



Full wwPDB EM Validation Report ⓘ

Mar 9, 2026 – 05:52 PM UTC

PDB ID : 7SC9 / pdb_00007sc9
EMDB ID : EMD-25030
Title : Synechocystis PCC 6803 Phycobilisome core, complex with OCP
Authors : Sauer, P.V.; Sutter, M.; Dominguez-Martin, M.A.; Kirst, H.; Kerfeld, C.A.
Deposited on : 2021-09-27
Resolution : 2.60 Å(reported)

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

We welcome your comments at validation@mail.wwpdb.org

A user guide is available at

<https://www.wwpdb.org/validation/2017/EMValidationReportHelp>

with specific help available everywhere you see the ⓘ symbol.

The types of validation reports are described at

<http://www.wwpdb.org/validation/2017/FAQs#types>.

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

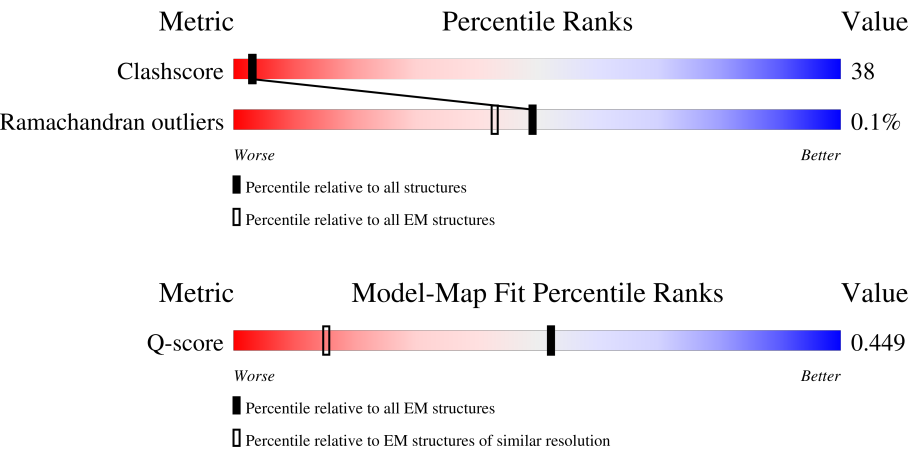
EMDB validation analysis : 0.0.1.dev132
Mogul : 2022.3.0, CSD as543be (2022)
MolProbity : 4-5-2 with Phenix2.0
Buster-report : wwPDB partial adaption of 1.1.7 (2018)
Percentile statistics : 20250101.v01 (using entries in the PDB archive January 1st 2025)
EM percentile statistics : 202505.v01 (Using data in the EMDB archive up until May 2025)
MapQ : 1.9.13
Ideal geometry (proteins) : Engh & Huber (2001)
Ideal geometry (DNA, RNA) : Parkinson et al. (1996)
Validation Pipeline (wwPDB-VP) : 2.49

1 Overall quality at a glance i

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

The reported resolution of this entry is 2.60 Å.

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



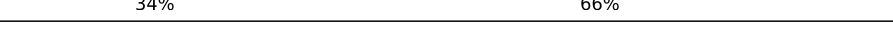
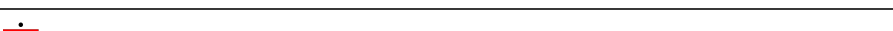
Metric	Whole archive (#Entries)	EM structures (#Entries)	Similar EM resolution (#Entries, resolution range(Å))
Clashscore	229148	23984	-
Ramachandran outliers	224038	23583	-
Q-score	-	25397	8728 (2.10 - 3.10)

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	AA	161	87% 41% 58% .
1	AC	161	87% 37% 63% .
1	AH	161	50% 50% .
1	AJ	161	49% 50% .
1	AN	161	43% 56% .

Continued on next page...

Continued from previous page...

Mol	Chain	Length	Quality of chain
1	AP	161	 53% 46%
1	AR	161	 42% 57%
1	AV	161	 34% 65%
1	AX	161	 47% 52%
1	AZ	161	 38% 61%
1	BI	161	 50% 50% 60%
1	BK	161	 39% 61% 61%
1	BP	161	 46% 53%
1	BR	161	 48% 51%
1	BV	161	 41% 58%
1	BX	161	 53% 47%
1	BZ	161	 46% 53%
1	CC	161	 39% 61%
1	CE	161	 34% 66%
1	CG	161	 48% 52%
1	CR	161	 37% 63% 6%
1	CT	161	 44% 55%
1	CV	161	 47% 52%
1	CY	161	 26% 73% 11%
1	DA	161	 48% 52%
1	DC	161	 52% 47%
1	DJ	161	 30% 70% 8%
1	DL	161	 37% 61%
1	DN	161	 49% 50%
1	DQ	161	 31% 68% 7%

Continued on next page...

Continued from previous page...

Mol	Chain	Length	Quality of chain
1	DS	161	
1	DU	161	
2	AB	161	
2	AD	161	
2	AF	161	
2	AI	161	
2	AL	161	
2	AO	161	
2	AQ	161	
2	AS	161	
2	AU	161	
2	AW	161	
2	AY	161	
2	BJ	161	
2	BL	161	
2	BN	161	
2	BQ	161	
2	BT	161	
2	BW	161	
2	BY	161	
2	CA	161	
2	CD	161	
2	CF	161	
2	CH	161	
2	CS	161	

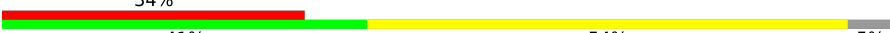
Continued on next page...

Continued from previous page...

Mol	Chain	Length	Quality of chain
2	CU	161	
2	CW	161	
2	CZ	161	
2	DB	161	
2	DD	161	
2	DK	161	
2	DM	161	
2	DO	161	
2	DR	161	
2	DT	161	
2	DV	161	
3	AE	161	
3	BM	161	
4	AK	169	
4	BS	169	
5	BA	67	
5	BB	67	
5	CJ	67	
5	CK	67	
5	DF	67	
5	DX	67	
6	BD	896	
6	CM	896	
7	BF	121	
7	CO	121	

Continued on next page...

Continued from previous page...

Mol	Chain	Length	Quality of chain
8	BG	249	 11% 12% 77%
8	CP	249	 12% 10% 77%
8	DG	249	 8% 13% 9% 78%
8	DH	249	 13% 8% 78%
8	DY	249	 9% 10% 12% 78%
8	DZ	249	 12% 10% 78%
9	BH	317	 34% 41% 54% 5%
9	CQ	317	 29% 40% 54% 5%
9	DI	317	 34% 28% 65% 7%
9	EA	317	 29% 31% 63% 7%

2 Entry composition [i](#)

There are 11 unique types of molecules in this entry. The entry contains 117470 atoms, of which 208 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 Allophycocyanin alpha chain.

Mol	Chain	Residues	Atoms					AltConf	Trace
1	AA	160	Total	C	N	O	S	0	0
			1210	754	207	242	7		
1	AC	160	Total	C	N	O	S	0	0
			1210	754	207	242	7		
1	AH	160	Total	C	N	O	S	0	0
			1210	754	207	242	7		
1	AJ	160	Total	C	N	O	S	0	0
			1210	754	207	242	7		
1	AN	160	Total	C	N	O	S	0	0
			1210	754	207	242	7		
1	AP	160	Total	C	N	O	S	0	0
			1210	754	207	242	7		
1	AR	160	Total	C	N	O	S	0	0
			1210	754	207	242	7		
1	AV	160	Total	C	N	O	S	0	0
			1210	754	207	242	7		
1	AX	160	Total	C	N	O	S	0	0
			1210	754	207	242	7		
1	AZ	160	Total	C	N	O	S	0	0
			1210	754	207	242	7		
1	BI	160	Total	C	N	O	S	0	0
			1210	754	207	242	7		
1	BK	160	Total	C	N	O	S	0	0
			1210	754	207	242	7		
1	BP	160	Total	C	N	O	S	0	0
			1210	754	207	242	7		
1	BR	160	Total	C	N	O	S	0	0
			1210	754	207	242	7		
1	BV	160	Total	C	N	O	S	0	0
			1210	754	207	242	7		
1	BX	160	Total	C	N	O	S	0	0
			1210	754	207	242	7		
1	BZ	160	Total	C	N	O	S	0	0
			1210	754	207	242	7		

Continued on next page...

Continued from previous page...

Mol	Chain	Residues	Atoms					AltConf	Trace
1	CC	160	Total	C	N	O	S	0	0
			1210	754	207	242	7		
1	CE	160	Total	C	N	O	S	0	0
			1210	754	207	242	7		
1	CG	160	Total	C	N	O	S	0	0
			1210	754	207	242	7		
1	CR	160	Total	C	N	O	S	0	0
			1210	754	207	242	7		
1	CT	160	Total	C	N	O	S	0	0
			1210	754	207	242	7		
1	CV	160	Total	C	N	O	S	0	0
			1210	754	207	242	7		
1	CY	160	Total	C	N	O	S	0	0
			1210	754	207	242	7		
1	DA	160	Total	C	N	O	S	0	0
			1210	754	207	242	7		
1	DC	160	Total	C	N	O	S	0	0
			1210	754	207	242	7		
1	DJ	160	Total	C	N	O	S	0	0
			1210	754	207	242	7		
1	DL	160	Total	C	N	O	S	0	0
			1210	754	207	242	7		
1	DN	160	Total	C	N	O	S	0	0
			1210	754	207	242	7		
1	DQ	160	Total	C	N	O	S	0	0
			1210	754	207	242	7		
1	DS	160	Total	C	N	O	S	0	0
			1210	754	207	242	7		
1	DU	160	Total	C	N	O	S	0	0
			1210	754	207	242	7		

- Molecule 2 is a protein called Allophycocyanin beta chain.

Mol	Chain	Residues	Atoms					AltConf	Trace
2	AB	161	Total	C	N	O	S	0	0
			1206	757	202	240	7		
2	AD	161	Total	C	N	O	S	0	0
			1206	757	202	240	7		
2	AF	161	Total	C	N	O	S	0	0
			1206	757	202	240	7		
2	AI	161	Total	C	N	O	S	0	0
			1206	757	202	240	7		

Continued on next page...

Continued from previous page...

Mol	Chain	Residues	Atoms					AltConf	Trace
2	AL	161	Total	C	N	O	S	0	0
			1206	757	202	240	7		
2	AO	161	Total	C	N	O	S	0	0
			1206	757	202	240	7		
2	AQ	161	Total	C	N	O	S	0	0
			1206	757	202	240	7		
2	AS	161	Total	C	N	O	S	0	0
			1206	757	202	240	7		
2	AU	161	Total	C	N	O	S	0	0
			1206	757	202	240	7		
2	AW	161	Total	C	N	O	S	0	0
			1206	757	202	240	7		
2	AY	161	Total	C	N	O	S	0	0
			1206	757	202	240	7		
2	BJ	161	Total	C	N	O	S	0	0
			1206	757	202	240	7		
2	BL	161	Total	C	N	O	S	0	0
			1206	757	202	240	7		
2	BN	161	Total	C	N	O	S	0	0
			1206	757	202	240	7		
2	BQ	161	Total	C	N	O	S	0	0
			1206	757	202	240	7		
2	BT	161	Total	C	N	O	S	0	0
			1206	757	202	240	7		
2	BW	161	Total	C	N	O	S	0	0
			1206	757	202	240	7		
2	BY	161	Total	C	N	O	S	0	0
			1206	757	202	240	7		
2	CA	161	Total	C	N	O	S	0	0
			1206	757	202	240	7		
2	CD	161	Total	C	N	O	S	0	0
			1206	757	202	240	7		
2	CF	161	Total	C	N	O	S	0	0
			1206	757	202	240	7		
2	CH	161	Total	C	N	O	S	0	0
			1206	757	202	240	7		
2	CS	161	Total	C	N	O	S	0	0
			1206	757	202	240	7		
2	CU	161	Total	C	N	O	S	0	0
			1206	757	202	240	7		
2	CW	161	Total	C	N	O	S	0	0
			1206	757	202	240	7		

Continued on next page...

Continued from previous page...

Mol	Chain	Residues	Atoms					AltConf	Trace
2	CZ	161	Total	C	N	O	S	0	0
			1206	757	202	240	7		
2	DB	161	Total	C	N	O	S	0	0
			1206	757	202	240	7		
2	DD	161	Total	C	N	O	S	0	0
			1206	757	202	240	7		
2	DK	161	Total	C	N	O	S	0	0
			1206	757	202	240	7		
2	DM	161	Total	C	N	O	S	0	0
			1206	757	202	240	7		
2	DO	161	Total	C	N	O	S	0	0
			1206	757	202	240	7		
2	DR	161	Total	C	N	O	S	0	0
			1206	757	202	240	7		
2	DT	161	Total	C	N	O	S	0	0
			1206	757	202	240	7		
2	DV	161	Total	C	N	O	S	0	0
			1206	757	202	240	7		

- Molecule 3 is a protein called Allophycocyanin subunit alpha-B.

Mol	Chain	Residues	Atoms					AltConf	Trace
3	AE	160	Total	C	N	O	S	0	0
			1254	797	212	241	4		
3	BM	160	Total	C	N	O	S	0	0
			1254	797	212	241	4		

- Molecule 4 is a protein called Allophycocyanin subunit beta-18.

Mol	Chain	Residues	Atoms					AltConf	Trace
4	AK	169	Total	C	N	O	S	0	0
			1322	825	229	259	9		
4	BS	169	Total	C	N	O	S	0	0
			1322	825	229	259	9		

- Molecule 5 is a protein called Phycobilisome 7.8 kDa linker polypeptide, allophycocyanin-associated, core.

Mol	Chain	Residues	Atoms					AltConf	Trace
5	BA	67	Total	C	N	O	S	0	0
			546	343	104	94	5		

Continued on next page...

Continued from previous page...

Mol	Chain	Residues	Atoms					AltConf	Trace
5	BB	67	Total	C	N	O	S	0	0
			546	343	104	94	5		
5	CJ	67	Total	C	N	O	S	0	0
			546	343	104	94	5		
5	CK	67	Total	C	N	O	S	0	0
			546	343	104	94	5		
5	DF	67	Total	C	N	O	S	0	0
			546	343	104	94	5		
5	DX	67	Total	C	N	O	S	0	0
			546	343	104	94	5		

There are 6 discrepancies between the modelled and reference sequences:

Chain	Residue	Modelled	Actual	Comment	Reference
BA	36	TRP	SER	conflict	UNP Q01950
BB	36	TRP	SER	conflict	UNP Q01950
CJ	36	TRP	SER	conflict	UNP Q01950
CK	36	TRP	SER	conflict	UNP Q01950
DF	36	TRP	SER	conflict	UNP Q01950
DX	36	TRP	SER	conflict	UNP Q01950

- Molecule 6 is a protein called Phycobiliprotein ApcE.

Mol	Chain	Residues	Atoms					AltConf	Trace
6	BD	850	Total	C	N	O	S	0	0
			6761	4299	1183	1262	17		
6	CM	850	Total	C	N	O	S	0	0
			6761	4299	1183	1262	17		

- Molecule 7 is a protein called Sll1873 protein.

Mol	Chain	Residues	Atoms					AltConf	Trace
7	BF	36	Total	C	N	O	S	0	0
			277	172	55	49	1		
7	CO	36	Total	C	N	O	S	0	0
			277	172	55	49	1		

- Molecule 8 is a protein called Phycobilisome rod-core linker polypeptide CpcG.

Mol	Chain	Residues	Atoms					AltConf	Trace
8	BG	57	Total	C	N	O	S	0	0
			451	282	87	80	2		

Continued on next page...

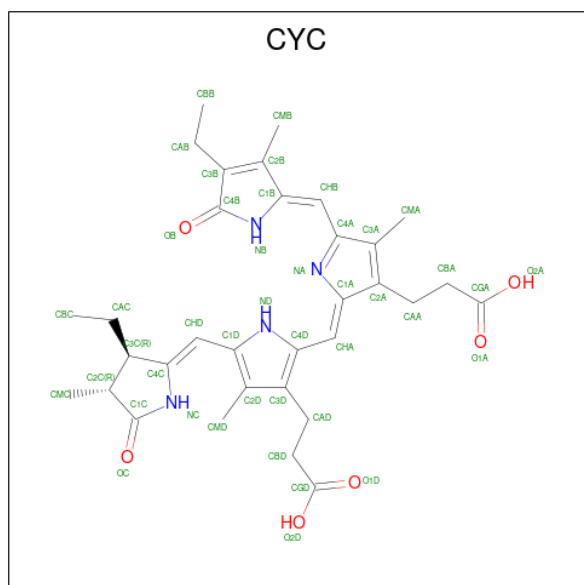
Continued from previous page...

Mol	Chain	Residues	Atoms					AltConf	Trace
8	CP	57	Total	C	N	O	S	0	0
			451	282	87	80	2		
8	DG	54	Total	C	N	O	S	0	0
			426	266	83	75	2		
8	DH	54	Total	C	N	O	S	0	0
			426	266	83	75	2		
8	DY	54	Total	C	N	O	S	0	0
			426	266	83	75	2		
8	DZ	54	Total	C	N	O	S	0	0
			426	266	83	75	2		

- Molecule 9 is a protein called Orange carotenoid-binding protein.

Mol	Chain	Residues	Atoms					AltConf	Trace
9	BH	300	Total	C	N	O	S	0	0
			2302	1474	388	429	11		
9	CQ	300	Total	C	N	O	S	0	0
			2302	1474	388	429	11		
9	DI	296	Total	C	N	O	S	0	0
			2280	1459	387	423	11		
9	EA	296	Total	C	N	O	S	0	0
			2280	1459	387	423	11		

- Molecule 10 is PHYCOCYANOBILIN (CCD ID: CYC) (formula: $C_{33}H_{40}N_4O_6$) (labeled as "Ligand of Interest" by depositor).



Mol	Chain	Residues	Atoms				AltConf
10	AA	1	Total	C	N	O	0
			43	33	4	6	
10	AB	1	Total	C	N	O	0
			43	33	4	6	
10	AC	1	Total	C	N	O	0
			43	33	4	6	
10	AD	1	Total	C	N	O	0
			43	33	4	6	
10	AE	1	Total	C	N	O	0
			43	33	4	6	
10	AF	1	Total	C	N	O	0
			43	33	4	6	
10	AH	1	Total	C	N	O	0
			43	33	4	6	
10	AI	1	Total	C	N	O	0
			43	33	4	6	
10	AJ	1	Total	C	N	O	0
			43	33	4	6	
10	AK	1	Total	C	N	O	0
			43	33	4	6	
10	AL	1	Total	C	N	O	0
			43	33	4	6	
10	AN	1	Total	C	N	O	0
			43	33	4	6	
10	AO	1	Total	C	N	O	0
			43	33	4	6	
10	AP	1	Total	C	N	O	0
			43	33	4	6	
10	AQ	1	Total	C	N	O	0
			43	33	4	6	
10	AR	1	Total	C	N	O	0
			43	33	4	6	
10	AU	1	Total	C	N	O	0
			43	33	4	6	
10	AV	1	Total	C	N	O	0
			43	33	4	6	
10	AW	1	Total	C	N	O	0
			43	33	4	6	
10	AX	1	Total	C	N	O	0
			43	33	4	6	
10	AY	1	Total	C	N	O	0
			43	33	4	6	
10	AZ	1	Total	C	N	O	0
			43	33	4	6	

Continued on next page...

Continued from previous page...

Mol	Chain	Residues	Atoms				AltConf
10	BD	1	Total 43	C 33	N 4	O 6	0
10	BD	1	Total 43	C 33	N 4	O 6	0
10	BI	1	Total 43	C 33	N 4	O 6	0
10	BJ	1	Total 43	C 33	N 4	O 6	0
10	BK	1	Total 43	C 33	N 4	O 6	0
10	BL	1	Total 43	C 33	N 4	O 6	0
10	BM	1	Total 43	C 33	N 4	O 6	0
10	BN	1	Total 43	C 33	N 4	O 6	0
10	BP	1	Total 43	C 33	N 4	O 6	0
10	BQ	1	Total 43	C 33	N 4	O 6	0
10	BR	1	Total 43	C 33	N 4	O 6	0
10	BS	1	Total 43	C 33	N 4	O 6	0
10	BT	1	Total 43	C 33	N 4	O 6	0
10	BV	1	Total 43	C 33	N 4	O 6	0
10	BW	1	Total 43	C 33	N 4	O 6	0
10	BX	1	Total 43	C 33	N 4	O 6	0
10	BY	1	Total 43	C 33	N 4	O 6	0
10	BZ	1	Total 43	C 33	N 4	O 6	0
10	CC	1	Total 43	C 33	N 4	O 6	0
10	CD	1	Total 43	C 33	N 4	O 6	0
10	CE	1	Total 43	C 33	N 4	O 6	0

Continued on next page...

Continued from previous page...

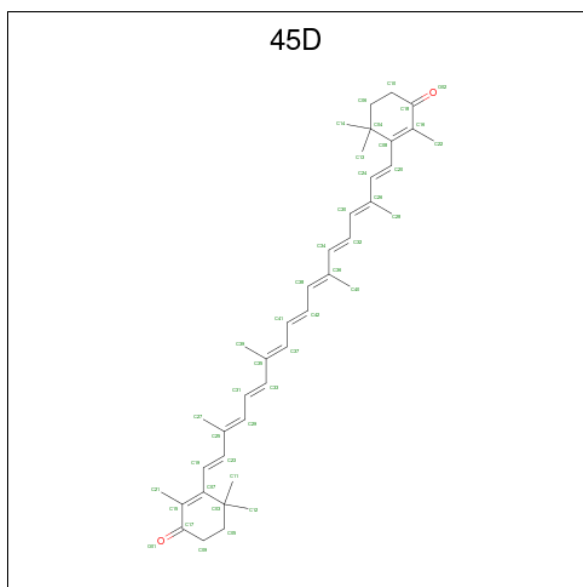
Mol	Chain	Residues	Atoms				AltConf
10	CF	1	Total 43	C 33	N 4	O 6	0
10	CG	1	Total 43	C 33	N 4	O 6	0
10	CH	1	Total 43	C 33	N 4	O 6	0
10	CM	1	Total 43	C 33	N 4	O 6	0
10	CM	1	Total 43	C 33	N 4	O 6	0
10	CR	1	Total 43	C 33	N 4	O 6	0
10	CS	1	Total 43	C 33	N 4	O 6	0
10	CT	1	Total 43	C 33	N 4	O 6	0
10	CU	1	Total 43	C 33	N 4	O 6	0
10	CV	1	Total 43	C 33	N 4	O 6	0
10	CW	1	Total 43	C 33	N 4	O 6	0
10	CY	1	Total 43	C 33	N 4	O 6	0
10	CZ	1	Total 43	C 33	N 4	O 6	0
10	DA	1	Total 43	C 33	N 4	O 6	0
10	DB	1	Total 43	C 33	N 4	O 6	0
10	DC	1	Total 43	C 33	N 4	O 6	0
10	DD	1	Total 43	C 33	N 4	O 6	0
10	DJ	1	Total 43	C 33	N 4	O 6	0
10	DK	1	Total 43	C 33	N 4	O 6	0
10	DL	1	Total 43	C 33	N 4	O 6	0
10	DM	1	Total 43	C 33	N 4	O 6	0

Continued on next page...

Continued from previous page...

Mol	Chain	Residues	Atoms				AltConf
10	DN	1	Total	C	N	O	0
			43	33	4	6	
10	DO	1	Total	C	N	O	0
			43	33	4	6	
10	DQ	1	Total	C	N	O	0
			43	33	4	6	
10	DR	1	Total	C	N	O	0
			43	33	4	6	
10	DS	1	Total	C	N	O	0
			43	33	4	6	
10	DT	1	Total	C	N	O	0
			43	33	4	6	
10	DU	1	Total	C	N	O	0
			43	33	4	6	
10	DV	1	Total	C	N	O	0
			43	33	4	6	

- Molecule 11 is beta,beta-carotene-4,4'-dione (CCD ID: 45D) (formula: $C_{40}H_{52}O_2$) (labeled as "Ligand of Interest" by depositor).



Mol	Chain	Residues	Atoms				AltConf
11	BH	1	Total	C	H	O	0
			94	40	52	2	
11	CQ	1	Total	C	H	O	0
			94	40	52	2	
11	DI	1	Total	C	H	O	0
			94	40	52	2	

Continued on next page...

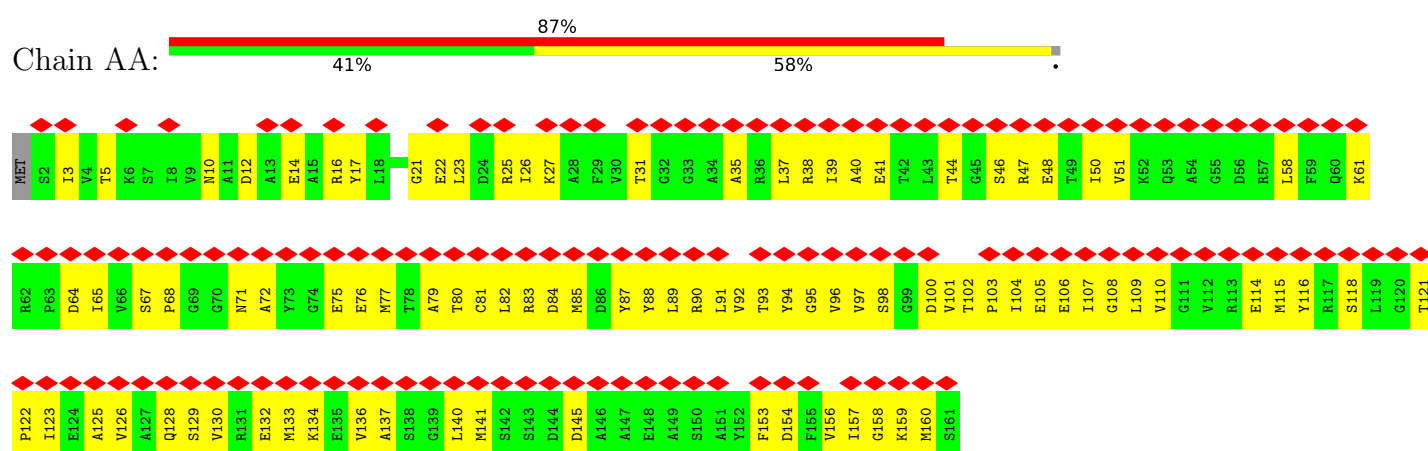
Continued from previous page...

Mol	Chain	Residues	Atoms				AltConf
			Total	C	H	O	
11	EA	1	94	40	52	2	0

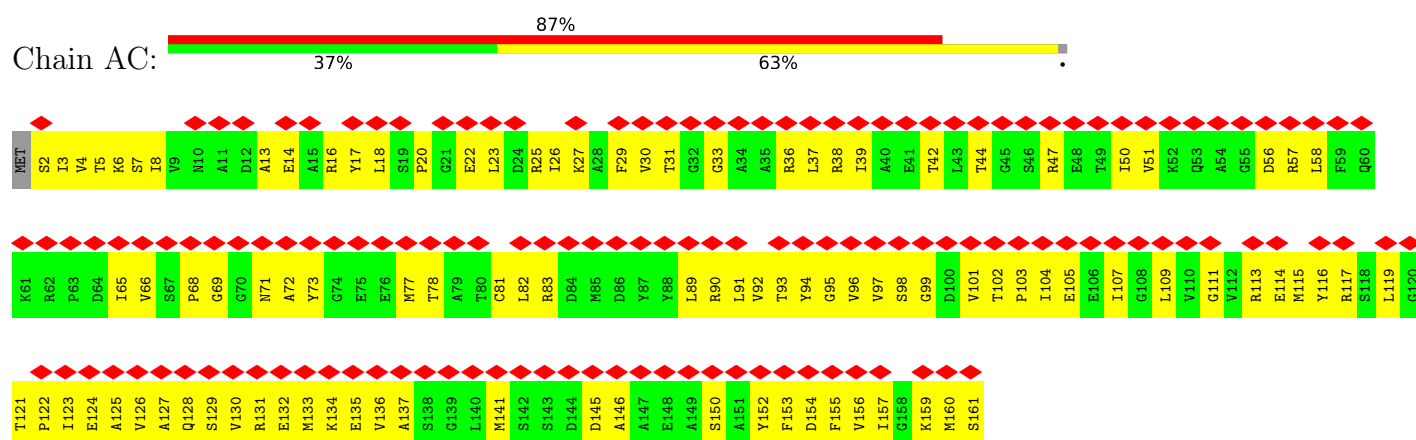
3 Residue-property plots

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

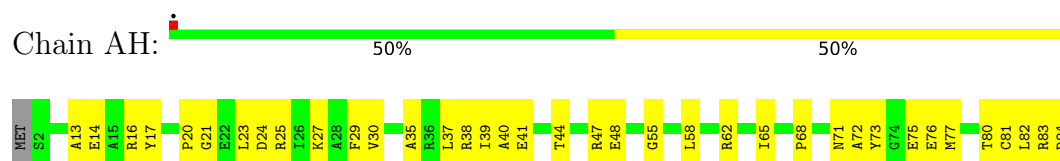
• Molecule 1: Allophycocyanin alpha chain



• Molecule 1: Allophycocyanin alpha chain

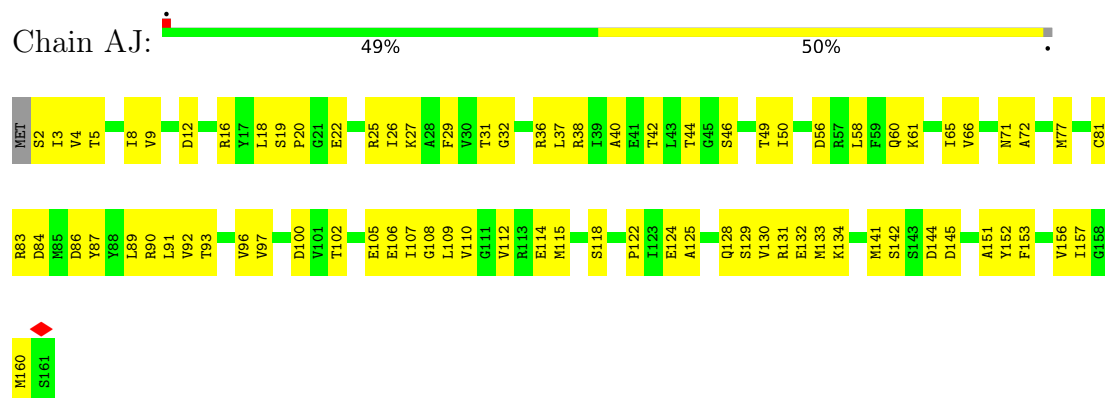


• Molecule 1: Allophycocyanin alpha chain

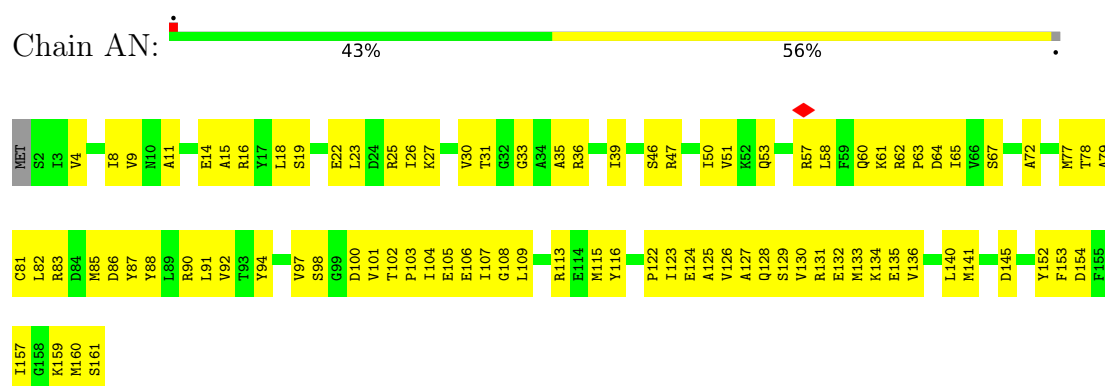




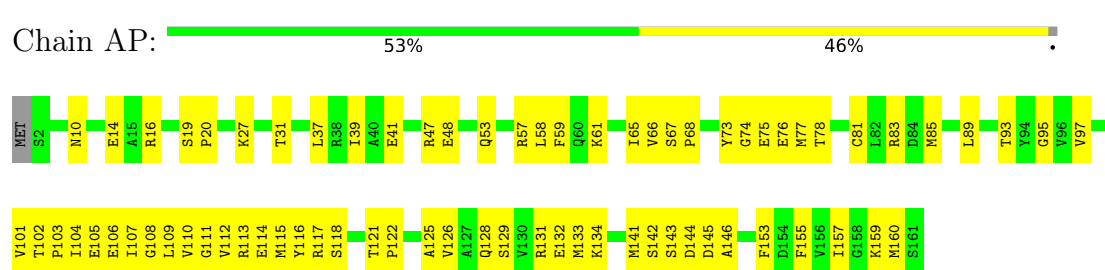
• Molecule 1: Allophycocyanin alpha chain



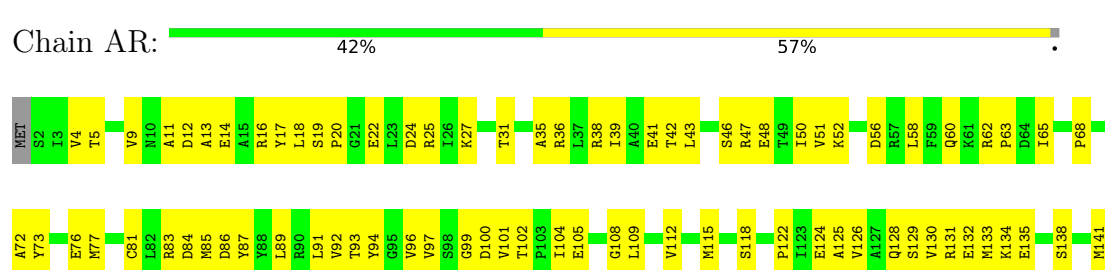
• Molecule 1: Allophycocyanin alpha chain



• Molecule 1: Allophycocyanin alpha chain



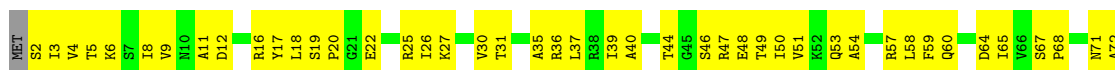
• Molecule 1: Allophycocyanin alpha chain





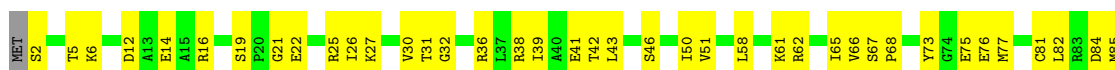
• Molecule 1: Allophycocyanin alpha chain

Chain AV: 34% 65%



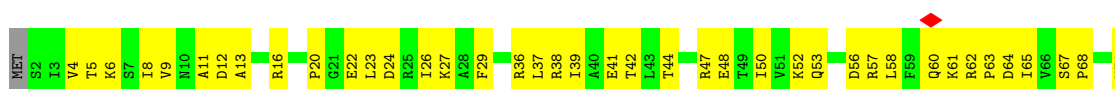
• Molecule 1: Allophycocyanin alpha chain

Chain AX: 47% 52%



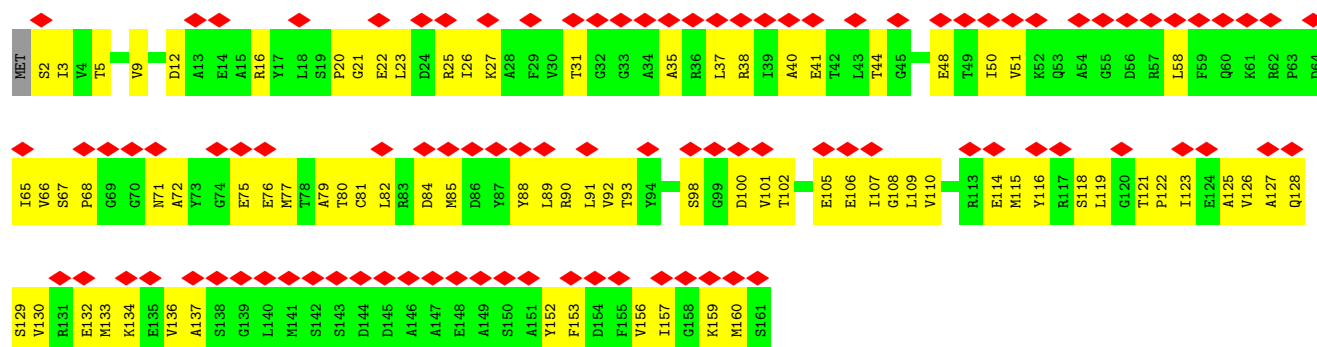
• Molecule 1: Allophycocyanin alpha chain

Chain AZ: 38% 61%



• Molecule 1: Allophycocyanin alpha chain

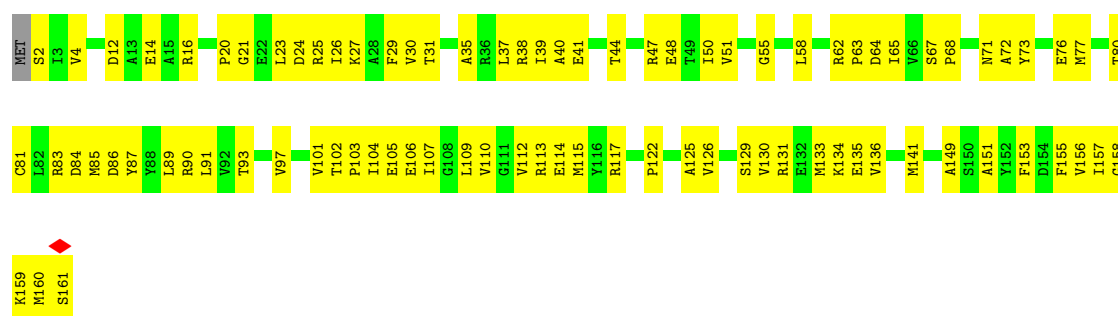
Chain BI: 60% 50% 50%



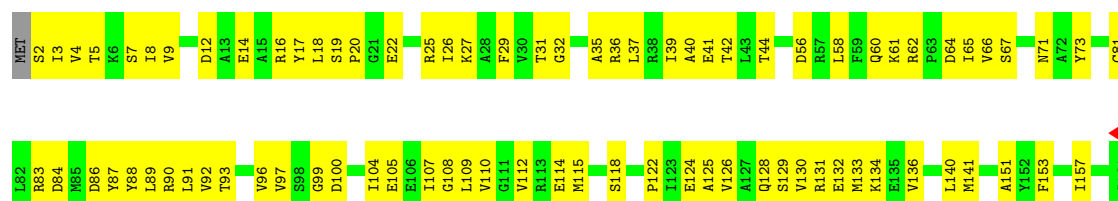
• Molecule 1: Allophycocyanin alpha chain



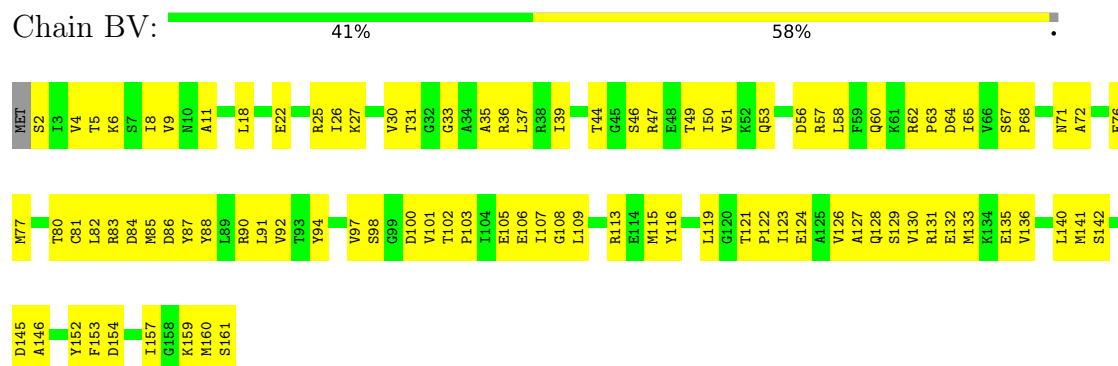
• Molecule 1: Allophycocyanin alpha chain



• Molecule 1: Allophycocyanin alpha chain



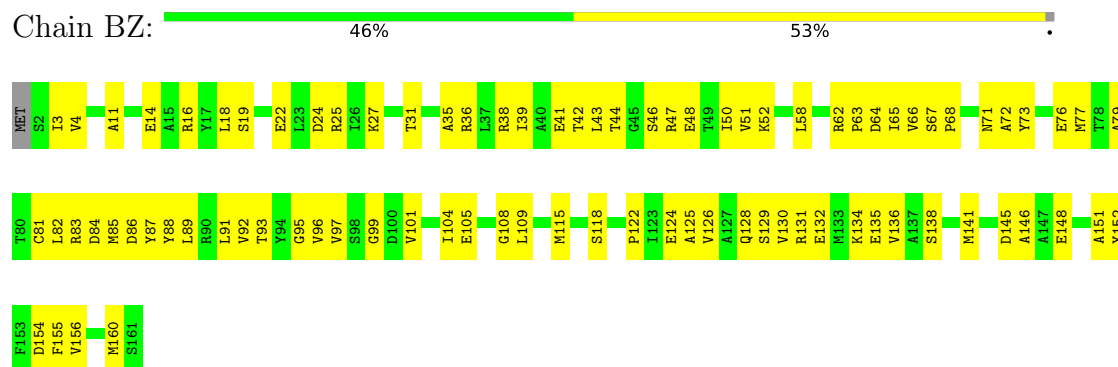
● Molecule 1: Allophycocyanin alpha chain



● Molecule 1: Allophycocyanin alpha chain



● Molecule 1: Allophycocyanin alpha chain

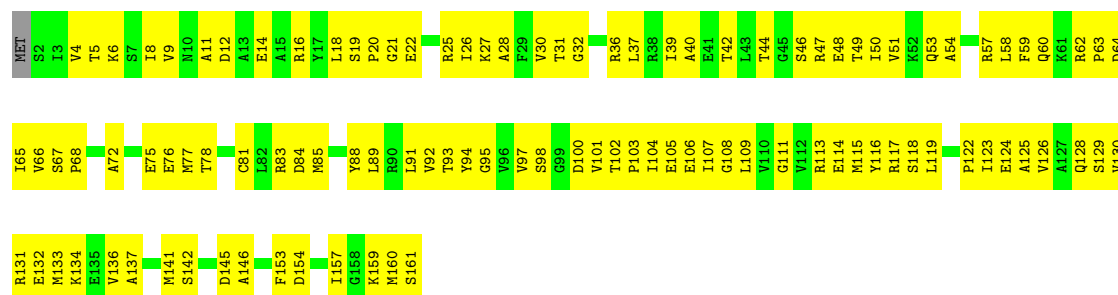


● Molecule 1: Allophycocyanin alpha chain



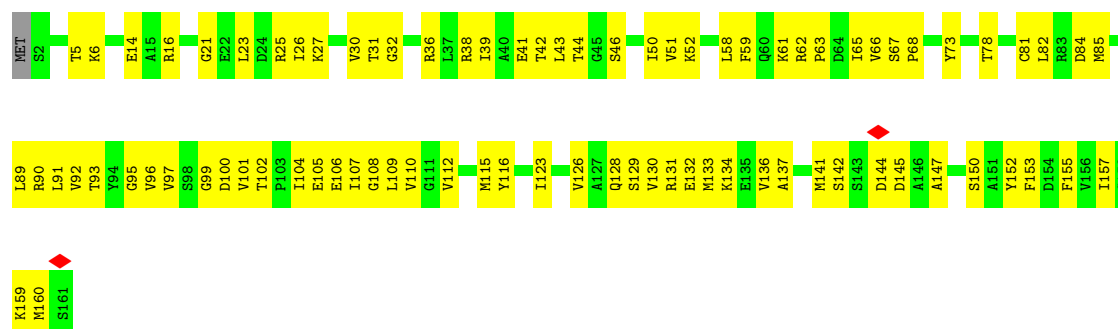
- Molecule 1: Allophycocyanin alpha chain

Chain CE:  34% 66%



- Molecule 1: Allophycocyanin alpha chain

Chain CG:  48% 52%

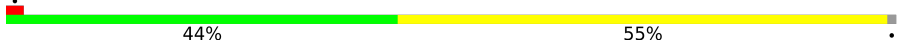


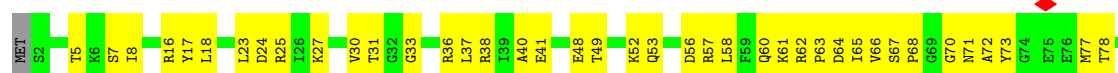
- Molecule 1: Allophycocyanin alpha chain

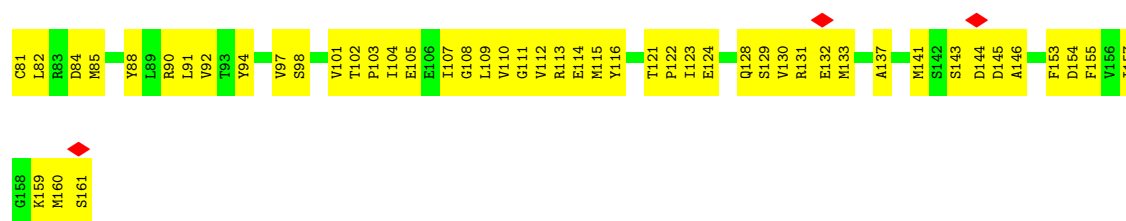
Chain CR:  6% 37% 63%



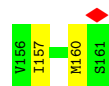
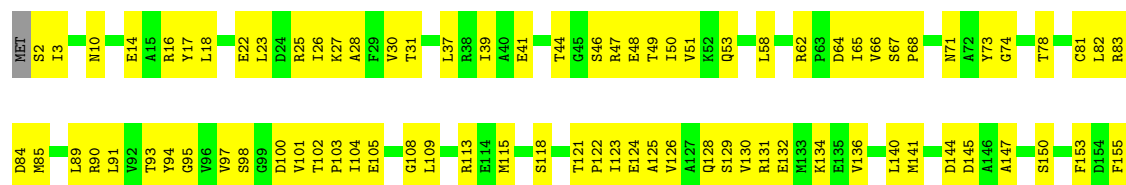
- Molecule 1: Allophycocyanin alpha chain

Chain CT:  44% 55%

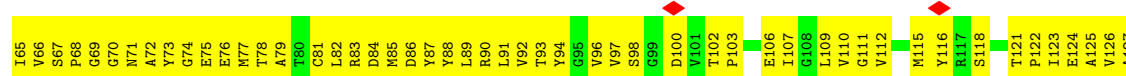
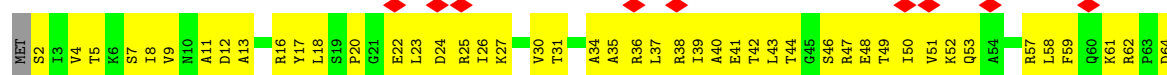




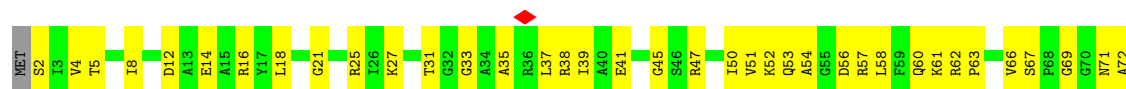
• Molecule 1: Allophycocyanin alpha chain



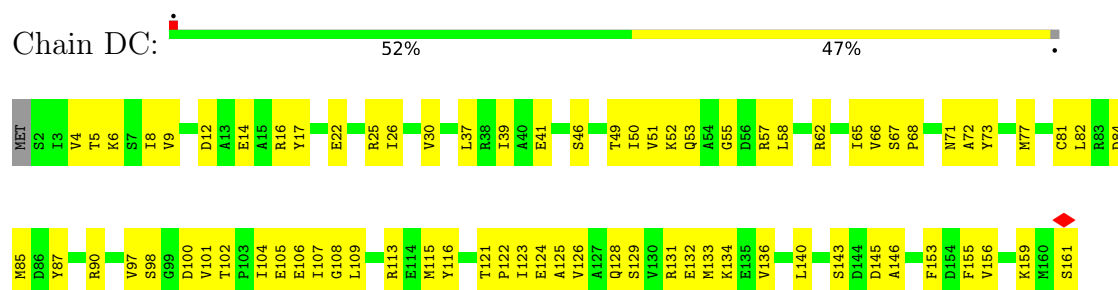
• Molecule 1: Allophycocyanin alpha chain



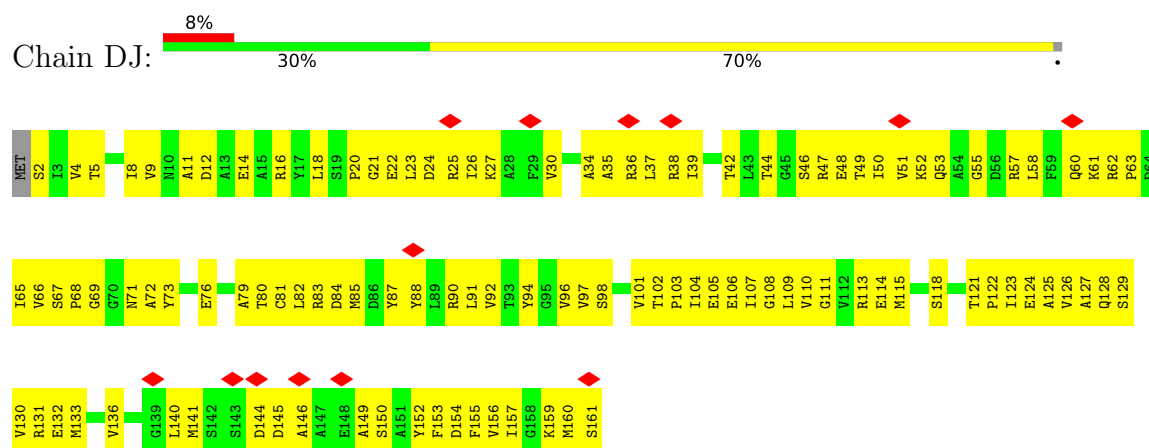
• Molecule 1: Allophycocyanin alpha chain



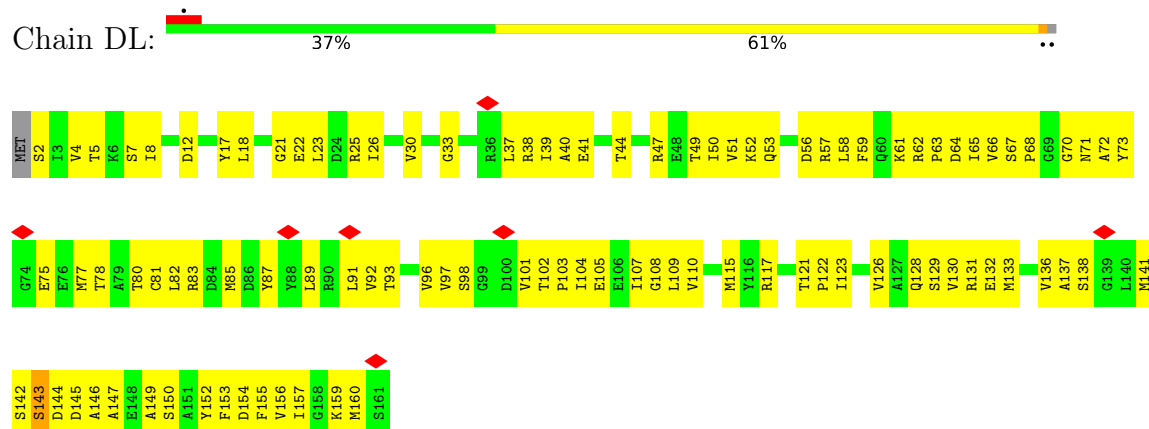
- Molecule 1: Allophycocyanin alpha chain



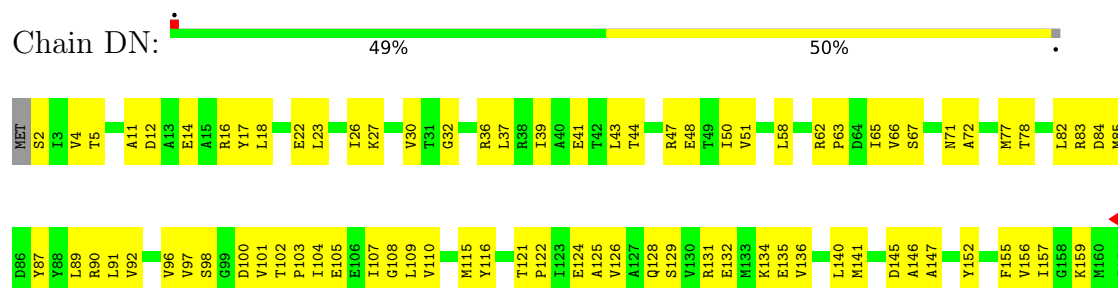
- Molecule 1: Allophycocyanin alpha chain



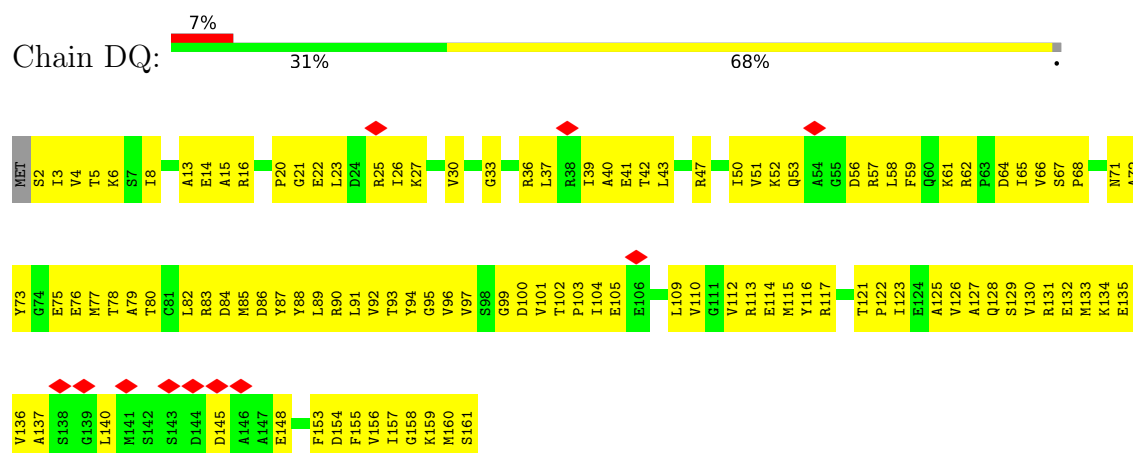
- Molecule 1: Allophycocyanin alpha chain



- Molecule 1: Allophycocyanin alpha chain



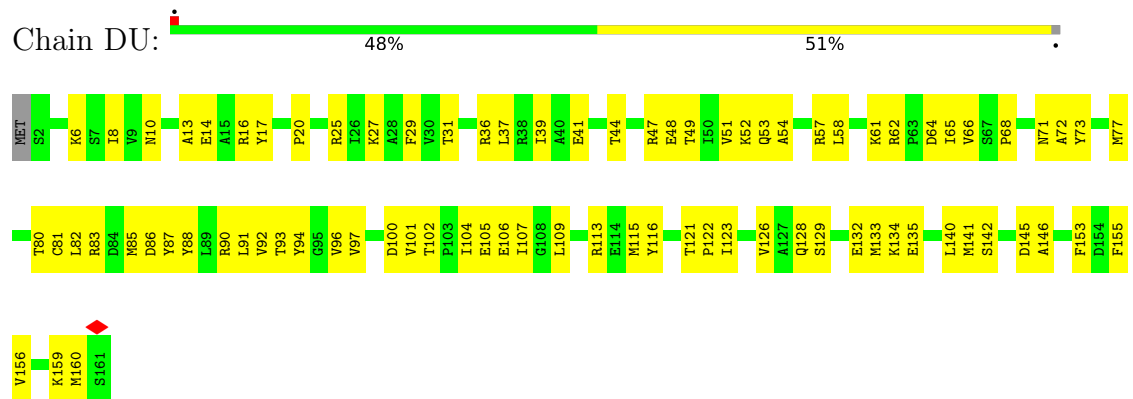
- Molecule 1: Allophycocyanin alpha chain



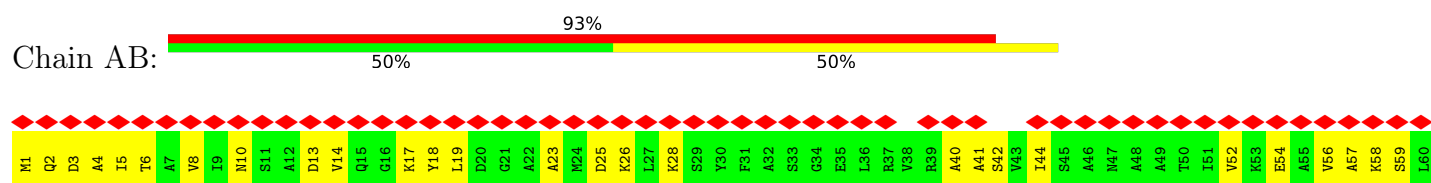
- Molecule 1: Allophycocyanin alpha chain

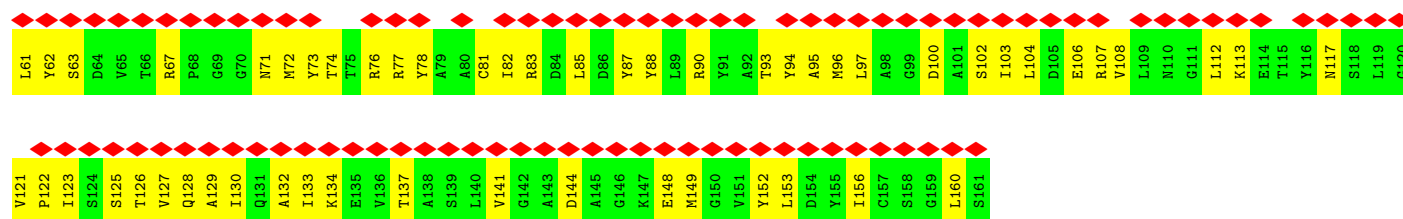


- Molecule 1: Allophycocyanin alpha chain

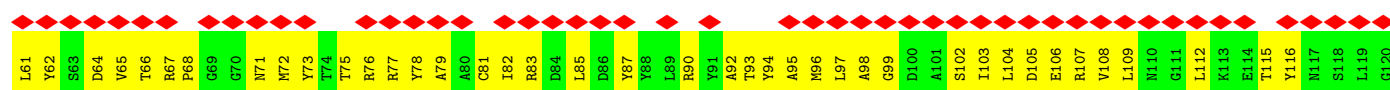


- Molecule 2: Allophycocyanin beta chain

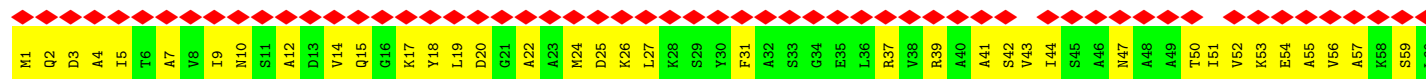




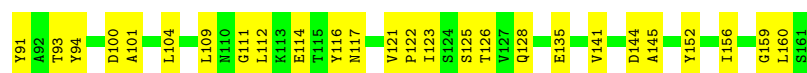
• Molecule 2: Allophycocyanin beta chain



• Molecule 2: Allophycocyanin beta chain

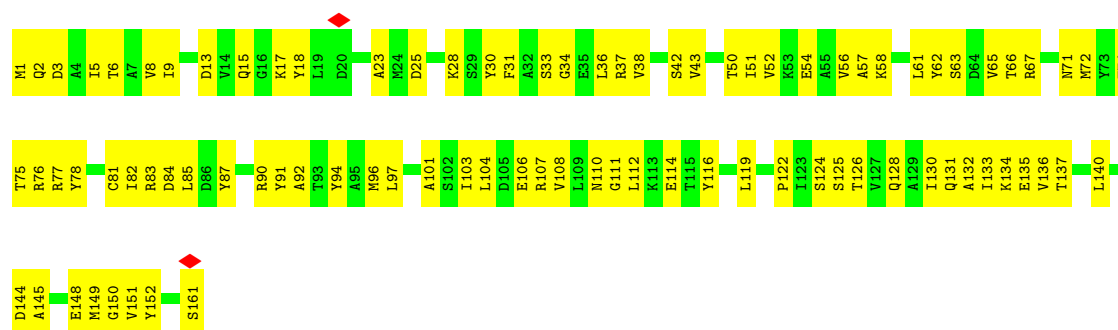


• Molecule 2: Allophycocyanin beta chain



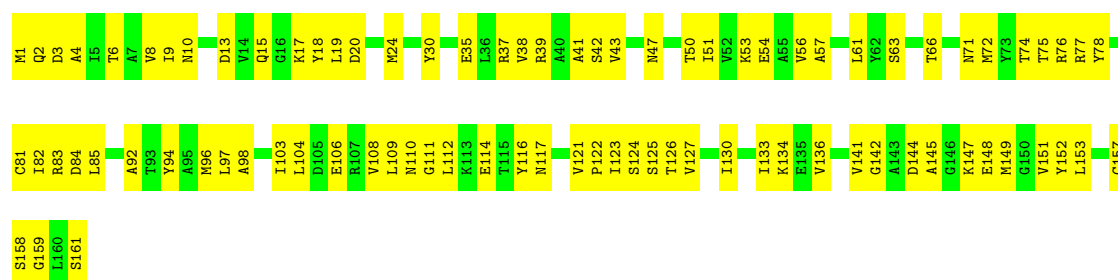
• Molecule 2: Allophycocyanin beta chain





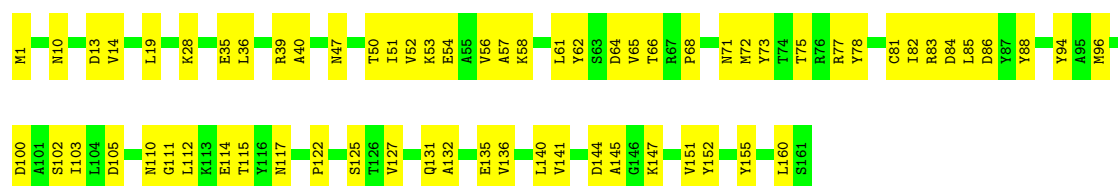
• Molecule 2: Allophycocyanin beta chain

Chain AO: 47% 53%



• Molecule 2: Allophycocyanin beta chain

Chain AQ: 59% 41%

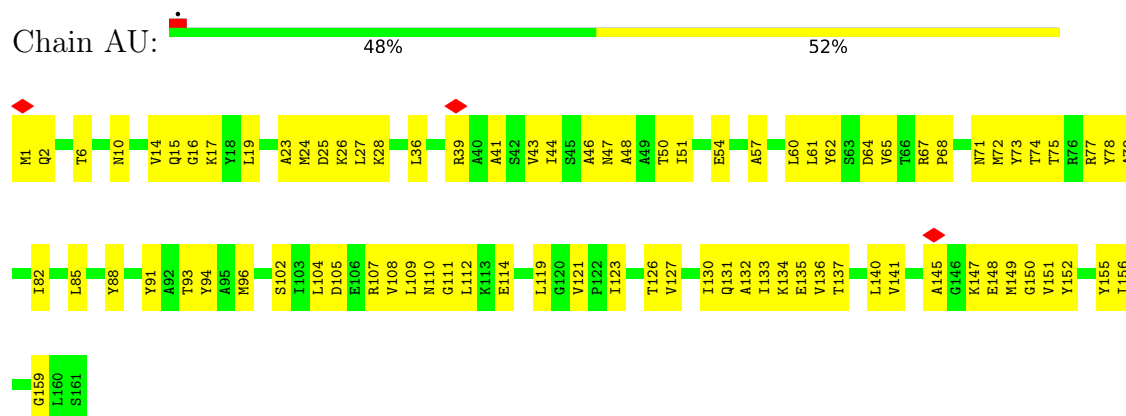


• Molecule 2: Allophycocyanin beta chain

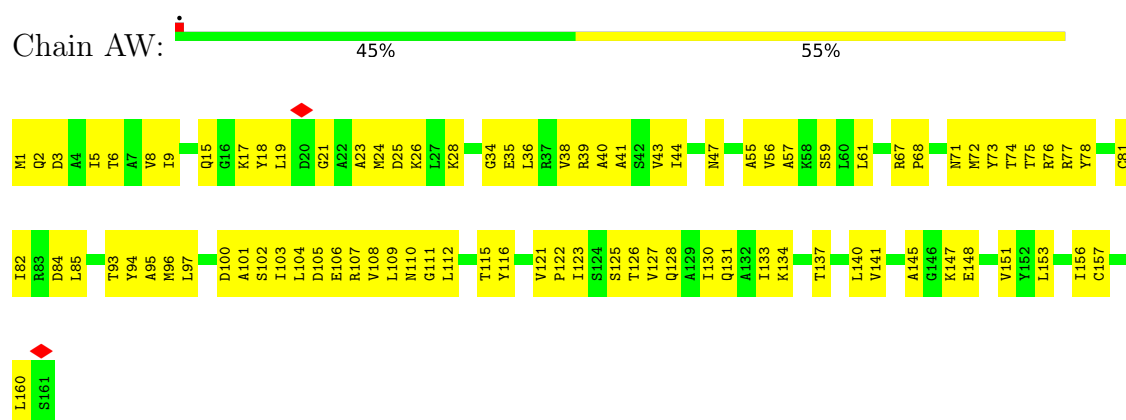
Chain AS: 41% 59%



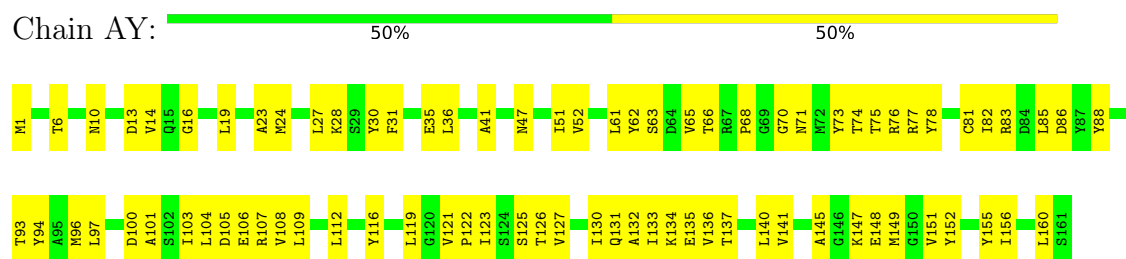
• Molecule 2: Allophycocyanin beta chain



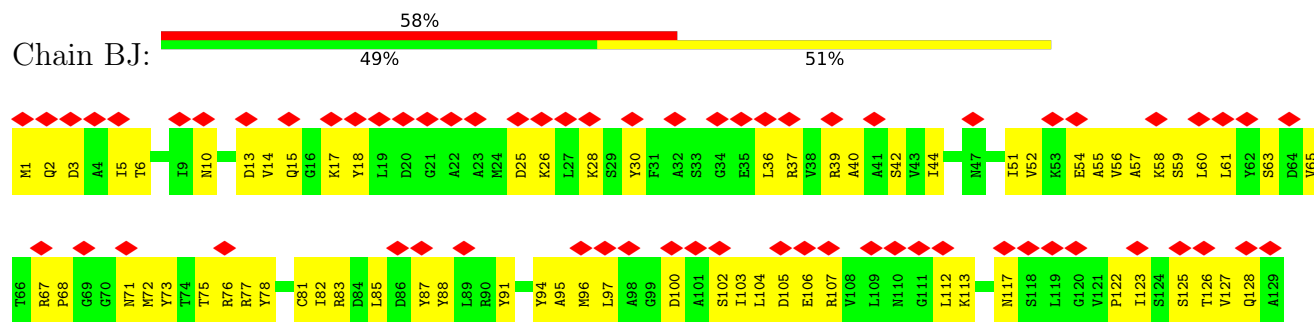
• Molecule 2: Allophycocyanin beta chain

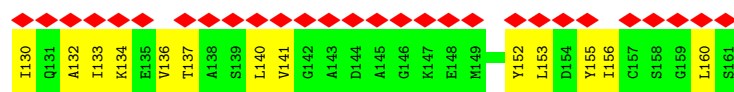


• Molecule 2: Allophycocyanin beta chain

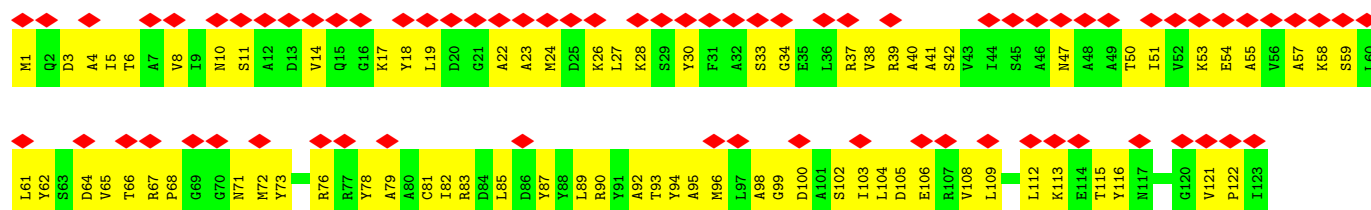


• Molecule 2: Allophycocyanin beta chain

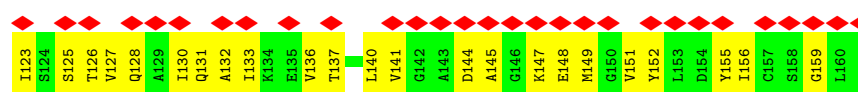
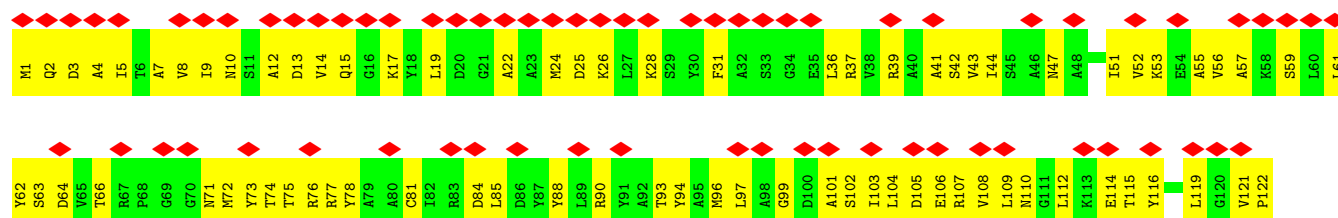




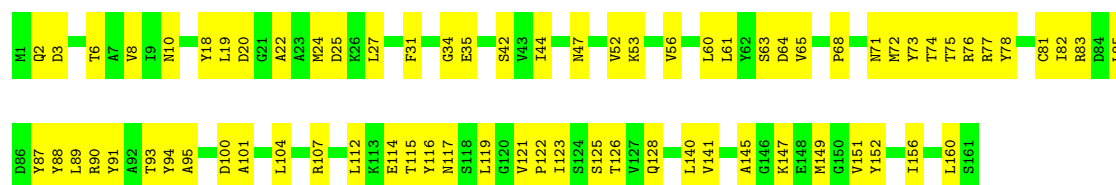
• Molecule 2: Allophycocyanin beta chain



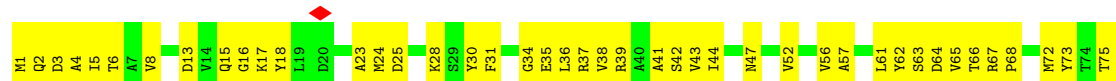
• Molecule 2: Allophycocyanin beta chain

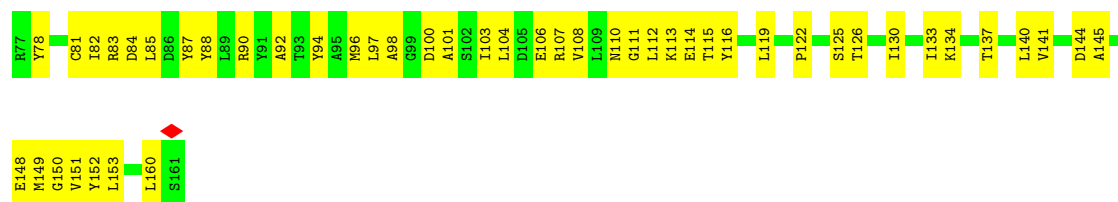


• Molecule 2: Allophycocyanin beta chain

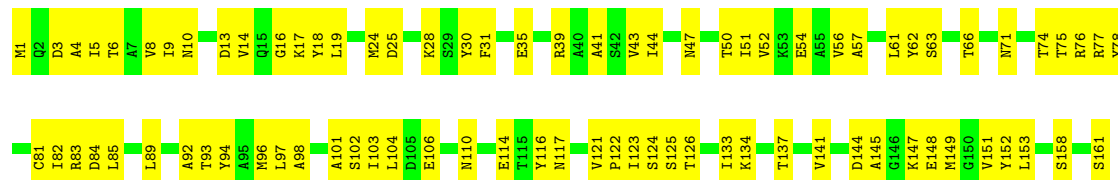


• Molecule 2: Allophycocyanin beta chain

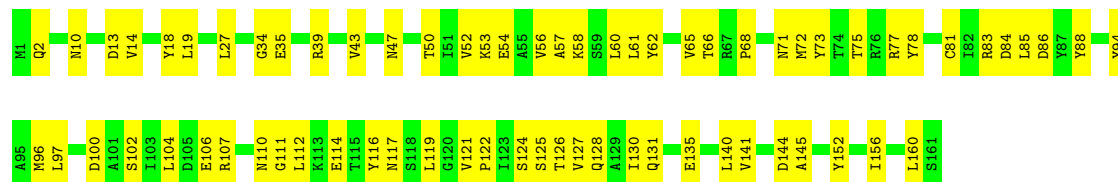




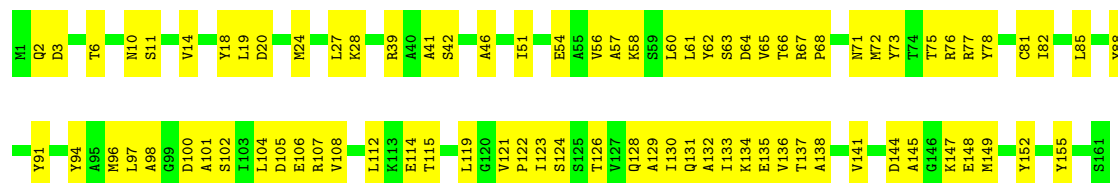
• Molecule 2: Allophycocyanin beta chain



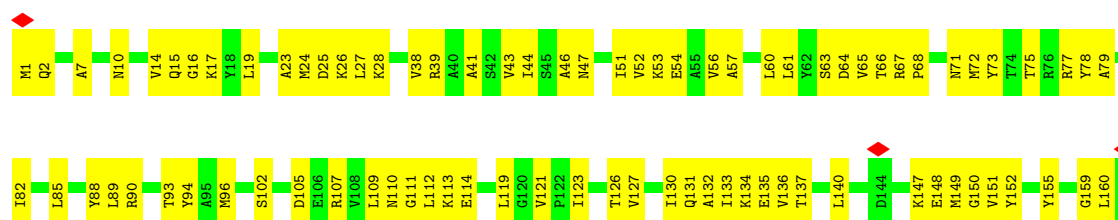
• Molecule 2: Allophycocyanin beta chain



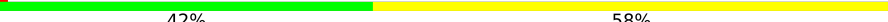
• Molecule 2: Allophycocyanin beta chain

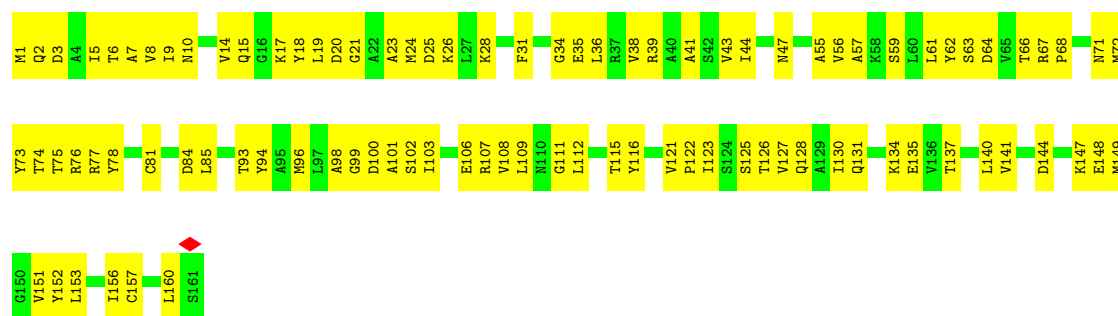


• Molecule 2: Allophycocyanin beta chain



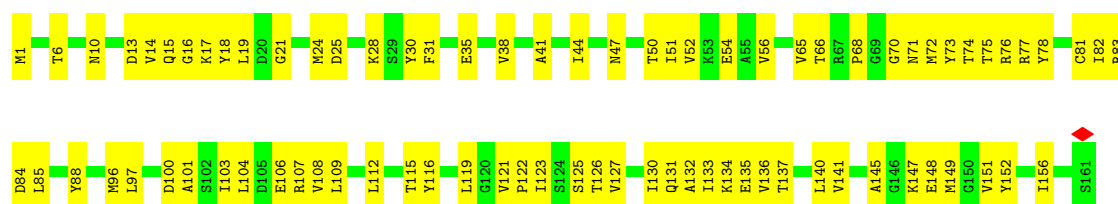
• Molecule 2: Allophycocyanin beta chain

Chain CF:  42% 58%



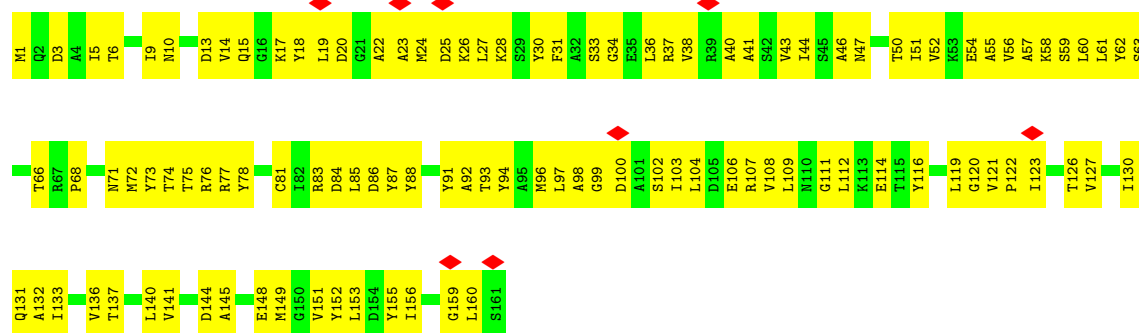
• Molecule 2: Allophycocyanin beta chain

Chain CH:  50% 50%



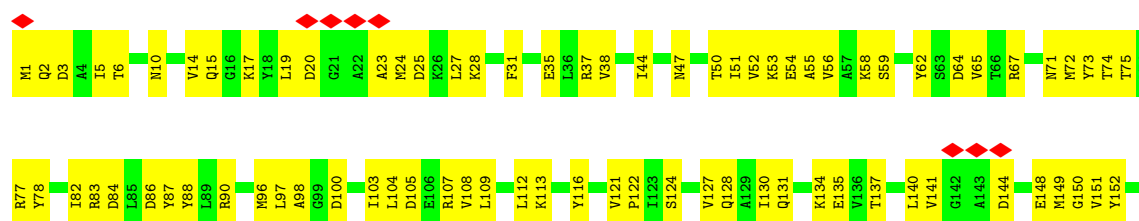
• Molecule 2: Allophycocyanin beta chain

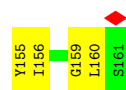
Chain CS:  5% 32% 68%



• Molecule 2: Allophycocyanin beta chain

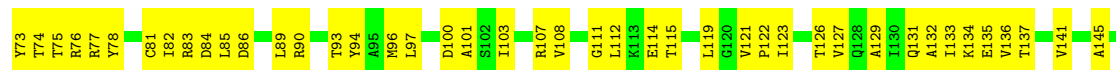
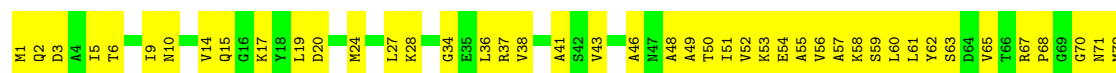
Chain CU:  6% 48% 52%





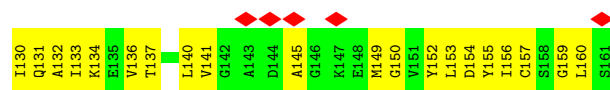
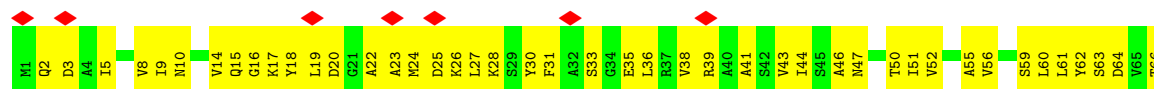
• Molecule 2: Allophycocyanin beta chain

Chain CW: 43% 57%



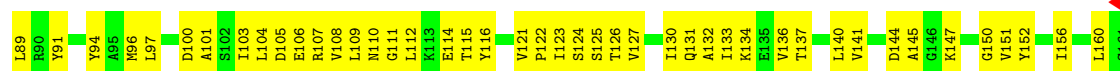
• Molecule 2: Allophycocyanin beta chain

Chain CZ: 8% 31% 69%



• Molecule 2: Allophycocyanin beta chain

Chain DB: 47% 53%

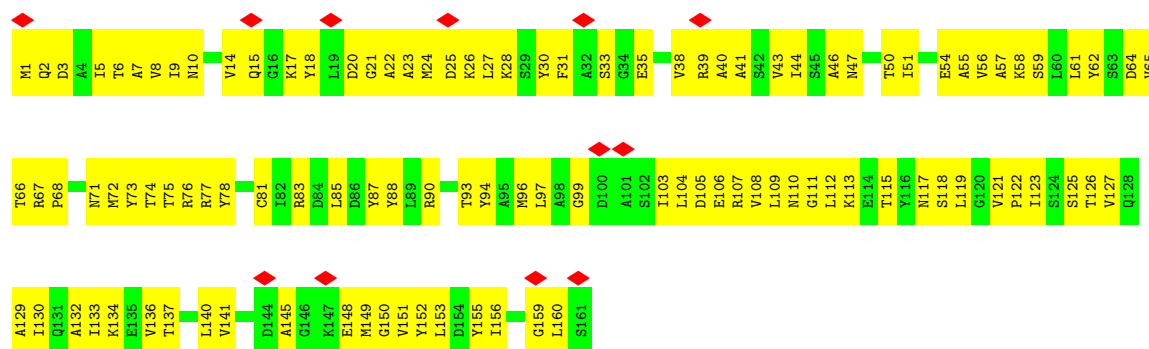


• Molecule 2: Allophycocyanin beta chain

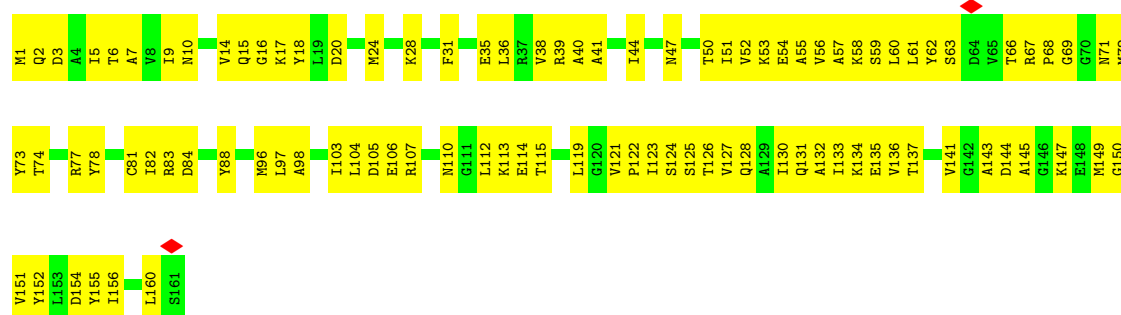
Chain DD: 43% 57%



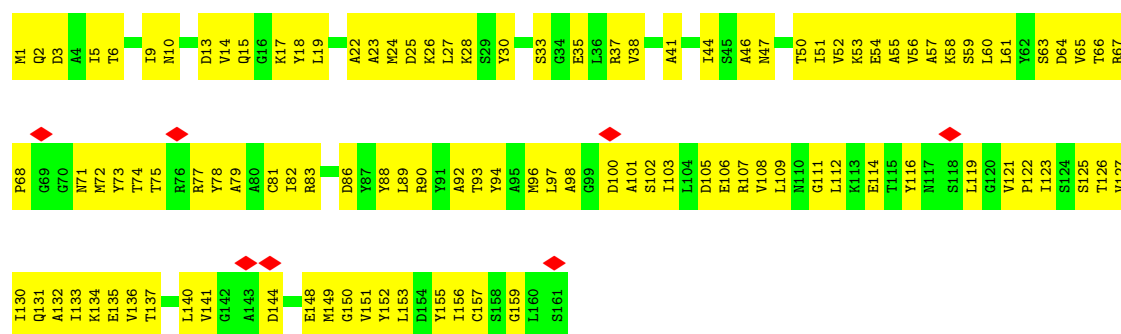




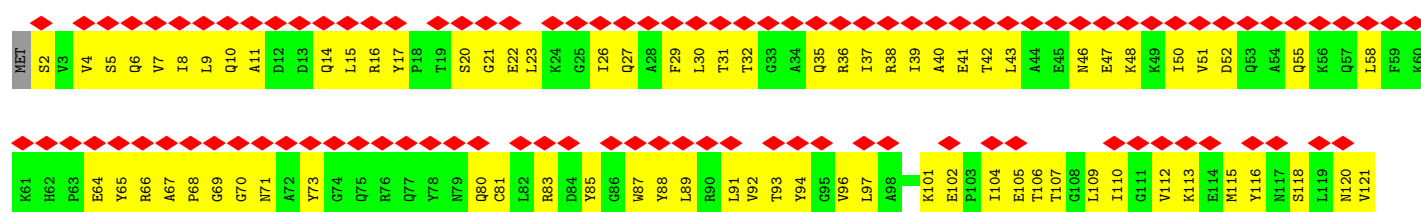
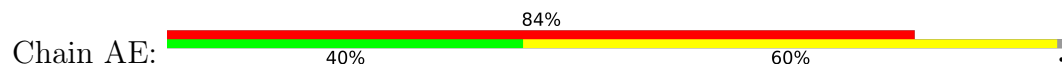
• Molecule 2: Allophycocyanin beta chain



• Molecule 2: Allophycocyanin beta chain

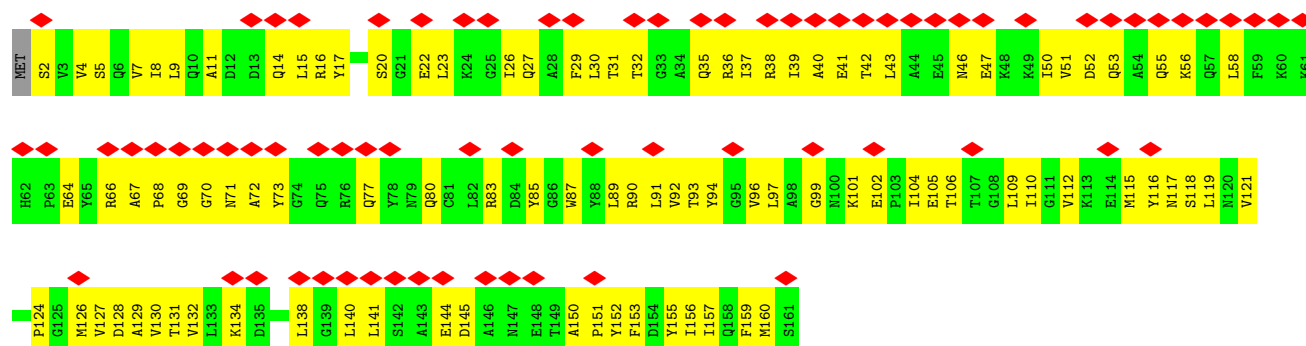


• Molecule 3: Allophycocyanin subunit alpha-B

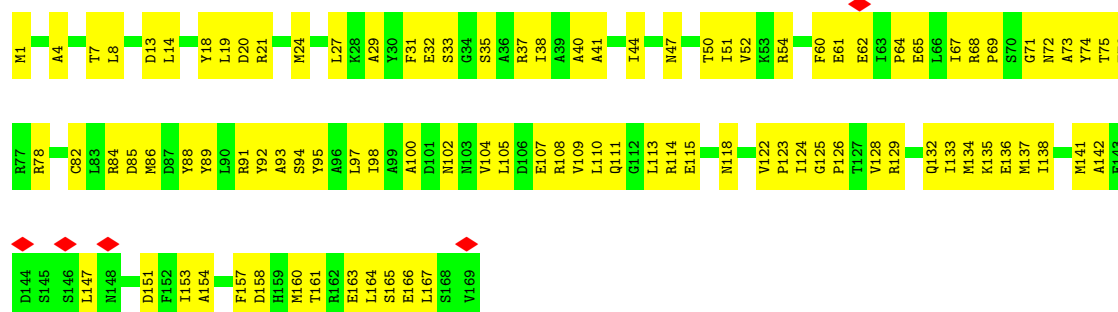




• Molecule 3: Allophycocyanin subunit alpha-B



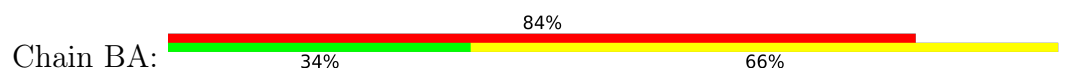
• Molecule 4: Allophycocyanin subunit beta-18

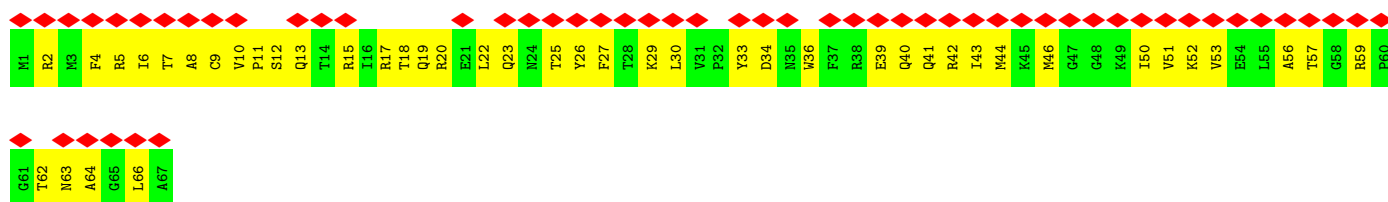


• Molecule 4: Allophycocyanin subunit beta-18

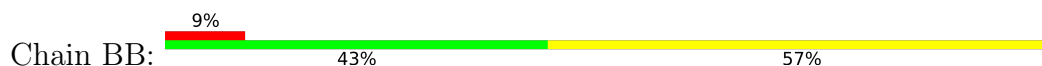


• Molecule 5: Phycobilisome 7.8 kDa linker polypeptide, allophycocyanin-associated, core





- Molecule 5: Phycobilisome 7.8 kDa linker polypeptide, allophycocyanin-associated, core



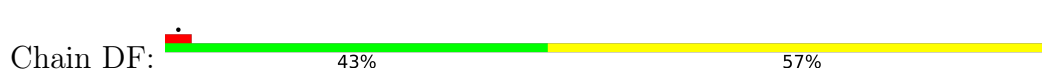
- Molecule 5: Phycobilisome 7.8 kDa linker polypeptide, allophycocyanin-associated, core



- Molecule 5: Phycobilisome 7.8 kDa linker polypeptide, allophycocyanin-associated, core

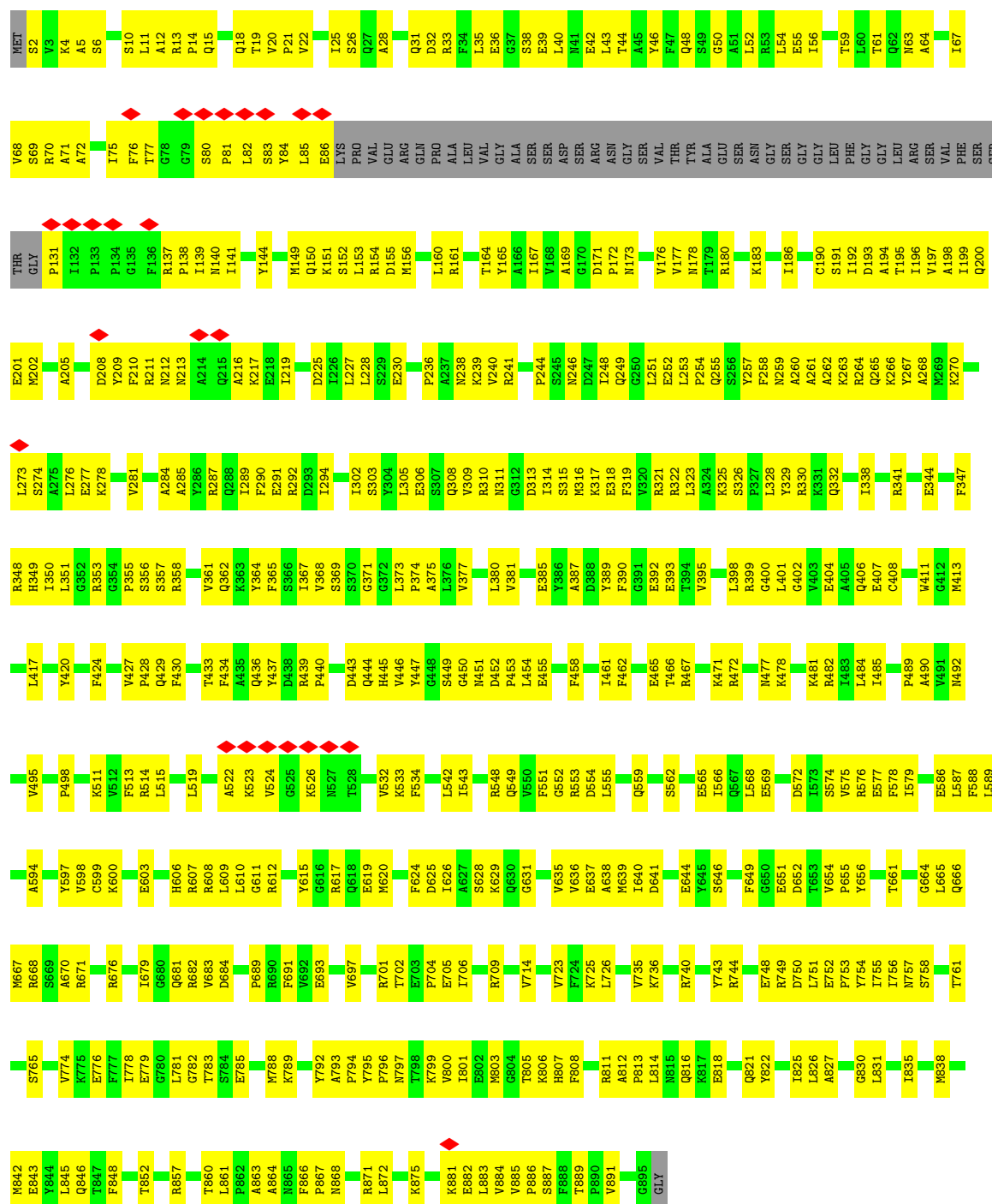


- Molecule 5: Phycobilisome 7.8 kDa linker polypeptide, allophycocyanin-associated, core



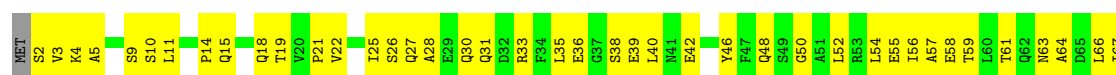
- Molecule 6: Phycobiliprotein ApcE

Chain BD: 

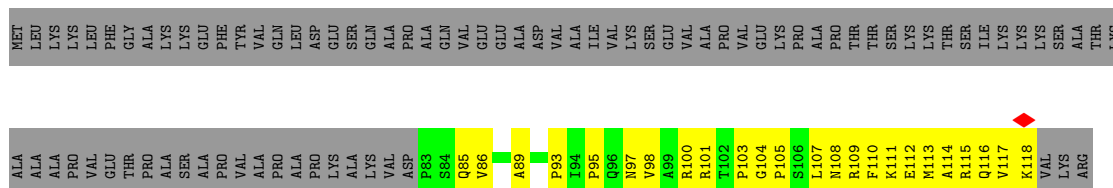


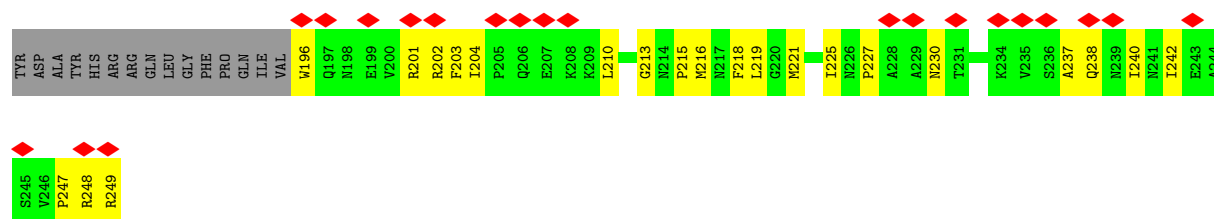
• Molecule 6: Phycobiliprotein ApcE

Chain CM: 

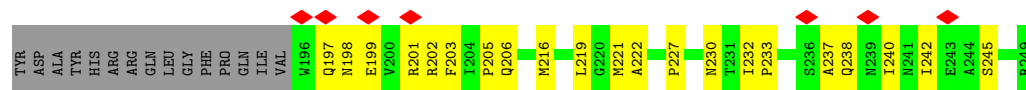
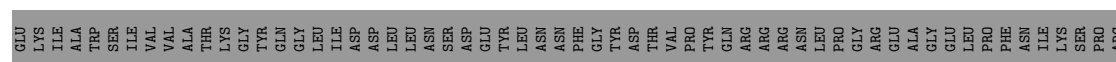
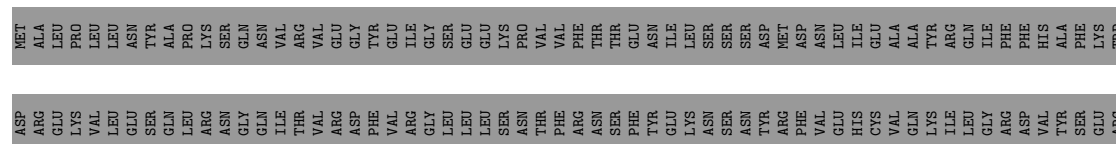


- Molecule 7: Sll1873 protein

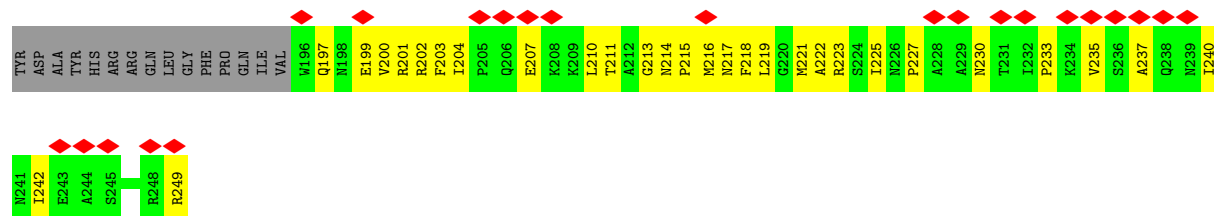
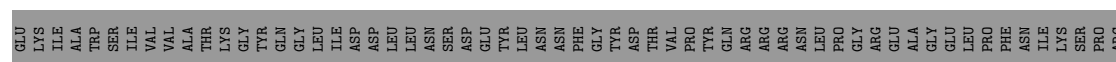
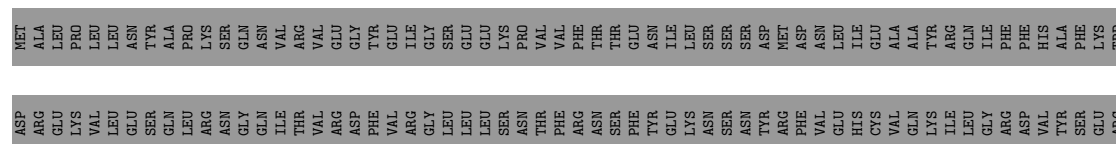




