



Full wwPDB EM Validation Report ⓘ

Aug 10, 2022 – 03:30 pm BST

PDB ID : 7PI5
EMDB ID : EMD-13430
Title : Unstacked stretched Dunaliella PSII
Authors : Caspy, I.; Fadeeva, M.; Mazor, Y.; Nelson, N.
Deposited on : 2021-08-19
Resolution : 2.78 Å (reported)
Based on initial model : 6KAC

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:

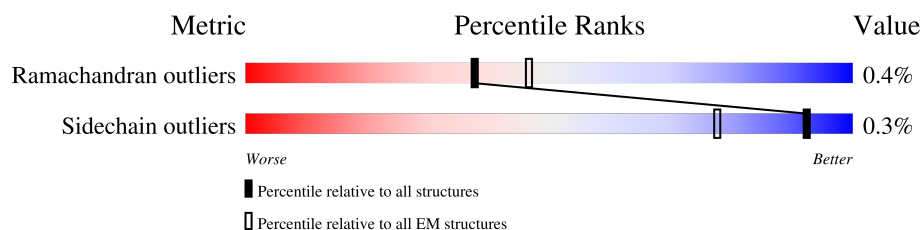
EMDB validation analysis : 0.0.1.dev8
Mogul : 1.8.4, CSD as541be (2020)
MolProbity : 4.02b-467
buster-report : 1.1.7 (2018)
Percentile statistics : 20191225.v01 (using entries in the PDB archive December 25th 2019)
Ideal geometry (proteins) : Engh & Huber (2001)
Ideal geometry (DNA, RNA) : Parkinson et al. (1996)
Validation Pipeline (wwPDB-VP) : 2.29

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 2.78 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.



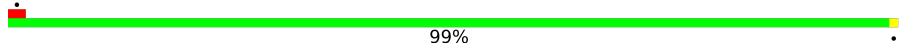
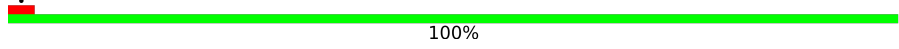
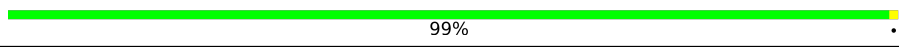
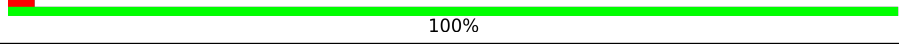
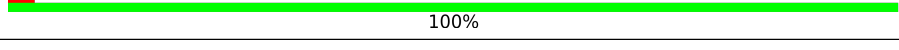
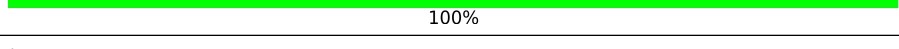
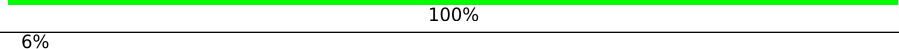
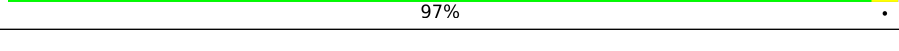
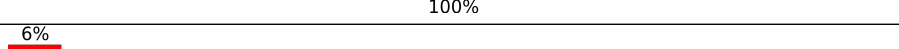
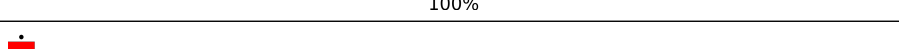
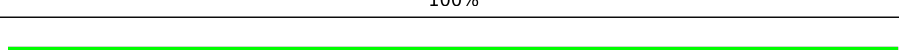
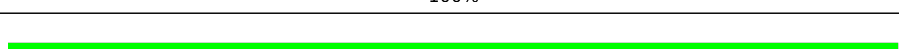
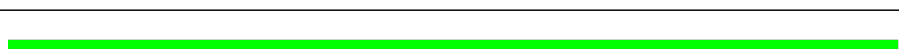
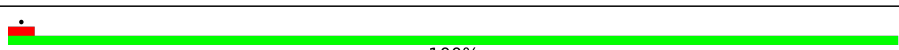
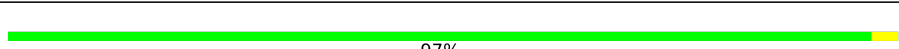

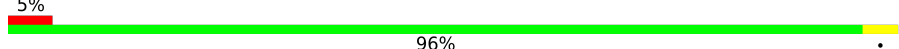
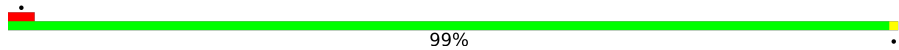
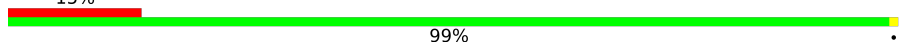
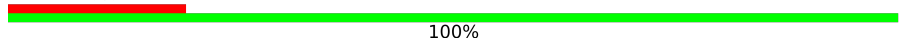
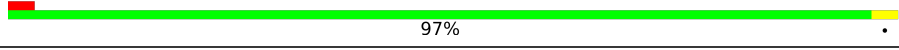
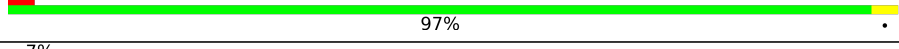
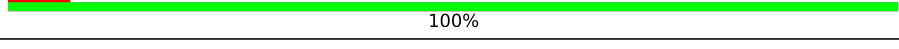
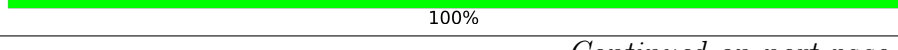

Metric	Whole archive (#Entries)	EM structures (#Entries)
Ramachandran outliers	154571	4023
Sidechain outliers	154315	3826

The table below summarises the geometric issues observed across the polymeric chains and their fit to the map. The red, orange, yellow and green segments of the bar indicate the fraction of residues that contain outliers for ≥ 3 , 2, 1 and 0 types of geometric quality criteria respectively. A grey segment represents the fraction of residues that are not modelled. The numeric value for each fraction is indicated below the corresponding segment, with a dot representing fractions $\leq 5\%$. 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	336	<div> <div>9%</div> <div>99%</div> <div>.</div> </div>
1	a	336	<div> <div>8%</div> <div>99%</div> <div>.</div> </div>
2	B	484	<div> <div>100%</div> </div>
2	b	484	<div> <div>100%</div> </div>
3	V	32	<div> <div>100%</div> </div>
3	v	32	<div> <div>100%</div> </div>
4	C	449	<div> <div>99%</div> <div>.</div> </div>
4	c	449	<div> <div>99%</div> <div>.</div> </div>
5	D	348	<div> <div>99%</div> <div>.</div> </div>

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Mol	Chain	Length	Quality of chain
5	d	348	 99%
6	E	76	 100%
6	e	76	 99%
7	F	31	 100%
7	f	31	 100%
8	H	67	 100%
8	h	67	 100%
9	I	35	 6%97%
9	i	35	 100%
10	J	36	 6%100%
10	j	36	 100%
11	K	37	 100%
11	k	37	 100%
12	L	38	 100%
12	l	38	 100%
13	M	32	 97%
13	m	32	 97%
14	O	238	 5%96%
14	o	238	 99%
15	P	187	 15%99%
15	p	187	 20%100%
16	T	30	 97%
16	t	30	 97%
17	W	44	 7%100%
17	w	44	 100%

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Mol	Chain	Length	Quality of chain
18	X	30	 100%
18	x	30	 100%
19	Z	61	 98%
19	z	61	 100%
20	N	222	 99%
20	n	222	 100%
21	G	221	 99%
21	g	221	 100%
22	R	202	 24% 100%
22	r	202	 26% 100%
23	S	243	 98%
23	s	243	 99%
24	Y	222	 99%
24	y	222	 99%
25	U	27	 7% 100%
25	u	27	 100%

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
29	CLA	A	405	X	-	-	-
29	CLA	A	406	X	-	-	-
29	CLA	A	407	X	-	-	-
29	CLA	A	410	X	-	-	-
29	CLA	B	602	X	-	-	-
29	CLA	B	603	X	-	-	-
29	CLA	B	604	X	-	-	-
29	CLA	B	605	X	-	-	-
29	CLA	B	606	X	-	-	-
29	CLA	B	607	X	-	-	-

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Mol	Type	Chain	Res	Chirality	Geometry	Clashes	Electron density
29	CLA	B	608	X	-	-	-
29	CLA	B	609	X	-	-	-
29	CLA	B	610	X	-	-	-
29	CLA	B	611	X	-	-	-
29	CLA	B	612	X	-	-	-
29	CLA	B	613	X	-	-	-
29	CLA	B	614	X	-	-	-
29	CLA	B	615	X	-	-	-
29	CLA	B	616	X	-	-	-
29	CLA	B	617	X	-	-	-
29	CLA	C	501	X	-	-	-
29	CLA	C	502	X	-	-	-
29	CLA	C	503	X	-	-	-
29	CLA	C	504	X	-	-	-
29	CLA	C	505	X	-	-	-
29	CLA	C	506	X	-	-	-
29	CLA	C	507	X	-	-	-
29	CLA	C	508	X	-	-	-
29	CLA	C	509	X	-	-	-
29	CLA	C	510	X	-	-	-
29	CLA	C	511	X	-	-	-
29	CLA	C	512	X	-	-	-
29	CLA	C	513	X	-	-	-
29	CLA	D	402	X	-	-	-
29	CLA	D	403	X	-	-	-
29	CLA	G	602	X	-	-	-
29	CLA	G	603	X	-	-	-
29	CLA	G	604	X	-	-	-
29	CLA	G	610	X	-	-	-
29	CLA	G	611	X	-	-	-
29	CLA	G	612	X	-	-	-
29	CLA	G	613	X	-	-	-
29	CLA	G	614	X	-	-	-
29	CLA	N	602	X	-	-	-
29	CLA	N	603	X	-	-	-
29	CLA	N	604	X	-	-	-
29	CLA	N	610	X	-	-	-
29	CLA	N	611	X	-	-	-
29	CLA	N	612	X	-	-	-
29	CLA	N	613	X	-	-	-
29	CLA	N	614	X	-	-	-
29	CLA	R	602	X	-	-	-

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Mol	Type	Chain	Res	Chirality	Geometry	Clashes	Electron density
29	CLA	R	603	X	-	-	-
29	CLA	R	604	X	-	-	-
29	CLA	R	608	X	-	-	-
29	CLA	R	609	X	-	-	-
29	CLA	R	610	X	-	-	-
29	CLA	R	611	X	-	-	-
29	CLA	R	612	X	-	-	-
29	CLA	R	613	X	-	-	-
29	CLA	S	602	X	-	-	-
29	CLA	S	603	X	-	-	-
29	CLA	S	604	X	-	-	-
29	CLA	S	605	X	-	-	-
29	CLA	S	609	X	-	-	-
29	CLA	S	610	X	-	-	-
29	CLA	S	611	X	-	-	-
29	CLA	S	612	X	-	-	-
29	CLA	S	613	X	-	-	-
29	CLA	S	614	X	-	-	-
29	CLA	S	617	X	-	-	-
29	CLA	Y	602	X	-	-	-
29	CLA	Y	603	X	-	-	-
29	CLA	Y	604	X	-	-	-
29	CLA	Y	608	X	-	-	-
29	CLA	Y	610	X	-	-	-
29	CLA	Y	611	X	-	-	-
29	CLA	Y	612	X	-	-	-
29	CLA	Y	613	X	-	-	-
29	CLA	Y	614	X	-	-	-
29	CLA	a	405	X	-	-	-
29	CLA	a	406	X	-	-	-
29	CLA	a	407	X	-	-	-
29	CLA	a	410	X	-	-	-
29	CLA	b	602	X	-	-	-
29	CLA	b	603	X	-	-	-
29	CLA	b	604	X	-	-	-
29	CLA	b	605	X	-	-	-
29	CLA	b	606	X	-	-	-
29	CLA	b	607	X	-	-	-
29	CLA	b	608	X	-	-	-
29	CLA	b	609	X	-	-	-
29	CLA	b	610	X	-	-	-
29	CLA	b	611	X	-	-	-

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Mol	Type	Chain	Res	Chirality	Geometry	Clashes	Electron density
29	CLA	b	612	X	-	-	-
29	CLA	b	613	X	-	-	-
29	CLA	b	614	X	-	-	-
29	CLA	b	615	X	-	-	-
29	CLA	b	616	X	-	-	-
29	CLA	b	617	X	-	-	-
29	CLA	c	501	X	-	-	-
29	CLA	c	502	X	-	-	-
29	CLA	c	503	X	-	-	-
29	CLA	c	504	X	-	-	-
29	CLA	c	505	X	-	-	-
29	CLA	c	506	X	-	-	-
29	CLA	c	507	X	-	-	-
29	CLA	c	508	X	-	-	-
29	CLA	c	509	X	-	-	-
29	CLA	c	510	X	-	-	-
29	CLA	c	511	X	-	-	-
29	CLA	c	512	X	-	-	-
29	CLA	c	513	X	-	-	-
29	CLA	d	402	X	-	-	-
29	CLA	d	403	X	-	-	-
29	CLA	g	602	X	-	-	-
29	CLA	g	603	X	-	-	-
29	CLA	g	604	X	-	-	-
29	CLA	g	610	X	-	-	-
29	CLA	g	611	X	-	-	-
29	CLA	g	612	X	-	-	-
29	CLA	g	613	X	-	-	-
29	CLA	g	614	X	-	-	-
29	CLA	n	602	X	-	-	-
29	CLA	n	603	X	-	-	-
29	CLA	n	604	X	-	-	-
29	CLA	n	610	X	-	-	-
29	CLA	n	611	X	-	-	-
29	CLA	n	612	X	-	-	-
29	CLA	n	613	X	-	-	-
29	CLA	n	614	X	-	-	-
29	CLA	r	602	X	-	-	-
29	CLA	r	603	X	-	-	-
29	CLA	r	604	X	-	-	-
29	CLA	r	608	X	-	-	-
29	CLA	r	609	X	-	-	-

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Mol	Type	Chain	Res	Chirality	Geometry	Clashes	Electron density
29	CLA	r	610	X	-	-	-
29	CLA	r	611	X	-	-	-
29	CLA	r	612	X	-	-	-
29	CLA	r	613	X	-	-	-
29	CLA	s	602	X	-	-	-
29	CLA	s	603	X	-	-	-
29	CLA	s	604	X	-	-	-
29	CLA	s	605	X	-	-	-
29	CLA	s	609	X	-	-	-
29	CLA	s	610	X	-	-	-
29	CLA	s	611	X	-	-	-
29	CLA	s	612	X	-	-	-
29	CLA	s	613	X	-	-	-
29	CLA	s	614	X	-	-	-
29	CLA	s	617	X	-	-	-
29	CLA	y	602	X	-	-	-
29	CLA	y	603	X	-	-	-
29	CLA	y	604	X	-	-	-
29	CLA	y	608	X	-	-	-
29	CLA	y	610	X	-	-	-
29	CLA	y	611	X	-	-	-
29	CLA	y	612	X	-	-	-
29	CLA	y	613	X	-	-	-
29	CLA	y	614	X	-	-	-
35	C7Z	B	620	X	-	-	-
35	C7Z	b	620	X	-	-	-
40	LMK	C	527	X	-	-	-
40	LMK	c	627	X	-	-	-
44	RRX	H	101	X	-	-	-
44	RRX	h	101	X	-	-	-
45	CHL	G	601	X	-	-	-
45	CHL	G	605	X	-	-	-
45	CHL	G	606	X	-	-	-
45	CHL	G	607	X	-	-	-
45	CHL	G	608	X	-	-	-
45	CHL	G	609	X	-	-	-
45	CHL	N	601	X	-	-	-
45	CHL	N	605	X	-	-	-
45	CHL	N	606	X	-	-	-
45	CHL	N	607	X	-	-	-
45	CHL	N	608	X	-	-	-
45	CHL	N	609	X	-	-	-

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Mol	Type	Chain	Res	Chirality	Geometry	Clashes	Electron density
45	CHL	R	606	X	-	-	-
45	CHL	R	607	X	-	-	-
45	CHL	S	601	X	-	-	-
45	CHL	S	606	X	-	-	-
45	CHL	S	607	X	-	-	-
45	CHL	S	608	X	-	-	-
45	CHL	Y	601	X	-	-	-
45	CHL	Y	605	X	-	-	-
45	CHL	Y	606	X	-	-	-
45	CHL	Y	607	X	-	-	-
45	CHL	Y	609	X	-	-	-
45	CHL	g	601	X	-	-	-
45	CHL	g	605	X	-	-	-
45	CHL	g	606	X	-	-	-
45	CHL	g	607	X	-	-	-
45	CHL	g	608	X	-	-	-
45	CHL	g	609	X	-	-	-
45	CHL	n	601	X	-	-	-
45	CHL	n	605	X	-	-	-
45	CHL	n	606	X	-	-	-
45	CHL	n	607	X	-	-	-
45	CHL	n	608	X	-	-	-
45	CHL	n	609	X	-	-	-
45	CHL	r	606	X	-	-	-
45	CHL	r	607	X	-	-	-
45	CHL	s	601	X	-	-	-
45	CHL	s	606	X	-	-	-
45	CHL	s	607	X	-	-	-
45	CHL	s	608	X	-	-	-
45	CHL	y	601	X	-	-	-
45	CHL	y	605	X	-	-	-
45	CHL	y	606	X	-	-	-
45	CHL	y	607	X	-	-	-
45	CHL	y	609	X	-	-	-
46	LUT	S	620	X	-	-	-
46	LUT	s	620	X	-	-	-
47	XAT	G	622	X	-	-	-
47	XAT	N	622	X	-	-	-
47	XAT	R	621	X	-	-	-
47	XAT	Y	622	X	-	-	-
47	XAT	g	622	X	-	-	-
47	XAT	n	622	X	-	-	-

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Mol	Type	Chain	Res	Chirality	Geometry	Clashes	Electron density
47	XAT	r	622	X	-	-	-
47	XAT	y	622	X	-	-	-

2 Entry composition

There are 52 unique types of molecules in this entry. The entry contains 76287 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 II protein D1.

Mol	Chain	Residues	Atoms					AltConf	Trace
1	A	336	Total	C	N	O	S	1	0
			2638	1721	432	468	17		
1	a	336	Total	C	N	O	S	1	0
			2638	1721	432	468	17		

- Molecule 2 is a protein called Photosystem II CP47 reaction center protein.

Mol	Chain	Residues	Atoms					AltConf	Trace
2	B	484	Total	C	N	O	S	0	0
			3783	2480	630	663	10		
2	b	484	Total	C	N	O	S	0	0
			3783	2480	630	663	10		

There are 4 discrepancies between the modelled and reference sequences:

Chain	Residue	Modelled	Actual	Comment	Reference
B	298	VAL	LEU	variant	UNP D0FY05
B	415	SER	LEU	variant	UNP D0FY05
b	298	VAL	LEU	variant	UNP D0FY05
b	415	SER	LEU	variant	UNP D0FY05

- Molecule 3 is a protein called Photosystem II reaction center protein Ycf12.

Mol	Chain	Residues	Atoms				AltConf	Trace
3	V	32	Total	C	N	O	0	0
			227	152	37	38		
3	v	32	Total	C	N	O	0	0
			227	152	37	38		

- Molecule 4 is a protein called Photosystem II CP43 reaction center protein.

Mol	Chain	Residues	Atoms					AltConf	Trace
4	C	449	Total	C	N	O	S	0	0
			3483	2282	581	607	13		
4	c	449	Total	C	N	O	S	0	0
			3483	2282	581	607	13		

- Molecule 5 is a protein called Photosystem II D2 protein.

Mol	Chain	Residues	Atoms					AltConf	Trace
5	D	348	Total	C	N	O	S	0	0
			2766	1824	454	477	11		
5	d	348	Total	C	N	O	S	0	0
			2766	1824	454	477	11		

- Molecule 6 is a protein called Cytochrome b559 subunit alpha.

Mol	Chain	Residues	Atoms				AltConf	Trace
6	E	76	Total	C	N	O	0	0
			621	404	102	115		
6	e	76	Total	C	N	O	0	0
			621	404	102	115		

- Molecule 7 is a protein called Cytochrome b559 subunit beta.

Mol	Chain	Residues	Atoms					AltConf	Trace
7	F	31	Total	C	N	O	S	0	0
			252	172	42	37	1		
7	f	31	Total	C	N	O	S	0	0
			252	172	42	37	1		

- Molecule 8 is a protein called Photosystem II reaction center protein H.

Mol	Chain	Residues	Atoms					AltConf	Trace
8	H	67	Total	C	N	O	S	0	0
			503	334	76	92	1		
8	h	67	Total	C	N	O	S	0	0
			503	334	76	92	1		

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

Mol	Chain	Residues	Atoms					AltConf	Trace
9	I	35	Total	C	N	O	S	0	0
			279	190	42	46	1		

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Mol	Chain	Residues	Atoms					AltConf	Trace
9	i	35	Total	C	N	O	S	0	0
			279	190	42	46	1		

- Molecule 10 is a protein called Photosystem II reaction center protein J.

Mol	Chain	Residues	Atoms					AltConf	Trace
10	J	36	Total	C	N	O		0	0
			266	183	40	43			
10	j	36	Total	C	N	O		0	0
			266	183	40	43			

There are 4 discrepancies between the modelled and reference sequences:

Chain	Residue	Modelled	Actual	Comment	Reference
J	7	ILE	THR	conflict	UNP A0A1C8XRM8
J	42	LEU	GLN	conflict	UNP A0A1C8XRM8
j	7	ILE	THR	conflict	UNP A0A1C8XRM8
j	42	LEU	GLN	conflict	UNP A0A1C8XRM8

- Molecule 11 is a protein called Photosystem II reaction center protein K.

Mol	Chain	Residues	Atoms					AltConf	Trace
11	K	37	Total	C	N	O		0	0
			297	207	43	47			
11	k	37	Total	C	N	O		0	0
			297	207	43	47			

- Molecule 12 is a protein called Photosystem II reaction center protein L.

Mol	Chain	Residues	Atoms					AltConf	Trace
12	L	38	Total	C	N	O	S	0	0
			313	209	51	52	1		
12	l	38	Total	C	N	O	S	0	0
			313	209	51	52	1		

- Molecule 13 is a protein called Photosystem II reaction center protein M.

Mol	Chain	Residues	Atoms					AltConf	Trace
13	M	32	Total	C	N	O		0	0
			243	164	34	45			
13	m	32	Total	C	N	O		0	0
			243	164	34	45			

There are 2 discrepancies between the modelled and reference sequences:

Chain	Residue	Modelled	Actual	Comment	Reference
M	9	THR	ILE	variant	UNP D0FXZ3
m	9	THR	ILE	variant	UNP D0FXZ3

- Molecule 14 is a protein called PsbO.

Mol	Chain	Residues	Atoms					AltConf	Trace
14	O	238	Total	C	N	O	S	0	0
			1820	1149	295	370	6		
14	o	238	Total	C	N	O	S	0	0
			1820	1149	295	370	6		

- Molecule 15 is a protein called PsbP.

Mol	Chain	Residues	Atoms					AltConf	Trace
15	P	187	Total	C	N	O	S	0	0
			1444	916	242	285	1		
15	p	187	Total	C	N	O	S	0	0
			1444	916	242	285	1		

- Molecule 16 is a protein called Photosystem II reaction center protein T.

Mol	Chain	Residues	Atoms					AltConf	Trace
16	T	30	Total	C	N	O	S	0	0
			247	171	36	39	1		
16	t	30	Total	C	N	O	S	0	0
			247	171	36	39	1		

- Molecule 17 is a protein called PSII 6.1 kDa protein.

Mol	Chain	Residues	Atoms					AltConf	Trace
17	W	44	Total	C	N	O	S	0	0
			332	215	53	63	1		
17	w	44	Total	C	N	O	S	0	0
			332	215	53	63	1		

There are 4 discrepancies between the modelled and reference sequences:

Chain	Residue	Modelled	Actual	Comment	Reference
W	65	MET	LEU	conflict	UNP A0A7S3QU88
W	96	TYR	PHE	conflict	UNP A0A7S3QU88
w	65	MET	LEU	conflict	UNP A0A7S3QU88

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Chain	Residue	Modelled	Actual	Comment	Reference
w	96	TYR	PHE	conflict	UNP A0A7S3QU88

- Molecule 18 is a protein called Hypothetical protein.

Mol	Chain	Residues	Atoms				AltConf	Trace
18	X	30	Total	C	N	O	0	0
			201	132	32	37		
18	x	30	Total	C	N	O	0	0
			201	132	32	37		

There are 2 discrepancies between the modelled and reference sequences:

Chain	Residue	Modelled	Actual	Comment	Reference
X	81	VAL	THR	conflict	UNP A0A7S3VKF3
x	81	VAL	THR	conflict	UNP A0A7S3VKF3

- Molecule 19 is a protein called Photosystem II reaction center protein Z.

Mol	Chain	Residues	Atoms					AltConf	Trace
19	Z	61	Total	C	N	O	S	0	0
			457	312	68	76	1		
19	z	61	Total	C	N	O	S	0	0
			457	312	68	76	1		

- Molecule 20 is a protein called LHCII M3.

Mol	Chain	Residues	Atoms					AltConf	Trace
20	N	222	Total	C	N	O	S	0	0
			1703	1100	277	321	5		
20	n	222	Total	C	N	O	S	0	0
			1703	1100	277	321	5		

- Molecule 21 is a protein called Chlorophyll a-b binding protein, chloroplastic.

Mol	Chain	Residues	Atoms					AltConf	Trace
21	G	221	Total	C	N	O	S	0	0
			1680	1085	271	321	3		
21	g	221	Total	C	N	O	S	0	0
			1680	1085	271	321	3		

There are 2 discrepancies between the modelled and reference sequences:

Chain	Residue	Modelled	Actual	Comment	Reference
G	180	ALA	PRO	conflict	UNP A1XKU7
g	180	ALA	PRO	conflict	UNP A1XKU7

- Molecule 22 is a protein called CP29.

Mol	Chain	Residues	Atoms					AltConf	Trace
22	R	202	Total	C	N	O	S	0	0
			1533	974	258	298	3		
22	r	202	Total	C	N	O	S	0	0
			1533	974	258	298	3		

- Molecule 23 is a protein called CP26.

Mol	Chain	Residues	Atoms					AltConf	Trace
23	S	242	Total	C	N	O	S	0	0
			1849	1195	297	354	3		
23	s	243	Total	C	N	O	S	0	0
			1856	1200	298	355	3		

There are 14 discrepancies between the modelled and reference sequences:

Chain	Residue	Modelled	Actual	Comment	Reference
S	56	ASP	GLU	conflict	UNP A0A7S3VRZ8
S	119	ILE	LEU	conflict	UNP A0A7S3VRZ8
S	209	LYS	GLN	conflict	UNP A0A7S3VRZ8
S	244	ILE	VAL	conflict	UNP A0A7S3VRZ8
S	245	ALA	GLY	conflict	UNP A0A7S3VRZ8
S	264	ILE	PHE	conflict	UNP A0A7S3VRZ8
S	268	LEU	ILE	conflict	UNP A0A7S3VRZ8
s	56	ASP	GLU	conflict	UNP A0A7S3VRZ8
s	119	ILE	LEU	conflict	UNP A0A7S3VRZ8
s	209	LYS	GLN	conflict	UNP A0A7S3VRZ8
s	244	ILE	VAL	conflict	UNP A0A7S3VRZ8
s	245	ALA	GLY	conflict	UNP A0A7S3VRZ8
s	264	ILE	PHE	conflict	UNP A0A7S3VRZ8
s	268	LEU	ILE	conflict	UNP A0A7S3VRZ8

- Molecule 24 is a protein called LHCII M1.

Mol	Chain	Residues	Atoms					AltConf	Trace
24	Y	222	Total	C	N	O	S	0	0
			1667	1080	272	312	3		

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Mol	Chain	Residues	Atoms					AltConf	Trace
24	y	222	Total	C	N	O	S	0	0
			1667	1080	272	312	3		

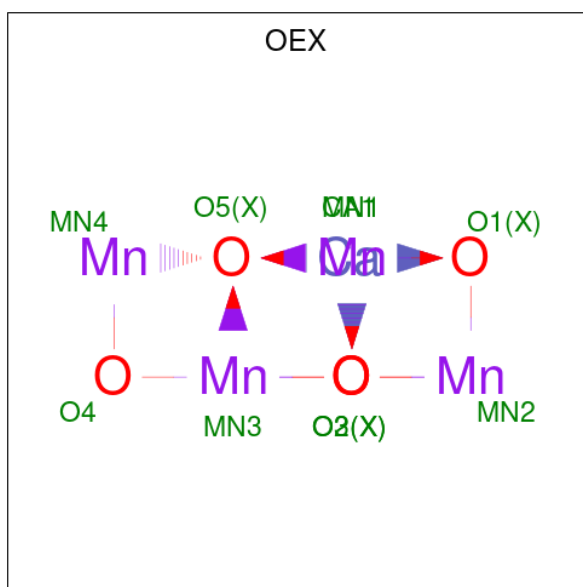
There are 16 discrepancies between the modelled and reference sequences:

Chain	Residue	Modelled	Actual	Comment	Reference
Y	49	PHE	TYR	conflict	UNP A0A6S8N9J6
Y	52	SER	ALA	conflict	UNP A0A6S8N9J6
Y	73	THR	SER	conflict	UNP A0A6S8N9J6
Y	81	THR	ASN	conflict	UNP A0A6S8N9J6
Y	123	ILE	VAL	conflict	UNP A0A6S8N9J6
Y	220	LEU	PHE	conflict	UNP A0A6S8N9J6
Y	235	GLN	THR	conflict	UNP A0A6S8N9J6
Y	259	THR	SER	conflict	UNP A0A6S8N9J6
y	49	PHE	TYR	conflict	UNP A0A6S8N9J6
y	52	SER	ALA	conflict	UNP A0A6S8N9J6
y	73	THR	SER	conflict	UNP A0A6S8N9J6
y	81	THR	ASN	conflict	UNP A0A6S8N9J6
y	123	ILE	VAL	conflict	UNP A0A6S8N9J6
y	220	LEU	PHE	conflict	UNP A0A6S8N9J6
y	235	GLN	THR	conflict	UNP A0A6S8N9J6
y	259	THR	SER	conflict	UNP A0A6S8N9J6

- Molecule 25 is a protein called PsbU.

Mol	Chain	Residues	Atoms					AltConf	Trace
25	U	27	Total	C	N	O	S	0	0
			224	134	42	47	1		
25	u	27	Total	C	N	O	S	0	0
			224	134	42	47	1		

- Molecule 26 is CA-MN4-O5 CLUSTER (three-letter code: OEX) (formula: CaMn_4O_5).



Mol	Chain	Residues	Atoms				AltConf
26	A	1	Total	Ca	Mn	O	0
			10	1	4	5	
26	a	1	Total	Ca	Mn	O	0
			10	1	4	5	

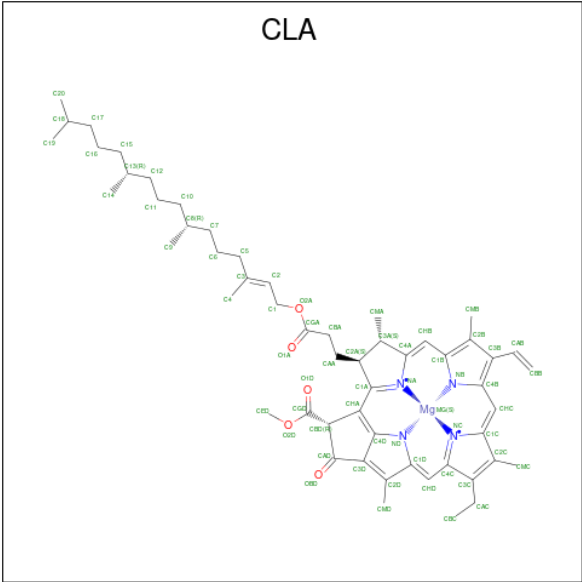
- Molecule 27 is FE (II) ION (three-letter code: FE2) (formula: Fe).

Mol	Chain	Residues	Atoms		AltConf
27	A	1	Total	Fe	0
			1	1	
27	a	1	Total	Fe	0
			1	1	

- Molecule 28 is CHLORIDE ION (three-letter code: CL) (formula: Cl).

Mol	Chain	Residues	Atoms		AltConf
28	A	2	Total	Cl	0
			2	2	
28	a	2	Total	Cl	0
			2	2	

- Molecule 29 is CHLOROPHYLL A (three-letter code: CLA) (formula: C₅₅H₇₂MgN₄O₅).



Mol	Chain	Residues	Atoms					AltConf
29	A	1	Total	C	Mg	N	O	0
			239	199	4	16	20	
29	A	1	Total	C	Mg	N	O	0
			239	199	4	16	20	
29	A	1	Total	C	Mg	N	O	0
			239	199	4	16	20	
29	A	1	Total	C	Mg	N	O	0
			239	199	4	16	20	
29	B	1	Total	C	Mg	N	O	0
			1040	880	16	64	80	
29	B	1	Total	C	Mg	N	O	0
			1040	880	16	64	80	
29	B	1	Total	C	Mg	N	O	0
			1040	880	16	64	80	
29	B	1	Total	C	Mg	N	O	0
			1040	880	16	64	80	
29	B	1	Total	C	Mg	N	O	0
			1040	880	16	64	80	
29	B	1	Total	C	Mg	N	O	0
			1040	880	16	64	80	
29	B	1	Total	C	Mg	N	O	0
			1040	880	16	64	80	
29	B	1	Total	C	Mg	N	O	0
			1040	880	16	64	80	
29	B	1	Total	C	Mg	N	O	0
			1040	880	16	64	80	

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Mol	Chain	Residues	Atoms					AltConf
29	B	1	Total	C	Mg	N	O	0
			1040	880	16	64	80	
29	B	1	Total	C	Mg	N	O	0
			1040	880	16	64	80	
29	B	1	Total	C	Mg	N	O	0
			1040	880	16	64	80	
29	B	1	Total	C	Mg	N	O	0
			1040	880	16	64	80	
29	B	1	Total	C	Mg	N	O	0
			1040	880	16	64	80	
29	C	1	Total	C	Mg	N	O	0
			845	715	13	52	65	
29	C	1	Total	C	Mg	N	O	0
			845	715	13	52	65	
29	C	1	Total	C	Mg	N	O	0
			845	715	13	52	65	
29	C	1	Total	C	Mg	N	O	0
			845	715	13	52	65	
29	C	1	Total	C	Mg	N	O	0
			845	715	13	52	65	
29	C	1	Total	C	Mg	N	O	0
			845	715	13	52	65	
29	C	1	Total	C	Mg	N	O	0
			845	715	13	52	65	
29	C	1	Total	C	Mg	N	O	0
			845	715	13	52	65	
29	C	1	Total	C	Mg	N	O	0
			845	715	13	52	65	
29	C	1	Total	C	Mg	N	O	0
			845	715	13	52	65	
29	C	1	Total	C	Mg	N	O	0
			845	715	13	52	65	
29	D	1	Total	C	Mg	N	O	0
			130	110	2	8	10	
29	D	1	Total	C	Mg	N	O	0
			130	110	2	8	10	

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Mol	Chain	Residues	Atoms					AltConf
29	N	1	Total 468	C 388	Mg 8	N 32	O 40	0
29	N	1	Total 468	C 388	Mg 8	N 32	O 40	0
29	N	1	Total 468	C 388	Mg 8	N 32	O 40	0
29	N	1	Total 468	C 388	Mg 8	N 32	O 40	0
29	N	1	Total 468	C 388	Mg 8	N 32	O 40	0
29	N	1	Total 468	C 388	Mg 8	N 32	O 40	0
29	N	1	Total 468	C 388	Mg 8	N 32	O 40	0
29	N	1	Total 468	C 388	Mg 8	N 32	O 40	0
29	N	1	Total 468	C 388	Mg 8	N 32	O 40	0
29	G	1	Total 446	C 368	Mg 8	N 32	O 38	0
29	G	1	Total 446	C 368	Mg 8	N 32	O 38	0
29	G	1	Total 446	C 368	Mg 8	N 32	O 38	0
29	G	1	Total 446	C 368	Mg 8	N 32	O 38	0
29	G	1	Total 446	C 368	Mg 8	N 32	O 38	0
29	G	1	Total 446	C 368	Mg 8	N 32	O 38	0
29	G	1	Total 446	C 368	Mg 8	N 32	O 38	0
29	G	1	Total 446	C 368	Mg 8	N 32	O 38	0
29	R	1	Total 501	C 411	Mg 9	N 36	O 45	0
29	R	1	Total 501	C 411	Mg 9	N 36	O 45	0
29	R	1	Total 501	C 411	Mg 9	N 36	O 45	0
29	R	1	Total 501	C 411	Mg 9	N 36	O 45	0
29	R	1	Total 501	C 411	Mg 9	N 36	O 45	0

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Mol	Chain	Residues	Atoms					AltConf
29	R	1	Total	C	Mg	N	O	0
			501	411	9	36	45	
29	R	1	Total	C	Mg	N	O	0
			501	411	9	36	45	
29	R	1	Total	C	Mg	N	O	0
			501	411	9	36	45	
29	R	1	Total	C	Mg	N	O	0
			501	411	9	36	45	
29	S	1	Total	C	Mg	N	O	0
			625	515	11	44	55	
29	S	1	Total	C	Mg	N	O	0
			625	515	11	44	55	
29	S	1	Total	C	Mg	N	O	0
			625	515	11	44	55	
29	S	1	Total	C	Mg	N	O	0
			625	515	11	44	55	
29	S	1	Total	C	Mg	N	O	0
			625	515	11	44	55	
29	S	1	Total	C	Mg	N	O	0
			625	515	11	44	55	
29	S	1	Total	C	Mg	N	O	0
			625	515	11	44	55	
29	S	1	Total	C	Mg	N	O	0
			625	515	11	44	55	
29	S	1	Total	C	Mg	N	O	0
			625	515	11	44	55	
29	Y	1	Total	C	Mg	N	O	0
			570	480	9	36	45	
29	Y	1	Total	C	Mg	N	O	0
			570	480	9	36	45	
29	Y	1	Total	C	Mg	N	O	0
			570	480	9	36	45	
29	Y	1	Total	C	Mg	N	O	0
			570	480	9	36	45	
29	Y	1	Total	C	Mg	N	O	0
			570	480	9	36	45	

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Mol	Chain	Residues	Atoms					AltConf
29	Y	1	Total	C	Mg	N	O	0
			570	480	9	36	45	
29	Y	1	Total	C	Mg	N	O	0
			570	480	9	36	45	
29	Y	1	Total	C	Mg	N	O	0
			570	480	9	36	45	
29	a	1	Total	C	Mg	N	O	0
			239	199	4	16	20	
29	a	1	Total	C	Mg	N	O	0
			239	199	4	16	20	
29	a	1	Total	C	Mg	N	O	0
			239	199	4	16	20	
29	a	1	Total	C	Mg	N	O	0
			239	199	4	16	20	
29	b	1	Total	C	Mg	N	O	0
			1040	880	16	64	80	
29	b	1	Total	C	Mg	N	O	0
			1040	880	16	64	80	
29	b	1	Total	C	Mg	N	O	0
			1040	880	16	64	80	
29	b	1	Total	C	Mg	N	O	0
			1040	880	16	64	80	
29	b	1	Total	C	Mg	N	O	0
			1040	880	16	64	80	
29	b	1	Total	C	Mg	N	O	0
			1040	880	16	64	80	
29	b	1	Total	C	Mg	N	O	0
			1040	880	16	64	80	
29	b	1	Total	C	Mg	N	O	0
			1040	880	16	64	80	
29	b	1	Total	C	Mg	N	O	0
			1040	880	16	64	80	
29	b	1	Total	C	Mg	N	O	0
			1040	880	16	64	80	
29	b	1	Total	C	Mg	N	O	0
			1040	880	16	64	80	
29	b	1	Total	C	Mg	N	O	0
			1040	880	16	64	80	
29	b	1	Total	C	Mg	N	O	0
			1040	880	16	64	80	

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Mol	Chain	Residues	Atoms					AltConf
29	b	1	Total	C	Mg	N	O	0
			1040	880	16	64	80	
29	b	1	Total	C	Mg	N	O	0
			1040	880	16	64	80	
29	c	1	Total	C	Mg	N	O	0
			845	715	13	52	65	
29	c	1	Total	C	Mg	N	O	0
			845	715	13	52	65	
29	c	1	Total	C	Mg	N	O	0
			845	715	13	52	65	
29	c	1	Total	C	Mg	N	O	0
			845	715	13	52	65	
29	c	1	Total	C	Mg	N	O	0
			845	715	13	52	65	
29	c	1	Total	C	Mg	N	O	0
			845	715	13	52	65	
29	c	1	Total	C	Mg	N	O	0
			845	715	13	52	65	
29	c	1	Total	C	Mg	N	O	0
			845	715	13	52	65	
29	c	1	Total	C	Mg	N	O	0
			845	715	13	52	65	
29	c	1	Total	C	Mg	N	O	0
			845	715	13	52	65	
29	c	1	Total	C	Mg	N	O	0
			845	715	13	52	65	
29	d	1	Total	C	Mg	N	O	0
			130	110	2	8	10	
29	d	1	Total	C	Mg	N	O	0
			130	110	2	8	10	
29	n	1	Total	C	Mg	N	O	0
			468	388	8	32	40	
29	n	1	Total	C	Mg	N	O	0
			468	388	8	32	40	
29	n	1	Total	C	Mg	N	O	0
			468	388	8	32	40	
29	n	1	Total	C	Mg	N	O	0
			468	388	8	32	40	

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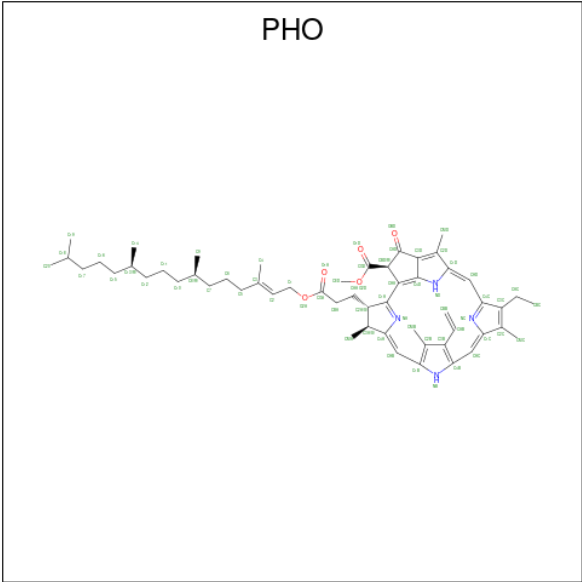
Mol	Chain	Residues	Atoms					AltConf
29	n	1	Total 468	C 388	Mg 8	N 32	O 40	0
29	n	1	Total 468	C 388	Mg 8	N 32	O 40	0
29	n	1	Total 468	C 388	Mg 8	N 32	O 40	0
29	n	1	Total 468	C 388	Mg 8	N 32	O 40	0
29	g	1	Total 446	C 368	Mg 8	N 32	O 38	0
29	g	1	Total 446	C 368	Mg 8	N 32	O 38	0
29	g	1	Total 446	C 368	Mg 8	N 32	O 38	0
29	g	1	Total 446	C 368	Mg 8	N 32	O 38	0
29	g	1	Total 446	C 368	Mg 8	N 32	O 38	0
29	g	1	Total 446	C 368	Mg 8	N 32	O 38	0
29	g	1	Total 446	C 368	Mg 8	N 32	O 38	0
29	g	1	Total 446	C 368	Mg 8	N 32	O 38	0
29	g	1	Total 446	C 368	Mg 8	N 32	O 38	0
29	r	1	Total 501	C 411	Mg 9	N 36	O 45	0
29	r	1	Total 501	C 411	Mg 9	N 36	O 45	0
29	r	1	Total 501	C 411	Mg 9	N 36	O 45	0
29	r	1	Total 501	C 411	Mg 9	N 36	O 45	0
29	r	1	Total 501	C 411	Mg 9	N 36	O 45	0
29	r	1	Total 501	C 411	Mg 9	N 36	O 45	0
29	r	1	Total 501	C 411	Mg 9	N 36	O 45	0
29	r	1	Total 501	C 411	Mg 9	N 36	O 45	0
29	r	1	Total 501	C 411	Mg 9	N 36	O 45	0

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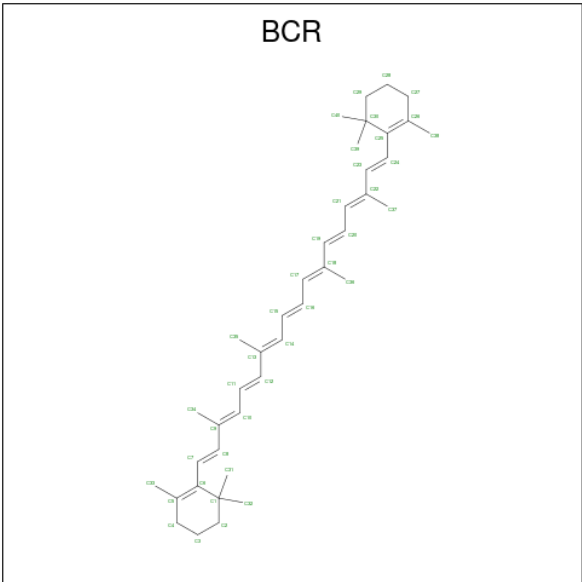
Mol	Chain	Residues	Atoms					AltConf
29	s	1	Total 625	C 515	Mg 11	N 44	O 55	0
29	s	1	Total 625	C 515	Mg 11	N 44	O 55	0
29	s	1	Total 625	C 515	Mg 11	N 44	O 55	0
29	s	1	Total 625	C 515	Mg 11	N 44	O 55	0
29	s	1	Total 625	C 515	Mg 11	N 44	O 55	0
29	s	1	Total 625	C 515	Mg 11	N 44	O 55	0
29	s	1	Total 625	C 515	Mg 11	N 44	O 55	0
29	s	1	Total 625	C 515	Mg 11	N 44	O 55	0
29	s	1	Total 625	C 515	Mg 11	N 44	O 55	0
29	s	1	Total 625	C 515	Mg 11	N 44	O 55	0
29	s	1	Total 625	C 515	Mg 11	N 44	O 55	0
29	y	1	Total 570	C 480	Mg 9	N 36	O 45	0
29	y	1	Total 570	C 480	Mg 9	N 36	O 45	0
29	y	1	Total 570	C 480	Mg 9	N 36	O 45	0
29	y	1	Total 570	C 480	Mg 9	N 36	O 45	0
29	y	1	Total 570	C 480	Mg 9	N 36	O 45	0
29	y	1	Total 570	C 480	Mg 9	N 36	O 45	0
29	y	1	Total 570	C 480	Mg 9	N 36	O 45	0
29	y	1	Total 570	C 480	Mg 9	N 36	O 45	0
29	y	1	Total 570	C 480	Mg 9	N 36	O 45	0

- Molecule 30 is PHEOPHYTIN A (three-letter code: PHO) (formula: C₅₅H₇₄N₄O₅).



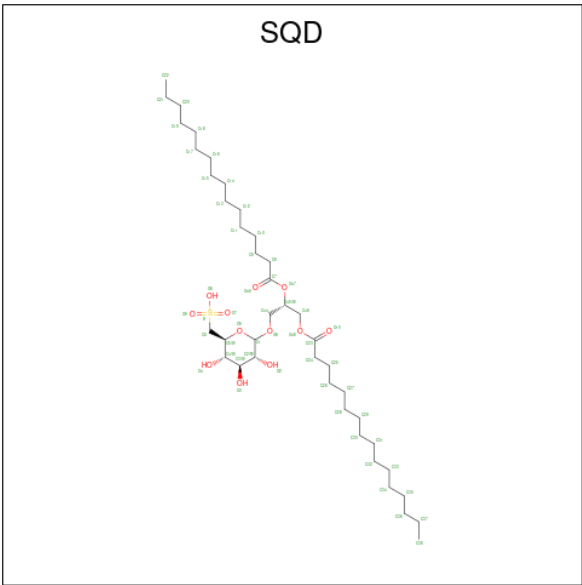
Mol	Chain	Residues	Atoms				AltConf
30	A	1	Total	C	N	O	0
			128	110	8	10	
30	A	1	Total	C	N	O	0
			128	110	8	10	
30	a	1	Total	C	N	O	0
			128	110	8	10	
30	a	1	Total	C	N	O	0
			128	110	8	10	

- Molecule 31 is BETA-CAROTENE (three-letter code: BCR) (formula: C₄₀H₅₆).



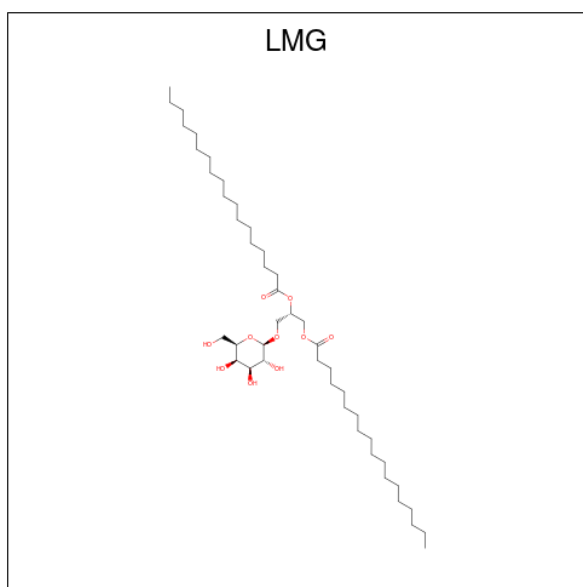
Mol	Chain	Residues	Atoms	AltConf
31	A	1	Total C 40 40	0
31	B	1	Total C 80 80	0
31	B	1	Total C 80 80	0
31	C	1	Total C 160 160	0
31	C	1	Total C 160 160	0
31	C	1	Total C 160 160	0
31	C	1	Total C 160 160	0
31	D	1	Total C 40 40	0
31	a	1	Total C 40 40	0
31	b	1	Total C 80 80	0
31	b	1	Total C 80 80	0
31	c	1	Total C 160 160	0
31	c	1	Total C 160 160	0
31	c	1	Total C 160 160	0
31	c	1	Total C 160 160	0
31	d	1	Total C 40 40	0

- Molecule 32 is 1,2-DI-O-ACYL-3-O-[6-DEOXY-6-SULFO-ALPHA-D-GLUCOPYRANOSYL]-SN-GLYCEROL (three-letter code: SQD) (formula: C₄₁H₇₈O₁₂S).



Mol	Chain	Residues	Atoms				AltConf
32	A	1	Total	C	O	S	0
			51	38	12	1	
32	B	1	Total	C	O	S	0
			54	41	12	1	
32	C	1	Total	C	O	S	0
			54	41	12	1	
32	a	1	Total	C	O	S	0
			51	38	12	1	
32	b	1	Total	C	O	S	0
			54	41	12	1	
32	c	1	Total	C	O	S	0
			54	41	12	1	

- Molecule 33 is 1,2-DISTEAROYL-MONOGALACTOSYL-DIGLYCERIDE (three-letter code: LMG) (formula: C₄₅H₈₆O₁₀).

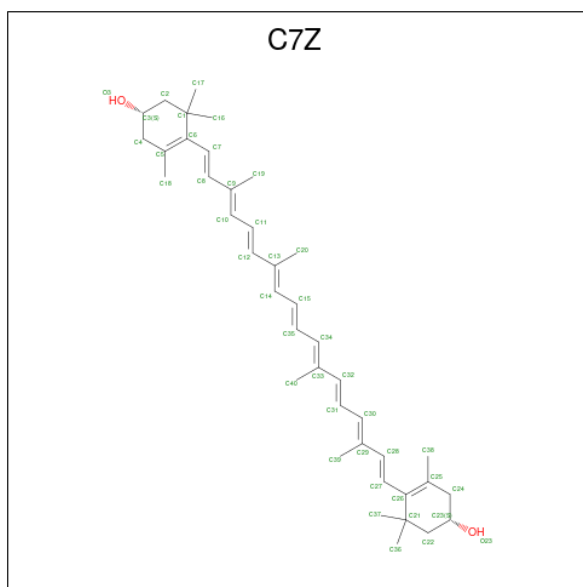


Mol	Chain	Residues	Atoms			AltConf
33	A	1	Total	C	O	0
			48	38	10	
33	B	1	Total	C	O	0
			44	34	10	
33	C	1	Total	C	O	0
			51	41	10	
33	D	1	Total	C	O	0
			46	36	10	
33	H	1	Total	C	O	0
			48	38	10	
33	J	1	Total	C	O	0
			45	35	10	
33	a	1	Total	C	O	0
			48	38	10	
33	b	1	Total	C	O	0
			44	34	10	
33	c	1	Total	C	O	0
			51	41	10	
33	d	1	Total	C	O	0
			46	36	10	
33	h	1	Total	C	O	0
			48	38	10	
33	j	1	Total	C	O	0
			45	35	10	

- Molecule 34 is SODIUM ION (three-letter code: NA) (formula: Na).

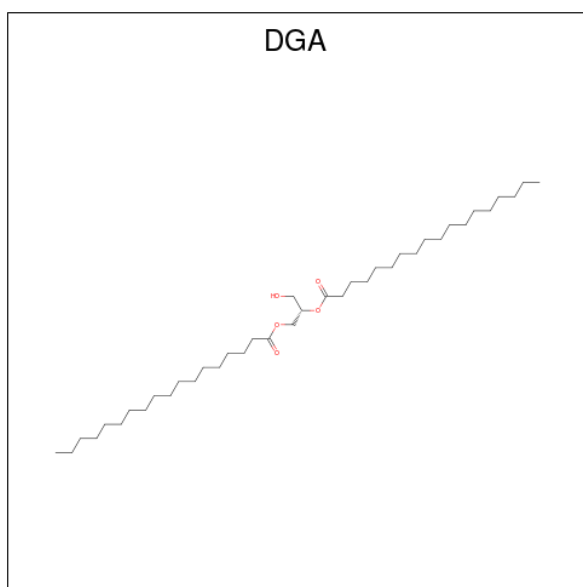
Mol	Chain	Residues	Atoms		AltConf
34	A	1	Total	Na	0
			1	1	
34	a	1	Total	Na	0
			1	1	

- Molecule 35 is (1 {S})-3,5,5-trimethyl-4-[(1 {E},3 {E},5 {E},7 {E},9 {E},11 {E},13 {E},15 {E},17 {E})-3,7,12,16-tetramethyl-18-[(4 {S})-2,6,6-trimethyl-4-oxidanyl-cyclohexen-1-yl]octadeca-1,3,5,7,9,11,13,15,17-nonaenyl]cyclohex-3-en-1-ol (three-letter code: C7Z) (formula: $C_{40}H_{56}O_2$).



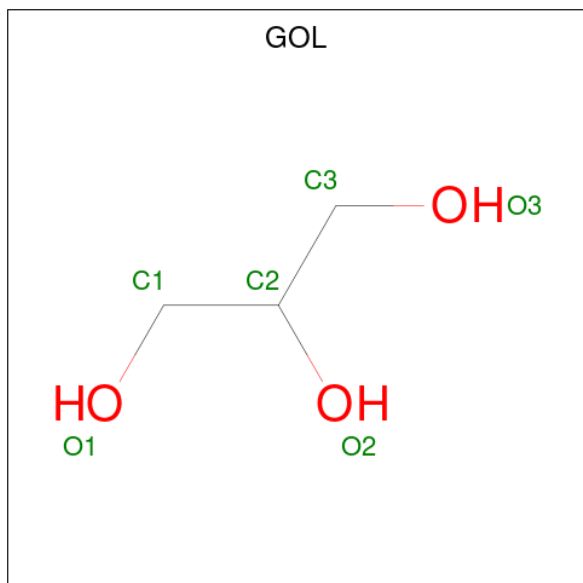
Mol	Chain	Residues	Atoms			AltConf
35	B	1	Total	C	O	0
			42	40	2	
35	b	1	Total	C	O	0
			42	40	2	

- Molecule 36 is DIACYL GLYCEROL (three-letter code: DGA) (formula: $C_{39}H_{76}O_5$).



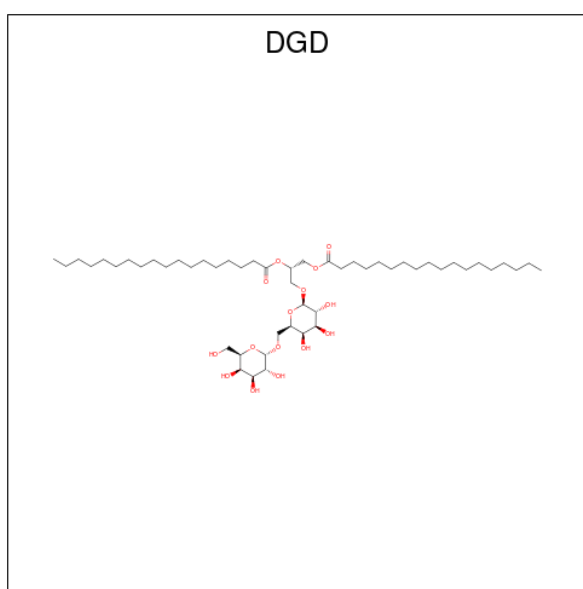
Mol	Chain	Residues	Atoms			AltConf
36	B	1	Total	C	O	0
			44	39	5	
36	C	1	Total	C	O	0
			44	39	5	
36	b	1	Total	C	O	0
			44	39	5	
36	c	1	Total	C	O	0
			44	39	5	

- Molecule 37 is GLYCEROL (three-letter code: GOL) (formula: $C_3H_8O_3$).



Mol	Chain	Residues	Atoms			AltConf
37	B	1	Total	C	O	0
			6	3	3	
37	b	1	Total	C	O	0
			12	6	6	
37	b	1	Total	C	O	0
			12	6	6	
37	y	1	Total	C	O	0
			6	3	3	

- Molecule 38 is DIGALACTOSYL DIACYL GLYCEROL (DGDG) (three-letter code: DGD) (formula: $C_{51}H_{96}O_{15}$).



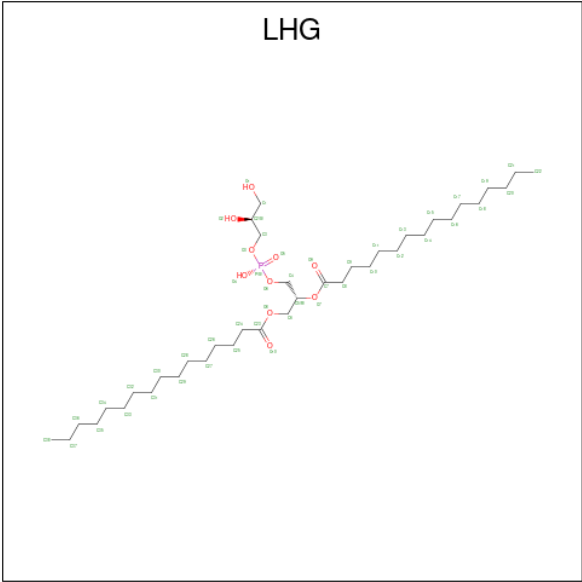
Mol	Chain	Residues	Atoms			AltConf
38	C	1	Total	C	O	0
			242	182	60	
38	C	1	Total	C	O	0
			242	182	60	
38	C	1	Total	C	O	0
			242	182	60	
38	C	1	Total	C	O	0
			242	182	60	
38	c	1	Total	C	O	0
			242	182	60	
38	c	1	Total	C	O	0
			242	182	60	
38	c	1	Total	C	O	0
			242	182	60	

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Mol	Chain	Residues	Atoms			AltConf
			Total	C	O	
38	c	1	242	182	60	0

- Molecule 39 is 1,2-DIPALMITOYL-PHOSPHATIDYL-GLYCEROLE (three-letter code: LHG) (formula: C₃₈H₇₅O₁₀P).



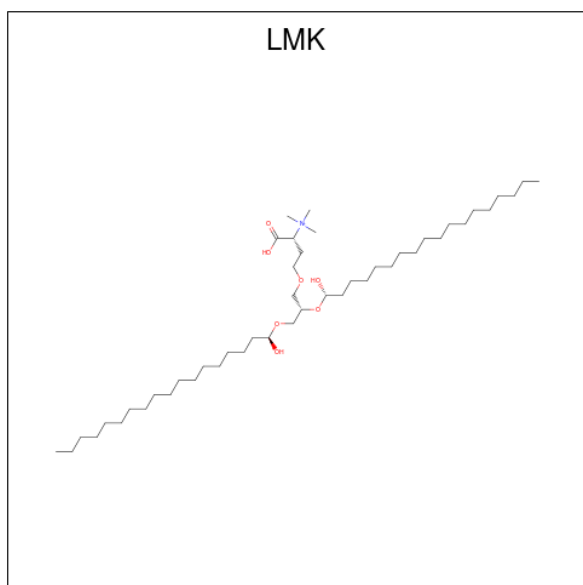
Mol	Chain	Residues	Atoms				AltConf
			Total	C	O	P	
39	C	1	47	36	10	1	0
39	D	1	132	99	30	3	0
39	D	1	132	99	30	3	0
39	D	1	132	99	30	3	0
39	L	1	49	38	10	1	0
39	N	1	49	38	10	1	0
39	G	1	49	38	10	1	0
39	S	1	45	34	10	1	0
39	Y	1	49	38	10	1	0
39	c	1	47	36	10	1	0

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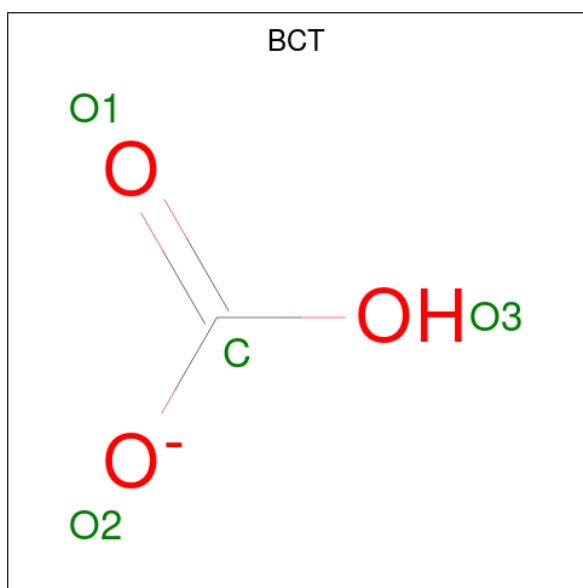
Mol	Chain	Residues	Atoms				AltConf
39	d	1	Total	C	O	P	0
			132	99	30	3	
39	d	1	Total	C	O	P	0
			132	99	30	3	
39	d	1	Total	C	O	P	0
			132	99	30	3	
39	l	1	Total	C	O	P	0
			49	38	10	1	
39	n	1	Total	C	O	P	0
			49	38	10	1	
39	g	1	Total	C	O	P	0
			49	38	10	1	
39	s	1	Total	C	O	P	0
			45	34	10	1	
39	y	1	Total	C	O	P	0
			49	38	10	1	

- Molecule 40 is trimethyl-[(2 {R})-1-oxidanyl-1-oxidanylidene-4-[(2 {S})-2-[(1 {S})-1-oxidanonyloctadecoxy]-3-[(1 {R})-1-oxidanyloctadecoxy]propoxy]butan-2-yl]azanium (three-letter code: LMK) (formula: C₄₆H₉₄NO₇).



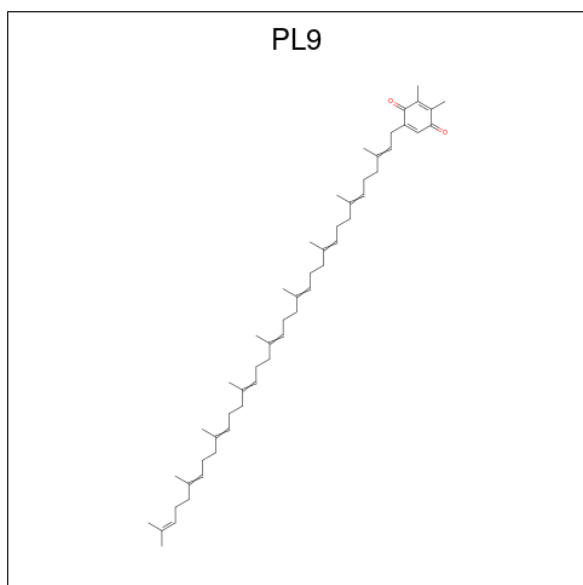
Mol	Chain	Residues	Atoms				AltConf
40	C	1	Total	C	N	O	0
			40	32	1	7	
40	c	1	Total	C	N	O	0
			40	32	1	7	

- Molecule 41 is BICARBONATE ION (three-letter code: BCT) (formula: CHO_3).



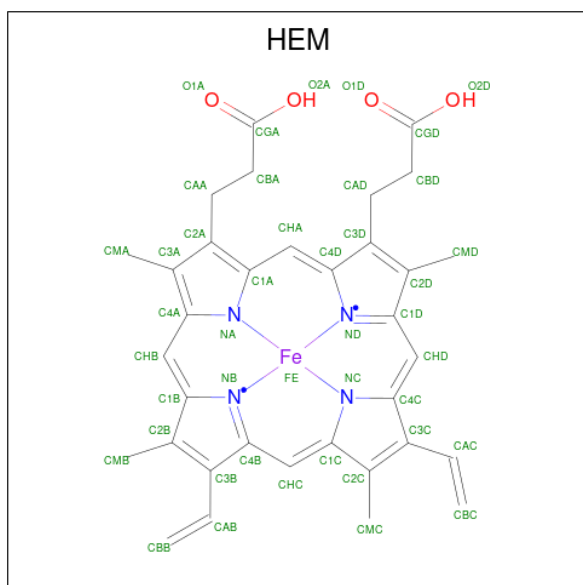
Mol	Chain	Residues	Atoms			AltConf
41	D	1	Total	C	O	0
			4	1	3	
41	d	1	Total	C	O	0
			4	1	3	

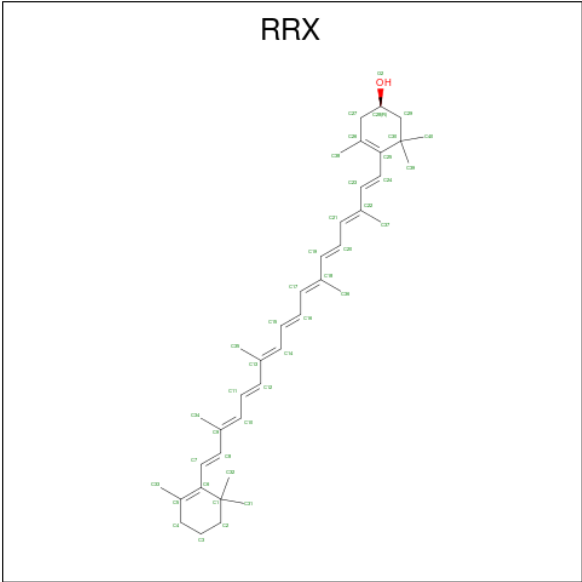
- Molecule 42 is 2,3-DIMETHYL-5-(3,7,11,15,19,23,27,31,35-NONAMETHYL-2,6,10,14,18,22,26,30,34-HEXATRIACONTANONAENYL-2,5-CYCLOHEXADIENE-1,4-DIONE-2,3-DIMETHYL-5-SOLANESYL-1,4-BENZOQUINONE (three-letter code: PL9) (formula: $\text{C}_{53}\text{H}_{80}\text{O}_2$).



Mol	Chain	Residues	Atoms			AltConf
42	D	1	Total	C	O	0
			55	53	2	
42	d	1	Total	C	O	0
			55	53	2	

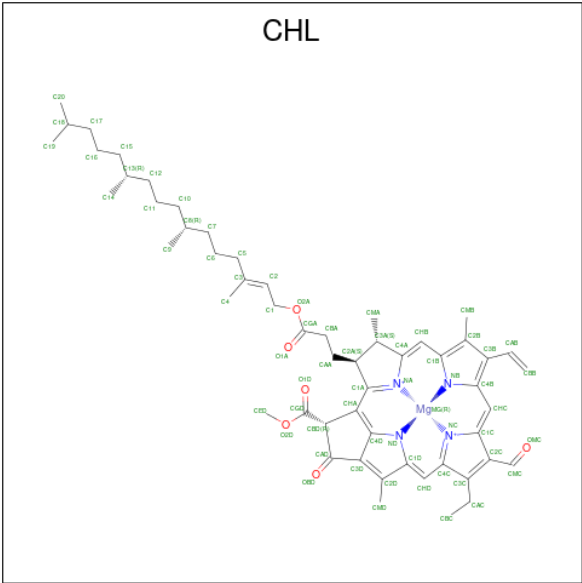
- Molecule 43 is PROTOPORPHYRIN IX CONTAINING FE (three-letter code: HEM) (formula: $C_{34}H_{32}FeN_4O_4$).





Mol	Chain	Residues	Atoms			AltConf
44	H	1	Total	C	O	0
			41	40	1	
44	h	1	Total	C	O	0
			41	40	1	

- Molecule 45 is CHLOROPHYLL B (three-letter code: CHL) (formula: C₅₅H₇₀MgN₄O₆).



Mol	Chain	Residues	Atoms					AltConf
45	N	1	Total	C	Mg	N	O	0
			380	314	6	24	36	
45	N	1	Total	C	Mg	N	O	0
			380	314	6	24	36	

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Mol	Chain	Residues	Atoms					AltConf
45	N	1	Total 380	C 314	Mg 6	N 24	O 36	0
45	N	1	Total 380	C 314	Mg 6	N 24	O 36	0
45	N	1	Total 380	C 314	Mg 6	N 24	O 36	0
45	N	1	Total 380	C 314	Mg 6	N 24	O 36	0
45	G	1	Total 324	C 260	Mg 6	N 24	O 34	0
45	G	1	Total 324	C 260	Mg 6	N 24	O 34	0
45	G	1	Total 324	C 260	Mg 6	N 24	O 34	0
45	G	1	Total 324	C 260	Mg 6	N 24	O 34	0
45	G	1	Total 324	C 260	Mg 6	N 24	O 34	0
45	G	1	Total 324	C 260	Mg 6	N 24	O 34	0
45	R	1	Total 94	C 74	Mg 2	N 8	O 10	0
45	R	1	Total 94	C 74	Mg 2	N 8	O 10	0
45	S	1	Total 194	C 154	Mg 4	N 16	O 20	0
45	S	1	Total 194	C 154	Mg 4	N 16	O 20	0
45	S	1	Total 194	C 154	Mg 4	N 16	O 20	0
45	S	1	Total 194	C 154	Mg 4	N 16	O 20	0
45	Y	1	Total 310	C 255	Mg 5	N 20	O 30	0
45	Y	1	Total 310	C 255	Mg 5	N 20	O 30	0
45	Y	1	Total 310	C 255	Mg 5	N 20	O 30	0
45	Y	1	Total 310	C 255	Mg 5	N 20	O 30	0
45	Y	1	Total 310	C 255	Mg 5	N 20	O 30	0

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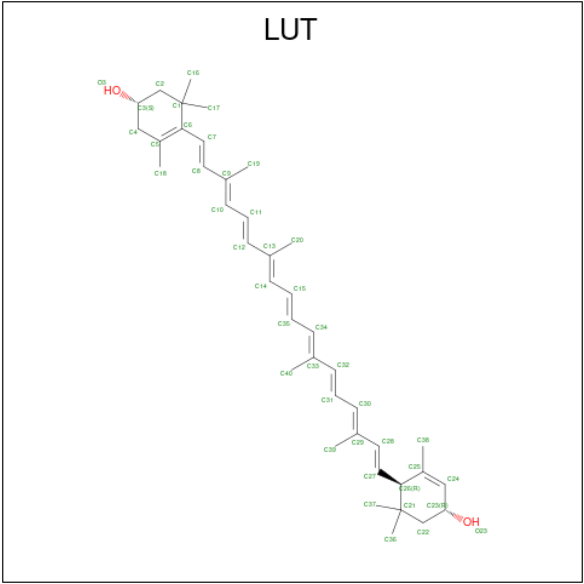
Mol	Chain	Residues	Atoms					AltConf
45	n	1	Total 380	C 314	Mg 6	N 24	O 36	0
45	n	1	Total 380	C 314	Mg 6	N 24	O 36	0
45	n	1	Total 380	C 314	Mg 6	N 24	O 36	0
45	n	1	Total 380	C 314	Mg 6	N 24	O 36	0
45	n	1	Total 380	C 314	Mg 6	N 24	O 36	0
45	n	1	Total 380	C 314	Mg 6	N 24	O 36	0
45	g	1	Total 324	C 260	Mg 6	N 24	O 34	0
45	g	1	Total 324	C 260	Mg 6	N 24	O 34	0
45	g	1	Total 324	C 260	Mg 6	N 24	O 34	0
45	g	1	Total 324	C 260	Mg 6	N 24	O 34	0
45	g	1	Total 324	C 260	Mg 6	N 24	O 34	0
45	g	1	Total 324	C 260	Mg 6	N 24	O 34	0
45	g	1	Total 324	C 260	Mg 6	N 24	O 34	0
45	r	1	Total 94	C 74	Mg 2	N 8	O 10	0
45	r	1	Total 94	C 74	Mg 2	N 8	O 10	0
45	s	1	Total 194	C 154	Mg 4	N 16	O 20	0
45	s	1	Total 194	C 154	Mg 4	N 16	O 20	0
45	s	1	Total 194	C 154	Mg 4	N 16	O 20	0
45	s	1	Total 194	C 154	Mg 4	N 16	O 20	0
45	y	1	Total 310	C 255	Mg 5	N 20	O 30	0
45	y	1	Total 310	C 255	Mg 5	N 20	O 30	0
45	y	1	Total 310	C 255	Mg 5	N 20	O 30	0

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Mol	Chain	Residues	Atoms					AltConf
45	y	1	Total	C	Mg	N	O	0
			310	255	5	20	30	
45	y	1	Total	C	Mg	N	O	0
			310	255	5	20	30	

- Molecule 46 is (3R,3'R,6S)-4,5-DIDEHYDRO-5,6-DIHYDRO-BETA,BETA-CAROTENE-3,3'-DIOL (three-letter code: LUT) (formula: C₄₀H₅₆O₂).



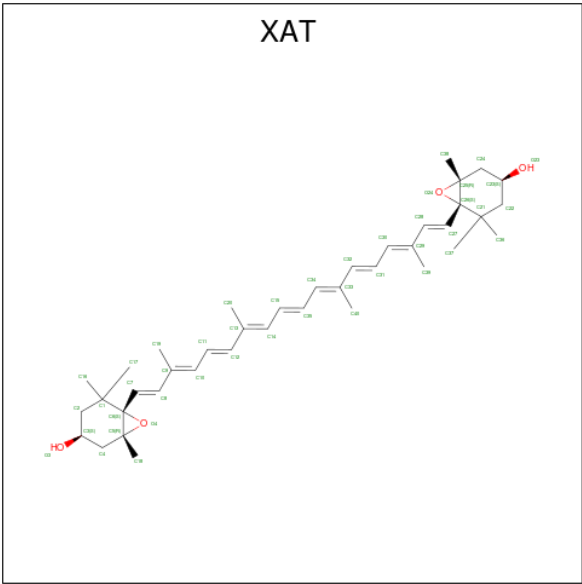
Mol	Chain	Residues	Atoms			AltConf
46	N	1	Total	C	O	0
			84	80	4	
46	N	1	Total	C	O	0
			84	80	4	
46	G	1	Total	C	O	0
			84	80	4	
46	G	1	Total	C	O	0
			84	80	4	
46	R	1	Total	C	O	0
			42	40	2	
46	S	1	Total	C	O	0
			84	80	4	
46	S	1	Total	C	O	0
			84	80	4	
46	Y	1	Total	C	O	0
			84	80	4	
46	Y	1	Total	C	O	0
			84	80	4	

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Mol	Chain	Residues	Atoms			AltConf
46	n	1	Total	C	O	0
			84	80	4	
46	n	1	Total	C	O	0
			84	80	4	
46	g	1	Total	C	O	0
			84	80	4	
46	g	1	Total	C	O	0
			84	80	4	
46	r	1	Total	C	O	0
			42	40	2	
46	s	1	Total	C	O	0
			84	80	4	
46	s	1	Total	C	O	0
			84	80	4	
46	y	1	Total	C	O	0
			84	80	4	
46	y	1	Total	C	O	0
			84	80	4	

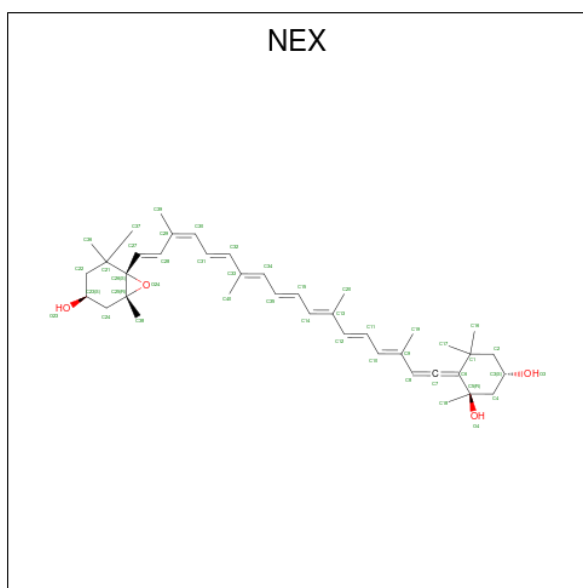
- Molecule 47 is (3S,5R,6S,3'S,5'R,6'S)-5,6,5',6'-DIEPOXY-5,6,5',6'- TETRAHYDRO-BETA ,BETA-CAROTENE-3,3'-DIOL (three-letter code: XAT) (formula: C₄₀H₅₆O₄).



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Mol	Chain	Residues	Atoms			AltConf
47	R	1	Total	C	O	0
			44	40	4	
47	Y	1	Total	C	O	0
			44	40	4	
47	n	1	Total	C	O	0
			44	40	4	
47	g	1	Total	C	O	0
			44	40	4	
47	r	1	Total	C	O	0
			44	40	4	
47	y	1	Total	C	O	0
			44	40	4	

- Molecule 48 is (1R,3R)-6-[(3E,5E,7E,9E,11E,13E,15E,17E)-18-[(1S,4R,6R)-4-HYDROXY-2,2,6-TRIMETHYL-7-OXABICYCLO[4.1.0]HEPT-1-YL]-3,7,12,16-TETRAMETHYLOCTA-DECA-1,3,5,7,9,11,13,15,17-NONAENYLIDENE}-1,5,5-TRIMETHYLCYCLOHEXANE-1,3-DIOL (three-letter code: NEX) (formula: C₄₀H₅₆O₄).



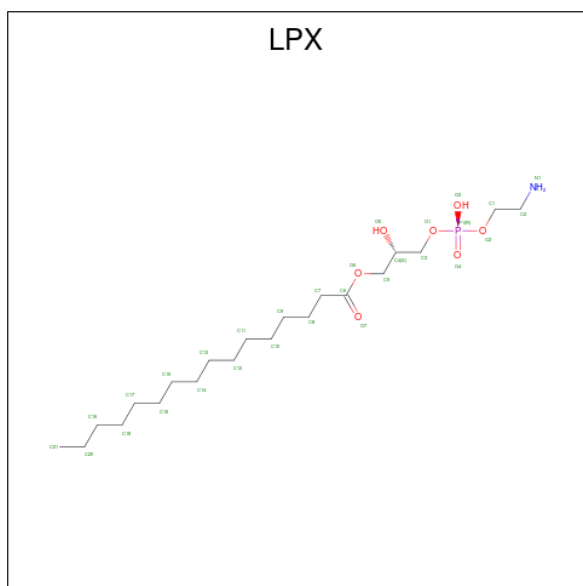
Mol	Chain	Residues	Atoms			AltConf
48	N	1	Total	C	O	0
			44	40	4	
48	G	1	Total	C	O	0
			44	40	4	
48	R	1	Total	C	O	0
			44	40	4	
48	S	1	Total	C	O	0
			44	40	4	

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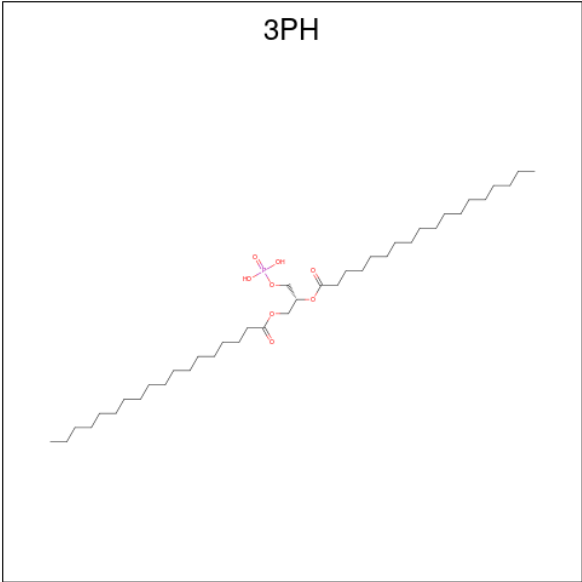
Mol	Chain	Residues	Atoms			AltConf
48	Y	1	Total	C	O	0
			44	40	4	
48	n	1	Total	C	O	0
			44	40	4	
48	g	1	Total	C	O	0
			44	40	4	
48	r	1	Total	C	O	0
			44	40	4	
48	s	1	Total	C	O	0
			44	40	4	
48	y	1	Total	C	O	0
			44	40	4	

- Molecule 49 is (2S)-3-{[(R)-(2-aminoethoxy)(hydroxy)phosphoryl]oxy}-2-hydroxypropyl hexadecanoate (three-letter code: LPX) (formula: C₂₁H₄₄NO₇P).



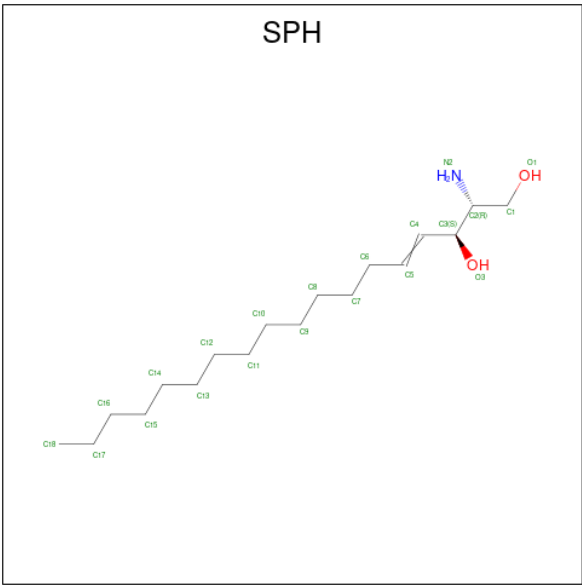
Mol	Chain	Residues	Atoms					AltConf
49	S	1	Total	C	N	O	P	0
			30	21	1	7	1	
49	s	1	Total	C	N	O	P	0
			30	21	1	7	1	

- Molecule 50 is 1,2-DIACYL-GLYCEROL-3-SN-PHOSPHATE (three-letter code: 3PH) (formula: C₃₉H₇₇O₈P).



Mol	Chain	Residues	Atoms				AltConf
50	S	1	Total	C	O	P	0
			48	39	8	1	
50	i	1	Total	C	O	P	0
			48	39	8	1	
50	s	1	Total	C	O	P	0
			48	39	8	1	

- Molecule 51 is SPHINGOSINE (three-letter code: SPH) (formula: C₁₈H₃₇NO₂).



Mol	Chain	Residues	Atoms				AltConf
51	Y	1	Total	C	N	O	0
			21	18	1	2	

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Mol	Chain	Residues	Atoms				AltConf
51	y	1	Total	C	N	O	0
			21	18	1	2	

- Molecule 52 is water.

Mol	Chain	Residues	Atoms		AltConf
52	A	43	Total	O	0
			43	43	
52	B	57	Total	O	0
			57	57	
52	C	55	Total	O	0
			55	55	
52	D	42	Total	O	0
			42	42	
52	E	7	Total	O	0
			7	7	
52	H	8	Total	O	0
			8	8	
52	I	4	Total	O	0
			4	4	
52	J	3	Total	O	0
			3	3	
52	K	2	Total	O	0
			2	2	
52	L	5	Total	O	0
			5	5	
52	M	4	Total	O	0
			4	4	
52	O	28	Total	O	0
			28	28	
52	P	10	Total	O	0
			10	10	
52	T	7	Total	O	0
			7	7	
52	W	5	Total	O	0
			5	5	
52	X	9	Total	O	0
			9	9	
52	Z	1	Total	O	0
			1	1	
52	N	5	Total	O	0
			5	5	

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Mol	Chain	Residues	Atoms		AltConf
52	G	4	Total 4	O 4	0
52	R	10	Total 10	O 10	0
52	S	8	Total 8	O 8	0
52	Y	14	Total 14	O 14	0
52	U	1	Total 1	O 1	0
52	a	59	Total 59	O 59	0
52	b	70	Total 70	O 70	0
52	v	3	Total 3	O 3	0
52	c	50	Total 50	O 50	0
52	d	41	Total 41	O 41	0
52	e	7	Total 7	O 7	0
52	f	1	Total 1	O 1	0
52	h	9	Total 9	O 9	0
52	i	3	Total 3	O 3	0
52	j	2	Total 2	O 2	0
52	k	1	Total 1	O 1	0
52	l	8	Total 8	O 8	0
52	m	4	Total 4	O 4	0
52	o	20	Total 20	O 20	0
52	p	15	Total 15	O 15	0
52	t	5	Total 5	O 5	0

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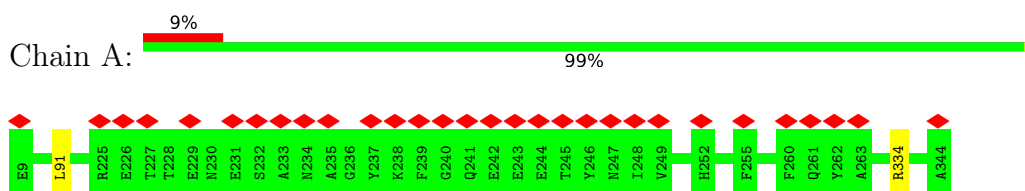
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Mol	Chain	Residues	Atoms		AltConf
52	w	3	Total 3	O 3	0
52	x	2	Total 2	O 2	0
52	n	13	Total 13	O 13	0
52	g	12	Total 12	O 12	0
52	r	15	Total 15	O 15	0
52	s	21	Total 21	O 21	0
52	y	22	Total 22	O 22	0

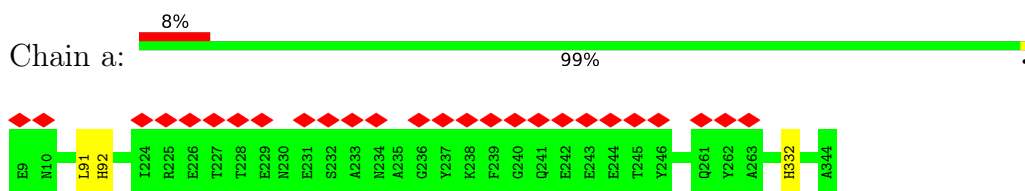
3 Residue-property plots [i](#)

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

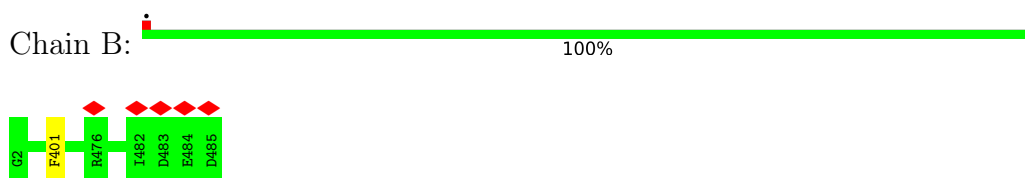
- Molecule 1: Photosystem II protein D1



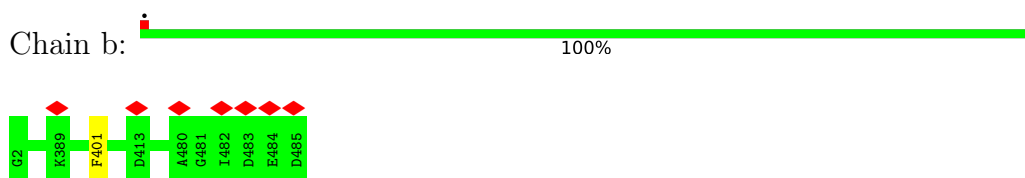
- Molecule 1: Photosystem II protein D1



- Molecule 2: Photosystem II CP47 reaction center protein



- Molecule 2: Photosystem II CP47 reaction center protein



- Molecule 3: Photosystem II reaction center protein Ycf12



There are no outlier residues recorded for this chain.

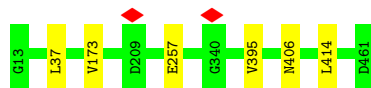
- Molecule 3: Photosystem II reaction center protein Ycf12

Chain v:  100%

There are no outlier residues recorded for this chain.

- Molecule 4: Photosystem II CP43 reaction center protein

Chain C:  99%



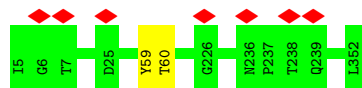
- Molecule 4: Photosystem II CP43 reaction center protein

Chain c:  99%



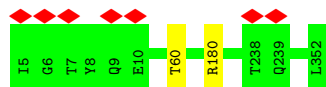
- Molecule 5: Photosystem II D2 protein

Chain D:  99%



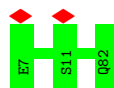
- Molecule 5: Photosystem II D2 protein

Chain d:  99%



- Molecule 6: Cytochrome b559 subunit alpha

Chain E:  100%



- Molecule 6: Cytochrome b559 subunit alpha

Chain e:  99%



- Molecule 7: Cytochrome b559 subunit beta

Chain F:  100%



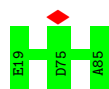
- Molecule 7: Cytochrome b559 subunit beta

Chain f:  100%



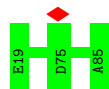
- Molecule 8: Photosystem II reaction center protein H

Chain H:  100%



- Molecule 8: Photosystem II reaction center protein H

Chain h:  100%



- Molecule 9: Photosystem II reaction center protein I

Chain I:  6% 97%



- Molecule 9: Photosystem II reaction center protein I

Chain i:  100%

There are no outlier residues recorded for this chain.

- Molecule 10: Photosystem II reaction center protein J

Chain J:  6% 100%



- Molecule 10: Photosystem II reaction center protein J

Chain j:  100%



- Molecule 11: Photosystem II reaction center protein K

Chain K:  100%

There are no outlier residues recorded for this chain.

- Molecule 11: Photosystem II reaction center protein K

Chain k:  100%

There are no outlier residues recorded for this chain.

- Molecule 12: Photosystem II reaction center protein L

Chain L:  100%

There are no outlier residues recorded for this chain.

- Molecule 12: Photosystem II reaction center protein L

Chain l:  100%



- Molecule 13: Photosystem II reaction center protein M

Chain M:  97%



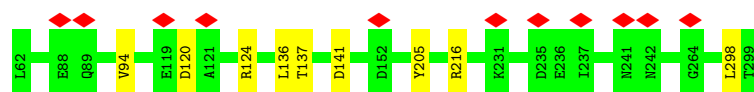
- Molecule 13: Photosystem II reaction center protein M

Chain m:  97%



- Molecule 14: PsbO

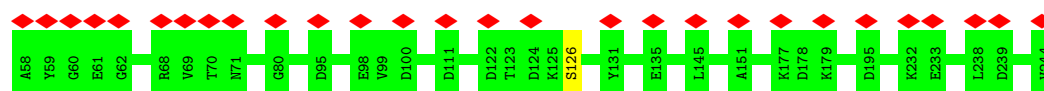
Chain O:  5% 96%



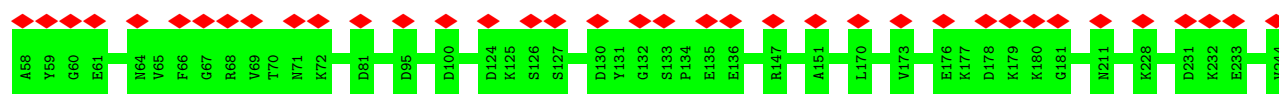
• Molecule 14: PsbO



• Molecule 15: PsbP



• Molecule 15: PsbP



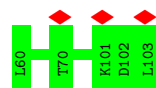
• Molecule 16: Photosystem II reaction center protein T



• Molecule 16: Photosystem II reaction center protein T

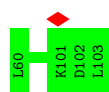


• Molecule 17: PSII 6.1 kDa protein



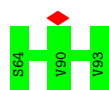
• Molecule 17: PSII 6.1 kDa protein

Chain w:  100%



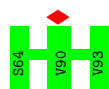
- Molecule 18: Hypothetical protein

Chain X:  100%



- Molecule 18: Hypothetical protein

Chain x:  100%



- Molecule 19: Photosystem II reaction center protein Z

Chain Z:  98%



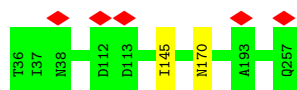
- Molecule 19: Photosystem II reaction center protein Z

Chain z:  100%

There are no outlier residues recorded for this chain.

- Molecule 20: LHCII M3

Chain N:  99%



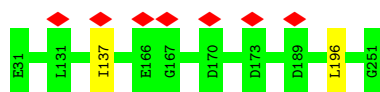
- Molecule 20: LHCII M3

Chain n:  100%



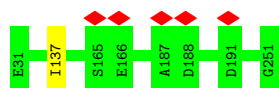
- Molecule 21: Chlorophyll a-b binding protein, chloroplastic

Chain G:  99%



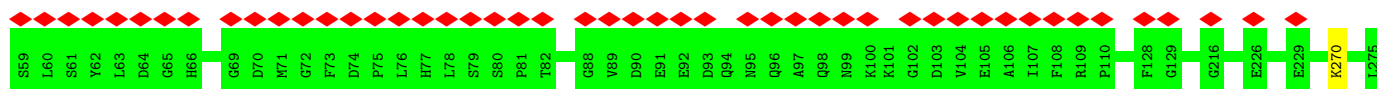
- Molecule 21: Chlorophyll a-b binding protein, chloroplastic

Chain g:  100%



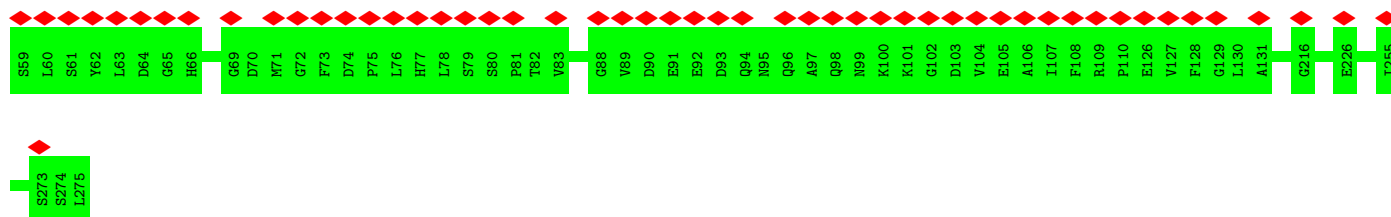
- Molecule 22: CP29

Chain R:  24%  100%



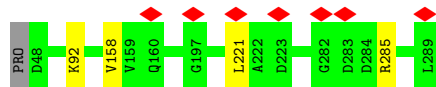
- Molecule 22: CP29

Chain r:  26%  100%



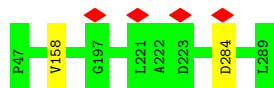
- Molecule 23: CP26

Chain S:  98%



- Molecule 23: CP26

Chain s:  99%



- Molecule 24: LHCII M1

Chain Y:  99%



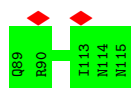
- Molecule 24: LHCII M1

Chain y:  99%



- Molecule 25: PsbU

Chain U:  7%  100%



- Molecule 25: PsbU

Chain u:  100%

There are no outlier residues recorded for this chain.

4 Experimental information

Property	Value	Source
EM reconstruction method	SINGLE PARTICLE	Depositor
Imposed symmetry	POINT, Not provided	
Number of particles used	23014	Depositor
Resolution determination method	FSC 0.143 CUT-OFF	Depositor
CTF correction method	NONE	Depositor
Microscope	FEI TITAN KRIOS	Depositor
Voltage (kV)	300	Depositor
Electron dose ($e^-/\text{\AA}^2$)	51.81	Depositor
Minimum defocus (nm)	Not provided	
Maximum defocus (nm)	Not provided	
Magnification	Not provided	
Image detector	GATAN K3 BIOQUANTUM (6k x 4k)	Depositor
Maximum map value	40.024	Depositor
Minimum map value	-28.419	Depositor
Average map value	0.000	Depositor
Map value standard deviation	1.000	Depositor
Recommended contour level	2.5	Depositor
Map size (\AA)	448.0, 448.0, 448.0	wwPDB
Map dimensions	500, 500, 500	wwPDB
Map angles ($^\circ$)	90.0, 90.0, 90.0	wwPDB
Pixel spacing (\AA)	0.896, 0.896, 0.896	Depositor

5 Model quality

5.1 Standard geometry

Bond lengths and bond angles in the following residue types are not validated in this section: OEX, HEM, LMG, 3PH, SPH, PHO, C7Z, CHL, RRX, BCT, XAT, PL9, LUT, CLA, LPX, DGA, NEX, LHG, DGD, NA, BCR, LMK, CL, SQD, FE2, GOL

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	0.29	0/2723	0.55	1/3715 (0.0%)
1	a	0.32	0/2723	0.57	1/3715 (0.0%)
2	B	0.30	0/3912	0.52	0/5327
2	b	0.30	0/3912	0.52	0/5327
3	V	0.25	0/228	0.57	0/311
3	v	0.24	0/228	0.61	0/311
4	C	0.29	0/3602	0.56	2/4913 (0.0%)
4	c	0.30	0/3602	0.58	1/4913 (0.0%)
5	D	0.30	0/2860	0.54	1/3899 (0.0%)
5	d	0.30	0/2860	0.53	0/3899
6	E	0.27	0/639	0.54	0/870
6	e	0.27	0/639	0.61	1/870 (0.1%)
7	F	0.27	0/259	0.54	0/351
7	f	0.26	0/259	0.49	0/351
8	H	0.27	0/513	0.53	0/703
8	h	0.29	0/513	0.52	0/703
9	I	0.30	0/287	0.50	0/386
9	i	0.29	0/287	0.51	0/386
10	J	0.25	0/272	0.46	0/369
10	j	0.25	0/272	0.55	0/369
11	K	0.32	0/308	0.52	0/423
11	k	0.34	0/308	0.58	0/423
12	L	0.27	0/321	0.45	0/435
12	l	0.27	0/321	0.47	0/435
13	M	0.26	0/246	0.48	0/335
13	m	0.25	0/246	0.46	0/335
14	O	0.31	0/1855	0.63	2/2505 (0.1%)
14	o	0.29	0/1855	0.59	1/2505 (0.0%)
15	P	0.28	0/1473	0.59	0/1988
15	p	0.27	0/1473	0.55	0/1988
16	T	0.29	0/254	0.50	0/342

Mol	Chain	Bond lengths		Bond angles	
		RMSZ	# Z >5	RMSZ	# Z >5
16	t	0.29	0/254	0.51	0/342
17	W	0.25	0/339	0.51	0/460
17	w	0.27	0/339	0.51	0/460
18	X	0.25	0/202	0.39	0/276
18	x	0.25	0/202	0.41	0/276
19	Z	0.27	0/469	0.44	0/641
19	z	0.26	0/469	0.47	0/641
20	N	0.27	0/1751	0.53	0/2386
20	n	0.27	0/1751	0.51	0/2386
21	G	0.28	0/1725	0.57	1/2348 (0.0%)
21	g	0.27	0/1725	0.50	0/2348
22	R	0.28	0/1561	0.56	0/2110
22	r	0.27	0/1561	0.56	0/2110
23	S	0.27	0/1895	0.52	1/2579 (0.0%)
23	s	0.27	0/1902	0.49	0/2587
24	Y	0.28	0/1715	0.57	1/2338 (0.0%)
24	y	0.28	0/1715	0.53	0/2338
25	U	0.29	0/224	0.70	0/298
25	u	0.35	0/224	0.67	0/298
All	All	0.29	0/59273	0.54	13/80624 (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
14	O	0	1

There are no bond length outliers.

All (13) bond angle outliers are listed below:

Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
24	Y	98	LEU	CA-CB-CG	8.32	134.44	115.30
21	G	196	LEU	CA-CB-CG	7.56	132.68	115.30
6	e	79	LEU	CA-CB-CG	7.15	131.74	115.30
4	C	37	LEU	CA-CB-CG	5.88	128.84	115.30
14	O	298	LEU	CA-CB-CG	5.84	128.74	115.30
1	a	91	LEU	CA-CB-CG	5.84	128.73	115.30
4	c	414	LEU	CA-CB-CG	5.78	128.59	115.30
23	S	221	LEU	CA-CB-CG	5.62	128.22	115.30
1	A	91	LEU	CA-CB-CG	5.51	127.98	115.30

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
14	o	141	ASP	CB-CG-OD2	5.21	122.99	118.30
14	O	141	ASP	CB-CG-OD2	5.18	122.96	118.30
5	D	59	TYR	C-N-CA	5.07	134.38	121.70
4	C	414	LEU	CA-CB-CG	5.06	126.94	115.30

There are no chirality outliers.

All (1) planarity outliers are listed below:

Mol	Chain	Res	Type	Group
14	O	136	LEU	Mainchain

5.2 Too-close contacts [i](#)

Due to software issues we are unable to calculate clashes - this section is therefore empty.

5.3 Torsion angles [i](#)

5.3.1 Protein backbone [i](#)

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

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

Mol	Chain	Analysed	Favoured	Allowed	Outliers	Percentiles	
1	A	335/336 (100%)	309 (92%)	26 (8%)	0	100	100
1	a	335/336 (100%)	310 (92%)	25 (8%)	0	100	100
2	B	482/484 (100%)	465 (96%)	16 (3%)	1 (0%)	47	76
2	b	482/484 (100%)	463 (96%)	18 (4%)	1 (0%)	47	76
3	V	30/32 (94%)	28 (93%)	2 (7%)	0	100	100
3	v	30/32 (94%)	28 (93%)	2 (7%)	0	100	100
4	C	447/449 (100%)	419 (94%)	25 (6%)	3 (1%)	22	50
4	c	447/449 (100%)	414 (93%)	29 (6%)	4 (1%)	17	44
5	D	346/348 (99%)	332 (96%)	13 (4%)	1 (0%)	41	70
5	d	346/348 (99%)	335 (97%)	10 (3%)	1 (0%)	41	70

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Mol	Chain	Analysed	Favoured	Allowed	Outliers	Percentiles	
6	E	74/76 (97%)	71 (96%)	3 (4%)	0	100	100
6	e	74/76 (97%)	70 (95%)	4 (5%)	0	100	100
7	F	29/31 (94%)	29 (100%)	0	0	100	100
7	f	29/31 (94%)	29 (100%)	0	0	100	100
8	H	65/67 (97%)	65 (100%)	0	0	100	100
8	h	65/67 (97%)	64 (98%)	1 (2%)	0	100	100
9	I	33/35 (94%)	33 (100%)	0	0	100	100
9	i	33/35 (94%)	33 (100%)	0	0	100	100
10	J	34/36 (94%)	34 (100%)	0	0	100	100
10	j	34/36 (94%)	34 (100%)	0	0	100	100
11	K	35/37 (95%)	35 (100%)	0	0	100	100
11	k	35/37 (95%)	35 (100%)	0	0	100	100
12	L	36/38 (95%)	35 (97%)	1 (3%)	0	100	100
12	l	36/38 (95%)	35 (97%)	1 (3%)	0	100	100
13	M	30/32 (94%)	29 (97%)	0	1 (3%)	4	11
13	m	30/32 (94%)	29 (97%)	0	1 (3%)	4	11
14	O	236/238 (99%)	208 (88%)	25 (11%)	3 (1%)	12	33
14	o	236/238 (99%)	214 (91%)	21 (9%)	1 (0%)	34	64
15	P	185/187 (99%)	169 (91%)	15 (8%)	1 (0%)	29	58
15	p	185/187 (99%)	172 (93%)	13 (7%)	0	100	100
16	T	28/30 (93%)	26 (93%)	1 (4%)	1 (4%)	3	10
16	t	28/30 (93%)	26 (93%)	1 (4%)	1 (4%)	3	10
17	W	42/44 (96%)	41 (98%)	1 (2%)	0	100	100
17	w	42/44 (96%)	41 (98%)	1 (2%)	0	100	100
18	X	28/30 (93%)	27 (96%)	1 (4%)	0	100	100
18	x	28/30 (93%)	28 (100%)	0	0	100	100
19	Z	59/61 (97%)	58 (98%)	1 (2%)	0	100	100
19	z	59/61 (97%)	58 (98%)	1 (2%)	0	100	100
20	N	220/222 (99%)	204 (93%)	15 (7%)	1 (0%)	29	58
20	n	220/222 (99%)	206 (94%)	13 (6%)	1 (0%)	29	58
21	G	219/221 (99%)	203 (93%)	15 (7%)	1 (0%)	29	58

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Mol	Chain	Analysed	Favoured	Allowed	Outliers	Percentiles	
21	g	219/221 (99%)	206 (94%)	12 (6%)	1 (0%)	29	58
22	R	198/202 (98%)	188 (95%)	10 (5%)	0	100	100
22	r	198/202 (98%)	185 (93%)	13 (7%)	0	100	100
23	S	240/243 (99%)	220 (92%)	18 (8%)	2 (1%)	19	47
23	s	239/243 (98%)	221 (92%)	16 (7%)	2 (1%)	19	47
24	Y	220/222 (99%)	210 (96%)	9 (4%)	1 (0%)	29	58
24	y	220/222 (99%)	211 (96%)	8 (4%)	1 (0%)	29	58
25	U	25/27 (93%)	25 (100%)	0	0	100	100
25	u	25/27 (93%)	24 (96%)	1 (4%)	0	100	100
All	All	7351/7456 (99%)	6934 (94%)	387 (5%)	30 (0%)	38	64

All (30) Ramachandran outliers are listed below:

Mol	Chain	Res	Type
4	C	257	GLU
4	C	395	VAL
14	O	94	VAL
15	P	126	SER
16	T	29	ILE
4	c	257	GLU
4	c	395	VAL
16	t	29	ILE
4	C	173	VAL
5	D	60	THR
14	O	205	TYR
20	N	145	ILE
4	c	173	VAL
5	d	60	THR
14	o	205	TYR
20	n	145	ILE
21	G	137	ILE
23	S	92	LYS
24	Y	146	ILE
21	g	137	ILE
2	B	401	PHE
13	M	3	VAL
2	b	401	PHE
13	m	3	VAL
24	y	146	ILE

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Mol	Chain	Res	Type
23	S	158	VAL
14	O	216	ARG
4	c	94	LEU
23	s	284	ASP
23	s	158	VAL

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	276/275 (100%)	275 (100%)	1 (0%)	91	96
1	a	276/275 (100%)	274 (99%)	2 (1%)	84	94
2	B	388/388 (100%)	388 (100%)	0	100	100
2	b	388/388 (100%)	388 (100%)	0	100	100
3	V	25/25 (100%)	25 (100%)	0	100	100
3	v	25/25 (100%)	25 (100%)	0	100	100
4	C	350/350 (100%)	349 (100%)	1 (0%)	92	97
4	c	350/350 (100%)	349 (100%)	1 (0%)	92	97
5	D	279/279 (100%)	279 (100%)	0	100	100
5	d	279/279 (100%)	278 (100%)	1 (0%)	91	96
6	E	68/68 (100%)	68 (100%)	0	100	100
6	e	68/68 (100%)	68 (100%)	0	100	100
7	F	25/25 (100%)	25 (100%)	0	100	100
7	f	25/25 (100%)	25 (100%)	0	100	100
8	H	56/56 (100%)	56 (100%)	0	100	100
8	h	56/56 (100%)	56 (100%)	0	100	100
9	I	31/31 (100%)	30 (97%)	1 (3%)	39	70
9	i	31/31 (100%)	31 (100%)	0	100	100
10	J	27/27 (100%)	27 (100%)	0	100	100
10	j	27/27 (100%)	27 (100%)	0	100	100

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Mol	Chain	Analysed	Rotameric	Outliers	Percentiles	
11	K	33/33 (100%)	33 (100%)	0	100	100
11	k	33/33 (100%)	33 (100%)	0	100	100
12	L	35/35 (100%)	35 (100%)	0	100	100
12	l	35/35 (100%)	35 (100%)	0	100	100
13	M	27/27 (100%)	27 (100%)	0	100	100
13	m	27/27 (100%)	27 (100%)	0	100	100
14	O	195/195 (100%)	192 (98%)	3 (2%)	65	87
14	o	195/195 (100%)	194 (100%)	1 (0%)	88	95
15	P	151/151 (100%)	151 (100%)	0	100	100
15	p	151/151 (100%)	151 (100%)	0	100	100
16	T	26/26 (100%)	26 (100%)	0	100	100
16	t	26/26 (100%)	26 (100%)	0	100	100
17	W	34/34 (100%)	34 (100%)	0	100	100
17	w	34/34 (100%)	34 (100%)	0	100	100
18	X	21/21 (100%)	21 (100%)	0	100	100
18	x	21/21 (100%)	21 (100%)	0	100	100
19	Z	50/50 (100%)	49 (98%)	1 (2%)	55	82
19	z	50/50 (100%)	50 (100%)	0	100	100
20	N	171/171 (100%)	170 (99%)	1 (1%)	86	95
20	n	171/171 (100%)	171 (100%)	0	100	100
21	G	168/168 (100%)	168 (100%)	0	100	100
21	g	168/168 (100%)	168 (100%)	0	100	100
22	R	158/158 (100%)	157 (99%)	1 (1%)	86	95
22	r	158/158 (100%)	158 (100%)	0	100	100
23	S	189/190 (100%)	188 (100%)	1 (0%)	88	95
23	s	190/190 (100%)	190 (100%)	0	100	100
24	Y	167/167 (100%)	166 (99%)	1 (1%)	86	95
24	y	167/167 (100%)	165 (99%)	2 (1%)	71	90
25	U	26/26 (100%)	26 (100%)	0	100	100
25	u	26/26 (100%)	26 (100%)	0	100	100
All	All	5953/5952 (100%)	5935 (100%)	18 (0%)	92	97

All (18) residues with a non-rotameric sidechain are listed below:

Mol	Chain	Res	Type
1	A	334	ARG
4	C	406	ASN
9	I	34	LYS
14	O	120	ASP
14	O	124	ARG
14	O	137	THR
19	Z	58	ASN
20	N	170	ASN
22	R	270	LYS
23	S	285	ARG
24	Y	46	ARG
1	a	92	HIS
1	a	332	HIS
4	c	406	ASN
5	d	180	ARG
14	o	124	ARG
24	y	76	LEU
24	y	149	GLN

Sometimes sidechains can be flipped to improve hydrogen bonding and reduce clashes. All (16) such sidechains are listed below:

Mol	Chain	Res	Type
1	A	187	GLN
1	A	303	ASN
2	B	26	HIS
4	C	16	GLN
4	C	406	ASN
8	H	69	ASN
24	Y	247	ASN
1	a	92	HIS
1	a	181	ASN
1	a	198	HIS
1	a	230	ASN
3	v	32	ASN
8	h	69	ASN
15	p	109	ASN
20	n	170	ASN
24	y	149	GLN

5.3.3 RNA ⓘ

There are no RNA molecules in this entry.

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

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

5.5 Carbohydrates ⓘ

There are no monosaccharides in this entry.

5.6 Ligand geometry ⓘ

Of 343 ligands modelled in this entry, 8 are monoatomic - leaving 335 for Mogul analysis.

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

Mol	Type	Chain	Res	Link	Bond lengths			Bond angles		
					Counts	RMSZ	# $ Z > 2$	Counts	RMSZ	# $ Z > 2$
40	LMK	C	527	-	38,39,53	1.48	2 (5%)	41,46,60	1.30	2 (4%)
33	LMG	C	521	-	51,51,55	1.06	6 (11%)	59,59,63	1.09	3 (5%)
48	NEX	G	623	-	38,46,46	3.31	10 (26%)	50,70,70	1.77	11 (22%)
29	CLA	N	612	-	45,53,73	1.63	9 (20%)	52,89,113	2.14	11 (21%)
29	CLA	y	610	-	65,73,73	1.35	8 (12%)	76,113,113	1.97	17 (22%)
39	LHG	C	525	-	46,46,48	0.39	0	49,52,54	1.03	2 (4%)
29	CLA	y	614	-	65,73,73	1.35	8 (12%)	76,113,113	1.95	16 (21%)
29	CLA	c	504	-	65,73,73	1.32	7 (10%)	76,113,113	2.05	19 (25%)
29	CLA	n	614	-	49,57,73	1.55	10 (20%)	55,93,113	2.30	18 (32%)
37	GOL	B	627	-	5,5,5	0.58	0	5,5,5	0.21	0
39	LHG	D	409	-	48,48,48	0.39	0	51,54,54	1.04	3 (5%)
48	NEX	r	623	-	38,46,46	3.32	9 (23%)	50,70,70	1.54	9 (18%)
45	CHL	y	601	24	66,74,74	0.79	2 (3%)	73,114,114	1.20	8 (10%)
29	CLA	G	613	-	65,73,73	1.35	9 (13%)	76,113,113	2.02	18 (23%)

Mol	Type	Chain	Res	Link	Bond lengths			Bond angles		
					Counts	RMSZ	# Z > 2	Counts	RMSZ	# Z > 2
31	BCR	C	516	-	41,41,41	1.85	4 (9%)	56,56,56	4.37	16 (28%)
31	BCR	a	411	-	41,41,41	1.83	4 (9%)	56,56,56	4.23	13 (23%)
45	CHL	N	609	-	66,74,74	0.78	2 (3%)	73,114,114	1.25	11 (15%)
51	SPH	y	625	-	19,20,20	0.64	0	18,21,21	1.10	1 (5%)
45	CHL	s	607	-	43,51,74	1.02	3 (6%)	45,86,114	1.44	7 (15%)
39	LHG	S	624	-	44,44,48	0.41	0	47,50,54	1.10	3 (6%)
46	LUT	s	621	-	42,43,43	2.32	1 (2%)	51,60,60	1.99	14 (27%)
46	LUT	y	620	-	42,43,43	2.33	1 (2%)	51,60,60	1.94	15 (29%)
32	SQD	a	412	-	50,51,54	0.81	0	59,62,65	0.91	2 (3%)
29	CLA	N	603	-	65,73,73	1.35	8 (12%)	76,113,113	2.04	21 (27%)
45	CHL	y	606	-	66,74,74	0.85	3 (4%)	73,114,114	1.22	10 (13%)
49	LPX	S	625	-	29,29,29	0.99	2 (6%)	31,33,33	0.98	1 (3%)
29	CLA	D	402	-	65,73,73	1.35	8 (12%)	76,113,113	1.93	15 (19%)
45	CHL	G	605	21	48,56,74	0.96	2 (4%)	51,92,114	1.38	10 (19%)
45	CHL	n	608	-	50,58,74	0.90	2 (4%)	52,94,114	1.41	11 (21%)
30	PHO	A	408	-	51,69,69	0.99	4 (7%)	47,99,99	1.15	4 (8%)
29	CLA	r	613	-	46,54,73	1.60	9 (19%)	53,90,113	2.19	15 (28%)
29	CLA	Y	610	-	65,73,73	1.35	8 (12%)	76,113,113	1.98	17 (22%)
29	CLA	a	406	-	65,73,73	1.33	6 (9%)	76,113,113	2.06	17 (22%)
29	CLA	R	610	-	60,68,73	1.42	9 (15%)	70,107,113	2.07	20 (28%)
29	CLA	r	610	-	60,68,73	1.40	10 (16%)	70,107,113	2.15	20 (28%)
36	DGA	B	625	-	43,43,43	1.13	2 (4%)	45,45,45	1.51	3 (6%)
33	LMG	B	622	-	44,44,55	0.85	2 (4%)	52,52,63	1.06	3 (5%)
45	CHL	y	607	-	66,74,74	0.77	2 (3%)	73,114,114	1.22	10 (13%)
41	BCT	D	401	27	2,3,3	1.27	0	2,3,3	2.64	2 (100%)
29	CLA	s	603	-	65,73,73	1.38	10 (15%)	76,113,113	1.89	14 (18%)
29	CLA	R	609	-	60,68,73	1.42	7 (11%)	70,107,113	2.02	15 (21%)
29	CLA	r	609	-	60,68,73	1.41	8 (13%)	70,107,113	2.02	14 (20%)
29	CLA	Y	604	-	65,73,73	1.35	9 (13%)	76,113,113	1.93	18 (23%)
29	CLA	C	506	-	65,73,73	1.36	8 (12%)	76,113,113	2.03	19 (25%)
29	CLA	s	611	-	65,73,73	1.36	7 (10%)	76,113,113	1.96	16 (21%)
47	XAT	n	622	-	39,47,47	0.70	1 (2%)	54,74,74	1.97	12 (22%)
29	CLA	R	603	-	60,68,73	1.41	7 (11%)	70,107,113	2.05	18 (25%)
39	LHG	L	101	-	48,48,48	0.40	0	51,54,54	0.92	2 (3%)
33	LMG	b	622	-	44,44,55	0.85	3 (6%)	52,52,63	1.07	3 (5%)

Mol	Type	Chain	Res	Link	Bond lengths			Bond angles		
					Counts	RMSZ	# Z > 2	Counts	RMSZ	# Z > 2
38	DGD	c	519	-	63,63,67	1.12	5 (7%)	77,77,81	1.03	3 (3%)
37	GOL	y	626	-	5,5,5	0.57	0	5,5,5	0.25	0
47	XAT	G	622	-	39,47,47	0.69	1 (2%)	54,74,74	1.90	13 (24%)
35	C7Z	B	620	-	43,43,43	5.41	26 (60%)	58,60,60	2.04	19 (32%)
45	CHL	g	601	21	66,74,74	0.82	3 (4%)	73,114,114	1.24	10 (13%)
29	CLA	S	605	-	50,58,73	1.58	9 (18%)	58,95,113	2.44	18 (31%)
45	CHL	Y	606	-	66,74,74	0.87	4 (6%)	73,114,114	1.16	9 (12%)
29	CLA	C	507	52	65,73,73	1.37	7 (10%)	76,113,113	1.96	18 (23%)
31	BCR	C	515	-	41,41,41	1.84	4 (9%)	56,56,56	4.22	13 (23%)
38	DGD	c	518	-	56,56,67	0.99	4 (7%)	70,70,81	1.05	3 (4%)
46	LUT	g	620	-	42,43,43	2.37	1 (2%)	51,60,60	1.95	11 (21%)
36	DGA	b	623	-	43,43,43	1.13	2 (4%)	45,45,45	1.51	3 (6%)
30	PHO	a	408	-	51,69,69	1.00	4 (7%)	47,99,99	1.15	6 (12%)
29	CLA	y	611	-	65,73,73	1.36	8 (12%)	76,113,113	1.94	16 (21%)
29	CLA	B	607	-	65,73,73	1.36	8 (12%)	76,113,113	2.00	18 (23%)
29	CLA	C	510	-	65,73,73	1.33	7 (10%)	76,113,113	2.06	16 (21%)
45	CHL	Y	607	-	66,74,74	0.76	2 (3%)	73,114,114	1.21	9 (12%)
29	CLA	S	603	-	65,73,73	1.39	10 (15%)	76,113,113	1.91	14 (18%)
39	LHG	Y	624	-	48,48,48	0.38	0	51,54,54	1.03	3 (5%)
45	CHL	N	601	-	66,74,74	0.82	3 (4%)	73,114,114	1.17	8 (10%)
29	CLA	n	612	-	45,53,73	1.63	8 (17%)	52,89,113	2.11	14 (26%)
29	CLA	G	602	-	65,73,73	1.35	9 (13%)	76,113,113	1.97	18 (23%)
29	CLA	c	510	-	65,73,73	1.33	8 (12%)	76,113,113	2.03	16 (21%)
29	CLA	Y	611	-	65,73,73	1.36	8 (12%)	76,113,113	1.89	13 (17%)
46	LUT	n	620	-	42,43,43	2.37	1 (2%)	51,60,60	1.82	14 (27%)
29	CLA	B	609	-	65,73,73	1.34	8 (12%)	76,113,113	2.10	17 (22%)
29	CLA	Y	614	-	65,73,73	1.35	9 (13%)	76,113,113	1.97	17 (22%)
29	CLA	G	612	-	43,51,73	1.67	8 (18%)	49,86,113	2.18	12 (24%)
45	CHL	g	607	-	50,58,74	0.88	2 (4%)	52,94,114	1.41	10 (19%)
29	CLA	s	617	-	50,58,73	1.53	9 (18%)	58,95,113	2.27	19 (32%)
29	CLA	c	508	-	65,73,73	1.34	7 (10%)	76,113,113	2.00	18 (23%)
43	HEM	F	101	7,6	41,50,50	1.55	3 (7%)	45,82,82	1.51	6 (13%)
29	CLA	B	608	-	65,73,73	1.33	7 (10%)	76,113,113	2.04	16 (21%)
29	CLA	c	507	52	65,73,73	1.37	9 (13%)	76,113,113	1.97	19 (25%)
29	CLA	r	602	-	60,68,73	1.41	9 (15%)	70,107,113	2.03	17 (24%)

Mol	Type	Chain	Res	Link	Bond lengths			Bond angles		
					Counts	RMSZ	# Z > 2	Counts	RMSZ	# Z > 2
44	RRX	H	101	-	42,42,42	4.87	24 (57%)	57,58,58	2.04	18 (31%)
29	CLA	c	509	-	65,73,73	1.32	7 (10%)	76,113,113	2.01	17 (22%)
29	CLA	C	508	-	65,73,73	1.34	7 (10%)	76,113,113	1.99	16 (21%)
33	LMG	d	411	-	46,46,55	0.92	3 (6%)	54,54,63	1.10	2 (3%)
45	CHL	g	608	-	44,52,74	0.99	3 (6%)	46,87,114	1.42	9 (19%)
36	DGA	c	524	-	43,43,43	1.13	3 (6%)	45,45,45	1.52	3 (6%)
47	XAT	Y	622	-	39,47,47	0.70	1 (2%)	54,74,74	3.70	16 (29%)
48	NEX	n	623	-	38,46,46	3.37	10 (26%)	50,70,70	1.71	12 (24%)
38	DGD	C	518	-	56,56,67	0.99	4 (7%)	70,70,81	1.04	3 (4%)
29	CLA	b	606	-	65,73,73	1.33	8 (12%)	76,113,113	2.05	14 (18%)
37	GOL	b	625	-	5,5,5	0.58	0	5,5,5	0.28	0
42	PL9	d	405	-	55,55,55	1.21	4 (7%)	68,69,69	1.49	12 (17%)
33	LMG	J	101	-	45,45,55	0.90	3 (6%)	53,53,63	1.00	2 (3%)
45	CHL	N	607	-	66,74,74	0.76	2 (3%)	73,114,114	1.27	10 (13%)
46	LUT	N	620	-	42,43,43	2.35	1 (2%)	51,60,60	1.92	13 (25%)
46	LUT	N	621	-	42,43,43	2.36	1 (2%)	51,60,60	1.92	13 (25%)
29	CLA	s	614	-	55,63,73	1.46	7 (12%)	64,101,113	2.17	15 (23%)
45	CHL	g	609	-	66,74,74	0.86	3 (4%)	73,114,114	1.23	10 (13%)
31	BCR	C	517	-	41,41,41	1.83	4 (9%)	56,56,56	4.23	14 (25%)
29	CLA	n	611	-	49,57,73	1.56	9 (18%)	55,93,113	2.27	15 (27%)
32	SQD	c	626	-	53,54,54	0.79	0	62,65,65	0.90	2 (3%)
32	SQD	A	412	-	50,51,54	0.84	0	59,62,65	0.94	3 (5%)
49	LPX	s	625	-	29,29,29	0.98	2 (6%)	31,33,33	0.98	1 (3%)
29	CLA	g	611	-	45,53,73	1.62	8 (17%)	52,89,113	2.20	13 (25%)
29	CLA	g	604	-	49,57,73	1.56	8 (16%)	55,93,113	2.27	17 (30%)
45	CHL	N	605	20	66,74,74	0.83	3 (4%)	73,114,114	1.20	10 (13%)
45	CHL	S	601	23	46,54,74	1.02	4 (8%)	49,90,114	1.35	9 (18%)
29	CLA	S	612	-	45,53,73	1.63	8 (17%)	52,89,113	2.14	12 (23%)
29	CLA	s	609	-	60,68,73	1.42	9 (15%)	70,107,113	2.00	16 (22%)
39	LHG	G	630	-	48,48,48	0.39	0	51,54,54	1.05	3 (5%)
29	CLA	R	604	-	49,57,73	1.52	8 (16%)	55,93,113	2.33	18 (32%)
29	CLA	s	610	-	65,73,73	1.37	9 (13%)	76,113,113	1.97	19 (25%)
29	CLA	c	502	-	65,73,73	1.36	7 (10%)	76,113,113	2.13	16 (21%)
33	LMG	a	413	-	48,48,55	0.99	5 (10%)	56,56,63	1.12	3 (5%)
29	CLA	G	604	-	49,57,73	1.56	8 (16%)	55,93,113	2.30	17 (30%)

Mol	Type	Chain	Res	Link	Bond lengths			Bond angles		
					Counts	RMSZ	# Z > 2	Counts	RMSZ	# Z > 2
39	LHG	y	624	-	48,48,48	0.38	0	51,54,54	1.02	3 (5%)
46	LUT	R	620	-	42,43,43	2.35	1 (2%)	51,60,60	2.16	14 (27%)
31	BCR	B	618	-	41,41,41	1.81	4 (9%)	56,56,56	4.29	15 (26%)
26	OEX	a	401	4,52,1	0,15,15	-	-	-		
29	CLA	N	611	-	49,57,73	1.57	10 (20%)	55,93,113	2.22	15 (27%)
29	CLA	G	603	-	65,73,73	1.34	8 (12%)	76,113,113	2.00	19 (25%)
29	CLA	c	506	-	65,73,73	1.37	8 (12%)	76,113,113	2.05	19 (25%)
46	LUT	g	621	-	42,43,43	2.36	1 (2%)	51,60,60	2.02	12 (23%)
29	CLA	b	604	-	65,73,73	1.36	8 (12%)	76,113,113	1.91	18 (23%)
48	NEX	y	623	-	38,46,46	3.37	10 (26%)	50,70,70	1.71	13 (26%)
29	CLA	b	615	-	65,73,73	1.34	7 (10%)	76,113,113	2.16	19 (25%)
45	CHL	G	606	-	50,58,74	0.99	4 (8%)	52,94,114	1.38	8 (15%)
45	CHL	N	608	-	50,58,74	0.89	2 (4%)	52,94,114	1.41	11 (21%)
29	CLA	s	605	-	50,58,73	1.57	9 (18%)	58,95,113	2.40	18 (31%)
29	CLA	S	617	-	50,58,73	1.55	9 (18%)	58,95,113	2.25	16 (27%)
29	CLA	D	403	-	65,73,73	1.38	8 (12%)	76,113,113	1.96	16 (21%)
45	CHL	G	609	-	66,74,74	0.86	3 (4%)	73,114,114	1.19	10 (13%)
33	LMG	A	413	-	48,48,55	0.99	5 (10%)	56,56,63	1.17	4 (7%)
29	CLA	B	615	-	65,73,73	1.34	8 (12%)	76,113,113	2.17	20 (26%)
45	CHL	R	607	-	50,58,74	0.96	3 (6%)	52,94,114	1.38	8 (15%)
45	CHL	r	606	-	44,52,74	1.02	3 (6%)	46,87,114	1.36	9 (19%)
45	CHL	S	606	-	44,52,74	1.02	3 (6%)	46,87,114	1.40	9 (19%)
29	CLA	y	613	-	65,73,73	1.35	8 (12%)	76,113,113	2.03	19 (25%)
45	CHL	s	608	-	61,69,74	0.86	3 (4%)	67,108,114	1.24	9 (13%)
29	CLA	B	606	-	65,73,73	1.33	8 (12%)	76,113,113	2.05	14 (18%)
37	GOL	b	624	-	5,5,5	0.59	0	5,5,5	0.22	0
39	LHG	g	624	-	48,48,48	0.39	0	51,54,54	1.04	3 (5%)
29	CLA	G	614	-	49,57,73	1.57	9 (18%)	55,93,113	2.28	16 (29%)
45	CHL	N	606	-	66,74,74	0.87	4 (6%)	73,114,114	1.17	9 (12%)
31	BCR	c	516	-	41,41,41	1.86	4 (9%)	56,56,56	4.30	15 (26%)
47	XAT	R	621	-	39,47,47	0.69	1 (2%)	54,74,74	2.22	17 (31%)
29	CLA	n	604	-	65,73,73	1.35	8 (12%)	76,113,113	2.05	20 (26%)
50	3PH	S	626	-	47,47,47	0.86	4 (8%)	51,52,52	1.13	2 (3%)
29	CLA	c	503	-	65,73,73	1.37	9 (13%)	76,113,113	2.01	18 (23%)
29	CLA	C	505	-	65,73,73	1.36	9 (13%)	76,113,113	2.03	16 (21%)

Mol	Type	Chain	Res	Link	Bond lengths			Bond angles		
					Counts	RMSZ	# Z > 2	Counts	RMSZ	# Z > 2
29	CLA	r	604	-	49,57,73	1.54	8 (16%)	55,93,113	2.28	17 (30%)
29	CLA	b	614	-	65,73,73	1.33	6 (9%)	76,113,113	2.00	18 (23%)
38	DGD	C	519	-	63,63,67	1.12	6 (9%)	77,77,81	1.06	4 (5%)
38	DGD	C	520	-	60,60,67	1.07	6 (10%)	74,74,81	0.98	2 (2%)
29	CLA	g	614	-	49,57,73	1.56	9 (18%)	55,93,113	2.26	15 (27%)
41	BCT	d	401	27	2,3,3	1.26	0	2,3,3	2.67	2 (100%)
48	NEX	Y	623	-	38,46,46	3.32	9 (23%)	50,70,70	1.80	11 (22%)
29	CLA	g	613	-	65,73,73	1.36	8 (12%)	76,113,113	2.02	18 (23%)
29	CLA	n	602	-	65,73,73	1.36	8 (12%)	76,113,113	1.97	22 (28%)
33	LMG	c	521	-	51,51,55	1.07	6 (11%)	59,59,63	1.10	3 (5%)
29	CLA	C	504	-	65,73,73	1.33	7 (10%)	76,113,113	2.05	18 (23%)
48	NEX	g	623	-	38,46,46	3.33	9 (23%)	50,70,70	1.81	13 (26%)
46	LUT	s	620	-	42,43,43	2.41	1 (2%)	51,60,60	2.12	16 (31%)
29	CLA	B	614	-	65,73,73	1.32	6 (9%)	76,113,113	2.01	18 (23%)
29	CLA	N	604	-	65,73,73	1.34	8 (12%)	76,113,113	2.05	20 (26%)
33	LMG	h	102	-	48,48,55	1.00	4 (8%)	56,56,63	1.09	2 (3%)
38	DGD	c	520	-	60,60,67	1.06	6 (10%)	74,74,81	0.98	2 (2%)
32	SQD	B	621	-	53,54,54	0.79	0	62,65,65	0.90	2 (3%)
29	CLA	S	610	-	65,73,73	1.37	8 (12%)	76,113,113	1.96	19 (25%)
45	CHL	s	606	-	44,52,74	1.01	3 (6%)	46,87,114	1.43	10 (21%)
31	BCR	C	514	-	41,41,41	1.84	4 (9%)	56,56,56	4.43	12 (21%)
47	XAT	g	622	-	39,47,47	0.69	1 (2%)	54,74,74	1.88	15 (27%)
36	DGA	C	524	-	43,43,43	1.13	3 (6%)	45,45,45	1.52	3 (6%)
45	CHL	S	608	-	61,69,74	0.87	3 (4%)	67,108,114	1.26	10 (14%)
29	CLA	Y	613	-	65,73,73	1.34	7 (10%)	76,113,113	2.03	20 (26%)
29	CLA	S	609	-	60,68,73	1.41	8 (13%)	70,107,113	2.04	17 (24%)
29	CLA	n	613	-	65,73,73	1.36	10 (15%)	76,113,113	2.01	16 (21%)
47	XAT	r	622	-	39,47,47	0.72	1 (2%)	54,74,74	2.18	19 (35%)
31	BCR	b	619	-	41,41,41	1.83	4 (9%)	56,56,56	4.38	17 (30%)
48	NEX	s	623	-	38,46,46	3.34	10 (26%)	50,70,70	1.72	11 (22%)
33	LMG	D	411	-	46,46,55	0.92	4 (8%)	54,54,63	1.11	2 (3%)
31	BCR	B	619	-	41,41,41	1.84	4 (9%)	56,56,56	4.36	18 (32%)
45	CHL	S	607	-	43,51,74	1.01	3 (6%)	45,86,114	1.45	8 (17%)
45	CHL	n	605	20	66,74,74	0.84	3 (4%)	73,114,114	1.18	10 (13%)
32	SQD	C	526	-	53,54,54	0.79	0	62,65,65	0.89	2 (3%)

Mol	Type	Chain	Res	Link	Bond lengths			Bond angles		
					Counts	RMSZ	# Z > 2	Counts	RMSZ	# Z > 2
47	XAT	N	622	-	39,47,47	0.69	1 (2%)	54,74,74	1.96	13 (24%)
29	CLA	S	614	-	55,63,73	1.47	8 (14%)	64,101,113	2.12	15 (23%)
30	PHO	A	409	-	51,69,69	1.00	3 (5%)	47,99,99	1.20	3 (6%)
38	DGD	c	523	-	67,67,67	1.18	7 (10%)	81,81,81	0.98	2 (2%)
39	LHG	c	625	-	46,46,48	0.41	0	49,52,54	1.00	2 (4%)
29	CLA	r	603	-	60,68,73	1.42	9 (15%)	70,107,113	2.03	17 (24%)
45	CHL	G	601	21	66,74,74	0.81	3 (4%)	73,114,114	1.24	11 (15%)
29	CLA	c	512	-	65,73,73	1.34	7 (10%)	76,113,113	1.96	19 (25%)
29	CLA	b	608	-	65,73,73	1.34	6 (9%)	76,113,113	2.05	17 (22%)
29	CLA	B	610	-	65,73,73	1.34	8 (12%)	76,113,113	1.99	16 (21%)
39	LHG	N	624	-	48,48,48	0.38	0	51,54,54	1.11	2 (3%)
29	CLA	n	603	-	65,73,73	1.35	8 (12%)	76,113,113	2.07	19 (25%)
29	CLA	N	614	-	49,57,73	1.55	8 (16%)	55,93,113	2.26	16 (29%)
46	LUT	G	621	-	42,43,43	2.37	1 (2%)	51,60,60	1.99	13 (25%)
29	CLA	g	612	-	43,51,73	1.67	8 (18%)	49,86,113	2.19	13 (26%)
48	NEX	S	622	-	38,46,46	3.28	9 (23%)	50,70,70	1.80	12 (24%)
29	CLA	C	512	-	65,73,73	1.35	7 (10%)	76,113,113	1.93	17 (22%)
45	CHL	Y	609	-	66,74,74	0.81	3 (4%)	73,114,114	1.23	12 (16%)
29	CLA	b	613	-	65,73,73	1.32	8 (12%)	76,113,113	1.96	17 (22%)
29	CLA	C	511	-	65,73,73	1.35	8 (12%)	76,113,113	2.11	19 (25%)
29	CLA	b	612	-	65,73,73	1.34	7 (10%)	76,113,113	2.05	17 (22%)
39	LHG	s	624	-	44,44,48	0.42	0	47,50,54	1.11	3 (6%)
29	CLA	S	611	-	65,73,73	1.37	8 (12%)	76,113,113	1.95	15 (19%)
29	CLA	B	612	-	65,73,73	1.33	8 (12%)	76,113,113	1.99	17 (22%)
29	CLA	c	511	-	65,73,73	1.34	7 (10%)	76,113,113	2.08	20 (26%)
45	CHL	G	607	-	50,58,74	0.87	2 (4%)	52,94,114	1.42	11 (21%)
39	LHG	D	408	-	43,43,48	0.41	0	46,49,54	1.14	3 (6%)
29	CLA	G	611	-	45,53,73	1.62	8 (17%)	52,89,113	2.19	15 (28%)
29	CLA	N	613	-	65,73,73	1.37	10 (15%)	76,113,113	1.99	16 (21%)
29	CLA	N	610	-	65,73,73	1.36	8 (12%)	76,113,113	2.01	18 (23%)
50	3PH	i	101	-	47,47,47	0.86	4 (8%)	51,52,52	1.13	2 (3%)
29	CLA	b	611	-	65,73,73	1.36	8 (12%)	76,113,113	1.96	16 (21%)
29	CLA	c	501	-	65,73,73	1.36	8 (12%)	76,113,113	2.08	18 (23%)
29	CLA	y	604	-	65,73,73	1.34	6 (9%)	76,113,113	1.95	18 (23%)
29	CLA	a	405	-	65,73,73	1.33	6 (9%)	76,113,113	2.01	19 (25%)

Mol	Type	Chain	Res	Link	Bond lengths			Bond angles		
					Counts	RMSZ	# Z > 2	Counts	RMSZ	# Z > 2
29	CLA	r	612	-	60,68,73	1.43	9 (15%)	70,107,113	2.03	17 (24%)
29	CLA	Y	602	-	65,73,73	1.36	9 (13%)	76,113,113	1.92	19 (25%)
39	LHG	l	101	-	48,48,48	0.40	0	51,54,54	0.92	2 (3%)
29	CLA	C	502	-	65,73,73	1.34	7 (10%)	76,113,113	2.04	16 (21%)
29	CLA	B	611	-	65,73,73	1.36	8 (12%)	76,113,113	1.95	15 (19%)
29	CLA	s	613	-	55,63,73	1.49	8 (14%)	64,101,113	2.32	16 (25%)
29	CLA	A	406	-	65,73,73	1.34	7 (10%)	76,113,113	2.03	16 (21%)
29	CLA	a	410	-	60,68,73	1.40	7 (11%)	70,107,113	2.11	17 (24%)
29	CLA	C	513	-	65,73,73	1.36	8 (12%)	76,113,113	2.02	19 (25%)
35	C7Z	b	620	-	43,43,43	5.40	26 (60%)	58,60,60	2.05	17 (29%)
29	CLA	C	503	-	65,73,73	1.36	9 (13%)	76,113,113	2.02	19 (25%)
29	CLA	a	407	-	49,57,73	1.56	8 (16%)	55,93,113	2.24	18 (32%)
45	CHL	R	606	-	44,52,74	1.02	3 (6%)	46,87,114	1.33	8 (17%)
45	CHL	y	605	24	46,54,74	1.00	2 (4%)	49,90,114	1.38	8 (16%)
29	CLA	R	613	-	46,54,73	1.61	8 (17%)	53,90,113	2.18	14 (26%)
29	CLA	S	604	-	55,63,73	1.45	7 (12%)	64,101,113	2.18	17 (26%)
29	CLA	c	513	-	65,73,73	1.35	7 (10%)	76,113,113	2.02	16 (21%)
29	CLA	y	602	-	65,73,73	1.35	8 (12%)	76,113,113	1.94	21 (27%)
29	CLA	R	608	-	60,68,73	1.43	9 (15%)	70,107,113	2.01	16 (22%)
46	LUT	n	621	-	42,43,43	2.33	1 (2%)	51,60,60	1.85	11 (21%)
29	CLA	d	402	-	65,73,73	1.35	8 (12%)	76,113,113	1.90	14 (18%)
33	LMG	j	101	-	45,45,55	0.89	3 (6%)	53,53,63	1.04	2 (3%)
29	CLA	C	501	-	65,73,73	1.36	9 (13%)	76,113,113	2.06	18 (23%)
29	CLA	S	613	-	55,63,73	1.48	8 (14%)	64,101,113	2.33	16 (25%)
29	CLA	y	608	-	50,58,73	1.55	8 (16%)	58,95,113	2.22	18 (31%)
45	CHL	n	609	-	66,74,74	0.78	2 (3%)	73,114,114	1.33	13 (17%)
29	CLA	B	605	-	65,73,73	1.39	8 (12%)	76,113,113	2.10	15 (19%)
46	LUT	G	620	-	42,43,43	2.35	1 (2%)	51,60,60	1.96	12 (23%)
45	CHL	Y	605	24	46,54,74	1.01	3 (6%)	49,90,114	1.41	10 (20%)
29	CLA	R	602	-	60,68,73	1.42	9 (15%)	70,107,113	2.10	18 (25%)
48	NEX	R	622	-	38,46,46	3.42	10 (26%)	50,70,70	1.69	11 (22%)
29	CLA	s	604	-	55,63,73	1.48	7 (12%)	64,101,113	2.07	14 (21%)
46	LUT	S	620	-	42,43,43	2.39	1 (2%)	51,60,60	2.09	12 (23%)
45	CHL	G	608	-	44,52,74	1.00	3 (6%)	46,87,114	1.41	9 (19%)
29	CLA	Y	603	-	65,73,73	1.34	8 (12%)	76,113,113	1.98	19 (25%)

Mol	Type	Chain	Res	Link	Bond lengths			Bond angles		
					Counts	RMSZ	# Z > 2	Counts	RMSZ	# Z > 2
31	BCR	b	618	-	41,41,41	1.82	4 (9%)	56,56,56	4.30	15 (26%)
32	SQD	b	621	-	53,54,54	0.79	0	62,65,65	0.90	2 (3%)
45	CHL	g	606	-	50,58,74	0.89	2 (4%)	52,94,114	1.39	9 (17%)
45	CHL	n	601	-	66,74,74	0.82	2 (3%)	73,114,114	1.21	8 (10%)
29	CLA	g	602	-	65,73,73	1.35	8 (12%)	76,113,113	1.99	19 (25%)
29	CLA	A	410	-	60,68,73	1.41	9 (15%)	70,107,113	2.10	16 (22%)
46	LUT	Y	620	-	42,43,43	2.36	1 (2%)	51,60,60	2.00	13 (25%)
39	LHG	d	408	-	43,43,48	0.41	0	46,49,54	1.14	3 (6%)
46	LUT	r	620	-	42,43,43	2.35	1 (2%)	51,60,60	2.20	15 (29%)
29	CLA	b	607	-	65,73,73	1.35	8 (12%)	76,113,113	2.00	19 (25%)
29	CLA	A	407	-	49,57,73	1.56	7 (14%)	55,93,113	2.25	16 (29%)
29	CLA	n	610	-	65,73,73	1.35	8 (12%)	76,113,113	2.03	17 (22%)
39	LHG	D	410	-	38,38,48	0.43	0	41,44,54	1.07	2 (4%)
39	LHG	d	410	-	38,38,48	0.41	0	41,44,54	1.15	3 (7%)
29	CLA	Y	608	-	50,58,73	1.55	8 (16%)	58,95,113	2.22	19 (32%)
29	CLA	b	616	-	65,73,73	1.35	8 (12%)	76,113,113	1.97	16 (21%)
29	CLA	B	602	-	65,73,73	1.37	9 (13%)	76,113,113	1.99	17 (22%)
46	LUT	y	621	-	42,43,43	2.35	1 (2%)	51,60,60	2.03	12 (23%)
29	CLA	C	509	-	65,73,73	1.33	6 (9%)	76,113,113	2.04	19 (25%)
45	CHL	g	605	21	48,56,74	0.95	3 (6%)	51,92,114	1.41	10 (19%)
29	CLA	y	603	-	65,73,73	1.33	8 (12%)	76,113,113	2.02	19 (25%)
29	CLA	c	505	-	65,73,73	1.36	7 (10%)	76,113,113	2.02	15 (19%)
45	CHL	r	607	-	50,58,74	0.94	3 (6%)	52,94,114	1.37	9 (17%)
45	CHL	n	607	-	66,74,74	0.76	2 (3%)	73,114,114	1.28	10 (13%)
29	CLA	b	602	-	65,73,73	1.37	9 (13%)	76,113,113	1.98	17 (22%)
29	CLA	A	405	-	65,73,73	1.32	6 (9%)	76,113,113	2.03	18 (23%)
30	PHO	a	409	-	51,69,69	0.98	3 (5%)	47,99,99	1.24	5 (10%)
45	CHL	y	609	-	66,74,74	0.83	3 (4%)	73,114,114	1.23	12 (16%)
29	CLA	g	603	-	65,73,73	1.34	8 (12%)	76,113,113	2.00	18 (23%)
29	CLA	y	612	-	65,73,73	1.36	8 (12%)	76,113,113	1.94	15 (19%)
31	BCR	A	411	-	41,41,41	1.84	4 (9%)	56,56,56	4.21	15 (26%)
29	CLA	S	602	-	60,68,73	1.41	8 (13%)	70,107,113	2.03	17 (24%)
26	OEX	A	401	52,1	0,15,15	-	-	-	-	-
39	LHG	d	409	-	48,48,48	0.39	0	51,54,54	1.03	3 (5%)
43	HEM	f	101	7,6	41,50,50	1.54	4 (9%)	45,82,82	1.55	7 (15%)

Mol	Type	Chain	Res	Link	Bond lengths			Bond angles		
					Counts	RMSZ	# Z > 2	Counts	RMSZ	# Z > 2
46	LUT	Y	621	-	42,43,43	2.34	1 (2%)	51,60,60	1.95	14 (27%)
29	CLA	G	610	-	65,73,73	1.35	8 (12%)	76,113,113	2.02	18 (23%)
48	NEX	N	623	-	38,46,46	3.33	9 (23%)	50,70,70	1.76	14 (28%)
44	RRX	h	101	-	42,42,42	4.87	24 (57%)	57,58,58	2.03	19 (33%)
31	BCR	D	404	-	41,41,41	1.85	4 (9%)	56,56,56	4.04	17 (30%)
29	CLA	N	602	-	65,73,73	1.36	7 (10%)	76,113,113	1.99	20 (26%)
29	CLA	b	610	-	65,73,73	1.34	7 (10%)	76,113,113	2.00	15 (19%)
45	CHL	Y	601	24	66,74,74	0.80	2 (3%)	73,114,114	1.20	9 (12%)
31	BCR	c	515	-	41,41,41	1.82	4 (9%)	56,56,56	4.22	13 (23%)
29	CLA	R	612	-	60,68,73	1.42	8 (13%)	70,107,113	2.07	16 (22%)
31	BCR	c	514	-	41,41,41	1.84	4 (9%)	56,56,56	4.38	14 (25%)
29	CLA	r	611	-	46,54,73	1.61	10 (21%)	53,90,113	2.12	14 (26%)
40	LMK	c	627	-	38,39,53	1.48	2 (5%)	41,46,60	1.33	2 (4%)
51	SPH	Y	625	-	19,20,20	0.63	0	18,21,21	1.10	1 (5%)
29	CLA	B	613	-	65,73,73	1.32	7 (10%)	76,113,113	1.95	16 (21%)
29	CLA	b	609	-	65,73,73	1.34	7 (10%)	76,113,113	2.14	17 (22%)
39	LHG	n	624	-	48,48,48	0.38	0	51,54,54	1.11	4 (7%)
45	CHL	n	606	-	66,74,74	0.89	4 (6%)	73,114,114	1.17	9 (12%)
29	CLA	d	403	-	65,73,73	1.38	8 (12%)	76,113,113	1.96	16 (21%)
29	CLA	r	608	-	60,68,73	1.43	8 (13%)	70,107,113	2.03	15 (21%)
29	CLA	b	603	-	65,73,73	1.36	8 (12%)	76,113,113	2.03	18 (23%)
29	CLA	s	602	-	60,68,73	1.38	8 (13%)	70,107,113	2.05	17 (24%)
47	XAT	y	622	-	39,47,47	0.68	1 (2%)	54,74,74	3.73	19 (35%)
50	3PH	s	626	-	47,47,47	0.86	4 (8%)	51,52,52	1.15	2 (3%)
33	LMG	H	102	-	48,48,55	1.00	5 (10%)	56,56,63	1.07	2 (3%)
46	LUT	S	621	-	42,43,43	2.31	1 (2%)	51,60,60	1.97	15 (29%)
29	CLA	b	605	-	65,73,73	1.38	8 (12%)	76,113,113	2.08	16 (21%)
29	CLA	B	604	-	65,73,73	1.36	9 (13%)	76,113,113	1.92	17 (22%)
29	CLA	b	617	-	65,73,73	1.35	7 (10%)	76,113,113	4.30	17 (22%)
29	CLA	Y	612	-	65,73,73	1.36	8 (12%)	76,113,113	1.97	17 (22%)
45	CHL	s	601	23	46,54,74	1.03	4 (8%)	49,90,114	1.37	9 (18%)
29	CLA	B	617	-	65,73,73	1.34	8 (12%)	76,113,113	4.31	17 (22%)
29	CLA	s	612	-	45,53,73	1.61	8 (17%)	52,89,113	2.16	15 (28%)
38	DGD	C	523	-	67,67,67	1.18	7 (10%)	81,81,81	0.98	2 (2%)
29	CLA	B	603	-	65,73,73	1.35	8 (12%)	76,113,113	2.00	18 (23%)

Mol	Type	Chain	Res	Link	Bond lengths			Bond angles		
					Counts	RMSZ	# Z > 2	Counts	RMSZ	# Z > 2
31	BCR	d	404	-	41,41,41	1.84	4 (9%)	56,56,56	4.11	17 (30%)
29	CLA	B	616	-	65,73,73	1.36	7 (10%)	76,113,113	1.97	15 (19%)
29	CLA	R	611	-	46,54,73	1.62	10 (21%)	53,90,113	2.10	14 (26%)
31	BCR	c	517	-	41,41,41	1.81	4 (9%)	56,56,56	4.10	19 (33%)
29	CLA	g	610	-	65,73,73	1.35	7 (10%)	76,113,113	2.01	18 (23%)
42	PL9	D	405	-	55,55,55	1.17	5 (9%)	68,69,69	1.50	13 (19%)

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
40	LMK	C	527	-	1/1/6/6	10/46/46/60	-
33	LMG	C	521	-	-	18/46/66/70	0/1/1/1
48	NEX	G	623	-	-	3/27/83/83	0/3/3/3
29	CLA	N	612	-	1/1/11/20	5/13/91/115	-
29	CLA	y	610	-	1/1/15/20	24/37/115/115	-
39	LHG	C	525	-	-	28/51/51/53	-
29	CLA	y	614	-	1/1/15/20	15/37/115/115	-
29	CLA	c	504	-	1/1/15/20	17/37/115/115	-
29	CLA	n	614	-	1/1/11/20	5/18/96/115	-
37	GOL	B	627	-	-	0/4/4/4	-
39	LHG	D	409	-	-	30/53/53/53	-
48	NEX	r	623	-	-	7/27/83/83	0/3/3/3
45	CHL	y	601	24	4/4/20/26	7/39/137/137	-
29	CLA	G	613	-	1/1/15/20	20/37/115/115	-
45	CHL	N	609	-	4/4/20/26	7/39/137/137	-
31	BCR	C	516	-	-	15/29/63/63	0/2/2/2
31	BCR	a	411	-	-	11/29/63/63	0/2/2/2
51	SPH	y	625	-	-	11/21/21/21	-
45	CHL	s	607	-	4/4/15/26	1/12/110/137	-
39	LHG	S	624	-	-	28/49/49/53	-
46	LUT	s	621	-	-	1/29/67/67	0/2/2/2
46	LUT	y	620	-	-	4/29/67/67	0/2/2/2
32	SQD	a	412	-	-	13/46/66/69	0/1/1/1

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Mol	Type	Chain	Res	Link	Chirals	Torsions	Rings
29	CLA	N	603	-	1/1/15/20	14/37/115/115	-
45	CHL	y	606	-	4/4/20/26	8/39/137/137	-
49	LPX	S	625	-	-	10/31/31/31	-
29	CLA	D	402	-	1/1/15/20	11/37/115/115	-
45	CHL	G	605	21	3/3/16/26	1/18/116/137	-
45	CHL	n	608	-	3/3/16/26	3/20/118/137	-
30	PHO	A	408	-	-	8/37/103/103	0/5/6/6
29	CLA	r	613	-	1/1/11/20	9/15/93/115	-
29	CLA	Y	610	-	1/1/15/20	22/37/115/115	-
29	CLA	a	406	-	1/1/15/20	13/37/115/115	-
29	CLA	R	610	-	1/1/14/20	10/31/109/115	-
29	CLA	r	610	-	1/1/14/20	11/31/109/115	-
36	DGA	B	625	-	-	24/45/45/45	-
33	LMG	B	622	-	-	11/39/59/70	0/1/1/1
45	CHL	y	607	-	4/4/20/26	5/39/137/137	-
29	CLA	s	603	-	1/1/15/20	15/37/115/115	-
29	CLA	R	609	-	1/1/14/20	15/31/109/115	-
29	CLA	r	609	-	1/1/14/20	18/31/109/115	-
29	CLA	Y	604	-	1/1/15/20	18/37/115/115	-
29	CLA	C	506	-	1/1/15/20	21/37/115/115	-
29	CLA	s	611	-	1/1/15/20	11/37/115/115	-
47	XAT	n	622	-	1/1/12/26	0/31/93/93	0/4/4/4
29	CLA	R	603	-	1/1/14/20	16/31/109/115	-
39	LHG	L	101	-	-	30/53/53/53	-
33	LMG	b	622	-	-	11/39/59/70	0/1/1/1
38	DGD	c	519	-	-	21/51/91/95	0/2/2/2
37	GOL	y	626	-	-	0/4/4/4	-
47	XAT	G	622	-	2/2/12/26	1/31/93/93	0/4/4/4
35	C7Z	B	620	-	1/1/12/26	9/29/67/67	0/2/2/2
45	CHL	g	601	21	4/4/20/26	12/39/137/137	-
29	CLA	S	605	-	1/1/12/20	12/19/97/115	-
45	CHL	Y	606	-	4/4/20/26	6/39/137/137	-
29	CLA	C	507	52	1/1/15/20	17/37/115/115	-
31	BCR	C	515	-	-	11/29/63/63	0/2/2/2

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Mol	Type	Chain	Res	Link	Chirals	Torsions	Rings
38	DGD	c	518	-	-	11/44/84/95	0/2/2/2
46	LUT	g	620	-	-	3/29/67/67	0/2/2/2
36	DGA	b	623	-	-	26/45/45/45	-
30	PHO	a	408	-	-	7/37/103/103	0/5/6/6
29	CLA	y	611	-	1/1/15/20	15/37/115/115	-
29	CLA	B	607	-	1/1/15/20	12/37/115/115	-
29	CLA	C	510	-	1/1/15/20	14/37/115/115	-
45	CHL	Y	607	-	4/4/20/26	4/39/137/137	-
29	CLA	S	603	-	1/1/15/20	15/37/115/115	-
39	LHG	Y	624	-	-	31/53/53/53	-
45	CHL	N	601	-	4/4/20/26	5/39/137/137	-
29	CLA	n	612	-	1/1/11/20	6/13/91/115	-
29	CLA	G	602	-	1/1/15/20	21/37/115/115	-
29	CLA	c	510	-	1/1/15/20	12/37/115/115	-
29	CLA	Y	611	-	1/1/15/20	17/37/115/115	-
46	LUT	n	620	-	-	6/29/67/67	0/2/2/2
29	CLA	B	609	-	1/1/15/20	14/37/115/115	-
29	CLA	Y	614	-	1/1/15/20	13/37/115/115	-
29	CLA	G	612	-	1/1/10/20	4/11/89/115	-
45	CHL	g	607	-	3/3/16/26	3/20/118/137	-
29	CLA	s	617	-	1/1/12/20	9/19/97/115	-
29	CLA	c	508	-	1/1/15/20	12/37/115/115	-
43	HEM	F	101	7,6	-	2/12/54/54	-
29	CLA	B	608	-	1/1/15/20	25/37/115/115	-
29	CLA	c	507	52	1/1/15/20	19/37/115/115	-
29	CLA	r	602	-	1/1/14/20	9/31/109/115	-
44	RRX	H	101	-	1/1/11/25	9/29/65/65	0/2/2/2
29	CLA	c	509	-	1/1/15/20	11/37/115/115	-
29	CLA	C	508	-	1/1/15/20	12/37/115/115	-
33	LMG	d	411	-	-	12/41/61/70	0/1/1/1
45	CHL	g	608	-	3/3/15/26	1/13/111/137	-
36	DGA	c	524	-	-	19/45/45/45	-
47	XAT	Y	622	-	1/1/12/26	3/31/93/93	0/4/4/4
48	NEX	n	623	-	-	2/27/83/83	0/3/3/3

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Mol	Type	Chain	Res	Link	Chirals	Torsions	Rings
38	DGD	C	518	-	-	8/44/84/95	0/2/2/2
29	CLA	b	606	-	1/1/15/20	12/37/115/115	-
37	GOL	b	625	-	-	1/4/4/4	-
42	PL9	d	405	-	-	10/53/73/73	0/1/1/1
33	LMG	J	101	-	-	12/40/60/70	0/1/1/1
45	CHL	N	607	-	4/4/20/26	7/39/137/137	-
46	LUT	N	620	-	-	3/29/67/67	0/2/2/2
46	LUT	N	621	-	-	2/29/67/67	0/2/2/2
29	CLA	s	614	-	1/1/13/20	8/25/103/115	-
45	CHL	g	609	-	4/4/20/26	6/39/137/137	-
31	BCR	C	517	-	-	8/29/63/63	0/2/2/2
29	CLA	n	611	-	1/1/11/20	10/18/96/115	-
32	SQD	c	626	-	-	16/49/69/69	0/1/1/1
32	SQD	A	412	-	-	12/46/66/69	0/1/1/1
49	LPX	s	625	-	-	10/31/31/31	-
29	CLA	g	611	-	1/1/11/20	4/13/91/115	-
29	CLA	g	604	-	1/1/11/20	12/18/96/115	-
45	CHL	N	605	20	4/4/20/26	10/39/137/137	-
45	CHL	S	601	23	3/3/16/26	3/15/113/137	-
29	CLA	S	612	-	1/1/11/20	5/13/91/115	-
29	CLA	s	609	-	1/1/14/20	14/31/109/115	-
39	LHG	G	630	-	-	31/53/53/53	-
29	CLA	R	604	-	1/1/11/20	12/18/96/115	-
29	CLA	s	610	-	1/1/15/20	20/37/115/115	-
29	CLA	c	502	-	1/1/15/20	20/37/115/115	-
33	LMG	a	413	-	-	13/43/63/70	0/1/1/1
29	CLA	G	604	-	1/1/11/20	8/18/96/115	-
39	LHG	y	624	-	-	30/53/53/53	-
46	LUT	R	620	-	-	8/29/67/67	0/2/2/2
31	BCR	B	618	-	-	13/29/63/63	0/2/2/2
29	CLA	N	611	-	1/1/11/20	10/18/96/115	-
29	CLA	G	603	-	1/1/15/20	18/37/115/115	-
29	CLA	c	506	-	1/1/15/20	22/37/115/115	-
46	LUT	g	621	-	-	2/29/67/67	0/2/2/2

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Mol	Type	Chain	Res	Link	Chirals	Torsions	Rings
29	CLA	b	604	-	1/1/15/20	14/37/115/115	-
48	NEX	y	623	-	-	3/27/83/83	0/3/3/3
29	CLA	b	615	-	1/1/15/20	12/37/115/115	-
45	CHL	G	606	-	4/4/16/26	5/20/118/137	-
45	CHL	N	608	-	3/3/16/26	3/20/118/137	-
29	CLA	s	605	-	1/1/12/20	8/19/97/115	-
29	CLA	S	617	-	1/1/12/20	10/19/97/115	-
29	CLA	D	403	-	1/1/15/20	12/37/115/115	-
45	CHL	G	609	-	4/4/20/26	9/39/137/137	-
33	LMG	A	413	-	-	14/43/63/70	0/1/1/1
29	CLA	B	615	-	1/1/15/20	8/37/115/115	-
45	CHL	R	607	-	3/3/16/26	5/20/118/137	-
45	CHL	r	606	-	3/3/15/26	1/13/111/137	-
45	CHL	S	606	-	3/3/15/26	0/13/111/137	-
29	CLA	y	613	-	1/1/15/20	19/37/115/115	-
45	CHL	s	608	-	4/4/19/26	1/33/131/137	-
29	CLA	B	606	-	1/1/15/20	12/37/115/115	-
37	GOL	b	624	-	-	0/4/4/4	-
39	LHG	g	624	-	-	28/53/53/53	-
29	CLA	G	614	-	1/1/11/20	10/18/96/115	-
45	CHL	N	606	-	4/4/20/26	4/39/137/137	-
31	BCR	c	516	-	-	13/29/63/63	0/2/2/2
47	XAT	R	621	-	1/1/12/26	11/31/93/93	0/4/4/4
29	CLA	n	604	-	1/1/15/20	16/37/115/115	-
50	3PH	S	626	-	-	19/49/49/49	-
29	CLA	c	503	-	1/1/15/20	20/37/115/115	-
29	CLA	C	505	-	1/1/15/20	12/37/115/115	-
29	CLA	r	604	-	1/1/11/20	12/18/96/115	-
29	CLA	b	614	-	1/1/15/20	10/37/115/115	-
38	DGD	C	519	-	-	19/51/91/95	0/2/2/2
38	DGD	C	520	-	-	12/48/88/95	0/2/2/2
29	CLA	g	614	-	1/1/11/20	9/18/96/115	-
48	NEX	Y	623	-	-	2/27/83/83	0/3/3/3
29	CLA	g	613	-	1/1/15/20	21/37/115/115	-

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Mol	Type	Chain	Res	Link	Chirals	Torsions	Rings
29	CLA	n	602	-	1/1/15/20	16/37/115/115	-
33	LMG	c	521	-	-	18/46/66/70	0/1/1/1
29	CLA	C	504	-	1/1/15/20	14/37/115/115	-
48	NEX	g	623	-	-	3/27/83/83	0/3/3/3
46	LUT	s	620	-	1/1/12/27	3/29/67/67	0/2/2/2
29	CLA	B	614	-	1/1/15/20	12/37/115/115	-
29	CLA	N	604	-	1/1/15/20	19/37/115/115	-
33	LMG	h	102	-	-	14/43/63/70	0/1/1/1
38	DGD	c	520	-	-	12/48/88/95	0/2/2/2
32	SQD	B	621	-	-	19/49/69/69	0/1/1/1
29	CLA	S	610	-	1/1/15/20	11/37/115/115	-
45	CHL	s	606	-	3/3/15/26	1/13/111/137	-
31	BCR	C	514	-	-	11/29/63/63	0/2/2/2
47	XAT	g	622	-	2/2/12/26	0/31/93/93	0/4/4/4
36	DGA	C	524	-	-	18/45/45/45	-
45	CHL	S	608	-	4/4/19/26	2/33/131/137	-
29	CLA	Y	613	-	1/1/15/20	21/37/115/115	-
29	CLA	S	609	-	1/1/14/20	8/31/109/115	-
29	CLA	n	613	-	1/1/15/20	14/37/115/115	-
47	XAT	r	622	-	1/1/12/26	11/31/93/93	0/4/4/4
31	BCR	b	619	-	-	8/29/63/63	0/2/2/2
48	NEX	s	623	-	-	13/27/83/83	0/3/3/3
33	LMG	D	411	-	-	8/41/61/70	0/1/1/1
31	BCR	B	619	-	-	11/29/63/63	0/2/2/2
45	CHL	S	607	-	4/4/15/26	1/12/110/137	-
45	CHL	n	605	20	4/4/20/26	8/39/137/137	-
32	SQD	C	526	-	-	19/49/69/69	0/1/1/1
47	XAT	N	622	-	1/1/12/26	1/31/93/93	0/4/4/4
29	CLA	S	614	-	1/1/13/20	8/25/103/115	-
30	PHO	A	409	-	-	10/37/103/103	0/5/6/6
38	DGD	c	523	-	-	18/55/95/95	0/2/2/2
39	LHG	c	625	-	-	27/51/51/53	-
29	CLA	r	603	-	1/1/14/20	17/31/109/115	-
45	CHL	G	601	21	4/4/20/26	8/39/137/137	-

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Mol	Type	Chain	Res	Link	Chirals	Torsions	Rings
29	CLA	c	512	-	1/1/15/20	19/37/115/115	-
29	CLA	b	608	-	1/1/15/20	25/37/115/115	-
29	CLA	B	610	-	1/1/15/20	15/37/115/115	-
39	LHG	N	624	-	-	30/53/53/53	-
29	CLA	n	603	-	1/1/15/20	22/37/115/115	-
29	CLA	N	614	-	1/1/11/20	7/18/96/115	-
46	LUT	G	621	-	-	2/29/67/67	0/2/2/2
29	CLA	g	612	-	1/1/10/20	4/11/89/115	-
48	NEX	S	622	-	-	11/27/83/83	0/3/3/3
29	CLA	C	512	-	1/1/15/20	17/37/115/115	-
45	CHL	Y	609	-	4/4/20/26	7/39/137/137	-
29	CLA	b	613	-	1/1/15/20	17/37/115/115	-
29	CLA	C	511	-	1/1/15/20	15/37/115/115	-
29	CLA	b	612	-	1/1/15/20	19/37/115/115	-
39	LHG	s	624	-	-	31/49/49/53	-
29	CLA	S	611	-	1/1/15/20	12/37/115/115	-
29	CLA	B	612	-	1/1/15/20	21/37/115/115	-
29	CLA	c	511	-	1/1/15/20	13/37/115/115	-
45	CHL	G	607	-	3/3/16/26	3/20/118/137	-
39	LHG	D	408	-	-	30/48/48/53	-
29	CLA	G	611	-	1/1/11/20	5/13/91/115	-
29	CLA	N	613	-	1/1/15/20	19/37/115/115	-
29	CLA	N	610	-	1/1/15/20	8/37/115/115	-
50	3PH	i	101	-	-	20/49/49/49	-
29	CLA	b	611	-	1/1/15/20	9/37/115/115	-
29	CLA	c	501	-	1/1/15/20	11/37/115/115	-
29	CLA	y	604	-	1/1/15/20	20/37/115/115	-
29	CLA	a	405	-	1/1/15/20	14/37/115/115	-
29	CLA	r	612	-	1/1/14/20	12/31/109/115	-
29	CLA	Y	602	-	1/1/15/20	15/37/115/115	-
39	LHG	l	101	-	-	29/53/53/53	-
29	CLA	C	502	-	1/1/15/20	14/37/115/115	-
29	CLA	B	611	-	1/1/15/20	9/37/115/115	-
29	CLA	s	613	-	1/1/13/20	10/25/103/115	-

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Mol	Type	Chain	Res	Link	Chirals	Torsions	Rings
29	CLA	A	406	-	1/1/15/20	13/37/115/115	-
29	CLA	a	410	-	1/1/14/20	9/31/109/115	-
29	CLA	C	513	-	1/1/15/20	19/37/115/115	-
35	C7Z	b	620	-	1/1/12/26	9/29/67/67	0/2/2/2
29	CLA	C	503	-	1/1/15/20	19/37/115/115	-
29	CLA	a	407	-	1/1/11/20	4/18/96/115	-
45	CHL	R	606	-	3/3/15/26	4/13/111/137	-
45	CHL	y	605	24	3/3/16/26	3/15/113/137	-
29	CLA	R	613	-	1/1/11/20	8/15/93/115	-
29	CLA	S	604	-	1/1/13/20	9/25/103/115	-
29	CLA	c	513	-	1/1/15/20	18/37/115/115	-
29	CLA	y	602	-	1/1/15/20	15/37/115/115	-
29	CLA	R	608	-	1/1/14/20	16/31/109/115	-
46	LUT	n	621	-	-	4/29/67/67	0/2/2/2
29	CLA	d	402	-	1/1/15/20	13/37/115/115	-
33	LMG	j	101	-	-	12/40/60/70	0/1/1/1
29	CLA	C	501	-	1/1/15/20	18/37/115/115	-
29	CLA	S	613	-	1/1/13/20	10/25/103/115	-
29	CLA	y	608	-	1/1/12/20	8/19/97/115	-
45	CHL	n	609	-	4/4/20/26	7/39/137/137	-
29	CLA	B	605	-	1/1/15/20	18/37/115/115	-
46	LUT	G	620	-	-	6/29/67/67	0/2/2/2
45	CHL	Y	605	24	3/3/16/26	1/15/113/137	-
29	CLA	R	602	-	1/1/14/20	13/31/109/115	-
48	NEX	R	622	-	-	8/27/83/83	0/3/3/3
29	CLA	s	604	-	1/1/13/20	12/25/103/115	-
46	LUT	S	620	-	1/1/12/27	4/29/67/67	0/2/2/2
45	CHL	G	608	-	3/3/15/26	0/13/111/137	-
29	CLA	Y	603	-	1/1/15/20	15/37/115/115	-
31	BCR	b	618	-	-	13/29/63/63	0/2/2/2
32	SQD	b	621	-	-	19/49/69/69	0/1/1/1
45	CHL	g	606	-	3/3/16/26	5/20/118/137	-
45	CHL	n	601	-	4/4/20/26	10/39/137/137	-
29	CLA	g	602	-	1/1/15/20	18/37/115/115	-

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Mol	Type	Chain	Res	Link	Chirals	Torsions	Rings
29	CLA	A	410	-	1/1/14/20	10/31/109/115	-
46	LUT	Y	620	-	-	4/29/67/67	0/2/2/2
39	LHG	d	408	-	-	29/48/48/53	-
46	LUT	r	620	-	-	9/29/67/67	0/2/2/2
29	CLA	b	607	-	1/1/15/20	14/37/115/115	-
29	CLA	A	407	-	1/1/11/20	4/18/96/115	-
29	CLA	n	610	-	1/1/15/20	12/37/115/115	-
39	LHG	D	410	-	-	26/43/43/53	-
39	LHG	d	410	-	-	33/43/43/53	-
29	CLA	Y	608	-	1/1/12/20	6/19/97/115	-
29	CLA	b	616	-	1/1/15/20	10/37/115/115	-
29	CLA	B	602	-	1/1/15/20	21/37/115/115	-
46	LUT	y	621	-	-	2/29/67/67	0/2/2/2
29	CLA	C	509	-	1/1/15/20	10/37/115/115	-
45	CHL	g	605	21	3/3/16/26	4/18/116/137	-
29	CLA	y	603	-	1/1/15/20	13/37/115/115	-
29	CLA	c	505	-	1/1/15/20	15/37/115/115	-
45	CHL	r	607	-	3/3/16/26	8/20/118/137	-
45	CHL	n	607	-	4/4/20/26	10/39/137/137	-
29	CLA	b	602	-	1/1/15/20	21/37/115/115	-
29	CLA	A	405	-	1/1/15/20	16/37/115/115	-
30	PHO	a	409	-	-	9/37/103/103	0/5/6/6
45	CHL	y	609	-	4/4/20/26	7/39/137/137	-
29	CLA	g	603	-	1/1/15/20	17/37/115/115	-
29	CLA	y	612	-	1/1/15/20	11/37/115/115	-
31	BCR	A	411	-	-	11/29/63/63	0/2/2/2
29	CLA	S	602	-	1/1/14/20	12/31/109/115	-
39	LHG	d	409	-	-	29/53/53/53	-
43	HEM	f	101	7,6	-	2/12/54/54	-
46	LUT	Y	621	-	-	2/29/67/67	0/2/2/2
29	CLA	G	610	-	1/1/15/20	13/37/115/115	-
48	NEX	N	623	-	-	2/27/83/83	0/3/3/3
44	RRX	h	101	-	1/1/11/25	10/29/65/65	0/2/2/2
31	BCR	D	404	-	-	11/29/63/63	0/2/2/2

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Mol	Type	Chain	Res	Link	Chirals	Torsions	Rings
29	CLA	N	602	-	1/1/15/20	17/37/115/115	-
29	CLA	b	610	-	1/1/15/20	15/37/115/115	-
45	CHL	Y	601	24	4/4/20/26	8/39/137/137	-
31	BCR	c	515	-	-	11/29/63/63	0/2/2/2
29	CLA	R	612	-	1/1/14/20	13/31/109/115	-
31	BCR	c	514	-	-	12/29/63/63	0/2/2/2
29	CLA	r	611	-	1/1/11/20	5/15/93/115	-
40	LMK	c	627	-	1/1/6/6	9/46/46/60	-
51	SPH	Y	625	-	-	10/21/21/21	-
29	CLA	B	613	-	1/1/15/20	15/37/115/115	-
29	CLA	b	609	-	1/1/15/20	15/37/115/115	-
39	LHG	n	624	-	-	27/53/53/53	-
45	CHL	n	606	-	4/4/20/26	5/39/137/137	-
29	CLA	d	403	-	1/1/15/20	13/37/115/115	-
29	CLA	r	608	-	1/1/14/20	13/31/109/115	-
29	CLA	b	603	-	1/1/15/20	16/37/115/115	-
29	CLA	s	602	-	1/1/14/20	10/31/109/115	-
47	XAT	y	622	-	1/1/12/26	1/31/93/93	0/4/4/4
50	3PH	s	626	-	-	27/49/49/49	-
33	LMG	H	102	-	-	12/43/63/70	0/1/1/1
46	LUT	S	621	-	-	4/29/67/67	0/2/2/2
29	CLA	b	605	-	1/1/15/20	20/37/115/115	-
29	CLA	B	604	-	1/1/15/20	17/37/115/115	-
29	CLA	b	617	-	1/1/15/20	16/37/115/115	-
29	CLA	Y	612	-	1/1/15/20	10/37/115/115	-
45	CHL	s	601	23	3/3/16/26	7/15/113/137	-
29	CLA	B	617	-	1/1/15/20	17/37/115/115	-
29	CLA	s	612	-	1/1/11/20	6/13/91/115	-
38	DGD	C	523	-	-	17/55/95/95	0/2/2/2
29	CLA	B	603	-	1/1/15/20	16/37/115/115	-
31	BCR	d	404	-	-	12/29/63/63	0/2/2/2
29	CLA	B	616	-	1/1/15/20	10/37/115/115	-
29	CLA	R	611	-	1/1/11/20	5/15/93/115	-
31	BCR	c	517	-	-	10/29/63/63	0/2/2/2

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Mol	Type	Chain	Res	Link	Chirals	Torsions	Rings
29	CLA	g	610	-	1/1/15/20	11/37/115/115	-
42	PL9	D	405	-	-	12/53/73/73	0/1/1/1

All (1846) bond length outliers are listed below:

Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
35	B	620	C7Z	C25-C26	16.16	1.62	1.34
35	b	620	C7Z	C25-C26	16.13	1.62	1.34
44	h	101	RRX	C26-C25	15.49	1.61	1.34
44	H	101	RRX	C26-C25	15.48	1.61	1.34
35	b	620	C7Z	C5-C6	15.32	1.61	1.34
35	B	620	C7Z	C5-C6	15.20	1.60	1.34
46	s	620	LUT	C24-C25	14.79	1.51	1.33
44	h	101	RRX	C5-C6	14.75	1.60	1.34
44	H	101	RRX	C5-C6	14.75	1.60	1.34
46	n	620	LUT	C24-C25	14.61	1.51	1.33
46	S	620	LUT	C24-C25	14.60	1.51	1.33
46	G	621	LUT	C24-C25	14.56	1.51	1.33
46	g	621	LUT	C24-C25	14.53	1.51	1.33
46	N	621	LUT	C24-C25	14.51	1.51	1.33
46	g	620	LUT	C24-C25	14.50	1.51	1.33
46	y	621	LUT	C24-C25	14.44	1.51	1.33
46	Y	620	LUT	C24-C25	14.43	1.51	1.33
46	R	620	LUT	C24-C25	14.42	1.51	1.33
46	N	620	LUT	C24-C25	14.41	1.51	1.33
46	r	620	LUT	C24-C25	14.39	1.51	1.33
46	G	620	LUT	C24-C25	14.37	1.51	1.33
46	y	620	LUT	C24-C25	14.32	1.51	1.33
46	Y	621	LUT	C24-C25	14.32	1.51	1.33
46	n	621	LUT	C24-C25	14.28	1.50	1.33
46	s	621	LUT	C24-C25	14.21	1.50	1.33
46	S	621	LUT	C24-C25	14.17	1.50	1.33
35	B	620	C7Z	C24-C23	11.64	1.72	1.52
35	b	620	C7Z	C24-C23	11.61	1.72	1.52
35	B	620	C7Z	C22-C23	-10.94	1.36	1.52
35	b	620	C7Z	C22-C23	-10.87	1.36	1.52
35	B	620	C7Z	C2-C3	-10.53	1.37	1.52
35	b	620	C7Z	C2-C3	-10.31	1.37	1.52
44	h	101	RRX	C29-C28	-10.10	1.37	1.52
44	H	101	RRX	C29-C28	-10.03	1.37	1.52
35	b	620	C7Z	C4-C3	8.49	1.67	1.52
35	B	620	C7Z	C4-C3	8.37	1.66	1.52

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
44	h	101	RRX	C27-C28	8.11	1.66	1.52
44	H	101	RRX	C27-C28	8.11	1.66	1.52
48	R	622	NEX	C10-C9	-8.03	1.25	1.35
48	R	622	NEX	C30-C29	-7.66	1.25	1.35
48	R	622	NEX	C14-C13	-7.62	1.25	1.35
48	y	623	NEX	C30-C29	-7.59	1.25	1.35
48	y	623	NEX	C10-C9	-7.55	1.25	1.35
48	n	623	NEX	C10-C9	-7.55	1.25	1.35
48	g	623	NEX	C34-C33	-7.54	1.25	1.35
48	g	623	NEX	C14-C13	-7.54	1.25	1.35
48	n	623	NEX	C14-C13	-7.53	1.25	1.35
48	n	623	NEX	C34-C33	-7.52	1.25	1.35
48	y	623	NEX	C14-C13	-7.51	1.25	1.35
48	R	622	NEX	C34-C33	-7.49	1.25	1.35
48	y	623	NEX	C34-C33	-7.48	1.25	1.35
48	Y	623	NEX	C30-C29	-7.48	1.25	1.35
48	s	623	NEX	C10-C9	-7.48	1.25	1.35
48	r	623	NEX	C14-C13	-7.47	1.25	1.35
48	r	623	NEX	C30-C29	-7.47	1.25	1.35
48	Y	623	NEX	C14-C13	-7.47	1.25	1.35
48	N	623	NEX	C34-C33	-7.46	1.25	1.35
48	N	623	NEX	C10-C9	-7.45	1.25	1.35
48	Y	623	NEX	C34-C33	-7.45	1.25	1.35
48	G	623	NEX	C34-C33	-7.44	1.25	1.35
48	S	622	NEX	C10-C9	-7.43	1.25	1.35
48	g	623	NEX	C30-C29	-7.42	1.25	1.35
48	G	623	NEX	C30-C29	-7.42	1.25	1.35
48	N	623	NEX	C30-C29	-7.42	1.25	1.35
48	n	623	NEX	C30-C29	-7.41	1.26	1.35
48	N	623	NEX	C14-C13	-7.41	1.26	1.35
48	r	623	NEX	C34-C33	-7.40	1.26	1.35
48	g	623	NEX	C10-C9	-7.39	1.26	1.35
48	G	623	NEX	C14-C13	-7.38	1.26	1.35
48	s	623	NEX	C14-C13	-7.38	1.26	1.35
48	s	623	NEX	C30-C29	-7.35	1.26	1.35
48	Y	623	NEX	C10-C9	-7.35	1.26	1.35
31	c	516	BCR	C10-C9	7.34	1.45	1.35
48	S	622	NEX	C14-C13	-7.31	1.26	1.35
48	s	623	NEX	C34-C33	-7.28	1.26	1.35
31	C	516	BCR	C10-C9	7.27	1.45	1.35
31	D	404	BCR	C10-C9	7.26	1.45	1.35
48	S	622	NEX	C34-C33	-7.25	1.26	1.35

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
48	S	622	NEX	C30-C29	-7.24	1.26	1.35
48	r	623	NEX	C10-C9	-7.23	1.26	1.35
48	G	623	NEX	C10-C9	-7.16	1.26	1.35
31	C	514	BCR	C10-C9	7.13	1.45	1.35
31	d	404	BCR	C10-C9	7.11	1.45	1.35
40	C	527	LMK	O3-C4	7.09	1.43	1.22
40	c	627	LMK	O3-C4	7.07	1.43	1.22
31	c	514	BCR	C10-C9	7.05	1.45	1.35
31	C	515	BCR	C10-C9	6.99	1.45	1.35
31	B	619	BCR	C10-C9	6.91	1.44	1.35
31	b	618	BCR	C10-C9	6.89	1.44	1.35
31	C	517	BCR	C10-C9	6.84	1.44	1.35
48	y	623	NEX	C35-C15	-6.81	1.18	1.36
31	c	515	BCR	C10-C9	6.81	1.44	1.35
31	A	411	BCR	C10-C9	6.81	1.44	1.35
48	R	622	NEX	C35-C15	-6.80	1.18	1.36
48	n	623	NEX	C35-C15	-6.80	1.18	1.36
31	b	619	BCR	C10-C9	6.80	1.44	1.35
31	B	618	BCR	C10-C9	6.79	1.44	1.35
48	g	623	NEX	C35-C15	-6.79	1.18	1.36
31	a	411	BCR	C10-C9	6.78	1.44	1.35
48	N	623	NEX	C35-C15	-6.75	1.18	1.36
48	Y	623	NEX	C35-C15	-6.75	1.18	1.36
48	r	623	NEX	C35-C15	-6.72	1.18	1.36
48	G	623	NEX	C35-C15	-6.72	1.18	1.36
48	s	623	NEX	C35-C15	-6.68	1.18	1.36
48	S	622	NEX	C35-C15	-6.67	1.18	1.36
31	c	517	BCR	C10-C9	6.53	1.44	1.35
29	B	616	CLA	MG-NA	6.47	2.21	2.06
29	S	605	CLA	MG-NA	6.46	2.21	2.06
29	s	605	CLA	MG-NA	6.46	2.21	2.06
29	s	603	CLA	MG-NA	6.45	2.21	2.06
29	r	612	CLA	MG-NA	6.45	2.21	2.06
29	b	616	CLA	MG-NA	6.44	2.21	2.06
44	h	101	RRX	C2-C3	-6.43	1.36	1.52
44	H	101	RRX	C2-C3	-6.43	1.36	1.52
29	r	611	CLA	MG-NA	6.42	2.21	2.06
29	N	613	CLA	MG-NA	6.42	2.21	2.06
29	B	605	CLA	MG-NA	6.41	2.21	2.06
29	G	612	CLA	MG-NA	6.41	2.21	2.06
29	s	609	CLA	MG-NA	6.41	2.21	2.06
29	n	611	CLA	MG-NA	6.41	2.21	2.06

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
29	R	611	CLA	MG-NA	6.40	2.21	2.06
29	R	603	CLA	MG-NA	6.40	2.21	2.06
29	b	602	CLA	MG-NA	6.40	2.21	2.06
29	R	612	CLA	MG-NA	6.39	2.21	2.06
29	b	605	CLA	MG-NA	6.39	2.21	2.06
29	S	603	CLA	MG-NA	6.39	2.21	2.06
29	Y	608	CLA	MG-NA	6.39	2.21	2.06
29	G	613	CLA	MG-NA	6.39	2.21	2.06
29	C	503	CLA	MG-NA	6.39	2.21	2.06
29	C	512	CLA	MG-NA	6.39	2.21	2.06
29	g	611	CLA	MG-NA	6.39	2.21	2.06
29	r	603	CLA	MG-NA	6.39	2.21	2.06
29	B	611	CLA	MG-NA	6.39	2.21	2.06
29	n	614	CLA	MG-NA	6.39	2.21	2.06
29	n	612	CLA	MG-NA	6.39	2.21	2.06
29	g	612	CLA	MG-NA	6.38	2.21	2.06
29	c	503	CLA	MG-NA	6.38	2.21	2.06
29	S	612	CLA	MG-NA	6.38	2.21	2.06
29	Y	612	CLA	MG-NA	6.38	2.21	2.06
29	s	613	CLA	MG-NA	6.38	2.21	2.06
29	S	611	CLA	MG-NA	6.38	2.21	2.06
29	n	613	CLA	MG-NA	6.38	2.21	2.06
29	r	608	CLA	MG-NA	6.37	2.21	2.06
29	b	611	CLA	MG-NA	6.37	2.21	2.06
29	C	501	CLA	MG-NA	6.37	2.21	2.06
29	s	612	CLA	MG-NA	6.37	2.21	2.06
29	S	617	CLA	MG-NA	6.37	2.21	2.06
29	G	611	CLA	MG-NA	6.37	2.21	2.06
29	d	403	CLA	MG-NA	6.37	2.21	2.06
29	R	610	CLA	MG-NA	6.37	2.21	2.06
29	R	602	CLA	MG-NA	6.36	2.21	2.06
29	B	602	CLA	MG-NA	6.36	2.21	2.06
29	R	613	CLA	MG-NA	6.36	2.21	2.06
29	S	609	CLA	MG-NA	6.36	2.21	2.06
29	g	613	CLA	MG-NA	6.36	2.21	2.06
29	N	611	CLA	MG-NA	6.36	2.21	2.06
29	C	513	CLA	MG-NA	6.36	2.21	2.06
29	N	612	CLA	MG-NA	6.36	2.21	2.06
29	c	507	CLA	MG-NA	6.35	2.21	2.06
29	c	501	CLA	MG-NA	6.35	2.21	2.06
29	D	403	CLA	MG-NA	6.35	2.21	2.06
29	G	604	CLA	MG-NA	6.35	2.21	2.06

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
29	r	602	CLA	MG-NA	6.35	2.21	2.06
29	g	614	CLA	MG-NA	6.35	2.21	2.06
29	y	611	CLA	MG-NA	6.35	2.21	2.06
29	y	612	CLA	MG-NA	6.34	2.21	2.06
29	c	511	CLA	MG-NA	6.34	2.21	2.06
29	y	613	CLA	MG-NA	6.34	2.21	2.06
29	c	502	CLA	MG-NA	6.34	2.21	2.06
29	g	604	CLA	MG-NA	6.34	2.21	2.06
29	C	511	CLA	MG-NA	6.34	2.21	2.06
29	Y	614	CLA	MG-NA	6.33	2.21	2.06
29	C	507	CLA	MG-NA	6.33	2.21	2.06
29	N	614	CLA	MG-NA	6.33	2.21	2.06
29	c	512	CLA	MG-NA	6.33	2.21	2.06
29	s	617	CLA	MG-NA	6.33	2.21	2.06
29	B	609	CLA	MG-NA	6.32	2.21	2.06
29	S	613	CLA	MG-NA	6.32	2.21	2.06
29	G	614	CLA	MG-NA	6.32	2.21	2.06
29	S	610	CLA	MG-NA	6.32	2.21	2.06
29	A	407	CLA	MG-NA	6.32	2.21	2.06
29	r	610	CLA	MG-NA	6.31	2.21	2.06
48	Y	623	NEX	C31-C32	-6.31	1.18	1.34
29	R	608	CLA	MG-NA	6.31	2.21	2.06
48	G	623	NEX	C11-C12	-6.31	1.18	1.34
29	s	610	CLA	MG-NA	6.31	2.21	2.06
29	Y	611	CLA	MG-NA	6.30	2.21	2.06
29	n	602	CLA	MG-NA	6.30	2.21	2.06
29	b	604	CLA	MG-NA	6.30	2.21	2.06
29	Y	613	CLA	MG-NA	6.30	2.21	2.06
29	c	506	CLA	MG-NA	6.30	2.21	2.06
29	g	602	CLA	MG-NA	6.30	2.21	2.06
29	s	611	CLA	MG-NA	6.30	2.21	2.06
29	y	614	CLA	MG-NA	6.30	2.21	2.06
29	y	602	CLA	MG-NA	6.30	2.21	2.06
48	s	623	NEX	C11-C12	-6.30	1.18	1.34
29	b	603	CLA	MG-NA	6.30	2.21	2.06
29	g	603	CLA	MG-NA	6.30	2.21	2.06
29	y	608	CLA	MG-NA	6.30	2.21	2.06
29	N	603	CLA	MG-NA	6.30	2.21	2.06
29	B	604	CLA	MG-NA	6.29	2.21	2.06
48	n	623	NEX	C11-C12	-6.29	1.18	1.34
29	Y	603	CLA	MG-NA	6.29	2.21	2.06
29	y	603	CLA	MG-NA	6.28	2.21	2.06

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
29	n	604	CLA	MG-NA	6.28	2.21	2.06
48	Y	623	NEX	C11-C12	-6.28	1.18	1.34
29	S	602	CLA	MG-NA	6.28	2.21	2.06
29	b	609	CLA	MG-NA	6.28	2.21	2.06
29	C	506	CLA	MG-NA	6.28	2.21	2.06
29	N	610	CLA	MG-NA	6.28	2.21	2.06
29	a	407	CLA	MG-NA	6.27	2.21	2.06
29	G	603	CLA	MG-NA	6.27	2.21	2.06
29	G	602	CLA	MG-NA	6.27	2.21	2.06
48	y	623	NEX	C11-C12	-6.27	1.18	1.34
29	N	604	CLA	MG-NA	6.26	2.21	2.06
29	N	602	CLA	MG-NA	6.26	2.21	2.06
48	y	623	NEX	C31-C32	-6.26	1.18	1.34
29	C	502	CLA	MG-NA	6.26	2.21	2.06
48	n	623	NEX	C31-C32	-6.26	1.18	1.34
29	c	513	CLA	MG-NA	6.25	2.21	2.06
29	r	613	CLA	MG-NA	6.25	2.21	2.06
48	s	623	NEX	C31-C32	-6.25	1.18	1.34
48	r	623	NEX	C31-C32	-6.25	1.18	1.34
29	B	603	CLA	MG-NA	6.25	2.21	2.06
29	C	509	CLA	MG-NA	6.25	2.21	2.06
29	B	607	CLA	MG-NA	6.25	2.21	2.06
29	g	610	CLA	MG-NA	6.24	2.21	2.06
48	N	623	NEX	C31-C32	-6.24	1.18	1.34
48	R	622	NEX	C31-C32	-6.24	1.18	1.34
29	r	604	CLA	MG-NA	6.24	2.21	2.06
29	s	604	CLA	MG-NA	6.24	2.21	2.06
29	Y	602	CLA	MG-NA	6.24	2.21	2.06
29	Y	604	CLA	MG-NA	6.23	2.21	2.06
29	S	614	CLA	MG-NA	6.23	2.21	2.06
29	C	505	CLA	MG-NA	6.23	2.21	2.06
48	g	623	NEX	C31-C32	-6.23	1.18	1.34
29	B	614	CLA	MG-NA	6.22	2.21	2.06
48	N	623	NEX	C11-C12	-6.22	1.18	1.34
29	A	406	CLA	MG-NA	6.22	2.21	2.06
48	G	623	NEX	C31-C32	-6.22	1.18	1.34
48	g	623	NEX	C11-C12	-6.22	1.18	1.34
29	n	603	CLA	MG-NA	6.21	2.21	2.06
48	S	622	NEX	C11-C12	-6.21	1.18	1.34
29	R	609	CLA	MG-NA	6.21	2.21	2.06
29	b	607	CLA	MG-NA	6.21	2.21	2.06
29	B	608	CLA	MG-NA	6.21	2.21	2.06

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
29	d	402	CLA	MG-NA	6.20	2.21	2.06
29	n	610	CLA	MG-NA	6.20	2.21	2.06
29	y	604	CLA	MG-NA	6.20	2.21	2.06
29	b	614	CLA	MG-NA	6.20	2.21	2.06
29	b	608	CLA	MG-NA	6.20	2.21	2.06
48	r	623	NEX	C11-C12	-6.20	1.18	1.34
29	Y	610	CLA	MG-NA	6.20	2.21	2.06
29	y	610	CLA	MG-NA	6.19	2.21	2.06
29	c	505	CLA	MG-NA	6.19	2.21	2.06
29	A	410	CLA	MG-NA	6.19	2.21	2.06
29	s	614	CLA	MG-NA	6.18	2.21	2.06
29	D	402	CLA	MG-NA	6.18	2.21	2.06
29	C	508	CLA	MG-NA	6.18	2.21	2.06
29	r	609	CLA	MG-NA	6.18	2.21	2.06
29	s	602	CLA	MG-NA	6.18	2.20	2.06
29	b	612	CLA	MG-NA	6.17	2.20	2.06
29	R	604	CLA	MG-NA	6.17	2.20	2.06
29	a	406	CLA	MG-NA	6.17	2.20	2.06
48	S	622	NEX	C31-C32	-6.17	1.18	1.34
29	b	615	CLA	MG-NA	6.16	2.20	2.06
29	C	510	CLA	MG-NA	6.16	2.20	2.06
29	c	509	CLA	MG-NA	6.16	2.20	2.06
29	a	410	CLA	MG-NA	6.15	2.20	2.06
29	S	604	CLA	MG-NA	6.15	2.20	2.06
29	B	612	CLA	MG-NA	6.14	2.20	2.06
29	c	510	CLA	MG-NA	6.13	2.20	2.06
29	B	610	CLA	MG-NA	6.13	2.20	2.06
29	B	615	CLA	MG-NA	6.13	2.20	2.06
29	c	508	CLA	MG-NA	6.08	2.20	2.06
29	b	617	CLA	MG-NA	6.08	2.20	2.06
29	B	617	CLA	MG-NA	6.06	2.20	2.06
29	C	504	CLA	MG-NA	6.06	2.20	2.06
29	b	610	CLA	MG-NA	6.06	2.20	2.06
29	a	405	CLA	MG-NA	6.05	2.20	2.06
29	b	613	CLA	MG-NA	6.05	2.20	2.06
29	A	405	CLA	MG-NA	6.04	2.20	2.06
29	G	610	CLA	MG-NA	6.04	2.20	2.06
29	c	504	CLA	MG-NA	6.03	2.20	2.06
48	R	622	NEX	C11-C12	-6.03	1.19	1.34
29	B	613	CLA	MG-NA	6.02	2.20	2.06
29	b	606	CLA	MG-NA	6.00	2.20	2.06
29	B	606	CLA	MG-NA	5.98	2.20	2.06

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
44	h	101	RRX	C1-C6	-5.88	1.45	1.53
48	R	622	NEX	C7-C8	5.83	1.41	1.32
31	D	404	BCR	C24-C23	5.79	1.50	1.33
31	d	404	BCR	C24-C23	5.78	1.50	1.33
44	H	101	RRX	C1-C6	-5.76	1.45	1.53
31	c	516	BCR	C24-C23	5.75	1.50	1.33
31	B	619	BCR	C24-C23	5.73	1.50	1.33
31	C	516	BCR	C24-C23	5.70	1.50	1.33
35	b	620	C7Z	C12-C13	5.70	1.58	1.45
31	c	517	BCR	C24-C23	5.66	1.50	1.33
31	C	517	BCR	C24-C23	5.65	1.50	1.33
35	B	620	C7Z	C12-C13	5.65	1.58	1.45
31	A	411	BCR	C24-C23	5.65	1.50	1.33
31	b	619	BCR	C24-C23	5.61	1.50	1.33
31	C	514	BCR	C24-C23	5.60	1.50	1.33
31	a	411	BCR	C24-C23	5.60	1.50	1.33
44	h	101	RRX	C30-C25	-5.58	1.46	1.53
31	C	515	BCR	C24-C23	5.58	1.49	1.33
44	H	101	RRX	C30-C25	-5.57	1.46	1.53
31	c	514	BCR	C24-C23	5.56	1.49	1.33
48	s	623	NEX	C7-C8	5.56	1.41	1.32
31	b	618	BCR	C24-C23	5.53	1.49	1.33
31	c	515	BCR	C24-C23	5.52	1.49	1.33
48	n	623	NEX	C7-C8	5.51	1.41	1.32
48	r	623	NEX	C7-C8	5.49	1.41	1.32
31	B	618	BCR	C24-C23	5.45	1.49	1.33
48	y	623	NEX	C7-C8	5.44	1.41	1.32
48	R	622	NEX	C28-C29	-5.32	1.34	1.45
31	c	517	BCR	C11-C12	-5.31	1.20	1.34
48	r	623	NEX	C28-C29	-5.30	1.34	1.45
31	C	517	BCR	C11-C12	-5.30	1.20	1.34
31	A	411	BCR	C11-C12	-5.29	1.20	1.34
48	N	623	NEX	C7-C8	5.29	1.40	1.32
31	B	618	BCR	C11-C12	-5.29	1.21	1.34
31	a	411	BCR	C11-C12	-5.28	1.21	1.34
31	b	619	BCR	C11-C12	-5.28	1.21	1.34
31	B	619	BCR	C11-C12	-5.27	1.21	1.34
31	b	618	BCR	C11-C12	-5.27	1.21	1.34
48	s	623	NEX	C28-C29	-5.27	1.34	1.45
31	c	515	BCR	C11-C12	-5.25	1.21	1.34
48	n	623	NEX	C28-C29	-5.25	1.34	1.45
48	N	623	NEX	C28-C29	-5.24	1.34	1.45

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
48	G	623	NEX	C7-C8	5.23	1.40	1.32
48	g	623	NEX	C7-C8	5.23	1.40	1.32
31	c	514	BCR	C11-C12	-5.22	1.21	1.34
48	Y	623	NEX	C28-C29	-5.22	1.34	1.45
48	y	623	NEX	C28-C29	-5.22	1.34	1.45
31	C	515	BCR	C11-C12	-5.21	1.21	1.34
48	g	623	NEX	C28-C29	-5.20	1.34	1.45
48	G	623	NEX	C28-C29	-5.19	1.34	1.45
48	S	622	NEX	C28-C29	-5.16	1.34	1.45
48	S	622	NEX	C7-C8	5.16	1.40	1.32
35	B	620	C7Z	C1-C6	-5.16	1.46	1.53
31	C	514	BCR	C11-C12	-5.15	1.21	1.34
31	d	404	BCR	C11-C12	-5.13	1.21	1.34
44	H	101	RRX	C2-C1	5.12	1.65	1.54
31	C	516	BCR	C11-C12	-5.09	1.21	1.34
44	h	101	RRX	C2-C1	5.09	1.65	1.54
31	c	516	BCR	C11-C12	-5.08	1.21	1.34
31	D	404	BCR	C11-C12	-5.00	1.21	1.34
35	b	620	C7Z	C28-C29	4.97	1.56	1.45
35	B	620	C7Z	C28-C29	4.96	1.56	1.45
35	b	620	C7Z	C1-C6	-4.94	1.47	1.53
48	Y	623	NEX	C7-C8	4.91	1.40	1.32
44	H	101	RRX	C19-C18	4.90	1.56	1.45
44	h	101	RRX	C19-C18	4.89	1.56	1.45
44	H	101	RRX	C8-C9	4.82	1.56	1.45
43	f	101	HEM	C3C-C2C	-4.74	1.33	1.40
44	h	101	RRX	C8-C9	4.73	1.56	1.45
43	F	101	HEM	C3C-C2C	-4.72	1.33	1.40
35	B	620	C7Z	C24-C25	-4.69	1.43	1.51
35	b	620	C7Z	C24-C25	-4.63	1.43	1.51
35	B	620	C7Z	C32-C33	4.61	1.55	1.45
35	b	620	C7Z	C32-C33	4.61	1.55	1.45
31	A	411	BCR	C16-C17	-4.46	1.29	1.43
31	a	411	BCR	C16-C17	-4.45	1.29	1.43
44	h	101	RRX	C27-C26	-4.43	1.44	1.51
31	c	517	BCR	C16-C17	-4.42	1.29	1.43
35	b	620	C7Z	C8-C9	4.41	1.55	1.45
44	H	101	RRX	C27-C26	-4.40	1.44	1.51
35	b	620	C7Z	C31-C30	4.39	1.57	1.43
31	c	515	BCR	C16-C17	-4.39	1.29	1.43
31	C	517	BCR	C16-C17	-4.38	1.29	1.43
35	B	620	C7Z	C8-C9	4.37	1.55	1.45

