	j	104	CLA	CMD-C2D	-2.13	1.46	1.50
18	i	101	CLA	CMD-C2D	-2.13	1.46	1.50
18	B	307	CLA	CMC-C2C	-2.13	1.46	1.50
18	b	816	CLA	MG-NB	-2.13	2.01	2.05
18	b	829	CLA	MG-ND	-2.13	2.01	2.05
18	B	309	CLA	MG-NB	-2.13	2.01	2.05
20	H	316	LHG	P-O6	2.13	1.67	1.59
26	E	302	A86	C13-C11	-2.13	1.45	1.49
18	a	816	CLA	CMD-C2D	-2.13	1.46	1.50
18	a	847	CLA	CMD-C2D	-2.13	1.46	1.50
18	a	841	CLA	C1B-C2B	2.13	1.48	1.43
18	B	309	CLA	C3B-C4B	2.13	1.48	1.42
18	b	812	CLA	CMB-C2B	-2.13	1.46	1.50
18	b	806	CLA	CHC-C1C	2.13	1.42	1.38
18	a	814	CLA	MG-NB	-2.13	2.01	2.05
18	E	312	CLA	C1B-NB	-2.13	1.35	1.37
18	b	821	CLA	C3B-C4B	2.13	1.48	1.42
18	b	845	CLA	MG-NB	-2.13	2.01	2.05
18	b	842	CLA	CMB-C2B	-2.13	1.46	1.50

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
18	a	844	CLA	CHC-C1C	2.13	1.42	1.38
18	a	824	CLA	MG-NB	-2.12	2.01	2.05
18	b	807	CLA	MG-NB	-2.12	2.01	2.05
18	C	314	CLA	MG-ND	-2.12	2.01	2.05
18	b	821	CLA	CHC-C1C	2.12	1.42	1.38
18	b	848	CLA	MG-NB	-2.12	2.01	2.05
23	C	319	LMG	C3-C2	-2.12	1.46	1.52
18	D	209	CLA	MG-NB	-2.12	2.01	2.05
18	a	821	CLA	CHC-C1C	2.12	1.42	1.38
18	a	813	CLA	C4B-NB	2.12	1.40	1.37
18	B	311	CLA	CMB-C2B	-2.12	1.46	1.50
18	D	211	CLA	CMC-C2C	-2.12	1.46	1.50
18	b	807	CLA	CMD-C2D	-2.12	1.46	1.50
18	a	853	CLA	CMD-C2D	-2.12	1.46	1.50
18	b	827	CLA	CMB-C2B	-2.12	1.46	1.50
18	b	844	CLA	CMD-C2D	-2.12	1.46	1.50
18	D	210	CLA	MG-ND	-2.12	2.01	2.05
18	a	839	CLA	MG-NB	-2.12	2.01	2.05
18	b	814	CLA	MG-NB	-2.12	2.01	2.05
18	H	314	CLA	C1B-C2B	2.12	1.48	1.43
18	f	204	CLA	CMD-C2D	-2.12	1.46	1.50
18	a	801	CLA	CAA-C2A	-2.12	1.50	1.54
18	b	818	CLA	CMD-C2D	-2.12	1.46	1.50
18	a	808	CLA	CHC-C1C	2.12	1.42	1.38
18	H	307	CLA	C1B-C2B	2.12	1.48	1.43
18	b	825	CLA	CMB-C2B	-2.12	1.46	1.50
18	a	808	CLA	CMC-C2C	-2.12	1.46	1.50
18	b	846	CLA	CMD-C2D	-2.12	1.46	1.50
18	D	211	CLA	MG-NB	-2.12	2.01	2.05
18	a	822	CLA	CMB-C2B	-2.11	1.46	1.50
18	b	820	CLA	C1B-C2B	2.11	1.48	1.43
18	D	217	CLA	C3B-C4B	2.11	1.48	1.42
18	b	815	CLA	CMD-C2D	-2.11	1.46	1.50
18	l	203	CLA	C1B-C2B	2.11	1.48	1.43
18	a	814	CLA	CMD-C2D	-2.11	1.46	1.50
18	a	815	CLA	MG-NB	-2.11	2.01	2.05
18	a	828	CLA	MG-NB	-2.11	2.01	2.05
18	b	844	CLA	MG-NB	-2.11	2.01	2.05
18	D	214	CLA	MG-NB	-2.11	2.01	2.05
18	E	310	CLA	MG-NB	-2.11	2.01	2.05
26	b	847	A86	O1-C15	-2.11	1.42	1.45
26	D	204	A86	C-C1	-2.11	1.46	1.50

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
18	D	209	CLA	CMC-C2C	-2.11	1.46	1.50
18	a	846	CLA	MG-NB	-2.11	2.01	2.05
26	D	206	A86	C15-C16	-2.11	1.52	1.55
18	a	806	CLA	CMD-C2D	-2.11	1.46	1.50
18	H	308	CLA	C1B-NB	-2.11	1.35	1.37
18	a	814	CLA	CMB-C2B	-2.11	1.46	1.50
18	H	307	CLA	C1B-NB	-2.11	1.35	1.37
27	l	206	ET4	C12-C13	2.11	1.50	1.46
26	C	304	A86	O1-C15	-2.11	1.42	1.45
18	b	813	CLA	CMD-C2D	-2.11	1.46	1.50
18	E	314	CLA	C3B-C4B	2.11	1.48	1.42
18	a	803	CLA	MG-ND	-2.11	2.01	2.05
18	B	311	CLA	MG-NB	-2.11	2.01	2.05
18	b	817	CLA	CMC-C2C	-2.11	1.46	1.50
26	B	305	A86	C-C1	-2.10	1.46	1.50
18	E	313	CLA	CMD-C2D	-2.10	1.46	1.50
18	b	810	CLA	MG-NB	-2.10	2.01	2.05
18	b	809	CLA	CMB-C2B	-2.10	1.46	1.50
18	b	814	CLA	CMD-C2D	-2.10	1.46	1.50
18	C	314	CLA	C3B-C4B	2.10	1.48	1.42
26	D	203	A86	O1-C15	-2.10	1.42	1.45
18	a	808	CLA	MG-NB	-2.10	2.01	2.05
18	a	824	CLA	CMD-C2D	-2.10	1.46	1.50
18	a	848	CLA	CMB-C2B	-2.10	1.46	1.50
18	a	805	CLA	CMB-C2B	-2.10	1.46	1.50
26	E	302	A86	O1-C15	-2.10	1.42	1.45
18	a	807	CLA	C1B-C2B	2.10	1.48	1.43
18	a	843	CLA	MG-NB	-2.10	2.01	2.05
18	b	805	CLA	CMB-C2B	-2.10	1.46	1.50
18	H	309	CLA	CMB-C2B	-2.10	1.46	1.50
18	b	822	CLA	CMD-C2D	-2.10	1.46	1.50
18	b	823	CLA	CMB-C2B	-2.10	1.46	1.50
18	B	307	CLA	MG-NB	-2.10	2.01	2.05
18	a	838	CLA	CMD-C2D	-2.10	1.46	1.50
18	E	314	CLA	CAA-C2A	-2.10	1.50	1.54
18	b	850	CLA	CMD-C2D	-2.10	1.46	1.50
20	a	832	LHG	P-O6	2.10	1.67	1.59
26	C	305	A86	C15-C16	-2.10	1.52	1.55
18	b	846	CLA	CMB-C2B	-2.10	1.46	1.50
18	a	823	CLA	MG-NB	-2.10	2.01	2.05
18	a	821	CLA	CMB-C2B	-2.10	1.46	1.50
26	B	305	A86	C15-C16	-2.10	1.52	1.55

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
18	a	826	CLA	CMB-C2B	-2.10	1.46	1.50
18	a	810	CLA	MG-NB	-2.10	2.01	2.05
18	E	311	CLA	MG-NB	-2.10	2.01	2.05
18	B	307	CLA	CMB-C2B	-2.10	1.46	1.50
18	E	312	CLA	C1B-C2B	2.10	1.48	1.43
18	a	801	CLA	C1B-NB	-2.10	1.35	1.37
18	a	812	CLA	CMD-C2D	-2.10	1.46	1.50
20	E	301	LHG	O8-C23	2.10	1.39	1.33
26	E	302	A86	C17-C16	-2.10	1.51	1.54
18	a	810	CLA	CMD-C2D	-2.10	1.46	1.50
18	E	309	CLA	CMD-C2D	-2.10	1.46	1.50
18	b	825	CLA	MG-NB	-2.10	2.01	2.05
18	j	104	CLA	MG-NB	-2.10	2.01	2.05
18	a	830	CLA	CMB-C2B	-2.09	1.46	1.50
18	a	811	CLA	CHC-C1C	2.09	1.42	1.38
18	b	844	CLA	CMB-C2B	-2.09	1.46	1.50
18	b	819	CLA	MG-NB	-2.09	2.01	2.05
18	a	848	CLA	CMD-C2D	-2.09	1.46	1.50
18	B	309	CLA	CMB-C2B	-2.09	1.46	1.50
18	D	213	CLA	CMD-C2D	-2.09	1.46	1.50
18	b	828	CLA	MG-NB	-2.09	2.01	2.05
18	E	316	CLA	CHC-C1C	2.09	1.42	1.38
18	D	208	CLA	C3B-C4B	2.09	1.48	1.42
18	a	816	CLA	MG-NB	-2.09	2.01	2.05
18	b	840	CLA	MG-NB	-2.09	2.01	2.05
18	b	825	CLA	CMD-C2D	-2.09	1.46	1.50
18	C	318	CLA	C4B-NB	2.09	1.40	1.37
27	l	206	ET4	C41-C34	-2.09	1.49	1.53
18	a	817	CLA	MG-NB	-2.09	2.01	2.05
18	b	819	CLA	CMB-C2B	-2.09	1.46	1.50
18	D	211	CLA	CMD-C2D	-2.09	1.46	1.50
18	H	310	CLA	C1B-NB	-2.09	1.35	1.37
18	a	806	CLA	CMB-C2B	-2.08	1.46	1.50
18	b	821	CLA	CMB-C2B	-2.08	1.46	1.50
23	E	318	LMG	O7-C10	2.08	1.40	1.34
18	b	848	CLA	CMB-C2B	-2.08	1.46	1.50
18	f	203	CLA	CMD-C2D	-2.08	1.46	1.50
18	H	305	CLA	MG-ND	-2.08	2.01	2.05
18	E	313	CLA	CMB-C2B	-2.08	1.46	1.50
18	C	312	CLA	CHC-C1C	2.08	1.42	1.38
18	C	314	CLA	C1B-NB	-2.08	1.35	1.37
18	H	309	CLA	CMD-C2D	-2.08	1.46	1.50

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
18	b	845	CLA	CMB-C2B	-2.08	1.46	1.50
18	a	816	CLA	CMB-C2B	-2.08	1.46	1.50
18	b	817	CLA	CMD-C2D	-2.08	1.46	1.50
18	b	811	CLA	C1B-C2B	2.08	1.48	1.43
18	a	847	CLA	CMC-C2C	-2.08	1.46	1.50
18	H	314	CLA	C4B-NB	2.08	1.40	1.37
18	a	846	CLA	CMB-C2B	-2.08	1.46	1.50
18	a	819	CLA	CMD-C2D	-2.08	1.46	1.50
26	H	304	A86	C15-C16	-2.08	1.52	1.55
18	b	840	CLA	CHC-C1C	2.08	1.42	1.38
18	b	849	CLA	MG-NB	-2.08	2.01	2.05
18	b	816	CLA	CHC-C1C	2.08	1.42	1.38
18	b	827	CLA	CMD-C2D	-2.08	1.46	1.50
18	E	313	CLA	MG-NB	-2.08	2.01	2.05
18	a	807	CLA	C1B-NB	-2.08	1.35	1.37
18	b	827	CLA	CMC-C2C	-2.08	1.46	1.50
18	b	838	CLA	CMB-C2B	-2.08	1.46	1.50
18	E	309	CLA	CMB-C2B	-2.08	1.46	1.50
18	a	827	CLA	MG-NB	-2.08	2.01	2.05
18	f	204	CLA	CMB-C2B	-2.08	1.46	1.50
18	D	215	CLA	MG-NB	-2.08	2.01	2.05
18	a	804	CLA	CMC-C2C	-2.07	1.46	1.50
18	D	214	CLA	CMB-C2B	-2.07	1.46	1.50
18	b	840	CLA	CMB-C2B	-2.07	1.46	1.50
18	H	309	CLA	MG-NB	-2.07	2.01	2.05
18	b	809	CLA	CMD-C2D	-2.07	1.46	1.50
18	b	808	CLA	CMD-C2D	-2.07	1.46	1.50
18	D	214	CLA	CMD-C2D	-2.07	1.46	1.50
18	a	839	CLA	CMD-C2D	-2.07	1.46	1.50
18	D	208	CLA	C1B-NB	-2.07	1.35	1.37
18	b	838	CLA	CMC-C2C	-2.07	1.46	1.50
18	D	215	CLA	CMB-C2B	-2.07	1.46	1.50
18	l	205	CLA	MG-NB	-2.07	2.01	2.05
18	i	101	CLA	CMB-C2B	-2.07	1.46	1.50
18	b	810	CLA	CMB-C2B	-2.07	1.46	1.50
18	D	209	CLA	CMB-C2B	-2.07	1.46	1.50
18	a	805	CLA	CMC-C2C	-2.07	1.46	1.50
18	a	853	CLA	CMB-C2B	-2.07	1.46	1.50
18	B	308	CLA	CMC-C2C	-2.07	1.46	1.50
18	a	812	CLA	MG-NB	-2.07	2.01	2.05
18	a	821	CLA	CMC-C2C	-2.07	1.46	1.50
18	a	828	CLA	CMD-C2D	-2.07	1.46	1.50

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
26	H	301	A86	C13-C11	-2.07	1.45	1.49
18	b	827	CLA	MG-NB	-2.07	2.01	2.05
18	B	309	CLA	CHC-C1C	2.07	1.42	1.38
18	a	811	CLA	CMD-C2D	-2.07	1.46	1.50
18	b	806	CLA	CMC-C2C	-2.07	1.46	1.50
28	E	307	DD6	O1-C20	-2.07	1.43	1.46
18	C	306	CLA	CMB-C2B	-2.07	1.46	1.50
18	l	202	CLA	C4B-NB	2.07	1.40	1.37
20	D	201	LHG	O7-C7	2.07	1.40	1.34
18	E	314	CLA	CHC-C1C	2.06	1.42	1.38
18	D	208	CLA	CHC-C1C	2.06	1.42	1.38
20	H	316	LHG	O7-C5	-2.06	1.41	1.46
18	E	311	CLA	CMB-C2B	-2.06	1.46	1.50
18	B	312	CLA	MG-NB	-2.06	2.01	2.05
18	a	823	CLA	CMD-C2D	-2.06	1.46	1.50
18	a	847	CLA	MG-NB	-2.06	2.01	2.05
18	B	308	CLA	MG-NB	-2.06	2.01	2.05
18	C	306	CLA	CMD-C2D	-2.06	1.46	1.50
18	f	202	CLA	MG-NB	-2.06	2.01	2.05
18	B	307	CLA	CMD-C2D	-2.06	1.46	1.50
18	a	808	CLA	CMB-C2B	-2.06	1.46	1.50
18	b	803	CLA	CHC-C4B	-2.06	1.34	1.39
18	b	817	CLA	MG-NB	-2.06	2.01	2.05
18	b	821	CLA	MG-NB	-2.06	2.01	2.05
18	C	314	CLA	CHC-C1C	2.06	1.42	1.38
18	b	818	CLA	CMB-C2B	-2.06	1.46	1.50
18	a	830	CLA	CMD-C2D	-2.06	1.46	1.50
18	b	826	CLA	CMD-C2D	-2.06	1.46	1.50
26	H	304	A86	C13-C11	-2.06	1.45	1.49
18	C	306	CLA	MG-NB	-2.06	2.01	2.05
18	a	846	CLA	CMD-C2D	-2.06	1.46	1.50
18	b	818	CLA	CMC-C2C	-2.06	1.46	1.50
18	b	823	CLA	CMD-C2D	-2.06	1.46	1.50
18	b	843	CLA	C4B-NB	2.06	1.40	1.37
18	f	203	CLA	CHC-C1C	2.06	1.42	1.38
18	E	315	CLA	CMD-C2D	-2.06	1.46	1.50
18	b	846	CLA	CMC-C2C	-2.06	1.46	1.50
28	D	205	DD6	C37-C36	-2.06	1.47	1.50
18	f	204	CLA	MG-NB	-2.06	2.01	2.05
18	a	810	CLA	CMC-C2C	-2.06	1.46	1.50
18	b	808	CLA	CHC-C1C	2.06	1.42	1.38
18	b	842	CLA	CMD-C2D	-2.05	1.46	1.50

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
18	f	203	CLA	CMC-C2C	-2.05	1.46	1.50
18	a	805	CLA	CHC-C1C	2.05	1.42	1.38
18	B	314	CLA	CHC-C1C	2.05	1.42	1.38
18	b	808	CLA	CMC-C2C	-2.05	1.46	1.50
18	b	822	CLA	CMC-C2C	-2.05	1.46	1.50
18	a	806	CLA	MG-NB	-2.05	2.01	2.05
18	a	823	CLA	CMB-C2B	-2.05	1.46	1.50
18	D	213	CLA	CMB-C2B	-2.05	1.46	1.50
18	a	809	CLA	CMB-C2B	-2.05	1.46	1.50
18	r	202	CLA	CMB-C2B	-2.05	1.46	1.50
18	D	211	CLA	CMB-C2B	-2.05	1.46	1.50
18	a	827	CLA	CMD-C2D	-2.05	1.46	1.50
28	B	303	DD6	C33-C34	-2.05	1.49	1.52
26	C	302	A86	C17-C16	-2.05	1.51	1.54
18	C	316	CLA	CHC-C1C	2.05	1.42	1.38
18	H	315	CLA	CMD-C2D	-2.05	1.46	1.50
18	b	842	CLA	MG-NB	-2.05	2.01	2.05
18	b	805	CLA	CMC-C2C	-2.05	1.46	1.50
18	a	825	CLA	CAC-C3C	-2.05	1.45	1.51
18	E	316	CLA	CMC-C2C	-2.05	1.46	1.50
18	a	813	CLA	MG-ND	-2.05	2.01	2.05
18	a	844	CLA	CMB-C2B	-2.05	1.46	1.50
18	b	850	CLA	CMB-C2B	-2.05	1.46	1.50
18	b	812	CLA	CMD-C2D	-2.05	1.46	1.50
18	a	817	CLA	CMB-C2B	-2.04	1.46	1.50
18	a	826	CLA	CMD-C2D	-2.04	1.46	1.50
18	b	824	CLA	C1B-C2B	2.04	1.47	1.43
18	a	814	CLA	CMC-C2C	-2.04	1.46	1.50
18	B	312	CLA	CMB-C2B	-2.04	1.46	1.50
18	a	810	CLA	CMB-C2B	-2.04	1.46	1.50
18	a	839	CLA	CMC-C2C	-2.04	1.46	1.50
18	j	104	CLA	CMB-C2B	-2.04	1.46	1.50
26	B	305	A86	C13-C11	-2.04	1.45	1.49
27	l	206	ET4	C20-C21	2.04	1.49	1.43
18	C	309	CLA	C1B-NB	-2.04	1.35	1.37
18	H	315	CLA	MG-NB	-2.04	2.01	2.05
18	r	202	CLA	CMD-C2D	-2.04	1.46	1.50
18	a	804	CLA	MG-NB	-2.04	2.01	2.05
18	b	837	CLA	C4B-NB	2.04	1.40	1.37
18	a	811	CLA	CMB-C2B	-2.04	1.46	1.50
26	B	304	A86	O1-C15	-2.04	1.42	1.45
18	a	836	CLA	MG-NB	-2.04	2.01	2.05

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
18	b	815	CLA	MG-NB	-2.04	2.01	2.05
18	a	847	CLA	CMB-C2B	-2.04	1.46	1.50
18	b	806	CLA	CMD-C2D	-2.04	1.46	1.50
18	b	817	CLA	CMB-C2B	-2.04	1.46	1.50
18	a	848	CLA	MG-NB	-2.04	2.01	2.05
18	a	813	CLA	C1B-NB	-2.04	1.35	1.37
18	b	841	CLA	CMC-C2C	-2.04	1.46	1.50
18	a	838	CLA	CMC-C2C	-2.04	1.46	1.50
18	b	809	CLA	CMC-C2C	-2.04	1.46	1.50
18	a	830	CLA	MG-NB	-2.04	2.01	2.05
26	C	302	A86	O4-C34	-2.04	1.41	1.46
18	b	838	CLA	MG-NB	-2.03	2.01	2.05
26	m	102	A86	C32-C31	-2.03	1.51	1.54
18	a	827	CLA	CMB-C2B	-2.03	1.46	1.50
21	f	201	BCR	C30-C25	-2.03	1.51	1.53
29	C	308	KC1	C4A-C3A	-2.03	1.40	1.44
18	a	824	CLA	CMB-C2B	-2.03	1.46	1.50
18	b	806	CLA	CMB-C2B	-2.03	1.46	1.50
18	E	313	CLA	CMC-C2C	-2.03	1.46	1.50
18	D	207	CLA	C3B-C4B	2.03	1.48	1.42
18	B	308	CLA	CMB-C2B	-2.03	1.46	1.50
23	l	207	LMG	O7-C10	2.03	1.40	1.34
18	C	315	CLA	CMD-C2D	-2.03	1.46	1.50
18	a	820	CLA	CHC-C1C	2.03	1.42	1.38
18	D	213	CLA	MG-NB	-2.03	2.01	2.05
18	E	315	CLA	CMC-C2C	-2.03	1.46	1.50
18	H	315	CLA	CMC-C2C	-2.03	1.46	1.50
18	b	815	CLA	CMB-C2B	-2.03	1.46	1.50
18	B	310	CLA	CMB-C2B	-2.03	1.46	1.50
18	B	312	CLA	CMD-C2D	-2.03	1.46	1.50
18	b	823	CLA	MG-NB	-2.03	2.01	2.05
18	b	829	CLA	C4B-NB	2.03	1.40	1.37
18	b	812	CLA	CMC-C2C	-2.03	1.46	1.50
18	a	821	CLA	CMD-C2D	-2.03	1.46	1.50
18	C	306	CLA	CMC-C2C	-2.03	1.46	1.50
18	C	316	CLA	C3B-C4B	2.03	1.48	1.42
18	b	830	CLA	CMB-C2B	-2.03	1.46	1.50
18	a	828	CLA	CMB-C2B	-2.03	1.46	1.50
18	b	821	CLA	CMC-C2C	-2.03	1.46	1.50
18	b	828	CLA	CMC-C2C	-2.03	1.46	1.50
18	b	822	CLA	MG-NB	-2.03	2.01	2.05
18	f	203	CLA	CMB-C2B	-2.03	1.46	1.50

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
18	b	801	CLA	CAC-C3C	-2.03	1.45	1.51
18	b	841	CLA	MG-NB	-2.02	2.01	2.05
18	C	317	CLA	C1B-NB	-2.02	1.35	1.37
18	E	311	CLA	CMC-C2C	-2.02	1.46	1.50
18	H	306	CLA	C1B-NB	-2.02	1.35	1.37
18	D	215	CLA	CMC-C2C	-2.02	1.46	1.50
18	C	312	CLA	C1B-NB	-2.02	1.35	1.37
18	b	826	CLA	CMB-C2B	-2.02	1.46	1.50
18	a	811	CLA	MG-NB	-2.02	2.01	2.05
18	D	207	CLA	C1B-C2B	2.02	1.47	1.43
26	r	203	A86	C15-C16	-2.02	1.52	1.55
18	a	855	CLA	CMB-C2B	-2.02	1.46	1.50
18	B	309	CLA	CMC-C2C	-2.02	1.46	1.50
18	C	311	CLA	CHC-C1C	2.02	1.42	1.38
18	b	814	CLA	CMC-C2C	-2.02	1.46	1.50
18	b	820	CLA	CHC-C4B	-2.02	1.34	1.39
18	b	820	CLA	C1B-NB	-2.02	1.35	1.37
18	a	804	CLA	CMB-C2B	-2.02	1.46	1.50
18	H	315	CLA	CMB-C2B	-2.02	1.46	1.50
18	C	310	CLA	C3B-C4B	2.02	1.48	1.42
18	l	205	CLA	CMB-C2B	-2.02	1.46	1.50
18	b	849	CLA	CMC-C2C	-2.02	1.46	1.50
18	C	315	CLA	MG-NB	-2.02	2.01	2.05
18	D	215	CLA	CMD-C2D	-2.02	1.46	1.50
18	b	801	CLA	C1B-C2B	2.01	1.47	1.43
23	D	202	LMG	O7-C10	2.01	1.40	1.34
18	C	312	CLA	MG-ND	-2.01	2.01	2.05
18	a	843	CLA	CMC-C2C	-2.01	1.46	1.50
18	C	316	CLA	MG-ND	-2.01	2.01	2.05
21	i	102	BCR	C1-C6	-2.01	1.51	1.53
18	b	830	CLA	CMC-C2C	-2.01	1.46	1.50
18	D	217	CLA	CMC-C2C	-2.01	1.46	1.50
18	b	819	CLA	CMC-C2C	-2.01	1.46	1.50
18	a	825	CLA	C1B-C2B	2.01	1.47	1.43
18	b	812	CLA	MG-NB	-2.01	2.01	2.05
18	b	829	CLA	C1B-NB	-2.01	1.35	1.37
18	a	806	CLA	CMC-C2C	-2.01	1.46	1.50
18	b	807	CLA	CMC-C2C	-2.01	1.46	1.50
18	B	310	CLA	CMC-C2C	-2.00	1.46	1.50
18	a	812	CLA	CMC-C2C	-2.00	1.46	1.50
18	b	815	CLA	CMC-C2C	-2.00	1.46	1.50
26	B	301	A86	C15-C16	-2.00	1.52	1.55

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
18	H	306	CLA	CAC-C3C	-2.00	1.46	1.51
26	m	102	A86	C-C1	-2.00	1.46	1.50
18	a	802	CLA	CMC-C2C	-2.00	1.46	1.50
18	D	207	CLA	CHC-C1C	2.00	1.42	1.38

All (1840) bond angle outliers are listed below:

Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
23	j	103	LMG	O2-C2-C1	-21.74	58.28	110.08
23	a	851	LMG	O2-C2-C1	-19.19	64.35	110.08
28	E	304	DD6	C21-C20-C15	-16.50	95.14	122.30
23	C	301	LMG	O2-C2-C1	15.53	147.09	110.08
21	b	833	BCR	C32-C1-C6	-14.58	87.39	110.24
26	r	203	A86	O1-C20-C19	-12.78	101.52	113.49
26	D	204	A86	C3-C4-C5	-12.14	98.69	123.52
28	B	303	DD6	C21-C20-C15	-11.60	103.21	122.30
26	D	203	A86	C25-C26-C27	-11.36	111.34	127.28
26	D	203	A86	O1-C20-C19	-11.03	103.15	113.49
28	D	205	DD6	C21-C20-C15	-10.90	104.36	122.30
26	r	203	A86	C20-O1-C15	10.90	65.71	61.03
22	a	849	SQD	O47-C7-O49	-10.71	111.97	125.70
26	H	304	A86	C20-O1-C15	10.40	65.50	61.03
26	D	203	A86	C20-O1-C15	10.37	65.48	61.03
26	C	305	A86	C20-O1-C15	10.28	65.45	61.03
23	C	319	LMG	O2-C2-C3	-10.20	86.32	110.38
26	C	302	A86	C20-O1-C15	10.11	65.38	61.03
26	D	206	A86	C20-O1-C15	9.96	65.31	61.03
26	B	302	A86	C20-O1-C15	9.92	65.29	61.03
26	H	301	A86	C20-O1-C15	9.79	65.23	61.03
26	C	304	A86	O1-C20-C19	-9.71	104.39	113.49
26	C	304	A86	C20-O1-C15	9.68	65.19	61.03
26	B	305	A86	C20-O1-C15	9.67	65.18	61.03
26	H	302	A86	O1-C20-C19	-9.65	104.45	113.49
26	H	302	A86	C3-C4-C5	-9.62	103.83	123.52
26	B	302	A86	C3-C4-C5	-9.59	103.90	123.52
26	B	302	A86	O1-C20-C19	-9.58	104.51	113.49
26	H	302	A86	C20-O1-C15	9.56	65.14	61.03
23	l	207	LMG	O2-C2-C1	9.55	132.83	110.08
26	m	102	A86	C20-O1-C15	9.42	65.08	61.03
26	C	305	A86	C3-C4-C5	-9.41	104.26	123.52
26	b	847	A86	C20-O1-C15	9.39	65.07	61.03
28	E	307	DD6	C14-C13-C11	-9.36	111.01	125.53

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
26	D	204	A86	C20-O1-C15	9.30	65.03	61.03
26	E	302	A86	C20-O1-C15	9.29	65.02	61.03
21	b	833	BCR	C32-C1-C31	-9.28	82.06	108.63
26	B	304	A86	C3-C4-C5	-9.09	104.92	123.52
26	D	204	A86	C4-C3-C2	9.02	141.98	123.52
26	D	206	A86	O1-C20-C19	-8.87	105.18	113.49
28	E	307	DD6	C3-C2-C1	-8.87	114.84	127.28
26	B	304	A86	C20-O1-C15	8.86	64.84	61.03
26	m	102	A86	C24-C1-C2	8.75	132.78	119.01
26	C	302	A86	C3-C4-C5	-8.60	105.93	123.52
26	C	305	A86	O1-C20-C19	-8.43	105.59	113.49
26	H	301	A86	O1-C20-C19	-8.40	105.62	113.49
26	B	305	A86	C3-C4-C5	-8.30	106.54	123.52
26	D	204	A86	C3-C2-C1	8.25	138.85	127.28
26	B	301	A86	C20-O1-C15	8.22	64.56	61.03
23	E	318	LMG	O2-C2-C3	-8.20	91.04	110.38
26	B	302	A86	C4-C3-C2	8.19	140.27	123.52
28	B	303	DD6	C21-C20-C19	8.17	123.42	114.24
26	H	302	A86	C4-C3-C2	8.16	140.22	123.52
26	m	102	A86	O1-C20-C19	-8.11	105.89	113.49
26	H	301	A86	C3-C4-C5	-8.09	106.97	123.52
26	H	304	A86	O1-C20-C19	-8.04	105.96	113.49
23	E	318	LMG	O2-C2-C1	-7.95	91.13	110.08
28	D	205	DD6	C8-C6-C5	7.66	131.05	119.01
28	D	205	DD6	O1-C20-C21	7.59	123.53	115.05
18	b	805	CLA	C4A-NA-C1A	7.58	110.14	106.68
26	H	304	A86	C3-C4-C5	-7.56	108.05	123.52
26	D	204	A86	O1-C20-C21	-7.55	106.61	115.05
26	E	302	A86	O1-C20-C19	-7.53	106.44	113.49
26	B	301	A86	O1-C20-C19	-7.47	106.49	113.49
18	a	854	CLA	C4A-NA-C1A	7.47	110.09	106.68
26	E	302	A86	C3-C4-C5	-7.44	108.30	123.52
18	b	829	CLA	C4A-NA-C1A	7.37	110.04	106.68
18	H	305	CLA	C4A-NA-C1A	7.35	110.03	106.68
18	H	310	CLA	C4A-NA-C1A	7.35	110.03	106.68
18	a	837	CLA	C4A-NA-C1A	7.35	110.03	106.68
26	D	204	A86	C4-C5-C6	7.34	137.57	127.28
26	r	203	A86	C34-O4-C38	-7.31	104.93	117.85
18	a	805	CLA	C4A-NA-C1A	7.29	110.01	106.68
18	b	820	CLA	C4A-NA-C1A	7.29	110.01	106.68
18	a	807	CLA	C4A-NA-C1A	7.25	109.99	106.68
18	a	826	CLA	C4A-NA-C1A	7.24	109.98	106.68

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
18	D	210	CLA	C4A-NA-C1A	7.23	109.98	106.68
18	b	824	CLA	C4A-NA-C1A	7.22	109.97	106.68
26	B	301	A86	C25-C26-C27	-7.20	117.18	127.28
18	b	803	CLA	C4A-NA-C1A	7.13	109.93	106.68
18	a	827	CLA	C4A-NA-C1A	7.12	109.93	106.68
18	a	803	CLA	C4A-NA-C1A	7.11	109.92	106.68
18	C	309	CLA	C4A-NA-C1A	7.09	109.91	106.68
18	E	313	CLA	C4A-NA-C1A	7.08	109.91	106.68
18	a	818	CLA	C4A-NA-C1A	7.07	109.90	106.68
18	b	840	CLA	C4A-NA-C1A	7.01	109.88	106.68
18	b	822	CLA	C4A-NA-C1A	7.01	109.88	106.68
23	a	851	LMG	O1-C1-C2	7.00	118.90	108.27
21	b	833	BCR	C31-C1-C6	7.00	121.21	110.24
26	B	302	A86	C3-C2-C1	6.99	137.09	127.28
18	b	815	CLA	C4A-NA-C1A	6.98	109.86	106.68
26	H	302	A86	C3-C2-C1	6.96	137.04	127.28
18	a	824	CLA	C4A-NA-C1A	6.96	109.85	106.68
18	b	844	CLA	C4A-NA-C1A	6.95	109.85	106.68
28	E	304	DD6	O1-C20-C19	6.95	120.00	113.49
18	b	827	CLA	C4A-NA-C1A	6.95	109.85	106.68
18	B	309	CLA	C4A-NA-C1A	6.94	109.85	106.68
18	H	307	CLA	C4A-NA-C1A	6.94	109.84	106.68
18	a	839	CLA	C4A-NA-C1A	6.94	109.84	106.68
18	b	804	CLA	C4A-NA-C1A	6.93	109.84	106.68
28	H	303	DD6	C24-C1-C2	6.93	129.91	119.01
18	b	838	CLA	C4A-NA-C1A	6.92	109.83	106.68
18	a	820	CLA	C4A-NA-C1A	6.91	109.83	106.68
18	b	850	CLA	C4A-NA-C1A	6.91	109.83	106.68
26	C	302	A86	O1-C20-C19	-6.90	107.02	113.49
18	H	313	CLA	C4A-NA-C1A	6.87	109.81	106.68
18	a	811	CLA	C4A-NA-C1A	6.86	109.81	106.68
18	a	822	CLA	C4A-NA-C1A	6.85	109.81	106.68
18	b	809	CLA	C4A-NA-C1A	6.85	109.80	106.68
18	C	317	CLA	C4A-NA-C1A	6.83	109.79	106.68
18	b	816	CLA	C4A-NA-C1A	6.81	109.79	106.68
18	a	819	CLA	C4A-NA-C1A	6.78	109.77	106.68
26	C	304	A86	C3-C2-C1	-6.78	117.77	127.28
18	b	849	CLA	C4A-NA-C1A	6.78	109.77	106.68
18	a	812	CLA	C4A-NA-C1A	6.76	109.77	106.68
26	B	305	A86	C41-C32-C31	-6.75	104.43	110.47
26	H	301	A86	C4-C3-C2	6.75	137.32	123.52
18	D	214	CLA	C4A-NA-C1A	6.73	109.75	106.68

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
18	H	315	CLA	C4A-NA-C1A	6.73	109.75	106.68
18	H	308	CLA	C4A-NA-C1A	6.73	109.75	106.68
26	b	847	A86	C3-C2-C1	-6.72	117.86	127.28
18	b	817	CLA	C4A-NA-C1A	6.71	109.74	106.68
18	B	310	CLA	C4A-NA-C1A	6.71	109.74	106.68
18	j	104	CLA	C4A-NA-C1A	6.70	109.74	106.68
18	b	813	CLA	C4A-NA-C1A	6.69	109.73	106.68
18	a	808	CLA	C4A-NA-C1A	6.68	109.73	106.68
18	a	813	CLA	C4A-NA-C1A	6.67	109.72	106.68
18	C	306	CLA	C4A-NA-C1A	6.67	109.72	106.68
18	b	842	CLA	C4A-NA-C1A	6.66	109.72	106.68
18	B	312	CLA	C4A-NA-C1A	6.65	109.71	106.68
18	B	308	CLA	C4A-NA-C1A	6.65	109.71	106.68
26	D	204	A86	O1-C20-C19	-6.64	107.27	113.49
18	a	825	CLA	C4A-NA-C1A	6.64	109.71	106.68
18	E	316	CLA	C4A-NA-C1A	6.63	109.70	106.68
18	a	843	CLA	C4A-NA-C1A	6.62	109.70	106.68
28	H	303	DD6	C21-C20-C19	6.62	121.68	114.24
18	a	814	CLA	C4A-NA-C1A	6.62	109.70	106.68
26	D	206	A86	C3-C2-C1	6.62	136.56	127.28
18	B	306	CLA	C4A-NA-C1A	6.61	109.69	106.68
18	b	811	CLA	C4A-NA-C1A	6.61	109.69	106.68
18	D	211	CLA	C4A-NA-C1A	6.61	109.69	106.68
18	B	307	CLA	C4A-NA-C1A	6.60	109.69	106.68
18	E	310	CLA	C4A-NA-C1A	6.60	109.69	106.68
26	B	304	A86	C4-C3-C2	6.60	137.03	123.52
18	a	840	CLA	C4A-NA-C1A	6.60	109.69	106.68
18	b	845	CLA	C4A-NA-C1A	6.59	109.69	106.68
18	a	830	CLA	C4A-NA-C1A	6.59	109.69	106.68
26	r	203	A86	C29-C30-C31	-6.59	169.89	177.66
18	b	818	CLA	C4A-NA-C1A	6.58	109.68	106.68
18	H	311	CLA	C4A-NA-C1A	6.58	109.68	106.68
26	B	305	A86	O1-C20-C19	-6.57	107.33	113.49
18	b	810	CLA	C4A-NA-C1A	6.57	109.68	106.68
18	b	843	CLA	C4A-NA-C1A	6.57	109.68	106.68
18	a	817	CLA	C4A-NA-C1A	6.57	109.68	106.68
18	b	823	CLA	C4A-NA-C1A	6.57	109.68	106.68
18	f	202	CLA	C4A-NA-C1A	6.57	109.68	106.68
18	a	816	CLA	C4A-NA-C1A	6.56	109.67	106.68
18	C	310	CLA	C4A-NA-C1A	6.56	109.67	106.68
18	a	828	CLA	C4A-NA-C1A	6.56	109.67	106.68
18	a	838	CLA	C4A-NA-C1A	6.55	109.67	106.68

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
18	D	209	CLA	C4A-NA-C1A	6.54	109.66	106.68
18	l	203	CLA	C4A-NA-C1A	6.54	109.66	106.68
26	D	206	A86	C3-C4-C5	-6.51	110.20	123.52
18	a	836	CLA	C4A-NA-C1A	6.51	109.65	106.68
18	b	808	CLA	C4A-NA-C1A	6.50	109.65	106.68
18	a	823	CLA	C4A-NA-C1A	6.49	109.64	106.68
18	f	204	CLA	C4A-NA-C1A	6.48	109.63	106.68
26	D	206	A86	C4-C3-C2	6.47	136.77	123.52
18	a	806	CLA	C4A-NA-C1A	6.47	109.63	106.68
18	b	830	CLA	C4A-NA-C1A	6.47	109.63	106.68
26	B	301	A86	C3-C2-C1	-6.47	118.20	127.28
18	a	844	CLA	C4A-NA-C1A	6.46	109.62	106.68
18	b	812	CLA	C4A-NA-C1A	6.46	109.62	106.68
18	E	312	CLA	C4A-NA-C1A	6.46	109.62	106.68
26	r	203	A86	C25-C24-C1	-6.45	108.66	126.36
18	b	807	CLA	C4A-NA-C1A	6.45	109.62	106.68
18	C	318	CLA	C4A-NA-C1A	6.45	109.62	106.68
18	a	855	CLA	C4A-NA-C1A	6.45	109.62	106.68
18	b	825	CLA	C4A-NA-C1A	6.45	109.62	106.68
23	a	851	LMG	O6-C1-O1	-6.44	94.82	110.04
18	a	821	CLA	C4A-NA-C1A	6.44	109.62	106.68
18	B	311	CLA	C4A-NA-C1A	6.43	109.61	106.68
18	b	806	CLA	C4A-NA-C1A	6.43	109.61	106.68
18	a	810	CLA	C4A-NA-C1A	6.43	109.61	106.68
18	b	821	CLA	C4A-NA-C1A	6.42	109.61	106.68
18	D	213	CLA	C4A-NA-C1A	6.42	109.61	106.68
18	b	814	CLA	C4A-NA-C1A	6.42	109.61	106.68
18	D	208	CLA	C4A-NA-C1A	6.41	109.60	106.68
18	b	841	CLA	C4A-NA-C1A	6.41	109.60	106.68
18	b	801	CLA	C4A-NA-C1A	6.39	109.59	106.68
18	C	307	CLA	C4A-NA-C1A	6.39	109.59	106.68
18	b	837	CLA	C4A-NA-C1A	6.39	109.59	106.68
18	b	819	CLA	C4A-NA-C1A	6.39	109.59	106.68
18	C	315	CLA	C4A-NA-C1A	6.39	109.59	106.68
18	H	309	CLA	C4A-NA-C1A	6.39	109.59	106.68
23	C	301	LMG	O2-C2-C3	-6.37	95.35	110.38
18	a	853	CLA	C4A-NA-C1A	6.37	109.59	106.68
18	C	311	CLA	C4A-NA-C1A	6.37	109.59	106.68
26	B	305	A86	C4-C3-C2	6.37	136.55	123.52
18	a	801	CLA	C4A-NA-C1A	6.36	109.58	106.68
18	E	309	CLA	C4A-NA-C1A	6.36	109.58	106.68
18	a	809	CLA	C4A-NA-C1A	6.35	109.58	106.68

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
18	D	212	CLA	C4A-NA-C1A	6.35	109.58	106.68
26	B	305	A86	O1-C20-C21	-6.34	107.96	115.05
18	a	847	CLA	C4A-NA-C1A	6.34	109.57	106.68
18	a	846	CLA	C4A-NA-C1A	6.34	109.57	106.68
27	l	206	ET4	C29-C22-C21	-6.32	111.91	124.18
18	D	215	CLA	C4A-NA-C1A	6.31	109.56	106.68
18	r	202	CLA	C4A-NA-C1A	6.31	109.56	106.68
18	a	804	CLA	C4A-NA-C1A	6.28	109.55	106.68
18	l	205	CLA	C4A-NA-C1A	6.28	109.55	106.68
26	C	305	A86	C4-C3-C2	6.28	136.38	123.52
26	H	304	A86	O1-C20-C21	-6.27	108.04	115.05
18	C	312	CLA	C4A-NA-C1A	6.26	109.53	106.68
21	l	201	BCR	C7-C8-C9	-6.26	116.98	126.23
28	H	303	DD6	C21-C20-C15	-6.26	112.00	122.30
26	r	203	A86	C3-C4-C5	-6.25	110.74	123.52
18	b	848	CLA	C4A-NA-C1A	6.24	109.53	106.68
18	a	848	CLA	C4A-NA-C1A	6.24	109.53	106.68
21	j	105	BCR	C24-C23-C22	-6.24	117.01	126.23
18	E	314	CLA	C4A-NA-C1A	6.23	109.52	106.68
21	b	836	BCR	C16-C17-C18	-6.22	118.55	127.28
18	E	311	CLA	C4A-NA-C1A	6.22	109.52	106.68
28	E	304	DD6	C4-C5-C6	-6.21	118.56	127.28
21	a	852	BCR	C15-C14-C13	-6.21	118.57	127.28
18	a	802	CLA	C4A-NA-C1A	6.19	109.50	106.68
18	f	203	CLA	C4A-NA-C1A	6.19	109.50	106.68
28	D	205	DD6	C21-C20-C19	6.18	121.18	114.24
18	E	315	CLA	C4A-NA-C1A	6.17	109.50	106.68
28	B	303	DD6	C24-C1-C2	6.16	128.70	119.01
18	b	826	CLA	C4A-NA-C1A	6.16	109.49	106.68
18	l	202	CLA	C4A-NA-C1A	6.15	109.49	106.68
18	b	828	CLA	C4A-NA-C1A	6.15	109.48	106.68
18	i	101	CLA	C4A-NA-C1A	6.14	109.48	106.68
18	D	217	CLA	C4A-NA-C1A	6.12	109.47	106.68
18	b	846	CLA	C4A-NA-C1A	6.10	109.46	106.68
26	D	204	A86	C9-C8-C6	-6.09	109.67	126.36
26	r	203	A86	C9-C10-C11	-6.08	109.45	126.64
28	E	307	DD6	C21-C20-C19	6.07	121.06	114.24
26	C	302	A86	C4-C3-C2	6.05	135.90	123.52
18	D	207	CLA	C4A-NA-C1A	6.05	109.44	106.68
26	C	302	A86	O1-C20-C21	-6.04	108.30	115.05
21	m	101	BCR	C7-C8-C9	-6.03	117.31	126.23
28	E	307	DD6	C24-C1-C2	6.03	128.50	119.01

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
18	a	842	CLA	C4A-NA-C1A	6.01	109.42	106.68
26	D	204	A86	C8-C6-C5	5.99	128.44	119.01
21	l	204	BCR	C11-C10-C9	-5.99	118.88	127.28
26	H	302	A86	O1-C20-C21	-5.98	108.36	115.05
21	a	834	BCR	C24-C23-C22	-5.97	117.40	126.23
26	B	302	A86	C9-C10-C11	-5.97	109.78	126.64
18	D	216	CLA	C4A-NA-C1A	5.96	109.40	106.68
26	C	302	A86	C9-C10-C11	-5.95	109.82	126.64
26	H	302	A86	C9-C10-C11	-5.95	109.83	126.64
26	B	302	A86	O1-C20-C21	-5.93	108.42	115.05
26	H	301	A86	O1-C20-C21	-5.91	108.44	115.05
18	a	815	CLA	C4A-NA-C1A	5.89	109.36	106.68
18	H	306	CLA	C4A-NA-C1A	5.88	109.36	106.68
26	D	206	A86	O1-C20-C21	-5.88	108.48	115.05
28	E	307	DD6	C13-C11-C10	5.87	128.25	119.01
18	a	841	CLA	C4A-NA-C1A	5.86	109.35	106.68
18	H	314	CLA	C4A-NA-C1A	5.86	109.35	106.68
26	D	203	A86	C9-C10-C11	-5.86	110.09	126.64
18	C	316	CLA	C4A-NA-C1A	5.85	109.35	106.68
28	C	303	DD6	C3-C2-C1	-5.84	119.09	127.28
18	C	314	CLA	C4A-NA-C1A	5.84	109.34	106.68
21	b	839	BCR	C11-C10-C9	-5.83	119.10	127.28
26	D	203	A86	O1-C20-C21	-5.74	108.63	115.05
26	B	304	A86	C9-C10-C11	-5.68	110.60	126.64
22	a	849	SQD	O8-S-C6	5.67	117.10	106.00
26	b	847	A86	C4-C5-C6	-5.66	119.34	127.28
21	j	105	BCR	C20-C21-C22	-5.66	119.34	127.28
26	H	304	A86	C4-C3-C2	5.63	135.04	123.52
26	D	206	A86	C34-O4-C38	-5.61	107.94	117.85
18	B	314	CLA	C4A-NA-C1A	5.60	109.23	106.68
26	B	305	A86	C9-C10-C11	-5.56	110.92	126.64
21	b	833	BCR	C11-C10-C9	-5.55	119.49	127.28
28	E	304	DD6	O1-C20-C21	5.54	121.24	115.05
21	l	204	BCR	C28-C27-C26	-5.54	104.17	114.06
26	b	847	A86	O1-C20-C21	-5.54	108.86	115.05
21	j	101	BCR	C24-C23-C22	-5.53	118.05	126.23
26	E	302	A86	O1-C20-C21	-5.52	108.88	115.05
26	r	203	A86	O1-C20-C21	-5.51	108.89	115.05
21	b	833	BCR	C32-C1-C2	-5.50	87.84	108.95
21	E	306	BCR	C16-C17-C18	-5.48	119.59	127.28
26	E	302	A86	C9-C10-C11	-5.47	111.17	126.64
23	a	851	LMG	O2-C2-C3	5.45	123.23	110.38

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
21	i	103	BCR	C24-C23-C22	-5.45	118.17	126.23
26	m	102	A86	C25-C24-C1	-5.45	111.42	126.36
26	C	305	A86	C9-C10-C11	-5.44	111.26	126.64
26	H	301	A86	C3-C2-C1	5.44	134.91	127.28
26	B	301	A86	O1-C20-C21	-5.41	109.00	115.05
29	B	313	KC1	C1D-ND-C4D	5.40	110.10	106.31
21	m	101	BCR	C16-C17-C18	-5.38	119.73	127.28
23	j	103	LMG	O6-C1-O1	-5.38	97.34	110.04
28	B	303	DD6	O1-C20-C21	5.34	121.02	115.05
26	r	203	A86	C4-C3-C2	5.32	134.40	123.52
26	E	302	A86	C4-C3-C2	5.29	134.35	123.52
21	b	832	BCR	C16-C17-C18	-5.29	119.86	127.28
26	B	304	A86	O1-C20-C21	-5.28	109.14	115.05
26	m	102	A86	O1-C20-C21	-5.27	109.16	115.05
28	D	205	DD6	C3-C2-C1	-5.27	119.89	127.28
22	a	849	SQD	O8-S-O9	-5.25	98.27	111.40
22	a	849	SQD	O8-S-O7	5.24	124.51	111.40
21	b	833	BCR	C15-C14-C13	-5.23	119.95	127.28
27	l	206	ET4	C19-C18-C17	5.22	127.23	119.01
21	f	201	BCR	C24-C23-C22	-5.19	118.56	126.23
21	l	201	BCR	C33-C5-C6	-5.19	118.82	124.48
26	H	304	A86	C9-C10-C11	-5.18	112.00	126.64
21	j	101	BCR	C20-C21-C22	-5.18	120.02	127.28
26	r	203	A86	O1-C20-C15	-5.17	57.15	59.23
21	f	205	BCR	C15-C14-C13	-5.16	120.04	127.28
29	C	313	KC1	C1D-ND-C4D	5.13	109.91	106.31
23	D	202	LMG	O2-C2-C1	5.12	122.28	110.08
21	b	833	BCR	C33-C5-C6	-5.12	118.90	124.48
29	B	313	KC1	O2D-CGD-CBD	5.11	120.16	111.23
29	H	312	KC1	C1D-ND-C4D	5.09	109.89	106.31
28	H	303	DD6	C-C1-C2	-5.07	114.61	122.82
26	D	206	A86	C9-C10-C11	-5.06	112.34	126.64
26	H	302	A86	O4-C38-C39	5.04	120.08	111.09
26	B	302	A86	O4-C38-C39	5.02	120.05	111.09
26	B	304	A86	O1-C20-C19	-5.01	108.80	113.49
26	C	305	A86	O4-C38-C39	5.00	120.00	111.09
26	C	305	A86	O1-C20-C21	-4.99	109.47	115.05
21	i	103	BCR	C15-C14-C13	-4.99	120.28	127.28
21	a	834	BCR	C20-C21-C22	-4.98	120.29	127.28
21	l	201	BCR	C38-C26-C25	-4.97	119.06	124.48
28	C	303	DD6	C4-C5-C6	-4.94	120.35	127.28
27	l	206	ET4	C24-C05-C06	-4.94	119.10	124.48

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
23	j	103	LMG	O1-C1-C2	4.94	115.77	108.27
26	H	301	A86	O4-C38-C39	4.92	119.87	111.09
21	m	101	BCR	C3-C4-C5	-4.92	105.28	114.06
29	C	308	KC1	C1D-ND-C4D	4.91	109.75	106.31
29	C	308	KC1	O2D-CGD-CBD	4.88	119.75	111.23
26	C	302	A86	O4-C38-C39	4.85	119.74	111.09
29	C	313	KC1	O2D-CGD-CBD	4.85	119.70	111.23
21	a	852	BCR	C16-C17-C18	-4.84	120.49	127.28
26	m	102	A86	C33-C32-C31	4.81	113.89	109.21
21	E	306	BCR	C20-C21-C22	-4.81	120.53	127.28
26	H	301	A86	C33-C32-C31	4.81	113.88	109.21
26	D	204	A86	O4-C38-C39	4.80	119.65	111.09
20	a	835	LHG	O7-C7-C8	4.80	119.65	111.09
28	E	303	DD6	C21-C20-C19	4.80	119.63	114.24
25	b	834	DGD	O2G-C1B-C2B	4.80	121.86	111.48
26	m	102	A86	C25-C26-C27	-4.76	120.60	127.28
21	a	852	BCR	C7-C8-C9	-4.76	119.19	126.23
21	a	850	BCR	C33-C5-C6	-4.76	119.29	124.48
26	B	305	A86	C33-C32-C31	4.75	113.83	109.21
26	m	102	A86	C7-C6-C5	-4.74	115.14	122.82
26	r	203	A86	C35-C34-C33	4.72	118.36	109.89
21	i	102	BCR	C20-C21-C22	-4.72	120.66	127.28
26	B	305	A86	O4-C38-C39	4.71	119.49	111.09
28	E	303	DD6	C9-C10-C11	-4.70	120.69	127.28
26	H	304	A86	C3-C2-C1	4.70	133.87	127.28
21	b	839	BCR	C38-C26-C25	-4.69	119.36	124.48
26	C	305	A86	O1-C20-C15	-4.69	57.34	59.23
26	B	301	A86	O4-C38-C39	4.67	119.42	111.09
26	b	847	A86	O1-C20-C15	-4.66	57.35	59.23
21	i	102	BCR	C16-C17-C18	-4.66	120.74	127.28
26	E	302	A86	O4-C38-C39	4.65	119.37	111.09
21	a	850	BCR	C24-C23-C22	-4.62	119.39	126.23
21	a	834	BCR	C16-C17-C18	-4.62	120.80	127.28
26	H	304	A86	O1-C20-C15	-4.61	57.37	59.23
26	D	203	A86	O4-C38-C39	4.61	119.31	111.09
21	j	105	BCR	C16-C17-C18	-4.60	120.82	127.28
26	m	102	A86	O4-C38-C39	4.59	119.28	111.09
28	E	303	DD6	C3-C2-C1	-4.59	120.84	127.28
28	E	304	DD6	C3-C2-C1	-4.57	120.86	127.28
21	E	305	BCR	C15-C14-C13	-4.57	120.86	127.28
23	j	103	LMG	O2-C2-C3	4.56	121.13	110.38
28	C	303	DD6	C25-C26-C27	-4.55	113.89	126.61

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
21	a	845	BCR	C33-C5-C6	-4.55	119.52	124.48
26	D	203	A86	C34-O4-C38	-4.53	109.84	117.85
18	b	824	CLA	C1-C2-C3	-4.53	118.78	126.20
26	H	301	A86	C9-C10-C11	-4.51	113.89	126.64
26	D	203	A86	C8-C6-C5	-4.49	111.95	119.01
26	r	203	A86	C25-C26-C27	4.48	133.56	127.28
28	D	205	DD6	C28-C27-C26	-4.48	115.48	124.18
21	f	205	BCR	C11-C10-C9	-4.47	121.01	127.28
23	C	301	LMG	C3-C4-C5	4.47	118.33	110.23
26	H	301	A86	O1-C20-C15	-4.47	57.43	59.23
26	C	304	A86	O1-C20-C21	-4.47	110.06	115.05
28	E	304	DD6	C24-C1-C2	4.46	126.03	119.01
28	E	308	DD6	C21-C20-C19	4.45	119.25	114.24
27	l	206	ET4	C30-C18-C17	-4.44	115.62	122.82
26	C	302	A86	O1-C20-C15	-4.44	57.44	59.23
26	m	102	A86	C-C1-C2	-4.43	115.64	122.82
28	D	205	DD6	C7-C6-C5	-4.43	115.64	122.82
21	m	101	BCR	C20-C21-C22	-4.43	121.07	127.28
21	i	102	BCR	C24-C23-C22	-4.42	119.70	126.23
18	b	820	CLA	O2D-CGD-O1D	-4.42	115.25	123.85
26	H	304	A86	O4-C38-C39	4.41	118.96	111.09
26	B	304	A86	O4-C38-C39	4.41	118.96	111.09
29	B	313	KC1	CHB-C1B-NB	4.41	131.01	124.80
23	j	103	LMG	O7-C10-C11	4.41	121.01	111.48
21	a	852	BCR	C20-C21-C22	-4.40	121.10	127.28
20	E	317	LHG	O7-C7-C8	4.40	121.00	111.48
21	l	204	BCR	C24-C23-C22	-4.40	119.73	126.23
26	B	304	A86	C3-C2-C1	4.39	133.43	127.28
18	H	313	CLA	C1-C2-C3	-4.39	119.01	126.20
28	C	303	DD6	C21-C20-C19	4.38	119.16	114.24
18	a	803	CLA	O2D-CGD-O1D	-4.38	115.33	123.85
26	r	203	A86	C10-C9-C8	4.38	135.88	123.20
18	b	801	CLA	O2D-CGD-O1D	-4.38	115.33	123.85
21	a	834	BCR	C15-C14-C13	-4.37	121.15	127.28
29	H	312	KC1	CHB-C1B-NB	4.35	130.93	124.80
20	b	835	LHG	O7-C7-C8	4.34	120.88	111.48
22	a	849	SQD	O9-S-O7	-4.34	99.72	113.82
26	C	305	A86	C3-C2-C1	4.32	133.34	127.28
26	D	206	A86	O4-C38-C39	4.32	118.79	111.09
28	E	303	DD6	C14-C13-C11	-4.32	118.83	125.53
21	f	201	BCR	C33-C5-C6	-4.32	119.78	124.48
21	j	101	BCR	C15-C14-C13	-4.31	121.23	127.28

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
23	D	202	LMG	O7-C10-C11	4.29	120.75	111.48
20	D	201	LHG	O7-C7-C8	4.28	120.74	111.48
26	B	305	A86	O1-C20-C15	-4.28	57.51	59.23
26	B	305	A86	C3-C2-C1	4.27	133.26	127.28
23	C	319	LMG	O7-C10-C11	4.26	120.70	111.48
26	m	102	A86	C-C1-C24	-4.26	111.58	118.09
26	D	206	A86	C10-C9-C8	4.25	135.53	123.20
26	D	206	A86	O1-C20-C15	-4.25	57.52	59.23
21	f	201	BCR	C20-C21-C22	-4.24	121.33	127.28
18	l	203	CLA	O2D-CGD-O1D	-4.24	115.60	123.85
21	a	833	BCR	C28-C27-C26	-4.24	106.50	114.06
26	B	305	A86	C35-C34-C33	4.23	117.49	109.89
26	D	204	A86	O1-C20-C15	-4.23	57.53	59.23
26	D	204	A86	C26-C25-C24	-4.22	110.96	123.20
21	i	103	BCR	C11-C10-C9	-4.21	121.37	127.28
26	C	304	A86	O1-C20-C15	-4.21	57.54	59.23
18	b	837	CLA	O2D-CGD-O1D	-4.21	115.66	123.85
21	b	832	BCR	C20-C21-C22	-4.21	121.38	127.28
26	H	302	A86	C10-C9-C8	4.20	135.38	123.20
29	H	312	KC1	O2D-CGD-CBD	4.20	118.58	111.23
26	B	302	A86	O1-C20-C15	-4.20	57.54	59.23
21	r	201	BCR	C3-C4-C5	-4.20	106.57	114.06
26	b	847	A86	O4-C38-C39	4.20	118.58	111.09
26	H	304	A86	C26-C25-C24	-4.20	111.04	123.20
26	B	302	A86	C10-C9-C8	4.20	135.36	123.20
26	B	305	A86	C10-C9-C8	4.20	135.36	123.20
28	C	303	DD6	C3-C4-C5	-4.19	114.94	123.52
28	E	303	DD6	C4-C5-C6	-4.19	121.40	127.28
23	a	851	LMG	O7-C10-C11	4.19	120.54	111.48
26	C	304	A86	O4-C38-C39	4.18	118.55	111.09
26	D	204	A86	C7-C6-C5	-4.17	116.06	122.82
26	E	302	A86	O1-C20-C15	-4.17	57.55	59.23
21	E	306	BCR	C24-C23-C22	-4.16	120.07	126.23
21	i	103	BCR	C20-C21-C22	-4.16	121.44	127.28
26	m	102	A86	O1-C20-C15	-4.13	57.57	59.23
20	a	832	LHG	O7-C7-C8	4.13	120.42	111.48
21	b	833	BCR	C24-C23-C22	-4.13	120.12	126.23
21	m	101	BCR	C24-C23-C22	-4.13	120.13	126.23
28	E	304	DD6	C-C1-C2	-4.13	116.13	122.82
21	b	833	BCR	C38-C26-C25	-4.12	119.99	124.48
20	E	301	LHG	O7-C7-C8	4.12	120.39	111.48
28	E	307	DD6	C7-C6-C5	-4.12	116.14	122.82

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
26	D	204	A86	C9-C10-C11	-4.11	115.01	126.64
21	m	101	BCR	C4-C5-C6	-4.11	117.15	122.70
28	C	303	DD6	C37-C36-C35	4.10	121.96	114.42
26	r	203	A86	C24-C1-C2	4.09	125.44	119.01
21	i	103	BCR	C7-C8-C9	-4.09	120.18	126.23
21	j	101	BCR	C11-C10-C9	-4.08	121.55	127.28
21	b	839	BCR	C1-C6-C5	-4.08	117.06	122.64
21	a	833	BCR	C16-C17-C18	-4.07	121.56	127.28
18	C	317	CLA	C3B-C4B-NB	-4.07	106.90	110.53
21	f	205	BCR	C7-C8-C9	-4.05	120.25	126.23
29	B	313	KC1	O1D-CGD-CBD	-4.05	116.54	124.52
28	B	303	DD6	C14-C13-C11	-4.04	119.26	125.53
26	C	304	A86	C4-C5-C6	-4.03	121.62	127.28
26	H	302	A86	O1-C20-C15	-4.03	57.61	59.23
21	b	832	BCR	C28-C27-C26	-4.01	106.91	114.06
26	C	302	A86	C3-C2-C1	4.01	132.90	127.28
23	C	301	LMG	O7-C10-C11	4.00	120.14	111.48
20	j	102	LHG	O7-C7-C8	4.00	120.14	111.48
26	b	847	A86	O1-C20-C19	-4.00	109.74	113.49
26	B	304	A86	C10-C9-C8	4.00	134.79	123.20
21	E	306	BCR	C15-C14-C13	-3.99	121.68	127.28
26	D	203	A86	C10-C9-C8	3.98	134.75	123.20
28	D	205	DD6	O1-C15-C14	-3.98	105.46	116.88
26	B	305	A86	C26-C25-C24	-3.98	111.66	123.20
29	C	308	KC1	CHB-C1B-NB	3.98	130.40	124.80
20	H	316	LHG	O7-C7-C8	3.97	120.06	111.48
26	H	301	A86	C41-C32-C31	-3.96	106.92	110.47
21	a	850	BCR	C20-C21-C22	-3.96	121.72	127.28
26	H	302	A86	C26-C25-C24	-3.96	111.73	123.20
28	E	304	DD6	O2-C18-C19	-3.95	101.66	109.75
26	D	203	A86	C7-C6-C8	3.95	124.13	118.09
26	B	302	A86	C26-C25-C24	-3.94	111.78	123.20
29	C	313	KC1	CHB-C1B-NB	3.94	130.35	124.80
26	D	203	A86	O1-C20-C15	-3.94	57.65	59.23
26	C	305	A86	C26-C25-C24	-3.93	111.80	123.20
29	C	308	KC1	CHC-C4B-NB	3.93	130.34	124.80
21	b	836	BCR	C3-C4-C5	-3.93	107.05	114.06
28	B	303	DD6	C4-C5-C6	-3.93	121.77	127.28
28	C	303	DD6	C14-C13-C11	-3.92	119.44	125.53
21	j	101	BCR	C16-C17-C18	-3.91	121.80	127.28
21	j	105	BCR	C15-C14-C13	-3.90	121.80	127.28
26	B	301	A86	O1-C20-C15	-3.90	57.66	59.23

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
23	l	207	LMG	O7-C10-C11	3.90	119.92	111.48
29	H	312	KC1	CHC-C4B-NB	3.90	130.29	124.80
21	a	850	BCR	C16-C17-C18	-3.90	121.81	127.28
21	i	103	BCR	C16-C17-C18	-3.89	121.82	127.28
28	D	205	DD6	C24-C1-C2	3.89	125.13	119.01
28	C	303	DD6	C15-C14-C13	-3.89	117.77	125.99
21	a	833	BCR	C15-C14-C13	-3.89	121.83	127.28
18	a	813	CLA	CAC-C3C-C4C	3.88	129.84	124.79
26	B	301	A86	C7-C6-C8	3.88	124.01	118.09
26	C	304	A86	C-C1-C2	-3.88	116.54	122.82
18	D	208	CLA	C1-C2-C3	-3.85	119.90	126.20
21	i	102	BCR	C7-C8-C9	-3.84	120.55	126.23
21	f	201	BCR	C1-C6-C5	-3.84	117.39	122.64
21	a	845	BCR	C38-C26-C25	-3.84	120.29	124.48
18	C	318	CLA	C3B-C4B-NB	-3.83	107.11	110.53
28	E	308	DD6	C4-C5-C6	-3.83	121.90	127.28
21	a	852	BCR	C24-C23-C22	-3.83	120.58	126.23
28	E	307	DD6	C12-C11-C10	-3.82	116.62	122.82
26	B	304	A86	C26-C25-C24	-3.82	112.14	123.20
21	l	201	BCR	C15-C14-C13	-3.82	121.93	127.28
26	C	305	A86	C34-O4-C38	-3.82	111.11	117.85
19	b	831	PQN	C11-C12-C13	-3.82	120.26	126.83
28	E	308	DD6	C3-C2-C1	-3.81	121.93	127.28
21	j	105	BCR	C7-C8-C9	-3.81	120.59	126.23
18	H	307	CLA	O2D-CGD-O1D	-3.81	116.43	123.85
21	l	204	BCR	C8-C9-C10	3.80	124.99	119.01
28	E	307	DD6	C25-C24-C1	-3.79	115.97	126.36
26	B	301	A86	C34-O4-C38	-3.79	111.16	117.85
21	j	105	BCR	C28-C27-C26	-3.78	107.31	114.06
18	a	825	CLA	O2D-CGD-O1D	-3.78	116.49	123.85
18	H	310	CLA	O2D-CGD-O1D	-3.78	116.49	123.85
26	m	102	A86	C9-C10-C11	-3.77	115.98	126.64
21	j	105	BCR	C11-C10-C9	-3.76	122.00	127.28
18	l	202	CLA	O2D-CGD-O1D	-3.75	116.54	123.85
28	E	304	DD6	C21-C20-C19	3.75	118.46	114.24
21	a	833	BCR	C24-C23-C22	-3.75	120.69	126.23
18	a	842	CLA	O2D-CGD-O1D	-3.75	116.56	123.85
18	H	308	CLA	O2D-CGD-O1D	-3.74	116.57	123.85
27	l	206	ET4	C27-C09-C10	-3.74	116.76	122.82
23	E	318	LMG	O7-C10-C11	3.74	119.57	111.48
29	C	313	KC1	CHC-C4B-NB	3.73	130.06	124.80
26	B	301	A86	C4-C5-C6	-3.73	122.05	127.28

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
21	a	852	BCR	C11-C10-C9	-3.73	122.05	127.28
18	a	803	CLA	O2D-CGD-CBD	3.73	117.74	111.23
21	E	305	BCR	C28-C27-C26	-3.73	107.41	114.06
21	a	852	BCR	C33-C5-C6	-3.72	120.43	124.48
28	E	308	DD6	C9-C10-C11	-3.71	122.07	127.28
21	f	205	BCR	C16-C17-C18	-3.71	122.07	127.28
20	a	831	LHG	O7-C7-C8	3.71	119.51	111.48
26	C	302	A86	C7-C6-C8	3.71	123.76	118.09
29	B	313	KC1	CHD-C1D-ND	3.71	129.61	124.05
22	a	849	SQD	O9-S-C6	-3.71	101.13	106.73
21	r	201	BCR	C16-C17-C18	-3.71	122.08	127.28
21	f	205	BCR	C20-C21-C22	-3.70	122.09	127.28
18	D	208	CLA	C3B-C4B-NB	-3.69	107.23	110.53
21	a	834	BCR	C33-C5-C6	-3.69	120.46	124.48
18	a	825	CLA	C3B-C4B-NB	-3.69	107.24	110.53
28	E	307	DD6	C4-C5-C6	-3.68	122.12	127.28
26	C	302	A86	C8-C6-C5	-3.68	113.22	119.01
18	D	208	CLA	O2D-CGD-O1D	-3.68	116.69	123.85
18	D	207	CLA	C3B-C4B-NB	-3.67	107.25	110.53
18	b	843	CLA	O2D-CGD-O1D	-3.67	116.71	123.85
18	C	314	CLA	C3B-C4B-NB	-3.66	107.26	110.53
26	D	206	A86	C35-C34-C33	3.66	116.45	109.89
18	a	813	CLA	O2D-CGD-O1D	-3.65	116.74	123.85
21	r	201	BCR	C24-C23-C22	-3.65	120.84	126.23
21	j	101	BCR	C33-C5-C6	-3.64	120.51	124.48
18	D	210	CLA	O2D-CGD-O1D	-3.64	116.77	123.85
18	a	818	CLA	O2D-CGD-O1D	-3.64	116.77	123.85
21	a	852	BCR	C38-C26-C25	-3.63	120.52	124.48
21	i	102	BCR	C11-C10-C9	-3.63	122.19	127.28
18	b	811	CLA	O2D-CGD-O1D	-3.62	116.79	123.85
18	a	841	CLA	O2D-CGD-O1D	-3.62	116.80	123.85
29	H	312	KC1	CHD-C1D-ND	3.61	129.47	124.05
21	a	850	BCR	C15-C14-C13	-3.61	122.22	127.28
18	a	807	CLA	O2D-CGD-O1D	-3.60	116.83	123.85
26	B	305	A86	C34-O4-C38	-3.60	111.50	117.85
21	j	105	BCR	C33-C5-C6	-3.59	120.56	124.48
26	H	304	A86	C41-C32-C31	-3.59	107.26	110.47
21	a	845	BCR	C7-C8-C9	-3.59	120.93	126.23
21	r	201	BCR	C20-C21-C22	-3.57	122.27	127.28
28	D	205	DD6	C34-C35-C36	-3.57	103.29	112.18
26	D	204	A86	C19-C18-C17	-3.57	104.12	110.79
18	D	207	CLA	O2D-CGD-O1D	-3.57	116.91	123.85

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
21	j	101	BCR	C38-C26-C25	-3.56	120.60	124.48
18	C	312	CLA	C3B-C4B-NB	-3.56	107.35	110.53
28	E	304	DD6	C37-C36-C35	3.56	120.96	114.42
20	B	315	LHG	O7-C7-C8	3.55	119.17	111.48
28	B	303	DD6	O1-C15-C14	-3.55	106.70	116.88
21	b	836	BCR	C7-C8-C9	-3.55	120.98	126.23
27	l	206	ET4	C07-C06-C05	-3.54	113.39	121.56
26	H	301	A86	C7-C6-C8	3.54	123.50	118.09
21	m	101	BCR	C2-C1-C6	3.54	115.59	110.44
18	H	311	CLA	O2D-CGD-O1D	-3.54	116.96	123.85
18	l	203	CLA	O2D-CGD-CBD	3.53	117.41	111.23
26	B	304	A86	O1-C20-C15	-3.53	57.81	59.23
21	i	103	BCR	C28-C27-C26	-3.53	107.76	114.06
21	a	845	BCR	C24-C23-C22	-3.53	121.02	126.23
29	C	313	KC1	CHD-C1D-ND	3.53	129.34	124.05
21	E	305	BCR	C33-C5-C6	-3.53	120.64	124.48
21	E	306	BCR	C33-C5-C6	-3.53	120.64	124.48
18	b	829	CLA	O2D-CGD-O1D	-3.52	116.99	123.85
21	a	834	BCR	C38-C26-C25	-3.52	120.64	124.48
21	b	833	BCR	C7-C8-C9	-3.52	121.03	126.23
21	l	204	BCR	C3-C4-C5	-3.51	107.80	114.06
21	b	836	BCR	C28-C27-C26	-3.51	107.80	114.06
18	H	311	CLA	C3B-C4B-NB	-3.51	107.40	110.53
18	a	837	CLA	O2D-CGD-O1D	-3.50	117.03	123.85
26	B	301	A86	C25-C24-C1	-3.50	116.77	126.36
21	a	833	BCR	C3-C4-C5	-3.50	107.82	114.06
26	E	302	A86	C26-C25-C24	-3.49	113.09	123.20
27	l	206	ET4	C01-C06-C05	-3.49	117.87	122.64
26	D	203	A86	C26-C25-C24	3.49	133.30	123.20
21	b	832	BCR	C15-C14-C13	-3.48	122.39	127.28
18	C	307	CLA	C3B-C4B-NB	-3.48	107.42	110.53
18	a	854	CLA	C3B-C4B-NB	-3.48	107.42	110.53
26	D	206	A86	C26-C25-C24	-3.48	113.12	123.20
23	C	319	LMG	O8-C28-C29	3.47	122.42	111.83
21	E	305	BCR	C11-C10-C9	-3.46	122.42	127.28
21	j	101	BCR	C7-C8-C9	-3.46	121.12	126.23
26	B	301	A86	C9-C10-C11	-3.46	116.87	126.64
21	a	850	BCR	C38-C26-C25	-3.46	120.71	124.48
26	H	304	A86	C7-C6-C8	3.45	123.36	118.09
26	C	305	A86	C7-C6-C8	3.45	123.36	118.09
21	E	306	BCR	C11-C10-C9	-3.45	122.45	127.28
19	a	829	PQN	C14-C13-C15	3.44	121.20	115.23

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
26	H	304	A86	C10-C9-C8	3.44	133.17	123.20
21	i	102	BCR	C15-C14-C13	-3.44	122.45	127.28
18	a	837	CLA	O2D-CGD-CBD	3.44	117.24	111.23
21	b	833	BCR	C2-C1-C6	3.44	115.44	110.44
29	C	308	KC1	CHD-C1D-ND	3.44	129.21	124.05
21	f	205	BCR	C24-C23-C22	-3.44	121.15	126.23
26	E	302	A86	C7-C6-C8	3.43	123.33	118.09
21	E	306	BCR	C38-C26-C25	-3.43	120.75	124.48
28	H	303	DD6	C3-C2-C1	-3.42	122.49	127.28
21	j	101	BCR	C28-C27-C26	-3.42	107.96	114.06
28	H	303	DD6	C34-C35-C36	-3.41	103.68	112.18
29	B	313	KC1	CHC-C4B-NB	3.41	129.60	124.80
18	l	202	CLA	C3B-C4B-NB	-3.41	107.49	110.53
28	B	303	DD6	C-C1-C2	-3.41	117.30	122.82
28	D	205	DD6	C9-C8-C6	-3.40	117.04	126.36
26	b	847	A86	C29-C30-C31	-3.40	173.65	177.66
18	a	821	CLA	O2D-CGD-O1D	-3.39	117.24	123.85
18	E	312	CLA	C3B-C4B-NB	-3.39	107.50	110.53
21	a	845	BCR	C11-C10-C9	-3.39	122.52	127.28
21	f	205	BCR	C33-C5-C6	-3.39	120.79	124.48
21	b	836	BCR	C20-C21-C22	-3.39	122.53	127.28
21	b	839	BCR	C24-C23-C22	-3.39	121.22	126.23
26	C	305	A86	C10-C9-C8	3.39	133.01	123.20
26	H	301	A86	C35-C34-C33	3.38	115.96	109.89
26	C	302	A86	C26-C25-C24	-3.37	113.43	123.20
21	l	204	BCR	C33-C5-C6	-3.37	120.81	124.48
21	f	205	BCR	C28-C27-C26	-3.37	108.06	114.06
26	E	302	A86	C10-C9-C8	3.36	132.94	123.20
28	E	307	DD6	C-C1-C24	-3.36	112.96	118.09
21	b	832	BCR	C3-C4-C5	-3.36	108.07	114.06
18	b	820	CLA	O2D-CGD-CBD	3.36	117.10	111.23
18	b	829	CLA	C3B-C4B-NB	-3.36	107.53	110.53
26	C	302	A86	C34-O4-C38	-3.36	111.92	117.85
20	B	315	LHG	O8-C23-C24	3.34	122.02	111.83
28	D	205	DD6	C4-C3-C2	3.34	130.35	123.52
21	b	832	BCR	C7-C8-C9	-3.34	121.30	126.23
18	a	840	CLA	C3B-C4B-NB	-3.33	107.55	110.53
26	b	847	A86	O-C13-C11	-3.33	113.86	121.04
18	H	306	CLA	O2D-CGD-O1D	-3.33	117.37	123.85
20	E	317	LHG	C5-O7-C7	-3.33	109.84	117.80
26	B	302	A86	O-C13-C11	-3.32	113.88	121.04
28	B	303	DD6	C15-C14-C13	-3.32	118.97	125.99

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
26	D	203	A86	C4-C3-C2	-3.32	116.73	123.52
21	b	833	BCR	C31-C1-C2	3.31	121.67	108.95
20	a	831	LHG	O8-C23-C24	3.31	121.93	111.83
29	B	313	KC1	C2D-C1D-ND	-3.30	106.90	110.33
18	C	314	CLA	O2D-CGD-O1D	-3.30	117.42	123.85
21	a	845	BCR	C16-C17-C18	-3.30	122.64	127.28
18	C	309	CLA	O2D-CGD-O1D	-3.30	117.42	123.85
21	a	833	BCR	C7-C8-C9	-3.29	121.37	126.23
18	b	837	CLA	C3B-C4B-NB	-3.29	107.59	110.53
18	a	828	CLA	O2D-CGD-O1D	-3.29	117.45	123.85
26	H	302	A86	O-C13-C11	-3.29	113.97	121.04
21	b	832	BCR	C11-C10-C9	-3.28	122.68	127.28
18	b	811	CLA	C3B-C4B-NB	-3.28	107.61	110.53
18	C	316	CLA	C3B-C4B-NB	-3.28	107.61	110.53
25	b	834	DGD	C2G-O2G-C1B	-3.27	109.96	117.80
21	j	105	BCR	C38-C26-C25	-3.27	120.91	124.48
21	b	832	BCR	C24-C23-C22	-3.27	121.39	126.23
21	i	103	BCR	C33-C5-C6	-3.27	120.92	124.48
18	E	314	CLA	C3B-C4B-NB	-3.27	107.61	110.53
26	C	302	A86	C10-C9-C8	3.26	132.66	123.20
21	b	833	BCR	C15-C16-C17	-3.26	116.84	123.52
18	a	842	CLA	O1D-CGD-CBD	3.26	130.95	124.52
18	b	824	CLA	C3B-C4B-NB	-3.26	107.62	110.53
18	b	840	CLA	O2D-CGD-O1D	-3.26	117.51	123.85
21	b	836	BCR	C11-C10-C9	-3.26	122.71	127.28
18	C	312	CLA	O2D-CGD-O1D	-3.26	117.51	123.85
21	E	306	BCR	C7-C8-C9	-3.25	121.43	126.23
18	a	820	CLA	O2D-CGD-O1D	-3.25	117.53	123.85
21	b	839	BCR	C3-C4-C5	-3.25	108.27	114.06
26	H	301	A86	C10-C9-C8	3.24	132.58	123.20
29	H	312	KC1	CHB-C1B-C2B	-3.24	118.76	125.49
28	H	303	DD6	C9-C10-C11	-3.23	122.75	127.28
21	b	836	BCR	C16-C15-C14	-3.23	116.92	123.52
18	b	837	CLA	O2D-CGD-CBD	3.22	116.86	111.23
28	D	205	DD6	C26-C25-C24	3.22	132.53	123.20
18	H	305	CLA	C3B-C4B-NB	-3.21	107.66	110.53
21	a	852	BCR	C28-C27-C26	-3.21	108.33	114.06
26	b	847	A86	C12-C11-C13	3.21	121.20	116.00
26	m	102	A86	C4-C3-C2	-3.20	116.96	123.52
26	m	102	A86	C8-C6-C5	3.20	124.05	119.01
18	E	314	CLA	O2D-CGD-O1D	-3.20	117.61	123.85
18	a	813	CLA	C3B-C4B-NB	-3.20	107.67	110.53

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
21	E	306	BCR	C3-C4-C5	-3.20	108.35	114.06
21	a	833	BCR	C38-C26-C25	-3.20	120.99	124.48
18	H	307	CLA	O2D-CGD-CBD	3.20	116.82	111.23
26	D	204	A86	C-C1-C24	3.20	122.97	118.09
18	a	809	CLA	O2D-CGD-O1D	-3.19	117.64	123.85
26	b	847	A86	C10-C9-C8	-3.19	113.96	123.20
18	l	203	CLA	C3B-C4B-NB	-3.19	107.68	110.53
18	B	314	CLA	C3B-C4B-NB	-3.19	107.69	110.53
21	E	305	BCR	C3-C4-C5	-3.19	108.38	114.06
26	D	204	A86	C24-C1-C2	-3.18	114.00	119.01
18	i	101	CLA	C1-C2-C3	-3.18	120.98	126.20
26	E	302	A86	C8-C6-C5	-3.18	114.01	119.01
26	B	304	A86	C34-O4-C38	-3.18	112.24	117.85
18	H	308	CLA	C3B-C4B-NB	-3.17	107.70	110.53
20	D	201	LHG	O8-C23-C24	3.17	121.50	111.83
18	H	314	CLA	O2D-CGD-O1D	-3.17	117.69	123.85
18	H	306	CLA	C3B-C4B-NB	-3.16	107.71	110.53
18	D	208	CLA	CAA-CBA-CGA	-3.15	104.25	113.21
18	C	311	CLA	C3B-C4B-NB	-3.15	107.72	110.53
28	B	303	DD6	C32-C33-C34	-3.15	106.69	113.59
20	E	301	LHG	O8-C23-C24	3.14	121.41	111.83
18	a	803	CLA	C3B-C4B-NB	-3.14	107.73	110.53
21	a	852	BCR	C11-C12-C13	-3.14	117.76	126.36
18	b	823	CLA	O2D-CGD-O1D	-3.13	117.75	123.85
18	b	848	CLA	O2D-CGD-O1D	-3.13	117.75	123.85
28	D	205	DD6	C7-C6-C8	-3.13	113.30	118.09
28	D	205	DD6	C19-C18-C17	-3.13	104.94	110.79
21	f	205	BCR	C38-C26-C25	-3.13	121.07	124.48
26	C	302	A86	C19-C18-C17	-3.12	104.96	110.79
18	a	804	CLA	O2D-CGD-O1D	-3.12	117.78	123.85
18	b	818	CLA	O2D-CGD-O1D	-3.12	117.78	123.85
18	a	813	CLA	C1-C2-C3	-3.11	121.09	126.20
21	m	101	BCR	C16-C15-C14	-3.11	117.15	123.52
18	b	844	CLA	O2D-CGD-O1D	-3.11	117.79	123.85
26	D	204	A86	C34-O4-C38	-3.11	112.35	117.85
18	a	842	CLA	C3B-C4B-NB	-3.11	107.75	110.53
21	b	839	BCR	C30-C25-C26	-3.10	118.40	122.64
26	E	302	A86	C34-O4-C38	-3.10	112.37	117.85
26	r	203	A86	O4-C38-C39	3.09	116.61	111.09
18	a	837	CLA	C3B-C4B-NB	-3.09	107.77	110.53
18	a	826	CLA	O2D-CGD-O1D	-3.09	117.84	123.85
18	B	312	CLA	C3B-C4B-NB	-3.09	107.78	110.53

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
18	D	216	CLA	O2D-CGD-O1D	-3.09	117.84	123.85
18	H	313	CLA	C3B-C4B-NB	-3.08	107.78	110.53
21	a	845	BCR	C15-C14-C13	-3.08	122.95	127.28
28	C	303	DD6	O1-C20-C19	-3.08	110.61	113.49
29	B	313	KC1	CHB-C1B-C2B	-3.08	119.09	125.49
18	b	804	CLA	O2D-CGD-O1D	-3.08	117.86	123.85
18	b	843	CLA	C3B-C4B-NB	-3.07	107.79	110.53
26	B	305	A86	C7-C6-C8	3.07	122.78	118.09
18	a	853	CLA	O2D-CGD-O1D	-3.07	117.87	123.85
18	b	814	CLA	C3B-C4B-NB	-3.07	107.79	110.53
21	E	305	BCR	C7-C8-C9	-3.07	121.70	126.23
28	B	303	DD6	C26-C25-C24	-3.07	114.32	123.20
26	C	302	A86	O-C13-C11	-3.06	114.44	121.04
18	B	310	CLA	O2D-CGD-O1D	-3.06	117.89	123.85
21	a	845	BCR	C29-C30-C25	-3.06	106.00	110.44
27	l	206	ET4	O40-C36-C35	3.05	115.98	109.75
21	l	204	BCR	C15-C14-C13	-3.05	123.00	127.28
23	D	202	LMG	O8-C28-C29	3.05	121.12	111.83
18	b	827	CLA	O2D-CGD-O1D	-3.04	117.92	123.85
28	H	303	DD6	C13-C11-C10	3.04	123.80	119.01
26	D	203	A86	O1-C15-C20	-3.04	56.87	59.45
21	f	201	BCR	C3-C4-C5	-3.04	108.63	114.06
21	b	839	BCR	C8-C9-C10	3.04	123.79	119.01
21	b	836	BCR	C33-C5-C6	-3.04	121.17	124.48
18	b	838	CLA	O2D-CGD-O1D	-3.04	117.93	123.85
23	l	207	LMG	O8-C28-C29	3.04	121.10	111.83
18	b	816	CLA	O2D-CGD-O1D	-3.04	117.93	123.85
26	H	301	A86	C25-C26-C27	-3.04	123.02	127.28
29	C	308	KC1	O1D-CGD-CBD	-3.03	118.53	124.52
26	B	304	A86	C21-C20-C15	3.03	133.15	123.35
18	b	805	CLA	O2D-CGD-O1D	-3.03	117.94	123.85
29	C	313	KC1	C2D-C1D-ND	-3.03	107.18	110.33
21	l	201	BCR	C24-C23-C22	-3.03	121.75	126.23
18	a	839	CLA	O2D-CGD-O1D	-3.03	117.95	123.85
18	j	104	CLA	CAA-C2A-C3A	-3.03	109.28	116.23
18	b	841	CLA	O2D-CGD-O1D	-3.03	117.95	123.85
21	m	101	BCR	C10-C11-C12	-3.03	114.42	123.20
18	b	807	CLA	O2D-CGD-O1D	-3.03	117.95	123.85
27	l	206	ET4	C16-C15-C14	3.03	129.71	123.52
18	C	307	CLA	O2D-CGD-O1D	-3.03	117.96	123.85
18	b	801	CLA	O2D-CGD-CBD	3.03	116.52	111.23
18	a	855	CLA	O2D-CGD-O1D	-3.02	117.96	123.85

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
18	H	313	CLA	O2D-CGD-O1D	-3.02	117.96	123.85
18	a	819	CLA	O2D-CGD-O1D	-3.02	117.97	123.85
18	a	844	CLA	O2D-CGD-O1D	-3.02	117.97	123.85
23	j	103	LMG	O8-C28-C29	3.02	121.04	111.83
27	l	206	ET4	C01-C06-C07	3.02	123.83	115.65
21	m	101	BCR	C34-C9-C10	-3.02	117.93	122.82
26	m	102	A86	C10-C9-C8	3.02	131.94	123.20
28	B	303	DD6	C3-C2-C1	-3.01	123.05	127.28
18	b	830	CLA	O2D-CGD-O1D	-3.01	117.98	123.85
23	C	301	LMG	O8-C28-C29	3.01	121.02	111.83
26	r	203	A86	C26-C25-C24	3.01	131.92	123.20
18	a	843	CLA	O2D-CGD-O1D	-3.01	117.99	123.85
21	i	103	BCR	C3-C4-C5	-3.01	108.69	114.06
23	E	318	LMG	O8-C28-C29	3.00	120.98	111.83
18	D	213	CLA	C3B-C4B-NB	-3.00	107.85	110.53
18	a	840	CLA	O2D-CGD-O1D	-3.00	118.01	123.85
18	i	101	CLA	C3B-C4B-NB	-2.99	107.86	110.53
21	l	201	BCR	C3-C4-C5	-2.99	108.72	114.06
18	a	818	CLA	C3B-C4B-NB	-2.99	107.86	110.53
26	m	102	A86	C34-O4-C38	-2.99	112.56	117.85
28	C	303	DD6	C-C1-C2	-2.99	117.97	122.82
26	b	847	A86	C41-C32-C31	-2.99	107.80	110.47
26	D	203	A86	C9-C8-C6	2.99	134.55	126.36
18	a	807	CLA	C3B-C4B-NB	-2.98	107.87	110.53
18	b	822	CLA	O2D-CGD-O1D	-2.98	118.05	123.85
18	b	801	CLA	C1-C2-C3	-2.98	121.31	126.20
18	a	815	CLA	O2D-CGD-O1D	-2.97	118.06	123.85
18	B	307	CLA	C3B-C4B-NB	-2.97	107.88	110.53
26	E	302	A86	C19-C18-C17	-2.97	105.23	110.79
18	a	830	CLA	O2D-CGD-O1D	-2.97	118.07	123.85
18	H	308	CLA	O2D-CGD-CBD	2.97	116.42	111.23
18	l	202	CLA	O2D-CGD-CBD	2.96	116.41	111.23
18	b	828	CLA	C3B-C4B-NB	-2.96	107.89	110.53
18	b	842	CLA	O2D-CGD-O1D	-2.96	118.08	123.85
21	b	839	BCR	C4-C5-C6	-2.96	118.70	122.70
21	b	833	BCR	C20-C21-C22	-2.96	123.13	127.28
20	E	317	LHG	O8-C23-C24	2.96	120.86	111.83
26	r	203	A86	C12-C11-C10	-2.96	116.48	123.67
18	b	850	CLA	O2D-CGD-O1D	-2.96	118.09	123.85
21	l	201	BCR	C16-C17-C18	-2.96	123.13	127.28
28	E	303	DD6	C21-C20-C15	-2.96	117.44	122.30
18	B	307	CLA	O2D-CGD-O1D	-2.96	118.10	123.85

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
23	E	318	LMG	O1-C7-C8	-2.95	103.63	110.82
18	D	214	CLA	O2D-CGD-O1D	-2.95	118.10	123.85
18	D	210	CLA	O2D-CGD-CBD	2.95	116.39	111.23
26	r	203	A86	O-C13-C14	-2.95	115.71	121.76
21	r	201	BCR	C15-C14-C13	-2.95	123.14	127.28
18	b	809	CLA	C3B-C4B-NB	-2.95	107.89	110.53
18	b	810	CLA	O2D-CGD-O1D	-2.95	118.10	123.85
18	a	854	CLA	O2A-CGA-O1A	-2.95	116.25	123.63
18	B	309	CLA	O2D-CGD-O1D	-2.95	118.11	123.85
18	b	826	CLA	O2D-CGD-O1D	-2.95	118.11	123.85
21	i	103	BCR	C38-C26-C25	-2.95	121.27	124.48
18	H	315	CLA	O2D-CGD-O1D	-2.95	118.11	123.85
18	b	812	CLA	O2D-CGD-O1D	-2.94	118.12	123.85
18	j	104	CLA	C3B-C4B-NB	-2.94	107.90	110.53
18	b	849	CLA	C3B-C4B-NB	-2.94	107.90	110.53
21	b	839	BCR	C16-C17-C18	-2.94	123.15	127.28
21	f	201	BCR	C28-C27-C26	-2.94	108.81	114.06
18	a	836	CLA	O2D-CGD-O1D	-2.94	118.13	123.85
21	f	201	BCR	C16-C17-C18	-2.94	123.16	127.28
18	a	816	CLA	O2D-CGD-O1D	-2.94	118.13	123.85
18	D	207	CLA	O2D-CGD-CBD	2.94	116.36	111.23
18	b	809	CLA	O2D-CGD-O1D	-2.94	118.13	123.85
18	a	806	CLA	O2D-CGD-O1D	-2.94	118.13	123.85
21	f	201	BCR	C1-C6-C7	2.94	123.61	115.65
29	C	308	KC1	CHC-C4B-C3B	-2.93	120.26	125.21
29	H	312	KC1	C2D-C1D-ND	-2.93	107.28	110.33
21	m	101	BCR	C28-C27-C26	-2.93	108.83	114.06
18	a	824	CLA	O2D-CGD-O1D	-2.93	118.14	123.85
28	E	304	DD6	C34-C35-C36	-2.93	104.88	112.18
18	D	211	CLA	O2D-CGD-O1D	-2.93	118.14	123.85
26	H	301	A86	C19-C18-C17	-2.93	105.31	110.79
28	E	307	DD6	C37-C36-C35	2.93	119.81	114.42
18	a	853	CLA	C3B-C4B-NB	-2.93	107.92	110.53
18	B	311	CLA	O2D-CGD-O1D	-2.93	118.15	123.85
18	B	312	CLA	O2D-CGD-O1D	-2.93	118.15	123.85
18	b	845	CLA	O2D-CGD-O1D	-2.93	118.15	123.85
25	b	834	DGD	O1G-C1A-C2A	2.93	120.76	111.83
21	j	101	BCR	C3-C4-C5	-2.92	108.84	114.06
18	E	316	CLA	O2D-CGD-O1D	-2.92	118.16	123.85
18	a	807	CLA	C1-C2-C3	-2.92	121.41	126.20
18	H	305	CLA	O2D-CGD-O1D	-2.92	118.16	123.85
18	a	823	CLA	O2D-CGD-O1D	-2.92	118.16	123.85

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
18	b	801	CLA	O2A-CGA-O1A	-2.92	116.33	123.63
18	C	311	CLA	CAA-CBA-CGA	-2.92	104.92	113.21
18	a	847	CLA	O2D-CGD-O1D	-2.92	118.17	123.85
18	a	848	CLA	O2D-CGD-O1D	-2.92	118.17	123.85
20	a	832	LHG	O8-C23-C24	2.92	120.00	111.15
18	b	805	CLA	CHB-C4A-NA	2.92	128.61	124.40
18	l	205	CLA	O2D-CGD-O1D	-2.91	118.17	123.85
26	C	305	A86	C19-C18-C17	-2.91	105.34	110.79
21	a	845	BCR	C3-C4-C5	-2.91	108.87	114.06
18	b	846	CLA	O2D-CGD-O1D	-2.91	118.19	123.85
18	a	822	CLA	O2D-CGD-O1D	-2.91	118.19	123.85
21	a	833	BCR	C20-C21-C22	-2.91	123.20	127.28
18	b	828	CLA	O2D-CGD-O1D	-2.91	118.19	123.85
18	b	826	CLA	CMB-C2B-C1B	-2.91	120.99	125.42
18	b	820	CLA	C3B-C4B-NB	-2.91	107.94	110.53
18	a	814	CLA	O2D-CGD-O1D	-2.91	118.19	123.85
26	C	305	A86	O-C13-C11	-2.91	114.78	121.04
28	D	205	DD6	C25-C24-C1	2.90	134.33	126.36
18	C	309	CLA	C3B-C4B-NB	-2.90	107.94	110.53
18	D	217	CLA	C3B-C4B-NB	-2.90	107.94	110.53
21	m	101	BCR	C8-C9-C10	2.90	123.57	119.01
18	a	842	CLA	O2A-CGA-O1A	-2.90	116.37	123.63
18	D	212	CLA	C3B-C4B-NB	-2.90	107.94	110.53
26	H	302	A86	C12-C11-C10	-2.90	116.62	123.67
18	B	311	CLA	C3B-C4B-NB	-2.90	107.94	110.53
21	E	305	BCR	C24-C23-C22	-2.90	121.95	126.23
18	D	216	CLA	C3B-C4B-NB	-2.90	107.94	110.53
18	E	309	CLA	O2D-CGD-O1D	-2.90	118.21	123.85
18	b	815	CLA	O2D-CGD-O1D	-2.90	118.21	123.85
18	j	104	CLA	O2D-CGD-O1D	-2.89	118.22	123.85
18	E	313	CLA	O2D-CGD-O1D	-2.89	118.22	123.85
18	a	841	CLA	O2D-CGD-CBD	2.89	116.29	111.23
18	a	812	CLA	O2D-CGD-O1D	-2.89	118.22	123.85
18	b	825	CLA	O2D-CGD-O1D	-2.89	118.22	123.85
18	H	309	CLA	O2D-CGD-O1D	-2.89	118.22	123.85
18	a	811	CLA	O2D-CGD-O1D	-2.89	118.23	123.85
18	a	810	CLA	O2D-CGD-O1D	-2.88	118.23	123.85
18	D	215	CLA	O2D-CGD-O1D	-2.88	118.23	123.85
21	E	305	BCR	C16-C17-C18	-2.88	123.23	127.28
18	D	210	CLA	C3B-C4B-NB	-2.88	107.96	110.53
18	b	806	CLA	O2D-CGD-O1D	-2.88	118.24	123.85
18	a	812	CLA	C3B-C4B-NB	-2.88	107.96	110.53

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
18	a	846	CLA	O2D-CGD-O1D	-2.88	118.24	123.85
18	a	805	CLA	O2D-CGD-O1D	-2.88	118.24	123.85
18	b	849	CLA	O2D-CGD-O1D	-2.88	118.24	123.85
18	C	306	CLA	O2D-CGD-O1D	-2.88	118.25	123.85
28	C	303	DD6	C10-C9-C8	-2.88	114.86	123.20
29	H	312	KC1	O1D-CGD-CBD	-2.88	118.85	124.52
18	a	802	CLA	O2D-CGD-O1D	-2.87	118.25	123.85
26	B	302	A86	C12-C11-C10	-2.87	116.69	123.67
21	l	204	BCR	C34-C9-C10	-2.87	118.17	122.82
18	D	215	CLA	CAA-C2A-C3A	-2.87	109.65	116.23
28	E	307	DD6	O1-C20-C19	-2.87	110.80	113.49
21	l	201	BCR	C15-C16-C17	-2.86	117.66	123.52
18	a	802	CLA	C3B-C4B-NB	-2.86	107.97	110.53
18	a	804	CLA	C3B-C4B-NB	-2.86	107.97	110.53
18	f	202	CLA	O2D-CGD-O1D	-2.86	118.28	123.85
21	a	833	BCR	C11-C10-C9	-2.86	123.26	127.28
18	a	819	CLA	C3B-C4B-NB	-2.86	107.97	110.53
18	E	315	CLA	C3B-C4B-NB	-2.86	107.97	110.53
21	l	204	BCR	C38-C26-C25	-2.86	121.36	124.48
18	D	212	CLA	O2D-CGD-O1D	-2.86	118.28	123.85
21	l	204	BCR	C20-C21-C22	-2.86	123.27	127.28
18	b	810	CLA	C3B-C4B-NB	-2.86	107.98	110.53
26	H	301	A86	C40-C32-C31	-2.86	107.91	110.47
21	a	834	BCR	C7-C8-C9	-2.86	122.01	126.23
18	b	819	CLA	O2D-CGD-O1D	-2.85	118.29	123.85
21	f	201	BCR	C33-C5-C4	2.85	119.67	113.60
18	E	315	CLA	O2D-CGD-O1D	-2.85	118.30	123.85
18	a	841	CLA	O2A-CGA-O1A	-2.85	116.51	123.63
18	D	213	CLA	O2D-CGD-O1D	-2.84	118.31	123.85
21	b	836	BCR	C24-C23-C22	-2.84	122.03	126.23
21	b	839	BCR	C34-C9-C10	-2.84	118.21	122.82
18	a	839	CLA	CHB-C4A-NA	2.84	128.50	124.40
18	H	314	CLA	C3B-C4B-NB	-2.84	107.99	110.53
18	b	808	CLA	O2D-CGD-O1D	-2.84	118.32	123.85
18	a	805	CLA	CHB-C4A-NA	2.84	128.50	124.40
28	H	303	DD6	C15-C14-C13	-2.84	119.99	125.99
29	C	308	KC1	C2D-C1D-ND	-2.84	107.38	110.33
26	r	203	A86	C7-C6-C8	2.84	122.42	118.09
18	b	814	CLA	O2D-CGD-O1D	-2.84	118.33	123.85
26	C	302	A86	O4-C38-O5	-2.83	117.53	122.99
18	b	817	CLA	O2D-CGD-O1D	-2.83	118.34	123.85
18	a	817	CLA	O2D-CGD-O1D	-2.83	118.35	123.85

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
18	E	310	CLA	C3B-C4B-NB	-2.83	108.01	110.53
21	a	845	BCR	C30-C25-C26	-2.83	118.77	122.64
26	B	301	A86	C3-C4-C5	-2.83	117.74	123.52
28	H	303	DD6	O1-C20-C19	-2.83	110.84	113.49
18	E	309	CLA	C3B-C4B-NB	-2.83	108.01	110.53
18	H	308	CLA	C1-C2-C3	-2.82	121.57	126.20
28	B	303	DD6	C4-C3-C2	-2.82	117.74	123.52
18	a	801	CLA	C4D-C3D-CAD	-2.82	105.04	108.11
18	a	808	CLA	O2D-CGD-O1D	-2.82	118.36	123.85
26	B	304	A86	O-C13-C11	-2.82	114.97	121.04
18	a	847	CLA	C3B-C4B-NB	-2.82	108.01	110.53
18	H	314	CLA	C1-C2-C3	-2.82	121.58	126.20
18	B	308	CLA	O2D-CGD-O1D	-2.82	118.36	123.85
23	D	202	LMG	O2-C2-C3	-2.82	103.74	110.38
18	a	855	CLA	C3B-C4B-NB	-2.82	108.02	110.53
26	B	305	A86	C19-C18-C17	-2.82	105.52	110.79
21	r	201	BCR	C38-C26-C25	-2.81	121.42	124.48
26	r	203	A86	C21-C20-C15	2.81	132.42	123.35
18	b	817	CLA	C3B-C4B-NB	-2.81	108.02	110.53
21	l	204	BCR	C38-C26-C27	2.81	119.58	113.60
18	a	810	CLA	C3B-C4B-NB	-2.80	108.03	110.53
18	b	838	CLA	CHB-C4A-NA	2.80	128.45	124.40
21	E	305	BCR	C38-C26-C25	-2.80	121.42	124.48
18	a	827	CLA	O2D-CGD-O1D	-2.80	118.39	123.85
26	B	304	A86	C12-C11-C10	-2.80	116.86	123.67
18	a	822	CLA	C3B-C4B-NB	-2.80	108.03	110.53
18	C	310	CLA	O2D-CGD-O1D	-2.80	118.40	123.85
18	b	846	CLA	C3B-C4B-NB	-2.80	108.03	110.53
29	C	308	KC1	C4B-CHC-C1C	-2.80	120.07	126.02
18	E	311	CLA	O2D-CGD-O1D	-2.80	118.40	123.85
18	f	203	CLA	O2D-CGD-O1D	-2.80	118.41	123.85
18	C	309	CLA	CHB-C4A-NA	2.79	128.43	124.40
21	E	305	BCR	C21-C20-C19	-2.79	115.11	123.20
18	E	310	CLA	O2D-CGD-O1D	-2.79	118.41	123.85
18	E	313	CLA	C3B-C4B-NB	-2.79	108.04	110.53
18	i	101	CLA	O2D-CGD-O1D	-2.79	118.42	123.85
18	b	837	CLA	O2A-CGA-O1A	-2.79	116.65	123.63
21	a	852	BCR	C20-C19-C18	-2.78	118.73	126.36
18	D	211	CLA	C3B-C4B-NB	-2.78	108.05	110.53
21	f	201	BCR	C7-C8-C9	-2.78	122.13	126.23
26	B	305	A86	O-C13-C14	-2.77	116.07	121.76
18	b	821	CLA	O2D-CGD-O1D	-2.77	118.45	123.85

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
18	a	801	CLA	C3B-C4B-NB	-2.77	108.05	110.53
26	H	302	A86	C7-C6-C8	2.77	122.33	118.09
26	B	302	A86	C7-C6-C8	2.77	122.32	118.09
21	b	833	BCR	C11-C12-C13	-2.77	118.78	126.36
18	b	848	CLA	C3B-C4B-NB	-2.77	108.06	110.53
18	C	306	CLA	C3B-C4B-NB	-2.77	108.06	110.53
18	f	204	CLA	C3B-C4B-NB	-2.77	108.06	110.53
29	C	313	KC1	CHB-C1B-C2B	-2.77	119.74	125.49
21	b	836	BCR	C20-C19-C18	-2.76	118.80	126.36
26	C	305	A86	C40-C32-C31	-2.76	108.00	110.47
20	j	102	LHG	O8-C23-C24	2.76	120.25	111.83
26	m	102	A86	C3-C4-C5	-2.76	117.88	123.52
18	f	202	CLA	C3B-C4B-NB	-2.76	108.07	110.53
28	D	205	DD6	C13-C11-C10	2.76	123.35	119.01
21	a	833	BCR	C38-C26-C27	2.76	119.47	113.60
23	a	851	LMG	O8-C28-C29	2.76	120.24	111.83
18	H	315	CLA	C3B-C4B-NB	-2.76	108.07	110.53
18	b	811	CLA	O2A-CGA-O1A	-2.76	116.74	123.63
18	C	318	CLA	CHB-C4A-NA	2.75	128.38	124.40
23	C	319	LMG	C1-O6-C5	-2.75	108.34	113.72
26	H	304	A86	C8-C6-C5	-2.75	114.68	119.01
18	r	202	CLA	C3B-C4B-NB	-2.75	108.07	110.53
18	a	855	CLA	CMB-C2B-C1B	-2.75	121.23	125.42
21	l	201	BCR	C34-C9-C10	-2.75	118.36	122.82
18	a	839	CLA	C3B-C4B-NB	-2.75	108.08	110.53
18	r	202	CLA	O2D-CGD-O1D	-2.75	118.50	123.85
18	B	314	CLA	CMA-C3A-C2A	-2.75	109.93	116.23
21	l	201	BCR	C21-C20-C19	-2.75	115.24	123.20
18	C	315	CLA	O2D-CGD-O1D	-2.74	118.51	123.85
18	a	826	CLA	CHB-C4A-NA	2.74	128.36	124.40
28	B	303	DD6	C37-C36-C31	-2.74	119.03	124.16
26	b	847	A86	C-C1-C24	2.74	122.28	118.09
29	B	313	KC1	C1D-CHD-C4C	-2.74	119.80	126.25
26	H	304	A86	O1-C15-C20	-2.74	57.13	59.45
18	b	844	CLA	C3B-C4B-NB	-2.74	108.09	110.53
21	f	201	BCR	C20-C19-C18	-2.74	118.86	126.36
28	E	308	DD6	C14-C13-C11	-2.73	121.29	125.53
18	E	313	CLA	CHB-C4A-NA	2.73	128.34	124.40
21	b	839	BCR	C15-C14-C13	-2.73	123.45	127.28
21	a	834	BCR	C28-C27-C26	-2.73	109.19	114.06
18	C	309	CLA	O2A-CGA-O1A	-2.73	116.80	123.63
26	r	203	A86	O1-C15-C20	-2.73	57.14	59.45

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
18	a	814	CLA	C3B-C4B-NB	-2.72	108.10	110.53
26	B	304	A86	O-C13-C14	-2.72	116.18	121.76
29	C	313	KC1	CHC-C4B-C3B	-2.72	120.62	125.21
23	C	301	LMG	C4-C3-C2	2.72	115.61	110.83
21	b	839	BCR	C21-C20-C19	-2.72	115.31	123.20
20	a	835	LHG	O8-C23-C24	2.72	120.12	111.83
26	C	305	A86	C8-C6-C5	-2.72	114.74	119.01
18	b	829	CLA	C1-C2-C3	-2.71	121.75	126.20
18	a	809	CLA	C3B-C4B-NB	-2.71	108.11	110.53
18	D	209	CLA	O2D-CGD-O1D	-2.71	118.57	123.85
18	a	816	CLA	C3B-C4B-NB	-2.71	108.11	110.53
18	b	825	CLA	C3B-C4B-NB	-2.70	108.12	110.53
28	B	303	DD6	C28-C27-C26	-2.70	118.93	124.18
21	a	833	BCR	C8-C7-C6	-2.70	119.78	127.00
21	l	204	BCR	C27-C26-C25	-2.70	119.06	122.70
20	b	835	LHG	O8-C23-C24	2.70	120.07	111.83
26	E	302	A86	C41-C32-C31	-2.70	108.06	110.47
21	i	102	BCR	C33-C5-C6	-2.70	121.54	124.48
18	l	205	CLA	C3B-C4B-NB	-2.70	108.12	110.53
18	a	854	CLA	O2D-CGD-O1D	-2.69	118.60	123.85
18	a	822	CLA	CHB-C4A-NA	2.69	128.29	124.40
26	B	302	A86	O1-C15-C20	-2.69	57.17	59.45
26	C	302	A86	C41-C32-C31	-2.69	108.06	110.47
18	a	811	CLA	CHB-C4A-NA	2.69	128.28	124.40
18	D	214	CLA	CHB-C4A-NA	2.69	128.28	124.40
29	H	312	KC1	C1D-CHD-C4C	-2.69	119.92	126.25
28	C	303	DD6	C28-C27-C26	-2.69	118.96	124.18
26	D	206	A86	O1-C15-C20	-2.69	57.17	59.45
18	f	204	CLA	O2D-CGD-O1D	-2.69	118.62	123.85
21	r	201	BCR	C33-C5-C6	-2.69	121.55	124.48
18	b	850	CLA	C3B-C4B-NB	-2.68	108.14	110.53
26	E	302	A86	O-C13-C11	-2.68	115.27	121.04
27	l	206	ET4	C34-C35-C36	-2.68	107.72	113.59
28	H	303	DD6	O1-C15-C14	-2.68	109.20	116.88
18	b	822	CLA	CHB-C4A-NA	2.68	128.26	124.40
26	C	302	A86	O1-C15-C20	-2.68	57.18	59.45
28	B	303	DD6	C-C1-C24	-2.68	114.00	118.09
21	a	845	BCR	C23-C24-C25	-2.68	119.84	127.00
18	b	824	CLA	O2D-CGD-O1D	-2.68	118.64	123.85
21	a	852	BCR	C3-C4-C5	-2.68	109.28	114.06
28	H	303	DD6	C4-C5-C6	-2.67	123.53	127.28
18	a	806	CLA	C3B-C4B-NB	-2.67	108.14	110.53