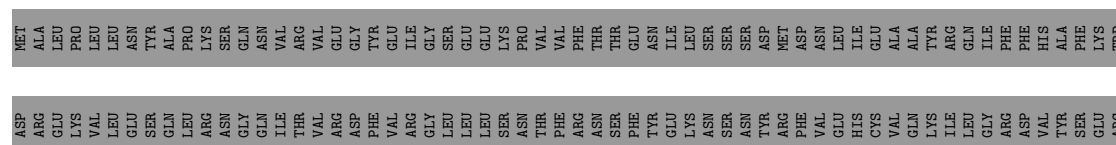
• Molecule 8: Phycobilisome rod-core linker polypeptide CpcG

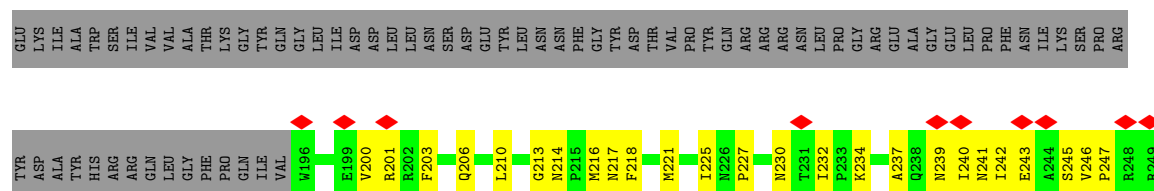


• Molecule 8: Phycobilisome rod-core linker polypeptide CpcG

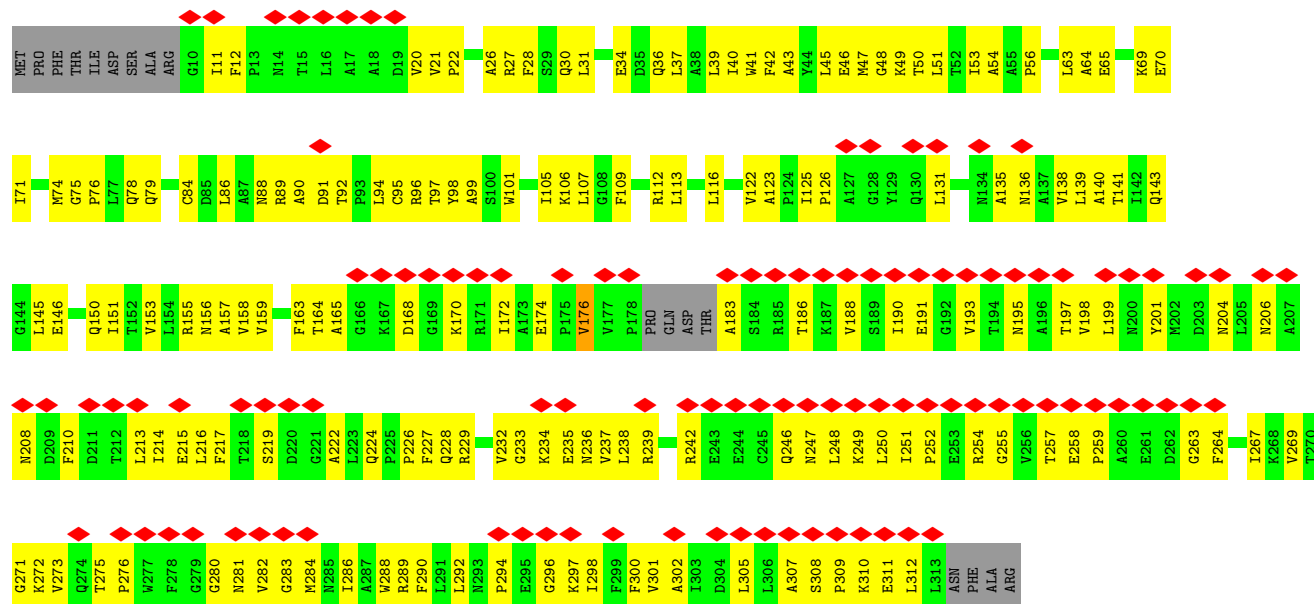
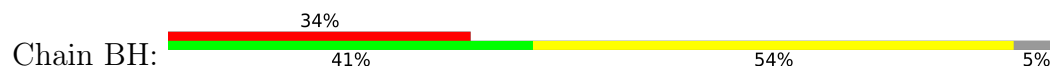


• Molecule 8: Phycobilisome rod-core linker polypeptide CpcG

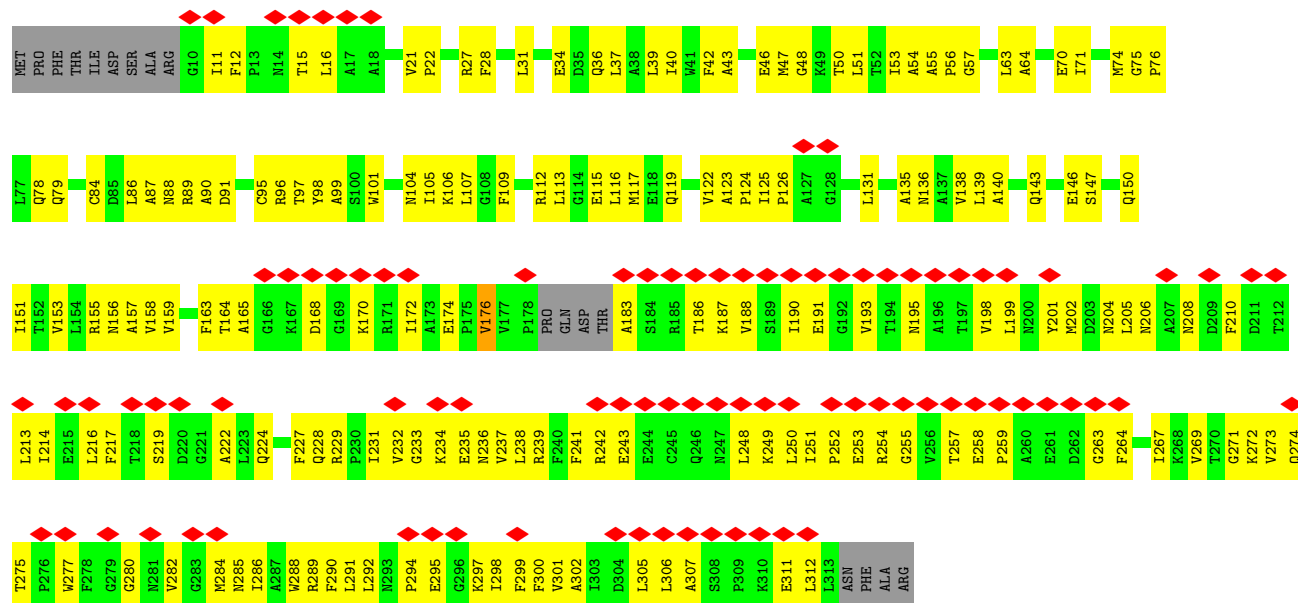




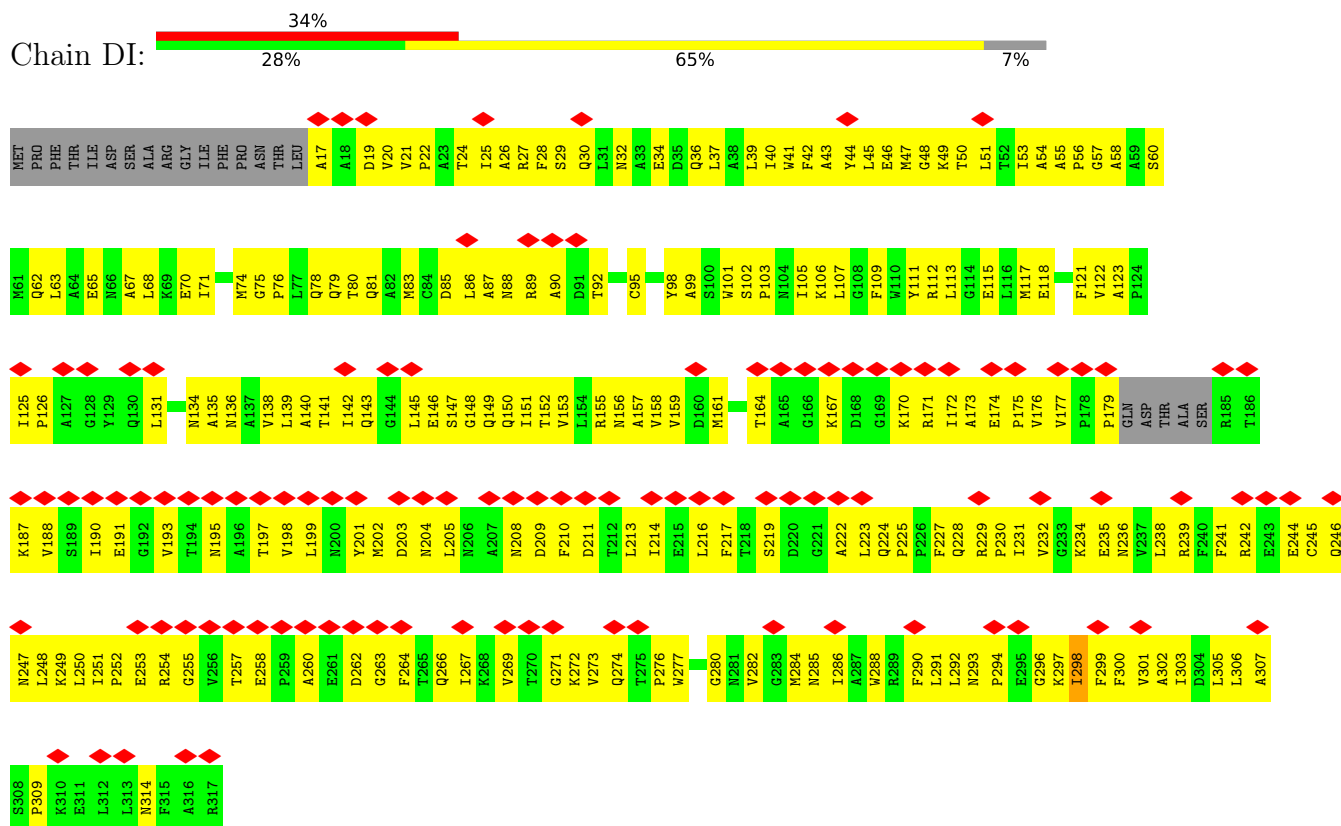
• Molecule 9: Orange carotenoid-binding protein



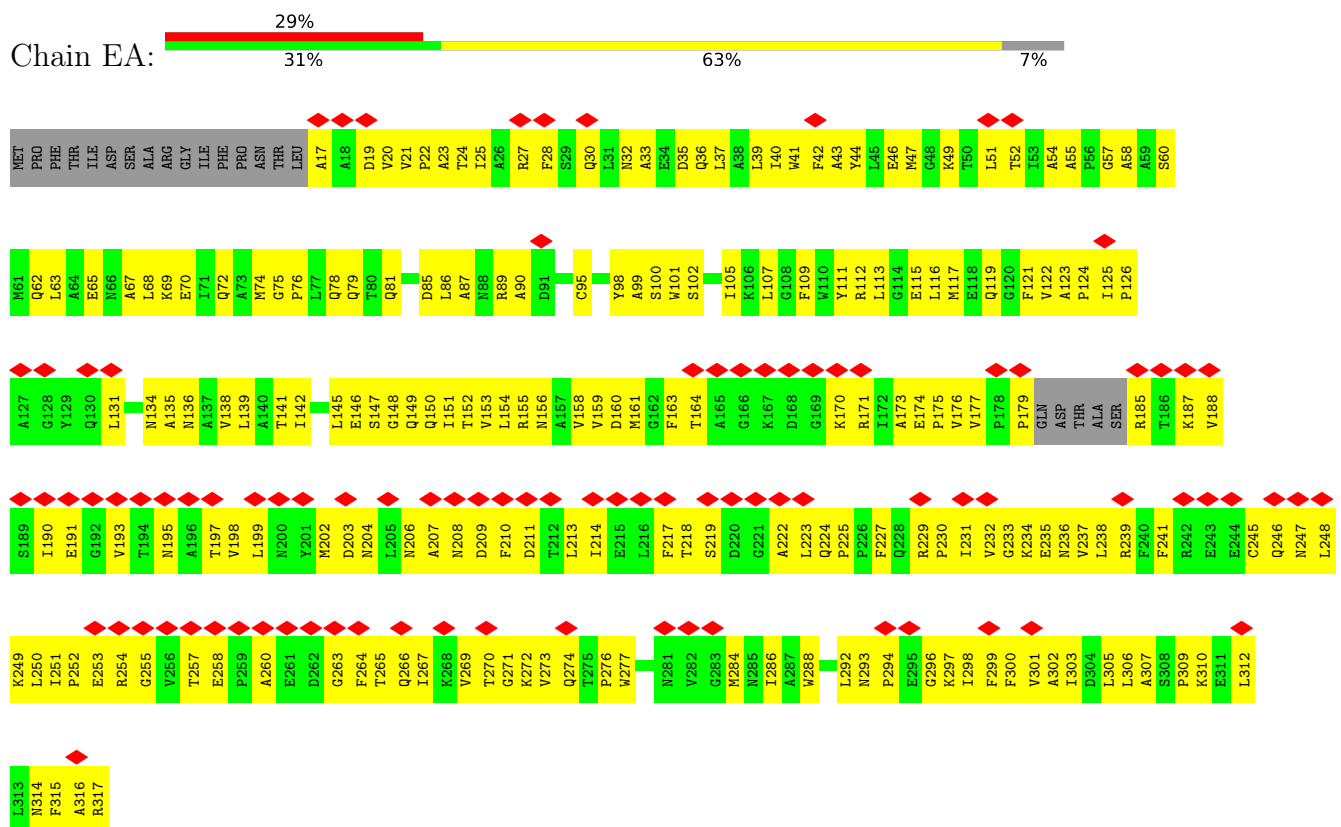
• Molecule 9: Orange carotenoid-binding protein



• Molecule 9: Orange carotenoid-binding protein



• Molecule 9: Orange carotenoid-binding protein



4 Experimental information

Property	Value	Source
EM reconstruction method	SINGLE PARTICLE	Depositor
Imposed symmetry	POINT, C1	Depositor
Number of particles used	340839	Depositor
Resolution determination method	FSC 0.143 CUT-OFF	Depositor
CTF correction method	PHASE FLIPPING AND AMPLITUDE CORRECTION	Depositor
Microscope	TFS KRIOS	Depositor
Voltage (kV)	300	Depositor
Electron dose ($e^-/\text{\AA}^2$)	50	Depositor
Minimum defocus (nm)	500	Depositor
Maximum defocus (nm)	1600	Depositor
Magnification	Not provided	
Image detector	GATAN K3 (6k x 4k)	Depositor
Maximum map value	2.199	Depositor
Minimum map value	-0.742	Depositor
Average map value	0.021	Depositor
Map value standard deviation	0.113	Depositor
Recommended contour level	0.437	Depositor
Map size (Å)	377.99997, 377.99997, 377.99997	wwPDB
Map dimensions	360, 360, 360	wwPDB
Map angles (°)	90.0, 90.0, 90.0	wwPDB
Pixel spacing (Å)	1.05, 1.05, 1.05	Depositor

5 Model quality

5.1 Standard geometry

Bond lengths and bond angles in the following residue types are not validated in this section: CYC, 45D

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	AA	0.10	0/1225	0.24	0/1652
1	AC	0.12	0/1225	0.29	0/1652
1	AH	0.16	0/1225	0.28	0/1652
1	AJ	0.20	0/1225	0.29	0/1652
1	AN	0.16	0/1225	0.33	0/1652
1	AP	0.18	0/1225	0.27	0/1652
1	AR	0.16	0/1225	0.28	0/1652
1	AV	0.14	0/1225	0.29	0/1652
1	AX	0.14	0/1225	0.27	0/1652
1	AZ	0.14	0/1225	0.24	0/1652
1	BI	0.11	0/1225	0.22	0/1652
1	BK	0.14	0/1225	0.32	0/1652
1	BP	0.16	0/1225	0.28	0/1652
1	BR	0.19	0/1225	0.29	0/1652
1	BV	0.17	0/1225	0.32	0/1652
1	BX	0.19	0/1225	0.26	0/1652
1	BZ	0.17	0/1225	0.27	0/1652
1	CC	0.14	0/1225	0.25	0/1652
1	CE	0.14	0/1225	0.28	0/1652
1	CG	0.16	0/1225	0.28	0/1652
1	CR	0.15	0/1225	0.32	0/1652
1	CT	0.15	0/1225	0.36	0/1652
1	CV	0.16	0/1225	0.30	0/1652
1	CY	0.16	0/1225	0.36	0/1652
1	DA	0.17	0/1225	0.35	0/1652
1	DC	0.18	0/1225	0.29	0/1652
1	DJ	0.15	0/1225	0.31	0/1652
1	DL	0.15	0/1225	0.33	0/1652
1	DN	0.16	0/1225	0.27	0/1652
1	DQ	0.15	0/1225	0.33	0/1652
1	DS	0.17	0/1225	0.33	0/1652
1	DU	0.18	0/1225	0.30	0/1652

Mol	Chain	Bond lengths		Bond angles	
		RMSZ	# Z >5	RMSZ	# Z >5
2	AB	0.11	0/1220	0.26	0/1650
2	AD	0.12	0/1220	0.29	0/1650
2	AF	0.10	0/1220	0.26	0/1650
2	AI	0.19	0/1220	0.27	0/1650
2	AL	0.16	0/1220	0.28	0/1650
2	AO	0.20	0/1220	0.30	0/1650
2	AQ	0.19	0/1220	0.26	0/1650
2	AS	0.18	0/1220	0.32	0/1650
2	AU	0.15	0/1220	0.31	0/1650
2	AW	0.14	0/1220	0.27	0/1650
2	AY	0.15	0/1220	0.29	0/1650
2	BJ	0.11	0/1220	0.26	0/1650
2	BL	0.13	0/1220	0.30	0/1650
2	BN	0.11	0/1220	0.24	0/1650
2	BQ	0.20	0/1220	0.27	0/1650
2	BT	0.18	0/1220	0.30	0/1650
2	BW	0.20	0/1220	0.30	0/1650
2	BY	0.19	0/1220	0.25	0/1650
2	CA	0.19	0/1220	0.36	0/1650
2	CD	0.15	0/1220	0.29	0/1650
2	CF	0.15	0/1220	0.28	0/1650
2	CH	0.15	0/1220	0.28	0/1650
2	CS	0.14	0/1220	0.35	0/1650
2	CU	0.16	0/1220	0.29	0/1650
2	CW	0.15	0/1220	0.28	0/1650
2	CZ	0.17	0/1220	0.36	0/1650
2	DB	0.18	0/1220	0.32	0/1650
2	DD	0.15	0/1220	0.28	0/1650
2	DK	0.14	0/1220	0.33	0/1650
2	DM	0.16	0/1220	0.29	0/1650
2	DO	0.15	0/1220	0.31	0/1650
2	DR	0.16	0/1220	0.32	0/1650
2	DT	0.17	0/1220	0.32	0/1650
2	DV	0.16	0/1220	0.37	0/1650
3	AE	0.13	0/1277	0.31	0/1730
3	BM	0.16	0/1277	0.36	0/1730
4	AK	0.18	0/1341	0.31	0/1813
4	BS	0.18	0/1341	0.31	0/1813
5	BA	0.13	0/555	0.28	0/743
5	BB	0.13	0/555	0.29	0/743
5	CJ	0.13	0/555	0.25	0/743
5	CK	0.13	0/555	0.27	0/743
5	DF	0.18	0/555	0.29	0/743

Mol	Chain	Bond lengths		Bond angles	
		RMSZ	# Z >5	RMSZ	# Z >5
5	DX	0.16	0/555	0.29	0/743
6	BD	0.18	0/6907	0.32	0/9337
6	CM	0.19	0/6907	0.33	0/9337
7	BF	0.12	0/283	0.26	0/381
7	CO	0.14	0/283	0.30	0/381
8	BG	0.14	0/459	0.31	0/620
8	CP	0.15	0/459	0.30	0/620
8	DG	0.14	0/434	0.33	0/587
8	DH	0.14	0/434	0.38	0/587
8	DY	0.14	0/434	0.31	0/587
8	DZ	0.15	0/434	0.30	0/587
9	BH	0.13	0/2349	0.31	0/3195
9	CQ	0.14	0/2349	0.31	0/3195
9	DI	0.12	0/2327	0.30	0/3164
9	EA	0.12	0/2327	0.29	0/3164
All	All	0.16	0/115632	0.30	0/156250

There are no bond length outliers.

There are no bond angle outliers.

There are no chirality outliers.

There are no planarity outliers.

5.2 Too-close contacts

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

Mol	Chain	Non-H	H(model)	H(added)	Clashes	Symm-Clashes
1	AA	1210	0	1210	109	0
1	AC	1210	0	1210	135	0
1	AH	1210	0	1210	104	0
1	AJ	1210	0	1210	90	0
1	AN	1210	0	1210	119	0
1	AP	1210	0	1210	83	0
1	AR	1210	0	1211	97	0
1	AV	1210	0	1210	125	0
1	AX	1210	0	1210	100	0
1	AZ	1210	0	1210	107	0

Continued on next page...

Continued from previous page...

Mol	Chain	Non-H	H(model)	H(added)	Clashes	Symm-Clashes
1	BI	1210	0	1210	86	0
1	BK	1210	0	1210	138	0
1	BP	1210	0	1210	99	0
1	BR	1210	0	1210	91	0
1	BV	1210	0	1210	121	0
1	BX	1210	0	1210	72	0
1	BZ	1210	0	1210	100	0
1	CC	1210	0	1210	111	0
1	CE	1210	0	1210	132	0
1	CG	1210	0	1210	105	0
1	CR	1210	0	1210	152	0
1	CT	1210	0	1210	105	0
1	CV	1210	0	1210	93	0
1	CY	1210	0	1210	186	0
1	DA	1210	0	1210	135	0
1	DC	1210	0	1210	80	0
1	DJ	1210	0	1210	156	0
1	DL	1210	0	1210	131	0
1	DN	1210	0	1210	85	0
1	DQ	1210	0	1210	169	0
1	DS	1210	0	1210	122	0
1	DU	1210	0	1210	102	0
2	AB	1206	0	1218	85	0
2	AD	1206	0	1218	125	0
2	AF	1206	0	1218	120	0
2	AI	1206	0	1218	76	0
2	AL	1206	0	1218	102	0
2	AO	1206	0	1218	93	0
2	AQ	1206	0	1218	59	0
2	AS	1206	0	1218	117	0
2	AU	1206	0	1218	98	0
2	AW	1206	0	1218	111	0
2	AY	1206	0	1218	91	0
2	BJ	1206	0	1218	85	0
2	BL	1206	0	1218	120	0
2	BN	1206	0	1218	124	0
2	BQ	1206	0	1218	83	0
2	BT	1206	0	1218	106	0
2	BW	1206	0	1218	99	0
2	BY	1206	0	1218	72	0
2	CA	1206	0	1218	102	0
2	CD	1206	0	1218	100	0

Continued on next page...

Continued from previous page...

Mol	Chain	Non-H	H(model)	H(added)	Clashes	Symm-Clashes
2	CF	1206	0	1218	129	0
2	CH	1206	0	1218	87	0
2	CS	1206	0	1218	139	0
2	CU	1206	0	1218	84	0
2	CW	1206	0	1218	117	0
2	CZ	1206	0	1218	152	0
2	DB	1206	0	1218	110	0
2	DD	1206	0	1218	108	0
2	DK	1206	0	1218	124	0
2	DM	1206	0	1218	101	0
2	DO	1206	0	1218	91	0
2	DR	1206	0	1218	141	0
2	DT	1206	0	1218	113	0
2	DV	1206	0	1218	134	0
3	AE	1254	0	1250	124	0
3	BM	1254	0	1250	129	0
4	AK	1322	0	1311	131	0
4	BS	1322	0	1311	123	0
5	BA	546	0	568	78	0
5	BB	546	0	568	43	0
5	CJ	546	0	568	67	0
5	CK	546	0	568	36	0
5	DF	546	0	568	54	0
5	DX	546	0	568	48	0
6	BD	6761	0	6739	544	0
6	CM	6761	0	6739	548	0
7	BF	277	0	283	37	0
7	CO	277	0	283	37	0
8	BG	451	0	465	34	0
8	CP	451	0	465	26	0
8	DG	426	0	437	48	0
8	DH	426	0	437	45	0
8	DY	426	0	437	40	0
8	DZ	426	0	437	51	0
9	BH	2302	0	2310	202	0
9	CQ	2302	0	2310	199	0
9	DI	2280	0	2286	259	0
9	EA	2280	0	2286	222	0
10	AA	43	0	37	15	0
10	AB	43	0	37	14	0
10	AC	43	0	37	10	0
10	AD	43	0	37	14	0

Continued on next page...

Continued from previous page...

Mol	Chain	Non-H	H(model)	H(added)	Clashes	Symm-Clashes
10	AE	43	0	37	9	0
10	AF	43	0	37	10	0
10	AH	43	0	37	14	0
10	AI	43	0	37	11	0
10	AJ	43	0	37	7	0
10	AK	43	0	37	14	0
10	AL	43	0	37	10	0
10	AN	43	0	37	11	0
10	AO	43	0	37	7	0
10	AP	43	0	37	10	0
10	AQ	43	0	37	8	0
10	AR	43	0	38	9	0
10	AU	43	0	37	14	0
10	AV	43	0	37	17	0
10	AW	43	0	37	10	0
10	AX	43	0	37	9	0
10	AY	43	0	37	15	0
10	AZ	43	0	37	8	0
10	BD	86	0	74	23	0
10	BI	43	0	37	15	0
10	BJ	43	0	37	11	0
10	BK	43	0	37	16	0
10	BL	43	0	37	15	0
10	BM	43	0	37	11	0
10	BN	43	0	37	16	0
10	BP	43	0	37	14	0
10	BQ	43	0	37	15	0
10	BR	43	0	37	6	0
10	BS	43	0	37	11	0
10	BT	43	0	37	10	0
10	BV	43	0	37	14	0
10	BW	43	0	37	8	0
10	BX	43	0	37	10	0
10	BY	43	0	37	9	0
10	BZ	43	0	37	12	0
10	CC	43	0	37	7	0
10	CD	43	0	37	10	0
10	CE	43	0	37	12	0
10	CF	43	0	37	10	0
10	CG	43	0	37	11	0
10	CH	43	0	37	13	0
10	CM	86	0	74	23	0

Continued on next page...

Continued from previous page...

Mol	Chain	Non-H	H(model)	H(added)	Clashes	Symm-Clashes
10	CR	43	0	37	7	0
10	CS	43	0	37	7	0
10	CT	43	0	37	13	0
10	CU	43	0	37	8	0
10	CV	43	0	37	11	0
10	CW	43	0	37	12	0
10	CY	43	0	37	12	0
10	CZ	43	0	37	15	0
10	DA	43	0	37	10	0
10	DB	43	0	37	12	0
10	DC	43	0	37	14	0
10	DD	43	0	37	19	0
10	DJ	43	0	37	13	0
10	DK	43	0	37	8	0
10	DL	43	0	37	11	0
10	DM	43	0	37	8	0
10	DN	43	0	37	15	0
10	DO	43	0	37	9	0
10	DQ	43	0	37	12	0
10	DR	43	0	37	14	0
10	DS	43	0	37	10	0
10	DT	43	0	37	13	0
10	DU	43	0	37	12	0
10	DV	43	0	37	12	0
11	BH	42	52	52	4	0
11	CQ	42	52	52	6	0
11	DI	42	52	52	14	0
11	EA	42	52	52	12	0
All	All	117262	208	117450	8971	0

The all-atom clashscore is defined as the number of clashes found per 1000 atoms (including hydrogen atoms). The all-atom clashscore for this structure is 38.

All (8971) close contacts within the same asymmetric unit are listed below, sorted by their clash magnitude.

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
9:BH:214:ILE:HD13	9:BH:238:LEU:HD22	1.27	1.12
9:CQ:214:ILE:HD13	9:CQ:238:LEU:HD22	1.28	1.12
6:CM:700:ILE:HG21	2:DB:147:LYS:HE2	1.36	1.06
2:DV:41:ALA:HB2	2:DV:97:LEU:HD21	1.37	1.04
1:CR:50:ILE:HD13	1:CR:136:VAL:HG12	1.39	1.04

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
6:CM:85:LEU:HG	6:CM:153:LEU:HD21	1.38	1.04
5:BA:6:ILE:HG22	5:BA:53:VAL:HG23	1.40	1.03
9:DI:224:GLN:HB3	9:DI:302:ALA:HA	1.41	1.00
9:EA:224:GLN:HB3	9:EA:302:ALA:HA	1.41	1.00
9:CQ:191:GLU:HB3	9:CQ:254:ARG:HG3	1.42	1.00
2:CA:137:THR:HG21	2:CA:149:MET:HG2	1.44	0.99
4:AK:105:LEU:HD11	4:AK:160:MET:HE3	1.41	0.99
1:DJ:58:LEU:HD22	1:DJ:129:SER:HB2	1.45	0.99
2:CZ:66:THR:HA	2:CZ:72:MET:HB3	1.45	0.99
1:AX:107:ILE:HD11	6:BD:532:VAL:HG22	1.45	0.98
1:BP:105:GLU:HA	1:BP:109:LEU:HB2	1.41	0.98
1:DN:23:LEU:HD12	2:DO:38:VAL:HG13	1.43	0.98
1:CY:98:SER:HA	2:CZ:5:ILE:HD11	1.43	0.98
1:BV:8:ILE:HB	2:BW:1:MET:HE1	1.47	0.97
9:DI:252:PRO:HA	9:DI:271:GLY:HA3	1.43	0.97
1:DQ:95:GLY:HA3	1:DQ:104:ILE:HD11	1.45	0.97
1:CT:85:MET:HE2	1:CT:129:SER:HB2	1.45	0.97
1:BK:66:VAL:HA	1:BK:72:ALA:HB3	1.45	0.96
2:DK:119:LEU:HD13	10:DK:200:CYC:HBD1	1.46	0.96
6:CM:195:THR:HG22	10:CM:902:CYC:H2C	1.44	0.96
1:CR:37:LEU:HD13	2:CS:28:LYS:HE3	1.47	0.96
1:CR:58:LEU:HD22	1:CR:129:SER:HB2	1.47	0.96
6:BD:756:ILE:HG21	1:CY:159:LYS:HG3	1.45	0.96
6:BD:273:LEU:HD22	6:BD:277:GLU:HG3	1.47	0.95
5:DF:6:ILE:HG22	5:DF:53:VAL:HG23	1.48	0.95
1:BK:109:LEU:HD21	1:BK:159:LYS:HG3	1.47	0.95
2:CZ:130:ILE:HG23	2:CZ:153:LEU:HD12	1.46	0.95
9:DI:25:ILE:HD11	9:DI:153:VAL:HA	1.45	0.95
9:EA:292:LEU:HG	9:EA:298:ILE:H	1.30	0.95
2:AI:8:VAL:HG21	2:AI:27:LEU:HD11	1.45	0.95
9:EA:76:PRO:HB3	9:EA:123:ALA:HB2	1.49	0.95
1:CE:30:VAL:HG21	2:CF:34:GLY:HA3	1.47	0.95
1:CG:112:VAL:HG21	1:CG:160:MET:HE1	1.48	0.95
5:DX:6:ILE:HG22	5:DX:53:VAL:HG23	1.47	0.95
1:AA:105:GLU:HA	1:AA:109:LEU:HB3	1.49	0.94
1:AH:58:LEU:HD22	1:AH:129:SER:HB2	1.49	0.94
6:CM:273:LEU:HD22	6:CM:277:GLU:HG3	1.46	0.94
1:DU:58:LEU:HD22	1:DU:129:SER:HB2	1.50	0.94
10:BT:200:CYC:HC	10:BT:200:CYC:HMD1	1.30	0.94
1:BR:8:ILE:HD11	4:BS:95:TYR:HB3	1.47	0.94
1:DA:41:GLU:HG3	2:DB:24:MET:HE2	1.46	0.94

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:DK:64:ASP:HA	2:DK:67:ARG:HB2	1.49	0.94
6:BD:455:GLU:HG3	6:BD:664:GLY:HA3	1.48	0.93
1:BK:7:SER:HB3	1:BK:25:ARG:HH12	1.32	0.93
2:DM:60:LEU:HD13	2:DM:72:MET:HE1	1.48	0.93
1:BI:105:GLU:HA	1:BI:109:LEU:HB3	1.51	0.93
6:CM:69:SER:HA	6:CM:82:LEU:HD21	1.48	0.93
1:DJ:130:VAL:HG13	1:DJ:157:ILE:HG12	1.49	0.93
6:BD:244:PRO:HG2	6:BD:248:ILE:HD13	1.50	0.93
9:DI:204:ASN:HB3	9:DI:209:ASP:HB3	1.50	0.93
2:BN:104:LEU:HD22	2:BN:156:ILE:HD11	1.48	0.92
2:DK:51:ILE:HG23	2:DK:136:VAL:HB	1.51	0.92
1:DQ:23:LEU:HD22	2:DR:38:VAL:HG23	1.52	0.92
2:DB:66:THR:HG21	10:DC:200:CYC:HBA1	1.51	0.92
2:AY:51:ILE:HD11	2:AY:140:LEU:HD23	1.52	0.92
1:BP:58:LEU:HD22	1:BP:129:SER:HB2	1.50	0.92
9:EA:25:ILE:HD11	9:EA:153:VAL:HA	1.49	0.92
6:CM:177:VAL:HG21	6:CM:260:ALA:HB2	1.52	0.92
2:CW:41:ALA:HB2	2:CW:97:LEU:HD21	1.50	0.92
1:DQ:130:VAL:HG13	1:DQ:157:ILE:HG22	1.52	0.92
9:BH:89:ARG:HB2	9:BH:170:LYS:HE3	1.53	0.91
2:AW:108:VAL:HG11	2:AW:156:ILE:HD11	1.48	0.91
1:CY:27:LYS:HD3	2:CZ:38:VAL:HG11	1.50	0.91
4:AK:113:LEU:HD13	4:AK:164:LEU:HD22	1.51	0.91
6:CM:455:GLU:HG3	6:CM:664:GLY:HA3	1.53	0.91
1:BZ:101:VAL:HG23	1:BZ:104:ILE:HD12	1.51	0.91
8:DH:216:MET:HE3	8:DY:204:ILE:HD11	1.53	0.91
6:BD:701:ARG:HB3	6:BD:706:ILE:HD11	1.53	0.91
2:CF:57:ALA:HA	2:CF:61:LEU:HD12	1.49	0.91
1:CR:76:GLU:HB3	8:DZ:247:PRO:HG3	1.50	0.91
1:AC:66:VAL:HA	1:AC:72:ALA:HB3	1.53	0.91
9:DI:219:SER:HA	9:DI:234:LYS:HE2	1.52	0.91
2:DB:124:SER:HB2	2:DT:124:SER:HB2	1.53	0.90
2:AS:60:LEU:HD13	2:AS:72:MET:HE1	1.52	0.90
1:DJ:37:LEU:HD23	2:DK:28:LYS:HE2	1.49	0.90
1:DU:72:ALA:HA	1:DU:77:MET:HB3	1.53	0.90
2:CZ:119:LEU:HD13	10:CZ:200:CYC:HBD1	1.53	0.90
1:AX:105:GLU:HA	1:AX:109:LEU:HB3	1.51	0.90
1:DS:41:GLU:HG3	2:DT:24:MET:HE2	1.53	0.90
1:DJ:50:ILE:HD11	1:DJ:140:LEU:HD11	1.54	0.90
1:AC:7:SER:HB3	1:AC:25:ARG:HH12	1.37	0.90
6:BD:788:MET:HA	6:BD:792:TYR:HB3	1.53	0.90

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
9:BH:191:GLU:HB3	9:BH:254:ARG:HG3	1.54	0.90
2:BJ:57:ALA:HA	2:BJ:61:LEU:HD12	1.53	0.90
9:CQ:252:PRO:HA	9:CQ:271:GLY:HA3	1.52	0.90
1:AR:101:VAL:HG23	1:AR:104:ILE:HD12	1.52	0.90
2:CZ:36:LEU:HD11	2:CZ:145:ALA:HB2	1.54	0.90
1:CV:105:GLU:HA	1:CV:109:LEU:HB2	1.54	0.89
2:DD:112:LEU:HG	2:DD:160:LEU:HD13	1.54	0.89
3:AE:105:GLU:HA	3:AE:109:LEU:HB2	1.54	0.89
6:CM:806:LYS:HD2	6:CM:872:LEU:HB3	1.53	0.89
2:AQ:54:GLU:HG3	9:BH:63:LEU:HD11	1.53	0.89
9:DI:292:LEU:HG	9:DI:298:ILE:H	1.36	0.89
6:BD:69:SER:HA	6:BD:82:LEU:HD21	1.54	0.89
1:BV:105:GLU:HA	1:BV:109:LEU:HB2	1.55	0.89
3:AE:39:ILE:HG21	3:AE:96:VAL:HG11	1.51	0.89
9:CQ:198:VAL:HG12	9:CQ:202:MET:HE1	1.55	0.88
1:CY:109:LEU:HD21	1:CY:159:LYS:HB3	1.55	0.88
2:AW:67:ARG:HG3	2:AW:68:PRO:HD2	1.54	0.88
1:CY:77:MET:HE3	8:DH:201:ARG:HB2	1.52	0.88
9:DI:44:TYR:HA	9:DI:47:MET:HE3	1.55	0.88
1:CR:23:LEU:HD12	2:CS:38:VAL:HG13	1.53	0.88
1:CE:58:LEU:HD22	1:CE:129:SER:HB3	1.54	0.88
1:CY:26:ILE:HD13	2:CZ:97:LEU:HD21	1.53	0.88
6:BD:85:LEU:HG	6:BD:153:LEU:HD21	1.53	0.88
1:CR:105:GLU:HA	1:CR:109:LEU:HB2	1.54	0.88
2:AY:71:ASN:HD22	2:AY:122:PRO:HD3	1.39	0.88
6:BD:19:THR:HG22	6:BD:169:ALA:HB1	1.56	0.87
1:CR:131:ARG:HG2	1:CR:157:ILE:HD13	1.55	0.87
1:DC:72:ALA:HA	1:DC:77:MET:HB3	1.54	0.87
1:AJ:105:GLU:HA	1:AJ:109:LEU:HB2	1.55	0.87
9:CQ:238:LEU:HD12	9:CQ:242:ARG:HH21	1.38	0.87
1:AN:9:VAL:HG23	2:AO:1:MET:HE2	1.57	0.87
8:DG:221:MET:HE3	1:DQ:52:LYS:HG2	1.55	0.87
9:EA:21:VAL:HG23	9:EA:22:PRO:HD3	1.55	0.87
1:AV:30:VAL:HG21	2:AW:34:GLY:HA3	1.55	0.87
8:DZ:237:ALA:HA	8:DZ:240:ILE:HD13	1.55	0.87
6:BD:439:ARG:HG2	6:BD:440:PRO:HD2	1.56	0.87
1:DJ:5:THR:HG23	2:DK:1:MET:HE3	1.57	0.87
6:BD:801:ILE:HD12	6:BD:822:TYR:HB3	1.53	0.86
1:DN:141:MET:HE3	1:DN:146:ALA:HA	1.57	0.86
9:EA:219:SER:HA	9:EA:234:LYS:HE2	1.55	0.86
2:BY:54:GLU:HG3	9:CQ:63:LEU:HD11	1.53	0.86

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:CD:137:THR:HG21	2:CD:149:MET:HG2	1.55	0.86
1:DQ:105:GLU:HA	1:DQ:109:LEU:HB2	1.55	0.86
6:CM:439:ARG:HG2	6:CM:440:PRO:HD2	1.57	0.86
2:DM:104:LEU:HD13	2:DM:156:ILE:HG13	1.57	0.86
2:CH:15:GLN:HG3	2:CH:17:LYS:HE2	1.58	0.85
1:CE:113:ARG:HD3	1:CE:123:ILE:HD13	1.58	0.85
2:DK:72:MET:HE3	2:DK:81:CYS:HB3	1.59	0.85
2:CF:24:MET:HE3	2:CF:28:LYS:HD3	1.58	0.85
2:DK:51:ILE:HG21	2:DK:137:THR:HG23	1.56	0.85
2:CH:51:ILE:HD11	2:CH:140:LEU:HD23	1.59	0.85
1:DN:85:MET:HE3	1:DN:129:SER:HB2	1.57	0.85
7:BF:113:MET:HA	7:BF:116:GLN:HE21	1.42	0.85
1:CT:131:ARG:HG3	1:CT:157:ILE:HD12	1.59	0.85
2:AD:122:PRO:HG2	10:AD:200:CYC:HMC1	1.57	0.85
1:BK:113:ARG:HH22	1:BP:117:ARG:HD3	1.41	0.85
6:CM:240:VAL:HG22	6:CM:252:GLU:HG3	1.58	0.84
10:CM:901:CYC:HMA3	10:CM:901:CYC:HB	1.41	0.84
1:DJ:50:ILE:HG13	1:DJ:140:LEU:HD21	1.59	0.84
9:BH:252:PRO:HA	9:BH:271:GLY:HA3	1.60	0.84
3:BM:97:LEU:HD13	2:BN:19:LEU:HD12	1.59	0.84
1:CV:65:ILE:HD12	10:CV:200:CYC:HBC2	1.57	0.84
2:DK:85:LEU:HD22	2:DK:133:ILE:HD11	1.57	0.84
1:DQ:72:ALA:HA	1:DQ:77:MET:HB3	1.59	0.84
2:DR:65:VAL:HG12	2:DR:72:MET:HB2	1.58	0.84
1:AV:58:LEU:HD22	1:AV:129:SER:HB3	1.59	0.84
1:CR:8:ILE:HG21	2:CS:1:MET:HE1	1.57	0.84
1:DQ:8:ILE:HD11	2:DR:94:TYR:HB3	1.59	0.84
1:CR:91:LEU:HD11	1:CR:104:ILE:HA	1.59	0.84
1:CV:131:ARG:HG2	1:CV:157:ILE:HD13	1.57	0.84
9:DI:284:MET:HG2	9:DI:305:LEU:HD23	1.59	0.84
1:AZ:4:VAL:HG22	1:AZ:26:ILE:HD12	1.60	0.84
1:BZ:141:MET:HB2	1:BZ:146:ALA:HB2	1.60	0.84
1:DL:137:ALA:HB1	1:DL:141:MET:HE2	1.60	0.84
2:AQ:14:VAL:HG12	6:BD:683:VAL:HG11	1.59	0.84
1:BV:113:ARG:HG2	1:BV:123:ILE:HD13	1.58	0.84
6:CM:353:ARG:HG2	6:CM:399:ARG:HH21	1.43	0.84
9:DI:195:ASN:HB3	9:DI:198:VAL:HG12	1.60	0.84
6:BD:676:ARG:HB2	6:BD:679:ILE:HG12	1.60	0.84
2:BN:107:ARG:HE	5:CJ:45:LYS:HG2	1.43	0.83
1:CY:8:ILE:HD13	1:CY:18:LEU:HD21	1.59	0.83
6:BD:845:LEU:HG	5:DF:42:ARG:HD2	1.61	0.83

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:CF:41:ALA:HB2	2:CF:96:MET:HE1	1.60	0.83
1:CV:41:GLU:HG3	2:CW:24:MET:HE2	1.60	0.83
1:CR:95:GLY:HA3	1:CR:104:ILE:HD11	1.59	0.83
1:DC:105:GLU:HA	1:DC:109:LEU:HB2	1.59	0.83
9:DI:121:PHE:HA	9:DI:276:PRO:HG2	1.59	0.83
8:BG:237:ALA:HA	8:BG:240:ILE:HD12	1.58	0.83
1:BZ:58:LEU:HD22	1:BZ:129:SER:HB3	1.60	0.83
6:CM:756:ILE:HG21	1:DQ:159:LYS:HG3	1.59	0.83
1:DA:51:VAL:HG22	1:DA:133:MET:HE1	1.61	0.83
2:DT:66:THR:HG21	10:DU:200:CYC:HBA1	1.59	0.83
9:EA:195:ASN:HB3	9:EA:198:VAL:HG12	1.59	0.83
9:BH:248:LEU:HA	9:BH:275:THR:HG21	1.59	0.83
2:BL:66:THR:HG21	10:BM:200:CYC:HBA1	1.61	0.83
2:BW:85:LEU:HD22	2:BW:133:ILE:HD11	1.61	0.83
6:CM:5:ALA:HB2	6:CM:436:GLN:HG3	1.59	0.83
2:DM:65:VAL:HG12	2:DM:72:MET:HB2	1.61	0.83
1:CG:105:GLU:HA	1:CG:109:LEU:HB3	1.59	0.83
5:CJ:19:GLN:HB2	6:CM:239:LYS:HG2	1.61	0.83
6:CM:703:GLU:HA	6:CM:706:ILE:HD12	1.61	0.83
2:AI:77:ARG:HG2	10:AI:200:CYC:HMD1	1.61	0.83
1:AN:105:GLU:HA	1:AN:109:LEU:HB2	1.59	0.83
9:BH:50:THR:HG21	9:BH:143:GLN:HE21	1.44	0.83
2:CW:52:VAL:HA	2:CW:133:ILE:HD11	1.61	0.83
6:BD:75:ILE:HD12	6:BD:197:VAL:HG23	1.59	0.82
1:BI:65:ILE:HG22	1:BI:72:ALA:HB3	1.61	0.82
10:BM:200:CYC:HMD1	10:BM:200:CYC:HC	1.43	0.82
6:CM:75:ILE:HD12	6:CM:197:VAL:HG23	1.59	0.82
6:CM:676:ARG:HB2	6:CM:679:ILE:HG12	1.61	0.82
1:DN:131:ARG:HG2	1:DN:157:ILE:HD13	1.61	0.82
9:EA:202:MET:HE1	9:EA:269:VAL:HG21	1.60	0.82
1:DL:115:MET:HE1	10:DL:200:CYC:HMB2	1.60	0.82
9:EA:252:PRO:HA	9:EA:271:GLY:HA3	1.61	0.82
1:AC:130:VAL:HA	1:AC:133:MET:HE3	1.60	0.82
2:CS:119:LEU:HD13	10:CS:200:CYC:HBD1	1.61	0.82
9:EA:147:SER:HA	9:EA:150:GLN:HE21	1.43	0.82
1:AC:102:THR:HG23	6:BD:276:LEU:HD22	1.61	0.82
6:CM:872:LEU:HG	6:CM:884:VAL:HG11	1.61	0.82
8:DH:240:ILE:HD11	1:DJ:79:ALA:HB2	1.62	0.82
2:DD:57:ALA:HA	2:DD:61:LEU:HD12	1.59	0.82
2:DR:66:THR:HG22	2:DR:72:MET:HG2	1.59	0.82
2:AF:104:LEU:HD22	2:AF:156:ILE:HD11	1.61	0.82

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:CZ:130:ILE:HA	2:CZ:133:ILE:HD12	1.61	0.82
2:DK:104:LEU:HD21	2:DK:156:ILE:HG13	1.61	0.82
5:BA:11:PRO:HA	5:BA:25:THR:HG21	1.59	0.82
6:BD:240:VAL:HG22	6:BD:252:GLU:HG3	1.60	0.82
2:BN:74:THR:HG23	2:BN:77:ARG:HH21	1.45	0.82
2:BN:77:ARG:HG2	10:BN:200:CYC:HAD1	1.60	0.82
2:BT:85:LEU:HD22	2:BT:133:ILE:HD11	1.60	0.82
1:AR:58:LEU:HD22	1:AR:129:SER:HB3	1.61	0.82
3:BM:39:ILE:HG21	3:BM:96:VAL:HG11	1.60	0.82
1:AA:115:MET:HE1	10:AA:200:CYC:HMB3	1.62	0.82
2:AD:5:ILE:HG22	2:AD:27:LEU:HD22	1.59	0.82
1:AN:102:THR:HA	1:AN:105:GLU:HG2	1.62	0.82
6:CM:845:LEU:HG	5:DX:42:ARG:HD2	1.59	0.82
2:AU:110:ASN:HD21	5:BB:49:LYS:HA	1.44	0.81
1:AN:50:ILE:HA	1:AN:136:VAL:HG11	1.61	0.81
9:BH:204:ASN:HB3	9:BH:213:LEU:HB2	1.62	0.81
1:AA:58:LEU:HD22	1:AA:129:SER:HB3	1.59	0.81
5:BA:29:LYS:HZ1	6:BD:318:GLU:H	1.28	0.81
1:BZ:18:LEU:HD12	2:CA:97:LEU:HD13	1.61	0.81
2:CF:77:ARG:HG2	10:CF:200:CYC:HAD1	1.62	0.81
8:DG:242:ILE:HD12	1:DL:82:LEU:HB2	1.63	0.81
1:DN:58:LEU:HD22	1:DN:129:SER:HB3	1.61	0.81
4:AK:1:MET:HE2	4:AK:104:VAL:HB	1.62	0.81
4:AK:88:TYR:OH	6:BD:252:GLU:N	2.13	0.81
2:BJ:106:GLU:HB2	5:CJ:57:THR:HG23	1.60	0.81
2:AD:83:ARG:HD2	5:BA:22:LEU:HD11	1.61	0.81
2:AW:76:ARG:HB2	1:AX:110:VAL:HG13	1.63	0.81
9:CQ:248:LEU:HA	9:CQ:275:THR:HG21	1.62	0.81
1:CE:62:ARG:HH12	1:CE:124:GLU:HG2	1.45	0.81
2:CU:104:LEU:HD22	2:CU:156:ILE:HD11	1.62	0.81
1:AJ:115:MET:HE1	10:AJ:200:CYC:HMB3	1.61	0.81
4:AK:105:LEU:HD21	4:AK:160:MET:HG3	1.62	0.81
1:DJ:50:ILE:HA	1:DJ:136:VAL:HG11	1.62	0.81
2:AB:106:GLU:HB2	5:BA:57:THR:HG23	1.61	0.81
1:BK:2:SER:HB3	1:BK:5:THR:HG23	1.62	0.81
6:CM:75:ILE:HG13	6:CM:198:ALA:HB2	1.61	0.81
2:DM:51:ILE:HD11	2:DM:140:LEU:HD23	1.62	0.81
2:DV:108:VAL:HG12	2:DV:109:LEU:HD23	1.63	0.80
1:BR:8:ILE:HD13	4:BS:1:MET:HE3	1.63	0.80
1:CG:27:LYS:HZ3	2:CH:35:GLU:HA	1.44	0.80
5:CK:10:VAL:HG12	5:CK:48:GLY:HA3	1.63	0.80

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:AC:131:ARG:HG2	1:AC:157:ILE:HD13	1.63	0.80
1:AX:27:LYS:HZ3	2:AY:35:GLU:HA	1.47	0.80
6:BD:75:ILE:HG13	6:BD:198:ALA:HB2	1.62	0.80
2:BT:72:MET:HE3	2:BT:81:CYS:HB3	1.64	0.80
1:DA:8:ILE:HD13	1:DA:18:LEU:HD11	1.63	0.80
2:DD:122:PRO:HG2	10:DD:200:CYC:HMC3	1.61	0.80
9:DI:140:ALA:HA	9:DI:143:GLN:HE21	1.46	0.80
1:CE:97:VAL:HG21	2:CF:19:LEU:HD11	1.63	0.80
9:CQ:50:THR:HG21	9:CQ:143:GLN:HE21	1.46	0.80
2:DR:71:ASN:HD22	2:DR:122:PRO:HD3	1.45	0.80
3:AE:97:LEU:HD13	2:AF:19:LEU:HD12	1.61	0.80
2:CW:126:THR:HB	10:CW:200:CYC:HMC1	1.63	0.80
1:DQ:58:LEU:HD22	1:DQ:129:SER:HB3	1.61	0.80
1:AN:85:MET:SD	1:AN:129:SER:OG	2.40	0.80
2:CH:24:MET:HG3	2:CH:28:LYS:HE2	1.64	0.80
2:DV:44:ILE:HD11	2:DV:89:LEU:HD11	1.63	0.80
1:AC:2:SER:HB3	1:AC:5:THR:HG23	1.63	0.80
1:CY:37:LEU:HD22	2:CZ:28:LYS:HE3	1.63	0.80
4:AK:123:PRO:HB2	4:AK:126:PRO:HD2	1.62	0.80
2:AL:56:VAL:HG12	2:AL:61:LEU:HG	1.63	0.80
2:BJ:103:ILE:HD11	2:BJ:107:ARG:HD2	1.63	0.80
2:DV:58:LYS:HA	8:DZ:206:GLN:HG3	1.62	0.80
1:AR:18:LEU:HD12	2:AS:97:LEU:HD13	1.64	0.80
4:AK:64:PRO:HG2	7:BF:105:PRO:HB2	1.64	0.80
5:BB:38:ARG:HE	6:BD:641:ASP:HB3	1.46	0.80
1:BR:89:LEU:HB2	1:BR:133:MET:HE1	1.63	0.80
2:AW:8:VAL:HG13	2:AW:23:ALA:HB1	1.64	0.79
10:BT:200:CYC:HAA1	6:CM:424:PHE:HZ	1.48	0.79
1:DL:58:LEU:HD22	1:DL:129:SER:HB3	1.63	0.79
2:DT:57:ALA:HA	2:DT:61:LEU:HD12	1.63	0.79
1:AR:4:VAL:HG11	2:AS:98:ALA:HA	1.63	0.79
2:AW:57:ALA:HA	2:AW:61:LEU:HD12	1.62	0.79
2:CS:104:LEU:HD23	2:CS:155:TYR:HD2	1.47	0.79
3:AE:71:ASN:HD21	3:AE:121:VAL:HG22	1.47	0.79
2:AL:57:ALA:HA	2:AL:61:LEU:HD12	1.63	0.79
2:AQ:110:ASN:HD22	6:BD:668:ARG:HG2	1.47	0.79
6:BD:141:ILE:HG23	6:BD:149:MET:HE2	1.64	0.79
1:BZ:105:GLU:HA	1:BZ:109:LEU:HB2	1.63	0.79
3:AE:58:LEU:HD22	3:AE:129:ALA:HB2	1.64	0.79
1:BK:58:LEU:HD22	1:BK:129:SER:HB3	1.62	0.79
3:BM:151:PRO:HB2	1:BR:20:PRO:HB3	1.64	0.79

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:CZ:59:SER:HB3	2:CZ:128:GLN:HG2	1.63	0.79
2:DK:106:GLU:HG3	5:DX:57:THR:HG23	1.63	0.79
1:AA:85:MET:HE1	1:AA:129:SER:HB2	1.61	0.79
1:AZ:115:MET:HE1	10:AZ:200:CYC:HMB3	1.65	0.79
2:DK:104:LEU:HD23	2:DK:155:TYR:HD2	1.47	0.79
4:AK:129:ARG:HH21	4:AK:133:ILE:HG13	1.47	0.79
1:AH:105:GLU:HA	1:AH:109:LEU:HB3	1.64	0.79
1:DA:61:LYS:HE3	9:DI:56:PRO:HB2	1.64	0.79
1:BX:115:MET:HE1	10:BX:200:CYC:HMB2	1.63	0.79
1:CG:115:MET:HE1	10:CG:200:CYC:HMB2	1.64	0.79
9:EA:37:LEU:HD12	9:EA:40:ILE:HD11	1.65	0.79
1:BK:102:THR:HG23	6:CM:276:LEU:HD22	1.64	0.79
1:AC:18:LEU:HD12	2:AD:97:LEU:HD13	1.65	0.78
2:AS:130:ILE:HA	2:AS:133:ILE:HD12	1.64	0.78
2:AL:1:MET:HE1	6:BD:25:ILE:HB	1.63	0.78
1:CG:109:LEU:HD21	1:CG:159:LYS:HG3	1.65	0.78
1:AA:72:ALA:HA	1:AA:77:MET:HB3	1.64	0.78
8:DG:221:MET:HE1	1:DQ:51:VAL:HG13	1.65	0.78
1:DJ:131:ARG:HG2	1:DJ:157:ILE:HD13	1.65	0.78
5:DX:5:ARG:HG2	5:DX:56:ALA:HB2	1.66	0.78
2:AU:57:ALA:HA	2:AU:61:LEU:HB2	1.64	0.78
6:CM:751:LEU:HD13	6:CM:755:ILE:HG21	1.64	0.78
1:DL:50:ILE:HA	1:DL:136:VAL:HG11	1.65	0.78
1:AJ:109:LEU:HD13	1:AJ:159:LYS:HG3	1.63	0.78
1:AC:95:GLY:HA3	1:AC:104:ILE:HD11	1.63	0.78
1:AR:39:ILE:HD13	1:AR:149:ALA:HB2	1.64	0.78
3:BM:71:ASN:HD21	3:BM:121:VAL:HG22	1.48	0.78
1:AX:58:LEU:HD22	1:AX:129:SER:HB3	1.66	0.78
2:BT:106:GLU:HG3	2:BT:107:ARG:HG2	1.64	0.78
2:CF:108:VAL:HG11	2:CF:156:ILE:HD11	1.66	0.78
2:DV:51:ILE:HG22	2:DV:133:ILE:HD11	1.66	0.78
4:BS:135:LYS:HD3	4:BS:157:PHE:HB2	1.66	0.78
2:AS:92:ALA:HB1	2:AS:149:MET:HE1	1.66	0.78
1:BX:105:GLU:HA	1:BX:109:LEU:HB2	1.65	0.78
2:DD:15:GLN:HB3	2:DD:17:LYS:HE3	1.64	0.78
1:DS:107:ILE:HG22	10:DS:200:CYC:HBB1	1.65	0.78
2:AO:77:ARG:HG2	10:AO:200:CYC:HAD1	1.66	0.77
6:BD:756:ILE:HD12	1:CY:109:LEU:HG	1.65	0.77
2:BW:77:ARG:HG2	10:BW:200:CYC:HAD1	1.67	0.77
1:AZ:109:LEU:HG	1:AZ:159:LYS:HG2	1.66	0.77
6:BD:519:LEU:HG	6:BD:522:ALA:HB2	1.66	0.77

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:BZ:11:ALA:HB2	1:BZ:18:LEU:HD23	1.66	0.77
1:CE:72:ALA:HA	1:CE:77:MET:HB3	1.66	0.77
1:DA:109:LEU:HD13	1:DA:159:LYS:HG3	1.64	0.77
2:AS:15:GLN:HB3	2:AS:17:LYS:HE2	1.66	0.77
1:BK:20:PRO:HG2	6:CM:172:PRO:HD3	1.67	0.77
2:BT:1:MET:HE1	6:CM:25:ILE:HB	1.65	0.77
1:BX:89:LEU:HB2	1:BX:133:MET:HE1	1.66	0.77
9:CQ:47:MET:HE1	9:CQ:54:ALA:HB2	1.65	0.77
1:AR:105:GLU:HA	1:AR:109:LEU:HB2	1.66	0.77
2:BL:5:ILE:HG22	2:BL:27:LEU:HD22	1.66	0.77
2:CD:2:GLN:HE21	2:CD:7:ALA:HA	1.50	0.77
9:CQ:248:LEU:HD13	9:CQ:250:LEU:HD21	1.67	0.77
1:CT:82:LEU:HB2	8:DY:242:ILE:HD12	1.67	0.77
2:DD:126:THR:HG22	10:DD:200:CYC:H3C	1.65	0.77
2:DK:130:ILE:HA	2:DK:133:ILE:HD12	1.65	0.77
1:BK:130:VAL:HA	1:BK:133:MET:HE3	1.66	0.77
6:CM:316:MET:HE2	6:CM:395:VAL:HG22	1.65	0.77
2:CH:112:LEU:HD13	10:CH:200:CYC:HMB2	1.66	0.77
8:DH:242:ILE:HD11	1:DJ:82:LEU:HB3	1.66	0.77
9:DI:17:ALA:HB3	9:DI:20:VAL:HG13	1.66	0.77
1:DN:41:GLU:HG3	2:DO:24:MET:HE2	1.67	0.77
1:DS:39:ILE:HD12	1:DS:141:MET:HE3	1.66	0.77
1:AC:134:LYS:HE3	1:AC:150:SER:HB2	1.66	0.77
2:AS:51:ILE:HG23	2:AS:136:VAL:HG12	1.65	0.77
1:AX:115:MET:HE1	10:AX:200:CYC:HMB2	1.67	0.77
1:AP:105:GLU:HA	1:AP:109:LEU:HB2	1.65	0.76
9:BH:232:VAL:HG11	9:EA:314:ASN:HD22	1.50	0.76
6:CM:782:GLY:HA3	6:CM:835:ILE:HG21	1.67	0.76
2:DR:134:LYS:HG3	2:DR:150:GLY:HA2	1.67	0.76
1:CE:50:ILE:HD13	1:CE:136:VAL:HG12	1.65	0.76
2:DB:104:LEU:HD22	2:DB:156:ILE:HD11	1.67	0.76
2:AB:57:ALA:HA	2:AB:61:LEU:HD12	1.68	0.76
10:AL:200:CYC:HBC3	10:AL:200:CYC:HHD	1.68	0.76
9:EA:117:MET:HE3	9:EA:124:PRO:HA	1.65	0.76
2:AY:93:THR:HA	2:AY:149:MET:HE1	1.66	0.76
6:BD:597:TYR:HB3	6:BD:600:LYS:HD2	1.68	0.76
8:CP:232:ILE:HD12	8:CP:233:PRO:HD2	1.67	0.76
9:DI:117:MET:HE1	11:DI:400:45D:H273	1.66	0.76
2:CD:60:LEU:HB3	2:CD:72:MET:HE1	1.68	0.76
1:CY:85:MET:HE3	1:CY:129:SER:HB2	1.66	0.76
2:CZ:51:ILE:HA	2:CZ:136:VAL:HG11	1.67	0.76

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:DS:115:MET:HE1	10:DS:200:CYC:HMB2	1.68	0.76
2:AW:71:ASN:HD22	2:AW:122:PRO:HD3	1.49	0.76
8:CP:226:ASN:HD21	9:CQ:146:GLU:HG3	1.50	0.76
4:AK:123:PRO:HB3	2:AS:64:ASP:HB3	1.66	0.76
6:BD:549:GLN:O	6:BD:608:ARG:HD2	1.85	0.76
2:BT:57:ALA:HA	2:BT:61:LEU:HD12	1.66	0.76
1:CG:131:ARG:HG3	1:CG:157:ILE:HD12	1.67	0.76
2:CS:57:ALA:HA	2:CS:61:LEU:HD12	1.67	0.76
8:DH:216:MET:HE2	8:DH:216:MET:HA	1.66	0.76
1:DQ:58:LEU:HD21	10:DQ:200:CYC:HBC3	1.68	0.76
1:DQ:65:ILE:HG22	1:DQ:72:ALA:HB3	1.67	0.76
1:DU:105:GLU:HA	1:DU:109:LEU:HB2	1.67	0.76
4:AK:74:TYR:OH	6:BD:161:ARG:NH2	2.18	0.76
1:AN:123:ILE:HD12	1:AV:117:ARG:HH12	1.51	0.76
2:DO:85:LEU:HG	10:DO:200:CYC:HBC1	1.68	0.76
10:BT:200:CYC:HBC3	10:BT:200:CYC:HHD	1.68	0.76
1:CC:109:LEU:HG	1:CC:159:LYS:HG2	1.67	0.76
1:DC:85:MET:HB3	1:DC:133:MET:HE3	1.68	0.76
4:AK:72:ASN:HD21	4:AK:122:VAL:HG22	1.49	0.75
2:AW:36:LEU:HD23	2:AW:39:ARG:HH12	1.50	0.75
1:AX:109:LEU:HD21	1:AX:159:LYS:HG3	1.67	0.75
5:BB:3:MET:N	5:BB:57:THR:OG1	2.19	0.75
9:CQ:214:ILE:HD11	9:CQ:237:VAL:HB	1.67	0.75
9:CQ:292:LEU:HD23	9:CQ:298:ILE:HA	1.66	0.75
9:DI:142:ILE:HD11	9:DI:153:VAL:HG11	1.68	0.75
2:DV:57:ALA:HA	2:DV:61:LEU:HD12	1.68	0.75
2:AD:51:ILE:HD11	2:AD:140:LEU:HD23	1.67	0.75
2:AL:85:LEU:HD22	2:AL:133:ILE:HD11	1.69	0.75
1:CE:85:MET:HE3	1:CE:129:SER:HB2	1.68	0.75
2:DK:51:ILE:HA	2:DK:136:VAL:HG11	1.68	0.75
10:DQ:200:CYC:HBB2	2:DV:74:THR:HA	1.67	0.75
2:BN:114:GLU:HG3	5:CJ:53:VAL:HG22	1.68	0.75
6:CM:401:LEU:HD13	6:CM:411:TRP:HH2	1.52	0.75
1:CY:16:ARG:O	2:CZ:94:TYR:OH	2.04	0.75
1:AC:109:LEU:HD21	1:AC:159:LYS:HG3	1.69	0.75
1:BK:65:ILE:HG21	10:BK:200:CYC:HBC2	1.68	0.75
6:CM:269:MET:HE3	6:CM:309:VAL:HG12	1.69	0.75
1:CR:130:VAL:HG13	1:CR:157:ILE:HG12	1.68	0.75
1:DL:105:GLU:HA	1:DL:109:LEU:HB2	1.67	0.75
9:EA:121:PHE:HA	9:EA:276:PRO:HG2	1.67	0.75
1:AN:64:ASP:HA	1:AN:67:SER:HB2	1.67	0.75

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:AV:101:VAL:HA	1:AV:104:ILE:HD12	1.67	0.75
1:BV:50:ILE:HA	1:BV:136:VAL:HG11	1.68	0.75
10:CD:200:CYC:HB	10:CD:200:CYC:HMA1	1.52	0.75
1:CG:6:LYS:HG2	6:CM:534:PHE:HZ	1.52	0.75
2:DV:60:LEU:HB3	2:DV:72:MET:HE1	1.67	0.75
2:AS:46:ALA:HB2	1:AV:154:ASP:HB3	1.68	0.75
2:AS:123:ILE:HA	2:AS:126:THR:HG22	1.68	0.75
1:BV:72:ALA:HA	1:BV:77:MET:HB3	1.69	0.75
1:CV:16:ARG:O	2:CW:94:TYR:OH	2.04	0.75
6:CM:19:THR:HG22	6:CM:169:ALA:HB1	1.68	0.75
6:CM:199:ILE:HD13	6:CM:227:LEU:HD21	1.69	0.75
6:CM:597:TYR:HB3	6:CM:600:LYS:HD2	1.69	0.75
1:AA:81:CYS:HA	10:AA:200:CYC:HHD	1.68	0.74
1:AC:141:MET:HG3	1:AC:146:ALA:HB2	1.68	0.74
6:BD:186:ILE:HG13	6:BD:190:CYS:HB3	1.67	0.74
3:BM:115:MET:HE2	10:BM:200:CYC:HB	1.50	0.74
2:DM:122:PRO:HG2	10:DM:200:CYC:HMC3	1.69	0.74
1:DU:85:MET:HE3	1:DU:133:MET:HE2	1.68	0.74
2:DV:81:CYS:HA	10:DV:200:CYC:HHD	1.68	0.74
5:DX:40:GLN:HG2	5:DX:44:MET:HE3	1.69	0.74
1:CG:58:LEU:HD22	1:CG:129:SER:HB3	1.69	0.74
6:CM:84:TYR:O	6:CM:150:GLN:NE2	2.20	0.74
9:BH:76:PRO:HB3	9:BH:123:ALA:HB2	1.70	0.74
6:BD:353:ARG:HG2	6:BD:399:ARG:HH21	1.50	0.74
1:BP:89:LEU:HB2	1:BP:133:MET:HE1	1.68	0.74
1:DS:105:GLU:HA	1:DS:109:LEU:HB2	1.68	0.74
1:AX:131:ARG:HG3	1:AX:157:ILE:HD12	1.68	0.74
2:BL:51:ILE:HD11	2:BL:140:LEU:HD23	1.69	0.74
9:DI:80:THR:HA	9:DI:83:MET:HE3	1.70	0.74
3:AE:20:SER:HB3	1:AJ:151:ALA:HB1	1.69	0.74
6:BD:726:LEU:HD11	6:BD:735:VAL:HG22	1.69	0.74
1:CY:122:PRO:HG2	1:CY:125:ALA:HB3	1.68	0.74
2:DK:104:LEU:HD11	2:DK:156:ILE:HD11	1.68	0.74
1:DQ:84:ASP:OD2	10:DQ:200:CYC:ND	2.21	0.74
6:BD:52:LEU:HD21	6:BD:216:ALA:HA	1.70	0.74
2:BY:14:VAL:HG12	6:CM:683:VAL:HG11	1.68	0.74
1:DN:51:VAL:HG12	1:DN:82:LEU:HD12	1.69	0.74
1:AC:105:GLU:HG2	1:AC:109:LEU:HD23	1.69	0.74
2:CW:71:ASN:HD21	2:CW:121:VAL:HG22	1.52	0.74
1:DA:84:ASP:OD2	10:DA:200:CYC:ND	2.21	0.74
5:CJ:19:GLN:NE2	6:CM:237:ALA:O	2.21	0.74

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:DD:51:ILE:HD13	2:DD:137:THR:HG22	1.70	0.74
2:CA:46:ALA:HB2	1:CE:154:ASP:HB3	1.70	0.73
2:CF:84:ASP:OD2	2:CF:116:TYR:OH	2.06	0.73
2:AB:104:LEU:HD22	2:AB:156:ILE:HD11	1.70	0.73
2:AF:74:THR:HG23	2:AF:77:ARG:HH21	1.52	0.73
6:BD:18:GLN:HB2	6:BD:173:ASN:HD22	1.51	0.73
9:BH:257:THR:HG23	9:BH:267:ILE:HG12	1.71	0.73
1:CG:36:ARG:NH1	1:CG:96:VAL:O	2.21	0.73
2:DB:112:LEU:HD13	10:DB:200:CYC:HMB3	1.70	0.73
1:DS:23:LEU:HD12	2:DT:38:VAL:HG13	1.69	0.73
2:AF:109:LEU:HD13	2:AF:159:GLY:HA3	1.69	0.73
2:AF:122:PRO:HB2	2:AF:125:SER:HB2	1.71	0.73
2:AI:77:ARG:HG3	10:AI:200:CYC:HAD1	1.71	0.73
1:AP:115:MET:HE1	10:AP:200:CYC:HMB3	1.70	0.73
1:AZ:81:CYS:HA	10:AZ:200:CYC:HHD	1.70	0.73
1:BP:115:MET:HE1	10:BP:200:CYC:HMB2	1.68	0.73
10:CW:200:CYC:HB	10:CW:200:CYC:HMA3	1.52	0.73
1:CY:83:ARG:NH1	1:CY:84:ASP:OD1	2.21	0.73
1:CG:97:VAL:HG21	2:CH:19:LEU:HD12	1.71	0.73
5:CJ:39:GLU:OE2	5:CJ:42:ARG:NH2	2.22	0.73
1:AH:16:ARG:O	2:AI:94:TYR:OH	2.07	0.73
8:BG:232:ILE:HD12	8:BG:233:PRO:HD2	1.70	0.73
10:BP:200:CYC:HBC3	10:BP:200:CYC:HHD	1.70	0.73
1:DJ:4:VAL:HG12	1:DJ:26:ILE:HD12	1.70	0.73
9:EA:74:MET:HE3	9:EA:78:GLN:HB2	1.69	0.73
6:BD:195:THR:HG22	10:BD:902:CYC:H2C	1.71	0.73
1:BK:131:ARG:HG2	1:BK:157:ILE:HD13	1.70	0.73
4:BS:44:ILE:HG22	4:BS:138:ILE:HD12	1.69	0.73
2:BT:88:TYR:OH	2:BT:116:TYR:OH	2.05	0.73
1:BZ:16:ARG:O	2:CA:94:TYR:OH	2.06	0.73
2:CS:1:MET:HG3	2:CS:103:ILE:HB	1.70	0.73
8:DG:210:LEU:HD13	8:DG:216:MET:HB3	1.70	0.73
3:AE:30:LEU:HD13	2:AF:31:PHE:HD1	1.54	0.73
10:AH:200:CYC:HBC3	10:AH:200:CYC:HHD	1.70	0.73
1:BV:106:GLU:OE1	6:CM:478:LYS:NZ	2.20	0.73
10:CE:200:CYC:HC	10:CE:200:CYC:HMD1	1.53	0.73
2:DK:105:ASP:HA	2:DK:109:LEU:HB2	1.70	0.73
1:AA:109:LEU:HG	1:AA:159:LYS:HG2	1.71	0.73
2:AD:40:ALA:HB3	2:AD:96:MET:HE1	1.70	0.73
5:BA:29:LYS:HE3	6:BD:318:GLU:HG2	1.68	0.73
2:BL:131:GLN:NE2	2:BL:135:GLU:OE2	2.22	0.73

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:CD:65:VAL:HG12	2:CD:72:MET:HE2	1.69	0.73
7:CO:114:ALA:HA	7:CO:117:VAL:HG12	1.70	0.73
1:CR:19:SER:OG	1:CR:22:GLU:OE1	2.05	0.73
2:CU:65:VAL:HG12	2:CU:72:MET:HB2	1.69	0.73
2:CW:71:ASN:O	2:CW:77:ARG:NE	2.22	0.73
9:EA:260:ALA:N	9:EA:264:PHE:O	2.19	0.73
9:EA:288:TRP:HD1	9:EA:303:ILE:HG12	1.54	0.73
2:AU:65:VAL:HG12	2:AU:72:MET:HE2	1.69	0.73
1:BI:5:THR:OG1	2:BJ:1:MET:SD	2.45	0.73
1:CY:50:ILE:HD13	1:CY:140:LEU:HD21	1.69	0.73
1:DA:50:ILE:HD13	1:DA:140:LEU:HD21	1.71	0.73
1:BR:26:ILE:HG22	4:BS:38:ILE:HD11	1.71	0.73
1:DC:109:LEU:HD13	1:DC:159:LYS:HG2	1.71	0.73
1:DL:141:MET:HE1	1:DL:149:ALA:HB3	1.70	0.73
6:CM:740:ARG:NH2	6:CM:750:ASP:OD2	2.22	0.72
9:CQ:204:ASN:HB3	9:CQ:213:LEU:HB2	1.71	0.72
1:DA:61:LYS:HA	9:DI:56:PRO:HD2	1.71	0.72
2:DV:65:VAL:HG12	2:DV:72:MET:HB2	1.70	0.72
2:AL:106:GLU:O	2:AL:110:ASN:ND2	2.21	0.72
2:AQ:117:ASN:O	6:BD:467:ARG:NH2	2.22	0.72
6:BD:236:PRO:HB2	6:BD:255:GLN:HB3	1.69	0.72
9:DI:208:ASN:HD21	9:DI:247:ASN:HA	1.54	0.72
1:AH:76:GLU:HG2	1:AH:77:MET:HE2	1.71	0.72
2:BW:30:TYR:OH	2:BW:97:LEU:O	2.07	0.72
9:DI:76:PRO:HB3	9:DI:123:ALA:HB2	1.71	0.72
1:BV:102:THR:HA	1:BV:105:GLU:HG2	1.70	0.72
2:CA:85:LEU:HD22	2:CA:133:ILE:HD11	1.72	0.72
5:CK:40:GLN:HG2	5:CK:44:MET:HE3	1.70	0.72
1:DJ:16:ARG:O	2:DK:94:TYR:OH	2.08	0.72
1:DJ:84:ASP:OD2	10:DJ:200:CYC:ND	2.22	0.72
6:BD:72:ALA:HB2	6:BD:156:MET:HE1	1.72	0.72
9:BH:214:ILE:HD11	9:BH:237:VAL:HB	1.71	0.72
2:CS:77:ARG:HB2	10:CS:200:CYC:HMD1	1.72	0.72
8:DH:227:PRO:HB2	1:DS:66:VAL:HG21	1.69	0.72
1:DU:16:ARG:O	2:DV:94:TYR:OH	2.08	0.72
1:AR:76:GLU:HG3	9:BH:11:ILE:HG21	1.72	0.72
2:AW:126:THR:HG22	10:AW:200:CYC:HMC1	1.72	0.72
1:BR:115:MET:HE1	10:BR:200:CYC:HMB3	1.70	0.72
9:CQ:224:GLN:NE2	9:CQ:228:GLN:O	2.22	0.72
2:DO:71:ASN:HD21	2:DO:121:VAL:HG22	1.53	0.72
2:AF:85:LEU:HG	10:AF:200:CYC:HBC1	1.72	0.72

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:AY:81:CYS:HA	10:AY:200:CYC:HHD	1.69	0.72
1:BK:73:TYR:H	1:BK:77:MET:HB2	1.55	0.72
3:BM:94:TYR:OH	2:BN:17:LYS:O	2.06	0.72
1:BZ:65:ILE:HG22	1:BZ:72:ALA:HB3	1.72	0.72
1:BZ:86:ASP:OD1	2:CA:18:TYR:OH	2.07	0.72
2:AD:76:ARG:NH1	3:AE:106:THR:O	2.23	0.72
2:AL:106:GLU:HG3	2:AL:107:ARG:HG2	1.70	0.72
2:BT:106:GLU:O	2:BT:110:ASN:ND2	2.21	0.72
6:CM:183:LYS:HG2	6:CM:192:ILE:HD12	1.70	0.72
1:AJ:9:VAL:HG11	6:BD:246:ASN:HB3	1.70	0.72
2:BJ:104:LEU:HD22	2:BJ:156:ILE:HD11	1.71	0.72
1:BV:39:ILE:HG12	1:BV:145:ASP:HB2	1.70	0.72
1:DU:109:LEU:HD13	1:DU:159:LYS:HG3	1.72	0.72
1:AN:8:ILE:HB	2:AO:1:MET:HE1	1.70	0.72
1:AX:85:MET:HB3	1:AX:133:MET:HE3	1.72	0.72
2:CD:134:LYS:HG3	2:CD:150:GLY:HA2	1.72	0.72
2:CH:81:CYS:HA	10:CH:200:CYC:HHD	1.71	0.72
6:BD:154:ARG:NH1	6:BD:155:ASP:OD1	2.23	0.71
1:CY:84:ASP:OD2	10:CY:200:CYC:ND	2.22	0.71
9:DI:32:ASN:HB3	9:DI:171:ARG:HH21	1.53	0.71
2:AY:24:MET:HG3	2:AY:28:LYS:HE2	1.71	0.71
2:BQ:60:LEU:O	2:BQ:63:SER:OG	2.08	0.71
1:BR:105:GLU:HA	1:BR:109:LEU:HB2	1.71	0.71
2:CF:103:ILE:HD11	2:CF:107:ARG:HD2	1.72	0.71
1:CG:85:MET:HB3	1:CG:133:MET:HE3	1.72	0.71
9:CQ:257:THR:HG23	9:CQ:267:ILE:HG12	1.72	0.71
2:DO:65:VAL:HG12	2:DO:72:MET:HB2	1.70	0.71
6:BD:292:ARG:NH2	6:BD:332:GLN:OE1	2.22	0.71
6:BD:749:ARG:NH1	10:DD:200:CYC:O2A	2.23	0.71
9:BH:206:ASN:O	9:BH:249:LYS:NZ	2.24	0.71
2:CZ:64:ASP:HA	2:CZ:67:ARG:HB2	1.71	0.71
2:DO:57:ALA:HA	2:DO:61:LEU:HD12	1.72	0.71
1:BK:16:ARG:O	2:BL:94:TYR:OH	2.08	0.71
2:BQ:75:THR:HG21	1:BR:112:VAL:HG12	1.72	0.71
4:BS:92:TYR:OH	6:CM:249:GLN:OE1	2.07	0.71
1:CY:82:LEU:HD11	8:DY:221:MET:HE1	1.71	0.71
1:DA:85:MET:HE2	1:DA:129:SER:HB2	1.71	0.71
2:DD:65:VAL:HG12	2:DD:72:MET:HB2	1.72	0.71
1:DJ:11:ALA:HB2	1:DJ:18:LEU:HD23	1.71	0.71
2:CU:15:GLN:HB3	2:CU:17:LYS:HE3	1.72	0.71
9:DI:85:ASP:HB3	9:DI:90:ALA:HB3	1.73	0.71

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:AR:100:ASP:OD2	1:AR:102:THR:OG1	2.06	0.71
1:AV:113:ARG:HD3	1:AV:123:ILE:HD13	1.72	0.71
2:CZ:51:ILE:HG21	2:CZ:137:THR:HG23	1.72	0.71
1:AV:72:ALA:HA	1:AV:77:MET:HB3	1.73	0.71
1:BI:85:MET:HB3	1:BI:133:MET:HE3	1.73	0.71
2:BN:57:ALA:HA	2:BN:61:LEU:HD12	1.72	0.71
1:BZ:128:GLN:OE1	1:BZ:131:ARG:NH1	2.24	0.71
2:CD:119:LEU:HD21	5:CK:33:TYR:HE2	1.56	0.71
1:CR:36:ARG:NH1	1:CR:96:VAL:O	2.22	0.71
2:AO:114:GLU:OE2	6:BD:443:ASP:N	2.23	0.71
2:AU:64:ASP:OD1	2:AU:67:ARG:NH1	2.23	0.71
2:AY:41:ALA:HB2	2:AY:96:MET:HE1	1.71	0.71
9:BH:183:ALA:N	9:BH:186:THR:HG1	1.88	0.71
1:BI:109:LEU:HG	1:BI:159:LYS:HG2	1.73	0.71
1:BI:115:MET:HE1	10:BI:200:CYC:HMB2	1.73	0.71
2:BJ:81:CYS:HA	10:BJ:200:CYC:HHD	1.73	0.71
3:BM:58:LEU:HD22	3:BM:129:ALA:HB2	1.72	0.71
1:BR:16:ARG:O	4:BS:95:TYR:OH	2.07	0.71
1:DN:12:ASP:OD1	2:DO:90:ARG:NH2	2.23	0.71
1:AC:113:ARG:NH2	1:AC:161:SER:OXT	2.24	0.71
1:AR:65:ILE:HG22	1:AR:72:ALA:HB3	1.71	0.71
6:BD:875:LYS:NZ	6:BD:882:GLU:OE2	2.24	0.71
9:BH:47:MET:HE1	9:BH:54:ALA:HB2	1.72	0.71
9:BH:224:GLN:NE2	9:BH:228:GLN:O	2.24	0.71
6:CM:292:ARG:NH2	6:CM:332:GLN:OE1	2.23	0.71
2:CS:62:TYR:OH	10:CT:200:CYC:O2D	2.08	0.71
1:AA:154:ASP:HA	1:AA:157:ILE:HG12	1.73	0.71
5:BB:10:VAL:HG12	5:BB:48:GLY:HA3	1.73	0.71
1:CR:7:SER:OG	1:CR:25:ARG:NH1	2.22	0.71
1:DA:52:LYS:HE2	8:DZ:221:MET:HG3	1.71	0.71
1:DJ:71:ASN:ND2	1:DJ:121:THR:OG1	2.24	0.71
1:AH:47:ARG:NH2	1:AH:86:ASP:OD2	2.24	0.70
9:BH:280:GLY:HA3	9:BH:284:MET:HG2	1.73	0.70
2:CD:110:ASN:HD21	5:CK:49:LYS:HA	1.55	0.70
6:CM:14:PRO:HG3	6:CM:401:LEU:HD21	1.72	0.70
2:CZ:71:ASN:HD22	2:CZ:122:PRO:HD3	1.56	0.70
3:AE:115:MET:HE2	10:AE:200:CYC:HB	1.55	0.70
10:AL:200:CYC:HAA1	6:BD:424:PHE:HZ	1.55	0.70
5:BB:40:GLN:HG2	5:BB:44:MET:HE3	1.73	0.70
6:BD:278:LYS:HD3	6:BD:310:ARG:HG3	1.73	0.70
6:BD:413:MET:HE2	6:BD:433:THR:HB	1.71	0.70

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
6:BD:714:VAL:N	10:DB:200:CYC:O1D	2.23	0.70
1:DJ:91:LEU:HD21	1:DJ:107:ILE:HG22	1.73	0.70
1:DN:128:GLN:OE1	1:DN:131:ARG:NH1	2.24	0.70
1:AJ:12:ASP:OD1	4:AK:91:ARG:NH1	2.23	0.70
2:AW:21:GLY:HA2	2:AW:24:MET:HB3	1.72	0.70
2:BT:13:ASP:OD1	6:CM:161:ARG:NH2	2.25	0.70
2:CZ:51:ILE:HG23	2:CZ:136:VAL:HB	1.73	0.70
1:DL:65:ILE:O	1:DL:72:ALA:N	2.22	0.70
2:AD:107:ARG:HD2	6:BD:270:LYS:HE3	1.74	0.70
2:AI:61:LEU:O	6:BD:682:ARG:NH1	2.23	0.70
2:AY:73:TYR:OH	1:AZ:90:ARG:NH2	2.24	0.70
2:BY:60:LEU:HB3	2:BY:72:MET:HE1	1.71	0.70
1:CE:131:ARG:HG2	1:CE:157:ILE:HD13	1.73	0.70
2:CF:76:ARG:HB2	1:CG:110:VAL:HG13	1.74	0.70
10:CH:200:CYC:O1A	5:CK:17:ARG:NH2	2.24	0.70
1:CR:3:ILE:HD12	1:CR:25:ARG:HG2	1.72	0.70
1:CR:73:TYR:HH	9:DI:147:SER:HG	1.33	0.70
1:DQ:59:PHE:HE1	10:DQ:200:CYC:HBC1	1.56	0.70
2:CZ:35:GLU:HA	2:CZ:38:VAL:HG12	1.74	0.70
1:DA:97:VAL:HG11	2:DB:19:LEU:HD13	1.73	0.70
2:DK:71:ASN:HD22	2:DK:121:VAL:HA	1.57	0.70
1:AC:73:TYR:H	1:AC:77:MET:HB2	1.54	0.70
2:AW:84:ASP:OD2	2:AW:116:TYR:OH	2.08	0.70
1:AX:36:ARG:NH1	1:AX:96:VAL:O	2.24	0.70
1:BK:134:LYS:HE3	1:BK:150:SER:HB2	1.73	0.70
3:BM:69:GLY:N	3:BM:73:TYR:OH	2.20	0.70
1:DC:16:ARG:O	2:DD:94:TYR:OH	2.10	0.70
2:DV:106:GLU:HG3	2:DV:107:ARG:HG3	1.71	0.70
2:AL:13:ASP:OD1	6:BD:161:ARG:NH2	2.24	0.70
1:BK:39:ILE:HD12	1:BK:145:ASP:OD1	1.92	0.70
1:CC:16:ARG:O	2:CD:94:TYR:OH	2.10	0.70
9:DI:45:LEU:O	9:DI:49:LYS:NZ	2.24	0.70
9:EA:208:ASN:HD21	9:EA:247:ASN:HA	1.56	0.70
2:AW:103:ILE:HD11	2:AW:107:ARG:HD2	1.72	0.70
2:BN:71:ASN:HD22	2:BN:122:PRO:HD3	1.56	0.70
2:BQ:8:VAL:HG21	2:BQ:27:LEU:HD11	1.72	0.70
5:BA:2:ARG:NH2	5:BA:34:ASP:OD1	2.21	0.69
1:CC:6:LYS:NZ	1:CC:100:ASP:OD2	2.24	0.69
1:CC:37:LEU:HD13	2:CD:28:LYS:HD3	1.74	0.69
2:CZ:3:ASP:HB3	2:CZ:98:ALA:HB1	1.74	0.69
1:DS:72:ALA:HA	1:DS:77:MET:HB3	1.73	0.69

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:AC:58:LEU:HD22	1:AC:129:SER:HB3	1.73	0.69
1:AC:81:CYS:HA	10:AC:200:CYC:HHD	1.74	0.69
2:AU:57:ALA:HB2	2:AU:61:LEU:HD12	1.75	0.69
2:AU:75:THR:HG23	1:AV:108:GLY:HA2	1.74	0.69
2:AU:94:TYR:OH	1:AZ:16:ARG:O	2.09	0.69
6:BD:447:TYR:OH	10:BD:901:CYC:O2A	2.09	0.69
1:BR:29:PHE:HA	1:BR:36:ARG:HH21	1.58	0.69
2:CA:54:GLU:O	2:CA:58:LYS:HG2	1.92	0.69
2:CW:81:CYS:HA	10:CW:200:CYC:HHD	1.74	0.69
2:DV:112:LEU:HD13	10:DV:200:CYC:HMB3	1.75	0.69
1:AA:85:MET:HB3	1:AA:133:MET:HE3	1.74	0.69
1:AJ:128:GLN:OE1	1:AJ:131:ARG:NH1	2.24	0.69
2:AU:107:ARG:NH1	1:AZ:12:ASP:OD2	2.25	0.69
9:BH:204:ASN:HB2	9:BH:213:LEU:HD13	1.74	0.69
10:BM:200:CYC:HB	10:BM:200:CYC:HMA3	1.56	0.69
1:BZ:151:ALA:HB1	1:CE:20:PRO:HB3	1.74	0.69
9:EA:284:MET:HG2	9:EA:305:LEU:HD23	1.74	0.69
2:BT:83:ARG:NH2	2:BT:84:ASP:OD1	2.26	0.69
1:BV:51:VAL:HG12	1:BV:85:MET:HG3	1.74	0.69
1:CC:128:GLN:NE2	1:CC:132:GLU:OE2	2.26	0.69
9:DI:235:GLU:OE1	9:DI:239:ARG:NH1	2.26	0.69
9:DI:269:VAL:HG23	9:DI:288:TRP:HE3	1.57	0.69
1:DJ:38:ARG:NH1	1:DJ:145:ASP:OD2	2.22	0.69
2:DK:108:VAL:HG23	10:DK:200:CYC:HAB1	1.74	0.69
2:DR:40:ALA:HB2	2:DR:145:ALA:HB1	1.73	0.69
3:AE:85:TYR:HE1	3:AE:129:ALA:HB1	1.58	0.69
1:AR:25:ARG:HG3	1:AV:25:ARG:HD3	1.73	0.69
2:AU:137:THR:HG21	2:AU:149:MET:HG2	1.74	0.69
2:CD:44:ILE:HD11	2:CD:137:THR:HG23	1.74	0.69
1:DJ:113:ARG:NH2	1:DJ:161:SER:O	2.25	0.69
1:DS:14:GLU:OE1	1:DS:16:ARG:NE	2.26	0.69
2:AD:131:GLN:NE2	2:AD:135:GLU:OE2	2.25	0.69
1:AN:81:CYS:HA	10:AN:200:CYC:HHD	1.73	0.69
4:BS:51:ILE:HG23	4:BS:137:MET:HE2	1.73	0.69
2:DB:71:ASN:HD22	2:DB:122:PRO:HD3	1.57	0.69
8:DH:238:GLN:HA	1:DJ:52:LYS:HE3	1.74	0.69
2:DM:73:TYR:OH	1:DN:90:ARG:NH2	2.25	0.69
3:AE:37:ILE:HG23	2:AF:24:MET:HE2	1.75	0.69
1:BK:117:ARG:HD2	1:BP:161:SER:HB2	1.74	0.69
1:BP:76:GLU:HG2	1:BP:77:MET:HE2	1.74	0.69
2:CD:75:THR:HG23	1:CE:108:GLY:HA2	1.75	0.69

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
6:CM:714:VAL:N	10:DT:200:CYC:O1D	2.23	0.69
1:CY:76:GLU:HA	8:DH:203:PHE:CZ	2.28	0.69
2:CZ:84:ASP:OD2	2:CZ:116:TYR:OH	2.08	0.69
2:AB:112:LEU:HD13	10:AB:200:CYC:HMB2	1.73	0.69
2:AL:148:GLU:OE2	7:BF:89:ALA:N	2.23	0.69
10:AP:200:CYC:HMD1	10:AP:200:CYC:HC	1.58	0.69
2:AW:112:LEU:HD11	10:AW:200:CYC:HMB3	1.74	0.69
9:BH:289:ARG:NH2	9:EA:307:ALA:O	2.26	0.69
2:BJ:112:LEU:HD13	10:BJ:200:CYC:HMB2	1.73	0.69
3:BM:110:ILE:HG22	5:CJ:16:ILE:HD11	1.74	0.69
2:BW:114:GLU:OE2	6:CM:443:ASP:N	2.25	0.69
1:BZ:72:ALA:HA	1:BZ:77:MET:HB3	1.75	0.69
1:CC:4:VAL:HG22	1:CC:26:ILE:HD12	1.75	0.69
2:CD:85:LEU:HG	10:CD:200:CYC:HBC1	1.73	0.69
1:DJ:12:ASP:OD2	2:DK:107:ARG:NE	2.26	0.69
1:DN:16:ARG:O	2:DO:94:TYR:OH	2.10	0.69
1:DQ:94:TYR:HD1	2:DR:9:ILE:HD12	1.58	0.69
9:EA:235:GLU:OE1	9:EA:239:ARG:NH1	2.26	0.69
1:AV:98:SER:HA	2:AW:5:ILE:HG21	1.74	0.69
2:BT:16:GLY:O	6:CM:161:ARG:NH1	2.25	0.69
2:CD:57:ALA:HB2	2:CD:61:LEU:HD12	1.75	0.69
2:CW:85:LEU:HG	10:CW:200:CYC:HBC1	1.73	0.69
1:CY:83:ARG:NH2	10:CY:200:CYC:O2A	2.25	0.69
10:CZ:200:CYC:HMD1	10:CZ:200:CYC:HC	1.56	0.69
1:DJ:105:GLU:HA	1:DJ:109:LEU:HB2	1.75	0.69
2:AO:158:SER:O	2:AO:161:SER:OG	2.06	0.69
2:AU:73:TYR:OH	1:AV:90:ARG:NH2	2.25	0.69
1:AX:85:MET:HE1	1:AX:129:SER:HB2	1.75	0.69
2:AY:83:ARG:NH2	10:AY:200:CYC:O2A	2.26	0.69
1:BP:65:ILE:HG22	1:BP:72:ALA:HB3	1.74	0.69
2:AQ:57:ALA:HA	2:AQ:61:LEU:HD12	1.74	0.68
8:BG:221:MET:O	8:BG:225:ILE:HG22	1.93	0.68
10:BV:200:CYC:O1A	2:CA:62:TYR:N	2.25	0.68
1:DS:95:GLY:HA3	1:DS:104:ILE:HD11	1.75	0.68
9:EA:47:MET:HE1	9:EA:54:ALA:HB2	1.74	0.68
3:AE:94:TYR:OH	2:AF:17:LYS:O	2.09	0.68
4:AK:47:ASN:ND2	4:AK:141:MET:SD	2.67	0.68
6:BD:84:TYR:O	6:BD:150:GLN:NE2	2.26	0.68
7:BF:114:ALA:HA	7:BF:117:VAL:HG12	1.76	0.68
9:BH:46:GLU:HB3	9:BH:50:THR:HG22	1.73	0.68
3:BM:16:ARG:O	2:BN:94:TYR:OH	2.11	0.68

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:BN:72:MET:HE3	2:BN:81:CYS:HB3	1.74	0.68
1:BX:65:ILE:HG13	1:BX:66:VAL:HG23	1.75	0.68
2:CA:61:LEU:O	7:CO:101:ARG:HG2	1.93	0.68
2:CF:76:ARG:NH1	10:CF:200:CYC:O2D	2.26	0.68
8:DG:219:LEU:HD13	8:DZ:203:PHE:HE2	1.57	0.68
1:AR:131:ARG:O	1:AR:135:GLU:HG2	1.94	0.68
1:BR:153:PHE:O	1:BR:157:ILE:HG12	1.93	0.68
1:BV:62:ARG:NH2	1:BV:124:GLU:OE2	2.26	0.68
8:CP:241:ASN:O	8:CP:245:SER:OG	2.07	0.68
2:CS:60:LEU:O	2:CS:63:SER:OG	2.10	0.68
1:CV:58:LEU:HD22	1:CV:129:SER:HB3	1.75	0.68
1:DS:76:GLU:HG3	8:DY:202:ARG:HA	1.75	0.68
5:BA:9:CYS:HB2	5:BA:26:TYR:HD1	1.58	0.68
2:CA:60:LEU:HD13	2:CA:72:MET:HE1	1.76	0.68
1:CC:71:ASN:ND2	1:CC:121:THR:OG1	2.26	0.68
6:CM:889:THR:HG22	2:DR:118:SER:HB2	1.75	0.68
9:CQ:191:GLU:O	9:CQ:255:GLY:N	2.27	0.68
1:CY:39:ILE:HA	1:CY:42:THR:HG22	1.76	0.68
1:DU:128:GLN:NE2	1:DU:132:GLU:OE2	2.27	0.68
9:EA:188:VAL:N	9:EA:203:ASP:OD1	2.24	0.68
5:BB:19:GLN:OE1	6:BD:514:ARG:NH1	2.27	0.68
6:CM:36:GLU:N	6:CM:39:GLU:OE1	2.27	0.68
2:AQ:83:ARG:NH2	10:AQ:200:CYC:O2A	2.26	0.68
6:BD:868:ASN:OD1	6:BD:871:ARG:NH2	2.26	0.68
1:BX:16:ARG:O	2:BY:94:TYR:OH	2.11	0.68
6:CM:28:ALA:HB2	6:CM:35:LEU:HD23	1.75	0.68
9:CQ:88:ASN:HD21	9:CQ:174:GLU:HB3	1.57	0.68
2:CS:104:LEU:O	2:CS:108:VAL:HG12	1.94	0.68
2:DK:51:ILE:HG13	2:DK:136:VAL:HG12	1.74	0.68
1:AA:25:ARG:HH21	1:AH:29:PHE:HD1	1.42	0.68
2:AD:8:VAL:HG11	2:AD:27:LEU:HD21	1.75	0.68
2:BY:83:ARG:NH1	2:BY:84:ASP:OD1	2.27	0.68
2:CA:119:LEU:HD11	10:CM:901:CYC:HAA2	1.76	0.68
2:CD:57:ALA:HA	2:CD:61:LEU:HB2	1.76	0.68
2:CW:1:MET:HE2	2:CW:103:ILE:HB	1.74	0.68
1:DJ:76:GLU:OE2	9:EA:155:ARG:NH1	2.26	0.68
2:AS:114:GLU:OE2	6:BD:449:SER:N	2.27	0.68
1:DN:105:GLU:HA	1:DN:109:LEU:HB2	1.75	0.68
1:AC:39:ILE:HD12	1:AC:141:MET:HE1	1.75	0.68
9:BH:191:GLU:O	9:BH:255:GLY:N	2.27	0.68
1:CC:81:CYS:HA	10:CC:200:CYC:HHD	1.74	0.68