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
35	B	620	C7Z	C31-C30	4.37	1.57	1.43
35	B	620	C7Z	C4-C5	-4.36	1.44	1.51
31	B	618	BCR	C16-C17	-4.34	1.30	1.43
38	c	518	DGD	O1G-C1A	4.33	1.46	1.33
38	C	523	DGD	O1G-C1A	4.33	1.46	1.33
31	b	618	BCR	C16-C17	-4.33	1.30	1.43
38	c	523	DGD	O1G-C1A	4.33	1.46	1.33
31	C	515	BCR	C16-C17	-4.32	1.30	1.43
31	b	619	BCR	C16-C17	-4.31	1.30	1.43
38	C	518	DGD	O1G-C1A	4.30	1.45	1.33
31	c	514	BCR	C16-C17	-4.30	1.30	1.43
44	H	101	RRX	C12-C13	4.29	1.55	1.45
35	b	620	C7Z	C4-C5	-4.29	1.44	1.51
31	C	514	BCR	C16-C17	-4.27	1.30	1.43
31	B	619	BCR	C16-C17	-4.26	1.30	1.43
44	h	101	RRX	C3-C4	4.25	1.65	1.52
42	d	405	PL9	C7-C3	-4.25	1.47	1.51
44	H	101	RRX	C3-C4	4.23	1.65	1.52
38	C	520	DGD	O1G-C1A	4.22	1.45	1.33
31	d	404	BCR	C16-C17	-4.21	1.30	1.43
31	C	516	BCR	C16-C17	-4.20	1.30	1.43
31	c	516	BCR	C16-C17	-4.20	1.30	1.43
35	B	620	C7Z	C11-C10	4.19	1.56	1.43
38	C	519	DGD	O1G-C1A	4.19	1.45	1.33
44	h	101	RRX	C12-C13	4.18	1.54	1.45
31	D	404	BCR	C16-C17	-4.17	1.30	1.43
38	c	520	DGD	O1G-C1A	4.17	1.45	1.33
38	c	519	DGD	O1G-C1A	4.16	1.45	1.33
35	b	620	C7Z	C11-C10	4.15	1.56	1.43
29	c	506	CLA	MG-ND	-4.12	1.97	2.05
29	b	609	CLA	MG-ND	-4.10	1.97	2.05
29	b	612	CLA	MG-ND	-4.10	1.97	2.05
29	b	604	CLA	MG-ND	-4.09	1.97	2.05
29	C	506	CLA	MG-ND	-4.07	1.97	2.05
29	B	609	CLA	MG-ND	-4.05	1.97	2.05
29	S	603	CLA	MG-ND	-4.02	1.97	2.05
42	D	405	PL9	C7-C3	-4.02	1.47	1.51
29	B	604	CLA	MG-ND	-4.02	1.97	2.05
40	C	527	LMK	O2-C4	4.01	1.43	1.30
29	B	612	CLA	MG-ND	-3.99	1.97	2.05
29	c	507	CLA	MG-ND	-3.98	1.97	2.05
29	C	508	CLA	MG-ND	-3.98	1.97	2.05

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
29	B	615	CLA	MG-ND	-3.96	1.97	2.05
29	s	603	CLA	MG-ND	-3.95	1.98	2.05
29	C	505	CLA	MG-ND	-3.95	1.98	2.05
29	b	617	CLA	MG-ND	-3.95	1.98	2.05
29	C	507	CLA	MG-ND	-3.94	1.98	2.05
40	c	627	LMK	O2-C4	3.94	1.43	1.30
29	c	505	CLA	MG-ND	-3.94	1.98	2.05
29	B	607	CLA	MG-ND	-3.93	1.98	2.05
29	B	617	CLA	MG-ND	-3.93	1.98	2.05
29	b	615	CLA	MG-ND	-3.92	1.98	2.05
29	c	508	CLA	MG-ND	-3.92	1.98	2.05
44	h	101	RRX	C20-C21	3.92	1.55	1.43
29	S	605	CLA	MG-ND	-3.91	1.98	2.05
29	Y	611	CLA	MG-ND	-3.90	1.98	2.05
29	A	410	CLA	MG-ND	-3.90	1.98	2.05
29	B	613	CLA	MG-ND	-3.90	1.98	2.05
29	B	605	CLA	MG-ND	-3.90	1.98	2.05
29	B	603	CLA	MG-ND	-3.90	1.98	2.05
29	s	611	CLA	MG-ND	-3.88	1.98	2.05
43	f	101	HEM	C3C-CAC	3.88	1.55	1.47
29	y	608	CLA	MG-ND	-3.88	1.98	2.05
44	H	101	RRX	C20-C21	3.87	1.55	1.43
29	C	502	CLA	MG-ND	-3.87	1.98	2.05
44	H	101	RRX	C23-C22	3.87	1.54	1.45
29	D	402	CLA	MG-ND	-3.87	1.98	2.05
29	N	612	CLA	MG-ND	-3.87	1.98	2.05
29	b	613	CLA	MG-ND	-3.87	1.98	2.05
29	b	603	CLA	MG-ND	-3.87	1.98	2.05
29	a	410	CLA	MG-ND	-3.86	1.98	2.05
29	b	607	CLA	MG-ND	-3.86	1.98	2.05
29	Y	612	CLA	MG-ND	-3.86	1.98	2.05
29	S	611	CLA	MG-ND	-3.86	1.98	2.05
29	a	405	CLA	MG-ND	-3.86	1.98	2.05
44	h	101	RRX	C23-C22	3.85	1.54	1.45
29	g	604	CLA	MG-ND	-3.85	1.98	2.05
29	b	605	CLA	MG-ND	-3.85	1.98	2.05
29	d	403	CLA	MG-ND	-3.85	1.98	2.05
29	c	513	CLA	MG-ND	-3.85	1.98	2.05
43	F	101	HEM	C3C-CAC	3.85	1.55	1.47
29	N	604	CLA	MG-ND	-3.85	1.98	2.05
29	r	609	CLA	MG-ND	-3.85	1.98	2.05
29	R	604	CLA	MG-ND	-3.84	1.98	2.05

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
29	r	604	CLA	MG-ND	-3.84	1.98	2.05
29	s	605	CLA	MG-ND	-3.84	1.98	2.05
29	D	403	CLA	MG-ND	-3.84	1.98	2.05
29	c	502	CLA	MG-ND	-3.84	1.98	2.05
29	d	402	CLA	MG-ND	-3.84	1.98	2.05
29	B	606	CLA	MG-ND	-3.84	1.98	2.05
29	b	606	CLA	MG-ND	-3.84	1.98	2.05
29	s	610	CLA	MG-ND	-3.83	1.98	2.05
29	y	614	CLA	MG-ND	-3.83	1.98	2.05
29	c	509	CLA	MG-ND	-3.83	1.98	2.05
29	y	612	CLA	MG-ND	-3.83	1.98	2.05
29	c	501	CLA	MG-ND	-3.83	1.98	2.05
29	S	613	CLA	MG-ND	-3.83	1.98	2.05
29	R	603	CLA	MG-ND	-3.83	1.98	2.05
29	b	611	CLA	MG-ND	-3.83	1.98	2.05
29	C	509	CLA	MG-ND	-3.83	1.98	2.05
29	n	613	CLA	MG-ND	-3.83	1.98	2.05
29	S	610	CLA	MG-ND	-3.83	1.98	2.05
29	b	608	CLA	MG-ND	-3.83	1.98	2.05
29	C	512	CLA	MG-ND	-3.83	1.98	2.05
29	A	407	CLA	MG-ND	-3.83	1.98	2.05
29	C	513	CLA	MG-ND	-3.83	1.98	2.05
29	c	512	CLA	MG-ND	-3.83	1.98	2.05
29	G	612	CLA	MG-ND	-3.82	1.98	2.05
29	A	405	CLA	MG-ND	-3.82	1.98	2.05
29	r	612	CLA	MG-ND	-3.82	1.98	2.05
29	Y	608	CLA	MG-ND	-3.82	1.98	2.05
29	A	406	CLA	MG-ND	-3.82	1.98	2.05
29	b	614	CLA	MG-ND	-3.82	1.98	2.05
29	R	609	CLA	MG-ND	-3.82	1.98	2.05
29	N	613	CLA	MG-ND	-3.81	1.98	2.05
29	n	602	CLA	MG-ND	-3.81	1.98	2.05
29	N	610	CLA	MG-ND	-3.81	1.98	2.05
29	N	602	CLA	MG-ND	-3.81	1.98	2.05
29	S	614	CLA	MG-ND	-3.81	1.98	2.05
29	g	612	CLA	MG-ND	-3.81	1.98	2.05
29	G	604	CLA	MG-ND	-3.80	1.98	2.05
29	Y	602	CLA	MG-ND	-3.80	1.98	2.05
29	Y	610	CLA	MG-ND	-3.80	1.98	2.05
29	s	614	CLA	MG-ND	-3.80	1.98	2.05
29	S	612	CLA	MG-ND	-3.80	1.98	2.05
29	r	608	CLA	MG-ND	-3.80	1.98	2.05

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
29	a	406	CLA	MG-ND	-3.80	1.98	2.05
29	n	604	CLA	MG-ND	-3.80	1.98	2.05
29	B	611	CLA	MG-ND	-3.79	1.98	2.05
29	C	501	CLA	MG-ND	-3.79	1.98	2.05
29	s	612	CLA	MG-ND	-3.78	1.98	2.05
29	a	407	CLA	MG-ND	-3.78	1.98	2.05
29	b	610	CLA	MG-ND	-3.78	1.98	2.05
29	B	610	CLA	MG-ND	-3.78	1.98	2.05
29	S	604	CLA	MG-ND	-3.78	1.98	2.05
29	C	504	CLA	MG-ND	-3.78	1.98	2.05
29	s	613	CLA	MG-ND	-3.78	1.98	2.05
35	B	620	C7Z	C27-C26	3.78	1.58	1.45
29	y	604	CLA	MG-ND	-3.78	1.98	2.05
29	n	612	CLA	MG-ND	-3.78	1.98	2.05
29	B	608	CLA	MG-ND	-3.78	1.98	2.05
29	S	617	CLA	MG-ND	-3.78	1.98	2.05
29	r	603	CLA	MG-ND	-3.78	1.98	2.05
29	C	503	CLA	MG-ND	-3.78	1.98	2.05
29	R	613	CLA	MG-ND	-3.78	1.98	2.05
29	c	504	CLA	MG-ND	-3.77	1.98	2.05
29	R	611	CLA	MG-ND	-3.77	1.98	2.05
29	y	611	CLA	MG-ND	-3.77	1.98	2.05
29	c	510	CLA	MG-ND	-3.77	1.98	2.05
44	h	101	RRX	C15-C14	3.77	1.55	1.43
29	S	602	CLA	MG-ND	-3.77	1.98	2.05
29	y	610	CLA	MG-ND	-3.77	1.98	2.05
29	g	602	CLA	MG-ND	-3.77	1.98	2.05
29	N	614	CLA	MG-ND	-3.77	1.98	2.05
29	r	611	CLA	MG-ND	-3.77	1.98	2.05
29	c	503	CLA	MG-ND	-3.76	1.98	2.05
35	b	620	C7Z	C15-C14	3.76	1.55	1.43
29	G	613	CLA	MG-ND	-3.76	1.98	2.05
29	n	614	CLA	MG-ND	-3.76	1.98	2.05
29	g	610	CLA	MG-ND	-3.76	1.98	2.05
29	C	511	CLA	MG-ND	-3.76	1.98	2.05
29	R	608	CLA	MG-ND	-3.76	1.98	2.05
35	b	620	C7Z	C22-C21	3.76	1.66	1.54
29	R	612	CLA	MG-ND	-3.75	1.98	2.05
29	N	611	CLA	MG-ND	-3.75	1.98	2.05
29	Y	604	CLA	MG-ND	-3.75	1.98	2.05
29	r	602	CLA	MG-ND	-3.75	1.98	2.05
29	B	614	CLA	MG-ND	-3.75	1.98	2.05

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
29	N	603	CLA	MG-ND	-3.75	1.98	2.05
29	g	611	CLA	MG-ND	-3.75	1.98	2.05
29	r	613	CLA	MG-ND	-3.75	1.98	2.05
35	b	620	C7Z	C27-C26	3.75	1.58	1.45
35	B	620	C7Z	C22-C21	3.75	1.66	1.54
29	g	613	CLA	MG-ND	-3.75	1.98	2.05
44	H	101	RRX	C15-C14	3.74	1.55	1.43
29	R	610	CLA	MG-ND	-3.74	1.98	2.05
29	G	611	CLA	MG-ND	-3.74	1.98	2.05
29	y	602	CLA	MG-ND	-3.74	1.98	2.05
29	c	508	CLA	C1C-NC	-3.74	1.32	1.37
29	r	610	CLA	MG-ND	-3.74	1.98	2.05
29	s	604	CLA	MG-ND	-3.74	1.98	2.05
29	n	611	CLA	MG-ND	-3.74	1.98	2.05
29	s	617	CLA	MG-ND	-3.74	1.98	2.05
29	n	603	CLA	MG-ND	-3.73	1.98	2.05
29	Y	614	CLA	MG-ND	-3.73	1.98	2.05
29	Y	613	CLA	MG-ND	-3.73	1.98	2.05
29	G	602	CLA	MG-ND	-3.73	1.98	2.05
29	G	610	CLA	MG-ND	-3.73	1.98	2.05
29	B	602	CLA	MG-ND	-3.73	1.98	2.05
29	B	616	CLA	MG-ND	-3.72	1.98	2.05
29	s	609	CLA	MG-ND	-3.72	1.98	2.05
29	y	613	CLA	MG-ND	-3.72	1.98	2.05
29	R	602	CLA	MG-ND	-3.72	1.98	2.05
29	S	609	CLA	MG-ND	-3.72	1.98	2.05
35	B	620	C7Z	C15-C14	3.72	1.55	1.43
29	C	508	CLA	C1C-NC	-3.72	1.32	1.37
29	g	614	CLA	MG-ND	-3.71	1.98	2.05
29	G	614	CLA	MG-ND	-3.71	1.98	2.05
29	s	602	CLA	MG-ND	-3.71	1.98	2.05
29	b	602	CLA	MG-ND	-3.70	1.98	2.05
29	b	617	CLA	C1C-NC	-3.70	1.32	1.37
29	G	603	CLA	MG-ND	-3.70	1.98	2.05
29	n	610	CLA	MG-ND	-3.68	1.98	2.05
29	c	511	CLA	MG-ND	-3.68	1.98	2.05
29	B	617	CLA	C1C-NC	-3.67	1.32	1.37
29	b	616	CLA	MG-ND	-3.65	1.98	2.05
29	g	603	CLA	MG-ND	-3.65	1.98	2.05
36	b	623	DGA	OG2-CB1	3.64	1.44	1.34
36	B	625	DGA	OG2-CB1	3.63	1.44	1.34
29	Y	603	CLA	MG-ND	-3.63	1.98	2.05

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
29	c	502	CLA	C1C-NC	-3.62	1.32	1.37
29	b	612	CLA	C1C-NC	-3.61	1.32	1.37
35	B	620	C7Z	C2-C1	3.60	1.66	1.54
29	b	610	CLA	C1C-NC	-3.60	1.32	1.37
29	G	610	CLA	C1C-NC	-3.60	1.32	1.37
29	B	605	CLA	C1C-NC	-3.59	1.32	1.37
35	b	620	C7Z	C2-C1	3.59	1.66	1.54
29	A	406	CLA	C1C-NC	-3.58	1.32	1.37
29	b	611	CLA	C1C-NC	-3.58	1.32	1.37
35	B	620	C7Z	C35-C34	3.57	1.54	1.43
35	b	620	C7Z	C35-C34	3.57	1.54	1.43
29	y	603	CLA	MG-ND	-3.55	1.98	2.05
29	C	510	CLA	MG-ND	-3.55	1.98	2.05
29	B	606	CLA	C1C-NC	-3.54	1.32	1.37
35	b	620	C7Z	C7-C6	3.54	1.57	1.45
29	N	602	CLA	C1C-NC	-3.51	1.32	1.37
29	c	506	CLA	C1C-NC	-3.50	1.32	1.37
29	B	611	CLA	C1C-NC	-3.49	1.32	1.37
35	B	620	C7Z	C7-C6	3.49	1.57	1.45
29	c	504	CLA	C1C-NC	-3.48	1.32	1.37
29	b	606	CLA	C1C-NC	-3.48	1.32	1.37
35	B	620	C7Z	C38-C25	3.47	1.56	1.50
29	b	605	CLA	C1C-NC	-3.47	1.32	1.37
42	d	405	PL9	C3-C4	-3.46	1.43	1.49
29	B	612	CLA	C1C-NC	-3.46	1.32	1.37
29	a	405	CLA	C1C-NC	-3.46	1.32	1.37
29	C	506	CLA	C1C-NC	-3.45	1.32	1.37
45	N	608	CHL	CBB-CAB	3.44	1.52	1.29
35	b	620	C7Z	C38-C25	3.44	1.56	1.50
45	Y	601	CHL	CBB-CAB	3.43	1.52	1.29
45	n	608	CHL	CBB-CAB	3.42	1.52	1.29
45	y	601	CHL	CBB-CAB	3.42	1.52	1.29
29	C	504	CLA	C1C-NC	-3.42	1.32	1.37
45	n	609	CHL	CBB-CAB	3.42	1.52	1.29
29	a	410	CLA	C1C-NC	-3.42	1.32	1.37
29	B	610	CLA	C1C-NC	-3.41	1.32	1.37
29	a	407	CLA	C1C-NC	-3.41	1.32	1.37
36	C	524	DGA	OG2-CB1	3.41	1.43	1.34
29	C	502	CLA	C1C-NC	-3.41	1.32	1.37
29	B	614	CLA	CBB-CAB	3.40	1.51	1.29
29	a	406	CLA	C1C-NC	-3.40	1.32	1.37
45	N	609	CHL	CBB-CAB	3.40	1.51	1.29

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
36	c	524	DGA	OG2-CB1	3.40	1.43	1.34
29	r	608	CLA	C1C-NC	-3.40	1.32	1.37
29	b	608	CLA	C1C-NC	-3.40	1.32	1.37
29	b	614	CLA	CBB-CAB	3.39	1.51	1.29
45	Y	605	CHL	CBB-CAB	3.39	1.51	1.29
45	y	605	CHL	CBB-CAB	3.39	1.51	1.29
29	c	503	CLA	C1C-NC	-3.38	1.32	1.37
45	n	601	CHL	CBB-CAB	3.38	1.51	1.29
45	N	601	CHL	CBB-CAB	3.38	1.51	1.29
29	n	602	CLA	CBB-CAB	3.38	1.51	1.29
45	G	605	CHL	CBB-CAB	3.38	1.51	1.29
45	G	609	CHL	CBB-CAB	3.38	1.51	1.29
29	R	602	CLA	CBB-CAB	3.38	1.51	1.29
29	C	509	CLA	CBB-CAB	3.38	1.51	1.29
45	y	609	CHL	CBB-CAB	3.37	1.51	1.29
29	N	610	CLA	C1C-NC	-3.37	1.32	1.37
29	R	608	CLA	C1C-NC	-3.37	1.32	1.37
29	r	610	CLA	CBB-CAB	3.37	1.51	1.29
29	N	602	CLA	CBB-CAB	3.37	1.51	1.29
29	y	602	CLA	CBB-CAB	3.37	1.51	1.29
29	r	602	CLA	CBB-CAB	3.37	1.51	1.29
29	C	507	CLA	C1C-NC	-3.37	1.32	1.37
29	g	604	CLA	CBB-CAB	3.37	1.51	1.29
29	A	405	CLA	C1C-NC	-3.37	1.32	1.37
29	A	405	CLA	CBB-CAB	3.37	1.51	1.29
45	G	601	CHL	CBB-CAB	3.37	1.51	1.29
29	n	602	CLA	C1C-NC	-3.37	1.32	1.37
29	R	613	CLA	CBB-CAB	3.37	1.51	1.29
29	y	603	CLA	CBB-CAB	3.37	1.51	1.29
29	g	610	CLA	CBB-CAB	3.37	1.51	1.29
29	B	602	CLA	CBB-CAB	3.37	1.51	1.29
29	C	508	CLA	CBB-CAB	3.37	1.51	1.29
29	b	608	CLA	CBB-CAB	3.37	1.51	1.29
29	n	610	CLA	CBB-CAB	3.37	1.51	1.29
29	a	405	CLA	CBB-CAB	3.36	1.51	1.29
29	B	602	CLA	C1C-NC	-3.36	1.32	1.37
29	B	604	CLA	CBB-CAB	3.36	1.51	1.29
45	g	608	CHL	CBB-CAB	3.36	1.51	1.29
29	G	603	CLA	CBB-CAB	3.36	1.51	1.29
29	N	604	CLA	CBB-CAB	3.36	1.51	1.29
29	s	605	CLA	CBB-CAB	3.36	1.51	1.29
29	s	614	CLA	CBB-CAB	3.36	1.51	1.29

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
29	r	613	CLA	CBB-CAB	3.36	1.51	1.29
44	h	101	RRX	C11-C10	3.36	1.53	1.43
45	G	608	CHL	CBB-CAB	3.36	1.51	1.29
29	s	604	CLA	CBB-CAB	3.36	1.51	1.29
29	c	509	CLA	CBB-CAB	3.36	1.51	1.29
29	r	612	CLA	CBB-CAB	3.36	1.51	1.29
29	n	612	CLA	CBB-CAB	3.36	1.51	1.29
29	R	612	CLA	CBB-CAB	3.36	1.51	1.29
45	g	609	CHL	CBB-CAB	3.36	1.51	1.29
29	R	603	CLA	CBB-CAB	3.36	1.51	1.29
29	b	610	CLA	CBB-CAB	3.36	1.51	1.29
45	Y	609	CHL	CBB-CAB	3.36	1.51	1.29
29	B	611	CLA	CBB-CAB	3.36	1.51	1.29
29	c	508	CLA	CBB-CAB	3.36	1.51	1.29
29	S	605	CLA	CBB-CAB	3.36	1.51	1.29
29	B	615	CLA	CBB-CAB	3.36	1.51	1.29
44	H	101	RRX	C11-C10	3.36	1.53	1.43
29	B	603	CLA	CBB-CAB	3.36	1.51	1.29
29	d	403	CLA	CBB-CAB	3.36	1.51	1.29
29	R	611	CLA	CBB-CAB	3.36	1.51	1.29
29	g	613	CLA	CBB-CAB	3.36	1.51	1.29
29	G	602	CLA	CBB-CAB	3.36	1.51	1.29
29	B	608	CLA	C1C-NC	-3.36	1.32	1.37
29	Y	611	CLA	C1C-NC	-3.36	1.32	1.37
29	s	617	CLA	CBB-CAB	3.36	1.51	1.29
29	N	614	CLA	CBB-CAB	3.36	1.51	1.29
45	g	605	CHL	CBB-CAB	3.36	1.51	1.29
29	B	612	CLA	CBB-CAB	3.36	1.51	1.29
29	G	612	CLA	CBB-CAB	3.36	1.51	1.29
29	S	614	CLA	CBB-CAB	3.36	1.51	1.29
29	c	509	CLA	C1C-NC	-3.36	1.32	1.37
45	n	606	CHL	C4B-NB	3.36	1.38	1.35
29	b	604	CLA	CBB-CAB	3.35	1.51	1.29
29	c	507	CLA	CBB-CAB	3.35	1.51	1.29
29	A	410	CLA	C1C-NC	-3.35	1.32	1.37
29	n	604	CLA	CBB-CAB	3.35	1.51	1.29
29	B	613	CLA	C1C-NC	-3.35	1.32	1.37
29	y	608	CLA	CBB-CAB	3.35	1.51	1.29
45	n	605	CHL	CBB-CAB	3.35	1.51	1.29
29	c	512	CLA	CBB-CAB	3.35	1.51	1.29
29	B	608	CLA	CBB-CAB	3.35	1.51	1.29
29	N	603	CLA	CBB-CAB	3.35	1.51	1.29

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
29	N	610	CLA	CBB-CAB	3.35	1.51	1.29
29	G	611	CLA	CBB-CAB	3.35	1.51	1.29
29	N	612	CLA	CBB-CAB	3.35	1.51	1.29
29	b	615	CLA	CBB-CAB	3.35	1.51	1.29
29	n	614	CLA	CBB-CAB	3.35	1.51	1.29
29	S	613	CLA	CBB-CAB	3.35	1.51	1.29
29	g	602	CLA	CBB-CAB	3.35	1.51	1.29
36	B	625	DGA	OG1-CA1	3.35	1.43	1.33
29	c	506	CLA	CBB-CAB	3.35	1.51	1.29
29	R	610	CLA	CBB-CAB	3.35	1.51	1.29
29	B	610	CLA	CBB-CAB	3.35	1.51	1.29
29	r	611	CLA	CBB-CAB	3.35	1.51	1.29
29	Y	604	CLA	CBB-CAB	3.35	1.51	1.29
29	A	407	CLA	C1C-NC	-3.35	1.32	1.37
29	d	402	CLA	C1C-NC	-3.35	1.32	1.37
29	C	506	CLA	CBB-CAB	3.35	1.51	1.29
29	S	617	CLA	CBB-CAB	3.35	1.51	1.29
29	c	510	CLA	CBB-CAB	3.35	1.51	1.29
29	S	609	CLA	CBB-CAB	3.35	1.51	1.29
29	Y	602	CLA	CBB-CAB	3.35	1.51	1.29
29	Y	614	CLA	CBB-CAB	3.35	1.51	1.29
29	Y	603	CLA	CBB-CAB	3.35	1.51	1.29
29	g	614	CLA	CBB-CAB	3.35	1.51	1.29
29	S	604	CLA	CBB-CAB	3.35	1.51	1.29
29	S	602	CLA	CBB-CAB	3.35	1.51	1.29
29	C	510	CLA	CBB-CAB	3.35	1.51	1.29
29	g	603	CLA	CBB-CAB	3.35	1.51	1.29
29	b	607	CLA	CBB-CAB	3.34	1.51	1.29
29	C	512	CLA	CBB-CAB	3.34	1.51	1.29
29	R	609	CLA	CBB-CAB	3.34	1.51	1.29
29	n	611	CLA	CBB-CAB	3.34	1.51	1.29
29	g	611	CLA	CBB-CAB	3.34	1.51	1.29
29	Y	610	CLA	CBB-CAB	3.34	1.51	1.29
29	y	604	CLA	CBB-CAB	3.34	1.51	1.29
29	C	502	CLA	CBB-CAB	3.34	1.51	1.29
29	s	609	CLA	CBB-CAB	3.34	1.51	1.29
29	r	604	CLA	CBB-CAB	3.34	1.51	1.29
29	n	610	CLA	C1C-NC	-3.34	1.32	1.37
29	b	616	CLA	CBB-CAB	3.34	1.51	1.29
29	R	608	CLA	CBB-CAB	3.34	1.51	1.29
45	N	605	CHL	CBB-CAB	3.34	1.51	1.29
29	b	612	CLA	CBB-CAB	3.34	1.51	1.29

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
29	r	609	CLA	CBB-CAB	3.34	1.51	1.29
29	y	614	CLA	CBB-CAB	3.34	1.51	1.29
29	Y	602	CLA	C1C-NC	-3.34	1.32	1.37
45	S	607	CHL	C4B-NB	3.34	1.38	1.35
29	y	613	CLA	CBB-CAB	3.34	1.51	1.29
29	G	604	CLA	CBB-CAB	3.34	1.51	1.29
29	Y	608	CLA	CBB-CAB	3.34	1.51	1.29
29	B	616	CLA	CBB-CAB	3.34	1.51	1.29
29	n	613	CLA	CBB-CAB	3.34	1.51	1.29
29	B	607	CLA	CBB-CAB	3.34	1.51	1.29
29	y	610	CLA	CBB-CAB	3.34	1.51	1.29
29	R	604	CLA	CBB-CAB	3.34	1.51	1.29
29	g	612	CLA	CBB-CAB	3.34	1.51	1.29
29	r	603	CLA	CBB-CAB	3.34	1.51	1.29
29	b	602	CLA	CBB-CAB	3.34	1.51	1.29
29	N	613	CLA	CBB-CAB	3.34	1.51	1.29
29	C	507	CLA	CBB-CAB	3.34	1.51	1.29
29	g	610	CLA	C1C-NC	-3.34	1.32	1.37
29	D	403	CLA	CBB-CAB	3.34	1.51	1.29
29	C	504	CLA	CBB-CAB	3.34	1.51	1.29
29	s	602	CLA	CBB-CAB	3.34	1.51	1.29
45	G	607	CHL	CBB-CAB	3.34	1.51	1.29
45	r	606	CHL	CBB-CAB	3.34	1.51	1.29
29	G	614	CLA	CBB-CAB	3.34	1.51	1.29
29	s	612	CLA	CBB-CAB	3.34	1.51	1.29
29	b	611	CLA	CBB-CAB	3.33	1.51	1.29
29	b	603	CLA	CBB-CAB	3.33	1.51	1.29
29	b	605	CLA	CBB-CAB	3.33	1.51	1.29
29	n	612	CLA	C1C-NC	-3.33	1.32	1.37
29	c	504	CLA	CBB-CAB	3.33	1.51	1.29
45	N	607	CHL	CBB-CAB	3.33	1.51	1.29
29	r	608	CLA	CBB-CAB	3.33	1.51	1.29
36	b	623	DGA	OG1-CA1	3.33	1.43	1.33
29	s	613	CLA	CBB-CAB	3.33	1.51	1.29
29	c	501	CLA	CBB-CAB	3.33	1.51	1.29
29	G	610	CLA	CBB-CAB	3.33	1.51	1.29
29	y	611	CLA	CBB-CAB	3.33	1.51	1.29
29	N	611	CLA	CBB-CAB	3.33	1.51	1.29
45	g	601	CHL	CBB-CAB	3.33	1.51	1.29
29	c	511	CLA	CBB-CAB	3.33	1.51	1.29
29	c	503	CLA	CBB-CAB	3.33	1.51	1.29
29	Y	612	CLA	CBB-CAB	3.33	1.51	1.29

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
29	S	610	CLA	CBB-CAB	3.33	1.51	1.29
29	Y	613	CLA	CBB-CAB	3.33	1.51	1.29
29	C	501	CLA	CBB-CAB	3.33	1.51	1.29
29	C	511	CLA	CBB-CAB	3.33	1.51	1.29
29	b	615	CLA	C1C-NC	-3.33	1.32	1.37
29	s	603	CLA	CBB-CAB	3.33	1.51	1.29
29	y	612	CLA	CBB-CAB	3.33	1.51	1.29
45	S	607	CHL	CBB-CAB	3.33	1.51	1.29
29	S	612	CLA	CBB-CAB	3.33	1.51	1.29
29	C	509	CLA	C1C-NC	-3.33	1.32	1.37
29	S	603	CLA	CBB-CAB	3.33	1.51	1.29
29	s	611	CLA	CBB-CAB	3.32	1.51	1.29
29	y	603	CLA	C1C-NC	-3.32	1.32	1.37
29	y	604	CLA	C1C-NC	-3.32	1.32	1.37
29	c	502	CLA	CBB-CAB	3.32	1.51	1.29
45	R	606	CHL	CBB-CAB	3.32	1.51	1.29
29	A	407	CLA	CBB-CAB	3.32	1.51	1.29
29	Y	610	CLA	C1C-NC	-3.32	1.32	1.37
29	G	613	CLA	CBB-CAB	3.32	1.51	1.29
29	A	406	CLA	CBB-CAB	3.32	1.51	1.29
29	s	610	CLA	CBB-CAB	3.32	1.51	1.29
29	B	613	CLA	CBB-CAB	3.32	1.51	1.29
29	a	407	CLA	CBB-CAB	3.32	1.51	1.29
29	S	611	CLA	CBB-CAB	3.32	1.51	1.29
29	n	603	CLA	C1C-NC	-3.32	1.32	1.37
29	b	613	CLA	CBB-CAB	3.32	1.51	1.29
29	D	402	CLA	CBB-CAB	3.32	1.51	1.29
29	n	603	CLA	CBB-CAB	3.32	1.51	1.29
29	b	606	CLA	CBB-CAB	3.32	1.51	1.29
29	C	503	CLA	CBB-CAB	3.31	1.51	1.29
29	C	513	CLA	CBB-CAB	3.31	1.51	1.29
29	c	513	CLA	C1C-NC	-3.31	1.32	1.37
45	s	606	CHL	CBB-CAB	3.31	1.51	1.29
45	s	607	CHL	C4B-NB	3.31	1.38	1.35
29	y	610	CLA	C1C-NC	-3.31	1.32	1.37
29	c	507	CLA	C1C-NC	-3.31	1.32	1.37
29	b	613	CLA	C1C-NC	-3.31	1.32	1.37
29	B	609	CLA	CBB-CAB	3.31	1.51	1.29
45	s	607	CHL	CBB-CAB	3.31	1.51	1.29
29	C	510	CLA	C1C-NC	-3.31	1.32	1.37
29	a	406	CLA	CBB-CAB	3.31	1.51	1.29
45	g	607	CHL	CBB-CAB	3.31	1.51	1.29

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
29	D	402	CLA	C1C-NC	-3.31	1.32	1.37
45	N	606	CHL	C4B-NB	3.30	1.38	1.35
29	c	505	CLA	C1C-NC	-3.30	1.32	1.37
29	s	604	CLA	C1C-NC	-3.30	1.32	1.37
29	y	602	CLA	C1C-NC	-3.30	1.32	1.37
36	C	524	DGA	OG1-CA1	3.30	1.43	1.33
45	s	608	CHL	CBB-CAB	3.30	1.51	1.29
29	B	605	CLA	CBB-CAB	3.30	1.51	1.29
29	S	604	CLA	C1C-NC	-3.30	1.32	1.37
29	B	604	CLA	C1C-NC	-3.30	1.32	1.37
29	C	505	CLA	C1C-NC	-3.30	1.32	1.37
45	S	606	CHL	CBB-CAB	3.30	1.51	1.29
29	G	602	CLA	C1C-NC	-3.29	1.32	1.37
29	B	606	CLA	CBB-CAB	3.29	1.51	1.29
29	c	513	CLA	CBB-CAB	3.29	1.51	1.29
45	r	607	CHL	CBB-CAB	3.29	1.51	1.29
29	c	505	CLA	CBB-CAB	3.29	1.51	1.29
45	S	608	CHL	CBB-CAB	3.29	1.51	1.29
29	b	617	CLA	CBB-CAB	3.29	1.51	1.29
29	Y	611	CLA	CBB-CAB	3.29	1.51	1.29
29	B	615	CLA	C1C-NC	-3.29	1.32	1.37
29	r	609	CLA	C1C-NC	-3.29	1.32	1.37
45	G	606	CHL	C4B-NB	3.29	1.38	1.35
36	c	524	DGA	OG1-CA1	3.29	1.42	1.33
45	Y	607	CHL	CBB-CAB	3.29	1.51	1.29
29	g	602	CLA	C1C-NC	-3.28	1.32	1.37
29	d	402	CLA	CBB-CAB	3.28	1.51	1.29
45	R	607	CHL	CBB-CAB	3.28	1.51	1.29
29	g	612	CLA	C1C-NC	-3.28	1.32	1.37
29	b	609	CLA	CBB-CAB	3.28	1.51	1.29
29	C	503	CLA	C1C-NC	-3.28	1.32	1.37
29	C	505	CLA	CBB-CAB	3.28	1.51	1.29
45	S	601	CHL	C4B-NB	3.28	1.38	1.35
29	c	501	CLA	C1C-NC	-3.28	1.32	1.37
29	n	604	CLA	C1C-NC	-3.28	1.32	1.37
29	b	604	CLA	C1C-NC	-3.28	1.32	1.37
45	G	605	CHL	C4B-NB	3.27	1.38	1.35
38	c	519	DGD	CDB-CCB	-3.27	1.33	1.51
45	s	601	CHL	C4B-NB	3.27	1.38	1.35
45	G	606	CHL	CBB-CAB	3.27	1.51	1.29
29	B	617	CLA	CBB-CAB	3.27	1.51	1.29
29	S	603	CLA	C1C-NC	-3.27	1.32	1.37

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
45	g	606	CHL	CBB-CAB	3.27	1.51	1.29
44	H	101	RRX	C24-C25	3.27	1.56	1.45
29	C	513	CLA	C1C-NC	-3.27	1.32	1.37
29	s	603	CLA	C1C-NC	-3.27	1.32	1.37
29	b	602	CLA	C1C-NC	-3.27	1.32	1.37
29	r	610	CLA	C1C-NC	-3.26	1.32	1.37
45	R	607	CHL	C4B-NB	3.26	1.38	1.35
29	B	607	CLA	C1C-NC	-3.26	1.32	1.37
42	D	405	PL9	C3-C4	-3.26	1.44	1.49
29	N	603	CLA	C1C-NC	-3.26	1.32	1.37
29	r	613	CLA	C1C-NC	-3.26	1.32	1.37
29	C	511	CLA	C1C-NC	-3.26	1.32	1.37
29	G	603	CLA	C1C-NC	-3.26	1.32	1.37
38	c	518	DGD	CDB-CCB	-3.26	1.33	1.51
29	R	613	CLA	C1C-NC	-3.26	1.32	1.37
29	b	614	CLA	C1C-NC	-3.26	1.32	1.37
33	b	622	LMG	C22-C21	-3.26	1.33	1.51
45	n	607	CHL	CBB-CAB	3.26	1.50	1.29
38	C	523	DGD	CAA-C9A	-3.25	1.33	1.51
29	y	608	CLA	C1C-NC	-3.25	1.32	1.37
45	y	606	CHL	CBB-CAB	3.25	1.50	1.29
45	S	601	CHL	CBB-CAB	3.25	1.50	1.29
38	C	519	DGD	CDB-CCB	-3.25	1.33	1.51
29	D	403	CLA	C1C-NC	-3.25	1.33	1.37
38	c	520	DGD	CAA-C9A	-3.25	1.33	1.51
38	C	523	DGD	CGA-CFA	-3.25	1.33	1.51
33	h	102	LMG	C37-C36	-3.25	1.33	1.51
38	C	520	DGD	CAB-C9B	-3.25	1.33	1.51
29	R	609	CLA	C1C-NC	-3.25	1.33	1.37
38	c	523	DGD	CAA-C9A	-3.25	1.33	1.51
33	B	622	LMG	C22-C21	-3.25	1.33	1.51
29	N	612	CLA	C1C-NC	-3.25	1.33	1.37
29	Y	612	CLA	C1C-NC	-3.25	1.33	1.37
38	c	523	DGD	CDB-CCB	-3.25	1.33	1.51
44	h	101	RRX	C24-C25	3.25	1.56	1.45
45	y	605	CHL	C4B-NB	3.25	1.38	1.35
38	C	520	DGD	CAA-C9A	-3.25	1.33	1.51
33	H	102	LMG	C19-C18	-3.24	1.33	1.51
45	Y	606	CHL	C4B-NB	3.24	1.38	1.35
38	c	520	DGD	CAB-C9B	-3.24	1.33	1.51
29	y	611	CLA	C1C-NC	-3.24	1.33	1.37
33	c	521	LMG	C22-C21	-3.24	1.33	1.51

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
33	c	521	LMG	C40-C39	-3.24	1.33	1.51
29	s	609	CLA	C1C-NC	-3.24	1.33	1.37
45	y	607	CHL	CBB-CAB	3.24	1.50	1.29
38	c	523	DGD	CGA-CFA	-3.24	1.33	1.51
29	b	616	CLA	C1C-NC	-3.24	1.33	1.37
38	C	519	DGD	CDA-CCA	-3.24	1.33	1.51
29	b	609	CLA	C1C-NC	-3.24	1.33	1.37
29	S	602	CLA	C1C-NC	-3.24	1.33	1.37
29	Y	604	CLA	C1C-NC	-3.24	1.33	1.37
45	y	606	CHL	C4B-NB	3.24	1.38	1.35
33	h	102	LMG	C19-C18	-3.24	1.33	1.51
29	d	403	CLA	C1C-NC	-3.24	1.33	1.37
38	c	519	DGD	CDA-CCA	-3.23	1.33	1.51
45	S	608	CHL	C4B-NB	3.23	1.38	1.35
29	b	607	CLA	C1C-NC	-3.23	1.33	1.37
33	C	521	LMG	C22-C21	-3.23	1.33	1.51
29	s	611	CLA	C1C-NC	-3.23	1.33	1.37
29	G	604	CLA	C1C-NC	-3.23	1.33	1.37
33	C	521	LMG	C40-C39	-3.23	1.33	1.51
33	H	102	LMG	C40-C39	-3.23	1.33	1.51
38	C	518	DGD	CDB-CCB	-3.23	1.33	1.51
38	C	523	DGD	CDB-CCB	-3.23	1.33	1.51
45	g	605	CHL	C4B-NB	3.23	1.38	1.35
29	g	611	CLA	C1C-NC	-3.23	1.33	1.37
29	g	613	CLA	C1C-NC	-3.23	1.33	1.37
29	g	603	CLA	C1C-NC	-3.23	1.33	1.37
29	s	610	CLA	C1C-NC	-3.23	1.33	1.37
33	b	622	LMG	C19-C18	-3.22	1.33	1.51
29	a	410	CLA	CBB-CAB	3.22	1.50	1.29
29	C	512	CLA	C1C-NC	-3.22	1.33	1.37
45	Y	606	CHL	CBB-CAB	3.22	1.50	1.29
29	N	613	CLA	C1C-NC	-3.22	1.33	1.37
44	h	101	RRX	C16-C17	3.22	1.53	1.43
38	c	523	DGD	CAB-C9B	-3.22	1.33	1.51
29	r	603	CLA	C1C-NC	-3.22	1.33	1.37
38	c	519	DGD	CAB-C9B	-3.22	1.33	1.51
29	G	612	CLA	C1C-NC	-3.22	1.33	1.37
33	B	622	LMG	C19-C18	-3.21	1.33	1.51
29	c	512	CLA	C1C-NC	-3.21	1.33	1.37
33	H	102	LMG	C37-C36	-3.21	1.33	1.51
29	Y	603	CLA	C1C-NC	-3.21	1.33	1.37
38	C	518	DGD	CAB-C9B	-3.21	1.33	1.51

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
38	c	518	DGD	CAB-C9B	-3.21	1.33	1.51
45	G	608	CHL	C4B-NB	3.21	1.38	1.35
33	J	101	LMG	C19-C18	-3.21	1.33	1.51
38	C	523	DGD	CDA-CCA	-3.21	1.33	1.51
38	C	523	DGD	CAB-C9B	-3.21	1.33	1.51
29	y	613	CLA	C1C-NC	-3.21	1.33	1.37
29	s	613	CLA	C1C-NC	-3.21	1.33	1.37
29	B	616	CLA	C1C-NC	-3.21	1.33	1.37
38	C	523	DGD	CGB-CFB	-3.21	1.33	1.51
33	h	102	LMG	C40-C39	-3.21	1.33	1.51
38	C	519	DGD	CAB-C9B	-3.21	1.33	1.51
33	d	411	LMG	C19-C18	-3.21	1.33	1.51
33	c	521	LMG	C19-C18	-3.20	1.33	1.51
33	A	413	LMG	C40-C39	-3.20	1.33	1.51
33	J	101	LMG	C22-C21	-3.20	1.33	1.51
33	c	521	LMG	C37-C36	-3.20	1.33	1.51
29	S	610	CLA	C1C-NC	-3.20	1.33	1.37
29	S	614	CLA	C1C-NC	-3.20	1.33	1.37
45	S	606	CHL	C4B-NB	3.20	1.38	1.35
29	A	410	CLA	CBB-CAB	3.20	1.50	1.29
33	j	101	LMG	C19-C18	-3.20	1.33	1.51
29	N	604	CLA	C1C-NC	-3.20	1.33	1.37
45	N	606	CHL	CBB-CAB	3.20	1.50	1.29
29	g	604	CLA	C1C-NC	-3.20	1.33	1.37
33	D	411	LMG	C19-C18	-3.20	1.33	1.51
33	d	411	LMG	C22-C21	-3.20	1.33	1.51
44	H	101	RRX	C16-C17	3.20	1.53	1.43
33	D	411	LMG	C22-C21	-3.20	1.33	1.51
33	a	413	LMG	C37-C36	-3.20	1.33	1.51
33	A	413	LMG	C37-C36	-3.20	1.33	1.51
44	h	101	RRX	C29-C30	3.20	1.64	1.54
29	Y	613	CLA	C1C-NC	-3.20	1.33	1.37
33	j	101	LMG	C22-C21	-3.20	1.33	1.51
33	C	521	LMG	C37-C36	-3.20	1.33	1.51
29	S	613	CLA	C1C-NC	-3.20	1.33	1.37
33	C	521	LMG	C19-C18	-3.20	1.33	1.51
38	c	523	DGD	CDA-CCA	-3.20	1.33	1.51
38	c	523	DGD	CGB-CFB	-3.20	1.33	1.51
29	r	604	CLA	C1C-NC	-3.19	1.33	1.37
29	R	602	CLA	C1C-NC	-3.19	1.33	1.37
45	s	601	CHL	CBB-CAB	3.19	1.50	1.29
29	c	510	CLA	C1C-NC	-3.19	1.33	1.37

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
45	g	601	CHL	C4B-NB	3.19	1.38	1.35
38	c	519	DGD	CAA-C9A	-3.19	1.33	1.51
29	R	610	CLA	C1C-NC	-3.19	1.33	1.37
33	a	413	LMG	C40-C39	-3.19	1.33	1.51
29	R	604	CLA	C1C-NC	-3.18	1.33	1.37
29	S	605	CLA	C1C-NC	-3.18	1.33	1.37
29	s	605	CLA	C1C-NC	-3.18	1.33	1.37
45	n	601	CHL	C4B-NB	3.18	1.38	1.35
44	H	101	RRX	C4-C5	-3.18	1.44	1.51
29	g	614	CLA	C1C-NC	-3.18	1.33	1.37
29	c	511	CLA	C1C-NC	-3.18	1.33	1.37
29	S	611	CLA	C1C-NC	-3.18	1.33	1.37
45	G	609	CHL	C4B-NB	3.18	1.38	1.35
45	y	609	CHL	C4B-NB	3.18	1.38	1.35
29	b	603	CLA	C1C-NC	-3.18	1.33	1.37
29	B	603	CLA	C1C-NC	-3.18	1.33	1.37
29	R	612	CLA	C1C-NC	-3.18	1.33	1.37
45	Y	605	CHL	C4B-NB	3.18	1.38	1.35
29	G	613	CLA	C1C-NC	-3.18	1.33	1.37
29	s	617	CLA	C1C-NC	-3.18	1.33	1.37
29	B	609	CLA	C1C-NC	-3.18	1.33	1.37
33	A	413	LMG	C19-C18	-3.18	1.33	1.51
45	r	607	CHL	C4B-NB	3.17	1.38	1.35
29	y	614	CLA	C1C-NC	-3.17	1.33	1.37
44	h	101	RRX	C4-C5	-3.17	1.44	1.51
29	C	501	CLA	C1C-NC	-3.17	1.33	1.37
29	G	611	CLA	C1C-NC	-3.17	1.33	1.37
29	n	614	CLA	C1C-NC	-3.17	1.33	1.37
38	C	519	DGD	CAA-C9A	-3.17	1.33	1.51
29	R	603	CLA	C1C-NC	-3.17	1.33	1.37
45	g	608	CHL	C4B-NB	3.17	1.38	1.35
44	H	101	RRX	C29-C30	3.17	1.64	1.54
29	G	614	CLA	C1C-NC	-3.17	1.33	1.37
29	r	602	CLA	C1C-NC	-3.17	1.33	1.37
29	n	613	CLA	C1C-NC	-3.16	1.33	1.37
29	y	612	CLA	C1C-NC	-3.16	1.33	1.37
33	a	413	LMG	C19-C18	-3.16	1.33	1.51
45	s	606	CHL	C4B-NB	3.16	1.38	1.35
38	c	520	DGD	CDB-CCB	-3.16	1.33	1.51
29	S	612	CLA	C1C-NC	-3.16	1.33	1.37
29	s	612	CLA	C1C-NC	-3.16	1.33	1.37
45	N	605	CHL	C4B-NB	3.16	1.38	1.35

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
29	S	609	CLA	C1C-NC	-3.16	1.33	1.37
29	S	617	CLA	C1C-NC	-3.16	1.33	1.37
29	r	612	CLA	C1C-NC	-3.16	1.33	1.37
45	s	608	CHL	C4B-NB	3.16	1.38	1.35
38	C	520	DGD	CDB-CCB	-3.15	1.33	1.51
45	G	601	CHL	C4B-NB	3.15	1.38	1.35
29	B	614	CLA	C1C-NC	-3.15	1.33	1.37
45	g	609	CHL	C4B-NB	3.15	1.38	1.35
29	Y	614	CLA	C1C-NC	-3.14	1.33	1.37
29	N	611	CLA	C1C-NC	-3.14	1.33	1.37
45	N	601	CHL	C4B-NB	3.13	1.38	1.35
29	s	602	CLA	C1C-NC	-3.13	1.33	1.37
29	r	611	CLA	C1C-NC	-3.13	1.33	1.37
45	R	606	CHL	C4B-NB	3.13	1.38	1.35
29	N	614	CLA	C1C-NC	-3.12	1.33	1.37
29	Y	608	CLA	C1C-NC	-3.12	1.33	1.37
45	r	606	CHL	C4B-NB	3.12	1.38	1.35
29	n	611	CLA	C1C-NC	-3.12	1.33	1.37
29	s	614	CLA	C1C-NC	-3.11	1.33	1.37
45	n	606	CHL	CBB-CAB	3.11	1.49	1.29
45	n	605	CHL	C4B-NB	3.10	1.38	1.35
29	R	611	CLA	C1C-NC	-3.10	1.33	1.37
45	Y	609	CHL	C4B-NB	3.08	1.38	1.35
44	H	101	RRX	C7-C6	3.03	1.55	1.45
44	h	101	RRX	C7-C6	3.00	1.55	1.45
45	y	607	CHL	C4B-NB	2.99	1.37	1.35
45	n	608	CHL	C4B-NB	2.99	1.37	1.35
29	B	605	CLA	C3B-C2B	-2.97	1.36	1.40
43	f	101	HEM	CAB-C3B	2.97	1.55	1.47
43	F	101	HEM	CAB-C3B	2.97	1.55	1.47
45	N	608	CHL	C4B-NB	2.94	1.37	1.35
45	g	606	CHL	C4B-NB	2.85	1.37	1.35
45	N	609	CHL	C4B-NB	2.83	1.37	1.35
45	y	601	CHL	C4B-NB	2.82	1.37	1.35
45	N	607	CHL	C4B-NB	2.82	1.37	1.35
29	d	402	CLA	CHC-C1C	2.81	1.42	1.35
49	S	625	LPX	P1-O1	2.81	1.70	1.59
45	n	609	CHL	C4B-NB	2.81	1.37	1.35
45	g	607	CHL	C4B-NB	2.80	1.37	1.35
29	R	610	CLA	CHC-C1C	2.80	1.42	1.35
45	Y	601	CHL	C4B-NB	2.79	1.37	1.35
45	G	607	CHL	C4B-NB	2.79	1.37	1.35

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
29	D	402	CLA	CHC-C1C	2.79	1.42	1.35
29	R	609	CLA	C3B-C2B	-2.78	1.36	1.40
29	R	602	CLA	CHC-C1C	2.76	1.42	1.35
29	S	613	CLA	C3B-C2B	-2.76	1.36	1.40
29	y	602	CLA	CHC-C1C	2.76	1.42	1.35
49	s	625	LPX	P1-O1	2.75	1.70	1.59
29	S	610	CLA	CHC-C1C	2.75	1.42	1.35
45	n	606	CHL	C3B-C2B	-2.74	1.36	1.40
45	Y	607	CHL	C4B-NB	2.73	1.37	1.35
29	C	513	CLA	CHC-C1C	2.73	1.42	1.35
45	n	607	CHL	C4B-NB	2.72	1.37	1.35
29	g	602	CLA	CHC-C1C	2.71	1.41	1.35
29	C	512	CLA	CHC-C1C	2.71	1.41	1.35
29	y	613	CLA	CHC-C1C	2.71	1.41	1.35
44	H	101	RRX	C32-C1	2.71	1.59	1.53
29	r	610	CLA	CHC-C1C	2.70	1.41	1.35
45	Y	606	CHL	C3B-C2B	-2.70	1.36	1.40
29	s	610	CLA	CHC-C1C	2.70	1.41	1.35
29	G	614	CLA	CHC-C1C	2.70	1.41	1.35
29	R	609	CLA	CHC-C1C	2.70	1.41	1.35
29	D	403	CLA	CHC-C1C	2.70	1.41	1.35
29	Y	602	CLA	CHC-C1C	2.70	1.41	1.35
29	G	602	CLA	CHC-C1C	2.70	1.41	1.35
44	h	101	RRX	C32-C1	2.70	1.59	1.53
29	C	501	CLA	CHC-C1C	2.69	1.41	1.35
29	r	613	CLA	CHC-C1C	2.69	1.41	1.35
29	n	614	CLA	CHC-C1C	2.69	1.41	1.35
29	r	609	CLA	CHC-C1C	2.68	1.41	1.35
29	Y	610	CLA	CHC-C1C	2.68	1.41	1.35
29	Y	608	CLA	CHC-C1C	2.67	1.41	1.35
29	c	512	CLA	CHC-C1C	2.67	1.41	1.35
29	d	403	CLA	CHC-C1C	2.67	1.41	1.35
29	R	611	CLA	CHC-C1C	2.67	1.41	1.35
29	g	604	CLA	CHC-C1C	2.67	1.41	1.35
29	s	611	CLA	C3B-C2B	-2.67	1.36	1.40
29	G	611	CLA	CHC-C1C	2.66	1.41	1.35
29	y	612	CLA	CHC-C1C	2.66	1.41	1.35
29	c	513	CLA	CHC-C1C	2.66	1.41	1.35
29	r	608	CLA	C3B-C2B	-2.66	1.36	1.40
29	S	611	CLA	C3B-C2B	-2.66	1.36	1.40
29	R	603	CLA	CHC-C1C	2.66	1.41	1.35
29	S	617	CLA	CHC-C1C	2.66	1.41	1.35

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
30	A	408	PHO	CAC-C3C	-2.66	1.47	1.52
29	g	610	CLA	CHC-C1C	2.66	1.41	1.35
29	n	610	CLA	CHC-C1C	2.65	1.41	1.35
29	c	501	CLA	CHC-C1C	2.65	1.41	1.35
29	s	612	CLA	CHC-C1C	2.65	1.41	1.35
29	S	602	CLA	CHC-C1C	2.65	1.41	1.35
29	N	604	CLA	CHC-C1C	2.65	1.41	1.35
29	y	610	CLA	CHC-C1C	2.65	1.41	1.35
29	r	603	CLA	CHC-C1C	2.65	1.41	1.35
29	b	602	CLA	CHC-C1C	2.65	1.41	1.35
29	r	611	CLA	CHC-C1C	2.65	1.41	1.35
29	y	604	CLA	CHC-C1C	2.65	1.41	1.35
30	A	409	PHO	CAC-C3C	-2.65	1.47	1.52
29	Y	613	CLA	CHC-C1C	2.64	1.41	1.35
29	r	604	CLA	CHC-C1C	2.64	1.41	1.35
29	S	614	CLA	CHC-C1C	2.64	1.41	1.35
29	N	610	CLA	CHC-C1C	2.64	1.41	1.35
29	n	611	CLA	CHC-C1C	2.64	1.41	1.35
29	g	614	CLA	CHC-C1C	2.64	1.41	1.35
29	n	602	CLA	CHC-C1C	2.64	1.41	1.35
29	B	613	CLA	CHC-C1C	2.63	1.41	1.35
29	N	613	CLA	CHC-C1C	2.63	1.41	1.35
29	g	613	CLA	CHC-C1C	2.63	1.41	1.35
29	R	608	CLA	CHC-C1C	2.63	1.41	1.35
29	s	613	CLA	C3B-C2B	-2.63	1.36	1.40
29	s	602	CLA	CHC-C1C	2.63	1.41	1.35
29	Y	604	CLA	CHC-C1C	2.63	1.41	1.35
29	S	609	CLA	CHC-C1C	2.63	1.41	1.35
29	Y	612	CLA	CHC-C1C	2.63	1.41	1.35
29	N	611	CLA	CHC-C1C	2.62	1.41	1.35
29	g	611	CLA	CHC-C1C	2.62	1.41	1.35
29	a	407	CLA	CHC-C1C	2.62	1.41	1.35
29	S	612	CLA	CHC-C1C	2.62	1.41	1.35
29	N	614	CLA	CHC-C1C	2.62	1.41	1.35
29	r	602	CLA	CHC-C1C	2.62	1.41	1.35
29	R	613	CLA	CHC-C1C	2.62	1.41	1.35
29	g	603	CLA	CHC-C1C	2.61	1.41	1.35
29	A	405	CLA	CHC-C1C	2.61	1.41	1.35
29	A	407	CLA	CHC-C1C	2.61	1.41	1.35
29	G	612	CLA	CHC-C1C	2.61	1.41	1.35
29	y	608	CLA	CHC-C1C	2.61	1.41	1.35
30	a	408	PHO	CAC-C3C	-2.61	1.47	1.52

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
29	B	611	CLA	CHC-C1C	2.61	1.41	1.35
29	C	511	CLA	CHC-C1C	2.61	1.41	1.35
29	Y	603	CLA	CHC-C1C	2.61	1.41	1.35
29	B	607	CLA	CHC-C1C	2.61	1.41	1.35
29	c	503	CLA	CHC-C1C	2.61	1.41	1.35
29	s	617	CLA	CHC-C1C	2.61	1.41	1.35
29	b	607	CLA	CHC-C1C	2.61	1.41	1.35
29	a	405	CLA	CHC-C1C	2.61	1.41	1.35
29	y	611	CLA	CHC-C1C	2.61	1.41	1.35
42	d	405	PL9	C6-C1	-2.61	1.43	1.48
29	n	613	CLA	CHC-C1C	2.61	1.41	1.35
29	s	613	CLA	CHC-C1C	2.60	1.41	1.35
29	s	609	CLA	CHC-C1C	2.60	1.41	1.35
29	c	511	CLA	CHC-C1C	2.60	1.41	1.35
29	s	605	CLA	CHC-C1C	2.60	1.41	1.35
29	R	604	CLA	CHC-C1C	2.60	1.41	1.35
29	G	604	CLA	CHC-C1C	2.60	1.41	1.35
29	g	612	CLA	CHC-C1C	2.60	1.41	1.35
29	S	605	CLA	CHC-C1C	2.60	1.41	1.35
29	N	612	CLA	CHC-C1C	2.60	1.41	1.35
29	c	503	CLA	C3B-C2B	-2.60	1.36	1.40
33	J	101	LMG	C37-C36	-2.60	1.33	1.51
45	s	601	CHL	C3B-C2B	-2.60	1.36	1.40
29	B	603	CLA	CHC-C1C	2.59	1.41	1.35
29	B	604	CLA	CHC-C1C	2.59	1.41	1.35
45	N	606	CHL	C3B-C2B	-2.59	1.36	1.40
29	b	603	CLA	CHC-C1C	2.59	1.41	1.35
29	n	604	CLA	CHC-C1C	2.59	1.41	1.35
29	d	403	CLA	C3B-C2B	-2.59	1.36	1.40
29	n	603	CLA	C3B-C2B	-2.59	1.36	1.40
33	j	101	LMG	C37-C36	-2.59	1.33	1.51
29	r	608	CLA	CHC-C1C	2.59	1.41	1.35
29	R	612	CLA	CHC-C1C	2.59	1.41	1.35
29	N	603	CLA	C3B-C2B	-2.59	1.36	1.40
29	r	612	CLA	CHC-C1C	2.58	1.41	1.35
29	R	608	CLA	C3B-C2B	-2.58	1.36	1.40
29	s	603	CLA	CHC-C1C	2.58	1.41	1.35
29	n	612	CLA	CHC-C1C	2.58	1.41	1.35
29	y	614	CLA	CHC-C1C	2.58	1.41	1.35
29	S	603	CLA	CHC-C1C	2.58	1.41	1.35
29	G	613	CLA	CHC-C1C	2.58	1.41	1.35
29	c	507	CLA	CHC-C1C	2.58	1.41	1.35

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
29	G	610	CLA	CHC-C1C	2.58	1.41	1.35
29	C	503	CLA	CHC-C1C	2.58	1.41	1.35
29	Y	611	CLA	CHC-C1C	2.57	1.41	1.35
29	B	615	CLA	CHC-C1C	2.57	1.41	1.35
42	D	405	PL9	C6-C1	-2.57	1.44	1.48
29	S	611	CLA	CHC-C1C	2.57	1.41	1.35
29	s	614	CLA	CHC-C1C	2.57	1.41	1.35
29	c	505	CLA	CHC-C1C	2.57	1.41	1.35
29	b	604	CLA	CHC-C1C	2.57	1.41	1.35
29	s	604	CLA	C3B-C2B	-2.56	1.36	1.40
30	a	409	PHO	CAC-C3C	-2.56	1.47	1.52
29	B	602	CLA	CHC-C1C	2.56	1.41	1.35
29	B	610	CLA	CHC-C1C	2.56	1.41	1.35
45	y	606	CHL	C3B-C2B	-2.56	1.36	1.40
29	b	610	CLA	CHC-C1C	2.56	1.41	1.35
29	C	510	CLA	CHC-C1C	2.56	1.41	1.35
29	b	616	CLA	CHC-C1C	2.55	1.41	1.35
29	b	611	CLA	CHC-C1C	2.55	1.41	1.35
48	s	623	NEX	C1-C6	-2.55	1.50	1.54
33	d	411	LMG	C37-C36	-2.55	1.33	1.51
29	C	503	CLA	C3B-C2B	-2.55	1.36	1.40
29	b	613	CLA	CHC-C1C	2.55	1.41	1.35
29	b	615	CLA	CHC-C1C	2.55	1.41	1.35
29	G	614	CLA	C3B-C2B	-2.55	1.36	1.40
29	s	610	CLA	C3B-C2B	-2.55	1.36	1.40
29	b	605	CLA	C3B-C2B	-2.55	1.36	1.40
29	n	603	CLA	CHC-C1C	2.54	1.41	1.35
29	B	614	CLA	CHC-C1C	2.54	1.41	1.35
29	C	504	CLA	CHC-C1C	2.54	1.41	1.35
29	n	611	CLA	C3B-C2B	-2.54	1.36	1.40
29	D	403	CLA	C3B-C2B	-2.54	1.36	1.40
35	b	620	C7Z	C18-C5	2.54	1.55	1.50
29	S	603	CLA	C3B-C2B	-2.54	1.36	1.40
29	G	603	CLA	CHC-C1C	2.54	1.41	1.35
29	Y	614	CLA	CHC-C1C	2.54	1.41	1.35
29	C	505	CLA	CHC-C1C	2.54	1.41	1.35
29	s	609	CLA	C3B-C2B	-2.53	1.36	1.40
29	A	410	CLA	CHC-C1C	2.53	1.41	1.35
29	S	605	CLA	C3B-C2B	-2.53	1.36	1.40
33	D	411	LMG	C37-C36	-2.53	1.33	1.51
50	s	626	3PH	O21-C2	-2.53	1.40	1.46
29	B	616	CLA	CHC-C1C	2.53	1.41	1.35

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
29	g	614	CLA	C3B-C2B	-2.52	1.36	1.40
29	y	603	CLA	CHC-C1C	2.52	1.41	1.35
29	b	608	CLA	CHC-C1C	2.52	1.41	1.35
29	N	603	CLA	CHC-C1C	2.52	1.41	1.35
29	c	510	CLA	CHC-C1C	2.52	1.41	1.35
29	n	612	CLA	C3B-C2B	-2.52	1.36	1.40
29	g	612	CLA	C3B-C2B	-2.51	1.36	1.40
29	s	604	CLA	CHC-C1C	2.51	1.41	1.35
29	S	604	CLA	CHC-C1C	2.51	1.41	1.35
29	S	613	CLA	CHC-C1C	2.51	1.41	1.35
35	B	620	C7Z	C18-C5	2.51	1.55	1.50
29	a	410	CLA	CHC-C1C	2.51	1.41	1.35
29	d	402	CLA	C3B-C2B	-2.51	1.36	1.40
29	s	603	CLA	C3B-C2B	-2.51	1.36	1.40
50	S	626	3PH	O21-C2	-2.51	1.40	1.46
29	Y	603	CLA	C3B-C2B	-2.50	1.36	1.40
29	C	507	CLA	CHC-C1C	2.50	1.41	1.35
29	B	608	CLA	CHC-C1C	2.50	1.41	1.35
29	C	501	CLA	C3B-C2B	-2.50	1.36	1.40
29	b	602	CLA	C3B-C2B	-2.50	1.36	1.40
29	r	609	CLA	C3B-C2B	-2.50	1.36	1.40
29	y	603	CLA	C3B-C2B	-2.50	1.36	1.40
29	b	614	CLA	CHC-C1C	2.49	1.41	1.35
29	G	603	CLA	C3B-C2B	-2.49	1.36	1.40
29	G	612	CLA	C3B-C2B	-2.49	1.36	1.40
29	c	504	CLA	CHC-C1C	2.49	1.41	1.35
35	b	620	C7Z	C21-C26	-2.49	1.50	1.53
29	b	617	CLA	CHC-C1C	2.49	1.41	1.35
29	B	617	CLA	C3B-C2B	-2.49	1.36	1.40
35	B	620	C7Z	C21-C26	-2.49	1.50	1.53
29	s	611	CLA	CHC-C1C	2.48	1.41	1.35
29	N	602	CLA	CHC-C1C	2.48	1.41	1.35
29	B	612	CLA	CHC-C1C	2.47	1.41	1.35
29	N	612	CLA	C3B-C2B	-2.47	1.36	1.40
29	B	602	CLA	C3B-C2B	-2.47	1.36	1.40
29	B	609	CLA	CHC-C1C	2.46	1.41	1.35
29	S	610	CLA	C3B-C2B	-2.46	1.37	1.40
50	i	101	3PH	O21-C2	-2.46	1.40	1.46
29	g	603	CLA	C3B-C2B	-2.46	1.37	1.40
50	S	626	3PH	O31-C31	2.45	1.40	1.33
29	c	506	CLA	C3B-C2B	-2.45	1.37	1.40
29	N	611	CLA	C3B-C2B	-2.45	1.37	1.40

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
29	R	611	CLA	C3B-C2B	-2.45	1.37	1.40
29	Y	614	CLA	C3B-C2B	-2.44	1.37	1.40
29	c	501	CLA	C3B-C2B	-2.44	1.37	1.40
29	C	506	CLA	CHC-C1C	2.44	1.41	1.35
29	r	611	CLA	C3B-C2B	-2.44	1.37	1.40
29	D	402	CLA	C3B-C2B	-2.44	1.37	1.40
29	r	612	CLA	C3B-C2B	-2.44	1.37	1.40
44	h	101	RRX	C35-C13	2.44	1.55	1.50
47	r	622	XAT	O24-C25	-2.43	1.42	1.46
29	G	610	CLA	C3B-C2B	-2.43	1.37	1.40
29	B	617	CLA	CHC-C1C	2.43	1.41	1.35
29	b	609	CLA	CHC-C1C	2.43	1.41	1.35
29	R	612	CLA	C3B-C2B	-2.43	1.37	1.40
29	C	510	CLA	C3B-C2B	-2.43	1.37	1.40
29	c	508	CLA	CHC-C1C	2.43	1.41	1.35
44	H	101	RRX	C35-C13	2.43	1.55	1.50
29	c	509	CLA	CHC-C1C	2.42	1.41	1.35
50	i	101	3PH	O31-C31	2.42	1.40	1.33
29	b	611	CLA	C3B-C2B	-2.42	1.37	1.40
47	g	622	XAT	O24-C25	-2.41	1.42	1.46
29	r	613	CLA	C3B-C2B	-2.41	1.37	1.40
35	B	620	C7Z	C20-C13	2.41	1.55	1.50
35	b	620	C7Z	C20-C13	2.41	1.55	1.50
45	S	601	CHL	C3B-C2B	-2.41	1.37	1.40
29	C	502	CLA	CHC-C1C	2.41	1.41	1.35
29	C	508	CLA	CHC-C1C	2.40	1.41	1.35
29	s	614	CLA	C3B-C2B	-2.40	1.37	1.40
45	s	606	CHL	C3B-C2B	-2.40	1.37	1.40
29	y	614	CLA	C3B-C2B	-2.40	1.37	1.40
50	s	626	3PH	O31-C31	2.39	1.40	1.33
29	Y	602	CLA	C3B-C2B	-2.39	1.37	1.40
29	G	604	CLA	C3B-C2B	-2.39	1.37	1.40
29	S	613	CLA	C1A-CHA	2.39	1.53	1.43
29	c	506	CLA	CHC-C1C	2.38	1.41	1.35
29	y	608	CLA	C3B-C2B	-2.38	1.37	1.40
29	C	509	CLA	CHC-C1C	2.38	1.41	1.35
29	s	613	CLA	C1A-CHA	2.38	1.53	1.43
29	Y	608	CLA	C3B-C2B	-2.38	1.37	1.40
29	b	612	CLA	CHC-C1C	2.38	1.41	1.35
29	a	406	CLA	CHC-C1C	2.37	1.41	1.35
29	B	611	CLA	C3B-C2B	-2.37	1.37	1.40
29	S	605	CLA	C1A-CHA	2.36	1.52	1.43

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
29	c	502	CLA	CHC-C1C	2.36	1.41	1.35
29	B	606	CLA	CHC-C1C	2.36	1.41	1.35
29	g	604	CLA	C3B-C2B	-2.36	1.37	1.40
29	b	606	CLA	CHC-C1C	2.36	1.41	1.35
29	c	507	CLA	C3B-C2B	-2.36	1.37	1.40
29	b	609	CLA	C1A-CHA	2.36	1.52	1.43
29	Y	610	CLA	C3B-C2B	-2.35	1.37	1.40
29	c	505	CLA	C3B-C2B	-2.35	1.37	1.40
29	R	610	CLA	C1C-C2C	2.35	1.49	1.44
45	S	606	CHL	C3B-C2B	-2.35	1.37	1.40
29	C	506	CLA	C3B-C2B	-2.35	1.37	1.40
29	C	507	CLA	C3B-C2B	-2.35	1.37	1.40
29	A	407	CLA	C3B-C2B	-2.35	1.37	1.40
29	y	612	CLA	C3B-C2B	-2.34	1.37	1.40
47	G	622	XAT	O24-C25	-2.34	1.42	1.46
29	A	410	CLA	C3B-C2B	-2.33	1.37	1.40
29	b	604	CLA	C3B-C2B	-2.33	1.37	1.40
29	g	611	CLA	C3B-C2B	-2.33	1.37	1.40
29	R	610	CLA	C3B-C2B	-2.33	1.37	1.40
29	s	605	CLA	C1A-CHA	2.33	1.52	1.43
29	S	612	CLA	C3B-C2B	-2.32	1.37	1.40
29	A	406	CLA	CHC-C1C	2.32	1.40	1.35
29	d	403	CLA	C1C-C2C	2.32	1.49	1.44
29	b	605	CLA	CHC-C1C	2.32	1.40	1.35
29	B	609	CLA	C1A-CHA	2.32	1.52	1.43
29	B	616	CLA	C1A-CHA	2.32	1.52	1.43
29	a	407	CLA	C3B-C2B	-2.32	1.37	1.40
29	B	608	CLA	C1A-CHA	2.30	1.52	1.43
29	b	605	CLA	C1A-CHA	2.30	1.52	1.43
47	Y	622	XAT	O24-C25	-2.30	1.42	1.46
29	G	611	CLA	C3B-C2B	-2.30	1.37	1.40
29	S	617	CLA	C3B-C2B	-2.29	1.37	1.40
29	n	602	CLA	C3B-C2B	-2.29	1.37	1.40
48	n	623	NEX	C1-C6	-2.29	1.50	1.54
30	A	409	PHO	CMD-C2D	-2.29	1.46	1.51
29	c	505	CLA	C1A-CHA	2.29	1.52	1.43
29	S	604	CLA	C3B-C2B	-2.29	1.37	1.40
29	B	605	CLA	CHC-C1C	2.29	1.40	1.35
29	s	605	CLA	C3B-C2B	-2.29	1.37	1.40
49	S	625	LPX	P1-O2	2.28	1.68	1.59
29	C	505	CLA	C1A-CHA	2.28	1.52	1.43
29	B	605	CLA	C1A-CHA	2.28	1.52	1.43

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
45	y	609	CHL	C3B-C2B	-2.28	1.37	1.40
29	b	615	CLA	C1A-CHA	2.28	1.52	1.43
47	R	621	XAT	O24-C25	-2.28	1.43	1.46
29	c	512	CLA	C1C-C2C	2.28	1.49	1.44
49	s	625	LPX	P1-O2	2.27	1.68	1.59
29	b	616	CLA	C1A-CHA	2.27	1.52	1.43
29	C	505	CLA	C3B-C2B	-2.27	1.37	1.40
29	S	614	CLA	C3B-C2B	-2.27	1.37	1.40
29	Y	611	CLA	C3B-C2B	-2.27	1.37	1.40
29	b	608	CLA	C1A-CHA	2.26	1.52	1.43
29	b	612	CLA	C1A-CHA	2.25	1.52	1.43
29	n	613	CLA	C1C-C2C	2.25	1.48	1.44
29	y	610	CLA	C3B-C2B	-2.25	1.37	1.40
29	S	602	CLA	C3B-C2B	-2.25	1.37	1.40
29	c	501	CLA	C1A-CHA	2.24	1.52	1.43
29	R	603	CLA	C1C-C2C	2.24	1.48	1.44
29	C	512	CLA	C1C-C2C	2.24	1.48	1.44
29	a	407	CLA	C1A-CHA	2.24	1.52	1.43
29	R	613	CLA	C3B-C2B	-2.24	1.37	1.40
29	r	608	CLA	C1A-CHA	2.24	1.52	1.43
29	C	501	CLA	C1A-CHA	2.24	1.52	1.43
29	S	602	CLA	C1C-C2C	2.24	1.48	1.44
29	D	403	CLA	C1C-C2C	2.23	1.48	1.44
29	B	604	CLA	C3B-C2B	-2.23	1.37	1.40
29	r	603	CLA	C1C-C2C	2.23	1.48	1.44
29	r	610	CLA	C3B-C2B	-2.23	1.37	1.40
29	N	604	CLA	C1C-C2C	2.23	1.48	1.44
29	C	507	CLA	C1A-CHA	2.23	1.52	1.43
29	b	617	CLA	C3B-C2B	-2.23	1.37	1.40
47	N	622	XAT	O24-C25	-2.22	1.43	1.46
29	n	604	CLA	C3B-C2B	-2.22	1.37	1.40
29	R	608	CLA	C1A-CHA	2.22	1.52	1.43
29	n	611	CLA	C1A-CHA	2.22	1.52	1.43
29	r	610	CLA	C1A-CHA	2.22	1.52	1.43
29	y	611	CLA	C1A-CHA	2.22	1.52	1.43
29	N	613	CLA	C1A-CHA	2.22	1.52	1.43
29	n	614	CLA	C1C-C2C	2.22	1.48	1.44
29	N	602	CLA	C3B-C2B	-2.22	1.37	1.40
29	B	615	CLA	C1A-CHA	2.22	1.52	1.43
29	B	614	CLA	C1A-CHA	2.22	1.52	1.43
29	Y	612	CLA	C3B-C2B	-2.22	1.37	1.40
29	y	611	CLA	C3B-C2B	-2.22	1.37	1.40

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
29	y	613	CLA	C1C-C2C	2.22	1.48	1.44
29	R	613	CLA	C1A-CHA	2.22	1.52	1.43
29	c	512	CLA	C1A-CHA	2.22	1.52	1.43
45	G	609	CHL	C3B-C2B	-2.22	1.37	1.40
29	R	612	CLA	C1A-CHA	2.22	1.52	1.43
45	s	601	CHL	CHC-C1C	2.22	1.40	1.35
29	A	406	CLA	C1A-CHA	2.21	1.52	1.43
29	C	512	CLA	C1A-CHA	2.21	1.52	1.43
29	s	609	CLA	C1A-CHA	2.21	1.52	1.43
29	n	610	CLA	C3B-C2B	-2.21	1.37	1.40
29	A	407	CLA	C1A-CHA	2.21	1.52	1.43
29	N	604	CLA	C1A-CHA	2.21	1.52	1.43
29	c	510	CLA	C3B-C2B	-2.21	1.37	1.40
29	y	613	CLA	C1A-CHA	2.21	1.52	1.43
29	N	613	CLA	C3B-C2B	-2.21	1.37	1.40
29	s	603	CLA	C1B-NB	2.21	1.37	1.35
47	n	622	XAT	O24-C25	-2.21	1.43	1.46
29	g	611	CLA	C1A-CHA	2.21	1.52	1.43
29	G	613	CLA	C3B-C2B	-2.21	1.37	1.40
29	G	614	CLA	C1C-C2C	2.20	1.48	1.44
29	Y	612	CLA	C1A-CHA	2.20	1.52	1.43
29	G	611	CLA	C1A-CHA	2.20	1.52	1.43
29	n	613	CLA	C3B-C2B	-2.20	1.37	1.40
29	S	609	CLA	C3B-C2B	-2.20	1.37	1.40
29	Y	613	CLA	C1A-CHA	2.20	1.52	1.43
29	S	611	CLA	C1A-CHA	2.20	1.52	1.43
42	d	405	PL9	C53-C6	-2.20	1.46	1.50
29	y	612	CLA	C1A-CHA	2.20	1.52	1.43
29	R	602	CLA	C1C-C2C	2.20	1.48	1.44
29	N	614	CLA	C3B-C2B	-2.20	1.37	1.40
29	Y	611	CLA	C1A-CHA	2.20	1.52	1.43
29	g	610	CLA	C1A-CHA	2.20	1.52	1.43
29	N	614	CLA	C1C-C2C	2.20	1.48	1.44
29	n	604	CLA	C1A-CHA	2.20	1.52	1.43
29	b	603	CLA	C1A-CHA	2.20	1.52	1.43
29	G	613	CLA	C1A-CHA	2.20	1.52	1.43
47	y	622	XAT	O24-C25	-2.19	1.43	1.46
29	C	504	CLA	C3B-C2B	-2.19	1.37	1.40
29	d	402	CLA	C1C-C2C	2.19	1.48	1.44
29	S	603	CLA	C1A-CHA	2.19	1.52	1.43
29	b	614	CLA	C1A-CHA	2.19	1.52	1.43
29	S	603	CLA	C1C-C2C	2.19	1.48	1.44

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
29	Y	603	CLA	C1A-CHA	2.19	1.52	1.43
29	r	612	CLA	C1A-CHA	2.19	1.52	1.43
29	C	502	CLA	C1A-CHA	2.19	1.52	1.43
29	n	614	CLA	C1A-CHA	2.19	1.52	1.43
29	C	513	CLA	C1C-C2C	2.19	1.48	1.44
29	c	502	CLA	C1A-CHA	2.19	1.52	1.43
29	S	617	CLA	C1C-C2C	2.19	1.48	1.44
29	s	610	CLA	C1C-C2C	2.19	1.48	1.44
29	b	612	CLA	C3B-C2B	-2.18	1.37	1.40
29	s	605	CLA	C1C-C2C	2.18	1.48	1.44
29	b	603	CLA	C3B-C2B	-2.18	1.37	1.40
29	y	603	CLA	C1A-CHA	2.18	1.52	1.43
29	C	501	CLA	C1C-C2C	2.18	1.48	1.44
29	N	611	CLA	C1A-CHA	2.18	1.52	1.43
29	B	612	CLA	C1A-CHA	2.18	1.52	1.43
29	s	611	CLA	C1A-CHA	2.18	1.52	1.43
29	N	612	CLA	C1A-CHA	2.18	1.52	1.43
29	N	614	CLA	C1A-CHA	2.18	1.52	1.43
29	g	613	CLA	C1A-CHA	2.18	1.52	1.43
29	R	611	CLA	C1A-CHA	2.18	1.52	1.43
29	b	602	CLA	C1C-C2C	2.18	1.48	1.44
29	S	612	CLA	C1A-CHA	2.18	1.52	1.43
29	a	406	CLA	C1A-CHA	2.18	1.52	1.43
29	s	617	CLA	C1A-CHA	2.18	1.52	1.43
30	a	409	PHO	CMC-C2C	-2.18	1.46	1.51
29	R	611	CLA	C1C-C2C	2.18	1.48	1.44
29	y	602	CLA	C1C-C2C	2.18	1.48	1.44
29	S	609	CLA	C1A-CHA	2.18	1.52	1.43
29	b	606	CLA	C3B-C2B	-2.18	1.37	1.40
29	b	607	CLA	C3B-C2B	-2.18	1.37	1.40
29	C	502	CLA	C3B-C2B	-2.18	1.37	1.40
29	n	613	CLA	C1A-CHA	2.17	1.52	1.43
29	G	604	CLA	C1A-CHA	2.17	1.52	1.43
45	n	606	CHL	CHC-C1C	2.17	1.40	1.35
29	g	614	CLA	C1A-CHA	2.17	1.52	1.43
29	b	604	CLA	C1A-CHA	2.17	1.52	1.43
29	r	603	CLA	C1A-CHA	2.17	1.52	1.43
29	r	611	CLA	C1A-CHA	2.17	1.52	1.43
29	R	611	CLA	C1B-NB	2.17	1.37	1.35
29	r	613	CLA	C1A-CHA	2.17	1.52	1.43
29	s	612	CLA	C1A-CHA	2.17	1.52	1.43
29	R	602	CLA	C3B-C2B	-2.17	1.37	1.40

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
45	S	607	CHL	C3B-C2B	-2.17	1.37	1.40
29	N	613	CLA	C1C-C2C	2.17	1.48	1.44
29	Y	602	CLA	C1C-C2C	2.17	1.48	1.44
29	s	604	CLA	C1A-CHA	2.17	1.52	1.43
29	g	602	CLA	C3B-C2B	-2.17	1.37	1.40
29	g	610	CLA	C3B-C2B	-2.17	1.37	1.40
29	B	611	CLA	C1C-C2C	2.17	1.48	1.44
42	D	405	PL9	C53-C6	-2.17	1.46	1.50
29	g	613	CLA	C3B-C2B	-2.17	1.37	1.40
29	n	610	CLA	C1A-CHA	2.17	1.52	1.43
29	b	610	CLA	C1A-CHA	2.17	1.52	1.43
29	y	612	CLA	C1C-C2C	2.17	1.48	1.44
29	C	509	CLA	C1A-CHA	2.17	1.52	1.43
29	y	614	CLA	C1A-CHA	2.17	1.52	1.43
29	b	609	CLA	C3B-C2B	-2.17	1.37	1.40
29	c	507	CLA	C1A-CHA	2.16	1.52	1.43
29	B	610	CLA	C3B-C2B	-2.16	1.37	1.40
45	s	607	CHL	C3B-C2B	-2.16	1.37	1.40
29	y	608	CLA	C1A-CHA	2.16	1.52	1.43
45	R	606	CHL	C3B-C2B	-2.16	1.37	1.40
45	R	607	CHL	C3B-C2B	-2.16	1.37	1.40
50	s	626	3PH	O21-C21	2.16	1.40	1.34
29	r	602	CLA	C3B-C2B	-2.16	1.37	1.40
29	g	612	CLA	C1A-CHA	2.16	1.52	1.43
29	G	612	CLA	C1A-CHA	2.16	1.52	1.43
29	D	402	CLA	C1C-C2C	2.16	1.48	1.44
29	N	611	CLA	C1C-C2C	2.16	1.48	1.44
29	G	614	CLA	C1A-CHA	2.16	1.52	1.43
29	R	603	CLA	C1A-CHA	2.16	1.52	1.43
50	i	101	3PH	O21-C21	2.16	1.40	1.34
29	b	616	CLA	C3B-C2B	-2.16	1.37	1.40
29	C	513	CLA	C1A-CHA	2.16	1.52	1.43
29	B	607	CLA	C3B-C2B	-2.16	1.37	1.40
29	a	410	CLA	C1A-CHA	2.16	1.52	1.43
29	Y	604	CLA	C3B-C2B	-2.15	1.37	1.40
29	S	617	CLA	C1A-CHA	2.15	1.52	1.43
36	c	524	DGA	OG2-CG2	-2.15	1.41	1.46
29	s	603	CLA	C1A-CHA	2.15	1.52	1.43
45	S	608	CHL	C3B-C2B	-2.15	1.37	1.40
29	s	602	CLA	C3B-C2B	-2.15	1.37	1.40
36	C	524	DGA	OG2-CG2	-2.15	1.41	1.46
29	y	602	CLA	C3B-C2B	-2.15	1.37	1.40

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
29	C	503	CLA	C1A-CHA	2.15	1.52	1.43
29	n	602	CLA	C1C-C2C	2.15	1.48	1.44
29	B	606	CLA	C3B-C2B	-2.15	1.37	1.40
29	A	410	CLA	C1A-CHA	2.15	1.52	1.43
29	G	614	CLA	C1B-NB	2.15	1.37	1.35
29	B	616	CLA	C3B-C2B	-2.15	1.37	1.40
29	S	602	CLA	C1A-CHA	2.15	1.52	1.43
29	g	604	CLA	C1A-CHA	2.15	1.52	1.43
29	C	508	CLA	C3B-C2B	-2.15	1.37	1.40
29	Y	608	CLA	C1A-CHA	2.15	1.52	1.43
29	Y	610	CLA	C1A-CHA	2.15	1.52	1.43
29	C	511	CLA	C1A-CHA	2.15	1.52	1.43
29	R	610	CLA	C1A-CHA	2.15	1.52	1.43
29	B	610	CLA	C1A-CHA	2.14	1.52	1.43
29	n	612	CLA	C1A-CHA	2.14	1.52	1.43
29	s	617	CLA	C3B-C2B	-2.14	1.37	1.40
29	r	611	CLA	C1C-C2C	2.14	1.48	1.44
29	B	602	CLA	C1A-CHA	2.14	1.52	1.43
29	Y	614	CLA	C1A-CHA	2.14	1.52	1.43
29	S	610	CLA	C1C-C2C	2.14	1.48	1.44
29	Y	608	CLA	C1C-C2C	2.14	1.48	1.44
29	B	607	CLA	C1A-CHA	2.14	1.52	1.43
29	b	610	CLA	C3B-C2B	-2.14	1.37	1.40
29	c	501	CLA	C1C-C2C	2.14	1.48	1.44
29	c	513	CLA	C1A-CHA	2.14	1.52	1.43
29	s	603	CLA	C1C-C2C	2.14	1.48	1.44
29	G	612	CLA	C1C-C2C	2.14	1.48	1.44
29	S	610	CLA	C1A-CHA	2.14	1.52	1.43
29	c	511	CLA	C1A-CHA	2.14	1.52	1.43
29	N	611	CLA	C1B-NB	2.14	1.37	1.35
29	c	513	CLA	C1C-C2C	2.14	1.48	1.44
29	y	610	CLA	C1A-CHA	2.14	1.52	1.43
50	S	626	3PH	O21-C21	2.13	1.40	1.34
29	N	603	CLA	C1A-CHA	2.13	1.52	1.43
29	b	611	CLA	C1C-C2C	2.13	1.48	1.44
29	d	403	CLA	C1A-CHA	2.13	1.52	1.43
29	r	602	CLA	C1C-C2C	2.13	1.48	1.44
29	s	613	CLA	C1C-C2C	2.13	1.48	1.44
29	G	602	CLA	C1C-C2C	2.13	1.48	1.44
29	g	603	CLA	C1C-C2C	2.13	1.48	1.44
29	g	612	CLA	C1C-C2C	2.13	1.48	1.44
29	b	602	CLA	C1A-CHA	2.13	1.52	1.43

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
29	G	603	CLA	C1C-C2C	2.13	1.48	1.44
29	c	508	CLA	C3B-C2B	-2.13	1.37	1.40
29	N	612	CLA	C1C-C2C	2.13	1.48	1.44
29	D	402	CLA	C3D-C4D	-2.13	1.39	1.44
29	s	614	CLA	C1A-CHA	2.13	1.51	1.43
29	B	603	CLA	C1A-CHA	2.13	1.51	1.43
29	B	611	CLA	C1A-CHA	2.13	1.51	1.43
29	b	603	CLA	C1C-C2C	2.13	1.48	1.44
50	s	626	3PH	O31-C3	-2.13	1.40	1.45
29	a	410	CLA	C3B-C2B	-2.13	1.37	1.40
29	s	602	CLA	C1C-C2C	2.13	1.48	1.44
29	s	617	CLA	C1C-C2C	2.13	1.48	1.44
29	y	603	CLA	C1C-C2C	2.13	1.48	1.44
29	D	403	CLA	C1A-CHA	2.13	1.51	1.43
29	b	607	CLA	C1A-CHA	2.13	1.51	1.43
29	R	609	CLA	C1A-CHA	2.13	1.51	1.43
29	s	610	CLA	C1A-CHA	2.13	1.51	1.43
29	c	503	CLA	C1A-CHA	2.12	1.51	1.43
29	B	604	CLA	C1A-CHA	2.12	1.51	1.43
29	N	610	CLA	C1A-CHA	2.12	1.51	1.43
29	r	604	CLA	C1A-CHA	2.12	1.51	1.43
29	R	602	CLA	C1A-CHA	2.12	1.51	1.43
30	A	409	PHO	CMC-C2C	-2.12	1.46	1.51
30	a	408	PHO	CMC-C2C	-2.12	1.46	1.51
30	a	409	PHO	CMD-C2D	-2.12	1.46	1.51
29	r	612	CLA	C1C-C2C	2.12	1.48	1.44
29	r	602	CLA	C1A-CHA	2.12	1.51	1.43
29	g	604	CLA	C1C-C2C	2.12	1.48	1.44
29	r	603	CLA	C3B-C2B	-2.12	1.37	1.40
29	S	603	CLA	C1B-NB	2.12	1.37	1.35
29	Y	604	CLA	C1A-CHA	2.12	1.51	1.43
29	g	602	CLA	C1A-CHA	2.12	1.51	1.43
29	C	504	CLA	C1A-CHA	2.12	1.51	1.43
29	A	410	CLA	C1B-NB	2.12	1.37	1.35
29	S	605	CLA	C1C-C2C	2.12	1.48	1.44
29	B	604	CLA	C3D-C4D	-2.11	1.39	1.44
29	y	604	CLA	C1A-CHA	2.11	1.51	1.43
35	b	620	C7Z	C40-C33	2.11	1.55	1.50
29	G	611	CLA	C1C-C2C	2.11	1.48	1.44
48	G	623	NEX	C1-C6	-2.11	1.51	1.54
29	c	504	CLA	C1A-CHA	2.11	1.51	1.43
29	r	604	CLA	C1C-C2C	2.11	1.48	1.44

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
29	R	604	CLA	C1A-CHA	2.11	1.51	1.43
29	G	613	CLA	C1C-C2C	2.11	1.48	1.44
29	N	610	CLA	C1C-C2C	2.11	1.48	1.44
29	R	608	CLA	C1C-C2C	2.11	1.48	1.44
29	G	610	CLA	C1A-CHA	2.10	1.51	1.43
29	n	603	CLA	C1C-C2C	2.10	1.48	1.44
35	B	620	C7Z	C40-C33	2.10	1.55	1.50
29	b	611	CLA	C1A-CHA	2.10	1.51	1.43
50	S	626	3PH	O31-C3	-2.10	1.40	1.45
29	g	614	CLA	C1B-NB	2.10	1.37	1.35
29	S	604	CLA	C1A-CHA	2.10	1.51	1.43
29	b	617	CLA	C3D-C4D	-2.10	1.39	1.44
29	Y	613	CLA	C1C-C2C	2.10	1.48	1.44
29	r	603	CLA	C1B-NB	2.10	1.37	1.35
29	A	410	CLA	C1C-C2C	2.10	1.48	1.44
30	A	408	PHO	CMD-C2D	-2.10	1.46	1.51
29	g	613	CLA	C1C-C2C	2.10	1.48	1.44
29	B	606	CLA	C3D-C4D	-2.09	1.39	1.44
29	s	609	CLA	C1B-NB	2.09	1.37	1.35
45	N	605	CHL	C3B-C2B	-2.09	1.37	1.40
45	g	609	CHL	C3B-C2B	-2.09	1.37	1.40
45	r	606	CHL	C3B-C2B	-2.09	1.37	1.40
43	f	101	HEM	FE-ND	2.09	2.07	1.96
29	r	609	CLA	C1A-CHA	2.09	1.51	1.43
29	G	602	CLA	C3B-C2B	-2.09	1.37	1.40
29	n	603	CLA	C1A-CHA	2.09	1.51	1.43
29	g	603	CLA	C1A-CHA	2.09	1.51	1.43
29	s	602	CLA	C1A-CHA	2.09	1.51	1.43
29	S	609	CLA	C1C-C2C	2.09	1.48	1.44
29	n	611	CLA	C1C-C2C	2.09	1.48	1.44
29	c	508	CLA	C1A-CHA	2.09	1.51	1.43
29	N	602	CLA	C1A-CHA	2.09	1.51	1.43
29	Y	603	CLA	C1C-C2C	2.09	1.48	1.44
45	N	601	CHL	C3B-C2B	-2.09	1.37	1.40
29	N	603	CLA	C1C-C2C	2.09	1.48	1.44
29	G	603	CLA	C1A-CHA	2.08	1.51	1.43
29	N	604	CLA	C3B-C2B	-2.08	1.37	1.40
29	R	612	CLA	C1C-C2C	2.08	1.48	1.44
29	g	614	CLA	C1C-C2C	2.08	1.48	1.44
29	S	612	CLA	C1C-C2C	2.08	1.48	1.44
29	c	504	CLA	C3B-C2B	-2.08	1.37	1.40
29	s	609	CLA	C1C-C2C	2.08	1.48	1.44

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
30	a	408	PHO	CMD-C2D	-2.08	1.46	1.51
29	Y	612	CLA	C1C-C2C	2.08	1.48	1.44
29	r	610	CLA	C1C-C2C	2.08	1.48	1.44
29	n	602	CLA	C1A-CHA	2.08	1.51	1.43
29	S	614	CLA	C1A-CHA	2.08	1.51	1.43
29	c	502	CLA	C3B-C2B	-2.08	1.37	1.40
29	C	506	CLA	C1A-CHA	2.08	1.51	1.43
29	Y	604	CLA	C1C-C2C	2.08	1.48	1.44
29	c	503	CLA	C1B-NB	2.08	1.37	1.35
29	B	602	CLA	C1C-C2C	2.08	1.48	1.44
29	y	610	CLA	C1C-C2C	2.07	1.48	1.44
45	s	608	CHL	C3B-C2B	-2.07	1.37	1.40
29	r	612	CLA	C1B-NB	2.07	1.37	1.35
29	G	604	CLA	C1C-C2C	2.07	1.48	1.44
29	B	606	CLA	C1A-CHA	2.07	1.51	1.43
30	A	408	PHO	CMC-C2C	-2.07	1.46	1.51
29	B	603	CLA	C1C-C2C	2.07	1.48	1.44
29	c	506	CLA	C1A-CHA	2.07	1.51	1.43
29	b	604	CLA	C3D-C4D	-2.07	1.39	1.44
29	s	612	CLA	C3B-C2B	-2.07	1.37	1.40
29	S	611	CLA	C1C-C2C	2.07	1.48	1.44
29	g	611	CLA	C1C-C2C	2.07	1.48	1.44
29	C	503	CLA	C1B-NB	2.07	1.37	1.35
29	C	508	CLA	C1A-CHA	2.07	1.51	1.43
29	g	602	CLA	C1C-C2C	2.07	1.48	1.44
29	b	606	CLA	C1A-CHA	2.07	1.51	1.43
29	R	602	CLA	C1B-NB	2.07	1.37	1.35
45	Y	609	CHL	C3B-C2B	-2.07	1.37	1.40
29	c	509	CLA	C1A-CHA	2.06	1.51	1.43
45	S	601	CHL	CHC-C1C	2.06	1.40	1.35
29	R	610	CLA	C1B-NB	2.06	1.37	1.35
29	b	613	CLA	C1A-CHA	2.06	1.51	1.43
29	n	613	CLA	C1B-NB	2.06	1.37	1.35
29	y	608	CLA	C1C-C2C	2.06	1.48	1.44
29	S	614	CLA	C1C-C2C	2.06	1.48	1.44
29	n	612	CLA	C1C-C2C	2.06	1.48	1.44
29	A	405	CLA	C1A-CHA	2.06	1.51	1.43
42	D	405	PL9	C52-C5	-2.06	1.46	1.50
29	B	617	CLA	C1A-CHA	2.06	1.51	1.43
45	G	601	CHL	C3B-C2B	-2.06	1.37	1.40
29	C	510	CLA	C1A-CHA	2.06	1.51	1.43
29	r	613	CLA	C1C-C2C	2.06	1.48	1.44

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
29	S	603	CLA	MG-NC	2.06	2.11	2.06
29	b	606	CLA	C3D-C4D	-2.05	1.39	1.44
29	a	405	CLA	C1A-CHA	2.05	1.51	1.43
29	G	613	CLA	C1B-NB	2.05	1.37	1.35
29	n	610	CLA	C1C-C2C	2.05	1.48	1.44
29	c	510	CLA	C3D-C4D	-2.05	1.39	1.44
45	G	608	CHL	C3B-C2B	-2.05	1.37	1.40
29	c	511	CLA	C1C-C2C	2.05	1.48	1.44
29	C	503	CLA	C1C-C2C	2.05	1.48	1.44
29	B	612	CLA	C3D-C4D	-2.05	1.39	1.44
45	r	607	CHL	C3B-C2B	-2.05	1.37	1.40
29	n	604	CLA	C1C-C2C	2.05	1.48	1.44
29	r	608	CLA	C1C-C2C	2.05	1.48	1.44
29	b	616	CLA	C1C-C2C	2.05	1.48	1.44
29	r	604	CLA	C3B-C2B	-2.05	1.37	1.40
29	y	614	CLA	C1C-C2C	2.05	1.48	1.44
29	N	610	CLA	C3B-C2B	-2.05	1.37	1.40
29	B	607	CLA	C3D-C4D	-2.05	1.39	1.44
29	s	605	CLA	MG-NC	2.05	2.11	2.06
29	R	613	CLA	C1C-C2C	2.05	1.48	1.44
29	b	602	CLA	C3D-C4D	-2.05	1.39	1.44
29	b	613	CLA	C3D-C4D	-2.05	1.39	1.44
29	b	605	CLA	MG-NC	2.05	2.11	2.06
50	i	101	3PH	O31-C3	-2.05	1.40	1.45
29	b	613	CLA	C3B-C2B	-2.05	1.37	1.40
45	g	601	CHL	C3B-C2B	-2.04	1.37	1.40
29	S	605	CLA	MG-NC	2.04	2.11	2.06
29	r	609	CLA	C3D-C4D	-2.04	1.39	1.44
45	n	605	CHL	C3B-C2B	-2.04	1.37	1.40
29	B	615	CLA	C1C-C2C	2.04	1.48	1.44
29	R	604	CLA	C1C-C2C	2.04	1.48	1.44
29	Y	604	CLA	C3D-C4D	-2.04	1.39	1.44
29	B	604	CLA	C1C-C2C	2.04	1.48	1.44
29	Y	614	CLA	C1C-C2C	2.04	1.48	1.44
45	G	606	CHL	C3B-C2B	-2.04	1.37	1.40
48	y	623	NEX	C1-C6	-2.04	1.51	1.54
29	N	613	CLA	C1B-NB	2.04	1.37	1.35
29	G	610	CLA	C1B-NB	2.04	1.37	1.35
29	n	613	CLA	MG-NC	2.04	2.11	2.06
29	c	503	CLA	C1C-C2C	2.04	1.48	1.44
29	B	609	CLA	C3B-C2B	-2.04	1.37	1.40
33	H	102	LMG	C22-C21	-2.04	1.33	1.49

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
29	B	602	CLA	C3D-C4D	-2.03	1.39	1.44
29	r	602	CLA	C1B-NB	2.03	1.37	1.35
29	B	609	CLA	C1C-C2C	2.03	1.48	1.44
45	g	605	CHL	C3B-C2B	-2.03	1.37	1.40
29	y	613	CLA	C1B-NB	2.03	1.37	1.35
29	b	607	CLA	C3D-C4D	-2.03	1.39	1.44
29	s	612	CLA	C1C-C2C	2.03	1.48	1.44
29	B	615	CLA	C3D-C4D	-2.03	1.39	1.44
33	h	102	LMG	C22-C21	-2.03	1.33	1.49
29	R	611	CLA	MG-NC	2.03	2.11	2.06
29	B	613	CLA	C1A-CHA	2.03	1.51	1.43
29	R	604	CLA	C3B-C2B	-2.03	1.37	1.40
29	n	611	CLA	MG-NC	2.03	2.11	2.06
29	y	611	CLA	C1B-NB	2.03	1.37	1.35
29	B	603	CLA	CHD-C1D	2.03	1.42	1.38
29	s	610	CLA	C3D-C4D	-2.03	1.39	1.44
29	C	511	CLA	C1C-C2C	2.03	1.48	1.44
29	C	513	CLA	C1B-NB	2.03	1.37	1.35
29	C	505	CLA	C3D-C4D	-2.03	1.39	1.44
29	r	610	CLA	C3D-C4D	-2.03	1.39	1.44
29	r	610	CLA	C1B-NB	2.02	1.37	1.35
29	c	510	CLA	C1A-CHA	2.02	1.51	1.43
38	c	520	DGD	CDA-CCA	-2.02	1.33	1.49
29	C	511	CLA	C3B-C2B	-2.02	1.37	1.40
29	C	506	CLA	C3D-C4D	-2.02	1.39	1.44
45	G	606	CHL	CHC-C1C	2.02	1.40	1.35
29	S	613	CLA	C1C-C2C	2.02	1.48	1.44
29	G	602	CLA	C1A-CHA	2.02	1.51	1.43
33	A	413	LMG	C22-C21	-2.02	1.33	1.49
29	B	605	CLA	MG-NC	2.02	2.11	2.06
29	C	505	CLA	C1C-C2C	2.02	1.48	1.44
45	N	606	CHL	CHC-C1C	2.02	1.40	1.35
38	c	518	DGD	CGB-CFB	-2.02	1.33	1.49
29	d	402	CLA	C3D-C4D	-2.02	1.39	1.44
29	S	617	CLA	MG-NC	2.02	2.11	2.06
29	G	602	CLA	C3D-C4D	-2.02	1.39	1.44
29	r	611	CLA	C1B-NB	2.02	1.37	1.35
29	c	507	CLA	C1C-C2C	2.02	1.48	1.44
29	Y	602	CLA	C1A-CHA	2.02	1.51	1.43
29	N	611	CLA	MG-NC	2.02	2.11	2.06
29	A	406	CLA	CHD-C1D	2.02	1.42	1.38
29	C	501	CLA	C3D-C4D	-2.01	1.39	1.44

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
29	c	507	CLA	C3D-C4D	-2.01	1.39	1.44
33	a	413	LMG	C22-C21	-2.01	1.33	1.49
29	R	608	CLA	C1B-NB	2.01	1.37	1.35
29	b	615	CLA	C1C-C2C	2.01	1.48	1.44
38	C	520	DGD	CDA-CCA	-2.01	1.33	1.49
29	B	610	CLA	C1B-NB	2.01	1.37	1.35
38	C	518	DGD	CGB-CFB	-2.01	1.33	1.49
29	n	614	CLA	C3B-C2B	-2.01	1.37	1.40
29	r	613	CLA	C1B-NB	2.01	1.37	1.35
38	C	520	DGD	CGB-CFB	-2.01	1.33	1.49
29	Y	614	CLA	C3D-C4D	-2.01	1.39	1.44
29	n	614	CLA	C1B-NB	2.01	1.37	1.35
30	A	408	PHO	CMB-C2B	-2.01	1.46	1.51
33	H	102	LMG	C43-C42	-2.01	1.33	1.49
33	c	521	LMG	C25-C24	-2.01	1.33	1.49
38	c	520	DGD	CGB-CFB	-2.01	1.33	1.49
29	Y	611	CLA	C1C-C2C	2.01	1.48	1.44
33	c	521	LMG	C43-C42	-2.01	1.33	1.49
33	A	413	LMG	C43-C42	-2.01	1.33	1.49
33	C	521	LMG	C43-C42	-2.01	1.33	1.49
29	y	602	CLA	C1A-CHA	2.01	1.51	1.43
29	Y	610	CLA	C1C-C2C	2.01	1.48	1.44
29	B	608	CLA	C3B-C2B	-2.01	1.37	1.40
45	g	608	CHL	C3B-C2B	-2.01	1.37	1.40
29	s	617	CLA	C1B-NB	2.01	1.37	1.35
48	R	622	NEX	O24-C25	-2.01	1.43	1.46
33	D	411	LMG	C25-C24	-2.01	1.33	1.49
29	n	614	CLA	MG-NC	2.01	2.11	2.06
30	a	408	PHO	CMB-C2B	-2.00	1.46	1.51
38	C	519	DGD	CGB-CFB	-2.00	1.33	1.49
29	B	617	CLA	C3D-C4D	-2.00	1.39	1.44
29	B	613	CLA	C3D-C4D	-2.00	1.39	1.44
29	N	612	CLA	MG-NC	2.00	2.11	2.06
29	B	612	CLA	C3B-C2B	-2.00	1.37	1.40
29	c	509	CLA	C3D-C4D	-2.00	1.39	1.44
45	Y	606	CHL	CHC-C1C	2.00	1.40	1.35
29	N	613	CLA	MG-NC	2.00	2.11	2.06
33	a	413	LMG	C43-C42	-2.00	1.33	1.49
33	C	521	LMG	C25-C24	-2.00	1.33	1.49
29	Y	602	CLA	C3D-C4D	-2.00	1.39	1.44
29	c	506	CLA	C3D-C4D	-2.00	1.39	1.44
29	a	407	CLA	C1C-C2C	2.00	1.48	1.44

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
29	r	611	CLA	MG-NC	2.00	2.11	2.06
29	s	603	CLA	MG-NC	2.00	2.11	2.06
33	b	622	LMG	C25-C24	-2.00	1.33	1.49
45	Y	605	CHL	C3B-C2B	-2.00	1.37	1.40

All (4140) bond angle outliers are listed below:

Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
29	B	617	CLA	C4-C3-C5	-22.25	77.84	115.27
29	b	617	CLA	C4-C3-C5	-22.23	77.87	115.27
29	b	617	CLA	C5-C3-C2	18.79	159.15	121.12
29	B	617	CLA	C5-C3-C2	18.73	159.01	121.12
31	C	517	BCR	C10-C11-C12	17.71	178.49	123.22
31	c	516	BCR	C10-C11-C12	17.71	178.47	123.22
31	C	516	BCR	C10-C11-C12	17.55	177.99	123.22
31	c	515	BCR	C10-C11-C12	17.52	177.89	123.22
31	d	404	BCR	C10-C11-C12	17.51	177.87	123.22
31	B	619	BCR	C10-C11-C12	17.49	177.80	123.22
31	b	618	BCR	C10-C11-C12	17.48	177.75	123.22
31	B	618	BCR	C10-C11-C12	17.47	177.73	123.22
31	b	619	BCR	C10-C11-C12	17.41	177.54	123.22
31	C	514	BCR	C10-C11-C12	17.25	177.05	123.22
31	c	514	BCR	C10-C11-C12	17.20	176.90	123.22
31	A	411	BCR	C10-C11-C12	17.15	176.75	123.22
31	a	411	BCR	C10-C11-C12	17.07	176.49	123.22
47	y	622	XAT	C37-C21-C36	-16.87	82.48	107.37
31	D	404	BCR	C10-C11-C12	16.82	175.70	123.22
31	c	517	BCR	C10-C11-C12	16.59	174.99	123.22
47	Y	622	XAT	C37-C21-C36	-16.58	82.91	107.37
29	B	617	CLA	C4-C3-C2	-16.55	81.23	123.68
29	b	617	CLA	C4-C3-C2	-16.50	81.35	123.68
31	C	515	BCR	C10-C11-C12	16.43	174.48	123.22
31	C	514	BCR	C16-C15-C14	14.95	154.10	123.47
31	C	515	BCR	C16-C15-C14	14.26	152.69	123.47
31	c	514	BCR	C16-C15-C14	14.21	152.59	123.47
31	c	514	BCR	C11-C10-C9	13.76	146.95	127.31
31	C	516	BCR	C11-C10-C9	13.63	146.76	127.31
31	C	514	BCR	C11-C10-C9	13.60	146.72	127.31
31	b	619	BCR	C16-C15-C14	13.44	151.00	123.47
31	c	516	BCR	C11-C10-C9	13.21	146.16	127.31
31	c	515	BCR	C16-C15-C14	13.06	150.23	123.47
31	b	618	BCR	C11-C10-C9	13.02	145.90	127.31

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
31	B	618	BCR	C11-C10-C9	12.97	145.82	127.31
31	B	619	BCR	C16-C15-C14	12.93	149.96	123.47
31	b	618	BCR	C16-C15-C14	12.90	149.90	123.47
31	B	618	BCR	C16-C15-C14	12.82	149.73	123.47
31	b	619	BCR	C21-C20-C19	12.80	163.15	123.22
31	C	516	BCR	C16-C15-C14	12.77	149.64	123.47
31	a	411	BCR	C11-C10-C9	12.73	145.48	127.31
31	B	619	BCR	C21-C20-C19	12.65	162.68	123.22
31	A	411	BCR	C11-C10-C9	12.64	145.35	127.31
47	Y	622	XAT	C37-C21-C22	-12.62	87.06	108.98
47	y	622	XAT	C37-C21-C22	-12.60	87.09	108.98
31	c	517	BCR	C21-C20-C19	12.54	162.35	123.22
31	a	411	BCR	C21-C20-C19	12.41	161.95	123.22
31	c	517	BCR	C16-C15-C14	12.33	148.73	123.47
31	A	411	BCR	C21-C20-C19	12.31	161.62	123.22
31	B	619	BCR	C11-C10-C9	12.18	144.70	127.31
31	C	517	BCR	C21-C20-C19	12.09	160.96	123.22
31	b	619	BCR	C11-C10-C9	12.03	144.47	127.31
31	c	516	BCR	C16-C15-C14	12.02	148.10	123.47
31	C	517	BCR	C16-C15-C14	11.97	147.99	123.47
31	d	404	BCR	C11-C10-C9	11.72	144.04	127.31
31	a	411	BCR	C16-C15-C14	11.66	147.36	123.47
31	C	515	BCR	C21-C20-C19	11.61	159.45	123.22
31	c	514	BCR	C21-C20-C19	11.59	159.38	123.22
31	A	411	BCR	C16-C15-C14	11.53	147.08	123.47
31	D	404	BCR	C16-C15-C14	11.49	147.02	123.47
31	c	515	BCR	C11-C10-C9	11.40	143.59	127.31
31	d	404	BCR	C16-C15-C14	11.38	146.78	123.47
31	D	404	BCR	C11-C10-C9	11.38	143.55	127.31
31	C	514	BCR	C21-C20-C19	11.37	158.69	123.22
31	C	517	BCR	C11-C10-C9	11.36	143.52	127.31
31	b	618	BCR	C21-C20-C19	11.11	157.89	123.22
31	c	516	BCR	C21-C20-C19	11.02	157.60	123.22
31	D	404	BCR	C11-C12-C13	10.97	157.24	126.42
31	B	618	BCR	C21-C20-C19	10.97	157.45	123.22
31	c	515	BCR	C21-C20-C19	10.78	156.86	123.22
31	a	411	BCR	C11-C12-C13	10.75	156.61	126.42
31	d	404	BCR	C11-C12-C13	10.74	156.60	126.42
31	C	515	BCR	C11-C10-C9	10.73	142.62	127.31
29	b	609	CLA	C4A-NA-C1A	10.65	111.49	106.71
31	c	515	BCR	C11-C12-C13	10.58	156.15	126.42
31	C	517	BCR	C11-C12-C13	10.51	155.95	126.42

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
31	A	411	BCR	C11-C12-C13	10.49	155.89	126.42
31	c	517	BCR	C11-C10-C9	10.39	142.14	127.31
31	C	516	BCR	C11-C12-C13	10.32	155.41	126.42
31	b	619	BCR	C11-C12-C13	10.25	155.20	126.42
31	C	515	BCR	C11-C12-C13	10.21	155.10	126.42
31	c	516	BCR	C11-C12-C13	10.20	155.06	126.42
31	C	516	BCR	C21-C20-C19	10.15	154.91	123.22
31	b	618	BCR	C11-C12-C13	10.12	154.85	126.42
31	B	619	BCR	C11-C12-C13	10.10	154.78	126.42
31	B	618	BCR	C11-C12-C13	9.92	154.29	126.42
29	S	613	CLA	C4A-NA-C1A	9.87	111.14	106.71
29	B	609	CLA	C4A-NA-C1A	9.86	111.14	106.71
29	B	608	CLA	C4A-NA-C1A	9.85	111.13	106.71
29	b	608	CLA	C4A-NA-C1A	9.82	111.12	106.71
29	b	605	CLA	C4A-NA-C1A	9.76	111.10	106.71
29	b	615	CLA	C4A-NA-C1A	9.75	111.09	106.71
29	b	610	CLA	C4A-NA-C1A	9.69	111.06	106.71
29	B	605	CLA	C4A-NA-C1A	9.69	111.06	106.71
29	c	505	CLA	C4A-NA-C1A	9.68	111.06	106.71
29	s	613	CLA	C4A-NA-C1A	9.68	111.06	106.71
31	c	517	BCR	C11-C12-C13	9.67	153.58	126.42
31	d	404	BCR	C21-C20-C19	9.65	153.34	123.22
29	B	617	CLA	C4A-NA-C1A	9.62	111.03	106.71
31	c	514	BCR	C11-C12-C13	9.62	153.45	126.42
29	b	613	CLA	C4A-NA-C1A	9.62	111.03	106.71
31	C	514	BCR	C11-C12-C13	9.61	153.42	126.42
29	C	505	CLA	C4A-NA-C1A	9.58	111.01	106.71
29	B	613	CLA	C4A-NA-C1A	9.54	111.00	106.71
29	c	502	CLA	C4A-NA-C1A	9.54	110.99	106.71
29	a	406	CLA	C4A-NA-C1A	9.53	110.99	106.71
29	b	606	CLA	C4A-NA-C1A	9.53	110.99	106.71
29	A	406	CLA	C4A-NA-C1A	9.52	110.99	106.71
29	B	610	CLA	C4A-NA-C1A	9.51	110.98	106.71
29	B	614	CLA	C4A-NA-C1A	9.49	110.97	106.71
29	b	612	CLA	C4A-NA-C1A	9.46	110.96	106.71
29	C	509	CLA	C4A-NA-C1A	9.45	110.95	106.71
29	C	502	CLA	C4A-NA-C1A	9.45	110.95	106.71
29	B	606	CLA	C4A-NA-C1A	9.44	110.95	106.71
29	s	605	CLA	C4A-NA-C1A	9.43	110.95	106.71
31	D	404	BCR	C21-C20-C19	9.42	152.62	123.22
29	R	604	CLA	C4A-NA-C1A	9.40	110.93	106.71
29	c	513	CLA	C4A-NA-C1A	9.38	110.92	106.71

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
29	G	610	CLA	C4A-NA-C1A	9.37	110.92	106.71
29	c	504	CLA	C4A-NA-C1A	9.37	110.92	106.71
29	c	509	CLA	C4A-NA-C1A	9.37	110.92	106.71
29	n	604	CLA	C4A-NA-C1A	9.34	110.90	106.71
29	y	613	CLA	C4A-NA-C1A	9.34	110.90	106.71
29	b	614	CLA	C4A-NA-C1A	9.33	110.90	106.71
29	B	615	CLA	C4A-NA-C1A	9.32	110.90	106.71
29	S	605	CLA	C4A-NA-C1A	9.32	110.89	106.71
29	y	603	CLA	C4A-NA-C1A	9.29	110.88	106.71
31	C	516	BCR	C20-C19-C18	9.28	152.47	126.42
29	n	613	CLA	C4A-NA-C1A	9.28	110.88	106.71
29	c	511	CLA	C4A-NA-C1A	9.25	110.86	106.71
29	N	604	CLA	C4A-NA-C1A	9.24	110.86	106.71
29	Y	613	CLA	C4A-NA-C1A	9.24	110.86	106.71
29	c	501	CLA	C4A-NA-C1A	9.22	110.85	106.71
29	N	613	CLA	C4A-NA-C1A	9.22	110.85	106.71
29	n	610	CLA	C4A-NA-C1A	9.22	110.85	106.71
29	a	410	CLA	C4A-NA-C1A	9.22	110.85	106.71
29	S	604	CLA	C4A-NA-C1A	9.20	110.84	106.71
29	r	604	CLA	C4A-NA-C1A	9.20	110.84	106.71
29	C	511	CLA	C4A-NA-C1A	9.19	110.84	106.71
29	r	610	CLA	C4A-NA-C1A	9.16	110.83	106.71
29	N	603	CLA	C4A-NA-C1A	9.16	110.82	106.71
29	A	410	CLA	C4A-NA-C1A	9.15	110.82	106.71
29	n	603	CLA	C4A-NA-C1A	9.14	110.81	106.71
29	N	612	CLA	C4A-NA-C1A	9.13	110.81	106.71
29	s	602	CLA	C4A-NA-C1A	9.12	110.81	106.71
29	a	405	CLA	C4A-NA-C1A	9.12	110.81	106.71
29	n	614	CLA	C4A-NA-C1A	9.12	110.80	106.71
29	B	616	CLA	C4A-NA-C1A	9.11	110.80	106.71
29	G	613	CLA	C4A-NA-C1A	9.10	110.80	106.71
29	C	510	CLA	C4A-NA-C1A	9.10	110.80	106.71
29	S	617	CLA	C4A-NA-C1A	9.10	110.80	106.71
29	Y	603	CLA	C4A-NA-C1A	9.08	110.79	106.71
29	C	501	CLA	C4A-NA-C1A	9.08	110.79	106.71
29	R	603	CLA	C4A-NA-C1A	9.07	110.78	106.71
29	C	504	CLA	C4A-NA-C1A	9.06	110.78	106.71
29	g	610	CLA	C4A-NA-C1A	9.06	110.78	106.71
29	A	407	CLA	C4A-NA-C1A	9.06	110.78	106.71
29	Y	611	CLA	C4A-NA-C1A	9.06	110.78	106.71
29	g	611	CLA	C4A-NA-C1A	9.05	110.78	106.71
29	G	614	CLA	C4A-NA-C1A	9.05	110.78	106.71

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
29	g	613	CLA	C4A-NA-C1A	9.04	110.77	106.71
29	C	507	CLA	C4A-NA-C1A	9.02	110.76	106.71
29	Y	610	CLA	C4A-NA-C1A	9.01	110.75	106.71
29	Y	612	CLA	C4A-NA-C1A	9.01	110.75	106.71
29	N	614	CLA	C4A-NA-C1A	9.00	110.75	106.71
29	A	405	CLA	C4A-NA-C1A	9.00	110.75	106.71
29	G	604	CLA	C4A-NA-C1A	8.99	110.75	106.71
29	b	603	CLA	C4A-NA-C1A	8.99	110.75	106.71
29	b	616	CLA	C4A-NA-C1A	8.99	110.75	106.71
29	s	617	CLA	C4A-NA-C1A	8.99	110.75	106.71
29	r	608	CLA	C4A-NA-C1A	8.98	110.74	106.71
29	B	612	CLA	C4A-NA-C1A	8.98	110.74	106.71
29	G	603	CLA	C4A-NA-C1A	8.97	110.74	106.71
29	c	512	CLA	C4A-NA-C1A	8.96	110.73	106.71
29	g	603	CLA	C4A-NA-C1A	8.95	110.73	106.71
29	c	507	CLA	C4A-NA-C1A	8.94	110.73	106.71
29	S	612	CLA	C4A-NA-C1A	8.93	110.72	106.71
29	c	508	CLA	C4A-NA-C1A	8.93	110.72	106.71
29	R	613	CLA	C4A-NA-C1A	8.92	110.72	106.71
29	n	611	CLA	C4A-NA-C1A	8.91	110.71	106.71
29	y	611	CLA	C4A-NA-C1A	8.91	110.71	106.71
29	G	611	CLA	C4A-NA-C1A	8.90	110.71	106.71
29	B	604	CLA	C4A-NA-C1A	8.89	110.70	106.71
29	R	608	CLA	C4A-NA-C1A	8.89	110.70	106.71
29	r	613	CLA	C4A-NA-C1A	8.89	110.70	106.71
29	y	610	CLA	C4A-NA-C1A	8.89	110.70	106.71
29	N	610	CLA	C4A-NA-C1A	8.88	110.70	106.71
29	c	510	CLA	C4A-NA-C1A	8.86	110.69	106.71
29	S	602	CLA	C4A-NA-C1A	8.86	110.69	106.71
29	r	603	CLA	C4A-NA-C1A	8.85	110.69	106.71
29	g	614	CLA	C4A-NA-C1A	8.84	110.68	106.71
29	C	508	CLA	C4A-NA-C1A	8.84	110.68	106.71
29	a	407	CLA	C4A-NA-C1A	8.82	110.67	106.71
29	g	612	CLA	C4A-NA-C1A	8.82	110.67	106.71
29	Y	614	CLA	C4A-NA-C1A	8.81	110.67	106.71
29	R	602	CLA	C4A-NA-C1A	8.80	110.66	106.71
29	B	602	CLA	C4A-NA-C1A	8.80	110.66	106.71
29	r	611	CLA	C4A-NA-C1A	8.79	110.66	106.71
29	s	609	CLA	C4A-NA-C1A	8.78	110.65	106.71
29	r	602	CLA	C4A-NA-C1A	8.78	110.65	106.71
29	y	608	CLA	C4A-NA-C1A	8.78	110.65	106.71
29	S	609	CLA	C4A-NA-C1A	8.77	110.65	106.71

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
29	s	604	CLA	C4A-NA-C1A	8.77	110.65	106.71
29	C	513	CLA	C4A-NA-C1A	8.77	110.65	106.71
29	g	602	CLA	C4A-NA-C1A	8.75	110.64	106.71
29	S	611	CLA	C4A-NA-C1A	8.74	110.64	106.71
29	s	612	CLA	C4A-NA-C1A	8.74	110.64	106.71
29	S	603	CLA	C4A-NA-C1A	8.73	110.63	106.71
29	y	614	CLA	C4A-NA-C1A	8.73	110.63	106.71
29	y	604	CLA	C4A-NA-C1A	8.73	110.63	106.71
29	c	506	CLA	C4A-NA-C1A	8.71	110.62	106.71
29	b	602	CLA	C4A-NA-C1A	8.71	110.62	106.71
29	Y	604	CLA	C4A-NA-C1A	8.71	110.62	106.71
29	C	503	CLA	C4A-NA-C1A	8.69	110.61	106.71
29	y	612	CLA	C4A-NA-C1A	8.69	110.61	106.71
29	R	612	CLA	C4A-NA-C1A	8.68	110.61	106.71
29	G	612	CLA	C4A-NA-C1A	8.68	110.61	106.71
29	B	603	CLA	C4A-NA-C1A	8.67	110.61	106.71
29	R	611	CLA	C4A-NA-C1A	8.64	110.59	106.71
29	C	512	CLA	C4A-NA-C1A	8.63	110.59	106.71
29	s	611	CLA	C4A-NA-C1A	8.63	110.59	106.71
29	Y	608	CLA	C4A-NA-C1A	8.61	110.58	106.71
29	b	607	CLA	C4A-NA-C1A	8.61	110.58	106.71
31	b	618	BCR	C20-C19-C18	8.60	150.58	126.42
29	R	609	CLA	C4A-NA-C1A	8.60	110.57	106.71
29	s	614	CLA	C4A-NA-C1A	8.60	110.57	106.71
29	b	617	CLA	C4A-NA-C1A	8.59	110.57	106.71
29	C	506	CLA	C4A-NA-C1A	8.59	110.57	106.71
29	g	604	CLA	C4A-NA-C1A	8.58	110.56	106.71
31	B	618	BCR	C20-C19-C18	8.58	150.52	126.42
29	b	604	CLA	C4A-NA-C1A	8.55	110.55	106.71
31	c	515	BCR	C20-C19-C18	8.55	150.44	126.42
29	B	607	CLA	C4A-NA-C1A	8.53	110.54	106.71
29	N	602	CLA	C4A-NA-C1A	8.53	110.54	106.71
29	S	614	CLA	C4A-NA-C1A	8.52	110.54	106.71
29	r	612	CLA	C4A-NA-C1A	8.51	110.53	106.71
29	N	611	CLA	C4A-NA-C1A	8.51	110.53	106.71
31	c	516	BCR	C20-C19-C18	8.50	150.28	126.42
29	S	610	CLA	C4A-NA-C1A	8.46	110.51	106.71
29	n	612	CLA	C4A-NA-C1A	8.45	110.50	106.71
29	c	503	CLA	C4A-NA-C1A	8.44	110.50	106.71
29	d	403	CLA	C4A-NA-C1A	8.44	110.50	106.71
29	G	602	CLA	C4A-NA-C1A	8.44	110.50	106.71
29	r	609	CLA	C4A-NA-C1A	8.42	110.49	106.71

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
29	s	603	CLA	C4A-NA-C1A	8.41	110.49	106.71
31	C	514	BCR	C20-C19-C18	8.35	149.89	126.42
29	D	402	CLA	C4A-NA-C1A	8.35	110.46	106.71
29	D	403	CLA	C4A-NA-C1A	8.28	110.43	106.71
29	s	610	CLA	C4A-NA-C1A	8.27	110.43	106.71
31	C	515	BCR	C20-C19-C18	8.26	149.62	126.42
29	d	402	CLA	C4A-NA-C1A	8.16	110.37	106.71
29	R	610	CLA	C4A-NA-C1A	8.13	110.36	106.71
29	b	611	CLA	C4A-NA-C1A	8.12	110.36	106.71
29	n	602	CLA	C4A-NA-C1A	8.10	110.35	106.71
29	y	602	CLA	C4A-NA-C1A	8.05	110.33	106.71
29	Y	602	CLA	C4A-NA-C1A	7.99	110.30	106.71
29	B	611	CLA	C4A-NA-C1A	7.98	110.29	106.71
31	c	514	BCR	C20-C19-C18	7.97	148.80	126.42
31	C	517	BCR	C20-C19-C18	7.65	147.92	126.42
31	B	619	BCR	C20-C19-C18	7.34	147.04	126.42
29	s	613	CLA	O2D-CGD-CBD	7.34	124.31	111.27
47	y	622	XAT	C36-C21-C22	7.34	121.73	108.98
29	S	613	CLA	O2D-CGD-CBD	7.28	124.20	111.27
31	b	619	BCR	C20-C19-C18	7.26	146.82	126.42
31	A	411	BCR	C20-C19-C18	7.18	146.58	126.42
29	c	502	CLA	O2A-C1-C2	7.17	127.49	108.64
47	Y	622	XAT	C36-C21-C22	7.13	121.38	108.98
31	c	517	BCR	C20-C19-C18	7.09	146.33	126.42
31	a	411	BCR	C20-C19-C18	7.09	146.32	126.42
47	y	622	XAT	C37-C21-C26	-7.05	91.02	110.05
47	Y	622	XAT	C37-C21-C26	-7.04	91.04	110.05
47	R	621	XAT	C15-C14-C13	-6.91	117.45	127.31
36	B	625	DGA	CDB-CCB-CBB	-6.76	80.12	114.42
29	s	605	CLA	O2D-CGD-CBD	6.75	123.27	111.27
36	b	623	DGA	CDB-CCB-CBB	-6.73	80.25	114.42
29	S	605	CLA	O2D-CGD-CBD	6.72	123.20	111.27
36	c	524	DGA	CDB-CCB-CBB	-6.67	80.55	114.42
36	C	524	DGA	CDB-CCB-CBB	-6.66	80.64	114.42
47	r	622	XAT	C15-C14-C13	-6.60	117.89	127.31
40	c	627	LMK	O2-C4-O3	-6.60	109.11	124.09
40	C	527	LMK	O2-C4-O3	-6.54	109.23	124.09
29	S	605	CLA	O2A-C1-C2	6.36	125.34	108.64
29	B	609	CLA	O2D-CGD-CBD	6.24	122.36	111.27
29	C	506	CLA	O2D-CGD-CBD	6.24	122.35	111.27
29	c	506	CLA	O2D-CGD-CBD	6.21	122.31	111.27
29	c	501	CLA	O2D-CGD-CBD	6.20	122.28	111.27

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
29	b	609	CLA	O2D-CGD-CBD	6.13	122.16	111.27
29	R	602	CLA	O2D-CGD-CBD	6.06	122.03	111.27
29	C	501	CLA	O2D-CGD-CBD	6.05	122.03	111.27
29	C	504	CLA	O2A-C1-C2	5.99	124.37	108.64
29	B	605	CLA	O2A-C1-C2	5.97	124.32	108.64
47	n	622	XAT	C31-C30-C29	-5.96	118.81	127.31
29	R	604	CLA	O2A-C1-C2	5.93	122.84	108.97
29	c	504	CLA	O2A-C1-C2	5.93	124.22	108.64
29	G	604	CLA	O2A-C1-C2	5.91	122.80	108.97
29	B	602	CLA	CMD-C2D-C1D	5.90	135.12	124.71
29	b	606	CLA	CMD-C2D-C1D	5.90	135.11	124.71
31	d	404	BCR	C20-C19-C18	5.90	142.98	126.42
29	A	405	CLA	O2A-C1-C2	5.89	124.12	108.64
29	s	614	CLA	O2A-C1-C2	5.89	124.11	108.64
29	c	503	CLA	CMD-C2D-C1D	5.88	135.07	124.71
29	C	513	CLA	O2A-C1-C2	5.87	124.06	108.64
29	b	602	CLA	CMD-C2D-C1D	5.87	135.05	124.71
29	C	511	CLA	CMD-C2D-C1D	5.87	135.05	124.71
29	s	610	CLA	O2A-C1-C2	5.86	124.05	108.64
29	B	606	CLA	CMD-C2D-C1D	5.85	135.03	124.71
29	c	510	CLA	CMD-C2D-C1D	5.85	135.02	124.71
29	c	513	CLA	O2A-C1-C2	5.83	123.95	108.64
29	n	613	CLA	O2A-C1-C2	5.82	123.94	108.64
47	r	622	XAT	C7-C8-C9	-5.81	116.52	125.53
29	c	511	CLA	CMD-C2D-C1D	5.80	134.94	124.71
29	C	501	CLA	CMD-C2D-C1D	5.80	134.93	124.71
29	b	605	CLA	O2A-C1-C2	5.78	123.83	108.64
29	C	510	CLA	CMD-C2D-C1D	5.76	134.87	124.71
29	S	614	CLA	CMD-C2D-C1D	5.76	134.86	124.71
29	G	614	CLA	CMD-C2D-C1D	5.76	134.86	124.71
29	r	613	CLA	O2D-CGD-CBD	5.76	121.50	111.27
29	C	503	CLA	CMD-C2D-C1D	5.75	134.84	124.71
46	y	620	LUT	C21-C26-C25	5.74	121.71	111.42
29	y	603	CLA	O2D-CGD-CBD	5.73	121.45	111.27
31	D	404	BCR	C20-C19-C18	5.73	142.50	126.42
46	s	620	LUT	C21-C26-C27	5.73	119.94	112.70
29	r	604	CLA	O2A-C1-C2	5.73	122.36	108.97
29	R	609	CLA	CMD-C2D-C1D	5.72	134.79	124.71
44	H	101	RRX	C15-C14-C13	-5.72	119.15	127.31
29	g	604	CLA	O2D-CGD-CBD	5.72	121.43	111.27
29	n	611	CLA	O2A-C1-C2	5.72	122.34	108.97
29	r	608	CLA	CMD-C2D-C1D	5.70	134.76	124.71

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
29	N	611	CLA	O2D-CGD-CBD	5.69	121.38	111.27
29	B	603	CLA	CMD-C2D-C1D	5.69	134.73	124.71
29	n	604	CLA	CMD-C2D-C1D	5.68	134.72	124.71
29	b	617	CLA	O2A-C1-C2	5.68	123.56	108.64
29	a	405	CLA	O2A-C1-C2	5.68	123.55	108.64
46	Y	620	LUT	C21-C26-C25	5.67	121.58	111.42
35	B	620	C7Z	C11-C10-C9	-5.67	119.22	127.31
47	y	622	XAT	C31-C30-C29	-5.67	119.22	127.31
29	R	612	CLA	CMD-C2D-C1D	5.67	134.71	124.71
29	r	609	CLA	CMD-C2D-C1D	5.67	134.70	124.71
47	R	621	XAT	C7-C8-C9	-5.66	116.74	125.53
29	s	604	CLA	CMD-C2D-C1D	5.66	134.69	124.71
29	S	604	CLA	CMD-C2D-C1D	5.66	134.69	124.71
29	g	613	CLA	CMD-C2D-C1D	5.64	134.66	124.71
29	s	614	CLA	CMD-C2D-C1D	5.64	134.66	124.71
29	B	613	CLA	O2D-CGD-CBD	5.64	121.29	111.27
29	n	611	CLA	O2D-CGD-CBD	5.64	121.29	111.27
29	c	503	CLA	O2D-CGD-CBD	5.64	121.28	111.27
46	G	621	LUT	C21-C26-C25	5.64	121.51	111.42
29	g	614	CLA	CMD-C2D-C1D	5.63	134.64	124.71
29	c	501	CLA	CMD-C2D-C1D	5.62	134.63	124.71
29	C	502	CLA	O2A-C1-C2	5.62	123.41	108.64
29	r	613	CLA	CMD-C2D-C1D	5.61	134.61	124.71
29	R	608	CLA	CMD-C2D-C1D	5.61	134.59	124.71
29	n	614	CLA	O2A-C1-C2	5.60	122.08	108.97
29	A	407	CLA	O2A-C1-C2	5.60	122.07	108.97
29	Y	603	CLA	O2D-CGD-CBD	5.60	121.22	111.27
29	b	608	CLA	O2A-C1-C2	5.60	123.35	108.64
29	s	610	CLA	CMD-C2D-C1D	5.60	134.58	124.71
29	g	604	CLA	CMD-C2D-C1D	5.59	134.57	124.71
29	N	604	CLA	O2D-CGD-CBD	5.59	121.21	111.27
29	r	610	CLA	CMD-C2D-C1D	5.59	134.57	124.71
29	R	603	CLA	CMD-C2D-C1D	5.58	134.56	124.71
29	B	607	CLA	O2D-CGD-CBD	5.58	121.19	111.27
29	g	603	CLA	CMD-C2D-C1D	5.58	134.55	124.71
29	G	604	CLA	CMD-C2D-C1D	5.58	134.54	124.71
29	a	406	CLA	CMD-C2D-C1D	5.57	134.54	124.71
29	G	613	CLA	CMD-C2D-C1D	5.57	134.53	124.71
47	N	622	XAT	C31-C30-C29	-5.57	119.36	127.31
29	C	509	CLA	CMD-C2D-C1D	5.57	134.53	124.71
29	R	602	CLA	CMD-C2D-C1D	5.57	134.53	124.71
29	b	613	CLA	O2D-CGD-CBD	5.57	121.16	111.27

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
29	Y	608	CLA	CMD-C2D-C1D	5.56	134.52	124.71
29	R	613	CLA	O2D-CGD-CBD	5.56	121.15	111.27
29	c	502	CLA	CMD-C2D-C1D	5.56	134.50	124.71
46	y	621	LUT	C21-C26-C25	5.55	121.37	111.42
29	r	603	CLA	CMD-C2D-C1D	5.55	134.50	124.71
29	B	608	CLA	O2A-C1-C2	5.55	123.23	108.64
29	y	604	CLA	CMD-C2D-C1D	5.55	134.49	124.71
29	c	509	CLA	O2D-CGD-CBD	5.55	121.13	111.27
29	R	604	CLA	CMD-C2D-C1D	5.54	134.48	124.71
29	B	605	CLA	CMD-C2D-C1D	5.54	134.48	124.71
29	B	607	CLA	CMD-C2D-C1D	5.54	134.48	124.71
29	C	513	CLA	CMD-C2D-C1D	5.54	134.48	124.71
29	G	604	CLA	O2D-CGD-CBD	5.53	121.10	111.27
46	g	621	LUT	C21-C26-C25	5.53	121.33	111.42
29	b	603	CLA	CMD-C2D-C1D	5.53	134.46	124.71
46	s	621	LUT	C21-C26-C25	5.53	121.33	111.42
29	Y	604	CLA	CMD-C2D-C1D	5.53	134.46	124.71
29	g	610	CLA	CMD-C2D-C1D	5.53	134.45	124.71
29	N	604	CLA	CMD-C2D-C1D	5.52	134.45	124.71
29	b	607	CLA	O2D-CGD-CBD	5.52	121.08	111.27
29	R	613	CLA	CMD-C2D-C1D	5.52	134.45	124.71
29	c	509	CLA	CMD-C2D-C1D	5.52	134.44	124.71
29	Y	614	CLA	CMD-C2D-C1D	5.52	134.44	124.71
29	S	610	CLA	CMD-C2D-C1D	5.52	134.44	124.71
29	r	612	CLA	CMD-C2D-C1D	5.52	134.44	124.71
29	a	407	CLA	O2A-C1-C2	5.52	121.88	108.97
29	n	604	CLA	O2D-CGD-CBD	5.52	121.07	111.27
44	h	101	RRX	C15-C14-C13	-5.52	119.44	127.31
29	r	602	CLA	CMD-C2D-C1D	5.51	134.43	124.71
29	A	406	CLA	CMD-C2D-C1D	5.51	134.43	124.71
29	y	608	CLA	CMD-C2D-C1D	5.51	134.42	124.71
29	r	611	CLA	CMD-C2D-C1D	5.51	134.42	124.71
29	c	507	CLA	CMD-C2D-C1D	5.51	134.42	124.71
29	b	615	CLA	O2A-C1-C2	5.50	123.10	108.64
29	g	602	CLA	O2A-C1-C2	5.50	123.09	108.64
29	A	405	CLA	CMD-C2D-C1D	5.50	134.40	124.71
29	g	602	CLA	O2D-CGD-CBD	5.50	121.03	111.27
29	d	403	CLA	O2D-CGD-CBD	5.49	121.03	111.27
29	C	506	CLA	CMD-C2D-C1D	5.49	134.39	124.71
29	r	609	CLA	O2A-C1-C2	5.49	123.07	108.64
29	N	604	CLA	O2A-C1-C2	5.49	123.06	108.64
29	B	615	CLA	O2A-C1-C2	5.48	123.04	108.64

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
29	D	402	CLA	O2A-C1-C2	5.47	123.02	108.64
29	n	603	CLA	O2A-C1-C2	5.47	123.02	108.64
29	G	610	CLA	CMD-C2D-C1D	5.47	134.35	124.71
29	G	602	CLA	O2A-C1-C2	5.47	123.01	108.64
29	G	602	CLA	CMD-C2D-C1D	5.47	134.35	124.71
46	n	621	LUT	C21-C26-C25	5.47	121.21	111.42
47	Y	622	XAT	C15-C14-C13	-5.47	119.51	127.31
46	Y	621	LUT	C21-C26-C25	5.46	121.20	111.42
29	B	617	CLA	O2A-C1-C2	5.46	122.98	108.64
29	b	607	CLA	CMD-C2D-C1D	5.46	134.33	124.71
29	C	503	CLA	O2D-CGD-CBD	5.46	120.97	111.27
35	b	620	C7Z	C11-C10-C9	-5.46	119.52	127.31
29	y	614	CLA	CMD-C2D-C1D	5.46	134.33	124.71
29	n	612	CLA	CMD-C2D-C1D	5.46	134.33	124.71
29	S	609	CLA	CMD-C2D-C1D	5.46	134.33	124.71
29	Y	613	CLA	CMD-C2D-C1D	5.45	134.33	124.71
29	n	604	CLA	O2A-C1-C2	5.45	122.97	108.64
29	R	610	CLA	CMD-C2D-C1D	5.45	134.32	124.71
29	S	613	CLA	CMD-C2D-C1D	5.45	134.31	124.71
29	G	603	CLA	CMD-C2D-C1D	5.45	134.31	124.71
29	c	506	CLA	CMD-C2D-C1D	5.45	134.31	124.71
29	s	613	CLA	CMD-C2D-C1D	5.45	134.31	124.71
29	C	507	CLA	CMD-C2D-C1D	5.44	134.30	124.71
29	R	604	CLA	O2D-CGD-CBD	5.44	120.94	111.27
29	y	613	CLA	CMD-C2D-C1D	5.44	134.30	124.71
29	C	509	CLA	O2D-CGD-CBD	5.44	120.93	111.27
29	G	611	CLA	O2D-CGD-CBD	5.43	120.92	111.27
29	b	603	CLA	O2A-C1-C2	5.43	122.91	108.64
29	G	610	CLA	O2A-C1-C2	5.43	122.90	108.64
29	A	410	CLA	O2D-CGD-CBD	5.43	120.91	111.27
29	N	603	CLA	O2D-CGD-CBD	5.43	120.91	111.27
29	n	603	CLA	O2D-CGD-CBD	5.43	120.91	111.27
29	R	609	CLA	O2A-C1-C2	5.42	122.89	108.64
29	B	614	CLA	CMD-C2D-C1D	5.42	134.27	124.71
29	B	603	CLA	O2A-C1-C2	5.42	122.89	108.64
29	Y	612	CLA	O2D-CGD-CBD	5.42	120.90	111.27
29	d	402	CLA	O2A-C1-C2	5.42	122.88	108.64
29	s	617	CLA	CMD-C2D-C1D	5.42	134.26	124.71
29	b	605	CLA	CMD-C2D-C1D	5.41	134.26	124.71
29	a	410	CLA	O2D-CGD-CBD	5.41	120.89	111.27
29	C	504	CLA	CMD-C2D-C1D	5.41	134.25	124.71
29	a	405	CLA	CMD-C2D-C1D	5.41	134.24	124.71

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
29	b	616	CLA	CMD-C2D-C1D	5.41	134.24	124.71
29	y	603	CLA	CMD-C2D-C1D	5.41	134.24	124.71
29	Y	603	CLA	CMD-C2D-C1D	5.41	134.24	124.71
29	g	610	CLA	O2A-C1-C2	5.40	122.84	108.64
46	r	620	LUT	C15-C14-C13	-5.40	119.60	127.31
29	g	604	CLA	O2A-C1-C2	5.40	121.59	108.97
29	y	602	CLA	O2D-CGD-CBD	5.39	120.85	111.27
29	b	614	CLA	CMD-C2D-C1D	5.39	134.22	124.71
29	r	604	CLA	CMD-C2D-C1D	5.39	134.22	124.71
29	G	611	CLA	CMD-C2D-C1D	5.39	134.21	124.71
46	r	620	LUT	C21-C26-C27	5.39	119.51	112.70
29	g	614	CLA	O2A-C1-C2	5.38	121.56	108.97
29	B	616	CLA	CMD-C2D-C1D	5.38	134.19	124.71
29	n	602	CLA	CMD-C2D-C1D	5.38	134.19	124.71
29	b	611	CLA	O2D-CGD-CBD	5.38	120.82	111.27
29	C	505	CLA	O2A-C1-C2	5.38	122.77	108.64
29	C	504	CLA	O2D-CGD-CBD	5.38	120.82	111.27
46	r	620	LUT	C35-C34-C33	-5.37	119.64	127.31
29	y	610	CLA	CMD-C2D-C1D	5.37	134.18	124.71
29	C	513	CLA	O2D-CGD-CBD	5.37	120.81	111.27
29	D	402	CLA	CMD-C2D-C1D	5.37	134.18	124.71
46	N	621	LUT	C21-C26-C25	5.37	121.04	111.42
29	N	603	CLA	CMD-C2D-C1D	5.37	134.17	124.71
46	S	620	LUT	C21-C26-C27	5.37	119.48	112.70
29	N	610	CLA	CMD-C2D-C1D	5.37	134.17	124.71
29	C	510	CLA	O2D-CGD-CBD	5.37	120.80	111.27
29	g	611	CLA	O2D-CGD-CBD	5.36	120.80	111.27
47	g	622	XAT	C31-C30-C29	-5.36	119.66	127.31
29	y	612	CLA	O2D-CGD-CBD	5.36	120.80	111.27
46	R	620	LUT	C15-C14-C13	-5.36	119.66	127.31
29	b	611	CLA	CMD-C2D-C1D	5.36	134.16	124.71
29	R	612	CLA	O2D-CGD-CBD	5.36	120.78	111.27
29	B	615	CLA	CMD-C2D-C1D	5.35	134.15	124.71
29	N	614	CLA	CMD-C2D-C1D	5.35	134.14	124.71
29	n	614	CLA	CMD-C2D-C1D	5.35	134.14	124.71
29	D	403	CLA	O2A-C1-C2	5.35	122.69	108.64
29	s	602	CLA	O2A-C1-C2	5.35	122.69	108.64
29	s	605	CLA	O2A-C1-C2	5.35	122.69	108.64
29	C	505	CLA	O2D-CGD-CBD	5.35	120.77	111.27
29	R	610	CLA	O2A-C1-C2	5.35	122.69	108.64
29	a	407	CLA	CMD-C2D-C1D	5.35	134.13	124.71
29	g	611	CLA	CMD-C2D-C1D	5.35	134.13	124.71

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
29	y	602	CLA	CMD-C2D-C1D	5.34	134.13	124.71
29	Y	608	CLA	O2A-C1-C2	5.34	122.68	108.64
29	b	610	CLA	CMD-C2D-C1D	5.34	134.12	124.71
29	b	608	CLA	CMD-C2D-C1D	5.34	134.12	124.71
29	n	610	CLA	CMD-C2D-C1D	5.34	134.12	124.71
29	n	611	CLA	CMD-C2D-C1D	5.34	134.12	124.71
47	N	622	XAT	C15-C14-C13	-5.34	119.70	127.31
29	C	510	CLA	O2A-C1-C2	5.33	122.65	108.64
29	g	612	CLA	O2D-CGD-CBD	5.33	120.75	111.27
29	y	611	CLA	O2D-CGD-CBD	5.33	120.74	111.27
29	D	403	CLA	O2D-CGD-CBD	5.32	120.73	111.27
29	c	510	CLA	O2D-CGD-CBD	5.32	120.73	111.27
29	b	606	CLA	O2D-CGD-CBD	5.32	120.73	111.27
29	s	605	CLA	CMD-C2D-C1D	5.32	134.09	124.71
29	Y	602	CLA	O2D-CGD-CBD	5.32	120.72	111.27
29	n	610	CLA	O2A-C1-C2	5.32	122.62	108.64
29	G	614	CLA	O2A-C1-C2	5.31	121.40	108.97
46	g	620	LUT	C21-C26-C25	5.31	120.94	111.42
29	g	603	CLA	O2D-CGD-CBD	5.31	120.71	111.27
29	c	513	CLA	O2D-CGD-CBD	5.31	120.70	111.27
29	R	611	CLA	CMD-C2D-C1D	5.31	134.06	124.71
29	S	611	CLA	CMD-C2D-C1D	5.31	134.06	124.71
29	C	511	CLA	O2D-CGD-CBD	5.30	120.69	111.27
29	G	602	CLA	O2D-CGD-CBD	5.30	120.69	111.27
29	N	602	CLA	O2D-CGD-CBD	5.30	120.68	111.27
29	B	606	CLA	O2D-CGD-CBD	5.30	120.68	111.27
47	n	622	XAT	C15-C14-C13	-5.30	119.75	127.31
29	B	610	CLA	CMD-C2D-C1D	5.30	134.04	124.71
29	c	507	CLA	O2D-CGD-CBD	5.29	120.68	111.27
29	C	511	CLA	O2A-C1-C2	5.29	122.55	108.64
29	B	611	CLA	O2D-CGD-CBD	5.29	120.67	111.27
29	c	504	CLA	CMD-C2D-C1D	5.29	134.04	124.71
29	n	603	CLA	CMD-C2D-C1D	5.29	134.03	124.71
29	N	614	CLA	O2D-CGD-CBD	5.29	120.67	111.27
29	N	613	CLA	CMD-C2D-C1D	5.28	134.03	124.71
29	d	402	CLA	CMD-C2D-C1D	5.28	134.03	124.71
29	b	610	CLA	O2D-CGD-CBD	5.28	120.66	111.27
29	r	604	CLA	O2D-CGD-CBD	5.28	120.66	111.27
29	a	410	CLA	CMD-C2D-C1D	5.28	134.01	124.71
29	b	615	CLA	CMD-C2D-C1D	5.28	134.01	124.71
29	N	613	CLA	O2A-C1-C2	5.27	122.49	108.64
29	B	608	CLA	CMD-C2D-C1D	5.27	134.00	124.71

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
29	c	511	CLA	O2A-C1-C2	5.27	122.48	108.64
29	S	610	CLA	O2A-C1-C2	5.27	122.48	108.64
29	B	611	CLA	CMD-C2D-C1D	5.27	134.00	124.71
46	N	620	LUT	C21-C26-C25	5.27	120.85	111.42
29	s	614	CLA	O2D-CGD-CBD	5.26	120.62	111.27
29	y	608	CLA	O2A-C1-C2	5.26	122.47	108.64
47	G	622	XAT	C31-C30-C29	-5.26	119.80	127.31
29	B	613	CLA	CMD-C2D-C1D	5.25	133.97	124.71
29	n	613	CLA	CMD-C2D-C1D	5.25	133.97	124.71
29	s	612	CLA	CMD-C2D-C1D	5.25	133.97	124.71
29	n	612	CLA	O2D-CGD-CBD	5.25	120.60	111.27
46	s	620	LUT	C21-C26-C25	5.25	120.82	111.42
29	r	610	CLA	O2A-C1-C2	5.25	122.43	108.64
29	N	611	CLA	CMD-C2D-C1D	5.25	133.96	124.71
29	B	602	CLA	O2D-CGD-CBD	5.25	120.59	111.27
29	Y	610	CLA	CMD-C2D-C1D	5.25	133.96	124.71
29	b	611	CLA	O2A-C1-C2	5.25	122.42	108.64
29	b	610	CLA	O2A-C1-C2	5.24	122.42	108.64
29	g	602	CLA	CMD-C2D-C1D	5.24	133.95	124.71
29	n	614	CLA	O2D-CGD-CBD	5.24	120.58	111.27
29	c	513	CLA	CMD-C2D-C1D	5.23	133.94	124.71
29	g	612	CLA	CMD-C2D-C1D	5.23	133.94	124.71
29	C	512	CLA	O2A-C1-C2	5.23	122.39	108.64
29	Y	610	CLA	O2D-CGD-CBD	5.23	120.57	111.27
29	G	612	CLA	O2D-CGD-CBD	5.23	120.56	111.27
29	n	602	CLA	O2D-CGD-CBD	5.23	120.56	111.27
29	b	602	CLA	O2D-CGD-CBD	5.23	120.56	111.27
29	s	609	CLA	CMD-C2D-C1D	5.23	133.93	124.71
29	c	501	CLA	O2A-C1-C2	5.23	122.38	108.64
29	s	612	CLA	O2D-CGD-CBD	5.23	120.55	111.27
29	b	613	CLA	CMD-C2D-C1D	5.22	133.91	124.71
29	B	610	CLA	O2A-C1-C2	5.22	122.35	108.64
29	s	602	CLA	CMD-C2D-C1D	5.22	133.91	124.71
29	c	510	CLA	O2A-C1-C2	5.22	122.35	108.64
29	G	603	CLA	O2D-CGD-CBD	5.22	120.54	111.27
29	b	612	CLA	O2A-C1-C2	5.21	122.34	108.64
29	d	403	CLA	O2A-C1-C2	5.21	122.34	108.64
29	s	611	CLA	CMD-C2D-C1D	5.21	133.90	124.71
29	r	612	CLA	O2D-CGD-CBD	5.21	120.52	111.27
29	b	608	CLA	O2D-CGD-CBD	5.21	120.52	111.27
29	R	610	CLA	O2D-CGD-CBD	5.21	120.52	111.27
29	N	602	CLA	CMD-C2D-C1D	5.20	133.89	124.71

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
29	C	501	CLA	O2A-C1-C2	5.20	122.31	108.64
29	B	608	CLA	O2D-CGD-CBD	5.20	120.51	111.27
29	b	604	CLA	CMD-C2D-C1D	5.20	133.88	124.71
29	r	608	CLA	O2A-C1-C2	5.20	122.31	108.64
29	R	602	CLA	O2A-C1-C2	5.20	122.30	108.64
29	r	602	CLA	O2D-CGD-CBD	5.20	120.50	111.27
29	B	610	CLA	O2D-CGD-CBD	5.20	120.50	111.27
29	C	507	CLA	O2D-CGD-CBD	5.20	120.50	111.27
29	c	508	CLA	CMD-C2D-C1D	5.20	133.87	124.71
29	R	603	CLA	O2D-CGD-CBD	5.19	120.49	111.27
29	b	603	CLA	O2D-CGD-CBD	5.19	120.49	111.27
29	c	505	CLA	O2D-CGD-CBD	5.18	120.48	111.27
29	y	611	CLA	CMD-C2D-C1D	5.18	133.85	124.71
29	c	504	CLA	O2D-CGD-CBD	5.18	120.48	111.27
29	c	511	CLA	O2D-CGD-CBD	5.18	120.48	111.27
29	S	602	CLA	O2A-C1-C2	5.18	122.25	108.64
29	S	617	CLA	CMD-C2D-C1D	5.18	133.84	124.71
29	c	512	CLA	O2A-C1-C2	5.18	122.25	108.64
46	N	621	LUT	C21-C26-C27	5.18	119.25	112.70
29	Y	614	CLA	O2D-CGD-CBD	5.18	120.47	111.27
29	N	614	CLA	O2A-C1-C2	5.17	121.07	108.97
29	B	605	CLA	O2D-CGD-CBD	5.17	120.45	111.27
29	g	613	CLA	O2A-C1-C2	5.17	122.22	108.64
29	B	614	CLA	O2A-C1-C2	5.17	122.22	108.64
29	S	602	CLA	CMD-C2D-C1D	5.16	133.81	124.71
29	C	502	CLA	CMD-C2D-C1D	5.16	133.81	124.71
29	c	505	CLA	O2A-C1-C2	5.16	122.19	108.64
29	y	614	CLA	O2D-CGD-CBD	5.16	120.43	111.27
29	A	410	CLA	CMD-C2D-C1D	5.16	133.80	124.71
29	S	617	CLA	O2D-CGD-CBD	5.15	120.43	111.27
29	D	403	CLA	CMD-C2D-C1D	5.15	133.79	124.71
29	c	503	CLA	O2A-C1-C2	5.15	122.17	108.64
29	G	612	CLA	CMD-C2D-C1D	5.14	133.78	124.71
29	B	604	CLA	CMD-C2D-C1D	5.14	133.78	124.71
29	G	610	CLA	O2D-CGD-CBD	5.14	120.41	111.27
29	c	508	CLA	O2A-C1-C2	5.14	122.14	108.64
29	B	616	CLA	O2D-CGD-CBD	5.14	120.39	111.27
29	B	611	CLA	O2A-C1-C2	5.13	122.13	108.64
29	y	612	CLA	CMD-C2D-C1D	5.13	133.76	124.71
29	S	609	CLA	O2D-CGD-CBD	5.13	120.39	111.27
29	G	613	CLA	O2A-C1-C2	5.13	122.11	108.64
29	S	604	CLA	O2D-CGD-CBD	5.13	120.38	111.27

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
29	Y	610	CLA	O2A-C1-C2	5.12	122.10	108.64
29	C	505	CLA	CMD-C2D-C1D	5.12	133.74	124.71
29	R	608	CLA	O2A-C1-C2	5.12	122.08	108.64
29	g	614	CLA	O2D-CGD-CBD	5.12	120.36	111.27
29	Y	602	CLA	CMD-C2D-C1D	5.12	133.73	124.71
29	C	508	CLA	O2A-C1-C2	5.12	122.08	108.64
29	a	410	CLA	O2A-C1-C2	5.11	122.07	108.64
47	g	622	XAT	C15-C14-C13	-5.11	120.02	127.31
29	S	617	CLA	O2A-C1-C2	5.11	122.06	108.64
29	A	410	CLA	O2A-C1-C2	5.11	122.06	108.64
29	S	613	CLA	O2A-C1-C2	5.11	122.06	108.64
29	r	612	CLA	O2A-C1-C2	5.11	122.06	108.64
29	Y	612	CLA	CMD-C2D-C1D	5.11	133.71	124.71
46	r	620	LUT	C31-C30-C29	-5.11	120.02	127.31
29	B	617	CLA	O2D-CGD-CBD	5.10	120.33	111.27
29	b	605	CLA	O2D-CGD-CBD	5.10	120.33	111.27
46	S	621	LUT	C21-C26-C25	5.10	120.55	111.42
29	y	610	CLA	O2A-C1-C2	5.10	122.04	108.64
29	b	606	CLA	O2A-C1-C2	5.10	122.03	108.64
29	C	508	CLA	CMD-C2D-C1D	5.09	133.69	124.71
29	S	604	CLA	O2A-C1-C2	5.09	122.02	108.64
29	b	614	CLA	O2A-C1-C2	5.09	122.01	108.64
29	b	616	CLA	O2D-CGD-CBD	5.09	120.31	111.27
29	N	602	CLA	O2A-C1-C2	5.09	122.01	108.64
29	s	611	CLA	O2D-CGD-CBD	5.09	120.31	111.27
29	B	609	CLA	CMD-C2D-C1D	5.09	133.68	124.71
29	S	605	CLA	CMD-C2D-C1D	5.08	133.67	124.71
29	S	612	CLA	CMD-C2D-C1D	5.08	133.67	124.71
29	c	505	CLA	CMD-C2D-C1D	5.08	133.67	124.71
46	G	621	LUT	C21-C26-C27	5.08	119.12	112.70
29	A	407	CLA	CMD-C2D-C1D	5.08	133.66	124.71
29	S	614	CLA	O2D-CGD-CBD	5.07	120.29	111.27
29	Y	613	CLA	O2A-C1-C2	5.07	121.96	108.64
46	R	620	LUT	C35-C34-C33	-5.07	120.08	127.31
29	B	606	CLA	O2A-C1-C2	5.06	121.94	108.64
46	g	621	LUT	C21-C26-C27	5.06	119.10	112.70
29	n	602	CLA	O2A-C1-C2	5.06	121.93	108.64
29	s	617	CLA	O2D-CGD-CBD	5.06	120.26	111.27
29	N	611	CLA	O2A-C1-C2	5.05	120.79	108.97
47	G	622	XAT	C15-C14-C13	-5.05	120.10	127.31
29	S	614	CLA	O2A-C1-C2	5.05	121.92	108.64
29	a	406	CLA	O2A-C1-C2	5.04	121.89	108.64

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
29	B	615	CLA	O2D-CGD-CBD	5.04	120.22	111.27
29	s	613	CLA	O2A-C1-C2	5.04	121.87	108.64
29	S	611	CLA	O2D-CGD-CBD	5.03	120.21	111.27
29	S	602	CLA	O2D-CGD-CBD	5.03	120.21	111.27
29	R	612	CLA	O2A-C1-C2	5.02	121.84	108.64
29	b	616	CLA	O2A-C1-C2	5.02	121.82	108.64
29	s	617	CLA	O2A-C1-C2	5.01	121.81	108.64
29	d	403	CLA	CMD-C2D-C1D	5.01	133.54	124.71
29	N	612	CLA	CMD-C2D-C1D	5.01	133.53	124.71
29	y	613	CLA	O2A-C1-C2	4.99	121.76	108.64
29	s	610	CLA	O2D-CGD-CBD	4.99	120.14	111.27
29	N	603	CLA	O2A-C1-C2	4.99	121.75	108.64
29	S	610	CLA	O2D-CGD-CBD	4.99	120.14	111.27
29	S	612	CLA	O2D-CGD-CBD	4.99	120.14	111.27
29	s	609	CLA	O2D-CGD-CBD	4.98	120.11	111.27
29	B	612	CLA	CMD-C2D-C1D	4.98	133.49	124.71
29	c	507	CLA	O2A-C1-C2	4.98	121.72	108.64
29	C	502	CLA	O2D-CGD-CBD	4.98	120.11	111.27
29	A	406	CLA	O2A-C1-C2	4.97	121.71	108.64
47	Y	622	XAT	C31-C30-C29	-4.97	120.22	127.31
29	N	610	CLA	O2D-CGD-CBD	4.97	120.09	111.27
29	y	610	CLA	O2D-CGD-CBD	4.96	120.09	111.27
35	b	620	C7Z	C15-C14-C13	-4.96	120.24	127.31
46	G	620	LUT	C21-C26-C25	4.95	120.29	111.42
29	G	614	CLA	O2D-CGD-CBD	4.95	120.06	111.27
29	R	609	CLA	O2D-CGD-CBD	4.94	120.05	111.27
29	r	603	CLA	O2D-CGD-CBD	4.94	120.05	111.27
29	r	609	CLA	O2D-CGD-CBD	4.94	120.05	111.27
29	b	604	CLA	O2D-CGD-CBD	4.94	120.05	111.27
29	Y	612	CLA	O2A-C1-C2	4.94	121.61	108.64
29	b	602	CLA	O2A-C1-C2	4.93	121.60	108.64
29	b	609	CLA	CMD-C2D-C1D	4.93	133.40	124.71
29	s	603	CLA	O2D-CGD-CBD	4.93	120.03	111.27
29	c	506	CLA	O2A-C1-C2	4.93	121.58	108.64
29	S	603	CLA	CMD-C2D-C1D	4.92	133.39	124.71
29	B	604	CLA	O2D-CGD-CBD	4.92	120.01	111.27
29	C	507	CLA	O2A-C1-C2	4.92	121.56	108.64
29	Y	608	CLA	O2D-CGD-CBD	4.92	120.01	111.27
29	b	617	CLA	O2D-CGD-CBD	4.91	120.00	111.27
29	D	402	CLA	O2D-CGD-CBD	4.91	120.00	111.27
29	c	512	CLA	O2D-CGD-CBD	4.91	120.00	111.27
29	b	615	CLA	O2D-CGD-CBD	4.91	120.00	111.27

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
29	B	616	CLA	O2A-C1-C2	4.91	121.54	108.64
29	n	610	CLA	O2D-CGD-CBD	4.91	119.99	111.27
29	d	402	CLA	O2D-CGD-CBD	4.91	119.98	111.27
29	N	612	CLA	O2D-CGD-CBD	4.90	119.98	111.27
29	B	603	CLA	O2D-CGD-CBD	4.90	119.98	111.27
29	b	612	CLA	CMD-C2D-C1D	4.90	133.35	124.71
48	R	622	NEX	C38-C25-C24	4.90	119.79	114.28
29	B	609	CLA	O2A-C1-C2	4.89	121.50	108.64
29	G	613	CLA	O2D-CGD-CBD	4.89	119.96	111.27
29	A	407	CLA	O2D-CGD-CBD	4.89	119.96	111.27
29	r	610	CLA	O2D-CGD-CBD	4.89	119.96	111.27
29	s	602	CLA	O2D-CGD-CBD	4.88	119.94	111.27
29	b	607	CLA	O2A-C1-C2	4.88	121.46	108.64
29	y	612	CLA	O2A-C1-C2	4.88	121.45	108.64
29	S	603	CLA	O2A-C1-C2	4.88	121.45	108.64
29	y	608	CLA	O2D-CGD-CBD	4.87	119.92	111.27
29	Y	604	CLA	O2D-CGD-CBD	4.87	119.92	111.27
29	y	613	CLA	O2D-CGD-CBD	4.87	119.92	111.27
29	B	612	CLA	O2A-C1-C2	4.87	121.43	108.64
48	r	623	NEX	C38-C25-C24	4.86	119.75	114.28
46	R	620	LUT	C11-C10-C9	-4.86	120.37	127.31
29	s	603	CLA	CMD-C2D-C1D	4.86	133.28	124.71
46	n	620	LUT	C21-C26-C25	4.86	120.12	111.42
29	B	612	CLA	O2D-CGD-CBD	4.85	119.89	111.27
29	a	406	CLA	O2D-CGD-CBD	4.85	119.88	111.27
29	Y	604	CLA	O2A-C1-C2	4.85	121.37	108.64
29	r	602	CLA	O2A-C1-C2	4.85	121.37	108.64
47	y	622	XAT	C15-C14-C13	-4.85	120.39	127.31
29	R	611	CLA	O2D-CGD-CBD	4.84	119.87	111.27
29	s	603	CLA	O2A-C1-C2	4.84	121.35	108.64
29	b	612	CLA	O2D-CGD-CBD	4.83	119.85	111.27
29	Y	613	CLA	O2D-CGD-CBD	4.83	119.85	111.27
39	n	624	LHG	O7-C7-C8	4.83	121.91	111.50
29	S	609	CLA	O2A-C1-C2	4.83	121.32	108.64
29	c	502	CLA	C1-C2-C3	-4.83	117.70	126.04
29	s	609	CLA	O2A-C1-C2	4.82	121.31	108.64
29	y	604	CLA	O2A-C1-C2	4.82	121.30	108.64
46	R	620	LUT	C21-C26-C27	4.82	118.79	112.70
29	N	613	CLA	O2D-CGD-CBD	4.81	119.82	111.27
29	y	602	CLA	O2A-C1-C2	4.81	121.28	108.64
29	B	607	CLA	O2A-C1-C2	4.81	121.28	108.64
46	S	620	LUT	C21-C26-C25	4.81	120.03	111.42

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
29	Y	611	CLA	O2A-C1-C2	4.81	121.26	108.64
29	S	611	CLA	O2A-C1-C2	4.80	121.26	108.64
29	r	608	CLA	O2D-CGD-CBD	4.80	119.80	111.27
29	N	610	CLA	O2A-C1-C2	4.80	121.25	108.64
35	B	620	C7Z	C18-C5-C6	-4.80	119.14	124.53
29	c	508	CLA	O2D-CGD-CBD	4.80	119.79	111.27
29	R	608	CLA	O2D-CGD-CBD	4.79	119.79	111.27
29	b	614	CLA	O2D-CGD-CBD	4.78	119.77	111.27
29	C	503	CLA	O2A-C1-C2	4.78	121.20	108.64
29	g	613	CLA	O2D-CGD-CBD	4.78	119.76	111.27
29	Y	611	CLA	O2D-CGD-CBD	4.77	119.75	111.27
29	s	604	CLA	O2D-CGD-CBD	4.77	119.75	111.27
29	B	602	CLA	O2A-C1-C2	4.77	121.17	108.64
39	N	624	LHG	O7-C7-C8	4.77	121.78	111.50
29	g	610	CLA	O2D-CGD-CBD	4.76	119.73	111.27
46	r	620	LUT	C11-C10-C9	-4.76	120.52	127.31
29	n	613	CLA	O2D-CGD-CBD	4.75	119.71	111.27
29	C	512	CLA	O2D-CGD-CBD	4.75	119.71	111.27
29	a	407	CLA	O2D-CGD-CBD	4.75	119.71	111.27
35	B	620	C7Z	C15-C14-C13	-4.75	120.53	127.31
48	N	623	NEX	C17-C1-C6	-4.75	106.22	110.47
29	s	611	CLA	O2A-C1-C2	4.75	121.11	108.64
29	b	613	CLA	O2A-C1-C2	4.74	121.09	108.64
42	D	405	PL9	C7-C3-C4	4.73	120.72	116.88
46	S	621	LUT	C35-C34-C33	-4.73	120.56	127.31
29	Y	614	CLA	O2A-C1-C2	4.73	121.07	108.64
29	C	508	CLA	O2D-CGD-CBD	4.72	119.66	111.27
48	N	623	NEX	C2-C1-C6	4.71	113.79	109.21
29	Y	611	CLA	CMD-C2D-C1D	4.71	133.01	124.71
29	C	506	CLA	O2A-C1-C2	4.70	120.99	108.64
29	r	611	CLA	O2D-CGD-CBD	4.69	119.60	111.27
48	s	623	NEX	C38-C25-C24	4.69	119.56	114.28
29	Y	602	CLA	O2A-C1-C2	4.69	120.95	108.64
29	b	617	CLA	CMD-C2D-C1D	4.68	132.96	124.71
48	S	622	NEX	C38-C25-C24	4.68	119.54	114.28
48	g	623	NEX	C38-C25-C24	4.67	119.54	114.28
29	y	604	CLA	O2D-CGD-CBD	4.67	119.57	111.27
29	C	509	CLA	O2A-C1-C2	4.67	120.91	108.64
29	y	611	CLA	O2A-C1-C2	4.66	120.89	108.64
47	N	622	XAT	C18-C5-C4	4.66	119.53	114.28
29	S	603	CLA	O2D-CGD-CBD	4.66	119.55	111.27
48	s	623	NEX	C2-C1-C6	4.66	113.74	109.21

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
48	n	623	NEX	C2-C1-C6	4.66	113.74	109.21
47	R	621	XAT	C38-C25-C24	4.65	119.51	114.28
29	B	617	CLA	CMD-C2D-C1D	4.65	132.90	124.71
29	B	614	CLA	O2D-CGD-CBD	4.64	119.51	111.27
48	Y	623	NEX	C2-C1-C6	4.63	113.71	109.21
29	b	609	CLA	O2A-C1-C2	4.63	120.80	108.64
47	N	622	XAT	C38-C25-C24	4.62	119.48	114.28
29	y	614	CLA	O2A-C1-C2	4.61	120.74	108.64
47	n	622	XAT	C38-C25-C24	4.60	119.46	114.28
29	C	512	CLA	CMD-C2D-C1D	4.60	132.82	124.71
29	c	512	CLA	CMD-C2D-C1D	4.59	132.80	124.71
47	n	622	XAT	C18-C5-C4	4.58	119.43	114.28
48	N	623	NEX	C38-C25-C24	4.58	119.43	114.28
46	Y	621	LUT	C21-C26-C27	4.58	118.48	112.70
48	S	622	NEX	C17-C1-C6	-4.57	106.38	110.47
47	G	622	XAT	C18-C5-C4	4.57	119.42	114.28
48	G	623	NEX	C38-C25-C24	4.57	119.42	114.28
29	r	603	CLA	O2A-C1-C2	4.55	120.60	108.64
29	A	406	CLA	O2D-CGD-CBD	4.55	119.35	111.27
46	S	621	LUT	C21-C26-C27	4.54	118.44	112.70
48	S	622	NEX	C2-C1-C6	4.54	113.62	109.21
29	B	604	CLA	O2A-C1-C2	4.53	120.54	108.64
46	s	621	LUT	C35-C34-C33	-4.53	120.85	127.31
48	G	623	NEX	C2-C1-C6	4.52	113.61	109.21
29	g	603	CLA	O2A-C1-C2	4.52	120.52	108.64
46	R	620	LUT	C31-C30-C29	-4.52	120.86	127.31
46	y	621	LUT	C35-C34-C33	-4.52	120.86	127.31
29	a	405	CLA	O2D-CGD-CBD	4.51	119.29	111.27
29	G	603	CLA	O2A-C1-C2	4.51	120.50	108.64
46	G	620	LUT	C21-C26-C27	4.51	118.40	112.70
46	Y	620	LUT	C21-C26-C27	4.50	118.39	112.70
48	Y	623	NEX	C38-C25-C24	4.49	119.33	114.28
29	b	604	CLA	O2A-C1-C2	4.48	120.42	108.64
42	d	405	PL9	C7-C3-C4	4.48	120.52	116.88
29	A	405	CLA	O2D-CGD-CBD	4.47	119.21	111.27
47	y	622	XAT	C18-C5-C4	4.46	119.30	114.28
48	y	623	NEX	C38-C25-C24	4.45	119.29	114.28
31	c	516	BCR	C33-C5-C6	-4.43	119.56	124.53
29	B	613	CLA	O2A-C1-C2	4.42	120.26	108.64
31	C	516	BCR	C33-C5-C6	-4.42	119.56	124.53
38	C	519	DGD	O2G-C1B-C2B	4.42	121.03	111.50
48	n	623	NEX	C38-C25-C24	4.42	119.25	114.28

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
46	S	620	LUT	C35-C34-C33	-4.41	121.02	127.31
48	g	623	NEX	C2-C1-C6	4.41	113.50	109.21
46	R	620	LUT	C7-C8-C9	-4.39	119.59	126.23
39	D	408	LHG	O7-C7-C8	4.38	120.94	111.50
46	n	621	LUT	C21-C26-C27	4.37	118.23	112.70
29	s	604	CLA	O2A-C1-C2	4.37	120.13	108.64
47	Y	622	XAT	C18-C5-C4	4.37	119.19	114.28
29	c	509	CLA	O2A-C1-C2	4.35	120.06	108.64
29	B	615	CLA	CAA-C2A-C3A	-4.33	100.91	112.78
47	y	622	XAT	C38-C25-C24	4.33	119.15	114.28
46	N	620	LUT	C21-C26-C27	4.33	118.17	112.70
39	d	408	LHG	O7-C7-C8	4.32	120.82	111.50
39	S	624	LHG	O7-C7-C8	4.32	120.81	111.50
29	Y	603	CLA	O2A-C1-C2	4.32	119.98	108.64
35	b	620	C7Z	C18-C5-C6	-4.31	119.68	124.53
29	R	603	CLA	O2A-C1-C2	4.30	119.94	108.64
46	g	620	LUT	C7-C8-C9	-4.29	119.75	126.23
46	N	621	LUT	C22-C23-C24	-4.29	106.86	111.74
46	y	620	LUT	C21-C26-C27	4.29	118.12	112.70
31	C	515	BCR	C15-C14-C13	-4.27	121.22	127.31
46	r	620	LUT	C7-C8-C9	-4.27	119.78	126.23
47	g	622	XAT	C18-C5-C4	4.27	119.08	114.28
39	G	630	LHG	O7-C7-C8	4.26	120.69	111.50
47	r	622	XAT	C38-C25-C24	4.26	119.08	114.28
46	g	621	LUT	C22-C23-C24	-4.25	106.90	111.74
46	s	620	LUT	C22-C23-C24	-4.25	106.91	111.74
45	g	609	CHL	CHD-C1D-ND	-4.24	120.56	124.45
38	C	520	DGD	O2G-C1B-C2B	4.23	120.61	111.50
39	s	624	LHG	O7-C7-C8	4.23	120.61	111.50
47	Y	622	XAT	C38-C25-C24	4.22	119.03	114.28
29	b	612	CLA	CAA-C2A-C3A	-4.22	101.22	112.78
33	a	413	LMG	O7-C10-C11	4.22	120.59	111.50
46	R	620	LUT	C21-C26-C25	4.22	118.97	111.42
38	c	519	DGD	O2G-C1B-C2B	4.21	120.58	111.50
48	Y	623	NEX	C17-C1-C6	-4.20	106.71	110.47
33	A	413	LMG	O7-C10-C11	4.20	120.55	111.50
38	C	523	DGD	O2G-C1B-C2B	4.18	120.51	111.50
29	b	615	CLA	CAA-C2A-C3A	-4.17	101.35	112.78
46	Y	620	LUT	C15-C14-C13	-4.17	121.36	127.31
44	H	101	RRX	C11-C10-C9	-4.17	121.36	127.31
47	G	622	XAT	C38-C25-C24	4.16	118.96	114.28
38	c	520	DGD	O2G-C1B-C2B	4.16	120.47	111.50

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
46	n	620	LUT	C21-C26-C27	4.14	117.94	112.70
46	y	621	LUT	C7-C8-C9	-4.14	119.97	126.23
47	n	622	XAT	C7-C8-C9	-4.13	119.11	125.53
46	y	621	LUT	C21-C26-C27	4.13	117.92	112.70
47	R	621	XAT	C18-C5-C4	4.13	118.92	114.28
48	y	623	NEX	C2-C1-C6	4.13	113.22	109.21
46	G	620	LUT	C7-C8-C9	-4.12	120.00	126.23
48	s	623	NEX	C17-C1-C6	-4.12	106.78	110.47
33	h	102	LMG	O7-C10-C11	4.12	120.38	111.50
29	y	603	CLA	O2A-C1-C2	4.12	119.46	108.64
48	R	622	NEX	C2-C1-C6	4.11	113.21	109.21
46	y	621	LUT	C22-C23-C24	-4.11	107.06	111.74
47	g	622	XAT	C38-C25-C24	4.11	118.90	114.28
46	s	620	LUT	C35-C34-C33	-4.11	121.45	127.31
39	g	624	LHG	O7-C7-C8	4.10	120.34	111.50
46	Y	620	LUT	C22-C23-C24	-4.10	107.08	111.74
50	i	101	3PH	O21-C21-C22	4.09	120.33	111.50
35	b	620	C7Z	C38-C25-C26	-4.09	119.93	124.53
46	Y	621	LUT	C7-C8-C9	-4.09	120.05	126.23
33	C	521	LMG	O7-C10-C11	4.09	120.31	111.50
47	Y	622	XAT	C36-C21-C26	4.08	121.06	110.05
38	c	523	DGD	O2G-C1B-C2B	4.08	120.30	111.50
39	d	410	LHG	O7-C7-C8	4.07	120.28	111.50
45	N	609	CHL	CHD-C1D-ND	-4.07	120.71	124.45
33	H	102	LMG	O7-C10-C11	4.07	120.26	111.50
29	c	502	CLA	O2D-CGD-CBD	4.06	118.48	111.27
33	c	521	LMG	O7-C10-C11	4.05	120.22	111.50
31	D	404	BCR	C19-C18-C17	4.03	125.13	118.94
44	h	101	RRX	C11-C10-C9	-4.01	121.58	127.31
50	S	626	3PH	O21-C21-C22	4.01	120.15	111.50
32	A	412	SQD	O7-S-C6	-4.00	102.18	106.94
46	Y	621	LUT	C22-C23-C24	-3.98	107.21	111.74
50	s	626	3PH	O21-C21-C22	3.98	120.08	111.50
36	C	524	DGA	OG2-CB1-CB2	3.98	120.07	111.50
31	d	404	BCR	C19-C18-C17	3.97	125.04	118.94
48	g	623	NEX	C27-C28-C29	-3.97	119.36	125.53
46	G	621	LUT	C22-C23-C24	-3.97	107.23	111.74
39	C	525	LHG	O7-C7-C8	3.96	120.05	111.50
36	b	623	DGA	OG2-CB1-CB2	3.96	120.03	111.50
45	g	601	CHL	CHD-C1D-ND	-3.96	120.82	124.45
46	g	621	LUT	C7-C8-C9	-3.96	120.26	126.23
46	S	620	LUT	C7-C8-C9	-3.95	120.27	126.23

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
46	G	620	LUT	C15-C14-C13	-3.95	121.68	127.31
47	y	622	XAT	C36-C21-C26	3.94	120.68	110.05
36	c	524	DGA	OG2-CB1-CB2	3.94	119.99	111.50
38	c	518	DGD	O2G-C1B-C2B	3.94	119.98	111.50
46	N	620	LUT	C35-C34-C33	-3.92	121.71	127.31
29	B	605	CLA	C2C-C1C-NC	3.91	113.63	109.97
39	c	625	LHG	O7-C7-C8	3.90	119.91	111.50
30	a	409	PHO	CMB-C2B-C3B	3.90	131.97	124.68
46	n	621	LUT	C35-C34-C33	-3.90	121.75	127.31
46	r	620	LUT	C21-C26-C25	3.90	118.40	111.42
45	y	601	CHL	CHD-C1D-ND	-3.90	120.87	124.45
45	Y	605	CHL	CHD-C1D-ND	-3.89	120.88	124.45
46	g	621	LUT	C35-C34-C33	-3.89	121.77	127.31
29	C	502	CLA	C1-C2-C3	-3.88	119.33	126.04
29	B	606	CLA	CHD-C1D-ND	-3.87	120.90	124.45
46	y	620	LUT	C22-C23-C24	-3.87	107.34	111.74
29	b	605	CLA	C2C-C1C-NC	3.87	113.60	109.97
36	B	625	DGA	OG2-CB1-CB2	3.86	119.81	111.50
35	B	620	C7Z	C38-C25-C26	-3.86	120.20	124.53
33	B	622	LMG	O7-C10-C11	3.85	119.79	111.50
33	j	101	LMG	O7-C10-C11	3.85	119.79	111.50
46	g	620	LUT	C21-C26-C27	3.84	117.55	112.70
45	n	605	CHL	CHD-C1D-ND	-3.83	120.93	124.45
45	N	605	CHL	CHD-C1D-ND	-3.82	120.94	124.45
39	D	409	LHG	O7-C7-C8	3.82	119.74	111.50
46	N	620	LUT	C15-C14-C13	-3.82	121.85	127.31
46	G	621	LUT	C7-C8-C9	-3.82	120.47	126.23
48	Y	623	NEX	C27-C28-C29	-3.81	119.61	125.53
39	Y	624	LHG	O7-C7-C8	3.81	119.72	111.50
29	S	617	CLA	C1-C2-C3	-3.81	120.59	126.75
43	f	101	HEM	C4B-CHC-C1C	3.81	127.59	122.56
46	S	620	LUT	C22-C23-C24	-3.80	107.41	111.74
45	Y	601	CHL	CHD-C1D-ND	-3.80	120.96	124.45
46	G	621	LUT	C35-C34-C33	-3.80	121.89	127.31
45	y	605	CHL	CHD-C1D-ND	-3.80	120.96	124.45
46	s	621	LUT	C21-C26-C27	3.79	117.50	112.70
33	b	622	LMG	O7-C10-C11	3.79	119.67	111.50
30	A	409	PHO	CMB-C2B-C3B	3.79	131.76	124.68
39	d	409	LHG	O7-C7-C8	3.78	119.65	111.50
44	H	101	RRX	C7-C8-C9	-3.78	120.53	126.23
33	d	411	LMG	O7-C10-C11	3.78	119.64	111.50
29	C	509	CLA	C2C-C1C-NC	3.77	113.50	109.97

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
35	b	620	C7Z	C1-C6-C5	-3.77	117.31	122.61
47	r	622	XAT	C18-C5-C4	3.77	118.52	114.28
29	S	613	CLA	O2D-CGD-O1D	-3.77	116.47	123.84
33	D	411	LMG	O7-C10-C11	3.77	119.62	111.50
29	s	617	CLA	C1-C2-C3	-3.76	120.66	126.75
29	A	406	CLA	C2C-C1C-NC	3.76	113.50	109.97
39	D	410	LHG	O7-C7-C8	3.76	119.60	111.50
46	N	621	LUT	C35-C34-C33	-3.75	121.96	127.31
29	s	613	CLA	C1-C2-C3	-3.74	119.57	126.04
33	J	101	LMG	O7-C10-C11	3.74	119.57	111.50
48	n	623	NEX	C27-C28-C29	-3.74	119.72	125.53
47	N	622	XAT	C7-C8-C9	-3.74	119.73	125.53
46	g	620	LUT	C35-C34-C33	-3.73	121.98	127.31
29	B	605	CLA	C1-C2-C3	-3.73	119.59	126.04
32	a	412	SQD	O7-S-C6	-3.72	102.51	106.94
29	b	606	CLA	CHD-C1D-ND	-3.72	121.03	124.45
45	S	608	CHL	CHD-C1D-ND	-3.72	121.04	124.45
46	g	621	LUT	C15-C14-C13	-3.72	122.01	127.31
46	n	621	LUT	C22-C23-C24	-3.71	107.52	111.74
45	G	601	CHL	CHD-C1D-ND	-3.71	121.04	124.45
40	c	627	LMK	O3-C4-C3	-3.71	110.25	122.98
45	s	608	CHL	CHD-C1D-ND	-3.70	121.05	124.45
29	s	613	CLA	O2D-CGD-O1D	-3.70	116.60	123.84
32	B	621	SQD	O7-S-C6	-3.70	102.54	106.94
32	b	621	SQD	O7-S-C6	-3.70	102.55	106.94
45	R	607	CHL	CHD-C1D-ND	-3.70	121.06	124.45
46	n	621	LUT	C15-C14-C13	-3.69	122.04	127.31
31	d	404	BCR	C36-C18-C17	-3.69	117.75	122.92
47	R	621	XAT	C19-C9-C10	-3.69	117.75	122.92
35	B	620	C7Z	C35-C34-C33	-3.69	122.05	127.31
29	S	613	CLA	C1-C2-C3	-3.69	119.67	126.04
46	N	620	LUT	C22-C23-C24	-3.69	107.55	111.74
46	s	621	LUT	C15-C14-C13	-3.68	122.05	127.31
46	y	621	LUT	C11-C10-C9	-3.68	122.05	127.31
45	r	606	CHL	CHD-C1D-ND	-3.68	121.07	124.45
45	y	609	CHL	CHD-C1D-ND	-3.68	121.07	124.45
31	B	619	BCR	C37-C22-C21	-3.67	117.78	122.92
45	G	609	CHL	CHD-C1D-ND	-3.67	121.08	124.45
45	g	605	CHL	CHD-C1D-ND	-3.67	121.08	124.45
39	y	624	LHG	O7-C7-C8	3.67	119.41	111.50
46	S	620	LUT	C18-C5-C6	-3.66	120.41	124.53
45	n	608	CHL	CHD-C1D-ND	-3.66	121.09	124.45

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
48	G	623	NEX	C27-C28-C29	-3.66	119.86	125.53
45	n	606	CHL	CHD-C1D-ND	-3.65	121.10	124.45
44	h	101	RRX	C33-C5-C4	3.64	120.61	113.62
46	g	621	LUT	C11-C10-C9	-3.64	122.12	127.31
38	C	518	DGD	O2G-C1B-C2B	3.64	119.34	111.50
46	s	621	LUT	C7-C8-C9	-3.63	120.75	126.23
45	s	607	CHL	CHD-C1D-ND	-3.62	121.12	124.45
46	S	620	LUT	C15-C14-C13	-3.62	122.14	127.31
47	R	621	XAT	C26-C27-C28	-3.62	118.35	125.99
47	r	622	XAT	C38-C25-C26	-3.61	116.20	122.26
45	Y	609	CHL	CHD-C1D-ND	-3.61	121.14	124.45
48	S	622	NEX	C27-C28-C29	-3.61	119.93	125.53
32	c	626	SQD	O7-S-C6	-3.61	102.65	106.94
46	Y	620	LUT	C7-C8-C9	-3.60	120.79	126.23
45	G	605	CHL	CHD-C1D-ND	-3.60	121.14	124.45
46	g	620	LUT	C15-C14-C13	-3.60	122.17	127.31
29	B	603	CLA	CHD-C1D-ND	-3.60	121.14	124.45
45	n	601	CHL	CHD-C1D-ND	-3.60	121.14	124.45
48	y	623	NEX	C27-C28-C29	-3.58	119.98	125.53
31	D	404	BCR	C36-C18-C17	-3.58	117.91	122.92
31	b	619	BCR	C33-C5-C4	3.57	120.47	113.62
40	C	527	LMK	O3-C4-C3	-3.57	110.73	122.98
31	c	517	BCR	C28-C27-C26	-3.57	107.70	114.08
48	g	623	NEX	C17-C1-C6	-3.57	107.28	110.47
31	D	404	BCR	C37-C22-C21	-3.57	117.93	122.92
32	C	526	SQD	O7-S-C6	-3.57	102.70	106.94
45	s	601	CHL	CHD-C1D-ND	-3.56	121.18	124.45
46	s	620	LUT	C35-C15-C14	-3.56	116.18	123.47
45	g	608	CHL	CHD-C1D-ND	-3.56	121.18	124.45
31	A	411	BCR	C33-C5-C6	-3.55	120.54	124.53
48	s	623	NEX	C27-C28-C29	-3.55	120.02	125.53
47	N	622	XAT	O24-C25-C24	3.55	116.05	113.38
45	N	608	CHL	CHD-C1D-ND	-3.55	121.19	124.45
31	B	619	BCR	C33-C5-C4	3.54	120.43	113.62
29	B	605	CLA	CMA-C3A-C4A	3.54	121.30	111.77
29	g	613	CLA	CMA-C3A-C4A	3.54	121.30	111.77
35	b	620	C7Z	C35-C34-C33	-3.54	122.25	127.31
45	y	606	CHL	C2C-C3C-C4C	3.54	109.01	106.49
46	g	620	LUT	C22-C23-C24	-3.54	107.71	111.74
44	H	101	RRX	C33-C5-C4	3.54	120.41	113.62
29	N	610	CLA	CAA-C2A-C3A	-3.54	103.09	112.78
29	S	604	CLA	CHD-C1D-ND	-3.54	121.20	124.45

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
46	G	620	LUT	C11-C10-C9	-3.53	122.27	127.31
29	c	506	CLA	CHD-C1D-ND	-3.53	121.21	124.45
29	c	506	CLA	C2C-C1C-NC	3.53	113.28	109.97
45	G	608	CHL	CHD-C1D-ND	-3.53	121.21	124.45
29	C	508	CLA	C1-C2-C3	-3.52	119.95	126.04
46	s	620	LUT	C7-C8-C9	-3.52	120.91	126.23
29	c	512	CLA	C2D-C1D-ND	3.52	112.70	110.10
31	a	411	BCR	C33-C5-C6	-3.51	120.58	124.53
45	S	607	CHL	CHD-C1D-ND	-3.51	121.22	124.45
29	C	510	CLA	CHD-C1D-ND	-3.51	121.23	124.45
46	N	621	LUT	C15-C14-C13	-3.51	122.30	127.31
29	B	610	CLA	C1-C2-C3	-3.50	119.99	126.04
47	y	622	XAT	C7-C8-C9	-3.50	120.10	125.53
46	y	620	LUT	C35-C34-C33	-3.50	122.32	127.31
47	G	622	XAT	C7-C8-C9	-3.49	120.11	125.53
35	B	620	C7Z	C1-C6-C5	-3.49	117.69	122.61
44	h	101	RRX	C30-C25-C26	-3.49	117.69	122.61
47	R	621	XAT	C11-C10-C9	-3.49	122.33	127.31
29	G	613	CLA	CMA-C3A-C4A	3.49	121.16	111.77
29	C	505	CLA	O2D-CGD-O1D	-3.49	117.02	123.84
44	H	101	RRX	C30-C25-C26	-3.49	117.70	122.61
29	n	604	CLA	C1-C2-C3	-3.49	120.01	126.04
47	r	622	XAT	C19-C9-C10	-3.49	118.04	122.92
29	s	614	CLA	CHD-C1D-ND	-3.48	121.25	124.45
29	C	506	CLA	C2C-C1C-NC	3.48	113.23	109.97
29	r	610	CLA	CMA-C3A-C4A	3.48	121.12	111.77
46	y	620	LUT	C15-C14-C13	-3.47	122.35	127.31
29	c	502	CLA	C2C-C1C-NC	3.47	113.22	109.97
48	N	623	NEX	C27-C28-C29	-3.47	120.14	125.53
29	C	511	CLA	CHD-C1D-ND	-3.47	121.27	124.45
29	C	504	CLA	C1-C2-C3	-3.47	120.05	126.04
29	a	406	CLA	C2C-C1C-NC	3.46	113.22	109.97
46	S	621	LUT	C35-C15-C14	-3.46	116.38	123.47
47	r	622	XAT	C20-C13-C14	-3.46	118.07	122.92
29	N	603	CLA	C1-C2-C3	-3.46	120.05	126.04
45	s	606	CHL	CHD-C1D-ND	-3.46	121.27	124.45
29	N	604	CLA	C1-C2-C3	-3.45	120.07	126.04
29	B	602	CLA	C2C-C1C-NC	3.45	113.21	109.97
31	C	514	BCR	C15-C14-C13	-3.45	122.38	127.31
31	d	404	BCR	C37-C22-C21	-3.45	118.09	122.92
31	D	404	BCR	C23-C22-C21	3.45	124.24	118.94
45	S	606	CHL	CHD-C1D-ND	-3.45	121.28	124.45

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
29	C	511	CLA	C1-C2-C3	-3.44	120.09	126.04
29	c	508	CLA	C1-C2-C3	-3.44	120.09	126.04
44	h	101	RRX	C7-C8-C9	-3.44	121.03	126.23
29	s	605	CLA	C1-C2-C3	-3.44	121.19	126.75
29	c	503	CLA	CHD-C1D-ND	-3.44	121.30	124.45
29	b	605	CLA	CMA-C3A-C4A	3.44	121.01	111.77
46	G	620	LUT	C35-C34-C33	-3.43	122.41	127.31
29	C	510	CLA	C1C-C2C-C3C	-3.43	103.35	106.96
47	R	621	XAT	C38-C25-C26	-3.43	116.51	122.26
29	g	610	CLA	C1-C2-C3	-3.43	120.11	126.04
29	B	615	CLA	CMB-C2B-C3B	3.43	131.09	124.68
46	S	620	LUT	C11-C10-C9	-3.43	122.42	127.31
29	S	614	CLA	CHD-C1D-ND	-3.43	121.31	124.45
29	b	610	CLA	C1-C2-C3	-3.43	120.12	126.04
29	a	406	CLA	CHD-C1D-ND	-3.42	121.31	124.45
29	c	511	CLA	CHD-C1D-ND	-3.42	121.31	124.45
29	c	510	CLA	C1-C2-C3	-3.42	120.12	126.04
29	c	501	CLA	C1-C2-C3	-3.42	120.12	126.04
29	C	507	CLA	C2C-C1C-NC	3.42	113.18	109.97
29	d	402	CLA	C2D-C1D-ND	3.42	112.62	110.10
29	B	612	CLA	CMB-C2B-C1B	-3.42	123.21	128.46
29	C	503	CLA	CHD-C1D-ND	-3.42	121.31	124.45
29	B	603	CLA	C1-C2-C3	-3.41	120.14	126.04
29	c	510	CLA	C1C-C2C-C3C	-3.41	103.37	106.96
29	s	611	CLA	C2C-C1C-NC	3.41	113.17	109.97
29	R	610	CLA	CMA-C3A-C4A	3.41	120.95	111.77
29	N	613	CLA	CMA-C3A-C4A	3.41	120.95	111.77
29	b	617	CLA	C1-C2-C3	-3.41	120.14	126.04
29	Y	613	CLA	C1-C2-C3	-3.41	120.14	126.04
48	G	623	NEX	C17-C1-C6	-3.41	107.42	110.47
46	S	621	LUT	C22-C23-C24	-3.41	107.86	111.74
29	c	505	CLA	O2D-CGD-O1D	-3.41	117.17	123.84
29	G	610	CLA	C1-C2-C3	-3.41	120.15	126.04
46	r	620	LUT	C31-C32-C33	-3.41	116.85	126.42
31	b	619	BCR	C37-C22-C21	-3.41	118.15	122.92
46	s	621	LUT	C18-C5-C6	-3.40	120.71	124.53
29	c	505	CLA	C1-C2-C3	-3.40	120.16	126.04
29	c	510	CLA	CHD-C1D-ND	-3.40	121.33	124.45
29	s	604	CLA	CHD-C1D-ND	-3.40	121.33	124.45
46	n	620	LUT	C22-C23-C24	-3.40	107.88	111.74
29	C	513	CLA	CHD-C1D-ND	-3.40	121.33	124.45
47	n	622	XAT	O24-C25-C24	3.40	115.93	113.38

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
29	C	510	CLA	C1-C2-C3	-3.39	120.18	126.04
29	C	506	CLA	CHD-C1D-ND	-3.39	121.34	124.45
48	Y	623	NEX	C39-C29-C30	-3.39	118.17	122.92
46	g	620	LUT	C11-C10-C9	-3.39	122.47	127.31
29	N	612	CLA	CMA-C3A-C4A	3.39	120.89	111.77
29	c	509	CLA	C2C-C1C-NC	3.39	113.15	109.97
29	b	603	CLA	C1-C2-C3	-3.39	120.18	126.04
46	n	620	LUT	C15-C14-C13	-3.39	122.48	127.31
47	r	622	XAT	O24-C25-C24	3.38	115.92	113.38
29	n	613	CLA	CMA-C3A-C4A	3.38	120.86	111.77
29	b	606	CLA	C1-C2-C3	-3.38	120.20	126.04
46	Y	621	LUT	C11-C10-C9	-3.38	122.49	127.31
29	b	616	CLA	C2C-C1C-NC	3.38	113.14	109.97
29	c	504	CLA	CHD-C1D-ND	-3.38	121.35	124.45
29	S	605	CLA	C2C-C1C-NC	3.38	113.13	109.97
29	C	512	CLA	C2D-C1D-ND	3.37	112.59	110.10
45	y	606	CHL	CHD-C1D-ND	-3.37	121.35	124.45
29	B	606	CLA	C2C-C1C-NC	3.37	113.13	109.97
29	B	609	CLA	C2C-C1C-NC	3.37	113.13	109.97
45	G	601	CHL	C2C-C3C-C4C	3.37	108.89	106.49
45	Y	606	CHL	CHD-C1D-ND	-3.37	121.36	124.45
47	r	622	XAT	C27-C28-C29	3.37	130.76	125.53
46	y	621	LUT	C15-C14-C13	-3.37	122.50	127.31
45	g	601	CHL	C3C-C4C-NC	-3.37	106.80	110.57
47	r	622	XAT	C26-C27-C28	-3.37	118.88	125.99
29	b	603	CLA	CHD-C1D-ND	-3.36	121.36	124.45
29	b	609	CLA	C2C-C1C-NC	3.36	113.12	109.97
45	S	601	CHL	CHD-C1D-ND	-3.36	121.36	124.45
29	C	504	CLA	CHD-C1D-ND	-3.36	121.37	124.45
45	y	606	CHL	C3C-C4C-NC	-3.36	106.81	110.57
29	b	605	CLA	C1-C2-C3	-3.36	120.24	126.04
45	G	607	CHL	CHD-C1D-ND	-3.35	121.37	124.45
29	r	610	CLA	C2D-C1D-ND	3.35	112.58	110.10
29	n	603	CLA	CHD-C1D-ND	-3.35	121.38	124.45
29	A	407	CLA	C2C-C1C-NC	3.35	113.11	109.97
29	y	613	CLA	CMA-C3A-C4A	3.35	120.76	111.77
47	y	622	XAT	O4-C5-C4	-3.34	110.87	113.38
29	r	608	CLA	C2C-C1C-NC	3.34	113.10	109.97
29	B	615	CLA	CBA-CAA-C2A	3.34	123.72	113.86
48	r	623	NEX	C2-C1-C6	3.34	112.45	109.21
29	y	603	CLA	C2D-C1D-ND	3.34	112.56	110.10
45	y	607	CHL	CHD-C1D-ND	-3.33	121.39	124.45

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
45	G	601	CHL	C3C-C4C-NC	-3.33	106.83	110.57
29	g	610	CLA	CHD-C1D-ND	-3.33	121.39	124.45
31	B	618	BCR	C23-C24-C25	-3.33	117.85	127.20
29	s	605	CLA	O2D-CGD-O1D	-3.33	117.33	123.84
44	h	101	RRX	C1-C6-C5	-3.33	117.93	122.61
29	b	609	CLA	CMA-C3A-C4A	3.33	120.71	111.77
29	R	610	CLA	C1-C2-C3	-3.32	120.30	126.04
29	A	405	CLA	CHD-C1D-ND	-3.32	121.40	124.45
29	c	505	CLA	C2C-C1C-NC	3.32	113.08	109.97
46	G	620	LUT	C31-C30-C29	-3.32	122.58	127.31
29	D	403	CLA	C1-C2-C3	-3.31	120.31	126.04
29	b	607	CLA	CHD-C1D-ND	-3.31	121.41	124.45
29	S	605	CLA	O2D-CGD-O1D	-3.31	117.36	123.84
45	s	607	CHL	C2C-C3C-C4C	3.31	108.85	106.49
47	R	621	XAT	C27-C28-C29	3.31	130.67	125.53
29	a	407	CLA	C2C-C1C-NC	3.31	113.07	109.97
29	B	612	CLA	CHD-C1D-ND	-3.31	121.41	124.45
29	G	612	CLA	C2C-C1C-NC	3.31	113.07	109.97
29	B	616	CLA	C2C-C1C-NC	3.31	113.07	109.97
29	c	503	CLA	CMA-C3A-C4A	3.31	120.66	111.77
29	C	502	CLA	C2C-C1C-NC	3.30	113.07	109.97
45	N	606	CHL	CHD-C1D-ND	-3.30	121.42	124.45
45	g	607	CHL	CHD-C1D-ND	-3.30	121.42	124.45
47	G	622	XAT	C38-C25-C26	-3.30	116.72	122.26
29	B	606	CLA	C1-C2-C3	-3.30	120.33	126.04
29	S	603	CLA	C2C-C1C-NC	3.30	113.06	109.97
29	s	602	CLA	CHD-C1D-ND	-3.30	121.42	124.45
44	H	101	RRX	C16-C17-C18	-3.30	122.60	127.31
29	g	603	CLA	C1-C2-C3	-3.30	120.33	126.04
45	G	606	CHL	CHD-C1D-ND	-3.30	121.42	124.45
29	S	605	CLA	C1-O2A-CGA	3.30	125.10	116.44
45	g	601	CHL	C2C-C3C-C4C	3.30	108.84	106.49
44	h	101	RRX	C20-C21-C22	-3.30	122.61	127.31
29	S	613	CLA	C2C-C1C-NC	3.29	113.06	109.97
45	s	607	CHL	CMA-C3A-C4A	3.29	120.62	111.77
45	y	605	CHL	C3C-C4C-NC	-3.29	106.88	110.57
29	a	406	CLA	C1-C2-C3	-3.29	120.35	126.04
29	y	608	CLA	C1-C2-C3	-3.29	121.44	126.75
45	Y	605	CHL	C3C-C4C-NC	-3.28	106.89	110.57
29	B	607	CLA	CHD-C1D-ND	-3.28	121.44	124.45
29	n	612	CLA	C2C-C1C-NC	3.28	113.05	109.97
29	G	610	CLA	CHD-C1D-ND	-3.28	121.44	124.45