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
18	a	808	CLA	C3B-C4B-NB	-2.67	108.14	110.53
18	E	311	CLA	CHB-C4A-NA	2.67	128.25	124.40
18	a	821	CLA	CHB-C4A-NA	2.67	128.25	124.40
28	D	205	DD6	C-C1-C2	-2.67	118.49	122.82
21	a	850	BCR	C30-C25-C26	-2.67	118.99	122.64
21	E	305	BCR	C20-C21-C22	-2.67	123.54	127.28
18	b	826	CLA	CHB-C4A-NA	2.67	128.25	124.40
21	r	201	BCR	C11-C10-C9	-2.67	123.54	127.28
18	C	310	CLA	C3B-C4B-NB	-2.66	108.15	110.53
28	E	307	DD6	C-C1-C2	-2.66	118.50	122.82
18	E	311	CLA	C3B-C4B-NB	-2.66	108.15	110.53
18	a	819	CLA	CHB-C4A-NA	2.66	128.24	124.40
26	C	305	A86	O4-C38-O5	-2.66	117.85	122.99
18	C	315	CLA	C3B-C4B-NB	-2.66	108.15	110.53
18	b	805	CLA	C3B-C4B-NB	-2.66	108.16	110.53
26	E	302	A86	C3-C2-C1	2.66	131.01	127.28
18	a	841	CLA	C3B-C4B-NB	-2.66	108.16	110.53
28	H	303	DD6	C23-C16-C22	2.66	111.23	107.37
29	C	313	KC1	O1D-CGD-CBD	-2.66	119.28	124.52
18	B	312	CLA	CHB-C4A-NA	2.66	128.23	124.40
21	E	305	BCR	C15-C16-C17	-2.66	118.08	123.52
18	b	814	CLA	CHB-C4A-NA	2.65	128.23	124.40
21	f	201	BCR	C16-C15-C14	-2.65	118.09	123.52
26	H	302	A86	O4-C38-O5	-2.65	117.88	122.99
26	B	302	A86	O4-C38-O5	-2.65	117.88	122.99
28	D	205	DD6	C41-C32-C31	-2.65	105.77	110.52
26	D	203	A86	C25-C24-C1	2.65	133.62	126.36
18	b	818	CLA	CHB-C4A-NA	2.65	128.22	124.40
18	B	308	CLA	CHB-C4A-NA	2.65	128.22	124.40
18	H	307	CLA	CAA-CBA-CGA	-2.64	105.70	113.21
18	b	821	CLA	C3B-C4B-NB	-2.64	108.17	110.53
21	i	102	BCR	C3-C4-C5	-2.64	109.34	114.06
26	C	305	A86	O1-C15-C20	-2.64	57.21	59.45
18	b	844	CLA	CHB-C4A-NA	2.64	128.21	124.40
26	D	203	A86	C19-C18-C17	-2.64	105.85	110.79
21	f	201	BCR	C10-C11-C12	-2.64	115.55	123.20
18	b	845	CLA	C3B-C4B-NB	-2.64	108.17	110.53
29	C	308	KC1	CHB-C1B-C2B	-2.64	120.00	125.49
26	C	302	A86	C4-C5-C6	2.64	130.98	127.28
18	a	810	CLA	CHB-C4A-NA	2.64	128.21	124.40
29	H	312	KC1	CHC-C4B-C3B	-2.64	120.76	125.21
21	r	201	BCR	C7-C8-C9	-2.64	122.33	126.23

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
18	a	817	CLA	C3B-C4B-NB	-2.64	108.18	110.53
18	H	309	CLA	C3B-C4B-NB	-2.63	108.18	110.53
18	a	844	CLA	C3B-C4B-NB	-2.63	108.18	110.53
22	a	849	SQD	O48-C23-C24	2.63	119.85	111.83
26	b	847	A86	C24-C1-C2	-2.63	114.88	119.01
21	a	833	BCR	C10-C11-C12	-2.63	115.59	123.20
28	E	308	DD6	O1-C20-C19	-2.62	111.03	113.49
28	E	308	DD6	C15-C14-C13	-2.62	120.45	125.99
21	l	201	BCR	C8-C9-C10	2.62	123.13	119.01
18	f	203	CLA	CHB-C4A-NA	2.62	128.18	124.40
18	b	813	CLA	O2D-CGD-O1D	-2.62	118.75	123.85
18	a	827	CLA	C3B-C4B-NB	-2.62	108.19	110.53
18	f	204	CLA	CHB-C4A-NA	2.62	128.18	124.40
27	l	206	ET4	C35-C34-C33	2.62	114.65	109.49
18	H	307	CLA	C1-C2-C3	-2.62	121.91	126.20
18	b	849	CLA	CHB-C4A-NA	2.62	128.18	124.40
18	b	840	CLA	O2A-CGA-O1A	-2.62	117.09	123.63
18	a	843	CLA	CHB-C4A-NA	2.61	128.17	124.40
29	C	313	KC1	C1D-CHD-C4C	-2.61	120.10	126.25
18	b	823	CLA	C3B-C4B-NB	-2.61	108.20	110.53
21	b	832	BCR	C38-C26-C25	-2.61	121.63	124.48
18	D	209	CLA	CHB-C4A-NA	2.61	128.17	124.40
18	f	203	CLA	C3B-C4B-NB	-2.61	108.20	110.53
18	a	824	CLA	C3B-C4B-NB	-2.61	108.20	110.53
26	D	204	A86	O4-C38-O5	-2.61	117.96	122.99
18	a	825	CLA	O1D-CGD-CBD	2.61	129.66	124.52
18	a	824	CLA	CHB-C4A-NA	2.60	128.16	124.40
18	C	316	CLA	CHB-C4A-NA	2.60	128.16	124.40
18	a	837	CLA	C1-C2-C3	-2.60	121.93	126.20
21	b	839	BCR	C33-C5-C4	2.60	119.15	113.60
18	b	808	CLA	C3B-C4B-NB	-2.60	108.21	110.53
18	b	804	CLA	CHB-C4A-NA	2.60	128.15	124.40
28	C	303	DD6	C33-C32-C31	-2.60	104.37	109.49
18	a	838	CLA	O2D-CGD-O1D	-2.60	118.79	123.85
18	H	310	CLA	CHB-C4A-NA	2.60	128.15	124.40
18	b	810	CLA	CHB-C4A-NA	2.60	128.15	124.40
21	a	845	BCR	C21-C20-C19	-2.59	115.69	123.20
21	i	102	BCR	C30-C25-C26	-2.59	119.09	122.64
18	a	843	CLA	CMB-C2B-C1B	-2.59	121.48	125.42
26	H	302	A86	O1-C15-C20	-2.59	57.26	59.45
18	C	312	CLA	C1-C2-C3	-2.59	121.96	126.20
18	b	809	CLA	CHB-C4A-NA	2.59	128.13	124.40

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
21	f	201	BCR	C38-C26-C25	-2.59	121.66	124.48
18	E	309	CLA	CHB-C4A-NA	2.58	128.13	124.40
18	b	830	CLA	C3B-C4B-NB	-2.58	108.23	110.53
21	l	201	BCR	C10-C11-C12	-2.58	115.72	123.20
18	b	812	CLA	CHB-C4A-NA	2.58	128.12	124.40
18	a	818	CLA	CHB-C4A-NA	2.58	128.12	124.40
18	B	309	CLA	CHB-C4A-NA	2.58	128.12	124.40
18	C	309	CLA	C1-C2-C3	-2.58	121.97	126.20
18	j	104	CLA	CHB-C4A-NA	2.58	128.12	124.40
18	b	842	CLA	C3B-C4B-NB	-2.58	108.23	110.53
18	a	846	CLA	C3B-C4B-NB	-2.58	108.23	110.53
26	H	301	A86	C34-O4-C38	-2.58	113.30	117.85
18	b	819	CLA	C3B-C4B-NB	-2.57	108.23	110.53
26	C	302	A86	C21-C20-C19	-2.57	111.35	114.24
18	a	816	CLA	CHB-C4A-NA	2.57	128.11	124.40
29	B	313	KC1	O2D-CGD-O1D	-2.57	118.84	123.85
21	b	833	BCR	C8-C7-C6	-2.57	120.13	127.00
18	b	803	CLA	C1-C2-C3	-2.57	121.99	126.20
26	C	304	A86	O1-C15-C20	-2.57	57.27	59.45
19	b	831	PQN	C14-C13-C15	2.57	119.69	115.23
18	b	806	CLA	C3B-C4B-NB	-2.57	108.24	110.53
18	D	208	CLA	O2D-CGD-CBD	2.57	115.72	111.23
18	D	211	CLA	CHB-C4A-NA	2.57	128.10	124.40
18	a	826	CLA	C3B-C4B-NB	-2.56	108.24	110.53
28	D	205	DD6	C4-C5-C6	2.56	130.87	127.28
26	B	305	A86	C29-C30-C31	-2.56	174.64	177.66
18	a	827	CLA	CHB-C4A-NA	2.56	128.10	124.40
18	a	855	CLA	CHB-C4A-NA	2.56	128.10	124.40
18	b	806	CLA	CHB-C4A-NA	2.56	128.10	124.40
28	H	303	DD6	C33-C32-C31	2.56	114.53	109.49
26	C	305	A86	C4-C5-C6	2.56	130.87	127.28
18	b	811	CLA	CHB-C4A-NA	2.56	128.09	124.40
26	B	301	A86	C28-C27-C26	-2.56	118.67	122.82
18	b	822	CLA	CMB-C2B-C1B	-2.56	121.53	125.42
18	D	213	CLA	CHB-C4A-NA	2.55	128.09	124.40
26	B	305	A86	C8-C6-C5	-2.55	114.99	119.01
18	E	312	CLA	CAA-CBA-CGA	-2.55	105.96	113.21
18	b	848	CLA	CHB-C4A-NA	2.55	128.08	124.40
18	a	803	CLA	CHB-C4A-NA	2.55	128.08	124.40
18	a	813	CLA	O2D-CGD-CBD	2.55	115.68	111.23
21	E	305	BCR	C8-C7-C6	-2.55	120.19	127.00
18	E	314	CLA	O2D-CGD-CBD	2.55	115.68	111.23

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
18	a	807	CLA	CBA-CAA-C2A	-2.55	106.22	113.79
18	l	203	CLA	O2A-CGA-O1A	-2.54	117.26	123.63
26	b	847	A86	C7-C6-C5	-2.54	118.70	122.82
26	D	203	A86	C28-C27-C26	-2.54	118.70	122.82
21	E	306	BCR	C33-C5-C4	2.54	119.02	113.60
18	C	306	CLA	CHB-C4A-NA	2.54	128.07	124.40
18	B	307	CLA	CHB-C4A-NA	2.54	128.06	124.40
18	a	814	CLA	CHB-C4A-NA	2.54	128.06	124.40
18	a	825	CLA	O2A-CGA-O1A	-2.54	117.28	123.63
28	D	205	DD6	C14-C13-C11	-2.54	121.59	125.53
21	b	836	BCR	C30-C25-C26	-2.54	119.17	122.64
26	C	304	A86	C3-C4-C5	-2.54	118.33	123.52
28	E	308	DD6	O1-C20-C21	-2.53	112.22	115.05
21	b	836	BCR	C27-C26-C25	-2.53	119.28	122.70
18	E	316	CLA	CMB-C2B-C1B	-2.53	121.56	125.42
26	r	203	A86	C21-C20-C19	-2.53	111.40	114.24
18	a	812	CLA	CHB-C4A-NA	2.53	128.06	124.40
18	b	842	CLA	CHB-C4A-NA	2.53	128.06	124.40
18	b	840	CLA	CHB-C4A-NA	2.53	128.05	124.40
18	H	314	CLA	CMB-C2B-C1B	-2.53	121.57	125.42
18	H	305	CLA	CHB-C4A-NA	2.53	128.05	124.40
26	B	305	A86	O1-C15-C20	-2.53	57.31	59.45
26	H	304	A86	C24-C1-C2	-2.53	115.03	119.01
18	l	202	CLA	O2A-CGA-O1A	-2.53	117.31	123.63
18	a	854	CLA	CHB-C4A-NA	2.53	128.04	124.40
26	H	302	A86	C40-C32-C31	-2.52	108.21	110.47
18	a	847	CLA	CHB-C4A-NA	2.52	128.04	124.40
18	a	827	CLA	O2A-CGA-O1A	-2.52	117.32	123.63
18	H	315	CLA	CHB-C4A-NA	2.52	128.04	124.40
26	E	302	A86	C12-C11-C10	-2.52	117.54	123.67
18	D	217	CLA	CMA-C3A-C2A	-2.52	110.45	116.23
18	a	815	CLA	C3B-C4B-NB	-2.52	108.28	110.53
18	b	813	CLA	C3B-C4B-NB	-2.52	108.28	110.53
18	D	209	CLA	C3B-C4B-NB	-2.52	108.28	110.53
21	a	852	BCR	C38-C26-C27	2.52	118.97	113.60
18	C	317	CLA	O2D-CGD-O1D	-2.52	118.94	123.85
26	D	206	A86	C19-C18-C17	-2.52	106.08	110.79
18	b	804	CLA	C3B-C4B-NB	-2.52	108.28	110.53
18	b	814	CLA	CMB-C2B-C1B	-2.52	121.58	125.42
26	B	302	A86	C40-C32-C31	-2.52	108.22	110.47
28	E	304	DD6	C4-C3-C2	-2.52	118.37	123.52
26	E	302	A86	C36-C31-C32	2.52	122.19	119.70

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
18	b	818	CLA	C3B-C4B-NB	-2.51	108.28	110.53
18	a	804	CLA	CHB-C4A-NA	2.51	128.03	124.40
19	b	831	PQN	C11-C3-C2	-2.51	120.58	124.89
19	b	831	PQN	C2M-C2-C3	-2.51	120.32	124.45
26	C	302	A86	C21-C20-C15	2.51	131.45	123.35
18	B	309	CLA	C3B-C4B-NB	-2.51	108.29	110.53
28	B	303	DD6	C9-C10-C11	-2.51	123.76	127.28
18	a	844	CLA	CHB-C4A-NA	2.51	128.02	124.40
18	H	313	CLA	CHB-C4A-NA	2.51	128.02	124.40
21	a	850	BCR	C7-C8-C9	-2.51	122.53	126.23
21	b	839	BCR	C30-C25-C24	2.51	122.45	115.65
26	D	203	A86	O4-C38-O5	-2.51	118.15	122.99
21	i	102	BCR	C1-C6-C5	-2.50	119.22	122.64
18	a	820	CLA	C3B-C4B-NB	-2.50	108.30	110.53
18	a	848	CLA	C3B-C4B-NB	-2.50	108.30	110.53
18	a	853	CLA	CHB-C4A-NA	2.50	128.01	124.40
18	b	843	CLA	CHB-C4A-NA	2.50	128.01	124.40
18	H	309	CLA	CHB-C4A-NA	2.50	128.01	124.40
26	H	301	A86	O1-C15-C20	-2.50	57.33	59.45
18	b	826	CLA	CMB-C2B-C3B	2.50	132.42	126.55
21	i	102	BCR	C27-C26-C25	-2.50	119.33	122.70
18	b	825	CLA	CHB-C4A-NA	2.50	128.00	124.40
26	B	305	A86	C9-C8-C6	2.49	133.20	126.36
28	E	308	DD6	O1-C15-C14	-2.49	109.73	116.88
18	l	205	CLA	CHB-C4A-NA	2.49	128.00	124.40
28	E	304	DD6	C37-C36-C31	-2.49	119.50	124.16
18	a	808	CLA	CHB-C4A-NA	2.49	128.00	124.40
29	C	308	KC1	C1D-CHD-C4C	-2.49	120.38	126.25
18	E	315	CLA	CHB-C4A-NA	2.49	127.99	124.40
26	C	304	A86	C26-C25-C24	-2.49	115.98	123.20
23	E	318	LMG	C8-O7-C10	-2.49	111.84	117.80
23	l	207	LMG	O2-C2-C3	-2.49	104.51	110.38
18	b	837	CLA	CHB-C4A-NA	2.49	127.99	124.40
18	b	838	CLA	C3B-C4B-NB	-2.49	108.31	110.53
18	b	840	CLA	C3B-C4B-NB	-2.49	108.31	110.53
21	E	305	BCR	C33-C5-C4	2.49	118.90	113.60
18	b	850	CLA	CHB-C4A-NA	2.49	127.99	124.40
26	m	102	A86	C12-C11-C10	-2.49	117.62	123.67
18	b	801	CLA	C3B-C4B-NB	-2.49	108.31	110.53
18	b	803	CLA	O2D-CGD-O1D	-2.48	119.01	123.85
26	H	301	A86	C26-C25-C24	-2.48	116.00	123.20
21	i	102	BCR	C33-C5-C4	2.48	118.89	113.60

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
18	B	311	CLA	CHB-C4A-NA	2.48	127.98	124.40
26	C	304	A86	C10-C9-C8	-2.48	116.02	123.20
18	b	820	CLA	CHB-C4A-NA	2.48	127.97	124.40
18	a	828	CLA	C3B-C4B-NB	-2.48	108.32	110.53
18	f	202	CLA	CHB-C4A-NA	2.48	127.97	124.40
18	H	310	CLA	C3B-C4B-NB	-2.47	108.32	110.53
26	m	102	A86	O1-C15-C20	-2.47	57.35	59.45
26	B	304	A86	O1-C15-C20	-2.47	57.35	59.45
29	H	312	KC1	C4B-CHC-C1C	-2.47	120.77	126.02
18	b	819	CLA	CHB-C4A-NA	2.47	127.97	124.40
26	C	305	A86	C41-C32-C31	-2.47	108.26	110.47
18	H	308	CLA	O2A-CGA-O1A	-2.47	117.45	123.63
18	B	308	CLA	C3B-C4B-NB	-2.47	108.33	110.53
21	E	306	BCR	C8-C7-C6	-2.47	120.41	127.00
26	E	302	A86	C4-C5-C6	2.47	130.74	127.28
18	a	823	CLA	CHB-C4A-NA	2.47	127.96	124.40
18	a	806	CLA	CHB-C4A-NA	2.47	127.96	124.40
18	b	812	CLA	C3B-C4B-NB	-2.46	108.33	110.53
21	b	836	BCR	C15-C14-C13	-2.46	123.82	127.28
18	a	837	CLA	CHB-C4A-NA	2.46	127.96	124.40
18	a	830	CLA	CHB-C4A-NA	2.46	127.95	124.40
18	b	815	CLA	CHB-C4A-NA	2.46	127.95	124.40
18	E	310	CLA	CHB-C4A-NA	2.46	127.95	124.40
26	H	301	A86	O4-C38-O5	-2.46	118.24	122.99
26	D	206	A86	C12-C11-C10	-2.46	117.69	123.67
18	a	815	CLA	CHB-C4A-NA	2.46	127.95	124.40
18	b	828	CLA	CHB-C4A-NA	2.46	127.95	124.40
18	b	823	CLA	C1-C2-C3	-2.46	122.17	126.20
18	b	817	CLA	CHB-C4A-NA	2.46	127.95	124.40
29	C	308	KC1	C2A-C3A-C4A	2.46	108.26	106.41
26	D	206	A86	C41-C32-C31	2.46	112.67	110.47
18	B	310	CLA	C3B-C4B-NB	-2.46	108.34	110.53
18	b	830	CLA	CHB-C4A-NA	2.46	127.94	124.40
18	a	817	CLA	CHB-C4A-NA	2.45	127.94	124.40
18	b	808	CLA	CHB-C4A-NA	2.45	127.94	124.40
18	a	836	CLA	CHB-C4A-NA	2.45	127.94	124.40
18	b	823	CLA	CHB-C4A-NA	2.45	127.94	124.40
18	H	307	CLA	CHB-C4A-NA	2.45	127.94	124.40
18	H	310	CLA	O2D-CGD-CBD	2.45	115.52	111.23
21	a	850	BCR	C37-C22-C21	-2.45	118.85	122.82
21	b	836	BCR	C38-C26-C27	2.45	118.82	113.60
21	a	833	BCR	C33-C5-C6	-2.45	121.81	124.48

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
21	b	839	BCR	C8-C7-C6	-2.45	120.46	127.00
26	B	305	A86	C12-C11-C10	-2.45	117.72	123.67
18	C	307	CLA	O2A-CGA-O1A	-2.45	117.51	123.63
18	a	838	CLA	CMB-C2B-C1B	-2.45	121.69	125.42
18	B	314	CLA	O2D-CGD-O1D	-2.45	119.08	123.85
18	a	821	CLA	C3B-C4B-NB	-2.45	108.35	110.53
18	D	214	CLA	C3B-C4B-NB	-2.45	108.35	110.53
21	b	839	BCR	C38-C26-C27	2.44	118.80	113.60
21	j	105	BCR	C20-C19-C18	-2.44	119.67	126.36
18	H	306	CLA	O2D-CGD-CBD	2.44	115.50	111.23
18	a	840	CLA	C11-C10-C8	-2.44	107.86	115.97
21	i	103	BCR	C15-C16-C17	-2.44	118.53	123.52
18	b	811	CLA	O2D-CGD-CBD	2.44	115.49	111.23
29	C	313	KC1	C4B-CHC-C1C	-2.44	120.84	126.02
18	D	210	CLA	CHB-C4A-NA	2.44	127.92	124.40
21	l	201	BCR	C11-C10-C9	-2.44	123.86	127.28
21	a	834	BCR	C10-C11-C12	-2.43	116.15	123.20
27	l	206	ET4	C15-C14-C13	2.43	130.69	127.28
18	E	312	CLA	O2D-CGD-O1D	-2.43	119.11	123.85
18	B	306	CLA	C3B-C4B-NB	-2.43	108.36	110.53
18	b	807	CLA	CHB-C4A-NA	2.43	127.91	124.40
18	E	312	CLA	CHB-C4A-NA	2.43	127.91	124.40
18	a	815	CLA	C1-C2-C3	-2.43	122.22	126.20
21	a	845	BCR	C20-C21-C22	-2.42	123.88	127.28
18	b	826	CLA	C3B-C4B-NB	-2.42	108.37	110.53
18	a	809	CLA	CHB-C4A-NA	2.42	127.90	124.40
18	b	845	CLA	CHB-C4A-NA	2.42	127.89	124.40
26	m	102	A86	C41-C32-C31	-2.42	108.31	110.47
18	a	848	CLA	CHB-C4A-NA	2.42	127.89	124.40
18	a	843	CLA	C3B-C4B-NB	-2.42	108.37	110.53
21	r	201	BCR	C2-C1-C6	2.42	113.95	110.44
18	C	315	CLA	CHB-C4A-NA	2.41	127.89	124.40
18	b	826	CLA	O2A-CGA-O1A	-2.41	117.59	123.63
18	a	825	CLA	CHB-C4A-NA	2.41	127.88	124.40
18	D	212	CLA	CHB-C4A-NA	2.41	127.88	124.40
26	C	305	A86	C36-C31-C32	-2.41	117.30	119.70
26	H	302	A86	C19-C18-C17	-2.41	106.28	110.79
18	a	855	CLA	CMB-C2B-C3B	2.41	132.22	126.55
21	E	306	BCR	C20-C19-C18	-2.41	119.76	126.36
18	H	307	CLA	C16-C15-C13	-2.41	107.96	115.97
18	H	311	CLA	O2D-CGD-CBD	2.41	115.44	111.23
18	b	820	CLA	O2A-CGA-O1A	-2.41	117.61	123.63

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
29	B	313	KC1	C4B-CHC-C1C	-2.40	120.91	126.02
18	a	846	CLA	CHB-C4A-NA	2.40	127.86	124.40
18	b	846	CLA	CHB-C4A-NA	2.40	127.86	124.40
18	C	316	CLA	O2D-CGD-O1D	-2.40	119.17	123.85
18	D	217	CLA	O2D-CGD-O1D	-2.40	119.17	123.85
18	a	813	CLA	CHB-C4A-NA	2.40	127.86	124.40
18	a	811	CLA	C3B-C4B-NB	-2.40	108.39	110.53
18	b	841	CLA	CHB-C4A-NA	2.40	127.86	124.40
26	B	301	A86	C21-C20-C15	2.40	131.09	123.35
21	b	832	BCR	C33-C5-C6	-2.40	121.87	124.48
18	b	816	CLA	CHB-C4A-NA	2.40	127.86	124.40
26	B	302	A86	C19-C18-C17	-2.40	106.31	110.79
18	b	822	CLA	C3B-C4B-NB	-2.39	108.39	110.53
26	H	304	A86	C12-C11-C10	-2.39	117.86	123.67
21	a	845	BCR	C38-C26-C27	2.39	118.70	113.60
26	E	302	A86	O1-C15-C20	-2.39	57.42	59.45
26	B	302	A86	C-C1-C24	2.39	121.74	118.09
18	a	830	CLA	C3B-C4B-NB	-2.39	108.40	110.53
21	b	836	BCR	C38-C26-C25	-2.39	121.88	124.48
18	C	311	CLA	CHB-C4A-NA	2.39	127.85	124.40
18	a	801	CLA	CBA-CAA-C2A	-2.39	106.69	113.79
21	r	201	BCR	C10-C11-C12	-2.39	116.28	123.20
26	C	305	A86	C12-C11-C10	-2.38	117.87	123.67
18	b	803	CLA	C3B-C4B-NB	-2.38	108.40	110.53
21	a	845	BCR	C10-C11-C12	-2.38	116.29	123.20
18	C	311	CLA	C1-C2-C3	-2.38	122.29	126.20
18	b	824	CLA	CHB-C4A-NA	2.38	127.84	124.40
26	m	102	A86	C19-C18-C17	-2.38	106.34	110.79
18	f	202	CLA	C1-C2-C3	-2.38	122.30	126.20
28	E	307	DD6	O1-C20-C21	-2.38	112.39	115.05
21	l	201	BCR	C23-C24-C25	-2.38	120.64	127.00
18	b	841	CLA	C3B-C4B-NB	-2.38	108.41	110.53
21	a	845	BCR	C16-C15-C14	-2.38	118.66	123.52
21	b	832	BCR	C16-C15-C14	-2.37	118.66	123.52
21	i	102	BCR	C20-C19-C18	-2.37	119.85	126.36
18	b	803	CLA	O2A-CGA-O1A	-2.37	117.69	123.63
26	D	204	A86	O-C13-C11	-2.37	115.93	121.04
21	b	836	BCR	C33-C5-C4	2.37	118.65	113.60
26	E	302	A86	O4-C38-O5	-2.37	118.41	122.99
21	i	102	BCR	C16-C15-C14	-2.37	118.67	123.52
21	j	105	BCR	C38-C26-C27	2.37	118.65	113.60
21	E	306	BCR	C16-C15-C14	-2.37	118.67	123.52

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
18	i	101	CLA	CHB-C4A-NA	2.37	127.82	124.40
26	D	204	A86	O1-C15-C20	-2.37	57.44	59.45
26	r	203	A86	O-C13-C11	-2.37	115.94	121.04
29	C	313	KC1	C2A-C3A-C4A	2.37	108.19	106.41
18	b	815	CLA	C3B-C4B-NB	-2.36	108.42	110.53
18	a	801	CLA	CHB-C4A-NA	2.36	127.81	124.40
21	i	102	BCR	C4-C5-C6	-2.36	119.51	122.70
18	a	837	CLA	O2A-CGA-O1A	-2.36	117.72	123.63
18	b	821	CLA	CHB-C4A-NA	2.36	127.81	124.40
26	H	304	A86	C-C1-C24	2.36	121.70	118.09
26	H	302	A86	C-C1-C24	2.36	121.69	118.09
26	C	305	A86	C21-C20-C19	-2.36	111.59	114.24
26	H	302	A86	O-C13-C14	-2.36	116.92	121.76
18	b	829	CLA	CAA-CBA-CGA	-2.36	106.51	113.21
18	D	215	CLA	CHB-C4A-NA	2.35	127.80	124.40
21	a	833	BCR	C27-C26-C25	-2.35	119.52	122.70
18	b	807	CLA	C3B-C4B-NB	-2.35	108.43	110.53
18	H	308	CLA	CHB-C4A-NA	2.35	127.80	124.40
18	a	823	CLA	C3B-C4B-NB	-2.35	108.43	110.53
18	a	838	CLA	CHB-C4A-NA	2.35	127.79	124.40
26	D	206	A86	O4-C38-O5	-2.35	118.45	122.99
28	D	205	DD6	C9-C10-C11	-2.35	123.98	127.28
21	m	101	BCR	C20-C19-C18	-2.35	119.92	126.36
26	C	304	A86	C24-C1-C2	2.35	122.70	119.01
23	C	301	LMG	C6-C5-C4	-2.35	107.26	113.02
26	b	847	A86	C25-C26-C27	-2.35	123.99	127.28
18	a	813	CLA	CAA-CBA-CGA	-2.34	106.55	113.21
18	a	801	CLA	O2A-CGA-O1A	-2.34	117.77	123.63
18	r	202	CLA	CHB-C4A-NA	2.34	127.78	124.40
21	m	101	BCR	C33-C5-C4	2.34	118.59	113.60
25	b	834	DGD	O2G-C1B-O1B	-2.34	118.23	123.70
18	a	820	CLA	O2D-CGD-CBD	2.34	115.32	111.23
21	a	833	BCR	C29-C30-C25	2.34	113.84	110.44
18	b	827	CLA	CHB-C4A-NA	2.34	127.77	124.40
18	a	842	CLA	CHB-C4A-NA	2.33	127.77	124.40
18	B	310	CLA	CHB-C4A-NA	2.33	127.77	124.40
18	B	306	CLA	CHB-C4A-NA	2.33	127.76	124.40
18	i	101	CLA	C11-C10-C8	-2.33	108.23	115.97
18	E	314	CLA	CHB-C4A-NA	2.33	127.76	124.40
28	E	307	DD6	C32-C33-C34	-2.33	108.49	113.59
18	b	843	CLA	CMB-C2B-C1B	-2.32	121.89	125.42
27	l	206	ET4	C42-C34-C33	-2.32	106.36	110.52

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
28	D	205	DD6	C33-C32-C31	2.32	114.06	109.49
18	a	805	CLA	CMB-C2B-C1B	-2.32	121.89	125.42
26	H	304	A86	O-C13-C11	-2.31	116.06	121.04
26	B	304	A86	C7-C6-C8	2.31	121.62	118.09
29	B	313	KC1	CHC-C4B-C3B	-2.31	121.31	125.21
21	b	832	BCR	C20-C19-C18	-2.31	120.03	126.36
21	f	205	BCR	C11-C12-C13	-2.31	120.03	126.36
26	C	302	A86	C12-C11-C10	-2.31	118.06	123.67
18	a	836	CLA	C3B-C4B-NB	-2.31	108.47	110.53
26	r	203	A86	C28-C27-C26	2.31	126.56	122.82
18	H	307	CLA	C3B-C4B-NB	-2.31	108.47	110.53
18	b	829	CLA	CHB-C4A-NA	2.31	127.73	124.40
21	a	845	BCR	C30-C25-C24	2.31	121.91	115.65
18	B	314	CLA	CAA-C2A-C3A	-2.31	110.94	116.23
29	C	308	KC1	O2D-CGD-O1D	-2.30	119.36	123.85
26	D	204	A86	C21-C20-C19	-2.30	111.65	114.24
21	a	845	BCR	C8-C7-C6	-2.30	120.84	127.00
18	a	828	CLA	CHB-C4A-NA	2.30	127.72	124.40
26	B	302	A86	O-C13-C14	-2.30	117.04	121.76
18	a	805	CLA	C3B-C4B-NB	-2.30	108.47	110.53
18	a	811	CLA	C1-C2-C3	-2.30	123.04	126.76
26	D	206	A86	C7-C6-C8	2.30	121.60	118.09
18	a	820	CLA	CHB-C4A-NA	2.30	127.72	124.40
21	b	833	BCR	C21-C20-C19	-2.30	116.55	123.20
18	D	208	CLA	C7-C6-C5	-2.30	107.14	113.26
26	H	302	A86	C8-C6-C5	-2.29	115.40	119.01
21	b	839	BCR	C23-C24-C25	-2.29	120.88	127.00
21	b	836	BCR	C23-C24-C25	-2.29	120.88	127.00
21	a	834	BCR	C8-C7-C6	-2.29	120.88	127.00
18	l	205	CLA	CMB-C2B-C1B	-2.29	121.93	125.42
21	f	205	BCR	C3-C4-C5	-2.29	109.97	114.06
18	a	854	CLA	O1D-CGD-CBD	2.29	129.03	124.52
18	b	813	CLA	CHB-C4A-NA	2.29	127.70	124.40
26	B	305	A86	O-C13-C11	-2.29	116.11	121.04
26	D	204	A86	C21-C20-C15	2.29	130.73	123.35
18	a	808	CLA	O2A-CGA-O1A	-2.28	117.93	123.63
21	j	101	BCR	C8-C7-C6	-2.28	120.92	127.00
21	b	839	BCR	C7-C8-C9	-2.28	122.87	126.23
18	H	307	CLA	O2A-CGA-O1A	-2.28	117.94	123.63
21	E	306	BCR	C28-C27-C26	-2.28	110.00	114.06
18	C	310	CLA	CHB-C4A-NA	2.27	127.68	124.40
29	C	313	KC1	O2D-CGD-O1D	-2.27	119.42	123.85

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
26	H	304	A86	C34-O4-C38	-2.27	113.83	117.85
18	E	316	CLA	CHB-C4A-NA	2.27	127.68	124.40
18	a	802	CLA	C1-C2-C3	-2.27	122.48	126.20
26	H	304	A86	C19-C18-C17	-2.27	106.55	110.79
26	m	102	A86	O-C13-C11	-2.27	116.16	121.04
18	a	802	CLA	CHB-C4A-NA	2.27	127.67	124.40
21	f	205	BCR	C15-C16-C17	-2.26	118.89	123.52
21	a	833	BCR	C21-C20-C19	-2.26	116.66	123.20
21	l	204	BCR	C8-C7-C6	-2.26	120.97	127.00
26	E	302	A86	C40-C32-C31	-2.26	108.45	110.47
18	b	840	CLA	O2D-CGD-CBD	2.26	115.17	111.23
18	D	215	CLA	C3B-C4B-NB	-2.26	108.52	110.53
21	b	832	BCR	C38-C26-C27	2.26	118.41	113.60
28	C	303	DD6	C23-C16-C22	2.26	110.65	107.37
21	l	204	BCR	C33-C5-C4	2.25	118.40	113.60
28	B	303	DD6	C10-C9-C8	-2.25	116.67	123.20
21	E	306	BCR	C38-C26-C27	2.25	118.39	113.60
18	D	207	CLA	CHB-C4A-NA	2.25	127.65	124.40
28	E	307	DD6	C7-C6-C8	2.25	121.52	118.09
21	f	205	BCR	C38-C26-C27	2.25	118.39	113.60
26	H	301	A86	C21-C20-C19	-2.25	111.72	114.24
21	a	834	BCR	C11-C10-C9	-2.25	124.13	127.28
18	a	843	CLA	CMB-C2B-C3B	2.24	131.83	126.55
18	a	803	CLA	CAA-CBA-CGA	-2.24	106.84	113.21
21	r	201	BCR	C16-C15-C14	-2.24	118.93	123.52
18	l	202	CLA	CMB-C2B-C1B	-2.24	122.01	125.42
18	a	815	CLA	CMB-C2B-C1B	-2.24	122.01	125.42
18	D	216	CLA	CHB-C4A-NA	2.24	127.63	124.40
28	E	304	DD6	O2-C18-C17	-2.24	105.17	109.75
18	a	838	CLA	CMB-C2B-C3B	2.23	131.80	126.55
18	b	823	CLA	O2A-CGA-O1A	-2.23	118.05	123.63
26	B	302	A86	C8-C6-C5	-2.23	115.50	119.01
28	E	304	DD6	C10-C9-C8	-2.23	116.75	123.20
18	C	317	CLA	CHB-C4A-NA	2.23	127.61	124.40
26	D	204	A86	C41-C32-C31	-2.23	108.48	110.47
20	B	315	LHG	O8-C23-O10	-2.23	118.06	123.63
18	C	316	CLA	CAA-C2A-C3A	-2.23	111.13	116.23
18	H	313	CLA	CBA-CAA-C2A	-2.22	107.17	113.79
18	b	822	CLA	CMB-C2B-C3B	2.22	131.78	126.55
18	a	853	CLA	C1-C2-C3	-2.22	122.55	126.20
18	C	314	CLA	CAA-C2A-C3A	-2.22	111.14	116.23
18	l	203	CLA	C11-C10-C8	-2.22	108.58	115.97

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
26	H	301	A86	C8-C6-C5	-2.22	115.52	119.01
18	b	829	CLA	O2D-CGD-CBD	2.22	115.11	111.23
26	C	304	A86	O4-C38-O5	-2.22	118.71	122.99
18	C	312	CLA	CHB-C4A-NA	2.22	127.60	124.40
21	r	201	BCR	C21-C20-C19	-2.22	116.77	123.20
18	a	839	CLA	O2A-CGA-O1A	-2.22	118.08	123.63
26	r	203	A86	C-C1-C24	-2.22	114.70	118.09
18	b	821	CLA	C1-C2-C3	-2.22	123.18	126.76
21	j	101	BCR	C33-C5-C4	2.22	118.32	113.60
18	l	202	CLA	CHB-C1B-NB	2.21	127.37	124.05
28	D	205	DD6	C37-C36-C31	-2.21	120.02	124.16
19	a	829	PQN	C11-C3-C2	-2.21	121.09	124.89
18	C	307	CLA	CHB-C4A-NA	2.21	127.59	124.40
22	a	849	SQD	O7-S-C6	2.21	110.07	106.73
18	a	818	CLA	O2D-CGD-CBD	2.21	115.10	111.23
18	H	310	CLA	CAA-CBA-CGA	-2.21	106.92	113.21
18	a	840	CLA	CHB-C4A-NA	2.21	127.59	124.40
18	D	217	CLA	CHB-C4A-NA	2.21	127.59	124.40
18	b	808	CLA	O2A-CGA-O1A	-2.21	118.10	123.63
18	b	812	CLA	O2A-CGA-O1A	-2.21	118.10	123.63
27	l	206	ET4	C26-C01-C06	-2.21	106.78	110.24
26	b	847	A86	O1-C15-C20	-2.21	57.58	59.45
21	r	201	BCR	C8-C7-C6	-2.21	121.10	127.00
21	a	833	BCR	C33-C5-C4	2.21	118.30	113.60
27	l	206	ET4	C15-C16-C17	2.21	128.03	123.52
20	b	835	LHG	O8-C6-C5	-2.20	102.04	108.40
18	C	317	CLA	O2D-CGD-CBD	2.20	115.08	111.23
18	D	208	CLA	O2A-CGA-O1A	-2.20	118.12	123.63
18	C	307	CLA	O2D-CGD-CBD	2.20	115.08	111.23
18	E	315	CLA	O2A-CGA-O1A	-2.20	118.12	123.63
26	r	203	A86	O2-C18-C19	2.20	114.25	109.75
18	l	203	CLA	CHB-C4A-NA	2.20	127.58	124.40
28	H	303	DD6	C37-C36-C35	2.20	118.47	114.42
18	E	314	CLA	CAA-CBA-CGA	-2.20	106.62	112.49
18	a	807	CLA	O2D-CGD-CBD	2.20	115.07	111.23
26	C	304	A86	C19-C18-C17	-2.20	106.68	110.79
18	b	805	CLA	C1-C2-C3	-2.19	122.60	126.20
18	b	810	CLA	O2A-CGA-O1A	-2.19	118.14	123.63
28	C	303	DD6	O1-C15-C14	-2.19	110.59	116.88
18	b	804	CLA	O2D-CGD-CBD	2.19	115.06	111.23
21	r	201	BCR	C33-C5-C4	2.19	118.27	113.60
18	b	845	CLA	O2A-CGA-O1A	-2.19	118.15	123.63

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
18	D	207	CLA	O2A-CGA-O1A	-2.19	118.15	123.63
18	f	202	CLA	CMB-C2B-C1B	-2.19	122.09	125.42
18	C	311	CLA	O2D-CGD-O1D	-2.19	119.59	123.85
26	B	301	A86	O4-C38-O5	-2.19	118.77	122.99
20	a	831	LHG	C26-C25-C24	2.19	121.17	113.13
21	a	834	BCR	C38-C26-C27	2.19	118.26	113.60
18	a	846	CLA	O2A-CGA-O1A	-2.19	118.16	123.63
18	b	804	CLA	CAA-CBA-CGA	-2.19	106.66	112.49
18	b	841	CLA	CMB-C2B-C1B	-2.18	122.10	125.42
18	a	828	CLA	C1-C2-C3	-2.18	122.62	126.20
18	C	307	CLA	CMB-C2B-C1B	-2.18	122.10	125.42
18	E	309	CLA	CMB-C2B-C1B	-2.18	122.10	125.42
18	a	840	CLA	O2D-CGD-CBD	2.18	115.04	111.23
21	a	852	BCR	C12-C13-C14	2.18	122.43	119.01
18	a	853	CLA	O2A-CGA-O1A	-2.18	118.18	123.63
26	D	203	A86	C3-C4-C5	-2.18	119.07	123.52
18	a	836	CLA	O2A-CGA-O1A	-2.18	118.19	123.63
18	b	816	CLA	O2A-CGA-O1A	-2.18	118.19	123.63
18	E	309	CLA	O2A-CGA-O1A	-2.17	118.19	123.63
28	E	307	DD6	C15-C14-C13	2.17	130.59	125.99
18	a	841	CLA	C1-C2-C3	-2.17	122.64	126.20
18	b	842	CLA	O2A-CGA-O1A	-2.17	118.20	123.63
21	a	833	BCR	C15-C16-C17	-2.17	119.08	123.52
18	b	850	CLA	O2A-CGA-O1A	-2.17	118.21	123.63
26	B	302	A86	C4-C5-C6	2.17	130.32	127.28
21	a	834	BCR	C20-C19-C18	-2.17	120.42	126.36
26	H	304	A86	O4-C38-O5	-2.17	118.81	122.99
18	b	814	CLA	CMB-C2B-C3B	2.17	131.64	126.55
28	E	308	DD6	C4-C3-C2	-2.17	119.09	123.52
28	H	303	DD6	C28-C27-C26	-2.16	119.98	124.18
18	H	307	CLA	CAA-C2A-C1A	-2.16	104.88	111.97
28	C	303	DD6	C25-C24-C1	-2.16	120.44	126.36
26	B	302	A86	C9-C8-C6	2.16	132.29	126.36
18	H	306	CLA	CHB-C4A-NA	2.16	127.52	124.40
18	b	814	CLA	O2A-CGA-O1A	-2.16	118.22	123.63
28	H	303	DD6	C12-C11-C10	-2.16	119.32	122.82
18	E	313	CLA	C1-C2-C3	-2.16	122.66	126.20
21	b	833	BCR	C16-C17-C18	-2.16	124.25	127.28
18	C	307	CLA	C1-C2-C3	-2.15	122.67	126.20
18	b	830	CLA	O2A-CGA-O1A	-2.15	118.24	123.63
18	D	210	CLA	O2A-CGA-O1A	-2.15	118.24	123.63
18	a	815	CLA	O2A-CGA-O1A	-2.15	118.24	123.63

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
18	D	209	CLA	O2A-CGA-O1A	-2.15	118.24	123.63
18	a	837	CLA	CGD-CBD-CAD	-2.15	103.88	110.85
21	b	839	BCR	C33-C5-C6	-2.15	122.14	124.48
18	l	202	CLA	CHB-C4A-NA	2.15	127.50	124.40
18	E	310	CLA	O2A-CGA-O1A	-2.15	118.25	123.63
18	b	827	CLA	O2A-CGA-O1A	-2.15	118.25	123.63
28	E	307	DD6	C37-C36-C31	-2.15	120.14	124.16
18	H	314	CLA	CHB-C4A-NA	2.15	127.50	124.40
18	a	804	CLA	O2A-CGA-O1A	-2.15	118.25	123.63
21	f	205	BCR	C33-C5-C4	2.15	118.17	113.60
21	f	205	BCR	C23-C24-C25	-2.15	121.26	127.00
18	H	307	CLA	CMB-C2B-C1B	-2.15	122.15	125.42
18	a	810	CLA	O2A-CGA-O1A	-2.15	118.26	123.63
18	b	846	CLA	O2A-CGA-O1A	-2.14	118.26	123.63
18	b	803	CLA	CHC-C4B-NB	2.14	127.27	124.05
18	C	314	CLA	CHB-C4A-NA	2.14	127.49	124.40
18	H	314	CLA	O2A-CGA-O1A	-2.14	118.27	123.63
26	B	305	A86	C-C1-C24	2.14	121.36	118.09
26	H	302	A86	C9-C8-C6	2.14	132.24	126.36
18	b	813	CLA	C1-C2-C3	-2.14	122.69	126.20
18	a	804	CLA	CMB-C2B-C1B	-2.14	122.16	125.42
21	i	103	BCR	C38-C26-C27	2.14	118.16	113.60
29	H	312	KC1	O2D-CGD-O1D	-2.14	119.68	123.85
18	a	818	CLA	CMB-C2B-C1B	-2.14	122.16	125.42
21	l	201	BCR	C33-C5-C4	2.14	118.16	113.60
26	B	305	A86	C4-C5-C6	2.14	130.28	127.28
26	H	301	A86	C12-C11-C10	-2.14	118.48	123.67
18	H	310	CLA	O2A-CGA-O1A	-2.13	118.29	123.63
18	a	825	CLA	C1-C2-C3	-2.13	122.70	126.20
18	b	828	CLA	CMB-C2B-C1B	-2.13	122.17	125.42
18	a	855	CLA	O2A-CGA-O1A	-2.13	118.30	123.63
18	E	312	CLA	C1-C2-C3	-2.13	122.71	126.20
21	f	201	BCR	C23-C22-C21	2.13	122.36	119.01
18	a	822	CLA	O2A-CGA-O1A	-2.13	118.30	123.63
18	H	314	CLA	CMB-C2B-C3B	2.13	131.55	126.55
28	E	303	DD6	C37-C36-C31	-2.13	120.18	124.16
26	C	305	A86	C35-C34-C33	2.13	113.71	109.89
26	H	302	A86	C4-C5-C6	2.13	130.26	127.28
23	D	202	LMG	O7-C10-O9	-2.13	118.74	123.70
18	a	824	CLA	CMB-C2B-C1B	-2.13	122.18	125.42
18	b	821	CLA	O2A-CGA-O1A	-2.12	118.32	123.63
27	l	206	ET4	C27-C09-C08	2.12	121.33	118.09

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
18	a	820	CLA	O2A-CGA-O1A	-2.12	118.32	123.63
18	b	827	CLA	C3B-C4B-NB	-2.12	108.64	110.53
28	H	303	DD6	C19-C18-C17	-2.12	106.82	110.79
18	a	824	CLA	O2A-CGA-O1A	-2.12	118.32	123.63
21	i	102	BCR	C10-C11-C12	-2.12	117.06	123.20
18	H	311	CLA	CHB-C4A-NA	2.12	127.46	124.40
18	B	308	CLA	O2A-CGA-O1A	-2.12	118.33	123.63
18	H	306	CLA	O2A-CGA-O1A	-2.12	118.33	123.63
18	b	817	CLA	O2A-CGA-O1A	-2.12	118.33	123.63
21	b	839	BCR	C16-C15-C14	-2.12	119.19	123.52
18	b	818	CLA	O2A-CGA-O1A	-2.12	118.33	123.63
18	D	214	CLA	O2A-CGA-O1A	-2.12	118.33	123.63
26	m	102	A86	O-C13-C14	-2.12	117.42	121.76
18	b	812	CLA	CMB-C2B-C1B	-2.12	122.20	125.42
21	a	850	BCR	C35-C13-C14	-2.12	119.39	122.82
21	E	305	BCR	C36-C18-C19	2.12	121.32	118.09
18	E	316	CLA	CMB-C2B-C3B	2.11	131.52	126.55
21	a	845	BCR	C33-C5-C4	2.11	118.10	113.60
21	i	102	BCR	C8-C7-C6	-2.11	121.36	127.00
18	b	828	CLA	O2A-CGA-O1A	-2.11	118.35	123.63
18	a	819	CLA	O2A-CGA-O1A	-2.11	118.35	123.63
18	b	838	CLA	O2A-CGA-O1A	-2.11	118.36	123.63
18	D	213	CLA	O2A-CGA-O1A	-2.11	118.36	123.63
23	a	851	LMG	O6-C5-C4	2.11	113.50	109.70
27	l	206	ET4	C28-C13-C14	-2.11	119.41	122.82
18	b	820	CLA	CAA-CBA-CGA	-2.11	107.23	113.21
18	C	309	CLA	CAA-CBA-CGA	-2.10	107.23	113.21
26	B	305	A86	O4-C38-O5	-2.10	118.93	122.99
26	B	301	A86	C10-C9-C8	2.10	129.30	123.20
18	b	843	CLA	O1D-CGD-CBD	2.10	128.67	124.52
18	a	803	CLA	O2A-CGA-O1A	-2.10	118.37	123.63
21	j	101	BCR	C38-C26-C27	2.10	118.08	113.60
18	b	806	CLA	O2A-CGA-O1A	-2.10	118.37	123.63
26	B	304	A86	O4-C38-O5	-2.10	118.94	122.99
28	E	307	DD6	C8-C6-C5	2.10	122.31	119.01
18	a	830	CLA	O2A-CGA-O1A	-2.10	118.37	123.63
18	a	807	CLA	CAA-CBA-CGA	-2.10	107.25	113.21
18	f	203	CLA	O2A-CGA-O1A	-2.10	118.38	123.63
21	E	305	BCR	C29-C30-C25	2.10	113.49	110.44
26	D	206	A86	C9-C8-C6	2.10	132.11	126.36
28	C	303	DD6	C21-C20-C15	-2.10	118.85	122.30
26	m	102	A86	C21-C20-C15	2.10	130.12	123.35

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
26	H	304	A86	O-C13-C14	-2.10	117.47	121.76
26	D	206	A86	O-C13-C11	-2.09	116.53	121.04
18	B	311	CLA	O2A-CGA-O1A	-2.09	118.39	123.63
23	E	318	LMG	O3-C3-C2	-2.09	105.45	110.38
21	a	852	BCR	C33-C5-C4	2.09	118.05	113.60
18	a	814	CLA	O2A-CGA-O1A	-2.09	118.40	123.63
26	r	203	A86	O4-C34-C35	2.09	112.98	107.64
18	H	313	CLA	O2A-CGA-O1A	-2.09	118.41	123.63
26	H	301	A86	C21-C20-C15	2.09	130.09	123.35
21	a	833	BCR	C4-C5-C6	-2.09	119.89	122.70
18	B	314	CLA	CHB-C1B-NB	2.09	127.18	124.05
18	b	849	CLA	C1-C2-C3	-2.08	122.78	126.20
18	a	818	CLA	C1-C2-C3	-2.08	122.79	126.20
21	f	201	BCR	C8-C7-C6	-2.08	121.44	127.00
18	f	202	CLA	O2A-CGA-O1A	-2.08	118.43	123.63
18	b	818	CLA	C1-C2-C3	-2.08	122.79	126.20
18	a	825	CLA	C6-C7-C8	-2.08	109.06	115.97
18	a	819	CLA	C1-C2-C3	-2.08	122.79	126.20
18	b	807	CLA	O2A-CGA-O1A	-2.07	118.44	123.63
18	a	826	CLA	O2A-CGA-O1A	-2.07	118.44	123.63
18	a	807	CLA	CHB-C4A-NA	2.07	127.39	124.40
18	C	314	CLA	O2D-CGD-CBD	2.07	114.85	111.23
18	a	824	CLA	C1-C2-C3	-2.07	122.81	126.20
21	a	833	BCR	C39-C30-C25	-2.07	107.00	110.24
20	H	316	LHG	O8-C23-C24	2.07	121.16	112.38
18	E	311	CLA	C2A-C1A-CHA	2.07	127.46	123.87
21	E	305	BCR	C23-C24-C25	-2.07	121.47	127.00
26	C	304	A86	C12-C11-C13	2.07	119.35	116.00
18	b	808	CLA	C1-C2-C3	-2.07	122.81	126.20
20	E	301	LHG	C26-C25-C24	2.07	120.72	113.13
18	D	208	CLA	CHB-C4A-NA	2.07	127.38	124.40
18	a	841	CLA	CAA-CBA-CGA	-2.07	107.34	113.21
21	f	201	BCR	C2-C1-C6	-2.07	107.44	110.44
21	a	834	BCR	C34-C9-C8	2.06	121.24	118.09
18	b	849	CLA	O2A-CGA-O1A	-2.06	118.46	123.63
18	D	214	CLA	CMB-C2B-C1B	-2.06	122.28	125.42
18	E	313	CLA	O2A-CGA-O1A	-2.06	118.47	123.63
21	b	833	BCR	C38-C26-C27	2.06	118.00	113.60
21	a	833	BCR	C23-C24-C25	-2.06	121.49	127.00
18	D	207	CLA	CHB-C1B-NB	2.06	127.14	124.05
21	m	101	BCR	C38-C26-C25	-2.06	122.23	124.48
26	H	302	A86	C34-O4-C38	-2.06	114.21	117.85

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
21	a	852	BCR	C23-C24-C25	-2.06	121.50	127.00
18	a	801	CLA	C1-C2-C3	-2.06	122.82	126.20
18	a	825	CLA	CHB-C1B-NB	2.06	127.14	124.05
18	C	309	CLA	O2D-CGD-CBD	2.06	114.83	111.23
26	B	302	A86	C34-O4-C38	-2.06	114.21	117.85
18	a	817	CLA	O2A-CGA-O1A	-2.06	118.48	123.63
18	B	314	CLA	C2A-C3A-C4A	2.06	104.00	101.59
28	E	307	DD6	C4-C3-C2	2.05	127.72	123.52
18	B	311	CLA	CAC-C3C-C4C	2.05	127.46	124.79
18	b	825	CLA	O2A-CGA-O1A	-2.05	118.05	123.33
18	a	843	CLA	C2A-C1A-CHA	2.05	127.43	123.87
18	b	819	CLA	O2A-CGA-O1A	-2.05	118.50	123.63
26	C	305	A86	C-C1-C24	2.05	121.22	118.09
20	E	317	LHG	O8-C23-O10	-2.05	118.50	123.63
18	b	844	CLA	O2A-CGA-O1A	-2.05	118.50	123.63
18	a	813	CLA	CAC-C3C-C2C	-2.05	123.80	127.56
18	E	312	CLA	C11-C12-C13	-2.05	109.16	115.97
18	b	801	CLA	CMB-C2B-C1B	-2.05	122.30	125.42
18	a	828	CLA	O2A-CGA-O1A	-2.05	118.51	123.63
18	a	805	CLA	CMB-C2B-C3B	2.05	131.37	126.55
23	j	103	LMG	O6-C5-C6	2.05	111.51	106.44
18	b	805	CLA	O2A-CGA-O1A	-2.05	118.51	123.63
21	b	832	BCR	C27-C26-C25	-2.04	119.94	122.70
18	a	827	CLA	C1-C2-C3	-2.04	122.85	126.20
29	B	313	KC1	CAB-C3B-C4B	2.04	129.70	124.82
18	C	312	CLA	CHB-C1B-NB	2.04	127.11	124.05
18	b	803	CLA	CHB-C4A-NA	2.04	127.35	124.40
18	b	803	CLA	CBA-CAA-C2A	-2.04	107.72	113.79
26	E	302	A86	C9-C8-C6	2.04	131.96	126.36
18	a	811	CLA	CMB-C2B-C1B	-2.04	122.31	125.42
18	D	217	CLA	CAA-C2A-C3A	-2.04	111.55	116.23
18	a	820	CLA	CMB-C2B-C1B	-2.04	122.31	125.42
21	b	833	BCR	C33-C5-C4	2.04	117.94	113.60
28	E	308	DD6	C28-C27-C26	-2.04	120.22	124.18
18	H	305	CLA	O2D-CGD-CBD	2.04	114.79	111.23
18	a	821	CLA	O2A-CGA-O1A	-2.04	118.53	123.63
18	i	101	CLA	O2A-CGA-O1A	-2.04	118.53	123.63
21	i	103	BCR	C33-C5-C4	2.04	117.94	113.60
26	C	302	A86	C9-C8-C6	2.04	131.94	126.36
18	b	829	CLA	C11-C10-C8	-2.04	109.20	115.97
21	a	850	BCR	C8-C7-C6	-2.03	121.56	127.00
20	E	317	LHG	O7-C7-O9	-2.03	118.95	123.70

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
18	H	307	CLA	CMB-C2B-C3B	2.03	131.33	126.55
18	a	826	CLA	O2D-CGD-CBD	2.03	114.78	111.23
18	E	309	CLA	CMB-C2B-C3B	2.03	131.33	126.55
18	b	824	CLA	O2A-CGA-O1A	-2.03	118.55	123.63
18	D	216	CLA	O1D-CGD-CBD	2.03	128.53	124.52
18	a	825	CLA	C6-C5-C3	-2.03	108.52	113.47
18	a	848	CLA	O2A-CGA-O1A	-2.03	118.55	123.63
18	l	205	CLA	CMB-C2B-C3B	2.03	131.32	126.55
18	a	847	CLA	O2A-CGA-O1A	-2.03	118.56	123.63
18	a	841	CLA	CMB-C2B-C1B	-2.03	122.33	125.42
18	B	309	CLA	O2A-CGA-O1A	-2.03	118.56	123.63
18	H	305	CLA	C1-C2-C3	-2.03	122.88	126.20
18	a	806	CLA	O2A-CGA-O1A	-2.02	118.56	123.63
18	r	202	CLA	O2A-CGA-O1A	-2.02	118.12	123.33
28	H	303	DD6	O4-C34-C33	-2.02	105.61	109.75
18	b	809	CLA	O2A-CGA-O1A	-2.02	118.57	123.63
21	a	834	BCR	C16-C15-C14	-2.02	119.38	123.52
18	a	801	CLA	C1D-ND-C4D	2.02	107.73	106.31
18	a	814	CLA	C1-C2-C3	-2.02	122.88	126.20
18	B	307	CLA	O2A-CGA-O1A	-2.02	118.58	123.63
18	a	807	CLA	C3A-C2A-C1A	2.02	104.36	101.34
18	a	838	CLA	C3B-C4B-NB	-2.02	108.73	110.53
18	a	844	CLA	O2A-CGA-O1A	-2.02	118.58	123.63
18	H	311	CLA	C6-C5-C3	-2.02	108.56	113.47
18	b	841	CLA	O2A-CGA-O1A	-2.02	118.58	123.63
21	a	845	BCR	C2-C3-C4	-2.02	106.84	111.28
19	a	829	PQN	C2M-C2-C3	-2.01	121.14	124.45
21	a	852	BCR	C15-C16-C17	-2.01	119.40	123.52
26	b	847	A86	C19-C18-C17	-2.01	107.03	110.79
18	b	824	CLA	CMB-C2B-C1B	-2.01	122.36	125.42
18	C	312	CLA	O2A-CGA-O1A	-2.01	118.61	123.63
21	l	204	BCR	C35-C13-C14	-2.01	119.57	122.82
29	H	312	KC1	CBA-CAA-C2A	-2.00	117.41	125.45
21	l	201	BCR	C32-C1-C6	-2.00	107.10	110.24
18	C	310	CLA	O1D-CGD-CBD	2.00	128.47	124.52
28	E	303	DD6	C15-C14-C13	-2.00	121.76	125.99
18	a	813	CLA	CHB-C1B-NB	2.00	127.06	124.05
18	a	822	CLA	C1-C2-C3	-2.00	123.52	126.76
18	a	811	CLA	O2A-CGA-O1A	-2.00	118.62	123.63
28	E	308	DD6	C37-C36-C35	2.00	118.10	114.42
18	a	824	CLA	CMB-C2B-C3B	2.00	131.26	126.55
18	D	214	CLA	O1D-CGD-CBD	2.00	128.46	124.52

All (142) chirality outliers are listed below:

Mol	Chain	Res	Type	Atom
18	a	801	CLA	ND
18	a	802	CLA	ND
18	a	803	CLA	ND
18	a	804	CLA	ND
18	a	805	CLA	ND
18	a	806	CLA	ND
18	a	807	CLA	ND
18	a	808	CLA	ND
18	a	809	CLA	ND
18	a	810	CLA	ND
18	a	811	CLA	ND
18	a	812	CLA	ND
18	a	813	CLA	ND
18	a	814	CLA	ND
18	a	815	CLA	ND
18	a	816	CLA	ND
18	a	817	CLA	ND
18	a	818	CLA	ND
18	a	819	CLA	ND
18	a	820	CLA	ND
18	a	821	CLA	ND
18	a	822	CLA	ND
18	a	823	CLA	ND
18	a	824	CLA	ND
18	a	825	CLA	ND
18	a	826	CLA	ND
18	a	827	CLA	ND
18	a	828	CLA	ND
18	a	830	CLA	ND
18	a	836	CLA	ND
18	a	837	CLA	ND
18	a	838	CLA	ND
18	a	839	CLA	ND
18	a	840	CLA	ND
18	a	841	CLA	ND
18	a	842	CLA	ND
18	a	843	CLA	ND
18	a	844	CLA	ND
18	a	846	CLA	ND
18	a	847	CLA	ND
18	a	848	CLA	ND
18	a	853	CLA	ND

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Mol	Chain	Res	Type	Atom
18	a	854	CLA	ND
18	a	855	CLA	ND
18	b	801	CLA	ND
18	b	803	CLA	ND
18	b	804	CLA	ND
18	b	805	CLA	ND
18	b	806	CLA	ND
18	b	807	CLA	ND
18	b	808	CLA	ND
18	b	809	CLA	ND
18	b	810	CLA	ND
18	b	811	CLA	ND
18	b	812	CLA	ND
18	b	813	CLA	ND
18	b	814	CLA	ND
18	b	815	CLA	ND
18	b	816	CLA	ND
18	b	817	CLA	ND
18	b	818	CLA	ND
18	b	819	CLA	ND
18	b	820	CLA	ND
18	b	821	CLA	ND
18	b	822	CLA	ND
18	b	823	CLA	ND
18	b	824	CLA	ND
18	b	825	CLA	ND
18	b	826	CLA	ND
18	b	827	CLA	ND
18	b	828	CLA	ND
18	b	829	CLA	ND
18	b	830	CLA	ND
18	b	837	CLA	ND
18	b	838	CLA	ND
18	b	840	CLA	ND
18	b	841	CLA	ND
18	b	842	CLA	ND
18	b	843	CLA	ND
18	b	844	CLA	ND
18	b	845	CLA	ND
18	b	846	CLA	ND
18	b	848	CLA	ND
18	b	849	CLA	ND

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Mol	Chain	Res	Type	Atom
18	b	850	CLA	ND
18	f	202	CLA	ND
18	f	203	CLA	ND
18	f	204	CLA	ND
18	i	101	CLA	ND
18	j	104	CLA	ND
18	l	202	CLA	ND
18	l	203	CLA	ND
18	l	205	CLA	ND
18	r	202	CLA	ND
18	B	306	CLA	ND
18	B	307	CLA	ND
18	B	308	CLA	ND
18	B	309	CLA	ND
18	B	310	CLA	ND
18	B	311	CLA	ND
18	B	312	CLA	ND
18	B	314	CLA	ND
18	C	306	CLA	ND
18	C	307	CLA	ND
18	C	309	CLA	ND
18	C	310	CLA	ND
18	C	311	CLA	ND
18	C	312	CLA	ND
18	C	314	CLA	ND
18	C	315	CLA	ND
18	C	316	CLA	ND
18	C	317	CLA	ND
18	C	318	CLA	ND
18	D	207	CLA	ND
18	D	208	CLA	ND
18	D	209	CLA	ND
18	D	210	CLA	ND
18	D	211	CLA	ND
18	D	212	CLA	ND
18	D	213	CLA	ND
18	D	214	CLA	ND
18	D	215	CLA	ND
18	D	216	CLA	ND
18	D	217	CLA	ND
18	E	309	CLA	ND
18	E	310	CLA	ND

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Mol	Chain	Res	Type	Atom
18	E	311	CLA	ND
18	E	312	CLA	ND
18	E	313	CLA	ND
18	E	314	CLA	ND
18	E	315	CLA	ND
18	E	316	CLA	ND
18	H	305	CLA	ND
18	H	306	CLA	ND
18	H	307	CLA	ND
18	H	308	CLA	ND
18	H	309	CLA	ND
18	H	310	CLA	ND
18	H	311	CLA	ND
18	H	313	CLA	ND
18	H	314	CLA	ND
18	H	315	CLA	ND

All (2059) torsion outliers are listed below:

Mol	Chain	Res	Type	Atoms
18	a	801	CLA	CBD-CGD-O2D-CED
18	a	801	CLA	O1D-CGD-O2D-CED
18	a	803	CLA	CHA-CBD-CGD-O1D
18	a	803	CLA	CHA-CBD-CGD-O2D
18	a	804	CLA	C1A-C2A-CAA-CBA
18	a	804	CLA	C3A-C2A-CAA-CBA
18	a	805	CLA	C3A-C2A-CAA-CBA
18	a	808	CLA	C4B-C3B-CAB-CBB
18	a	809	CLA	CAD-CBD-CGD-O2D
18	a	813	CLA	C1A-C2A-CAA-CBA
18	a	813	CLA	CHA-CBD-CGD-O1D
18	a	813	CLA	CHA-CBD-CGD-O2D
18	a	814	CLA	C3A-C2A-CAA-CBA
18	a	815	CLA	C1A-C2A-CAA-CBA
18	a	815	CLA	C3A-C2A-CAA-CBA
18	a	816	CLA	C1A-C2A-CAA-CBA
18	a	816	CLA	C3A-C2A-CAA-CBA
18	a	820	CLA	CHA-CBD-CGD-O1D
18	a	820	CLA	CHA-CBD-CGD-O2D
18	a	821	CLA	C1A-C2A-CAA-CBA
18	a	821	CLA	C3A-C2A-CAA-CBA
18	a	823	CLA	C1A-C2A-CAA-CBA

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Mol	Chain	Res	Type	Atoms
18	a	823	CLA	C3A-C2A-CAA-CBA
18	a	824	CLA	C1A-C2A-CAA-CBA
18	a	828	CLA	CAD-CBD-CGD-O1D
18	a	828	CLA	CAD-CBD-CGD-O2D
18	a	837	CLA	C1A-C2A-CAA-CBA
18	a	837	CLA	C3A-C2A-CAA-CBA
18	a	837	CLA	CHA-CBD-CGD-O1D
18	a	837	CLA	CHA-CBD-CGD-O2D
18	a	838	CLA	C1A-C2A-CAA-CBA
18	a	839	CLA	CHA-CBD-CGD-O1D
18	a	839	CLA	CHA-CBD-CGD-O2D
18	a	841	CLA	CAD-CBD-CGD-O1D
18	a	841	CLA	CAD-CBD-CGD-O2D
18	a	841	CLA	C11-C12-C13-C14
18	a	844	CLA	C1A-C2A-CAA-CBA
18	a	844	CLA	C3A-C2A-CAA-CBA
18	a	846	CLA	C1A-C2A-CAA-CBA
18	a	846	CLA	C3A-C2A-CAA-CBA
18	a	853	CLA	CAD-CBD-CGD-O1D
18	a	853	CLA	CAD-CBD-CGD-O2D
18	a	854	CLA	CBD-CGD-O2D-CED
18	a	854	CLA	C2-C3-C5-C6
18	a	854	CLA	C4-C3-C5-C6
18	b	803	CLA	CBD-CGD-O2D-CED
18	b	804	CLA	CHA-CBD-CGD-O1D
18	b	804	CLA	CHA-CBD-CGD-O2D
18	b	805	CLA	C3A-C2A-CAA-CBA
18	b	806	CLA	CBD-CGD-O2D-CED
18	b	809	CLA	C1A-C2A-CAA-CBA
18	b	812	CLA	O1A-CGA-O2A-C1
18	b	813	CLA	CAD-CBD-CGD-O1D
18	b	813	CLA	CAD-CBD-CGD-O2D
18	b	814	CLA	C1A-C2A-CAA-CBA
18	b	814	CLA	C3A-C2A-CAA-CBA
18	b	817	CLA	C1A-C2A-CAA-CBA
18	b	817	CLA	C3A-C2A-CAA-CBA
18	b	818	CLA	C1A-C2A-CAA-CBA
18	b	818	CLA	C3A-C2A-CAA-CBA
18	b	819	CLA	C1A-C2A-CAA-CBA
18	b	819	CLA	C3A-C2A-CAA-CBA
18	b	820	CLA	CHA-CBD-CGD-O1D
18	b	820	CLA	CHA-CBD-CGD-O2D

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Mol	Chain	Res	Type	Atoms
18	b	820	CLA	C11-C12-C13-C14
18	b	821	CLA	C1A-C2A-CAA-CBA
18	b	821	CLA	C3A-C2A-CAA-CBA
18	b	822	CLA	C1A-C2A-CAA-CBA
18	b	822	CLA	C3A-C2A-CAA-CBA
18	b	823	CLA	C1A-C2A-CAA-CBA
18	b	823	CLA	CHA-CBD-CGD-O1D
18	b	823	CLA	CHA-CBD-CGD-O2D
18	b	824	CLA	C1A-C2A-CAA-CBA
18	b	828	CLA	C4B-C3B-CAB-CBB
18	b	840	CLA	CHA-CBD-CGD-O1D
18	b	840	CLA	CHA-CBD-CGD-O2D
18	b	841	CLA	C1A-C2A-CAA-CBA
18	b	841	CLA	C3A-C2A-CAA-CBA
18	b	841	CLA	CBD-CGD-O2D-CED
18	b	844	CLA	C1A-C2A-CAA-CBA
18	b	845	CLA	C1A-C2A-CAA-CBA
18	b	845	CLA	C3A-C2A-CAA-CBA
18	b	848	CLA	CHA-CBD-CGD-O1D
18	b	848	CLA	CHA-CBD-CGD-O2D
18	b	849	CLA	C4-C3-C5-C6
18	f	202	CLA	CBD-CGD-O2D-CED
18	l	203	CLA	C2-C3-C5-C6
18	l	203	CLA	C4-C3-C5-C6
18	l	203	CLA	C11-C10-C8-C9
18	r	202	CLA	C1A-C2A-CAA-CBA
18	r	202	CLA	C3A-C2A-CAA-CBA
18	r	202	CLA	CBD-CGD-O2D-CED
18	B	306	CLA	C1A-C2A-CAA-CBA
18	B	306	CLA	C3A-C2A-CAA-CBA
18	B	309	CLA	C4-C3-C5-C6
18	B	310	CLA	CBA-CGA-O2A-C1
18	B	310	CLA	C4B-C3B-CAB-CBB
18	B	310	CLA	CBD-CGD-O2D-CED
18	B	310	CLA	O1D-CGD-O2D-CED
18	B	311	CLA	CBD-CGD-O2D-CED
18	B	312	CLA	C3A-C2A-CAA-CBA
18	B	314	CLA	CBD-CGD-O2D-CED
18	C	310	CLA	CBA-CGA-O2A-C1
18	C	310	CLA	C2B-C3B-CAB-CBB
18	C	310	CLA	C4B-C3B-CAB-CBB
18	C	310	CLA	CBD-CGD-O2D-CED

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Mol	Chain	Res	Type	Atoms
18	C	311	CLA	CBD-CGD-O2D-CED
18	C	312	CLA	C3A-C2A-CAA-CBA
18	C	312	CLA	C4B-C3B-CAB-CBB
18	C	315	CLA	C1A-C2A-CAA-CBA
18	C	316	CLA	C2B-C3B-CAB-CBB
18	C	316	CLA	C4B-C3B-CAB-CBB
18	C	317	CLA	C1A-C2A-CAA-CBA
18	C	318	CLA	C1A-C2A-CAA-CBA
18	C	318	CLA	C3A-C2A-CAA-CBA
18	D	210	CLA	CHA-CBD-CGD-O1D
18	D	210	CLA	CHA-CBD-CGD-O2D
18	D	215	CLA	CBD-CGD-O2D-CED
18	D	216	CLA	C1A-C2A-CAA-CBA
18	D	216	CLA	C3A-C2A-CAA-CBA
18	E	310	CLA	C4B-C3B-CAB-CBB
18	E	311	CLA	CBA-CGA-O2A-C1
18	E	311	CLA	O1A-CGA-O2A-C1
18	E	311	CLA	C4B-C3B-CAB-CBB
18	E	315	CLA	CAD-CBD-CGD-O1D
18	E	315	CLA	CAD-CBD-CGD-O2D
18	H	305	CLA	C1A-C2A-CAA-CBA
18	H	305	CLA	C3A-C2A-CAA-CBA
18	H	311	CLA	C1A-C2A-CAA-CBA
18	H	311	CLA	C3A-C2A-CAA-CBA
18	H	313	CLA	CHA-CBD-CGD-O1D
18	H	313	CLA	CHA-CBD-CGD-O2D
18	H	315	CLA	CBD-CGD-O2D-CED
18	H	315	CLA	O1D-CGD-O2D-CED
19	a	829	PQN	C12-C13-C15-C16
19	a	829	PQN	C14-C13-C15-C16
20	a	832	LHG	C3-O3-P-O4
20	a	832	LHG	C3-O3-P-O6
20	a	832	LHG	C8-C7-O7-C5
20	B	315	LHG	O1-C1-C2-C3
20	B	315	LHG	C3-O3-P-O4
20	B	315	LHG	C3-O3-P-O6
20	B	315	LHG	C4-O6-P-O4
20	D	201	LHG	C3-O3-P-O4
20	D	201	LHG	C3-O3-P-O5
20	D	201	LHG	C3-O3-P-O6
20	E	301	LHG	C2-C3-O3-P
20	H	316	LHG	C4-O6-P-O3

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Mol	Chain	Res	Type	Atoms
20	H	316	LHG	C4-O6-P-O4
20	H	316	LHG	C4-O6-P-O5
20	H	316	LHG	C8-C7-O7-C5
21	a	834	BCR	C37-C22-C23-C24
21	a	845	BCR	C23-C24-C25-C26
21	a	845	BCR	C23-C24-C25-C30
21	a	850	BCR	C1-C6-C7-C8
21	a	850	BCR	C5-C6-C7-C8
21	a	852	BCR	C5-C6-C7-C8
21	b	836	BCR	C7-C8-C9-C10
21	b	836	BCR	C7-C8-C9-C34
21	b	836	BCR	C19-C20-C21-C22
21	b	836	BCR	C21-C22-C23-C24
21	b	836	BCR	C37-C22-C23-C24
21	b	839	BCR	C23-C24-C25-C26
21	b	839	BCR	C23-C24-C25-C30
21	f	201	BCR	C5-C6-C7-C8
21	f	201	BCR	C17-C18-C19-C20
21	f	201	BCR	C36-C18-C19-C20
21	f	201	BCR	C21-C22-C23-C24
21	f	201	BCR	C37-C22-C23-C24
21	i	103	BCR	C21-C22-C23-C24
21	i	103	BCR	C37-C22-C23-C24
21	j	105	BCR	C21-C22-C23-C24
21	j	105	BCR	C37-C22-C23-C24
21	l	204	BCR	C21-C22-C23-C24
21	l	204	BCR	C37-C22-C23-C24
21	m	101	BCR	C7-C8-C9-C10
22	a	849	SQD	O49-C7-O47-C45
22	a	849	SQD	O5-C5-C6-S
22	a	849	SQD	C5-C6-S-O7
23	a	851	LMG	O7-C8-C9-O8
23	j	103	LMG	O10-C28-O8-C9
23	j	103	LMG	C29-C28-O8-C9
23	l	207	LMG	O6-C1-O1-C7
23	l	207	LMG	O7-C8-C9-O8
23	l	207	LMG	C11-C10-O7-C8
23	C	301	LMG	O6-C1-O1-C7
23	E	318	LMG	O7-C8-C9-O8
26	b	847	A86	C-C1-C24-C25
26	b	847	A86	C2-C1-C24-C25
26	b	847	A86	C11-C10-C9-C8

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Mol	Chain	Res	Type	Atoms
26	B	301	A86	C26-C27-C29-C30
26	B	301	A86	C28-C27-C29-C30
26	B	301	A86	C39-C38-O4-C34
26	B	302	A86	C39-C38-O4-C34
26	B	305	A86	C1-C2-C3-C4
26	C	302	A86	C39-C38-O4-C34
26	C	304	A86	C35-C34-O4-C38
26	C	305	A86	C1-C2-C3-C4
26	C	305	A86	O5-C38-O4-C34
26	D	203	A86	C2-C1-C24-C25
26	D	203	A86	O-C13-C14-C15
26	D	203	A86	O5-C38-O4-C34
26	D	206	A86	C1-C2-C3-C4
26	D	206	A86	C39-C38-O4-C34
26	D	206	A86	O5-C38-O4-C34
26	E	302	A86	C1-C2-C3-C4
26	H	301	A86	O5-C38-O4-C34
26	H	302	A86	C39-C38-O4-C34
27	I	206	ET4	C05-C06-C07-C08
28	E	303	DD6	C10-C11-C13-C14
28	E	304	DD6	C13-C14-C15-C16
28	E	304	DD6	C13-C14-C15-O1
28	E	307	DD6	C13-C14-C15-C16
28	E	307	DD6	C13-C14-C15-C20
28	E	307	DD6	C13-C14-C15-O1
28	E	307	DD6	C5-C6-C8-C9
28	E	307	DD6	C7-C6-C8-C9
29	B	313	KC1	C2A-CAA-CBA-CGA
29	B	313	KC1	CBD-CGD-O2D-CED
29	C	308	KC1	C3A-C2A-CAA-CBA
29	C	308	KC1	C2B-C3B-CAB-CBB
29	C	308	KC1	C2A-CAA-CBA-CGA
29	C	308	KC1	CHA-CBD-CGD-O1D
29	C	308	KC1	CHA-CBD-CGD-O2D
29	C	313	KC1	C3A-C2A-CAA-CBA
29	C	313	KC1	C2B-C3B-CAB-CBB
18	B	311	CLA	O1D-CGD-O2D-CED
18	B	314	CLA	O1D-CGD-O2D-CED
18	C	311	CLA	O1D-CGD-O2D-CED
18	C	314	CLA	O1D-CGD-O2D-CED
26	m	102	A86	C39-C38-O4-C34
26	B	301	A86	O5-C38-O4-C34

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Mol	Chain	Res	Type	Atoms
26	B	302	A86	O5-C38-O4-C34
26	D	203	A86	C39-C38-O4-C34
26	E	302	A86	C39-C38-O4-C34
26	H	301	A86	C39-C38-O4-C34
26	H	302	A86	O5-C38-O4-C34
18	a	813	CLA	C2C-C3C-CAC-CBC
18	a	802	CLA	O1D-CGD-O2D-CED
18	b	806	CLA	O1D-CGD-O2D-CED
18	j	104	CLA	O1D-CGD-O2D-CED
26	C	302	A86	O5-C38-O4-C34
26	C	304	A86	C39-C38-O4-C34
18	a	802	CLA	CBD-CGD-O2D-CED
18	a	821	CLA	CBD-CGD-O2D-CED
18	a	853	CLA	CBD-CGD-O2D-CED
18	b	813	CLA	CBD-CGD-O2D-CED
18	j	104	CLA	CBD-CGD-O2D-CED
18	B	307	CLA	CBD-CGD-O2D-CED
18	B	312	CLA	CBD-CGD-O2D-CED
18	C	314	CLA	CBD-CGD-O2D-CED
18	C	315	CLA	CBD-CGD-O2D-CED
18	D	211	CLA	CBD-CGD-O2D-CED
18	D	217	CLA	CBD-CGD-O2D-CED
18	E	315	CLA	O1A-CGA-O2A-C1
20	j	102	LHG	O10-C23-O8-C6
18	B	310	CLA	O1A-CGA-O2A-C1
18	C	310	CLA	O1A-CGA-O2A-C1
26	C	305	A86	C39-C38-O4-C34
26	E	302	A86	C35-C34-O4-C38
18	b	813	CLA	O1D-CGD-O2D-CED
18	D	217	CLA	O1D-CGD-O2D-CED
18	E	315	CLA	CBA-CGA-O2A-C1
20	j	102	LHG	C24-C23-O8-C6
26	H	304	A86	C39-C38-O4-C34
18	E	309	CLA	CBD-CGD-O2D-CED
18	a	804	CLA	O1A-CGA-O2A-C1
18	a	811	CLA	O1A-CGA-O2A-C1
18	a	813	CLA	O1A-CGA-O2A-C1
18	a	815	CLA	O1A-CGA-O2A-C1
18	a	817	CLA	O1A-CGA-O2A-C1
18	a	838	CLA	O1A-CGA-O2A-C1
18	a	839	CLA	O1A-CGA-O2A-C1
18	b	813	CLA	O1A-CGA-O2A-C1

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Mol	Chain	Res	Type	Atoms
18	b	826	CLA	O1A-CGA-O2A-C1
18	b	849	CLA	O1A-CGA-O2A-C1
18	E	313	CLA	O1A-CGA-O2A-C1
18	H	305	CLA	O1A-CGA-O2A-C1
20	a	835	LHG	O10-C23-O8-C6
23	C	319	LMG	O10-C28-O8-C9
18	a	813	CLA	C4C-C3C-CAC-CBC
18	C	317	CLA	C2C-C3C-CAC-CBC
18	a	854	CLA	O1D-CGD-O2D-CED
18	C	310	CLA	O1D-CGD-O2D-CED
18	D	215	CLA	O1D-CGD-O2D-CED
18	b	803	CLA	O1D-CGD-O2D-CED
18	f	202	CLA	O1D-CGD-O2D-CED
18	r	202	CLA	O1D-CGD-O2D-CED
26	E	302	A86	O5-C38-O4-C34
18	E	315	CLA	CBD-CGD-O2D-CED
20	a	832	LHG	O9-C7-O7-C5
20	H	316	LHG	O9-C7-O7-C5
23	l	207	LMG	O9-C10-O7-C8
18	a	807	CLA	C3-C5-C6-C7
18	a	837	CLA	C3-C5-C6-C7
18	a	839	CLA	C3-C5-C6-C7
18	b	806	CLA	C3-C5-C6-C7
18	b	822	CLA	C3-C5-C6-C7
18	b	837	CLA	C3-C5-C6-C7
18	b	846	CLA	C3-C5-C6-C7
18	C	309	CLA	C3-C5-C6-C7
18	C	312	CLA	C3-C5-C6-C7
18	E	309	CLA	C3-C5-C6-C7
18	b	841	CLA	O1D-CGD-O2D-CED
18	a	804	CLA	CBA-CGA-O2A-C1
18	a	811	CLA	CBA-CGA-O2A-C1
18	a	813	CLA	CBA-CGA-O2A-C1
18	a	817	CLA	CBA-CGA-O2A-C1
18	a	839	CLA	CBA-CGA-O2A-C1
18	b	812	CLA	CBA-CGA-O2A-C1
18	b	813	CLA	CBA-CGA-O2A-C1
18	b	826	CLA	CBA-CGA-O2A-C1
18	b	849	CLA	CBA-CGA-O2A-C1
18	H	305	CLA	CBA-CGA-O2A-C1
26	B	305	A86	C39-C38-O4-C34
18	a	811	CLA	CBD-CGD-O2D-CED

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Mol	Chain	Res	Type	Atoms
18	a	828	CLA	CBD-CGD-O2D-CED
18	a	841	CLA	CBD-CGD-O2D-CED
18	a	842	CLA	CBD-CGD-O2D-CED
18	a	844	CLA	CBD-CGD-O2D-CED
18	b	815	CLA	CBD-CGD-O2D-CED
18	b	816	CLA	CBD-CGD-O2D-CED
18	b	817	CLA	CBD-CGD-O2D-CED
18	b	822	CLA	CBD-CGD-O2D-CED
18	C	317	CLA	CBD-CGD-O2D-CED
18	C	318	CLA	CBD-CGD-O2D-CED
18	E	313	CLA	CBD-CGD-O2D-CED
18	D	211	CLA	O1D-CGD-O2D-CED
18	H	305	CLA	C2C-C3C-CAC-CBC
29	B	313	KC1	CAA-CBA-CGA-O1A
18	a	848	CLA	C4-C3-C5-C6
18	b	829	CLA	C4-C3-C5-C6
18	b	843	CLA	C4-C3-C5-C6
18	H	314	CLA	C4-C3-C5-C6
18	a	848	CLA	C2-C3-C5-C6
18	b	849	CLA	C2-C3-C5-C6
18	B	309	CLA	C2-C3-C5-C6
18	H	314	CLA	C2-C3-C5-C6
18	b	830	CLA	CBD-CGD-O2D-CED
18	C	315	CLA	O1D-CGD-O2D-CED
26	m	102	A86	O5-C38-O4-C34
18	a	809	CLA	C2A-CAA-CBA-CGA
18	b	801	CLA	C2A-CAA-CBA-CGA
18	D	213	CLA	C2A-CAA-CBA-CGA
18	H	305	CLA	C2A-CAA-CBA-CGA
18	b	844	CLA	O1A-CGA-O2A-C1
18	a	836	CLA	C3-C5-C6-C7
18	b	840	CLA	C3-C5-C6-C7
18	D	213	CLA	C3-C5-C6-C7
18	H	307	CLA	C3-C5-C6-C7
18	H	311	CLA	C3-C5-C6-C7
18	a	803	CLA	CBA-CGA-O2A-C1
18	a	815	CLA	CBA-CGA-O2A-C1
18	a	822	CLA	CBA-CGA-O2A-C1
18	a	825	CLA	CBA-CGA-O2A-C1
18	a	838	CLA	CBA-CGA-O2A-C1
18	a	847	CLA	CBA-CGA-O2A-C1
18	a	853	CLA	CBA-CGA-O2A-C1

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Mol	Chain	Res	Type	Atoms
18	b	821	CLA	CBA-CGA-O2A-C1
18	B	308	CLA	CBA-CGA-O2A-C1
18	D	207	CLA	CBA-CGA-O2A-C1
18	E	313	CLA	CBA-CGA-O2A-C1
20	a	835	LHG	C24-C23-O8-C6
23	C	319	LMG	C29-C28-O8-C9
20	a	835	LHG	C24-C25-C26-C27
20	D	201	LHG	C24-C25-C26-C27
20	E	301	LHG	C24-C25-C26-C27
26	b	847	A86	C39-C38-O4-C34
26	b	847	A86	C24-C25-C26-C27
26	r	203	A86	C1-C2-C3-C4
26	r	203	A86	C24-C25-C26-C27
26	B	302	A86	C1-C2-C3-C4
26	B	304	A86	C1-C2-C3-C4
26	C	302	A86	C1-C2-C3-C4
26	H	301	A86	C1-C2-C3-C4
26	H	302	A86	C1-C2-C3-C4
26	H	304	A86	C1-C2-C3-C4
28	C	303	DD6	C1-C2-C3-C4
18	a	822	CLA	O1A-CGA-O2A-C1
18	a	847	CLA	O1A-CGA-O2A-C1
18	a	848	CLA	O1A-CGA-O2A-C1
18	a	803	CLA	C5-C6-C7-C8
18	b	815	CLA	CBA-CGA-O2A-C1
20	a	831	LHG	C24-C25-C26-C27
20	b	835	LHG	C24-C25-C26-C27
23	a	851	LMG	C31-C32-C33-C34
23	E	318	LMG	C4-C5-C6-O5
29	B	313	KC1	CAA-CBA-CGA-O2A
18	a	801	CLA	C3-C5-C6-C7
18	b	814	CLA	C3-C5-C6-C7
18	a	823	CLA	CBD-CGD-O2D-CED
18	b	805	CLA	CBD-CGD-O2D-CED
18	b	837	CLA	CBD-CGD-O2D-CED
20	D	201	LHG	C15-C16-C17-C18
20	D	201	LHG	C30-C31-C32-C33
18	C	317	CLA	C4C-C3C-CAC-CBC
20	a	835	LHG	O2-C2-C3-O3
18	a	853	CLA	O1D-CGD-O2D-CED
18	a	841	CLA	CBA-CGA-O2A-C1
18	b	844	CLA	CBA-CGA-O2A-C1

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Mol	Chain	Res	Type	Atoms
18	B	311	CLA	CBA-CGA-O2A-C1
18	E	312	CLA	CBA-CGA-O2A-C1
26	r	203	A86	C39-C38-O4-C34
18	B	307	CLA	O1D-CGD-O2D-CED
26	r	203	A86	O5-C38-O4-C34
26	C	304	A86	O5-C38-O4-C34
18	C	309	CLA	CBD-CGD-O2D-CED
18	H	313	CLA	CBD-CGD-O2D-CED
18	H	314	CLA	CBD-CGD-O2D-CED
20	b	835	LHG	C8-C7-O7-C5
18	a	803	CLA	O1A-CGA-O2A-C1
18	E	312	CLA	O1A-CGA-O2A-C1
18	a	821	CLA	O1D-CGD-O2D-CED
18	B	312	CLA	O1D-CGD-O2D-CED
18	B	311	CLA	C3-C5-C6-C7
18	a	806	CLA	CBD-CGD-O2D-CED
18	a	836	CLA	CBD-CGD-O2D-CED
18	a	837	CLA	CBD-CGD-O2D-CED
18	b	850	CLA	CBD-CGD-O2D-CED
18	a	848	CLA	CBA-CGA-O2A-C1
18	a	841	CLA	C4-C3-C5-C6
18	a	847	CLA	C4-C3-C5-C6
18	B	311	CLA	C4-C3-C5-C6
18	D	207	CLA	C4-C3-C5-C6
18	H	310	CLA	C4-C3-C5-C6
18	a	825	CLA	C2-C3-C5-C6
18	a	841	CLA	C2-C3-C5-C6
18	b	829	CLA	C2-C3-C5-C6
18	b	843	CLA	C2-C3-C5-C6
18	B	311	CLA	C2-C3-C5-C6
18	D	207	CLA	C2-C3-C5-C6
18	H	307	CLA	C2-C3-C5-C6
18	H	310	CLA	C2-C3-C5-C6
18	a	825	CLA	O1A-CGA-O2A-C1
18	a	841	CLA	O1A-CGA-O2A-C1
18	b	821	CLA	O1A-CGA-O2A-C1
18	B	311	CLA	O1A-CGA-O2A-C1
18	D	207	CLA	O1A-CGA-O2A-C1
20	b	835	LHG	O9-C7-O7-C5
18	a	822	CLA	CBD-CGD-O2D-CED
18	b	826	CLA	CBD-CGD-O2D-CED
18	a	847	CLA	C2A-CAA-CBA-CGA

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Mol	Chain	Res	Type	Atoms
18	a	853	CLA	O1A-CGA-O2A-C1
18	B	308	CLA	O1A-CGA-O2A-C1
23	D	202	LMG	O6-C1-O1-C7
18	E	309	CLA	O1D-CGD-O2D-CED
18	b	842	CLA	CBA-CGA-O2A-C1
18	a	805	CLA	CBD-CGD-O2D-CED
18	a	824	CLA	CBD-CGD-O2D-CED
18	a	848	CLA	CBD-CGD-O2D-CED
18	a	855	CLA	CBD-CGD-O2D-CED
18	D	208	CLA	CBD-CGD-O2D-CED
18	b	815	CLA	O1A-CGA-O2A-C1
23	E	318	LMG	O6-C5-C6-O5
20	b	835	LHG	C27-C28-C29-C30
20	D	201	LHG	C11-C12-C13-C14
20	D	201	LHG	C13-C14-C15-C16
23	j	103	LMG	O6-C5-C6-O5
18	a	812	CLA	CBD-CGD-O2D-CED
18	b	818	CLA	CBD-CGD-O2D-CED
18	b	848	CLA	CBD-CGD-O2D-CED
20	B	315	LHG	C8-C7-O7-C5
20	b	835	LHG	C13-C14-C15-C16
18	H	313	CLA	C3-C5-C6-C7
18	E	315	CLA	O1D-CGD-O2D-CED
26	b	847	A86	C1-C2-C3-C4
18	b	842	CLA	O1A-CGA-O2A-C1
18	H	305	CLA	C4C-C3C-CAC-CBC
20	a	835	LHG	C1-C2-C3-O3
20	B	315	LHG	C1-C2-C3-O3
20	D	201	LHG	C1-C2-C3-O3
20	H	316	LHG	C1-C2-C3-O3
18	C	318	CLA	O1D-CGD-O2D-CED
18	a	806	CLA	CBA-CGA-O2A-C1
18	a	824	CLA	CBA-CGA-O2A-C1
18	b	801	CLA	CBA-CGA-O2A-C1
18	E	309	CLA	CBA-CGA-O2A-C1
20	E	301	LHG	C24-C23-O8-C6
23	l	207	LMG	C29-C28-O8-C9
25	b	834	DGD	O6E-C5E-C6E-O5E
23	j	103	LMG	C4-C5-C6-O5
23	l	207	LMG	C21-C22-C23-C24
23	C	319	LMG	C11-C12-C13-C14
18	E	313	CLA	O1D-CGD-O2D-CED

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Mol	Chain	Res	Type	Atoms
18	a	825	CLA	C4-C3-C5-C6
18	H	307	CLA	C4-C3-C5-C6
19	b	831	PQN	C14-C13-C15-C16
18	a	847	CLA	C2-C3-C5-C6
18	b	820	CLA	C3-C5-C6-C7
18	a	847	CLA	CBD-CGD-O2D-CED
18	l	205	CLA	CBD-CGD-O2D-CED
18	a	813	CLA	C6-C7-C8-C9
18	a	813	CLA	C11-C12-C13-C14
18	a	813	CLA	C14-C13-C15-C16
18	a	854	CLA	C6-C7-C8-C9
18	b	803	CLA	C11-C12-C13-C14
18	b	837	CLA	C11-C10-C8-C9
18	b	843	CLA	C11-C10-C8-C9
18	i	101	CLA	C11-C10-C8-C9
18	B	308	CLA	C11-C12-C13-C14
18	C	311	CLA	C11-C10-C8-C9
18	D	207	CLA	C11-C10-C8-C9
18	D	208	CLA	C11-C10-C8-C9
18	H	306	CLA	C11-C12-C13-C14
18	H	307	CLA	C6-C7-C8-C9
18	H	313	CLA	C6-C7-C8-C9
18	H	313	CLA	C14-C13-C15-C16
18	b	815	CLA	O1D-CGD-O2D-CED
29	H	312	KC1	O1D-CGD-O2D-CED
23	l	207	LMG	C2-C1-O1-C7
23	D	202	LMG	C2-C1-O1-C7
23	C	319	LMG	C13-C14-C15-C16
20	H	316	LHG	O2-C2-C3-O3
18	a	824	CLA	O1A-CGA-O2A-C1
18	b	822	CLA	O1D-CGD-O2D-CED
21	a	852	BCR	C11-C12-C13-C35
21	b	833	BCR	C7-C8-C9-C34
21	j	101	BCR	C37-C22-C23-C24
21	m	101	BCR	C7-C8-C9-C34
21	m	101	BCR	C37-C22-C23-C24
26	b	847	A86	C7-C6-C8-C9
26	B	301	A86	C-C1-C24-C25
26	D	203	A86	C-C1-C24-C25
28	C	303	DD6	C12-C11-C13-C14
28	E	303	DD6	C12-C11-C13-C14
28	E	304	DD6	C12-C11-C13-C14

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Mol	Chain	Res	Type	Atoms
28	H	303	DD6	C12-C11-C13-C14
21	a	834	BCR	C21-C22-C23-C24
21	a	852	BCR	C11-C12-C13-C14
21	j	101	BCR	C21-C22-C23-C24
21	m	101	BCR	C21-C22-C23-C24
26	b	847	A86	C5-C6-C8-C9
26	B	301	A86	C2-C1-C24-C25
28	C	303	DD6	C10-C11-C13-C14
28	E	304	DD6	C10-C11-C13-C14
28	H	303	DD6	C10-C11-C13-C14
23	C	301	LMG	C4-C5-C6-O5
18	b	824	CLA	CBD-CGD-O2D-CED
18	C	312	CLA	CBD-CGD-O2D-CED
18	H	314	CLA	CBA-CGA-O2A-C1
23	D	202	LMG	C29-C28-O8-C9
18	a	813	CLA	C10-C11-C12-C13
23	C	319	LMG	C28-C29-C30-C31
23	E	318	LMG	C16-C17-C18-C19
18	a	811	CLA	O1D-CGD-O2D-CED
18	a	828	CLA	O1D-CGD-O2D-CED
18	a	841	CLA	O1D-CGD-O2D-CED
26	H	304	A86	O5-C38-O4-C34
18	a	810	CLA	C15-C16-C17-C18
18	a	813	CLA	C13-C15-C16-C17
18	a	825	CLA	C8-C10-C11-C12
18	a	840	CLA	C15-C16-C17-C18
18	a	841	CLA	C5-C6-C7-C8
20	B	315	LHG	O1-C1-C2-O2
23	D	202	LMG	C28-C29-C30-C31
18	a	817	CLA	CBD-CGD-O2D-CED
29	H	312	KC1	C2C-C3C-CAC-CBC
18	a	855	CLA	C12-C13-C15-C16
18	b	801	CLA	C11-C10-C8-C7
18	b	803	CLA	C12-C13-C15-C16
18	C	309	CLA	C11-C10-C8-C7
18	a	809	CLA	CBA-CGA-O2A-C1
18	a	855	CLA	CBA-CGA-O2A-C1
20	E	301	LHG	C34-C35-C36-C37
28	C	303	DD6	C3-C4-C5-C6
22	a	849	SQD	O5-C1-O6-C44
22	a	849	SQD	O6-C1-O5-C5
20	B	315	LHG	O9-C7-O7-C5

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Mol	Chain	Res	Type	Atoms
18	H	311	CLA	C8-C10-C11-C12
18	C	311	CLA	C10-C11-C12-C13
18	H	313	CLA	C10-C11-C12-C13
20	b	835	LHG	C7-C8-C9-C10
20	B	315	LHG	C7-C8-C9-C10
23	a	851	LMG	C10-C11-C12-C13
23	j	103	LMG	C10-C11-C12-C13
18	a	806	CLA	O1A-CGA-O2A-C1
18	b	801	CLA	O1A-CGA-O2A-C1
20	E	301	LHG	O10-C23-O8-C6
23	l	207	LMG	O10-C28-O8-C9
18	a	827	CLA	CBD-CGD-O2D-CED
18	b	817	CLA	O1D-CGD-O2D-CED
18	a	854	CLA	C5-C6-C7-C8
18	b	803	CLA	C10-C11-C12-C13
18	b	811	CLA	C10-C11-C12-C13
18	b	850	CLA	C13-C15-C16-C17
18	H	306	CLA	C8-C10-C11-C12
18	a	841	CLA	C2A-CAA-CBA-CGA
18	b	808	CLA	C2A-CAA-CBA-CGA
18	b	828	CLA	C2A-CAA-CBA-CGA
18	B	306	CLA	C2A-CAA-CBA-CGA
18	C	310	CLA	C2A-CAA-CBA-CGA
18	D	216	CLA	C2A-CAA-CBA-CGA
18	a	813	CLA	C8-C10-C11-C12
18	a	836	CLA	C8-C10-C11-C12
18	b	824	CLA	C13-C15-C16-C17
18	D	213	CLA	C8-C10-C11-C12
18	E	309	CLA	C5-C6-C7-C8
18	E	312	CLA	C5-C6-C7-C8
18	E	312	CLA	C8-C10-C11-C12
18	H	307	CLA	C8-C10-C11-C12
23	l	207	LMG	C10-C11-C12-C13
23	C	301	LMG	C28-C29-C30-C31
18	E	309	CLA	O1A-CGA-O2A-C1
18	a	844	CLA	O1D-CGD-O2D-CED
18	b	816	CLA	O1D-CGD-O2D-CED
18	a	806	CLA	C8-C10-C11-C12
18	a	813	CLA	C15-C16-C17-C18
18	a	838	CLA	C5-C6-C7-C8
18	a	840	CLA	C13-C15-C16-C17
18	b	807	CLA	C10-C11-C12-C13

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Mol	Chain	Res	Type	Atoms
18	b	823	CLA	C5-C6-C7-C8
18	b	843	CLA	C10-C11-C12-C13
18	b	849	CLA	C13-C15-C16-C17
18	l	203	CLA	C8-C10-C11-C12
18	D	208	CLA	C8-C10-C11-C12
18	E	312	CLA	C15-C16-C17-C18
22	a	849	SQD	C28-C29-C30-C31
20	B	315	LHG	O2-C2-C3-O3
18	b	817	CLA	CBA-CGA-O2A-C1
18	b	842	CLA	C5-C6-C7-C8
18	b	845	CLA	C8-C10-C11-C12
18	C	307	CLA	C13-C15-C16-C17
18	b	830	CLA	O1D-CGD-O2D-CED
18	D	216	CLA	CBA-CGA-O2A-C1
23	C	301	LMG	C11-C12-C13-C14
18	b	837	CLA	O1D-CGD-O2D-CED
18	C	317	CLA	O1D-CGD-O2D-CED
18	a	827	CLA	C13-C15-C16-C17
18	a	836	CLA	C13-C15-C16-C17
18	a	841	CLA	C15-C16-C17-C18
18	b	820	CLA	C13-C15-C16-C17
18	b	829	CLA	C13-C15-C16-C17
18	H	307	CLA	C5-C6-C7-C8
18	H	311	CLA	C5-C6-C7-C8
18	b	823	CLA	CBD-CGD-O2D-CED
23	D	202	LMG	O10-C28-O8-C9
18	a	823	CLA	O1D-CGD-O2D-CED
18	a	842	CLA	O1D-CGD-O2D-CED
18	a	801	CLA	C15-C16-C17-C18
18	a	815	CLA	C5-C6-C7-C8
18	a	843	CLA	C5-C6-C7-C8
18	b	805	CLA	C15-C16-C17-C18
18	b	811	CLA	C8-C10-C11-C12
18	H	313	CLA	C15-C16-C17-C18
18	a	836	CLA	CBA-CGA-O2A-C1
18	b	808	CLA	CBA-CGA-O2A-C1
20	D	201	LHG	C8-C7-O7-C5
29	H	312	KC1	C2A-CAA-CBA-CGA
20	D	201	LHG	C26-C27-C28-C29
18	f	202	CLA	C3-C5-C6-C7
26	B	305	A86	O5-C38-O4-C34
21	a	852	BCR	C9-C10-C11-C12

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Mol	Chain	Res	Type	Atoms
26	B	301	A86	C24-C25-C26-C27
28	C	303	DD6	C24-C25-C26-C27
18	b	817	CLA	C8-C10-C11-C12
20	D	201	LHG	O9-C7-O7-C5
18	b	819	CLA	C2A-CAA-CBA-CGA
18	b	838	CLA	C2A-CAA-CBA-CGA
18	b	830	CLA	CBA-CGA-O2A-C1
18	f	203	CLA	CBA-CGA-O2A-C1
18	i	101	CLA	CBA-CGA-O2A-C1
18	D	213	CLA	CBA-CGA-O2A-C1
18	B	306	CLA	C5-C6-C7-C8
18	C	312	CLA	C10-C11-C12-C13
18	D	207	CLA	C8-C10-C11-C12
18	D	213	CLA	C10-C11-C12-C13
18	E	315	CLA	C5-C6-C7-C8
18	H	306	CLA	C13-C15-C16-C17
18	H	313	CLA	C13-C15-C16-C17
18	b	843	CLA	CBD-CGD-O2D-CED
18	a	855	CLA	O1A-CGA-O2A-C1
18	b	805	CLA	O1D-CGD-O2D-CED
18	a	843	CLA	C15-C16-C17-C18
18	b	803	CLA	C8-C10-C11-C12
18	b	829	CLA	C5-C6-C7-C8
23	C	301	LMG	O6-C5-C6-O5
18	b	820	CLA	C8-C10-C11-C12
18	E	312	CLA	C10-C11-C12-C13
23	D	202	LMG	C35-C36-C37-C38
29	H	312	KC1	CAA-CBA-CGA-O2A
22	a	849	SQD	C23-C24-C25-C26
18	b	827	CLA	CBD-CGD-O2D-CED
18	H	313	CLA	O1D-CGD-O2D-CED
18	a	809	CLA	O1A-CGA-O2A-C1
18	H	314	CLA	O1A-CGA-O2A-C1
18	a	806	CLA	O1D-CGD-O2D-CED
18	b	821	CLA	CBD-CGD-O2D-CED
20	E	301	LHG	C10-C11-C12-C13
20	b	835	LHG	O2-C2-C3-O3
20	D	201	LHG	O2-C2-C3-O3
18	a	826	CLA	CBD-CGD-O2D-CED
20	D	201	LHG	C2-C3-O3-P
21	a	850	BCR	C37-C22-C23-C24
21	b	839	BCR	C7-C8-C9-C34

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Mol	Chain	Res	Type	Atoms
18	l	202	CLA	C2C-C3C-CAC-CBC
21	a	850	BCR	C21-C22-C23-C24
21	b	833	BCR	C7-C8-C9-C10
21	b	839	BCR	C7-C8-C9-C10
18	a	836	CLA	O1A-CGA-O2A-C1
18	a	808	CLA	C2A-CAA-CBA-CGA
18	b	811	CLA	C2A-CAA-CBA-CGA
18	b	842	CLA	C2A-CAA-CBA-CGA
18	D	209	CLA	C2A-CAA-CBA-CGA
20	b	835	LHG	O1-C1-C2-C3
18	b	811	CLA	CBD-CGD-O2D-CED
18	b	850	CLA	O1D-CGD-O2D-CED
18	l	202	CLA	CBA-CGA-O2A-C1
20	H	316	LHG	C6-C5-O7-C7
26	D	204	A86	C1-C2-C3-C4
18	a	801	CLA	C16-C17-C18-C19
18	a	825	CLA	C11-C12-C13-C14
18	b	842	CLA	C16-C17-C18-C20
20	a	831	LHG	C11-C12-C13-C14
18	a	836	CLA	O1D-CGD-O2D-CED
18	H	314	CLA	O1D-CGD-O2D-CED
18	a	827	CLA	C15-C16-C17-C18
23	E	318	LMG	C11-C10-O7-C8
18	C	309	CLA	O1D-CGD-O2D-CED
18	a	822	CLA	O1D-CGD-O2D-CED
18	b	826	CLA	O1D-CGD-O2D-CED
18	a	803	CLA	C10-C11-C12-C13
18	b	845	CLA	C13-C15-C16-C17
18	E	316	CLA	CBD-CGD-O2D-CED
18	a	841	CLA	C2-C1-O2A-CGA
18	a	825	CLA	C11-C12-C13-C15
18	a	854	CLA	C16-C17-C18-C19
18	b	842	CLA	C16-C17-C18-C19
18	C	311	CLA	C11-C12-C13-C15
26	b	847	A86	O5-C38-O4-C34
18	D	213	CLA	O1A-CGA-O2A-C1
20	B	315	LHG	C15-C16-C17-C18
20	b	835	LHG	C14-C15-C16-C17
20	b	835	LHG	C34-C35-C36-C37
20	j	102	LHG	C30-C31-C32-C33
20	D	201	LHG	C18-C19-C20-C21
23	j	103	LMG	C30-C31-C32-C33

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Mol	Chain	Res	Type	Atoms
23	C	319	LMG	C17-C18-C19-C20
18	a	837	CLA	O1D-CGD-O2D-CED
23	E	318	LMG	C10-C11-C12-C13
18	a	801	CLA	C8-C10-C11-C12
20	E	301	LHG	C25-C26-C27-C28
18	b	808	CLA	O1A-CGA-O2A-C1
18	b	817	CLA	O1A-CGA-O2A-C1
18	b	837	CLA	CBA-CGA-O2A-C1
20	E	301	LHG	C17-C18-C19-C20
20	H	316	LHG	C15-C16-C17-C18
23	l	207	LMG	C20-C21-C22-C23
23	C	319	LMG	C20-C21-C22-C23
23	E	318	LMG	C11-C12-C13-C14
25	b	834	DGD	C8A-C9A-CAA-CBA
25	b	834	DGD	CEB-CFB-CGB-CHB
20	b	835	LHG	O1-C1-C2-O2
28	H	303	DD6	C27-C29-C30-C31
20	a	831	LHG	C11-C10-C9-C8
20	b	835	LHG	C26-C27-C28-C29
25	b	834	DGD	C6A-C7A-C8A-C9A
25	b	834	DGD	C6B-C7B-C8B-C9B
18	a	805	CLA	O1D-CGD-O2D-CED
18	a	801	CLA	C4B-C3B-CAB-CBB
18	a	812	CLA	C4B-C3B-CAB-CBB
18	a	814	CLA	C4B-C3B-CAB-CBB
18	a	816	CLA	C4B-C3B-CAB-CBB
18	a	821	CLA	C4B-C3B-CAB-CBB
18	a	826	CLA	C4B-C3B-CAB-CBB
18	b	809	CLA	C4B-C3B-CAB-CBB
18	b	813	CLA	C4B-C3B-CAB-CBB
18	b	827	CLA	C4B-C3B-CAB-CBB
18	b	846	CLA	C4B-C3B-CAB-CBB
18	b	849	CLA	C4B-C3B-CAB-CBB
18	j	104	CLA	C4B-C3B-CAB-CBB
18	B	307	CLA	C4B-C3B-CAB-CBB
18	C	306	CLA	C4B-C3B-CAB-CBB
18	C	311	CLA	C4B-C3B-CAB-CBB
18	C	317	CLA	C4B-C3B-CAB-CBB
18	D	207	CLA	C4B-C3B-CAB-CBB
18	D	211	CLA	C4B-C3B-CAB-CBB
20	b	835	LHG	C23-C24-C25-C26
25	b	834	DGD	C2G-C1G-O1G-C1A