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
6:CM:702:THR:O	6:CM:705:GLU:HG3	1.94	0.68
1:DJ:115:MET:HE1	10:DJ:200:CYC:HMB3	1.74	0.68
9:EA:54:ALA:HA	11:EA:400:45D:H221	1.75	0.68
2:AB:100:ASP:OD2	2:AB:102:SER:OG	2.10	0.67
2:AL:94:TYR:OH	6:BD:33:ARG:O	2.12	0.67
2:CA:65:VAL:HG12	2:CA:72:MET:HE3	1.76	0.67
1:DA:85:MET:HG2	1:DA:133:MET:HE3	1.74	0.67
9:DI:224:GLN:NE2	9:DI:228:GLN:O	2.26	0.67
9:EA:89:ARG:HH21	9:EA:160:ASP:HB3	1.59	0.67
10:AV:200:CYC:HMD1	10:AV:200:CYC:HC	1.58	0.67
2:BJ:77:ARG:HG2	10:BJ:200:CYC:HAD1	1.75	0.67
2:BN:88:TYR:OH	2:BN:116:TYR:OH	2.12	0.67
9:CQ:76:PRO:HB3	9:CQ:123:ALA:HB2	1.75	0.67
2:DR:104:LEU:HD13	2:DR:156:ILE:HG13	1.76	0.67
1:AA:44:THR:OG1	1:AA:47:ARG:NH2	2.26	0.67
2:AY:108:VAL:HG12	2:AY:109:LEU:HD23	1.77	0.67
6:BD:792:TYR:OH	1:CY:12:ASP:O	2.12	0.67
9:BH:292:LEU:HD23	9:BH:298:ILE:HA	1.75	0.67
1:BP:122:PRO:HG2	1:BP:125:ALA:HB3	1.76	0.67
10:CF:200:CYC:HHA	10:CF:200:CYC:HBA2	1.75	0.67
8:CP:237:ALA:HA	8:CP:240:ILE:HD12	1.76	0.67
1:CR:106:GLU:HG3	5:DF:66:LEU:HD22	1.76	0.67
1:CV:83:ARG:NH1	10:CV:200:CYC:O2A	2.24	0.67
1:DN:84:ASP:OD2	10:DN:200:CYC:ND	2.27	0.67
1:AH:122:PRO:HG2	1:AH:125:ALA:HB3	1.76	0.67
1:AR:76:GLU:OE2	9:BH:155:ARG:NH2	2.24	0.67
1:BK:7:SER:HB3	1:BK:25:ARG:NH1	2.07	0.67
2:BT:30:TYR:OH	2:BT:97:LEU:O	2.13	0.67
2:CS:94:TYR:HB3	2:CS:103:ILE:HD13	1.76	0.67
2:CW:2:GLN:OE1	2:CW:10:ASN:ND2	2.27	0.67
2:CZ:61:LEU:HD11	10:DA:200:CYC:CAA	2.24	0.67
1:DA:97:VAL:HG21	2:DB:9:ILE:HD11	1.76	0.67
2:DK:19:LEU:HB3	2:DK:24:MET:HE3	1.76	0.67
1:AP:41:GLU:OE2	2:DT:143:ALA:N	2.25	0.67
1:BI:58:LEU:HD22	1:BI:129:SER:HB3	1.76	0.67
2:BJ:76:ARG:NH1	10:BJ:200:CYC:O1D	2.27	0.67
2:BL:76:ARG:NH1	3:BM:106:THR:O	2.28	0.67
1:BX:128:GLN:NE2	1:BX:132:GLU:OE2	2.26	0.67
2:BY:83:ARG:NH2	10:BY:200:CYC:O2A	2.27	0.67
6:CM:697:VAL:HG21	6:CM:701:ARG:HE	1.60	0.67
9:CQ:264:PHE:HD1	9:CQ:294:PRO:HD3	1.59	0.67

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:CU:77:ARG:NH1	10:CU:200:CYC:O1D	2.27	0.67
2:CW:41:ALA:CB	2:CW:97:LEU:HD21	2.24	0.67
5:BA:19:GLN:HB2	6:BD:239:LYS:HG2	1.75	0.67
6:BD:323:LEU:O	6:BD:326:SER:OG	2.09	0.67
9:BH:84:CYS:O	9:BH:88:ASN:ND2	2.21	0.67
10:BM:200:CYC:HMA3	10:BM:200:CYC:NB	2.09	0.67
2:DK:84:ASP:OD2	2:DK:116:TYR:OH	2.11	0.67
2:DM:113:LYS:NZ	2:DM:160:LEU:O	2.28	0.67
1:DS:66:VAL:HA	1:DS:72:ALA:O	1.93	0.67
1:AC:117:ARG:HD2	1:AH:161:SER:HB2	1.75	0.67
1:AC:122:PRO:HG2	1:AC:125:ALA:HB3	1.76	0.67
1:AN:106:GLU:OE1	6:BD:478:LYS:NZ	2.24	0.67
1:AR:81:CYS:HA	10:AR:200:CYC:HHD	1.75	0.67
2:AW:24:MET:HE3	2:AW:28:LYS:HD3	1.75	0.67
6:BD:799:LYS:HB2	6:BD:883:LEU:HD21	1.75	0.67
1:BP:14:GLU:OE1	1:BP:16:ARG:NE	2.22	0.67
2:BQ:104:LEU:HD22	2:BQ:156:ILE:HD11	1.75	0.67
1:CG:61:LYS:HG3	9:CQ:56:PRO:HG2	1.75	0.67
6:CM:400:GLY:O	6:CM:406:GLN:NE2	2.28	0.67
2:CW:60:LEU:O	2:CW:63:SER:OG	2.10	0.67
2:CZ:134:LYS:NZ	2:CZ:154:ASP:OD1	2.26	0.67
9:DI:223:LEU:HD23	9:DI:225:PRO:HD3	1.77	0.67
1:DS:76:GLU:HB2	8:DY:203:PHE:CD2	2.30	0.67
1:AV:44:THR:O	1:AV:47:ARG:NH1	2.27	0.67
1:BP:113:ARG:HG2	1:BP:117:ARG:HH21	1.60	0.67
1:CE:44:THR:O	1:CE:47:ARG:NH1	2.26	0.67
1:DJ:125:ALA:HB3	10:DJ:200:CYC:HMC2	1.76	0.67
1:DN:11:ALA:HB2	1:DN:18:LEU:HD23	1.75	0.67
1:DS:83:ARG:NH1	1:DS:84:ASP:OD1	2.22	0.67
2:AI:128:GLN:NE2	6:BD:681:GLN:OE1	2.28	0.67
1:AV:12:ASP:OD2	2:AW:107:ARG:NH1	2.28	0.67
1:AV:16:ARG:NH1	1:AV:17:TYR:O	2.27	0.67
1:BK:50:ILE:HD13	1:BK:136:VAL:HG12	1.75	0.67
2:BQ:128:GLN:NE2	6:CM:681:GLN:OE1	2.27	0.67
6:CM:247:ASP:HB2	6:CM:248:ILE:HD12	1.75	0.67
9:CQ:222:ALA:HB2	9:CQ:232:VAL:HG12	1.76	0.67
1:AC:50:ILE:HD13	1:AC:136:VAL:HG12	1.76	0.67
2:AO:30:TYR:OH	2:AO:97:LEU:O	2.12	0.67
1:AV:122:PRO:HG2	1:AV:125:ALA:HB3	1.77	0.67
1:CE:84:ASP:OD2	1:CE:116:TYR:OH	2.09	0.67
2:CH:106:GLU:OE2	2:CH:107:ARG:NH1	2.28	0.67

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:DD:83:ARG:NH2	2:DD:84:ASP:OD1	2.25	0.67
1:DJ:126:VAL:HG22	10:DJ:200:CYC:HMC3	1.77	0.67
9:EA:267:ILE:HG22	9:EA:269:VAL:HG13	1.77	0.67
3:AE:81:CYS:HA	10:AE:200:CYC:HHD	1.77	0.66
1:BK:68:PRO:HA	1:BK:73:TYR:HE1	1.60	0.66
1:BP:47:ARG:NH2	1:BP:86:ASP:OD2	2.28	0.66
1:BZ:115:MET:HE1	10:BZ:200:CYC:HMB3	1.75	0.66
1:DC:58:LEU:HD22	1:DC:129:SER:HB2	1.76	0.66
1:DJ:88:TYR:HE2	1:DJ:160:MET:HE1	1.60	0.66
1:DQ:39:ILE:HD11	1:DQ:145:ASP:HB3	1.77	0.66
2:AB:76:ARG:NH1	10:AB:200:CYC:O1D	2.23	0.66
1:BR:128:GLN:OE1	1:BR:131:ARG:NH1	2.29	0.66
2:BT:18:TYR:CD2	6:CM:61:THR:HG22	2.30	0.66
2:BT:37:ARG:NH1	2:BT:96:MET:O	2.27	0.66
2:CA:58:LYS:HB3	7:CO:98:VAL:HG11	1.77	0.66
2:CF:8:VAL:HG13	2:CF:23:ALA:HB1	1.77	0.66
6:CM:805:THR:HA	6:CM:838:MET:HE1	1.77	0.66
1:AX:107:ILE:HG22	10:AX:200:CYC:HBB1	1.76	0.66
1:AZ:71:ASN:ND2	1:AZ:121:THR:OG1	2.28	0.66
6:BD:152:SER:CB	10:BD:902:CYC:H3C	2.26	0.66
2:BL:108:VAL:O	2:BL:112:LEU:HD23	1.96	0.66
1:CC:115:MET:HE1	10:CC:200:CYC:HMB2	1.76	0.66
6:CM:50:GLY:O	6:CM:54:LEU:HG	1.95	0.66
6:CM:210:PHE:O	6:CM:211:ARG:HD3	1.96	0.66
1:CR:134:LYS:NZ	1:CR:154:ASP:OD1	2.27	0.66
1:CT:17:TYR:CE1	2:CU:90:ARG:HD3	2.31	0.66
9:DI:190:ILE:HG21	9:DI:193:VAL:HB	1.78	0.66
2:DT:105:ASP:OD2	2:DT:155:TYR:OH	2.11	0.66
1:AC:8:ILE:HD11	2:AD:98:ALA:HB2	1.77	0.66
9:BH:264:PHE:HD1	9:BH:294:PRO:HD3	1.60	0.66
1:BV:9:VAL:HG23	2:BW:1:MET:HE2	1.78	0.66
6:CM:278:LYS:HD3	6:CM:310:ARG:HG3	1.77	0.66
2:DT:112:LEU:HD13	10:DT:200:CYC:HMB3	1.78	0.66
1:AC:83:ARG:NH2	10:AC:200:CYC:O2A	2.28	0.66
3:AE:67:ALA:HB1	3:AE:68:PRO:HD2	1.76	0.66
1:AJ:16:ARG:O	4:AK:95:TYR:OH	2.11	0.66
6:BD:576:ARG:NH2	6:BD:640:ILE:O	2.22	0.66
2:BL:57:ALA:HA	2:BL:61:LEU:HG	1.76	0.66
2:BN:84:ASP:OD2	2:BN:116:TYR:OH	2.11	0.66
1:BR:8:ILE:HG21	4:BS:1:MET:HE3	1.78	0.66
1:CR:87:TYR:HA	1:CR:90:ARG:HH12	1.60	0.66

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:CS:152:TYR:O	2:CS:156:ILE:HD12	1.96	0.66
1:CV:71:ASN:OD1	1:CV:121:THR:OG1	2.05	0.66
1:AA:39:ILE:HG12	1:AA:145:ASP:HB3	1.76	0.66
2:AU:67:ARG:HG3	2:AU:68:PRO:HD2	1.78	0.66
2:AU:85:LEU:HD22	2:AU:133:ILE:HD11	1.76	0.66
1:AV:19:SER:N	1:AV:22:GLU:OE1	2.24	0.66
10:AY:200:CYC:O1A	5:BB:17:ARG:NH2	2.26	0.66
1:BK:85:MET:HE2	1:BK:129:SER:HB2	1.77	0.66
10:BM:200:CYC:HBD2	10:BM:200:CYC:HMD3	1.76	0.66
4:BS:54:ARG:HB2	4:BS:137:MET:HE1	1.78	0.66
2:CF:21:GLY:HA2	2:CF:24:MET:HB3	1.76	0.66
6:CM:892:VAL:HG21	2:DT:2:GLN:HB2	1.77	0.66
1:CT:91:LEU:HD11	1:CT:107:ILE:HG21	1.76	0.66
2:CW:70:GLY:O	2:CW:77:ARG:NH2	2.28	0.66
1:CY:36:ARG:NH1	1:CY:148:GLU:OE2	2.28	0.66
2:DD:73:TYR:O	2:DD:77:ARG:HB2	1.96	0.66
1:DJ:80:THR:HG23	1:DJ:83:ARG:HH21	1.59	0.66
1:DJ:128:GLN:OE1	1:DJ:131:ARG:NH1	2.28	0.66
2:DM:83:ARG:NH2	10:DM:200:CYC:O1A	2.29	0.66
1:AJ:66:VAL:HG11	1:BX:68:PRO:HG2	1.78	0.66
2:AS:58:LYS:HD3	7:BF:100:ARG:HA	1.78	0.66
6:BD:400:GLY:O	6:BD:406:GLN:NE2	2.29	0.66
3:BM:47:GLU:HG2	3:BM:89:LEU:HD23	1.78	0.66
1:BV:90:ARG:NH2	2:BW:13:ASP:OD1	2.27	0.66
2:CH:126:THR:HB	10:CH:200:CYC:HMC1	1.77	0.66
2:CS:10:ASN:ND2	5:DF:64:ALA:O	2.29	0.66
2:CU:122:PRO:HG2	10:CU:200:CYC:HMC3	1.76	0.66
1:DJ:50:ILE:HG13	1:DJ:136:VAL:HG12	1.76	0.66
1:DJ:141:MET:HE1	1:DJ:149:ALA:HB3	1.78	0.66
1:DQ:50:ILE:HD12	1:DQ:136:VAL:HG12	1.77	0.66
2:DR:35:GLU:HA	2:DR:38:VAL:HG12	1.77	0.66
1:DS:91:LEU:HD11	1:DS:107:ILE:HG21	1.78	0.66
2:AB:81:CYS:HA	10:AB:200:CYC:HHD	1.76	0.66
2:AS:63:SER:H	2:AS:66:THR:HG22	1.60	0.66
2:AS:71:ASN:HD22	2:AS:122:PRO:HD3	1.60	0.66
5:BA:40:GLN:O	5:BA:44:MET:HG3	1.96	0.66
2:BJ:51:ILE:HG22	2:BJ:133:ILE:HD11	1.78	0.66
6:CM:649:PHE:CE1	6:CM:655:PRO:HA	2.31	0.66
1:DA:142:SER:O	1:DA:146:ALA:N	2.25	0.66
9:DI:21:VAL:HG23	9:DI:153:VAL:HG23	1.76	0.66
9:DI:37:LEU:HD12	9:DI:40:ILE:HD11	1.78	0.66

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
9:DI:250:LEU:HD22	9:DI:288:TRP:HZ2	1.59	0.66
1:DL:5:THR:OG1	2:DM:3:ASP:OD1	2.14	0.66
3:AE:80:GLN:OE1	3:AE:83:ARG:NH2	2.29	0.66
2:AO:114:GLU:OE1	6:BD:439:ARG:NH2	2.29	0.66
2:AU:36:LEU:HD21	2:AU:145:ALA:HB2	1.78	0.66
2:BL:40:ALA:HB3	2:BL:96:MET:HE1	1.77	0.66
4:BS:85:ASP:OD2	4:BS:117:TYR:OH	2.12	0.66
1:BV:83:ARG:NH2	10:BV:200:CYC:O2A	2.29	0.66
1:CY:38:ARG:NH1	1:CY:145:ASP:OD2	2.29	0.66
1:CY:85:MET:HE2	1:CY:133:MET:HE2	1.77	0.66
1:DA:37:LEU:HD11	2:DB:27:LEU:HD22	1.78	0.66
2:AB:8:VAL:HG13	2:AB:23:ALA:HB1	1.78	0.66
2:AD:93:THR:O	2:AD:96:MET:HG3	1.96	0.66
1:AR:128:GLN:OE1	1:AR:131:ARG:NH1	2.29	0.66
1:BK:25:ARG:CD	6:CM:42:GLU:HG3	2.26	0.66
2:BL:8:VAL:HG11	2:BL:27:LEU:HD21	1.76	0.66
10:BP:200:CYC:O2D	2:BT:62:TYR:OH	2.09	0.66
2:CD:112:LEU:HD21	10:CD:200:CYC:C1B	2.26	0.66
6:CM:302:ILE:HG22	6:CM:305:LEU:H	1.61	0.66
6:CM:436:GLN:NE2	6:CM:440:PRO:O	2.26	0.66
2:CS:51:ILE:HG21	2:CS:137:THR:HG23	1.77	0.66
2:CZ:104:LEU:O	2:CZ:108:VAL:N	2.28	0.66
1:DU:6:LYS:NZ	1:DU:102:THR:OG1	2.28	0.66
2:AF:71:ASN:HD22	2:AF:121:VAL:HA	1.61	0.65
1:AN:62:ARG:NH2	1:AN:124:GLU:OE2	2.29	0.65
2:AS:85:LEU:HD22	2:AS:133:ILE:HD11	1.77	0.65
1:BZ:38:ARG:NH1	1:BZ:145:ASP:OD2	2.29	0.65
2:CA:85:LEU:HG	10:CM:901:CYC:HBC1	1.78	0.65
2:CF:14:VAL:HG12	5:CK:63:ASN:HD22	1.60	0.65
1:CG:51:VAL:HG22	1:CG:133:MET:HE2	1.79	0.65
2:DM:44:ILE:HD13	2:DM:149:MET:HE2	1.77	0.65
1:DQ:86:ASP:O	1:DQ:90:ARG:HG3	1.96	0.65
2:DR:85:LEU:HD13	2:DR:88:TYR:HD2	1.60	0.65
1:DS:98:SER:HA	2:DT:5:ILE:HG21	1.78	0.65
9:EA:86:LEU:HD21	9:EA:95:CYS:HA	1.78	0.65
9:EA:112:ARG:NH1	9:EA:115:GLU:OE1	2.28	0.65
4:AK:84:ARG:NH2	10:AK:200:CYC:O1A	2.29	0.65
1:AV:44:THR:HG22	2:AW:18:TYR:HD2	1.61	0.65
9:CQ:63:LEU:HD12	9:CQ:105:ILE:HD13	1.78	0.65
9:CQ:206:ASN:O	9:CQ:249:LYS:NZ	2.29	0.65
1:CR:141:MET:HE1	1:CR:146:ALA:HA	1.77	0.65

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:DN:107:ILE:HG22	10:DN:200:CYC:HBB1	1.77	0.65
1:AN:50:ILE:HG12	1:AN:136:VAL:HG12	1.77	0.65
6:BD:52:LEU:HD11	6:BD:216:ALA:HB2	1.76	0.65
3:BM:105:GLU:HA	3:BM:109:LEU:HB2	1.78	0.65
1:BV:86:ASP:OD2	7:CO:109:ARG:NH2	2.29	0.65
2:BY:85:LEU:HG	10:BY:200:CYC:HBC1	1.79	0.65
1:CC:128:GLN:OE1	1:CC:131:ARG:NH1	2.29	0.65
6:CM:213:ASN:O	6:CM:217:LYS:N	2.30	0.65
1:CR:101:VAL:HA	1:CR:104:ILE:HD13	1.78	0.65
1:CT:5:THR:HG23	2:CU:1:MET:HE2	1.77	0.65
2:DM:106:GLU:O	2:DM:110:ASN:ND2	2.29	0.65
1:DN:4:VAL:HG22	1:DN:26:ILE:HD12	1.77	0.65
9:EA:191:GLU:O	9:EA:255:GLY:N	2.29	0.65
10:AC:200:CYC:HMD1	10:AC:200:CYC:HC	1.62	0.65
1:AR:14:GLU:OE1	1:AR:16:ARG:NE	2.23	0.65
1:AZ:128:GLN:OE1	1:AZ:131:ARG:NH1	2.29	0.65
9:BH:219:SER:HA	9:BH:234:LYS:HD2	1.79	0.65
1:BI:21:GLY:O	1:BI:25:ARG:HG2	1.96	0.65
1:BR:40:ALA:O	1:BR:44:THR:HG23	1.97	0.65
1:BZ:131:ARG:O	1:BZ:135:GLU:HG2	1.97	0.65
6:CM:52:LEU:HD21	6:CM:216:ALA:HA	1.78	0.65
2:CU:44:ILE:HD13	2:CU:149:MET:HE2	1.78	0.65
1:CY:94:TYR:OH	2:CZ:17:LYS:O	2.09	0.65
1:DL:131:ARG:NH2	2:DV:47:ASN:OD1	2.28	0.65
4:AK:65:GLU:OE1	4:AK:65:GLU:N	2.30	0.65
10:AU:200:CYC:HAA1	5:BB:36:TRP:HD1	1.60	0.65
2:AY:76:ARG:HB2	1:AZ:110:VAL:HG13	1.78	0.65
6:BD:302:ILE:HG22	6:BD:305:LEU:H	1.60	0.65
10:BL:200:CYC:HMA1	5:CJ:22:LEU:HD22	1.76	0.65
3:BM:124:PRO:HA	3:BM:127:VAL:HG12	1.78	0.65
2:BT:15:GLN:NE2	7:CO:97:ASN:O	2.30	0.65
2:DV:122:PRO:HG2	10:DV:200:CYC:HMC3	1.78	0.65
9:EA:269:VAL:HG23	9:EA:288:TRP:HE3	1.59	0.65
10:AE:200:CYC:HBA1	10:AE:200:CYC:HMA3	1.79	0.65
1:BK:69:GLY:N	1:BK:73:TYR:OH	2.23	0.65
1:BK:105:GLU:HA	1:BK:109:LEU:HB2	1.79	0.65
2:CU:73:TYR:OH	1:CV:90:ARG:NH2	2.30	0.65
2:AB:106:GLU:HG3	2:AB:107:ARG:HG3	1.79	0.65
1:AX:51:VAL:HG22	1:AX:133:MET:HE2	1.79	0.65
3:BM:20:SER:HB3	1:BR:151:ALA:HB1	1.78	0.65
1:CC:65:ILE:HG22	1:CC:72:ALA:HB3	1.78	0.65

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
6:CM:273:LEU:O	6:CM:278:LYS:NZ	2.23	0.65
2:DT:137:THR:O	2:DT:141:VAL:HG22	1.97	0.65
1:AC:29:PHE:HE1	1:AC:99:GLY:HA3	1.61	0.65
2:AW:76:ARG:NH1	10:AW:200:CYC:O2D	2.29	0.65
2:AW:95:ALA:HB2	2:AW:104:LEU:HD21	1.78	0.65
6:BD:861:LEU:HD11	2:DB:87:TYR:CE2	2.32	0.65
2:BN:93:THR:HA	2:BN:149:MET:HE1	1.77	0.65
2:BQ:61:LEU:O	6:CM:682:ARG:NH1	2.26	0.65
2:CD:71:ASN:HD22	2:CD:121:VAL:HA	1.62	0.65
2:CF:63:SER:O	2:CF:67:ARG:HG3	1.97	0.65
2:CS:126:THR:O	2:CS:130:ILE:HG12	1.96	0.65
2:CW:51:ILE:HA	2:CW:136:VAL:HG11	1.77	0.65
8:DG:238:GLN:HA	1:DL:52:LYS:HD2	1.79	0.65
2:DR:129:ALA:O	2:DR:133:ILE:HD12	1.97	0.65
1:AJ:153:PHE:O	1:AJ:157:ILE:HG12	1.96	0.65
6:BD:736:LYS:HE3	6:BD:740:ARG:HD2	1.76	0.65
1:CG:6:LYS:HG3	1:CG:100:ASP:OD2	1.97	0.65
6:CM:323:LEU:O	6:CM:326:SER:OG	2.13	0.65
1:CV:14:GLU:OE1	1:CV:16:ARG:NE	2.30	0.65
1:CY:40:ALA:HB2	1:CY:97:VAL:HG22	1.76	0.65
1:DJ:21:GLY:O	1:DJ:25:ARG:HG3	1.97	0.65
1:DJ:50:ILE:HG23	1:DJ:136:VAL:HB	1.78	0.65
1:DL:98:SER:HA	2:DM:5:ILE:HG21	1.78	0.65
2:DM:127:VAL:O	2:DM:131:GLN:HG2	1.97	0.65
4:AK:68:ARG:HB3	4:AK:69:PRO:HD2	1.78	0.65
2:AS:114:GLU:HG3	6:BD:471:LYS:HB2	1.78	0.65
1:AZ:105:GLU:HA	1:AZ:109:LEU:HB3	1.79	0.65
8:BG:216:MET:HE3	9:BH:12:PHE:HB2	1.78	0.65
4:BS:123:PRO:HB2	4:BS:126:PRO:HD2	1.79	0.65
9:CQ:204:ASN:HB2	9:CQ:213:LEU:HD13	1.79	0.65
2:CZ:61:LEU:HD11	10:DA:200:CYC:HAA1	1.78	0.65
1:DC:128:GLN:NE2	1:DC:132:GLU:OE2	2.30	0.65
1:DL:154:ASP:CG	2:DV:46:ALA:HB2	2.22	0.65
1:AC:97:VAL:HG11	2:AD:19:LEU:HD12	1.79	0.64
2:AW:111:GLY:O	2:AW:115:THR:OG1	2.15	0.64
1:AZ:65:ILE:HG22	1:AZ:72:ALA:HB3	1.80	0.64
7:BF:104:GLY:O	7:BF:111:LYS:NZ	2.24	0.64
2:BN:22:ALA:O	2:BN:26:LYS:NZ	2.29	0.64
2:CW:90:ARG:NH1	2:CW:94:TYR:OH	2.29	0.64
8:DH:232:ILE:HD12	8:DH:233:PRO:HD2	1.79	0.64
9:DI:111:TYR:HB2	11:DI:400:45D:H301	1.77	0.64

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:DJ:141:MET:HE1	1:DJ:149:ALA:CB	2.27	0.64
1:DQ:90:ARG:HG2	2:DR:18:TYR:CZ	2.32	0.64
1:DQ:154:ASP:HA	1:DQ:157:ILE:HG12	1.79	0.64
2:DR:123:ILE:HG23	2:DR:160:LEU:HD23	1.78	0.64
2:AI:65:VAL:HG12	2:AI:72:MET:HB2	1.79	0.64
1:AZ:129:SER:O	1:AZ:133:MET:HG3	1.97	0.64
2:BL:33:SER:O	2:BL:37:ARG:HG3	1.97	0.64
2:BL:83:ARG:NH1	10:BL:200:CYC:O2A	2.31	0.64
4:BS:84:ARG:NH2	10:BS:200:CYC:O1A	2.30	0.64
2:BY:110:ASN:HD22	6:CM:668:ARG:HG2	1.62	0.64
2:CA:3:ASP:OD1	2:CA:6:THR:OG1	2.12	0.64
1:CG:52:LYS:NZ	8:CP:240:ILE:O	2.30	0.64
2:CZ:30:TYR:O	2:CZ:33:SER:OG	2.13	0.64
1:DL:138:SER:HA	1:DL:146:ALA:HB1	1.79	0.64
1:DQ:76:GLU:HA	8:DZ:203:PHE:CZ	2.31	0.64
2:DR:5:ILE:O	2:DR:9:ILE:HG12	1.96	0.64
1:DU:71:ASN:ND2	1:DU:121:THR:OG1	2.30	0.64
9:EA:111:TYR:HB2	11:EA:400:45D:H301	1.78	0.64
4:AK:88:TYR:HH	6:BD:252:GLU:H	1.43	0.64
2:BJ:100:ASP:OD2	2:BJ:102:SER:OG	2.12	0.64
2:BT:2:GLN:HB2	2:BT:6:THR:CG2	2.28	0.64
1:BX:58:LEU:HD22	1:BX:129:SER:HB3	1.78	0.64
1:BZ:97:VAL:HG21	2:CA:19:LEU:HD12	1.78	0.64
1:CG:107:ILE:HG22	10:CG:200:CYC:HBB1	1.80	0.64
9:CQ:46:GLU:HB3	9:CQ:50:THR:HG22	1.78	0.64
1:CV:39:ILE:HG12	1:CV:145:ASP:HB3	1.80	0.64
1:CV:50:ILE:HA	1:CV:136:VAL:HG11	1.77	0.64
5:DF:42:ARG:NH1	5:DF:46:MET:SD	2.70	0.64
1:DQ:58:LEU:HD22	1:DQ:129:SER:CB	2.28	0.64
1:DQ:122:PRO:HG2	1:DQ:125:ALA:HB3	1.78	0.64
2:DR:126:THR:HG23	10:DR:200:CYC:HBC3	1.78	0.64
4:AK:108:ARG:HD3	6:BD:249:GLN:HB3	1.80	0.64
6:BD:676:ARG:HB2	6:BD:679:ILE:CG1	2.27	0.64
1:CC:110:VAL:HG13	2:CH:76:ARG:HB2	1.79	0.64
1:CE:19:SER:N	1:CE:22:GLU:OE1	2.25	0.64
6:CM:603:GLU:OE2	6:CM:607:ARG:NE	2.23	0.64
6:CM:748:GLU:OE1	6:CM:877:THR:OG1	2.12	0.64
6:CM:765:SER:OG	10:DK:200:CYC:O2A	2.14	0.64
6:CM:792:TYR:CD1	6:CM:831:LEU:HD13	2.32	0.64
6:CM:868:ASN:OD1	6:CM:871:ARG:NH2	2.29	0.64
9:CQ:84:CYS:O	9:CQ:88:ASN:ND2	2.21	0.64

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:CT:68:PRO:HD2	8:DY:230:ASN:HD21	1.61	0.64
2:DB:122:PRO:HB2	2:DB:125:SER:HB2	1.77	0.64
1:DL:44:THR:O	1:DL:47:ARG:NH1	2.30	0.64
2:DM:104:LEU:HD22	2:DM:156:ILE:HD11	1.79	0.64
1:DQ:78:THR:O	1:DQ:82:LEU:HG	1.98	0.64
1:AC:7:SER:HB3	1:AC:25:ARG:NH1	2.12	0.64
2:AI:75:THR:HG21	1:AJ:112:VAL:HG12	1.78	0.64
1:AJ:40:ALA:O	1:AJ:44:THR:HG23	1.98	0.64
2:AL:30:TYR:OH	2:AL:97:LEU:O	2.12	0.64
6:BD:28:ALA:HB2	6:BD:35:LEU:HD23	1.79	0.64
6:BD:137:ARG:NH1	6:BD:138:PRO:O	2.31	0.64
2:BL:93:THR:O	2:BL:96:MET:HG3	1.98	0.64
1:CC:29:PHE:O	1:CC:36:ARG:NH1	2.30	0.64
1:CE:37:LEU:HD21	2:CF:24:MET:HE1	1.79	0.64
6:CM:814:LEU:N	6:CM:818:GLU:OE1	2.27	0.64
1:AC:16:ARG:O	2:AD:94:TYR:OH	2.15	0.64
2:AS:3:ASP:OD1	2:AS:6:THR:OG1	2.08	0.64
1:AV:137:ALA:O	1:AV:141:MET:HG3	1.98	0.64
6:CM:609:LEU:HD13	6:CM:640:ILE:HD11	1.80	0.64
1:CV:22:GLU:O	1:CV:26:ILE:HD12	1.98	0.64
1:CV:128:GLN:OE1	1:CV:131:ARG:NH1	2.29	0.64
9:DI:250:LEU:HD23	9:DI:273:VAL:HG22	1.79	0.64
2:DR:73:TYR:OH	1:DS:90:ARG:NH2	2.30	0.64
1:AJ:58:LEU:HD22	1:AJ:129:SER:HB3	1.79	0.64
1:AR:62:ARG:HG2	1:AR:65:ILE:HG12	1.79	0.64
2:AU:48:ALA:HA	2:AU:51:ILE:HD13	1.79	0.64
2:CF:1:MET:N	5:CK:67:ALA:O	2.31	0.64
2:CH:71:ASN:HD22	2:CH:122:PRO:HD3	1.61	0.64
9:CQ:284:MET:HE1	9:CQ:307:ALA:HB2	1.80	0.64
1:DA:100:ASP:OD2	1:DA:102:THR:HG23	1.97	0.64
2:DB:137:THR:O	2:DB:141:VAL:HG12	1.98	0.64
9:DI:39:LEU:HD12	9:DI:139:LEU:HD12	1.80	0.64
1:DQ:65:ILE:HG21	10:DQ:200:CYC:HBC2	1.80	0.64
1:AC:20:PRO:HG2	6:BD:172:PRO:HD3	1.80	0.64
4:AK:135:LYS:HD3	4:AK:157:PHE:HB2	1.79	0.64
2:CF:108:VAL:HG11	2:CF:156:ILE:CD1	2.28	0.64
1:CG:81:CYS:HA	10:CG:200:CYC:HHD	1.80	0.64
6:CM:726:LEU:HD11	6:CM:735:VAL:HG22	1.78	0.64
2:DR:62:TYR:OH	10:DS:200:CYC:O2D	2.08	0.64
2:DV:47:ASN:HB2	2:DV:51:ILE:HD11	1.78	0.64
2:AU:54:GLU:OE1	2:AU:136:VAL:HG21	1.98	0.64

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:AV:11:ALA:HB2	1:AV:18:LEU:HD23	1.80	0.64
6:BD:609:LEU:HD13	6:BD:640:ILE:HD11	1.79	0.64
6:CM:52:LEU:HD11	6:CM:216:ALA:HB2	1.79	0.64
2:DK:60:LEU:O	2:DK:63:SER:OG	2.15	0.64
1:DN:58:LEU:HD22	1:DN:129:SER:CB	2.28	0.64
2:DR:85:LEU:CD2	10:DR:200:CYC:HBC1	2.27	0.64
1:AA:5:THR:OG1	2:AB:1:MET:SD	2.53	0.64
1:AR:84:ASP:OD2	10:AR:200:CYC:ND	2.28	0.64
6:BD:377:VAL:O	6:BD:381:VAL:HG22	1.98	0.64
2:BL:24:MET:O	2:BL:28:LYS:HG2	1.98	0.64
1:BR:9:VAL:HG11	6:CM:246:ASN:HB3	1.80	0.64
2:CD:130:ILE:HA	2:CD:133:ILE:HD12	1.80	0.64
5:CJ:17:ARG:HG3	6:CM:240:VAL:HB	1.80	0.64
9:CQ:205:LEU:HA	9:CQ:210:PHE:HE1	1.63	0.64
9:CQ:210:PHE:HA	9:CQ:213:LEU:HB3	1.79	0.64
1:DA:77:MET:HE3	8:DG:201:ARG:HB3	1.80	0.64
5:DF:43:ILE:HG21	5:DF:50:ILE:HD11	1.80	0.64
9:DI:204:ASN:HB2	9:DI:213:LEU:HD21	1.80	0.64
1:DU:48:GLU:O	1:DU:52:LYS:HG2	1.98	0.64
3:AE:37:ILE:O	3:AE:41:GLU:HG2	1.98	0.63
1:AJ:65:ILE:HG13	1:AJ:66:VAL:HG23	1.80	0.63
2:AW:75:THR:HG23	10:AX:200:CYC:HAB2	1.79	0.63
5:BB:38:ARG:NH2	6:BD:641:ASP:O	2.24	0.63
1:BK:77:MET:SD	10:BK:200:CYC:O2D	2.55	0.63
2:BL:109:LEU:HD23	2:BL:159:GLY:HA3	1.79	0.63
2:BT:101:ALA:HB1	2:BT:104:LEU:HD12	1.80	0.63
1:CC:129:SER:O	1:CC:133:MET:HG3	1.98	0.63
5:CJ:9:CYS:HB2	5:CJ:26:TYR:HD1	1.62	0.63
9:DI:68:LEU:HD13	9:DI:112:ARG:HG2	1.78	0.63
9:DI:87:ALA:O	9:DI:89:ARG:NH1	2.32	0.63
1:DL:50:ILE:HG13	1:DL:136:VAL:CG1	2.29	0.63
9:EA:117:MET:HE1	9:EA:125:ILE:HG23	1.80	0.63
10:AF:200:CYC:OB	5:BA:41:GLN:NE2	2.31	0.63
6:BD:5:ALA:HB2	6:BD:436:GLN:HG3	1.79	0.63
6:BD:199:ILE:HD13	6:BD:227:LEU:CD2	2.29	0.63
2:BN:109:LEU:HD13	2:BN:159:GLY:HA3	1.78	0.63
4:BS:72:ASN:HD22	4:BS:122:VAL:HA	1.63	0.63
2:BT:88:TYR:HH	2:BT:116:TYR:HH	1.44	0.63
2:BW:1:MET:SD	2:BW:103:ILE:HD12	2.38	0.63
1:CE:137:ALA:O	1:CE:141:MET:HG3	1.98	0.63
6:CM:225:ASP:O	6:CM:228:LEU:HG	1.99	0.63

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:AI:60:LEU:O	2:AI:63:SER:OG	2.12	0.63
1:AP:58:LEU:HD22	1:AP:129:SER:HB3	1.79	0.63
1:AP:89:LEU:HB2	1:AP:133:MET:HE1	1.79	0.63
1:AX:81:CYS:HA	10:AX:200:CYC:HHD	1.79	0.63
2:BL:72:MET:HE1	2:BL:81:CYS:HB3	1.80	0.63
1:BP:93:THR:O	1:BP:97:VAL:HG23	1.98	0.63
1:BR:97:VAL:HG21	4:BS:19:LEU:HD12	1.81	0.63
2:CF:127:VAL:O	2:CF:131:GLN:HG2	1.99	0.63
2:CU:37:ARG:NH1	2:CU:148:GLU:OE2	2.31	0.63
9:DI:155:ARG:O	9:DI:159:VAL:HG23	1.97	0.63
1:AA:12:ASP:OD2	2:AB:107:ARG:NH1	2.26	0.63
1:AX:142:SER:OG	1:AX:144:ASP:OD1	2.10	0.63
2:BJ:2:GLN:HG2	5:CJ:65:GLY:HA3	1.80	0.63
1:BV:35:ALA:O	1:BV:39:ILE:HD12	1.98	0.63
2:BW:4:ALA:O	2:BW:8:VAL:HG23	1.99	0.63
2:BW:158:SER:O	2:BW:161:SER:OG	2.12	0.63
10:BX:200:CYC:HMD1	10:BX:200:CYC:HC	1.62	0.63
1:CG:109:LEU:HD21	1:CG:159:LYS:CG	2.28	0.63
6:CM:867:PRO:HB3	2:DT:114:GLU:OE1	1.98	0.63
2:CZ:20:ASP:O	2:CZ:24:MET:HG2	1.99	0.63
1:DJ:96:VAL:HG22	1:DJ:152:TYR:HD2	1.63	0.63
4:AK:72:ASN:HD22	4:AK:123:PRO:HD2	1.63	0.63
6:BD:316:MET:HE2	6:BD:395:VAL:HG22	1.79	0.63
1:CC:92:VAL:O	1:CC:96:VAL:HG13	1.99	0.63
8:CP:221:MET:O	8:CP:225:ILE:HG22	1.99	0.63
1:CR:75:GLU:HA	1:CR:78:THR:HG22	1.81	0.63
2:CU:71:ASN:HB3	10:CU:200:CYC:OC	1.99	0.63
2:CZ:96:MET:HA	2:CZ:152:TYR:HE2	1.63	0.63
1:DN:122:PRO:HD2	10:DN:200:CYC:OC	1.99	0.63
2:DT:66:THR:HG21	10:DU:200:CYC:CBA	2.28	0.63
9:EA:224:GLN:HE22	9:EA:230:PRO:HG3	1.62	0.63
1:AV:97:VAL:HG11	2:AW:19:LEU:CD1	2.29	0.63
1:AV:109:LEU:HD22	1:AV:159:LYS:HB3	1.81	0.63
1:BK:10:ASN:ND2	6:CM:27:GLN:OE1	2.32	0.63
1:BK:105:GLU:HA	1:BK:109:LEU:CB	2.29	0.63
1:BK:105:GLU:HG2	1:BK:109:LEU:HD23	1.81	0.63
4:BS:64:PRO:HG2	7:CO:105:PRO:HB2	1.80	0.63
6:CM:15:GLN:HB3	6:CM:260:ALA:O	1.99	0.63
6:CM:489:PRO:HD2	6:CM:492:ASN:HD22	1.63	0.63
2:CS:37:ARG:HH12	2:CS:99:GLY:HA2	1.62	0.63
1:CY:44:THR:HG22	2:CZ:18:TYR:HD2	1.64	0.63

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:DB:20:ASP:O	2:DB:24:MET:HG2	1.99	0.63
9:DI:62:GLN:HA	9:DI:65:GLU:OE1	1.99	0.63
1:DU:57:ARG:HH22	1:DU:135:GLU:HB3	1.63	0.63
2:AD:36:LEU:HD11	2:AD:144:ASP:HB2	1.80	0.63
2:AD:83:ARG:HD2	5:BA:22:LEU:CD1	2.29	0.63
1:AR:138:SER:HA	1:AR:146:ALA:HB1	1.80	0.63
2:AU:74:THR:HA	10:AV:200:CYC:HBB2	1.81	0.63
1:AV:54:ALA:CB	1:AV:133:MET:HG2	2.28	0.63
2:AY:1:MET:HE2	2:AY:103:ILE:HB	1.79	0.63
6:BD:347:PHE:CE1	6:BD:380:LEU:HD21	2.34	0.63
6:BD:454:LEU:HD22	6:BD:667:MET:HE1	1.80	0.63
1:BI:98:SER:HA	2:BJ:5:ILE:HG21	1.79	0.63
2:BJ:15:GLN:OE1	2:BJ:17:LYS:NZ	2.31	0.63
1:BK:2:SER:HA	1:BK:100:ASP:HB3	1.79	0.63
10:BQ:200:CYC:HMA3	10:BQ:200:CYC:HB	1.63	0.63
2:BT:37:ARG:HG3	7:CO:86:VAL:HG23	1.81	0.63
2:BY:116:TYR:HE2	2:BY:160:LEU:HD21	1.64	0.63
1:CC:89:LEU:HB2	1:CC:133:MET:HE1	1.81	0.63
2:CH:83:ARG:NH2	10:CH:200:CYC:O2A	2.32	0.63
2:CS:83:ARG:NH2	2:CS:84:ASP:OD1	2.29	0.63
2:CU:62:TYR:OH	10:CV:200:CYC:O2D	2.13	0.63
2:CW:15:GLN:HG2	2:CW:17:LYS:HG2	1.81	0.63
1:DL:50:ILE:HG13	1:DL:136:VAL:HB	1.81	0.63
2:DO:81:CYS:HA	10:DO:200:CYC:HHD	1.81	0.63
2:DT:20:ASP:O	2:DT:24:MET:HG2	1.99	0.63
3:AE:37:ILE:HG23	2:AF:24:MET:CE	2.29	0.63
1:AH:68:PRO:HA	1:AH:73:TYR:CD2	2.33	0.63
2:AI:71:ASN:ND2	2:AI:121:VAL:HA	2.14	0.63
2:AS:63:SER:O	2:AS:67:ARG:HG3	1.99	0.63
1:AV:71:ASN:OD1	1:AV:121:THR:OG1	2.16	0.63
2:AW:73:TYR:OH	1:AX:90:ARG:NH2	2.30	0.63
1:AZ:92:VAL:O	1:AZ:96:VAL:HG13	1.99	0.63
9:BH:89:ARG:HD3	9:BH:170:LYS:HD2	1.80	0.63
1:BK:83:ARG:NH1	10:BK:200:CYC:O2A	2.32	0.63
1:BK:141:MET:HG3	1:BK:146:ALA:HB2	1.80	0.63
1:BP:68:PRO:HA	1:BP:73:TYR:CD2	2.33	0.63
1:BR:12:ASP:OD1	4:BS:91:ARG:NH1	2.27	0.63
1:DQ:77:MET:CE	8:DZ:201:ARG:HB2	2.28	0.63
2:DR:24:MET:O	2:DR:28:LYS:HG2	1.99	0.63
1:AA:105:GLU:HG3	1:AA:110:VAL:HG23	1.79	0.63
1:AA:115:MET:HG3	2:AF:78:TYR:CD1	2.34	0.63

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:AF:57:ALA:HA	2:AF:61:LEU:HD12	1.79	0.63
1:AN:113:ARG:HG2	1:AN:123:ILE:HD13	1.81	0.63
9:BH:236:ASN:OD1	9:BH:239:ARG:NH2	2.32	0.63
1:BR:86:ASP:OD1	4:BS:18:TYR:OH	2.11	0.63
2:BY:71:ASN:O	2:BY:77:ARG:HD3	1.98	0.63
2:CF:75:THR:HG23	10:CG:200:CYC:HAB2	1.80	0.63
2:CH:108:VAL:HG12	2:CH:109:LEU:HD23	1.81	0.63
9:CQ:239:ARG:NH2	1:DA:41:GLU:HB3	2.14	0.63
2:DB:51:ILE:HD11	2:DB:140:LEU:HD23	1.81	0.63
1:DL:141:MET:HE3	1:DL:146:ALA:HA	1.81	0.63
2:DO:93:THR:HA	2:DO:149:MET:HE1	1.80	0.63
1:DQ:105:GLU:O	1:DQ:110:VAL:HG23	1.98	0.63
1:AH:109:LEU:HG	1:AH:159:LYS:HG2	1.81	0.62
1:AR:11:ALA:HB2	1:AR:18:LEU:HD23	1.79	0.62
10:AY:200:CYC:HMA1	10:AY:200:CYC:NB	2.14	0.62
1:BX:122:PRO:HG2	1:BX:125:ALA:HB3	1.80	0.62
1:CC:68:PRO:HG2	1:CV:66:VAL:HG21	1.81	0.62
1:CE:12:ASP:OD2	2:CF:107:ARG:NH1	2.31	0.62
1:DA:66:VAL:HA	1:DA:72:ALA:O	1.98	0.62
2:DM:78:TYR:CD1	1:DN:115:MET:HG3	2.34	0.62
2:DO:127:VAL:O	2:DO:131:GLN:HG2	1.99	0.62
2:DV:88:TYR:OH	2:DV:112:LEU:HD21	1.99	0.62
2:AL:72:MET:HE3	2:AL:81:CYS:HB3	1.79	0.62
2:AO:74:THR:HG23	1:AP:107:ILE:HA	1.78	0.62
2:AU:2:GLN:NE2	2:AU:10:ASN:OD1	2.24	0.62
2:AY:76:ARG:HB2	1:AZ:110:VAL:CG1	2.29	0.62
6:BD:152:SER:HB2	10:BD:902:CYC:H3C	1.80	0.62
2:BN:55:ALA:O	2:BN:59:SER:OG	2.14	0.62
4:BS:44:ILE:HD12	4:BS:94:SER:HB2	1.80	0.62
2:CA:51:ILE:HG23	2:CA:136:VAL:HG12	1.80	0.62
1:CE:54:ALA:CB	1:CE:133:MET:HG2	2.29	0.62
1:CR:14:GLU:OE1	1:CR:16:ARG:NE	2.27	0.62
1:CY:4:VAL:HG13	2:CZ:97:LEU:HD23	1.81	0.62
1:CY:118:SER:O	2:DD:53:LYS:NZ	2.24	0.62
2:CZ:2:GLN:N	2:CZ:102:SER:OG	2.25	0.62
2:CZ:108:VAL:HG12	2:CZ:109:LEU:HD23	1.81	0.62
1:DA:38:ARG:NH1	1:DA:145:ASP:OD2	2.32	0.62
1:DL:8:ILE:HD13	1:DL:18:LEU:HD11	1.79	0.62
2:DO:58:LYS:NZ	9:EA:100:SER:O	2.29	0.62
2:DT:104:LEU:HD22	2:DT:156:ILE:HD11	1.81	0.62
2:AQ:105:ASP:OD2	2:AQ:155:TYR:OH	2.10	0.62

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
5:BA:13:GLN:NE2	5:BA:46:MET:O	2.32	0.62
5:BB:39:GLU:O	5:BB:43:ILE:HG12	2.00	0.62
6:BD:706:ILE:HG13	2:DD:15:GLN:HE21	1.64	0.62
6:BD:797:ASN:HB2	10:CZ:200:CYC:OB	2.00	0.62
1:BI:12:ASP:OD2	2:BJ:107:ARG:NH1	2.29	0.62
1:BP:16:ARG:O	2:BQ:94:TYR:OH	2.16	0.62
1:BZ:77:MET:HE2	9:CQ:11:ILE:HD13	1.81	0.62
1:CC:38:ARG:O	1:CC:42:THR:HG23	1.99	0.62
1:CC:110:VAL:CG1	2:CH:76:ARG:HB2	2.30	0.62
1:CE:53:GLN:O	1:CE:57:ARG:HD3	1.98	0.62
6:CM:621:ASN:ND2	10:CM:901:CYC:O1A	2.32	0.62
1:CY:100:ASP:OD2	1:CY:102:THR:OG1	2.10	0.62
1:CY:127:ALA:HA	1:CY:160:MET:HE1	1.81	0.62
2:DB:134:LYS:HG3	2:DB:150:GLY:HA2	1.80	0.62
8:DG:218:PHE:HE1	1:DQ:83:ARG:HH11	1.46	0.62
1:AC:105:GLU:HA	1:AC:109:LEU:CB	2.29	0.62
2:AI:104:LEU:HD11	2:AI:152:TYR:HB3	1.82	0.62
1:AV:134:LYS:HD3	1:AV:153:PHE:HB3	1.81	0.62
2:BL:62:TYR:HE2	3:BM:80:GLN:HG3	1.65	0.62
2:BN:37:ARG:NH1	2:BN:148:GLU:OE2	2.33	0.62
2:BN:122:PRO:HB2	2:BN:125:SER:HB2	1.81	0.62
2:BT:81:CYS:HA	10:BT:200:CYC:HBC3	1.80	0.62
2:BW:134:LYS:HB2	2:BW:153:LEU:HD13	1.80	0.62
1:CC:114:GLU:HG3	1:CC:117:ARG:HH21	1.65	0.62
1:CG:84:ASP:OD2	10:CG:200:CYC:ND	2.32	0.62
1:CY:90:ARG:HG2	2:CZ:18:TYR:CZ	2.33	0.62
2:DR:76:ARG:NH2	10:DR:200:CYC:O1D	2.27	0.62
2:AW:74:THR:OG1	2:AW:77:ARG:NH2	2.27	0.62
1:BK:122:PRO:HG2	1:BK:125:ALA:HB3	1.79	0.62
2:CD:64:ASP:OD1	2:CD:67:ARG:NH1	2.31	0.62
2:CF:2:GLN:O	2:CF:102:SER:OG	2.18	0.62
6:CM:749:ARG:HH12	10:DV:200:CYC:HBA1	1.63	0.62
2:DB:126:THR:O	2:DB:130:ILE:HG13	2.00	0.62
1:DQ:39:ILE:HA	1:DQ:42:THR:HG22	1.82	0.62
5:DX:42:ARG:NH1	5:DX:46:MET:SD	2.73	0.62
10:AD:200:CYC:HAA2	5:BA:11:PRO:HB3	1.81	0.62
3:AE:69:GLY:N	3:AE:73:TYR:OH	2.26	0.62
1:AH:40:ALA:HB2	1:AH:97:VAL:HG22	1.81	0.62
1:AR:83:ARG:NH2	10:AR:200:CYC:O2A	2.32	0.62
1:BI:115:MET:HG3	2:BN:78:TYR:CD1	2.34	0.62
1:BK:96:VAL:HA	1:BK:152:TYR:HE2	1.65	0.62

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:BT:17:LYS:O	6:CM:165:TYR:OH	2.18	0.62
2:CA:67:ARG:O	2:CA:73:TYR:HB2	1.99	0.62
1:CC:7:SER:HB2	1:CC:18:LEU:HD11	1.80	0.62
6:CM:754:TYR:OH	2:DV:83:ARG:NH1	2.33	0.62
9:EA:253:GLU:OE2	9:EA:272:LYS:NZ	2.30	0.62
1:AC:8:ILE:CD1	2:AD:98:ALA:HB2	2.30	0.62
1:AP:95:GLY:HA3	1:AP:104:ILE:HD11	1.82	0.62
10:AU:200:CYC:HB	10:AU:200:CYC:HMA1	1.65	0.62
1:AX:109:LEU:HD21	1:AX:159:LYS:CG	2.29	0.62
6:BD:827:ALA:HB2	2:CZ:87:TYR:OH	1.99	0.62
9:BH:235:GLU:CG	1:DS:45:GLY:HA2	2.30	0.62
1:BK:35:ALA:O	1:BK:39:ILE:HG12	1.98	0.62
2:BQ:85:LEU:HG	10:BQ:200:CYC:HBC1	1.81	0.62
1:BR:18:LEU:HD13	4:BS:98:ILE:HD12	1.80	0.62
4:BS:76:THR:HG23	10:CM:902:CYC:OB	1.99	0.62
4:BS:142:ALA:HB1	4:BS:147:LEU:HD12	1.81	0.62
1:BV:154:ASP:HB3	2:CD:46:ALA:HB2	1.82	0.62
1:CC:105:GLU:HA	1:CC:109:LEU:HB3	1.82	0.62
2:CH:71:ASN:HD22	2:CH:121:VAL:HA	1.64	0.62
2:CH:71:ASN:ND2	2:CH:121:VAL:HA	2.14	0.62
9:CQ:289:ARG:NH2	9:DI:307:ALA:O	2.33	0.62
1:CR:77:MET:HA	1:CR:80:THR:HG22	1.81	0.62
1:DA:50:ILE:HD11	1:DA:136:VAL:HG13	1.82	0.62
1:DA:98:SER:HA	2:DB:5:ILE:CD1	2.29	0.62
3:AE:29:PHE:HA	3:AE:32:THR:HG22	1.82	0.62
3:AE:85:TYR:CE1	3:AE:129:ALA:HB1	2.35	0.62
4:AK:44:ILE:HG22	4:AK:138:ILE:HD12	1.80	0.62
1:AN:19:SER:HB3	1:AN:22:GLU:HG3	1.81	0.62
1:AN:108:GLY:HA2	2:AS:75:THR:HG23	1.82	0.62
2:AQ:71:ASN:O	2:AQ:77:ARG:HD3	2.00	0.62
1:CC:89:LEU:O	1:CC:93:THR:HG23	2.00	0.62
2:DB:71:ASN:HB3	10:DB:200:CYC:OC	2.00	0.62
2:DD:112:LEU:HD13	10:DD:200:CYC:HMB1	1.82	0.62
9:DI:260:ALA:N	9:DI:264:PHE:O	2.21	0.62
9:EA:188:VAL:HG11	9:EA:252:PRO:HG2	1.81	0.62
4:AK:51:ILE:HD13	4:AK:138:ILE:CD1	2.30	0.62
4:AK:64:PRO:CG	7:BF:105:PRO:HB2	2.30	0.62
6:BD:612:ARG:NH2	6:BD:644:GLU:OE1	2.33	0.62
2:BL:112:LEU:HA	2:BL:115:THR:HG22	1.82	0.62
2:BT:94:TYR:OH	6:CM:33:ARG:O	2.17	0.62
1:BV:84:ASP:OD2	10:BV:200:CYC:ND	2.33	0.62

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:CS:91:TYR:HE1	2:CS:107:ARG:HD3	1.65	0.62
1:CV:18:LEU:HD12	1:CV:22:GLU:HB3	1.81	0.62
1:CY:92:VAL:HG11	1:CY:153:PHE:CD1	2.34	0.62
1:DL:143:SER:OG	1:DL:144:ASP:N	2.29	0.62
2:DM:62:TYR:OH	10:DN:200:CYC:O2D	2.16	0.62
1:DS:104:ILE:HG22	1:DS:109:LEU:HD23	1.82	0.62
2:DT:3:ASP:H	2:DT:6:THR:HG1	1.47	0.62
2:DT:123:ILE:HG23	2:DT:160:LEU:HG	1.81	0.62
2:AQ:83:ARG:NH1	2:AQ:84:ASP:OD1	2.32	0.62
1:AV:83:ARG:NH1	10:AV:200:CYC:O1A	2.23	0.62
4:BS:92:TYR:CE2	10:BS:200:CYC:HBB1	2.35	0.62
1:BV:56:ASP:OD1	7:CO:117:VAL:HG23	2.00	0.62
6:CM:71:ALA:O	6:CM:75:ILE:HG12	2.00	0.62
6:CM:861:LEU:HD23	10:DT:200:CYC:OB	1.99	0.62
2:CS:104:LEU:HD21	2:CS:156:ILE:CD1	2.30	0.62
1:CY:115:MET:SD	2:DD:78:TYR:HB3	2.40	0.62
1:DA:58:LEU:HD22	1:DA:129:SER:HB3	1.82	0.62
1:DJ:37:LEU:HD11	2:DK:24:MET:CE	2.30	0.62
2:DK:41:ALA:HB2	2:DK:97:LEU:HD21	1.82	0.62
3:AE:131:THR:HG22	3:AE:157:ILE:HD13	1.82	0.61
1:AR:101:VAL:HG21	1:AR:155:PHE:CD2	2.35	0.61
1:AR:122:PRO:HG2	1:AR:125:ALA:HB3	1.82	0.61
1:AR:151:ALA:HB1	1:AV:20:PRO:HB3	1.82	0.61
6:BD:225:ASP:O	6:BD:228:LEU:HG	1.99	0.61
6:BD:385:GLU:OE2	6:BD:399:ARG:NH2	2.33	0.61
6:BD:702:THR:HB	2:DT:151:VAL:HG22	1.82	0.61
6:BD:801:ILE:CD1	6:BD:822:TYR:HB3	2.29	0.61
2:BJ:1:MET:CE	2:BJ:103:ILE:HD13	2.30	0.61
3:BM:109:LEU:HD22	3:BM:159:PHE:HB3	1.81	0.61
6:CM:182:LEU:HD12	6:CM:185:VAL:HG23	1.81	0.61
9:DI:107:LEU:HD22	9:DI:155:ARG:HG2	1.82	0.61
2:DK:36:LEU:HA	2:DK:39:ARG:HG2	1.81	0.61
2:DR:35:GLU:HA	2:DR:38:VAL:CG1	2.30	0.61
1:DS:134:LYS:HE3	1:DS:150:SER:OG	1.99	0.61
9:EA:25:ILE:HD13	9:EA:153:VAL:HG13	1.82	0.61
1:AC:25:ARG:CD	6:BD:42:GLU:HG3	2.30	0.61
1:AJ:115:MET:HE1	10:AJ:200:CYC:CMB	2.29	0.61
4:AK:51:ILE:HA	4:AK:137:MET:HE1	1.82	0.61
1:AZ:38:ARG:O	1:AZ:42:THR:HG23	2.00	0.61
6:BD:792:TYR:HE2	1:CY:13:ALA:HA	1.65	0.61
1:BK:33:GLY:HA2	1:BK:36:ARG:HE	1.65	0.61

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:BQ:71:ASN:ND2	2:BQ:121:VAL:HA	2.14	0.61
1:BV:64:ASP:HA	1:BV:67:SER:HB2	1.83	0.61
1:BV:94:TYR:OH	2:BW:17:LYS:O	2.17	0.61
2:CH:51:ILE:HD13	2:CH:137:THR:HG22	1.83	0.61
5:CJ:9:CYS:HB2	5:CJ:26:TYR:CD1	2.35	0.61
1:CY:124:GLU:OE1	1:CY:124:GLU:N	2.31	0.61
2:DB:2:GLN:NE2	2:DB:7:ALA:HA	2.15	0.61
8:DH:237:ALA:O	8:DH:240:ILE:HG22	1.99	0.61
1:DQ:5:THR:HA	1:DQ:8:ILE:HG22	1.82	0.61
9:EA:37:LEU:HG	11:EA:400:45D:H291	1.83	0.61
9:EA:223:LEU:HB3	9:EA:231:ILE:HG22	1.82	0.61
1:AX:81:CYS:O	1:AX:85:MET:HG2	2.01	0.61
6:BD:702:THR:O	6:BD:706:ILE:HG12	2.00	0.61
2:CD:111:GLY:HA2	2:CD:114:GLU:OE2	1.99	0.61
2:CH:76:ARG:NH2	10:CH:200:CYC:O2D	2.31	0.61
6:CM:377:VAL:O	6:CM:381:VAL:HG22	2.00	0.61
6:CM:576:ARG:NH2	6:CM:640:ILE:O	2.26	0.61
1:CR:85:MET:HB3	1:CR:133:MET:HE3	1.81	0.61
2:CW:57:ALA:HA	2:CW:61:LEU:HD12	1.81	0.61
10:CY:200:CYC:CBA	2:DD:61:LEU:HD22	2.30	0.61
1:DA:27:LYS:HE3	2:DB:35:GLU:OE1	2.01	0.61
9:DI:223:LEU:HB3	9:DI:231:ILE:HG22	1.82	0.61
2:DM:6:THR:HG22	2:DM:10:ASN:HD21	1.65	0.61
1:DU:27:LYS:HZ3	2:DV:35:GLU:HA	1.65	0.61
9:EA:142:ILE:HD11	9:EA:153:VAL:HG11	1.81	0.61
1:AC:113:ARG:HH22	1:AH:117:ARG:HD3	1.66	0.61
1:AH:105:GLU:HA	1:AH:109:LEU:CB	2.30	0.61
2:AI:8:VAL:HG21	2:AI:27:LEU:CD1	2.26	0.61
1:AJ:32:GLY:O	1:AJ:36:ARG:HG3	2.00	0.61
4:AK:61:GLU:OE1	4:AK:61:GLU:N	2.33	0.61
4:AK:97:LEU:CD2	4:AK:153:ILE:HA	2.30	0.61
4:AK:109:VAL:HG11	4:AK:160:MET:HE2	1.81	0.61
1:AP:81:CYS:HA	10:AP:200:CYC:HHD	1.82	0.61
6:BD:195:THR:CG2	10:BD:902:CYC:H2C	2.29	0.61
2:BQ:114:GLU:HG3	6:CM:472:ARG:NH1	2.15	0.61
2:CA:63:SER:O	2:CA:67:ARG:HG3	2.00	0.61
1:CR:95:GLY:CA	1:CR:104:ILE:HD11	2.30	0.61
1:CR:127:ALA:HA	1:CR:160:MET:HE1	1.82	0.61
1:CY:116:TYR:HB2	1:CY:123:ILE:HD11	1.81	0.61
2:CZ:127:VAL:O	2:CZ:131:GLN:HG2	1.99	0.61
8:DG:210:LEU:HD22	8:DG:216:MET:HG3	1.82	0.61

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:DR:85:LEU:HD21	10:DR:200:CYC:HBC1	1.80	0.61
9:EA:21:VAL:O	9:EA:25:ILE:HG12	1.99	0.61
1:AC:68:PRO:HA	1:AC:73:TYR:CE1	2.36	0.61
2:AD:108:VAL:O	2:AD:112:LEU:HD23	2.00	0.61
3:AE:35:GLN:O	3:AE:39:ILE:HG12	2.01	0.61
1:AP:128:GLN:NE2	1:AP:132:GLU:OE2	2.28	0.61
2:AU:93:THR:HA	2:AU:149:MET:HE1	1.82	0.61
5:BB:35:ASN:ND2	6:BD:637:GLU:OE2	2.33	0.61
6:BD:14:PRO:HG3	6:BD:401:LEU:CD2	2.31	0.61
6:BD:35:LEU:HD22	6:BD:39:GLU:HB3	1.83	0.61
6:BD:792:TYR:CE2	1:CY:13:ALA:HA	2.36	0.61
1:BK:123:ILE:HG13	1:BK:160:MET:O	2.00	0.61
4:BS:119:SER:OG	6:CM:10:SER:O	2.10	0.61
2:BT:97:LEU:HD13	6:CM:35:LEU:HD12	1.82	0.61
1:BZ:38:ARG:O	1:BZ:42:THR:HG23	2.00	0.61
1:CC:90:ARG:NH2	2:CH:73:TYR:OH	2.33	0.61
2:DM:20:ASP:O	2:DM:24:MET:HG2	2.01	0.61
1:DS:78:THR:O	1:DS:82:LEU:HG	2.00	0.61
2:DT:131:GLN:O	2:DT:135:GLU:HG2	2.00	0.61
1:AR:24:ASP:HA	1:AR:27:LYS:HG2	1.83	0.61
2:AU:71:ASN:HD22	2:AU:121:VAL:HA	1.65	0.61
2:BJ:78:TYR:O	2:BJ:82:ILE:HG12	2.00	0.61
3:BM:50:ILE:HD11	3:BM:140:LEU:HB2	1.80	0.61
1:BZ:97:VAL:HG11	2:CA:27:LEU:HD13	1.83	0.61
1:CC:128:GLN:O	1:CC:132:GLU:HG2	2.00	0.61
1:CG:128:GLN:NE2	1:CG:132:GLU:OE2	2.33	0.61
6:CM:447:TYR:OH	10:CM:901:CYC:O2A	2.10	0.61
2:CS:1:MET:N	5:DF:67:ALA:O	2.23	0.61
1:CT:105:GLU:HA	1:CT:109:LEU:HB2	1.81	0.61
2:CU:131:GLN:O	2:CU:135:GLU:HG2	2.00	0.61
1:DC:46:SER:O	1:DC:50:ILE:HG12	1.99	0.61
2:DR:71:ASN:HD22	2:DR:121:VAL:HA	1.65	0.61
9:EA:21:VAL:CG2	9:EA:22:PRO:HD3	2.27	0.61
9:EA:293:ASN:OD1	9:EA:299:PHE:HB2	2.01	0.61
1:AA:79:ALA:O	1:AA:82:LEU:HG	2.00	0.61
1:AA:85:MET:HE2	10:AA:200:CYC:HBC1	1.81	0.61
2:AL:8:VAL:CG1	2:AL:23:ALA:HB1	2.31	0.61
1:AR:86:ASP:OD1	2:AS:18:TYR:OH	2.10	0.61
9:BH:288:TRP:HB3	9:BH:290:PHE:CE1	2.35	0.61
10:BJ:200:CYC:HAA1	6:CM:311:ASN:HB3	1.83	0.61
1:BV:46:SER:HB3	1:BV:50:ILE:HG13	1.81	0.61

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:CG:128:GLN:O	1:CG:132:GLU:HG2	2.01	0.61
6:CM:274:SER:N	6:CM:277:GLU:OE2	2.33	0.61
6:CM:769:ASN:HB2	10:DK:200:CYC:HBA1	1.82	0.61
6:CM:774:VAL:O	6:CM:778:ILE:HG12	2.01	0.61
1:CR:92:VAL:HG11	1:CR:153:PHE:CE1	2.36	0.61
2:CZ:126:THR:O	2:CZ:130:ILE:HD12	2.00	0.61
1:DQ:94:TYR:CD1	2:DR:9:ILE:HD12	2.34	0.61
1:DQ:105:GLU:HA	1:DQ:109:LEU:HD12	1.81	0.61
2:AB:78:TYR:O	2:AB:82:ILE:HG12	2.01	0.61
1:AC:38:ARG:O	1:AC:42:THR:HG23	2.01	0.61
1:AX:5:THR:HG23	2:AY:1:MET:SD	2.40	0.61
1:AZ:134:LYS:HD3	1:AZ:153:PHE:HB3	1.81	0.61
2:BN:104:LEU:HD11	2:BN:152:TYR:HB3	1.82	0.61
4:BS:97:LEU:HD21	4:BS:153:ILE:HA	1.82	0.61
6:CM:191:SER:O	6:CM:195:THR:HG23	2.00	0.61
6:CM:751:LEU:CD1	6:CM:755:ILE:HD13	2.30	0.61
9:CQ:191:GLU:H	9:CQ:254:ARG:HA	1.66	0.61
9:CQ:238:LEU:O	9:CQ:242:ARG:HG2	2.01	0.61
1:DJ:129:SER:O	1:DJ:133:MET:HG3	2.01	0.61
1:DN:65:ILE:HD12	10:DN:200:CYC:HBC2	1.80	0.61
1:DQ:132:GLU:O	1:DQ:136:VAL:HG23	2.00	0.61
2:DR:41:ALA:N	2:DR:96:MET:HE1	2.16	0.61
2:AB:106:GLU:HB2	5:BA:57:THR:CG2	2.31	0.61
2:AO:50:THR:O	2:AO:54:GLU:HG3	2.01	0.61
2:AY:71:ASN:ND2	2:AY:121:VAL:HA	2.16	0.61
1:AZ:91:LEU:HB3	1:AZ:104:ILE:HG23	1.82	0.61
1:AZ:131:ARG:HA	1:AZ:157:ILE:HD11	1.83	0.61
3:BM:52:ASP:O	3:BM:55:GLN:HG3	2.01	0.61
2:BQ:77:ARG:HG3	10:BQ:200:CYC:HAD1	1.81	0.61
6:CM:208:ASP:O	6:CM:211:ARG:HD2	1.99	0.61
6:CM:244:PRO:HG2	6:CM:248:ILE:HD13	1.81	0.61
1:DA:56:ASP:OD1	1:DA:60:GLN:NE2	2.33	0.61
9:DI:205:LEU:HG	9:DI:210:PHE:CE1	2.35	0.61
2:DO:104:LEU:HD13	2:DO:156:ILE:HG13	1.82	0.61
2:AD:72:MET:HE1	2:AD:81:CYS:HB3	1.82	0.61
10:AI:200:CYC:O2A	6:BD:358:ARG:NH1	2.30	0.61
2:AL:18:TYR:CD2	6:BD:61:THR:HG22	2.36	0.61
2:AL:50:THR:O	2:AL:54:GLU:HG2	1.99	0.61
2:AO:4:ALA:O	2:AO:8:VAL:HG23	2.00	0.61
6:BD:2:SER:O	6:BD:4:LYS:NZ	2.33	0.61
6:BD:797:ASN:O	6:BD:801:ILE:HG12	2.01	0.61

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
7:BF:114:ALA:O	7:BF:118:LYS:HG2	2.01	0.61
1:BI:105:GLU:HG3	1:BI:110:VAL:HG23	1.82	0.61
2:BL:103:ILE:HG23	2:BL:104:LEU:HD22	1.83	0.61
3:BM:5:SER:HB3	2:BN:3:ASP:OD2	2.01	0.61
4:BS:97:LEU:CD2	4:BS:153:ILE:HA	2.31	0.61
2:BW:62:TYR:OH	10:BX:200:CYC:O2D	2.15	0.61
2:CF:112:LEU:HD11	10:CF:200:CYC:HMB3	1.83	0.61
1:CG:38:ARG:HA	1:CG:41:GLU:OE1	2.00	0.61
7:CO:105:PRO:HA	7:CO:108:ASN:OD1	2.00	0.61
9:CQ:84:CYS:SG	9:CQ:176:VAL:HG12	2.40	0.61
1:CT:48:GLU:O	1:CT:52:LYS:HG2	2.00	0.61
1:CT:143:SER:OG	1:CT:144:ASP:N	2.31	0.61
2:CU:47:ASN:ND2	2:CU:140:LEU:HD21	2.16	0.61
1:DA:98:SER:HA	2:DB:5:ILE:HD11	1.83	0.61
2:DD:112:LEU:HD13	10:DD:200:CYC:CMB	2.31	0.61
1:DS:50:ILE:HD11	1:DS:137:ALA:HB2	1.82	0.61
2:AB:1:MET:HG3	2:AB:103:ILE:HB	1.83	0.60
3:AE:65:TYR:HA	3:AE:70:GLY:HA3	1.83	0.60
3:AE:92:VAL:HG21	3:AE:153:PHE:CE1	2.36	0.60
1:AR:129:SER:O	1:AR:133:MET:HG3	2.01	0.60
2:AS:87:TYR:OH	6:BD:628:SER:HB2	2.00	0.60
1:AV:4:VAL:O	1:AV:8:ILE:HG12	2.00	0.60
9:BH:188:VAL:HG21	9:BH:252:PRO:HG2	1.83	0.60
9:BH:273:VAL:HB	9:BH:286:ILE:HD12	1.83	0.60
1:BI:125:ALA:HB3	10:BI:200:CYC:HMC2	1.83	0.60
1:BZ:4:VAL:HG11	2:CA:98:ALA:HA	1.83	0.60
6:CM:131:PRO:HD2	6:CM:200:GLN:HG2	1.81	0.60
1:CR:154:ASP:HB3	2:CZ:46:ALA:HB2	1.82	0.60
1:CY:58:LEU:HD22	1:CY:129:SER:HB3	1.83	0.60
2:DD:72:MET:HE2	2:DD:78:TYR:CE2	2.36	0.60
2:DK:73:TYR:O	2:DK:77:ARG:HB2	2.02	0.60
10:DM:200:CYC:O2A	5:DX:17:ARG:NH2	2.28	0.60
5:DX:2:ARG:H	5:DX:57:THR:HG22	1.65	0.60
2:AD:112:LEU:HA	2:AD:115:THR:HG22	1.83	0.60
1:AH:115:MET:HG3	2:AL:78:TYR:CD1	2.36	0.60
2:AL:126:THR:O	2:AL:130:ILE:HG13	2.01	0.60
1:AP:48:GLU:OE1	1:AP:48:GLU:N	2.29	0.60
1:BR:128:GLN:O	1:BR:132:GLU:HG3	2.00	0.60
1:BV:98:SER:HB3	2:BW:9:ILE:CD1	2.31	0.60
1:BX:115:MET:HE1	10:BX:200:CYC:CMB	2.30	0.60
2:CH:15:GLN:HG3	2:CH:17:LYS:CE	2.30	0.60

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
9:CQ:188:VAL:HG11	9:CQ:202:MET:CE	2.31	0.60
2:CU:20:ASP:O	2:CU:24:MET:HG2	1.99	0.60
2:DO:93:THR:CA	2:DO:149:MET:HE1	2.31	0.60
2:DR:113:LYS:O	2:DR:117:ASN:ND2	2.28	0.60
1:DU:39:ILE:HG12	1:DU:145:ASP:HB3	1.82	0.60
1:DU:97:VAL:HG21	2:DV:19:LEU:HD12	1.83	0.60
2:DV:109:LEU:HD13	2:DV:159:GLY:HA3	1.83	0.60
2:DV:127:VAL:O	2:DV:131:GLN:HG3	1.99	0.60
4:AK:113:LEU:HD21	10:AK:200:CYC:CMB	2.30	0.60
1:AZ:116:TYR:HE2	1:AZ:126:VAL:HG11	1.65	0.60
5:BA:2:ARG:NH1	5:BA:33:TYR:HB3	2.15	0.60
9:BH:116:LEU:HB3	9:BH:122:VAL:HG23	1.83	0.60
1:CE:4:VAL:HG12	1:CE:26:ILE:HG23	1.84	0.60
6:CM:18:GLN:HB2	6:CM:173:ASN:HD22	1.66	0.60
1:CR:68:PRO:HA	1:CR:73:TYR:CD2	2.37	0.60
1:CY:96:VAL:HG22	1:CY:152:TYR:CD2	2.36	0.60
2:DT:61:LEU:HD22	10:DU:200:CYC:HBA2	1.82	0.60
1:AJ:89:LEU:O	1:AJ:93:THR:HG23	2.00	0.60
2:AW:74:THR:HG23	1:AX:107:ILE:HA	1.83	0.60
1:AZ:68:PRO:HG2	1:DN:66:VAL:HG21	1.84	0.60
5:BA:9:CYS:HB2	5:BA:26:TYR:CD1	2.36	0.60
6:BD:843:GLU:OE2	6:BD:857:ARG:NH2	2.35	0.60
1:BR:122:PRO:HG2	1:BR:125:ALA:HB3	1.82	0.60
6:CM:796:PRO:CG	2:DR:110:ASN:HB3	2.31	0.60
6:CM:836:GLY:O	6:CM:840:ASN:ND2	2.34	0.60
1:CV:50:ILE:HG12	1:CV:136:VAL:HG12	1.83	0.60
1:CY:127:ALA:CB	1:CY:161:SER:HB2	2.32	0.60
1:CY:129:SER:O	1:CY:133:MET:HG3	2.02	0.60
2:CZ:51:ILE:HA	2:CZ:136:VAL:CG1	2.30	0.60
1:DJ:88:TYR:O	1:DJ:92:VAL:HG23	2.02	0.60
2:DK:37:ARG:HD2	2:DK:96:MET:O	2.00	0.60
1:DL:66:VAL:HG12	1:DL:78:THR:HG22	1.84	0.60
1:DQ:22:GLU:O	1:DQ:26:ILE:HG12	2.01	0.60
1:DS:27:LYS:NZ	2:DT:35:GLU:OE1	2.33	0.60
2:DV:51:ILE:HG12	2:DV:140:LEU:HD13	1.83	0.60
9:EA:214:ILE:O	9:EA:234:LYS:NZ	2.26	0.60
3:AE:101:LYS:HD2	3:AE:155:TYR:CZ	2.37	0.60
4:AK:76:THR:HG23	10:BD:902:CYC:OB	2.02	0.60
2:AW:96:MET:HG2	2:AW:148:GLU:OE1	2.01	0.60
1:BK:14:GLU:OE2	6:CM:263:LYS:HB2	2.01	0.60
1:BK:71:ASN:HB3	10:BK:200:CYC:HMD2	1.83	0.60

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:BT:114:GLU:HA	6:CM:466:THR:O	2.02	0.60
1:BZ:122:PRO:HG2	1:BZ:125:ALA:HB3	1.83	0.60
2:CA:147:LYS:HE3	7:CO:93:PRO:HG3	1.82	0.60
1:CE:94:TYR:OH	2:CF:17:LYS:O	2.20	0.60
2:CS:71:ASN:ND2	2:CS:120:GLY:O	2.34	0.60
1:CV:104:ILE:HG22	1:CV:109:LEU:HD23	1.81	0.60
1:DA:58:LEU:HD22	1:DA:129:SER:CB	2.32	0.60
1:DA:78:THR:O	1:DA:82:LEU:HG	2.01	0.60
9:DI:21:VAL:HG23	9:DI:153:VAL:CG2	2.31	0.60
1:AC:94:TYR:OH	2:AD:17:LYS:O	2.18	0.60
2:AF:126:THR:O	2:AF:130:ILE:HG12	2.01	0.60
4:AK:97:LEU:HD21	4:AK:153:ILE:HA	1.82	0.60
1:AZ:128:GLN:O	1:AZ:132:GLU:HG2	2.02	0.60
6:BD:458:PHE:HB3	6:BD:552:GLY:HA3	1.83	0.60
6:BD:551:PHE:CE2	6:BD:555:LEU:HD11	2.37	0.60
9:BH:222:ALA:HB2	9:BH:232:VAL:HG12	1.82	0.60
1:BI:58:LEU:HD22	1:BI:129:SER:CB	2.31	0.60
1:BV:33:GLY:HA2	1:BV:36:ARG:CD	2.32	0.60
1:BZ:83:ARG:NH2	10:BZ:200:CYC:O2A	2.29	0.60
1:BZ:156:VAL:O	1:BZ:160:MET:HG3	2.01	0.60
6:CM:195:THR:CG2	10:CM:902:CYC:H2C	2.25	0.60
9:CQ:219:SER:HA	9:CQ:234:LYS:HD2	1.84	0.60
2:CS:66:THR:CG2	10:CT:200:CYC:HMA2	2.31	0.60
2:DB:3:ASP:OD1	2:DB:6:THR:OG1	2.12	0.60
1:DC:50:ILE:HD11	1:DC:140:LEU:HD21	1.84	0.60
1:DQ:112:VAL:O	1:DQ:115:MET:HB2	2.02	0.60
2:AD:61:LEU:HD22	10:AE:200:CYC:HBA2	1.82	0.60
2:AL:65:VAL:HG12	2:AL:72:MET:HB3	1.83	0.60
1:AN:92:VAL:HG21	1:AN:153:PHE:CE1	2.36	0.60
1:AR:38:ARG:O	1:AR:42:THR:HG23	2.02	0.60
2:AU:126:THR:O	2:AU:130:ILE:HG13	2.00	0.60
2:BL:105:ASP:HA	2:BL:109:LEU:HB2	1.83	0.60
2:BW:71:ASN:HD22	2:BW:121:VAL:HA	1.65	0.60
1:CG:107:ILE:HG13	6:CM:531:SER:HB2	1.83	0.60
2:CH:30:TYR:OH	2:CH:97:LEU:O	2.18	0.60
9:CQ:74:MET:O	9:CQ:79:GLN:NE2	2.34	0.60
1:CV:26:ILE:O	1:CV:30:VAL:HG22	2.02	0.60
1:DA:76:GLU:OE1	8:DG:202:ARG:HA	2.00	0.60
1:DC:12:ASP:OD2	2:DD:107:ARG:NH1	2.34	0.60
2:DD:44:ILE:CD1	2:DD:149:MET:HE3	2.32	0.60
9:DI:205:LEU:HG	9:DI:210:PHE:HE1	1.66	0.60

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:DK:148:GLU:O	2:DK:151:VAL:HG12	2.00	0.60
2:AI:89:LEU:O	2:AI:93:THR:HG23	2.02	0.60
2:AU:19:LEU:CD1	1:AZ:97:VAL:HG21	2.31	0.60
1:BI:122:PRO:HD2	10:BI:200:CYC:OC	2.02	0.60
2:BL:109:LEU:CD2	2:BL:159:GLY:HA3	2.31	0.60
3:BM:92:VAL:HG21	3:BM:153:PHE:CE1	2.37	0.60
1:BR:58:LEU:N	1:BR:132:GLU:OE1	2.34	0.60
2:BT:141:VAL:HG12	2:BT:145:ALA:HB3	1.84	0.60
1:CC:134:LYS:HD3	1:CC:153:PHE:HB3	1.83	0.60
1:CE:11:ALA:HB2	1:CE:18:LEU:HD23	1.84	0.60
1:CY:81:CYS:O	1:CY:85:MET:HG3	2.02	0.60
1:DJ:50:ILE:CG1	1:DJ:140:LEU:HD21	2.31	0.60
10:DL:200:CYC:NC	10:DL:200:CYC:HMD1	2.16	0.60
1:AA:3:ILE:CG2	1:AH:25:ARG:HD3	2.32	0.60
2:AB:8:VAL:CG1	2:AB:23:ALA:HB1	2.32	0.60
1:AC:69:GLY:N	1:AC:73:TYR:OH	2.24	0.60
3:AE:46:ASN:ND2	3:AE:140:LEU:HD11	2.17	0.60
2:AW:108:VAL:HG11	2:AW:156:ILE:CD1	2.28	0.60
1:AX:97:VAL:HG21	2:AY:19:LEU:HD12	1.83	0.60
6:BD:173:ASN:HD21	6:BD:260:ALA:HA	1.67	0.60
6:BD:754:TYR:OH	2:DD:83:ARG:NH1	2.34	0.60
1:BI:72:ALA:HA	1:BI:77:MET:HB3	1.84	0.60
1:BI:134:LYS:HD3	1:BI:153:PHE:HB3	1.83	0.60
3:BM:38:ARG:O	3:BM:42:THR:HG22	2.01	0.60
3:BM:71:ASN:ND2	3:BM:121:VAL:HG22	2.16	0.60
1:BP:2:SER:OG	2:BQ:3:ASP:OD2	2.15	0.60
1:BZ:62:ARG:HG2	1:BZ:65:ILE:HG12	1.83	0.60
6:CM:447:TYR:HB2	6:CM:451:ASN:HD22	1.67	0.60
1:DC:50:ILE:HD13	1:DC:136:VAL:HG12	1.84	0.60
1:DJ:14:GLU:HB3	1:DJ:16:ARG:CD	2.32	0.60
2:DK:72:MET:HE2	2:DK:78:TYR:CD2	2.37	0.60
2:DK:75:THR:HA	1:DL:115:MET:HE3	1.83	0.60
2:DK:96:MET:HA	2:DK:152:TYR:HE2	1.65	0.60
1:DU:29:PHE:O	1:DU:36:ARG:NH2	2.27	0.60
2:DV:148:GLU:O	2:DV:151:VAL:HG12	2.02	0.60
9:EA:68:LEU:HD13	9:EA:112:ARG:HG2	1.84	0.60
9:EA:223:LEU:HD23	9:EA:225:PRO:HD3	1.83	0.60
1:AH:141:MET:HE1	1:AH:149:ALA:HB3	1.84	0.60
1:AJ:97:VAL:HG21	4:AK:19:LEU:CD1	2.32	0.60
2:AS:57:ALA:HA	2:AS:61:LEU:HG	1.84	0.60
6:BD:84:TYR:HD1	6:BD:150:GLN:HE21	1.50	0.60

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
9:BH:84:CYS:SG	9:BH:176:VAL:HG12	2.41	0.60
1:BK:122:PRO:HD2	10:BK:200:CYC:OC	2.01	0.60
1:BK:123:ILE:HA	1:BK:126:VAL:HG22	1.84	0.60
2:BN:147:LYS:O	2:BN:151:VAL:HG23	2.02	0.60
2:BT:65:VAL:HG12	2:BT:72:MET:HB3	1.82	0.60
1:CE:122:PRO:HG2	1:CE:125:ALA:HB3	1.82	0.60
2:CF:56:VAL:HG12	2:CF:61:LEU:HG	1.84	0.60
9:CQ:288:TRP:HB3	9:CQ:290:PHE:CE1	2.37	0.60
1:CR:89:LEU:O	1:CR:93:THR:HG23	2.02	0.60
2:CS:15:GLN:HB2	2:CS:17:LYS:HE3	1.84	0.60
1:CY:62:ARG:NH2	1:CY:64:ASP:OD2	2.34	0.60
1:DA:50:ILE:CD1	1:DA:136:VAL:HG13	2.32	0.60
9:DI:54:ALA:HA	11:DI:400:45D:H221	1.83	0.60
9:DI:134:ASN:O	9:DI:138:VAL:HG23	2.01	0.60
2:DM:127:VAL:HG13	2:DM:157:CYS:SG	2.41	0.60
1:DU:14:GLU:OE1	1:DU:16:ARG:NE	2.33	0.60
1:DU:27:LYS:HE2	2:DV:38:VAL:HG11	1.84	0.60
2:DV:111:GLY:HA2	2:DV:114:GLU:OE2	2.01	0.60
1:AJ:58:LEU:N	1:AJ:132:GLU:OE1	2.34	0.59
1:AN:85:MET:HG3	1:AN:133:MET:SD	2.42	0.59
2:AS:65:VAL:CG1	2:AS:72:MET:HE3	2.32	0.59
1:AV:44:THR:HG22	2:AW:18:TYR:CD2	2.37	0.59
1:AV:104:ILE:HG22	1:AV:109:LEU:HD12	1.84	0.59
1:AX:128:GLN:O	1:AX:132:GLU:HG2	2.02	0.59
6:BD:14:PRO:HG3	6:BD:401:LEU:HD23	1.84	0.59
9:BH:21:VAL:HG13	9:BH:153:VAL:HG23	1.83	0.59
3:BM:40:ALA:HB2	3:BM:97:LEU:HD11	1.83	0.59
1:BV:65:ILE:HD12	10:BV:200:CYC:OC	2.01	0.59
2:CD:61:LEU:HD21	2:CD:78:TYR:OH	2.02	0.59
2:CD:105:ASP:OD2	2:CD:155:TYR:OH	2.14	0.59
6:CM:574:SER:HB2	6:CM:651:GLU:O	2.02	0.59
9:CQ:193:VAL:HG21	9:CQ:269:VAL:HG22	1.84	0.59
2:CS:47:ASN:O	2:CS:51:ILE:HD12	2.02	0.59
2:CU:127:VAL:O	2:CU:131:GLN:HG2	2.01	0.59
1:CY:126:VAL:HG13	1:CY:160:MET:SD	2.41	0.59
1:DJ:101:VAL:HB	1:DJ:155:PHE:CE2	2.37	0.59
1:DL:33:GLY:O	1:DL:37:LEU:HG	2.02	0.59
1:DQ:131:ARG:O	1:DQ:135:GLU:HG3	2.02	0.59
2:DV:90:ARG:NH1	2:DV:94:TYR:OH	2.34	0.59
5:DX:2:ARG:O	5:DX:33:TYR:HB2	2.02	0.59
1:AC:68:PRO:HA	1:AC:73:TYR:HE1	1.67	0.59

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:AD:105:ASP:HA	2:AD:109:LEU:HB2	1.84	0.59
2:AF:51:ILE:HD11	2:AF:140:LEU:HD23	1.84	0.59
4:AK:20:ASP:O	4:AK:24:MET:HG2	2.01	0.59
2:AU:85:LEU:HG	10:AU:200:CYC:HBC1	1.84	0.59
2:BL:64:ASP:OD1	2:BL:67:ARG:NH1	2.33	0.59
10:BL:200:CYC:CAA	5:CJ:11:PRO:HB3	2.32	0.59
2:BQ:89:LEU:O	2:BQ:93:THR:HG23	2.02	0.59
2:BQ:114:GLU:N	2:BQ:114:GLU:OE1	2.35	0.59
2:BT:98:ALA:HB2	6:CM:25:ILE:HD13	1.84	0.59
1:BZ:101:VAL:HG21	1:BZ:155:PHE:CD2	2.37	0.59
1:CC:16:ARG:NH1	1:CC:22:GLU:OE1	2.26	0.59
2:CD:67:ARG:HG3	2:CD:68:PRO:HD2	1.82	0.59
1:CR:106:GLU:HG3	5:DF:66:LEU:CD2	2.33	0.59
2:CS:66:THR:HG21	10:CT:200:CYC:HMA2	1.84	0.59
1:CY:18:LEU:HD12	1:CY:22:GLU:HB3	1.85	0.59
1:DA:67:SER:OG	8:DZ:230:ASN:HB2	2.02	0.59
1:DL:17:TYR:CE1	2:DM:90:ARG:HD3	2.37	0.59
2:DR:20:ASP:O	2:DR:24:MET:HG2	2.01	0.59
2:DV:92:ALA:HB3	2:DV:149:MET:HE1	1.85	0.59
3:AE:26:ILE:O	3:AE:30:LEU:HD23	2.03	0.59
1:AH:93:THR:O	1:AH:97:VAL:HG23	2.02	0.59
2:AO:134:LYS:HA	2:AO:153:LEU:HD13	1.84	0.59
2:AY:30:TYR:OH	2:AY:97:LEU:O	2.18	0.59
9:BH:251:ILE:N	9:BH:272:LYS:O	2.33	0.59
2:BJ:122:PRO:HB2	2:BJ:125:SER:OG	2.02	0.59
2:BL:71:ASN:HD22	2:BL:121:VAL:HA	1.66	0.59
1:CG:97:VAL:HG21	2:CH:19:LEU:CD1	2.33	0.59
6:CM:35:LEU:HD22	6:CM:39:GLU:CB	2.31	0.59
6:CM:758:SER:O	6:CM:761:THR:OG1	2.18	0.59
6:CM:823:ASN:ND2	10:DR:200:CYC:O2A	2.24	0.59
2:CW:55:ALA:O	2:CW:59:SER:OG	2.13	0.59
1:CY:44:THR:HG22	2:CZ:18:TYR:CD2	2.37	0.59
2:CZ:62:TYR:OH	10:DA:200:CYC:O2D	2.16	0.59
1:DA:50:ILE:HD12	1:DA:136:VAL:HG22	1.84	0.59
2:DB:36:LEU:HD11	2:DB:144:ASP:HB2	1.84	0.59
2:DV:71:ASN:O	2:DV:77:ARG:HD3	2.01	0.59
1:AJ:5:THR:HG23	4:AK:1:MET:SD	2.43	0.59
4:AK:29:ALA:O	4:AK:32:GLU:HG3	2.03	0.59
2:AL:15:GLN:NE2	7:BF:97:ASN:O	2.36	0.59
1:AR:152:TYR:O	1:AR:156:VAL:HG23	2.03	0.59
2:BN:96:MET:HA	2:BN:152:TYR:HE2	1.67	0.59

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:BZ:48:GLU:OE1	1:BZ:48:GLU:N	2.30	0.59
1:CE:65:ILE:CG2	1:CE:72:ALA:HB3	2.33	0.59
1:CE:105:GLU:HA	1:CE:109:LEU:HB2	1.85	0.59
9:CQ:21:VAL:HG13	9:CQ:153:VAL:HG23	1.84	0.59
2:CW:111:GLY:O	5:DF:40:GLN:NE2	2.35	0.59
9:DI:19:ASP:C	9:DI:22:PRO:HD2	2.26	0.59
9:DI:21:VAL:CG1	9:DI:22:PRO:HD3	2.31	0.59
2:DO:71:ASN:O	2:DO:77:ARG:NE	2.28	0.59
2:DR:109:LEU:CD1	2:DR:159:GLY:HA3	2.33	0.59
1:AA:132:GLU:O	1:AA:136:VAL:HG23	2.02	0.59
9:BH:229:ARG:HG2	9:EA:227:PHE:HB3	1.84	0.59
3:BM:67:ALA:HB1	3:BM:68:PRO:HD2	1.84	0.59
4:BS:64:PRO:CB	7:CO:105:PRO:HB2	2.32	0.59
2:BW:96:MET:HG3	2:BW:148:GLU:OE1	2.01	0.59
2:CA:88:TYR:OH	2:CA:112:LEU:HD21	2.02	0.59
1:CE:44:THR:HG22	2:CF:18:TYR:CD2	2.37	0.59
5:CK:39:GLU:O	5:CK:43:ILE:HG12	2.02	0.59
6:CM:612:ARG:NE	6:CM:644:GLU:OE2	2.35	0.59
2:CW:53:LYS:NZ	2:CW:54:GLU:OE2	2.36	0.59
1:DJ:108:GLY:HA2	2:DO:75:THR:HG23	1.84	0.59
1:DL:78:THR:O	1:DL:82:LEU:HG	2.02	0.59
1:DQ:71:ASN:HD22	1:DQ:121:THR:HA	1.68	0.59
8:DZ:242:ILE:HG23	8:DZ:246:VAL:CG2	2.31	0.59
9:EA:224:GLN:NE2	9:EA:230:PRO:HG3	2.18	0.59
2:AD:66:THR:HG21	10:AE:200:CYC:HBA1	1.85	0.59
3:AE:32:THR:O	3:AE:36:ARG:HG2	2.03	0.59
2:AO:126:THR:HB	10:AO:200:CYC:HMC2	1.83	0.59
1:AR:35:ALA:O	1:AR:39:ILE:HG13	2.02	0.59
1:AR:89:LEU:O	1:AR:93:THR:HG23	2.03	0.59
2:AU:112:LEU:HD21	10:AU:200:CYC:C4B	2.33	0.59
5:BA:2:ARG:HA	5:BA:57:THR:O	2.03	0.59
6:BD:70:ARG:HG2	6:BD:201:GLU:OE2	2.02	0.59
6:BD:253:LEU:HD12	6:BD:254:PRO:HD2	1.85	0.59
3:BM:30:LEU:CD1	2:BN:31:PHE:HA	2.33	0.59
1:BV:80:THR:O	1:BV:83:ARG:HG2	2.03	0.59
1:BX:95:GLY:HA3	1:BX:104:ILE:HD11	1.84	0.59
1:BX:102:THR:O	1:BX:106:GLU:HG2	2.03	0.59
1:CC:72:ALA:HA	1:CC:77:MET:HB3	1.85	0.59
9:CQ:27:ARG:O	9:CQ:31:LEU:HG	2.02	0.59
2:CS:23:ALA:O	2:CS:27:LEU:HG	2.02	0.59
2:CW:3:ASP:H	2:CW:6:THR:HG1	1.48	0.59

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:CZ:94:TYR:HB3	2:CZ:103:ILE:HD13	1.83	0.59
1:DC:85:MET:HB3	1:DC:133:MET:CE	2.32	0.59
8:DG:215:PRO:HG2	1:DQ:80:THR:HG22	1.84	0.59
9:DI:250:LEU:CD2	9:DI:273:VAL:HG22	2.33	0.59
1:DJ:22:GLU:O	1:DJ:26:ILE:HG12	2.02	0.59
2:DT:71:ASN:HD22	2:DT:122:PRO:HD3	1.66	0.59
2:AD:24:MET:O	2:AD:28:LYS:HG2	2.01	0.59
2:AD:131:GLN:O	2:AD:134:LYS:HB2	2.03	0.59
3:AE:52:ASP:O	3:AE:55:GLN:HG3	2.02	0.59
2:AI:104:LEU:HD22	2:AI:156:ILE:HD11	1.84	0.59
2:AL:1:MET:SD	2:AL:103:ILE:HD13	2.43	0.59
1:AP:89:LEU:O	1:AP:93:THR:HG23	2.02	0.59
1:AV:89:LEU:O	1:AV:93:THR:HG23	2.03	0.59
1:AX:97:VAL:HG21	2:AY:19:LEU:CD1	2.31	0.59
9:BH:286:ILE:HG12	9:BH:305:LEU:HD23	1.85	0.59
1:BK:47:ARG:O	1:BK:51:VAL:HG23	2.03	0.59
1:BP:83:ARG:NH2	10:BP:200:CYC:O2A	2.36	0.59
1:BP:89:LEU:HB2	1:BP:133:MET:CE	2.33	0.59
1:BV:108:GLY:HA2	2:CA:75:THR:CG2	2.32	0.59
2:BY:75:THR:HG23	1:BZ:108:GLY:HA2	1.84	0.59
2:CA:91:TYR:OH	2:CA:107:ARG:NE	2.36	0.59
1:CC:56:ASP:O	1:CC:60:GLN:HG2	2.03	0.59
1:CC:76:GLU:O	1:CC:80:THR:HG23	2.03	0.59
6:CM:317:LYS:NZ	6:CM:381:VAL:O	2.26	0.59
1:CT:17:TYR:HE1	2:CU:90:ARG:HD3	1.67	0.59
1:CV:41:GLU:CG	2:CW:24:MET:HE2	2.32	0.59
1:CY:37:LEU:HG	2:CZ:24:MET:HE3	1.85	0.59
1:DL:72:ALA:O	1:DL:78:THR:HG23	2.03	0.59
1:DQ:14:GLU:OE1	1:DQ:16:ARG:NE	2.25	0.59
2:AB:103:ILE:CD1	2:AB:107:ARG:HD2	2.33	0.59
1:AH:41:GLU:HA	1:AH:44:THR:HG22	1.83	0.59
2:AS:119:LEU:HB3	2:AS:121:VAL:HG23	1.85	0.59
1:AV:53:GLN:O	1:AV:57:ARG:HD3	2.02	0.59
6:BD:35:LEU:HD22	6:BD:39:GLU:CB	2.32	0.59
6:BD:611:GLY:HA3	6:BD:656:TYR:O	2.02	0.59
2:BJ:63:SER:O	2:BJ:67:ARG:HD3	2.03	0.59
3:BM:97:LEU:CD1	2:BN:19:LEU:HD12	2.31	0.59
2:BT:1:MET:SD	2:BT:103:ILE:HD13	2.42	0.59
2:CA:71:ASN:HD22	2:CA:121:VAL:HA	1.68	0.59
2:CD:10:ASN:O	2:CD:14:VAL:HG23	2.03	0.59
1:CG:5:THR:HG23	2:CH:1:MET:SD	2.42	0.59

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
6:CM:210:PHE:CE2	6:CM:212:ASN:HB2	2.38	0.59
6:CM:551:PHE:CE2	6:CM:555:LEU:HD11	2.38	0.59
6:CM:797:ASN:HB2	10:DR:200:CYC:OB	2.02	0.59
1:CR:82:LEU:CD1	8:DZ:242:ILE:HD11	2.33	0.59
2:CW:20:ASP:O	2:CW:24:MET:HG2	2.03	0.59
1:CY:136:VAL:HG12	1:CY:140:LEU:HD23	1.84	0.59
2:DB:66:THR:HG21	10:DC:200:CYC:CBA	2.30	0.59
2:DB:75:THR:HG23	10:DC:200:CYC:HBB2	1.83	0.59
1:DJ:118:SER:O	2:DO:53:LYS:NZ	2.27	0.59
2:DM:15:GLN:HG3	2:DM:17:LYS:NZ	2.17	0.59
1:DN:39:ILE:HG12	1:DN:145:ASP:HB3	1.85	0.59
2:DO:60:LEU:HD13	2:DO:72:MET:SD	2.43	0.59
1:DQ:130:VAL:CG1	1:DQ:157:ILE:HG22	2.29	0.59
2:DR:72:MET:HE3	2:DR:78:TYR:HA	1.85	0.59
5:DX:19:GLN:O	5:DX:21:GLU:HG2	2.03	0.59
1:AC:96:VAL:HA	1:AC:152:TYR:HE2	1.67	0.59
2:AL:144:ASP:OD1	2:AL:145:ALA:N	2.36	0.59
2:AO:83:ARG:HH21	6:BD:490:ALA:HB1	1.68	0.59
2:AU:61:LEU:HD22	10:AV:200:CYC:HBA1	1.84	0.59
1:AZ:89:LEU:O	1:AZ:93:THR:HG23	2.02	0.59
4:BS:76:THR:HG21	6:CM:182:LEU:HB2	1.85	0.59
1:CC:67:SER:O	1:CC:73:TYR:HB2	2.03	0.59
1:CE:89:LEU:O	1:CE:93:THR:HG23	2.03	0.59
2:CH:65:VAL:HA	2:CH:70:GLY:HA3	1.84	0.59
8:DG:204:ILE:HD11	8:DZ:216:MET:SD	2.43	0.59
2:DM:71:ASN:ND2	2:DM:121:VAL:HA	2.18	0.59
1:DQ:79:ALA:HB3	8:DZ:203:PHE:CE1	2.38	0.59
1:DS:72:ALA:HA	1:DS:77:MET:CB	2.33	0.59
2:AO:77:ARG:HB3	10:AO:200:CYC:HMD1	1.85	0.59
10:AU:200:CYC:O2A	5:BB:37:PHE:HB2	2.02	0.59
1:AV:37:LEU:HD21	2:AW:24:MET:HE1	1.84	0.59
1:AX:38:ARG:HA	1:AX:41:GLU:OE1	2.03	0.59
6:BD:139:ILE:HD11	6:BD:191:SER:HB3	1.83	0.59
9:BH:107:LEU:HG	11:BH:400:45D:C30	2.33	0.59
1:BI:72:ALA:HB2	10:BI:200:CYC:HMD3	1.85	0.59
1:BK:38:ARG:O	1:BK:42:THR:HG23	2.03	0.59
2:BN:126:THR:O	2:BN:130:ILE:HG12	2.03	0.59
10:BN:200:CYC:HBA1	5:CJ:33:TYR:CZ	2.38	0.59
1:BP:110:VAL:CG1	2:BT:76:ARG:HG3	2.33	0.59
4:BS:75:THR:HB	6:CM:178:ASN:OD1	2.03	0.59
4:BS:105:LEU:HD11	4:BS:160:MET:CG	2.33	0.59