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
29	N	602	CLA	C2C-C1C-NC	3.28	113.04	109.97
29	r	612	CLA	C2C-C1C-NC	3.27	113.04	109.97
45	G	606	CHL	CMA-C3A-C4A	3.27	120.57	111.77
45	g	606	CHL	C2C-C3C-C4C	3.27	108.82	106.49
45	r	607	CHL	C2C-C3C-C4C	3.27	108.82	106.49
29	C	505	CLA	C2C-C1C-NC	3.27	113.03	109.97
29	G	603	CLA	C2C-C1C-NC	3.27	113.03	109.97
45	N	609	CHL	C3C-C4C-NC	-3.27	106.91	110.57
47	Y	622	XAT	C6-C7-C8	-3.27	119.08	125.99
45	R	607	CHL	C3C-C4C-NC	-3.27	106.91	110.57
29	C	501	CLA	C1-C2-C3	-3.27	120.39	126.04
46	s	621	LUT	C22-C23-C24	-3.26	108.03	111.74
29	r	613	CLA	CHD-C1D-ND	-3.26	121.45	124.45
29	y	610	CLA	C1-C2-C3	-3.26	120.40	126.04
29	s	604	CLA	C2C-C1C-NC	3.26	113.03	109.97
29	c	506	CLA	O2D-CGD-O1D	-3.26	117.46	123.84
45	N	601	CHL	CHD-C1D-ND	-3.26	121.46	124.45
29	s	602	CLA	C2D-C1D-ND	3.26	112.51	110.10
45	S	607	CHL	C2C-C3C-C4C	3.26	108.81	106.49
29	Y	611	CLA	C2C-C1C-NC	3.26	113.02	109.97
31	C	516	BCR	C34-C9-C10	-3.26	118.36	122.92
45	S	607	CHL	CMA-C3A-C4A	3.25	120.52	111.77
35	b	620	C7Z	C2-C3-C4	3.25	114.76	110.30
29	G	611	CLA	C2C-C1C-NC	3.25	113.02	109.97
29	b	606	CLA	C2C-C1C-NC	3.25	113.02	109.97
29	b	609	CLA	C2D-C1D-ND	3.25	112.50	110.10
29	R	612	CLA	C2C-C1C-NC	3.25	113.02	109.97
29	G	614	CLA	CHD-C1D-ND	-3.25	121.47	124.45
29	c	507	CLA	C2C-C1C-NC	3.25	113.01	109.97
45	n	609	CHL	CHD-C1D-ND	-3.25	121.47	124.45
29	y	604	CLA	CHD-C1D-ND	-3.25	121.47	124.45
46	G	621	LUT	C18-C5-C6	-3.25	120.88	124.53
45	n	601	CHL	C2C-C3C-C4C	3.24	108.80	106.49
29	R	610	CLA	CHD-C1D-ND	-3.24	121.47	124.45
29	A	405	CLA	CMB-C2B-C3B	3.24	130.75	124.68
29	s	617	CLA	CHD-C1D-ND	-3.24	121.47	124.45
29	g	611	CLA	C2C-C1C-NC	3.24	113.01	109.97
29	y	610	CLA	CHD-C1D-ND	-3.24	121.47	124.45
46	s	620	LUT	C15-C14-C13	-3.24	122.69	127.31
45	r	607	CHL	C3C-C4C-NC	-3.24	106.94	110.57
30	A	408	PHO	CMB-C2B-C3B	3.24	130.74	124.68
29	S	612	CLA	C2C-C1C-NC	3.24	113.00	109.97

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
29	b	615	CLA	CMB-C2B-C3B	3.23	130.73	124.68
29	C	503	CLA	CMA-C3A-C4A	3.23	120.47	111.77
29	b	615	CLA	C1-C2-C3	-3.23	120.45	126.04
31	C	514	BCR	C33-C5-C6	-3.23	120.90	124.53
46	G	620	LUT	C18-C5-C6	-3.23	120.90	124.53
45	R	607	CHL	C2C-C3C-C4C	3.23	108.79	106.49
29	B	611	CLA	C2C-C1C-NC	3.23	113.00	109.97
29	D	402	CLA	C2D-C1D-ND	3.23	112.48	110.10
47	g	622	XAT	C7-C8-C9	-3.23	120.52	125.53
29	a	405	CLA	CHD-C1D-ND	-3.23	121.49	124.45
29	b	611	CLA	C2C-C1C-NC	3.23	112.99	109.97
29	a	410	CLA	CHD-C1D-ND	-3.22	121.49	124.45
29	b	614	CLA	C2C-C1C-NC	3.22	112.99	109.97
29	s	610	CLA	CHD-C1D-ND	-3.22	121.49	124.45
29	S	611	CLA	C2C-C1C-NC	3.22	112.99	109.97
29	g	613	CLA	CHD-C1D-ND	-3.22	121.50	124.45
31	c	514	BCR	C15-C14-C13	-3.22	122.72	127.31
29	c	511	CLA	C1-C2-C3	-3.22	120.48	126.04
29	S	604	CLA	C1-C2-C3	-3.22	120.48	126.04
29	s	612	CLA	C2C-C1C-NC	3.22	112.98	109.97
45	N	607	CHL	CHD-C1D-ND	-3.22	121.50	124.45
43	f	101	HEM	C1B-NB-C4B	3.22	108.39	105.07
29	R	608	CLA	C2C-C1C-NC	3.22	112.98	109.97
29	N	603	CLA	CHD-C1D-ND	-3.21	121.50	124.45
45	g	606	CHL	CHD-C1D-ND	-3.21	121.50	124.45
29	n	603	CLA	C2D-C1D-ND	3.21	112.47	110.10
29	n	610	CLA	CHD-C1D-ND	-3.21	121.50	124.45
29	Y	608	CLA	C1-C2-C3	-3.21	121.56	126.75
29	B	609	CLA	O2D-CGD-O1D	-3.21	117.56	123.84
29	c	508	CLA	CMB-C2B-C1B	-3.21	123.53	128.46
29	B	612	CLA	CMB-C2B-C3B	3.21	130.68	124.68
29	a	405	CLA	CMB-C2B-C3B	3.21	130.68	124.68
29	b	609	CLA	C1-C2-C3	-3.21	120.50	126.04
29	S	617	CLA	CMA-C3A-C4A	3.21	120.39	111.77
29	c	501	CLA	CHD-C1D-ND	-3.21	121.51	124.45
29	c	508	CLA	CHD-C1D-ND	-3.20	121.51	124.45
29	g	614	CLA	CHD-C1D-ND	-3.20	121.51	124.45
29	B	614	CLA	C2D-C1D-ND	3.20	112.46	110.10
45	n	609	CHL	C1-O2A-CGA	3.20	124.85	116.44
45	n	601	CHL	C3C-C4C-NC	-3.20	106.98	110.57
48	G	623	NEX	C39-C29-C30	-3.20	118.44	122.92
29	A	406	CLA	C1-C2-C3	-3.20	120.51	126.04

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
29	B	614	CLA	C2C-C1C-NC	3.20	112.97	109.97
45	n	607	CHL	CHB-C4A-NA	3.20	128.93	124.51
29	c	504	CLA	C1-C2-C3	-3.20	120.51	126.04
42	D	405	PL9	C7-C3-C2	-3.20	119.10	123.30
47	g	622	XAT	C38-C25-C26	-3.20	116.90	122.26
46	R	620	LUT	C31-C32-C33	-3.19	117.44	126.42
29	Y	614	CLA	C2C-C1C-NC	3.19	112.96	109.97
29	n	610	CLA	C1-C2-C3	-3.19	120.53	126.04
29	s	605	CLA	C2C-C1C-NC	3.19	112.96	109.97
29	B	615	CLA	CHD-C1D-ND	-3.19	121.52	124.45
29	Y	614	CLA	CHD-C1D-ND	-3.19	121.52	124.45
29	y	603	CLA	CHD-C1D-ND	-3.19	121.52	124.45
45	Y	609	CHL	CMA-C3A-C4A	3.19	120.34	111.77
29	b	602	CLA	C2C-C1C-NC	3.19	112.96	109.97
29	g	614	CLA	C2C-C1C-NC	3.19	112.96	109.97
29	Y	604	CLA	CHD-C1D-ND	-3.19	121.53	124.45
29	b	609	CLA	O2D-CGD-O1D	-3.18	117.61	123.84
48	R	622	NEX	C17-C1-C6	-3.18	107.62	110.47
46	s	620	LUT	C18-C5-C6	-3.18	120.95	124.53
46	n	620	LUT	C35-C34-C33	-3.18	122.77	127.31
29	g	603	CLA	CHD-C1D-ND	-3.18	121.53	124.45
29	y	613	CLA	CHD-C1D-ND	-3.18	121.53	124.45
45	G	607	CHL	C3C-C4C-NC	-3.18	107.01	110.57
29	n	614	CLA	CHD-C1D-ND	-3.18	121.53	124.45
48	G	623	NEX	C31-C30-C29	3.18	131.84	127.31
45	y	607	CHL	C2C-C3C-C4C	3.18	108.75	106.49
44	h	101	RRX	C4-C5-C6	-3.18	118.12	122.73
45	g	605	CHL	C2C-C3C-C4C	3.17	108.75	106.49
29	y	614	CLA	C2C-C1C-NC	3.17	112.95	109.97
29	r	611	CLA	C2C-C1C-NC	3.17	112.94	109.97
45	Y	609	CHL	C3C-C4C-NC	-3.17	107.01	110.57
29	B	615	CLA	C1-C2-C3	-3.17	120.56	126.04
29	y	602	CLA	CHD-C1D-ND	-3.17	121.54	124.45
43	F	101	HEM	C4B-CHC-C1C	3.17	126.74	122.56
29	N	603	CLA	C2C-C1C-NC	3.17	112.94	109.97
29	b	611	CLA	C1-C2-C3	-3.17	120.56	126.04
45	N	606	CHL	C2C-C3C-C4C	3.17	108.75	106.49
29	R	604	CLA	CHD-C1D-ND	-3.17	121.54	124.45
29	C	506	CLA	O2D-CGD-O1D	-3.17	117.64	123.84
45	G	606	CHL	C3C-C4C-NC	-3.17	107.02	110.57
31	C	516	BCR	C23-C24-C25	-3.17	118.31	127.20
46	g	621	LUT	C18-C5-C6	-3.17	120.97	124.53

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
29	b	602	CLA	CHD-C1D-ND	-3.17	121.55	124.45
29	s	613	CLA	C2C-C1C-NC	3.16	112.94	109.97
29	y	611	CLA	C2C-C1C-NC	3.16	112.94	109.97
29	r	603	CLA	CMA-C3A-C4A	3.16	120.28	111.77
29	A	410	CLA	CHD-C1D-ND	-3.16	121.55	124.45
29	C	501	CLA	CHD-C1D-ND	-3.16	121.55	124.45
29	R	613	CLA	C2C-C1C-NC	3.16	112.94	109.97
31	a	411	BCR	C23-C24-C25	-3.16	118.32	127.20
31	c	514	BCR	C33-C5-C6	-3.16	120.98	124.53
29	s	617	CLA	C2C-C1C-NC	3.16	112.93	109.97
41	d	401	BCT	O2-C-O1	-3.16	111.34	119.55
45	s	607	CHL	C3C-C4C-NC	-3.16	107.03	110.57
29	Y	610	CLA	CHD-C1D-ND	-3.16	121.55	124.45
44	H	101	RRX	C4-C5-C6	-3.16	118.14	122.73
45	s	608	CHL	C2C-C3C-C4C	3.16	108.74	106.49
45	N	608	CHL	CMA-C3A-C4A	3.16	120.26	111.77
45	N	609	CHL	CMA-C3A-C4A	3.16	120.26	111.77
29	c	501	CLA	O2D-CGD-O1D	-3.16	117.66	123.84
29	C	510	CLA	C2D-C1D-ND	3.16	112.43	110.10
29	n	611	CLA	C2C-C1C-NC	3.16	112.93	109.97
29	N	614	CLA	CHD-C1D-ND	-3.16	121.55	124.45
29	y	614	CLA	CHD-C1D-ND	-3.16	121.55	124.45
44	H	101	RRX	C20-C21-C22	-3.16	122.81	127.31
47	Y	622	XAT	O4-C5-C4	-3.16	111.01	113.38
29	n	602	CLA	CHD-C1D-ND	-3.16	121.55	124.45
45	r	607	CHL	CHD-C1D-ND	-3.16	121.55	124.45
46	G	621	LUT	C15-C14-C13	-3.16	122.81	127.31
29	S	609	CLA	CHD-C1D-ND	-3.15	121.56	124.45
47	n	622	XAT	C38-C25-C26	-3.15	116.98	122.26
29	g	602	CLA	C1-C2-C3	-3.15	120.59	126.04
29	C	511	CLA	C2C-C1C-NC	3.15	112.92	109.97
29	G	613	CLA	CHD-C1D-ND	-3.15	121.56	124.45
47	N	622	XAT	C38-C25-C26	-3.15	116.98	122.26
29	S	617	CLA	C2C-C1C-NC	3.15	112.92	109.97
45	G	608	CHL	C2C-C3C-C4C	3.15	108.73	106.49
29	g	612	CLA	C2C-C1C-NC	3.15	112.92	109.97
45	g	605	CHL	C3C-C4C-NC	-3.15	107.04	110.57
29	S	610	CLA	CHD-C1D-ND	-3.15	121.56	124.45
29	C	503	CLA	C2C-C1C-NC	3.15	112.92	109.97
48	y	623	NEX	C39-C29-C30	-3.15	118.52	122.92
31	B	619	BCR	C33-C5-C6	-3.15	121.00	124.53
29	c	510	CLA	C2C-C1C-NC	3.14	112.92	109.97

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
29	S	604	CLA	C2C-C1C-NC	3.14	112.92	109.97
29	S	605	CLA	C1-C2-C3	-3.14	121.67	126.75
29	s	603	CLA	C2C-C1C-NC	3.14	112.92	109.97
29	R	603	CLA	CMA-C3A-C4A	3.14	120.22	111.77
29	N	610	CLA	CHD-C1D-ND	-3.14	121.57	124.45
29	R	603	CLA	CHD-C1D-ND	-3.14	121.57	124.45
29	r	602	CLA	CHD-C1D-ND	-3.14	121.57	124.45
45	y	609	CHL	CMA-C3A-C4A	3.14	120.22	111.77
29	N	614	CLA	C2C-C1C-NC	3.14	112.91	109.97
29	B	611	CLA	CHD-C1D-ND	-3.14	121.57	124.45
29	c	502	CLA	C1-O2A-CGA	3.14	124.67	116.44
29	r	609	CLA	C1-C2-C3	-3.14	120.62	126.04
29	G	602	CLA	CHD-C1D-ND	-3.14	121.57	124.45
48	g	623	NEX	C39-C29-C30	-3.14	118.53	122.92
41	D	401	BCT	O2-C-O1	-3.13	111.42	119.55
29	r	603	CLA	C2C-C1C-NC	3.13	112.91	109.97
29	c	508	CLA	CMB-C2B-C3B	3.13	130.54	124.68
29	b	615	CLA	CBA-CAA-C2A	3.13	123.11	113.86
29	Y	608	CLA	CHD-C1D-ND	-3.13	121.58	124.45
29	n	603	CLA	C1-C2-C3	-3.13	120.63	126.04
29	C	509	CLA	C1-C2-C3	-3.13	120.63	126.04
29	R	609	CLA	CHD-C1D-ND	-3.13	121.58	124.45
29	b	614	CLA	C2D-C1D-ND	3.13	112.41	110.10
29	r	602	CLA	C2C-C1C-NC	3.13	112.90	109.97
45	n	605	CHL	C3C-C4C-NC	-3.13	107.06	110.57
44	h	101	RRX	C16-C17-C18	-3.13	122.85	127.31
29	C	509	CLA	CHD-C1D-ND	-3.13	121.58	124.45
29	n	604	CLA	CHD-C1D-ND	-3.13	121.58	124.45
29	C	502	CLA	C2D-C1D-ND	3.13	112.41	110.10
46	S	620	LUT	C35-C15-C14	-3.12	117.07	123.47
29	a	410	CLA	C2C-C1C-NC	3.12	112.90	109.97
29	B	611	CLA	C1-C2-C3	-3.12	120.64	126.04
29	b	604	CLA	C2C-C1C-NC	3.12	112.90	109.97
48	S	622	NEX	C39-C29-C30	-3.12	118.55	122.92
29	N	611	CLA	C2C-C1C-NC	3.12	112.90	109.97
45	g	606	CHL	CMA-C3A-C4A	3.12	120.16	111.77
45	g	607	CHL	C3C-C4C-NC	-3.12	107.07	110.57
29	y	612	CLA	C2C-C1C-NC	3.12	112.89	109.97
45	G	607	CHL	CHB-C4A-NA	3.12	128.82	124.51
45	y	605	CHL	C2C-C3C-C4C	3.12	108.71	106.49
45	y	609	CHL	C3C-C4C-NC	-3.12	107.07	110.57
43	F	101	HEM	C3B-C2B-C1B	3.12	108.80	106.49

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
29	A	406	CLA	CHD-C1D-ND	-3.12	121.59	124.45
29	R	613	CLA	CHD-C1D-ND	-3.12	121.59	124.45
45	s	601	CHL	C3C-C4C-NC	-3.12	107.08	110.57
29	Y	613	CLA	CMA-C3A-C4A	3.12	120.15	111.77
46	y	621	LUT	C35-C15-C14	-3.12	117.09	123.47
29	R	611	CLA	C2C-C1C-NC	3.12	112.89	109.97
45	g	608	CHL	C2C-C3C-C4C	3.12	108.71	106.49
29	b	611	CLA	CHD-C1D-ND	-3.12	121.59	124.45
29	b	610	CLA	C2D-C1D-ND	3.11	112.40	110.10
31	C	515	BCR	C33-C5-C6	-3.11	121.03	124.53
45	N	605	CHL	C3C-C4C-NC	-3.11	107.08	110.57
46	Y	621	LUT	C18-C5-C6	-3.11	121.03	124.53
29	b	603	CLA	C2C-C1C-NC	3.11	112.89	109.97
29	n	603	CLA	C1C-C2C-C3C	-3.11	103.69	106.96
29	Y	612	CLA	C2C-C1C-NC	3.11	112.89	109.97
29	A	410	CLA	C1-C2-C3	-3.11	120.66	126.04
31	C	517	BCR	C28-C27-C26	-3.11	108.52	114.08
45	n	608	CHL	CMA-C3A-C4A	3.11	120.13	111.77
29	r	603	CLA	CHD-C1D-ND	-3.11	121.60	124.45
43	f	101	HEM	C3B-C2B-C1B	3.11	108.79	106.49
29	G	604	CLA	C2C-C1C-NC	3.11	112.89	109.97
29	R	602	CLA	CHD-C1D-ND	-3.11	121.60	124.45
29	g	603	CLA	C2C-C1C-NC	3.11	112.88	109.97
29	Y	613	CLA	CHD-C1D-ND	-3.11	121.60	124.45
31	b	619	BCR	C33-C5-C6	-3.11	121.04	124.53
29	C	508	CLA	CMB-C2B-C1B	-3.10	123.69	128.46
29	r	610	CLA	CHD-C1D-ND	-3.10	121.60	124.45
29	N	613	CLA	C2C-C1C-NC	3.10	112.88	109.97
29	b	614	CLA	CHD-C1D-ND	-3.10	121.60	124.45
45	Y	606	CHL	C2C-C3C-C4C	3.10	108.70	106.49
29	c	503	CLA	C2C-C1C-NC	3.10	112.88	109.97
29	B	614	CLA	CHD-C1D-ND	-3.10	121.61	124.45
45	S	608	CHL	C2C-C3C-C4C	3.10	108.70	106.49
29	Y	603	CLA	CHD-C1D-ND	-3.10	121.61	124.45
46	S	621	LUT	C7-C8-C9	-3.10	121.55	126.23
45	n	609	CHL	C3C-C4C-NC	-3.10	107.10	110.57
29	S	609	CLA	C2C-C1C-NC	3.10	112.87	109.97
29	C	510	CLA	C2C-C1C-NC	3.09	112.87	109.97
46	s	620	LUT	C38-C25-C24	-3.09	116.94	123.56
29	G	603	CLA	CHD-C1D-ND	-3.09	121.61	124.45
29	b	615	CLA	C2C-C1C-NC	3.09	112.87	109.97
45	s	608	CHL	C3C-C4C-NC	-3.09	107.10	110.57

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
29	d	403	CLA	C1-C2-C3	-3.09	120.69	126.04
45	S	607	CHL	C3C-C4C-NC	-3.09	107.10	110.57
45	g	605	CHL	CMA-C3A-C4A	3.09	120.08	111.77
29	R	612	CLA	CHD-C1D-ND	-3.09	121.61	124.45
45	N	601	CHL	C2C-C3C-C4C	3.09	108.69	106.49
45	Y	607	CHL	CHB-C4A-NA	3.09	128.78	124.51
45	R	606	CHL	CMA-C3A-C4A	3.09	120.07	111.77
45	N	608	CHL	C3C-C4C-NC	-3.09	107.11	110.57
39	L	101	LHG	O7-C7-C8	3.09	118.15	111.50
29	b	610	CLA	CHD-C1D-ND	-3.08	121.62	124.45
29	Y	603	CLA	C2D-C1D-ND	3.08	112.38	110.10
29	B	609	CLA	C1-C2-C3	-3.08	120.71	126.04
29	G	613	CLA	C2C-C1C-NC	3.08	112.86	109.97
29	N	612	CLA	C2C-C1C-NC	3.08	112.86	109.97
29	y	608	CLA	CHD-C1D-ND	-3.08	121.62	124.45
45	S	608	CHL	C3C-C4C-NC	-3.08	107.12	110.57
45	n	608	CHL	C3C-C4C-NC	-3.08	107.12	110.57
29	Y	602	CLA	CHD-C1D-ND	-3.08	121.62	124.45
29	r	604	CLA	CHD-C1D-ND	-3.08	121.62	124.45
45	G	608	CHL	C3C-C4C-NC	-3.08	107.12	110.57
29	b	614	CLA	C1-C2-C3	-3.08	120.72	126.04
29	B	609	CLA	CMA-C3A-C4A	3.08	120.04	111.77
45	G	605	CHL	CMA-C3A-C4A	3.08	120.04	111.77
29	s	617	CLA	CMA-C3A-C4A	3.07	120.04	111.77
29	N	610	CLA	O2A-CGA-CBA	3.07	121.55	111.91
46	g	620	LUT	C18-C5-C6	-3.07	121.08	124.53
29	B	615	CLA	O2D-CGD-O1D	-3.07	117.83	123.84
29	C	501	CLA	CMA-C3A-C4A	3.07	120.03	111.77
29	n	614	CLA	C2C-C1C-NC	3.07	112.85	109.97
44	h	101	RRX	C23-C24-C25	-3.07	118.58	127.20
39	l	101	LHG	O7-C7-C8	3.07	118.12	111.50
29	B	610	CLA	CHD-C1D-ND	-3.07	121.63	124.45
29	d	402	CLA	CHD-C1D-ND	-3.07	121.63	124.45
29	S	603	CLA	C1-C2-C3	-3.07	120.73	126.04
29	B	617	CLA	C2C-C1C-NC	3.07	112.85	109.97
29	A	407	CLA	CMA-C3A-C4A	3.07	120.02	111.77
29	C	508	CLA	C2C-C1C-NC	3.07	112.85	109.97
45	Y	609	CHL	C2C-C3C-C4C	3.07	108.67	106.49
29	S	602	CLA	C1-C2-C3	-3.07	120.74	126.04
45	n	607	CHL	C1-C2-C3	-3.07	120.74	126.04
29	G	602	CLA	CMA-C3A-C4A	3.07	120.01	111.77
45	g	609	CHL	C4D-CHA-C1A	3.06	124.98	121.25

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
45	n	607	CHL	C1-O2A-CGA	3.06	124.48	116.44
29	g	613	CLA	C2C-C1C-NC	3.06	112.84	109.97
45	N	607	CHL	CHB-C4A-NA	3.06	128.75	124.51
29	n	603	CLA	C2C-C1C-NC	3.06	112.84	109.97
29	B	614	CLA	C1-C2-C3	-3.06	120.75	126.04
45	G	605	CHL	C2C-C3C-C4C	3.06	108.67	106.49
47	Y	622	XAT	C7-C8-C9	-3.06	120.78	125.53
42	d	405	PL9	C7-C3-C2	-3.06	119.28	123.30
29	s	602	CLA	C1-C2-C3	-3.06	120.75	126.04
29	b	615	CLA	CHD-C1D-ND	-3.06	121.64	124.45
29	g	613	CLA	C1-C2-C3	-3.06	120.76	126.04
29	G	613	CLA	C1-C2-C3	-3.06	120.76	126.04
29	c	503	CLA	C1-C2-C3	-3.06	120.76	126.04
31	c	515	BCR	C33-C5-C6	-3.05	121.10	124.53
29	B	615	CLA	C2C-C1C-NC	3.05	112.83	109.97
29	B	615	CLA	CMB-C2B-C1B	-3.05	123.77	128.46
29	G	614	CLA	C2C-C1C-NC	3.05	112.83	109.97
29	b	615	CLA	O2D-CGD-O1D	-3.05	117.87	123.84
29	b	608	CLA	C2C-C1C-NC	3.05	112.83	109.97
29	R	612	CLA	C1-C2-C3	-3.05	120.77	126.04
29	n	604	CLA	C2C-C1C-NC	3.05	112.83	109.97
45	S	601	CHL	C3C-C4C-NC	-3.05	107.15	110.57
29	S	610	CLA	C1-C2-C3	-3.05	120.77	126.04
29	c	507	CLA	CHD-C1D-ND	-3.05	121.65	124.45
45	Y	606	CHL	C3C-C4C-NC	-3.05	107.15	110.57
45	G	606	CHL	C2C-C3C-C4C	3.05	108.66	106.49
42	d	405	PL9	C7-C8-C9	-3.05	121.72	126.79
29	C	508	CLA	CHD-C1D-ND	-3.05	121.65	124.45
29	c	509	CLA	C1-C2-C3	-3.05	120.77	126.04
45	g	608	CHL	CMA-C3A-C4A	3.05	119.96	111.77
29	N	614	CLA	CMA-C3A-C4A	3.04	119.95	111.77
46	Y	620	LUT	C18-C5-C6	-3.04	121.11	124.53
29	b	612	CLA	CMB-C2B-C1B	-3.04	123.79	128.46
29	G	614	CLA	CMA-C3A-C4A	3.04	119.95	111.77
45	g	608	CHL	C3C-C4C-NC	-3.04	107.16	110.57
29	a	407	CLA	CMA-C3A-C4A	3.04	119.95	111.77
29	D	403	CLA	CHD-C1D-ND	-3.04	121.66	124.45
29	c	513	CLA	CHD-C1D-ND	-3.04	121.66	124.45
29	G	602	CLA	C1-C2-C3	-3.04	120.78	126.04
29	N	604	CLA	C2D-C1D-ND	3.04	112.34	110.10
29	Y	610	CLA	C1-C2-C3	-3.04	120.79	126.04
45	G	607	CHL	C2C-C3C-C4C	3.04	108.66	106.49

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
29	b	607	CLA	CAA-C2A-C3A	-3.04	104.46	112.78
29	G	603	CLA	C2D-C1D-ND	3.04	112.34	110.10
29	A	410	CLA	CMA-C3A-C4A	3.04	119.94	111.77
47	R	621	XAT	O4-C5-C4	-3.04	111.10	113.38
46	G	620	LUT	C22-C23-C24	-3.04	108.28	111.74
45	N	609	CHL	C2C-C3C-C4C	3.04	108.65	106.49
29	N	604	CLA	CHD-C1D-ND	-3.04	121.66	124.45
45	N	601	CHL	C3C-C4C-NC	-3.04	107.17	110.57
29	r	611	CLA	CMA-C3A-C4A	3.04	119.93	111.77
29	s	610	CLA	C1-O2A-CGA	3.04	124.41	116.44
29	R	603	CLA	C2C-C1C-NC	3.03	112.81	109.97
29	Y	613	CLA	C2C-C1C-NC	3.03	112.81	109.97
29	c	511	CLA	C2C-C1C-NC	3.03	112.81	109.97
45	N	605	CHL	C2C-C3C-C4C	3.03	108.65	106.49
29	a	405	CLA	C2D-C1D-ND	3.03	112.34	110.10
29	a	407	CLA	CHD-C1D-ND	-3.03	121.67	124.45
29	A	406	CLA	C1C-C2C-C3C	-3.03	103.77	106.96
45	G	601	CHL	CMA-C3A-C4A	3.03	119.92	111.77
29	c	509	CLA	CHD-C1D-ND	-3.03	121.67	124.45
45	y	605	CHL	CMA-C3A-C4A	3.03	119.92	111.77
29	r	608	CLA	CHD-C1D-ND	-3.03	121.67	124.45
29	g	604	CLA	C2C-C1C-NC	3.03	112.81	109.97
45	y	607	CHL	C3C-C4C-NC	-3.03	107.17	110.57
45	Y	605	CHL	CMA-C3A-C4A	3.03	119.92	111.77
29	C	509	CLA	C1C-C2C-C3C	-3.03	103.77	106.96
29	y	613	CLA	C2D-C1D-ND	3.03	112.34	110.10
29	R	608	CLA	CHD-C1D-ND	-3.03	121.67	124.45
29	C	508	CLA	CMB-C2B-C3B	3.03	130.34	124.68
29	A	410	CLA	C2C-C1C-NC	3.03	112.81	109.97
43	F	101	HEM	C1B-NB-C4B	3.03	108.20	105.07
33	d	411	LMG	O8-C28-C29	3.02	121.40	111.91
29	y	613	CLA	C1-C2-C3	-3.02	120.81	126.04
45	n	609	CHL	CMA-C3A-C4A	3.02	119.90	111.77
29	B	608	CLA	C2C-C1C-NC	3.02	112.81	109.97
29	s	614	CLA	C2C-C1C-NC	3.02	112.81	109.97
46	s	621	LUT	C35-C15-C14	-3.02	117.28	123.47
45	S	608	CHL	CMA-C3A-C4A	3.02	119.89	111.77
45	N	607	CHL	C3C-C4C-NC	-3.02	107.18	110.57
29	g	604	CLA	CHD-C1D-ND	-3.02	121.68	124.45
30	a	409	PHO	O2D-CGD-O1D	-3.02	117.94	123.84
29	S	604	CLA	C1C-C2C-C3C	-3.02	103.79	106.96
45	g	609	CHL	C1B-CHB-C4A	-3.02	124.14	130.12

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
29	G	611	CLA	CHD-C1D-ND	-3.02	121.68	124.45
45	R	607	CHL	CMA-C3A-C4A	3.01	119.88	111.77
48	y	623	NEX	C17-C1-C6	-3.01	107.77	110.47
29	r	612	CLA	CHD-C1D-ND	-3.01	121.69	124.45
29	r	610	CLA	C1-C2-C3	-3.01	120.83	126.04
31	d	404	BCR	C23-C22-C21	3.01	123.56	118.94
29	B	607	CLA	CMA-C3A-C4A	3.01	119.87	111.77
45	N	605	CHL	CMA-C3A-C4A	3.01	119.87	111.77
29	S	614	CLA	C2C-C1C-NC	3.01	112.79	109.97
29	R	602	CLA	C2D-C1D-ND	3.01	112.32	110.10
29	y	603	CLA	C1C-C2C-C3C	-3.01	103.79	106.96
29	S	611	CLA	CHD-C1D-ND	-3.01	121.69	124.45
29	a	406	CLA	C1C-C2C-C3C	-3.01	103.79	106.96
29	b	614	CLA	C1C-C2C-C3C	-3.01	103.79	106.96
29	c	504	CLA	C2D-C1D-ND	3.01	112.32	110.10
29	G	603	CLA	C1C-C2C-C3C	-3.01	103.79	106.96
45	n	607	CHL	C3C-C4C-NC	-3.01	107.20	110.57
47	y	622	XAT	C6-C7-C8	-3.01	119.63	125.99
29	n	614	CLA	CMA-C3A-C4A	3.01	119.86	111.77
47	Y	622	XAT	C38-C25-C26	-3.01	117.22	122.26
44	H	101	RRX	C1-C6-C5	-3.01	118.38	122.61
44	H	101	RRX	C23-C24-C25	-3.01	118.76	127.20
29	B	607	CLA	CAA-C2A-C3A	-3.01	104.55	112.78
45	n	609	CHL	C2C-C3C-C4C	3.00	108.63	106.49
48	g	623	NEX	C20-C13-C14	-3.00	118.71	122.92
29	Y	610	CLA	C2D-C1D-ND	3.00	112.32	110.10
29	B	604	CLA	C2C-C1C-NC	3.00	112.79	109.97
29	Y	603	CLA	C2C-C1C-NC	3.00	112.79	109.97
45	r	606	CHL	CMA-C3A-C4A	3.00	119.84	111.77
29	s	613	CLA	C2D-C1D-ND	3.00	112.32	110.10
29	b	612	CLA	C2C-C1C-NC	3.00	112.78	109.97
29	s	609	CLA	C2C-C1C-NC	3.00	112.78	109.97
45	N	608	CHL	C2C-C3C-C4C	3.00	108.63	106.49
29	b	607	CLA	O2A-CGA-CBA	3.00	121.32	111.91
29	c	502	CLA	CHD-C1D-ND	-3.00	121.70	124.45
29	B	607	CLA	O2A-CGA-CBA	3.00	121.32	111.91
45	s	601	CHL	C2C-C3C-C4C	3.00	108.63	106.49
45	n	607	CHL	CHD-C1D-ND	-3.00	121.70	124.45
29	D	403	CLA	C2C-C1C-NC	3.00	112.78	109.97
44	h	101	RRX	C33-C5-C6	-3.00	121.16	124.53
29	B	610	CLA	C2D-C1D-ND	3.00	112.31	110.10
45	n	606	CHL	C3C-C4C-NC	-3.00	107.21	110.57

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
29	N	602	CLA	CHD-C1D-ND	-2.99	121.70	124.45
45	g	607	CHL	C2C-C3C-C4C	2.99	108.62	106.49
29	s	611	CLA	CHD-C1D-ND	-2.99	121.70	124.45
29	R	604	CLA	C2D-C1D-ND	2.99	112.31	110.10
29	R	610	CLA	CAA-C2A-C3A	-2.99	104.59	112.78
44	H	101	RRX	C38-C26-C27	2.99	119.90	114.36
29	C	512	CLA	C2C-C1C-NC	2.99	112.77	109.97
29	N	611	CLA	CMA-C3A-C4A	2.99	119.81	111.77
29	B	614	CLA	C1C-C2C-C3C	-2.99	103.81	106.96
47	r	622	XAT	C11-C10-C9	-2.99	123.05	127.31
29	S	617	CLA	CHD-C1D-ND	-2.99	121.71	124.45
44	H	101	RRX	C21-C20-C19	-2.99	113.90	123.22
29	N	613	CLA	O2A-CGA-CBA	2.98	121.28	111.91
45	g	606	CHL	C3C-C4C-NC	-2.98	107.22	110.57
29	g	610	CLA	C2C-C1C-NC	2.98	112.77	109.97
45	s	608	CHL	CMA-C3A-C4A	2.98	119.79	111.77
29	r	604	CLA	C2D-C1D-ND	2.98	112.30	110.10
29	c	501	CLA	CMA-C3A-C4A	2.98	119.78	111.77
29	G	604	CLA	CHD-C1D-ND	-2.98	121.72	124.45
45	N	607	CHL	CMA-C3A-C4A	2.98	119.78	111.77
45	G	605	CHL	C3C-C4C-NC	-2.98	107.23	110.57
29	n	611	CLA	CMA-C3A-C4A	2.98	119.78	111.77
29	s	612	CLA	CHD-C1D-ND	-2.98	121.72	124.45
29	S	614	CLA	C1-C2-C3	-2.98	120.89	126.04
45	S	606	CHL	CMA-C3A-C4A	2.98	119.78	111.77
29	n	612	CLA	CHD-C1D-ND	-2.98	121.72	124.45
48	n	623	NEX	C39-C29-C30	-2.98	118.75	122.92
29	y	608	CLA	C2C-C1C-NC	2.98	112.76	109.97
29	s	609	CLA	CMA-C3A-C4A	2.98	119.77	111.77
29	R	611	CLA	CMA-C3A-C4A	2.97	119.77	111.77
46	y	620	LUT	C10-C11-C12	-2.97	113.94	123.22
48	R	622	NEX	C27-C28-C29	-2.97	120.92	125.53
45	y	606	CHL	CMA-C3A-C4A	2.97	119.75	111.77
29	s	612	CLA	C2D-C1D-ND	2.97	112.29	110.10
45	N	607	CHL	C2C-C3C-C4C	2.97	108.61	106.49
29	y	604	CLA	C2C-C1C-NC	2.97	112.75	109.97
29	N	610	CLA	CMA-C3A-C4A	2.97	119.75	111.77
29	B	613	CLA	CHD-C1D-ND	-2.97	121.73	124.45
29	A	410	CLA	C2D-C1D-ND	2.96	112.29	110.10
29	C	502	CLA	CMA-C3A-C4A	2.96	119.74	111.77
29	y	602	CLA	C2D-C1D-ND	2.96	112.29	110.10
29	C	507	CLA	CHD-C1D-ND	-2.96	121.73	124.45

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
45	g	609	CHL	C1-O2A-CGA	2.96	124.21	116.44
29	n	613	CLA	C2C-C1C-NC	2.96	112.75	109.97
30	a	408	PHO	CMB-C2B-C3B	2.96	130.22	124.68
29	G	610	CLA	C2D-C1D-ND	2.96	112.29	110.10
29	S	612	CLA	C2D-C1D-ND	2.96	112.29	110.10
45	N	606	CHL	C3C-C4C-NC	-2.96	107.25	110.57
29	B	609	CLA	C1C-C2C-C3C	-2.96	103.84	106.96
45	n	607	CHL	C2C-C3C-C4C	2.96	108.60	106.49
45	G	608	CHL	CMA-C3A-C4A	2.96	119.73	111.77
29	r	609	CLA	C2C-C1C-NC	2.96	112.74	109.97
45	Y	601	CHL	CMA-C3A-C4A	2.96	119.72	111.77
48	r	623	NEX	C27-C28-C29	-2.96	120.94	125.53
45	s	606	CHL	C3C-C4C-NC	-2.96	107.25	110.57
29	d	403	CLA	CHD-C1D-ND	-2.96	121.74	124.45
29	s	609	CLA	CHD-C1D-ND	-2.96	121.74	124.45
45	g	607	CHL	CHB-C4A-NA	2.96	128.60	124.51
43	F	101	HEM	C4C-CHD-C1D	2.95	126.45	122.56
29	r	609	CLA	CHD-C1D-ND	-2.95	121.74	124.45
29	N	603	CLA	C1C-C2C-C3C	-2.95	103.85	106.96
29	n	602	CLA	CMA-C3A-C4A	2.95	119.70	111.77
29	a	410	CLA	C1-C2-C3	-2.95	120.94	126.04
45	n	606	CHL	C2C-C3C-C4C	2.95	108.59	106.49
29	d	403	CLA	C2C-C1C-NC	2.95	112.73	109.97
29	y	602	CLA	C1-C2-C3	-2.95	120.94	126.04
45	R	606	CHL	CHD-C1D-ND	-2.95	121.75	124.45
29	R	602	CLA	O2D-CGD-O1D	-2.95	118.08	123.84
29	R	611	CLA	CHD-C1D-ND	-2.95	121.75	124.45
29	B	608	CLA	O2D-CGD-O1D	-2.95	118.08	123.84
46	y	620	LUT	C7-C8-C9	-2.94	121.78	126.23
29	S	605	CLA	O2A-CGA-CBA	2.94	121.15	111.91
29	S	613	CLA	C2D-C1D-ND	2.94	112.27	110.10
45	n	608	CHL	C2C-C3C-C4C	2.94	108.59	106.49
29	S	605	CLA	CMA-C3A-C4A	2.94	119.69	111.77
48	S	622	NEX	C31-C30-C29	2.94	131.51	127.31
31	A	411	BCR	C23-C24-C25	-2.94	118.94	127.20
39	d	408	LHG	O8-C23-C24	2.94	121.14	111.91
29	B	603	CLA	C2C-C1C-NC	2.94	112.73	109.97
29	y	603	CLA	C2C-C1C-NC	2.94	112.73	109.97
45	r	607	CHL	CMA-C3A-C4A	2.94	119.68	111.77
29	a	410	CLA	CMA-C3A-C4A	2.94	119.67	111.77
45	n	601	CHL	CMA-C3A-C4A	2.94	119.67	111.77
45	n	605	CHL	C2C-C3C-C4C	2.94	108.58	106.49

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
48	R	622	NEX	C26-C27-C28	-2.94	119.78	125.99
48	r	623	NEX	C26-C27-C28	-2.94	119.78	125.99
45	g	601	CHL	CMA-C3A-C4A	2.94	119.67	111.77
47	g	622	XAT	O4-C5-C4	-2.94	111.17	113.38
29	c	504	CLA	C2C-C1C-NC	2.94	112.72	109.97
29	s	603	CLA	C1-C2-C3	-2.94	120.97	126.04
29	Y	604	CLA	C2C-C1C-NC	2.94	112.72	109.97
29	c	512	CLA	C2C-C1C-NC	2.94	112.72	109.97
29	c	502	CLA	CMA-C3A-C4A	2.93	119.66	111.77
46	r	620	LUT	C2-C3-C4	-2.93	106.29	110.30
29	s	605	CLA	CMA-C3A-C4A	2.93	119.65	111.77
29	r	611	CLA	CHD-C1D-ND	-2.93	121.76	124.45
45	n	606	CHL	CMA-C3A-C4A	2.93	119.65	111.77
44	h	101	RRX	C21-C20-C19	-2.93	114.07	123.22
29	n	614	CLA	C2D-C1D-ND	2.93	112.26	110.10
29	g	603	CLA	C2D-C1D-ND	2.93	112.26	110.10
29	N	611	CLA	CHD-C1D-ND	-2.93	121.76	124.45
29	S	609	CLA	CMA-C3A-C4A	2.93	119.64	111.77
29	n	611	CLA	CHD-C1D-ND	-2.93	121.76	124.45
29	A	407	CLA	CHD-C1D-ND	-2.93	121.76	124.45
29	g	614	CLA	CMA-C3A-C4A	2.93	119.64	111.77
31	b	619	BCR	C4-C5-C6	-2.92	118.48	122.73
29	S	613	CLA	CMA-C3A-C4A	2.92	119.63	111.77
35	b	620	C7Z	C8-C7-C6	-2.92	118.99	127.20
29	g	611	CLA	C2D-C1D-ND	2.92	112.26	110.10
46	Y	621	LUT	C15-C14-C13	-2.92	123.14	127.31
45	N	606	CHL	CMA-C3A-C4A	2.92	119.63	111.77
29	B	602	CLA	CHD-C1D-ND	-2.92	121.77	124.45
29	n	610	CLA	C2D-C1D-ND	2.92	112.26	110.10
29	n	612	CLA	C2D-C1D-ND	2.92	112.26	110.10
29	D	403	CLA	CMA-C3A-C4A	2.92	119.62	111.77
29	c	501	CLA	C2C-C1C-NC	2.92	112.71	109.97
29	b	612	CLA	CMB-C2B-C3B	2.92	130.14	124.68
45	g	609	CHL	CMA-C3A-C4A	2.92	119.62	111.77
29	g	602	CLA	CHD-C1D-ND	-2.92	121.77	124.45
29	R	602	CLA	C2C-C1C-NC	2.92	112.71	109.97
46	S	620	LUT	C38-C25-C24	-2.92	117.31	123.56
29	A	407	CLA	C2D-C1D-ND	2.92	112.25	110.10
29	B	610	CLA	C2C-C1C-NC	2.92	112.70	109.97
29	R	610	CLA	C2D-C1D-ND	2.92	112.25	110.10
29	b	613	CLA	CHD-C1D-ND	-2.92	121.78	124.45
29	C	504	CLA	C2D-C1D-ND	2.91	112.25	110.10

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
29	C	501	CLA	O2D-CGD-O1D	-2.91	118.14	123.84
29	G	610	CLA	C2C-C1C-NC	2.91	112.70	109.97
29	B	608	CLA	CHD-C1D-ND	-2.91	121.78	124.45
45	g	607	CHL	CMA-C3A-C4A	2.91	119.60	111.77
29	Y	608	CLA	CMA-C3A-C4A	2.91	119.60	111.77
29	s	613	CLA	CMA-C3A-C4A	2.91	119.60	111.77
29	Y	612	CLA	C2D-C1D-ND	2.91	112.25	110.10
29	S	602	CLA	C2C-C1C-NC	2.91	112.70	109.97
29	Y	603	CLA	C1C-C2C-C3C	-2.91	103.89	106.96
29	g	603	CLA	CMA-C3A-C4A	2.91	119.60	111.77
45	n	609	CHL	CHB-C4A-NA	2.91	128.54	124.51
29	s	610	CLA	CAA-C2A-C3A	-2.91	104.81	112.78
29	s	602	CLA	C1C-C2C-C3C	-2.91	103.90	106.96
29	y	613	CLA	C2C-C1C-NC	2.91	112.70	109.97
29	G	610	CLA	CAA-C2A-C3A	-2.91	104.81	112.78
31	d	404	BCR	C34-C9-C10	-2.91	118.85	122.92
29	y	611	CLA	C1-C2-C3	-2.91	121.02	126.04
29	G	612	CLA	CMA-C3A-C4A	2.91	119.58	111.77
47	R	621	XAT	C20-C13-C14	-2.91	118.85	122.92
29	C	501	CLA	C2C-C1C-NC	2.90	112.69	109.97
29	N	602	CLA	C1-C2-C3	-2.90	121.02	126.04
29	S	602	CLA	C2D-C1D-ND	2.90	112.24	110.10
29	y	610	CLA	C2D-C1D-ND	2.90	112.24	110.10
29	N	610	CLA	C2C-C1C-NC	2.90	112.69	109.97
29	Y	608	CLA	C2C-C1C-NC	2.90	112.69	109.97
29	C	503	CLA	C1-C2-C3	-2.90	121.02	126.04
29	S	617	CLA	C2D-C1D-ND	2.90	112.24	110.10
39	D	408	LHG	O8-C23-C24	2.90	121.00	111.91
29	C	503	CLA	C2D-C1D-ND	2.90	112.24	110.10
48	n	623	NEX	C17-C1-C6	-2.90	107.88	110.47
29	G	611	CLA	C2D-C1D-ND	2.90	112.24	110.10
29	c	513	CLA	C2D-C1D-ND	2.90	112.24	110.10
29	G	611	CLA	CMA-C3A-C4A	2.90	119.56	111.77
29	S	612	CLA	CHD-C1D-ND	-2.90	121.79	124.45
29	s	602	CLA	C2C-C1C-NC	2.90	112.69	109.97
29	b	602	CLA	C1-C2-C3	-2.89	121.04	126.04
29	b	608	CLA	O2D-CGD-O1D	-2.89	118.18	123.84
45	S	606	CHL	C3C-C4C-NC	-2.89	107.33	110.57
29	B	612	CLA	CAA-C2A-C3A	-2.89	104.86	112.78
29	R	609	CLA	C2C-C1C-NC	2.89	112.68	109.97
29	n	610	CLA	C2C-C1C-NC	2.89	112.68	109.97
29	B	616	CLA	C2D-C1D-ND	2.89	112.23	110.10

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
29	D	402	CLA	CHD-C1D-ND	-2.89	121.80	124.45
29	b	608	CLA	CHD-C1D-ND	-2.89	121.80	124.45
29	a	407	CLA	C2D-C1D-ND	2.89	112.23	110.10
29	s	617	CLA	C2D-C1D-ND	2.89	112.23	110.10
29	A	405	CLA	CMB-C2B-C1B	-2.89	124.02	128.46
48	Y	623	NEX	C31-C30-C29	2.89	131.43	127.31
29	y	614	CLA	CMA-C3A-C4A	2.89	119.53	111.77
45	n	607	CHL	CMA-C3A-C4A	2.89	119.53	111.77
29	Y	614	CLA	CMA-C3A-C4A	2.88	119.52	111.77
45	Y	607	CHL	C3C-C4C-NC	-2.88	107.34	110.57
29	N	602	CLA	CMA-C3A-C4A	2.88	119.52	111.77
29	C	502	CLA	CHD-C1D-ND	-2.88	121.81	124.45
29	N	604	CLA	C2C-C1C-NC	2.88	112.67	109.97
35	b	620	C7Z	C22-C23-C24	2.88	114.25	110.30
29	n	611	CLA	O2D-CGD-O1D	-2.88	118.21	123.84
47	G	622	XAT	C26-C27-C28	-2.88	119.91	125.99
29	C	504	CLA	C2C-C1C-NC	2.88	112.67	109.97
29	n	603	CLA	O2D-CGD-O1D	-2.88	118.21	123.84
29	b	607	CLA	CMA-C3A-C4A	2.88	119.50	111.77
29	s	609	CLA	C2D-C1D-ND	2.88	112.22	110.10
29	d	403	CLA	CMA-C3A-C4A	2.88	119.50	111.77
45	S	601	CHL	CMA-C3A-C4A	2.88	119.50	111.77
45	y	601	CHL	CMA-C3A-C4A	2.88	119.50	111.77
45	y	607	CHL	CMA-C3A-C4A	2.87	119.50	111.77
29	S	602	CLA	CHD-C1D-ND	-2.87	121.81	124.45
29	g	611	CLA	CHD-C1D-ND	-2.87	121.81	124.45
29	C	507	CLA	C1C-C2C-C3C	-2.87	103.93	106.96
29	c	508	CLA	C2C-C1C-NC	2.87	112.67	109.97
29	N	603	CLA	C2D-C1D-ND	2.87	112.22	110.10
29	N	613	CLA	C2D-C1D-ND	2.87	112.22	110.10
33	D	411	LMG	O8-C28-C29	2.87	120.92	111.91
29	C	510	CLA	C1D-ND-C4D	-2.87	104.30	106.33
29	s	611	CLA	C1C-C2C-C3C	-2.87	103.94	106.96
29	B	602	CLA	C1-C2-C3	-2.87	121.08	126.04
29	a	410	CLA	C2D-C1D-ND	2.87	112.22	110.10
45	N	607	CHL	C1-O2A-CGA	2.87	123.97	116.44
29	B	617	CLA	C1-C2-C3	-2.87	121.08	126.04
29	B	612	CLA	C2C-C1C-NC	2.87	112.66	109.97
38	C	523	DGD	O1G-C1A-C2A	2.87	120.91	111.91
29	G	603	CLA	C1-C2-C3	-2.87	121.08	126.04
46	Y	621	LUT	C35-C15-C14	-2.87	117.60	123.47
45	Y	607	CHL	C2C-C3C-C4C	2.87	108.53	106.49

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
29	r	613	CLA	C2C-C1C-NC	2.87	112.66	109.97
29	N	603	CLA	CMA-C3A-C4A	2.87	119.48	111.77
29	n	602	CLA	C1-C2-C3	-2.87	121.08	126.04
29	n	602	CLA	C2C-C1C-NC	2.87	112.66	109.97
29	G	612	CLA	C2D-C1D-ND	2.87	112.22	110.10
29	C	513	CLA	C1-C2-C3	-2.86	121.09	126.04
29	B	611	CLA	C2D-C1D-ND	2.86	112.22	110.10
33	A	413	LMG	C8-O7-C10	-2.86	110.74	117.79
29	b	616	CLA	CHD-C1D-ND	-2.86	121.82	124.45
29	a	405	CLA	CMB-C2B-C1B	-2.86	124.06	128.46
38	c	518	DGD	O1G-C1A-C2A	2.86	120.89	111.91
29	n	603	CLA	CAA-C2A-C3A	-2.86	104.94	112.78
46	S	621	LUT	C18-C5-C6	-2.86	121.31	124.53
31	B	619	BCR	C4-C5-C6	-2.86	118.58	122.73
29	r	610	CLA	C1D-ND-C4D	-2.86	104.30	106.33
35	b	620	C7Z	C31-C30-C29	-2.86	123.23	127.31
29	C	512	CLA	CMB-C2B-C3B	2.86	130.03	124.68
29	s	603	CLA	C2D-C1D-ND	2.86	112.21	110.10
47	g	622	XAT	C6-C7-C8	-2.86	119.95	125.99
29	R	609	CLA	C1-C2-C3	-2.86	121.10	126.04
29	g	602	CLA	C2D-C1D-ND	2.86	112.21	110.10
45	y	601	CHL	CHB-C4A-NA	2.86	128.46	124.51
29	r	608	CLA	C1-C2-C3	-2.86	121.11	126.04
29	a	405	CLA	CAA-C2A-C3A	-2.85	104.96	112.78
31	B	619	BCR	C36-C18-C17	-2.85	118.92	122.92
47	N	622	XAT	C6-C7-C8	-2.85	119.96	125.99
30	A	409	PHO	O1D-CGD-CBD	2.85	129.48	124.74
29	y	612	CLA	C2D-C1D-ND	2.85	112.20	110.10
29	a	407	CLA	C1C-C2C-C3C	-2.85	103.96	106.96
29	B	607	CLA	C2C-C1C-NC	2.85	112.64	109.97
42	D	405	PL9	C7-C8-C9	-2.85	122.05	126.79
45	Y	605	CHL	C2C-C3C-C4C	2.85	108.52	106.49
29	y	610	CLA	C2C-C1C-NC	2.85	112.64	109.97
29	c	509	CLA	C1C-C2C-C3C	-2.85	103.97	106.96
29	b	609	CLA	C1C-C2C-C3C	-2.84	103.97	106.96
29	g	603	CLA	C1C-C2C-C3C	-2.84	103.97	106.96
29	B	608	CLA	C2D-C1D-ND	2.84	112.20	110.10
29	Y	602	CLA	CMA-C3A-C4A	2.84	119.40	111.77
29	b	608	CLA	C2D-C1D-ND	2.84	112.19	110.10
29	S	610	CLA	C2D-C1D-ND	2.84	112.19	110.10
29	d	402	CLA	C1D-ND-C4D	-2.83	104.32	106.33
31	c	515	BCR	C23-C24-C25	-2.83	119.25	127.20

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
29	S	609	CLA	C2D-C1D-ND	2.83	112.19	110.10
29	A	407	CLA	C1C-C2C-C3C	-2.83	103.98	106.96
29	b	605	CLA	C1C-C2C-C3C	-2.83	103.98	106.96
29	c	507	CLA	C1C-C2C-C3C	-2.83	103.98	106.96
29	G	603	CLA	CMA-C3A-C4A	2.83	119.38	111.77
38	c	523	DGD	O1G-C1A-C2A	2.83	120.79	111.91
29	G	614	CLA	C2D-C1D-ND	2.83	112.19	110.10
47	y	622	XAT	C38-C25-C26	-2.83	117.52	122.26
29	r	610	CLA	OBD-CAD-C3D	-2.83	121.72	128.52
29	N	613	CLA	C1-C2-C3	-2.83	121.15	126.04
29	Y	614	CLA	C1-C2-C3	-2.83	121.16	126.04
44	h	101	RRX	C15-C16-C17	-2.83	117.69	123.47
29	g	611	CLA	CMA-C3A-C4A	2.82	119.36	111.77
29	B	606	CLA	C1C-C2C-C3C	-2.82	103.99	106.96
29	b	602	CLA	CMA-C3A-C4A	2.82	119.36	111.77
46	y	621	LUT	C38-C25-C24	-2.82	117.52	123.56
29	N	614	CLA	C2D-C1D-ND	2.82	112.18	110.10
29	B	615	CLA	C1-O2A-CGA	2.82	123.85	116.44
48	N	623	NEX	C39-C29-C30	-2.82	118.97	122.92
29	n	604	CLA	C2D-C1D-ND	2.82	112.18	110.10
31	b	618	BCR	C23-C24-C25	-2.82	119.28	127.20
29	B	602	CLA	CMA-C3A-C4A	2.82	119.35	111.77
29	r	602	CLA	C2D-C1D-ND	2.82	112.18	110.10
29	c	512	CLA	CMB-C2B-C3B	2.82	129.95	124.68
44	H	101	RRX	C15-C16-C17	-2.82	117.70	123.47
38	C	518	DGD	O1G-C1A-C2A	2.82	120.75	111.91
30	A	409	PHO	O2D-CGD-O1D	-2.82	118.33	123.84
29	b	607	CLA	C2C-C1C-NC	2.82	112.61	109.97
29	S	603	CLA	C2D-C1D-ND	2.82	112.18	110.10
46	G	621	LUT	C11-C10-C9	-2.82	123.29	127.31
29	d	403	CLA	C2D-C1D-ND	2.81	112.18	110.10
29	C	513	CLA	C2D-C1D-ND	2.81	112.18	110.10
48	G	623	NEX	C38-C25-C26	-2.81	117.55	122.26
29	n	602	CLA	C2D-C1D-ND	2.81	112.18	110.10
29	g	612	CLA	C2D-C1D-ND	2.81	112.18	110.10
29	b	612	CLA	C1-C2-C3	-2.81	121.18	126.04
29	r	608	CLA	CMA-C3A-C4A	2.81	119.33	111.77
46	y	620	LUT	C38-C25-C24	-2.81	117.55	123.56
45	s	601	CHL	CMA-C3A-C4A	2.81	119.32	111.77
47	R	621	XAT	C31-C30-C29	-2.81	123.30	127.31
46	g	620	LUT	C31-C30-C29	-2.81	123.31	127.31
29	Y	602	CLA	C2D-C1D-ND	2.81	112.17	110.10

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
29	r	604	CLA	C2C-C1C-NC	2.81	112.60	109.97
43	f	101	HEM	C4C-CHD-C1D	2.81	126.26	122.56
45	y	601	CHL	C1-C2-C3	-2.80	121.19	126.04
29	G	612	CLA	CHD-C1D-ND	-2.80	121.88	124.45
48	s	623	NEX	C39-C29-C30	-2.80	119.00	122.92
29	r	610	CLA	CAA-C2A-C3A	-2.80	105.10	112.78
35	B	620	C7Z	C22-C23-C24	2.80	114.14	110.30
45	N	601	CHL	CMA-C3A-C4A	2.80	119.31	111.77
29	n	613	CLA	CHD-C1D-ND	-2.80	121.88	124.45
45	s	606	CHL	CMA-C3A-C4A	2.80	119.30	111.77
29	D	402	CLA	C1D-ND-C4D	-2.80	104.34	106.33
29	b	612	CLA	CHD-C1D-ND	-2.80	121.88	124.45
30	A	408	PHO	O2D-CGD-O1D	-2.80	118.36	123.84
29	n	610	CLA	CAA-C2A-C3A	-2.80	105.11	112.78
29	C	501	CLA	C2D-C1D-ND	2.80	112.17	110.10
29	D	403	CLA	C2D-C1D-ND	2.80	112.17	110.10
29	B	612	CLA	C1-C2-C3	-2.80	121.20	126.04
31	D	404	BCR	C34-C9-C10	-2.80	119.00	122.92
29	b	606	CLA	C1C-C2C-C3C	-2.80	104.02	106.96
29	N	610	CLA	C2D-C1D-ND	2.80	112.17	110.10
29	Y	613	CLA	C2D-C1D-ND	2.80	112.17	110.10
45	Y	607	CHL	C1-O2A-CGA	2.80	123.78	116.44
29	y	611	CLA	CHD-C1D-ND	-2.80	121.89	124.45
45	S	601	CHL	C2C-C3C-C4C	2.79	108.48	106.49
29	C	504	CLA	O2D-CGD-O1D	-2.79	118.38	123.84
35	B	620	C7Z	C8-C7-C6	-2.79	119.36	127.20
29	C	511	CLA	O2A-CGA-CBA	2.79	120.67	111.91
29	R	604	CLA	C2C-C1C-NC	2.79	112.59	109.97
45	Y	606	CHL	CMA-C3A-C4A	2.79	119.28	111.77
29	R	602	CLA	C1-C2-C3	-2.79	121.22	126.04
29	s	610	CLA	C2D-C1D-ND	2.79	112.16	110.10
46	Y	620	LUT	C38-C25-C24	-2.79	117.59	123.56
29	G	613	CLA	C2D-C1D-ND	2.79	112.16	110.10
29	R	611	CLA	C2D-C1D-ND	2.79	112.16	110.10
29	s	614	CLA	C1C-C2C-C3C	-2.79	104.02	106.96
29	b	604	CLA	CHD-C1D-ND	-2.79	121.89	124.45
29	r	613	CLA	C2D-C1D-ND	2.79	112.16	110.10
47	g	622	XAT	C26-C27-C28	-2.79	120.10	125.99
29	b	617	CLA	C2C-C1C-NC	2.79	112.58	109.97
29	S	614	CLA	C1C-C2C-C3C	-2.79	104.03	106.96
33	c	521	LMG	O8-C28-C29	2.79	120.65	111.91
46	g	621	LUT	C31-C30-C29	-2.79	123.33	127.31

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
45	G	609	CHL	C1-O2A-CGA	2.79	123.75	116.44
45	g	606	CHL	C1-O2A-CGA	2.78	123.75	116.44
29	c	509	CLA	O2D-CGD-O1D	-2.78	118.39	123.84
29	C	513	CLA	CMA-C3A-C4A	2.78	119.25	111.77
29	B	617	CLA	O2A-CGA-CBA	2.78	120.64	111.91
29	A	405	CLA	C2D-C1D-ND	2.78	112.16	110.10
29	N	613	CLA	CHD-C1D-ND	-2.78	121.90	124.45
29	c	504	CLA	O2D-CGD-O1D	-2.78	118.40	123.84
29	y	604	CLA	C1-C2-C3	-2.78	121.23	126.04
29	B	617	CLA	C2D-C1D-ND	2.78	112.15	110.10
29	c	511	CLA	O2A-CGA-CBA	2.78	120.63	111.91
29	y	608	CLA	CMA-C3A-C4A	2.78	119.24	111.77
29	g	613	CLA	C2D-C1D-ND	2.78	112.15	110.10
31	B	619	BCR	C23-C22-C21	2.78	123.20	118.94
29	r	602	CLA	C1-C2-C3	-2.78	121.24	126.04
31	c	514	BCR	C23-C24-C25	-2.78	119.40	127.20
45	y	609	CHL	C2C-C3C-C4C	2.78	108.47	106.49
29	a	410	CLA	C1C-C2C-C3C	-2.78	104.04	106.96
47	G	622	XAT	C6-C7-C8	-2.77	120.13	125.99
47	G	622	XAT	O24-C25-C24	2.77	115.47	113.38
45	G	609	CHL	C1B-CHB-C4A	-2.77	124.62	130.12
29	g	612	CLA	CMA-C3A-C4A	2.77	119.23	111.77
39	D	409	LHG	O8-C23-C24	2.77	120.61	111.91
29	G	604	CLA	C2D-C1D-ND	2.77	112.15	110.10
29	Y	611	CLA	C1-C2-C3	-2.77	121.25	126.04
29	n	613	CLA	O2A-CGA-CBA	2.77	120.61	111.91
49	s	625	LPX	O3-P1-O4	2.77	125.95	112.24
29	r	603	CLA	C1-C2-C3	-2.77	121.25	126.04
29	b	616	CLA	C2D-C1D-ND	2.77	112.15	110.10
45	y	601	CHL	C3C-C4C-NC	-2.77	107.47	110.57
31	c	515	BCR	C33-C5-C4	2.77	118.94	113.62
29	a	406	CLA	CMB-C2B-C3B	2.77	129.86	124.68
45	R	606	CHL	CHB-C4A-NA	2.77	128.34	124.51
29	B	605	CLA	C1C-C2C-C3C	-2.77	104.05	106.96
29	s	604	CLA	C1C-C2C-C3C	-2.77	104.05	106.96
49	S	625	LPX	O3-P1-O4	2.77	125.92	112.24
38	C	519	DGD	O1G-C1A-C2A	2.77	120.59	111.91
29	G	602	CLA	C2D-C1D-ND	2.77	112.14	110.10
29	c	507	CLA	O2D-CGD-O1D	-2.77	118.43	123.84
29	g	612	CLA	CHD-C1D-ND	-2.77	121.91	124.45
29	Y	611	CLA	C2D-C1D-ND	2.77	112.14	110.10
45	Y	607	CHL	CMA-C3A-C4A	2.77	119.20	111.77

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
29	R	602	CLA	CMA-C3A-C4A	2.76	119.20	111.77
31	c	516	BCR	C34-C9-C10	-2.76	119.05	122.92
29	s	611	CLA	C1-C2-C3	-2.76	121.27	126.04
39	d	410	LHG	O8-C23-C24	2.76	120.57	111.91
29	n	613	CLA	C2D-C1D-ND	2.76	112.14	110.10
29	y	608	CLA	C2D-C1D-ND	2.76	112.14	110.10
29	b	616	CLA	C1-C2-C3	-2.76	121.27	126.04
29	c	511	CLA	CMA-C3A-C4A	2.76	119.19	111.77
29	S	610	CLA	C2C-C1C-NC	2.76	112.56	109.97
29	s	617	CLA	C1C-C2C-C3C	-2.76	104.06	106.96
29	B	615	CLA	C1C-C2C-C3C	-2.75	104.06	106.96
29	C	512	CLA	CHD-C1D-ND	-2.75	121.92	124.45
29	y	612	CLA	CHD-C1D-ND	-2.75	121.92	124.45
29	B	613	CLA	O2D-CGD-O1D	-2.75	118.45	123.84
31	b	619	BCR	C28-C27-C26	-2.75	109.16	114.08
29	b	616	CLA	C1C-C2C-C3C	-2.75	104.06	106.96
29	Y	608	CLA	C2D-C1D-ND	2.75	112.13	110.10
29	b	611	CLA	C2D-C1D-ND	2.75	112.13	110.10
29	n	603	CLA	CMA-C3A-C4A	2.75	119.17	111.77
29	a	410	CLA	CAA-C2A-C3A	-2.75	105.24	112.78
29	b	603	CLA	C1C-C2C-C3C	-2.75	104.06	106.96
45	S	606	CHL	C2C-C3C-C4C	2.75	108.45	106.49
29	c	501	CLA	C2D-C1D-ND	2.75	112.13	110.10
46	n	620	LUT	C10-C11-C12	-2.75	114.64	123.22
29	c	512	CLA	C1-C2-C3	-2.75	121.29	126.04
29	s	614	CLA	C1-O2A-CGA	2.75	123.66	116.44
29	s	610	CLA	C2C-C1C-NC	2.75	112.55	109.97
29	B	602	CLA	C1C-C2C-C3C	-2.75	104.07	106.96
44	H	101	RRX	C33-C5-C6	-2.75	121.44	124.53
45	N	607	CHL	C1-C2-C3	-2.75	121.29	126.04
29	c	505	CLA	CHD-C1D-ND	-2.75	121.93	124.45
29	g	604	CLA	C2D-C1D-ND	2.75	112.13	110.10
46	Y	621	LUT	C38-C25-C24	-2.75	117.68	123.56
31	C	515	BCR	C33-C5-C4	2.75	118.89	113.62
42	d	405	PL9	C40-C39-C41	2.74	119.89	115.27
29	d	403	CLA	O2D-CGD-O1D	-2.74	118.47	123.84
29	b	615	CLA	C1-O2A-CGA	2.74	123.64	116.44
45	r	607	CHL	CHB-C4A-NA	2.74	128.31	124.51
45	y	607	CHL	CHB-C4A-NA	2.74	128.31	124.51
29	Y	614	CLA	C1C-C2C-C3C	-2.74	104.07	106.96
29	c	513	CLA	C2C-C1C-NC	2.74	112.54	109.97
46	Y	620	LUT	C15-C35-C34	-2.74	117.86	123.47

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
46	y	621	LUT	C18-C5-C6	-2.74	121.45	124.53
46	Y	620	LUT	C11-C10-C9	-2.74	123.40	127.31
29	g	602	CLA	CMA-C3A-C4A	2.74	119.14	111.77
35	B	620	C7Z	C31-C30-C29	-2.74	123.40	127.31
46	N	621	LUT	C11-C10-C9	-2.74	123.40	127.31
29	s	603	CLA	CMA-C3A-C4A	2.74	119.14	111.77
29	y	612	CLA	C1-C2-C3	-2.74	121.30	126.04
31	A	411	BCR	C38-C26-C25	-2.74	121.45	124.53
45	G	601	CHL	CHB-C4A-NA	2.74	128.30	124.51
29	R	613	CLA	C2D-C1D-ND	2.74	112.12	110.10
45	N	607	CHL	C4A-NA-C1A	2.74	107.94	106.71
29	A	406	CLA	CMA-C3A-C4A	2.74	119.13	111.77
29	c	513	CLA	C1-C2-C3	-2.74	121.31	126.04
29	N	602	CLA	O2A-CGA-CBA	2.74	120.49	111.91
46	Y	620	LUT	C10-C11-C12	-2.73	114.68	123.22
29	n	610	CLA	O2A-CGA-CBA	2.73	120.49	111.91
33	C	521	LMG	O8-C28-C29	2.73	120.49	111.91
46	Y	621	LUT	C31-C30-C29	-2.73	123.41	127.31
45	Y	601	CHL	CHB-C4A-NA	2.73	128.29	124.51
29	b	613	CLA	O2D-CGD-O1D	-2.73	118.50	123.84
29	a	410	CLA	O2D-CGD-O1D	-2.73	118.50	123.84
47	n	622	XAT	C6-C7-C8	-2.73	120.22	125.99
29	g	604	CLA	O2D-CGD-O1D	-2.73	118.50	123.84
29	R	608	CLA	C2D-C1D-ND	2.73	112.11	110.10
29	S	611	CLA	CMA-C3A-C4A	2.73	119.11	111.77
29	S	610	CLA	O2A-CGA-CBA	2.73	120.47	111.91
29	y	614	CLA	C2D-C1D-ND	2.73	112.11	110.10
45	Y	601	CHL	C1-C2-C3	-2.73	121.33	126.04
39	D	410	LHG	O8-C23-C24	2.73	120.46	111.91
29	N	611	CLA	C2D-C1D-ND	2.73	112.11	110.10
38	C	520	DGD	O1G-C1A-C2A	2.73	120.46	111.91
29	c	505	CLA	C1C-C2C-C3C	-2.73	104.09	106.96
29	S	605	CLA	C2D-C1D-ND	2.72	112.11	110.10
29	N	611	CLA	O2D-CGD-O1D	-2.72	118.51	123.84
29	B	609	CLA	C2D-C1D-ND	2.72	112.11	110.10
46	R	620	LUT	C2-C3-C4	-2.72	106.58	110.30
29	r	609	CLA	O2A-CGA-CBA	2.72	120.45	111.91
29	r	612	CLA	C2D-C1D-ND	2.72	112.11	110.10
29	B	604	CLA	CHD-C1D-ND	-2.72	121.95	124.45
31	A	411	BCR	C33-C5-C4	2.72	118.84	113.62
29	g	610	CLA	CMA-C3A-C4A	2.72	119.08	111.77
29	C	507	CLA	O2D-CGD-O1D	-2.72	118.52	123.84

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
31	b	619	BCR	C27-C26-C25	-2.72	118.78	122.73
45	Y	609	CHL	C1-O2A-CGA	2.72	123.57	116.44
45	s	606	CHL	C2C-C3C-C4C	2.72	108.43	106.49
29	A	410	CLA	CAA-C2A-C3A	-2.72	105.34	112.78
48	g	623	NEX	C31-C30-C29	2.72	131.19	127.31
29	B	616	CLA	CHD-C1D-ND	-2.72	121.96	124.45
48	s	623	NEX	C38-C25-C26	-2.72	117.71	122.26
29	b	615	CLA	CMB-C2B-C1B	-2.72	124.29	128.46
30	a	408	PHO	O2D-CGD-O1D	-2.71	118.53	123.84
29	C	506	CLA	C1C-C2C-C3C	-2.71	104.10	106.96
29	R	608	CLA	C1C-C2C-C3C	-2.71	104.10	106.96
29	A	410	CLA	O2D-CGD-O1D	-2.71	118.53	123.84
48	g	623	NEX	C38-C25-C26	-2.71	117.71	122.26
39	L	101	LHG	O8-C23-C24	2.71	120.41	111.91
29	N	604	CLA	C1C-C2C-C3C	-2.71	104.11	106.96
29	c	501	CLA	C1C-C2C-C3C	-2.71	104.11	106.96
46	s	621	LUT	C38-C25-C24	-2.71	117.76	123.56
31	B	618	BCR	C15-C14-C13	-2.71	123.44	127.31
29	g	610	CLA	C2D-C1D-ND	2.71	112.10	110.10
29	N	614	CLA	C1C-C2C-C3C	-2.71	104.11	106.96
29	c	506	CLA	C1C-C2C-C3C	-2.71	104.11	106.96
31	b	619	BCR	C36-C18-C17	-2.71	119.13	122.92
29	s	614	CLA	O2A-CGA-CBA	2.71	120.40	111.91
29	y	614	CLA	C1C-C2C-C3C	-2.71	104.11	106.96
29	B	616	CLA	O2D-CGD-O1D	-2.70	118.55	123.84
29	r	610	CLA	O2A-CGA-CBA	2.70	120.39	111.91
29	C	513	CLA	C2C-C1C-NC	2.70	112.50	109.97
29	b	615	CLA	C1C-C2C-C3C	-2.70	104.11	106.96
33	j	101	LMG	O8-C28-C29	2.70	120.39	111.91
29	y	611	CLA	O2D-CGD-O1D	-2.70	118.55	123.84
45	G	609	CHL	CMA-C3A-C4A	2.70	119.04	111.77
29	C	505	CLA	C1-C2-C3	-2.70	121.37	126.04
29	g	602	CLA	C2C-C1C-NC	2.70	112.50	109.97
29	N	602	CLA	C1C-C2C-C3C	-2.70	104.12	106.96
29	c	503	CLA	C2D-C1D-ND	2.70	112.09	110.10
29	B	615	CLA	OBD-CAD-C3D	-2.70	122.02	128.52
31	c	514	BCR	C36-C18-C17	-2.70	119.14	122.92
29	n	614	CLA	C1C-C2C-C3C	-2.70	104.12	106.96
39	S	624	LHG	O8-C23-C24	2.70	120.38	111.91
29	n	611	CLA	C2D-C1D-ND	2.70	112.09	110.10
29	b	602	CLA	C1C-C2C-C3C	-2.70	104.12	106.96
29	A	406	CLA	CMB-C2B-C3B	2.70	129.73	124.68

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
29	Y	602	CLA	C2C-C1C-NC	2.70	112.50	109.97
38	c	519	DGD	O1G-C1A-C2A	2.70	120.38	111.91
46	N	620	LUT	C7-C8-C9	-2.70	122.16	126.23
29	b	610	CLA	O2D-CGD-O1D	-2.70	118.57	123.84
29	C	511	CLA	C2D-C1D-ND	2.70	112.09	110.10
29	S	611	CLA	C1C-C2C-C3C	-2.70	104.12	106.96
29	b	605	CLA	O2D-CGD-O1D	-2.69	118.57	123.84
46	n	621	LUT	C38-C25-C24	-2.69	117.80	123.56
29	a	406	CLA	C2D-C1D-ND	2.69	112.09	110.10
46	N	620	LUT	C31-C30-C29	-2.69	123.47	127.31
29	D	403	CLA	O2D-CGD-O1D	-2.69	118.58	123.84
46	N	620	LUT	C10-C11-C12	-2.69	114.82	123.22
29	r	603	CLA	C2D-C1D-ND	2.69	112.09	110.10
29	S	604	CLA	C2D-C1D-ND	2.69	112.09	110.10
29	y	602	CLA	CMA-C3A-C4A	2.69	119.00	111.77
39	l	101	LHG	O8-C23-C24	2.69	120.35	111.91
29	G	602	CLA	C2C-C1C-NC	2.69	112.49	109.97
29	g	614	CLA	C2D-C1D-ND	2.69	112.08	110.10
45	G	609	CHL	C3C-C4C-NC	-2.68	107.56	110.57
29	R	603	CLA	C1C-C2C-C3C	-2.68	104.13	106.96
29	A	410	CLA	C1C-C2C-C3C	-2.68	104.14	106.96
29	y	614	CLA	C1-C2-C3	-2.68	121.40	126.04
29	S	611	CLA	C2D-C1D-ND	2.68	112.08	110.10
29	g	614	CLA	C1C-C2C-C3C	-2.68	104.14	106.96
48	n	623	NEX	C16-C1-C6	-2.68	108.07	110.47
44	h	101	RRX	C38-C26-C27	2.68	119.32	114.36
33	a	413	LMG	C8-O7-C10	-2.68	111.19	117.79
29	R	612	CLA	C1C-C2C-C3C	-2.68	104.14	106.96
29	n	612	CLA	C1C-C2C-C3C	-2.68	104.14	106.96
46	N	620	LUT	C11-C10-C9	-2.68	123.49	127.31
29	G	612	CLA	C1C-C2C-C3C	-2.68	104.14	106.96
29	s	611	CLA	CMA-C3A-C4A	2.68	118.97	111.77
29	S	613	CLA	C1C-C2C-C3C	-2.68	104.14	106.96
31	B	619	BCR	C28-C27-C26	-2.68	109.30	114.08
29	S	617	CLA	C1C-C2C-C3C	-2.68	104.14	106.96
38	C	519	DGD	C2G-O2G-C1B	-2.68	111.20	117.79
29	C	511	CLA	C1C-C2C-C3C	-2.68	104.14	106.96
29	n	602	CLA	O2A-CGA-CBA	2.67	120.30	111.91
30	a	408	PHO	O1D-CGD-CBD	2.67	129.19	124.74
29	G	613	CLA	C1C-C2C-C3C	-2.67	104.14	106.96
29	b	602	CLA	O2D-CGD-O1D	-2.67	118.61	123.84
29	b	617	CLA	O2A-CGA-CBA	2.67	120.30	111.91

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
29	B	605	CLA	CHA-C4D-ND	2.67	138.09	132.50
29	G	611	CLA	C1C-C2C-C3C	-2.67	104.15	106.96
45	g	601	CHL	CHB-C4A-NA	2.67	128.21	124.51
31	B	618	BCR	C3-C4-C5	-2.67	109.31	114.08
45	Y	601	CHL	C3C-C4C-NC	-2.67	107.58	110.57
29	c	511	CLA	C2D-C1D-ND	2.67	112.07	110.10
29	R	613	CLA	CMA-C3A-C4A	2.67	118.95	111.77
29	B	616	CLA	C1-C2-C3	-2.67	121.43	126.04
29	r	603	CLA	C1C-C2C-C3C	-2.67	104.15	106.96
47	G	622	XAT	O4-C5-C4	-2.67	111.38	113.38
29	G	614	CLA	C1C-C2C-C3C	-2.67	104.15	106.96
29	C	511	CLA	CMA-C3A-C4A	2.67	118.94	111.77
29	C	513	CLA	O2D-CGD-O1D	-2.67	118.62	123.84
29	r	613	CLA	O2D-CGD-O1D	-2.67	118.62	123.84
45	n	607	CHL	C4A-NA-C1A	2.67	107.91	106.71
29	B	613	CLA	C2D-C1D-ND	2.67	112.07	110.10
29	g	611	CLA	C1C-C2C-C3C	-2.67	104.15	106.96
29	Y	603	CLA	CMA-C3A-C4A	2.67	118.94	111.77
29	Y	602	CLA	O2A-CGA-CBA	2.67	120.27	111.91
29	R	612	CLA	C2D-C1D-ND	2.67	112.07	110.10
45	N	601	CHL	CHB-C4A-NA	2.66	128.20	124.51
29	b	611	CLA	O2D-CGD-O1D	-2.66	118.63	123.84
29	B	616	CLA	C1C-C2C-C3C	-2.66	104.16	106.96
29	S	602	CLA	C1C-C2C-C3C	-2.66	104.16	106.96
29	r	612	CLA	C1C-C2C-C3C	-2.66	104.16	106.96
29	Y	613	CLA	CMB-C2B-C3B	2.66	129.66	124.68
31	b	619	BCR	C23-C22-C21	2.66	123.03	118.94
29	B	611	CLA	C1C-C2C-C3C	-2.66	104.16	106.96
38	c	518	DGD	O6D-C5D-C6D	2.66	112.04	106.67
39	s	624	LHG	C5-O7-C7	-2.66	111.24	117.79
29	b	603	CLA	CMA-C3A-C4A	2.66	118.93	111.77
31	a	411	BCR	C35-C13-C12	2.66	122.27	118.08
39	c	625	LHG	O8-C23-C24	2.66	120.26	111.91
39	s	624	LHG	O8-C23-C24	2.66	120.26	111.91
29	C	505	CLA	CHD-C1D-ND	-2.66	122.01	124.45
29	S	612	CLA	C1C-C2C-C3C	-2.66	104.16	106.96
29	S	603	CLA	C1C-C2C-C3C	-2.66	104.16	106.96
29	c	511	CLA	C1C-C2C-C3C	-2.66	104.16	106.96
48	Y	623	NEX	C38-C25-C26	-2.66	117.81	122.26
29	S	609	CLA	C1C-C2C-C3C	-2.66	104.16	106.96
29	G	610	CLA	CMA-C3A-C4A	2.66	118.91	111.77
43	F	101	HEM	C4D-ND-C1D	2.66	107.82	105.07

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
35	b	620	C7Z	C27-C28-C29	-2.66	122.22	126.23
45	Y	607	CHL	CHD-C1D-ND	-2.66	122.01	124.45
31	c	517	BCR	C36-C18-C17	-2.65	119.20	122.92
36	C	524	DGA	OG1-CA1-CA2	2.65	120.24	111.91
29	C	513	CLA	CMB-C2B-C3B	2.65	129.65	124.68
31	C	517	BCR	C31-C1-C6	-2.65	105.99	110.30
47	R	621	XAT	C8-C9-C10	2.65	123.01	118.94
29	c	512	CLA	C1C-C2C-C3C	-2.65	104.17	106.96
29	s	605	CLA	O2A-CGA-CBA	2.65	120.22	111.91
29	s	613	CLA	O1D-CGD-CBD	-2.65	119.06	124.48
36	c	524	DGA	OG1-CA1-CA2	2.65	120.22	111.91
45	G	607	CHL	CMA-C3A-C4A	2.65	118.89	111.77
31	b	619	BCR	C1-C6-C5	-2.65	118.88	122.61
29	g	603	CLA	CAA-C2A-C3A	-2.65	105.53	112.78
45	s	606	CHL	C4D-CHA-C1A	2.65	124.47	121.25
35	b	620	C7Z	C28-C27-C26	-2.65	119.77	127.20
31	a	411	BCR	C38-C26-C25	-2.65	121.56	124.53
48	S	622	NEX	C38-C25-C26	-2.65	117.83	122.26
29	C	503	CLA	C1C-C2C-C3C	-2.65	104.17	106.96
29	Y	612	CLA	CHD-C1D-ND	-2.65	122.02	124.45
29	b	607	CLA	O2D-CGD-O1D	-2.65	118.67	123.84
29	b	614	CLA	O2D-CGD-O1D	-2.64	118.67	123.84
29	c	512	CLA	CHD-C1D-ND	-2.64	122.02	124.45
29	N	612	CLA	C2D-C1D-ND	2.64	112.05	110.10
29	R	609	CLA	O2A-CGA-CBA	2.64	120.20	111.91
31	C	517	BCR	C23-C24-C25	-2.64	119.78	127.20
29	B	602	CLA	CMD-C2D-C3D	-2.64	121.54	127.61
29	b	607	CLA	CMB-C2B-C3B	2.64	129.62	124.68
29	b	610	CLA	C2C-C1C-NC	2.64	112.45	109.97
29	y	602	CLA	O2A-CGA-CBA	2.64	120.20	111.91
29	B	602	CLA	O2D-CGD-O1D	-2.64	118.67	123.84
29	b	602	CLA	CMD-C2D-C3D	-2.64	121.54	127.61
29	g	613	CLA	C1C-C2C-C3C	-2.64	104.18	106.96
29	r	608	CLA	C1C-C2C-C3C	-2.64	104.18	106.96
29	r	608	CLA	C2D-C1D-ND	2.64	112.05	110.10
31	c	516	BCR	C37-C22-C23	2.64	122.23	118.08
48	y	623	NEX	C38-C25-C26	-2.64	117.84	122.26
46	n	620	LUT	C31-C30-C29	-2.64	123.55	127.31
29	B	610	CLA	O2D-CGD-O1D	-2.64	118.68	123.84
29	R	608	CLA	CMA-C3A-C4A	2.64	118.86	111.77
29	C	509	CLA	O2D-CGD-O1D	-2.64	118.68	123.84
29	s	611	CLA	O2D-CGD-O1D	-2.64	118.68	123.84

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
38	c	520	DGD	O1G-C1A-C2A	2.64	120.18	111.91
45	N	601	CHL	C1-O2A-CGA	2.64	123.36	116.44
29	Y	604	CLA	C1-C2-C3	-2.64	121.48	126.04
45	y	609	CHL	C1-O2A-CGA	2.64	123.36	116.44
29	Y	613	CLA	C1C-C2C-C3C	-2.63	104.19	106.96
29	s	605	CLA	C2D-C1D-ND	2.63	112.05	110.10
29	G	603	CLA	CAA-C2A-C3A	-2.63	105.57	112.78
29	s	605	CLA	CHA-C4D-ND	2.63	138.01	132.50
29	c	506	CLA	CMA-C3A-C4A	2.63	118.85	111.77
46	S	621	LUT	C38-C25-C24	-2.63	117.93	123.56
45	y	607	CHL	C1-O2A-CGA	2.63	123.35	116.44
29	B	613	CLA	C2C-C1C-NC	2.63	112.44	109.97
39	d	409	LHG	O8-C23-C24	2.63	120.17	111.91
29	C	505	CLA	C1C-C2C-C3C	-2.63	104.19	106.96
29	S	611	CLA	C1-C2-C3	-2.63	121.50	126.04
29	s	614	CLA	C1-C2-C3	-2.63	121.50	126.04
29	n	610	CLA	CMA-C3A-C4A	2.63	118.84	111.77
29	b	611	CLA	C1C-C2C-C3C	-2.63	104.19	106.96
31	b	618	BCR	C15-C14-C13	-2.63	123.56	127.31
47	N	622	XAT	C18-C5-C6	-2.63	117.86	122.26
29	d	403	CLA	C1C-C2C-C3C	-2.63	104.19	106.96
29	r	609	CLA	C1C-C2C-C3C	-2.63	104.19	106.96
50	s	626	3PH	O31-C31-C32	2.63	120.15	111.91
29	Y	604	CLA	C2D-C1D-ND	2.63	112.04	110.10
29	Y	614	CLA	C2D-C1D-ND	2.63	112.04	110.10
39	S	624	LHG	C5-O7-C7	-2.62	111.33	117.79
48	n	623	NEX	C1-C2-C3	2.62	119.57	113.64
29	S	602	CLA	CMA-C3A-C4A	2.62	118.82	111.77
47	n	622	XAT	C18-C5-C6	-2.62	117.86	122.26
29	s	612	CLA	C1C-C2C-C3C	-2.62	104.20	106.96
29	s	603	CLA	CHD-C1D-ND	-2.62	122.04	124.45
29	S	613	CLA	CHA-C4D-ND	2.62	137.98	132.50
29	C	501	CLA	C1C-C2C-C3C	-2.62	104.20	106.96
29	r	611	CLA	C2D-C1D-ND	2.62	112.04	110.10
38	C	518	DGD	O6D-C5D-C6D	2.62	111.96	106.67
29	C	505	CLA	C1-O2A-CGA	2.62	123.32	116.44
29	c	513	CLA	CMA-C3A-C4A	2.62	118.82	111.77
29	c	510	CLA	CMD-C2D-C3D	-2.62	121.59	127.61
29	Y	612	CLA	O2D-CGD-O1D	-2.62	118.72	123.84
48	Y	623	NEX	C4-C3-C2	2.62	115.83	110.77
29	y	612	CLA	C1C-C2C-C3C	-2.62	104.20	106.96
29	C	505	CLA	O2A-CGA-CBA	2.62	120.13	111.91

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
29	A	405	CLA	C6-C5-C3	-2.62	106.59	113.45
29	C	512	CLA	C1C-C2C-C3C	-2.62	104.20	106.96
29	y	613	CLA	C1C-C2C-C3C	-2.62	104.20	106.96
46	N	621	LUT	C38-C25-C24	-2.62	117.96	123.56
29	Y	610	CLA	C2C-C1C-NC	2.62	112.42	109.97
29	N	614	CLA	O2D-CGD-O1D	-2.62	118.72	123.84
45	Y	607	CHL	C4A-NA-C1A	2.62	107.88	106.71
29	B	605	CLA	O2D-CGD-O1D	-2.62	118.72	123.84
29	c	507	CLA	C2D-C1D-ND	2.62	112.03	110.10
42	D	405	PL9	C40-C39-C41	2.62	119.67	115.27
29	B	609	CLA	CHA-C4D-ND	2.62	137.97	132.50
29	B	607	CLA	CMB-C2B-C3B	2.62	129.57	124.68
31	a	411	BCR	C33-C5-C4	2.62	118.64	113.62
48	G	623	NEX	C1-C2-C3	2.62	119.55	113.64
29	b	613	CLA	C2C-C1C-NC	2.62	112.42	109.97
31	C	514	BCR	C36-C18-C17	-2.61	119.26	122.92
29	g	602	CLA	O2D-CGD-O1D	-2.61	118.73	123.84
29	b	616	CLA	O2D-CGD-O1D	-2.61	118.73	123.84
29	y	614	CLA	O2D-CGD-O1D	-2.61	118.73	123.84
29	y	613	CLA	O2A-CGA-CBA	2.61	120.11	111.91
29	c	513	CLA	O2A-CGA-CBA	2.61	120.11	111.91
35	B	620	C7Z	C28-C27-C26	-2.61	119.87	127.20
42	D	405	PL9	C22-C23-C24	-2.61	121.37	127.66
50	i	101	3PH	O31-C31-C32	2.61	120.10	111.91
31	c	514	BCR	C38-C26-C25	-2.61	121.60	124.53
29	G	604	CLA	O2D-CGD-O1D	-2.61	118.74	123.84
29	b	604	CLA	C1C-C2C-C3C	-2.61	104.21	106.96
29	b	604	CLA	CMA-C3A-C4A	2.61	118.78	111.77
29	r	602	CLA	C1C-C2C-C3C	-2.61	104.22	106.96
31	c	515	BCR	C34-C9-C10	-2.61	119.27	122.92
29	B	606	CLA	O2D-CGD-O1D	-2.61	118.74	123.84
29	r	609	CLA	CHA-C4D-ND	2.61	137.95	132.50
29	s	613	CLA	CHA-C4D-ND	2.61	137.95	132.50
29	S	604	CLA	O2D-CGD-O1D	-2.60	118.75	123.84
29	y	603	CLA	O2D-CGD-O1D	-2.60	118.75	123.84
48	R	622	NEX	C20-C13-C14	-2.60	119.28	122.92
29	g	604	CLA	C1C-C2C-C3C	-2.60	104.22	106.96
45	G	608	CHL	CHB-C4A-NA	2.60	128.11	124.51
29	S	605	CLA	C1C-C2C-C3C	-2.60	104.22	106.96
29	b	608	CLA	CMA-C3A-C4A	2.60	118.77	111.77
29	b	605	CLA	CHA-C4D-ND	2.60	137.94	132.50
29	Y	603	CLA	O2D-CGD-O1D	-2.60	118.75	123.84

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
46	S	620	LUT	C31-C30-C29	-2.60	123.60	127.31
31	A	411	BCR	C35-C13-C12	2.60	122.17	118.08
31	B	619	BCR	C1-C6-C5	-2.60	118.95	122.61
29	S	605	CLA	CHA-C4D-ND	2.60	137.94	132.50
29	g	611	CLA	O2D-CGD-O1D	-2.60	118.75	123.84
45	s	601	CHL	CHB-C4A-NA	2.60	128.11	124.51
29	a	406	CLA	CMA-C3A-C4A	2.60	118.76	111.77
29	B	609	CLA	CMB-C2B-C3B	2.60	129.54	124.68
29	R	613	CLA	O2D-CGD-O1D	-2.60	118.76	123.84
29	C	502	CLA	O2D-CGD-O1D	-2.60	118.76	123.84
45	y	609	CHL	C1B-CHB-C4A	-2.60	124.97	130.12
29	y	603	CLA	CMA-C3A-C4A	2.60	118.75	111.77
29	Y	602	CLA	O2D-CGD-O1D	-2.60	118.76	123.84
29	R	610	CLA	C2C-C1C-NC	2.60	112.40	109.97
29	B	611	CLA	O2D-CGD-O1D	-2.60	118.76	123.84
29	r	610	CLA	C2C-C1C-NC	2.59	112.40	109.97
29	R	608	CLA	C1-C2-C3	-2.59	121.56	126.04
29	R	603	CLA	C2D-C1D-ND	2.59	112.02	110.10
29	b	615	CLA	C2D-C1D-ND	2.59	112.02	110.10
29	C	505	CLA	CHA-C4D-ND	2.59	137.92	132.50
45	G	609	CHL	CHC-C1C-NC	2.59	128.13	124.20
29	B	607	CLA	O2D-CGD-O1D	-2.59	118.77	123.84
33	c	521	LMG	C8-O7-C10	-2.59	111.41	117.79
46	g	620	LUT	C38-C25-C24	-2.59	118.02	123.56
45	y	601	CHL	C1-O2A-CGA	2.59	123.24	116.44
29	B	612	CLA	C2D-C1D-ND	2.59	112.01	110.10
29	y	612	CLA	O2D-CGD-O1D	-2.59	118.78	123.84
29	B	603	CLA	CMA-C3A-C4A	2.59	118.73	111.77
29	N	610	CLA	C1C-C2C-C3C	-2.59	104.24	106.96
29	n	602	CLA	C1C-C2C-C3C	-2.59	104.24	106.96
29	s	613	CLA	C1C-C2C-C3C	-2.59	104.24	106.96
31	c	517	BCR	C23-C24-C25	-2.59	119.94	127.20
29	s	611	CLA	C2D-C1D-ND	2.59	112.01	110.10
29	R	609	CLA	CHA-C4D-ND	2.59	137.91	132.50
29	s	605	CLA	C1C-C2C-C3C	-2.59	104.24	106.96
46	r	620	LUT	C3-C4-C5	-2.59	106.70	111.85
46	r	620	LUT	C11-C12-C13	-2.59	119.15	126.42
31	B	619	BCR	C19-C18-C17	2.58	122.91	118.94
29	b	603	CLA	C2D-C1D-ND	2.58	112.01	110.10
29	G	611	CLA	O2D-CGD-O1D	-2.58	118.79	123.84
29	S	603	CLA	CHD-C1D-ND	-2.58	122.08	124.45
33	J	101	LMG	O8-C28-C29	2.58	120.00	111.91

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
29	c	505	CLA	CHA-C4D-ND	2.58	137.90	132.50
29	D	403	CLA	C1C-C2C-C3C	-2.58	104.24	106.96
29	G	604	CLA	C1C-C2C-C3C	-2.58	104.24	106.96
29	B	608	CLA	CMA-C3A-C4A	2.58	118.70	111.77
29	b	617	CLA	CHD-C1D-ND	-2.58	122.08	124.45
29	B	604	CLA	CMA-C3A-C4A	2.58	118.70	111.77
29	Y	610	CLA	O2A-CGA-CBA	2.58	120.00	111.91
29	y	611	CLA	C2D-C1D-ND	2.58	112.00	110.10
31	C	514	BCR	C23-C24-C25	-2.58	119.97	127.20
29	Y	611	CLA	CMA-C3A-C4A	2.57	118.69	111.77
29	B	615	CLA	CHA-C4D-ND	2.57	137.88	132.50
31	c	515	BCR	C15-C14-C13	-2.57	123.64	127.31
45	N	609	CHL	C1B-CHB-C4A	-2.57	125.02	130.12
29	B	602	CLA	CHA-C4D-ND	2.57	137.88	132.50
29	b	606	CLA	CMD-C2D-C3D	-2.57	121.69	127.61
29	C	507	CLA	C2D-C1D-ND	2.57	112.00	110.10
29	n	612	CLA	O2D-CGD-O1D	-2.57	118.81	123.84
33	C	521	LMG	C8-O7-C10	-2.57	111.46	117.79
47	r	622	XAT	C8-C9-C10	2.57	122.89	118.94
29	y	604	CLA	CMB-C2B-C3B	2.57	129.49	124.68
29	Y	612	CLA	C1-C2-C3	-2.57	121.59	126.04
29	b	606	CLA	O2D-CGD-O1D	-2.57	118.81	123.84
29	s	613	CLA	CHD-C1D-ND	-2.57	122.09	124.45
29	Y	604	CLA	C1C-C2C-C3C	-2.57	104.25	106.96
29	g	612	CLA	C1C-C2C-C3C	-2.57	104.25	106.96
29	C	511	CLA	O2D-CGD-O1D	-2.57	118.81	123.84
29	y	602	CLA	O2D-CGD-O1D	-2.57	118.81	123.84
50	S	626	3PH	O31-C31-C32	2.57	119.97	111.91
29	N	602	CLA	C2D-C1D-ND	2.57	112.00	110.10
31	B	618	BCR	C36-C18-C17	-2.57	119.32	122.92
43	F	101	HEM	CHC-C4B-C3B	2.57	128.50	124.57
29	C	501	CLA	CHA-C4D-ND	2.57	137.87	132.50
29	r	609	CLA	CMD-C2D-C3D	-2.57	121.71	127.61
29	g	612	CLA	O2D-CGD-O1D	-2.57	118.82	123.84
29	s	609	CLA	C1-C2-C3	-2.56	121.61	126.04
29	Y	611	CLA	C1C-C2C-C3C	-2.56	104.26	106.96
29	g	604	CLA	CHA-C4D-ND	2.56	137.86	132.50
29	b	613	CLA	C2D-C1D-ND	2.56	111.99	110.10
45	g	609	CHL	C3C-C4C-NC	-2.56	107.70	110.57
46	Y	620	LUT	C31-C30-C29	-2.56	123.66	127.31
29	B	603	CLA	C1C-C2C-C3C	-2.56	104.27	106.96
29	c	506	CLA	C1-C2-C3	-2.56	121.62	126.04

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
31	C	516	BCR	C36-C18-C17	-2.56	119.34	122.92
29	c	512	CLA	C1D-ND-C4D	-2.56	104.52	106.33
29	S	609	CLA	CAA-C2A-C3A	-2.56	105.77	112.78
29	n	604	CLA	C1C-C2C-C3C	-2.56	104.27	106.96
29	c	501	CLA	CHA-C4D-ND	2.56	137.85	132.50
45	R	607	CHL	C1-O2A-CGA	2.56	123.15	116.44
31	d	404	BCR	C33-C5-C4	2.56	118.53	113.62
45	n	608	CHL	C1B-CHB-C4A	-2.56	125.06	130.12
29	B	603	CLA	CMB-C2B-C3B	2.56	129.46	124.68
39	y	624	LHG	O8-C23-C24	2.55	119.92	111.91
29	c	513	CLA	O2D-CGD-O1D	-2.55	118.84	123.84
29	Y	612	CLA	C1C-C2C-C3C	-2.55	104.27	106.96
29	b	609	CLA	CHA-C4D-ND	2.55	137.84	132.50
31	C	515	BCR	C23-C24-C25	-2.55	120.03	127.20
29	C	506	CLA	C1-C2-C3	-2.55	121.63	126.04
29	y	604	CLA	C2D-C1D-ND	2.55	111.99	110.10
29	B	604	CLA	CHA-C4D-ND	2.55	137.84	132.50
29	b	604	CLA	CHA-C4D-ND	2.55	137.84	132.50
29	C	513	CLA	O2A-CGA-CBA	2.55	119.92	111.91
45	Y	605	CHL	C1B-CHB-C4A	-2.55	125.06	130.12
29	c	502	CLA	C2D-C1D-ND	2.55	111.98	110.10
45	S	607	CHL	CHB-C4A-NA	2.55	128.04	124.51
29	y	611	CLA	C1C-C2C-C3C	-2.55	104.28	106.96
29	R	612	CLA	CHA-C4D-ND	2.55	137.83	132.50
29	B	606	CLA	CMD-C2D-C3D	-2.55	121.75	127.61
45	R	606	CHL	C4A-NA-C1A	2.55	107.85	106.71
31	C	517	BCR	C36-C18-C17	-2.55	119.35	122.92
31	c	517	BCR	C31-C1-C6	-2.55	106.17	110.30
29	s	602	CLA	C1D-ND-C4D	-2.55	104.53	106.33
29	y	603	CLA	C1D-ND-C4D	-2.55	104.53	106.33
45	g	609	CHL	CHD-C4C-C3C	2.55	128.59	124.84
29	b	603	CLA	O2D-CGD-O1D	-2.55	118.86	123.84
44	h	101	RRX	C8-C7-C6	-2.55	120.05	127.20
45	r	607	CHL	C1-O2A-CGA	2.55	123.13	116.44
29	R	613	CLA	C1C-C2C-C3C	-2.55	104.28	106.96
33	a	413	LMG	O8-C28-C29	2.55	119.90	111.91
29	D	402	CLA	CMC-C2C-C1C	2.54	128.91	125.04
29	N	612	CLA	CHD-C1D-ND	-2.54	122.12	124.45
29	N	604	CLA	CHA-C4D-ND	2.54	137.82	132.50
29	N	604	CLA	O2D-CGD-O1D	-2.54	118.86	123.84
45	r	606	CHL	C3C-C4C-NC	-2.54	107.72	110.57
29	N	612	CLA	C1C-C2C-C3C	-2.54	104.28	106.96

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
47	r	622	XAT	O4-C5-C4	-2.54	111.47	113.38
29	C	508	CLA	CMA-C3A-C4A	2.54	118.61	111.77
45	n	605	CHL	CHB-C4A-NA	2.54	128.03	124.51
29	r	611	CLA	C1C-C2C-C3C	-2.54	104.28	106.96
29	c	506	CLA	CMB-C2B-C1B	-2.54	124.56	128.46
29	n	604	CLA	O2D-CGD-O1D	-2.54	118.87	123.84
29	n	604	CLA	CHA-C4D-ND	2.54	137.81	132.50
29	d	403	CLA	O2A-CGA-CBA	2.54	119.88	111.91
29	c	507	CLA	C1-C2-C3	-2.54	121.65	126.04
29	y	613	CLA	CMB-C2B-C3B	2.54	129.43	124.68
29	r	612	CLA	O2A-CGA-CBA	2.54	119.88	111.91
29	N	613	CLA	C1C-C2C-C3C	-2.54	104.29	106.96
29	s	610	CLA	CHA-C4D-ND	2.54	137.81	132.50
29	b	602	CLA	CHA-C4D-ND	2.54	137.81	132.50
31	a	411	BCR	C36-C18-C17	-2.54	119.37	122.92
44	h	101	RRX	C24-C23-C22	-2.54	122.40	126.23
29	b	608	CLA	CMB-C2B-C3B	2.54	129.42	124.68
29	C	507	CLA	CHA-C4D-ND	2.54	137.80	132.50
29	y	602	CLA	C2C-C1C-NC	2.54	112.35	109.97
29	C	506	CLA	CMA-C3A-C4A	2.53	118.59	111.77
45	G	606	CHL	CHB-C4A-NA	2.53	128.02	124.51
45	S	601	CHL	CHB-C4A-NA	2.53	128.02	124.51
29	n	614	CLA	O2D-CGD-O1D	-2.53	118.88	123.84
29	S	613	CLA	O1D-CGD-CBD	-2.53	119.30	124.48
29	c	511	CLA	O2D-CGD-O1D	-2.53	118.89	123.84
29	b	615	CLA	CHA-C4D-ND	2.53	137.80	132.50
29	Y	614	CLA	O2D-CGD-O1D	-2.53	118.89	123.84
45	G	606	CHL	C1-O2A-CGA	2.53	123.09	116.44
29	y	604	CLA	C1C-C2C-C3C	-2.53	104.29	106.96
29	Y	612	CLA	CHA-C4D-ND	2.53	137.80	132.50
29	S	612	CLA	CMA-C3A-C4A	2.53	118.57	111.77
29	R	609	CLA	C1C-C2C-C3C	-2.53	104.30	106.96
29	d	402	CLA	O2A-CGA-CBA	2.53	119.84	111.91
42	d	405	PL9	C22-C23-C24	-2.53	121.57	127.66
29	R	611	CLA	C1C-C2C-C3C	-2.53	104.30	106.96
29	R	612	CLA	O2D-CGD-O1D	-2.53	118.90	123.84
29	Y	608	CLA	C1C-C2C-C3C	-2.53	104.30	106.96
47	R	621	XAT	C18-C5-C6	-2.53	118.03	122.26
46	s	620	LUT	C11-C10-C9	-2.53	123.71	127.31
29	C	512	CLA	C1-C2-C3	-2.52	121.68	126.04
45	G	607	CHL	C4A-NA-C1A	2.52	107.84	106.71
29	n	612	CLA	CMA-C3A-C4A	2.52	118.56	111.77

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
48	G	623	NEX	C16-C1-C6	-2.52	108.21	110.47
29	B	603	CLA	CHA-C4D-ND	2.52	137.78	132.50
29	s	605	CLA	CHD-C1D-ND	-2.52	122.14	124.45
29	s	604	CLA	C1-C2-C3	-2.52	121.68	126.04
29	s	614	CLA	O2D-CGD-O1D	-2.52	118.91	123.84
29	b	606	CLA	C2D-C1D-ND	2.52	111.96	110.10
45	g	608	CHL	CHB-C4A-NA	2.52	128.00	124.51
45	g	605	CHL	C4A-NA-C1A	2.52	107.84	106.71
29	N	611	CLA	C1C-C2C-C3C	-2.52	104.31	106.96
29	b	617	CLA	C2D-C1D-ND	2.52	111.96	110.10
29	g	603	CLA	C1D-ND-C4D	-2.52	104.54	106.33
45	G	609	CHL	CHD-C4C-C3C	2.52	128.54	124.84
39	G	630	LHG	O8-C23-C24	2.52	119.82	111.91
29	N	602	CLA	CAA-C2A-C3A	-2.52	105.88	112.78
29	Y	608	CLA	CHA-C4D-ND	2.52	137.77	132.50
43	f	101	HEM	CHC-C4B-C3B	2.52	128.43	124.57
33	A	413	LMG	O8-C28-C29	2.52	119.81	111.91
29	n	613	CLA	C1-O2A-CGA	2.52	123.05	116.44
48	S	622	NEX	C1-C2-C3	2.52	119.33	113.64
29	n	611	CLA	C1C-C2C-C3C	-2.52	104.31	106.96
29	G	604	CLA	CMA-C3A-C4A	2.52	118.53	111.77
29	S	603	CLA	CMA-C3A-C4A	2.51	118.53	111.77
29	C	509	CLA	C2D-C1D-ND	2.51	111.96	110.10
45	n	605	CHL	C1-O2A-CGA	2.51	123.04	116.44
29	y	608	CLA	CHA-C4D-ND	2.51	137.76	132.50
29	S	609	CLA	C1-C2-C3	-2.51	121.70	126.04
29	S	610	CLA	CHA-C4D-ND	2.51	137.76	132.50
29	c	513	CLA	CMB-C2B-C3B	2.51	129.38	124.68
29	y	610	CLA	O2A-CGA-CBA	2.51	119.79	111.91
29	n	610	CLA	C1C-C2C-C3C	-2.51	104.32	106.96
29	b	617	CLA	OBD-CAD-C3D	-2.51	122.48	128.52
45	N	609	CHL	C1-O2A-CGA	2.51	123.03	116.44
35	B	620	C7Z	C21-C26-C25	-2.51	119.08	122.61
31	b	619	BCR	C8-C7-C6	-2.51	120.16	127.20
29	n	613	CLA	C1C-C2C-C3C	-2.51	104.32	106.96
29	B	613	CLA	CHA-C4D-ND	2.51	137.75	132.50
29	n	611	CLA	CHA-C4D-ND	2.51	137.75	132.50
29	s	611	CLA	CHA-C4D-ND	2.51	137.75	132.50
29	B	604	CLA	O2D-CGD-O1D	-2.51	118.94	123.84
29	N	603	CLA	O2D-CGD-O1D	-2.51	118.94	123.84
29	y	612	CLA	CHA-C4D-ND	2.51	137.74	132.50
29	Y	612	CLA	O2A-CGA-CBA	2.51	119.78	111.91

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
45	n	601	CHL	CHB-C4A-NA	2.51	127.98	124.51
39	g	624	LHG	O8-C23-C24	2.51	119.78	111.91
29	R	604	CLA	CHA-C4D-ND	2.51	137.74	132.50
45	Y	601	CHL	C4A-NA-C1A	2.51	107.83	106.71
29	C	503	CLA	O1D-CGD-CBD	-2.51	119.36	124.48
46	R	620	LUT	C22-C23-C24	-2.51	108.89	111.74
29	c	510	CLA	O2D-CGD-O1D	-2.51	118.94	123.84
29	s	612	CLA	O2D-CGD-O1D	-2.50	118.94	123.84
46	n	620	LUT	C18-C5-C6	-2.50	121.72	124.53
29	r	602	CLA	CMA-C3A-C4A	2.50	118.50	111.77
29	c	510	CLA	C2D-C1D-ND	2.50	111.95	110.10
29	B	612	CLA	O2D-CGD-O1D	-2.50	118.94	123.84
29	Y	611	CLA	O2D-CGD-O1D	-2.50	118.95	123.84
29	B	607	CLA	CHA-C4D-ND	2.50	137.73	132.50
45	S	608	CHL	C1-O2A-CGA	2.50	123.00	116.44
29	R	613	CLA	CHA-C4D-ND	2.50	137.73	132.50
46	G	621	LUT	C10-C11-C12	-2.50	115.42	123.22
45	S	606	CHL	C1B-CHB-C4A	-2.50	125.17	130.12
29	c	506	CLA	CMB-C2B-C3B	2.50	129.35	124.68
29	r	608	CLA	CHA-C4D-ND	2.50	137.73	132.50
29	C	501	CLA	CMD-C2D-C3D	-2.50	121.87	127.61
29	S	614	CLA	CHA-C4D-ND	2.50	137.72	132.50
29	c	507	CLA	CHA-C4D-ND	2.50	137.72	132.50
46	N	620	LUT	C38-C25-C24	-2.50	118.21	123.56
29	S	612	CLA	O2D-CGD-O1D	-2.50	118.95	123.84
45	S	607	CHL	C4A-NA-C1A	2.50	107.83	106.71
45	s	606	CHL	C1B-CHB-C4A	-2.50	125.17	130.12
35	B	620	C7Z	C27-C28-C29	-2.50	122.46	126.23
29	G	602	CLA	O2D-CGD-O1D	-2.50	118.96	123.84
29	g	603	CLA	O2D-CGD-O1D	-2.50	118.96	123.84
29	B	617	CLA	O2D-CGD-O1D	-2.50	118.96	123.84
29	b	611	CLA	CHA-C4D-ND	2.50	137.72	132.50
29	c	510	CLA	CHA-C4D-ND	2.50	137.72	132.50
29	r	612	CLA	CHA-C4D-ND	2.50	137.72	132.50
31	b	618	BCR	C36-C18-C17	-2.49	119.43	122.92
29	R	609	CLA	CMD-C2D-C3D	-2.49	121.88	127.61
45	y	601	CHL	C4A-NA-C1A	2.49	107.83	106.71
29	c	504	CLA	CMB-C2B-C3B	2.49	129.34	124.68
29	b	606	CLA	CHA-C4D-ND	2.49	137.71	132.50
29	G	612	CLA	O2D-CGD-O1D	-2.49	118.96	123.84
45	N	609	CHL	CHD-C4C-C3C	2.49	128.50	124.84
29	c	503	CLA	C1C-C2C-C3C	-2.49	104.34	106.96

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
45	n	605	CHL	CMA-C3A-C4A	2.49	118.47	111.77
29	G	604	CLA	CHA-C4D-ND	2.49	137.71	132.50
31	B	619	BCR	C27-C26-C25	-2.49	119.11	122.73
45	g	601	CHL	C1-O2A-CGA	2.49	122.98	116.44
29	C	504	CLA	C1C-C2C-C3C	-2.49	104.34	106.96
29	B	613	CLA	CMB-C2B-C3B	2.49	129.34	124.68
43	f	101	HEM	C4D-ND-C1D	2.49	107.65	105.07
29	b	607	CLA	CHA-C4D-ND	2.49	137.71	132.50
29	b	608	CLA	CHA-C4D-ND	2.49	137.71	132.50
29	b	614	CLA	CHA-C4D-ND	2.49	137.71	132.50
29	b	613	CLA	C1-C2-C3	-2.49	121.74	126.04
29	g	613	CLA	O2A-CGA-CBA	2.49	119.72	111.91
29	s	605	CLA	O1D-CGD-CBD	-2.49	119.39	124.48
29	B	604	CLA	C1C-C2C-C3C	-2.49	104.34	106.96
31	c	516	BCR	C23-C24-C25	-2.49	120.21	127.20
29	Y	610	CLA	CMA-C3A-C4A	2.49	118.46	111.77
29	S	614	CLA	CMD-C2D-C3D	-2.49	121.89	127.61
29	c	505	CLA	CMA-C3A-C4A	2.49	118.46	111.77
29	C	512	CLA	CMB-C2B-C1B	-2.49	124.64	128.46
29	Y	603	CLA	CHA-C4D-ND	2.49	137.70	132.50
29	R	604	CLA	O2D-CGD-O1D	-2.49	118.98	123.84
29	c	503	CLA	CMD-C2D-C3D	-2.49	121.90	127.61
29	b	613	CLA	CHA-C4D-ND	2.48	137.70	132.50
29	y	608	CLA	C1C-C2C-C3C	-2.48	104.34	106.96
31	d	404	BCR	C38-C26-C27	2.48	118.39	113.62
48	R	622	NEX	C19-C9-C10	-2.48	119.44	122.92
44	H	101	RRX	C24-C23-C22	-2.48	122.48	126.23
45	s	607	CHL	CHB-C4A-NA	2.48	127.94	124.51
29	c	509	CLA	CMB-C2B-C3B	2.48	129.32	124.68
29	c	511	CLA	CHA-C4D-ND	2.48	137.69	132.50
29	R	602	CLA	O2A-CGA-CBA	2.48	119.69	111.91
45	N	608	CHL	CHB-C4A-NA	2.48	127.94	124.51
29	C	511	CLA	CHA-C4D-ND	2.48	137.69	132.50
45	r	606	CHL	C4A-NA-C1A	2.48	107.82	106.71
29	D	402	CLA	CHA-C4D-ND	2.48	137.69	132.50
29	s	614	CLA	CHA-C4D-ND	2.48	137.69	132.50
29	R	602	CLA	CHA-C4D-ND	2.48	137.68	132.50
29	y	610	CLA	C1C-C2C-C3C	-2.48	104.35	106.96
29	s	612	CLA	CMA-C3A-C4A	2.48	118.43	111.77
29	G	602	CLA	O2A-CGA-CBA	2.48	119.68	111.91
29	B	614	CLA	O2D-CGD-O1D	-2.48	119.00	123.84
46	G	621	LUT	C35-C15-C14	-2.48	118.40	123.47

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
29	S	614	CLA	C2D-C1D-ND	2.48	111.93	110.10
45	n	601	CHL	C1-C2-C3	-2.48	121.76	126.04
29	r	604	CLA	CHA-C4D-ND	2.48	137.68	132.50
48	G	623	NEX	C20-C13-C14	-2.48	119.45	122.92
45	Y	609	CHL	C1B-CHB-C4A	-2.48	125.21	130.12
45	R	606	CHL	C3C-C4C-NC	-2.48	107.80	110.57
29	C	505	CLA	CMA-C3A-C4A	2.47	118.42	111.77
29	b	612	CLA	C2A-C1A-CHA	2.47	128.19	123.86
29	N	603	CLA	CHA-C4D-ND	2.47	137.68	132.50
29	N	612	CLA	CHA-C4D-ND	2.47	137.68	132.50
29	S	611	CLA	CHA-C4D-ND	2.47	137.68	132.50
31	C	515	BCR	C34-C9-C10	-2.47	119.46	122.92
29	c	503	CLA	O1D-CGD-CBD	-2.47	119.42	124.48
29	c	509	CLA	CHA-C4D-ND	2.47	137.67	132.50
39	D	409	LHG	C5-O7-C7	-2.47	111.71	117.79
29	S	605	CLA	O1D-CGD-CBD	-2.47	119.43	124.48
29	g	611	CLA	CHA-C4D-ND	2.47	137.66	132.50
29	d	402	CLA	C1-C2-C3	-2.47	121.77	126.04
39	C	525	LHG	O8-C23-C24	2.47	119.65	111.91
31	A	411	BCR	C36-C18-C17	-2.47	119.47	122.92
48	N	623	NEX	C38-C25-C26	-2.47	118.12	122.26
29	y	608	CLA	C1-O2A-CGA	2.47	122.92	116.44
29	b	604	CLA	CMB-C2B-C3B	2.47	129.29	124.68
29	c	508	CLA	C2D-C1D-ND	2.47	111.92	110.10
29	b	607	CLA	C1C-C2C-C3C	-2.47	104.36	106.96
29	r	611	CLA	CHA-C4D-ND	2.47	137.66	132.50
36	B	625	DGA	OG1-CA1-CA2	2.47	119.64	111.91
46	S	621	LUT	C10-C11-C12	-2.47	115.52	123.22
29	g	613	CLA	CHA-C4D-ND	2.47	137.66	132.50
29	Y	613	CLA	O2A-CGA-CBA	2.47	119.64	111.91
45	S	606	CHL	C4D-CHA-C1A	2.47	124.25	121.25
29	g	610	CLA	O2A-CGA-CBA	2.46	119.64	111.91
29	G	613	CLA	CHA-C4D-ND	2.46	137.66	132.50
29	B	607	CLA	C1C-C2C-C3C	-2.46	104.36	106.96
29	b	612	CLA	CHA-C1A-NA	-2.46	120.75	126.40
46	s	620	LUT	C10-C11-C12	-2.46	115.53	123.22
29	C	510	CLA	CMD-C2D-C3D	-2.46	121.94	127.61
29	S	603	CLA	CHA-C4D-ND	2.46	137.65	132.50
29	B	608	CLA	CMB-C2B-C3B	2.46	129.29	124.68
29	G	613	CLA	O2A-CGA-CBA	2.46	119.64	111.91
47	r	622	XAT	C18-C5-C6	-2.46	118.13	122.26
29	B	608	CLA	CHA-C4D-ND	2.46	137.65	132.50

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
29	b	603	CLA	O2A-CGA-CBA	2.46	119.63	111.91
29	C	512	CLA	C1D-ND-C4D	-2.46	104.59	106.33
29	C	510	CLA	O2D-CGD-O1D	-2.46	119.03	123.84
29	s	617	CLA	O2D-CGD-O1D	-2.46	119.03	123.84
45	s	606	CHL	CHB-C4A-NA	2.46	127.92	124.51
29	c	512	CLA	CMB-C2B-C1B	-2.46	124.68	128.46
45	N	606	CHL	C1-O2A-CGA	2.46	122.90	116.44
29	R	608	CLA	CHA-C4D-ND	2.46	137.64	132.50
29	y	614	CLA	CHA-C4D-ND	2.46	137.64	132.50
29	c	509	CLA	CBC-CAC-C3C	-2.46	105.65	112.43
39	Y	624	LHG	O8-C23-C24	2.46	119.62	111.91
29	R	603	CLA	C1-C2-C3	-2.46	121.79	126.04
39	N	624	LHG	O8-C23-C24	2.46	119.62	111.91
45	Y	601	CHL	C1-O2A-CGA	2.46	122.89	116.44
29	s	603	CLA	C1C-C2C-C3C	-2.46	104.37	106.96
29	b	612	CLA	O2D-CGD-O1D	-2.46	119.04	123.84
29	D	403	CLA	O2A-CGA-CBA	2.46	119.61	111.91
29	B	606	CLA	CHA-C4D-ND	2.46	137.64	132.50
29	S	612	CLA	CHA-C4D-ND	2.46	137.64	132.50
46	N	620	LUT	C18-C5-C6	-2.46	121.77	124.53
29	S	617	CLA	O2D-CGD-O1D	-2.46	119.04	123.84
29	s	612	CLA	CHA-C4D-ND	2.45	137.63	132.50
29	c	508	CLA	CMA-C3A-C4A	2.45	118.37	111.77
29	C	509	CLA	CHA-C4D-ND	2.45	137.63	132.50
29	C	506	CLA	C2D-C1D-ND	2.45	111.91	110.10
29	y	611	CLA	CHA-C4D-ND	2.45	137.63	132.50
29	b	613	CLA	C1C-C2C-C3C	-2.45	104.38	106.96
29	c	504	CLA	C1C-C2C-C3C	-2.45	104.38	106.96
29	s	605	CLA	C1-O2A-CGA	2.45	122.88	116.44
29	B	611	CLA	CHA-C4D-ND	2.45	137.63	132.50
29	R	603	CLA	CHA-C4D-ND	2.45	137.63	132.50
45	n	605	CHL	C1-C2-C3	-2.45	121.80	126.04
29	S	611	CLA	O2D-CGD-O1D	-2.45	119.05	123.84
29	g	612	CLA	CHA-C4D-ND	2.45	137.63	132.50
29	s	604	CLA	CHA-C4D-ND	2.45	137.63	132.50
29	B	603	CLA	CMD-C2D-C3D	-2.45	121.98	127.61
29	c	511	CLA	CMB-C2B-C3B	2.45	129.26	124.68
29	S	613	CLA	CHD-C1D-ND	-2.45	122.20	124.45
29	C	506	CLA	CMB-C2B-C3B	2.45	129.26	124.68
29	N	611	CLA	CHA-C4D-ND	2.45	137.62	132.50
29	B	614	CLA	CHA-C4D-ND	2.45	137.62	132.50
29	y	602	CLA	CHA-C4D-ND	2.45	137.62	132.50

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
29	r	612	CLA	O2D-CGD-O1D	-2.45	119.05	123.84
31	c	516	BCR	C38-C26-C25	-2.45	121.78	124.53
29	R	602	CLA	C1C-C2C-C3C	-2.45	104.38	106.96
29	G	614	CLA	CHA-C4D-ND	2.45	137.62	132.50
31	C	517	BCR	C33-C5-C6	-2.45	121.78	124.53
29	y	603	CLA	CHA-C4D-ND	2.45	137.62	132.50
46	G	621	LUT	C31-C30-C29	-2.45	123.82	127.31
47	N	622	XAT	O4-C5-C4	-2.44	111.55	113.38
29	n	613	CLA	CHA-C4D-ND	2.44	137.61	132.50
48	r	623	NEX	C39-C29-C30	-2.44	119.50	122.92
33	b	622	LMG	O8-C28-C29	2.44	119.58	111.91
29	C	511	CLA	CMD-C2D-C3D	-2.44	121.99	127.61
47	r	622	XAT	C31-C30-C29	-2.44	123.82	127.31
29	R	612	CLA	O2A-CGA-CBA	2.44	119.57	111.91
45	y	609	CHL	C4D-CHA-C1A	2.44	124.22	121.25
31	c	514	BCR	C34-C9-C10	-2.44	119.50	122.92
29	Y	611	CLA	CHA-C4D-ND	2.44	137.61	132.50
29	r	612	CLA	C1-C2-C3	-2.44	121.82	126.04
46	n	621	LUT	C10-C11-C12	-2.44	115.60	123.22
29	B	604	CLA	O2A-CGA-CBA	2.44	119.57	111.91
46	S	621	LUT	C3-C4-C5	-2.44	106.99	111.85
46	G	621	LUT	C38-C25-C24	-2.44	118.34	123.56
33	B	622	LMG	O8-C28-C29	2.44	119.56	111.91
29	B	616	CLA	CHA-C4D-ND	2.44	137.60	132.50
29	G	611	CLA	CHA-C4D-ND	2.44	137.60	132.50
48	n	623	NEX	C31-C30-C29	2.44	130.79	127.31
29	G	612	CLA	CHA-C4D-ND	2.44	137.60	132.50
29	S	609	CLA	CHA-C4D-ND	2.44	137.60	132.50
29	N	602	CLA	O2D-CGD-O1D	-2.44	119.07	123.84
29	S	614	CLA	O2D-CGD-O1D	-2.44	119.07	123.84
29	D	402	CLA	O2A-CGA-CBA	2.44	119.56	111.91
29	S	609	CLA	O2D-CGD-O1D	-2.44	119.07	123.84
29	Y	613	CLA	CHA-C4D-ND	2.44	137.60	132.50
29	Y	604	CLA	CMA-C3A-C4A	2.44	118.32	111.77
29	R	610	CLA	CMC-C2C-C1C	2.44	128.75	125.04
39	d	409	LHG	C5-O7-C7	-2.44	111.79	117.79
29	B	609	CLA	CMB-C2B-C1B	-2.44	124.72	128.46
29	b	605	CLA	CHD-C1D-ND	-2.44	122.22	124.45
29	S	610	CLA	CMA-C3A-C4A	2.44	118.32	111.77
29	d	403	CLA	CHA-C4D-ND	2.43	137.59	132.50
29	r	602	CLA	O2D-CGD-O1D	-2.43	119.08	123.84
29	s	614	CLA	CMD-C2D-C3D	-2.43	122.02	127.61

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
29	B	606	CLA	O2A-CGA-CBA	2.43	119.54	111.91
29	C	510	CLA	CHA-C4D-ND	2.43	137.59	132.50
29	r	613	CLA	CHA-C4D-ND	2.43	137.59	132.50
48	R	622	NEX	C40-C33-C34	-2.43	119.52	122.92
29	R	610	CLA	C1C-C2C-C3C	-2.43	104.40	106.96
45	N	605	CHL	CHB-C4A-NA	2.43	127.87	124.51
29	y	612	CLA	O2A-CGA-CBA	2.43	119.54	111.91
44	H	101	RRX	C8-C7-C6	-2.43	120.38	127.20
48	y	623	NEX	C4-C3-C2	2.43	115.47	110.77
29	N	603	CLA	OBD-CAD-C3D	-2.43	122.67	128.52
31	c	516	BCR	C38-C26-C27	2.43	118.28	113.62
29	N	613	CLA	CHA-C4D-ND	2.43	137.58	132.50
29	c	509	CLA	C2D-C1D-ND	2.43	111.89	110.10
46	R	620	LUT	C3-C4-C5	-2.43	107.01	111.85
29	Y	602	CLA	C1C-C2C-C3C	-2.43	104.40	106.96
29	r	604	CLA	O2D-CGD-O1D	-2.43	119.09	123.84
29	s	614	CLA	C2D-C1D-ND	2.43	111.89	110.10
29	n	603	CLA	CHA-C4D-ND	2.43	137.58	132.50
45	Y	609	CHL	CHB-C4A-NA	2.43	127.87	124.51
45	g	605	CHL	C4D-CHA-C1A	2.43	124.20	121.25
29	r	602	CLA	CHA-C4D-ND	2.43	137.58	132.50
29	y	613	CLA	C1-O2A-CGA	2.43	122.81	116.44
29	S	614	CLA	O2A-CGA-CBA	2.43	119.53	111.91
45	G	609	CHL	C4D-CHA-C1A	2.43	124.20	121.25
29	S	605	CLA	CHD-C1D-ND	-2.43	122.22	124.45
29	Y	610	CLA	O1D-CGD-CBD	-2.43	119.52	124.48
29	s	604	CLA	CMD-C2D-C3D	-2.42	122.04	127.61
29	r	612	CLA	CMA-C3A-C4A	2.42	118.29	111.77
29	b	603	CLA	CHA-C4D-ND	2.42	137.57	132.50
29	Y	604	CLA	CHA-C4D-ND	2.42	137.56	132.50
47	g	622	XAT	C19-C9-C10	-2.42	119.53	122.92
39	Y	624	LHG	C6-C5-C4	-2.42	106.06	111.79
29	B	605	CLA	CMD-C2D-C3D	-2.42	122.04	127.61
29	a	410	CLA	CHA-C4D-ND	2.42	137.56	132.50
33	b	622	LMG	O1-C7-C8	-2.42	105.06	110.90
29	n	602	CLA	CHA-C4D-ND	2.42	137.56	132.50
29	g	603	CLA	CHA-C4D-ND	2.42	137.56	132.50
29	n	614	CLA	CHA-C4D-ND	2.42	137.56	132.50
29	s	610	CLA	CMA-C3A-C4A	2.42	118.28	111.77
45	r	606	CHL	CHB-C4A-NA	2.42	127.86	124.51
29	r	603	CLA	CHA-C4D-ND	2.42	137.56	132.50
45	n	609	CHL	CAA-C2A-C1A	2.42	119.90	111.97

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
47	G	622	XAT	C19-C9-C10	-2.42	119.53	122.92
35	b	620	C7Z	C21-C26-C25	-2.42	119.21	122.61
29	b	604	CLA	O2A-CGA-CBA	2.42	119.50	111.91
29	R	611	CLA	CHA-C4D-ND	2.42	137.56	132.50
29	r	604	CLA	C1C-C2C-C3C	-2.42	104.42	106.96
46	n	621	LUT	C35-C15-C14	-2.42	118.52	123.47
48	R	622	NEX	C38-C25-C26	-2.42	118.21	122.26
29	c	502	CLA	CHA-C4D-ND	2.42	137.55	132.50
29	g	614	CLA	CHA-C4D-ND	2.42	137.55	132.50
29	G	602	CLA	CHA-C4D-ND	2.41	137.55	132.50
29	y	610	CLA	CHA-C4D-ND	2.41	137.55	132.50
33	h	102	LMG	O8-C28-C29	2.41	119.48	111.91
31	D	404	BCR	C38-C26-C27	2.41	118.25	113.62
29	c	503	CLA	CHA-C4D-ND	2.41	137.55	132.50
29	C	513	CLA	C1C-C2C-C3C	-2.41	104.42	106.96
29	s	609	CLA	O2D-CGD-O1D	-2.41	119.12	123.84
29	N	614	CLA	CHA-C4D-ND	2.41	137.55	132.50
29	S	617	CLA	CHA-C4D-ND	2.41	137.54	132.50
29	G	603	CLA	C1D-ND-C4D	-2.41	104.62	106.33
29	g	602	CLA	CHA-C4D-ND	2.41	137.54	132.50
29	c	509	CLA	O2A-CGA-CBA	2.41	119.47	111.91
29	C	508	CLA	C2D-C1D-ND	2.41	111.88	110.10
29	y	604	CLA	O2D-CGD-O1D	-2.41	119.13	123.84
45	n	608	CHL	CHB-C4A-NA	2.41	127.84	124.51
29	B	605	CLA	CHD-C1D-ND	-2.41	122.24	124.45
31	C	515	BCR	C38-C26-C25	-2.41	121.82	124.53
29	a	407	CLA	CHA-C4D-ND	2.41	137.53	132.50
29	b	604	CLA	CMB-C2B-C1B	-2.41	124.76	128.46
29	g	610	CLA	C1C-C2C-C3C	-2.41	104.43	106.96
46	g	621	LUT	C35-C15-C14	-2.41	118.54	123.47
29	b	604	CLA	O2D-CGD-O1D	-2.41	119.13	123.84
29	s	603	CLA	CHA-C4D-ND	2.41	137.53	132.50
29	D	403	CLA	CHA-C4D-ND	2.41	137.53	132.50
29	Y	614	CLA	CHA-C4D-ND	2.41	137.53	132.50
29	b	616	CLA	CHA-C4D-ND	2.41	137.53	132.50
29	b	617	CLA	O2D-CGD-O1D	-2.41	119.13	123.84
29	s	604	CLA	O2D-CGD-O1D	-2.41	119.13	123.84
29	B	602	CLA	C2D-C1D-ND	2.41	111.88	110.10
48	R	622	NEX	C39-C29-C30	-2.41	119.55	122.92
29	A	406	CLA	CHA-C4D-ND	2.41	137.53	132.50
29	c	501	CLA	CMD-C2D-C3D	-2.41	122.08	127.61
29	r	608	CLA	CMD-C2D-C3D	-2.40	122.08	127.61

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
45	G	601	CHL	C1-O2A-CGA	2.40	122.75	116.44
29	c	511	CLA	CMD-C2D-C3D	-2.40	122.08	127.61
29	y	611	CLA	CMA-C3A-C4A	2.40	118.23	111.77
29	s	610	CLA	C1C-C2C-C3C	-2.40	104.43	106.96
29	a	410	CLA	O2A-CGA-CBA	2.40	119.45	111.91
31	C	516	BCR	C38-C26-C27	2.40	118.23	113.62
29	N	610	CLA	CHA-C4D-ND	2.40	137.53	132.50
29	y	604	CLA	CHA-C4D-ND	2.40	137.53	132.50
29	A	407	CLA	CHA-C4D-ND	2.40	137.53	132.50
29	b	608	CLA	C1C-C2C-C3C	-2.40	104.43	106.96
29	Y	611	CLA	CHD-C1D-ND	-2.40	122.25	124.45
29	C	506	CLA	CHA-C4D-ND	2.40	137.52	132.50
29	N	602	CLA	CHA-C4D-ND	2.40	137.52	132.50
29	B	603	CLA	O2A-CGA-CBA	2.40	119.44	111.91
46	N	621	LUT	C18-C5-C6	-2.40	121.83	124.53
45	s	608	CHL	CHB-C4A-NA	2.40	127.83	124.51
29	B	610	CLA	CHA-C4D-ND	2.40	137.52	132.50
29	b	606	CLA	O2A-CGA-CBA	2.40	119.44	111.91
45	y	606	CHL	C1-O2A-CGA	2.40	122.74	116.44
29	C	513	CLA	CHA-C4D-ND	2.40	137.52	132.50
29	Y	610	CLA	CHA-C4D-ND	2.40	137.52	132.50
29	a	406	CLA	O2A-CGA-CBA	2.40	119.43	111.91
45	g	606	CHL	CHB-C4A-NA	2.40	127.83	124.51
29	g	614	CLA	O2D-CGD-O1D	-2.40	119.15	123.84
29	B	606	CLA	C2D-C1D-ND	2.40	111.87	110.10
31	d	404	BCR	C4-C5-C6	-2.40	119.25	122.73
29	S	602	CLA	CHA-C4D-ND	2.40	137.51	132.50
48	S	622	NEX	C40-C33-C34	-2.40	119.56	122.92
29	C	502	CLA	C1C-C2C-C3C	-2.40	104.44	106.96
29	c	513	CLA	CHA-C4D-ND	2.40	137.51	132.50
45	n	608	CHL	C1-O2A-CGA	2.40	122.73	116.44
29	c	506	CLA	CHA-C4D-ND	2.40	137.51	132.50
29	s	609	CLA	CHA-C4D-ND	2.39	137.51	132.50
29	s	617	CLA	O2A-CGA-CBA	2.39	119.42	111.91
48	Y	623	NEX	C16-C1-C6	-2.39	108.33	110.47
29	C	503	CLA	CHA-C4D-ND	2.39	137.51	132.50
45	G	605	CHL	CHB-C4A-NA	2.39	127.82	124.51
29	y	602	CLA	CMB-C2B-C3B	2.39	129.16	124.68
29	D	402	CLA	C1-C2-C3	-2.39	121.90	126.04
29	C	512	CLA	CMA-C3A-C4A	2.39	118.20	111.77
29	A	407	CLA	CAA-C2A-C3A	-2.39	106.23	112.78
29	s	602	CLA	O2D-CGD-O1D	-2.39	119.16	123.84

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
29	G	614	CLA	CMD-C2D-C3D	-2.39	122.11	127.61
29	C	502	CLA	CHA-C4D-ND	2.39	137.50	132.50
29	n	612	CLA	CHA-C4D-ND	2.39	137.50	132.50
29	s	617	CLA	CHA-C4D-ND	2.39	137.50	132.50
29	a	406	CLA	CHA-C4D-ND	2.39	137.50	132.50
29	r	608	CLA	O2D-CGD-O1D	-2.39	119.16	123.84
29	y	613	CLA	CHA-C4D-ND	2.39	137.50	132.50
45	y	606	CHL	CHB-C4A-NA	2.39	127.82	124.51
29	Y	604	CLA	O2D-CGD-O1D	-2.39	119.17	123.84
29	S	604	CLA	CHA-C4D-ND	2.39	137.50	132.50
29	G	610	CLA	C1D-ND-C4D	-2.39	104.64	106.33
29	g	610	CLA	CHA-C4D-ND	2.39	137.50	132.50
31	C	516	BCR	C31-C1-C6	-2.39	106.43	110.30
29	B	612	CLA	CMA-C3A-C4A	2.39	118.19	111.77
29	b	610	CLA	CHA-C4D-ND	2.38	137.49	132.50
29	R	604	CLA	C1C-C2C-C3C	-2.38	104.45	106.96
29	C	511	CLA	CHA-C1A-NA	-2.38	120.94	126.40
29	Y	603	CLA	C1-C2-C3	-2.38	121.92	126.04
29	C	503	CLA	CMD-C2D-C3D	-2.38	122.13	127.61
46	R	620	LUT	C11-C12-C13	-2.38	119.72	126.42
31	b	618	BCR	C33-C5-C6	-2.38	121.85	124.53
29	a	405	CLA	CHA-C4D-ND	2.38	137.49	132.50
29	d	402	CLA	CHA-C4D-ND	2.38	137.49	132.50
31	B	619	BCR	C8-C7-C6	-2.38	120.51	127.20
45	G	609	CHL	CHB-C4A-NA	2.38	127.81	124.51
29	g	602	CLA	O2A-CGA-CBA	2.38	119.38	111.91
29	y	610	CLA	CMA-C3A-C4A	2.38	118.17	111.77
29	R	610	CLA	C1D-ND-C4D	-2.38	104.64	106.33
29	G	603	CLA	CHA-C4D-ND	2.38	137.48	132.50
29	C	509	CLA	CBC-CAC-C3C	-2.38	105.87	112.43
29	Y	602	CLA	CAA-C2A-C3A	-2.38	106.26	112.78
31	c	517	BCR	C3-C4-C5	-2.38	109.83	114.08
36	b	623	DGA	OG1-CA1-CA2	2.38	119.38	111.91
45	N	608	CHL	C4A-NA-C1A	2.38	107.78	106.71
29	s	609	CLA	C1C-C2C-C3C	-2.38	104.45	106.96
45	S	608	CHL	C1-C2-C3	-2.38	121.93	126.04
31	b	618	BCR	C3-C4-C5	-2.38	109.83	114.08
31	C	514	BCR	C33-C5-C4	2.38	118.19	113.62
29	S	610	CLA	C1C-C2C-C3C	-2.38	104.46	106.96
46	S	621	LUT	C15-C14-C13	-2.38	123.92	127.31
29	b	608	CLA	O2A-CGA-CBA	2.38	119.37	111.91
45	n	606	CHL	C1-O2A-CGA	2.38	122.68	116.44

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
45	G	607	CHL	C1-O2A-CGA	2.38	122.68	116.44
45	S	608	CHL	C1B-CHB-C4A	-2.37	125.41	130.12
29	B	610	CLA	O2A-CGA-CBA	2.37	119.36	111.91
29	Y	608	CLA	O2D-CGD-O1D	-2.37	119.20	123.84
29	b	612	CLA	CHA-C4D-ND	2.37	137.47	132.50
29	b	612	CLA	C2D-C1D-ND	2.37	111.85	110.10
42	D	405	PL9	C20-C19-C21	2.37	119.26	115.27
45	n	608	CHL	C4D-CHA-C1A	2.37	124.14	121.25
48	y	623	NEX	C19-C9-C10	-2.37	119.60	122.92
45	S	608	CHL	CHB-C4A-NA	2.37	127.79	124.51
29	c	513	CLA	C1C-C2C-C3C	-2.37	104.46	106.96
29	S	604	CLA	O2A-CGA-CBA	2.37	119.35	111.91
45	s	608	CHL	C1-O2A-CGA	2.37	122.67	116.44
45	Y	606	CHL	CHB-C4A-NA	2.37	127.79	124.51
29	A	410	CLA	CHA-C4D-ND	2.37	137.46	132.50
29	c	506	CLA	C2D-C1D-ND	2.37	111.85	110.10
29	s	604	CLA	C2D-C1D-ND	2.37	111.85	110.10
45	Y	606	CHL	C1-C2-C3	-2.37	121.95	126.04
29	C	504	CLA	CHA-C4D-ND	2.37	137.45	132.50
29	n	610	CLA	CHA-C4D-ND	2.37	137.45	132.50
39	d	408	LHG	C6-C5-C4	-2.37	106.19	111.79
29	S	602	CLA	O2D-CGD-O1D	-2.37	119.21	123.84
45	n	601	CHL	C1-O2A-CGA	2.37	122.65	116.44
45	G	606	CHL	C1B-CHB-C4A	-2.36	125.44	130.12
46	N	621	LUT	C2-C3-C4	-2.36	107.07	110.30
29	r	610	CLA	CHA-C4D-ND	2.36	137.44	132.50
29	R	603	CLA	O1D-CGD-CBD	-2.36	119.65	124.48
29	G	603	CLA	O2D-CGD-O1D	-2.36	119.22	123.84
29	C	506	CLA	CMB-C2B-C1B	-2.36	124.83	128.46
29	N	603	CLA	CAA-C2A-C3A	-2.36	106.31	112.78
31	d	404	BCR	C27-C26-C25	-2.36	119.30	122.73
29	R	610	CLA	CHA-C4D-ND	2.36	137.44	132.50
46	Y	621	LUT	C10-C11-C12	-2.36	115.85	123.22
29	a	405	CLA	C2C-C1C-NC	2.36	112.18	109.97
29	B	613	CLA	C1C-C2C-C3C	-2.36	104.47	106.96
29	b	617	CLA	CHA-C4D-ND	2.36	137.44	132.50
29	R	603	CLA	CMB-C2B-C3B	2.36	129.09	124.68
29	y	603	CLA	CAA-C2A-C3A	-2.36	106.31	112.78
47	n	622	XAT	C39-C29-C30	-2.36	119.62	122.92
48	n	623	NEX	C40-C33-C34	-2.36	119.62	122.92
29	B	610	CLA	C1C-C2C-C3C	-2.36	104.48	106.96
29	A	405	CLA	CHA-C4D-ND	2.36	137.43	132.50

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
29	R	612	CLA	CMD-C2D-C3D	-2.36	122.19	127.61
29	c	505	CLA	O2A-CGA-CBA	2.36	119.31	111.91
47	r	622	XAT	C31-C32-C33	2.36	133.04	126.42
29	Y	603	CLA	O2A-CGA-CBA	2.36	119.31	111.91
29	C	501	CLA	C1D-ND-C4D	-2.36	104.66	106.33
46	S	621	LUT	C16-C1-C6	-2.36	106.48	110.30
29	s	609	CLA	CAA-C2A-C3A	-2.36	106.33	112.78
29	R	609	CLA	C2D-C1D-ND	2.36	111.84	110.10
45	r	607	CHL	C4A-NA-C1A	2.36	107.77	106.71
29	S	617	CLA	O2A-CGA-CBA	2.36	119.30	111.91
29	Y	613	CLA	O2D-CGD-O1D	-2.35	119.23	123.84
29	S	604	CLA	CMD-C2D-C3D	-2.35	122.20	127.61
29	b	602	CLA	C2D-C1D-ND	2.35	111.84	110.10
29	r	613	CLA	C1C-C2C-C3C	-2.35	104.48	106.96
29	Y	602	CLA	CHA-C4D-ND	2.35	137.42	132.50
47	r	622	XAT	C30-C31-C32	-2.35	115.87	123.22
29	b	607	CLA	CMB-C2B-C1B	-2.35	124.85	128.46
29	G	614	CLA	O2D-CGD-O1D	-2.35	119.24	123.84
29	g	614	CLA	CMD-C2D-C3D	-2.35	122.20	127.61
29	c	505	CLA	C2D-C1D-ND	2.35	111.84	110.10
29	R	610	CLA	O1D-CGD-CBD	-2.35	119.67	124.48
29	c	504	CLA	CHA-C4D-ND	2.35	137.42	132.50
29	b	610	CLA	O2A-CGA-CBA	2.35	119.28	111.91
48	N	623	NEX	C1-C2-C3	2.35	118.95	113.64
29	G	602	CLA	C1C-C2C-C3C	-2.35	104.49	106.96
29	B	617	CLA	CHA-C4D-ND	2.35	137.41	132.50
29	B	617	CLA	CHD-C1D-ND	-2.35	122.30	124.45
29	y	608	CLA	O2D-CGD-O1D	-2.35	119.25	123.84
45	Y	609	CHL	CHD-C4C-C3C	2.35	128.29	124.84
45	n	609	CHL	C4A-NA-C1A	2.35	107.76	106.71
29	y	604	CLA	CMD-C2D-C3D	-2.35	122.21	127.61
29	n	602	CLA	O2D-CGD-O1D	-2.35	119.25	123.84
45	N	608	CHL	C1B-CHB-C4A	-2.35	125.47	130.12
29	c	506	CLA	CHA-C1A-NA	-2.35	121.02	126.40
43	f	101	HEM	CBA-CAA-C2A	-2.35	108.61	112.62
29	b	603	CLA	C1-O2A-CGA	2.35	122.60	116.44
29	G	610	CLA	O2A-CGA-CBA	2.35	119.27	111.91
48	y	623	NEX	C5-C4-C3	2.35	114.52	111.75
29	C	503	CLA	O2A-CGA-CBA	2.35	119.27	111.91
29	b	605	CLA	CMD-C2D-C3D	-2.35	122.22	127.61
29	R	611	CLA	O2D-CGD-O1D	-2.35	119.25	123.84
29	B	608	CLA	O2A-CGA-CBA	2.34	119.27	111.91

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
45	S	606	CHL	CHB-C4A-NA	2.34	127.75	124.51
47	y	622	XAT	C18-C5-C6	-2.34	118.33	122.26
29	A	406	CLA	C2D-C1D-ND	2.34	111.83	110.10
46	y	621	LUT	C10-C11-C12	-2.34	115.91	123.22
45	y	609	CHL	CHB-C4A-NA	2.34	127.75	124.51
45	g	605	CHL	CHB-C4A-NA	2.34	127.75	124.51
29	B	611	CLA	O2A-CGA-CBA	2.34	119.26	111.91
29	C	506	CLA	CHA-C1A-NA	-2.34	121.03	126.40
29	B	607	CLA	CMD-C2D-C3D	-2.34	122.23	127.61
29	B	608	CLA	C1C-C2C-C3C	-2.34	104.50	106.96
29	s	602	CLA	CHA-C4D-ND	2.34	137.40	132.50
45	N	606	CHL	CHB-C4A-NA	2.34	127.75	124.51
46	n	620	LUT	C38-C25-C24	-2.34	118.55	123.56
29	R	609	CLA	CHA-C1A-NA	-2.34	121.04	126.40
29	r	609	CLA	CHA-C1A-NA	-2.34	121.04	126.40
29	c	509	CLA	CMD-C2D-C3D	-2.34	122.23	127.61
29	g	613	CLA	CMD-C2D-C3D	-2.34	122.23	127.61
47	N	622	XAT	C19-C9-C10	-2.34	119.65	122.92
29	B	612	CLA	O2A-CGA-CBA	2.34	119.25	111.91
45	N	608	CHL	C4D-CHA-C1A	2.34	124.09	121.25
47	y	622	XAT	O24-C25-C38	-2.34	112.25	115.06
29	a	406	CLA	CMB-C2B-C1B	-2.34	124.87	128.46
33	B	622	LMG	O1-C7-C8	-2.34	105.26	110.90
31	B	618	BCR	C2-C1-C6	2.34	114.08	110.48
29	a	410	CLA	CMB-C2B-C3B	2.34	129.05	124.68
29	C	508	CLA	CHA-C4D-ND	2.34	137.39	132.50
29	C	501	CLA	O2A-CGA-CBA	2.34	119.24	111.91
29	Y	603	CLA	C1D-ND-C4D	-2.34	104.68	106.33
29	b	614	CLA	CMB-C2B-C3B	2.33	129.05	124.68
29	B	612	CLA	C1C-C2C-C3C	-2.33	104.50	106.96
46	g	620	LUT	C10-C11-C12	-2.33	115.93	123.22
46	y	620	LUT	C30-C31-C32	-2.33	115.93	123.22
46	s	620	LUT	C31-C30-C29	-2.33	123.98	127.31
30	a	409	PHO	O1D-CGD-CBD	2.33	128.62	124.74
29	y	612	CLA	CMA-C3A-C4A	2.33	118.04	111.77
33	H	102	LMG	O8-C28-C29	2.33	119.23	111.91
29	C	507	CLA	CHA-C1A-NA	-2.33	121.06	126.40
29	a	407	CLA	CAA-C2A-C3A	-2.33	106.39	112.78
29	c	507	CLA	CHA-C1A-NA	-2.33	121.06	126.40
29	R	603	CLA	CMD-C2D-C3D	-2.33	122.26	127.61
29	s	603	CLA	CHA-C1A-NA	-2.33	121.07	126.40
29	y	613	CLA	O2D-CGD-O1D	-2.33	119.29	123.84

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
29	B	604	CLA	C2D-C1D-ND	2.33	111.82	110.10
29	N	604	CLA	CMA-C3A-C4A	2.33	118.02	111.77
29	c	503	CLA	O2D-CGD-O1D	-2.33	119.29	123.84
29	y	602	CLA	C1C-C2C-C3C	-2.32	104.51	106.96
45	G	607	CHL	CMB-C2B-C1B	-2.32	124.89	128.46
29	C	511	CLA	CMB-C2B-C3B	2.32	129.03	124.68
46	g	621	LUT	C38-C25-C24	-2.32	118.59	123.56
45	s	606	CHL	CHD-C4C-C3C	2.32	128.26	124.84
31	D	404	BCR	C1-C6-C5	-2.32	119.34	122.61
29	r	609	CLA	O2D-CGD-O1D	-2.32	119.30	123.84
48	r	623	NEX	C38-C25-C26	-2.32	118.37	122.26
47	y	622	XAT	C40-C33-C34	-2.32	119.67	122.92
29	d	402	CLA	O2D-CGD-O1D	-2.32	119.30	123.84
29	A	410	CLA	O2A-CGA-CBA	2.32	119.19	111.91
45	n	608	CHL	CMB-C2B-C1B	-2.32	124.90	128.46
29	r	604	CLA	C1D-ND-C4D	-2.32	104.69	106.33
29	c	512	CLA	CHA-C4D-ND	2.32	137.35	132.50
31	D	404	BCR	C33-C5-C4	2.32	118.07	113.62
29	b	607	CLA	C2D-C1D-ND	2.32	111.81	110.10
29	s	603	CLA	O2D-CGD-O1D	-2.32	119.31	123.84
35	B	620	C7Z	C30-C31-C32	-2.32	115.99	123.22
29	S	611	CLA	O2A-CGA-CBA	2.32	119.17	111.91
31	B	618	BCR	C33-C5-C6	-2.31	121.93	124.53
29	b	613	CLA	CMB-C2B-C3B	2.31	129.01	124.68
45	R	607	CHL	CHB-C4A-NA	2.31	127.71	124.51
45	G	601	CHL	C1-C2-C3	-2.31	122.04	126.04
29	r	602	CLA	O2A-CGA-CBA	2.31	119.17	111.91
29	c	511	CLA	CHA-C1A-NA	-2.31	121.10	126.40
46	N	621	LUT	C35-C15-C14	-2.31	118.73	123.47
46	y	620	LUT	C39-C29-C28	2.31	121.72	118.08
29	B	611	CLA	CHA-C1A-NA	-2.31	121.10	126.40
45	g	607	CHL	CMB-C2B-C1B	-2.31	124.91	128.46
29	b	609	CLA	CMB-C2B-C3B	2.31	129.00	124.68
29	Y	604	CLA	CMD-C2D-C3D	-2.31	122.30	127.61
31	c	514	BCR	C33-C5-C4	2.31	118.06	113.62
29	Y	608	CLA	O2A-CGA-CBA	2.31	119.16	111.91
30	A	408	PHO	O1D-CGD-CBD	2.31	128.59	124.74
32	C	526	SQD	O3-C3-C2	-2.31	105.01	110.35
29	c	507	CLA	CMA-C3A-C4A	2.31	117.98	111.77
29	r	611	CLA	CMD-C2D-C3D	-2.31	122.30	127.61
29	B	615	CLA	C2D-C1D-ND	2.31	111.81	110.10
29	A	406	CLA	CMD-C2D-C3D	-2.31	122.30	127.61

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
29	C	510	CLA	O2A-CGA-CBA	2.31	119.15	111.91
29	r	611	CLA	O2D-CGD-O1D	-2.31	119.33	123.84
29	G	610	CLA	O1D-CGD-CBD	-2.31	119.76	124.48
29	c	510	CLA	O2A-CGA-CBA	2.31	119.15	111.91
45	n	609	CHL	CMB-C2B-C1B	-2.31	124.92	128.46
29	c	508	CLA	CHA-C4D-ND	2.31	137.32	132.50
48	s	623	NEX	C26-C27-C28	-2.31	121.12	125.99
45	g	606	CHL	CMB-C2B-C1B	-2.31	124.92	128.46
29	G	610	CLA	CHA-C4D-ND	2.31	137.32	132.50
29	g	603	CLA	CMD-C2D-C3D	-2.30	122.31	127.61
45	s	608	CHL	C1-C2-C3	-2.30	122.06	126.04
31	b	619	BCR	C19-C18-C17	2.30	122.47	118.94
29	R	608	CLA	CMD-C2D-C3D	-2.30	122.32	127.61
44	h	101	RRX	C38-C26-C25	-2.30	121.94	124.53
31	C	514	BCR	C34-C9-C10	-2.30	119.70	122.92
46	s	620	LUT	C39-C29-C28	2.30	121.70	118.08
29	g	613	CLA	O2D-CGD-O1D	-2.30	119.34	123.84
29	R	604	CLA	C1D-ND-C4D	-2.30	104.70	106.33
42	d	405	PL9	C20-C19-C21	2.30	119.14	115.27
42	d	405	PL9	C27-C28-C29	-2.30	122.12	127.66
29	A	405	CLA	C2C-C1C-NC	2.30	112.13	109.97
29	n	604	CLA	CMD-C2D-C3D	-2.30	122.33	127.61
48	N	623	NEX	C4-C3-C2	2.30	115.21	110.77
46	s	620	LUT	C31-C32-C33	-2.30	119.96	126.42
29	a	405	CLA	CMA-C3A-C4A	2.30	117.95	111.77
29	S	603	CLA	CHA-C1A-NA	-2.30	121.14	126.40
45	N	607	CHL	CMB-C2B-C1B	-2.30	124.94	128.46
45	y	601	CHL	CMB-C2B-C1B	-2.30	124.94	128.46
29	d	402	CLA	CAC-C3C-C4C	2.30	127.79	124.81
29	N	604	CLA	CMB-C2B-C3B	2.30	128.97	124.68
32	A	412	SQD	O3-C3-C2	-2.29	105.05	110.35
29	N	610	CLA	CMB-C2B-C3B	2.29	128.97	124.68
29	n	602	CLA	CMB-C2B-C3B	2.29	128.97	124.68
45	y	607	CHL	CMB-C2B-C1B	-2.29	124.94	128.46
46	G	620	LUT	C38-C25-C24	-2.29	118.66	123.56
29	c	502	CLA	CMD-C2D-C3D	-2.29	122.35	127.61
45	n	607	CHL	CMB-C2B-C1B	-2.29	124.95	128.46
45	y	605	CHL	CHB-C4A-NA	2.29	127.68	124.51
29	r	613	CLA	CMD-C2D-C3D	-2.29	122.35	127.61
45	Y	607	CHL	CMB-C2B-C1B	-2.29	124.95	128.46
45	g	606	CHL	C4D-CHA-C1A	2.29	124.03	121.25
46	Y	620	LUT	C35-C34-C33	-2.29	124.05	127.31

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
29	y	603	CLA	O1D-CGD-CBD	-2.29	119.81	124.48
29	C	512	CLA	CHA-C4D-ND	2.29	137.28	132.50
46	y	621	LUT	C16-C1-C6	-2.28	106.59	110.30
29	n	603	CLA	C1D-ND-C4D	-2.28	104.71	106.33
31	C	517	BCR	C34-C9-C10	-2.28	119.72	122.92
29	a	406	CLA	CMD-C2D-C3D	-2.28	122.36	127.61
32	c	626	SQD	O3-C3-C2	-2.28	105.07	110.35
45	N	609	CHL	CMB-C2B-C1B	-2.28	124.95	128.46
29	C	509	CLA	CMD-C2D-C3D	-2.28	122.36	127.61
45	N	608	CHL	CMB-C2B-C1B	-2.28	124.95	128.46
45	Y	601	CHL	CMB-C2B-C1B	-2.28	124.95	128.46
47	g	622	XAT	O24-C25-C24	2.28	115.10	113.38
29	N	602	CLA	CMB-C2B-C3B	2.28	128.95	124.68
29	d	402	CLA	C2C-C1C-NC	2.28	112.11	109.97
29	r	613	CLA	CMA-C3A-C4A	2.28	117.91	111.77
29	G	602	CLA	C1D-ND-C4D	-2.28	104.71	106.33
29	B	613	CLA	C1-C2-C3	-2.28	122.10	126.04
29	A	405	CLA	C1-O2A-CGA	2.28	122.43	116.44
29	G	610	CLA	C1C-C2C-C3C	-2.28	104.56	106.96
29	R	604	CLA	CMB-C2B-C3B	2.28	128.94	124.68
29	C	503	CLA	C1D-ND-C4D	-2.28	104.72	106.33
29	c	510	CLA	C1D-ND-C4D	-2.28	104.72	106.33
45	n	609	CHL	CHD-C4C-C3C	2.28	128.19	124.84
29	c	503	CLA	O2A-CGA-CBA	2.28	119.06	111.91
45	n	606	CHL	C4D-CHA-C1A	2.28	124.02	121.25
29	S	603	CLA	O2D-CGD-O1D	-2.28	119.38	123.84
29	N	613	CLA	O2D-CGD-O1D	-2.28	119.38	123.84
29	B	616	CLA	CHA-C1A-NA	-2.28	121.18	126.40
46	S	621	LUT	C11-C10-C9	-2.28	124.06	127.31
29	n	610	CLA	C1D-ND-C4D	-2.28	104.72	106.33
29	D	402	CLA	O2D-CGD-O1D	-2.28	119.39	123.84
45	y	606	CHL	C1-C2-C3	-2.28	122.11	126.04
29	B	614	CLA	CMB-C2B-C3B	2.28	128.94	124.68
29	y	602	CLA	C1D-ND-C4D	-2.28	104.72	106.33
45	Y	606	CHL	C1-O2A-CGA	2.28	122.42	116.44
29	r	603	CLA	CMD-C2D-C3D	-2.28	122.38	127.61
29	B	612	CLA	CHA-C1A-NA	-2.28	121.19	126.40
45	G	605	CHL	C4D-CHA-C1A	2.28	124.02	121.25
48	N	623	NEX	C19-C9-C10	-2.28	119.74	122.92
29	Y	612	CLA	CHA-C1A-NA	-2.27	121.19	126.40
31	c	516	BCR	C31-C1-C6	-2.27	106.61	110.30
29	c	510	CLA	CAA-C2A-C3A	-2.27	106.55	112.78

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
29	N	604	CLA	O2A-CGA-CBA	2.27	119.04	111.91
29	B	607	CLA	CMB-C2B-C1B	-2.27	124.97	128.46
31	b	618	BCR	C34-C9-C10	-2.27	119.74	122.92
29	C	504	CLA	O2A-CGA-CBA	2.27	119.04	111.91
39	n	624	LHG	O8-C23-C24	2.27	119.04	111.91
29	n	612	CLA	CHA-C1A-NA	-2.27	121.19	126.40
35	B	620	C7Z	C7-C8-C9	-2.27	122.80	126.23
29	C	501	CLA	O1D-CGD-CBD	-2.27	119.83	124.48
31	C	516	BCR	C37-C22-C23	2.27	121.66	118.08
29	s	611	CLA	O2A-CGA-CBA	2.27	119.04	111.91
29	C	504	CLA	CMB-C2B-C3B	2.27	128.93	124.68
45	n	606	CHL	C1B-CHB-C4A	-2.27	125.62	130.12
45	N	608	CHL	C1-O2A-CGA	2.27	122.40	116.44
29	B	614	CLA	C1D-ND-C4D	-2.27	104.72	106.33
29	b	611	CLA	CHA-C1A-NA	-2.27	121.20	126.40
29	N	604	CLA	OBD-CAD-C3D	-2.27	123.06	128.52
29	y	613	CLA	C1D-ND-C4D	-2.27	104.72	106.33
42	d	405	PL9	C31-C32-C33	-2.27	104.42	111.88
29	C	507	CLA	C1-C2-C3	-2.27	122.12	126.04
29	y	614	CLA	O2A-CGA-CBA	2.27	119.03	111.91
29	B	611	CLA	CMA-C3A-C4A	2.27	117.87	111.77
29	b	616	CLA	CHA-C1A-NA	-2.27	121.20	126.40
48	N	623	NEX	C40-C33-C34	-2.27	119.75	122.92
45	N	609	CHL	CHB-C4A-NA	2.27	127.65	124.51
29	g	614	CLA	O2A-CGA-CBA	2.27	119.02	111.91
29	g	610	CLA	CMD-C2D-C3D	-2.27	122.40	127.61
45	y	609	CHL	CHD-C4C-C3C	2.27	128.17	124.84
29	y	604	CLA	CMA-C3A-C4A	2.27	117.86	111.77
31	c	517	BCR	C2-C1-C6	2.27	113.97	110.48
29	R	603	CLA	O2A-CGA-CBA	2.27	119.02	111.91
42	d	405	PL9	O1-C4-C3	-2.27	118.22	120.72
29	b	607	CLA	CMD-C2D-C3D	-2.27	122.40	127.61
29	R	602	CLA	C1D-ND-C4D	-2.26	104.73	106.33
29	S	602	CLA	C1D-ND-C4D	-2.26	104.73	106.33
29	Y	602	CLA	C1D-ND-C4D	-2.26	104.73	106.33
48	g	623	NEX	C1-C2-C3	2.26	118.76	113.64
29	c	512	CLA	O2A-CGA-CBA	2.26	119.01	111.91
45	g	606	CHL	C1B-CHB-C4A	-2.26	125.63	130.12
29	y	610	CLA	O1D-CGD-CBD	-2.26	119.85	124.48
45	g	605	CHL	C1B-CHB-C4A	-2.26	125.64	130.12
32	B	621	SQD	O3-C3-C2	-2.26	105.12	110.35
39	d	410	LHG	C5-O7-C7	-2.26	112.22	117.79

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
29	B	603	CLA	O2D-CGD-O1D	-2.26	119.42	123.84
29	G	604	CLA	CMD-C2D-C3D	-2.26	122.42	127.61
29	b	604	CLA	CMD-C2D-C3D	-2.26	122.42	127.61
29	s	617	CLA	C1D-ND-C4D	-2.26	104.73	106.33
29	g	602	CLA	C1C-C2C-C3C	-2.26	104.58	106.96
31	c	517	BCR	C15-C14-C13	-2.26	124.09	127.31
32	a	412	SQD	O3-C3-C2	-2.26	105.13	110.35
29	Y	610	CLA	C1C-C2C-C3C	-2.26	104.58	106.96
29	C	510	CLA	CAA-C2A-C3A	-2.26	106.60	112.78
29	G	613	CLA	CMD-C2D-C3D	-2.26	122.42	127.61
29	Y	614	CLA	CMD-C2D-C3D	-2.26	122.42	127.61
29	c	501	CLA	C1D-ND-C4D	-2.26	104.73	106.33
47	g	622	XAT	C18-C5-C6	-2.26	118.48	122.26
29	r	610	CLA	C3D-C2D-C1D	-2.26	102.75	105.83
29	R	604	CLA	CMD-C2D-C3D	-2.26	122.42	127.61
29	A	406	CLA	O2A-CGA-CBA	2.25	118.98	111.91
29	d	403	CLA	CAA-C2A-C3A	-2.25	106.61	112.78
29	R	609	CLA	O2D-CGD-O1D	-2.25	119.43	123.84
29	c	512	CLA	O2D-CGD-O1D	-2.25	119.43	123.84
45	n	606	CHL	CHB-C4A-NA	2.25	127.63	124.51
29	B	602	CLA	O2A-CGA-CBA	2.25	118.98	111.91
48	Y	623	NEX	C1-C2-C3	2.25	118.73	113.64
29	r	612	CLA	CHA-C1A-NA	-2.25	121.24	126.40
29	B	603	CLA	C2D-C1D-ND	2.25	111.76	110.10
29	C	507	CLA	CMA-C3A-C4A	2.25	117.83	111.77
29	g	604	CLA	O2A-CGA-CBA	2.25	118.97	111.91
45	g	601	CHL	C1-C2-C3	-2.25	122.15	126.04
48	y	623	NEX	C31-C30-C29	2.25	130.52	127.31
29	c	503	CLA	CHA-C1A-NA	-2.25	121.24	126.40
39	D	408	LHG	C6-C5-C4	-2.25	106.47	111.79
29	C	505	CLA	C2D-C1D-ND	2.25	111.76	110.10
29	r	603	CLA	O2A-CGA-CBA	2.25	118.97	111.91
45	S	608	CHL	C4D-CHA-C1A	2.25	123.99	121.25
48	g	623	NEX	C16-C1-C6	-2.25	108.46	110.47
29	C	507	CLA	C1-O2A-CGA	2.25	122.34	116.44
29	r	613	CLA	O1D-CGD-CBD	-2.25	119.88	124.48
45	G	608	CHL	C4A-NA-C1A	2.25	107.72	106.71
45	y	605	CHL	C1B-CHB-C4A	-2.25	125.66	130.12
45	g	607	CHL	C4D-CHA-C1A	2.25	123.98	121.25
29	y	603	CLA	C1-C2-C3	-2.25	122.16	126.04
30	a	408	PHO	C1-C2-C3	-2.25	122.16	126.04
42	D	405	PL9	C31-C32-C33	-2.25	104.49	111.88

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
48	y	623	NEX	C20-C13-C14	-2.25	119.77	122.92
45	g	607	CHL	C1-O2A-CGA	2.25	122.34	116.44
46	s	621	LUT	C10-C11-C12	-2.25	116.20	123.22
46	y	620	LUT	C11-C10-C9	-2.25	124.10	127.31
29	G	614	CLA	O2A-CGA-CBA	2.25	118.96	111.91
29	R	608	CLA	O2D-CGD-O1D	-2.25	119.45	123.84
46	n	621	LUT	C8-C7-C6	-2.25	120.89	127.20
29	s	612	CLA	C1D-ND-C4D	-2.25	104.74	106.33
29	C	509	CLA	O2A-CGA-CBA	2.24	118.95	111.91
29	b	608	CLA	C1-C2-C3	-2.24	122.16	126.04
29	s	614	CLA	CHA-C1A-NA	-2.24	121.26	126.40
29	c	501	CLA	O2A-CGA-CBA	2.24	118.95	111.91
29	R	602	CLA	O1D-CGD-CBD	-2.24	119.89	124.48
46	n	620	LUT	C7-C8-C9	-2.24	122.84	126.23
45	N	609	CHL	C4D-CHA-C1A	2.24	123.98	121.25
45	G	605	CHL	C4A-NA-C1A	2.24	107.71	106.71
29	n	604	CLA	O2A-CGA-CBA	2.24	118.94	111.91
29	c	512	CLA	CMA-C3A-C4A	2.24	117.80	111.77
46	s	621	LUT	C11-C10-C9	-2.24	124.11	127.31
29	c	504	CLA	O2A-CGA-CBA	2.24	118.94	111.91
29	Y	614	CLA	O2A-CGA-CBA	2.24	118.94	111.91
29	R	603	CLA	CHA-C1A-NA	-2.24	121.27	126.40
29	g	602	CLA	CMB-C2B-C3B	2.24	128.87	124.68
29	N	612	CLA	O2D-CGD-O1D	-2.24	119.46	123.84
47	n	622	XAT	C19-C9-C10	-2.24	119.79	122.92
48	r	623	NEX	C40-C33-C34	-2.24	119.79	122.92
46	Y	621	LUT	C35-C34-C33	-2.24	124.12	127.31
29	g	613	CLA	C1-O2A-CGA	2.24	122.31	116.44
29	b	611	CLA	O2A-CGA-CBA	2.24	118.93	111.91
29	g	604	CLA	CMD-C2D-C3D	-2.24	122.47	127.61
29	b	605	CLA	O2A-CGA-CBA	2.23	118.92	111.91
45	n	609	CHL	C4D-CHA-C1A	2.23	123.97	121.25
39	y	624	LHG	C6-C5-C4	-2.23	106.50	111.79
32	b	621	SQD	O3-C3-C2	-2.23	105.19	110.35
45	Y	605	CHL	CHB-C4A-NA	2.23	127.60	124.51
29	B	613	CLA	O2A-CGA-CBA	2.23	118.91	111.91
29	n	613	CLA	O2D-CGD-O1D	-2.23	119.47	123.84
48	n	623	NEX	C38-C25-C26	-2.23	118.52	122.26
47	N	622	XAT	C39-C29-C30	-2.23	119.80	122.92
29	n	610	CLA	O1D-CGD-CBD	-2.23	119.92	124.48
29	R	608	CLA	CHA-C1A-NA	-2.23	121.29	126.40
31	D	404	BCR	C27-C26-C25	-2.23	119.49	122.73

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
29	n	614	CLA	CHA-C1A-NA	-2.23	121.29	126.40
29	A	405	CLA	CMD-C2D-C3D	-2.23	122.48	127.61
29	R	613	CLA	CHA-C1A-NA	-2.23	121.29	126.40
31	B	618	BCR	C33-C5-C4	2.23	117.90	113.62
31	b	618	BCR	C2-C1-C6	2.23	113.91	110.48
33	A	413	LMG	C7-O1-C1	-2.23	109.38	113.74
29	s	617	CLA	CHA-C1A-NA	-2.23	121.29	126.40
29	C	509	CLA	CMB-C2B-C3B	2.23	128.85	124.68
29	r	612	CLA	CMD-C2D-C3D	-2.23	122.49	127.61
29	b	615	CLA	C2A-C1A-CHA	2.23	127.75	123.86
29	N	604	CLA	O1D-CGD-CBD	-2.23	119.93	124.48
29	B	604	CLA	CMB-C2B-C3B	2.23	128.84	124.68
35	b	620	C7Z	C19-C9-C10	-2.23	119.80	122.92
29	b	603	CLA	CMD-C2D-C3D	-2.23	122.49	127.61
29	D	403	CLA	CAA-C2A-C3A	-2.23	106.68	112.78
29	r	603	CLA	CHA-C1A-NA	-2.23	121.30	126.40
45	y	609	CHL	C1-C2-C3	-2.23	122.19	126.04
35	B	620	C7Z	C19-C9-C10	-2.23	119.81	122.92
35	b	620	C7Z	C30-C31-C32	-2.23	116.27	123.22
29	R	613	CLA	CMD-C2D-C3D	-2.23	122.49	127.61
29	c	504	CLA	C1D-ND-C4D	-2.23	104.75	106.33
46	y	620	LUT	C18-C5-C6	-2.23	122.03	124.53
29	C	507	CLA	O2A-CGA-CBA	2.23	118.89	111.91
29	b	603	CLA	CMB-C2B-C3B	2.23	128.84	124.68
31	c	517	BCR	C35-C13-C12	2.22	121.58	118.08
29	Y	612	CLA	CMA-C3A-C4A	2.22	117.75	111.77
29	A	407	CLA	C1D-ND-C4D	-2.22	104.76	106.33
29	c	502	CLA	CHA-C1A-NA	-2.22	121.31	126.40
29	A	406	CLA	CMB-C2B-C1B	-2.22	125.05	128.46
31	C	517	BCR	C33-C5-C4	2.22	117.88	113.62
29	C	504	CLA	CAA-C2A-C3A	-2.22	106.69	112.78
42	D	405	PL9	C27-C28-C29	-2.22	122.31	127.66
29	Y	613	CLA	C1-O2A-CGA	2.22	122.27	116.44
47	g	622	XAT	C20-C13-C14	-2.22	119.81	122.92
29	B	607	CLA	C2D-C1D-ND	2.22	111.74	110.10
29	b	613	CLA	O2A-CGA-CBA	2.22	118.87	111.91
29	R	612	CLA	CMA-C3A-C4A	2.22	117.74	111.77
29	r	604	CLA	CMA-C3A-C4A	2.22	117.74	111.77
29	B	604	CLA	CMD-C2D-C3D	-2.22	122.51	127.61
29	B	613	CLA	CMD-C2D-C3D	-2.22	122.51	127.61
29	y	612	CLA	CHA-C1A-NA	-2.22	121.32	126.40
29	B	615	CLA	CMD-C2D-C3D	-2.22	122.51	127.61

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
29	s	610	CLA	CMD-C2D-C3D	-2.22	122.51	127.61
29	S	610	CLA	O1D-CGD-CBD	-2.22	119.95	124.48
47	n	622	XAT	C40-C33-C34	-2.22	119.82	122.92
29	B	613	CLA	C1D-ND-C4D	-2.22	104.76	106.33
29	R	604	CLA	O2A-CGA-CBA	2.22	118.86	111.91
47	r	622	XAT	C12-C13-C14	2.22	122.34	118.94
29	Y	608	CLA	CMD-C2D-C3D	-2.22	122.51	127.61
29	c	507	CLA	C3D-C2D-C1D	-2.22	102.81	105.83
29	b	614	CLA	O2A-CGA-CBA	2.22	118.86	111.91
29	S	605	CLA	CHA-C1A-NA	-2.22	121.32	126.40
31	c	516	BCR	C33-C5-C4	2.22	117.87	113.62
29	b	604	CLA	C2D-C1D-ND	2.22	111.74	110.10
47	y	622	XAT	C19-C9-C10	-2.22	119.82	122.92
29	G	613	CLA	O2D-CGD-O1D	-2.21	119.51	123.84
29	b	610	CLA	C1D-ND-C4D	-2.21	104.76	106.33
29	n	603	CLA	OBD-CAD-C3D	-2.21	123.19	128.52
45	r	606	CHL	C1B-CHB-C4A	-2.21	125.73	130.12
29	B	614	CLA	C1-O2A-CGA	2.21	122.25	116.44
29	C	512	CLA	O2D-CGD-O1D	-2.21	119.51	123.84
45	y	606	CHL	C4D-CHA-C1A	2.21	123.94	121.25
31	C	517	BCR	C35-C13-C12	2.21	121.56	118.08
29	c	501	CLA	CAA-C2A-C3A	-2.21	106.72	112.78
29	y	603	CLA	O2A-CGA-CBA	2.21	118.85	111.91
29	A	407	CLA	O2D-CGD-O1D	-2.21	119.51	123.84
45	g	609	CHL	CHC-C1C-NC	2.21	127.56	124.20
45	g	608	CHL	C4A-NA-C1A	2.21	107.70	106.71
47	G	622	XAT	C18-C5-C6	-2.21	118.56	122.26
29	R	604	CLA	CMA-C3A-C4A	2.21	117.71	111.77
30	A	408	PHO	CMC-C2C-C3C	2.21	129.11	124.94
29	b	611	CLA	CMA-C3A-C4A	2.21	117.71	111.77
29	b	602	CLA	O2A-CGA-CBA	2.21	118.84	111.91
29	s	612	CLA	CHA-C1A-NA	-2.21	121.34	126.40
29	B	605	CLA	O2A-CGA-CBA	2.21	118.83	111.91
29	N	610	CLA	C1D-ND-C4D	-2.21	104.77	106.33
29	C	513	CLA	CMD-C2D-C3D	-2.21	122.54	127.61
29	S	605	CLA	C2A-C1A-CHA	2.21	127.72	123.86
29	B	612	CLA	CHA-C4D-ND	2.21	137.11	132.50
29	R	602	CLA	CMD-C2D-C3D	-2.21	122.54	127.61
47	y	622	XAT	C39-C29-C30	-2.21	119.83	122.92
29	C	502	CLA	C3D-C2D-C1D	-2.20	102.82	105.83
29	Y	610	CLA	C1D-ND-C4D	-2.20	104.77	106.33
29	y	614	CLA	CMD-C2D-C3D	-2.20	122.54	127.61

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
45	n	608	CHL	C4A-NA-C1A	2.20	107.70	106.71
48	s	623	NEX	C4-C3-C2	2.20	115.03	110.77
44	H	101	RRX	C35-C13-C14	-2.20	119.84	122.92
29	G	602	CLA	CMD-C2D-C3D	-2.20	122.55	127.61
31	C	516	BCR	C27-C26-C25	-2.20	119.53	122.73
29	a	407	CLA	C1D-ND-C4D	-2.20	104.77	106.33
29	n	603	CLA	CBC-CAC-C3C	-2.20	106.36	112.43
42	D	405	PL9	O1-C4-C3	-2.20	118.29	120.72
46	Y	620	LUT	C30-C31-C32	-2.20	116.34	123.22
29	S	614	CLA	CHA-C1A-NA	-2.20	121.36	126.40
29	N	604	CLA	C3D-C2D-C1D	-2.20	102.83	105.83
31	b	618	BCR	C33-C5-C4	2.20	117.85	113.62
29	c	504	CLA	CAA-C2A-C3A	-2.20	106.75	112.78
45	y	607	CHL	C1-C2-C3	-2.20	122.24	126.04
29	b	613	CLA	CMD-C2D-C3D	-2.20	122.55	127.61
29	S	602	CLA	O2A-CGA-CBA	2.20	118.81	111.91
29	r	602	CLA	CMD-C2D-C3D	-2.20	122.55	127.61
29	B	603	CLA	CHA-C1A-NA	-2.20	121.36	126.40
29	n	611	CLA	CHA-C1A-NA	-2.20	121.36	126.40
29	g	604	CLA	CMA-C3A-C4A	2.20	117.68	111.77
45	G	601	CHL	C4A-NA-C1A	2.20	107.69	106.71
29	a	405	CLA	C3D-C2D-C1D	-2.20	102.83	105.83
29	Y	604	CLA	CAA-C2A-C3A	-2.20	106.76	112.78
29	R	612	CLA	CHA-C1A-NA	-2.20	121.37	126.40
29	g	604	CLA	CHA-C1A-NA	-2.20	121.37	126.40
38	c	519	DGD	C2G-O2G-C1B	-2.20	112.39	117.79
29	r	603	CLA	O2D-CGD-O1D	-2.20	119.55	123.84
29	R	610	CLA	CHA-C1A-NA	-2.20	121.37	126.40
29	S	609	CLA	CHA-C1A-NA	-2.19	121.38	126.40
29	C	512	CLA	O2A-CGA-CBA	2.19	118.78	111.91
29	r	613	CLA	C1D-ND-C4D	-2.19	104.78	106.33
31	d	404	BCR	C1-C6-C5	-2.19	119.53	122.61
29	A	405	CLA	CAC-C3C-C4C	2.19	127.65	124.81
46	Y	621	LUT	C16-C1-C6	-2.19	106.75	110.30
46	s	621	LUT	C1-C6-C5	-2.19	119.53	122.61
29	C	506	CLA	O1D-CGD-CBD	-2.19	120.01	124.48
29	r	610	CLA	O1D-CGD-CBD	-2.19	120.01	124.48
29	S	614	CLA	CMB-C2B-C3B	2.19	128.77	124.68
29	n	603	CLA	C3D-C2D-C1D	-2.19	102.85	105.83
29	B	610	CLA	C1D-ND-C4D	-2.19	104.78	106.33
45	G	606	CHL	CMB-C2B-C1B	-2.19	125.10	128.46
29	s	609	CLA	CHA-C1A-NA	-2.19	121.39	126.40

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
29	c	507	CLA	C1-O2A-CGA	2.19	122.18	116.44
46	n	621	LUT	C19-C9-C8	2.19	121.52	118.08
29	Y	604	CLA	CMB-C2B-C3B	2.18	128.77	124.68
29	c	506	CLA	CMD-C2D-C3D	-2.18	122.59	127.61
29	b	602	CLA	C1D-ND-C4D	-2.18	104.78	106.33
29	n	614	CLA	C1D-ND-C4D	-2.18	104.78	106.33
29	g	610	CLA	C1D-ND-C4D	-2.18	104.78	106.33
29	y	608	CLA	CAA-C2A-C3A	-2.18	106.80	112.78
29	G	612	CLA	CHA-C1A-NA	-2.18	121.40	126.40
45	g	607	CHL	C1B-CHB-C4A	-2.18	125.79	130.12
29	N	611	CLA	CAA-C2A-C3A	-2.18	106.80	112.78
29	b	609	CLA	C3D-C2D-C1D	-2.18	102.85	105.83
29	B	609	CLA	CHD-C1D-ND	-2.18	122.45	124.45
29	S	611	CLA	CHA-C1A-NA	-2.18	121.40	126.40
29	r	613	CLA	CHA-C1A-NA	-2.18	121.40	126.40
45	G	605	CHL	C1B-CHB-C4A	-2.18	125.80	130.12
29	y	608	CLA	CMD-C2D-C3D	-2.18	122.60	127.61
29	b	612	CLA	C1C-C2C-C3C	-2.18	104.66	106.96
31	B	618	BCR	C34-C9-C10	-2.18	119.87	122.92
29	G	614	CLA	C1D-ND-C4D	-2.18	104.79	106.33
29	y	610	CLA	C1D-ND-C4D	-2.18	104.79	106.33
29	g	610	CLA	CAA-C2A-C3A	-2.18	106.81	112.78
29	N	611	CLA	CHA-C1A-NA	-2.18	121.41	126.40
31	b	618	BCR	C38-C26-C25	-2.18	122.08	124.53
45	s	608	CHL	C1B-CHB-C4A	-2.18	125.80	130.12
29	s	612	CLA	CMB-C2B-C3B	2.18	128.75	124.68
46	G	620	LUT	C10-C11-C12	-2.18	116.42	123.22
29	b	616	CLA	CMD-C2D-C3D	-2.18	122.60	127.61
29	B	607	CLA	O1D-CGD-CBD	-2.18	120.03	124.48
29	B	603	CLA	C1-O2A-CGA	2.18	122.16	116.44
29	b	614	CLA	C1D-ND-C4D	-2.18	104.79	106.33
29	c	503	CLA	C1D-ND-C4D	-2.18	104.79	106.33
29	S	610	CLA	CHA-C1A-NA	-2.18	121.41	126.40
45	n	605	CHL	C1B-CHB-C4A	-2.18	125.80	130.12
29	r	610	CLA	C1C-C2C-C3C	-2.18	104.67	106.96
29	Y	603	CLA	O1D-CGD-CBD	-2.18	120.03	124.48
29	C	506	CLA	CMD-C2D-C3D	-2.18	122.61	127.61
45	r	607	CHL	CHD-C4C-C3C	2.18	128.04	124.84
48	g	623	NEX	C19-C9-C10	-2.18	119.88	122.92
29	Y	613	CLA	CMD-C2D-C3D	-2.17	122.61	127.61
29	c	512	CLA	C1-O2A-CGA	2.17	122.15	116.44
29	R	611	CLA	CHA-C1A-NA	-2.17	121.42	126.40

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
29	r	608	CLA	CHA-C1A-NA	-2.17	121.42	126.40
29	S	609	CLA	CMD-C2D-C3D	-2.17	122.61	127.61
29	S	610	CLA	CMD-C2D-C3D	-2.17	122.61	127.61
29	R	610	CLA	O2A-CGA-CBA	2.17	118.73	111.91
44	H	101	RRX	C38-C26-C25	-2.17	122.09	124.53
29	N	612	CLA	CHA-C1A-NA	-2.17	121.42	126.40
29	B	614	CLA	C3D-C2D-C1D	-2.17	102.87	105.83
30	a	408	PHO	CMC-C2C-C3C	2.17	129.04	124.94
29	Y	608	CLA	C1-O2A-CGA	2.17	122.14	116.44
29	C	502	CLA	CHA-C1A-NA	-2.17	121.43	126.40
45	N	605	CHL	C1B-CHB-C4A	-2.17	125.82	130.12
29	A	405	CLA	C1D-ND-C4D	-2.17	104.79	106.33
29	n	612	CLA	C1D-ND-C4D	-2.17	104.79	106.33
45	s	608	CHL	CMB-C2B-C1B	-2.17	125.13	128.46
48	g	623	NEX	C4-C3-C2	2.17	114.96	110.77
29	c	506	CLA	O2A-CGA-CBA	2.17	118.71	111.91
48	N	623	NEX	C26-C27-C28	-2.17	121.41	125.99
29	C	503	CLA	CHA-C1A-NA	-2.17	121.43	126.40
29	s	610	CLA	CHA-C1A-NA	-2.17	121.43	126.40
29	c	501	CLA	O1D-CGD-CBD	-2.17	120.05	124.48
29	y	608	CLA	CHA-C1A-NA	-2.17	121.44	126.40
48	s	623	NEX	C31-C30-C29	2.17	130.40	127.31
29	s	617	CLA	CAA-C2A-C3A	-2.17	106.84	112.78
29	N	613	CLA	C1D-ND-C4D	-2.17	104.80	106.33
51	Y	625	SPH	C3-C4-C5	-2.17	119.96	124.79
46	s	620	LUT	C20-C13-C12	2.17	121.49	118.08
29	a	407	CLA	O2D-CGD-O1D	-2.17	119.60	123.84
45	Y	605	CHL	C4D-CHA-C1A	2.17	123.89	121.25
29	s	610	CLA	C1-C2-C3	-2.17	122.30	126.04
29	y	611	CLA	CHA-C1A-NA	-2.17	121.44	126.40
29	n	604	CLA	O1D-CGD-CBD	-2.17	120.05	124.48
29	N	614	CLA	O2A-CGA-CBA	2.16	118.70	111.91
29	B	608	CLA	CMB-C2B-C1B	-2.16	125.14	128.46
29	b	603	CLA	CHA-C1A-NA	-2.16	121.44	126.40
29	C	505	CLA	CMD-C2D-C3D	-2.16	122.64	127.61
46	N	621	LUT	C10-C11-C12	-2.16	116.46	123.22
29	y	603	CLA	C3D-C2D-C1D	-2.16	102.88	105.83
29	Y	602	CLA	C1-C2-C3	-2.16	122.30	126.04
29	A	405	CLA	CAA-C2A-C3A	-2.16	106.86	112.78
45	g	608	CHL	CMB-C2B-C1B	-2.16	125.14	128.46
29	G	610	CLA	CMD-C2D-C3D	-2.16	122.64	127.61
29	B	609	CLA	O1D-CGD-CBD	-2.16	120.06	124.48

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
29	G	603	CLA	CMD-C2D-C3D	-2.16	122.64	127.61
29	c	507	CLA	CMB-C2B-C3B	2.16	128.72	124.68
29	b	608	CLA	CMB-C2B-C1B	-2.16	125.14	128.46
39	n	624	LHG	C6-C5-C4	-2.16	106.68	111.79
29	r	604	CLA	CMD-C2D-C3D	-2.16	122.65	127.61
29	g	614	CLA	C1D-ND-C4D	-2.16	104.80	106.33
29	g	610	CLA	CHA-C1A-NA	-2.16	121.45	126.40
29	D	402	CLA	CMD-C2D-C3D	-2.16	122.65	127.61
31	c	515	BCR	C38-C26-C25	-2.16	122.10	124.53
29	C	503	CLA	CAA-C2A-C3A	-2.16	106.87	112.78
29	C	509	CLA	CHA-C1A-NA	-2.16	121.45	126.40
29	S	612	CLA	CHA-C1A-NA	-2.16	121.45	126.40
29	s	617	CLA	CMD-C2D-C3D	-2.16	122.65	127.61
47	N	622	XAT	C26-C27-C28	-2.16	121.43	125.99
29	G	611	CLA	CHA-C1A-NA	-2.16	121.46	126.40
29	g	604	CLA	O1D-CGD-CBD	-2.16	120.07	124.48
45	g	607	CHL	C4A-NA-C1A	2.16	107.67	106.71
29	r	604	CLA	CMB-C2B-C3B	2.16	128.71	124.68
45	s	606	CHL	CMB-C2B-C1B	-2.15	125.15	128.46
29	y	604	CLA	C1D-ND-C4D	-2.15	104.81	106.33
46	G	620	LUT	C31-C32-C33	-2.15	120.37	126.42
29	y	604	CLA	O2A-CGA-CBA	2.15	118.67	111.91
29	c	509	CLA	CMB-C2B-C1B	-2.15	125.15	128.46
29	C	512	CLA	CHA-C1A-NA	-2.15	121.47	126.40
29	Y	604	CLA	O2A-CGA-CBA	2.15	118.66	111.91
29	B	611	CLA	C3D-C2D-C1D	-2.15	102.89	105.83
29	g	611	CLA	CHA-C1A-NA	-2.15	121.47	126.40
29	r	602	CLA	CHA-C1A-NA	-2.15	121.47	126.40
29	r	602	CLA	C1D-ND-C4D	-2.15	104.81	106.33
29	G	603	CLA	CHA-C1A-NA	-2.15	121.47	126.40
29	B	607	CLA	C1-C2-C3	-2.15	122.32	126.04
45	y	607	CHL	C1B-CHB-C4A	-2.15	125.86	130.12
29	r	604	CLA	O2A-CGA-CBA	2.15	118.66	111.91
29	N	610	CLA	O1D-CGD-CBD	-2.15	120.08	124.48
29	R	604	CLA	O1D-CGD-CBD	-2.15	120.08	124.48
29	c	502	CLA	C1C-C2C-C3C	-2.15	104.70	106.96
29	C	504	CLA	C1D-ND-C4D	-2.15	104.81	106.33
45	y	606	CHL	CMB-C2B-C1B	-2.15	125.16	128.46
45	Y	601	CHL	CHC-C1C-NC	2.15	127.46	124.20
29	Y	603	CLA	CHA-C1A-NA	-2.15	121.48	126.40
45	Y	606	CHL	C4D-CHA-C1A	2.15	123.86	121.25
51	y	625	SPH	C3-C4-C5	-2.15	120.00	124.79

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
29	c	509	CLA	CHA-C1A-NA	-2.15	121.48	126.40
45	S	601	CHL	C1B-CHB-C4A	-2.15	125.86	130.12
29	S	613	CLA	C3D-C2D-C1D	-2.15	102.90	105.83
47	Y	622	XAT	C18-C5-C6	-2.15	118.66	122.26
45	R	606	CHL	CMB-C2B-C1B	-2.15	125.17	128.46
31	c	517	BCR	C33-C5-C6	-2.15	122.12	124.53
29	C	508	CLA	CHA-C1A-NA	-2.15	121.48	126.40
29	r	610	CLA	CMD-C2D-C3D	-2.15	122.68	127.61
29	s	605	CLA	CMD-C2D-C3D	-2.15	122.68	127.61
29	R	613	CLA	O1D-CGD-CBD	-2.15	120.09	124.48
46	n	620	LUT	C15-C35-C34	-2.15	119.08	123.47
29	b	611	CLA	C3D-C2D-C1D	-2.14	102.90	105.83
29	R	610	CLA	CMD-C2D-C3D	-2.14	122.68	127.61
45	R	607	CHL	CMB-C2B-C1B	-2.14	125.17	128.46
31	D	404	BCR	C4-C5-C6	-2.14	119.62	122.73
47	g	622	XAT	C40-C33-C34	-2.14	119.92	122.92
29	S	602	CLA	CMC-C2C-C1C	2.14	128.30	125.04
45	G	609	CHL	CMB-C2B-C1B	-2.14	125.17	128.46
29	S	617	CLA	CHA-C1A-NA	-2.14	121.49	126.40
45	Y	605	CHL	C4A-NA-C1A	2.14	107.67	106.71
45	N	605	CHL	CMB-C2B-C1B	-2.14	125.17	128.46
45	Y	609	CHL	C4D-CHA-C1A	2.14	123.86	121.25
29	S	612	CLA	C1D-ND-C4D	-2.14	104.81	106.33
47	g	622	XAT	C39-C29-C30	-2.14	119.92	122.92
29	c	505	CLA	CMD-C2D-C3D	-2.14	122.69	127.61
45	Y	609	CHL	CMB-C2B-C1B	-2.14	125.17	128.46
29	C	507	CLA	C3D-C2D-C1D	-2.14	102.91	105.83
29	s	610	CLA	C3D-C2D-C1D	-2.14	102.91	105.83
29	r	611	CLA	CHA-C1A-NA	-2.14	121.50	126.40
29	b	604	CLA	C1-C2-C3	-2.14	122.34	126.04
29	b	610	CLA	C3D-C2D-C1D	-2.14	102.91	105.83
29	s	613	CLA	C3D-C2D-C1D	-2.14	102.91	105.83
29	N	603	CLA	C1D-ND-C4D	-2.14	104.81	106.33
29	N	611	CLA	O1D-CGD-CBD	-2.14	120.11	124.48
29	a	405	CLA	C1D-ND-C4D	-2.14	104.82	106.33
29	n	604	CLA	CMA-C3A-C4A	2.14	117.52	111.77
29	B	614	CLA	CHA-C1A-NA	-2.14	121.50	126.40
29	B	616	CLA	CMD-C2D-C3D	-2.14	122.70	127.61
29	S	604	CLA	CBC-CAC-C3C	-2.14	106.54	112.43
29	Y	603	CLA	CMD-C2D-C3D	-2.14	122.70	127.61
45	S	608	CHL	CMB-C2B-C1B	-2.14	125.18	128.46
29	c	505	CLA	C1-O2A-CGA	2.14	122.05	116.44

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
29	n	613	CLA	CMD-C2D-C3D	-2.14	122.70	127.61
29	s	605	CLA	CHA-C1A-NA	-2.14	121.51	126.40
45	n	605	CHL	CMB-C2B-C1B	-2.13	125.18	128.46
29	y	610	CLA	CHA-C1A-NA	-2.13	121.51	126.40
29	B	602	CLA	C1D-ND-C4D	-2.13	104.82	106.33
29	c	507	CLA	O2A-CGA-CBA	2.13	118.60	111.91
29	B	608	CLA	C1-C2-C3	-2.13	122.35	126.04
45	y	609	CHL	CMB-C2B-C1B	-2.13	125.19	128.46
29	n	612	CLA	CMD-C2D-C3D	-2.13	122.71	127.61
29	Y	608	CLA	CHA-C1A-NA	-2.13	121.51	126.40
45	N	605	CHL	C1-O2A-CGA	2.13	122.04	116.44
29	n	602	CLA	CAA-C2A-C3A	-2.13	106.94	112.78
45	S	606	CHL	CMB-C2B-C1B	-2.13	125.19	128.46
45	N	609	CHL	C1-C2-C3	-2.13	122.36	126.04
31	C	516	BCR	C33-C5-C4	2.13	117.71	113.62
47	R	621	XAT	C30-C31-C32	-2.13	116.57	123.22
46	s	620	LUT	C36-C21-C26	2.13	112.77	109.55
45	Y	606	CHL	CMB-C2B-C1B	-2.13	125.19	128.46
29	D	403	CLA	CHA-C1A-NA	-2.13	121.52	126.40
29	g	602	CLA	CHA-C1A-NA	-2.13	121.52	126.40
29	b	609	CLA	O2A-CGA-CBA	2.13	118.59	111.91
29	C	503	CLA	O2D-CGD-O1D	-2.13	119.67	123.84
45	s	607	CHL	CMB-C2B-C1B	-2.13	125.19	128.46
29	Y	604	CLA	C1D-ND-C4D	-2.13	104.82	106.33
29	g	613	CLA	C1D-ND-C4D	-2.13	104.82	106.33
45	g	609	CHL	CHB-C4A-NA	2.13	127.45	124.51
45	Y	605	CHL	CMB-C2B-C1B	-2.13	125.19	128.46
46	y	620	LUT	C20-C13-C12	2.13	121.43	118.08
29	s	604	CLA	CHA-C1A-NA	-2.13	121.53	126.40
29	y	611	CLA	CAA-C2A-C3A	-2.13	106.96	112.78
39	g	624	LHG	C6-C5-C4	-2.13	106.76	111.79
29	N	604	CLA	CMD-C2D-C3D	-2.13	122.72	127.61
45	s	601	CHL	CMB-C2B-C1B	-2.13	125.20	128.46
29	Y	612	CLA	CMB-C2B-C3B	2.13	128.65	124.68
45	N	606	CHL	CMB-C2B-C1B	-2.12	125.20	128.46
29	s	610	CLA	O1D-CGD-CBD	-2.12	120.14	124.48
29	s	609	CLA	C1D-ND-C4D	-2.12	104.83	106.33
45	g	608	CHL	C4D-CHA-C1A	2.12	123.83	121.25
45	N	606	CHL	C1-C2-C3	-2.12	122.37	126.04
29	b	609	CLA	CHD-C1D-ND	-2.12	122.50	124.45
29	b	614	CLA	CHA-C1A-NA	-2.12	121.54	126.40
29	N	610	CLA	CMD-C2D-C3D	-2.12	122.73	127.61

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
29	y	613	CLA	CMD-C2D-C3D	-2.12	122.73	127.61
45	n	606	CHL	CMB-C2B-C1B	-2.12	125.20	128.46
29	Y	613	CLA	CMB-C2B-C1B	-2.12	125.20	128.46
45	g	601	CHL	CMB-C2B-C1B	-2.12	125.20	128.46
29	R	602	CLA	C3D-C2D-C1D	-2.12	102.94	105.83
45	r	606	CHL	CHC-C1C-NC	2.12	127.42	124.20
29	N	603	CLA	O1D-CGD-CBD	-2.12	120.14	124.48
45	G	605	CHL	CMB-C2B-C1B	-2.12	125.20	128.46
45	r	607	CHL	CMB-C2B-C1B	-2.12	125.20	128.46
29	B	616	CLA	C2A-C1A-CHA	2.12	127.57	123.86
29	Y	611	CLA	CHA-C1A-NA	-2.12	121.54	126.40
45	r	606	CHL	CMB-C2B-C1B	-2.12	125.20	128.46
46	n	620	LUT	C11-C10-C9	-2.12	124.28	127.31
29	s	602	CLA	O2A-CGA-CBA	2.12	118.56	111.91
29	s	603	CLA	O2A-CGA-CBA	2.12	118.56	111.91
29	B	616	CLA	C1D-ND-C4D	-2.12	104.83	106.33
29	n	613	CLA	C1D-ND-C4D	-2.12	104.83	106.33
29	b	604	CLA	CHA-C1A-NA	-2.12	121.54	126.40
29	g	612	CLA	CHA-C1A-NA	-2.12	121.54	126.40
35	B	620	C7Z	C35-C15-C14	-2.12	119.13	123.47
29	C	504	CLA	CMD-C2D-C3D	-2.12	122.74	127.61
29	r	609	CLA	C2D-C1D-ND	2.12	111.67	110.10
45	N	605	CHL	C4D-CHA-C1A	2.12	123.83	121.25
29	S	604	CLA	C1D-ND-C4D	-2.12	104.83	106.33
45	G	608	CHL	CMB-C2B-C1B	-2.12	125.21	128.46
29	Y	610	CLA	CHA-C1A-NA	-2.12	121.55	126.40
29	G	604	CLA	CHA-C1A-NA	-2.12	121.55	126.40
47	r	622	XAT	C39-C29-C28	2.12	121.41	118.08
45	N	601	CHL	CMB-C2B-C1B	-2.12	125.21	128.46
45	G	601	CHL	CMB-C2B-C1B	-2.12	125.21	128.46
29	n	602	CLA	CMD-C2D-C3D	-2.12	122.75	127.61
45	n	601	CHL	CMB-C2B-C1B	-2.12	125.21	128.46
29	d	403	CLA	CHA-C1A-NA	-2.12	121.55	126.40
29	n	614	CLA	CAA-C2A-C3A	-2.12	106.98	112.78
29	B	610	CLA	CHA-C1A-NA	-2.12	121.55	126.40
45	S	601	CHL	CMB-C2B-C1B	-2.12	125.21	128.46
29	G	611	CLA	CMD-C2D-C3D	-2.12	122.75	127.61
29	C	501	CLA	CAA-C2A-C3A	-2.12	106.98	112.78
29	n	602	CLA	CMB-C2B-C1B	-2.12	125.21	128.46
29	B	602	CLA	CHA-C1A-NA	-2.12	121.55	126.40
29	S	610	CLA	C3D-C2D-C1D	-2.11	102.94	105.83
29	c	504	CLA	CMB-C2B-C1B	-2.11	125.21	128.46