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Mol	Chain	Res	Type	Atoms
18	b	822	CLA	C16-C17-C18-C19
18	E	313	CLA	C6-C7-C8-C9
18	a	823	CLA	C2A-CAA-CBA-CGA
18	b	822	CLA	C2A-CAA-CBA-CGA
18	b	837	CLA	C2A-CAA-CBA-CGA
20	b	835	LHG	C30-C31-C32-C33
18	a	807	CLA	C5-C6-C7-C8
18	b	806	CLA	C13-C15-C16-C17
18	D	208	CLA	C5-C6-C7-C8
20	j	102	LHG	C8-C7-O7-C5
23	C	319	LMG	C11-C10-O7-C8
23	C	319	LMG	C29-C30-C31-C32
18	a	802	CLA	C12-C13-C15-C16
18	b	842	CLA	C6-C7-C8-C10
18	b	845	CLA	C12-C13-C15-C16
18	D	208	CLA	C11-C12-C13-C15
18	H	311	CLA	C6-C7-C8-C10
20	H	316	LHG	C11-C12-C13-C14
18	H	313	CLA	C5-C6-C7-C8
18	i	101	CLA	O1A-CGA-O2A-C1
18	a	806	CLA	C3A-C2A-CAA-CBA
18	a	824	CLA	C3A-C2A-CAA-CBA
18	a	838	CLA	C3A-C2A-CAA-CBA
18	b	808	CLA	C3A-C2A-CAA-CBA
18	b	810	CLA	C3A-C2A-CAA-CBA
18	b	824	CLA	C3A-C2A-CAA-CBA
18	b	825	CLA	C3A-C2A-CAA-CBA
18	b	844	CLA	C3A-C2A-CAA-CBA
18	f	204	CLA	C3A-C2A-CAA-CBA
18	D	209	CLA	C3A-C2A-CAA-CBA
18	D	211	CLA	C3A-C2A-CAA-CBA
18	H	314	CLA	C3A-C2A-CAA-CBA
20	a	831	LHG	C26-C27-C28-C29
19	b	831	PQN	C12-C13-C15-C16
18	a	837	CLA	C5-C6-C7-C8
18	b	837	CLA	C10-C11-C12-C13
18	a	824	CLA	O1D-CGD-O2D-CED
28	E	304	DD6	C24-C25-C26-C27
18	a	840	CLA	C16-C17-C18-C19
18	E	313	CLA	C6-C7-C8-C10
18	b	830	CLA	O1A-CGA-O2A-C1
18	f	203	CLA	O1A-CGA-O2A-C1

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Mol	Chain	Res	Type	Atoms
20	b	835	LHG	C29-C30-C31-C32
20	j	102	LHG	C9-C10-C11-C12
25	b	834	DGD	CAB-CBB-CCB-CDB
18	b	843	CLA	C5-C6-C7-C8
18	b	829	CLA	CBD-CGD-O2D-CED
20	a	831	LHG	C29-C30-C31-C32
20	E	301	LHG	C14-C15-C16-C17
20	b	835	LHG	C1-C2-C3-O3
18	a	855	CLA	O1D-CGD-O2D-CED
18	a	844	CLA	CBA-CGA-O2A-C1
18	B	306	CLA	CBA-CGA-O2A-C1
23	a	851	LMG	C29-C28-O8-C9
20	b	835	LHG	C4-C5-C6-O8
20	E	301	LHG	C4-C5-C6-O8
25	b	834	DGD	O1G-C1G-C2G-C3G
20	E	301	LHG	C29-C30-C31-C32
18	a	827	CLA	C3-C5-C6-C7
18	D	208	CLA	C3-C5-C6-C7
23	a	851	LMG	C28-C29-C30-C31
18	a	807	CLA	C8-C10-C11-C12
20	j	102	LHG	C26-C27-C28-C29
20	B	315	LHG	C10-C11-C12-C13
22	a	849	SQD	C24-C25-C26-C27
23	D	202	LMG	C32-C33-C34-C35
23	E	318	LMG	C33-C34-C35-C36
25	b	834	DGD	C7B-C8B-C9B-CAB
20	B	315	LHG	C11-C10-C9-C8
23	a	851	LMG	C30-C31-C32-C33
18	a	848	CLA	O1D-CGD-O2D-CED
20	a	831	LHG	C15-C16-C17-C18
23	C	319	LMG	C30-C31-C32-C33
18	a	854	CLA	C15-C16-C17-C18
20	a	831	LHG	C31-C32-C33-C34
18	C	311	CLA	C11-C12-C13-C14
18	l	202	CLA	O1A-CGA-O2A-C1
18	a	808	CLA	C2B-C3B-CAB-CBB
18	a	812	CLA	C2B-C3B-CAB-CBB
18	a	814	CLA	C2B-C3B-CAB-CBB
18	a	816	CLA	C2B-C3B-CAB-CBB
18	a	821	CLA	C2B-C3B-CAB-CBB
18	a	826	CLA	C2B-C3B-CAB-CBB
18	b	809	CLA	C2B-C3B-CAB-CBB

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Mol	Chain	Res	Type	Atoms
18	b	813	CLA	C2B-C3B-CAB-CBB
18	b	827	CLA	C2B-C3B-CAB-CBB
18	b	828	CLA	C2B-C3B-CAB-CBB
18	b	846	CLA	C2B-C3B-CAB-CBB
18	b	849	CLA	C2B-C3B-CAB-CBB
18	j	104	CLA	C2B-C3B-CAB-CBB
18	B	310	CLA	C2B-C3B-CAB-CBB
18	C	306	CLA	C2B-C3B-CAB-CBB
18	C	311	CLA	C2B-C3B-CAB-CBB
18	C	312	CLA	C2B-C3B-CAB-CBB
18	C	317	CLA	C2B-C3B-CAB-CBB
18	D	211	CLA	C2B-C3B-CAB-CBB
18	E	310	CLA	C2B-C3B-CAB-CBB
18	E	311	CLA	C2B-C3B-CAB-CBB
21	a	834	BCR	C23-C24-C25-C26
21	a	834	BCR	C23-C24-C25-C30
21	a	852	BCR	C1-C6-C7-C8
21	b	833	BCR	C1-C6-C7-C8
21	b	833	BCR	C5-C6-C7-C8
21	f	201	BCR	C1-C6-C7-C8
21	i	103	BCR	C23-C24-C25-C26
21	i	103	BCR	C23-C24-C25-C30
21	l	201	BCR	C1-C6-C7-C8
21	l	201	BCR	C5-C6-C7-C8
21	l	204	BCR	C23-C24-C25-C26
21	l	204	BCR	C23-C24-C25-C30
21	r	201	BCR	C1-C6-C7-C8
21	E	305	BCR	C23-C24-C25-C26
21	E	305	BCR	C23-C24-C25-C30
27	l	206	ET4	C01-C06-C07-C08
20	E	301	LHG	C9-C10-C11-C12
18	a	801	CLA	CBA-CGA-O2A-C1
18	H	308	CLA	C5-C6-C7-C8
20	B	315	LHG	C26-C27-C28-C29
20	a	831	LHG	C14-C15-C16-C17
18	a	839	CLA	C2A-CAA-CBA-CGA
18	D	210	CLA	C2A-CAA-CBA-CGA
18	b	849	CLA	C15-C16-C17-C18
20	j	102	LHG	C10-C11-C12-C13
18	b	837	CLA	O1A-CGA-O2A-C1
20	B	315	LHG	C28-C29-C30-C31
20	D	201	LHG	C11-C10-C9-C8

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Mol	Chain	Res	Type	Atoms
20	j	102	LHG	O9-C7-O7-C5
23	E	318	LMG	O9-C10-O7-C8
18	b	818	CLA	O1D-CGD-O2D-CED
18	a	840	CLA	C16-C17-C18-C20
20	a	835	LHG	C30-C31-C32-C33
23	a	851	LMG	C11-C12-C13-C14
25	b	834	DGD	C3A-C4A-C5A-C6A
25	b	834	DGD	CDB-CEB-CFB-CGB
18	a	818	CLA	CBA-CGA-O2A-C1
18	a	807	CLA	C14-C13-C15-C16
20	a	832	LHG	C9-C10-C11-C12
20	D	201	LHG	C10-C11-C12-C13
20	a	835	LHG	C31-C32-C33-C34
18	B	308	CLA	C10-C11-C12-C13
18	C	312	CLA	C8-C10-C11-C12
18	D	217	CLA	C2C-C3C-CAC-CBC
20	b	835	LHG	C10-C11-C12-C13
20	D	201	LHG	C28-C29-C30-C31
20	H	316	LHG	C13-C14-C15-C16
20	D	201	LHG	C16-C17-C18-C19
23	C	319	LMG	C31-C32-C33-C34
18	b	846	CLA	CBA-CGA-O2A-C1
20	E	301	LHG	C8-C7-O7-C5
23	C	301	LMG	C11-C10-O7-C8
23	D	202	LMG	C11-C10-O7-C8
18	a	807	CLA	C10-C11-C12-C13
18	H	308	CLA	C8-C10-C11-C12
20	E	301	LHG	O9-C7-O7-C5
23	C	319	LMG	O9-C10-O7-C8
23	D	202	LMG	O9-C10-O7-C8
20	a	831	LHG	C28-C29-C30-C31
22	a	849	SQD	C26-C27-C28-C29
23	j	103	LMG	C31-C32-C33-C34
18	B	306	CLA	O1A-CGA-O2A-C1
18	b	840	CLA	C5-C6-C7-C8
18	H	307	CLA	C13-C15-C16-C17
21	a	852	BCR	C36-C18-C19-C20
21	j	105	BCR	C36-C18-C19-C20
21	l	204	BCR	C7-C8-C9-C34
20	E	301	LHG	C12-C13-C14-C15
20	E	301	LHG	C7-C8-C9-C10
21	l	204	BCR	C7-C8-C9-C10

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Mol	Chain	Res	Type	Atoms
23	l	207	LMG	C29-C30-C31-C32
23	a	851	LMG	C4-C5-C6-O5
18	b	820	CLA	C10-C11-C12-C13
18	b	816	CLA	C2A-CAA-CBA-CGA
18	b	819	CLA	CBD-CGD-O2D-CED
20	a	831	LHG	C13-C14-C15-C16
23	C	319	LMG	C16-C17-C18-C19
18	b	823	CLA	C4-C3-C5-C6
18	B	312	CLA	C4-C3-C5-C6
29	H	312	KC1	C2B-C3B-CAB-CBB
23	j	103	LMG	C14-C15-C16-C17
25	b	834	DGD	C3B-C4B-C5B-C6B
18	a	842	CLA	C2C-C3C-CAC-CBC
20	E	301	LHG	C19-C20-C21-C22
18	a	844	CLA	O1A-CGA-O2A-C1
23	a	851	LMG	O10-C28-O8-C9
18	b	805	CLA	C3-C5-C6-C7
18	a	854	CLA	C10-C11-C12-C13
18	D	209	CLA	CBA-CGA-O2A-C1
18	a	840	CLA	C10-C11-C12-C13
29	C	308	KC1	C4B-C3B-CAB-CBB
29	C	313	KC1	C4B-C3B-CAB-CBB
29	C	313	KC1	C2A-CAA-CBA-CGA
18	b	822	CLA	C16-C17-C18-C20
18	C	307	CLA	C15-C16-C17-C18
18	a	812	CLA	O1D-CGD-O2D-CED
18	b	848	CLA	O1D-CGD-O2D-CED
18	D	208	CLA	O1D-CGD-O2D-CED
29	B	313	KC1	O1D-CGD-O2D-CED
18	a	841	CLA	C3-C5-C6-C7
20	E	301	LHG	O7-C5-C6-O8
23	C	301	LMG	O7-C8-C9-O8
23	l	207	LMG	C13-C14-C15-C16
18	H	306	CLA	CBA-CGA-O2A-C1
23	C	319	LMG	C12-C13-C14-C15
18	l	203	CLA	C13-C15-C16-C17
20	b	835	LHG	C31-C32-C33-C34
18	b	803	CLA	C5-C6-C7-C8
20	a	835	LHG	C2-C3-O3-P
20	H	316	LHG	C11-C10-C9-C8
18	l	202	CLA	C4C-C3C-CAC-CBC
23	E	318	LMG	C13-C14-C15-C16

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Mol	Chain	Res	Type	Atoms
18	b	846	CLA	C4-C3-C5-C6
18	D	213	CLA	C4-C3-C5-C6
23	C	301	LMG	O9-C10-O7-C8
18	a	814	CLA	C2A-CAA-CBA-CGA
18	a	816	CLA	C2A-CAA-CBA-CGA
18	a	854	CLA	C2A-CAA-CBA-CGA
18	b	841	CLA	C2A-CAA-CBA-CGA
18	H	309	CLA	C2A-CAA-CBA-CGA
20	E	301	LHG	C11-C12-C13-C14
18	a	847	CLA	O1D-CGD-O2D-CED
23	l	207	LMG	C18-C19-C20-C21
18	a	801	CLA	O1A-CGA-O2A-C1
18	a	806	CLA	C3-C5-C6-C7
18	a	844	CLA	C3-C5-C6-C7
18	E	315	CLA	C3-C5-C6-C7
20	j	102	LHG	C34-C35-C36-C37
18	a	805	CLA	C1A-C2A-CAA-CBA
18	a	806	CLA	C1A-C2A-CAA-CBA
18	a	814	CLA	C1A-C2A-CAA-CBA
18	b	805	CLA	C1A-C2A-CAA-CBA
18	b	808	CLA	C1A-C2A-CAA-CBA
18	b	810	CLA	C1A-C2A-CAA-CBA
18	b	813	CLA	C1A-C2A-CAA-CBA
18	b	825	CLA	C1A-C2A-CAA-CBA
18	b	827	CLA	C1A-C2A-CAA-CBA
18	b	849	CLA	C1A-C2A-CAA-CBA
18	f	204	CLA	C1A-C2A-CAA-CBA
18	B	312	CLA	C1A-C2A-CAA-CBA
18	C	310	CLA	C1A-C2A-CAA-CBA
18	C	312	CLA	C1A-C2A-CAA-CBA
18	D	209	CLA	C1A-C2A-CAA-CBA
18	D	211	CLA	C1A-C2A-CAA-CBA
18	E	309	CLA	C1A-C2A-CAA-CBA
18	H	314	CLA	C1A-C2A-CAA-CBA
18	a	825	CLA	C5-C6-C7-C8
18	H	314	CLA	C8-C10-C11-C12
20	j	102	LHG	C25-C26-C27-C28
20	D	201	LHG	C34-C35-C36-C37
18	b	817	CLA	C3-C5-C6-C7
18	l	203	CLA	C3-C5-C6-C7
18	a	802	CLA	C6-C7-C8-C10
18	a	805	CLA	C12-C13-C15-C16

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Mol	Chain	Res	Type	Atoms
18	a	806	CLA	C11-C12-C13-C15
18	a	809	CLA	C12-C13-C15-C16
18	a	826	CLA	C11-C12-C13-C15
18	a	837	CLA	C6-C7-C8-C10
18	a	844	CLA	C11-C12-C13-C15
18	a	854	CLA	C11-C10-C8-C7
18	a	855	CLA	C11-C10-C8-C7
18	b	803	CLA	C11-C10-C8-C7
18	b	820	CLA	C11-C12-C13-C15
18	b	822	CLA	C11-C12-C13-C15
18	b	830	CLA	C11-C12-C13-C15
18	b	843	CLA	C11-C10-C8-C7
18	b	850	CLA	C11-C12-C13-C15
18	f	202	CLA	C11-C10-C8-C7
18	f	203	CLA	C12-C13-C15-C16
18	B	308	CLA	C12-C13-C15-C16
18	D	214	CLA	C6-C7-C8-C10
18	H	306	CLA	C12-C13-C15-C16
18	H	307	CLA	C6-C7-C8-C10
18	H	313	CLA	C11-C10-C8-C7
18	H	313	CLA	C12-C13-C15-C16
18	a	853	CLA	C6-C7-C8-C9
18	H	306	CLA	C5-C6-C7-C8
18	a	818	CLA	O1A-CGA-O2A-C1
23	E	318	LMG	C30-C31-C32-C33
18	C	315	CLA	C3A-C2A-CAA-CBA
18	C	317	CLA	C3A-C2A-CAA-CBA
18	a	815	CLA	C15-C16-C17-C18
18	a	841	CLA	C8-C10-C11-C12
18	H	308	CLA	C15-C16-C17-C18
20	a	831	LHG	C33-C34-C35-C36
18	D	208	CLA	C4-C3-C5-C6
18	D	208	CLA	C2-C3-C5-C6
23	C	319	LMG	C32-C33-C34-C35
18	D	209	CLA	O1A-CGA-O2A-C1
18	b	807	CLA	C2A-CAA-CBA-CGA
18	E	312	CLA	C2A-CAA-CBA-CGA
18	E	315	CLA	C2A-CAA-CBA-CGA
18	H	307	CLA	C2A-CAA-CBA-CGA
18	a	803	CLA	C11-C10-C8-C9
18	a	808	CLA	C14-C13-C15-C16
18	a	809	CLA	C14-C13-C15-C16

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Mol	Chain	Res	Type	Atoms
18	a	828	CLA	C14-C13-C15-C16
18	a	837	CLA	C6-C7-C8-C9
18	a	840	CLA	C11-C12-C13-C14
18	a	842	CLA	C14-C13-C15-C16
18	a	854	CLA	C11-C10-C8-C9
18	a	855	CLA	C11-C10-C8-C9
18	b	803	CLA	C11-C10-C8-C9
18	b	805	CLA	C6-C7-C8-C9
18	b	822	CLA	C11-C12-C13-C14
18	b	842	CLA	C11-C12-C13-C14
18	b	850	CLA	C11-C12-C13-C14
18	f	203	CLA	C11-C12-C13-C14
18	B	308	CLA	C14-C13-C15-C16
18	D	208	CLA	C6-C7-C8-C9
18	D	210	CLA	C6-C7-C8-C9
18	D	214	CLA	C6-C7-C8-C9
23	j	103	LMG	C11-C12-C13-C14
18	l	205	CLA	O1D-CGD-O2D-CED
18	E	310	CLA	CBA-CGA-O2A-C1
23	C	301	LMG	C2-C1-O1-C7
18	b	824	CLA	C10-C11-C12-C13
18	C	309	CLA	C5-C6-C7-C8
20	E	301	LHG	C11-C10-C9-C8
20	a	835	LHG	C4-C5-C6-O8
23	a	851	LMG	O1-C7-C8-C9
23	a	851	LMG	C7-C8-C9-O8
23	l	207	LMG	C7-C8-C9-O8
23	D	202	LMG	C7-C8-C9-O8
25	b	834	DGD	C9B-CAB-CBB-CCB
18	b	824	CLA	O1D-CGD-O2D-CED
18	C	312	CLA	O1D-CGD-O2D-CED
18	b	819	CLA	C13-C15-C16-C17
18	B	306	CLA	C8-C10-C11-C12
18	b	819	CLA	CBA-CGA-O2A-C1
18	a	841	CLA	C16-C17-C18-C19
25	b	834	DGD	C8B-C9B-CAB-CBB
18	b	846	CLA	O1A-CGA-O2A-C1
20	B	315	LHG	C24-C25-C26-C27
18	a	817	CLA	O1D-CGD-O2D-CED
23	E	318	LMG	C17-C18-C19-C20
18	a	809	CLA	C4-C3-C5-C6
18	H	311	CLA	C4-C3-C5-C6

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Mol	Chain	Res	Type	Atoms
18	b	846	CLA	C2-C3-C5-C6
18	b	829	CLA	C16-C17-C18-C20
29	H	312	KC1	C4C-C3C-CAC-CBC
18	H	306	CLA	O1A-CGA-O2A-C1
20	a	831	LHG	C16-C17-C18-C19
18	b	814	CLA	C2A-CAA-CBA-CGA
18	D	216	CLA	O1A-CGA-O2A-C1
18	D	207	CLA	C3-C5-C6-C7
23	j	103	LMG	C28-C29-C30-C31
18	a	830	CLA	CBA-CGA-O2A-C1
18	H	307	CLA	CBA-CGA-O2A-C1
20	E	301	LHG	C30-C31-C32-C33
26	D	203	A86	C1-C2-C3-C4
18	a	853	CLA	C6-C7-C8-C10
18	H	308	CLA	C16-C17-C18-C20
18	a	827	CLA	O1D-CGD-O2D-CED
18	b	803	CLA	C3-C5-C6-C7
18	b	811	CLA	C3-C5-C6-C7
18	b	823	CLA	O1D-CGD-O2D-CED
18	b	807	CLA	C5-C6-C7-C8
20	E	301	LHG	C27-C28-C29-C30
23	j	103	LMG	C29-C30-C31-C32
18	a	846	CLA	CBA-CGA-O2A-C1
18	b	840	CLA	C2-C3-C5-C6
23	E	318	LMG	C15-C16-C17-C18
20	a	831	LHG	C34-C35-C36-C37
20	H	316	LHG	C12-C13-C14-C15
23	j	103	LMG	C15-C16-C17-C18
23	C	301	LMG	C31-C32-C33-C34
20	j	102	LHG	C15-C16-C17-C18
23	D	202	LMG	O7-C8-C9-O8
18	a	810	CLA	CBA-CGA-O2A-C1
18	b	840	CLA	CBA-CGA-O2A-C1
23	C	319	LMG	C21-C22-C23-C24
18	b	850	CLA	CAA-CBA-CGA-O2A
23	C	319	LMG	C18-C19-C20-C21
23	D	202	LMG	C30-C31-C32-C33
20	B	315	LHG	C31-C32-C33-C34
18	a	854	CLA	C16-C17-C18-C20
20	j	102	LHG	C12-C13-C14-C15
18	a	820	CLA	CBA-CGA-O2A-C1
20	E	317	LHG	C24-C23-O8-C6

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Mol	Chain	Res	Type	Atoms
22	a	849	SQD	C6-C5-O5-C1
18	H	309	CLA	CBD-CGD-O2D-CED
18	b	824	CLA	C5-C6-C7-C8
20	b	835	LHG	C19-C20-C21-C22
20	D	201	LHG	C35-C36-C37-C38
25	b	834	DGD	C7A-C8A-C9A-CAA
18	H	313	CLA	C16-C17-C18-C20
18	a	842	CLA	C4-C3-C5-C6
18	b	840	CLA	C4-C3-C5-C6
18	E	310	CLA	O1A-CGA-O2A-C1
18	b	838	CLA	C2-C3-C5-C6
18	H	308	CLA	C2-C3-C5-C6
18	a	821	CLA	CBA-CGA-O2A-C1
18	b	810	CLA	CBA-CGA-O2A-C1
18	b	843	CLA	O1D-CGD-O2D-CED
18	a	802	CLA	C6-C7-C8-C9
18	a	805	CLA	C14-C13-C15-C16
18	a	806	CLA	C11-C12-C13-C14
18	a	815	CLA	C6-C7-C8-C9
18	a	826	CLA	C11-C12-C13-C14
18	a	844	CLA	C14-C13-C15-C16
18	a	846	CLA	C11-C10-C8-C9
18	a	855	CLA	C6-C7-C8-C9
18	b	824	CLA	C6-C7-C8-C9
18	b	824	CLA	C11-C12-C13-C14
18	b	843	CLA	C11-C12-C13-C14
18	f	202	CLA	C11-C10-C8-C9
18	f	203	CLA	C14-C13-C15-C16
18	E	310	CLA	C6-C7-C8-C9
18	H	306	CLA	C14-C13-C15-C16
18	b	844	CLA	C5-C6-C7-C8
18	a	822	CLA	C4B-C3B-CAB-CBB
18	a	823	CLA	C4B-C3B-CAB-CBB
18	a	844	CLA	C4B-C3B-CAB-CBB
18	b	803	CLA	C4B-C3B-CAB-CBB
18	b	817	CLA	C4B-C3B-CAB-CBB
18	b	821	CLA	C4B-C3B-CAB-CBB
18	b	823	CLA	C4B-C3B-CAB-CBB
18	b	825	CLA	C4B-C3B-CAB-CBB
18	b	840	CLA	C4B-C3B-CAB-CBB
18	b	848	CLA	C4B-C3B-CAB-CBB
18	f	204	CLA	C4B-C3B-CAB-CBB

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Mol	Chain	Res	Type	Atoms
18	B	312	CLA	C4B-C3B-CAB-CBB
18	C	309	CLA	C4B-C3B-CAB-CBB
18	D	209	CLA	C4B-C3B-CAB-CBB
18	E	315	CLA	C4B-C3B-CAB-CBB
18	H	308	CLA	C4B-C3B-CAB-CBB
18	H	315	CLA	C4B-C3B-CAB-CBB
20	E	301	LHG	C16-C17-C18-C19
18	b	812	CLA	C5-C6-C7-C8
18	b	843	CLA	C3-C5-C6-C7
23	j	103	LMG	C17-C18-C19-C20
18	a	808	CLA	C12-C13-C15-C16
18	a	813	CLA	C12-C13-C15-C16
18	a	815	CLA	C6-C7-C8-C10
18	a	819	CLA	C11-C12-C13-C15
18	a	828	CLA	C12-C13-C15-C16
18	a	840	CLA	C11-C12-C13-C15
18	a	841	CLA	C11-C12-C13-C15
18	a	842	CLA	C12-C13-C15-C16
18	a	844	CLA	C12-C13-C15-C16
18	a	846	CLA	C11-C10-C8-C7
18	a	855	CLA	C6-C7-C8-C10
18	b	807	CLA	C11-C12-C13-C15
18	b	824	CLA	C11-C12-C13-C15
18	b	842	CLA	C11-C12-C13-C15
18	b	843	CLA	C11-C12-C13-C15
18	f	203	CLA	C11-C12-C13-C15
18	l	203	CLA	C6-C7-C8-C10
18	B	308	CLA	C11-C10-C8-C7
18	D	207	CLA	C11-C10-C8-C7
18	D	208	CLA	C11-C10-C8-C7
18	D	210	CLA	C6-C7-C8-C10
18	E	310	CLA	C6-C7-C8-C10
18	H	313	CLA	C6-C7-C8-C10
18	a	847	CLA	C13-C15-C16-C17
18	b	819	CLA	C15-C16-C17-C18
20	E	301	LHG	C13-C14-C15-C16
18	a	813	CLA	C3A-C2A-CAA-CBA
18	a	819	CLA	C4-C3-C5-C6
18	a	839	CLA	C3A-C2A-CAA-CBA
18	b	809	CLA	C3A-C2A-CAA-CBA
18	b	823	CLA	C3A-C2A-CAA-CBA
18	l	202	CLA	C3A-C2A-CAA-CBA

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Mol	Chain	Res	Type	Atoms
18	a	803	CLA	C15-C16-C17-C18
18	a	840	CLA	C2-C3-C5-C6
18	a	842	CLA	C2-C3-C5-C6
18	E	310	CLA	C2-C3-C5-C6
18	b	819	CLA	O1A-CGA-O2A-C1
18	a	826	CLA	O1D-CGD-O2D-CED
18	a	837	CLA	CBA-CGA-O2A-C1
18	b	820	CLA	CBA-CGA-O2A-C1
25	b	834	DGD	C2A-C1A-O1G-C1G
21	f	201	BCR	C19-C20-C21-C22
26	B	301	A86	C11-C10-C9-C8
26	B	304	A86	C3-C4-C5-C6
18	a	801	CLA	C16-C17-C18-C20
18	b	811	CLA	O1D-CGD-O2D-CED
23	l	207	LMG	C14-C15-C16-C17
18	b	812	CLA	C11-C12-C13-C14
18	H	309	CLA	CBA-CGA-O2A-C1
18	b	845	CLA	C10-C11-C12-C13
18	b	849	CLA	CBD-CGD-O2D-CED
18	b	829	CLA	C2A-CAA-CBA-CGA
20	H	316	LHG	C4-C5-C6-O8
18	H	305	CLA	C6-C7-C8-C10
18	b	818	CLA	C16-C17-C18-C19
18	H	313	CLA	C16-C17-C18-C19
18	b	822	CLA	C10-C11-C12-C13
18	b	827	CLA	O1D-CGD-O2D-CED
25	b	834	DGD	C2B-C3B-C4B-C5B
18	a	840	CLA	C4-C3-C5-C6
18	b	820	CLA	C4-C3-C5-C6
18	b	838	CLA	C4-C3-C5-C6
18	E	310	CLA	C4-C3-C5-C6
18	H	308	CLA	C4-C3-C5-C6
20	a	835	LHG	C32-C33-C34-C35
18	a	841	CLA	C16-C17-C18-C20
18	a	843	CLA	C16-C17-C18-C19
18	b	829	CLA	C16-C17-C18-C19
20	H	316	LHG	O6-C4-C5-O7
18	b	821	CLA	O1D-CGD-O2D-CED
21	l	204	BCR	C1-C6-C7-C8
18	B	306	CLA	C14-C13-C15-C16
20	b	835	LHG	C28-C29-C30-C31
20	E	301	LHG	C28-C29-C30-C31

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Mol	Chain	Res	Type	Atoms
18	a	806	CLA	C15-C16-C17-C18
19	b	831	PQN	C18-C20-C21-C22
18	a	805	CLA	C15-C16-C17-C18
18	a	820	CLA	C5-C6-C7-C8
22	a	849	SQD	O47-C45-C46-O48
23	a	851	LMG	O1-C7-C8-O7
20	b	835	LHG	C33-C34-C35-C36
18	a	830	CLA	O1A-CGA-O2A-C1
18	E	316	CLA	O1D-CGD-O2D-CED
18	B	306	CLA	C12-C13-C15-C16
20	D	201	LHG	C19-C20-C21-C22
18	H	307	CLA	O1A-CGA-O2A-C1
18	a	802	CLA	C2C-C3C-CAC-CBC
20	E	301	LHG	C15-C16-C17-C18
18	b	822	CLA	C15-C16-C17-C18
18	b	850	CLA	C5-C6-C7-C8
18	a	819	CLA	C2-C3-C5-C6
18	a	843	CLA	C16-C17-C18-C20
18	b	818	CLA	C16-C17-C18-C20
23	j	103	LMG	C13-C14-C15-C16
18	f	203	CLA	CBD-CGD-O2D-CED
18	a	819	CLA	C11-C12-C13-C14
18	a	838	CLA	C11-C10-C8-C9
18	a	844	CLA	C11-C12-C13-C14
18	b	830	CLA	C11-C12-C13-C14
18	B	308	CLA	C11-C10-C8-C9
18	H	310	CLA	C11-C10-C8-C9
20	a	831	LHG	C12-C13-C14-C15
18	b	820	CLA	C16-C17-C18-C20
18	a	828	CLA	C5-C6-C7-C8
18	b	817	CLA	C2A-CAA-CBA-CGA
20	E	301	LHG	C18-C19-C20-C21
18	b	806	CLA	CBA-CGA-O2A-C1
21	m	101	BCR	C19-C20-C21-C22
20	a	831	LHG	C35-C36-C37-C38
18	i	101	CLA	C11-C12-C13-C14
20	a	831	LHG	C32-C33-C34-C35
18	B	307	CLA	C15-C16-C17-C18
18	a	846	CLA	O1A-CGA-O2A-C1
18	a	843	CLA	C3-C5-C6-C7
18	b	814	CLA	CBA-CGA-O2A-C1
20	D	201	LHG	C24-C23-O8-C6

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Mol	Chain	Res	Type	Atoms
18	a	855	CLA	C8-C10-C11-C12
18	a	838	CLA	C16-C17-C18-C20
18	H	308	CLA	C16-C17-C18-C19
23	D	202	LMG	C17-C18-C19-C20
18	b	830	CLA	C8-C10-C11-C12
21	b	833	BCR	C37-C22-C23-C24
18	a	813	CLA	C11-C12-C13-C15
18	a	820	CLA	C11-C12-C13-C15
18	a	836	CLA	C12-C13-C15-C16
18	a	838	CLA	C11-C10-C8-C7
18	b	805	CLA	C11-C10-C8-C7
18	b	806	CLA	C11-C12-C13-C15
18	b	812	CLA	C6-C7-C8-C10
18	b	824	CLA	C6-C7-C8-C10
18	f	202	CLA	C12-C13-C15-C16
18	B	306	CLA	C6-C7-C8-C10
18	B	311	CLA	C6-C7-C8-C10
18	E	309	CLA	C12-C13-C15-C16
18	H	310	CLA	C11-C10-C8-C7
20	b	835	LHG	C12-C13-C14-C15
20	j	102	LHG	C14-C15-C16-C17
18	b	850	CLA	C10-C11-C12-C13
18	C	311	CLA	C5-C6-C7-C8
21	j	105	BCR	C17-C18-C19-C20
18	a	810	CLA	O1A-CGA-O2A-C1
18	a	821	CLA	O1A-CGA-O2A-C1
18	a	854	CLA	C8-C10-C11-C12
18	b	804	CLA	C2A-CAA-CBA-CGA
18	b	845	CLA	C2A-CAA-CBA-CGA
18	a	848	CLA	C11-C10-C8-C9
18	a	820	CLA	O1A-CGA-O2A-C1
18	a	837	CLA	O1A-CGA-O2A-C1
18	b	810	CLA	O1A-CGA-O2A-C1
18	b	820	CLA	O1A-CGA-O2A-C1
18	b	840	CLA	O1A-CGA-O2A-C1
18	a	813	CLA	C3-C5-C6-C7
18	b	842	CLA	C3-C5-C6-C7
18	b	849	CLA	C3-C5-C6-C7
18	a	825	CLA	CBD-CGD-O2D-CED
18	b	820	CLA	O2A-C1-C2-C3
18	H	306	CLA	C16-C17-C18-C19
25	b	834	DGD	O1A-C1A-O1G-C1G

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Mol	Chain	Res	Type	Atoms
18	a	826	CLA	C3-C5-C6-C7
25	b	834	DGD	C4B-C5B-C6B-C7B
20	a	835	LHG	O6-C4-C5-O7
18	a	817	CLA	O2A-C1-C2-C3
23	E	318	LMG	O6-C1-O1-C7
18	b	819	CLA	O1D-CGD-O2D-CED
20	a	832	LHG	C4-C5-C6-O8
23	C	301	LMG	C7-C8-C9-O8
23	E	318	LMG	O1-C7-C8-C9
23	E	318	LMG	C7-C8-C9-O8
20	a	832	LHG	C10-C11-C12-C13
18	a	809	CLA	C10-C11-C12-C13
18	a	821	CLA	C5-C6-C7-C8
29	B	313	KC1	C4B-C3B-CAB-CBB
29	H	312	KC1	C4B-C3B-CAB-CBB
18	b	806	CLA	O1A-CGA-O2A-C1
18	b	828	CLA	C13-C15-C16-C17
18	a	817	CLA	C2A-CAA-CBA-CGA
18	b	823	CLA	C2-C3-C5-C6
18	b	801	CLA	C5-C6-C7-C8
20	H	316	LHG	O7-C5-C6-O8
23	E	318	LMG	O1-C7-C8-O7
18	a	843	CLA	C11-C12-C13-C14
18	b	807	CLA	C6-C7-C8-C9
18	l	203	CLA	C6-C7-C8-C9
18	H	306	CLA	C6-C7-C8-C9
18	b	820	CLA	C16-C17-C18-C19
23	a	851	LMG	O6-C5-C6-O5
20	D	201	LHG	O10-C23-O8-C6
18	b	828	CLA	C5-C6-C7-C8
26	D	204	A86	C13-C14-C15-O1
26	H	301	A86	C13-C14-C15-O1
22	a	849	SQD	C31-C32-C33-C34
20	D	201	LHG	C29-C30-C31-C32
18	b	843	CLA	C15-C16-C17-C18
20	E	317	LHG	O10-C23-O8-C6
18	a	848	CLA	C3-C5-C6-C7
18	a	807	CLA	C16-C17-C18-C20
18	b	814	CLA	C16-C17-C18-C20
18	B	307	CLA	C16-C17-C18-C20
18	H	310	CLA	C11-C12-C13-C15
18	a	803	CLA	C13-C15-C16-C17

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Mol	Chain	Res	Type	Atoms
18	H	306	CLA	C2A-CAA-CBA-CGA
18	b	828	CLA	C16-C17-C18-C20
18	H	306	CLA	C16-C17-C18-C20
20	j	102	LHG	C31-C32-C33-C34
18	b	814	CLA	O1A-CGA-O2A-C1
18	b	845	CLA	C3-C5-C6-C7
18	D	214	CLA	C11-C10-C8-C9
18	b	843	CLA	C16-C17-C18-C19
18	a	808	CLA	C15-C16-C17-C18
18	B	312	CLA	C8-C10-C11-C12
18	H	309	CLA	CAA-CBA-CGA-O2A
18	l	202	CLA	O2A-C1-C2-C3
18	a	828	CLA	C1A-C2A-CAA-CBA
18	b	848	CLA	C1A-C2A-CAA-CBA
18	B	311	CLA	C4B-C3B-CAB-CBB
18	C	315	CLA	C4B-C3B-CAB-CBB
18	C	318	CLA	C4B-C3B-CAB-CBB
18	b	829	CLA	O1D-CGD-O2D-CED
18	b	849	CLA	O1D-CGD-O2D-CED
18	b	810	CLA	C16-C17-C18-C20
20	D	201	LHG	C33-C34-C35-C36
20	a	831	LHG	O6-C4-C5-C6
20	E	301	LHG	O6-C4-C5-C6
18	a	824	CLA	C2-C3-C5-C6
18	a	814	CLA	C11-C10-C8-C7
18	a	838	CLA	C12-C13-C15-C16
18	a	843	CLA	C11-C12-C13-C15
18	b	814	CLA	C11-C12-C13-C15
18	B	308	CLA	C11-C12-C13-C15
18	B	309	CLA	C12-C13-C15-C16
18	B	312	CLA	C12-C13-C15-C16
18	C	309	CLA	C6-C7-C8-C10
20	H	316	LHG	C14-C15-C16-C17
18	a	838	CLA	C16-C17-C18-C19
18	b	828	CLA	C16-C17-C18-C19
18	C	307	CLA	C16-C17-C18-C19
20	b	835	LHG	C11-C10-C9-C8
18	i	101	CLA	CBD-CGD-O2D-CED
18	C	314	CLA	C2C-C3C-CAC-CBC
18	a	821	CLA	C4-C3-C5-C6
20	a	831	LHG	O6-C4-C5-O7
20	E	301	LHG	O6-C4-C5-O7

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Mol	Chain	Res	Type	Atoms
18	H	309	CLA	O1D-CGD-O2D-CED
18	a	827	CLA	C2A-CAA-CBA-CGA
18	a	820	CLA	C11-C12-C13-C14
18	a	836	CLA	C14-C13-C15-C16
18	b	805	CLA	C11-C10-C8-C9
18	b	806	CLA	C11-C12-C13-C14
18	b	812	CLA	C6-C7-C8-C9
18	b	824	CLA	C11-C10-C8-C9
18	f	202	CLA	C14-C13-C15-C16
18	B	306	CLA	C6-C7-C8-C9
18	B	311	CLA	C6-C7-C8-C9
18	E	309	CLA	C14-C13-C15-C16
21	j	105	BCR	C19-C20-C21-C22
18	a	817	CLA	C1-C2-C3-C4
18	D	209	CLA	C1-C2-C3-C4
29	C	313	KC1	C1A-C2A-CAA-CBA
18	B	308	CLA	C3-C5-C6-C7
18	b	846	CLA	C15-C16-C17-C18
20	a	832	LHG	O7-C5-C6-O8
20	b	835	LHG	O7-C5-C6-O8
22	a	849	SQD	O6-C44-C45-O47
25	b	834	DGD	O1G-C1G-C2G-O2G
18	a	801	CLA	C5-C6-C7-C8
20	B	315	LHG	C30-C31-C32-C33
22	a	849	SQD	C44-C45-C46-O48
18	a	813	CLA	C5-C6-C7-C8
18	a	809	CLA	C2-C3-C5-C6
18	B	312	CLA	C2-C3-C5-C6
18	D	213	CLA	C2-C3-C5-C6
18	a	816	CLA	CBD-CGD-O2D-CED
18	a	824	CLA	CAD-CBD-CGD-O2D
18	r	202	CLA	CAD-CBD-CGD-O2D
18	B	309	CLA	CAD-CBD-CGD-O2D
18	C	316	CLA	CAD-CBD-CGD-O2D
18	H	314	CLA	CAD-CBD-CGD-O2D
26	b	847	A86	C13-C14-C15-C20
18	a	836	CLA	C5-C6-C7-C8
18	f	203	CLA	O1D-CGD-O2D-CED
18	b	828	CLA	CBA-CGA-O2A-C1
18	a	841	CLA	C10-C11-C12-C13
18	b	817	CLA	C11-C12-C13-C15
18	i	101	CLA	C11-C12-C13-C15

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Mol	Chain	Res	Type	Atoms
18	C	311	CLA	C3-C5-C6-C7
18	H	310	CLA	C3-C5-C6-C7
20	D	201	LHG	C9-C10-C11-C12
23	C	301	LMG	C29-C30-C31-C32
18	a	809	CLA	CAD-CBD-CGD-O1D
18	a	816	CLA	CHA-CBD-CGD-O1D
18	a	816	CLA	CHA-CBD-CGD-O2D
18	a	821	CLA	CHA-CBD-CGD-O1D
18	a	821	CLA	CHA-CBD-CGD-O2D
18	a	824	CLA	CAD-CBD-CGD-O1D
18	b	812	CLA	CHA-CBD-CGD-O1D
18	b	812	CLA	CHA-CBD-CGD-O2D
18	b	815	CLA	CHA-CBD-CGD-O1D
18	b	815	CLA	CHA-CBD-CGD-O2D
18	b	816	CLA	CHA-CBD-CGD-O1D
18	b	838	CLA	CHA-CBD-CGD-O1D
18	b	838	CLA	CHA-CBD-CGD-O2D
18	r	202	CLA	CAD-CBD-CGD-O1D
18	B	309	CLA	CAD-CBD-CGD-O1D
18	C	309	CLA	CHA-CBD-CGD-O2D
18	C	316	CLA	CAD-CBD-CGD-O1D
18	H	307	CLA	CHA-CBD-CGD-O1D
18	H	307	CLA	CHA-CBD-CGD-O2D
18	H	314	CLA	CAD-CBD-CGD-O1D
20	a	832	LHG	C3-O3-P-O5
20	a	832	LHG	C4-O6-P-O3
20	a	835	LHG	C4-O6-P-O5
20	b	835	LHG	C3-O3-P-O4
20	b	835	LHG	C3-O3-P-O6
20	B	315	LHG	C4-O6-P-O3
20	B	315	LHG	C4-O6-P-O5
20	D	201	LHG	C4-O6-P-O3
20	D	201	LHG	C4-O6-P-O4
20	D	201	LHG	C4-O6-P-O5
20	E	301	LHG	C3-O3-P-O5
20	E	301	LHG	C3-O3-P-O6
20	E	301	LHG	C4-O6-P-O3
20	E	301	LHG	C4-O6-P-O5
21	a	852	BCR	C19-C20-C21-C22
26	B	302	A86	C3-C4-C5-C6
26	H	302	A86	C3-C4-C5-C6
18	D	211	CLA	CBA-CGA-O2A-C1

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Mol	Chain	Res	Type	Atoms
18	a	801	CLA	C2B-C3B-CAB-CBB
18	B	307	CLA	C2B-C3B-CAB-CBB
18	B	311	CLA	C2B-C3B-CAB-CBB
18	C	318	CLA	C2B-C3B-CAB-CBB
18	D	207	CLA	C2B-C3B-CAB-CBB
20	a	835	LHG	C5-C4-O6-P
18	r	202	CLA	C2C-C3C-CAC-CBC
18	b	828	CLA	O1A-CGA-O2A-C1
18	a	848	CLA	C11-C10-C8-C7
20	b	835	LHG	C6-C5-O7-C7
26	b	847	A86	C1-C24-C25-C26
26	m	102	A86	C6-C8-C9-C10
26	r	203	A86	C6-C8-C9-C10
26	B	301	A86	C6-C8-C9-C10
26	B	302	A86	C6-C8-C9-C10
26	B	304	A86	C6-C8-C9-C10
26	B	305	A86	C6-C8-C9-C10
26	C	302	A86	C6-C8-C9-C10
26	C	305	A86	C6-C8-C9-C10
26	D	203	A86	C1-C24-C25-C26
26	D	203	A86	C6-C8-C9-C10
26	D	206	A86	C6-C8-C9-C10
26	E	302	A86	C6-C8-C9-C10
26	H	301	A86	C6-C8-C9-C10
26	H	302	A86	C6-C8-C9-C10
26	H	304	A86	C6-C8-C9-C10
28	D	205	DD6	C1-C24-C25-C26
28	E	307	DD6	C6-C8-C9-C10
18	a	810	CLA	C8-C10-C11-C12
18	b	814	CLA	C16-C17-C18-C19
18	B	307	CLA	C16-C17-C18-C19
25	b	834	DGD	C4E-C5E-C6E-O5E
20	a	835	LHG	O6-C4-C5-C6
18	D	208	CLA	C13-C15-C16-C17
18	a	840	CLA	C14-C13-C15-C16
18	a	855	CLA	C14-C13-C15-C16
18	b	803	CLA	C14-C13-C15-C16
18	b	807	CLA	C11-C12-C13-C14
18	b	845	CLA	C14-C13-C15-C16
18	B	309	CLA	C6-C7-C8-C9
18	C	309	CLA	C6-C7-C8-C9
18	D	208	CLA	C14-C13-C15-C16

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Mol	Chain	Res	Type	Atoms
18	H	314	CLA	C6-C7-C8-C9
18	a	813	CLA	C6-C7-C8-C10
18	b	803	CLA	C11-C12-C13-C15
18	b	823	CLA	C11-C10-C8-C7
23	D	202	LMG	C11-C12-C13-C14
20	E	317	LHG	O6-C4-C5-O7
18	C	312	CLA	O1A-CGA-O2A-C1
18	H	313	CLA	C4-C3-C5-C6
18	H	311	CLA	C2-C3-C5-C6
18	b	810	CLA	C16-C17-C18-C19
18	b	846	CLA	C5-C6-C7-C8
23	C	319	LMG	C34-C35-C36-C37
20	a	835	LHG	O7-C5-C6-O8
23	C	301	LMG	O1-C7-C8-O7
18	b	846	CLA	CAA-CBA-CGA-O2A
18	a	825	CLA	C2A-CAA-CBA-CGA
18	b	818	CLA	C13-C15-C16-C17
18	b	811	CLA	CBA-CGA-O2A-C1
18	a	813	CLA	C2-C1-O2A-CGA
18	i	101	CLA	C2-C1-O2A-CGA
20	B	315	LHG	C9-C10-C11-C12
23	C	301	LMG	O1-C7-C8-C9
18	a	826	CLA	C10-C11-C12-C13
23	j	103	LMG	C8-C7-O1-C1
20	b	835	LHG	C35-C36-C37-C38
18	a	803	CLA	C2A-CAA-CBA-CGA
18	a	844	CLA	C2A-CAA-CBA-CGA
18	C	307	CLA	C2A-CAA-CBA-CGA
18	H	308	CLA	C2A-CAA-CBA-CGA
28	E	304	DD6	C1-C2-C3-C4
18	a	844	CLA	C4-C3-C5-C6
18	b	805	CLA	C4-C3-C5-C6
18	b	808	CLA	C15-C16-C17-C18
18	a	821	CLA	C2-C3-C5-C6
18	a	815	CLA	C10-C11-C12-C13
18	a	825	CLA	O1D-CGD-O2D-CED
20	a	835	LHG	C25-C26-C27-C28
26	r	203	A86	C13-C14-C15-C16
18	a	814	CLA	C11-C10-C8-C9
18	a	838	CLA	C14-C13-C15-C16
18	a	840	CLA	C11-C10-C8-C9
18	b	814	CLA	C11-C12-C13-C14

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Mol	Chain	Res	Type	Atoms
18	B	309	CLA	C14-C13-C15-C16
18	B	312	CLA	C14-C13-C15-C16
18	i	101	CLA	O1D-CGD-O2D-CED
18	b	811	CLA	O1A-CGA-O2A-C1
18	a	809	CLA	C4B-C3B-CAB-CBB
18	a	839	CLA	C4B-C3B-CAB-CBB
18	a	853	CLA	C4B-C3B-CAB-CBB
18	b	842	CLA	C4B-C3B-CAB-CBB
18	f	203	CLA	C4B-C3B-CAB-CBB
18	D	208	CLA	C4B-C3B-CAB-CBB
18	H	309	CLA	C4B-C3B-CAB-CBB
20	a	831	LHG	C27-C28-C29-C30
18	C	312	CLA	CBA-CGA-O2A-C1
18	H	309	CLA	O1A-CGA-O2A-C1
18	H	313	CLA	C2-C3-C5-C6
25	b	834	DGD	CBB-CCB-CDB-CEB
18	a	801	CLA	C11-C12-C13-C15
18	a	854	CLA	C6-C7-C8-C10
18	b	817	CLA	C11-C10-C8-C7
18	B	306	CLA	C11-C12-C13-C15
18	E	312	CLA	C12-C13-C15-C16
18	a	816	CLA	O1D-CGD-O2D-CED
18	C	311	CLA	C2C-C3C-CAC-CBC
21	f	205	BCR	C35-C13-C14-C15
21	l	201	BCR	C11-C10-C9-C34
21	m	101	BCR	C11-C10-C9-C34
26	b	847	A86	C-C1-C2-C3
26	m	102	A86	C-C1-C2-C3
26	m	102	A86	C4-C5-C6-C7
26	D	204	A86	C4-C5-C6-C7
28	D	205	DD6	C4-C5-C6-C7
28	E	307	DD6	C9-C10-C11-C12
28	E	307	DD6	C4-C5-C6-C7
18	a	807	CLA	C13-C15-C16-C17
18	a	837	CLA	C2-C1-O2A-CGA
18	b	841	CLA	C2-C1-O2A-CGA
18	b	845	CLA	C2-C1-O2A-CGA
18	D	207	CLA	C2-C1-O2A-CGA
18	H	305	CLA	C2-C1-O2A-CGA
20	a	831	LHG	C25-C26-C27-C28
18	l	203	CLA	C15-C16-C17-C18
21	b	833	BCR	C21-C22-C23-C24

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Mol	Chain	Res	Type	Atoms
18	b	801	CLA	C4-C3-C5-C6
18	b	837	CLA	C4-C3-C5-C6
20	D	201	LHG	C32-C33-C34-C35
18	H	310	CLA	C2C-C3C-CAC-CBC
18	D	209	CLA	O2A-C1-C2-C3
18	f	204	CLA	C2A-CAA-CBA-CGA
18	B	307	CLA	C2A-CAA-CBA-CGA
23	E	318	LMG	C28-C29-C30-C31
18	b	850	CLA	CAA-CBA-CGA-O1A
18	B	309	CLA	C15-C16-C17-C18
18	E	310	CLA	C10-C11-C12-C13
18	b	801	CLA	C6-C7-C8-C9
18	b	806	CLA	C11-C10-C8-C9
18	b	850	CLA	C6-C7-C8-C9
18	B	306	CLA	C11-C12-C13-C14
18	C	307	CLA	C6-C7-C8-C9
18	C	312	CLA	C6-C7-C8-C9
18	E	310	CLA	C14-C13-C15-C16
18	D	217	CLA	C4C-C3C-CAC-CBC
20	a	831	LHG	C9-C10-C11-C12
18	B	312	CLA	C2A-CAA-CBA-CGA
29	H	312	KC1	CAA-CBA-CGA-O1A
18	a	808	CLA	C1A-C2A-CAA-CBA
18	a	818	CLA	C1A-C2A-CAA-CBA
18	a	836	CLA	C1A-C2A-CAA-CBA
18	a	839	CLA	C1A-C2A-CAA-CBA
18	l	202	CLA	C1A-C2A-CAA-CBA
18	E	311	CLA	C1A-C2A-CAA-CBA
21	f	205	BCR	C12-C13-C14-C15
21	l	201	BCR	C11-C10-C9-C8
21	m	101	BCR	C11-C10-C9-C8
26	b	847	A86	C24-C1-C2-C3
26	m	102	A86	C24-C1-C2-C3
26	m	102	A86	C4-C5-C6-C8
26	D	204	A86	C4-C5-C6-C8
28	D	205	DD6	C4-C5-C6-C8
28	E	307	DD6	C9-C10-C11-C13
28	E	307	DD6	C4-C5-C6-C8
18	E	310	CLA	C15-C16-C17-C18
18	a	809	CLA	C2B-C3B-CAB-CBB
18	a	822	CLA	C2B-C3B-CAB-CBB
18	a	828	CLA	C2B-C3B-CAB-CBB

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Mol	Chain	Res	Type	Atoms
18	a	839	CLA	C2B-C3B-CAB-CBB
18	a	853	CLA	C2B-C3B-CAB-CBB
18	b	816	CLA	C2B-C3B-CAB-CBB
18	b	821	CLA	C2B-C3B-CAB-CBB
18	b	840	CLA	C2B-C3B-CAB-CBB
18	f	203	CLA	C2B-C3B-CAB-CBB
18	C	315	CLA	C2B-C3B-CAB-CBB
18	D	209	CLA	C2B-C3B-CAB-CBB
18	D	213	CLA	C2B-C3B-CAB-CBB
18	D	215	CLA	C2B-C3B-CAB-CBB
18	E	315	CLA	C2B-C3B-CAB-CBB
18	H	308	CLA	C2B-C3B-CAB-CBB
18	H	309	CLA	C2B-C3B-CAB-CBB
21	l	204	BCR	C5-C6-C7-C8
21	r	201	BCR	C5-C6-C7-C8
21	r	201	BCR	C23-C24-C25-C30
25	b	834	DGD	C1B-C2B-C3B-C4B
18	a	809	CLA	C3-C5-C6-C7
18	b	842	CLA	C10-C11-C12-C13
18	a	837	CLA	C4-C3-C5-C6
18	b	803	CLA	C4-C3-C5-C6
20	H	316	LHG	O6-C4-C5-C6
23	a	851	LMG	O9-C10-O7-C8
18	a	803	CLA	C11-C12-C13-C15
18	a	825	CLA	C6-C7-C8-C10
18	a	826	CLA	C11-C10-C8-C7
18	a	840	CLA	C11-C10-C8-C7
18	b	826	CLA	C11-C10-C8-C7
18	b	837	CLA	C11-C10-C8-C7
18	l	203	CLA	C11-C10-C8-C7
18	C	311	CLA	C11-C10-C8-C7
18	b	850	CLA	C16-C17-C18-C19
18	a	810	CLA	C2A-CAA-CBA-CGA
18	a	813	CLA	C2A-CAA-CBA-CGA
23	j	103	LMG	O7-C8-C9-O8
18	a	808	CLA	C8-C10-C11-C12
18	b	849	CLA	C10-C11-C12-C13
23	E	318	LMG	C18-C19-C20-C21
18	b	844	CLA	C3-C5-C6-C7
21	f	201	BCR	C7-C8-C9-C34
18	B	306	CLA	C4-C3-C5-C6
18	B	308	CLA	C4-C3-C5-C6

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Mol	Chain	Res	Type	Atoms
18	b	805	CLA	C2-C3-C5-C6
18	b	820	CLA	C2-C3-C5-C6
18	b	837	CLA	C2-C3-C5-C6
25	b	834	DGD	C2A-C3A-C4A-C5A
20	j	102	LHG	C16-C17-C18-C19
18	l	203	CLA	C16-C17-C18-C19
18	r	202	CLA	C4C-C3C-CAC-CBC
23	D	202	LMG	C20-C21-C22-C23
18	a	801	CLA	C11-C12-C13-C14
18	a	815	CLA	C11-C12-C13-C14
18	b	829	CLA	C8-C10-C11-C12
27	l	206	ET4	C19-C20-C21-C22
18	b	828	CLA	C15-C16-C17-C18
20	b	835	LHG	C5-C4-O6-P
18	a	814	CLA	CAA-CBA-CGA-O2A
18	b	801	CLA	C2-C3-C5-C6
18	b	837	CLA	C8-C10-C11-C12
18	E	312	CLA	C13-C15-C16-C17
18	b	812	CLA	C3-C5-C6-C7
25	b	834	DGD	C9A-CAA-CBA-CCA
18	a	854	CLA	C2C-C3C-CAC-CBC
18	b	844	CLA	CBD-CGD-O2D-CED
18	a	840	CLA	C8-C10-C11-C12
25	b	834	DGD	C2B-C1B-O2G-C2G
18	C	311	CLA	C4C-C3C-CAC-CBC
20	a	835	LHG	C23-C24-C25-C26
23	C	319	LMG	C15-C16-C17-C18
18	a	823	CLA	CAA-CBA-CGA-O2A
18	H	315	CLA	CAA-CBA-CGA-O2A
18	b	821	CLA	C2A-CAA-CBA-CGA
18	b	824	CLA	C2C-C3C-CAC-CBC
25	b	834	DGD	O1B-C1B-O2G-C2G
28	E	308	DD6	C27-C29-C30-C31
18	C	309	CLA	C4-C3-C5-C6
20	a	831	LHG	C17-C18-C19-C20
18	C	306	CLA	CAA-CBA-CGA-O2A
18	a	801	CLA	CAA-CBA-CGA-O2A
18	a	810	CLA	C4B-C3B-CAB-CBB
18	a	828	CLA	C4B-C3B-CAB-CBB
18	b	805	CLA	C4B-C3B-CAB-CBB
18	b	808	CLA	C4B-C3B-CAB-CBB
18	b	830	CLA	C4B-C3B-CAB-CBB

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Mol	Chain	Res	Type	Atoms
18	l	203	CLA	C4B-C3B-CAB-CBB
18	D	213	CLA	C4B-C3B-CAB-CBB
18	D	215	CLA	C4B-C3B-CAB-CBB
18	b	840	CLA	O1D-CGD-O2D-CED
18	H	307	CLA	C10-C11-C12-C13
18	i	101	CLA	C10-C11-C12-C13
18	B	311	CLA	C13-C15-C16-C17
18	C	309	CLA	C4C-C3C-CAC-CBC
23	a	851	LMG	C11-C10-O7-C8
23	l	207	LMG	C12-C13-C14-C15
20	B	315	LHG	C13-C14-C15-C16
23	a	851	LMG	C32-C33-C34-C35
18	a	847	CLA	C11-C10-C8-C9
18	b	806	CLA	C14-C13-C15-C16
18	b	843	CLA	C14-C13-C15-C16
18	E	309	CLA	C6-C7-C8-C9
18	a	805	CLA	CAA-CBA-CGA-O2A
18	a	827	CLA	C5-C6-C7-C8
18	B	309	CLA	C2A-CAA-CBA-CGA
18	E	309	CLA	C2-C1-O2A-CGA
18	b	846	CLA	C16-C17-C18-C20
18	H	310	CLA	C10-C11-C12-C13
18	a	855	CLA	C3A-C2A-CAA-CBA
18	b	843	CLA	C3A-C2A-CAA-CBA
18	b	849	CLA	C3A-C2A-CAA-CBA
18	E	313	CLA	C3A-C2A-CAA-CBA
18	E	316	CLA	C3A-C2A-CAA-CBA
18	a	812	CLA	CAA-CBA-CGA-O2A
18	a	823	CLA	CAA-CBA-CGA-O1A
20	B	315	LHG	C23-C24-C25-C26
18	b	848	CLA	C5-C6-C7-C8
23	E	318	LMG	C31-C32-C33-C34
18	b	822	CLA	O1A-CGA-O2A-C1
18	H	315	CLA	CAA-CBA-CGA-O1A
18	b	810	CLA	C15-C16-C17-C18
18	a	843	CLA	C13-C15-C16-C17
20	j	102	LHG	C18-C19-C20-C21
18	H	310	CLA	C11-C12-C13-C14
18	a	818	CLA	C2C-C3C-CAC-CBC
18	b	806	CLA	C2A-CAA-CBA-CGA
20	a	832	LHG	O2-C2-C3-O3
18	b	806	CLA	C10-C11-C12-C13

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Mol	Chain	Res	Type	Atoms
18	b	807	CLA	C16-C17-C18-C19
28	E	303	DD6	C13-C14-C15-O1
18	C	306	CLA	CAA-CBA-CGA-O1A
18	E	314	CLA	O1D-CGD-O2D-CED
20	a	831	LHG	C30-C31-C32-C33
18	b	822	CLA	CBA-CGA-O2A-C1
18	b	840	CLA	CBD-CGD-O2D-CED
18	D	214	CLA	O1A-CGA-O2A-C1
20	H	316	LHG	C17-C18-C19-C20
18	b	826	CLA	C2A-CAA-CBA-CGA
26	D	206	A86	C3-C4-C5-C6
26	b	847	A86	C28-C27-C29-C30
22	a	849	SQD	C25-C26-C27-C28
18	a	836	CLA	C15-C16-C17-C18
18	a	838	CLA	C13-C15-C16-C17
18	a	812	CLA	CAA-CBA-CGA-O1A
18	D	214	CLA	O1D-CGD-O2D-CED
18	D	214	CLA	CBA-CGA-O2A-C1
20	E	317	LHG	C16-C17-C18-C19
18	a	807	CLA	C16-C17-C18-C19
18	b	824	CLA	CAA-CBA-CGA-O2A
20	E	317	LHG	O6-C4-C5-C6
21	a	852	BCR	C17-C18-C19-C20
26	D	206	A86	C13-C14-C15-O1
18	b	824	CLA	C2A-CAA-CBA-CGA
18	a	815	CLA	C11-C12-C13-C15
18	a	847	CLA	C11-C10-C8-C7
18	b	806	CLA	C12-C13-C15-C16
18	b	829	CLA	C11-C10-C8-C7
18	E	309	CLA	C6-C7-C8-C10
18	E	310	CLA	C12-C13-C15-C16
18	D	211	CLA	O1A-CGA-O2A-C1
18	b	844	CLA	O1D-CGD-O2D-CED
18	a	810	CLA	C2B-C3B-CAB-CBB
18	a	817	CLA	C2B-C3B-CAB-CBB
18	a	823	CLA	C2B-C3B-CAB-CBB
18	a	830	CLA	C2B-C3B-CAB-CBB
18	a	841	CLA	C2B-C3B-CAB-CBB
18	a	844	CLA	C2B-C3B-CAB-CBB
18	a	846	CLA	C2B-C3B-CAB-CBB
18	a	847	CLA	C2B-C3B-CAB-CBB
18	a	848	CLA	C2B-C3B-CAB-CBB

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Mol	Chain	Res	Type	Atoms
18	b	803	CLA	C2B-C3B-CAB-CBB
18	b	805	CLA	C2B-C3B-CAB-CBB
18	b	808	CLA	C2B-C3B-CAB-CBB
18	b	817	CLA	C2B-C3B-CAB-CBB
18	b	819	CLA	C2B-C3B-CAB-CBB
18	b	823	CLA	C2B-C3B-CAB-CBB
18	b	825	CLA	C2B-C3B-CAB-CBB
18	b	826	CLA	C2B-C3B-CAB-CBB
18	b	830	CLA	C2B-C3B-CAB-CBB
18	b	838	CLA	C2B-C3B-CAB-CBB
18	b	842	CLA	C2B-C3B-CAB-CBB
18	b	843	CLA	C2B-C3B-CAB-CBB
18	b	845	CLA	C2B-C3B-CAB-CBB
18	b	848	CLA	C2B-C3B-CAB-CBB
18	f	204	CLA	C2B-C3B-CAB-CBB
18	l	203	CLA	C2B-C3B-CAB-CBB
18	B	312	CLA	C2B-C3B-CAB-CBB
18	C	309	CLA	C2B-C3B-CAB-CBB
18	D	208	CLA	C2B-C3B-CAB-CBB
18	D	212	CLA	C2B-C3B-CAB-CBB
18	E	313	CLA	C2B-C3B-CAB-CBB
18	H	315	CLA	C2B-C3B-CAB-CBB
20	a	832	LHG	O8-C23-C24-C25
21	b	839	BCR	C1-C6-C7-C8
21	b	839	BCR	C5-C6-C7-C8
21	r	201	BCR	C23-C24-C25-C26
18	b	806	CLA	C2-C1-O2A-CGA
18	b	829	CLA	C2-C1-O2A-CGA
18	b	838	CLA	C2-C1-O2A-CGA
18	b	846	CLA	C13-C15-C16-C17
20	a	835	LHG	C27-C28-C29-C30
18	a	813	CLA	CAA-CBA-CGA-O2A
18	B	306	CLA	C10-C11-C12-C13
18	b	813	CLA	C2-C3-C5-C6
18	a	826	CLA	C2A-CAA-CBA-CGA
18	b	813	CLA	C2A-CAA-CBA-CGA
22	a	849	SQD	C33-C34-C35-C36
18	b	829	CLA	C2C-C3C-CAC-CBC
18	l	203	CLA	O1A-CGA-O2A-C1
18	B	306	CLA	CAA-CBA-CGA-O2A
18	a	808	CLA	CAA-CBA-CGA-O2A
18	b	817	CLA	CAA-CBA-CGA-O2A

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Mol	Chain	Res	Type	Atoms
18	D	209	CLA	CAA-CBA-CGA-O2A
18	D	211	CLA	CAA-CBA-CGA-O2A
20	a	835	LHG	C29-C30-C31-C32
18	D	214	CLA	C11-C10-C8-C7
18	D	214	CLA	CBD-CGD-O2D-CED
18	b	813	CLA	C4-C3-C5-C6
18	a	844	CLA	C2-C3-C5-C6
18	b	817	CLA	C2-C3-C5-C6
20	D	201	LHG	C17-C18-C19-C20
23	D	202	LMG	C10-C11-C12-C13
18	b	844	CLA	CAA-CBA-CGA-O2A
18	C	309	CLA	C10-C11-C12-C13
18	a	802	CLA	C14-C13-C15-C16
18	a	826	CLA	C11-C10-C8-C9
18	i	101	CLA	C6-C7-C8-C9
18	l	203	CLA	C11-C12-C13-C14
18	a	811	CLA	CAA-CBA-CGA-O2A
20	a	831	LHG	O8-C23-C24-C25
18	b	846	CLA	C16-C17-C18-C19
23	D	202	LMG	C21-C22-C23-C24
18	a	810	CLA	C1A-C2A-CAA-CBA
18	a	817	CLA	C4B-C3B-CAB-CBB
18	a	830	CLA	C4B-C3B-CAB-CBB
18	a	843	CLA	C1A-C2A-CAA-CBA
18	a	846	CLA	C4B-C3B-CAB-CBB
18	a	847	CLA	C4B-C3B-CAB-CBB
18	a	848	CLA	C4B-C3B-CAB-CBB
18	b	807	CLA	C4B-C3B-CAB-CBB
18	b	816	CLA	C4B-C3B-CAB-CBB
18	b	819	CLA	C4B-C3B-CAB-CBB
18	b	838	CLA	C4B-C3B-CAB-CBB
18	b	845	CLA	C4B-C3B-CAB-CBB
18	B	306	CLA	C4B-C3B-CAB-CBB
18	C	314	CLA	C4B-C3B-CAB-CBB
18	D	212	CLA	C4B-C3B-CAB-CBB
18	E	310	CLA	C1A-C2A-CAA-CBA
18	E	313	CLA	C4B-C3B-CAB-CBB
18	E	314	CLA	C4B-C3B-CAB-CBB
26	C	302	A86	O-C13-C14-C15
18	a	817	CLA	CAA-CBA-CGA-O2A
18	b	808	CLA	CAA-CBA-CGA-O2A
20	B	315	LHG	O7-C7-C8-C9

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Mol	Chain	Res	Type	Atoms
18	a	815	CLA	C2-C3-C5-C6
18	a	830	CLA	C2-C3-C5-C6
18	E	314	CLA	CBD-CGD-O2D-CED
20	b	835	LHG	O10-C23-O8-C6
18	l	203	CLA	CBA-CGA-O2A-C1
18	a	842	CLA	C4C-C3C-CAC-CBC
18	b	849	CLA	CAA-CBA-CGA-O2A
18	r	202	CLA	C2A-CAA-CBA-CGA
18	f	203	CLA	C5-C6-C7-C8
18	a	830	CLA	C4-C3-C5-C6
18	b	817	CLA	C4-C3-C5-C6
18	a	803	CLA	C2-C1-O2A-CGA
18	a	842	CLA	C2-C1-O2A-CGA
18	b	824	CLA	C2-C1-O2A-CGA
18	a	805	CLA	C6-C7-C8-C10
18	a	815	CLA	C11-C10-C8-C7
18	b	819	CLA	C11-C10-C8-C7
18	b	843	CLA	C12-C13-C15-C16
18	E	312	CLA	C11-C10-C8-C7
18	a	808	CLA	C5-C6-C7-C8
18	C	309	CLA	C2C-C3C-CAC-CBC
18	b	848	CLA	O1A-CGA-O2A-C1
22	a	849	SQD	O48-C23-C24-C25
20	j	102	LHG	C29-C30-C31-C32
18	b	848	CLA	CBA-CGA-O2A-C1
26	C	304	A86	C24-C25-C26-C27
26	D	203	A86	C3-C4-C5-C6
18	E	309	CLA	C2A-CAA-CBA-CGA
18	b	845	CLA	C16-C17-C18-C20
18	a	802	CLA	C15-C16-C17-C18
18	a	847	CLA	CAA-CBA-CGA-O2A
18	a	839	CLA	C10-C11-C12-C13
18	a	843	CLA	C3A-C2A-CAA-CBA
18	E	311	CLA	C3A-C2A-CAA-CBA
18	D	209	CLA	CAA-CBA-CGA-O1A
18	b	824	CLA	CAA-CBA-CGA-O1A
18	b	807	CLA	C16-C17-C18-C20
18	E	316	CLA	C2A-CAA-CBA-CGA
18	a	805	CLA	CBA-CGA-O2A-C1
20	b	835	LHG	C24-C23-O8-C6
18	r	202	CLA	CAA-CBA-CGA-O2A
18	a	803	CLA	C14-C13-C15-C16

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Mol	Chain	Res	Type	Atoms
18	a	815	CLA	C11-C10-C8-C9
18	a	825	CLA	C6-C7-C8-C9
18	b	819	CLA	C11-C10-C8-C9
18	b	823	CLA	C11-C10-C8-C9
18	b	826	CLA	C11-C10-C8-C9
18	b	842	CLA	C6-C7-C8-C9
18	a	817	CLA	CAA-CBA-CGA-O1A
18	b	817	CLA	CAA-CBA-CGA-O1A
18	D	211	CLA	CAA-CBA-CGA-O1A
18	H	309	CLA	CAA-CBA-CGA-O1A
18	b	814	CLA	C13-C15-C16-C17
18	H	307	CLA	O1D-CGD-O2D-CED
18	b	808	CLA	C13-C15-C16-C17
18	b	819	CLA	C10-C11-C12-C13
18	a	815	CLA	C4-C3-C5-C6
18	b	841	CLA	C4-C3-C5-C6
18	a	808	CLA	CAA-CBA-CGA-O1A
18	b	803	CLA	C2-C3-C5-C6
26	b	847	A86	C26-C27-C29-C30
26	D	203	A86	C26-C27-C29-C30
29	C	308	KC1	C1A-C2A-CAA-CBA
18	r	202	CLA	CAA-CBA-CGA-O1A
20	a	831	LHG	O10-C23-C24-C25
18	a	816	CLA	CAA-CBA-CGA-O2A
20	a	832	LHG	O10-C23-C24-C25
23	C	319	LMG	O7-C8-C9-O8
18	a	811	CLA	CAA-CBA-CGA-O1A
18	B	306	CLA	CAA-CBA-CGA-O1A
20	B	315	LHG	O9-C7-C8-C9
18	D	214	CLA	CAA-CBA-CGA-O2A
23	E	318	LMG	O7-C10-C11-C12
20	j	102	LHG	C17-C18-C19-C20
18	a	806	CLA	C16-C17-C18-C20
22	a	849	SQD	O6-C44-C45-C46
26	B	301	A86	C12-C11-C13-O
18	a	813	CLA	CAA-CBA-CGA-O1A
26	D	206	A86	C24-C25-C26-C27
28	E	307	DD6	C11-C10-C9-C8
22	a	849	SQD	O10-C23-C24-C25
18	a	804	CLA	CAD-CBD-CGD-O2D
18	a	808	CLA	CAD-CBD-CGD-O2D
18	b	826	CLA	CAD-CBD-CGD-O2D

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Mol	Chain	Res	Type	Atoms
18	b	837	CLA	CAD-CBD-CGD-O2D
18	b	846	CLA	CAD-CBD-CGD-O2D
18	E	314	CLA	CAD-CBD-CGD-O2D
29	H	312	KC1	CAD-CBD-CGD-O2D
18	b	842	CLA	C15-C16-C17-C18
18	a	807	CLA	CAA-CBA-CGA-O2A
18	a	822	CLA	CAA-CBA-CGA-O2A
18	b	803	CLA	CAA-CBA-CGA-O2A
18	b	806	CLA	CAA-CBA-CGA-O2A
18	D	210	CLA	O1A-CGA-O2A-C1
25	b	834	DGD	O6D-C5D-C6D-O5D
18	l	205	CLA	CAA-CBA-CGA-O2A
18	b	822	CLA	CAA-CBA-CGA-O2A
18	a	805	CLA	O1A-CGA-O2A-C1
18	a	806	CLA	C16-C17-C18-C19
18	a	820	CLA	C12-C13-C15-C16
18	a	805	CLA	C2-C3-C5-C6
18	a	847	CLA	CAA-CBA-CGA-O1A
18	b	844	CLA	CAA-CBA-CGA-O1A
18	b	849	CLA	CAA-CBA-CGA-O1A
18	H	305	CLA	O1D-CGD-O2D-CED

There are no ring outliers.

188 monomers are involved in 611 short contacts:

Mol	Chain	Res	Type	Clashes	Symm-Clashes
18	a	809	CLA	4	0
21	i	103	BCR	5	0
21	f	201	BCR	4	0
26	b	847	A86	4	0
18	b	849	CLA	5	0
18	C	312	CLA	10	0
18	D	217	CLA	3	0
18	b	814	CLA	7	0
18	a	815	CLA	6	0
18	a	816	CLA	1	0
20	b	835	LHG	5	0
18	E	311	CLA	1	0
18	a	825	CLA	5	0
18	a	828	CLA	2	0
21	m	101	BCR	5	0
18	b	817	CLA	6	0

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Mol	Chain	Res	Type	Clashes	Symm-Clashes
18	B	311	CLA	4	0
26	m	102	A86	5	0
18	a	814	CLA	2	0
18	E	310	CLA	1	0
18	a	802	CLA	12	0
23	l	207	LMG	1	0
18	b	846	CLA	2	0
18	H	314	CLA	5	0
18	a	838	CLA	2	0
18	b	830	CLA	1	0
18	H	310	CLA	3	0
21	r	201	BCR	2	0
18	a	844	CLA	5	0
23	E	318	LMG	5	0
20	B	315	LHG	3	0
18	B	308	CLA	1	0
18	a	823	CLA	1	0
18	a	817	CLA	1	0
29	B	313	KC1	5	0
18	a	839	CLA	1	0
18	D	207	CLA	3	0
18	b	818	CLA	3	0
18	B	312	CLA	7	0
18	B	307	CLA	4	0
18	b	811	CLA	1	0
18	l	202	CLA	4	0
18	b	828	CLA	1	0
18	a	808	CLA	5	0
18	b	805	CLA	4	0
21	E	306	BCR	7	0
23	C	319	LMG	1	0
18	H	306	CLA	8	0
18	a	807	CLA	14	0
20	D	201	LHG	6	0
21	a	852	BCR	14	0
18	a	806	CLA	4	0
18	b	838	CLA	4	0
18	a	801	CLA	3	0
18	D	214	CLA	5	0
21	j	105	BCR	11	0
18	B	309	CLA	2	0
18	a	805	CLA	2	0

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Mol	Chain	Res	Type	Clashes	Symm-Clashes
26	H	302	A86	17	0
21	b	832	BCR	3	0
18	C	316	CLA	10	0
18	a	846	CLA	1	0
18	b	825	CLA	3	0
18	b	824	CLA	5	0
21	j	101	BCR	6	0
24	c	101	SF4	2	0
18	b	844	CLA	2	0
18	B	314	CLA	3	0
18	E	315	CLA	3	0
18	D	215	CLA	5	0
18	H	307	CLA	4	0
18	a	830	CLA	3	0
18	b	829	CLA	6	0
21	a	833	BCR	17	0
18	a	841	CLA	7	0
18	b	812	CLA	2	0
26	C	304	A86	1	0
18	C	310	CLA	3	0
18	a	854	CLA	3	0
26	D	203	A86	3	0
18	a	847	CLA	3	0
20	a	831	LHG	2	0
21	b	839	BCR	9	0
18	b	823	CLA	7	0
18	f	202	CLA	1	0
24	c	102	SF4	5	0
18	a	811	CLA	1	0
18	H	309	CLA	16	0
18	a	804	CLA	7	0
18	b	806	CLA	3	0
18	b	821	CLA	3	0
21	E	305	BCR	2	0
18	a	820	CLA	2	0
24	b	802	SF4	2	0
18	a	848	CLA	5	0
18	a	837	CLA	5	0
26	B	304	A86	2	0
18	E	309	CLA	5	0
21	f	205	BCR	2	0
28	D	205	DD6	3	0

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Mol	Chain	Res	Type	Clashes	Symm-Clashes
19	a	829	PQN	1	0
23	D	202	LMG	1	0
18	H	313	CLA	6	0
18	b	841	CLA	3	0
18	a	819	CLA	1	0
28	H	303	DD6	3	0
29	H	312	KC1	1	0
21	l	201	BCR	3	0
18	C	317	CLA	8	0
18	b	827	CLA	2	0
18	D	211	CLA	2	0
18	b	843	CLA	5	0
18	b	804	CLA	1	0
20	j	102	LHG	5	0
26	C	305	A86	4	0
18	H	311	CLA	6	0
18	l	203	CLA	6	0
18	b	813	CLA	18	0
18	C	307	CLA	6	0
18	b	837	CLA	2	0
19	b	831	PQN	4	0
26	r	203	A86	1	0
29	C	308	KC1	4	0
18	b	820	CLA	5	0
18	B	306	CLA	13	0
18	H	305	CLA	3	0
18	b	803	CLA	4	0
18	b	840	CLA	3	0
18	b	848	CLA	3	0
28	C	303	DD6	1	0
18	b	850	CLA	6	0
21	b	836	BCR	4	0
18	b	808	CLA	2	0
18	D	208	CLA	9	0
18	a	810	CLA	3	0
18	H	308	CLA	4	0
21	b	833	BCR	3	0
18	b	809	CLA	2	0
18	f	204	CLA	1	0
18	E	316	CLA	6	0
18	j	104	CLA	1	0
18	a	818	CLA	2	0

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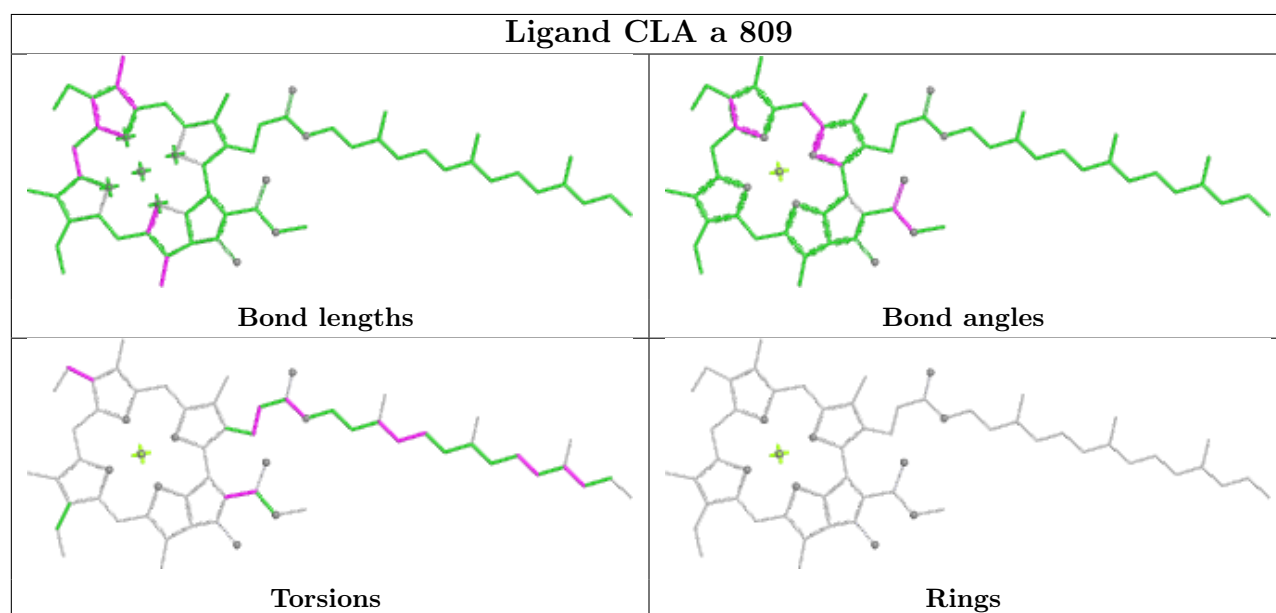
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18	a	840	CLA	2	0
18	D	209	CLA	3	0
20	E	301	LHG	3	0
18	D	212	CLA	1	0
18	C	315	CLA	2	0
21	a	834	BCR	4	0
18	b	842	CLA	6	0
18	C	314	CLA	3	0
22	a	849	SQD	7	0
18	b	819	CLA	4	0
18	i	101	CLA	1	0
18	b	822	CLA	3	0
18	f	203	CLA	1	0
18	a	842	CLA	5	0
21	a	850	BCR	4	0
28	E	307	DD6	1	0
18	a	836	CLA	3	0
18	b	810	CLA	2	0
18	a	821	CLA	1	0
21	l	204	BCR	8	0
18	C	318	CLA	11	0
20	a	832	LHG	3	0
21	i	102	BCR	9	0
26	B	302	A86	3	0
18	b	845	CLA	7	0
18	a	827	CLA	3	0
18	D	210	CLA	1	0
20	a	835	LHG	1	0
23	a	851	LMG	2	0
18	b	807	CLA	3	0
18	a	813	CLA	8	0
18	a	826	CLA	1	0
18	C	309	CLA	6	0
18	C	311	CLA	6	0
18	D	216	CLA	4	0
18	E	313	CLA	2	0
18	b	801	CLA	10	0
18	r	202	CLA	2	0
18	a	855	CLA	5	0
18	E	312	CLA	7	0
21	a	845	BCR	4	0
18	D	213	CLA	2	0

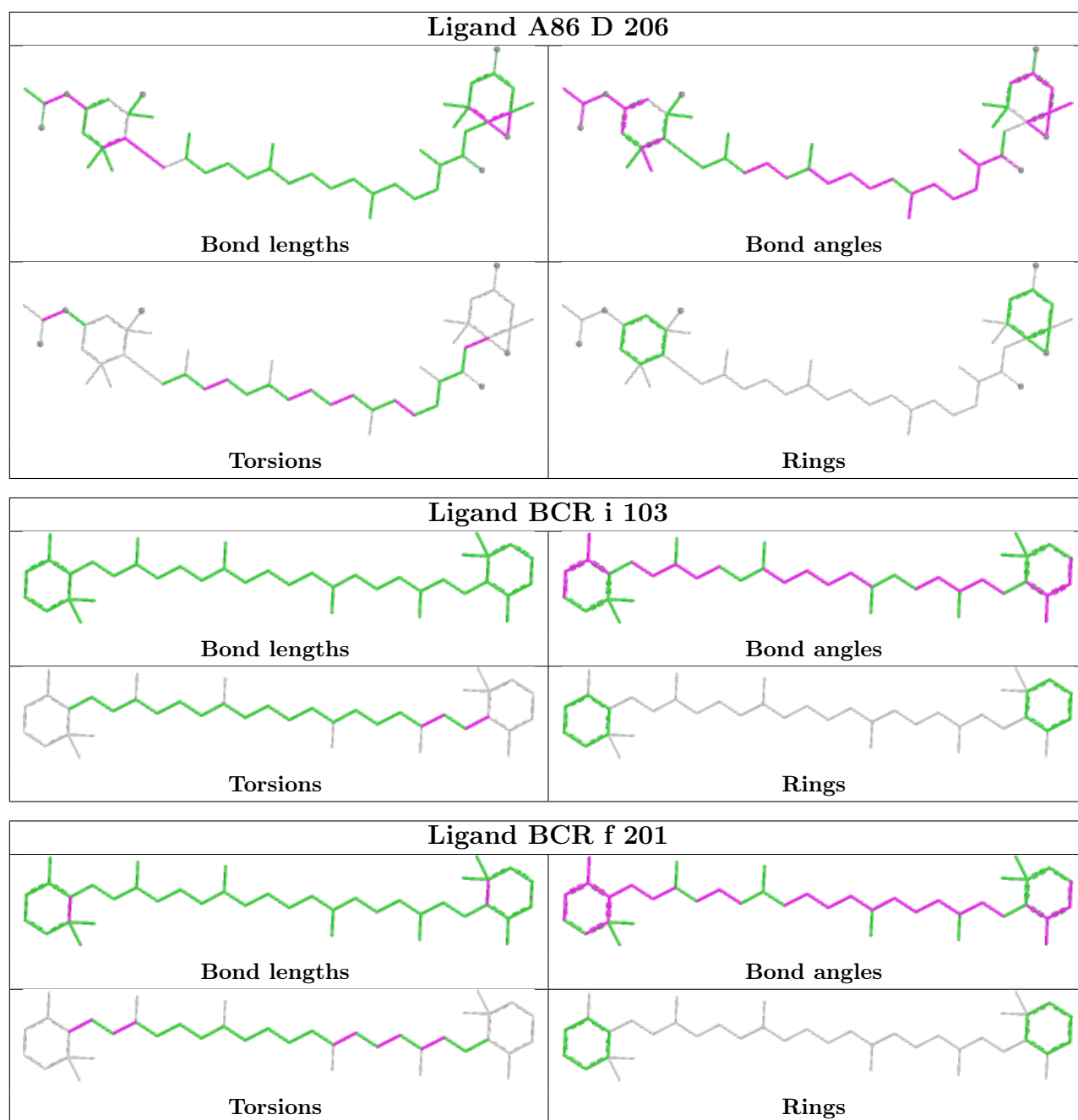
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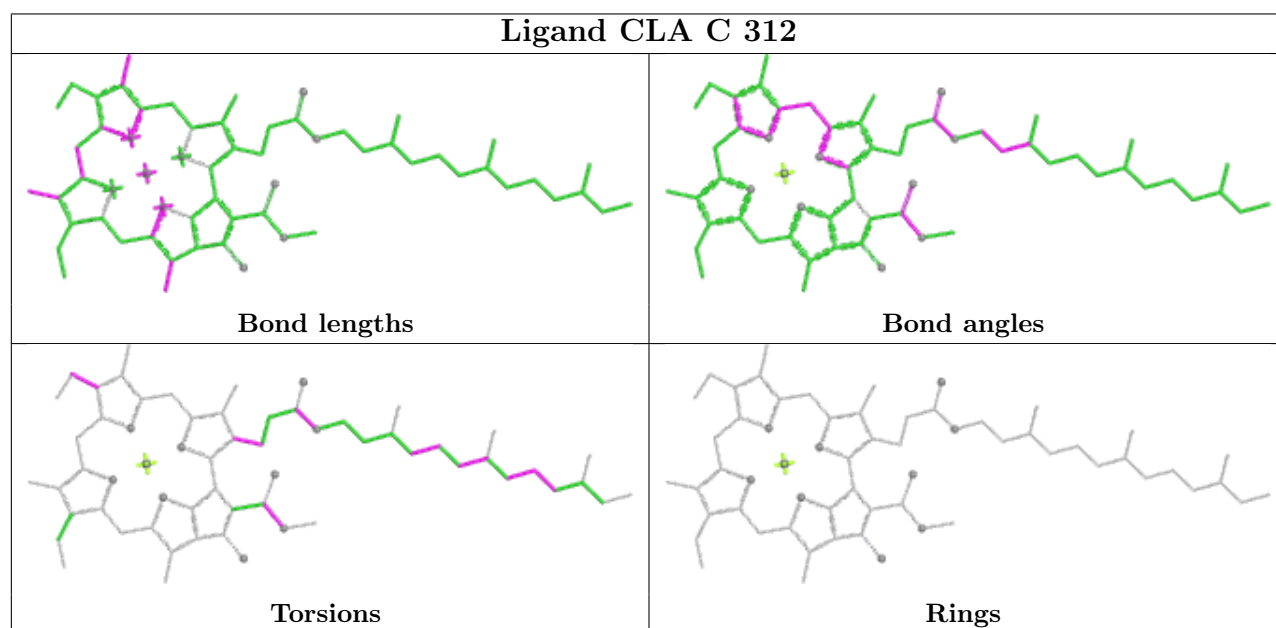
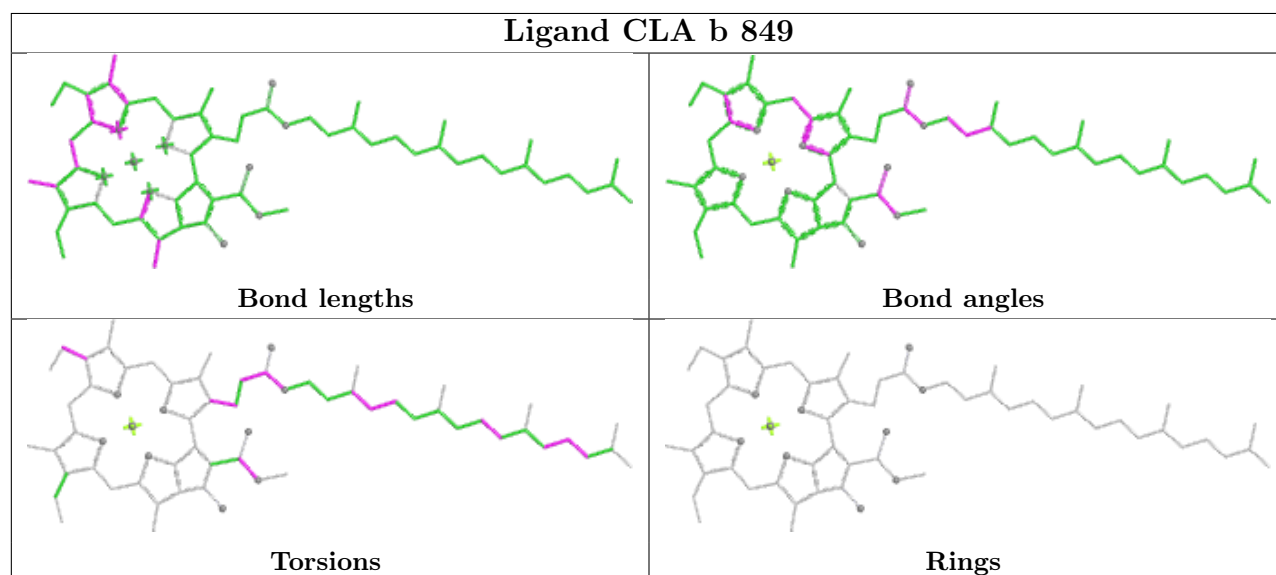
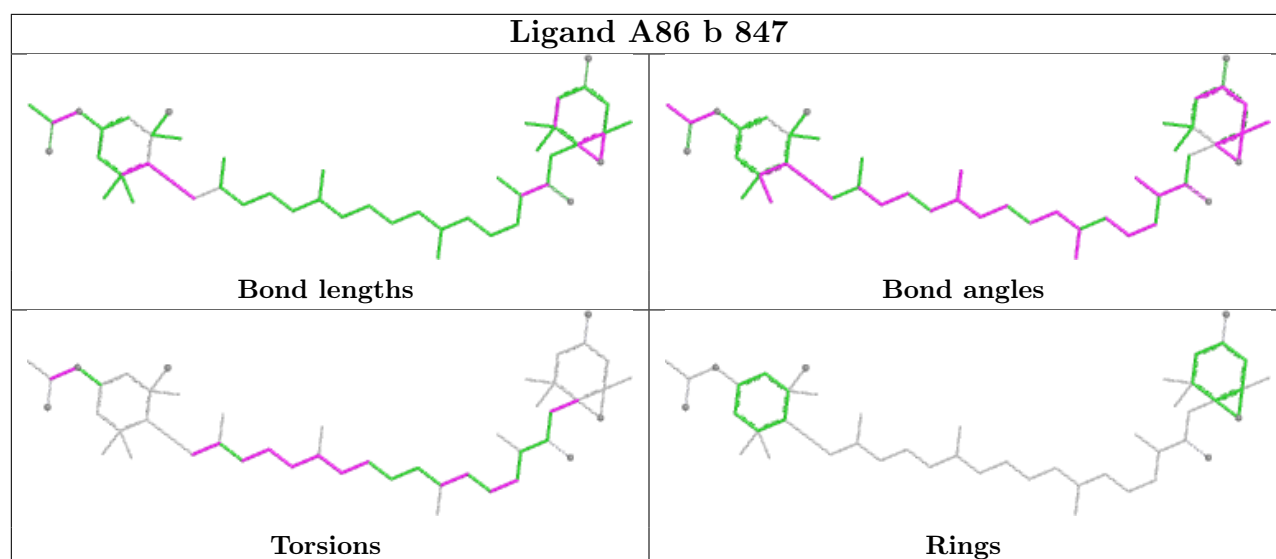
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Mol	Chain	Res	Type	Clashes	Symm-Clashes
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28	E	304	DD6	1	0
18	a	803	CLA	12	0
26	B	301	A86	2	0

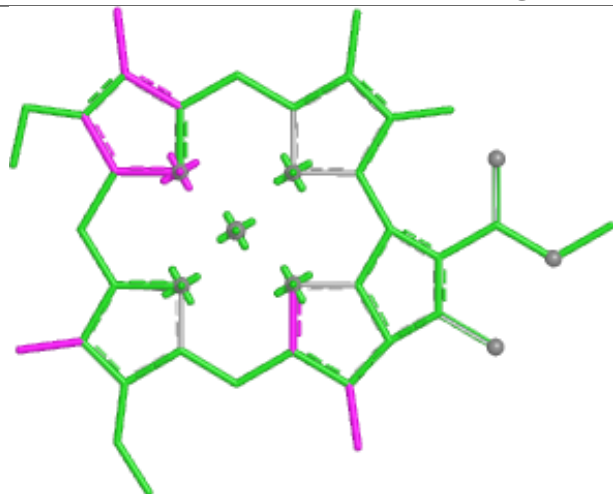
The following is a two-dimensional graphical depiction of Mogul quality analysis of bond lengths, bond angles, torsion angles, and ring geometry for all instances of the Ligand of Interest. In addition, ligands with molecular weight > 250 and outliers as shown on the validation Tables will also be included. For torsion angles, if less than 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.