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:BW:110:ASN:ND2	6:CM:481:LYS:O	2.34	0.59
2:CF:108:VAL:HG12	2:CF:109:LEU:HD23	1.84	0.59
6:CM:389:TYR:OH	6:CM:404:GLU:OE1	2.16	0.59
1:CR:3:ILE:HD12	1:CR:25:ARG:CG	2.32	0.59
1:CR:68:PRO:HG2	8:DZ:230:ASN:HD21	1.67	0.59
2:CW:68:PRO:HA	2:CW:73:TYR:CD2	2.38	0.59
8:DG:215:PRO:CG	1:DQ:80:THR:HG22	2.33	0.59
1:DJ:92:VAL:O	1:DJ:96:VAL:HG23	2.03	0.59
1:DQ:127:ALA:HA	1:DQ:160:MET:HE1	1.84	0.59
2:DT:96:MET:HA	2:DT:152:TYR:CE2	2.38	0.59
9:EA:134:ASN:O	9:EA:138:VAL:HG23	2.03	0.59
1:AH:71:ASN:OD1	10:AH:200:CYC:NC	2.35	0.58
1:AH:112:VAL:CG2	2:AL:75:THR:HG21	2.32	0.58
2:AI:73:TYR:HH	1:AJ:90:ARG:HH22	1.49	0.58
1:AX:131:ARG:HG3	1:AX:157:ILE:CD1	2.33	0.58
1:AZ:67:SER:O	1:AZ:73:TYR:HB2	2.03	0.58
1:AZ:156:VAL:O	1:AZ:160:MET:HG2	2.03	0.58
9:BH:21:VAL:HG13	9:BH:153:VAL:CG2	2.33	0.58
2:BL:113:LYS:HD2	2:BL:160:LEU:O	2.02	0.58
3:BM:30:LEU:HD13	2:BN:31:PHE:HD1	1.68	0.58
2:BT:1:MET:HE2	6:CM:26:SER:OG	2.03	0.58
1:BV:26:ILE:O	1:BV:30:VAL:HG13	2.03	0.58
2:CF:1:MET:HE3	2:CF:103:ILE:HB	1.83	0.58
6:CM:317:LYS:HD3	6:CM:392:GLU:OE2	2.03	0.58
6:CM:709:ARG:HD3	2:DT:69:GLY:O	2.03	0.58
9:CQ:300:PHE:CZ	9:CQ:302:ALA:HB2	2.39	0.58
1:CV:125:ALA:HB3	10:CV:200:CYC:HMC2	1.84	0.58
2:CW:123:ILE:HG23	2:CW:160:LEU:HG	1.85	0.58
1:DA:122:PRO:HG2	1:DA:125:ALA:HB3	1.84	0.58
1:DJ:50:ILE:HA	1:DJ:136:VAL:CG1	2.33	0.58
2:DM:73:TYR:O	2:DM:77:ARG:HB2	2.03	0.58
2:DO:61:LEU:HD21	2:DO:78:TYR:OH	2.03	0.58
2:DR:22:ALA:O	2:DR:26:LYS:HE3	2.03	0.58
2:DT:3:ASP:N	2:DT:6:THR:OG1	2.29	0.58
2:DV:149:MET:HE3	2:DV:153:LEU:HD11	1.85	0.58
2:AL:71:ASN:OD1	10:AL:200:CYC:HMD2	2.03	0.58
1:AX:14:GLU:OE1	1:AX:16:ARG:NE	2.27	0.58
6:BD:543:ILE:HG21	6:BD:565:GLU:HG2	1.85	0.58
2:BN:114:GLU:OE1	5:CJ:52:LYS:HA	2.03	0.58
4:BS:65:GLU:OE1	4:BS:65:GLU:N	2.36	0.58
10:BT:200:CYC:HMD1	10:BT:200:CYC:NC	2.10	0.58

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:CA:58:LYS:HB3	7:CO:98:VAL:CG1	2.33	0.58
2:CF:74:THR:HG23	1:CG:107:ILE:HA	1.84	0.58
2:CH:10:ASN:O	2:CH:14:VAL:HG23	2.04	0.58
2:CS:33:SER:O	2:CS:37:ARG:HG2	2.03	0.58
2:CS:153:LEU:HA	2:CS:156:ILE:HD13	1.85	0.58
2:CU:78:TYR:CD1	1:CV:115:MET:HG3	2.37	0.58
10:CY:200:CYC:HBA2	2:DD:61:LEU:HD22	1.84	0.58
9:DI:188:VAL:N	9:DI:203:ASP:OD1	2.30	0.58
2:DM:88:TYR:CD2	2:DM:130:ILE:HD11	2.38	0.58
1:DS:33:GLY:O	1:DS:37:LEU:HD23	2.02	0.58
8:DZ:242:ILE:HG23	8:DZ:246:VAL:HG21	1.85	0.58
9:EA:249:LYS:O	9:EA:274:GLN:N	2.36	0.58
2:AB:10:ASN:O	2:AB:14:VAL:HG23	2.03	0.58
1:AC:2:SER:HB3	1:AC:5:THR:CG2	2.33	0.58
2:AD:33:SER:O	2:AD:37:ARG:HG3	2.03	0.58
2:AI:2:GLN:HG2	2:AI:6:THR:HG22	1.85	0.58
2:AU:111:GLY:HA2	2:AU:114:GLU:OE2	2.02	0.58
2:AW:47:ASN:OD1	2:AW:140:LEU:HD21	2.03	0.58
1:BK:68:PRO:HA	1:BK:73:TYR:CE1	2.38	0.58
2:BL:85:LEU:HG	10:BL:200:CYC:HBC1	1.85	0.58
3:BM:50:ILE:HD11	3:BM:140:LEU:CB	2.33	0.58
3:BM:101:LYS:HD2	3:BM:155:TYR:CZ	2.37	0.58
1:BV:64:ASP:HB2	1:CE:64:ASP:HB2	1.85	0.58
2:BW:71:ASN:HD22	2:BW:122:PRO:HD3	1.68	0.58
1:BZ:81:CYS:HA	10:BZ:200:CYC:HHD	1.85	0.58
1:BZ:84:ASP:OD2	10:BZ:200:CYC:ND	2.32	0.58
2:CA:123:ILE:HA	2:CA:126:THR:HG22	1.84	0.58
1:CG:62:ARG:O	1:CG:65:ILE:HG12	2.03	0.58
6:CM:309:VAL:HG21	6:CM:319:PHE:CG	2.38	0.58
1:CR:148:GLU:OE1	1:CR:148:GLU:N	2.37	0.58
2:CZ:91:TYR:CE2	10:CZ:200:CYC:HBB1	2.39	0.58
1:DC:8:ILE:CD1	2:DD:98:ALA:HB2	2.33	0.58
9:DI:223:LEU:HG	9:DI:301:VAL:HG13	1.85	0.58
9:EA:250:LEU:CD2	9:EA:273:VAL:HG22	2.33	0.58
4:AK:108:ARG:HB3	6:BD:249:GLN:OE1	2.03	0.58
2:AS:58:LYS:HB3	7:BF:98:VAL:HG11	1.85	0.58
1:AX:107:ILE:HD11	6:BD:532:VAL:CG2	2.29	0.58
9:BH:298:ILE:HD11	9:BH:300:PHE:O	2.03	0.58
2:BT:122:PRO:HB2	2:BT:125:SER:HB2	1.85	0.58
6:CM:154:ARG:NH1	6:CM:155:ASP:OD1	2.37	0.58
6:CM:485:ILE:HD12	6:CM:665:LEU:HD13	1.85	0.58

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:DD:1:MET:N	2:DD:106:GLU:OE1	2.27	0.58
2:DD:61:LEU:HA	2:DD:66:THR:HG21	1.86	0.58
2:DK:2:GLN:N	2:DK:102:SER:OG	2.35	0.58
2:DR:112:LEU:HA	2:DR:115:THR:HG22	1.85	0.58
1:DS:58:LEU:HD22	1:DS:129:SER:CB	2.34	0.58
1:DU:86:ASP:OD1	2:DV:18:TYR:OH	2.19	0.58
9:EA:218:THR:HG23	9:EA:297:LYS:O	2.02	0.58
2:AD:96:MET:HB2	2:AD:148:GLU:OE1	2.03	0.58
1:AH:89:LEU:HB2	1:AH:133:MET:CE	2.33	0.58
2:AU:61:LEU:HD21	2:AU:78:TYR:OH	2.04	0.58
6:BD:860:THR:OG1	10:DB:200:CYC:O1A	2.21	0.58
2:BJ:137:THR:O	2:BJ:141:VAL:HG12	2.04	0.58
2:BL:10:ASN:O	2:BL:14:VAL:HG13	2.03	0.58
1:BP:87:TYR:HB3	10:BP:200:CYC:HBB3	1.85	0.58
1:BV:37:LEU:HG	2:BW:24:MET:HG3	1.84	0.58
2:CA:104:LEU:HD12	2:CA:155:TYR:HD2	1.69	0.58
2:CD:75:THR:CG2	1:CE:108:GLY:HA2	2.33	0.58
6:CM:555:LEU:HD22	6:CM:587:LEU:HD21	1.84	0.58
9:CQ:250:LEU:CD2	9:CQ:273:VAL:HG23	2.34	0.58
2:CW:84:ASP:OD2	10:CW:200:CYC:ND	2.36	0.58
2:DD:57:ALA:HA	2:DD:61:LEU:CD1	2.31	0.58
2:DD:127:VAL:O	2:DD:131:GLN:HG2	2.04	0.58
2:DT:122:PRO:HB2	2:DT:125:SER:HB2	1.84	0.58
2:DT:124:SER:O	2:DT:128:GLN:HG2	2.04	0.58
1:DU:27:LYS:HD2	2:DV:35:GLU:OE1	2.04	0.58
3:AE:97:LEU:CD1	2:AF:19:LEU:HD12	2.33	0.58
1:AJ:109:LEU:HD13	1:AJ:159:LYS:CG	2.34	0.58
2:AL:2:GLN:HB2	2:AL:6:THR:CG2	2.33	0.58
1:AP:122:PRO:HG2	1:AP:125:ALA:HB3	1.85	0.58
1:AR:101:VAL:HG22	1:AR:104:ILE:HB	1.85	0.58
2:AW:76:ARG:HB2	1:AX:110:VAL:CG1	2.32	0.58
6:BD:864:ALA:C	6:BD:867:PRO:HD2	2.28	0.58
1:BP:81:CYS:HA	10:BP:200:CYC:HHD	1.85	0.58
2:BQ:77:ARG:HG2	10:BQ:200:CYC:HMD1	1.85	0.58
1:BR:32:GLY:O	1:BR:36:ARG:HG3	2.03	0.58
2:BT:113:LYS:NZ	2:CA:105:ASP:OD2	2.30	0.58
1:BV:122:PRO:O	1:BV:126:VAL:HG23	2.04	0.58
10:CE:200:CYC:HHA	10:CE:200:CYC:HBD1	1.86	0.58
1:CG:51:VAL:HG22	1:CG:133:MET:CE	2.33	0.58
1:CG:105:GLU:HA	1:CG:109:LEU:CB	2.30	0.58
6:CM:253:LEU:HD12	6:CM:254:PRO:HD2	1.85	0.58

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
6:CM:477:ASN:O	6:CM:617:ARG:HD3	2.03	0.58
1:CR:85:MET:HB3	1:CR:133:MET:CE	2.34	0.58
1:DC:121:THR:HG23	10:DC:200:CYC:C1C	2.32	0.58
1:DJ:76:GLU:OE1	1:DJ:76:GLU:N	2.32	0.58
1:DL:57:ARG:O	1:DL:61:LYS:HG2	2.03	0.58
1:DL:141:MET:CE	1:DL:146:ALA:HA	2.33	0.58
1:DQ:58:LEU:CD2	10:DQ:200:CYC:HBC3	2.33	0.58
1:DQ:76:GLU:O	1:DQ:80:THR:HG23	2.03	0.58
1:DS:113:ARG:HH11	1:DS:123:ILE:HD13	1.69	0.58
5:DX:2:ARG:HA	5:DX:57:THR:O	2.04	0.58
3:AE:30:LEU:CD1	2:AF:31:PHE:HA	2.34	0.58
3:AE:65:TYR:HA	3:AE:71:ASN:H	1.67	0.58
9:BH:63:LEU:HD12	9:BH:105:ILE:CD1	2.34	0.58
3:BM:112:VAL:O	3:BM:115:MET:HB3	2.04	0.58
2:BY:127:VAL:O	2:BY:131:GLN:HG2	2.04	0.58
1:CC:131:ARG:HG3	1:CC:157:ILE:HD13	1.85	0.58
10:CH:200:CYC:HBD2	10:CH:200:CYC:HHA	1.83	0.58
6:CM:225:ASP:HA	6:CM:228:LEU:CD2	2.34	0.58
7:CO:104:GLY:O	7:CO:111:LYS:NZ	2.35	0.58
2:CS:54:GLU:O	2:CS:58:LYS:HG2	2.04	0.58
2:CS:148:GLU:O	2:CS:151:VAL:HG12	2.02	0.58
1:CT:102:THR:OG1	1:CT:103:PRO:HD3	2.03	0.58
1:CY:17:TYR:OH	2:CZ:89:LEU:HD22	2.03	0.58
2:DB:68:PRO:HD3	1:DC:87:TYR:OH	2.04	0.58
8:DG:219:LEU:HD13	8:DZ:203:PHE:CE2	2.38	0.58
9:EA:99:ALA:HB1	9:EA:164:THR:HG22	1.86	0.58
1:AJ:66:VAL:CG1	1:BX:68:PRO:HG2	2.33	0.58
6:BD:5:ALA:HB3	6:BD:437:TYR:CD1	2.39	0.58
6:BD:160:LEU:O	6:BD:164:THR:HG23	2.04	0.58
2:BL:112:LEU:HD21	10:BL:200:CYC:CAB	2.34	0.58
2:CA:71:ASN:ND2	2:CA:121:VAL:HA	2.19	0.58
2:CD:19:LEU:HB2	2:CD:24:MET:HE2	1.86	0.58
1:CG:131:ARG:HG3	1:CG:157:ILE:CD1	2.34	0.58
6:CM:14:PRO:HG3	6:CM:401:LEU:CD2	2.32	0.58
6:CM:152:SER:CB	10:CM:902:CYC:H3C	2.34	0.58
6:CM:198:ALA:O	6:CM:202:MET:HG3	2.03	0.58
6:CM:264:ARG:O	6:CM:266:LYS:NZ	2.27	0.58
6:CM:318:GLU:OE1	6:CM:321:ARG:NE	2.30	0.58
6:CM:748:GLU:C	6:CM:749:ARG:HD3	2.29	0.58
6:CM:826:LEU:HD23	6:CM:830:GLY:O	2.03	0.58
6:CM:882:GLU:N	6:CM:882:GLU:OE1	2.36	0.58

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
9:CQ:187:LYS:HE2	9:CQ:199:LEU:HD21	1.86	0.58
2:DB:104:LEU:O	2:DB:108:VAL:HB	2.04	0.58
2:DD:65:VAL:HA	2:DD:70:GLY:HA3	1.86	0.58
5:DF:6:ILE:HG21	5:DF:36:TRP:CZ3	2.39	0.58
1:AA:134:LYS:HD3	1:AA:153:PHE:HB3	1.86	0.58
2:AB:13:ASP:OD2	5:BA:63:ASN:ND2	2.34	0.58
2:AB:122:PRO:HB2	2:AB:125:SER:OG	2.04	0.58
3:AE:104:ILE:HG22	3:AE:109:LEU:HG	1.86	0.58
2:AI:114:GLU:OE1	2:AI:114:GLU:N	2.36	0.58
4:AK:21:ARG:HE	4:BS:21:ARG:HE	1.51	0.58
2:AS:71:ASN:ND2	2:AS:121:VAL:HA	2.19	0.58
2:AW:72:MET:HE3	2:AW:81:CYS:HB3	1.84	0.58
2:AW:127:VAL:O	2:AW:131:GLN:HG2	2.04	0.58
5:BA:39:GLU:OE2	5:BA:42:ARG:NH2	2.25	0.58
6:BD:75:ILE:HG13	6:BD:198:ALA:CB	2.33	0.58
6:BD:709:ARG:HD3	2:DB:69:GLY:O	2.03	0.58
3:BM:134:LYS:HD3	3:BM:153:PHE:HB3	1.85	0.58
4:BS:111:GLN:NE2	6:CM:437:TYR:HB3	2.19	0.58
1:BV:4:VAL:O	1:BV:8:ILE:HG12	2.03	0.58
1:BZ:19:SER:OG	1:BZ:22:GLU:HG3	2.04	0.58
1:BZ:126:VAL:HG11	1:BZ:160:MET:HE1	1.85	0.58
9:CQ:289:ARG:NH1	9:DI:306:LEU:HD12	2.19	0.58
1:DC:62:ARG:O	1:DC:65:ILE:HG12	2.03	0.58
5:DF:3:MET:HE2	5:DF:30:LEU:HB3	1.85	0.58
8:DG:215:PRO:HB2	8:DZ:203:PHE:CD1	2.39	0.58
9:DI:107:LEU:HB3	11:DI:400:45D:H141	1.86	0.58
2:DK:2:GLN:HB3	2:DK:7:ALA:HB2	1.84	0.58
2:DT:2:GLN:NE2	2:DT:7:ALA:HA	2.19	0.58
2:DT:63:SER:O	2:DT:67:ARG:HG3	2.04	0.58
1:AC:27:LYS:O	1:AC:31:THR:HG23	2.03	0.58
3:AE:27:GLN:O	3:AE:31:THR:HG23	2.04	0.58
3:AE:104:ILE:HG21	3:AE:156:ILE:HD11	1.86	0.58
10:AH:200:CYC:O2D	2:AL:62:TYR:OH	2.12	0.58
4:AK:51:ILE:HD13	4:AK:138:ILE:HD13	1.86	0.58
1:AN:53:GLN:O	1:AN:57:ARG:HD3	2.04	0.58
1:AN:91:LEU:HD21	1:AN:107:ILE:HB	1.85	0.58
2:AW:147:LYS:O	2:AW:151:VAL:HG23	2.04	0.58
1:AX:50:ILE:HG13	1:AX:136:VAL:CG1	2.34	0.58
1:AZ:76:GLU:O	1:AZ:80:THR:HG23	2.03	0.58
6:BD:85:LEU:CG	6:BD:153:LEU:HD21	2.29	0.58
2:BL:57:ALA:HA	2:BL:61:LEU:CG	2.33	0.58

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:BL:131:GLN:O	2:BL:134:LYS:HB2	2.03	0.58
4:BS:68:ARG:HD2	7:CO:105:PRO:HG2	1.86	0.58
1:BV:39:ILE:CG1	1:BV:145:ASP:HB2	2.34	0.58
2:BY:65:VAL:HG12	2:BY:72:MET:HB2	1.84	0.58
1:CG:14:GLU:OE1	1:CG:16:ARG:NE	2.24	0.58
9:CQ:311:GLU:OE1	9:CQ:312:LEU:HD22	2.04	0.58
2:CU:55:ALA:O	2:CU:59:SER:OG	2.10	0.58
2:DD:85:LEU:HG	10:DD:200:CYC:HBC1	1.85	0.58
1:DL:97:VAL:HG21	2:DM:19:LEU:HD11	1.86	0.58
1:DL:154:ASP:OD2	2:DV:46:ALA:HB2	2.03	0.58
1:DU:39:ILE:HG22	1:DU:96:VAL:HG11	1.84	0.58
1:AC:123:ILE:HA	1:AC:126:VAL:HG22	1.85	0.57
2:AD:109:LEU:CD2	2:AD:159:GLY:HA3	2.34	0.57
2:AS:107:ARG:HD3	6:BD:594:ALA:O	2.04	0.57
9:BH:70:GLU:O	9:BH:74:MET:HG3	2.04	0.57
1:BI:22:GLU:O	1:BI:26:ILE:HG12	2.04	0.57
2:BJ:51:ILE:HG22	2:BJ:133:ILE:CD1	2.33	0.57
2:BQ:107:ARG:HG2	6:CM:337:PHE:O	2.04	0.57
1:BR:115:MET:HE1	10:BR:200:CYC:CMB	2.34	0.57
1:BV:22:GLU:O	1:BV:26:ILE:HG13	2.04	0.57
2:CF:126:THR:HG22	10:CF:200:CYC:HMC1	1.86	0.57
6:CM:199:ILE:HD13	6:CM:227:LEU:CD2	2.34	0.57
2:DD:71:ASN:O	2:DD:77:ARG:HD3	2.04	0.57
4:AK:113:LEU:HA	6:BD:408:CYS:SG	2.44	0.57
1:AN:92:VAL:HG11	1:AN:153:PHE:CE2	2.38	0.57
1:AV:39:ILE:HG12	1:AV:145:ASP:HB3	1.86	0.57
1:AX:62:ARG:O	1:AX:65:ILE:HG12	2.04	0.57
5:BB:15:ARG:NH1	5:BB:17:ARG:HG2	2.19	0.57
9:BH:26:ALA:O	9:BH:30:GLN:HG2	2.03	0.57
1:BI:84:ASP:OD2	10:BI:200:CYC:ND	2.37	0.57
2:BQ:76:ARG:CZ	1:BR:110:VAL:HG11	2.35	0.57
1:BV:33:GLY:HA2	1:BV:36:ARG:HD2	1.85	0.57
2:CD:85:LEU:HD22	2:CD:133:ILE:HD11	1.84	0.57
2:CS:84:ASP:OD2	2:CS:116:TYR:OH	2.15	0.57
2:CS:137:THR:O	2:CS:141:VAL:HG22	2.04	0.57
1:CT:68:PRO:HD2	8:DY:230:ASN:ND2	2.18	0.57
2:CU:148:GLU:O	2:CU:151:VAL:HG12	2.04	0.57
1:CY:126:VAL:HG22	1:CY:160:MET:HE2	1.86	0.57
8:DH:238:GLN:HA	1:DJ:52:LYS:CE	2.33	0.57
9:DI:102:SER:OG	9:DI:105:ILE:HG12	2.04	0.57
9:DI:191:GLU:HB2	9:DI:254:ARG:CB	2.34	0.57

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:DJ:20:PRO:HA	1:DJ:23:LEU:HD21	1.86	0.57
1:DJ:96:VAL:HA	1:DJ:152:TYR:CE2	2.39	0.57
2:DM:71:ASN:HB3	10:DM:200:CYC:OC	2.04	0.57
2:DO:110:ASN:HD22	5:DX:49:LYS:HA	1.68	0.57
1:DQ:37:LEU:HD12	2:DR:24:MET:CE	2.34	0.57
1:AJ:84:ASP:OD2	10:AJ:200:CYC:ND	2.35	0.57
4:AK:1:MET:HE2	4:AK:104:VAL:CB	2.34	0.57
1:AR:141:MET:HB2	1:AR:146:ALA:HB2	1.86	0.57
2:AW:24:MET:HE3	2:AW:28:LYS:CD	2.34	0.57
9:BH:75:GLY:O	9:BH:79:GLN:HG3	2.03	0.57
1:BI:153:PHE:O	1:BI:157:ILE:HD12	2.04	0.57
2:BJ:88:TYR:OH	2:BJ:112:LEU:HD21	2.04	0.57
1:CG:38:ARG:O	1:CG:42:THR:HG23	2.04	0.57
6:CM:63:ASN:O	6:CM:67:ILE:HG12	2.05	0.57
6:CM:70:ARG:HG2	6:CM:201:GLU:OE2	2.05	0.57
9:CQ:188:VAL:HG11	9:CQ:202:MET:HE3	1.86	0.57
1:CV:58:LEU:HD22	1:CV:129:SER:CB	2.33	0.57
2:DB:96:MET:HA	2:DB:152:TYR:CE2	2.39	0.57
1:DN:50:ILE:CG1	1:DN:140:LEU:HD21	2.34	0.57
2:DV:57:ALA:HA	2:DV:61:LEU:CD1	2.35	0.57
1:AV:50:ILE:CD1	1:AV:136:VAL:HG12	2.35	0.57
6:BD:321:ARG:O	6:BD:325:LYS:HG2	2.04	0.57
6:BD:447:TYR:CZ	10:BD:901:CYC:HAA1	2.40	0.57
6:BD:519:LEU:HD13	6:BD:569:GLU:O	2.04	0.57
6:BD:793:ALA:O	2:CZ:107:ARG:HD3	2.04	0.57
9:BH:289:ARG:HB3	9:BH:302:ALA:HB3	1.86	0.57
2:BQ:2:GLN:HG2	2:BQ:6:THR:HG22	1.86	0.57
5:CK:6:ILE:HD13	5:CK:36:TRP:HZ3	1.69	0.57
6:CM:702:THR:HB	2:DB:151:VAL:HG22	1.85	0.57
2:CS:57:ALA:O	8:DY:249:ARG:NH1	2.38	0.57
1:CT:91:LEU:HD21	1:CT:107:ILE:HG22	1.87	0.57
1:CY:58:LEU:HD22	1:CY:129:SER:CB	2.34	0.57
2:CZ:108:VAL:O	2:CZ:112:LEU:HB2	2.04	0.57
1:DC:107:ILE:HG22	10:DC:200:CYC:HBB1	1.85	0.57
1:DQ:94:TYR:CD1	2:DR:9:ILE:HG23	2.40	0.57
2:DR:57:ALA:HA	2:DR:61:LEU:HB2	1.86	0.57
2:DR:71:ASN:ND2	2:DR:121:VAL:HA	2.18	0.57
2:DR:122:PRO:HB2	2:DR:125:SER:HB2	1.86	0.57
2:DT:15:GLN:HG3	2:DT:17:LYS:HG3	1.87	0.57
8:DZ:210:LEU:HD11	8:DZ:214:ASN:HB3	1.84	0.57
2:AB:61:LEU:HD22	10:AC:200:CYC:HAA1	1.87	0.57

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:AC:25:ARG:HB3	6:BD:42:GLU:HA	1.86	0.57
3:AE:124:PRO:HA	3:AE:127:VAL:HG12	1.86	0.57
10:AN:200:CYC:O1D	7:BF:100:ARG:NH2	2.37	0.57
1:AR:48:GLU:OE1	1:AR:48:GLU:N	2.33	0.57
2:AW:78:TYR:O	2:AW:82:ILE:HG12	2.05	0.57
3:BM:85:TYR:CE1	3:BM:129:ALA:HB1	2.40	0.57
2:BN:71:ASN:ND2	2:BN:121:VAL:HA	2.19	0.57
1:BP:41:GLU:HA	1:BP:44:THR:HG22	1.85	0.57
1:BR:97:VAL:HG21	4:BS:19:LEU:CD1	2.34	0.57
6:CM:549:GLN:O	6:CM:608:ARG:HD2	2.04	0.57
9:CQ:46:GLU:HB3	9:CQ:50:THR:CG2	2.34	0.57
9:CQ:75:GLY:O	9:CQ:79:GLN:HG3	2.04	0.57
9:CQ:205:LEU:HD12	9:CQ:210:PHE:CE1	2.38	0.57
9:CQ:264:PHE:CD1	9:CQ:294:PRO:HD3	2.38	0.57
1:CT:130:VAL:HA	1:CT:133:MET:HE3	1.87	0.57
2:CU:15:GLN:CB	2:CU:17:LYS:HE3	2.34	0.57
2:CZ:5:ILE:O	2:CZ:9:ILE:HG12	2.03	0.57
1:DA:72:ALA:HB2	10:DA:200:CYC:HMD3	1.87	0.57
1:DJ:44:THR:O	1:DJ:47:ARG:HG2	2.04	0.57
2:DV:65:VAL:HG12	2:DV:72:MET:CB	2.35	0.57
1:AA:21:GLY:O	1:AA:25:ARG:HG2	2.05	0.57
1:AA:64:ASP:OD1	1:AA:65:ILE:HD12	2.05	0.57
1:AC:25:ARG:CG	6:BD:42:GLU:HG3	2.34	0.57
1:AH:112:VAL:HG22	2:AL:75:THR:HG21	1.86	0.57
4:AK:85:ASP:OD2	10:AK:200:CYC:ND	2.35	0.57
4:AK:109:VAL:O	4:AK:113:LEU:HG	2.04	0.57
1:AR:72:ALA:HA	1:AR:77:MET:HB3	1.86	0.57
1:AV:27:LYS:HD2	2:AW:35:GLU:OE1	2.04	0.57
1:AX:105:GLU:HA	1:AX:109:LEU:CB	2.29	0.57
1:AZ:87:TYR:O	1:AZ:91:LEU:HG	2.04	0.57
6:BD:11:LEU:HD23	6:BD:13:ARG:HD2	1.86	0.57
6:BD:348:ARG:NH2	6:BD:398:LEU:HD11	2.19	0.57
6:BD:736:LYS:HE3	6:BD:740:ARG:CG	2.35	0.57
6:BD:806:LYS:HB2	6:BD:872:LEU:HD13	1.86	0.57
3:BM:27:GLN:O	3:BM:31:THR:HG23	2.04	0.57
1:BP:71:ASN:OD1	10:BP:200:CYC:NC	2.38	0.57
2:BW:75:THR:HG23	10:BX:200:CYC:HAB2	1.86	0.57
1:CE:83:ARG:NH1	10:CE:200:CYC:O1A	2.31	0.57
10:CW:200:CYC:HMA2	5:DF:37:PHE:HD1	1.69	0.57
2:CZ:22:ALA:HA	2:CZ:25:ASP:OD2	2.05	0.57
2:CZ:134:LYS:HG3	2:CZ:150:GLY:HA2	1.85	0.57

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
9:DI:213:LEU:HD13	9:DI:216:LEU:HB3	1.86	0.57
1:DN:32:GLY:O	1:DN:36:ARG:HD3	2.05	0.57
2:DO:88:TYR:OH	2:DO:112:LEU:HD21	2.04	0.57
1:DS:22:GLU:O	1:DS:26:ILE:HG12	2.05	0.57
1:DS:76:GLU:OE1	1:DS:76:GLU:N	2.32	0.57
1:DS:131:ARG:O	1:DS:135:GLU:HG2	2.04	0.57
2:AD:3:ASP:OD1	2:AD:98:ALA:HB1	2.05	0.57
3:AE:47:GLU:HG2	3:AE:89:LEU:HD23	1.86	0.57
1:AN:154:ASP:HB3	2:AU:46:ALA:HB2	1.85	0.57
9:BH:201:TYR:CD1	9:BH:213:LEU:HD11	2.40	0.57
1:BI:126:VAL:O	1:BI:130:VAL:HG23	2.05	0.57
2:BL:50:THR:O	2:BL:54:GLU:HG2	2.05	0.57
3:BM:4:VAL:HG13	2:BN:97:LEU:CD2	2.33	0.57
2:BN:85:LEU:HG	10:BN:200:CYC:HBC1	1.87	0.57
2:BQ:76:ARG:HD3	6:CM:362:GLN:OE1	2.04	0.57
2:BT:148:GLU:O	2:BT:151:VAL:HG22	2.05	0.57
1:BV:108:GLY:HA2	2:CA:75:THR:HG23	1.85	0.57
2:CA:123:ILE:HD12	6:CM:9:SER:O	2.05	0.57
2:CF:71:ASN:HD22	2:CF:122:PRO:HD3	1.70	0.57
5:CJ:18:THR:HG22	6:CM:240:VAL:O	2.03	0.57
9:CQ:188:VAL:HG21	9:CQ:252:PRO:HG2	1.87	0.57
2:DD:143:ALA:O	2:DD:147:LYS:HG2	2.05	0.57
2:DM:47:ASN:ND2	2:DM:140:LEU:HD21	2.20	0.57
2:DR:87:TYR:CD2	10:DR:200:CYC:HBB3	2.39	0.57
8:DY:211:THR:O	8:DY:217:ASN:ND2	2.38	0.57
1:AC:128:GLN:OE1	1:AC:131:ARG:HD2	2.05	0.57
3:AE:92:VAL:HG11	3:AE:153:PHE:CE2	2.38	0.57
2:AI:2:GLN:NE2	2:AI:10:ASN:OD1	2.38	0.57
2:AO:1:MET:SD	2:AO:103:ILE:HD12	2.45	0.57
1:AR:126:VAL:O	1:AR:130:VAL:HG23	2.04	0.57
2:AS:100:ASP:OD1	2:AS:101:ALA:N	2.37	0.57
1:AZ:58:LEU:HD22	1:AZ:129:SER:CB	2.35	0.57
5:BA:2:ARG:CZ	5:BA:33:TYR:HB3	2.35	0.57
9:BH:190:ILE:HD13	9:BH:252:PRO:CB	2.35	0.57
1:BI:100:ASP:CG	1:BP:21:GLY:HA3	2.30	0.57
1:BI:132:GLU:O	1:BI:136:VAL:HG23	2.04	0.57
1:BK:50:ILE:CD1	1:BK:136:VAL:HG12	2.35	0.57
1:BV:8:ILE:HD12	2:BW:94:TYR:HD1	1.69	0.57
1:BV:90:ARG:NH1	2:BW:16:GLY:O	2.38	0.57
1:BX:66:VAL:HG12	1:DC:68:PRO:HG2	1.87	0.57
1:BX:89:LEU:O	1:BX:93:THR:HG23	2.05	0.57

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:CA:68:PRO:HA	2:CA:73:TYR:CG	2.39	0.57
2:CH:96:MET:HG3	2:CH:148:GLU:CD	2.30	0.57
6:CM:625:ASP:OD1	6:CM:629:LYS:HE2	2.05	0.57
1:CR:101:VAL:HA	1:CR:104:ILE:CD1	2.35	0.57
2:CW:68:PRO:HA	2:CW:73:TYR:CG	2.39	0.57
2:DD:113:LYS:NZ	2:DD:160:LEU:O	2.37	0.57
5:DF:2:ARG:O	5:DF:33:TYR:HB2	2.04	0.57
5:DF:59:ARG:HD2	5:DF:60:PRO:HD2	1.87	0.57
1:DJ:25:ARG:HG2	1:DQ:25:ARG:NE	2.19	0.57
2:DK:137:THR:O	2:DK:141:VAL:HG22	2.05	0.57
2:DT:55:ALA:O	2:DT:59:SER:OG	2.08	0.57
2:DT:113:LYS:HE2	2:DT:160:LEU:O	2.04	0.57
2:DV:132:ALA:O	2:DV:136:VAL:HG23	2.04	0.57
9:EA:19:ASP:C	9:EA:22:PRO:HD2	2.30	0.57
1:AA:25:ARG:HE	1:AH:29:PHE:HB2	1.70	0.57
1:AH:137:ALA:HB1	1:AH:141:MET:HE2	1.87	0.57
1:AP:66:VAL:HG12	1:DU:68:PRO:HG2	1.86	0.57
2:AW:110:ASN:HD21	5:BB:56:ALA:HB1	1.69	0.57
1:AX:32:GLY:O	1:AX:36:ARG:HG3	2.04	0.57
1:AX:38:ARG:O	1:AX:42:THR:HG23	2.04	0.57
1:AX:39:ILE:HG23	1:AX:141:MET:SD	2.45	0.57
6:BD:248:ILE:HG12	6:BD:356:SER:HB2	1.86	0.57
9:BH:63:LEU:HD12	9:BH:105:ILE:HD13	1.87	0.57
9:BH:229:ARG:HG2	9:EA:227:PHE:CB	2.35	0.57
2:BJ:10:ASN:O	2:BJ:14:VAL:HG23	2.04	0.57
4:BS:113:LEU:HD23	6:CM:408:CYS:SG	2.45	0.57
2:BT:56:VAL:O	2:BT:61:LEU:HG	2.03	0.57
1:CC:96:VAL:HG12	1:CC:152:TYR:CD2	2.40	0.57
2:CF:122:PRO:O	2:CF:126:THR:HG23	2.04	0.57
2:CH:50:THR:O	2:CH:54:GLU:HG3	2.05	0.57
6:CM:847:THR:HG21	6:CM:857:ARG:CZ	2.35	0.57
8:CP:197:GLN:OE1	8:CP:197:GLN:N	2.36	0.57
2:CS:72:MET:HE2	2:CS:81:CYS:SG	2.45	0.57
1:CY:43:LEU:HD21	1:CY:137:ALA:HB1	1.87	0.57
2:CZ:73:TYR:OH	1:DA:90:ARG:NH2	2.31	0.57
9:DI:24:THR:HG21	9:DI:142:ILE:HD11	1.86	0.57
2:DM:15:GLN:CG	2:DM:17:LYS:HG2	2.35	0.57
1:DQ:89:LEU:O	1:DQ:93:THR:HG23	2.05	0.57
2:DR:123:ILE:HG23	2:DR:160:LEU:CD2	2.35	0.57
1:DS:90:ARG:NH1	2:DT:16:GLY:O	2.38	0.57
1:AA:16:ARG:O	2:AB:94:TYR:OH	2.23	0.57

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
4:AK:84:ARG:HD2	6:BD:252:GLU:O	2.04	0.57
4:AK:123:PRO:HB3	2:AS:64:ASP:CB	2.33	0.57
1:AN:8:ILE:CB	2:AO:1:MET:HE1	2.35	0.57
2:AW:74:THR:HG1	2:AW:77:ARG:HH21	1.51	0.57
1:AZ:88:TYR:O	1:AZ:92:VAL:HG23	2.05	0.57
5:BA:15:ARG:HD2	5:BA:22:LEU:HD12	1.85	0.57
6:BD:199:ILE:HD13	6:BD:227:LEU:HD21	1.87	0.57
6:BD:225:ASP:HA	6:BD:228:LEU:CD2	2.35	0.57
9:BH:42:PHE:HD2	9:BH:131:LEU:HD13	1.69	0.57
9:BH:246:GLN:NE2	9:BH:247:ASN:OD1	2.30	0.57
3:BM:90:ARG:NH1	2:BN:13:ASP:OD1	2.36	0.57
2:BQ:123:ILE:HA	2:BQ:126:THR:HG22	1.87	0.57
2:BT:84:ASP:OD2	2:BT:116:TYR:OH	2.23	0.57
2:BY:75:THR:CG2	1:BZ:108:GLY:HA2	2.35	0.57
1:BZ:115:MET:HE1	10:BZ:200:CYC:CMB	2.35	0.57
1:CE:72:ALA:HB2	10:CE:200:CYC:HMD1	1.87	0.57
2:CF:36:LEU:HA	2:CF:39:ARG:HH12	1.69	0.57
5:CK:6:ILE:HG21	5:CK:36:TRP:CZ3	2.40	0.57
9:CQ:70:GLU:O	9:CQ:74:MET:HG3	2.04	0.57
2:CS:20:ASP:O	2:CS:24:MET:HG2	2.05	0.57
2:CS:22:ALA:HA	2:CS:25:ASP:OD2	2.05	0.57
1:CY:72:ALA:HA	1:CY:77:MET:HB3	1.86	0.57
1:CY:88:TYR:O	1:CY:92:VAL:HG23	2.05	0.57
2:CZ:61:LEU:HG	2:CZ:62:TYR:CD2	2.39	0.57
2:CZ:87:TYR:CD2	10:CZ:200:CYC:HBB3	2.39	0.57
2:DD:132:ALA:O	2:DD:136:VAL:HG23	2.05	0.57
8:DG:221:MET:HE3	1:DQ:52:LYS:HA	1.87	0.57
9:DI:276:PRO:HA	9:DI:280:GLY:O	2.05	0.57
2:DM:85:LEU:HD22	2:DM:133:ILE:HD11	1.87	0.57
1:DQ:101:VAL:HA	1:DQ:104:ILE:CD1	2.35	0.57
1:DQ:123:ILE:HD12	1:DQ:123:ILE:H	1.69	0.57
1:DS:91:LEU:HD21	1:DS:107:ILE:HG22	1.85	0.57
1:DS:130:VAL:HA	1:DS:133:MET:HE3	1.86	0.57
1:AA:65:ILE:CG2	1:AA:72:ALA:HB3	2.35	0.56
2:AD:116:TYR:HB3	2:AD:121:VAL:HB	1.86	0.56
2:AQ:52:VAL:HG21	2:AQ:86:ASP:OD1	2.05	0.56
2:AQ:85:LEU:HG	10:AQ:200:CYC:HBC1	1.87	0.56
1:AR:134:LYS:NZ	1:AR:154:ASP:OD1	2.26	0.56
6:BD:401:LEU:HD13	6:BD:411:TRP:HH2	1.69	0.56
9:BH:37:LEU:HD12	9:BH:40:ILE:HD11	1.87	0.56
3:BM:80:GLN:OE1	3:BM:83:ARG:NH2	2.37	0.56

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
4:BS:109:VAL:O	4:BS:113:LEU:HG	2.05	0.56
1:BV:63:PRO:HG2	1:CE:68:PRO:O	2.05	0.56
2:CD:57:ALA:CB	2:CD:61:LEU:HD12	2.35	0.56
6:CM:290:PHE:O	6:CM:292:ARG:NH1	2.38	0.56
2:DB:112:LEU:HD13	10:DB:200:CYC:CMB	2.35	0.56
5:DF:2:ARG:HG2	5:DF:55:LEU:HD11	1.87	0.56
9:DI:42:PHE:HD2	9:DI:131:LEU:HD13	1.70	0.56
9:DI:43:ALA:O	9:DI:51:LEU:HD12	2.05	0.56
2:DM:96:MET:HA	2:DM:152:TYR:CE2	2.40	0.56
1:DQ:8:ILE:HD11	2:DR:94:TYR:CB	2.31	0.56
9:EA:146:GLU:O	9:EA:150:GLN:HG3	2.05	0.56
1:AA:3:ILE:HG23	1:AH:25:ARG:HD3	1.86	0.56
2:AB:85:LEU:HD21	10:AB:200:CYC:HBC1	1.88	0.56
2:AD:87:TYR:HE2	5:BA:22:LEU:HD13	1.70	0.56
2:AI:85:LEU:HG	10:AI:200:CYC:HBC1	1.86	0.56
1:AP:76:GLU:HG2	1:AP:77:MET:HE2	1.87	0.56
2:AS:138:ALA:HA	2:AS:141:VAL:HG12	1.87	0.56
2:AW:125:SER:HA	2:AW:128:GLN:CD	2.30	0.56
6:BD:886:PRO:HG3	2:CZ:114:GLU:OE1	2.05	0.56
9:BH:217:PHE:HA	9:BH:298:ILE:HG23	1.86	0.56
9:BH:311:GLU:OE1	9:BH:312:LEU:HD22	2.05	0.56
2:BJ:130:ILE:HA	2:BJ:133:ILE:HG22	1.86	0.56
1:BK:100:ASP:OD2	1:BK:102:THR:OG1	2.14	0.56
2:BN:63:SER:H	2:BN:66:THR:HG22	1.70	0.56
2:BN:72:MET:HE2	2:BN:78:TYR:CD2	2.40	0.56
2:BT:85:LEU:HD22	2:BT:133:ILE:CD1	2.33	0.56
1:BV:8:ILE:HD12	2:BW:94:TYR:CD1	2.39	0.56
2:CA:107:ARG:O	6:CM:597:TYR:HA	2.05	0.56
6:CM:727:VAL:CG1	5:DX:20:ARG:HD3	2.35	0.56
9:CQ:210:PHE:CD2	9:CQ:241:PHE:HB3	2.40	0.56
2:CS:3:ASP:N	2:CS:6:THR:OG1	2.37	0.56
2:CS:127:VAL:O	2:CS:131:GLN:HG2	2.04	0.56
2:CW:93:THR:O	2:CW:97:LEU:HD23	2.05	0.56
1:CY:110:VAL:CG1	2:DD:76:ARG:HB2	2.35	0.56
1:DC:85:MET:O	1:DC:133:MET:HE1	2.06	0.56
5:DF:5:ARG:HB3	5:DF:30:LEU:HD23	1.87	0.56
5:DF:6:ILE:O	5:DF:28:THR:HA	2.05	0.56
8:DH:240:ILE:HD11	8:DH:245:SER:HB2	1.87	0.56
2:DK:39:ARG:O	2:DK:43:VAL:HG23	2.04	0.56
1:DQ:39:ILE:HD11	1:DQ:145:ASP:CB	2.35	0.56
1:DQ:116:TYR:CE2	1:DQ:126:VAL:HG11	2.41	0.56

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:DR:43:VAL:O	2:DR:46:ALA:HB3	2.05	0.56
1:DS:62:ARG:HG2	9:EA:57:GLY:HA3	1.86	0.56
2:DV:54:GLU:O	2:DV:58:LYS:HG2	2.05	0.56
9:EA:210:PHE:CG	9:EA:241:PHE:HB3	2.39	0.56
1:AA:65:ILE:HG22	1:AA:72:ALA:HB3	1.86	0.56
1:AA:71:ASN:HD22	1:AA:121:THR:HA	1.69	0.56
3:AE:9:LEU:HG	2:AF:1:MET:SD	2.45	0.56
2:AF:15:GLN:HB3	2:AF:17:LYS:HZ3	1.69	0.56
1:AH:81:CYS:HA	10:AH:200:CYC:HHD	1.88	0.56
1:AH:113:ARG:HG2	1:AH:117:ARG:HH21	1.68	0.56
2:AI:76:ARG:CZ	1:AJ:110:VAL:HG11	2.36	0.56
2:AI:141:VAL:HG12	2:AI:145:ALA:HB3	1.87	0.56
1:AJ:128:GLN:O	1:AJ:132:GLU:HG3	2.05	0.56
2:AQ:65:VAL:HG12	2:AQ:72:MET:HB2	1.87	0.56
2:AQ:78:TYR:O	2:AQ:82:ILE:HG12	2.05	0.56
1:AR:19:SER:OG	1:AR:22:GLU:HG3	2.05	0.56
1:AR:36:ARG:NH1	1:AR:99:GLY:HA2	2.21	0.56
1:AR:87:TYR:O	1:AR:91:LEU:HG	2.05	0.56
2:AS:91:TYR:OH	2:AS:107:ARG:NE	2.33	0.56
2:AU:68:PRO:HG3	2:AU:73:TYR:CE1	2.41	0.56
2:AU:147:LYS:O	2:AU:151:VAL:HG23	2.04	0.56
1:AV:12:ASP:OD2	2:AW:107:ARG:HD3	2.05	0.56
1:AX:112:VAL:HG21	1:AX:160:MET:HE1	1.88	0.56
2:AY:96:MET:HG3	2:AY:148:GLU:CD	2.31	0.56
1:AZ:53:GLN:HE22	1:AZ:57:ARG:HH21	1.53	0.56
5:BA:30:LEU:HD23	6:BD:313:ASP:O	2.05	0.56
6:BD:555:LEU:HD22	6:BD:587:LEU:HD21	1.87	0.56
4:BS:135:LYS:HB2	4:BS:157:PHE:CG	2.41	0.56
1:BZ:152:TYR:O	1:BZ:156:VAL:HG23	2.05	0.56
2:CH:1:MET:HE3	2:CH:103:ILE:HB	1.86	0.56
6:CM:574:SER:OG	6:CM:577:GLU:OE1	2.22	0.56
6:CM:586:GLU:OE1	6:CM:586:GLU:N	2.38	0.56
8:CP:214:ASN:O	8:CP:217:ASN:ND2	2.27	0.56
8:CP:232:ILE:HD12	8:CP:233:PRO:CD	2.35	0.56
1:CR:76:GLU:HA	8:DZ:245:SER:O	2.06	0.56
1:CR:127:ALA:O	1:CR:131:ARG:HG3	2.06	0.56
2:CS:104:LEU:HD23	2:CS:155:TYR:CD2	2.35	0.56
1:CY:82:LEU:HD22	8:DY:218:PHE:HB3	1.87	0.56
1:CY:93:THR:O	1:CY:97:VAL:HG23	2.05	0.56
1:CY:111:GLY:O	1:CY:115:MET:HG2	2.05	0.56
1:CY:126:VAL:O	1:CY:130:VAL:HG23	2.05	0.56

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:CZ:28:LYS:HE2	2:CZ:28:LYS:HA	1.87	0.56
2:DB:15:GLN:HG3	2:DB:17:LYS:NZ	2.21	0.56
1:DC:37:LEU:HD13	2:DD:28:LYS:HG2	1.86	0.56
2:DD:93:THR:HA	2:DD:149:MET:HE1	1.87	0.56
9:DI:39:LEU:CD1	9:DI:139:LEU:HB2	2.36	0.56
9:DI:245:CYS:O	9:DI:246:GLN:NE2	2.38	0.56
2:DR:15:GLN:HB3	2:DR:17:LYS:HE3	1.86	0.56
2:DR:44:ILE:HD13	2:DR:149:MET:HE3	1.88	0.56
2:DR:62:TYR:OH	10:DS:200:CYC:HAD2	2.04	0.56
1:DS:38:ARG:HD2	1:DS:41:GLU:OE1	2.05	0.56
1:DS:77:MET:O	1:DS:80:THR:OG1	2.19	0.56
2:DT:1:MET:SD	2:DT:103:ILE:HD12	2.46	0.56
2:DT:74:THR:O	2:DT:77:ARG:N	2.35	0.56
8:DY:222:ALA:HA	8:DY:225:ILE:HD12	1.88	0.56
9:EA:190:ILE:HD11	9:EA:202:MET:SD	2.45	0.56
9:EA:202:MET:CE	9:EA:269:VAL:HG21	2.33	0.56
9:EA:245:CYS:O	9:EA:246:GLN:NE2	2.37	0.56
2:AD:68:PRO:HG3	2:AD:73:TYR:CE1	2.40	0.56
2:AF:72:MET:HE2	2:AF:78:TYR:CE2	2.40	0.56
2:AS:67:ARG:O	2:AS:73:TYR:HB2	2.06	0.56
6:BD:309:VAL:HG21	6:BD:319:PHE:CG	2.40	0.56
6:BD:489:PRO:HD2	6:BD:492:ASN:HD22	1.70	0.56
6:BD:871:ARG:HE	6:BD:884:VAL:HG23	1.71	0.56
9:BH:89:ARG:HD3	9:BH:170:LYS:CD	2.35	0.56
9:BH:98:TYR:HA	9:BH:101:TRP:CE3	2.40	0.56
2:BJ:106:GLU:HG3	2:BJ:107:ARG:HG3	1.87	0.56
3:BM:29:PHE:HA	3:BM:32:THR:HG22	1.85	0.56
3:BM:101:LYS:HD2	3:BM:155:TYR:CE2	2.40	0.56
2:BQ:60:LEU:HD13	2:BQ:72:MET:SD	2.46	0.56
1:BX:76:GLU:OE1	1:BX:76:GLU:N	2.37	0.56
2:CA:78:TYR:O	2:CA:82:ILE:HG12	2.04	0.56
2:CF:76:ARG:HB2	1:CG:110:VAL:CG1	2.35	0.56
6:CM:239:LYS:NZ	6:CM:403:VAL:O	2.37	0.56
9:CQ:21:VAL:HG13	9:CQ:153:VAL:CG2	2.36	0.56
9:CQ:285:ASN:HB2	9:CQ:306:LEU:HB2	1.87	0.56
1:CR:67:SER:HB3	8:DZ:230:ASN:ND2	2.20	0.56
2:CU:113:LYS:NZ	2:CU:160:LEU:O	2.25	0.56
1:DC:50:ILE:CD1	1:DC:140:LEU:HD21	2.35	0.56
9:DI:195:ASN:HB3	9:DI:198:VAL:CG1	2.34	0.56
1:DQ:145:ASP:O	1:DQ:148:GLU:HG3	2.04	0.56
1:DS:128:GLN:O	1:DS:132:GLU:HG2	2.06	0.56

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:AB:1:MET:CE	2:AB:103:ILE:HD13	2.36	0.56
2:AF:76:ARG:NH2	5:BA:62:THR:O	2.38	0.56
1:AJ:86:ASP:OD1	4:AK:18:TYR:OH	2.20	0.56
1:AN:92:VAL:HG11	1:AN:153:PHE:CZ	2.41	0.56
1:AR:52:LYS:HE2	8:BG:225:ILE:HB	1.86	0.56
2:AU:28:LYS:HD3	1:AZ:37:LEU:HD13	1.87	0.56
6:BD:753:PRO:HB3	2:DD:79:ALA:HB1	1.87	0.56
9:BH:216:LEU:HA	9:BH:297:LYS:HD3	1.87	0.56
1:BK:8:ILE:CD1	2:BL:98:ALA:HB2	2.36	0.56
3:BM:131:THR:HG22	3:BM:157:ILE:HD13	1.87	0.56
1:BP:115:MET:HE1	10:BP:200:CYC:CMB	2.35	0.56
2:BQ:112:LEU:HG	2:BQ:160:LEU:HD21	1.87	0.56
2:BT:15:GLN:OE1	7:CO:99:ALA:N	2.35	0.56
1:CC:23:LEU:HD22	2:CD:38:VAL:HG13	1.88	0.56
1:CC:87:TYR:O	1:CC:91:LEU:HG	2.06	0.56
1:CC:116:TYR:HE2	1:CC:126:VAL:HG11	1.71	0.56
1:CE:91:LEU:HD21	1:CE:107:ILE:HG21	1.87	0.56
6:CM:5:ALA:HB3	6:CM:437:TYR:CD1	2.41	0.56
6:CM:745:GLN:NE2	6:CM:807:HIS:O	2.30	0.56
1:CR:3:ILE:HG22	1:CR:100:ASP:HB3	1.87	0.56
1:DC:97:VAL:HG21	2:DD:19:LEU:HD12	1.88	0.56
2:DD:57:ALA:HB1	8:DH:201:ARG:HH22	1.71	0.56
1:DJ:46:SER:O	1:DJ:49:THR:OG1	2.19	0.56
2:DM:106:GLU:HG2	2:DM:107:ARG:HG3	1.86	0.56
9:EA:87:ALA:HA	9:EA:161:MET:CE	2.35	0.56
1:AA:97:VAL:HG21	2:AB:19:LEU:CD1	2.36	0.56
10:AB:200:CYC:O2A	6:BD:310:ARG:HD2	2.06	0.56
2:AD:10:ASN:O	2:AD:14:VAL:HG13	2.06	0.56
2:AD:144:ASP:OD1	2:AD:145:ALA:N	2.39	0.56
2:AI:76:ARG:HD3	6:BD:362:GLN:OE1	2.05	0.56
1:AJ:58:LEU:HD22	1:AJ:129:SER:CB	2.35	0.56
2:AO:109:LEU:HD13	2:AO:159:GLY:HA3	1.86	0.56
1:AP:58:LEU:HD22	1:AP:129:SER:CB	2.36	0.56
2:AW:8:VAL:CG1	2:AW:23:ALA:HB1	2.34	0.56
1:AZ:6:LYS:NZ	1:AZ:100:ASP:OD2	2.26	0.56
1:AZ:65:ILE:HG22	1:AZ:72:ALA:CB	2.35	0.56
9:BH:45:LEU:O	9:BH:49:LYS:NZ	2.37	0.56
9:BH:46:GLU:HB3	9:BH:50:THR:CG2	2.35	0.56
10:BJ:200:CYC:HBD1	10:BJ:200:CYC:HHA	1.86	0.56
1:BK:37:LEU:HD23	2:BL:28:LYS:HD3	1.88	0.56
1:BP:91:LEU:HD21	1:BP:107:ILE:HG21	1.88	0.56

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:BP:141:MET:HE1	1:BP:149:ALA:CB	2.35	0.56
2:BT:1:MET:HE1	6:CM:25:ILE:CB	2.33	0.56
1:BZ:50:ILE:HA	1:BZ:136:VAL:HG11	1.87	0.56
1:CE:97:VAL:HG11	2:CF:19:LEU:HD12	1.87	0.56
6:CM:560:ARG:NH2	6:CM:565:GLU:OE1	2.36	0.56
9:CQ:227:PHE:CD1	9:DI:229:ARG:HA	2.40	0.56
1:CT:8:ILE:HG12	1:CT:18:LEU:HD11	1.88	0.56
1:CV:81:CYS:O	1:CV:85:MET:HG2	2.05	0.56
1:DA:61:LYS:HB3	9:DI:57:GLY:CA	2.36	0.56
1:DJ:92:VAL:HG11	1:DJ:153:PHE:CD2	2.40	0.56
2:DR:65:VAL:HG12	2:DR:72:MET:CB	2.34	0.56
2:DR:104:LEU:HD12	2:DR:155:TYR:HD2	1.71	0.56
2:AB:137:THR:O	2:AB:141:VAL:HG12	2.06	0.56
1:AV:39:ILE:HG23	1:AV:141:MET:HE3	1.88	0.56
1:AV:58:LEU:HD22	1:AV:129:SER:CB	2.33	0.56
2:AW:72:MET:HE2	2:AW:78:TYR:CD2	2.41	0.56
6:BD:612:ARG:NE	6:BD:644:GLU:OE2	2.39	0.56
8:BG:196:TRP:O	8:BG:200:VAL:HG23	2.06	0.56
1:BK:4:VAL:HG23	2:BL:3:ASP:OD2	2.06	0.56
1:BK:141:MET:HE2	1:BK:146:ALA:HA	1.88	0.56
2:BL:68:PRO:HG3	2:BL:73:TYR:CE1	2.41	0.56
1:BR:58:LEU:HD22	1:BR:129:SER:HB3	1.88	0.56
1:BV:47:ARG:O	1:BV:51:VAL:HG22	2.05	0.56
10:CD:200:CYC:O2A	5:CK:37:PHE:HB2	2.06	0.56
6:CM:357:SER:O	6:CM:361:VAL:HG23	2.06	0.56
2:CU:137:THR:O	2:CU:141:VAL:HG22	2.05	0.56
1:CV:126:VAL:HG12	1:CV:160:MET:CE	2.36	0.56
1:CY:39:ILE:O	1:CY:42:THR:HG22	2.04	0.56
1:CY:65:ILE:HD12	10:CY:200:CYC:OC	2.05	0.56
1:CY:115:MET:HE3	2:DD:79:ALA:HB2	1.88	0.56
1:DC:55:GLY:HA2	1:DC:85:MET:HE1	1.88	0.56
9:DI:81:GLN:HA	9:DI:176:VAL:HG21	1.86	0.56
9:DI:141:THR:O	9:DI:145:LEU:HG	2.05	0.56
9:DI:232:VAL:O	9:DI:236:ASN:ND2	2.30	0.56
9:DI:258:GLU:OE1	9:DI:266:GLN:HG3	2.06	0.56
2:DM:101:ALA:HB2	2:DM:152:TYR:CD1	2.40	0.56
2:DM:131:GLN:O	2:DM:135:GLU:HG2	2.05	0.56
1:DQ:66:VAL:HG12	1:DQ:72:ALA:O	2.06	0.56
1:DU:27:LYS:O	1:DU:31:THR:HG23	2.06	0.56
9:EA:272:LYS:HA	9:EA:284:MET:O	2.06	0.56
2:AF:77:ARG:HG2	10:AF:200:CYC:HAD1	1.87	0.56

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:AP:113:ARG:NH1	1:AP:160:MET:O	2.38	0.56
1:AP:145:ASP:HB2	2:DT:39:ARG:HH22	1.69	0.56
2:AQ:105:ASP:OD1	6:BD:671:ARG:HD3	2.06	0.56
2:AS:41:ALA:N	2:AS:96:MET:HE1	2.21	0.56
1:AV:50:ILE:HD11	1:AV:140:LEU:HD22	1.87	0.56
6:BD:453:PRO:HG3	6:BD:462:PHE:CZ	2.40	0.56
2:BL:83:ARG:HD2	5:CJ:22:LEU:HG	1.87	0.56
3:BM:43:LEU:HD12	3:BM:93:THR:HG22	1.87	0.56
2:BN:104:LEU:O	2:BN:108:VAL:HB	2.06	0.56
10:BQ:200:CYC:HBB2	6:CM:338:ILE:CD1	2.35	0.56
2:BT:3:ASP:H	2:BT:6:THR:HG22	1.71	0.56
1:BV:50:ILE:HG12	1:BV:136:VAL:HG12	1.87	0.56
2:BW:101:ALA:HB1	2:BW:104:LEU:HD12	1.88	0.56
1:BX:39:ILE:HG12	1:BX:145:ASP:HB3	1.87	0.56
2:CF:39:ARG:O	2:CF:43:VAL:HG23	2.06	0.56
1:CG:153:PHE:O	1:CG:157:ILE:HG12	2.05	0.56
5:CJ:2:ARG:NH1	5:CJ:66:LEU:HD22	2.21	0.56
6:CM:75:ILE:HD12	6:CM:197:VAL:CG2	2.33	0.56
6:CM:747:PHE:HA	6:CM:791:PHE:CE2	2.41	0.56
6:CM:796:PRO:HG3	2:DR:110:ASN:HB3	1.86	0.56
9:CQ:116:LEU:HB3	9:CQ:122:VAL:HG23	1.88	0.56
2:CS:51:ILE:HG21	2:CS:137:THR:CG2	2.35	0.56
1:CV:104:ILE:HG22	1:CV:109:LEU:CD2	2.36	0.56
1:CV:128:GLN:O	1:CV:132:GLU:HG2	2.06	0.56
2:CW:83:ARG:NH1	10:CW:200:CYC:O1A	2.30	0.56
1:DA:33:GLY:O	1:DA:37:LEU:HD23	2.05	0.56
2:DD:61:LEU:HD21	2:DD:78:TYR:OH	2.06	0.56
1:DL:131:ARG:HG3	1:DL:157:ILE:HD12	1.88	0.56
1:DQ:50:ILE:HD11	1:DQ:137:ALA:N	2.21	0.56
1:DQ:153:PHE:HA	1:DQ:156:VAL:HG12	1.86	0.56
2:DR:54:GLU:O	2:DR:58:LYS:HG2	2.06	0.56
2:DR:96:MET:HA	2:DR:152:TYR:CE2	2.40	0.56
2:DV:122:PRO:O	2:DV:126:THR:HG23	2.06	0.56
9:EA:131:LEU:HG	9:EA:135:ALA:HB3	1.88	0.56
2:AB:61:LEU:HD22	10:AC:200:CYC:CAA	2.36	0.56
1:AC:33:GLY:HA2	1:AC:36:ARG:HG2	1.87	0.56
2:AI:25:ASP:OD2	9:BH:165:ALA:HB1	2.06	0.56
4:AK:72:ASN:HB3	10:AK:200:CYC:OC	2.06	0.56
2:AS:132:ALA:O	2:AS:136:VAL:HG23	2.06	0.56
10:AV:200:CYC:HMD1	10:AV:200:CYC:NC	2.20	0.56
2:AW:55:ALA:O	2:AW:59:SER:OG	2.14	0.56

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:AW:122:PRO:O	2:AW:126:THR:HG23	2.06	0.56
6:BD:264:ARG:O	6:BD:266:LYS:NZ	2.26	0.56
6:BD:701:ARG:HB3	6:BD:706:ILE:CD1	2.32	0.56
1:BK:27:LYS:O	1:BK:31:THR:HG23	2.05	0.56
3:BM:38:ARG:HA	3:BM:41:GLU:CD	2.31	0.56
2:BN:115:THR:CG2	5:CJ:53:VAL:HG21	2.36	0.56
2:BT:87:TYR:CD2	10:BT:200:CYC:HBB3	2.40	0.56
2:BW:3:ASP:OD1	2:BW:6:THR:OG1	2.13	0.56
5:CJ:2:ARG:HH12	5:CJ:66:LEU:HD22	1.71	0.56
6:CM:35:LEU:HD22	6:CM:39:GLU:HB3	1.88	0.56
6:CM:805:THR:HG22	6:CM:811:ARG:O	2.06	0.56
1:CR:68:PRO:HA	1:CR:73:TYR:CG	2.40	0.56
2:CU:73:TYR:O	2:CU:77:ARG:HB2	2.05	0.56
2:CU:112:LEU:HD13	10:CU:200:CYC:HMB2	1.87	0.56
1:CY:85:MET:CE	1:CY:133:MET:HE2	2.36	0.56
2:DD:50:THR:O	2:DD:54:GLU:HG2	2.06	0.56
9:DI:46:GLU:HB3	9:DI:50:THR:HB	1.87	0.56
9:DI:190:ILE:CG2	9:DI:193:VAL:HB	2.35	0.56
9:DI:288:TRP:HD1	9:DI:303:ILE:HG12	1.71	0.56
1:DJ:115:MET:HE1	10:DJ:200:CYC:CMB	2.36	0.56
2:DK:43:VAL:HG12	2:DK:140:LEU:HG	1.88	0.56
2:DO:126:THR:O	2:DO:130:ILE:HG12	2.05	0.56
2:DV:23:ALA:O	2:DV:27:LEU:HD13	2.06	0.56
2:DV:51:ILE:HG12	2:DV:140:LEU:CD1	2.36	0.56
1:AH:65:ILE:HG22	1:AH:72:ALA:HB3	1.87	0.56
1:AH:76:GLU:CG	1:AH:77:MET:HE2	2.35	0.56
2:AL:122:PRO:HB2	2:AL:125:SER:HB2	1.87	0.56
2:AS:131:GLN:O	2:AS:134:LYS:HB2	2.06	0.56
2:AW:39:ARG:O	2:AW:43:VAL:HG23	2.05	0.56
5:BB:6:ILE:HG21	5:BB:36:TRP:CZ3	2.41	0.56
1:BK:58:LEU:HD22	1:BK:129:SER:CB	2.34	0.56
3:BM:104:ILE:HG22	3:BM:109:LEU:HG	1.88	0.56
2:BQ:77:ARG:CG	10:BQ:200:CYC:HMD1	2.36	0.56
4:BS:73:ALA:HA	4:BS:78:ARG:HB3	1.86	0.56
2:BW:56:VAL:HG12	2:BW:61:LEU:HG	1.88	0.56
2:CA:39:ARG:HA	2:CA:39:ARG:NH1	2.21	0.56
2:CA:119:LEU:HD12	10:CM:901:CYC:HBD1	1.88	0.56
6:CM:15:GLN:NE2	6:CM:262:ALA:HB2	2.21	0.56
6:CM:751:LEU:HD12	6:CM:755:ILE:HD13	1.86	0.56
1:CR:37:LEU:CD1	2:CS:28:LYS:HE3	2.29	0.56
2:CS:43:VAL:CG1	2:CS:141:VAL:HG12	2.36	0.56

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:DC:4:VAL:HG22	1:DC:26:ILE:HD12	1.88	0.56
2:DO:137:THR:O	2:DO:141:VAL:HG22	2.05	0.56
2:DR:51:ILE:HG22	2:DR:133:ILE:HG23	1.86	0.56
5:DX:5:ARG:HB3	5:DX:30:LEU:HD23	1.87	0.56
9:EA:214:ILE:HA	9:EA:217:PHE:CE1	2.41	0.56
2:AD:5:ILE:HG23	2:AD:30:TYR:HD2	1.71	0.55
2:AD:57:ALA:HA	2:AD:61:LEU:HD12	1.89	0.55
3:AE:48:LYS:HA	3:AE:51:VAL:HG12	1.87	0.55
3:AE:50:ILE:HG12	3:AE:140:LEU:HD13	1.88	0.55
2:AO:57:ALA:HA	2:AO:61:LEU:HD12	1.87	0.55
1:AP:105:GLU:O	1:AP:110:VAL:HG23	2.06	0.55
2:AQ:115:THR:HG23	6:BD:461:ILE:HG13	1.88	0.55
2:AS:68:PRO:HA	2:AS:73:TYR:CG	2.42	0.55
5:BB:30:LEU:HD12	6:BD:572:ASP:HB3	1.88	0.55
6:BD:210:PHE:CE2	6:BD:212:ASN:HB2	2.40	0.55
9:BH:227:PHE:CD1	9:EA:229:ARG:HA	2.41	0.55
3:BM:26:ILE:O	3:BM:30:LEU:HD23	2.06	0.55
4:BS:33:SER:O	4:BS:37:ARG:HG3	2.05	0.55
1:CC:97:VAL:HG21	2:CD:19:LEU:CD1	2.36	0.55
1:CE:104:ILE:HG22	1:CE:109:LEU:HD12	1.88	0.55
2:CF:62:TYR:N	10:CG:200:CYC:O1A	2.32	0.55
2:CF:72:MET:HE2	2:CF:78:TYR:CD2	2.41	0.55
2:CF:131:GLN:O	2:CF:135:GLU:HG2	2.06	0.55
1:CR:106:GLU:OE1	5:DF:64:ALA:HA	2.06	0.55
9:DI:41:TRP:NE1	9:DI:45:LEU:HD11	2.22	0.55
1:DJ:90:ARG:HD2	2:DK:17:LYS:C	2.31	0.55
1:DQ:87:TYR:O	1:DQ:91:LEU:HG	2.06	0.55
2:DR:148:GLU:O	2:DR:151:VAL:HG12	2.06	0.55
2:DT:15:GLN:CG	2:DT:17:LYS:HG3	2.35	0.55
8:DY:211:THR:N	8:DY:214:ASN:OD1	2.39	0.55
9:EA:248:LEU:HD11	9:EA:277:TRP:CZ3	2.41	0.55
1:AA:118:SER:O	2:AF:53:LYS:HD3	2.06	0.55
1:AC:50:ILE:CD1	1:AC:136:VAL:HG12	2.36	0.55
2:AF:15:GLN:HB3	2:AF:17:LYS:NZ	2.21	0.55
1:AJ:2:SER:N	1:AJ:100:ASP:OD2	2.39	0.55
1:AR:92:VAL:HG11	1:AR:153:PHE:CZ	2.42	0.55
2:AS:107:ARG:O	6:BD:597:TYR:HA	2.06	0.55
2:AW:101:ALA:HB1	2:AW:104:LEU:HD12	1.88	0.55
5:BA:2:ARG:HE	5:BA:66:LEU:HD12	1.71	0.55
5:BB:29:LYS:HG3	5:BB:31:VAL:HG13	1.88	0.55
6:BD:329:TYR:OH	6:BD:349:HIS:ND1	2.26	0.55

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
9:BH:39:LEU:CD1	9:BH:139:LEU:HB2	2.36	0.55
10:BJ:200:CYC:O2A	6:CM:310:ARG:HD2	2.06	0.55
2:BL:87:TYR:CE2	5:CJ:22:LEU:HB2	2.41	0.55
2:BL:142:GLY:O	2:BL:146:GLY:N	2.39	0.55
4:BS:41:ALA:HB1	4:BS:98:ILE:HD11	1.88	0.55
2:CF:122:PRO:HB2	2:CF:125:SER:HB2	1.86	0.55
9:CQ:190:ILE:HD13	9:CQ:252:PRO:HB2	1.88	0.55
2:CU:78:TYR:O	2:CU:82:ILE:HG12	2.06	0.55
1:CV:115:MET:CE	10:CV:200:CYC:HMA3	2.35	0.55
9:DI:68:LEU:HD22	9:DI:112:ARG:HG2	1.86	0.55
9:DI:267:ILE:HG22	9:DI:269:VAL:HG13	1.89	0.55
9:DI:272:LYS:HA	9:DI:284:MET:O	2.05	0.55
2:DK:33:SER:O	2:DK:37:ARG:HG2	2.06	0.55
1:DL:102:THR:OG1	1:DL:103:PRO:HD3	2.06	0.55
1:DN:141:MET:HG2	1:DN:145:ASP:HB2	1.88	0.55
2:DR:137:THR:HG21	2:DR:149:MET:HG2	1.88	0.55
2:DT:103:ILE:HG13	2:DT:107:ARG:HD2	1.89	0.55
10:DT:200:CYC:HBD1	10:DT:200:CYC:HHA	1.87	0.55
2:DV:56:VAL:HA	2:DV:59:SER:OG	2.06	0.55
2:DV:71:ASN:ND2	2:DV:121:VAL:HA	2.21	0.55
9:EA:141:THR:O	9:EA:145:LEU:HG	2.05	0.55
2:AI:83:ARG:HD2	6:BD:369:SER:CB	2.37	0.55
4:AK:88:TYR:CD2	10:AK:200:CYC:HBB3	2.41	0.55
1:AP:134:LYS:HD3	1:AP:153:PHE:HB3	1.87	0.55
1:AV:37:LEU:HD11	2:AW:24:MET:SD	2.47	0.55
1:AV:90:ARG:HG2	1:AV:94:TYR:CZ	2.41	0.55
1:AZ:93:THR:O	1:AZ:97:VAL:HG23	2.06	0.55
9:BH:276:PRO:HA	9:BH:281:ASN:OD1	2.05	0.55
2:BN:71:ASN:HD22	2:BN:121:VAL:HA	1.70	0.55
1:BP:89:LEU:O	1:BP:93:THR:HG23	2.06	0.55
2:BQ:104:LEU:HD11	2:BQ:152:TYR:HB3	1.88	0.55
4:BS:2:ARG:HB2	4:BS:103:ASN:HD22	1.71	0.55
1:BV:77:MET:SD	10:BV:200:CYC:O2D	2.64	0.55
1:CE:47:ARG:O	1:CE:51:VAL:HG22	2.06	0.55
1:CE:113:ARG:CD	1:CE:123:ILE:HD13	2.35	0.55
2:CF:96:MET:HG2	2:CF:148:GLU:OE1	2.06	0.55
2:CF:125:SER:HA	2:CF:128:GLN:CD	2.31	0.55
1:CG:107:ILE:HD11	6:CM:532:VAL:CG2	2.37	0.55
2:CH:122:PRO:HB2	2:CH:125:SER:OG	2.06	0.55
2:CS:68:PRO:HA	2:CS:73:TYR:CG	2.40	0.55
1:DA:77:MET:HG3	8:DG:201:ARG:HB2	1.88	0.55

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:DA:85:MET:HG2	1:DA:133:MET:CE	2.36	0.55
2:DD:106:GLU:HG3	2:DD:107:ARG:HG2	1.87	0.55
1:DJ:38:ARG:O	1:DJ:42:THR:HG23	2.07	0.55
2:DK:51:ILE:HG21	2:DK:137:THR:CG2	2.32	0.55
1:AA:114:GLU:OE1	1:AA:114:GLU:N	2.40	0.55
1:AC:65:ILE:HD11	1:AC:122:PRO:HG3	1.88	0.55
1:AC:105:GLU:HA	1:AC:109:LEU:HB3	1.88	0.55
2:AI:123:ILE:HA	2:AI:126:THR:HG22	1.88	0.55
1:AN:98:SER:HB3	2:AO:9:ILE:CD1	2.37	0.55
1:AN:109:LEU:HD11	1:AN:156:VAL:HG22	1.88	0.55
1:AP:102:THR:O	1:AP:106:GLU:HG2	2.06	0.55
2:AY:119:LEU:HD11	10:AY:200:CYC:HAA2	1.88	0.55
6:BD:63:ASN:O	6:BD:67:ILE:HG12	2.06	0.55
6:BD:213:ASN:O	6:BD:217:LYS:N	2.39	0.55
4:BS:118:ASN:HB3	2:CA:122:PRO:HA	1.87	0.55
2:BT:110:ASN:OD1	2:BT:111:GLY:N	2.39	0.55
10:BX:200:CYC:HMD1	10:BX:200:CYC:NC	2.21	0.55
1:BZ:44:THR:O	1:BZ:47:ARG:HD3	2.07	0.55
2:CF:36:LEU:HA	2:CF:39:ARG:NH1	2.22	0.55
1:CR:71:ASN:OD1	10:CR:200:CYC:HBD2	2.06	0.55
2:CU:87:TYR:OH	5:DF:19:GLN:O	2.23	0.55
1:CV:122:PRO:HD2	10:CV:200:CYC:OC	2.06	0.55
2:DB:104:LEU:HD13	2:DB:156:ILE:HG13	1.87	0.55
1:DL:91:LEU:HD11	1:DL:107:ILE:HG21	1.88	0.55
2:DM:124:SER:O	2:DM:128:GLN:HG3	2.06	0.55
1:DQ:72:ALA:HA	1:DQ:77:MET:CB	2.34	0.55
1:DU:122:PRO:O	1:DU:126:VAL:HG23	2.06	0.55
2:DV:22:ALA:HA	2:DV:25:ASP:OD2	2.06	0.55
1:AA:126:VAL:O	1:AA:130:VAL:HG23	2.06	0.55
2:AB:1:MET:HE2	2:AB:103:ILE:HD13	1.88	0.55
2:AB:3:ASP:N	2:AB:6:THR:OG1	2.39	0.55
1:AH:44:THR:O	1:AH:47:ARG:HG2	2.06	0.55
4:AK:33:SER:O	4:AK:37:ARG:HG3	2.06	0.55
1:AN:47:ARG:O	1:AN:51:VAL:HG22	2.05	0.55
2:AS:78:TYR:O	2:AS:82:ILE:HG12	2.07	0.55
1:AV:27:LYS:O	1:AV:30:VAL:HG12	2.06	0.55
6:BD:350:ILE:HD12	6:BD:380:LEU:CD1	2.36	0.55
9:BH:214:ILE:HG21	9:BH:238:LEU:CD2	2.36	0.55
9:BH:300:PHE:CZ	9:BH:302:ALA:HB2	2.41	0.55
2:BJ:61:LEU:HD22	10:BK:200:CYC:HAA1	1.88	0.55
3:BM:127:VAL:O	3:BM:131:THR:HG23	2.06	0.55

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:BV:91:LEU:HD21	1:BV:107:ILE:HB	1.88	0.55
1:BZ:134:LYS:NZ	1:BZ:154:ASP:OD1	2.33	0.55
2:CD:123:ILE:HA	2:CD:126:THR:HG22	1.89	0.55
2:CF:71:ASN:ND2	2:CF:122:PRO:HD3	2.21	0.55
6:CM:452:ASP:OD1	6:CM:607:ARG:NH2	2.40	0.55
9:CQ:107:LEU:HD23	11:CQ:400:45D:H141	1.87	0.55
1:CR:38:ARG:O	1:CR:42:THR:HG23	2.07	0.55
1:CT:62:ARG:O	1:CT:65:ILE:HG12	2.05	0.55
1:DC:71:ASN:HB3	10:DC:200:CYC:OC	2.06	0.55
1:DL:67:SER:O	1:DL:73:TYR:HB2	2.06	0.55
2:DO:104:LEU:HD13	2:DO:156:ILE:CG1	2.37	0.55
1:DU:62:ARG:O	1:DU:65:ILE:HG12	2.06	0.55
2:DV:73:TYR:O	2:DV:77:ARG:HB2	2.07	0.55
2:DV:123:ILE:O	2:DV:127:VAL:HG23	2.06	0.55
9:EA:33:ALA:HB3	9:EA:174:GLU:HB2	1.89	0.55
9:EA:42:PHE:HD2	9:EA:131:LEU:HD13	1.72	0.55
9:EA:146:GLU:HB3	9:EA:149:GLN:OE1	2.07	0.55
3:AE:127:VAL:O	3:AE:131:THR:HG23	2.07	0.55
2:AI:109:LEU:HD13	2:AI:159:GLY:HA3	1.89	0.55
2:AQ:100:ASP:OD1	2:AQ:102:SER:OG	2.12	0.55
10:AQ:200:CYC:HHA	10:AQ:200:CYC:HBA2	1.88	0.55
2:AW:24:MET:HG3	2:AW:28:LYS:NZ	2.21	0.55
2:AW:105:ASP:OD1	2:AW:109:LEU:HD12	2.07	0.55
1:AX:153:PHE:O	1:AX:157:ILE:HG12	2.07	0.55
6:BD:481:LYS:NZ	6:BD:495:VAL:HG22	2.21	0.55
6:BD:626:ILE:HG22	6:BD:635:VAL:HG22	1.88	0.55
9:BH:250:LEU:HD22	9:BH:286:ILE:HD12	1.88	0.55
4:BS:113:LEU:HA	6:CM:408:CYS:SG	2.47	0.55
2:CA:138:ALA:HA	2:CA:141:VAL:HG12	1.88	0.55
6:CM:261:ALA:HB1	6:CM:402:GLY:HA3	1.87	0.55
9:CQ:295:GLU:OE1	9:CQ:297:LYS:HG2	2.06	0.55
10:CU:200:CYC:OB	5:DF:20:ARG:HA	2.07	0.55
1:CY:30:VAL:HA	2:CZ:31:PHE:CD1	2.42	0.55
1:CY:67:SER:O	1:CY:73:TYR:HB2	2.05	0.55
1:CY:110:VAL:HG13	2:DD:76:ARG:HB2	1.89	0.55
2:CZ:62:TYR:OH	10:DA:200:CYC:HAD2	2.05	0.55
2:DK:43:VAL:HG11	2:DK:141:VAL:HA	1.86	0.55
2:DK:72:MET:HE3	2:DK:81:CYS:CB	2.36	0.55
1:DL:50:ILE:HD11	1:DL:137:ALA:HB2	1.88	0.55
2:DR:51:ILE:CG2	2:DR:133:ILE:HG23	2.36	0.55
2:DV:152:TYR:O	2:DV:156:ILE:HG13	2.06	0.55

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:AD:3:ASP:H	2:AD:6:THR:HB	1.72	0.55
3:AE:134:LYS:HD3	3:AE:153:PHE:HB3	1.89	0.55
1:AJ:122:PRO:HG2	1:AJ:125:ALA:HB3	1.88	0.55
4:AK:64:PRO:CB	7:BF:105:PRO:HB2	2.36	0.55
1:AN:22:GLU:O	1:AN:26:ILE:HG13	2.07	0.55
2:AQ:14:VAL:CG1	6:BD:683:VAL:HG11	2.34	0.55
2:AU:110:ASN:ND2	5:BB:49:LYS:HA	2.20	0.55
1:AV:113:ARG:NH2	1:AV:161:SER:O	2.38	0.55
9:BH:288:TRP:HB3	9:BH:290:PHE:HE1	1.71	0.55
1:BI:20:PRO:HG3	1:BP:151:ALA:HB1	1.89	0.55
1:BK:25:ARG:HD3	6:CM:42:GLU:HG3	1.89	0.55
2:BL:132:ALA:O	2:BL:136:VAL:HG23	2.07	0.55
2:BN:15:GLN:HB3	2:BN:17:LYS:NZ	2.22	0.55
4:BS:20:ASP:O	4:BS:24:MET:HG2	2.07	0.55
4:BS:68:ARG:HB3	4:BS:69:PRO:HD2	1.87	0.55
2:BW:85:LEU:HD22	2:BW:133:ILE:CD1	2.35	0.55
1:CC:37:LEU:HD13	2:CD:28:LYS:CD	2.35	0.55
1:CE:12:ASP:OD2	2:CF:107:ARG:HD3	2.06	0.55
1:CG:115:MET:HE1	10:CG:200:CYC:CMB	2.33	0.55
9:CQ:34:GLU:HG2	9:CQ:174:GLU:HG2	1.89	0.55
2:CU:96:MET:HA	2:CU:152:TYR:CE2	2.42	0.55
1:CY:20:PRO:HA	1:CY:23:LEU:HD21	1.89	0.55
2:CZ:91:TYR:HB3	2:CZ:104:LEU:CD2	2.37	0.55
9:DI:47:MET:HE1	11:DI:400:45D:C34	2.36	0.55
9:DI:202:MET:SD	9:DI:252:PRO:HG2	2.46	0.55
9:EA:24:THR:HG21	9:EA:142:ILE:HD11	1.88	0.55
1:AA:137:ALA:HB1	1:AA:141:MET:HE3	1.89	0.55
2:AF:144:ASP:OD1	2:AF:144:ASP:N	2.40	0.55
2:AO:71:ASN:ND2	2:AO:121:VAL:HA	2.21	0.55
1:AX:101:VAL:HB	1:AX:155:PHE:CE2	2.42	0.55
6:BD:71:ALA:O	6:BD:75:ILE:HG12	2.06	0.55
6:BD:199:ILE:HG21	6:BD:227:LEU:HD22	1.89	0.55
6:BD:689:PRO:O	6:BD:693:GLU:HG3	2.06	0.55
2:BJ:106:GLU:HB2	5:CJ:57:THR:CG2	2.33	0.55
2:BL:22:ALA:O	2:BL:26:LYS:HG3	2.07	0.55
3:BM:66:ARG:NH2	3:BM:73:TYR:O	2.37	0.55
10:CD:200:CYC:HMA1	10:CD:200:CYC:NB	2.19	0.55
2:CH:112:LEU:HD11	2:CH:116:TYR:CZ	2.42	0.55
5:CJ:2:ARG:HG2	5:CJ:58:GLY:HA3	1.89	0.55
2:CU:88:TYR:CD2	2:CU:130:ILE:HD11	2.42	0.55
1:CY:85:MET:HA	1:CY:133:MET:HE1	1.87	0.55

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:DC:39:ILE:HG12	1:DC:145:ASP:HB3	1.87	0.55
1:DC:116:TYR:HD2	1:DC:123:ILE:HD12	1.70	0.55
9:DI:50:THR:HG21	9:DI:143:GLN:OE1	2.06	0.55
1:DQ:122:PRO:HD2	10:DQ:200:CYC:OC	2.07	0.55
2:DR:30:TYR:O	2:DR:33:SER:OG	2.18	0.55
2:DV:50:THR:O	2:DV:53:LYS:HB3	2.07	0.55
9:EA:190:ILE:CG2	9:EA:193:VAL:HB	2.36	0.55
1:AC:33:GLY:HA2	1:AC:36:ARG:CD	2.36	0.55
1:AH:62:ARG:HG2	1:AH:65:ILE:HG12	1.87	0.55
1:AH:109:LEU:CG	1:AH:159:LYS:HG2	2.37	0.55
2:AL:8:VAL:HG13	2:AL:23:ALA:HB1	1.87	0.55
1:AN:116:TYR:OH	10:AN:200:CYC:HHB	2.06	0.55
2:AO:85:LEU:HD22	2:AO:133:ILE:CD1	2.36	0.55
1:AV:72:ALA:HB2	10:AV:200:CYC:CMD	2.36	0.55
2:AW:56:VAL:HG12	2:AW:61:LEU:HG	1.87	0.55
1:AX:67:SER:O	1:AX:73:TYR:HB2	2.07	0.55
1:AX:115:MET:HE1	10:AX:200:CYC:CMB	2.35	0.55
5:BB:5:ARG:NE	5:BB:54:GLU:OE2	2.29	0.55
6:BD:765:SER:OG	10:CS:200:CYC:NB	2.30	0.55
1:BI:66:VAL:HA	1:BI:72:ALA:O	2.07	0.55
3:BM:7:VAL:HG11	3:BM:26:ILE:HD11	1.89	0.55
1:BP:83:ARG:HD2	8:CP:206:GLN:NE2	2.21	0.55
2:BT:39:ARG:O	2:BT:43:VAL:HG23	2.06	0.55
2:BT:134:LYS:HB2	2:BT:153:LEU:HD13	1.87	0.55
1:BV:27:LYS:O	1:BV:31:THR:HG23	2.07	0.55
1:BV:90:ARG:HG3	2:BW:18:TYR:CE1	2.42	0.55
2:CA:57:ALA:HA	2:CA:61:LEU:HD12	1.89	0.55
1:CE:4:VAL:O	1:CE:8:ILE:HG12	2.07	0.55
9:CQ:107:LEU:HG	11:CQ:400:45D:C30	2.37	0.55
1:CT:154:ASP:HB2	2:DD:46:ALA:CB	2.37	0.55
1:CV:62:ARG:O	1:CV:65:ILE:HG12	2.07	0.55
1:CY:57:ARG:O	1:CY:61:LYS:HG2	2.07	0.55
2:CZ:73:TYR:O	2:CZ:77:ARG:HB2	2.07	0.55
1:DA:61:LYS:CE	9:DI:56:PRO:HB2	2.37	0.55
2:DB:2:GLN:NE2	2:DB:10:ASN:OD1	2.37	0.55
1:DC:101:VAL:HB	1:DC:155:PHE:CE2	2.42	0.55
1:DJ:25:ARG:CD	1:DQ:25:ARG:HG3	2.37	0.55
2:DK:10:ASN:O	2:DK:14:VAL:HG13	2.07	0.55
2:DT:2:GLN:HE21	2:DT:10:ASN:HD22	1.55	0.55
2:DV:1:MET:HG3	2:DV:103:ILE:HB	1.87	0.55
1:AC:116:TYR:CE2	1:AC:126:VAL:HG21	2.42	0.55

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:AF:115:THR:HG23	5:BA:53:VAL:HG11	1.88	0.55
1:AH:156:VAL:HG12	1:AH:160:MET:HE3	1.89	0.55
1:AJ:19:SER:OG	1:AJ:22:GLU:HG3	2.06	0.55
1:AJ:56:ASP:O	1:AJ:60:GLN:HG2	2.07	0.55
2:AY:122:PRO:HB2	2:AY:125:SER:OG	2.07	0.55
6:BD:481:LYS:HE3	6:BD:615:TYR:CE2	2.41	0.55
9:BH:71:ILE:HD11	9:BH:94:LEU:CD2	2.36	0.55
9:BH:190:ILE:CG2	9:BH:193:VAL:HB	2.37	0.55
2:BQ:147:LYS:O	2:BQ:151:VAL:HG23	2.07	0.55
1:BR:83:ARG:NH2	1:BR:84:ASP:OD1	2.33	0.55
4:BS:51:ILE:HD13	4:BS:138:ILE:HD13	1.89	0.55
2:BT:56:VAL:HG12	2:BT:61:LEU:HG	1.89	0.55
1:BX:12:ASP:OD2	2:BY:107:ARG:NE	2.39	0.55
1:CC:50:ILE:HG12	1:CC:136:VAL:HG12	1.89	0.55
1:CC:65:ILE:HG22	1:CC:72:ALA:CB	2.37	0.55
1:CE:27:LYS:O	1:CE:30:VAL:HG12	2.06	0.55
9:CQ:242:ARG:HG3	9:CQ:243:GLU:OE1	2.07	0.55
1:CT:64:ASP:OD2	1:CY:70:GLY:HA2	2.06	0.55
1:CV:102:THR:OG1	1:CV:103:PRO:HD3	2.06	0.55
1:CV:130:VAL:HG21	1:CV:160:MET:HE1	1.88	0.55
1:CY:75:GLU:OE2	8:DY:223:ARG:HG2	2.07	0.55
2:DD:61:LEU:O	8:DH:205:PRO:HB3	2.07	0.55
9:DI:190:ILE:HB	9:DI:193:VAL:O	2.07	0.55
1:DQ:50:ILE:HD12	1:DQ:136:VAL:CG1	2.36	0.55
2:DR:64:ASP:HA	2:DR:67:ARG:HB2	1.88	0.55
2:DR:109:LEU:HD13	2:DR:159:GLY:HA3	1.89	0.55
1:DU:113:ARG:NH2	1:DU:160:MET:O	2.39	0.55
2:DV:90:ARG:HG2	2:DV:94:TYR:CZ	2.42	0.55
9:EA:292:LEU:HD23	9:EA:293:ASN:O	2.07	0.55
1:AA:116:TYR:CE2	1:AA:126:VAL:HG21	2.42	0.54
2:AD:57:ALA:HA	2:AD:61:LEU:CD1	2.36	0.54
2:AI:111:GLY:HA2	6:BD:427:VAL:HG22	1.90	0.54
4:AK:50:THR:HB	4:AK:54:ARG:NH1	2.22	0.54
2:AL:84:ASP:OD2	2:AL:116:TYR:OH	2.26	0.54
2:AY:126:THR:HB	10:AY:200:CYC:HMC1	1.89	0.54
6:BD:261:ALA:HB1	6:BD:402:GLY:HA3	1.87	0.54
2:BN:10:ASN:O	2:BN:14:VAL:HG23	2.07	0.54
2:BN:15:GLN:HB3	2:BN:17:LYS:HZ3	1.72	0.54
2:BT:36:LEU:CD1	7:CO:85:GLN:HB2	2.36	0.54
2:BT:97:LEU:CD1	6:CM:35:LEU:HD12	2.38	0.54
2:BW:84:ASP:OD2	10:BW:200:CYC:ND	2.40	0.54

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:CC:39:ILE:HG23	1:CC:141:MET:SD	2.47	0.54
1:CG:81:CYS:O	1:CG:85:MET:HG2	2.07	0.54
6:CM:131:PRO:O	6:CM:132:ILE:HD13	2.07	0.54
6:CM:706:ILE:HG12	2:DV:15:GLN:HE21	1.71	0.54
6:CM:796:PRO:O	6:CM:800:VAL:HG23	2.07	0.54
9:CQ:48:GLY:HA2	9:CQ:53:ILE:HG22	1.89	0.54
9:CQ:190:ILE:HD13	9:CQ:252:PRO:CB	2.37	0.54
1:DA:62:ARG:HD3	9:DI:58:ALA:CA	2.37	0.54
2:DB:112:LEU:HD11	2:DB:116:TYR:CE2	2.43	0.54
9:DI:41:TRP:NE1	9:DI:125:ILE:HD13	2.22	0.54
2:DR:104:LEU:HD13	2:DR:156:ILE:CG1	2.37	0.54
10:DR:200:CYC:HC	10:DR:200:CYC:HMD1	1.71	0.54
1:DU:104:ILE:HG21	1:DU:156:VAL:HG22	1.89	0.54
2:DV:126:THR:O	2:DV:130:ILE:HG13	2.07	0.54
2:DV:131:GLN:O	2:DV:134:LYS:HB3	2.06	0.54
2:AF:127:VAL:O	2:AF:131:GLN:HG2	2.07	0.54
2:AI:62:TYR:OH	10:AJ:200:CYC:HAD2	2.07	0.54
10:AO:200:CYC:HAA1	6:BD:485:ILE:HG23	1.89	0.54
2:AS:62:TYR:CE2	7:BF:103:PRO:HG3	2.42	0.54
2:AU:57:ALA:CB	2:AU:61:LEU:HD12	2.36	0.54
1:AV:83:ARG:NH2	1:AV:84:ASP:OD1	2.30	0.54
1:AX:85:MET:HB3	1:AX:133:MET:CE	2.38	0.54
5:BA:26:TYR:OH	5:BA:51:VAL:HG21	2.07	0.54
6:BD:5:ALA:HB3	6:BD:437:TYR:HD1	1.72	0.54
6:BD:72:ALA:CB	6:BD:156:MET:HE1	2.37	0.54
6:BD:799:LYS:HE2	6:BD:803:MET:SD	2.47	0.54
1:BK:92:VAL:O	1:BK:96:VAL:HG23	2.07	0.54
3:BM:89:LEU:O	3:BM:93:THR:HG23	2.07	0.54
2:BN:96:MET:HA	2:BN:152:TYR:CE2	2.42	0.54
2:BT:63:SER:O	2:BT:67:ARG:HG3	2.06	0.54
1:BV:83:ARG:HA	7:CO:110:PHE:CZ	2.42	0.54
1:BV:123:ILE:HD12	1:CE:117:ARG:HH22	1.71	0.54
2:BW:126:THR:HB	10:BW:200:CYC:HMC2	1.88	0.54
1:BX:58:LEU:HD22	1:BX:129:SER:CB	2.36	0.54
1:BZ:89:LEU:O	1:BZ:93:THR:HG23	2.07	0.54
2:CF:77:ARG:HB3	10:CF:200:CYC:HMD1	1.88	0.54
1:CG:58:LEU:HD22	1:CG:129:SER:CB	2.35	0.54
6:CM:501:VAL:HG21	6:CM:647:ASP:HB3	1.89	0.54
6:CM:526:LYS:HE2	6:CM:533:LYS:HB2	1.89	0.54
1:CV:95:GLY:HA3	1:CV:104:ILE:HD11	1.88	0.54
2:CW:114:GLU:HG2	5:DF:53:VAL:HG12	1.87	0.54

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:CY:92:VAL:O	1:CY:96:VAL:HG23	2.07	0.54
1:DA:41:GLU:CG	2:DB:24:MET:HE2	2.31	0.54
1:DC:51:VAL:HG12	1:DC:133:MET:CE	2.37	0.54
9:DI:107:LEU:HB3	11:DI:400:45D:C14	2.37	0.54
2:DK:43:VAL:CG1	2:DK:140:LEU:HG	2.38	0.54
1:DU:41:GLU:HG3	2:DV:24:MET:HE2	1.88	0.54
1:DU:121:THR:HG23	10:DU:200:CYC:C1C	2.37	0.54
1:AA:79:ALA:HA	1:AA:82:LEU:CD2	2.38	0.54
2:AB:123:ILE:O	2:AB:127:VAL:HG23	2.06	0.54
2:AD:64:ASP:HA	2:AD:67:ARG:NH1	2.23	0.54
2:AF:96:MET:HA	2:AF:152:TYR:CE2	2.43	0.54
1:AH:101:VAL:HB	1:AH:155:PHE:CE2	2.42	0.54
1:AN:39:ILE:HG12	1:AN:145:ASP:HB2	1.88	0.54
2:AO:74:THR:HG22	2:AO:76:ARG:HG2	1.90	0.54
2:AQ:75:THR:HG22	1:AR:115:MET:HE1	1.90	0.54
1:AR:122:PRO:O	1:AR:126:VAL:HG23	2.07	0.54
2:AU:1:MET:HE2	1:AZ:8:ILE:HG22	1.90	0.54
2:BJ:123:ILE:O	2:BJ:127:VAL:HG23	2.06	0.54
2:BL:65:VAL:C	2:BL:72:MET:HB3	2.31	0.54
3:BM:112:VAL:HG21	3:BM:160:MET:HE1	1.88	0.54
2:CA:71:ASN:HD22	2:CA:122:PRO:HD3	1.71	0.54
1:CC:116:TYR:CE2	1:CC:126:VAL:HG21	2.42	0.54
6:CM:777:PHE:CE1	6:CM:781:LEU:HD11	2.42	0.54
6:CM:795:TYR:HB3	6:CM:799:LYS:HB3	1.89	0.54
6:CM:820:GLN:HG2	10:DR:200:CYC:O1A	2.08	0.54
9:CQ:273:VAL:HB	9:CQ:286:ILE:CD1	2.38	0.54
1:CV:67:SER:O	1:CV:73:TYR:HB2	2.07	0.54
1:CV:121:THR:HG23	10:CV:200:CYC:C1C	2.38	0.54
2:CW:71:ASN:HA	2:CW:77:ARG:HH21	1.72	0.54
2:CW:103:ILE:O	2:CW:107:ARG:HB2	2.07	0.54
2:CZ:51:ILE:HG21	2:CZ:137:THR:CG2	2.37	0.54
1:DC:128:GLN:OE1	1:DC:131:ARG:NH1	2.40	0.54
5:DF:19:GLN:O	5:DF:21:GLU:HG2	2.07	0.54
5:DF:66:LEU:H	5:DF:66:LEU:HD23	1.71	0.54
9:DI:68:LEU:O	9:DI:71:ILE:HG22	2.08	0.54
1:DJ:20:PRO:HA	1:DJ:23:LEU:CD2	2.37	0.54
1:DL:2:SER:HG	2:DM:3:ASP:CG	2.15	0.54
2:DO:73:TYR:O	2:DO:77:ARG:HB2	2.06	0.54
2:DR:30:TYR:HE1	2:DR:99:GLY:HA3	1.73	0.54
9:EA:257:THR:HG23	9:EA:267:ILE:HG12	1.89	0.54
1:AC:122:PRO:HG2	1:AC:125:ALA:CB	2.37	0.54