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
29	C	506	CLA	O2A-CGA-CBA	2.11	118.54	111.91
29	a	406	CLA	O2D-CGD-O1D	-2.11	119.70	123.84
29	Y	602	CLA	CMC-C2C-C1C	2.11	128.26	125.04
29	G	602	CLA	CMB-C2B-C3B	2.11	128.63	124.68
29	G	604	CLA	O1D-CGD-CBD	-2.11	120.16	124.48
29	C	508	CLA	C1C-C2C-C3C	-2.11	104.73	106.96
45	y	609	CHL	C4A-NA-C1A	2.11	107.66	106.71
29	y	614	CLA	CHA-C1A-NA	-2.11	121.56	126.40
29	A	410	CLA	C1D-ND-C4D	-2.11	104.83	106.33
46	n	620	LUT	C19-C9-C8	2.11	121.41	118.08
29	y	608	CLA	O2A-CGA-CBA	2.11	118.54	111.91
45	g	605	CHL	CMB-C2B-C1B	-2.11	125.22	128.46
29	b	617	CLA	CHA-C1A-NA	-2.11	121.56	126.40
29	N	603	CLA	CHA-C1A-NA	-2.11	121.56	126.40
29	N	614	CLA	CHA-C1A-NA	-2.11	121.56	126.40
29	Y	602	CLA	C1-O2A-CGA	2.11	121.98	116.44
45	R	607	CHL	C1B-CHB-C4A	-2.11	125.93	130.12
48	y	623	NEX	C26-C27-C28	-2.11	121.53	125.99
45	y	605	CHL	CMB-C2B-C1B	-2.11	125.22	128.46
29	n	604	CLA	C3D-C2D-C1D	-2.11	102.95	105.83
29	Y	602	CLA	CMB-C2B-C3B	2.11	128.63	124.68
29	s	610	CLA	O2D-CGD-O1D	-2.11	119.71	123.84
29	s	611	CLA	CHA-C1A-NA	-2.11	121.57	126.40
29	N	603	CLA	CMD-C2D-C3D	-2.11	122.76	127.61
29	C	513	CLA	CMB-C2B-C1B	-2.11	125.22	128.46
46	N	621	LUT	C16-C1-C6	-2.11	106.88	110.30
29	n	604	CLA	CMB-C2B-C3B	2.11	128.62	124.68
46	r	620	LUT	C35-C15-C14	-2.11	119.15	123.47
29	b	605	CLA	C2D-C1D-ND	2.11	111.66	110.10
29	C	511	CLA	CAA-CBA-CGA	-2.11	107.09	113.25
44	h	101	RRX	C1-C6-C7	2.11	121.74	115.78
29	r	610	CLA	CHA-C1A-NA	-2.11	121.57	126.40
29	C	511	CLA	C3D-C2D-C1D	-2.11	102.95	105.83
29	g	604	CLA	C3D-C2D-C1D	-2.11	102.95	105.83
29	C	510	CLA	O1D-CGD-CBD	-2.11	120.17	124.48
47	G	622	XAT	C39-C29-C30	-2.11	119.97	122.92
29	n	602	CLA	C1D-ND-C4D	-2.11	104.84	106.33
29	g	602	CLA	C1D-ND-C4D	-2.11	104.84	106.33
31	c	517	BCR	C27-C26-C25	-2.11	119.67	122.73
29	y	602	CLA	C3D-C2D-C1D	-2.11	102.96	105.83
29	y	603	CLA	CHA-C1A-NA	-2.11	121.58	126.40
45	S	601	CHL	C4A-NA-C1A	2.10	107.65	106.71

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
29	N	604	CLA	CHA-C1A-NA	-2.10	121.58	126.40
29	c	507	CLA	CMD-C2D-C3D	-2.10	122.78	127.61
29	s	611	CLA	CAA-C2A-C3A	-2.10	107.02	112.78
29	y	610	CLA	CMD-C2D-C3D	-2.10	122.78	127.61
45	Y	607	CHL	C1-C2-C3	-2.10	122.41	126.04
29	B	617	CLA	C1C-C2C-C3C	-2.10	104.75	106.96
45	G	608	CHL	C1B-CHB-C4A	-2.10	125.95	130.12
45	Y	605	CHL	CHD-C4C-C3C	2.10	127.93	124.84
29	s	613	CLA	CMD-C2D-C3D	-2.10	122.78	127.61
29	b	606	CLA	CMB-C2B-C3B	2.10	128.61	124.68
45	Y	609	CHL	C1-C2-C3	-2.10	122.41	126.04
29	a	407	CLA	CHA-C1A-NA	-2.10	121.59	126.40
29	c	508	CLA	O2D-CGD-O1D	-2.10	119.73	123.84
29	S	603	CLA	O2A-CGA-CBA	2.10	118.50	111.91
29	n	612	CLA	C3D-C2D-C1D	-2.10	102.96	105.83
29	y	604	CLA	CMB-C2B-C1B	-2.10	125.23	128.46
29	B	605	CLA	C2D-C1D-ND	2.10	111.65	110.10
45	G	605	CHL	C1-O2A-CGA	2.10	122.83	116.73
29	c	511	CLA	CAA-CBA-CGA	-2.10	107.12	113.25
29	b	615	CLA	CMD-C2D-C3D	-2.10	122.78	127.61
29	C	507	CLA	CMD-C2D-C3D	-2.10	122.78	127.61
29	b	613	CLA	C1D-ND-C4D	-2.10	104.84	106.33
29	r	603	CLA	C1D-ND-C4D	-2.10	104.84	106.33
29	S	613	CLA	CMD-C2D-C3D	-2.10	122.79	127.61
29	b	614	CLA	C3D-C2D-C1D	-2.10	102.97	105.83
29	b	605	CLA	C1-O2A-CGA	2.10	121.95	116.44
29	c	502	CLA	O2A-CGA-CBA	2.10	118.49	111.91
45	G	601	CHL	C1B-CHB-C4A	-2.10	125.96	130.12
45	y	606	CHL	C1B-CHB-C4A	-2.10	125.96	130.12
29	n	602	CLA	O1D-CGD-CBD	-2.10	120.19	124.48
29	C	511	CLA	C1D-ND-C4D	-2.10	104.84	106.33
29	c	513	CLA	C1D-ND-C4D	-2.10	104.84	106.33
29	y	611	CLA	CMB-C2B-C3B	2.10	128.60	124.68
29	c	508	CLA	C1C-C2C-C3C	-2.10	104.75	106.96
29	Y	608	CLA	C3D-C2D-C1D	-2.10	102.97	105.83
29	Y	614	CLA	CHA-C1A-NA	-2.10	121.60	126.40
29	G	612	CLA	C1D-ND-C4D	-2.10	104.85	106.33
29	B	609	CLA	C1-O2A-CGA	2.10	121.94	116.44
29	A	410	CLA	CMB-C2B-C3B	2.10	128.60	124.68
29	B	617	CLA	CHA-C1A-NA	-2.10	121.60	126.40
29	R	611	CLA	CMD-C2D-C3D	-2.09	122.80	127.61
29	a	405	CLA	CAC-C3C-C4C	2.09	127.53	124.81

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
29	c	511	CLA	C3D-C2D-C1D	-2.09	102.97	105.83
29	y	613	CLA	C3D-C2D-C1D	-2.09	102.97	105.83
29	S	602	CLA	CHA-C1A-NA	-2.09	121.60	126.40
29	a	407	CLA	C3D-C2D-C1D	-2.09	102.97	105.83
29	g	611	CLA	C3D-C2D-C1D	-2.09	102.97	105.83
45	g	601	CHL	C1B-CHB-C4A	-2.09	125.97	130.12
29	N	614	CLA	CMD-C2D-C3D	-2.09	122.80	127.61
31	c	515	BCR	C35-C13-C12	2.09	121.37	118.08
48	S	622	NEX	C4-C3-C2	2.09	114.81	110.77
29	n	602	CLA	CHA-C1A-NA	-2.09	121.61	126.40
29	g	610	CLA	O1D-CGD-CBD	-2.09	120.20	124.48
29	Y	603	CLA	CAA-C2A-C3A	-2.09	107.05	112.78
31	c	516	BCR	C30-C25-C24	2.09	121.70	115.78
45	g	609	CHL	CMB-C2B-C1B	-2.09	125.25	128.46
29	n	611	CLA	CMD-C2D-C3D	-2.09	122.80	127.61
29	b	602	CLA	CHA-C1A-NA	-2.09	121.61	126.40
29	g	614	CLA	CHA-C1A-NA	-2.09	121.61	126.40
29	S	609	CLA	C1D-ND-C4D	-2.09	104.85	106.33
29	n	610	CLA	CMD-C2D-C3D	-2.09	122.81	127.61
45	G	601	CHL	CHD-C4C-C3C	2.09	127.91	124.84
48	N	623	NEX	O24-C25-C38	-2.09	112.55	115.06
29	r	610	CLA	C6-C7-C8	-2.09	109.17	115.92
29	B	614	CLA	O2A-CGA-CBA	2.09	118.46	111.91
29	S	611	CLA	CMD-C2D-C3D	-2.09	122.81	127.61
29	r	611	CLA	C1D-ND-C4D	-2.09	104.85	106.33
29	B	603	CLA	CMB-C2B-C1B	-2.09	125.25	128.46
29	g	603	CLA	CHA-C1A-NA	-2.09	121.61	126.40
29	y	608	CLA	C3D-C2D-C1D	-2.09	102.98	105.83
46	N	620	LUT	C35-C15-C14	-2.09	119.20	123.47
29	b	609	CLA	O1D-CGD-CBD	-2.09	120.21	124.48
29	C	509	CLA	CMA-C3A-C4A	2.09	117.38	111.77
29	c	502	CLA	O2D-CGD-O1D	-2.09	119.76	123.84
45	n	609	CHL	C1B-CHB-C4A	-2.09	125.98	130.12
29	a	410	CLA	C1D-ND-C4D	-2.09	104.85	106.33
29	A	407	CLA	CHA-C1A-NA	-2.09	121.62	126.40
29	b	614	CLA	CMD-C2D-C3D	-2.09	122.82	127.61
45	S	607	CHL	CMB-C2B-C1B	-2.09	125.26	128.46
29	C	508	CLA	O2A-CGA-CBA	2.08	118.45	111.91
29	g	612	CLA	C1D-ND-C4D	-2.08	104.85	106.33
29	C	513	CLA	C3D-C2D-C1D	-2.08	102.99	105.83
45	r	606	CHL	C4D-CHA-C1A	2.08	123.79	121.25
29	N	613	CLA	CHA-C1A-NA	-2.08	121.62	126.40

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
29	A	407	CLA	C1-O2A-CGA	2.08	121.91	116.44
29	s	602	CLA	CMB-C2B-C3B	2.08	128.58	124.68
39	G	630	LHG	C6-C5-C4	-2.08	106.86	111.79
45	R	606	CHL	C3A-C2A-C1A	2.08	104.46	101.34
29	s	602	CLA	C3D-C2D-C1D	-2.08	102.99	105.83
29	s	611	CLA	CMD-C2D-C3D	-2.08	122.82	127.61
46	R	620	LUT	C20-C13-C14	-2.08	120.00	122.92
29	G	603	CLA	O2A-CGA-CBA	2.08	118.44	111.91
29	n	614	CLA	C3D-C2D-C1D	-2.08	102.99	105.83
29	c	506	CLA	O1D-CGD-CBD	-2.08	120.22	124.48
42	D	405	PL9	C12-C13-C14	-2.08	122.65	127.66
29	b	612	CLA	CMA-C3A-C4A	2.08	117.37	111.77
48	s	623	NEX	C40-C33-C34	-2.08	120.01	122.92
29	b	609	CLA	C1-O2A-CGA	2.08	121.90	116.44
48	N	623	NEX	C20-C13-C14	-2.08	120.01	122.92
29	B	606	CLA	CMB-C2B-C3B	2.08	128.57	124.68
45	y	607	CHL	C4D-CHA-C1A	2.08	123.78	121.25
29	c	508	CLA	CHA-C1A-NA	-2.08	121.64	126.40
29	N	602	CLA	C6-C5-C3	-2.08	108.00	113.45
29	S	610	CLA	CAA-C2A-C3A	-2.08	107.09	112.78
29	C	513	CLA	C1D-ND-C4D	-2.08	104.86	106.33
45	n	608	CHL	CHD-C4C-C3C	2.08	127.89	124.84
29	y	604	CLA	CHA-C1A-NA	-2.08	121.64	126.40
29	R	610	CLA	C3D-C2D-C1D	-2.08	103.00	105.83
29	a	405	CLA	O2D-CGD-O1D	-2.08	119.78	123.84
29	N	602	CLA	CMB-C2B-C1B	-2.08	125.27	128.46
29	R	611	CLA	C1D-ND-C4D	-2.08	104.86	106.33
46	R	620	LUT	C40-C33-C34	-2.08	120.02	122.92
48	N	623	NEX	C31-C30-C29	2.08	130.27	127.31
29	n	614	CLA	C1-O2A-CGA	2.08	121.89	116.44
29	b	608	CLA	CMD-C2D-C3D	-2.08	122.84	127.61
46	r	620	LUT	C22-C23-C24	-2.08	109.38	111.74
29	C	502	CLA	C1D-ND-C4D	-2.07	104.86	106.33
29	G	604	CLA	C1D-ND-C4D	-2.07	104.86	106.33
29	b	607	CLA	O1D-CGD-CBD	-2.07	120.24	124.48
29	g	602	CLA	O1D-CGD-CBD	-2.07	120.24	124.48
29	G	603	CLA	O1D-CGD-CBD	-2.07	120.24	124.48
29	C	506	CLA	C3D-C2D-C1D	-2.07	103.00	105.83
46	r	620	LUT	C20-C13-C14	-2.07	120.02	122.92
29	N	613	CLA	CMD-C2D-C3D	-2.07	122.84	127.61
45	y	605	CHL	C4A-NA-C1A	2.07	107.64	106.71
29	s	604	CLA	CMA-C3A-C4A	2.07	117.34	111.77

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
29	c	512	CLA	CHA-C1A-NA	-2.07	121.65	126.40
29	B	610	CLA	C3D-C2D-C1D	-2.07	103.00	105.83
29	r	611	CLA	CAA-C2A-C3A	-2.07	107.11	112.78
29	G	610	CLA	C3D-C2D-C1D	-2.07	103.00	105.83
29	s	605	CLA	C2A-C1A-CHA	2.07	127.48	123.86
29	N	602	CLA	O1D-CGD-CBD	-2.07	120.25	124.48
45	N	601	CHL	C1-C2-C3	-2.07	122.46	126.04
42	d	405	PL9	C36-C34-C33	-2.07	116.93	121.12
46	s	621	LUT	C31-C32-C33	-2.07	120.60	126.42
46	Y	621	LUT	C2-C3-C4	-2.07	107.47	110.30
29	R	603	CLA	C1D-ND-C4D	-2.07	104.86	106.33
29	a	410	CLA	CMD-C2D-C3D	-2.07	122.85	127.61
29	A	405	CLA	O2D-CGD-O1D	-2.07	119.79	123.84
45	G	607	CHL	C1B-CHB-C4A	-2.07	126.02	130.12
41	d	401	BCT	O3-C-O1	-2.07	114.18	119.55
29	R	610	CLA	O2D-CGD-O1D	-2.07	119.79	123.84
46	N	621	LUT	C7-C8-C9	-2.07	123.11	126.23
29	Y	614	CLA	CAA-C2A-C3A	-2.07	107.11	112.78
29	C	504	CLA	C3D-C2D-C1D	-2.07	103.01	105.83
45	g	605	CHL	C1-O2A-CGA	2.07	122.74	116.73
29	b	607	CLA	C11-C10-C8	-2.07	109.23	115.92
29	r	604	CLA	O1D-CGD-CBD	-2.07	120.25	124.48
29	A	406	CLA	CHA-C1A-NA	-2.07	121.66	126.40
29	n	604	CLA	CHA-C1A-NA	-2.07	121.66	126.40
45	g	608	CHL	C1B-CHB-C4A	-2.07	126.02	130.12
29	g	613	CLA	CHA-C1A-NA	-2.07	121.66	126.40
29	N	603	CLA	O2A-CGA-CBA	2.07	118.39	111.91
48	S	622	NEX	C26-C27-C28	-2.07	121.62	125.99
35	b	620	C7Z	C18-C5-C4	2.07	118.18	114.36
29	d	402	CLA	CMD-C2D-C3D	-2.07	122.86	127.61
29	G	613	CLA	C1D-ND-C4D	-2.07	104.87	106.33
29	B	613	CLA	O1D-CGD-CBD	-2.07	120.26	124.48
29	B	614	CLA	CMD-C2D-C3D	-2.07	122.86	127.61
45	N	607	CHL	C1B-CHB-C4A	-2.07	126.03	130.12
29	c	511	CLA	C1-O2A-CGA	2.07	121.86	116.44
29	G	613	CLA	C1-O2A-CGA	2.07	121.86	116.44
29	B	604	CLA	CHA-C1A-NA	-2.06	121.67	126.40
29	C	508	CLA	O2D-CGD-O1D	-2.06	119.80	123.84
29	S	604	CLA	CMA-C3A-C4A	2.06	117.32	111.77
45	n	605	CHL	C4D-CHA-C1A	2.06	123.76	121.25
47	Y	622	XAT	C39-C29-C30	-2.06	120.03	122.92
35	B	620	C7Z	C2-C3-C4	2.06	113.13	110.30

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
29	a	406	CLA	C1D-ND-C4D	-2.06	104.87	106.33
29	b	616	CLA	C1D-ND-C4D	-2.06	104.87	106.33
29	r	602	CLA	C3D-C2D-C1D	-2.06	103.02	105.83
29	n	603	CLA	O2A-CGA-CBA	2.06	118.38	111.91
35	b	620	C7Z	C10-C11-C12	-2.06	116.78	123.22
45	g	601	CHL	CHD-C4C-C3C	2.06	127.87	124.84
29	n	614	CLA	CMD-C2D-C3D	-2.06	122.87	127.61
47	R	621	XAT	C39-C29-C28	2.06	121.33	118.08
29	b	610	CLA	CHA-C1A-NA	-2.06	121.68	126.40
29	g	610	CLA	CMB-C2B-C3B	2.06	128.53	124.68
29	Y	610	CLA	C3D-C2D-C1D	-2.06	103.02	105.83
42	d	405	PL9	C12-C13-C14	-2.06	122.70	127.66
29	c	508	CLA	CAA-C2A-C3A	-2.06	107.14	112.78
29	R	604	CLA	CHA-C1A-NA	-2.06	121.68	126.40
29	G	602	CLA	CHA-C1A-NA	-2.06	121.68	126.40
48	r	623	NEX	C20-C13-C14	-2.06	120.04	122.92
29	b	615	CLA	CHA-C1A-NA	-2.06	121.68	126.40
29	y	611	CLA	CMD-C2D-C3D	-2.06	122.88	127.61
29	n	602	CLA	CMC-C2C-C1C	2.06	128.17	125.04
29	B	615	CLA	CHA-C1A-NA	-2.06	121.68	126.40
29	y	603	CLA	CMD-C2D-C3D	-2.06	122.88	127.61
31	c	517	BCR	C34-C9-C10	-2.06	120.04	122.92
29	s	610	CLA	C1D-ND-C4D	-2.06	104.87	106.33
29	b	607	CLA	C1-C2-C3	-2.06	122.48	126.04
48	Y	623	NEX	C20-C13-C14	-2.06	120.04	122.92
29	s	613	CLA	O2A-CGA-CBA	2.06	118.36	111.91
45	G	607	CHL	CHD-C4C-C3C	2.06	127.86	124.84
29	N	610	CLA	O2D-CGD-O1D	-2.06	119.82	123.84
29	R	602	CLA	CHA-C1A-NA	-2.06	121.69	126.40
29	c	508	CLA	O2A-CGA-CBA	2.06	118.36	111.91
46	S	621	LUT	C20-C13-C12	2.06	121.32	118.08
46	r	620	LUT	C40-C33-C34	-2.06	120.04	122.92
48	n	623	NEX	C20-C13-C14	-2.06	120.04	122.92
29	c	501	CLA	CHA-C1A-NA	-2.06	121.69	126.40
29	b	617	CLA	CMB-C2B-C3B	2.06	128.52	124.68
29	C	503	CLA	C3D-C2D-C1D	-2.06	103.03	105.83
48	R	622	NEX	O24-C25-C38	-2.05	112.59	115.06
29	G	614	CLA	C3D-C2D-C1D	-2.05	103.03	105.83
29	S	617	CLA	C3D-C2D-C1D	-2.05	103.03	105.83
45	S	607	CHL	C1B-CHB-C4A	-2.05	126.05	130.12
29	S	610	CLA	C1D-ND-C4D	-2.05	104.88	106.33
31	d	404	BCR	C33-C5-C6	-2.05	122.22	124.53

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
45	Y	609	CHL	C4A-NA-C1A	2.05	107.63	106.71
29	a	407	CLA	CMD-C2D-C3D	-2.05	122.89	127.61
29	g	611	CLA	CMD-C2D-C3D	-2.05	122.89	127.61
29	G	611	CLA	C1D-ND-C4D	-2.05	104.88	106.33
29	b	602	CLA	CAA-C2A-C3A	-2.05	107.16	112.78
29	B	608	CLA	CMD-C2D-C3D	-2.05	122.89	127.61
29	a	405	CLA	C6-C5-C3	-2.05	108.08	113.45
29	G	610	CLA	O2D-CGD-O1D	-2.05	119.83	123.84
29	S	605	CLA	CAA-C2A-C3A	-2.05	107.16	112.78
29	G	611	CLA	O1D-CGD-CBD	-2.05	120.29	124.48
29	c	503	CLA	C3D-C2D-C1D	-2.05	103.03	105.83
48	y	623	NEX	C1-C2-C3	2.05	118.27	113.64
29	G	613	CLA	CHA-C1A-NA	-2.05	121.70	126.40
29	c	511	CLA	C1D-ND-C4D	-2.05	104.88	106.33
29	b	605	CLA	CHA-C1A-NA	-2.05	121.71	126.40
47	R	621	XAT	C12-C13-C14	2.05	122.08	118.94
29	C	501	CLA	CHA-C1A-NA	-2.05	121.71	126.40
46	S	621	LUT	C31-C30-C29	-2.05	124.39	127.31
29	S	604	CLA	CMB-C2B-C3B	2.05	128.51	124.68
29	S	617	CLA	C1D-ND-C4D	-2.05	104.88	106.33
29	y	610	CLA	C3D-C2D-C1D	-2.05	103.04	105.83
29	N	602	CLA	CMD-C2D-C3D	-2.05	122.91	127.61
46	y	620	LUT	C15-C35-C34	-2.05	119.28	123.47
29	C	504	CLA	CMA-C3A-C4A	2.05	117.27	111.77
29	D	402	CLA	C2C-C1C-NC	2.05	111.89	109.97
29	b	608	CLA	C3D-C2D-C1D	-2.05	103.04	105.83
29	y	602	CLA	CMD-C2D-C3D	-2.05	122.91	127.61
29	c	504	CLA	CMD-C2D-C3D	-2.05	122.91	127.61
31	C	516	BCR	C38-C26-C25	-2.04	122.23	124.53
29	N	602	CLA	CHA-C1A-NA	-2.04	121.72	126.40
29	Y	613	CLA	CHA-C1A-NA	-2.04	121.72	126.40
45	N	605	CHL	CHD-C4C-C3C	2.04	127.84	124.84
29	R	608	CLA	O2A-CGA-CBA	2.04	118.32	111.91
45	s	607	CHL	C4A-NA-C1A	2.04	107.62	106.71
29	s	609	CLA	CMD-C2D-C3D	-2.04	122.91	127.61
29	B	617	CLA	C1D-ND-C4D	-2.04	104.88	106.33
29	r	613	CLA	C3D-C2D-C1D	-2.04	103.04	105.83
29	B	605	CLA	CHA-C1A-NA	-2.04	121.72	126.40
29	g	612	CLA	CMD-C2D-C3D	-2.04	122.91	127.61
46	y	620	LUT	C35-C15-C14	-2.04	119.29	123.47
29	G	604	CLA	C3D-C2D-C1D	-2.04	103.04	105.83
29	Y	613	CLA	C1D-ND-C4D	-2.04	104.88	106.33

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
48	g	623	NEX	C40-C33-C34	-2.04	120.06	122.92
29	r	604	CLA	CHA-C1A-NA	-2.04	121.72	126.40
31	a	411	BCR	C19-C18-C17	2.04	122.07	118.94
29	y	613	CLA	CMB-C2B-C1B	-2.04	125.33	128.46
29	G	603	CLA	C3D-C2D-C1D	-2.04	103.05	105.83
29	c	513	CLA	C3D-C2D-C1D	-2.04	103.05	105.83
29	y	602	CLA	CAA-C2A-C3A	-2.04	107.19	112.78
46	g	621	LUT	C10-C11-C12	-2.04	116.85	123.22
45	G	607	CHL	C4D-CHA-C1A	2.04	123.73	121.25
29	G	611	CLA	C3D-C2D-C1D	-2.04	103.05	105.83
29	b	603	CLA	C3D-C2D-C1D	-2.04	103.05	105.83
45	S	601	CHL	C4D-CHA-C1A	2.04	123.73	121.25
29	N	611	CLA	CMD-C2D-C3D	-2.04	122.92	127.61
29	G	613	CLA	C3D-C2D-C1D	-2.04	103.05	105.83
29	a	405	CLA	CMD-C2D-C3D	-2.04	122.92	127.61
29	B	604	CLA	C1-C2-C3	-2.04	122.52	126.04
29	D	403	CLA	C1D-ND-C4D	-2.04	104.89	106.33
29	N	614	CLA	C1D-ND-C4D	-2.04	104.89	106.33
29	r	603	CLA	CMB-C2B-C3B	2.04	128.49	124.68
29	b	604	CLA	CAA-C2A-C3A	-2.04	107.20	112.78
29	R	612	CLA	O1D-CGD-CBD	-2.04	120.32	124.48
45	N	606	CHL	C4D-CHA-C1A	2.04	123.73	121.25
29	g	604	CLA	CMB-C2B-C3B	2.04	128.49	124.68
29	A	405	CLA	CAA-CBA-CGA	-2.04	107.30	113.25
29	G	604	CLA	OBD-CAD-C3D	-2.04	123.62	128.52
29	n	611	CLA	C1-O2A-CGA	2.03	121.78	116.44
29	B	615	CLA	C2A-C1A-CHA	2.03	127.42	123.86
29	n	603	CLA	CHA-C1A-NA	-2.03	121.74	126.40
29	R	603	CLA	O2D-CGD-O1D	-2.03	119.86	123.84
29	c	504	CLA	C3D-C2D-C1D	-2.03	103.06	105.83
29	c	508	CLA	CMD-C2D-C3D	-2.03	122.94	127.61
45	S	606	CHL	CHD-C4C-C3C	2.03	127.83	124.84
32	A	412	SQD	O5-C1-O6	-2.03	105.16	109.97
29	R	613	CLA	C3D-C2D-C1D	-2.03	103.06	105.83
29	g	602	CLA	C3D-C2D-C1D	-2.03	103.06	105.83
29	s	602	CLA	CHA-C1A-NA	-2.03	121.74	126.40
29	Y	603	CLA	C3D-C2D-C1D	-2.03	103.06	105.83
29	b	611	CLA	CMD-C2D-C3D	-2.03	122.94	127.61
29	r	608	CLA	O2A-CGA-CBA	2.03	118.28	111.91
48	S	622	NEX	C16-C1-C6	-2.03	108.65	110.47
29	n	604	CLA	C1D-ND-C4D	-2.03	104.89	106.33
29	S	609	CLA	C3D-C2D-C1D	-2.03	103.06	105.83

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
29	N	603	CLA	CBC-CAC-C3C	-2.03	106.83	112.43
45	s	601	CHL	C4A-NA-C1A	2.03	107.62	106.71
29	g	603	CLA	O1D-CGD-CBD	-2.03	120.33	124.48
29	s	612	CLA	CMD-C2D-C3D	-2.03	122.94	127.61
29	S	602	CLA	CMD-C2D-C3D	-2.03	122.94	127.61
30	a	409	PHO	C1-C2-C3	-2.03	122.53	126.04
29	s	602	CLA	CMA-C3A-C4A	2.03	117.23	111.77
29	y	602	CLA	CMB-C2B-C1B	-2.03	125.34	128.46
29	N	614	CLA	C3D-C2D-C1D	-2.03	103.06	105.83
29	c	512	CLA	C3D-C2D-C1D	-2.03	103.06	105.83
29	a	407	CLA	O2A-CGA-CBA	2.03	118.28	111.91
31	c	517	BCR	C33-C5-C4	2.03	117.52	113.62
29	n	602	CLA	C3D-C2D-C1D	-2.03	103.06	105.83
29	c	510	CLA	O1D-CGD-CBD	-2.03	120.33	124.48
29	Y	613	CLA	C3D-C2D-C1D	-2.03	103.06	105.83
31	B	618	BCR	C31-C1-C6	-2.03	107.01	110.30
29	N	604	CLA	C1D-ND-C4D	-2.03	104.89	106.33
29	D	402	CLA	CAC-C3C-C4C	2.03	127.44	124.81
29	N	603	CLA	C3D-C2D-C1D	-2.03	103.06	105.83
29	y	602	CLA	O1D-CGD-CBD	-2.03	120.34	124.48
29	c	504	CLA	CMA-C3A-C4A	2.03	117.22	111.77
29	B	610	CLA	CMD-C2D-C3D	-2.03	122.95	127.61
29	n	613	CLA	CHA-C1A-NA	-2.03	121.76	126.40
41	D	401	BCT	O3-C-O1	-2.03	114.29	119.55
29	B	604	CLA	CAA-C2A-C3A	-2.03	107.23	112.78
29	r	612	CLA	C1D-ND-C4D	-2.03	104.90	106.33
29	a	405	CLA	CHA-C1A-NA	-2.03	121.76	126.40
31	D	404	BCR	C38-C26-C25	-2.03	122.25	124.53
48	r	623	NEX	O24-C25-C38	-2.02	112.63	115.06
29	y	612	CLA	C1D-ND-C4D	-2.02	104.90	106.33
29	G	614	CLA	CHA-C1A-NA	-2.02	121.76	126.40
29	y	602	CLA	CHA-C1A-NA	-2.02	121.76	126.40
45	R	606	CHL	CHC-C1C-NC	2.02	127.27	124.20
29	Y	602	CLA	CHA-C1A-NA	-2.02	121.76	126.40
46	G	621	LUT	C20-C13-C12	2.02	121.27	118.08
35	B	620	C7Z	C18-C5-C4	2.02	118.10	114.36
29	d	403	CLA	C1D-ND-C4D	-2.02	104.90	106.33
42	D	405	PL9	O2-C1-C6	2.02	124.09	120.59
29	B	602	CLA	CAA-C2A-C3A	-2.02	107.24	112.78
29	b	613	CLA	O1D-CGD-CBD	-2.02	120.35	124.48
29	r	612	CLA	C3D-C2D-C1D	-2.02	103.07	105.83
29	b	616	CLA	CMA-C3A-C4A	2.02	117.20	111.77

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
29	n	610	CLA	C3D-C2D-C1D	-2.02	103.07	105.83
45	s	606	CHL	CHC-C1C-NC	2.02	127.27	124.20
29	b	610	CLA	CMD-C2D-C3D	-2.02	122.97	127.61
29	B	609	CLA	CHA-C1A-NA	-2.02	121.77	126.40
29	C	513	CLA	CHA-C1A-NA	-2.02	121.77	126.40
45	N	608	CHL	CHD-C4C-C3C	2.02	127.81	124.84
31	A	411	BCR	C37-C22-C23	2.02	121.26	118.08
29	n	611	CLA	C3D-C2D-C1D	-2.02	103.08	105.83
44	h	101	RRX	C35-C13-C14	-2.02	120.10	122.92
31	c	517	BCR	C29-C30-C25	2.02	113.59	110.48
29	Y	614	CLA	C1D-ND-C4D	-2.02	104.90	106.33
39	n	624	LHG	O7-C7-O9	-2.02	118.83	123.70
29	G	602	CLA	O1D-CGD-CBD	-2.02	120.36	124.48
29	C	509	CLA	CMB-C2B-C1B	-2.02	125.36	128.46
42	d	405	PL9	O2-C1-C2	-2.02	117.16	121.78
29	Y	604	CLA	CHA-C1A-NA	-2.02	121.78	126.40
48	s	623	NEX	C1-C2-C3	2.02	118.19	113.64
29	s	617	CLA	CMB-C2B-C3B	2.01	128.45	124.68
29	C	505	CLA	CMB-C2B-C3B	2.01	128.44	124.68
46	n	620	LUT	C8-C7-C6	-2.01	121.55	127.20
29	C	507	CLA	CMB-C2B-C3B	2.01	128.44	124.68
46	N	620	LUT	C39-C29-C28	2.01	121.25	118.08
45	n	607	CHL	CHA-C1A-NA	-2.01	121.79	126.40
29	C	502	CLA	CAC-C3C-C4C	2.01	127.42	124.81
29	y	614	CLA	C1D-ND-C4D	-2.01	104.91	106.33
29	s	612	CLA	C3D-C2D-C1D	-2.01	103.09	105.83
46	S	620	LUT	C10-C11-C12	-2.01	116.94	123.22
30	a	408	PHO	C1B-NB-C4B	2.01	111.22	107.09
29	R	608	CLA	C3D-C2D-C1D	-2.01	103.09	105.83
29	n	614	CLA	CMB-C2B-C3B	2.01	128.44	124.68
29	Y	608	CLA	CAA-C2A-C3A	-2.01	107.27	112.78
29	g	602	CLA	CMD-C2D-C3D	-2.01	122.99	127.61
35	B	620	C7Z	C4-C5-C6	-2.01	116.37	120.85
29	Y	608	CLA	C1D-ND-C4D	-2.01	104.91	106.33
29	S	610	CLA	O2D-CGD-O1D	-2.01	119.91	123.84
29	Y	610	CLA	O2D-CGD-O1D	-2.01	119.91	123.84
29	s	617	CLA	C3D-C2D-C1D	-2.01	103.09	105.83
29	n	604	CLA	OBD-CAD-C3D	-2.01	123.68	128.52
29	Y	612	CLA	C3D-C2D-C1D	-2.01	103.09	105.83
29	g	613	CLA	CAA-C2A-C3A	-2.01	107.28	112.78
29	A	407	CLA	C3D-C2D-C1D	-2.01	103.09	105.83
42	D	405	PL9	C36-C34-C33	-2.01	117.05	121.12

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
29	c	506	CLA	C1-O2A-CGA	2.01	121.71	116.44
38	C	519	DGD	O2G-C1B-O1B	-2.01	118.85	123.70
29	b	614	CLA	CMB-C2B-C1B	-2.01	125.38	128.46
48	G	623	NEX	C26-C27-C28	-2.01	121.75	125.99
30	a	409	PHO	C1B-NB-C4B	2.01	111.21	107.09
45	n	609	CHL	CHA-C1A-NA	-2.01	121.80	126.40
45	s	601	CHL	C1B-CHB-C4A	-2.01	126.14	130.12
29	Y	612	CLA	O1D-CGD-CBD	-2.01	120.38	124.48
29	g	603	CLA	O2A-CGA-CBA	2.01	118.20	111.91
31	C	515	BCR	C3-C4-C5	-2.01	110.50	114.08
42	D	405	PL9	O2-C1-C2	-2.01	117.18	121.78
29	S	613	CLA	C2A-C1A-CHA	2.01	127.37	123.86
29	c	507	CLA	C2A-C1A-CHA	2.01	127.37	123.86
31	c	514	BCR	C29-C28-C27	2.00	115.86	111.38
29	R	611	CLA	CAA-C2A-C3A	-2.00	107.29	112.78
46	n	621	LUT	C11-C10-C9	-2.00	124.45	127.31
31	A	411	BCR	C12-C13-C14	-2.00	115.86	118.94
31	A	411	BCR	C19-C18-C17	2.00	122.02	118.94
29	a	407	CLA	C1-O2A-CGA	2.00	121.70	116.44
29	R	604	CLA	C3D-C2D-C1D	-2.00	103.10	105.83
45	G	608	CHL	C4D-CHA-C1A	2.00	123.69	121.25
31	B	619	BCR	C30-C25-C26	-2.00	119.79	122.61
47	y	622	XAT	C26-C27-C28	-2.00	121.76	125.99
29	C	509	CLA	O1D-CGD-CBD	-2.00	120.39	124.48
29	b	616	CLA	C2A-C1A-CHA	2.00	127.36	123.86
29	b	613	CLA	CHA-C1A-NA	-2.00	121.81	126.40
48	n	623	NEX	O24-C25-C38	-2.00	112.66	115.06
29	N	610	CLA	C3D-C2D-C1D	-2.00	103.10	105.83
46	s	621	LUT	C39-C29-C28	2.00	121.23	118.08
29	R	609	CLA	CAA-C2A-C3A	-2.00	107.30	112.78
47	Y	622	XAT	C40-C33-C34	-2.00	120.12	122.92
45	s	601	CHL	C4D-CHA-C1A	2.00	123.68	121.25

All (343) chirality outliers are listed below:

Mol	Chain	Res	Type	Atom
29	A	405	CLA	ND
29	A	406	CLA	ND
29	A	407	CLA	ND
29	A	410	CLA	ND
29	B	602	CLA	ND
29	B	603	CLA	ND

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Mol	Chain	Res	Type	Atom
29	B	604	CLA	ND
29	B	605	CLA	ND
29	B	606	CLA	ND
29	B	607	CLA	ND
29	B	608	CLA	ND
29	B	609	CLA	ND
29	B	610	CLA	ND
29	B	611	CLA	ND
29	B	612	CLA	ND
29	B	613	CLA	ND
29	B	614	CLA	ND
29	B	615	CLA	ND
29	B	616	CLA	ND
29	B	617	CLA	ND
29	C	501	CLA	ND
29	C	502	CLA	ND
29	C	503	CLA	ND
29	C	504	CLA	ND
29	C	505	CLA	ND
29	C	506	CLA	ND
29	C	507	CLA	ND
29	C	508	CLA	ND
29	C	509	CLA	ND
29	C	510	CLA	ND
29	C	511	CLA	ND
29	C	512	CLA	ND
29	C	513	CLA	ND
29	D	402	CLA	ND
29	D	403	CLA	ND
29	N	602	CLA	ND
29	N	603	CLA	ND
29	N	604	CLA	ND
29	N	610	CLA	ND
29	N	611	CLA	ND
29	N	612	CLA	ND
29	N	613	CLA	ND
29	N	614	CLA	ND
29	G	602	CLA	ND
29	G	603	CLA	ND
29	G	604	CLA	ND
29	G	610	CLA	ND
29	G	611	CLA	ND

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Mol	Chain	Res	Type	Atom
29	G	612	CLA	ND
29	G	613	CLA	ND
29	G	614	CLA	ND
29	R	602	CLA	ND
29	R	603	CLA	ND
29	R	604	CLA	ND
29	R	608	CLA	ND
29	R	609	CLA	ND
29	R	610	CLA	ND
29	R	611	CLA	ND
29	R	612	CLA	ND
29	R	613	CLA	ND
29	S	602	CLA	ND
29	S	603	CLA	ND
29	S	604	CLA	ND
29	S	605	CLA	ND
29	S	609	CLA	ND
29	S	610	CLA	ND
29	S	611	CLA	ND
29	S	612	CLA	ND
29	S	613	CLA	ND
29	S	614	CLA	ND
29	S	617	CLA	ND
29	Y	602	CLA	ND
29	Y	603	CLA	ND
29	Y	604	CLA	ND
29	Y	608	CLA	ND
29	Y	610	CLA	ND
29	Y	611	CLA	ND
29	Y	612	CLA	ND
29	Y	613	CLA	ND
29	Y	614	CLA	ND
29	a	405	CLA	ND
29	a	406	CLA	ND
29	a	407	CLA	ND
29	a	410	CLA	ND
29	b	602	CLA	ND
29	b	603	CLA	ND
29	b	604	CLA	ND
29	b	605	CLA	ND
29	b	606	CLA	ND
29	b	607	CLA	ND

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Mol	Chain	Res	Type	Atom
29	b	608	CLA	ND
29	b	609	CLA	ND
29	b	610	CLA	ND
29	b	611	CLA	ND
29	b	612	CLA	ND
29	b	613	CLA	ND
29	b	614	CLA	ND
29	b	615	CLA	ND
29	b	616	CLA	ND
29	b	617	CLA	ND
29	c	501	CLA	ND
29	c	502	CLA	ND
29	c	503	CLA	ND
29	c	504	CLA	ND
29	c	505	CLA	ND
29	c	506	CLA	ND
29	c	507	CLA	ND
29	c	508	CLA	ND
29	c	509	CLA	ND
29	c	510	CLA	ND
29	c	511	CLA	ND
29	c	512	CLA	ND
29	c	513	CLA	ND
29	d	402	CLA	ND
29	d	403	CLA	ND
29	n	602	CLA	ND
29	n	603	CLA	ND
29	n	604	CLA	ND
29	n	610	CLA	ND
29	n	611	CLA	ND
29	n	612	CLA	ND
29	n	613	CLA	ND
29	n	614	CLA	ND
29	g	602	CLA	ND
29	g	603	CLA	ND
29	g	604	CLA	ND
29	g	610	CLA	ND
29	g	611	CLA	ND
29	g	612	CLA	ND
29	g	613	CLA	ND
29	g	614	CLA	ND
29	r	602	CLA	ND

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Mol	Chain	Res	Type	Atom
29	r	603	CLA	ND
29	r	604	CLA	ND
29	r	608	CLA	ND
29	r	609	CLA	ND
29	r	610	CLA	ND
29	r	611	CLA	ND
29	r	612	CLA	ND
29	r	613	CLA	ND
29	s	602	CLA	ND
29	s	603	CLA	ND
29	s	604	CLA	ND
29	s	605	CLA	ND
29	s	609	CLA	ND
29	s	610	CLA	ND
29	s	611	CLA	ND
29	s	612	CLA	ND
29	s	613	CLA	ND
29	s	614	CLA	ND
29	s	617	CLA	ND
29	y	602	CLA	ND
29	y	603	CLA	ND
29	y	604	CLA	ND
29	y	608	CLA	ND
29	y	610	CLA	ND
29	y	611	CLA	ND
29	y	612	CLA	ND
29	y	613	CLA	ND
29	y	614	CLA	ND
35	B	620	C7Z	C3
35	b	620	C7Z	C3
40	C	527	LMK	C3
40	c	627	LMK	C3
44	H	101	RRX	C28
44	h	101	RRX	C28
45	N	601	CHL	NA
45	N	601	CHL	C8
45	N	601	CHL	NC
45	N	601	CHL	ND
45	N	605	CHL	NA
45	N	605	CHL	C8
45	N	605	CHL	NC
45	N	605	CHL	ND

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Mol	Chain	Res	Type	Atom
45	N	606	CHL	NA
45	N	606	CHL	C8
45	N	606	CHL	NC
45	N	606	CHL	ND
45	N	607	CHL	NA
45	N	607	CHL	C8
45	N	607	CHL	NC
45	N	607	CHL	ND
45	N	608	CHL	NA
45	N	608	CHL	NC
45	N	608	CHL	ND
45	N	609	CHL	NA
45	N	609	CHL	C8
45	N	609	CHL	NC
45	N	609	CHL	ND
45	G	601	CHL	NA
45	G	601	CHL	C8
45	G	601	CHL	NC
45	G	601	CHL	ND
45	G	605	CHL	NA
45	G	605	CHL	NC
45	G	605	CHL	ND
45	G	606	CHL	NA
45	G	606	CHL	NC
45	G	606	CHL	C3A
45	G	606	CHL	ND
45	G	607	CHL	NA
45	G	607	CHL	NC
45	G	607	CHL	ND
45	G	608	CHL	NA
45	G	608	CHL	NC
45	G	608	CHL	ND
45	G	609	CHL	NA
45	G	609	CHL	C8
45	G	609	CHL	NC
45	G	609	CHL	ND
45	R	606	CHL	NA
45	R	606	CHL	NC
45	R	606	CHL	ND
45	R	607	CHL	NA
45	R	607	CHL	NC
45	R	607	CHL	ND

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Mol	Chain	Res	Type	Atom
45	S	601	CHL	NA
45	S	601	CHL	NC
45	S	601	CHL	ND
45	S	606	CHL	NA
45	S	606	CHL	NC
45	S	606	CHL	ND
45	S	607	CHL	NA
45	S	607	CHL	NC
45	S	607	CHL	C3A
45	S	607	CHL	ND
45	S	608	CHL	NA
45	S	608	CHL	C8
45	S	608	CHL	NC
45	S	608	CHL	ND
45	Y	601	CHL	NA
45	Y	601	CHL	C8
45	Y	601	CHL	NC
45	Y	601	CHL	ND
45	Y	605	CHL	NA
45	Y	605	CHL	NC
45	Y	605	CHL	ND
45	Y	606	CHL	NA
45	Y	606	CHL	C8
45	Y	606	CHL	NC
45	Y	606	CHL	ND
45	Y	607	CHL	NA
45	Y	607	CHL	C8
45	Y	607	CHL	NC
45	Y	607	CHL	ND
45	Y	609	CHL	NA
45	Y	609	CHL	C8
45	Y	609	CHL	NC
45	Y	609	CHL	ND
45	n	601	CHL	NA
45	n	601	CHL	C8
45	n	601	CHL	NC
45	n	601	CHL	ND
45	n	605	CHL	NA
45	n	605	CHL	C8
45	n	605	CHL	NC
45	n	605	CHL	ND
45	n	606	CHL	NA

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Mol	Chain	Res	Type	Atom
45	n	606	CHL	C8
45	n	606	CHL	NC
45	n	606	CHL	ND
45	n	607	CHL	NA
45	n	607	CHL	C8
45	n	607	CHL	NC
45	n	607	CHL	ND
45	n	608	CHL	NA
45	n	608	CHL	NC
45	n	608	CHL	ND
45	n	609	CHL	NA
45	n	609	CHL	C8
45	n	609	CHL	NC
45	n	609	CHL	ND
45	g	601	CHL	NA
45	g	601	CHL	C8
45	g	601	CHL	NC
45	g	601	CHL	ND
45	g	605	CHL	NA
45	g	605	CHL	NC
45	g	605	CHL	ND
45	g	606	CHL	NA
45	g	606	CHL	NC
45	g	606	CHL	ND
45	g	607	CHL	NA
45	g	607	CHL	NC
45	g	607	CHL	ND
45	g	608	CHL	NA
45	g	608	CHL	NC
45	g	608	CHL	ND
45	g	609	CHL	NA
45	g	609	CHL	C8
45	g	609	CHL	NC
45	g	609	CHL	ND
45	r	606	CHL	NA
45	r	606	CHL	NC
45	r	606	CHL	ND
45	r	607	CHL	NA
45	r	607	CHL	NC
45	r	607	CHL	ND
45	s	601	CHL	NA
45	s	601	CHL	NC

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Mol	Chain	Res	Type	Atom
45	s	601	CHL	ND
45	s	606	CHL	NA
45	s	606	CHL	NC
45	s	606	CHL	ND
45	s	607	CHL	NA
45	s	607	CHL	NC
45	s	607	CHL	C3A
45	s	607	CHL	ND
45	s	608	CHL	NA
45	s	608	CHL	C8
45	s	608	CHL	NC
45	s	608	CHL	ND
45	y	601	CHL	NA
45	y	601	CHL	C8
45	y	601	CHL	NC
45	y	601	CHL	ND
45	y	605	CHL	NA
45	y	605	CHL	NC
45	y	605	CHL	ND
45	y	606	CHL	NA
45	y	606	CHL	C8
45	y	606	CHL	NC
45	y	606	CHL	ND
45	y	607	CHL	NA
45	y	607	CHL	C8
45	y	607	CHL	NC
45	y	607	CHL	ND
45	y	609	CHL	NA
45	y	609	CHL	C8
45	y	609	CHL	NC
45	y	609	CHL	ND
46	S	620	LUT	C26
46	s	620	LUT	C26
47	N	622	XAT	C6
47	G	622	XAT	C26
47	G	622	XAT	C6
47	R	621	XAT	C26
47	Y	622	XAT	C6
47	n	622	XAT	C6
47	g	622	XAT	C26
47	g	622	XAT	C6
47	r	622	XAT	C26

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Mol	Chain	Res	Type	Atom
47	y	622	XAT	C6

All (3918) torsion outliers are listed below:

Mol	Chain	Res	Type	Atoms
29	A	405	CLA	CBD-CGD-O2D-CED
29	A	406	CLA	C1A-C2A-CAA-CBA
29	A	406	CLA	C3A-C2A-CAA-CBA
29	A	406	CLA	CHA-CBD-CGD-O1D
29	A	406	CLA	CHA-CBD-CGD-O2D
29	A	407	CLA	CHA-CBD-CGD-O1D
29	A	407	CLA	CHA-CBD-CGD-O2D
29	B	602	CLA	CHA-CBD-CGD-O1D
29	B	602	CLA	CHA-CBD-CGD-O2D
29	B	602	CLA	CBD-CGD-O2D-CED
29	B	603	CLA	CHA-CBD-CGD-O1D
29	B	603	CLA	CHA-CBD-CGD-O2D
29	B	603	CLA	CBD-CGD-O2D-CED
29	B	604	CLA	CBD-CGD-O2D-CED
29	B	604	CLA	C2-C3-C5-C6
29	B	604	CLA	C4-C3-C5-C6
29	B	605	CLA	C2-C1-O2A-CGA
29	B	605	CLA	CHA-CBD-CGD-O1D
29	B	605	CLA	CHA-CBD-CGD-O2D
29	B	605	CLA	CAD-CBD-CGD-O1D
29	B	605	CLA	CAD-CBD-CGD-O2D
29	B	605	CLA	C2-C3-C5-C6
29	B	605	CLA	C4-C3-C5-C6
29	B	606	CLA	C2-C3-C5-C6
29	B	606	CLA	C4-C3-C5-C6
29	B	607	CLA	CHA-CBD-CGD-O1D
29	B	607	CLA	CHA-CBD-CGD-O2D
29	B	607	CLA	CBD-CGD-O2D-CED
29	B	608	CLA	C1A-C2A-CAA-CBA
29	B	608	CLA	C3A-C2A-CAA-CBA
29	B	608	CLA	CHA-CBD-CGD-O1D
29	B	608	CLA	CHA-CBD-CGD-O2D
29	B	608	CLA	CAD-CBD-CGD-O1D
29	B	608	CLA	CAD-CBD-CGD-O2D
29	B	608	CLA	CBD-CGD-O2D-CED
29	B	608	CLA	C2-C3-C5-C6
29	B	608	CLA	C4-C3-C5-C6

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Mol	Chain	Res	Type	Atoms
29	B	609	CLA	C1A-C2A-CAA-CBA
29	B	609	CLA	CHA-CBD-CGD-O1D
29	B	609	CLA	CHA-CBD-CGD-O2D
29	B	610	CLA	C1A-C2A-CAA-CBA
29	B	610	CLA	C3A-C2A-CAA-CBA
29	B	610	CLA	CHA-CBD-CGD-O1D
29	B	610	CLA	CHA-CBD-CGD-O2D
29	B	610	CLA	CAD-CBD-CGD-O1D
29	B	610	CLA	CBD-CGD-O2D-CED
29	B	612	CLA	C1A-C2A-CAA-CBA
29	B	612	CLA	CHA-CBD-CGD-O1D
29	B	612	CLA	CHA-CBD-CGD-O2D
29	B	612	CLA	CBD-CGD-O2D-CED
29	B	612	CLA	C2-C3-C5-C6
29	B	612	CLA	C4-C3-C5-C6
29	B	613	CLA	C1A-C2A-CAA-CBA
29	B	613	CLA	C2-C1-O2A-CGA
29	B	614	CLA	C2-C1-O2A-CGA
29	B	614	CLA	CBD-CGD-O2D-CED
29	B	615	CLA	C1A-C2A-CAA-CBA
29	B	615	CLA	CAD-CBD-CGD-O1D
29	B	615	CLA	CAD-CBD-CGD-O2D
29	B	617	CLA	O1A-CGA-O2A-C1
29	B	617	CLA	CBD-CGD-O2D-CED
29	C	501	CLA	CHA-CBD-CGD-O1D
29	C	501	CLA	CHA-CBD-CGD-O2D
29	C	501	CLA	C2-C3-C5-C6
29	C	501	CLA	C4-C3-C5-C6
29	C	502	CLA	CHA-CBD-CGD-O1D
29	C	502	CLA	CHA-CBD-CGD-O2D
29	C	502	CLA	CAD-CBD-CGD-O1D
29	C	503	CLA	CBD-CGD-O2D-CED
29	C	503	CLA	C2-C3-C5-C6
29	C	503	CLA	C4-C3-C5-C6
29	C	504	CLA	CHA-CBD-CGD-O1D
29	C	504	CLA	CHA-CBD-CGD-O2D
29	C	504	CLA	C2-C3-C5-C6
29	C	504	CLA	C4-C3-C5-C6
29	C	505	CLA	CAD-CBD-CGD-O1D
29	C	505	CLA	CAD-CBD-CGD-O2D
29	C	506	CLA	C1A-C2A-CAA-CBA
29	C	507	CLA	CBD-CGD-O2D-CED

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Mol	Chain	Res	Type	Atoms
29	C	508	CLA	CHA-CBD-CGD-O1D
29	C	508	CLA	CHA-CBD-CGD-O2D
29	C	509	CLA	C2-C1-O2A-CGA
29	C	511	CLA	C1A-C2A-CAA-CBA
29	C	511	CLA	CBD-CGD-O2D-CED
29	C	512	CLA	CHA-CBD-CGD-O1D
29	C	512	CLA	CHA-CBD-CGD-O2D
29	C	513	CLA	C1A-C2A-CAA-CBA
29	C	513	CLA	C2-C3-C5-C6
29	C	513	CLA	C4-C3-C5-C6
29	D	403	CLA	CBD-CGD-O2D-CED
29	N	610	CLA	C1A-C2A-CAA-CBA
29	N	610	CLA	CHA-CBD-CGD-O1D
29	N	610	CLA	CHA-CBD-CGD-O2D
29	N	610	CLA	CBD-CGD-O2D-CED
29	N	611	CLA	CHA-CBD-CGD-O1D
29	N	611	CLA	CHA-CBD-CGD-O2D
29	N	612	CLA	C1A-C2A-CAA-CBA
29	N	612	CLA	C3A-C2A-CAA-CBA
29	N	613	CLA	CHA-CBD-CGD-O1D
29	N	613	CLA	CHA-CBD-CGD-O2D
29	G	602	CLA	C1A-C2A-CAA-CBA
29	G	602	CLA	C3A-C2A-CAA-CBA
29	G	603	CLA	CBD-CGD-O2D-CED
29	G	604	CLA	C1A-C2A-CAA-CBA
29	G	610	CLA	CBD-CGD-O2D-CED
29	G	614	CLA	CBD-CGD-O2D-CED
29	R	603	CLA	C2-C1-O2A-CGA
29	R	604	CLA	C1A-C2A-CAA-CBA
29	R	608	CLA	C3A-C2A-CAA-CBA
29	R	608	CLA	CHA-CBD-CGD-O1D
29	R	608	CLA	CHA-CBD-CGD-O2D
29	R	609	CLA	CBD-CGD-O2D-CED
29	R	610	CLA	C1A-C2A-CAA-CBA
29	R	610	CLA	CHA-CBD-CGD-O1D
29	R	610	CLA	CHA-CBD-CGD-O2D
29	R	611	CLA	CBA-CGA-O2A-C1
29	R	611	CLA	O1A-CGA-O2A-C1
29	R	612	CLA	C2-C1-O2A-CGA
29	R	613	CLA	CBA-CGA-O2A-C1
29	R	613	CLA	CHA-CBD-CGD-O1D
29	R	613	CLA	CHA-CBD-CGD-O2D

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Mol	Chain	Res	Type	Atoms
29	S	602	CLA	C3A-C2A-CAA-CBA
29	S	602	CLA	CHA-CBD-CGD-O1D
29	S	602	CLA	CHA-CBD-CGD-O2D
29	S	603	CLA	CBD-CGD-O2D-CED
29	S	604	CLA	CHA-CBD-CGD-O1D
29	S	604	CLA	CHA-CBD-CGD-O2D
29	S	605	CLA	C1A-C2A-CAA-CBA
29	S	605	CLA	C2-C1-O2A-CGA
29	S	605	CLA	CHA-CBD-CGD-O1D
29	S	605	CLA	CHA-CBD-CGD-O2D
29	S	609	CLA	CBD-CGD-O2D-CED
29	S	610	CLA	CHA-CBD-CGD-O1D
29	S	610	CLA	CHA-CBD-CGD-O2D
29	S	610	CLA	CBD-CGD-O2D-CED
29	S	611	CLA	CHA-CBD-CGD-O1D
29	S	611	CLA	CHA-CBD-CGD-O2D
29	S	612	CLA	CBD-CGD-O2D-CED
29	S	613	CLA	CHA-CBD-CGD-O1D
29	S	613	CLA	CHA-CBD-CGD-O2D
29	S	617	CLA	CBD-CGD-O2D-CED
29	Y	603	CLA	C2-C1-O2A-CGA
29	Y	604	CLA	CBD-CGD-O2D-CED
29	Y	610	CLA	CBD-CGD-O2D-CED
29	Y	611	CLA	CHA-CBD-CGD-O1D
29	Y	611	CLA	CHA-CBD-CGD-O2D
29	Y	613	CLA	CHA-CBD-CGD-O1D
29	Y	613	CLA	CHA-CBD-CGD-O2D
29	Y	614	CLA	CBD-CGD-O2D-CED
29	a	405	CLA	CBD-CGD-O2D-CED
29	a	406	CLA	C1A-C2A-CAA-CBA
29	a	406	CLA	C3A-C2A-CAA-CBA
29	a	406	CLA	CHA-CBD-CGD-O1D
29	a	406	CLA	CHA-CBD-CGD-O2D
29	a	407	CLA	CHA-CBD-CGD-O1D
29	a	407	CLA	CHA-CBD-CGD-O2D
29	b	602	CLA	CHA-CBD-CGD-O1D
29	b	602	CLA	CHA-CBD-CGD-O2D
29	b	602	CLA	CBD-CGD-O2D-CED
29	b	603	CLA	CHA-CBD-CGD-O1D
29	b	603	CLA	CHA-CBD-CGD-O2D
29	b	604	CLA	CBD-CGD-O2D-CED
29	b	604	CLA	C4-C3-C5-C6

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Mol	Chain	Res	Type	Atoms
29	b	605	CLA	CHA-CBD-CGD-O1D
29	b	605	CLA	CHA-CBD-CGD-O2D
29	b	605	CLA	CAD-CBD-CGD-O1D
29	b	605	CLA	CAD-CBD-CGD-O2D
29	b	605	CLA	C2-C3-C5-C6
29	b	605	CLA	C4-C3-C5-C6
29	b	606	CLA	C2-C3-C5-C6
29	b	606	CLA	C4-C3-C5-C6
29	b	607	CLA	CHA-CBD-CGD-O1D
29	b	607	CLA	CHA-CBD-CGD-O2D
29	b	607	CLA	CBD-CGD-O2D-CED
29	b	608	CLA	C1A-C2A-CAA-CBA
29	b	608	CLA	C3A-C2A-CAA-CBA
29	b	608	CLA	CHA-CBD-CGD-O1D
29	b	608	CLA	CHA-CBD-CGD-O2D
29	b	608	CLA	CAD-CBD-CGD-O1D
29	b	608	CLA	CAD-CBD-CGD-O2D
29	b	608	CLA	CBD-CGD-O2D-CED
29	b	608	CLA	C2-C3-C5-C6
29	b	608	CLA	C4-C3-C5-C6
29	b	609	CLA	C1A-C2A-CAA-CBA
29	b	609	CLA	C3A-C2A-CAA-CBA
29	b	609	CLA	CBD-CGD-O2D-CED
29	b	610	CLA	C3A-C2A-CAA-CBA
29	b	610	CLA	CHA-CBD-CGD-O1D
29	b	610	CLA	CHA-CBD-CGD-O2D
29	b	610	CLA	CAD-CBD-CGD-O1D
29	b	610	CLA	CBD-CGD-O2D-CED
29	b	612	CLA	C1A-C2A-CAA-CBA
29	b	612	CLA	CHA-CBD-CGD-O1D
29	b	612	CLA	CHA-CBD-CGD-O2D
29	b	613	CLA	C2-C1-O2A-CGA
29	b	615	CLA	C1A-C2A-CAA-CBA
29	b	615	CLA	CAD-CBD-CGD-O1D
29	b	615	CLA	CAD-CBD-CGD-O2D
29	b	616	CLA	CHA-CBD-CGD-O1D
29	b	616	CLA	CHA-CBD-CGD-O2D
29	b	617	CLA	CBD-CGD-O2D-CED
29	c	501	CLA	C1A-C2A-CAA-CBA
29	c	501	CLA	CHA-CBD-CGD-O1D
29	c	501	CLA	CHA-CBD-CGD-O2D
29	c	502	CLA	CBD-CGD-O2D-CED

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Mol	Chain	Res	Type	Atoms
29	c	503	CLA	C4-C3-C5-C6
29	c	504	CLA	CHA-CBD-CGD-O1D
29	c	504	CLA	CHA-CBD-CGD-O2D
29	c	504	CLA	CAD-CBD-CGD-O1D
29	c	504	CLA	C2-C3-C5-C6
29	c	504	CLA	C4-C3-C5-C6
29	c	505	CLA	CAD-CBD-CGD-O1D
29	c	505	CLA	CAD-CBD-CGD-O2D
29	c	506	CLA	C1A-C2A-CAA-CBA
29	c	508	CLA	CHA-CBD-CGD-O1D
29	c	508	CLA	CHA-CBD-CGD-O2D
29	c	509	CLA	C2-C1-O2A-CGA
29	c	511	CLA	C1A-C2A-CAA-CBA
29	c	511	CLA	CBD-CGD-O2D-CED
29	c	512	CLA	CHA-CBD-CGD-O1D
29	c	513	CLA	C1A-C2A-CAA-CBA
29	c	513	CLA	C2-C3-C5-C6
29	c	513	CLA	C4-C3-C5-C6
29	d	403	CLA	CBD-CGD-O2D-CED
29	n	603	CLA	C3A-C2A-CAA-CBA
29	n	603	CLA	CBA-CGA-O2A-C1
29	n	603	CLA	O1A-CGA-O2A-C1
29	n	603	CLA	CHA-CBD-CGD-O1D
29	n	603	CLA	CHA-CBD-CGD-O2D
29	n	603	CLA	CAD-CBD-CGD-O1D
29	n	603	CLA	C2-C3-C5-C6
29	n	603	CLA	C4-C3-C5-C6
29	n	604	CLA	C1A-C2A-CAA-CBA
29	n	610	CLA	C1A-C2A-CAA-CBA
29	n	610	CLA	CHA-CBD-CGD-O1D
29	n	610	CLA	CHA-CBD-CGD-O2D
29	n	610	CLA	CBD-CGD-O2D-CED
29	n	611	CLA	C1A-C2A-CAA-CBA
29	n	611	CLA	C3A-C2A-CAA-CBA
29	n	611	CLA	CHA-CBD-CGD-O1D
29	n	611	CLA	CHA-CBD-CGD-O2D
29	n	612	CLA	CBD-CGD-O2D-CED
29	n	613	CLA	CHA-CBD-CGD-O1D
29	n	613	CLA	CHA-CBD-CGD-O2D
29	n	613	CLA	CBD-CGD-O2D-CED
29	n	614	CLA	C1A-C2A-CAA-CBA
29	n	614	CLA	CBD-CGD-O2D-CED

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Mol	Chain	Res	Type	Atoms
29	g	602	CLA	C1A-C2A-CAA-CBA
29	g	602	CLA	C3A-C2A-CAA-CBA
29	g	610	CLA	CHA-CBD-CGD-O2D
29	g	610	CLA	CBD-CGD-O2D-CED
29	g	613	CLA	CHA-CBD-CGD-O1D
29	g	613	CLA	CHA-CBD-CGD-O2D
29	g	614	CLA	C2A-CAA-CBA-CGA
29	g	614	CLA	CBD-CGD-O2D-CED
29	r	608	CLA	C3A-C2A-CAA-CBA
29	r	608	CLA	CHA-CBD-CGD-O1D
29	r	608	CLA	CHA-CBD-CGD-O2D
29	r	609	CLA	CBD-CGD-O2D-CED
29	r	610	CLA	C1A-C2A-CAA-CBA
29	r	610	CLA	C2-C3-C5-C6
29	r	610	CLA	C4-C3-C5-C6
29	r	612	CLA	C2-C1-O2A-CGA
29	r	613	CLA	CBA-CGA-O2A-C1
29	r	613	CLA	CBD-CGD-O2D-CED
29	s	602	CLA	C1A-C2A-CAA-CBA
29	s	602	CLA	C3A-C2A-CAA-CBA
29	s	602	CLA	CHA-CBD-CGD-O1D
29	s	602	CLA	CHA-CBD-CGD-O2D
29	s	604	CLA	CHA-CBD-CGD-O1D
29	s	604	CLA	CHA-CBD-CGD-O2D
29	s	605	CLA	C1A-C2A-CAA-CBA
29	s	605	CLA	CHA-CBD-CGD-O1D
29	s	605	CLA	CHA-CBD-CGD-O2D
29	s	609	CLA	C2-C3-C5-C6
29	s	609	CLA	C4-C3-C5-C6
29	s	610	CLA	CHA-CBD-CGD-O1D
29	s	610	CLA	CHA-CBD-CGD-O2D
29	s	610	CLA	CBD-CGD-O2D-CED
29	s	611	CLA	CHA-CBD-CGD-O1D
29	s	611	CLA	CHA-CBD-CGD-O2D
29	s	613	CLA	CHA-CBD-CGD-O1D
29	s	613	CLA	CHA-CBD-CGD-O2D
29	s	614	CLA	C1A-C2A-CAA-CBA
29	s	617	CLA	CBD-CGD-O2D-CED
29	y	603	CLA	C3A-C2A-CAA-CBA
29	y	604	CLA	CHA-CBD-CGD-O1D
29	y	604	CLA	CHA-CBD-CGD-O2D
29	y	610	CLA	CBD-CGD-O2D-CED

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Mol	Chain	Res	Type	Atoms
29	y	614	CLA	C2-C1-O2A-CGA
30	A	409	PHO	C3A-C2A-CAA-CBA
31	A	411	BCR	C11-C10-C9-C8
31	A	411	BCR	C11-C10-C9-C34
31	A	411	BCR	C10-C11-C12-C13
31	A	411	BCR	C11-C12-C13-C35
31	A	411	BCR	C17-C18-C19-C20
31	A	411	BCR	C36-C18-C19-C20
31	A	411	BCR	C23-C24-C25-C30
31	B	618	BCR	C1-C6-C7-C8
31	B	618	BCR	C7-C8-C9-C10
31	B	618	BCR	C7-C8-C9-C34
31	B	618	BCR	C11-C10-C9-C8
31	B	618	BCR	C11-C10-C9-C34
31	B	618	BCR	C10-C11-C12-C13
31	B	618	BCR	C17-C18-C19-C20
31	B	618	BCR	C36-C18-C19-C20
31	B	618	BCR	C37-C22-C23-C24
31	B	619	BCR	C11-C10-C9-C8
31	B	619	BCR	C11-C10-C9-C34
31	B	619	BCR	C10-C11-C12-C13
31	B	619	BCR	C21-C22-C23-C24
31	B	619	BCR	C37-C22-C23-C24
31	C	514	BCR	C11-C10-C9-C8
31	C	514	BCR	C11-C10-C9-C34
31	C	514	BCR	C9-C10-C11-C12
31	C	514	BCR	C10-C11-C12-C13
31	C	514	BCR	C13-C14-C15-C16
31	C	514	BCR	C17-C18-C19-C20
31	C	514	BCR	C36-C18-C19-C20
31	C	515	BCR	C7-C8-C9-C10
31	C	515	BCR	C7-C8-C9-C34
31	C	515	BCR	C11-C10-C9-C8
31	C	515	BCR	C11-C10-C9-C34
31	C	515	BCR	C17-C18-C19-C20
31	C	515	BCR	C36-C18-C19-C20
31	C	515	BCR	C23-C24-C25-C30
31	C	516	BCR	C1-C6-C7-C8
31	C	516	BCR	C5-C6-C7-C8
31	C	516	BCR	C7-C8-C9-C10
31	C	516	BCR	C7-C8-C9-C34
31	C	516	BCR	C11-C10-C9-C8

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Mol	Chain	Res	Type	Atoms
31	C	516	BCR	C11-C10-C9-C34
31	C	516	BCR	C10-C11-C12-C13
31	C	516	BCR	C17-C18-C19-C20
31	C	516	BCR	C36-C18-C19-C20
31	C	516	BCR	C21-C22-C23-C24
31	C	516	BCR	C37-C22-C23-C24
31	C	517	BCR	C11-C10-C9-C8
31	C	517	BCR	C11-C10-C9-C34
31	C	517	BCR	C10-C11-C12-C13
31	C	517	BCR	C17-C18-C19-C20
31	C	517	BCR	C36-C18-C19-C20
31	D	404	BCR	C11-C10-C9-C8
31	D	404	BCR	C11-C10-C9-C34
31	D	404	BCR	C10-C11-C12-C13
31	a	411	BCR	C11-C10-C9-C8
31	a	411	BCR	C11-C10-C9-C34
31	a	411	BCR	C10-C11-C12-C13
31	a	411	BCR	C17-C18-C19-C20
31	a	411	BCR	C36-C18-C19-C20
31	a	411	BCR	C23-C24-C25-C30
31	b	618	BCR	C7-C8-C9-C10
31	b	618	BCR	C7-C8-C9-C34
31	b	618	BCR	C11-C10-C9-C8
31	b	618	BCR	C11-C10-C9-C34
31	b	618	BCR	C10-C11-C12-C13
31	b	618	BCR	C17-C18-C19-C20
31	b	618	BCR	C36-C18-C19-C20
31	b	619	BCR	C11-C10-C9-C8
31	b	619	BCR	C11-C10-C9-C34
31	b	619	BCR	C10-C11-C12-C13
31	b	619	BCR	C21-C22-C23-C24
31	b	619	BCR	C37-C22-C23-C24
31	c	514	BCR	C11-C10-C9-C8
31	c	514	BCR	C11-C10-C9-C34
31	c	514	BCR	C9-C10-C11-C12
31	c	514	BCR	C10-C11-C12-C13
31	c	514	BCR	C17-C18-C19-C20
31	c	514	BCR	C36-C18-C19-C20
31	c	515	BCR	C7-C8-C9-C34
31	c	515	BCR	C11-C10-C9-C8
31	c	515	BCR	C11-C10-C9-C34
31	c	515	BCR	C10-C11-C12-C13

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Mol	Chain	Res	Type	Atoms
31	c	515	BCR	C17-C18-C19-C20
31	c	515	BCR	C36-C18-C19-C20
31	c	516	BCR	C1-C6-C7-C8
31	c	516	BCR	C5-C6-C7-C8
31	c	516	BCR	C7-C8-C9-C10
31	c	516	BCR	C7-C8-C9-C34
31	c	516	BCR	C11-C10-C9-C8
31	c	516	BCR	C11-C10-C9-C34
31	c	516	BCR	C10-C11-C12-C13
31	c	516	BCR	C21-C22-C23-C24
31	c	516	BCR	C37-C22-C23-C24
31	c	516	BCR	C23-C24-C25-C30
31	c	517	BCR	C11-C10-C9-C8
31	c	517	BCR	C11-C10-C9-C34
31	c	517	BCR	C17-C18-C19-C20
31	c	517	BCR	C36-C18-C19-C20
31	d	404	BCR	C11-C10-C9-C8
31	d	404	BCR	C11-C10-C9-C34
31	d	404	BCR	C10-C11-C12-C13
32	A	412	SQD	C2-C1-O6-C44
32	A	412	SQD	O5-C1-O6-C44
32	A	412	SQD	O5-C5-C6-S
32	B	621	SQD	O5-C1-O6-C44
32	B	621	SQD	C8-C7-O47-C45
32	B	621	SQD	C5-C6-S-O7
32	B	621	SQD	C5-C6-S-O8
32	B	621	SQD	C5-C6-S-O9
32	C	526	SQD	O5-C5-C6-S
32	C	526	SQD	C5-C6-S-O7
32	C	526	SQD	C5-C6-S-O8
32	C	526	SQD	C5-C6-S-O9
32	a	412	SQD	O5-C1-O6-C44
32	a	412	SQD	O5-C5-C6-S
32	b	621	SQD	O5-C1-O6-C44
32	b	621	SQD	C8-C7-O47-C45
32	b	621	SQD	C5-C6-S-O7
32	b	621	SQD	C5-C6-S-O8
32	b	621	SQD	C5-C6-S-O9
32	c	626	SQD	O5-C5-C6-S
32	c	626	SQD	C5-C6-S-O7
32	c	626	SQD	C5-C6-S-O8
32	c	626	SQD	C5-C6-S-O9

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Mol	Chain	Res	Type	Atoms
33	A	413	LMG	O9-C10-O7-C8
33	A	413	LMG	C11-C10-O7-C8
33	B	622	LMG	C9-C8-O7-C10
33	B	622	LMG	C11-C10-O7-C8
33	C	521	LMG	C2-C1-O1-C7
33	C	521	LMG	O6-C1-O1-C7
33	C	521	LMG	C11-C10-O7-C8
33	b	622	LMG	O9-C10-O7-C8
33	c	521	LMG	C2-C1-O1-C7
33	c	521	LMG	O6-C1-O1-C7
33	c	521	LMG	C11-C10-O7-C8
33	j	101	LMG	O7-C8-C9-O8
35	B	620	C7Z	C7-C8-C9-C19
35	B	620	C7Z	C7-C8-C9-C10
35	B	620	C7Z	C11-C12-C13-C20
35	B	620	C7Z	C11-C12-C13-C14
35	B	620	C7Z	C27-C28-C29-C30
35	B	620	C7Z	C27-C28-C29-C39
35	b	620	C7Z	C7-C8-C9-C19
35	b	620	C7Z	C7-C8-C9-C10
35	b	620	C7Z	C11-C12-C13-C20
35	b	620	C7Z	C11-C12-C13-C14
35	b	620	C7Z	C27-C28-C29-C30
35	b	620	C7Z	C27-C28-C29-C39
36	B	625	DGA	OB1-CB1-OG2-CG2
36	B	625	DGA	CG1-CG2-CG3-OXT
36	B	625	DGA	OG2-CG2-CG3-OXT
36	b	623	DGA	OB1-CB1-OG2-CG2
38	C	519	DGD	C2E-C1E-O5D-C6D
38	C	519	DGD	O6E-C1E-O5D-C6D
38	C	520	DGD	C2B-C1B-O2G-C2G
38	C	523	DGD	C2A-C1A-O1G-C1G
38	C	523	DGD	O1A-C1A-O1G-C1G
38	C	523	DGD	O1B-C1B-O2G-C2G
38	C	523	DGD	C2E-C1E-O5D-C6D
38	C	523	DGD	O6E-C1E-O5D-C6D
38	c	519	DGD	C2E-C1E-O5D-C6D
38	c	519	DGD	O6E-C1E-O5D-C6D
38	c	520	DGD	C2B-C1B-O2G-C2G
38	c	523	DGD	C2E-C1E-O5D-C6D
38	c	523	DGD	O6E-C1E-O5D-C6D
39	C	525	LHG	C1-C2-C3-O3

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Mol	Chain	Res	Type	Atoms
39	C	525	LHG	C3-O3-P-O4
39	C	525	LHG	C4-O6-P-O4
39	D	408	LHG	C1-C2-C3-O3
39	D	408	LHG	C4-O6-P-O4
39	D	409	LHG	O2-C2-C3-O3
39	D	409	LHG	C3-O3-P-O5
39	D	409	LHG	C3-O3-P-O6
39	D	409	LHG	C4-O6-P-O5
39	D	410	LHG	C1-C2-C3-O3
39	D	410	LHG	C3-O3-P-O5
39	D	410	LHG	C4-O6-P-O4
39	L	101	LHG	O1-C1-C2-C3
39	L	101	LHG	C1-C2-C3-O3
39	L	101	LHG	C4-O6-P-O3
39	L	101	LHG	C4-O6-P-O5
39	N	624	LHG	O9-C7-O7-C5
39	N	624	LHG	C8-C7-O7-C5
39	G	630	LHG	O1-C1-C2-C3
39	G	630	LHG	C1-C2-C3-O3
39	G	630	LHG	C4-O6-P-O4
39	S	624	LHG	C8-C7-O7-C5
39	Y	624	LHG	C1-C2-C3-O3
39	Y	624	LHG	O2-C2-C3-O3
39	Y	624	LHG	C4-O6-P-O3
39	Y	624	LHG	C4-O6-P-O4
39	Y	624	LHG	C4-O6-P-O5
39	c	625	LHG	C3-O3-P-O6
39	c	625	LHG	C4-O6-P-O3
39	c	625	LHG	C4-O6-P-O4
39	c	625	LHG	C4-O6-P-O5
39	d	408	LHG	O1-C1-C2-C3
39	d	408	LHG	C4-O6-P-O4
39	d	408	LHG	C4-O6-P-O5
39	d	409	LHG	C1-C2-C3-O3
39	d	409	LHG	C4-O6-P-O5
39	d	410	LHG	O1-C1-C2-C3
39	d	410	LHG	C3-O3-P-O4
39	d	410	LHG	C3-O3-P-O5
39	d	410	LHG	C4-O6-P-O5
39	d	410	LHG	O7-C5-C6-O8
39	l	101	LHG	O1-C1-C2-C3
39	l	101	LHG	C1-C2-C3-O3

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Mol	Chain	Res	Type	Atoms
39	l	101	LHG	C4-O6-P-O3
39	l	101	LHG	C4-O6-P-O5
39	n	624	LHG	O9-C7-O7-C5
39	n	624	LHG	C8-C7-O7-C5
39	g	624	LHG	O1-C1-C2-C3
39	g	624	LHG	C1-C2-C3-O3
39	g	624	LHG	C4-O6-P-O4
39	g	624	LHG	O7-C5-C6-O8
39	s	624	LHG	O1-C1-C2-C3
39	s	624	LHG	C8-C7-O7-C5
39	y	624	LHG	C1-C2-C3-O3
39	y	624	LHG	C4-O6-P-O4
39	y	624	LHG	C4-O6-P-O5
40	C	527	LMK	O9-C10-O7-C8
40	C	527	LMK	O10-C28-O8-C9
40	c	627	LMK	O9-C10-C11-C12
40	c	627	LMK	O9-C10-O7-C8
42	D	405	PL9	C12-C13-C14-C15
42	D	405	PL9	C12-C13-C14-C16
42	d	405	PL9	C12-C13-C14-C15
44	H	101	RRX	C37-C22-C23-C24
44	H	101	RRX	C21-C22-C23-C24
44	H	101	RRX	C7-C8-C9-C34
44	h	101	RRX	C37-C22-C23-C24
44	h	101	RRX	C21-C22-C23-C24
44	h	101	RRX	C7-C8-C9-C10
44	h	101	RRX	C7-C8-C9-C34
45	G	601	CHL	CHA-CBD-CGD-O1D
45	G	601	CHL	CHA-CBD-CGD-O2D
45	G	601	CHL	C2-C3-C5-C6
45	G	601	CHL	C4-C3-C5-C6
45	G	606	CHL	C1A-C2A-CAA-CBA
45	G	607	CHL	C1A-C2A-CAA-CBA
45	R	606	CHL	CHA-CBD-CGD-O1D
45	R	606	CHL	CHA-CBD-CGD-O2D
45	R	606	CHL	CAD-CBD-CGD-O1D
45	R	606	CHL	CAD-CBD-CGD-O2D
45	R	607	CHL	CHA-CBD-CGD-O1D
45	R	607	CHL	CHA-CBD-CGD-O2D
45	S	607	CHL	C1A-C2A-CAA-CBA
45	Y	601	CHL	CHA-CBD-CGD-O1D
45	Y	601	CHL	CHA-CBD-CGD-O2D

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Mol	Chain	Res	Type	Atoms
45	n	609	CHL	C1A-C2A-CAA-CBA
45	g	601	CHL	CHA-CBD-CGD-O1D
45	g	601	CHL	CHA-CBD-CGD-O2D
45	g	601	CHL	C2-C3-C5-C6
45	g	601	CHL	C4-C3-C5-C6
45	g	605	CHL	CHA-CBD-CGD-O1D
45	g	606	CHL	C1A-C2A-CAA-CBA
45	s	607	CHL	C1A-C2A-CAA-CBA
45	y	601	CHL	CHA-CBD-CGD-O1D
45	y	601	CHL	CHA-CBD-CGD-O2D
45	y	601	CHL	C2-C3-C5-C6
45	y	601	CHL	C4-C3-C5-C6
46	N	621	LUT	C21-C26-C27-C28
46	G	620	LUT	C25-C26-C27-C28
46	G	621	LUT	C21-C26-C27-C28
46	R	620	LUT	C7-C8-C9-C10
46	S	620	LUT	C21-C26-C27-C28
46	S	620	LUT	C27-C28-C29-C30
46	S	620	LUT	C27-C28-C29-C39
46	S	621	LUT	C25-C26-C27-C28
46	Y	620	LUT	C27-C28-C29-C39
46	Y	621	LUT	C21-C26-C27-C28
46	n	620	LUT	C1-C6-C7-C8
46	n	620	LUT	C25-C26-C27-C28
46	n	621	LUT	C21-C26-C27-C28
46	g	620	LUT	C27-C28-C29-C30
46	g	620	LUT	C27-C28-C29-C39
46	g	621	LUT	C21-C26-C27-C28
46	s	620	LUT	C21-C26-C27-C28
46	s	620	LUT	C25-C26-C27-C28
47	R	621	XAT	C9-C10-C11-C12
47	R	621	XAT	C10-C11-C12-C13
47	R	621	XAT	O24-C26-C27-C28
47	R	621	XAT	C26-C27-C28-C29
47	Y	622	XAT	C27-C28-C29-C30
47	Y	622	XAT	C27-C28-C29-C39
47	r	622	XAT	C7-C8-C9-C10
47	r	622	XAT	C7-C8-C9-C19
47	r	622	XAT	C9-C10-C11-C12
47	r	622	XAT	C10-C11-C12-C13
47	r	622	XAT	C20-C13-C14-C15
47	r	622	XAT	O24-C26-C27-C28

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Mol	Chain	Res	Type	Atoms
47	r	622	XAT	C26-C27-C28-C29
48	R	622	NEX	C12-C13-C14-C15
48	R	622	NEX	C20-C13-C14-C15
48	R	622	NEX	C14-C15-C35-C34
48	S	622	NEX	C11-C10-C9-C8
48	S	622	NEX	C11-C10-C9-C19
48	S	622	NEX	C10-C11-C12-C13
48	S	622	NEX	C11-C12-C13-C14
48	S	622	NEX	C11-C12-C13-C20
48	r	623	NEX	C12-C13-C14-C15
48	r	623	NEX	C20-C13-C14-C15
48	r	623	NEX	C30-C31-C32-C33
48	s	623	NEX	C10-C11-C12-C13
48	s	623	NEX	C11-C12-C13-C14
48	s	623	NEX	C11-C12-C13-C20
49	S	625	LPX	C3-C4-C5-O6
49	S	625	LPX	O1-C3-C4-C5
49	s	625	LPX	O1-C3-C4-O5
50	S	626	3PH	C22-C21-O21-C2
50	i	101	3PH	C1-O11-P-O13
50	i	101	3PH	C1-O11-P-O14
50	s	626	3PH	C1-O11-P-O13
50	s	626	3PH	C1-O11-P-O14
50	s	626	3PH	C1-O11-P-O12
50	s	626	3PH	O21-C2-C3-O31
50	s	626	3PH	C22-C21-O21-C2
51	Y	625	SPH	C1-C2-C3-O3
51	Y	625	SPH	C1-C2-C3-C4
51	Y	625	SPH	N2-C2-C3-O3
51	Y	625	SPH	N2-C2-C3-C4
51	y	625	SPH	O1-C1-C2-N2
51	y	625	SPH	C1-C2-C3-O3
51	y	625	SPH	C1-C2-C3-C4
51	y	625	SPH	N2-C2-C3-O3
51	y	625	SPH	N2-C2-C3-C4
29	S	603	CLA	O1D-CGD-O2D-CED
29	c	502	CLA	O1D-CGD-O2D-CED
29	r	611	CLA	O1D-CGD-O2D-CED
29	N	612	CLA	O1D-CGD-O2D-CED
29	R	603	CLA	O1D-CGD-O2D-CED
29	R	611	CLA	O1D-CGD-O2D-CED
29	b	612	CLA	O1D-CGD-O2D-CED

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Mol	Chain	Res	Type	Atoms
29	g	613	CLA	O1D-CGD-O2D-CED
29	r	603	CLA	O1D-CGD-O2D-CED
29	s	603	CLA	O1D-CGD-O2D-CED
29	y	604	CLA	O1D-CGD-O2D-CED
29	A	410	CLA	CBD-CGD-O2D-CED
29	B	606	CLA	CBD-CGD-O2D-CED
29	B	609	CLA	CBD-CGD-O2D-CED
29	B	611	CLA	CBD-CGD-O2D-CED
29	B	616	CLA	CBD-CGD-O2D-CED
29	C	502	CLA	CBD-CGD-O2D-CED
29	C	504	CLA	CBD-CGD-O2D-CED
29	C	509	CLA	CBD-CGD-O2D-CED
29	C	513	CLA	CBD-CGD-O2D-CED
29	N	602	CLA	CBD-CGD-O2D-CED
29	N	612	CLA	CBD-CGD-O2D-CED
29	N	613	CLA	CBD-CGD-O2D-CED
29	N	614	CLA	CBD-CGD-O2D-CED
29	G	602	CLA	CBD-CGD-O2D-CED
29	G	604	CLA	CBD-CGD-O2D-CED
29	G	611	CLA	CBD-CGD-O2D-CED
29	G	612	CLA	CBD-CGD-O2D-CED
29	G	613	CLA	CBD-CGD-O2D-CED
29	R	602	CLA	CBD-CGD-O2D-CED
29	R	603	CLA	CBD-CGD-O2D-CED
29	R	608	CLA	CBD-CGD-O2D-CED
29	R	611	CLA	CBD-CGD-O2D-CED
29	R	612	CLA	CBD-CGD-O2D-CED
29	R	613	CLA	CBD-CGD-O2D-CED
29	S	604	CLA	CBD-CGD-O2D-CED
29	S	605	CLA	CBD-CGD-O2D-CED
29	S	613	CLA	CBD-CGD-O2D-CED
29	S	614	CLA	CBD-CGD-O2D-CED
29	Y	602	CLA	CBD-CGD-O2D-CED
29	Y	603	CLA	CBD-CGD-O2D-CED
29	Y	608	CLA	CBD-CGD-O2D-CED
29	Y	612	CLA	CBD-CGD-O2D-CED
29	Y	613	CLA	CBD-CGD-O2D-CED
29	a	410	CLA	CBD-CGD-O2D-CED
29	b	603	CLA	CBD-CGD-O2D-CED
29	b	606	CLA	CBD-CGD-O2D-CED
29	b	611	CLA	CBD-CGD-O2D-CED
29	b	612	CLA	CBD-CGD-O2D-CED

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Mol	Chain	Res	Type	Atoms
29	b	614	CLA	CBD-CGD-O2D-CED
29	c	501	CLA	CBD-CGD-O2D-CED
29	c	503	CLA	CBD-CGD-O2D-CED
29	c	504	CLA	CBD-CGD-O2D-CED
29	c	507	CLA	CBD-CGD-O2D-CED
29	c	509	CLA	CBD-CGD-O2D-CED
29	c	510	CLA	CBD-CGD-O2D-CED
29	c	513	CLA	CBD-CGD-O2D-CED
29	d	402	CLA	CBD-CGD-O2D-CED
29	n	602	CLA	CBD-CGD-O2D-CED
29	n	611	CLA	CBD-CGD-O2D-CED
29	g	602	CLA	CBD-CGD-O2D-CED
29	g	603	CLA	CBD-CGD-O2D-CED
29	g	604	CLA	CBD-CGD-O2D-CED
29	g	611	CLA	CBD-CGD-O2D-CED
29	g	612	CLA	CBD-CGD-O2D-CED
29	g	613	CLA	CBD-CGD-O2D-CED
29	r	602	CLA	CBD-CGD-O2D-CED
29	r	603	CLA	CBD-CGD-O2D-CED
29	r	604	CLA	CBD-CGD-O2D-CED
29	r	608	CLA	CBD-CGD-O2D-CED
29	r	611	CLA	CBD-CGD-O2D-CED
29	r	612	CLA	CBD-CGD-O2D-CED
29	s	603	CLA	CBD-CGD-O2D-CED
29	s	604	CLA	CBD-CGD-O2D-CED
29	s	609	CLA	CBD-CGD-O2D-CED
29	s	612	CLA	CBD-CGD-O2D-CED
29	s	613	CLA	CBD-CGD-O2D-CED
29	s	614	CLA	CBD-CGD-O2D-CED
29	y	603	CLA	CBD-CGD-O2D-CED
29	y	604	CLA	CBD-CGD-O2D-CED
29	y	608	CLA	CBD-CGD-O2D-CED
29	y	612	CLA	CBD-CGD-O2D-CED
29	y	613	CLA	CBD-CGD-O2D-CED
29	y	614	CLA	CBD-CGD-O2D-CED
30	A	409	PHO	CBD-CGD-O2D-CED
29	A	406	CLA	O1A-CGA-O2A-C1
29	C	504	CLA	O1A-CGA-O2A-C1
29	G	614	CLA	O1A-CGA-O2A-C1
29	S	605	CLA	O1A-CGA-O2A-C1
29	S	613	CLA	O1A-CGA-O2A-C1
29	S	617	CLA	O1A-CGA-O2A-C1

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Mol	Chain	Res	Type	Atoms
29	a	406	CLA	O1A-CGA-O2A-C1
29	b	603	CLA	O1A-CGA-O2A-C1
29	b	617	CLA	O1A-CGA-O2A-C1
29	c	504	CLA	O1A-CGA-O2A-C1
29	s	603	CLA	O1A-CGA-O2A-C1
29	s	605	CLA	O1A-CGA-O2A-C1
29	s	614	CLA	O1A-CGA-O2A-C1
29	y	603	CLA	O1A-CGA-O2A-C1
33	j	101	LMG	O10-C28-O8-C9
38	c	523	DGD	O1A-C1A-O1G-C1G
29	R	613	CLA	O1A-CGA-O2A-C1
29	r	611	CLA	O1A-CGA-O2A-C1
29	r	613	CLA	O1A-CGA-O2A-C1
29	A	405	CLA	O1D-CGD-O2D-CED
29	B	606	CLA	O1D-CGD-O2D-CED
29	D	403	CLA	O1D-CGD-O2D-CED
29	N	610	CLA	O1D-CGD-O2D-CED
29	N	613	CLA	O1D-CGD-O2D-CED
29	G	613	CLA	O1D-CGD-O2D-CED
29	R	608	CLA	O1D-CGD-O2D-CED
29	Y	608	CLA	O1D-CGD-O2D-CED
29	a	405	CLA	O1D-CGD-O2D-CED
29	b	606	CLA	O1D-CGD-O2D-CED
29	b	617	CLA	O1D-CGD-O2D-CED
29	d	403	CLA	O1D-CGD-O2D-CED
29	r	608	CLA	O1D-CGD-O2D-CED
29	s	610	CLA	O1D-CGD-O2D-CED
29	y	608	CLA	O1D-CGD-O2D-CED
29	y	613	CLA	O1D-CGD-O2D-CED
29	r	611	CLA	CBA-CGA-O2A-C1
29	B	602	CLA	O1D-CGD-O2D-CED
29	B	604	CLA	O1D-CGD-O2D-CED
29	B	607	CLA	O1D-CGD-O2D-CED
29	B	608	CLA	O1D-CGD-O2D-CED
29	B	612	CLA	O1D-CGD-O2D-CED
29	B	614	CLA	O1D-CGD-O2D-CED
29	C	502	CLA	O1D-CGD-O2D-CED
29	C	503	CLA	O1D-CGD-O2D-CED
29	C	507	CLA	O1D-CGD-O2D-CED
29	C	511	CLA	O1D-CGD-O2D-CED
29	G	614	CLA	O1D-CGD-O2D-CED
29	R	609	CLA	O1D-CGD-O2D-CED

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Mol	Chain	Res	Type	Atoms
29	S	609	CLA	O1D-CGD-O2D-CED
29	S	612	CLA	O1D-CGD-O2D-CED
29	S	614	CLA	O1D-CGD-O2D-CED
29	Y	613	CLA	O1D-CGD-O2D-CED
29	b	604	CLA	O1D-CGD-O2D-CED
29	b	607	CLA	O1D-CGD-O2D-CED
29	c	511	CLA	O1D-CGD-O2D-CED
29	n	612	CLA	O1D-CGD-O2D-CED
29	n	613	CLA	O1D-CGD-O2D-CED
29	n	614	CLA	O1D-CGD-O2D-CED
29	y	610	CLA	O1D-CGD-O2D-CED
29	A	406	CLA	CBA-CGA-O2A-C1
29	C	504	CLA	CBA-CGA-O2A-C1
29	C	505	CLA	CBA-CGA-O2A-C1
29	G	614	CLA	CBA-CGA-O2A-C1
29	S	605	CLA	CBA-CGA-O2A-C1
29	S	617	CLA	CBA-CGA-O2A-C1
29	a	406	CLA	CBA-CGA-O2A-C1
29	b	617	CLA	CBA-CGA-O2A-C1
29	c	504	CLA	CBA-CGA-O2A-C1
29	s	603	CLA	CBA-CGA-O2A-C1
29	s	605	CLA	CBA-CGA-O2A-C1
29	s	614	CLA	CBA-CGA-O2A-C1
29	y	603	CLA	CBA-CGA-O2A-C1
38	c	523	DGD	C2A-C1A-O1G-C1G
29	B	613	CLA	CBD-CGD-O2D-CED
29	C	501	CLA	CBD-CGD-O2D-CED
29	C	510	CLA	CBD-CGD-O2D-CED
29	D	402	CLA	CBD-CGD-O2D-CED
29	N	604	CLA	CBD-CGD-O2D-CED
29	N	611	CLA	CBD-CGD-O2D-CED
29	R	604	CLA	CBD-CGD-O2D-CED
29	R	610	CLA	CBD-CGD-O2D-CED
29	S	611	CLA	CBD-CGD-O2D-CED
29	Y	611	CLA	CBD-CGD-O2D-CED
29	b	613	CLA	CBD-CGD-O2D-CED
29	b	616	CLA	CBD-CGD-O2D-CED
29	n	603	CLA	CBD-CGD-O2D-CED
29	n	604	CLA	CBD-CGD-O2D-CED
29	s	605	CLA	CBD-CGD-O2D-CED
29	s	611	CLA	CBD-CGD-O2D-CED
29	y	602	CLA	CBD-CGD-O2D-CED

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Mol	Chain	Res	Type	Atoms
29	B	603	CLA	O1A-CGA-O2A-C1
29	C	505	CLA	O1A-CGA-O2A-C1
29	C	511	CLA	O1A-CGA-O2A-C1
29	N	611	CLA	O1A-CGA-O2A-C1
29	Y	603	CLA	O1A-CGA-O2A-C1
29	Y	604	CLA	O1A-CGA-O2A-C1
29	Y	611	CLA	O1A-CGA-O2A-C1
29	c	505	CLA	O1A-CGA-O2A-C1
29	c	511	CLA	O1A-CGA-O2A-C1
29	g	602	CLA	O1A-CGA-O2A-C1
29	g	614	CLA	O1A-CGA-O2A-C1
29	s	602	CLA	O1A-CGA-O2A-C1
29	s	613	CLA	O1A-CGA-O2A-C1
29	y	604	CLA	O1A-CGA-O2A-C1
29	y	611	CLA	O1A-CGA-O2A-C1
33	J	101	LMG	O10-C28-O8-C9
50	S	626	3PH	O32-C31-O31-C3
50	s	626	3PH	O32-C31-O31-C3
29	B	610	CLA	O1D-CGD-O2D-CED
29	G	610	CLA	O1D-CGD-O2D-CED
29	S	610	CLA	O1D-CGD-O2D-CED
29	S	617	CLA	O1D-CGD-O2D-CED
29	Y	604	CLA	O1D-CGD-O2D-CED
29	Y	610	CLA	O1D-CGD-O2D-CED
29	b	602	CLA	O1D-CGD-O2D-CED
29	b	608	CLA	O1D-CGD-O2D-CED
29	b	609	CLA	O1D-CGD-O2D-CED
29	n	610	CLA	O1D-CGD-O2D-CED
29	g	614	CLA	O1D-CGD-O2D-CED
29	s	617	CLA	O1D-CGD-O2D-CED
29	B	603	CLA	O1D-CGD-O2D-CED
29	B	617	CLA	O1D-CGD-O2D-CED
29	G	603	CLA	O1D-CGD-O2D-CED
29	Y	614	CLA	O1D-CGD-O2D-CED
29	b	610	CLA	O1D-CGD-O2D-CED
29	g	610	CLA	O1D-CGD-O2D-CED
29	r	609	CLA	O1D-CGD-O2D-CED
29	r	613	CLA	O1D-CGD-O2D-CED
29	C	506	CLA	CBD-CGD-O2D-CED
29	N	603	CLA	CBD-CGD-O2D-CED
29	c	506	CLA	CBD-CGD-O2D-CED
29	N	614	CLA	O1D-CGD-O2D-CED

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Mol	Chain	Res	Type	Atoms
29	R	602	CLA	O1D-CGD-O2D-CED
29	b	603	CLA	O1D-CGD-O2D-CED
29	g	603	CLA	O1D-CGD-O2D-CED
29	s	609	CLA	O1D-CGD-O2D-CED
32	B	621	SQD	O49-C7-O47-C45
32	b	621	SQD	O49-C7-O47-C45
33	B	622	LMG	O9-C10-O7-C8
33	C	521	LMG	O9-C10-O7-C8
33	c	521	LMG	O9-C10-O7-C8
38	C	520	DGD	O1B-C1B-O2G-C2G
38	c	520	DGD	O1B-C1B-O2G-C2G
38	c	523	DGD	O1B-C1B-O2G-C2G
39	s	624	LHG	O9-C7-O7-C5
50	S	626	3PH	O22-C21-O21-C2
50	s	626	3PH	O22-C21-O21-C2
29	A	410	CLA	C3-C5-C6-C7
29	B	605	CLA	C3-C5-C6-C7
29	B	612	CLA	C3-C5-C6-C7
29	C	504	CLA	C3-C5-C6-C7
29	C	511	CLA	C3-C5-C6-C7
29	C	512	CLA	C3-C5-C6-C7
29	D	403	CLA	C3-C5-C6-C7
29	N	603	CLA	C3-C5-C6-C7
29	N	610	CLA	C3-C5-C6-C7
29	G	603	CLA	C3-C5-C6-C7
29	S	609	CLA	C3-C5-C6-C7
29	Y	613	CLA	C3-C5-C6-C7
29	b	602	CLA	C3-C5-C6-C7
29	b	605	CLA	C3-C5-C6-C7
29	b	608	CLA	C3-C5-C6-C7
29	b	610	CLA	C3-C5-C6-C7
29	b	612	CLA	C3-C5-C6-C7
29	c	504	CLA	C3-C5-C6-C7
29	c	512	CLA	C3-C5-C6-C7
29	n	602	CLA	C3-C5-C6-C7
29	r	609	CLA	C3-C5-C6-C7
29	s	609	CLA	C3-C5-C6-C7
29	y	611	CLA	C3-C5-C6-C7
30	a	409	PHO	C3-C5-C6-C7
29	B	617	CLA	CBA-CGA-O2A-C1
29	C	511	CLA	CBA-CGA-O2A-C1
29	N	603	CLA	CBA-CGA-O2A-C1

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Mol	Chain	Res	Type	Atoms
29	S	613	CLA	CBA-CGA-O2A-C1
29	Y	604	CLA	CBA-CGA-O2A-C1
29	Y	611	CLA	CBA-CGA-O2A-C1
29	Y	612	CLA	CBA-CGA-O2A-C1
29	b	603	CLA	CBA-CGA-O2A-C1
29	c	505	CLA	CBA-CGA-O2A-C1
29	c	511	CLA	CBA-CGA-O2A-C1
29	r	612	CLA	CBA-CGA-O2A-C1
29	s	613	CLA	CBA-CGA-O2A-C1
29	y	604	CLA	CBA-CGA-O2A-C1
29	y	611	CLA	CBA-CGA-O2A-C1
33	J	101	LMG	C29-C28-O8-C9
33	j	101	LMG	C29-C28-O8-C9
50	S	626	3PH	C32-C31-O31-C3
50	s	626	3PH	C32-C31-O31-C3
33	b	622	LMG	C11-C10-O7-C8
36	B	625	DGA	CB2-CB1-OG2-CG2
36	b	623	DGA	CB2-CB1-OG2-CG2
38	C	523	DGD	C2B-C1B-O2G-C2G
38	c	523	DGD	C2B-C1B-O2G-C2G
29	G	604	CLA	O1D-CGD-O2D-CED
29	r	602	CLA	O1D-CGD-O2D-CED
29	y	611	CLA	CBD-CGD-O2D-CED
29	s	617	CLA	O1A-CGA-O2A-C1
50	i	101	3PH	O32-C31-O31-C3
29	R	602	CLA	C4-C3-C5-C6
29	R	602	CLA	C2-C3-C5-C6
29	c	503	CLA	C2-C3-C5-C6
29	r	610	CLA	CBD-CGD-O2D-CED
29	N	612	CLA	C2A-CAA-CBA-CGA
29	N	614	CLA	C2A-CAA-CBA-CGA
29	G	604	CLA	C2A-CAA-CBA-CGA
29	G	613	CLA	C2A-CAA-CBA-CGA
29	R	609	CLA	C2A-CAA-CBA-CGA
29	S	617	CLA	C2A-CAA-CBA-CGA
29	b	612	CLA	C2A-CAA-CBA-CGA
29	n	603	CLA	C2A-CAA-CBA-CGA
29	n	611	CLA	C2A-CAA-CBA-CGA
29	g	613	CLA	C2A-CAA-CBA-CGA
29	r	602	CLA	C2A-CAA-CBA-CGA
29	r	609	CLA	C2A-CAA-CBA-CGA
29	s	617	CLA	C2A-CAA-CBA-CGA

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Mol	Chain	Res	Type	Atoms
45	G	605	CHL	C2A-CAA-CBA-CGA
45	G	606	CHL	C2A-CAA-CBA-CGA
45	Y	605	CHL	C2A-CAA-CBA-CGA
45	n	605	CHL	C2A-CAA-CBA-CGA
45	y	605	CHL	C2A-CAA-CBA-CGA
29	C	512	CLA	O1A-CGA-O2A-C1
29	S	603	CLA	O1A-CGA-O2A-C1
29	c	512	CLA	O1A-CGA-O2A-C1
29	g	613	CLA	O1A-CGA-O2A-C1
33	A	413	LMG	C17-C18-C19-C20
33	A	413	LMG	C35-C36-C37-C38
33	A	413	LMG	C38-C39-C40-C41
33	B	622	LMG	C17-C18-C19-C20
33	C	521	LMG	C17-C18-C19-C20
33	C	521	LMG	C35-C36-C37-C38
33	D	411	LMG	C17-C18-C19-C20
33	D	411	LMG	C20-C21-C22-C23
33	H	102	LMG	C38-C39-C40-C41
33	J	101	LMG	C17-C18-C19-C20
33	J	101	LMG	C20-C21-C22-C23
33	a	413	LMG	C17-C18-C19-C20
33	a	413	LMG	C35-C36-C37-C38
33	a	413	LMG	C38-C39-C40-C41
33	b	622	LMG	C17-C18-C19-C20
33	c	521	LMG	C17-C18-C19-C20
33	c	521	LMG	C35-C36-C37-C38
33	d	411	LMG	C17-C18-C19-C20
33	d	411	LMG	C20-C21-C22-C23
33	h	102	LMG	C38-C39-C40-C41
33	j	101	LMG	C17-C18-C19-C20
33	j	101	LMG	C20-C21-C22-C23
38	C	518	DGD	C8B-C9B-CAB-CBB
38	C	519	DGD	C8A-C9A-CAA-CBA
38	C	519	DGD	C8B-C9B-CAB-CBB
38	C	520	DGD	CBB-CCB-CDB-CEB
38	C	523	DGD	CBA-CCA-CDA-CEA
38	C	523	DGD	CEB-CFB-CGB-CHB
38	c	518	DGD	C8B-C9B-CAB-CBB
38	c	519	DGD	C8A-C9A-CAA-CBA
38	c	519	DGD	C8B-C9B-CAB-CBB
38	c	520	DGD	CBB-CCB-CDB-CEB
38	c	523	DGD	CBA-CCA-CDA-CEA

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Mol	Chain	Res	Type	Atoms
38	c	523	DGD	CEB-CFB-CGB-CHB
29	B	608	CLA	C3-C5-C6-C7
29	N	602	CLA	C3-C5-C6-C7
29	Y	610	CLA	C3-C5-C6-C7
29	c	502	CLA	C3-C5-C6-C7
29	d	403	CLA	C3-C5-C6-C7
29	y	610	CLA	C3-C5-C6-C7
30	A	409	PHO	C3-C5-C6-C7
29	B	603	CLA	CBA-CGA-O2A-C1
29	B	605	CLA	CBA-CGA-O2A-C1
29	D	403	CLA	CBA-CGA-O2A-C1
29	N	611	CLA	CBA-CGA-O2A-C1
29	R	603	CLA	CBA-CGA-O2A-C1
29	R	608	CLA	CBA-CGA-O2A-C1
29	R	612	CLA	CBA-CGA-O2A-C1
29	Y	603	CLA	CBA-CGA-O2A-C1
29	a	405	CLA	CBA-CGA-O2A-C1
29	b	605	CLA	CBA-CGA-O2A-C1
29	g	602	CLA	CBA-CGA-O2A-C1
29	g	614	CLA	CBA-CGA-O2A-C1
29	r	603	CLA	CBA-CGA-O2A-C1
29	r	608	CLA	CBA-CGA-O2A-C1
29	r	609	CLA	CBA-CGA-O2A-C1
29	s	602	CLA	CBA-CGA-O2A-C1
29	s	609	CLA	CBA-CGA-O2A-C1
29	s	617	CLA	CBA-CGA-O2A-C1
29	B	611	CLA	O1D-CGD-O2D-CED
29	C	504	CLA	O1D-CGD-O2D-CED
29	N	602	CLA	O1D-CGD-O2D-CED
29	S	604	CLA	O1D-CGD-O2D-CED
29	b	614	CLA	O1D-CGD-O2D-CED
29	c	503	CLA	O1D-CGD-O2D-CED
29	c	513	CLA	O1D-CGD-O2D-CED
29	r	612	CLA	O1D-CGD-O2D-CED
29	s	604	CLA	O1D-CGD-O2D-CED
29	C	509	CLA	O1D-CGD-O2D-CED
29	G	612	CLA	O1D-CGD-O2D-CED
29	R	612	CLA	O1D-CGD-O2D-CED
29	b	611	CLA	O1D-CGD-O2D-CED
29	c	507	CLA	O1D-CGD-O2D-CED
29	c	509	CLA	O1D-CGD-O2D-CED
29	n	602	CLA	O1D-CGD-O2D-CED

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Mol	Chain	Res	Type	Atoms
29	s	614	CLA	O1D-CGD-O2D-CED
29	y	612	CLA	O1D-CGD-O2D-CED
39	S	624	LHG	O9-C7-O7-C5
29	D	403	CLA	O1A-CGA-O2A-C1
29	N	603	CLA	O1A-CGA-O2A-C1
29	G	602	CLA	O1A-CGA-O2A-C1
29	R	612	CLA	O1A-CGA-O2A-C1
29	S	609	CLA	O1A-CGA-O2A-C1
29	Y	612	CLA	O1A-CGA-O2A-C1
29	a	405	CLA	O1A-CGA-O2A-C1
29	c	502	CLA	O1A-CGA-O2A-C1
29	d	403	CLA	O1A-CGA-O2A-C1
29	r	612	CLA	O1A-CGA-O2A-C1
29	s	609	CLA	O1A-CGA-O2A-C1
29	S	613	CLA	O1D-CGD-O2D-CED
29	c	504	CLA	O1D-CGD-O2D-CED
29	y	614	CLA	O1D-CGD-O2D-CED
47	R	621	XAT	C13-C14-C15-C35
47	r	622	XAT	C13-C14-C15-C35
30	a	409	PHO	CBD-CGD-O2D-CED
29	B	609	CLA	O1D-CGD-O2D-CED
29	B	616	CLA	O1D-CGD-O2D-CED
29	g	612	CLA	O1D-CGD-O2D-CED
39	C	525	LHG	O2-C2-C3-O3
39	D	410	LHG	O2-C2-C3-O3
39	G	630	LHG	O2-C2-C3-O3
39	c	625	LHG	O2-C2-C3-O3
39	d	409	LHG	O2-C2-C3-O3
39	g	624	LHG	O2-C2-C3-O3
39	y	624	LHG	O2-C2-C3-O3
49	S	625	LPX	O1-C3-C4-O5
29	B	610	CLA	C3-C5-C6-C7
29	C	502	CLA	C3-C5-C6-C7
29	c	508	CLA	C3-C5-C6-C7
29	c	511	CLA	C3-C5-C6-C7
29	y	613	CLA	C3-C5-C6-C7
29	B	608	CLA	CBA-CGA-O2A-C1
29	B	613	CLA	CBA-CGA-O2A-C1
29	C	506	CLA	CBA-CGA-O2A-C1
29	C	512	CLA	CBA-CGA-O2A-C1
29	G	602	CLA	CBA-CGA-O2A-C1
29	G	603	CLA	CBA-CGA-O2A-C1

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Mol	Chain	Res	Type	Atoms
29	G	613	CLA	CBA-CGA-O2A-C1
29	S	602	CLA	CBA-CGA-O2A-C1
29	S	609	CLA	CBA-CGA-O2A-C1
29	Y	613	CLA	CBA-CGA-O2A-C1
29	b	608	CLA	CBA-CGA-O2A-C1
29	c	502	CLA	CBA-CGA-O2A-C1
29	g	604	CLA	CBA-CGA-O2A-C1
29	y	602	CLA	CBA-CGA-O2A-C1
29	y	612	CLA	CBA-CGA-O2A-C1
38	c	519	DGD	C2A-C1A-O1G-C1G
50	i	101	3PH	C32-C31-O31-C3
29	B	605	CLA	O1A-CGA-O2A-C1
29	G	613	CLA	O1A-CGA-O2A-C1
29	R	603	CLA	O1A-CGA-O2A-C1
29	R	608	CLA	O1A-CGA-O2A-C1
29	S	602	CLA	O1A-CGA-O2A-C1
29	b	605	CLA	O1A-CGA-O2A-C1
29	r	603	CLA	O1A-CGA-O2A-C1
29	r	608	CLA	O1A-CGA-O2A-C1
29	A	410	CLA	O1D-CGD-O2D-CED
29	G	611	CLA	O1D-CGD-O2D-CED
29	Y	612	CLA	O1D-CGD-O2D-CED
29	d	402	CLA	O1D-CGD-O2D-CED
29	g	602	CLA	O1D-CGD-O2D-CED
29	g	604	CLA	O1D-CGD-O2D-CED
29	s	612	CLA	O1D-CGD-O2D-CED
29	s	613	CLA	O1D-CGD-O2D-CED
29	y	603	CLA	O1D-CGD-O2D-CED
30	A	409	PHO	O1D-CGD-O2D-CED
49	S	625	LPX	O5-C4-C5-O6
38	c	519	DGD	C2B-C1B-O2G-C2G
29	Y	602	CLA	O1D-CGD-O2D-CED
48	r	623	NEX	C14-C15-C35-C34
29	C	513	CLA	O1D-CGD-O2D-CED
29	Y	603	CLA	O1D-CGD-O2D-CED
39	l	101	LHG	C11-C10-C9-C8
29	S	603	CLA	CBA-CGA-O2A-C1
29	c	512	CLA	CBA-CGA-O2A-C1
29	d	403	CLA	CBA-CGA-O2A-C1
29	g	613	CLA	CBA-CGA-O2A-C1
29	G	603	CLA	O1A-CGA-O2A-C1
29	b	608	CLA	O1A-CGA-O2A-C1

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Mol	Chain	Res	Type	Atoms
29	r	609	CLA	O1A-CGA-O2A-C1
39	L	101	LHG	C11-C10-C9-C8
29	b	612	CLA	C4-C3-C5-C6
29	c	501	CLA	C4-C3-C5-C6
45	Y	601	CHL	C4-C3-C5-C6
45	n	601	CHL	C4-C3-C5-C6
29	b	604	CLA	C2-C3-C5-C6
29	b	612	CLA	C2-C3-C5-C6
45	Y	601	CHL	C2-C3-C5-C6
45	n	601	CHL	C2-C3-C5-C6
29	R	602	CLA	C2A-CAA-CBA-CGA
29	g	604	CLA	C2A-CAA-CBA-CGA
29	r	603	CLA	C2A-CAA-CBA-CGA
45	g	605	CHL	C2A-CAA-CBA-CGA
29	R	613	CLA	O1D-CGD-O2D-CED
29	a	410	CLA	O1D-CGD-O2D-CED
29	B	613	CLA	O1A-CGA-O2A-C1
29	C	506	CLA	O1A-CGA-O2A-C1
29	C	510	CLA	O1A-CGA-O2A-C1
29	R	609	CLA	O1A-CGA-O2A-C1
29	g	604	CLA	O1A-CGA-O2A-C1
29	y	602	CLA	O1A-CGA-O2A-C1
29	y	612	CLA	O1A-CGA-O2A-C1
42	d	405	PL9	C34-C36-C37-C38
29	A	405	CLA	CBA-CGA-O2A-C1
29	B	606	CLA	CBA-CGA-O2A-C1
29	C	510	CLA	CBA-CGA-O2A-C1
29	R	604	CLA	CBA-CGA-O2A-C1
29	R	609	CLA	CBA-CGA-O2A-C1
29	Y	602	CLA	CBA-CGA-O2A-C1
29	b	614	CLA	CBA-CGA-O2A-C1
29	c	506	CLA	CBA-CGA-O2A-C1
29	r	604	CLA	CBA-CGA-O2A-C1
39	d	410	LHG	C25-C26-C27-C28
29	n	611	CLA	O1D-CGD-O2D-CED
29	r	604	CLA	O1D-CGD-O2D-CED
29	D	402	CLA	O1D-CGD-O2D-CED
29	G	602	CLA	O1D-CGD-O2D-CED
29	R	610	CLA	O1D-CGD-O2D-CED
29	S	605	CLA	O1D-CGD-O2D-CED
29	b	616	CLA	O1D-CGD-O2D-CED
29	c	501	CLA	O1D-CGD-O2D-CED

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Mol	Chain	Res	Type	Atoms
29	c	510	CLA	O1D-CGD-O2D-CED
29	g	611	CLA	O1D-CGD-O2D-CED
29	B	608	CLA	O1A-CGA-O2A-C1
29	Y	613	CLA	O1A-CGA-O2A-C1
38	c	519	DGD	O1A-C1A-O1G-C1G
29	R	604	CLA	O1D-CGD-O2D-CED
29	S	611	CLA	O1D-CGD-O2D-CED
29	s	605	CLA	O1D-CGD-O2D-CED
29	C	505	CLA	CBD-CGD-O2D-CED
29	c	505	CLA	CBD-CGD-O2D-CED
39	D	409	LHG	C1-C2-C3-O3
38	c	519	DGD	O1B-C1B-O2G-C2G
29	B	606	CLA	O1A-CGA-O2A-C1
29	Y	602	CLA	O1A-CGA-O2A-C1
29	b	606	CLA	O1A-CGA-O2A-C1
29	b	611	CLA	O1A-CGA-O2A-C1
29	c	506	CLA	O1A-CGA-O2A-C1
29	S	604	CLA	C3-C5-C6-C7
29	n	610	CLA	C3-C5-C6-C7
29	N	611	CLA	O1D-CGD-O2D-CED
29	A	410	CLA	CBA-CGA-O2A-C1
29	B	611	CLA	CBA-CGA-O2A-C1
29	B	614	CLA	CBA-CGA-O2A-C1
29	C	509	CLA	CBA-CGA-O2A-C1
29	N	614	CLA	CBA-CGA-O2A-C1
29	b	602	CLA	CBA-CGA-O2A-C1
29	b	606	CLA	CBA-CGA-O2A-C1
29	b	611	CLA	CBA-CGA-O2A-C1
29	b	613	CLA	CBA-CGA-O2A-C1
29	c	503	CLA	CBA-CGA-O2A-C1
29	c	509	CLA	CBA-CGA-O2A-C1
29	c	510	CLA	CBA-CGA-O2A-C1
29	g	603	CLA	CBA-CGA-O2A-C1
29	s	611	CLA	CBA-CGA-O2A-C1
29	y	613	CLA	CBA-CGA-O2A-C1
38	C	519	DGD	C2A-C1A-O1G-C1G
39	c	625	LHG	C24-C23-O8-C6
31	B	618	BCR	C9-C10-C11-C12
35	b	620	C7Z	C9-C10-C11-C12
29	C	502	CLA	C5-C6-C7-C8
29	b	612	CLA	C13-C15-C16-C17
29	c	509	CLA	C10-C11-C12-C13

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Mol	Chain	Res	Type	Atoms
29	g	603	CLA	O1A-CGA-O2A-C1
49	s	625	LPX	C3-C4-C5-O6
29	C	510	CLA	O1D-CGD-O2D-CED
29	y	602	CLA	O1D-CGD-O2D-CED
39	S	624	LHG	O6-C4-C5-O7
29	A	410	CLA	C10-C11-C12-C13
29	N	602	CLA	C5-C6-C7-C8
29	G	602	CLA	C13-C15-C16-C17
29	c	502	CLA	C8-C10-C11-C12
29	c	506	CLA	C8-C10-C11-C12
29	r	612	CLA	C8-C10-C11-C12
39	S	624	LHG	O2-C2-C3-O3
32	b	621	SQD	C23-C24-C25-C26
36	c	524	DGA	CB1-CB2-CB3-CB4
38	c	519	DGD	C1B-C2B-C3B-C4B
39	C	525	LHG	C7-C8-C9-C10
32	C	526	SQD	C2-C1-O6-C44
32	a	412	SQD	C2-C1-O6-C44
32	c	626	SQD	C2-C1-O6-C44
38	C	519	DGD	C2D-C1D-O3G-C3G
38	c	519	DGD	C2D-C1D-O3G-C3G
39	G	630	LHG	O7-C5-C6-O8
29	B	611	CLA	O1A-CGA-O2A-C1
29	b	613	CLA	O1A-CGA-O2A-C1
29	y	613	CLA	O1A-CGA-O2A-C1
29	g	602	CLA	C4-C3-C5-C6
29	B	603	CLA	C14-C13-C15-C16
29	C	506	CLA	C11-C12-C13-C14
29	C	507	CLA	C11-C10-C8-C9
29	C	507	CLA	C14-C13-C15-C16
29	C	513	CLA	C11-C10-C8-C9
29	G	602	CLA	C14-C13-C15-C16
29	R	603	CLA	C11-C10-C8-C9
29	b	603	CLA	C14-C13-C15-C16
29	c	502	CLA	C6-C7-C8-C9
29	c	502	CLA	C11-C12-C13-C14
29	c	506	CLA	C11-C12-C13-C14
29	c	507	CLA	C11-C10-C8-C9
29	c	507	CLA	C14-C13-C15-C16
29	c	513	CLA	C11-C10-C8-C9
29	g	602	CLA	C14-C13-C15-C16
29	r	603	CLA	C11-C10-C8-C9

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Mol	Chain	Res	Type	Atoms
30	A	409	PHO	C6-C7-C8-C9
30	a	409	PHO	C6-C7-C8-C9
45	Y	601	CHL	C11-C10-C8-C9
45	Y	607	CHL	C11-C10-C8-C9
45	n	605	CHL	C14-C13-C15-C16
45	n	607	CHL	C14-C13-C15-C16
45	y	609	CHL	C11-C12-C13-C14
29	C	501	CLA	O1D-CGD-O2D-CED
29	b	613	CLA	O1D-CGD-O2D-CED
29	C	507	CLA	C5-C6-C7-C8
29	C	507	CLA	C15-C16-C17-C18
29	c	504	CLA	C15-C16-C17-C18
29	g	613	CLA	C10-C11-C12-C13
29	c	505	CLA	C2A-CAA-CBA-CGA
45	r	607	CHL	C2A-CAA-CBA-CGA
44	H	101	RRX	C36-C18-C19-C20
44	h	101	RRX	C36-C18-C19-C20
46	R	620	LUT	C7-C8-C9-C19
46	r	620	LUT	C7-C8-C9-C19
46	r	620	LUT	C31-C32-C33-C40
44	H	101	RRX	C17-C18-C19-C20
44	h	101	RRX	C17-C18-C19-C20
46	r	620	LUT	C7-C8-C9-C10
49	s	625	LPX	O5-C4-C5-O6
38	C	519	DGD	C2B-C1B-O2G-C2G
32	B	621	SQD	C23-C24-C25-C26
36	C	524	DGA	CB1-CB2-CB3-CB4
38	c	518	DGD	C1A-C2A-C3A-C4A
39	D	408	LHG	C7-C8-C9-C10
39	d	408	LHG	C7-C8-C9-C10
42	D	405	PL9	C47-C48-C49-C51
29	B	614	CLA	O1A-CGA-O2A-C1
29	b	602	CLA	O1A-CGA-O2A-C1
29	c	510	CLA	O1A-CGA-O2A-C1
38	C	519	DGD	O1A-C1A-O1G-C1G
29	C	501	CLA	C8-C10-C11-C12
29	Y	611	CLA	C10-C11-C12-C13
29	a	410	CLA	C8-C10-C11-C12
29	b	610	CLA	C15-C16-C17-C18
29	n	603	CLA	C10-C11-C12-C13
29	g	610	CLA	C8-C10-C11-C12
29	r	608	CLA	C8-C10-C11-C12

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Mol	Chain	Res	Type	Atoms
29	Y	611	CLA	O1D-CGD-O2D-CED
29	n	603	CLA	O1D-CGD-O2D-CED
29	C	508	CLA	C3-C5-C6-C7
29	R	602	CLA	CBA-CGA-O2A-C1
29	Y	614	CLA	CBA-CGA-O2A-C1
29	a	410	CLA	CBA-CGA-O2A-C1
29	y	614	CLA	CBA-CGA-O2A-C1
29	B	602	CLA	C5-C6-C7-C8
29	B	611	CLA	C15-C16-C17-C18
29	B	612	CLA	C15-C16-C17-C18
29	C	502	CLA	C8-C10-C11-C12
29	C	502	CLA	C13-C15-C16-C17
29	C	503	CLA	C10-C11-C12-C13
29	C	503	CLA	C15-C16-C17-C18
29	C	508	CLA	C10-C11-C12-C13
29	N	603	CLA	C5-C6-C7-C8
29	N	603	CLA	C13-C15-C16-C17
29	N	604	CLA	C10-C11-C12-C13
29	Y	602	CLA	C15-C16-C17-C18
29	Y	610	CLA	C5-C6-C7-C8
29	b	606	CLA	C15-C16-C17-C18
29	b	617	CLA	C5-C6-C7-C8
29	c	507	CLA	C5-C6-C7-C8
29	c	512	CLA	C13-C15-C16-C17
29	g	602	CLA	C13-C15-C16-C17
29	y	603	CLA	C13-C15-C16-C17
29	y	604	CLA	C15-C16-C17-C18
29	y	610	CLA	C13-C15-C16-C17
29	y	613	CLA	C15-C16-C17-C18
39	Y	624	LHG	C33-C34-C35-C36
39	D	408	LHG	C23-C24-C25-C26
39	N	624	LHG	C7-C8-C9-C10
39	d	408	LHG	C23-C24-C25-C26
29	n	604	CLA	O1D-CGD-O2D-CED
29	s	611	CLA	O1D-CGD-O2D-CED
29	c	509	CLA	O1A-CGA-O2A-C1
29	B	605	CLA	C8-C10-C11-C12
29	B	606	CLA	C15-C16-C17-C18
29	B	607	CLA	C8-C10-C11-C12
29	B	607	CLA	C10-C11-C12-C13
29	B	617	CLA	C5-C6-C7-C8
29	D	403	CLA	C15-C16-C17-C18

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Mol	Chain	Res	Type	Atoms
29	N	604	CLA	C13-C15-C16-C17
29	R	612	CLA	C10-C11-C12-C13
29	S	603	CLA	C13-C15-C16-C17
29	S	613	CLA	C5-C6-C7-C8
29	Y	604	CLA	C15-C16-C17-C18
29	Y	611	CLA	C5-C6-C7-C8
29	Y	613	CLA	C15-C16-C17-C18
29	b	602	CLA	C5-C6-C7-C8
29	b	602	CLA	C15-C16-C17-C18
29	b	604	CLA	C15-C16-C17-C18
29	b	607	CLA	C13-C15-C16-C17
29	b	616	CLA	C5-C6-C7-C8
29	c	505	CLA	C13-C15-C16-C17
29	c	507	CLA	C15-C16-C17-C18
29	n	602	CLA	C5-C6-C7-C8
29	s	613	CLA	C5-C6-C7-C8
29	y	603	CLA	C15-C16-C17-C18
29	y	604	CLA	C5-C6-C7-C8
29	y	610	CLA	C5-C6-C7-C8
29	y	611	CLA	C8-C10-C11-C12
29	y	613	CLA	C8-C10-C11-C12
39	Y	624	LHG	O1-C1-C2-O2
29	A	410	CLA	O1A-CGA-O2A-C1
38	c	518	DGD	C1B-C2B-C3B-C4B
39	C	525	LHG	C23-C24-C25-C26
39	D	410	LHG	C7-C8-C9-C10
39	c	625	LHG	C23-C24-C25-C26
39	d	410	LHG	C7-C8-C9-C10
49	s	625	LPX	C6-C7-C8-C9
29	A	410	CLA	C8-C10-C11-C12
29	B	608	CLA	C15-C16-C17-C18
29	B	616	CLA	C5-C6-C7-C8
29	B	617	CLA	C13-C15-C16-C17
29	C	501	CLA	C13-C15-C16-C17
29	R	602	CLA	C8-C10-C11-C12
29	R	612	CLA	C8-C10-C11-C12
29	Y	603	CLA	C5-C6-C7-C8
29	Y	610	CLA	C13-C15-C16-C17
29	Y	614	CLA	C8-C10-C11-C12
29	b	608	CLA	C15-C16-C17-C18
29	b	613	CLA	C15-C16-C17-C18
29	c	503	CLA	C15-C16-C17-C18

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Mol	Chain	Res	Type	Atoms
29	c	505	CLA	C5-C6-C7-C8
29	n	604	CLA	C10-C11-C12-C13
29	r	608	CLA	C5-C6-C7-C8
29	s	603	CLA	C5-C6-C7-C8
29	s	610	CLA	C5-C6-C7-C8
29	s	610	CLA	C15-C16-C17-C18
29	C	507	CLA	CBA-CGA-O2A-C1
29	S	614	CLA	CBA-CGA-O2A-C1
29	B	613	CLA	O1D-CGD-O2D-CED
29	N	603	CLA	O1D-CGD-O2D-CED
29	B	602	CLA	C2-C1-O2A-CGA
29	B	608	CLA	C2-C1-O2A-CGA
29	B	612	CLA	C2-C1-O2A-CGA
29	C	512	CLA	C2-C1-O2A-CGA
29	N	614	CLA	C2-C1-O2A-CGA
29	Y	614	CLA	C2-C1-O2A-CGA
29	b	602	CLA	C2-C1-O2A-CGA
29	b	605	CLA	C2-C1-O2A-CGA
29	b	608	CLA	C2-C1-O2A-CGA
29	b	614	CLA	C2-C1-O2A-CGA
29	c	506	CLA	C2-C1-O2A-CGA
29	c	512	CLA	C2-C1-O2A-CGA
29	n	613	CLA	C2-C1-O2A-CGA
29	g	614	CLA	C2-C1-O2A-CGA
29	r	602	CLA	C2-C1-O2A-CGA
29	r	603	CLA	C2-C1-O2A-CGA
29	s	610	CLA	C2-C1-O2A-CGA
29	s	614	CLA	C2-C1-O2A-CGA
29	s	617	CLA	C2-C1-O2A-CGA
29	y	603	CLA	C2-C1-O2A-CGA
42	D	405	PL9	C42-C43-C44-C46
29	B	604	CLA	C5-C6-C7-C8
29	B	613	CLA	C5-C6-C7-C8
29	B	613	CLA	C13-C15-C16-C17
29	C	504	CLA	C13-C15-C16-C17
29	C	505	CLA	C5-C6-C7-C8
29	Y	613	CLA	C8-C10-C11-C12
29	b	612	CLA	C15-C16-C17-C18
29	b	614	CLA	C10-C11-C12-C13
29	c	503	CLA	C5-C6-C7-C8
29	d	402	CLA	C15-C16-C17-C18
29	d	403	CLA	C8-C10-C11-C12

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Mol	Chain	Res	Type	Atoms
29	g	613	CLA	C15-C16-C17-C18
39	D	410	LHG	C23-C24-C25-C26
39	n	624	LHG	C7-C8-C9-C10
39	s	624	LHG	C23-C24-C25-C26
43	F	101	HEM	C3D-CAD-CBD-CGD
29	Y	614	CLA	C10-C11-C12-C13
29	n	603	CLA	C8-C10-C11-C12
29	n	613	CLA	C8-C10-C11-C12
29	y	612	CLA	C13-C15-C16-C17
29	y	612	CLA	C15-C16-C17-C18
45	n	601	CHL	C8-C10-C11-C12
29	r	610	CLA	O1D-CGD-O2D-CED
29	B	603	CLA	C12-C13-C15-C16
29	B	604	CLA	C6-C7-C8-C10
29	c	506	CLA	C11-C12-C13-C15
29	s	609	CLA	C11-C10-C8-C7
29	A	405	CLA	O1A-CGA-O2A-C1
29	N	614	CLA	O1A-CGA-O2A-C1
31	b	618	BCR	C9-C10-C11-C12
31	c	517	BCR	C9-C10-C11-C12
44	H	101	RRX	C19-C20-C21-C22
44	h	101	RRX	C19-C20-C21-C22
46	G	621	LUT	C29-C30-C31-C32
46	n	621	LUT	C29-C30-C31-C32
29	B	612	CLA	C2A-CAA-CBA-CGA
29	R	603	CLA	C2A-CAA-CBA-CGA
29	S	613	CLA	C2A-CAA-CBA-CGA
29	S	614	CLA	C2A-CAA-CBA-CGA
29	Y	610	CLA	C2A-CAA-CBA-CGA
29	a	405	CLA	C2A-CAA-CBA-CGA
29	s	610	CLA	C2A-CAA-CBA-CGA
29	N	604	CLA	O1D-CGD-O2D-CED
29	y	611	CLA	O1D-CGD-O2D-CED
29	B	607	CLA	C13-C15-C16-C17
29	B	607	CLA	C15-C16-C17-C18
29	B	614	CLA	C10-C11-C12-C13
29	C	505	CLA	C13-C15-C16-C17
29	N	613	CLA	C5-C6-C7-C8
29	N	613	CLA	C8-C10-C11-C12
29	R	609	CLA	C5-C6-C7-C8
29	S	609	CLA	C10-C11-C12-C13
29	Y	612	CLA	C13-C15-C16-C17

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Mol	Chain	Res	Type	Atoms
29	b	608	CLA	C10-C11-C12-C13
29	b	614	CLA	C5-C6-C7-C8
29	c	506	CLA	C5-C6-C7-C8
29	n	602	CLA	C15-C16-C17-C18
29	r	602	CLA	C8-C10-C11-C12
29	y	611	CLA	C5-C6-C7-C8
29	y	614	CLA	C8-C10-C11-C12
29	y	614	CLA	C13-C15-C16-C17
29	C	507	CLA	O1A-CGA-O2A-C1
29	C	509	CLA	O1A-CGA-O2A-C1
29	c	503	CLA	O1A-CGA-O2A-C1
29	S	602	CLA	CBD-CGD-O2D-CED
33	b	622	LMG	O6-C1-O1-C7
33	j	101	LMG	O6-C1-O1-C7
38	c	519	DGD	O6D-C1D-O3G-C3G
29	B	603	CLA	C10-C11-C12-C13
29	B	614	CLA	C5-C6-C7-C8
29	C	506	CLA	C5-C6-C7-C8
29	C	506	CLA	C8-C10-C11-C12
29	Y	602	CLA	C8-C10-C11-C12
31	C	515	BCR	C10-C11-C12-C13
31	c	517	BCR	C10-C11-C12-C13
48	R	622	NEX	C30-C31-C32-C33
33	D	411	LMG	C35-C36-C37-C38
33	d	411	LMG	C35-C36-C37-C38
29	c	506	CLA	O1D-CGD-O2D-CED
39	D	408	LHG	O2-C2-C3-O3
39	L	101	LHG	O2-C2-C3-O3
39	l	101	LHG	O2-C2-C3-O3
38	C	519	DGD	O1B-C1B-O2G-C2G
29	n	603	CLA	C3-C5-C6-C7
29	y	614	CLA	C3-C5-C6-C7
29	B	602	CLA	C13-C15-C16-C17
29	B	610	CLA	C13-C15-C16-C17
29	B	613	CLA	C15-C16-C17-C18
29	C	502	CLA	C15-C16-C17-C18
29	G	610	CLA	C5-C6-C7-C8
29	R	602	CLA	C5-C6-C7-C8
29	S	602	CLA	C10-C11-C12-C13
29	S	610	CLA	C8-C10-C11-C12
29	b	603	CLA	C10-C11-C12-C13
29	c	508	CLA	C10-C11-C12-C13

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Mol	Chain	Res	Type	Atoms
29	g	603	CLA	C10-C11-C12-C13
29	y	611	CLA	C10-C11-C12-C13
29	y	614	CLA	C10-C11-C12-C13
45	n	607	CHL	C8-C10-C11-C12
29	B	612	CLA	CBA-CGA-O2A-C1
29	R	602	CLA	O1A-CGA-O2A-C1
29	R	604	CLA	O1A-CGA-O2A-C1
29	Y	614	CLA	O1A-CGA-O2A-C1
29	a	410	CLA	O1A-CGA-O2A-C1
29	b	614	CLA	O1A-CGA-O2A-C1
29	r	604	CLA	O1A-CGA-O2A-C1
29	s	611	CLA	O1A-CGA-O2A-C1
29	y	614	CLA	O1A-CGA-O2A-C1
39	c	625	LHG	O10-C23-O8-C6
39	s	624	LHG	C7-C8-C9-C10
29	B	604	CLA	C13-C15-C16-C17
29	C	505	CLA	C10-C11-C12-C13
29	C	506	CLA	C13-C15-C16-C17
29	C	507	CLA	C8-C10-C11-C12
29	C	513	CLA	C10-C11-C12-C13
29	R	608	CLA	C5-C6-C7-C8
29	Y	610	CLA	C15-C16-C17-C18
29	Y	613	CLA	C13-C15-C16-C17
29	b	603	CLA	C13-C15-C16-C17
29	c	507	CLA	C8-C10-C11-C12
29	c	507	CLA	C13-C15-C16-C17
29	c	513	CLA	C10-C11-C12-C13
29	g	603	CLA	C8-C10-C11-C12
45	Y	606	CHL	C15-C16-C17-C18
33	a	413	LMG	C11-C10-O7-C8
39	G	630	LHG	C8-C7-O7-C5
38	c	518	DGD	CAB-CBB-CCB-CDB
29	B	604	CLA	C8-C10-C11-C12
29	C	503	CLA	C5-C6-C7-C8
29	C	506	CLA	C15-C16-C17-C18
29	b	607	CLA	C8-C10-C11-C12
29	b	607	CLA	C10-C11-C12-C13
29	b	608	CLA	C13-C15-C16-C17
29	d	403	CLA	C15-C16-C17-C18
29	g	610	CLA	C15-C16-C17-C18
29	y	604	CLA	C8-C10-C11-C12
39	C	525	LHG	C3-O3-P-O6

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Mol	Chain	Res	Type	Atoms
39	C	525	LHG	C4-O6-P-O3
39	D	408	LHG	C3-O3-P-O6
39	D	408	LHG	C4-O6-P-O3
39	D	410	LHG	C3-O3-P-O6
39	D	410	LHG	C4-O6-P-O3
39	L	101	LHG	C3-O3-P-O6
39	N	624	LHG	C4-O6-P-O3
39	G	630	LHG	C4-O6-P-O3
39	S	624	LHG	C4-O6-P-O3
39	d	408	LHG	C4-O6-P-O3
39	d	409	LHG	C3-O3-P-O6
39	d	410	LHG	C3-O3-P-O6
39	l	101	LHG	C3-O3-P-O6
39	n	624	LHG	C4-O6-P-O3
39	g	624	LHG	C4-O6-P-O3
39	s	624	LHG	C4-O6-P-O3
39	y	624	LHG	C4-O6-P-O3
49	S	625	LPX	C1-O2-P1-O1
49	s	625	LPX	C1-O2-P1-O1
39	G	630	LHG	C7-C8-C9-C10
29	C	510	CLA	C3-C5-C6-C7
29	B	602	CLA	CBA-CGA-O2A-C1
29	n	602	CLA	CBA-CGA-O2A-C1
38	c	520	DGD	C2A-C1A-O1G-C1G
39	d	410	LHG	C24-C23-O8-C6
29	B	610	CLA	C15-C16-C17-C18
29	C	509	CLA	C10-C11-C12-C13
29	b	611	CLA	C15-C16-C17-C18
29	c	506	CLA	C13-C15-C16-C17
29	S	614	CLA	O1A-CGA-O2A-C1
29	C	506	CLA	O1D-CGD-O2D-CED
39	c	625	LHG	C1-C2-C3-O3
33	a	413	LMG	O9-C10-O7-C8
39	G	630	LHG	O9-C7-O7-C5
29	n	613	CLA	C4-C3-C5-C6
45	y	607	CHL	C4-C3-C5-C6
29	C	505	CLA	C2A-CAA-CBA-CGA
29	N	602	CLA	C2A-CAA-CBA-CGA
29	Y	602	CLA	C2A-CAA-CBA-CGA
29	Y	613	CLA	C2A-CAA-CBA-CGA
29	n	602	CLA	C2A-CAA-CBA-CGA
29	s	613	CLA	C2A-CAA-CBA-CGA

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Mol	Chain	Res	Type	Atoms
29	y	602	CLA	C2A-CAA-CBA-CGA
29	y	613	CLA	C2A-CAA-CBA-CGA
29	B	612	CLA	C16-C17-C18-C20
29	c	505	CLA	C16-C17-C18-C19
29	C	503	CLA	CBA-CGA-O2A-C1
29	c	507	CLA	CBA-CGA-O2A-C1
29	s	604	CLA	CBA-CGA-O2A-C1
38	c	518	DGD	CCB-CDB-CEB-CFB
46	g	621	LUT	C29-C30-C31-C32
46	y	621	LUT	C29-C30-C31-C32
48	S	622	NEX	C13-C14-C15-C35
39	L	101	LHG	C7-C8-C9-C10
39	S	624	LHG	C23-C24-C25-C26
36	b	623	DGA	CA6-CA7-CA8-CA9
36	c	524	DGA	CB2-CB1-OG2-CG2
39	g	624	LHG	C8-C7-O7-C5
39	y	624	LHG	C8-C7-O7-C5
50	i	101	3PH	C22-C21-O21-C2
29	c	506	CLA	C15-C16-C17-C18
29	n	610	CLA	C5-C6-C7-C8
47	R	621	XAT	C20-C13-C14-C15
47	R	621	XAT	C40-C33-C34-C35
48	N	623	NEX	C39-C29-C30-C31
48	G	623	NEX	C39-C29-C30-C31
48	R	622	NEX	C39-C29-C30-C31
48	R	622	NEX	C40-C33-C34-C35
48	S	622	NEX	C40-C33-C34-C35
48	Y	623	NEX	C39-C29-C30-C31
48	n	623	NEX	C39-C29-C30-C31
48	g	623	NEX	C39-C29-C30-C31
48	r	623	NEX	C39-C29-C30-C31
48	s	623	NEX	C11-C10-C9-C19
48	s	623	NEX	C40-C33-C34-C35
48	y	623	NEX	C39-C29-C30-C31
32	A	412	SQD	C30-C31-C32-C33
33	B	622	LMG	C15-C16-C17-C18
39	G	630	LHG	C31-C32-C33-C34
39	S	624	LHG	C30-C31-C32-C33
39	c	625	LHG	C24-C25-C26-C27
39	d	410	LHG	C26-C27-C28-C29
39	g	624	LHG	C13-C14-C15-C16
29	G	602	CLA	C16-C17-C18-C19

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Mol	Chain	Res	Type	Atoms
29	R	609	CLA	C11-C12-C13-C15
29	S	602	CLA	C11-C12-C13-C15
29	S	613	CLA	C6-C7-C8-C9
29	Y	613	CLA	C16-C17-C18-C20
29	c	513	CLA	C16-C17-C18-C19
29	n	604	CLA	C16-C17-C18-C19
29	r	609	CLA	C11-C12-C13-C14
29	r	612	CLA	C11-C12-C13-C15
29	y	604	CLA	C16-C17-C18-C19
29	y	610	CLA	C16-C17-C18-C20
32	A	412	SQD	C14-C15-C16-C17
36	c	524	DGA	CB5-CB6-CB7-CB8
39	N	624	LHG	C15-C16-C17-C18
39	G	630	LHG	C10-C11-C12-C13
39	Y	624	LHG	C30-C31-C32-C33
39	d	408	LHG	C13-C14-C15-C16
39	d	409	LHG	C26-C27-C28-C29
39	g	624	LHG	C31-C32-C33-C34
39	s	624	LHG	C30-C31-C32-C33
50	S	626	3PH	C3E-C3F-C3G-C3H
36	c	524	DGA	OB1-CB1-OG2-CG2
39	g	624	LHG	O9-C7-O7-C5
50	i	101	3PH	O22-C21-O21-C2
29	B	609	CLA	C15-C16-C17-C18
39	g	624	LHG	C7-C8-C9-C10
50	s	626	3PH	C21-C22-C23-C24
36	B	625	DGA	CA6-CA7-CA8-CA9
36	B	625	DGA	CDA-CEA-CFA-CGA
36	C	524	DGA	CDA-CEA-CFA-CGA
39	y	624	LHG	C11-C10-C9-C8
50	s	626	3PH	C3E-C3F-C3G-C3H
51	Y	625	SPH	C10-C11-C12-C13
39	D	408	LHG	C13-C14-C15-C16
39	N	624	LHG	C13-C14-C15-C16
39	n	624	LHG	C13-C14-C15-C16
39	N	624	LHG	O2-C2-C3-O3
36	C	524	DGA	CB5-CB6-CB7-CB8
38	C	520	DGD	C3B-C4B-C5B-C6B
39	D	408	LHG	C33-C34-C35-C36
39	l	101	LHG	C33-C34-C35-C36
39	n	624	LHG	C26-C27-C28-C29
29	a	405	CLA	C3-C5-C6-C7

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Mol	Chain	Res	Type	Atoms
39	y	624	LHG	C7-C8-C9-C10
49	S	625	LPX	C6-C7-C8-C9
47	R	621	XAT	C12-C13-C14-C15
47	R	621	XAT	C32-C33-C34-C35
47	r	622	XAT	C12-C13-C14-C15
48	N	623	NEX	C28-C29-C30-C31
48	G	623	NEX	C28-C29-C30-C31
48	R	622	NEX	C28-C29-C30-C31
48	R	622	NEX	C32-C33-C34-C35
48	S	622	NEX	C32-C33-C34-C35
48	Y	623	NEX	C28-C29-C30-C31
48	n	623	NEX	C28-C29-C30-C31
48	g	623	NEX	C28-C29-C30-C31
48	r	623	NEX	C28-C29-C30-C31
48	s	623	NEX	C11-C10-C9-C8
48	s	623	NEX	C32-C33-C34-C35
48	y	623	NEX	C28-C29-C30-C31
29	S	604	CLA	CBA-CGA-O2A-C1
29	S	610	CLA	CBA-CGA-O2A-C1
38	C	520	DGD	C2A-C1A-O1G-C1G
33	c	521	LMG	C29-C30-C31-C32
36	c	524	DGA	CDA-CEA-CFA-CGA
36	c	524	DGA	CB2-CB3-CB4-CB5
39	N	624	LHG	C26-C27-C28-C29
39	d	408	LHG	C28-C29-C30-C31
39	d	409	LHG	C29-C30-C31-C32
51	y	625	SPH	C7-C8-C9-C10
29	G	613	CLA	C8-C10-C11-C12
29	C	503	CLA	O1A-CGA-O2A-C1
29	B	617	CLA	C16-C17-C18-C19
29	C	507	CLA	C16-C17-C18-C19
29	C	507	CLA	C16-C17-C18-C20
29	G	613	CLA	C16-C17-C18-C19
29	c	507	CLA	C16-C17-C18-C19
29	s	613	CLA	C6-C7-C8-C10
29	y	611	CLA	C16-C17-C18-C19
29	y	613	CLA	C16-C17-C18-C19
29	c	510	CLA	C4-C3-C5-C6
32	a	412	SQD	C14-C15-C16-C17
36	B	625	DGA	CB7-CB8-CB9-CAB
39	L	101	LHG	C33-C34-C35-C36
39	c	625	LHG	C28-C29-C30-C31

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Mol	Chain	Res	Type	Atoms
29	c	510	CLA	C2-C3-C5-C6
45	y	607	CHL	C2-C3-C5-C6
29	A	405	CLA	C11-C12-C13-C14
29	C	511	CLA	C11-C10-C8-C9
29	Y	610	CLA	C11-C12-C13-C14
29	Y	613	CLA	C6-C7-C8-C9
29	Y	613	CLA	C11-C12-C13-C14
29	b	602	CLA	C6-C7-C8-C9
29	b	616	CLA	C11-C10-C8-C9
29	c	502	CLA	C14-C13-C15-C16
29	g	603	CLA	C14-C13-C15-C16
29	y	604	CLA	C11-C12-C13-C14
29	y	610	CLA	C11-C12-C13-C14
29	y	613	CLA	C11-C12-C13-C14
45	g	609	CHL	C11-C12-C13-C14
39	l	101	LHG	C7-C8-C9-C10
32	B	621	SQD	C30-C31-C32-C33
32	B	621	SQD	C32-C33-C34-C35
32	b	621	SQD	C32-C33-C34-C35
33	C	521	LMG	C29-C30-C31-C32
33	j	101	LMG	C31-C32-C33-C34
39	D	408	LHG	C28-C29-C30-C31
39	D	409	LHG	C26-C27-C28-C29
39	d	408	LHG	C33-C34-C35-C36
29	C	512	CLA	C8-C10-C11-C12
29	N	610	CLA	C15-C16-C17-C18
29	c	512	CLA	C15-C16-C17-C18
29	r	608	CLA	C10-C11-C12-C13
29	b	602	CLA	C2A-CAA-CBA-CGA
29	B	602	CLA	O1A-CGA-O2A-C1
29	c	507	CLA	O1A-CGA-O2A-C1
39	d	410	LHG	O10-C23-O8-C6
39	C	525	LHG	O1-C1-C2-C3
39	D	408	LHG	O1-C1-C2-C3
39	D	409	LHG	O1-C1-C2-C3
39	D	410	LHG	O1-C1-C2-C3
39	N	624	LHG	O1-C1-C2-C3
39	S	624	LHG	O1-C1-C2-C3
39	Y	624	LHG	O1-C1-C2-C3
39	d	409	LHG	O1-C1-C2-C3
39	n	624	LHG	O1-C1-C2-C3
39	y	624	LHG	O1-C1-C2-C3

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Mol	Chain	Res	Type	Atoms
31	A	411	BCR	C11-C12-C13-C14
31	c	515	BCR	C7-C8-C9-C10
44	H	101	RRX	C7-C8-C9-C10
29	Y	614	CLA	C3-C5-C6-C7
36	C	524	DGA	OB1-CB1-OG2-CG2
39	d	408	LHG	O9-C7-O7-C5
39	y	624	LHG	O9-C7-O7-C5
29	D	403	CLA	C13-C15-C16-C17
36	C	524	DGA	CB2-CB1-OG2-CG2
39	d	408	LHG	C8-C7-O7-C5
32	c	626	SQD	C31-C32-C33-C34
36	C	524	DGA	CA7-CA8-CA9-CAA
38	c	520	DGD	C3B-C4B-C5B-C6B
39	D	409	LHG	C29-C30-C31-C32
39	D	409	LHG	C31-C32-C33-C34
39	L	101	LHG	C13-C14-C15-C16
39	Y	624	LHG	C28-C29-C30-C31
38	C	518	DGD	C1B-C2B-C3B-C4B
39	G	630	LHG	C23-C24-C25-C26
39	S	624	LHG	C7-C8-C9-C10
39	d	410	LHG	C23-C24-C25-C26
33	B	622	LMG	C18-C19-C20-C21
33	c	521	LMG	C15-C16-C17-C18
33	c	521	LMG	C16-C17-C18-C19
36	B	625	DGA	CA4-CA5-CA6-CA7
36	b	623	DGA	CDB-CEB-CFB-CGB
38	C	523	DGD	C9B-CAB-CBB-CCB
39	l	101	LHG	C13-C14-C15-C16
39	y	624	LHG	C11-C12-C13-C14
50	S	626	3PH	C22-C23-C24-C25
50	S	626	3PH	C27-C28-C29-C2A
29	s	604	CLA	O1A-CGA-O2A-C1
29	R	608	CLA	C11-C12-C13-C15
29	R	609	CLA	C11-C12-C13-C14
29	R	612	CLA	C11-C12-C13-C15
29	S	613	CLA	C6-C7-C8-C10
29	Y	604	CLA	C16-C17-C18-C19
29	Y	604	CLA	C16-C17-C18-C20
29	b	605	CLA	C16-C17-C18-C19
29	b	617	CLA	C16-C17-C18-C19
29	r	609	CLA	C11-C12-C13-C15
29	y	602	CLA	C16-C17-C18-C19

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Mol	Chain	Res	Type	Atoms
32	C	526	SQD	O5-C1-O6-C44
29	C	506	CLA	C10-C11-C12-C13
32	b	621	SQD	C26-C27-C28-C29
39	D	409	LHG	C28-C29-C30-C31
39	Y	624	LHG	C11-C10-C9-C8
39	c	625	LHG	C11-C12-C13-C14
39	y	624	LHG	C33-C34-C35-C36
36	b	623	DGA	CEA-CFA-CGA-CHA
36	c	524	DGA	CB3-CB4-CB5-CB6
39	Y	624	LHG	C11-C12-C13-C14
39	y	624	LHG	C26-C27-C28-C29
39	g	624	LHG	C23-C24-C25-C26
50	i	101	3PH	C31-C32-C33-C34
29	B	612	CLA	C8-C10-C11-C12
29	G	610	CLA	C8-C10-C11-C12
29	G	613	CLA	C15-C16-C17-C18
29	n	610	CLA	C13-C15-C16-C17
29	n	602	CLA	O1A-CGA-O2A-C1
33	b	622	LMG	C15-C16-C17-C18
36	B	625	DGA	CB3-CB4-CB5-CB6
39	C	525	LHG	C31-C32-C33-C34
39	G	630	LHG	C11-C12-C13-C14
29	b	614	CLA	C3-C5-C6-C7
29	c	507	CLA	C3-C5-C6-C7
29	N	602	CLA	CBA-CGA-O2A-C1
36	b	623	DGA	CCA-CDA-CEA-CFA
36	b	623	DGA	CB7-CB8-CB9-CAB
39	S	624	LHG	C11-C12-C13-C14
29	c	505	CLA	O1D-CGD-O2D-CED
30	a	409	PHO	O1D-CGD-O2D-CED
29	B	609	CLA	C3A-C2A-CAA-CBA
29	B	612	CLA	C3A-C2A-CAA-CBA
29	B	615	CLA	C3A-C2A-CAA-CBA
29	C	513	CLA	C3A-C2A-CAA-CBA
29	N	603	CLA	C3A-C2A-CAA-CBA
29	G	603	CLA	C3A-C2A-CAA-CBA
29	G	611	CLA	C3A-C2A-CAA-CBA
29	Y	603	CLA	C3A-C2A-CAA-CBA
29	a	405	CLA	C3A-C2A-CAA-CBA
29	b	612	CLA	C3A-C2A-CAA-CBA
29	b	615	CLA	C3A-C2A-CAA-CBA
29	c	506	CLA	C3A-C2A-CAA-CBA

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Mol	Chain	Res	Type	Atoms
29	c	513	CLA	C3A-C2A-CAA-CBA
29	n	604	CLA	C3A-C2A-CAA-CBA
29	n	614	CLA	C3A-C2A-CAA-CBA
29	g	603	CLA	C3A-C2A-CAA-CBA
29	g	611	CLA	C3A-C2A-CAA-CBA
45	G	607	CHL	C3A-C2A-CAA-CBA
45	Y	606	CHL	C3A-C2A-CAA-CBA
45	n	609	CHL	C3A-C2A-CAA-CBA
45	g	606	CHL	C3A-C2A-CAA-CBA
45	y	606	CHL	C3A-C2A-CAA-CBA
29	b	602	CLA	C8-C10-C11-C12
29	r	603	CLA	C8-C10-C11-C12
31	c	514	BCR	C13-C14-C15-C16
32	b	621	SQD	C30-C31-C32-C33
33	C	521	LMG	C15-C16-C17-C18
36	C	524	DGA	CA6-CA7-CA8-CA9
36	C	524	DGA	CB2-CB3-CB4-CB5
36	C	524	DGA	CB3-CB4-CB5-CB6
36	b	623	DGA	CB3-CB4-CB5-CB6
39	D	408	LHG	C29-C30-C31-C32
39	L	101	LHG	C11-C12-C13-C14
39	n	624	LHG	C15-C16-C17-C18
38	c	520	DGD	O1A-C1A-O1G-C1G
29	G	613	CLA	C16-C17-C18-C20
29	R	608	CLA	C11-C12-C13-C14
29	R	612	CLA	C11-C12-C13-C14
29	S	602	CLA	C11-C12-C13-C14
29	n	604	CLA	C16-C17-C18-C20
29	g	602	CLA	C16-C17-C18-C19
29	g	602	CLA	C16-C17-C18-C20
29	r	612	CLA	C11-C12-C13-C14
29	s	613	CLA	C6-C7-C8-C9
29	y	604	CLA	C16-C17-C18-C20
32	C	526	SQD	C11-C12-C13-C14
33	J	101	LMG	C31-C32-C33-C34
39	S	624	LHG	C9-C10-C11-C12
39	c	625	LHG	C31-C32-C33-C34
39	d	408	LHG	C29-C30-C31-C32
39	y	624	LHG	C16-C17-C18-C19
50	S	626	3PH	C23-C24-C25-C26
29	C	505	CLA	O1D-CGD-O2D-CED
39	d	408	LHG	C4-C5-C6-O8

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Mol	Chain	Res	Type	Atoms
39	y	624	LHG	C31-C32-C33-C34
50	i	101	3PH	C23-C24-C25-C26
50	s	626	3PH	C25-C26-C27-C28
29	A	405	CLA	O2A-C1-C2-C3
45	n	601	CHL	C3-C5-C6-C7
39	n	624	LHG	C28-C29-C30-C31
29	B	614	CLA	C15-C16-C17-C18
29	b	604	CLA	CBA-CGA-O2A-C1
29	c	501	CLA	C2-C3-C5-C6
29	n	613	CLA	C2-C3-C5-C6
45	N	601	CHL	C2-C3-C5-C6
39	D	408	LHG	C8-C7-O7-C5
36	b	623	DGA	CA4-CA5-CA6-CA7
39	D	409	LHG	C11-C12-C13-C14
39	G	630	LHG	C28-C29-C30-C31
39	d	409	LHG	C11-C12-C13-C14
50	S	626	3PH	C24-C25-C26-C27
39	C	525	LHG	O1-C1-C2-O2
39	D	409	LHG	O1-C1-C2-O2
39	D	410	LHG	O1-C1-C2-O2
39	L	101	LHG	O1-C1-C2-O2
39	N	624	LHG	O1-C1-C2-O2
39	S	624	LHG	O1-C1-C2-O2
39	d	408	LHG	O1-C1-C2-O2
39	d	409	LHG	O1-C1-C2-O2
39	l	101	LHG	O1-C1-C2-O2
39	n	624	LHG	O1-C1-C2-O2
39	y	624	LHG	O1-C1-C2-O2
40	C	527	LMK	O9-C10-C11-C12
29	C	502	CLA	C10-C11-C12-C13
39	D	408	LHG	C25-C26-C27-C28
39	Y	624	LHG	C16-C17-C18-C19
39	d	409	LHG	C28-C29-C30-C31
39	l	101	LHG	C11-C12-C13-C14
39	y	624	LHG	C13-C14-C15-C16
50	s	626	3PH	C26-C27-C28-C29
29	B	612	CLA	O1A-CGA-O2A-C1
29	G	602	CLA	C16-C17-C18-C20
29	Y	613	CLA	C16-C17-C18-C19
29	y	602	CLA	C16-C17-C18-C20
29	y	610	CLA	C16-C17-C18-C19
39	N	624	LHG	C11-C12-C13-C14

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Mol	Chain	Res	Type	Atoms
29	B	603	CLA	C13-C15-C16-C17
29	c	503	CLA	C10-C11-C12-C13
29	c	505	CLA	C10-C11-C12-C13
39	c	625	LHG	C34-C35-C36-C37
39	G	630	LHG	C13-C14-C15-C16
39	l	101	LHG	C34-C35-C36-C37
29	s	603	CLA	C15-C16-C17-C18
36	c	524	DGA	CB9-CAB-CBB-CCB
39	D	409	LHG	C13-C14-C15-C16
39	d	408	LHG	C11-C12-C13-C14
39	d	409	LHG	C31-C32-C33-C34
39	n	624	LHG	C11-C12-C13-C14
39	D	408	LHG	O9-C7-O7-C5
29	A	406	CLA	C2-C1-O2A-CGA
29	B	607	CLA	C2-C1-O2A-CGA
29	C	502	CLA	C2-C1-O2A-CGA
29	C	503	CLA	C2-C1-O2A-CGA
29	C	506	CLA	C2-C1-O2A-CGA
29	N	603	CLA	C2-C1-O2A-CGA
29	G	602	CLA	C2-C1-O2A-CGA
29	G	603	CLA	C2-C1-O2A-CGA
29	G	614	CLA	C2-C1-O2A-CGA
29	R	604	CLA	C2-C1-O2A-CGA
29	R	609	CLA	C2-C1-O2A-CGA
29	S	611	CLA	C2-C1-O2A-CGA
29	S	617	CLA	C2-C1-O2A-CGA
29	Y	604	CLA	C2-C1-O2A-CGA
29	Y	608	CLA	C2-C1-O2A-CGA
29	a	405	CLA	C2-C1-O2A-CGA
29	a	406	CLA	C2-C1-O2A-CGA
29	a	407	CLA	C2-C1-O2A-CGA
29	b	612	CLA	C2-C1-O2A-CGA
29	b	616	CLA	C2-C1-O2A-CGA
29	c	502	CLA	C2-C1-O2A-CGA
29	g	604	CLA	C2-C1-O2A-CGA
29	r	604	CLA	C2-C1-O2A-CGA
29	r	609	CLA	C2-C1-O2A-CGA
29	s	603	CLA	C2-C1-O2A-CGA
29	s	611	CLA	C2-C1-O2A-CGA
29	y	604	CLA	C2-C1-O2A-CGA
29	y	608	CLA	C2-C1-O2A-CGA
39	C	525	LHG	C13-C14-C15-C16

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Mol	Chain	Res	Type	Atoms
29	C	507	CLA	C13-C15-C16-C17
29	N	602	CLA	C15-C16-C17-C18
29	c	506	CLA	C10-C11-C12-C13
29	N	602	CLA	O1A-CGA-O2A-C1
29	S	604	CLA	O1A-CGA-O2A-C1
33	J	101	LMG	C29-C30-C31-C32
39	g	624	LHG	C25-C26-C27-C28
39	g	624	LHG	C28-C29-C30-C31
39	d	409	LHG	C23-C24-C25-C26
29	Y	611	CLA	C3-C5-C6-C7
31	A	411	BCR	C23-C24-C25-C26
31	B	618	BCR	C5-C6-C7-C8
31	C	514	BCR	C1-C6-C7-C8
31	C	514	BCR	C5-C6-C7-C8
31	C	515	BCR	C23-C24-C25-C26
31	C	517	BCR	C23-C24-C25-C26
31	C	517	BCR	C23-C24-C25-C30
31	D	404	BCR	C1-C6-C7-C8
31	D	404	BCR	C5-C6-C7-C8
31	a	411	BCR	C23-C24-C25-C26
31	b	618	BCR	C1-C6-C7-C8
31	b	618	BCR	C5-C6-C7-C8
31	c	514	BCR	C1-C6-C7-C8
31	c	514	BCR	C5-C6-C7-C8
31	c	516	BCR	C23-C24-C25-C26
31	c	517	BCR	C23-C24-C25-C26
31	c	517	BCR	C23-C24-C25-C30
31	d	404	BCR	C1-C6-C7-C8
31	d	404	BCR	C5-C6-C7-C8
31	d	404	BCR	C23-C24-C25-C30
46	n	620	LUT	C5-C6-C7-C8
39	L	101	LHG	C34-C35-C36-C37
39	G	630	LHG	C11-C10-C9-C8
50	S	626	3PH	C25-C26-C27-C28
29	B	614	CLA	C8-C10-C11-C12
29	b	605	CLA	C8-C10-C11-C12
29	b	609	CLA	C8-C10-C11-C12
29	d	403	CLA	C10-C11-C12-C13
29	g	603	CLA	C15-C16-C17-C18
39	d	409	LHG	C8-C7-O7-C5
36	B	625	DGA	CAA-CBA-CCA-CDA
36	c	524	DGA	CA6-CA7-CA8-CA9

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Mol	Chain	Res	Type	Atoms
39	D	408	LHG	C30-C31-C32-C33
39	G	630	LHG	C25-C26-C27-C28
50	S	626	3PH	C35-C36-C37-C38
29	S	610	CLA	O1A-CGA-O2A-C1
38	C	520	DGD	O1A-C1A-O1G-C1G
36	C	524	DGA	CB4-CB5-CB6-CB7
39	Y	624	LHG	C13-C14-C15-C16
29	c	502	CLA	C4-C3-C5-C6
45	N	601	CHL	C4-C3-C5-C6
45	G	609	CHL	C4-C3-C5-C6
29	A	405	CLA	C11-C12-C13-C15
29	B	602	CLA	C6-C7-C8-C10
29	B	602	CLA	C11-C10-C8-C7
29	C	501	CLA	C12-C13-C15-C16
29	C	506	CLA	C11-C12-C13-C15
29	C	507	CLA	C11-C12-C13-C15
29	C	513	CLA	C11-C10-C8-C7
29	N	613	CLA	C2-C3-C5-C6
29	R	603	CLA	C11-C10-C8-C7
29	Y	604	CLA	C11-C10-C8-C7
29	Y	613	CLA	C11-C12-C13-C15
29	b	602	CLA	C6-C7-C8-C10
29	b	603	CLA	C12-C13-C15-C16
29	b	604	CLA	C6-C7-C8-C10
29	b	616	CLA	C11-C10-C8-C7
29	c	502	CLA	C12-C13-C15-C16
29	c	506	CLA	C11-C10-C8-C7
29	c	507	CLA	C11-C12-C13-C15
29	c	513	CLA	C11-C10-C8-C7
29	g	603	CLA	C12-C13-C15-C16
29	s	611	CLA	C6-C7-C8-C10
29	y	604	CLA	C11-C12-C13-C15
29	y	610	CLA	C11-C12-C13-C15
29	y	613	CLA	C11-C12-C13-C15
45	Y	601	CHL	C11-C10-C8-C7
45	Y	607	CHL	C11-C10-C8-C7
45	Y	609	CHL	C11-C12-C13-C15
45	n	607	CHL	C11-C12-C13-C15
29	B	614	CLA	C3-C5-C6-C7
29	b	604	CLA	O1A-CGA-O2A-C1
39	D	408	LHG	C26-C27-C28-C29
29	A	406	CLA	C5-C6-C7-C8

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Mol	Chain	Res	Type	Atoms
29	B	610	CLA	C5-C6-C7-C8
29	a	406	CLA	C5-C6-C7-C8
29	b	608	CLA	C5-C6-C7-C8
29	b	608	CLA	C8-C10-C11-C12
29	y	612	CLA	C8-C10-C11-C12
31	c	514	BCR	C15-C16-C17-C18
35	B	620	C7Z	C9-C10-C11-C12
46	S	621	LUT	C29-C30-C31-C32
46	s	621	LUT	C29-C30-C31-C32
29	B	609	CLA	C16-C17-C18-C19
29	Y	610	CLA	C16-C17-C18-C20
39	D	409	LHG	O9-C7-O7-C5
39	Y	624	LHG	O9-C7-O7-C5
39	D	409	LHG	C23-C24-C25-C26
29	B	604	CLA	CBA-CGA-O2A-C1
29	C	508	CLA	CBA-CGA-O2A-C1
29	S	611	CLA	CBA-CGA-O2A-C1
29	r	602	CLA	CBA-CGA-O2A-C1
38	C	518	DGD	C2A-C1A-O1G-C1G
38	c	518	DGD	C2A-C1A-O1G-C1G
36	c	524	DGA	CB7-CB8-CB9-CAB
38	c	523	DGD	C9B-CAB-CBB-CCB
39	D	409	LHG	C33-C34-C35-C36
29	R	613	CLA	C2A-CAA-CBA-CGA
29	S	602	CLA	C2A-CAA-CBA-CGA
29	r	612	CLA	C2A-CAA-CBA-CGA
29	y	610	CLA	C2A-CAA-CBA-CGA
30	A	408	PHO	C2A-CAA-CBA-CGA
29	B	615	CLA	C8-C10-C11-C12
29	c	512	CLA	C8-C10-C11-C12
29	s	609	CLA	C8-C10-C11-C12
33	d	411	LMG	C11-C12-C13-C14
36	b	623	DGA	CDA-CEA-CFA-CGA
39	n	624	LHG	C25-C26-C27-C28
50	i	101	3PH	C33-C34-C35-C36
50	i	101	3PH	C3E-C3F-C3G-C3H
39	c	625	LHG	C7-C8-C9-C10
29	B	608	CLA	C10-C11-C12-C13
29	B	616	CLA	C8-C10-C11-C12
29	c	502	CLA	C5-C6-C7-C8
29	g	610	CLA	C5-C6-C7-C8
32	A	412	SQD	C11-C12-C13-C14

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Mol	Chain	Res	Type	Atoms
39	d	409	LHG	C13-C14-C15-C16
39	y	624	LHG	C30-C31-C32-C33
32	B	621	SQD	C25-C26-C27-C28
39	L	101	LHG	C28-C29-C30-C31
39	d	408	LHG	C30-C31-C32-C33
29	c	508	CLA	CBA-CGA-O2A-C1
29	R	603	CLA	C11-C12-C13-C15
29	y	611	CLA	C16-C17-C18-C20
29	C	511	CLA	C10-C11-C12-C13
29	c	501	CLA	C5-C6-C7-C8
29	s	610	CLA	C10-C11-C12-C13
36	C	524	DGA	CA4-CA5-CA6-CA7
39	D	408	LHG	C27-C28-C29-C30
39	l	101	LHG	C9-C10-C11-C12
36	B	625	DGA	CB1-CB2-CB3-CB4
32	A	412	SQD	C8-C7-O47-C45
39	D	409	LHG	C8-C7-O7-C5
39	Y	624	LHG	C8-C7-O7-C5
39	c	625	LHG	C8-C7-O7-C5
39	d	410	LHG	C8-C7-O7-C5
39	D	410	LHG	O6-C4-C5-O7
32	b	621	SQD	C24-C25-C26-C27
39	d	408	LHG	C34-C35-C36-C37
47	r	622	XAT	C14-C15-C35-C34
29	A	405	CLA	C8-C10-C11-C12
29	N	610	CLA	C13-C15-C16-C17
29	Y	604	CLA	C8-C10-C11-C12
29	b	614	CLA	C15-C16-C17-C18
29	y	602	CLA	C13-C15-C16-C17
29	y	603	CLA	C8-C10-C11-C12
33	C	521	LMG	C16-C17-C18-C19
39	l	101	LHG	C28-C29-C30-C31
39	c	625	LHG	O9-C7-O7-C5
29	c	513	CLA	C3-C5-C6-C7
36	B	625	DGA	CBB-CAB-CB9-CB8
50	i	101	3PH	C2D-C2E-C2F-C2G
33	J	101	LMG	O7-C8-C9-O8
33	a	413	LMG	O1-C7-C8-O7
36	C	524	DGA	CB7-CB8-CB9-CAB
38	c	520	DGD	CAB-CBB-CCB-CDB
39	N	624	LHG	C28-C29-C30-C31
39	d	408	LHG	C26-C27-C28-C29

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Mol	Chain	Res	Type	Atoms
39	s	624	LHG	C9-C10-C11-C12
50	i	101	3PH	C22-C23-C24-C25
29	B	617	CLA	C16-C17-C18-C20
29	c	507	CLA	C16-C17-C18-C20
29	y	613	CLA	C16-C17-C18-C20
33	a	413	LMG	C16-C17-C18-C19
38	C	520	DGD	CAB-CBB-CCB-CDB
39	c	625	LHG	C11-C10-C9-C8
39	n	624	LHG	C34-C35-C36-C37
33	C	521	LMG	O6-C5-C6-O5
29	A	410	CLA	C5-C6-C7-C8
29	Y	612	CLA	C8-C10-C11-C12
29	N	613	CLA	C4-C3-C5-C6
29	c	502	CLA	C2-C3-C5-C6
29	g	602	CLA	C2-C3-C5-C6
42	D	405	PL9	C4-C3-C7-C8
42	d	405	PL9	C4-C3-C7-C8
32	B	621	SQD	C26-C27-C28-C29
29	B	602	CLA	C6-C7-C8-C9
29	B	602	CLA	C11-C10-C8-C9
29	B	604	CLA	C6-C7-C8-C9
29	C	501	CLA	C14-C13-C15-C16
29	C	503	CLA	C14-C13-C15-C16
29	C	506	CLA	C11-C10-C8-C9
29	C	507	CLA	C11-C12-C13-C14
29	G	603	CLA	C14-C13-C15-C16
29	b	604	CLA	C6-C7-C8-C9
29	c	507	CLA	C11-C12-C13-C14
29	c	512	CLA	C11-C12-C13-C14
29	g	603	CLA	C11-C12-C13-C14
29	g	613	CLA	C11-C10-C8-C9
29	s	609	CLA	C6-C7-C8-C9
29	s	609	CLA	C11-C10-C8-C9
29	s	611	CLA	C6-C7-C8-C9
45	N	607	CHL	C11-C12-C13-C14
39	C	525	LHG	C11-C12-C13-C14
29	A	405	CLA	C3-C5-C6-C7
29	C	506	CLA	C3-C5-C6-C7
29	B	604	CLA	C2A-CAA-CBA-CGA
29	R	608	CLA	C2A-CAA-CBA-CGA
29	b	604	CLA	C2A-CAA-CBA-CGA
29	r	604	CLA	C2A-CAA-CBA-CGA

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Mol	Chain	Res	Type	Atoms
45	N	606	CHL	C2A-CAA-CBA-CGA
45	N	608	CHL	C2A-CAA-CBA-CGA
45	Y	607	CHL	C2A-CAA-CBA-CGA
45	y	607	CHL	C2A-CAA-CBA-CGA
39	S	624	LHG	C12-C13-C14-C15
39	S	624	LHG	C28-C29-C30-C31
33	H	102	LMG	O6-C5-C6-O5
46	G	620	LUT	C27-C28-C29-C39
46	R	620	LUT	C31-C32-C33-C40
29	B	608	CLA	C5-C6-C7-C8
39	D	408	LHG	C11-C12-C13-C14
39	S	624	LHG	C11-C10-C9-C8
50	s	626	3PH	C32-C33-C34-C35
29	B	604	CLA	O1A-CGA-O2A-C1
29	S	611	CLA	O1A-CGA-O2A-C1
29	A	407	CLA	C1A-C2A-CAA-CBA
29	A	410	CLA	C1A-C2A-CAA-CBA
29	B	602	CLA	C1A-C2A-CAA-CBA
29	B	604	CLA	C1A-C2A-CAA-CBA
29	B	607	CLA	C1A-C2A-CAA-CBA
29	C	501	CLA	C1A-C2A-CAA-CBA
29	C	503	CLA	C1A-C2A-CAA-CBA
29	N	602	CLA	C1A-C2A-CAA-CBA
29	N	603	CLA	C1A-C2A-CAA-CBA
29	N	604	CLA	C1A-C2A-CAA-CBA
29	N	611	CLA	C1A-C2A-CAA-CBA
29	N	613	CLA	C1A-C2A-CAA-CBA
29	N	614	CLA	C1A-C2A-CAA-CBA
29	G	603	CLA	C1A-C2A-CAA-CBA
29	G	610	CLA	C1A-C2A-CAA-CBA
29	G	611	CLA	C1A-C2A-CAA-CBA
29	R	603	CLA	C1A-C2A-CAA-CBA
29	R	608	CLA	C1A-C2A-CAA-CBA
29	R	609	CLA	C1A-C2A-CAA-CBA
29	R	613	CLA	C1A-C2A-CAA-CBA
29	S	602	CLA	C1A-C2A-CAA-CBA
29	S	609	CLA	C1A-C2A-CAA-CBA
29	S	610	CLA	C1A-C2A-CAA-CBA
29	Y	602	CLA	C1A-C2A-CAA-CBA
29	Y	603	CLA	C1A-C2A-CAA-CBA
29	Y	610	CLA	C1A-C2A-CAA-CBA
29	Y	611	CLA	C1A-C2A-CAA-CBA

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Mol	Chain	Res	Type	Atoms
29	a	405	CLA	C1A-C2A-CAA-CBA
29	a	407	CLA	C1A-C2A-CAA-CBA
29	a	410	CLA	C1A-C2A-CAA-CBA
29	b	602	CLA	C1A-C2A-CAA-CBA
29	b	604	CLA	C1A-C2A-CAA-CBA
29	b	607	CLA	C1A-C2A-CAA-CBA
29	b	610	CLA	C1A-C2A-CAA-CBA
29	c	502	CLA	C1A-C2A-CAA-CBA
29	c	503	CLA	C1A-C2A-CAA-CBA
29	n	603	CLA	C1A-C2A-CAA-CBA
29	n	613	CLA	C1A-C2A-CAA-CBA
29	g	603	CLA	C1A-C2A-CAA-CBA
29	g	610	CLA	C1A-C2A-CAA-CBA
29	g	611	CLA	C1A-C2A-CAA-CBA
29	r	603	CLA	C1A-C2A-CAA-CBA
29	r	604	CLA	C1A-C2A-CAA-CBA
29	r	608	CLA	C1A-C2A-CAA-CBA
29	r	611	CLA	C1A-C2A-CAA-CBA
29	r	613	CLA	C1A-C2A-CAA-CBA
29	s	604	CLA	C1A-C2A-CAA-CBA
29	s	609	CLA	C1A-C2A-CAA-CBA
29	s	610	CLA	C1A-C2A-CAA-CBA
29	s	617	CLA	C1A-C2A-CAA-CBA
29	y	603	CLA	C1A-C2A-CAA-CBA
29	y	608	CLA	C1A-C2A-CAA-CBA
29	y	610	CLA	C1A-C2A-CAA-CBA
29	y	611	CLA	C1A-C2A-CAA-CBA
45	Y	606	CHL	C1A-C2A-CAA-CBA
45	y	606	CHL	C1A-C2A-CAA-CBA
29	R	603	CLA	C11-C12-C13-C14
29	b	605	CLA	C16-C17-C18-C20
29	b	617	CLA	C16-C17-C18-C20
32	A	412	SQD	O49-C7-O47-C45
39	d	409	LHG	O9-C7-O7-C5
39	d	410	LHG	O9-C7-O7-C5
39	L	101	LHG	C29-C30-C31-C32
31	C	517	BCR	C9-C10-C11-C12
46	N	621	LUT	C29-C30-C31-C32
46	S	620	LUT	C29-C30-C31-C32
46	s	620	LUT	C29-C30-C31-C32
29	B	609	CLA	C8-C10-C11-C12
29	R	603	CLA	C8-C10-C11-C12

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Mol	Chain	Res	Type	Atoms
29	Y	602	CLA	C5-C6-C7-C8
29	Y	604	CLA	C13-C15-C16-C17
29	b	613	CLA	C10-C11-C12-C13
29	R	610	CLA	C3-C5-C6-C7
39	C	525	LHG	C2-C3-O3-P
29	B	607	CLA	C5-C6-C7-C8
29	B	609	CLA	C10-C11-C12-C13
29	D	403	CLA	C8-C10-C11-C12
29	b	609	CLA	C15-C16-C17-C18
39	C	525	LHG	O6-C4-C5-C6
39	D	409	LHG	O6-C4-C5-C6
39	N	624	LHG	O6-C4-C5-C6
39	G	630	LHG	O6-C4-C5-C6
39	S	624	LHG	O6-C4-C5-C6
39	d	409	LHG	O6-C4-C5-C6
39	g	624	LHG	O6-C4-C5-C6
50	S	626	3PH	O11-C1-C2-C3
32	C	526	SQD	C30-C31-C32-C33
38	C	519	DGD	C2B-C3B-C4B-C5B
39	d	408	LHG	C25-C26-C27-C28
29	B	602	CLA	C10-C11-C12-C13
29	B	608	CLA	C13-C15-C16-C17
29	c	505	CLA	C16-C17-C18-C20
29	c	513	CLA	C16-C17-C18-C20
39	d	409	LHG	C33-C34-C35-C36
39	s	624	LHG	C11-C10-C9-C8
33	j	101	LMG	O6-C5-C6-O5
33	A	413	LMG	C18-C19-C20-C21
39	y	624	LHG	C32-C33-C34-C35
29	r	610	CLA	C3-C5-C6-C7
39	L	101	LHG	C9-C10-C11-C12
33	h	102	LMG	O6-C5-C6-O5
49	s	625	LPX	O1-C3-C4-C5
29	B	602	CLA	C4-C3-C5-C6
36	b	623	DGA	CB9-CAB-CBB-CCB
29	C	503	CLA	C13-C15-C16-C17
29	C	509	CLA	C8-C10-C11-C12
29	b	617	CLA	C10-C11-C12-C13
29	y	604	CLA	C10-C11-C12-C13
38	C	518	DGD	O1A-C1A-O1G-C1G
38	c	518	DGD	O1A-C1A-O1G-C1G
32	c	626	SQD	C27-C28-C29-C30

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Mol	Chain	Res	Type	Atoms
33	b	622	LMG	C18-C19-C20-C21
33	j	101	LMG	C11-C12-C13-C14
39	l	101	LHG	C29-C30-C31-C32
29	N	604	CLA	C2A-CAA-CBA-CGA
29	R	612	CLA	C2A-CAA-CBA-CGA
33	C	521	LMG	O1-C7-C8-C9
33	j	101	LMG	C7-C8-C9-O8
38	c	523	DGD	O1G-C1G-C2G-C3G
39	N	624	LHG	C34-C35-C36-C37
39	Y	624	LHG	C4-C5-C6-O8
39	c	625	LHG	C4-C5-C6-O8
39	s	624	LHG	C4-C5-C6-O8
40	c	627	LMK	O1-C7-C8-C9
50	s	626	3PH	C1-C2-C3-O31
29	B	603	CLA	C5-C6-C7-C8
29	g	602	CLA	C5-C6-C7-C8
29	s	602	CLA	C8-C10-C11-C12
29	y	613	CLA	C13-C15-C16-C17
45	n	607	CHL	C15-C16-C17-C18
36	b	623	DGA	CB5-CB6-CB7-CB8
39	C	525	LHG	C30-C31-C32-C33
29	r	602	CLA	O1A-CGA-O2A-C1
32	A	412	SQD	C45-C44-O6-C1
38	c	519	DGD	C5D-C6D-O5D-C1E
40	C	527	LMK	C8-C9-O8-C28
29	S	602	CLA	O1D-CGD-O2D-CED
39	D	408	LHG	C34-C35-C36-C37
29	Y	612	CLA	C10-C11-C12-C13
29	b	614	CLA	C8-C10-C11-C12
29	c	511	CLA	C15-C16-C17-C18
39	C	525	LHG	C34-C35-C36-C37
39	c	625	LHG	C13-C14-C15-C16
39	s	624	LHG	C29-C30-C31-C32
29	C	508	CLA	O1A-CGA-O2A-C1
39	s	624	LHG	C11-C12-C13-C14
39	s	624	LHG	C28-C29-C30-C31
29	C	507	CLA	C3-C5-C6-C7
29	s	610	CLA	C3-C5-C6-C7
32	c	626	SQD	O5-C1-O6-C44
33	B	622	LMG	O6-C1-O1-C7
29	b	604	CLA	C5-C6-C7-C8
29	b	612	CLA	C10-C11-C12-C13

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Mol	Chain	Res	Type	Atoms
29	n	610	CLA	C15-C16-C17-C18
42	D	405	PL9	C34-C36-C37-C38
51	y	625	SPH	C10-C11-C12-C13
33	D	411	LMG	O6-C5-C6-O5
33	c	521	LMG	O6-C5-C6-O5
39	G	630	LHG	O1-C1-C2-O2
39	d	410	LHG	O1-C1-C2-O2
39	g	624	LHG	O1-C1-C2-O2
39	s	624	LHG	O1-C1-C2-O2
38	c	519	DGD	C9A-CAA-CBA-CCA
50	S	626	3PH	C37-C38-C39-C3A
45	S	608	CHL	C5-C6-C7-C8
45	n	607	CHL	C10-C11-C12-C13
32	C	526	SQD	C7-C8-C9-C10
33	b	622	LMG	C28-C29-C30-C31
39	n	624	LHG	C31-C32-C33-C34
38	c	520	DGD	CCB-CDB-CEB-CFB
50	i	101	3PH	C34-C35-C36-C37
29	R	610	CLA	C5-C6-C7-C8
29	c	504	CLA	C8-C10-C11-C12
29	c	513	CLA	C15-C16-C17-C18
45	n	601	CHL	C10-C11-C12-C13
45	y	606	CHL	C15-C16-C17-C18
33	d	411	LMG	O6-C5-C6-O5
29	B	605	CLA	C16-C17-C18-C19
29	c	504	CLA	C10-C11-C12-C13
45	G	601	CHL	C10-C11-C12-C13
33	b	622	LMG	C9-C8-O7-C10
36	B	625	DGA	CG1-CG2-OG2-CB1
36	b	623	DGA	CG1-CG2-OG2-CB1
29	b	616	CLA	C13-C15-C16-C17
45	s	608	CHL	C5-C6-C7-C8
29	A	405	CLA	C2-C1-O2A-CGA
29	A	407	CLA	C2-C1-O2A-CGA
29	B	603	CLA	C2-C1-O2A-CGA
29	B	611	CLA	C2-C1-O2A-CGA
29	B	615	CLA	C2-C1-O2A-CGA
29	N	613	CLA	C2-C1-O2A-CGA
29	b	607	CLA	C2-C1-O2A-CGA
45	n	607	CHL	C2-C1-O2A-CGA
39	S	624	LHG	C26-C27-C28-C29
39	s	624	LHG	C31-C32-C33-C34

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Mol	Chain	Res	Type	Atoms
38	C	519	DGD	C9A-CAA-CBA-CCA
29	b	607	CLA	C5-C6-C7-C8
29	y	602	CLA	C8-C10-C11-C12
36	c	524	DGA	CA4-CA5-CA6-CA7
39	N	624	LHG	C25-C26-C27-C28
39	Y	624	LHG	C32-C33-C34-C35
39	d	410	LHG	C28-C29-C30-C31
39	s	624	LHG	C34-C35-C36-C37
29	Y	610	CLA	CBA-CGA-O2A-C1
29	s	610	CLA	CBA-CGA-O2A-C1
39	C	525	LHG	C24-C23-O8-C6
29	B	609	CLA	C16-C17-C18-C20
29	Y	610	CLA	C16-C17-C18-C19
38	C	519	DGD	C9B-CAB-CBB-CCB
39	S	624	LHG	C31-C32-C33-C34
38	c	518	DGD	O6D-C5D-C6D-O5D
29	r	609	CLA	C10-C11-C12-C13
36	C	524	DGA	CAB-CBB-CCB-CDB
29	c	508	CLA	O1A-CGA-O2A-C1
29	s	610	CLA	O1A-CGA-O2A-C1
36	c	524	DGA	CB4-CB5-CB6-CB7
39	D	410	LHG	C30-C31-C32-C33
33	B	622	LMG	C28-C29-C30-C31
29	B	606	CLA	C5-C6-C7-C8
29	B	608	CLA	C8-C10-C11-C12
29	b	609	CLA	C13-C15-C16-C17
29	b	613	CLA	C13-C15-C16-C17
29	n	604	CLA	C13-C15-C16-C17
39	N	624	LHG	C31-C32-C33-C34
29	N	613	CLA	CAA-CBA-CGA-O2A
33	A	413	LMG	O1-C7-C8-O7
38	C	523	DGD	O1G-C1G-C2G-O2G
50	i	101	3PH	O21-C2-C3-O31
29	C	501	CLA	C5-C6-C7-C8
39	N	624	LHG	C23-C24-C25-C26
38	C	520	DGD	CCB-CDB-CEB-CFB
39	D	409	LHG	C34-C35-C36-C37
29	B	602	CLA	C2-C3-C5-C6
29	B	613	CLA	C6-C7-C8-C10
29	C	501	CLA	C11-C10-C8-C7
29	C	503	CLA	C12-C13-C15-C16
29	C	506	CLA	C11-C10-C8-C7

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Mol	Chain	Res	Type	Atoms
29	C	506	CLA	C12-C13-C15-C16
29	C	508	CLA	C11-C10-C8-C7
29	C	508	CLA	C12-C13-C15-C16
29	C	511	CLA	C11-C10-C8-C7
29	D	402	CLA	C11-C10-C8-C7
29	D	402	CLA	C11-C12-C13-C15
29	N	613	CLA	C12-C13-C15-C16
29	G	603	CLA	C11-C12-C13-C15
29	G	603	CLA	C12-C13-C15-C16
29	G	613	CLA	C6-C7-C8-C10
29	S	603	CLA	C6-C7-C8-C10
29	Y	610	CLA	C11-C12-C13-C15
29	Y	611	CLA	C11-C10-C8-C7
29	Y	613	CLA	C6-C7-C8-C10
29	b	602	CLA	C11-C10-C8-C7
29	b	602	CLA	C11-C12-C13-C15
29	c	506	CLA	C12-C13-C15-C16
29	c	510	CLA	C6-C7-C8-C10
29	c	511	CLA	C11-C12-C13-C15
29	c	512	CLA	C11-C12-C13-C15
29	c	513	CLA	C6-C7-C8-C10
29	c	513	CLA	C11-C12-C13-C15
29	g	603	CLA	C11-C12-C13-C15
29	g	613	CLA	C11-C10-C8-C7
29	r	603	CLA	C11-C10-C8-C7
29	s	610	CLA	C6-C7-C8-C10
29	s	610	CLA	C11-C10-C8-C7
29	y	602	CLA	C6-C7-C8-C10
29	y	604	CLA	C12-C13-C15-C16
29	y	611	CLA	C11-C10-C8-C7
29	y	611	CLA	C12-C13-C15-C16
29	y	612	CLA	C11-C12-C13-C15
42	D	405	PL9	C43-C44-C46-C47
45	N	607	CHL	C11-C10-C8-C7
45	N	609	CHL	C11-C10-C8-C7
45	G	609	CHL	C11-C12-C13-C15
45	n	605	CHL	C12-C13-C15-C16
45	n	606	CHL	C11-C10-C8-C7
45	n	607	CHL	C12-C13-C15-C16
45	g	601	CHL	C11-C12-C13-C15
45	y	606	CHL	C11-C10-C8-C7
45	y	609	CHL	C11-C12-C13-C15

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Mol	Chain	Res	Type	Atoms
29	B	608	CLA	C14-C13-C15-C16
29	B	617	CLA	C11-C12-C13-C14
29	C	501	CLA	C11-C10-C8-C9
29	C	508	CLA	C11-C10-C8-C9
29	C	512	CLA	C11-C10-C8-C9
29	C	513	CLA	C11-C12-C13-C14
29	D	402	CLA	C6-C7-C8-C9
29	N	602	CLA	C14-C13-C15-C16
29	N	613	CLA	C14-C13-C15-C16
29	G	602	CLA	C6-C7-C8-C9
29	G	603	CLA	C11-C12-C13-C14
29	G	613	CLA	C6-C7-C8-C9
29	S	611	CLA	C6-C7-C8-C9
29	Y	604	CLA	C11-C12-C13-C14
29	Y	611	CLA	C11-C10-C8-C9
29	a	405	CLA	C11-C12-C13-C14
29	b	602	CLA	C11-C10-C8-C9
29	c	506	CLA	C11-C10-C8-C9
29	c	508	CLA	C11-C10-C8-C9
29	c	508	CLA	C14-C13-C15-C16
29	c	509	CLA	C6-C7-C8-C9
29	c	510	CLA	C6-C7-C8-C9
29	c	512	CLA	C14-C13-C15-C16
29	c	513	CLA	C11-C12-C13-C14
29	n	603	CLA	C6-C7-C8-C9
29	g	613	CLA	C6-C7-C8-C9
29	g	613	CLA	C14-C13-C15-C16
29	s	610	CLA	C11-C10-C8-C9
29	y	602	CLA	C6-C7-C8-C9
29	y	604	CLA	C14-C13-C15-C16
29	y	611	CLA	C11-C10-C8-C9
45	G	609	CHL	C11-C12-C13-C14
45	Y	609	CHL	C14-C13-C15-C16
45	n	607	CHL	C11-C12-C13-C14
45	y	606	CHL	C11-C10-C8-C9
31	A	411	BCR	C9-C10-C11-C12
31	B	619	BCR	C9-C10-C11-C12
29	d	402	CLA	C13-C15-C16-C17
29	B	605	CLA	C2A-CAA-CBA-CGA
30	a	408	PHO	C2A-CAA-CBA-CGA
45	N	605	CHL	C2A-CAA-CBA-CGA
45	g	606	CHL	C2A-CAA-CBA-CGA

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Mol	Chain	Res	Type	Atoms
39	c	625	LHG	C35-C36-C37-C38
29	n	610	CLA	C16-C17-C18-C20
46	Y	620	LUT	C27-C28-C29-C30
36	c	524	DGA	CA7-CA8-CA9-CAA
29	C	513	CLA	C3-C5-C6-C7
29	b	603	CLA	C5-C6-C7-C8
39	D	410	LHG	C26-C27-C28-C29
39	L	101	LHG	C35-C36-C37-C38
42	d	405	PL9	C42-C43-C44-C46
29	b	610	CLA	C13-C15-C16-C17
45	n	609	CHL	C8-C10-C11-C12
39	l	101	LHG	C25-C26-C27-C28
29	s	602	CLA	CBD-CGD-O2D-CED
29	Y	603	CLA	C15-C16-C17-C18
39	D	410	LHG	O6-C4-C5-C6
39	d	410	LHG	O6-C4-C5-C6
39	l	101	LHG	O6-C4-C5-C6
39	n	624	LHG	O6-C4-C5-C6
39	n	624	LHG	C23-C24-C25-C26
29	y	610	CLA	CBA-CGA-O2A-C1
50	s	626	3PH	C39-C3A-C3B-C3C
29	G	610	CLA	C4-C3-C5-C6
45	N	609	CHL	C4-C3-C5-C6
42	d	405	PL9	C13-C14-C16-C17
33	H	102	LMG	C28-C29-C30-C31
50	S	626	3PH	C3D-C3E-C3F-C3G
29	B	604	CLA	C10-C11-C12-C13
29	Y	614	CLA	C5-C6-C7-C8
32	c	626	SQD	C32-C33-C34-C35
33	D	411	LMG	C11-C12-C13-C14
39	L	101	LHG	C25-C26-C27-C28
49	s	625	LPX	C14-C15-C16-C17
29	Y	604	CLA	C5-C6-C7-C8
29	b	613	CLA	C8-C10-C11-C12
39	s	624	LHG	C24-C23-O8-C6
39	C	525	LHG	C10-C11-C12-C13
50	s	626	3PH	C3F-C3G-C3H-C3I
29	B	613	CLA	C3A-C2A-CAA-CBA
29	C	506	CLA	C3A-C2A-CAA-CBA
29	N	604	CLA	C3A-C2A-CAA-CBA
29	N	613	CLA	C3A-C2A-CAA-CBA
29	G	604	CLA	C3A-C2A-CAA-CBA

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Mol	Chain	Res	Type	Atoms
29	R	604	CLA	C3A-C2A-CAA-CBA
29	b	613	CLA	C3A-C2A-CAA-CBA
29	n	613	CLA	C3A-C2A-CAA-CBA
29	r	604	CLA	C3A-C2A-CAA-CBA
29	r	610	CLA	C3A-C2A-CAA-CBA
29	s	614	CLA	C3A-C2A-CAA-CBA
30	a	408	PHO	C3A-C2A-CAA-CBA
45	Y	601	CHL	C3A-C2A-CAA-CBA
36	C	524	DGA	CB9-CAB-CBB-CCB
39	n	624	LHG	C35-C36-C37-C38
50	S	626	3PH	C3F-C3G-C3H-C3I
31	a	411	BCR	C9-C10-C11-C12
46	Y	621	LUT	C33-C34-C35-C15
39	C	525	LHG	C35-C36-C37-C38
39	d	410	LHG	C33-C34-C35-C36
39	l	101	LHG	C35-C36-C37-C38
29	G	610	CLA	C13-C15-C16-C17
39	d	408	LHG	C27-C28-C29-C30
29	y	608	CLA	CBA-CGA-O2A-C1
36	B	625	DGA	CA8-CA9-CAA-CBA
38	c	519	DGD	C9B-CAB-CBB-CCB
32	B	621	SQD	C44-C45-C46-O48
32	b	621	SQD	C44-C45-C46-O48
33	A	413	LMG	C7-C8-C9-O8
33	J	101	LMG	C7-C8-C9-O8
33	a	413	LMG	O1-C7-C8-C9
39	D	410	LHG	C4-C5-C6-O8
39	G	630	LHG	C4-C5-C6-O8
39	d	410	LHG	C4-C5-C6-O8
39	l	101	LHG	C4-C5-C6-O8
40	C	527	LMK	O1-C7-C8-C9
46	y	621	LUT	C21-C26-C27-C28
36	B	625	DGA	CCB-CDB-CEB-CFB
36	b	623	DGA	CFB-CGB-CHB-CIB
39	y	624	LHG	C9-C10-C11-C12
36	C	524	DGA	CA8-CA9-CAA-CBA
51	y	625	SPH	C6-C7-C8-C9
29	C	501	CLA	C3-C5-C6-C7
29	Y	610	CLA	O1A-CGA-O2A-C1
45	N	609	CHL	C2-C3-C5-C6
39	D	410	LHG	C33-C34-C35-C36
39	S	624	LHG	C34-C35-C36-C37

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Mol	Chain	Res	Type	Atoms
50	i	101	3PH	C25-C26-C27-C28
29	N	602	CLA	C13-C15-C16-C17
40	C	527	LMK	C9-C8-O7-C10
40	c	627	LMK	C9-C8-O7-C10
36	b	623	DGA	CB1-CB2-CB3-CB4
39	C	525	LHG	O10-C23-O8-C6
29	G	614	CLA	C2A-CAA-CBA-CGA
29	b	605	CLA	C2A-CAA-CBA-CGA
39	D	408	LHG	O1-C1-C2-O2
39	d	409	LHG	C25-C26-C27-C28
39	D	408	LHG	O6-C4-C5-O7
39	D	409	LHG	O6-C4-C5-O7
39	d	410	LHG	O6-C4-C5-O7
39	l	101	LHG	O6-C4-C5-O7
39	n	624	LHG	O6-C4-C5-O7
39	g	624	LHG	O6-C4-C5-O7
39	s	624	LHG	O6-C4-C5-O7
38	c	518	DGD	C4D-C5D-C6D-O5D
39	N	624	LHG	C24-C23-O8-C6
33	h	102	LMG	C28-C29-C30-C31
29	B	612	CLA	C16-C17-C18-C19
51	y	625	SPH	C11-C10-C9-C8
39	N	624	LHG	C35-C36-C37-C38
51	Y	625	SPH	C11-C12-C13-C14
29	G	613	CLA	C3-C5-C6-C7
32	B	621	SQD	O47-C45-C46-O48
32	b	621	SQD	O47-C45-C46-O48
33	C	521	LMG	O1-C7-C8-O7
38	C	519	DGD	O1G-C1G-C2G-O2G
39	Y	624	LHG	O7-C5-C6-O8
39	c	625	LHG	O7-C5-C6-O8
39	l	101	LHG	O7-C5-C6-O8
29	C	512	CLA	C13-C15-C16-C17
29	R	602	CLA	C10-C11-C12-C13
29	y	614	CLA	C5-C6-C7-C8
29	s	602	CLA	O1D-CGD-O2D-CED
32	B	621	SQD	C10-C11-C12-C13
32	a	412	SQD	C28-C29-C30-C31
33	J	101	LMG	O6-C1-O1-C7
38	C	519	DGD	O6D-C1D-O3G-C3G
29	A	405	CLA	C13-C15-C16-C17
29	C	511	CLA	C15-C16-C17-C18

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Mol	Chain	Res	Type	Atoms
29	b	604	CLA	C13-C15-C16-C17
36	b	623	DGA	CG1-CG2-CG3-OXT
29	A	410	CLA	C2-C1-O2A-CGA
29	a	410	CLA	C2-C1-O2A-CGA
29	b	617	CLA	C2-C1-O2A-CGA
30	a	409	PHO	C2-C1-O2A-CGA
36	B	625	DGA	CB9-CAB-CBB-CCB
39	N	624	LHG	C16-C17-C18-C19
39	n	624	LHG	C16-C17-C18-C19
39	g	624	LHG	C11-C12-C13-C14
29	c	511	CLA	C13-C15-C16-C17
29	d	403	CLA	C13-C15-C16-C17
29	y	610	CLA	C10-C11-C12-C13
29	B	606	CLA	C6-C7-C8-C9
29	B	613	CLA	C6-C7-C8-C9
29	B	616	CLA	C14-C13-C15-C16
29	C	506	CLA	C14-C13-C15-C16
29	C	508	CLA	C14-C13-C15-C16
29	C	512	CLA	C14-C13-C15-C16
29	D	402	CLA	C11-C12-C13-C14
29	N	604	CLA	C6-C7-C8-C9
29	N	604	CLA	C14-C13-C15-C16
29	b	604	CLA	C14-C13-C15-C16
29	b	605	CLA	C11-C12-C13-C14
29	c	506	CLA	C14-C13-C15-C16
29	d	402	CLA	C6-C7-C8-C9
29	g	602	CLA	C6-C7-C8-C9
29	y	602	CLA	C11-C10-C8-C9
45	N	606	CHL	C11-C12-C13-C14
45	G	609	CHL	C14-C13-C15-C16
45	g	601	CHL	C11-C12-C13-C14
39	D	410	LHG	C28-C29-C30-C31
32	B	621	SQD	C24-C25-C26-C27
50	i	101	3PH	C38-C39-C3A-C3B
50	s	626	3PH	C38-C39-C3A-C3B
29	C	503	CLA	C8-C10-C11-C12
29	s	609	CLA	C10-C11-C12-C13
30	a	408	PHO	C1A-C2A-CAA-CBA
39	D	409	LHG	C2-C3-O3-P
39	d	409	LHG	C2-C3-O3-P
29	A	406	CLA	C2A-CAA-CBA-CGA
29	C	506	CLA	C2A-CAA-CBA-CGA

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Mol	Chain	Res	Type	Atoms
29	R	604	CLA	C2A-CAA-CBA-CGA
29	c	506	CLA	C2A-CAA-CBA-CGA
45	n	607	CHL	C2A-CAA-CBA-CGA
29	B	605	CLA	C16-C17-C18-C20
31	C	516	BCR	C23-C24-C25-C26
31	C	516	BCR	C23-C24-C25-C30
31	D	404	BCR	C23-C24-C25-C26
31	D	404	BCR	C23-C24-C25-C30
31	c	515	BCR	C23-C24-C25-C26
31	c	515	BCR	C23-C24-C25-C30
31	d	404	BCR	C23-C24-C25-C26
35	B	620	C7Z	C21-C26-C27-C28
35	b	620	C7Z	C21-C26-C27-C28
44	H	101	RRX	C23-C24-C25-C30
44	h	101	RRX	C23-C24-C25-C30
46	N	620	LUT	C1-C6-C7-C8
46	R	620	LUT	C1-C6-C7-C8
46	R	620	LUT	C5-C6-C7-C8
46	n	621	LUT	C1-C6-C7-C8
46	n	621	LUT	C5-C6-C7-C8
46	r	620	LUT	C1-C6-C7-C8
46	r	620	LUT	C5-C6-C7-C8
29	C	510	CLA	C15-C16-C17-C18
32	B	621	SQD	C18-C19-C20-C21
33	a	413	LMG	C30-C31-C32-C33
33	H	102	LMG	O7-C10-C11-C12
33	h	102	LMG	O7-C10-C11-C12
42	d	405	PL9	C47-C48-C49-C51
36	b	623	DGA	CBB-CAB-CB9-CB8
31	B	618	BCR	C21-C22-C23-C24
46	R	620	LUT	C31-C32-C33-C34
46	r	620	LUT	C31-C32-C33-C34
29	y	614	CLA	C15-C16-C17-C18
48	s	623	NEX	C14-C15-C35-C34
39	s	624	LHG	C13-C14-C15-C16
39	D	409	LHG	C19-C20-C21-C22
29	Y	610	CLA	C8-C10-C11-C12
29	A	405	CLA	C10-C11-C12-C13
29	C	512	CLA	C5-C6-C7-C8
29	C	512	CLA	C15-C16-C17-C18
39	D	408	LHG	O6-C4-C5-C6
39	L	101	LHG	O6-C4-C5-C6