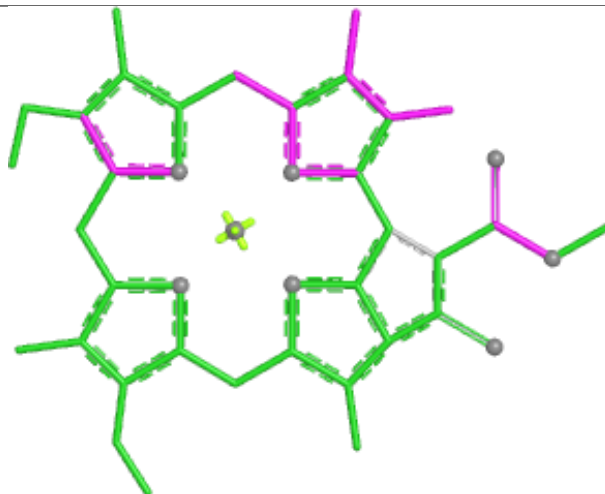




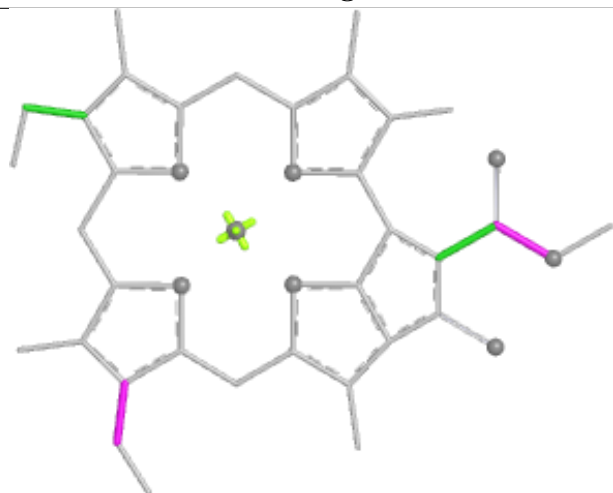
Ligand CLA D 217



Bond lengths



Bond angles

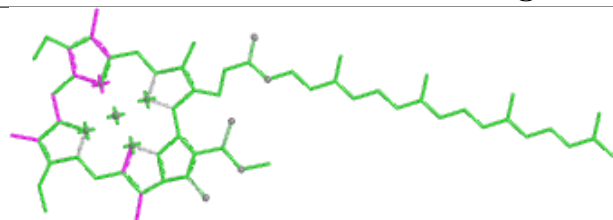


Torsions

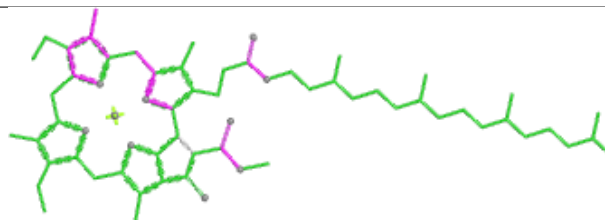


Rings

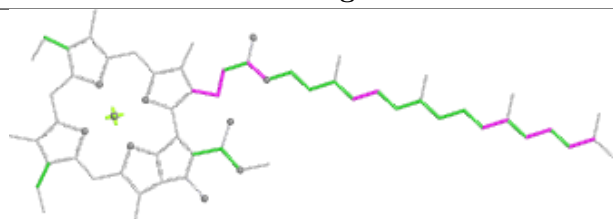
Ligand CLA b 814



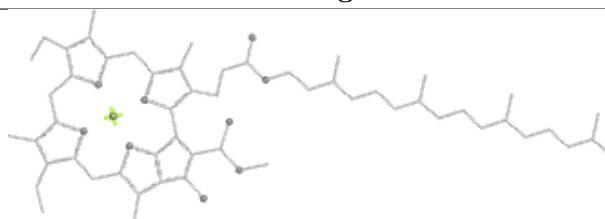
Bond lengths



Bond angles

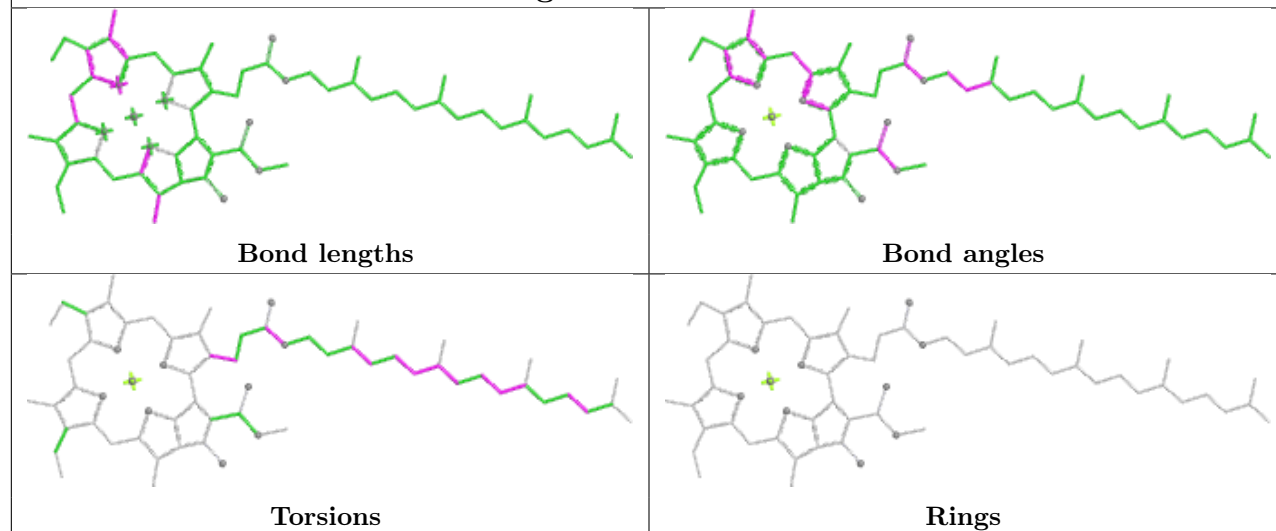


Torsions

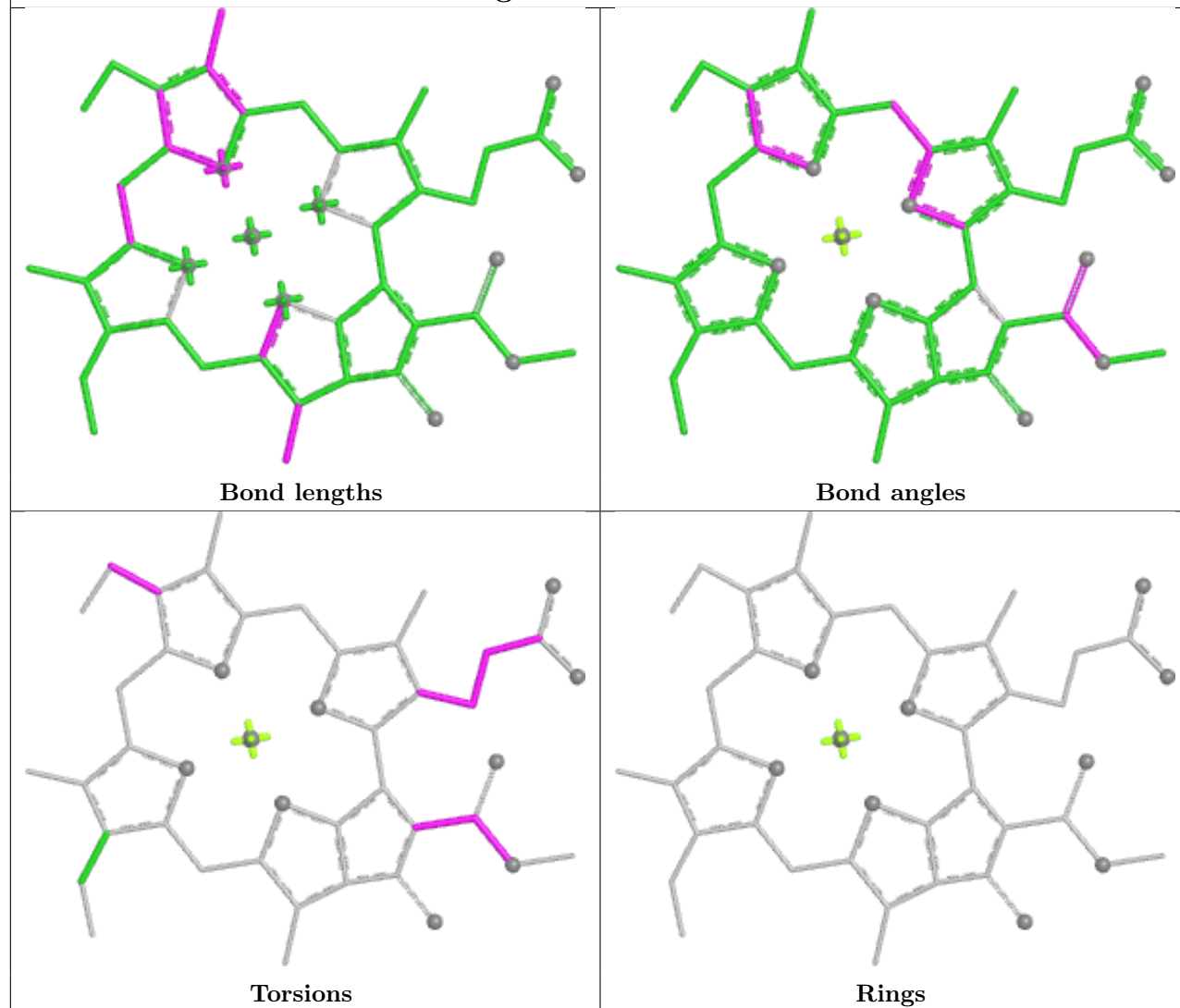


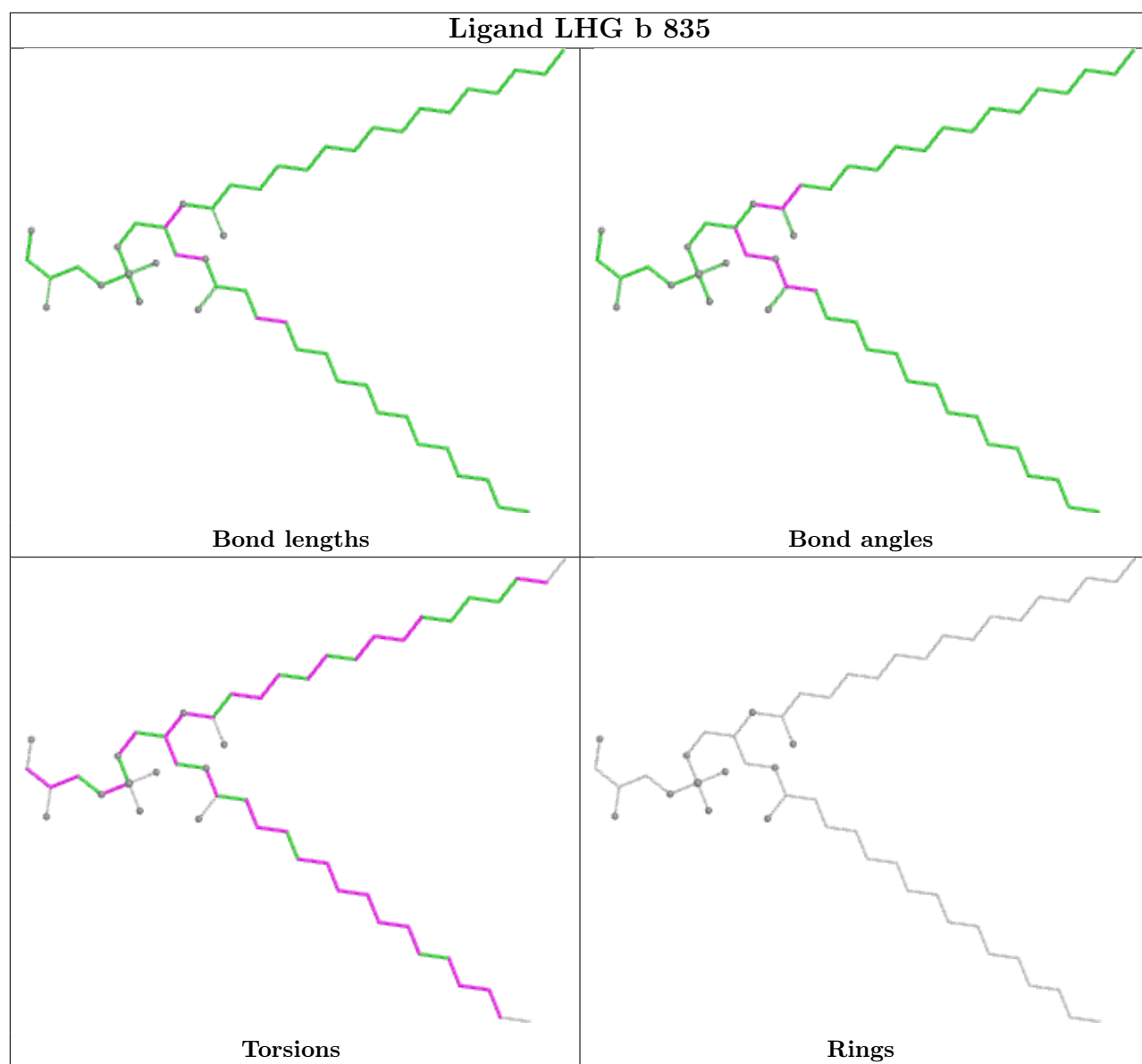
Rings

Ligand CLA a 815

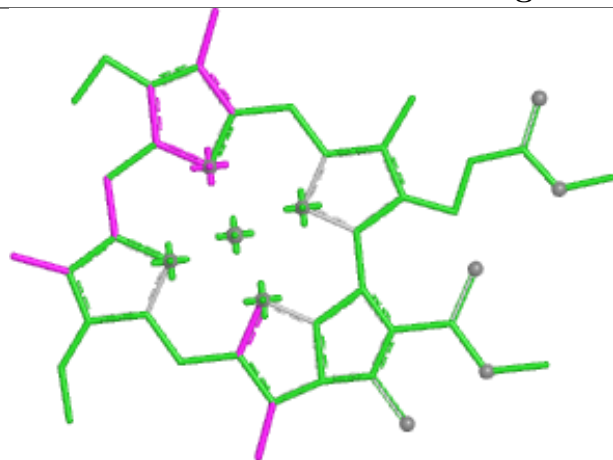


Ligand CLA a 816

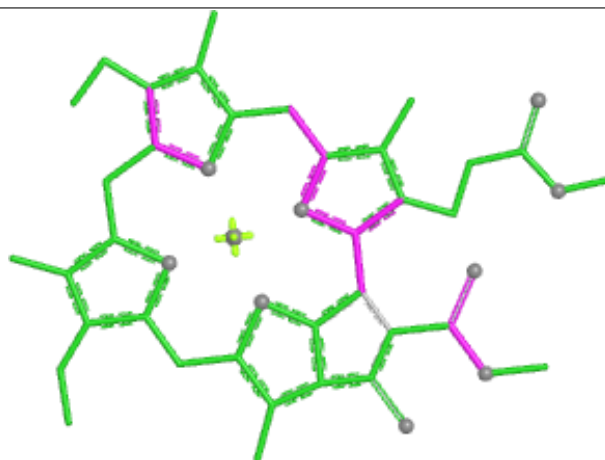




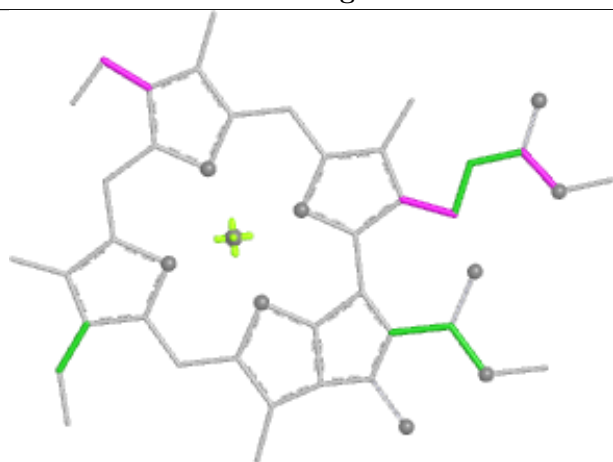
Ligand CLA E 311



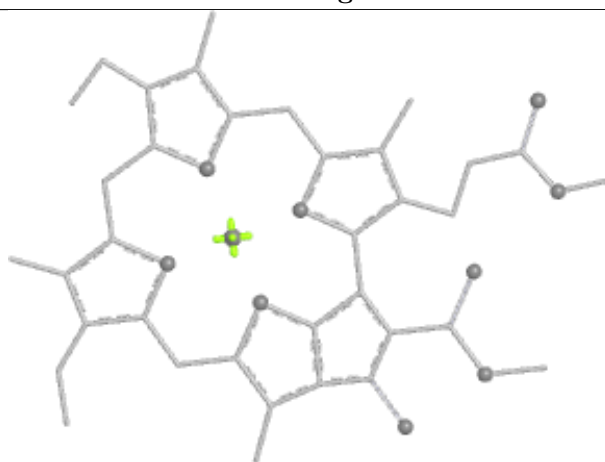
Bond lengths



Bond angles

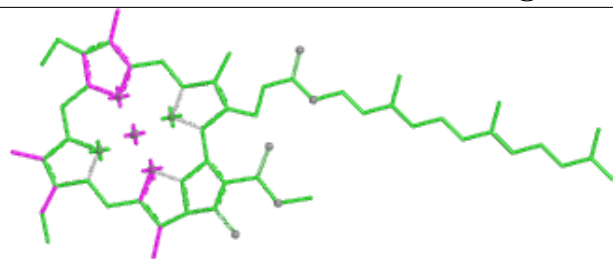


Torsions

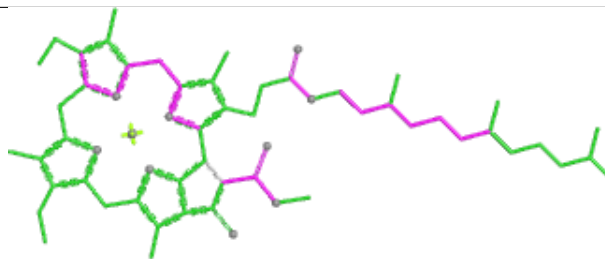


Rings

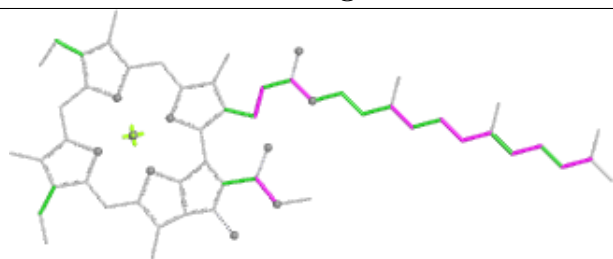
Ligand CLA a 825



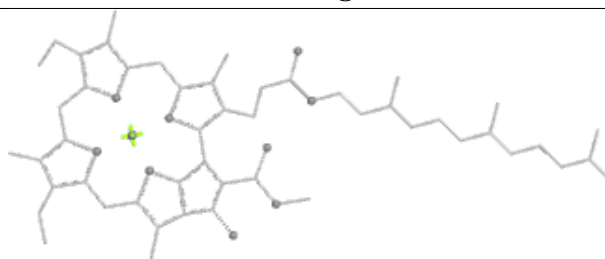
Bond lengths



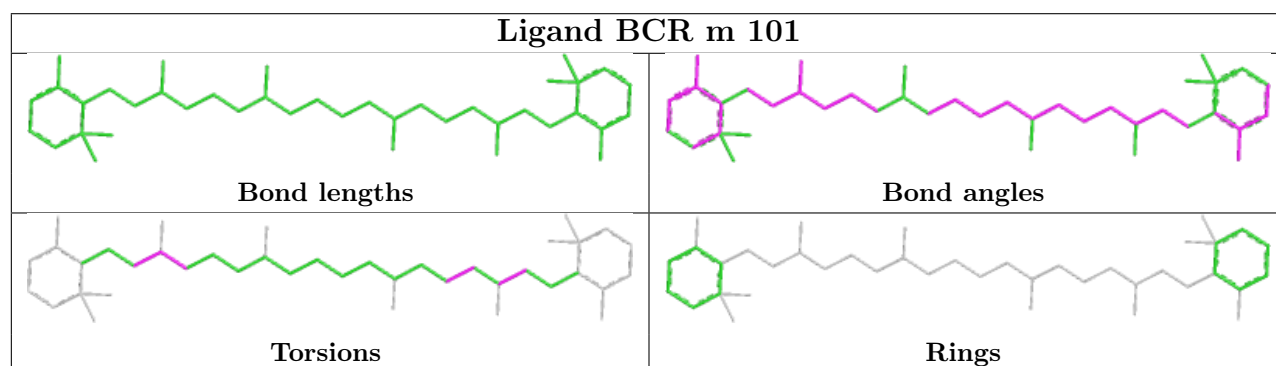
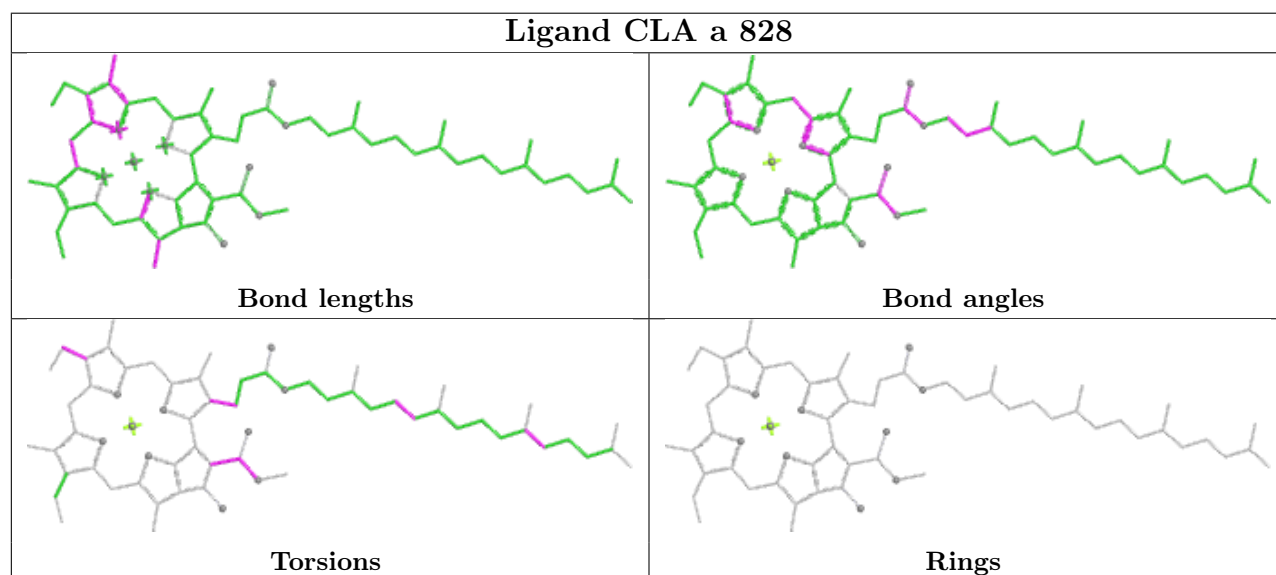
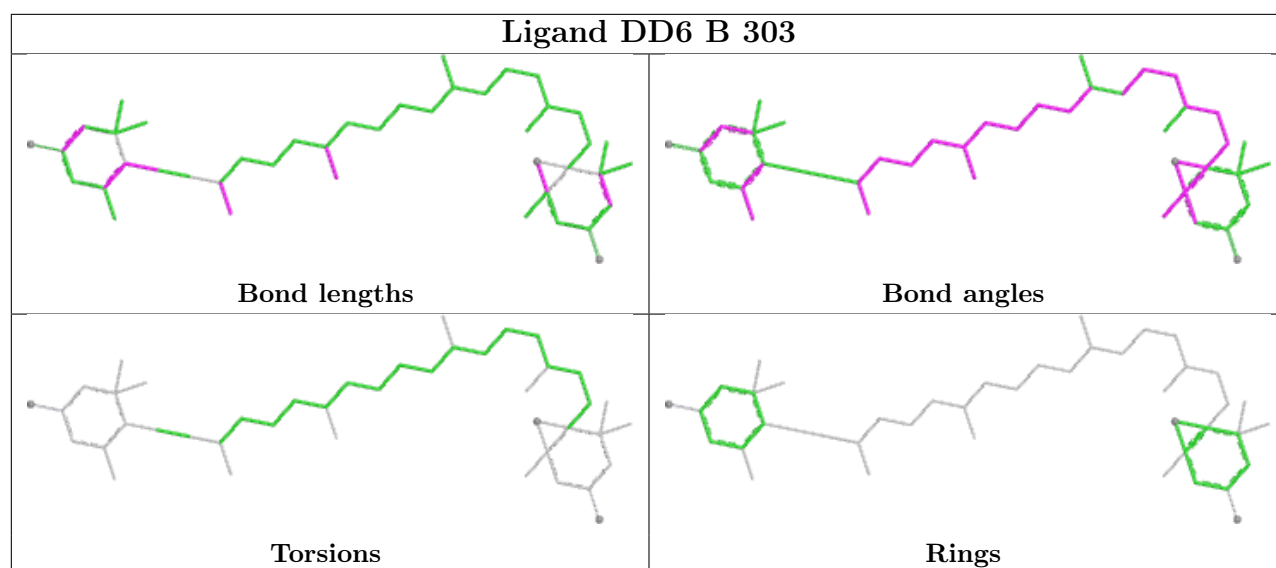
Bond angles



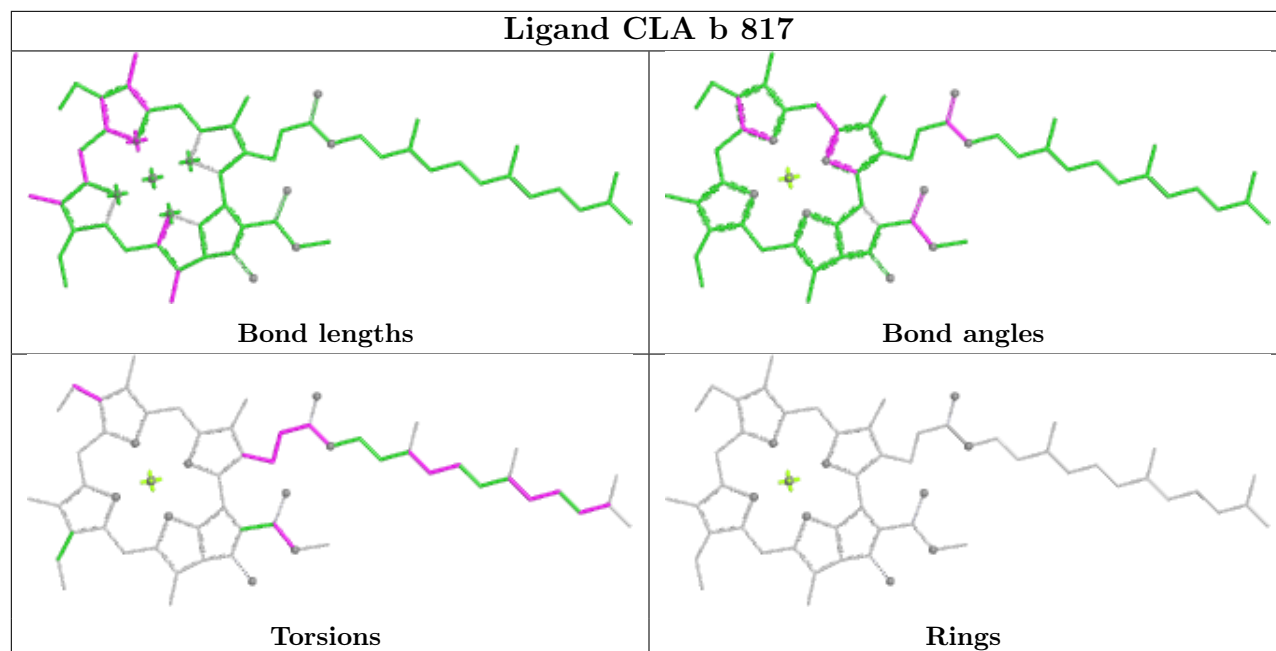
Torsions



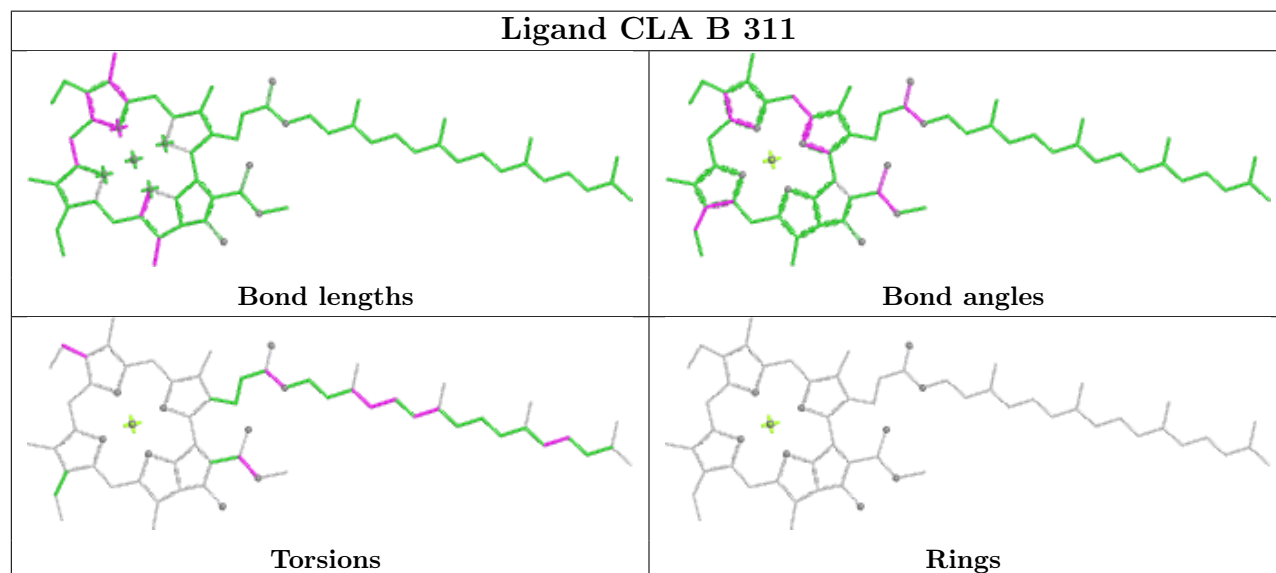
Rings



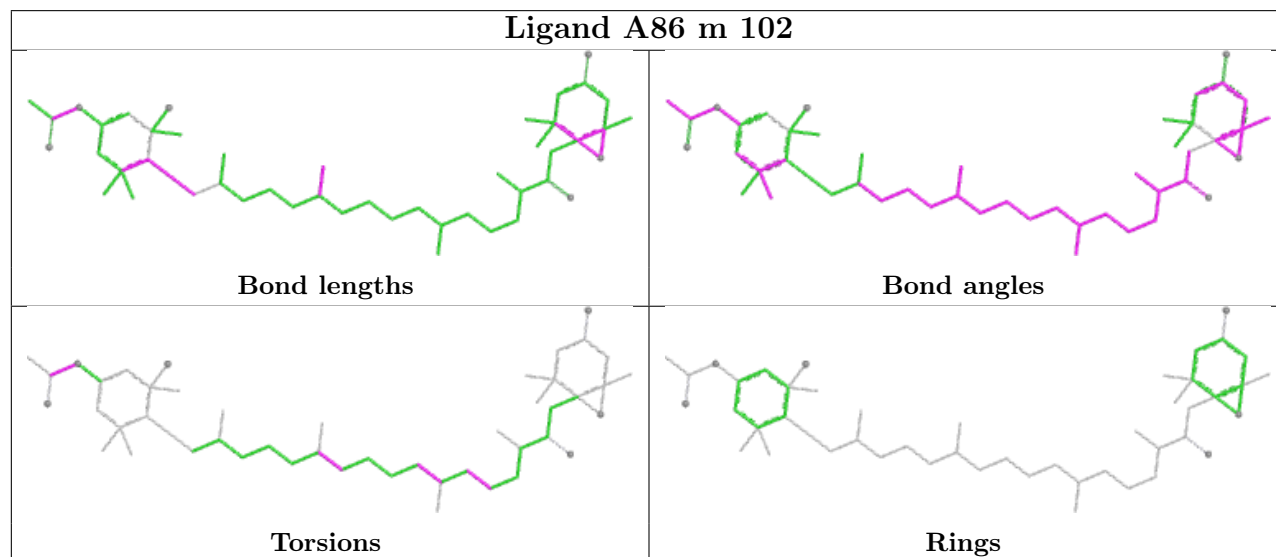
Ligand CLA b 817



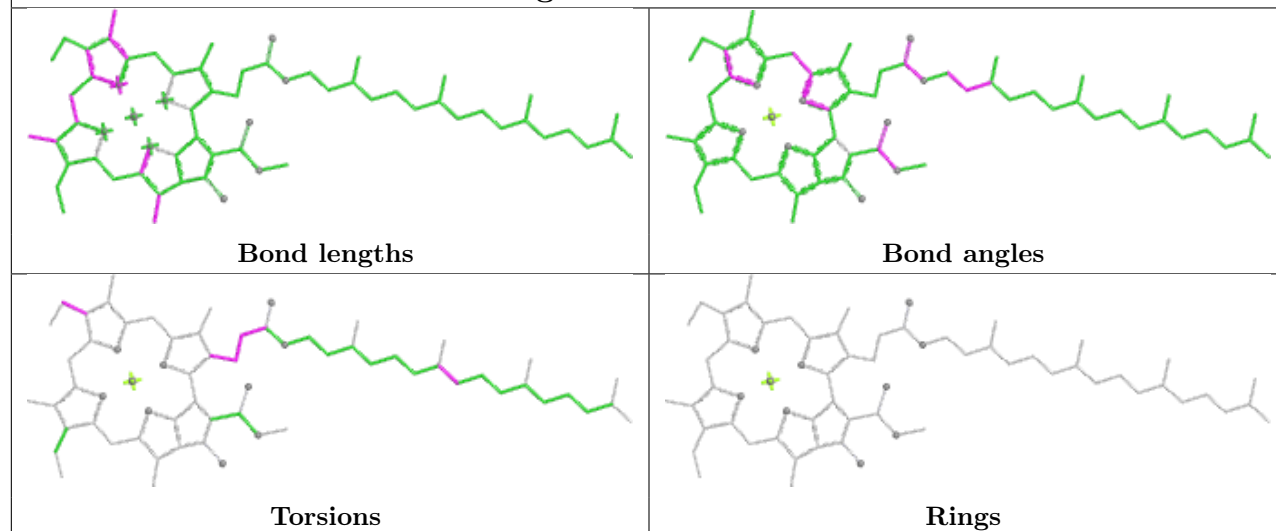
Ligand CLA B 311



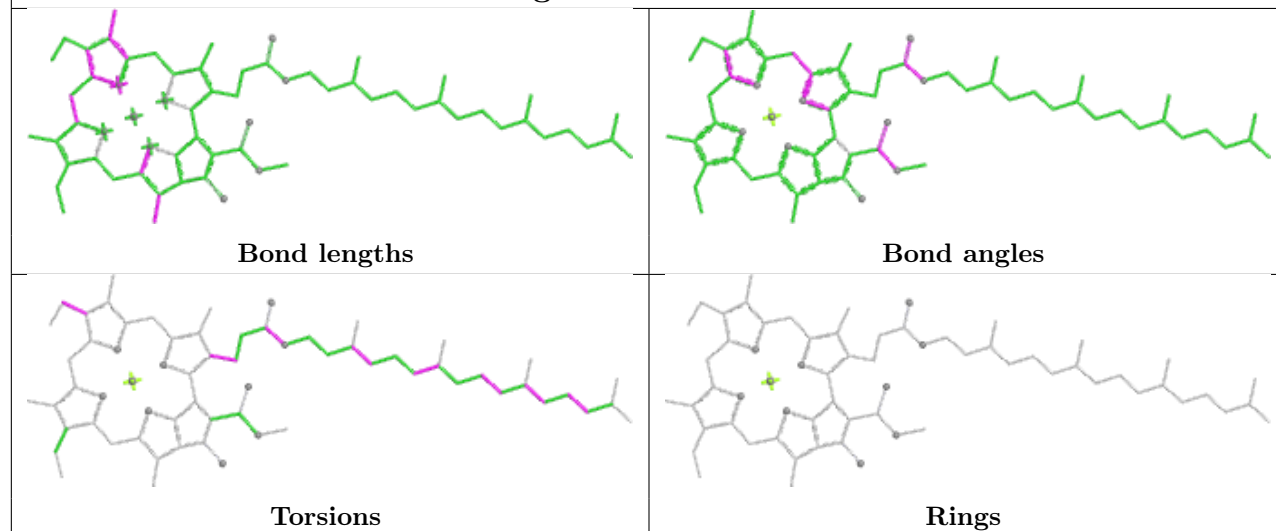
Ligand A86 m 102



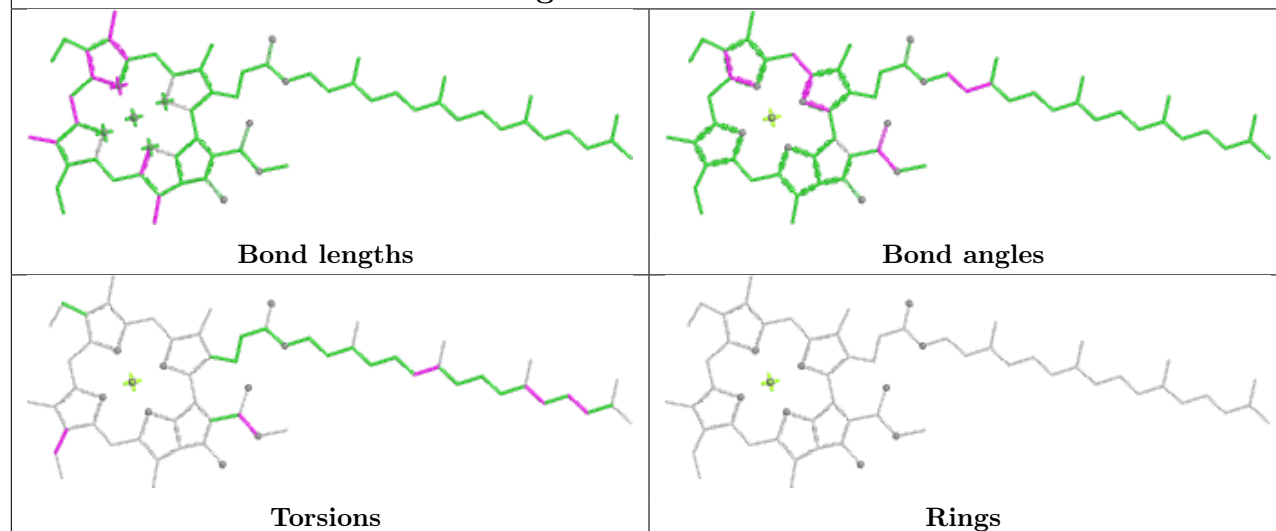
Ligand CLA a 814

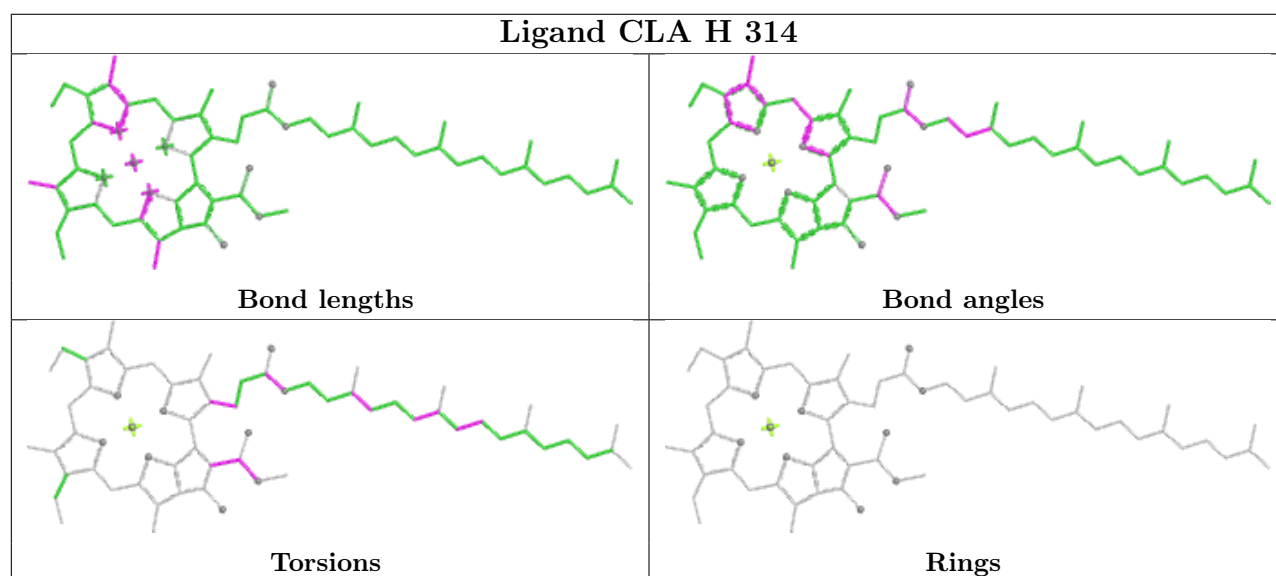
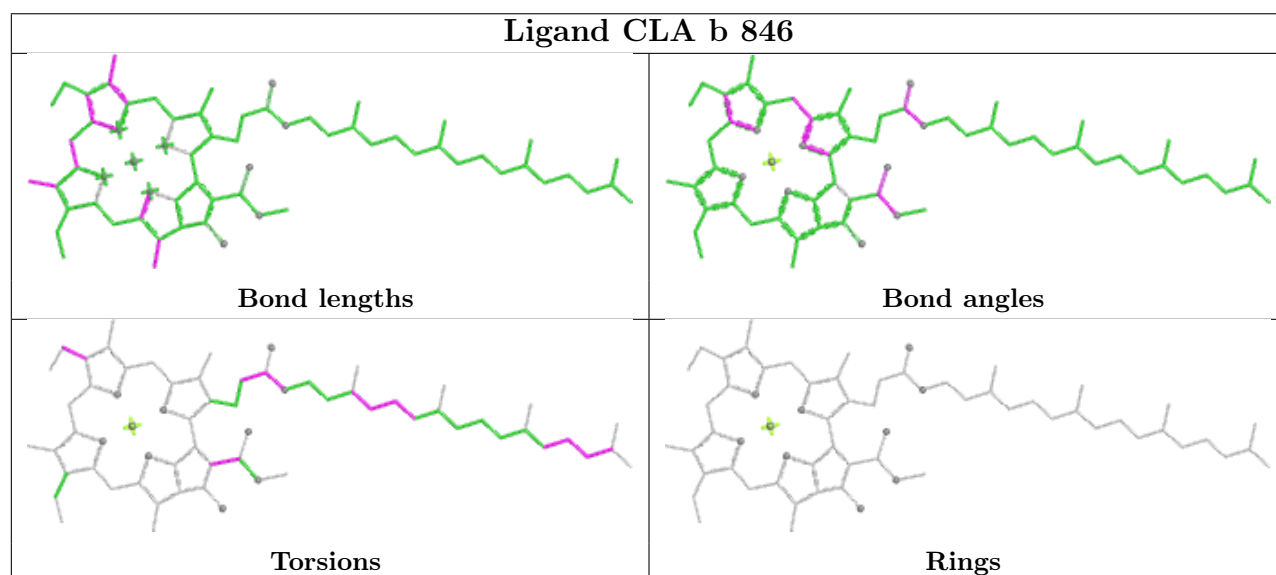
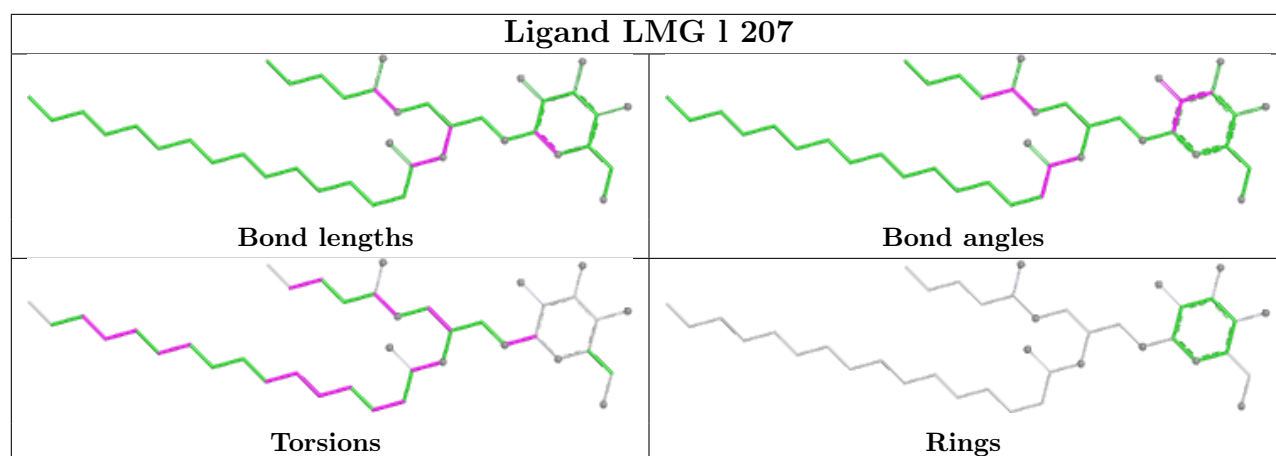


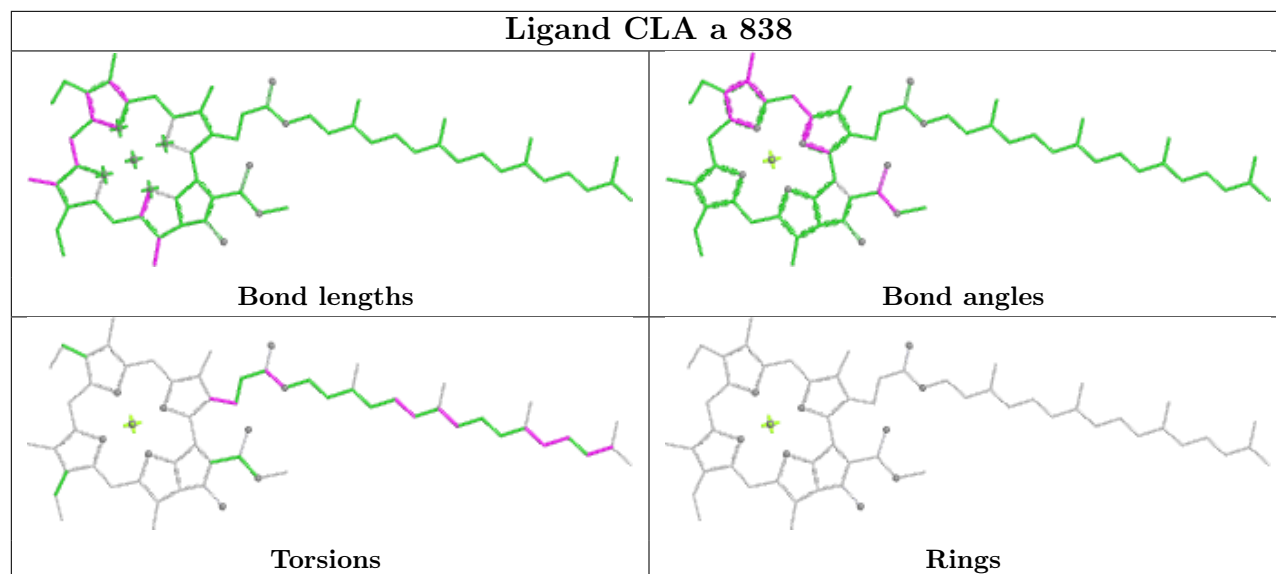
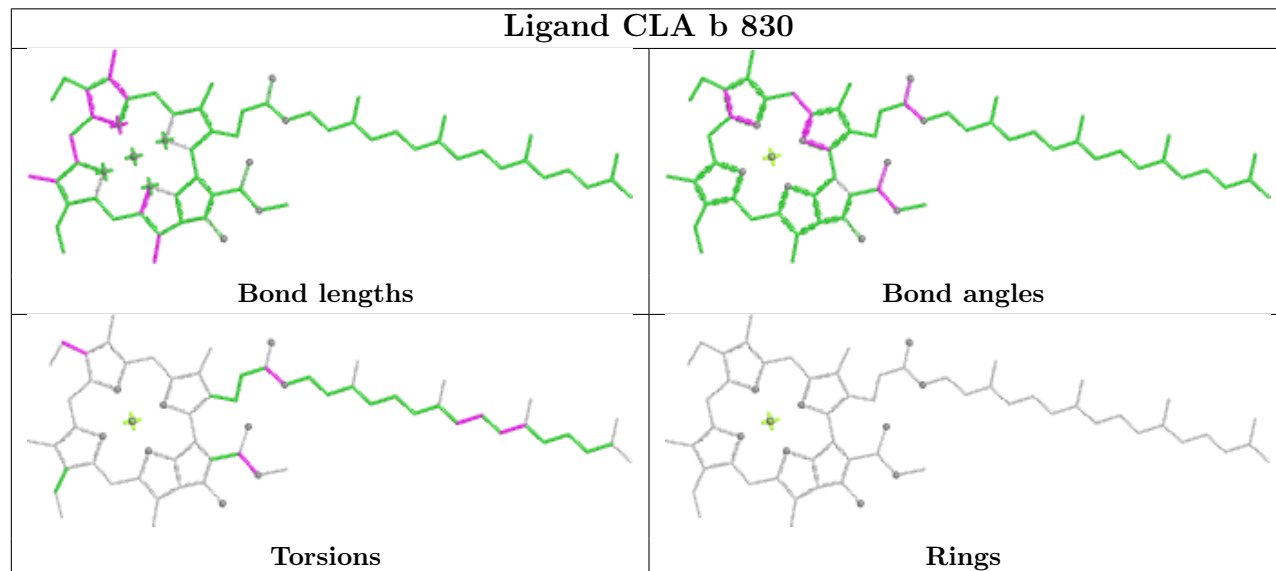
Ligand CLA E 310



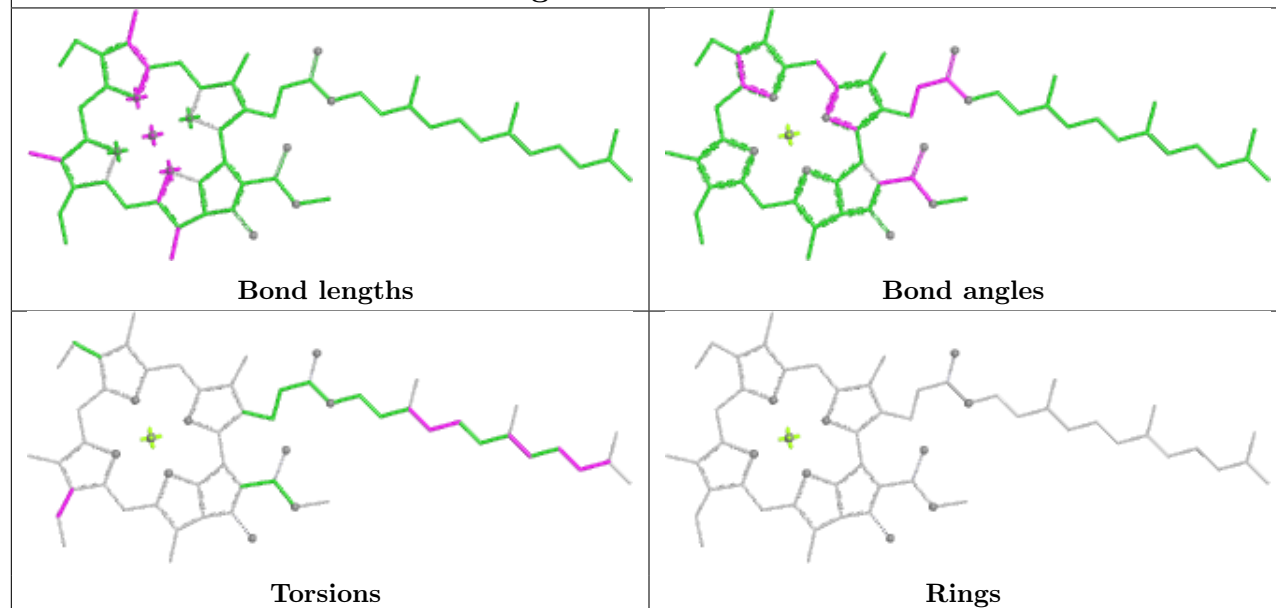
Ligand CLA a 802



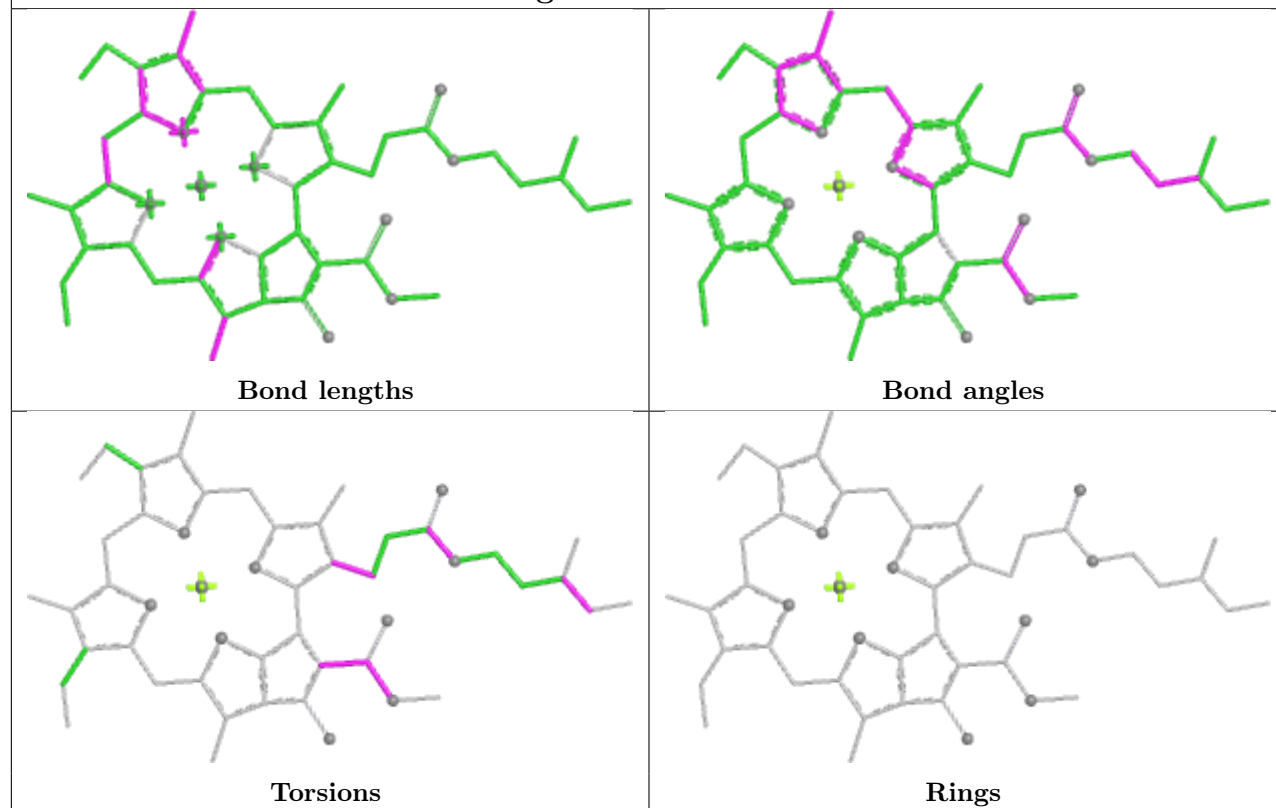


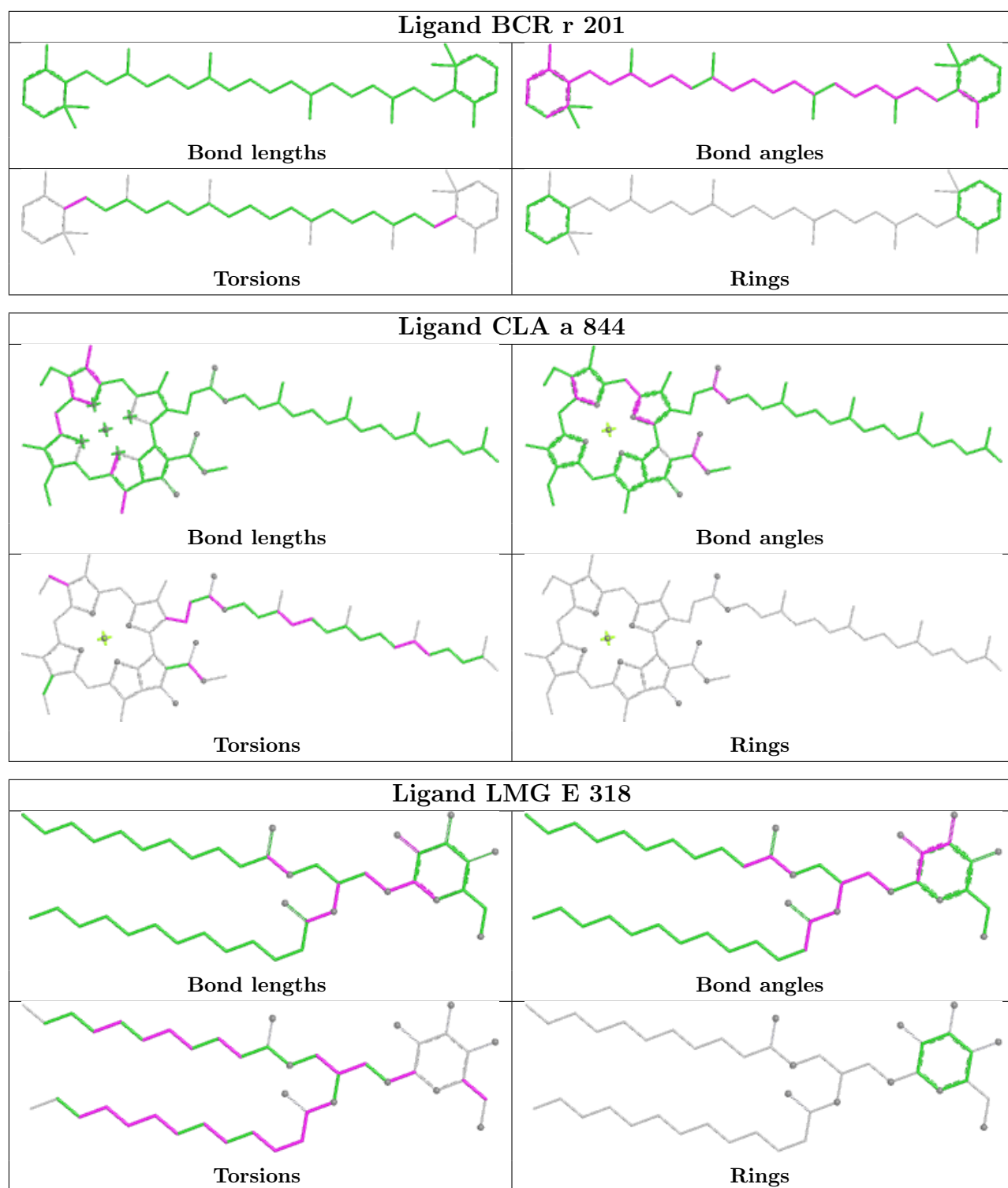
Ligand CLA a 838**Ligand CLA b 830**

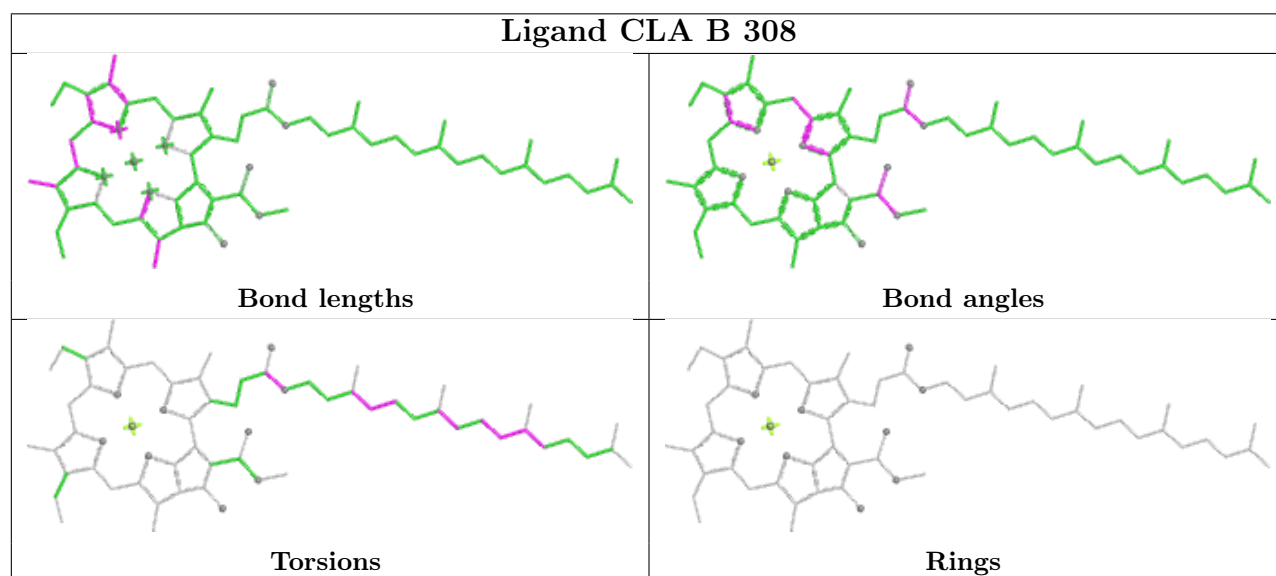
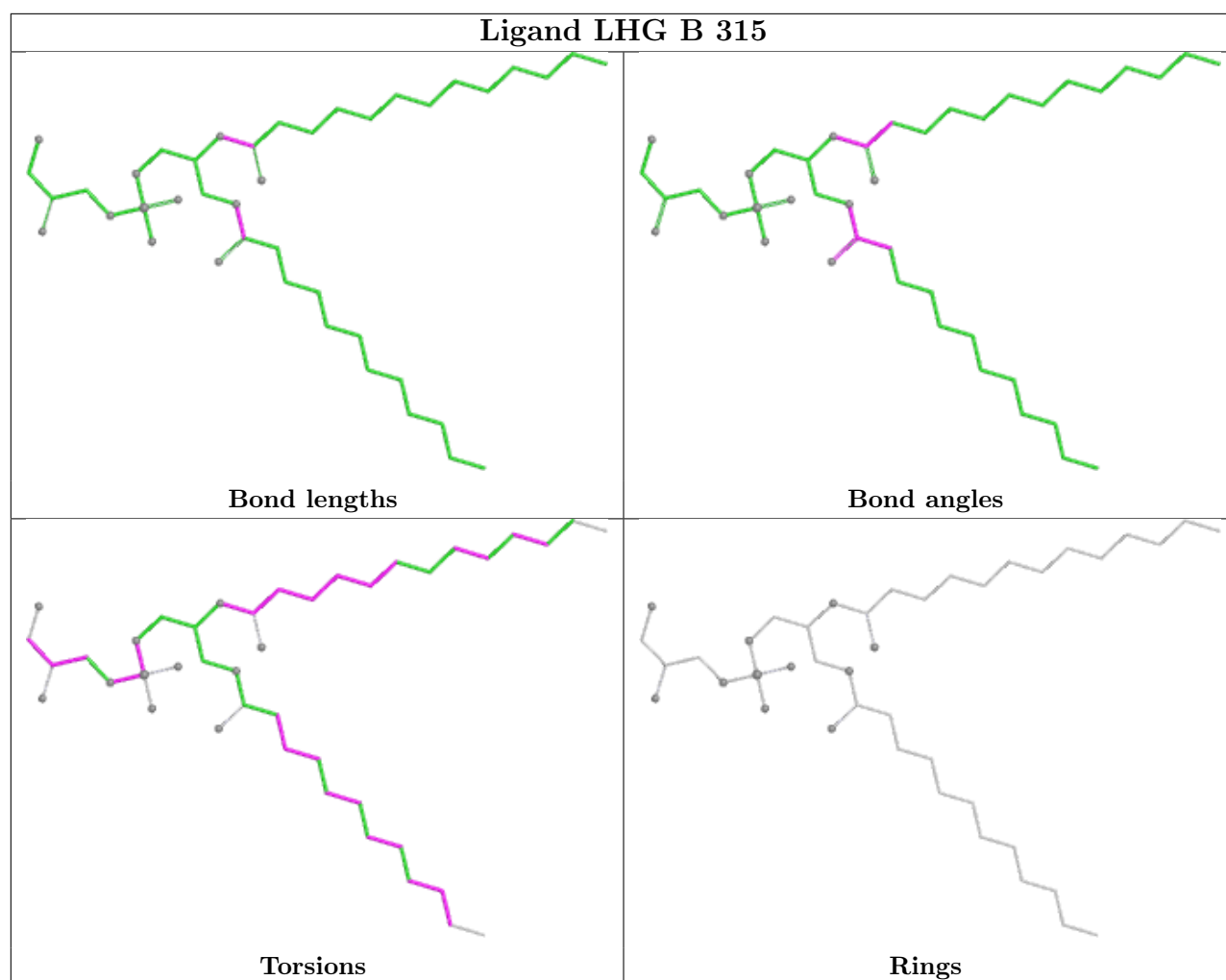
Ligand CLA H 310



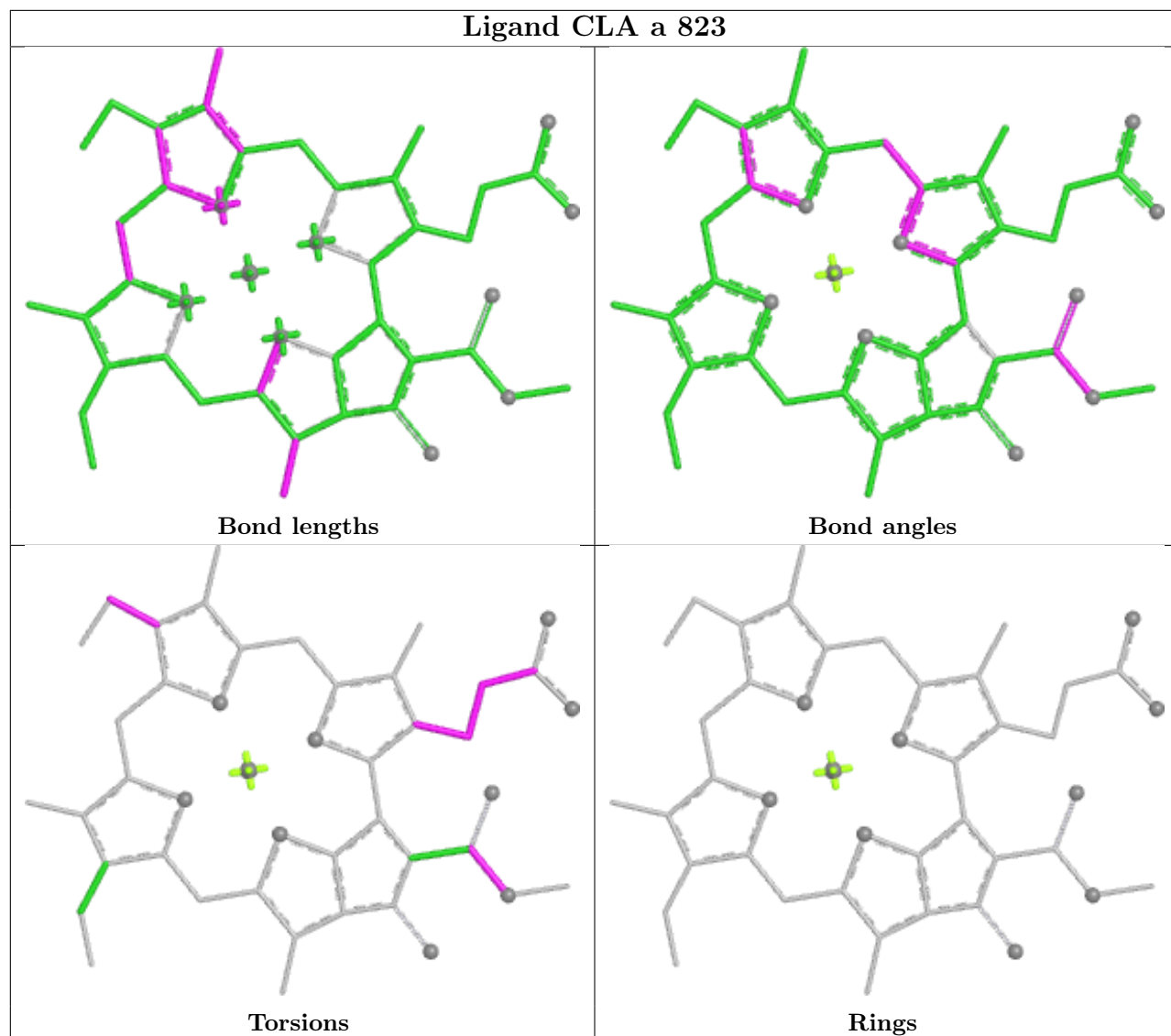
Ligand CLA a 824

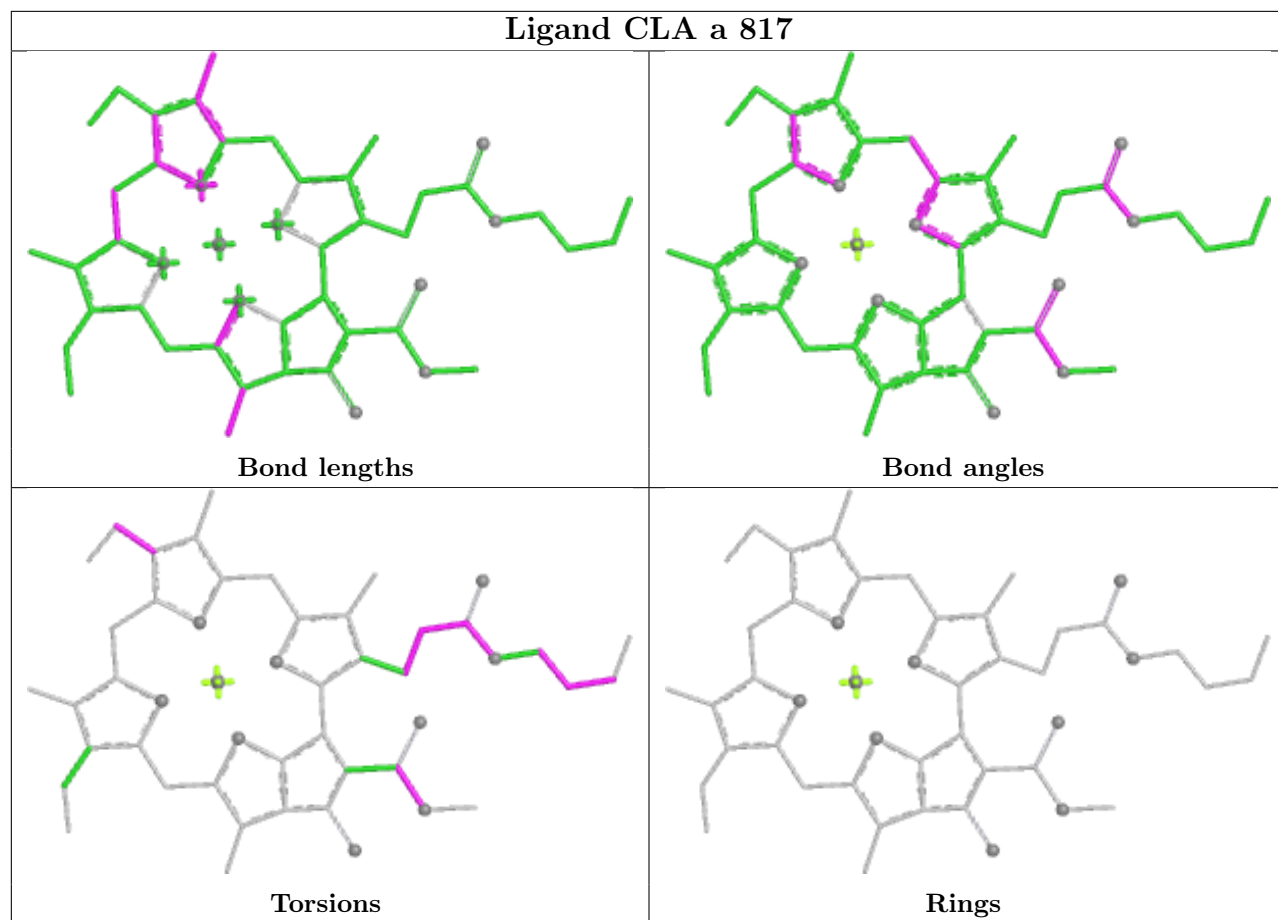




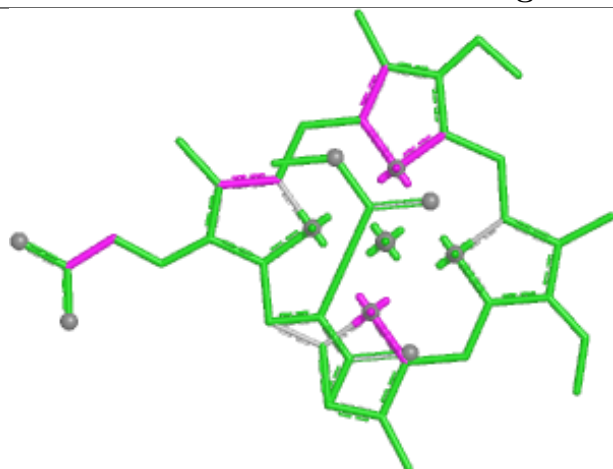


Ligand CLA a 823

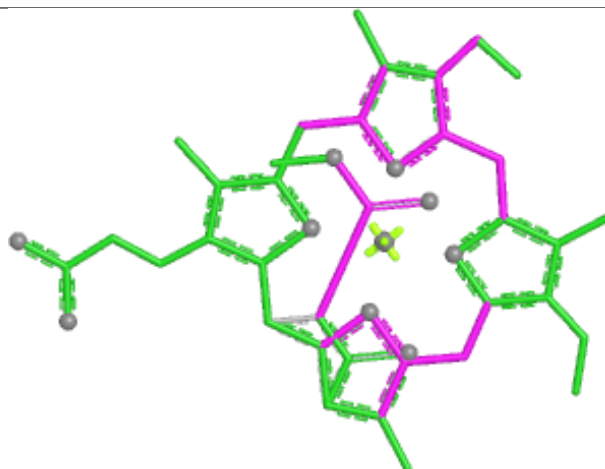




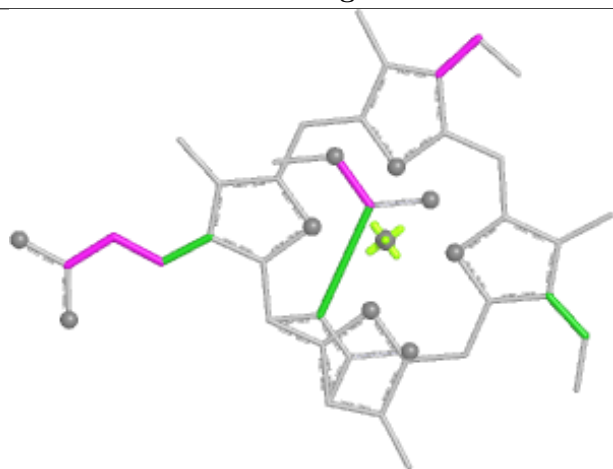
Ligand KC1 B 313



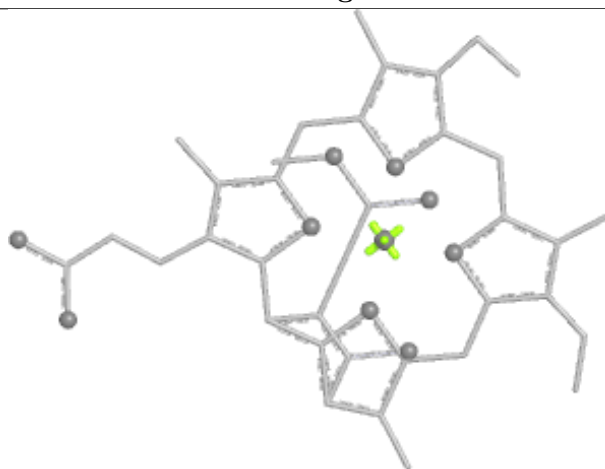
Bond lengths



Bond angles

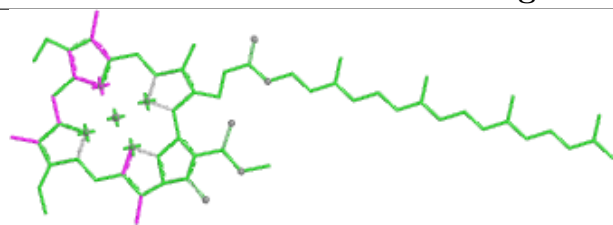


Torsions

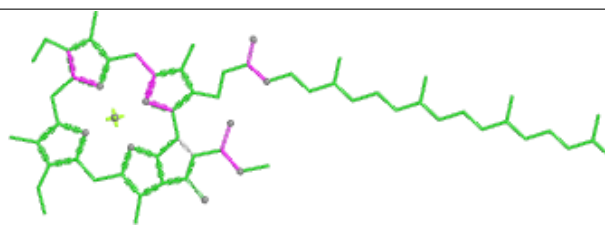


Rings

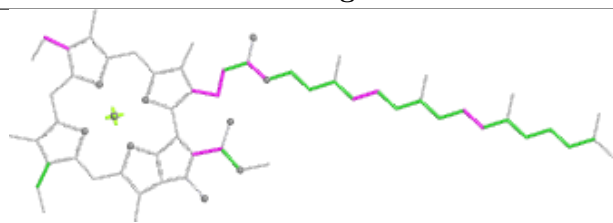
Ligand CLA a 839



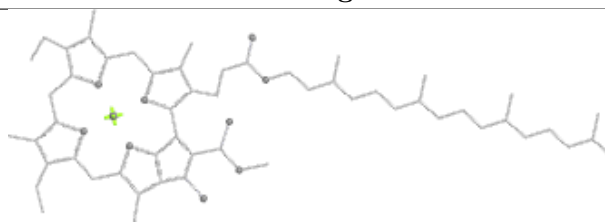
Bond lengths



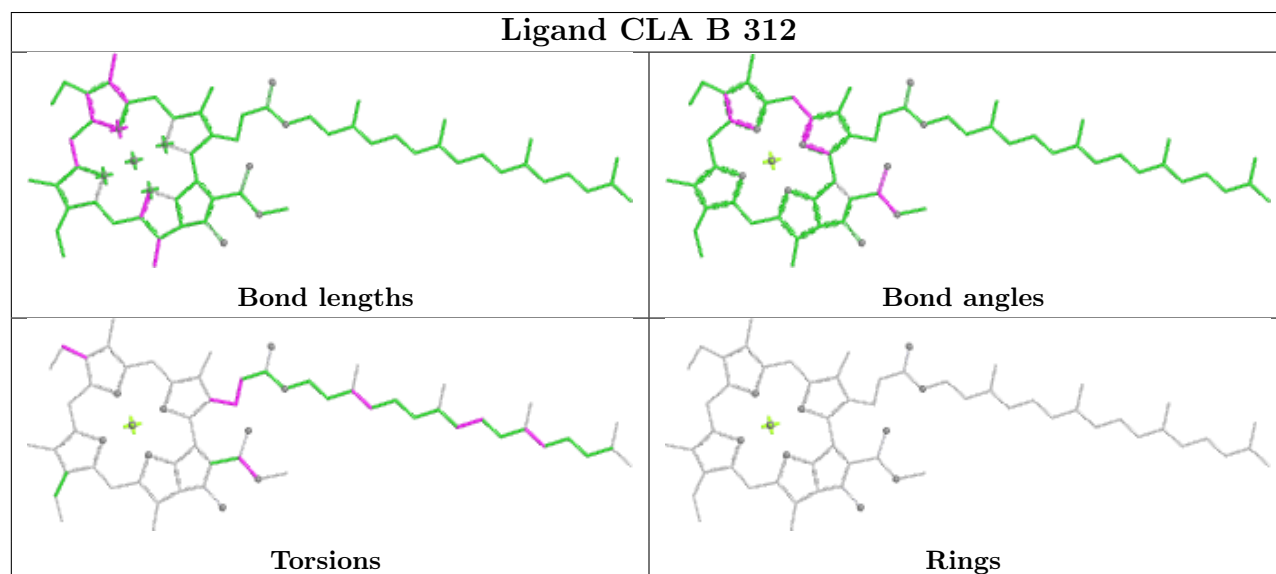
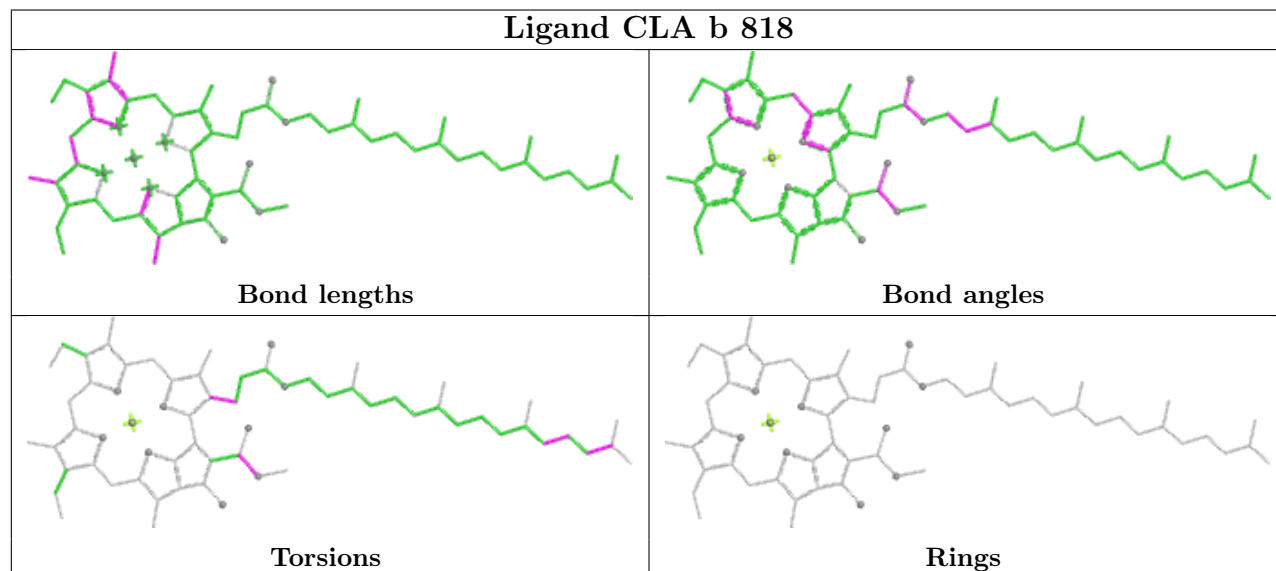
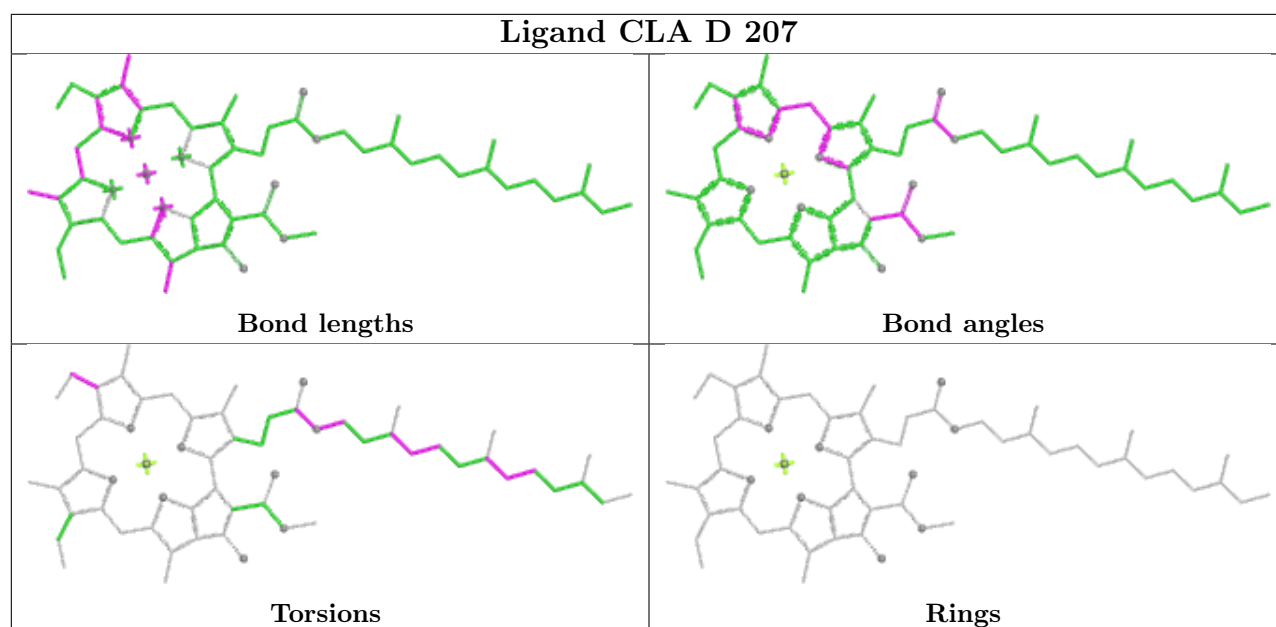
Bond angles

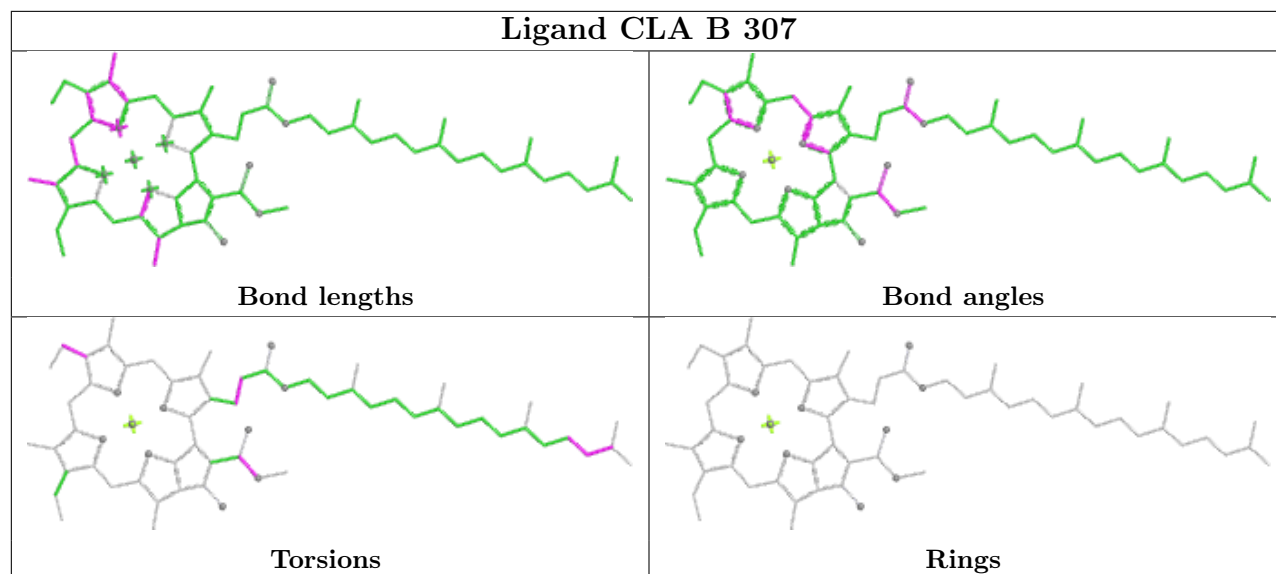
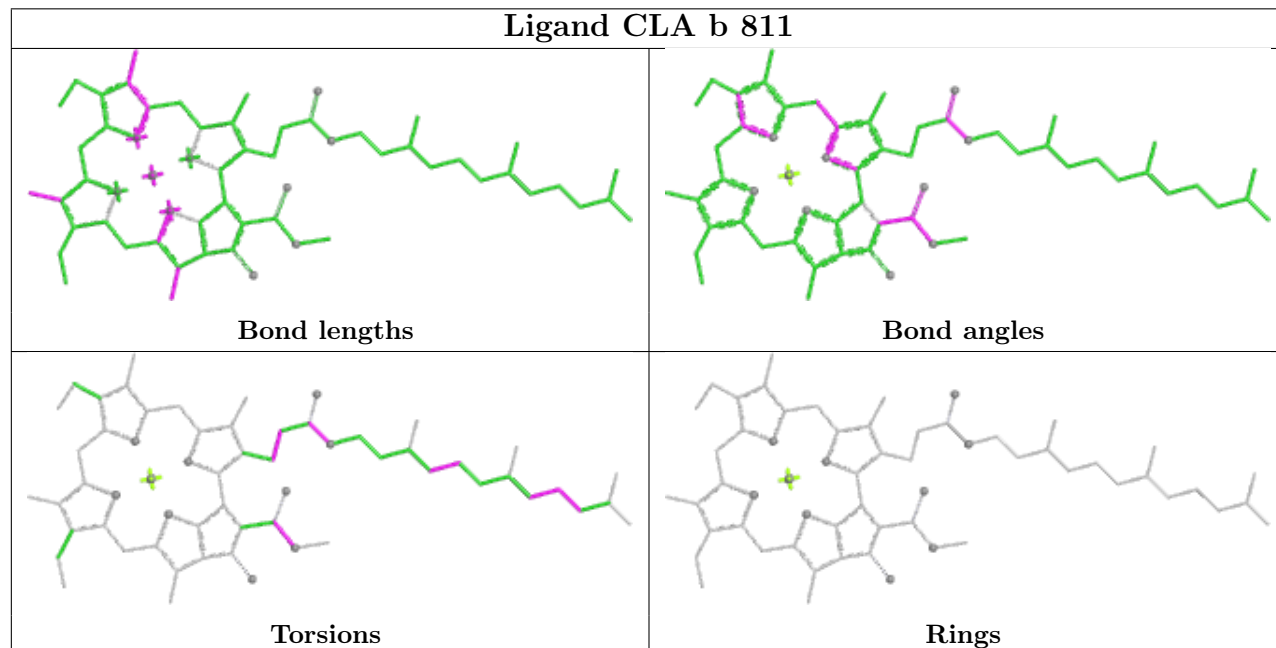


Torsions

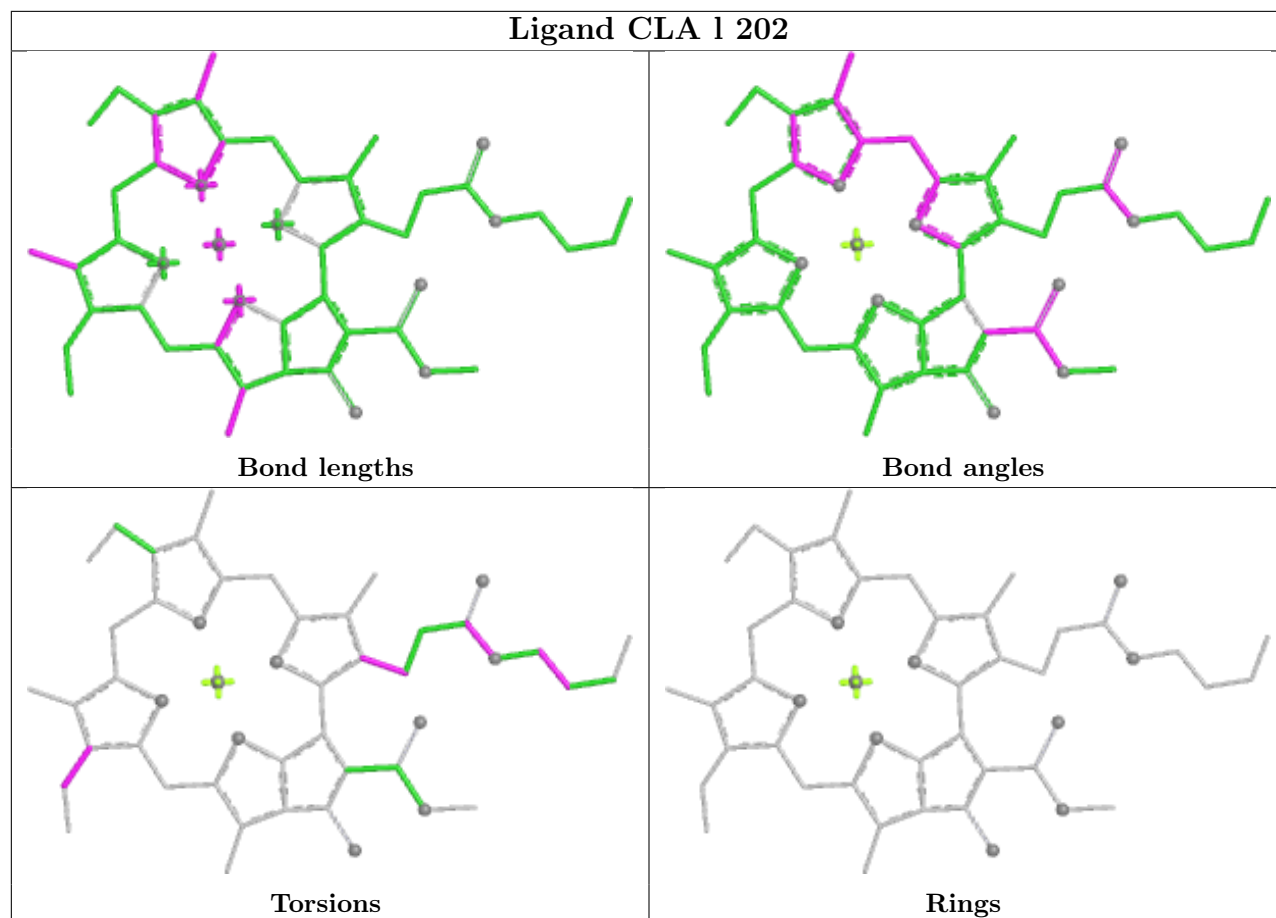


Rings

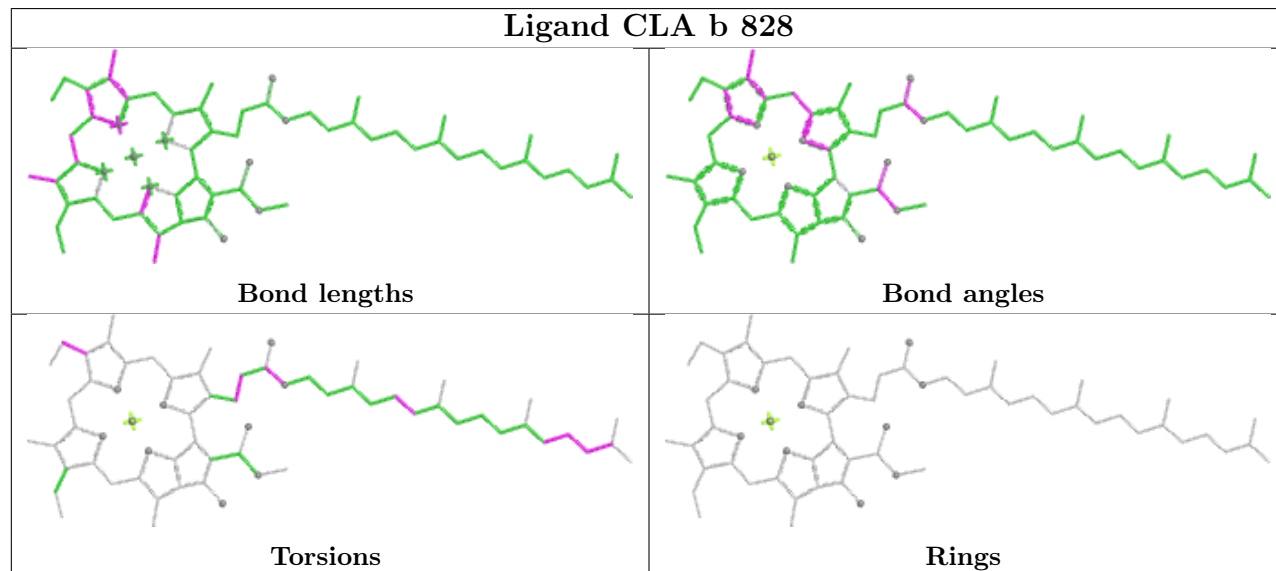


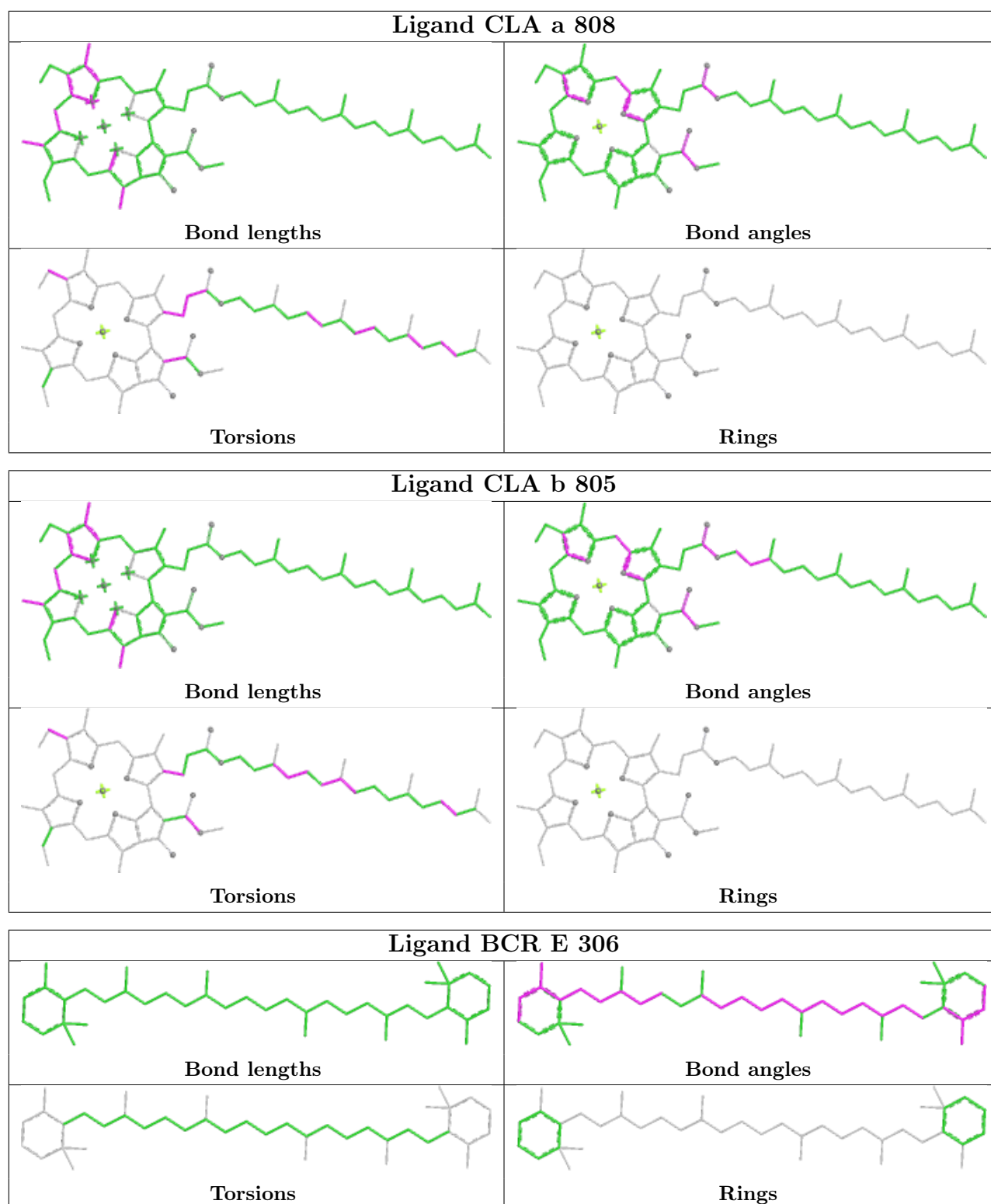
Ligand CLA B 307**Ligand CLA b 811**

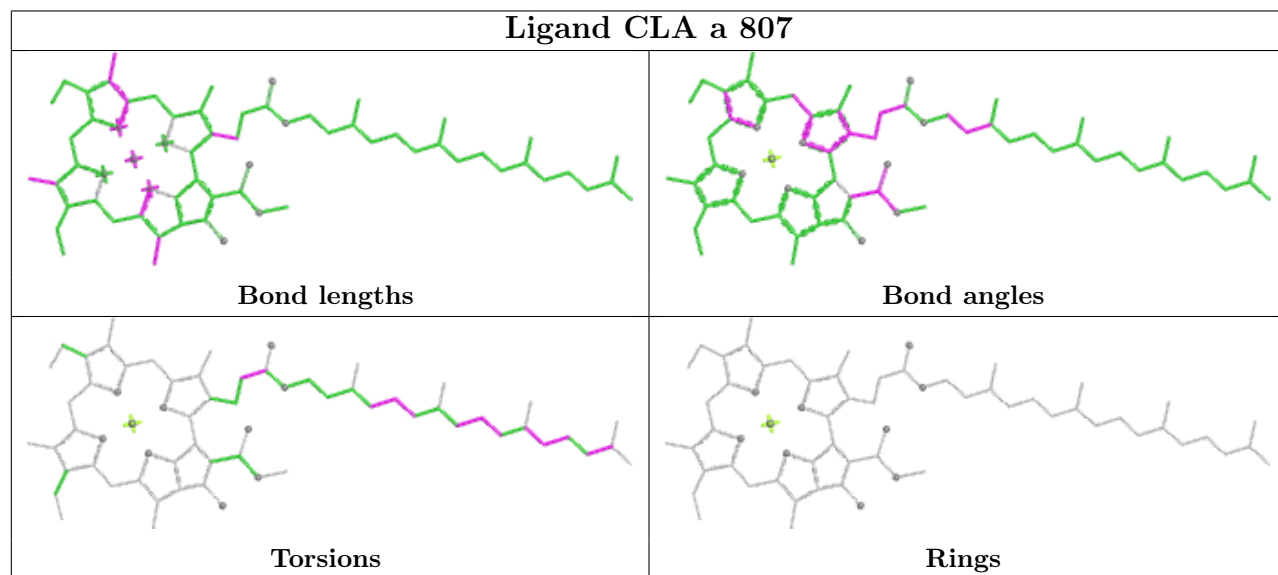
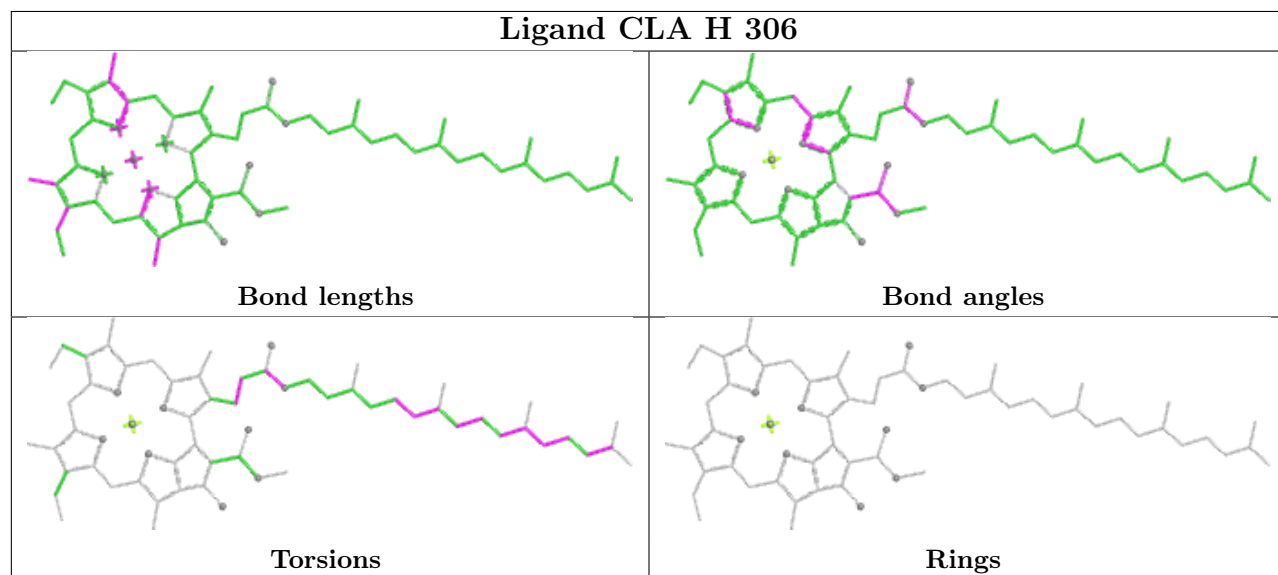
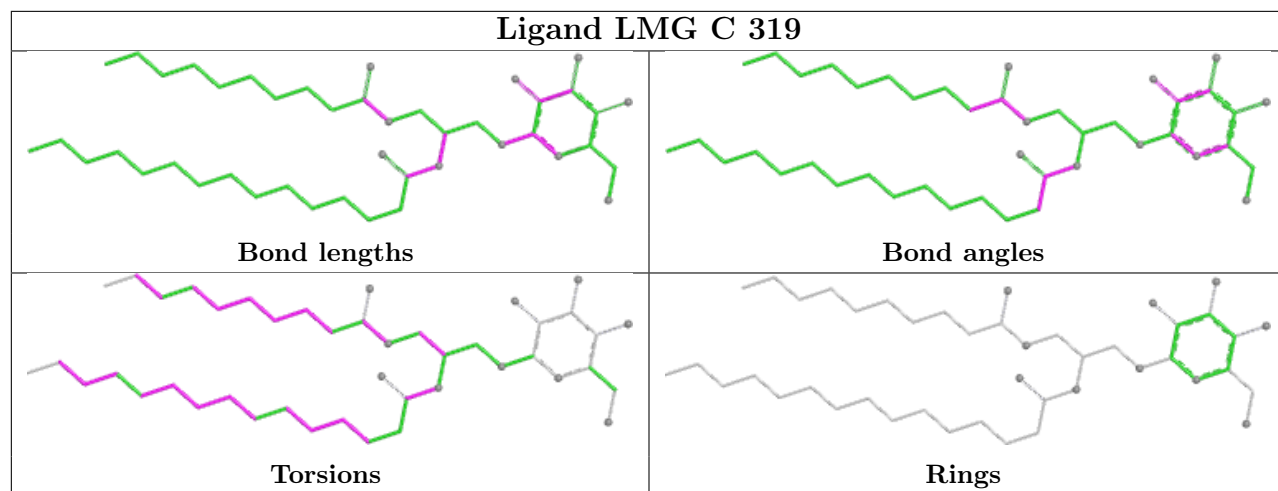
Ligand CLA 1 202

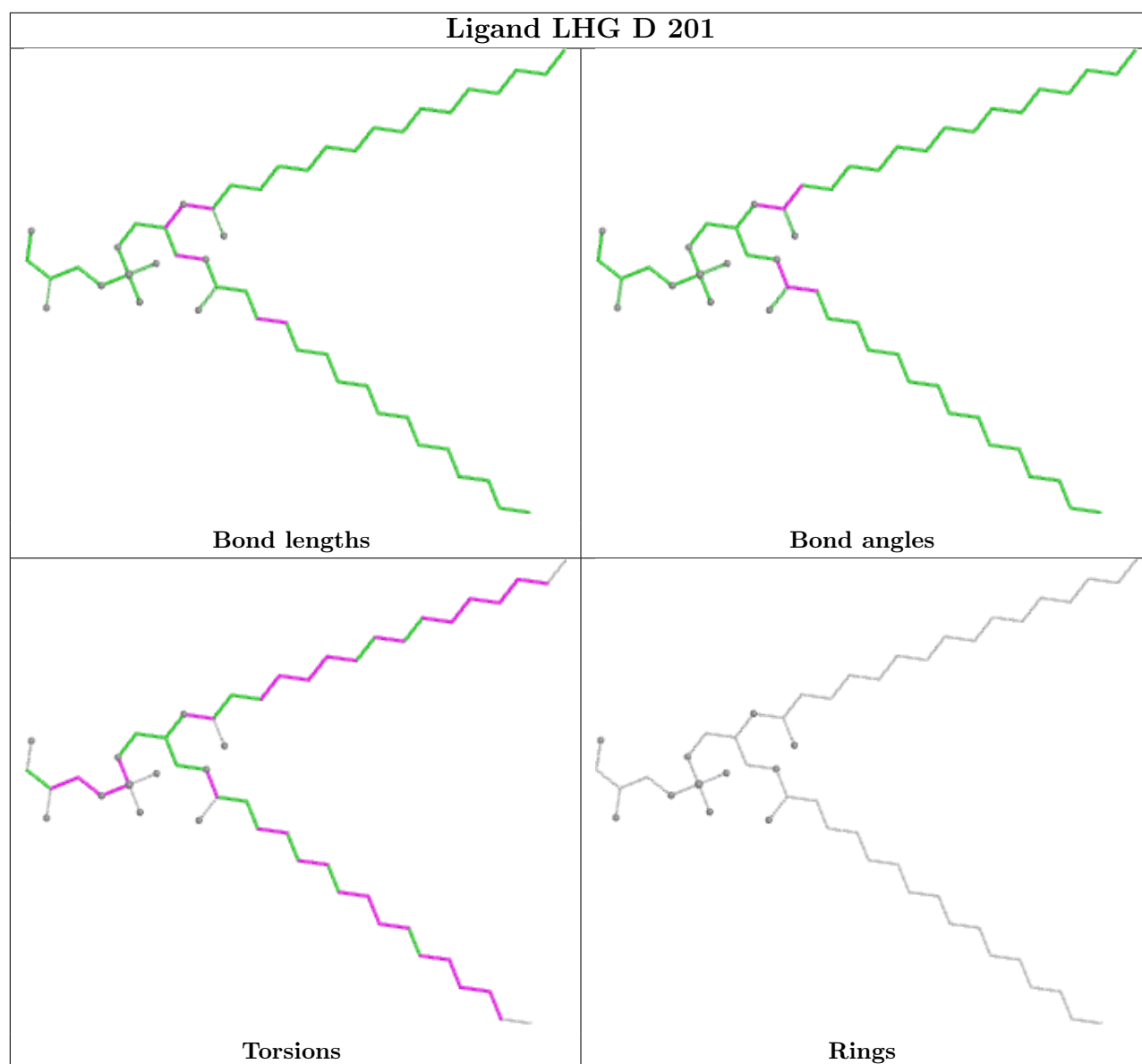


Ligand CLA b 828

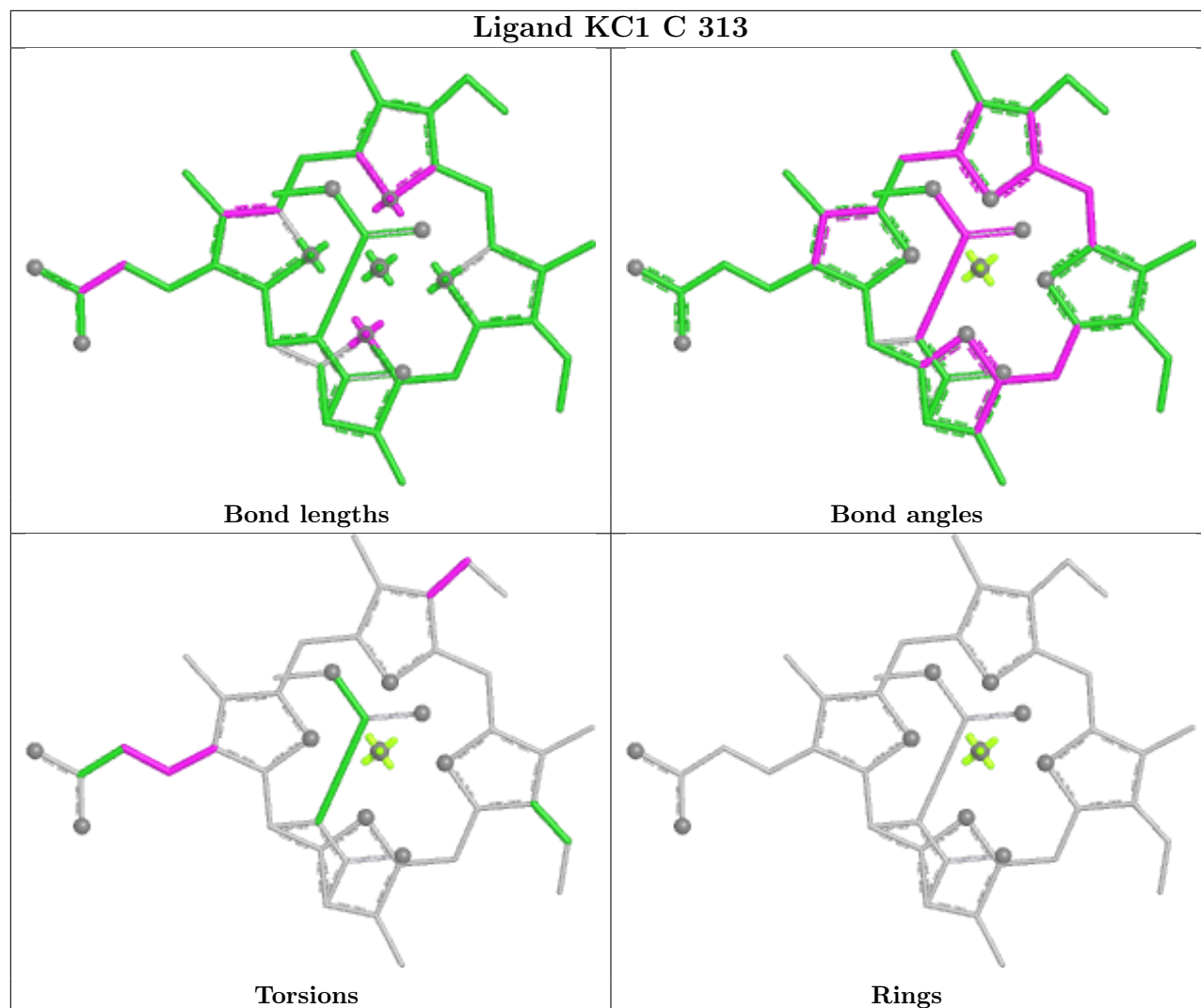




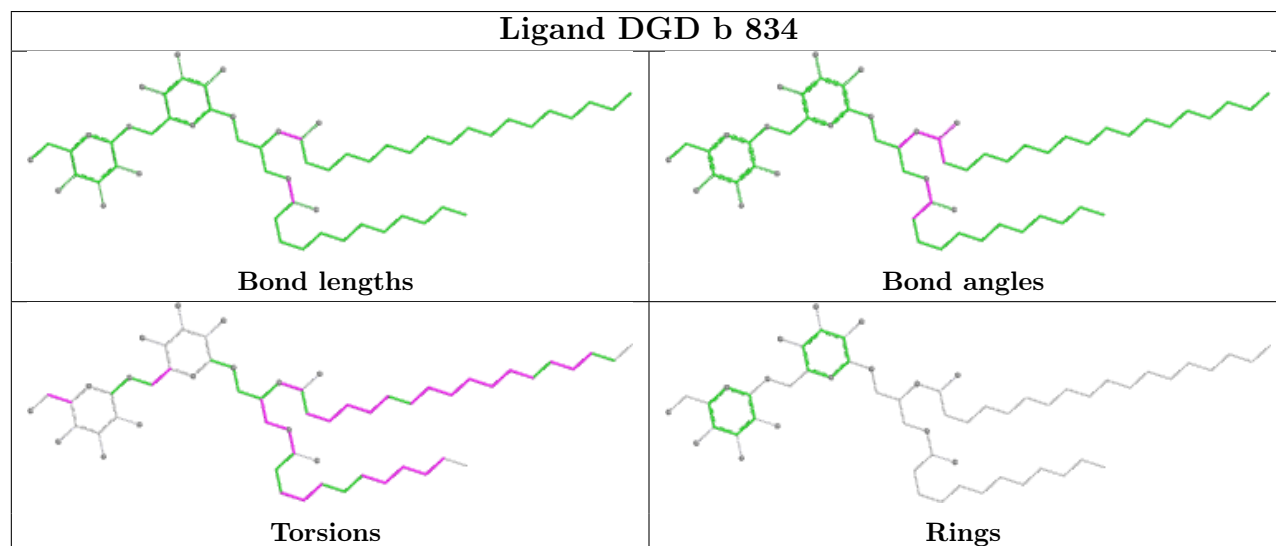




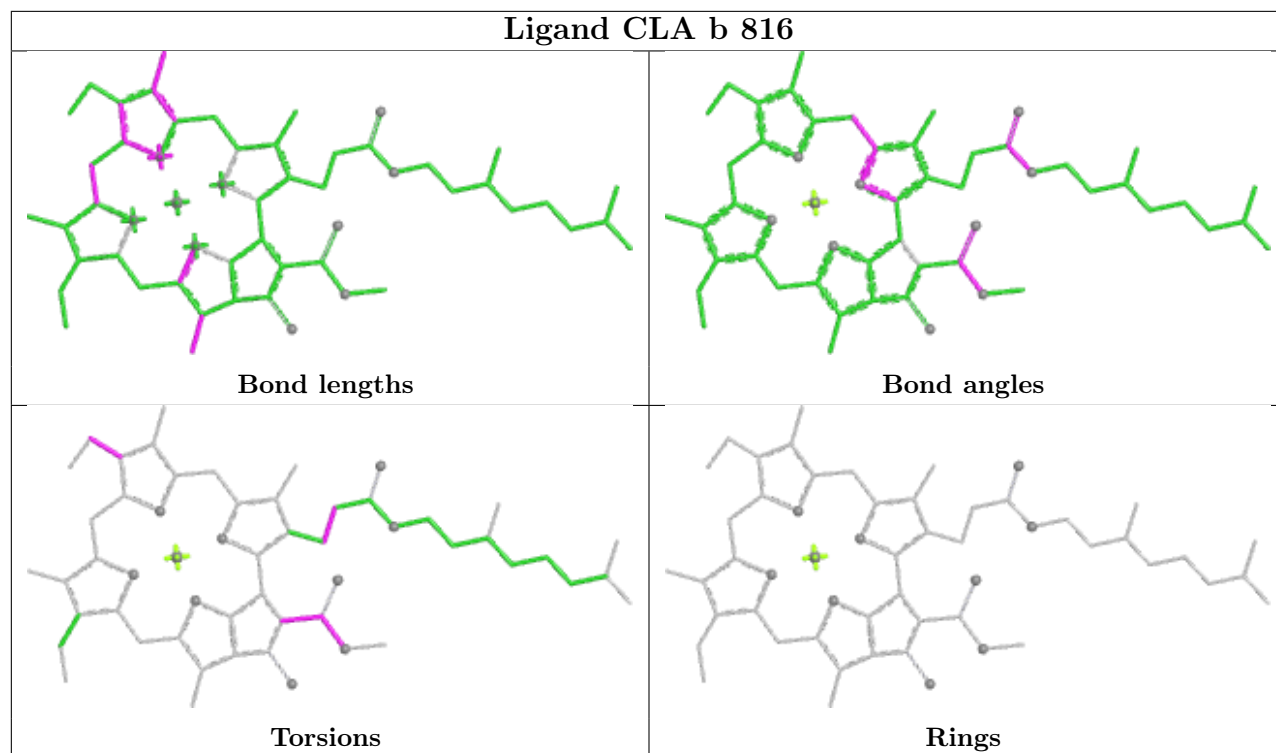
Ligand KC1 C 313



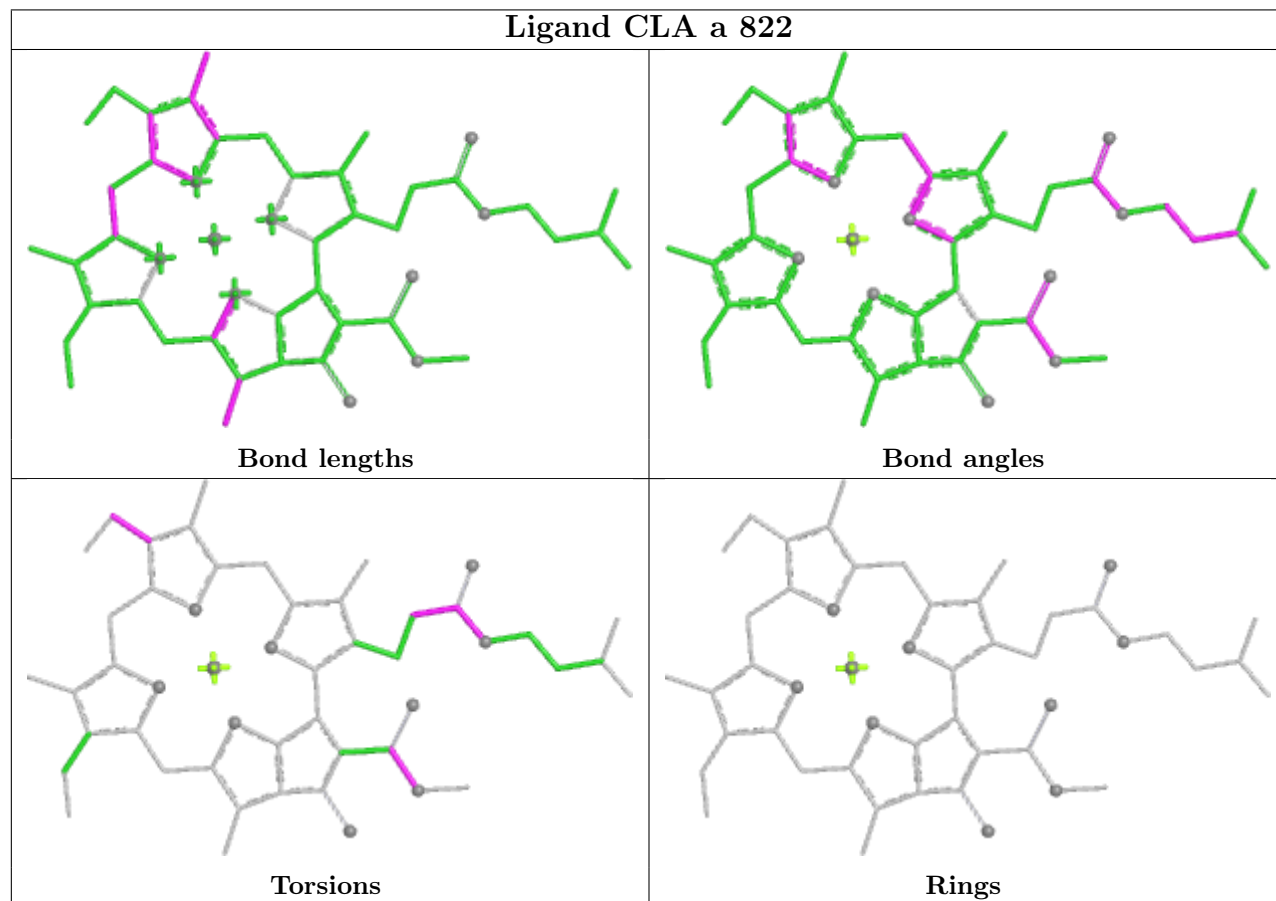
Ligand DGD b 834

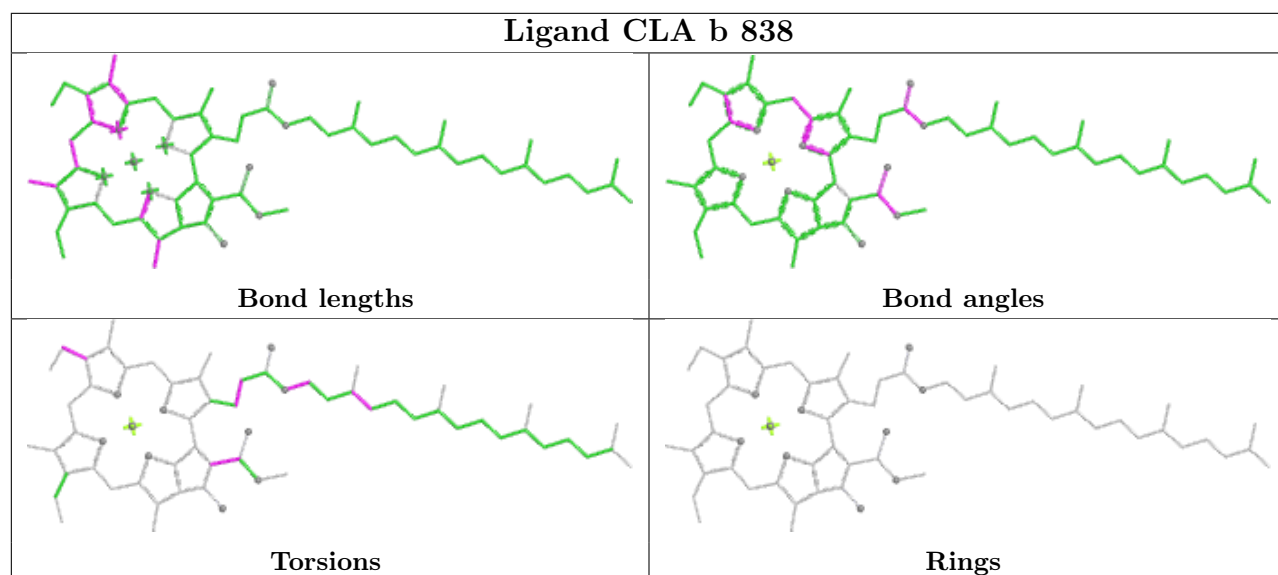
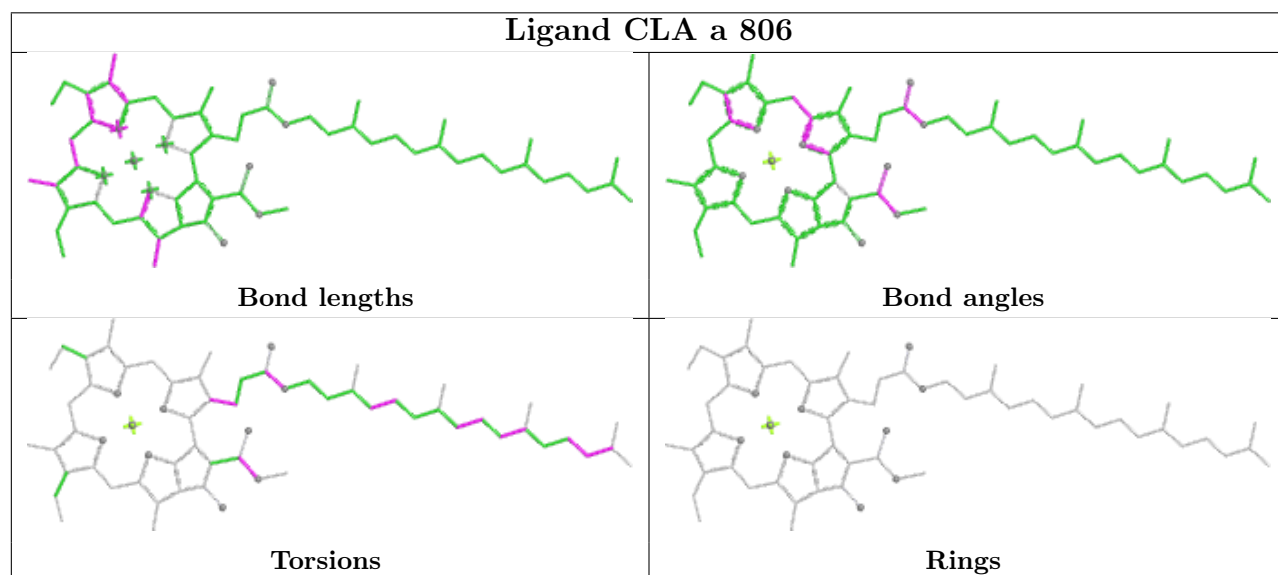
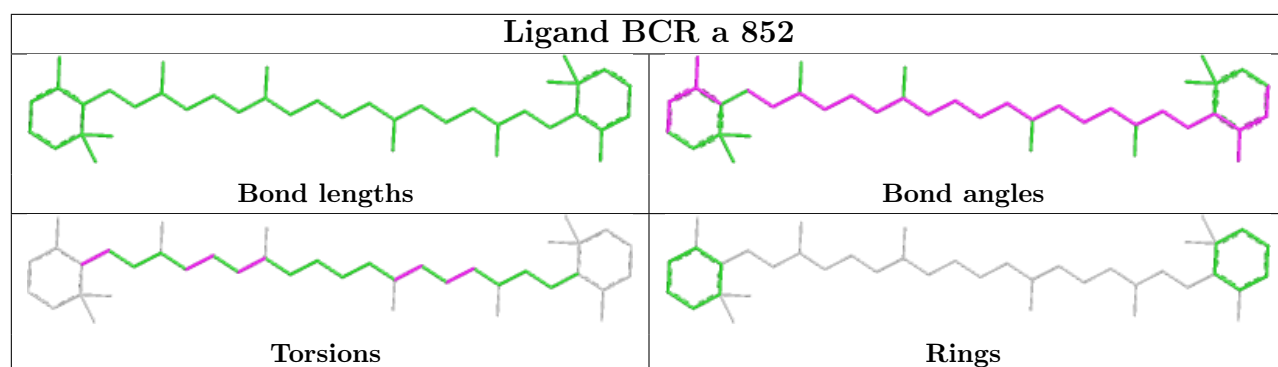