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:AD:132:ALA:O	2:AD:136:VAL:HG23	2.08	0.54
2:AF:39:ARG:NE	1:CC:142:SER:HA	2.22	0.54
10:AF:200:CYC:O2A	5:BA:33:TYR:O	2.25	0.54
2:AI:144:ASP:OD1	2:AI:145:ALA:N	2.41	0.54
4:AK:72:ASN:ND2	4:AK:122:VAL:HA	2.22	0.54
2:AS:54:GLU:O	2:AS:58:LYS:HG2	2.07	0.54
2:AS:58:LYS:HB3	7:BF:98:VAL:CG1	2.38	0.54
6:BD:198:ALA:O	6:BD:202:MET:HG3	2.08	0.54
6:BD:348:ARG:HH21	6:BD:398:LEU:HD11	1.73	0.54
6:BD:797:ASN:OD1	6:BD:826:LEU:HD13	2.07	0.54
9:BH:99:ALA:HB1	9:BH:164:THR:HG22	1.88	0.54
9:BH:289:ARG:NH1	9:EA:306:LEU:HD12	2.23	0.54
2:BJ:122:PRO:HB2	2:BJ:125:SER:HG	1.72	0.54
1:BR:81:CYS:HA	10:BR:200:CYC:HHD	1.90	0.54
2:BW:85:LEU:HD21	10:BW:200:CYC:HBC1	1.90	0.54
2:BW:114:GLU:HA	2:BW:117:ASN:HB2	1.90	0.54
1:CG:89:LEU:O	1:CG:93:THR:HG23	2.07	0.54
5:CJ:6:ILE:HG21	5:CJ:36:TRP:CZ3	2.42	0.54
6:CM:450:GLY:O	6:CM:600:LYS:HE3	2.08	0.54
7:CO:103:PRO:HB3	7:CO:107:LEU:HD12	1.90	0.54
2:CS:108:VAL:HG13	2:CS:109:LEU:HG	1.89	0.54
1:CY:20:PRO:HA	1:CY:23:LEU:HG	1.87	0.54
2:CZ:47:ASN:HB2	2:CZ:50:THR:HB	1.88	0.54
2:DD:68:PRO:HG3	2:DD:73:TYR:CE1	2.42	0.54
5:DF:6:ILE:HD13	5:DF:36:TRP:HZ3	1.72	0.54
1:DJ:2:SER:OG	2:DK:3:ASP:OD2	2.10	0.54
1:DL:18:LEU:HD23	1:DL:22:GLU:OE1	2.07	0.54
1:DL:104:ILE:HG21	1:DL:156:VAL:HG22	1.88	0.54
10:DO:200:CYC:HMA3	10:DO:200:CYC:HBA1	1.90	0.54
1:DQ:85:MET:SD	1:DQ:133:MET:HE3	2.46	0.54
1:DS:97:VAL:HG21	2:DT:9:ILE:HD11	1.90	0.54
2:DV:44:ILE:HD11	2:DV:89:LEU:CD1	2.34	0.54
8:DY:197:GLN:OE1	8:DY:199:GLU:HB2	2.07	0.54
9:EA:269:VAL:CG2	9:EA:288:TRP:HB2	2.37	0.54
2:AF:71:ASN:ND2	2:AF:121:VAL:HA	2.22	0.54
2:AL:83:ARG:NH2	2:AL:84:ASP:OD1	2.36	0.54
1:AN:72:ALA:HA	1:AN:77:MET:HB3	1.88	0.54
2:AO:84:ASP:OD2	10:AO:200:CYC:ND	2.41	0.54
2:AO:141:VAL:HG12	2:AO:145:ALA:HB3	1.90	0.54
2:AU:85:LEU:HD22	2:AU:133:ILE:CD1	2.38	0.54
2:AU:132:ALA:HA	2:AU:135:GLU:OE1	2.07	0.54

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:AZ:13:ALA:HA	5:BB:45:LYS:NZ	2.22	0.54
1:AZ:142:SER:HB2	2:BN:39:ARG:CZ	2.38	0.54
6:BD:792:TYR:CG	6:BD:831:LEU:HD22	2.43	0.54
8:BG:232:ILE:HD12	8:BG:233:PRO:CD	2.37	0.54
9:BH:208:ASN:HA	9:BH:210:PHE:CE1	2.42	0.54
1:BK:22:GLU:HA	1:BK:25:ARG:HG2	1.90	0.54
3:BM:50:ILE:HG12	3:BM:140:LEU:HD13	1.89	0.54
1:BR:136:VAL:O	1:BR:140:LEU:HG	2.08	0.54
1:BV:115:MET:HG3	2:CA:78:TYR:HB3	1.88	0.54
2:BW:71:ASN:ND2	2:BW:121:VAL:HA	2.22	0.54
2:BY:126:THR:O	2:BY:130:ILE:HG13	2.08	0.54
1:BZ:46:SER:O	1:BZ:50:ILE:HD12	2.07	0.54
1:CE:37:LEU:HD11	2:CF:24:MET:SD	2.47	0.54
2:CH:88:TYR:OH	2:CH:112:LEU:HD21	2.07	0.54
5:CJ:14:THR:O	5:CJ:18:THR:HG23	2.07	0.54
6:CM:131:PRO:HG2	6:CM:200:GLN:HG2	1.90	0.54
6:CM:321:ARG:O	6:CM:325:LYS:HG2	2.06	0.54
6:CM:752:GLU:HB2	6:CM:753:PRO:HD3	1.90	0.54
1:CT:72:ALA:O	1:CT:78:THR:HG23	2.08	0.54
2:CW:65:VAL:HG12	2:CW:72:MET:CB	2.37	0.54
1:CY:20:PRO:HA	1:CY:23:LEU:CD2	2.38	0.54
2:CZ:96:MET:HA	2:CZ:152:TYR:CE2	2.42	0.54
9:DI:25:ILE:HD13	9:DI:153:VAL:HG13	1.89	0.54
9:DI:48:GLY:HA2	9:DI:53:ILE:HG22	1.89	0.54
1:DQ:57:ARG:O	1:DQ:61:LYS:HG2	2.07	0.54
10:DQ:200:CYC:HBB3	2:DV:73:TYR:CE1	2.42	0.54
2:DT:36:LEU:HD21	2:DT:145:ALA:HA	1.90	0.54
1:AA:85:MET:CE	1:AA:129:SER:HB2	2.36	0.54
1:AJ:124:GLU:OE1	1:AJ:124:GLU:N	2.37	0.54
2:AL:87:TYR:CD2	10:AL:200:CYC:HBB3	2.42	0.54
2:AQ:75:THR:CG2	1:AR:108:GLY:HA2	2.38	0.54
2:AS:72:MET:HE2	2:AS:81:CYS:SG	2.48	0.54
1:AX:84:ASP:OD2	10:AX:200:CYC:ND	2.38	0.54
1:AX:90:ARG:NH1	2:AY:16:GLY:HA2	2.23	0.54
1:AX:96:VAL:HG12	1:AX:152:TYR:CD2	2.43	0.54
1:AX:107:ILE:HG12	2:AY:13:ASP:OD2	2.07	0.54
9:BH:75:GLY:O	9:BH:78:GLN:HG2	2.08	0.54
9:BH:250:LEU:HD23	9:BH:273:VAL:HG23	1.89	0.54
1:BI:121:THR:HG23	10:BI:200:CYC:C1C	2.37	0.54
1:BK:25:ARG:HB3	6:CM:42:GLU:HA	1.89	0.54
2:BN:109:LEU:CD1	2:BN:159:GLY:HA3	2.38	0.54

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
4:BS:29:ALA:O	4:BS:32:GLU:HG3	2.07	0.54
4:BS:37:ARG:HH12	4:BS:152:PHE:HB2	1.73	0.54
4:BS:129:ARG:HH21	4:BS:133:ILE:HG13	1.73	0.54
1:BZ:138:SER:HA	1:BZ:146:ALA:HB1	1.88	0.54
2:CF:147:LYS:O	2:CF:151:VAL:HG23	2.07	0.54
6:CM:577:GLU:OE1	6:CM:577:GLU:N	2.40	0.54
9:CQ:36:GLN:O	9:CQ:40:ILE:HG23	2.08	0.54
1:CR:71:ASN:HD22	1:CR:121:THR:HA	1.71	0.54
2:CZ:63:SER:H	2:CZ:66:THR:HG22	1.73	0.54
1:DC:14:GLU:OE1	1:DC:16:ARG:NE	2.41	0.54
8:DH:203:PHE:CD1	8:DY:215:PRO:HB2	2.43	0.54
9:DI:151:ILE:HD11	11:DI:400:45D:H201	1.89	0.54
9:DI:292:LEU:HD23	9:DI:293:ASN:O	2.07	0.54
2:DM:72:MET:HE2	2:DM:81:CYS:SG	2.47	0.54
1:DQ:71:ASN:ND2	1:DQ:121:THR:HA	2.23	0.54
1:DS:115:MET:HE1	10:DS:200:CYC:CMB	2.35	0.54
1:DU:129:SER:O	1:DU:133:MET:HG3	2.08	0.54
2:DV:5:ILE:O	2:DV:9:ILE:HG12	2.08	0.54
9:EA:75:GLY:O	9:EA:78:GLN:HG2	2.08	0.54
3:AE:124:PRO:O	3:AE:127:VAL:HG12	2.08	0.54
2:AF:84:ASP:OD2	2:AF:116:TYR:OH	2.17	0.54
2:AF:104:LEU:O	2:AF:108:VAL:HB	2.08	0.54
2:AF:114:GLU:HG3	5:BA:53:VAL:HG12	1.88	0.54
1:AH:156:VAL:O	1:AH:160:MET:HG3	2.08	0.54
2:AL:96:MET:HA	2:AL:152:TYR:CE2	2.43	0.54
1:AV:50:ILE:HD11	1:AV:140:LEU:CD2	2.38	0.54
2:AW:122:PRO:HB2	2:AW:125:SER:HB2	1.88	0.54
1:AX:58:LEU:HD22	1:AX:129:SER:CB	2.36	0.54
1:AZ:131:ARG:HG3	1:AZ:157:ILE:HD13	1.90	0.54
1:BK:128:GLN:OE1	1:BK:131:ARG:HD2	2.08	0.54
2:BL:64:ASP:HA	2:BL:67:ARG:NH1	2.23	0.54
3:BM:30:LEU:HD13	2:BN:31:PHE:CD1	2.42	0.54
3:BM:68:PRO:HA	3:BM:73:TYR:CE1	2.43	0.54
1:CE:72:ALA:CA	1:CE:77:MET:HB3	2.38	0.54
1:CG:107:ILE:HG12	2:CH:13:ASP:OD2	2.07	0.54
6:CM:193:ASP:O	6:CM:197:VAL:HG22	2.08	0.54
6:CM:826:LEU:HA	6:CM:830:GLY:O	2.06	0.54
6:CM:848:PHE:O	6:CM:852:THR:HB	2.08	0.54
1:CR:28:ALA:O	1:CR:31:THR:HG22	2.07	0.54
1:CR:113:ARG:NH2	1:CR:161:SER:O	2.35	0.54
2:CU:53:LYS:HD3	1:CV:118:SER:O	2.08	0.54

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:CY:37:LEU:O	1:CY:40:ALA:HB3	2.07	0.54
2:CZ:39:ARG:HH12	2:CZ:141:VAL:HA	1.73	0.54
2:CZ:126:THR:HA	10:CZ:200:CYC:HMC1	1.89	0.54
1:DA:50:ILE:HD11	1:DA:137:ALA:N	2.23	0.54
1:DA:53:GLN:HB2	1:DA:57:ARG:NH1	2.22	0.54
9:DI:210:PHE:CG	9:DI:241:PHE:HB3	2.42	0.54
1:DJ:47:ARG:O	1:DJ:51:VAL:HG12	2.08	0.54
9:EA:233:GLY:O	9:EA:237:VAL:HG23	2.08	0.54
2:AB:40:ALA:O	2:AB:44:ILE:HG12	2.07	0.54
2:AB:104:LEU:HD13	2:AB:156:ILE:HG13	1.88	0.54
2:AD:5:ILE:HG23	2:AD:30:TYR:CD2	2.43	0.54
2:AD:83:ARG:NE	5:BA:22:LEU:HD21	2.22	0.54
3:AE:11:ALA:HB1	3:AE:16:ARG:O	2.08	0.54
1:AN:134:LYS:NZ	1:AN:154:ASP:OD1	2.26	0.54
2:AO:147:LYS:O	2:AO:151:VAL:HG23	2.07	0.54
1:AR:115:MET:HE1	10:AR:200:CYC:HMB3	1.89	0.54
1:AX:21:GLY:O	1:AX:25:ARG:HG3	2.07	0.54
2:AY:65:VAL:HA	2:AY:70:GLY:HA3	1.90	0.54
10:AY:200:CYC:HBD2	10:AY:200:CYC:HHA	1.90	0.54
6:BD:355:PRO:HB3	6:BD:361:VAL:HG22	1.90	0.54
1:BP:30:VAL:HG11	2:BQ:34:GLY:HA3	1.88	0.54
1:BP:101:VAL:HB	1:BP:155:PHE:CE2	2.42	0.54
1:BZ:63:PRO:HG2	8:CP:230:ASN:HD21	1.73	0.54
1:CE:65:ILE:HG23	1:CE:72:ALA:HB3	1.90	0.54
9:CQ:31:LEU:HD11	9:CQ:138:VAL:HG21	1.90	0.54
9:CQ:39:LEU:CD1	9:CQ:139:LEU:HB2	2.38	0.54
9:CQ:217:PHE:HA	9:CQ:298:ILE:CG2	2.38	0.54
2:CS:100:ASP:OD2	2:CS:102:SER:HB3	2.08	0.54
1:CT:84:ASP:OD2	10:CT:200:CYC:ND	2.41	0.54
2:CW:52:VAL:HA	2:CW:133:ILE:CD1	2.36	0.54
8:DG:249:ARG:HB2	2:DK:62:TYR:HB2	1.90	0.54
8:DH:227:PRO:HB3	1:DS:59:PHE:CD2	2.42	0.54
9:DI:25:ILE:CD1	9:DI:156:ASN:HB2	2.38	0.54
1:DJ:109:LEU:HD12	1:DJ:159:LYS:HB3	1.88	0.54
2:DK:62:TYR:OH	10:DL:200:CYC:HAD2	2.08	0.54
1:DN:102:THR:OG1	1:DN:103:PRO:HD3	2.07	0.54
2:DO:20:ASP:O	2:DO:24:MET:HG2	2.06	0.54
1:DQ:92:VAL:HG21	1:DQ:153:PHE:CE1	2.43	0.54
1:AC:91:LEU:HD21	1:AC:107:ILE:HG21	1.89	0.54
10:AC:200:CYC:HBB3	10:AC:200:CYC:HMB3	1.90	0.54
3:AE:66:ARG:NH2	3:AE:73:TYR:O	2.41	0.54

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:AJ:97:VAL:HG21	4:AK:19:LEU:HD12	1.90	0.54
1:AN:33:GLY:HA2	1:AN:36:ARG:HD2	1.90	0.54
1:AN:108:GLY:HA2	2:AS:75:THR:CG2	2.37	0.54
1:AR:56:ASP:CA	8:BG:225:ILE:HD11	2.37	0.54
1:AZ:22:GLU:O	1:AZ:26:ILE:HG12	2.07	0.54
1:AZ:68:PRO:HA	1:AZ:73:TYR:CG	2.43	0.54
6:BD:371:GLY:HA3	6:BD:375:ALA:HB2	1.89	0.54
6:BD:889:THR:HG22	2:CZ:118:SER:HB2	1.89	0.54
9:BH:156:ASN:O	9:BH:159:VAL:HG12	2.08	0.54
1:BI:81:CYS:O	1:BI:85:MET:HG2	2.07	0.54
1:BK:22:GLU:HG2	6:CM:42:GLU:OE2	2.08	0.54
1:BK:132:GLU:O	1:BK:136:VAL:HG23	2.08	0.54
3:BM:119:LEU:HD13	10:BM:200:CYC:CHA	2.38	0.54
4:BS:51:ILE:HA	4:BS:137:MET:HE1	1.90	0.54
2:BT:52:VAL:HG11	2:BT:82:ILE:HG23	1.89	0.54
2:BW:39:ARG:O	2:BW:43:VAL:HG23	2.07	0.54
2:CA:60:LEU:CD1	2:CA:72:MET:HE1	2.38	0.54
2:CA:85:LEU:HD22	2:CA:133:ILE:CD1	2.38	0.54
1:CC:58:LEU:HD22	1:CC:129:SER:CB	2.38	0.54
1:CC:93:THR:O	1:CC:97:VAL:HG23	2.08	0.54
1:CE:39:ILE:HG12	1:CE:145:ASP:HB3	1.89	0.54
2:CF:15:GLN:HB2	2:CF:17:LYS:NZ	2.23	0.54
2:CS:75:THR:HG22	1:CT:108:GLY:HA2	1.90	0.54
1:CT:53:GLN:O	1:CT:57:ARG:HG3	2.08	0.54
1:CV:37:LEU:HD22	2:CW:24:MET:CE	2.38	0.54
1:CY:77:MET:CE	8:DH:201:ARG:HB2	2.30	0.54
1:DA:97:VAL:CG2	2:DB:9:ILE:HD11	2.38	0.54
2:DB:1:MET:HE2	2:DB:103:ILE:HB	1.90	0.54
2:DD:116:TYR:HE2	2:DD:160:LEU:HD21	1.73	0.54
1:DJ:25:ARG:NE	1:DQ:25:ARG:HG3	2.23	0.54
1:DQ:115:MET:HG3	2:DV:75:THR:HG22	1.90	0.54
2:DT:72:MET:HE2	2:DT:78:TYR:CE2	2.43	0.54
10:AA:200:CYC:HAA1	2:AF:61:LEU:HD22	1.90	0.54
2:AD:57:ALA:HA	2:AD:61:LEU:HG	1.89	0.54
3:AE:30:LEU:HD12	2:AF:31:PHE:O	2.08	0.54
2:AF:109:LEU:CD1	2:AF:159:GLY:HA3	2.37	0.54
2:AF:147:LYS:O	2:AF:151:VAL:HG23	2.08	0.54
2:AL:25:ASP:HA	2:AL:28:LYS:HZ2	1.73	0.54
6:BD:274:SER:N	6:BD:277:GLU:OE2	2.36	0.54
6:BD:350:ILE:HD12	6:BD:380:LEU:HD13	1.90	0.54
9:BH:227:PHE:CE1	9:EA:230:PRO:HD3	2.43	0.54

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:BI:119:LEU:HA	2:BN:53:LYS:HZ3	1.73	0.54
2:BJ:30:TYR:OH	2:BJ:97:LEU:O	2.25	0.54
2:BL:83:ARG:NH1	5:CJ:22:LEU:HD21	2.23	0.54
1:BR:26:ILE:HD13	4:BS:98:ILE:HG21	1.90	0.54
10:BV:200:CYC:HBB2	2:CA:73:TYR:CE1	2.43	0.54
2:BW:25:ASP:HA	2:BW:28:LYS:NZ	2.23	0.54
2:BW:74:THR:HG22	2:BW:76:ARG:HG2	1.90	0.54
2:CA:147:LYS:HE3	7:CO:93:PRO:CG	2.38	0.54
1:CE:109:LEU:HD22	1:CE:159:LYS:HB3	1.90	0.54
6:CM:864:ALA:C	6:CM:867:PRO:HD2	2.33	0.54
9:CQ:98:TYR:HA	9:CQ:101:TRP:CE3	2.43	0.54
2:CS:145:ALA:HA	2:CS:148:GLU:OE1	2.08	0.54
2:CW:72:MET:HE2	2:CW:78:TYR:CE2	2.43	0.54
1:CY:20:PRO:HA	1:CY:23:LEU:CG	2.38	0.54
1:DA:132:GLU:O	1:DA:136:VAL:HG12	2.08	0.54
2:DM:104:LEU:HD12	2:DM:155:TYR:HD2	1.72	0.54
1:DQ:64:ASP:HA	1:DQ:67:SER:OG	2.08	0.54
2:DR:75:THR:HG22	1:DS:115:MET:SD	2.48	0.54
2:DR:137:THR:O	2:DR:141:VAL:HG22	2.08	0.54
1:DU:61:LYS:NZ	1:DU:132:GLU:OE2	2.37	0.54
2:DV:47:ASN:O	2:DV:51:ILE:HG13	2.08	0.54
8:DZ:243:GLU:OE1	8:DZ:243:GLU:N	2.41	0.54
1:AC:96:VAL:HA	1:AC:152:TYR:CE2	2.43	0.53
1:AH:35:ALA:O	1:AH:39:ILE:HG13	2.07	0.53
2:AL:81:CYS:HA	10:AL:200:CYC:HHD	1.90	0.53
1:AR:126:VAL:HG11	1:AR:160:MET:HE1	1.90	0.53
1:AV:91:LEU:HD21	1:AV:107:ILE:CG2	2.38	0.53
1:AX:6:LYS:HG2	6:BD:534:PHE:HZ	1.72	0.53
1:AX:85:MET:CE	1:AX:129:SER:HB2	2.37	0.53
6:BD:210:PHE:CZ	6:BD:212:ASN:HB2	2.43	0.53
6:BD:248:ILE:HG12	6:BD:356:SER:CB	2.38	0.53
2:BL:3:ASP:OD1	2:BL:98:ALA:HB1	2.09	0.53
1:BV:2:SER:O	1:BV:5:THR:OG1	2.17	0.53
2:CF:23:ALA:HA	2:CF:26:LYS:HG2	1.89	0.53
1:CG:26:ILE:O	1:CG:30:VAL:HG23	2.08	0.53
2:CH:25:ASP:HA	2:CH:28:LYS:HE3	1.90	0.53
5:CJ:40:GLN:HG2	5:CJ:44:MET:CE	2.38	0.53
6:CM:736:LYS:O	6:CM:740:ARG:HG3	2.08	0.53
9:CQ:42:PHE:HD2	9:CQ:131:LEU:HD13	1.74	0.53
1:CR:8:ILE:CD1	2:CS:98:ALA:HB2	2.38	0.53
1:CR:73:TYR:O	1:CR:77:MET:HB2	2.08	0.53

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:CR:131:ARG:HG2	1:CR:157:ILE:CD1	2.33	0.53
2:CS:47:ASN:HB3	2:CS:50:THR:HB	1.90	0.53
2:CW:54:GLU:HG3	9:DI:63:LEU:CD2	2.38	0.53
2:CZ:5:ILE:O	2:CZ:8:VAL:HG22	2.08	0.53
2:CZ:8:VAL:HG21	2:CZ:27:LEU:HD21	1.89	0.53
2:CZ:19:LEU:HD12	2:CZ:23:ALA:HB1	1.90	0.53
2:CZ:149:MET:O	2:CZ:153:LEU:HD23	2.08	0.53
1:DA:102:THR:OG1	1:DA:103:PRO:HD3	2.07	0.53
8:DH:227:PRO:CB	1:DS:66:VAL:HG21	2.38	0.53
1:DS:104:ILE:HG22	1:DS:109:LEU:CD2	2.38	0.53
2:DT:83:ARG:NH1	10:DT:200:CYC:O1A	2.41	0.53
5:DX:62:THR:O	5:DX:66:LEU:HD11	2.08	0.53
1:AH:62:ARG:NH2	1:AH:124:GLU:OE1	2.38	0.53
2:AI:8:VAL:CG2	2:AI:27:LEU:HD11	2.30	0.53
4:AK:60:PHE:O	4:AK:64:PRO:HA	2.08	0.53
1:AV:72:ALA:CA	1:AV:77:MET:HB3	2.38	0.53
2:AW:1:MET:N	5:BB:67:ALA:O	2.41	0.53
9:BH:190:ILE:HD11	9:BH:254:ARG:O	2.09	0.53
2:BJ:103:ILE:CD1	2:BJ:107:ARG:HD2	2.35	0.53
2:BL:39:ARG:NH1	2:BL:39:ARG:HA	2.23	0.53
3:BM:104:ILE:HD13	3:BM:156:ILE:HD11	1.88	0.53
2:BQ:78:TYR:O	2:BQ:82:ILE:HG12	2.08	0.53
2:BQ:117:ASN:O	6:CM:439:ARG:NH1	2.40	0.53
4:BS:64:PRO:HB2	7:CO:105:PRO:HB2	1.89	0.53
1:BX:97:VAL:HG21	2:BY:19:LEU:HD13	1.91	0.53
1:CG:142:SER:OG	1:CG:144:ASP:OD1	2.19	0.53
2:CH:41:ALA:HB2	2:CH:96:MET:HE1	1.90	0.53
5:CK:41:GLN:HG3	5:CK:45:LYS:HD2	1.91	0.53
6:CM:77:THR:O	6:CM:140:ASN:HA	2.08	0.53
6:CM:160:LEU:O	6:CM:164:THR:HG23	2.08	0.53
6:CM:701:ARG:O	6:CM:706:ILE:HD11	2.09	0.53
1:CR:75:GLU:HG3	9:DI:152:THR:OG1	2.07	0.53
2:CS:96:MET:HA	2:CS:152:TYR:CE2	2.43	0.53
1:CT:88:TYR:OH	1:CT:160:MET:HE1	2.08	0.53
1:CV:134:LYS:HE3	1:CV:150:SER:CB	2.38	0.53
1:CY:86:ASP:O	1:CY:90:ARG:HG3	2.09	0.53
1:CY:102:THR:O	1:CY:106:GLU:HG2	2.08	0.53
1:DA:14:GLU:OE1	1:DA:16:ARG:NE	2.29	0.53
10:DC:200:CYC:NC	10:DC:200:CYC:HMD1	2.23	0.53
8:DH:219:LEU:HD22	1:DS:79:ALA:CB	2.39	0.53
2:DM:112:LEU:HD11	2:DM:116:TYR:CZ	2.43	0.53

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:DR:68:PRO:HA	2:DR:73:TYR:CG	2.42	0.53
2:DV:105:ASP:OD1	2:DV:155:TYR:OH	2.12	0.53
1:AC:89:LEU:O	1:AC:93:THR:HG23	2.08	0.53
3:AE:4:VAL:HG13	2:AF:97:LEU:HD23	1.91	0.53
3:AE:71:ASN:ND2	3:AE:121:VAL:HG22	2.21	0.53
2:AF:55:ALA:O	2:AF:59:SER:OG	2.16	0.53
1:AH:82:LEU:HD13	8:BG:203:PHE:HB2	1.90	0.53
2:AL:96:MET:SD	2:AL:149:MET:HG2	2.49	0.53
10:AP:200:CYC:HMD1	10:AP:200:CYC:NC	2.21	0.53
2:AS:96:MET:HA	2:AS:152:TYR:CE2	2.43	0.53
2:AU:36:LEU:CD2	2:AU:145:ALA:HB2	2.38	0.53
1:AZ:41:GLU:HA	1:AZ:44:THR:HG22	1.91	0.53
6:BD:268:ALA:HB3	6:BD:270:LYS:NZ	2.24	0.53
6:BD:353:ARG:HG2	6:BD:399:ARG:NH2	2.22	0.53
6:BD:682:ARG:NH2	6:BD:684:ASP:OD2	2.32	0.53
6:BD:861:LEU:HG	10:DB:200:CYC:OB	2.08	0.53
1:BI:116:TYR:CE2	1:BI:126:VAL:HG21	2.43	0.53
2:BJ:3:ASP:N	2:BJ:6:THR:OG1	2.42	0.53
2:BL:47:ASN:O	2:BL:51:ILE:HD12	2.08	0.53
1:BP:110:VAL:HG11	2:BT:76:ARG:HG3	1.91	0.53
2:BT:47:ASN:OD1	2:BT:140:LEU:HD21	2.08	0.53
2:BY:61:LEU:HD22	10:BZ:200:CYC:HAA1	1.90	0.53
10:BY:200:CYC:HHA	10:BY:200:CYC:HBA2	1.89	0.53
2:CA:119:LEU:HB3	2:CA:121:VAL:HG23	1.89	0.53
2:CH:1:MET:CE	2:CH:103:ILE:HB	2.38	0.53
1:CR:79:ALA:C	8:DZ:246:VAL:HG22	2.33	0.53
1:CR:82:LEU:HD13	8:DZ:242:ILE:HD11	1.90	0.53
2:CS:51:ILE:HG23	2:CS:136:VAL:HB	1.91	0.53
2:CU:3:ASP:OD1	2:CU:6:THR:N	2.27	0.53
1:CV:89:LEU:O	1:CV:93:THR:HG23	2.09	0.53
2:CW:96:MET:HA	2:CW:152:TYR:CE2	2.44	0.53
1:DA:47:ARG:O	1:DA:50:ILE:HG22	2.08	0.53
1:DA:159:LYS:HE2	1:DA:159:LYS:HA	1.90	0.53
5:DF:2:ARG:HA	5:DF:57:THR:O	2.08	0.53
9:DI:201:TYR:HA	9:DI:213:LEU:HD11	1.89	0.53
1:DJ:85:MET:HE2	1:DJ:129:SER:OG	2.09	0.53
2:DM:1:MET:SD	2:DM:103:ILE:HD12	2.48	0.53
1:DQ:113:ARG:NH2	1:DQ:161:SER:O	2.33	0.53
2:AD:71:ASN:HB3	10:AD:200:CYC:OC	2.08	0.53
2:AI:2:GLN:HG2	2:AI:6:THR:CG2	2.39	0.53
1:AN:27:LYS:O	1:AN:31:THR:HG23	2.08	0.53

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:AQ:50:THR:O	2:AQ:54:GLU:HG2	2.09	0.53
1:AR:56:ASP:HA	8:BG:225:ILE:HD11	1.90	0.53
1:AX:26:ILE:O	1:AX:30:VAL:HG23	2.09	0.53
1:AX:109:LEU:HD11	1:AX:159:LYS:CB	2.38	0.53
1:AZ:96:VAL:HG12	1:AZ:152:TYR:CD2	2.43	0.53
6:BD:72:ALA:HB1	6:BD:81:PRO:CB	2.39	0.53
6:BD:76:PHE:CD2	6:BD:149:MET:HE1	2.44	0.53
6:BD:351:LEU:O	6:BD:399:ARG:NH2	2.41	0.53
6:BD:353:ARG:NH2	6:BD:385:GLU:OE1	2.38	0.53
6:BD:553:ARG:HG3	6:BD:554:ASP:O	2.08	0.53
1:BK:111:GLY:HA2	1:BK:114:GLU:OE2	2.08	0.53
2:BN:114:GLU:CG	5:CJ:53:VAL:HG22	2.37	0.53
2:BW:44:ILE:HD13	2:BW:149:MET:HE2	1.90	0.53
1:CC:11:ALA:HB1	1:CC:16:ARG:O	2.08	0.53
1:CC:131:ARG:O	1:CC:135:GLU:HG2	2.08	0.53
10:CC:200:CYC:HMB3	10:CC:200:CYC:HBB3	1.90	0.53
2:CF:8:VAL:CG1	2:CF:23:ALA:HB1	2.36	0.53
1:CG:36:ARG:NH2	1:CG:99:GLY:HA2	2.23	0.53
7:CO:114:ALA:O	7:CO:117:VAL:HG12	2.08	0.53
1:CR:104:ILE:H	1:CR:104:ILE:HD12	1.72	0.53
1:DC:134:LYS:HD3	1:DC:153:PHE:HB3	1.91	0.53
2:DD:124:SER:O	2:DD:128:GLN:HG3	2.08	0.53
5:DF:2:ARG:CG	5:DF:55:LEU:HD11	2.38	0.53
8:DH:232:ILE:CD1	8:DH:233:PRO:HD2	2.37	0.53
9:DI:99:ALA:HB1	9:DI:164:THR:CG2	2.38	0.53
1:DQ:95:GLY:CA	1:DQ:104:ILE:HD11	2.31	0.53
1:DS:61:LYS:HG2	9:EA:57:GLY:CA	2.39	0.53
2:DV:10:ASN:O	2:DV:14:VAL:HG23	2.08	0.53
2:DV:41:ALA:CB	2:DV:97:LEU:HD21	2.25	0.53
9:EA:211:ASP:O	9:EA:214:ILE:HG22	2.09	0.53
1:AA:37:LEU:O	1:AA:41:GLU:HG3	2.09	0.53
2:AD:53:LYS:NZ	3:AE:120:ASN:OD1	2.29	0.53
3:AE:4:VAL:HG13	2:AF:97:LEU:CD2	2.39	0.53
4:AK:52:VAL:HG13	4:AK:86:MET:HG2	1.90	0.53
2:AL:42:SER:OG	6:BD:40:LEU:HD13	2.07	0.53
1:AR:41:GLU:OE2	2:AS:24:MET:HG3	2.08	0.53
2:AU:1:MET:SD	1:AZ:5:THR:HG23	2.49	0.53
1:AX:85:MET:O	1:AX:133:MET:HE1	2.08	0.53
6:BD:788:MET:HE1	1:CY:9:VAL:CG1	2.38	0.53
10:BK:200:CYC:HBB3	10:BK:200:CYC:HMB3	1.90	0.53
2:BL:144:ASP:OD1	2:BL:145:ALA:N	2.41	0.53

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:BM:4:VAL:HG13	2:BN:97:LEU:HD23	1.91	0.53
2:BT:87:TYR:CG	10:BT:200:CYC:HBB3	2.44	0.53
2:BY:100:ASP:OD1	2:BY:102:SER:OG	2.13	0.53
2:CD:2:GLN:HB3	2:CD:102:SER:HB3	1.91	0.53
6:CM:353:ARG:HB3	6:CM:399:ARG:HE	1.74	0.53
6:CM:776:GLU:OE1	6:CM:776:GLU:N	2.41	0.53
9:CQ:277:TRP:HD1	9:CQ:305:LEU:HD21	1.73	0.53
9:CQ:292:LEU:HD23	9:CQ:298:ILE:CA	2.36	0.53
1:CT:71:ASN:HB3	10:CT:200:CYC:OC	2.07	0.53
2:CW:123:ILE:HA	2:CW:126:THR:HG22	1.91	0.53
1:CY:71:ASN:HD21	1:CY:121:THR:HG22	1.72	0.53
2:DD:22:ALA:O	2:DD:26:LYS:HG3	2.07	0.53
9:DI:224:GLN:OE1	9:DI:230:PRO:HG3	2.09	0.53
9:DI:314:ASN:HD22	9:DI:314:ASN:C	2.12	0.53
1:DJ:25:ARG:HG2	1:DQ:25:ARG:HE	1.72	0.53
2:DK:47:ASN:O	2:DK:51:ILE:HD12	2.08	0.53
2:DM:44:ILE:O	2:DM:48:ALA:HB2	2.09	0.53
1:DU:10:ASN:O	1:DU:14:GLU:HG3	2.08	0.53
9:EA:125:ILE:HG22	11:EA:400:45D:C25	2.39	0.53
10:AA:200:CYC:CAA	2:AF:61:LEU:HD22	2.38	0.53
1:AC:47:ARG:O	1:AC:51:VAL:HG23	2.08	0.53
1:AC:137:ALA:O	1:AC:141:MET:HG2	2.08	0.53
3:AE:6:GLN:O	3:AE:10:GLN:HG2	2.09	0.53
1:AN:122:PRO:HB2	1:AN:125:ALA:CB	2.39	0.53
5:BB:19:GLN:O	5:BB:21:GLU:HG2	2.08	0.53
8:BG:214:ASN:O	8:BG:217:ASN:ND2	2.31	0.53
9:BH:239:ARG:HA	9:BH:242:ARG:HG2	1.90	0.53
9:BH:311:GLU:CD	9:BH:312:LEU:H	2.17	0.53
1:BI:71:ASN:HD22	1:BI:121:THR:HA	1.74	0.53
2:BL:1:MET:SD	2:BL:103:ILE:HD13	2.48	0.53
2:BN:41:ALA:N	2:BN:96:MET:HE1	2.23	0.53
2:BN:59:SER:O	2:BN:128:GLN:NE2	2.42	0.53
2:BQ:71:ASN:HD22	2:BQ:122:PRO:HD2	1.73	0.53
1:BV:71:ASN:ND2	1:BV:119:LEU:O	2.41	0.53
1:BX:68:PRO:HG3	1:BX:73:TYR:CE1	2.43	0.53
1:BX:131:ARG:HA	1:BX:157:ILE:HD11	1.90	0.53
2:CD:96:MET:HA	2:CD:152:TYR:CE2	2.43	0.53
1:CE:141:MET:HE1	1:CE:145:ASP:CB	2.37	0.53
2:CF:24:MET:HG3	2:CF:28:LYS:NZ	2.23	0.53
9:CQ:300:PHE:CE2	9:CQ:302:ALA:HB2	2.43	0.53
9:CQ:311:GLU:CD	9:CQ:312:LEU:H	2.17	0.53

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:DJ:90:ARG:NH2	2:DO:73:TYR:OH	2.39	0.53
2:DM:54:GLU:O	2:DM:58:LYS:HG2	2.09	0.53
2:DO:111:GLY:HA2	2:DO:114:GLU:OE2	2.08	0.53
2:DT:123:ILE:O	2:DT:127:VAL:HG23	2.07	0.53
1:AA:35:ALA:HA	1:AA:38:ARG:NH1	2.23	0.53
2:AI:61:LEU:HD21	2:AI:78:TYR:OH	2.08	0.53
4:AK:92:TYR:CE2	10:AK:200:CYC:HBB1	2.44	0.53
2:AL:90:ARG:HD3	6:BD:33:ARG:C	2.34	0.53
1:AN:23:LEU:HD13	2:AO:42:SER:HB2	1.90	0.53
1:AR:85:MET:CE	1:AR:133:MET:HE2	2.39	0.53
2:AS:39:ARG:NH1	2:AS:39:ARG:HA	2.23	0.53
1:AX:102:THR:HG21	6:BD:534:PHE:CE1	2.44	0.53
5:BB:26:TYR:HE1	5:BB:51:VAL:HG21	1.73	0.53
6:BD:664:GLY:O	6:BD:668:ARG:HG3	2.08	0.53
1:BI:80:THR:HG21	2:BN:62:TYR:HE1	1.73	0.53
3:BM:47:GLU:HG2	3:BM:89:LEU:CD2	2.38	0.53
1:BR:114:GLU:OE1	1:BR:114:GLU:N	2.41	0.53
2:BT:96:MET:HA	2:BT:152:TYR:CE2	2.44	0.53
1:CC:41:GLU:HA	1:CC:44:THR:HG22	1.90	0.53
1:CC:90:ARG:NH1	2:CD:16:GLY:O	2.41	0.53
1:CC:93:THR:O	1:CC:96:VAL:HG22	2.09	0.53
1:CG:92:VAL:HG21	1:CG:153:PHE:CE1	2.44	0.53
2:CH:47:ASN:O	2:CH:51:ILE:HG13	2.07	0.53
1:CY:37:LEU:HD21	2:CZ:24:MET:HB3	1.90	0.53
1:DA:12:ASP:OD2	2:DB:107:ARG:HD3	2.08	0.53
1:DA:85:MET:CE	1:DA:129:SER:HB2	2.39	0.53
2:DK:126:THR:HG21	2:DK:160:LEU:HD13	1.90	0.53
1:DL:122:PRO:HD2	10:DL:200:CYC:OC	2.07	0.53
2:DM:122:PRO:HG2	10:DM:200:CYC:CMC	2.38	0.53
9:EA:195:ASN:HB3	9:EA:198:VAL:CG1	2.32	0.53
1:AC:105:GLU:HA	1:AC:109:LEU:HB2	1.91	0.53
2:AD:1:MET:SD	2:AD:103:ILE:HD13	2.49	0.53
2:AI:68:PRO:O	2:AO:124:SER:HB2	2.09	0.53
10:AL:200:CYC:HAA1	6:BD:424:PHE:CZ	2.39	0.53
2:AQ:147:LYS:O	2:AQ:151:VAL:HG23	2.09	0.53
1:AV:91:LEU:HD21	1:AV:107:ILE:HG21	1.91	0.53
2:AW:23:ALA:HA	2:AW:26:LYS:HG2	1.90	0.53
2:AY:88:TYR:CE2	2:AY:130:ILE:HD11	2.44	0.53
1:AZ:75:GLU:OE2	1:AZ:76:GLU:HG2	2.09	0.53
5:BA:6:ILE:HG21	5:BA:36:TRP:CZ3	2.43	0.53
2:BL:66:THR:HG21	10:BM:200:CYC:CBA	2.37	0.53

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:BN:101:ALA:HB1	2:BN:104:LEU:HD12	1.91	0.53
1:BR:14:GLU:OE1	1:BR:16:ARG:NE	2.26	0.53
2:BW:74:THR:CG2	2:BW:76:ARG:HG2	2.38	0.53
10:CE:200:CYC:HMD1	10:CE:200:CYC:NC	2.21	0.53
10:CF:200:CYC:HHA	10:CF:200:CYC:CBA	2.38	0.53
2:CW:15:GLN:HG3	2:CW:17:LYS:NZ	2.24	0.53
10:CW:200:CYC:HMA3	10:CW:200:CYC:NB	2.22	0.53
2:CZ:59:SER:CB	2:CZ:128:GLN:HG2	2.35	0.53
2:DB:101:ALA:HB2	2:DB:152:TYR:CD1	2.44	0.53
2:DD:116:TYR:CD2	2:DD:160:LEU:HD11	2.44	0.53
9:DI:36:GLN:O	9:DI:40:ILE:HG23	2.08	0.53
9:DI:44:TYR:OH	9:DI:118:GLU:OE1	2.16	0.53
9:DI:75:GLY:O	9:DI:78:GLN:HG2	2.08	0.53
9:DI:214:ILE:HD13	9:DI:238:LEU:HB2	1.90	0.53
1:DJ:51:VAL:HG23	1:DJ:85:MET:HB3	1.89	0.53
2:DM:71:ASN:O	2:DM:77:ARG:HD3	2.09	0.53
1:DN:126:VAL:HG22	10:DN:200:CYC:HMC2	1.89	0.53
2:DR:105:ASP:OD2	2:DR:155:TYR:OH	2.25	0.53
2:DV:109:LEU:HD21	2:DV:156:ILE:HG23	1.91	0.53
8:DZ:214:ASN:OD1	8:DZ:216:MET:HG2	2.09	0.53
9:EA:36:GLN:O	9:EA:40:ILE:HG23	2.09	0.53
1:AA:85:MET:O	1:AA:133:MET:HE1	2.09	0.53
1:AC:58:LEU:HD22	1:AC:129:SER:CB	2.37	0.53
2:AO:92:ALA:CB	2:AO:149:MET:HE3	2.38	0.53
1:AZ:56:ASP:O	1:AZ:60:GLN:HG2	2.09	0.53
6:BD:314:ILE:HG21	6:BD:322:ARG:HH12	1.74	0.53
6:BD:620:MET:O	6:BD:624:PHE:HB2	2.08	0.53
1:BK:96:VAL:HA	1:BK:152:TYR:CE2	2.42	0.53
3:BM:85:TYR:HE1	3:BM:129:ALA:HB1	1.74	0.53
10:BM:200:CYC:HMD1	10:BM:200:CYC:NC	2.19	0.53
2:BN:76:ARG:NE	5:CJ:66:LEU:HD11	2.24	0.53
2:BN:127:VAL:O	2:BN:131:GLN:HG2	2.09	0.53
2:BQ:83:ARG:HD2	6:CM:369:SER:CB	2.39	0.53
1:BX:105:GLU:OE2	1:BX:159:LYS:NZ	2.41	0.53
1:CC:5:THR:HG23	2:CD:1:MET:SD	2.49	0.53
1:CC:115:MET:HE1	10:CC:200:CYC:CMB	2.37	0.53
6:CM:543:ILE:HG12	6:CM:568:LEU:HD23	1.91	0.53
6:CM:597:TYR:HD2	6:CM:600:LYS:HG3	1.74	0.53
9:CQ:258:GLU:HB2	9:CQ:259:PRO:HD2	1.91	0.53
1:CT:101:VAL:HB	1:CT:155:PHE:CE2	2.44	0.53
1:CV:16:ARG:O	2:CW:90:ARG:HD2	2.09	0.53

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:CY:50:ILE:HD12	1:CY:136:VAL:CG1	2.39	0.53
2:CZ:73:TYR:O	2:CZ:74:THR:OG1	2.24	0.53
9:DI:211:ASP:O	9:DI:214:ILE:HG22	2.08	0.53
2:DK:56:VAL:HG12	2:DK:61:LEU:HG	1.91	0.53
1:DN:141:MET:HE3	1:DN:146:ALA:CA	2.34	0.53
2:DR:40:ALA:CB	2:DR:145:ALA:HB1	2.39	0.53
9:EA:33:ALA:CB	9:EA:174:GLU:HB2	2.39	0.53
2:AD:22:ALA:O	2:AD:26:LYS:HG3	2.09	0.53
2:AF:10:ASN:O	2:AF:14:VAL:HG23	2.09	0.53
1:AJ:3:ILE:HG21	1:AJ:25:ARG:CZ	2.39	0.53
1:AP:153:PHE:O	1:AP:157:ILE:HG13	2.08	0.53
1:AR:73:TYR:CD2	9:BH:151:ILE:HD13	2.44	0.53
1:AV:65:ILE:CG2	1:AV:72:ALA:HB3	2.39	0.53
1:AZ:116:TYR:CE2	1:AZ:126:VAL:HG21	2.44	0.53
6:BD:50:GLY:O	6:BD:54:LEU:HG	2.09	0.53
9:BH:90:ALA:O	9:BH:163:PHE:HB2	2.09	0.53
10:BQ:200:CYC:HB	10:BQ:200:CYC:CMA	2.22	0.53
4:BS:64:PRO:CG	7:CO:105:PRO:HB2	2.39	0.53
2:BT:144:ASP:HB2	7:CO:86:VAL:CG1	2.39	0.53
2:BY:60:LEU:HB3	2:BY:72:MET:CE	2.38	0.53
1:BZ:101:VAL:HG22	1:BZ:104:ILE:HB	1.91	0.53
1:BZ:151:ALA:CB	1:CE:20:PRO:HB3	2.39	0.53
1:CC:131:ARG:HA	1:CC:157:ILE:HD11	1.91	0.53
1:CC:141:MET:HE3	1:CC:145:ASP:C	2.34	0.53
6:CM:821:GLN:O	6:CM:825:ILE:HG13	2.09	0.53
9:CQ:28:PHE:CD1	9:CQ:157:ALA:HB1	2.44	0.53
10:CS:200:CYC:NB	10:CS:200:CYC:HMA1	2.23	0.53
1:CT:122:PRO:HD2	10:CT:200:CYC:OC	2.08	0.53
5:DF:59:ARG:CD	5:DF:60:PRO:HD2	2.39	0.53
9:DI:187:LYS:HD3	9:DI:203:ASP:OD2	2.09	0.53
2:DM:104:LEU:O	2:DM:108:VAL:HB	2.09	0.53
1:DU:134:LYS:HD3	1:DU:153:PHE:HB3	1.91	0.53
2:DV:68:PRO:HG3	2:DV:73:TYR:CE1	2.44	0.53
2:AD:131:GLN:HA	2:AD:134:LYS:HD3	1.91	0.52
3:AE:89:LEU:O	3:AE:93:THR:HG23	2.09	0.52
3:AE:92:VAL:HG11	3:AE:153:PHE:CZ	2.45	0.52
2:AI:78:TYR:CD1	1:AJ:115:MET:HG3	2.43	0.52
1:AN:97:VAL:HG21	2:AO:19:LEU:HD12	1.91	0.52
1:AP:114:GLU:HG3	1:AP:117:ARG:NH2	2.24	0.52
1:AR:52:LYS:CE	8:BG:225:ILE:HB	2.40	0.52
1:AZ:58:LEU:HD22	1:AZ:129:SER:HB3	1.91	0.52

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
6:BD:413:MET:HE2	6:BD:433:THR:CB	2.37	0.52
6:BD:795:TYR:HB3	6:BD:799:LYS:HB3	1.90	0.52
9:BH:190:ILE:HD13	9:BH:252:PRO:HB2	1.90	0.52
2:BL:3:ASP:H	2:BL:6:THR:HB	1.73	0.52
2:BL:92:ALA:CB	2:BL:149:MET:HE3	2.39	0.52
2:BT:36:LEU:HD12	7:CO:85:GLN:HB2	1.91	0.52
1:BV:98:SER:HA	2:BW:5:ILE:HG21	1.91	0.52
1:CE:98:SER:HA	2:CF:5:ILE:HG21	1.91	0.52
1:CG:39:ILE:HG23	1:CG:141:MET:SD	2.49	0.52
6:CM:69:SER:CA	6:CM:82:LEU:HD21	2.32	0.52
9:CQ:63:LEU:HD12	9:CQ:105:ILE:CD1	2.38	0.52
9:CQ:75:GLY:O	9:CQ:78:GLN:HG2	2.09	0.52
9:CQ:264:PHE:CD2	9:DI:309:PRO:HG3	2.44	0.52
1:CR:85:MET:HE2	1:CR:133:MET:HE3	1.91	0.52
2:CS:1:MET:HE3	2:CS:103:ILE:HB	1.90	0.52
2:CS:46:ALA:HB2	1:CY:154:ASP:OD2	2.08	0.52
1:CT:68:PRO:HA	1:CT:73:TYR:CG	2.45	0.52
1:DA:61:LYS:HD2	9:DI:56:PRO:HD2	1.91	0.52
1:DA:72:ALA:HA	1:DA:77:MET:HB3	1.91	0.52
9:DI:131:LEU:HG	9:DI:135:ALA:HB3	1.91	0.52
1:DL:96:VAL:HA	1:DL:152:TYR:CZ	2.44	0.52
1:DN:14:GLU:OE1	1:DN:16:ARG:NE	2.38	0.52
1:DQ:102:THR:O	1:DQ:105:GLU:HG2	2.10	0.52
2:DT:57:ALA:HA	2:DT:61:LEU:CD1	2.38	0.52
2:DT:73:TYR:O	2:DT:77:ARG:HB2	2.09	0.52
9:EA:258:GLU:OE1	9:EA:266:GLN:HG3	2.09	0.52
1:AA:89:LEU:O	1:AA:93:THR:HG23	2.09	0.52
1:AA:100:ASP:CG	1:AH:21:GLY:HA3	2.33	0.52
1:AC:128:GLN:O	1:AC:131:ARG:HB2	2.10	0.52
3:AE:151:PRO:HB2	1:AJ:20:PRO:HB3	1.92	0.52
2:AF:115:THR:CG2	5:BA:53:VAL:HG11	2.40	0.52
1:AH:89:LEU:O	1:AH:93:THR:HG23	2.08	0.52
1:AH:110:VAL:CG1	2:AL:76:ARG:HG3	2.39	0.52
1:AJ:4:VAL:HG12	1:AJ:8:ILE:HD12	1.92	0.52
4:AK:142:ALA:HB1	4:AK:147:LEU:HD12	1.90	0.52
2:AL:2:GLN:HB2	2:AL:6:THR:HG23	1.90	0.52
1:AP:68:PRO:HG3	1:AP:73:TYR:CE1	2.44	0.52
1:AP:76:GLU:OE1	1:AP:76:GLU:N	2.39	0.52
2:AU:44:ILE:HG21	2:AU:93:THR:OG1	2.08	0.52
2:AY:63:SER:H	2:AY:66:THR:HG22	1.74	0.52
6:BD:316:MET:CE	6:BD:395:VAL:HG22	2.40	0.52

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
9:BH:31:LEU:HD11	9:BH:138:VAL:HG21	1.92	0.52
1:BK:33:GLY:O	1:BK:36:ARG:HG2	2.09	0.52
1:BK:97:VAL:HG21	2:BL:19:LEU:HD11	1.90	0.52
2:BL:96:MET:HB2	2:BL:148:GLU:OE1	2.09	0.52
2:BL:105:ASP:OD2	2:BL:155:TYR:OH	2.25	0.52
3:BM:87:TRP:O	3:BM:91:LEU:HD13	2.10	0.52
1:BP:27:LYS:HG3	2:BQ:35:GLU:OE1	2.09	0.52
1:BR:89:LEU:HB2	1:BR:133:MET:CE	2.37	0.52
1:BV:5:THR:HA	2:BW:1:MET:HE3	1.91	0.52
1:BZ:76:GLU:HG3	9:CQ:11:ILE:HG21	1.91	0.52
1:CE:30:VAL:HA	2:CF:31:PHE:CD1	2.44	0.52
1:CE:44:THR:HG22	2:CF:18:TYR:HD2	1.74	0.52
1:CE:50:ILE:CD1	1:CE:136:VAL:HG12	2.37	0.52
1:CE:93:THR:O	1:CE:97:VAL:HG13	2.09	0.52
1:CG:109:LEU:HD11	1:CG:159:LYS:CB	2.39	0.52
6:CM:183:LYS:O	6:CM:186:ILE:HG22	2.10	0.52
2:CS:10:ASN:O	2:CS:14:VAL:HG23	2.08	0.52
1:CV:101:VAL:HB	1:CV:155:PHE:CE2	2.44	0.52
1:DA:53:GLN:CG	1:DA:136:VAL:HG21	2.40	0.52
1:DJ:25:ARG:HG2	1:DQ:25:ARG:HG3	1.90	0.52
1:DJ:124:GLU:OE1	1:DJ:124:GLU:N	2.34	0.52
1:DQ:27:LYS:HZ3	2:DR:39:ARG:HG2	1.73	0.52
10:DQ:200:CYC:CBB	2:DV:74:THR:HA	2.38	0.52
2:DR:57:ALA:HA	2:DR:61:LEU:CB	2.38	0.52
1:DS:38:ARG:HA	1:DS:41:GLU:OE1	2.09	0.52
2:DT:147:LYS:O	2:DT:151:VAL:HG23	2.10	0.52
5:DX:5:ARG:CG	5:DX:56:ALA:HB2	2.37	0.52
8:DZ:239:ASN:OD1	8:DZ:240:ILE:HD12	2.09	0.52
9:EA:288:TRP:CD1	9:EA:303:ILE:HG12	2.39	0.52
3:AE:101:LYS:HE3	1:AJ:16:ARG:HE	1.73	0.52
4:AK:54:ARG:HB2	4:AK:137:MET:HE1	1.91	0.52
10:AN:200:CYC:HBB2	2:AS:73:TYR:CE1	2.44	0.52
1:AV:102:THR:HB	1:AV:103:PRO:HD3	1.91	0.52
1:AX:109:LEU:HD11	1:AX:159:LYS:HB3	1.92	0.52
2:AY:47:ASN:O	2:AY:51:ILE:HG13	2.09	0.52
6:BD:481:LYS:HE3	6:BD:615:TYR:CD2	2.44	0.52
6:BD:691:PHE:CG	1:DC:82:LEU:HB3	2.45	0.52
1:BK:16:ARG:HG2	1:BK:17:TYR:O	2.09	0.52
1:BK:152:TYR:O	1:BK:156:VAL:HG23	2.10	0.52
2:BL:112:LEU:HD21	10:BL:200:CYC:HAB1	1.91	0.52
3:BM:14:GLN:HB3	3:BM:16:ARG:HG2	1.90	0.52

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:BT:31:PHE:CD1	6:CM:50:GLY:HA3	2.44	0.52
1:BV:97:VAL:HG21	2:BW:19:LEU:HD12	1.90	0.52
2:BY:52:VAL:O	2:BY:56:VAL:HG13	2.09	0.52
1:CC:12:ASP:OD2	2:CD:107:ARG:NH1	2.43	0.52
1:CC:89:LEU:HB2	1:CC:133:MET:CE	2.39	0.52
2:CD:24:MET:O	2:CD:28:LYS:HG2	2.09	0.52
6:CM:158:TRP:HB3	10:CM:902:CYC:HMB2	1.92	0.52
6:CM:171:ASP:OD2	6:CM:173:ASN:HB2	2.09	0.52
6:CM:282:ILE:HG21	6:CM:306:GLU:HG2	1.91	0.52
6:CM:794:PRO:HA	2:DR:106:GLU:HG3	1.91	0.52
10:CM:901:CYC:HMA3	10:CM:901:CYC:NB	2.18	0.52
2:CS:30:TYR:HE1	2:CS:99:GLY:HA3	1.73	0.52
2:CS:88:TYR:OH	2:CS:116:TYR:OH	2.28	0.52
2:CU:122:PRO:HG2	10:CU:200:CYC:CMC	2.40	0.52
10:CY:200:CYC:HAB2	2:DD:74:THR:HA	1.92	0.52
1:DA:83:ARG:HG2	8:DZ:218:PHE:CE2	2.44	0.52
2:DD:137:THR:O	2:DD:141:VAL:HG22	2.10	0.52
8:DG:215:PRO:HB2	8:DZ:203:PHE:CE1	2.44	0.52
1:DJ:67:SER:O	1:DJ:73:TYR:HB2	2.09	0.52
2:DK:85:LEU:HG	10:DK:200:CYC:HBC1	1.91	0.52
1:DL:33:GLY:HA3	2:DM:31:PHE:CE2	2.44	0.52
1:DL:50:ILE:HG13	1:DL:136:VAL:CB	2.38	0.52
1:DL:66:VAL:HA	1:DL:72:ALA:HB3	1.91	0.52
2:DM:123:ILE:O	2:DM:127:VAL:HG23	2.08	0.52
1:DN:97:VAL:HG11	2:DO:19:LEU:HD12	1.91	0.52
1:DQ:92:VAL:HG11	1:DQ:153:PHE:CZ	2.44	0.52
2:DR:55:ALA:CB	2:DR:133:ILE:HG13	2.39	0.52
3:AE:105:GLU:HA	3:AE:109:LEU:CB	2.34	0.52
1:AH:126:VAL:O	1:AH:130:VAL:HG23	2.09	0.52
2:AQ:53:LYS:HE3	2:AQ:54:GLU:OE2	2.10	0.52
6:BD:347:PHE:CD1	6:BD:380:LEU:HD21	2.44	0.52
6:BD:551:PHE:HE1	6:BD:588:PHE:HB2	1.74	0.52
6:BD:609:LEU:C	6:BD:610:LEU:HD12	2.34	0.52
6:BD:631:GLY:O	6:BD:635:VAL:HG23	2.09	0.52
6:BD:801:ILE:O	6:BD:805:THR:HG22	2.09	0.52
7:BF:113:MET:HA	7:BF:116:GLN:NE2	2.20	0.52
9:BH:65:GLU:OE1	9:BH:69:LYS:HE2	2.10	0.52
2:BL:37:ARG:HH21	2:BL:99:GLY:HA2	1.73	0.52
4:BS:166:GLU:HB3	2:CA:77:ARG:HH22	1.74	0.52
2:BW:77:ARG:HB3	10:BW:200:CYC:HMD1	1.92	0.52
1:BX:111:GLY:HA2	1:BX:114:GLU:OE1	2.10	0.52

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:BZ:39:ILE:HG23	1:BZ:141:MET:SD	2.49	0.52
2:CF:103:ILE:CD1	2:CF:107:ARG:HD2	2.40	0.52
1:CG:21:GLY:O	1:CG:25:ARG:HG3	2.08	0.52
1:CG:50:ILE:HG13	1:CG:136:VAL:CG1	2.38	0.52
2:CS:55:ALA:O	2:CS:59:SER:OG	2.07	0.52
1:CT:27:LYS:NZ	2:CU:38:VAL:HB	2.24	0.52
1:CY:47:ARG:HD2	2:CZ:18:TYR:CE1	2.44	0.52
2:DB:123:ILE:O	2:DB:127:VAL:HG23	2.09	0.52
9:DI:25:ILE:HD12	9:DI:156:ASN:HB2	1.89	0.52
10:DM:200:CYC:OB	5:DX:20:ARG:HA	2.08	0.52
2:DO:3:ASP:H	2:DO:6:THR:HG1	1.56	0.52
2:DO:83:ARG:NH2	2:DO:84:ASP:OD1	2.28	0.52
9:EA:40:ILE:HG13	11:EA:400:45D:H371	1.91	0.52
1:AC:111:GLY:HA2	1:AC:114:GLU:OE2	2.09	0.52
2:AD:142:GLY:O	2:AD:146:GLY:N	2.43	0.52
1:AR:63:PRO:HG2	8:BG:230:ASN:HD21	1.73	0.52
2:AS:84:ASP:OD2	2:AS:116:TYR:OH	2.13	0.52
2:AY:75:THR:HG22	1:AZ:107:ILE:O	2.09	0.52
1:AZ:141:MET:HE3	1:AZ:145:ASP:C	2.34	0.52
1:BV:132:GLU:HA	1:BV:135:GLU:OE1	2.08	0.52
2:BW:96:MET:HG3	2:BW:148:GLU:CD	2.34	0.52
2:BY:117:ASN:O	6:CM:467:ARG:NH2	2.42	0.52
1:CC:6:LYS:NZ	1:CC:102:THR:OG1	2.43	0.52
1:CC:88:TYR:O	1:CC:92:VAL:HG23	2.09	0.52
5:CK:19:GLN:O	5:CK:21:GLU:HG2	2.08	0.52
6:CM:273:LEU:HB3	6:CM:277:GLU:HG2	1.90	0.52
6:CM:401:LEU:HD13	6:CM:411:TRP:CH2	2.39	0.52
1:CR:154:ASP:CB	2:CZ:46:ALA:HB2	2.38	0.52
1:CT:18:LEU:HD13	2:CU:97:LEU:HD23	1.91	0.52
1:CT:56:ASP:OD1	8:DY:237:ALA:HB3	2.10	0.52
1:CT:65:ILE:O	1:CT:72:ALA:N	2.38	0.52
2:CW:127:VAL:O	2:CW:131:GLN:HG3	2.10	0.52
2:CZ:52:VAL:HG23	2:CZ:85:LEU:HD13	1.92	0.52
10:CZ:200:CYC:HMD1	10:CZ:200:CYC:NC	2.24	0.52
1:DA:105:GLU:O	1:DA:110:VAL:HG23	2.09	0.52
9:DI:32:ASN:O	9:DI:36:GLN:HG3	2.10	0.52
2:DK:41:ALA:CA	2:DK:96:MET:HE1	2.38	0.52
1:DQ:92:VAL:O	1:DQ:96:VAL:HG13	2.09	0.52
1:DS:102:THR:OG1	1:DS:103:PRO:HD3	2.10	0.52
2:DT:10:ASN:O	2:DT:14:VAL:HG13	2.09	0.52
9:EA:89:ARG:NE	9:EA:160:ASP:OD1	2.43	0.52

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:AE:4:VAL:O	3:AE:8:ILE:HG12	2.09	0.52
3:AE:115:MET:HE2	10:AE:200:CYC:NB	2.23	0.52
1:AJ:105:GLU:O	1:AJ:110:VAL:HG23	2.08	0.52
1:AN:132:GLU:HA	1:AN:135:GLU:OE1	2.10	0.52
2:AO:39:ARG:O	2:AO:43:VAL:HG23	2.10	0.52
1:AR:18:LEU:HD12	2:AS:97:LEU:CD1	2.36	0.52
1:AR:115:MET:HE1	10:AR:200:CYC:CMB	2.40	0.52
5:BB:23:GLN:OE1	5:BB:23:GLN:N	2.38	0.52
6:BD:244:PRO:HD2	6:BD:248:ILE:HB	1.92	0.52
9:BH:131:LEU:HG	9:BH:135:ALA:HB3	1.92	0.52
1:BV:58:LEU:HD13	1:BV:129:SER:CA	2.39	0.52
1:BV:101:VAL:HG12	1:BV:152:TYR:HD1	1.75	0.52
1:BZ:35:ALA:O	1:BZ:39:ILE:HG13	2.10	0.52
2:CD:63:SER:H	2:CD:66:THR:HG22	1.75	0.52
2:CD:65:VAL:CG1	2:CD:72:MET:HE2	2.36	0.52
1:CE:113:ARG:NH2	1:CE:161:SER:OXT	2.37	0.52
2:CF:123:ILE:O	2:CF:127:VAL:HG23	2.10	0.52
1:CG:85:MET:O	1:CG:133:MET:HE1	2.10	0.52
6:CM:186:ILE:O	6:CM:190:CYS:HB3	2.09	0.52
6:CM:453:PRO:HG3	6:CM:462:PHE:CE1	2.45	0.52
1:CT:113:ARG:HD3	1:CT:123:ILE:HD13	1.91	0.52
2:CW:15:GLN:CG	2:CW:17:LYS:HG2	2.40	0.52
1:CY:38:ARG:HA	1:CY:41:GLU:OE1	2.09	0.52
1:DA:92:VAL:HG11	1:DA:153:PHE:CE2	2.44	0.52
2:DD:23:ALA:O	2:DD:27:LEU:HG	2.09	0.52
9:DI:67:ALA:HB3	9:DI:101:TRP:HH2	1.75	0.52
9:DI:117:MET:CE	11:DI:400:45D:H273	2.37	0.52
2:DK:71:ASN:ND2	2:DK:121:VAL:HA	2.24	0.52
1:DL:128:GLN:O	1:DL:132:GLU:HG2	2.08	0.52
1:DQ:126:VAL:HG13	1:DQ:160:MET:SD	2.49	0.52
5:DX:6:ILE:HG22	5:DX:53:VAL:CG2	2.30	0.52
1:AA:115:MET:HE3	1:AA:116:TYR:CE1	2.45	0.52
1:AC:132:GLU:O	1:AC:136:VAL:HG23	2.09	0.52
3:AE:14:GLN:HB3	3:AE:16:ARG:HG2	1.91	0.52
2:AF:71:ASN:O	2:AF:77:ARG:HD2	2.10	0.52
1:AH:109:LEU:CD2	1:AH:159:LYS:HG2	2.39	0.52
1:AH:131:ARG:HG3	1:AH:157:ILE:HD13	1.92	0.52
2:AL:25:ASP:HA	2:AL:28:LYS:NZ	2.25	0.52
1:AN:8:ILE:HD11	2:AO:97:LEU:HD13	1.92	0.52
2:AQ:122:PRO:HB2	2:AQ:125:SER:HB2	1.92	0.52
10:AQ:200:CYC:HHA	10:AQ:200:CYC:HBD1	1.91	0.52

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:AR:81:CYS:SG	10:AR:200:CYC:HAC2	2.50	0.52
2:AU:123:ILE:HA	2:AU:126:THR:HG22	1.91	0.52
2:AW:2:GLN:O	2:AW:102:SER:OG	2.27	0.52
2:AY:10:ASN:O	2:AY:14:VAL:HG23	2.10	0.52
7:BF:112:GLU:O	7:BF:116:GLN:HG2	2.09	0.52
8:BG:226:ASN:HD21	9:BH:146:GLU:HG3	1.75	0.52
9:BH:210:PHE:HA	9:BH:213:LEU:HB3	1.90	0.52
9:BH:233:GLY:O	9:BH:237:VAL:HG23	2.10	0.52
1:BK:39:ILE:HD12	1:BK:145:ASP:CG	2.35	0.52
2:BL:112:LEU:HD21	10:BL:200:CYC:C3B	2.40	0.52
2:BN:5:ILE:O	2:BN:8:VAL:HG22	2.10	0.52
1:BP:109:LEU:HD22	1:BP:159:LYS:HG2	1.90	0.52
6:CM:52:LEU:HA	6:CM:55:GLU:OE1	2.09	0.52
6:CM:607:ARG:HG2	6:CM:613:PRO:HA	1.92	0.52
6:CM:745:GLN:OE1	6:CM:855:TYR:HA	2.10	0.52
1:CT:61:LYS:NZ	1:CT:132:GLU:OE2	2.43	0.52
1:CV:46:SER:HB2	1:CV:140:LEU:HD23	1.91	0.52
1:CV:58:LEU:HD13	1:CV:129:SER:N	2.25	0.52
1:CY:127:ALA:HB2	1:CY:161:SER:HB2	1.92	0.52
1:CY:137:ALA:HB1	1:CY:141:MET:CE	2.38	0.52
1:DJ:83:ARG:NH2	10:DJ:200:CYC:O2A	2.42	0.52
2:DK:91:TYR:CD2	2:DK:108:VAL:HB	2.43	0.52
1:DL:71:ASN:HB3	10:DL:200:CYC:OC	2.09	0.52
1:DN:22:GLU:O	1:DN:26:ILE:HG12	2.10	0.52
1:DQ:116:TYR:HD2	1:DQ:123:ILE:HG13	1.74	0.52
2:DR:15:GLN:CB	2:DR:17:LYS:HE3	2.40	0.52
2:DV:96:MET:HA	2:DV:152:TYR:CE2	2.45	0.52
1:AC:4:VAL:HG23	2:AD:3:ASP:OD2	2.09	0.52
2:AD:109:LEU:HD23	2:AD:159:GLY:HA3	1.92	0.52
3:AE:7:VAL:HG13	3:AE:22:GLU:HB3	1.91	0.52
2:AL:1:MET:HE2	6:BD:26:SER:OG	2.10	0.52
2:AO:85:LEU:HD22	2:AO:133:ILE:HD11	1.92	0.52
1:AP:100:ASP:OD2	1:AP:102:THR:OG1	2.15	0.52
1:AR:39:ILE:HD11	1:AR:148:GLU:OE2	2.09	0.52
6:BD:177:VAL:HG21	6:BD:260:ALA:HB2	1.91	0.52
6:BD:290:PHE:O	6:BD:292:ARG:NH1	2.43	0.52
1:BI:27:LYS:O	1:BI:31:THR:HG23	2.10	0.52
1:BI:90:ARG:HD3	2:BJ:18:TYR:CD1	2.45	0.52
2:BJ:40:ALA:O	2:BJ:44:ILE:HG12	2.10	0.52
2:BJ:61:LEU:HD21	2:BJ:78:TYR:OH	2.09	0.52
1:BK:71:ASN:ND2	10:BK:200:CYC:OC	2.33	0.52

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:BM:110:ILE:HG22	5:CJ:16:ILE:CD1	2.37	0.52
2:BN:4:ALA:HB2	2:BN:99:GLY:O	2.10	0.52
2:BN:44:ILE:CD1	2:BN:149:MET:HE3	2.40	0.52
1:BR:2:SER:N	1:BR:100:ASP:OD2	2.43	0.52
2:BY:116:TYR:CD2	2:BY:160:LEU:HD11	2.44	0.52
1:CE:102:THR:O	1:CE:106:GLU:HG2	2.10	0.52
10:CF:200:CYC:HB	6:CM:566:ILE:HD12	1.74	0.52
5:CJ:12:SER:CB	5:CJ:15:ARG:HG3	2.40	0.52
6:CM:291:GLU:HG2	6:CM:348:ARG:NH1	2.24	0.52
6:CM:365:PHE:HA	6:CM:368:VAL:HG12	1.92	0.52
2:CS:43:VAL:HG11	2:CS:141:VAL:HG12	1.91	0.52
1:CT:63:PRO:HG2	1:CY:68:PRO:O	2.09	0.52
1:CY:112:VAL:HA	1:CY:115:MET:CG	2.40	0.52
2:DD:108:VAL:HG22	10:DD:200:CYC:CBB	2.40	0.52
1:DJ:53:GLN:O	1:DJ:57:ARG:HG3	2.09	0.52
2:DM:85:LEU:HD22	2:DM:133:ILE:CD1	2.40	0.52
2:DR:59:SER:OG	2:DR:132:ALA:HB2	2.10	0.52
3:AE:47:GLU:OE1	2:AF:18:TYR:OH	2.24	0.52
2:AF:96:MET:HA	2:AF:152:TYR:HE2	1.74	0.52
4:AK:52:VAL:HG13	4:AK:86:MET:CG	2.40	0.52
4:AK:161:THR:O	4:AK:165:SER:CB	2.57	0.52
2:AL:52:VAL:O	2:AL:56:VAL:HG23	2.10	0.52
1:AN:127:ALA:HB1	1:AN:161:SER:HB3	1.91	0.52
1:AR:46:SER:O	1:AR:50:ILE:HD12	2.08	0.52
1:AR:156:VAL:O	1:AR:160:MET:HG3	2.10	0.52
1:AZ:131:ARG:O	1:AZ:135:GLU:HG2	2.10	0.52
9:BH:91:ASP:OD1	9:BH:96:ARG:NH2	2.38	0.52
9:BH:264:PHE:CD1	9:BH:294:PRO:HD3	2.42	0.52
1:BK:117:ARG:HD2	1:BP:161:SER:CB	2.40	0.52
1:BP:38:ARG:HA	1:BP:41:GLU:OE1	2.10	0.52
4:BS:124:ILE:HD11	4:BS:164:LEU:HD12	1.91	0.52
1:BZ:97:VAL:HG21	2:CA:19:LEU:CD1	2.39	0.52
2:CA:19:LEU:CB	2:CA:24:MET:HE2	2.40	0.52
1:CC:113:ARG:NH1	1:CC:160:MET:O	2.42	0.52
6:CM:749:ARG:NH1	10:DV:200:CYC:HBA1	2.25	0.52
9:CQ:217:PHE:HA	9:CQ:298:ILE:HG23	1.91	0.52
9:CQ:227:PHE:CE1	9:DI:230:PRO:HD3	2.45	0.52
2:CS:1:MET:CG	2:CS:103:ILE:HB	2.39	0.52
1:CT:27:LYS:NZ	2:CU:35:GLU:HA	2.25	0.52
1:CT:154:ASP:HB2	2:DD:46:ALA:HB1	1.92	0.52
1:CY:23:LEU:O	1:CY:27:LYS:HG2	2.10	0.52

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:DD:71:ASN:HD22	2:DD:121:VAL:HA	1.75	0.52
2:DD:123:ILE:O	2:DD:127:VAL:HG23	2.10	0.52
9:DI:27:ARG:HA	9:DI:30:GLN:OE1	2.10	0.52
1:DJ:65:ILE:HG22	1:DJ:72:ALA:HB3	1.92	0.52
1:DN:115:MET:HE1	10:DN:200:CYC:C1B	2.39	0.52
1:DU:57:ARG:NH2	1:DU:135:GLU:HB3	2.24	0.52
5:DX:3:MET:H	5:DX:57:THR:HB	1.75	0.52
9:EA:42:PHE:CD2	9:EA:131:LEU:HD13	2.45	0.52
2:AB:77:ARG:HB3	10:AB:200:CYC:HMD1	1.91	0.52
10:AD:200:CYC:CHB	5:BA:22:LEU:HD23	2.39	0.52
1:AH:13:ALA:HB2	6:BD:330:ARG:HD2	1.91	0.52
1:AJ:89:LEU:HB2	1:AJ:133:MET:CE	2.40	0.52
2:AL:63:SER:O	2:AL:67:ARG:HG3	2.10	0.52
1:AN:33:GLY:HA2	1:AN:36:ARG:CD	2.40	0.52
2:AO:74:THR:CG2	2:AO:76:ARG:HG2	2.40	0.52
1:AP:115:MET:HE1	10:AP:200:CYC:CMB	2.40	0.52
2:AY:41:ALA:HB2	2:AY:96:MET:CE	2.38	0.52
5:BA:29:LYS:HE2	6:BD:392:GLU:OE1	2.10	0.52
6:BD:52:LEU:HA	6:BD:55:GLU:OE1	2.09	0.52
6:BD:183:LYS:HG2	6:BD:192:ILE:HD12	1.92	0.52
9:BH:300:PHE:CE2	9:BH:302:ALA:HB2	2.45	0.52
1:BI:118:SER:O	2:BN:53:LYS:HD3	2.10	0.52
4:BS:138:ILE:HG22	4:BS:153:ILE:HG21	1.92	0.52
1:BV:2:SER:OG	2:BW:3:ASP:OD2	2.24	0.52
1:BZ:73:TYR:CD2	9:CQ:151:ILE:HD13	2.45	0.52
2:CF:34:GLY:O	2:CF:38:VAL:HG23	2.10	0.52
2:CH:119:LEU:HD11	10:CH:200:CYC:HAA2	1.91	0.52
8:CP:196:TRP:O	8:CP:200:VAL:HG23	2.10	0.52
1:CR:116:TYR:HD2	1:CR:123:ILE:HG22	1.74	0.52
1:CT:105:GLU:HG2	1:CT:110:VAL:HG23	1.92	0.52
1:DA:72:ALA:HB2	10:DA:200:CYC:CMD	2.39	0.52
5:DF:5:ARG:HD2	5:DF:54:GLU:OE2	2.09	0.52
9:DI:34:GLU:HB2	9:DI:174:GLU:CG	2.40	0.52
1:DL:104:ILE:HG22	1:DL:109:LEU:CD1	2.40	0.52
1:DN:128:GLN:O	1:DN:132:GLU:HG2	2.10	0.52
1:DQ:89:LEU:HD22	2:DR:18:TYR:OH	2.10	0.52
1:DS:26:ILE:O	1:DS:30:VAL:HG22	2.09	0.52
10:DS:200:CYC:O1D	8:DY:201:ARG:NE	2.42	0.52
2:AF:72:MET:HE2	2:AF:78:TYR:CD2	2.45	0.51
2:AF:90:ARG:HG2	2:AF:94:TYR:CZ	2.45	0.51
2:AL:91:TYR:CE2	10:AL:200:CYC:HBB1	2.45	0.51

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:AR:97:VAL:HG11	2:AS:27:LEU:HD13	1.92	0.51
2:AS:92:ALA:CB	2:AS:149:MET:HE1	2.38	0.51
2:AW:71:ASN:ND2	2:AW:121:VAL:HA	2.25	0.51
1:AZ:93:THR:O	1:AZ:96:VAL:HG22	2.10	0.51
6:BD:444:GLN:NE2	6:BD:472:ARG:O	2.20	0.51
6:BD:736:LYS:HE3	6:BD:740:ARG:CD	2.39	0.51
9:BH:74:MET:O	9:BH:79:GLN:NE2	2.43	0.51
10:BI:200:CYC:HMB3	10:BI:200:CYC:HBB3	1.90	0.51
3:BM:36:ARG:HD2	3:BM:96:VAL:O	2.09	0.51
1:BP:141:MET:HE1	1:BP:149:ALA:HB3	1.92	0.51
2:BQ:8:VAL:HG21	2:BQ:27:LEU:CD1	2.37	0.51
4:BS:40:ALA:O	4:BS:44:ILE:HG12	2.10	0.51
2:BW:5:ILE:HD11	2:BW:31:PHE:CE1	2.45	0.51
1:BX:97:VAL:HG21	2:BY:19:LEU:CD1	2.40	0.51
1:BX:143:SER:OG	1:CV:147:ALA:HB2	2.10	0.51
2:BY:104:LEU:HD22	2:BY:156:ILE:HD11	1.92	0.51
2:CD:2:GLN:HB3	2:CD:102:SER:CB	2.40	0.51
2:CD:68:PRO:HG3	2:CD:73:TYR:CE1	2.45	0.51
6:CM:210:PHE:CZ	6:CM:212:ASN:HB2	2.45	0.51
6:CM:210:PHE:HZ	6:CM:216:ALA:HB1	1.74	0.51
6:CM:371:GLY:HA3	6:CM:375:ALA:HB2	1.92	0.51
9:CQ:42:PHE:CD2	9:CQ:131:LEU:HD13	2.45	0.51
9:CQ:195:ASN:O	9:CQ:199:LEU:HG	2.10	0.51
1:CR:8:ILE:HD13	2:CS:1:MET:HE1	1.91	0.51
2:DB:112:LEU:HD11	2:DB:116:TYR:CZ	2.45	0.51
9:DI:214:ILE:O	9:DI:234:LYS:NZ	2.31	0.51
1:DL:115:MET:HE1	10:DL:200:CYC:CMB	2.38	0.51
1:DQ:128:GLN:O	1:DQ:132:GLU:HG2	2.10	0.51
1:AA:22:GLU:O	1:AA:26:ILE:HG12	2.11	0.51
10:AC:200:CYC:HMD1	10:AC:200:CYC:NC	2.24	0.51
3:AE:35:GLN:HG3	3:AE:38:ARG:NH2	2.26	0.51
1:AH:105:GLU:O	1:AH:110:VAL:HG23	2.10	0.51
2:AL:36:LEU:CD1	7:BF:85:GLN:HB2	2.40	0.51
2:AS:51:ILE:HG12	2:AS:140:LEU:HD12	1.93	0.51
2:AS:105:ASP:OD1	2:AS:109:LEU:HD12	2.10	0.51
2:AW:78:TYR:CE1	1:AX:115:MET:HG3	2.46	0.51
1:AX:91:LEU:CD1	1:AX:108:GLY:HA3	2.40	0.51
6:BD:15:GLN:HB3	6:BD:260:ALA:O	2.10	0.51
6:BD:167:ILE:CD1	6:BD:219:ILE:HG22	2.40	0.51
6:BD:303:SER:HA	6:BD:306:GLU:HB2	1.92	0.51
6:BD:774:VAL:O	6:BD:778:ILE:HG12	2.10	0.51

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
6:BD:788:MET:CA	6:BD:792:TYR:HB3	2.32	0.51
9:BH:264:PHE:CE2	9:EA:309:PRO:HG3	2.45	0.51
2:BQ:81:CYS:HA	10:BQ:200:CYC:CHD	2.40	0.51
1:BR:56:ASP:O	1:BR:60:GLN:HG2	2.10	0.51
4:BS:62:GLU:OE1	4:BS:133:ILE:HD11	2.11	0.51
2:CA:115:THR:O	2:CA:119:LEU:HD23	2.10	0.51
6:CM:551:PHE:HE1	6:CM:588:PHE:HB2	1.75	0.51
9:CQ:250:LEU:HD22	9:CQ:286:ILE:CD1	2.41	0.51
1:CR:114:GLU:OE1	1:CR:114:GLU:N	2.43	0.51
1:CV:2:SER:N	1:CV:100:ASP:OD2	2.44	0.51
1:CY:76:GLU:HG2	8:DH:202:ARG:HA	1.92	0.51
1:DA:51:VAL:HA	1:DA:133:MET:CE	2.41	0.51
1:DA:126:VAL:O	1:DA:130:VAL:HG23	2.10	0.51
2:DB:2:GLN:O	2:DB:100:ASP:HB3	2.10	0.51
8:DG:210:LEU:HD22	8:DG:216:MET:CG	2.39	0.51
9:DI:24:THR:HG21	9:DI:142:ILE:CD1	2.40	0.51
9:DI:70:GLU:O	9:DI:74:MET:HG3	2.10	0.51
2:DK:60:LEU:HD22	2:DK:65:VAL:HG11	1.92	0.51
2:DK:96:MET:HA	2:DK:152:TYR:CE2	2.45	0.51
2:DR:3:ASP:N	2:DR:6:THR:OG1	2.43	0.51
2:DR:145:ALA:HA	2:DR:148:GLU:OE1	2.10	0.51
1:DU:47:ARG:O	1:DU:51:VAL:HG22	2.10	0.51
1:DU:71:ASN:HB3	10:DU:200:CYC:OC	2.09	0.51
9:EA:107:LEU:HG	11:EA:400:45D:C30	2.40	0.51
1:AA:27:LYS:O	1:AA:31:THR:HG23	2.09	0.51
2:AF:37:ARG:NH1	2:AF:99:GLY:HA2	2.25	0.51
2:AL:37:ARG:HG3	7:BF:86:VAL:CG2	2.39	0.51
2:AO:92:ALA:HB3	2:AO:149:MET:HE3	1.92	0.51
1:AP:128:GLN:O	1:AP:132:GLU:HG2	2.10	0.51
2:AQ:71:ASN:HB3	10:AQ:200:CYC:OC	2.10	0.51
2:AU:41:ALA:N	2:AU:96:MET:HE1	2.25	0.51
2:AU:47:ASN:O	2:AU:51:ILE:HD12	2.11	0.51
1:AX:147:ALA:O	1:AX:150:SER:OG	2.19	0.51
1:AZ:44:THR:O	1:AZ:47:ARG:HG2	2.10	0.51
6:BD:697:VAL:CG2	6:BD:701:ARG:HE	2.23	0.51
6:BD:860:THR:HG22	6:BD:861:LEU:HD12	1.91	0.51
2:BJ:1:MET:HE2	2:BJ:103:ILE:HD13	1.92	0.51
1:BK:22:GLU:O	1:BK:26:ILE:HG12	2.10	0.51
2:BN:141:VAL:HG12	2:BN:145:ALA:HB3	1.91	0.51
4:BS:92:TYR:HE2	10:BS:200:CYC:HBB1	1.75	0.51
1:CC:48:GLU:O	1:CC:52:LYS:HD3	2.11	0.51

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:CC:122:PRO:O	1:CC:126:VAL:HG23	2.10	0.51
1:CE:123:ILE:HG12	1:CE:160:MET:O	2.09	0.51
1:CE:128:GLN:NE2	1:CE:132:GLU:OE2	2.43	0.51
2:CF:24:MET:HE3	2:CF:28:LYS:CD	2.35	0.51
6:CM:253:LEU:HD11	6:CM:257:TYR:CD2	2.45	0.51
6:CM:532:VAL:O	6:CM:532:VAL:HG12	2.10	0.51
6:CM:760:PHE:HE1	6:CM:784:SER:HB2	1.76	0.51
6:CM:806:LYS:NZ	6:CM:872:LEU:O	2.27	0.51
9:CQ:126:PRO:HD3	11:CQ:400:45D:H211	1.92	0.51
9:CQ:250:LEU:O	9:CQ:251:ILE:HD13	2.11	0.51
1:CR:8:ILE:HD12	2:CS:98:ALA:HB2	1.92	0.51
1:CY:123:ILE:HA	1:CY:126:VAL:HG12	1.90	0.51
2:CZ:112:LEU:HA	2:CZ:115:THR:HG22	1.91	0.51
1:DA:77:MET:CG	8:DG:201:ARG:HB2	2.41	0.51
1:DA:104:ILE:HG22	1:DA:109:LEU:HD12	1.91	0.51
2:DB:133:ILE:O	2:DB:137:THR:HG22	2.11	0.51
2:DD:65:VAL:HG12	2:DD:72:MET:CB	2.38	0.51
8:DH:219:LEU:HD21	8:DY:203:PHE:CZ	2.44	0.51
9:DI:171:ARG:HG3	9:DI:172:ILE:HG12	1.92	0.51
2:DR:10:ASN:O	2:DR:14:VAL:HG23	2.10	0.51
2:DR:66:THR:HA	2:DR:72:MET:HB3	1.92	0.51
9:EA:85:ASP:HB3	9:EA:90:ALA:HB3	1.93	0.51
1:AC:26:ILE:O	1:AC:30:VAL:HG23	2.11	0.51
1:AC:44:THR:O	1:AC:47:ARG:HG2	2.10	0.51
2:AF:141:VAL:HG12	2:AF:145:ALA:HB3	1.92	0.51
2:AI:52:VAL:O	2:AI:56:VAL:HG23	2.10	0.51
2:AI:53:LYS:HG3	1:AJ:118:SER:O	2.11	0.51
2:AI:75:THR:HG23	1:AJ:108:GLY:HA2	1.91	0.51
1:AR:5:THR:O	1:AR:9:VAL:HG23	2.09	0.51
2:AU:112:LEU:HD21	10:AU:200:CYC:NB	2.26	0.51
10:AU:200:CYC:HMA1	10:AU:200:CYC:NB	2.25	0.51
1:AV:95:GLY:HA3	1:AV:104:ILE:HD11	1.91	0.51
9:BH:74:MET:HE2	9:BH:78:GLN:HB2	1.92	0.51
3:BM:11:ALA:HB1	3:BM:16:ARG:O	2.11	0.51
2:BN:101:ALA:CB	2:BN:104:LEU:HD12	2.40	0.51
1:BR:87:TYR:O	1:BR:91:LEU:HG	2.11	0.51
2:BY:57:ALA:HA	2:BY:61:LEU:HD12	1.93	0.51
1:BZ:124:GLU:OE1	1:BZ:124:GLU:N	2.36	0.51
2:CD:119:LEU:O	5:CK:60:PRO:HB3	2.11	0.51
2:CF:44:ILE:HD13	2:CF:149:MET:CE	2.41	0.51
1:CG:85:MET:HB3	1:CG:133:MET:CE	2.38	0.51

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
6:CM:205:ALA:HA	6:CM:208:ASP:OD2	2.11	0.51
6:CM:353:ARG:HG2	6:CM:399:ARG:NH2	2.20	0.51
6:CM:385:GLU:OE2	6:CM:399:ARG:NH2	2.44	0.51
1:CT:131:ARG:HE	1:CT:157:ILE:HG21	1.75	0.51
2:CU:105:ASP:OD2	2:CU:155:TYR:OH	2.27	0.51
1:CV:10:ASN:O	1:CV:14:GLU:HG3	2.10	0.51
1:CV:134:LYS:HD3	1:CV:153:PHE:CB	2.41	0.51
2:DK:2:GLN:HE21	2:DK:7:ALA:HA	1.76	0.51
2:DK:75:THR:HA	1:DL:115:MET:CE	2.40	0.51
2:DK:83:ARG:NH2	2:DK:84:ASP:OD1	2.43	0.51
1:DL:58:LEU:HD22	1:DL:129:SER:CB	2.36	0.51
2:DM:101:ALA:HB2	2:DM:152:TYR:HD1	1.73	0.51
1:DQ:64:ASP:OD1	1:DQ:65:ILE:HD12	2.09	0.51
2:DR:1:MET:SD	2:DR:103:ILE:HD12	2.50	0.51
2:DR:134:LYS:HA	2:DR:153:LEU:HD13	1.92	0.51
2:DV:93:THR:O	2:DV:97:LEU:HD23	2.10	0.51
9:EA:79:GLN:CD	9:EA:122:VAL:HG23	2.35	0.51
9:EA:89:ARG:NE	9:EA:160:ASP:O	2.42	0.51
1:AC:17:TYR:CE1	2:AD:90:ARG:HA	2.45	0.51
2:AD:71:ASN:HD22	2:AD:121:VAL:HA	1.76	0.51
2:AF:81:CYS:HA	10:AF:200:CYC:HHD	1.92	0.51
10:AF:200:CYC:HAB2	5:BA:41:GLN:NE2	2.26	0.51
4:AK:84:ARG:HD3	6:BD:254:PRO:HD3	1.92	0.51
2:AL:85:LEU:HD22	2:AL:133:ILE:CD1	2.37	0.51
1:AN:127:ALA:CB	1:AN:161:SER:HB3	2.40	0.51
2:AO:83:ARG:NH2	6:BD:490:ALA:HB1	2.26	0.51
1:AP:97:VAL:HG21	2:AQ:19:LEU:HD13	1.92	0.51
2:AS:58:LYS:HD2	7:BF:98:VAL:HG12	1.93	0.51
2:AS:65:VAL:HG11	2:AS:72:MET:HE3	1.92	0.51
1:AV:119:LEU:HD12	10:AV:200:CYC:HBD1	1.92	0.51
2:AY:1:MET:CE	2:AY:103:ILE:HB	2.40	0.51
9:BH:71:ILE:HD13	9:BH:74:MET:SD	2.50	0.51
1:BI:116:TYR:HD2	1:BI:123:ILE:HD12	1.73	0.51
1:BK:128:GLN:O	1:BK:131:ARG:HB2	2.10	0.51
1:BP:105:GLU:O	1:BP:110:VAL:HG23	2.11	0.51
1:BP:109:LEU:HD22	1:BP:159:LYS:CG	2.41	0.51
1:BR:27:LYS:O	1:BR:31:THR:HG23	2.09	0.51
2:BY:71:ASN:HD21	2:BY:121:VAL:HG22	1.76	0.51
2:CA:39:ARG:HH12	2:CA:42:SER:HB3	1.75	0.51
1:CG:101:VAL:HB	1:CG:155:PHE:CE2	2.45	0.51
6:CM:269:MET:CE	6:CM:309:VAL:HG12	2.39	0.51

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
6:CM:691:PHE:HB3	1:DU:82:LEU:HB2	1.92	0.51
6:CM:794:PRO:HG2	6:CM:795:TYR:CE2	2.46	0.51
1:CT:30:VAL:HG23	2:CU:31:PHE:HA	1.91	0.51
2:CW:131:GLN:O	2:CW:135:GLU:HG2	2.10	0.51
1:DA:52:LYS:HD3	8:DZ:225:ILE:HD11	1.92	0.51
2:DB:10:ASN:O	2:DB:14:VAL:HG13	2.10	0.51
9:DI:191:GLU:O	9:DI:255:GLY:N	2.38	0.51
9:DI:250:LEU:HD22	9:DI:288:TRP:CZ2	2.43	0.51
1:DL:22:GLU:O	1:DL:26:ILE:HG13	2.10	0.51
1:DQ:2:SER:O	1:DQ:5:THR:HG22	2.11	0.51
2:DR:126:THR:O	2:DR:130:ILE:HG13	2.11	0.51
1:DU:8:ILE:CD1	2:DV:98:ALA:HB2	2.41	0.51
9:EA:264:PHE:CD1	9:EA:294:PRO:HD3	2.46	0.51
2:AD:39:ARG:NH1	2:AD:39:ARG:HA	2.26	0.51
2:AD:124:SER:O	2:AD:128:GLN:HG2	2.10	0.51
2:AF:123:ILE:O	2:AF:127:VAL:HG23	2.11	0.51
1:AJ:18:LEU:HD12	1:AJ:18:LEU:H	1.76	0.51
2:AL:1:MET:HE1	6:BD:25:ILE:CB	2.36	0.51
1:AN:27:LYS:HD3	2:AO:38:VAL:HG11	1.91	0.51
1:AZ:83:ARG:NH2	10:AZ:200:CYC:O2A	2.43	0.51
1:AZ:111:GLY:HA2	1:AZ:114:GLU:OE1	2.11	0.51
1:AZ:142:SER:HB2	2:BN:39:ARG:NE	2.26	0.51
6:BD:261:ALA:CB	6:BD:402:GLY:HA3	2.41	0.51
6:BD:316:MET:SD	6:BD:395:VAL:HG22	2.50	0.51
6:BD:752:GLU:HB2	6:BD:753:PRO:HD3	1.92	0.51
6:BD:779:GLU:O	6:BD:783:THR:HG23	2.10	0.51
9:BH:28:PHE:CD1	9:BH:157:ALA:HB1	2.46	0.51
9:BH:42:PHE:CD2	9:BH:131:LEU:HD13	2.45	0.51
9:BH:217:PHE:HA	9:BH:298:ILE:CG2	2.41	0.51
1:BK:129:SER:HA	1:BK:132:GLU:OE1	2.10	0.51
2:BL:95:ALA:HB2	2:BL:104:LEU:CD2	2.41	0.51
2:BN:56:VAL:O	2:BN:61:LEU:HG	2.09	0.51
2:BN:81:CYS:HA	10:BN:200:CYC:HHD	1.91	0.51
4:BS:41:ALA:CB	4:BS:98:ILE:HD11	2.41	0.51
4:BS:105:LEU:HD11	4:BS:160:MET:HG3	1.91	0.51
1:BV:27:LYS:HE3	2:BW:35:GLU:OE1	2.11	0.51
2:CF:3:ASP:OD1	2:CF:6:THR:OG1	2.12	0.51
2:CF:71:ASN:ND2	2:CF:121:VAL:HA	2.26	0.51
1:CG:32:GLY:O	1:CG:36:ARG:HG3	2.11	0.51
2:CH:131:GLN:O	2:CH:134:LYS:HB3	2.11	0.51
6:CM:830:GLY:HA2	1:DQ:15:ALA:HB2	1.93	0.51

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
9:CQ:99:ALA:HB1	9:CQ:164:THR:HG22	1.92	0.51
2:CS:5:ILE:HD12	2:CS:27:LEU:HD22	1.91	0.51
2:CW:54:GLU:HG3	9:DI:63:LEU:HD21	1.92	0.51
2:CZ:3:ASP:HB2	2:CZ:98:ALA:O	2.11	0.51
1:DA:66:VAL:O	1:DA:73:TYR:HA	2.11	0.51
2:DB:105:ASP:HA	2:DB:109:LEU:HB2	1.92	0.51
8:DG:210:LEU:HD22	8:DG:216:MET:CB	2.40	0.51
9:DI:125:ILE:HB	9:DI:126:PRO:HD2	1.93	0.51
1:DJ:76:GLU:OE1	9:EA:152:THR:OG1	2.25	0.51
1:DL:50:ILE:HG13	1:DL:136:VAL:HG12	1.93	0.51
10:DL:200:CYC:HBB3	10:DL:200:CYC:HMB1	1.92	0.51
2:DM:41:ALA:CB	2:DM:97:LEU:HD12	2.40	0.51
1:DQ:153:PHE:O	1:DQ:157:ILE:HG23	2.11	0.51
2:DR:43:VAL:CG1	2:DR:141:VAL:HG12	2.41	0.51
2:DR:44:ILE:CD1	2:DR:149:MET:HE3	2.41	0.51
2:DT:71:ASN:ND2	2:DT:121:VAL:HA	2.26	0.51
2:DT:126:THR:O	2:DT:130:ILE:HG13	2.11	0.51
2:DV:111:GLY:HA2	2:DV:114:GLU:CD	2.36	0.51
8:DZ:210:LEU:CD1	8:DZ:216:MET:HB2	2.41	0.51
2:AB:103:ILE:HD11	2:AB:107:ARG:HD2	1.91	0.51
2:AD:37:ARG:NH2	2:AD:99:GLY:HA2	2.25	0.51
2:AO:20:ASP:OD1	2:AO:20:ASP:N	2.42	0.51
1:AP:97:VAL:HG21	2:AQ:19:LEU:CD1	2.41	0.51
2:AS:63:SER:H	2:AS:66:THR:CG2	2.24	0.51
6:BD:144:TYR:OH	10:BD:902:CYC:HAC1	2.10	0.51
9:BH:208:ASN:OD1	9:BH:247:ASN:HA	2.11	0.51
2:BJ:13:ASP:OD2	5:CJ:63:ASN:ND2	2.36	0.51
2:BL:137:THR:O	2:BL:141:VAL:HG22	2.11	0.51
1:BP:68:PRO:HA	1:BP:73:TYR:CE2	2.46	0.51
1:BP:115:MET:HG3	2:BT:78:TYR:CD1	2.45	0.51
2:BQ:107:ARG:NH1	6:CM:334:PHE:O	2.43	0.51
2:BT:18:TYR:OH	6:CM:160:LEU:HD23	2.10	0.51
2:BT:41:ALA:CB	2:BT:97:LEU:HD11	2.41	0.51
1:BZ:24:ASP:HA	1:BZ:27:LYS:HG2	1.93	0.51
2:CA:96:MET:HA	2:CA:152:TYR:CE2	2.45	0.51
1:CC:58:LEU:HD22	1:CC:129:SER:HB3	1.92	0.51
2:CD:54:GLU:OE1	2:CD:136:VAL:HG21	2.10	0.51
2:CH:96:MET:HA	2:CH:152:TYR:CE2	2.46	0.51
9:CQ:172:ILE:H	9:CQ:172:ILE:HD12	1.76	0.51
1:CR:72:ALA:HA	1:CR:77:MET:HB3	1.92	0.51
2:CU:112:LEU:HD11	2:CU:116:TYR:CZ	2.45	0.51

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:CW:73:TYR:O	2:CW:77:ARG:HB2	2.10	0.51
2:CZ:112:LEU:HD22	2:CZ:160:LEU:HD13	1.93	0.51
1:DA:66:VAL:HG21	8:DZ:227:PRO:CG	2.40	0.51
2:DB:156:ILE:O	2:DB:160:LEU:HD23	2.10	0.51
5:DF:27:PHE:CE2	5:DF:29:LYS:HE3	2.46	0.51
9:DI:291:LEU:HD23	9:DI:299:PHE:HD2	1.75	0.51
2:DK:43:VAL:HB	2:DK:141:VAL:HG12	1.92	0.51
2:DM:2:GLN:O	2:DM:100:ASP:HB3	2.11	0.51
2:DM:65:VAL:HG12	2:DM:72:MET:CB	2.36	0.51
2:DM:104:LEU:HD13	2:DM:156:ILE:CG1	2.36	0.51
2:DR:15:GLN:HB2	2:DR:17:LYS:HG2	1.93	0.51
1:DS:50:ILE:HG13	1:DS:136:VAL:HG13	1.93	0.51
1:DS:113:ARG:NH2	1:DS:161:SER:O	2.33	0.51
9:EA:102:SER:OG	9:EA:105:ILE:HG12	2.11	0.51
1:AA:85:MET:HB3	1:AA:133:MET:CE	2.39	0.51
2:AD:75:THR:HG22	3:AE:107:THR:O	2.11	0.51
10:AD:200:CYC:HMA3	10:AD:200:CYC:O1A	2.10	0.51
2:AI:77:ARG:CG	10:AI:200:CYC:HAD1	2.40	0.51
4:AK:41:ALA:HA	4:AK:44:ILE:HD11	1.93	0.51
4:AK:50:THR:HB	4:AK:54:ARG:HH12	1.76	0.51
2:AL:5:ILE:O	2:AL:9:ILE:HG13	2.11	0.51
2:AO:3:ASP:HA	2:AO:98:ALA:HB1	1.93	0.51
1:AV:96:VAL:HG12	1:AV:152:TYR:CD2	2.46	0.51
5:BA:29:LYS:NZ	6:BD:315:SER:OG	2.22	0.51
6:BD:788:MET:HB2	6:BD:792:TYR:HD2	1.76	0.51
9:BH:36:GLN:O	9:BH:40:ILE:HG23	2.10	0.51
1:BI:3:ILE:CG2	1:BP:25:ARG:HD3	2.41	0.51
3:BM:40:ALA:CB	3:BM:97:LEU:HD11	2.40	0.51
2:BN:144:ASP:OD1	2:BN:144:ASP:N	2.43	0.51
1:BP:35:ALA:O	1:BP:39:ILE:HG13	2.10	0.51
10:BQ:200:CYC:HBB2	6:CM:338:ILE:HD11	1.93	0.51
1:BR:18:LEU:HD13	4:BS:98:ILE:CD1	2.40	0.51
1:BV:53:GLN:O	1:BV:57:ARG:HD3	2.10	0.51
1:CC:18:LEU:HD12	1:CC:22:GLU:HB3	1.93	0.51
1:CE:81:CYS:HA	10:CE:200:CYC:HHD	1.93	0.51
1:CE:97:VAL:HG11	2:CF:19:LEU:CD1	2.39	0.51
6:CM:316:MET:CE	6:CM:395:VAL:HG22	2.37	0.51
1:CT:81:CYS:SG	10:CT:200:CYC:HMD3	2.51	0.51
1:CV:91:LEU:O	1:CV:104:ILE:HG12	2.10	0.51
1:DA:130:VAL:HB	1:DA:157:ILE:CD1	2.41	0.51
8:DG:230:ASN:ND2	1:DL:68:PRO:HD2	2.26	0.51