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Mol	Chain	Res	Type	Atoms
36	B	625	DGA	CB5-CB6-CB7-CB8
39	D	409	LHG	C24-C25-C26-C27
29	B	603	CLA	C6-C7-C8-C10
29	B	605	CLA	C11-C12-C13-C15
29	B	608	CLA	C12-C13-C15-C16
29	B	616	CLA	C12-C13-C15-C16
29	C	507	CLA	C11-C10-C8-C7
29	C	512	CLA	C11-C10-C8-C7
29	C	512	CLA	C12-C13-C15-C16
29	C	513	CLA	C6-C7-C8-C10
29	C	513	CLA	C11-C12-C13-C15
29	D	403	CLA	C11-C12-C13-C15
29	N	602	CLA	C11-C10-C8-C7
29	N	604	CLA	C6-C7-C8-C10
29	N	613	CLA	C11-C12-C13-C15
29	G	602	CLA	C6-C7-C8-C10
29	S	611	CLA	C6-C7-C8-C10
29	Y	602	CLA	C6-C7-C8-C10
29	Y	604	CLA	C11-C12-C13-C15
29	Y	614	CLA	C11-C10-C8-C7
29	a	405	CLA	C11-C12-C13-C15
29	b	617	CLA	C11-C12-C13-C15
29	c	504	CLA	C6-C7-C8-C10
29	c	507	CLA	C12-C13-C15-C16
29	c	508	CLA	C11-C10-C8-C7
29	c	508	CLA	C12-C13-C15-C16
29	c	512	CLA	C12-C13-C15-C16
29	n	603	CLA	C6-C7-C8-C10
29	n	604	CLA	C6-C7-C8-C10
29	n	613	CLA	C12-C13-C15-C16
29	g	602	CLA	C6-C7-C8-C10
29	g	613	CLA	C6-C7-C8-C10
29	g	613	CLA	C12-C13-C15-C16
29	s	603	CLA	C11-C12-C13-C15
45	N	605	CHL	C11-C10-C8-C7
45	N	605	CHL	C12-C13-C15-C16
45	N	606	CHL	C11-C12-C13-C15
45	Y	609	CHL	C12-C13-C15-C16
45	G	601	CHL	C3-C5-C6-C7
33	B	622	LMG	C14-C15-C16-C17
31	C	514	BCR	C15-C16-C17-C18
31	C	516	BCR	C9-C10-C11-C12

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Mol	Chain	Res	Type	Atoms
31	C	516	BCR	C13-C14-C15-C16
31	b	619	BCR	C9-C10-C11-C12
31	b	619	BCR	C13-C14-C15-C16
31	c	515	BCR	C19-C20-C21-C22
44	H	101	RRX	C9-C10-C11-C12
44	h	101	RRX	C9-C10-C11-C12
46	g	620	LUT	C29-C30-C31-C32
48	s	623	NEX	C13-C14-C15-C35
39	d	409	LHG	C19-C20-C21-C22
29	B	612	CLA	C10-C11-C12-C13
29	R	609	CLA	C10-C11-C12-C13
29	r	609	CLA	CAA-CBA-CGA-O2A
29	S	604	CLA	C2A-CAA-CBA-CGA
29	c	512	CLA	C2A-CAA-CBA-CGA
29	n	604	CLA	C2A-CAA-CBA-CGA
29	r	608	CLA	C2A-CAA-CBA-CGA
29	s	614	CLA	C2A-CAA-CBA-CGA
40	C	527	LMK	O1-C1-C2-C3
29	C	509	CLA	C5-C6-C7-C8
48	r	623	NEX	C40-C33-C34-C35
48	s	623	NEX	C20-C13-C14-C15
50	s	626	3PH	C23-C24-C25-C26
38	c	523	DGD	O6D-C5D-C6D-O5D
45	g	609	CHL	C3-C5-C6-C7
29	n	610	CLA	C16-C17-C18-C19
42	D	405	PL9	C47-C48-C49-C50
36	c	524	DGA	CA8-CA9-CAA-CBA
39	s	624	LHG	C12-C13-C14-C15
50	S	626	3PH	C38-C39-C3A-C3B
50	s	626	3PH	C34-C35-C36-C37
29	B	613	CLA	CAD-CBD-CGD-O2D
29	B	616	CLA	CAD-CBD-CGD-O2D
29	B	617	CLA	CAD-CBD-CGD-O2D
29	C	502	CLA	CAD-CBD-CGD-O2D
29	N	602	CLA	CAD-CBD-CGD-O2D
29	G	603	CLA	CAD-CBD-CGD-O2D
29	G	614	CLA	CAD-CBD-CGD-O2D
29	c	509	CLA	CAD-CBD-CGD-O2D
29	n	602	CLA	CAD-CBD-CGD-O2D
38	C	523	DGD	C1G-C2G-O2G-C1B
38	c	523	DGD	C1G-C2G-O2G-C1B
45	Y	606	CHL	CAD-CBD-CGD-O2D

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Mol	Chain	Res	Type	Atoms
48	S	622	NEX	C7-C8-C9-C19
48	s	623	NEX	C7-C8-C9-C19
32	C	526	SQD	C32-C33-C34-C35
33	A	413	LMG	C29-C30-C31-C32
29	B	609	CLA	C5-C6-C7-C8
29	N	604	CLA	C15-C16-C17-C18
29	b	615	CLA	C8-C10-C11-C12
29	g	603	CLA	C13-C15-C16-C17
39	Y	624	LHG	C25-C26-C27-C28
29	r	603	CLA	C4-C3-C5-C6
29	r	609	CLA	C4-C3-C5-C6
33	b	622	LMG	C14-C15-C16-C17
33	a	413	LMG	O6-C1-O1-C7
29	r	609	CLA	C2-C3-C5-C6
45	n	605	CHL	C2-C3-C5-C6
33	c	521	LMG	O1-C7-C8-C9
38	c	520	DGD	O1G-C1G-C2G-C3G
39	D	408	LHG	C4-C5-C6-O8
39	L	101	LHG	C4-C5-C6-O8
39	S	624	LHG	C4-C5-C6-O8
39	g	624	LHG	C4-C5-C6-O8
29	y	610	CLA	O1A-CGA-O2A-C1
39	L	101	LHG	O6-C4-C5-O7
39	N	624	LHG	O6-C4-C5-O7
39	c	625	LHG	O6-C4-C5-O7
29	B	602	CLA	C8-C10-C11-C12
43	F	101	HEM	C4B-C3B-CAB-CBB
43	f	101	HEM	C4B-C3B-CAB-CBB
36	b	623	DGA	CA8-CA9-CAA-CBA
36	b	623	DGA	CCB-CDB-CEB-CFB
39	y	624	LHG	C28-C29-C30-C31
39	S	624	LHG	C1-C2-C3-O3
39	d	410	LHG	C1-C2-C3-O3
29	N	604	CLA	CHA-CBD-CGD-O1D
29	N	604	CLA	CHA-CBD-CGD-O2D
29	G	602	CLA	CHA-CBD-CGD-O1D
29	G	602	CLA	CHA-CBD-CGD-O2D
29	R	604	CLA	CHA-CBD-CGD-O1D
29	R	604	CLA	CHA-CBD-CGD-O2D
29	R	612	CLA	CHA-CBD-CGD-O1D
29	R	612	CLA	CHA-CBD-CGD-O2D
29	Y	603	CLA	CHA-CBD-CGD-O1D

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Mol	Chain	Res	Type	Atoms
29	Y	603	CLA	CHA-CBD-CGD-O2D
29	Y	604	CLA	CHA-CBD-CGD-O1D
29	Y	612	CLA	CHA-CBD-CGD-O1D
29	Y	612	CLA	CHA-CBD-CGD-O2D
29	c	512	CLA	CHA-CBD-CGD-O2D
29	g	604	CLA	CHA-CBD-CGD-O1D
29	g	604	CLA	CHA-CBD-CGD-O2D
29	g	610	CLA	CHA-CBD-CGD-O1D
29	r	612	CLA	CHA-CBD-CGD-O1D
29	r	612	CLA	CHA-CBD-CGD-O2D
29	r	613	CLA	CHA-CBD-CGD-O1D
29	r	613	CLA	CHA-CBD-CGD-O2D
29	y	612	CLA	CHA-CBD-CGD-O1D
29	y	612	CLA	CHA-CBD-CGD-O2D
40	C	527	LMK	C2-C3-C4-O2
40	c	627	LMK	C2-C3-C4-O2
45	N	601	CHL	CHA-CBD-CGD-O1D
45	n	601	CHL	CHA-CBD-CGD-O1D
45	g	605	CHL	CHA-CBD-CGD-O2D
39	D	409	LHG	C25-C26-C27-C28
29	y	608	CLA	O1A-CGA-O2A-C1
32	c	626	SQD	C28-C29-C30-C31
33	A	413	LMG	O7-C8-C9-O8
33	H	102	LMG	O7-C8-C9-O8
38	c	519	DGD	O1G-C1G-C2G-O2G
39	S	624	LHG	O7-C5-C6-O8
40	c	627	LMK	O1-C7-C8-O7
29	B	610	CLA	CBA-CGA-O2A-C1
39	d	410	LHG	C30-C31-C32-C33
29	G	613	CLA	C5-C6-C7-C8
29	b	606	CLA	C5-C6-C7-C8
38	c	523	DGD	CCB-CDB-CEB-CFB
39	Y	624	LHG	C31-C32-C33-C34
29	b	608	CLA	C16-C17-C18-C20
29	C	509	CLA	C13-C15-C16-C17
45	n	605	CHL	C4-C3-C5-C6
39	s	624	LHG	O10-C23-O8-C6
39	L	101	LHG	C17-C18-C19-C20
29	B	615	CLA	C11-C12-C13-C14
29	D	403	CLA	C11-C12-C13-C14
29	N	613	CLA	C11-C12-C13-C14
29	G	613	CLA	C11-C10-C8-C9

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Mol	Chain	Res	Type	Atoms
29	b	603	CLA	C6-C7-C8-C9
29	b	606	CLA	C6-C7-C8-C9
29	c	503	CLA	C11-C12-C13-C14
45	N	605	CHL	C11-C10-C8-C9
45	N	609	CHL	C14-C13-C15-C16
45	Y	606	CHL	C11-C10-C8-C9
45	n	601	CHL	C11-C10-C8-C9
39	D	410	LHG	C11-C10-C9-C8
39	N	624	LHG	O10-C23-O8-C6
39	Y	624	LHG	C9-C10-C11-C12
50	i	101	3PH	C3B-C3C-C3D-C3E
32	c	626	SQD	C4-C5-C6-S
29	n	602	CLA	C16-C17-C18-C19
29	a	406	CLA	C2A-CAA-CBA-CGA
29	b	617	CLA	C2A-CAA-CBA-CGA
45	N	607	CHL	C2A-CAA-CBA-CGA
39	n	624	LHG	O8-C23-C24-C25
39	g	624	LHG	C12-C13-C14-C15
46	G	620	LUT	C27-C28-C29-C30
32	c	626	SQD	C12-C13-C14-C15
33	H	102	LMG	C15-C16-C17-C18
29	R	611	CLA	C1A-C2A-CAA-CBA
29	Y	604	CLA	C1A-C2A-CAA-CBA
29	Y	608	CLA	C1A-C2A-CAA-CBA
29	b	613	CLA	C1A-C2A-CAA-CBA
29	n	602	CLA	C1A-C2A-CAA-CBA
29	y	604	CLA	C1A-C2A-CAA-CBA
45	R	607	CHL	C1A-C2A-CAA-CBA
45	y	605	CHL	C1A-C2A-CAA-CBA
29	n	603	CLA	C5-C6-C7-C8
29	r	603	CLA	C10-C11-C12-C13
39	l	101	LHG	C24-C25-C26-C27
29	N	611	CLA	C2-C1-O2A-CGA
29	S	603	CLA	C2-C1-O2A-CGA
29	S	614	CLA	C2-C1-O2A-CGA
29	b	611	CLA	C2-C1-O2A-CGA
29	c	503	CLA	C2-C1-O2A-CGA
29	n	614	CLA	C2-C1-O2A-CGA
29	g	603	CLA	C2-C1-O2A-CGA
39	C	525	LHG	C28-C29-C30-C31
49	S	625	LPX	C12-C13-C14-C15
31	c	515	BCR	C9-C10-C11-C12

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Mol	Chain	Res	Type	Atoms
46	R	620	LUT	C29-C30-C31-C32
39	D	409	LHG	C4-O6-P-O3
39	d	410	LHG	C4-O6-P-O3
49	S	625	LPX	C3-O1-P1-O2
32	C	526	SQD	C13-C14-C15-C16
39	D	410	LHG	C2-C3-O3-P
39	d	410	LHG	C2-C3-O3-P
50	s	626	3PH	C2-C1-O11-P
45	G	609	CHL	C2-C3-C5-C6
32	b	621	SQD	C10-C11-C12-C13
39	n	624	LHG	C11-C10-C9-C8
39	s	624	LHG	C33-C34-C35-C36
29	B	610	CLA	O1A-CGA-O2A-C1
39	D	408	LHG	C3-O3-P-O5
39	D	410	LHG	C3-O3-P-O4
39	L	101	LHG	C3-O3-P-O5
39	N	624	LHG	C4-O6-P-O4
39	S	624	LHG	C4-O6-P-O5
39	c	625	LHG	C3-O3-P-O4
39	d	409	LHG	C3-O3-P-O5
39	d	410	LHG	C4-O6-P-O4
39	l	101	LHG	C3-O3-P-O5
39	n	624	LHG	C4-O6-P-O4
39	s	624	LHG	C4-O6-P-O5
49	S	625	LPX	C1-O2-P1-O4
49	s	625	LPX	C1-O2-P1-O4
29	N	604	CLA	C16-C17-C18-C19
29	b	611	CLA	C16-C17-C18-C20
29	b	615	CLA	C16-C17-C18-C19
29	r	603	CLA	C11-C12-C13-C15
38	c	523	DGD	C7B-C8B-C9B-CAB
30	A	408	PHO	CBA-CGA-O2A-C1
39	c	625	LHG	O6-C4-C5-C6
39	G	630	LHG	C35-C36-C37-C38
33	a	413	LMG	C28-C29-C30-C31
29	r	608	CLA	C3-C5-C6-C7
33	J	101	LMG	C11-C12-C13-C14
33	A	413	LMG	C34-C35-C36-C37
29	C	504	CLA	CAD-CBD-CGD-O1D
29	G	604	CLA	CAD-CBD-CGD-O1D
29	n	612	CLA	CAD-CBD-CGD-O1D
29	g	604	CLA	CAD-CBD-CGD-O1D

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Mol	Chain	Res	Type	Atoms
45	N	601	CHL	CAD-CBD-CGD-O1D
45	n	601	CHL	CAD-CBD-CGD-O1D
45	s	601	CHL	CAD-CBD-CGD-O1D
49	S	625	LPX	C2-C1-O2-P1
49	s	625	LPX	C2-C1-O2-P1
38	C	518	DGD	C2B-C3B-C4B-C5B
45	N	605	CHL	C3-C5-C6-C7
29	R	608	CLA	C8-C10-C11-C12
39	g	624	LHG	C35-C36-C37-C38
38	c	523	DGD	C4D-C5D-C6D-O5D
29	B	604	CLA	C11-C12-C13-C15
29	B	615	CLA	C11-C12-C13-C15
29	B	617	CLA	C11-C10-C8-C7
29	C	510	CLA	C6-C7-C8-C10
29	C	510	CLA	C11-C10-C8-C7
29	G	602	CLA	C12-C13-C15-C16
29	G	603	CLA	C11-C10-C8-C7
29	G	613	CLA	C11-C10-C8-C7
29	R	608	CLA	C6-C7-C8-C10
29	S	603	CLA	C11-C10-C8-C7
29	Y	611	CLA	C12-C13-C15-C16
29	a	406	CLA	C11-C10-C8-C7
29	b	603	CLA	C6-C7-C8-C10
29	b	605	CLA	C11-C12-C13-C15
29	b	613	CLA	C11-C12-C13-C15
29	b	615	CLA	C11-C12-C13-C15
29	c	502	CLA	C6-C7-C8-C10
29	c	507	CLA	C11-C10-C8-C7
29	c	512	CLA	C6-C7-C8-C10
29	c	512	CLA	C11-C10-C8-C7
29	d	402	CLA	C11-C12-C13-C15
29	d	403	CLA	C11-C12-C13-C15
29	g	602	CLA	C12-C13-C15-C16
29	s	609	CLA	C6-C7-C8-C10
29	y	610	CLA	C11-C10-C8-C7
29	y	614	CLA	C11-C10-C8-C7
30	A	409	PHO	C6-C7-C8-C10
30	a	409	PHO	C6-C7-C8-C10
39	C	525	LHG	O6-C4-C5-O7
39	G	630	LHG	O6-C4-C5-O7
45	G	601	CHL	C11-C12-C13-C15
45	Y	606	CHL	C11-C10-C8-C7

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Mol	Chain	Res	Type	Atoms
45	n	606	CHL	C11-C12-C13-C15
46	R	620	LUT	C25-C26-C27-C28
46	r	620	LUT	C25-C26-C27-C28
50	S	626	3PH	O11-C1-C2-O21
50	s	626	3PH	O11-C1-C2-O21
36	B	625	DGA	CA5-CA6-CA7-CA8
36	B	625	DGA	CDB-CEB-CFB-CGB
45	g	601	CHL	C10-C11-C12-C13
36	c	524	DGA	CAB-CBB-CCB-CDB
38	c	519	DGD	C5A-C6A-C7A-C8A
29	g	613	CLA	C8-C10-C11-C12
33	d	411	LMG	C28-C29-C30-C31
39	d	410	LHG	O2-C2-C3-O3
39	c	625	LHG	C25-C26-C27-C28
29	B	613	CLA	C2A-CAA-CBA-CGA
29	C	501	CLA	C2A-CAA-CBA-CGA
29	b	607	CLA	C2A-CAA-CBA-CGA
45	n	607	CHL	CAA-CBA-CGA-O2A
32	A	412	SQD	C44-C45-C46-O48
33	A	413	LMG	O1-C7-C8-C9
39	Y	624	LHG	C34-C35-C36-C37
32	A	412	SQD	O47-C45-C46-O48
38	c	519	DGD	O2G-C2G-C3G-O3G
39	D	410	LHG	O7-C5-C6-O8
39	L	101	LHG	O7-C5-C6-O8
39	d	408	LHG	O7-C5-C6-O8
39	s	624	LHG	O7-C5-C6-O8
39	y	624	LHG	O7-C5-C6-O8
40	C	527	LMK	O1-C7-C8-O7
33	h	102	LMG	C15-C16-C17-C18
40	c	627	LMK	C8-C9-O8-C28
39	g	624	LHG	O8-C23-C24-C25
29	c	513	CLA	C8-C10-C11-C12
29	G	602	CLA	C3-C5-C6-C7
29	n	602	CLA	C8-C10-C11-C12
36	b	623	DGA	CAA-CBA-CCA-CDA
29	Y	602	CLA	C10-C11-C12-C13
29	s	610	CLA	C8-C10-C11-C12
29	B	603	CLA	C6-C7-C8-C9
29	B	605	CLA	C11-C12-C13-C14
29	C	501	CLA	C11-C12-C13-C14
29	C	513	CLA	C6-C7-C8-C9

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Mol	Chain	Res	Type	Atoms
29	N	602	CLA	C11-C10-C8-C9
29	N	603	CLA	C11-C12-C13-C14
29	Y	602	CLA	C6-C7-C8-C9
29	Y	614	CLA	C11-C10-C8-C9
29	b	606	CLA	C11-C12-C13-C14
29	b	612	CLA	C11-C12-C13-C14
29	c	504	CLA	C6-C7-C8-C9
29	c	509	CLA	C11-C12-C13-C14
29	c	512	CLA	C11-C10-C8-C9
29	d	402	CLA	C11-C12-C13-C14
29	n	604	CLA	C6-C7-C8-C9
29	n	613	CLA	C14-C13-C15-C16
29	s	603	CLA	C11-C12-C13-C14
45	N	607	CHL	C11-C10-C8-C9
45	N	609	CHL	C11-C10-C8-C9
45	Y	609	CHL	C11-C12-C13-C14
45	n	606	CHL	C11-C10-C8-C9
45	y	601	CHL	C11-C10-C8-C9
29	G	610	CLA	C3-C5-C6-C7
29	Y	610	CLA	C10-C11-C12-C13
42	D	405	PL9	C14-C16-C17-C18
42	d	405	PL9	C39-C41-C42-C43
30	A	408	PHO	O1A-CGA-O2A-C1
39	N	624	LHG	O8-C23-C24-C25
31	D	404	BCR	C18-C19-C20-C21
31	d	404	BCR	C18-C19-C20-C21
31	B	619	BCR	C11-C12-C13-C35
46	G	620	LUT	C31-C32-C33-C40
29	c	508	CLA	C8-C10-C11-C12
50	s	626	3PH	C31-C32-C33-C34
36	b	623	DGA	CA7-CA8-CA9-CAA
39	d	409	LHG	C34-C35-C36-C37
36	B	625	DGA	CCA-CDA-CEA-CFA
42	D	405	PL9	C45-C44-C46-C47
29	B	617	CLA	C10-C11-C12-C13
29	C	504	CLA	C10-C11-C12-C13
29	Y	611	CLA	C8-C10-C11-C12
39	d	409	LHG	C17-C18-C19-C20
29	b	617	CLA	C15-C16-C17-C18
29	N	611	CLA	C1-C2-C3-C4
29	G	604	CLA	C1-C2-C3-C4
29	G	614	CLA	C1-C2-C3-C4

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Mol	Chain	Res	Type	Atoms
29	n	611	CLA	C1-C2-C3-C4
29	g	604	CLA	C1-C2-C3-C4
29	g	614	CLA	C1-C2-C3-C4
45	N	608	CHL	C1-C2-C3-C4
45	G	606	CHL	C1-C2-C3-C4
45	G	607	CHL	C1-C2-C3-C4
45	n	608	CHL	C1-C2-C3-C4
45	g	606	CHL	C1-C2-C3-C4
45	g	607	CHL	C1-C2-C3-C4
45	r	607	CHL	C1-C2-C3-C4
29	C	505	CLA	C3-C5-C6-C7
38	C	518	DGD	CAB-CBB-CCB-CDB
39	S	624	LHG	C13-C14-C15-C16
29	A	405	CLA	C5-C6-C7-C8
39	N	624	LHG	C6-C5-O7-C7
50	s	626	3PH	O11-C1-C2-C3
29	B	607	CLA	C2A-CAA-CBA-CGA
29	N	613	CLA	C2A-CAA-CBA-CGA
29	S	610	CLA	C2A-CAA-CBA-CGA
29	c	501	CLA	C2A-CAA-CBA-CGA
29	y	603	CLA	C2A-CAA-CBA-CGA
45	n	606	CHL	C2A-CAA-CBA-CGA
29	b	603	CLA	C2-C1-O2A-CGA
29	d	403	CLA	C2-C1-O2A-CGA
38	C	519	DGD	C6B-C7B-C8B-C9B
33	C	521	LMG	C10-C11-C12-C13
29	R	609	CLA	CAA-CBA-CGA-O2A
29	G	603	CLA	C15-C16-C17-C18
39	d	410	LHG	C31-C32-C33-C34
39	L	101	LHG	C2-C3-O3-P
32	b	621	SQD	C14-C15-C16-C17
44	h	101	RRX	C15-C16-C17-C18
48	g	623	NEX	C13-C14-C15-C35
29	N	604	CLA	O1A-CGA-O2A-C1
39	d	409	LHG	O6-C4-C5-O7
39	L	101	LHG	C24-C25-C26-C27
30	a	408	PHO	C16-C17-C18-C19
45	R	607	CHL	O2A-C1-C2-C3
29	R	603	CLA	C4-C3-C5-C6
29	c	506	CLA	C4-C3-C5-C6
31	c	517	BCR	C5-C6-C7-C8
35	b	620	C7Z	C1-C6-C7-C8

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Mol	Chain	Res	Type	Atoms
46	N	620	LUT	C5-C6-C7-C8
29	G	610	CLA	C2-C3-C5-C6
45	N	607	CHL	CAA-CBA-CGA-O2A
29	b	610	CLA	C5-C6-C7-C8
39	D	408	LHG	C11-C10-C9-C8
29	B	608	CLA	C16-C17-C18-C20
29	b	608	CLA	C16-C17-C18-C19
29	r	603	CLA	C11-C12-C13-C14
39	l	101	LHG	C17-C18-C19-C20
29	B	617	CLA	C2A-CAA-CBA-CGA
33	c	521	LMG	O1-C7-C8-O7
38	c	523	DGD	O1G-C1G-C2G-O2G
29	Y	613	CLA	C10-C11-C12-C13
39	d	410	LHG	C11-C10-C9-C8
29	N	604	CLA	CBA-CGA-O2A-C1
39	N	624	LHG	C3-O3-P-O6
39	G	630	LHG	C3-O3-P-O6
39	S	624	LHG	C3-O3-P-O6
39	Y	624	LHG	C3-O3-P-O6
39	d	408	LHG	C3-O3-P-O6
39	d	409	LHG	C4-O6-P-O3
39	n	624	LHG	C3-O3-P-O6
39	g	624	LHG	C3-O3-P-O6
39	s	624	LHG	C3-O3-P-O6
39	y	624	LHG	C3-O3-P-O6
49	s	625	LPX	C3-O1-P1-O2
32	C	526	SQD	C12-C13-C14-C15
29	Y	613	CLA	C5-C6-C7-C8
30	a	409	PHO	CHA-CBD-CGD-O1D
30	a	409	PHO	CHA-CBD-CGD-O2D
36	B	625	DGA	CB4-CB5-CB6-CB7
33	H	102	LMG	C7-C8-C9-O8
50	i	101	3PH	C1-C2-C3-O31
39	C	525	LHG	C33-C34-C35-C36
39	N	624	LHG	C11-C10-C9-C8
29	B	613	CLA	C12-C13-C15-C16
29	B	617	CLA	C11-C12-C13-C15
29	N	603	CLA	C11-C12-C13-C15
29	b	606	CLA	C11-C12-C13-C15
29	r	603	CLA	C2-C3-C5-C6
29	S	614	CLA	CAA-CBA-CGA-O2A
32	b	621	SQD	C25-C26-C27-C28

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Mol	Chain	Res	Type	Atoms
29	C	510	CLA	C6-C7-C8-C9
29	D	402	CLA	C11-C10-C8-C9
29	G	603	CLA	C11-C10-C8-C9
29	S	603	CLA	C6-C7-C8-C9
29	Y	604	CLA	C11-C10-C8-C9
29	Y	611	CLA	C14-C13-C15-C16
29	b	602	CLA	C11-C12-C13-C14
29	b	612	CLA	C14-C13-C15-C16
29	b	615	CLA	C11-C12-C13-C14
29	b	617	CLA	C11-C12-C13-C14
29	c	511	CLA	C11-C12-C13-C14
29	c	513	CLA	C6-C7-C8-C9
29	d	403	CLA	C11-C12-C13-C14
29	s	610	CLA	C6-C7-C8-C9
29	y	611	CLA	C14-C13-C15-C16
29	y	612	CLA	C11-C12-C13-C14
45	N	605	CHL	C14-C13-C15-C16
31	C	515	BCR	C9-C10-C11-C12
31	c	516	BCR	C9-C10-C11-C12
29	C	513	CLA	C16-C17-C18-C19
29	b	613	CLA	C16-C17-C18-C19
39	n	624	LHG	C24-C23-O8-C6
29	D	403	CLA	C10-C11-C12-C13
29	C	513	CLA	C8-C10-C11-C12
33	A	413	LMG	C30-C31-C32-C33
46	n	620	LUT	C31-C32-C33-C40
29	n	602	CLA	C16-C17-C18-C20
30	a	408	PHO	C16-C17-C18-C20
39	d	409	LHG	C24-C25-C26-C27
29	b	606	CLA	C8-C10-C11-C12
33	d	411	LMG	C12-C13-C14-C15
50	s	626	3PH	C3D-C3E-C3F-C3G
39	Y	624	LHG	C29-C30-C31-C32
39	d	408	LHG	C11-C10-C9-C8
29	N	611	CLA	O2A-C1-C2-C3
29	s	604	CLA	C4-C3-C5-C6
33	c	521	LMG	C28-C29-C30-C31
29	n	611	CLA	O1A-CGA-O2A-C1
29	N	604	CLA	C16-C17-C18-C20
29	b	611	CLA	C16-C17-C18-C19
29	n	611	CLA	CBA-CGA-O2A-C1
30	a	408	PHO	CBA-CGA-O2A-C1

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Mol	Chain	Res	Type	Atoms
29	y	610	CLA	C15-C16-C17-C18
50	s	626	3PH	C22-C23-C24-C25
29	B	611	CLA	C2A-CAA-CBA-CGA
29	C	510	CLA	C2A-CAA-CBA-CGA
31	B	619	BCR	C13-C14-C15-C16
31	d	404	BCR	C9-C10-C11-C12
46	n	620	LUT	C29-C30-C31-C32
46	r	620	LUT	C9-C10-C11-C12
46	r	620	LUT	C29-C30-C31-C32
46	y	620	LUT	C29-C30-C31-C32
39	s	624	LHG	O6-C4-C5-C6
42	D	405	PL9	C39-C41-C42-C43
29	y	602	CLA	C10-C11-C12-C13
39	y	624	LHG	C34-C35-C36-C37
47	R	621	XAT	C30-C31-C32-C33
48	y	623	NEX	C30-C31-C32-C33
33	C	521	LMG	C28-C29-C30-C31
29	Y	611	CLA	C16-C17-C18-C19
29	G	602	CLA	C4-C3-C5-C6
29	b	607	CLA	C4-C3-C5-C6
29	n	602	CLA	C13-C15-C16-C17
39	n	624	LHG	C33-C34-C35-C36
39	n	624	LHG	O10-C23-O8-C6
36	b	623	DGA	CB2-CB3-CB4-CB5
29	B	606	CLA	C8-C10-C11-C12
29	C	504	CLA	C8-C10-C11-C12
29	S	610	CLA	C2-C1-O2A-CGA
29	Y	602	CLA	C2-C1-O2A-CGA
29	y	602	CLA	C2-C1-O2A-CGA
30	A	409	PHO	C2-C1-O2A-CGA
39	l	101	LHG	C26-C27-C28-C29
38	c	520	DGD	C4B-C5B-C6B-C7B
29	C	502	CLA	C2A-CAA-CBA-CGA
29	C	512	CLA	C2A-CAA-CBA-CGA
29	G	610	CLA	C2A-CAA-CBA-CGA
29	b	611	CLA	C2A-CAA-CBA-CGA
29	g	602	CLA	C2A-CAA-CBA-CGA
29	r	613	CLA	C2A-CAA-CBA-CGA
33	h	102	LMG	O7-C8-C9-O8
39	S	624	LHG	C2-C3-O3-P
39	l	101	LHG	C2-C3-O3-P
33	D	411	LMG	C32-C33-C34-C35

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Mol	Chain	Res	Type	Atoms
29	C	511	CLA	C3A-C2A-CAA-CBA
29	S	605	CLA	C3A-C2A-CAA-CBA
45	y	601	CHL	C3A-C2A-CAA-CBA
33	h	102	LMG	C12-C13-C14-C15
39	D	409	LHG	C17-C18-C19-C20
46	Y	620	LUT	C33-C34-C35-C15
29	C	510	CLA	C4-C3-C5-C6
39	C	525	LHG	C9-C10-C11-C12
50	i	101	3PH	C27-C28-C29-C2A
29	c	510	CLA	C5-C6-C7-C8
29	B	616	CLA	C6-C7-C8-C9
29	C	510	CLA	C11-C10-C8-C9
29	G	602	CLA	C11-C10-C8-C9
29	b	609	CLA	C11-C12-C13-C14
29	b	617	CLA	C6-C7-C8-C9
29	c	511	CLA	C6-C7-C8-C9
29	n	603	CLA	C11-C12-C13-C14
29	y	610	CLA	C11-C10-C8-C9
29	y	614	CLA	C11-C10-C8-C9
45	N	605	CHL	C11-C12-C13-C14
45	N	606	CHL	C14-C13-C15-C16
45	n	609	CHL	C14-C13-C15-C16
45	g	601	CHL	C11-C10-C8-C9
45	y	607	CHL	C11-C12-C13-C14
29	c	503	CLA	C16-C17-C18-C19
45	s	601	CHL	CAA-CBA-CGA-O2A
29	c	509	CLA	C8-C10-C11-C12
39	C	525	LHG	C25-C26-C27-C28
39	d	410	LHG	C34-C35-C36-C37
31	D	404	BCR	C20-C21-C22-C37
31	d	404	BCR	C20-C21-C22-C37
38	c	519	DGD	C1G-C2G-C3G-O3G
39	y	624	LHG	C4-C5-C6-O8
48	S	622	NEX	C39-C29-C30-C31
48	s	623	NEX	C39-C29-C30-C31
33	h	102	LMG	O9-C10-O7-C8
32	c	626	SQD	C9-C10-C11-C12
39	C	525	LHG	C12-C13-C14-C15
45	s	606	CHL	C2A-CAA-CBA-CGA
29	C	511	CLA	C2A-CAA-CBA-CGA
29	b	615	CLA	C16-C17-C18-C20
38	C	518	DGD	O6E-C1E-O5D-C6D

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Mol	Chain	Res	Type	Atoms
29	r	609	CLA	C8-C10-C11-C12
45	S	601	CHL	CAA-CBA-CGA-O1A
47	G	622	XAT	C14-C15-C35-C34
45	N	607	CHL	C8-C10-C11-C12
33	H	102	LMG	C12-C13-C14-C15
51	Y	625	SPH	C11-C10-C9-C8
39	n	624	LHG	C6-C5-O7-C7
51	Y	625	SPH	O3-C3-C4-C5
29	b	605	CLA	C15-C16-C17-C18
45	y	609	CHL	C4-C3-C5-C6
29	D	403	CLA	C1A-C2A-CAA-CBA
29	Y	614	CLA	C1A-C2A-CAA-CBA
29	d	403	CLA	C1A-C2A-CAA-CBA
29	g	614	CLA	C1A-C2A-CAA-CBA
29	y	602	CLA	C1A-C2A-CAA-CBA
45	N	609	CHL	C1A-C2A-CAA-CBA
45	Y	601	CHL	C1A-C2A-CAA-CBA
29	A	406	CLA	C11-C10-C8-C7
29	C	510	CLA	C2-C3-C5-C6
29	R	609	CLA	C6-C7-C8-C10
29	b	608	CLA	C12-C13-C15-C16
29	c	503	CLA	C6-C7-C8-C10
29	c	510	CLA	C11-C10-C8-C7
29	d	402	CLA	C11-C10-C8-C7
29	y	604	CLA	C11-C10-C8-C7
45	N	607	CHL	C11-C12-C13-C15
29	C	508	CLA	C8-C10-C11-C12
29	G	613	CLA	C13-C15-C16-C17
29	Y	603	CLA	C10-C11-C12-C13
29	c	501	CLA	C3-C5-C6-C7
31	C	514	BCR	C19-C20-C21-C22
31	C	515	BCR	C19-C20-C21-C22
39	L	101	LHG	C31-C32-C33-C34
30	a	408	PHO	O1A-CGA-O2A-C1
29	g	603	CLA	C3-C5-C6-C7
29	A	405	CLA	C2A-CAA-CBA-CGA
29	G	602	CLA	C2A-CAA-CBA-CGA
29	g	610	CLA	C2A-CAA-CBA-CGA
29	c	509	CLA	C13-C15-C16-C17
29	c	512	CLA	C5-C6-C7-C8
38	C	523	DGD	C6A-C7A-C8A-C9A
39	D	410	LHG	C34-C35-C36-C37

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Mol	Chain	Res	Type	Atoms
39	G	630	LHG	C33-C34-C35-C36
45	N	608	CHL	O2A-C1-C2-C3
45	n	608	CHL	O2A-C1-C2-C3
29	N	613	CLA	C10-C11-C12-C13
38	c	519	DGD	C6A-C7A-C8A-C9A
36	b	623	DGA	CA5-CA6-CA7-CA8
39	G	630	LHG	O8-C23-C24-C25
29	n	602	CLA	C4-C3-C5-C6
33	d	411	LMG	C32-C33-C34-C35
29	s	603	CLA	C8-C10-C11-C12
38	C	520	DGD	C1A-C2A-C3A-C4A
32	C	526	SQD	C31-C32-C33-C34
43	f	101	HEM	C3D-CAD-CBD-CGD
45	s	601	CHL	CAA-CBA-CGA-O1A
29	g	603	CLA	C5-C6-C7-C8
39	G	630	LHG	C9-C10-C11-C12
39	l	101	LHG	C31-C32-C33-C34
29	a	406	CLA	C3-C5-C6-C7
45	g	601	CHL	C3-C5-C6-C7
31	D	404	BCR	C20-C21-C22-C23
31	d	404	BCR	C20-C21-C22-C23
48	S	622	NEX	C28-C29-C30-C31
48	s	623	NEX	C28-C29-C30-C31
29	s	602	CLA	C10-C11-C12-C13
32	a	412	SQD	C27-C28-C29-C30
38	C	519	DGD	O2G-C2G-C3G-O3G
39	D	408	LHG	O7-C5-C6-O8
29	D	402	CLA	CBA-CGA-O2A-C1
32	a	412	SQD	C32-C33-C34-C35
39	G	630	LHG	C16-C17-C18-C19
50	s	626	3PH	C33-C34-C35-C36
31	a	411	BCR	C19-C20-C21-C22
31	c	514	BCR	C19-C20-C21-C22
46	Y	620	LUT	C29-C30-C31-C32
33	h	102	LMG	C19-C20-C21-C22
29	D	402	CLA	O1A-CGA-O2A-C1
33	C	521	LMG	O10-C28-O8-C9
39	N	624	LHG	C1-C2-C3-O3
29	a	406	CLA	C15-C16-C17-C18
45	N	609	CHL	C8-C10-C11-C12
29	R	602	CLA	C3-C5-C6-C7
29	C	506	CLA	C4-C3-C5-C6

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Mol	Chain	Res	Type	Atoms
29	N	613	CLA	CAA-CBA-CGA-O1A
29	N	602	CLA	C2-C1-O2A-CGA
29	b	615	CLA	C2-C1-O2A-CGA
29	c	501	CLA	C2-C1-O2A-CGA
45	G	609	CHL	C2-C1-O2A-CGA
45	g	607	CHL	C2-C1-O2A-CGA
29	c	506	CLA	C2-C3-C5-C6
29	s	604	CLA	C2-C3-C5-C6
39	D	409	LHG	C16-C17-C18-C19
39	d	409	LHG	C18-C19-C20-C21
45	S	601	CHL	CAA-CBA-CGA-O2A
29	C	512	CLA	C11-C12-C13-C14
29	c	504	CLA	C14-C13-C15-C16
29	r	610	CLA	C10-C11-C12-C13
45	g	608	CHL	C2A-CAA-CBA-CGA
33	c	521	LMG	O10-C28-O8-C9
29	A	406	CLA	C3-C5-C6-C7
29	S	614	CLA	C5-C6-C7-C8
51	y	625	SPH	O1-C1-C2-C3
29	d	402	CLA	CAA-CBA-CGA-O2A
33	H	102	LMG	C19-C20-C21-C22
39	N	624	LHG	C33-C34-C35-C36
45	g	605	CHL	C2-C1-O2A-CGA
29	B	602	CLA	C16-C17-C18-C19
38	c	520	DGD	C9B-CAB-CBB-CCB
38	c	523	DGD	CDA-CEA-CFA-CGA
31	B	618	BCR	C23-C24-C25-C30
31	B	619	BCR	C23-C24-C25-C30
31	b	618	BCR	C23-C24-C25-C30
31	c	514	BCR	C23-C24-C25-C30
31	c	517	BCR	C1-C6-C7-C8
35	B	620	C7Z	C1-C6-C7-C8
46	S	621	LUT	C1-C6-C7-C8
46	y	620	LUT	C1-C6-C7-C8
46	y	620	LUT	C5-C6-C7-C8
29	Y	612	CLA	C5-C6-C7-C8
29	s	611	CLA	CAA-CBA-CGA-O2A
39	C	525	LHG	O8-C23-C24-C25
32	B	621	SQD	O6-C44-C45-C46
32	b	621	SQD	O6-C44-C45-C46
33	h	102	LMG	C7-C8-C9-O8
31	b	619	BCR	C15-C16-C17-C18

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Mol	Chain	Res	Type	Atoms
46	G	620	LUT	C29-C30-C31-C32
47	r	622	XAT	C33-C34-C35-C15
47	y	622	XAT	C29-C30-C31-C32
29	B	617	CLA	C4-C3-C5-C6
29	N	602	CLA	C4-C3-C5-C6
29	c	511	CLA	C4-C3-C5-C6
29	y	610	CLA	C4-C3-C5-C6
45	n	609	CHL	C4-C3-C5-C6
31	B	619	BCR	C11-C12-C13-C14
46	G	620	LUT	C31-C32-C33-C34
38	C	519	DGD	C5A-C6A-C7A-C8A
29	R	603	CLA	C2-C3-C5-C6
48	G	623	NEX	C14-C15-C35-C34
39	L	101	LHG	C26-C27-C28-C29
29	D	402	CLA	C15-C16-C17-C18
29	g	613	CLA	C5-C6-C7-C8
29	b	613	CLA	C2A-CAA-CBA-CGA
33	C	521	LMG	C29-C28-O8-C9
29	g	610	CLA	C3-C5-C6-C7
29	n	612	CLA	CAA-CBA-CGA-O2A
29	B	604	CLA	C15-C16-C17-C18
29	G	610	CLA	C15-C16-C17-C18
39	D	410	LHG	C29-C30-C31-C32
29	C	512	CLA	C11-C12-C13-C15
29	n	603	CLA	C11-C12-C13-C15
45	n	605	CHL	C11-C12-C13-C15
45	g	609	CHL	C11-C12-C13-C15
45	y	609	CHL	C2-C3-C5-C6
33	c	521	LMG	C29-C28-O8-C9
33	A	413	LMG	C33-C34-C35-C36
33	h	102	LMG	C11-C12-C13-C14
36	b	623	DGA	OG2-CG2-CG3-OXT
36	c	524	DGA	OG2-CG2-CG3-OXT
31	A	411	BCR	C19-C20-C21-C22
31	D	404	BCR	C13-C14-C15-C16
46	N	620	LUT	C29-C30-C31-C32
29	D	402	CLA	CAA-CBA-CGA-O2A
30	A	408	PHO	C16-C17-C18-C19
32	b	621	SQD	C18-C19-C20-C21
29	S	603	CLA	C10-C11-C12-C13
38	c	520	DGD	O1G-C1G-C2G-O2G
39	s	624	LHG	C35-C36-C37-C38

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Mol	Chain	Res	Type	Atoms
29	Y	603	CLA	C13-C15-C16-C17
45	g	606	CHL	O2A-C1-C2-C3
45	r	607	CHL	O2A-C1-C2-C3
29	b	605	CLA	CAA-CBA-CGA-O2A
29	y	613	CLA	CAA-CBA-CGA-O2A
39	d	410	LHG	O8-C23-C24-C25
45	r	607	CHL	CAA-CBA-CGA-O2A
33	d	411	LMG	C29-C30-C31-C32
39	D	408	LHG	C31-C32-C33-C34
29	y	613	CLA	C10-C11-C12-C13
29	B	608	CLA	C16-C17-C18-C19
36	c	524	DGA	CA2-CA1-OG1-CG1
38	C	520	DGD	O6D-C5D-C6D-O5D
29	S	604	CLA	C4-C3-C5-C6
29	b	609	CLA	C4-C3-C5-C6
29	b	617	CLA	C4-C3-C5-C6
29	c	505	CLA	C4-C3-C5-C6
29	s	611	CLA	C4-C3-C5-C6
42	d	405	PL9	C20-C19-C21-C22
45	Y	609	CHL	C4-C3-C5-C6
29	A	406	CLA	C15-C16-C17-C18
38	C	520	DGD	C9A-CAA-CBA-CCA
29	N	602	CLA	C2-C3-C5-C6
29	n	602	CLA	C2-C3-C5-C6
30	A	408	PHO	C2-C3-C5-C6
42	d	405	PL9	C18-C19-C21-C22
33	b	622	LMG	O7-C10-C11-C12
39	Y	624	LHG	C15-C16-C17-C18
29	B	604	CLA	C11-C12-C13-C14
29	R	608	CLA	C6-C7-C8-C9
29	S	603	CLA	C11-C10-C8-C9
29	a	406	CLA	C11-C10-C8-C9
29	b	613	CLA	C11-C12-C13-C14
29	c	506	CLA	C6-C7-C8-C9
29	c	512	CLA	C6-C7-C8-C9
29	n	604	CLA	C11-C10-C8-C9
45	y	609	CHL	C11-C10-C8-C9
51	Y	625	SPH	C2-C3-C4-C5
29	R	610	CLA	C3A-C2A-CAA-CBA
29	n	610	CLA	C3A-C2A-CAA-CBA
29	g	604	CLA	C3A-C2A-CAA-CBA
29	s	605	CLA	C3A-C2A-CAA-CBA

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Mol	Chain	Res	Type	Atoms
45	R	607	CHL	C3A-C2A-CAA-CBA
45	g	601	CHL	C3A-C2A-CAA-CBA
45	s	601	CHL	C3A-C2A-CAA-CBA
45	y	605	CHL	C3A-C2A-CAA-CBA
39	d	409	LHG	C16-C17-C18-C19
39	g	624	LHG	C16-C17-C18-C19
45	Y	607	CHL	C10-C11-C12-C13
29	b	610	CLA	O1A-CGA-O2A-C1
36	c	524	DGA	OA1-CA1-OG1-CG1
29	b	609	CLA	CAA-CBA-CGA-O2A
32	a	412	SQD	O47-C7-C8-C9
33	B	622	LMG	O7-C10-C11-C12
39	l	101	LHG	O7-C7-C8-C9
50	s	626	3PH	C36-C37-C38-C39
29	S	612	CLA	CAA-CBA-CGA-O2A
29	B	610	CLA	CAD-CBD-CGD-O2D
29	C	509	CLA	CAD-CBD-CGD-O2D
29	C	510	CLA	CAD-CBD-CGD-O2D
29	N	603	CLA	CAD-CBD-CGD-O2D
29	G	611	CLA	CAD-CBD-CGD-O2D
29	S	612	CLA	CAD-CBD-CGD-O2D
29	b	610	CLA	CAD-CBD-CGD-O2D
29	b	617	CLA	CAD-CBD-CGD-O2D
29	c	503	CLA	CAD-CBD-CGD-O2D
29	c	504	CLA	CAD-CBD-CGD-O2D
29	n	603	CLA	CAD-CBD-CGD-O2D
29	y	614	CLA	CAD-CBD-CGD-O2D
30	A	408	PHO	CAD-CBD-CGD-O2D
40	c	627	LMK	C1-C2-C3-N4
45	S	601	CHL	CAD-CBD-CGD-O2D
45	r	606	CHL	CAD-CBD-CGD-O2D
45	y	606	CHL	CAD-CBD-CGD-O2D
29	C	513	CLA	C16-C17-C18-C20
29	y	610	CLA	C8-C10-C11-C12
29	B	617	CLA	C2-C1-O2A-CGA
38	c	519	DGD	O6D-C5D-C6D-O5D
39	D	410	LHG	O8-C23-C24-C25
33	d	411	LMG	C15-C16-C17-C18
39	D	410	LHG	C31-C32-C33-C34
39	G	630	LHG	C34-C35-C36-C37
42	D	405	PL9	C20-C19-C21-C22
29	b	609	CLA	C16-C17-C18-C19

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Mol	Chain	Res	Type	Atoms
39	g	624	LHG	C33-C34-C35-C36
29	b	602	CLA	C2-C3-C5-C6
29	S	617	CLA	CAA-CBA-CGA-O2A
29	Y	613	CLA	CAA-CBA-CGA-O2A
33	c	521	LMG	O7-C10-C11-C12
32	C	526	SQD	C27-C28-C29-C30
38	C	520	DGD	C9B-CAB-CBB-CCB
46	n	620	LUT	C31-C32-C33-C34
38	C	519	DGD	O1G-C1G-C2G-C3G
38	C	523	DGD	O1G-C1G-C2G-C3G
39	c	625	LHG	C5-C4-O6-P
40	C	527	LMK	C11-C10-O7-C8
47	N	622	XAT	O24-C26-C27-C28
47	Y	622	XAT	O24-C26-C27-C28
29	n	612	CLA	CAA-CBA-CGA-O1A
39	Y	624	LHG	O6-C4-C5-O7
29	C	503	CLA	CAA-CBA-CGA-O2A
29	G	613	CLA	CAA-CBA-CGA-O2A
36	B	625	DGA	OG2-CB1-CB2-CB3
29	n	613	CLA	C16-C17-C18-C19
39	S	624	LHG	C35-C36-C37-C38
36	C	524	DGA	CA5-CA6-CA7-CA8
29	S	605	CLA	C2A-CAA-CBA-CGA
29	b	605	CLA	C13-C15-C16-C17
39	G	630	LHG	C32-C33-C34-C35
29	a	410	CLA	C3-C5-C6-C7
39	D	409	LHG	C27-C28-C29-C30
29	B	611	CLA	CHA-CBD-CGD-O1D
29	B	611	CLA	CHA-CBD-CGD-O2D
29	C	507	CLA	CHA-CBD-CGD-O2D
29	G	604	CLA	CHA-CBD-CGD-O1D
29	G	610	CLA	CHA-CBD-CGD-O1D
29	G	610	CLA	CHA-CBD-CGD-O2D
29	G	612	CLA	CHA-CBD-CGD-O1D
29	G	612	CLA	CHA-CBD-CGD-O2D
29	G	613	CLA	CHA-CBD-CGD-O1D
29	G	613	CLA	CHA-CBD-CGD-O2D
29	R	602	CLA	CHA-CBD-CGD-O1D
29	R	602	CLA	CHA-CBD-CGD-O2D
29	R	603	CLA	CHA-CBD-CGD-O1D
29	R	603	CLA	CHA-CBD-CGD-O2D
29	R	609	CLA	CHA-CBD-CGD-O1D

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Mol	Chain	Res	Type	Atoms
29	R	609	CLA	CHA-CBD-CGD-O2D
29	S	603	CLA	CHA-CBD-CGD-O1D
29	S	603	CLA	CHA-CBD-CGD-O2D
29	S	617	CLA	CHA-CBD-CGD-O1D
29	S	617	CLA	CHA-CBD-CGD-O2D
29	Y	602	CLA	CHA-CBD-CGD-O1D
29	Y	604	CLA	CHA-CBD-CGD-O2D
29	Y	608	CLA	CHA-CBD-CGD-O1D
29	Y	608	CLA	CHA-CBD-CGD-O2D
29	Y	610	CLA	CHA-CBD-CGD-O1D
29	Y	610	CLA	CHA-CBD-CGD-O2D
29	b	609	CLA	CHA-CBD-CGD-O1D
29	b	609	CLA	CHA-CBD-CGD-O2D
29	b	615	CLA	CHA-CBD-CGD-O1D
29	b	615	CLA	CHA-CBD-CGD-O2D
29	c	502	CLA	CHA-CBD-CGD-O1D
29	c	502	CLA	CHA-CBD-CGD-O2D
29	c	507	CLA	CHA-CBD-CGD-O1D
29	c	507	CLA	CHA-CBD-CGD-O2D
29	d	402	CLA	CHA-CBD-CGD-O2D
29	n	604	CLA	CHA-CBD-CGD-O1D
29	n	604	CLA	CHA-CBD-CGD-O2D
29	n	612	CLA	CHA-CBD-CGD-O1D
29	g	602	CLA	CHA-CBD-CGD-O2D
29	g	612	CLA	CHA-CBD-CGD-O1D
29	g	612	CLA	CHA-CBD-CGD-O2D
29	r	602	CLA	CHA-CBD-CGD-O1D
29	r	602	CLA	CHA-CBD-CGD-O2D
29	r	603	CLA	CHA-CBD-CGD-O2D
29	r	604	CLA	CHA-CBD-CGD-O1D
29	r	604	CLA	CHA-CBD-CGD-O2D
29	r	609	CLA	CHA-CBD-CGD-O1D
29	r	609	CLA	CHA-CBD-CGD-O2D
29	r	610	CLA	CHA-CBD-CGD-O1D
29	r	610	CLA	CHA-CBD-CGD-O2D
29	s	603	CLA	CHA-CBD-CGD-O2D
29	s	612	CLA	CHA-CBD-CGD-O1D
29	s	612	CLA	CHA-CBD-CGD-O2D
29	s	617	CLA	CHA-CBD-CGD-O1D
29	s	617	CLA	CHA-CBD-CGD-O2D
29	y	603	CLA	CHA-CBD-CGD-O1D
29	y	603	CLA	CHA-CBD-CGD-O2D

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Mol	Chain	Res	Type	Atoms
29	y	608	CLA	CHA-CBD-CGD-O1D
29	y	608	CLA	CHA-CBD-CGD-O2D
29	y	610	CLA	CHA-CBD-CGD-O1D
29	y	610	CLA	CHA-CBD-CGD-O2D
29	y	613	CLA	CHA-CBD-CGD-O1D
29	y	613	CLA	CHA-CBD-CGD-O2D
31	d	404	BCR	C13-C14-C15-C16
45	N	601	CHL	CHA-CBD-CGD-O2D
45	G	606	CHL	CHA-CBD-CGD-O1D
45	G	606	CHL	CHA-CBD-CGD-O2D
45	G	609	CHL	CHA-CBD-CGD-O1D
45	G	609	CHL	CHA-CBD-CGD-O2D
45	Y	609	CHL	CHA-CBD-CGD-O2D
45	n	601	CHL	CHA-CBD-CGD-O2D
45	g	609	CHL	CHA-CBD-CGD-O1D
45	g	609	CHL	CHA-CBD-CGD-O2D
45	r	607	CHL	CHA-CBD-CGD-O1D
45	r	607	CHL	CHA-CBD-CGD-O2D
45	s	601	CHL	CHA-CBD-CGD-O1D
45	y	609	CHL	CHA-CBD-CGD-O1D
45	y	609	CHL	CHA-CBD-CGD-O2D
29	s	612	CLA	CAA-CBA-CGA-O2A
29	B	608	CLA	CAA-CBA-CGA-O2A
29	g	613	CLA	CAA-CBA-CGA-O2A
29	c	505	CLA	C2-C3-C5-C6
45	n	609	CHL	C2-C3-C5-C6
29	s	604	CLA	C3-C5-C6-C7
39	Y	624	LHG	O6-C4-C5-C6
39	d	408	LHG	O6-C4-C5-C6
29	S	612	CLA	CAA-CBA-CGA-O1A
39	d	408	LHG	C31-C32-C33-C34
29	s	604	CLA	C6-C7-C8-C9
42	d	405	PL9	C2-C3-C7-C8
29	B	605	CLA	CAA-CBA-CGA-O2A
29	B	612	CLA	CAA-CBA-CGA-O2A
29	S	611	CLA	CAA-CBA-CGA-O2A
29	b	608	CLA	CAA-CBA-CGA-O2A
29	n	604	CLA	CAA-CBA-CGA-O2A
32	C	526	SQD	O47-C7-C8-C9
33	C	521	LMG	O7-C10-C11-C12
33	J	101	LMG	O7-C10-C11-C12
32	a	412	SQD	O6-C44-C45-O47

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Mol	Chain	Res	Type	Atoms
40	c	627	LMK	O7-C8-C9-O8
29	n	603	CLA	C15-C16-C17-C18
29	A	405	CLA	CAA-CBA-CGA-O2A
29	B	603	CLA	CAA-CBA-CGA-O2A
29	B	609	CLA	CAA-CBA-CGA-O2A
29	R	604	CLA	CAA-CBA-CGA-O2A
36	b	623	DGA	OG2-CB1-CB2-CB3
39	D	408	LHG	O7-C7-C8-C9
45	n	605	CHL	CAA-CBA-CGA-O2A
30	A	409	PHO	CHA-CBD-CGD-O1D
30	A	409	PHO	CHA-CBD-CGD-O2D
29	a	405	CLA	C8-C10-C11-C12
29	a	405	CLA	C15-C16-C17-C18
29	b	610	CLA	CBA-CGA-O2A-C1
32	C	526	SQD	O48-C23-C24-C25
36	C	524	DGA	OG1-CA1-CA2-CA3
32	a	412	SQD	C11-C12-C13-C14
39	g	624	LHG	C34-C35-C36-C37
39	s	624	LHG	C25-C26-C27-C28
29	r	610	CLA	C8-C10-C11-C12
29	G	602	CLA	C2-C3-C5-C6
29	S	610	CLA	C11-C12-C13-C15
29	S	611	CLA	C11-C12-C13-C15
29	Y	603	CLA	C11-C12-C13-C15
29	b	607	CLA	C2-C3-C5-C6
29	s	603	CLA	C12-C13-C15-C16
29	y	610	CLA	C6-C7-C8-C10
45	Y	609	CHL	C2-C3-C5-C6
29	Y	611	CLA	C16-C17-C18-C20
38	c	523	DGD	C6A-C7A-C8A-C9A
29	b	603	CLA	CAA-CBA-CGA-O2A
32	A	412	SQD	C10-C11-C12-C13
29	A	406	CLA	C11-C10-C8-C9
29	B	606	CLA	C11-C12-C13-C14
29	B	617	CLA	C11-C10-C8-C9
29	Y	610	CLA	C11-C10-C8-C9
29	b	608	CLA	C14-C13-C15-C16
29	c	503	CLA	C6-C7-C8-C9
31	B	619	BCR	C19-C20-C21-C22
31	b	618	BCR	C13-C14-C15-C16
31	c	516	BCR	C19-C20-C21-C22
39	D	409	LHG	C11-C10-C9-C8

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Mol	Chain	Res	Type	Atoms
50	S	626	3PH	C2E-C2F-C2G-C2H
29	R	612	CLA	CAA-CBA-CGA-O2A
29	S	603	CLA	CAA-CBA-CGA-O2A
32	a	412	SQD	C4-C5-C6-S
39	G	630	LHG	C19-C20-C21-C22
33	h	102	LMG	C11-C10-O7-C8
29	B	610	CLA	C2A-CAA-CBA-CGA
29	B	614	CLA	C2A-CAA-CBA-CGA
29	G	613	CLA	CAA-CBA-CGA-O1A
29	r	609	CLA	CAA-CBA-CGA-O1A
45	N	605	CHL	CAA-CBA-CGA-O2A
39	y	624	LHG	C14-C15-C16-C17
32	C	526	SQD	C9-C10-C11-C12
29	B	602	CLA	CAA-CBA-CGA-O2A
39	d	408	LHG	C10-C11-C12-C13
39	d	410	LHG	C27-C28-C29-C30
31	a	411	BCR	C7-C8-C9-C10
31	a	411	BCR	C11-C12-C13-C14
47	R	621	XAT	C7-C8-C9-C10
29	g	604	CLA	C1A-C2A-CAA-CBA
29	r	609	CLA	C1A-C2A-CAA-CBA
45	N	605	CHL	C1A-C2A-CAA-CBA
45	G	601	CHL	C1A-C2A-CAA-CBA
45	g	601	CHL	C1A-C2A-CAA-CBA
45	g	607	CHL	C1A-C2A-CAA-CBA
45	r	607	CHL	C1A-C2A-CAA-CBA
45	s	601	CHL	C1A-C2A-CAA-CBA
45	y	601	CHL	C1A-C2A-CAA-CBA
29	b	605	CLA	CAA-CBA-CGA-O1A
45	r	607	CHL	CAA-CBA-CGA-O1A
29	c	503	CLA	CAA-CBA-CGA-O2A
29	S	609	CLA	C8-C10-C11-C12
33	H	102	LMG	C29-C30-C31-C32
33	d	411	LMG	C19-C20-C21-C22
29	S	617	CLA	CAA-CBA-CGA-O1A
33	B	622	LMG	O9-C10-C11-C12
39	d	410	LHG	O10-C23-C24-C25
45	N	605	CHL	CAA-CBA-CGA-O1A
32	a	412	SQD	C12-C13-C14-C15
39	d	408	LHG	C12-C13-C14-C15
38	c	519	DGD	O1G-C1G-C2G-C3G
29	s	610	CLA	CAA-CBA-CGA-O2A

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Mol	Chain	Res	Type	Atoms
39	y	624	LHG	O8-C23-C24-C25
29	B	603	CLA	CAA-CBA-CGA-O1A
29	y	613	CLA	CAA-CBA-CGA-O1A
32	C	526	SQD	O49-C7-C8-C9
33	a	413	LMG	C29-C30-C31-C32
30	A	408	PHO	C4-C3-C5-C6
39	g	624	LHG	C10-C11-C12-C13
51	y	625	SPH	C12-C13-C14-C15
29	B	605	CLA	CAA-CBA-CGA-O1A
29	B	608	CLA	CAA-CBA-CGA-O1A
33	C	521	LMG	O9-C10-C11-C12
33	b	622	LMG	O9-C10-C11-C12
50	S	626	3PH	C29-C2A-C2B-C2C
29	C	511	CLA	C13-C15-C16-C17
29	R	608	CLA	C10-C11-C12-C13
39	N	624	LHG	C3-O3-P-O5
39	G	630	LHG	C3-O3-P-O5
39	Y	624	LHG	C3-O3-P-O5
39	d	408	LHG	C3-O3-P-O5
39	g	624	LHG	C3-O3-P-O5
39	y	624	LHG	C3-O3-P-O5
29	B	609	CLA	CAA-CBA-CGA-O1A
29	C	503	CLA	CAA-CBA-CGA-O1A
29	b	603	CLA	CAA-CBA-CGA-O1A
29	b	609	CLA	CAA-CBA-CGA-O1A
36	B	625	DGA	OB1-CB1-CB2-CB3
29	C	508	CLA	CAA-CBA-CGA-O2A
29	s	612	CLA	CAA-CBA-CGA-O1A
46	S	621	LUT	C5-C6-C7-C8
33	j	101	LMG	C30-C31-C32-C33
29	B	616	CLA	C10-C11-C12-C13
29	B	612	CLA	CAA-CBA-CGA-O1A
29	b	608	CLA	CAA-CBA-CGA-O1A
33	c	521	LMG	O9-C10-C11-C12
29	c	503	CLA	C8-C10-C11-C12
39	D	409	LHG	C18-C19-C20-C21
45	s	601	CHL	C2A-CAA-CBA-CGA
29	g	613	CLA	CAA-CBA-CGA-O1A
39	D	410	LHG	O10-C23-C24-C25
39	L	101	LHG	C19-C20-C21-C22
39	Y	624	LHG	C23-C24-C25-C26
39	y	624	LHG	C23-C24-C25-C26

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Mol	Chain	Res	Type	Atoms
29	Y	613	CLA	CAA-CBA-CGA-O1A
33	J	101	LMG	O9-C10-C11-C12
29	b	610	CLA	C4-C3-C5-C6
30	a	409	PHO	C4-C3-C5-C6
32	a	412	SQD	C10-C11-C12-C13
29	B	614	CLA	CAD-CBD-CGD-O1D
29	C	511	CLA	CAD-CBD-CGD-O1D
29	a	410	CLA	CAD-CBD-CGD-O1D
29	b	613	CLA	CAD-CBD-CGD-O1D
29	c	502	CLA	CAD-CBD-CGD-O1D
29	c	510	CLA	CAD-CBD-CGD-O1D
29	d	402	CLA	CAD-CBD-CGD-O1D
29	r	603	CLA	CAD-CBD-CGD-O1D
29	r	613	CLA	CAD-CBD-CGD-O1D
29	s	603	CLA	CAD-CBD-CGD-O1D
29	y	610	CLA	CAD-CBD-CGD-O1D
32	B	621	SQD	O5-C5-C6-S
32	b	621	SQD	O5-C5-C6-S
45	G	609	CHL	CAD-CBD-CGD-O1D
29	R	604	CLA	CAA-CBA-CGA-O1A
29	n	604	CLA	CAA-CBA-CGA-O1A
36	b	623	DGA	OB1-CB1-CB2-CB3
39	D	408	LHG	O9-C7-C8-C9
29	S	605	CLA	CAA-CBA-CGA-O2A
29	g	614	CLA	CAA-CBA-CGA-O2A
29	B	612	CLA	C11-C12-C13-C14
29	C	503	CLA	C6-C7-C8-C9
29	C	504	CLA	C6-C7-C8-C9
29	S	611	CLA	C11-C12-C13-C14
29	Y	603	CLA	C11-C12-C13-C14
29	c	510	CLA	C11-C10-C8-C9
29	d	402	CLA	C11-C10-C8-C9
29	n	613	CLA	C11-C12-C13-C14
29	y	604	CLA	C11-C10-C8-C9
29	y	610	CLA	C6-C7-C8-C9
29	y	613	CLA	C6-C7-C8-C9
45	n	606	CHL	C11-C12-C13-C14
45	n	609	CHL	C11-C10-C8-C9
45	g	609	CHL	C14-C13-C15-C16
32	B	621	SQD	C14-C15-C16-C17
37	b	625	GOL	O2-C2-C3-O3
33	a	413	LMG	C11-C12-C13-C14

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Mol	Chain	Res	Type	Atoms
33	H	102	LMG	O9-C10-C11-C12
33	h	102	LMG	O9-C10-C11-C12
36	C	524	DGA	OA1-CA1-CA2-CA3
29	C	505	CLA	CAA-CBA-CGA-O2A
29	R	610	CLA	CAA-CBA-CGA-O2A
29	r	604	CLA	CAA-CBA-CGA-O2A
29	s	603	CLA	CAA-CBA-CGA-O2A
45	y	606	CHL	CAA-CBA-CGA-O2A
39	s	624	LHG	C10-C11-C12-C13
29	s	610	CLA	CAA-CBA-CGA-O1A
45	n	605	CHL	CAA-CBA-CGA-O1A
29	C	513	CLA	C15-C16-C17-C18
29	Y	602	CLA	C13-C15-C16-C17
29	N	604	CLA	CAA-CBA-CGA-O2A
29	G	614	CLA	CAA-CBA-CGA-O2A
33	D	411	LMG	O7-C10-C11-C12
33	d	411	LMG	O7-C10-C11-C12
33	j	101	LMG	O7-C10-C11-C12
38	c	518	DGD	O1G-C1A-C2A-C3A
29	N	603	CLA	C10-C11-C12-C13
29	b	612	CLA	C5-C6-C7-C8
29	g	613	CLA	C3-C5-C6-C7
29	A	405	CLA	CAA-CBA-CGA-O1A
36	B	625	DGA	CA7-CA8-CA9-CAA
29	R	610	CLA	C4-C3-C5-C6
29	Y	610	CLA	C4-C3-C5-C6
29	b	602	CLA	C4-C3-C5-C6
29	b	616	CLA	C4-C3-C5-C6
29	g	610	CLA	C4-C3-C5-C6
30	A	409	PHO	C4-C3-C5-C6
29	B	616	CLA	C15-C16-C17-C18
36	c	524	DGA	CA2-CA3-CA4-CA5
29	B	606	CLA	C11-C12-C13-C15
29	C	501	CLA	C11-C12-C13-C15
29	C	503	CLA	C6-C7-C8-C10
29	Y	610	CLA	C6-C7-C8-C10
29	Y	610	CLA	C11-C10-C8-C7
29	b	607	CLA	C11-C12-C13-C15
29	b	609	CLA	C11-C12-C13-C15
29	b	612	CLA	C11-C12-C13-C15
29	b	612	CLA	C12-C13-C15-C16
29	b	613	CLA	C6-C7-C8-C10

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Mol	Chain	Res	Type	Atoms
29	c	503	CLA	C11-C10-C8-C7
29	c	511	CLA	C3A-C2A-CAA-CBA
29	g	613	CLA	C11-C12-C13-C15
45	S	608	CHL	C11-C10-C8-C7
45	n	601	CHL	C11-C10-C8-C7
29	S	603	CLA	CAA-CBA-CGA-O1A
39	y	624	LHG	O10-C23-C24-C25
38	C	523	DGD	C4D-C5D-C6D-O5D
38	C	523	DGD	O6D-C5D-C6D-O5D
29	Y	614	CLA	CAA-CBA-CGA-O2A
29	c	505	CLA	CAA-CBA-CGA-O2A
29	y	614	CLA	CAA-CBA-CGA-O2A
38	C	523	DGD	O1G-C1A-C2A-C3A
39	Y	624	LHG	O8-C23-C24-C25
39	d	408	LHG	O7-C7-C8-C9
39	s	624	LHG	O8-C23-C24-C25
29	C	511	CLA	C8-C10-C11-C12
51	Y	625	SPH	C4-C5-C6-C7
33	c	521	LMG	C30-C31-C32-C33
38	C	519	DGD	C1B-C2B-C3B-C4B
31	b	618	BCR	C21-C22-C23-C24
46	y	620	LUT	C27-C28-C29-C30
29	c	503	CLA	CAA-CBA-CGA-O1A
32	c	626	SQD	O49-C7-C8-C9
45	y	606	CHL	CAA-CBA-CGA-O1A
29	s	604	CLA	C6-C7-C8-C10
30	A	408	PHO	C16-C17-C18-C20
29	G	603	CLA	CAA-CBA-CGA-O2A
39	L	101	LHG	O7-C7-C8-C9
33	H	102	LMG	C16-C17-C18-C19
38	C	518	DGD	C6A-C7A-C8A-C9A
29	a	405	CLA	C13-C15-C16-C17
29	B	602	CLA	CAA-CBA-CGA-O1A
29	N	604	CLA	CAA-CBA-CGA-O1A
29	G	603	CLA	CAA-CBA-CGA-O1A
29	G	614	CLA	CAA-CBA-CGA-O1A
29	S	605	CLA	CAA-CBA-CGA-O1A
29	s	603	CLA	CAA-CBA-CGA-O1A
32	C	526	SQD	O10-C23-C24-C25
33	D	411	LMG	O9-C10-C11-C12
38	C	523	DGD	O1A-C1A-C2A-C3A
38	c	518	DGD	O1A-C1A-C2A-C3A

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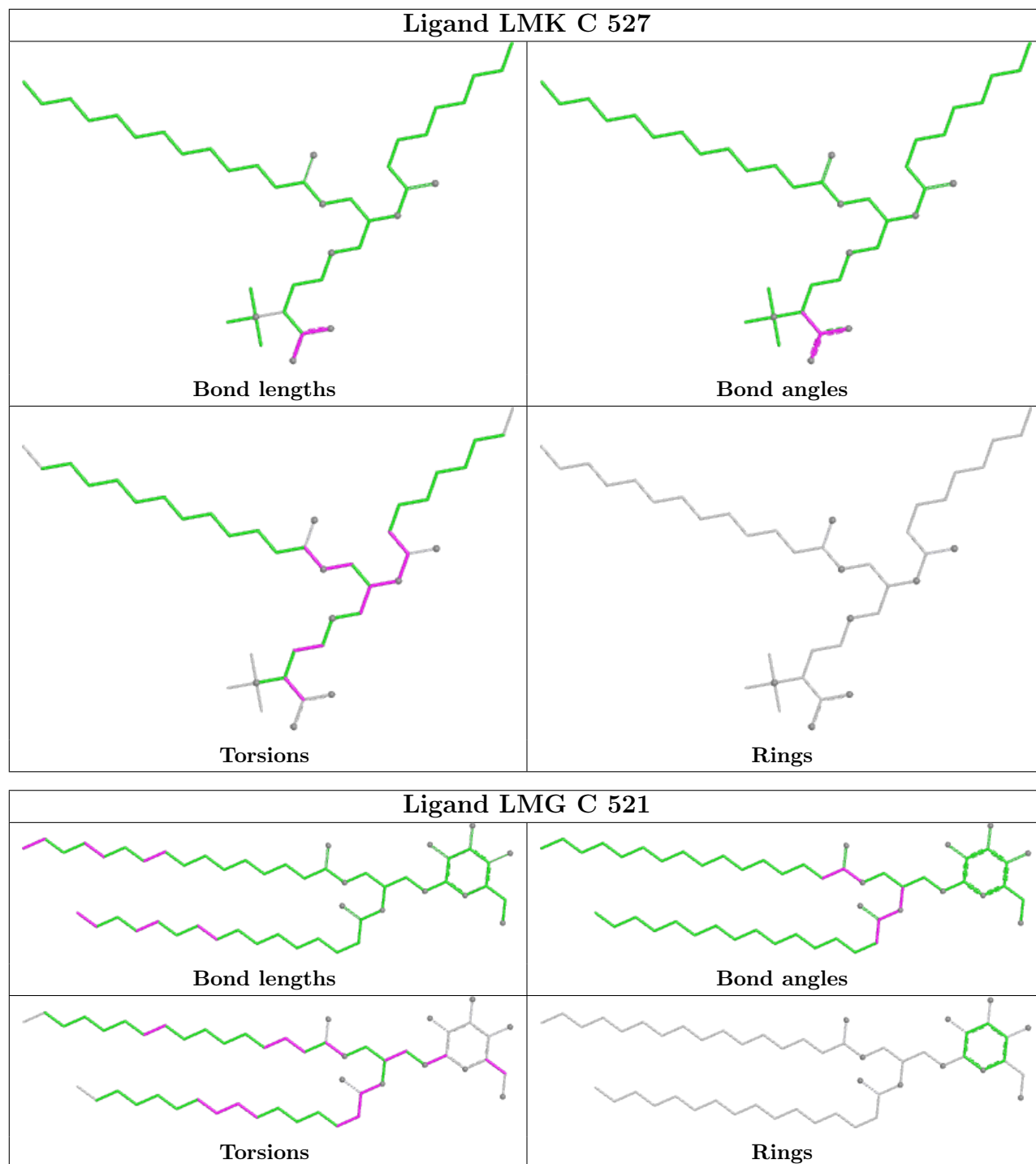
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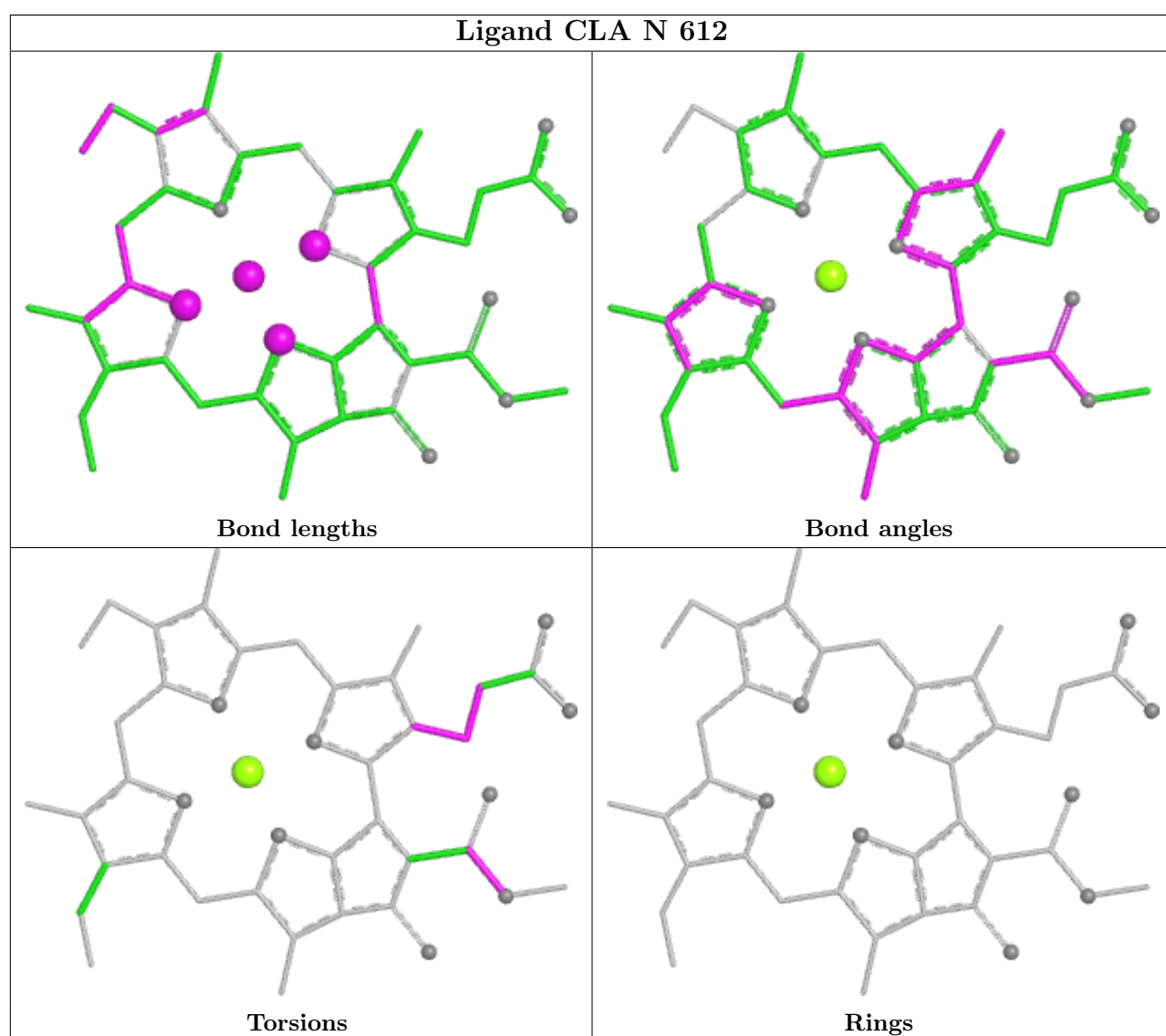
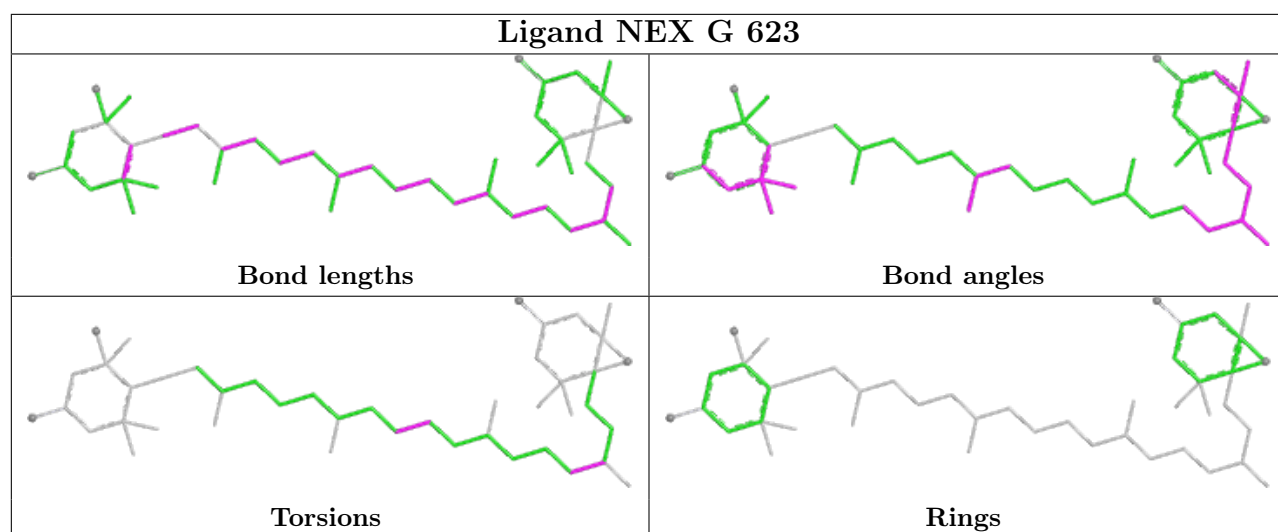
Mol	Chain	Res	Type	Atoms
29	r	612	CLA	C5-C6-C7-C8
45	g	601	CHL	C8-C10-C11-C12
45	y	607	CHL	C10-C11-C12-C13
33	h	102	LMG	C16-C17-C18-C19
50	i	101	3PH	C2C-C2D-C2E-C2F
29	c	502	CLA	CAA-CBA-CGA-O2A
29	c	508	CLA	CAA-CBA-CGA-O2A
32	c	626	SQD	O47-C7-C8-C9
39	S	624	LHG	O8-C23-C24-C25
39	d	409	LHG	C35-C36-C37-C38
29	r	604	CLA	CAA-CBA-CGA-O1A
45	n	608	CHL	C2A-CAA-CBA-CGA
29	d	402	CLA	O1A-CGA-O2A-C1
32	c	626	SQD	C11-C12-C13-C14
39	s	624	LHG	O10-C23-C24-C25
39	d	410	LHG	C10-C11-C12-C13
29	C	513	CLA	CAA-CBA-CGA-O2A

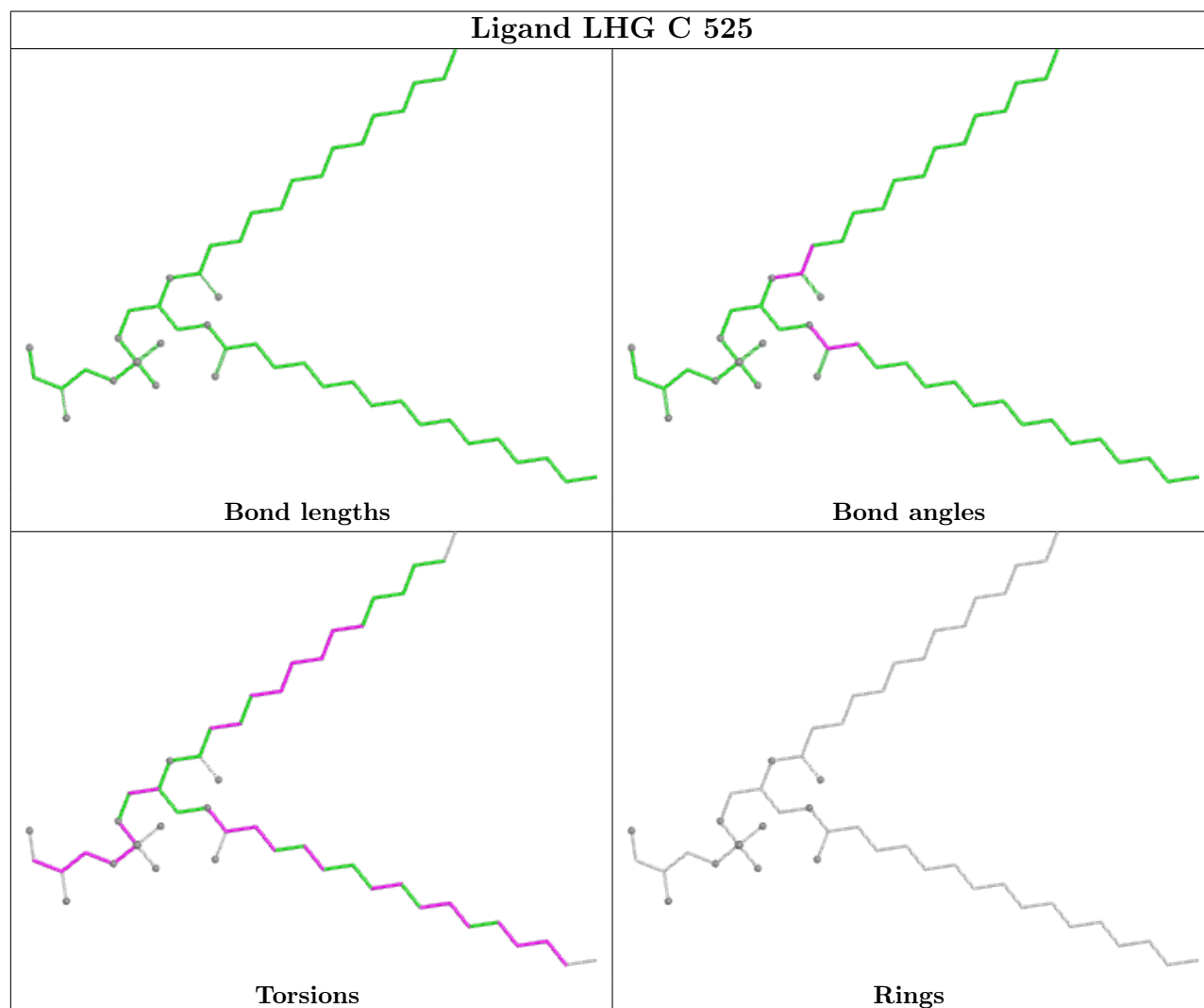
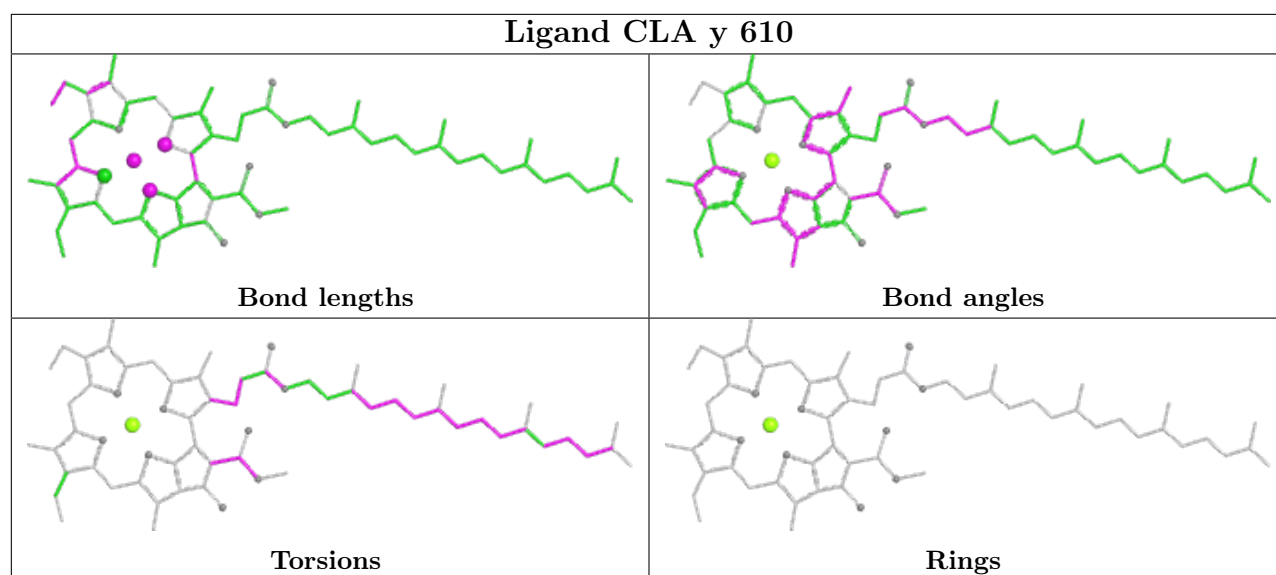
There are no ring outliers.

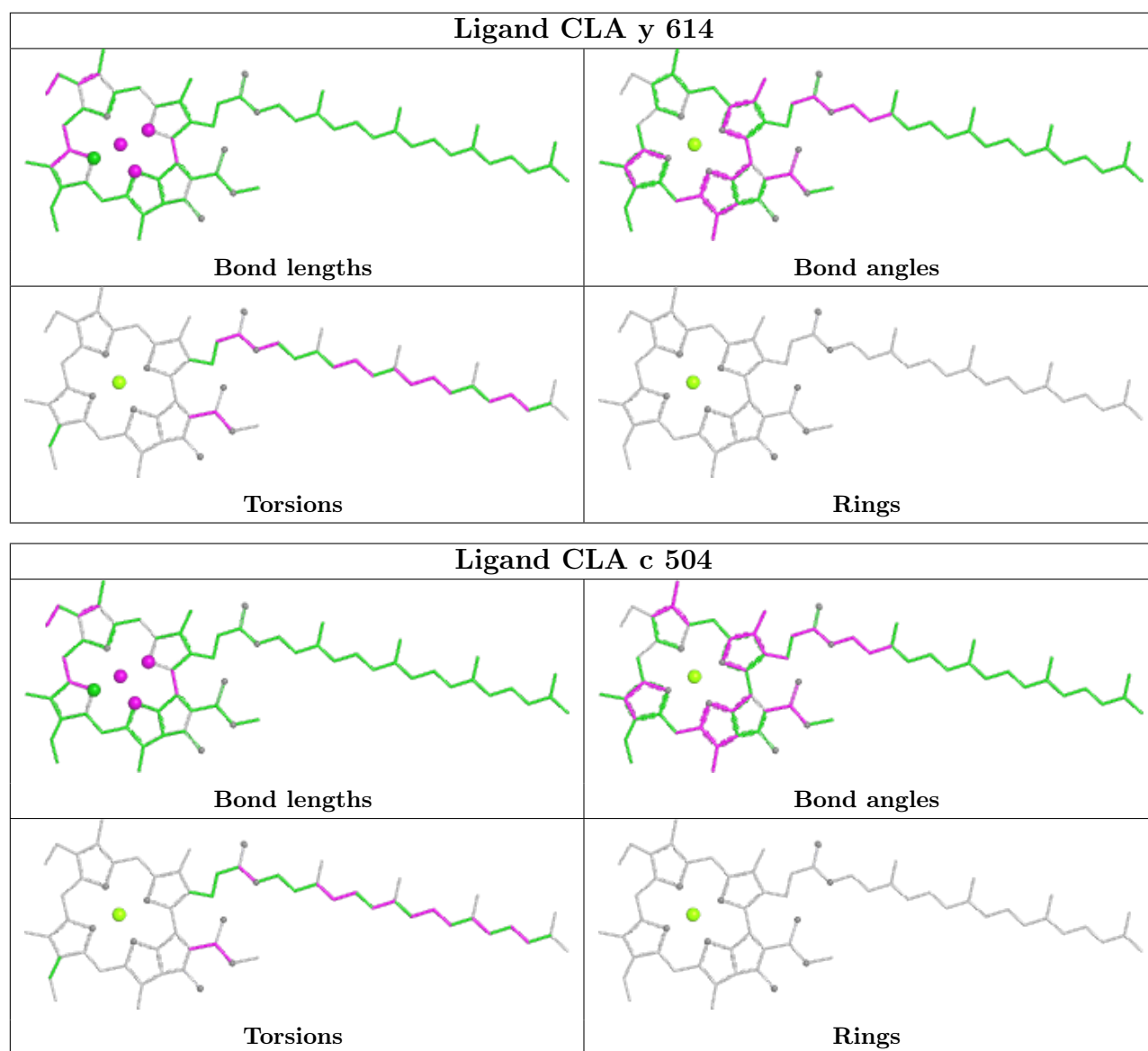
No monomer is involved in short contacts.

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.

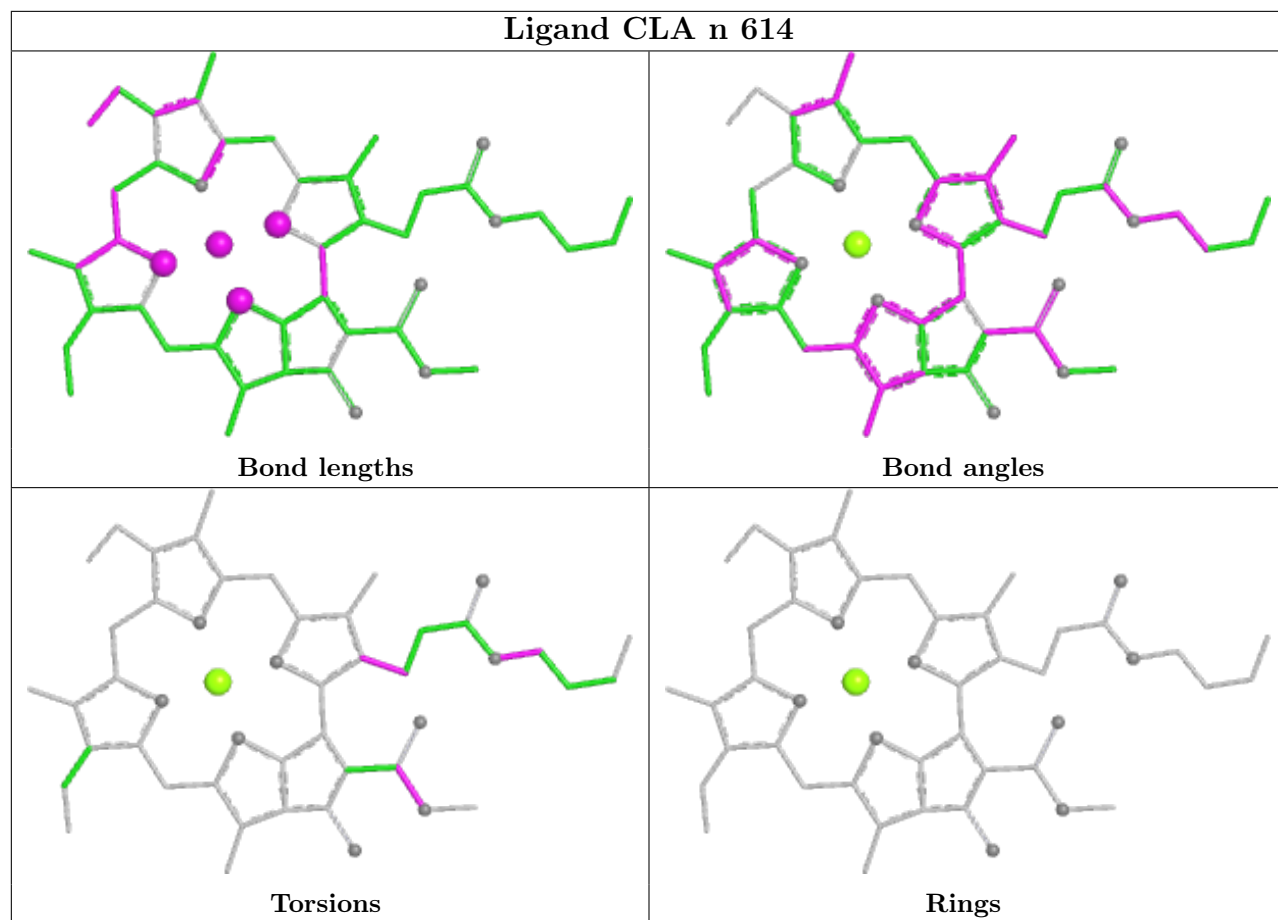


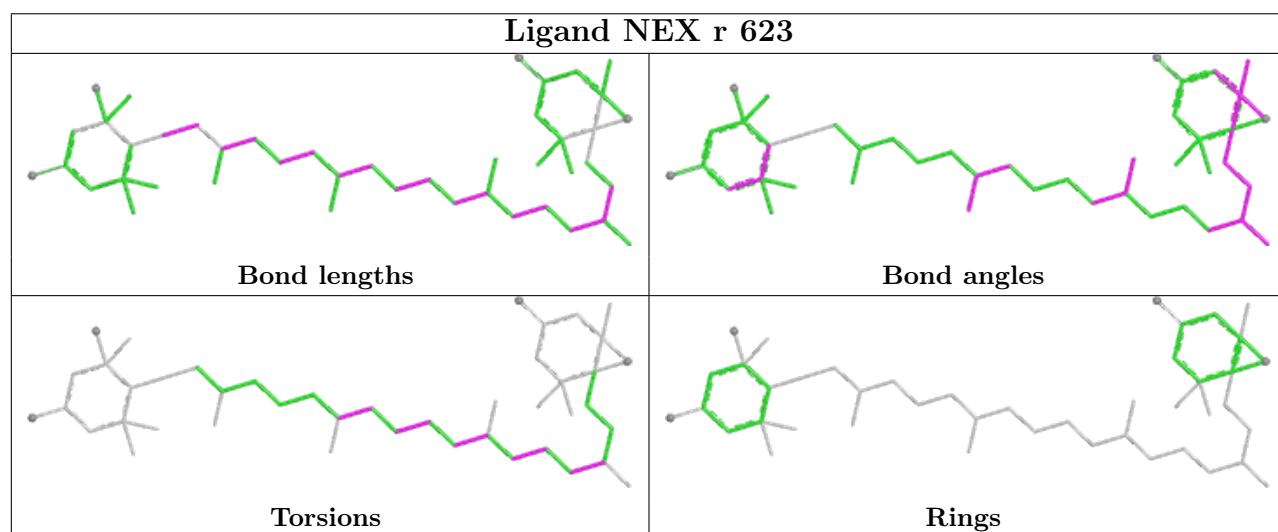
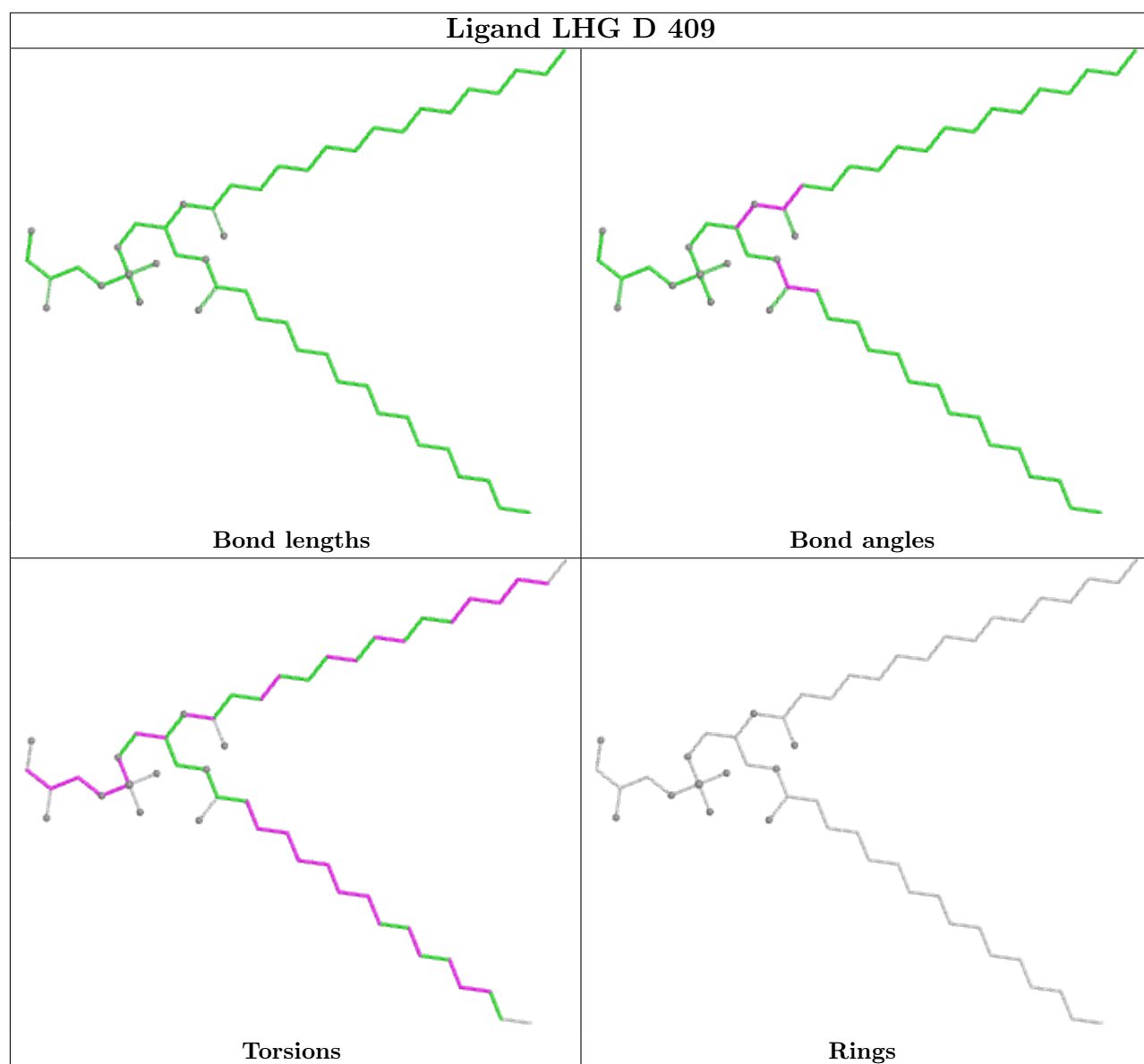


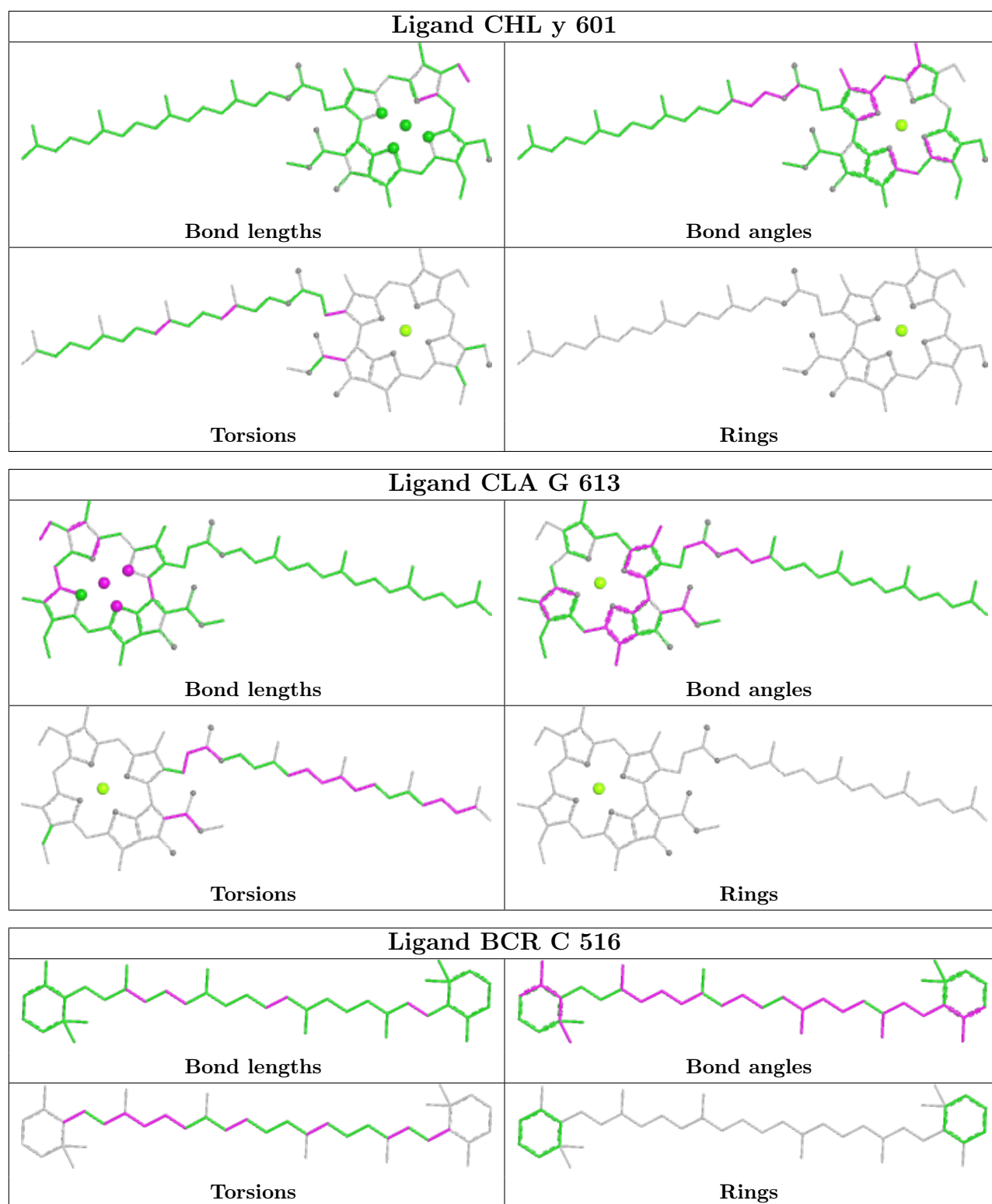


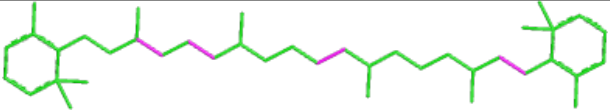
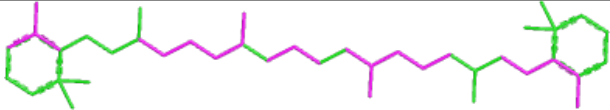
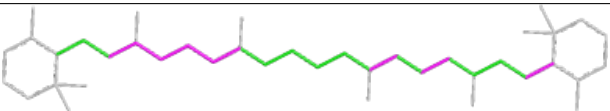
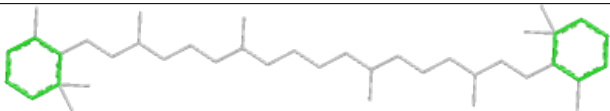
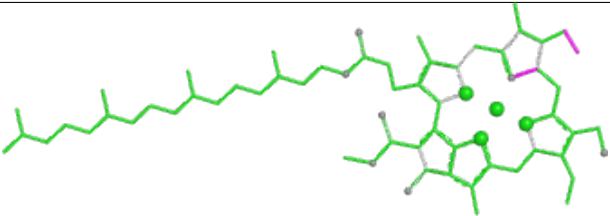
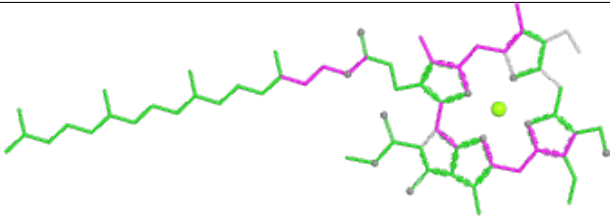
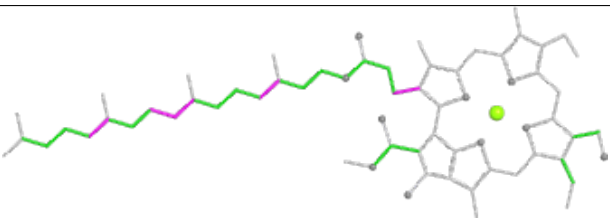
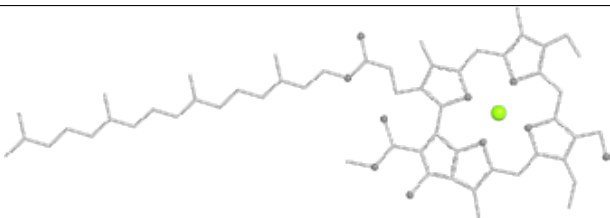
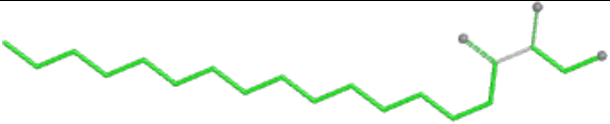
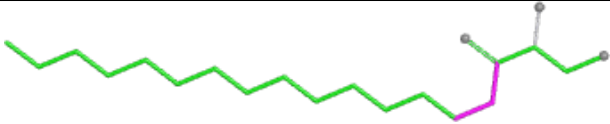
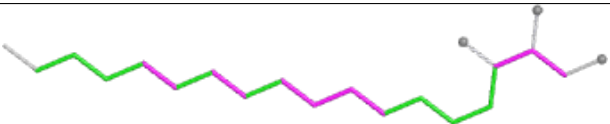
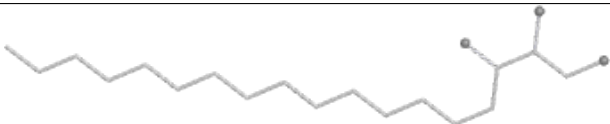


Ligand CLA n 614

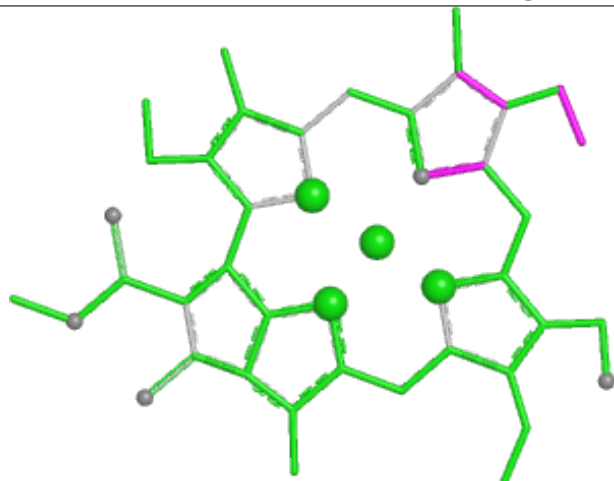




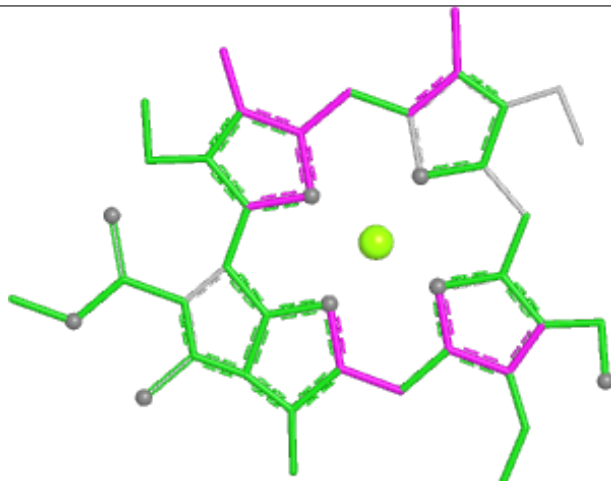


Ligand BCR a 411	
 <p>Bond lengths</p>	 <p>Bond angles</p>
 <p>Torsions</p>	 <p>Rings</p>
Ligand CHL N 609	
 <p>Bond lengths</p>	 <p>Bond angles</p>
 <p>Torsions</p>	 <p>Rings</p>
Ligand SPH y 625	
 <p>Bond lengths</p>	 <p>Bond angles</p>
 <p>Torsions</p>	 <p>Rings</p>

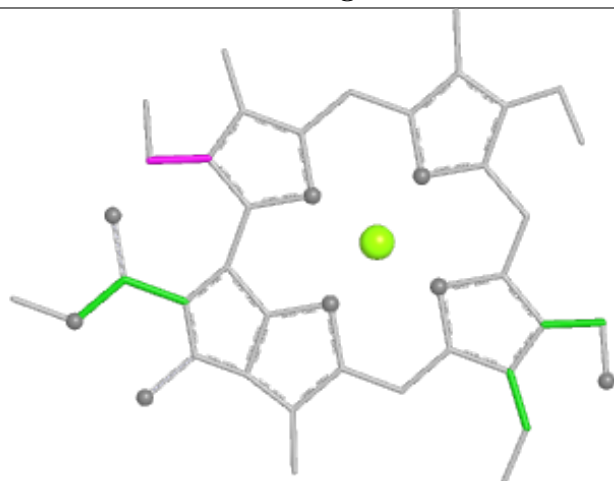
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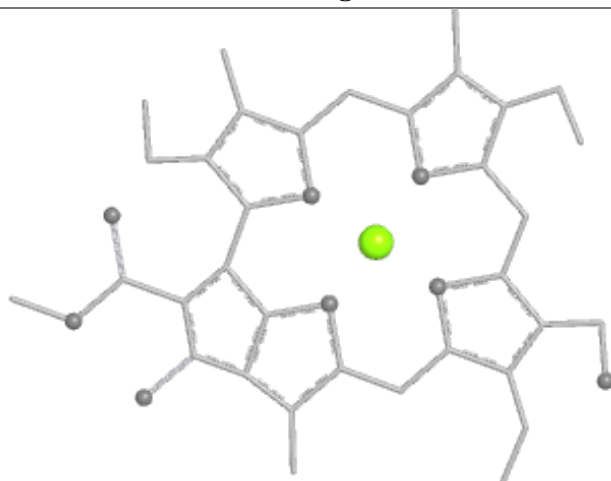
Bond lengths



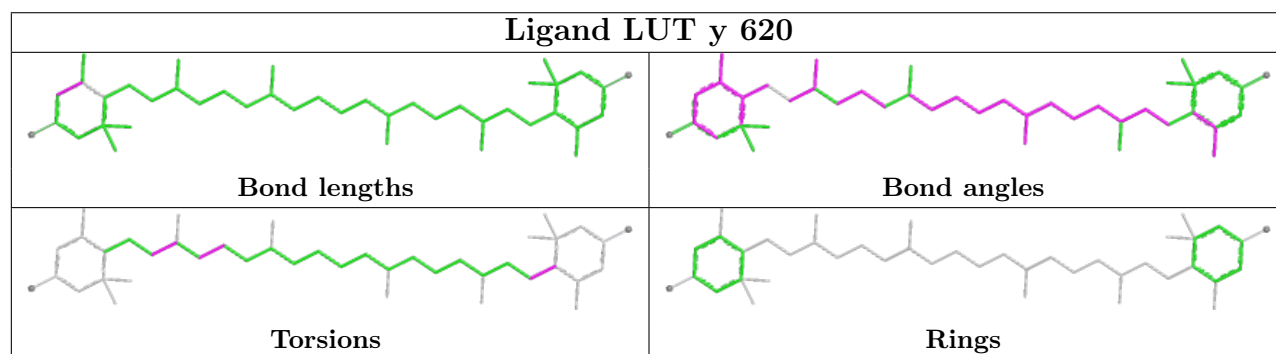
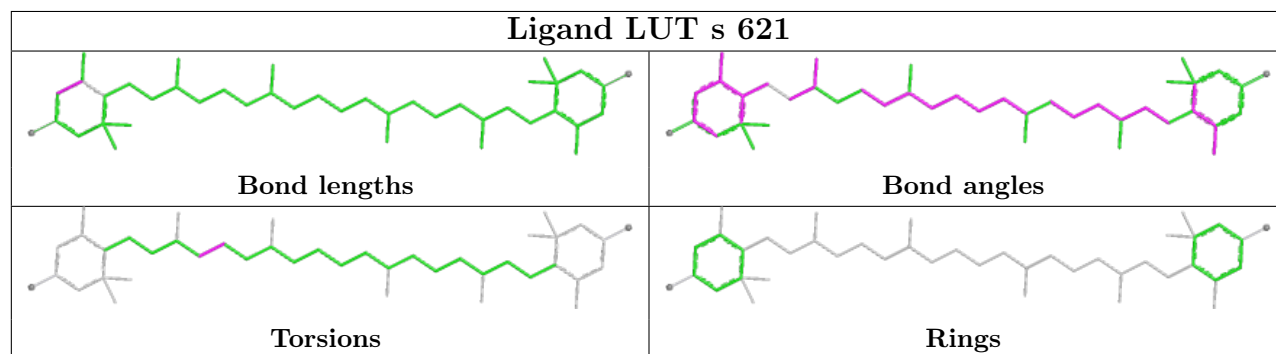
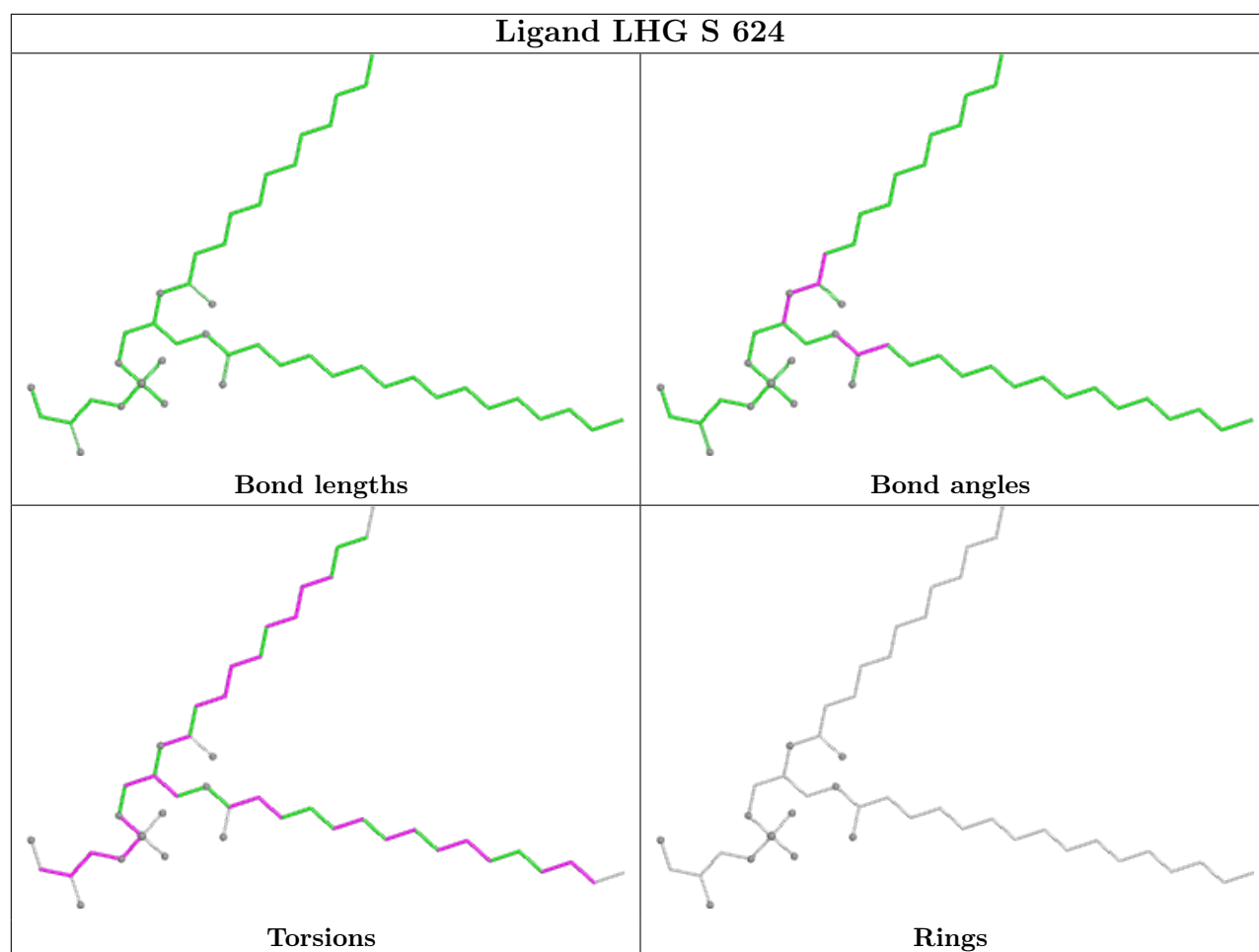
Bond angles

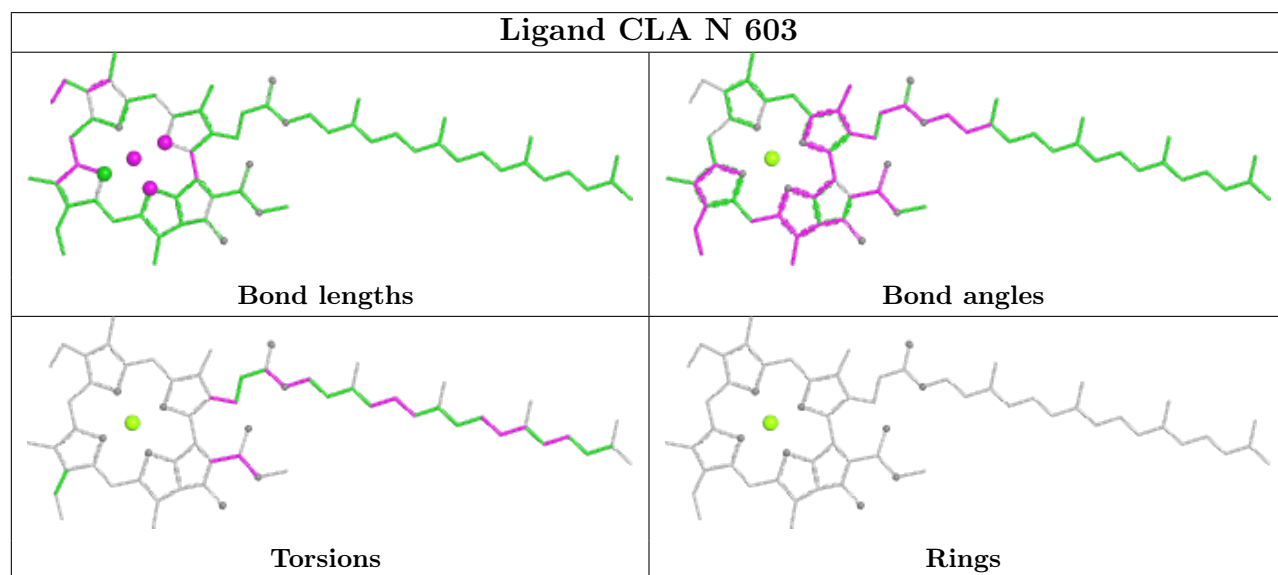
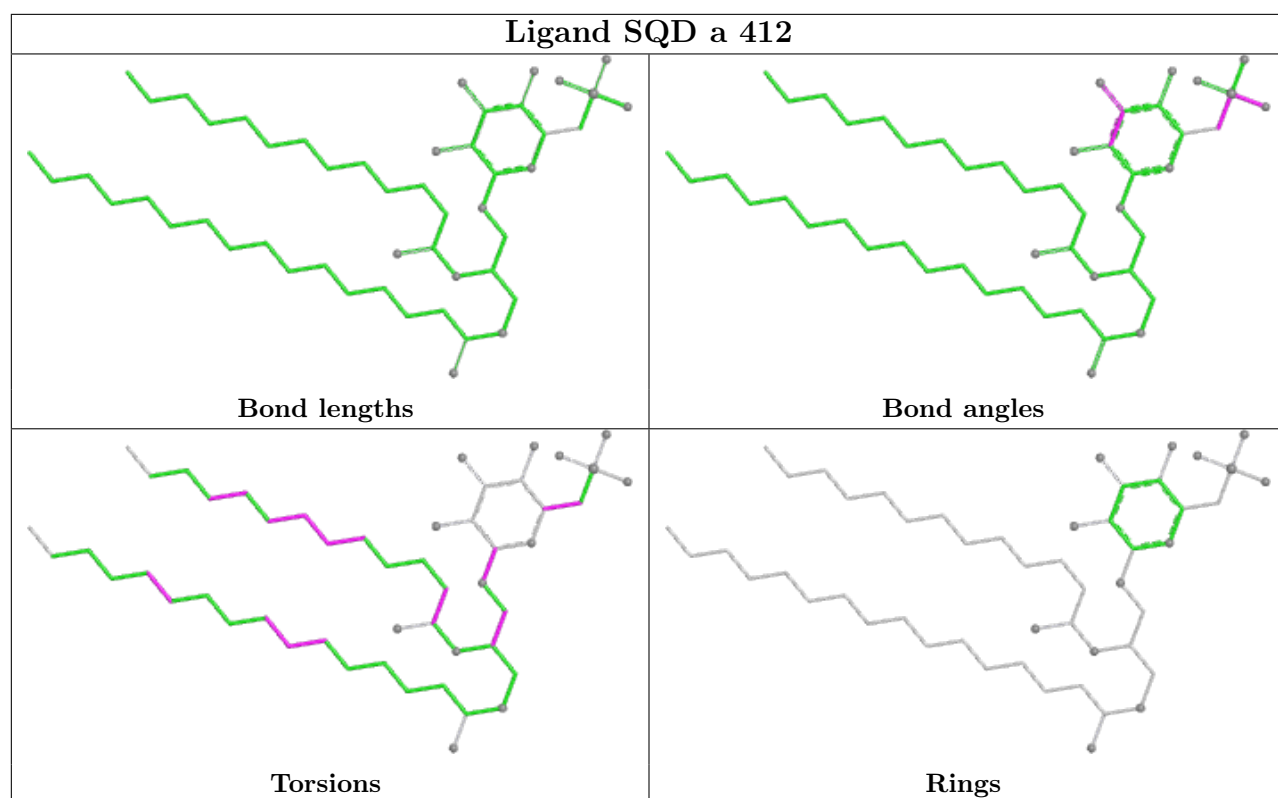


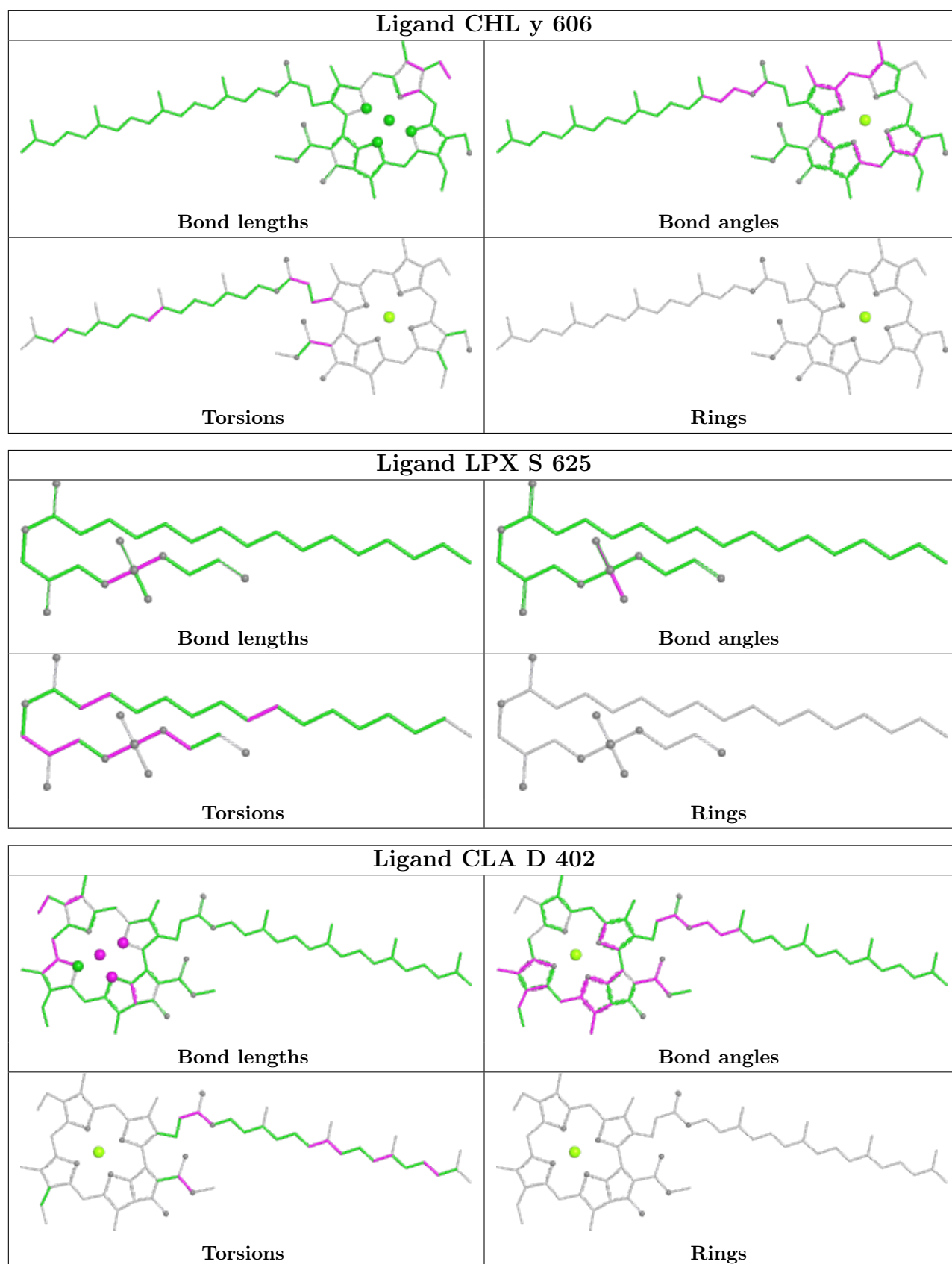
Torsions

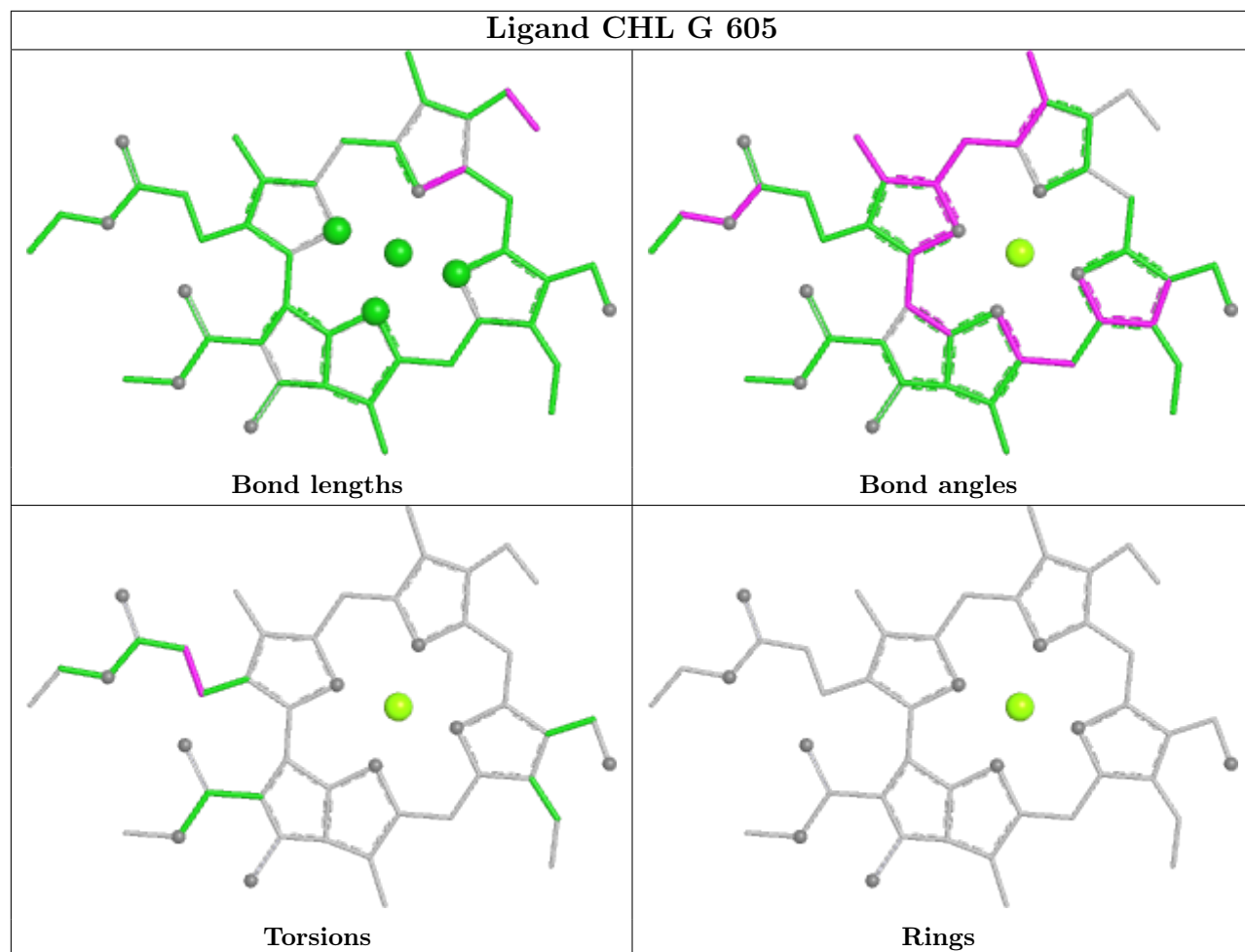


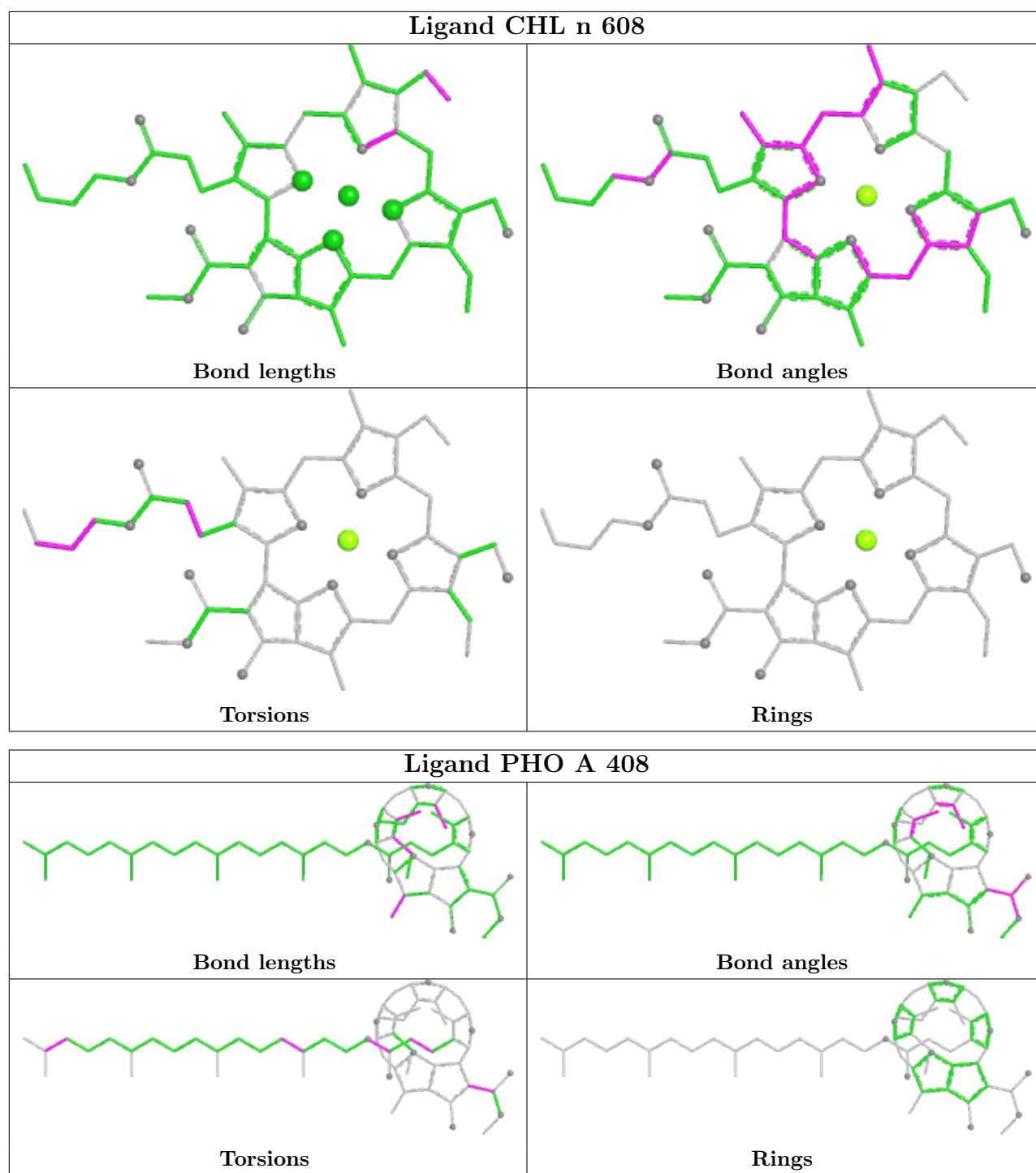
Rings



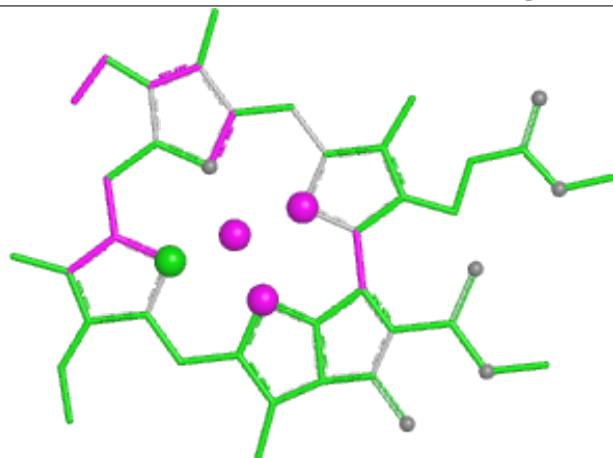




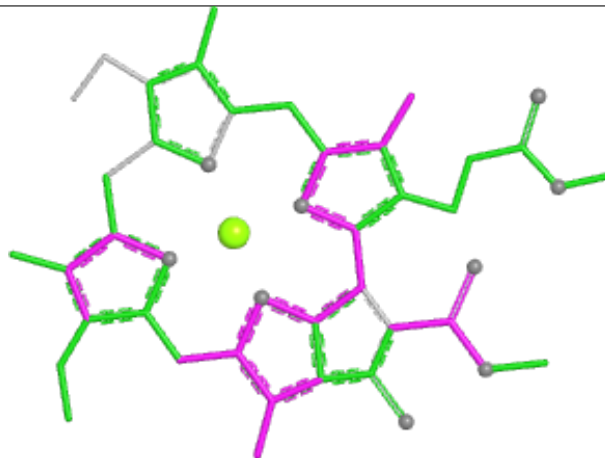




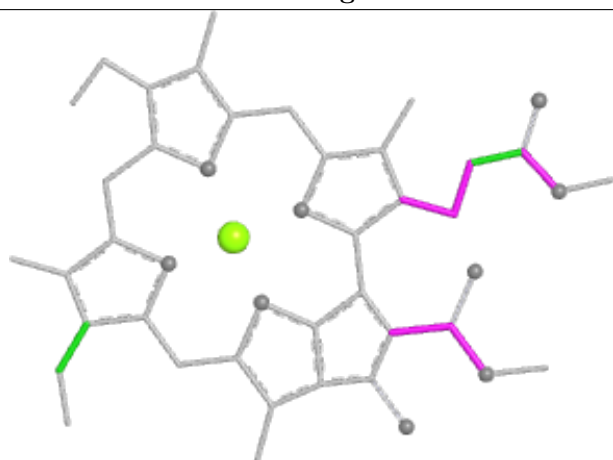
Ligand CLA r 613



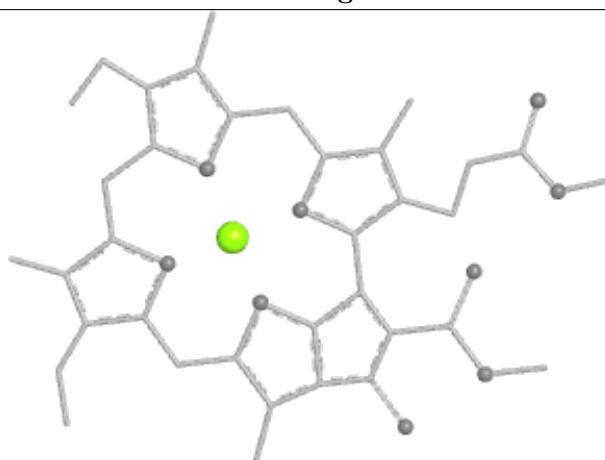
Bond lengths



Bond angles

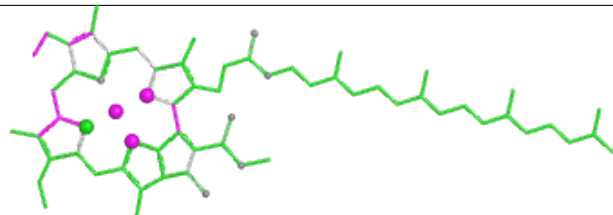


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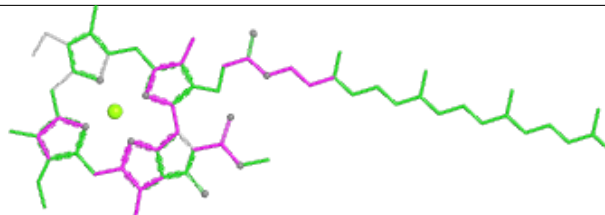


Rings

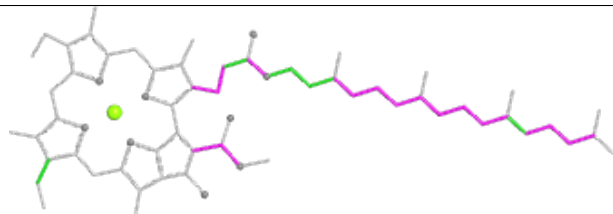
Ligand CLA Y 610



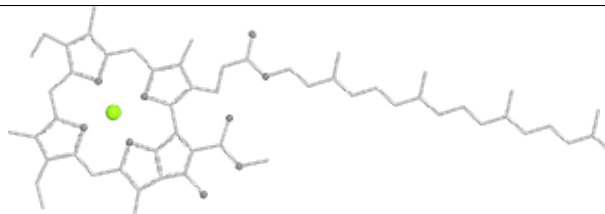
Bond lengths



Bond angles

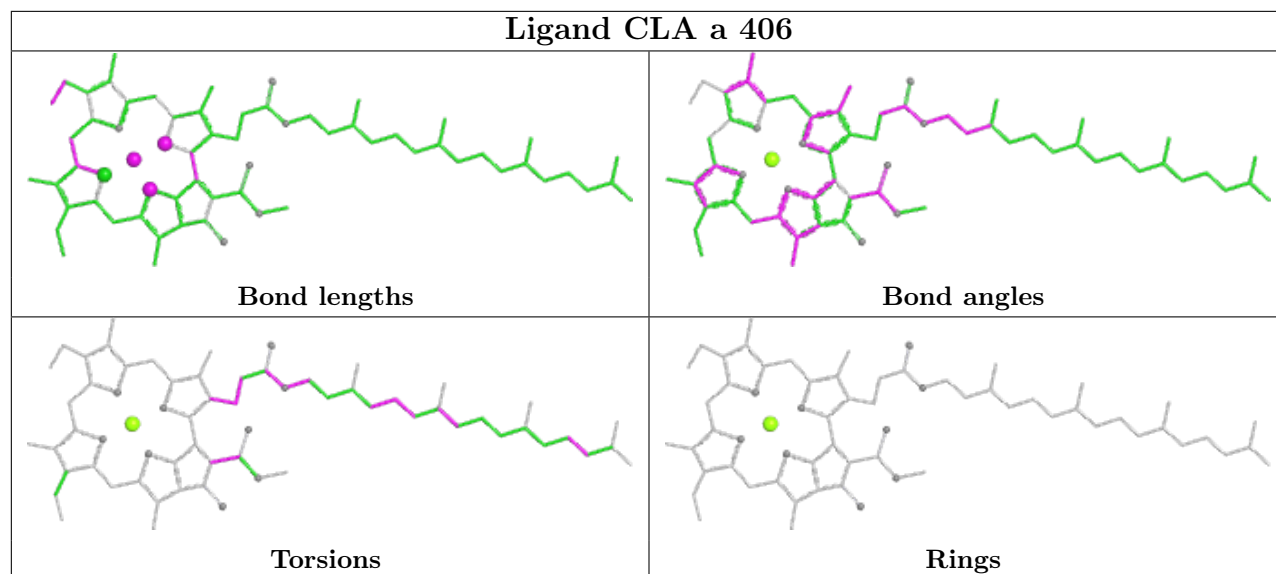


Torsions

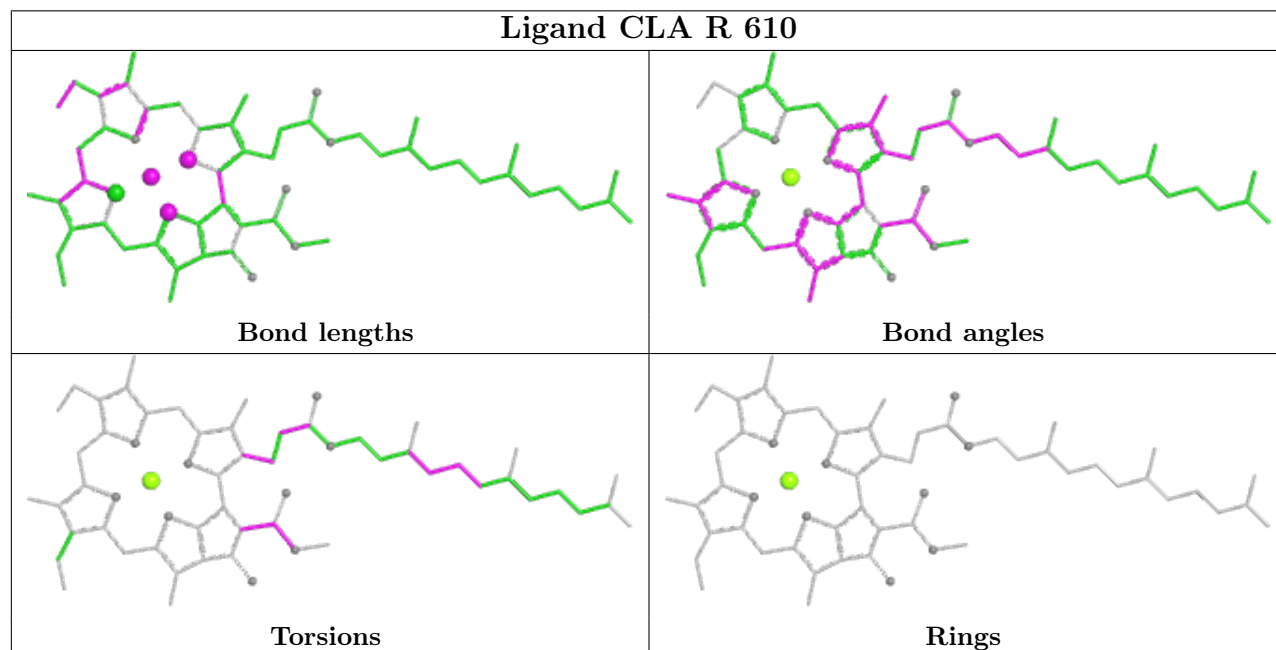


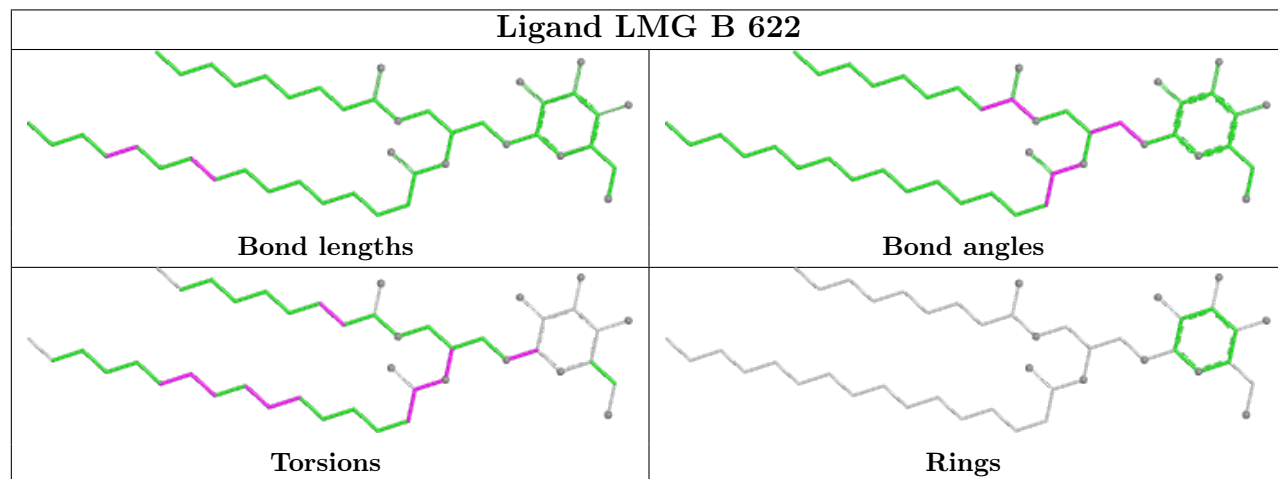
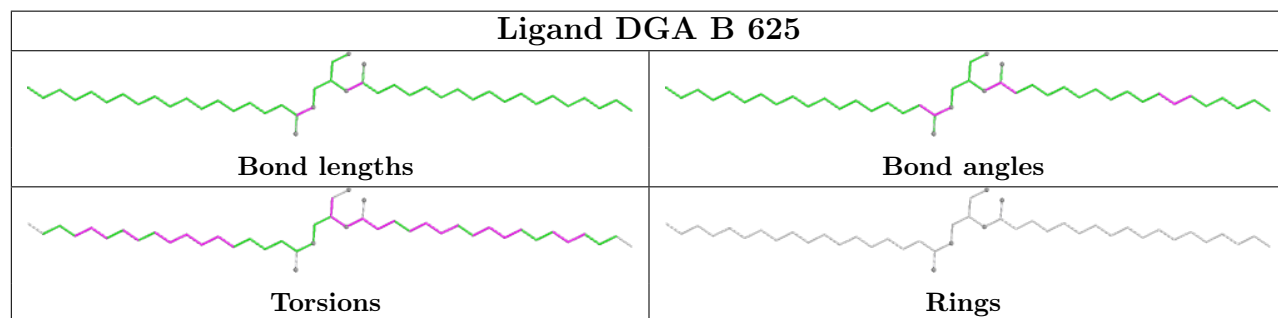
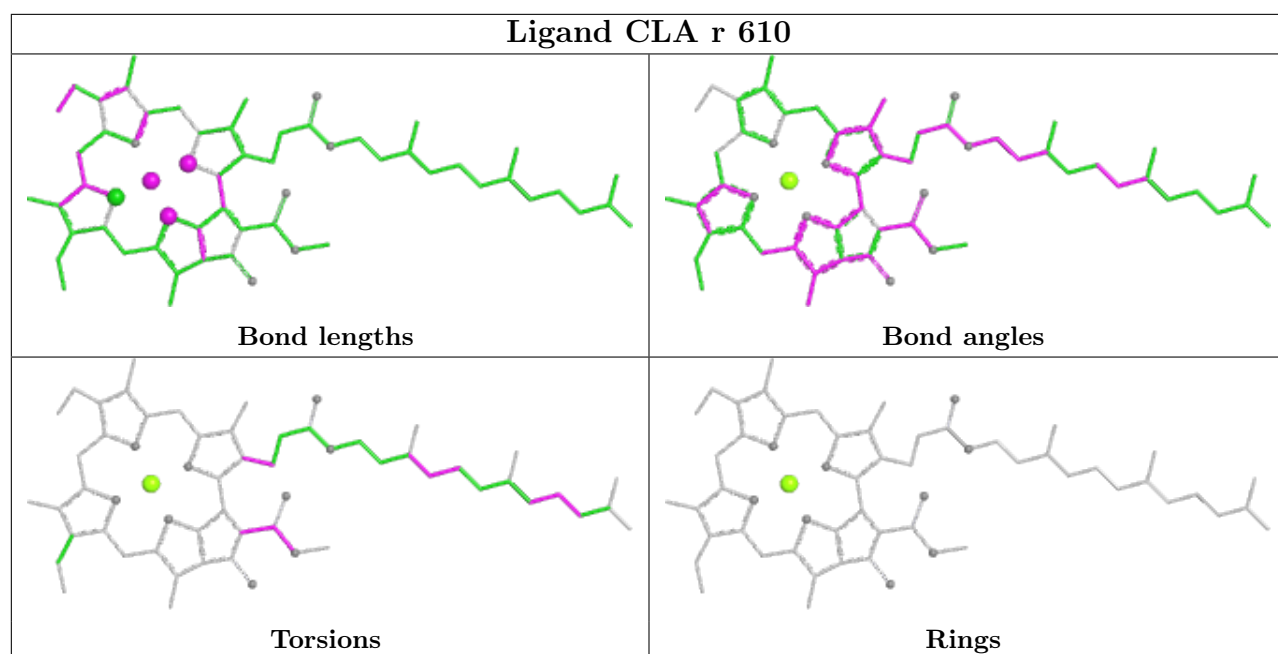
Rings

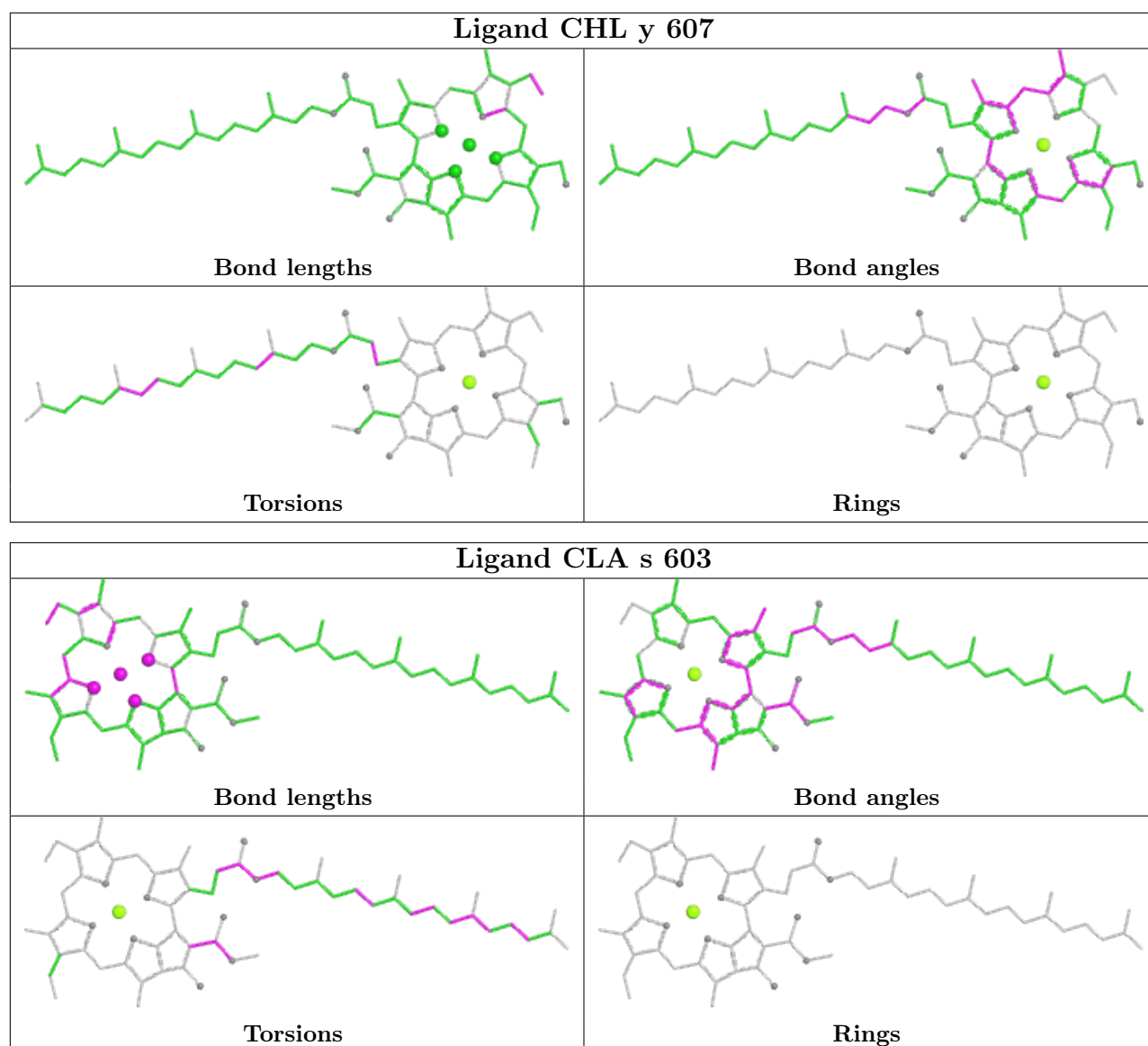
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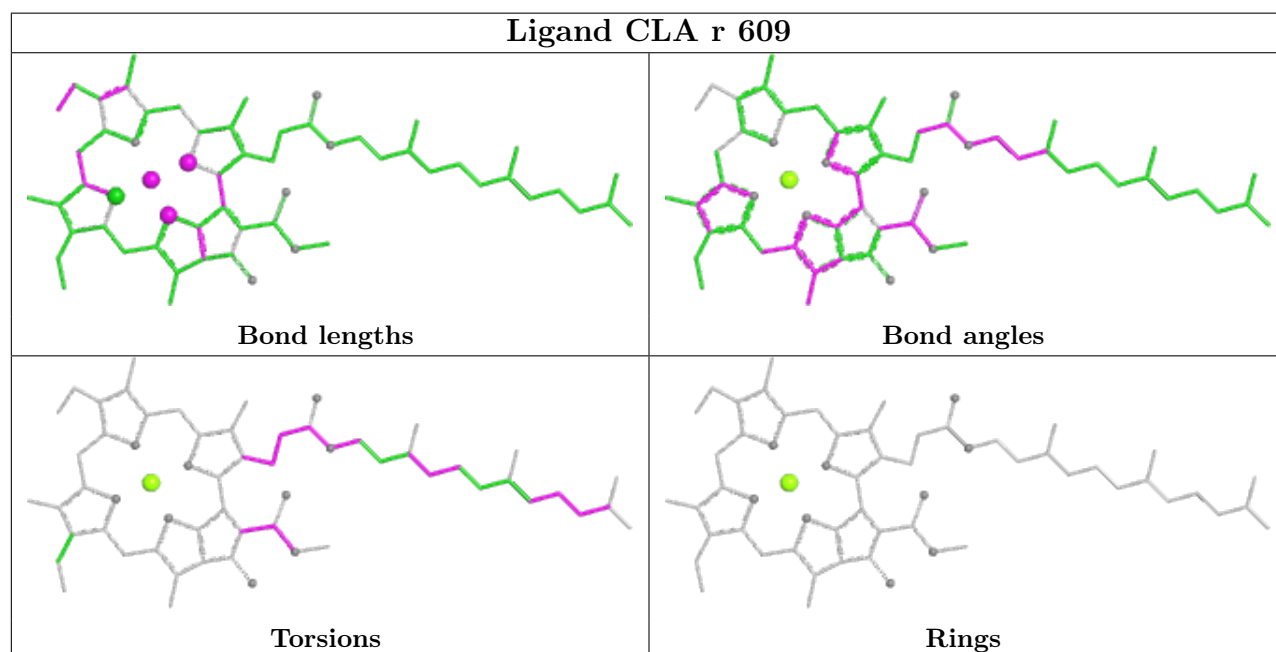
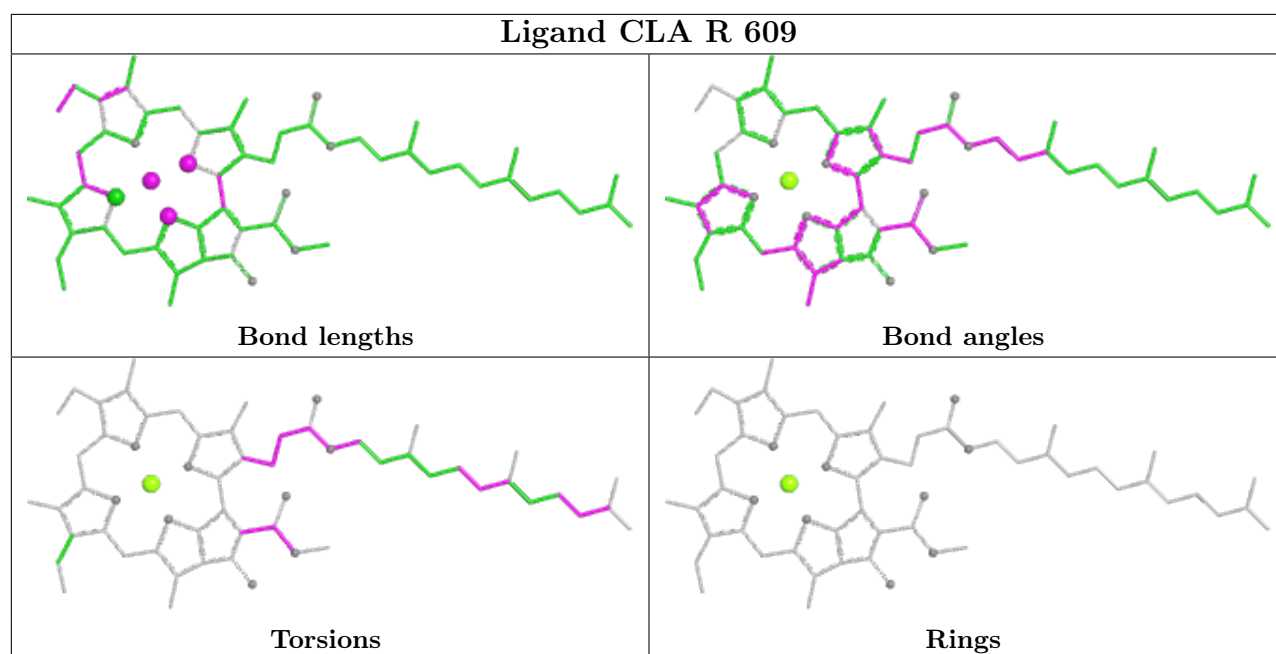


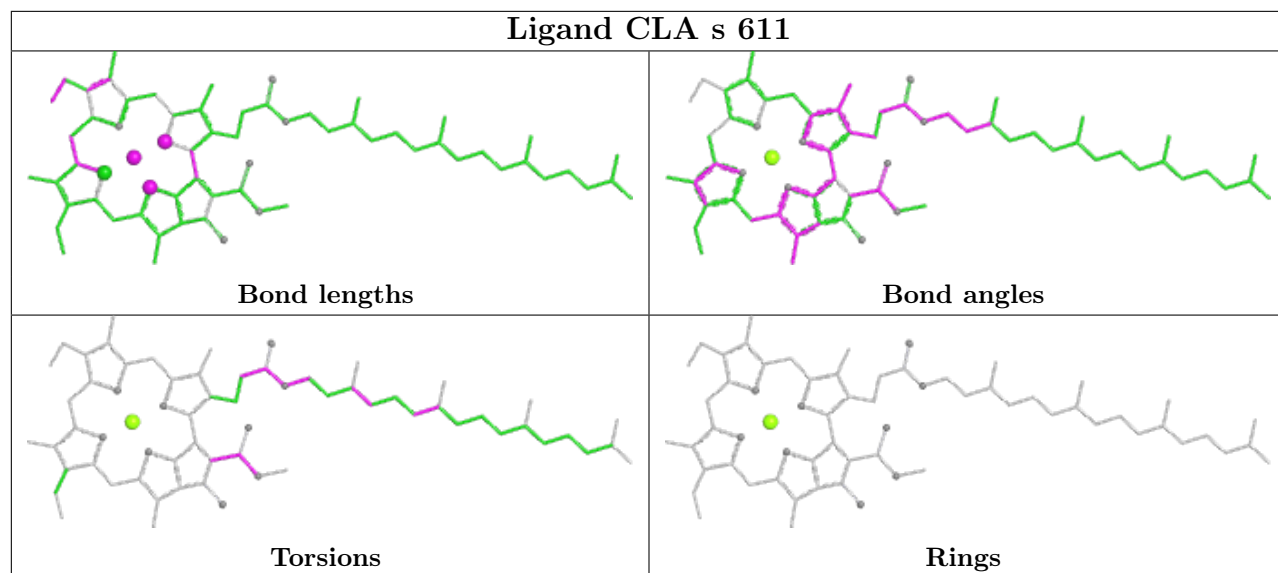
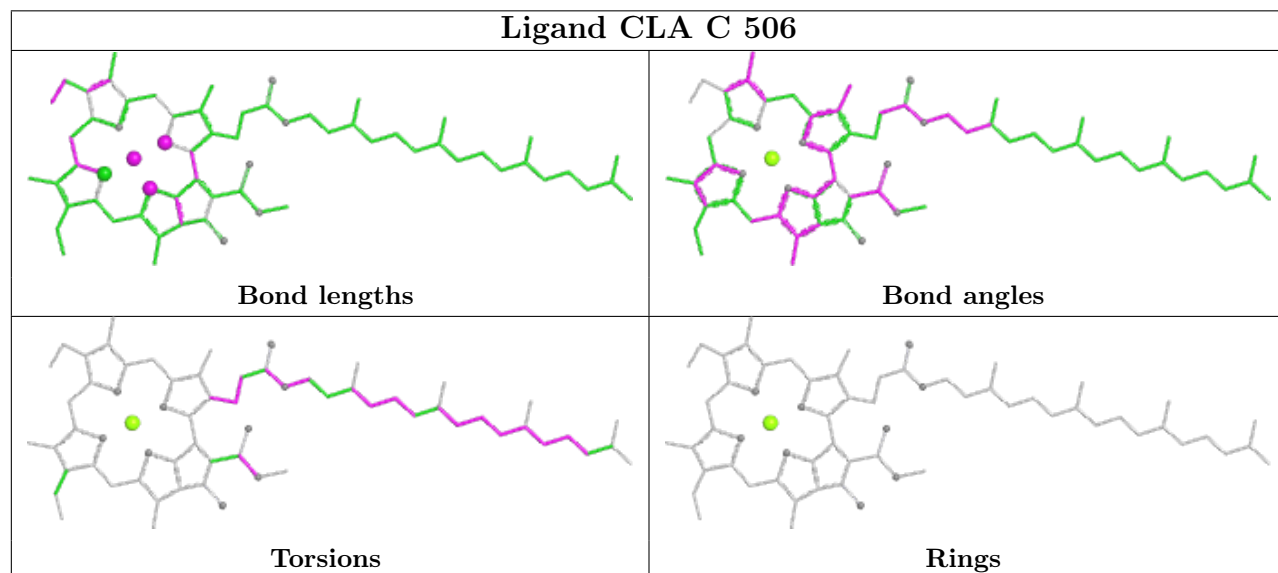
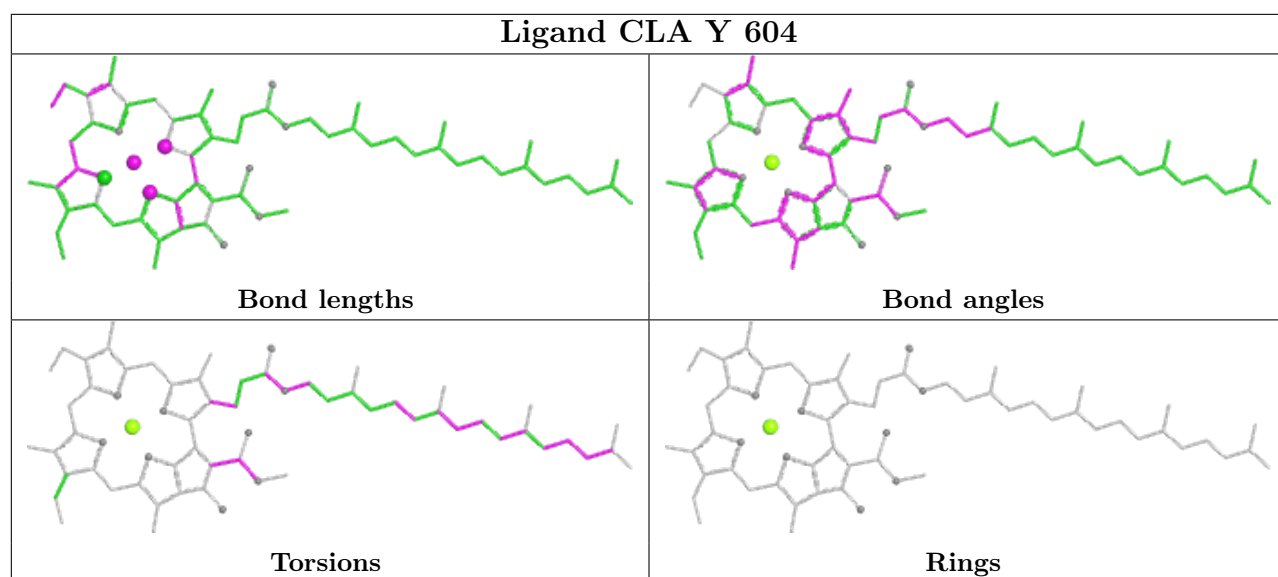
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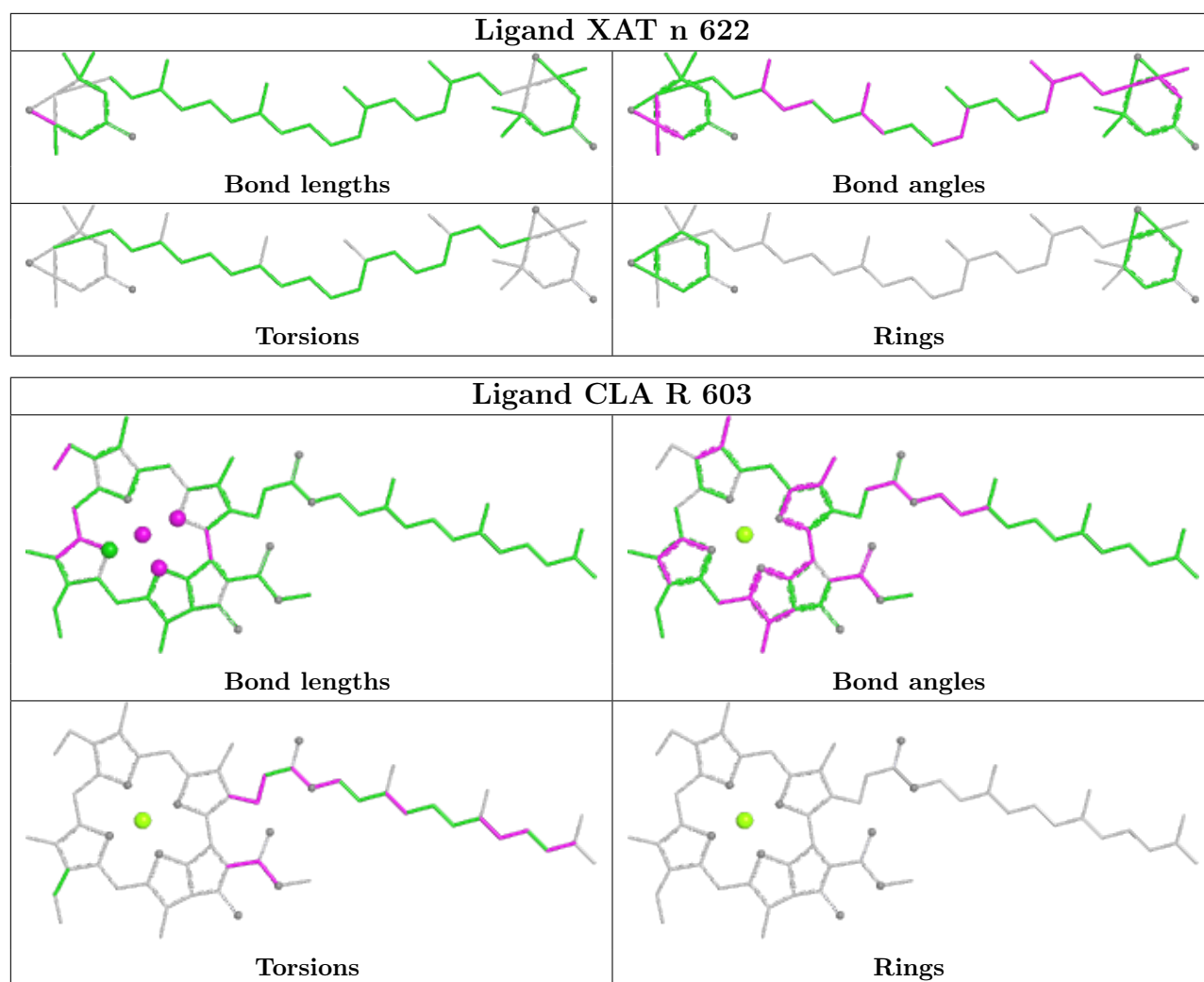


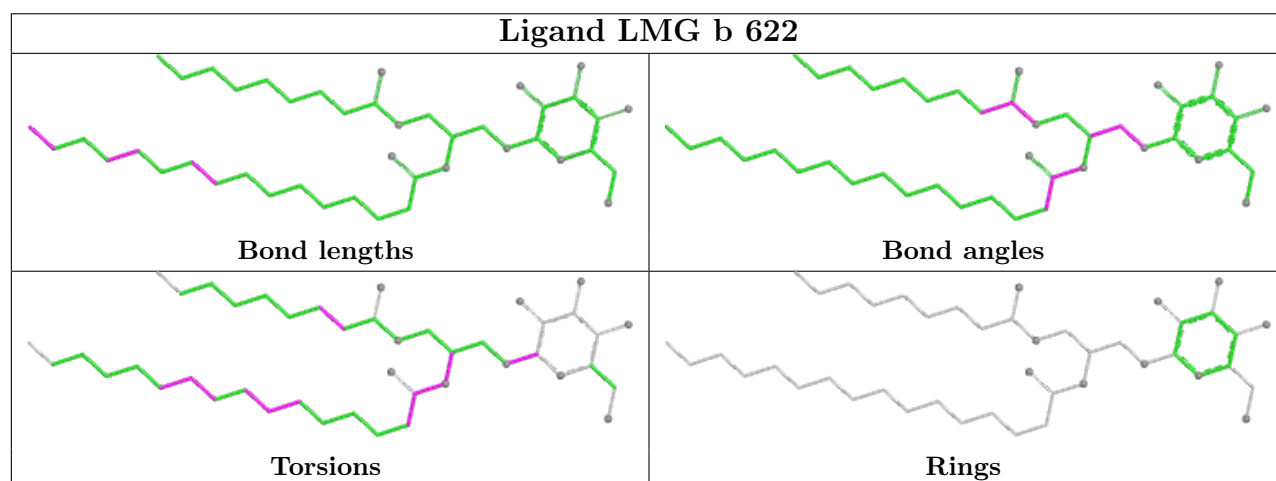
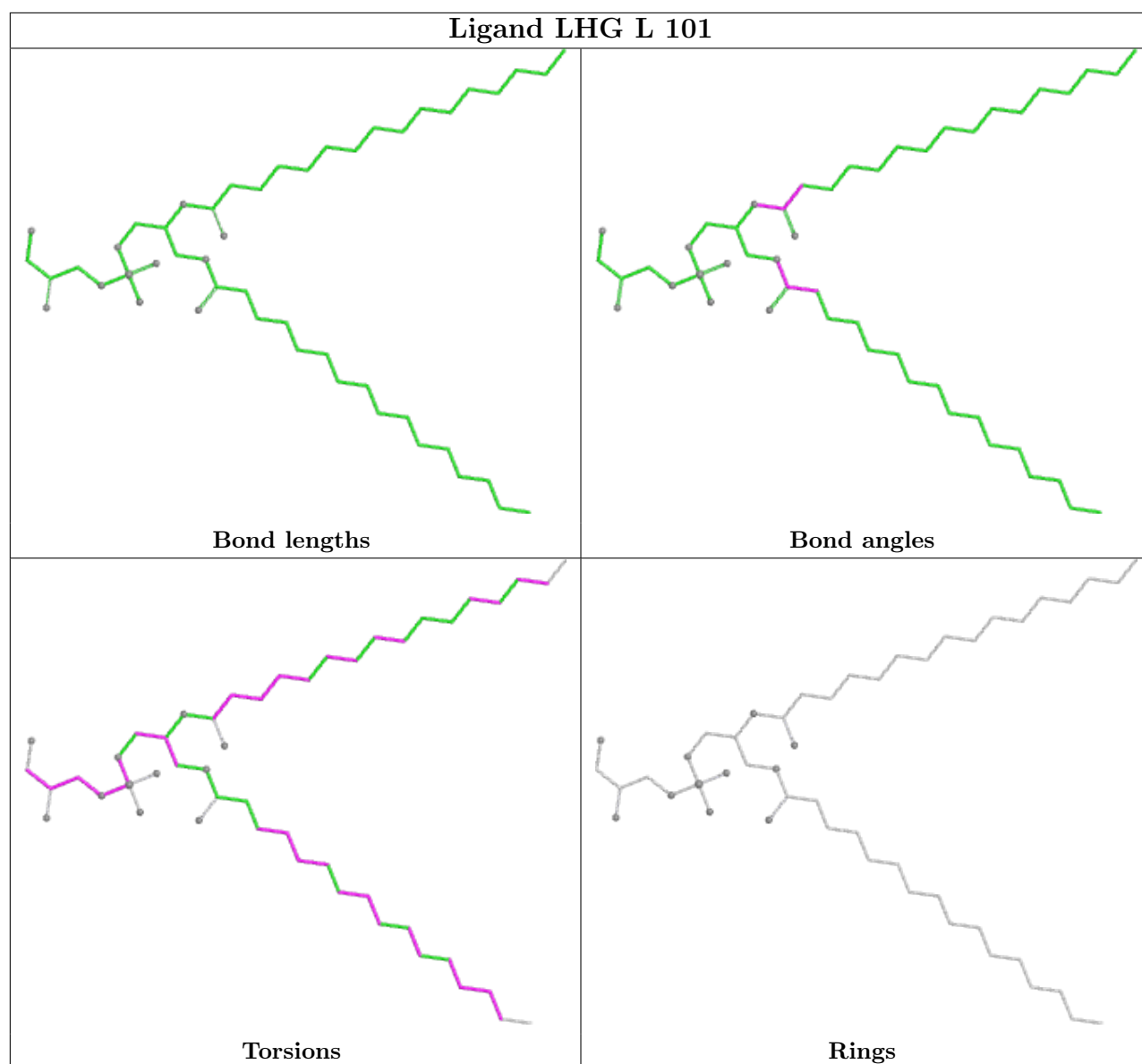


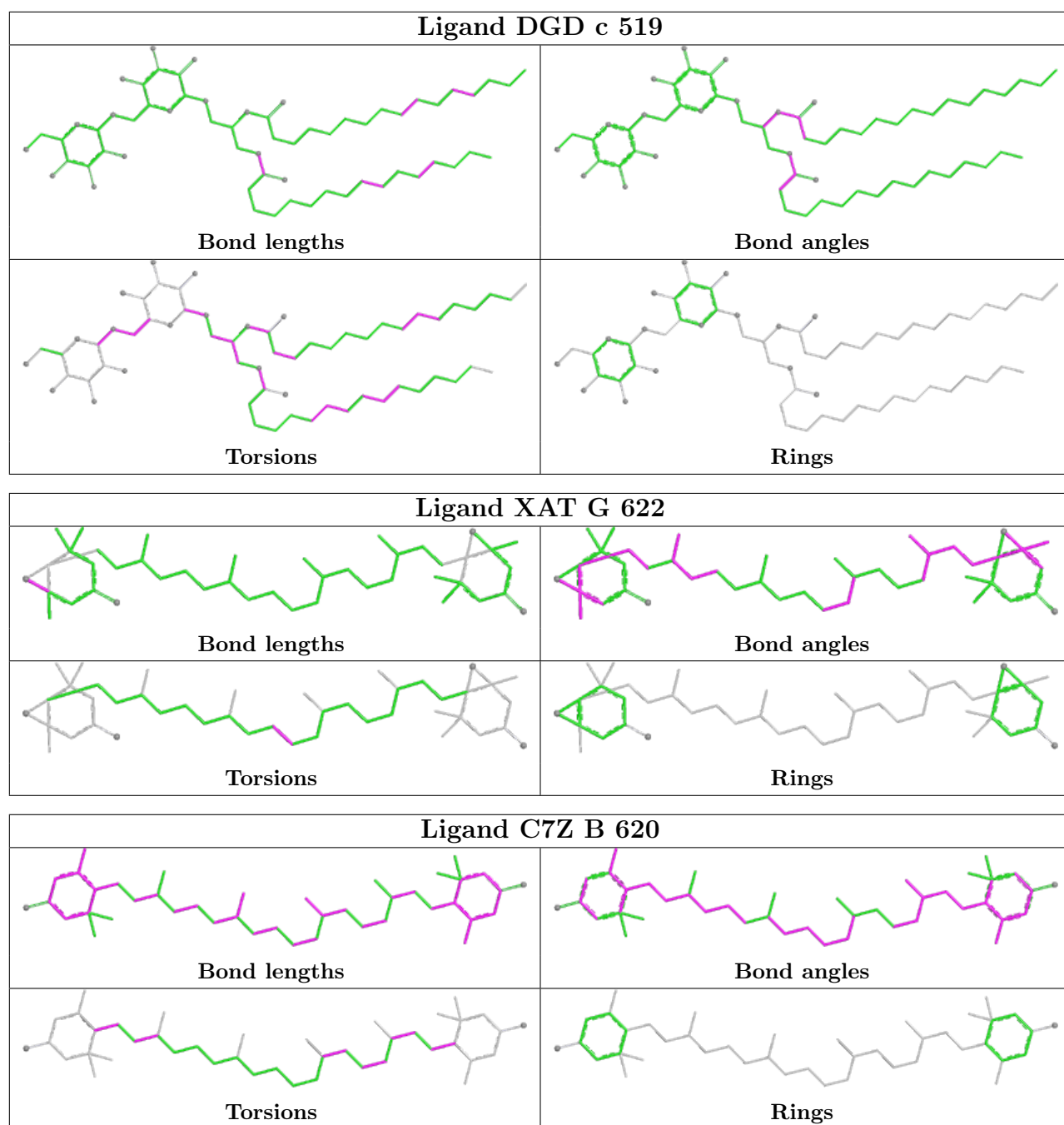


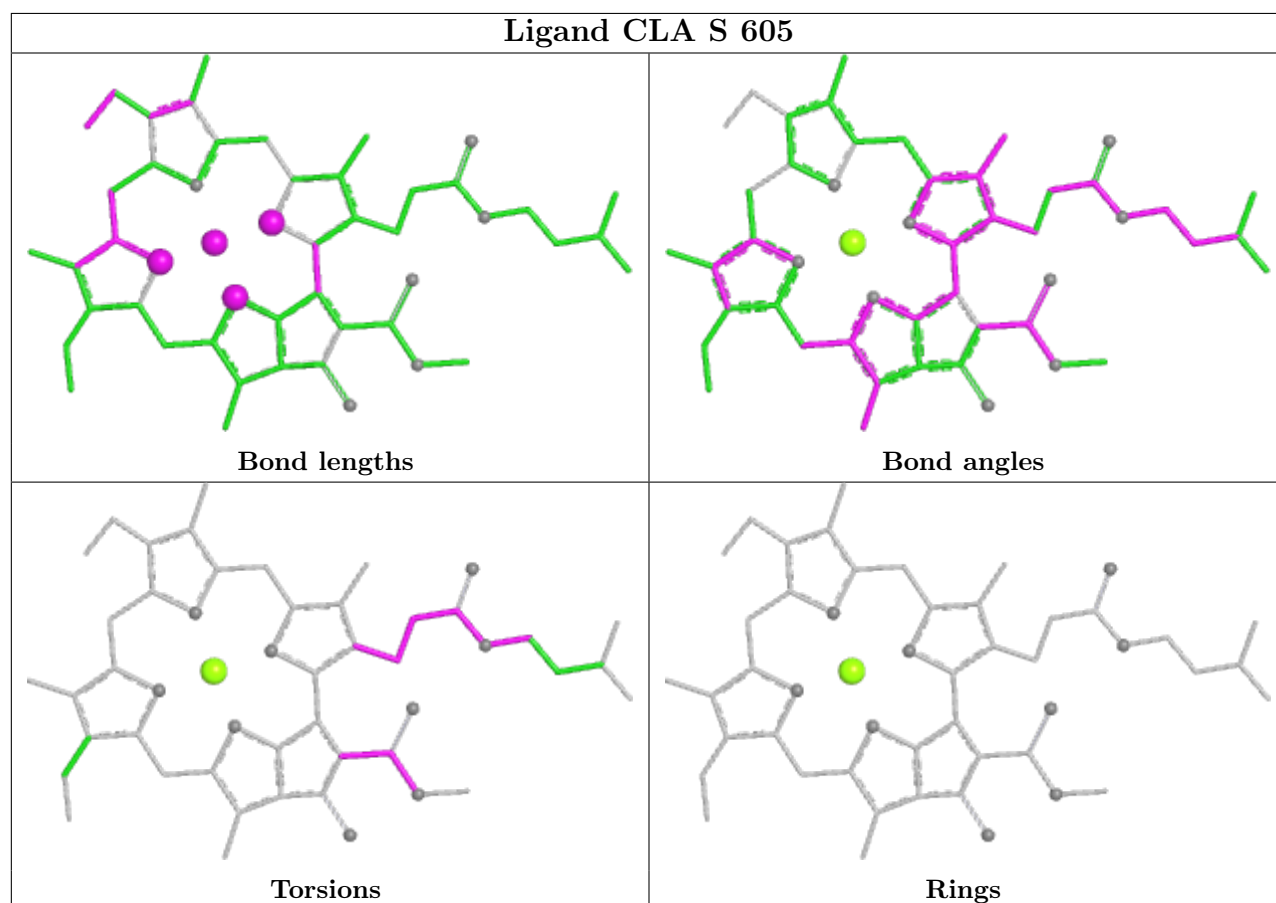
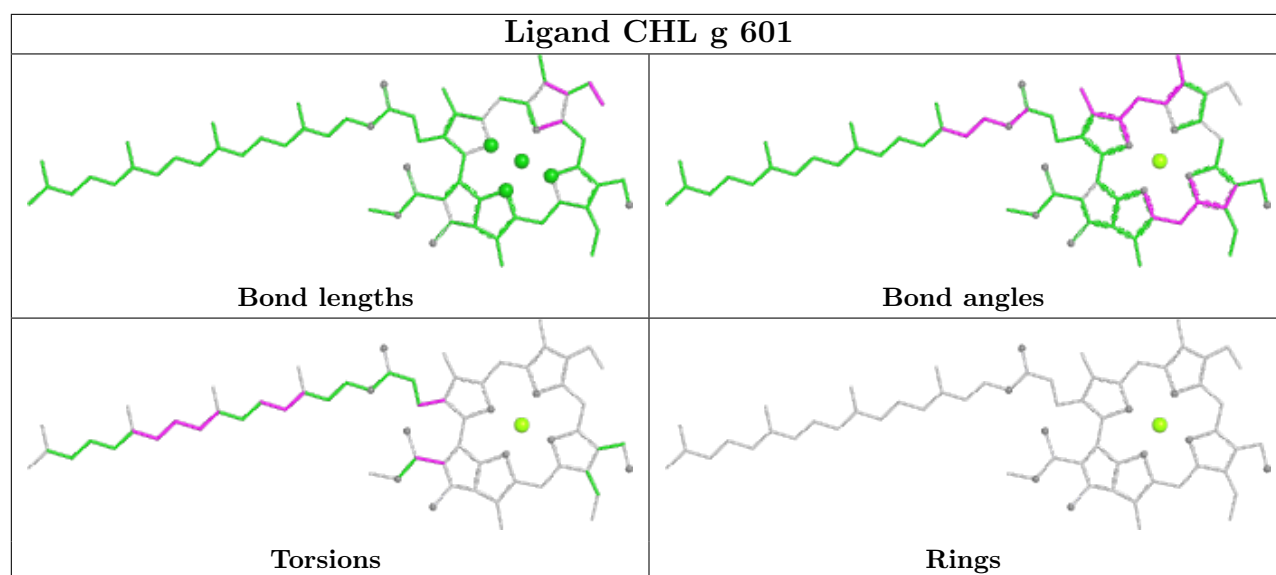


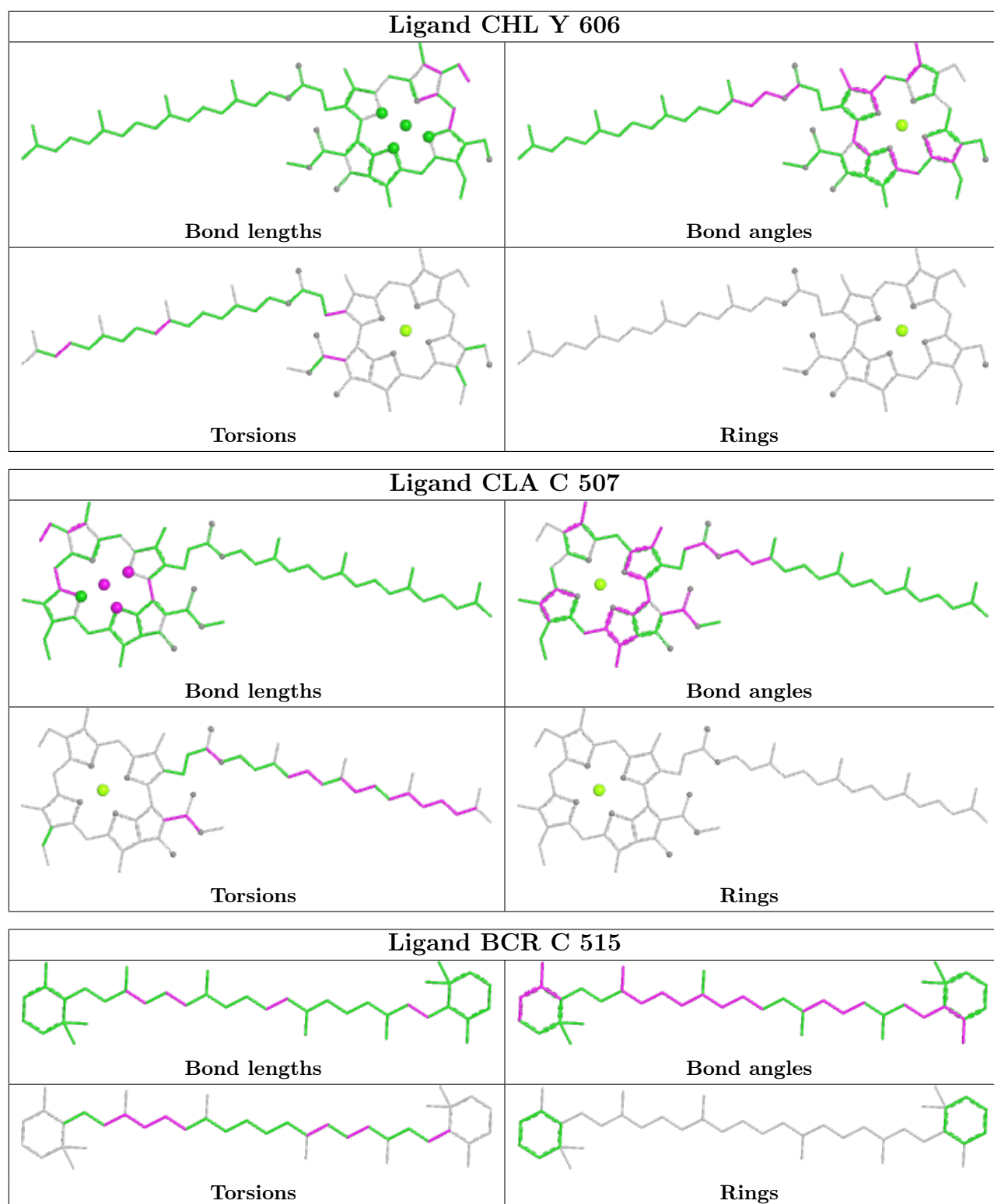


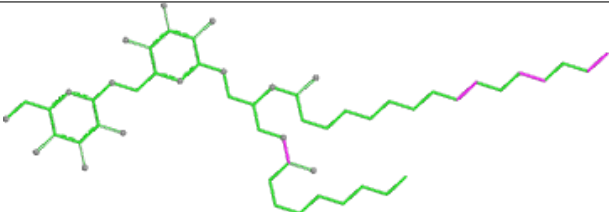
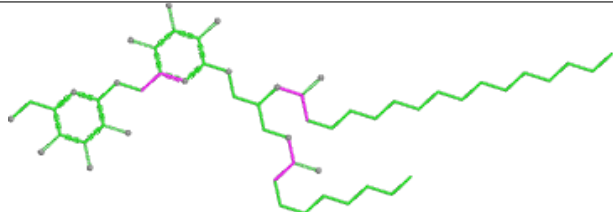
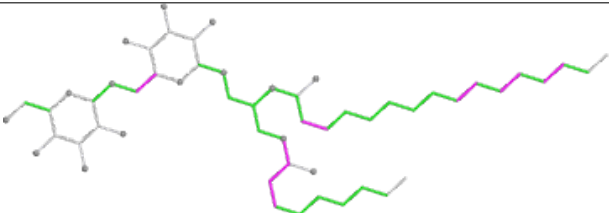
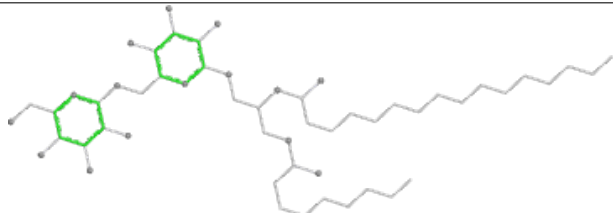


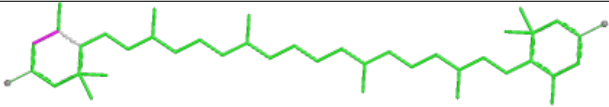
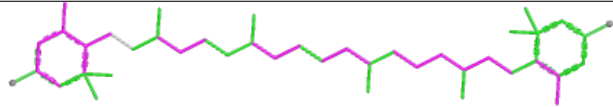
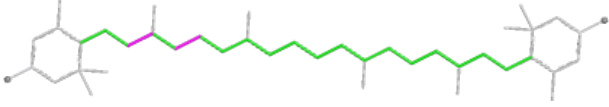
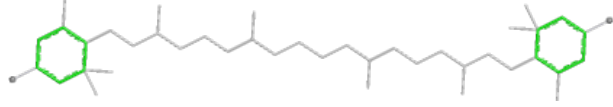


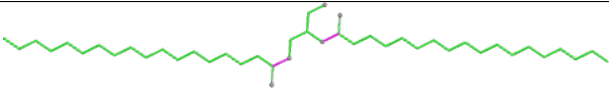
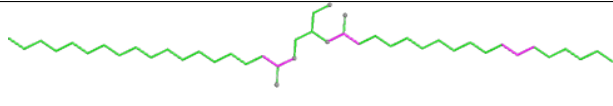
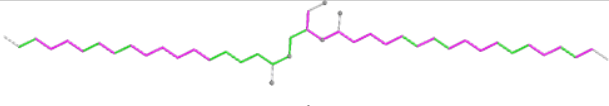
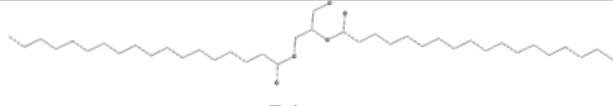


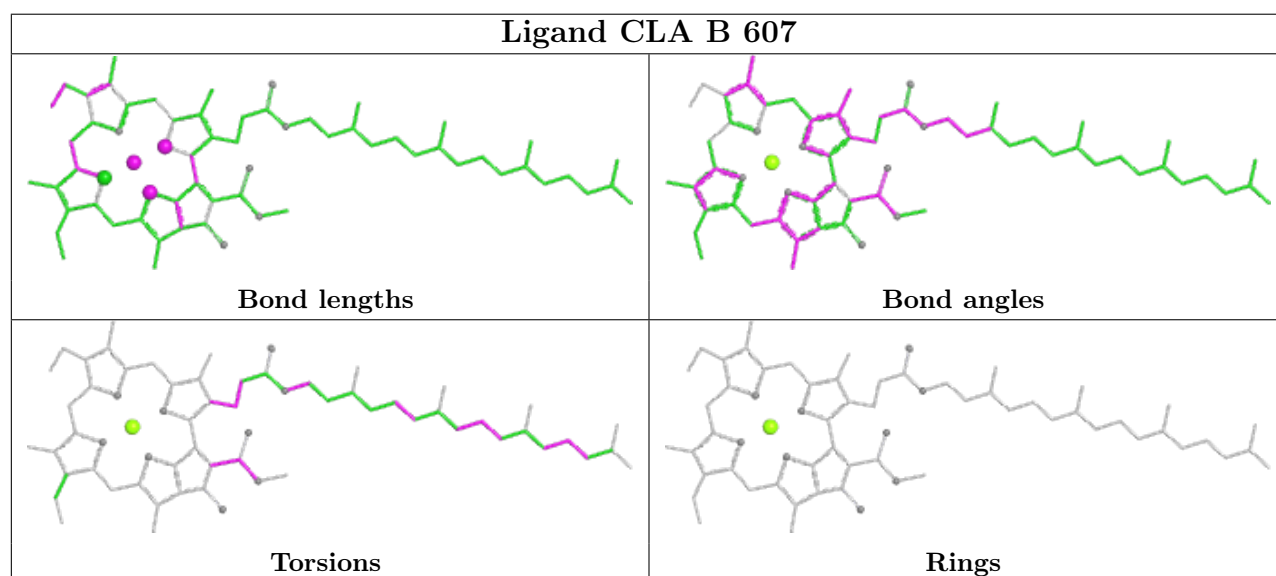
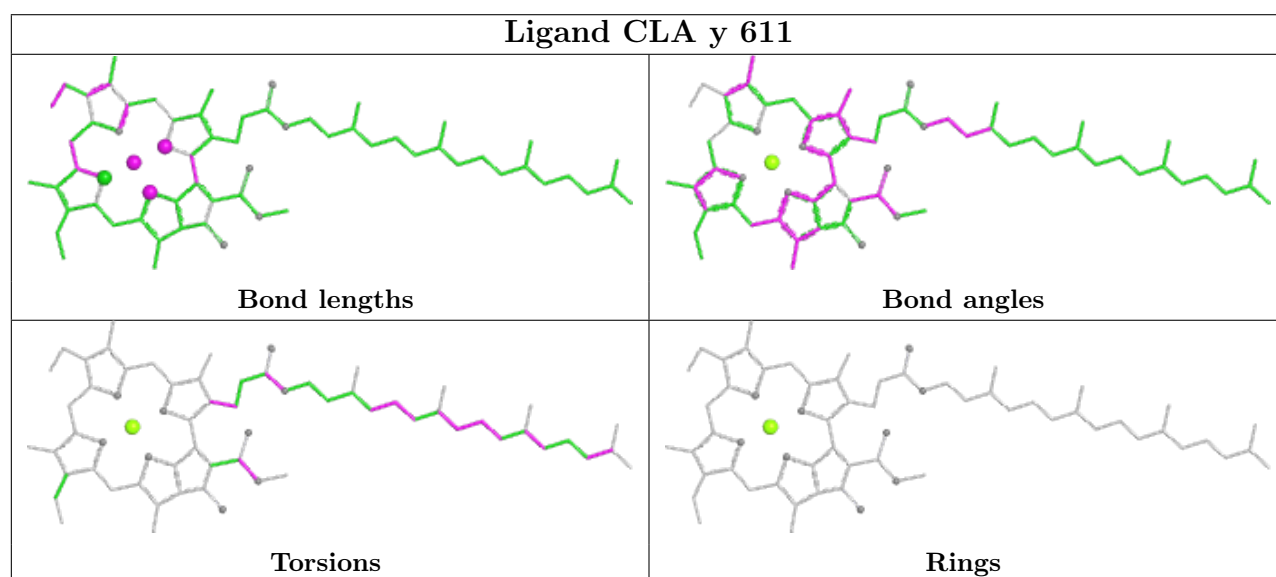
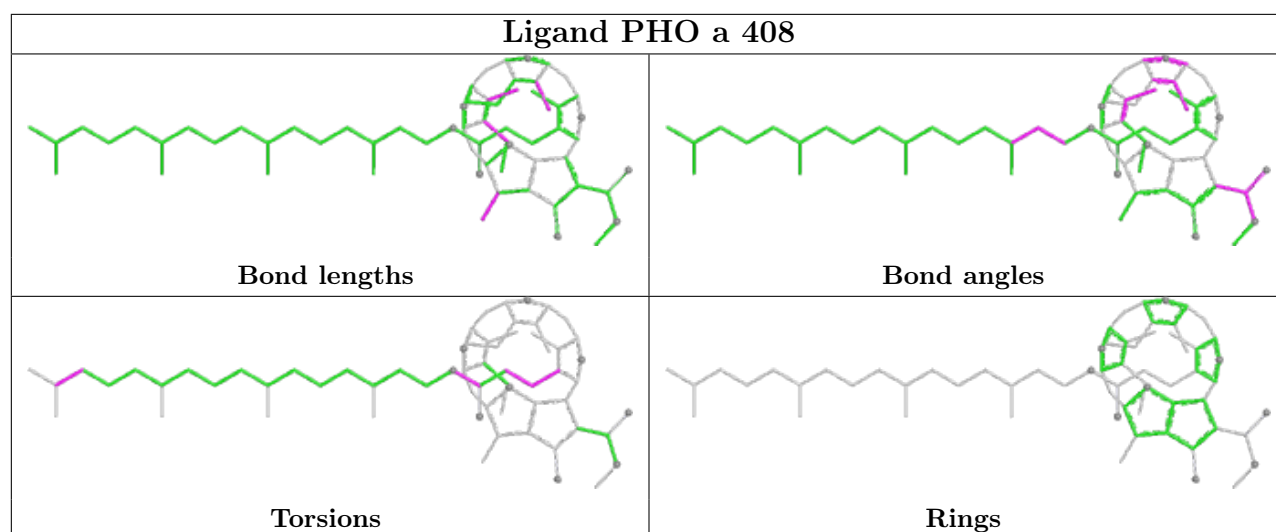