Ligand CLA b 816

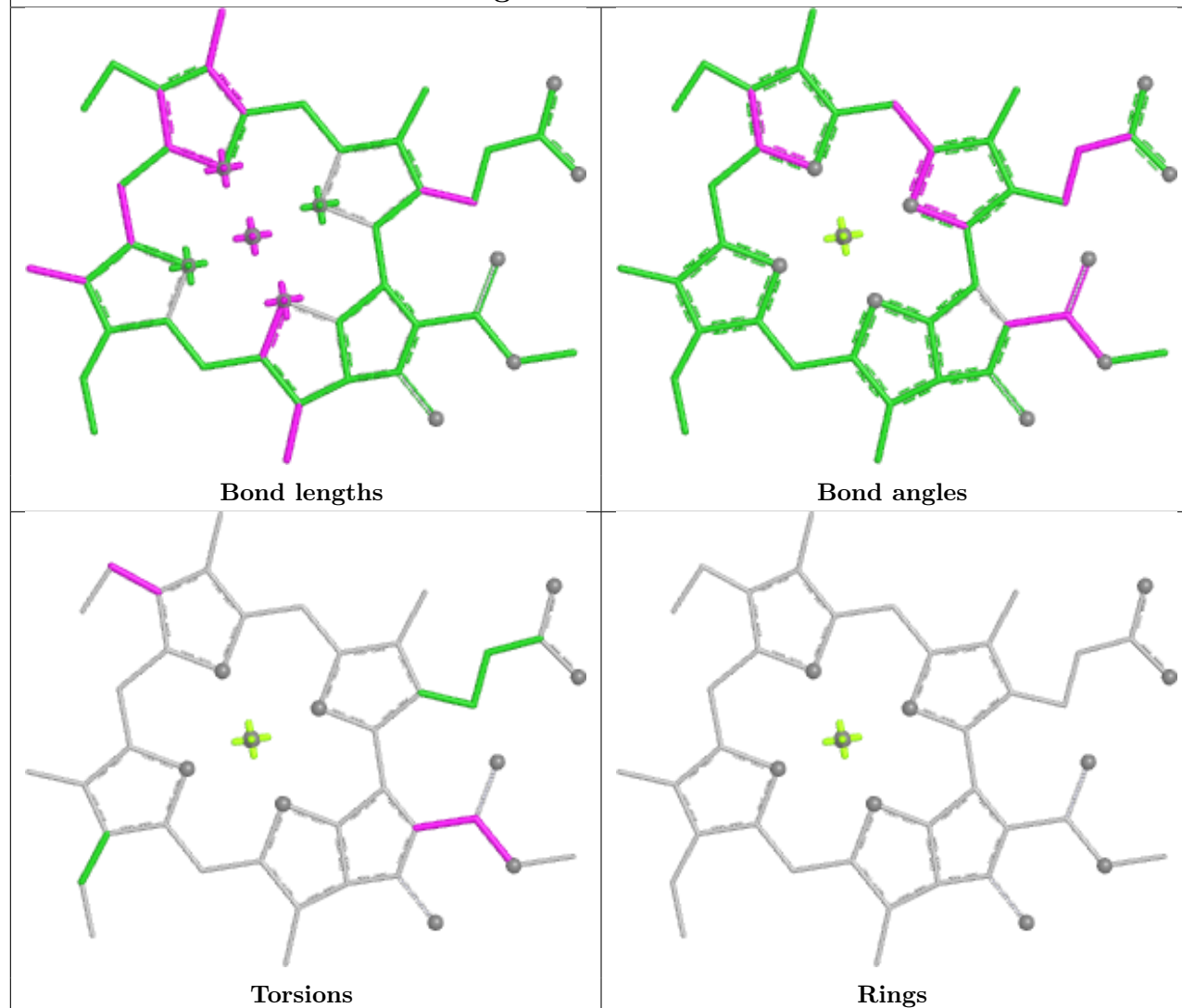


Ligand CLA a 822

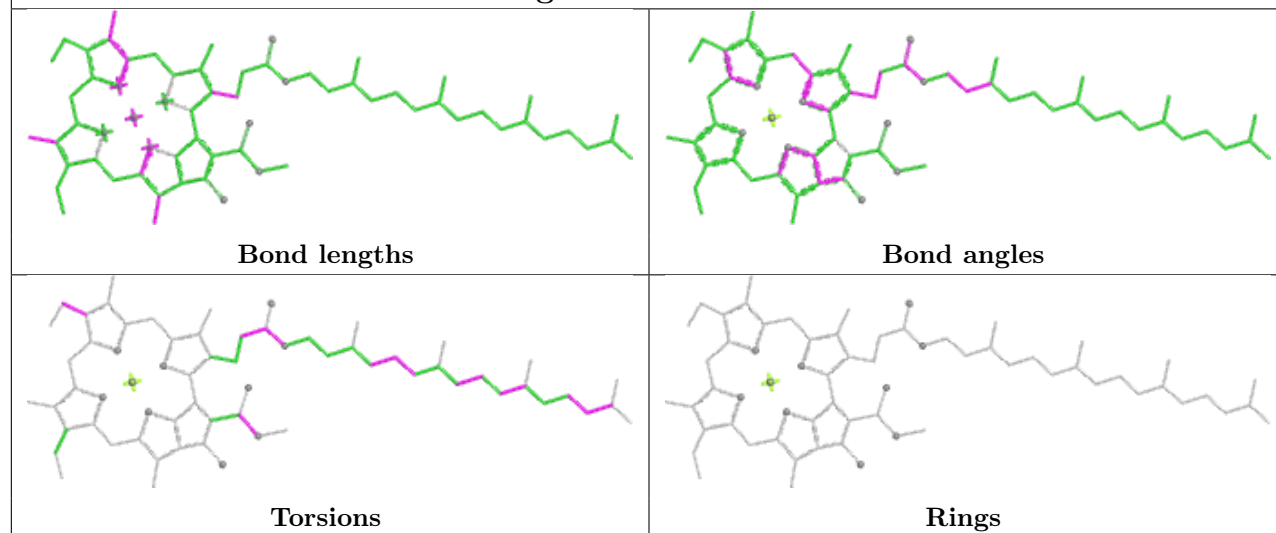


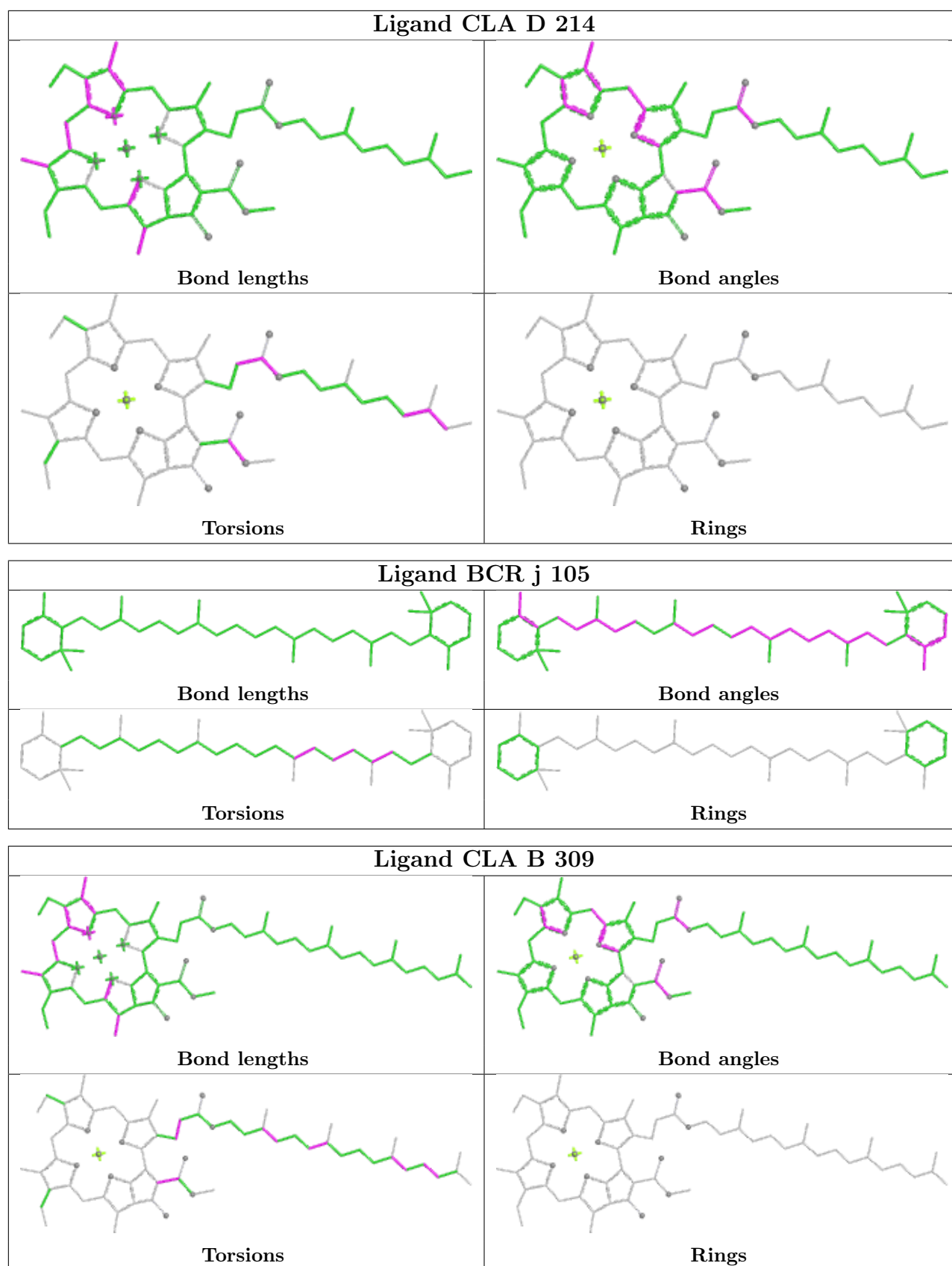


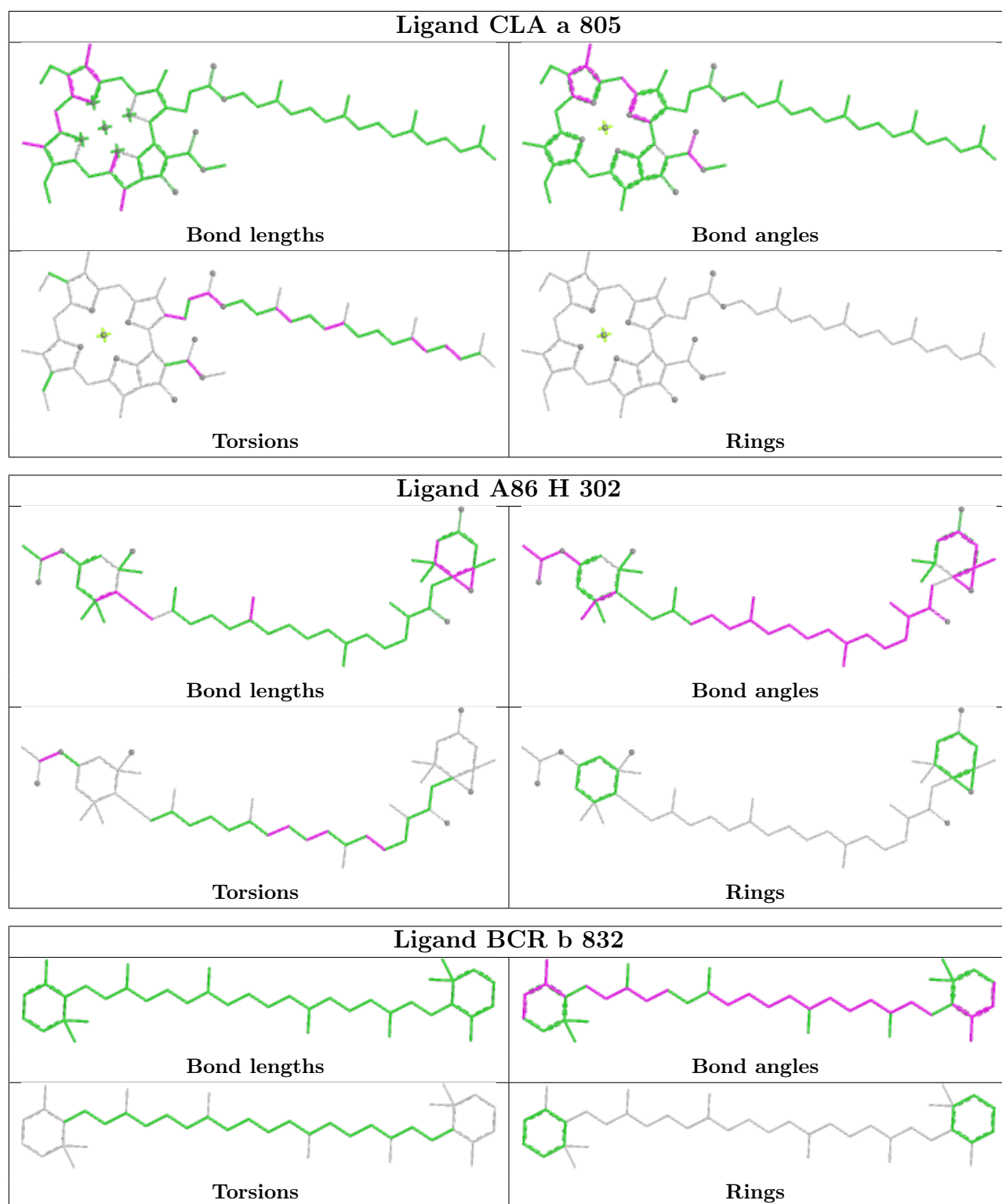
Ligand CLA E 314



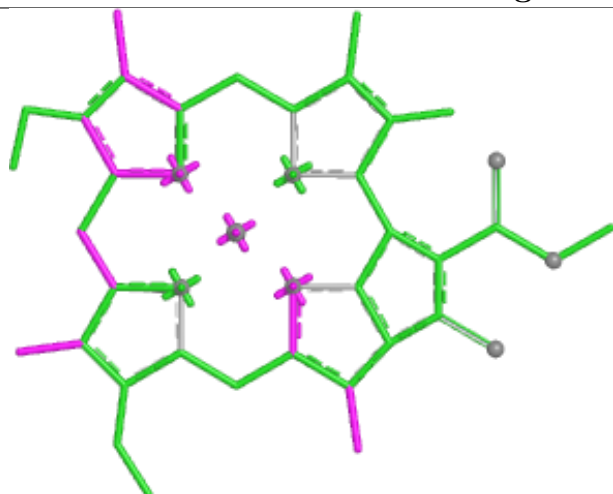
Ligand CLA a 801



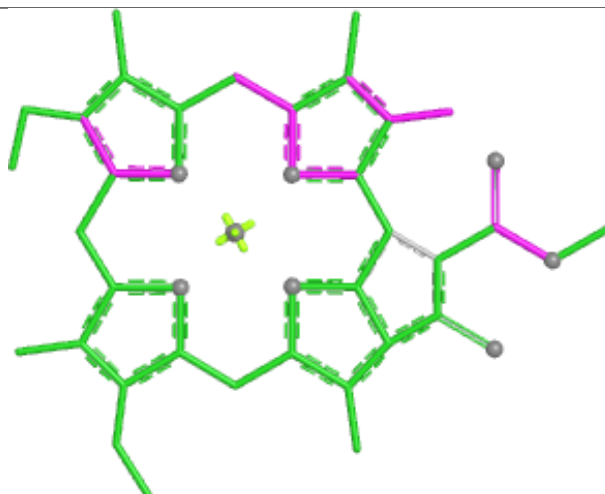




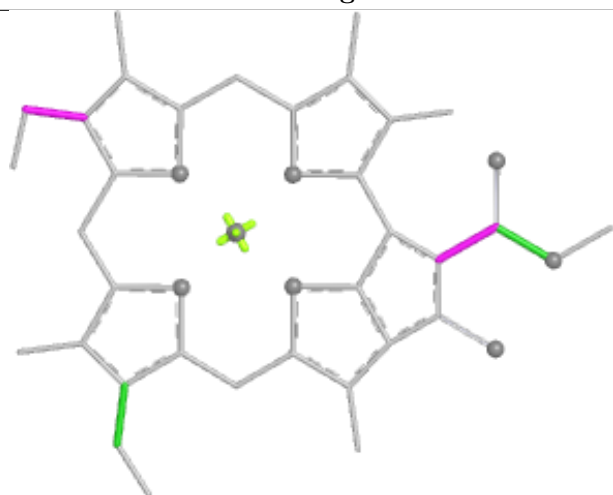
Ligand CLA C 316



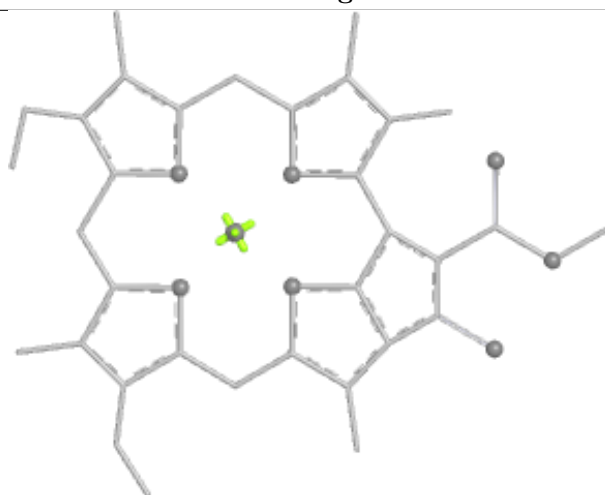
Bond lengths



Bond angles

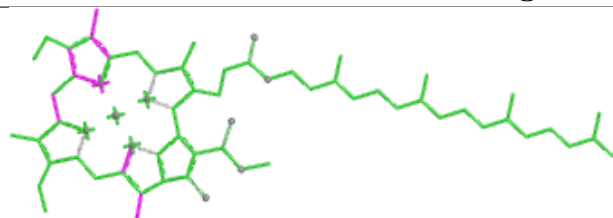


Torsions

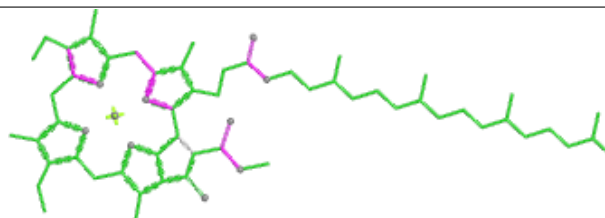


Rings

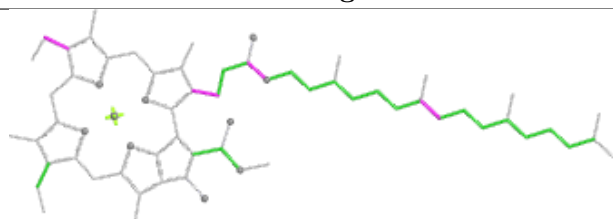
Ligand CLA a 846



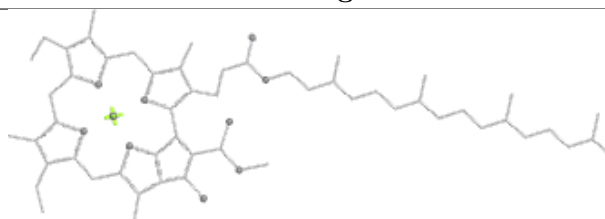
Bond lengths



Bond angles

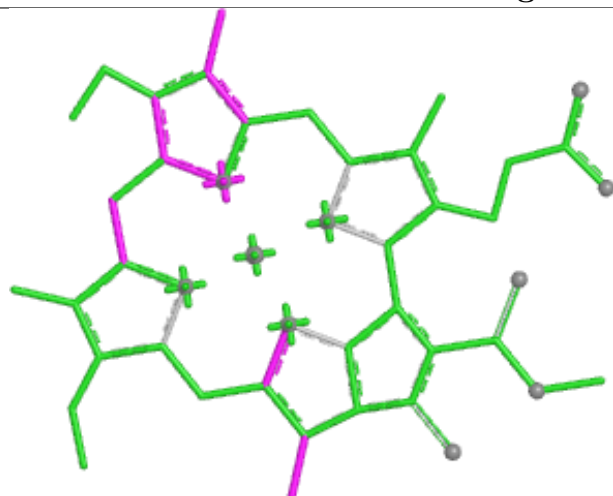


Torsions

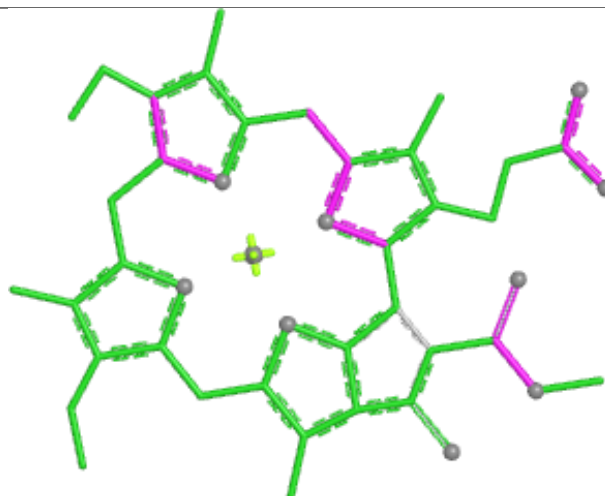


Rings

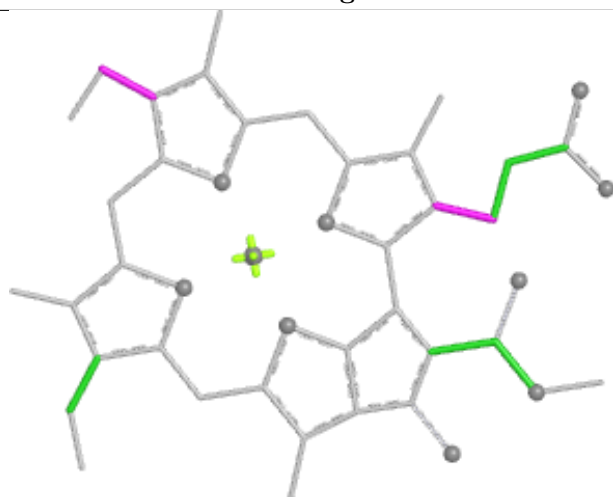
Ligand CLA b 825



Bond lengths



Bond angles

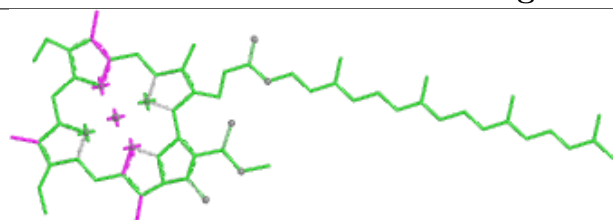


Torsions

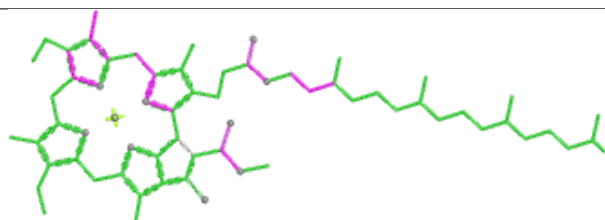


Rings

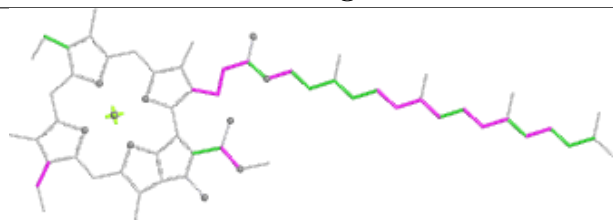
Ligand CLA b 824



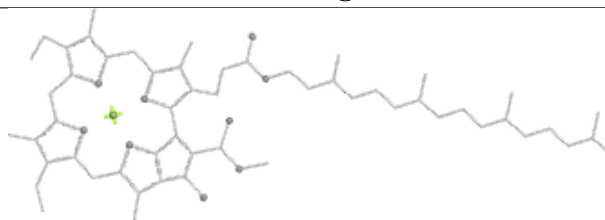
Bond lengths



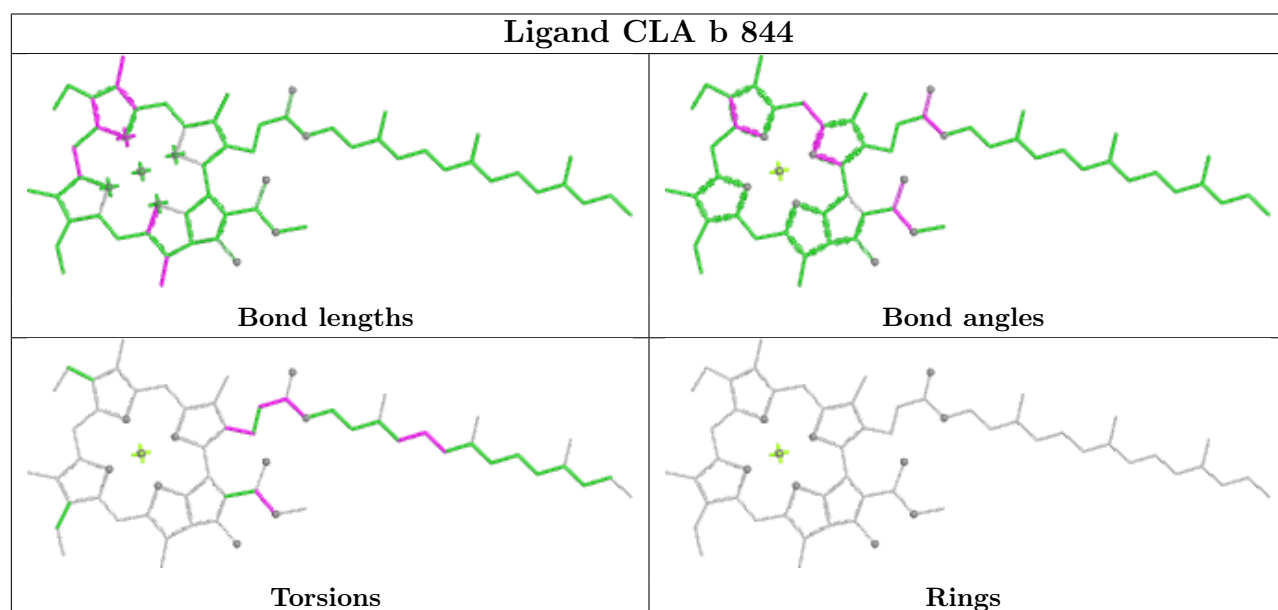
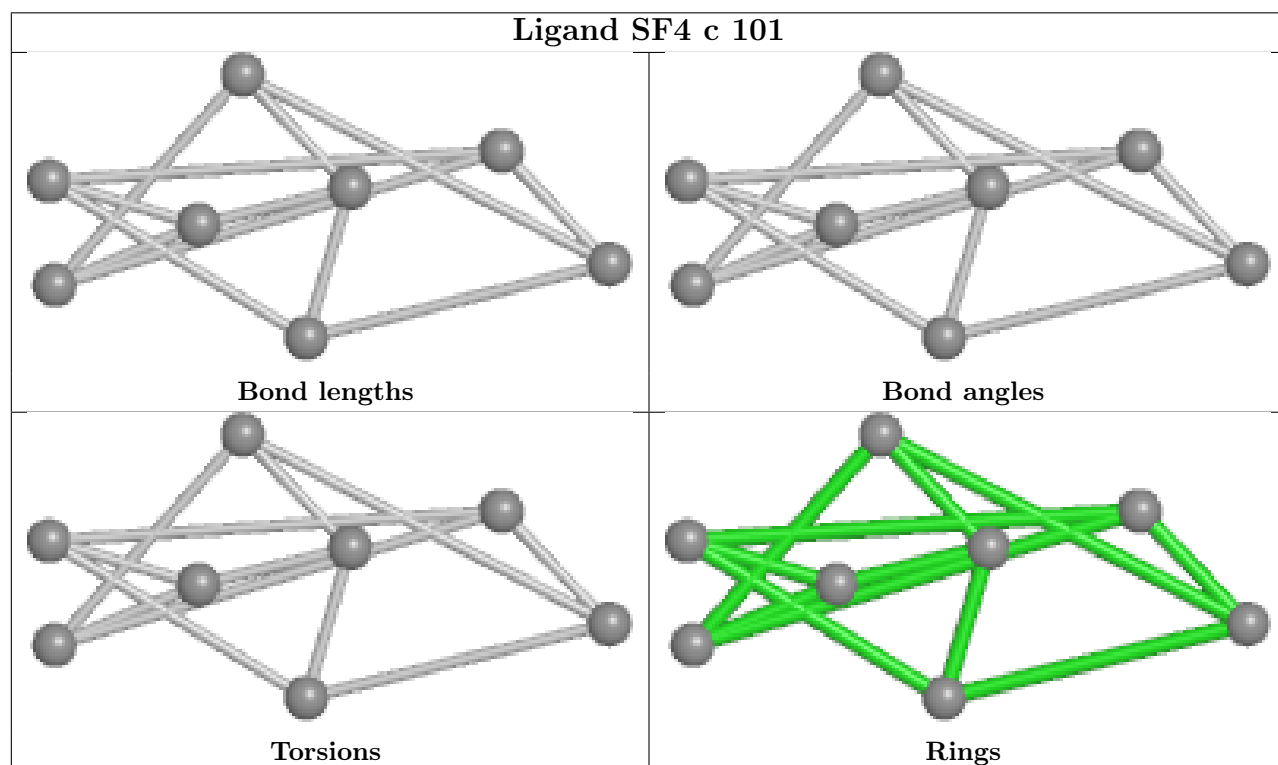
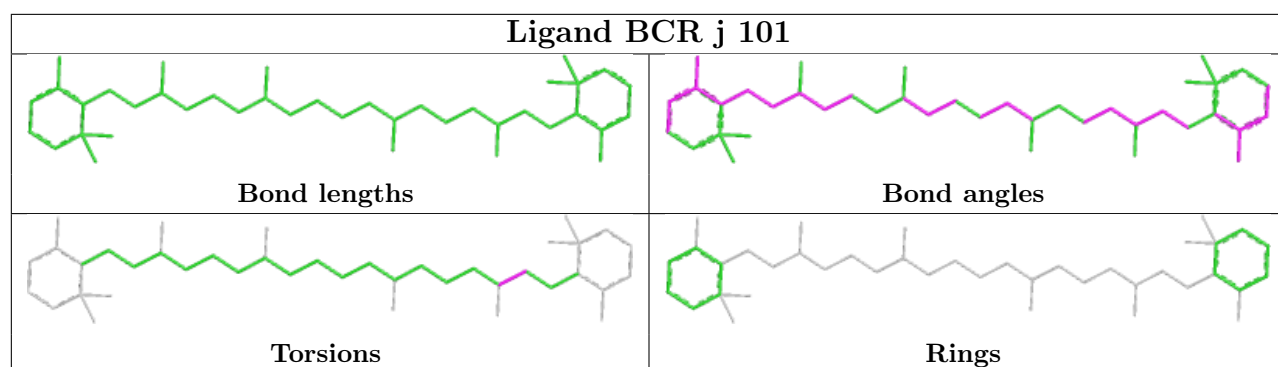
Bond angles



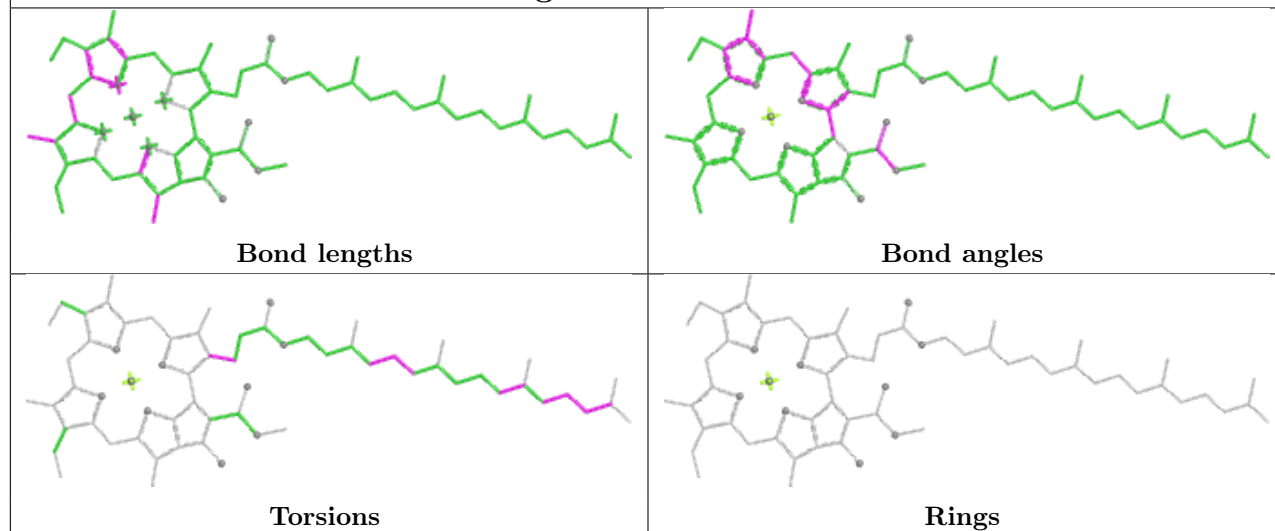
Torsions



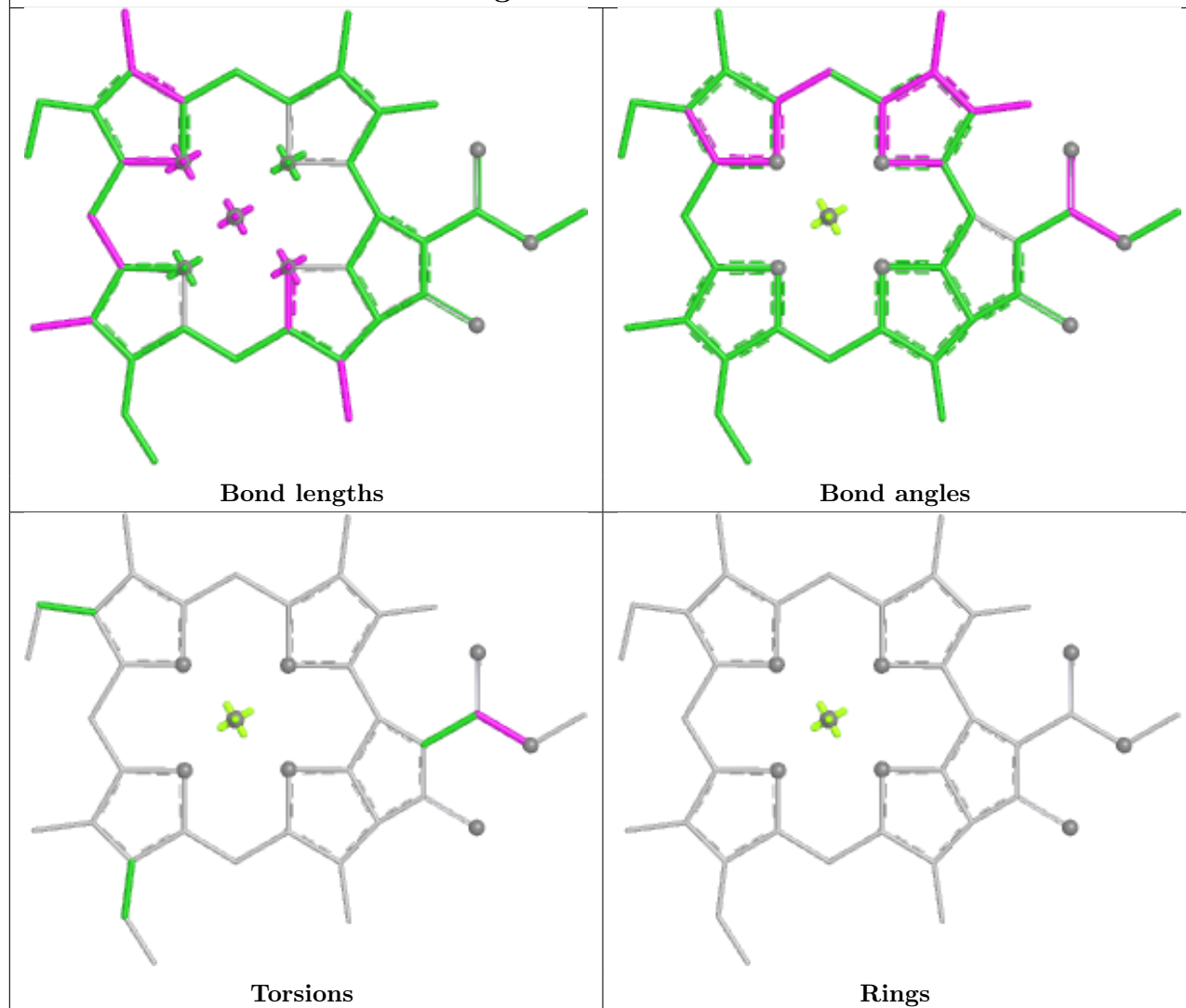
Rings



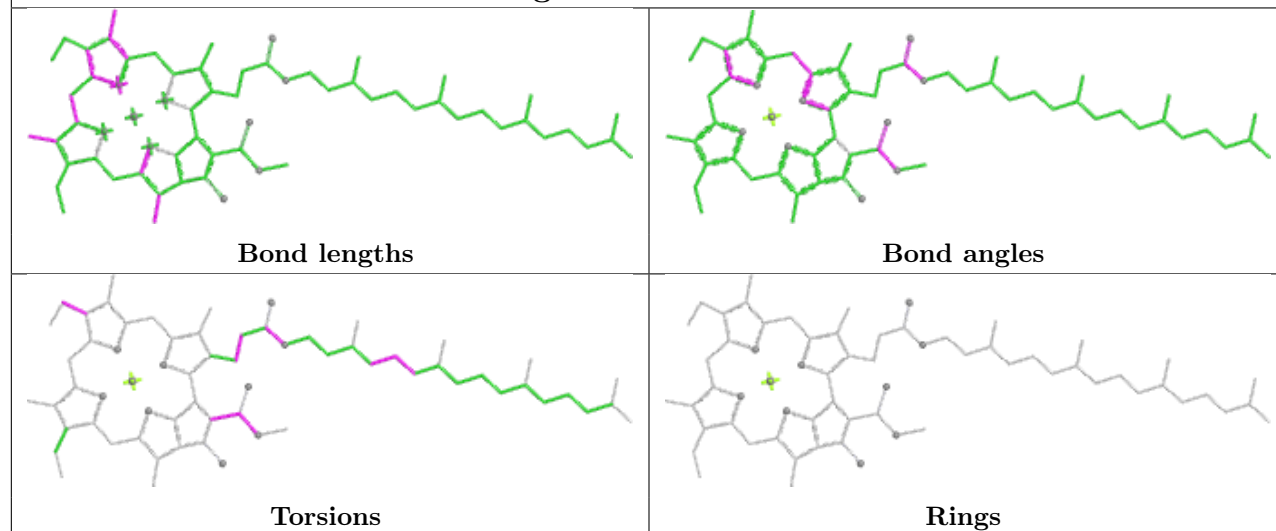
Ligand CLA a 843



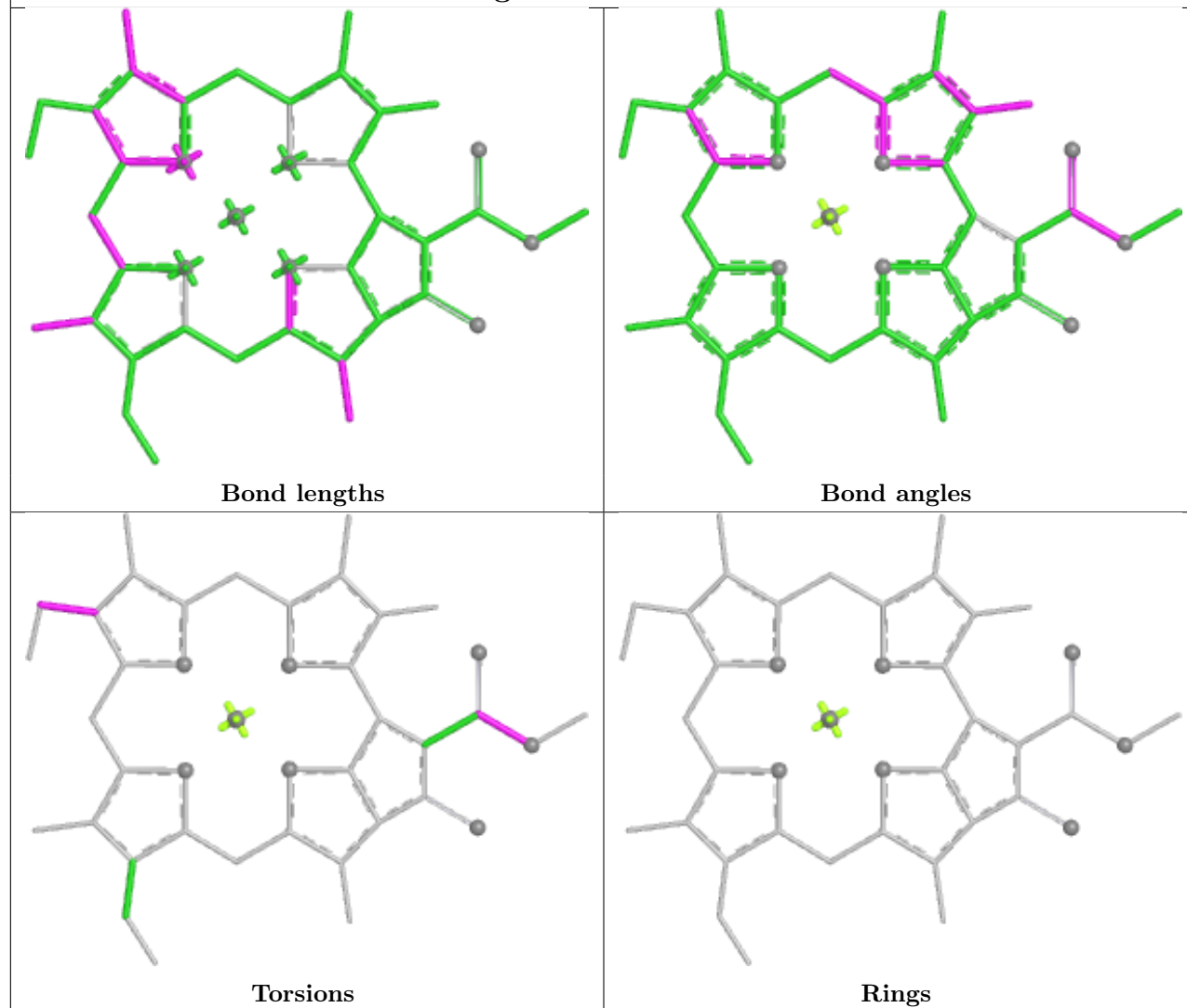
Ligand CLA B 314

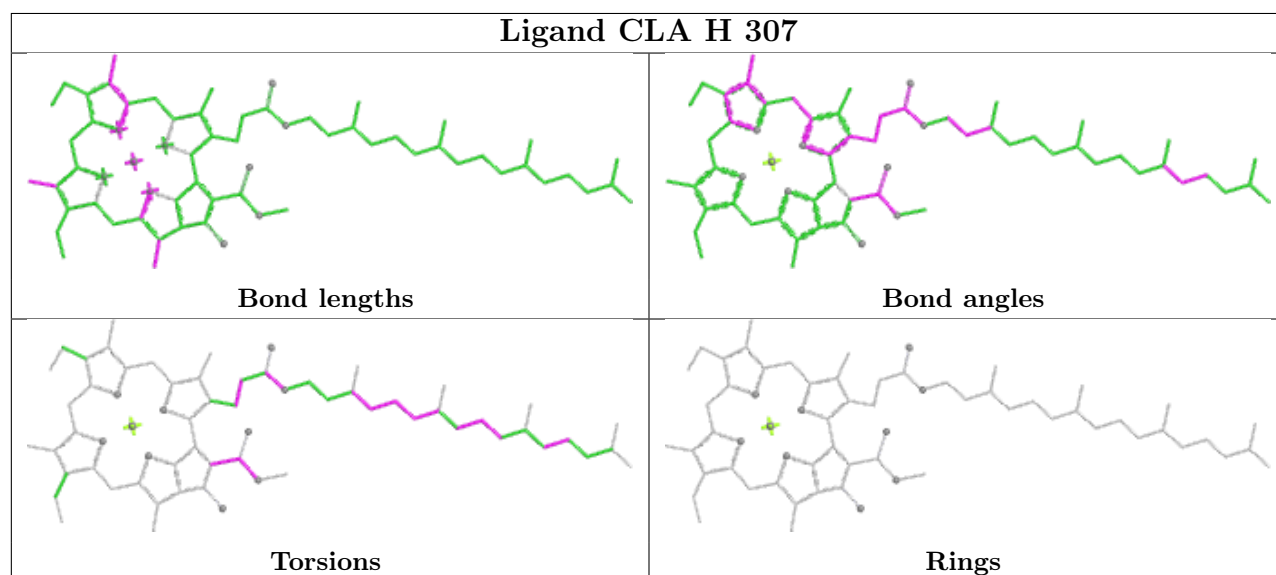
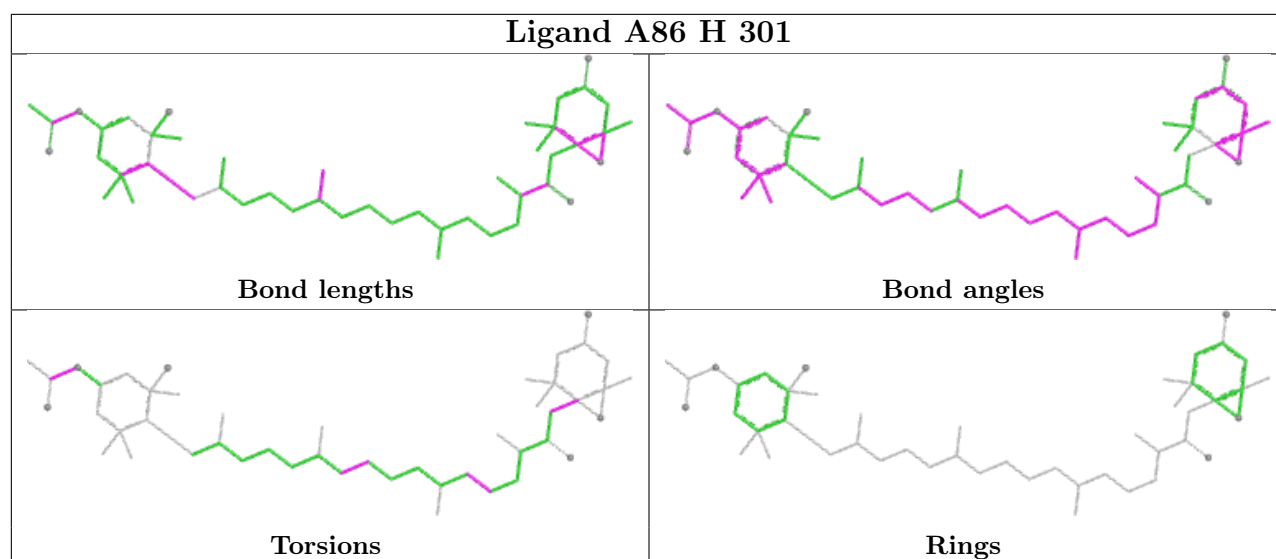


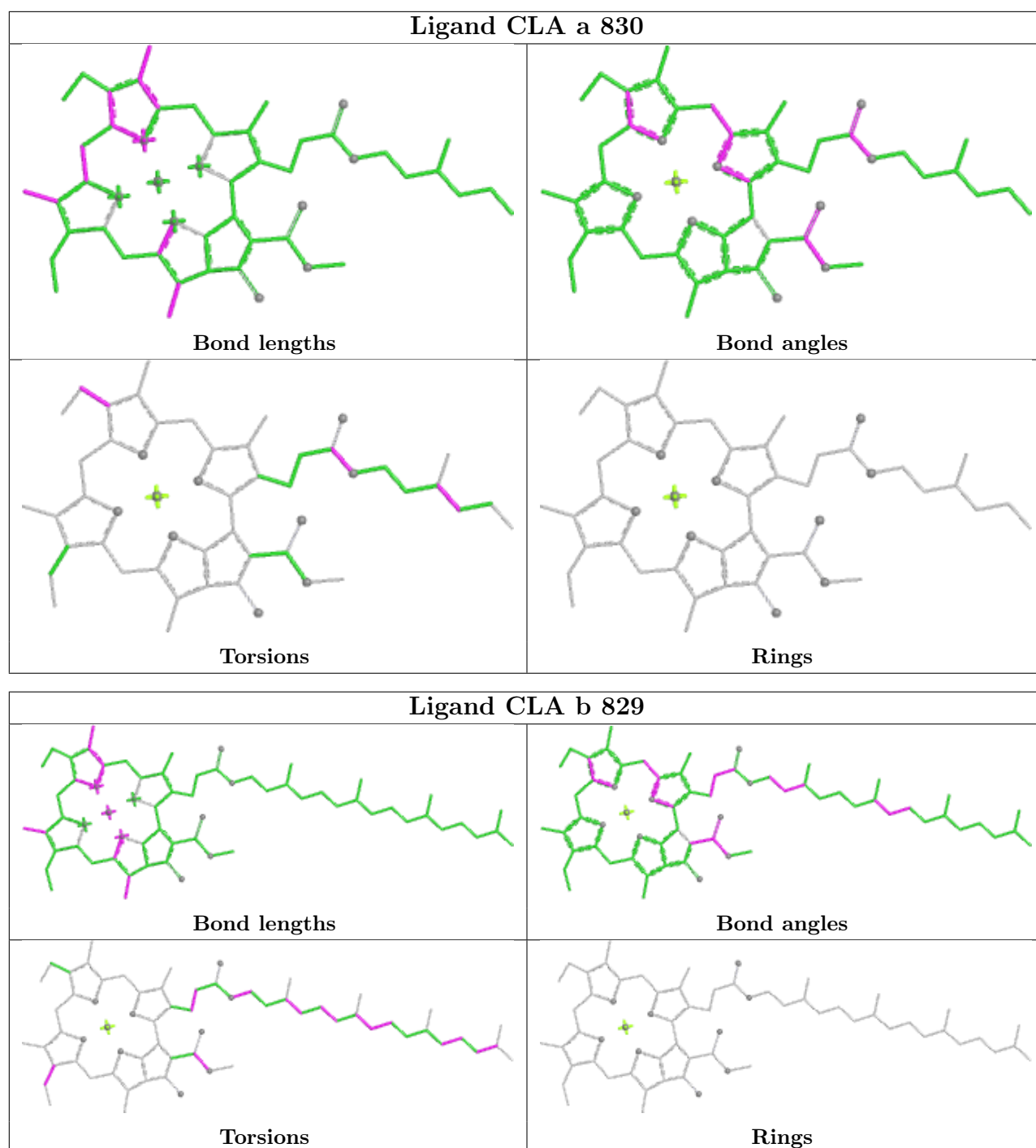
Ligand CLA E 315



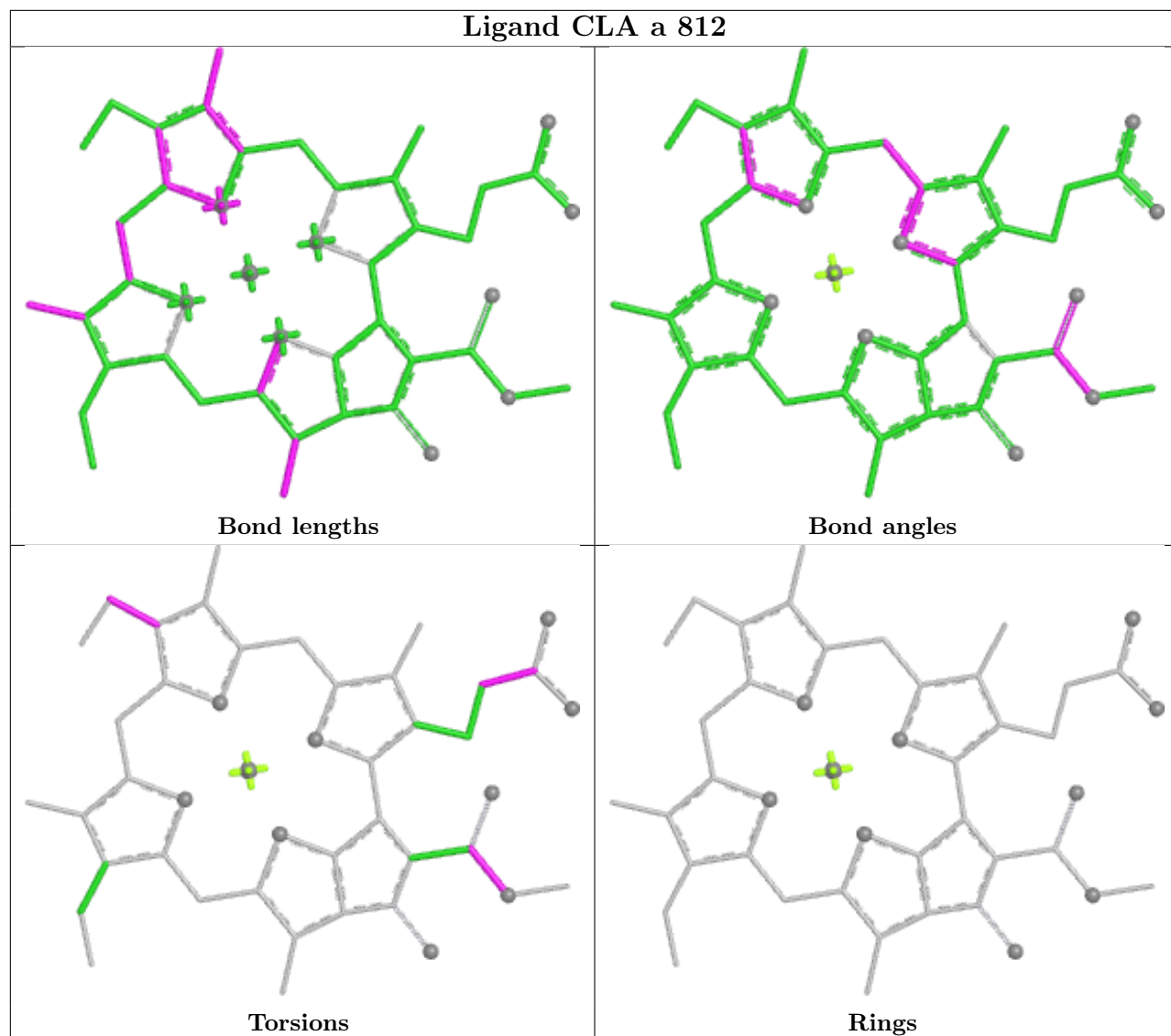
Ligand CLA D 215



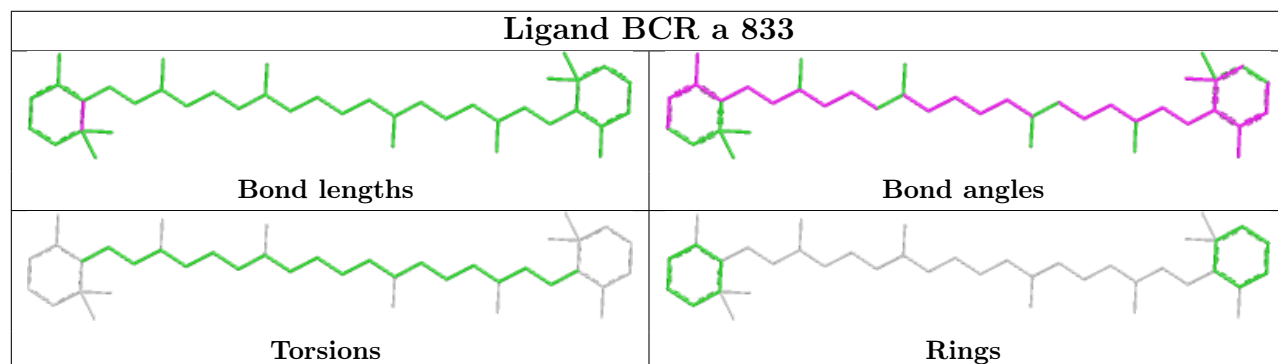




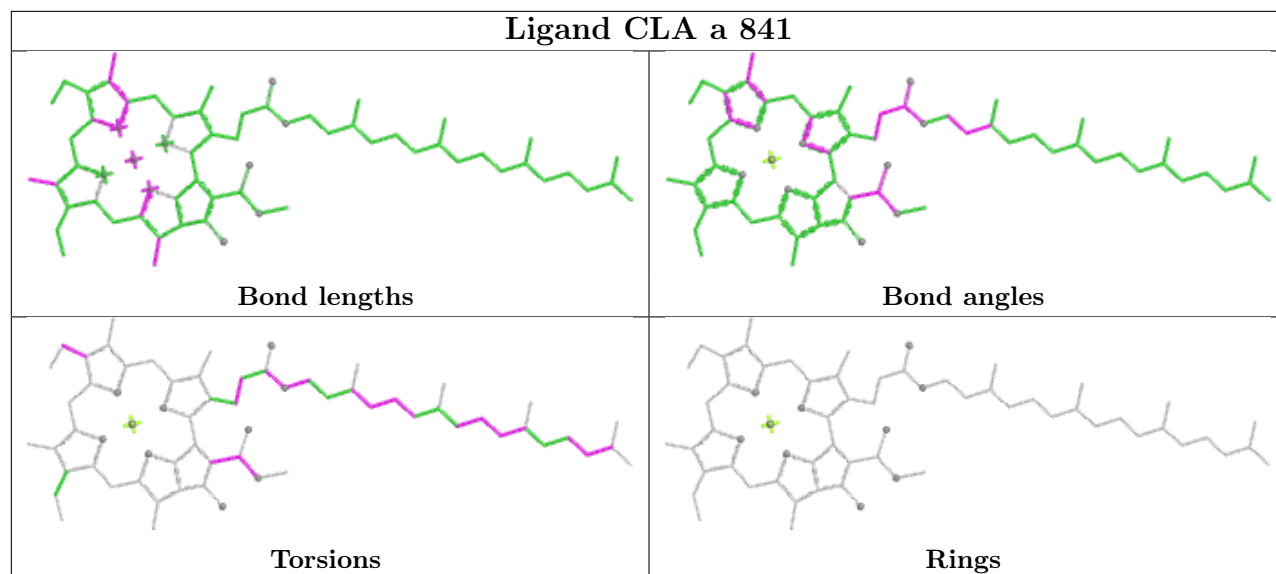
Ligand CLA a 812



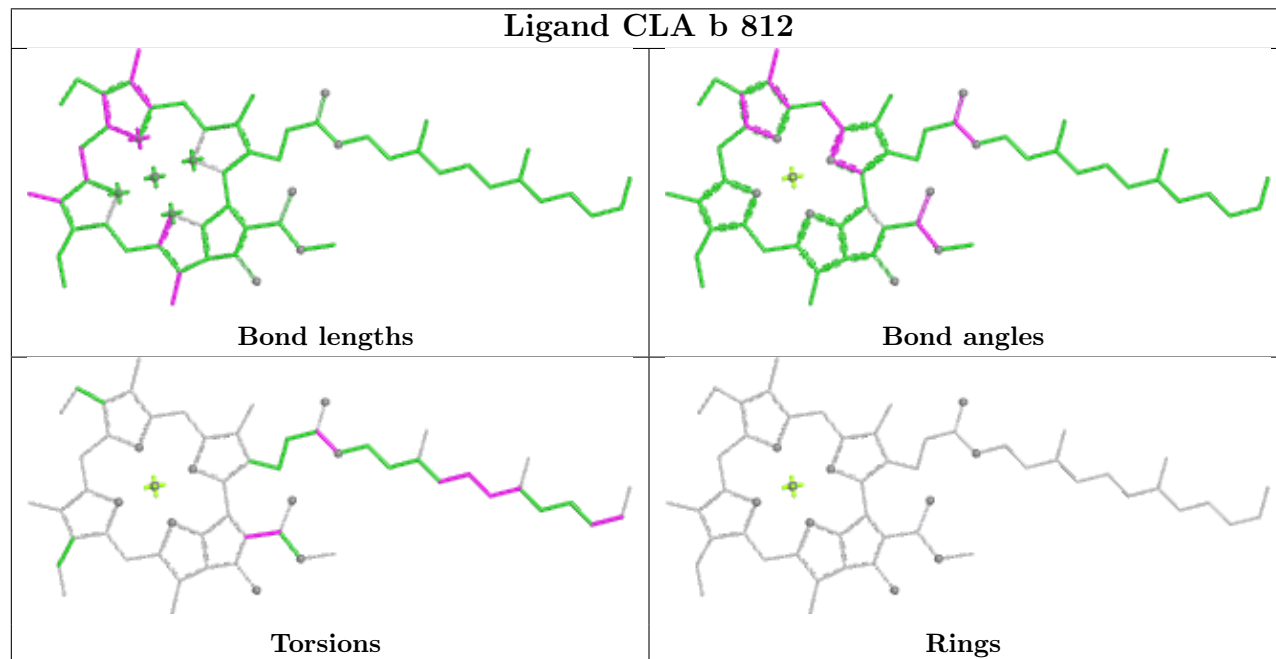
Ligand BCR a 833

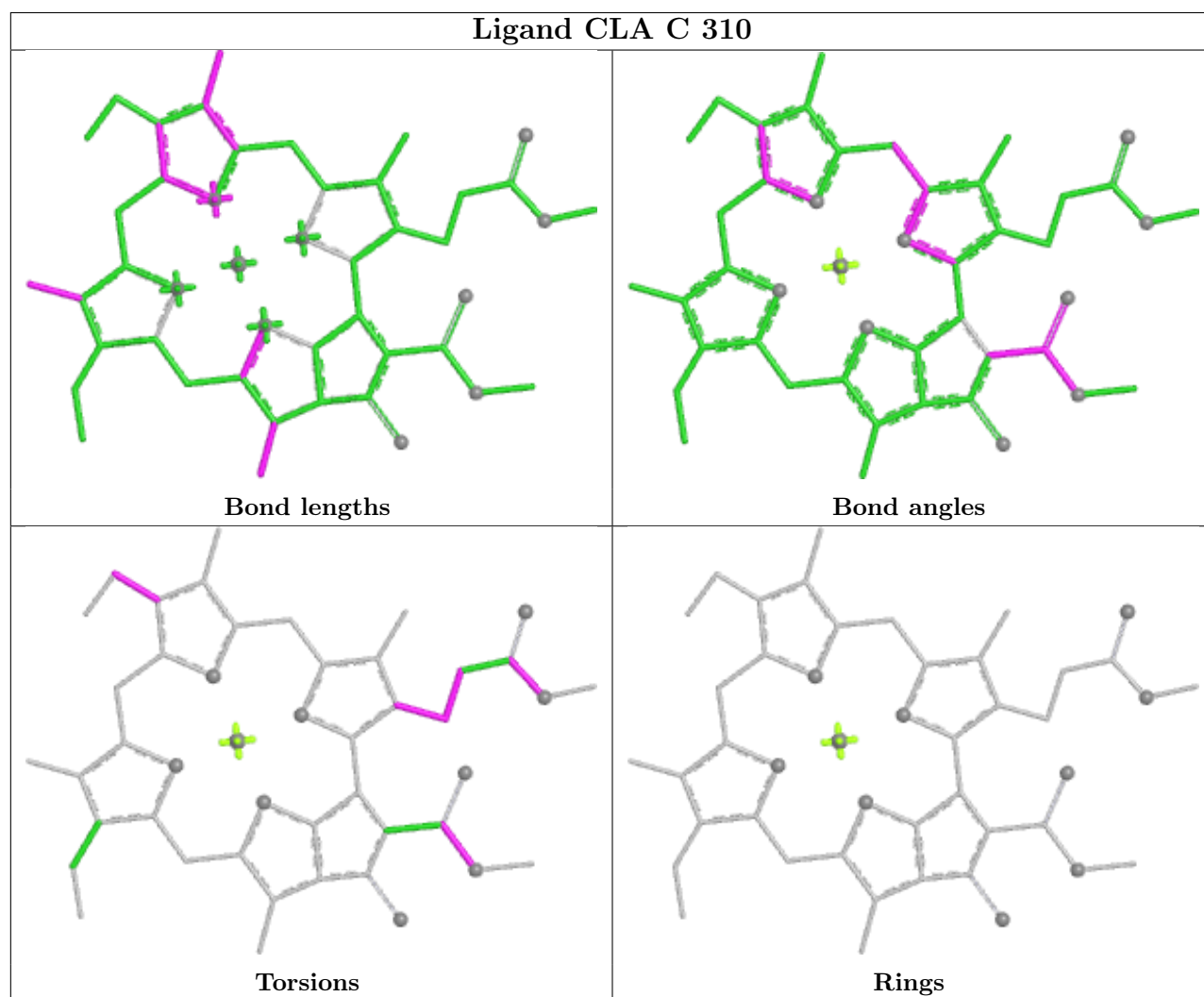
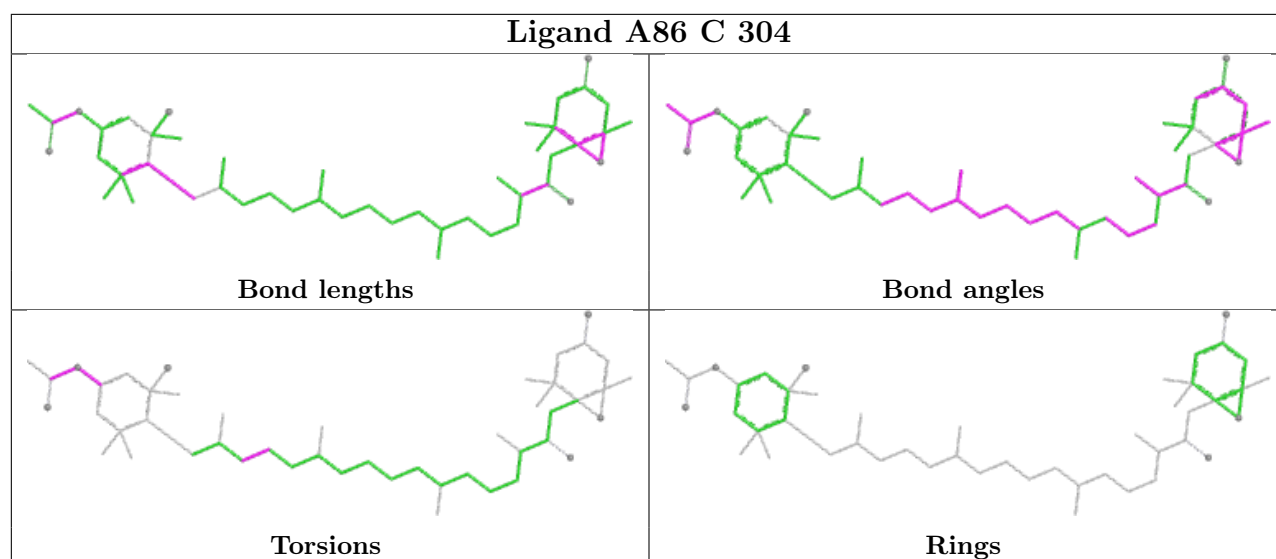


Ligand CLA a 841

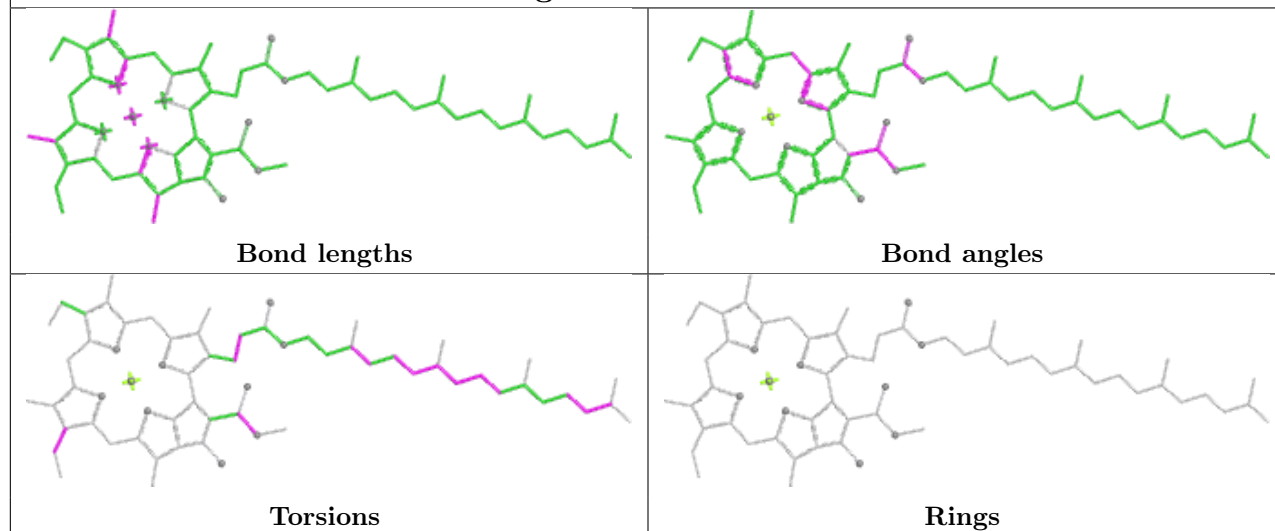


Ligand CLA b 812

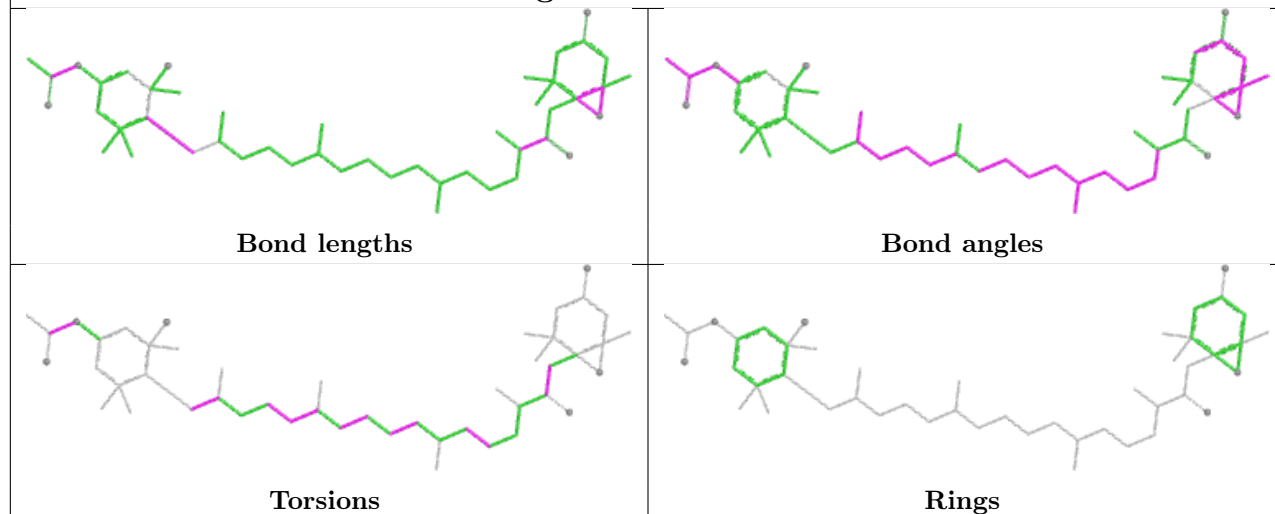




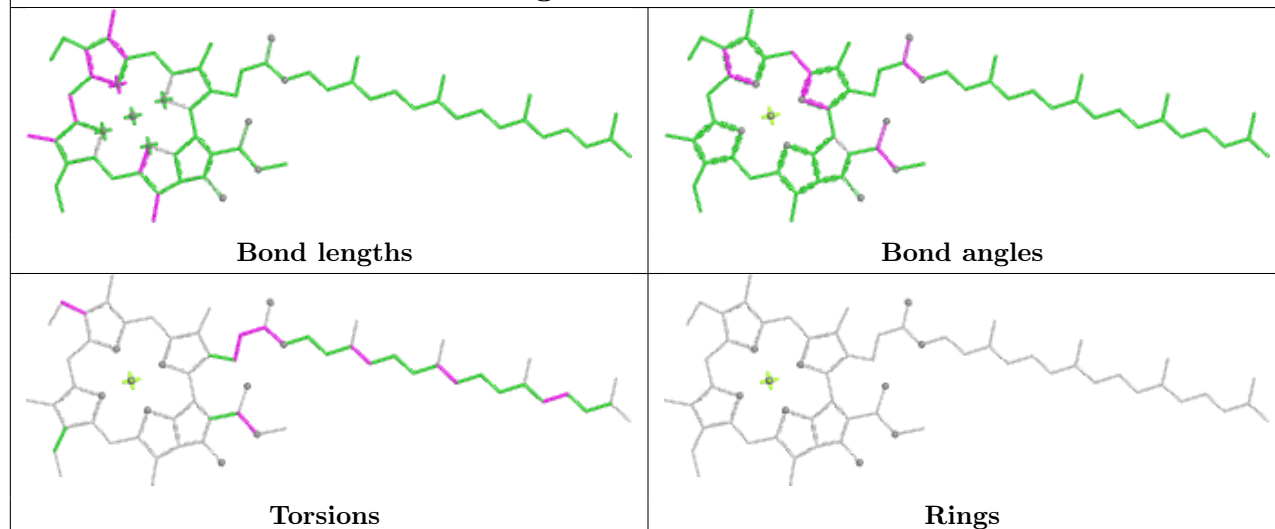
Ligand CLA a 854

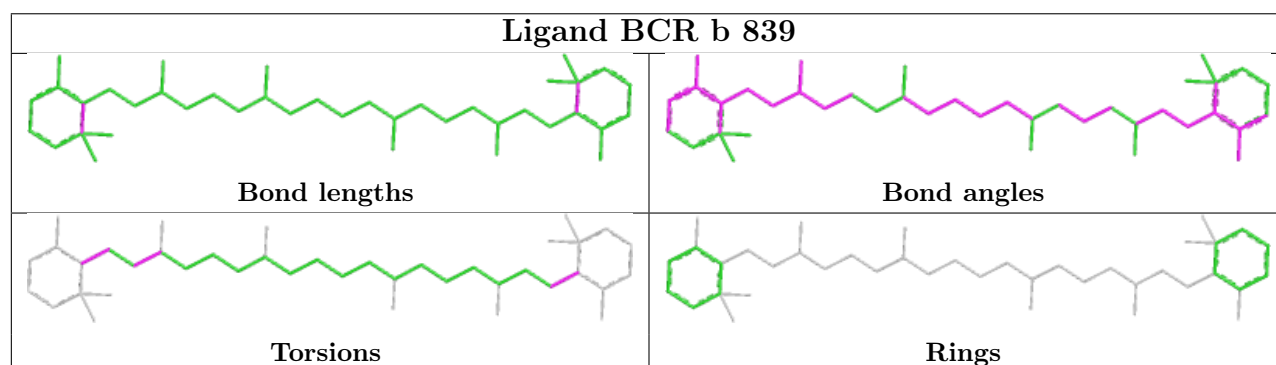
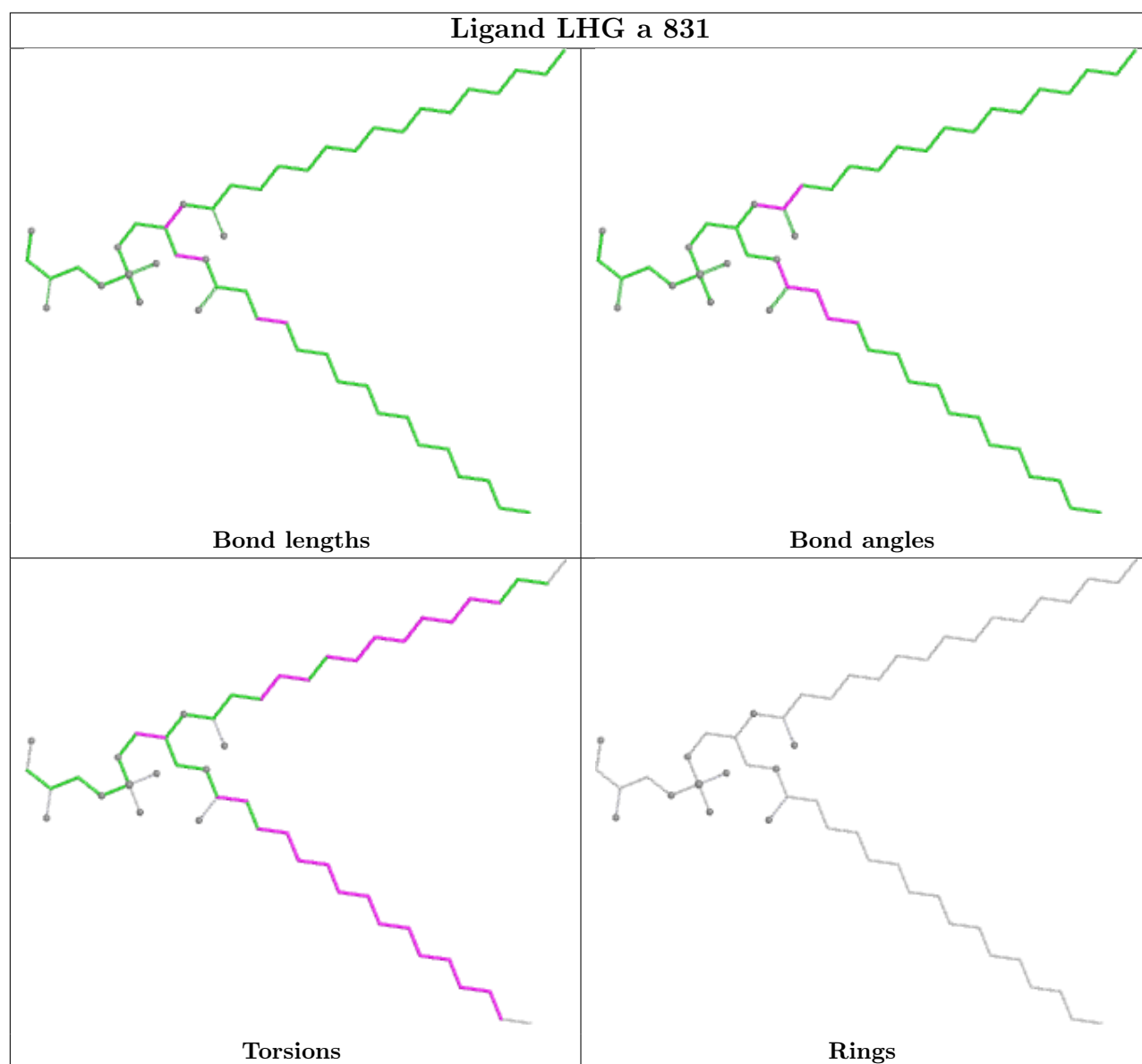


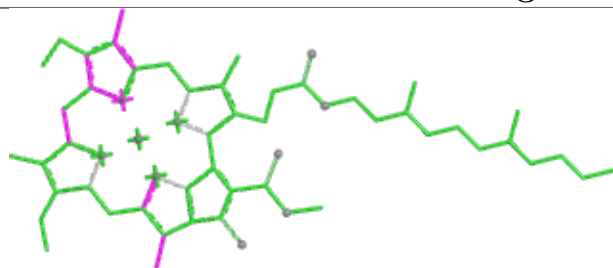
Ligand A86 D 203



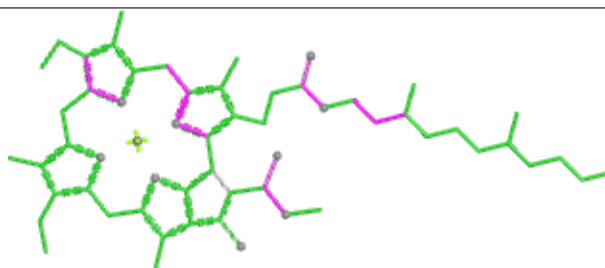
Ligand CLA a 847



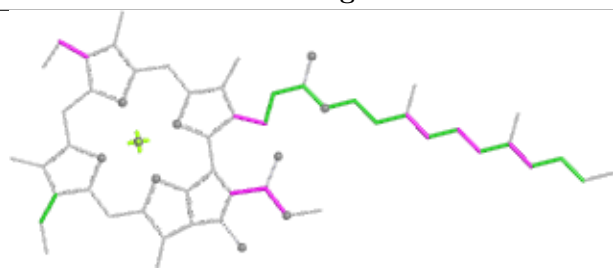


Ligand CLA b 823

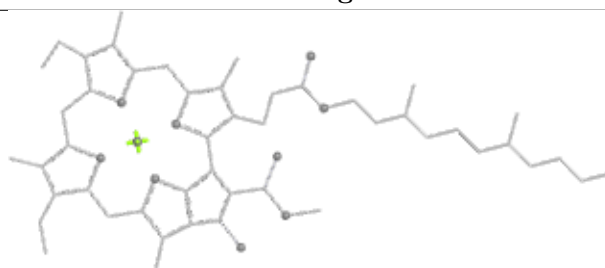
Bond lengths



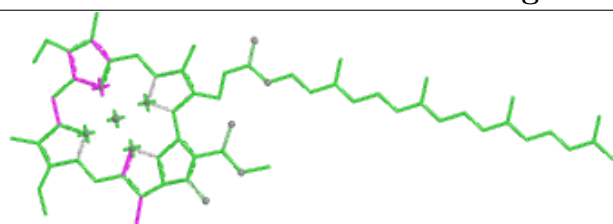
Bond angles



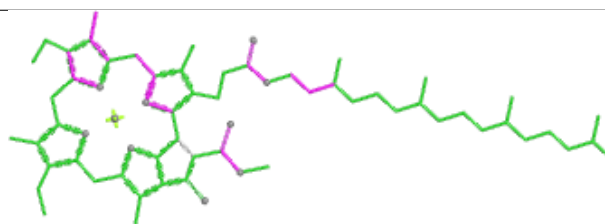
Torsions



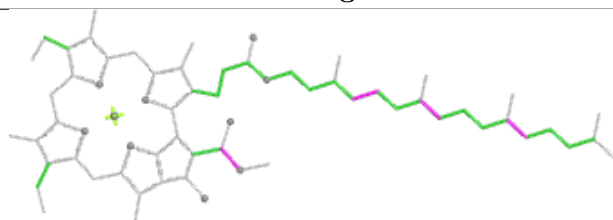
Rings

Ligand CLA f 202

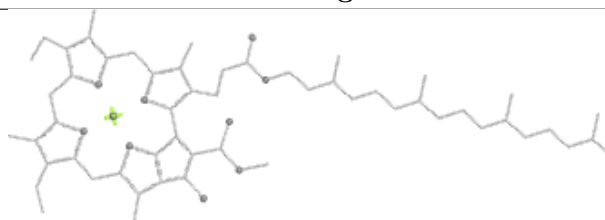
Bond lengths



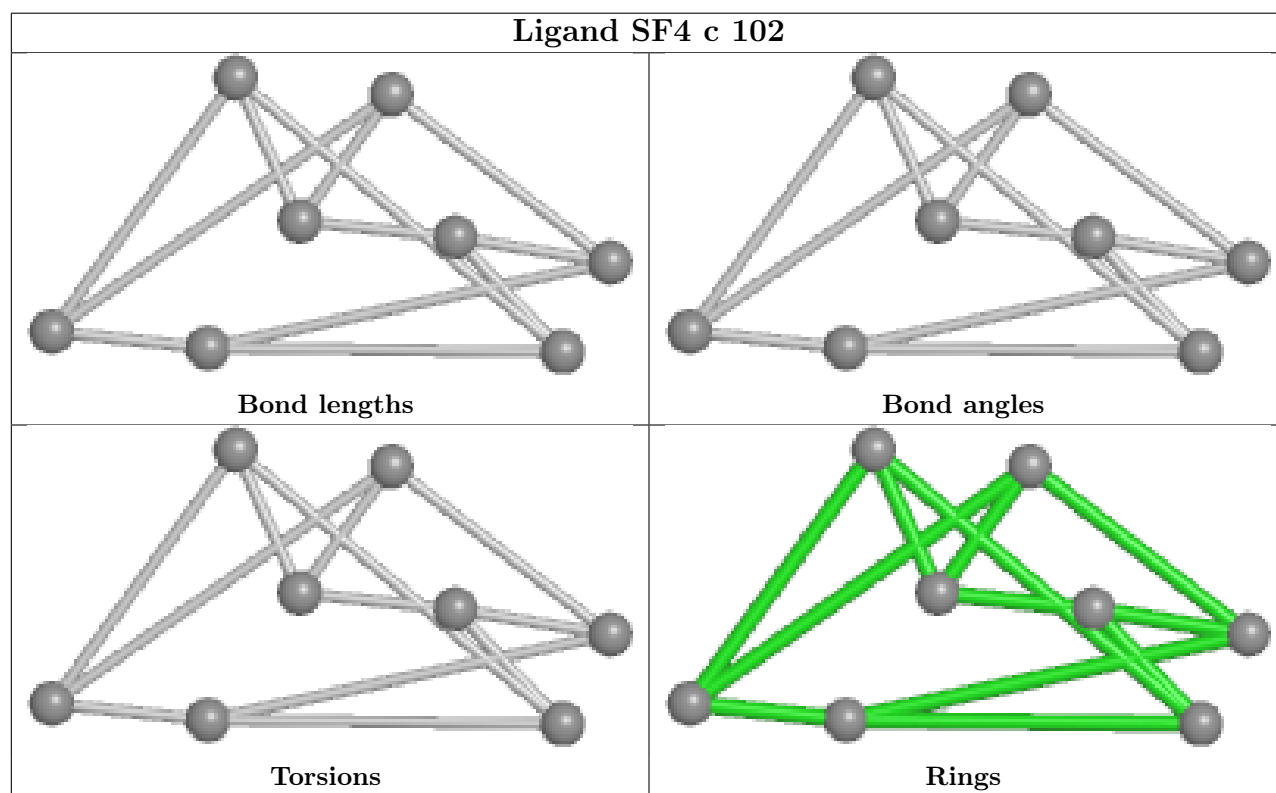
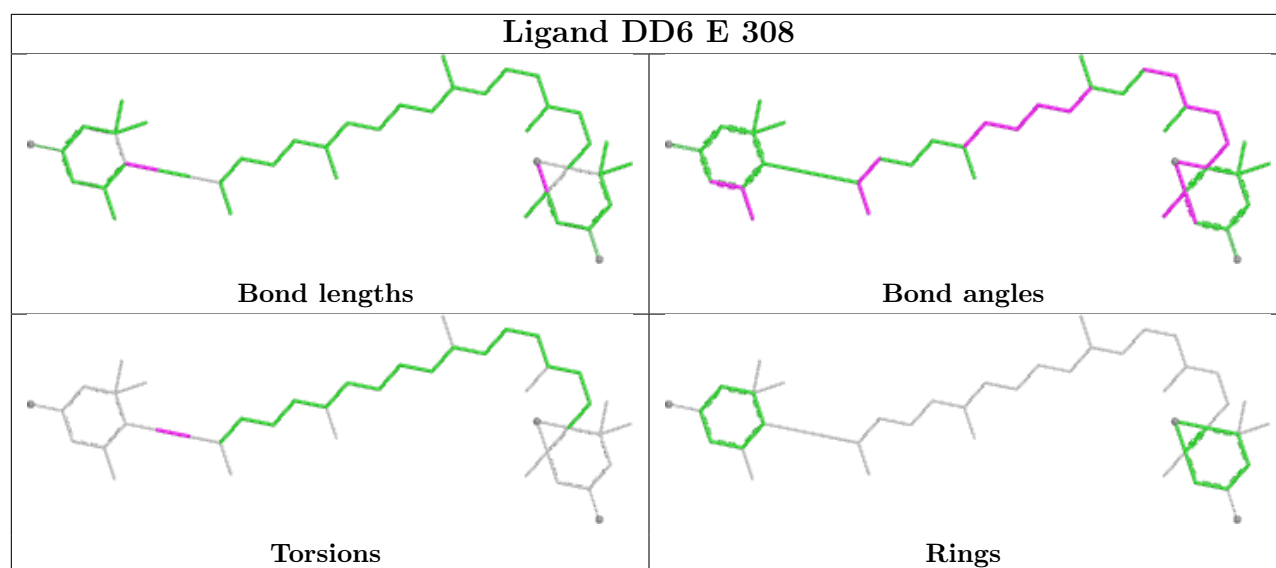
Bond angles