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
9:DI:263:GLY:C	9:DI:294:PRO:HB3	2.36	0.51
2:DO:122:PRO:HG2	10:DO:200:CYC:HMC3	1.92	0.51
1:DQ:62:ARG:NH2	1:DQ:64:ASP:OD2	2.44	0.51
5:DX:54:GLU:N	5:DX:54:GLU:OE1	2.44	0.51
10:AB:200:CYC:HMA1	6:BD:311:ASN:HB3	1.93	0.51
1:AC:8:ILE:HD12	2:AD:1:MET:HE1	1.92	0.51
1:AH:89:LEU:HB2	1:AH:133:MET:HE1	1.93	0.51
1:AH:141:MET:HE1	1:AH:149:ALA:CB	2.41	0.51
2:AI:88:TYR:OH	2:AI:112:LEU:HD21	2.10	0.51
2:AL:97:LEU:CD1	6:BD:35:LEU:HD12	2.41	0.51
1:AN:8:ILE:HD12	2:AO:94:TYR:CD1	2.46	0.51
2:AW:15:GLN:HB2	2:AW:17:LYS:NZ	2.26	0.51
6:BD:811:ARG:NH2	6:BD:814:LEU:HG	2.26	0.51
9:BH:20:VAL:HG22	9:BH:145:LEU:HD21	1.92	0.51
9:BH:21:VAL:HB	9:BH:22:PRO:HD3	1.92	0.51
9:BH:42:PHE:CE2	9:BH:131:LEU:HB2	2.46	0.51
1:BI:114:GLU:OE1	1:BI:114:GLU:N	2.41	0.51
1:BK:126:VAL:CG1	10:BK:200:CYC:HMC2	2.40	0.51
2:BL:72:MET:HE3	2:BL:78:TYR:CE1	2.45	0.51
2:BN:96:MET:HG2	2:BN:148:GLU:OE1	2.11	0.51
1:BP:114:GLU:OE2	6:CM:299:SER:OG	2.16	0.51
1:BP:126:VAL:O	1:BP:130:VAL:HG23	2.11	0.51
1:BR:8:ILE:CD1	4:BS:95:TYR:HB3	2.30	0.51
1:BR:126:VAL:HG22	10:BR:200:CYC:HMC1	1.93	0.51
1:BX:89:LEU:HB2	1:BX:133:MET:CE	2.37	0.51
2:BY:62:TYR:CE2	9:CQ:11:ILE:HA	2.45	0.51
1:CE:98:SER:HB3	2:CF:9:ILE:CD1	2.41	0.51
1:CG:59:PHE:CZ	1:CG:78:THR:HG23	2.46	0.51
1:CG:85:MET:SD	1:CG:129:SER:HB2	2.51	0.51
9:CQ:90:ALA:O	9:CQ:163:PHE:HB2	2.10	0.51
9:CQ:195:ASN:OD1	9:CQ:198:VAL:HB	2.11	0.51
1:CV:134:LYS:HD3	1:CV:153:PHE:HB3	1.93	0.51
2:CZ:113:LYS:HG2	2:CZ:160:LEU:O	2.11	0.51
9:DI:41:TRP:CZ2	9:DI:45:LEU:HD21	2.46	0.51
1:DL:63:PRO:O	1:DL:67:SER:OG	2.23	0.51
2:DO:78:TYR:O	2:DO:82:ILE:HG12	2.11	0.51
9:EA:126:PRO:HD3	11:EA:400:45D:H211	1.93	0.51
3:AE:58:LEU:HD22	3:AE:129:ALA:CB	2.36	0.51
1:AH:27:LYS:HG3	2:AI:35:GLU:OE1	2.11	0.51
1:AX:122:PRO:O	1:AX:126:VAL:HG23	2.11	0.51
1:AZ:71:ASN:HD22	1:AZ:121:THR:HA	1.76	0.51

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
6:BD:137:ARG:HD3	6:BD:138:PRO:O	2.10	0.51
1:BI:89:LEU:O	1:BI:93:THR:HG23	2.11	0.51
2:BL:79:ALA:HA	2:BL:82:ILE:HG12	1.91	0.51
1:BR:18:LEU:CD1	4:BS:98:ILE:HD12	2.41	0.51
1:BR:88:TYR:O	1:BR:92:VAL:HG23	2.11	0.51
1:BX:128:GLN:O	1:BX:132:GLU:HG2	2.11	0.51
1:BZ:71:ASN:HD21	10:BZ:200:CYC:HBD2	1.76	0.51
6:CM:268:ALA:HB3	6:CM:270:LYS:NZ	2.26	0.51
6:CM:413:MET:HE1	6:CM:434:PHE:CE1	2.46	0.51
6:CM:447:TYR:CE1	10:CM:901:CYC:HAA1	2.46	0.51
9:CQ:21:VAL:HB	9:CQ:22:PRO:HD3	1.93	0.51
9:CQ:274:GLN:HA	9:CQ:282:VAL:HA	1.92	0.51
9:CQ:284:MET:SD	9:CQ:307:ALA:HA	2.51	0.51
2:CS:28:LYS:HE2	2:CS:28:LYS:HA	1.93	0.51
2:CS:109:LEU:CD1	2:CS:159:GLY:HA3	2.41	0.51
1:CY:85:MET:HG2	10:CY:200:CYC:HBC1	1.93	0.51
1:DA:47:ARG:O	1:DA:51:VAL:HG23	2.11	0.51
9:DI:266:GLN:HA	9:DI:290:PHE:O	2.10	0.51
1:DJ:104:ILE:HG22	1:DJ:109:LEU:HD23	1.93	0.51
2:DK:20:ASP:O	2:DK:24:MET:HG2	2.11	0.51
2:DK:51:ILE:HG23	2:DK:136:VAL:CB	2.34	0.51
2:DT:52:VAL:HA	2:DT:133:ILE:HD11	1.93	0.51
1:AA:23:LEU:HD13	2:AB:42:SER:OG	2.11	0.50
1:AA:98:SER:HA	2:AB:5:ILE:HG21	1.92	0.50
2:AD:137:THR:O	2:AD:141:VAL:HG22	2.11	0.50
2:AF:63:SER:H	2:AF:66:THR:CG2	2.24	0.50
6:BD:597:TYR:CB	6:BD:600:LYS:HD2	2.39	0.50
7:BF:111:LYS:O	7:BF:115:ARG:HG3	2.11	0.50
9:BH:248:LEU:HD13	9:BH:250:LEU:HD21	1.93	0.50
10:BL:200:CYC:O1A	5:CJ:12:SER:N	2.31	0.50
2:BN:112:LEU:HD22	10:BN:200:CYC:HMB1	1.94	0.50
1:BP:44:THR:O	1:BP:47:ARG:HG2	2.11	0.50
1:BP:156:VAL:HG12	1:BP:160:MET:HE3	1.93	0.50
1:BR:89:LEU:O	1:BR:93:THR:HG23	2.11	0.50
4:BS:72:ASN:ND2	4:BS:122:VAL:HA	2.24	0.50
2:BY:52:VAL:HG21	2:BY:86:ASP:OD1	2.11	0.50
1:BZ:82:LEU:HD13	8:CP:221:MET:SD	2.51	0.50
2:CF:3:ASP:CB	2:CF:98:ALA:HB1	2.41	0.50
6:CM:210:PHE:C	6:CM:211:ARG:HD3	2.36	0.50
6:CM:799:LYS:HB2	6:CM:883:LEU:HD21	1.92	0.50
2:CS:61:LEU:HD22	10:CT:200:CYC:HBA1	1.93	0.50

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:CT:115:MET:HE3	1:CT:116:TYR:CE1	2.46	0.50
1:CT:137:ALA:O	1:CT:141:MET:HG2	2.11	0.50
2:CZ:129:ALA:O	2:CZ:133:ILE:HG13	2.11	0.50
1:DL:68:PRO:HA	1:DL:73:TYR:CG	2.45	0.50
2:DM:36:LEU:HD13	2:DM:144:ASP:OD2	2.11	0.50
1:DN:41:GLU:CG	2:DO:24:MET:HE2	2.37	0.50
2:DR:50:THR:HG22	2:DR:54:GLU:OE1	2.10	0.50
1:AN:11:ALA:HB2	1:AN:18:LEU:HD23	1.93	0.50
1:AP:53:GLN:O	1:AP:57:ARG:HG3	2.10	0.50
2:AS:20:ASP:O	2:AS:24:MET:HG2	2.11	0.50
1:AV:105:GLU:HA	1:AV:109:LEU:HB2	1.94	0.50
1:AV:122:PRO:HG2	1:AV:125:ALA:CB	2.41	0.50
2:AW:56:VAL:O	2:AW:61:LEU:HG	2.11	0.50
2:AW:127:VAL:HG22	2:AW:157:CYS:SG	2.51	0.50
10:AY:200:CYC:HMA1	10:AY:200:CYC:HB	1.75	0.50
1:AZ:101:VAL:HG12	1:AZ:152:TYR:HD1	1.77	0.50
6:BD:20:VAL:HB	6:BD:21:PRO:HD3	1.93	0.50
6:BD:33:ARG:NH1	6:BD:36:GLU:HG2	2.26	0.50
6:BD:401:LEU:HD13	6:BD:411:TRP:CH2	2.46	0.50
6:BD:526:LYS:HD3	6:BD:533:LYS:HE2	1.93	0.50
6:BD:574:SER:HB2	6:BD:651:GLU:O	2.11	0.50
1:BI:23:LEU:HD13	2:BJ:42:SER:OG	2.11	0.50
1:BI:152:TYR:O	1:BI:156:VAL:HG23	2.11	0.50
1:BK:76:GLU:O	1:BK:80:THR:HG23	2.11	0.50
1:BK:105:GLU:HG2	1:BK:109:LEU:CD2	2.41	0.50
1:BR:92:VAL:O	1:BR:96:VAL:HG23	2.11	0.50
1:BX:18:LEU:HD22	2:BY:97:LEU:HD13	1.93	0.50
1:BZ:39:ILE:HD11	1:BZ:148:GLU:OE2	2.10	0.50
1:CG:90:ARG:NH1	2:CH:16:GLY:HA2	2.26	0.50
6:CM:515:LEU:HD11	6:CM:568:LEU:HD11	1.94	0.50
6:CM:743:TYR:OH	6:CM:760:PHE:HB2	2.10	0.50
6:CM:881:LYS:NZ	2:DR:106:GLU:OE2	2.43	0.50
9:CQ:229:ARG:HG2	9:DI:227:PHE:CB	2.41	0.50
1:CR:82:LEU:HB2	8:DZ:242:ILE:HG12	1.93	0.50
1:CR:85:MET:HE3	1:CR:129:SER:OG	2.12	0.50
2:CS:41:ALA:N	2:CS:96:MET:HE1	2.26	0.50
2:CS:61:LEU:HD22	10:CT:200:CYC:CBA	2.41	0.50
1:CV:128:GLN:NE2	1:CV:132:GLU:OE2	2.39	0.50
2:CW:43:VAL:O	2:CW:46:ALA:HB3	2.12	0.50
2:CW:119:LEU:HD21	5:DF:33:TYR:HE2	1.76	0.50
2:CZ:41:ALA:N	2:CZ:96:MET:HE1	2.25	0.50

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:DA:103:PRO:O	1:DA:107:ILE:HG13	2.11	0.50
2:DB:64:ASP:OD1	2:DB:64:ASP:N	2.45	0.50
2:DB:111:GLY:HA2	2:DB:114:GLU:CD	2.36	0.50
8:DH:203:PHE:CZ	8:DY:219:LEU:HD13	2.46	0.50
1:DL:89:LEU:HD13	1:DL:133:MET:HE2	1.94	0.50
1:DN:37:LEU:HD23	2:DO:28:LYS:HE3	1.93	0.50
1:DQ:77:MET:HE2	8:DZ:201:ARG:HB2	1.92	0.50
1:DQ:127:ALA:CA	1:DQ:160:MET:HE1	2.41	0.50
10:DQ:200:CYC:HBA2	2:DV:61:LEU:HD22	1.93	0.50
2:DR:94:TYR:HB3	2:DR:103:ILE:HD13	1.93	0.50
1:DS:41:GLU:HG3	2:DT:24:MET:CE	2.35	0.50
2:DV:55:ALA:HB2	2:DV:133:ILE:HD13	1.93	0.50
5:DX:3:MET:HG2	5:DX:32:PRO:HA	1.92	0.50
2:AB:95:ALA:HB2	2:AB:104:LEU:HG	1.94	0.50
2:AD:65:VAL:C	2:AD:72:MET:HB3	2.36	0.50
2:AF:44:ILE:HG21	2:AF:93:THR:OG1	2.12	0.50
4:AK:40:ALA:O	4:AK:44:ILE:HG12	2.10	0.50
2:AL:3:ASP:H	2:AL:6:THR:HG22	1.76	0.50
2:AS:90:ARG:NH1	2:AS:94:TYR:OH	2.37	0.50
2:AS:119:LEU:CB	2:AS:121:VAL:HG23	2.41	0.50
1:AZ:144:ASP:H	2:BN:39:ARG:NH2	2.09	0.50
5:BA:5:ARG:HB2	5:BA:56:ALA:HB2	1.93	0.50
9:BH:216:LEU:O	9:BH:298:ILE:HG22	2.11	0.50
9:BH:280:GLY:HA3	9:BH:284:MET:CG	2.41	0.50
1:BI:35:ALA:HA	1:BI:38:ARG:NH1	2.25	0.50
1:BI:156:VAL:CG1	1:BI:160:MET:HE3	2.41	0.50
1:BK:132:GLU:HA	1:BK:135:GLU:OE1	2.11	0.50
3:BM:35:GLN:O	3:BM:39:ILE:HG12	2.11	0.50
1:BV:92:VAL:HG11	1:BV:153:PHE:CE2	2.46	0.50
2:CA:126:THR:HA	10:CM:901:CYC:HMC1	1.93	0.50
1:CC:23:LEU:CD2	2:CD:38:VAL:HG13	2.42	0.50
1:CC:109:LEU:CD1	1:CC:160:MET:HE2	2.41	0.50
1:CE:98:SER:HB3	2:CF:9:ILE:HD11	1.94	0.50
2:CF:78:TYR:CD1	1:CG:115:MET:HG3	2.46	0.50
1:CG:85:MET:HE3	1:CG:133:MET:HE3	1.93	0.50
1:CT:5:THR:HG23	2:CU:1:MET:CE	2.41	0.50
1:CT:153:PHE:O	1:CT:157:ILE:HG12	2.10	0.50
2:CW:36:LEU:HD21	2:CW:145:ALA:HB2	1.91	0.50
2:CW:114:GLU:CG	5:DF:53:VAL:HG12	2.41	0.50
1:CY:49:THR:O	1:CY:53:GLN:HG3	2.11	0.50
1:CY:79:ALA:HB3	8:DH:203:PHE:CE1	2.47	0.50

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:CZ:47:ASN:ND2	2:CZ:140:LEU:HD13	2.27	0.50
1:DA:5:THR:HG23	2:DB:1:MET:SD	2.52	0.50
2:DD:148:GLU:O	2:DD:151:VAL:HG12	2.11	0.50
5:DF:59:ARG:O	5:DF:62:THR:HG23	2.12	0.50
9:DI:40:ILE:HG13	11:DI:400:45D:H371	1.92	0.50
1:DJ:39:ILE:HG12	1:DJ:145:ASP:CG	2.37	0.50
1:DJ:126:VAL:CG2	10:DJ:200:CYC:HMC3	2.41	0.50
1:DL:85:MET:HE3	1:DL:129:SER:HB2	1.92	0.50
1:DS:18:LEU:HD22	2:DT:97:LEU:CD2	2.41	0.50
1:DS:63:PRO:HD2	9:EA:55:ALA:HB1	1.93	0.50
2:DV:123:ILE:O	2:DV:126:THR:OG1	2.24	0.50
9:EA:248:LEU:O	9:EA:249:LYS:HE2	2.10	0.50
1:AA:90:ARG:HD3	2:AB:18:TYR:CD1	2.46	0.50
2:AD:61:LEU:HB3	2:AD:62:TYR:CD1	2.47	0.50
2:AF:119:LEU:HG	5:BA:4:PHE:HZ	1.75	0.50
1:AH:110:VAL:HG11	2:AL:76:ARG:HG3	1.92	0.50
2:AI:63:SER:HB3	6:BD:682:ARG:HB2	1.93	0.50
2:AL:97:LEU:HD13	6:BD:43:LEU:CD1	2.41	0.50
1:AN:83:ARG:HA	7:BF:110:PHE:CZ	2.47	0.50
2:AY:85:LEU:HG	10:AY:200:CYC:HBC1	1.93	0.50
2:AY:96:MET:HG3	2:AY:148:GLU:OE1	2.12	0.50
1:AZ:6:LYS:NZ	1:AZ:102:THR:OG1	2.44	0.50
1:AZ:115:MET:HE1	10:AZ:200:CYC:CMB	2.37	0.50
6:BD:452:ASP:OD1	6:BD:607:ARG:NH2	2.45	0.50
9:BH:248:LEU:HA	9:BH:275:THR:CG2	2.37	0.50
1:BK:20:PRO:HD2	6:CM:171:ASP:OD1	2.12	0.50
1:BR:84:ASP:OD2	10:BR:200:CYC:ND	2.42	0.50
2:BY:50:THR:O	2:BY:54:GLU:HG2	2.11	0.50
2:CA:65:VAL:CG1	2:CA:72:MET:HE3	2.41	0.50
1:CC:7:SER:HB2	1:CC:18:LEU:CD1	2.41	0.50
2:CD:132:ALA:HA	2:CD:135:GLU:OE1	2.12	0.50
1:CG:67:SER:O	1:CG:73:TYR:HB2	2.11	0.50
6:CM:86:GLU:OE1	6:CM:154:ARG:HB2	2.11	0.50
6:CM:156:MET:HB3	6:CM:202:MET:CE	2.41	0.50
6:CM:238:ASN:HB2	6:CM:252:GLU:OE2	2.11	0.50
6:CM:805:THR:HA	6:CM:838:MET:CE	2.41	0.50
7:CO:114:ALA:HA	7:CO:117:VAL:CG1	2.42	0.50
1:CR:87:TYR:HA	1:CR:90:ARG:NH1	2.26	0.50
1:CR:105:GLU:HG3	1:CR:155:PHE:HZ	1.77	0.50
1:CT:91:LEU:HD21	1:CT:107:ILE:CG2	2.42	0.50
1:CV:131:ARG:O	1:CV:134:LYS:HB3	2.12	0.50

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:CZ:2:GLN:O	2:CZ:100:ASP:HB3	2.12	0.50
2:DB:36:LEU:HD21	2:DB:145:ALA:HA	1.93	0.50
1:DC:71:ASN:HD21	10:DC:200:CYC:HBD2	1.76	0.50
1:DC:115:MET:HE3	1:DC:116:TYR:CE1	2.47	0.50
9:DI:148:GLY:O	9:DI:151:ILE:HG22	2.11	0.50
2:DM:64:ASP:OD1	2:DM:64:ASP:N	2.45	0.50
1:DQ:101:VAL:HG21	1:DQ:155:PHE:CZ	2.46	0.50
1:DS:130:VAL:HA	1:DS:133:MET:CE	2.41	0.50
2:DT:71:ASN:HB3	10:DT:200:CYC:OC	2.10	0.50
9:EA:63:LEU:HD11	9:EA:105:ILE:HG21	1.92	0.50
9:EA:223:LEU:HB3	9:EA:231:ILE:CG2	2.41	0.50
1:AC:132:GLU:HA	1:AC:135:GLU:OE1	2.12	0.50
2:AD:105:ASP:OD1	2:AD:109:LEU:HD22	2.12	0.50
2:AD:107:ARG:CD	6:BD:270:LYS:HE3	2.40	0.50
4:AK:68:ARG:HD2	7:BF:105:PRO:HG2	1.94	0.50
2:AL:81:CYS:HA	10:AL:200:CYC:HBC3	1.92	0.50
1:AN:87:TYR:HB3	10:AN:200:CYC:HBB3	1.94	0.50
1:AN:90:ARG:NH2	2:AS:73:TYR:OH	2.45	0.50
1:AN:102:THR:HA	1:AN:105:GLU:CG	2.40	0.50
2:AO:72:MET:HE2	2:AO:78:TYR:CE2	2.47	0.50
2:AO:134:LYS:CA	2:AO:153:LEU:HD13	2.42	0.50
2:AW:44:ILE:HG21	2:AW:93:THR:OG1	2.12	0.50
1:AX:27:LYS:O	1:AX:31:THR:HG23	2.12	0.50
1:BK:131:ARG:HE	1:BK:157:ILE:HG21	1.75	0.50
2:BL:37:ARG:NH2	2:BL:99:GLY:HA2	2.26	0.50
10:BQ:200:CYC:OB	6:CM:339:ASN:HB2	2.11	0.50
4:BS:51:ILE:HD13	4:BS:138:ILE:CD1	2.41	0.50
2:BY:100:ASP:OD1	2:BY:102:SER:N	2.44	0.50
1:BZ:126:VAL:HG22	10:BZ:200:CYC:HMC1	1.94	0.50
2:CF:73:TYR:OH	1:CG:90:ARG:NH2	2.43	0.50
2:CH:78:TYR:O	2:CH:82:ILE:HG12	2.11	0.50
6:CM:152:SER:OG	10:CM:902:CYC:H3C	2.12	0.50
6:CM:691:PHE:CB	1:DU:82:LEU:HB2	2.41	0.50
9:CQ:290:PHE:CD2	9:CQ:298:ILE:HD13	2.47	0.50
1:CR:132:GLU:HA	1:CR:135:GLU:OE1	2.11	0.50
2:CU:50:THR:O	2:CU:53:LYS:HB3	2.12	0.50
2:CU:103:ILE:CD1	2:CU:107:ARG:HG3	2.42	0.50
1:CV:125:ALA:CB	10:CV:200:CYC:HMC2	2.41	0.50
1:CY:134:LYS:HB2	1:CY:153:PHE:HB3	1.93	0.50
2:CZ:87:TYR:CG	10:CZ:200:CYC:HBB3	2.46	0.50
1:DA:33:GLY:HA3	2:DB:31:PHE:CE2	2.47	0.50

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
9:DI:28:PHE:CD2	9:DI:39:LEU:HD23	2.47	0.50
1:DL:68:PRO:HG3	1:DL:73:TYR:CE1	2.46	0.50
1:DL:141:MET:HB2	1:DL:146:ALA:HB2	1.92	0.50
2:DM:10:ASN:O	2:DM:14:VAL:HG13	2.10	0.50
2:DM:126:THR:O	2:DM:130:ILE:HG12	2.12	0.50
1:DN:83:ARG:NH1	10:DN:200:CYC:O2A	2.40	0.50
1:DQ:27:LYS:HD3	2:DR:38:VAL:HG11	1.94	0.50
1:DQ:33:GLY:O	1:DQ:37:LEU:HD23	2.12	0.50
2:DR:8:VAL:CG1	2:DR:23:ALA:HB1	2.40	0.50
2:AD:3:ASP:HB3	2:AD:98:ALA:O	2.12	0.50
2:AD:50:THR:O	2:AD:54:GLU:HG2	2.10	0.50
1:AJ:29:PHE:HA	1:AJ:36:ARG:HH21	1.76	0.50
1:AN:26:ILE:O	1:AN:30:VAL:HG13	2.11	0.50
1:AN:87:TYR:CD2	10:AN:200:CYC:HBB3	2.47	0.50
1:AP:142:SER:HB2	2:DT:39:ARG:NH1	2.27	0.50
2:AU:104:LEU:HD13	2:AU:156:ILE:HG13	1.94	0.50
1:AV:50:ILE:HD13	1:AV:136:VAL:HG12	1.93	0.50
1:AV:132:GLU:O	1:AV:136:VAL:HG23	2.12	0.50
5:BA:9:CYS:HA	5:BA:25:THR:O	2.11	0.50
6:BD:826:LEU:HD23	6:BD:830:GLY:O	2.11	0.50
6:BD:891:VAL:HG21	2:DB:105:ASP:OD1	2.12	0.50
9:BH:193:VAL:CG2	9:BH:255:GLY:HA3	2.42	0.50
9:BH:273:VAL:O	9:BH:282:VAL:HA	2.12	0.50
1:BI:115:MET:HE1	10:BI:200:CYC:CMB	2.41	0.50
1:BK:87:TYR:O	1:BK:91:LEU:HG	2.12	0.50
3:BM:92:VAL:HG11	3:BM:153:PHE:CE2	2.47	0.50
4:BS:116:THR:HG21	10:BS:200:CYC:CMA	2.42	0.50
2:BT:2:GLN:HB2	2:BT:6:THR:HG21	1.93	0.50
2:BT:8:VAL:CG1	2:BT:23:ALA:HB1	2.41	0.50
2:BT:42:SER:OG	6:CM:40:LEU:HD13	2.12	0.50
1:BZ:87:TYR:O	1:BZ:91:LEU:HG	2.11	0.50
1:CC:13:ALA:HA	5:CK:45:LYS:NZ	2.26	0.50
1:CC:156:VAL:O	1:CC:160:MET:HG2	2.12	0.50
2:CD:147:LYS:O	2:CD:151:VAL:HG23	2.12	0.50
1:CE:27:LYS:O	1:CE:31:THR:HG23	2.11	0.50
1:CE:30:VAL:HA	2:CF:31:PHE:HD1	1.76	0.50
2:CF:126:THR:O	2:CF:130:ILE:HG13	2.11	0.50
2:CH:72:MET:HG3	2:CH:78:TYR:HA	1.94	0.50
2:CH:137:THR:O	2:CH:141:VAL:HG22	2.12	0.50
6:CM:277:GLU:O	6:CM:281:VAL:HG23	2.12	0.50
6:CM:612:ARG:NH2	6:CM:644:GLU:OE1	2.44	0.50

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
9:CQ:280:GLY:HA3	9:CQ:284:MET:CG	2.41	0.50
2:CS:5:ILE:O	2:CS:9:ILE:HG12	2.12	0.50
2:CS:47:ASN:HB2	2:CS:51:ILE:HD11	1.92	0.50
1:CY:18:LEU:HD12	1:CY:22:GLU:CB	2.41	0.50
1:CY:53:GLN:HB3	1:CY:57:ARG:HH22	1.76	0.50
5:DF:27:PHE:HE2	5:DF:29:LYS:HE3	1.75	0.50
1:DL:2:SER:OG	2:DM:3:ASP:OD1	2.26	0.50
1:DN:50:ILE:HG12	1:DN:140:LEU:HD21	1.93	0.50
2:DR:3:ASP:OD1	2:DR:6:THR:OG1	2.30	0.50
1:DS:62:ARG:O	1:DS:65:ILE:HG12	2.11	0.50
1:DS:76:GLU:HB2	8:DY:203:PHE:HD2	1.77	0.50
2:DV:64:ASP:O	2:DV:67:ARG:HG2	2.11	0.50
9:EA:25:ILE:HD12	9:EA:156:ASN:HB2	1.92	0.50
9:EA:214:ILE:HD13	9:EA:238:LEU:HB2	1.93	0.50
3:AE:47:GLU:HG2	3:AE:89:LEU:CD2	2.42	0.50
10:AH:200:CYC:CAA	2:AL:61:LEU:HD22	2.42	0.50
2:AL:106:GLU:CG	2:AL:107:ARG:HG2	2.41	0.50
2:AQ:110:ASN:ND2	6:BD:668:ARG:HG2	2.22	0.50
2:AY:96:MET:HG3	2:AY:148:GLU:OE2	2.12	0.50
5:BA:7:THR:HB	5:BA:52:LYS:HB3	1.93	0.50
5:BB:30:LEU:N	6:BD:572:ASP:O	2.42	0.50
6:BD:782:GLY:HA3	6:BD:835:ILE:HG21	1.94	0.50
6:BD:788:MET:O	6:BD:793:ALA:N	2.36	0.50
6:BD:842:MET:O	6:BD:846:GLN:HG2	2.10	0.50
9:BH:76:PRO:CB	9:BH:123:ALA:HB2	2.40	0.50
1:BK:5:THR:HG22	2:BL:3:ASP:OD2	2.11	0.50
1:BK:39:ILE:HG23	1:BK:145:ASP:OD1	2.12	0.50
2:BL:57:ALA:HA	2:BL:61:LEU:CD1	2.42	0.50
10:BN:200:CYC:CGA	5:CJ:37:PHE:HB2	2.41	0.50
2:BQ:53:LYS:HG3	1:BR:118:SER:O	2.12	0.50
2:BQ:65:VAL:HG12	2:BQ:72:MET:HB2	1.93	0.50
1:BV:92:VAL:HG21	1:BV:153:PHE:CE1	2.46	0.50
1:CE:64:ASP:HA	1:CE:67:SER:HB2	1.94	0.50
1:CG:61:LYS:HG2	9:CQ:57:GLY:CA	2.42	0.50
6:CM:689:PRO:HB2	6:CM:692:VAL:HG23	1.93	0.50
9:CQ:298:ILE:HD11	9:CQ:300:PHE:O	2.12	0.50
1:CR:77:MET:HA	1:CR:80:THR:CG2	2.42	0.50
2:CS:78:TYR:CE1	1:CT:115:MET:HG3	2.47	0.50
2:CU:23:ALA:O	2:CU:27:LEU:HD23	2.12	0.50
2:CU:73:TYR:O	2:CU:74:THR:OG1	2.23	0.50
2:CW:46:ALA:HB2	1:DA:154:ASP:OD2	2.11	0.50

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:CY:47:ARG:HA	1:CY:50:ILE:HG22	1.94	0.50
9:DI:99:ALA:HB1	9:DI:164:THR:HG22	1.94	0.50
1:DL:72:ALA:C	1:DL:77:MET:HB2	2.36	0.50
1:DL:81:CYS:O	1:DL:85:MET:HG3	2.11	0.50
1:DN:5:THR:HG23	2:DO:1:MET:HE3	1.94	0.50
1:DS:76:GLU:HB2	8:DY:203:PHE:CE2	2.47	0.50
9:EA:42:PHE:O	9:EA:46:GLU:HG2	2.12	0.50
9:EA:138:VAL:O	9:EA:142:ILE:HG12	2.11	0.50
9:EA:177:VAL:HG23	11:EA:400:45D:O01	2.12	0.50
9:EA:213:LEU:HB3	9:EA:217:PHE:CE2	2.47	0.50
9:EA:286:ILE:HD11	9:EA:303:ILE:CG2	2.41	0.50
2:AB:96:MET:HA	2:AB:152:TYR:CE2	2.47	0.50
2:AD:47:ASN:O	2:AD:51:ILE:HD12	2.11	0.50
3:AE:2:SER:OG	2:AF:3:ASP:OD2	2.30	0.50
3:AE:112:VAL:O	3:AE:115:MET:HB3	2.12	0.50
2:AF:76:ARG:NH2	5:BA:66:LEU:HD21	2.27	0.50
4:AK:73:ALA:HA	4:AK:78:ARG:HB3	1.94	0.50
1:AN:105:GLU:HA	1:AN:109:LEU:CB	2.36	0.50
1:AN:126:VAL:O	1:AN:130:VAL:HG23	2.11	0.50
2:AO:37:ARG:NH1	2:AO:148:GLU:OE2	2.45	0.50
1:AP:14:GLU:HA	6:BD:548:ARG:NH2	2.27	0.50
2:AQ:127:VAL:O	2:AQ:131:GLN:HG2	2.11	0.50
1:AR:68:PRO:HA	1:AR:73:TYR:CD2	2.46	0.50
2:AW:77:ARG:HG2	10:AW:200:CYC:HAD1	1.94	0.50
2:AW:123:ILE:O	2:AW:127:VAL:HG23	2.12	0.50
6:BD:36:GLU:O	6:BD:40:LEU:HG	2.12	0.50
1:BI:128:GLN:O	1:BI:132:GLU:HG2	2.12	0.50
2:BJ:60:LEU:HD13	2:BJ:72:MET:SD	2.51	0.50
2:BL:11:SER:O	2:BL:14:VAL:HG22	2.11	0.50
2:BL:78:TYR:O	2:BL:82:ILE:HG23	2.12	0.50
2:BN:112:LEU:HD11	2:BN:116:TYR:CZ	2.47	0.50
1:BP:90:ARG:O	1:BP:93:THR:OG1	2.26	0.50
1:BP:112:VAL:CG2	2:BT:75:THR:HG21	2.42	0.50
4:BS:165:SER:OG	2:BW:14:VAL:HG11	2.11	0.50
1:BX:67:SER:O	1:BX:73:TYR:HB2	2.10	0.50
2:CA:132:ALA:HA	2:CA:135:GLU:OE1	2.12	0.50
1:CC:44:THR:O	1:CC:47:ARG:HG2	2.12	0.50
1:CC:141:MET:HE3	1:CC:146:ALA:HA	1.94	0.50
2:CD:127:VAL:O	2:CD:131:GLN:HG3	2.12	0.50
1:CE:62:ARG:NH2	1:CE:124:GLU:OE2	2.44	0.50
1:CE:62:ARG:NH1	1:CE:124:GLU:HG2	2.22	0.50

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:CE:91:LEU:HD21	1:CE:107:ILE:CG2	2.41	0.50
2:CF:134:LYS:HB2	2:CF:153:LEU:HD13	1.94	0.50
6:CM:241:ARG:HG3	6:CM:389:TYR:OH	2.11	0.50
6:CM:611:GLY:HA3	6:CM:656:TYR:O	2.12	0.50
2:CS:22:ALA:O	2:CS:26:LYS:HG2	2.12	0.50
1:CT:108:GLY:C	1:CT:109:LEU:HD22	2.37	0.50
2:DD:58:LYS:HE3	2:DD:132:ALA:HB1	1.94	0.50
9:DI:34:GLU:HB2	9:DI:174:GLU:HG3	1.92	0.50
9:DI:51:LEU:HD22	9:DI:142:ILE:HG22	1.94	0.50
1:DJ:65:ILE:HG22	1:DJ:72:ALA:CB	2.42	0.50
1:DL:18:LEU:O	1:DL:23:LEU:HG	2.12	0.50
2:DM:47:ASN:CG	2:DM:140:LEU:HD21	2.37	0.50
1:DS:39:ILE:HD11	1:DS:145:ASP:CG	2.37	0.50
1:AC:56:ASP:OD1	1:AC:57:ARG:N	2.44	0.50
2:AF:2:GLN:HB3	2:AF:102:SER:OG	2.11	0.50
2:AF:25:ASP:OD1	2:AF:26:LYS:N	2.45	0.50
1:AP:65:ILE:HG13	1:AP:66:VAL:HG23	1.93	0.50
2:AS:60:LEU:HD13	2:AS:72:MET:CE	2.35	0.50
1:AV:37:LEU:HD21	2:AW:24:MET:SD	2.52	0.50
6:BD:285:ALA:O	6:BD:289:ILE:HG12	2.12	0.50
6:BD:364:TYR:HA	6:BD:367:ILE:HD12	1.92	0.50
9:BH:195:ASN:ND2	9:BH:267:ILE:HD13	2.27	0.50
1:BK:126:VAL:HG13	10:BK:200:CYC:HMC2	1.94	0.50
1:BK:131:ARG:CG	1:BK:157:ILE:HD13	2.39	0.50
2:BN:63:SER:H	2:BN:66:THR:CG2	2.24	0.50
2:BQ:2:GLN:HG2	2:BQ:6:THR:CG2	2.42	0.50
2:BT:111:GLY:O	2:BT:115:THR:HG22	2.12	0.50
1:BV:115:MET:HE1	10:BV:200:CYC:CHB	2.41	0.50
2:BW:50:THR:O	2:BW:54:GLU:HG3	2.12	0.50
1:BZ:41:GLU:OE2	2:CA:24:MET:HG3	2.11	0.50
2:CA:100:ASP:OD1	2:CA:101:ALA:N	2.45	0.50
6:CM:694:LEU:HB3	1:DU:83:ARG:HD3	1.94	0.50
8:CP:219:LEU:O	8:CP:223:ARG:HG3	2.12	0.50
9:CQ:86:LEU:HD23	9:CQ:95:CYS:SG	2.52	0.50
9:CQ:267:ILE:HD12	9:CQ:292:LEU:HD12	1.94	0.50
2:CS:1:MET:SD	2:CS:103:ILE:HB	2.52	0.50
1:CY:65:ILE:HG22	1:CY:72:ALA:HB3	1.94	0.50
1:CY:79:ALA:HB3	8:DH:203:PHE:HE1	1.77	0.50
2:CZ:104:LEU:HD13	2:CZ:156:ILE:HG13	1.94	0.50
1:DA:61:LYS:O	9:DI:57:GLY:N	2.42	0.50
2:DB:1:MET:HE3	2:DB:103:ILE:HD12	1.94	0.50

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:DC:84:ASP:OD2	10:DC:200:CYC:ND	2.44	0.50
9:DI:222:ALA:O	9:DI:300:PHE:HA	2.12	0.50
2:DM:24:MET:O	2:DM:28:LYS:HG3	2.12	0.50
2:AB:125:SER:HA	2:AB:128:GLN:OE1	2.12	0.49
1:AC:117:ARG:HD2	1:AH:161:SER:CB	2.42	0.49
2:AD:51:ILE:HA	2:AD:136:VAL:HG11	1.93	0.49
3:AE:116:TYR:HD2	3:AE:121:VAL:HB	1.77	0.49
1:AH:23:LEU:HD13	2:AI:42:SER:OG	2.13	0.49
1:AN:87:TYR:CG	10:AN:200:CYC:HBB3	2.46	0.49
1:AN:97:VAL:HG21	2:AO:19:LEU:CD1	2.41	0.49
1:AP:57:ARG:O	1:AP:61:LYS:HG3	2.12	0.49
2:AU:127:VAL:O	2:AU:131:GLN:HG3	2.12	0.49
2:AW:34:GLY:O	2:AW:38:VAL:HG23	2.12	0.49
2:AY:74:THR:HB	2:AY:77:ARG:HG3	1.94	0.49
5:BA:46:MET:HE3	6:BD:387:ALA:HB3	1.92	0.49
9:BH:263:GLY:O	9:BH:294:PRO:HG3	2.12	0.49
1:BI:75:GLU:OE1	1:BI:75:GLU:N	2.45	0.49
1:BK:92:VAL:HG21	1:BK:153:PHE:CE1	2.47	0.49
2:BL:105:ASP:OD1	2:BL:109:LEU:HD22	2.12	0.49
1:BP:27:LYS:O	1:BP:31:THR:HG23	2.11	0.49
1:BZ:141:MET:CB	1:BZ:146:ALA:HB2	2.39	0.49
1:CE:8:ILE:HD12	2:CF:94:TYR:CD1	2.47	0.49
1:CE:32:GLY:O	1:CE:36:ARG:HG3	2.12	0.49
2:CF:127:VAL:HG22	2:CF:157:CYS:SG	2.52	0.49
6:CM:485:ILE:HG21	6:CM:665:LEU:HD13	1.94	0.49
6:CM:791:PHE:HA	6:CM:795:TYR:HE2	1.76	0.49
8:CP:198:ASN:HA	8:CP:201:ARG:NH2	2.26	0.49
2:DB:44:ILE:HG23	2:DB:89:LEU:HD21	1.94	0.49
2:DD:85:LEU:CG	10:DD:200:CYC:HBC1	2.41	0.49
9:DI:314:ASN:O	9:DI:314:ASN:ND2	2.41	0.49
1:DS:50:ILE:HD11	1:DS:137:ALA:CA	2.42	0.49
1:DS:111:GLY:HA2	1:DS:114:GLU:OE1	2.12	0.49
1:DU:47:ARG:HG3	2:DV:18:TYR:CZ	2.47	0.49
2:DV:58:LYS:HA	8:DZ:206:GLN:CG	2.38	0.49
9:EA:62:GLN:HA	9:EA:65:GLU:OE1	2.11	0.49
3:AE:50:ILE:HD11	3:AE:140:LEU:CB	2.42	0.49
3:AE:101:LYS:HE3	1:AJ:16:ARG:NE	2.27	0.49
3:AE:126:MET:HE2	10:AE:200:CYC:HBC3	1.95	0.49
1:AJ:81:CYS:HA	10:AJ:200:CYC:HHD	1.94	0.49
2:AL:65:VAL:HG12	2:AL:72:MET:CB	2.42	0.49
2:AL:87:TYR:CG	10:AL:200:CYC:HBB3	2.47	0.49

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:AL:97:LEU:HD12	6:BD:35:LEU:HD12	1.94	0.49
1:AP:111:GLY:HA2	1:AP:114:GLU:OE1	2.12	0.49
2:AQ:75:THR:HG23	1:AR:108:GLY:HA2	1.93	0.49
1:AR:17:TYR:OH	2:AS:89:LEU:HG	2.11	0.49
2:AW:3:ASP:OD1	2:AW:6:THR:OG1	2.28	0.49
2:AY:132:ALA:HA	2:AY:135:GLU:OE1	2.12	0.49
6:BD:751:LEU:O	6:BD:755:ILE:HG12	2.12	0.49
9:BH:43:ALA:O	9:BH:51:LEU:HD12	2.12	0.49
1:BX:85:MET:HE3	1:BX:129:SER:HB2	1.93	0.49
2:BY:10:ASN:O	2:BY:14:VAL:HG23	2.12	0.49
2:CD:57:ALA:CA	2:CD:61:LEU:HD12	2.42	0.49
5:CJ:15:ARG:HH12	5:CJ:23:GLN:HE21	1.60	0.49
6:CM:743:TYR:OH	6:CM:764:GLU:OE2	2.19	0.49
1:CR:109:LEU:O	1:CR:112:VAL:HG12	2.12	0.49
1:CT:78:THR:O	1:CT:82:LEU:HG	2.12	0.49
1:DA:63:PRO:O	1:DA:67:SER:N	2.43	0.49
2:DB:15:GLN:HG3	2:DB:17:LYS:HZ1	1.75	0.49
8:DG:196:TRP:CE3	8:DZ:232:ILE:HG21	2.47	0.49
1:DJ:5:THR:HA	1:DJ:8:ILE:HG12	1.93	0.49
2:DO:112:LEU:CD2	2:DO:160:LEU:HD13	2.42	0.49
2:DR:41:ALA:HB1	2:DR:97:LEU:HD11	1.94	0.49
1:DS:77:MET:HE2	1:DS:77:MET:HA	1.94	0.49
1:DS:89:LEU:O	1:DS:93:THR:HG23	2.12	0.49
9:EA:131:LEU:HD23	9:EA:136:ASN:HA	1.93	0.49
9:EA:232:VAL:O	9:EA:236:ASN:ND2	2.31	0.49
1:AA:51:VAL:HG23	1:AA:85:MET:HB3	1.93	0.49
1:AA:76:GLU:O	1:AA:80:THR:HG23	2.12	0.49
2:AD:96:MET:HA	2:AD:152:TYR:CE2	2.48	0.49
3:AE:7:VAL:CG1	3:AE:22:GLU:HB3	2.41	0.49
2:AF:123:ILE:HG23	2:AF:160:LEU:CD2	2.42	0.49
2:AU:2:GLN:NE2	2:AU:6:THR:HG22	2.27	0.49
2:AU:96:MET:HA	2:AU:152:TYR:CE2	2.47	0.49
2:AW:71:ASN:ND2	2:AW:122:PRO:HD3	2.22	0.49
1:AX:61:LYS:HG2	9:BH:56:PRO:HG2	1.94	0.49
5:BA:20:ARG:HH21	6:BD:389:TYR:HA	1.76	0.49
9:BH:107:LEU:HD23	11:BH:400:45D:H141	1.94	0.49
2:BJ:104:LEU:CD1	2:BJ:152:TYR:HB3	2.43	0.49
1:BR:65:ILE:HG13	1:BR:66:VAL:HG23	1.94	0.49
2:BT:101:ALA:CB	2:BT:104:LEU:HD12	2.42	0.49
1:BV:103:PRO:O	1:BV:107:ILE:HG12	2.13	0.49
2:BW:74:THR:HG23	1:BX:107:ILE:O	2.13	0.49

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:CE:59:PHE:CZ	1:CE:78:THR:HG23	2.47	0.49
2:CF:74:THR:OG1	2:CF:77:ARG:NH2	2.32	0.49
2:CF:74:THR:HG1	2:CF:77:ARG:HH21	1.53	0.49
6:CM:248:ILE:HG12	6:CM:356:SER:HB2	1.94	0.49
2:CS:36:LEU:HD21	2:CS:145:ALA:HB2	1.94	0.49
2:CZ:35:GLU:O	2:CZ:38:VAL:HG12	2.13	0.49
1:DA:51:VAL:HA	1:DA:133:MET:HE2	1.93	0.49
1:DA:61:LYS:HB3	9:DI:57:GLY:HA3	1.95	0.49
1:DA:61:LYS:O	1:DA:62:ARG:HG2	2.12	0.49
5:DF:6:ILE:HD13	5:DF:36:TRP:CZ3	2.46	0.49
9:DI:248:LEU:O	9:DI:249:LYS:HE2	2.12	0.49
2:DK:123:ILE:O	2:DK:127:VAL:HG23	2.12	0.49
1:DN:101:VAL:HB	1:DN:155:PHE:CE2	2.48	0.49
1:DQ:79:ALA:HB3	8:DZ:203:PHE:HE1	1.77	0.49
2:DT:24:MET:O	2:DT:28:LYS:HG3	2.12	0.49
1:DU:27:LYS:NZ	2:DV:38:VAL:HB	2.27	0.49
9:EA:27:ARG:HA	9:EA:30:GLN:OE1	2.12	0.49
9:EA:86:LEU:CD2	9:EA:95:CYS:HA	2.40	0.49
2:AB:4:ALA:O	2:AB:8:VAL:HG23	2.12	0.49
10:AD:200:CYC:CGA	5:BA:12:SER:HB3	2.42	0.49
3:AE:23:LEU:HD13	2:AF:42:SER:OG	2.12	0.49
1:AH:102:THR:O	1:AH:106:GLU:HG2	2.12	0.49
1:AH:122:PRO:O	1:AH:126:VAL:HG23	2.11	0.49
2:AS:19:LEU:CB	2:AS:24:MET:HE2	2.42	0.49
1:AX:36:ARG:NH2	1:AX:99:GLY:HA2	2.26	0.49
2:AY:63:SER:O	2:AY:66:THR:HG22	2.12	0.49
6:BD:15:GLN:NE2	6:BD:18:GLN:HA	2.27	0.49
6:BD:251:LEU:HD11	6:BD:407:GLU:HB3	1.93	0.49
7:BF:107:LEU:O	7:BF:111:LYS:HG3	2.12	0.49
1:BI:16:ARG:O	2:BJ:94:TYR:OH	2.30	0.49
2:BJ:1:MET:HE3	2:BJ:103:ILE:HD13	1.92	0.49
2:BJ:61:LEU:HD22	10:BK:200:CYC:CAA	2.43	0.49
3:BM:38:ARG:HA	3:BM:41:GLU:OE1	2.11	0.49
1:BR:64:ASP:OD1	1:BR:64:ASP:N	2.43	0.49
1:BV:49:THR:O	1:BV:53:GLN:HG3	2.12	0.49
1:BV:87:TYR:CD2	10:BV:200:CYC:HBB3	2.47	0.49
2:CA:19:LEU:HB2	2:CA:24:MET:HE2	1.94	0.49
2:CD:41:ALA:N	2:CD:96:MET:HE1	2.28	0.49
2:CD:44:ILE:CD1	2:CD:137:THR:HG23	2.40	0.49
2:CD:64:ASP:HA	2:CD:67:ARG:HH11	1.77	0.49
6:CM:55:GLU:O	6:CM:59:THR:HG23	2.13	0.49

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
6:CM:86:GLU:OE2	6:CM:150:GLN:HB3	2.12	0.49
6:CM:749:ARG:HH12	10:DV:200:CYC:CGA	2.25	0.49
6:CM:791:PHE:HA	6:CM:795:TYR:CE2	2.47	0.49
10:CW:200:CYC:HB	10:CW:200:CYC:CMA	2.23	0.49
2:CZ:134:LYS:HG3	2:CZ:150:GLY:CA	2.43	0.49
1:DA:134:LYS:HD3	1:DA:153:PHE:HB3	1.94	0.49
2:DB:22:ALA:O	2:DB:26:LYS:HG3	2.13	0.49
2:DB:124:SER:CB	2:DT:124:SER:HB2	2.36	0.49
2:DD:5:ILE:HD11	2:DD:31:PHE:HE1	1.76	0.49
2:DD:44:ILE:HD12	2:DD:149:MET:HE3	1.94	0.49
1:DJ:65:ILE:O	1:DJ:72:ALA:HB3	2.12	0.49
1:DJ:118:SER:O	2:DO:53:LYS:HG3	2.11	0.49
2:DK:36:LEU:O	2:DK:39:ARG:HG3	2.12	0.49
1:DL:64:ASP:O	1:DL:70:GLY:HA3	2.11	0.49
1:DN:78:THR:O	1:DN:82:LEU:HD23	2.11	0.49
1:DQ:36:ARG:NH1	1:DQ:99:GLY:HA2	2.27	0.49
1:DS:85:MET:HE3	1:DS:129:SER:HB2	1.95	0.49
2:DT:115:THR:O	2:DT:119:LEU:HD23	2.12	0.49
9:EA:68:LEU:CD1	9:EA:112:ARG:HG2	2.43	0.49
9:EA:107:LEU:HD21	11:EA:400:45D:H283	1.94	0.49
9:EA:223:LEU:HG	9:EA:301:VAL:HG13	1.93	0.49
1:AA:58:LEU:HD22	1:AA:129:SER:CB	2.37	0.49
1:AC:39:ILE:HD11	1:AC:145:ASP:CG	2.38	0.49
1:AC:127:ALA:O	1:AC:131:ARG:HG3	2.13	0.49
2:AI:2:GLN:O	2:AI:100:ASP:HB3	2.12	0.49
1:AJ:27:LYS:O	1:AJ:31:THR:HG23	2.12	0.49
4:AK:86:MET:HE3	10:AK:200:CYC:HBC1	1.94	0.49
1:AR:38:ARG:NH1	1:AR:145:ASP:OD2	2.44	0.49
2:AS:65:VAL:HG12	2:AS:72:MET:HB3	1.95	0.49
2:AU:39:ARG:O	2:AU:43:VAL:HG23	2.13	0.49
6:BD:308:GLN:HB3	6:BD:314:ILE:HG12	1.93	0.49
6:BD:649:PHE:CE1	6:BD:655:PRO:HA	2.48	0.49
2:BJ:68:PRO:HA	2:BJ:73:TYR:CG	2.48	0.49
1:BK:71:ASN:O	1:BK:77:MET:HB3	2.12	0.49
2:BT:96:MET:SD	2:BT:149:MET:HG2	2.53	0.49
1:BV:76:GLU:HB3	7:CO:103:PRO:HG2	1.94	0.49
1:BV:127:ALA:CB	1:BV:161:SER:HB3	2.43	0.49
2:BW:10:ASN:O	2:BW:14:VAL:HG22	2.12	0.49
2:BW:83:ARG:HH21	6:CM:490:ALA:HB1	1.77	0.49
1:CG:46:SER:O	1:CG:50:ILE:HG22	2.13	0.49
1:CR:33:GLY:HA3	2:CS:31:PHE:CD2	2.48	0.49

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:CU:65:VAL:HG12	2:CU:72:MET:CB	2.40	0.49
2:DB:126:THR:HG22	10:DB:200:CYC:H3C	1.95	0.49
9:DI:86:LEU:HD21	9:DI:95:CYS:HA	1.95	0.49
1:DJ:98:SER:HA	2:DK:5:ILE:HG21	1.93	0.49
2:DK:40:ALA:O	2:DK:44:ILE:HG13	2.12	0.49
2:DK:41:ALA:CB	2:DK:97:LEU:HD21	2.42	0.49
1:DL:109:LEU:HD13	1:DL:159:LYS:HG3	1.94	0.49
2:DO:71:ASN:ND2	2:DO:121:VAL:HA	2.28	0.49
2:DT:88:TYR:OH	2:DT:112:LEU:HD21	2.11	0.49
2:DV:60:LEU:CB	2:DV:72:MET:HE1	2.38	0.49
1:AA:128:GLN:O	1:AA:132:GLU:HG2	2.13	0.49
2:AD:23:ALA:O	2:AD:27:LEU:HG	2.12	0.49
3:AE:42:THR:HG21	3:AE:141:LEU:HD21	1.94	0.49
1:AH:152:TYR:O	1:AH:156:VAL:HG23	2.12	0.49
2:AI:44:ILE:HG21	2:AI:93:THR:CG2	2.41	0.49
4:AK:113:LEU:HD23	6:BD:408:CYS:SG	2.53	0.49
2:AO:74:THR:HG23	1:AP:107:ILE:O	2.12	0.49
2:AS:68:PRO:HA	2:AS:73:TYR:CD1	2.48	0.49
2:AU:2:GLN:HB2	2:AU:102:SER:HB3	1.94	0.49
1:AV:68:PRO:HA	1:AV:73:TYR:CD1	2.48	0.49
6:BD:86:GLU:OE2	6:BD:150:GLN:HB3	2.12	0.49
6:BD:626:ILE:HD13	6:BD:638:ALA:CB	2.42	0.49
9:BH:217:PHE:CE2	9:BH:237:VAL:HG11	2.47	0.49
9:BH:263:GLY:C	9:BH:294:PRO:HG3	2.37	0.49
1:BK:134:LYS:HE3	1:BK:150:SER:CB	2.41	0.49
2:BL:67:ARG:HG3	2:BL:68:PRO:HD2	1.95	0.49
3:BM:134:LYS:O	3:BM:138:LEU:HG	2.12	0.49
2:BT:133:ILE:O	2:BT:137:THR:HG23	2.12	0.49
2:BY:53:LYS:HE3	2:BY:54:GLU:OE2	2.12	0.49
1:CE:14:GLU:HB2	1:CE:16:ARG:HG2	1.94	0.49
1:CE:39:ILE:CD1	1:CE:145:ASP:HB3	2.42	0.49
2:CF:3:ASP:HB3	2:CF:98:ALA:HB1	1.94	0.49
5:CJ:18:THR:HA	6:CM:240:VAL:H	1.77	0.49
5:CK:2:ARG:NH2	5:CK:66:LEU:HD13	2.28	0.49
6:CM:151:LYS:CD	6:CM:154:ARG:HH21	2.25	0.49
6:CM:793:ALA:HB3	6:CM:794:PRO:HD3	1.94	0.49
9:CQ:273:VAL:HB	9:CQ:286:ILE:HD11	1.94	0.49
9:CQ:289:ARG:HB3	9:CQ:302:ALA:HB3	1.95	0.49
1:CR:34:ALA:HA	2:CS:28:LYS:NZ	2.28	0.49
2:CS:78:TYR:HE1	1:CT:115:MET:HA	1.77	0.49
2:DB:74:THR:O	2:DB:77:ARG:N	2.38	0.49

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:DC:122:PRO:HD2	10:DC:200:CYC:HMC1	1.95	0.49
2:DD:122:PRO:HG2	10:DD:200:CYC:CMC	2.39	0.49
2:DD:132:ALA:HA	2:DD:135:GLU:OE1	2.12	0.49
9:DI:204:ASN:O	9:DI:209:ASP:N	2.26	0.49
9:DI:274:GLN:HA	9:DI:282:VAL:O	2.13	0.49
2:DO:107:ARG:NH2	5:DX:45:LYS:HG2	2.27	0.49
1:DS:50:ILE:HD11	1:DS:137:ALA:CB	2.42	0.49
1:DS:58:LEU:HD22	1:DS:129:SER:HB3	1.95	0.49
9:EA:222:ALA:HB3	9:EA:299:PHE:O	2.12	0.49
1:AA:122:PRO:HB2	1:AA:125:ALA:HB3	1.95	0.49
1:AC:92:VAL:O	1:AC:96:VAL:HG23	2.13	0.49
2:AD:72:MET:HE3	2:AD:78:TYR:CE1	2.48	0.49
2:AD:147:LYS:O	2:AD:151:VAL:HG23	2.12	0.49
2:AF:88:TYR:OH	2:AF:116:TYR:OH	2.30	0.49
1:AN:33:GLY:HA2	1:AN:36:ARG:HG2	1.95	0.49
1:AN:86:ASP:OD2	7:BF:109:ARG:NH2	2.46	0.49
2:AO:3:ASP:N	2:AO:6:THR:OG1	2.46	0.49
1:AZ:141:MET:HE3	1:AZ:146:ALA:HA	1.95	0.49
9:BH:140:ALA:HA	9:BH:143:GLN:OE1	2.12	0.49
1:BP:62:ARG:HG2	1:BP:65:ILE:HG12	1.94	0.49
1:BR:3:ILE:HD11	1:BR:99:GLY:O	2.13	0.49
4:BS:86:MET:HE3	10:BS:200:CYC:HBC1	1.94	0.49
1:BX:75:GLU:OE1	1:BX:75:GLU:N	2.44	0.49
2:CA:71:ASN:HD21	2:CA:121:VAL:HG22	1.77	0.49
1:CC:71:ASN:HD22	1:CC:121:THR:HA	1.78	0.49
2:CF:77:ARG:CB	10:CF:200:CYC:HMD1	2.42	0.49
6:CM:804:GLY:O	6:CM:838:MET:HE1	2.12	0.49
1:CR:57:ARG:O	1:CR:61:LYS:HG2	2.13	0.49
1:CY:75:GLU:O	1:CY:78:THR:OG1	2.27	0.49
2:DD:96:MET:HA	2:DD:152:TYR:CE2	2.48	0.49
5:DF:6:ILE:HG22	5:DF:53:VAL:CG2	2.33	0.49
8:DG:221:MET:CE	1:DQ:51:VAL:HG13	2.39	0.49
1:DJ:131:ARG:CG	1:DJ:157:ILE:HD13	2.39	0.49
2:DK:92:ALA:HB1	2:DK:149:MET:HE1	1.95	0.49
2:DO:62:TYR:H	2:DO:66:THR:HG21	1.76	0.49
2:DR:47:ASN:HB2	2:DR:50:THR:HB	1.94	0.49
1:DS:88:TYR:CZ	1:DS:160:MET:HE1	2.46	0.49
1:DU:39:ILE:HG23	1:DU:141:MET:HE1	1.95	0.49
8:DY:237:ALA:HA	8:DY:240:ILE:HD13	1.95	0.49
9:EA:210:PHE:HB3	9:EA:241:PHE:HB3	1.94	0.49
1:AC:65:ILE:HD11	1:AC:122:PRO:CG	2.43	0.49

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:AC:131:ARG:HA	1:AC:157:ILE:HD11	1.95	0.49
3:AE:50:ILE:CD1	3:AE:140:LEU:HD13	2.43	0.49
4:AK:88:TYR:HH	6:BD:252:GLU:N	2.03	0.49
2:AL:1:MET:HB3	2:AL:106:GLU:OE2	2.13	0.49
2:AL:78:TYR:O	2:AL:82:ILE:HG12	2.12	0.49
1:AN:58:LEU:HD22	1:AN:129:SER:HB2	1.95	0.49
1:AN:107:ILE:HD12	2:AO:13:ASP:OD2	2.12	0.49
2:AO:72:MET:HE2	2:AO:78:TYR:CD2	2.48	0.49
2:AO:134:LYS:HB2	2:AO:153:LEU:HD13	1.95	0.49
1:AP:59:PHE:CD1	1:AP:66:VAL:HG21	2.48	0.49
2:AU:1:MET:HE1	1:AZ:9:VAL:HG22	1.94	0.49
1:AX:92:VAL:O	1:AX:96:VAL:HG13	2.13	0.49
1:AZ:81:CYS:O	1:AZ:85:MET:HG2	2.13	0.49
6:BD:542:LEU:HD11	6:BD:654:VAL:HG23	1.94	0.49
6:BD:885:VAL:HG13	6:BD:885:VAL:O	2.13	0.49
9:BH:235:GLU:HG2	1:DS:45:GLY:HA2	1.94	0.49
1:BI:127:ALA:HB1	1:BI:157:ILE:HG23	1.94	0.49
1:BK:33:GLY:HA2	1:BK:36:ARG:NE	2.28	0.49
1:BK:122:PRO:HG2	1:BK:125:ALA:CB	2.43	0.49
2:BL:23:ALA:O	2:BL:27:LEU:HG	2.13	0.49
1:BP:156:VAL:O	1:BP:160:MET:HG3	2.13	0.49
2:BT:72:MET:HE3	2:BT:81:CYS:CB	2.41	0.49
2:CD:65:VAL:HG12	2:CD:72:MET:HB2	1.94	0.49
1:CE:58:LEU:CD2	1:CE:129:SER:HB3	2.36	0.49
1:CE:115:MET:O	1:CE:119:LEU:HD23	2.13	0.49
1:CG:102:THR:HG21	6:CM:534:PHE:CE1	2.47	0.49
2:CH:41:ALA:HB2	2:CH:96:MET:CE	2.43	0.49
6:CM:793:ALA:O	2:DR:107:ARG:HD3	2.12	0.49
9:CQ:263:GLY:C	9:CQ:294:PRO:HG3	2.37	0.49
1:CY:2:SER:HB3	1:CY:5:THR:HG22	1.94	0.49
1:CY:11:ALA:HA	1:CY:16:ARG:CZ	2.43	0.49
1:CY:30:VAL:HA	2:CZ:31:PHE:HD1	1.76	0.49
1:CY:52:LYS:HG2	8:DY:225:ILE:HG12	1.95	0.49
1:CY:112:VAL:HA	1:CY:115:MET:HG2	1.95	0.49
2:CZ:47:ASN:O	2:CZ:51:ILE:HD12	2.12	0.49
1:DA:4:VAL:O	1:DA:8:ILE:HG12	2.13	0.49
1:DC:22:GLU:O	1:DC:26:ILE:HG12	2.12	0.49
2:DD:73:TYR:O	2:DD:74:THR:OG1	2.24	0.49
9:DI:80:THR:HG22	9:DI:176:VAL:HG22	1.94	0.49
9:DI:248:LEU:HD21	9:DI:277:TRP:CE3	2.48	0.49
1:DJ:102:THR:HB	1:DJ:103:PRO:HD3	1.95	0.49

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:DN:62:ARG:O	1:DN:65:ILE:HG12	2.12	0.49
1:AC:25:ARG:NE	6:BD:42:GLU:OE2	2.42	0.49
1:AC:103:PRO:O	1:AC:107:ILE:HG12	2.13	0.49
1:AC:111:GLY:HA2	1:AC:114:GLU:CD	2.38	0.49
2:AD:82:ILE:HA	2:AD:85:LEU:HD12	1.95	0.49
3:AE:138:LEU:HD22	3:AE:146:ALA:HB1	1.95	0.49
2:AF:12:ALA:HB1	2:AF:17:LYS:O	2.13	0.49
1:AH:68:PRO:HA	1:AH:73:TYR:CE2	2.47	0.49
1:AJ:26:ILE:HD13	4:AK:98:ILE:HG21	1.95	0.49
4:AK:67:ILE:HG21	10:BD:902:CYC:CBA	2.42	0.49
4:AK:125:GLY:O	4:AK:128:VAL:HG22	2.13	0.49
1:AN:9:VAL:CG2	2:AO:1:MET:HE2	2.36	0.49
2:AO:1:MET:N	2:AO:106:GLU:OE1	2.44	0.49
2:AO:2:GLN:OE1	2:AO:10:ASN:ND2	2.28	0.49
2:AU:36:LEU:HA	2:AU:39:ARG:HG2	1.94	0.49
1:AX:39:ILE:O	1:AX:43:LEU:HG	2.13	0.49
5:BA:4:PHE:HE1	5:BA:33:TYR:HD1	1.61	0.49
5:BA:8:ALA:HB2	5:BA:50:ILE:HG22	1.94	0.49
5:BA:43:ILE:HG21	5:BA:50:ILE:HG23	1.94	0.49
6:BD:44:THR:O	6:BD:48:GLN:HG3	2.13	0.49
6:BD:205:ALA:HA	6:BD:208:ASP:OD2	2.13	0.49
9:BH:290:PHE:CD2	9:BH:298:ILE:HD13	2.48	0.49
2:BJ:78:TYR:OH	1:BK:119:LEU:HD11	2.11	0.49
2:BJ:125:SER:HA	2:BJ:128:GLN:OE1	2.12	0.49
1:CC:101:VAL:HG12	1:CC:152:TYR:HD1	1.78	0.49
2:CF:67:ARG:O	2:CF:73:TYR:HB2	2.13	0.49
2:CH:147:LYS:O	2:CH:151:VAL:HG23	2.12	0.49
5:CJ:12:SER:OG	5:CJ:15:ARG:HG3	2.13	0.49
6:CM:863:ALA:HB2	2:DT:107:ARG:O	2.12	0.49
9:CQ:43:ALA:O	9:CQ:51:LEU:HD12	2.13	0.49
9:CQ:263:GLY:O	9:CQ:294:PRO:HG3	2.13	0.49
1:CY:47:ARG:O	1:CY:51:VAL:HG12	2.12	0.49
1:CY:66:VAL:HA	1:CY:73:TYR:HA	1.95	0.49
1:CY:109:LEU:HD21	1:CY:159:LYS:CB	2.35	0.49
2:CZ:23:ALA:HA	2:CZ:26:LYS:HE3	1.93	0.49
1:DJ:26:ILE:O	1:DJ:30:VAL:HG13	2.13	0.49
2:DO:63:SER:H	2:DO:66:THR:HG22	1.77	0.49
1:DQ:87:TYR:OH	2:DV:68:PRO:HD3	2.12	0.49
10:DR:200:CYC:HC	10:DR:200:CYC:CMD	2.26	0.49
1:DS:12:ASP:OD2	2:DT:107:ARG:HD3	2.12	0.49
1:DS:113:ARG:NH1	1:DS:160:MET:O	2.46	0.49

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:DT:62:TYR:OH	10:DU:200:CYC:O2D	2.17	0.49
2:DV:79:ALA:HA	2:DV:82:ILE:HG12	1.94	0.49
2:DV:130:ILE:HA	2:DV:133:ILE:HG22	1.94	0.49
2:DV:134:LYS:HG3	2:DV:150:GLY:HA2	1.94	0.49
9:EA:24:THR:HG21	9:EA:142:ILE:CD1	2.43	0.49
2:AB:93:THR:HA	2:AB:149:MET:HE1	1.94	0.49
1:AC:123:ILE:HA	1:AC:126:VAL:CG2	2.43	0.49
2:AD:87:TYR:CE2	5:BA:22:LEU:HD13	2.47	0.49
2:AD:93:THR:N	2:AD:149:MET:HE1	2.28	0.49
2:AD:103:ILE:HG23	2:AD:104:LEU:HD22	1.95	0.49
2:AF:132:ALA:O	2:AF:136:VAL:HG23	2.13	0.49
1:AJ:134:LYS:HD3	1:AJ:153:PHE:HB3	1.94	0.49
4:AK:113:LEU:HD21	10:AK:200:CYC:HMB3	1.95	0.49
1:AV:131:ARG:HG2	1:AV:157:ILE:HD13	1.95	0.49
2:AW:24:MET:HG3	2:AW:28:LYS:HZ3	1.78	0.49
2:AW:36:LEU:HD23	2:AW:39:ARG:NH1	2.24	0.49
2:AW:77:ARG:CB	10:AW:200:CYC:HMD1	2.43	0.49
6:BD:155:ASP:OD1	10:BD:902:CYC:HHB	2.13	0.49
6:BD:455:GLU:HG3	6:BD:664:GLY:CA	2.34	0.49
1:BI:85:MET:O	1:BI:133:MET:HE1	2.12	0.49
1:BK:44:THR:O	1:BK:47:ARG:HG2	2.12	0.49
1:BK:111:GLY:HA2	1:BK:114:GLU:CD	2.38	0.49
2:BL:4:ALA:HB3	2:BL:30:TYR:CD2	2.48	0.49
1:BV:33:GLY:O	1:BV:37:LEU:HD13	2.13	0.49
1:BV:88:TYR:CE2	1:BV:160:MET:HE1	2.47	0.49
2:BY:124:SER:O	2:BY:128:GLN:HG3	2.13	0.49
1:BZ:97:VAL:CG1	2:CA:27:LEU:HD13	2.42	0.49
2:CF:21:GLY:O	2:CF:24:MET:HB3	2.13	0.49
1:CG:27:LYS:O	1:CG:31:THR:HG23	2.12	0.49
6:CM:247:ASP:HB3	6:CM:357:SER:OG	2.13	0.49
6:CM:251:LEU:HD11	6:CM:407:GLU:HB3	1.93	0.49
6:CM:864:ALA:HB2	2:DT:110:ASN:O	2.13	0.49
1:CR:131:ARG:O	1:CR:134:LYS:HB3	2.13	0.49
2:CW:14:VAL:HG23	2:CW:15:GLN:OE1	2.13	0.49
2:CW:50:THR:O	2:CW:54:GLU:HG2	2.12	0.49
1:CY:122:PRO:HG2	1:CY:125:ALA:CB	2.41	0.49
1:DA:109:LEU:HD13	1:DA:159:LYS:CG	2.37	0.49
5:DF:40:GLN:O	5:DF:44:MET:HG3	2.13	0.49
9:DI:214:ILE:CD1	9:DI:238:LEU:HB2	2.42	0.49
1:DJ:20:PRO:HA	1:DJ:23:LEU:HG	1.93	0.49
1:DJ:48:GLU:HA	1:DJ:51:VAL:HG12	1.95	0.49

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:DL:17:TYR:CD1	2:DM:90:ARG:HD3	2.48	0.49
2:DM:44:ILE:CD1	2:DM:149:MET:HE2	2.42	0.49
2:DM:112:LEU:HD13	10:DM:200:CYC:HMB2	1.94	0.49
1:DN:39:ILE:HD11	1:DN:145:ASP:HA	1.95	0.49
1:DQ:72:ALA:HB2	10:DQ:200:CYC:CMD	2.42	0.49
2:DT:78:TYR:CD1	1:DU:115:MET:HG3	2.47	0.49
1:DU:90:ARG:O	1:DU:93:THR:OG1	2.27	0.49
2:AB:77:ARG:CB	10:AB:200:CYC:HMD1	2.43	0.48
1:AC:25:ARG:HD3	6:BD:42:GLU:HG3	1.95	0.48
1:AC:97:VAL:HG11	2:AD:19:LEU:CD1	2.41	0.48
1:AC:129:SER:HA	1:AC:132:GLU:OE1	2.13	0.48
1:AH:16:ARG:NH1	1:AH:17:TYR:O	2.45	0.48
2:AL:31:PHE:CE1	6:BD:50:GLY:HA3	2.48	0.48
1:AN:65:ILE:HD11	1:AN:122:PRO:HG2	1.95	0.48
2:AQ:47:ASN:CB	2:AQ:140:LEU:HD21	2.42	0.48
1:AR:77:MET:HE2	9:BH:11:ILE:HD13	1.95	0.48
2:AS:123:ILE:HA	2:AS:126:THR:CG2	2.40	0.48
2:AU:137:THR:CG2	2:AU:149:MET:HG2	2.43	0.48
2:AW:103:ILE:CD1	2:AW:107:ARG:HD2	2.40	0.48
6:BD:285:ALA:HA	6:BD:395:VAL:HG11	1.93	0.48
6:BD:744:ARG:HG3	6:BD:750:ASP:OD1	2.13	0.48
6:BD:751:LEU:HD13	6:BD:755:ILE:HD13	1.95	0.48
1:BR:37:LEU:HD11	4:BS:31:PHE:HE2	1.78	0.48
4:BS:109:VAL:HG12	4:BS:113:LEU:HD11	1.95	0.48
1:BV:113:ARG:HD3	1:CE:117:ARG:NH2	2.28	0.48
2:CD:71:ASN:ND2	2:CD:121:VAL:HA	2.26	0.48
1:CE:66:VAL:HA	1:CE:72:ALA:O	2.12	0.48
1:CE:119:LEU:HD12	10:CE:200:CYC:HBD1	1.94	0.48
1:CG:30:VAL:HG13	2:CH:31:PHE:HA	1.95	0.48
6:CM:136:PHE:CD1	6:CM:197:VAL:HG11	2.48	0.48
6:CM:513:PHE:HB3	6:CM:542:LEU:HD13	1.94	0.48
6:CM:809:LEU:HG	6:CM:838:MET:HE3	1.95	0.48
1:CR:25:ARG:HA	1:CY:25:ARG:NE	2.28	0.48
1:CR:134:LYS:HA	1:CR:153:PHE:CD2	2.48	0.48
2:CS:24:MET:O	2:CS:28:LYS:HG2	2.13	0.48
2:CS:68:PRO:HA	2:CS:73:TYR:CD2	2.47	0.48
2:CW:57:ALA:CB	2:CW:61:LEU:HD12	2.42	0.48
8:DH:216:MET:HE2	8:DH:219:LEU:HD23	1.94	0.48
8:DH:230:ASN:HB2	1:DS:67:SER:OG	2.13	0.48
9:DI:149:GLN:O	9:DI:153:VAL:HG23	2.13	0.48
1:DJ:20:PRO:HA	1:DJ:23:LEU:CG	2.43	0.48

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:DJ:144:ASP:OD1	1:DJ:145:ASP:N	2.46	0.48
1:DN:98:SER:OG	1:DN:100:ASP:OD1	2.18	0.48
1:DS:5:THR:O	1:DS:9:VAL:HG23	2.12	0.48
2:DT:68:PRO:HD3	1:DU:87:TYR:OH	2.13	0.48
9:EA:115:GLU:O	9:EA:119:GLN:HG3	2.13	0.48
1:AA:10:ASN:O	1:AA:14:GLU:HG3	2.13	0.48
1:AA:92:VAL:O	1:AA:96:VAL:HG23	2.13	0.48
2:AF:39:ARG:NH2	1:CC:142:SER:HA	2.27	0.48
1:AH:115:MET:HE1	10:AH:200:CYC:CMB	2.43	0.48
2:AL:134:LYS:HD2	2:AL:150:GLY:HA3	1.95	0.48
2:AO:142:GLY:HA2	1:BR:41:GLU:OE1	2.13	0.48
2:AS:65:VAL:HG12	2:AS:72:MET:CB	2.42	0.48
2:AS:113:LYS:NZ	6:BD:10:SER:OG	2.26	0.48
2:AU:28:LYS:CD	1:AZ:37:LEU:HD13	2.42	0.48
1:AV:71:ASN:ND2	1:AV:119:LEU:O	2.31	0.48
1:AV:88:TYR:CZ	1:AV:160:MET:HE1	2.47	0.48
2:AY:36:LEU:HD21	2:AY:145:ALA:HB2	1.93	0.48
2:AY:123:ILE:HA	2:AY:126:THR:HG22	1.96	0.48
5:BB:41:GLN:HG3	5:BB:45:LYS:HD2	1.94	0.48
1:BK:76:GLU:HG2	1:BK:77:MET:CE	2.44	0.48
1:BK:78:THR:O	1:BK:82:LEU:HD23	2.13	0.48
2:BL:5:ILE:HG23	2:BL:30:TYR:HD2	1.78	0.48
1:BV:33:GLY:O	1:BV:36:ARG:HG2	2.12	0.48
2:BW:5:ILE:HD11	2:BW:31:PHE:HE1	1.78	0.48
1:BX:101:VAL:HB	1:BX:155:PHE:CE2	2.48	0.48
1:BZ:52:LYS:HD3	8:CP:221:MET:HG3	1.95	0.48
1:BZ:52:LYS:HE3	8:CP:224:SER:OG	2.14	0.48
2:CA:123:ILE:HA	2:CA:126:THR:CG2	2.42	0.48
6:CM:251:LEU:HB3	6:CM:405:ALA:HB1	1.95	0.48
6:CM:597:TYR:CE2	6:CM:599:CYS:HB2	2.48	0.48
1:CR:155:PHE:O	1:CR:159:LYS:HG2	2.13	0.48
1:CT:33:GLY:O	1:CT:37:LEU:HD23	2.13	0.48
1:CV:39:ILE:HG23	1:CV:141:MET:SD	2.53	0.48
2:CW:65:VAL:HG12	2:CW:72:MET:HB2	1.95	0.48
1:CY:26:ILE:CD1	2:CZ:97:LEU:HD21	2.35	0.48
1:CY:134:LYS:HA	1:CY:153:PHE:CD2	2.47	0.48
1:DA:12:ASP:OD2	2:DB:107:ARG:NH1	2.35	0.48
1:DC:49:THR:HG22	1:DC:53:GLN:OE1	2.13	0.48
2:DD:122:PRO:HG2	2:DD:125:SER:HB2	1.95	0.48
9:DI:190:ILE:HD11	9:DI:202:MET:SD	2.53	0.48
1:DJ:62:ARG:HB3	1:DJ:65:ILE:HG12	1.96	0.48

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:DM:90:ARG:O	2:DM:93:THR:HG22	2.12	0.48
1:DQ:76:GLU:HA	8:DZ:203:PHE:CE1	2.48	0.48
1:DS:90:ARG:HG3	2:DT:18:TYR:CD1	2.48	0.48
1:AC:13:ALA:HB1	6:BD:265:GLN:HB3	1.96	0.48
2:AD:112:LEU:HD21	10:AD:200:CYC:C4B	2.43	0.48
1:AH:62:ARG:NH1	1:AH:124:GLU:OE2	2.44	0.48
1:AH:109:LEU:HD21	1:AH:159:LYS:HG2	1.94	0.48
10:AI:200:CYC:HB	10:AI:200:CYC:CMA	2.27	0.48
4:AK:62:GLU:CD	4:AK:133:ILE:HD11	2.38	0.48
1:AN:105:GLU:CA	1:AN:109:LEU:HB2	2.37	0.48
2:AQ:96:MET:HA	2:AQ:152:TYR:CE2	2.49	0.48
1:AX:19:SER:HB3	1:AX:22:GLU:HG3	1.94	0.48
1:AZ:85:MET:O	1:AZ:133:MET:HE1	2.14	0.48
1:AZ:113:ARG:NH1	1:AZ:160:MET:O	2.46	0.48
5:BB:23:GLN:O	5:BB:27:PHE:HB3	2.12	0.48
9:BH:195:ASN:O	9:BH:199:LEU:HG	2.14	0.48
2:BJ:126:THR:O	2:BJ:130:ILE:HG12	2.12	0.48
3:BM:51:VAL:HG23	3:BM:85:TYR:HB3	1.95	0.48
2:BN:72:MET:HE2	2:BN:78:TYR:CE2	2.48	0.48
2:BN:126:THR:HG22	10:BN:200:CYC:HMC1	1.95	0.48
1:BV:65:ILE:HG22	1:BV:72:ALA:HB3	1.94	0.48
1:BV:82:LEU:O	1:BV:85:MET:HB3	2.13	0.48
2:BW:147:LYS:O	2:BW:151:VAL:HG23	2.14	0.48
1:BZ:88:TYR:O	1:BZ:92:VAL:HG23	2.12	0.48
1:BZ:122:PRO:O	1:BZ:126:VAL:HG23	2.14	0.48
1:CC:71:ASN:OD1	10:CC:200:CYC:HBD2	2.13	0.48
2:CD:19:LEU:CB	2:CD:24:MET:HE2	2.43	0.48
1:CE:65:ILE:HD11	1:CE:125:ALA:HB1	1.95	0.48
1:CE:75:GLU:OE2	1:CE:76:GLU:HG3	2.13	0.48
6:CM:155:ASP:OD1	10:CM:902:CYC:HHB	2.13	0.48
9:CQ:288:TRP:HB3	9:CQ:290:PHE:HE1	1.75	0.48
2:CW:54:GLU:O	2:CW:58:LYS:HG2	2.12	0.48
9:DI:224:GLN:HB2	9:DI:300:PHE:CE1	2.48	0.48
1:DJ:34:ALA:HA	2:DK:28:LYS:NZ	2.28	0.48
1:DJ:57:ARG:HA	1:DJ:60:GLN:OE1	2.13	0.48
1:DJ:65:ILE:HG21	10:DJ:200:CYC:H2C	1.96	0.48
1:DN:126:VAL:CG2	10:DN:200:CYC:HMC2	2.43	0.48
9:EA:173:ALA:O	9:EA:175:PRO:HD3	2.14	0.48
9:EA:190:ILE:HG21	9:EA:193:VAL:HB	1.95	0.48
2:AB:77:ARG:HG2	10:AB:200:CYC:HAD1	1.96	0.48
3:AE:94:TYR:CG	2:AF:9:ILE:HG23	2.49	0.48

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
4:AK:107:GLU:HA	4:AK:111:GLN:OE1	2.13	0.48
1:AN:8:ILE:CG2	2:AO:1:MET:HE1	2.43	0.48
2:AO:47:ASN:O	2:AO:51:ILE:HG13	2.13	0.48
1:AR:85:MET:HA	1:AR:133:MET:HE1	1.95	0.48
2:AU:57:ALA:CA	2:AU:61:LEU:HD12	2.42	0.48
1:AX:36:ARG:NH2	1:AX:152:TYR:OH	2.47	0.48
6:BD:55:GLU:O	6:BD:59:THR:HG23	2.14	0.48
6:BD:167:ILE:HD12	6:BD:219:ILE:HG22	1.95	0.48
6:BD:811:ARG:NH1	6:BD:812:ALA:O	2.47	0.48
10:BD:901:CYC:HMA1	10:BD:901:CYC:NB	2.29	0.48
9:BH:28:PHE:CD2	9:BH:39:LEU:HD23	2.47	0.48
2:BJ:52:VAL:O	2:BJ:56:VAL:HG23	2.13	0.48
2:BJ:71:ASN:HB3	10:BJ:200:CYC:OC	2.12	0.48
2:BJ:130:ILE:O	2:BJ:133:ILE:HG22	2.13	0.48
1:BK:121:THR:HG23	10:BK:200:CYC:C1C	2.43	0.48
2:BL:71:ASN:CG	10:BL:200:CYC:HMD2	2.39	0.48
10:BQ:200:CYC:O1A	6:CM:358:ARG:NH2	2.39	0.48
2:BT:18:TYR:HD2	6:CM:61:THR:HG22	1.73	0.48
10:BT:200:CYC:HAA1	6:CM:424:PHE:CZ	2.37	0.48
1:BV:8:ILE:HG23	2:BW:94:TYR:CD1	2.49	0.48
2:BY:122:PRO:HB2	2:BY:125:SER:HB2	1.95	0.48
10:CD:200:CYC:HB	10:CD:200:CYC:CMA	2.23	0.48
2:CF:78:TYR:CE1	1:CG:115:MET:HG3	2.47	0.48
2:CH:44:ILE:HD13	2:CH:149:MET:HE2	1.94	0.48
2:CH:68:PRO:HG3	2:CH:73:TYR:CE1	2.49	0.48
6:CM:77:THR:CG2	6:CM:138:PRO:HB2	2.44	0.48
6:CM:399:ARG:HA	6:CM:403:VAL:HG11	1.95	0.48
6:CM:751:LEU:HD12	6:CM:751:LEU:O	2.14	0.48
6:CM:829:GLN:HB2	6:CM:833:ALA:HB3	1.95	0.48
9:CQ:190:ILE:HD12	9:CQ:253:GLU:C	2.38	0.48
2:CS:111:GLY:O	2:CS:114:GLU:HG3	2.14	0.48
2:CW:2:GLN:HB2	2:CW:100:ASP:OD2	2.13	0.48
2:CW:50:THR:HG21	9:DI:62:GLN:HE21	1.79	0.48
2:CW:76:ARG:NH2	5:DF:62:THR:O	2.46	0.48
2:CZ:60:LEU:HD13	2:CZ:72:MET:HE1	1.95	0.48
2:CZ:111:GLY:O	2:CZ:115:THR:HG22	2.14	0.48
2:CZ:112:LEU:HD11	10:CZ:200:CYC:HMB1	1.95	0.48
1:DA:51:VAL:CG2	1:DA:133:MET:HE1	2.39	0.48
2:DB:68:PRO:HA	2:DB:73:TYR:CD1	2.49	0.48
9:DI:262:ASP:O	9:DI:294:PRO:HG3	2.13	0.48
1:DL:159:LYS:HE2	1:DL:159:LYS:HA	1.95	0.48

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:DN:37:LEU:CD2	2:DO:28:LYS:HE3	2.43	0.48
2:DR:2:GLN:OE1	2:DR:7:ALA:HB2	2.13	0.48
2:DR:81:CYS:O	2:DR:85:LEU:HD23	2.12	0.48
1:DS:88:TYR:OH	1:DS:160:MET:HE1	2.13	0.48
1:DS:96:VAL:HA	1:DS:152:TYR:CE2	2.49	0.48
1:DS:122:PRO:HD2	10:DS:200:CYC:OC	2.13	0.48
5:DX:6:ILE:O	5:DX:28:THR:HA	2.13	0.48
9:EA:21:VAL:HG13	9:EA:149:GLN:HG3	1.95	0.48
9:EA:25:ILE:CD1	9:EA:156:ASN:HB2	2.44	0.48
9:EA:185:ARG:NH1	9:EA:207:ALA:HB1	2.28	0.48
9:EA:206:ASN:OD1	9:EA:250:LEU:N	2.31	0.48
1:AA:84:ASP:OD2	10:AA:200:CYC:ND	2.40	0.48
1:AC:33:GLY:O	1:AC:36:ARG:HG2	2.13	0.48
1:AC:65:ILE:HD12	10:AC:200:CYC:OC	2.13	0.48
1:AN:33:GLY:HA2	1:AN:36:ARG:CG	2.44	0.48
1:AN:57:ARG:O	1:AN:60:GLN:HG3	2.13	0.48
2:AY:137:THR:O	2:AY:141:VAL:HG22	2.13	0.48
6:BD:803:MET:HE3	6:BD:807:HIS:CE1	2.49	0.48
9:BH:273:VAL:HB	9:BH:286:ILE:CD1	2.42	0.48
10:BS:200:CYC:HBA2	6:CM:401:LEU:HD11	1.95	0.48
1:BV:115:MET:HG2	2:CA:78:TYR:CD1	2.48	0.48
1:CC:75:GLU:OE2	1:CC:76:GLU:HG2	2.13	0.48
1:CG:116:TYR:HD2	1:CG:123:ILE:HD12	1.78	0.48
6:CM:5:ALA:HB3	6:CM:437:TYR:HD1	1.78	0.48
6:CM:244:PRO:HD2	6:CM:248:ILE:HB	1.96	0.48
6:CM:519:LEU:HD13	6:CM:569:GLU:O	2.13	0.48
6:CM:872:LEU:CG	6:CM:884:VAL:HG11	2.39	0.48
9:CQ:109:PHE:CZ	9:CQ:113:LEU:HD11	2.48	0.48
9:CQ:204:ASN:HD22	9:CQ:213:LEU:HA	1.79	0.48
9:CQ:249:LYS:O	9:CQ:250:LEU:HD23	2.14	0.48
2:CS:47:ASN:HB2	2:CS:51:ILE:CD1	2.42	0.48
1:CT:128:GLN:O	1:CT:132:GLU:HG2	2.14	0.48
1:CY:87:TYR:O	1:CY:91:LEU:HG	2.14	0.48
2:DB:122:PRO:HD2	10:DB:200:CYC:HMC3	1.95	0.48
9:DI:146:GLU:H	9:DI:149:GLN:NE2	2.12	0.48
2:DM:22:ALA:O	2:DM:26:LYS:HG3	2.13	0.48
2:DO:124:SER:O	2:DO:128:GLN:HG3	2.13	0.48
1:DQ:21:GLY:O	1:DQ:25:ARG:HG2	2.13	0.48
1:DQ:104:ILE:H	1:DQ:104:ILE:HD12	1.77	0.48
2:AQ:62:TYR:CE2	9:BH:11:ILE:HA	2.48	0.48
2:AS:71:ASN:O	2:AS:77:ARG:HD3	2.13	0.48

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:AV:102:THR:O	1:AV:106:GLU:HG2	2.13	0.48
1:AV:105:GLU:OE2	1:AV:159:LYS:HE2	2.13	0.48
2:AW:125:SER:HA	2:AW:128:GLN:OE1	2.12	0.48
1:AX:115:MET:HE3	1:AX:116:TYR:CE1	2.49	0.48
2:AY:6:THR:HG23	6:BD:532:VAL:CG1	2.44	0.48
5:BB:3:MET:N	5:BB:57:THR:HG1	2.10	0.48
9:BH:27:ARG:O	9:BH:31:LEU:HG	2.13	0.48
1:BK:63:PRO:O	1:BK:67:SER:N	2.40	0.48
3:BM:4:VAL:HG22	3:BM:26:ILE:HG23	1.96	0.48
2:BN:1:MET:SD	2:BN:1:MET:N	2.81	0.48
2:BN:39:ARG:O	2:BN:43:VAL:HG23	2.13	0.48
1:BP:91:LEU:HB3	1:BP:104:ILE:HG23	1.96	0.48
10:BP:200:CYC:CGA	2:BT:66:THR:HG21	2.43	0.48
1:BV:27:LYS:O	1:BV:30:VAL:HG22	2.14	0.48
2:BW:3:ASP:HA	2:BW:98:ALA:HB1	1.95	0.48
2:BW:52:VAL:O	2:BW:56:VAL:HG23	2.14	0.48
2:CA:76:ARG:HH21	6:CM:478:LYS:HD3	1.78	0.48
2:CD:52:VAL:O	2:CD:56:VAL:HG22	2.13	0.48
2:CF:14:VAL:HG12	5:CK:63:ASN:ND2	2.27	0.48
2:CF:21:GLY:CA	2:CF:24:MET:HB3	2.43	0.48
2:CF:55:ALA:O	2:CF:59:SER:OG	2.13	0.48
1:CG:81:CYS:HA	10:CG:200:CYC:CHD	2.43	0.48
6:CM:609:LEU:C	6:CM:610:LEU:HD12	2.39	0.48
6:CM:651:GLU:OE1	6:CM:651:GLU:N	2.46	0.48
6:CM:756:ILE:HB	1:DQ:159:LYS:HZ1	1.78	0.48
1:CR:11:ALA:HA	1:CR:16:ARG:CZ	2.43	0.48
2:CS:112:LEU:HG	2:CS:160:LEU:HD21	1.94	0.48
1:CT:33:GLY:HA3	2:CU:31:PHE:CE2	2.49	0.48
2:CW:51:ILE:HG12	2:CW:136:VAL:HG12	1.96	0.48
2:CW:71:ASN:ND2	2:CW:121:VAL:HA	2.29	0.48
2:CZ:75:THR:HG22	10:DA:200:CYC:HMB2	1.96	0.48
1:DA:16:ARG:O	2:DB:94:TYR:OH	2.31	0.48
2:DD:68:PRO:HG3	2:DD:73:TYR:CZ	2.49	0.48
2:DD:144:ASP:OD1	2:DD:145:ALA:N	2.44	0.48
10:DD:200:CYC:HMB3	10:DD:200:CYC:HBB3	1.95	0.48
8:DH:221:MET:HE3	1:DS:52:LYS:HB2	1.94	0.48
8:DH:237:ALA:O	1:DJ:52:LYS:HE2	2.12	0.48
1:DJ:88:TYR:CE2	1:DJ:160:MET:HE1	2.44	0.48
2:DK:19:LEU:HD12	2:DK:23:ALA:HB1	1.96	0.48
2:DK:122:PRO:HG2	2:DK:125:SER:HB2	1.94	0.48
2:DO:84:ASP:OD2	2:DO:116:TYR:OH	2.16	0.48

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:DQ:39:ILE:HD11	1:DQ:145:ASP:CG	2.38	0.48
2:DR:73:TYR:O	2:DR:74:THR:OG1	2.24	0.48
2:DR:85:LEU:HD12	2:DR:133:ILE:HD13	1.95	0.48
9:EA:30:GLN:O	9:EA:171:ARG:NH1	2.47	0.48
9:EA:286:ILE:HD11	9:EA:303:ILE:HG23	1.95	0.48
2:AB:71:ASN:HB3	10:AB:200:CYC:OC	2.14	0.48
2:AB:104:LEU:HD11	2:AB:152:TYR:HB3	1.96	0.48
2:AF:71:ASN:HD22	2:AF:122:PRO:HD3	1.77	0.48
1:AH:38:ARG:HA	1:AH:41:GLU:OE1	2.13	0.48
1:AP:105:GLU:OE2	1:AP:159:LYS:NZ	2.46	0.48
10:AQ:200:CYC:CMA	10:AQ:200:CYC:HB	2.26	0.48
1:AV:100:ASP:OD1	1:AV:101:VAL:N	2.46	0.48
2:AY:61:LEU:HD22	10:AZ:200:CYC:HAA1	1.96	0.48
6:BD:543:ILE:HG23	6:BD:578:PHE:CE2	2.49	0.48
6:BD:555:LEU:HB3	6:BD:559:GLN:HB2	1.95	0.48
6:BD:755:ILE:HD12	6:BD:757:ASN:C	2.39	0.48
9:BH:98:TYR:O	9:BH:106:LYS:HE3	2.14	0.48
2:BL:100:ASP:OD2	2:BL:102:SER:HB3	2.14	0.48
2:BN:76:ARG:CZ	5:CJ:66:LEU:HD11	2.44	0.48
2:BQ:52:VAL:O	2:BQ:56:VAL:HG23	2.13	0.48
2:BQ:71:ASN:HD22	2:BQ:121:VAL:HA	1.78	0.48
4:BS:66:LEU:HD11	4:BS:126:PRO:HB3	1.96	0.48
1:CC:24:ASP:HA	1:CC:27:LYS:HG2	1.96	0.48
6:CM:526:LYS:HD3	6:CM:533:LYS:HE2	1.96	0.48
6:CM:856:ARG:NH2	6:CM:870:GLU:OE1	2.46	0.48
9:CQ:187:LYS:HD3	9:CQ:188:VAL:O	2.13	0.48
1:CR:113:ARG:HH11	1:CR:123:ILE:HD12	1.77	0.48
1:CT:92:VAL:HA	1:CT:104:ILE:HD11	1.96	0.48
2:CU:2:GLN:O	2:CU:100:ASP:HB3	2.14	0.48
2:CW:10:ASN:O	2:CW:14:VAL:HG13	2.13	0.48
2:CW:112:LEU:CD2	2:CW:160:LEU:HD13	2.43	0.48
1:CY:7:SER:HB3	1:CY:18:LEU:HD11	1.95	0.48
2:DB:9:ILE:HD13	2:DB:19:LEU:HD21	1.96	0.48
2:DD:102:SER:O	2:DD:106:GLU:HG2	2.13	0.48
9:DI:250:LEU:O	9:DI:251:ILE:HD13	2.13	0.48
1:DJ:152:TYR:HE1	1:DQ:20:PRO:HB2	1.79	0.48
2:DM:36:LEU:HA	2:DM:39:ARG:HG2	1.95	0.48
1:DS:100:ASP:OD2	1:DS:102:THR:HG23	2.13	0.48
1:DU:107:ILE:HG12	2:DV:13:ASP:OD2	2.14	0.48
9:EA:43:ALA:O	9:EA:51:LEU:HD12	2.12	0.48
1:AC:29:PHE:CE1	1:AC:99:GLY:HA3	2.45	0.48

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:AO:123:ILE:O	2:AO:127:VAL:HG23	2.14	0.48
1:AV:8:ILE:HD12	2:AW:94:TYR:CD1	2.49	0.48
1:AZ:39:ILE:HD11	1:AZ:145:ASP:OD1	2.14	0.48
5:BA:10:VAL:HG12	5:BA:27:PHE:HE2	1.78	0.48
6:BD:86:GLU:OE1	6:BD:154:ARG:HB2	2.13	0.48
6:BD:776:GLU:OE2	5:DF:29:LYS:HD3	2.13	0.48
6:BD:792:TYR:HA	6:BD:800:VAL:HG22	1.94	0.48
1:BK:94:TYR:OH	2:BL:17:LYS:O	2.31	0.48
3:BM:141:LEU:HD13	3:BM:145:ASP:OD2	2.14	0.48
1:BR:124:GLU:OE1	1:BR:124:GLU:N	2.39	0.48
1:BV:8:ILE:CB	2:BW:1:MET:HE1	2.31	0.48
2:BW:75:THR:HG23	10:BX:200:CYC:CAB	2.43	0.48
2:BY:111:GLY:HA2	2:BY:114:GLU:OE1	2.14	0.48
2:BY:116:TYR:CE2	2:BY:160:LEU:HD21	2.47	0.48
2:CA:76:ARG:NH2	6:CM:478:LYS:HD3	2.28	0.48
2:CA:107:ARG:HD3	6:CM:594:ALA:O	2.14	0.48
1:CE:85:MET:CE	1:CE:129:SER:HB2	2.41	0.48
2:CF:8:VAL:HG22	2:CF:26:LYS:CD	2.44	0.48
2:CF:61:LEU:HD22	10:CG:200:CYC:CAA	2.44	0.48
1:CG:101:VAL:HG12	1:CG:152:TYR:HD1	1.78	0.48
5:CK:3:MET:O	5:CK:55:LEU:HD12	2.14	0.48
6:CM:75:ILE:HG13	6:CM:198:ALA:CB	2.38	0.48
6:CM:131:PRO:CD	6:CM:200:GLN:HG2	2.43	0.48
6:CM:799:LYS:HB2	6:CM:883:LEU:CD2	2.43	0.48
6:CM:891:VAL:CG1	2:DT:106:GLU:HB3	2.44	0.48
9:CQ:131:LEU:HG	9:CQ:135:ALA:HB3	1.94	0.48
9:CQ:201:TYR:HA	9:CQ:213:LEU:CD1	2.44	0.48
2:CW:111:GLY:O	2:CW:115:THR:OG1	2.15	0.48
1:CY:66:VAL:HG12	1:CY:72:ALA:O	2.14	0.48
1:DC:6:LYS:NZ	1:DC:100:ASP:OD2	2.47	0.48
2:DD:10:ASN:O	2:DD:14:VAL:HG23	2.12	0.48
2:DD:67:ARG:HD3	8:DY:214:ASN:ND2	2.28	0.48
9:DI:213:LEU:HA	9:DI:216:LEU:HB3	1.96	0.48
1:DJ:57:ARG:NE	1:DJ:132:GLU:OE1	2.47	0.48
2:DK:23:ALA:HA	2:DK:26:LYS:HG2	1.96	0.48
1:DL:85:MET:HE2	1:DL:133:MET:SD	2.54	0.48
2:DR:47:ASN:O	2:DR:51:ILE:HD12	2.13	0.48
1:DU:14:GLU:HB2	1:DU:16:ARG:HG2	1.96	0.48
1:DU:49:THR:HG22	1:DU:53:GLN:OE1	2.14	0.48
1:DU:66:VAL:O	1:DU:73:TYR:HA	2.14	0.48
1:AA:85:MET:HE2	10:AA:200:CYC:CBC	2.44	0.48

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:AD:87:TYR:CD2	5:BA:22:LEU:HD22	2.49	0.48
1:AH:37:LEU:HD22	2:AI:24:MET:SD	2.53	0.48
1:AN:4:VAL:O	1:AN:8:ILE:HG12	2.13	0.48
2:AO:63:SER:H	2:AO:66:THR:HG22	1.78	0.48
2:AQ:10:ASN:O	2:AQ:14:VAL:HG23	2.14	0.48
1:AZ:128:GLN:NE2	1:AZ:132:GLU:OE2	2.43	0.48
5:BA:10:VAL:HG13	5:BA:10:VAL:O	2.14	0.48
6:BD:866:PHE:HD2	2:DB:115:THR:HG23	1.79	0.48
9:BH:48:GLY:HA2	9:BH:53:ILE:HG22	1.96	0.48
9:BH:168:ASP:OD1	9:BH:168:ASP:N	2.46	0.48
9:BH:290:PHE:CE1	9:BH:301:VAL:HG23	2.49	0.48
2:BJ:1:MET:N	5:CJ:67:ALA:OXT	2.46	0.48
1:BK:16:ARG:NH1	1:BK:17:TYR:O	2.47	0.48
1:BK:126:VAL:O	1:BK:130:VAL:HG23	2.13	0.48
3:BM:58:LEU:HD13	3:BM:129:ALA:CA	2.44	0.48
2:BN:1:MET:HE3	2:BN:103:ILE:HD12	1.96	0.48
2:BN:76:ARG:NH1	10:BN:200:CYC:O2D	2.46	0.48
2:BQ:68:PRO:O	2:BW:124:SER:HB2	2.14	0.48
4:BS:63:ILE:CG2	4:BS:66:LEU:HG	2.43	0.48
1:CC:116:TYR:CE2	1:CC:126:VAL:HG11	2.48	0.48
2:CD:54:GLU:CD	2:CD:136:VAL:HG21	2.39	0.48
2:CD:109:LEU:CD2	2:CD:159:GLY:HA3	2.44	0.48
1:CE:37:LEU:HD21	2:CF:24:MET:SD	2.54	0.48
2:CH:132:ALA:HA	2:CH:135:GLU:OE1	2.14	0.48
6:CM:553:ARG:HG3	6:CM:554:ASP:O	2.13	0.48
6:CM:603:GLU:O	6:CM:607:ARG:HG3	2.14	0.48
9:CQ:187:LYS:HG2	9:CQ:199:LEU:HD22	1.95	0.48
1:CR:27:LYS:NZ	2:CS:38:VAL:HB	2.29	0.48
2:CS:62:TYR:OH	10:CT:200:CYC:HAD2	2.14	0.48
1:CY:107:ILE:O	2:DD:74:THR:HB	2.14	0.48
2:CZ:81:CYS:O	2:CZ:85:LEU:HG	2.14	0.48
2:CZ:133:ILE:O	2:CZ:137:THR:HG23	2.14	0.48
2:DB:76:ARG:NH2	1:DC:106:GLU:OE1	2.47	0.48
2:DD:122:PRO:O	2:DD:126:THR:HG23	2.13	0.48
1:DJ:5:THR:O	1:DJ:9:VAL:HG23	2.13	0.48
2:DK:121:VAL:HG13	2:DK:122:PRO:HD2	1.96	0.48
2:DM:15:GLN:HG2	2:DM:17:LYS:HG2	1.96	0.48
1:DN:131:ARG:O	1:DN:134:LYS:HB3	2.13	0.48
2:DO:103:ILE:HG13	2:DO:107:ARG:HD3	1.96	0.48
1:DS:97:VAL:CG2	2:DT:9:ILE:HD11	2.44	0.48
2:DT:40:ALA:O	2:DT:44:ILE:HG13	2.13	0.48