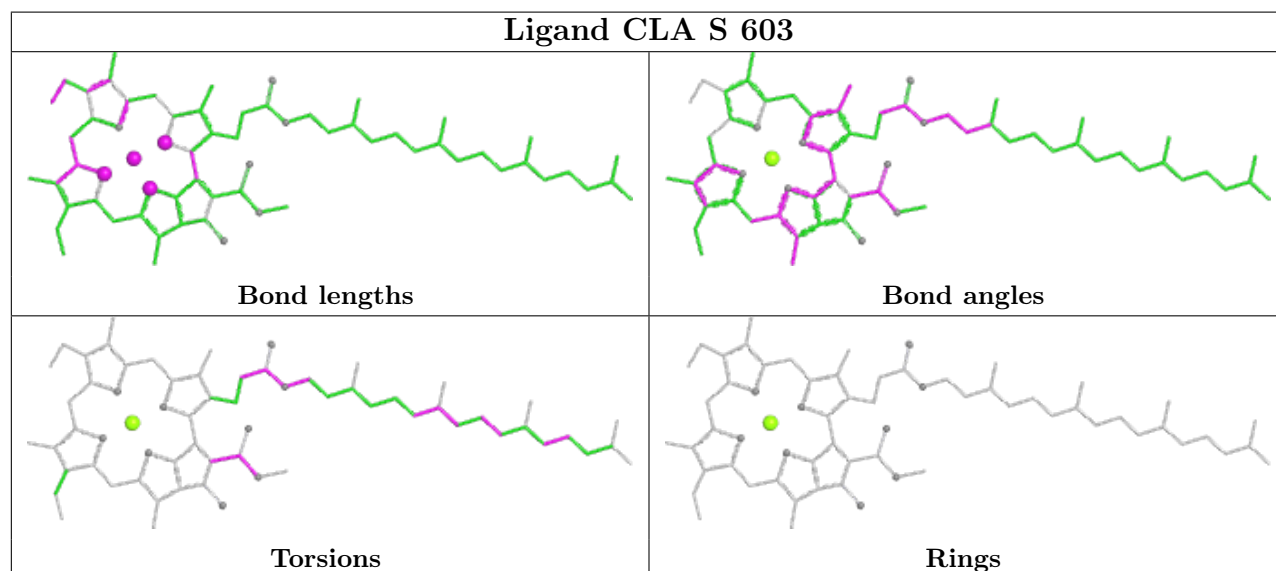
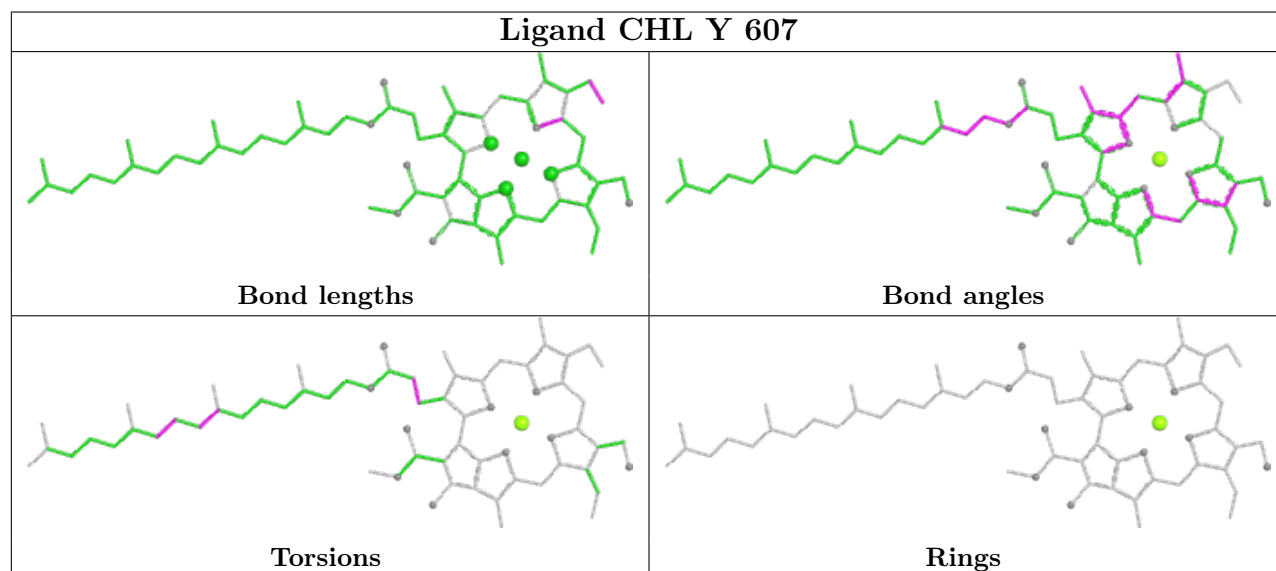
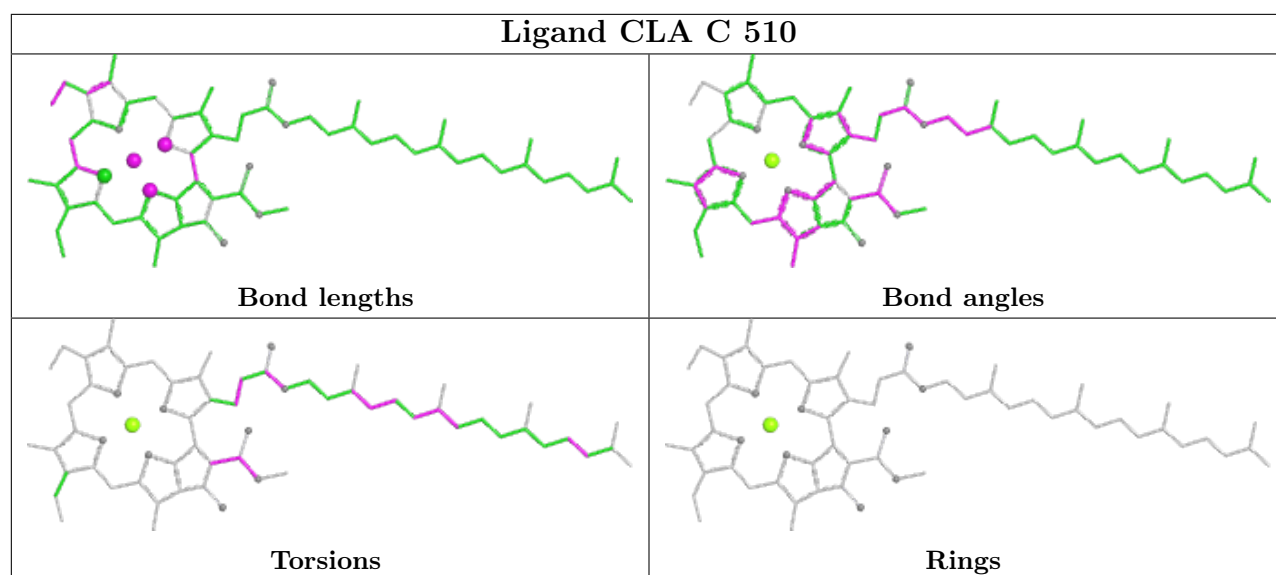


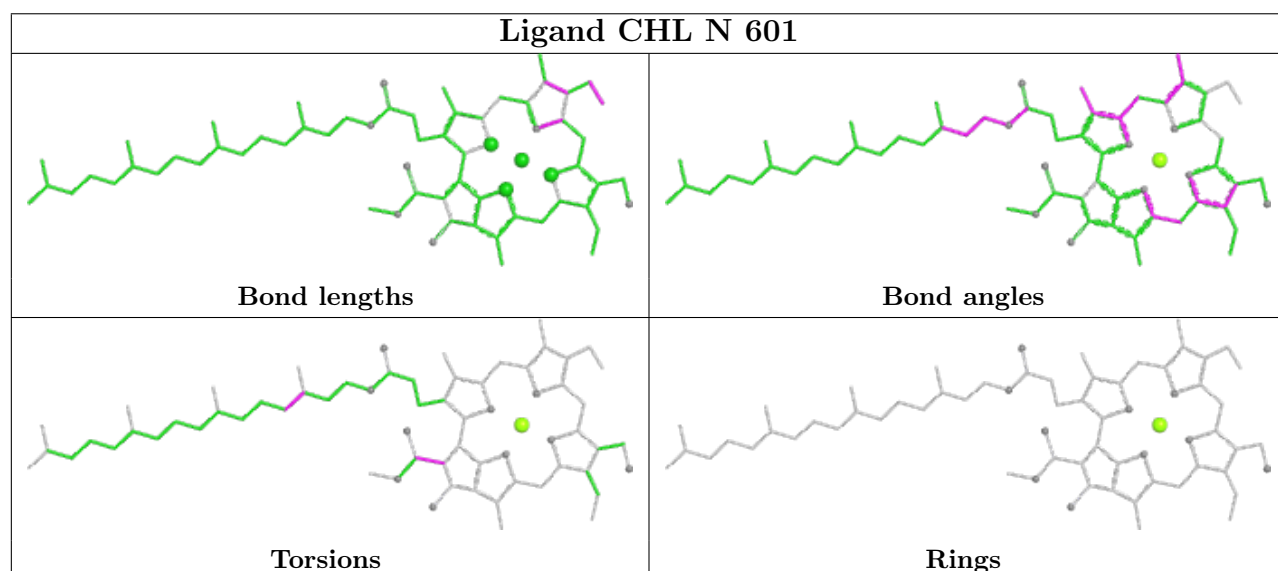
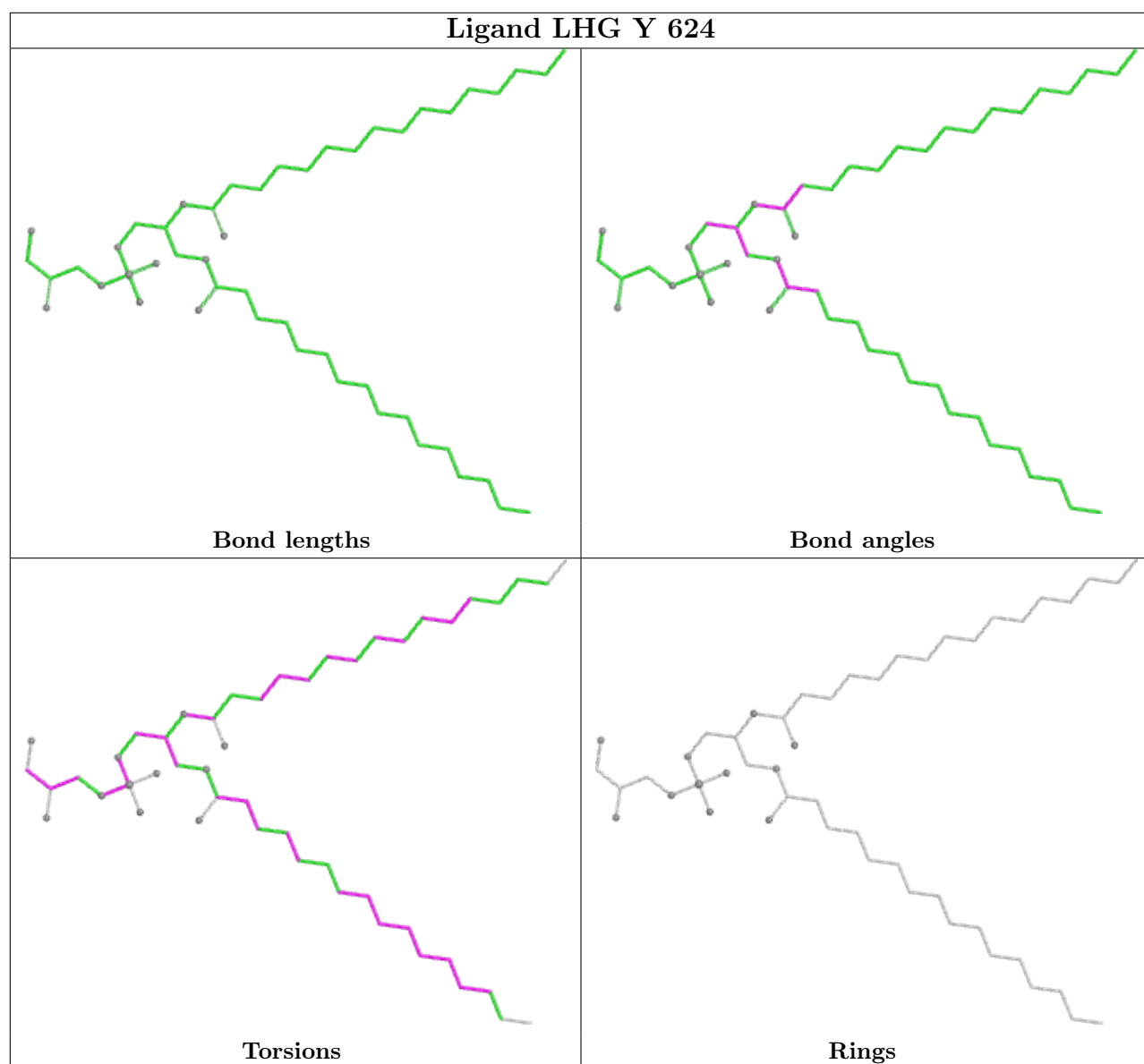
Ligand DGD c 518	
	
Bond lengths	Bond angles
	
Torsions	Rings

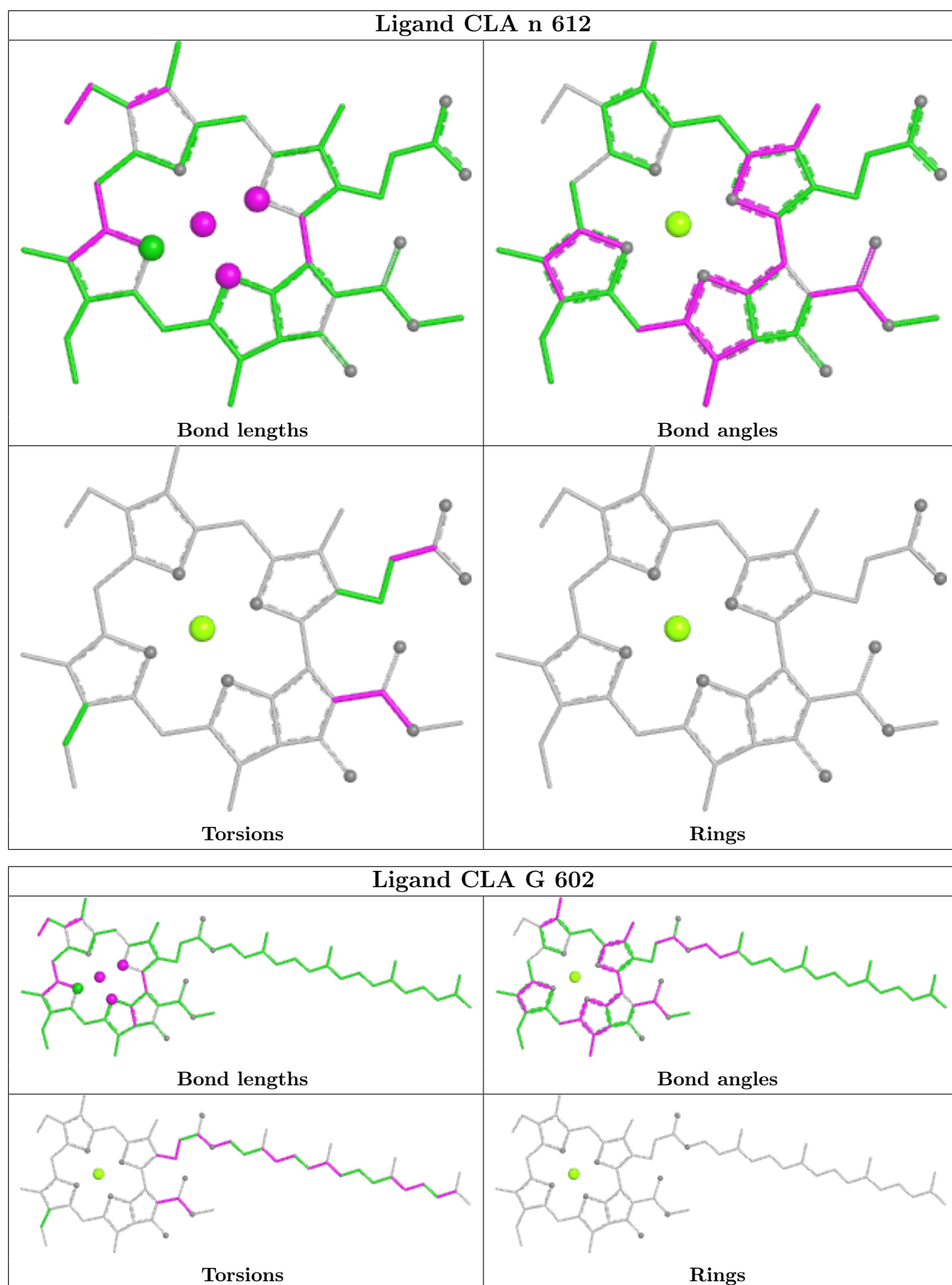
Ligand LUT g 620	
	
Bond lengths	Bond angles
	
Torsions	Rings

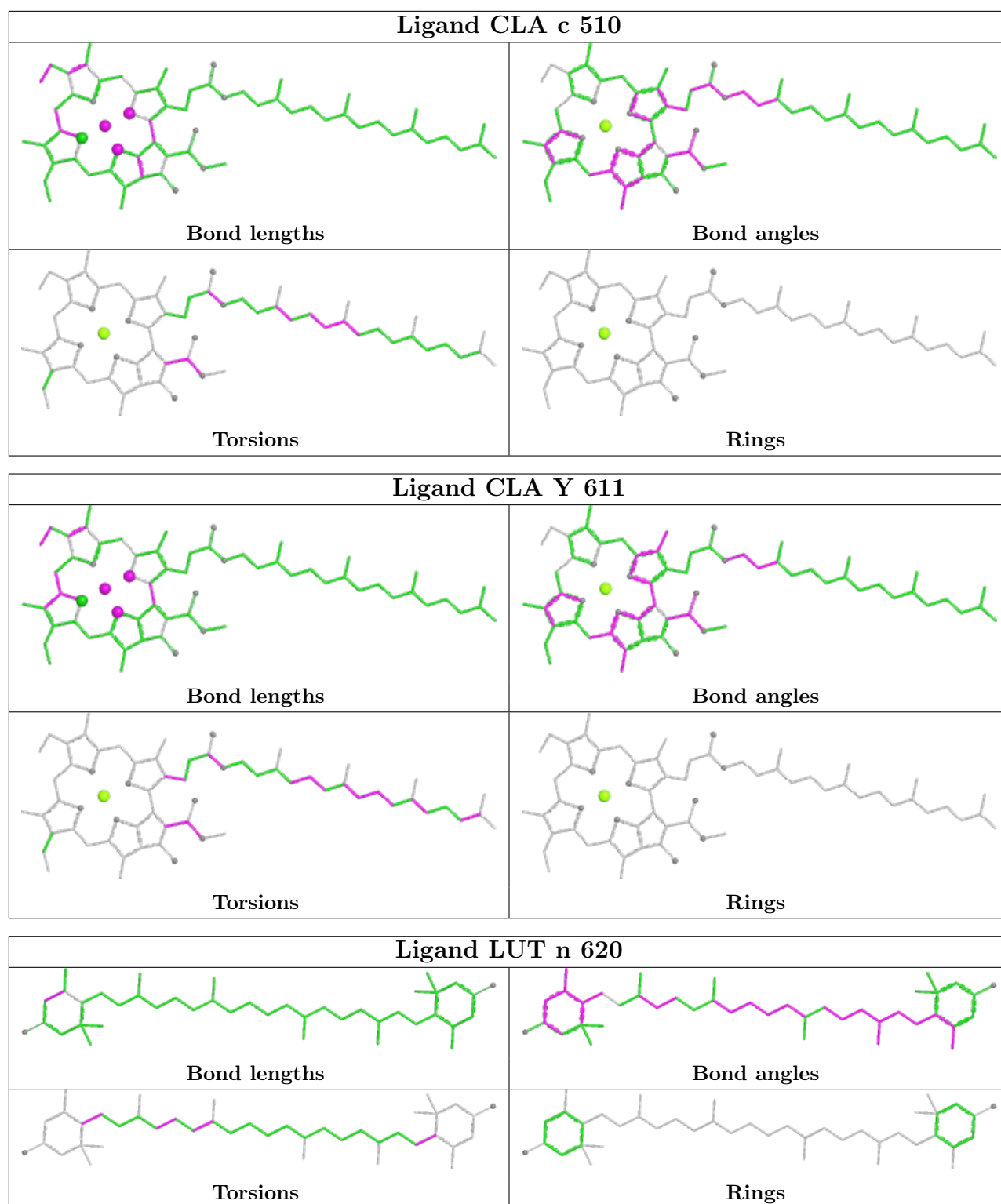
Ligand DGA b 623	
	
Bond lengths	Bond angles
	
Torsions	Rings

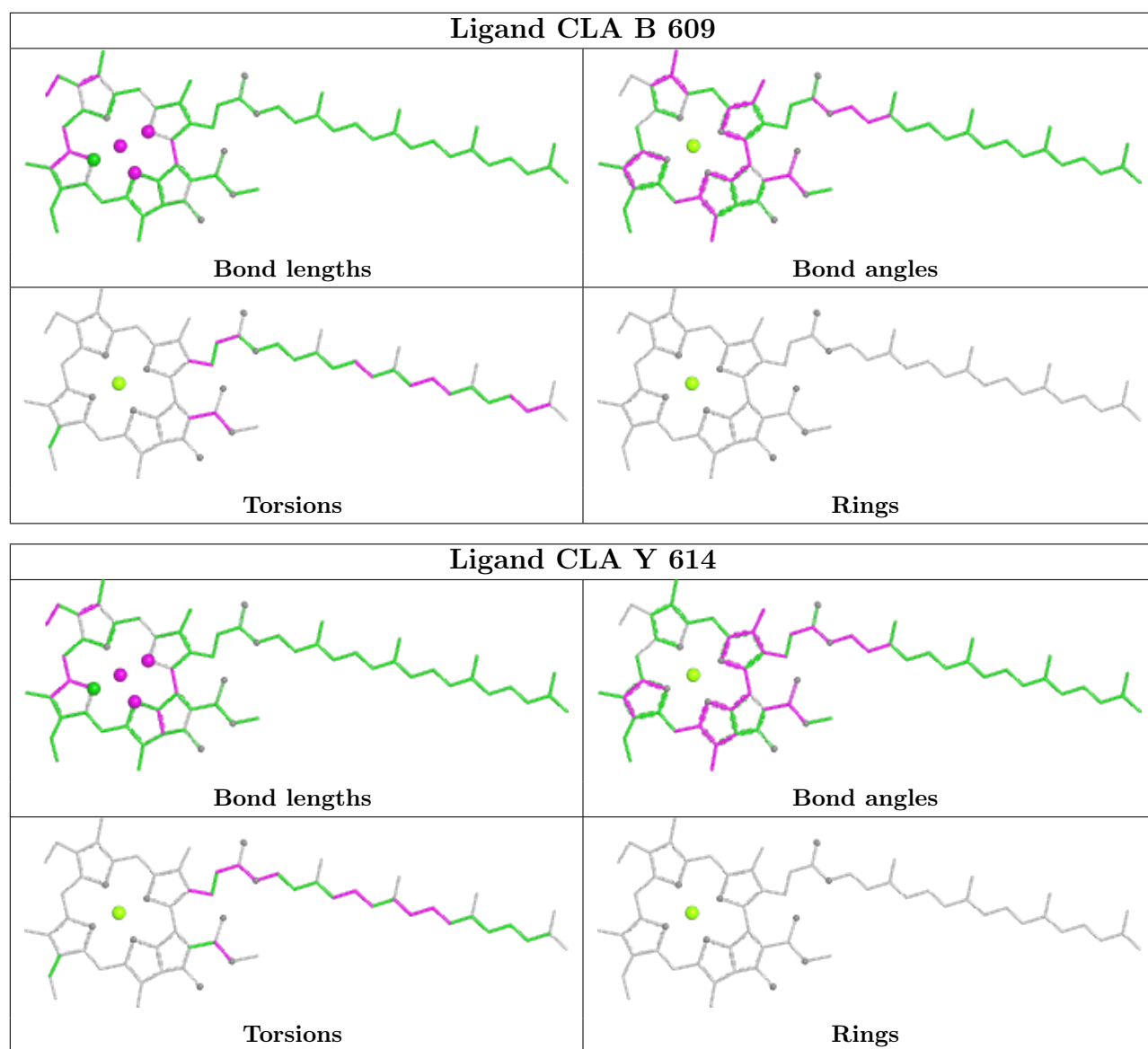


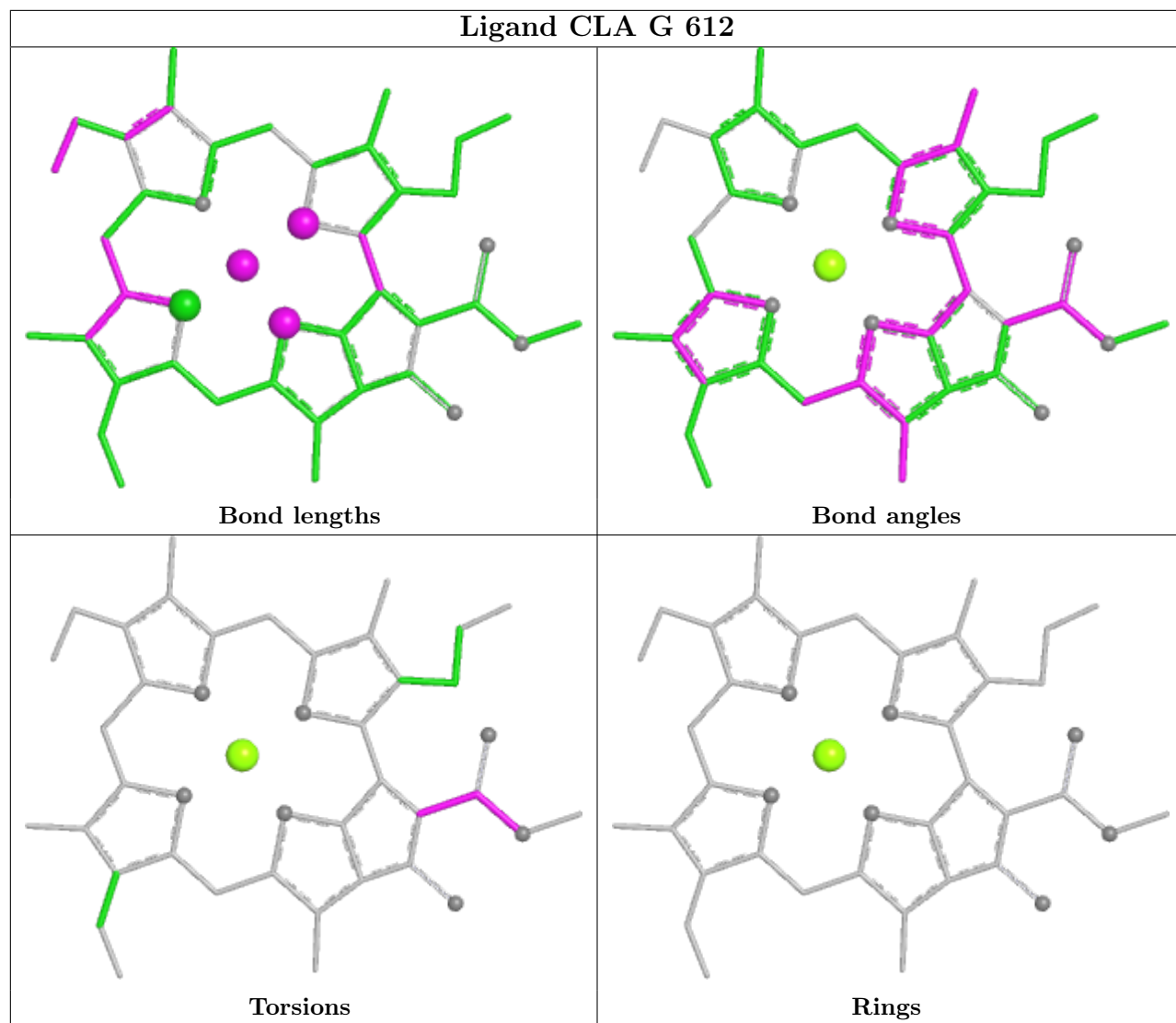


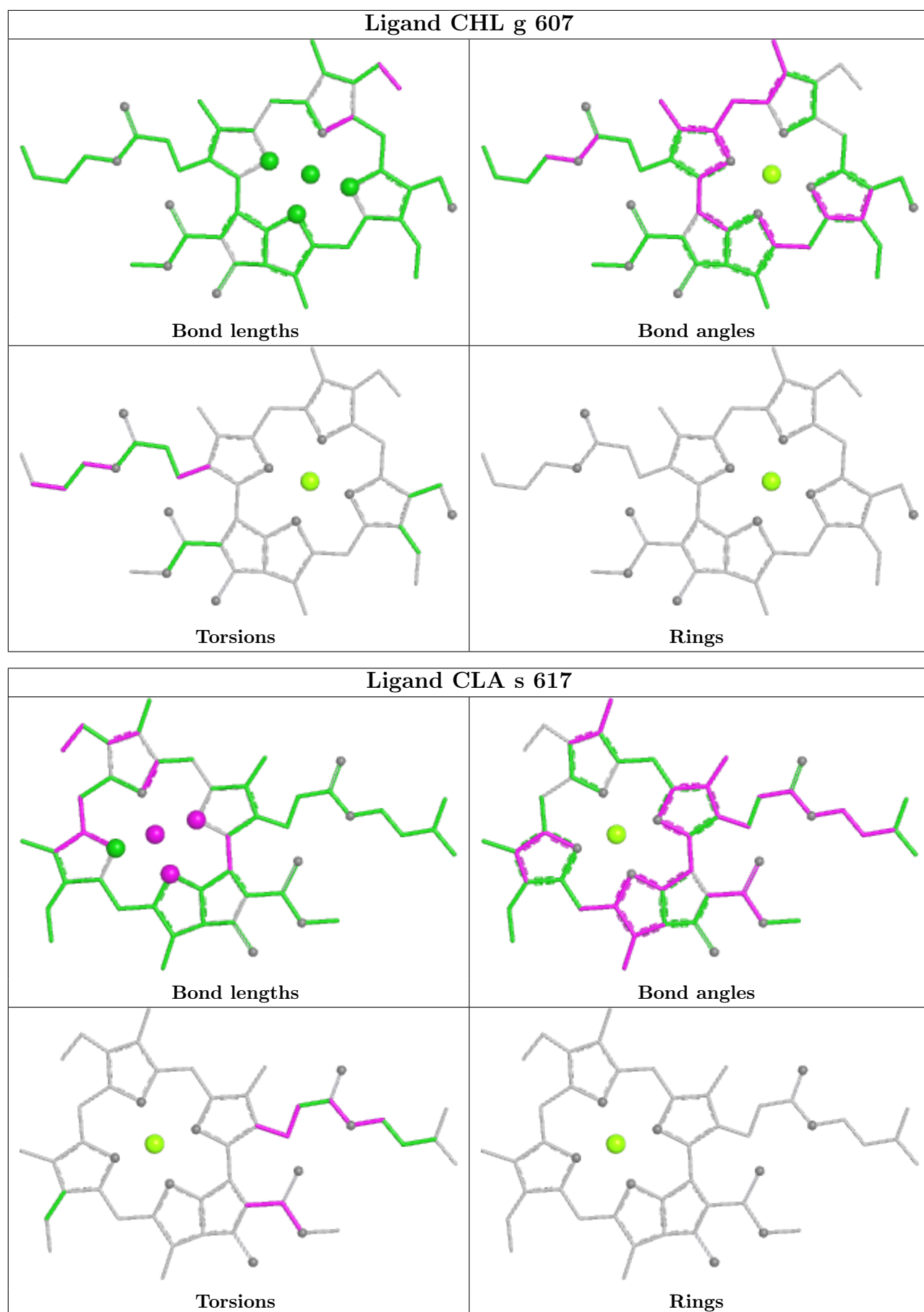




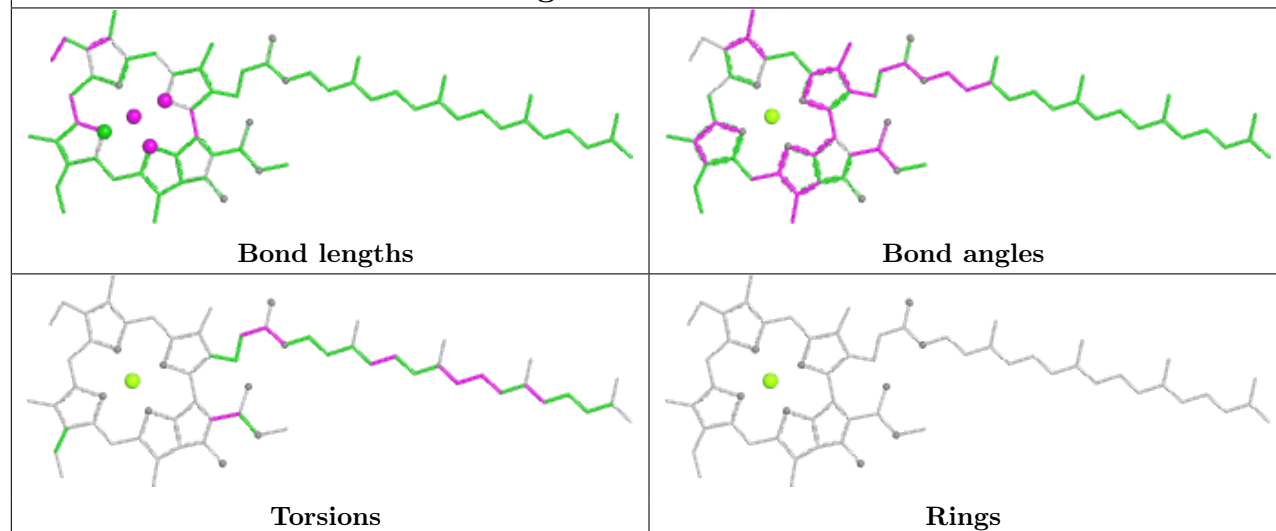




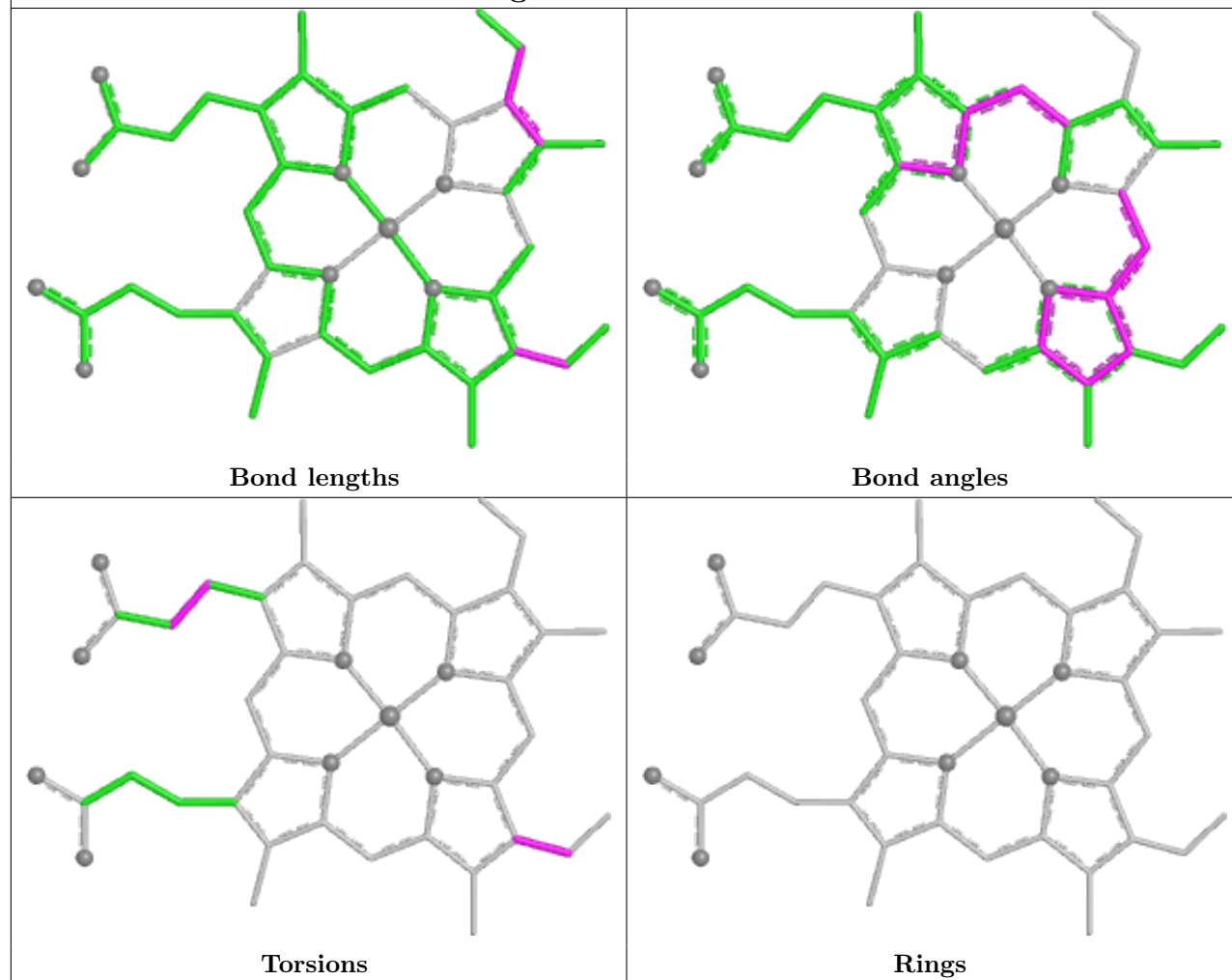


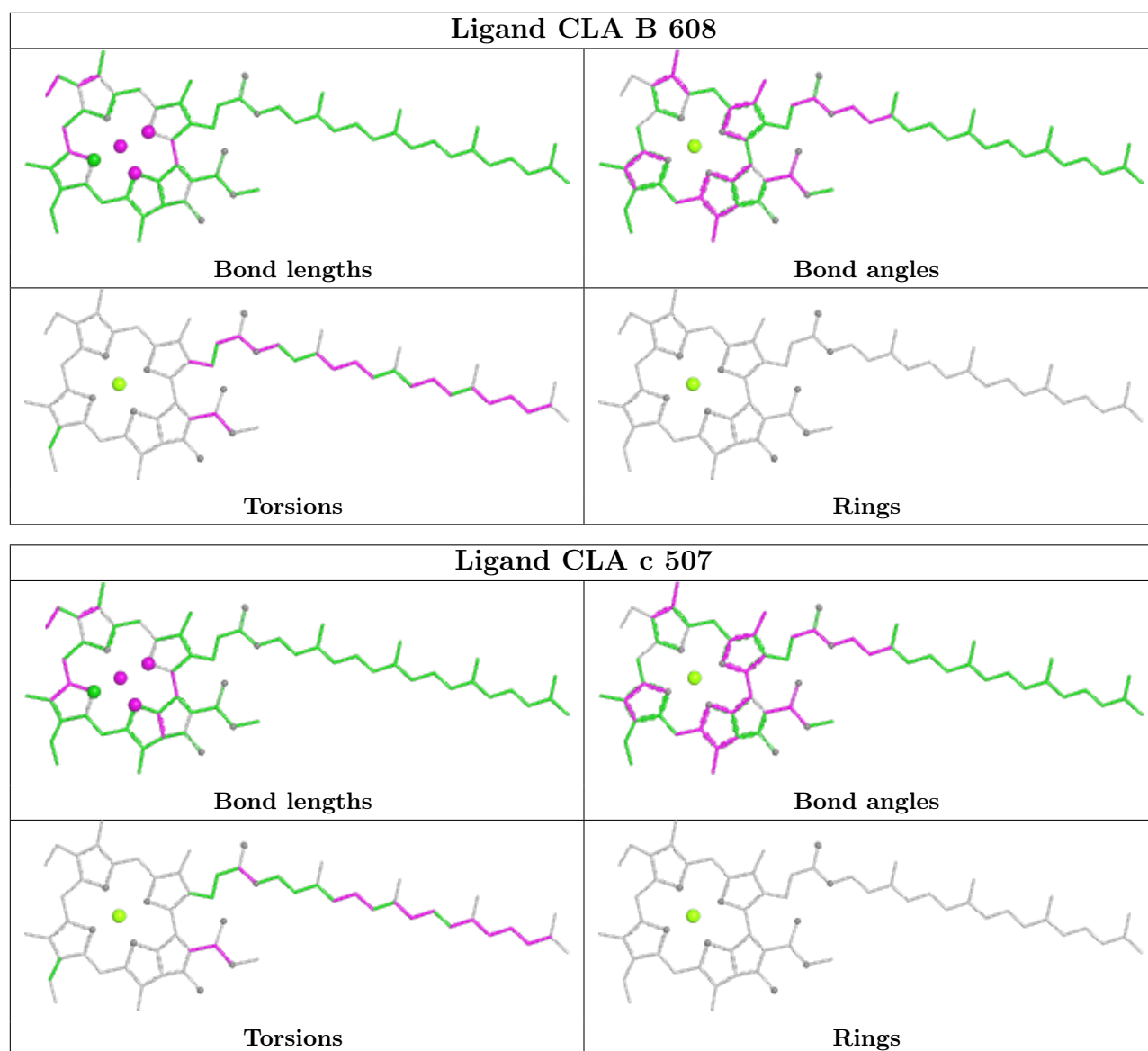


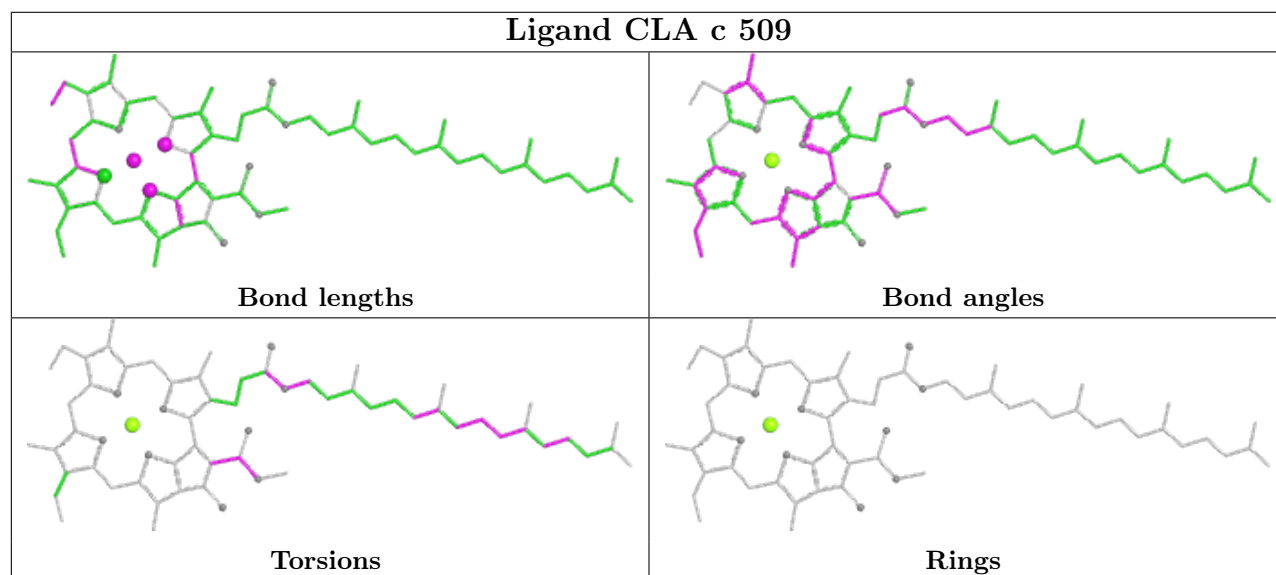
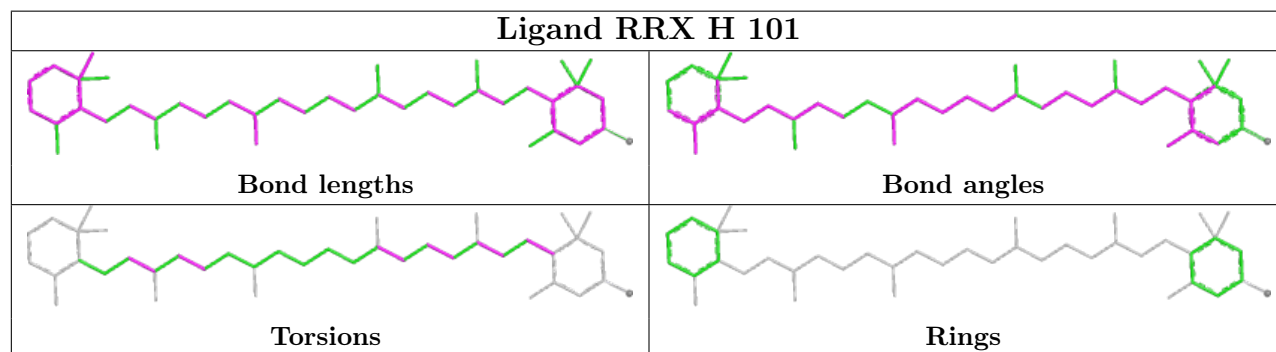
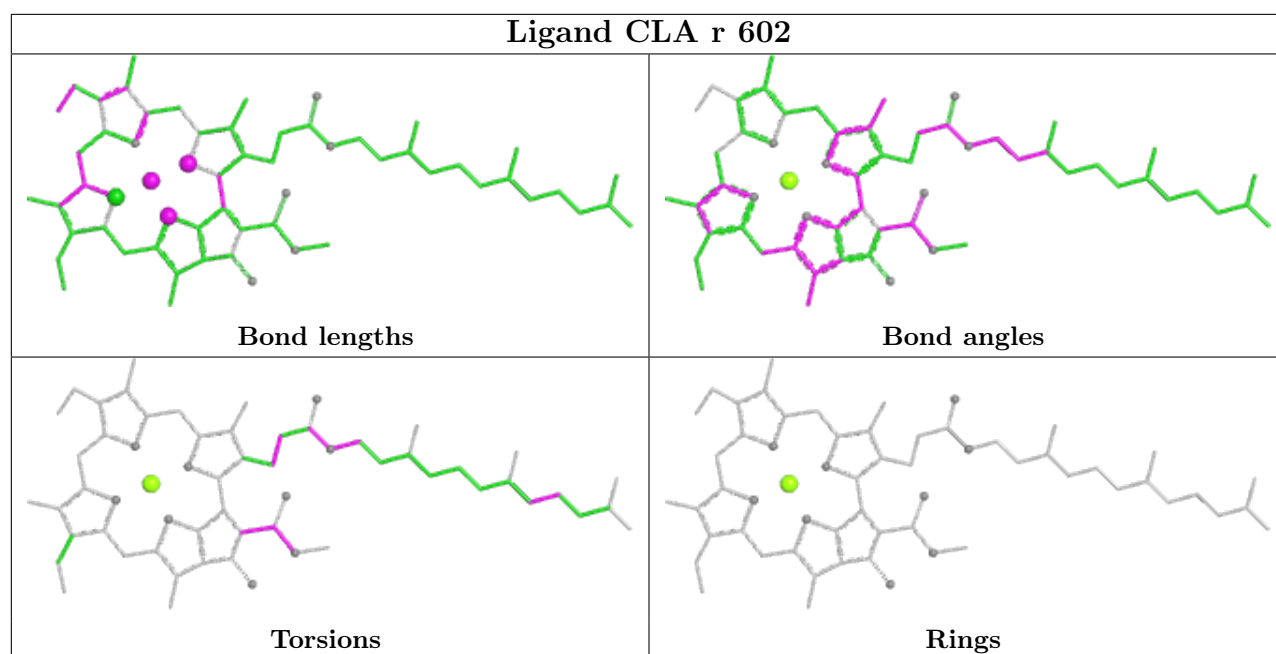
Ligand CLA c 508

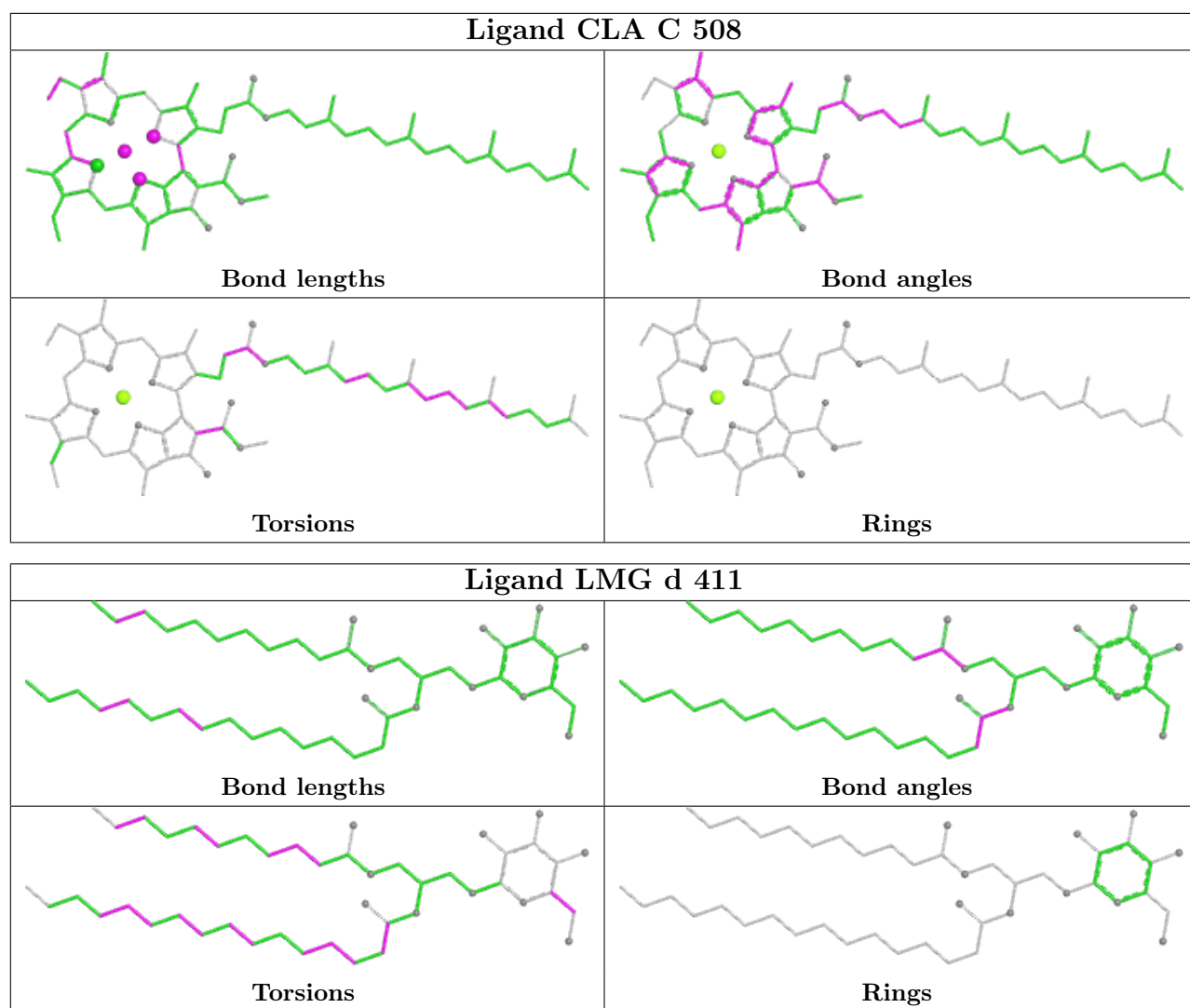


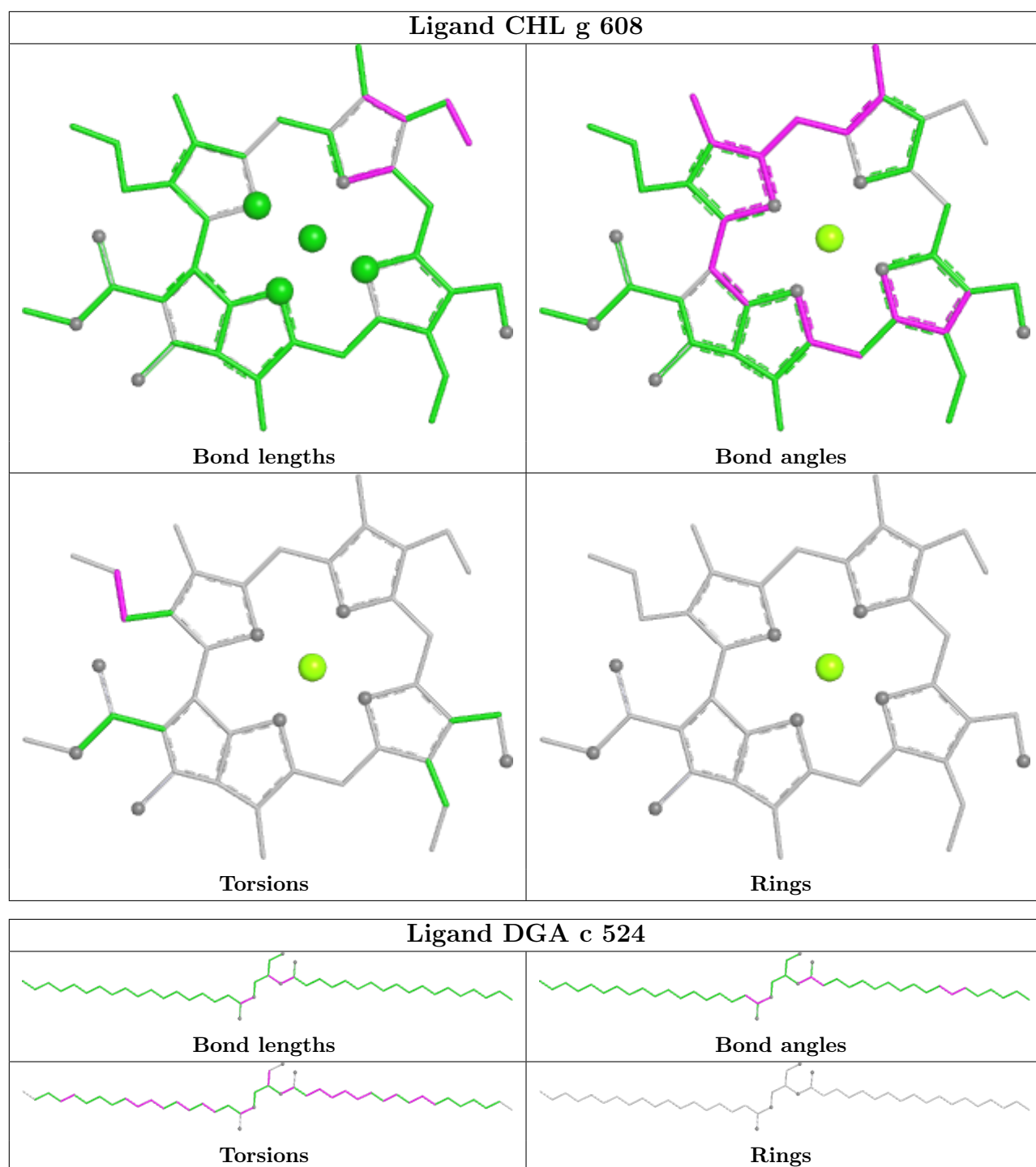
Ligand HEM F 101

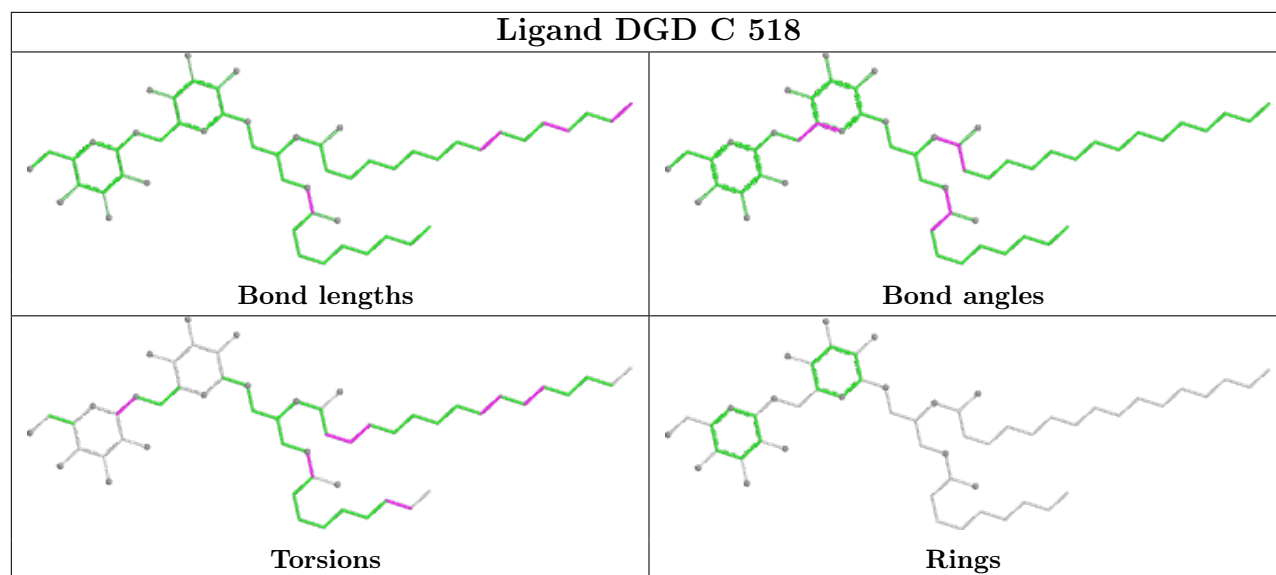
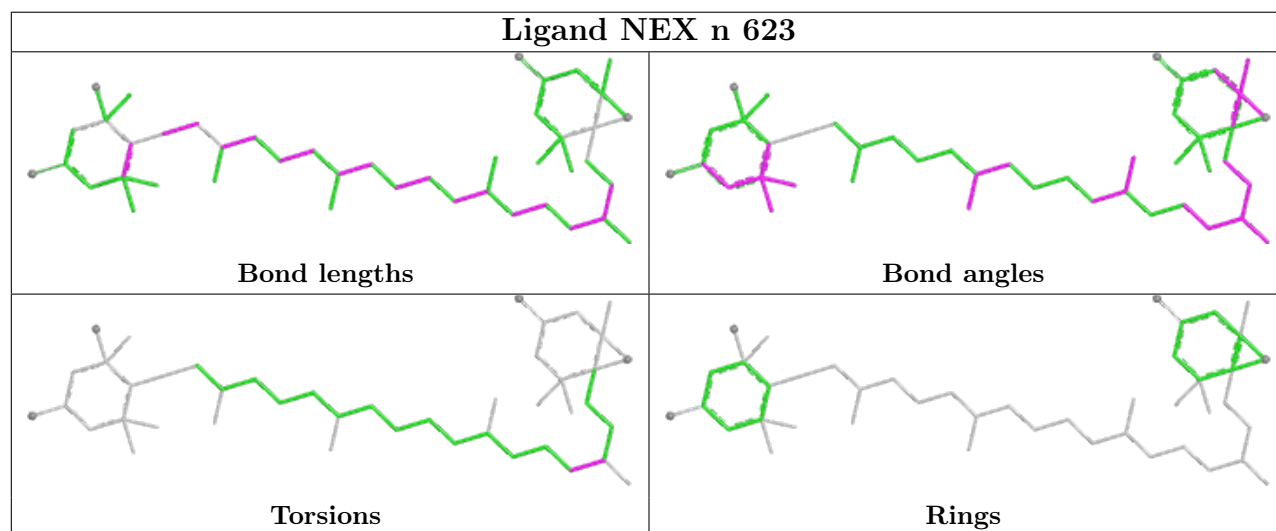
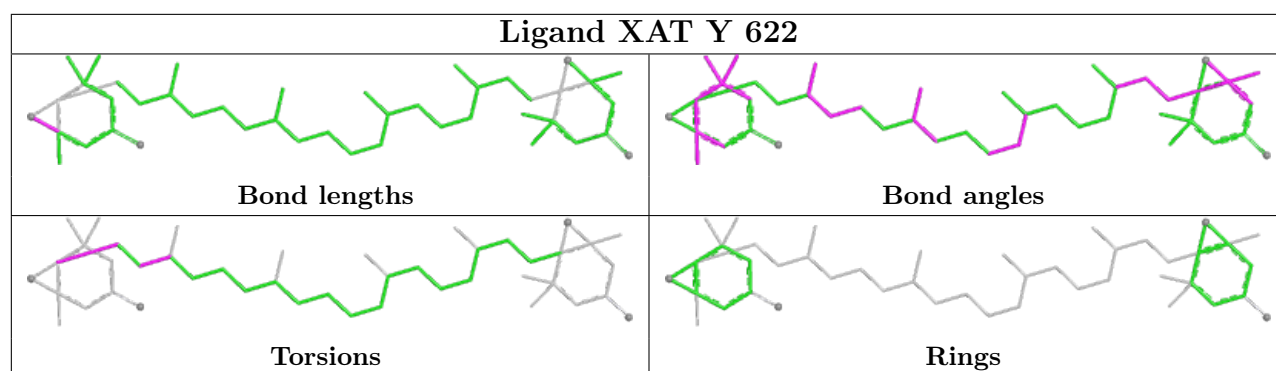


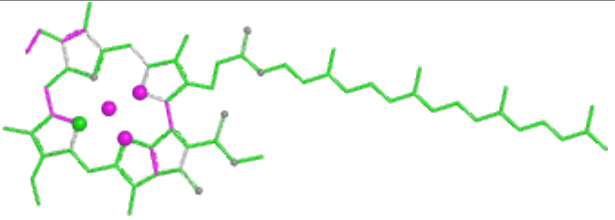
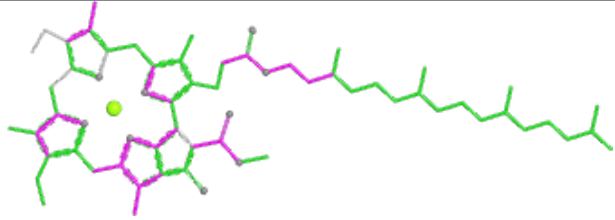
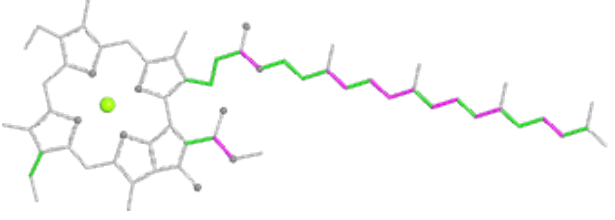
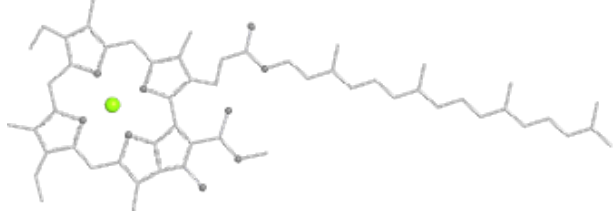
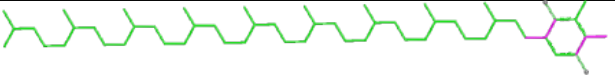
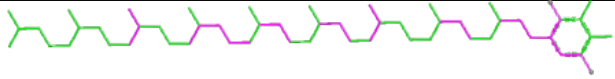
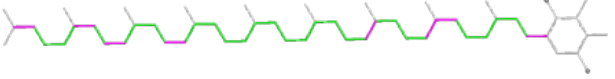
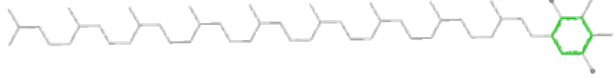
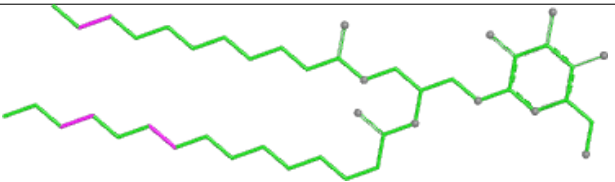
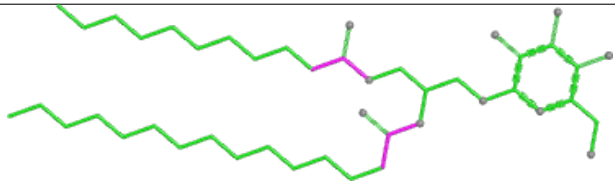
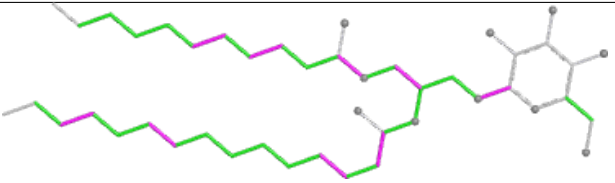
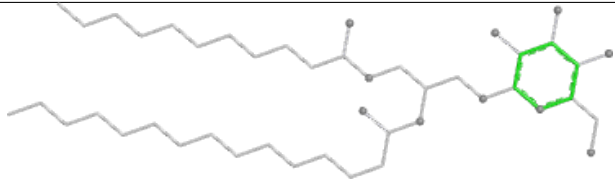


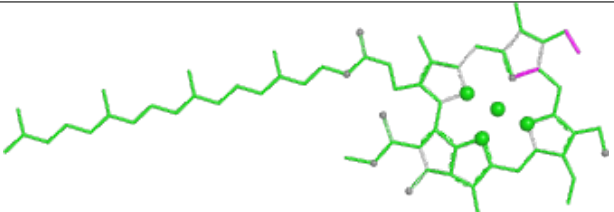
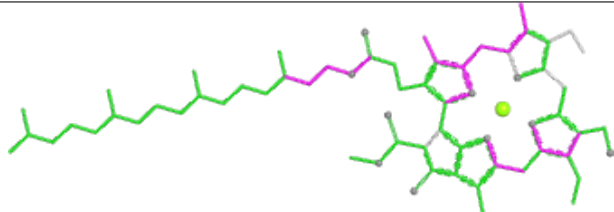
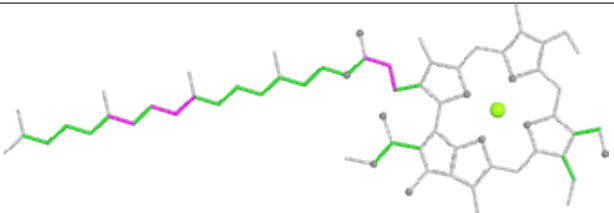
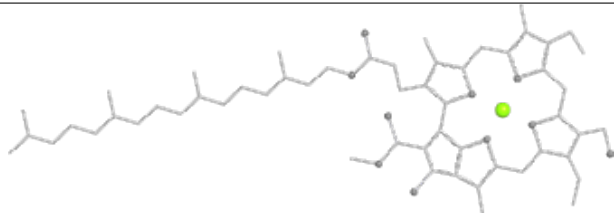


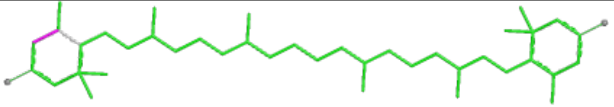
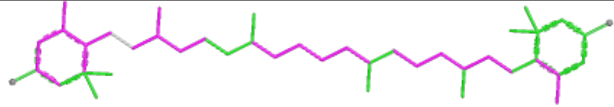
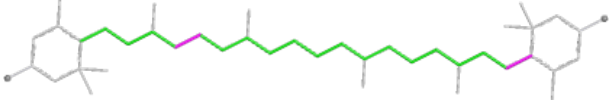
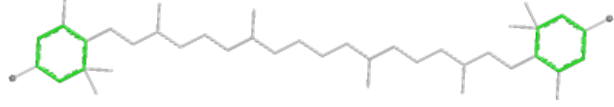


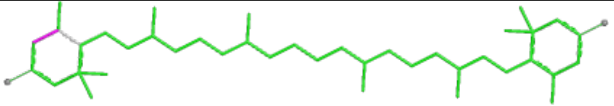
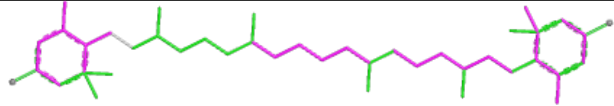
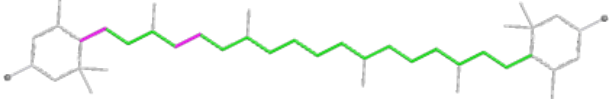
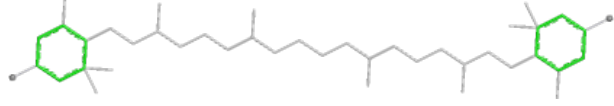


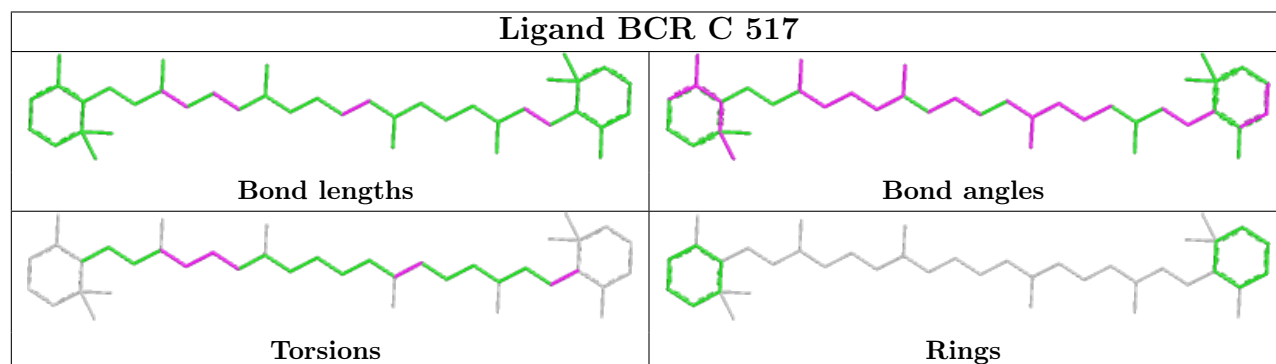
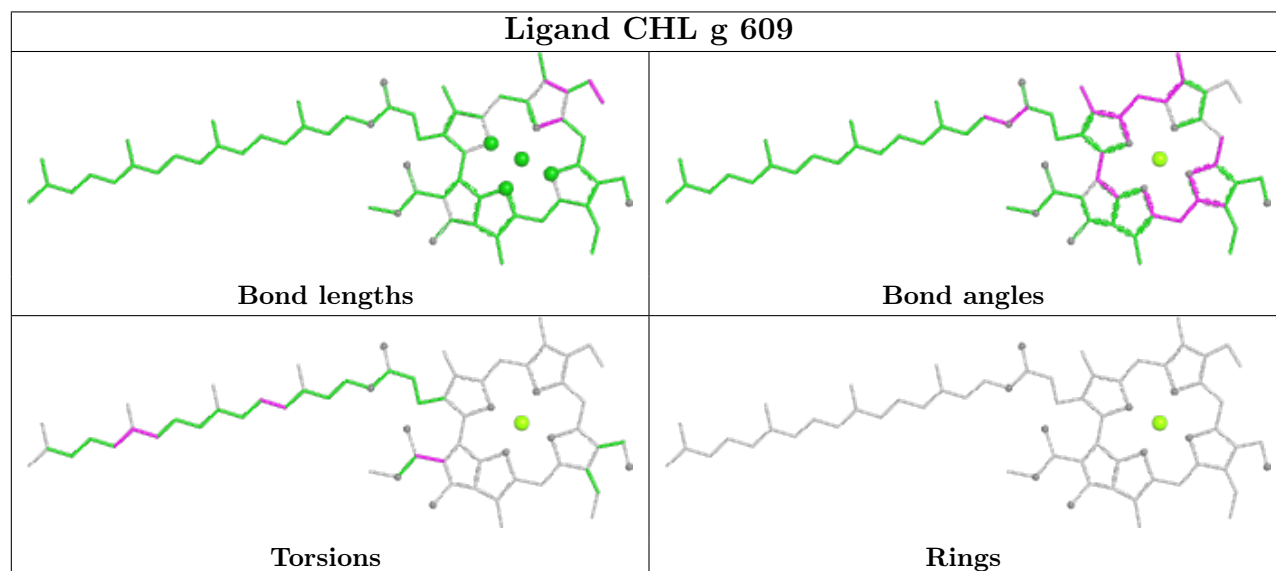
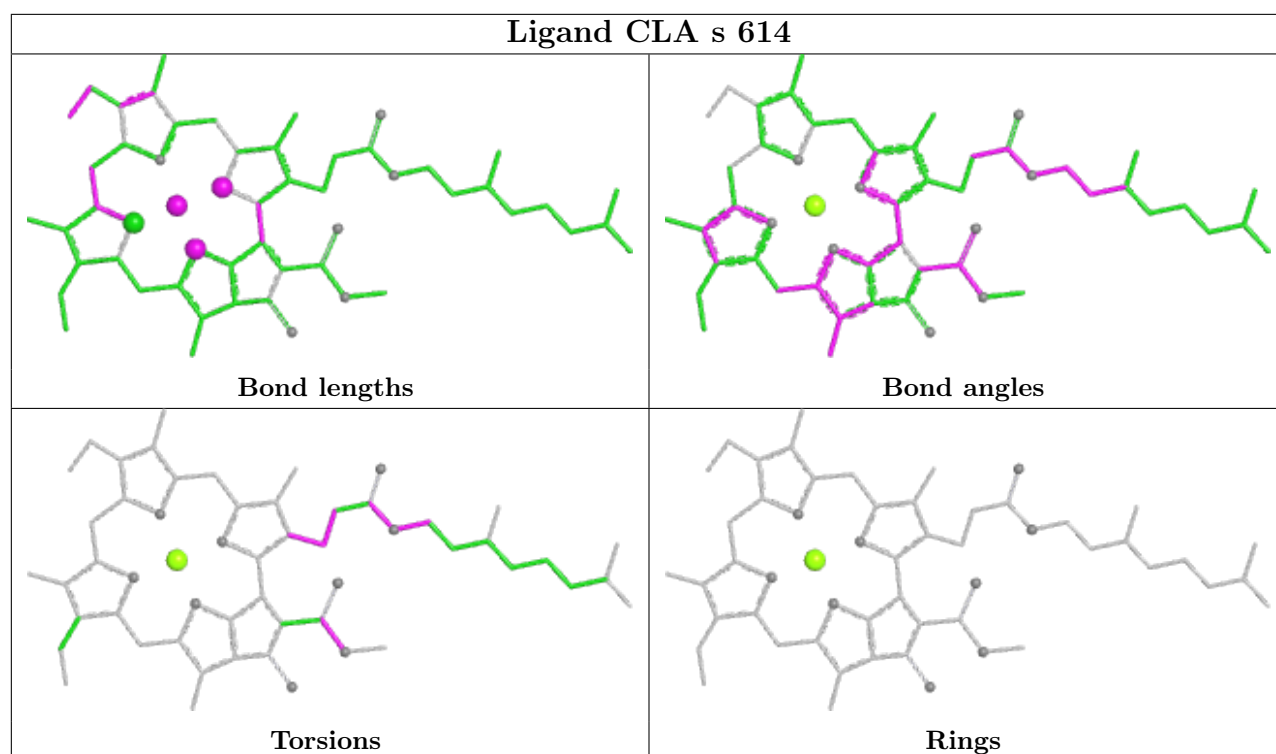


Ligand CLA b 606	
 <p>Bond lengths</p>	 <p>Bond angles</p>
 <p>Torsions</p>	 <p>Rings</p>
Ligand PL9 d 405	
 <p>Bond lengths</p>	 <p>Bond angles</p>
 <p>Torsions</p>	 <p>Rings</p>
Ligand LMG J 101	
 <p>Bond lengths</p>	 <p>Bond angles</p>
 <p>Torsions</p>	 <p>Rings</p>

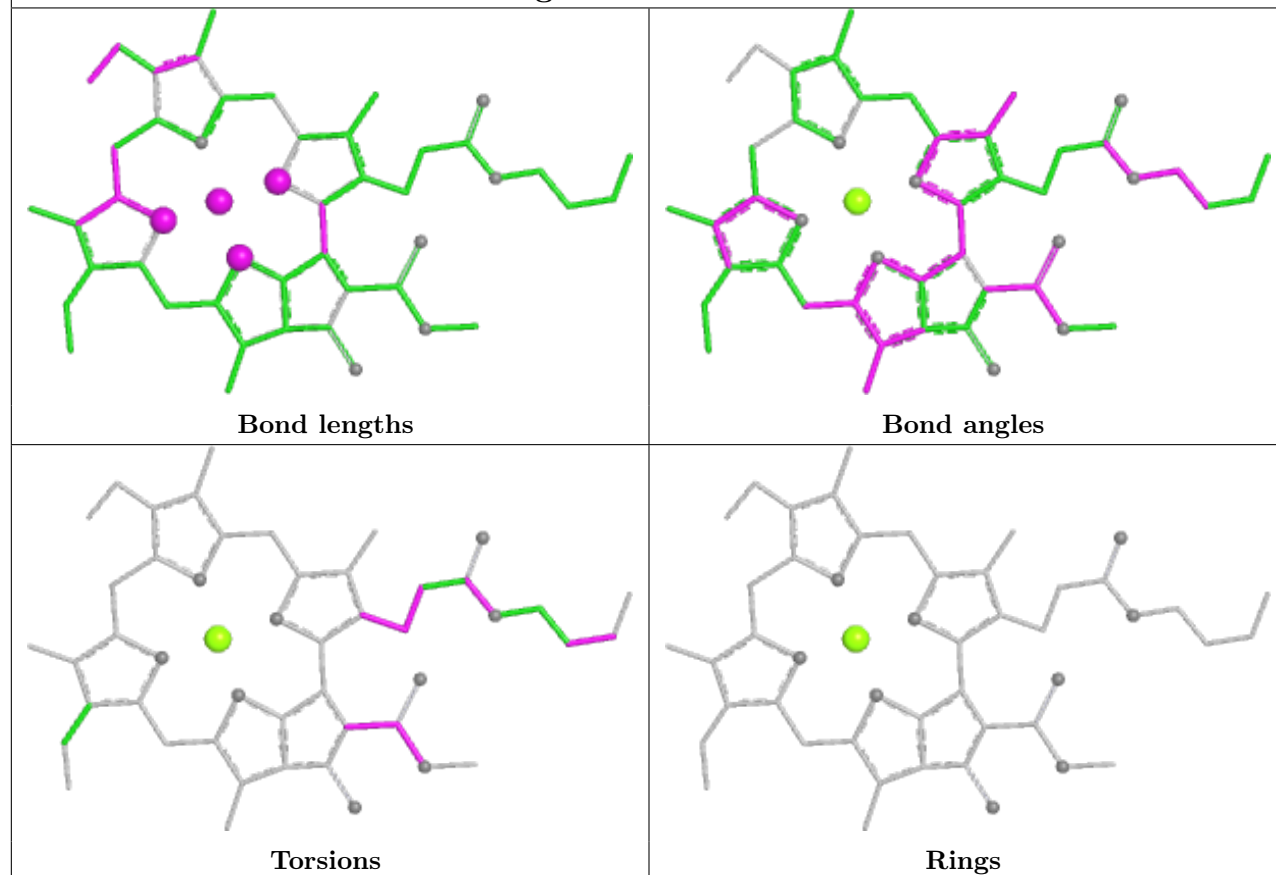
Ligand CHL N 607	
	
Bond lengths	Bond angles
	
Torsions	Rings

Ligand LUT N 620	
	
Bond lengths	Bond angles
	
Torsions	Rings

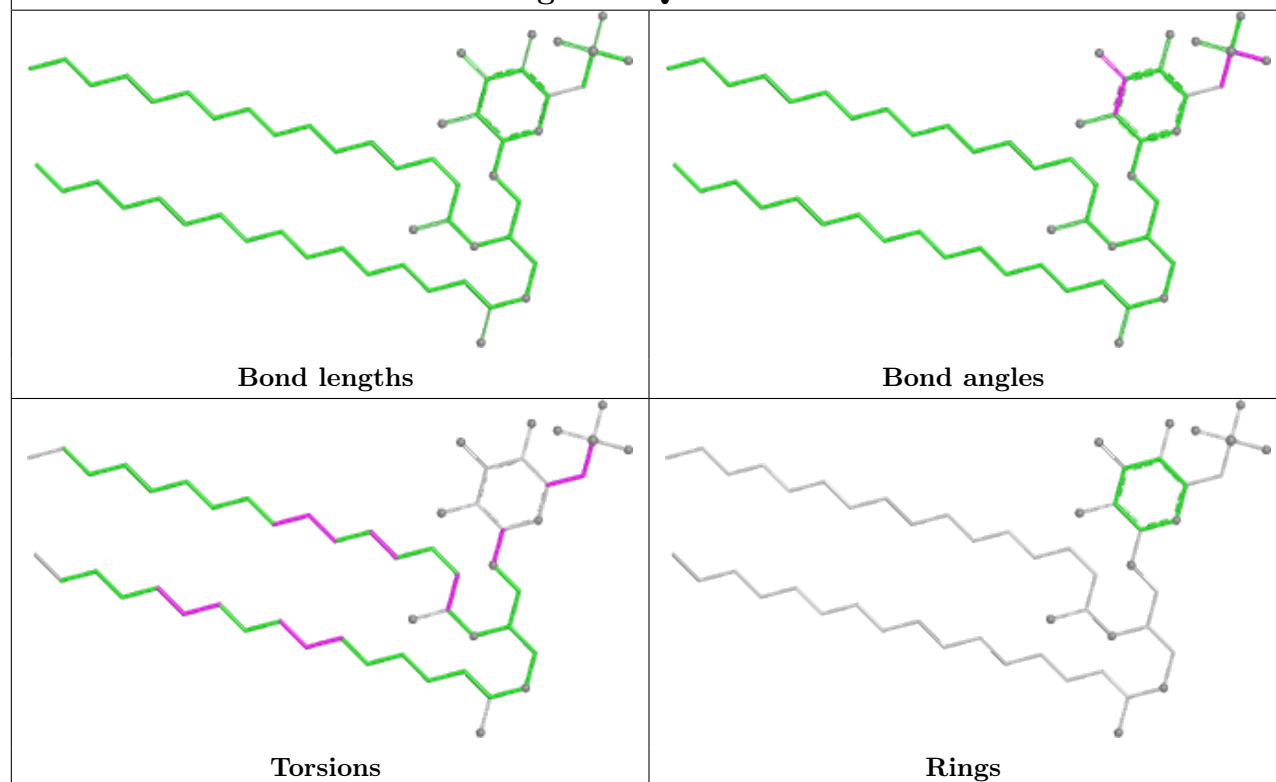
Ligand LUT N 621	
	
Bond lengths	Bond angles
	
Torsions	Rings

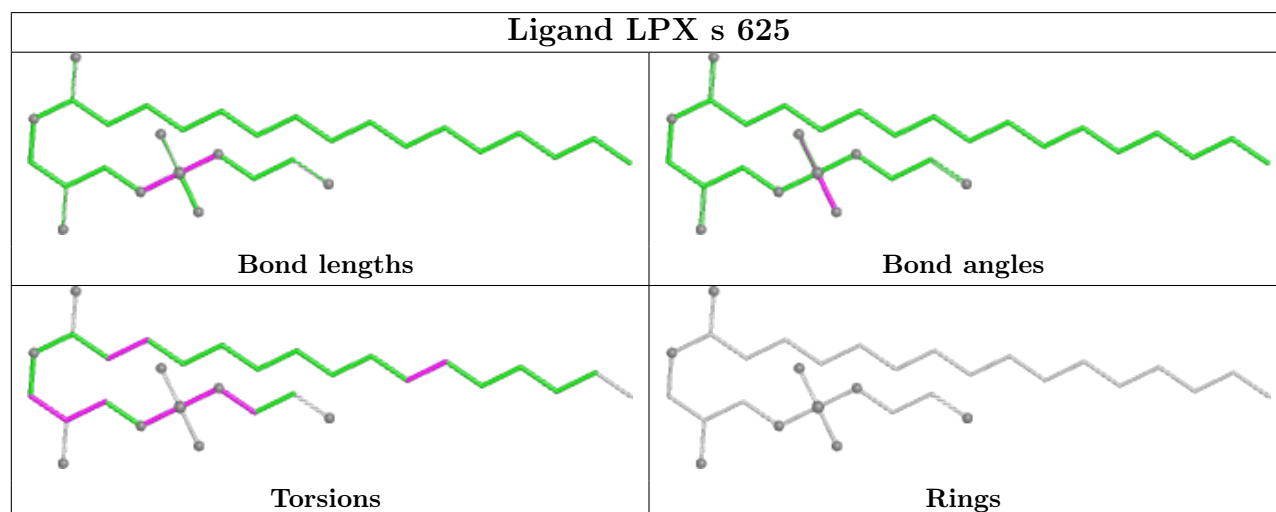
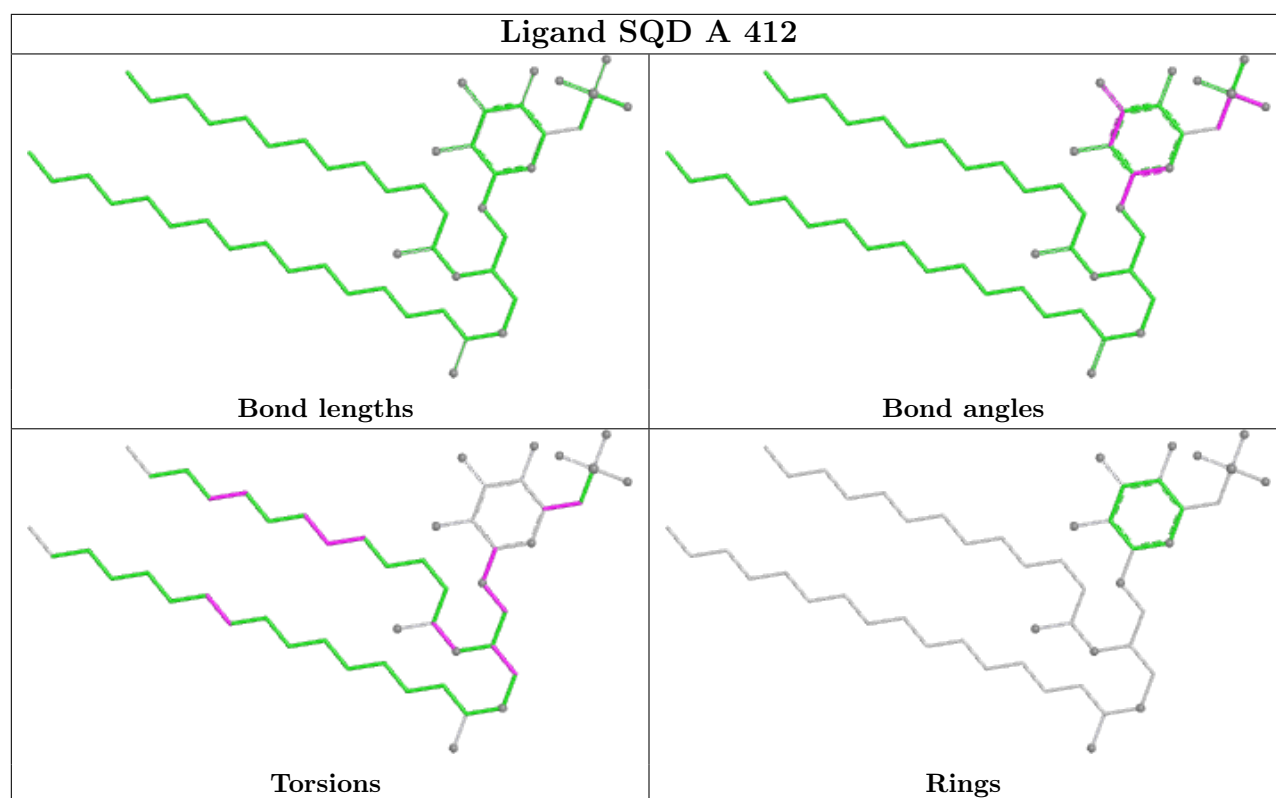


Ligand CLA n 611

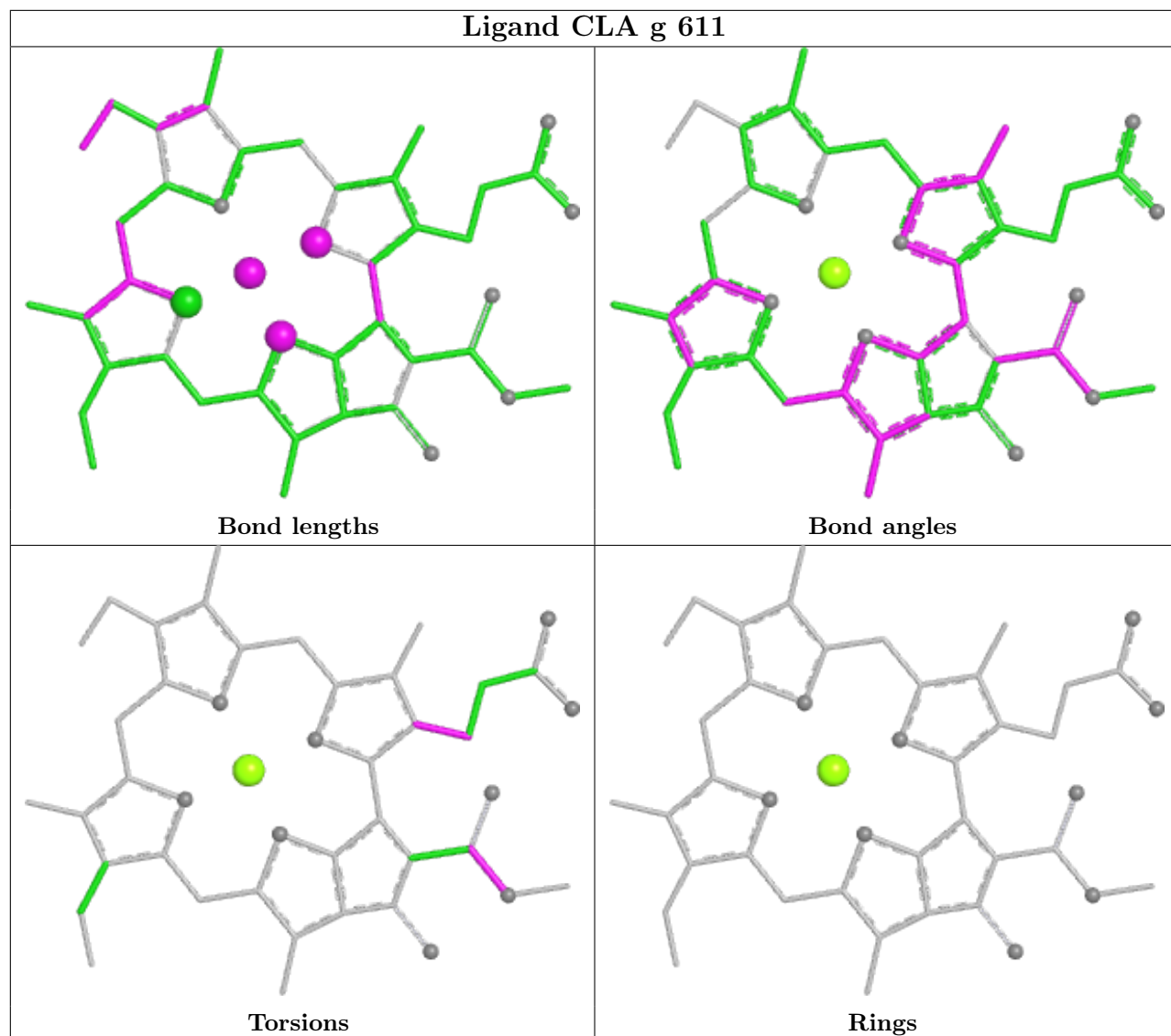


Ligand SQD c 626

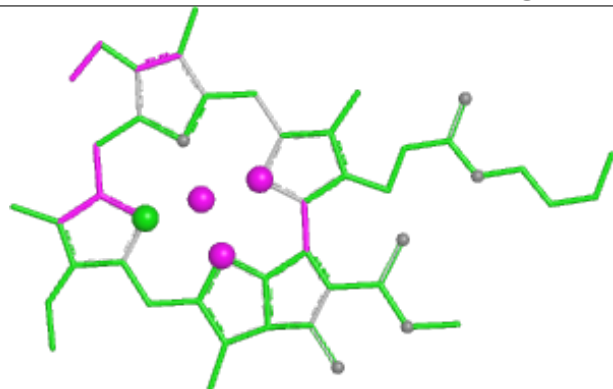




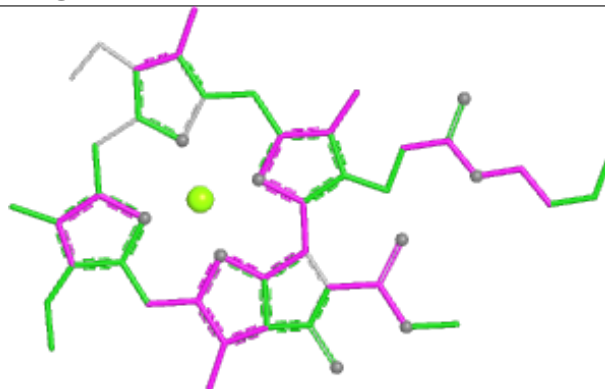
Ligand CLA g 611



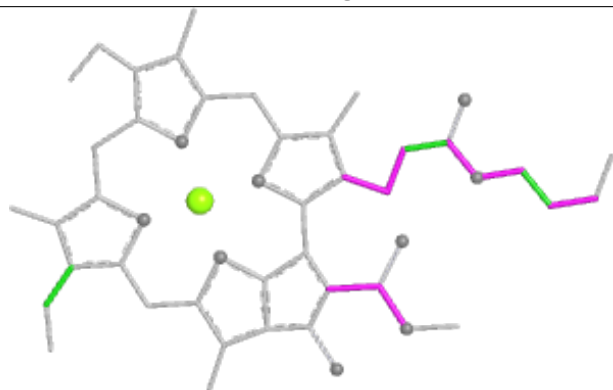
Ligand CLA g 604



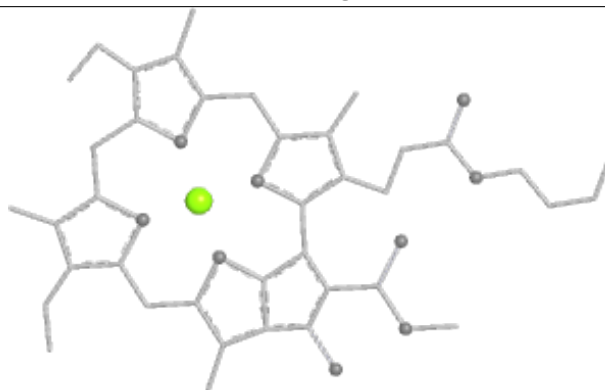
Bond lengths



Bond angles

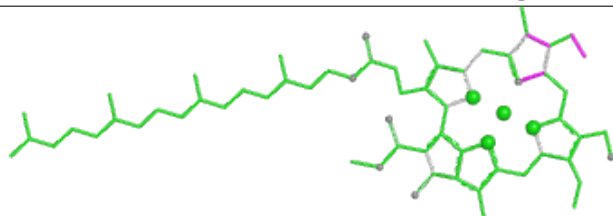


Torsions

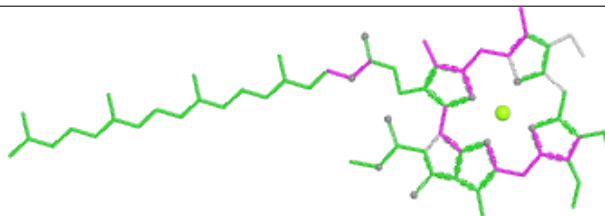


Rings

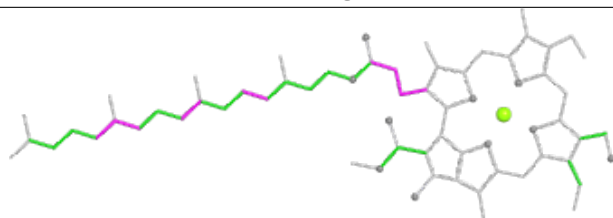
Ligand CHL N 605



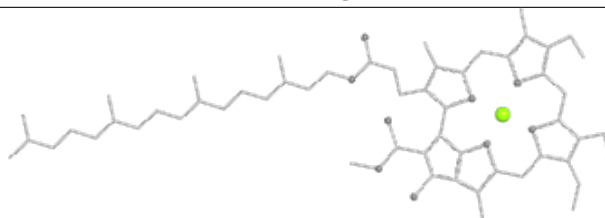
Bond lengths



Bond angles

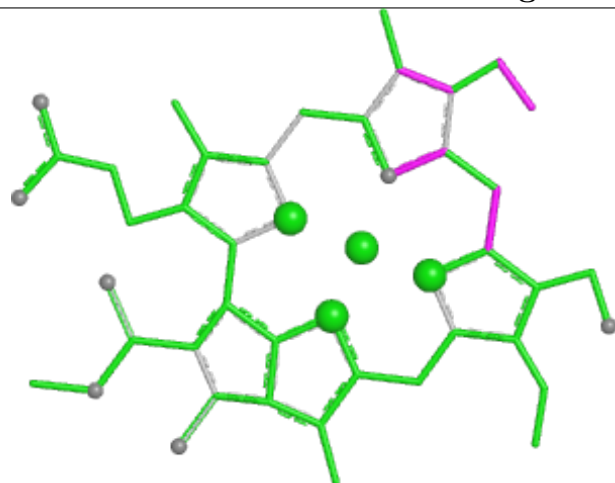


Torsions

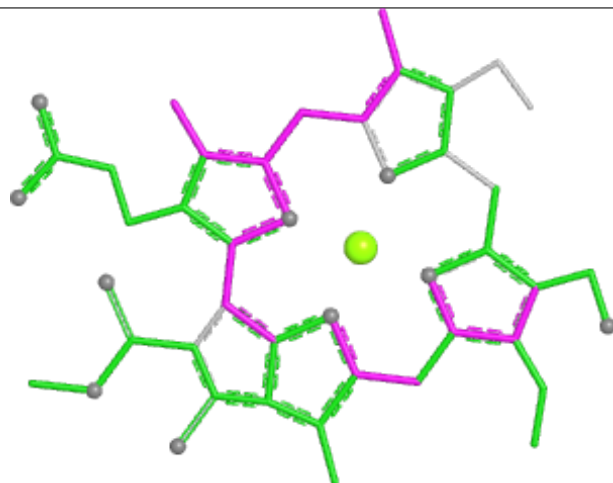


Rings

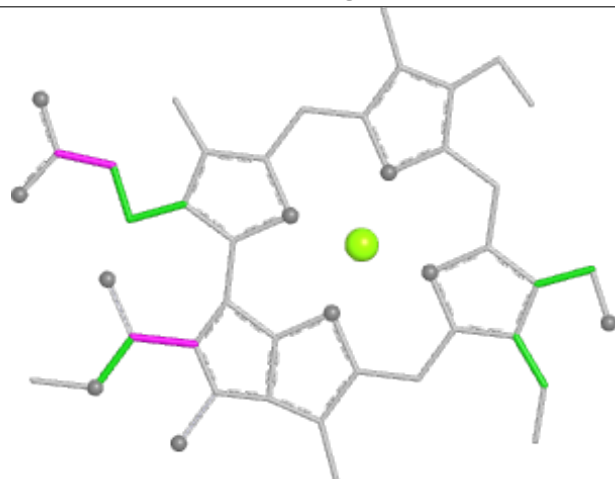
Ligand CHL S 601



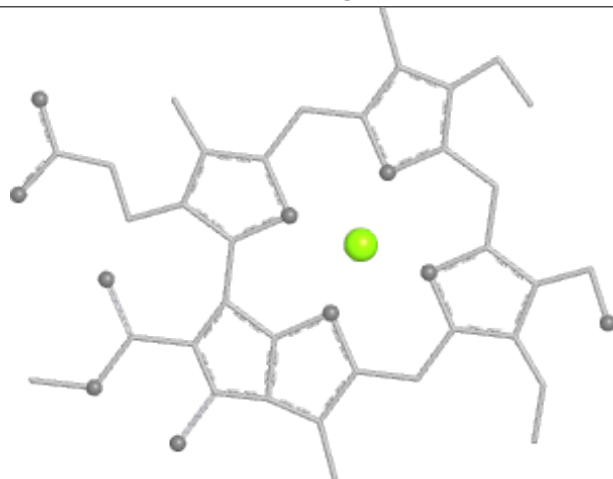
Bond lengths



Bond angles

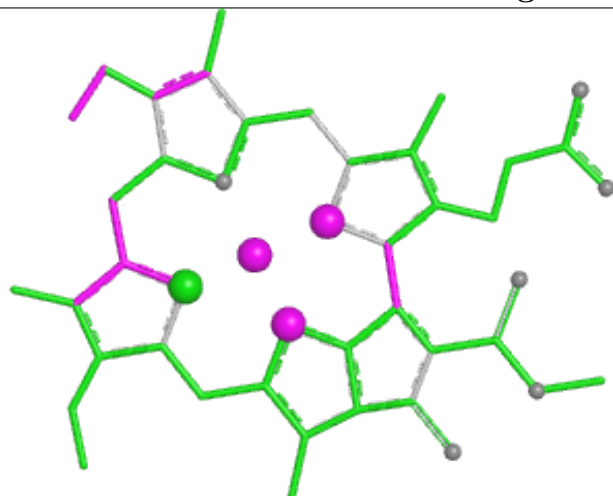


Torsions

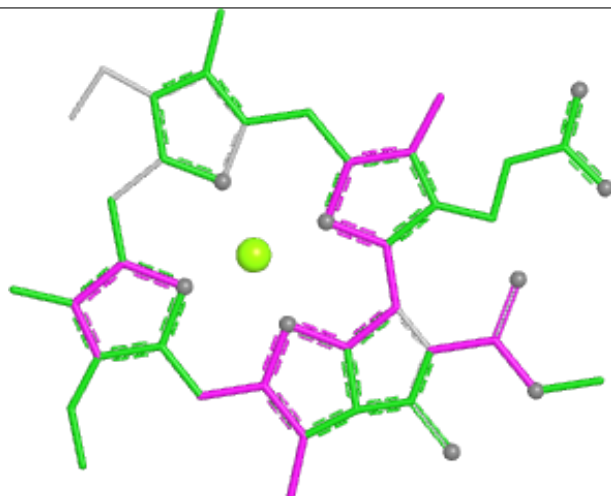


Rings

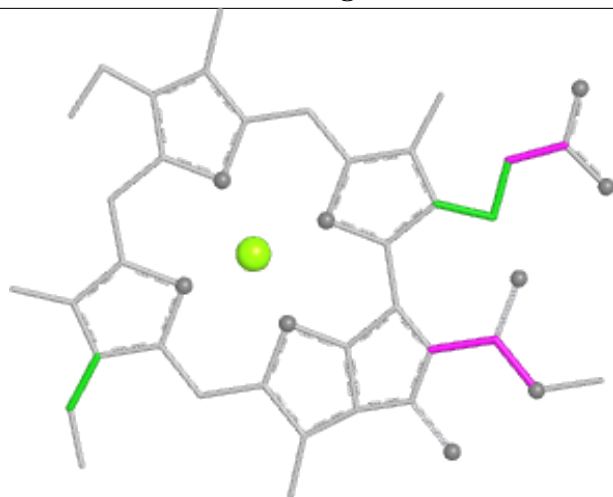
Ligand CLA S 612



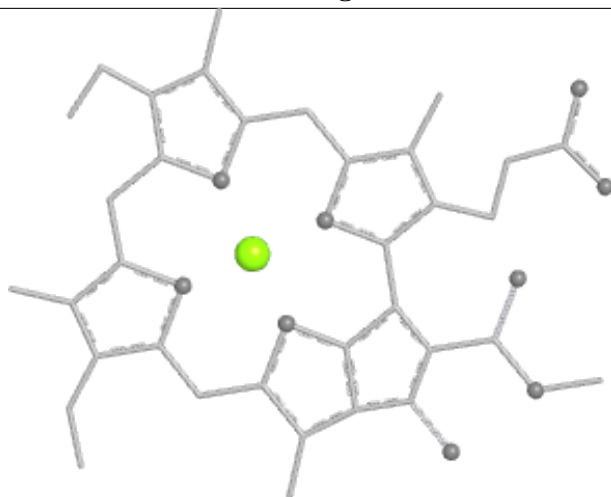
Bond lengths



Bond angles

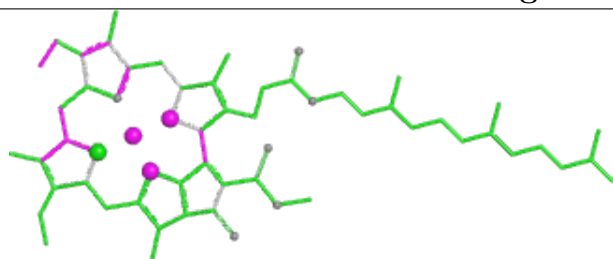


Torsions

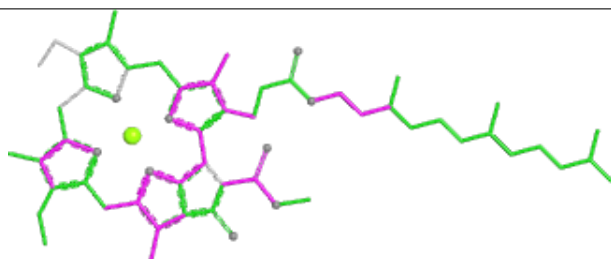


Rings

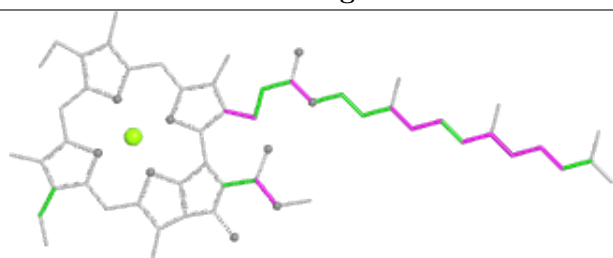
Ligand CLA s 609



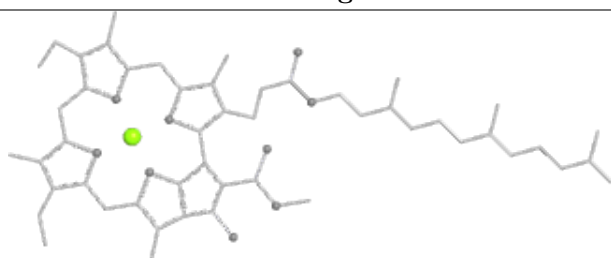
Bond lengths



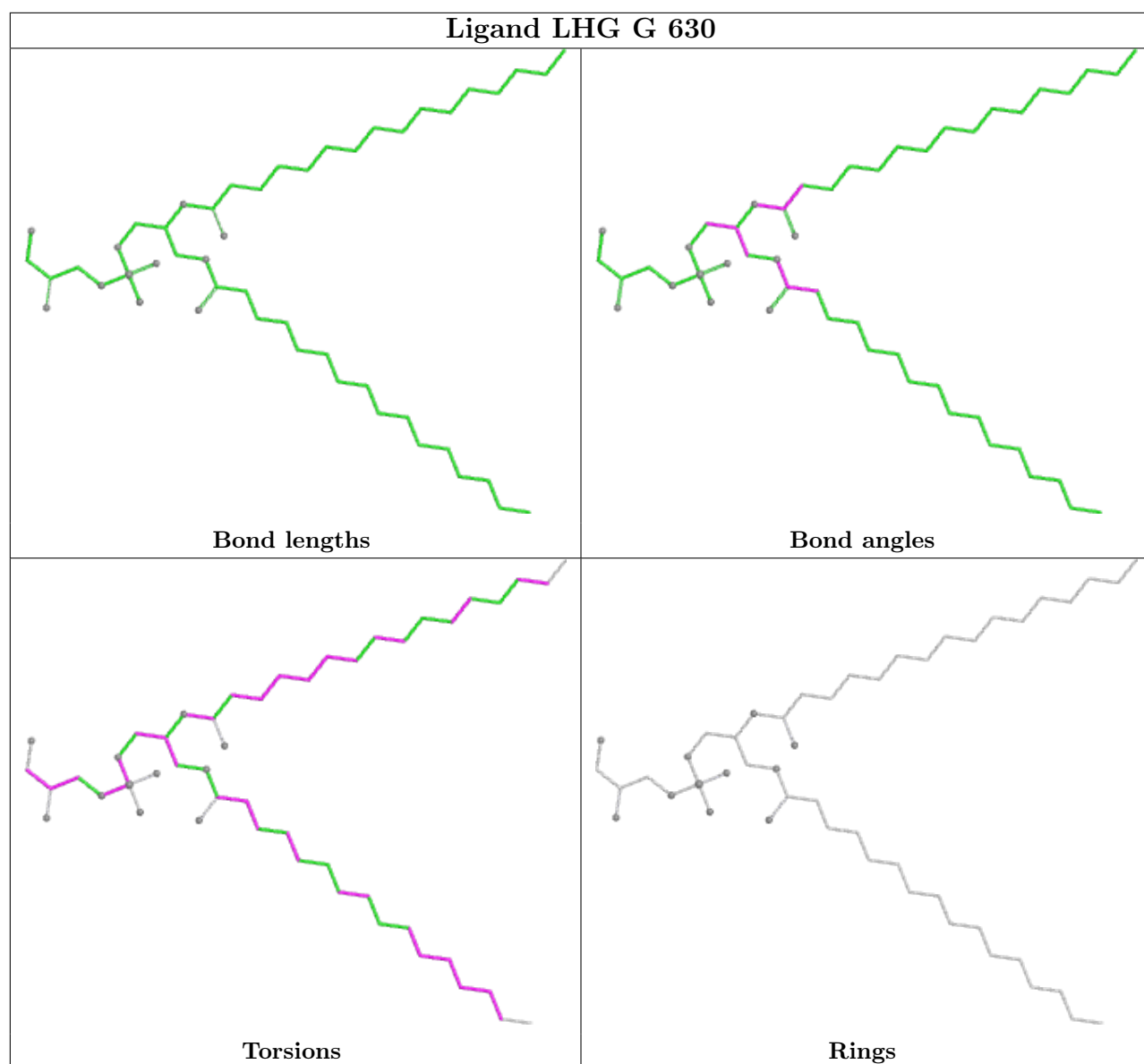
Bond angles



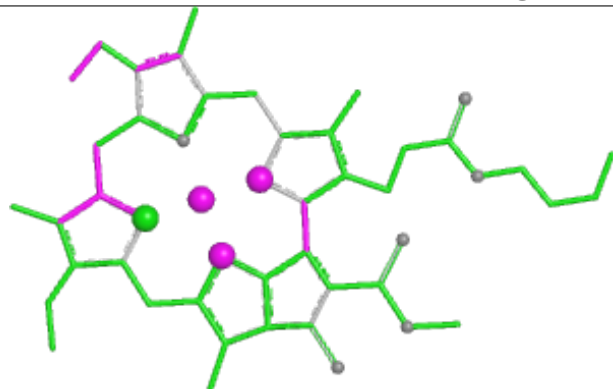
Torsions



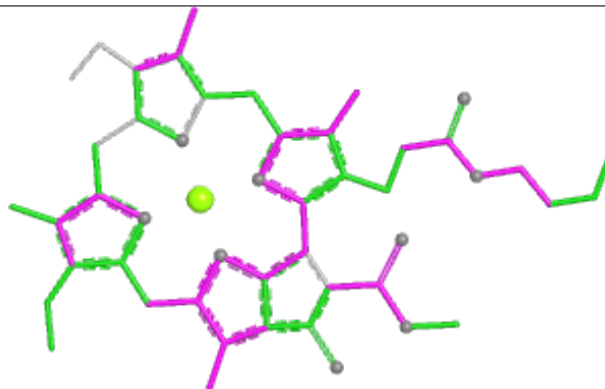
Rings



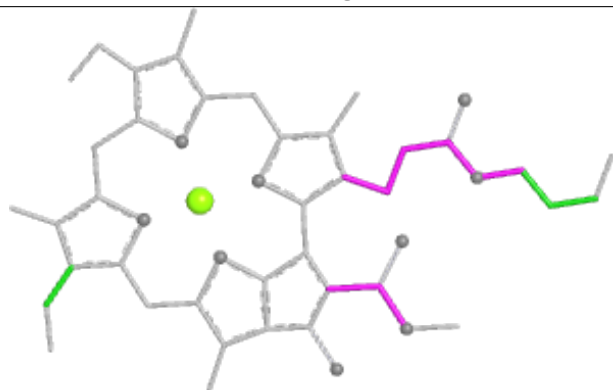
Ligand CLA R 604



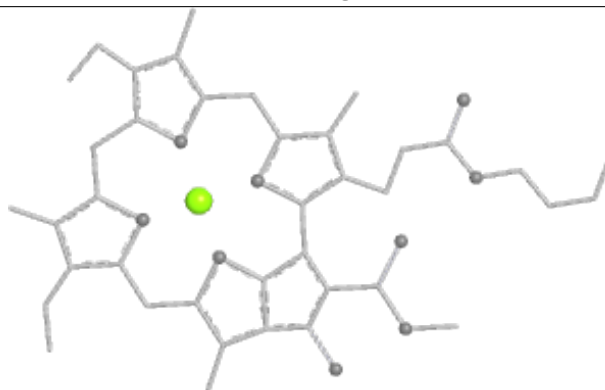
Bond lengths



Bond angles

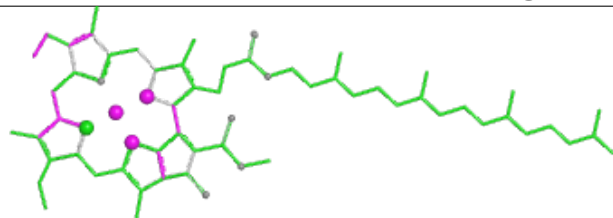


Torsions

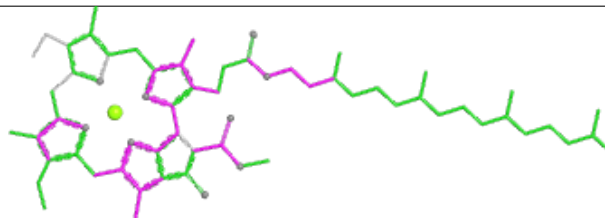


Rings

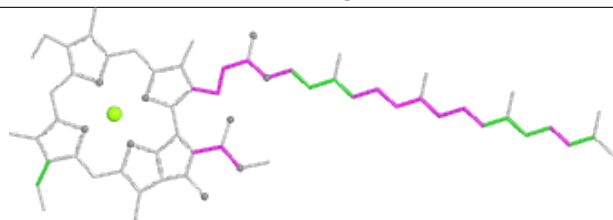
Ligand CLA s 610



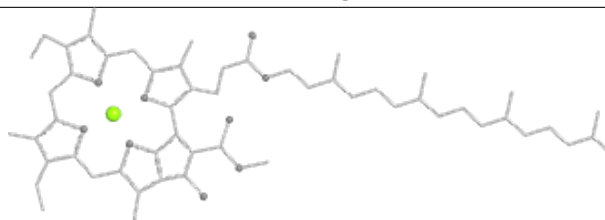
Bond lengths



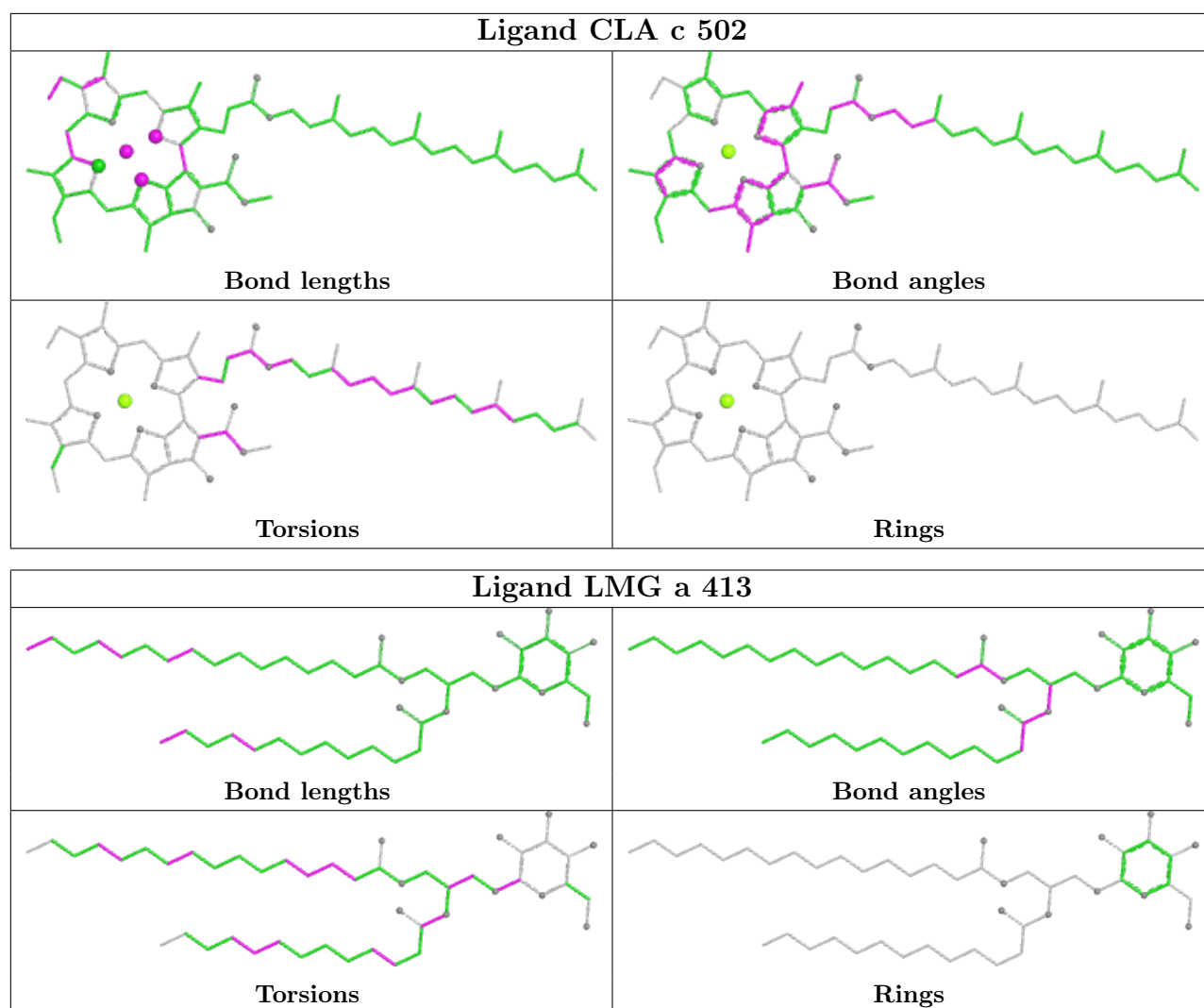
Bond angles

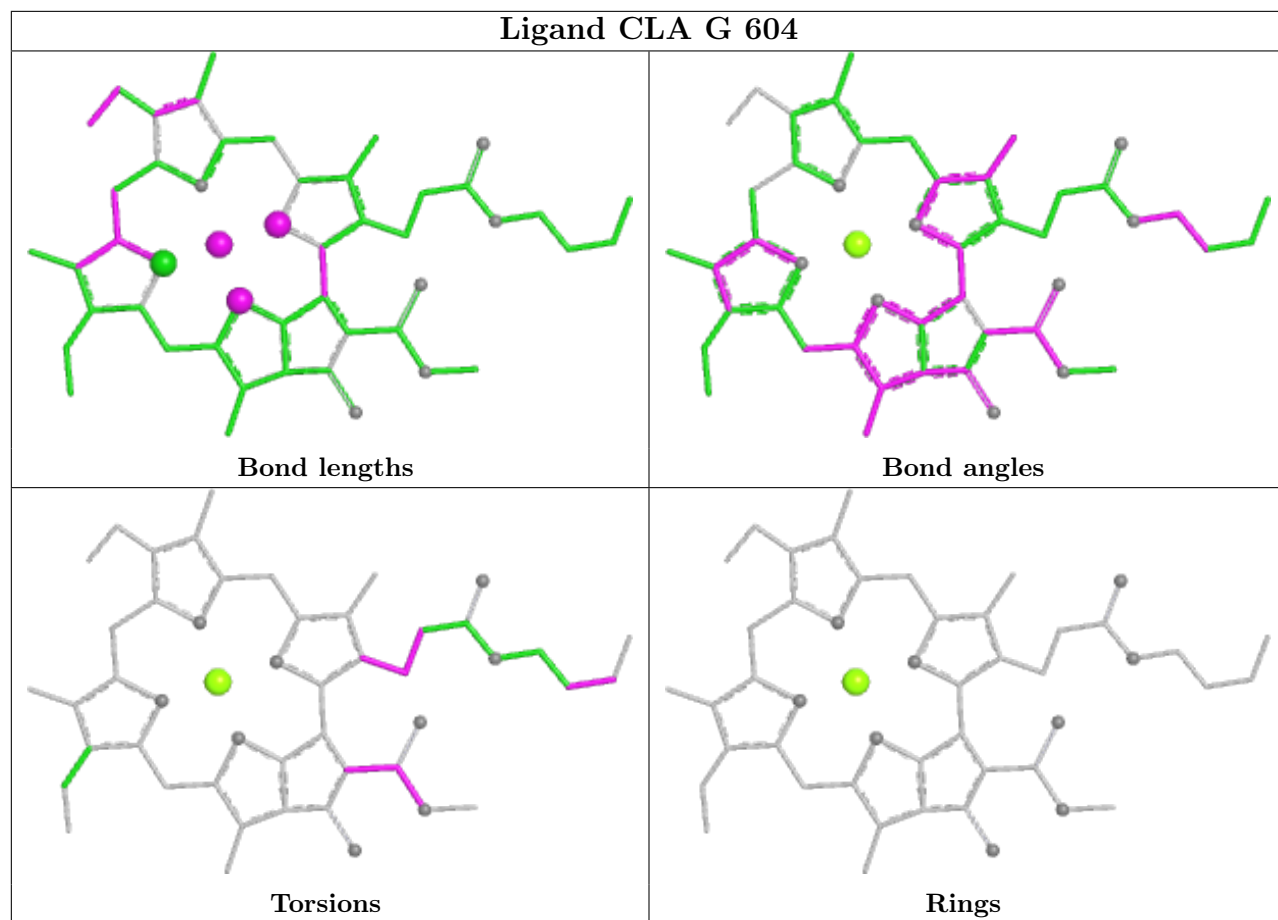


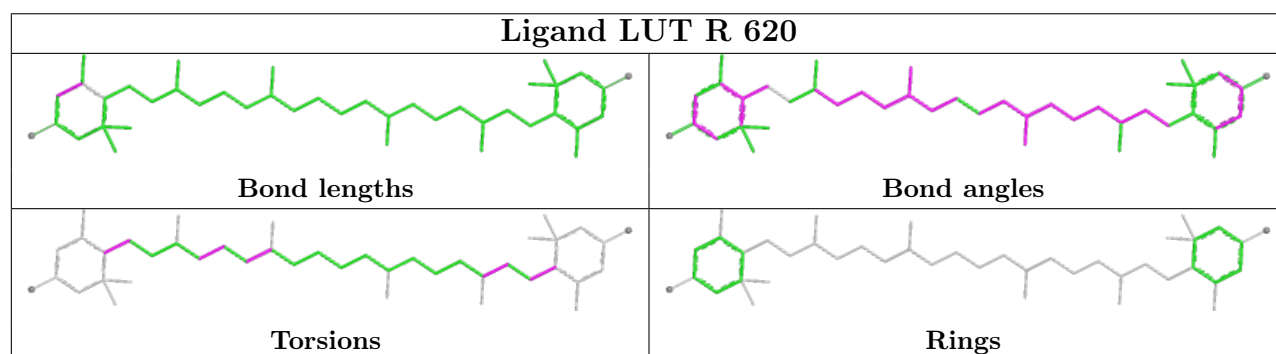
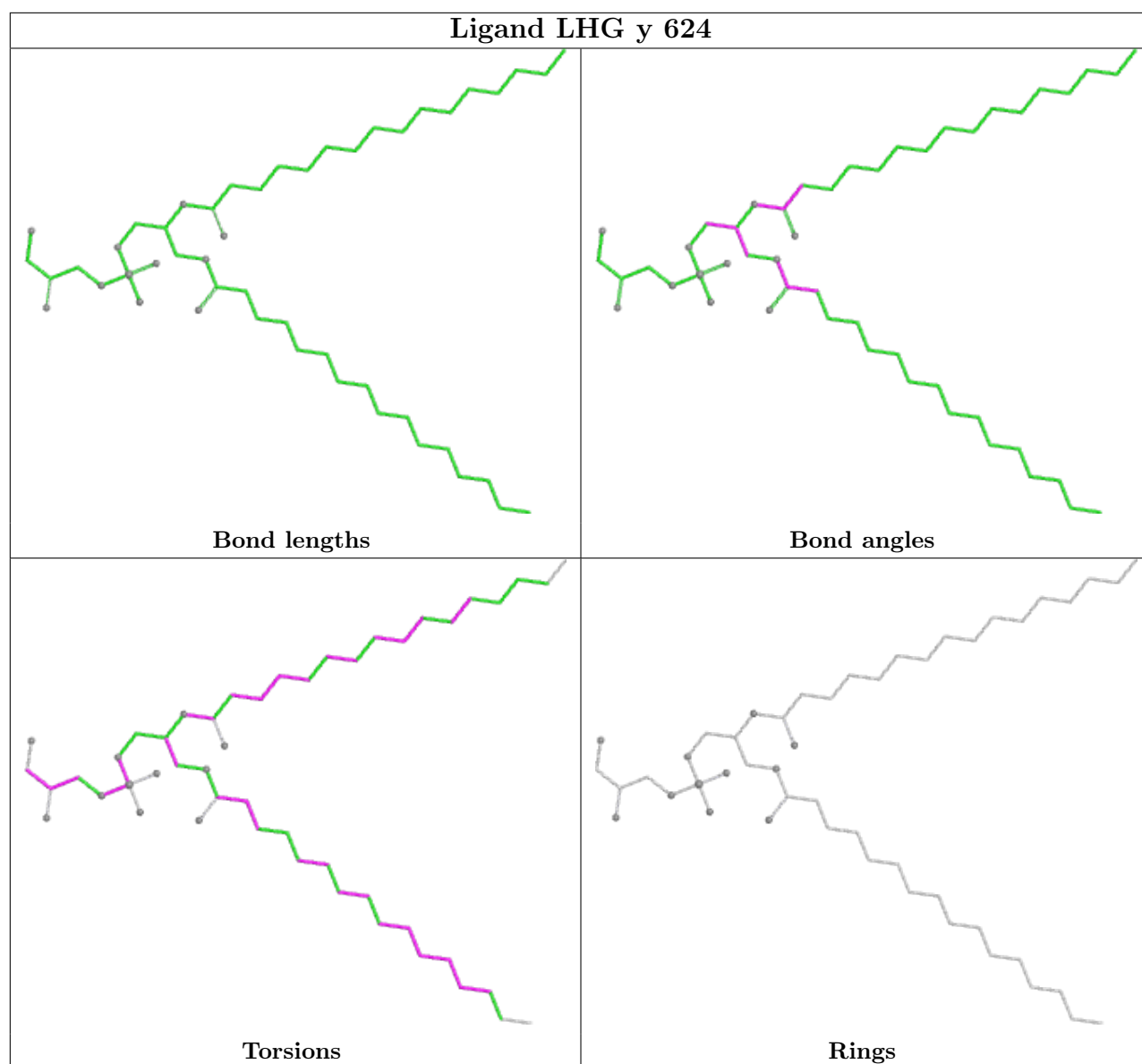
Torsions

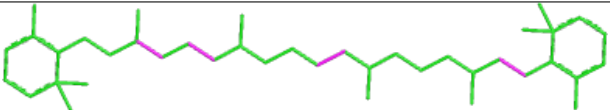
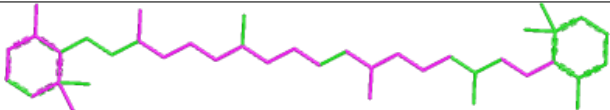
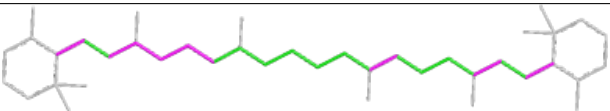
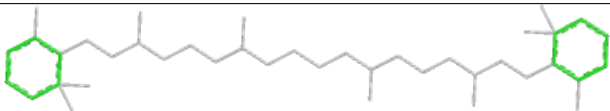
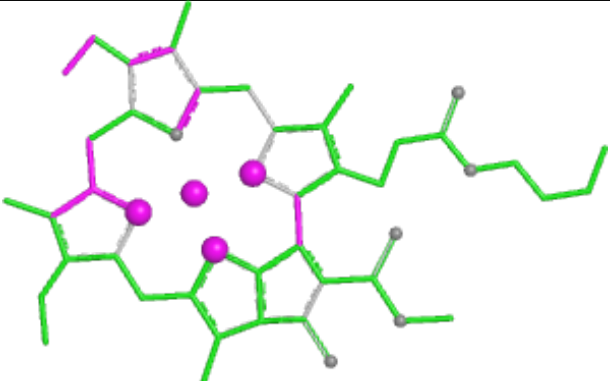
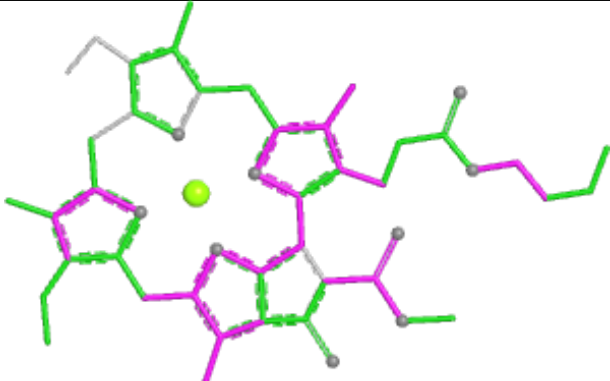
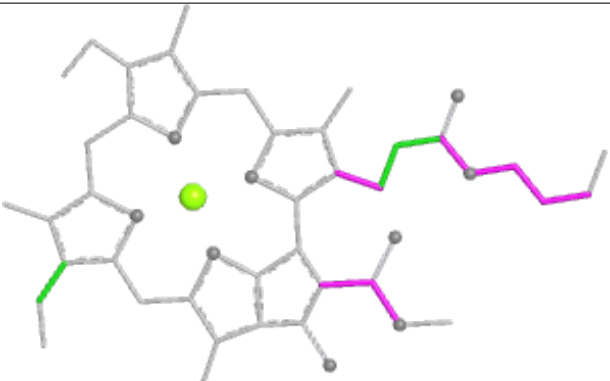
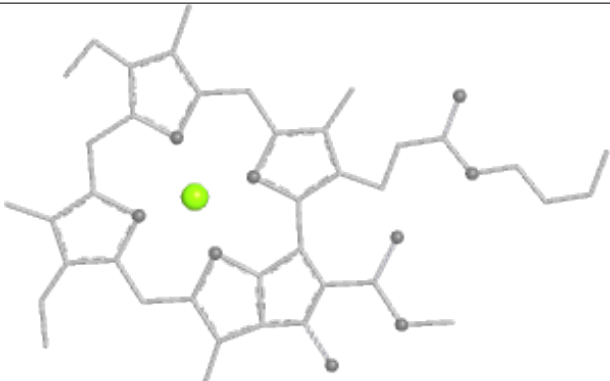


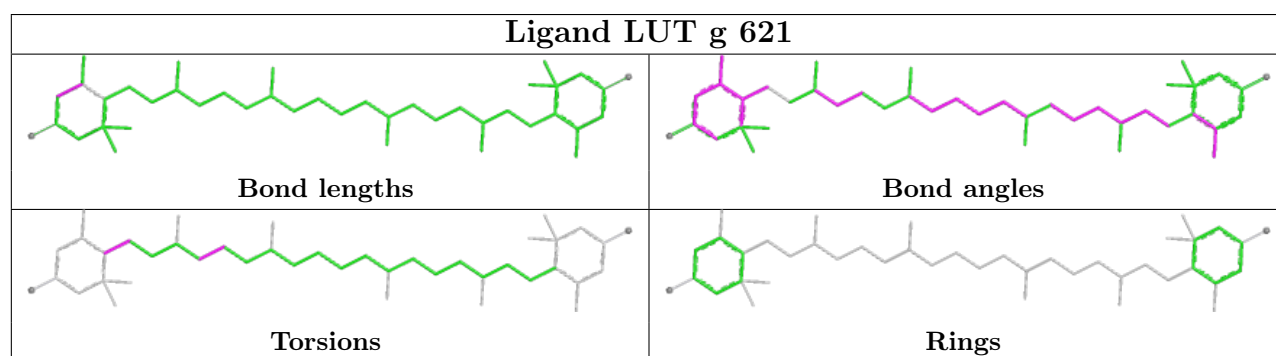
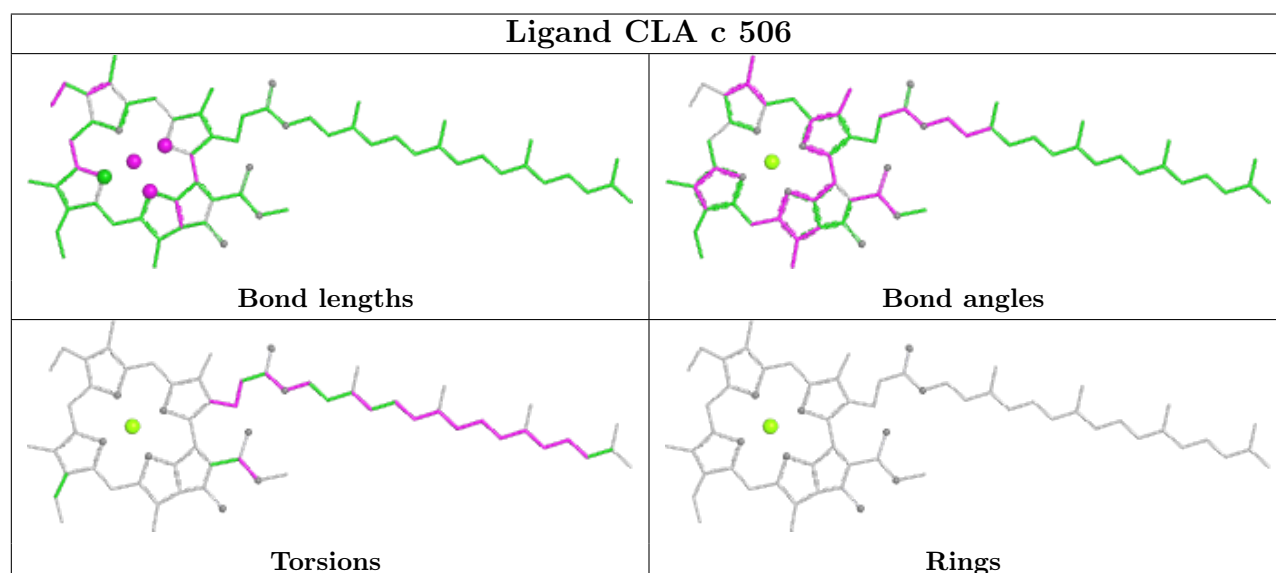
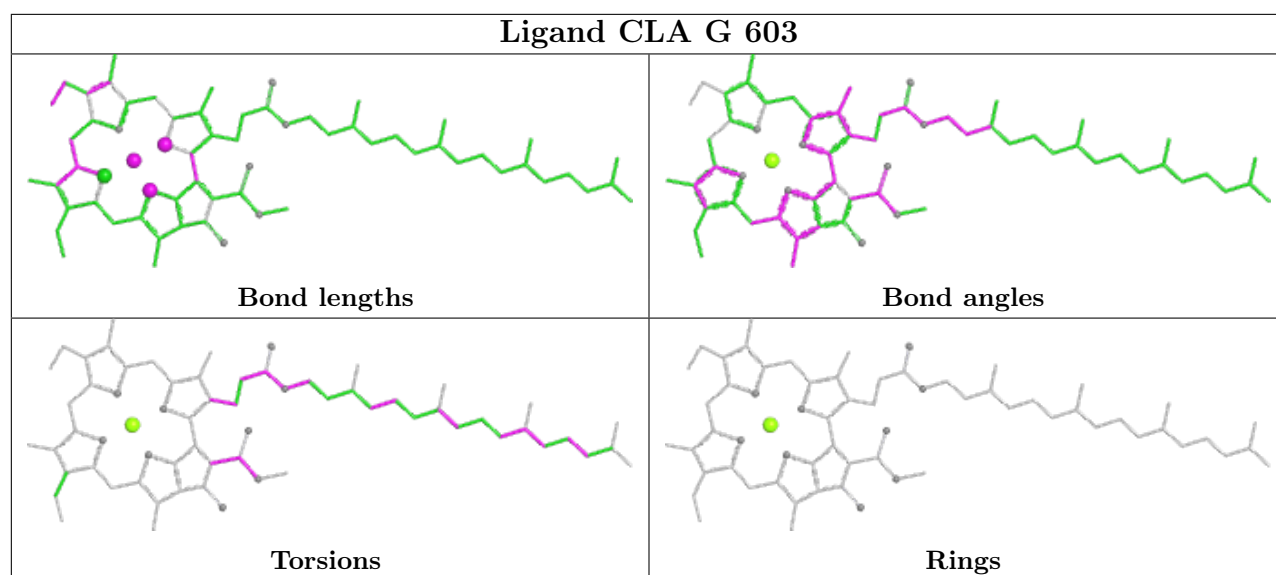
Rings

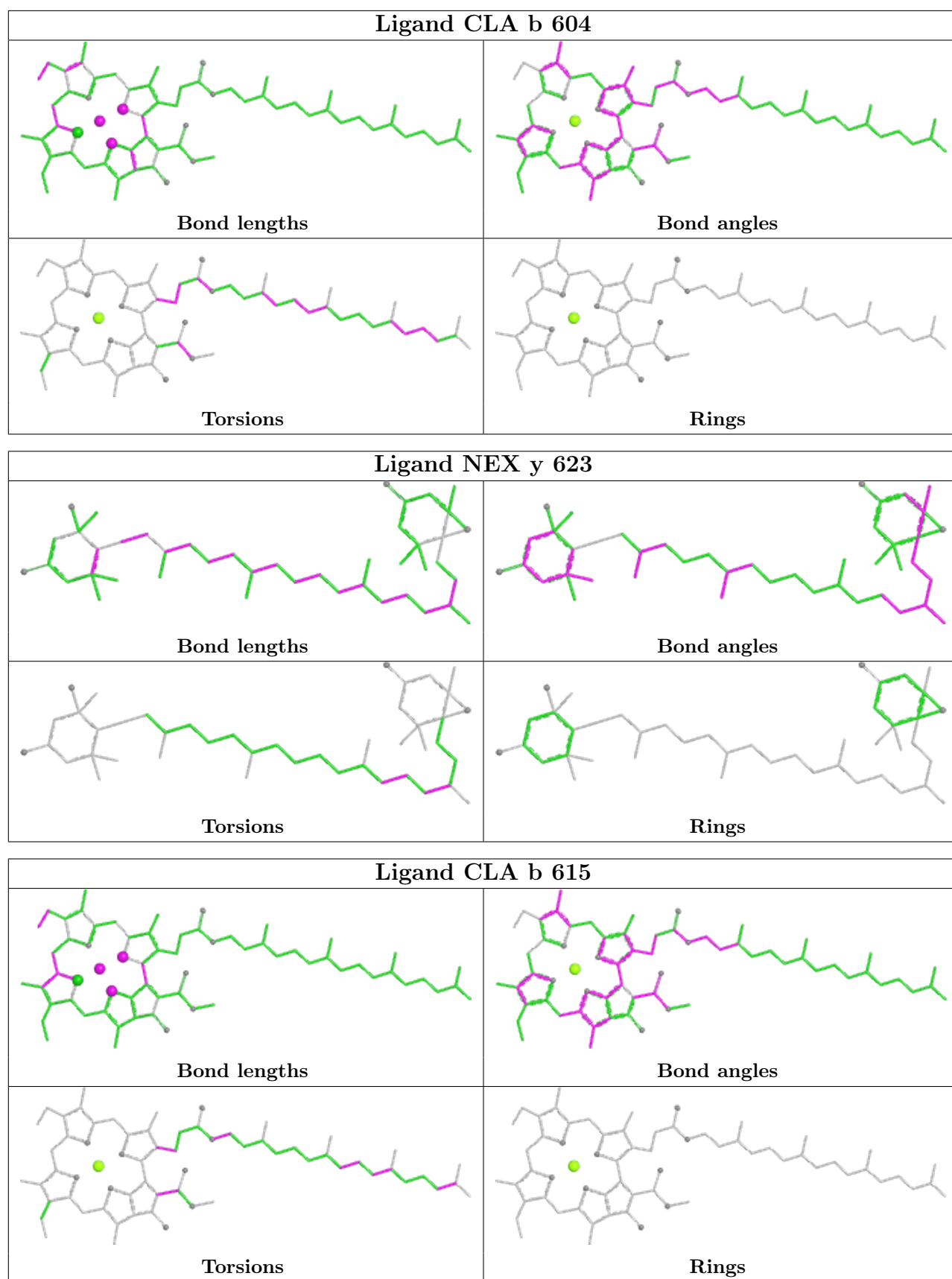


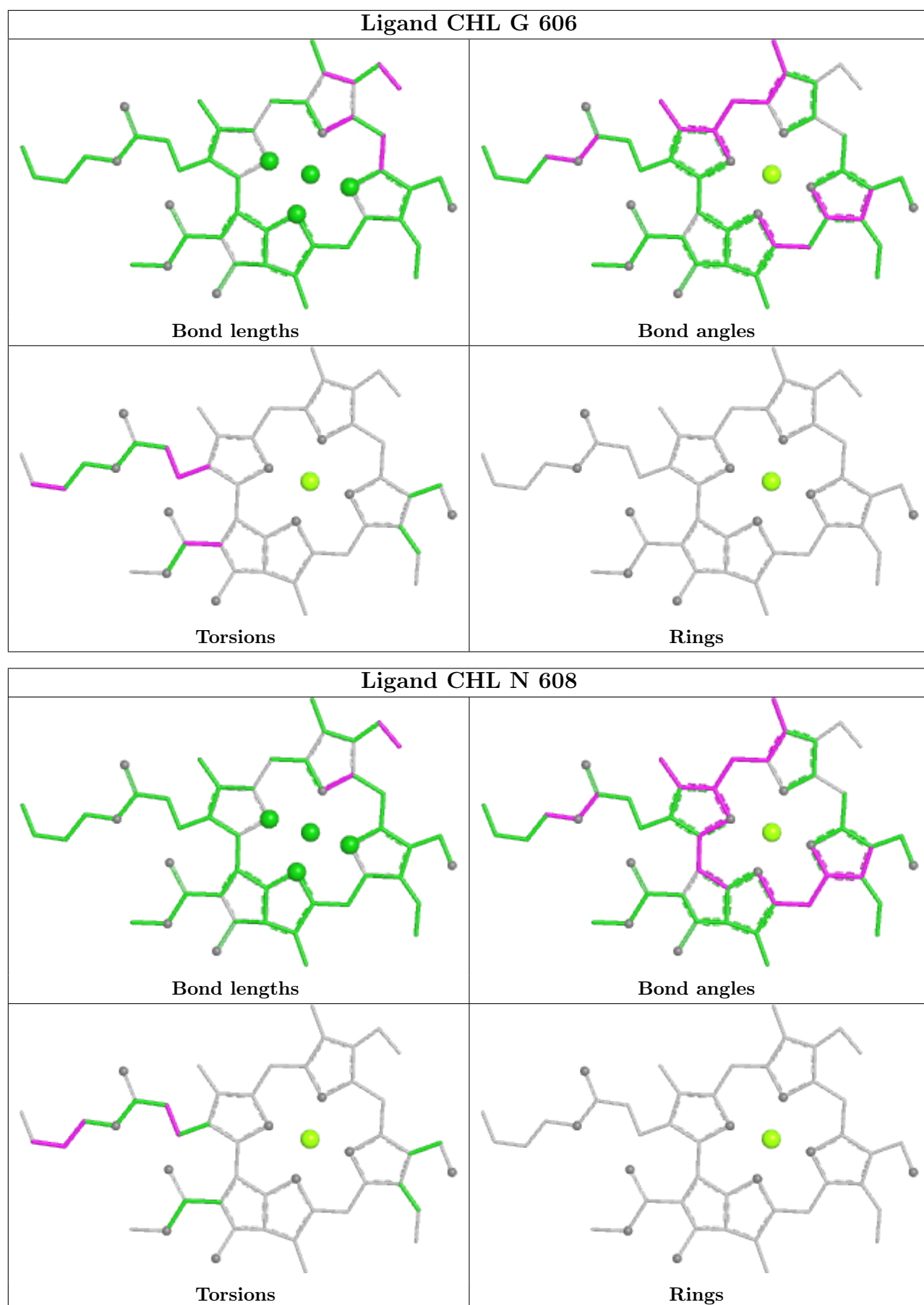


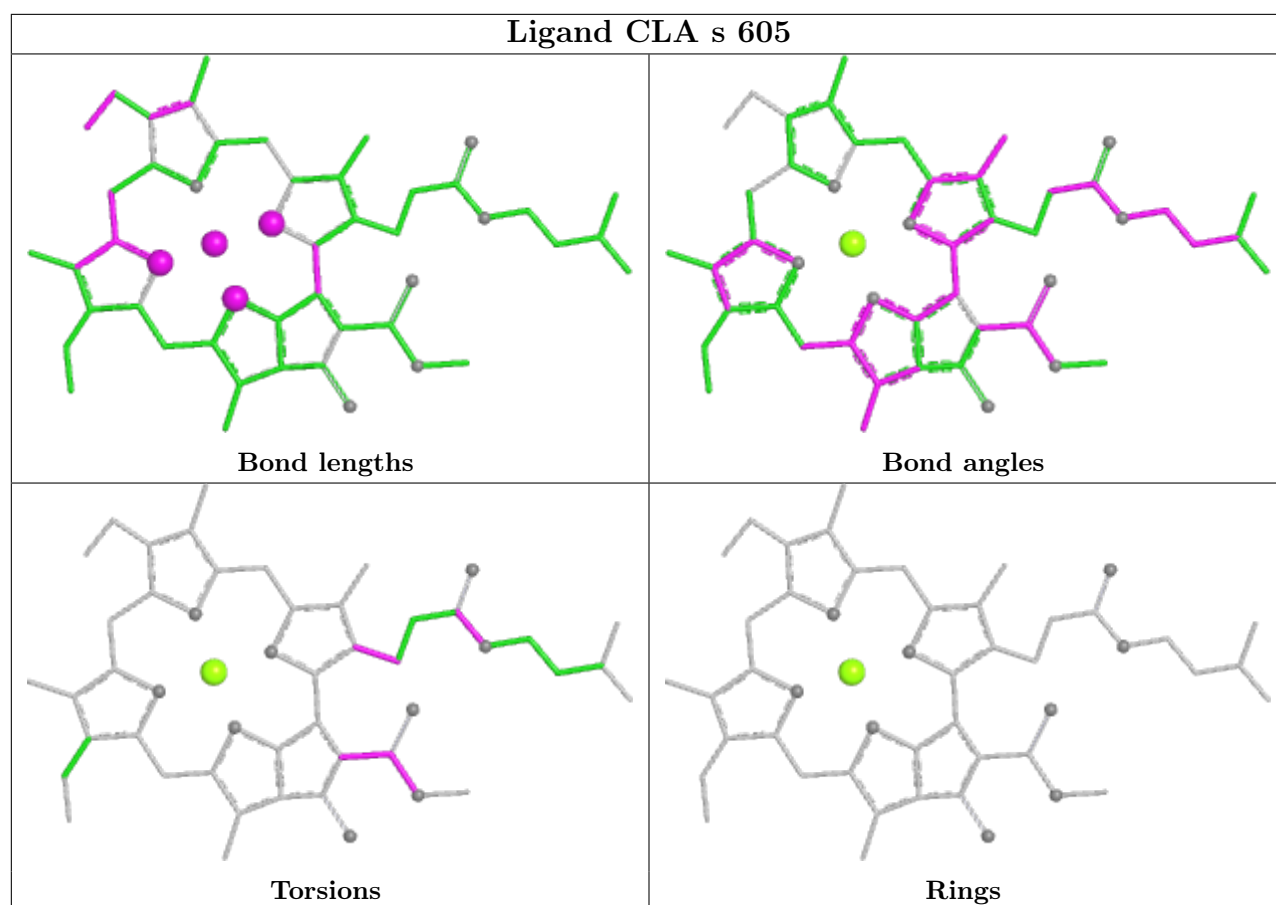


Ligand BCR B 618	
 <p>Bond lengths</p>	 <p>Bond angles</p>
 <p>Torsions</p>	 <p>Rings</p>
Ligand CLA N 611	
 <p>Bond lengths</p>	 <p>Bond angles</p>
 <p>Torsions</p>	 <p>Rings</p>

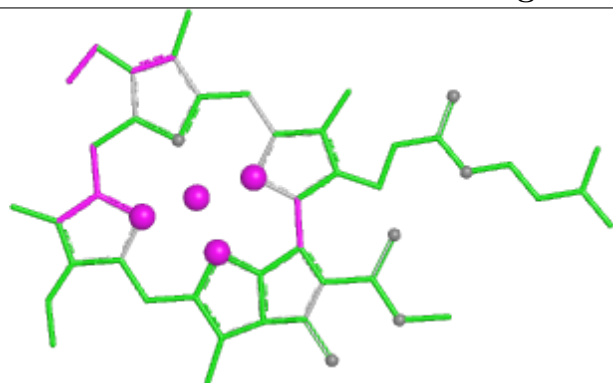




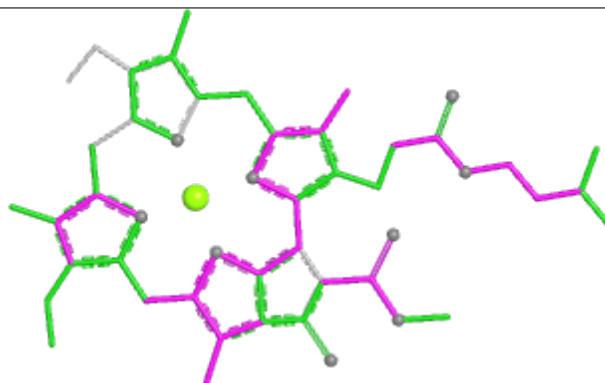




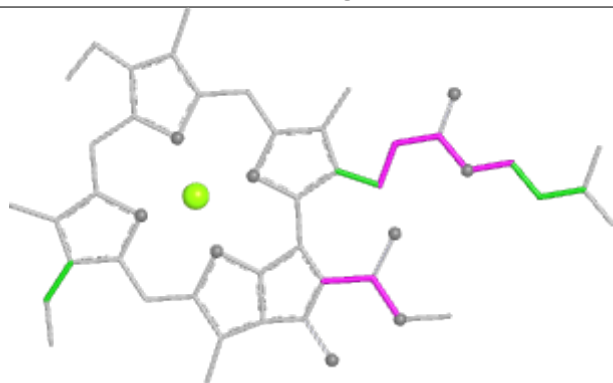
Ligand CLA S 617



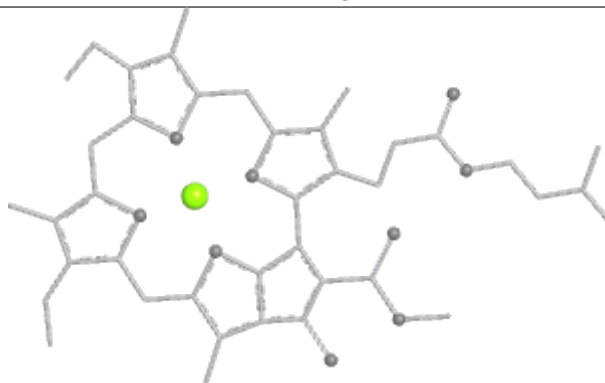
Bond lengths



Bond angles

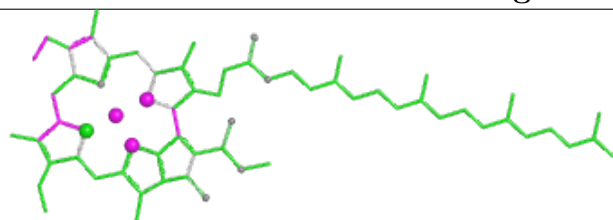


Torsions

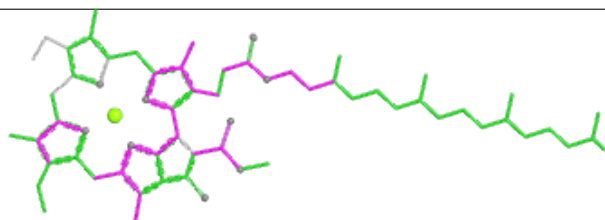


Rings

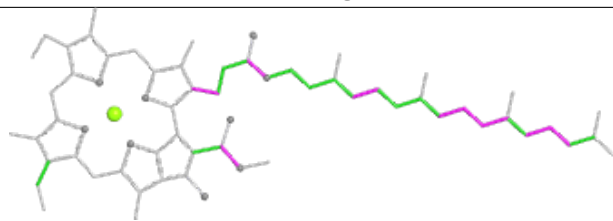
Ligand CLA D 403



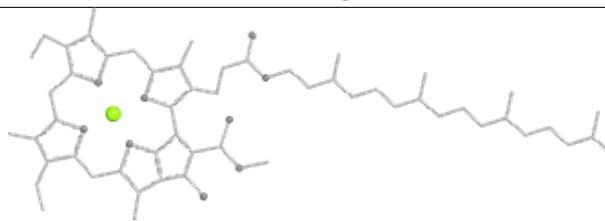
Bond lengths



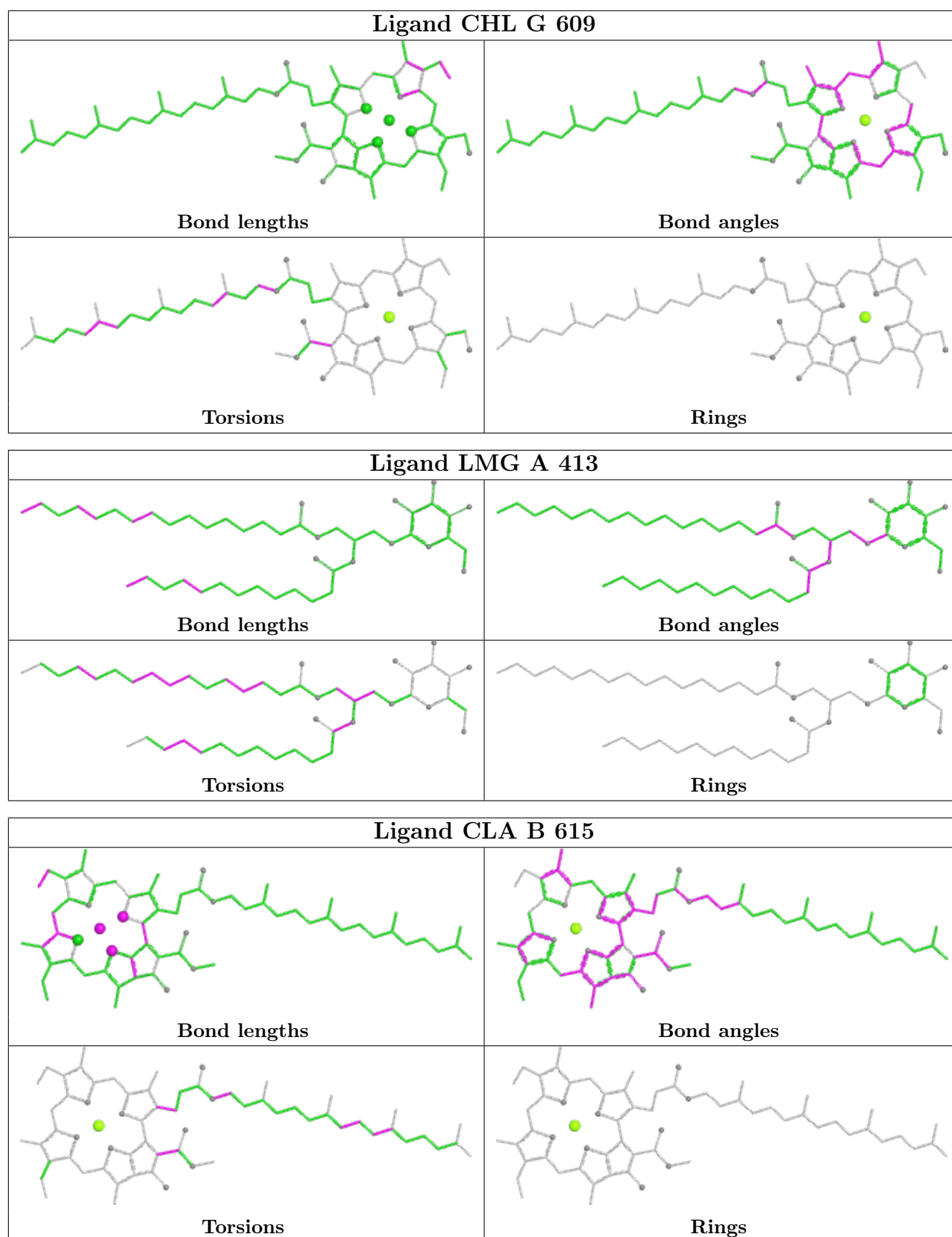
Bond angles

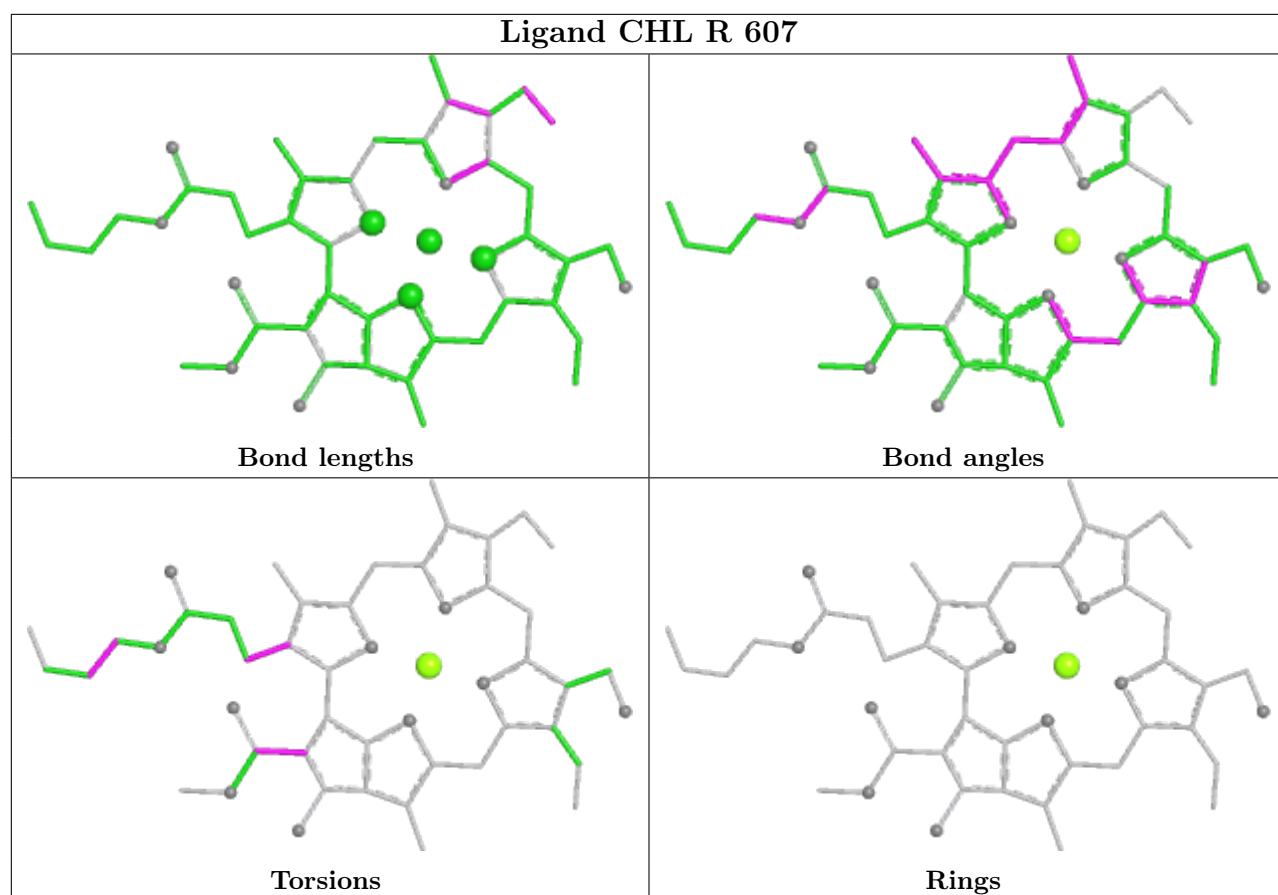


Torsions

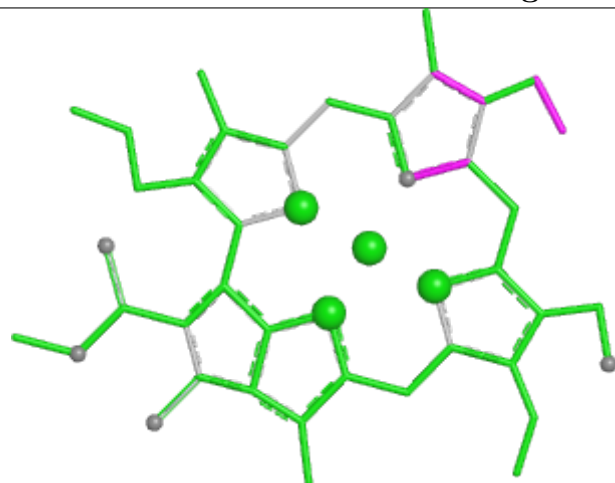


Rings

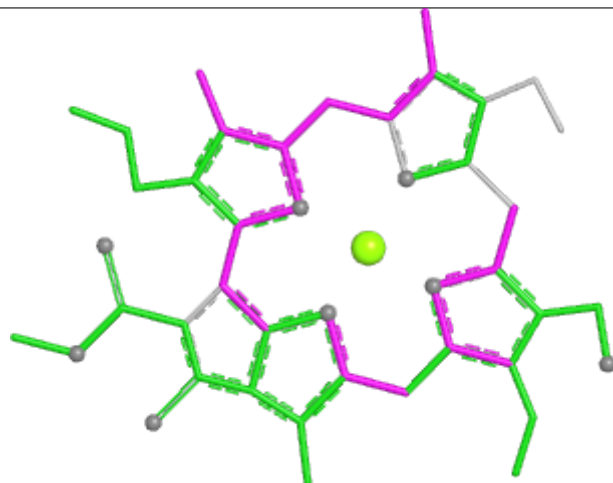




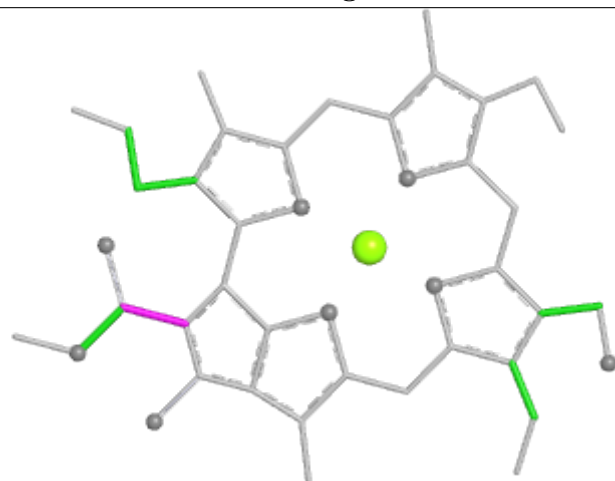
Ligand CHL r 606



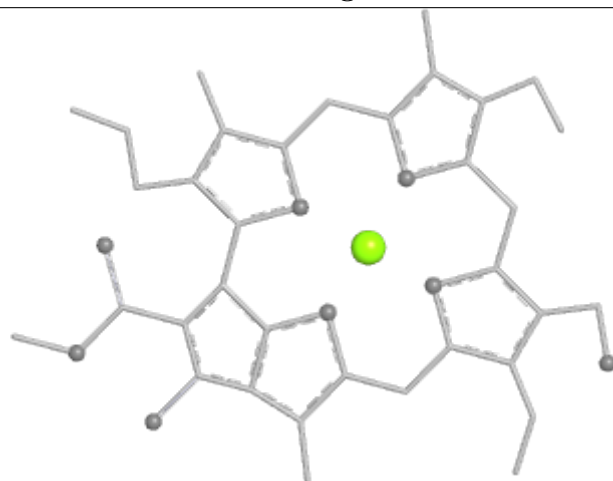
Bond lengths



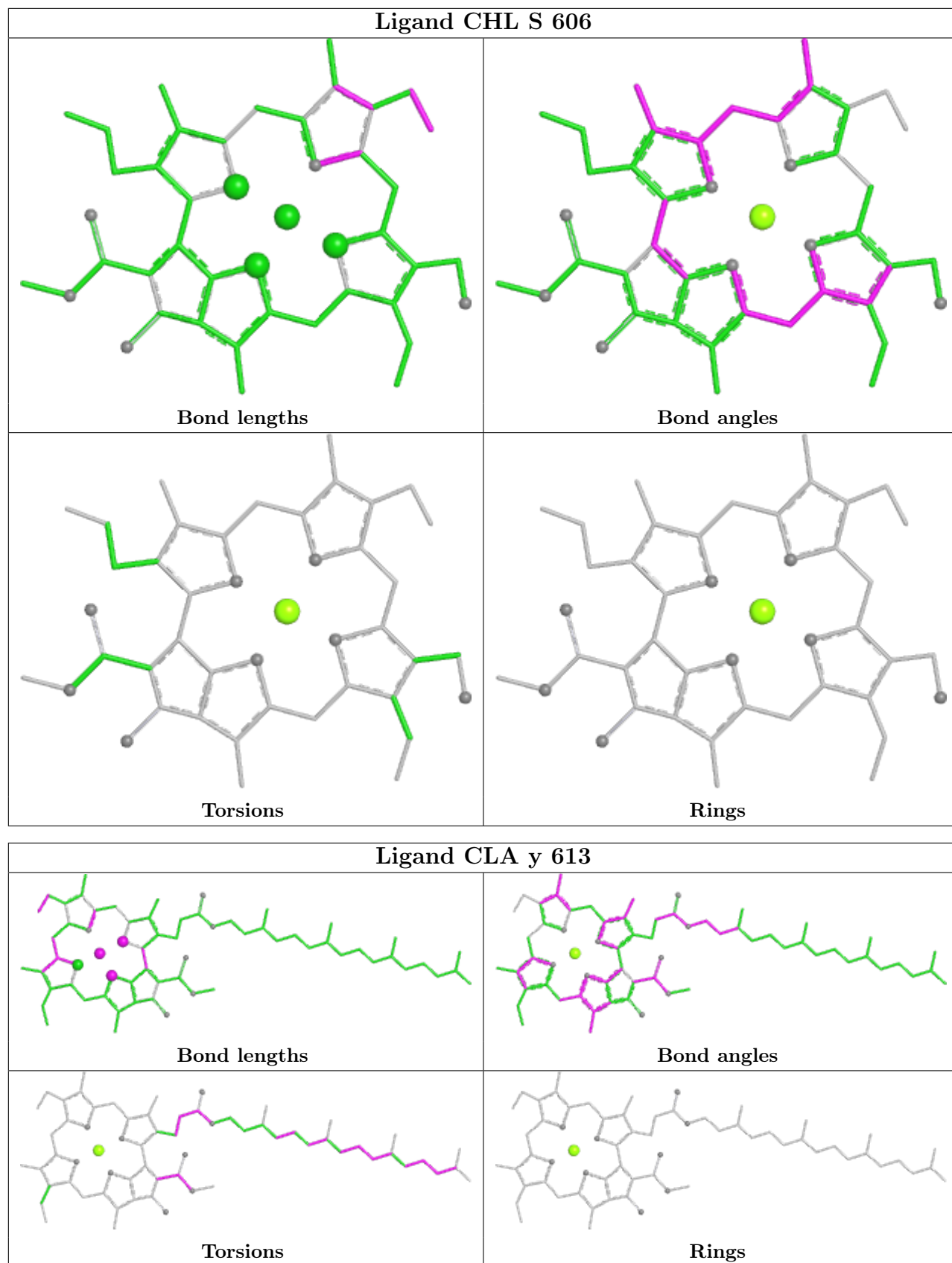
Bond angles

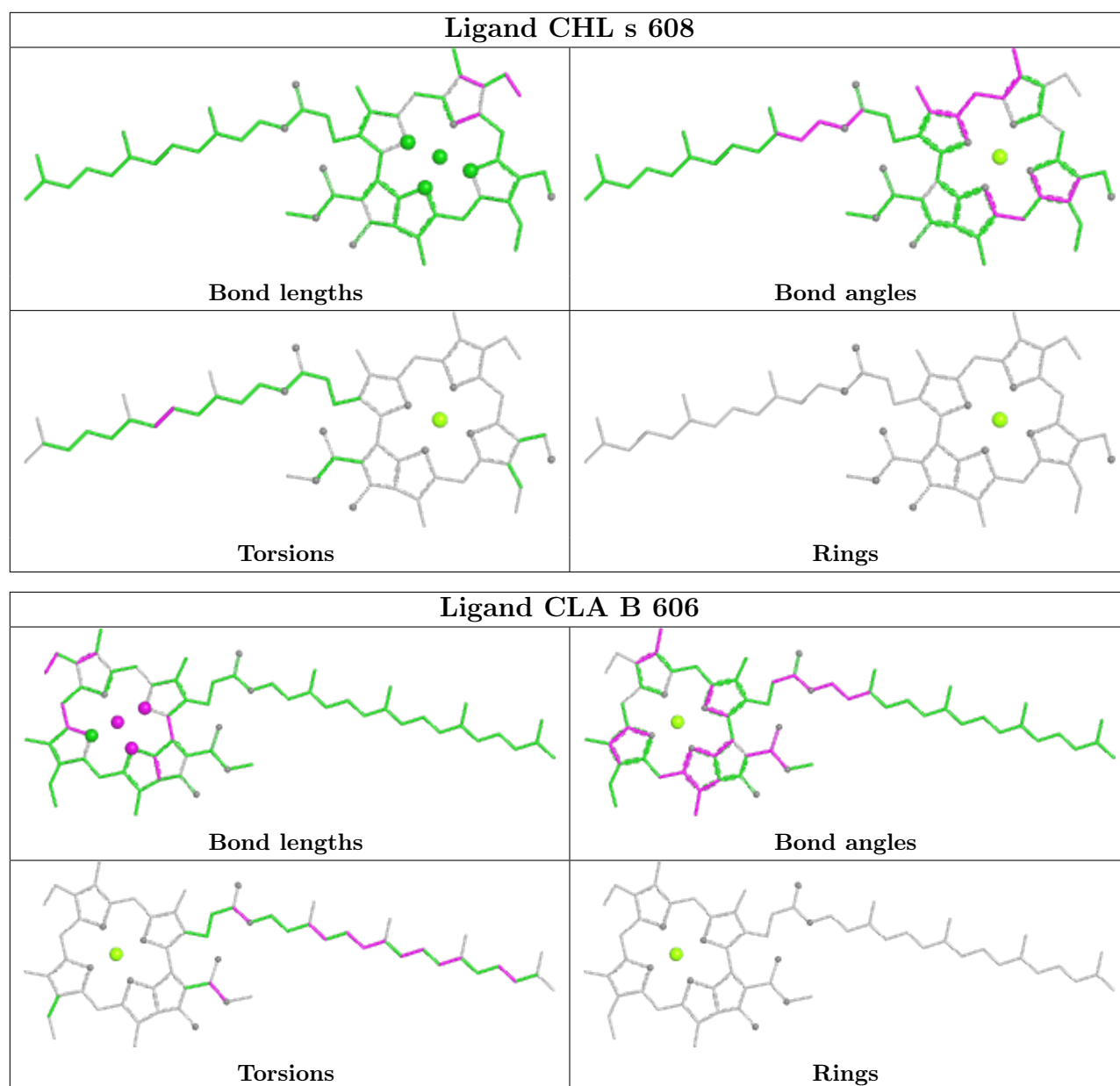


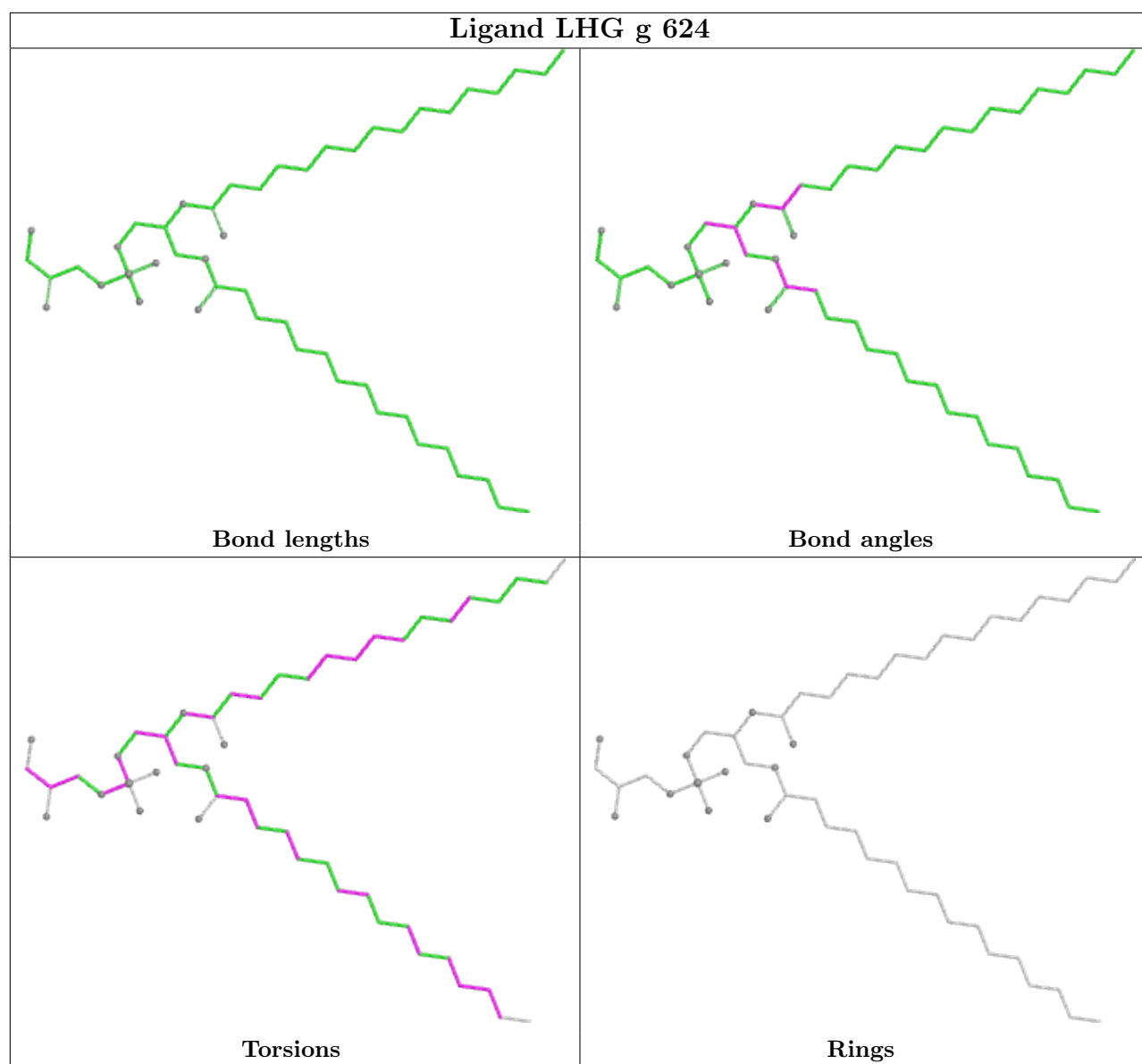
Torsions



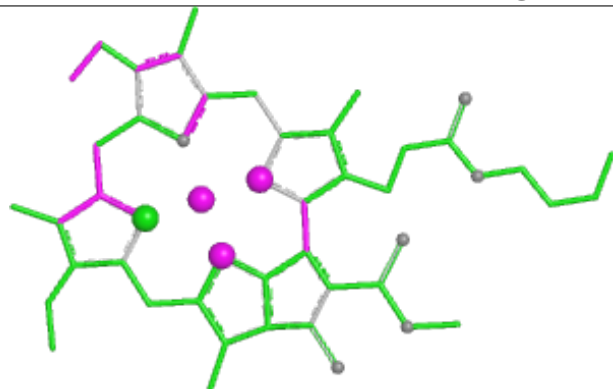
Rings



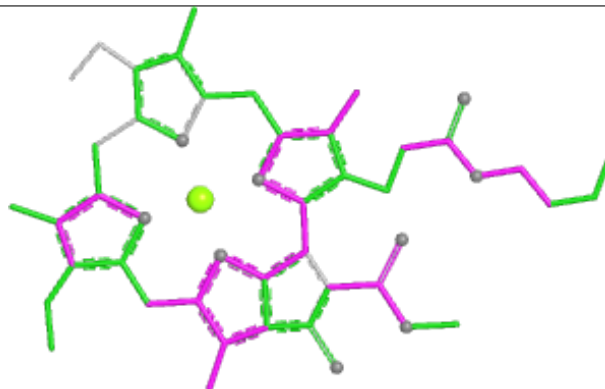




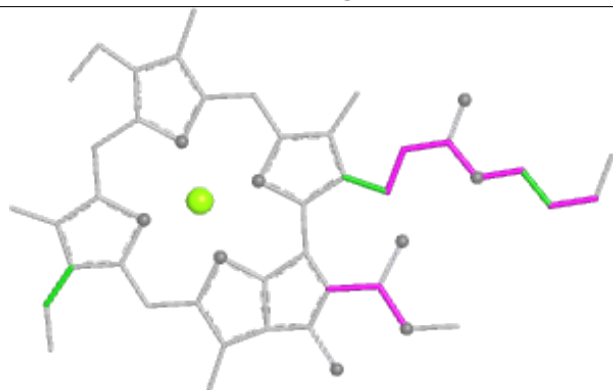
Ligand CLA G 614



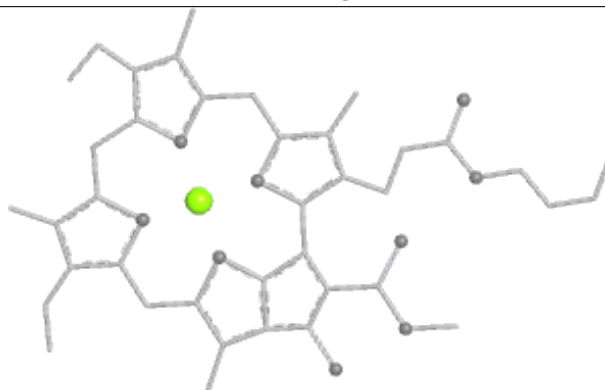
Bond lengths



Bond angles

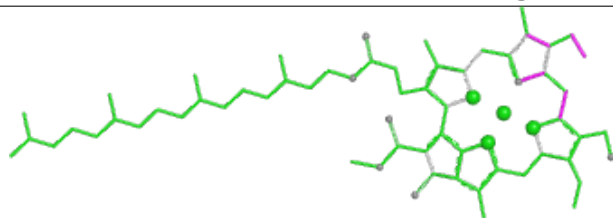


Torsions

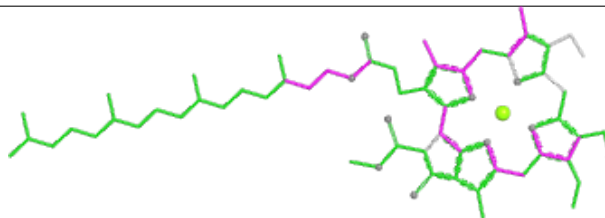


Rings

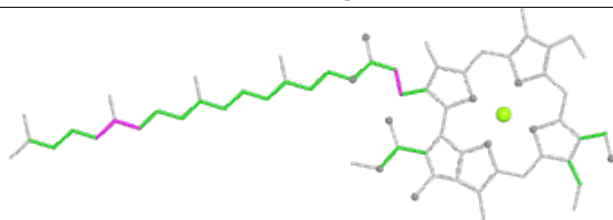
Ligand CHL N 606



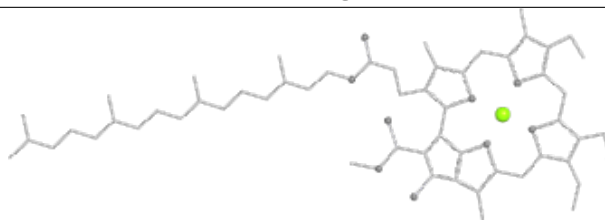
Bond lengths



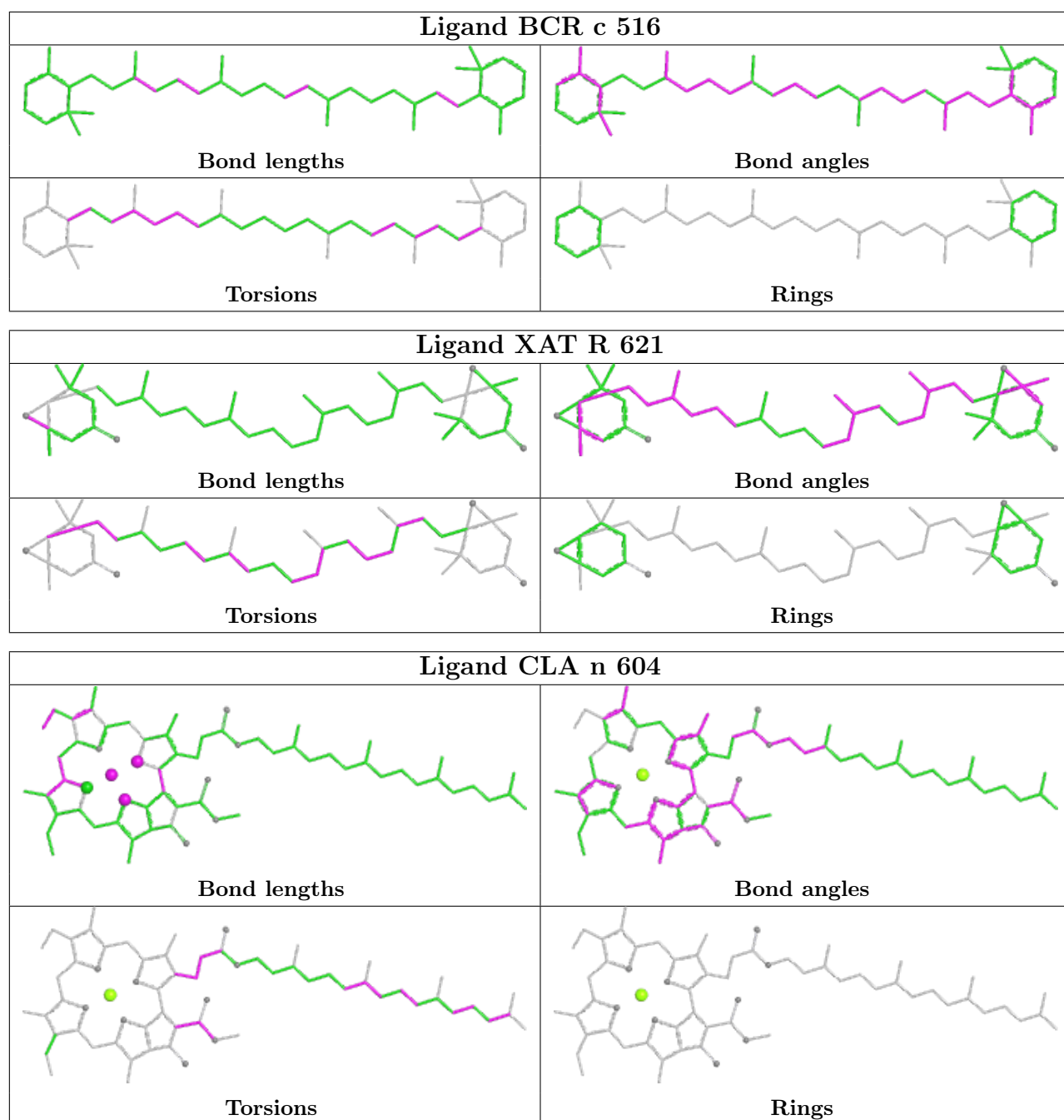
Bond angles

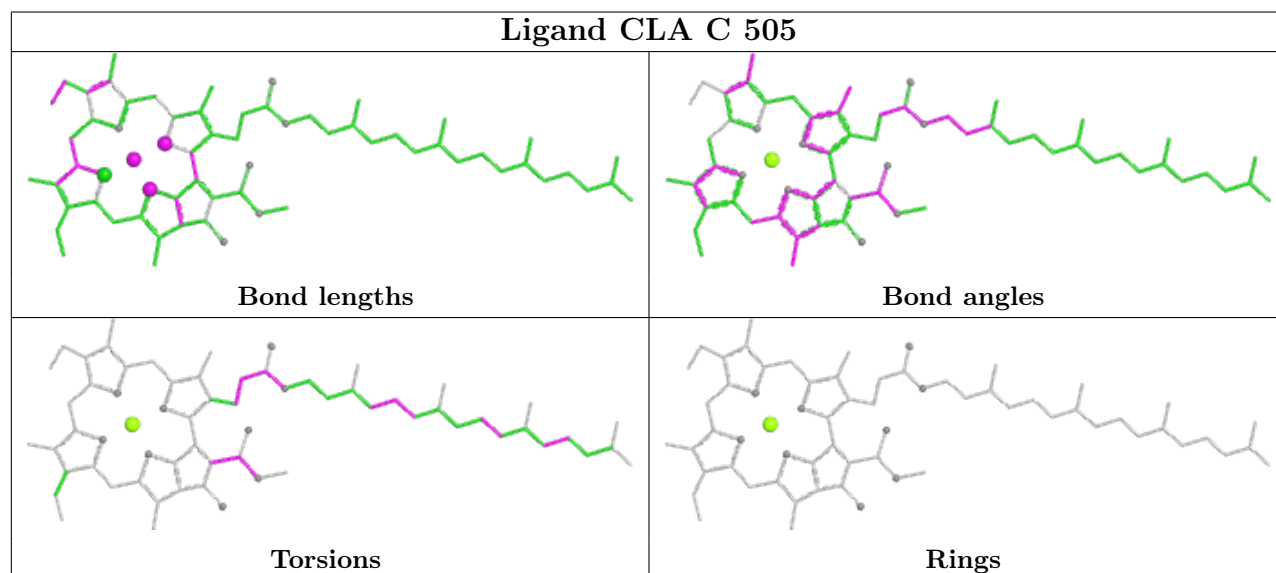
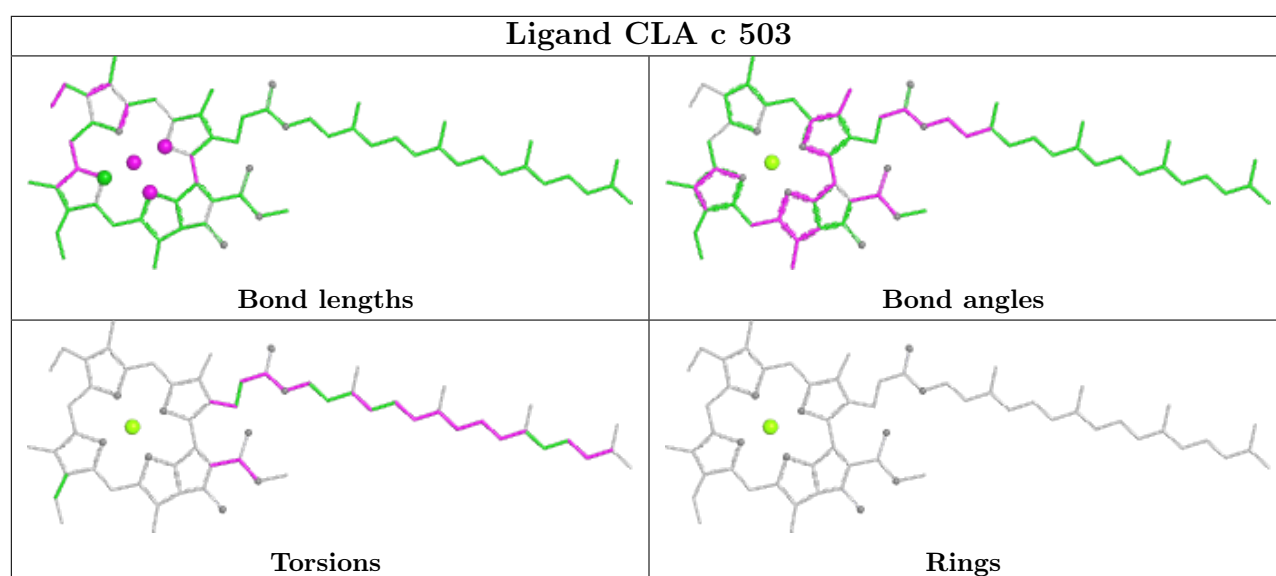
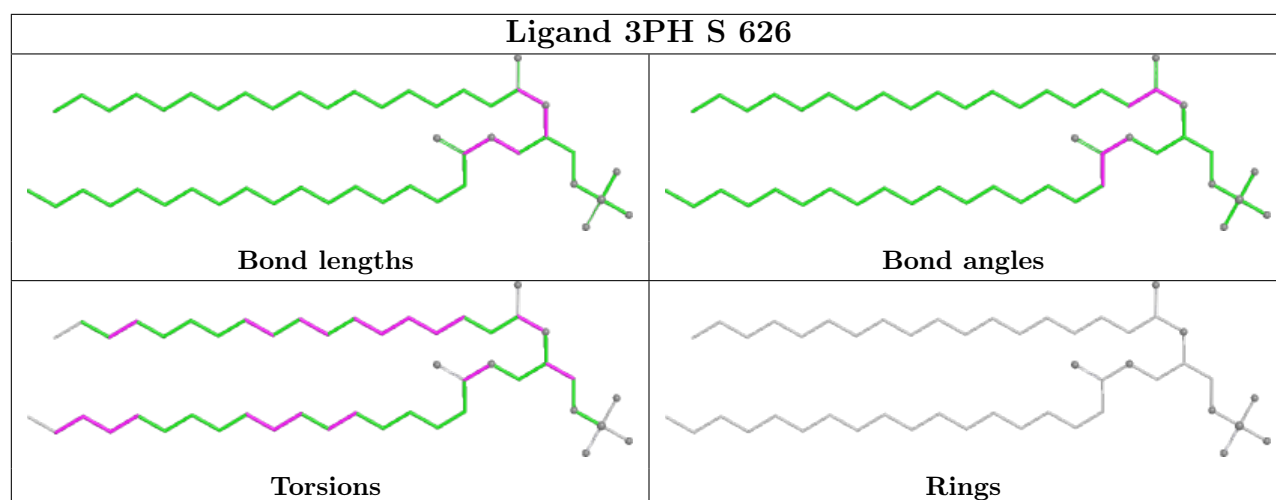