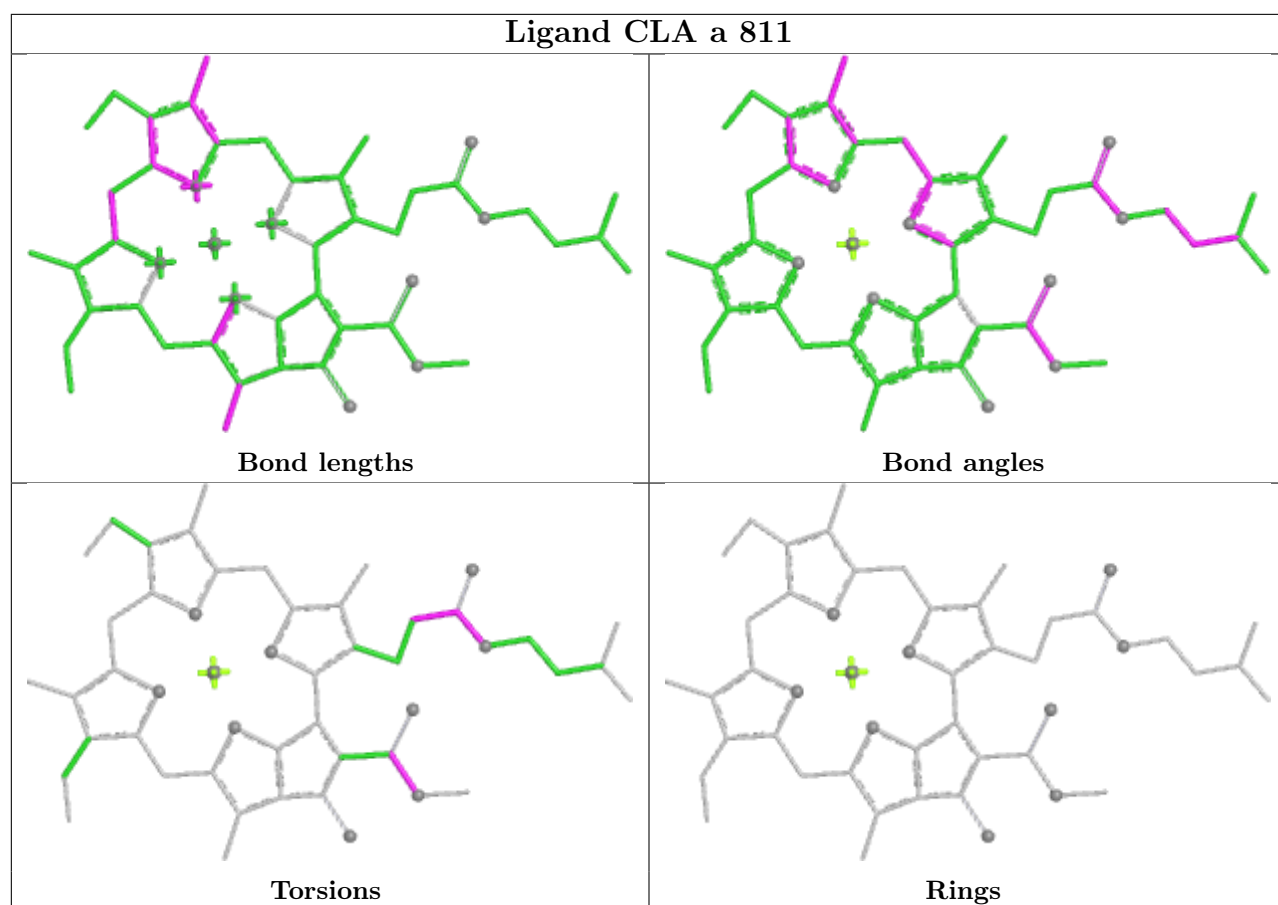


Torsions

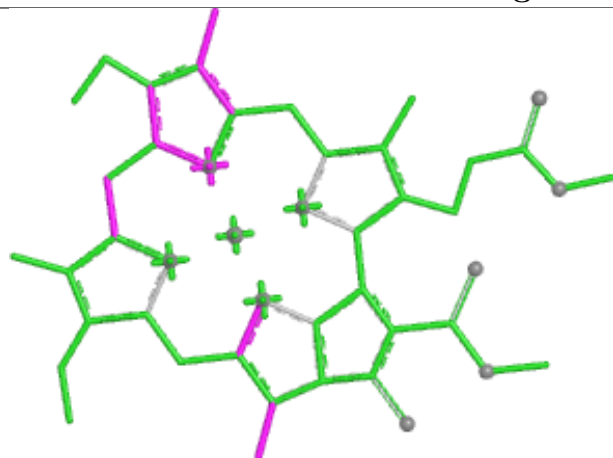


Rings

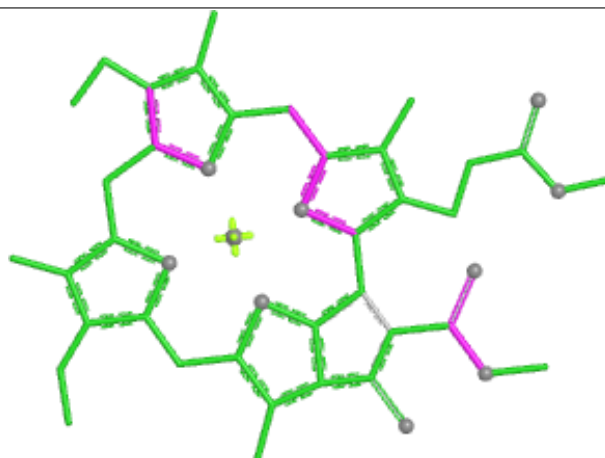




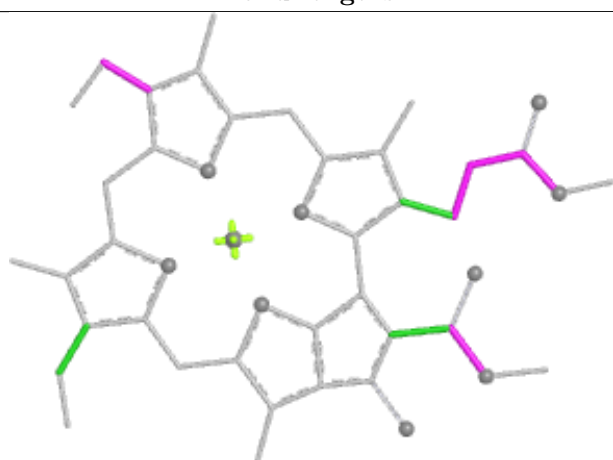
Ligand CLA H 309



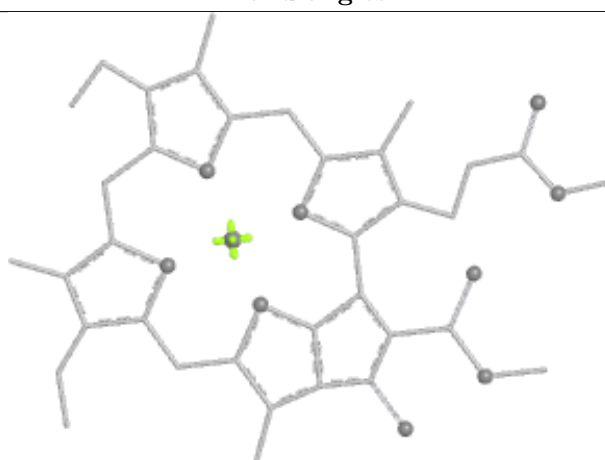
Bond lengths



Bond angles

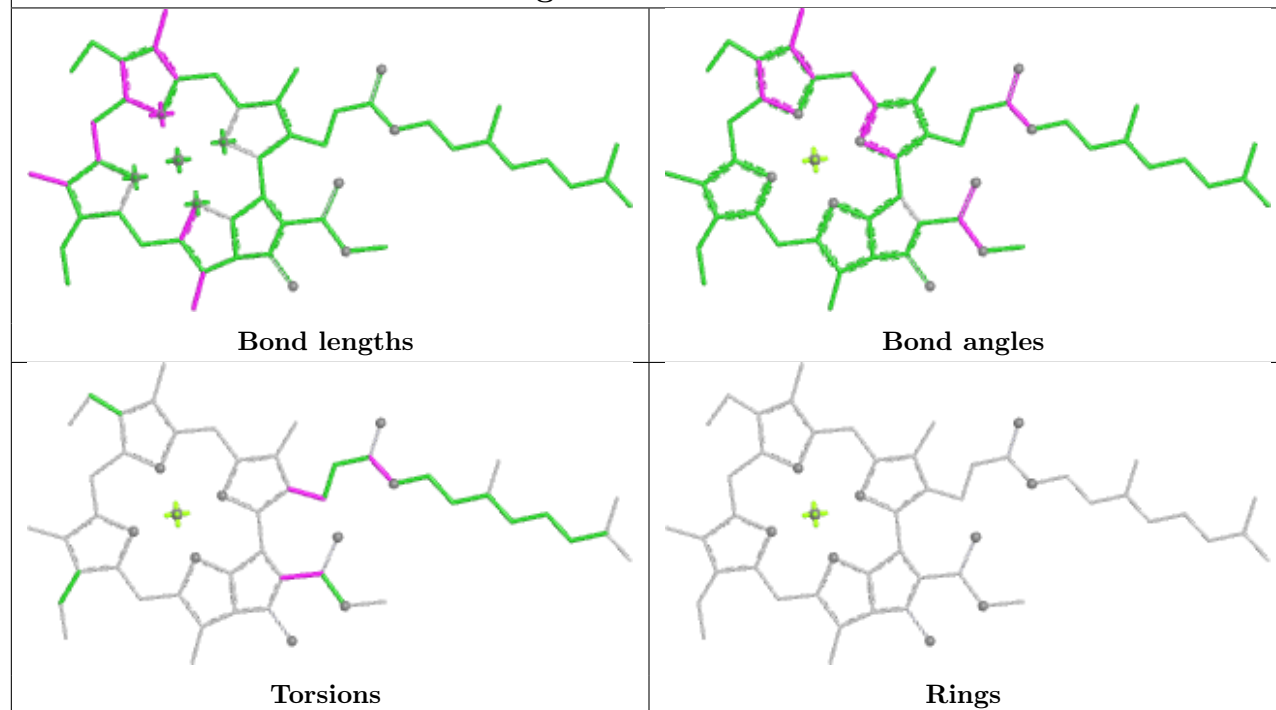


Torsions

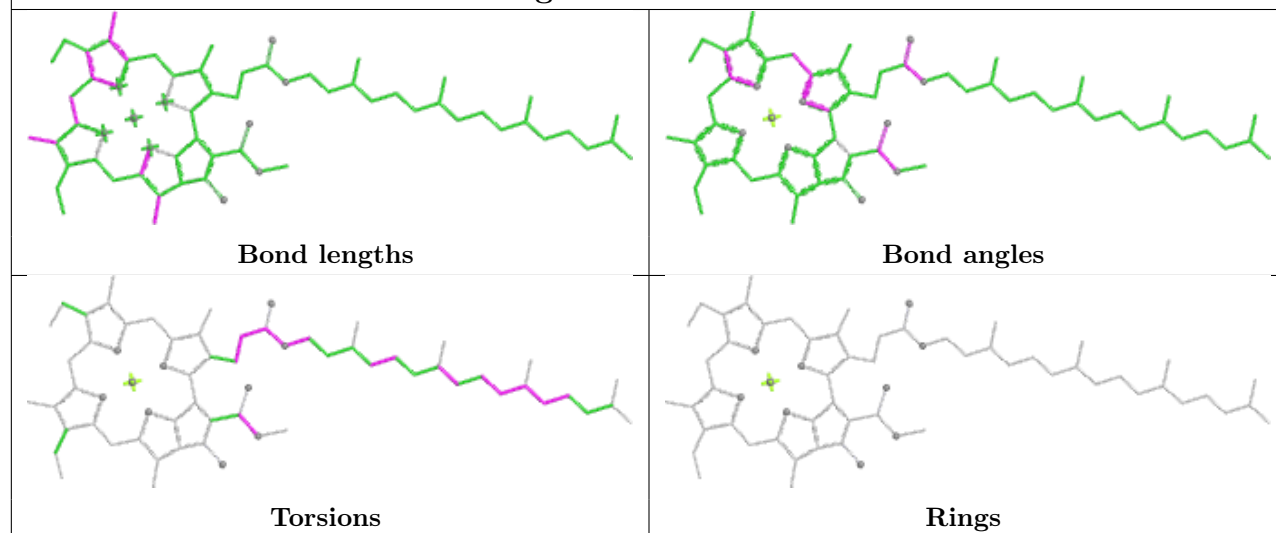


Rings

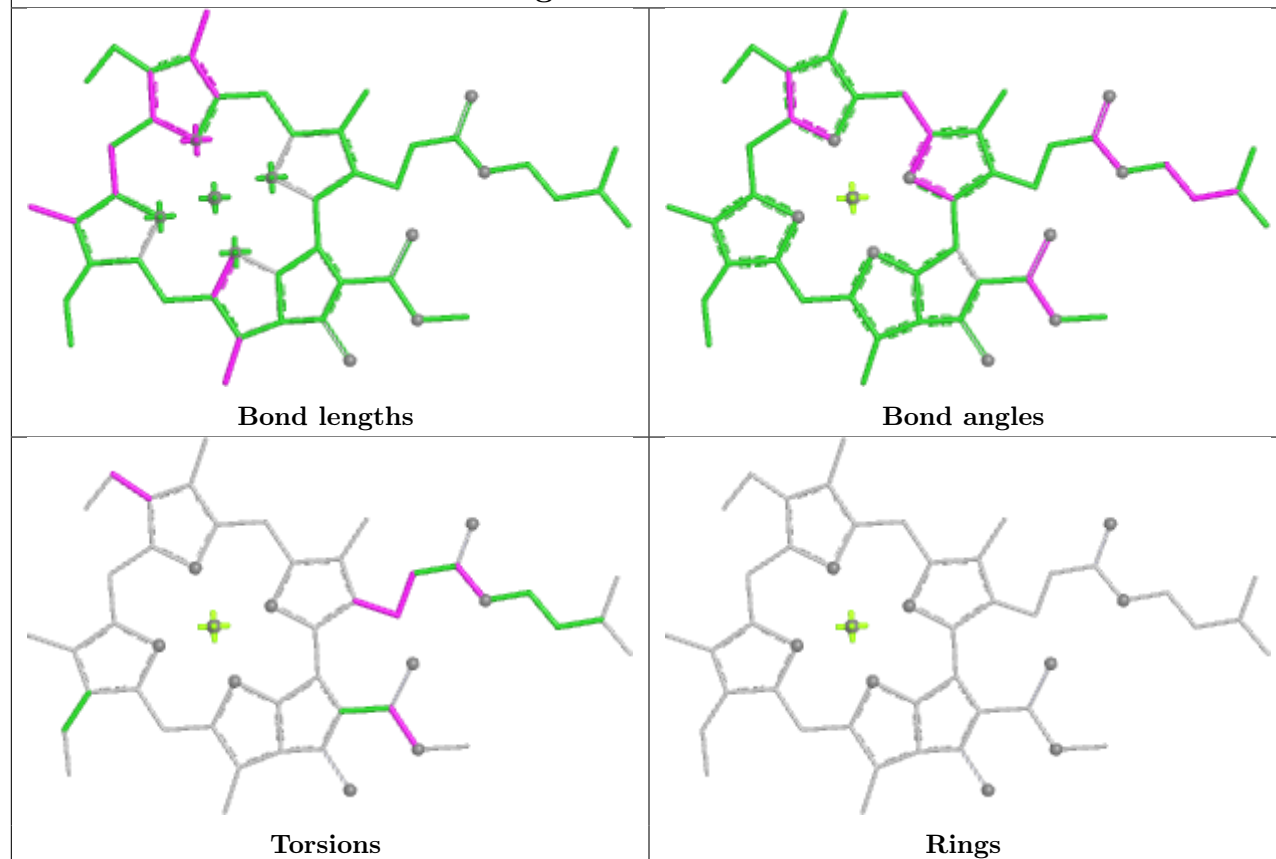
Ligand CLA a 804



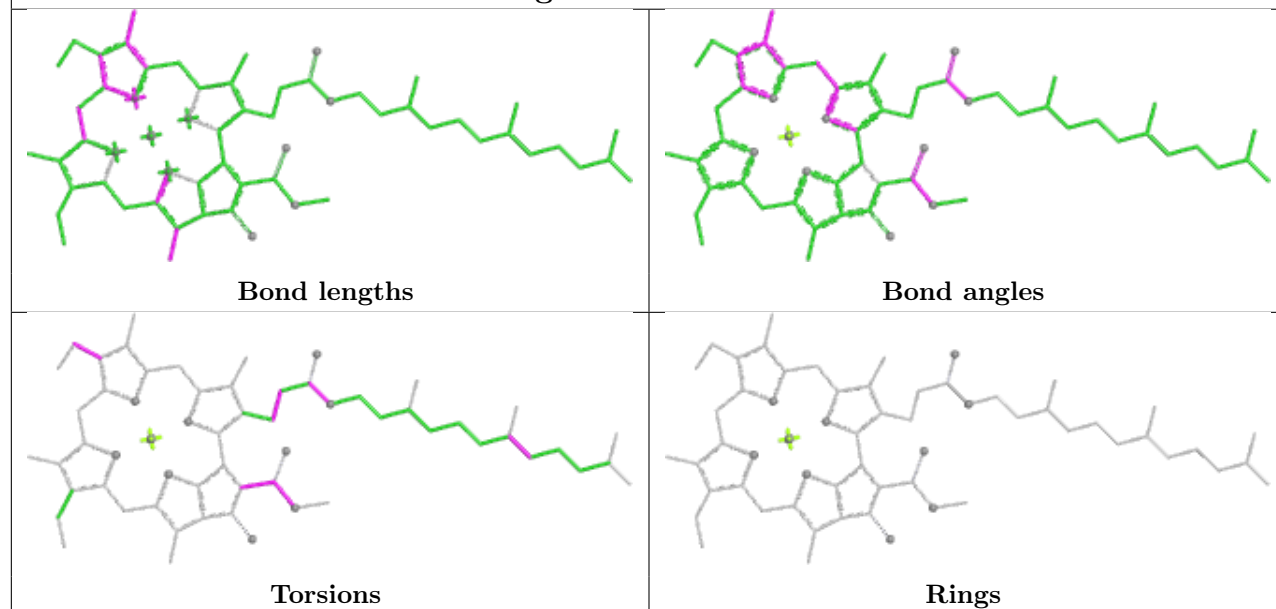
Ligand CLA b 806

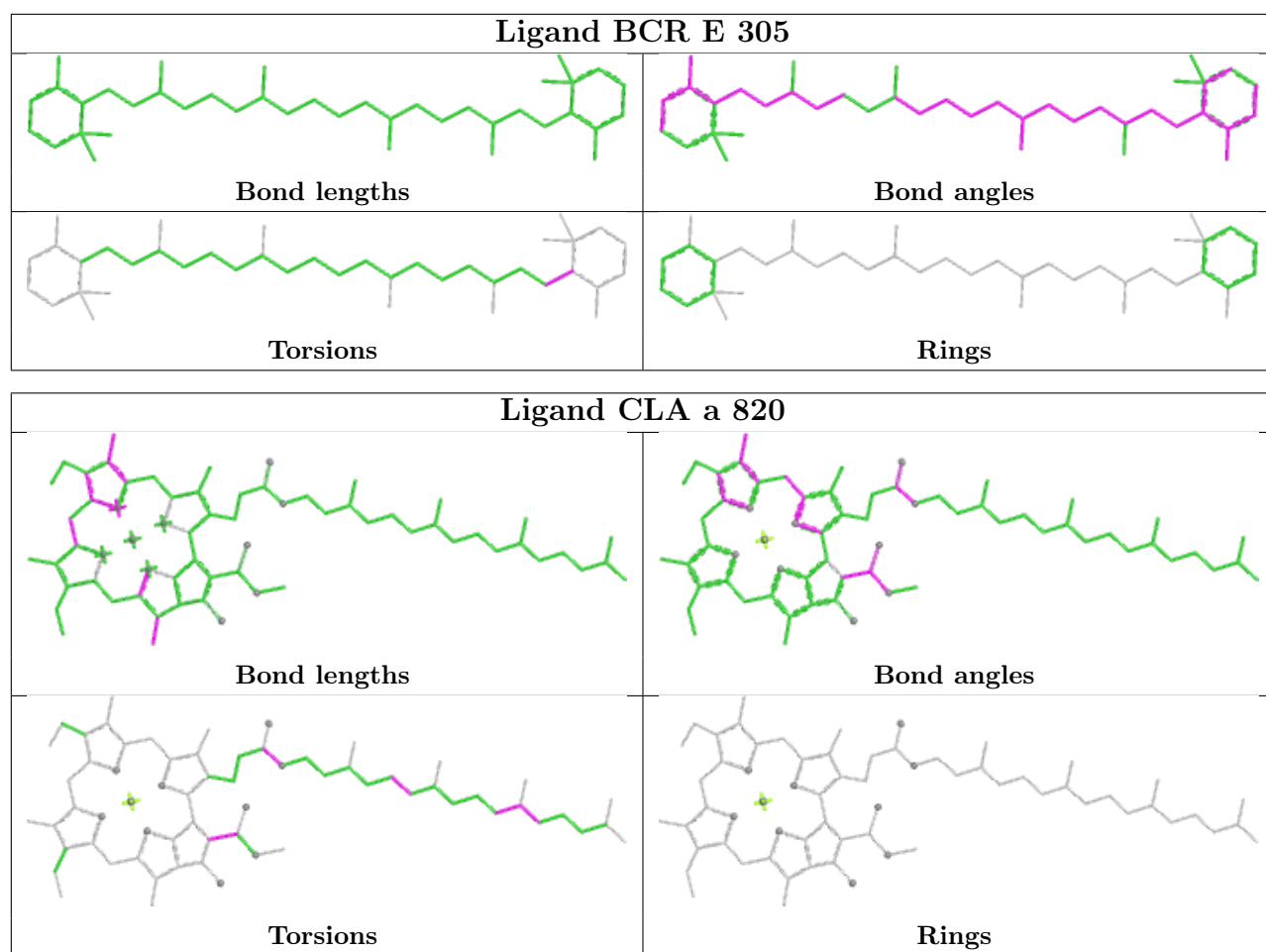


Ligand CLA b 821

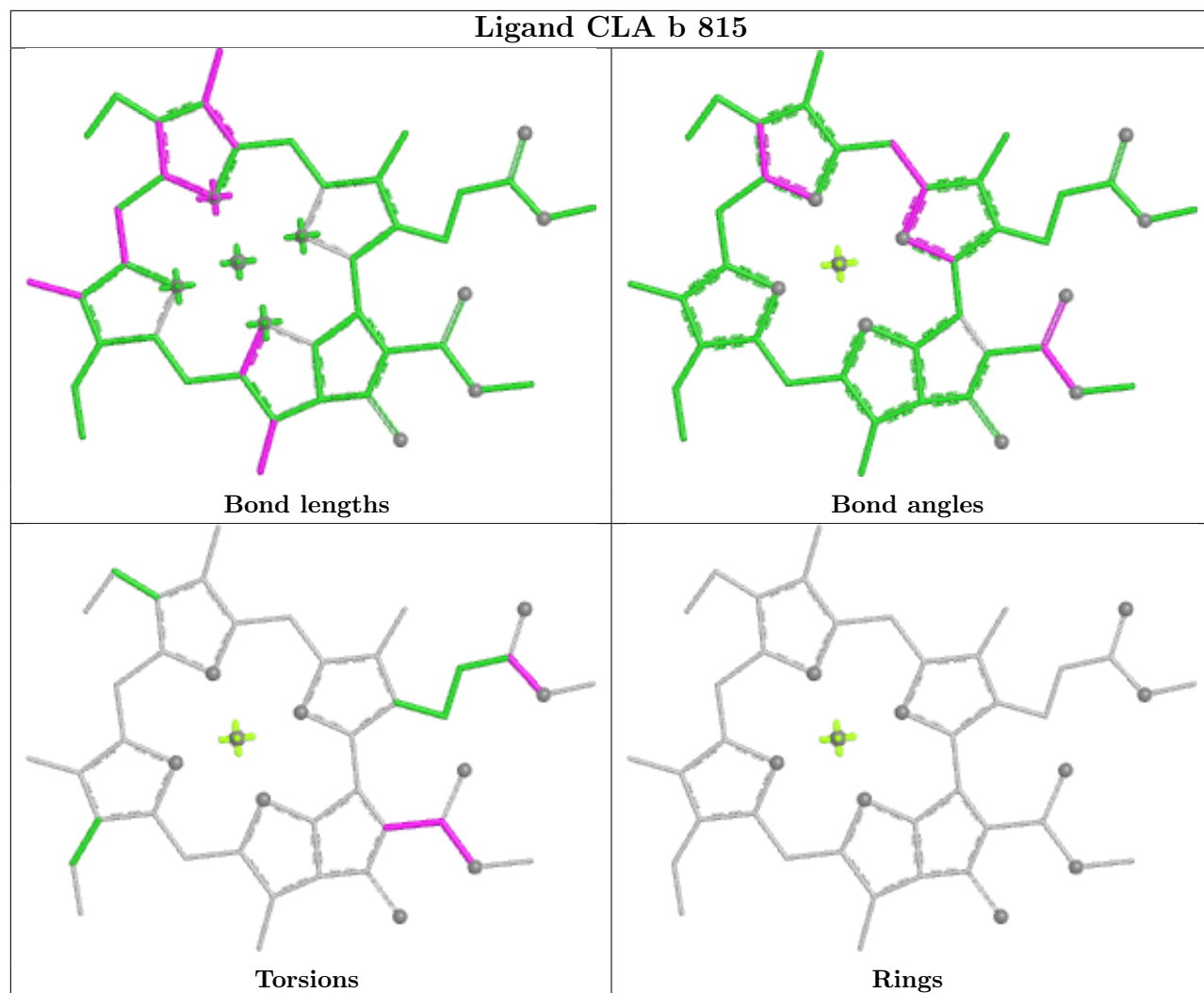


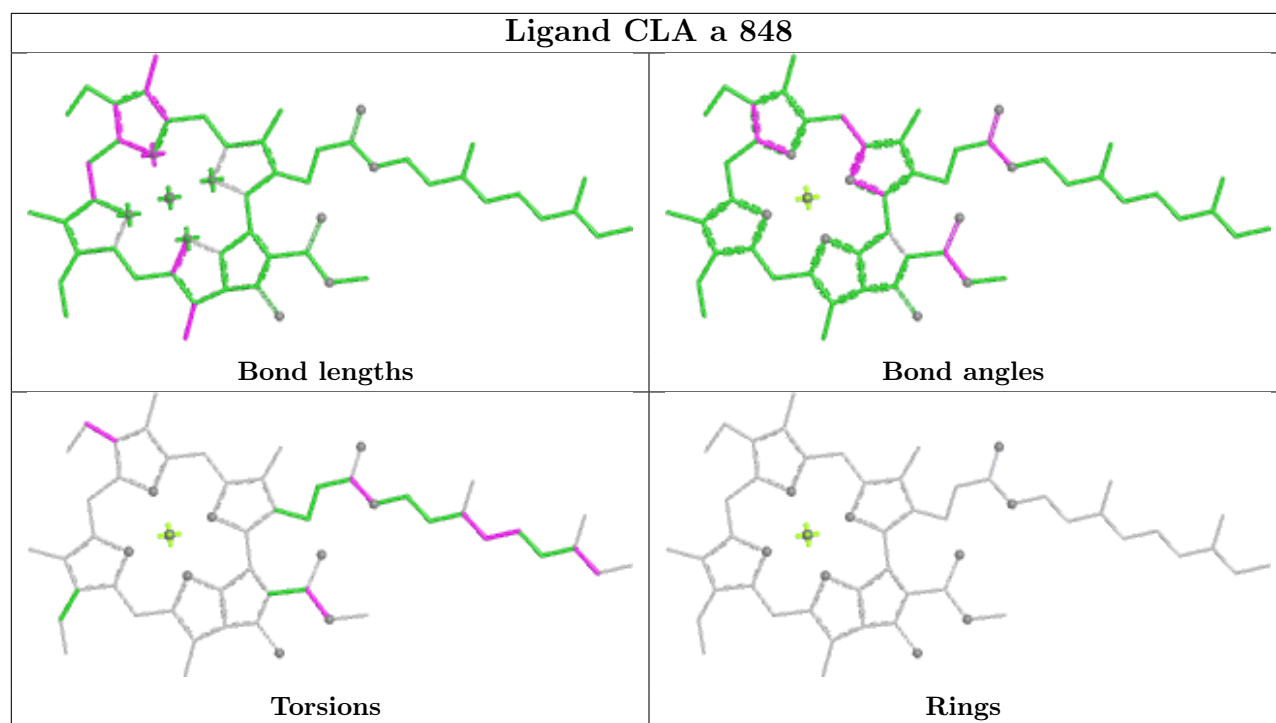
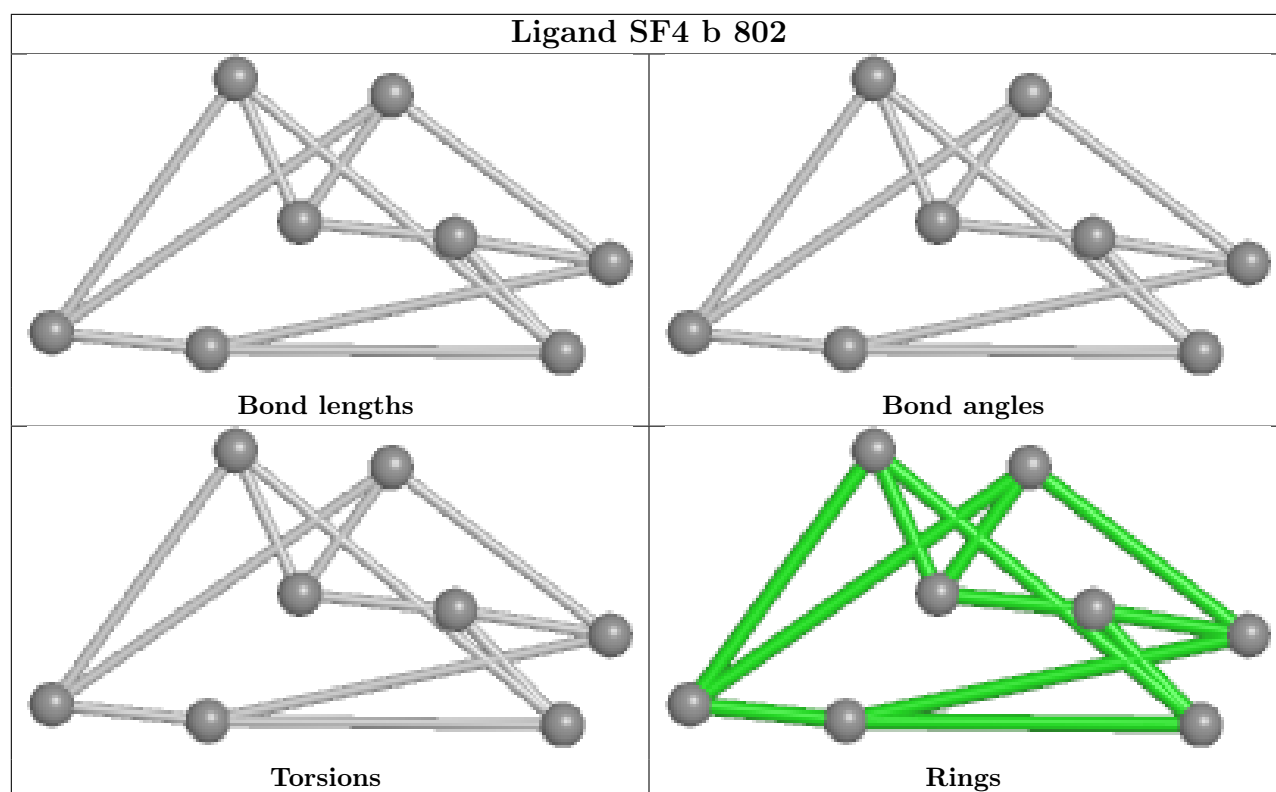
Ligand CLA b 826



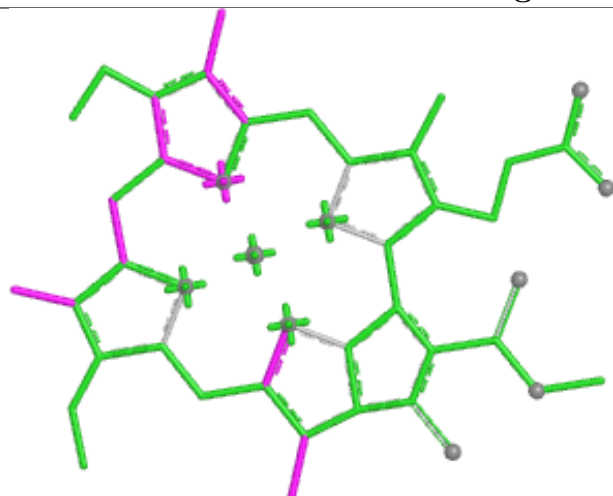


Ligand CLA b 815

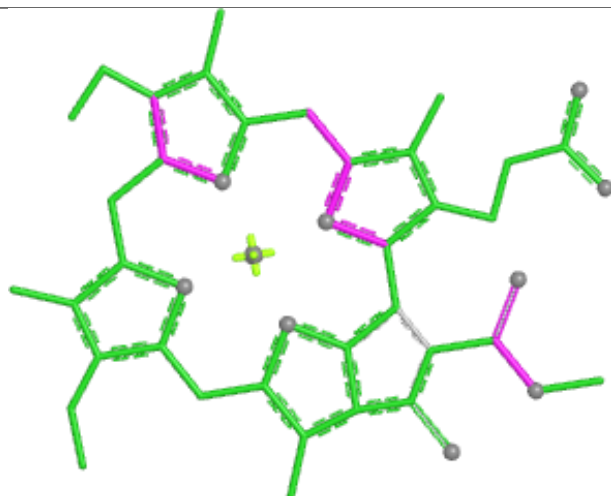




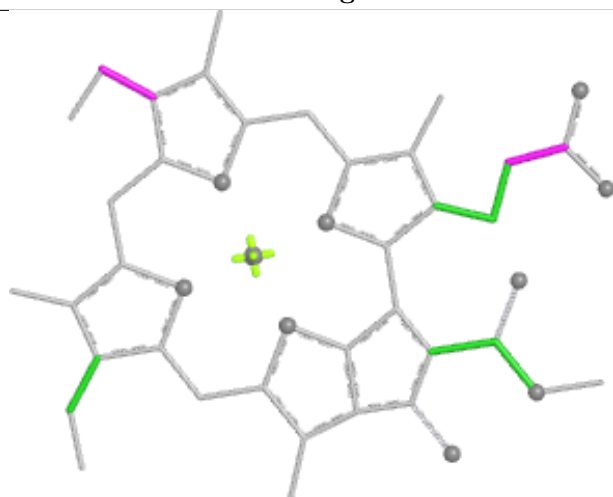
Ligand CLA C 306



Bond lengths



Bond angles

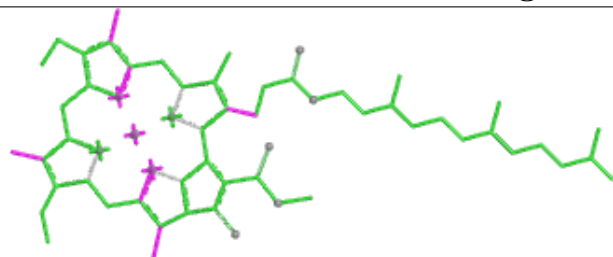


Torsions

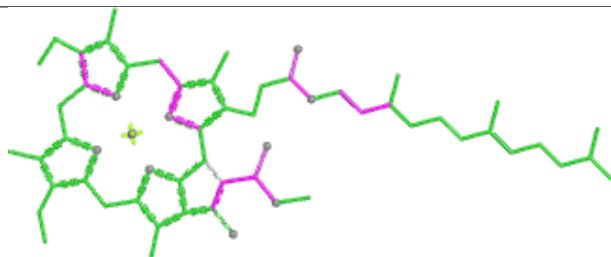


Rings

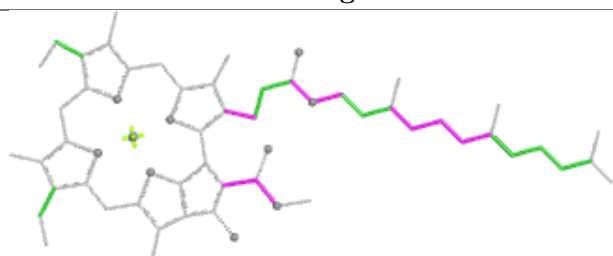
Ligand CLA a 837



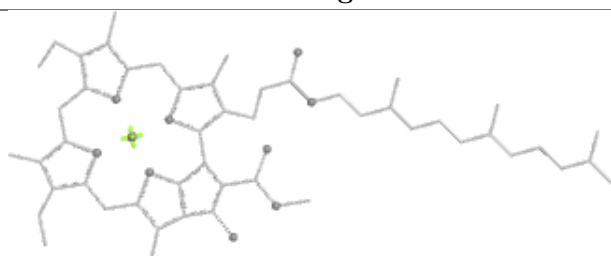
Bond lengths



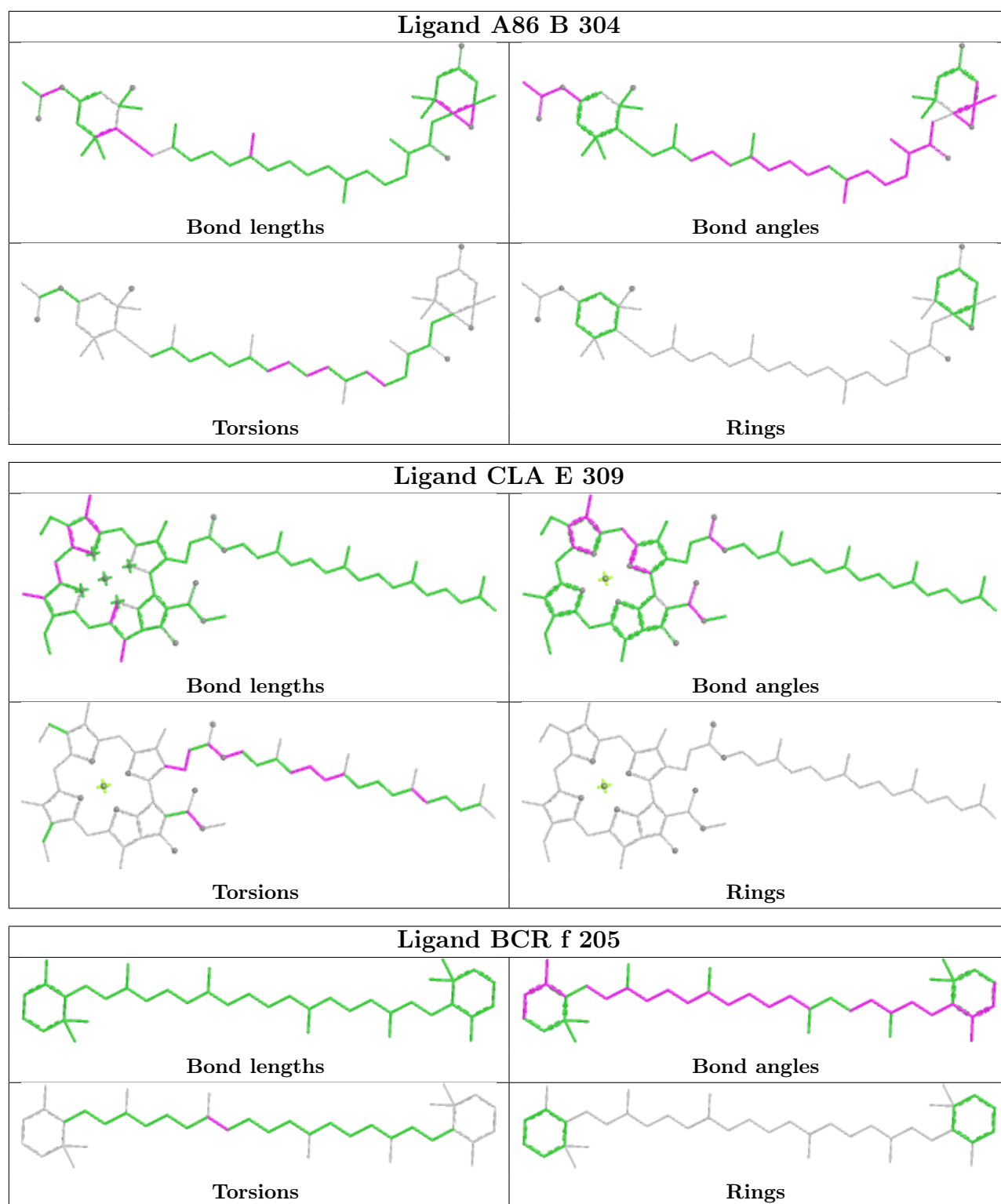
Bond angles

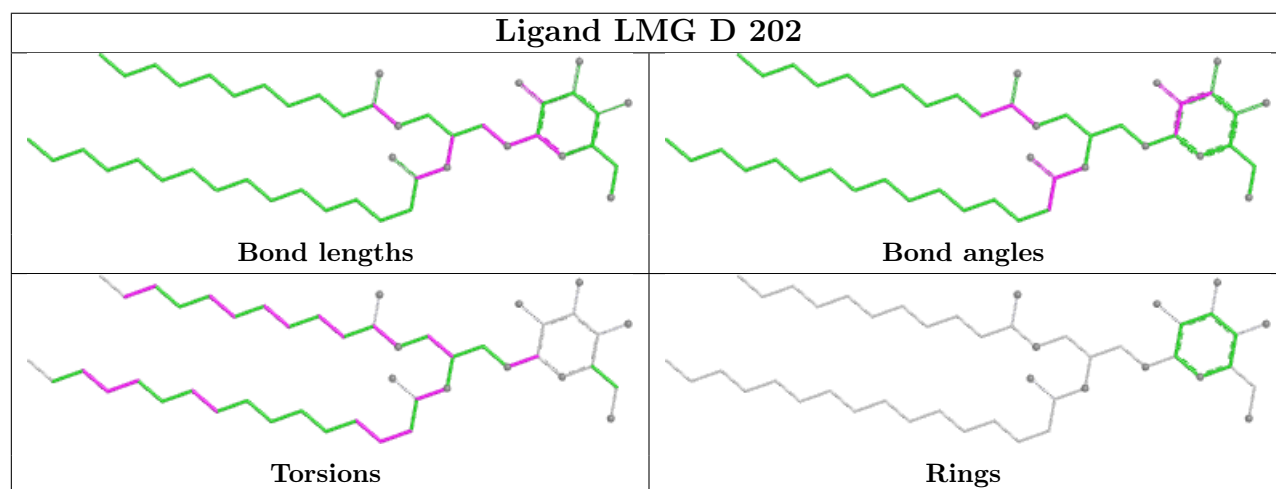
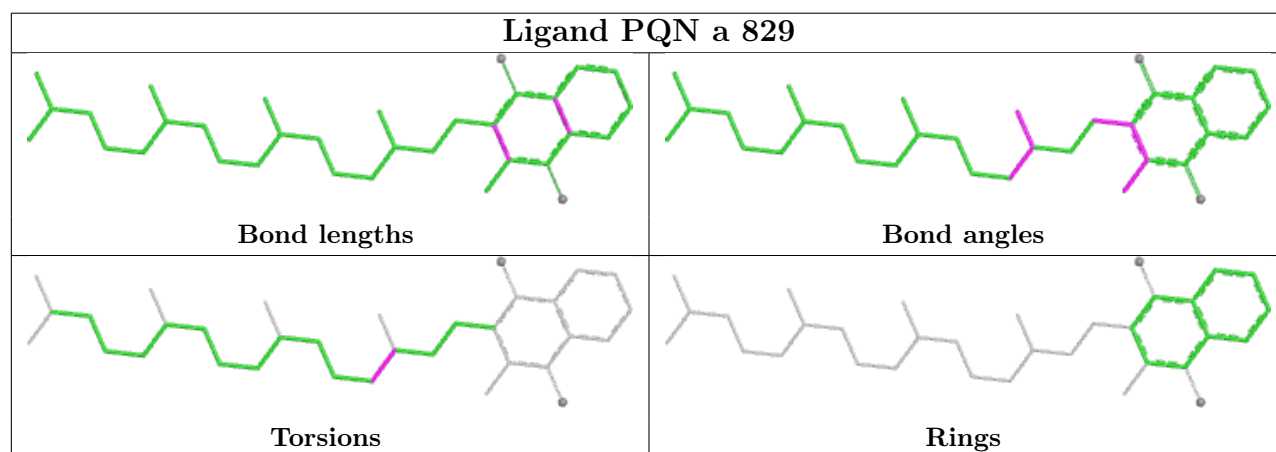
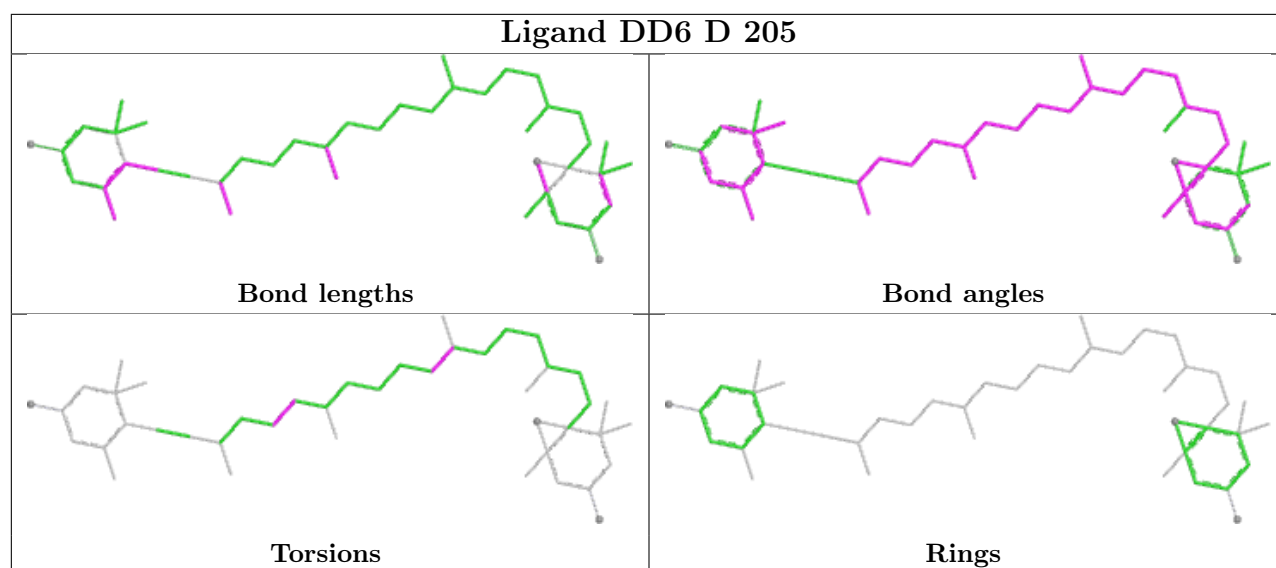


Torsions

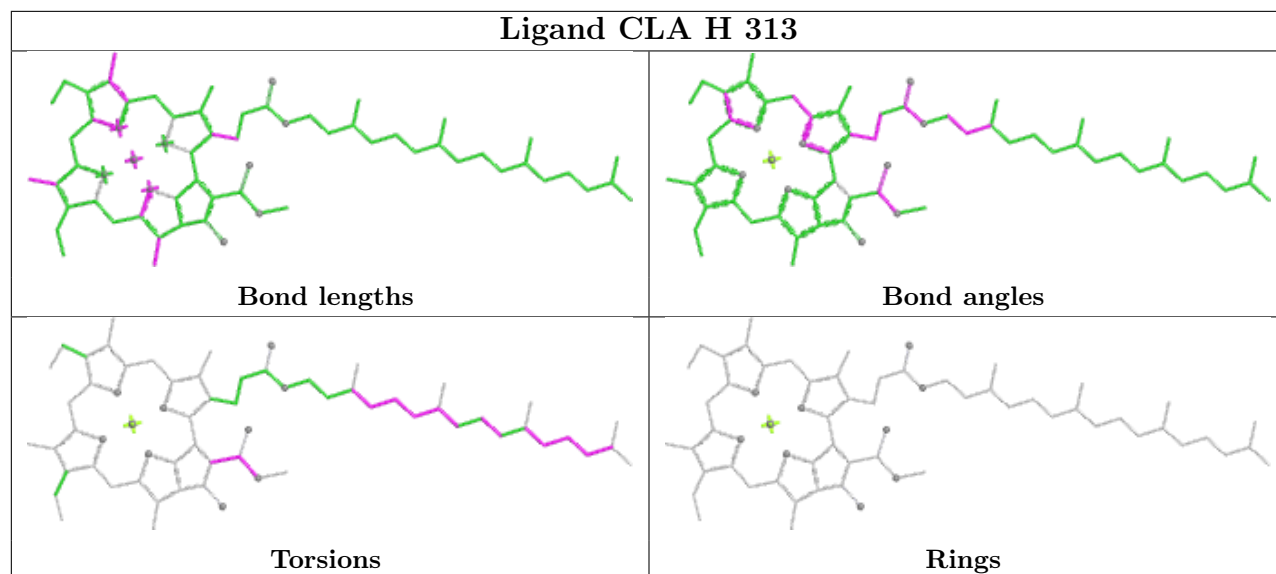


Rings

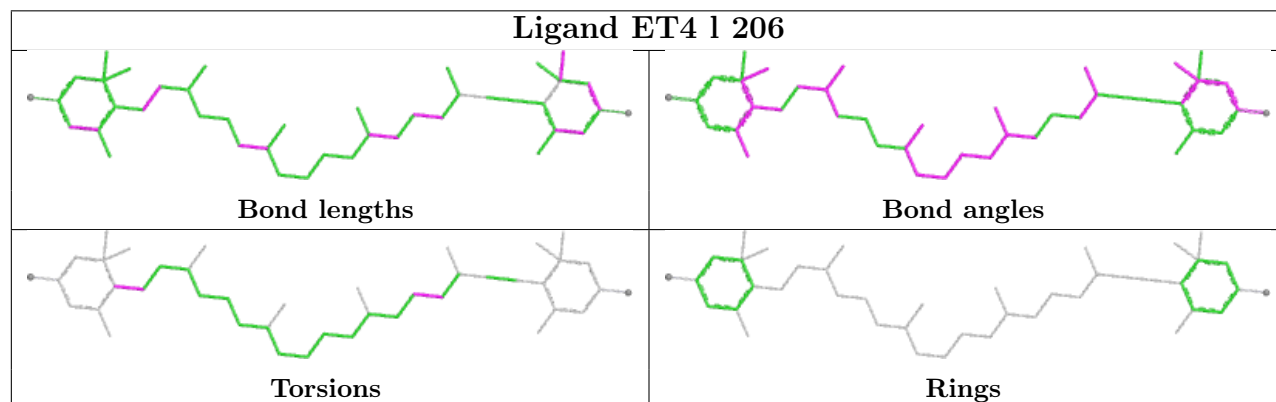




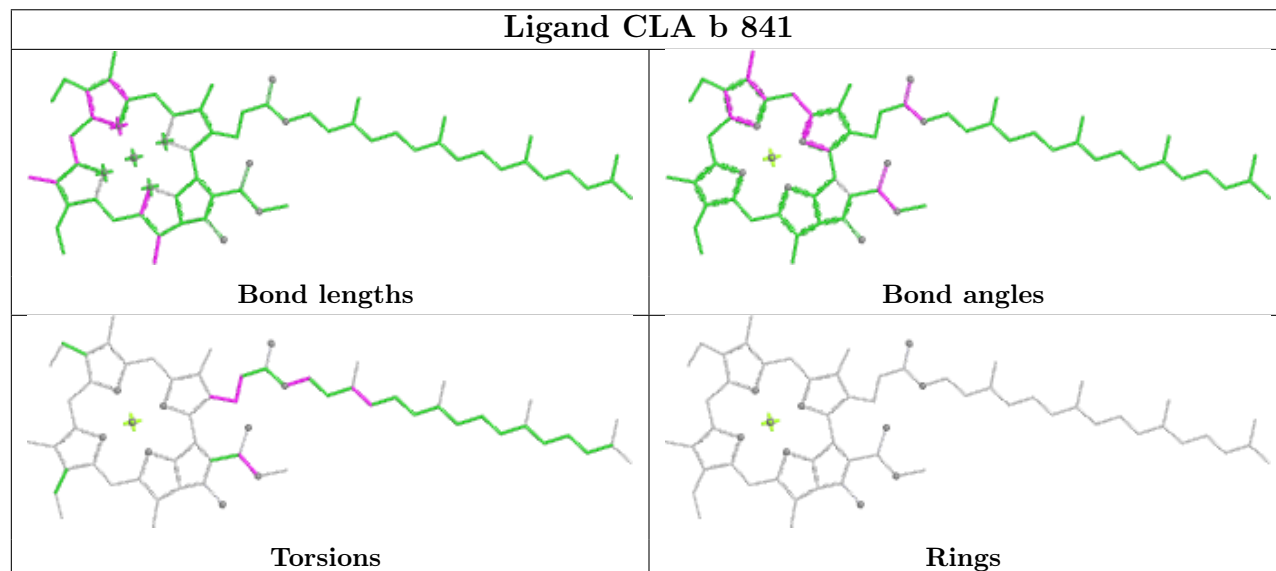
Ligand CLA H 313

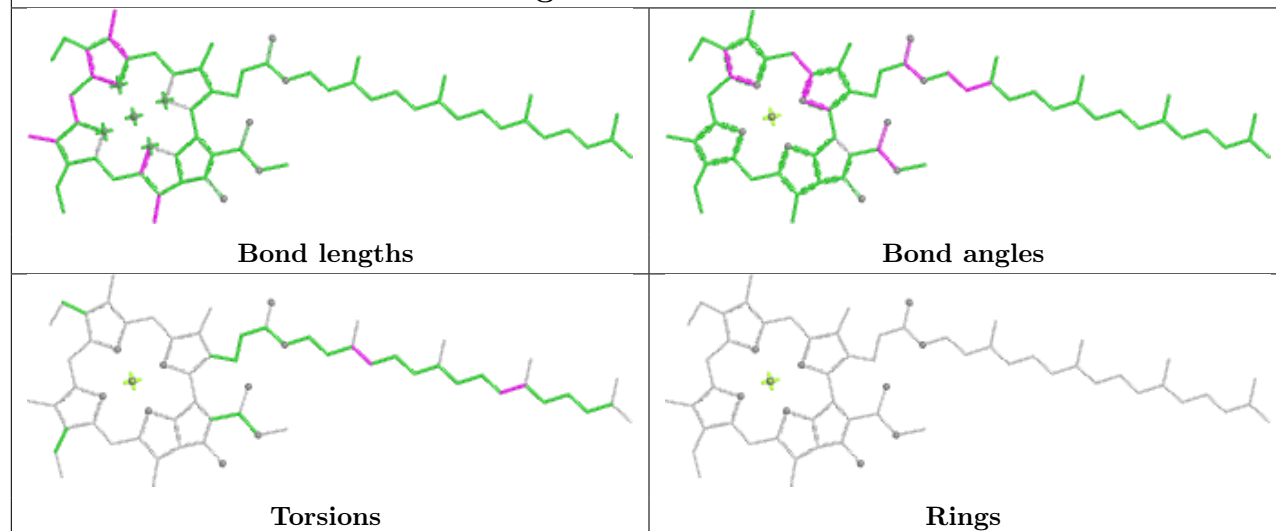
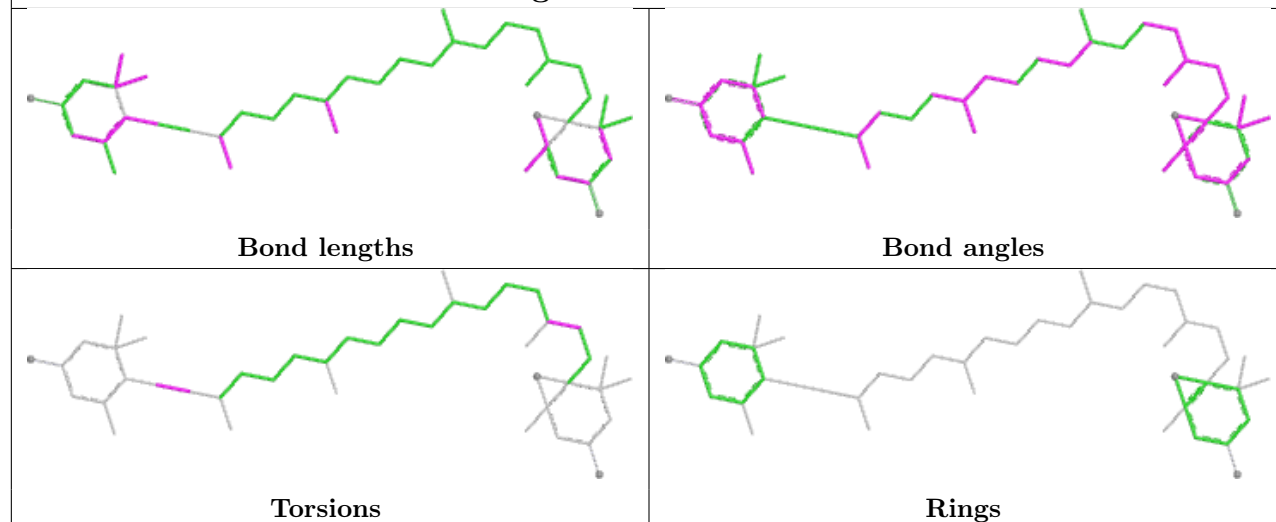


Ligand ET4 l 206

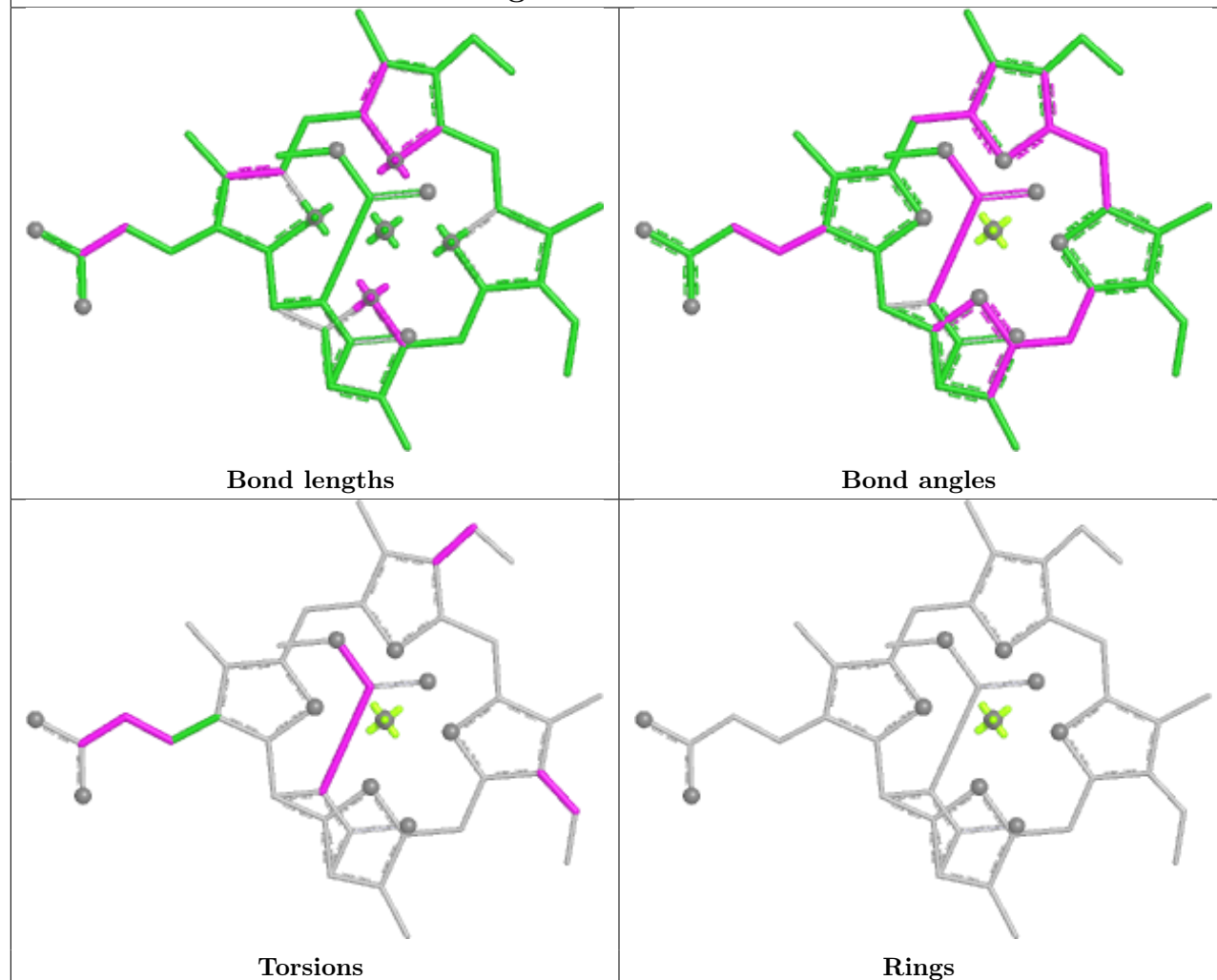


Ligand CLA b 841

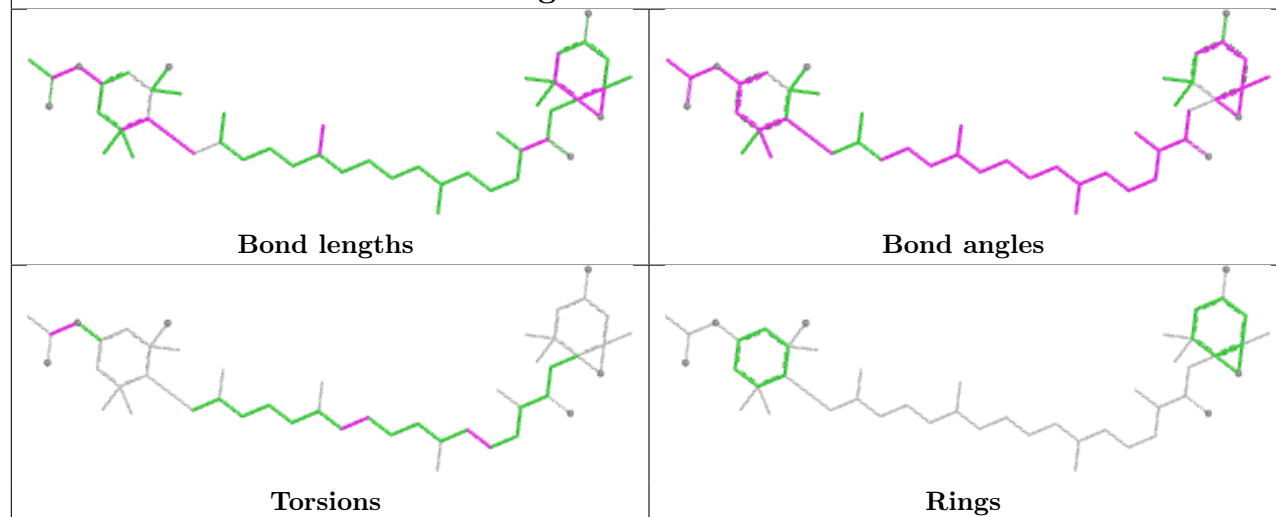


Ligand CLA a 819**Ligand DD6 H 303**

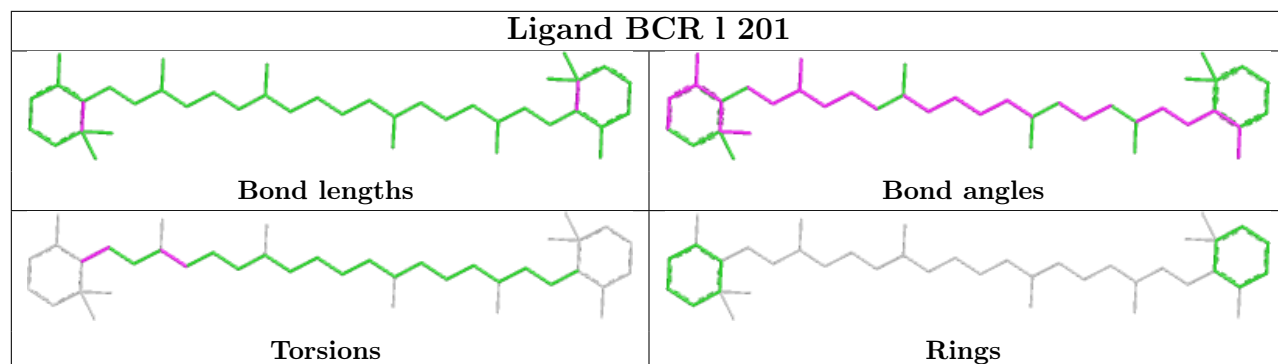
Ligand KC1 H 312



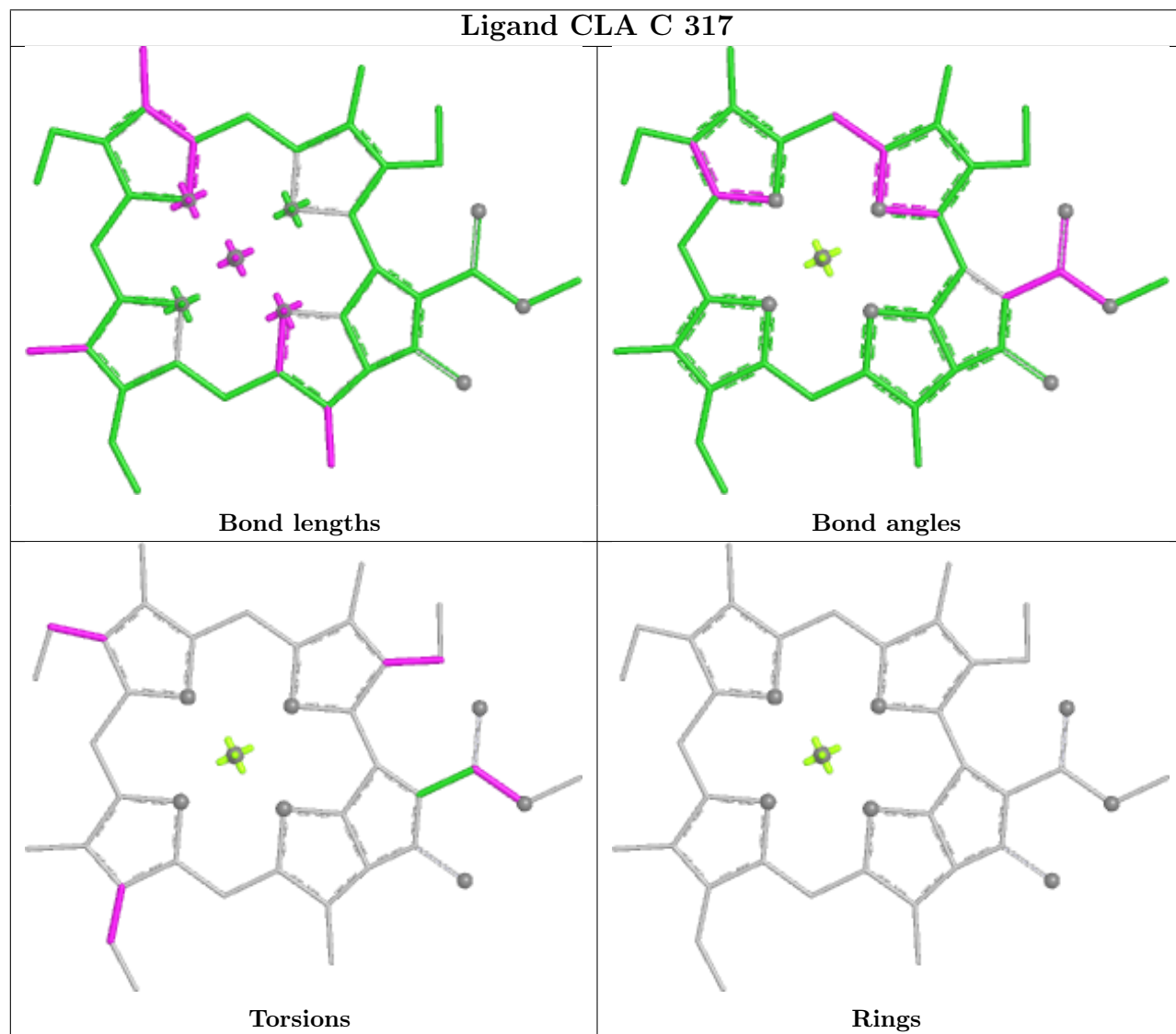
Ligand A86 B 305



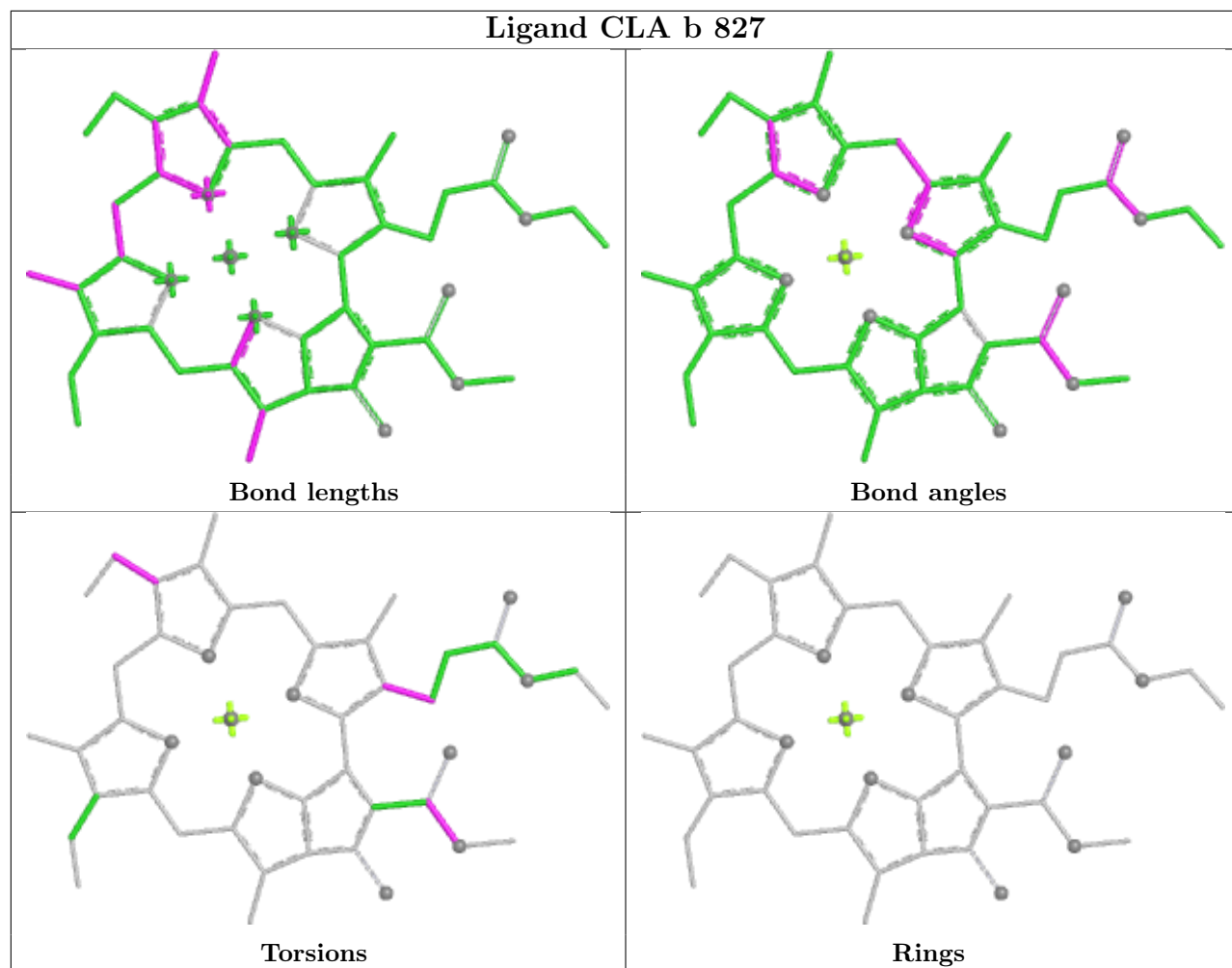
Ligand BCR 1 201



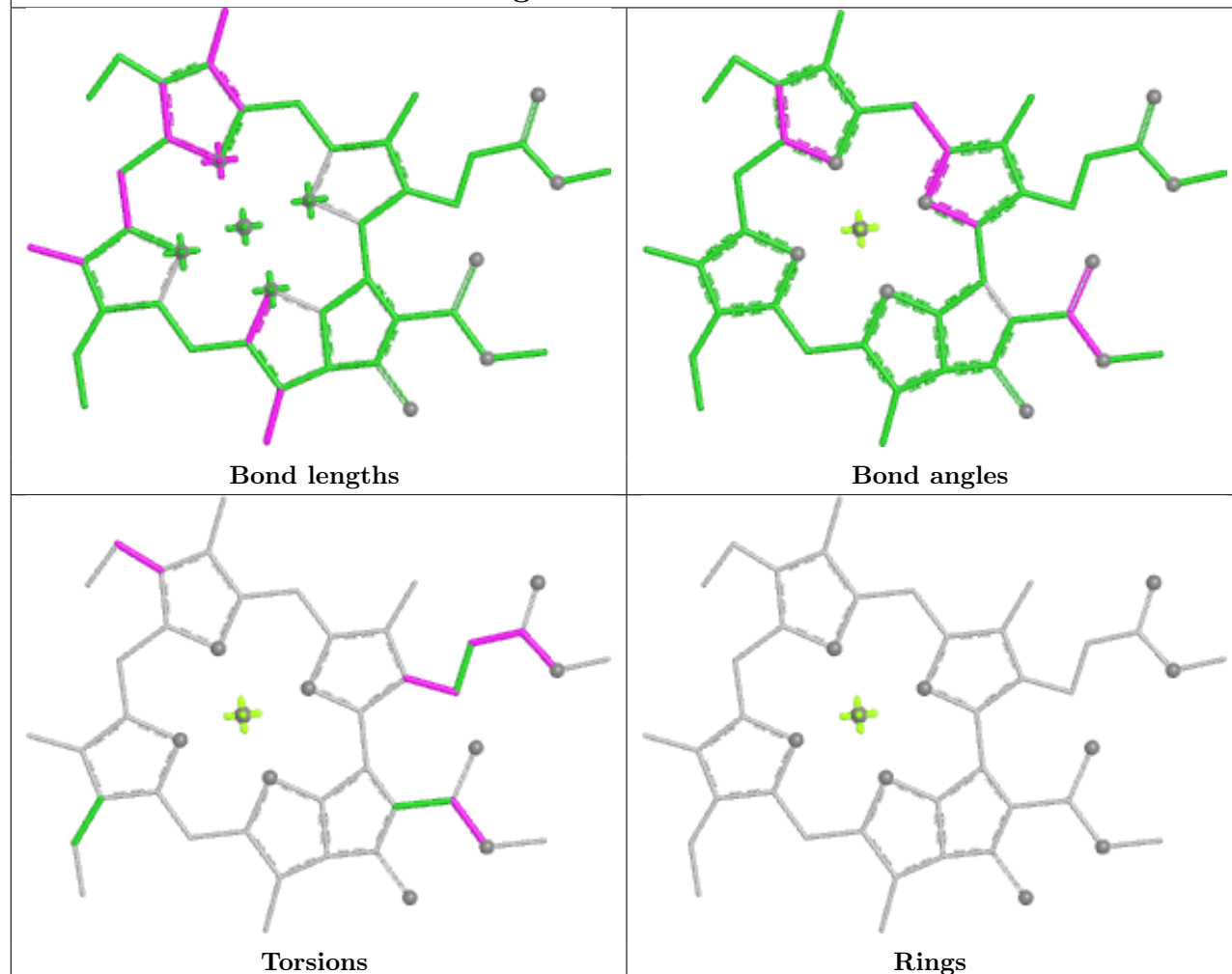
Ligand CLA C 317



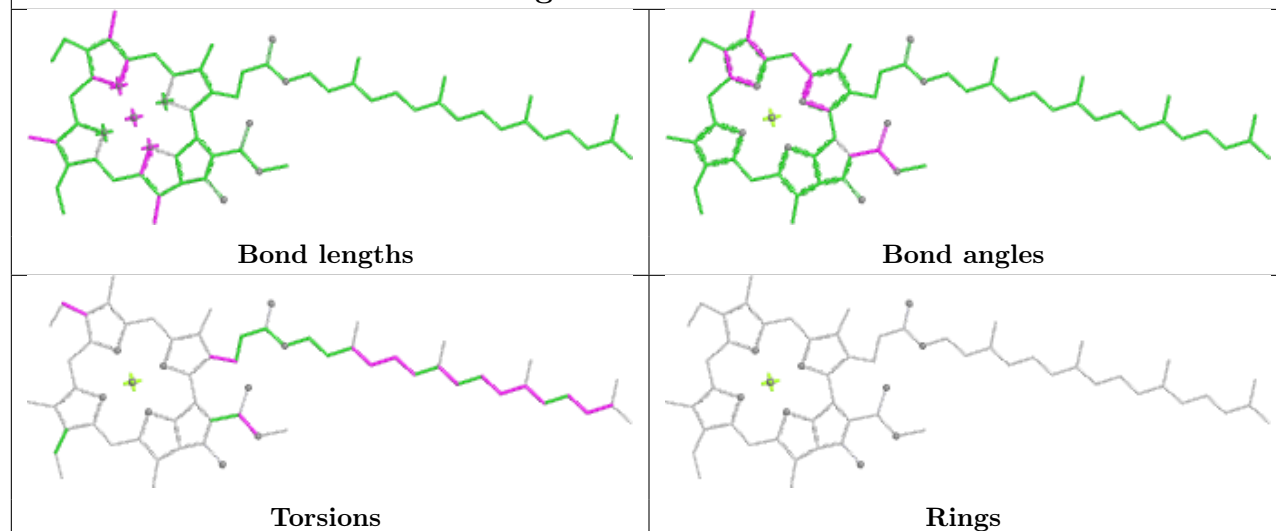
Ligand CLA b 827

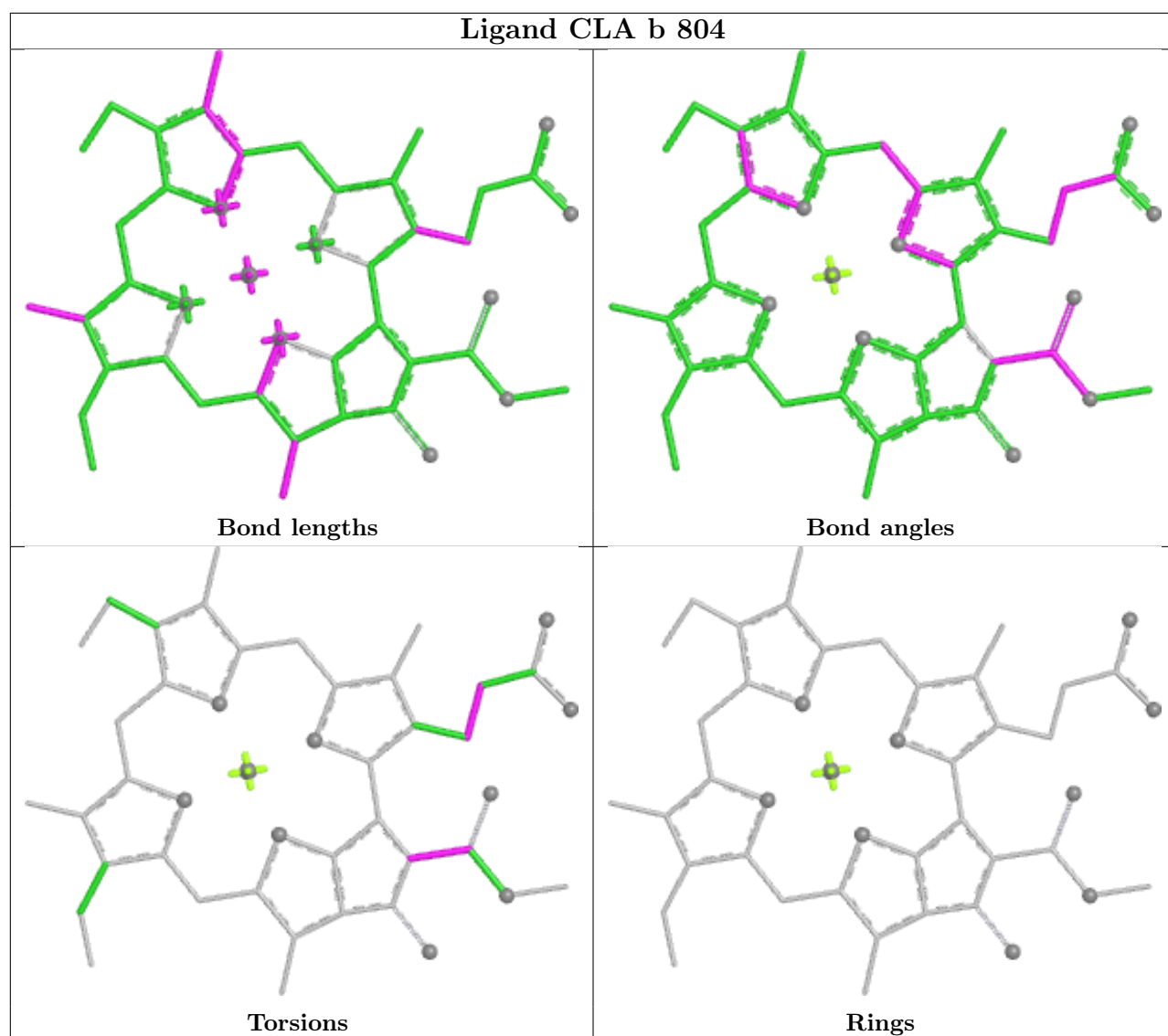


Ligand CLA D 211

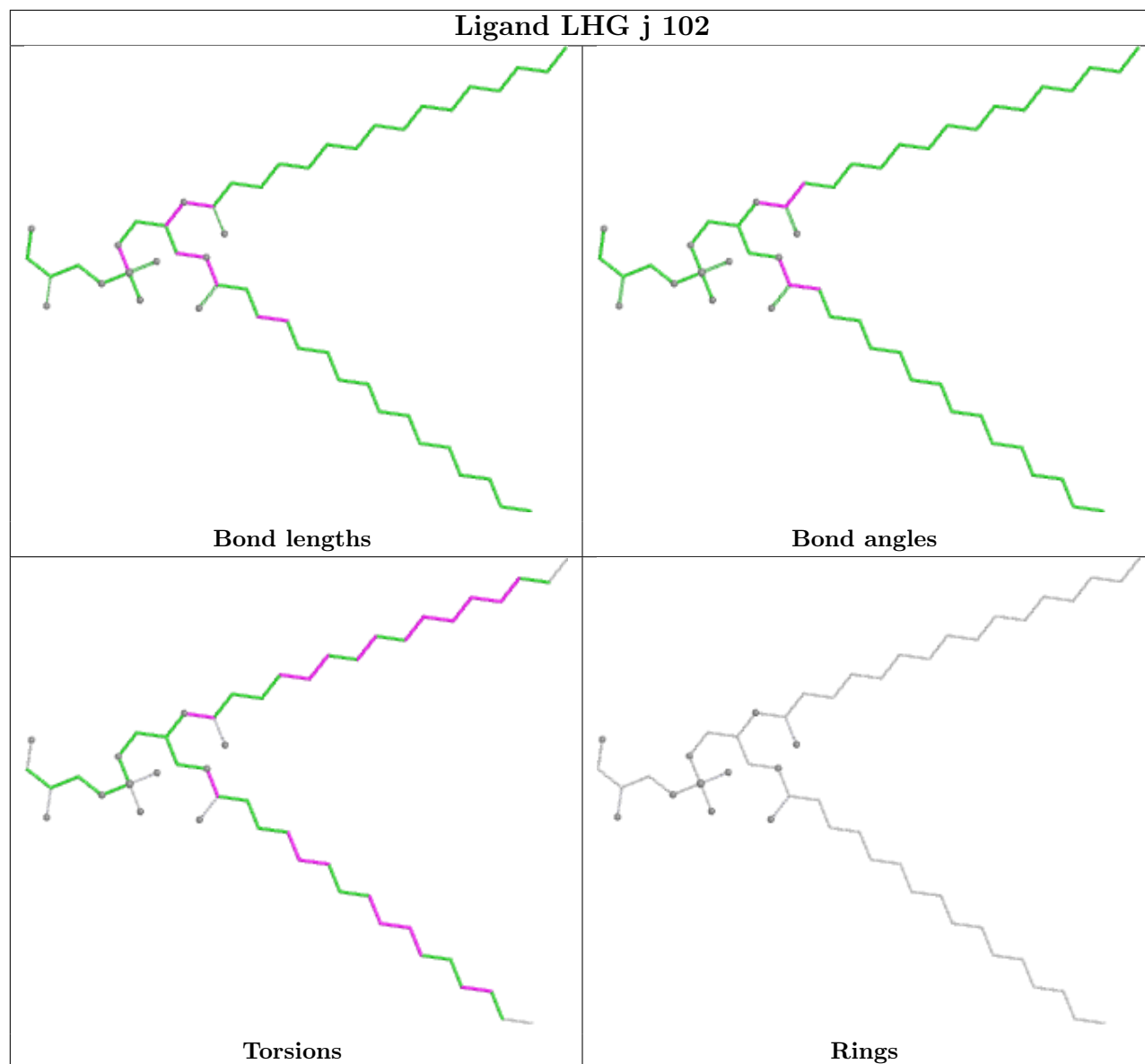


Ligand CLA b 843

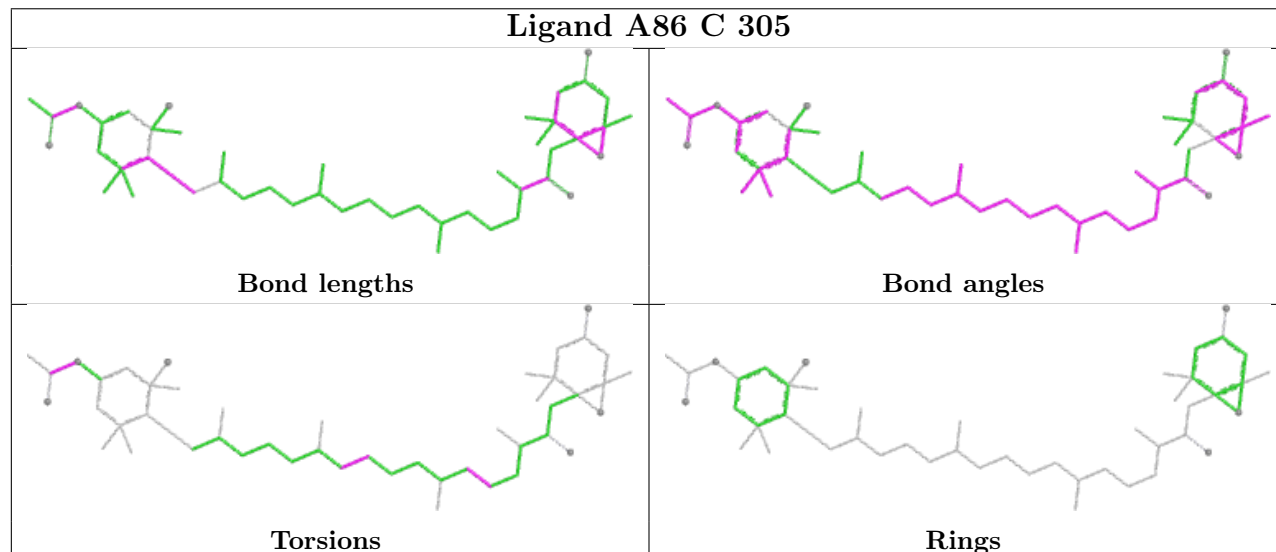


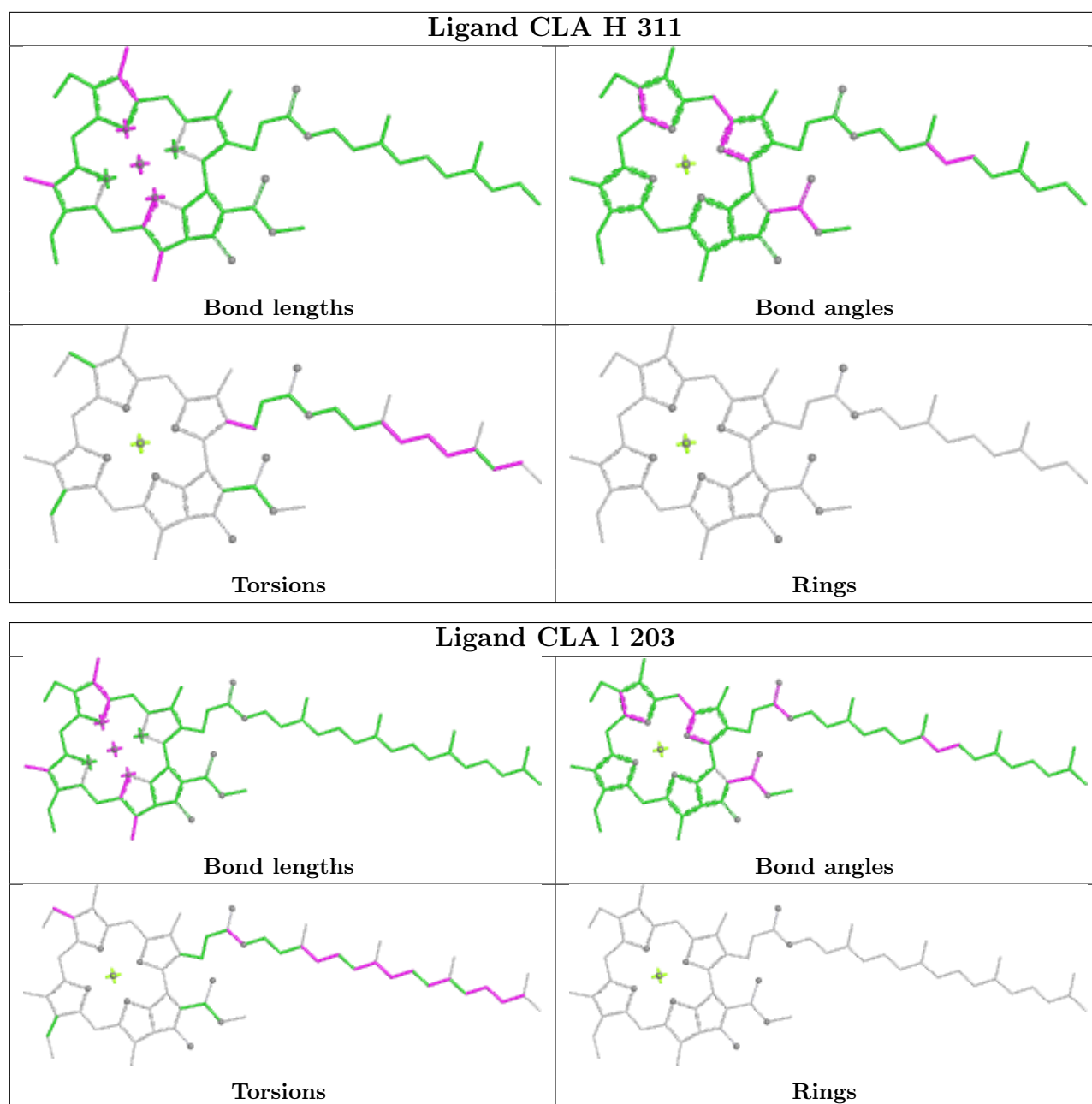


Ligand LHG j 102

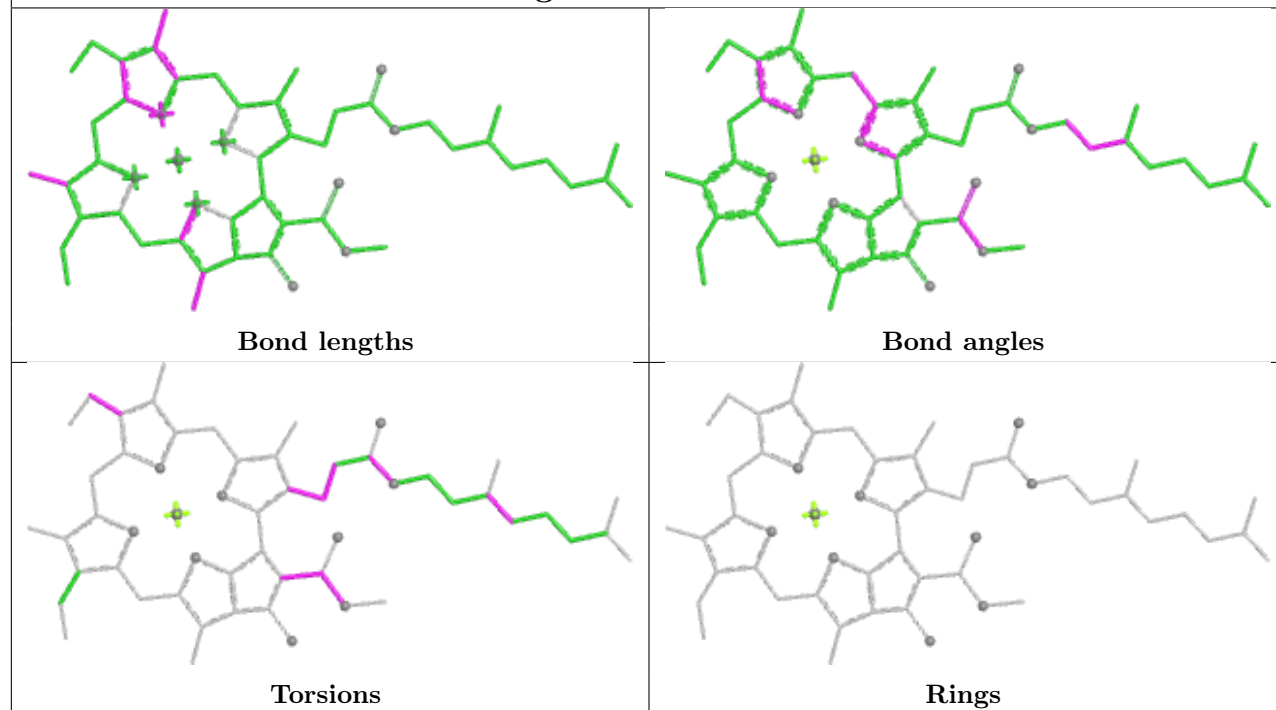


Ligand A86 C 305

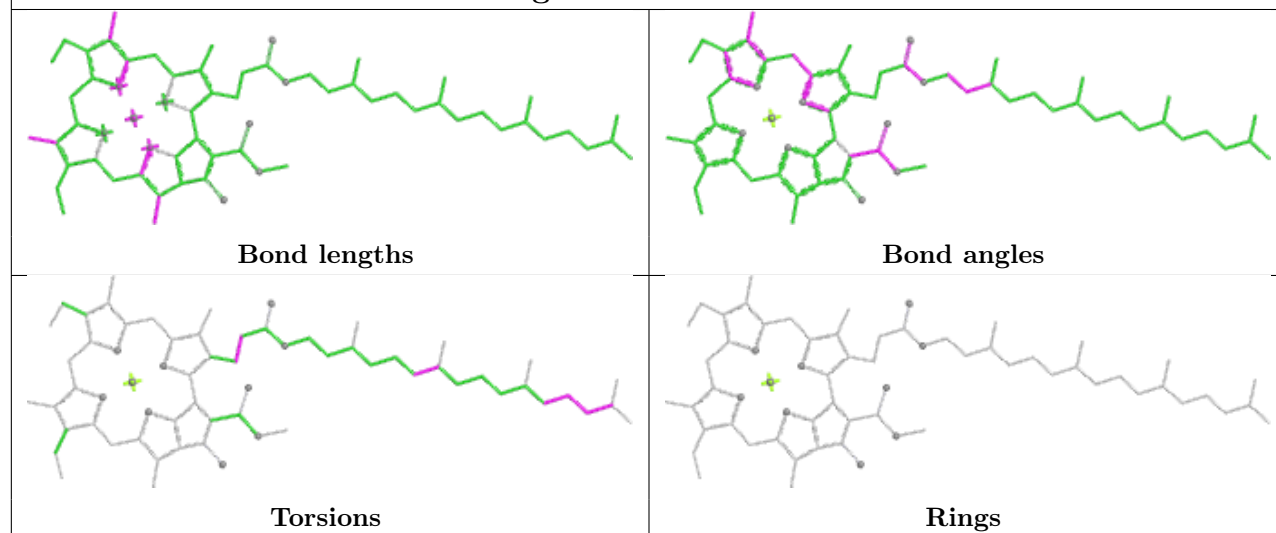


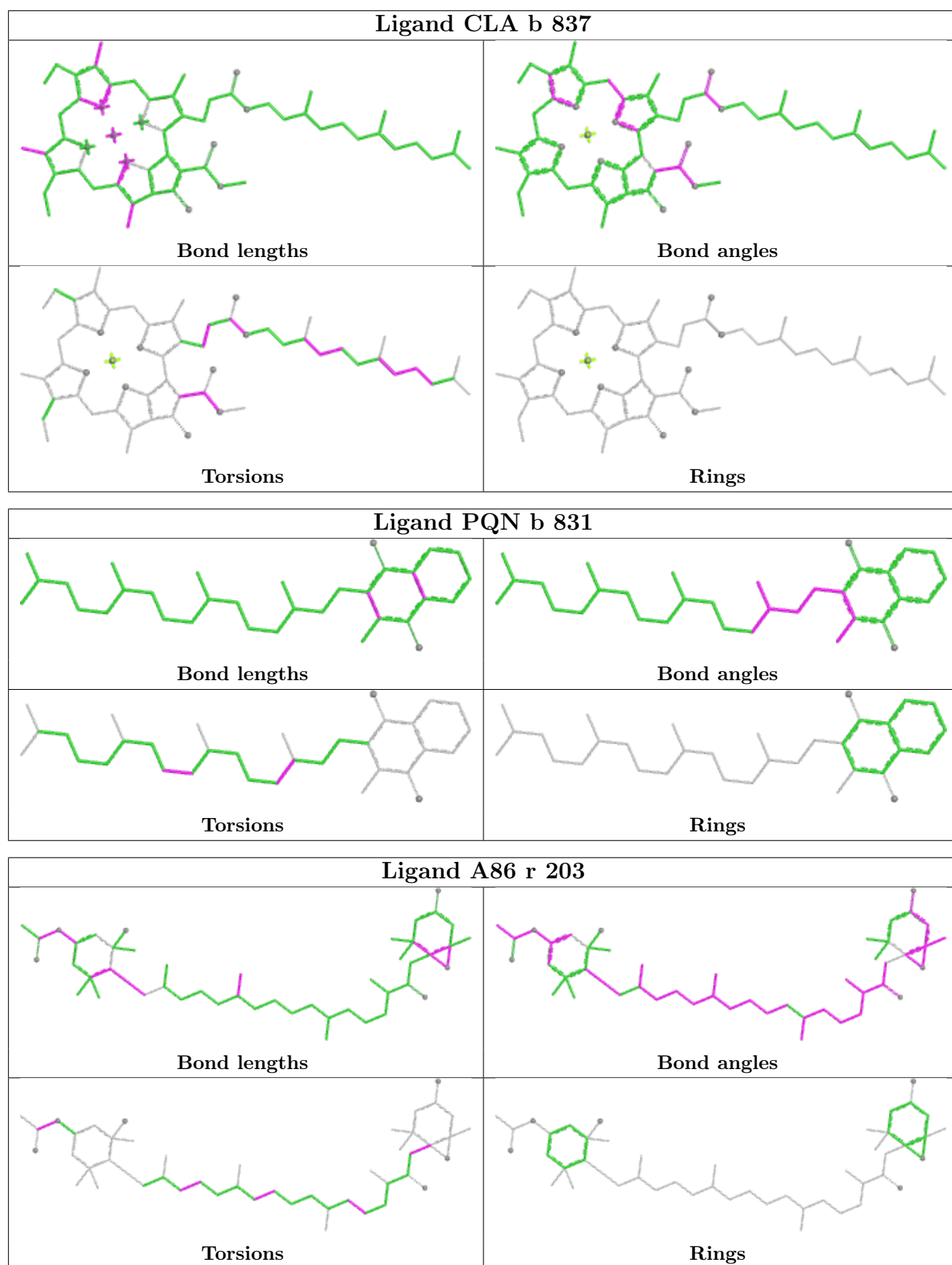


Ligand CLA b 813

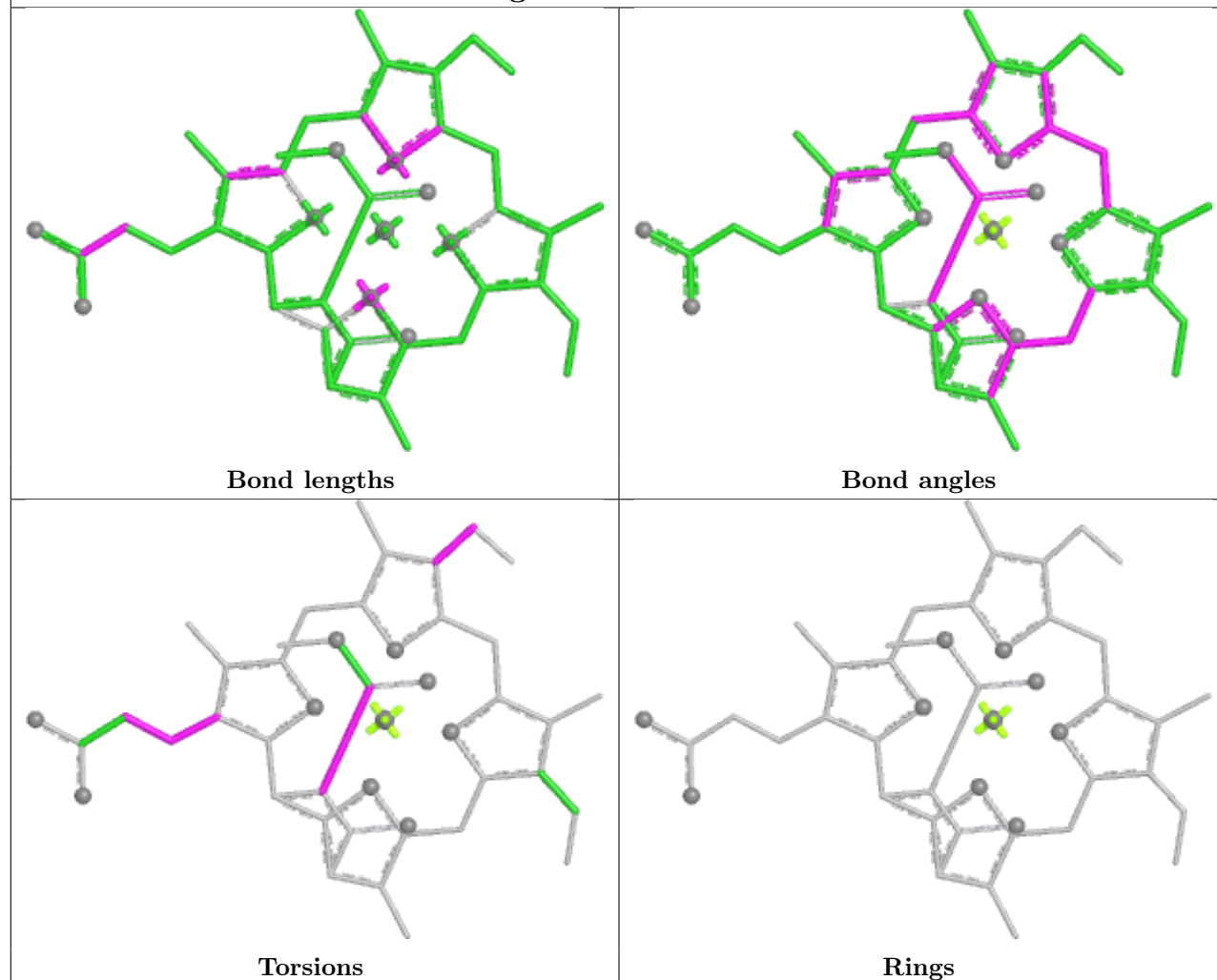


Ligand CLA C 307

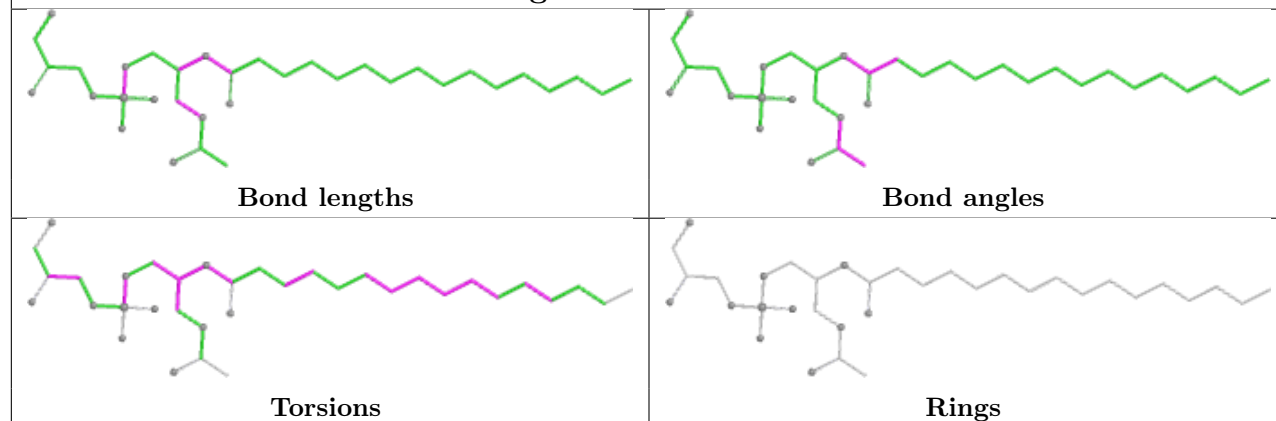


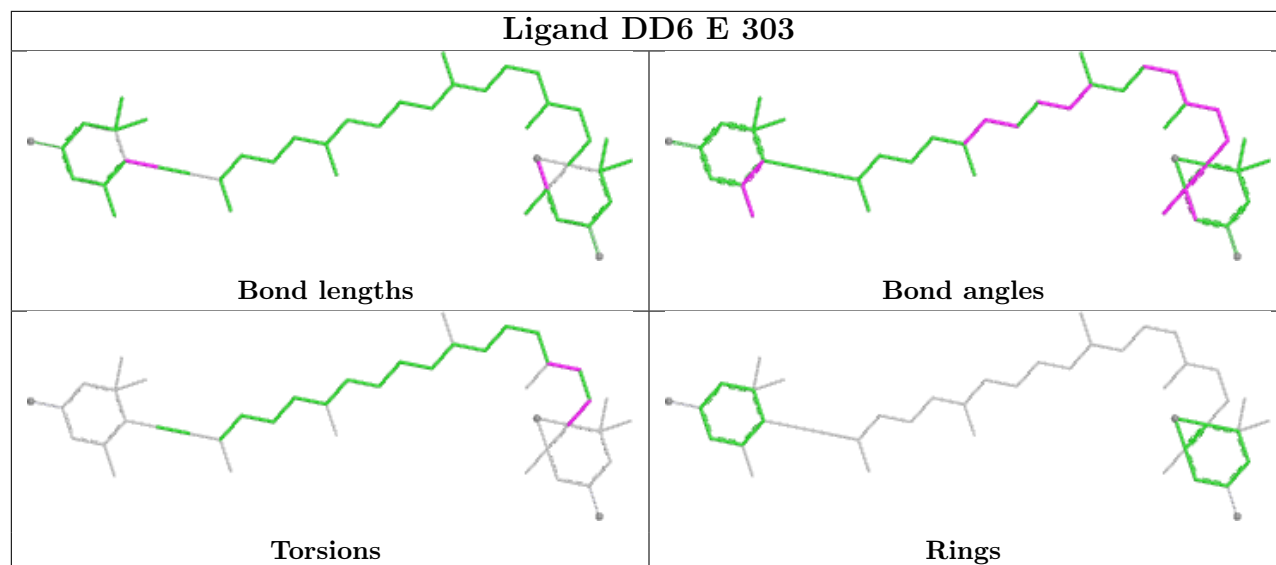
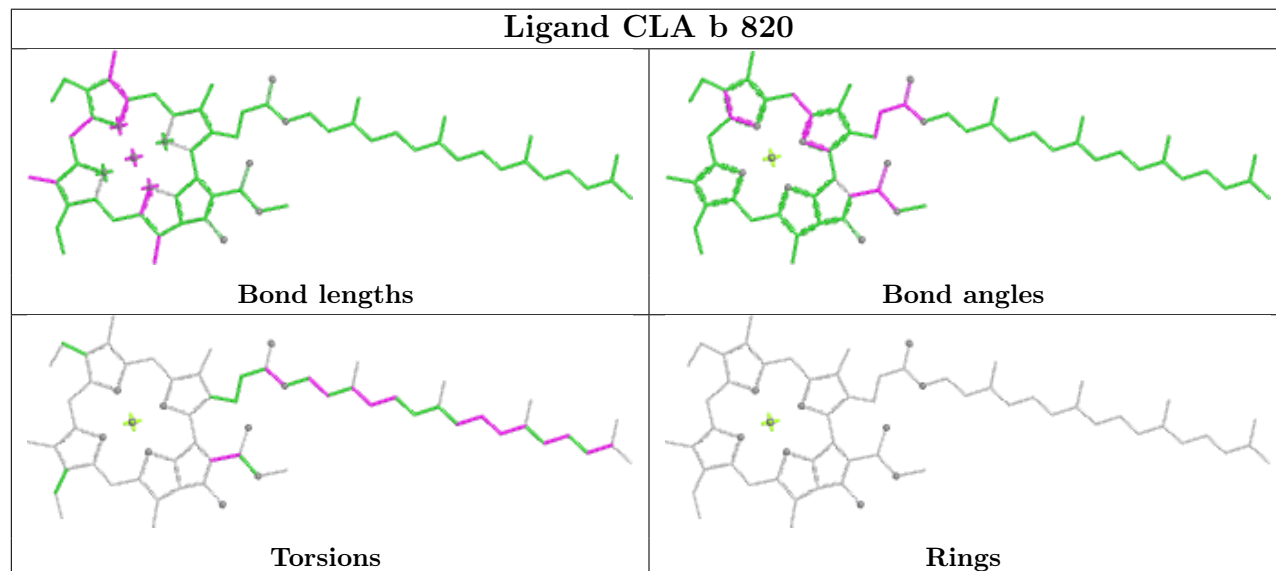


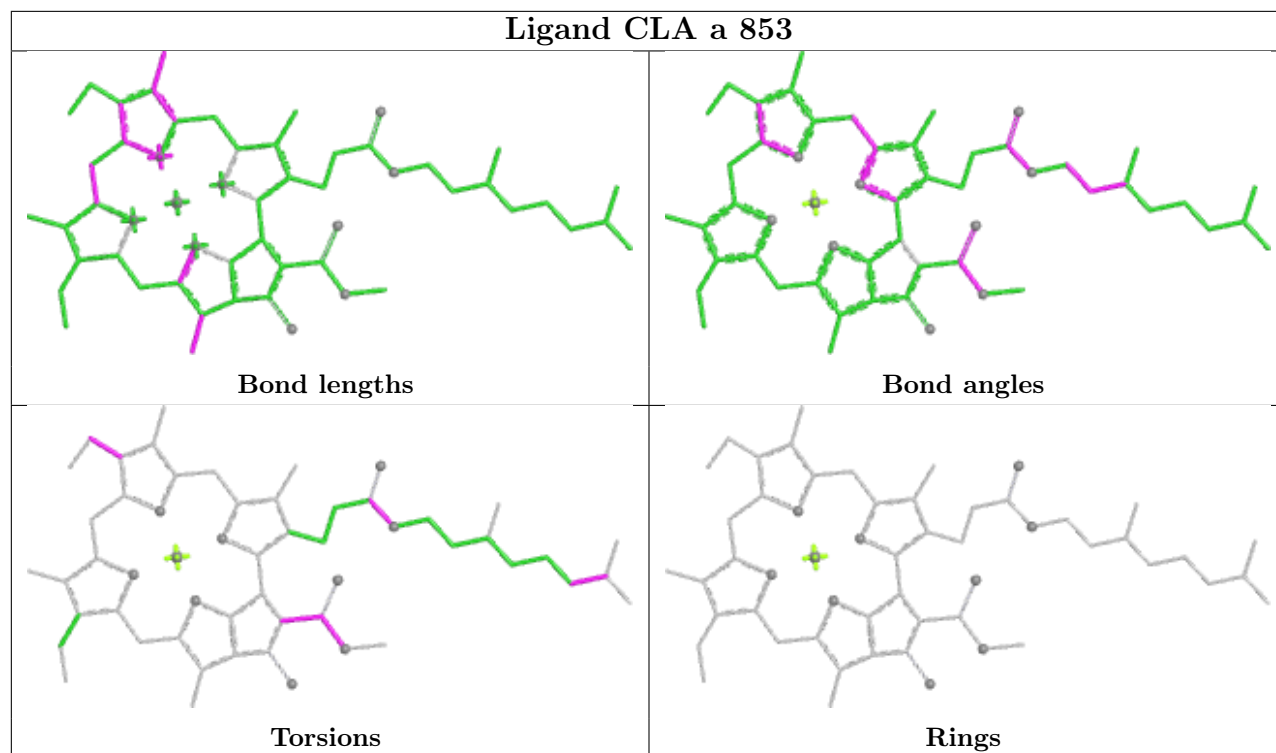
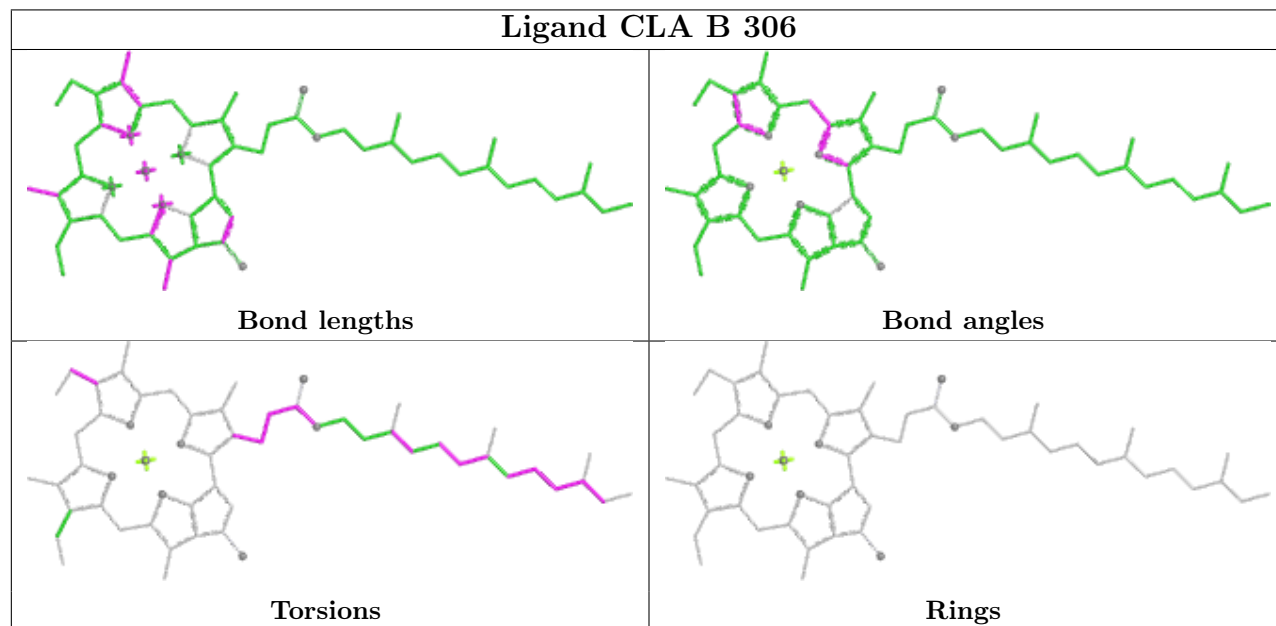
Ligand KC1 C 308



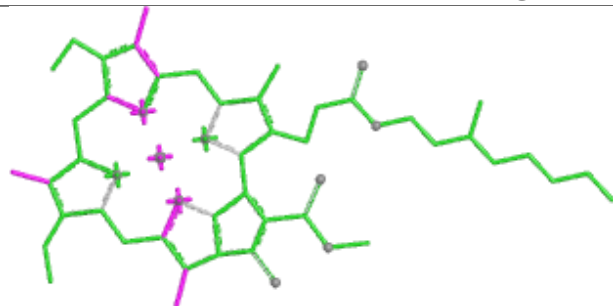
Ligand LHG H 316



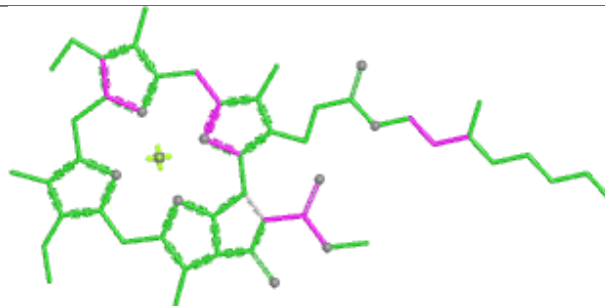
Ligand DD6 E 303**Ligand CLA b 820**

Ligand CLA a 853**Ligand CLA B 306**

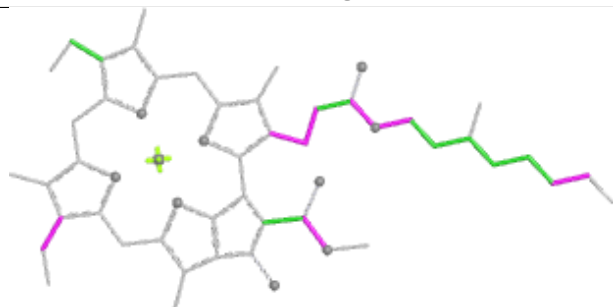
Ligand CLA H 305



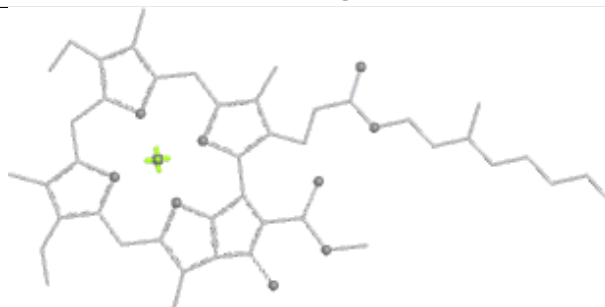
Bond lengths



Bond angles

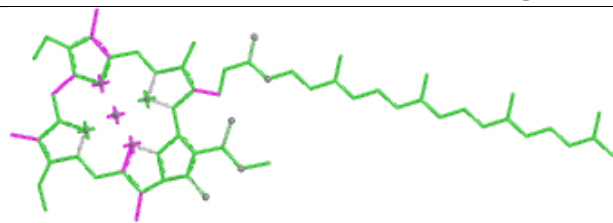


Torsions

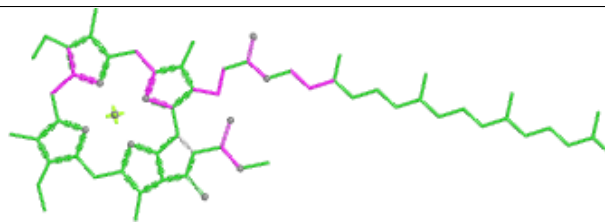


Rings

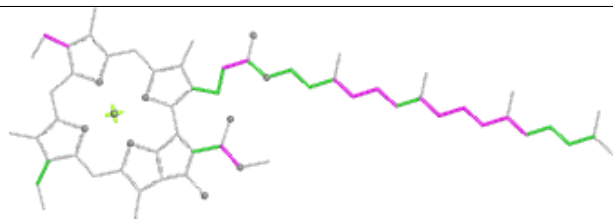
Ligand CLA b 803



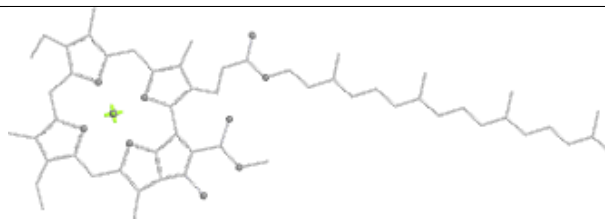
Bond lengths



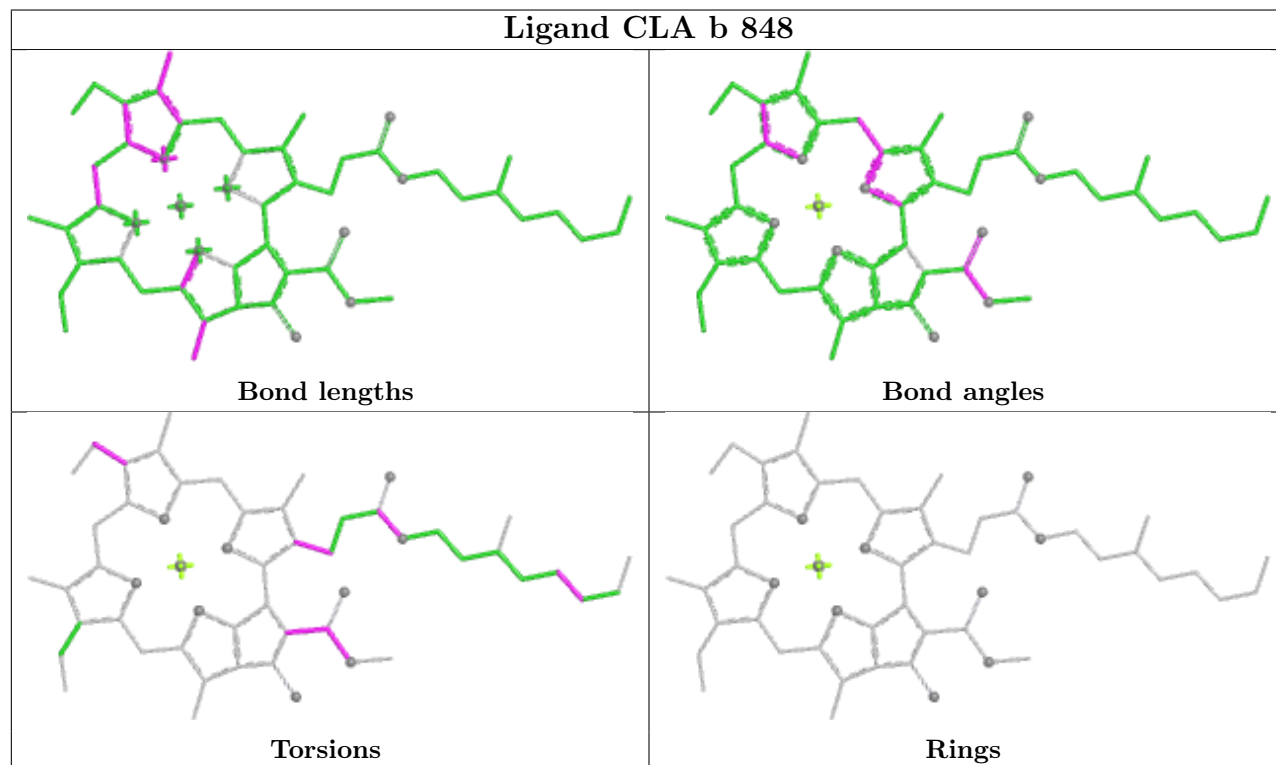
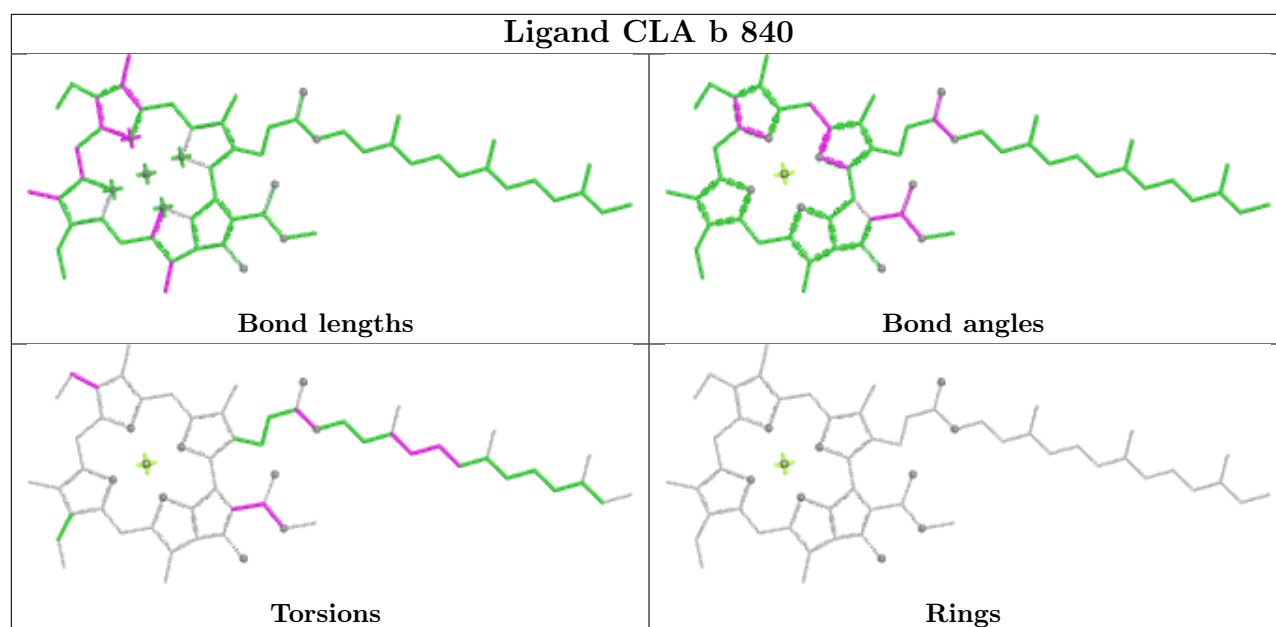
Bond angles