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:DT:132:ALA:O	2:DT:136:VAL:HG23	2.13	0.48
2:DV:102:SER:HA	2:DV:105:ASP:OD2	2.13	0.48
5:DX:5:ARG:HB3	5:DX:30:LEU:CD2	2.43	0.48
9:EA:204:ASN:HB3	9:EA:209:ASP:HB3	1.96	0.48
1:AA:157:ILE:HG13	1:AA:158:GLY:N	2.28	0.48
2:AD:133:ILE:O	2:AD:137:THR:HG22	2.13	0.48
3:AE:141:LEU:HD13	3:AE:145:ASP:OD2	2.14	0.48
1:AH:30:VAL:HG11	2:AI:34:GLY:HA3	1.96	0.48
1:AH:129:SER:O	1:AH:133:MET:HG3	2.14	0.48
1:AJ:108:GLY:O	1:AJ:109:LEU:HD23	2.14	0.48
1:AN:105:GLU:HA	1:AN:109:LEU:HD12	1.96	0.48
1:AR:12:ASP:OD2	2:AS:91:TYR:OH	2.19	0.48
1:AR:77:MET:HE2	9:BH:11:ILE:CD1	2.44	0.48
2:AU:75:THR:CG2	1:AV:108:GLY:HA2	2.42	0.48
2:AU:134:LYS:HG3	2:AU:150:GLY:HA2	1.96	0.48
1:AV:3:ILE:HD12	1:AV:6:LYS:HE3	1.96	0.48
1:AV:97:VAL:HG11	2:AW:19:LEU:HD11	1.96	0.48
2:AY:132:ALA:O	2:AY:136:VAL:HG23	2.13	0.48
6:BD:788:MET:HB3	6:BD:831:LEU:HD23	1.95	0.48
1:BI:130:VAL:HG12	1:BI:157:ILE:HD11	1.95	0.48
10:BK:200:CYC:HBC2	10:BK:200:CYC:H2C	1.75	0.48
3:BM:4:VAL:O	3:BM:8:ILE:HG12	2.14	0.48
2:BN:71:ASN:O	2:BN:77:ARG:HD2	2.14	0.48
1:BR:17:TYR:OH	4:BS:90:LEU:HD23	2.14	0.48
4:BS:86:MET:HG2	4:BS:134:MET:SD	2.54	0.48
4:BS:105:LEU:O	4:BS:109:VAL:HB	2.14	0.48
2:BY:88:TYR:OH	2:BY:112:LEU:HD21	2.14	0.48
1:BZ:39:ILE:O	1:BZ:43:LEU:HG	2.13	0.48
2:CD:39:ARG:O	2:CD:43:VAL:HG23	2.14	0.48
2:CD:78:TYR:O	2:CD:82:ILE:HG22	2.14	0.48
2:CF:125:SER:HA	2:CF:128:GLN:OE1	2.14	0.48
1:CG:109:LEU:HD11	1:CG:159:LYS:HB3	1.94	0.48
2:CH:132:ALA:O	2:CH:136:VAL:HG23	2.13	0.48
5:CJ:15:ARG:NH2	5:CJ:23:GLN:HB2	2.29	0.48
6:CM:76:PHE:CD2	6:CM:149:MET:HE1	2.49	0.48
6:CM:133:PRO:HB3	6:CM:136:PHE:CE1	2.49	0.48
6:CM:278:LYS:CD	6:CM:310:ARG:HG3	2.42	0.48
6:CM:335:GLU:HB3	6:CM:336:PRO:HD3	1.96	0.48
6:CM:348:ARG:HB3	6:CM:417:LEU:HD21	1.95	0.48
1:CR:75:GLU:OE2	9:DI:152:THR:HG21	2.13	0.48
1:CR:107:ILE:HD13	2:CS:13:ASP:OD2	2.14	0.48

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:CS:106:GLU:HG3	5:DF:57:THR:CG2	2.44	0.48
2:CU:52:VAL:HG21	2:CU:86:ASP:OD1	2.14	0.48
2:CW:57:ALA:CA	2:CW:61:LEU:HD12	2.44	0.48
1:DJ:39:ILE:HG23	1:DJ:141:MET:SD	2.54	0.48
1:DJ:104:ILE:HG21	1:DJ:156:VAL:HG22	1.95	0.48
2:DK:104:LEU:O	2:DK:108:VAL:HG12	2.14	0.48
2:DK:127:VAL:O	2:DK:131:GLN:HG2	2.12	0.48
2:DO:50:THR:HG21	9:EA:62:GLN:NE2	2.29	0.48
2:DO:88:TYR:CD2	2:DO:130:ILE:HD11	2.48	0.48
2:DO:114:GLU:CG	5:DX:53:VAL:HG12	2.44	0.48
2:DR:112:LEU:HA	2:DR:115:THR:CG2	2.44	0.48
2:DV:2:GLN:OE1	2:DV:2:GLN:N	2.47	0.48
9:EA:44:TYR:HA	9:EA:47:MET:HG3	1.96	0.48
1:AA:103:PRO:HA	1:AA:106:GLU:HG2	1.96	0.47
2:AB:88:TYR:OH	2:AB:112:LEU:HD21	2.13	0.47
2:AF:123:ILE:HD12	2:AF:160:LEU:HD21	1.96	0.47
1:AN:39:ILE:HG23	1:AN:141:MET:CE	2.43	0.47
1:AN:115:MET:HE1	10:AN:200:CYC:CHB	2.43	0.47
2:AO:114:GLU:HG3	6:BD:443:ASP:HB3	1.96	0.47
1:AP:67:SER:O	1:AP:73:TYR:HB2	2.13	0.47
2:AS:144:ASP:OD1	2:AS:145:ALA:N	2.46	0.47
1:AV:11:ALA:CB	1:AV:18:LEU:HD23	2.44	0.47
1:AZ:72:ALA:HA	1:AZ:77:MET:HB3	1.95	0.47
6:BD:291:GLU:HG2	6:BD:348:ARG:NH1	2.29	0.47
6:BD:743:TYR:HE1	6:BD:781:LEU:HD21	1.78	0.47
9:BH:215:GLU:OE1	9:BH:215:GLU:N	2.46	0.47
10:BI:200:CYC:O2A	2:BN:62:TYR:N	2.32	0.47
2:BL:54:GLU:O	2:BL:58:LYS:HG2	2.13	0.47
3:BM:101:LYS:HE3	1:BR:16:ARG:CD	2.44	0.47
1:CE:30:VAL:HG23	2:CF:31:PHE:HA	1.96	0.47
2:CF:137:THR:O	2:CF:141:VAL:HG22	2.13	0.47
6:CM:77:THR:O	6:CM:141:ILE:HD12	2.12	0.47
6:CM:744:ARG:NH2	1:DU:13:ALA:O	2.45	0.47
6:CM:786:LEU:O	6:CM:790:GLU:HG2	2.13	0.47
9:CQ:89:ARG:HD3	9:CQ:170:LYS:HG2	1.96	0.47
9:CQ:233:GLY:O	9:CQ:237:VAL:HG23	2.13	0.47
1:CR:69:GLY:HA3	9:DI:55:ALA:HB3	1.96	0.47
1:CR:80:THR:HG21	10:CR:200:CYC:HAD1	1.95	0.47
2:CS:104:LEU:HD21	2:CS:156:ILE:HD12	1.95	0.47
2:CU:47:ASN:CG	2:CU:140:LEU:HD21	2.39	0.47
1:CV:3:ILE:CD1	1:CV:25:ARG:HE	2.27	0.47

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:CV:98:SER:OG	1:CV:100:ASP:OD1	2.24	0.47
2:CW:85:LEU:CG	10:CW:200:CYC:HBC1	2.42	0.47
2:CW:101:ALA:HB1	2:CW:155:TYR:CE2	2.49	0.47
1:DC:67:SER:O	1:DC:73:TYR:HB2	2.13	0.47
2:DD:116:TYR:CE2	2:DD:160:LEU:HD21	2.49	0.47
5:DF:5:ARG:HA	5:DF:29:LYS:O	2.13	0.47
5:DF:23:GLN:OE1	5:DF:23:GLN:N	2.42	0.47
9:DI:21:VAL:O	9:DI:25:ILE:HG12	2.14	0.47
9:DI:28:PHE:CE1	9:DI:157:ALA:HB1	2.48	0.47
1:DJ:127:ALA:HA	1:DJ:130:VAL:HG12	1.95	0.47
2:DK:43:VAL:CG1	2:DK:141:VAL:HG12	2.44	0.47
1:DL:103:PRO:O	1:DL:107:ILE:HG12	2.13	0.47
2:DM:137:THR:O	2:DM:141:VAL:HG22	2.13	0.47
1:DQ:115:MET:SD	2:DV:78:TYR:HB3	2.54	0.47
1:DQ:122:PRO:HG2	1:DQ:125:ALA:CB	2.44	0.47
2:DT:54:GLU:O	2:DT:58:LYS:HG2	2.13	0.47
9:EA:293:ASN:HB2	9:EA:296:GLY:CA	2.43	0.47
1:AA:153:PHE:O	1:AA:157:ILE:HG23	2.15	0.47
2:AB:83:ARG:HG2	2:AB:87:TYR:CE2	2.49	0.47
1:AC:3:ILE:HG22	1:AC:26:ILE:HD13	1.96	0.47
2:AF:122:PRO:HB2	2:AF:125:SER:CB	2.42	0.47
4:AK:114:ARG:HE	4:AK:124:ILE:HD13	1.80	0.47
1:AN:101:VAL:HG12	1:AN:152:TYR:HD1	1.79	0.47
1:AP:81:CYS:HA	10:AP:200:CYC:CHD	2.43	0.47
1:AP:141:MET:HG3	1:AP:146:ALA:HB2	1.96	0.47
2:AU:109:LEU:HD22	2:AU:159:GLY:HA3	1.97	0.47
2:AW:71:ASN:O	2:AW:77:ARG:HD2	2.13	0.47
2:AY:112:LEU:CD1	10:AY:200:CYC:HMB3	2.44	0.47
1:AZ:122:PRO:O	1:AZ:126:VAL:HG23	2.14	0.47
6:BD:80:SER:HB3	6:BD:83:SER:OG	2.14	0.47
6:BD:180:ARG:HB2	6:BD:230:GLU:OE1	2.14	0.47
6:BD:355:PRO:HD2	6:BD:430:PHE:CE1	2.49	0.47
6:BD:705:GLU:HB2	2:DT:154:ASP:OD2	2.14	0.47
8:BG:216:MET:CE	9:BH:12:PHE:HB2	2.44	0.47
9:BH:204:ASN:CB	9:BH:213:LEU:HB2	2.40	0.47
9:BH:290:PHE:HD2	9:BH:292:LEU:HD21	1.78	0.47
2:BJ:77:ARG:CB	10:BJ:200:CYC:HMD1	2.44	0.47
2:BJ:85:LEU:HD21	10:BJ:200:CYC:HBC1	1.96	0.47
1:BK:56:ASP:OD1	1:BK:57:ARG:N	2.47	0.47
2:BN:51:ILE:HD11	2:BN:140:LEU:HD23	1.95	0.47
1:BR:4:VAL:HG11	4:BS:99:ALA:HA	1.96	0.47

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:BX:122:PRO:O	1:BX:126:VAL:HG23	2.14	0.47
2:BY:110:ASN:ND2	6:CM:668:ARG:HG2	2.29	0.47
1:CC:3:ILE:HG12	1:CC:25:ARG:HG3	1.95	0.47
1:CE:134:LYS:HD3	1:CE:153:PHE:HB3	1.96	0.47
1:CG:102:THR:O	1:CG:106:GLU:HG2	2.14	0.47
2:CH:106:GLU:HG3	2:CH:107:ARG:HG3	1.94	0.47
9:CQ:156:ASN:O	9:CQ:159:VAL:HG12	2.13	0.47
9:CQ:214:ILE:O	9:CQ:217:PHE:HB2	2.14	0.47
1:CR:154:ASP:CG	2:CZ:46:ALA:HB2	2.39	0.47
1:CT:18:LEU:HD13	2:CU:97:LEU:CD2	2.43	0.47
1:CT:88:TYR:CZ	1:CT:160:MET:HE1	2.49	0.47
1:CT:141:MET:SD	1:CT:146:ALA:HA	2.54	0.47
1:CV:84:ASP:OD2	10:CV:200:CYC:ND	2.45	0.47
2:CW:52:VAL:HG21	2:CW:86:ASP:OD1	2.14	0.47
2:CW:121:VAL:HG13	2:CW:122:PRO:HD2	1.96	0.47
1:CY:66:VAL:HG12	1:CY:78:THR:HG22	1.95	0.47
1:DC:104:ILE:HG21	1:DC:156:VAL:HG22	1.96	0.47
5:DF:6:ILE:HG21	5:DF:36:TRP:CH2	2.49	0.47
9:DI:264:PHE:CE2	9:DI:293:ASN:HA	2.50	0.47
1:DJ:4:VAL:HG12	1:DJ:26:ILE:CD1	2.40	0.47
1:DL:87:TYR:HB3	10:DL:200:CYC:HAB2	1.97	0.47
1:DL:105:GLU:O	1:DL:110:VAL:HG23	2.14	0.47
1:DS:4:VAL:HG22	1:DS:26:ILE:HD12	1.96	0.47
2:DV:2:GLN:HB2	2:DV:6:THR:OG1	2.14	0.47
2:DV:68:PRO:HA	2:DV:73:TYR:CG	2.49	0.47
1:AA:95:GLY:HA3	1:AA:104:ILE:HD11	1.95	0.47
2:AD:2:GLN:HB2	2:AD:6:THR:HB	1.96	0.47
1:AJ:61:LYS:HD2	1:AJ:132:GLU:OE2	2.15	0.47
1:AN:14:GLU:OE1	1:AN:16:ARG:NE	2.34	0.47
1:AP:68:PRO:HG2	1:BR:66:VAL:HG11	1.96	0.47
2:AU:50:THR:O	2:AU:54:GLU:HG3	2.14	0.47
1:AV:84:ASP:OD2	10:AV:200:CYC:ND	2.43	0.47
2:AY:105:ASP:O	6:BD:523:LYS:NZ	2.43	0.47
1:AZ:29:PHE:O	1:AZ:36:ARG:NH1	2.47	0.47
6:BD:31:GLN:HG2	6:BD:267:TYR:OH	2.13	0.47
6:BD:85:LEU:HA	6:BD:153:LEU:CD1	2.44	0.47
6:BD:482:ARG:HG3	6:BD:615:TYR:O	2.14	0.47
10:BD:901:CYC:HMA1	10:BD:901:CYC:HB	1.79	0.47
7:BF:109:ARG:HG3	7:BF:110:PHE:N	2.28	0.47
2:BN:77:ARG:CB	10:BN:200:CYC:HMD1	2.44	0.47
1:BR:39:ILE:O	1:BR:42:THR:OG1	2.30	0.47

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:BX:107:ILE:HD13	2:BY:13:ASP:OD2	2.15	0.47
2:CA:63:SER:H	2:CA:66:THR:CG2	2.26	0.47
2:CD:23:ALA:O	2:CD:27:LEU:HG	2.15	0.47
2:CH:108:VAL:HG12	2:CH:109:LEU:CD2	2.44	0.47
6:CM:70:ARG:HH22	6:CM:205:ALA:HB1	1.79	0.47
9:CQ:291:LEU:HD23	9:CQ:299:PHE:CE2	2.49	0.47
1:CT:63:PRO:O	1:CT:66:VAL:HG12	2.14	0.47
10:CT:200:CYC:HMB1	10:CT:200:CYC:HBB3	1.96	0.47
2:CW:126:THR:CG2	2:CW:160:LEU:HD21	2.43	0.47
1:CY:72:ALA:HA	1:CY:77:MET:CB	2.45	0.47
1:CY:85:MET:HE3	1:CY:129:SER:CB	2.41	0.47
1:CY:131:ARG:O	1:CY:135:GLU:HG3	2.14	0.47
2:DD:108:VAL:HG22	10:DD:200:CYC:HBB1	1.95	0.47
5:DF:59:ARG:HB3	5:DF:62:THR:CG2	2.44	0.47
9:DI:42:PHE:CD2	9:DI:131:LEU:HD13	2.47	0.47
9:DI:167:LYS:HE2	9:DI:167:LYS:HA	1.96	0.47
9:DI:285:ASN:CG	9:DI:306:LEU:HB2	2.39	0.47
1:DL:37:LEU:O	1:DL:40:ALA:HB3	2.15	0.47
1:DQ:88:TYR:CD1	1:DQ:91:LEU:HD12	2.50	0.47
1:DS:96:VAL:HA	1:DS:152:TYR:HE2	1.78	0.47
2:DT:51:ILE:HG22	2:DT:133:ILE:HD13	1.96	0.47
2:DT:134:LYS:HG3	2:DT:150:GLY:HA2	1.95	0.47
1:DU:80:THR:CG2	1:DU:83:ARG:HH21	2.28	0.47
1:DU:85:MET:HA	1:DU:133:MET:HE1	1.96	0.47
9:EA:19:ASP:O	9:EA:22:PRO:HD2	2.15	0.47
2:AD:57:ALA:HA	2:AD:61:LEU:CG	2.44	0.47
3:AE:92:VAL:HA	3:AE:104:ILE:HD11	1.95	0.47
2:AF:63:SER:H	2:AF:66:THR:HG22	1.79	0.47
2:AI:68:PRO:HG3	2:AI:73:TYR:CE1	2.50	0.47
2:AI:90:ARG:O	2:AI:93:THR:OG1	2.30	0.47
1:AJ:38:ARG:O	1:AJ:42:THR:HG23	2.14	0.47
2:AO:53:LYS:HD2	1:AP:118:SER:O	2.14	0.47
2:AS:143:ALA:O	2:AS:147:LYS:HG2	2.13	0.47
1:AV:54:ALA:HB1	1:AV:133:MET:HG2	1.95	0.47
1:AX:51:VAL:HG22	1:AX:133:MET:CE	2.42	0.47
6:BD:273:LEU:O	6:BD:278:LYS:NZ	2.31	0.47
6:BD:776:GLU:N	6:BD:776:GLU:OE1	2.47	0.47
6:BD:845:LEU:CG	5:DF:42:ARG:HD2	2.39	0.47
1:BK:126:VAL:HG23	1:BK:160:MET:HE2	1.96	0.47
1:CC:17:TYR:OH	2:CD:89:LEU:HD22	2.14	0.47
2:CD:65:VAL:HG12	2:CD:72:MET:CB	2.45	0.47

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:CH:52:VAL:O	2:CH:56:VAL:HG22	2.14	0.47
2:CH:112:LEU:HA	2:CH:115:THR:HG22	1.96	0.47
5:CJ:10:VAL:HG12	5:CJ:27:PHE:HE2	1.79	0.47
6:CM:3:VAL:HG12	6:CM:5:ALA:H	1.78	0.47
9:CQ:125:ILE:HG13	11:CQ:400:45D:H191	1.95	0.47
9:CQ:187:LYS:HE2	9:CQ:199:LEU:CD2	2.44	0.47
9:CQ:224:GLN:HB3	9:CQ:302:ALA:HA	1.97	0.47
2:CU:5:ILE:HG23	2:CU:27:LEU:HD11	1.96	0.47
1:CY:103:PRO:O	1:CY:107:ILE:HG12	2.14	0.47
1:CY:115:MET:HE1	2:DD:79:ALA:HA	1.95	0.47
2:CZ:19:LEU:HB3	2:CZ:24:MET:SD	2.55	0.47
2:CZ:51:ILE:HG13	2:CZ:136:VAL:HG12	1.97	0.47
1:DA:35:ALA:O	1:DA:39:ILE:HG12	2.14	0.47
1:DC:108:GLY:O	1:DC:109:LEU:HD23	2.14	0.47
2:DD:119:LEU:HD13	10:DD:200:CYC:HBD1	1.96	0.47
8:DG:247:PRO:HB2	2:DK:62:TYR:CG	2.49	0.47
9:DI:170:LYS:NZ	9:DI:173:ALA:HB2	2.29	0.47
9:DI:201:TYR:O	9:DI:205:LEU:HD13	2.14	0.47
1:DJ:46:SER:O	1:DJ:50:ILE:HD12	2.15	0.47
2:DK:52:VAL:O	2:DK:56:VAL:HG23	2.14	0.47
2:DK:88:TYR:CD2	2:DK:130:ILE:HD11	2.50	0.47
1:DL:97:VAL:HG11	2:DM:19:LEU:CD1	2.44	0.47
1:DN:122:PRO:HG2	1:DN:125:ALA:HB3	1.96	0.47
2:DO:63:SER:O	2:DO:67:ARG:HG3	2.14	0.47
2:DO:112:LEU:HD23	2:DO:160:LEU:HD13	1.95	0.47
5:DX:15:ARG:NH1	5:DX:17:ARG:HG2	2.30	0.47
9:EA:32:ASN:O	9:EA:36:GLN:HG3	2.15	0.47
9:EA:43:ALA:O	9:EA:47:MET:HG3	2.13	0.47
1:AA:122:PRO:HB2	1:AA:125:ALA:CB	2.45	0.47
2:AB:72:MET:HG3	10:AB:200:CYC:OC	2.14	0.47
3:AE:104:ILE:HD13	3:AE:156:ILE:HD11	1.95	0.47
1:AH:38:ARG:NH1	1:AH:145:ASP:OD2	2.35	0.47
1:AJ:60:GLN:HB3	1:CC:61:LYS:NZ	2.28	0.47
1:AJ:109:LEU:HA	1:AJ:112:VAL:HG11	1.96	0.47
2:AL:5:ILE:HG21	6:BD:169:ALA:HA	1.95	0.47
1:AN:88:TYR:CE2	1:AN:160:MET:HE1	2.49	0.47
1:AV:5:THR:O	1:AV:9:VAL:HG13	2.13	0.47
1:AV:109:LEU:HD13	1:AV:159:LYS:HG3	1.96	0.47
2:AW:8:VAL:HG22	2:AW:26:LYS:HD2	1.96	0.47
2:AY:131:GLN:O	2:AY:134:LYS:HB3	2.13	0.47
9:BH:193:VAL:HG23	9:BH:255:GLY:HA3	1.97	0.47

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:BL:53:LYS:HG3	3:BM:118:SER:O	2.13	0.47
2:BL:104:LEU:HD12	2:BL:156:ILE:HG12	1.96	0.47
2:BQ:141:VAL:HG12	2:BQ:145:ALA:HB3	1.96	0.47
2:BT:108:VAL:O	2:BT:112:LEU:HB2	2.15	0.47
1:BV:57:ARG:HA	1:BV:60:GLN:HE21	1.79	0.47
1:CG:95:GLY:HA3	1:CG:104:ILE:HD11	1.96	0.47
5:CJ:4:PHE:O	5:CJ:31:VAL:HG12	2.14	0.47
6:CM:318:GLU:O	6:CM:322:ARG:HG3	2.13	0.47
6:CM:749:ARG:HH12	10:DV:200:CYC:CBA	2.26	0.47
6:CM:751:LEU:HD22	6:CM:760:PHE:HE2	1.80	0.47
1:CR:18:LEU:HD22	2:CS:97:LEU:HD13	1.97	0.47
1:CR:157:ILE:O	1:CR:161:SER:HB2	2.15	0.47
1:CT:109:LEU:HD12	1:CT:159:LYS:HB3	1.97	0.47
2:CU:108:VAL:O	2:CU:112:LEU:HB2	2.14	0.47
10:CY:200:CYC:HHA	10:CY:200:CYC:HBD1	1.97	0.47
1:DA:75:GLU:N	1:DA:75:GLU:OE1	2.46	0.47
2:DB:81:CYS:HB3	10:DB:200:CYC:HBC1	1.64	0.47
2:DD:24:MET:O	2:DD:28:LYS:HG3	2.15	0.47
8:DG:237:ALA:HB3	1:DL:56:ASP:OD1	2.15	0.47
2:DO:2:GLN:OE1	2:DO:10:ASN:ND2	2.48	0.47
2:DO:76:ARG:HH22	5:DX:63:ASN:HA	1.79	0.47
1:DQ:47:ARG:O	1:DQ:50:ILE:HG22	2.15	0.47
1:DQ:116:TYR:CD2	1:DQ:123:ILE:HG13	2.49	0.47
2:DR:8:VAL:HG11	2:DR:27:LEU:CD1	2.44	0.47
1:DS:76:GLU:HG3	8:DY:202:ARG:HD2	1.97	0.47
9:EA:155:ARG:O	9:EA:159:VAL:HG13	2.14	0.47
1:AC:14:GLU:OE2	6:BD:263:LYS:HB2	2.14	0.47
2:AL:110:ASN:OD1	2:AL:111:GLY:N	2.47	0.47
2:AQ:96:MET:HE3	2:AQ:96:MET:HB3	1.81	0.47
1:AV:47:ARG:HB2	1:AV:89:LEU:HD23	1.96	0.47
6:BD:171:ASP:OD2	6:BD:173:ASN:HB2	2.15	0.47
6:BD:453:PRO:HD2	6:BD:607:ARG:CZ	2.45	0.47
9:BH:238:LEU:HD12	9:BH:242:ARG:HH21	1.80	0.47
1:BI:115:MET:HE3	1:BI:116:TYR:CE1	2.49	0.47
1:BI:130:VAL:HB	1:BI:157:ILE:HG13	1.96	0.47
2:BJ:88:TYR:CD2	2:BJ:130:ILE:HD11	2.50	0.47
1:BP:102:THR:O	1:BP:106:GLU:HG2	2.15	0.47
1:BV:116:TYR:OH	10:BV:200:CYC:HHB	2.14	0.47
6:CM:186:ILE:HD11	10:CM:902:CYC:NC	2.30	0.47
6:CM:331:LYS:HA	6:CM:335:GLU:OE1	2.15	0.47
6:CM:799:LYS:CE	6:CM:803:MET:HE3	2.45	0.47

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
9:CQ:37:LEU:HD12	9:CQ:40:ILE:HD11	1.97	0.47
1:CR:19:SER:O	1:CR:23:LEU:HD23	2.15	0.47
2:CU:71:ASN:HD22	2:CU:121:VAL:HA	1.79	0.47
1:CY:34:ALA:HA	2:CZ:28:LYS:NZ	2.30	0.47
1:CY:39:ILE:HA	1:CY:42:THR:CG2	2.44	0.47
1:CY:50:ILE:HD12	1:CY:136:VAL:HB	1.96	0.47
2:CZ:44:ILE:CG2	2:CZ:89:LEU:HD21	2.45	0.47
2:DB:123:ILE:H	2:DB:123:ILE:HD12	1.79	0.47
10:DB:200:CYC:HHA	10:DB:200:CYC:HBD1	1.96	0.47
9:DI:222:ALA:HB3	9:DI:299:PHE:O	2.14	0.47
9:DI:292:LEU:HG	9:DI:298:ILE:N	2.17	0.47
1:DL:93:THR:HA	1:DL:96:VAL:HG12	1.97	0.47
1:DL:121:THR:HG23	10:DL:200:CYC:C1C	2.44	0.47
1:DL:132:GLU:O	1:DL:136:VAL:HG23	2.15	0.47
1:DU:94:TYR:CG	2:DV:9:ILE:HG23	2.49	0.47
2:DV:132:ALA:HA	2:DV:135:GLU:OE1	2.15	0.47
5:DX:41:GLN:O	5:DX:45:LYS:HG3	2.14	0.47
8:DZ:210:LEU:HD11	8:DZ:216:MET:HB2	1.96	0.47
1:AA:50:ILE:HD12	1:AA:136:VAL:CG1	2.45	0.47
1:AA:126:VAL:HG22	10:AA:200:CYC:HMC1	1.97	0.47
2:AB:73:TYR:OH	2:AD:13:ASP:OD1	2.22	0.47
2:AB:133:ILE:O	2:AB:137:THR:HG22	2.15	0.47
10:AB:200:CYC:HHA	10:AB:200:CYC:HBD1	1.95	0.47
1:AC:22:GLU:HA	1:AC:25:ARG:HG2	1.96	0.47
3:AE:102:GLU:O	3:AE:106:THR:HG23	2.15	0.47
2:AF:5:ILE:HG13	2:AF:27:LEU:CD1	2.44	0.47
2:AF:39:ARG:HE	1:CC:142:SER:HA	1.78	0.47
2:AF:44:ILE:CD1	2:AF:149:MET:HE3	2.45	0.47
2:AF:112:LEU:HD22	10:AF:200:CYC:HMB1	1.96	0.47
2:AI:71:ASN:HD22	2:AI:122:PRO:CD	2.28	0.47
2:AI:83:ARG:HD2	6:BD:369:SER:HB3	1.97	0.47
4:AK:4:ALA:O	4:AK:7:THR:HG22	2.13	0.47
1:AP:68:PRO:HG2	1:BR:66:VAL:CG1	2.44	0.47
1:AP:126:VAL:HG12	1:AP:160:MET:HE1	1.96	0.47
1:AP:144:ASP:N	1:AP:144:ASP:OD1	2.48	0.47
1:AR:27:LYS:O	1:AR:31:THR:HG23	2.15	0.47
1:AR:85:MET:HE3	1:AR:133:MET:HE2	1.97	0.47
2:AS:147:LYS:HE3	7:BF:93:PRO:HD3	1.97	0.47
2:AU:24:MET:CE	1:AZ:41:GLU:HB2	2.44	0.47
1:AV:37:LEU:HD21	2:AW:24:MET:CE	2.45	0.47
1:AV:65:ILE:HG22	1:AV:72:ALA:HB3	1.96	0.47

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:AX:82:LEU:HB2	8:BG:242:ILE:HD12	1.95	0.47
1:AX:102:THR:O	1:AX:106:GLU:HG2	2.15	0.47
2:AY:75:THR:HG23	1:AZ:110:VAL:O	2.15	0.47
1:AZ:5:THR:O	1:AZ:9:VAL:HG23	2.15	0.47
1:AZ:64:ASP:HA	1:AZ:67:SER:HB2	1.95	0.47
1:AZ:77:MET:SD	10:AZ:200:CYC:HAD1	2.55	0.47
6:BD:72:ALA:HB1	6:BD:81:PRO:HB2	1.95	0.47
6:BD:399:ARG:HD2	6:BD:404:GLU:OE2	2.14	0.47
6:BD:612:ARG:NH2	6:BD:619:GLU:OE2	2.30	0.47
9:BH:97:THR:HG22	9:BH:101:TRP:CZ2	2.50	0.47
9:BH:214:ILE:HG21	9:BH:238:LEU:HD21	1.96	0.47
9:BH:280:GLY:O	9:BH:284:MET:HG2	2.15	0.47
1:BI:116:TYR:HE2	1:BI:126:VAL:HG11	1.80	0.47
1:BI:126:VAL:HG12	1:BI:160:MET:SD	2.55	0.47
2:BJ:37:ARG:HD3	2:BJ:96:MET:O	2.14	0.47
2:BL:87:TYR:HE2	5:CJ:22:LEU:HB2	1.80	0.47
2:BL:133:ILE:O	2:BL:137:THR:HG22	2.14	0.47
3:BM:115:MET:HE2	10:BM:200:CYC:NB	2.25	0.47
2:BQ:44:ILE:HD13	2:BQ:149:MET:HE2	1.96	0.47
2:BQ:88:TYR:OH	2:BQ:116:TYR:OH	2.23	0.47
1:BR:58:LEU:HD22	1:BR:129:SER:CB	2.44	0.47
4:BS:44:ILE:HG22	4:BS:138:ILE:CD1	2.43	0.47
4:BS:56:ALA:HB1	4:BS:60:PHE:CE2	2.49	0.47
2:BT:90:ARG:HG2	2:BT:94:TYR:CZ	2.50	0.47
2:BW:96:MET:HA	2:BW:152:TYR:CE2	2.50	0.47
1:BZ:64:ASP:HA	1:BZ:67:SER:HB2	1.97	0.47
2:CA:63:SER:H	2:CA:66:THR:HG22	1.80	0.47
2:CH:74:THR:HB	2:CH:77:ARG:HG3	1.96	0.47
2:CH:88:TYR:CE2	2:CH:130:ILE:HD11	2.49	0.47
5:CK:6:ILE:HD13	5:CK:36:TRP:CZ3	2.49	0.47
6:CM:80:SER:HB3	6:CM:83:SER:OG	2.15	0.47
6:CM:152:SER:O	6:CM:156:MET:HG3	2.14	0.47
6:CM:513:PHE:HB3	6:CM:542:LEU:CD1	2.45	0.47
6:CM:625:ASP:CG	6:CM:629:LYS:HE2	2.40	0.47
6:CM:666:GLN:O	6:CM:670:ALA:HB3	2.15	0.47
6:CM:847:THR:HG21	6:CM:857:ARG:NH2	2.30	0.47
6:CM:867:PRO:HG3	2:DT:115:THR:OG1	2.15	0.47
1:CR:91:LEU:CD1	1:CR:104:ILE:HA	2.37	0.47
1:CR:101:VAL:HG13	1:CY:20:PRO:HG2	1.96	0.47
2:CS:91:TYR:CE1	2:CS:107:ARG:HD3	2.47	0.47
2:CS:109:LEU:HD13	2:CS:159:GLY:HA3	1.96	0.47

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:CT:25:ARG:HH11	1:DC:25:ARG:HG2	1.79	0.47
2:DB:73:TYR:O	2:DB:74:THR:OG1	2.32	0.47
1:DC:5:THR:O	1:DC:9:VAL:HG23	2.14	0.47
1:DC:66:VAL:HA	1:DC:72:ALA:O	2.14	0.47
2:DD:108:VAL:HA	10:DD:200:CYC:CBB	2.45	0.47
9:DI:19:ASP:O	9:DI:22:PRO:HD2	2.14	0.47
1:DJ:16:ARG:C	2:DK:90:ARG:HD2	2.39	0.47
2:DK:2:GLN:H	2:DK:102:SER:CB	2.27	0.47
2:DK:112:LEU:CD1	10:DK:200:CYC:HMB3	2.45	0.47
1:DL:8:ILE:CD1	1:DL:18:LEU:HD11	2.45	0.47
2:DM:60:LEU:HD11	2:DM:129:ALA:HB2	1.96	0.47
2:DM:73:TYR:O	2:DM:74:THR:OG1	2.25	0.47
1:DN:97:VAL:HG21	2:DO:19:LEU:HD11	1.96	0.47
2:DR:105:ASP:OD1	2:DR:109:LEU:HD12	2.15	0.47
1:DS:123:ILE:HG12	1:DS:160:MET:HG3	1.96	0.47
1:AC:157:ILE:O	1:AC:161:SER:OG	2.22	0.47
2:AD:54:GLU:O	2:AD:58:LYS:HG2	2.15	0.47
1:AH:55:GLY:N	1:AH:85:MET:HE1	2.30	0.47
2:AI:101:ALA:HB2	2:AI:152:TYR:CD1	2.50	0.47
1:AN:126:VAL:CG2	10:AN:200:CYC:HMC2	2.44	0.47
2:AS:90:ARG:HG2	2:AS:94:TYR:CZ	2.49	0.47
1:AV:47:ARG:O	1:AV:51:VAL:HG22	2.14	0.47
2:AW:57:ALA:O	8:BG:249:ARG:NH2	2.47	0.47
1:AX:121:THR:HG23	10:AX:200:CYC:C1C	2.44	0.47
2:AY:116:TYR:HB3	2:AY:123:ILE:HD11	1.96	0.47
9:BH:40:ILE:HG13	11:BH:400:45D:H371	1.96	0.47
1:BI:48:GLU:O	1:BI:51:VAL:HG12	2.15	0.47
1:BK:17:TYR:CE1	2:BL:90:ARG:HG3	2.50	0.47
2:BL:57:ALA:HA	2:BL:61:LEU:HD12	1.97	0.47
3:BM:46:ASN:ND2	3:BM:140:LEU:HD21	2.30	0.47
2:BN:4:ALA:HB2	2:BN:99:GLY:C	2.40	0.47
2:BQ:119:LEU:HD23	6:CM:435:ALA:HB2	1.96	0.47
4:BS:113:LEU:HD21	10:BS:200:CYC:HMB1	1.96	0.47
2:BT:18:TYR:CE2	6:CM:161:ARG:HA	2.50	0.47
2:BY:78:TYR:CD1	1:BZ:115:MET:HG3	2.49	0.47
5:CJ:10:VAL:O	5:CJ:10:VAL:HG13	2.14	0.47
6:CM:682:ARG:NH2	6:CM:684:ASP:OD2	2.47	0.47
6:CM:842:MET:O	6:CM:846:GLN:HG3	2.15	0.47
9:CQ:115:GLU:HG2	9:CQ:119:GLN:HE22	1.79	0.47
9:CQ:204:ASN:CB	9:CQ:213:LEU:HB2	2.41	0.47
10:CR:200:CYC:CAA	2:CW:61:LEU:HD22	2.44	0.47

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:CS:85:LEU:HD23	2:CS:88:TYR:CD2	2.50	0.47
2:CS:92:ALA:CB	2:CS:149:MET:HE3	2.44	0.47
1:CV:17:TYR:OH	2:CW:89:LEU:HG	2.15	0.47
1:CV:27:LYS:O	1:CV:31:THR:HG23	2.15	0.47
2:DB:71:ASN:ND2	2:DB:121:VAL:HA	2.30	0.47
2:DD:84:ASP:OD2	10:DD:200:CYC:ND	2.47	0.47
9:DI:21:VAL:HG13	9:DI:22:PRO:HD3	1.95	0.47
9:DI:117:MET:HE1	11:DI:400:45D:H311	1.97	0.47
1:DJ:16:ARG:HA	2:DK:90:ARG:HD2	1.97	0.47
1:DJ:35:ALA:O	1:DJ:39:ILE:HG13	2.14	0.47
1:DJ:102:THR:O	1:DJ:106:GLU:HG3	2.14	0.47
1:DJ:113:ARG:NH1	1:DS:117:ARG:HD2	2.29	0.47
2:DK:106:GLU:HG3	5:DX:57:THR:CG2	2.41	0.47
1:DL:39:ILE:HD11	1:DL:145:ASP:HB3	1.97	0.47
1:DL:53:GLN:O	1:DL:57:ARG:HD3	2.15	0.47
1:DL:61:LYS:O	1:DL:62:ARG:HG2	2.15	0.47
1:DQ:41:GLU:HG2	2:DR:24:MET:HE2	1.97	0.47
8:DY:202:ARG:NH2	8:DY:204:ILE:HB	2.30	0.47
9:EA:17:ALA:O	9:EA:20:VAL:HG22	2.14	0.47
9:EA:21:VAL:CG1	9:EA:149:GLN:HG3	2.45	0.47
1:AC:102:THR:HB	1:AC:103:PRO:HD3	1.96	0.47
2:AD:96:MET:HE2	2:AD:148:GLU:OE1	2.15	0.47
2:AD:106:GLU:HG3	2:AD:107:ARG:HD3	1.97	0.47
1:AH:71:ASN:HB3	10:AH:200:CYC:OC	2.14	0.47
1:AN:102:THR:HB	1:AN:103:PRO:HD3	1.96	0.47
2:AQ:88:TYR:OH	2:AQ:112:LEU:HD21	2.15	0.47
2:AW:41:ALA:N	2:AW:96:MET:HE1	2.30	0.47
2:AW:78:TYR:CD1	1:AX:115:MET:HG3	2.50	0.47
2:AY:108:VAL:HG12	2:AY:109:LEU:CD2	2.43	0.47
1:AZ:20:PRO:HA	1:AZ:23:LEU:HD12	1.97	0.47
6:BD:477:ASN:O	6:BD:617:ARG:HD3	2.15	0.47
6:BD:543:ILE:HG12	6:BD:568:LEU:HD23	1.97	0.47
2:BN:132:ALA:O	2:BN:136:VAL:HG23	2.15	0.47
1:BP:40:ALA:HB2	1:BP:97:VAL:HG22	1.96	0.47
1:BP:76:GLU:CG	1:BP:77:MET:HE2	2.43	0.47
2:BQ:2:GLN:NE2	2:BQ:10:ASN:OD1	2.47	0.47
2:BQ:101:ALA:HB2	2:BQ:152:TYR:CD1	2.49	0.47
2:BW:41:ALA:N	2:BW:96:MET:HE1	2.30	0.47
1:BX:68:PRO:HA	1:BX:73:TYR:CG	2.50	0.47
2:CA:148:GLU:HG3	2:CA:152:TYR:CE2	2.50	0.47
1:CG:63:PRO:HB2	8:CP:232:ILE:HD13	1.97	0.47

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
6:CM:788:MET:HE3	6:CM:788:MET:HB3	1.79	0.47
9:CQ:183:ALA:N	9:CQ:186:THR:HG1	2.12	0.47
2:CS:119:LEU:HD13	10:CS:200:CYC:CBD	2.41	0.47
1:CT:161:SER:OG	2:DD:49:ALA:HB3	2.15	0.47
2:DK:2:GLN:OE1	5:DX:65:GLY:HA3	2.14	0.47
1:DU:37:LEU:HD13	2:DV:28:LYS:HG3	1.97	0.47
2:DV:126:THR:HG22	10:DV:200:CYC:HMC1	1.96	0.47
2:AB:73:TYR:HB3	2:AB:77:ARG:NH2	2.30	0.47
1:AC:25:ARG:HG3	1:AC:26:ILE:N	2.29	0.47
2:AD:95:ALA:HB2	2:AD:104:LEU:CD2	2.44	0.47
3:AE:36:ARG:HD2	3:AE:96:VAL:O	2.15	0.47
3:AE:68:PRO:HA	3:AE:73:TYR:OH	2.15	0.47
4:AK:113:LEU:CD1	4:AK:164:LEU:HD22	2.36	0.47
2:AQ:58:LYS:HE2	2:AQ:135:GLU:OE1	2.14	0.47
1:AX:89:LEU:O	1:AX:93:THR:HG23	2.15	0.47
6:BD:31:GLN:HB3	6:BD:33:ARG:HE	1.79	0.47
6:BD:626:ILE:HG21	6:BD:635:VAL:HA	1.96	0.47
6:BD:891:VAL:HG11	2:DB:105:ASP:OD1	2.15	0.47
1:BI:40:ALA:O	1:BI:44:THR:HG22	2.14	0.47
3:BM:112:VAL:HA	3:BM:115:MET:CB	2.45	0.47
4:BS:167:LEU:HD12	2:BW:10:ASN:ND2	2.30	0.47
2:BT:34:GLY:O	2:BT:38:VAL:HG23	2.14	0.47
1:BV:11:ALA:HB2	1:BV:18:LEU:HD23	1.97	0.47
1:BV:101:VAL:HG11	1:CC:20:PRO:HG2	1.96	0.47
1:BX:72:ALA:HA	1:BX:77:MET:HB3	1.97	0.47
1:CG:66:VAL:HG21	8:CP:235:VAL:HG21	1.96	0.47
2:CH:24:MET:HE3	2:CH:28:LYS:CD	2.45	0.47
2:CH:77:ARG:HB3	10:CH:200:CYC:HMD1	1.97	0.47
6:CM:273:LEU:HB3	6:CM:277:GLU:CG	2.44	0.47
6:CM:444:GLN:HE21	6:CM:448:GLY:HA2	1.80	0.47
2:CS:36:LEU:HD11	2:CS:144:ASP:CG	2.40	0.47
2:CW:56:VAL:HG12	2:CW:85:LEU:HD12	1.97	0.47
2:CW:60:LEU:HD11	2:CW:129:ALA:HB2	1.97	0.47
2:CW:90:ARG:HH11	2:CW:94:TYR:HH	1.58	0.47
2:CZ:41:ALA:CA	2:CZ:96:MET:HE1	2.45	0.47
2:CZ:55:ALA:HA	2:CZ:132:ALA:HB1	1.97	0.47
2:CZ:123:ILE:H	2:CZ:123:ILE:HD12	1.80	0.47
2:DB:106:GLU:OE1	2:DB:107:ARG:HG3	2.14	0.47
1:DC:97:VAL:HG21	2:DD:19:LEU:CD1	2.45	0.47
8:DH:222:ALA:HB1	1:DS:78:THR:OG1	2.15	0.47
9:DI:21:VAL:HG12	9:DI:22:PRO:HD3	1.97	0.47

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
9:DI:191:GLU:HB2	9:DI:254:ARG:HA	1.97	0.47
1:DL:141:MET:SD	1:DL:146:ALA:HA	2.55	0.47
1:DU:81:CYS:O	1:DU:85:MET:HG3	2.15	0.47
2:DV:52:VAL:HG21	2:DV:86:ASP:OD1	2.15	0.47
9:EA:263:GLY:C	9:EA:294:PRO:HB3	2.40	0.47
1:AA:88:TYR:O	1:AA:92:VAL:HG23	2.15	0.46
1:AC:92:VAL:HG11	1:AC:153:PHE:CZ	2.50	0.46
2:AD:11:SER:O	2:AD:14:VAL:HG22	2.16	0.46
3:AE:50:ILE:HD11	3:AE:140:LEU:HD13	1.97	0.46
1:AH:91:LEU:HB3	1:AH:104:ILE:HG23	1.97	0.46
1:AJ:19:SER:HB2	1:AJ:20:PRO:HD2	1.96	0.46
4:AK:78:ARG:NE	10:AK:200:CYC:O1D	2.48	0.46
2:AL:74:THR:OG1	2:AL:77:ARG:HG3	2.15	0.46
1:AN:8:ILE:HG23	2:AO:94:TYR:CD1	2.49	0.46
1:AP:10:ASN:OD1	6:BD:511:LYS:HE3	2.15	0.46
1:AR:39:ILE:O	1:AR:43:LEU:HG	2.16	0.46
10:AU:200:CYC:HAA1	5:BB:36:TRP:CD1	2.45	0.46
1:AV:57:ARG:O	1:AV:60:GLN:NE2	2.48	0.46
1:AV:123:ILE:HG12	1:AV:160:MET:O	2.15	0.46
2:AY:96:MET:HA	2:AY:152:TYR:CE2	2.50	0.46
6:BD:19:THR:OG1	6:BD:22:VAL:HG23	2.14	0.46
6:BD:543:ILE:HG23	6:BD:578:PHE:CZ	2.50	0.46
9:BH:126:PRO:HD3	11:BH:400:45D:H211	1.96	0.46
2:BL:152:TYR:O	2:BL:156:ILE:HG13	2.15	0.46
3:BM:116:TYR:CE2	3:BM:126:MET:HE1	2.50	0.46
2:BT:31:PHE:CE1	6:CM:50:GLY:HA3	2.50	0.46
2:BY:112:LEU:HG	2:BY:160:LEU:CD1	2.45	0.46
10:BY:200:CYC:HB	10:BY:200:CYC:CMA	2.28	0.46
1:CE:37:LEU:HD21	2:CF:24:MET:CE	2.45	0.46
1:CE:141:MET:HE1	1:CE:145:ASP:HB2	1.96	0.46
1:CG:96:VAL:HG12	1:CG:152:TYR:CD2	2.50	0.46
6:CM:10:SER:HB2	6:CM:415:GLN:NE2	2.29	0.46
6:CM:187:GLU:OE1	6:CM:192:ILE:HG12	2.15	0.46
6:CM:515:LEU:HG	6:CM:542:LEU:CD2	2.45	0.46
1:CR:26:ILE:O	1:CR:30:VAL:HG13	2.14	0.46
1:CT:49:THR:HG22	1:CT:53:GLN:OE1	2.15	0.46
1:CT:62:ARG:NH1	1:CT:124:GLU:OE2	2.44	0.46
1:CV:123:ILE:HG12	1:CV:160:MET:O	2.15	0.46
1:DC:109:LEU:HD13	1:DC:159:LYS:CG	2.42	0.46
8:DG:248:ARG:NH1	2:DK:67:ARG:HD3	2.30	0.46
1:DJ:37:LEU:HD11	2:DK:24:MET:HE1	1.97	0.46

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:DL:38:ARG:HA	1:DL:41:GLU:OE1	2.15	0.46
2:DM:93:THR:HA	2:DM:149:MET:SD	2.55	0.46
1:DN:72:ALA:HA	1:DN:77:MET:HB3	1.97	0.46
1:DN:107:ILE:CG2	10:DN:200:CYC:HBB1	2.45	0.46
2:DR:56:VAL:O	2:DR:61:LEU:HB2	2.14	0.46
2:DT:73:TYR:O	2:DT:74:THR:OG1	2.33	0.46
2:DT:122:PRO:HD2	10:DT:200:CYC:HMC3	1.96	0.46
1:AC:130:VAL:HA	1:AC:133:MET:CE	2.37	0.46
2:AF:4:ALA:HB2	2:AF:99:GLY:O	2.15	0.46
2:AF:123:ILE:HD12	2:AF:160:LEU:CD2	2.45	0.46
1:AH:91:LEU:O	1:AH:104:ILE:HG12	2.16	0.46
4:AK:41:ALA:CB	4:AK:98:ILE:HD11	2.45	0.46
1:AN:91:LEU:O	1:AN:104:ILE:HG12	2.16	0.46
1:AV:93:THR:HA	1:AV:96:VAL:HG22	1.97	0.46
2:AY:68:PRO:HG3	2:AY:73:TYR:CE1	2.49	0.46
1:AZ:71:ASN:OD1	10:AZ:200:CYC:HBD2	2.15	0.46
6:BD:244:PRO:CG	6:BD:248:ILE:HD13	2.34	0.46
6:BD:365:PHE:HA	6:BD:368:VAL:HG12	1.97	0.46
6:BD:788:MET:HE1	1:CY:9:VAL:HG12	1.97	0.46
2:BL:3:ASP:HA	2:BL:100:ASP:HB3	1.96	0.46
3:BM:131:THR:HG22	3:BM:157:ILE:CD1	2.45	0.46
2:BQ:90:ARG:O	2:BQ:93:THR:OG1	2.27	0.46
1:BR:134:LYS:HB2	1:BR:153:PHE:HB3	1.98	0.46
4:BS:72:ASN:OD1	10:BS:200:CYC:HMD2	2.15	0.46
1:BV:65:ILE:HG23	10:BV:200:CYC:OC	2.15	0.46
2:BY:85:LEU:CG	10:BY:200:CYC:HBC1	2.44	0.46
1:CC:17:TYR:CE2	2:CD:90:ARG:HG3	2.50	0.46
6:CM:175:ILE:HD12	6:CM:227:LEU:HD12	1.96	0.46
6:CM:283:LYS:O	6:CM:287:ARG:HG3	2.14	0.46
7:CO:111:LYS:O	7:CO:115:ARG:HG3	2.15	0.46
9:CQ:117:MET:HE3	9:CQ:124:PRO:HA	1.95	0.46
9:CQ:140:ALA:HA	9:CQ:143:GLN:OE1	2.16	0.46
9:CQ:248:LEU:CA	9:CQ:275:THR:HG21	2.40	0.46
1:CR:36:ARG:NH1	1:CR:99:GLY:HA2	2.30	0.46
1:CR:108:GLY:HA2	2:CW:75:THR:HG23	1.97	0.46
1:CR:140:LEU:HD23	1:CR:140:LEU:O	2.15	0.46
1:CT:111:GLY:HA2	1:CT:114:GLU:OE1	2.15	0.46
1:CY:115:MET:HE1	2:DD:79:ALA:CA	2.45	0.46
2:CZ:109:LEU:CD2	2:CZ:159:GLY:HA3	2.44	0.46
1:DJ:20:PRO:O	1:DJ:23:LEU:HG	2.15	0.46
10:DM:200:CYC:HBB2	5:DX:20:ARG:HG3	1.96	0.46

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:DN:132:GLU:HA	1:DN:135:GLU:OE1	2.16	0.46
2:DO:68:PRO:HG3	2:DO:73:TYR:CE1	2.51	0.46
1:DS:63:PRO:HD2	9:EA:55:ALA:CB	2.45	0.46
1:DS:113:ARG:HG2	1:DS:123:ILE:CD1	2.45	0.46
1:DS:122:PRO:HG2	1:DS:125:ALA:HB3	1.98	0.46
1:DU:41:GLU:CD	2:DV:24:MET:HE2	2.40	0.46
1:DU:116:TYR:HD2	1:DU:123:ILE:HD12	1.79	0.46
1:DU:142:SER:O	1:DU:146:ALA:N	2.46	0.46
2:DV:15:GLN:HB3	2:DV:17:LYS:HG3	1.96	0.46
9:EA:260:ALA:HB3	9:EA:264:PHE:CB	2.46	0.46
9:EA:265:THR:HG22	9:EA:267:ILE:HG13	1.98	0.46
2:AD:68:PRO:HA	2:AD:73:TYR:CG	2.50	0.46
2:AF:50:THR:O	2:AF:54:GLU:HG2	2.15	0.46
1:AH:37:LEU:O	1:AH:40:ALA:HB3	2.15	0.46
4:AK:113:LEU:HD21	10:AK:200:CYC:HMB1	1.95	0.46
1:AN:126:VAL:HG22	10:AN:200:CYC:HMC2	1.97	0.46
2:AQ:51:ILE:HD11	2:AQ:140:LEU:HD23	1.96	0.46
2:AQ:52:VAL:O	2:AQ:56:VAL:HG13	2.15	0.46
1:AR:126:VAL:O	1:AR:129:SER:OG	2.28	0.46
1:AV:4:VAL:HG12	1:AV:26:ILE:HG23	1.96	0.46
1:AV:126:VAL:O	1:AV:129:SER:OG	2.27	0.46
10:AV:200:CYC:HHA	10:AV:200:CYC:HBA2	1.96	0.46
2:AY:68:PRO:HA	2:AY:73:TYR:CG	2.50	0.46
1:AZ:89:LEU:HB2	1:AZ:133:MET:HE1	1.97	0.46
5:BB:38:ARG:HE	6:BD:641:ASP:CB	2.24	0.46
6:BD:177:VAL:HG22	6:BD:259:ASN:HB2	1.96	0.46
6:BD:186:ILE:O	6:BD:190:CYS:HB3	2.15	0.46
6:BD:199:ILE:CG2	6:BD:227:LEU:HD22	2.45	0.46
6:BD:513:PHE:HB3	6:BD:542:LEU:CD1	2.46	0.46
6:BD:606:HIS:HB2	6:BD:639:MET:CE	2.45	0.46
6:BD:811:ARG:NE	6:BD:843:GLU:OE2	2.42	0.46
9:BH:131:LEU:HD23	9:BH:136:ASN:HA	1.96	0.46
1:BI:125:ALA:CB	10:BI:200:CYC:HMC2	2.45	0.46
2:BJ:51:ILE:HD11	2:BJ:140:LEU:HD23	1.97	0.46
1:BK:113:ARG:NH2	1:BP:117:ARG:HD3	2.21	0.46
2:BL:112:LEU:HA	2:BL:115:THR:CG2	2.45	0.46
2:BL:112:LEU:HD11	10:BL:200:CYC:C1B	2.46	0.46
1:BP:71:ASN:HB3	10:BP:200:CYC:OC	2.15	0.46
1:BP:84:ASP:HB2	10:BP:200:CYC:HBC3	1.97	0.46
2:BQ:71:ASN:OD1	2:BQ:77:ARG:HD3	2.15	0.46
2:BY:96:MET:HE3	2:BY:96:MET:HB3	1.76	0.46

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:BY:141:VAL:HG12	2:BY:145:ALA:HB3	1.97	0.46
1:BZ:27:LYS:O	1:BZ:31:THR:HG23	2.15	0.46
2:CD:112:LEU:HD23	2:CD:112:LEU:HA	1.66	0.46
1:CE:5:THR:O	1:CE:9:VAL:HG13	2.15	0.46
2:CF:111:GLY:O	2:CF:115:THR:OG1	2.27	0.46
1:CG:50:ILE:HD11	1:CG:137:ALA:HB2	1.98	0.46
1:CG:107:ILE:HD11	6:CM:532:VAL:HG22	1.96	0.46
1:CG:116:TYR:CE2	1:CG:126:VAL:HG21	2.50	0.46
2:CH:123:ILE:HA	2:CH:126:THR:HG22	1.97	0.46
6:CM:348:ARG:HD2	6:CM:417:LEU:HG	1.98	0.46
9:CQ:28:PHE:CD2	9:CQ:39:LEU:HD23	2.50	0.46
9:CQ:147:SER:HA	9:CQ:150:GLN:OE1	2.15	0.46
2:CU:1:MET:SD	2:CU:103:ILE:HD13	2.56	0.46
2:CW:126:THR:HG21	2:CW:160:LEU:HD21	1.96	0.46
1:CY:137:ALA:HB1	1:CY:141:MET:HE3	1.97	0.46
2:CZ:24:MET:O	2:CZ:28:LYS:HG2	2.14	0.46
9:DI:286:ILE:HD11	9:DI:303:ILE:CG2	2.45	0.46
1:DJ:5:THR:HG23	2:DK:1:MET:CE	2.37	0.46
1:DL:117:ARG:HD3	1:DQ:113:ARG:NH1	2.30	0.46
2:DM:15:GLN:HG3	2:DM:17:LYS:HG2	1.98	0.46
2:DO:10:ASN:O	2:DO:14:VAL:HG13	2.15	0.46
1:DQ:75:GLU:O	1:DQ:78:THR:OG1	2.27	0.46
1:DS:85:MET:HE2	1:DS:133:MET:HG3	1.97	0.46
2:DT:60:LEU:O	2:DT:63:SER:OG	2.29	0.46
9:EA:177:VAL:HG12	9:EA:179:PRO:HD2	1.97	0.46
9:EA:248:LEU:HD11	9:EA:277:TRP:CE3	2.49	0.46
1:AA:40:ALA:O	1:AA:44:THR:HG22	2.15	0.46
3:AE:30:LEU:HD13	2:AF:31:PHE:CD1	2.43	0.46
2:AF:39:ARG:O	2:AF:43:VAL:HG23	2.15	0.46
2:AL:56:VAL:HG12	2:AL:61:LEU:CG	2.38	0.46
1:AR:97:VAL:HG21	2:AS:19:LEU:HD12	1.97	0.46
2:AU:88:TYR:CE1	10:AU:200:CYC:HMB1	2.51	0.46
6:BD:450:GLY:O	6:BD:600:LYS:HE3	2.16	0.46
6:BD:736:LYS:O	6:BD:740:ARG:HG3	2.15	0.46
6:BD:754:TYR:HH	10:DD:200:CYC:CGA	2.26	0.46
1:BP:91:LEU:O	1:BP:104:ILE:HG12	2.15	0.46
1:BR:19:SER:OG	1:BR:22:GLU:HG3	2.15	0.46
2:CA:20:ASP:O	2:CA:24:MET:HG2	2.16	0.46
1:CC:91:LEU:O	1:CC:104:ILE:HG12	2.15	0.46
1:CE:83:ARG:NH2	1:CE:84:ASP:OD1	2.32	0.46
2:CH:68:PRO:HA	2:CH:73:TYR:CG	2.51	0.46

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:CH:96:MET:HG3	2:CH:148:GLU:OE2	2.16	0.46
5:CK:6:ILE:O	5:CK:28:THR:HA	2.14	0.46
9:CQ:250:LEU:HD22	9:CQ:273:VAL:HG23	1.96	0.46
2:CU:112:LEU:HD22	10:CU:200:CYC:HMB2	1.96	0.46
2:CW:48:ALA:O	2:CW:52:VAL:HG23	2.16	0.46
1:DC:116:TYR:HE2	1:DC:126:VAL:HG11	1.79	0.46
9:DI:81:GLN:HA	9:DI:176:VAL:CG2	2.45	0.46
9:DI:146:GLU:HG2	8:DZ:234:LYS:NZ	2.31	0.46
9:DI:146:GLU:HA	8:DZ:234:LYS:HZ2	1.81	0.46
9:DI:238:LEU:HD21	9:DI:242:ARG:NH2	2.31	0.46
10:DJ:200:CYC:O2D	2:DO:62:TYR:OH	2.23	0.46
10:DJ:200:CYC:HAA1	2:DO:61:LEU:HD22	1.98	0.46
2:DK:78:TYR:HE1	1:DL:115:MET:HA	1.80	0.46
2:DK:104:LEU:HD23	2:DK:155:TYR:CD2	2.38	0.46
1:DN:17:TYR:CZ	2:DO:90:ARG:HG3	2.51	0.46
10:DO:200:CYC:HAD2	5:DX:33:TYR:OH	2.16	0.46
2:DR:40:ALA:HB2	2:DR:145:ALA:CB	2.42	0.46
2:DT:68:PRO:HA	2:DT:73:TYR:CD1	2.50	0.46
1:DU:44:THR:HG23	2:DV:18:TYR:CD2	2.50	0.46
1:DU:71:ASN:HD22	1:DU:121:THR:HA	1.81	0.46
2:DV:73:TYR:O	2:DV:74:THR:OG1	2.23	0.46
9:EA:187:LYS:HE2	9:EA:187:LYS:HA	1.98	0.46
9:EA:231:ILE:CG1	9:EA:236:ASN:HB3	2.46	0.46
9:EA:248:LEU:HD21	9:EA:277:TRP:CE3	2.50	0.46
2:AF:39:ARG:CZ	1:CC:142:SER:HA	2.46	0.46
1:AH:134:LYS:HB2	1:AH:153:PHE:HB3	1.96	0.46
1:AJ:109:LEU:HD13	1:AJ:159:LYS:CB	2.46	0.46
4:AK:44:ILE:CG2	4:AK:138:ILE:HD12	2.46	0.46
1:AN:46:SER:HB3	1:AN:50:ILE:HG13	1.98	0.46
1:AP:110:VAL:HG11	6:BD:489:PRO:HB3	1.98	0.46
2:AS:2:GLN:HG3	2:AS:100:ASP:OD2	2.16	0.46
6:BD:666:GLN:O	6:BD:670:ALA:HB3	2.15	0.46
6:BD:866:PHE:HB3	6:BD:867:PRO:HD3	1.98	0.46
9:BH:20:VAL:CG2	9:BH:145:LEU:HD21	2.45	0.46
9:BH:86:LEU:HD23	9:BH:95:CYS:SG	2.56	0.46
1:BK:26:ILE:O	1:BK:30:VAL:HG23	2.14	0.46
2:BL:122:PRO:HG2	2:BL:125:SER:OG	2.16	0.46
3:BM:144:GLU:OE1	3:BM:144:GLU:N	2.40	0.46
2:BN:25:ASP:OD1	2:BN:26:LYS:N	2.48	0.46
2:BN:37:ARG:NH1	2:BN:99:GLY:HA2	2.30	0.46
2:BQ:78:TYR:CD1	1:BR:115:MET:HG3	2.51	0.46

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:BR:18:LEU:HD11	4:BS:95:TYR:HD1	1.79	0.46
1:BV:113:ARG:CG	1:BV:123:ILE:HD13	2.38	0.46
2:CD:25:ASP:OD1	2:CD:26:LYS:N	2.48	0.46
1:CG:67:SER:HB3	1:CG:68:PRO:HD2	1.97	0.46
2:CH:145:ALA:O	2:CH:148:GLU:HG3	2.15	0.46
5:CJ:12:SER:HB3	5:CJ:15:ARG:HG3	1.96	0.46
6:CM:559:GLN:HB3	6:CM:587:LEU:HD22	1.96	0.46
6:CM:727:VAL:HG13	5:DX:20:ARG:HD3	1.96	0.46
9:CQ:273:VAL:HG22	9:CQ:275:THR:H	1.80	0.46
2:CS:71:ASN:ND2	2:CS:122:PRO:HD3	2.30	0.46
2:CS:71:ASN:CB	2:CS:122:PRO:HD3	2.46	0.46
1:CT:27:LYS:HZ2	2:CU:35:GLU:CD	2.24	0.46
1:CY:76:GLU:HG3	8:DH:203:PHE:CE2	2.50	0.46
2:CZ:132:ALA:O	2:CZ:136:VAL:HG23	2.16	0.46
1:DA:18:LEU:HD22	2:DB:97:LEU:CD2	2.46	0.46
1:DA:27:LYS:O	1:DA:31:THR:HG23	2.15	0.46
5:DF:9:CYS:HA	5:DF:25:THR:O	2.15	0.46
5:DF:29:LYS:NZ	5:DF:39:GLU:OE2	2.48	0.46
9:DI:177:VAL:HG12	9:DI:179:PRO:HD2	1.97	0.46
1:DJ:110:VAL:CG1	2:DO:76:ARG:HB2	2.45	0.46
1:DL:80:THR:HG21	10:DL:200:CYC:HAD2	1.98	0.46
2:DM:6:THR:HG22	2:DM:10:ASN:ND2	2.29	0.46
1:DN:109:LEU:HD22	1:DN:159:LYS:CG	2.45	0.46
1:DQ:71:ASN:HD21	1:DQ:122:PRO:HD2	1.79	0.46
2:DR:51:ILE:HG21	2:DR:137:THR:OG1	2.16	0.46
2:DR:78:TYR:CE1	1:DS:115:MET:HA	2.51	0.46
2:DT:122:PRO:O	2:DT:126:THR:HG23	2.16	0.46
8:DY:207:GLU:HG2	8:DY:207:GLU:O	2.15	0.46
9:EA:28:PHE:CD2	9:EA:39:LEU:HD23	2.51	0.46
9:EA:76:PRO:CB	9:EA:123:ALA:HB2	2.35	0.46
9:EA:252:PRO:HA	9:EA:271:GLY:CA	2.37	0.46
3:AE:39:ILE:CG2	3:AE:96:VAL:HG11	2.34	0.46
3:AE:131:THR:HG22	3:AE:157:ILE:CD1	2.45	0.46
2:AL:18:TYR:HD2	6:BD:61:THR:HG22	1.77	0.46
2:AL:90:ARG:NH1	6:BD:32:ASP:OD1	2.48	0.46
2:AO:123:ILE:O	2:AO:126:THR:HG22	2.16	0.46
2:AQ:144:ASP:OD1	2:AQ:144:ASP:N	2.48	0.46
1:AR:97:VAL:CG1	2:AS:27:LEU:HD13	2.46	0.46
2:AU:61:LEU:HD22	10:AV:200:CYC:CAA	2.46	0.46
2:AW:21:GLY:CA	2:AW:24:MET:HB3	2.42	0.46
6:BD:821:GLN:O	6:BD:825:ILE:HG13	2.15	0.46

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
9:BH:190:ILE:HG21	9:BH:193:VAL:HB	1.98	0.46
1:BI:126:VAL:CG2	10:BI:200:CYC:HMC3	2.44	0.46
2:BJ:83:ARG:HG2	2:BJ:87:TYR:CE2	2.50	0.46
3:BM:9:LEU:HD23	2:BN:1:MET:HE1	1.97	0.46
3:BM:43:LEU:CD1	3:BM:93:THR:HG22	2.45	0.46
2:BQ:2:GLN:O	2:BQ:100:ASP:HB3	2.15	0.46
2:BW:78:TYR:CD1	1:BX:115:MET:HG3	2.51	0.46
2:CA:62:TYR:CE2	7:CO:103:PRO:HG3	2.51	0.46
2:CA:102:SER:O	2:CA:106:GLU:HG2	2.16	0.46
2:CD:54:GLU:CD	2:CD:136:VAL:HG11	2.40	0.46
2:CD:88:TYR:OH	2:CD:112:LEU:HD13	2.15	0.46
6:CM:410:ASN:HA	6:CM:413:MET:HE2	1.96	0.46
1:CR:134:LYS:HA	1:CR:153:PHE:CE2	2.51	0.46
2:CS:19:LEU:HB2	2:CS:24:MET:HE3	1.97	0.46
1:CT:90:ARG:HD2	2:CU:17:LYS:C	2.41	0.46
2:CU:3:ASP:HB2	2:CU:98:ALA:O	2.15	0.46
1:CV:49:THR:O	1:CV:53:GLN:HG3	2.16	0.46
1:CY:74:GLY:O	1:CY:78:THR:HG23	2.15	0.46
2:DB:111:GLY:O	2:DB:115:THR:OG1	2.28	0.46
9:DI:26:ALA:O	9:DI:29:SER:HB3	2.15	0.46
1:DJ:48:GLU:HA	1:DJ:51:VAL:CG1	2.46	0.46
1:DJ:85:MET:HG2	1:DJ:133:MET:HE2	1.97	0.46
1:DJ:153:PHE:O	1:DJ:157:ILE:HG13	2.15	0.46
2:DK:19:LEU:HB3	2:DK:24:MET:CE	2.45	0.46
1:DL:4:VAL:HG23	2:DM:97:LEU:HD23	1.97	0.46
2:DO:44:ILE:HG21	2:DO:93:THR:OG1	2.16	0.46
1:DQ:101:VAL:HA	1:DQ:104:ILE:HD13	1.97	0.46
1:DS:50:ILE:CG1	1:DS:136:VAL:HG13	2.45	0.46
1:DU:52:LYS:HE2	1:DU:52:LYS:HA	1.96	0.46
1:DU:128:GLN:O	1:DU:132:GLU:HG2	2.15	0.46
2:DV:15:GLN:CB	2:DV:17:LYS:HG3	2.45	0.46
2:DV:149:MET:HE3	2:DV:153:LEU:CD1	2.46	0.46
8:DY:214:ASN:HB2	8:DY:217:ASN:ND2	2.31	0.46
8:DZ:210:LEU:HD21	8:DZ:217:ASN:N	2.31	0.46
9:EA:46:GLU:O	9:EA:49:LYS:HG2	2.15	0.46
2:AB:62:TYR:OH	10:AC:200:CYC:O2D	2.26	0.46
1:AC:25:ARG:HD3	6:BD:38:SER:O	2.15	0.46
1:AC:73:TYR:H	1:AC:77:MET:CB	2.26	0.46
2:AD:34:GLY:O	2:AD:38:VAL:HG23	2.16	0.46
3:AE:43:LEU:O	3:AE:47:GLU:HG3	2.16	0.46
1:AH:21:GLY:O	1:AH:25:ARG:HG3	2.16	0.46

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:AJ:27:LYS:HZ3	4:AK:35:SER:HA	1.81	0.46
1:AN:79:ALA:HB1	7:BF:110:PHE:HB3	1.98	0.46
1:AN:128:GLN:NE2	1:AN:132:GLU:OE2	2.49	0.46
10:AW:200:CYC:HB	6:BD:566:ILE:HD12	1.78	0.46
2:AY:62:TYR:H	2:AY:66:THR:HG21	1.81	0.46
6:BD:241:ARG:HG3	6:BD:389:TYR:OH	2.16	0.46
6:BD:347:PHE:O	6:BD:351:LEU:N	2.37	0.46
1:BI:37:LEU:HD13	2:BJ:28:LYS:NZ	2.30	0.46
1:BP:4:VAL:HG22	1:BP:26:ILE:HG23	1.98	0.46
1:BP:97:VAL:HG21	2:BQ:19:LEU:CD1	2.46	0.46
1:BP:97:VAL:HG21	2:BQ:19:LEU:HD13	1.97	0.46
2:BQ:53:LYS:HE2	1:BR:118:SER:O	2.16	0.46
4:BS:109:VAL:CG1	4:BS:113:LEU:HD11	2.46	0.46
1:BV:116:TYR:HD1	1:BV:121:THR:HB	1.80	0.46
2:CA:91:TYR:CE1	2:CA:107:ARG:HG3	2.51	0.46
2:CF:116:TYR:CE2	2:CF:160:LEU:HD21	2.51	0.46
6:CM:731:ASP:O	6:CM:735:VAL:HG23	2.15	0.46
6:CM:799:LYS:HD3	6:CM:883:LEU:CD2	2.45	0.46
1:CR:122:PRO:HG2	1:CR:125:ALA:HB3	1.98	0.46
2:CS:15:GLN:CB	2:CS:17:LYS:HE3	2.46	0.46
2:CS:51:ILE:CG1	2:CS:140:LEU:HD12	2.46	0.46
1:CT:109:LEU:HD12	1:CT:159:LYS:CB	2.46	0.46
2:CU:52:VAL:O	2:CU:56:VAL:HG12	2.16	0.46
2:CU:109:LEU:CD1	2:CU:159:GLY:HA3	2.45	0.46
2:CW:85:LEU:HD11	2:CW:129:ALA:HB1	1.97	0.46
1:CY:4:VAL:O	1:CY:8:ILE:HG12	2.16	0.46
1:CY:7:SER:HB3	1:CY:18:LEU:CD1	2.46	0.46
1:CY:57:ARG:HD2	1:CY:132:GLU:OE1	2.15	0.46
1:CY:83:ARG:HA	8:DY:218:PHE:CE2	2.51	0.46
1:CY:106:GLU:O	2:DD:76:ARG:NH1	2.48	0.46
2:CZ:97:LEU:HD23	2:CZ:97:LEU:O	2.15	0.46
1:DA:93:THR:HG22	2:DB:19:LEU:HD12	1.98	0.46
5:DF:4:PHE:O	5:DF:31:VAL:N	2.23	0.46
9:DI:79:GLN:CD	9:DI:122:VAL:HG23	2.41	0.46
9:DI:121:PHE:CA	9:DI:276:PRO:HG2	2.40	0.46
9:DI:146:GLU:HA	8:DZ:234:LYS:NZ	2.30	0.46
2:DK:2:GLN:O	2:DK:100:ASP:HB3	2.15	0.46
1:DL:12:ASP:OD1	2:DM:107:ARG:NH1	2.48	0.46
1:DL:109:LEU:HD22	1:DL:159:LYS:CB	2.45	0.46
1:DQ:47:ARG:O	1:DQ:51:VAL:HG12	2.16	0.46
2:DR:24:MET:HE3	2:DR:28:LYS:HZ1	1.80	0.46

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:DV:51:ILE:HG22	2:DV:133:ILE:CD1	2.40	0.46
1:AA:93:THR:O	1:AA:97:VAL:HG23	2.15	0.46
1:AA:123:ILE:HG13	1:AA:160:MET:O	2.15	0.46
1:AC:126:VAL:HG23	1:AC:160:MET:HE2	1.98	0.46
2:AF:105:ASP:HB2	2:AF:155:TYR:OH	2.16	0.46
2:AI:91:TYR:HB3	2:AI:104:LEU:HD23	1.98	0.46
1:AJ:107:ILE:HD13	4:AK:13:ASP:CG	2.41	0.46
4:AK:123:PRO:HG3	2:AS:64:ASP:OD2	2.16	0.46
2:AL:161:SER:O	2:AS:155:TYR:OH	2.28	0.46
1:AN:94:TYR:CG	2:AO:9:ILE:HG23	2.50	0.46
1:AP:16:ARG:O	2:AQ:94:TYR:OH	2.32	0.46
2:AU:108:VAL:O	2:AU:112:LEU:HG	2.16	0.46
1:AV:27:LYS:O	1:AV:31:THR:HG23	2.15	0.46
1:AV:36:ARG:HG2	1:AV:148:GLU:OE2	2.16	0.46
6:BD:447:TYR:OH	10:BD:901:CYC:HAA1	2.16	0.46
6:BD:485:ILE:HD12	6:BD:665:LEU:HD13	1.98	0.46
6:BD:796:PRO:O	6:BD:800:VAL:HG23	2.16	0.46
1:BK:123:ILE:HD12	1:BK:126:VAL:HG21	1.98	0.46
3:BM:99:GLY:O	3:BM:152:TYR:OH	2.30	0.46
2:BQ:20:ASP:OD1	2:BQ:22:ALA:N	2.49	0.46
2:BQ:71:ASN:HD22	2:BQ:122:PRO:CD	2.29	0.46
1:BV:115:MET:CG	2:CA:78:TYR:HB3	2.46	0.46
2:BW:63:SER:H	2:BW:66:THR:HG22	1.80	0.46
1:BX:26:ILE:O	1:BX:30:VAL:HG23	2.16	0.46
1:BZ:81:CYS:O	1:BZ:85:MET:HG3	2.16	0.46
2:CD:123:ILE:HA	2:CD:126:THR:CG2	2.45	0.46
2:CH:119:LEU:CD1	10:CH:200:CYC:HAA2	2.46	0.46
2:CH:126:THR:CB	10:CH:200:CYC:HMC1	2.46	0.46
6:CM:576:ARG:NH1	6:CM:646:SER:OG	2.43	0.46
6:CM:799:LYS:HE2	6:CM:803:MET:HE3	1.98	0.46
6:CM:806:LYS:HD3	6:CM:872:LEU:HD22	1.98	0.46
8:CP:233:PRO:O	8:CP:235:VAL:HG13	2.16	0.46
8:CP:234:LYS:HE2	8:CP:234:LYS:HA	1.97	0.46
9:CQ:40:ILE:HG13	11:CQ:400:45D:H371	1.97	0.46
1:CR:50:ILE:CD1	1:CR:136:VAL:HG12	2.27	0.46
1:CR:101:VAL:CG1	1:CY:20:PRO:HG2	2.45	0.46
1:CT:23:LEU:HD22	2:CU:38:VAL:HG13	1.96	0.46
1:CT:66:VAL:HG13	8:DY:233:PRO:HG3	1.98	0.46
2:CU:75:THR:HG23	1:CV:108:GLY:HA2	1.98	0.46
1:CV:113:ARG:HG2	1:CV:123:ILE:CD1	2.46	0.46
1:CY:37:LEU:CD2	2:CZ:28:LYS:HE3	2.42	0.46

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:CZ:76:ARG:HB2	1:DA:110:VAL:HB	1.98	0.46
8:DG:230:ASN:HD21	1:DL:68:PRO:HD2	1.80	0.46
8:DH:197:GLN:HG2	8:DH:198:ASN:N	2.31	0.46
1:DQ:2:SER:OG	1:DQ:4:VAL:HG12	2.15	0.46
1:DQ:50:ILE:HG21	1:DQ:89:LEU:HD11	1.97	0.46
1:DQ:76:GLU:HG2	8:DZ:203:PHE:H	1.81	0.46
9:EA:223:LEU:HA	9:EA:301:VAL:CG1	2.45	0.46
1:AA:137:ALA:O	1:AA:141:MET:HG2	2.16	0.46
1:AA:156:VAL:CG1	1:AA:160:MET:HE3	2.45	0.46
1:AC:90:ARG:O	1:AC:93:THR:OG1	2.28	0.46
1:AC:94:TYR:HA	1:AC:97:VAL:HG22	1.98	0.46
2:AD:15:GLN:CB	2:AD:17:LYS:HE2	2.46	0.46
2:AI:64:ASP:OD1	2:AO:124:SER:OG	2.08	0.46
4:AK:151:ASP:OD1	4:AK:151:ASP:N	2.46	0.46
10:AU:200:CYC:HB	10:AU:200:CYC:CMA	2.27	0.46
1:AZ:61:LYS:HG3	1:AZ:62:ARG:HG2	1.98	0.46
6:BD:284:ALA:HA	6:BD:287:ARG:CZ	2.46	0.46
6:BD:636:VAL:O	6:BD:640:ILE:HG12	2.15	0.46
6:BD:848:PHE:O	6:BD:852:THR:HB	2.16	0.46
9:BH:280:GLY:HA2	9:BH:310:LYS:NZ	2.31	0.46
1:BK:17:TYR:OH	2:BL:89:LEU:HD22	2.16	0.46
3:BM:7:VAL:HG13	3:BM:22:GLU:HB3	1.98	0.46
3:BM:94:TYR:CG	2:BN:9:ILE:HG23	2.51	0.46
2:BN:25:ASP:HA	2:BN:28:LYS:HE3	1.98	0.46
2:BN:52:VAL:O	2:BN:56:VAL:HG23	2.16	0.46
1:BP:64:ASP:HA	1:BP:67:SER:OG	2.16	0.46
1:BP:103:PRO:HA	1:BP:106:GLU:HG2	1.97	0.46
2:BQ:68:PRO:HG3	2:BQ:73:TYR:CE1	2.51	0.46
4:BS:14:LEU:HD13	2:BW:161:SER:HB3	1.97	0.46
1:BV:98:SER:HB3	2:BW:9:ILE:HD12	1.98	0.46
1:BX:27:LYS:HG3	2:BY:35:GLU:OE2	2.16	0.46
2:BY:107:ARG:HB3	6:CM:457:GLN:NE2	2.31	0.46
1:CE:88:TYR:CZ	1:CE:160:MET:HE1	2.51	0.46
2:CF:85:LEU:HG	10:CF:200:CYC:HBC1	1.98	0.46
2:CH:6:THR:HG23	6:CM:532:VAL:CG1	2.46	0.46
6:CM:2:SER:O	6:CM:4:LYS:NZ	2.41	0.46
6:CM:86:GLU:H	6:CM:153:LEU:CD2	2.29	0.46
6:CM:831:LEU:O	6:CM:835:ILE:HD12	2.16	0.46
10:CM:901:CYC:HBD1	10:CM:901:CYC:HHA	1.98	0.46
9:CQ:229:ARG:HG2	9:DI:227:PHE:HB3	1.98	0.46
1:CT:108:GLY:O	1:CT:109:LEU:HD22	2.16	0.46

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:CV:23:LEU:O	1:CV:27:LYS:HG2	2.16	0.46
1:CV:98:SER:HA	2:CW:5:ILE:HG21	1.97	0.46
2:CZ:68:PRO:HG3	2:CZ:73:TYR:CZ	2.50	0.46
2:DB:12:ALA:O	2:DB:15:GLN:HG2	2.15	0.46
2:DK:62:TYR:OH	10:DL:200:CYC:O2D	2.14	0.46
2:DO:71:ASN:HD22	2:DO:122:PRO:HD2	1.81	0.46
1:DQ:3:ILE:O	1:DQ:6:LYS:HG2	2.16	0.46
1:DQ:68:PRO:HA	1:DQ:73:TYR:CD2	2.50	0.46
1:DQ:105:GLU:CA	1:DQ:109:LEU:HD12	2.45	0.46
1:DQ:114:GLU:OE1	1:DQ:117:ARG:NE	2.40	0.46
2:DR:112:LEU:HD13	10:DR:200:CYC:HMB3	1.97	0.46
1:DS:131:ARG:HA	1:DS:157:ILE:HD11	1.97	0.46
1:DU:39:ILE:CG2	1:DU:96:VAL:HG11	2.46	0.46
9:EA:39:LEU:HD13	9:EA:139:LEU:HB2	1.97	0.46
9:EA:112:ARG:O	9:EA:116:LEU:HG	2.16	0.46
1:AA:116:TYR:HE2	1:AA:126:VAL:HG11	1.80	0.46
1:AC:126:VAL:O	1:AC:130:VAL:HG23	2.15	0.46
3:AE:110:ILE:HG12	5:BA:17:ARG:HH11	1.81	0.46
2:AI:71:ASN:HD22	2:AI:121:VAL:HA	1.81	0.46
1:AP:27:LYS:HG3	2:AQ:35:GLU:OE2	2.16	0.46
1:AP:66:VAL:CG1	1:DU:68:PRO:HG2	2.46	0.46
2:AU:54:GLU:CD	2:AU:136:VAL:HG21	2.41	0.46
2:AU:75:THR:OG1	1:AV:110:VAL:O	2.28	0.46
10:AU:200:CYC:HAD2	5:BB:33:TYR:OH	2.16	0.46
1:AV:27:LYS:HZ3	2:AW:38:VAL:HB	1.80	0.46
2:AW:108:VAL:HG12	2:AW:109:LEU:HD23	1.97	0.46
1:AX:30:VAL:HG13	2:AY:31:PHE:HA	1.98	0.46
2:AY:147:LYS:O	2:AY:151:VAL:HG23	2.16	0.46
1:AZ:132:GLU:O	1:AZ:136:VAL:HG23	2.15	0.46
5:BA:59:ARG:O	5:BA:62:THR:HG23	2.16	0.46
6:BD:484:LEU:HD12	6:BD:615:TYR:HE1	1.81	0.46
6:BD:842:MET:HE1	5:DF:46:MET:CG	2.46	0.46
7:BF:105:PRO:HA	7:BF:108:ASN:OD1	2.16	0.46
9:BH:89:ARG:HD3	9:BH:170:LYS:CE	2.46	0.46
9:BH:286:ILE:HG12	9:BH:305:LEU:CD2	2.45	0.46
1:BK:117:ARG:HD2	1:BP:161:SER:CA	2.46	0.46
1:BP:86:ASP:OD1	2:BQ:18:TYR:OH	2.30	0.46
4:BS:108:ARG:HB3	6:CM:249:GLN:OE1	2.16	0.46
1:BV:90:ARG:HG3	2:BW:18:TYR:CD1	2.51	0.46
1:BX:74:GLY:O	1:BX:78:THR:HG23	2.16	0.46
2:CA:24:MET:O	2:CA:28:LYS:HG3	2.16	0.46

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:CE:134:LYS:HE2	1:CE:154:ASP:OD1	2.16	0.46
2:CF:93:THR:CA	2:CF:149:MET:HE1	2.46	0.46
6:CM:453:PRO:HD2	6:CM:607:ARG:CZ	2.46	0.46
6:CM:611:GLY:O	6:CM:658:ARG:N	2.46	0.46
6:CM:681:GLN:HG2	6:CM:682:ARG:N	2.30	0.46
6:CM:845:LEU:CG	5:DX:42:ARG:HD2	2.37	0.46
6:CM:891:VAL:CG2	2:DT:106:GLU:HB3	2.46	0.46
9:CQ:98:TYR:OH	9:CQ:158:VAL:O	2.34	0.46
9:CQ:170:LYS:HD3	9:CQ:170:LYS:HA	1.77	0.46
1:CR:7:SER:HG	1:CR:25:ARG:HH11	1.57	0.46
1:CR:25:ARG:HB2	1:CY:25:ARG:HG3	1.96	0.46
2:CW:34:GLY:O	2:CW:38:VAL:HG23	2.16	0.46
1:CY:77:MET:HE2	8:DH:201:ARG:O	2.15	0.46
2:CZ:130:ILE:CG2	2:CZ:153:LEU:HD12	2.33	0.46
2:DD:114:GLU:OE1	2:DD:114:GLU:N	2.43	0.46
9:DI:74:MET:CE	9:DI:78:GLN:HB2	2.46	0.46
1:DJ:36:ARG:NH1	1:DJ:152:TYR:OH	2.49	0.46
1:DN:96:VAL:HA	1:DN:152:TYR:CE2	2.51	0.46
2:DR:5:ILE:HD11	2:DR:31:PHE:HE1	1.80	0.46
2:DR:73:TYR:O	2:DR:77:ARG:HB2	2.16	0.46
2:DV:133:ILE:O	2:DV:137:THR:HG22	2.15	0.46
9:EA:107:LEU:CD1	9:EA:158:VAL:HG11	2.45	0.46
1:AA:115:MET:HE1	10:AA:200:CYC:CMB	2.40	0.45
2:AB:126:THR:O	2:AB:130:ILE:HG12	2.15	0.45
2:AD:104:LEU:HB3	2:AD:109:LEU:HD13	1.98	0.45
3:AE:87:TRP:O	3:AE:91:LEU:HD13	2.16	0.45
2:AF:47:ASN:HB2	2:AF:140:LEU:HD21	1.97	0.45
1:AH:156:VAL:CG1	1:AH:160:MET:HE3	2.46	0.45
1:AJ:38:ARG:NH1	1:AJ:145:ASP:OD2	2.46	0.45
2:AL:108:VAL:O	6:BD:420:TYR:OH	2.29	0.45
1:AN:115:MET:HG2	2:AS:78:TYR:HB3	1.96	0.45
1:AP:101:VAL:HB	1:AP:155:PHE:CE2	2.51	0.45
2:AU:148:GLU:HG3	2:AU:152:TYR:CE2	2.50	0.45
1:AV:72:ALA:HB2	10:AV:200:CYC:HMD1	1.97	0.45
2:AY:96:MET:HB2	2:AY:149:MET:HE2	1.97	0.45
6:BD:597:TYR:HB3	6:BD:600:LYS:CD	2.43	0.45
1:BK:88:TYR:O	1:BK:92:VAL:HG23	2.16	0.45
2:BL:95:ALA:HB2	2:BL:104:LEU:HD21	1.98	0.45
3:BM:43:LEU:O	3:BM:47:GLU:HG3	2.16	0.45
1:BR:3:ILE:HG21	1:BR:25:ARG:CZ	2.46	0.45
2:BT:44:ILE:HD13	2:BT:149:MET:HE2	1.98	0.45

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:BT:56:VAL:HG21	2:BT:82:ILE:CD1	2.46	0.45
1:BX:59:PHE:CD1	1:BX:66:VAL:HG21	2.50	0.45
2:BY:119:LEU:HD11	10:BY:200:CYC:HAA2	1.98	0.45
1:BZ:36:ARG:NH1	1:BZ:99:GLY:HA2	2.30	0.45
1:CC:81:CYS:HA	10:CC:200:CYC:CHD	2.41	0.45
1:CE:95:GLY:HA3	1:CE:104:ILE:HD11	1.97	0.45
1:CE:100:ASP:OD1	1:CE:101:VAL:N	2.49	0.45
2:CF:96:MET:HA	2:CF:152:TYR:CE2	2.51	0.45
6:CM:64:ALA:O	6:CM:68:VAL:HG22	2.16	0.45
6:CM:514:ARG:HA	6:CM:652:ASP:O	2.16	0.45
6:CM:777:PHE:O	6:CM:781:LEU:HG	2.14	0.45
9:CQ:51:LEU:HD23	9:CQ:150:GLN:NE2	2.30	0.45
9:CQ:252:PRO:HA	9:CQ:271:GLY:CA	2.33	0.45
1:DA:39:ILE:CD1	1:DA:145:ASP:HB3	2.46	0.45
2:DB:81:CYS:SG	10:DB:200:CYC:HMD3	2.56	0.45
2:DD:126:THR:O	2:DD:130:ILE:HG13	2.15	0.45
8:DH:203:PHE:CE2	8:DY:219:LEU:HD22	2.51	0.45
9:DI:40:ILE:O	11:DI:400:45D:H401	2.16	0.45
10:DJ:200:CYC:HHA	10:DJ:200:CYC:HBD1	1.99	0.45
1:DL:126:VAL:O	1:DL:130:VAL:HG23	2.16	0.45
2:DM:15:GLN:HG3	2:DM:17:LYS:HZ3	1.81	0.45
10:DO:200:CYC:HB	10:DO:200:CYC:HMA1	1.80	0.45
2:DV:51:ILE:HA	2:DV:54:GLU:OE2	2.16	0.45
5:DX:7:THR:HA	5:DX:27:PHE:O	2.16	0.45
1:AA:102:THR:HB	1:AA:103:PRO:HD3	1.99	0.45
2:AB:96:MET:HG3	2:AB:148:GLU:HG3	1.98	0.45
2:AD:83:ARG:CZ	5:BA:22:LEU:HD21	2.46	0.45
10:AD:200:CYC:HAA1	5:BA:25:THR:HG21	1.97	0.45
2:AI:117:ASN:O	6:BD:439:ARG:NH1	2.49	0.45
1:AJ:87:TYR:O	1:AJ:91:LEU:HG	2.16	0.45
1:AN:15:ALA:HB3	6:BD:498:PRO:HG3	1.98	0.45
1:AN:39:ILE:HG23	1:AN:141:MET:HE1	1.98	0.45
1:AN:91:LEU:CD2	1:AN:107:ILE:HB	2.45	0.45
1:AP:102:THR:HB	1:AP:103:PRO:HD3	1.98	0.45
2:AU:91:TYR:HE2	10:AU:200:CYC:HBB1	1.81	0.45
2:AU:123:ILE:HA	2:AU:126:THR:CG2	2.46	0.45
2:AW:36:LEU:HA	2:AW:39:ARG:NH1	2.30	0.45
1:AX:109:LEU:HD21	1:AX:159:LYS:CB	2.46	0.45
6:BD:413:MET:HE2	6:BD:433:THR:CG2	2.46	0.45
2:BL:40:ALA:HB3	2:BL:96:MET:CE	2.45	0.45
3:BM:9:LEU:CD2	2:BN:1:MET:HE1	2.47	0.45

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:BM:72:ALA:HA	3:BM:77:GLN:CB	2.45	0.45
3:BM:124:PRO:O	3:BM:127:VAL:HG12	2.16	0.45
1:BP:50:ILE:HA	1:BP:136:VAL:HG11	1.97	0.45
2:BT:25:ASP:HA	2:BT:28:LYS:NZ	2.32	0.45
1:BV:57:ARG:O	1:BV:60:GLN:HG3	2.17	0.45
2:CD:47:ASN:O	2:CD:51:ILE:HD12	2.16	0.45
2:CD:137:THR:CG2	2:CD:149:MET:HG2	2.38	0.45
1:CE:111:GLY:HA2	1:CE:114:GLU:OE1	2.15	0.45
2:CH:84:ASP:OD2	2:CH:116:TYR:OH	2.17	0.45
6:CM:84:TYR:HD2	6:CM:150:GLN:HE21	1.63	0.45
6:CM:210:PHE:CZ	6:CM:216:ALA:HB1	2.51	0.45
6:CM:355:PRO:HB3	6:CM:361:VAL:HG22	1.98	0.45
9:CQ:168:ASP:N	9:CQ:168:ASP:OD1	2.49	0.45
2:CS:75:THR:HG23	1:CT:110:VAL:O	2.15	0.45
2:CU:25:ASP:HA	2:CU:28:LYS:NZ	2.30	0.45
2:CU:71:ASN:ND2	2:CU:121:VAL:HA	2.31	0.45
1:CV:113:ARG:HG2	1:CV:123:ILE:HD13	1.98	0.45
2:CW:49:ALA:HB3	1:DA:161:SER:OG	2.17	0.45
2:DD:56:VAL:HG21	2:DD:82:ILE:CD1	2.46	0.45
9:DI:103:PRO:HB3	9:DI:159:VAL:HG22	1.98	0.45
9:DI:140:ALA:HA	9:DI:143:GLN:NE2	2.23	0.45
9:DI:188:VAL:O	9:DI:199:LEU:HD22	2.17	0.45
2:DK:124:SER:O	2:DK:128:GLN:HG3	2.16	0.45
1:DQ:40:ALA:HB2	1:DQ:97:VAL:CG2	2.46	0.45
1:DQ:88:TYR:HD1	1:DQ:91:LEU:HD12	1.79	0.45
2:DT:1:MET:HE3	2:DT:103:ILE:HB	1.98	0.45
1:AC:16:ARG:HG2	1:AC:17:TYR:O	2.15	0.45
2:AD:79:ALA:HA	2:AD:82:ILE:HG12	1.98	0.45
10:AD:200:CYC:CAA	5:BA:11:PRO:HB3	2.45	0.45
3:AE:50:ILE:CG1	3:AE:140:LEU:HD13	2.46	0.45
2:AF:2:GLN:HE22	2:AF:7:ALA:HB2	1.81	0.45
2:AL:18:TYR:OH	6:BD:160:LEU:HD23	2.16	0.45
2:AL:37:ARG:HD2	2:AL:96:MET:O	2.15	0.45
1:AP:27:LYS:O	1:AP:31:THR:HG23	2.16	0.45
1:AP:122:PRO:HG2	1:AP:125:ALA:CB	2.46	0.45
2:AS:88:TYR:OH	2:AS:112:LEU:HD21	2.17	0.45
2:AS:132:ALA:HA	2:AS:135:GLU:OE1	2.15	0.45
2:AW:125:SER:HA	2:AW:128:GLN:NE2	2.31	0.45
1:AZ:101:VAL:HB	1:AZ:155:PHE:CE2	2.51	0.45
1:AZ:116:TYR:CD2	1:AZ:126:VAL:HG21	2.52	0.45
1:AZ:126:VAL:O	1:AZ:130:VAL:HG23	2.16	0.45

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
5:BA:23:GLN:NE2	6:BD:390:PHE:O	2.35	0.45
6:BD:77:THR:CG2	6:BD:138:PRO:HB2	2.45	0.45
6:BD:151:LYS:CD	6:BD:154:ARG:HH21	2.29	0.45
6:BD:597:TYR:CE2	6:BD:599:CYS:HB2	2.51	0.45
9:BH:308:SER:HB2	9:BH:309:PRO:HD2	1.98	0.45
1:BK:14:GLU:HG3	1:BK:16:ARG:HD2	1.98	0.45
2:BL:96:MET:HA	2:BL:152:TYR:CE2	2.52	0.45
10:BL:200:CYC:OB	5:CJ:24:ASN:HA	2.15	0.45
3:BM:30:LEU:HD12	2:BN:31:PHE:HA	1.96	0.45
2:BN:2:GLN:NE2	2:BN:7:ALA:HA	2.31	0.45
10:BV:200:CYC:HBD1	10:BV:200:CYC:HHA	1.97	0.45
1:CC:111:GLY:HA2	1:CC:114:GLU:OE1	2.16	0.45
1:CE:92:VAL:HG11	1:CE:153:PHE:CE2	2.51	0.45
6:CM:21:PRO:O	6:CM:25:ILE:HD12	2.17	0.45
6:CM:73:ASN:OD1	6:CM:81:PRO:HD2	2.16	0.45
9:CQ:79:GLN:OE1	9:CQ:122:VAL:HA	2.17	0.45
2:CS:44:ILE:HG21	2:CS:93:THR:OG1	2.17	0.45
1:DA:153:PHE:O	1:DA:157:ILE:HG12	2.16	0.45
1:DC:41:GLU:OE2	2:DD:24:MET:HG2	2.16	0.45
9:DI:253:GLU:OE1	9:DI:254:ARG:NH1	2.49	0.45
9:DI:300:PHE:CZ	9:DI:302:ALA:HB2	2.52	0.45
2:DT:51:ILE:HG22	2:DT:133:ILE:CD1	2.46	0.45
2:DV:64:ASP:HA	2:DV:67:ARG:HE	1.81	0.45
2:DV:68:PRO:HG3	2:DV:73:TYR:CZ	2.51	0.45
1:AA:48:GLU:O	1:AA:51:VAL:HG12	2.16	0.45
1:AA:61:LYS:HE2	1:DC:57:ARG:NH2	2.32	0.45
1:AA:109:LEU:CG	1:AA:159:LYS:HG2	2.45	0.45
2:AD:68:PRO:HD3	3:AE:87:TRP:CH2	2.51	0.45
3:AE:81:CYS:HA	10:AE:200:CYC:HAC1	1.86	0.45
10:AH:200:CYC:CGA	2:AL:66:THR:HG21	2.46	0.45
2:AI:74:THR:OG1	2:AI:77:ARG:HB2	2.16	0.45
1:AJ:141:MET:HE2	1:AJ:141:MET:HB3	1.81	0.45
4:AK:142:ALA:CB	4:AK:147:LEU:HD12	2.47	0.45
2:AS:62:TYR:O	7:BF:101:ARG:NH2	2.49	0.45
2:AW:106:GLU:OE1	2:AW:107:ARG:HG3	2.16	0.45
1:AX:82:LEU:HD13	8:BG:242:ILE:HD11	1.99	0.45
2:AY:10:ASN:OD1	6:BD:532:VAL:HG23	2.16	0.45
2:AY:51:ILE:HG21	2:AY:137:THR:HG22	1.98	0.45
2:AY:133:ILE:O	2:AY:137:THR:HG23	2.17	0.45
5:BA:8:ALA:CB	5:BA:50:ILE:HG22	2.45	0.45
6:BD:240:VAL:HG22	6:BD:252:GLU:CG	2.39	0.45