Torsions

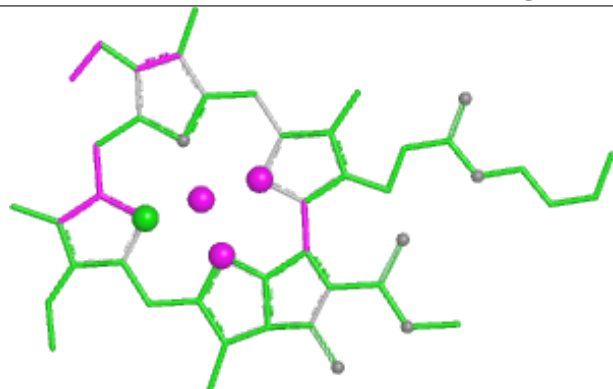


Rings

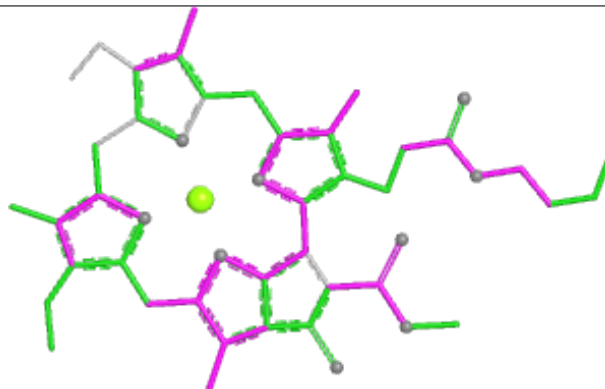




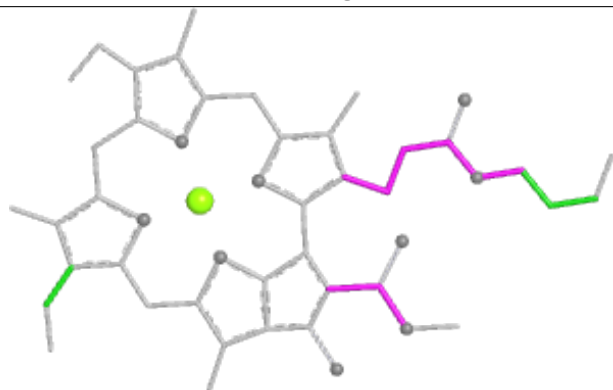
Ligand CLA r 604



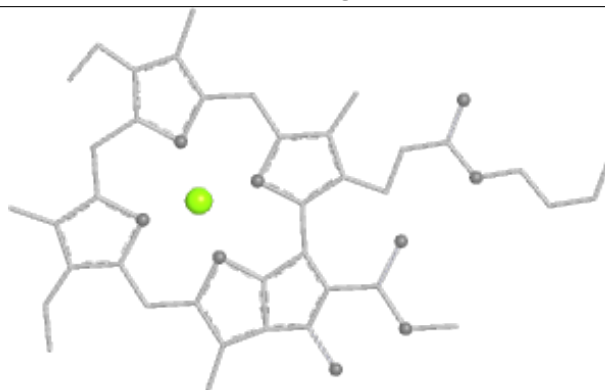
Bond lengths



Bond angles

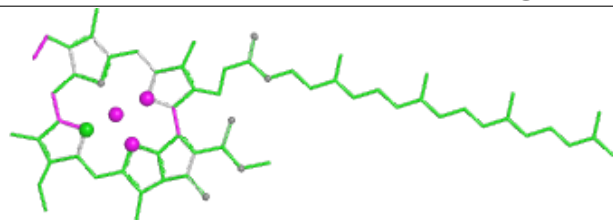


Torsions

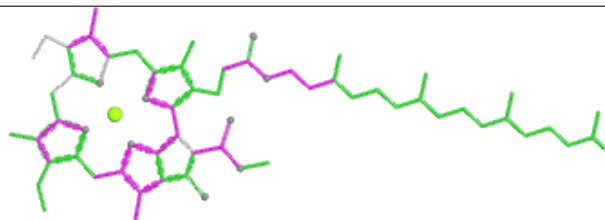


Rings

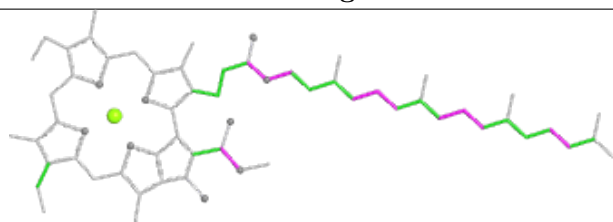
Ligand CLA b 614



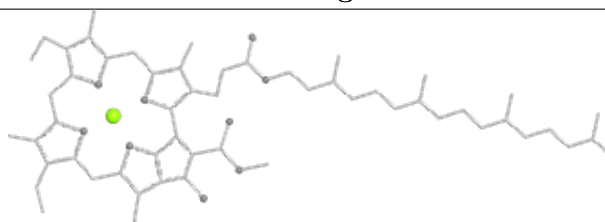
Bond lengths



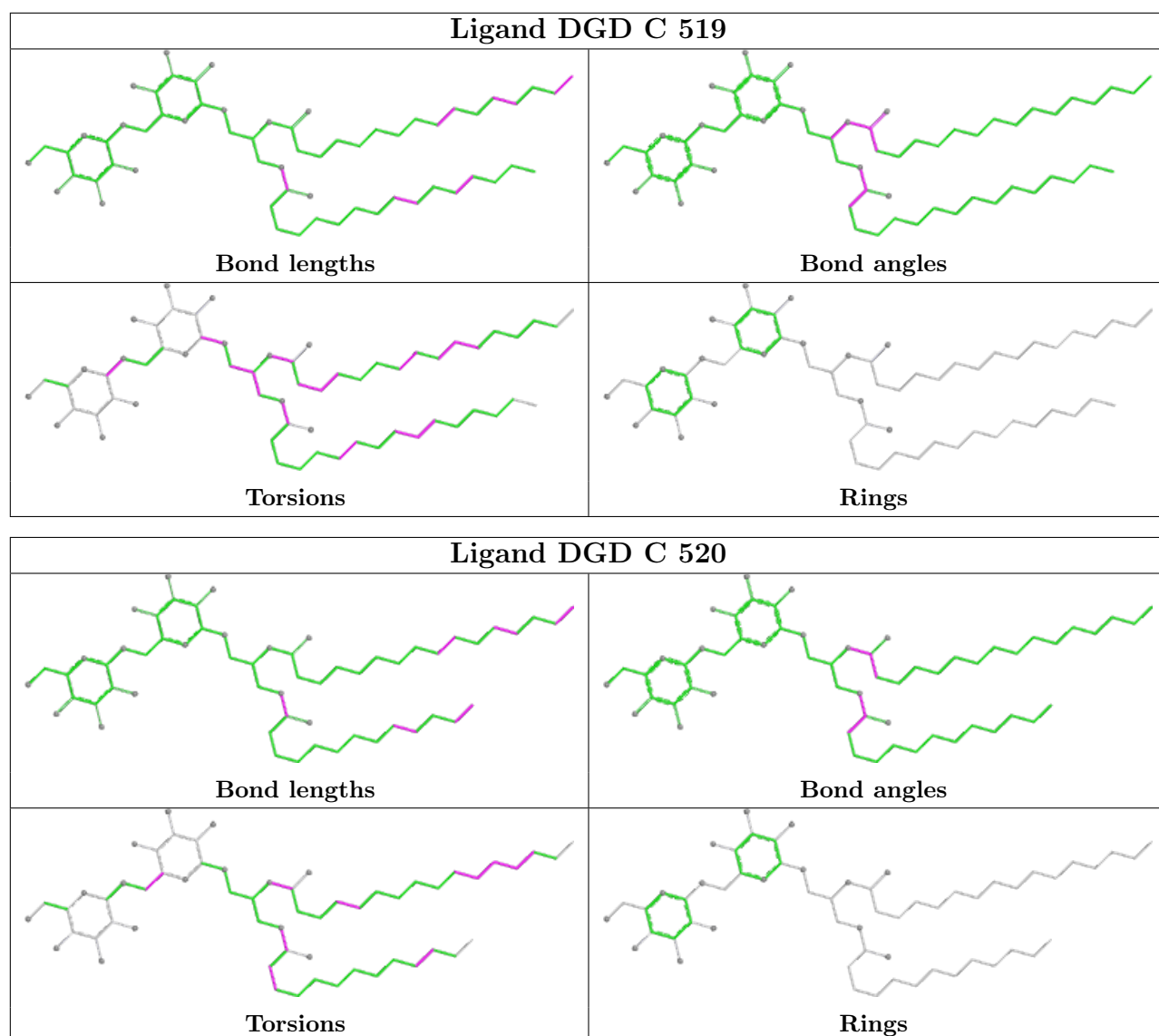
Bond angles

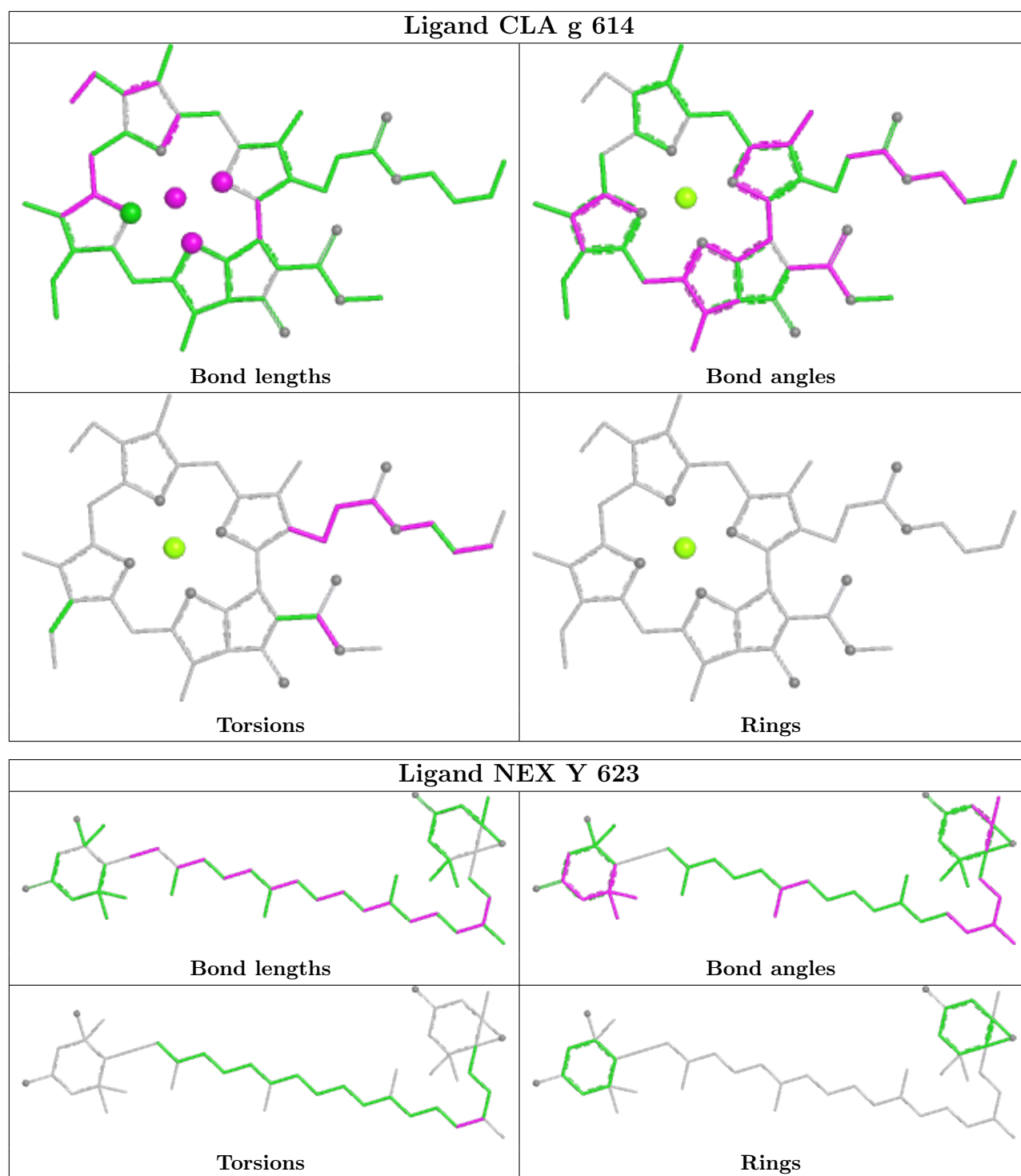


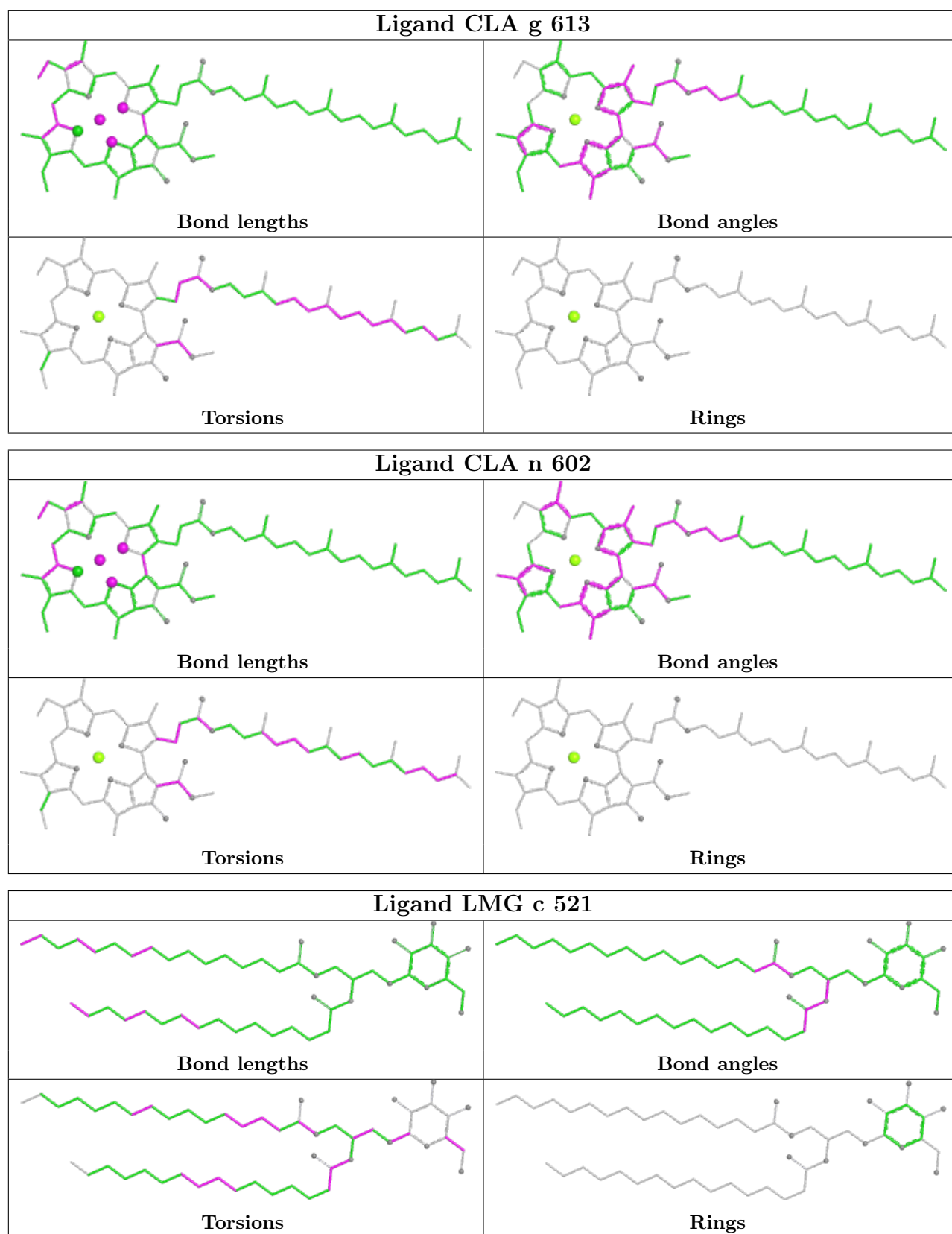
Torsions

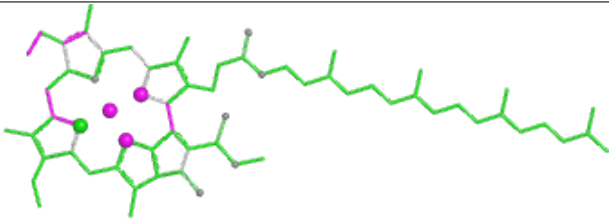
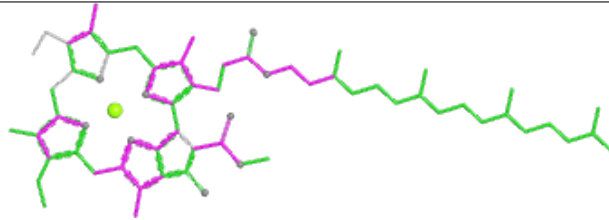
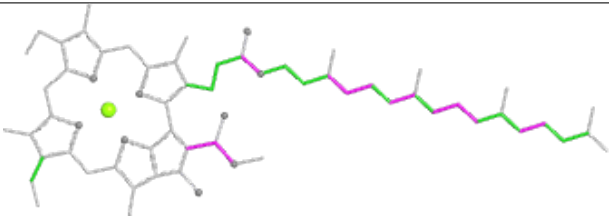
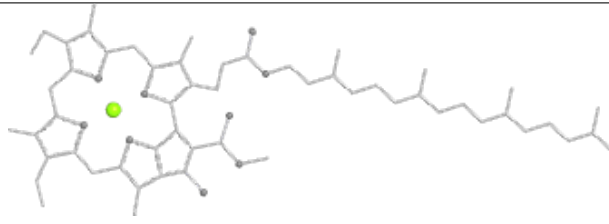


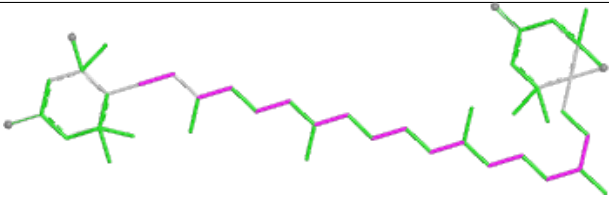
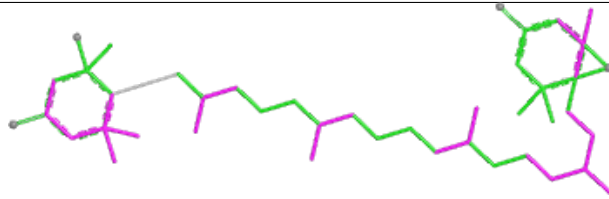
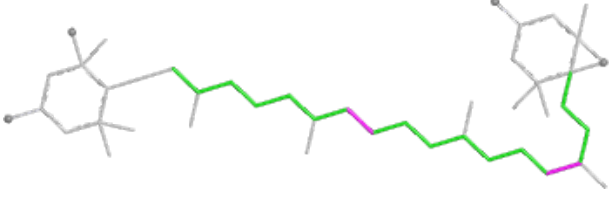
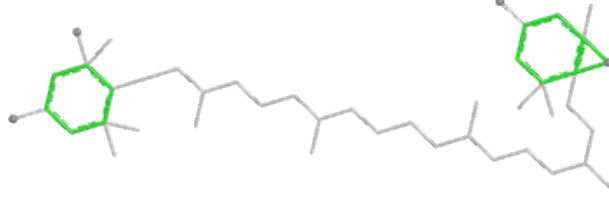
Rings

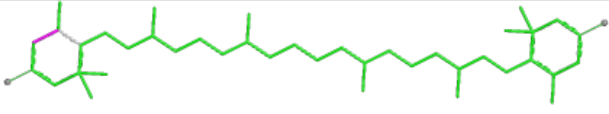
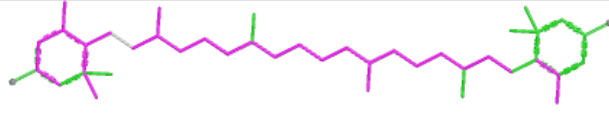
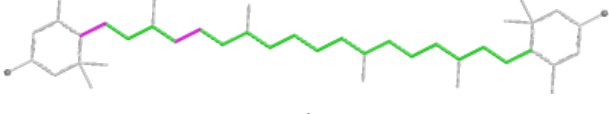
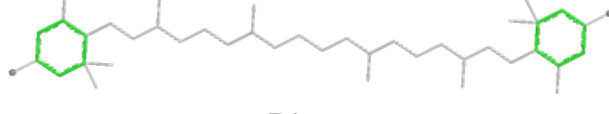


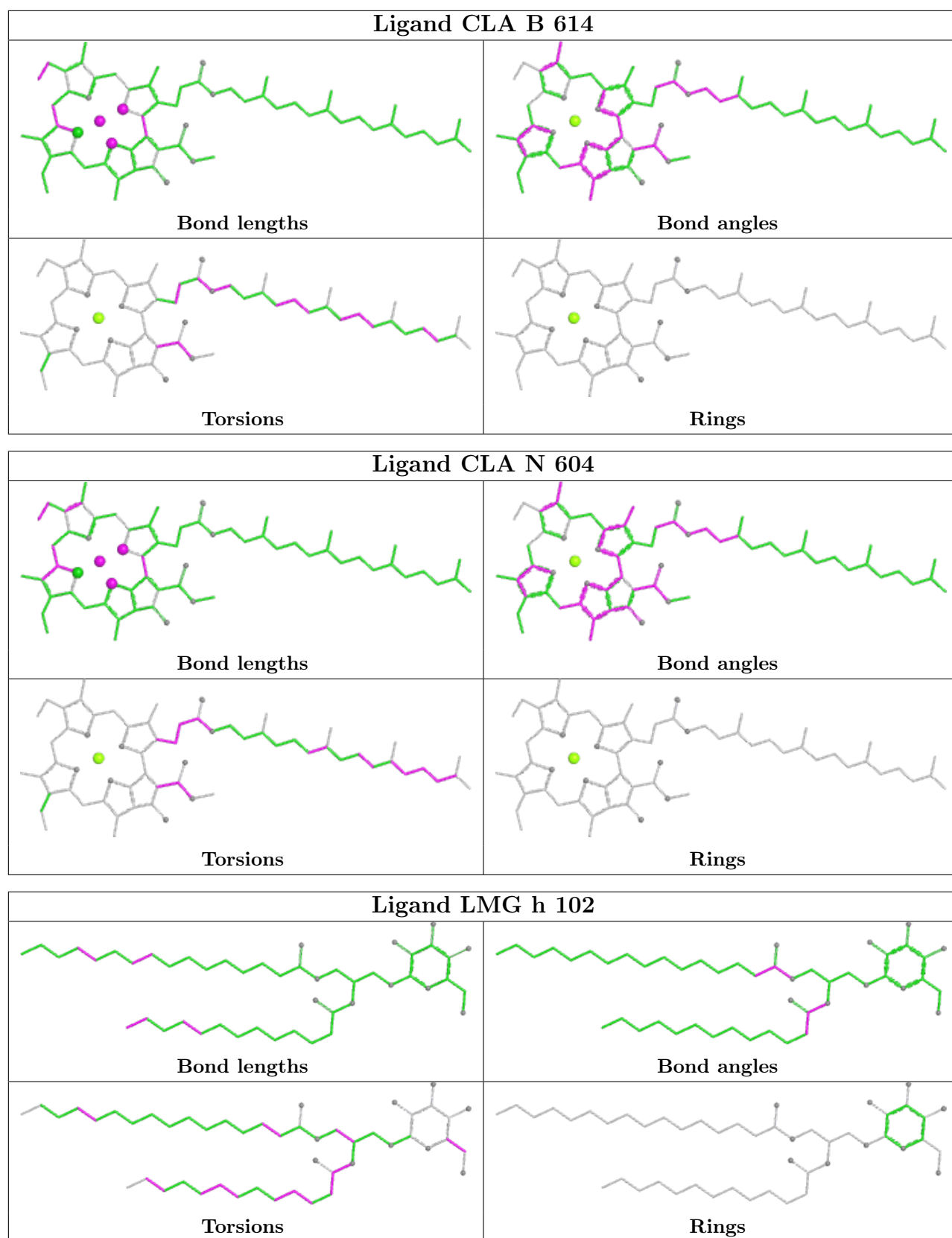


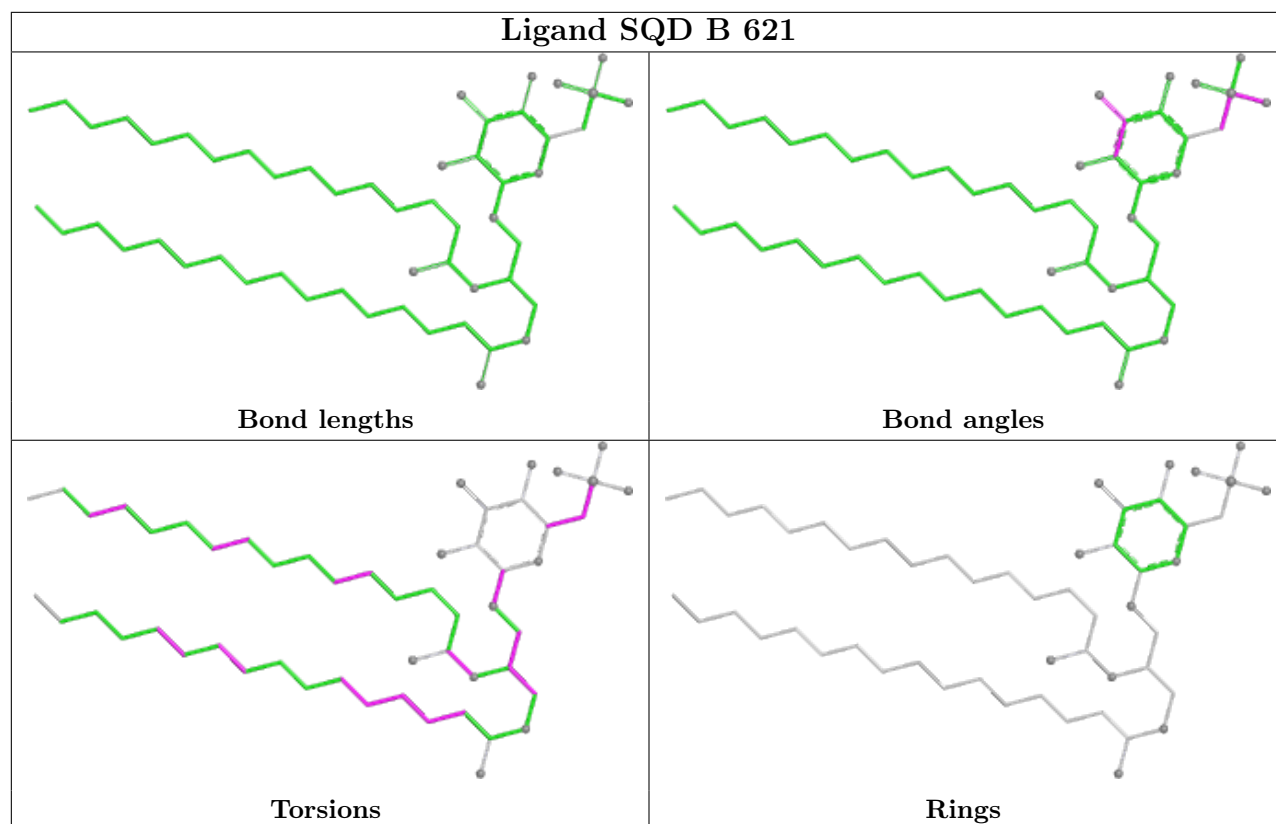
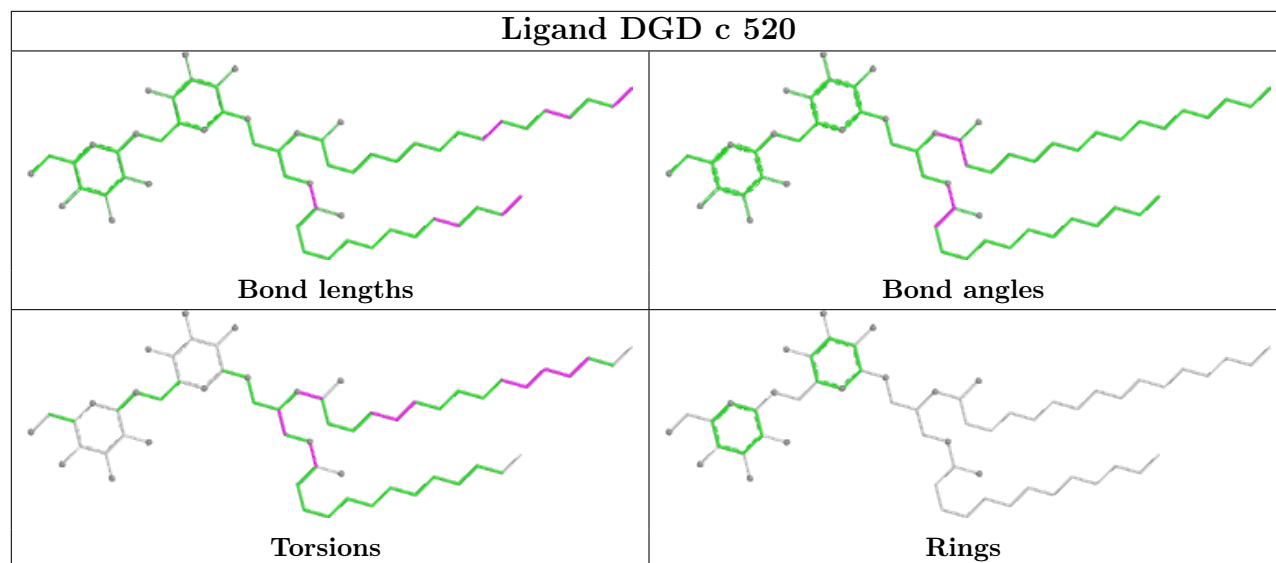


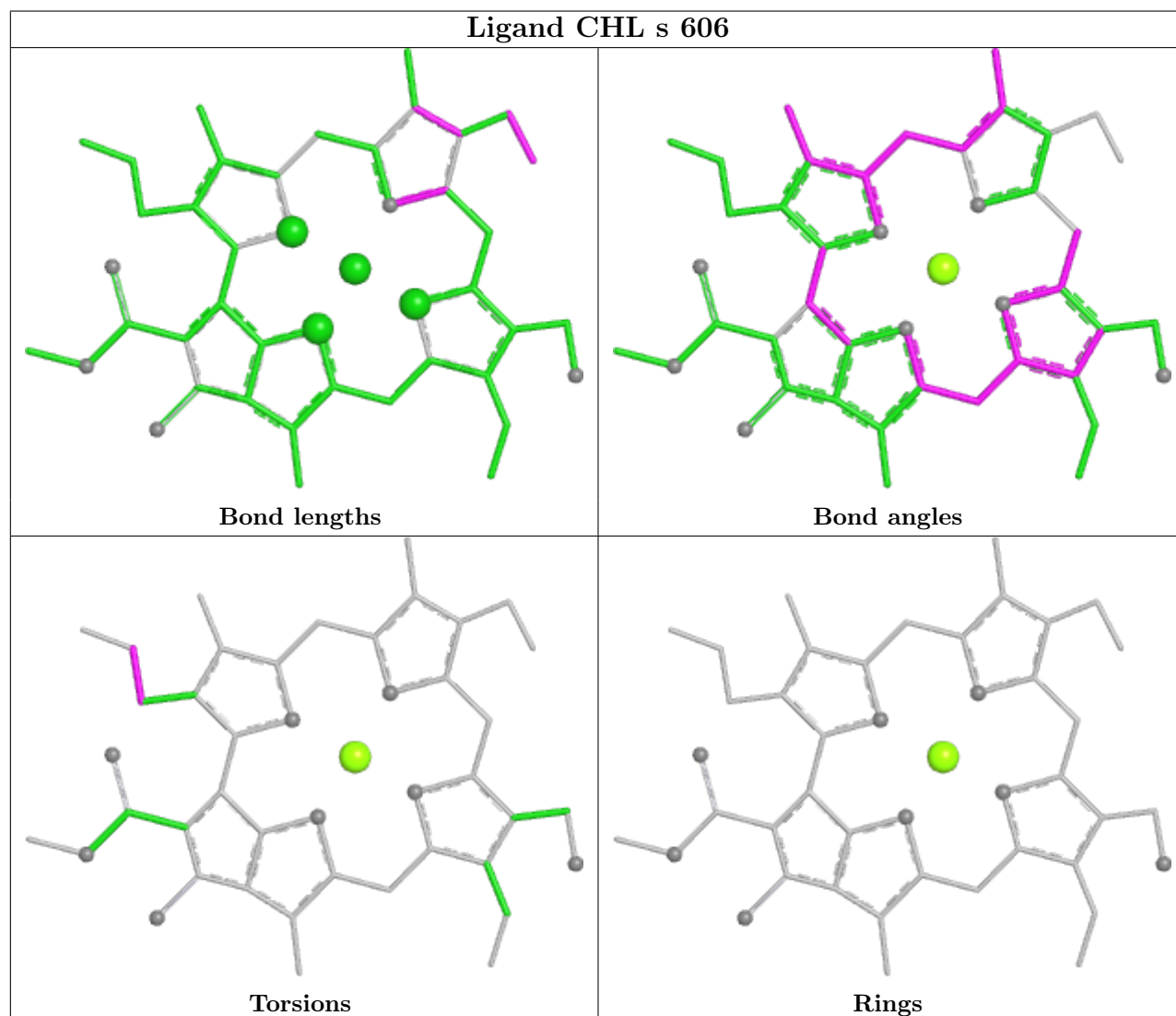
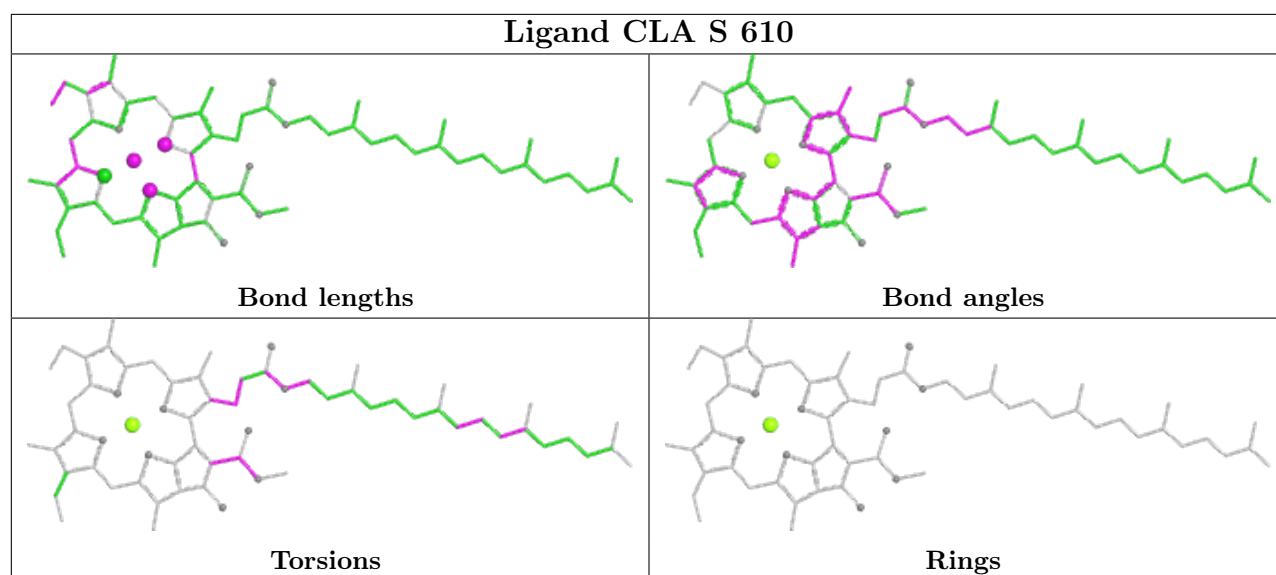
Ligand CLA C 504	
	
Bond lengths	Bond angles
	
Torsions	Rings

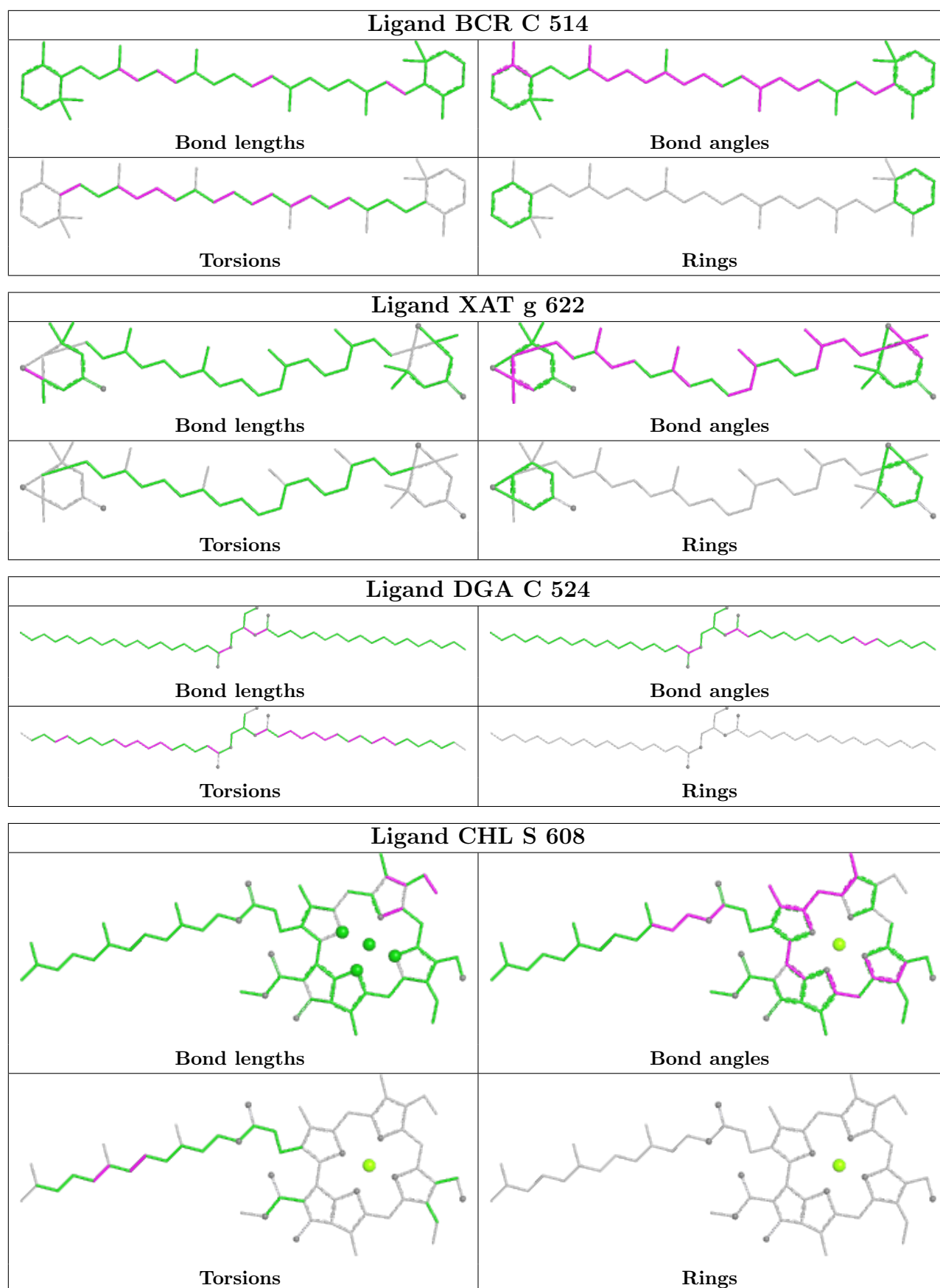
Ligand NEX g 623	
	
Bond lengths	Bond angles
	
Torsions	Rings

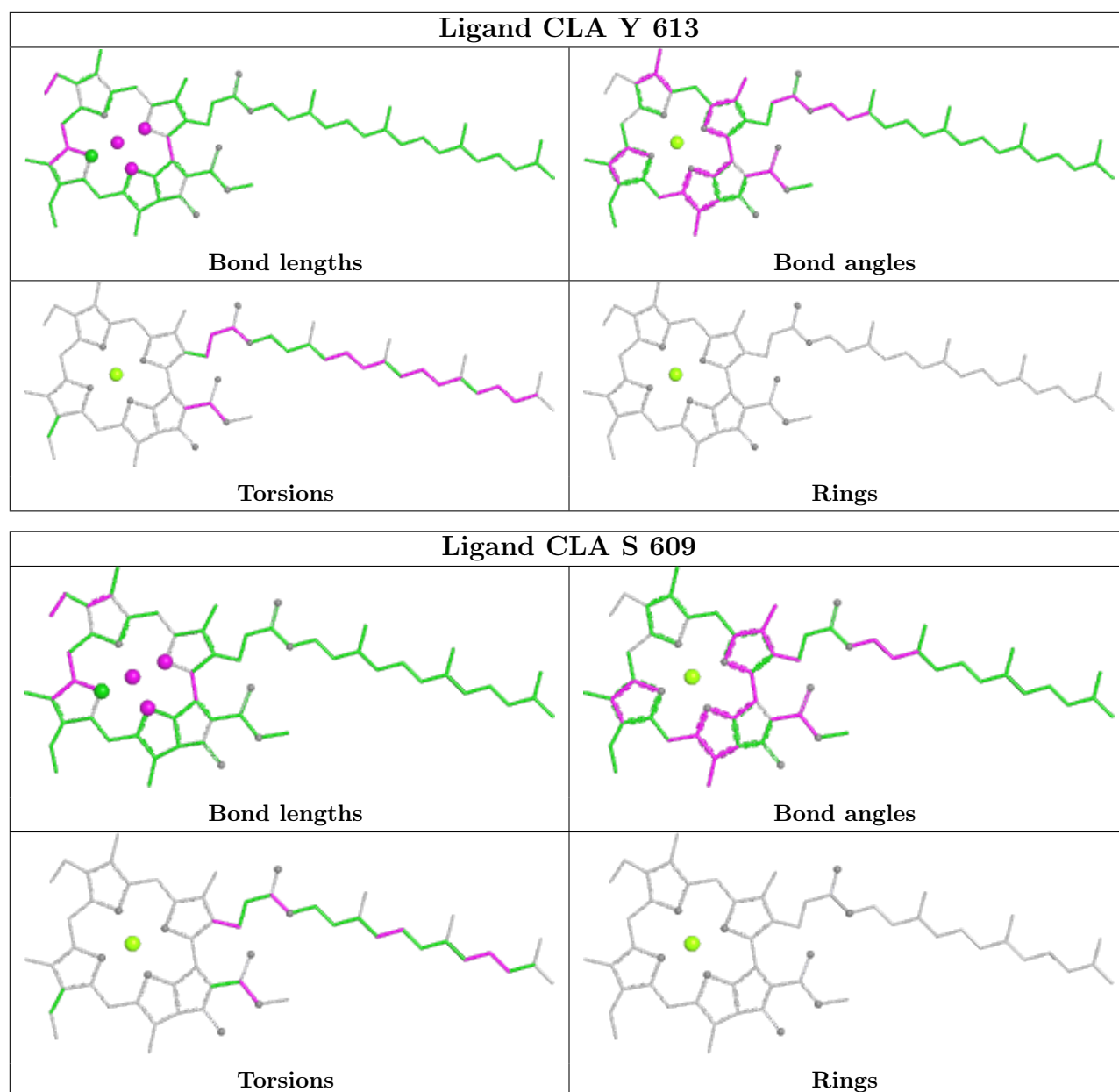
Ligand LUT s 620	
	
Bond lengths	Bond angles
	
Torsions	Rings

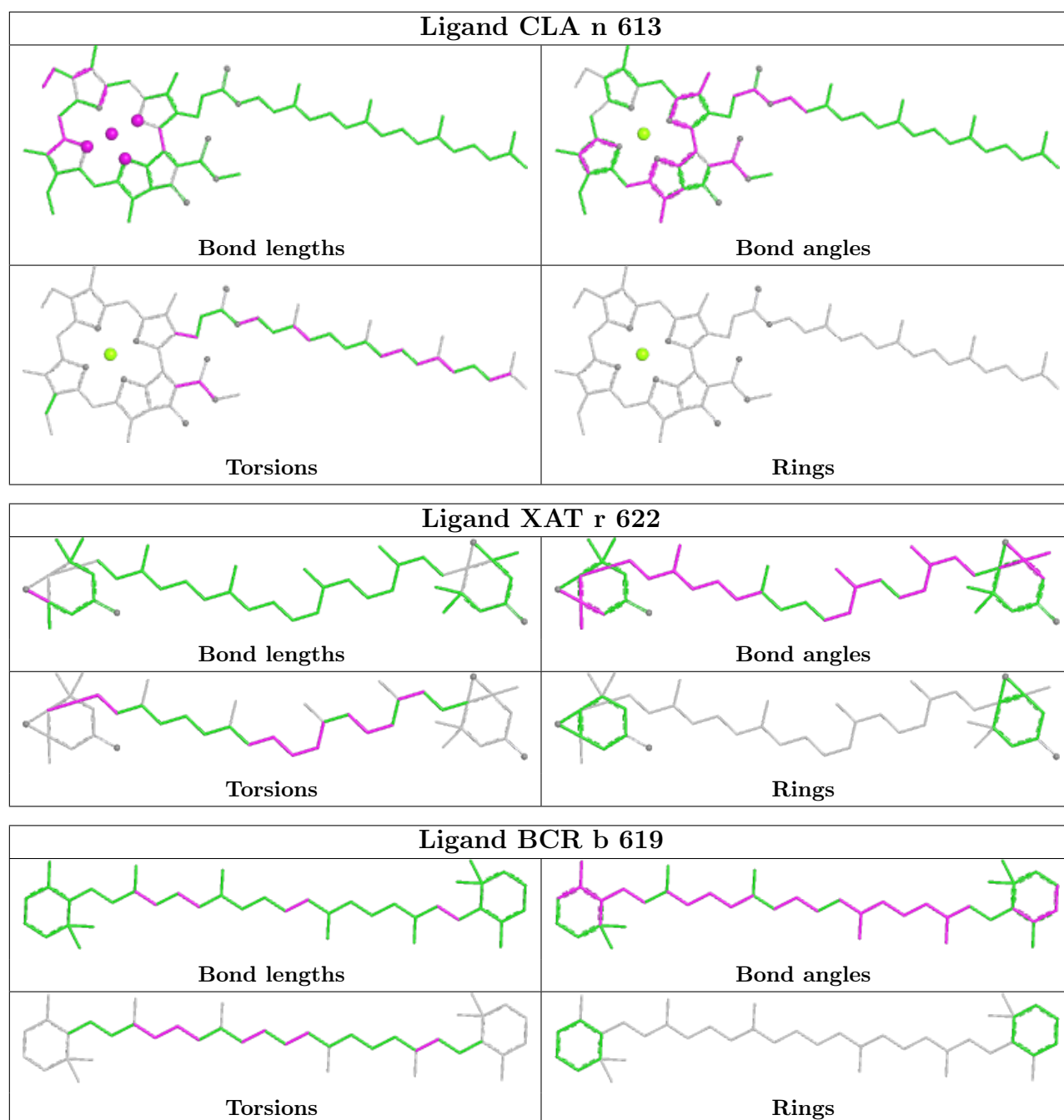


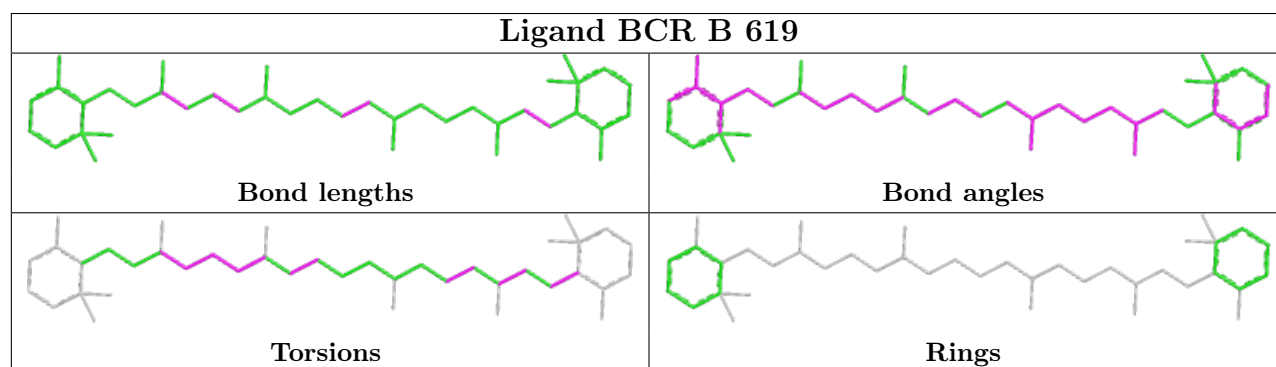
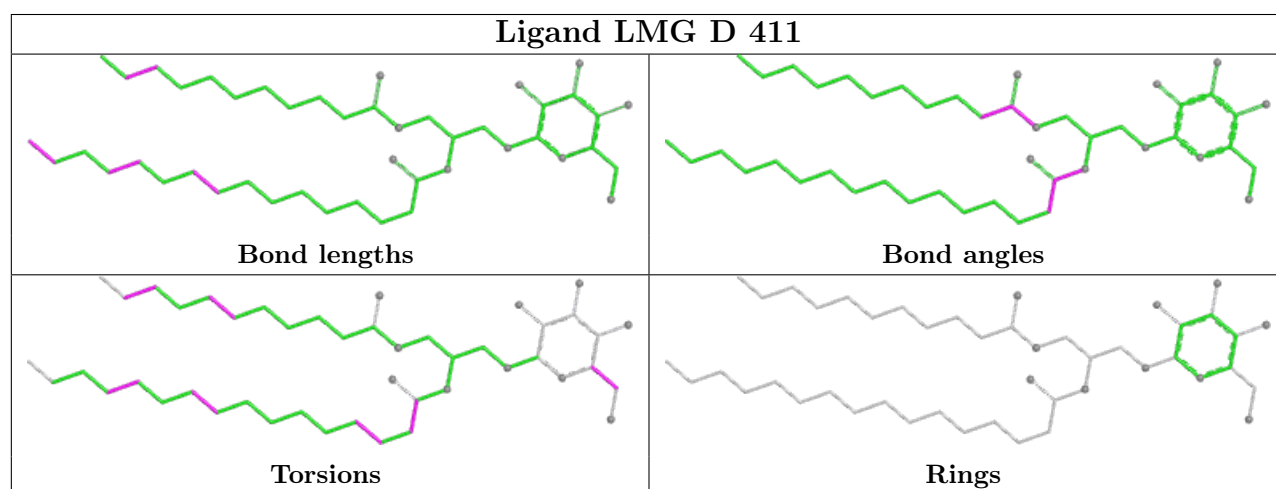
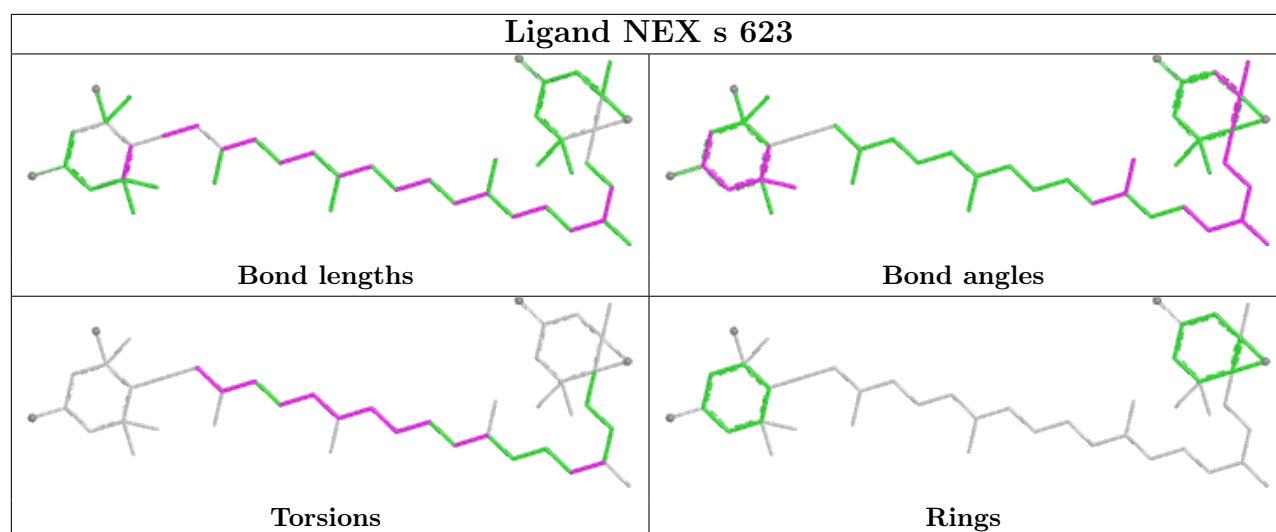


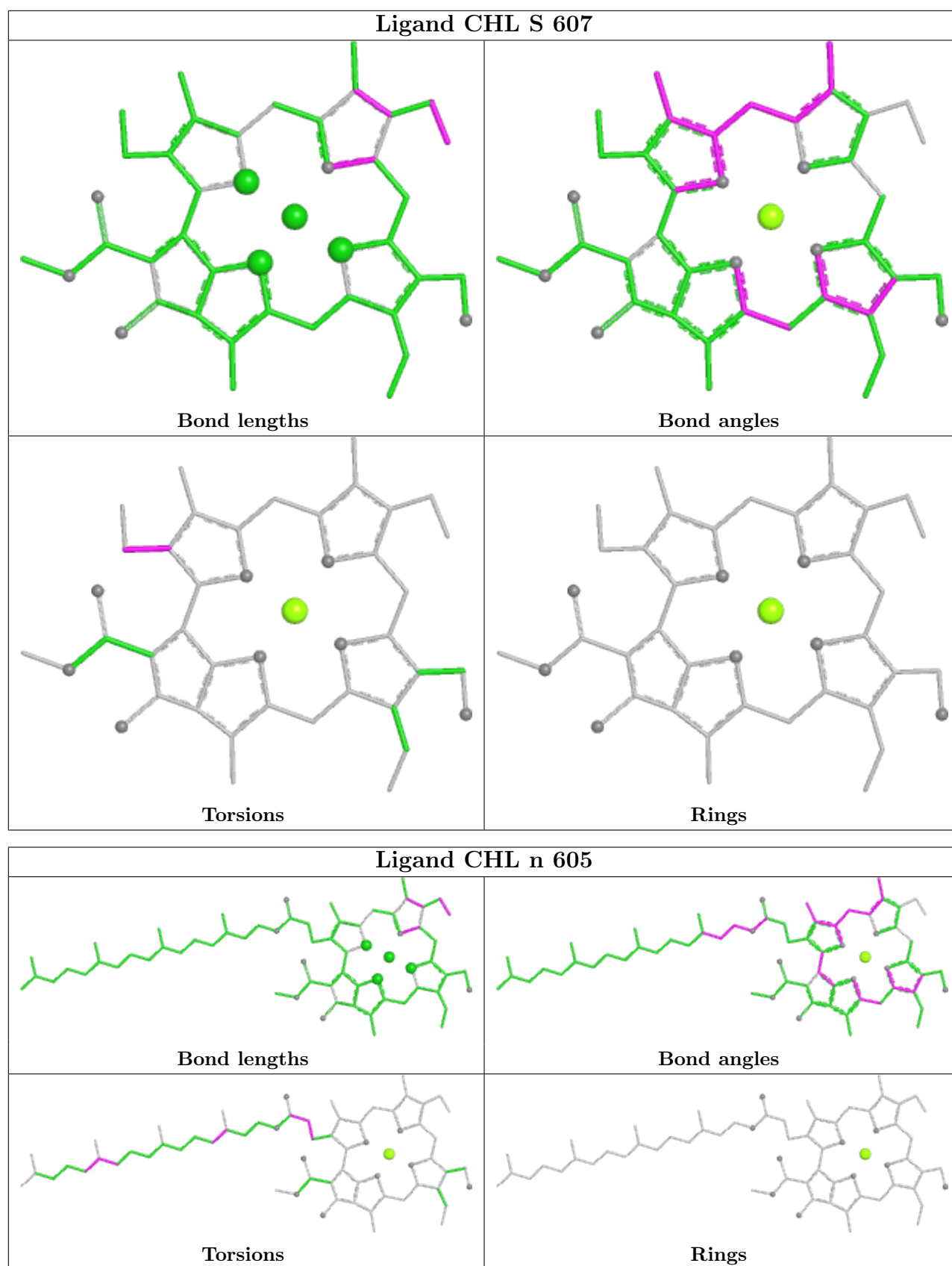


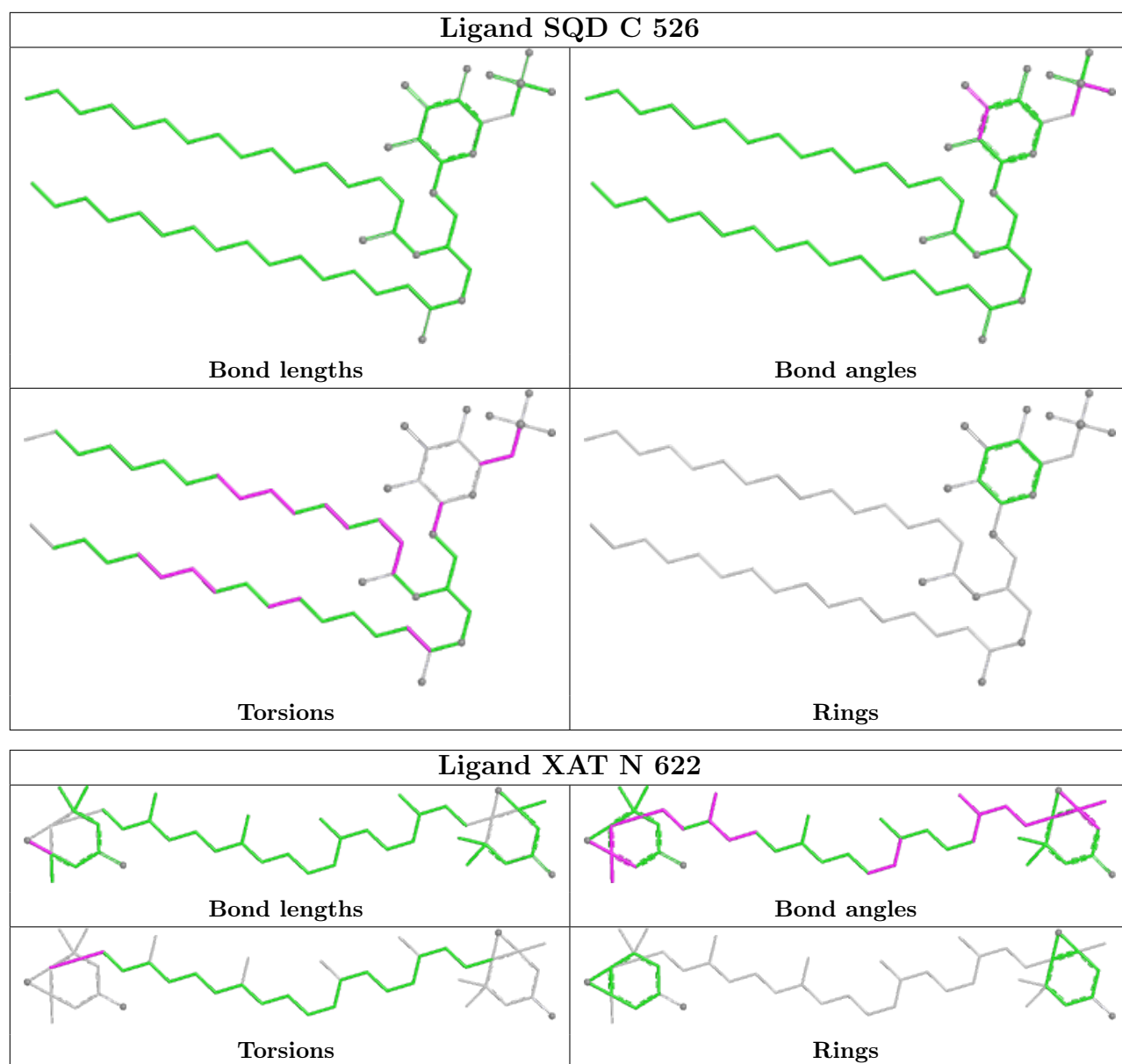


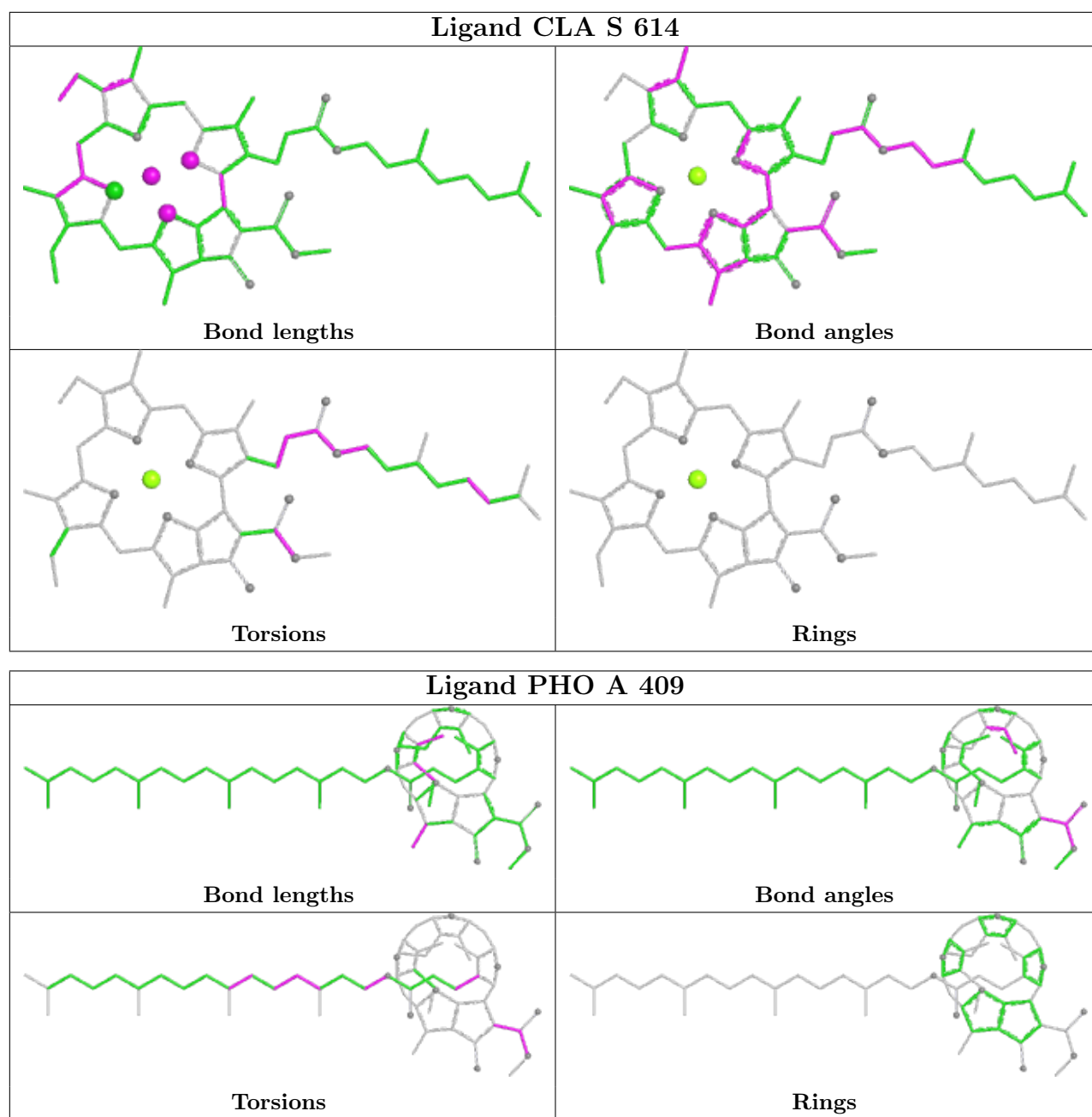


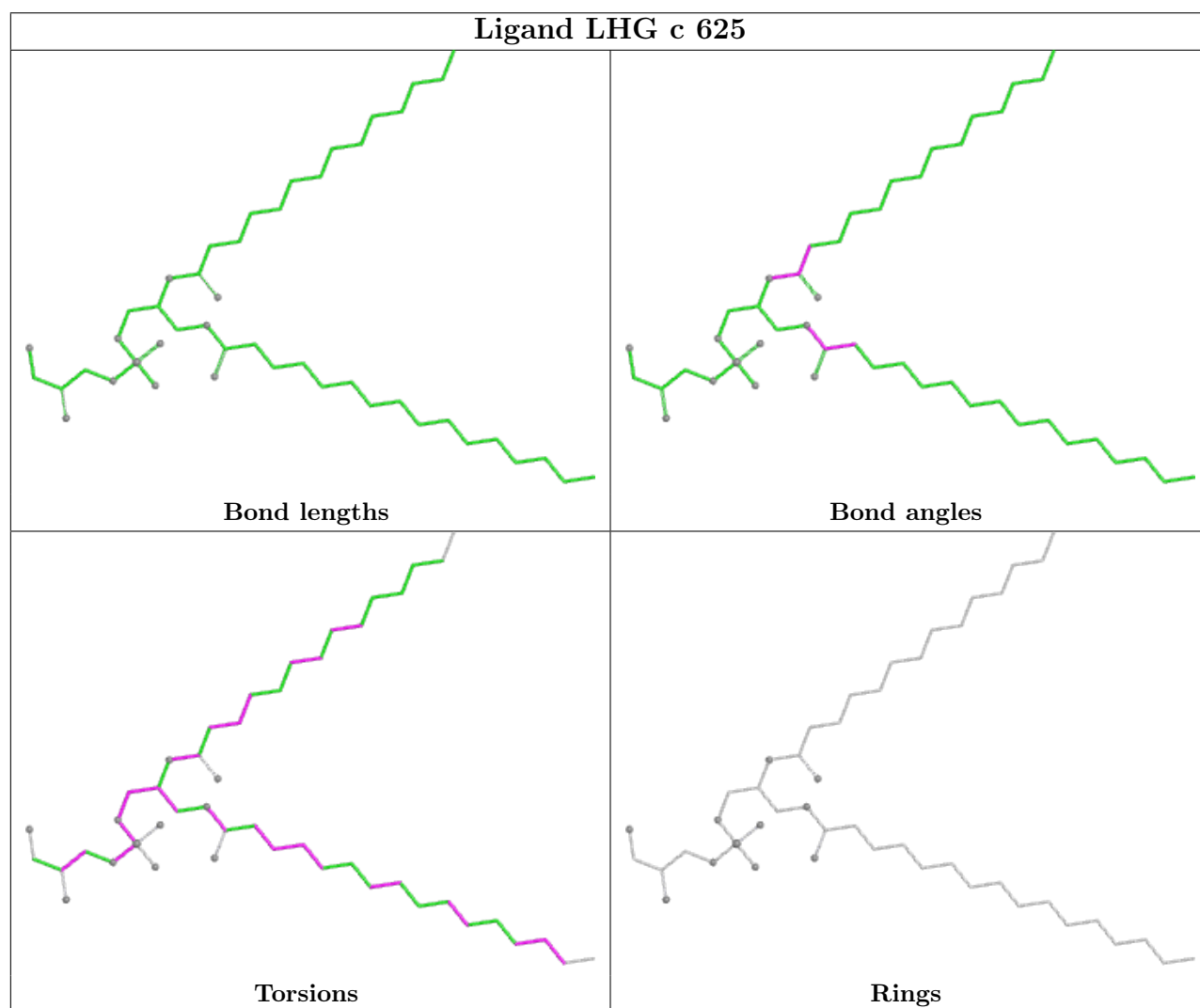
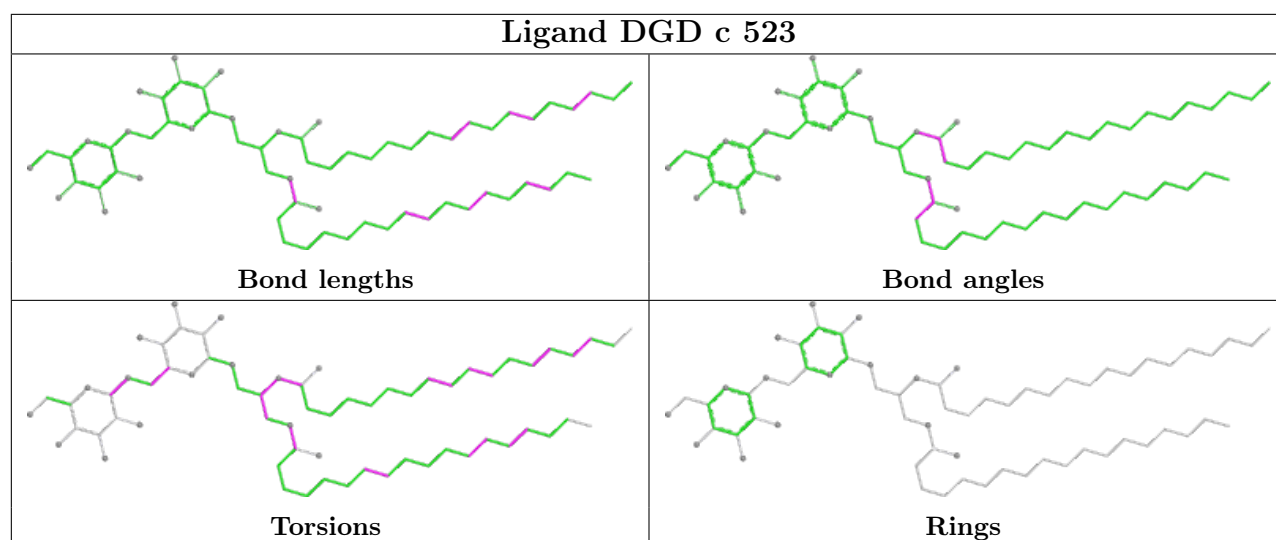


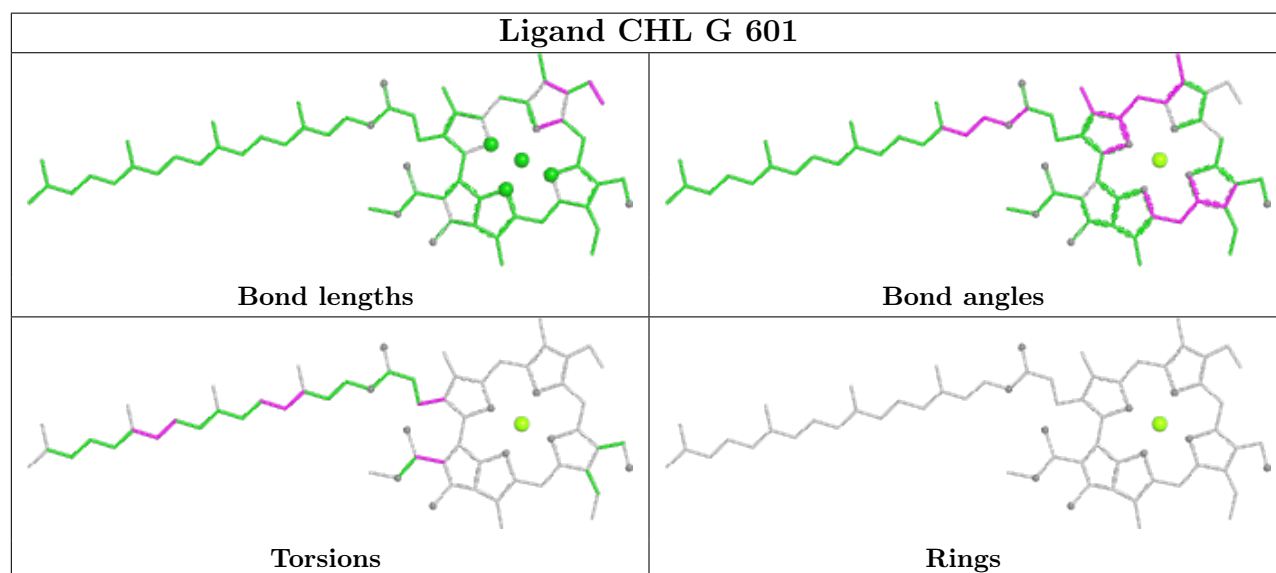
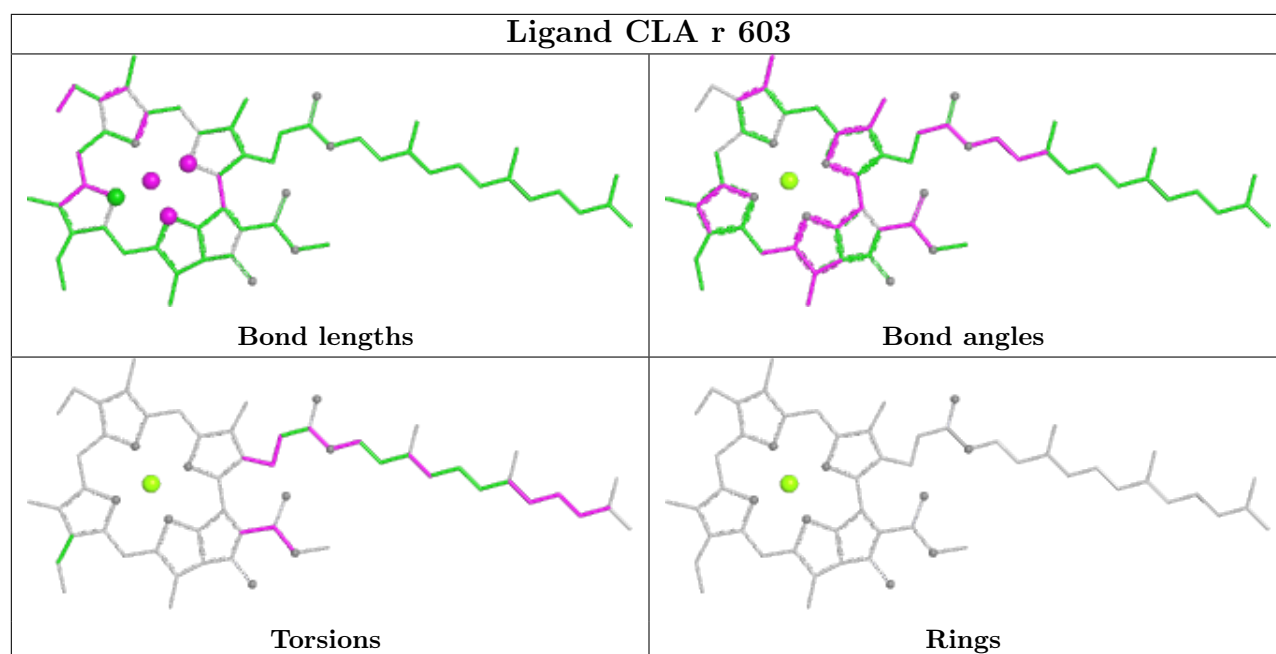


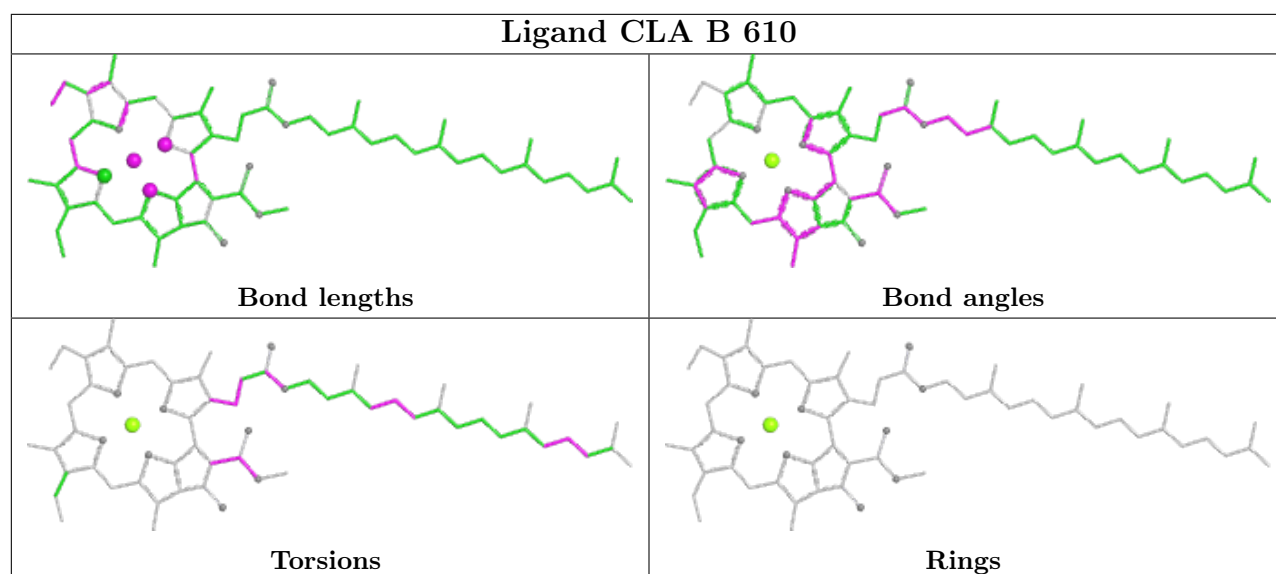
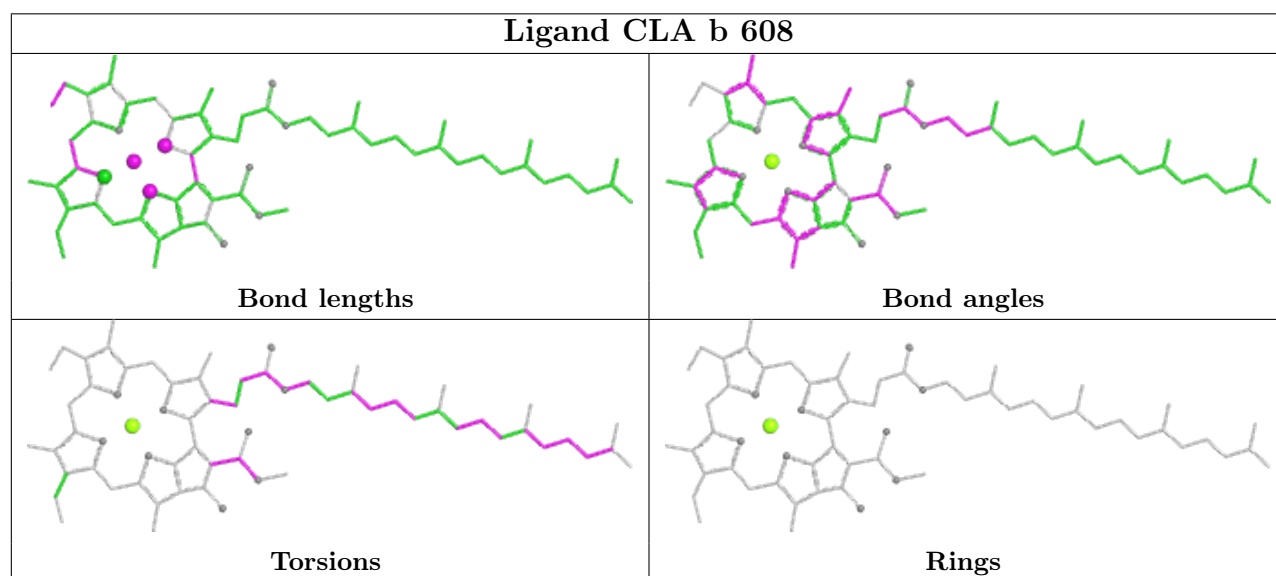
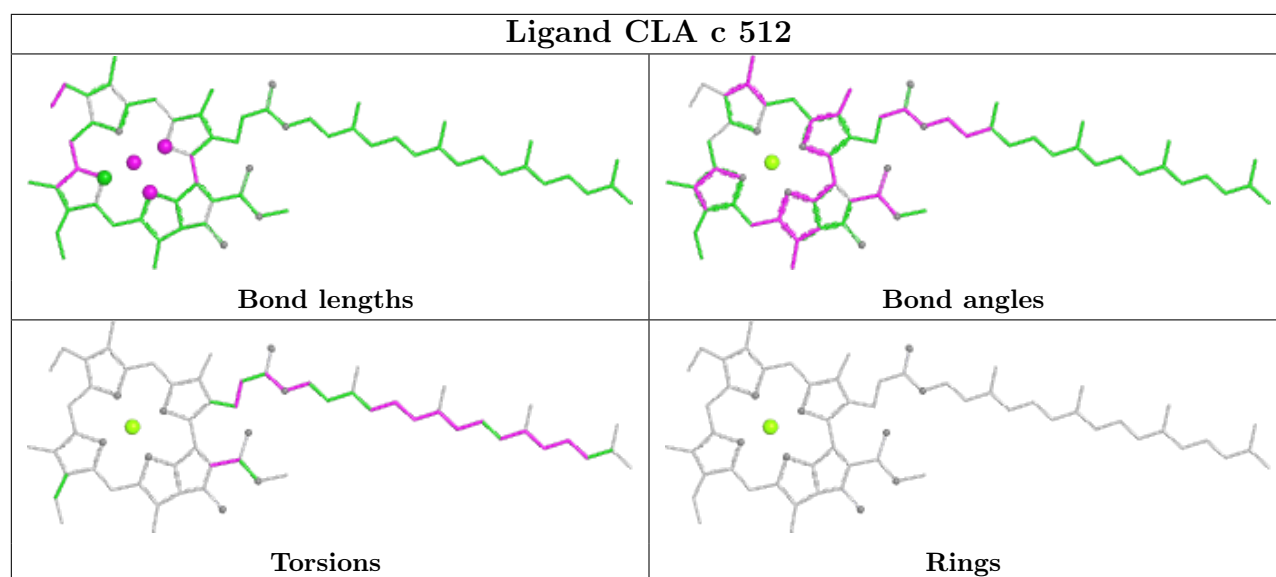


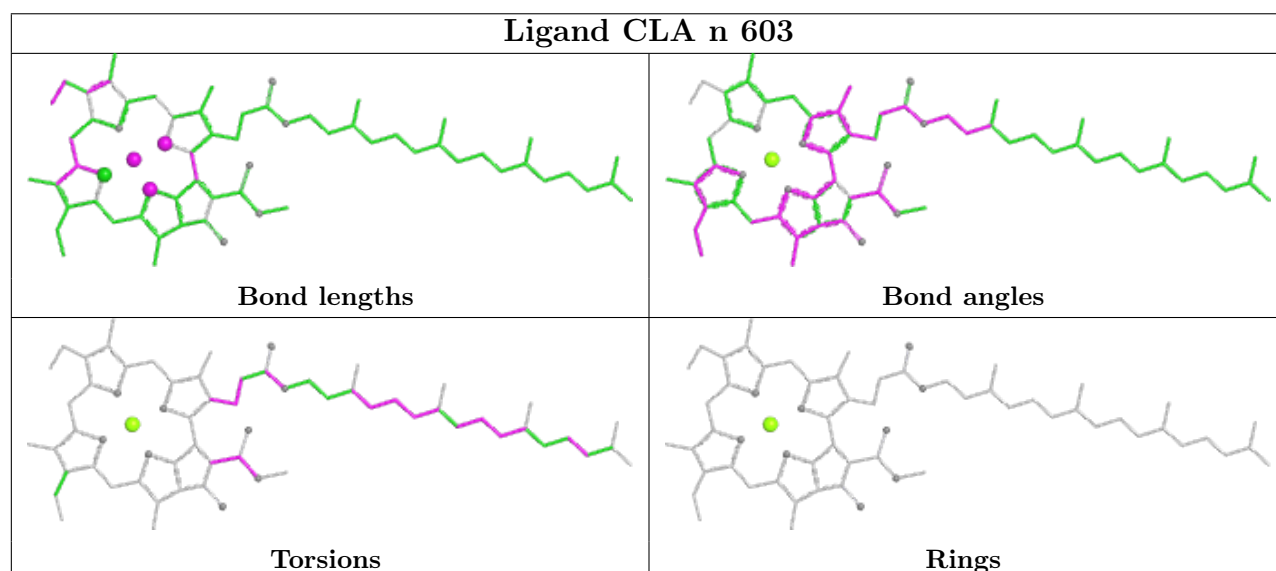
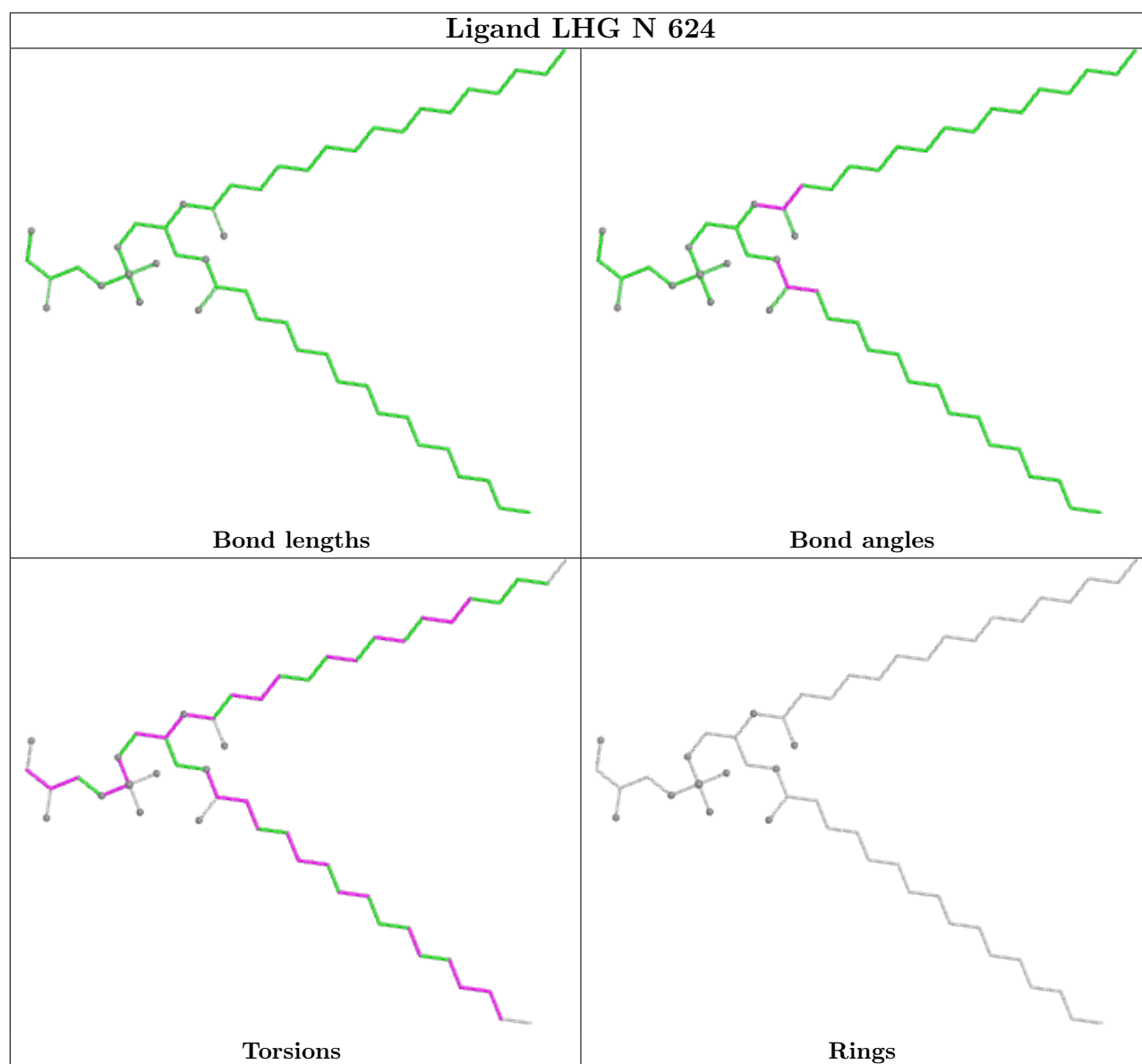


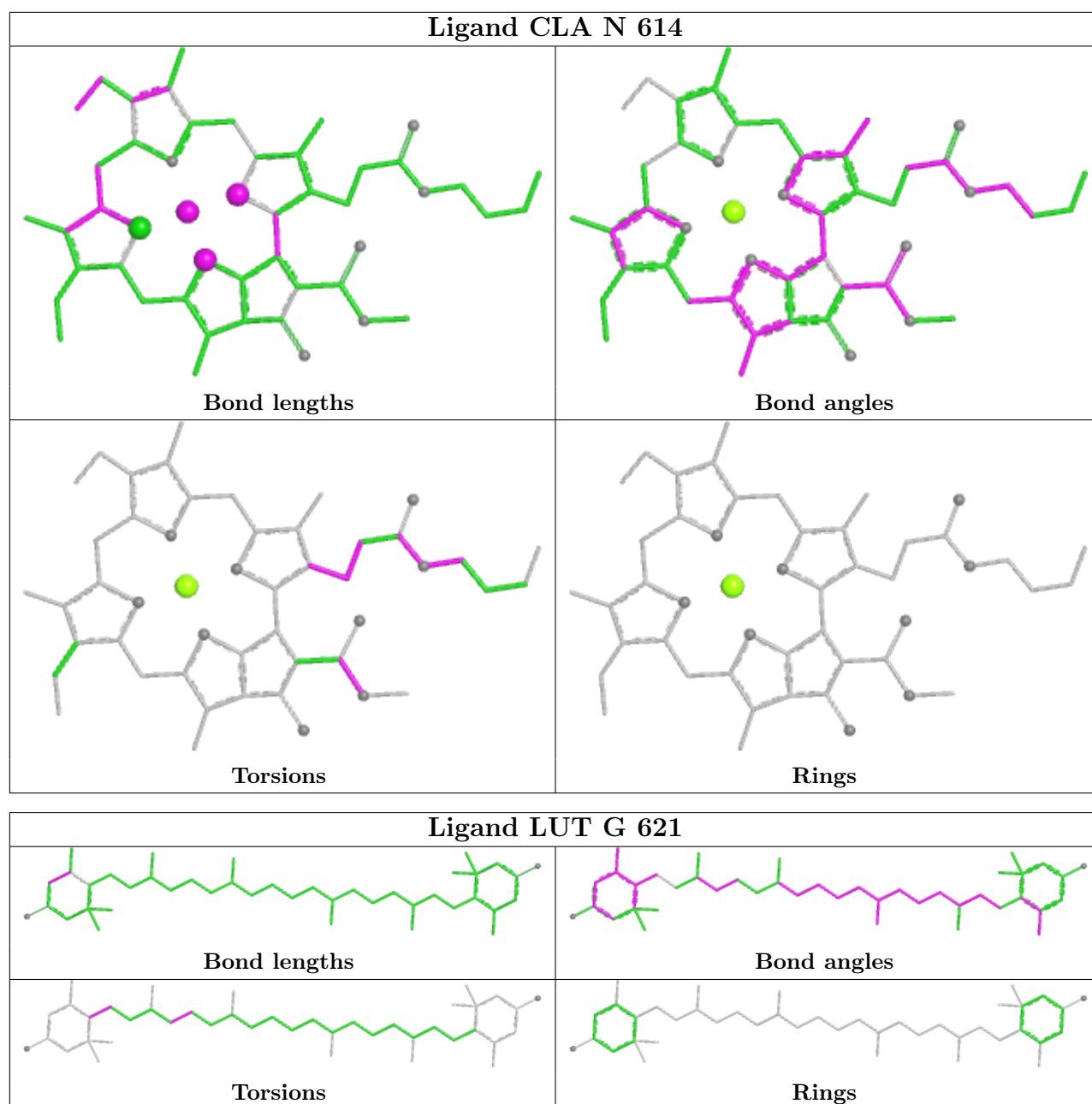




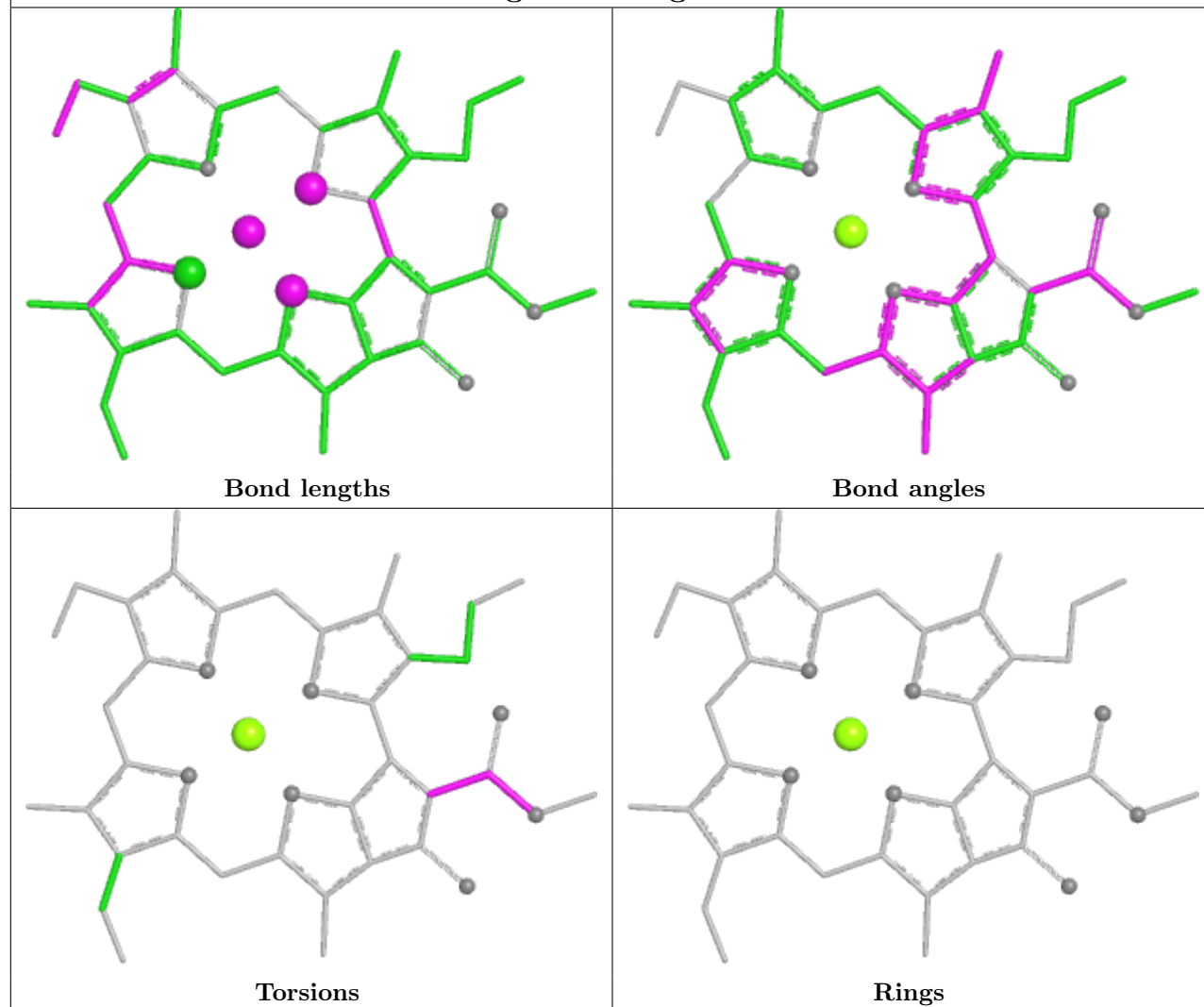




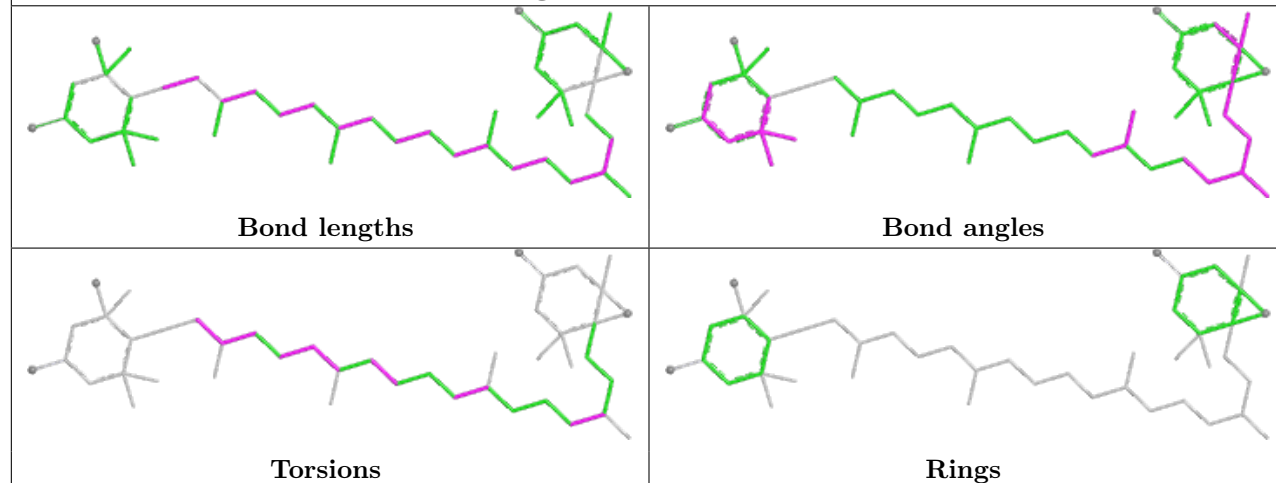


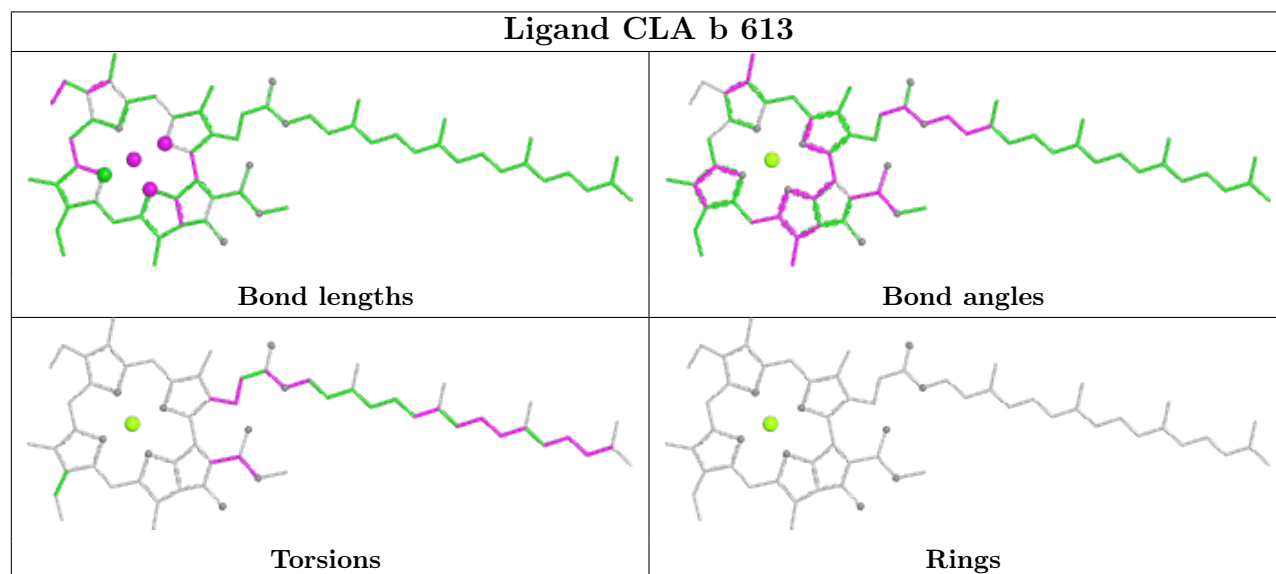
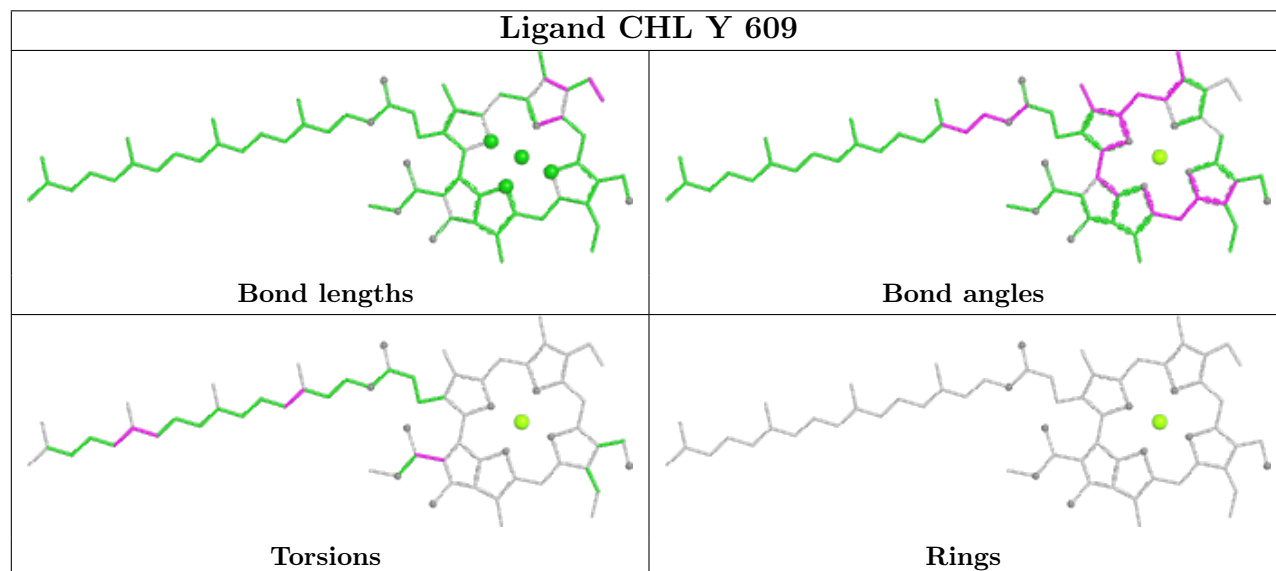
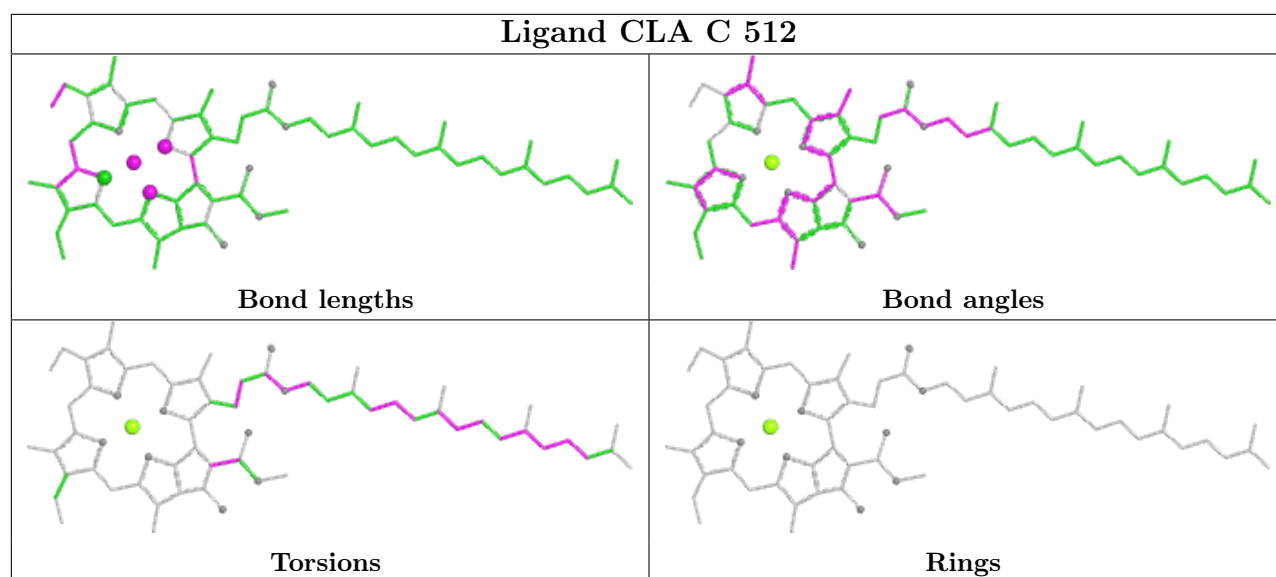


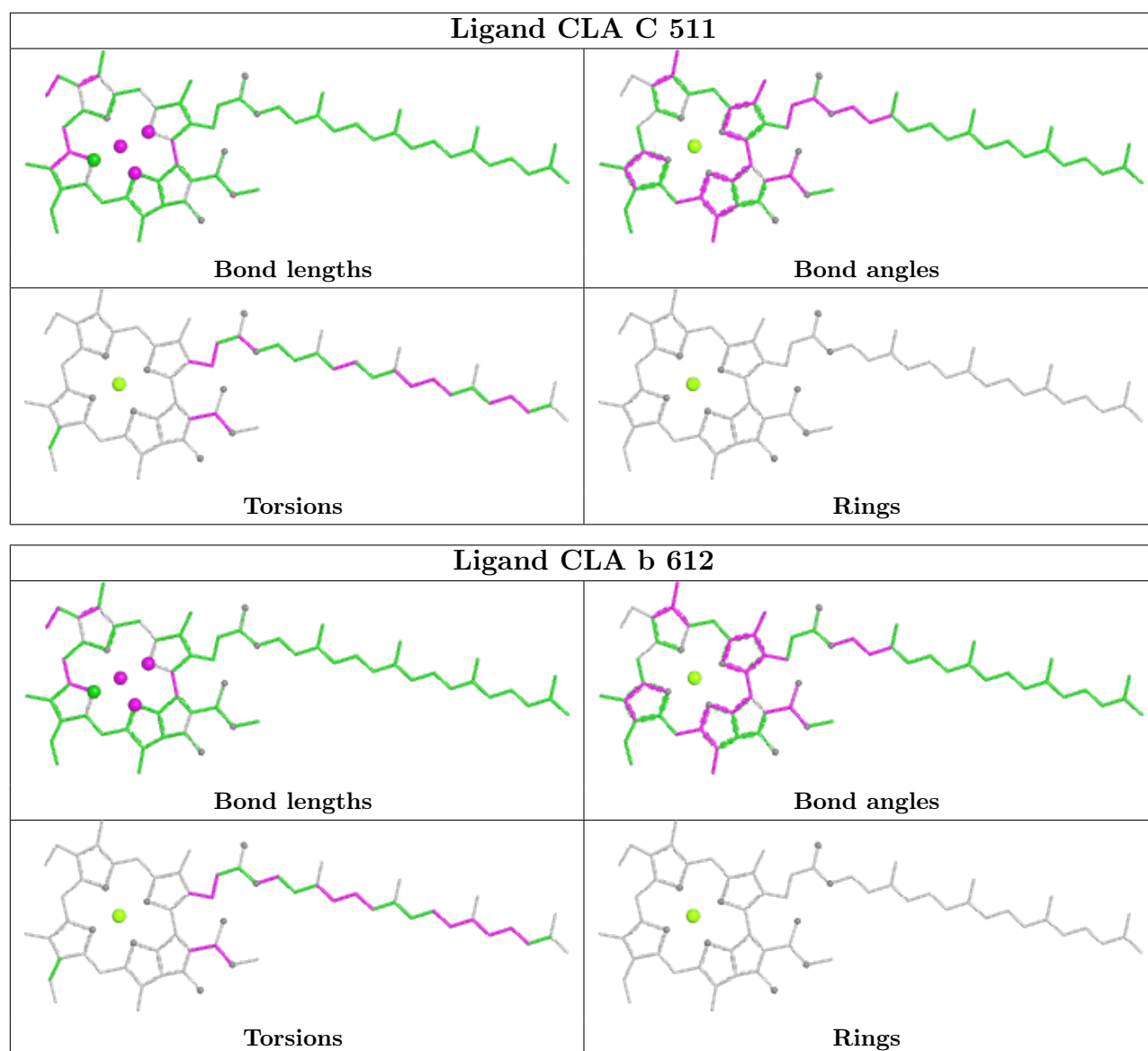
Ligand CLA g 612

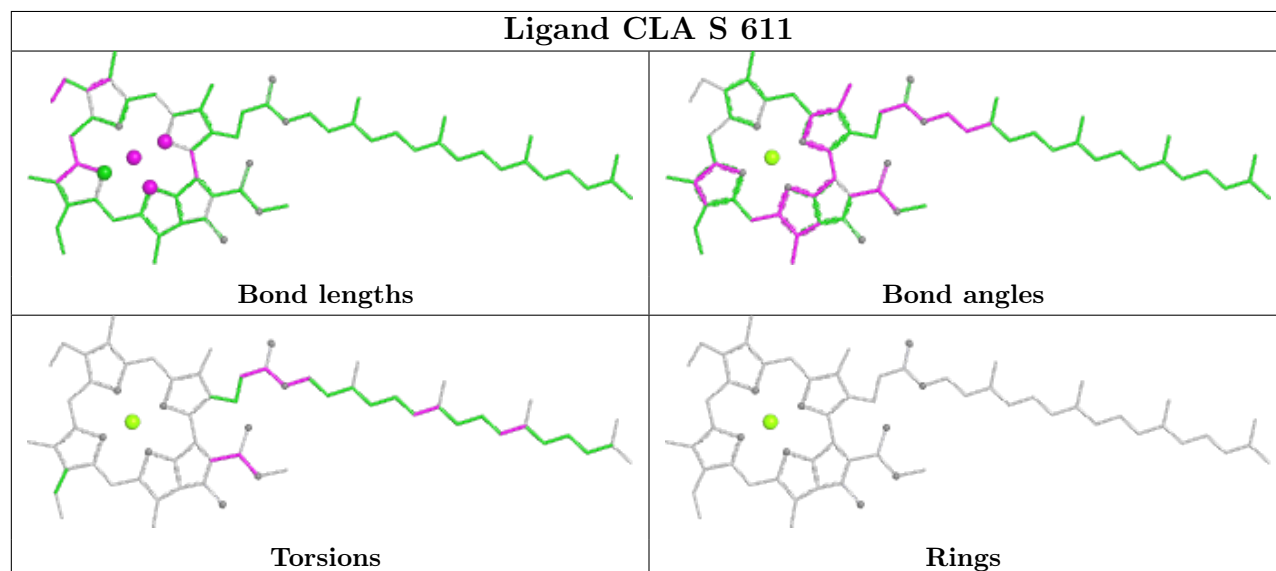
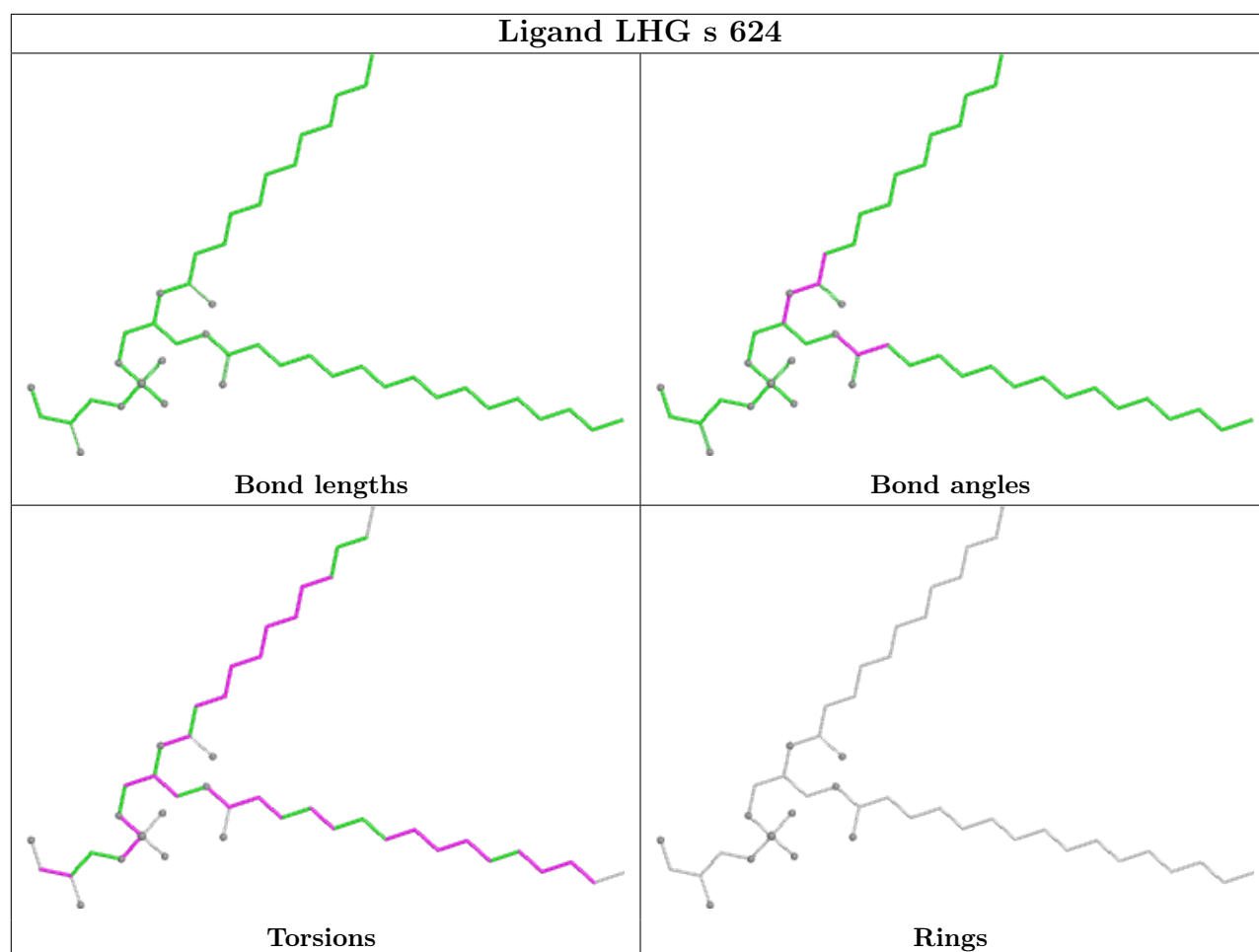


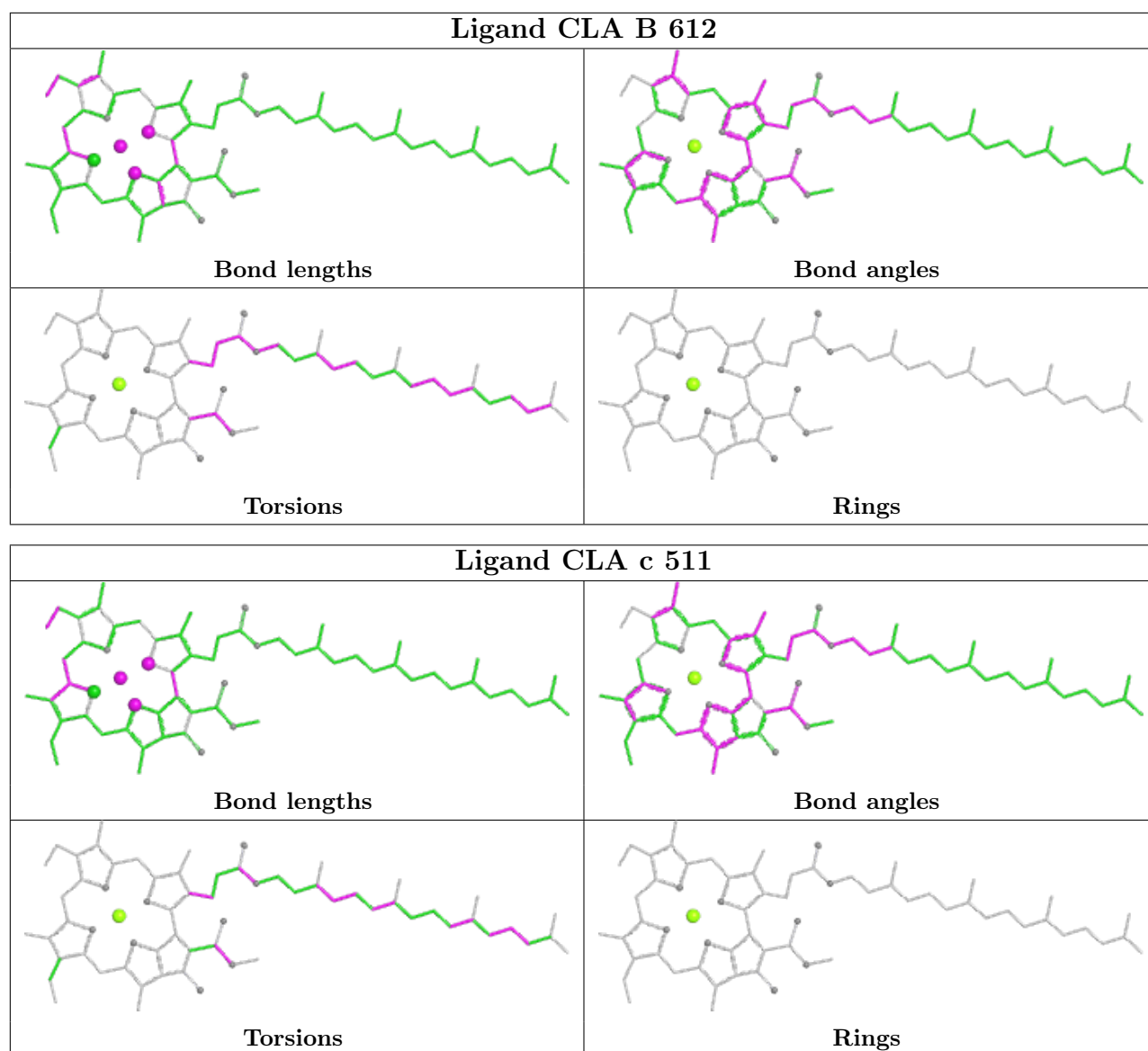
Ligand NEX S 622

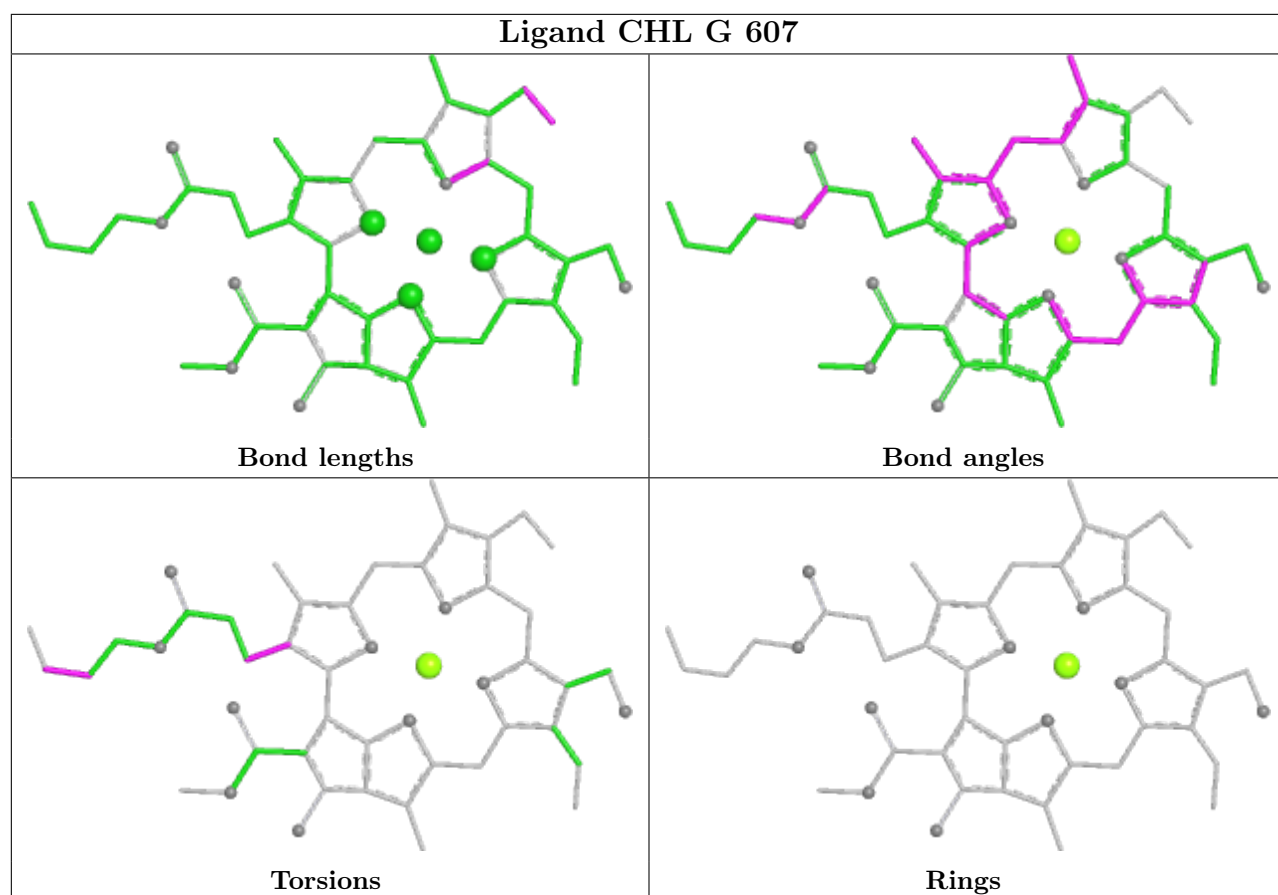


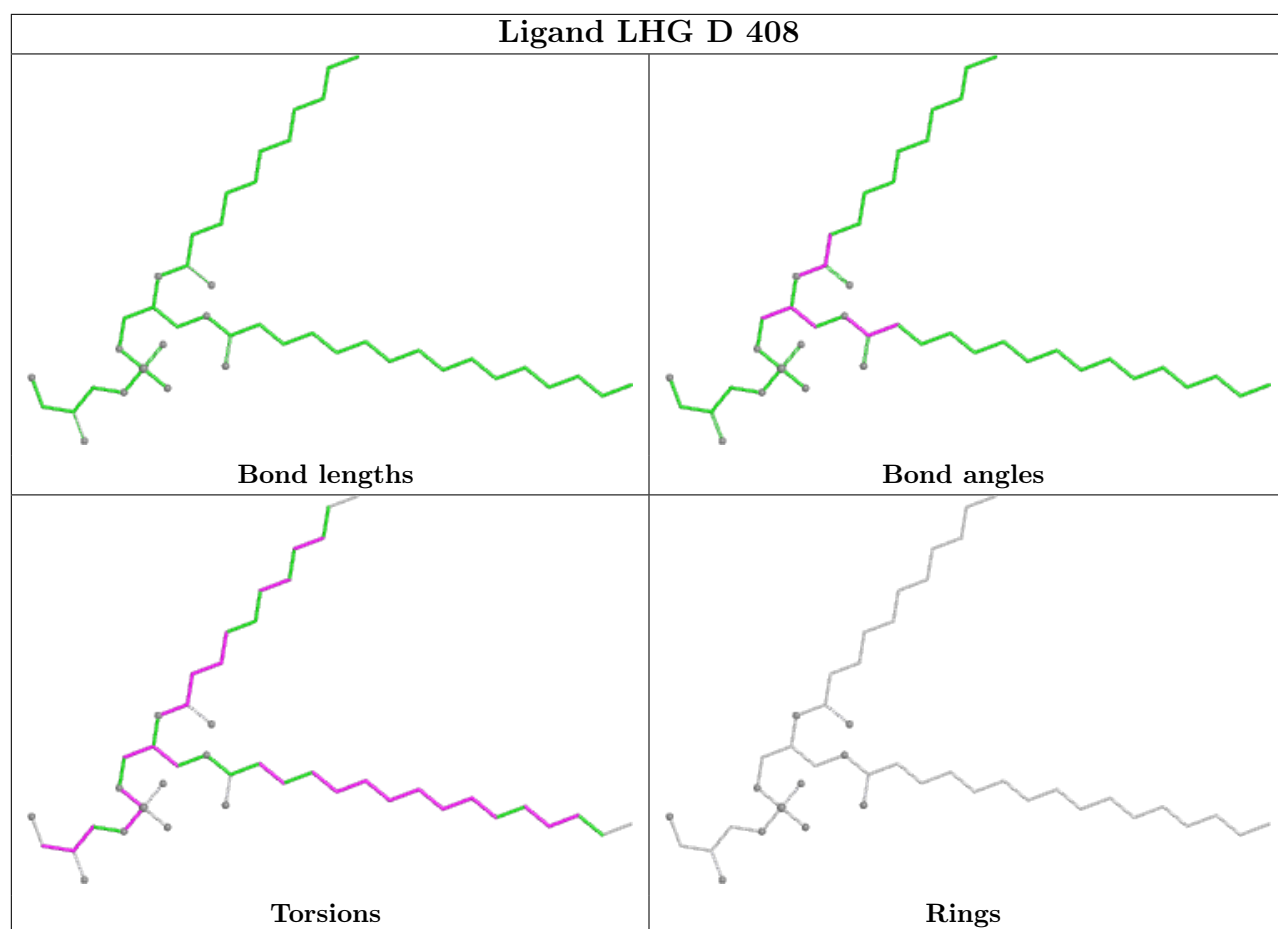


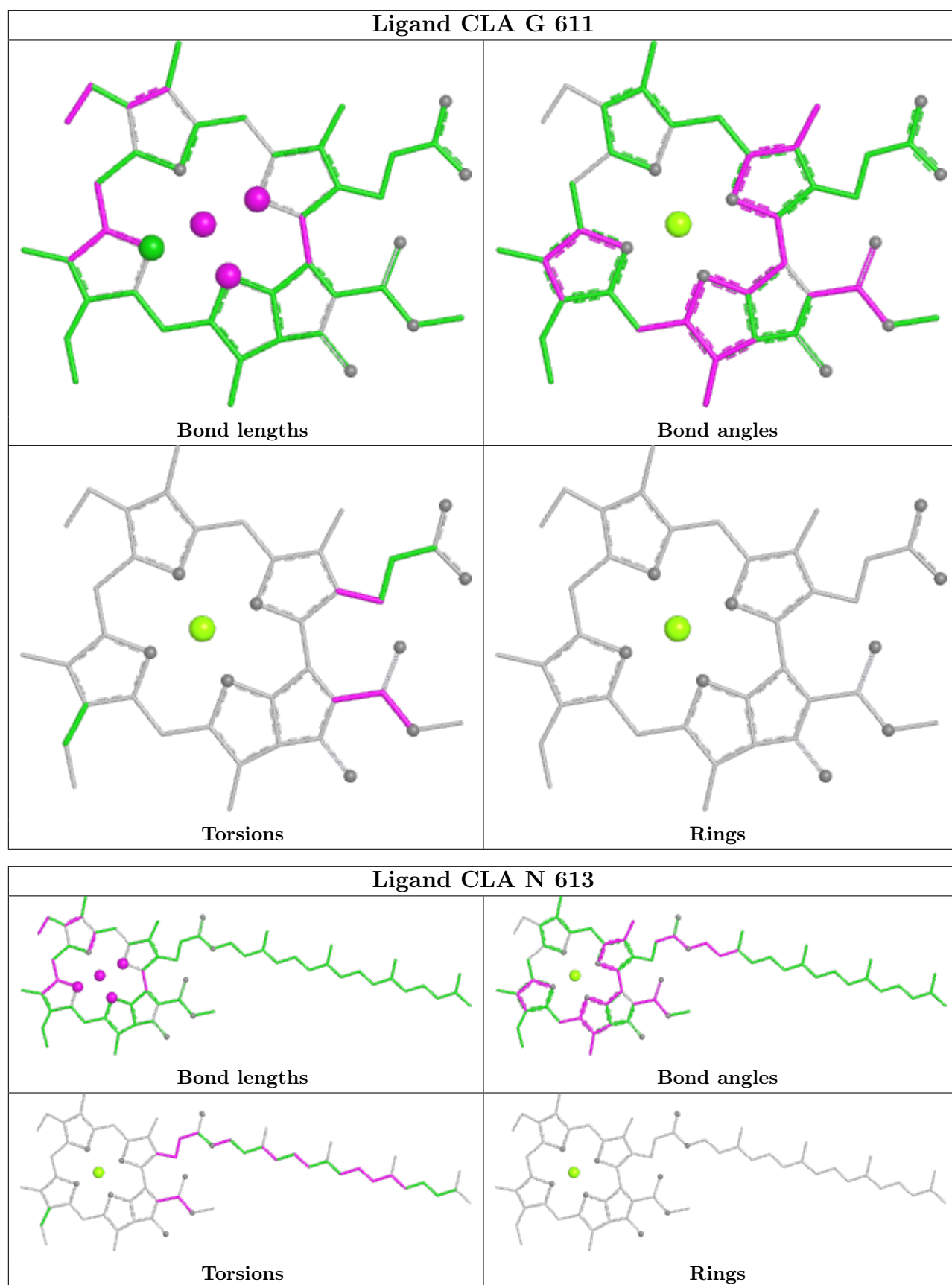


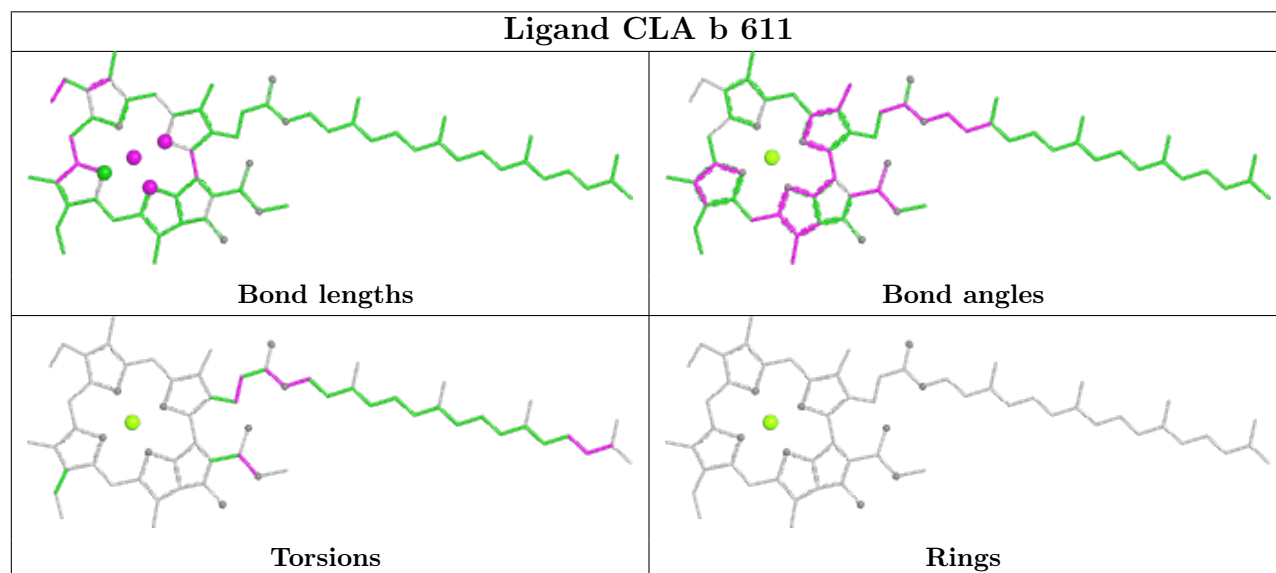
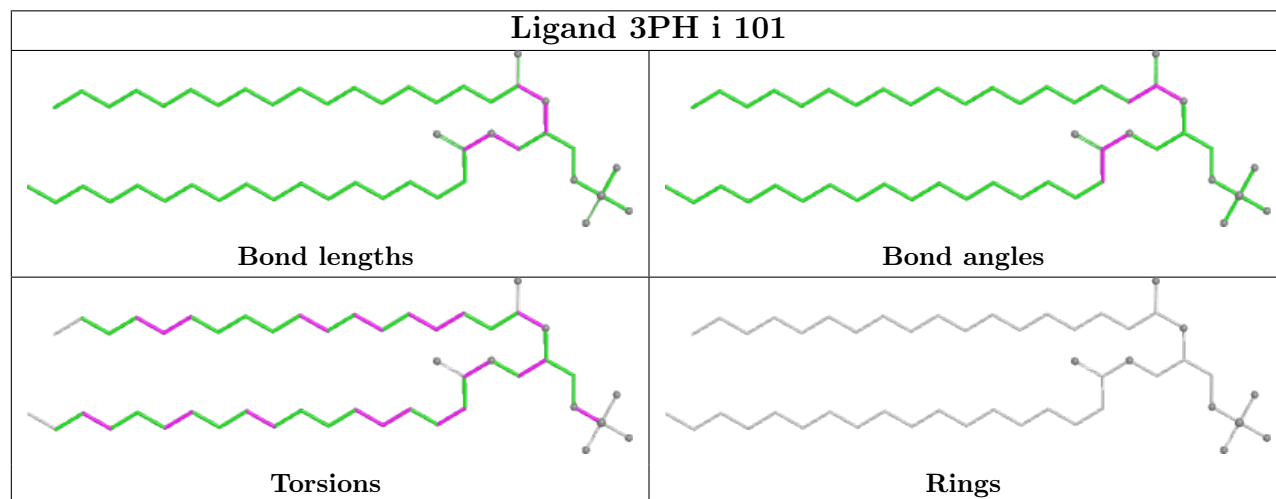
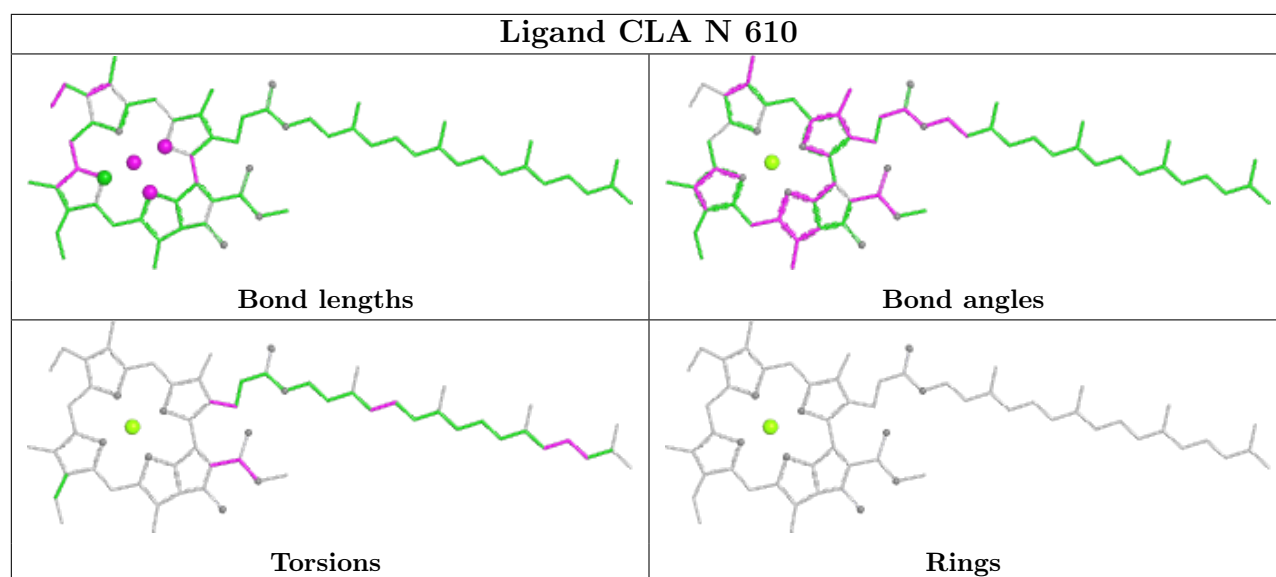


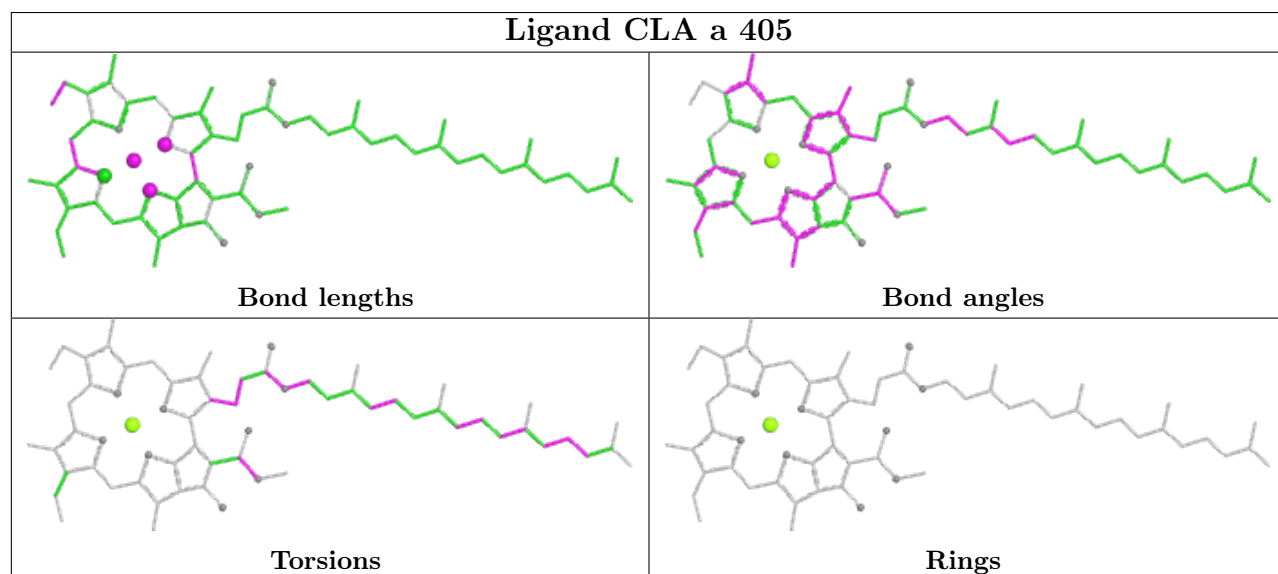
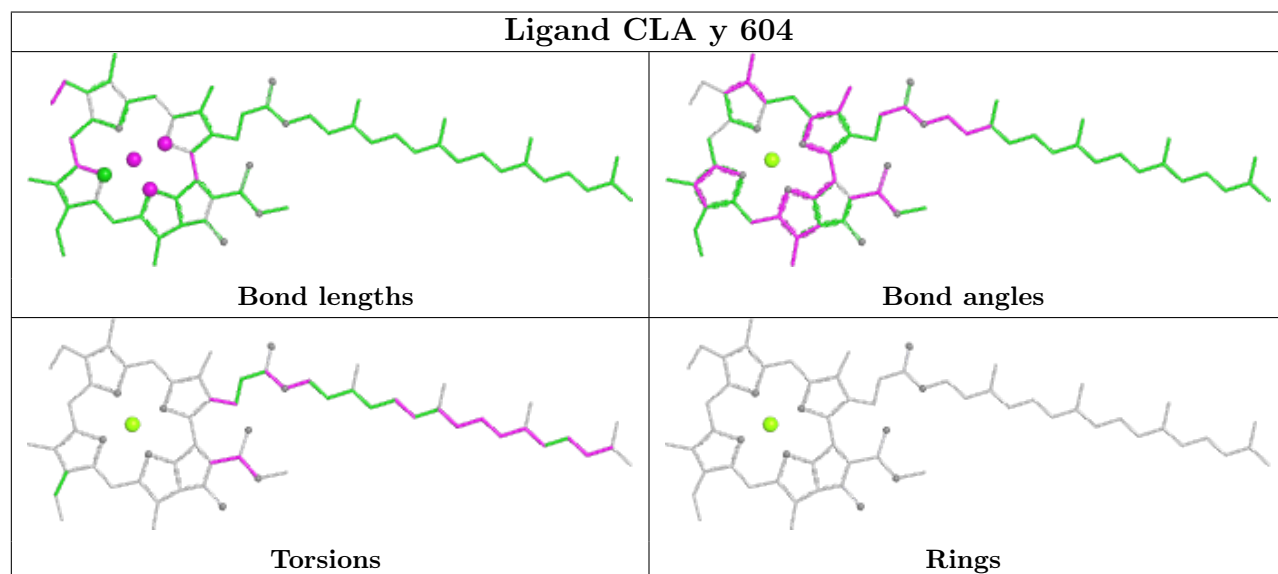
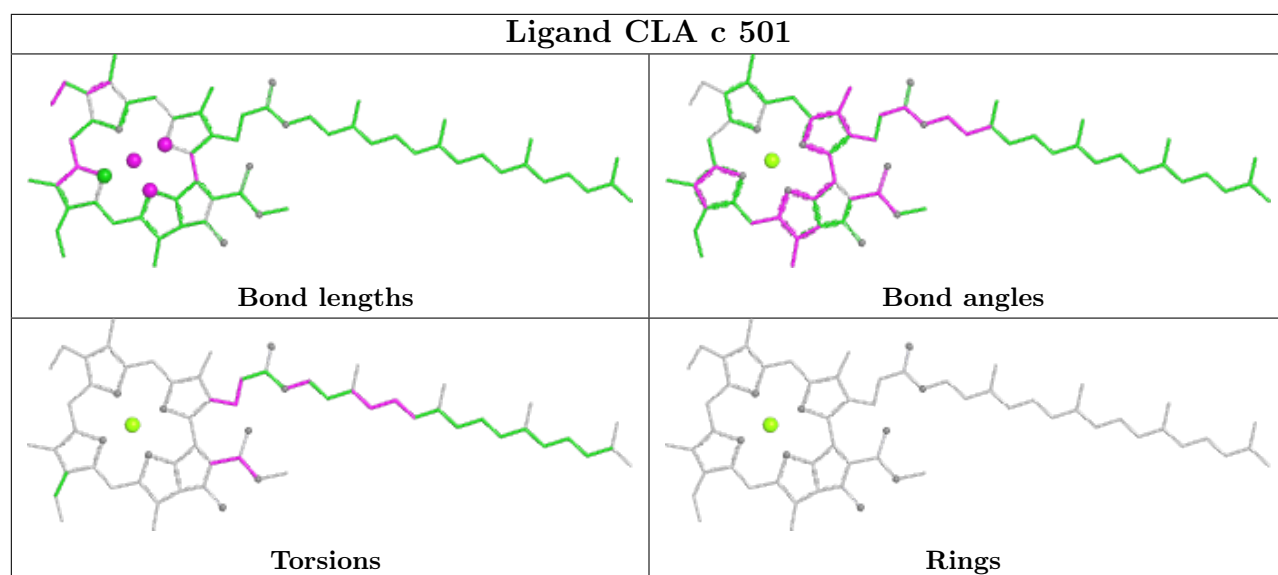


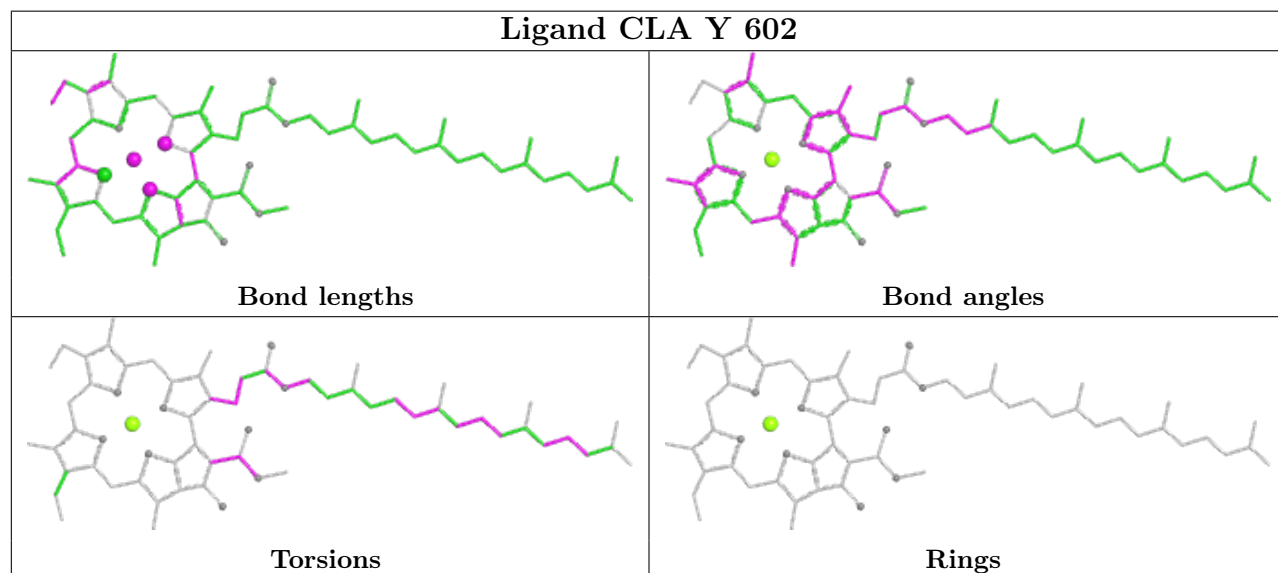
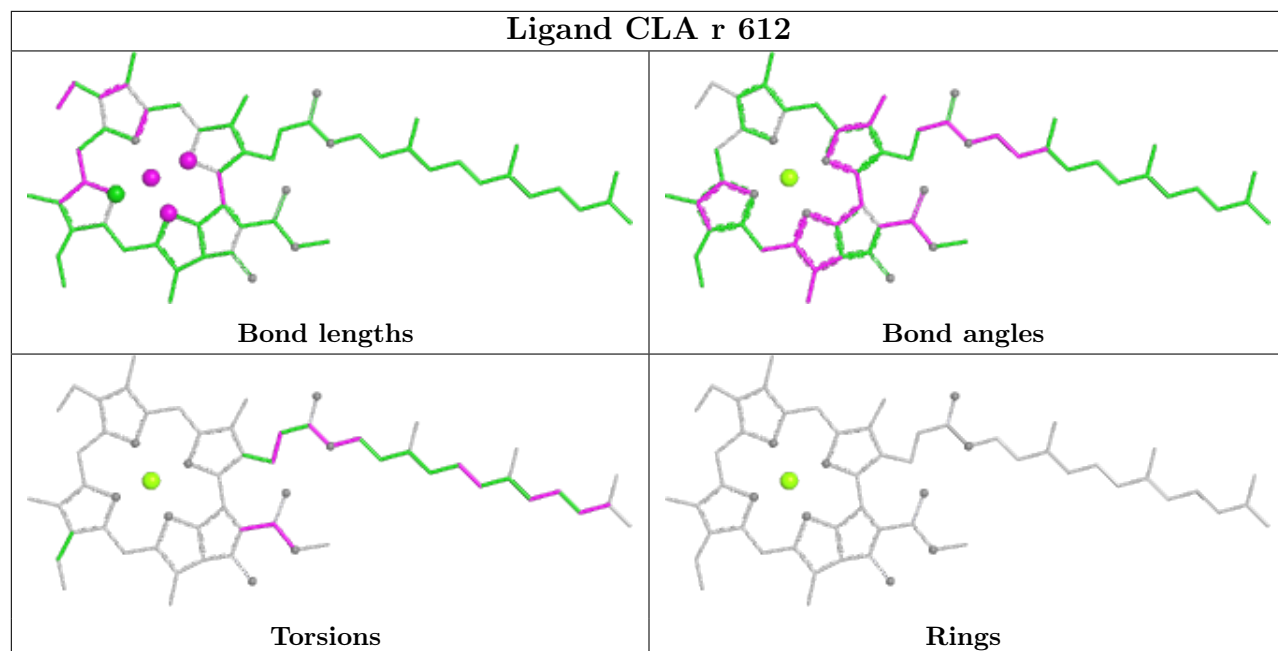


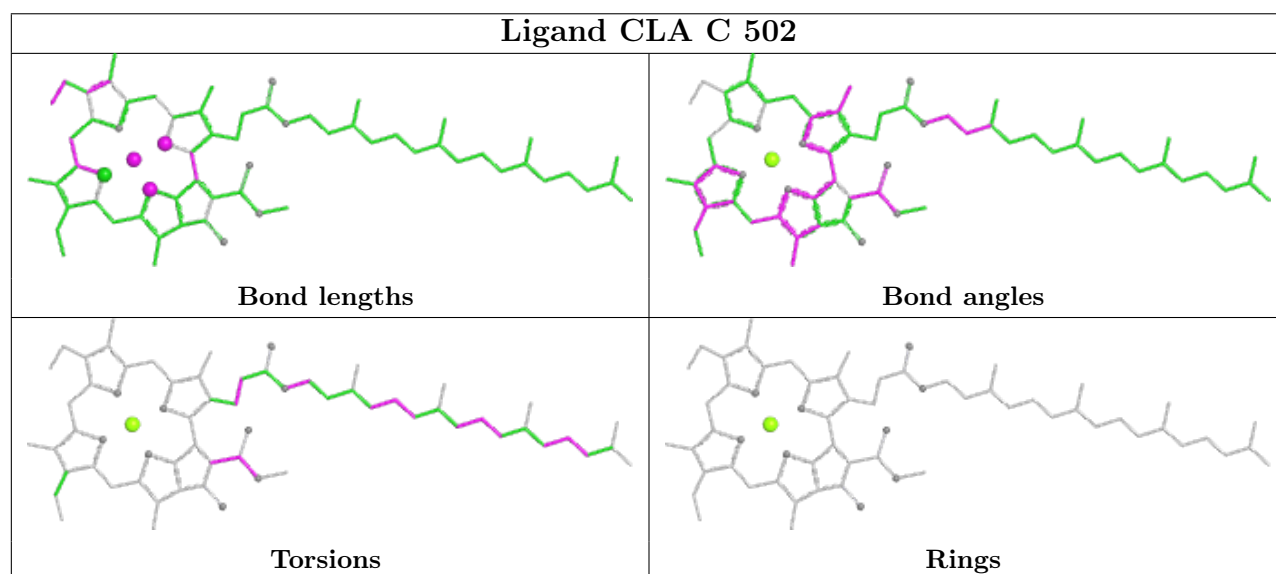
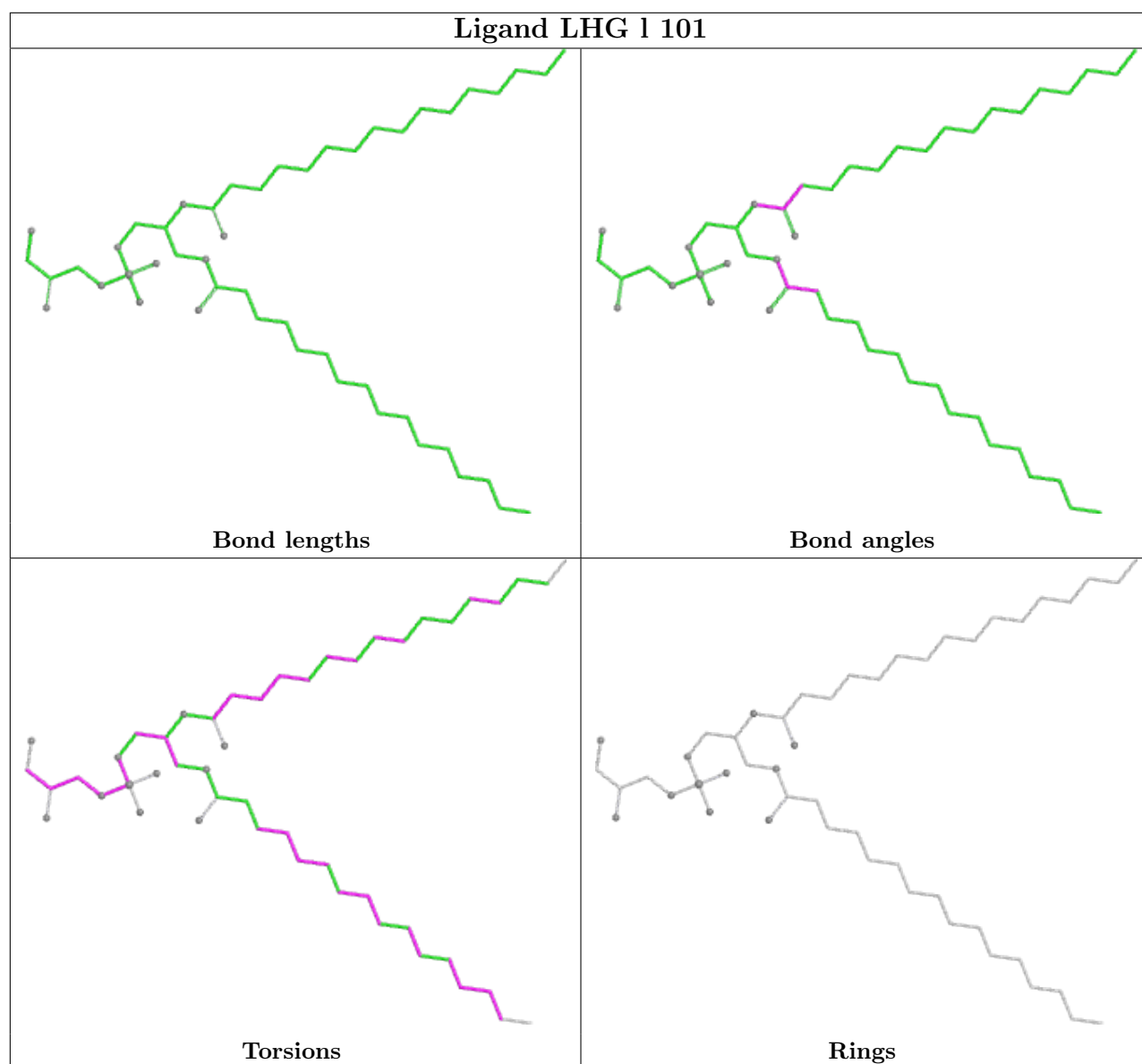


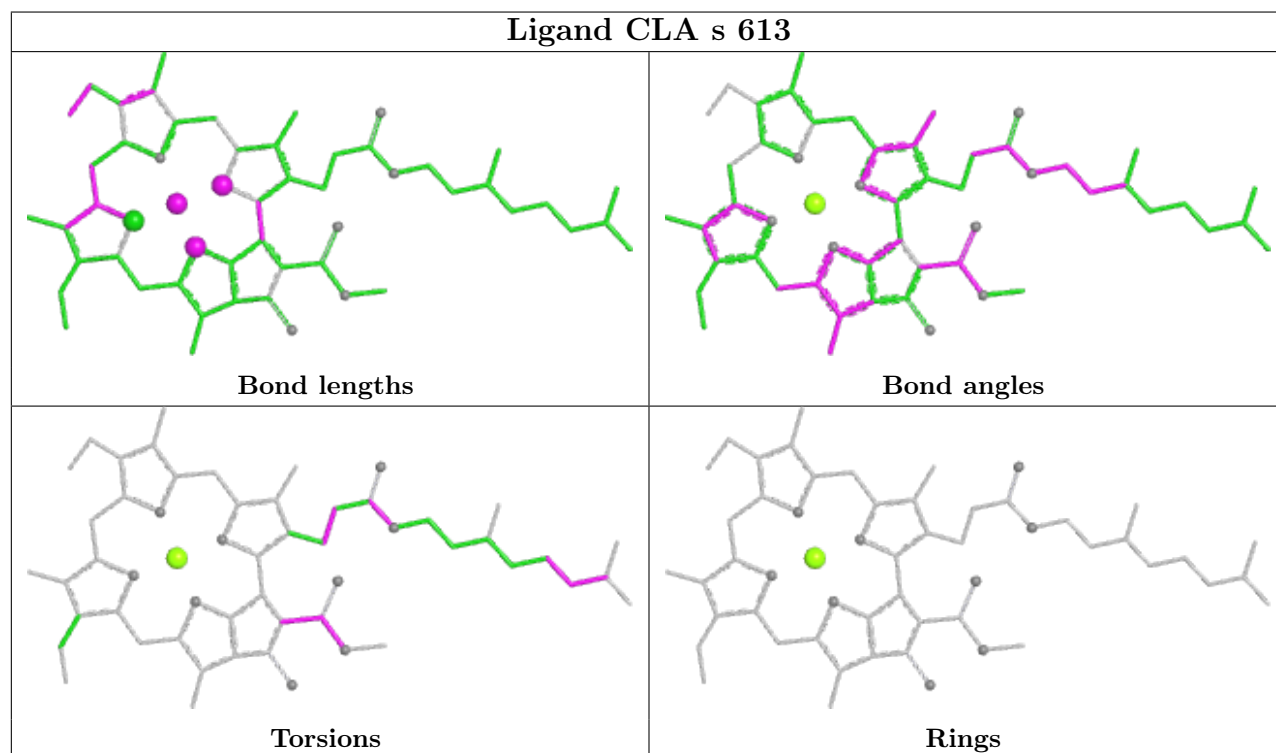
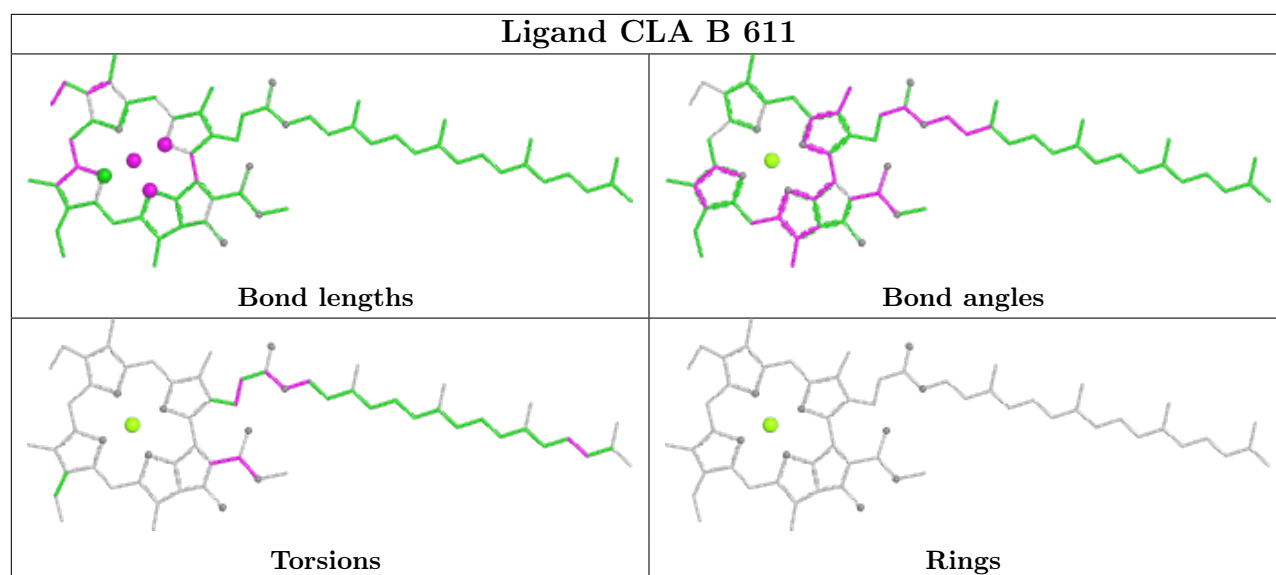


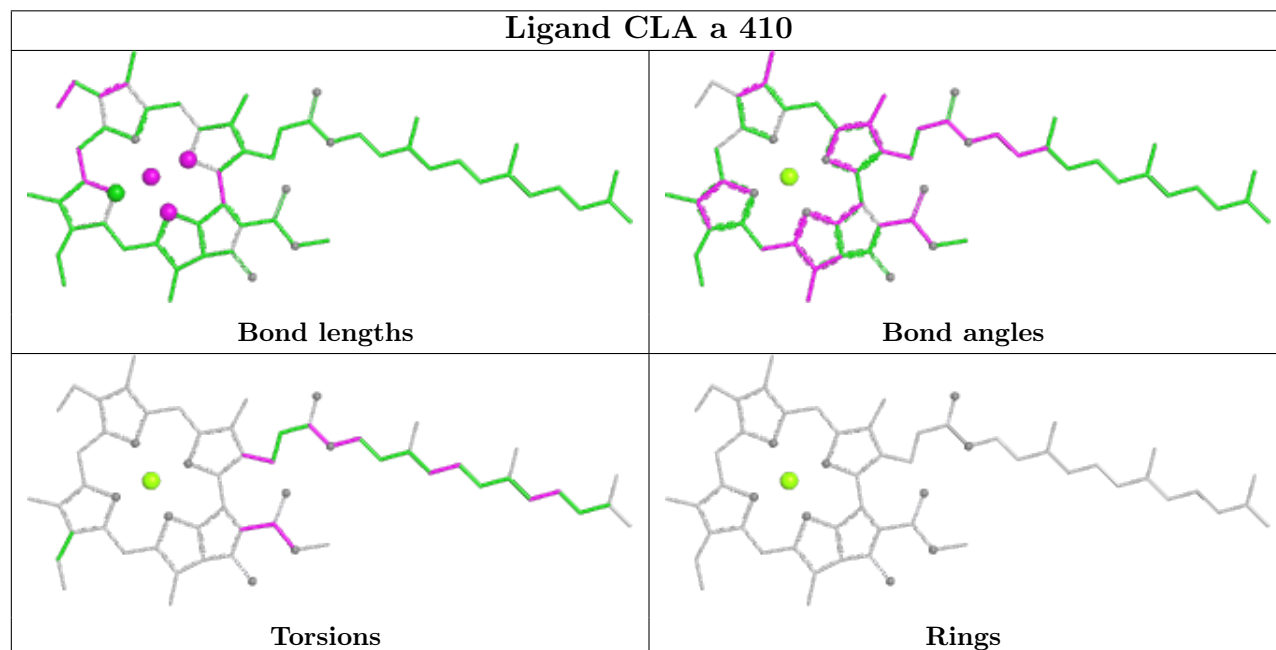
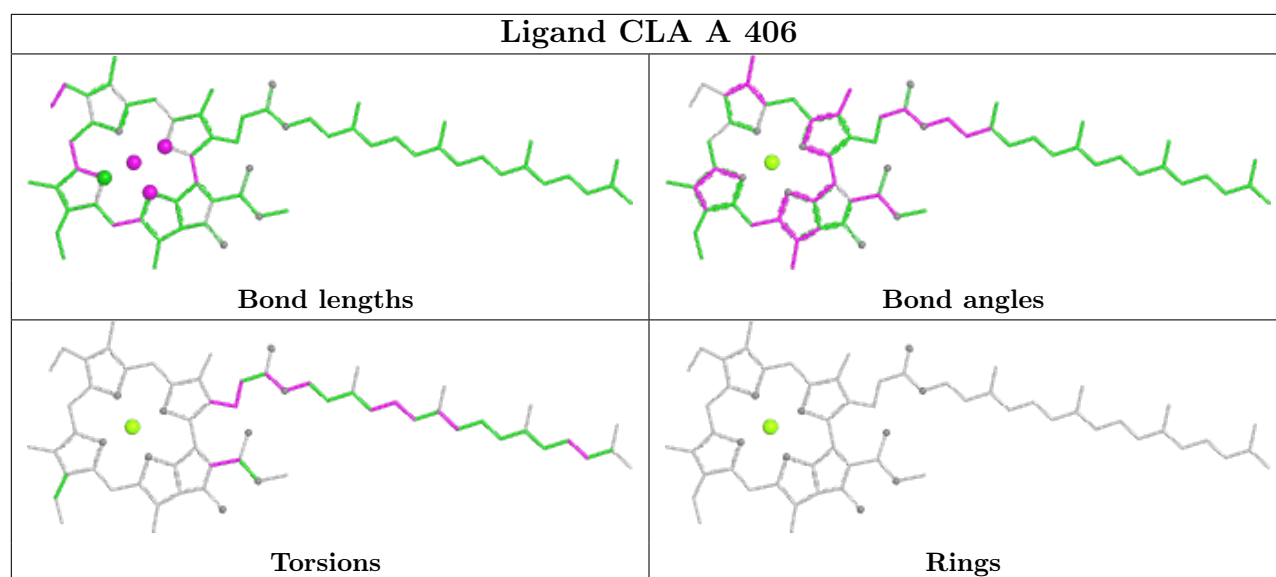


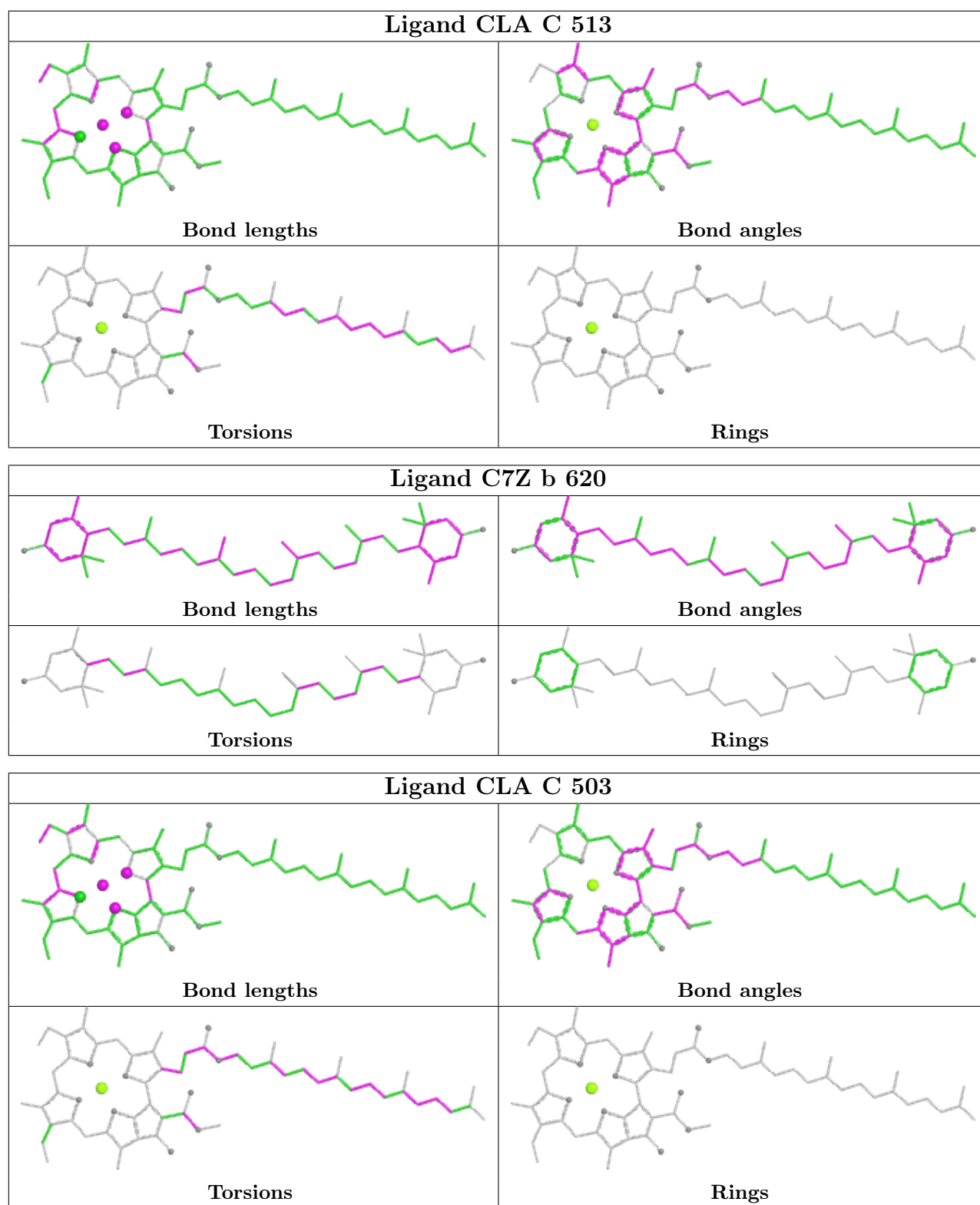


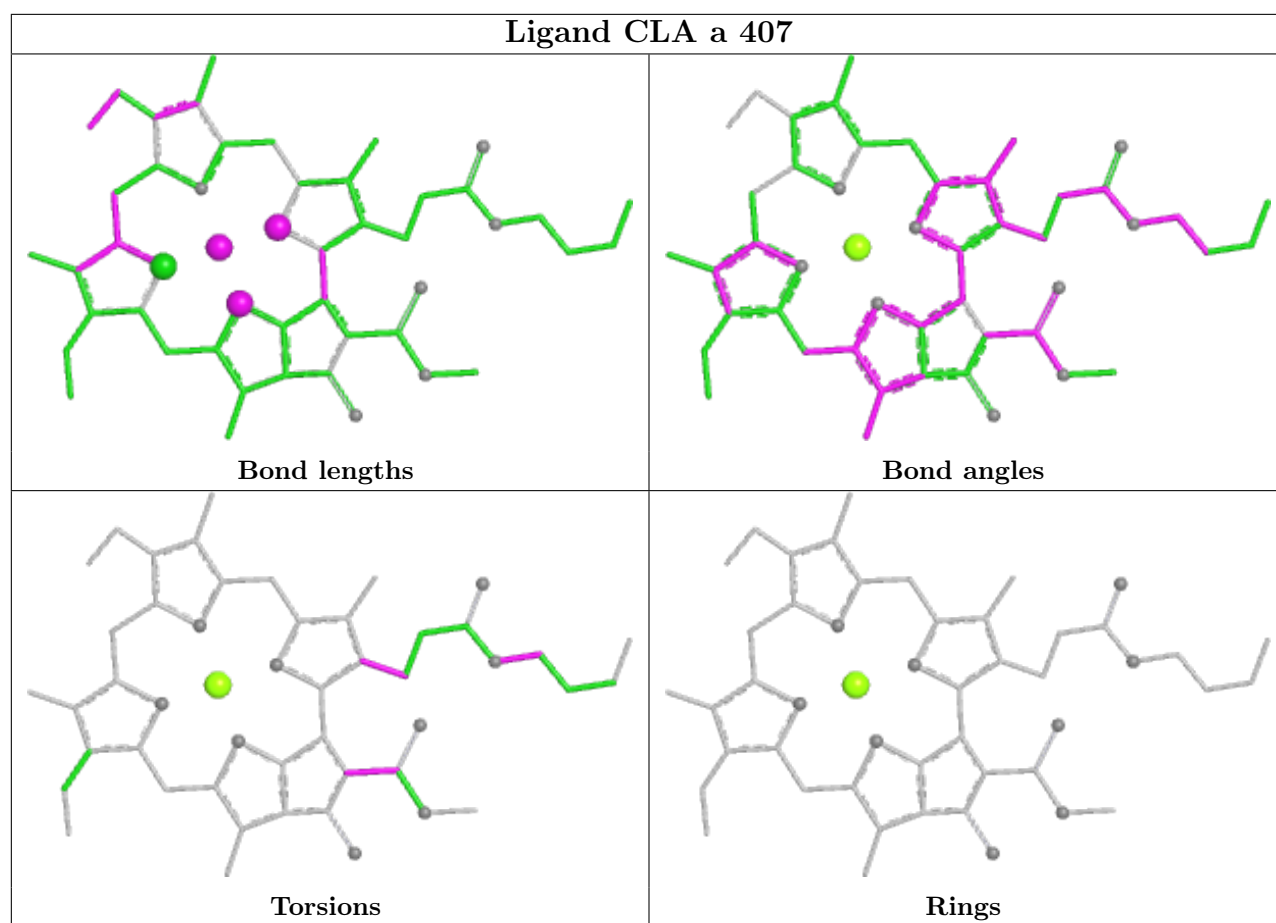


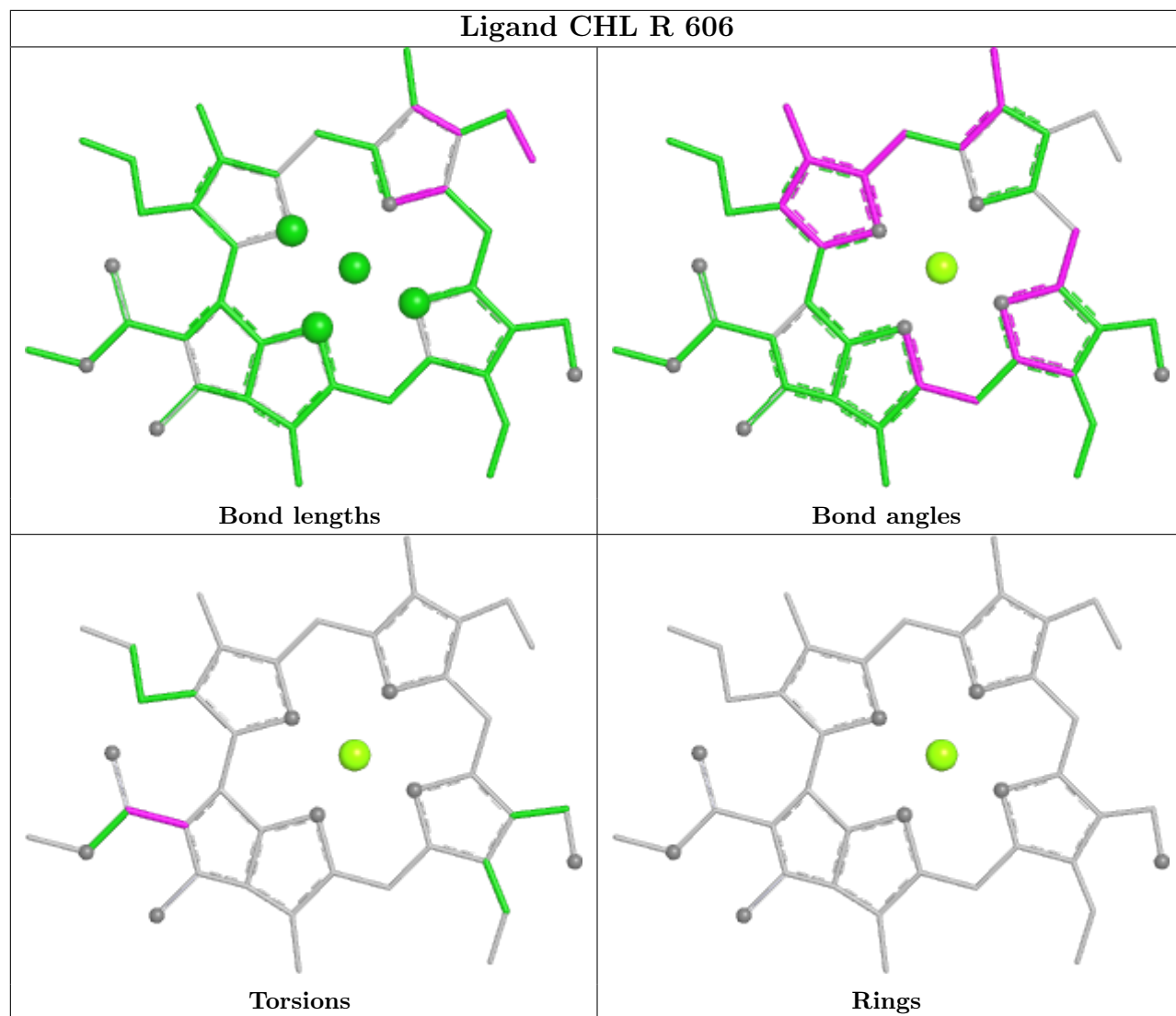


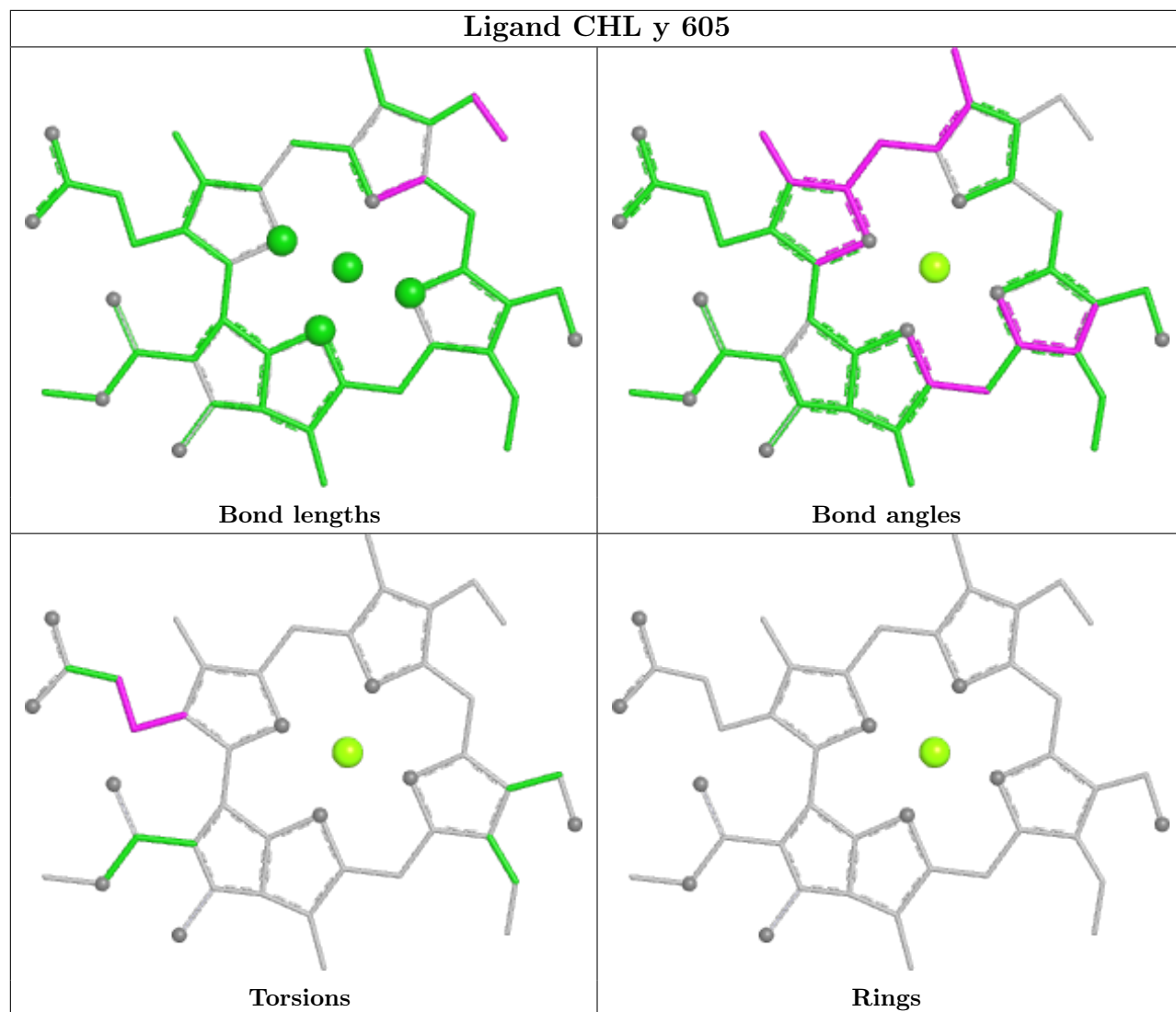


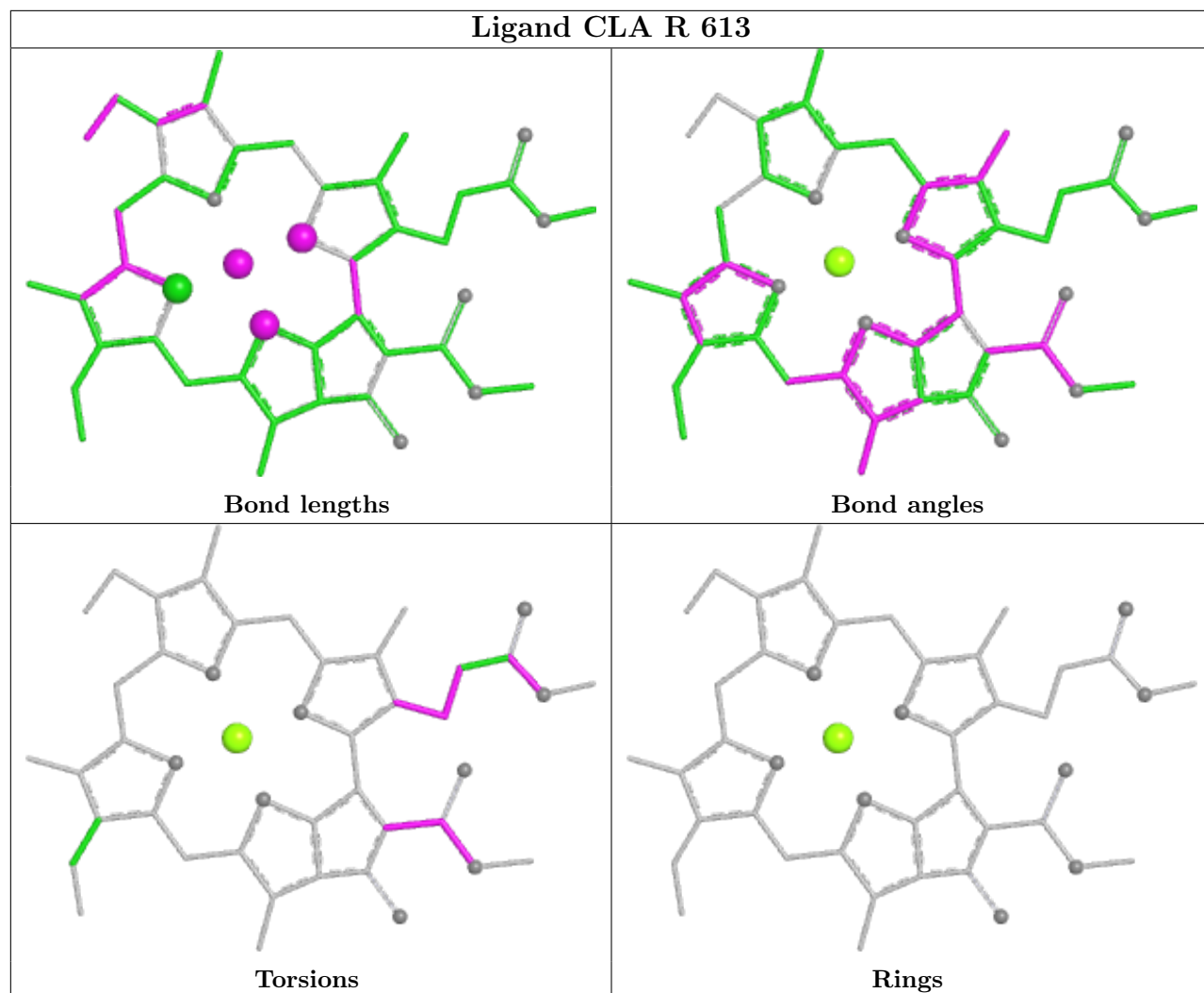


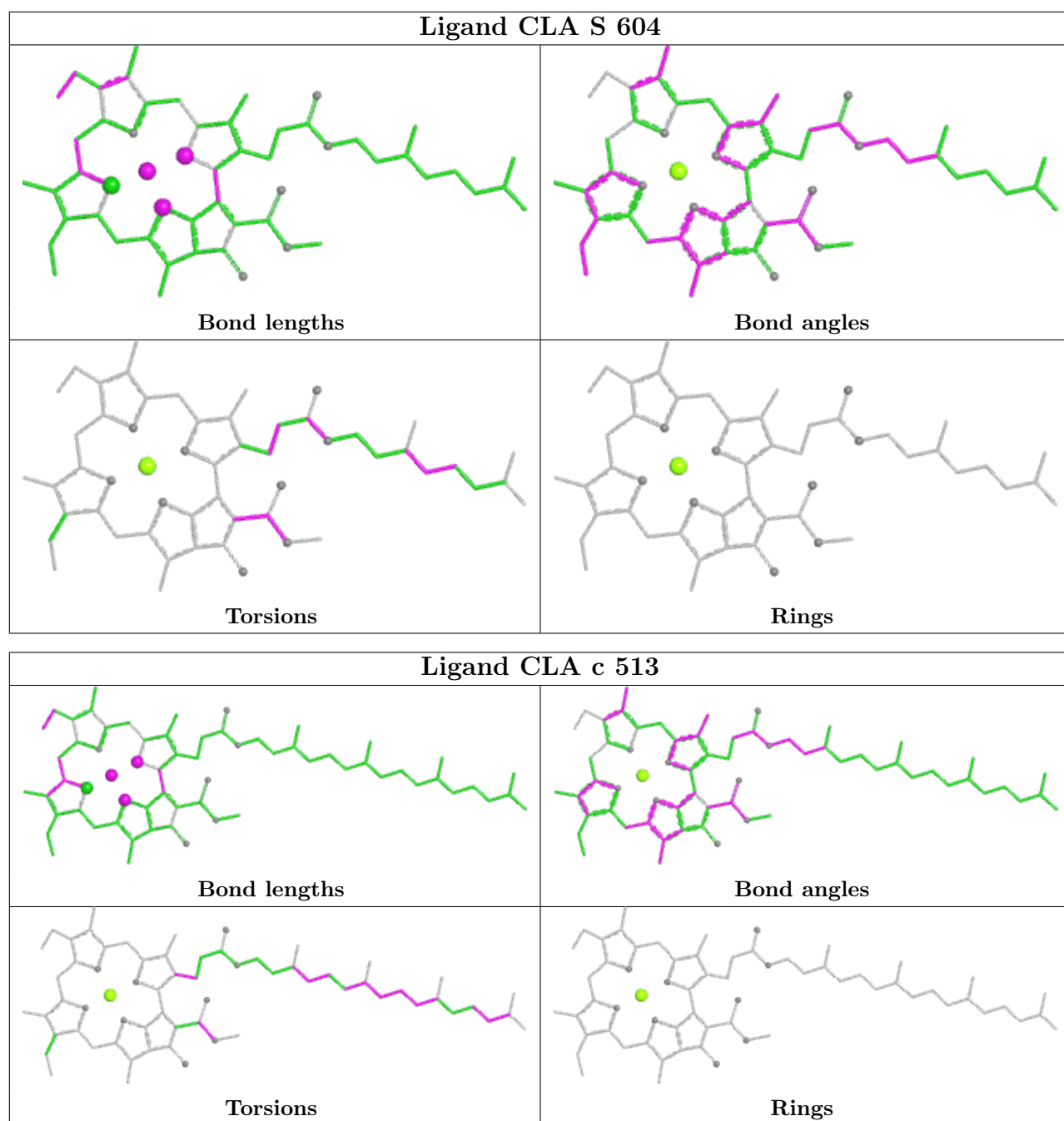


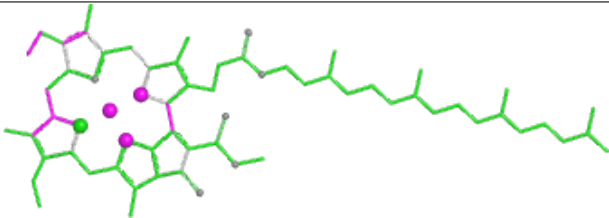
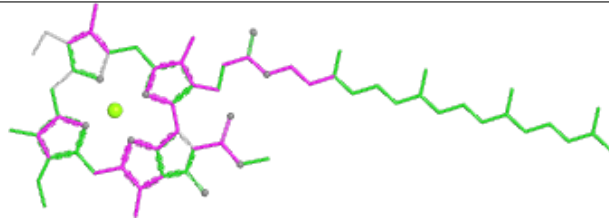
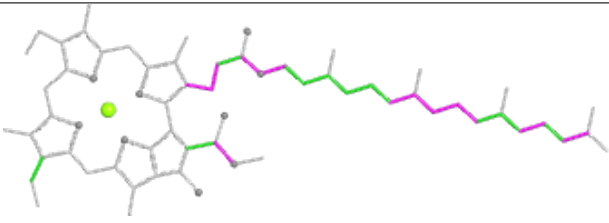
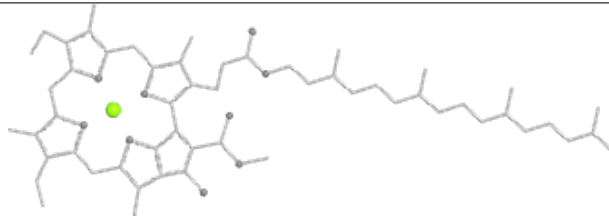


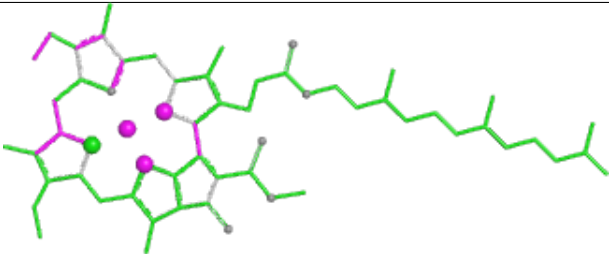
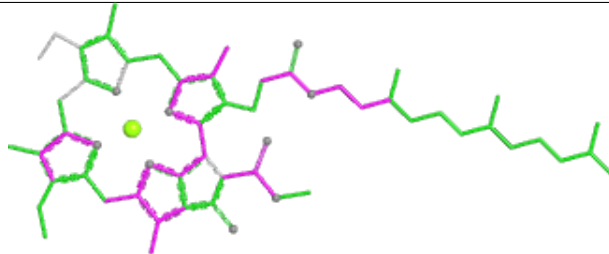
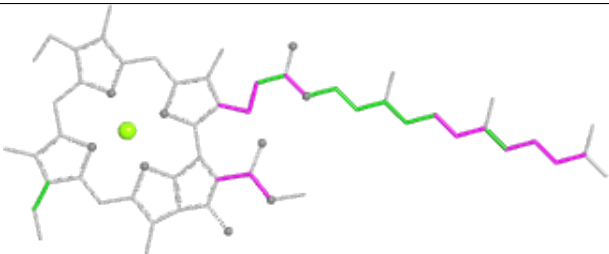
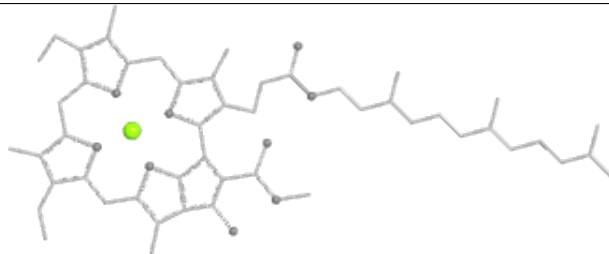


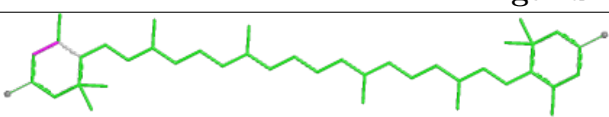
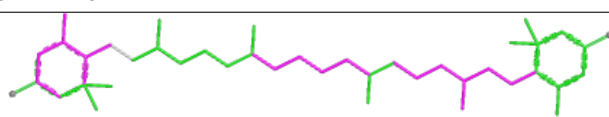
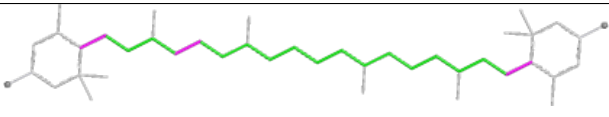
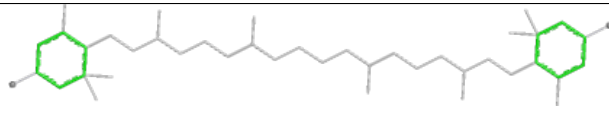


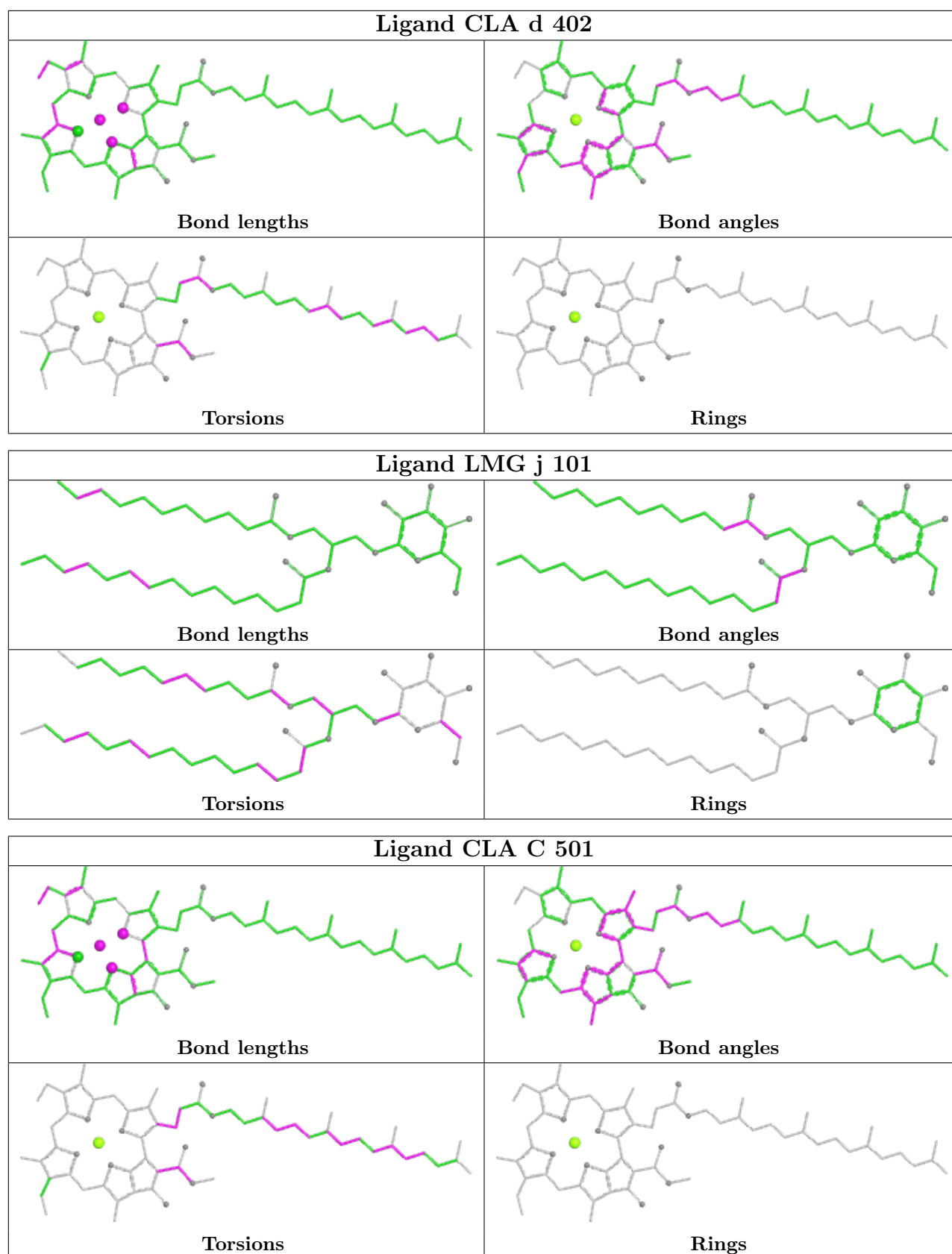




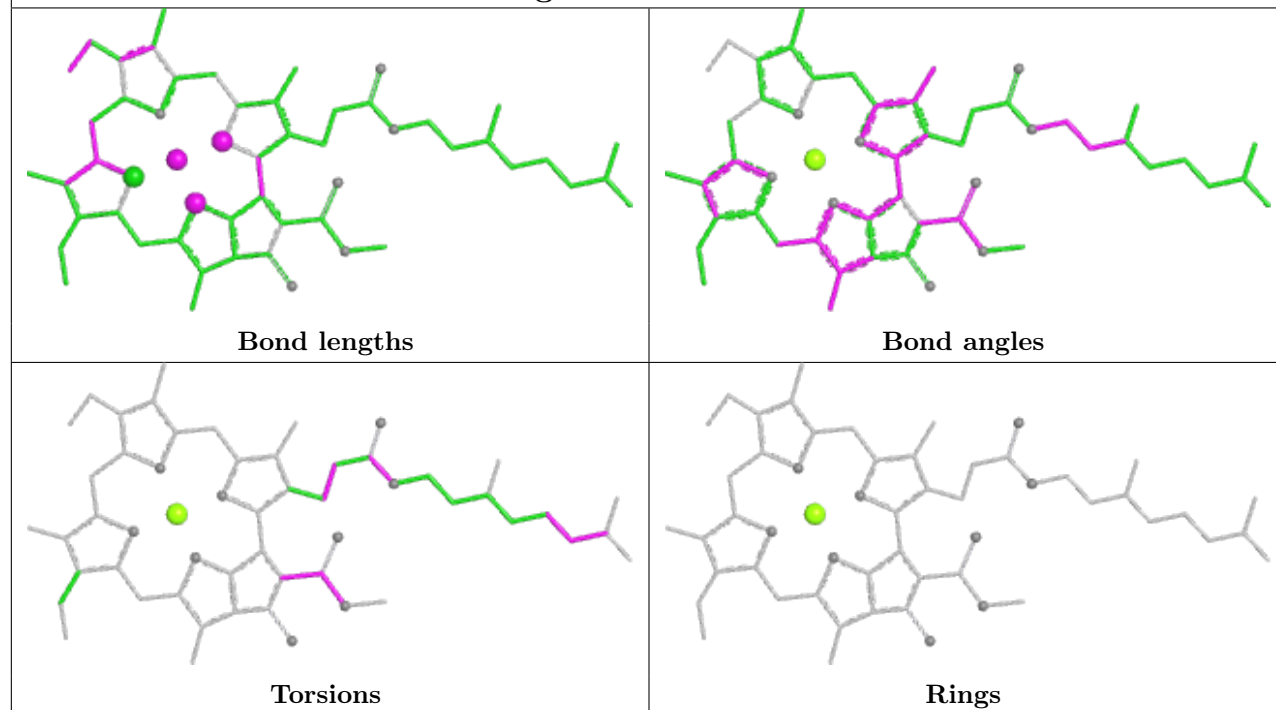
Ligand CLA y 602	
	
Bond lengths	Bond angles
	
Torsions	Rings

Ligand CLA R 608	
	
Bond lengths	Bond angles
	
Torsions	Rings

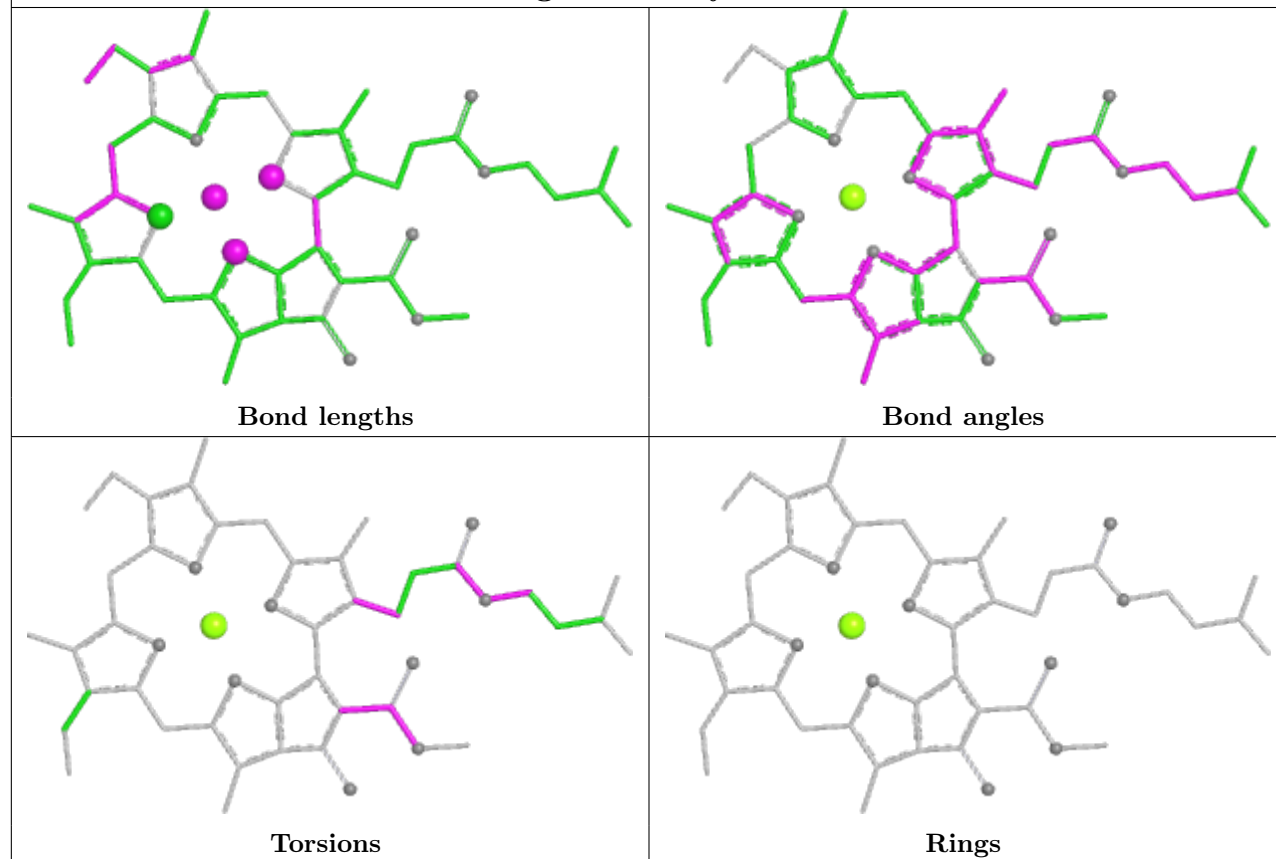
Ligand LUT n 621	
	
Bond lengths	Bond angles
	
Torsions	Rings

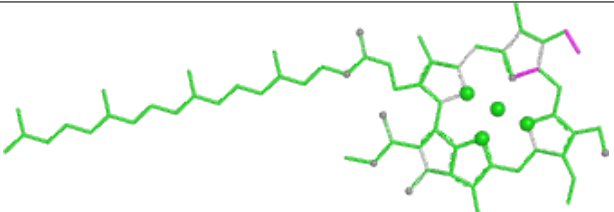
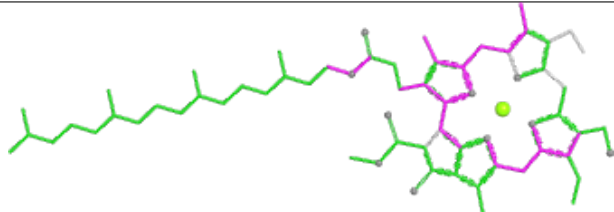
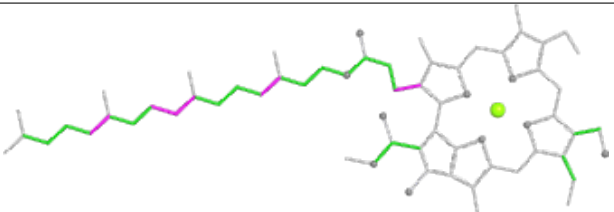
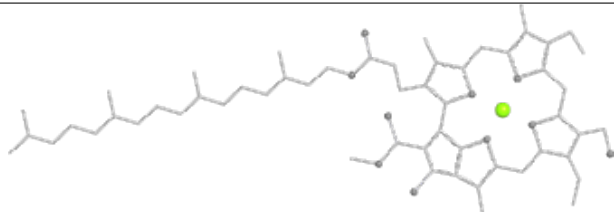


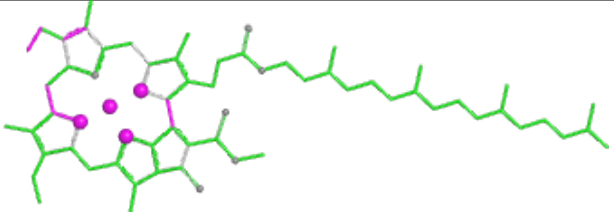
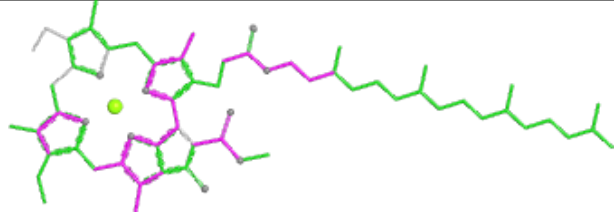
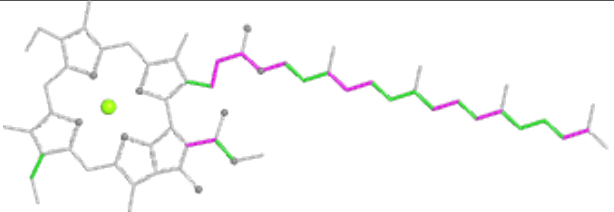
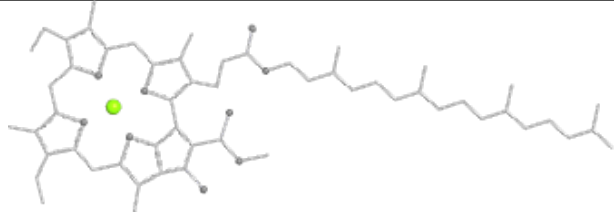
Ligand CLA S 613

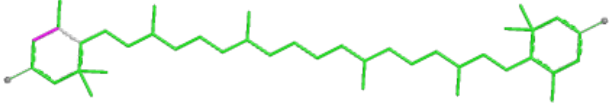
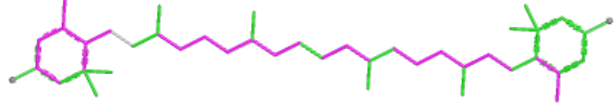
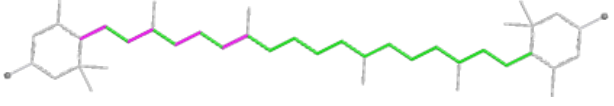
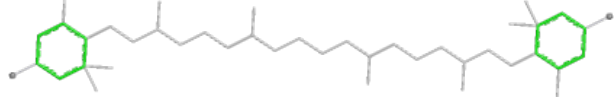