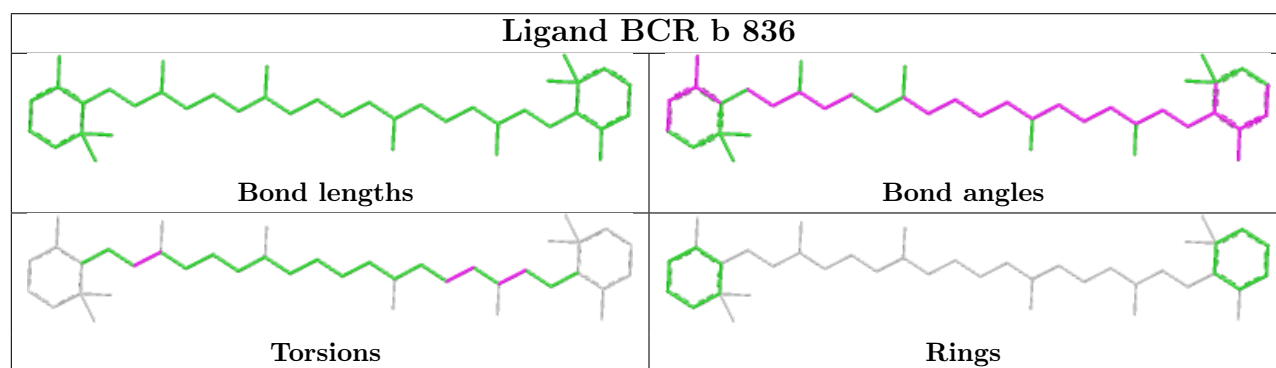
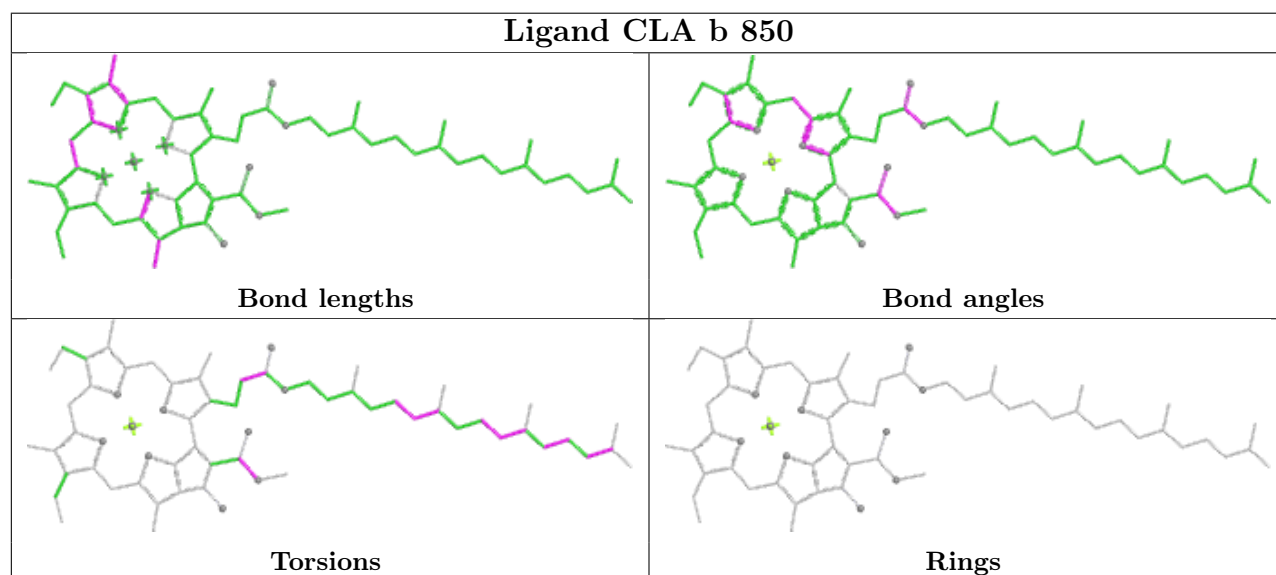
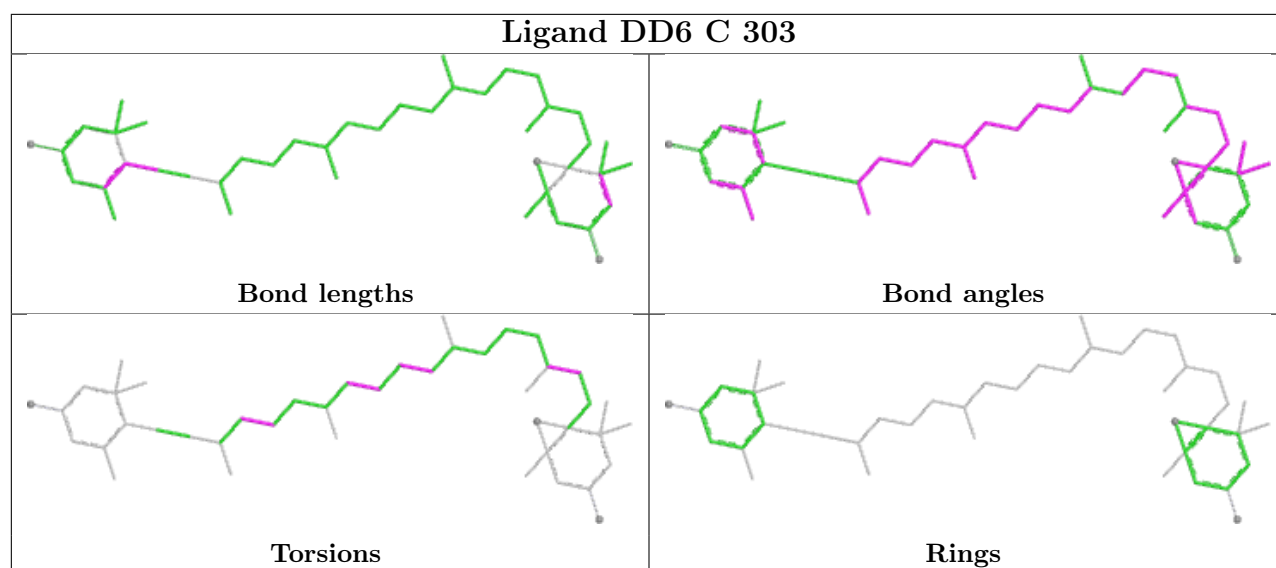


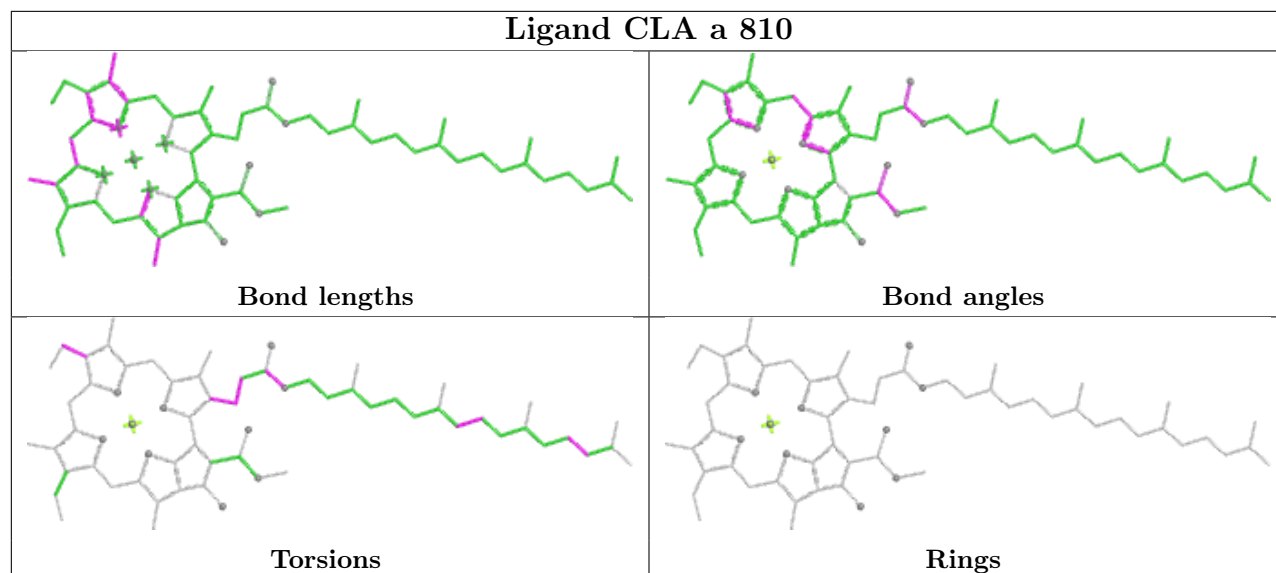
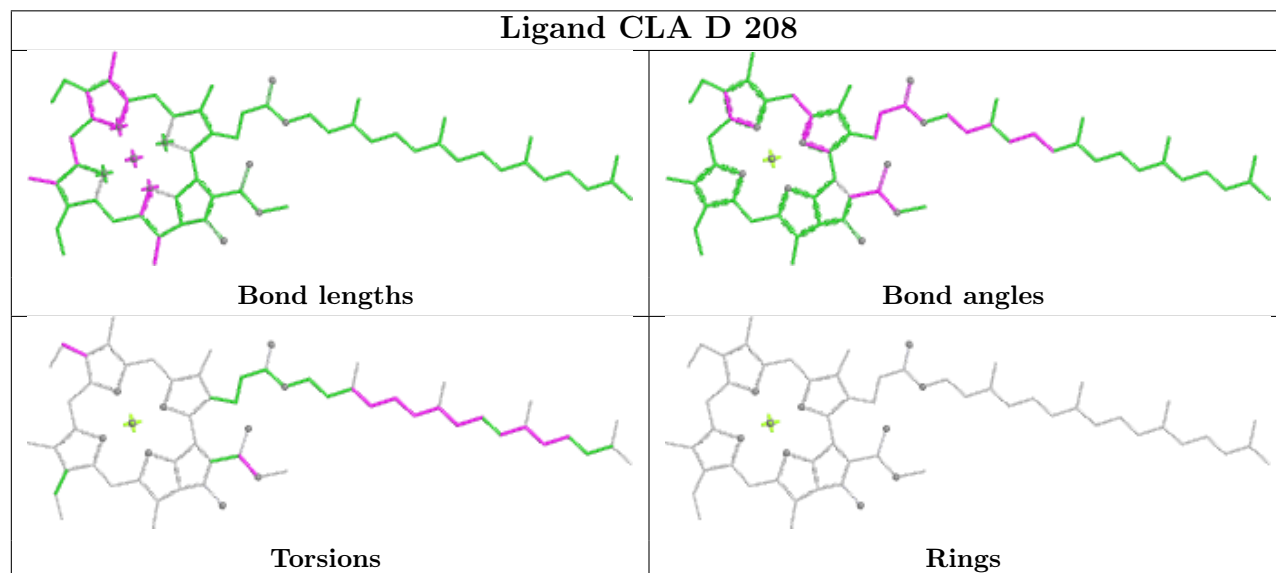
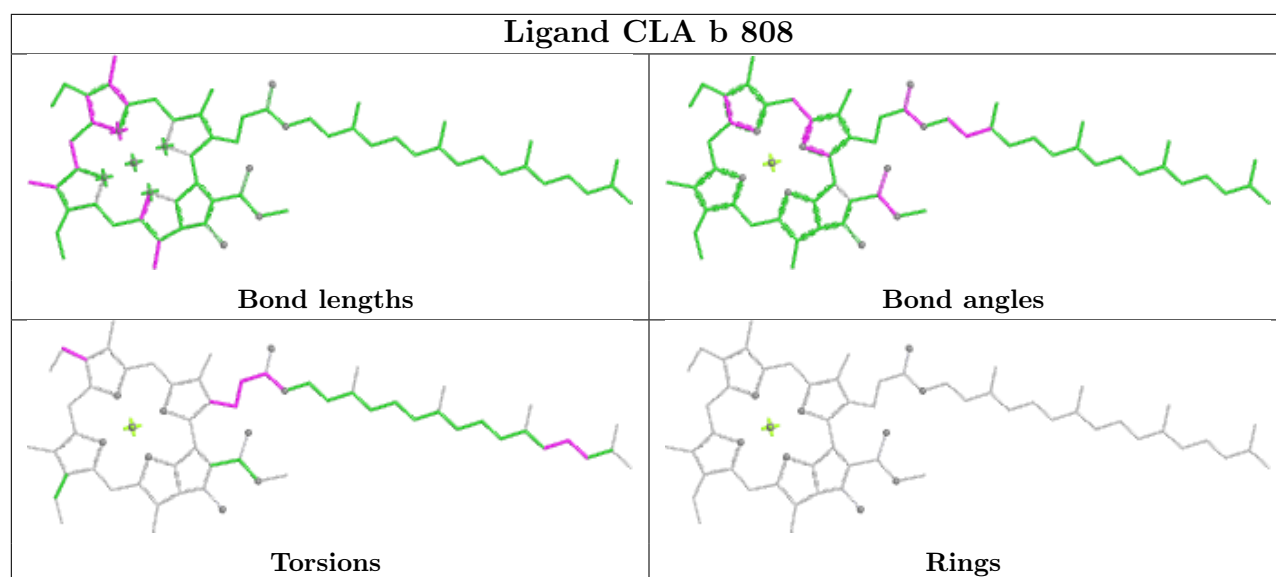
Torsions



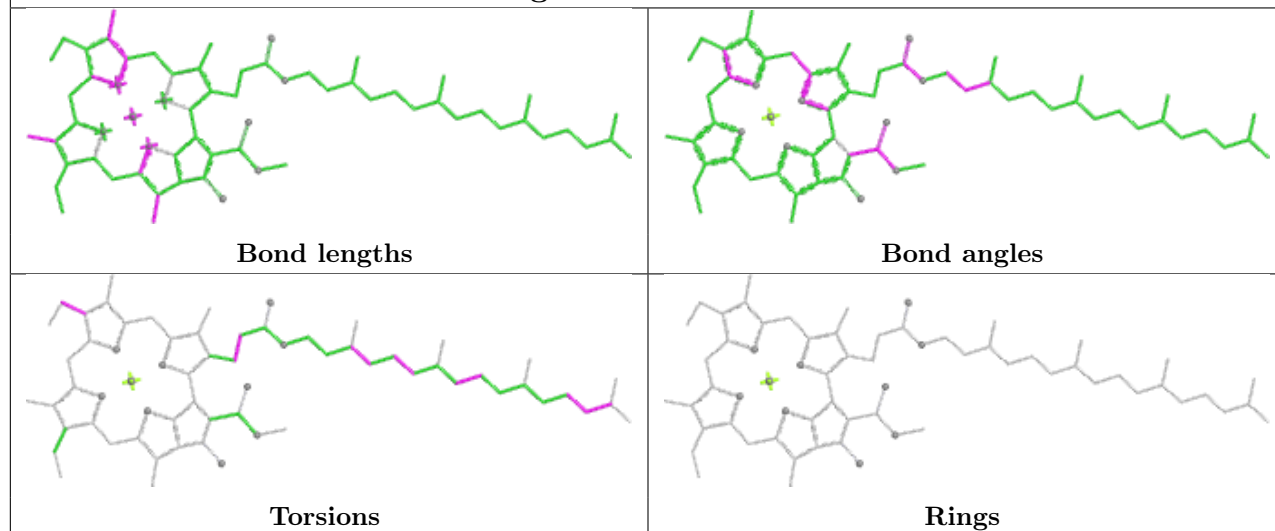
Rings



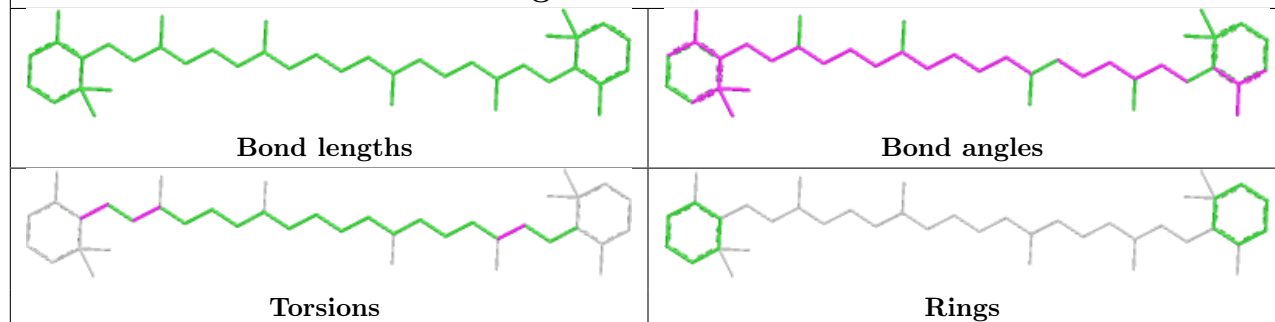




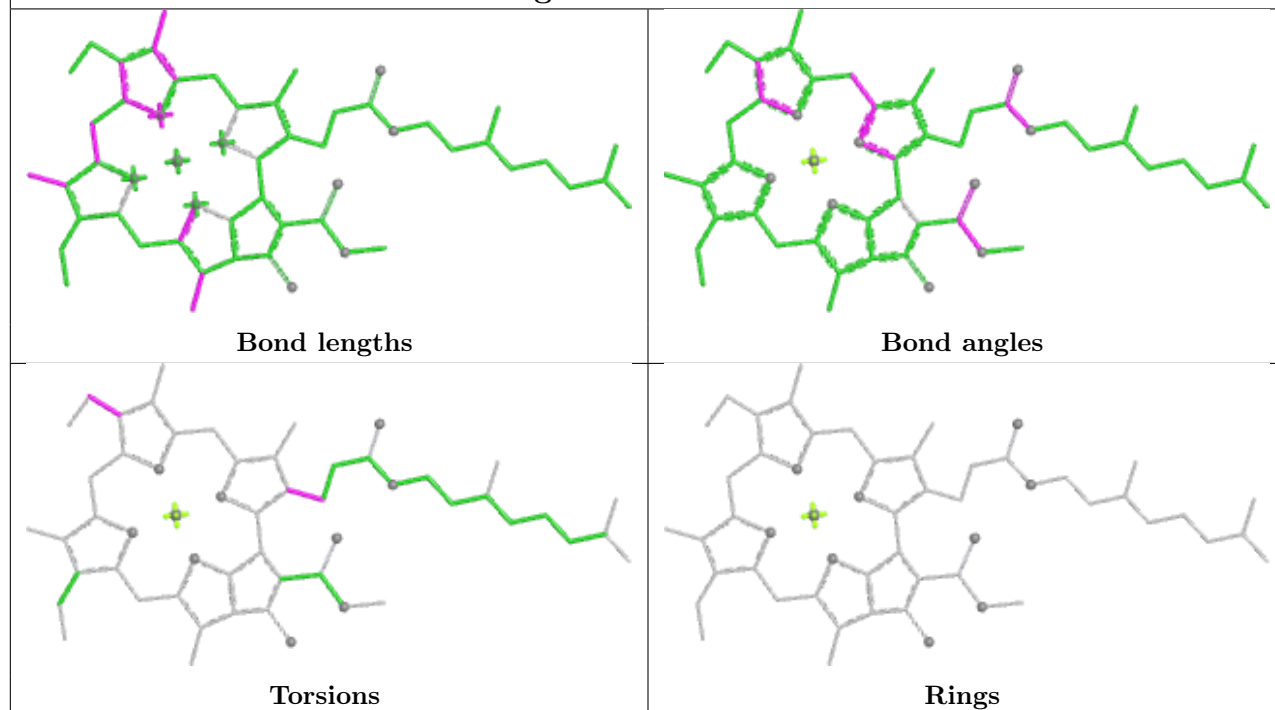
Ligand CLA H 308

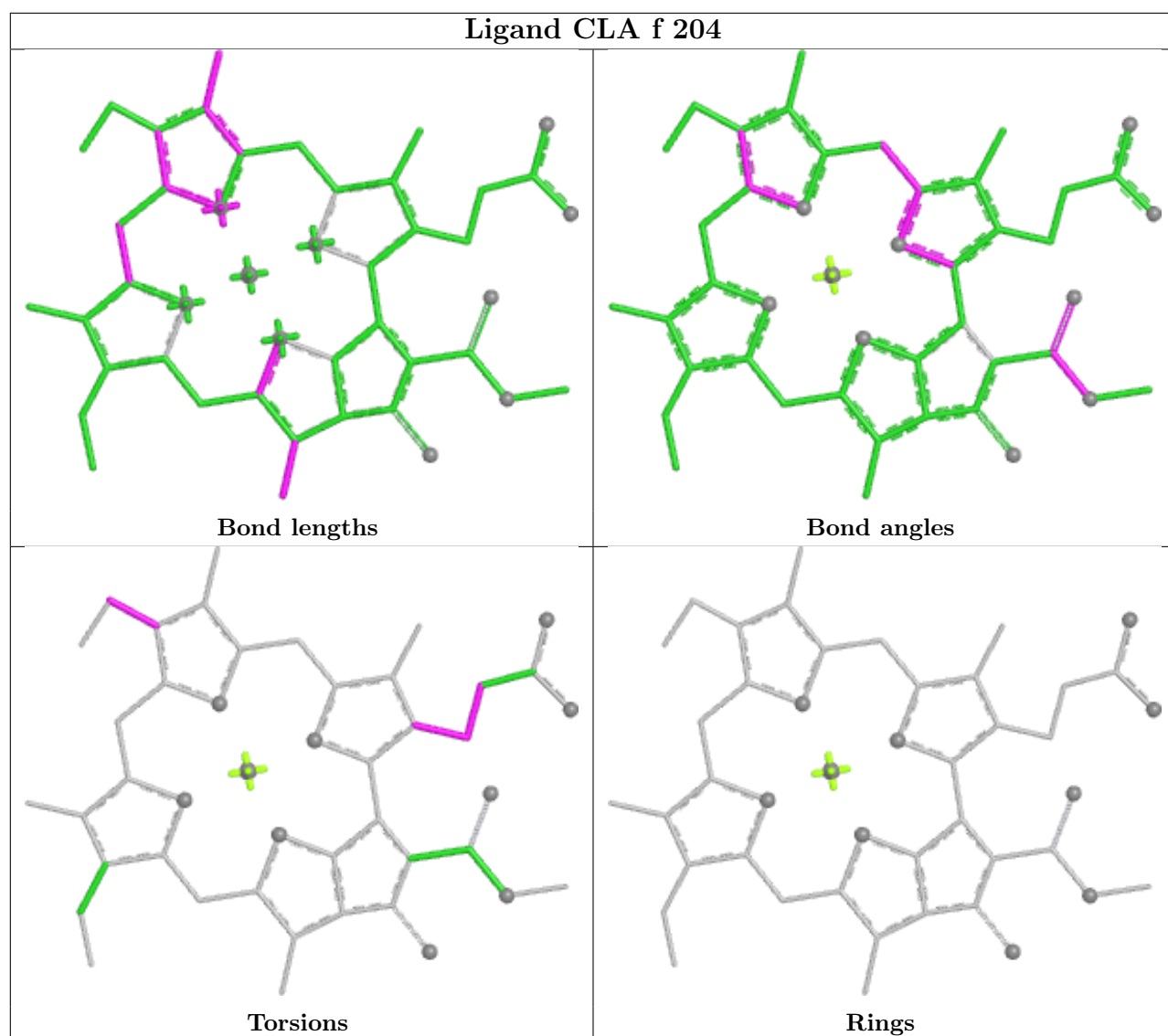


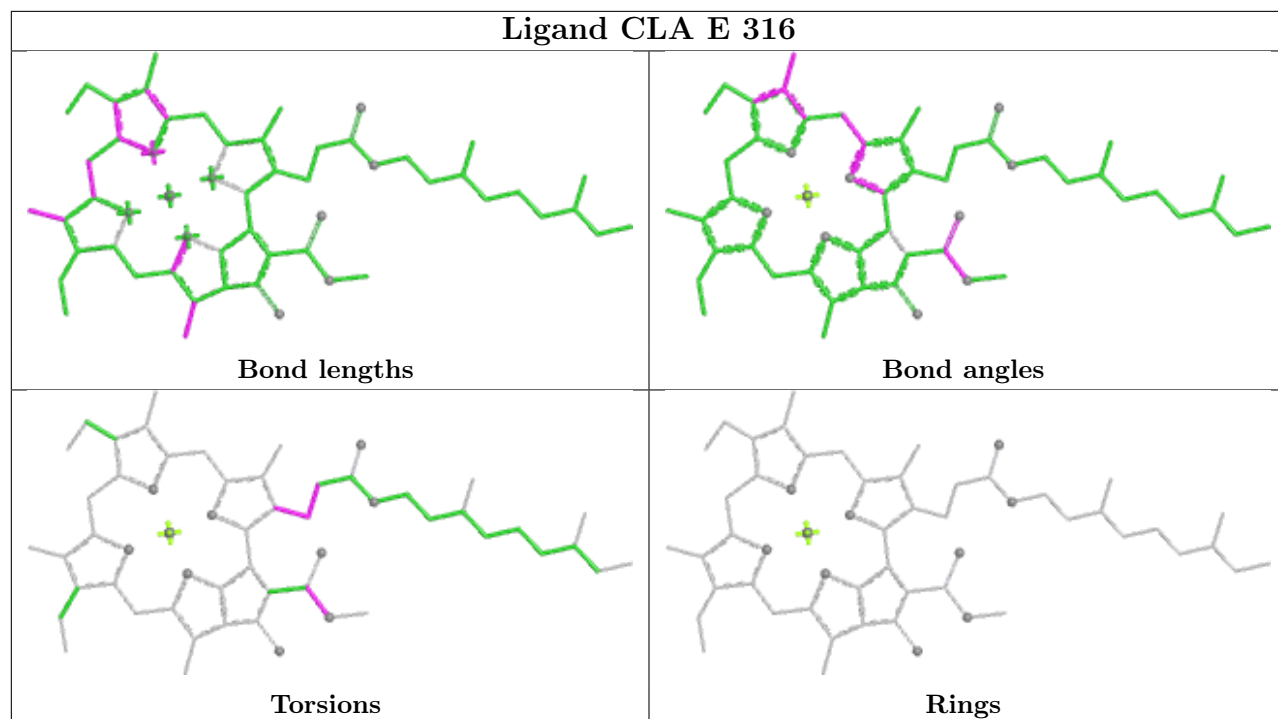
Ligand BCR b 833



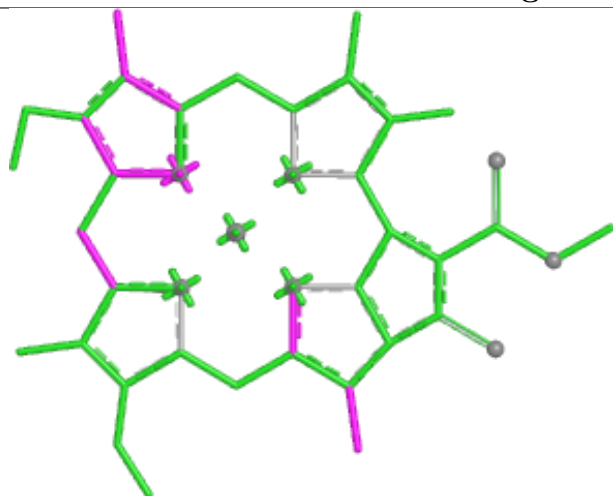
Ligand CLA b 809



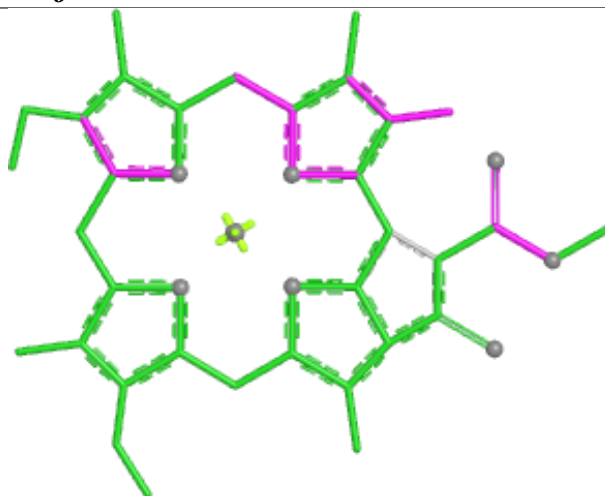




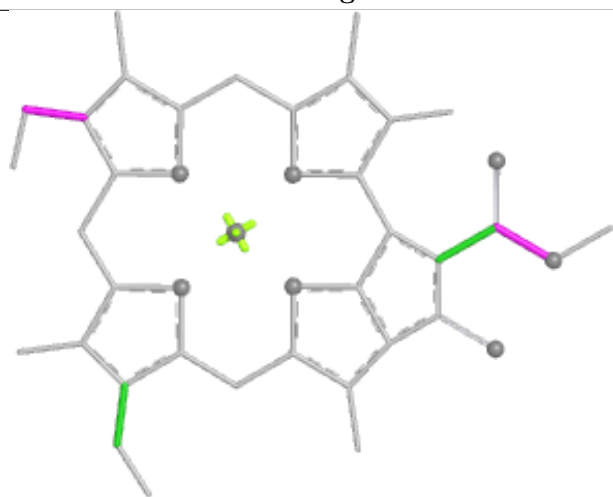
Ligand CLA j 104



Bond lengths



Bond angles

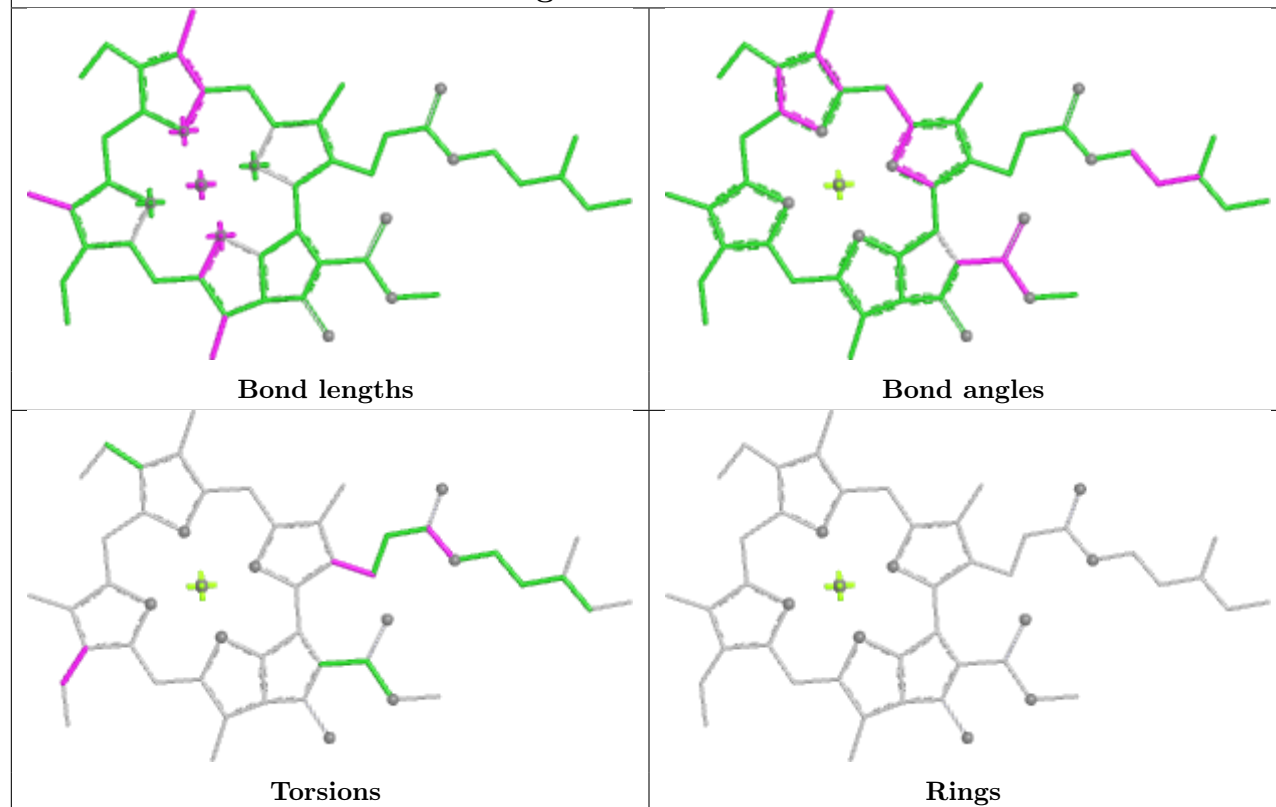


Torsions

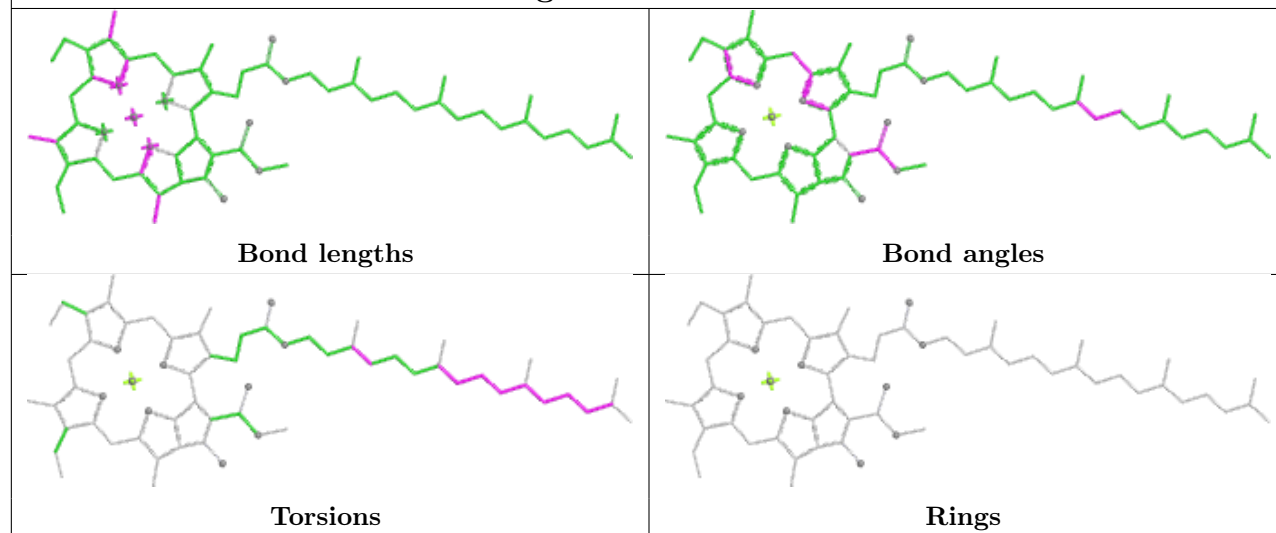


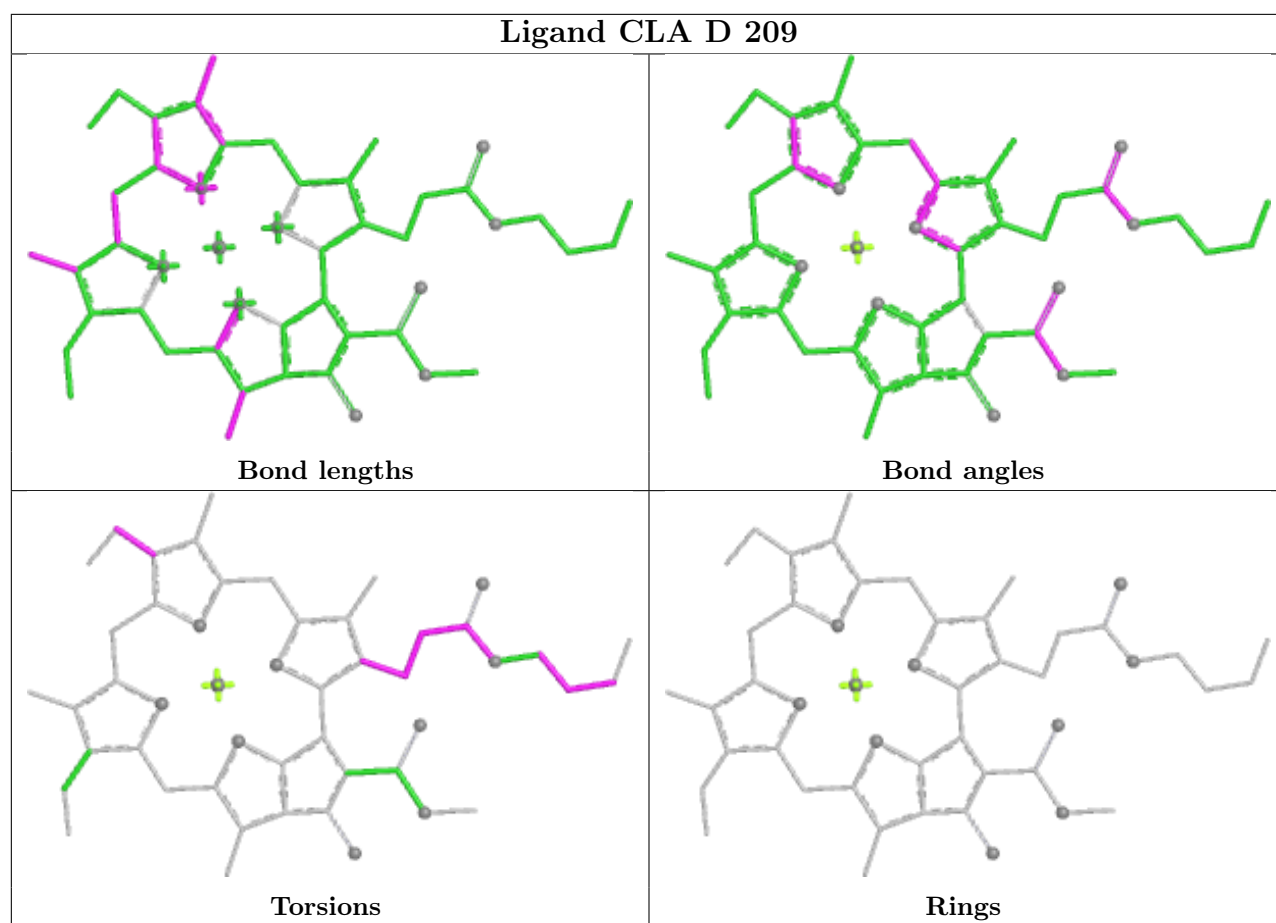
Rings

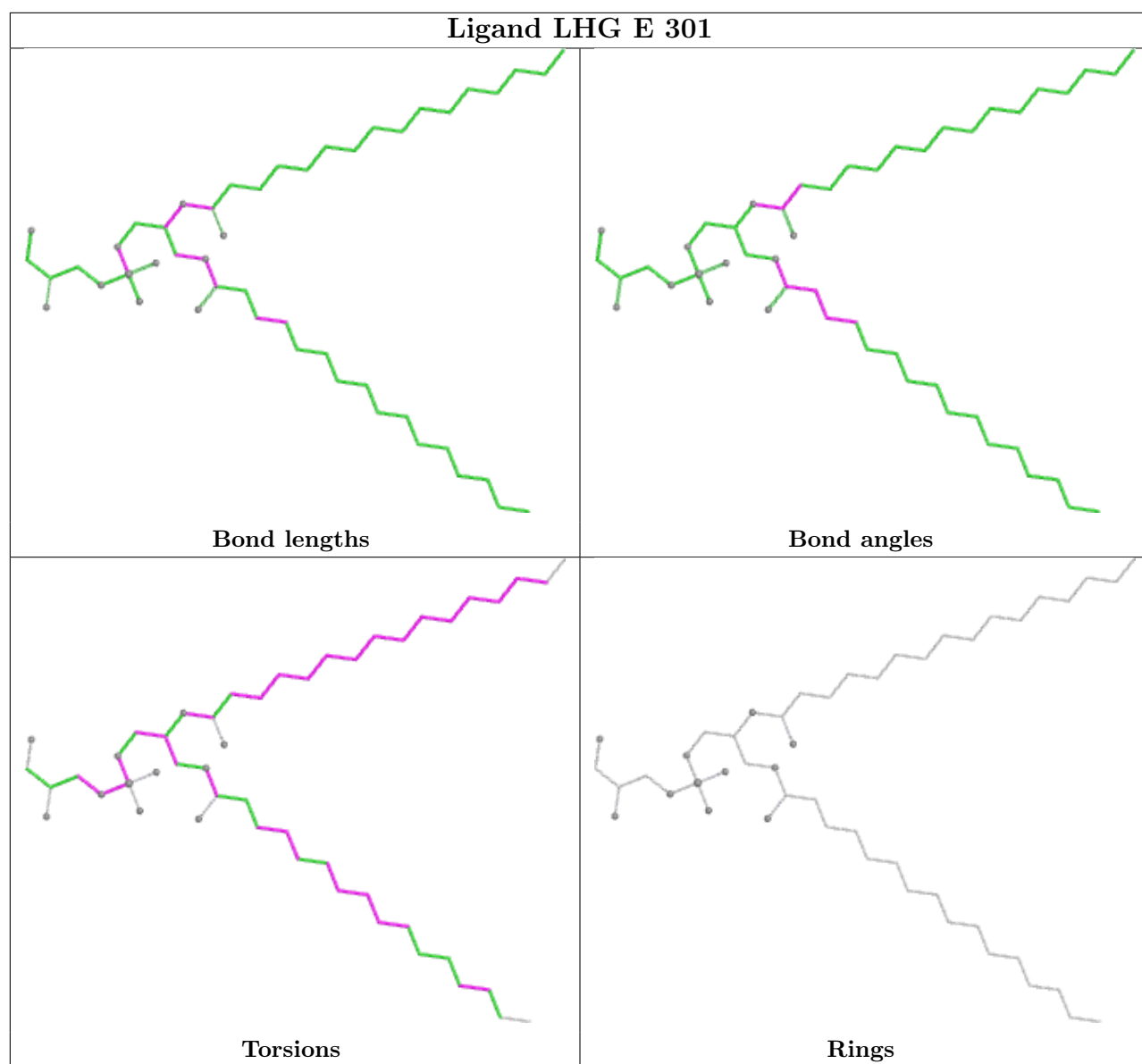
Ligand CLA a 818

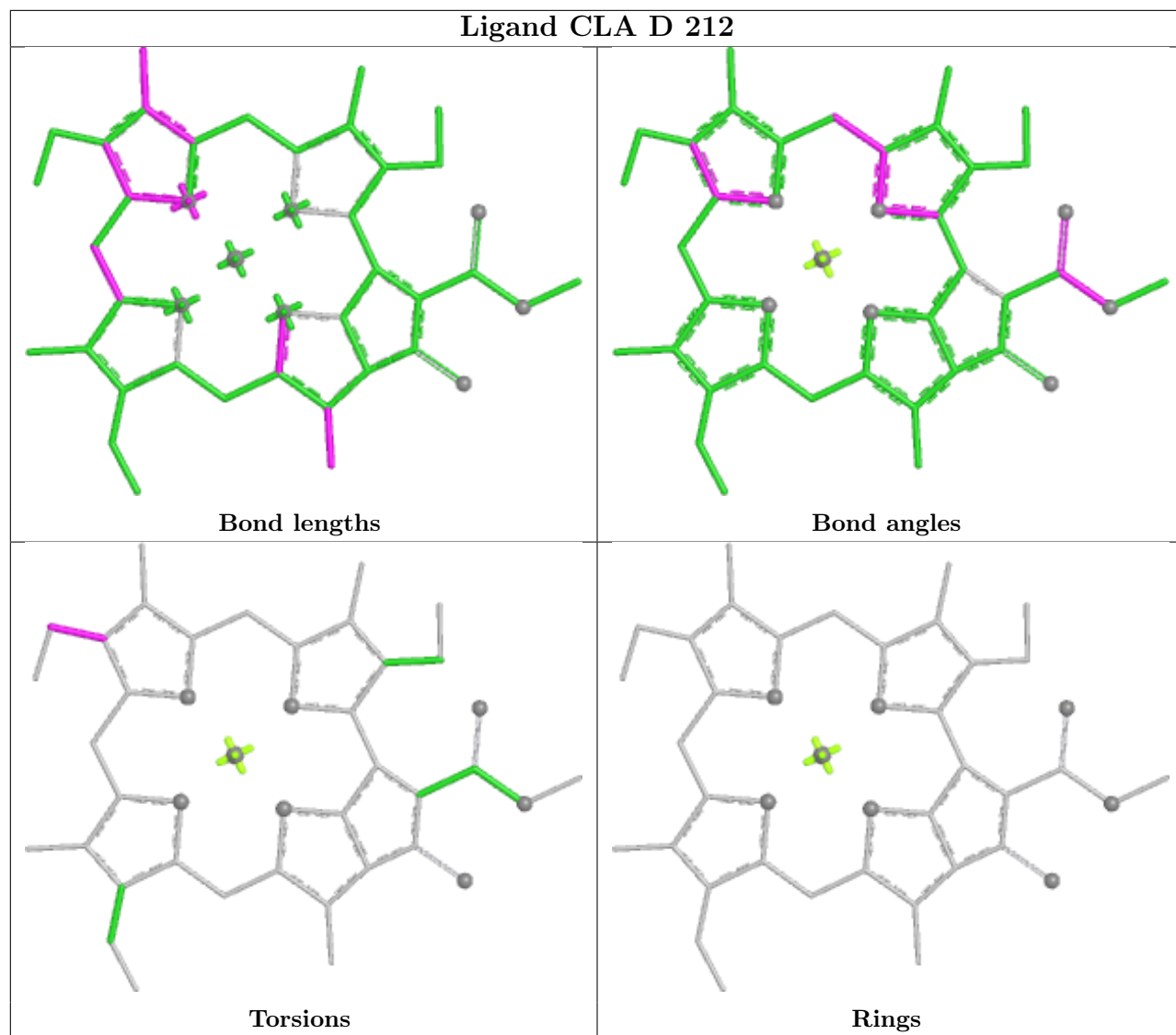


Ligand CLA a 840

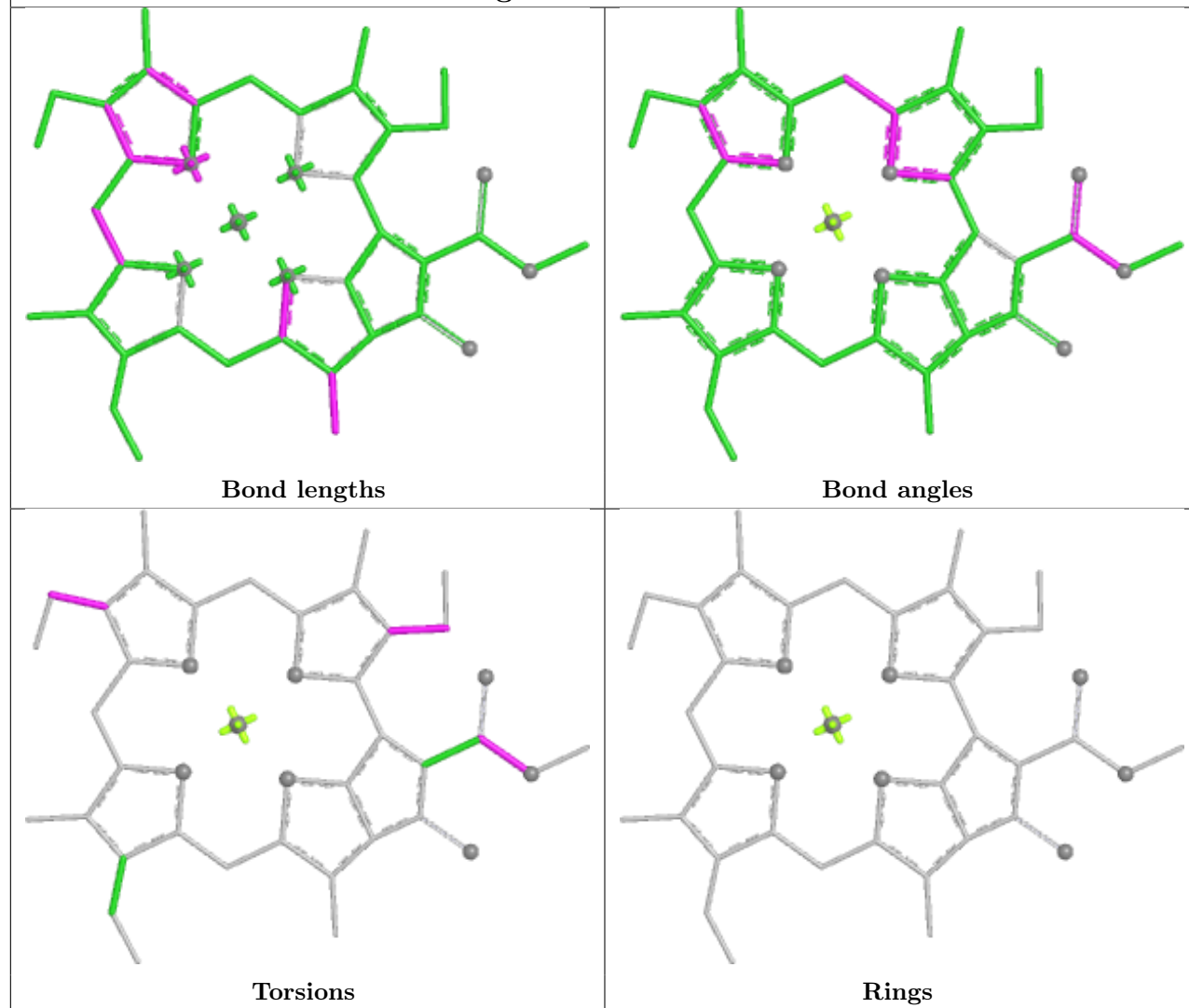




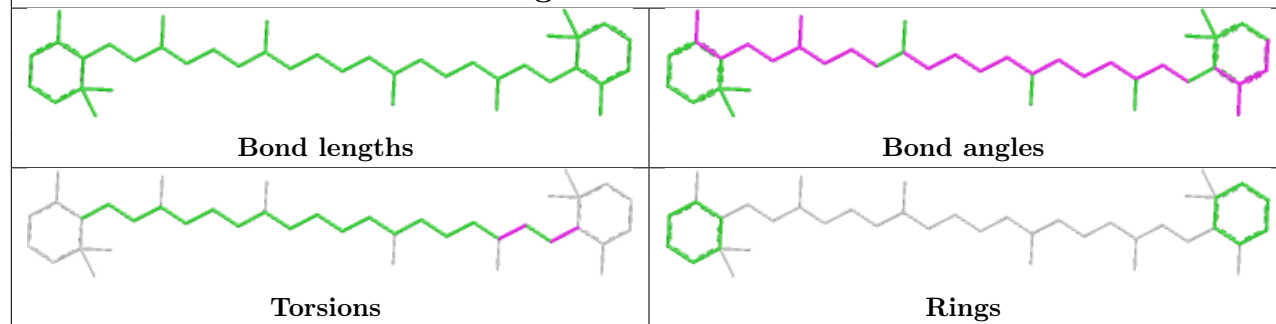


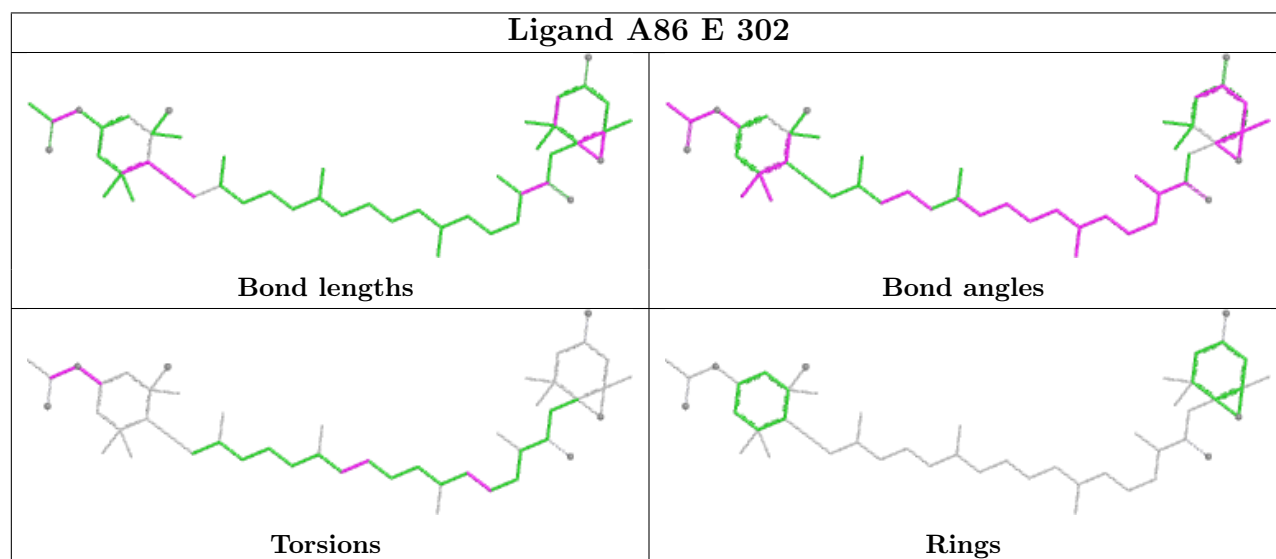
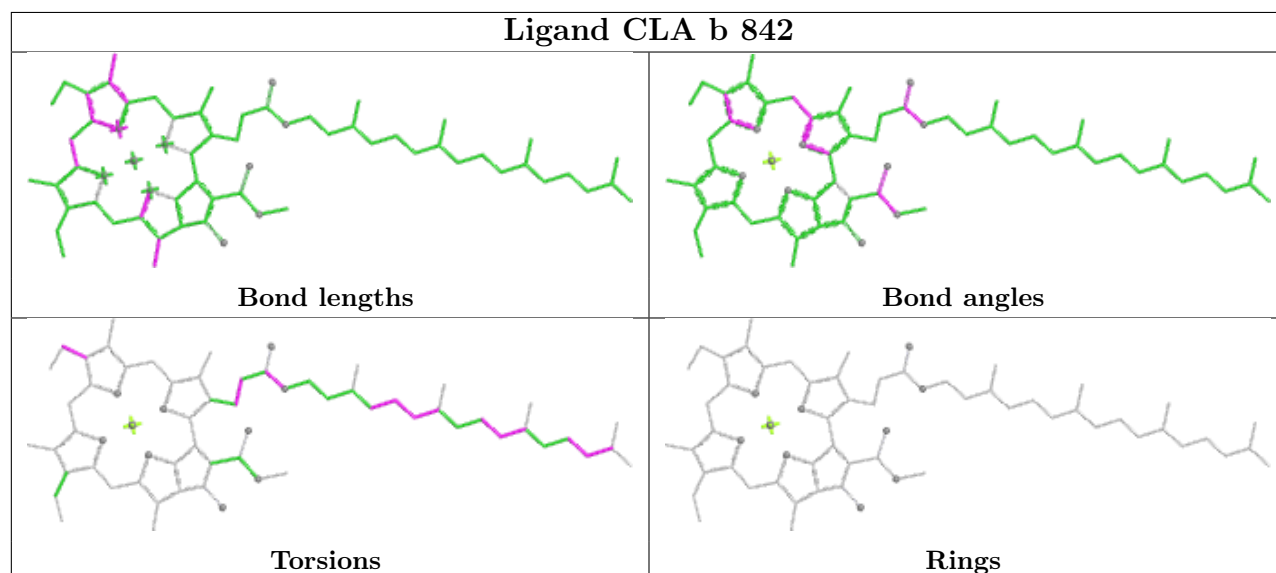
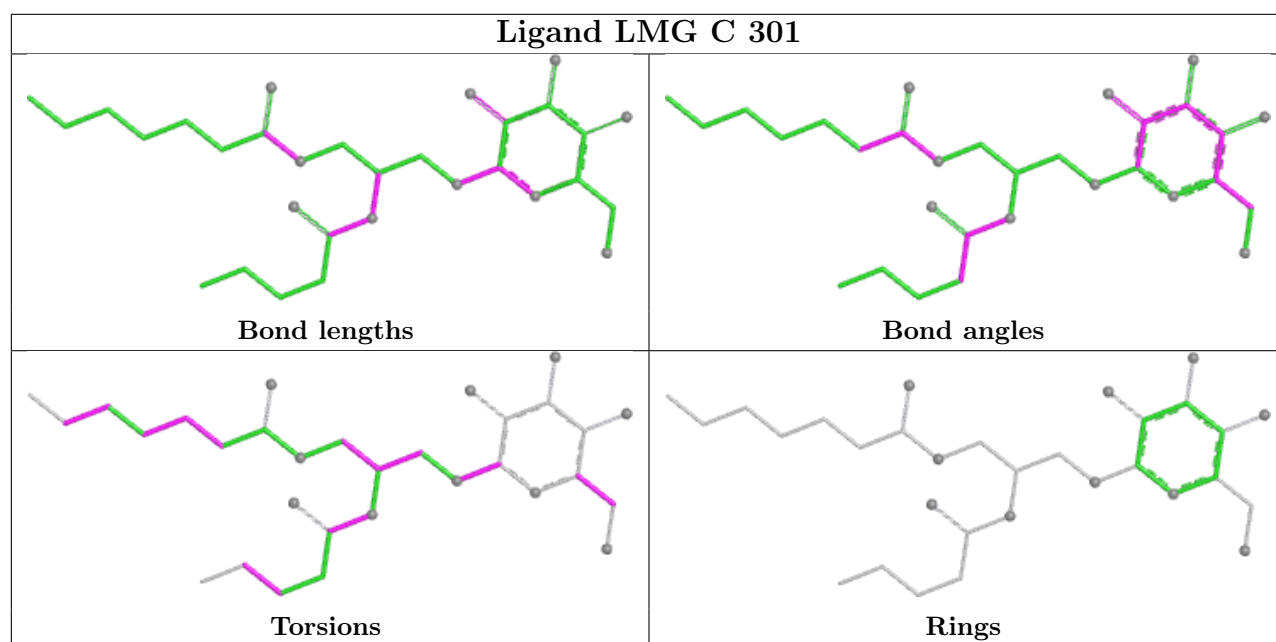


Ligand CLA C 315

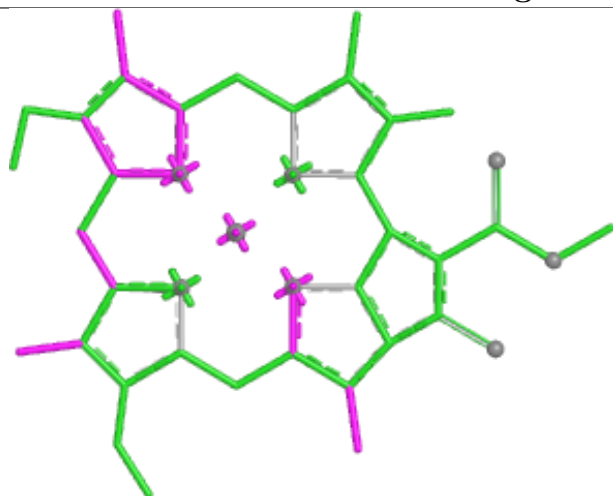


Ligand BCR a 834

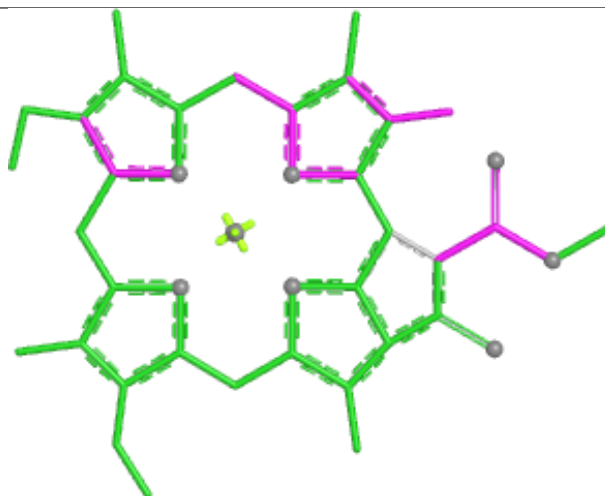




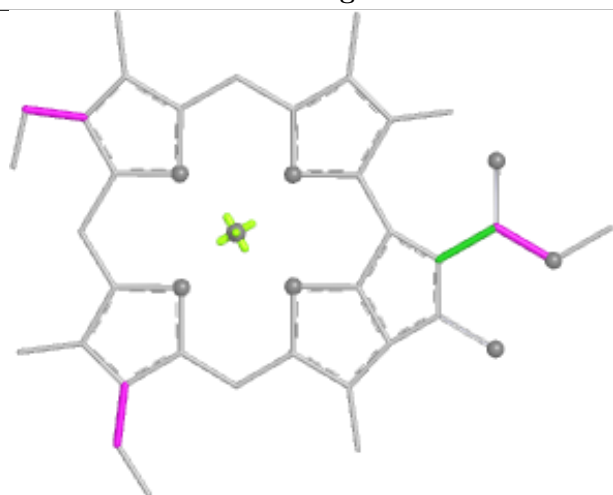
Ligand CLA C 314



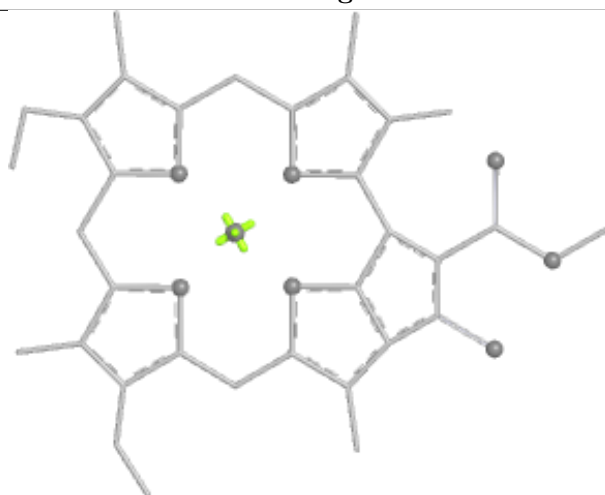
Bond lengths



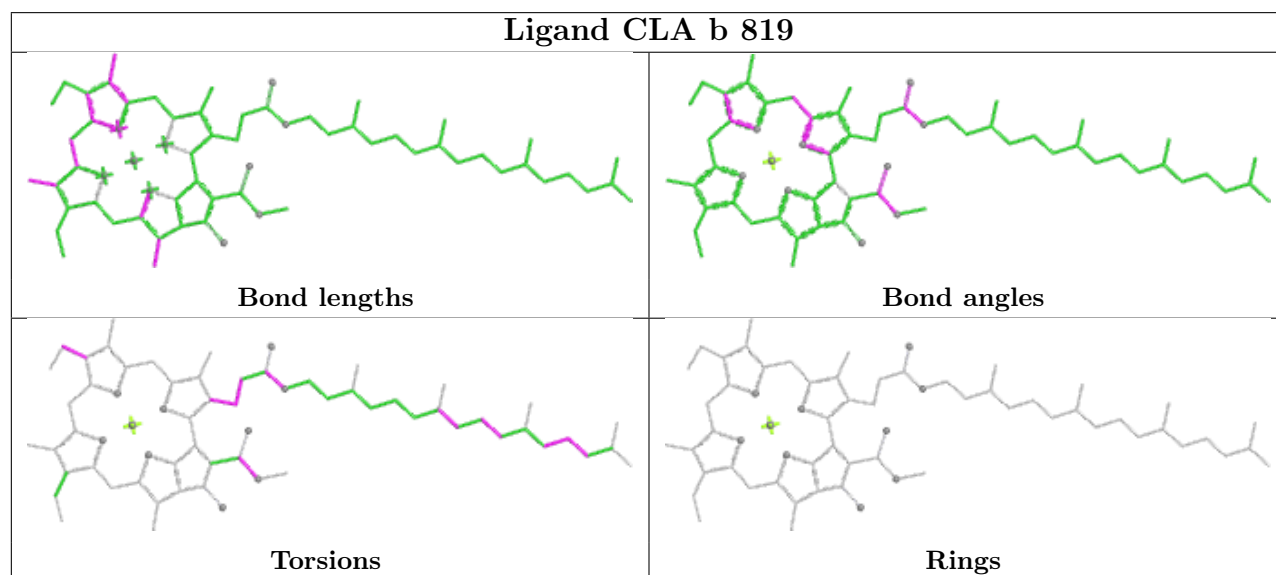
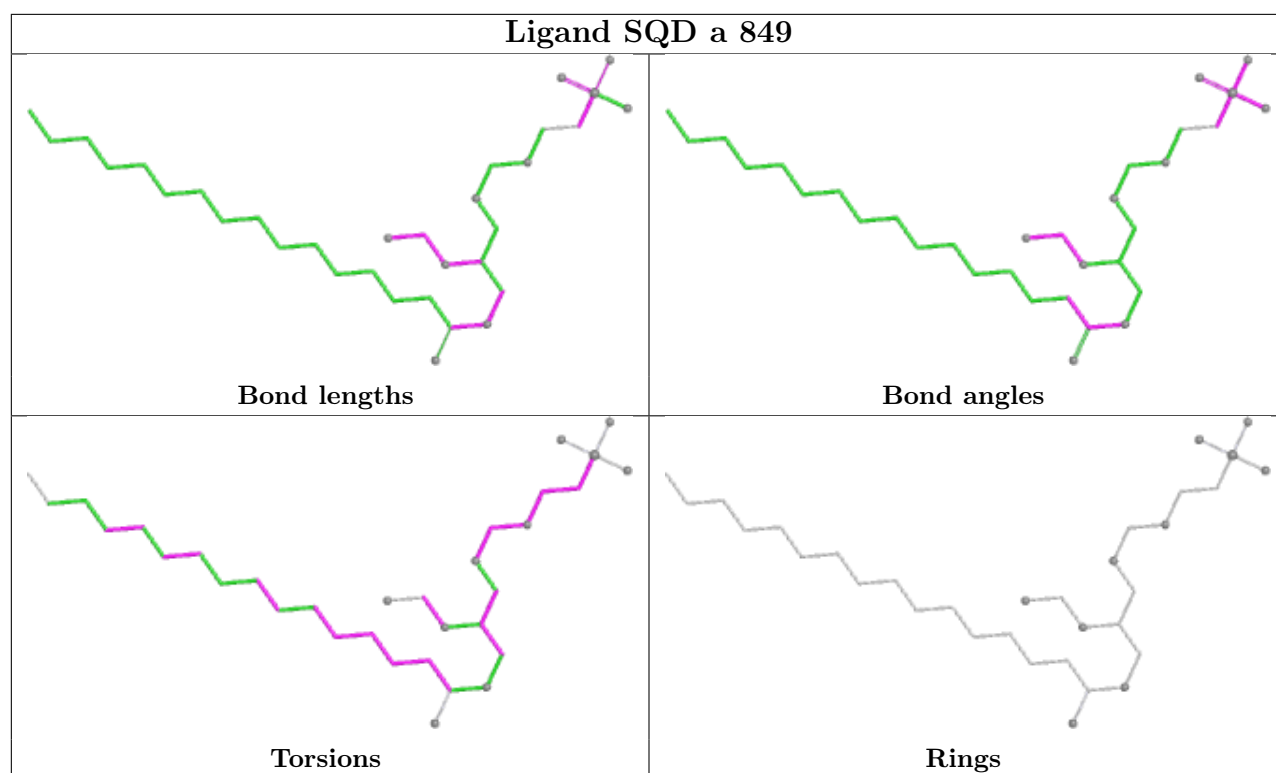
Bond angles



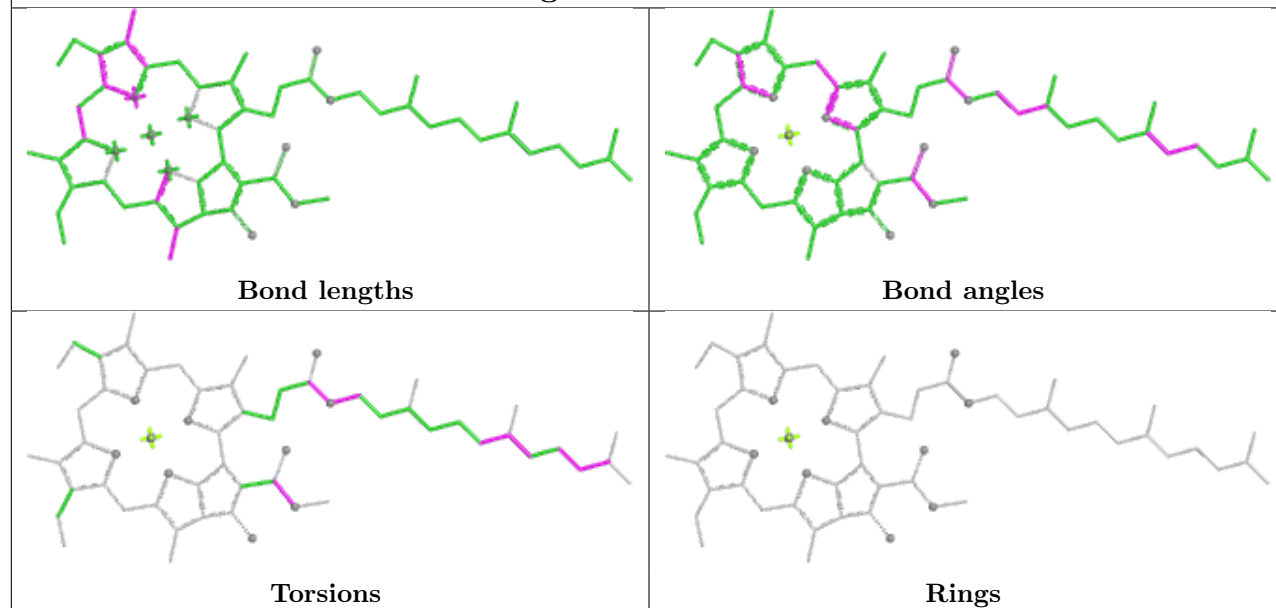
Torsions



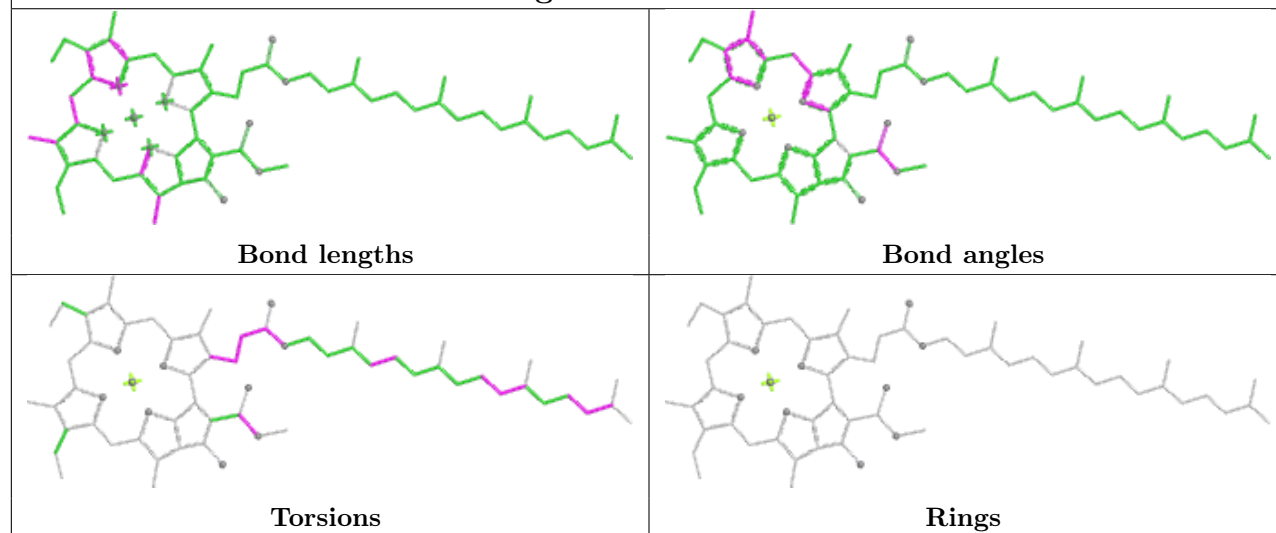
Rings



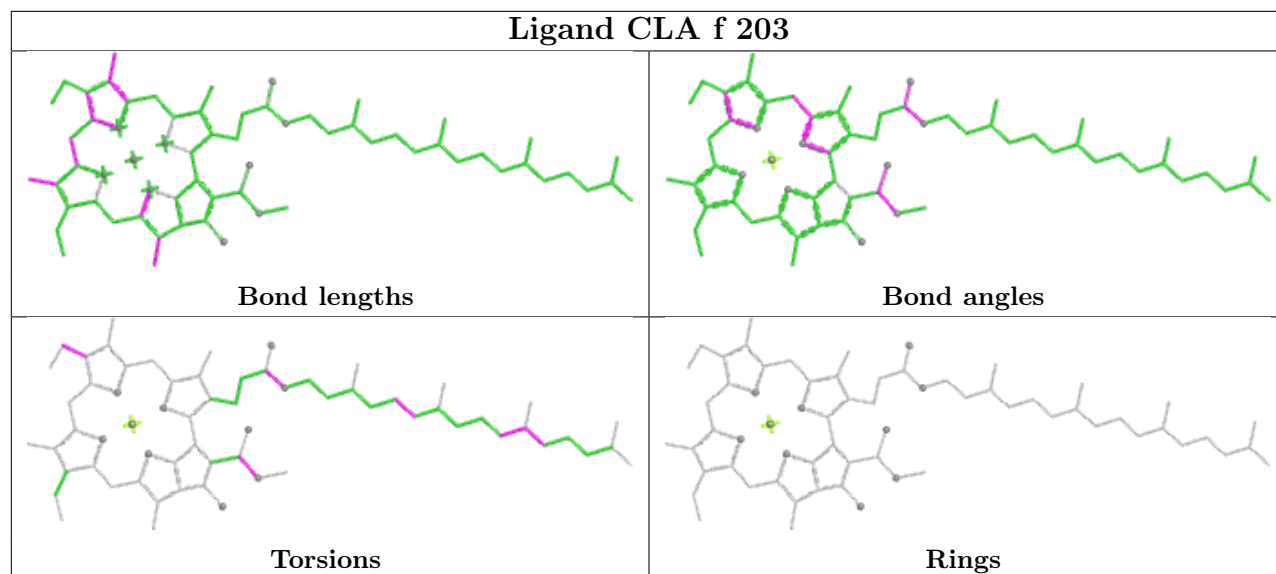
Ligand CLA i 101



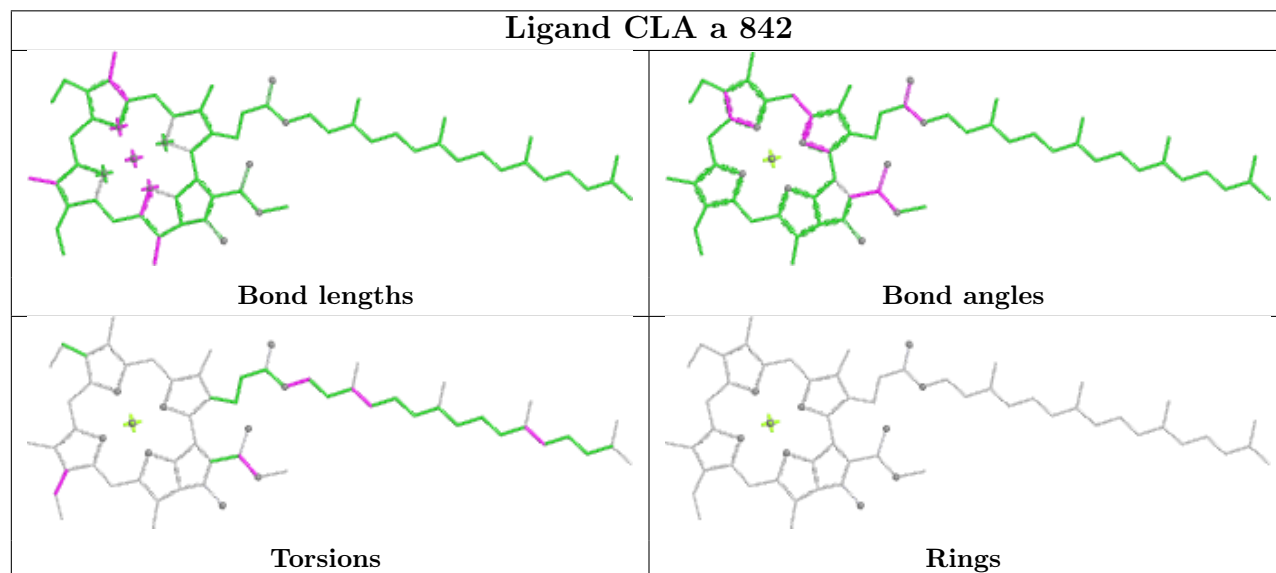
Ligand CLA b 822



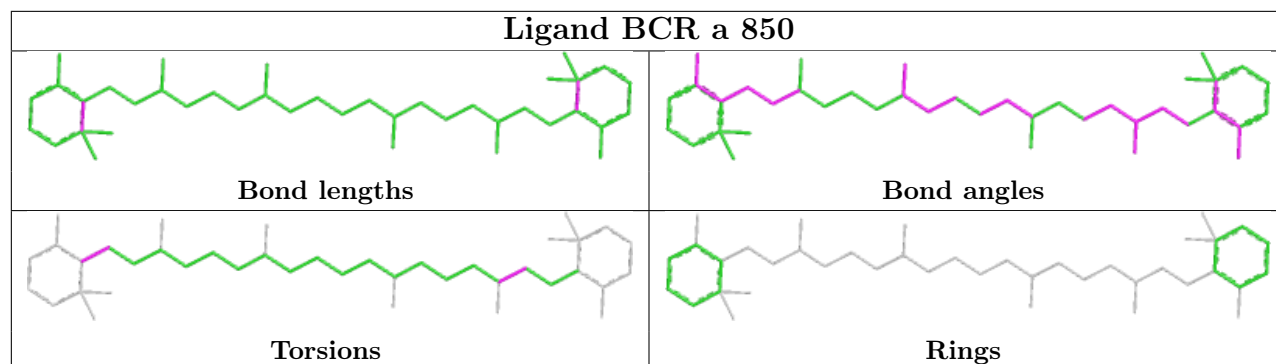
Ligand CLA f 203



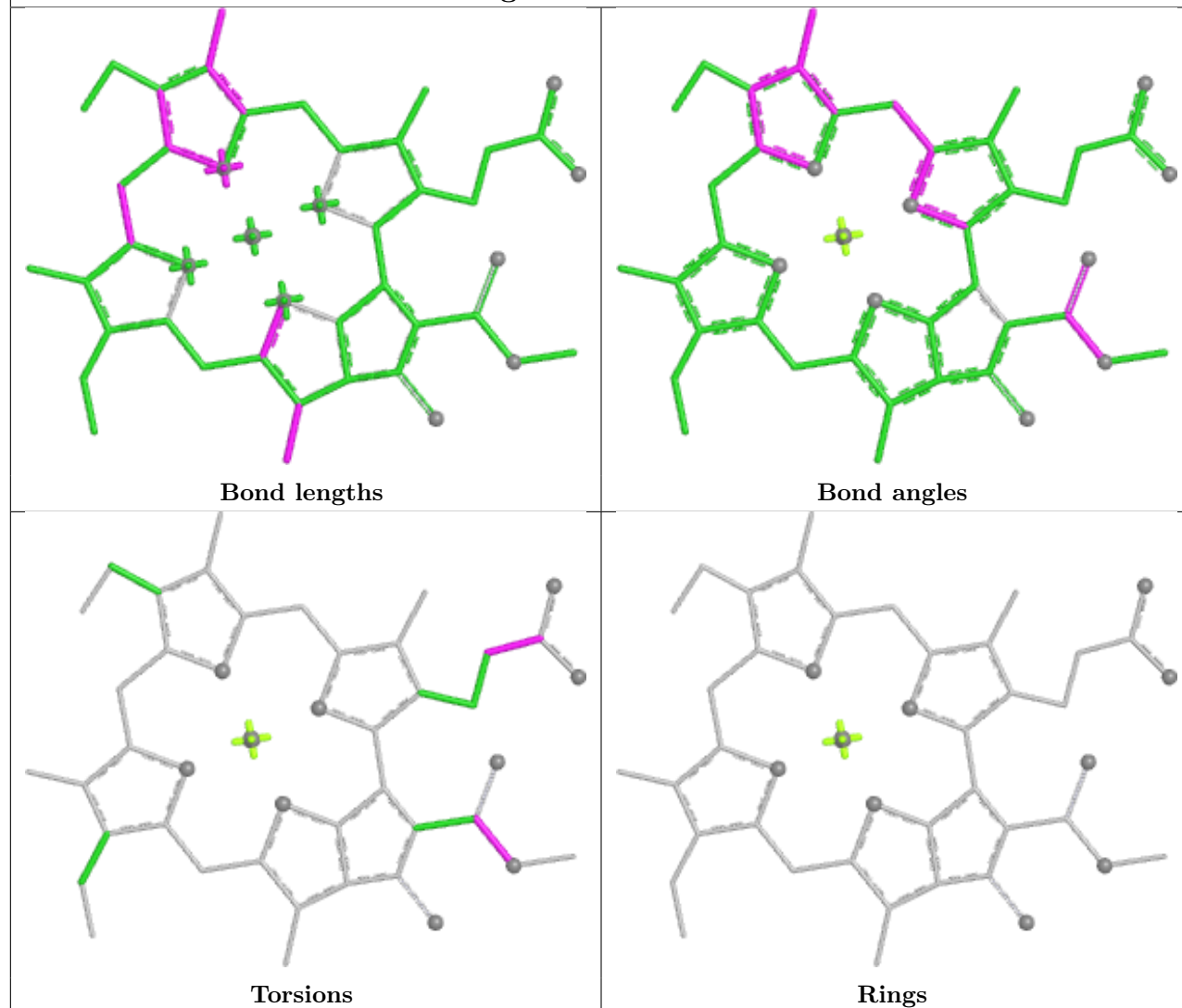
Ligand CLA a 842



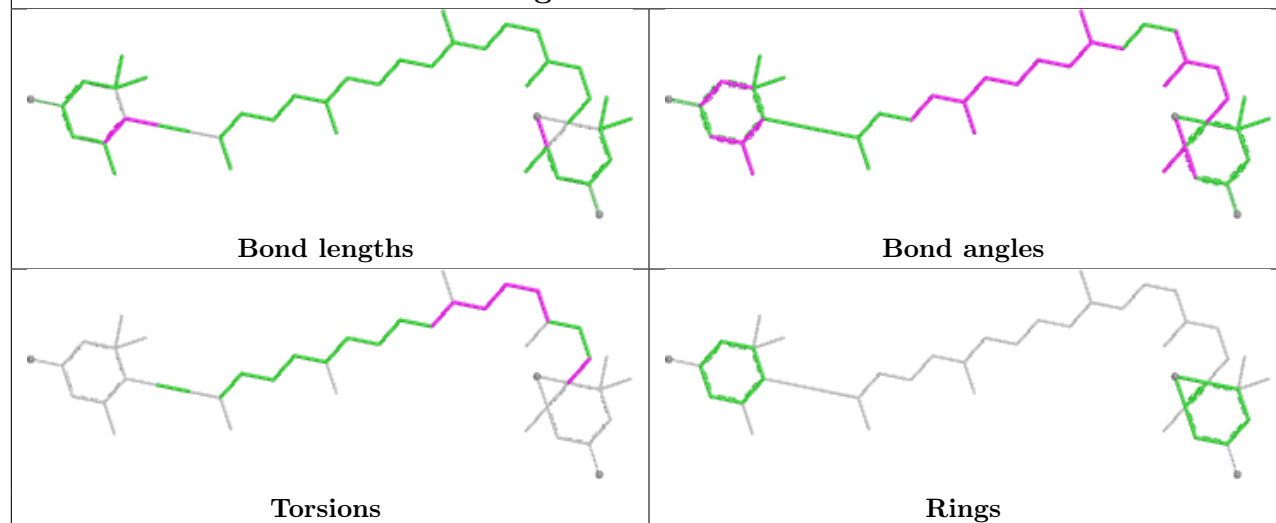
Ligand BCR a 850

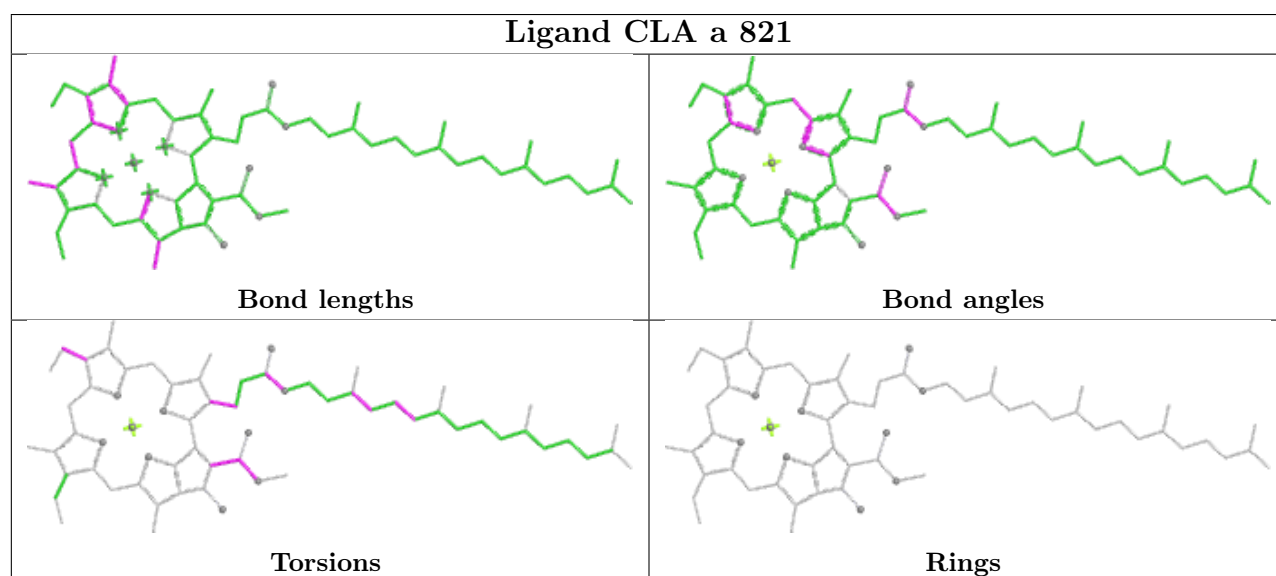
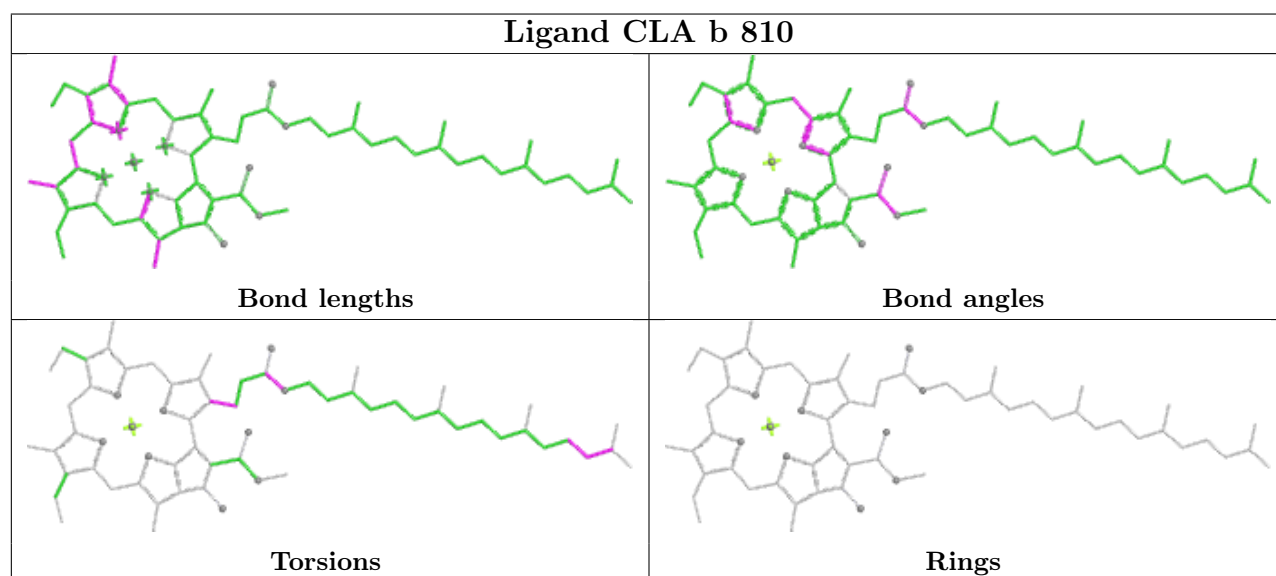
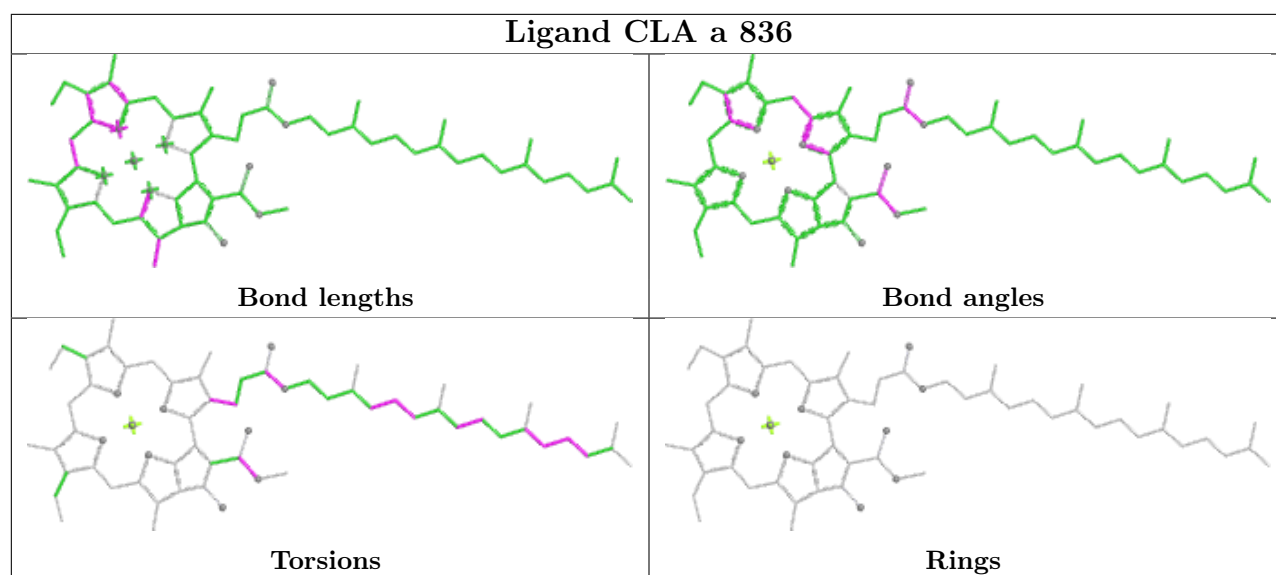


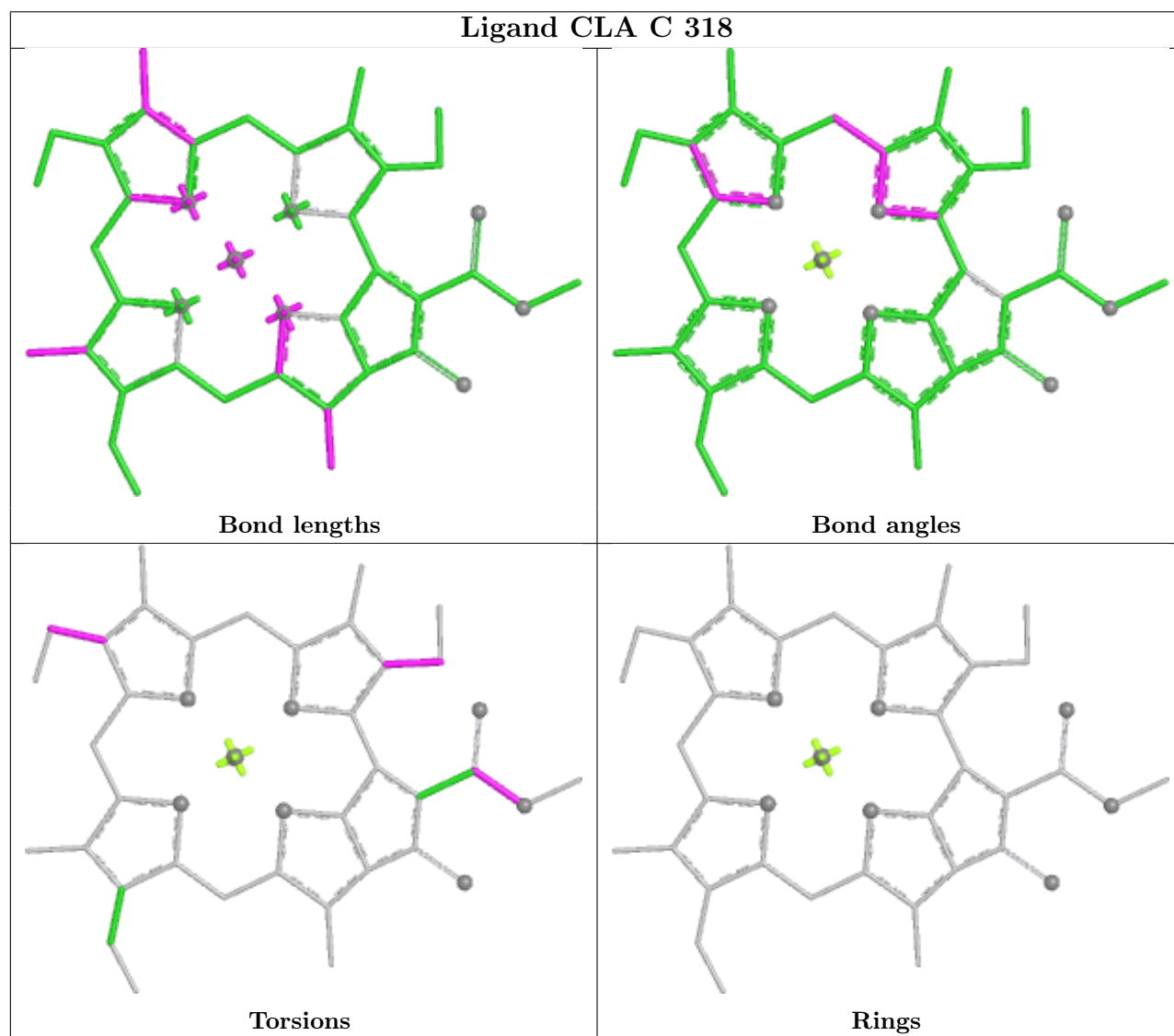
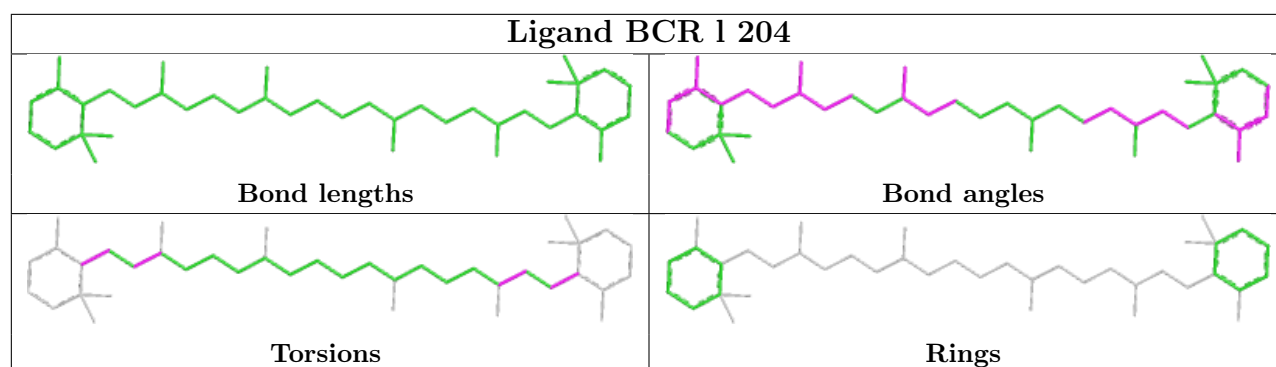
Ligand CLA 1 205

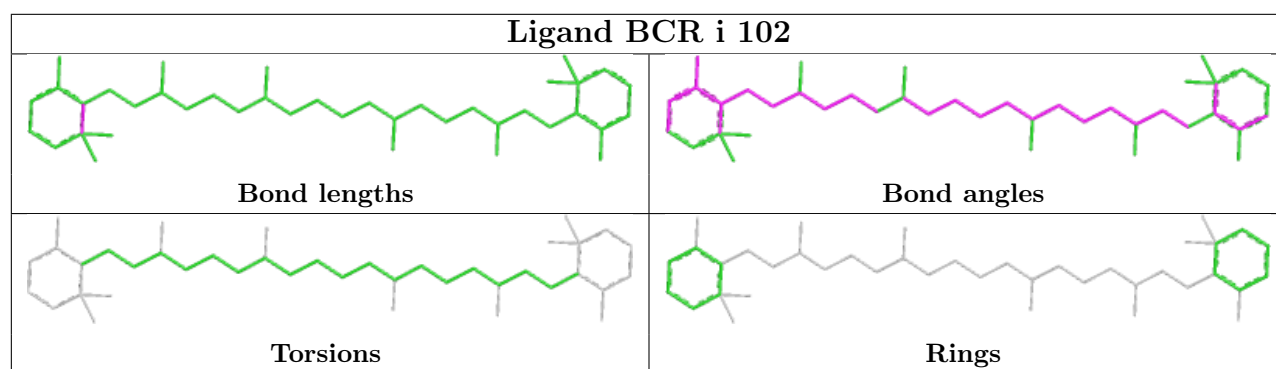
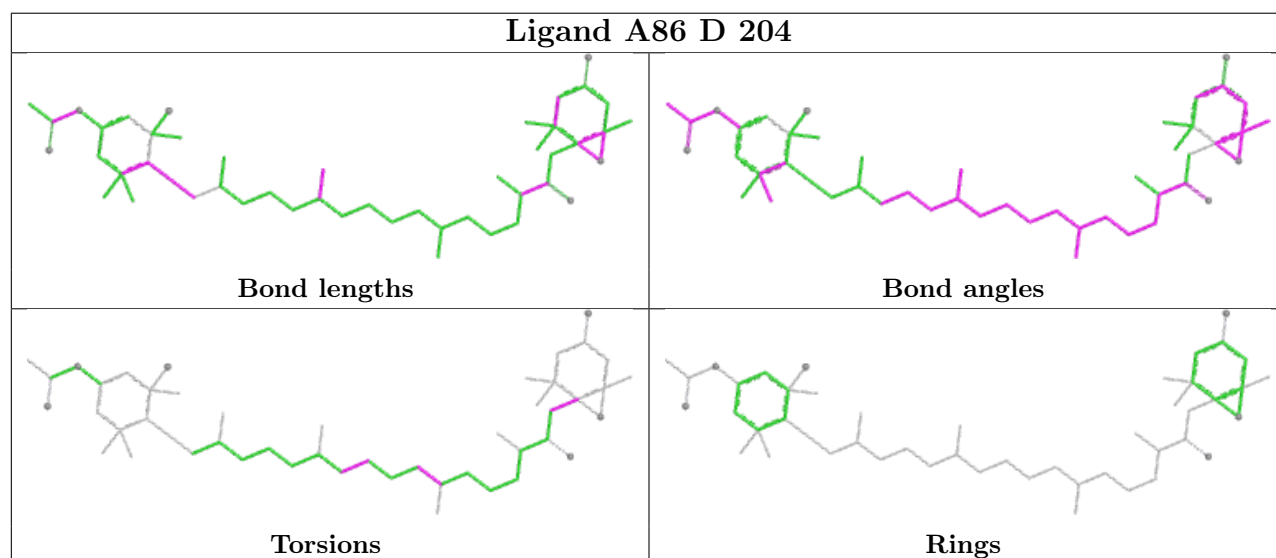
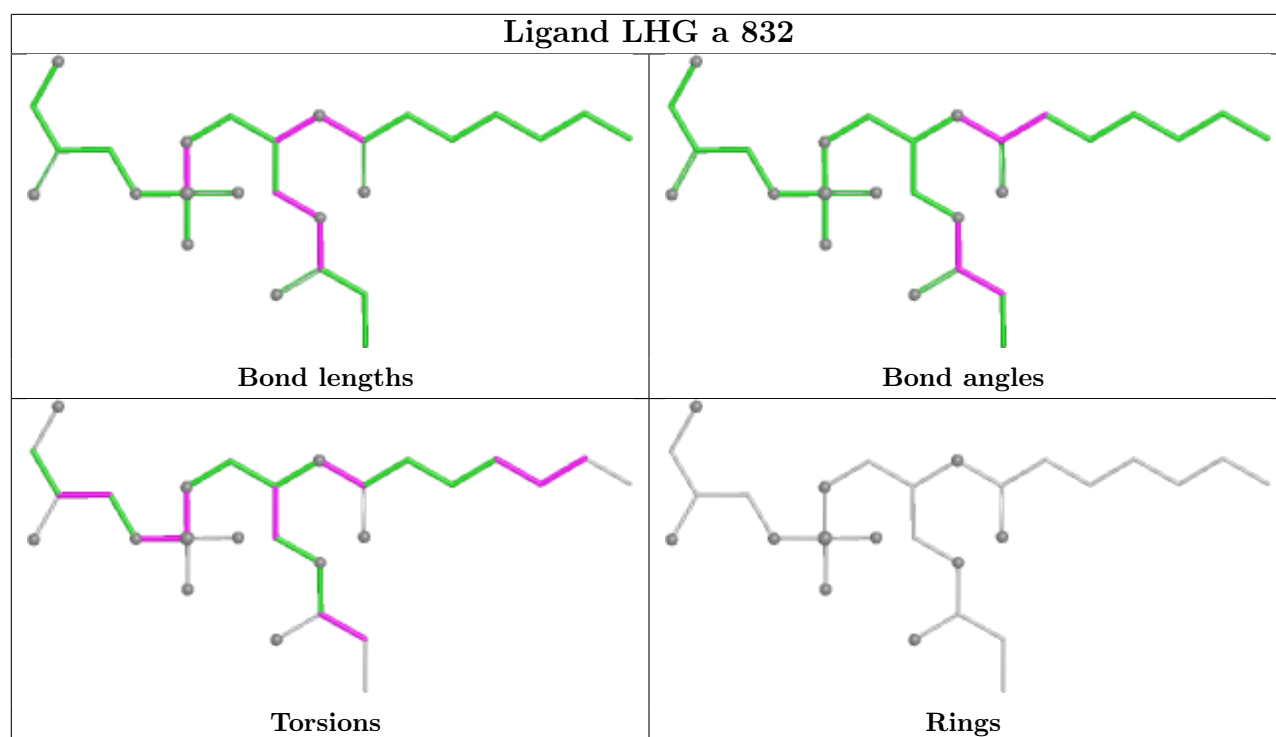


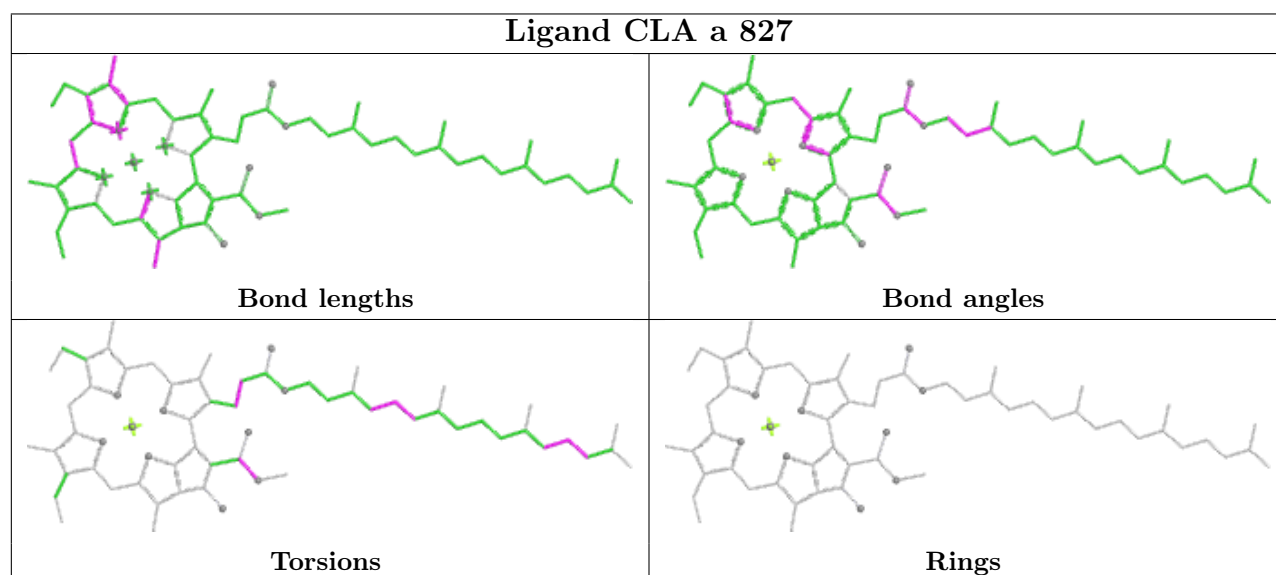
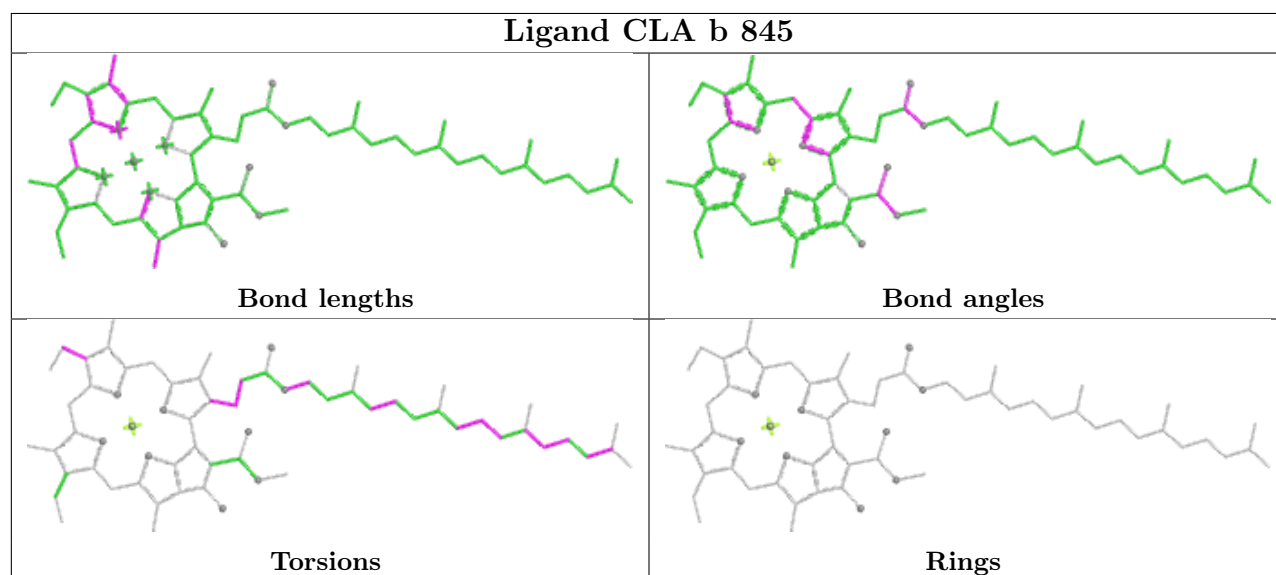
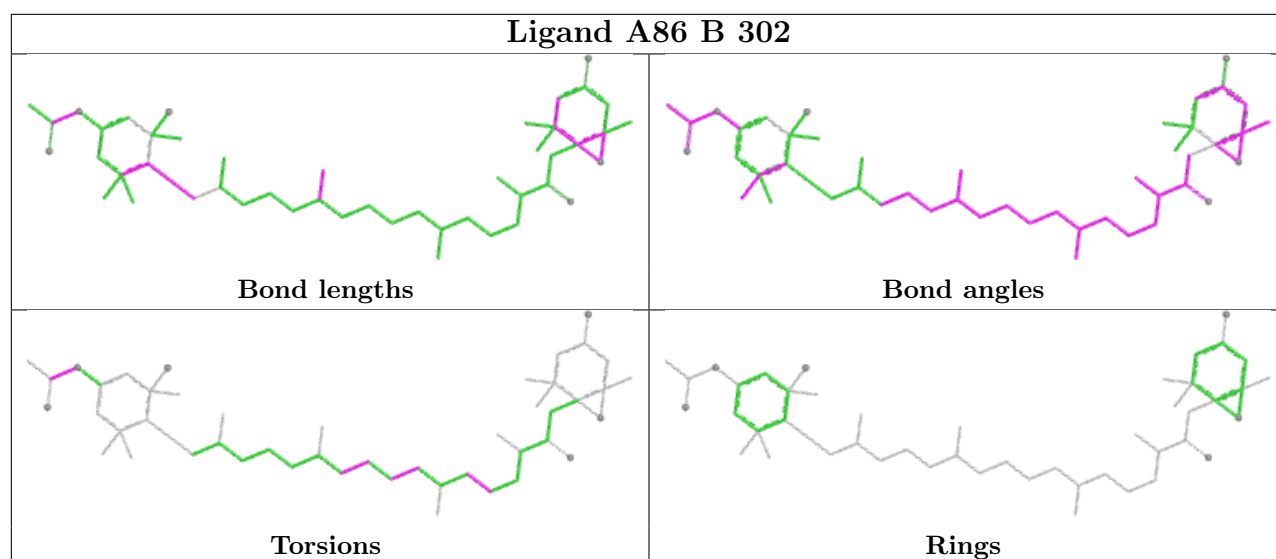
Ligand DD6 E 307