Ligand CLA y 608

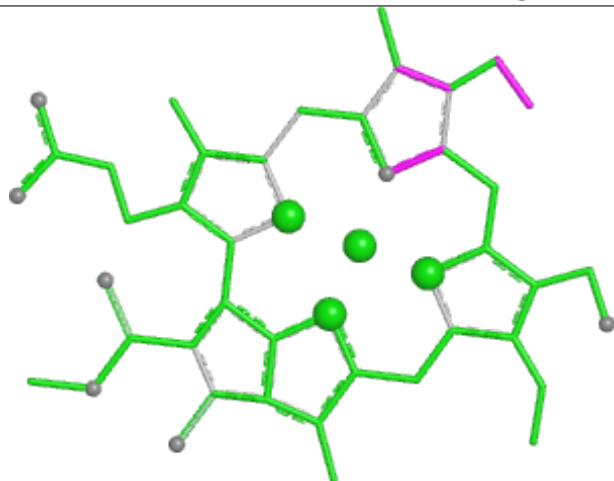


Ligand CHL n 609	
	
Bond lengths	Bond angles
	
Torsions	Rings

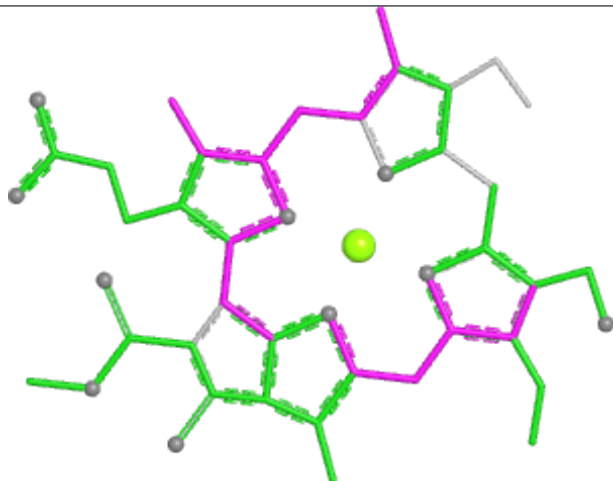
Ligand CLA B 605	
	
Bond lengths	Bond angles
	
Torsions	Rings

Ligand LUT G 620	
	
Bond lengths	Bond angles
	
Torsions	Rings

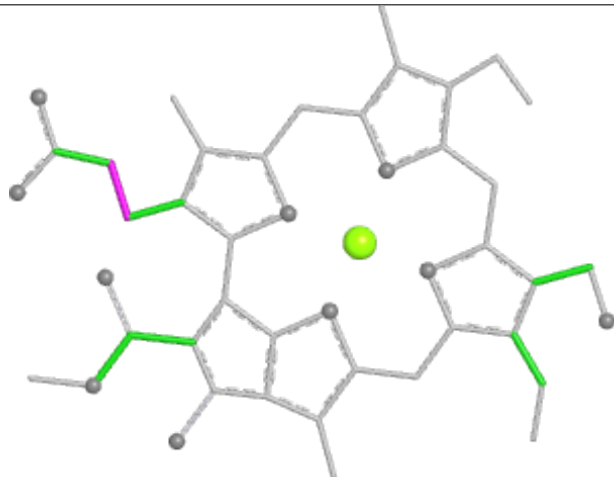
Ligand CHL Y 605



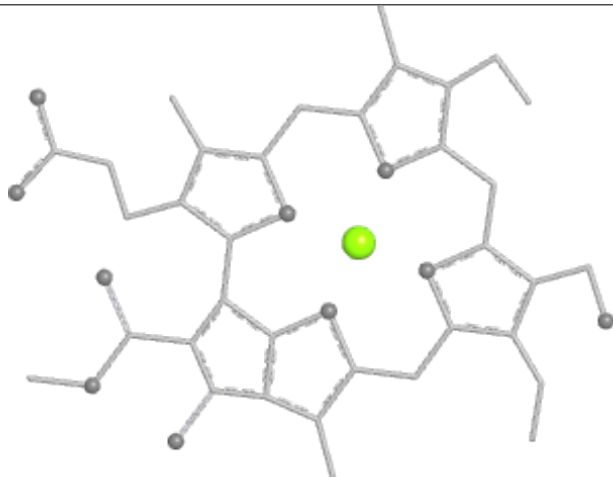
Bond lengths



Bond angles

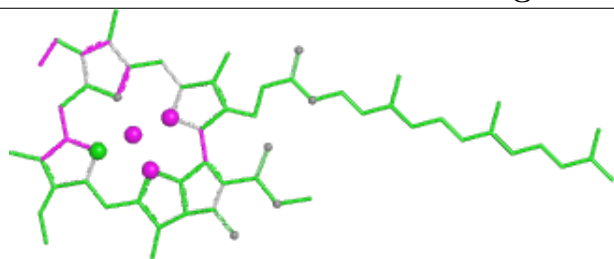


Torsions

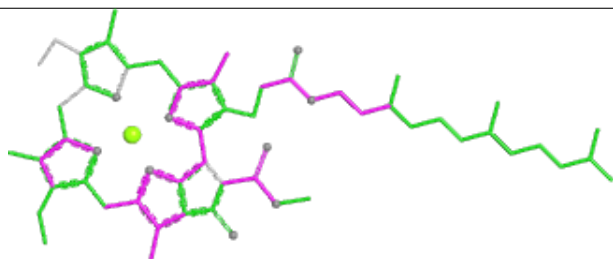


Rings

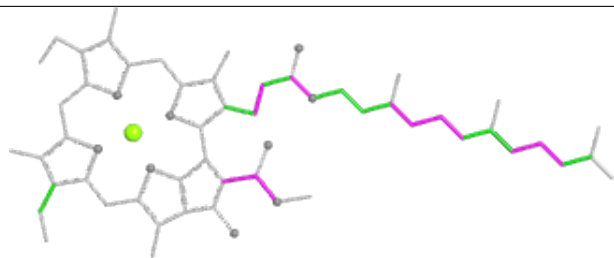
Ligand CLA R 602



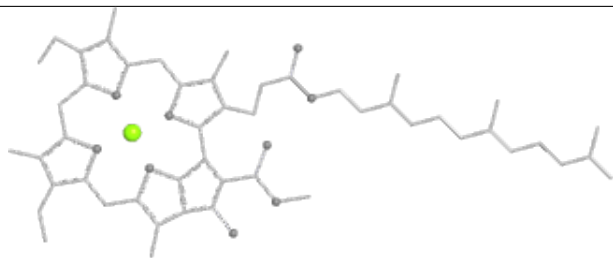
Bond lengths



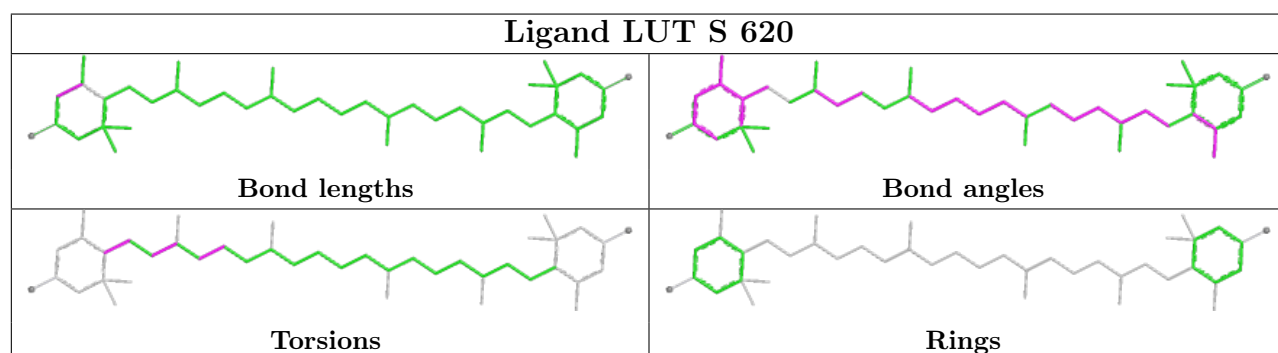
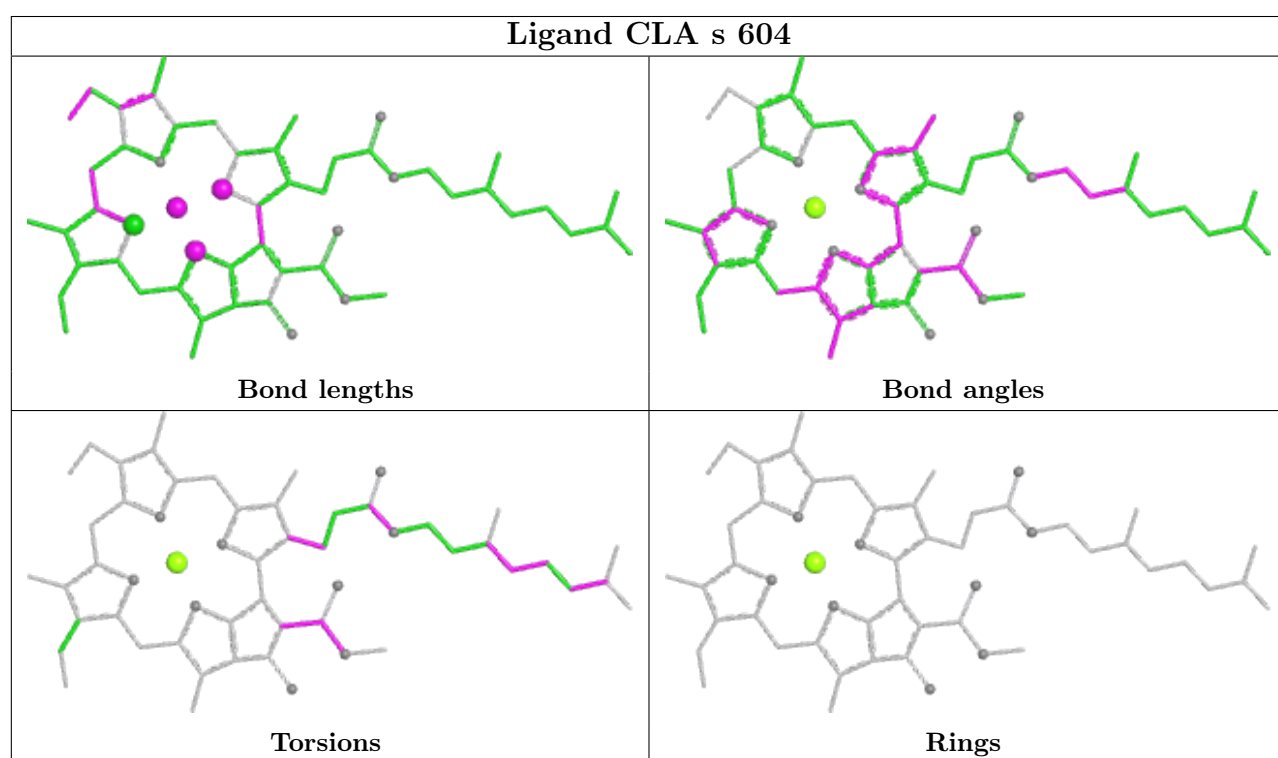
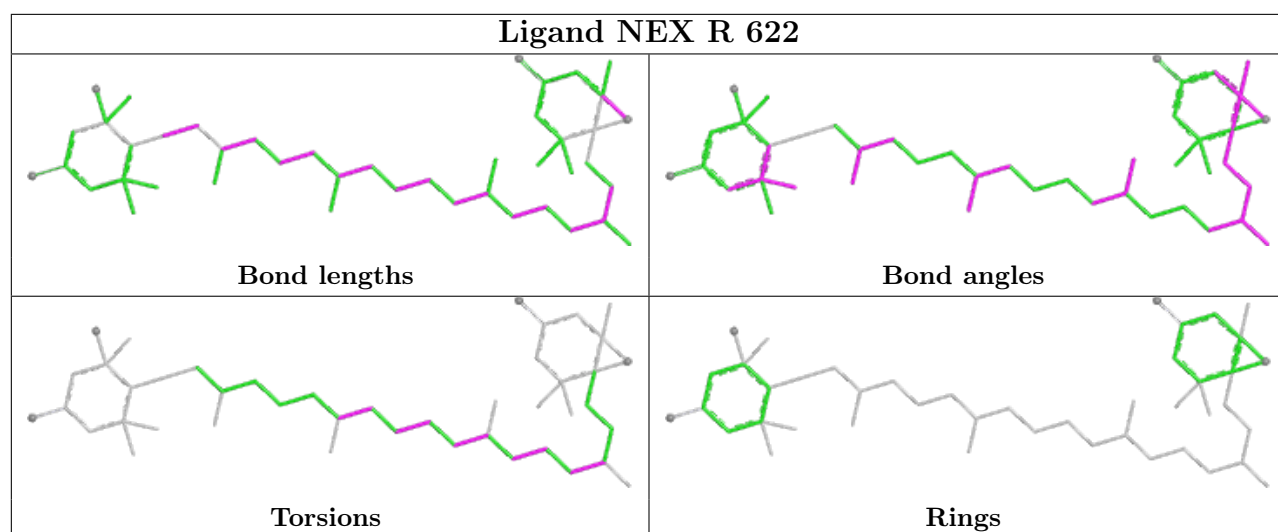
Bond angles

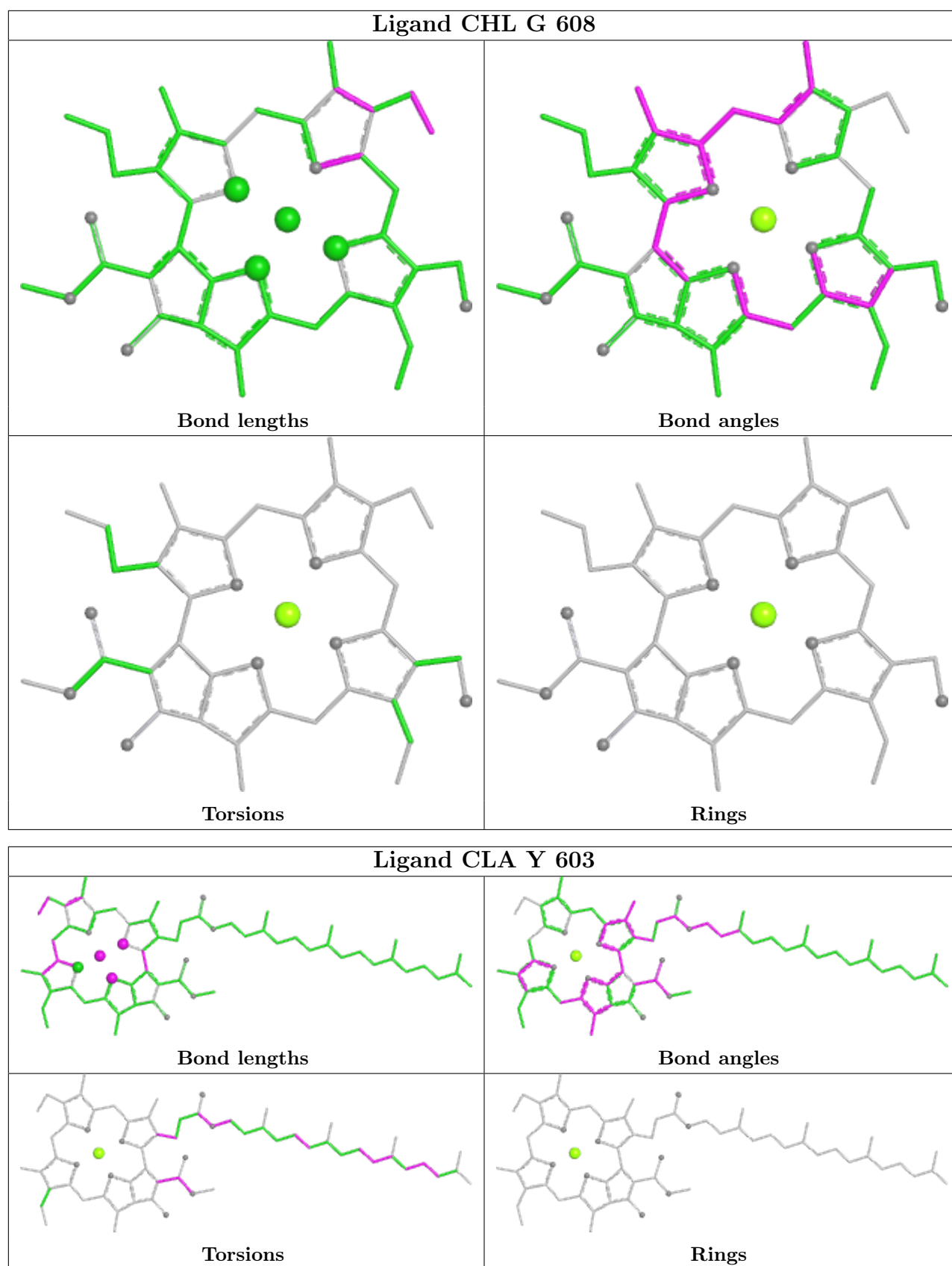


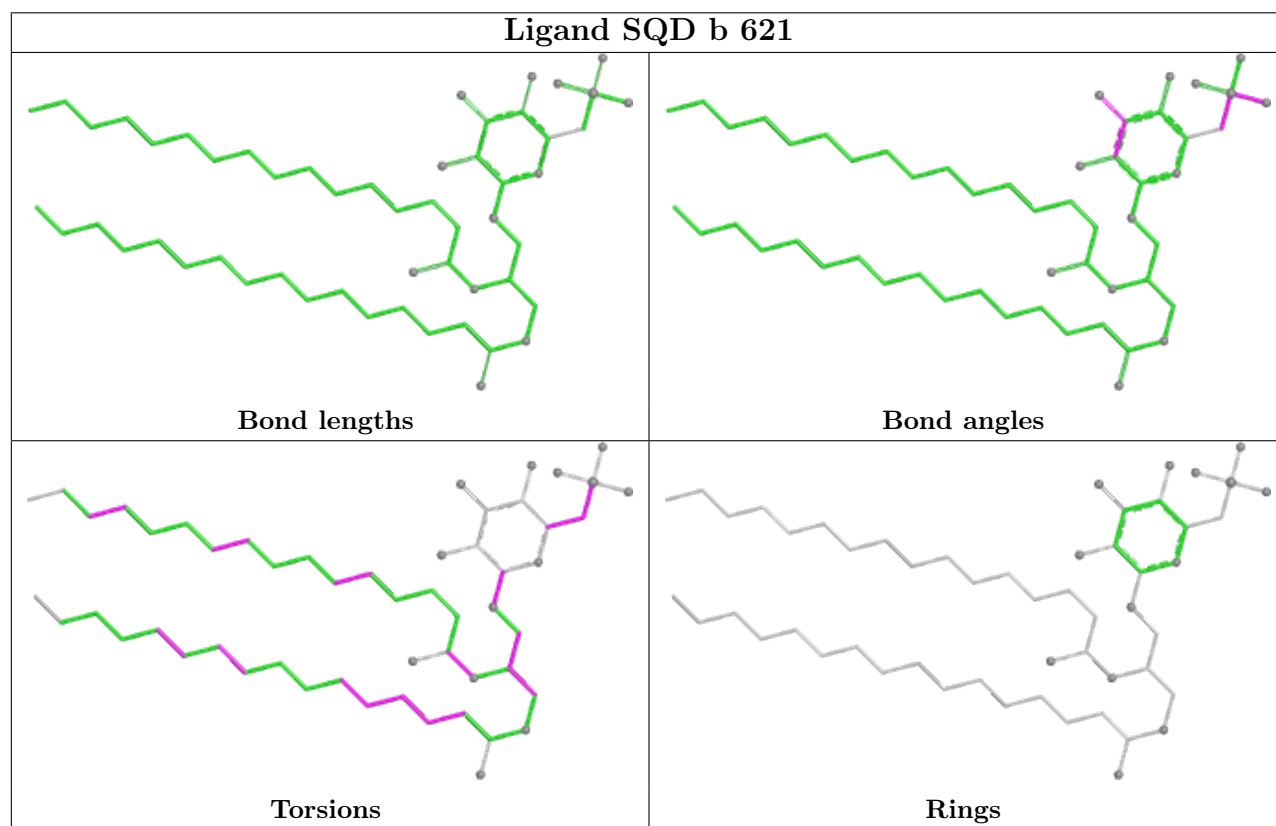
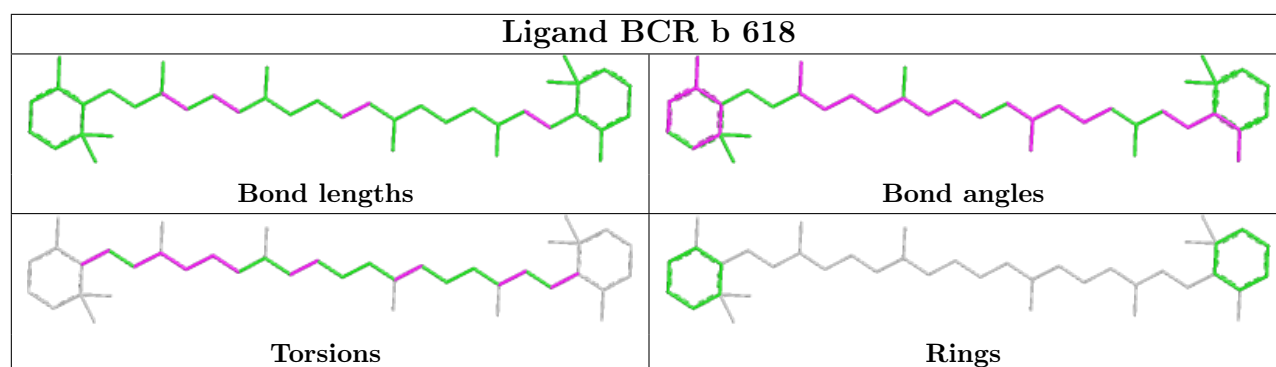
Torsions

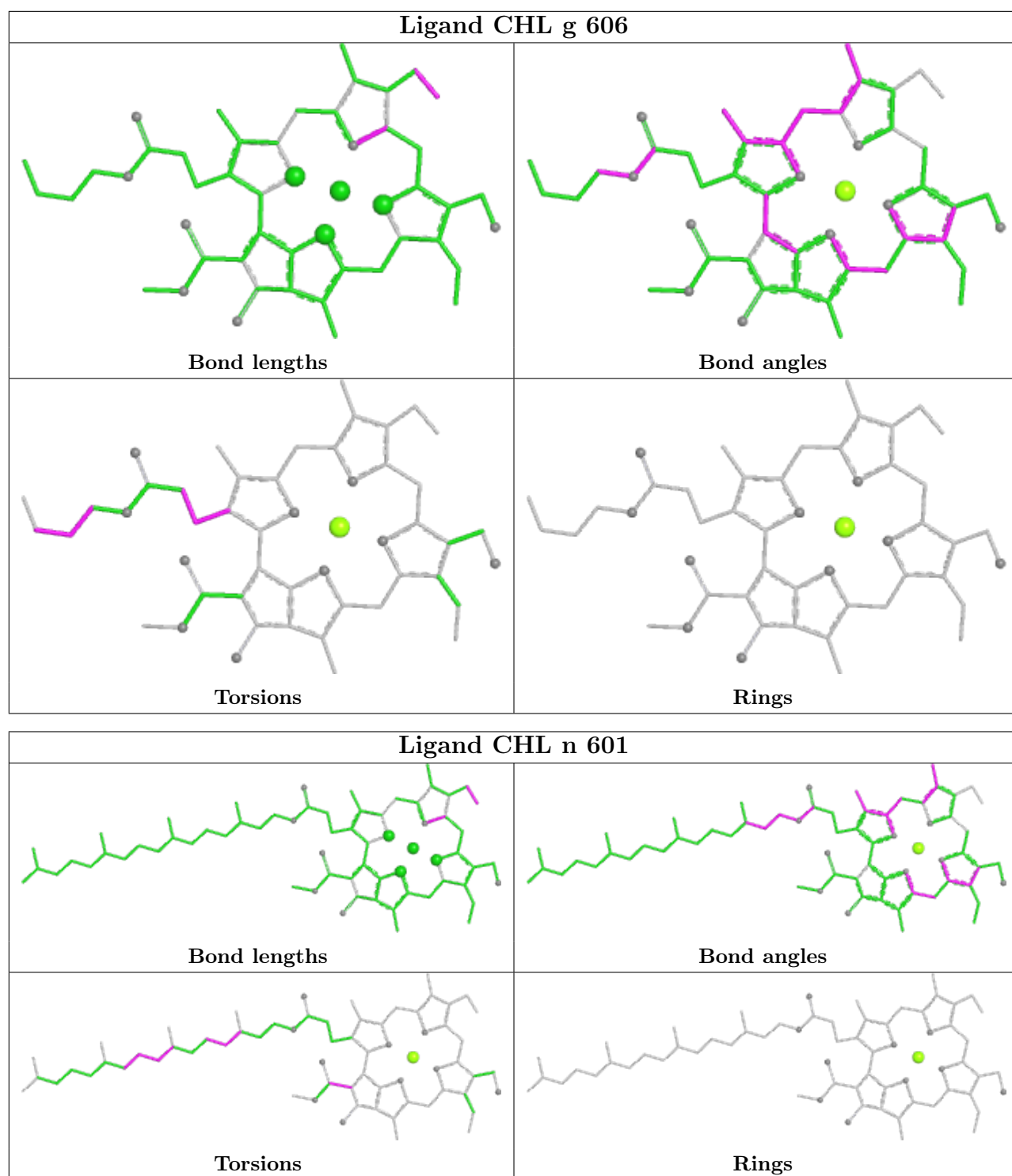


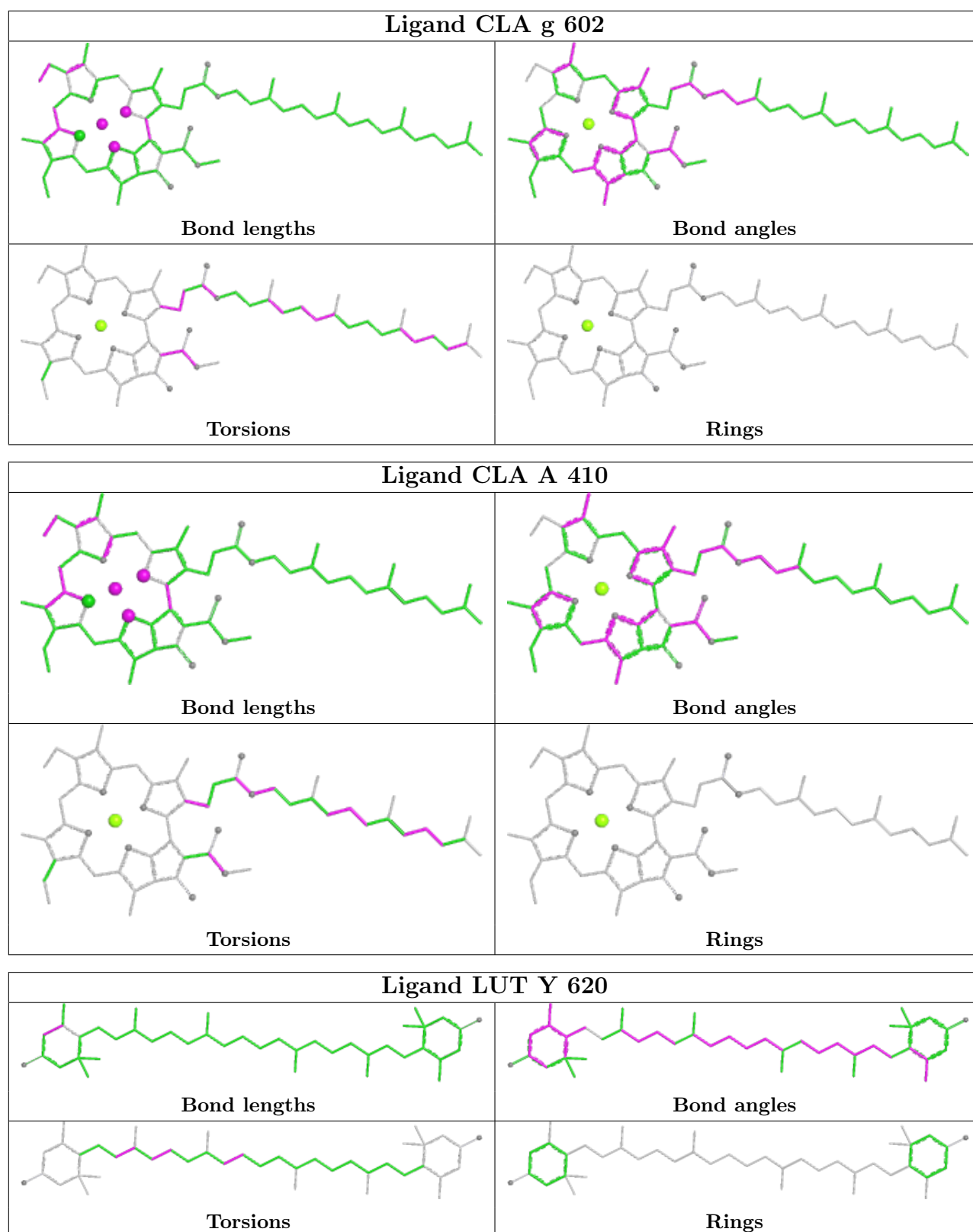
Rings

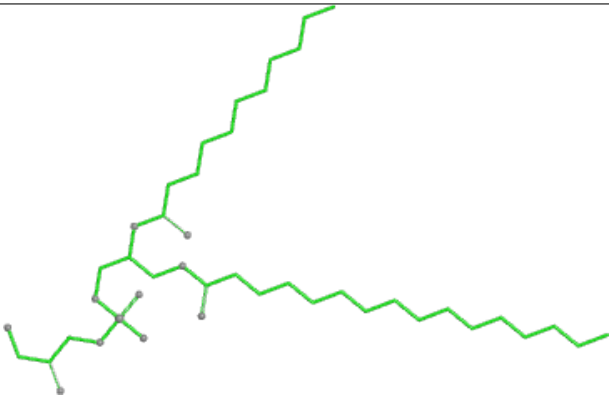
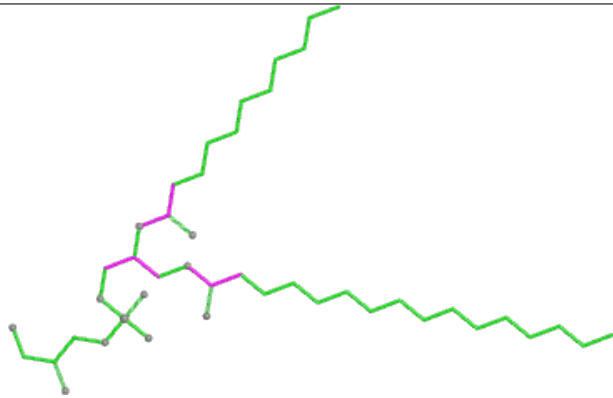
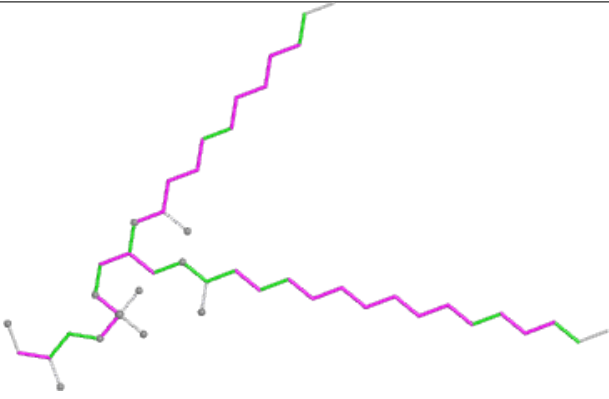
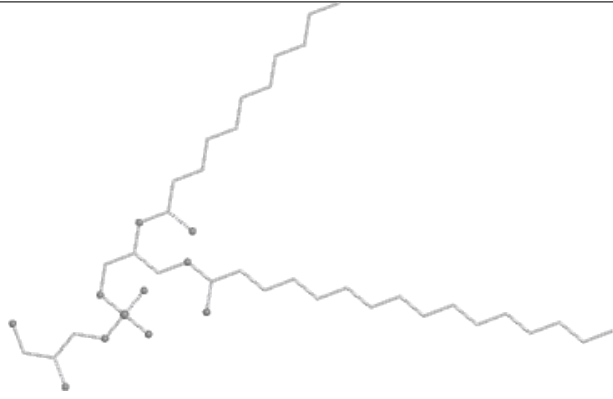


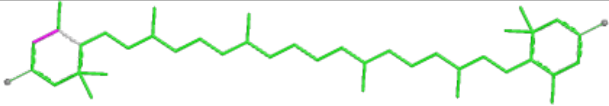
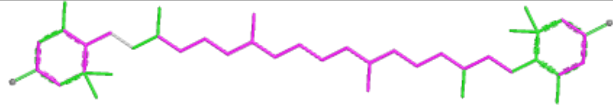
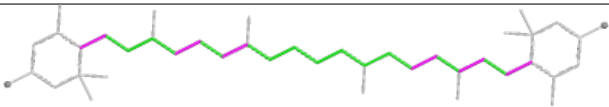
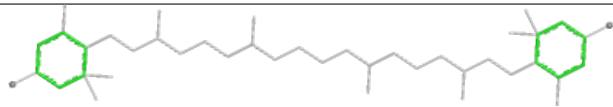


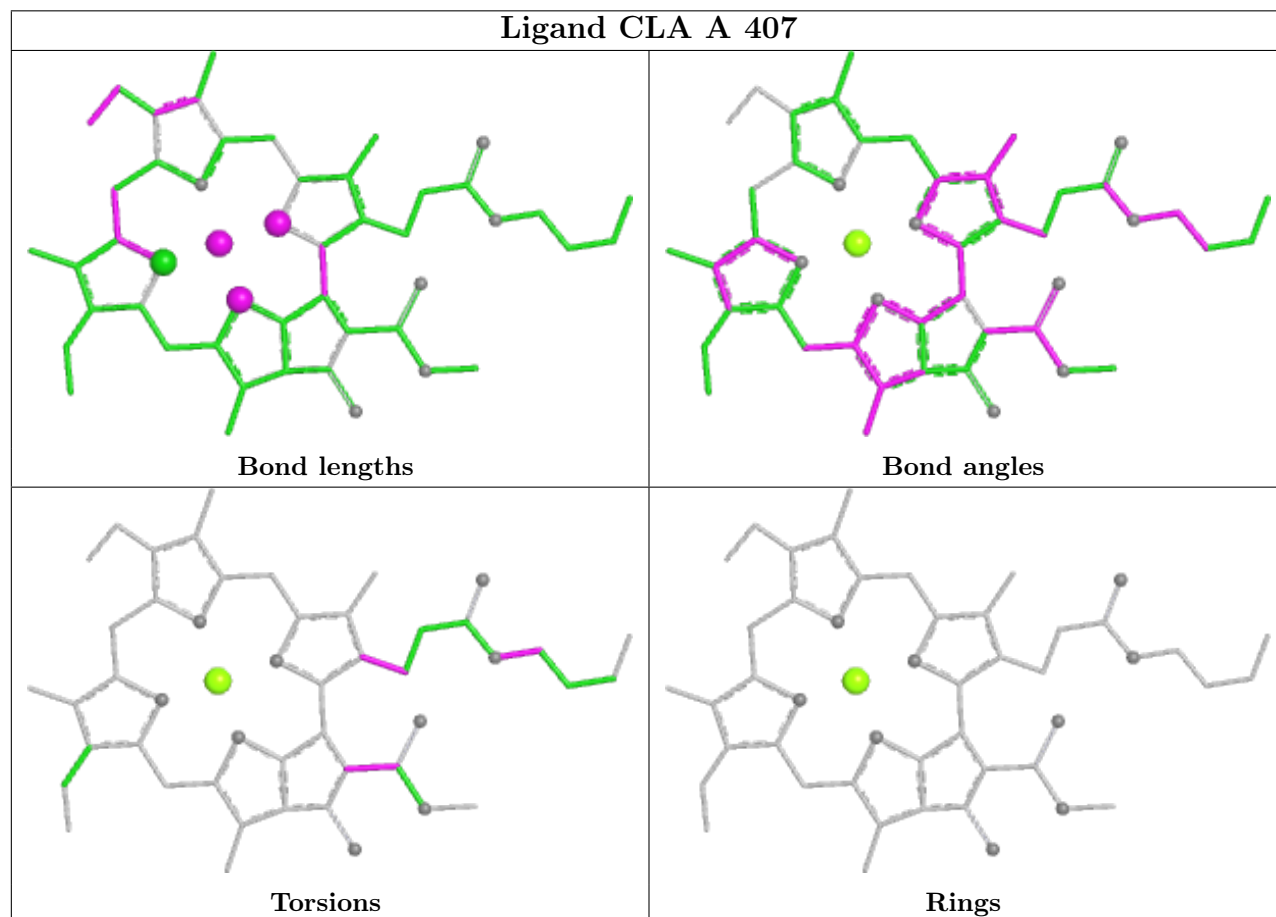
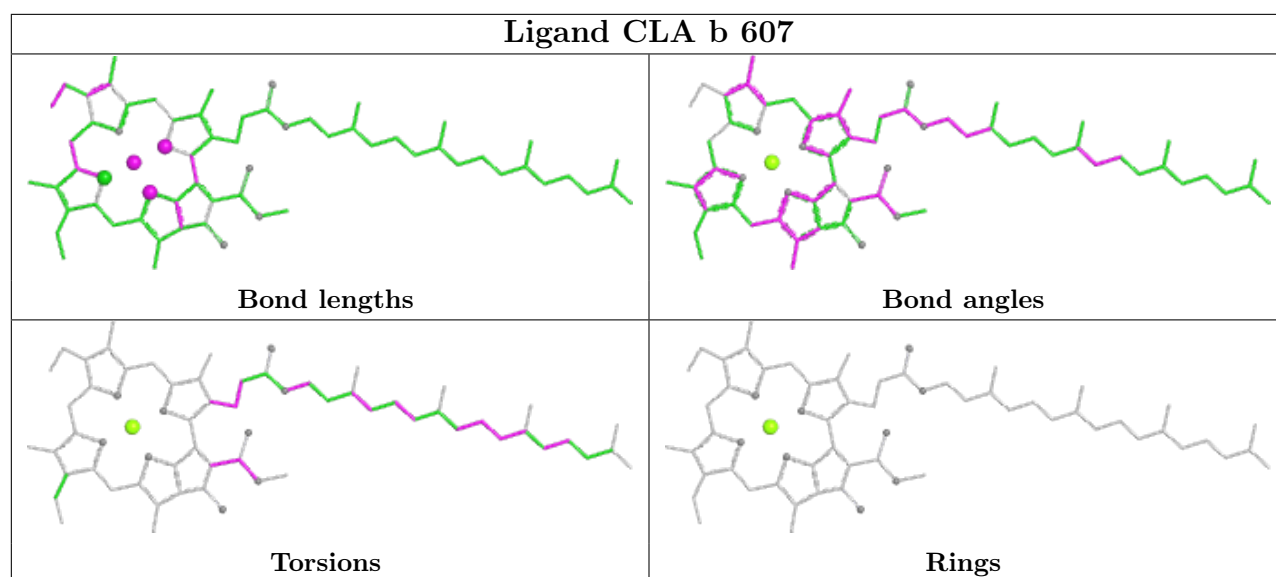


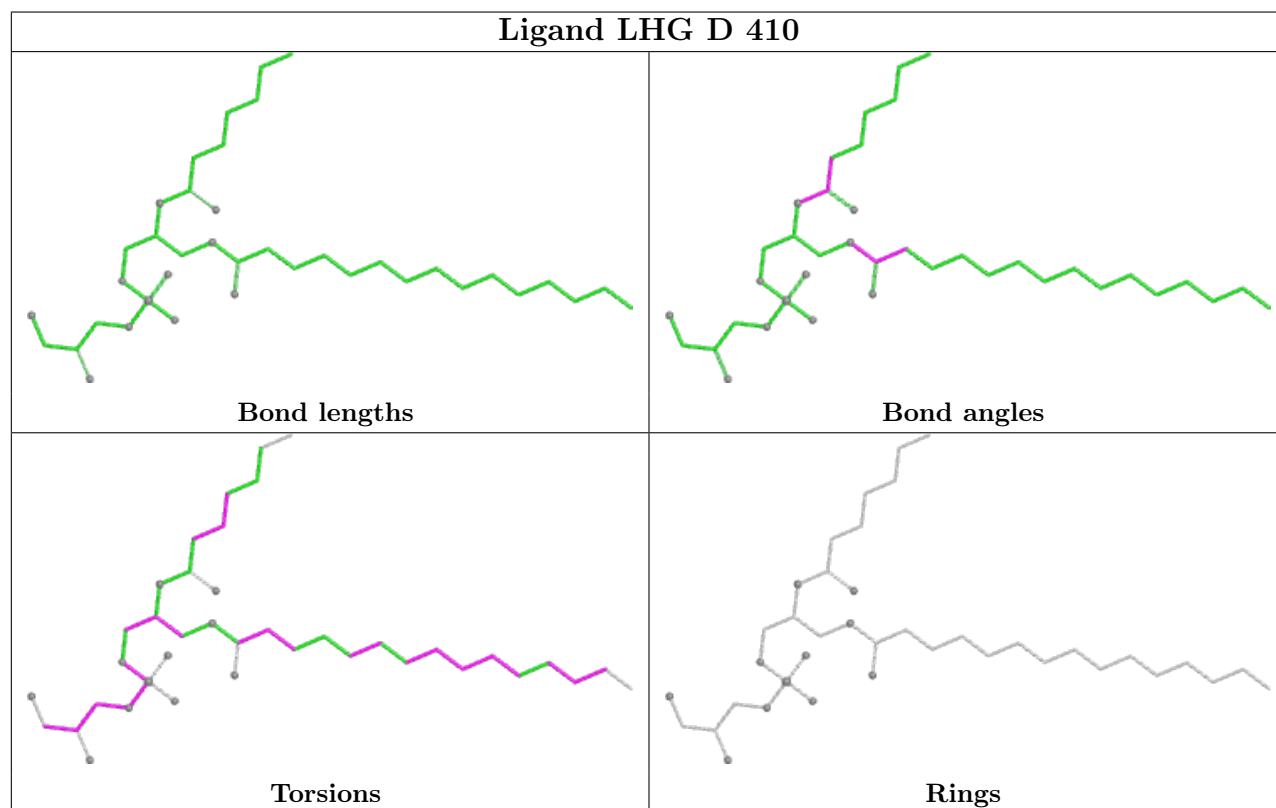
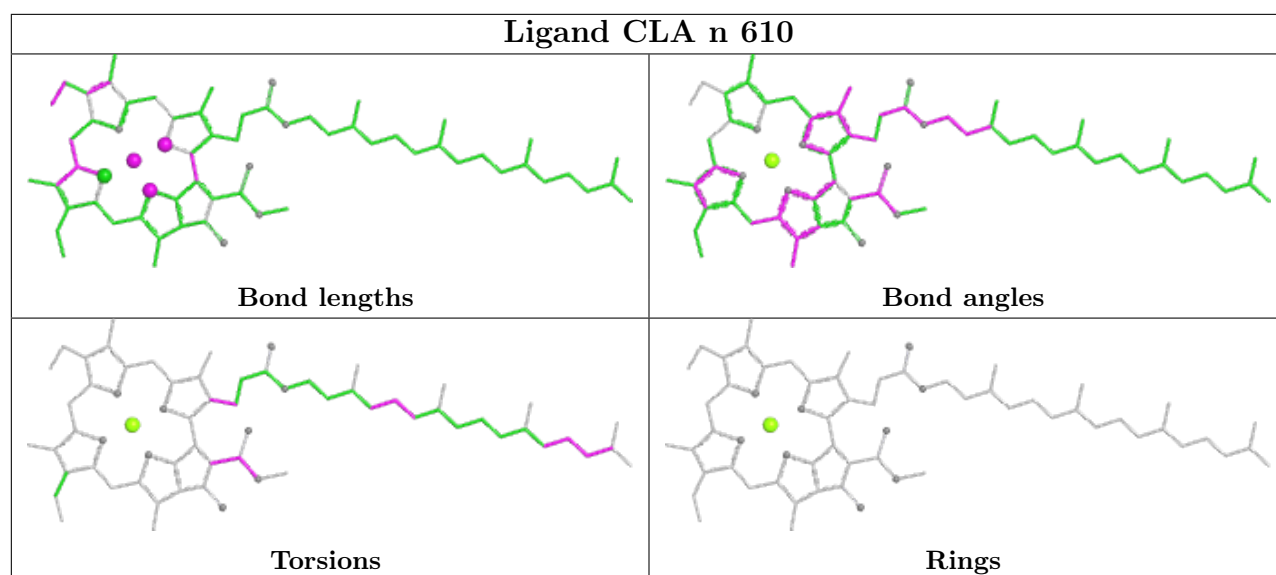


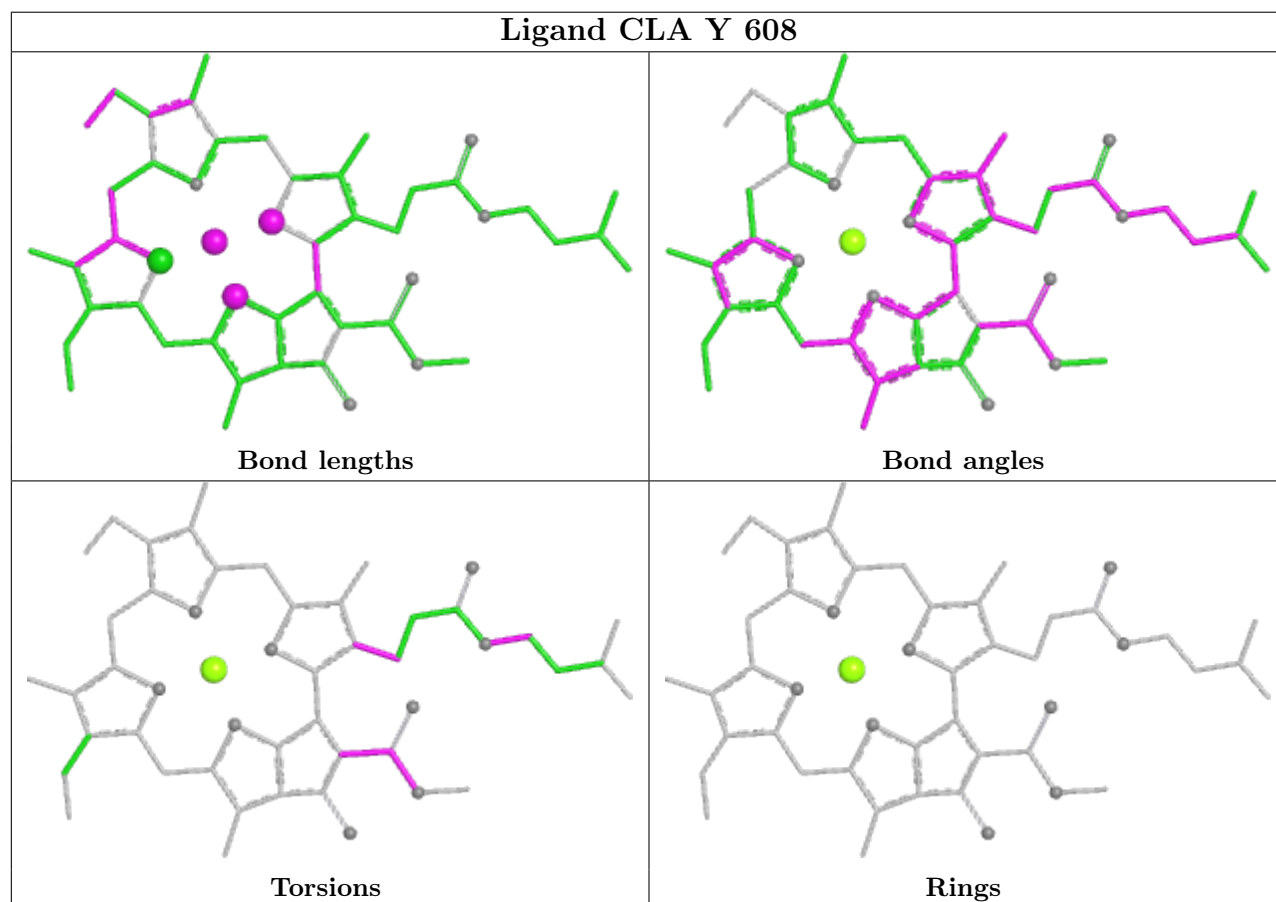
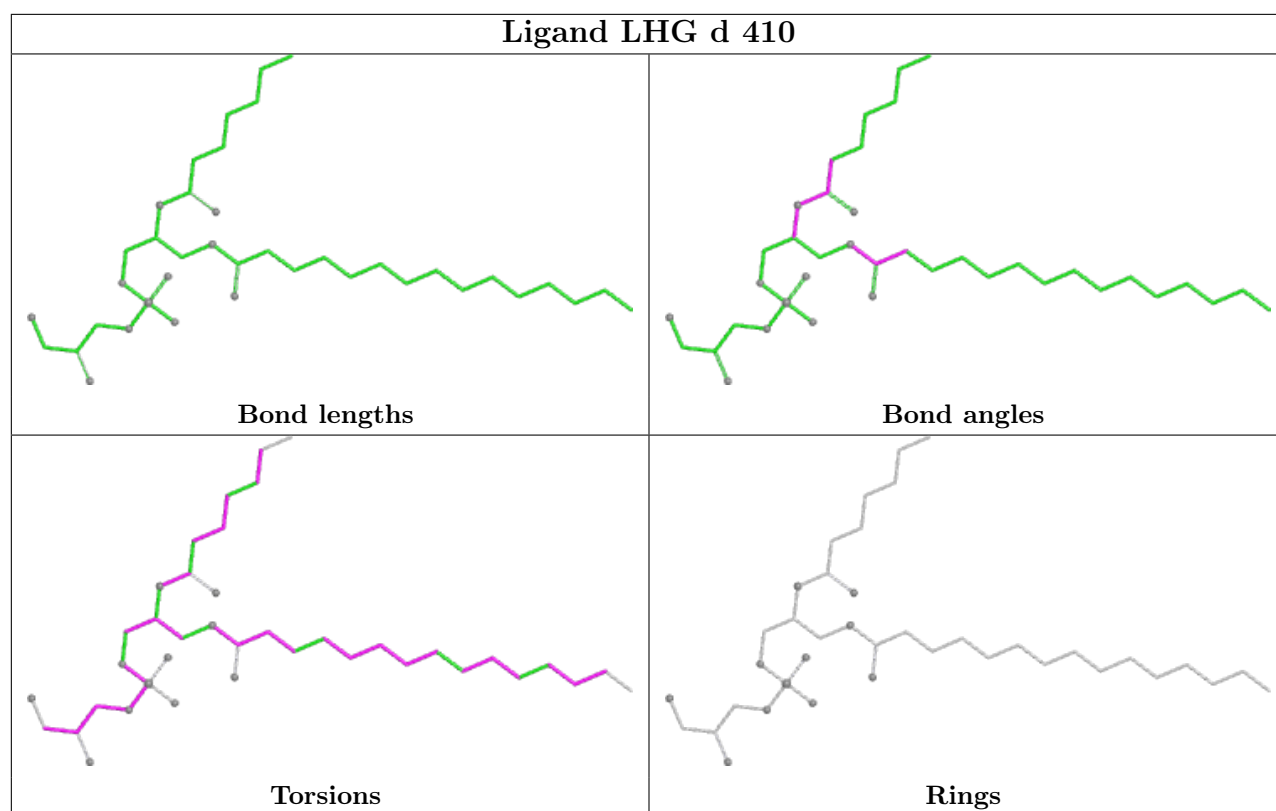


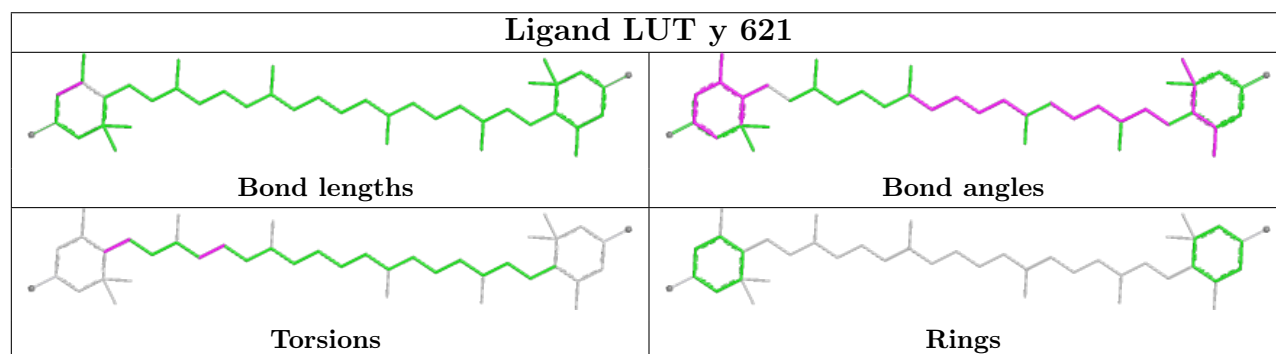
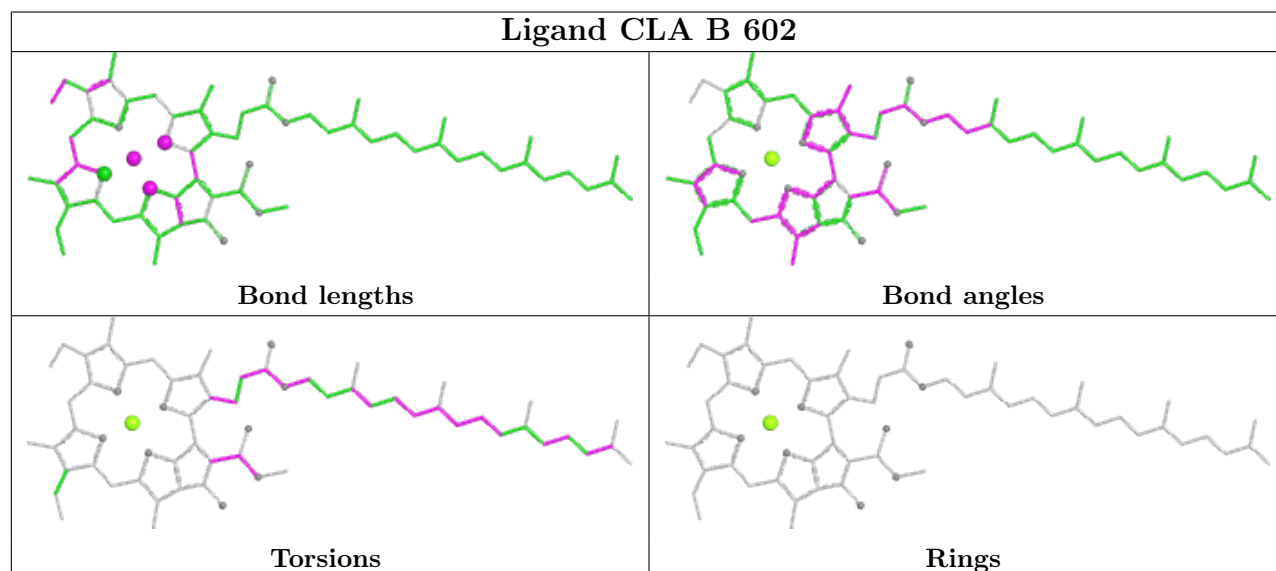
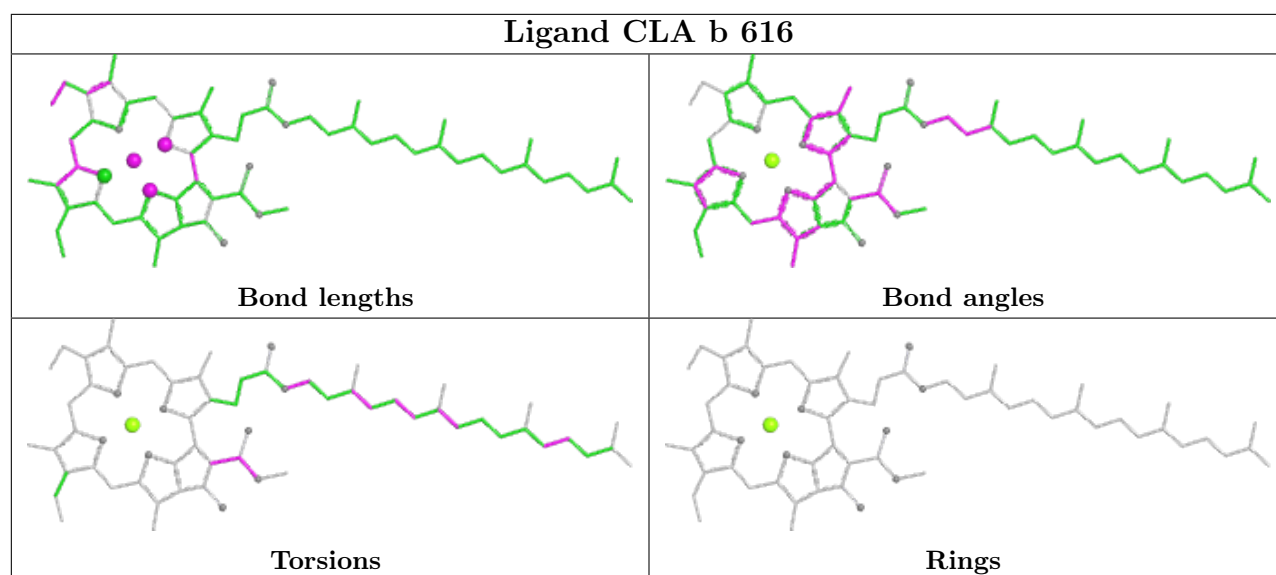
Ligand LHG d 408	
	
Bond lengths	Bond angles
	
Torsions	Rings

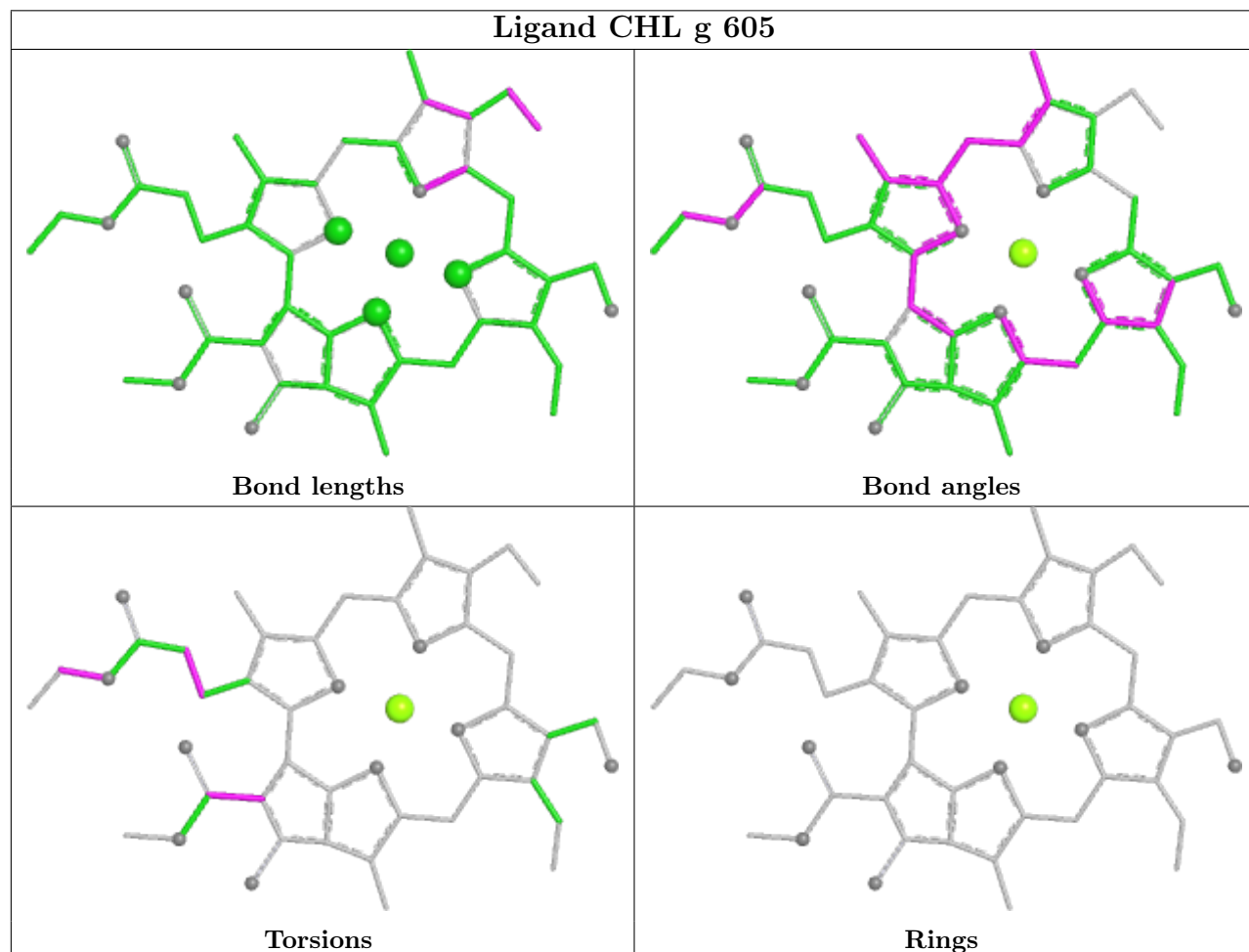
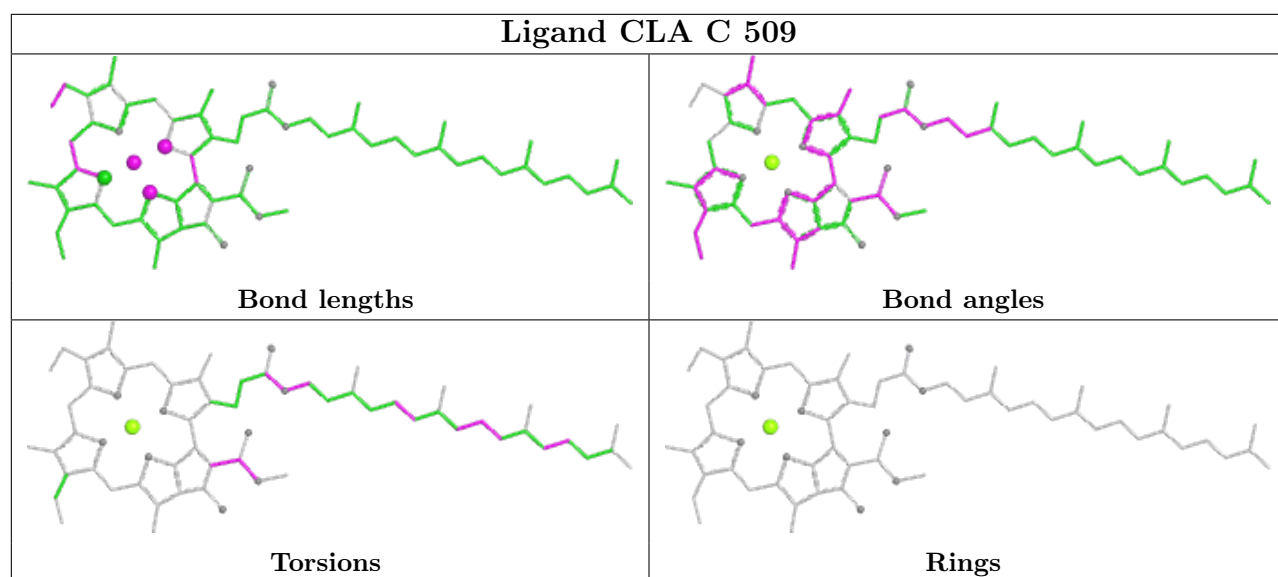
Ligand LUT r 620	
	
Bond lengths	Bond angles
	
Torsions	Rings

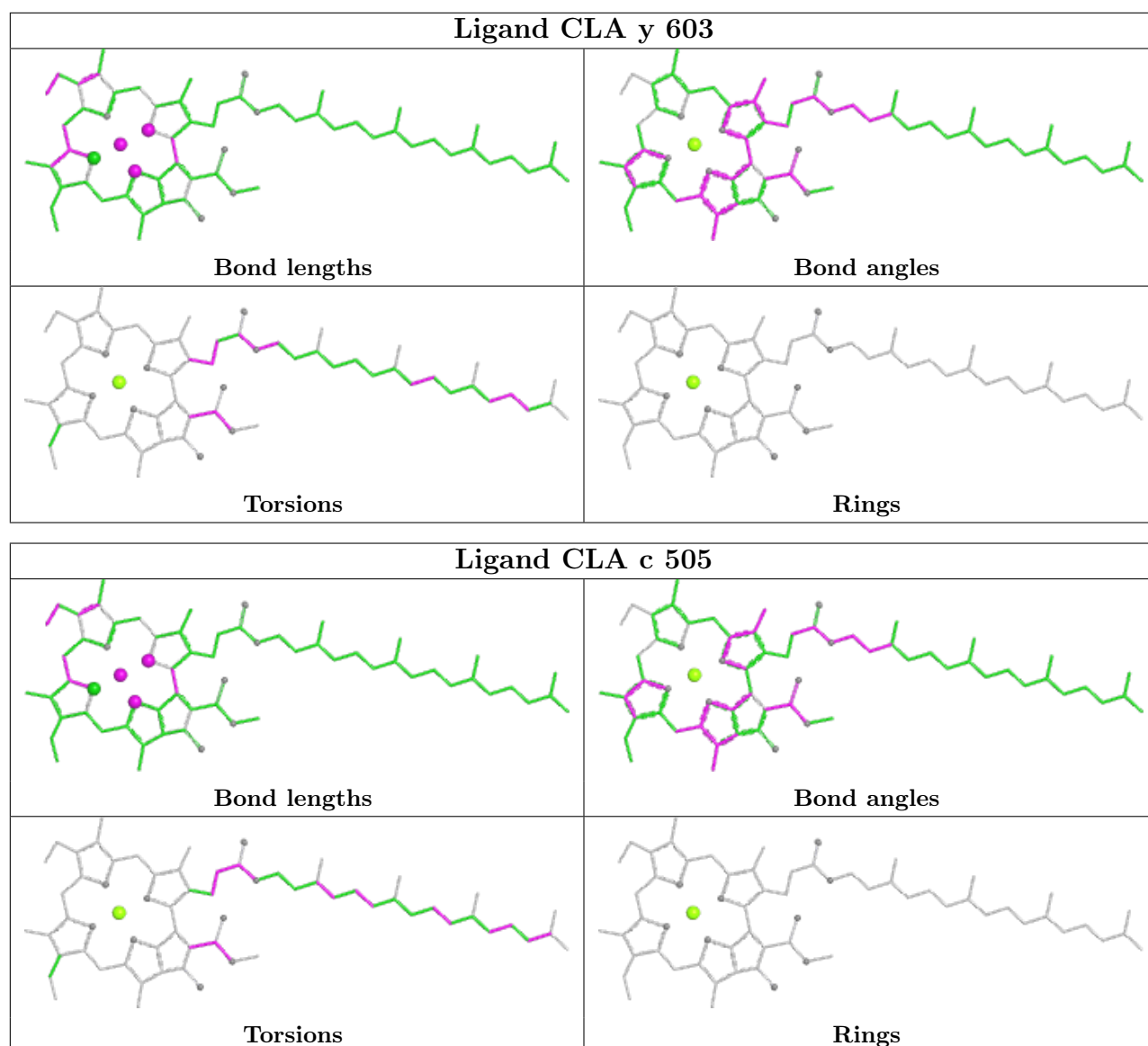


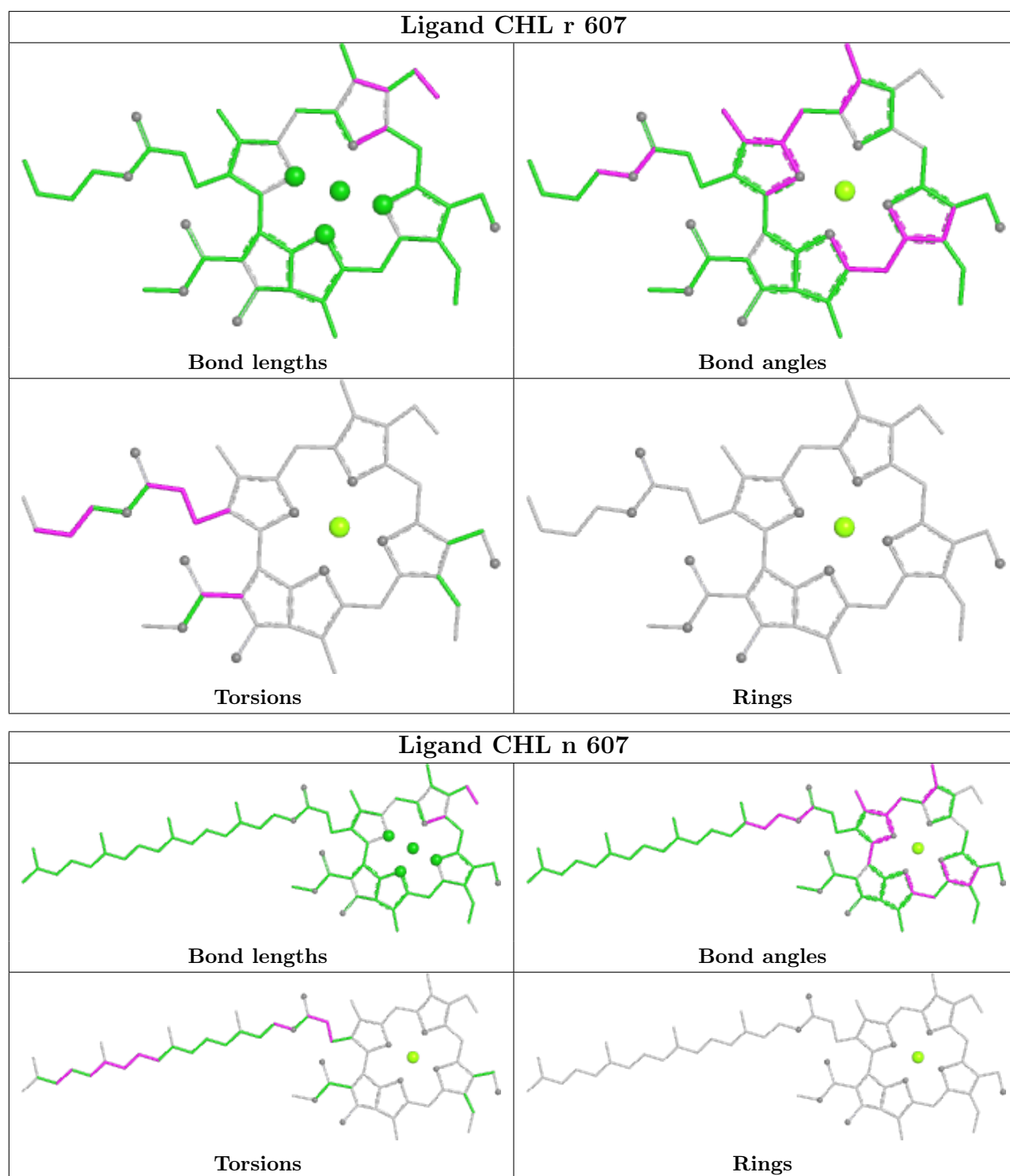


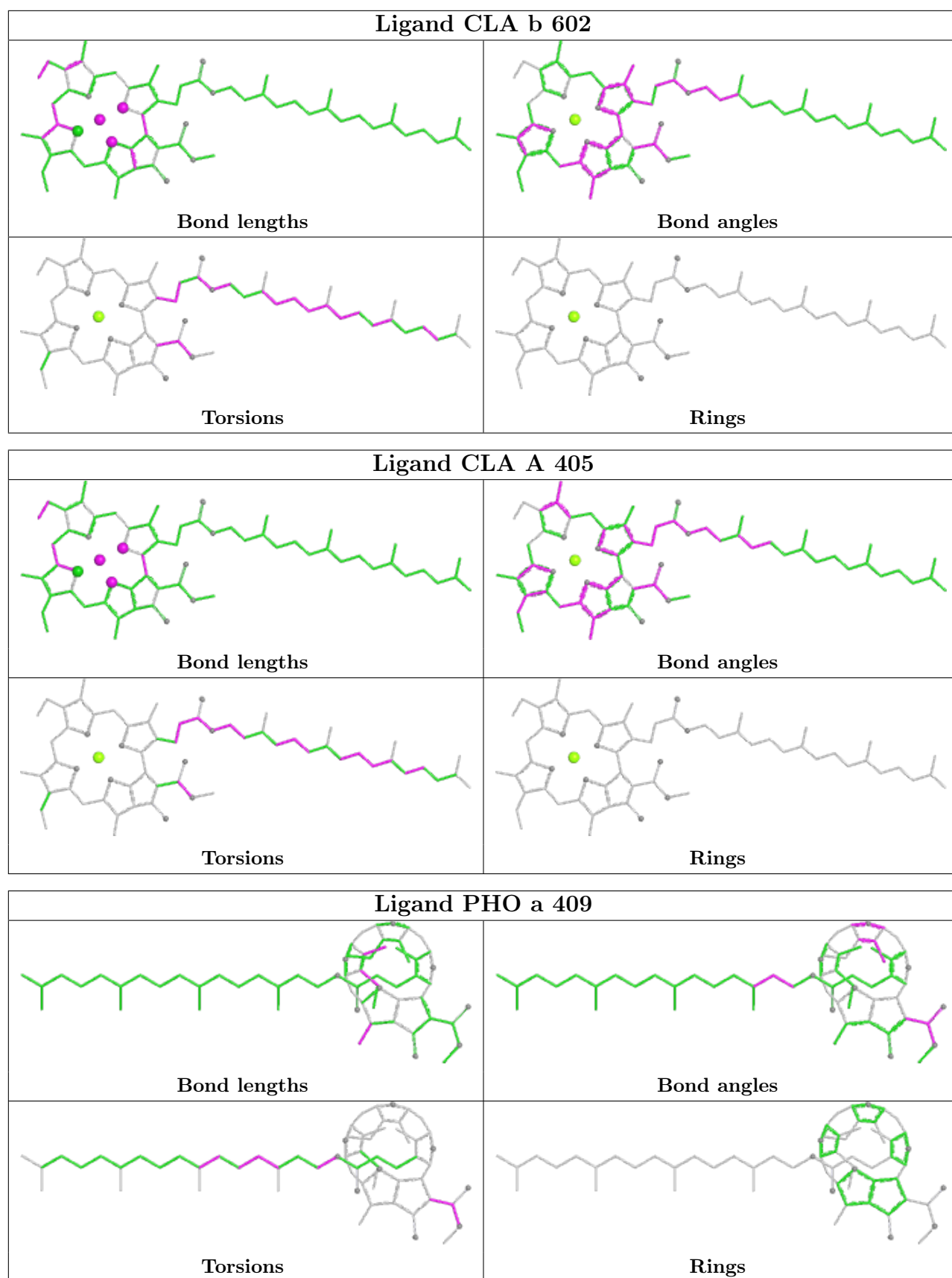


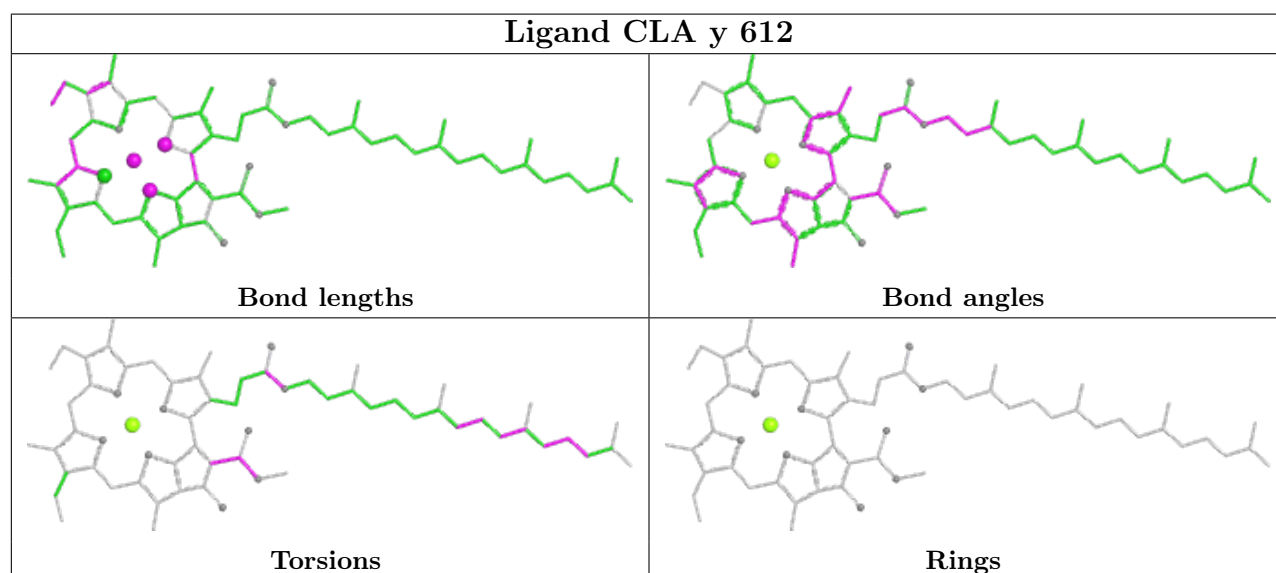
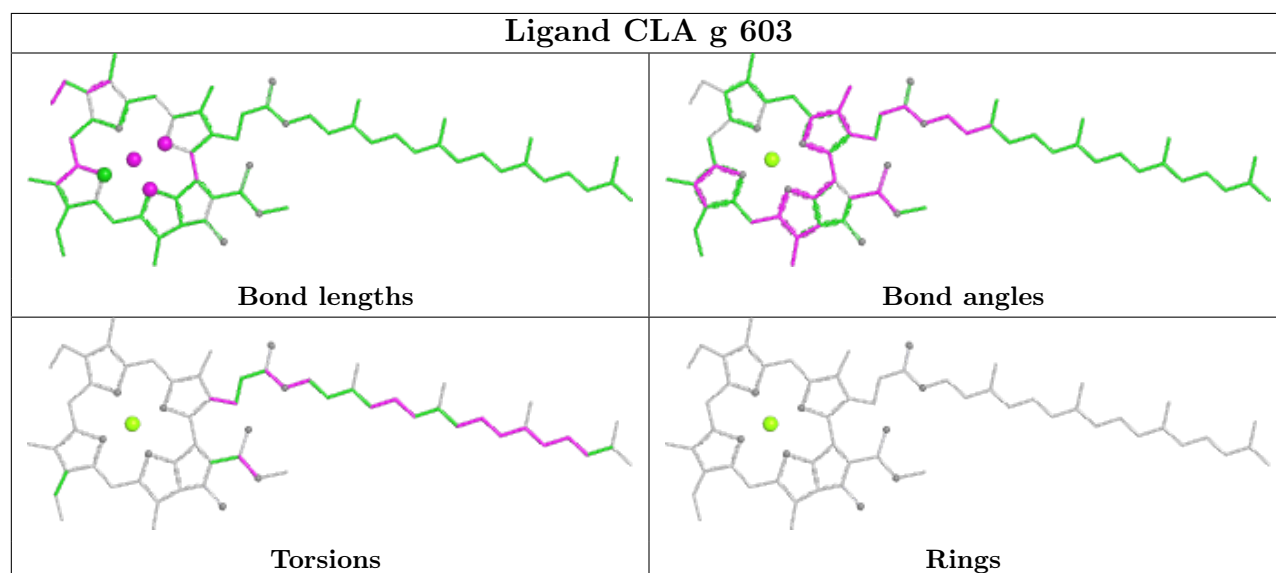
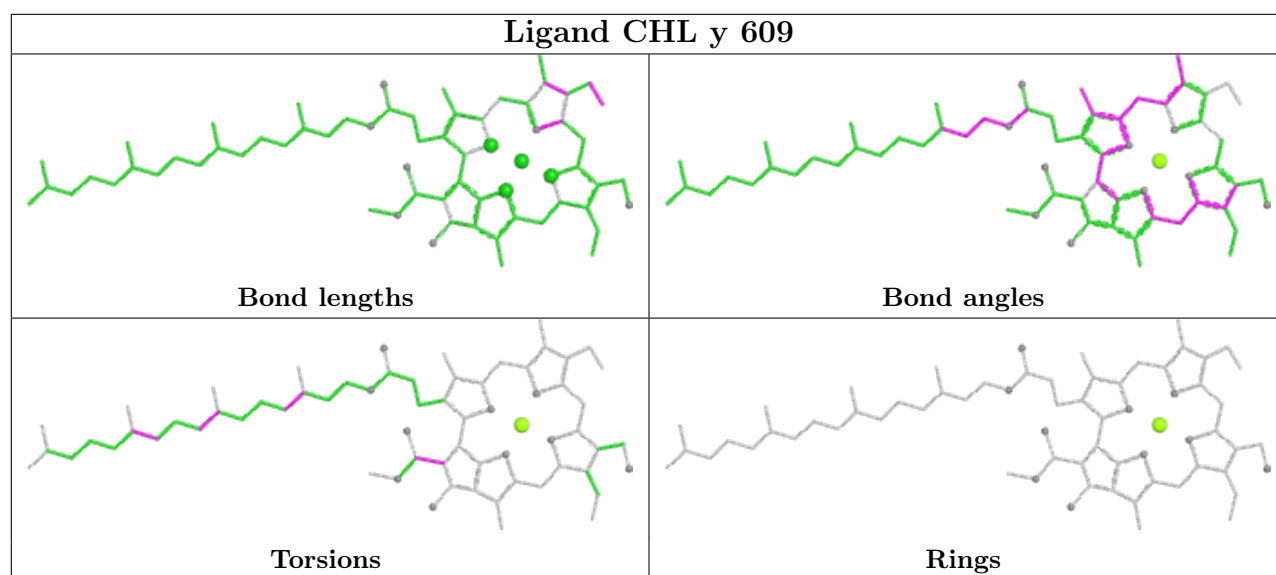


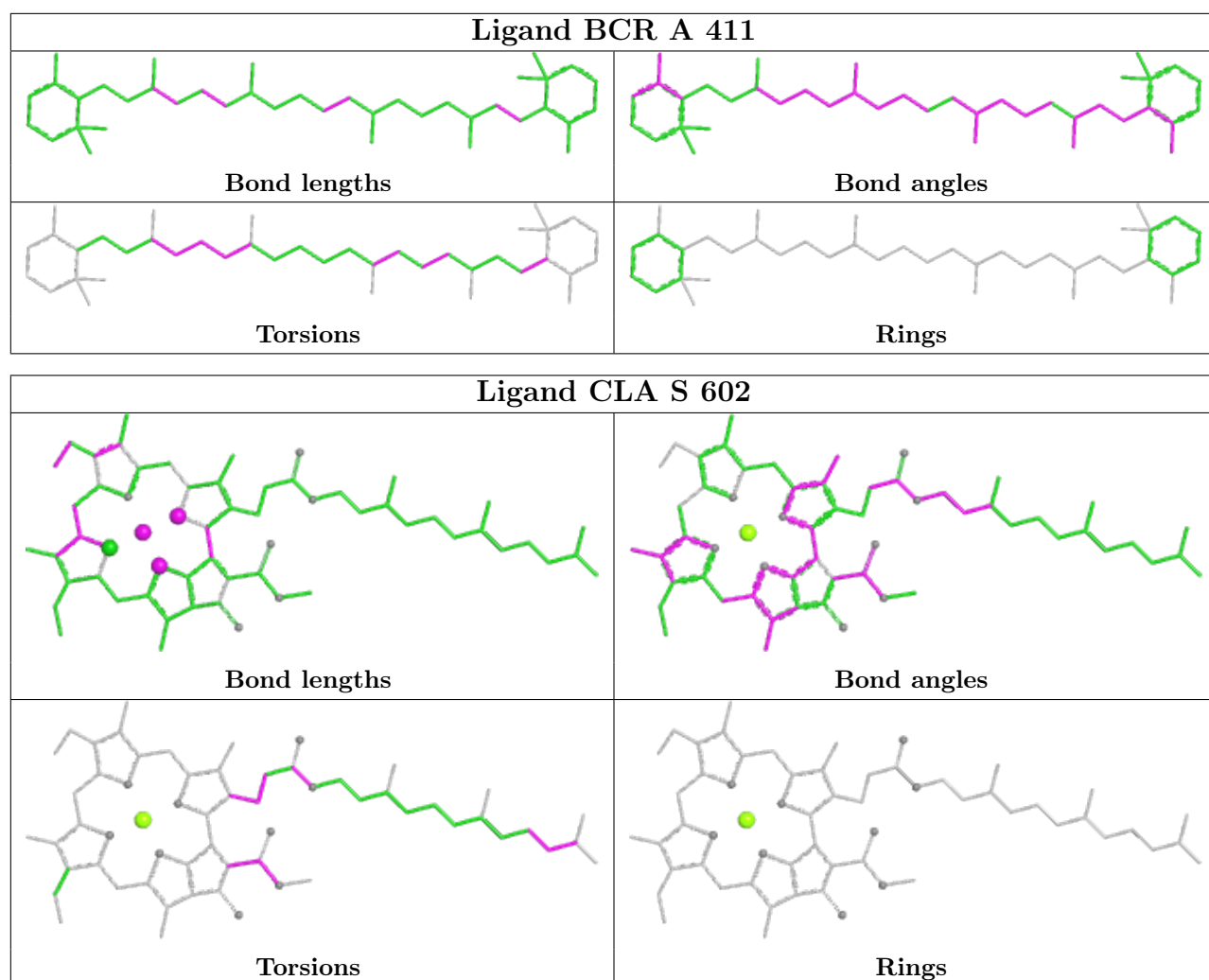


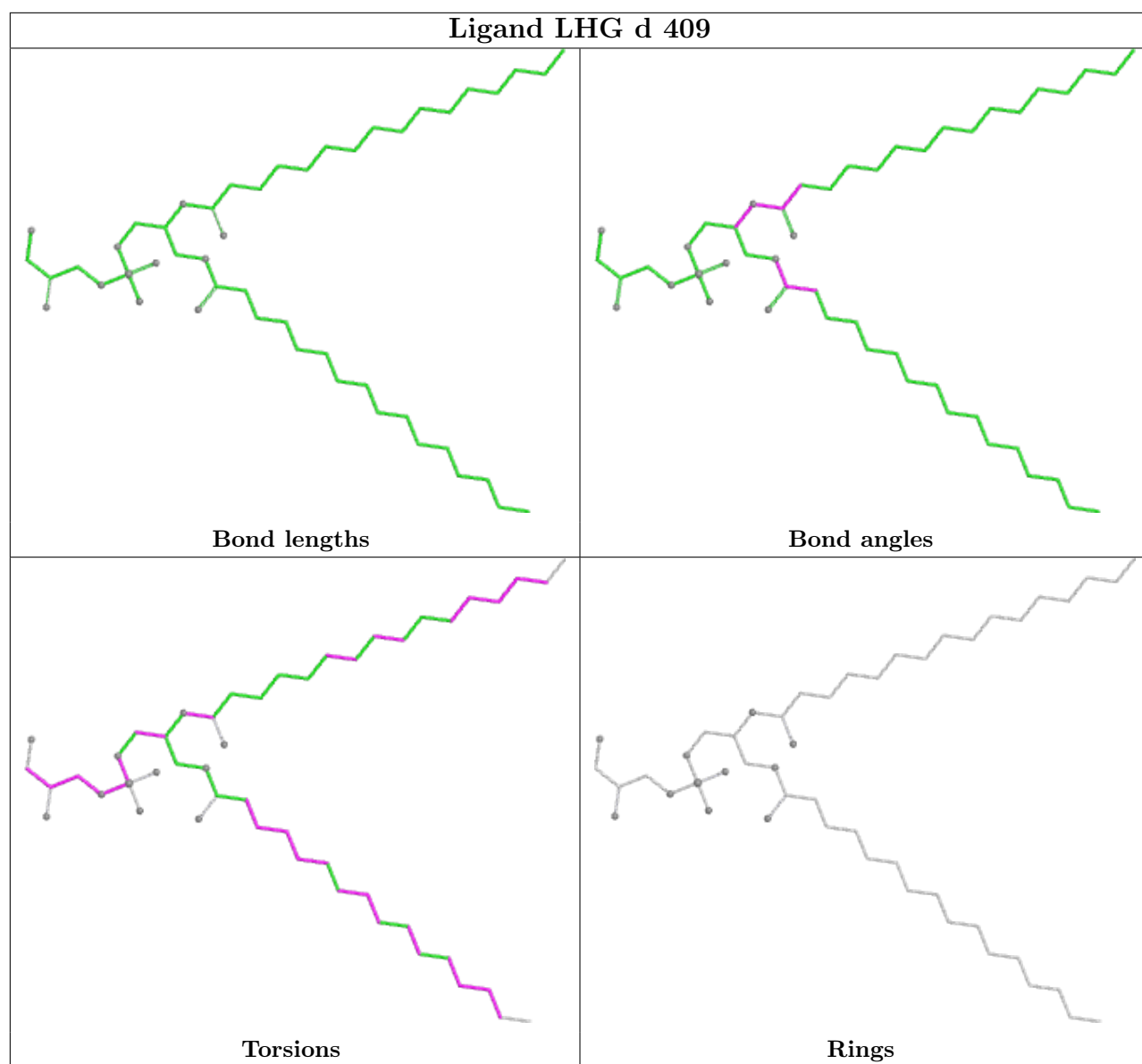




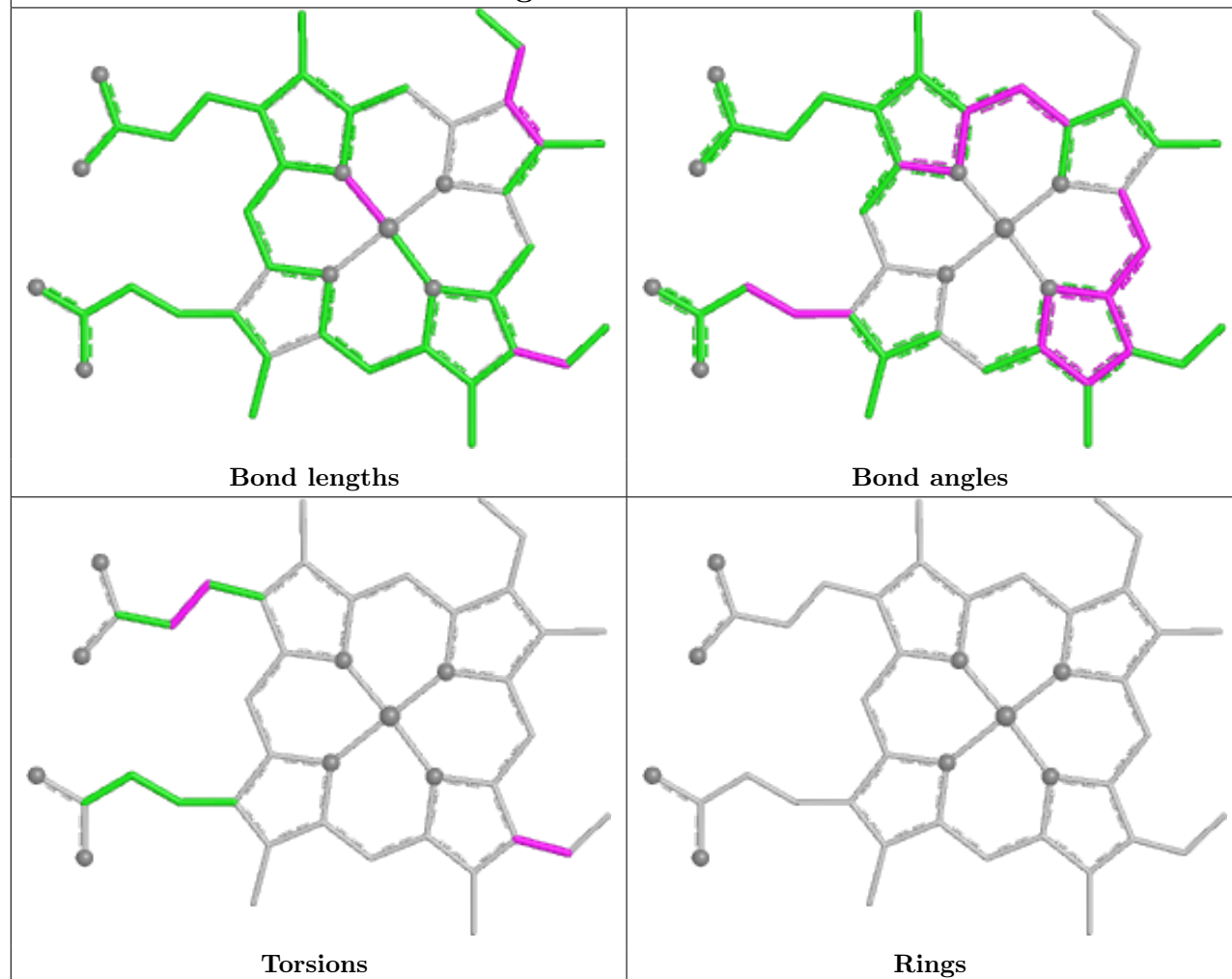




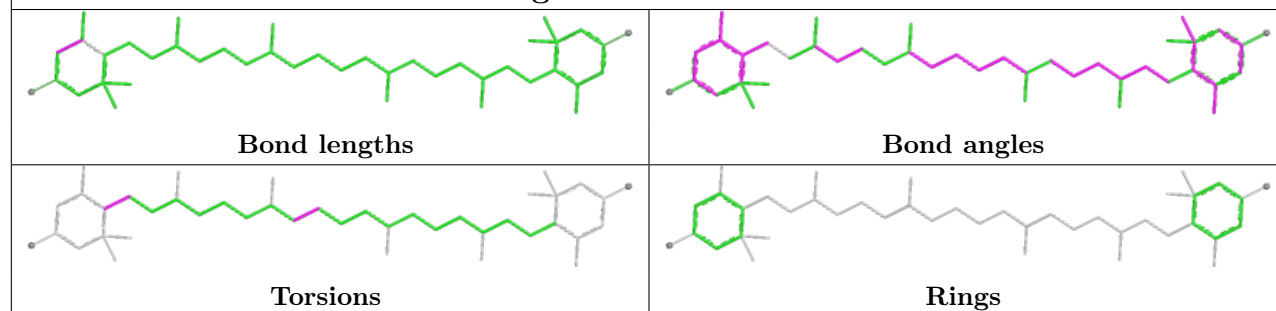


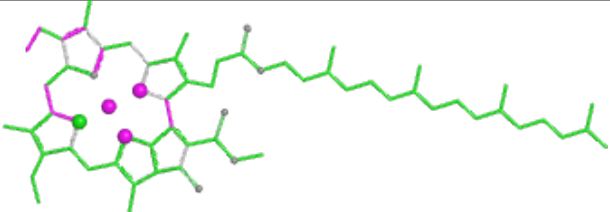
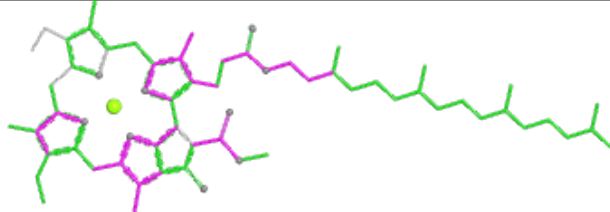
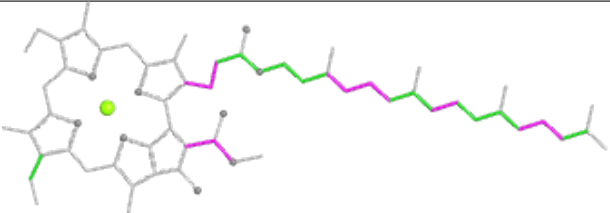
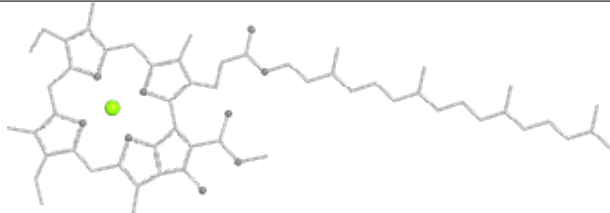
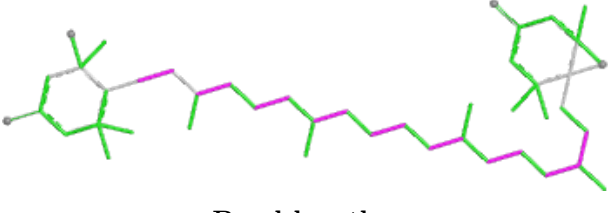
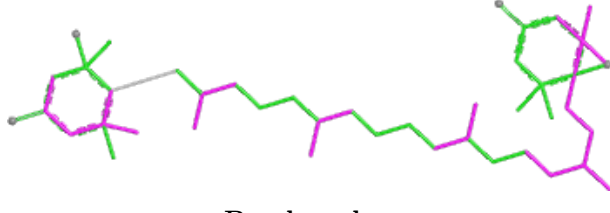

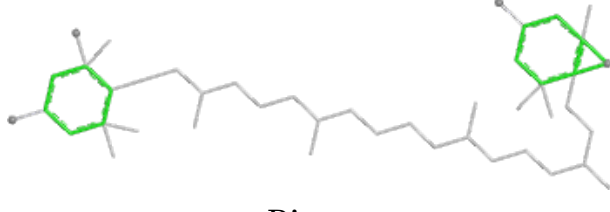
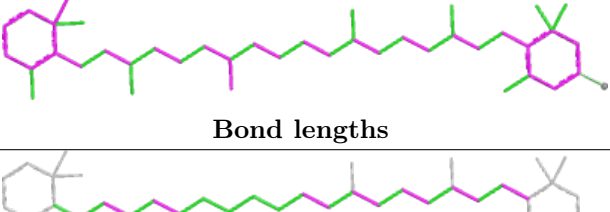
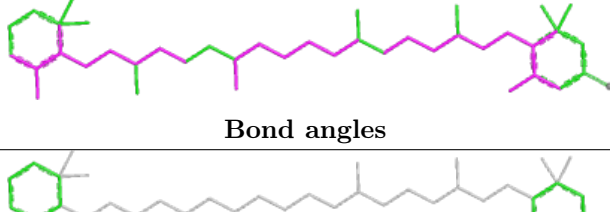
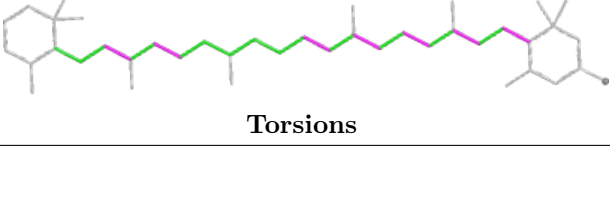
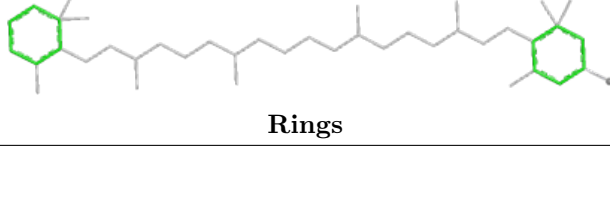


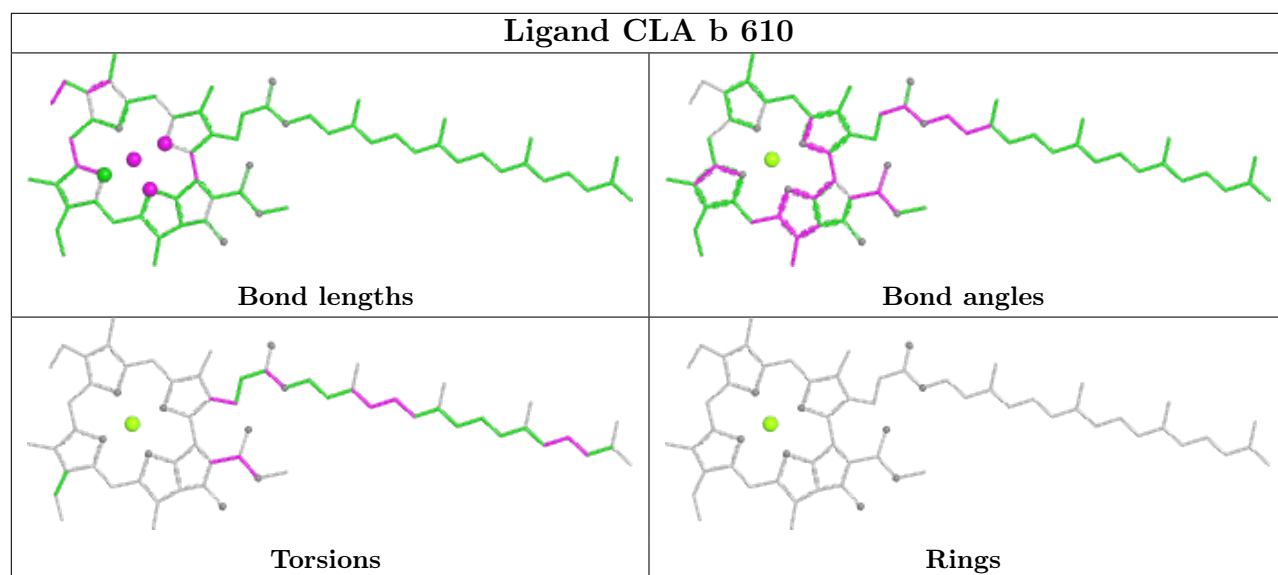
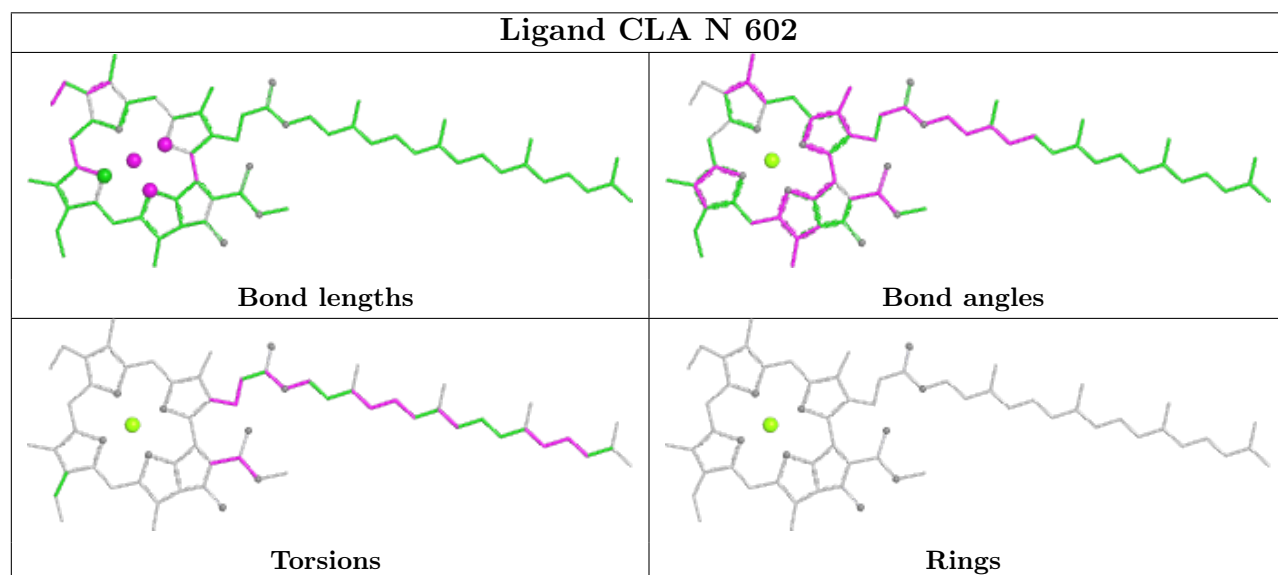
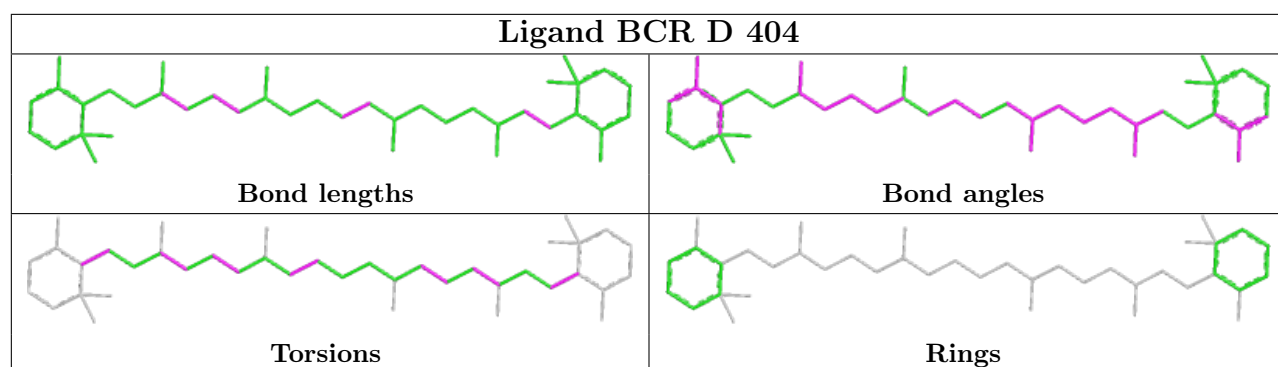
Ligand HEM f 101

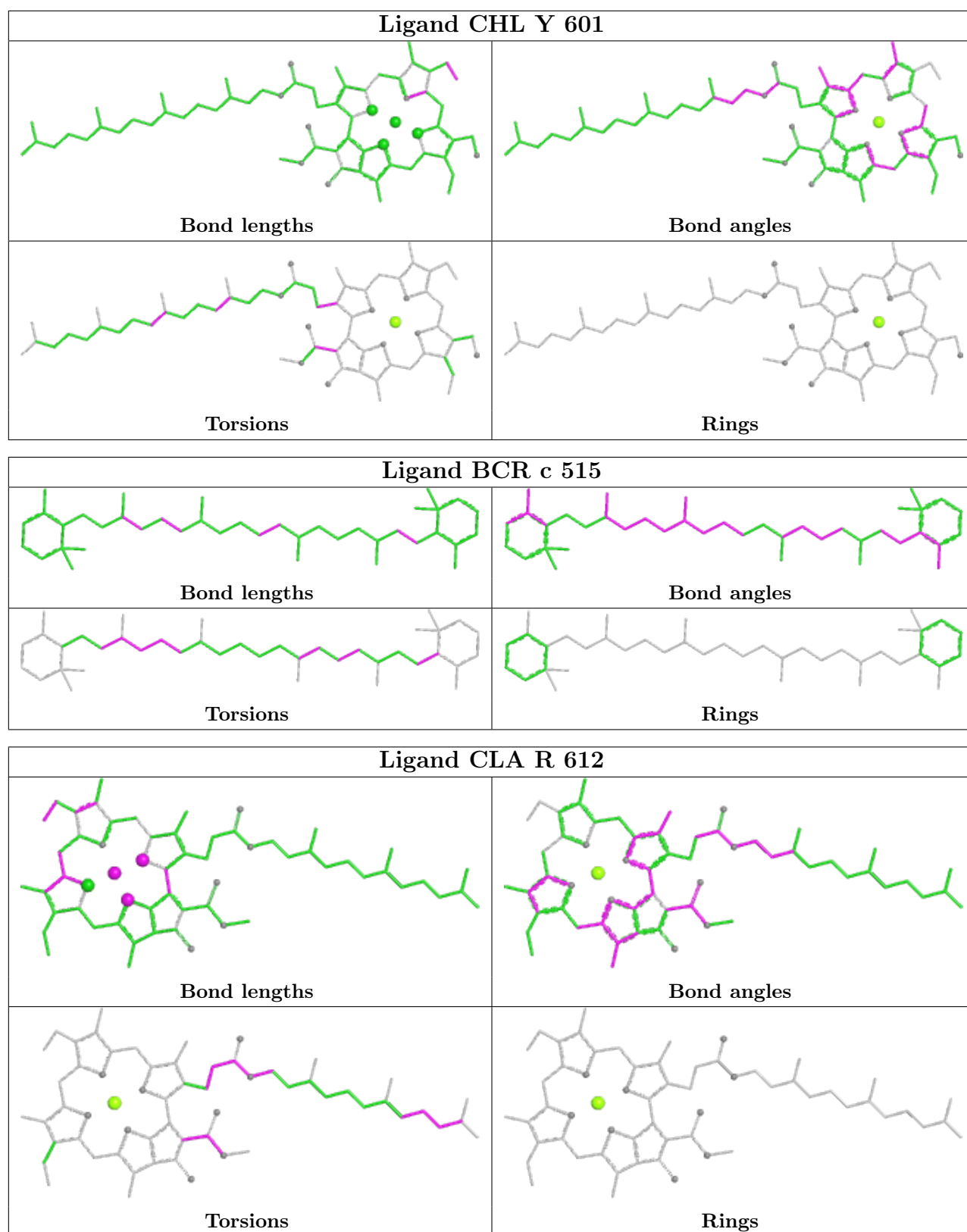


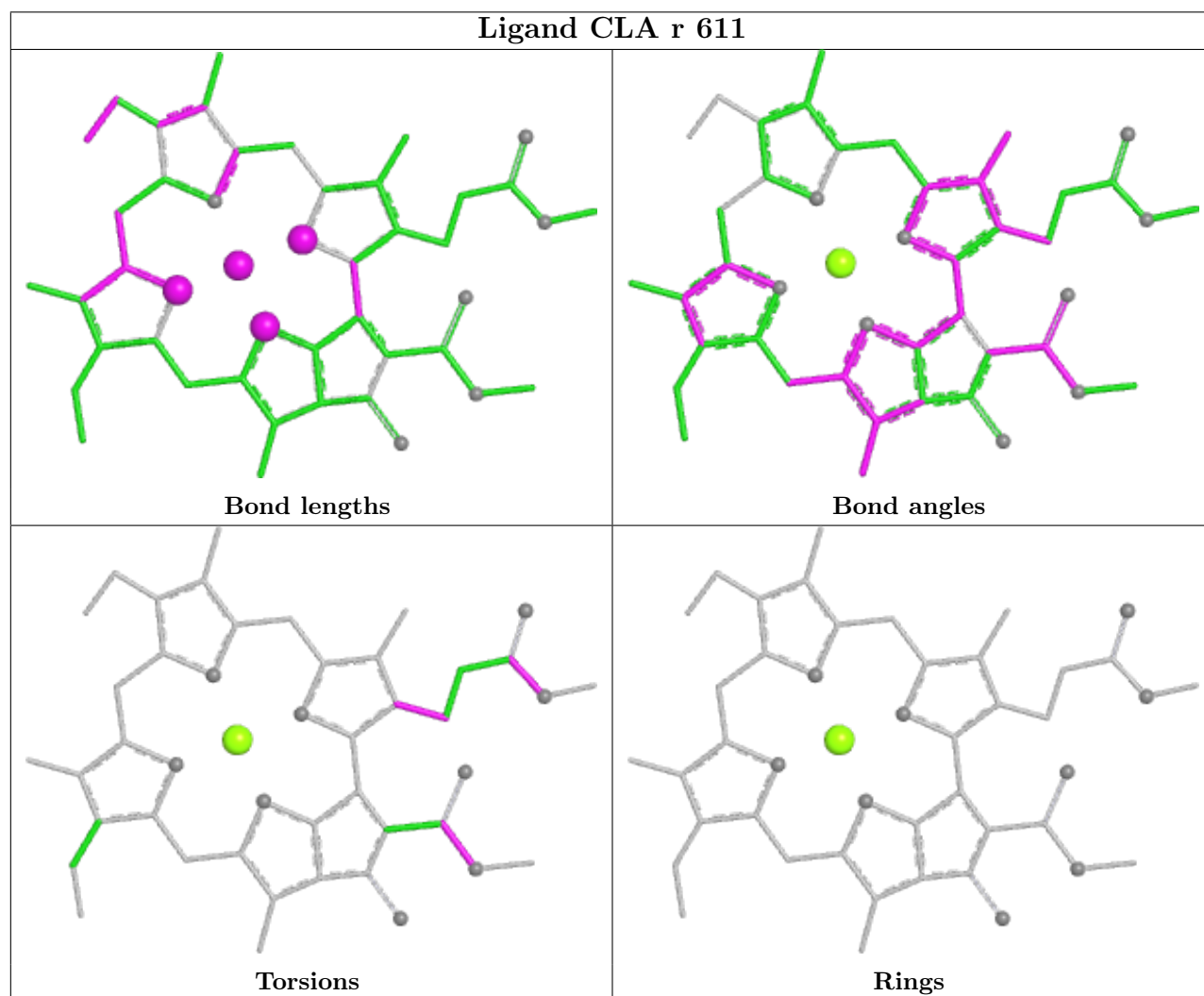
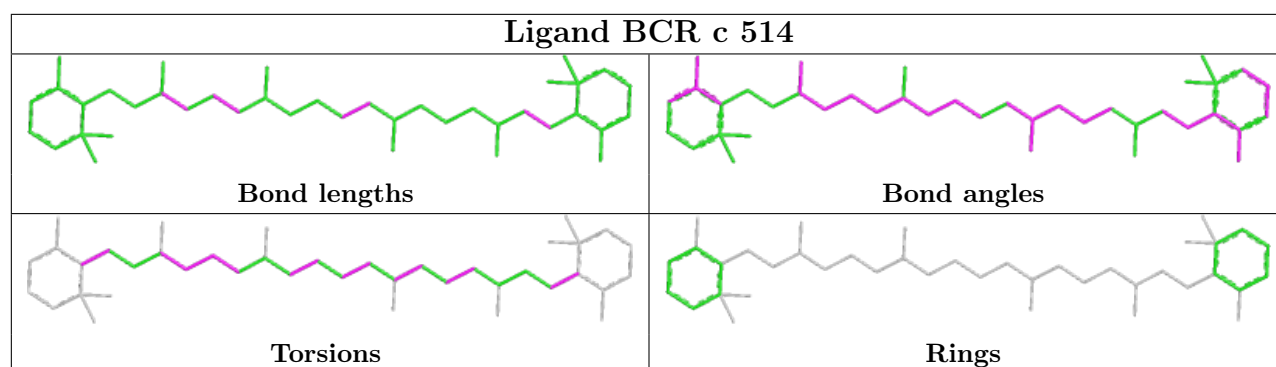
Ligand LUT Y 621

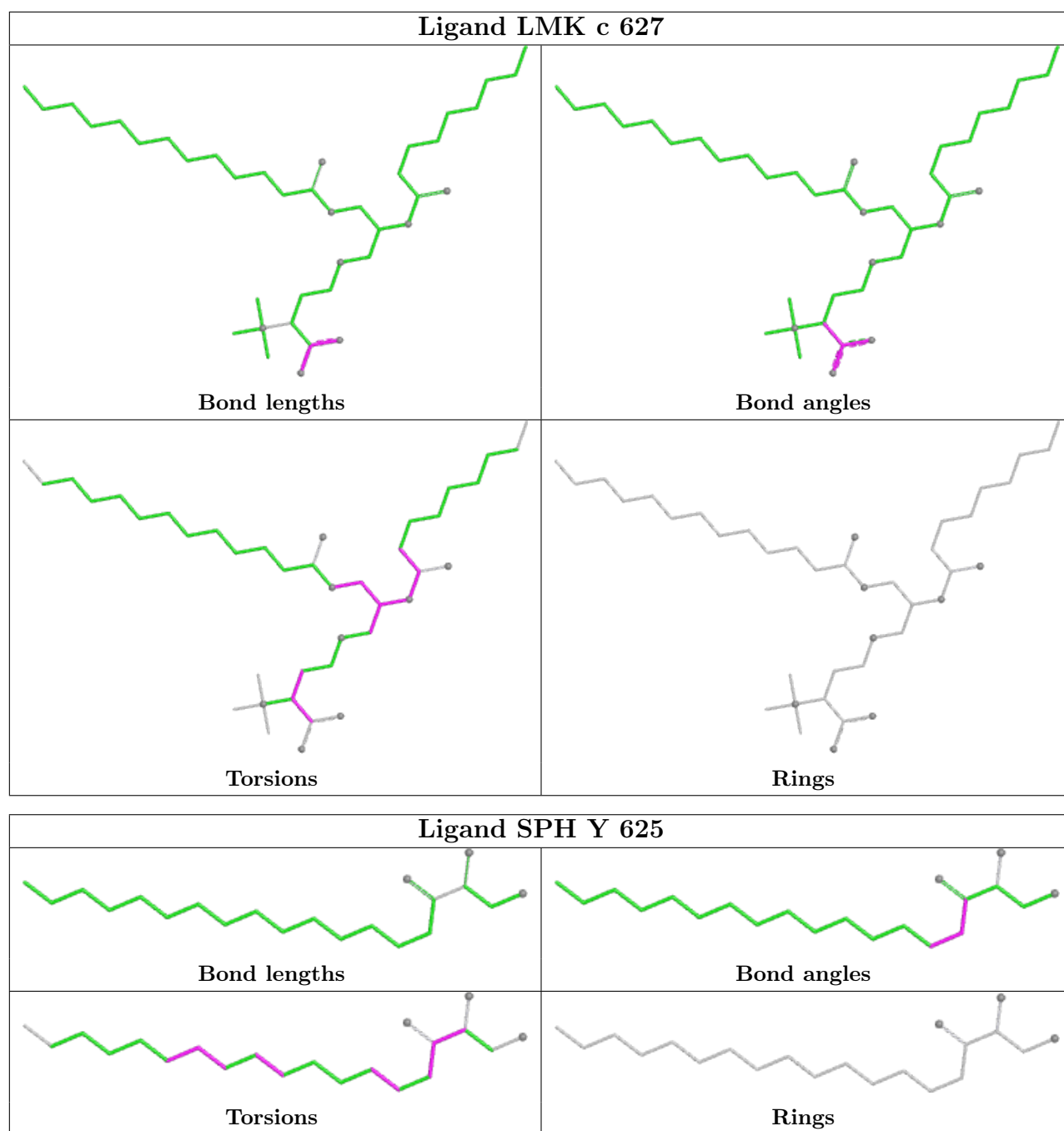


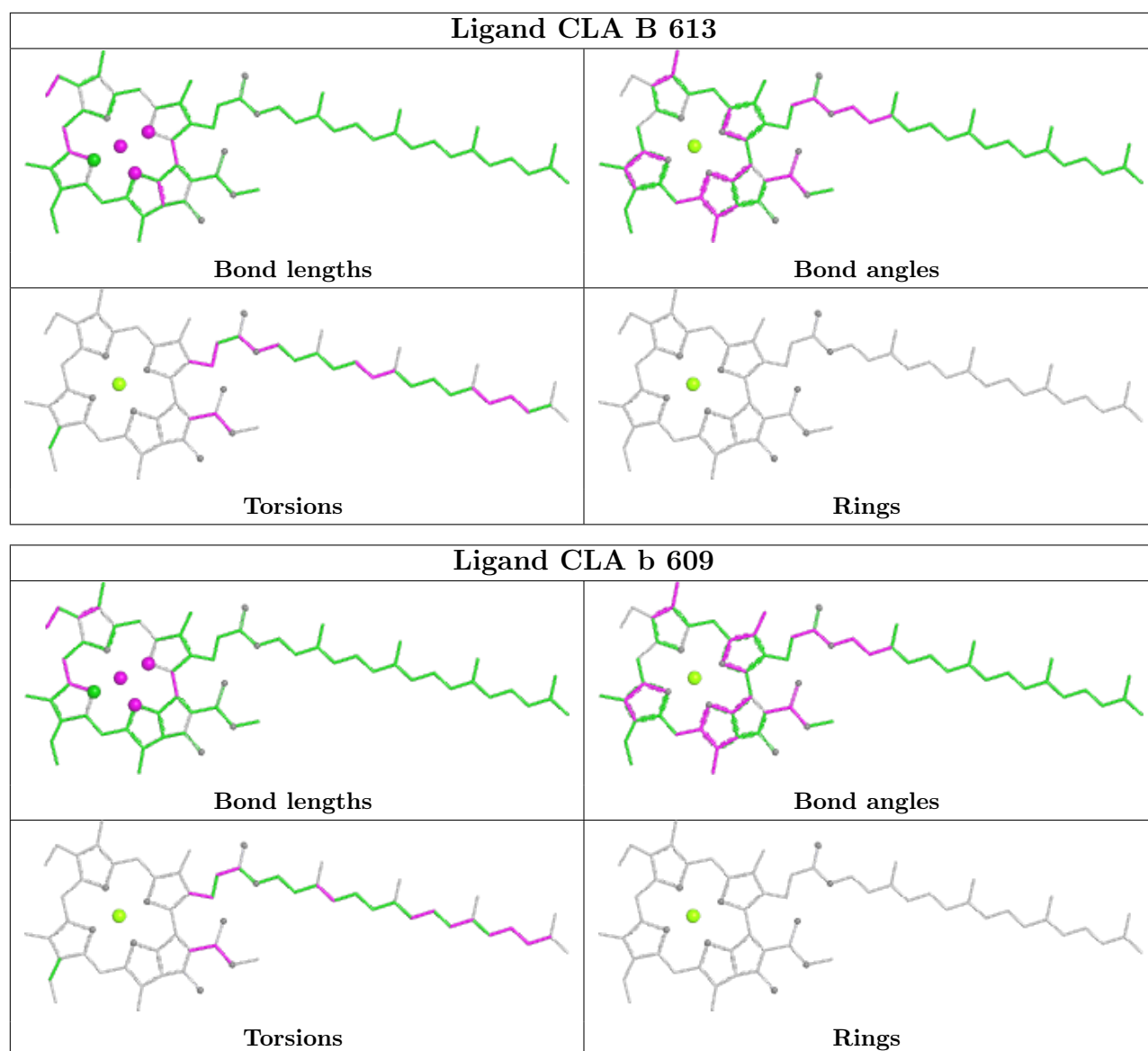
Ligand CLA G 610	
 <p>Bond lengths</p>	 <p>Bond angles</p>
 <p>Torsions</p>	 <p>Rings</p>
Ligand NEX N 623	
 <p>Bond lengths</p>	 <p>Bond angles</p>
 <p>Torsions</p>	 <p>Rings</p>
Ligand RRX h 101	
 <p>Bond lengths</p>	 <p>Bond angles</p>
 <p>Torsions</p>	 <p>Rings</p>

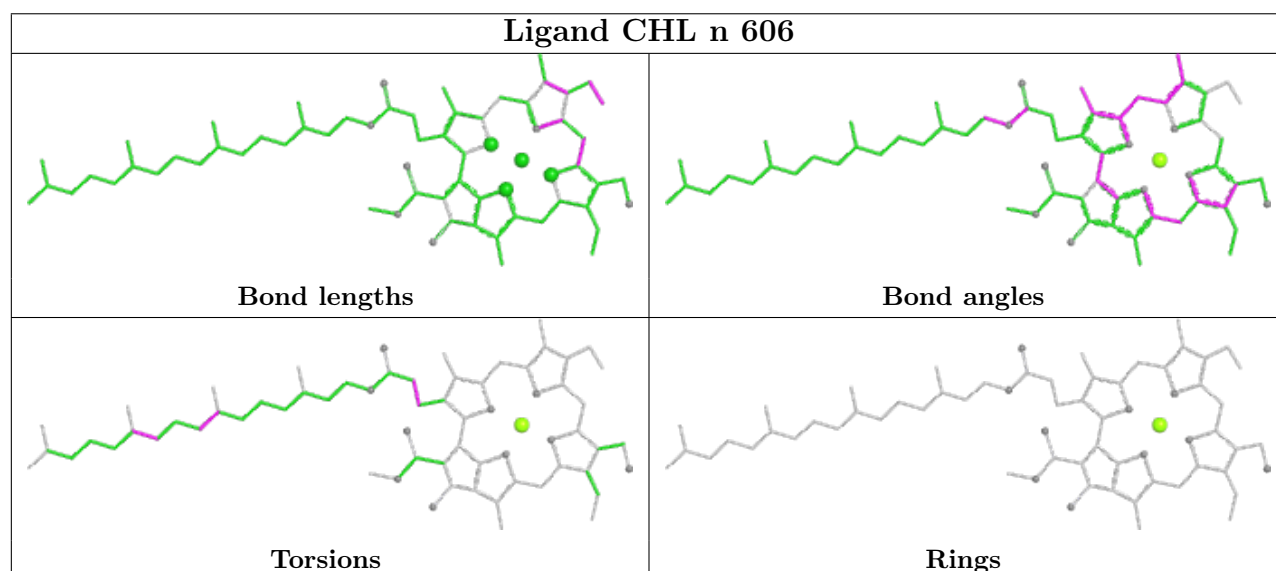
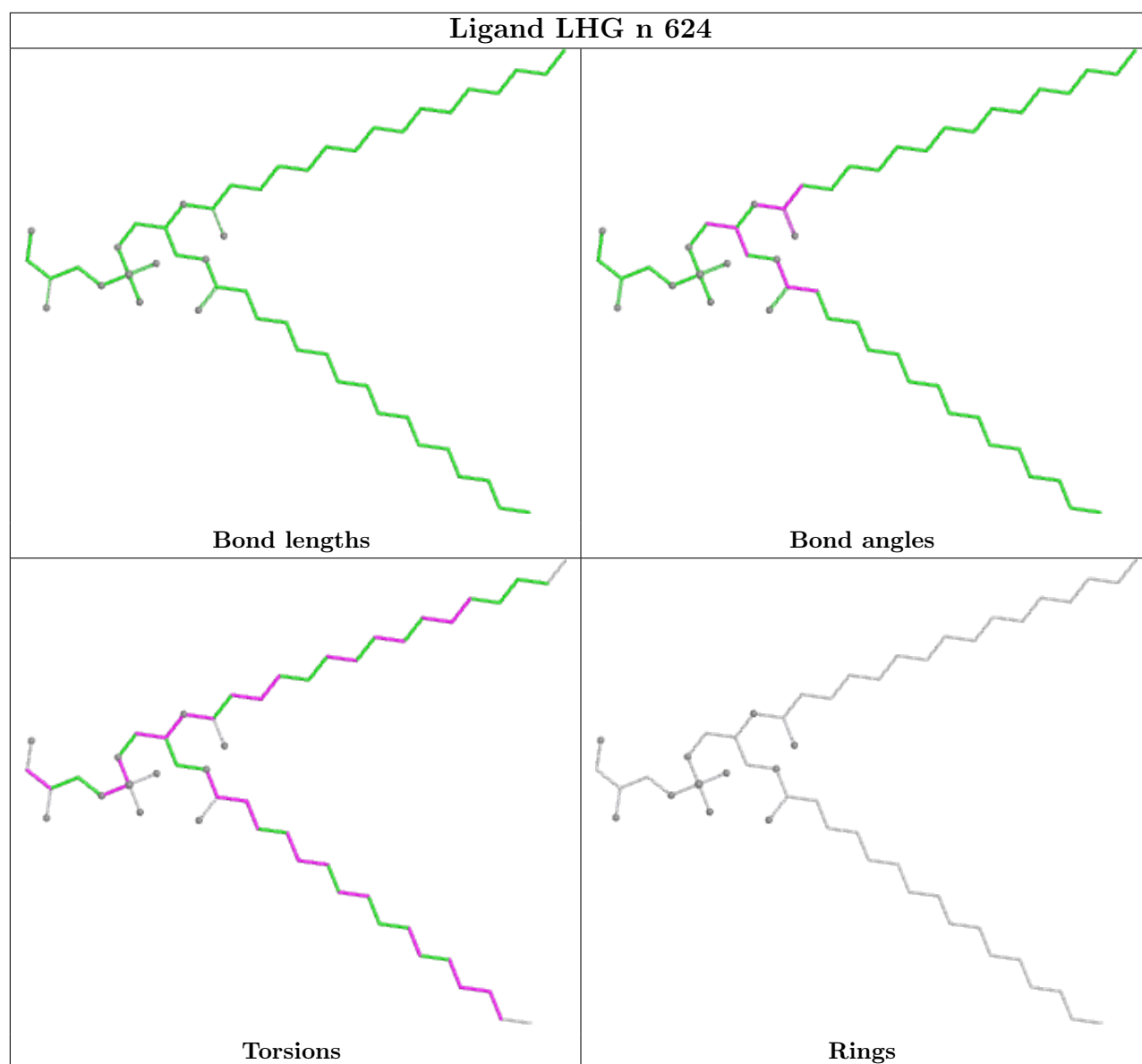


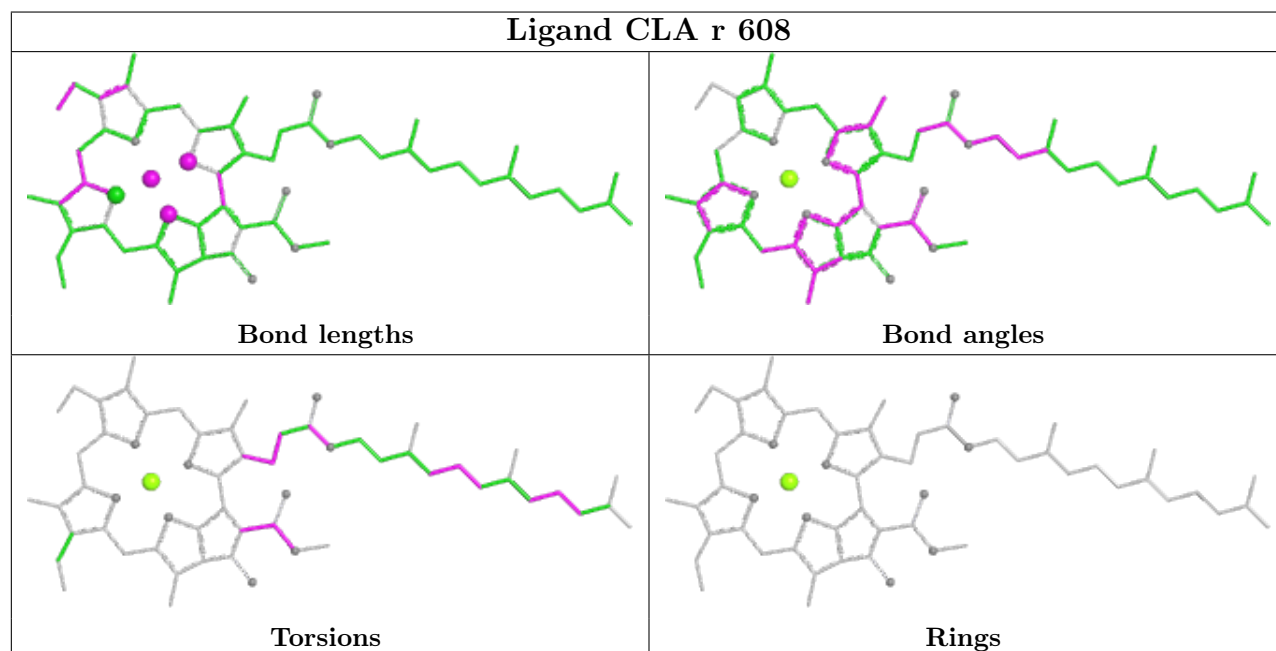
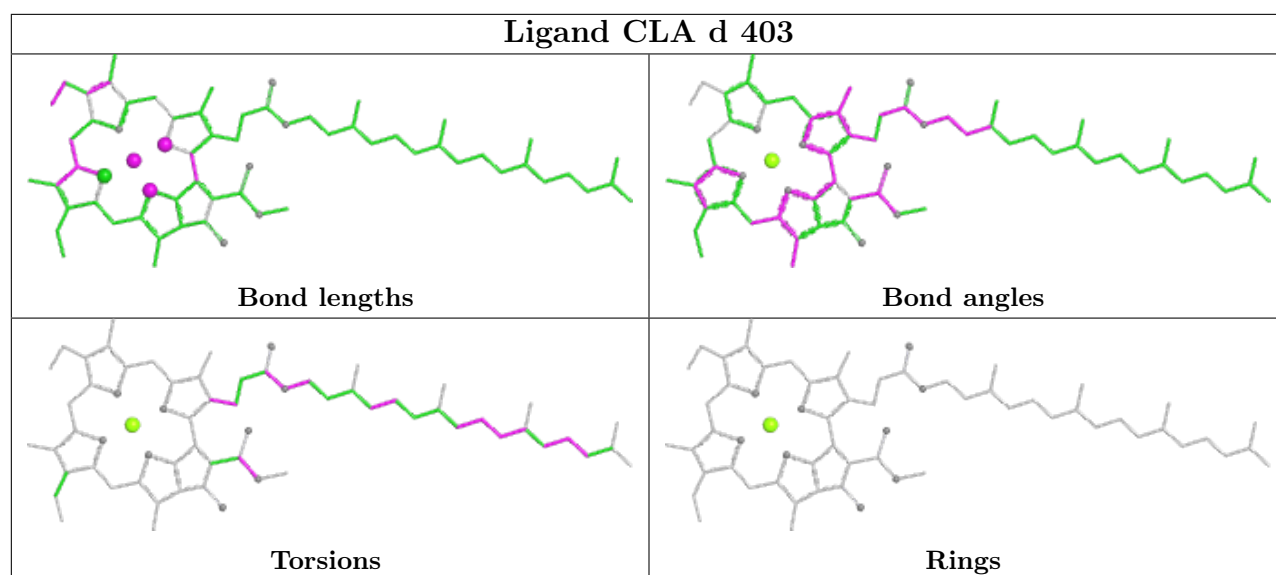


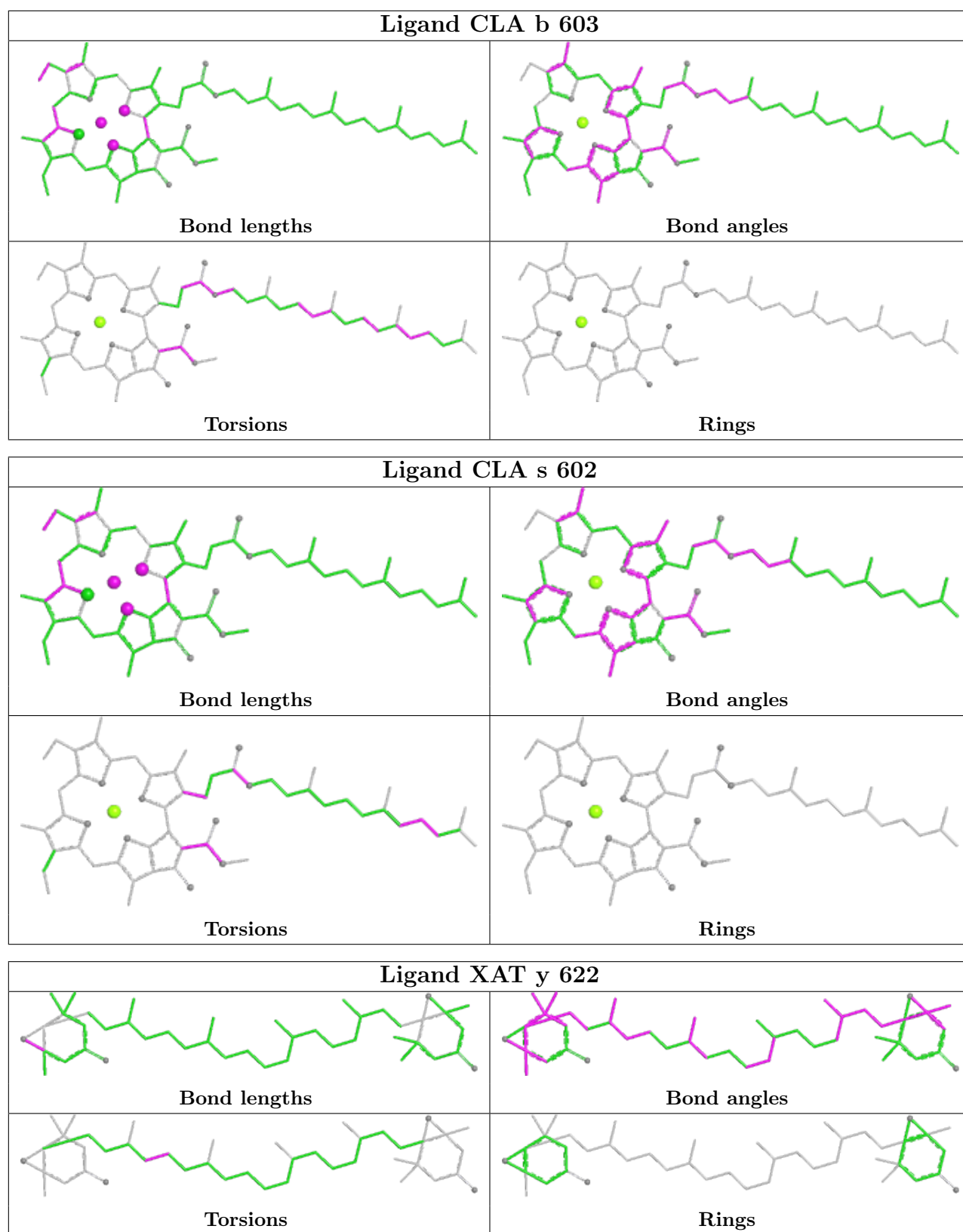


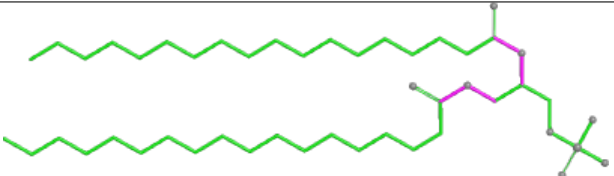
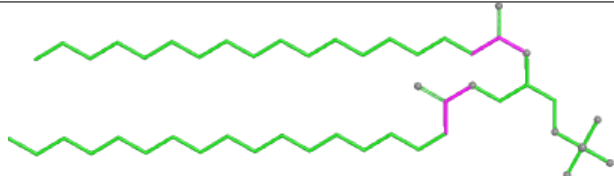
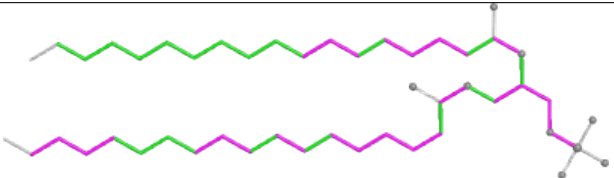
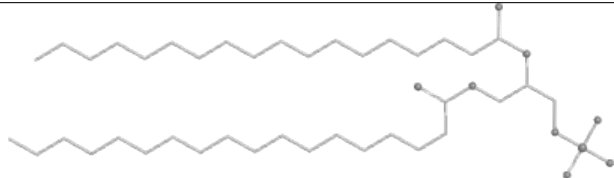
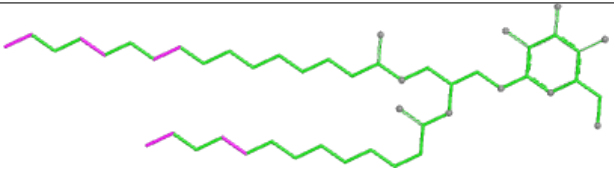
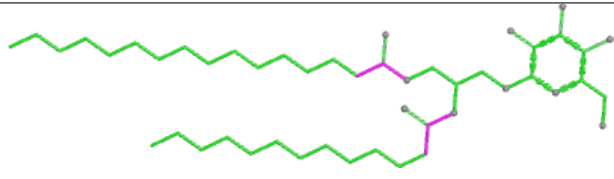
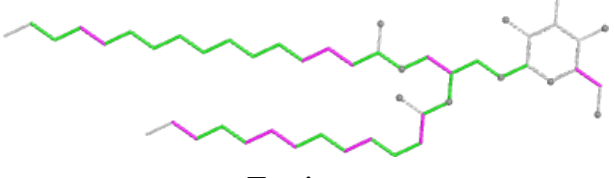
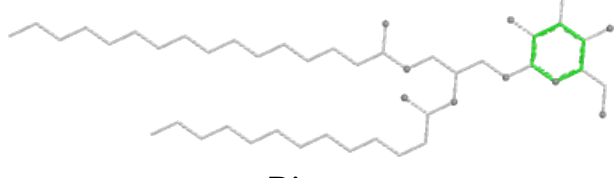
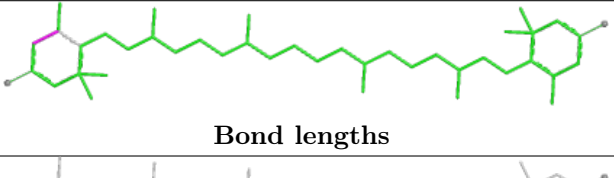
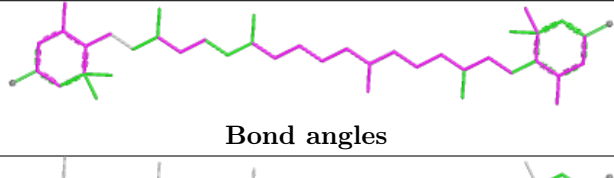




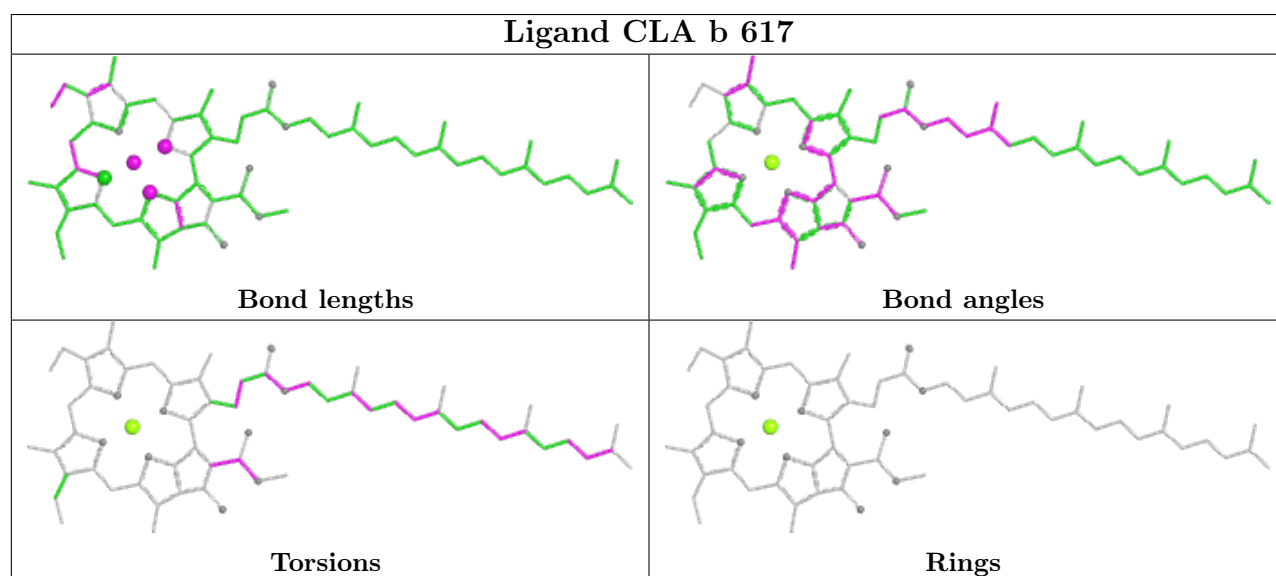
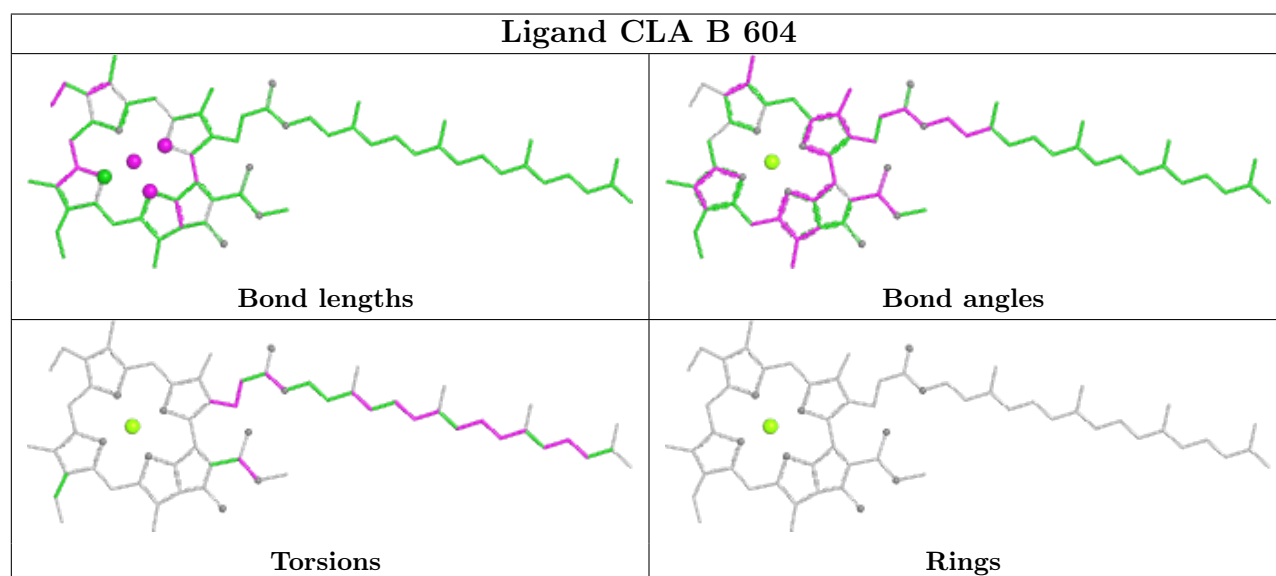
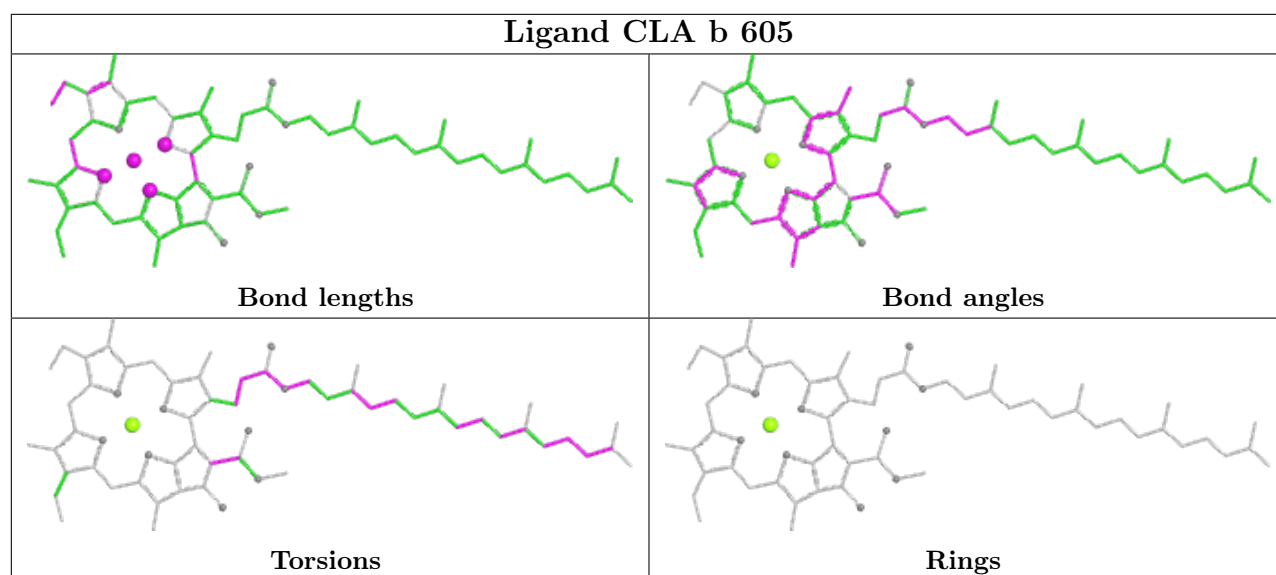


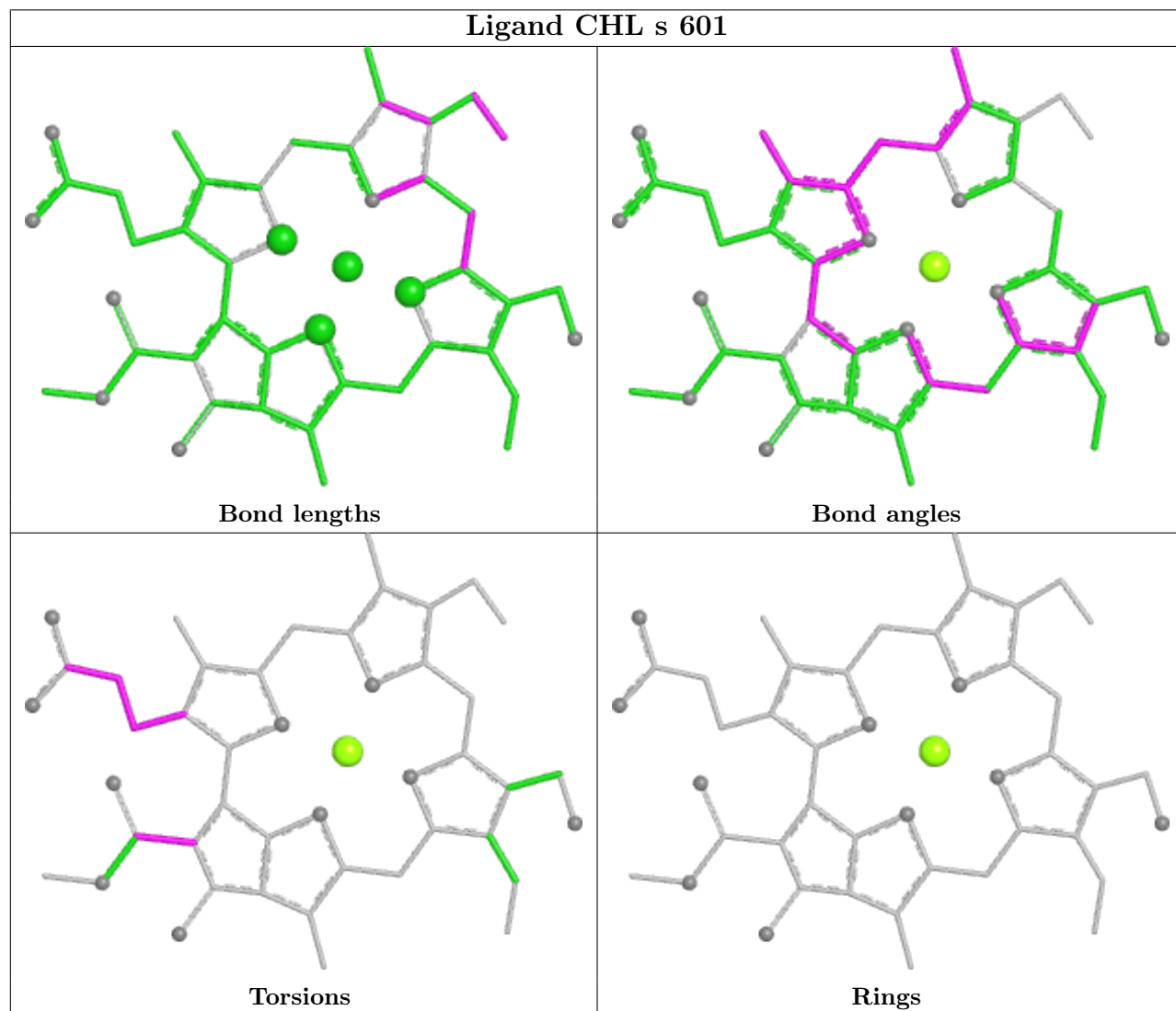
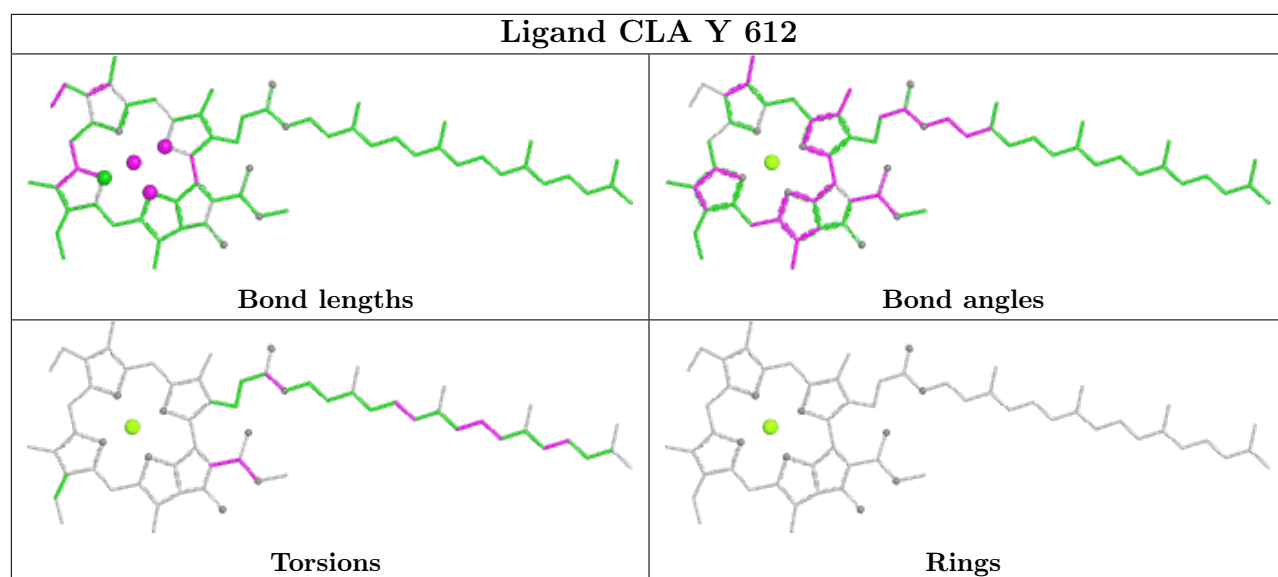


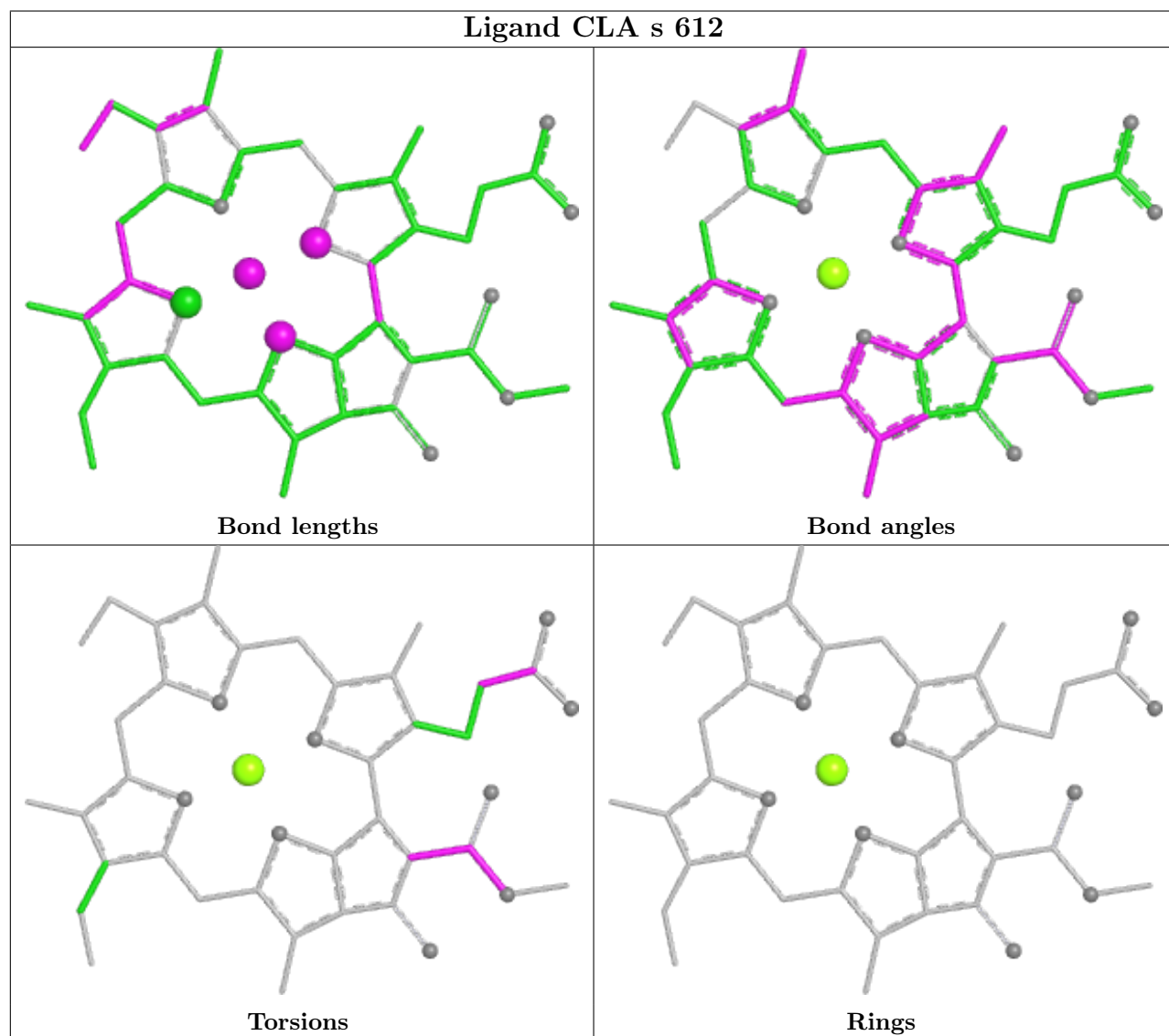
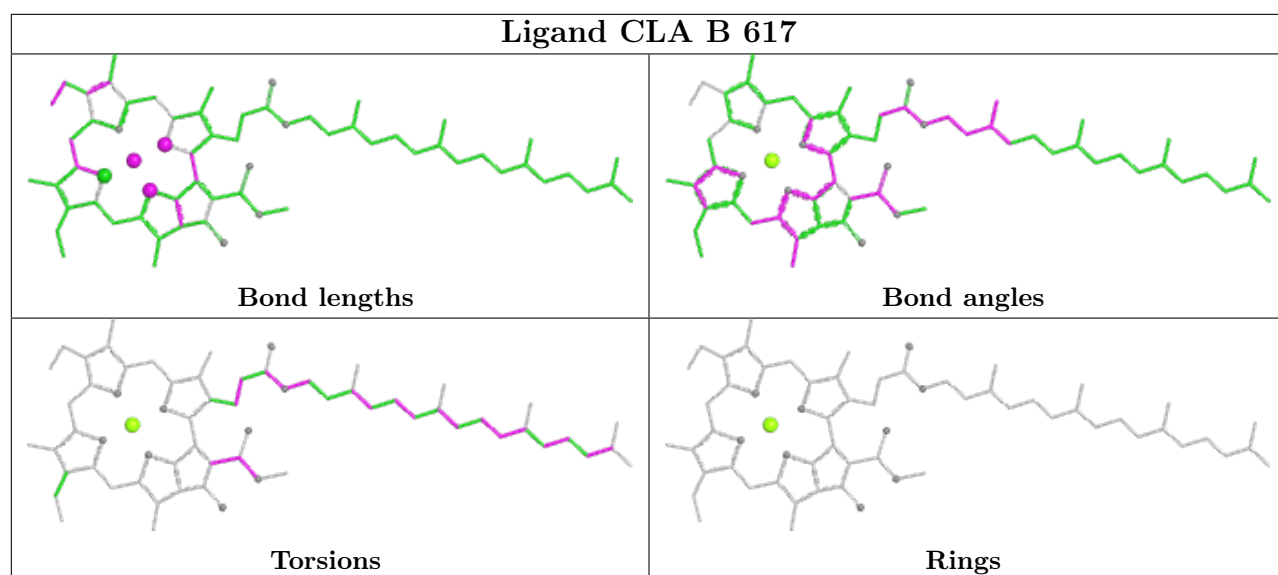


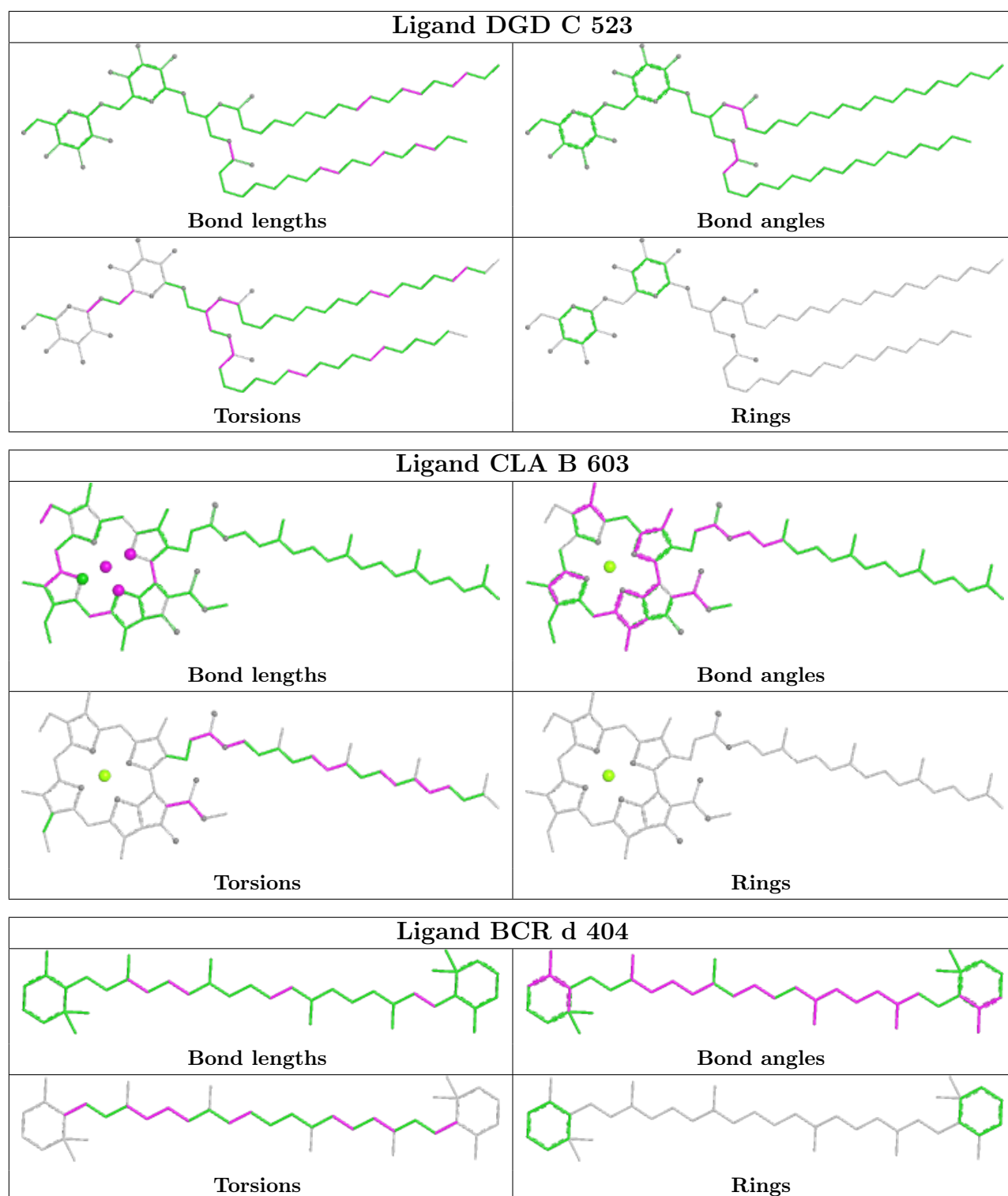


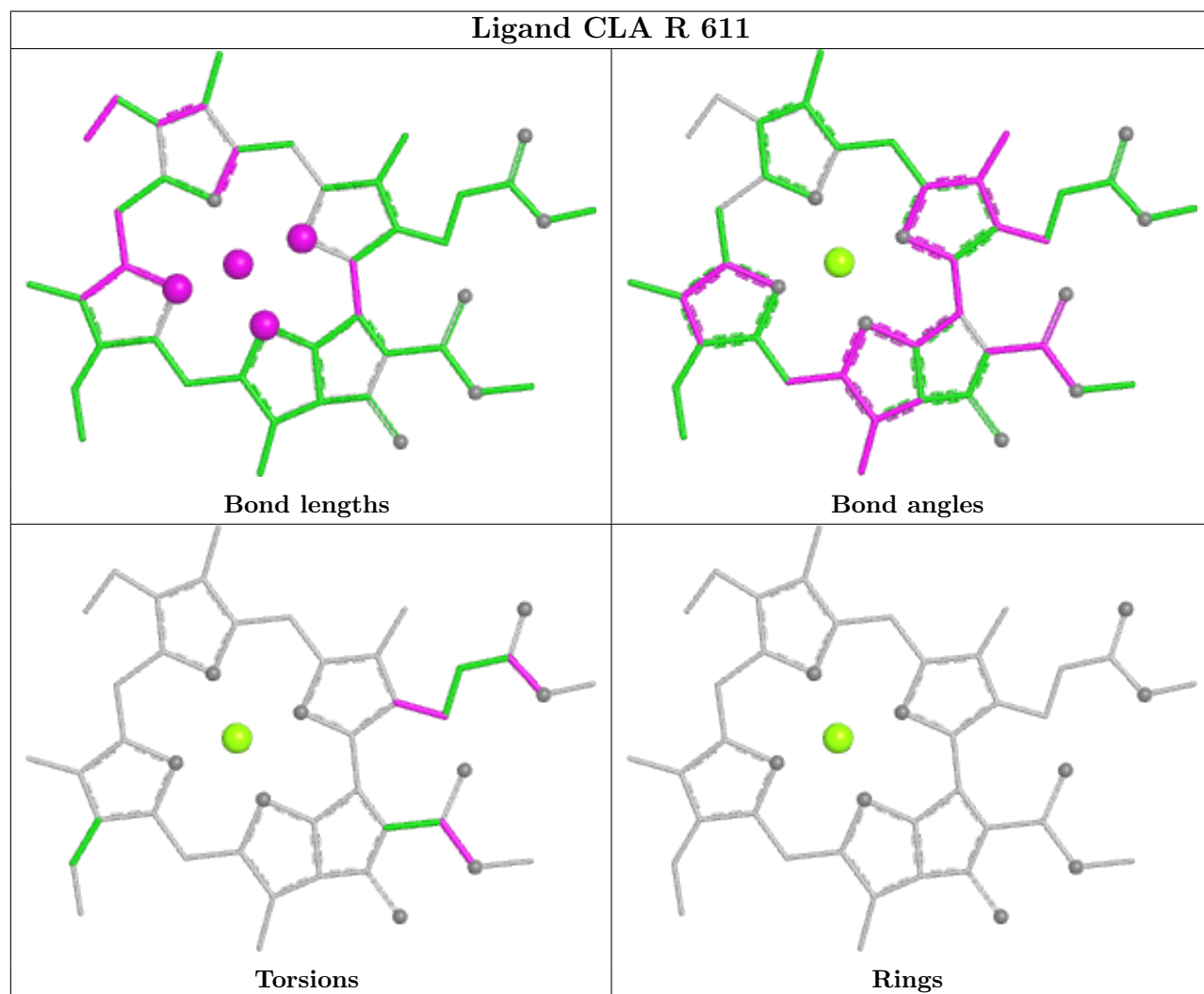
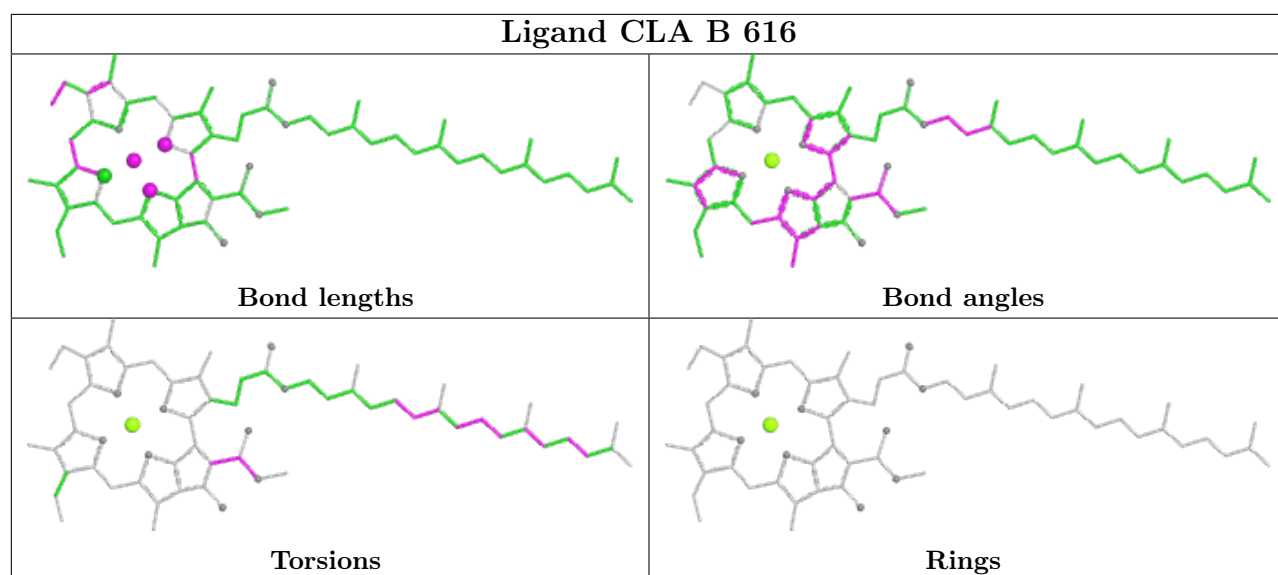
Ligand 3PH s 626	
 <p>Bond lengths</p>	 <p>Bond angles</p>
 <p>Torsions</p>	 <p>Rings</p>
Ligand LMG H 102	
 <p>Bond lengths</p>	 <p>Bond angles</p>
 <p>Torsions</p>	 <p>Rings</p>
Ligand LUT S 621	
 <p>Bond lengths</p>	 <p>Bond angles</p>
 <p>Torsions</p>	 <p>Rings</p>

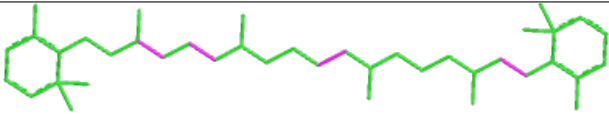
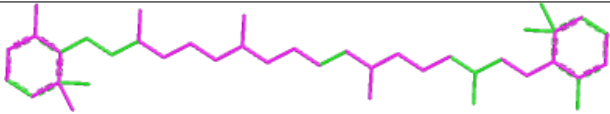
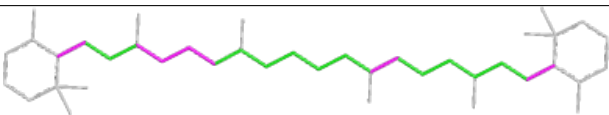
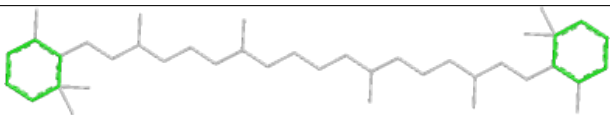


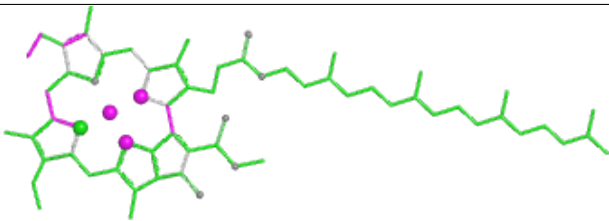
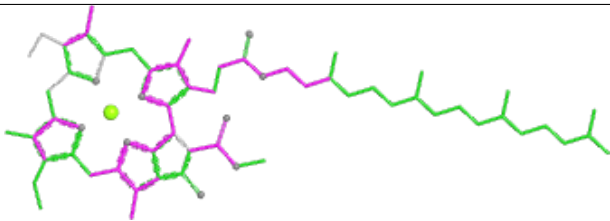
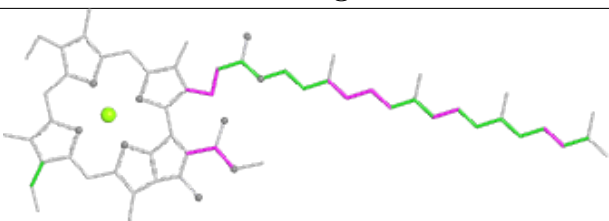
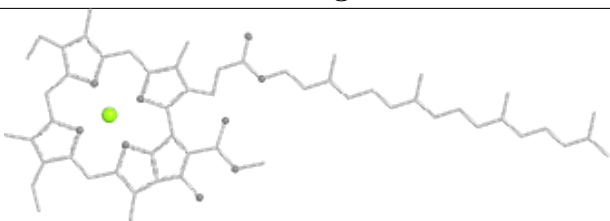


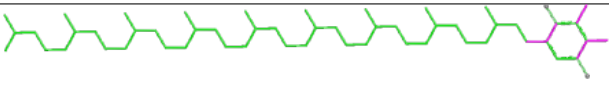
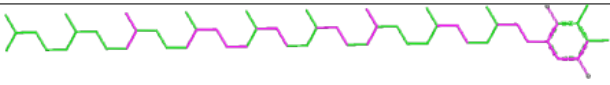
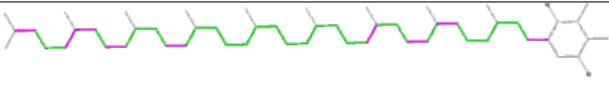
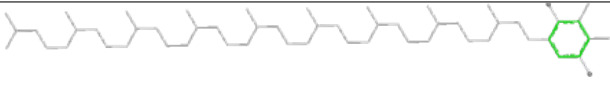






Ligand BCR c 517	
	Bond lengths
	Bond angles
	Torsions
	Rings

Ligand CLA g 610	
	Bond lengths
	Bond angles
	Torsions
	Rings

Ligand PL9 D 405	
	Bond lengths
	Bond angles
	Torsions
	Rings

5.7 Other polymers ⓘ

There are no such residues in this entry.

5.8 Polymer linkage issues ⓘ

The following chains have linkage breaks:

Mol	Chain	Number of breaks
22	R	1

Continued on next page...

Continued from previous page...

Mol	Chain	Number of breaks
22	r	1
23	s	1

All chain breaks are listed below:

Model	Chain	Residue-1	Atom-1	Residue-2	Atom-2	Distance (Å)
1	R	110:PRO	C	126:GLU	N	17.87
1	r	110:PRO	C	126:GLU	N	17.80
1	s	285:ARG	C	286:VAL	N	3.20

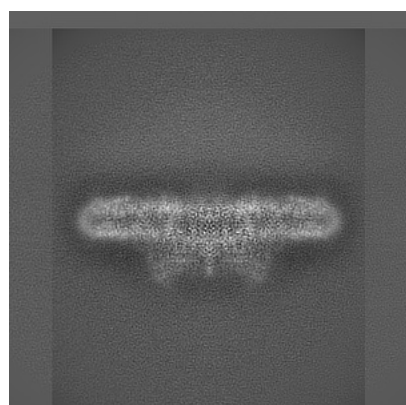
6 Map visualisation [i](#)

This section contains visualisations of the EMDB entry EMD-13430. These allow visual inspection of the internal detail of the map and identification of artifacts.

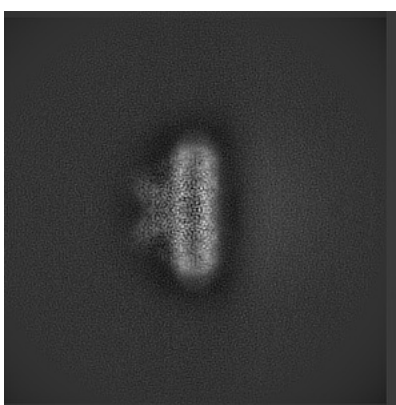
No raw map or half-maps were deposited for this entry and therefore no images, graphs, etc. pertaining to the raw map can be shown.

6.1 Orthogonal projections [i](#)

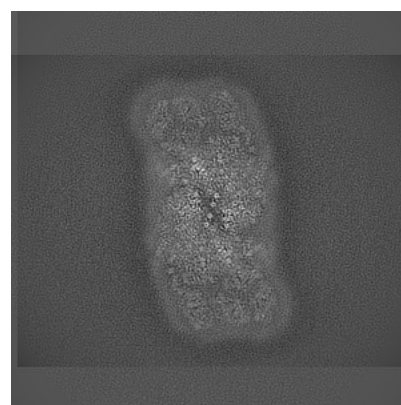
6.1.1 Primary map



X



Y

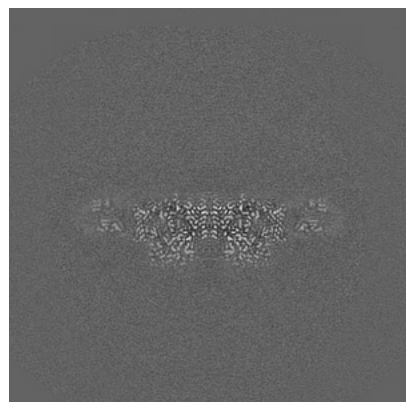


Z

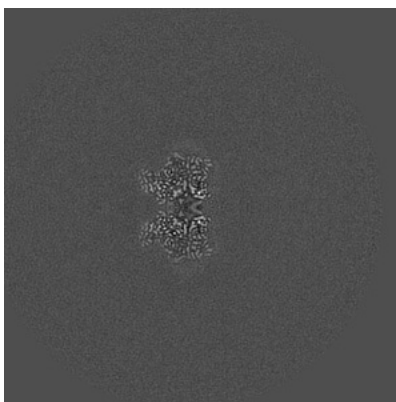
The images above show the map projected in three orthogonal directions.

6.2 Central slices [i](#)

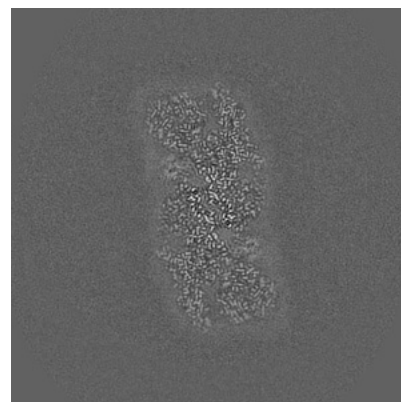
6.2.1 Primary map



X Index: 250



Y Index: 250

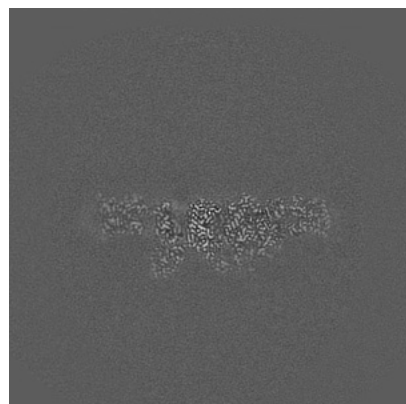


Z Index: 250

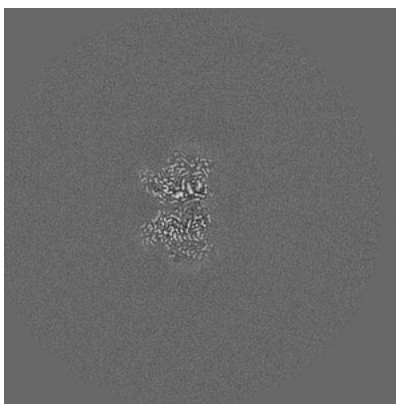
The images above show central slices of the map in three orthogonal directions.

6.3 Largest variance slices [i](#)

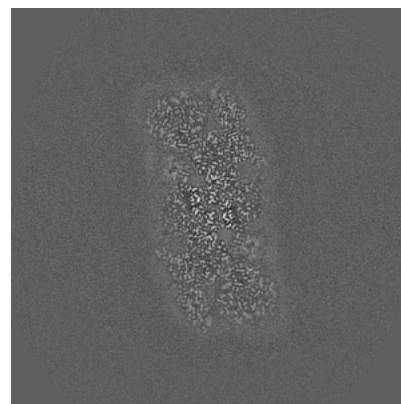
6.3.1 Primary map



X Index: 268



Y Index: 248

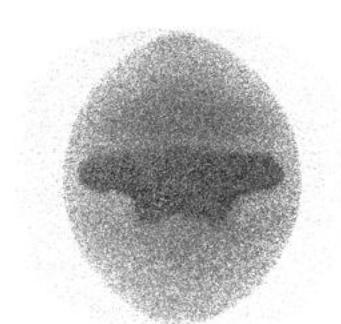


Z Index: 249

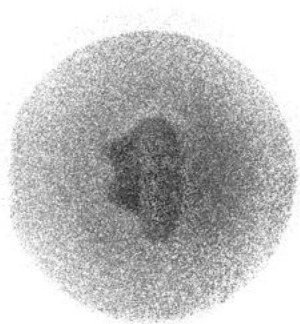
The images above show the largest variance slices of the map in three orthogonal directions.

6.4 Orthogonal surface views [i](#)

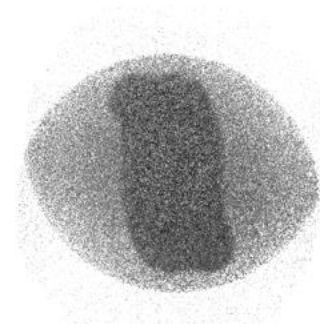
6.4.1 Primary map



X



Y



Z

The images above show the 3D surface view of the map at the recommended contour level 2.5. These images, in conjunction with the slice images, may facilitate assessment of whether an appropriate contour level has been provided.

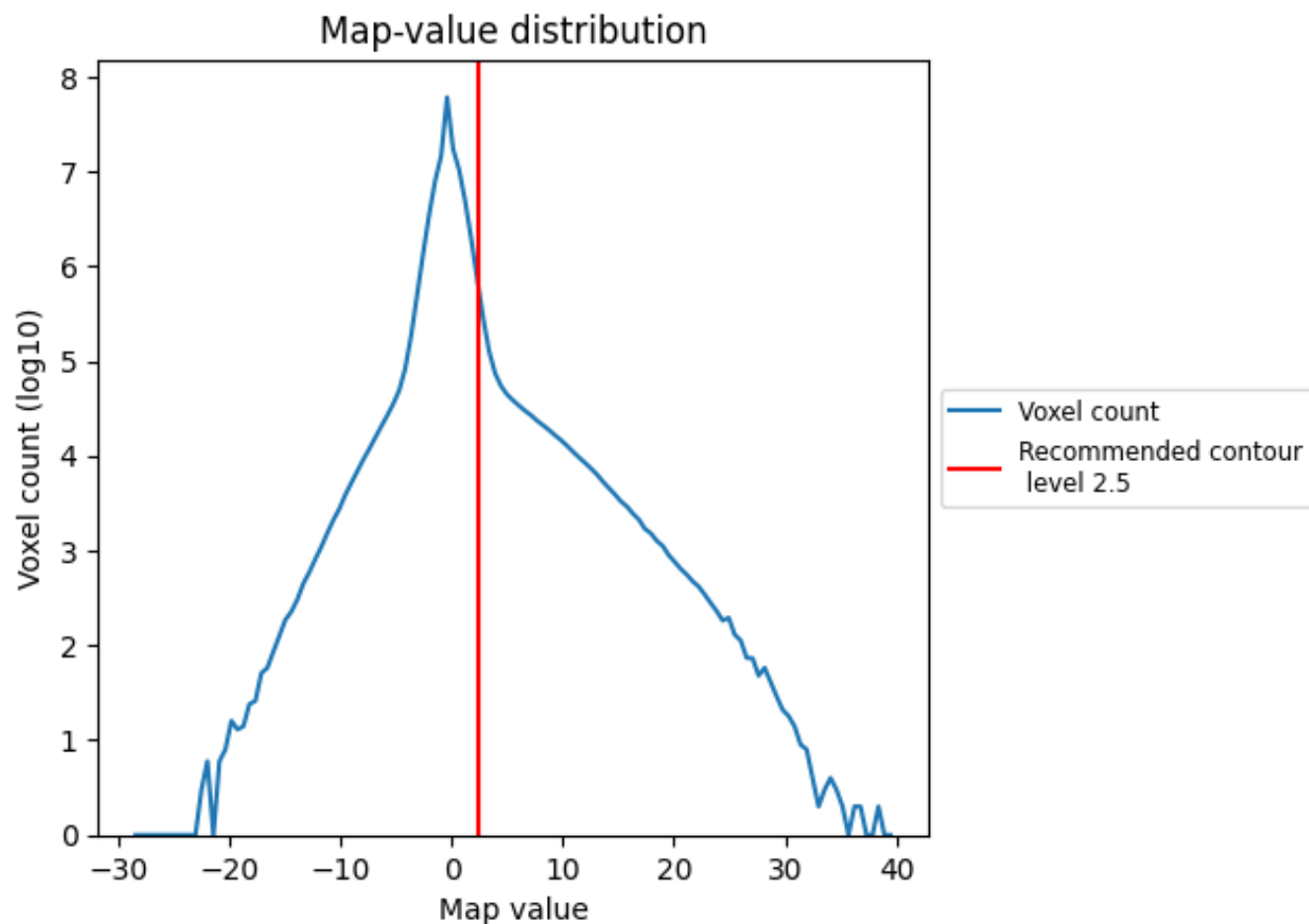
6.5 Mask visualisation

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

7 Map analysis [i](#)

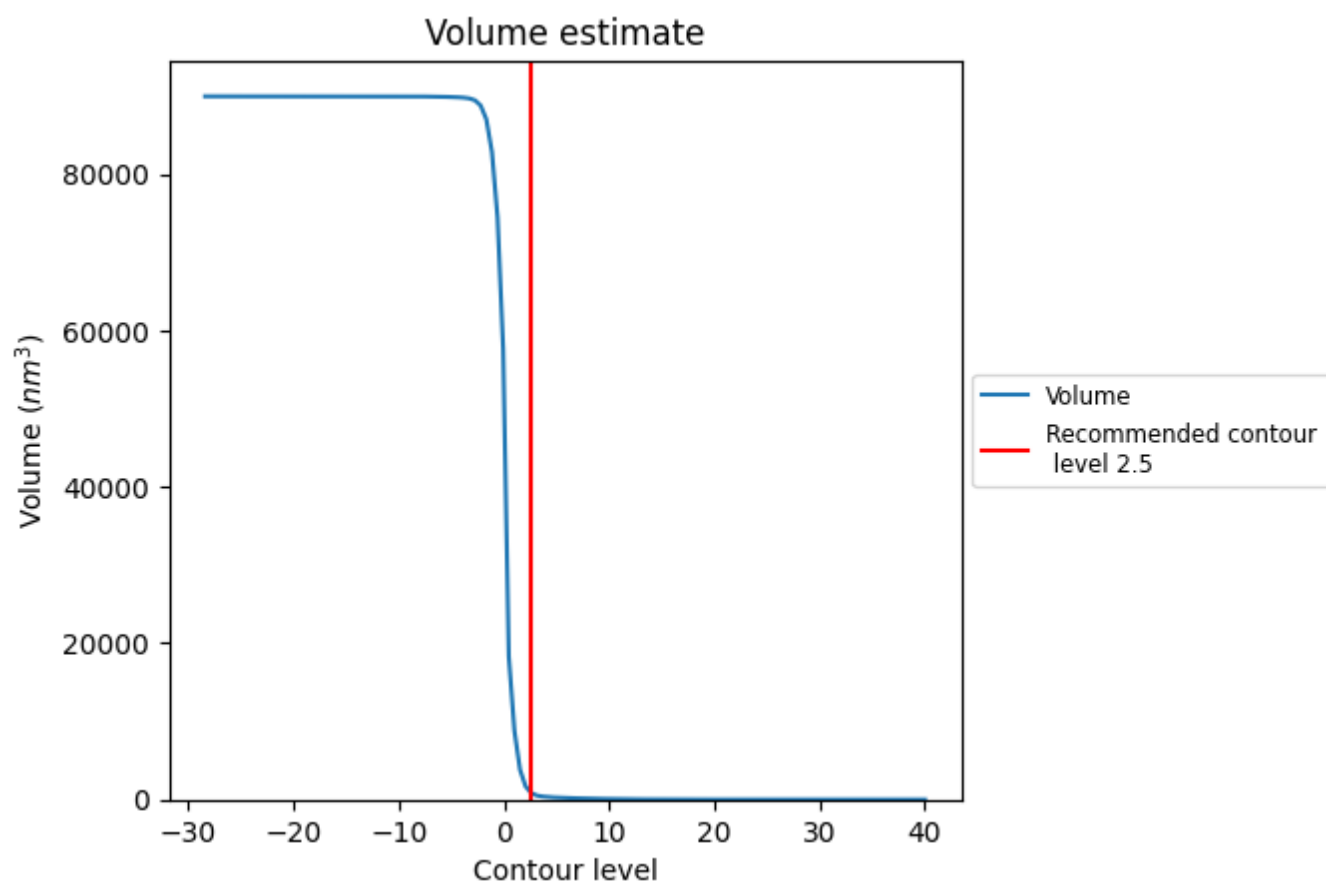
This section contains the results of statistical analysis of the map.

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.

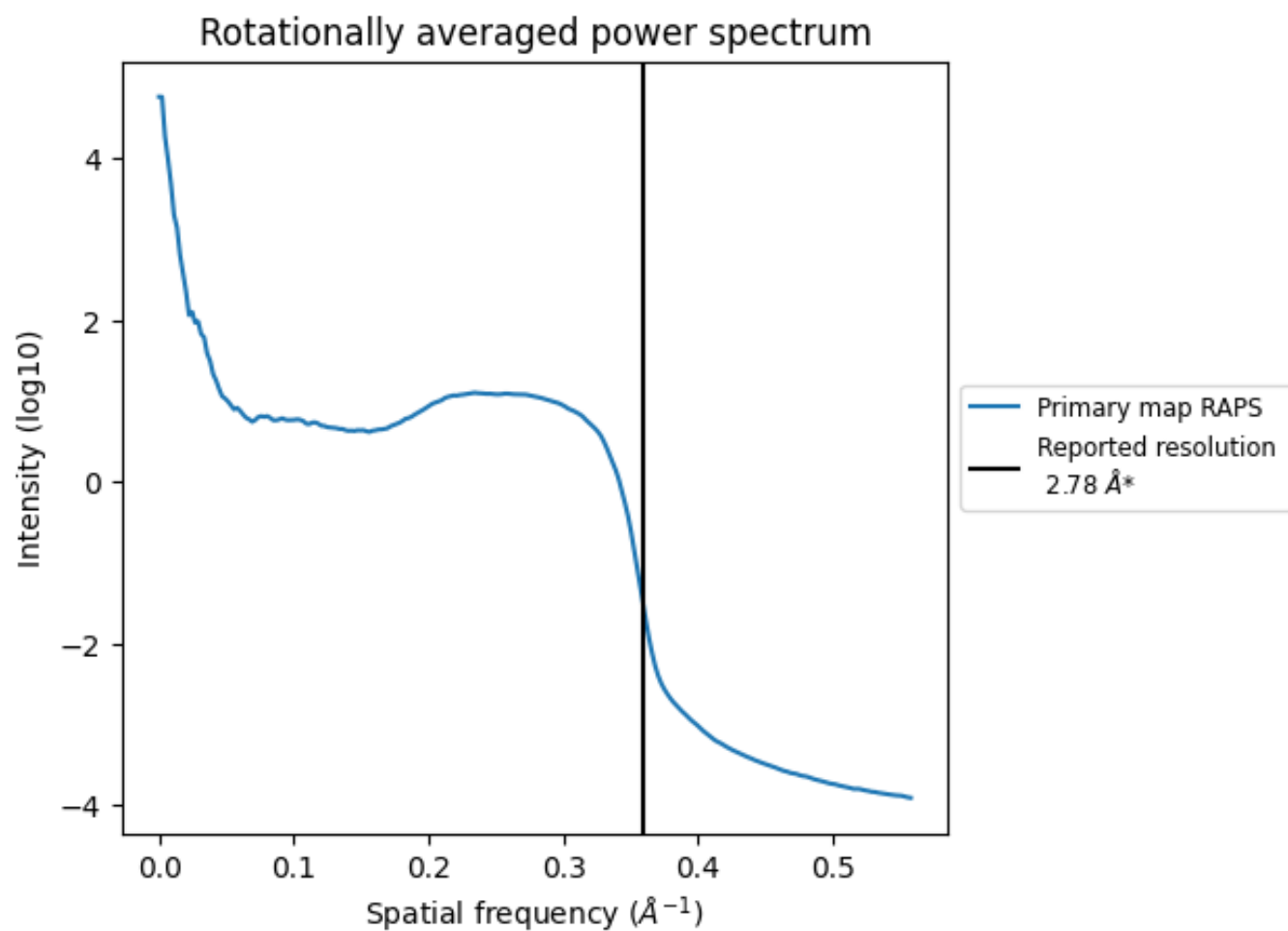
7.2 Volume estimate [i](#)



The volume at the recommended contour level is 980 nm³; this corresponds to an approximate mass of 885 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 ⓘ

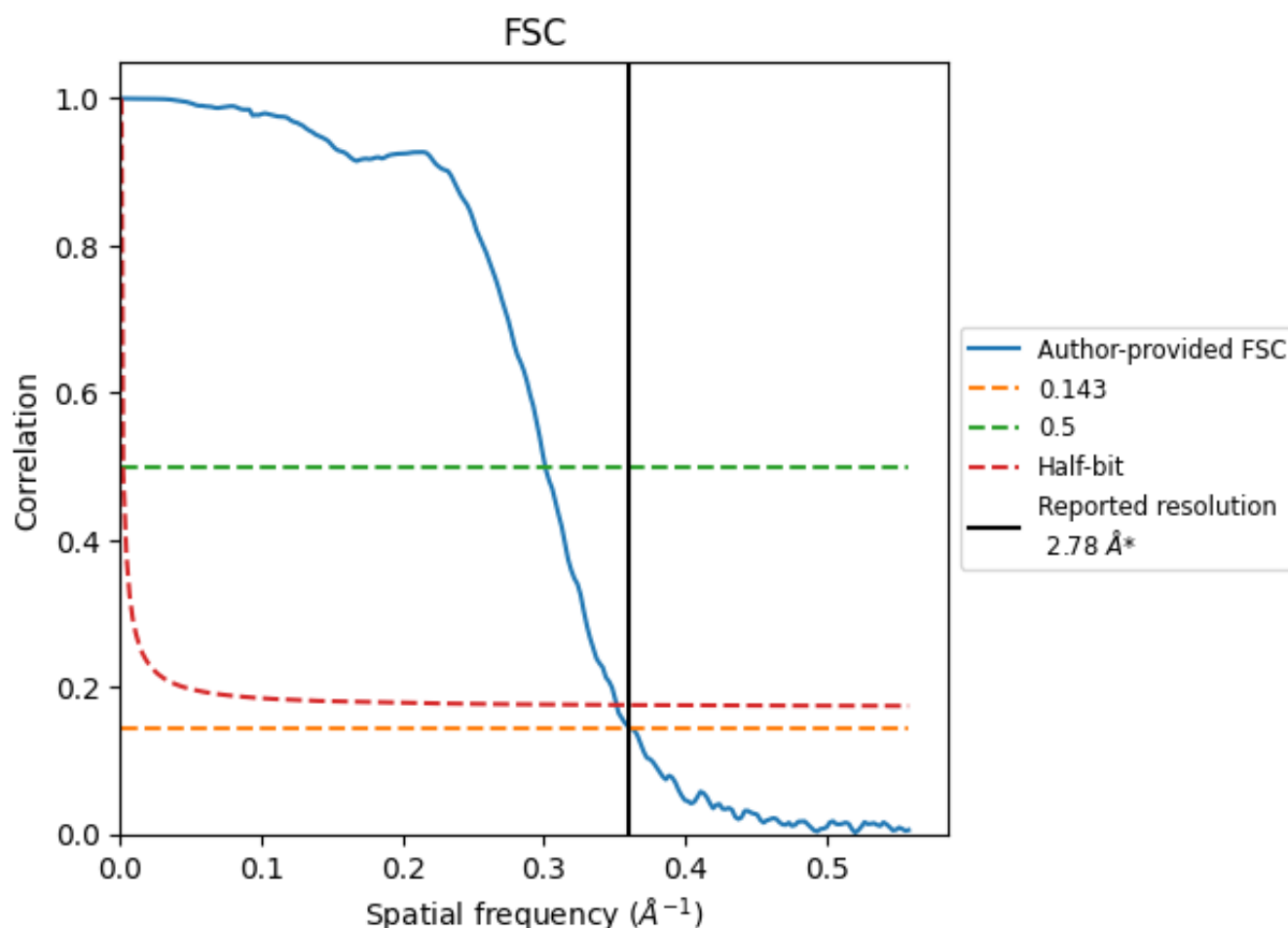


*Reported resolution corresponds to spatial frequency of 0.360 Å⁻¹

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 0.360 Å⁻¹

8.2 Resolution estimates [i](#)

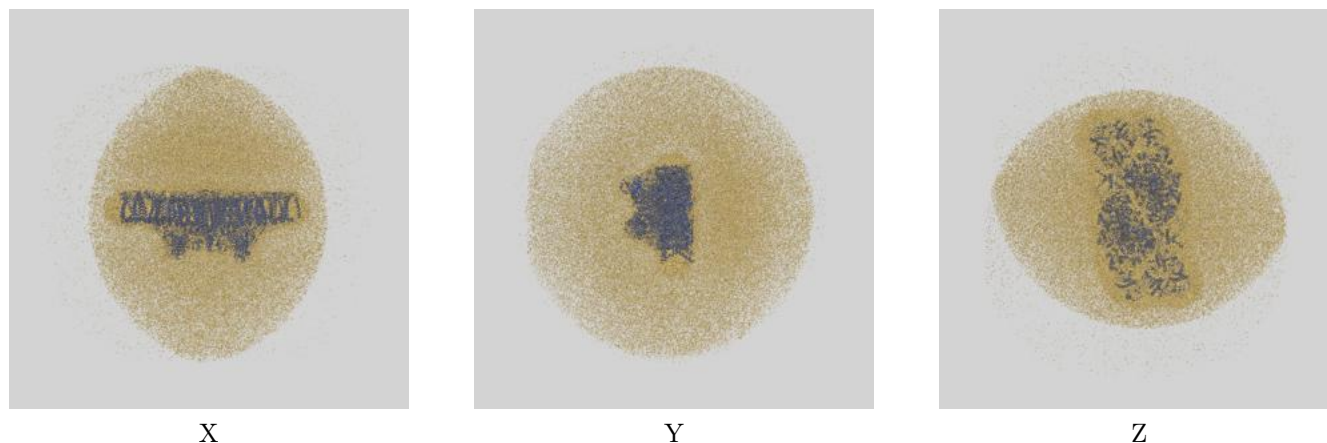
Resolution estimate (Å)	Estimation criterion (FSC cut-off)		
	0.143	0.5	Half-bit
Reported by author	2.78	-	-
Author-provided FSC curve	2.77	3.32	2.85
Unmasked-calculated*	-	-	-

*Resolution estimate based on FSC curve calculated by comparison of deposited half-maps.

9 Map-model fit [i](#)

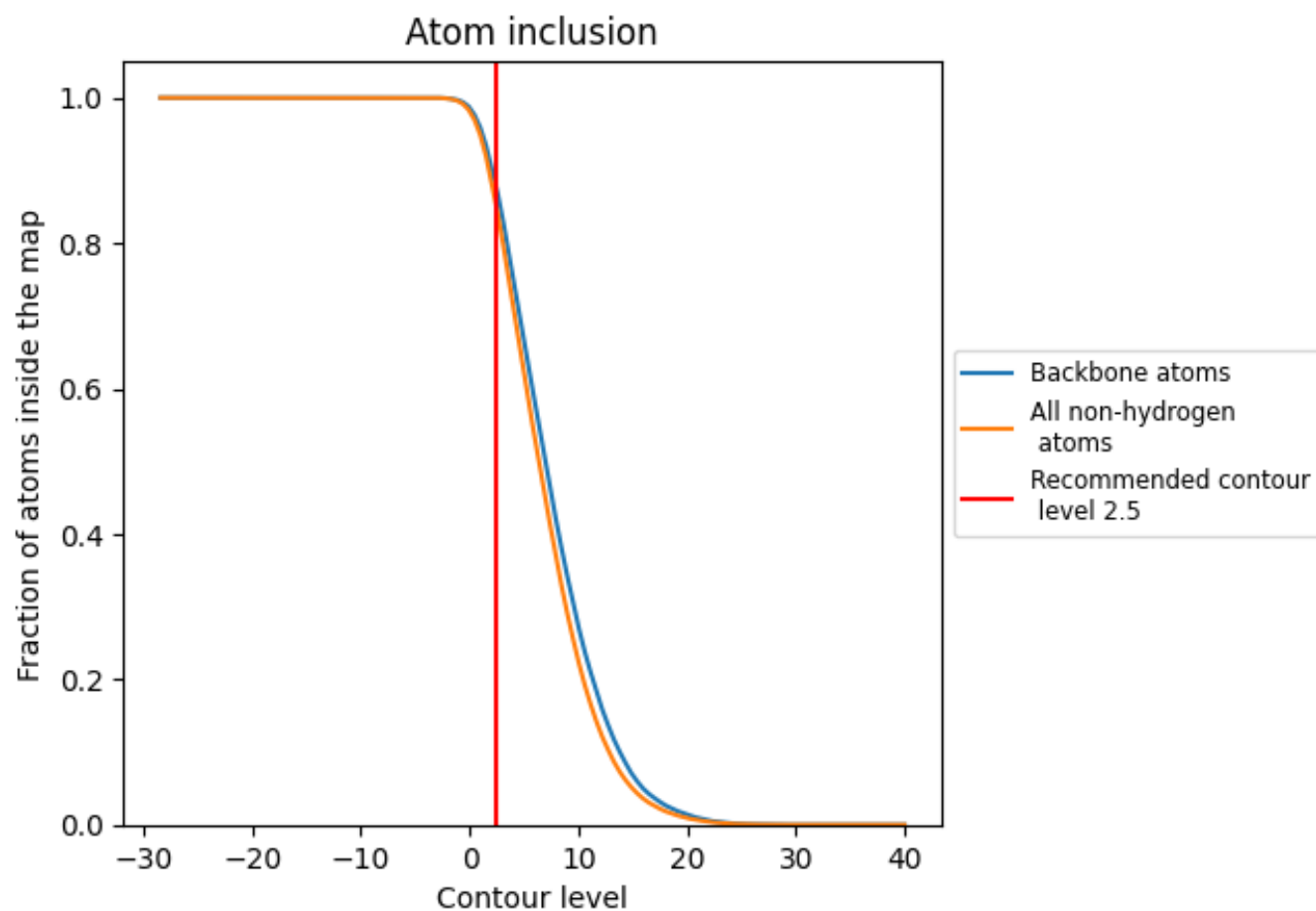
This section contains information regarding the fit between EMDB map EMD-13430 and PDB model 7PI5. Per-residue inclusion information can be found in section [3](#) on page [49](#).

9.1 Map-model overlay [i](#)



The images above show the 3D surface view of the map at the recommended contour level 2.5 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 Atom inclusion [i](#)



At the recommended contour level, 87% of all backbone atoms, 85% of all non-hydrogen atoms, are inside the map.