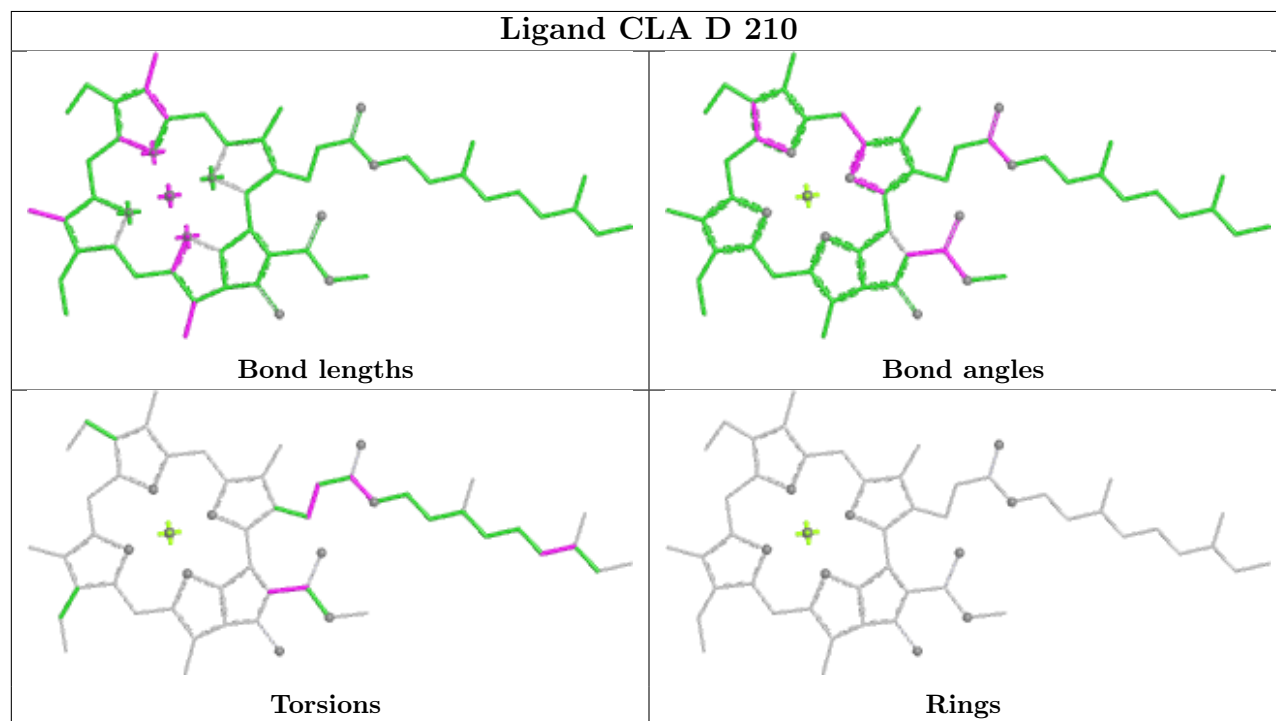
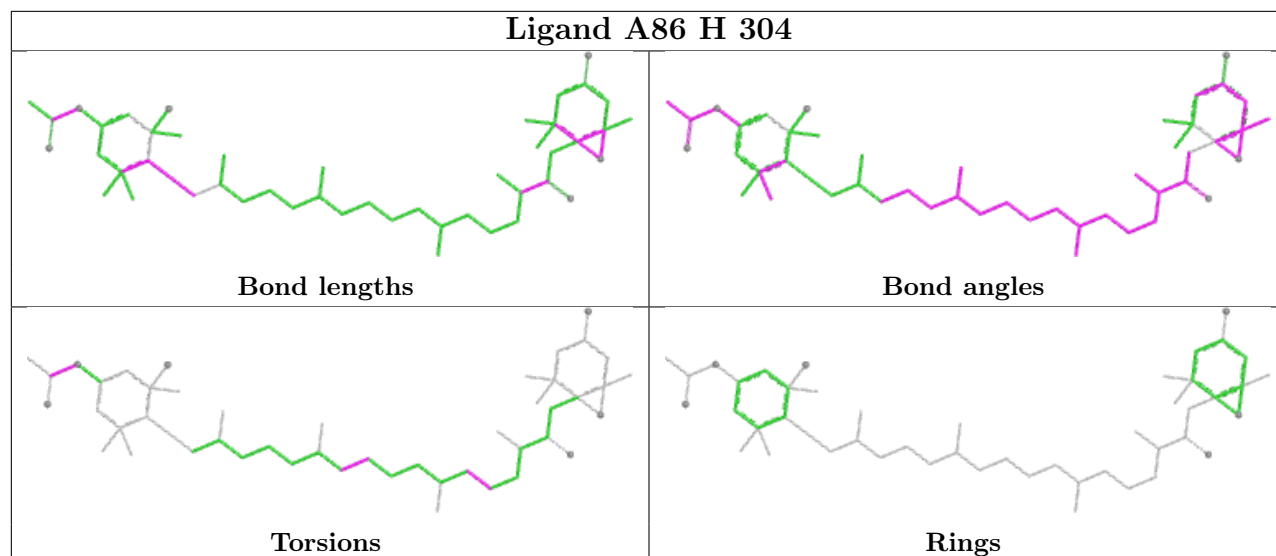




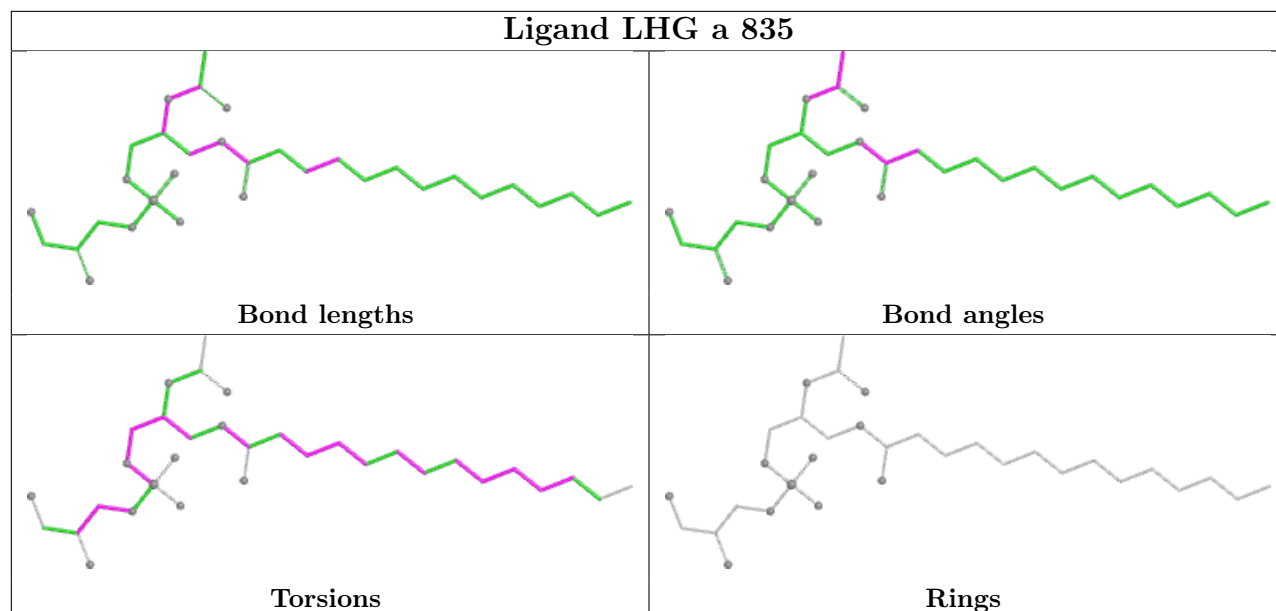




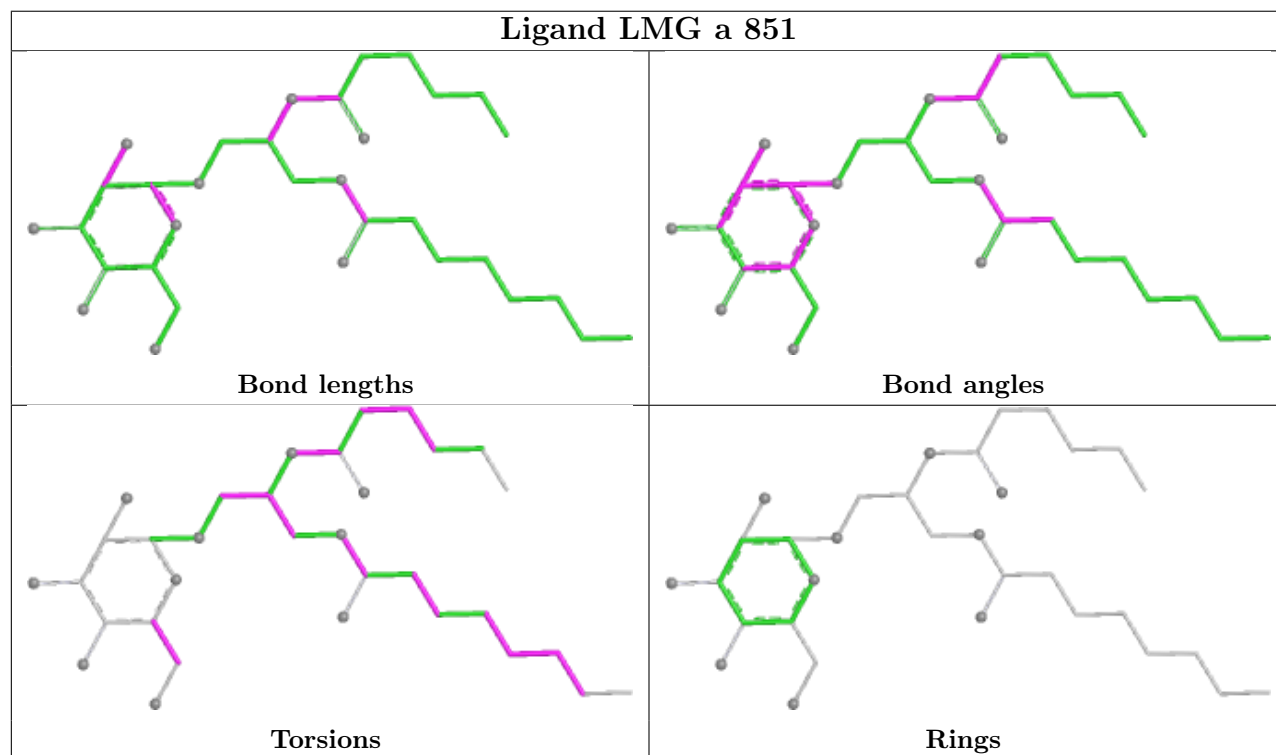




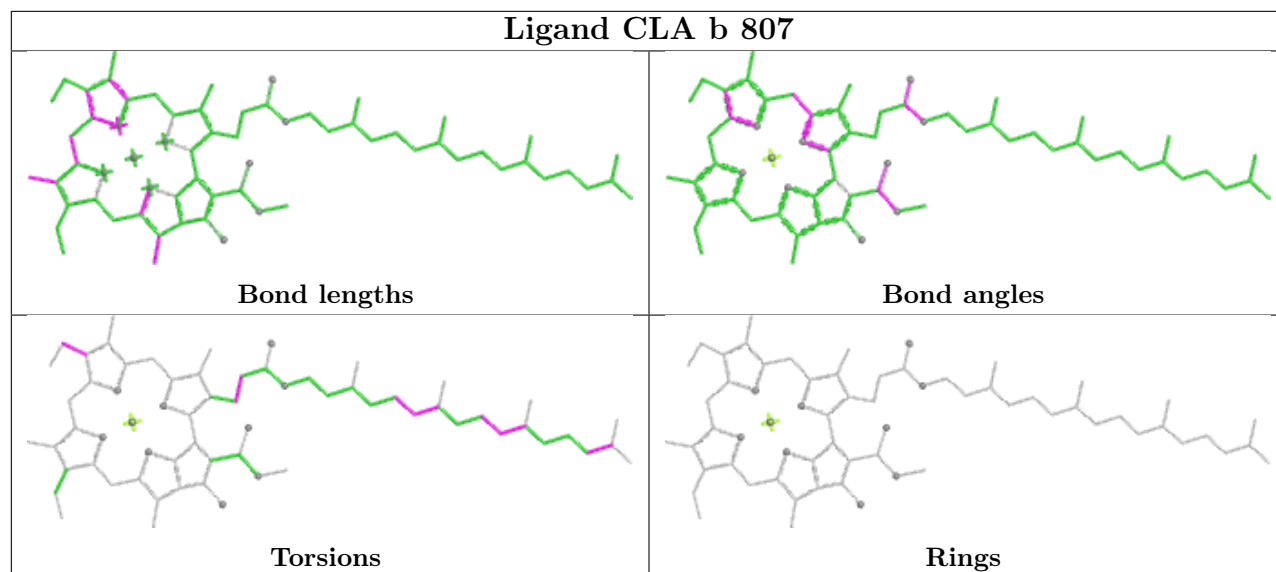
Ligand LHG a 835



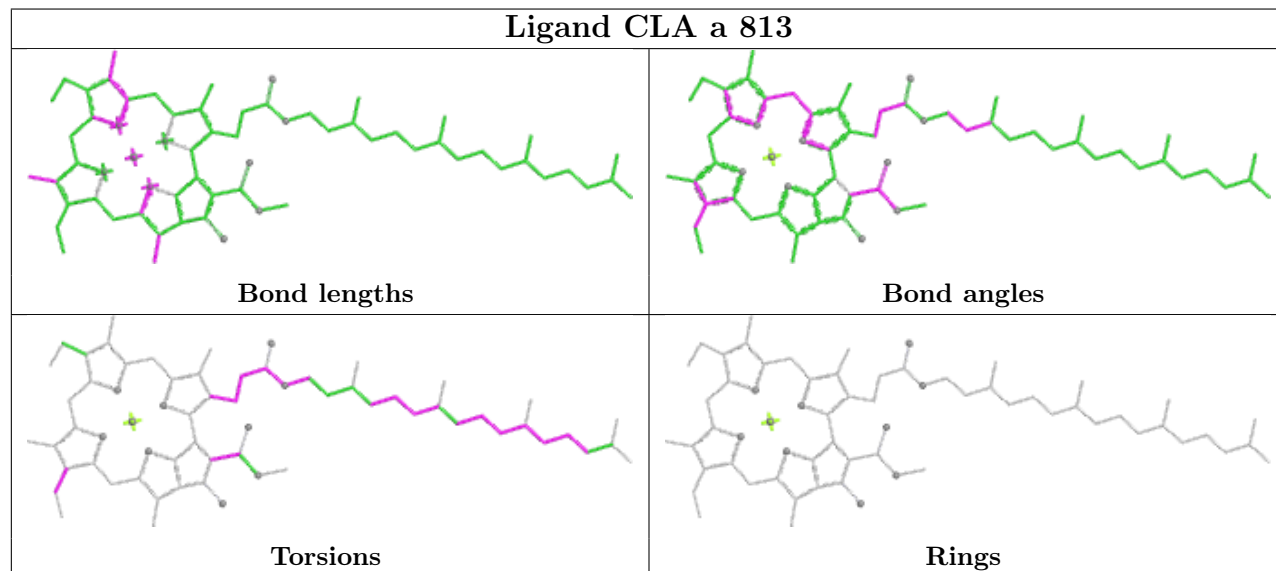
Ligand LMG a 851



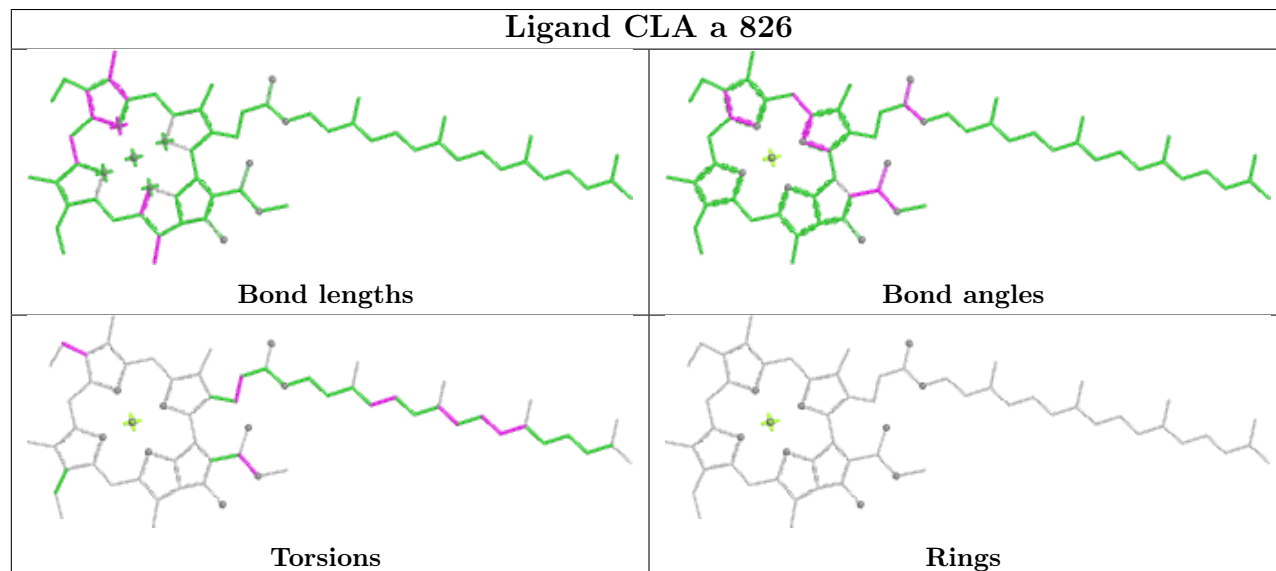
Ligand CLA b 807

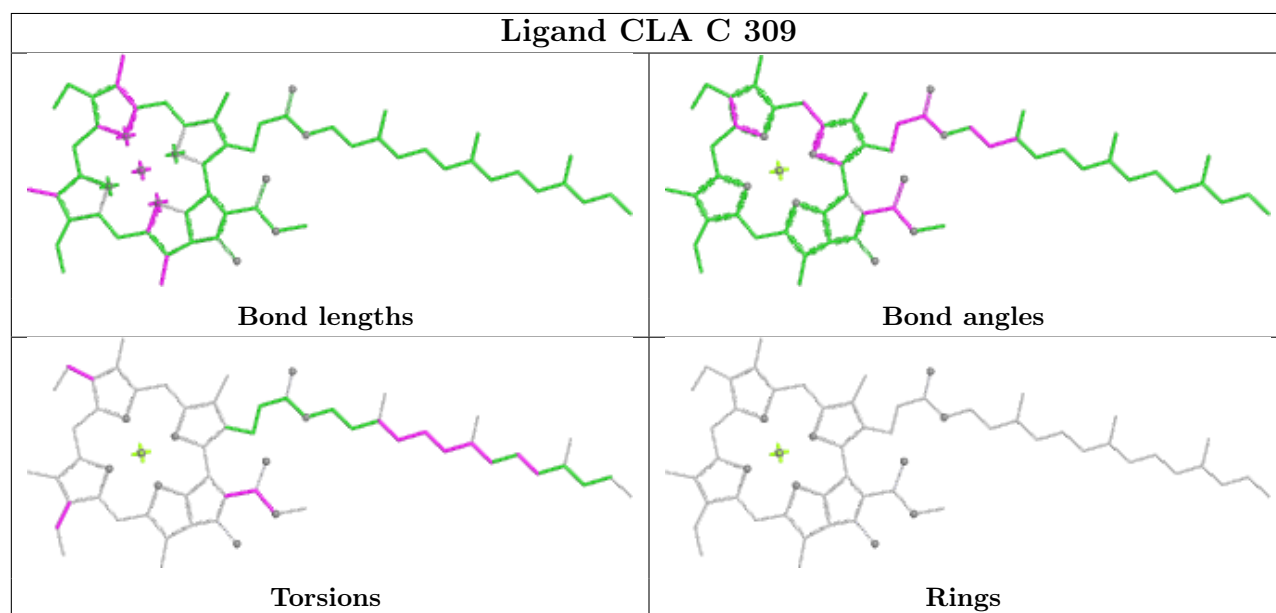
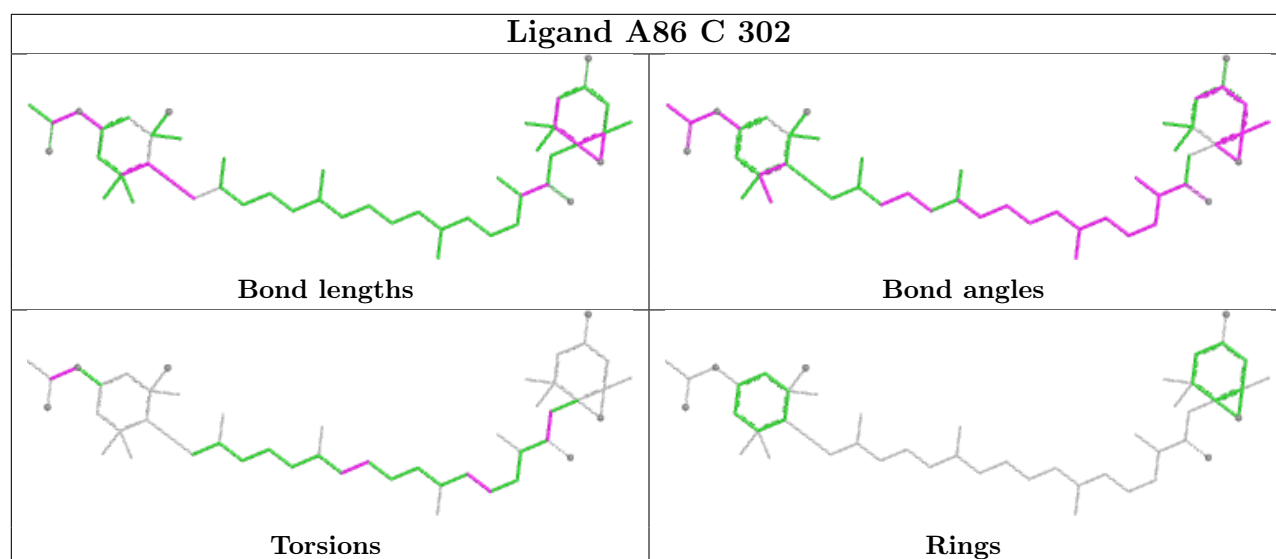


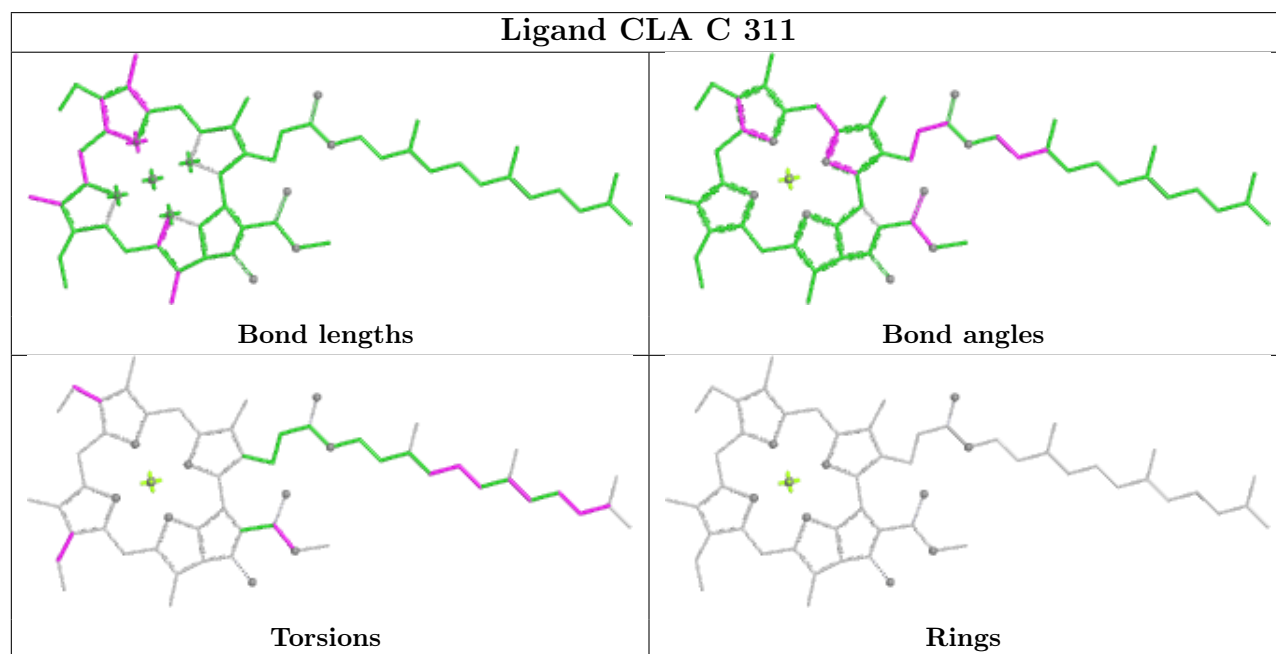
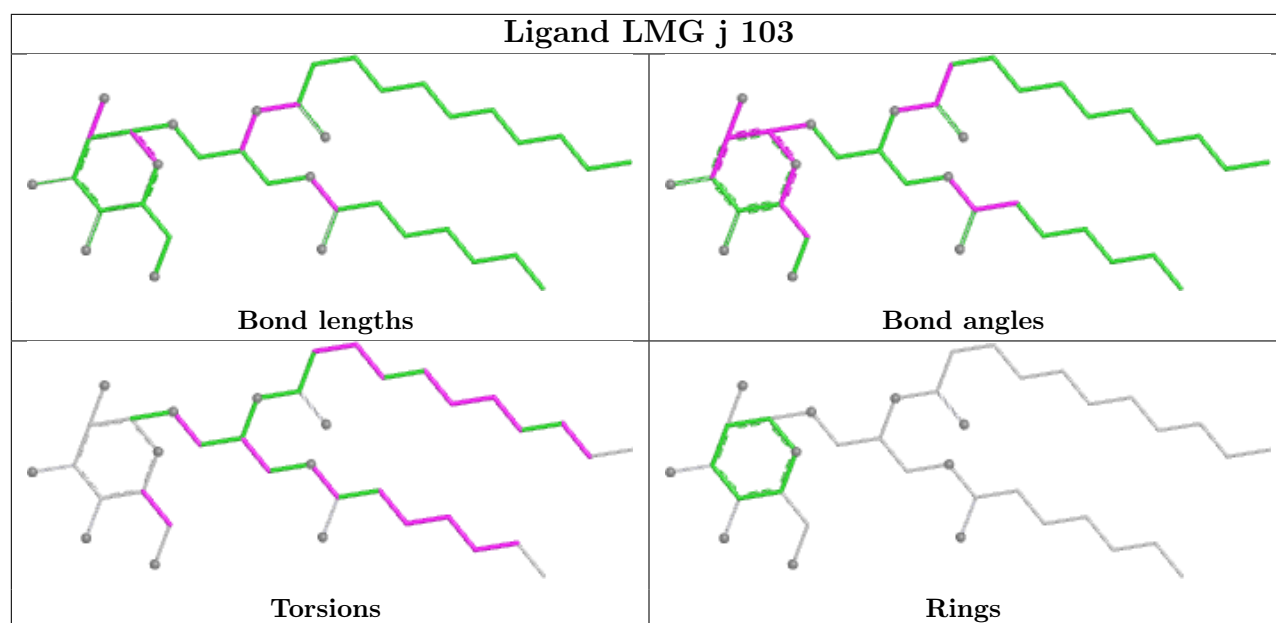
Ligand CLA a 813



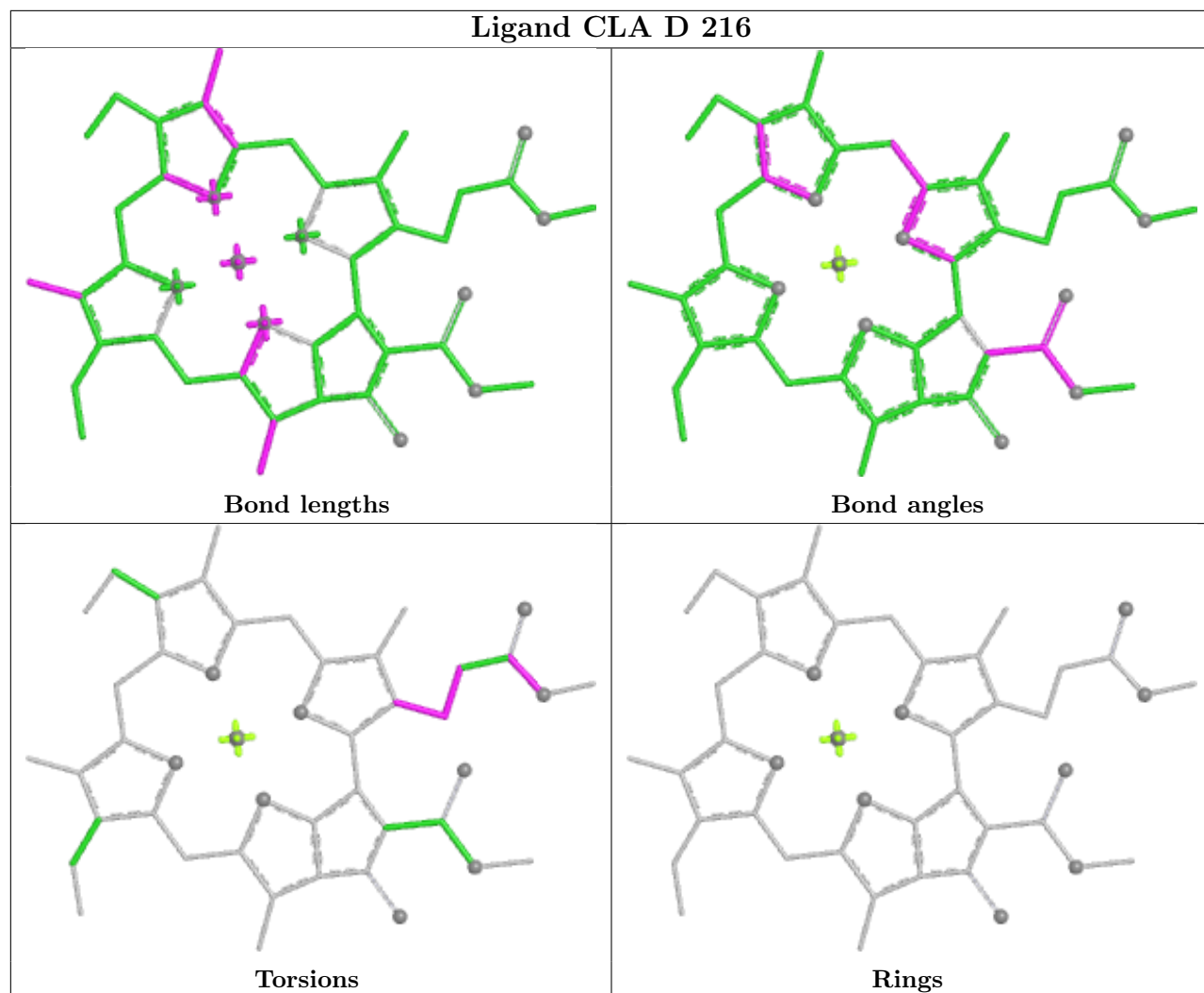
Ligand CLA a 826



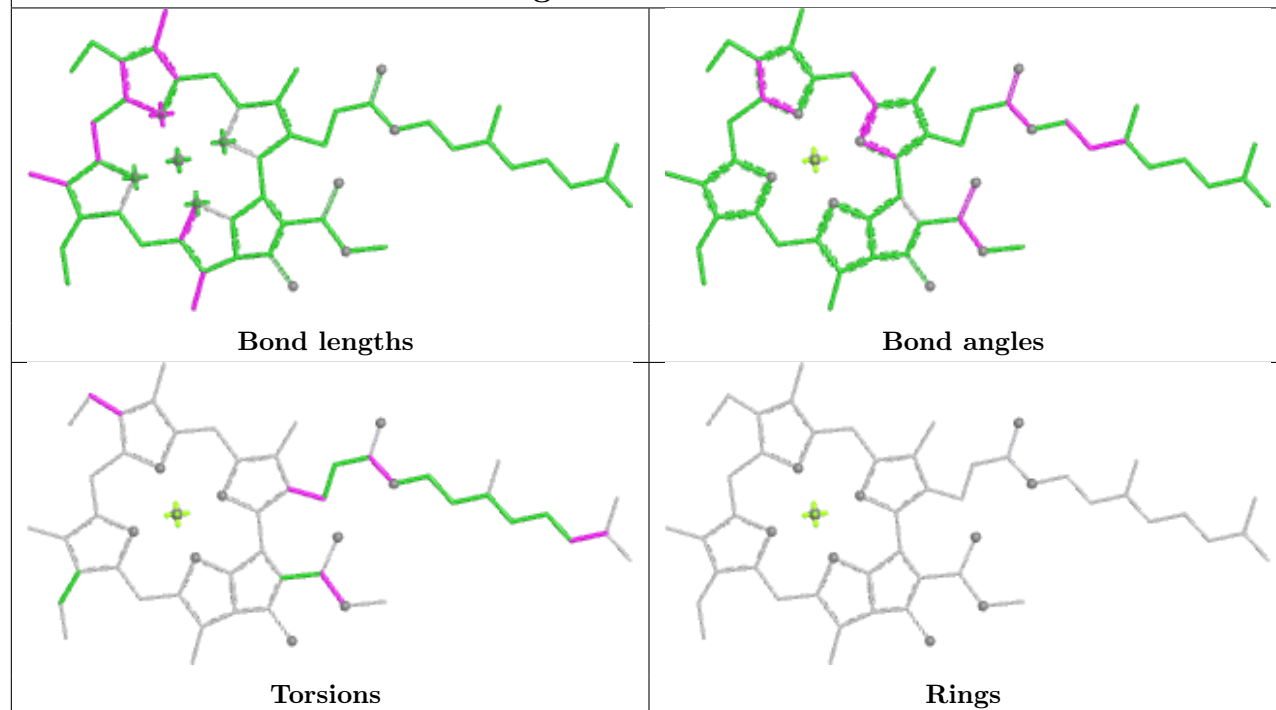




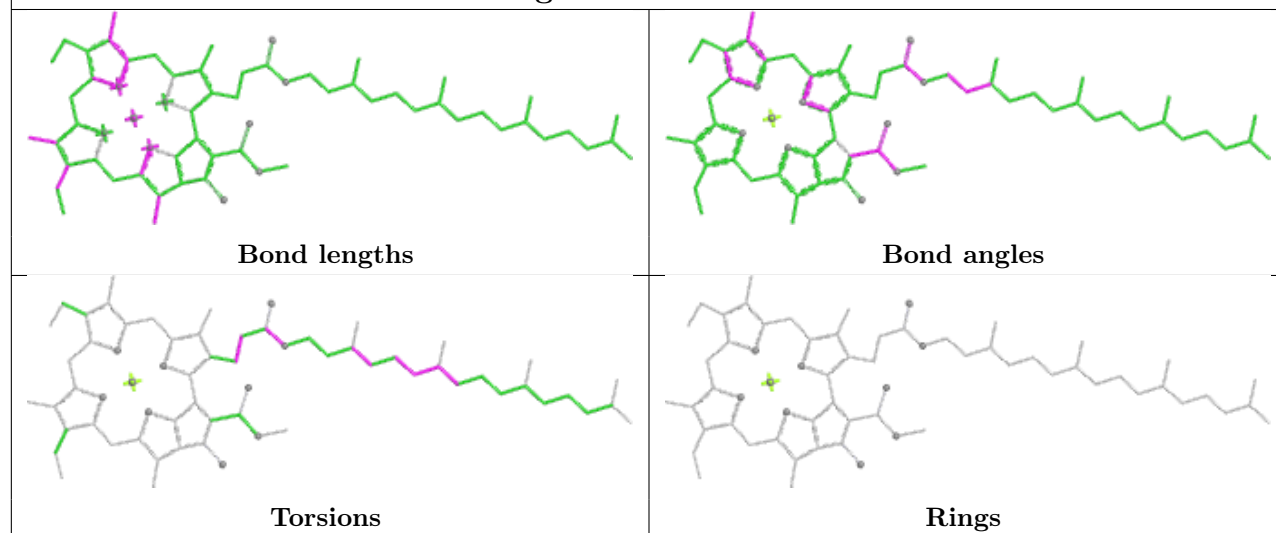
Ligand CLA D 216



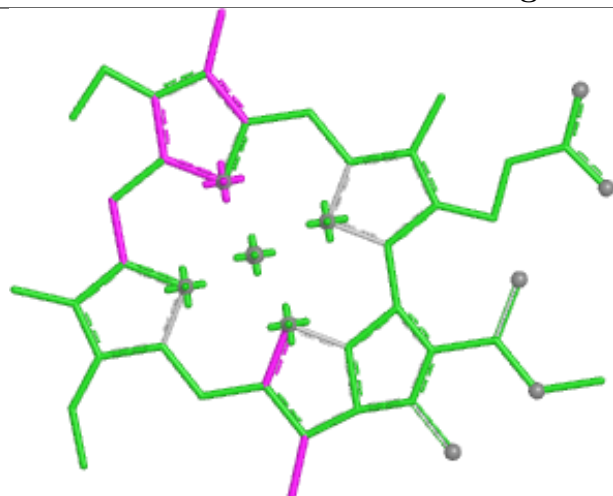
Ligand CLA E 313



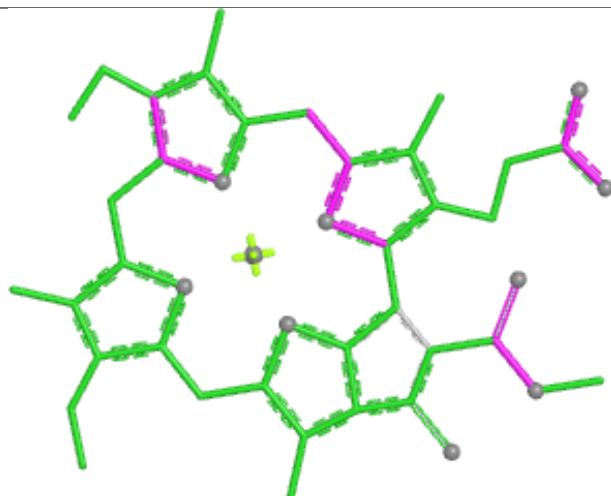
Ligand CLA b 801



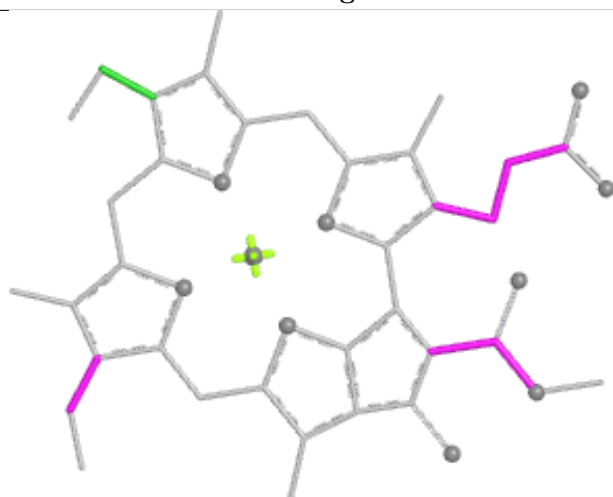
Ligand CLA r 202



Bond lengths



Bond angles

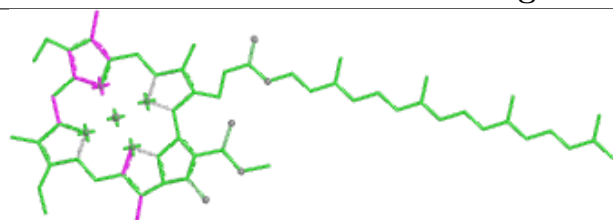


Torsions

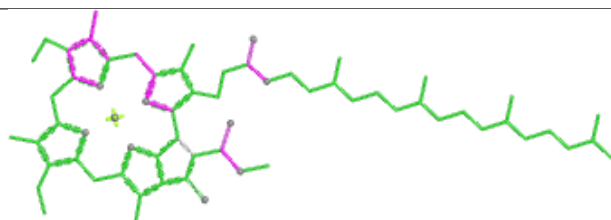


Rings

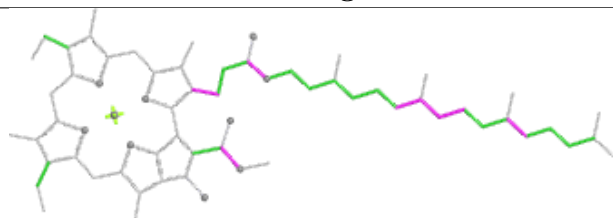
Ligand CLA a 855



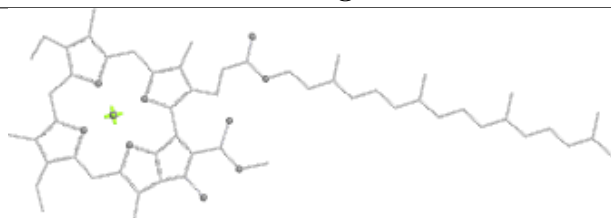
Bond lengths



Bond angles

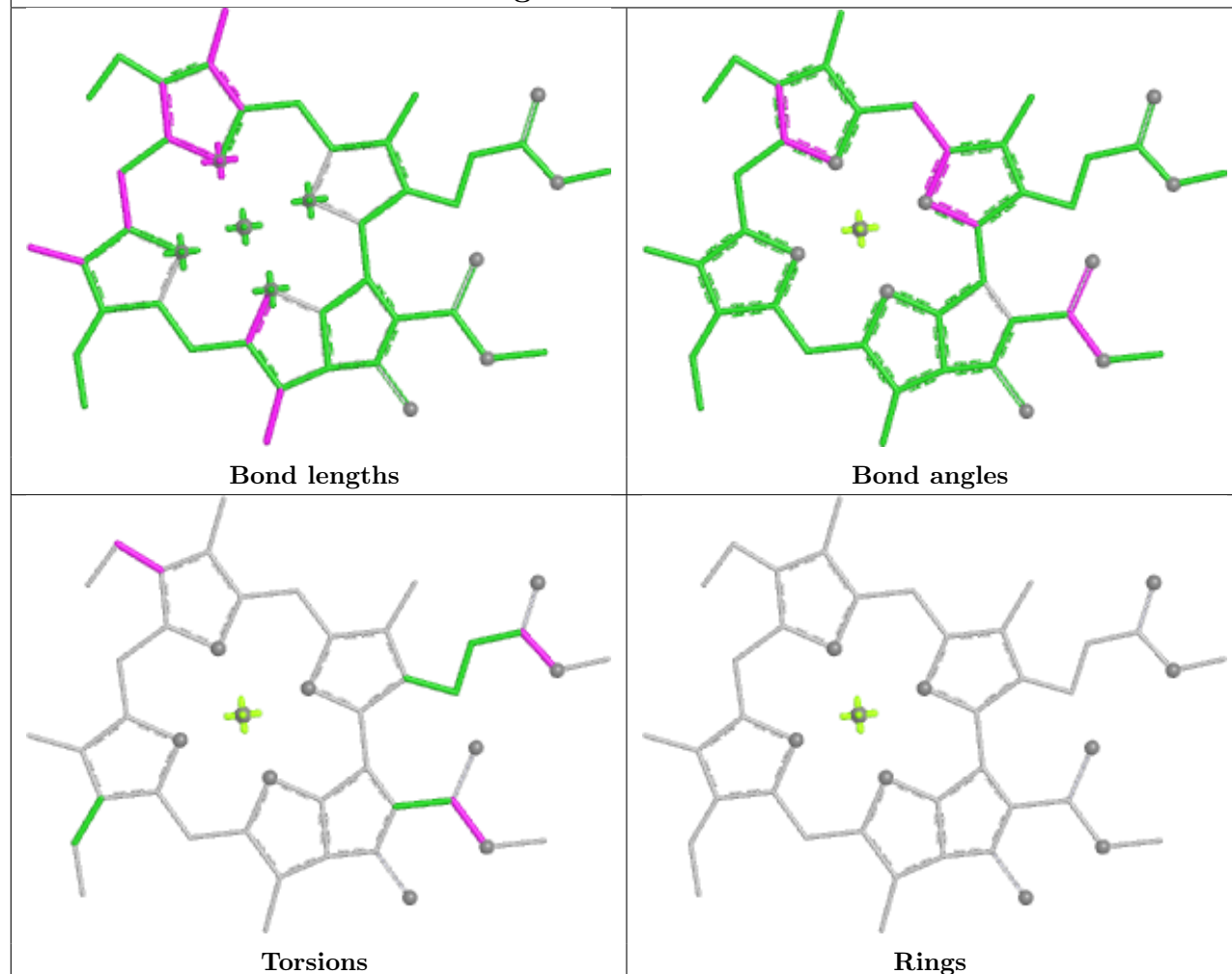


Torsions

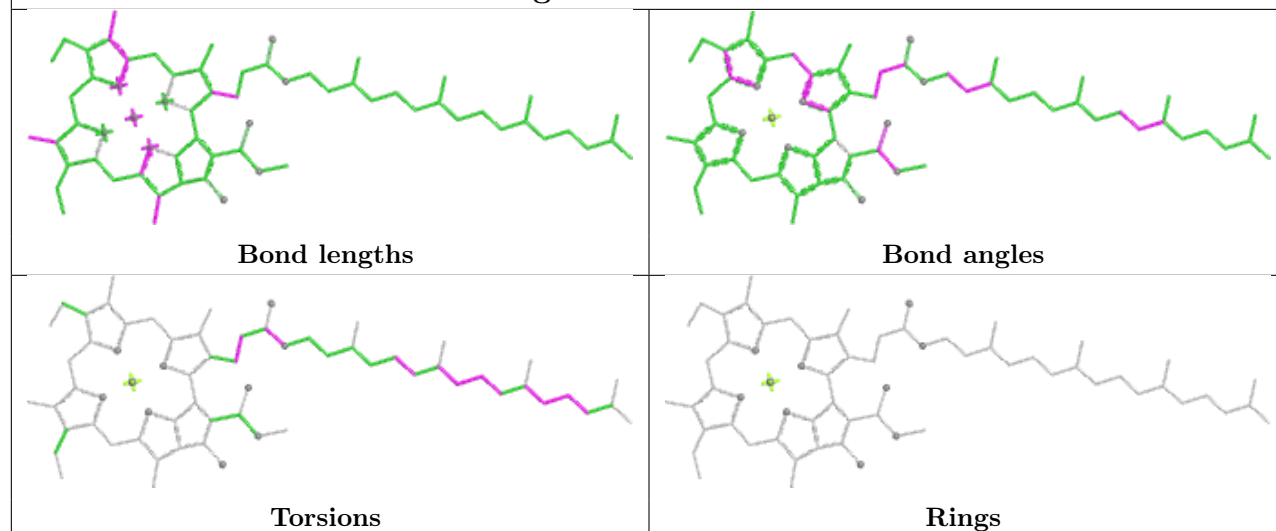


Rings

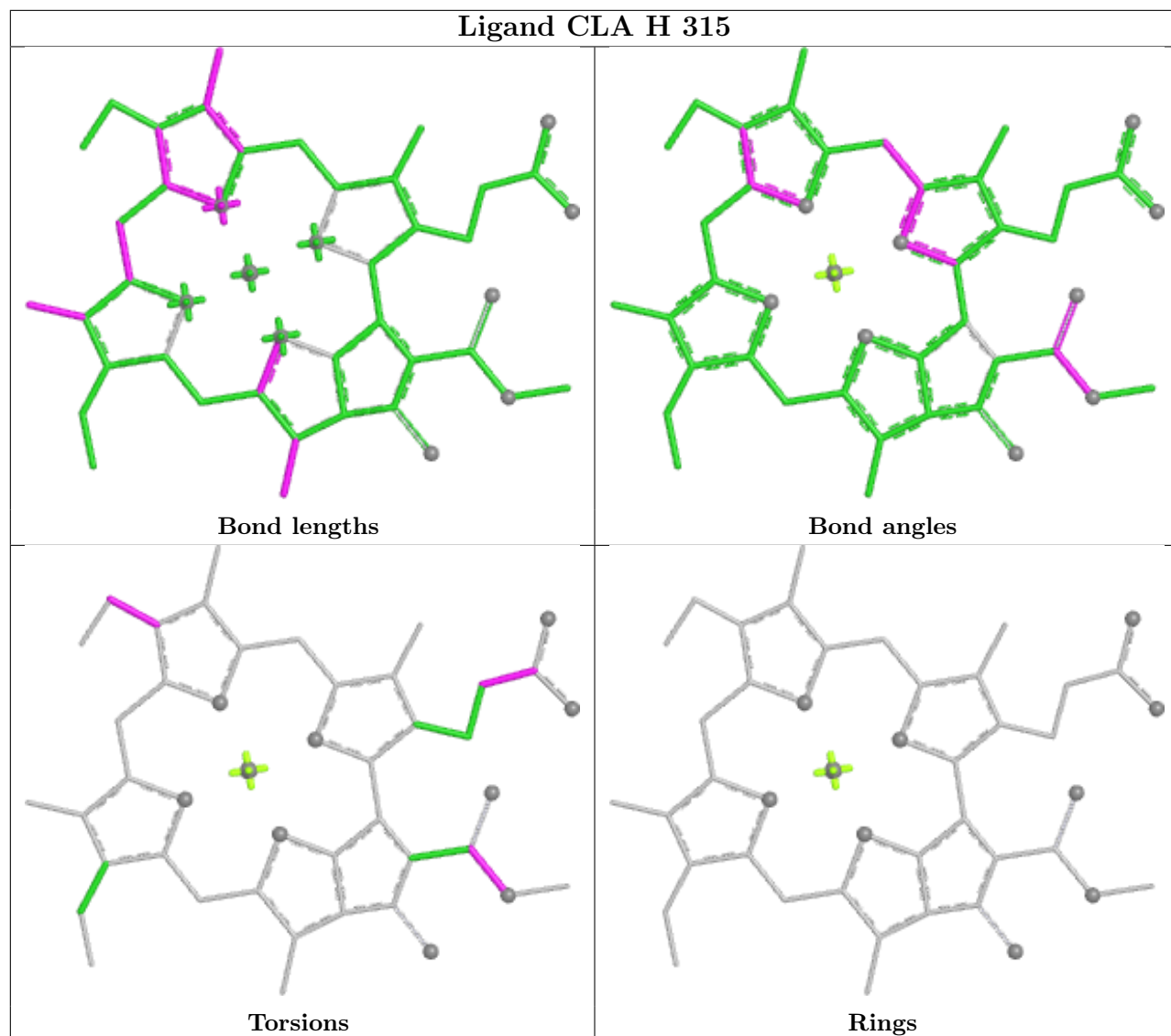
Ligand CLA B 310



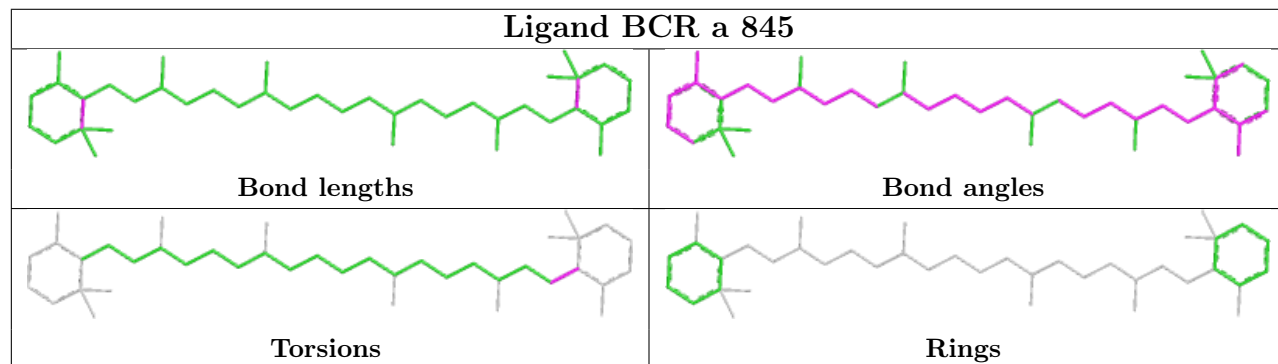
Ligand CLA E 312

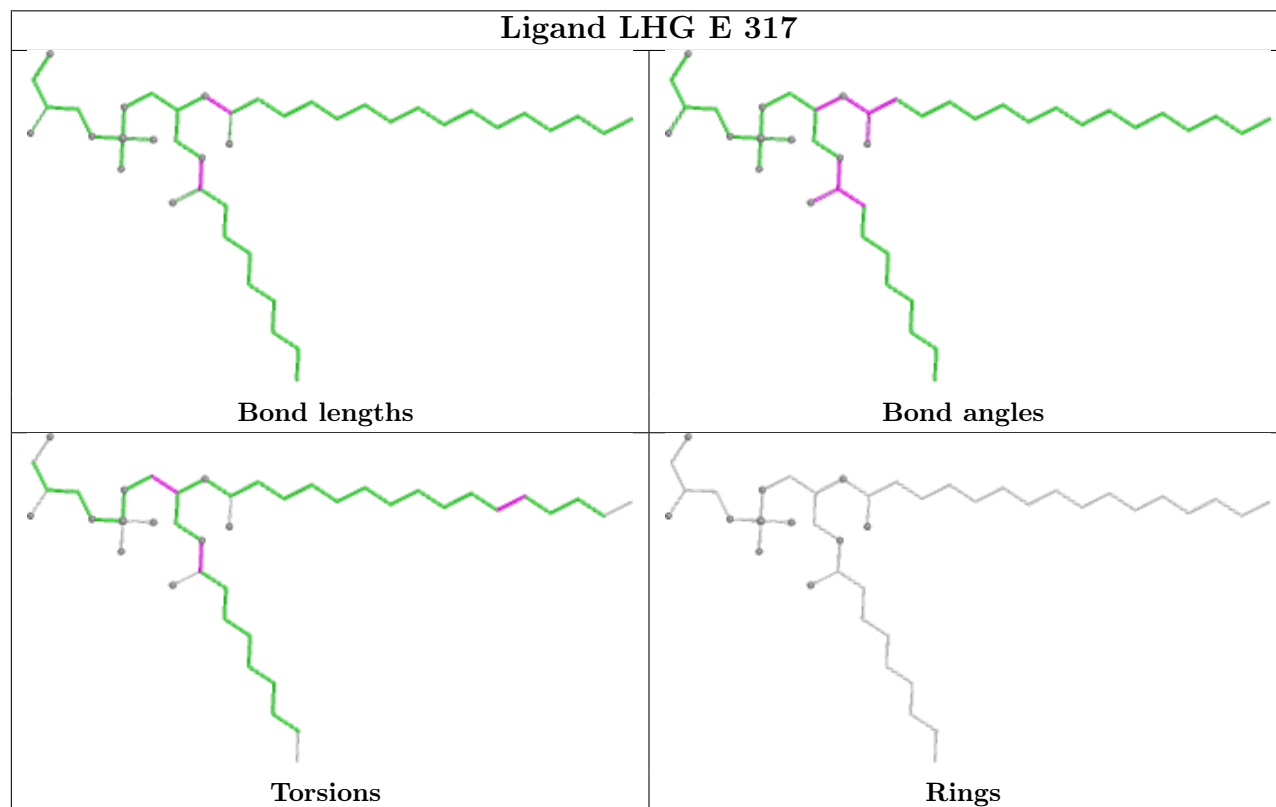
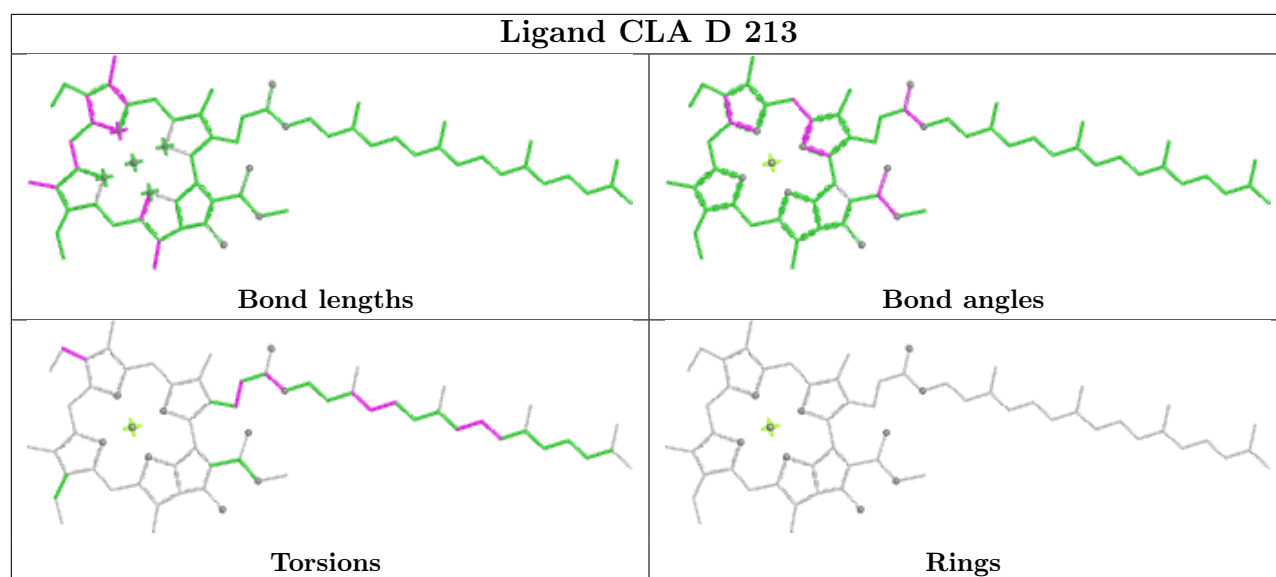


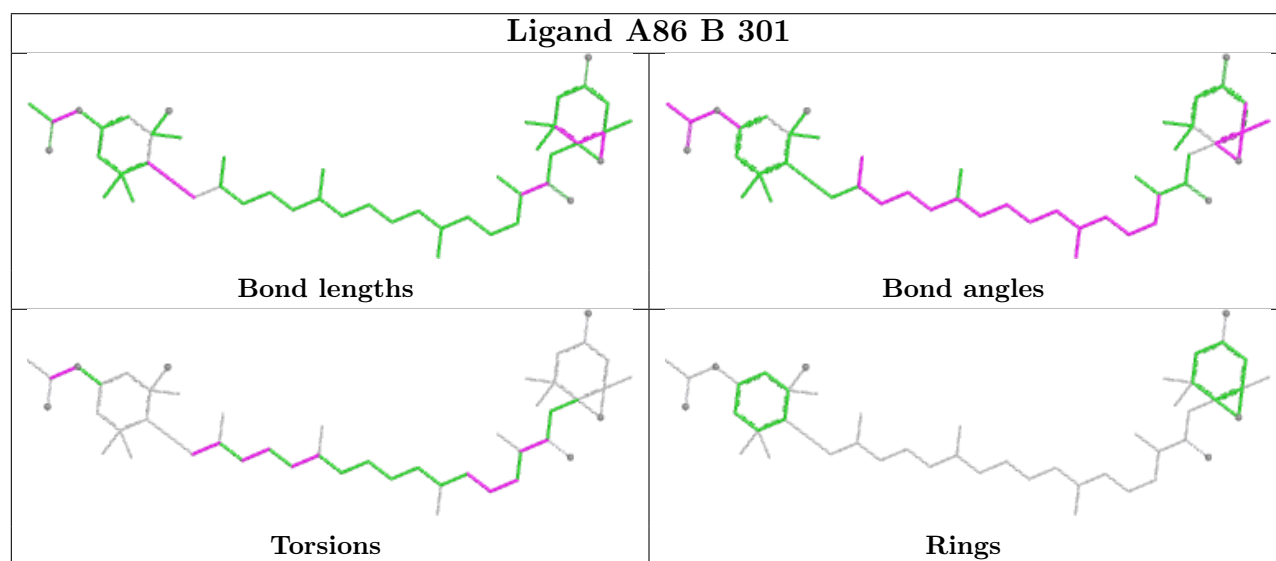
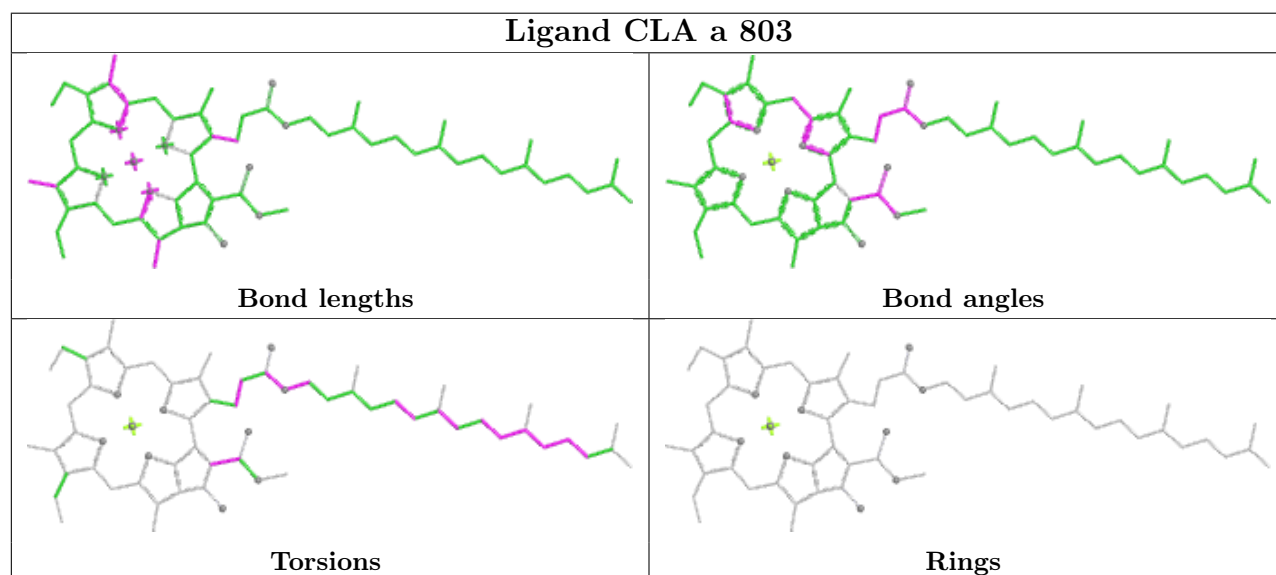
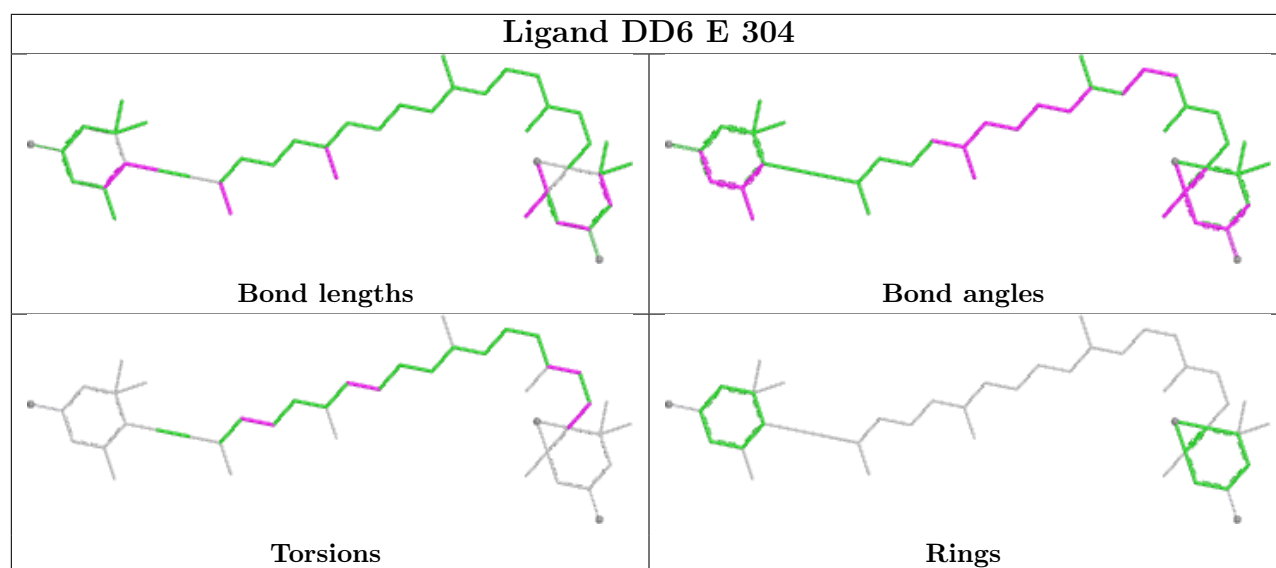
Ligand CLA H 315



Ligand BCR a 845







5.7 Other polymers [i](#)

There are no such residues in this entry.

5.8 Polymer linkage issues [i](#)

There are no chain breaks in this entry.

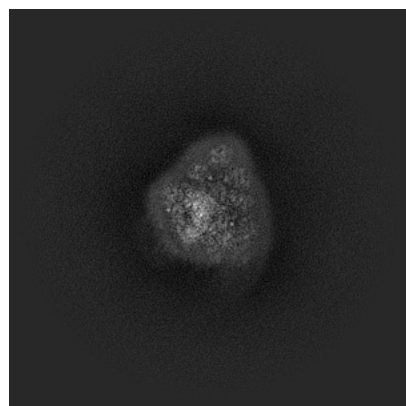
6 Map visualisation [i](#)

This section contains visualisations of the EMDB entry EMD-60044. These allow visual inspection of the internal detail of the map and identification of artifacts.

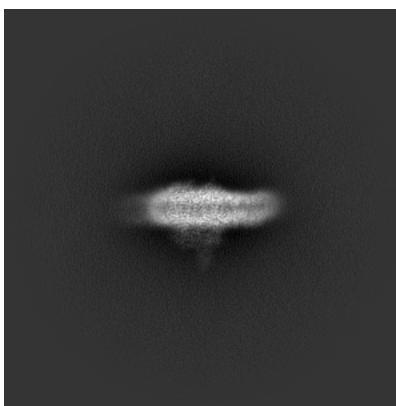
Images derived from a raw map, generated by summing the deposited half-maps, are presented below the corresponding image components of the primary map to allow further visual inspection and comparison with those of the primary map.

6.1 Orthogonal projections [i](#)

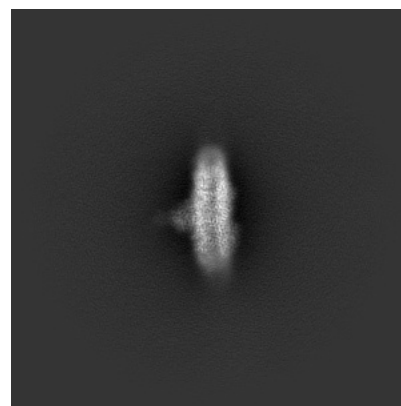
6.1.1 Primary map



X

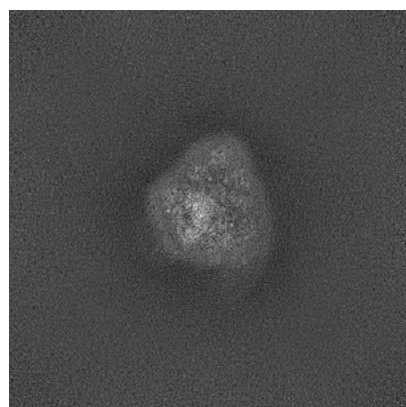


Y

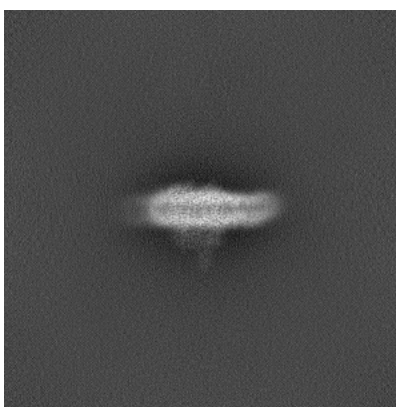


Z

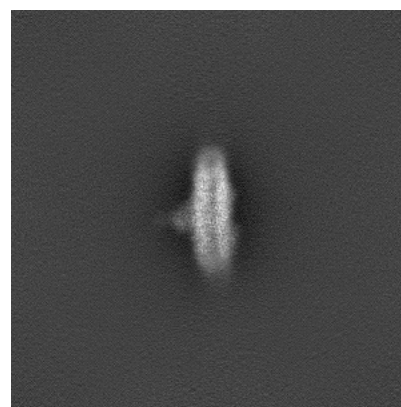
6.1.2 Raw map



X



Y

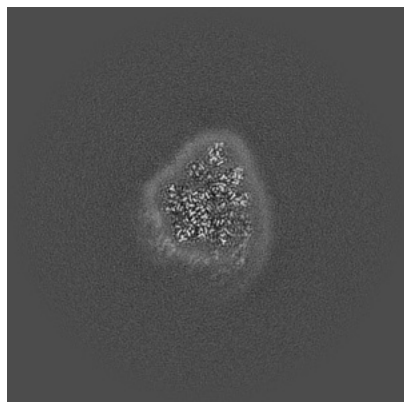


Z

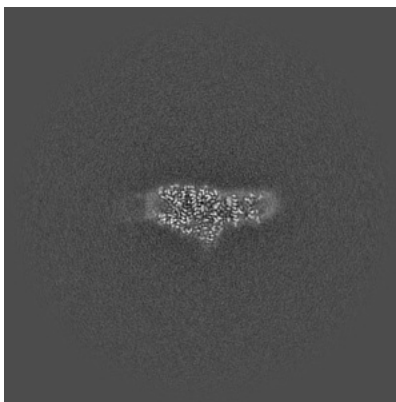
The images above show the map projected in three orthogonal directions.

6.2 Central slices [i](#)

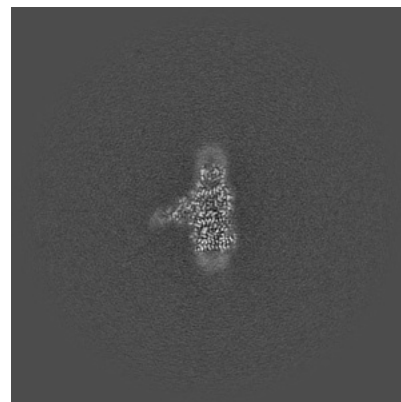
6.2.1 Primary map



X Index: 256

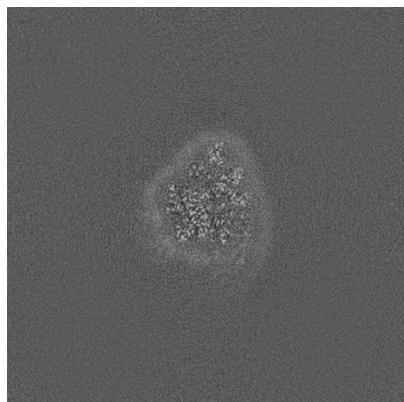


Y Index: 256

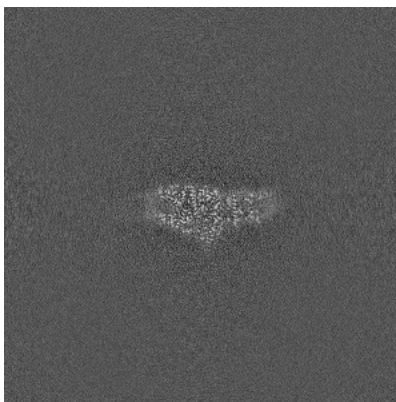


Z Index: 256

6.2.2 Raw map



X Index: 256



Y Index: 256

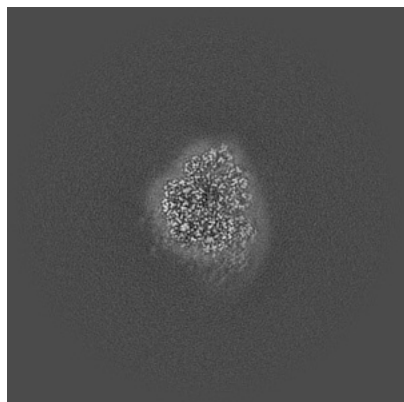


Z Index: 256

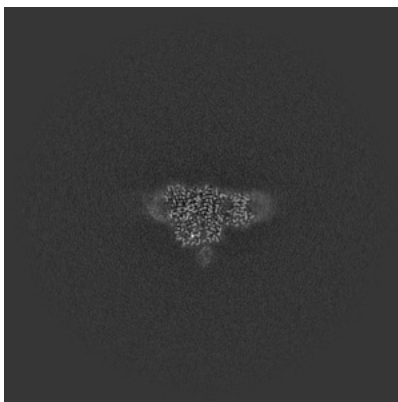
The images above show central slices of the map in three orthogonal directions.

6.3 Largest variance slices [i](#)

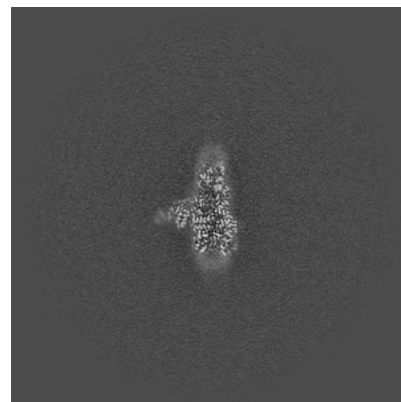
6.3.1 Primary map



X Index: 244

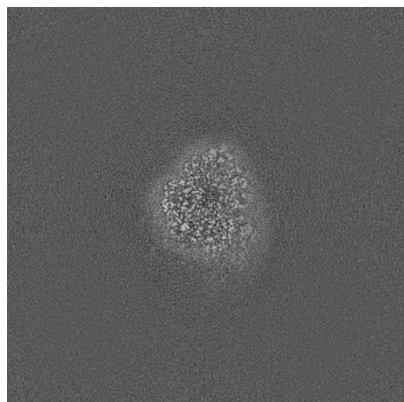


Y Index: 240

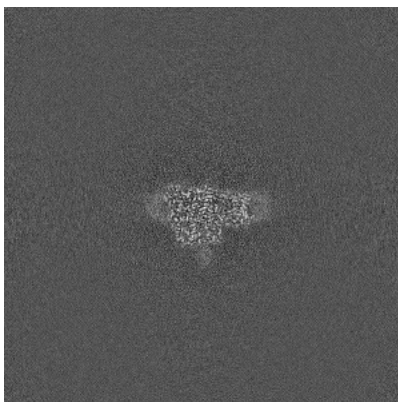


Z Index: 262

6.3.2 Raw map



X Index: 244



Y Index: 239

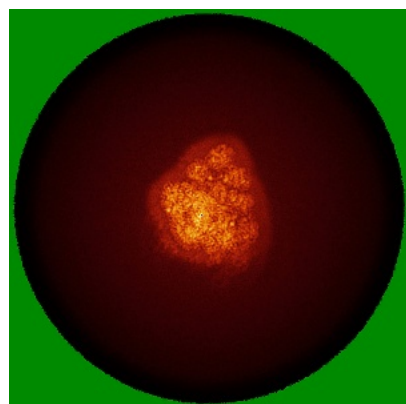


Z Index: 262

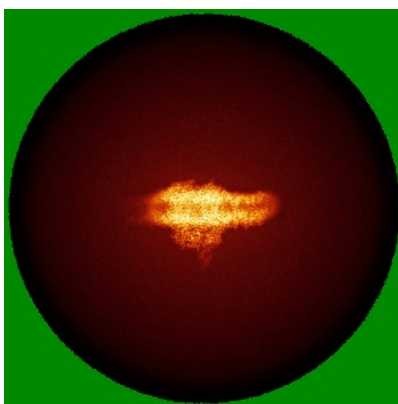
The images above show the largest variance slices of the map in three orthogonal directions.

6.4 Orthogonal standard-deviation projections (False-color) [i](#)

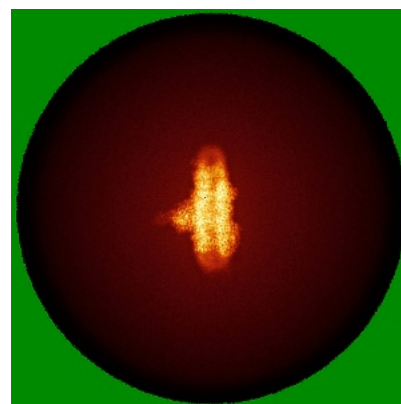
6.4.1 Primary map



X

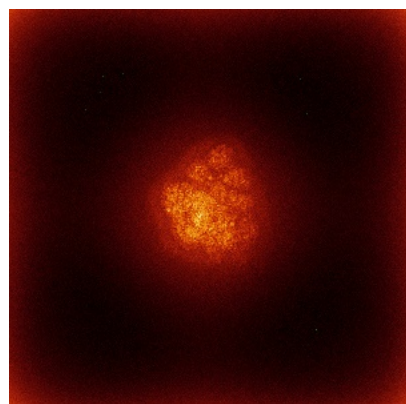


Y

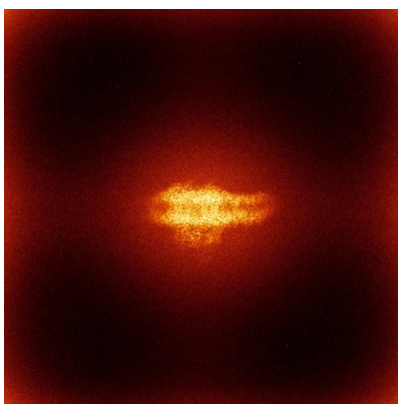


Z

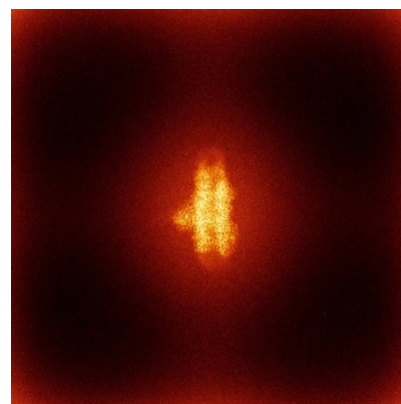
6.4.2 Raw map



X



Y

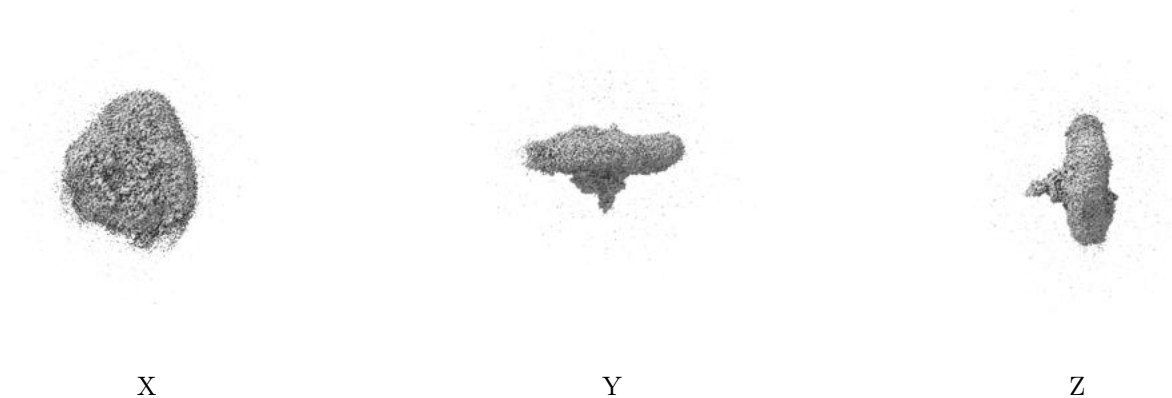


Z

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.

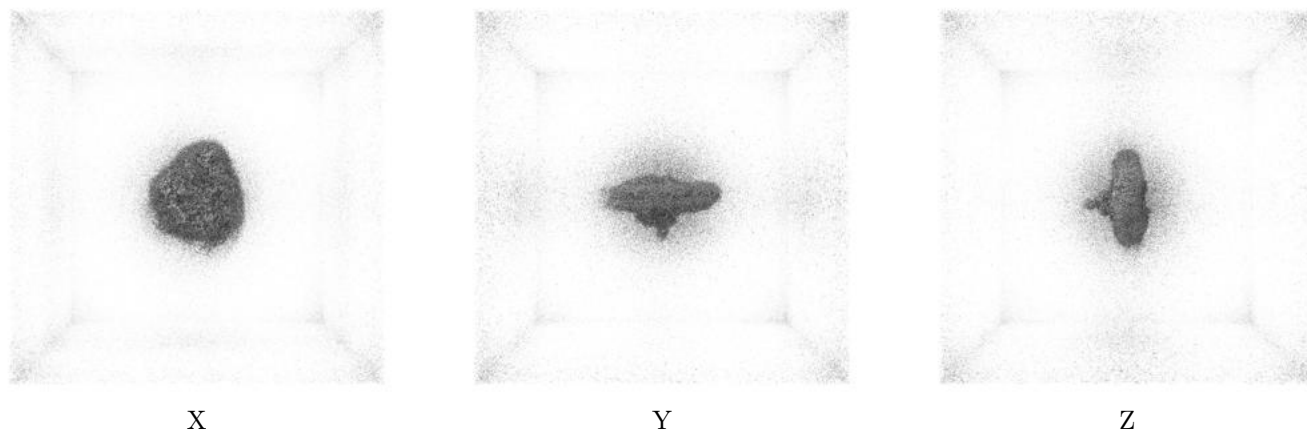
6.5 Orthogonal surface views [i](#)

6.5.1 Primary map



The images above show the 3D surface view of the map at the recommended contour level 0.08. 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



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.

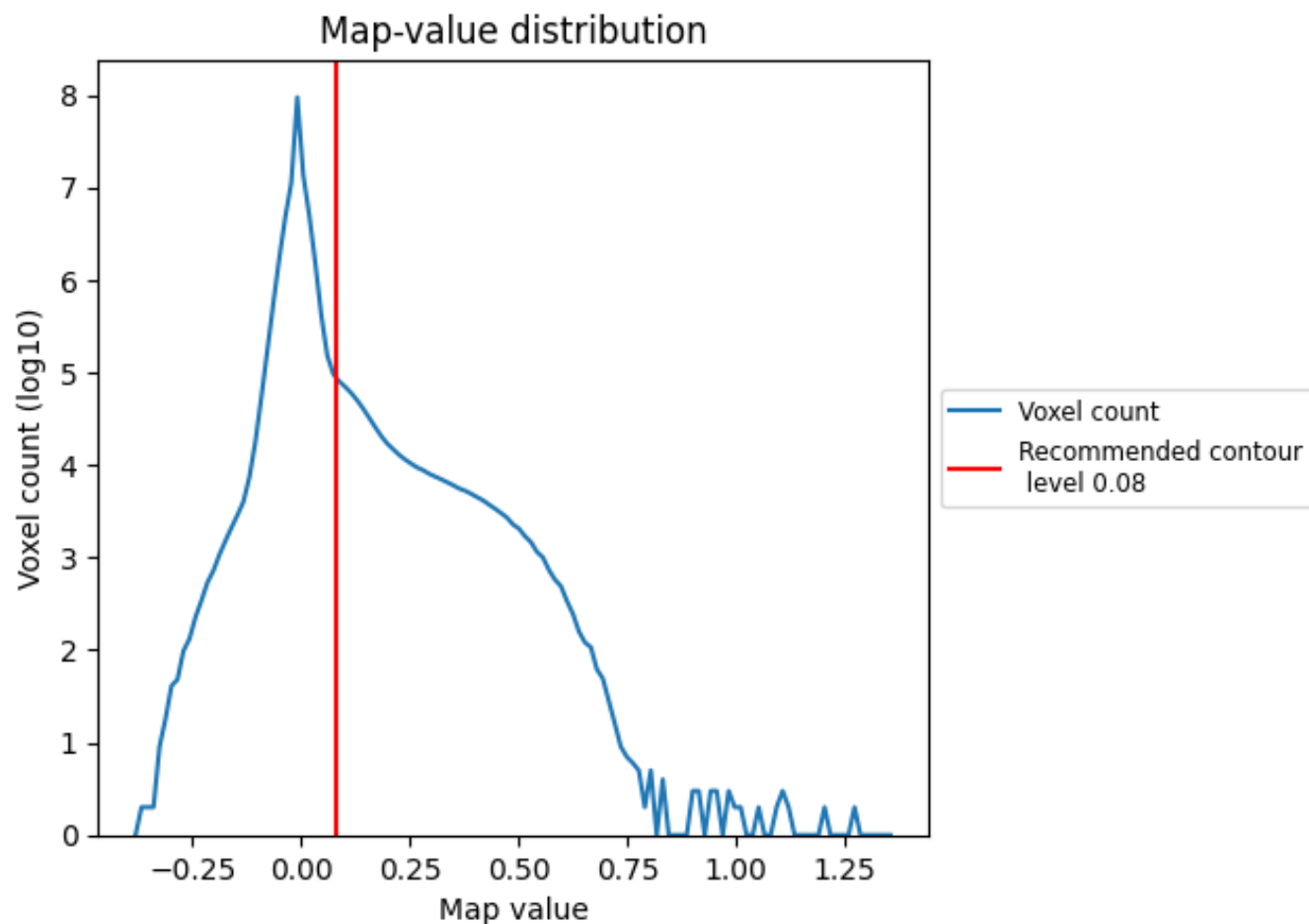
6.6 Mask visualisation [i](#)

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

7 Map analysis [i](#)

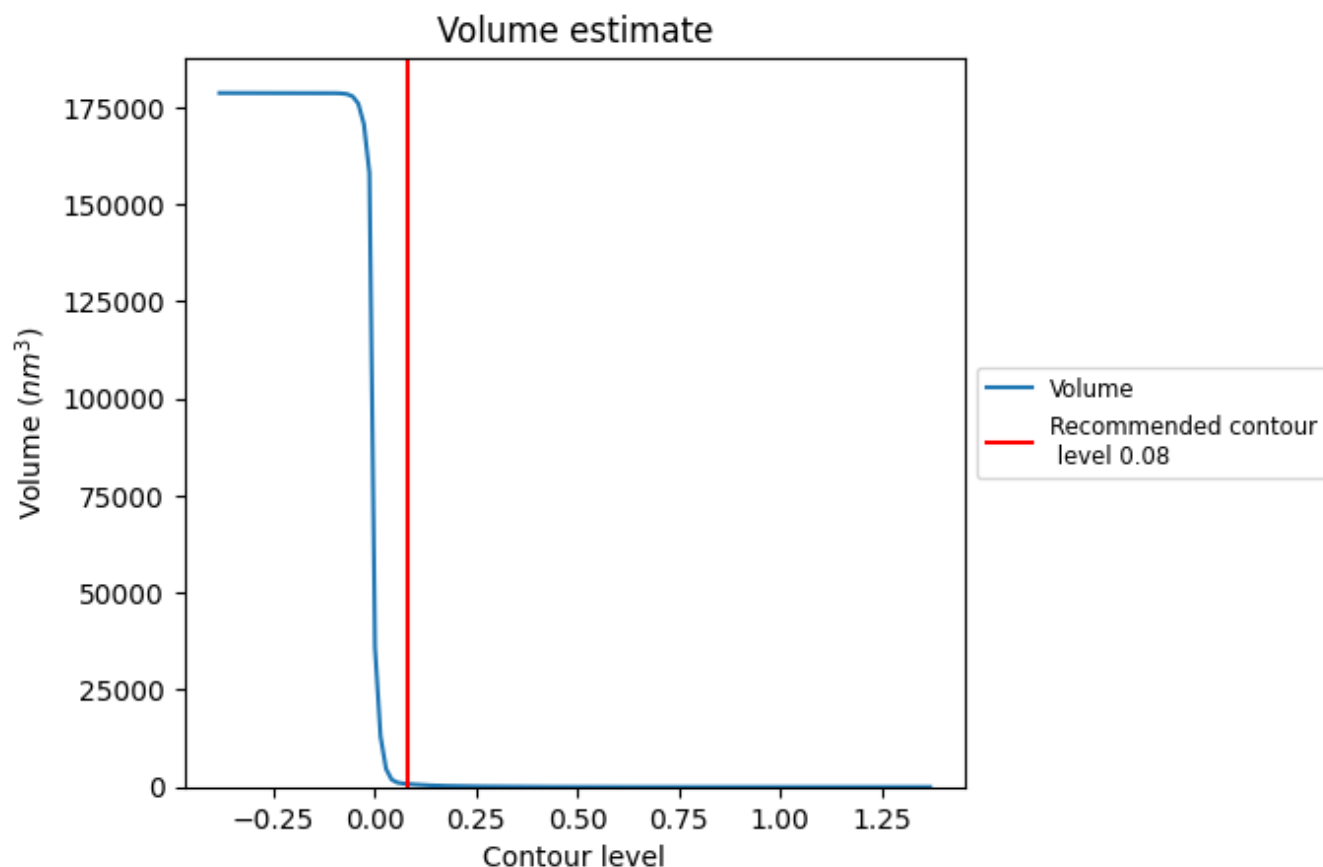
This section contains the results of statistical analysis of the map.

7.1 Map-value distribution [i](#)



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.

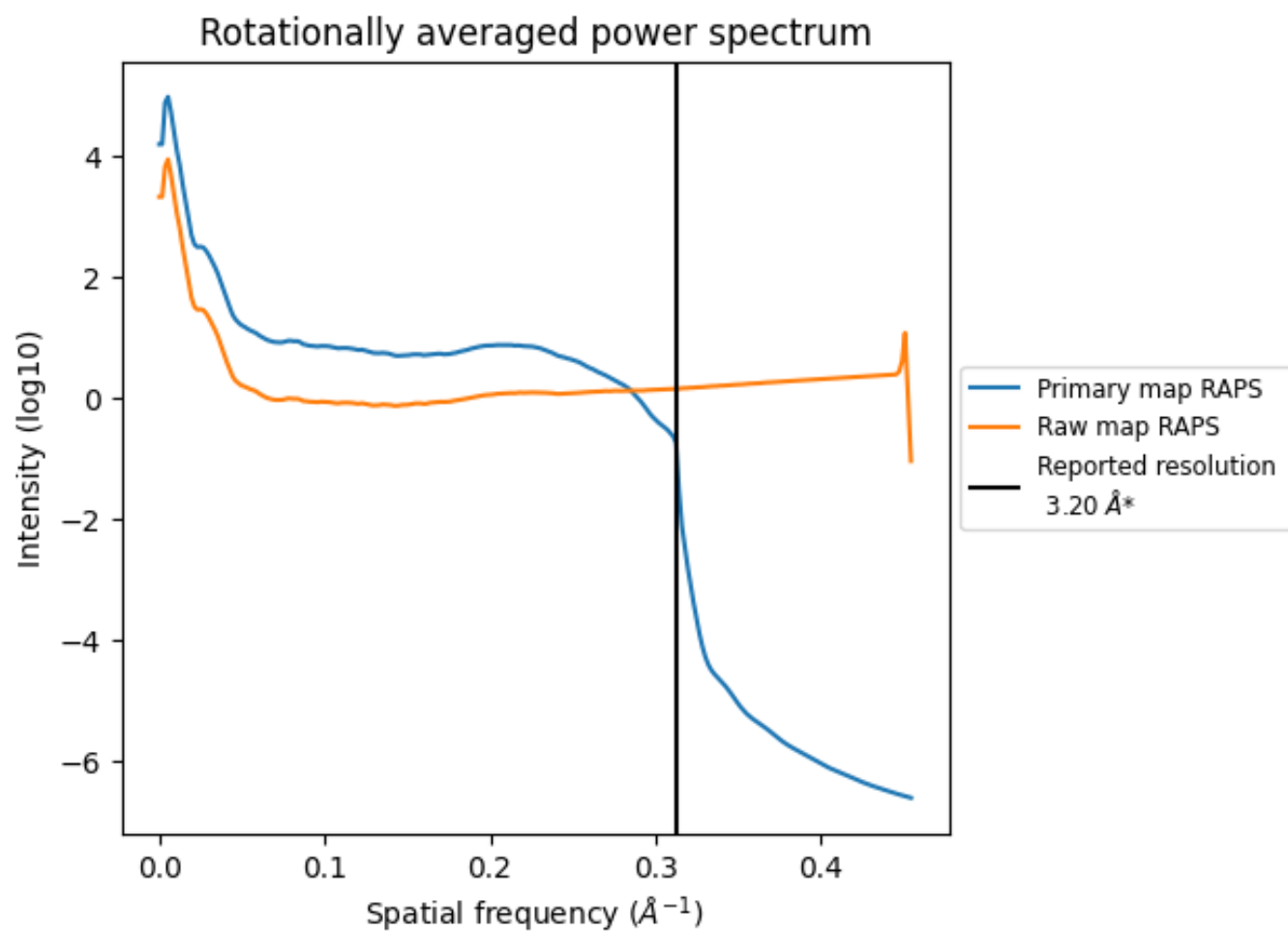
7.2 Volume estimate [i](#)



The volume at the recommended contour level is 799 nm³; this corresponds to an approximate mass of 722 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](#)

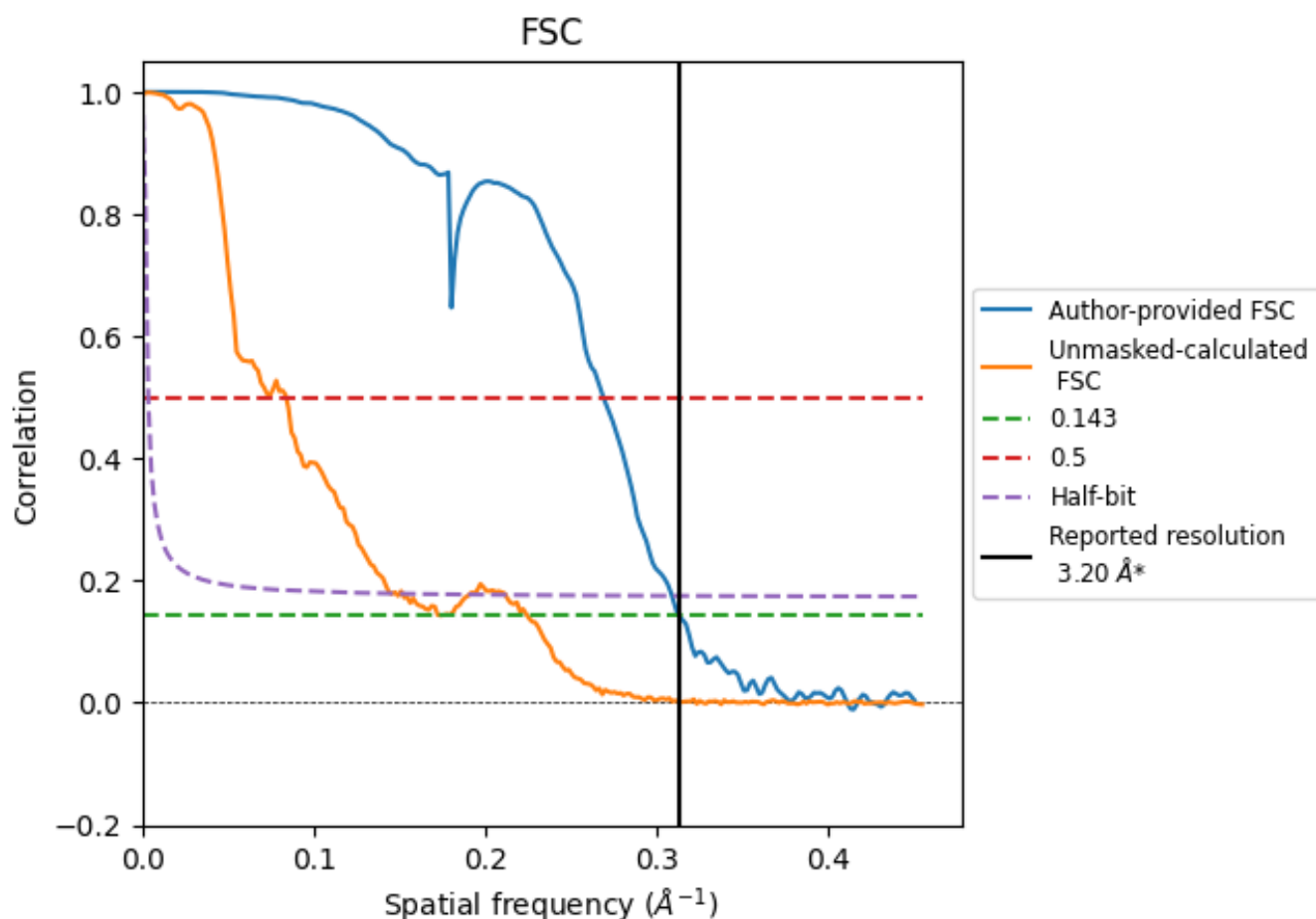


*Reported resolution corresponds to spatial frequency of 0.312 Å⁻¹

8 Fourier-Shell correlation [i](#)

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 [i](#)



*Reported resolution corresponds to spatial frequency of 0.312 $\text{\AA}^{-1}$

8.2 Resolution estimates [i](#)

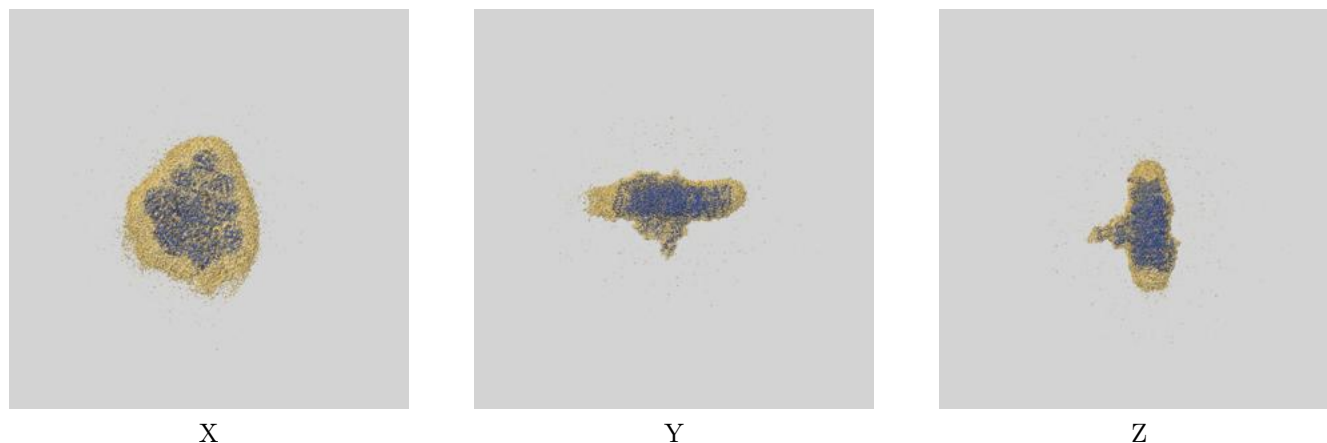
Resolution estimate (Å)	Estimation criterion (FSC cut-off)		
	0.143	0.5	Half-bit
Reported by author	3.20	-	-
Author-provided FSC curve	3.20	3.72	3.24
Unmasked-calculated*	5.75	11.93	6.88

*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 5.75 differs from the reported value 3.2 by more than 10 %

9 Map-model fit [i](#)

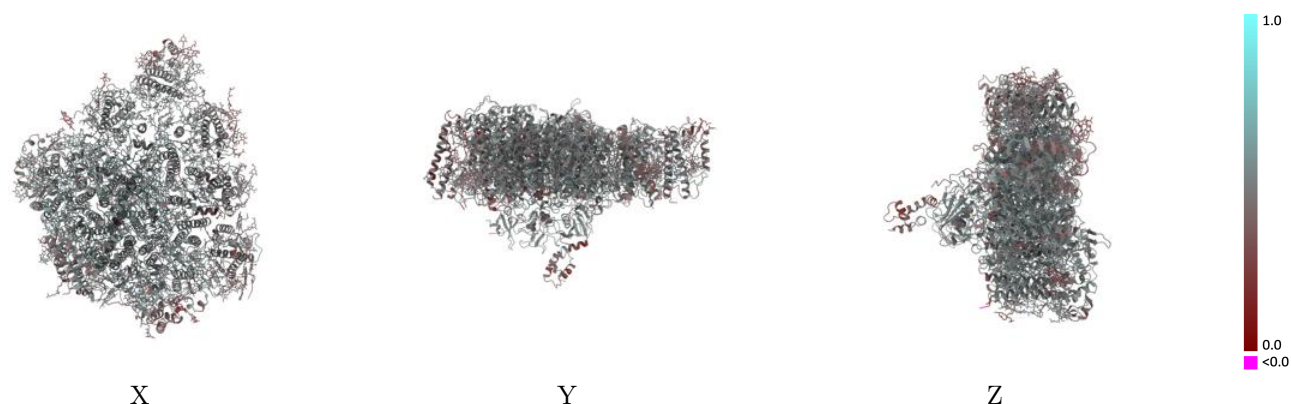
This section contains information regarding the fit between EMDB map EMD-60044 and PDB model 8ZET. Per-residue inclusion information can be found in [section 3](#) on [page 27](#).

9.1 Map-model overlay [i](#)



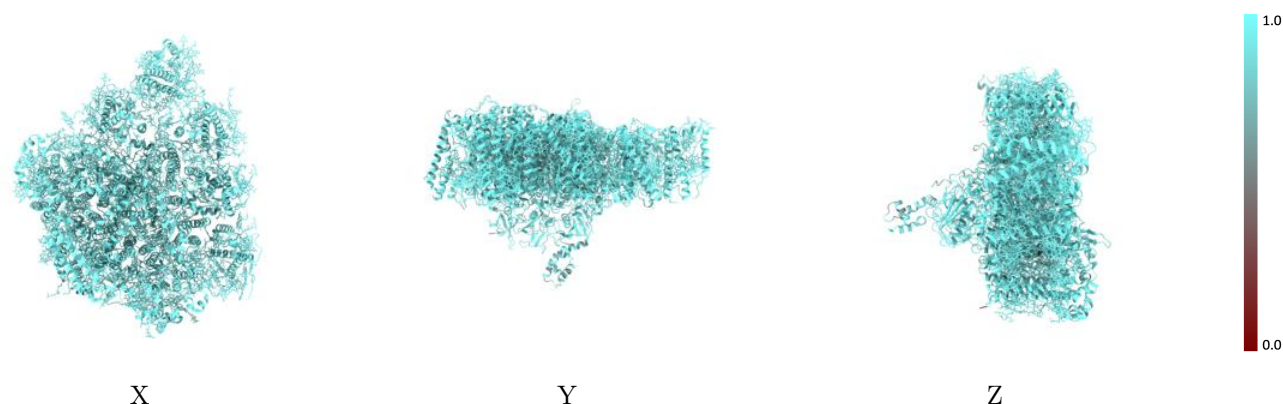
The images above show the 3D surface view of the map at the recommended contour level 0.08 at 50% transparency in yellow overlaid with a ribbon representation of the model coloured in blue. These images allow for the visual assessment of the quality of fit between the atomic model and the map.

9.2 Q-score mapped to coordinate model [i](#)



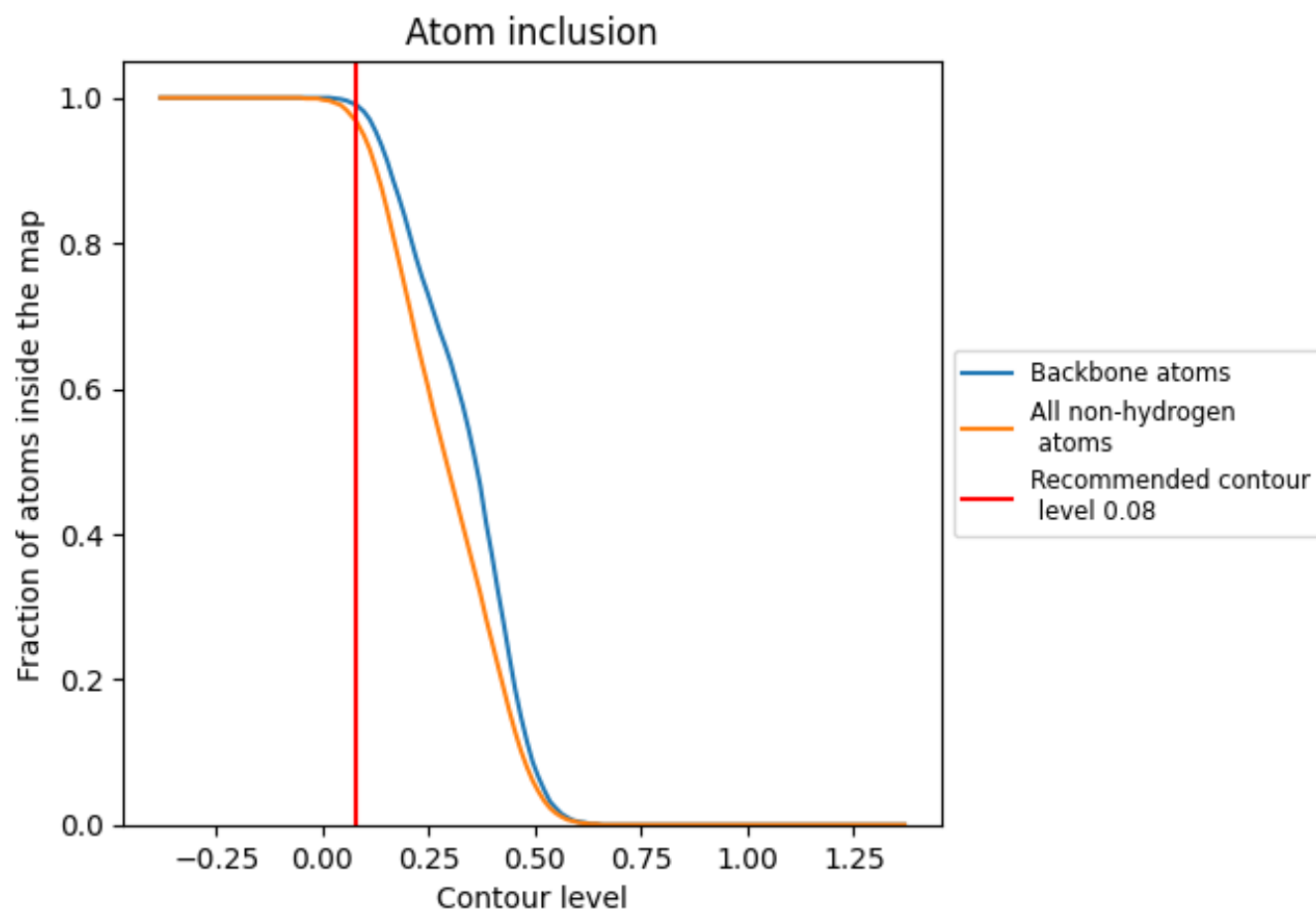
The images above show the model with each residue coloured according to its Q-score. This shows their resolvability in the map with higher Q-score values reflecting better resolvability. Please note: Q-score is calculating the resolvability of atoms, and thus high values are only expected at resolutions 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 (0.08).

9.4 Atom inclusion [i](#)



At the recommended contour level, 99% of all backbone atoms, 97% 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 (0.08) and Q-score for the entire model and for each chain.

Chain	Atom inclusion	Q-score
All	0.9670	0.4850
B	0.9740	0.4570
C	0.9780	0.4520
D	0.9670	0.4720
E	0.9700	0.4710
H	0.9600	0.4360
a	0.9740	0.5120
b	0.9690	0.5060
c	0.9880	0.5100
d	0.9790	0.5070
e	0.9510	0.4720
f	0.9650	0.4760
g	0.8680	0.4000
i	0.9640	0.5130
j	0.9490	0.4710
l	0.9770	0.5060
m	0.9440	0.4770
r	0.9330	0.3970

1.0

0.0

<0.0