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
6:BD:294:ILE:HD12	6:BD:328:LEU:HD21	1.98	0.45
6:BD:445:HIS:HE1	6:BD:620:MET:HE2	1.81	0.45
6:BD:598:VAL:H	10:BD:901:CYC:HBB3	1.81	0.45
6:BD:863:ALA:HB2	2:DB:107:ARG:O	2.16	0.45
2:BJ:113:LYS:HE2	2:BJ:117:ASN:OD1	2.17	0.45
2:BL:41:ALA:HB2	2:BL:96:MET:SD	2.56	0.45
2:BL:106:GLU:CD	6:CM:273:LEU:HD23	2.41	0.45
2:BQ:44:ILE:HG21	2:BQ:93:THR:CG2	2.46	0.45
2:BT:34:GLY:HA2	2:BT:37:ARG:HH21	1.80	0.45
2:BW:56:VAL:HG21	2:BW:82:ILE:CD1	2.46	0.45
2:BW:93:THR:HA	2:BW:149:MET:HE1	1.99	0.45
10:CD:200:CYC:HAA1	5:CK:36:TRP:HD1	1.82	0.45
1:CE:102:THR:HB	1:CE:103:PRO:HD3	1.99	0.45
2:CF:36:LEU:HD22	2:CF:39:ARG:HH12	1.81	0.45
1:CR:22:GLU:HG3	1:CR:25:ARG:NH1	2.32	0.45
1:CR:61:LYS:HE2	1:CR:61:LYS:HA	1.98	0.45
2:CS:126:THR:HG21	2:CS:160:LEU:HD13	1.99	0.45
1:CT:38:ARG:HA	1:CT:41:GLU:OE1	2.16	0.45
10:CW:200:CYC:HMA2	5:DF:37:PHE:CD1	2.51	0.45
1:DA:85:MET:HE3	10:DA:200:CYC:HBC1	1.98	0.45
9:DI:17:ALA:HB3	9:DI:20:VAL:CG1	2.41	0.45
9:DI:37:LEU:HA	9:DI:40:ILE:HG12	1.99	0.45
9:DI:42:PHE:CD1	9:DI:45:LEU:HD12	2.52	0.45
9:DI:103:PRO:HB3	9:DI:159:VAL:CG2	2.47	0.45
1:DJ:141:MET:HE3	1:DJ:146:ALA:HA	1.98	0.45
2:DO:114:GLU:HG2	5:DX:53:VAL:HG12	1.98	0.45
1:DQ:47:ARG:C	1:DQ:50:ILE:HG22	2.42	0.45
2:DT:112:LEU:HD13	10:DT:200:CYC:CMB	2.46	0.45
9:EA:117:MET:SD	11:EA:400:45D:H392	2.57	0.45
2:AB:96:MET:SD	2:AB:149:MET:HE2	2.57	0.45
2:AD:95:ALA:HB2	2:AD:104:LEU:HD23	1.98	0.45
1:AH:80:THR:HG23	1:AH:83:ARG:HH21	1.82	0.45
4:AK:51:ILE:HD13	4:AK:138:ILE:HD11	1.97	0.45
2:AL:124:SER:O	2:AL:128:GLN:HG3	2.16	0.45
2:AO:116:TYR:HB3	2:AO:121:VAL:HB	1.99	0.45
1:AP:19:SER:OG	1:AP:20:PRO:HD2	2.17	0.45
1:AR:134:LYS:HB2	1:AR:153:PHE:HB3	1.98	0.45
2:AU:25:ASP:OD1	2:AU:26:LYS:N	2.49	0.45
2:AW:137:THR:O	2:AW:141:VAL:HG22	2.16	0.45
1:AZ:57:ARG:HD2	1:AZ:132:GLU:OE1	2.16	0.45
5:BA:6:ILE:HD13	5:BA:36:TRP:HZ3	1.80	0.45

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
6:BD:12:ALA:C	6:BD:13:ARG:HD3	2.42	0.45
6:BD:131:PRO:HD2	6:BD:200:GLN:HB3	1.98	0.45
6:BD:191:SER:HB3	6:BD:194:ALA:HB3	1.98	0.45
6:BD:210:PHE:C	6:BD:211:ARG:HG3	2.42	0.45
8:BG:234:LYS:HA	8:BG:234:LYS:HE2	1.97	0.45
8:BG:237:ALA:HA	8:BG:240:ILE:CD1	2.38	0.45
9:BH:74:MET:CE	9:BH:78:GLN:HB2	2.45	0.45
9:BH:238:LEU:HD12	9:BH:242:ARG:NH2	2.32	0.45
1:BK:116:TYR:CE2	1:BK:126:VAL:HG11	2.52	0.45
2:BL:34:GLY:O	2:BL:38:VAL:HG23	2.16	0.45
2:BL:95:ALA:HB1	2:BL:152:TYR:CD1	2.51	0.45
3:BM:32:THR:O	3:BM:36:ARG:HG2	2.16	0.45
2:BQ:83:ARG:HD2	6:CM:369:SER:HB3	1.99	0.45
1:BR:108:GLY:O	1:BR:112:VAL:HG11	2.17	0.45
1:BZ:36:ARG:NH1	1:BZ:96:VAL:O	2.44	0.45
1:BZ:79:ALA:HB2	8:CP:219:LEU:HA	1.98	0.45
2:CA:131:GLN:O	2:CA:134:LYS:HB2	2.17	0.45
1:CG:85:MET:C	1:CG:133:MET:HE1	2.41	0.45
1:CG:130:VAL:HG12	1:CG:157:ILE:HD11	1.98	0.45
6:CM:355:PRO:CB	6:CM:361:VAL:HG22	2.47	0.45
6:CM:446:VAL:HB	6:CM:617:ARG:HG3	1.98	0.45
9:CQ:71:ILE:HD13	9:CQ:74:MET:SD	2.57	0.45
9:CQ:91:ASP:OD1	9:CQ:96:ARG:NH2	2.46	0.45
9:CQ:97:THR:HG22	9:CQ:101:TRP:CZ2	2.50	0.45
9:CQ:139:LEU:HD23	9:CQ:143:GLN:OE1	2.17	0.45
9:CQ:201:TYR:CD2	9:CQ:213:LEU:HD11	2.52	0.45
9:CQ:239:ARG:HH21	1:DA:41:GLU:HB3	1.80	0.45
1:CR:23:LEU:CD1	2:CS:38:VAL:HG13	2.36	0.45
1:CT:109:LEU:O	1:CT:112:VAL:HG12	2.17	0.45
1:CV:108:GLY:C	1:CV:109:LEU:HD22	2.42	0.45
1:CY:23:LEU:HD12	1:CY:24:ASP:N	2.32	0.45
1:CY:130:VAL:HB	1:CY:157:ILE:HG12	1.98	0.45
2:CZ:41:ALA:HB2	2:CZ:96:MET:HE1	1.98	0.45
2:DK:149:MET:O	2:DK:153:LEU:HG	2.17	0.45
1:DL:21:GLY:O	1:DL:25:ARG:HG2	2.15	0.45
1:DN:2:SER:N	1:DN:100:ASP:OD2	2.49	0.45
2:DO:1:MET:CG	2:DO:103:ILE:HB	2.46	0.45
2:DR:21:GLY:HA2	2:DR:24:MET:HB2	1.99	0.45
2:DR:44:ILE:HG21	2:DR:93:THR:OG1	2.16	0.45
1:DS:91:LEU:HD21	1:DS:107:ILE:CG2	2.47	0.45
1:DU:27:LYS:NZ	2:DV:35:GLU:HA	2.31	0.45

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:AA:75:GLU:OE1	1:AA:75:GLU:N	2.45	0.45
1:AC:134:LYS:NZ	1:AC:154:ASP:HB3	2.31	0.45
3:AE:104:ILE:HG22	3:AE:109:LEU:CD1	2.47	0.45
2:AF:61:LEU:HD21	2:AF:78:TYR:OH	2.16	0.45
1:AH:77:MET:SD	10:AH:200:CYC:HAD1	2.57	0.45
1:AJ:108:GLY:O	1:AJ:112:VAL:HG11	2.17	0.45
4:AK:52:VAL:HG13	4:AK:86:MET:HB3	1.99	0.45
1:AN:9:VAL:HG23	2:AO:1:MET:CE	2.38	0.45
1:AV:83:ARG:HD2	1:AV:87:TYR:OH	2.17	0.45
1:AV:93:THR:O	1:AV:97:VAL:HG13	2.16	0.45
5:BB:55:LEU:HG	5:BB:57:THR:O	2.17	0.45
6:BD:253:LEU:HD11	6:BD:257:TYR:CG	2.51	0.45
6:BD:305:LEU:O	6:BD:309:VAL:HG23	2.17	0.45
6:BD:462:PHE:CZ	6:BD:600:LYS:HB3	2.52	0.45
6:BD:725:LYS:HG2	6:BD:852:THR:OG1	2.17	0.45
9:BH:37:LEU:CD1	9:BH:40:ILE:HD11	2.46	0.45
10:BI:200:CYC:HBC2	10:BI:200:CYC:H2C	1.67	0.45
2:BJ:54:GLU:O	2:BJ:58:LYS:HG2	2.17	0.45
2:BJ:102:SER:O	2:BJ:106:GLU:HG2	2.17	0.45
1:BK:90:ARG:HB2	2:BL:18:TYR:CE1	2.50	0.45
2:BN:55:ALA:CB	2:BN:133:ILE:HD13	2.46	0.45
1:BP:65:ILE:HG23	10:BP:200:CYC:OC	2.17	0.45
2:BT:106:GLU:CG	2:BT:107:ARG:HG2	2.40	0.45
2:BW:25:ASP:HA	2:BW:28:LYS:HZ3	1.82	0.45
1:BX:44:THR:HG23	2:BY:18:TYR:HB3	1.98	0.45
1:BX:98:SER:OG	1:BX:103:PRO:HG2	2.17	0.45
1:BX:134:LYS:HD3	1:BX:153:PHE:HB3	1.98	0.45
2:CA:72:MET:HE2	2:CA:81:CYS:SG	2.57	0.45
1:CE:126:VAL:O	1:CE:129:SER:OG	2.29	0.45
2:CH:96:MET:HG3	2:CH:148:GLU:OE1	2.17	0.45
6:CM:18:GLN:CB	6:CM:173:ASN:HD22	2.28	0.45
9:CQ:88:ASN:OD1	9:CQ:174:GLU:N	2.50	0.45
1:CR:94:TYR:O	1:CR:97:VAL:HG12	2.17	0.45
1:CR:94:TYR:OH	2:CS:17:LYS:O	2.29	0.45
10:CR:200:CYC:HAD2	2:CW:62:TYR:OH	2.17	0.45
2:CS:76:ARG:HH22	10:CS:200:CYC:CGD	2.30	0.45
1:CV:23:LEU:HB3	2:CW:38:VAL:HG13	1.99	0.45
2:CW:57:ALA:HA	2:CW:61:LEU:HB2	1.98	0.45
2:CZ:78:TYR:CE2	1:DA:115:MET:HG3	2.51	0.45
1:DA:97:VAL:HG11	2:DB:19:LEU:CD1	2.45	0.45
5:DF:1:MET:HE2	5:DF:32:PRO:HB3	1.99	0.45

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
9:DI:55:ALA:H	11:DI:400:45D:H221	1.81	0.45
9:DI:217:PHE:CB	9:DI:298:ILE:HB	2.47	0.45
1:DJ:156:VAL:O	1:DJ:160:MET:HG2	2.16	0.45
2:DM:6:THR:O	2:DM:10:ASN:ND2	2.50	0.45
2:DO:134:LYS:HG3	2:DO:150:GLY:HA2	1.97	0.45
2:DR:103:ILE:O	2:DR:107:ARG:HB2	2.16	0.45
1:DS:155:PHE:O	1:DS:159:LYS:HG2	2.17	0.45
2:DT:78:TYR:O	2:DT:82:ILE:HG12	2.17	0.45
1:DU:87:TYR:O	1:DU:91:LEU:HG	2.17	0.45
9:EA:149:GLN:O	9:EA:153:VAL:HG23	2.16	0.45
9:EA:274:GLN:N	9:EA:274:GLN:OE1	2.50	0.45
1:AA:67:SER:HB3	1:AA:68:PRO:HD2	1.99	0.45
1:AC:23:LEU:HD22	2:AD:38:VAL:HG13	1.99	0.45
1:AH:80:THR:CG2	1:AH:83:ARG:HH21	2.29	0.45
4:AK:41:ALA:HB1	4:AK:98:ILE:HD11	1.97	0.45
4:AK:64:PRO:HB2	7:BF:105:PRO:HB2	1.98	0.45
4:AK:67:ILE:HG21	10:BD:902:CYC:HBA1	1.99	0.45
2:AL:31:PHE:HD2	6:BD:54:LEU:HD11	1.82	0.45
2:AO:77:ARG:CB	10:AO:200:CYC:HMD1	2.47	0.45
1:AR:126:VAL:HG22	10:AR:200:CYC:HMC2	1.99	0.45
1:AX:46:SER:O	1:AX:50:ILE:HG22	2.16	0.45
6:BD:515:LEU:HG	6:BD:542:LEU:CD2	2.46	0.45
6:BD:575:VAL:O	6:BD:579:ILE:HG12	2.15	0.45
6:BD:625:ASP:OD1	6:BD:629:LYS:HE2	2.17	0.45
6:BD:785:GLU:HB3	6:BD:789:LYS:NZ	2.32	0.45
6:BD:818:GLU:HG2	6:BD:822:TYR:CE2	2.52	0.45
9:BH:139:LEU:HD23	9:BH:143:GLN:OE1	2.17	0.45
1:BI:51:VAL:HG23	1:BI:133:MET:CE	2.47	0.45
1:BI:119:LEU:HA	2:BN:53:LYS:NZ	2.30	0.45
1:BV:65:ILE:CG2	1:BV:72:ALA:HB3	2.45	0.45
2:BW:78:TYR:CE1	1:BX:115:MET:HG3	2.52	0.45
1:CC:85:MET:SD	1:CC:129:SER:HB2	2.57	0.45
1:CE:122:PRO:HG2	1:CE:125:ALA:CB	2.47	0.45
1:CE:129:SER:O	1:CE:133:MET:HG3	2.15	0.45
2:CF:47:ASN:CG	2:CF:140:LEU:HD21	2.42	0.45
2:CF:64:ASP:HA	2:CF:67:ARG:HD2	1.99	0.45
2:CF:72:MET:HE3	2:CF:81:CYS:HB3	1.97	0.45
2:CH:24:MET:HE3	2:CH:28:LYS:HD3	1.98	0.45
6:CM:576:ARG:HG3	6:CM:645:TYR:HE1	1.80	0.45
1:DA:56:ASP:O	1:DA:60:GLN:HG3	2.16	0.45
1:DA:83:ARG:NH2	1:DA:84:ASP:OD1	2.27	0.45

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:DB:127:VAL:CG1	2:DB:131:GLN:HE22	2.30	0.45
1:DC:115:MET:HE1	10:DC:200:CYC:CMB	2.47	0.45
2:DD:119:LEU:CD1	10:DD:200:CYC:HBD1	2.47	0.45
9:DI:223:LEU:HB3	9:DI:231:ILE:CG2	2.46	0.45
2:DK:41:ALA:HA	2:DK:96:MET:HE1	1.99	0.45
1:DL:59:PHE:CE1	1:DL:65:ILE:HD11	2.52	0.45
1:DQ:37:LEU:HG	2:DR:28:LYS:CE	2.47	0.45
1:DQ:85:MET:HG3	1:DQ:133:MET:HE2	1.99	0.45
1:DQ:157:ILE:HG13	1:DQ:158:GLY:N	2.31	0.45
8:DZ:237:ALA:HA	8:DZ:240:ILE:CD1	2.39	0.45
2:AF:4:ALA:HB2	2:AF:99:GLY:C	2.42	0.45
1:AH:84:ASP:HB2	10:AH:200:CYC:HBC3	1.99	0.45
4:AK:76:THR:HG23	10:BD:902:CYC:C4B	2.47	0.45
4:AK:105:LEU:O	4:AK:109:VAL:HB	2.17	0.45
1:AN:115:MET:CG	2:AS:78:TYR:HB3	2.46	0.45
2:AO:78:TYR:CD1	1:AP:115:MET:HG3	2.51	0.45
1:AR:124:GLU:OE1	1:AR:124:GLU:N	2.40	0.45
2:AU:51:ILE:HG22	2:AU:133:ILE:HG23	1.99	0.45
1:AV:113:ARG:CD	1:AV:123:ILE:HD13	2.41	0.45
2:AW:57:ALA:HA	2:AW:61:LEU:CD1	2.40	0.45
2:AY:88:TYR:OH	2:AY:112:LEU:HD21	2.17	0.45
1:AZ:11:ALA:HA	1:AZ:16:ARG:NH1	2.32	0.45
5:BB:6:ILE:O	5:BB:28:THR:HA	2.16	0.45
6:BD:139:ILE:HD11	6:BD:191:SER:CB	2.47	0.45
6:BD:351:LEU:HG	6:BD:380:LEU:HD22	1.99	0.45
6:BD:559:GLN:HB3	6:BD:587:LEU:HD22	1.97	0.45
6:BD:577:GLU:OE1	6:BD:577:GLU:N	2.49	0.45
1:BI:79:ALA:O	1:BI:82:LEU:HG	2.17	0.45
2:BJ:55:ALA:O	2:BJ:59:SER:OG	2.29	0.45
4:BS:153:ILE:O	4:BS:156:PRO:HD2	2.16	0.45
1:BV:6:LYS:HG2	1:BV:100:ASP:OD2	2.17	0.45
2:BW:123:ILE:O	2:BW:126:THR:HG22	2.17	0.45
1:BX:107:ILE:HG22	10:BX:200:CYC:HBB1	1.99	0.45
1:BZ:76:GLU:HG2	1:BZ:77:MET:N	2.31	0.45
2:CD:60:LEU:HD21	10:CD:200:CYC:HMC3	1.97	0.45
6:CM:597:TYR:CD2	6:CM:600:LYS:HG3	2.52	0.45
1:CR:47:ARG:O	1:CR:51:VAL:HG12	2.17	0.45
1:CR:65:ILE:O	1:CR:72:ALA:HB3	2.17	0.45
1:DA:12:ASP:OD1	2:DB:91:TYR:OH	2.22	0.45
1:DC:101:VAL:HB	1:DC:155:PHE:CD2	2.52	0.45
8:DG:242:ILE:HG23	8:DG:242:ILE:O	2.17	0.45

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
9:DI:89:ARG:CZ	9:DI:161:MET:HA	2.46	0.45
9:DI:209:ASP:O	9:DI:213:LEU:HD23	2.17	0.45
2:DR:43:VAL:HG13	2:DR:141:VAL:HG12	1.99	0.45
1:DU:104:ILE:HG21	1:DU:156:VAL:CG2	2.47	0.45
9:EA:60:SER:OG	9:EA:105:ILE:HD11	2.17	0.45
9:EA:107:LEU:HB3	11:EA:400:45D:H141	1.98	0.45
9:EA:187:LYS:HD3	9:EA:203:ASP:OD2	2.17	0.45
2:AD:87:TYR:CE2	5:BA:22:LEU:HD22	2.51	0.45
3:AE:5:SER:N	2:AF:3:ASP:OD2	2.50	0.45
3:AE:88:TYR:O	3:AE:92:VAL:HG23	2.17	0.45
2:AF:111:GLY:O	2:AF:115:THR:HG23	2.17	0.45
1:AH:30:VAL:HG13	2:AI:31:PHE:HA	1.99	0.45
1:AJ:83:ARG:NH1	10:AJ:200:CYC:O2A	2.47	0.45
4:AK:82:CYS:HA	10:AK:200:CYC:HHD	1.98	0.45
4:AK:135:LYS:HD2	4:AK:154:ALA:HA	1.98	0.45
2:AL:17:LYS:HG3	2:AL:18:TYR:O	2.15	0.45
2:AO:126:THR:O	2:AO:130:ILE:HG13	2.16	0.45
2:AS:88:TYR:CD2	2:AS:130:ILE:HD11	2.52	0.45
2:AW:77:ARG:HB3	10:AW:200:CYC:HMD1	1.97	0.45
2:AW:127:VAL:HG13	2:AW:157:CYS:SG	2.57	0.45
2:AY:119:LEU:CD1	10:AY:200:CYC:HAA2	2.47	0.45
5:BA:10:VAL:C	5:BA:25:THR:HB	2.41	0.45
6:BD:15:GLN:OE1	6:BD:262:ALA:HB2	2.16	0.45
6:BD:69:SER:CA	6:BD:82:LEU:HD21	2.35	0.45
6:BD:455:GLU:OE2	6:BD:661:THR:OG1	2.23	0.45
9:BH:109:PHE:CZ	9:BH:113:LEU:HD11	2.52	0.45
9:BH:195:ASN:OD1	9:BH:198:VAL:HB	2.17	0.45
1:BK:93:THR:O	1:BK:97:VAL:HG22	2.16	0.45
2:BQ:25:ASP:OD2	9:CQ:165:ALA:HB1	2.16	0.45
2:BT:4:ALA:O	2:BT:8:VAL:HG23	2.17	0.45
2:BW:122:PRO:HB2	2:BW:125:SER:HB2	1.99	0.45
2:BY:106:GLU:OE2	6:CM:674:SER:HB2	2.16	0.45
1:CC:4:VAL:HA	1:CC:26:ILE:HD11	1.98	0.45
2:CF:10:ASN:O	2:CF:14:VAL:HG22	2.17	0.45
2:CF:44:ILE:HD13	2:CF:149:MET:HE2	1.97	0.45
2:CF:66:THR:HG21	10:CG:200:CYC:O1A	2.16	0.45
1:CG:82:LEU:HB2	8:CP:242:ILE:HD12	1.99	0.45
1:CG:155:PHE:O	1:CG:159:LYS:HG2	2.16	0.45
5:CJ:40:GLN:HG3	5:CJ:50:ILE:HD11	1.99	0.45
6:CM:144:TYR:CE2	6:CM:148:ASN:HB3	2.51	0.45
6:CM:274:SER:OG	6:CM:277:GLU:OE1	2.33	0.45

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
6:CM:291:GLU:HG2	6:CM:348:ARG:HH12	1.82	0.45
6:CM:477:ASN:OD1	6:CM:480:THR:OG1	2.34	0.45
1:CV:28:ALA:O	1:CV:31:THR:OG1	2.29	0.45
8:DG:238:GLN:OE1	1:DL:52:LYS:HG2	2.17	0.45
9:DI:20:VAL:HB	9:DI:145:LEU:HD21	1.98	0.45
9:DI:151:ILE:O	9:DI:155:ARG:HG3	2.16	0.45
1:DL:4:VAL:HG23	2:DM:97:LEU:CD2	2.47	0.45
2:DO:72:MET:HE2	2:DO:78:TYR:CD2	2.51	0.45
1:DS:73:TYR:HD2	8:DY:200:VAL:HG23	1.82	0.45
2:DT:81:CYS:HB3	10:DT:200:CYC:HBC1	1.77	0.45
1:DU:17:TYR:OH	2:DV:89:LEU:HG	2.17	0.45
1:AA:102:THR:O	1:AA:106:GLU:HG2	2.16	0.45
3:AE:16:ARG:O	2:AF:94:TYR:OH	2.28	0.45
3:AE:50:ILE:HD11	3:AE:140:LEU:HB2	1.99	0.45
2:AF:22:ALA:O	2:AF:26:LYS:NZ	2.25	0.45
1:AJ:102:THR:O	1:AJ:106:GLU:HG2	2.17	0.45
1:AJ:142:SER:OG	1:AJ:144:ASP:OD1	2.27	0.45
2:AL:56:VAL:O	2:AL:61:LEU:HG	2.17	0.45
2:AS:129:ALA:O	2:AS:133:ILE:HG13	2.17	0.45
1:AV:129:SER:O	1:AV:133:MET:HG3	2.17	0.45
1:AZ:24:ASP:HA	1:AZ:27:LYS:HG2	1.99	0.45
6:BD:860:THR:C	6:BD:861:LEU:HD12	2.41	0.45
9:BH:250:LEU:O	9:BH:251:ILE:HD13	2.17	0.45
1:BI:37:LEU:CD1	2:BJ:28:LYS:HD3	2.46	0.45
2:BJ:112:LEU:CD2	2:BJ:160:LEU:HD21	2.46	0.45
1:BK:105:GLU:HA	1:BK:109:LEU:HB3	1.99	0.45
3:BM:37:ILE:O	3:BM:40:ALA:HB3	2.17	0.45
3:BM:117:ASN:HB3	6:CM:234:PRO:HG2	1.98	0.45
3:BM:126:MET:HE3	10:BM:200:CYC:H3C	1.99	0.45
4:BS:60:PHE:O	4:BS:64:PRO:HA	2.17	0.45
2:BT:81:CYS:HA	10:BT:200:CYC:HHD	1.98	0.45
2:BW:134:LYS:HA	2:BW:153:LEU:HD13	1.99	0.45
1:BZ:87:TYR:CG	10:BZ:200:CYC:HBB3	2.52	0.45
1:BZ:87:TYR:CD2	10:BZ:200:CYC:HBB3	2.51	0.45
1:CC:5:THR:O	1:CC:9:VAL:HG23	2.17	0.45
1:CC:134:LYS:HB2	1:CC:153:PHE:CG	2.52	0.45
2:CD:93:THR:HA	2:CD:149:MET:HE1	1.99	0.45
2:CF:153:LEU:O	2:CF:156:ILE:HG22	2.16	0.45
1:CG:61:LYS:HG2	9:CQ:57:GLY:N	2.32	0.45
1:CG:109:LEU:HD21	1:CG:159:LYS:CB	2.47	0.45
6:CM:63:ASN:HB2	6:CM:66:LEU:HB3	1.99	0.45

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
6:CM:176:VAL:HG22	6:CM:230:GLU:CD	2.42	0.45
6:CM:338:ILE:CG2	6:CM:341:ARG:H	2.30	0.45
6:CM:804:GLY:C	6:CM:838:MET:HE1	2.41	0.45
6:CM:866:PHE:O	6:CM:870:GLU:HG2	2.17	0.45
9:CQ:216:LEU:O	9:CQ:298:ILE:HG22	2.17	0.45
1:CR:104:ILE:HG22	1:CR:109:LEU:HD23	1.98	0.45
2:CS:56:VAL:HG12	2:CS:61:LEU:HG	1.99	0.45
2:CU:144:ASP:OD1	2:CU:144:ASP:N	2.48	0.45
1:CV:44:THR:O	1:CV:47:ARG:HG2	2.17	0.45
2:CW:122:PRO:O	2:CW:126:THR:HG22	2.17	0.45
2:DD:57:ALA:HB1	8:DH:201:ARG:NH2	2.31	0.45
1:DJ:81:CYS:SG	1:DJ:85:MET:HE3	2.57	0.45
1:DJ:115:MET:HG3	2:DO:78:TYR:CD1	2.52	0.45
1:DN:47:ARG:O	1:DN:51:VAL:HG23	2.17	0.45
2:DO:50:THR:HG21	9:EA:60:SER:HB2	1.99	0.45
1:DQ:37:LEU:HG	2:DR:28:LYS:HE3	1.99	0.45
1:DS:133:MET:HA	1:DS:136:VAL:HG12	1.99	0.45
1:DU:17:TYR:CZ	2:DV:90:ARG:HA	2.52	0.45
1:DU:80:THR:HG23	1:DU:83:ARG:HH21	1.81	0.45
2:DV:41:ALA:HA	2:DV:44:ILE:HG22	1.99	0.45
3:AE:8:ILE:HG13	2:AF:1:MET:CE	2.47	0.44
2:AF:20:ASP:O	2:AF:24:MET:HG3	2.17	0.44
2:AF:104:LEU:HD11	2:AF:152:TYR:HB3	1.99	0.44
2:AL:36:LEU:HD12	7:BF:85:GLN:HB2	1.99	0.44
1:AN:8:ILE:CD1	1:AN:18:LEU:HD11	2.47	0.44
1:AP:61:LYS:HE2	1:AP:128:GLN:NE2	2.32	0.44
2:AS:65:VAL:HG12	2:AS:72:MET:HE3	1.98	0.44
2:AY:127:VAL:O	2:AY:131:GLN:HG3	2.16	0.44
6:BD:453:PRO:HG3	6:BD:462:PHE:CE1	2.52	0.44
6:BD:551:PHE:CD2	6:BD:555:LEU:HD21	2.52	0.44
6:BD:793:ALA:HB3	6:BD:794:PRO:HD3	1.98	0.44
1:BP:24:ASP:OD1	1:BP:25:ARG:N	2.50	0.44
2:BW:81:CYS:HA	10:BW:200:CYC:HHD	1.99	0.44
2:BW:96:MET:HG3	2:BW:148:GLU:OE2	2.17	0.44
1:BX:122:PRO:HG2	1:BX:125:ALA:CB	2.45	0.44
2:BY:60:LEU:CB	2:BY:72:MET:HE1	2.43	0.44
1:BZ:18:LEU:HD12	2:CA:97:LEU:CD1	2.41	0.44
1:CE:58:LEU:HD13	1:CE:129:SER:N	2.31	0.44
6:CM:156:MET:HB3	6:CM:202:MET:HE2	1.99	0.44
6:CM:167:ILE:CD1	6:CM:219:ILE:HG22	2.47	0.44
6:CM:285:ALA:O	6:CM:289:ILE:HG12	2.16	0.44

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
6:CM:358:ARG:HB2	6:CM:434:PHE:HB3	1.99	0.44
1:CT:104:ILE:HG22	1:CT:109:LEU:HD23	1.98	0.44
1:CT:143:SER:O	1:CT:146:ALA:HB3	2.17	0.44
2:CW:37:ARG:HD3	2:CW:96:MET:O	2.17	0.44
1:CY:90:ARG:NH2	2:CZ:16:GLY:HA2	2.32	0.44
2:CZ:72:MET:O	2:CZ:72:MET:HG3	2.18	0.44
1:DA:131:ARG:N	1:DA:157:ILE:HD11	2.32	0.44
1:DC:122:PRO:HB2	1:DC:125:ALA:CB	2.47	0.44
9:DI:109:PHE:CZ	9:DI:113:LEU:HD11	2.52	0.44
1:DJ:23:LEU:HD12	1:DJ:24:ASP:N	2.32	0.44
1:DJ:58:LEU:HD22	1:DJ:129:SER:CB	2.32	0.44
2:DM:15:GLN:HG3	2:DM:17:LYS:HZ2	1.82	0.44
1:DQ:40:ALA:HB2	1:DQ:97:VAL:HG22	1.99	0.44
1:DS:108:GLY:C	1:DS:109:LEU:HD22	2.42	0.44
9:EA:67:ALA:HA	9:EA:70:GLU:OE1	2.16	0.44
9:EA:90:ALA:O	9:EA:163:PHE:HB2	2.17	0.44
1:AA:17:TYR:CE2	2:AB:90:ARG:HA	2.52	0.44
1:AA:50:ILE:HD11	1:AA:137:ALA:N	2.32	0.44
1:AA:91:LEU:O	1:AA:104:ILE:HG12	2.17	0.44
2:AB:113:LYS:HE2	2:AB:117:ASN:OD1	2.17	0.44
1:AC:37:LEU:HD11	2:AD:24:MET:SD	2.58	0.44
1:AC:101:VAL:HA	1:AC:104:ILE:HD12	2.00	0.44
3:AE:42:THR:CG2	3:AE:141:LEU:HD21	2.47	0.44
2:AI:76:ARG:HB2	1:AJ:110:VAL:CG1	2.47	0.44
2:AL:58:LYS:HB3	2:AL:58:LYS:HE2	1.65	0.44
1:AN:100:ASP:OD2	1:AN:102:THR:OG1	2.28	0.44
1:AN:122:PRO:HB2	1:AN:125:ALA:HB3	1.99	0.44
2:AO:85:LEU:HB3	2:AO:133:ILE:HD11	1.99	0.44
2:AO:112:LEU:HD11	2:AO:116:TYR:CZ	2.52	0.44
1:AP:75:GLU:OE1	1:AP:75:GLU:N	2.44	0.44
2:AU:72:MET:HG2	2:AU:78:TYR:CD2	2.53	0.44
2:AW:40:ALA:HB2	2:AW:145:ALA:HB1	1.99	0.44
1:AX:104:ILE:HG21	1:AX:156:VAL:HG22	1.99	0.44
2:AY:96:MET:CB	2:AY:149:MET:HE2	2.46	0.44
6:BD:18:GLN:O	6:BD:171:ASP:HB3	2.17	0.44
6:BD:75:ILE:HD12	6:BD:197:VAL:CG2	2.39	0.44
6:BD:691:PHE:HE2	1:DC:52:LYS:HE3	1.82	0.44
6:BD:748:GLU:C	6:BD:749:ARG:HD3	2.43	0.44
9:BH:258:GLU:HB2	9:BH:259:PRO:HD2	1.98	0.44
9:BH:292:LEU:HD23	9:BH:298:ILE:CD1	2.47	0.44
1:BI:101:VAL:HG13	1:BP:20:PRO:HG2	1.98	0.44

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:BK:25:ARG:HD3	6:CM:38:SER:O	2.17	0.44
2:BL:121:VAL:HG13	10:BL:200:CYC:NC	2.32	0.44
2:BQ:81:CYS:HA	10:BQ:200:CYC:HHD	1.98	0.44
1:BV:123:ILE:CD1	1:CE:117:ARG:HH22	2.30	0.44
1:BV:159:LYS:HB3	1:BV:159:LYS:HE3	1.78	0.44
1:BX:105:GLU:O	1:BX:110:VAL:HG23	2.17	0.44
2:CD:113:LYS:HD3	2:CD:160:LEU:O	2.17	0.44
2:CD:148:GLU:HG3	2:CD:152:TYR:CE2	2.52	0.44
2:CF:25:ASP:HA	2:CF:28:LYS:NZ	2.31	0.44
5:CK:6:ILE:O	5:CK:6:ILE:HG13	2.17	0.44
6:CM:141:ILE:HG13	6:CM:149:MET:HE2	1.99	0.44
7:CO:114:ALA:O	7:CO:118:LYS:HG2	2.17	0.44
9:CQ:205:LEU:HA	9:CQ:210:PHE:CE1	2.49	0.44
1:CT:27:LYS:O	1:CT:31:THR:HG23	2.17	0.44
1:CV:71:ASN:HB3	10:CV:200:CYC:OC	2.17	0.44
2:CW:63:SER:O	2:CW:67:ARG:HG3	2.16	0.44
1:DA:77:MET:HE3	8:DG:201:ARG:CB	2.45	0.44
2:DB:73:TYR:O	2:DB:77:ARG:HB2	2.17	0.44
2:DB:101:ALA:HB2	2:DB:152:TYR:HD1	1.81	0.44
1:DN:63:PRO:O	1:DN:67:SER:OG	2.29	0.44
1:DS:71:ASN:HB3	10:DS:200:CYC:OC	2.17	0.44
1:DS:109:LEU:O	1:DS:112:VAL:HG12	2.17	0.44
1:DU:116:TYR:CE2	1:DU:126:VAL:HG11	2.52	0.44
5:DX:40:GLN:CG	5:DX:44:MET:HE3	2.43	0.44
9:EA:253:GLU:HG3	9:EA:271:GLY:HA2	2.00	0.44
1:AC:3:ILE:HG22	1:AC:26:ILE:CD1	2.47	0.44
1:AC:101:VAL:HB	1:AC:155:PHE:CE2	2.51	0.44
2:AD:71:ASN:O	2:AD:77:ARG:HB3	2.17	0.44
4:AK:166:GLU:HG3	2:AS:69:GLY:O	2.17	0.44
1:AN:86:ASP:OD1	2:AO:18:TYR:OH	2.17	0.44
1:AN:94:TYR:OH	2:AO:17:LYS:O	2.32	0.44
2:AS:57:ALA:HA	2:AS:61:LEU:CG	2.45	0.44
1:AV:2:SER:OG	1:AV:98:SER:HB2	2.16	0.44
1:AV:85:MET:HG3	10:AV:200:CYC:HBC1	1.98	0.44
2:AW:85:LEU:HD21	10:AW:200:CYC:HBC1	2.00	0.44
2:AY:52:VAL:HG21	2:AY:86:ASP:OD1	2.18	0.44
2:AY:93:THR:HA	2:AY:149:MET:CE	2.43	0.44
1:AZ:48:GLU:O	1:AZ:52:LYS:HD3	2.17	0.44
6:BD:238:ASN:HB2	6:BD:252:GLU:OE2	2.17	0.44
6:BD:373:LEU:HB3	6:BD:374:PRO:HD3	2.00	0.44
9:BH:47:MET:HA	9:BH:51:LEU:HB2	1.99	0.44

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
9:BH:257:THR:CG2	9:BH:267:ILE:HG12	2.45	0.44
1:BI:88:TYR:O	1:BI:92:VAL:HG23	2.16	0.44
3:BM:58:LEU:HD22	3:BM:85:TYR:OH	2.18	0.44
2:BN:61:LEU:HD21	2:BN:78:TYR:OH	2.17	0.44
1:CG:115:MET:HE3	1:CG:116:TYR:CE1	2.52	0.44
6:CM:19:THR:OG1	6:CM:22:VAL:HG23	2.17	0.44
6:CM:183:LYS:HG2	6:CM:192:ILE:CD1	2.44	0.44
9:CQ:112:ARG:NH2	9:CQ:116:LEU:HD21	2.31	0.44
1:CR:127:ALA:HA	1:CR:130:VAL:HG12	1.98	0.44
2:CS:3:ASP:HB2	2:CS:98:ALA:O	2.17	0.44
1:CT:7:SER:C	1:CT:18:LEU:HD21	2.43	0.44
1:CT:66:VAL:HA	1:CT:72:ALA:HB3	1.99	0.44
2:CU:10:ASN:O	2:CU:14:VAL:HG13	2.17	0.44
1:CV:64:ASP:OD1	1:CV:64:ASP:N	2.47	0.44
2:CW:78:TYR:O	2:CW:82:ILE:HG12	2.17	0.44
2:DB:87:TYR:HB3	10:DB:200:CYC:HBB2	1.98	0.44
1:DC:81:CYS:HB3	10:DC:200:CYC:HBC1	1.33	0.44
9:DI:17:ALA:O	9:DI:20:VAL:HG22	2.18	0.44
2:DK:3:ASP:HB2	2:DK:98:ALA:O	2.16	0.44
1:DL:17:TYR:HE1	2:DM:90:ARG:HD3	1.81	0.44
1:DL:152:TYR:O	1:DL:156:VAL:HG23	2.17	0.44
2:DM:76:ARG:HG3	1:DN:110:VAL:HG11	2.00	0.44
2:DM:85:LEU:HD13	2:DM:133:ILE:HD11	1.98	0.44
1:DN:14:GLU:HB2	1:DN:16:ARG:HG2	1.97	0.44
1:DN:115:MET:CE	10:DN:200:CYC:HMA3	2.48	0.44
1:DU:92:VAL:HG21	1:DU:153:PHE:CE1	2.52	0.44
1:AC:33:GLY:HA2	1:AC:36:ARG:CG	2.47	0.44
2:AD:53:LYS:HG3	3:AE:118:SER:O	2.17	0.44
2:AF:41:ALA:N	2:AF:96:MET:HE1	2.33	0.44
2:AF:133:ILE:O	2:AF:137:THR:HG22	2.18	0.44
1:AH:47:ARG:HG3	1:AH:48:GLU:N	2.33	0.44
2:AI:4:ALA:O	2:AI:8:VAL:HG22	2.18	0.44
2:AL:34:GLY:O	2:AL:38:VAL:HG23	2.17	0.44
1:AN:81:CYS:HA	10:AN:200:CYC:CHD	2.44	0.44
2:AO:81:CYS:HA	10:AO:200:CYC:HHD	1.99	0.44
2:AS:71:ASN:O	2:AS:77:ARG:HB3	2.18	0.44
2:AU:119:LEU:HD11	10:AU:200:CYC:HAA2	1.98	0.44
1:AV:35:ALA:O	1:AV:39:ILE:HG13	2.18	0.44
6:BD:52:LEU:HD11	6:BD:216:ALA:CB	2.44	0.44
6:BD:155:ASP:CG	10:BD:902:CYC:HHB	2.42	0.44
6:BD:597:TYR:HD2	6:BD:600:LYS:HG3	1.82	0.44

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
6:BD:751:LEU:O	6:BD:751:LEU:HD12	2.18	0.44
1:BK:102:THR:HB	1:BK:103:PRO:HD3	1.98	0.44
2:BL:54:GLU:O	2:BL:58:LYS:HE3	2.17	0.44
1:BR:18:LEU:HD22	4:BS:98:ILE:HD13	2.00	0.44
4:BS:4:ALA:HA	4:BS:7:THR:HG22	2.00	0.44
1:BV:80:THR:HG23	1:BV:83:ARG:HE	1.82	0.44
2:BY:77:ARG:HG2	10:BY:200:CYC:HAD1	1.98	0.44
1:BZ:68:PRO:HA	1:BZ:73:TYR:CD2	2.53	0.44
1:BZ:128:GLN:O	1:BZ:132:GLU:HG2	2.18	0.44
2:CA:107:ARG:NH1	6:CM:596:HIS:O	2.48	0.44
2:CA:124:SER:HA	6:CM:11:LEU:HD11	1.99	0.44
10:CH:200:CYC:HMA1	5:CK:25:THR:HG21	1.99	0.44
6:CM:30:GLN:OE1	6:CM:265:GLN:HG3	2.17	0.44
6:CM:182:LEU:HD12	6:CM:185:VAL:CG2	2.47	0.44
6:CM:268:ALA:HB3	6:CM:270:LYS:HZ2	1.82	0.44
6:CM:451:ASN:OD1	6:CM:600:LYS:HA	2.16	0.44
6:CM:453:PRO:HG3	6:CM:462:PHE:CZ	2.53	0.44
6:CM:458:PHE:HB3	6:CM:552:GLY:C	2.42	0.44
1:CR:71:ASN:ND2	1:CR:121:THR:OG1	2.50	0.44
2:CS:121:VAL:HG13	2:CS:122:PRO:HD2	1.98	0.44
1:CT:33:GLY:HA2	1:CT:36:ARG:HD2	2.00	0.44
1:CT:37:LEU:O	1:CT:40:ALA:HB3	2.17	0.44
1:CT:72:ALA:C	1:CT:77:MET:HB2	2.42	0.44
1:CV:144:ASP:OD1	1:CV:145:ASP:N	2.50	0.44
2:CZ:22:ALA:O	2:CZ:26:LYS:HG3	2.18	0.44
2:DB:40:ALA:HB2	2:DB:145:ALA:HB1	1.99	0.44
2:DD:81:CYS:HA	10:DD:200:CYC:CHD	2.47	0.44
2:DK:154:ASP:HA	2:DK:157:CYS:SG	2.58	0.44
1:DL:77:MET:HE2	1:DL:77:MET:N	2.32	0.44
2:DR:1:MET:HE3	2:DR:103:ILE:HB	1.99	0.44
2:DR:112:LEU:CD1	10:DR:200:CYC:HMB3	2.48	0.44
2:DT:133:ILE:O	2:DT:137:THR:HG22	2.17	0.44
1:DU:85:MET:SD	1:DU:129:SER:OG	2.70	0.44
2:DV:144:ASP:N	2:DV:144:ASP:OD1	2.50	0.44
9:EA:185:ARG:CZ	9:EA:207:ALA:HB1	2.47	0.44
1:AA:25:ARG:HB3	1:AH:25:ARG:HD2	2.00	0.44
2:AB:2:GLN:O	2:AB:100:ASP:HB3	2.18	0.44
2:AB:57:ALA:HA	2:AB:61:LEU:CD1	2.44	0.44
1:AC:121:THR:HG23	10:AC:200:CYC:C1C	2.48	0.44
1:AH:24:ASP:OD1	1:AH:25:ARG:N	2.50	0.44
1:AN:11:ALA:CB	1:AN:18:LEU:HD23	2.47	0.44

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:AN:57:ARG:HA	1:AN:60:GLN:HG3	2.00	0.44
2:AQ:47:ASN:HB2	2:AQ:140:LEU:HD21	1.99	0.44
2:AS:52:VAL:HG21	2:AS:86:ASP:OD1	2.17	0.44
2:AU:72:MET:HG2	2:AU:78:TYR:HD2	1.83	0.44
1:AV:144:ASP:OD1	1:AV:145:ASP:N	2.51	0.44
2:AY:1:MET:HE3	2:AY:103:ILE:HD12	2.00	0.44
1:AZ:140:LEU:HD23	1:AZ:140:LEU:O	2.18	0.44
5:BB:3:MET:SD	5:BB:32:PRO:HA	2.57	0.44
6:BD:141:ILE:CG2	6:BD:149:MET:HE2	2.41	0.44
6:BD:761:THR:HG22	2:CS:87:TYR:OH	2.18	0.44
7:BF:114:ALA:O	7:BF:117:VAL:HG12	2.18	0.44
9:BH:201:TYR:HA	9:BH:213:LEU:CD1	2.48	0.44
1:BI:76:GLU:O	1:BI:80:THR:HG23	2.17	0.44
1:BI:102:THR:O	1:BI:106:GLU:HG2	2.18	0.44
1:BI:130:VAL:HG21	1:BI:160:MET:SD	2.56	0.44
2:BN:85:LEU:CG	10:BN:200:CYC:HBC1	2.47	0.44
4:BS:72:ASN:HD21	4:BS:122:VAL:HG22	1.82	0.44
2:BT:144:ASP:N	2:BT:144:ASP:OD1	2.49	0.44
1:BZ:11:ALA:HA	1:BZ:16:ARG:NH1	2.33	0.44
1:CE:72:ALA:HB2	10:CE:200:CYC:CMD	2.48	0.44
5:CJ:31:VAL:HG11	5:CJ:36:TRP:CE3	2.53	0.44
6:CM:701:ARG:HD2	6:CM:705:GLU:CD	2.42	0.44
9:CQ:64:ALA:HA	9:CQ:101:TRP:CZ2	2.52	0.44
9:CQ:98:TYR:OH	9:CQ:158:VAL:HG22	2.17	0.44
9:CQ:273:VAL:HG12	9:CQ:284:MET:O	2.18	0.44
2:CZ:102:SER:HA	2:CZ:105:ASP:OD1	2.18	0.44
2:DB:51:ILE:HG21	2:DB:89:LEU:HD11	1.99	0.44
1:DC:122:PRO:HB2	1:DC:125:ALA:HB3	1.98	0.44
8:DH:230:ASN:ND2	1:DJ:67:SER:HB3	2.33	0.44
9:DI:101:TRP:HB3	9:DI:105:ILE:HB	1.99	0.44
2:DK:1:MET:SD	2:DK:103:ILE:HB	2.58	0.44
1:DL:47:ARG:O	1:DL:51:VAL:HG12	2.17	0.44
2:DM:73:TYR:HH	1:DN:90:ARG:HH22	1.60	0.44
1:DQ:57:ARG:HH12	1:DQ:136:VAL:HG22	1.81	0.44
1:DQ:127:ALA:N	1:DQ:160:MET:HE1	2.33	0.44
1:DS:62:ARG:HD3	9:EA:58:ALA:N	2.32	0.44
2:DV:51:ILE:HA	2:DV:54:GLU:CD	2.42	0.44
1:AA:83:ARG:NH2	10:AA:200:CYC:O1A	2.48	0.44
2:AB:25:ASP:OD1	2:AB:26:LYS:N	2.51	0.44
2:AB:85:LEU:HD22	2:AB:133:ILE:HD11	1.99	0.44
1:AC:98:SER:CB	1:AC:103:PRO:HG2	2.48	0.44

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:AD:2:GLN:HB2	2:AD:6:THR:CG2	2.46	0.44
2:AF:85:LEU:CG	10:AF:200:CYC:HBC1	2.43	0.44
2:AF:122:PRO:O	2:AF:126:THR:HG23	2.17	0.44
4:AK:91:ARG:O	4:AK:94:SER:OG	2.31	0.44
10:AK:200:CYC:HBA2	6:BD:401:LEU:CD1	2.48	0.44
1:AN:46:SER:OG	1:AN:140:LEU:HD23	2.18	0.44
1:AP:89:LEU:HD13	1:AP:133:MET:HE2	2.00	0.44
1:AP:110:VAL:CG1	6:BD:489:PRO:HB3	2.48	0.44
2:AQ:111:GLY:HA2	2:AQ:114:GLU:OE1	2.17	0.44
2:AU:79:ALA:HA	2:AU:82:ILE:HG22	1.99	0.44
1:AV:64:ASP:HA	1:AV:67:SER:HB2	2.00	0.44
1:AV:109:LEU:O	1:AV:112:VAL:HG12	2.17	0.44
2:AW:41:ALA:CB	2:AW:97:LEU:HD21	2.48	0.44
5:BB:26:TYR:CE1	5:BB:51:VAL:HG21	2.52	0.44
6:BD:18:GLN:CB	6:BD:173:ASN:HD22	2.26	0.44
6:BD:77:THR:HG21	6:BD:138:PRO:HB2	2.00	0.44
6:BD:186:ILE:HG13	6:BD:190:CYS:CB	2.44	0.44
6:BD:348:ARG:HD2	6:BD:417:LEU:HG	1.99	0.44
6:BD:551:PHE:HE2	6:BD:555:LEU:HD11	1.83	0.44
6:BD:681:GLN:HG2	6:BD:682:ARG:N	2.32	0.44
8:BG:233:PRO:O	8:BG:235:VAL:HG13	2.18	0.44
9:BH:193:VAL:HG21	9:BH:269:VAL:HG22	1.99	0.44
9:BH:249:LYS:H	9:BH:275:THR:CB	2.30	0.44
2:BJ:96:MET:HA	2:BJ:152:TYR:CE2	2.53	0.44
1:BK:65:ILE:HG21	10:BK:200:CYC:H2C	1.98	0.44
1:BP:134:LYS:HB2	1:BP:153:PHE:HB3	1.99	0.44
1:BR:19:SER:HB2	1:BR:20:PRO:HD2	1.99	0.44
1:BR:35:ALA:O	1:BR:39:ILE:HG13	2.17	0.44
1:BV:81:CYS:HA	10:BV:200:CYC:CHD	2.48	0.44
2:CA:68:PRO:HA	2:CA:73:TYR:CD1	2.53	0.44
1:CC:4:VAL:HA	1:CC:26:ILE:CD1	2.47	0.44
1:CE:4:VAL:CG1	1:CE:26:ILE:HG23	2.46	0.44
1:CE:83:ARG:HH12	10:CE:200:CYC:CGA	2.27	0.44
1:CG:91:LEU:CD1	1:CG:108:GLY:HA3	2.47	0.44
6:CM:5:ALA:HB2	6:CM:436:GLN:CG	2.39	0.44
6:CM:56:ILE:HD11	6:CM:216:ALA:O	2.18	0.44
6:CM:697:VAL:CG2	6:CM:701:ARG:HE	2.28	0.44
7:CO:113:MET:HE3	7:CO:113:MET:HB2	1.83	0.44
9:CQ:47:MET:HA	9:CQ:51:LEU:HB2	2.00	0.44
9:CQ:104:ASN:OD1	9:CQ:155:ARG:HD3	2.17	0.44
2:CS:85:LEU:HD23	2:CS:85:LEU:HA	1.80	0.44

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:CV:74:GLY:O	1:CV:78:THR:HG23	2.18	0.44
1:CY:27:LYS:O	1:CY:31:THR:HG23	2.18	0.44
1:DC:109:LEU:HD23	1:DC:109:LEU:HA	1.78	0.44
9:DI:224:GLN:N	9:DI:301:VAL:O	2.50	0.44
1:DQ:5:THR:HA	1:DQ:8:ILE:CG2	2.46	0.44
2:DT:144:ASP:OD1	2:DT:145:ALA:N	2.47	0.44
8:DZ:221:MET:O	8:DZ:225:ILE:HG12	2.18	0.44
1:AA:25:ARG:NE	1:AH:29:PHE:HB2	2.32	0.44
1:AC:16:ARG:NH1	1:AC:17:TYR:O	2.51	0.44
1:AC:94:TYR:O	1:AC:97:VAL:HG22	2.18	0.44
1:AC:116:TYR:CD2	1:AC:126:VAL:HG21	2.53	0.44
2:AD:92:ALA:CB	2:AD:149:MET:HE3	2.48	0.44
2:AL:92:ALA:HB3	2:AL:149:MET:HE1	1.98	0.44
2:AO:75:THR:HG23	10:AP:200:CYC:CAB	2.48	0.44
1:AR:77:MET:SD	10:AR:200:CYC:HAD1	2.57	0.44
2:AS:85:LEU:HG	10:BD:901:CYC:HBC1	1.99	0.44
2:AU:51:ILE:HD12	2:AU:51:ILE:H	1.82	0.44
1:AV:50:ILE:HD12	1:AV:136:VAL:HG12	1.99	0.44
2:AW:153:LEU:O	2:AW:156:ILE:HG22	2.17	0.44
6:BD:56:ILE:HD11	6:BD:216:ALA:O	2.18	0.44
6:BD:788:MET:HB3	6:BD:831:LEU:CD2	2.47	0.44
9:BH:249:LYS:O	9:BH:273:VAL:HG23	2.17	0.44
1:BI:25:ARG:HH21	1:BP:29:PHE:HD1	1.65	0.44
1:BI:58:LEU:CD2	10:BI:200:CYC:HBC3	2.48	0.44
2:BL:82:ILE:HA	2:BL:85:LEU:HD12	1.99	0.44
3:BM:104:ILE:HG22	3:BM:109:LEU:CD1	2.47	0.44
2:BN:119:LEU:HG	5:CJ:4:PHE:HZ	1.83	0.44
1:BV:154:ASP:HB3	2:CD:46:ALA:CB	2.47	0.44
2:BW:41:ALA:CA	2:BW:96:MET:HE1	2.47	0.44
2:CH:44:ILE:CD1	2:CH:149:MET:HE2	2.47	0.44
6:CM:177:VAL:HG13	6:CM:256:SER:O	2.17	0.44
1:CR:85:MET:O	1:CR:133:MET:HE1	2.18	0.44
1:CR:105:GLU:O	1:CR:110:VAL:HG23	2.18	0.44
2:CU:54:GLU:O	2:CU:58:LYS:HG2	2.18	0.44
2:CW:123:ILE:O	2:CW:127:VAL:HG23	2.18	0.44
1:CY:26:ILE:O	1:CY:30:VAL:HG13	2.18	0.44
2:CZ:15:GLN:OE1	2:CZ:17:LYS:NZ	2.43	0.44
9:DI:32:ASN:HB3	9:DI:171:ARG:NH2	2.28	0.44
9:DI:60:SER:OG	9:DI:105:ILE:HD11	2.18	0.44
2:DK:17:LYS:HB3	2:DK:17:LYS:HE3	1.79	0.44
2:DK:44:ILE:HG21	2:DK:93:THR:OG1	2.17	0.44

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:DL:26:ILE:O	1:DL:30:VAL:HG13	2.18	0.44
1:DL:91:LEU:HD21	1:DL:107:ILE:CG2	2.48	0.44
2:DM:144:ASP:N	2:DM:144:ASP:OD1	2.50	0.44
1:DQ:93:THR:O	1:DQ:96:VAL:HG22	2.17	0.44
2:DR:1:MET:CE	2:DR:103:ILE:HB	2.47	0.44
1:DS:142:SER:O	1:DS:146:ALA:N	2.40	0.44
9:EA:39:LEU:CD1	9:EA:139:LEU:HB2	2.48	0.44
1:AA:37:LEU:CD1	2:AB:28:LYS:HD3	2.47	0.44
2:AB:85:LEU:CD2	10:AB:200:CYC:HBC1	2.47	0.44
2:AB:104:LEU:O	2:AB:108:VAL:HB	2.17	0.44
1:AC:33:GLY:HA2	1:AC:36:ARG:HE	1.83	0.44
2:AD:39:ARG:HA	2:AD:39:ARG:HH11	1.82	0.44
2:AD:78:TYR:HB3	3:AE:115:MET:SD	2.57	0.44
10:AD:200:CYC:HBA1	5:BA:12:SER:N	2.33	0.44
2:AF:112:LEU:HD11	2:AF:116:TYR:CZ	2.52	0.44
2:AI:75:THR:OG1	1:AJ:110:VAL:O	2.27	0.44
1:AJ:152:TYR:O	1:AJ:156:VAL:HG23	2.17	0.44
2:AL:112:LEU:HD11	2:AL:116:TYR:CE2	2.53	0.44
2:AO:111:GLY:HA2	2:AO:114:GLU:OE2	2.18	0.44
2:AU:44:ILE:HD11	2:AU:137:THR:HG23	2.00	0.44
2:AU:57:ALA:HA	2:AU:61:LEU:CB	2.42	0.44
1:AV:64:ASP:O	1:AV:67:SER:HB2	2.18	0.44
2:AY:83:ARG:NH1	5:BB:17:ARG:O	2.46	0.44
6:BD:273:LEU:HB3	6:BD:277:GLU:HG2	2.00	0.44
6:BD:761:THR:HG22	2:CS:83:ARG:HD2	1.99	0.44
7:BF:114:ALA:CA	7:BF:117:VAL:HG12	2.47	0.44
8:BG:219:LEU:O	8:BG:223:ARG:HG3	2.17	0.44
3:BM:68:PRO:HA	3:BM:73:TYR:HE1	1.82	0.44
3:BM:102:GLU:O	3:BM:106:THR:HG23	2.17	0.44
2:BN:47:ASN:HB2	2:BN:140:LEU:HD21	2.00	0.44
2:BQ:76:ARG:HB2	1:BR:110:VAL:CG1	2.48	0.44
2:BQ:95:ALA:HB2	2:BQ:104:LEU:HG	2.00	0.44
4:BS:167:LEU:HD12	2:BW:10:ASN:HD21	1.83	0.44
1:BV:81:CYS:HA	10:BV:200:CYC:HHD	1.98	0.44
1:BZ:52:LYS:HE2	8:CP:225:ILE:HB	2.00	0.44
1:BZ:58:LEU:HD22	1:BZ:129:SER:CB	2.39	0.44
1:BZ:65:ILE:HD11	1:BZ:122:PRO:HG3	2.00	0.44
5:CK:12:SER:HB3	5:CK:17:ARG:NH2	2.32	0.44
6:CM:159:PHE:CE1	10:CM:902:CYC:HBB2	2.53	0.44
6:CM:749:ARG:NH2	6:CM:877:THR:HG21	2.33	0.44
6:CM:809:LEU:O	6:CM:857:ARG:NH2	2.51	0.44

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
6:CM:866:PHE:HE2	2:DT:119:LEU:HD21	1.83	0.44
1:CT:27:LYS:HZ3	2:CU:35:GLU:HA	1.83	0.44
1:CY:92:VAL:HG11	1:CY:153:PHE:CE1	2.51	0.44
10:CY:200:CYC:O1D	2:DD:62:TYR:OH	2.32	0.44
9:DI:173:ALA:O	9:DI:175:PRO:HD3	2.17	0.44
9:DI:223:LEU:HG	9:DI:301:VAL:CG1	2.48	0.44
9:DI:288:TRP:CD1	9:DI:303:ILE:HG12	2.51	0.44
9:DI:291:LEU:HD23	9:DI:299:PHE:CD2	2.53	0.44
1:DJ:11:ALA:HA	1:DJ:16:ARG:NH1	2.33	0.44
2:DM:156:ILE:O	2:DM:160:LEU:HD23	2.18	0.44
2:DT:50:THR:HG22	2:DT:54:GLU:OE1	2.18	0.44
2:DV:122:PRO:HG2	10:DV:200:CYC:CMC	2.46	0.44
2:DV:137:THR:O	2:DV:141:VAL:HG22	2.17	0.44
9:EA:142:ILE:CD1	9:EA:153:VAL:HG11	2.46	0.44
3:AE:20:SER:CB	1:AJ:151:ALA:HB1	2.45	0.44
2:AF:114:GLU:CG	5:BA:53:VAL:HG12	2.47	0.44
1:AJ:130:VAL:HB	1:AJ:157:ILE:CD1	2.48	0.44
1:AN:27:LYS:HE3	2:AO:35:GLU:OE1	2.17	0.44
1:AN:83:ARG:HD3	7:BF:107:LEU:HD21	1.98	0.44
2:AO:144:ASP:OD1	2:AO:144:ASP:N	2.50	0.44
1:AP:89:LEU:HB2	1:AP:133:MET:CE	2.44	0.44
1:AP:107:ILE:HD13	2:AQ:13:ASP:OD2	2.17	0.44
2:AW:101:ALA:CB	2:AW:104:LEU:HD12	2.48	0.44
2:AW:134:LYS:HB2	2:AW:153:LEU:HD13	1.98	0.44
1:AZ:4:VAL:HA	1:AZ:26:ILE:CD1	2.48	0.44
5:BB:1:MET:HE2	5:BB:32:PRO:HB3	1.99	0.44
6:BD:626:ILE:HG22	6:BD:635:VAL:CG2	2.48	0.44
9:BH:214:ILE:O	9:BH:217:PHE:HB2	2.18	0.44
1:BI:65:ILE:CG2	1:BI:72:ALA:HB3	2.41	0.44
1:BI:85:MET:HB3	1:BI:133:MET:CE	2.43	0.44
1:BK:92:VAL:HG11	1:BK:153:PHE:CE1	2.53	0.44
3:BM:141:LEU:HD22	3:BM:145:ASP:OD2	2.18	0.44
1:BP:12:ASP:OD1	2:BQ:91:TYR:OH	2.27	0.44
1:BP:55:GLY:N	1:BP:85:MET:HE1	2.33	0.44
1:BP:122:PRO:HG2	1:BP:125:ALA:CB	2.47	0.44
2:BQ:107:ARG:C	6:CM:338:ILE:HD12	2.42	0.44
4:BS:37:ARG:HD3	4:BS:97:LEU:O	2.18	0.44
4:BS:110:LEU:HD13	4:BS:163:GLU:CB	2.48	0.44
2:BT:34:GLY:HA2	2:BT:37:ARG:NH2	2.33	0.44
1:BX:153:PHE:O	1:BX:157:ILE:HG13	2.18	0.44
2:CA:63:SER:HB3	7:CO:101:ARG:HH12	1.82	0.44

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:CA:114:GLU:HG3	6:CM:471:LYS:HB2	2.00	0.44
2:CD:51:ILE:HD12	2:CD:51:ILE:H	1.83	0.44
2:CF:125:SER:HA	2:CF:128:GLN:NE2	2.33	0.44
1:CG:134:LYS:HD3	1:CG:153:PHE:HB3	2.00	0.44
2:CH:1:MET:O	6:CM:534:PHE:HB2	2.18	0.44
5:CK:10:VAL:HG12	5:CK:48:GLY:CA	2.41	0.44
6:CM:248:ILE:CG1	6:CM:356:SER:HB2	2.47	0.44
6:CM:285:ALA:HB2	6:CM:316:MET:HE1	1.99	0.44
6:CM:792:TYR:HD1	6:CM:800:VAL:HG11	1.83	0.44
9:CQ:233:GLY:HA3	9:CQ:236:ASN:HD22	1.83	0.44
1:CR:141:MET:CE	1:CR:146:ALA:HA	2.46	0.44
1:CT:16:ARG:O	2:CU:90:ARG:HD2	2.18	0.44
1:CV:78:THR:O	1:CV:82:LEU:HG	2.18	0.44
2:CW:111:GLY:HA2	2:CW:114:GLU:OE2	2.17	0.44
10:CW:200:CYC:HAD2	5:DF:33:TYR:OH	2.17	0.44
2:CZ:60:LEU:HD13	2:CZ:72:MET:CE	2.47	0.44
1:DA:100:ASP:O	1:DA:103:PRO:HD2	2.17	0.44
1:DL:109:LEU:HD22	1:DL:159:LYS:HB2	1.99	0.44
1:DL:154:ASP:OD1	2:DV:46:ALA:HB2	2.18	0.44
2:DO:71:ASN:ND2	2:DO:121:VAL:HG22	2.25	0.44
1:DQ:115:MET:HA	1:DQ:115:MET:HE3	2.00	0.44
1:DQ:131:ARG:NH1	1:DQ:157:ILE:HD12	2.33	0.44
2:DR:67:ARG:O	2:DR:73:TYR:HB2	2.18	0.44
2:DR:78:TYR:CD1	1:DS:115:MET:HG3	2.51	0.44
2:DT:112:LEU:HD22	10:DT:200:CYC:HMB1	2.00	0.44
1:DU:66:VAL:HA	1:DU:72:ALA:O	2.18	0.44
2:DV:130:ILE:HB	2:DV:157:CYS:SG	2.58	0.44
8:DZ:200:VAL:HG23	8:DZ:200:VAL:O	2.18	0.44
9:EA:191:GLU:HB2	9:EA:254:ARG:HA	2.00	0.44
1:AA:94:TYR:OH	2:AB:17:LYS:O	2.36	0.43
1:AA:100:ASP:OD1	1:AA:101:VAL:N	2.49	0.43
1:AA:107:ILE:HD11	5:BA:64:ALA:HB2	1.99	0.43
1:AC:4:VAL:HG12	1:AC:26:ILE:HD12	2.00	0.43
1:AC:33:GLY:HA2	1:AC:36:ARG:NE	2.32	0.43
1:AC:115:MET:O	1:AC:119:LEU:HD13	2.18	0.43
2:AD:51:ILE:HD11	2:AD:140:LEU:CD2	2.40	0.43
10:AF:200:CYC:CMA	10:AF:200:CYC:HB	2.31	0.43
2:AI:77:ARG:HG2	10:AI:200:CYC:CMD	2.40	0.43
10:AI:200:CYC:OB	6:BD:338:ILE:HG22	2.18	0.43
2:AO:78:TYR:O	2:AO:82:ILE:HG12	2.18	0.43
1:AR:19:SER:HB2	1:AR:20:PRO:HD2	2.00	0.43

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:AR:94:TYR:OH	2:AS:17:LYS:O	2.35	0.43
2:AS:5:ILE:HG22	2:AS:9:ILE:CD1	2.48	0.43
10:AU:200:CYC:O2D	5:BB:33:TYR:OH	2.15	0.43
1:AX:12:ASP:OD2	2:AY:107:ARG:NE	2.51	0.43
6:BD:14:PRO:HG3	6:BD:401:LEU:HD21	1.99	0.43
6:BD:429:GLN:HG2	6:BD:429:GLN:O	2.18	0.43
6:BD:519:LEU:CG	6:BD:522:ALA:HB2	2.41	0.43
6:BD:702:THR:OG1	6:BD:704:PRO:HD2	2.17	0.43
9:BH:298:ILE:HG12	9:BH:301:VAL:HB	2.00	0.43
9:BH:311:GLU:O	9:BH:312:LEU:HB2	2.17	0.43
2:BL:65:VAL:O	2:BL:72:MET:HB3	2.16	0.43
3:BM:7:VAL:CG1	3:BM:22:GLU:HB3	2.48	0.43
3:BM:9:LEU:HD23	3:BM:9:LEU:HA	1.89	0.43
2:BQ:47:ASN:CG	2:BQ:140:LEU:HD21	2.43	0.43
2:BQ:85:LEU:CG	10:BQ:200:CYC:HBC1	2.47	0.43
1:BV:141:MET:HE3	1:BV:141:MET:HB2	1.94	0.43
2:BY:60:LEU:HD13	2:BY:72:MET:CE	2.48	0.43
2:BY:83:ARG:HG2	2:BY:83:ARG:HH11	1.83	0.43
1:BZ:25:ARG:HH21	1:CE:21:GLY:HA3	1.82	0.43
1:CC:110:VAL:O	2:CH:75:THR:HG23	2.18	0.43
1:CG:109:LEU:HD11	1:CG:159:LYS:HB2	2.00	0.43
6:CM:410:ASN:CB	6:CM:413:MET:HE3	2.48	0.43
1:CR:27:LYS:HD3	2:CS:38:VAL:HG11	2.00	0.43
1:CR:91:LEU:CD2	1:CR:104:ILE:HG23	2.47	0.43
2:CS:51:ILE:HG13	2:CS:140:LEU:HD12	1.99	0.43
2:CS:71:ASN:ND2	2:CS:121:VAL:HA	2.33	0.43
2:CS:133:ILE:O	2:CS:137:THR:HG23	2.18	0.43
1:CV:68:PRO:HA	1:CV:73:TYR:CG	2.53	0.43
2:CZ:137:THR:O	2:CZ:141:VAL:HG22	2.17	0.43
1:DA:51:VAL:HG12	8:DZ:221:MET:HE1	2.00	0.43
1:DA:63:PRO:HD2	9:DI:55:ALA:HB1	2.00	0.43
9:DI:217:PHE:HA	9:DI:297:LYS:O	2.18	0.43
1:DJ:113:ARG:CZ	1:DS:117:ARG:HD2	2.48	0.43
1:DJ:150:SER:O	1:DJ:154:ASP:HB2	2.18	0.43
2:DK:78:TYR:CE1	1:DL:115:MET:HG3	2.52	0.43
1:DL:93:THR:O	1:DL:97:VAL:HG13	2.19	0.43
1:DN:18:LEU:HD12	2:DO:97:LEU:HD13	1.99	0.43
10:DO:200:CYC:HMA1	10:DO:200:CYC:NB	2.33	0.43
1:DQ:43:LEU:CD2	1:DQ:93:THR:HG22	2.48	0.43
1:DQ:85:MET:HG3	1:DQ:133:MET:CE	2.47	0.43
1:DS:19:SER:O	1:DS:23:LEU:HD23	2.18	0.43

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:DT:44:ILE:HD13	2:DT:149:MET:HE2	1.98	0.43
2:DT:72:MET:HE2	2:DT:78:TYR:CD2	2.53	0.43
1:DU:54:ALA:CB	1:DU:85:MET:HE1	2.48	0.43
2:DV:81:CYS:HA	10:DV:200:CYC:CHD	2.45	0.43
8:DY:202:ARG:HH22	8:DY:204:ILE:HB	1.83	0.43
9:EA:193:VAL:HG23	9:EA:255:GLY:C	2.43	0.43
9:EA:223:LEU:CD2	9:EA:225:PRO:HD3	2.48	0.43
1:AA:87:TYR:CD2	10:AA:200:CYC:HBB3	2.52	0.43
2:AI:71:ASN:HD22	2:AI:122:PRO:HD2	1.82	0.43
1:AJ:49:THR:HG21	2:BW:54:GLU:CD	2.43	0.43
4:AK:72:ASN:ND2	4:AK:123:PRO:HD2	2.30	0.43
1:AP:85:MET:O	1:AP:133:MET:HE1	2.19	0.43
1:AP:131:ARG:HA	1:AP:157:ILE:HD11	1.99	0.43
2:AQ:40:ALA:HB2	2:AQ:145:ALA:HB1	2.00	0.43
1:AV:39:ILE:CD1	1:AV:145:ASP:HB3	2.48	0.43
1:AX:50:ILE:HD11	1:AX:137:ALA:HB2	1.99	0.43
1:AX:93:THR:O	1:AX:96:VAL:HG22	2.18	0.43
2:BJ:25:ASP:OD1	2:BJ:26:LYS:N	2.52	0.43
2:BJ:71:ASN:C	2:BJ:77:ARG:HE	2.27	0.43
2:BJ:95:ALA:HB2	2:BJ:104:LEU:HG	2.00	0.43
1:BK:153:PHE:O	1:BK:157:ILE:HG13	2.18	0.43
3:BM:17:TYR:CE1	2:BN:90:ARG:HG3	2.52	0.43
3:BM:128:ASP:O	3:BM:131:THR:OG1	2.25	0.43
1:CE:28:ALA:O	1:CE:31:THR:OG1	2.30	0.43
6:CM:31:GLN:HB3	6:CM:33:ARG:HE	1.82	0.43
6:CM:57:ALA:O	6:CM:61:THR:HG23	2.18	0.43
6:CM:86:GLU:H	6:CM:153:LEU:HD22	1.82	0.43
6:CM:563:VAL:O	6:CM:567:GLN:HG3	2.18	0.43
6:CM:702:THR:O	6:CM:706:ILE:HG13	2.18	0.43
6:CM:788:MET:O	6:CM:792:TYR:HB3	2.18	0.43
6:CM:824:GLN:HG3	2:DR:83:ARG:NH1	2.33	0.43
9:CQ:264:PHE:CE2	9:DI:309:PRO:HG3	2.52	0.43
1:CR:8:ILE:CD1	2:CS:1:MET:HE1	2.47	0.43
1:CR:154:ASP:O	2:CZ:46:ALA:HA	2.17	0.43
1:CV:124:GLU:OE1	1:CV:124:GLU:N	2.45	0.43
2:CW:59:SER:HB3	2:CW:132:ALA:HB2	2.00	0.43
1:CY:59:PHE:CE2	8:DY:227:PRO:HG3	2.53	0.43
1:DA:21:GLY:O	1:DA:25:ARG:HG2	2.18	0.43
1:DC:37:LEU:CD1	2:DD:28:LYS:HG2	2.47	0.43
9:DI:42:PHE:CE2	9:DI:131:LEU:HB2	2.53	0.43
9:DI:193:VAL:HG22	9:DI:257:THR:OG1	2.18	0.43

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
9:DI:264:PHE:CD1	9:DI:294:PRO:HD3	2.53	0.43
2:DM:104:LEU:HD12	2:DM:155:TYR:CD2	2.52	0.43
1:DN:47:ARG:HB2	1:DN:89:LEU:HD23	2.00	0.43
1:DN:71:ASN:C	1:DN:71:ASN:HD22	2.26	0.43
1:DN:71:ASN:HB3	10:DN:200:CYC:OC	2.18	0.43
2:DO:47:ASN:O	2:DO:51:ILE:HG13	2.17	0.43
2:DT:2:GLN:NE2	2:DT:10:ASN:HD22	2.14	0.43
2:DT:68:PRO:HA	2:DT:73:TYR:CG	2.52	0.43
10:DU:200:CYC:HB	10:DU:200:CYC:CMA	2.31	0.43
9:EA:197:THR:HG21	9:EA:292:LEU:HD11	2.00	0.43
1:AA:115:MET:HE2	1:AA:115:MET:HB3	1.91	0.43
1:AA:130:VAL:HB	1:AA:157:ILE:HG22	2.00	0.43
1:AC:131:ARG:HE	1:AC:157:ILE:CG2	2.31	0.43
3:AE:101:LYS:HD2	3:AE:155:TYR:OH	2.18	0.43
2:AF:87:TYR:HB3	10:AF:200:CYC:CBB	2.48	0.43
2:AI:62:TYR:OH	10:AJ:200:CYC:O2D	2.30	0.43
2:AI:84:ASP:OD2	2:AI:116:TYR:OH	2.36	0.43
4:AK:161:THR:O	4:AK:165:SER:HB3	2.19	0.43
2:AU:15:GLN:HB2	2:AU:17:LYS:NZ	2.33	0.43
1:AV:6:LYS:O	1:AV:9:VAL:HG22	2.17	0.43
2:AW:36:LEU:HA	2:AW:39:ARG:HH12	1.83	0.43
2:AY:41:ALA:CB	2:AY:96:MET:HE1	2.45	0.43
2:AY:123:ILE:O	2:AY:127:VAL:HG23	2.18	0.43
6:BD:278:LYS:CD	6:BD:310:ARG:HG3	2.44	0.43
6:BD:866:PHE:CD2	2:DB:115:THR:HG23	2.53	0.43
9:BH:34:GLU:HG2	9:BH:174:GLU:CG	2.48	0.43
9:BH:41:TRP:CZ2	9:BH:45:LEU:HD21	2.54	0.43
2:BJ:75:THR:OG1	1:BK:108:GLY:HA2	2.18	0.43
2:BL:147:LYS:O	2:BL:151:VAL:HG23	2.19	0.43
1:BP:47:ARG:HG3	1:BP:48:GLU:N	2.33	0.43
4:BS:41:ALA:HA	4:BS:44:ILE:HG12	2.00	0.43
4:BS:113:LEU:HD21	10:BS:200:CYC:CMB	2.48	0.43
2:BT:144:ASP:HB2	7:CO:86:VAL:HG12	1.99	0.43
1:BV:33:GLY:HA2	1:BV:36:ARG:HG2	1.99	0.43
2:BY:53:LYS:HD2	1:BZ:118:SER:O	2.17	0.43
2:CA:60:LEU:HD11	2:CA:129:ALA:HB2	2.00	0.43
1:CE:11:ALA:CB	1:CE:18:LEU:HD23	2.47	0.43
1:CE:42:THR:HG21	1:CE:141:MET:HB3	1.99	0.43
1:CE:58:LEU:HD22	1:CE:129:SER:CB	2.38	0.43
2:CH:51:ILE:HG21	2:CH:137:THR:HG22	2.01	0.43
5:CJ:40:GLN:O	5:CJ:44:MET:HG2	2.19	0.43

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
5:CK:40:GLN:CG	5:CK:44:MET:HE3	2.44	0.43
6:CM:556:TYR:HB2	6:CM:559:GLN:HG3	1.99	0.43
6:CM:627:ALA:HB2	6:CM:635:VAL:HG21	2.01	0.43
6:CM:806:LYS:HG3	6:CM:873:TYR:CE2	2.53	0.43
6:CM:887:SER:OG	2:DR:119:LEU:HD21	2.18	0.43
1:CR:39:ILE:HD11	1:CR:145:ASP:CB	2.48	0.43
1:CR:57:ARG:HB3	1:CR:132:GLU:OE2	2.18	0.43
1:CV:141:MET:HG2	1:CV:145:ASP:HB2	1.99	0.43
2:CW:96:MET:SD	2:CW:97:LEU:HD22	2.57	0.43
2:DD:58:LYS:HA	8:DH:206:GLN:OE1	2.18	0.43
9:DI:155:ARG:HA	9:DI:158:VAL:HG22	2.00	0.43
9:DI:191:GLU:HB2	9:DI:254:ARG:HB2	2.00	0.43
9:DI:293:ASN:HB2	9:DI:296:GLY:C	2.43	0.43
1:DJ:25:ARG:HG2	1:DQ:25:ARG:CD	2.47	0.43
10:DJ:200:CYC:H2C	10:DJ:200:CYC:HBC2	1.74	0.43
1:DL:21:GLY:HA3	1:DU:25:ARG:NH2	2.34	0.43
1:DL:75:GLU:HA	1:DL:78:THR:OG1	2.18	0.43
1:DN:5:THR:HA	2:DO:1:MET:HE1	1.99	0.43
1:DU:87:TYR:HB3	10:DU:200:CYC:CBB	2.48	0.43
2:AB:71:ASN:ND2	2:AB:121:VAL:HA	2.33	0.43
2:AB:123:ILE:H	2:AB:123:ILE:HD12	1.83	0.43
2:AD:1:MET:HA	2:AD:102:SER:OG	2.18	0.43
2:AF:114:GLU:HG3	5:BA:53:VAL:CG1	2.48	0.43
2:AI:122:PRO:HG2	2:AI:125:SER:OG	2.18	0.43
1:AJ:37:LEU:HD11	4:AK:31:PHE:CE2	2.54	0.43
4:AK:113:LEU:HB3	4:AK:164:LEU:HD13	2.01	0.43
2:AO:114:GLU:HA	2:AO:117:ASN:HB2	2.01	0.43
2:AQ:100:ASP:OD1	2:AQ:102:SER:N	2.47	0.43
1:AR:71:ASN:C	1:AR:71:ASN:HD22	2.27	0.43
2:AS:19:LEU:HB2	2:AS:24:MET:HE2	1.99	0.43
2:AS:148:GLU:HG3	2:AS:152:TYR:CE2	2.54	0.43
1:AV:115:MET:O	1:AV:119:LEU:HD23	2.19	0.43
2:AW:116:TYR:CE2	2:AW:160:LEU:HD21	2.54	0.43
2:AY:71:ASN:HD21	2:AY:121:VAL:HG22	1.83	0.43
6:BD:514:ARG:HA	6:BD:652:ASP:O	2.18	0.43
1:BI:91:LEU:HD21	1:BI:107:ILE:HG21	2.01	0.43
1:BI:126:VAL:HG22	10:BI:200:CYC:HMC3	1.99	0.43
2:BJ:36:LEU:HD13	2:BJ:39:ARG:HH22	1.82	0.43
1:BK:89:LEU:O	1:BK:93:THR:HG23	2.17	0.43
2:BL:64:ASP:O	2:BL:67:ARG:HB3	2.18	0.43
4:BS:120:LEU:HD21	10:BS:200:CYC:HAA2	2.00	0.43

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:BT:52:VAL:O	2:BT:56:VAL:HG23	2.19	0.43
2:BT:68:PRO:HA	2:BT:73:TYR:CD1	2.53	0.43
2:BT:90:ARG:HD3	6:CM:33:ARG:O	2.18	0.43
2:BT:126:THR:OG1	10:BT:200:CYC:HMC2	2.18	0.43
1:BX:83:ARG:NH2	1:BX:84:ASP:OD1	2.35	0.43
1:CG:6:LYS:HG2	6:CM:534:PHE:CZ	2.42	0.43
5:CJ:4:PHE:HB2	5:CJ:31:VAL:HG13	1.99	0.43
5:CJ:46:MET:HE2	6:CM:384:GLN:HA	2.00	0.43
6:CM:163:THR:HG21	6:CM:224:PHE:CZ	2.54	0.43
6:CM:274:SER:HG	6:CM:277:GLU:CD	2.27	0.43
1:CR:18:LEU:H	1:CR:18:LEU:HD12	1.83	0.43
1:CR:90:ARG:NH1	1:CR:90:ARG:HB3	2.33	0.43
2:CS:52:VAL:HG21	2:CS:86:ASP:OD1	2.18	0.43
2:CU:122:PRO:HD2	10:CU:200:CYC:C1C	2.48	0.43
1:CV:37:LEU:CD1	2:CW:28:LYS:HE3	2.48	0.43
2:CW:123:ILE:HA	2:CW:126:THR:CG2	2.47	0.43
1:CY:132:GLU:O	1:CY:136:VAL:HG23	2.18	0.43
2:CZ:81:CYS:HA	10:CZ:200:CYC:HAC1	1.76	0.43
2:CZ:112:LEU:CD1	10:CZ:200:CYC:HMB1	2.48	0.43
1:DC:109:LEU:HB3	1:DC:159:LYS:HG2	2.01	0.43
1:DC:143:SER:O	1:DC:146:ALA:HB3	2.18	0.43
9:DI:39:LEU:HD11	9:DI:139:LEU:HB2	1.99	0.43
9:DI:219:SER:HA	9:DI:234:LYS:CE	2.36	0.43
2:DK:76:ARG:NH2	10:DK:200:CYC:O2D	2.29	0.43
1:DL:7:SER:HB3	1:DL:18:LEU:CD2	2.49	0.43
2:DM:112:LEU:CD2	2:DM:160:LEU:HD21	2.49	0.43
10:DO:200:CYC:HB	10:DO:200:CYC:CMA	2.32	0.43
10:DR:200:CYC:HHA	10:DR:200:CYC:HBD1	2.01	0.43
1:DS:71:ASN:C	1:DS:71:ASN:HD22	2.26	0.43
2:DV:30:TYR:OH	2:DV:97:LEU:O	2.32	0.43
2:DV:116:TYR:CD1	2:DV:121:VAL:HG21	2.54	0.43
9:EA:248:LEU:HD21	9:EA:277:TRP:HB2	2.00	0.43
9:EA:316:ALA:O	9:EA:317:ARG:HG2	2.18	0.43
1:AA:105:GLU:OE1	1:AA:109:LEU:HD23	2.19	0.43
2:AD:53:LYS:HD3	3:AE:120:ASN:ND2	2.34	0.43
2:AD:67:ARG:HG3	2:AD:68:PRO:HD2	2.00	0.43
10:AH:200:CYC:HAA1	2:AL:61:LEU:HD22	2.01	0.43
1:AJ:27:LYS:HB2	4:AK:38:ILE:HD13	1.99	0.43
2:AL:148:GLU:O	2:AL:151:VAL:HG22	2.18	0.43
1:AN:57:ARG:HA	1:AN:60:GLN:CG	2.48	0.43
1:AN:154:ASP:HB3	2:AU:46:ALA:CB	2.48	0.43

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:AP:116:TYR:CD1	1:AP:121:THR:HB	2.54	0.43
1:AR:13:ALA:HB2	6:BD:589:LEU:HD11	1.99	0.43
1:AV:46:SER:O	1:AV:50:ILE:HG12	2.18	0.43
1:AX:85:MET:C	1:AX:133:MET:HE1	2.43	0.43
1:AZ:90:ARG:HG2	1:AZ:94:TYR:CE2	2.53	0.43
6:BD:321:ARG:HG2	6:BD:325:LYS:HE3	2.00	0.43
9:BH:125:ILE:HG13	9:BH:126:PRO:HD2	2.00	0.43
9:BH:146:GLU:O	9:BH:150:GLN:N	2.45	0.43
9:BH:172:ILE:H	9:BH:172:ILE:HD12	1.83	0.43
1:BI:50:ILE:HD12	1:BI:136:VAL:CG1	2.49	0.43
2:BJ:105:ASP:OD2	2:BJ:155:TYR:OH	2.36	0.43
2:BJ:133:ILE:O	2:BJ:137:THR:HG22	2.19	0.43
2:BL:8:VAL:CG1	2:BL:27:LEU:HD21	2.46	0.43
2:BN:56:VAL:HG12	2:BN:61:LEU:HG	2.00	0.43
2:BQ:104:LEU:CD1	2:BQ:152:TYR:HB3	2.48	0.43
2:BQ:123:ILE:O	2:BQ:126:THR:HG22	2.18	0.43
4:BS:63:ILE:HG22	4:BS:66:LEU:HG	2.00	0.43
4:BS:63:ILE:HD13	4:BS:126:PRO:HG3	2.01	0.43
1:BV:160:MET:HE3	1:BV:160:MET:HB2	1.86	0.43
1:BX:30:VAL:HG11	2:BY:34:GLY:HA3	2.01	0.43
1:BX:141:MET:HG3	1:BX:146:ALA:HB2	2.01	0.43
1:CG:23:LEU:HD22	2:CH:38:VAL:HG13	2.00	0.43
2:CH:122:PRO:O	2:CH:126:THR:HG22	2.18	0.43
5:CJ:6:ILE:HD13	5:CJ:36:TRP:CZ3	2.54	0.43
6:CM:269:MET:HB2	6:CM:281:VAL:HG11	2.00	0.43
8:CP:216:MET:HE3	9:CQ:12:PHE:HB2	2.00	0.43
2:CS:59:SER:HB3	2:CS:132:ALA:HB2	2.00	0.43
1:CT:68:PRO:HG3	1:CT:73:TYR:CE2	2.54	0.43
2:CU:65:VAL:HG12	2:CU:72:MET:CG	2.49	0.43
2:CW:73:TYR:O	2:CW:74:THR:OG1	2.23	0.43
1:CY:39:ILE:CA	1:CY:42:THR:HG22	2.46	0.43
1:DA:58:LEU:HD22	1:DA:129:SER:HB2	2.01	0.43
9:DI:92:THR:OG1	9:DI:95:CYS:SG	2.52	0.43
1:DJ:69:GLY:O	9:EA:58:ALA:HB2	2.18	0.43
2:DK:51:ILE:HD12	2:DK:51:ILE:H	1.83	0.43
1:DL:91:LEU:HD21	1:DL:107:ILE:HG22	2.00	0.43
2:DO:54:GLU:HG3	9:EA:63:LEU:CD2	2.49	0.43
2:DO:122:PRO:HG2	2:DO:125:SER:HB2	2.01	0.43
10:DQ:200:CYC:HBC2	10:DQ:200:CYC:H2C	1.70	0.43
2:DT:3:ASP:HB3	2:DT:98:ALA:HB1	2.01	0.43
1:DU:53:GLN:O	1:DU:57:ARG:HG3	2.18	0.43

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
5:DX:4:PHE:O	5:DX:30:LEU:HA	2.18	0.43
9:EA:250:LEU:HD23	9:EA:273:VAL:HG22	2.00	0.43
1:AC:128:GLN:NE2	1:AC:132:GLU:OE2	2.47	0.43
3:AE:47:GLU:CG	3:AE:89:LEU:HD23	2.48	0.43
10:AK:200:CYC:HBA2	6:BD:401:LEU:HD11	1.99	0.43
2:AL:31:PHE:CD1	6:BD:50:GLY:HA3	2.53	0.43
1:AN:8:ILE:HD13	1:AN:18:LEU:HD21	2.00	0.43
1:AN:58:LEU:HD13	1:AN:129:SER:N	2.34	0.43
2:AS:122:PRO:HB2	2:AS:125:SER:HB2	2.00	0.43
1:AX:93:THR:HA	1:AX:96:VAL:HG22	2.01	0.43
2:AY:77:ARG:HB3	10:AY:200:CYC:HMD1	2.01	0.43
1:AZ:116:TYR:CE2	1:AZ:126:VAL:HG11	2.50	0.43
5:BA:5:ARG:HE	5:BA:30:LEU:HD21	1.82	0.43
5:BA:18:THR:HG22	6:BD:240:VAL:O	2.19	0.43
6:BD:309:VAL:HG21	6:BD:319:PHE:CD1	2.53	0.43
9:BH:257:THR:HG23	9:BH:267:ILE:CG1	2.46	0.43
2:BJ:104:LEU:HD11	2:BJ:152:TYR:HB3	2.00	0.43
1:BK:6:LYS:HD2	1:BK:6:LYS:HA	1.84	0.43
1:BP:30:VAL:HG13	2:BQ:31:PHE:HA	2.00	0.43
2:BQ:64:ASP:OD1	2:BW:124:SER:OG	2.11	0.43
2:BQ:75:THR:HG23	1:BR:108:GLY:HA2	2.00	0.43
2:BQ:119:LEU:HD23	6:CM:435:ALA:CB	2.49	0.43
1:BR:5:THR:HG23	4:BS:1:MET:SD	2.58	0.43
1:BR:5:THR:HG23	4:BS:1:MET:HG3	2.01	0.43
2:BT:112:LEU:HD21	2:BT:160:LEU:CD2	2.49	0.43
1:BV:76:GLU:OE1	7:CO:111:LYS:HE3	2.19	0.43
2:BW:137:THR:O	2:BW:141:VAL:HG23	2.18	0.43
1:BZ:77:MET:HE1	10:BZ:200:CYC:CGD	2.48	0.43
2:CD:71:ASN:O	2:CD:77:ARG:HD2	2.19	0.43
2:CF:68:PRO:HA	2:CF:73:TYR:CG	2.53	0.43
2:CH:133:ILE:O	2:CH:137:THR:HG23	2.19	0.43
5:CK:30:LEU:HD12	6:CM:572:ASP:HB3	2.00	0.43
6:CM:141:ILE:HD12	6:CM:141:ILE:H	1.83	0.43
1:CT:66:VAL:HG21	8:DY:235:VAL:HG11	2.00	0.43
1:CY:20:PRO:O	1:CY:23:LEU:HG	2.17	0.43
1:DC:98:SER:OG	1:DC:100:ASP:OD1	2.23	0.43
1:DJ:115:MET:HG3	2:DO:78:TYR:CE1	2.53	0.43
1:DJ:130:VAL:CG1	1:DJ:157:ILE:HG12	2.36	0.43
2:DK:92:ALA:CB	2:DK:149:MET:HE1	2.49	0.43
1:DQ:26:ILE:O	1:DQ:30:VAL:HG13	2.18	0.43
1:DQ:79:ALA:HB3	8:DZ:203:PHE:CZ	2.54	0.43

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
9:EA:101:TRP:HB3	9:EA:105:ILE:HB	1.99	0.43
9:EA:155:ARG:HA	9:EA:158:VAL:HG22	2.00	0.43
2:AD:109:LEU:HD21	2:AD:159:GLY:HA3	2.00	0.43
3:AE:21:GLY:HA3	1:AJ:3:ILE:HD11	2.01	0.43
2:AF:52:VAL:O	2:AF:56:VAL:HG23	2.19	0.43
1:AH:81:CYS:HA	10:AH:200:CYC:HBC3	2.00	0.43
1:AH:85:MET:O	1:AH:133:MET:HE1	2.19	0.43
4:AK:1:MET:HB2	4:AK:104:VAL:HB	2.01	0.43
4:AK:51:ILE:HA	4:AK:137:MET:CE	2.47	0.43
2:AL:3:ASP:OD1	2:AL:6:THR:HG22	2.18	0.43
2:AQ:1:MET:SD	2:AQ:103:ILE:HD12	2.58	0.43
2:AU:16:GLY:O	1:AZ:90:ARG:NH1	2.48	0.43
2:AU:62:TYR:N	10:AV:200:CYC:O2A	2.50	0.43
1:AZ:109:LEU:HD13	1:AZ:160:MET:HE2	2.01	0.43
6:BD:85:LEU:HA	6:BD:153:LEU:CD2	2.48	0.43
6:BD:485:ILE:HG21	6:BD:665:LEU:HD13	1.99	0.43
6:BD:598:VAL:HB	10:BD:901:CYC:HBB3	2.00	0.43
6:BD:795:TYR:CE1	6:BD:881:LYS:HE3	2.54	0.43
9:BH:98:TYR:OH	9:BH:158:VAL:HG22	2.19	0.43
2:BJ:60:LEU:HD22	2:BJ:65:VAL:HG21	2.01	0.43
1:BK:76:GLU:HG2	1:BK:77:MET:HE3	2.00	0.43
2:BL:129:ALA:O	2:BL:133:ILE:HG13	2.18	0.43
3:BM:72:ALA:HA	3:BM:77:GLN:HB3	2.00	0.43
4:BS:61:GLU:OE1	4:BS:61:GLU:N	2.51	0.43
4:BS:82:CYS:HA	10:BS:200:CYC:HAC1	1.73	0.43
2:BT:31:PHE:HD2	6:CM:54:LEU:HD11	1.84	0.43
1:BV:101:VAL:HG12	1:BV:152:TYR:CD1	2.52	0.43
1:BV:128:GLN:HA	1:BV:131:ARG:HD2	1.99	0.43
1:BX:19:SER:OG	1:BX:20:PRO:HD2	2.18	0.43
2:BY:68:PRO:HG3	2:BY:73:TYR:CE1	2.53	0.43
1:BZ:126:VAL:O	1:BZ:130:VAL:HG23	2.18	0.43
2:CA:126:THR:O	2:CA:130:ILE:HG13	2.18	0.43
1:CC:140:LEU:O	1:CC:140:LEU:HD23	2.19	0.43
2:CD:75:THR:HG22	10:CE:200:CYC:C4B	2.49	0.43
1:CG:39:ILE:O	1:CG:43:LEU:HG	2.18	0.43
6:CM:353:ARG:HA	6:CM:418:PHE:HZ	1.83	0.43
7:CO:114:ALA:CA	7:CO:117:VAL:HG12	2.41	0.43
9:CQ:112:ARG:HD2	9:CQ:112:ARG:HA	1.85	0.43
1:CR:126:VAL:HG12	1:CR:160:MET:CE	2.49	0.43
1:CR:142:SER:O	1:CR:146:ALA:HB2	2.19	0.43
1:CT:63:PRO:HD2	1:CY:69:GLY:HA3	2.00	0.43

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:CU:160:LEU:HD23	2:CU:160:LEU:HA	1.86	0.43
2:CZ:85:LEU:HG	10:CZ:200:CYC:HBC1	2.00	0.43
2:CZ:101:ALA:HB1	2:CZ:155:TYR:CE2	2.53	0.43
1:DA:53:GLN:HG2	1:DA:136:VAL:HG21	1.99	0.43
2:DD:91:TYR:HB3	2:DD:104:LEU:HD23	2.00	0.43
9:DI:204:ASN:CB	9:DI:209:ASP:HB3	2.36	0.43
1:DJ:4:VAL:O	1:DJ:8:ILE:HG23	2.19	0.43
1:DJ:11:ALA:CB	1:DJ:18:LEU:HD23	2.44	0.43
1:DJ:50:ILE:CD1	1:DJ:140:LEU:HD21	2.48	0.43
1:DQ:33:GLY:HA3	2:DR:31:PHE:CD2	2.54	0.43
2:DR:74:THR:OG1	2:DR:77:ARG:HG3	2.19	0.43
1:DS:58:LEU:HD22	1:DS:129:SER:HB2	2.00	0.43
1:DU:27:LYS:HZ2	2:DV:35:GLU:CD	2.27	0.43
1:DU:115:MET:HE1	10:DU:200:CYC:CMB	2.48	0.43
5:DX:6:ILE:HD13	5:DX:36:TRP:CZ3	2.53	0.43
9:EA:23:ALA:O	9:EA:27:ARG:HG3	2.18	0.43
2:AB:71:ASN:C	2:AB:77:ARG:HE	2.27	0.43
1:AC:4:VAL:O	1:AC:8:ILE:HG13	2.19	0.43
2:AD:37:ARG:HH21	2:AD:99:GLY:HA2	1.84	0.43
1:AH:131:ARG:HA	1:AH:157:ILE:HD11	2.00	0.43
1:AJ:130:VAL:HG21	1:AJ:160:MET:SD	2.59	0.43
2:AL:91:TYR:O	2:AL:104:LEU:HD21	2.19	0.43
1:AN:131:ARG:CG	1:AN:157:ILE:HD13	2.49	0.43
2:AS:119:LEU:HD21	6:BD:447:TYR:CD1	2.54	0.43
1:AX:109:LEU:O	1:AX:112:VAL:HG12	2.19	0.43
1:AZ:50:ILE:HG12	1:AZ:136:VAL:HG12	2.00	0.43
1:AZ:91:LEU:O	1:AZ:104:ILE:HG12	2.18	0.43
6:BD:141:ILE:HD12	6:BD:141:ILE:H	1.84	0.43
6:BD:785:GLU:O	6:BD:788:MET:HG3	2.19	0.43
6:BD:864:ALA:HB2	2:DB:110:ASN:O	2.19	0.43
8:BG:198:ASN:HA	8:BG:201:ARG:NH2	2.33	0.43
1:BI:67:SER:HB3	1:BI:68:PRO:HD2	1.99	0.43
2:BJ:71:ASN:CG	2:BJ:122:PRO:HD3	2.43	0.43
2:BJ:134:LYS:HB2	2:BJ:153:LEU:HD13	2.01	0.43
2:BL:141:VAL:HG23	2:BL:145:ALA:HB3	2.00	0.43
3:BM:2:SER:O	3:BM:5:SER:OG	2.28	0.43
1:BP:37:LEU:O	1:BP:40:ALA:HB3	2.18	0.43
1:BP:130:VAL:HG21	1:BP:160:MET:HE3	2.01	0.43
4:BS:66:LEU:O	4:BS:73:ALA:HB3	2.18	0.43
1:CE:126:VAL:O	1:CE:130:VAL:HG23	2.19	0.43
2:CF:72:MET:HE2	2:CF:78:TYR:CE2	2.53	0.43

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:CF:96:MET:HE3	2:CF:96:MET:HB3	1.79	0.43
6:CM:79:GLY:N	6:CM:141:ILE:HD13	2.33	0.43
6:CM:203:ARG:HG3	6:CM:224:PHE:CB	2.48	0.43
2:CS:51:ILE:HD12	2:CS:51:ILE:H	1.83	0.43
1:CT:121:THR:HG23	10:CT:200:CYC:C1C	2.49	0.43
1:CY:51:VAL:HG22	1:CY:82:LEU:HG	2.00	0.43
2:CZ:36:LEU:CD1	2:CZ:145:ALA:HB2	2.36	0.43
1:DA:75:GLU:HA	1:DA:78:THR:OG1	2.19	0.43
2:DB:51:ILE:HD11	2:DB:140:LEU:CD2	2.45	0.43
10:DC:200:CYC:CMA	10:DC:200:CYC:HB	2.32	0.43
2:DD:112:LEU:CG	2:DD:160:LEU:HD13	2.36	0.43
9:DI:98:TYR:OH	9:DI:106:LYS:HB3	2.18	0.43
1:DJ:71:ASN:HD22	1:DJ:121:THR:HA	1.83	0.43
1:DJ:111:GLY:HA2	1:DJ:114:GLU:OE1	2.19	0.43
2:DK:91:TYR:CE2	2:DK:108:VAL:HB	2.53	0.43
2:DM:22:ALA:HB1	2:DM:26:LYS:NZ	2.34	0.43
2:DO:2:GLN:HB2	2:DO:102:SER:OG	2.19	0.43
2:DR:136:VAL:O	2:DR:140:LEU:HD23	2.19	0.43
1:DS:33:GLY:HA3	2:DT:31:PHE:CE2	2.54	0.43
1:DS:85:MET:HE2	1:DS:133:MET:CG	2.49	0.43
1:DS:90:ARG:HG3	2:DT:18:TYR:CE1	2.53	0.43
2:DT:103:ILE:O	2:DT:107:ARG:HB2	2.18	0.43
1:DU:116:TYR:HE2	1:DU:126:VAL:HG11	1.84	0.43
9:EA:251:ILE:HD12	9:EA:274:GLN:CD	2.44	0.43
1:AC:14:GLU:HG2	1:AC:16:ARG:HE	1.83	0.43
1:AC:78:THR:O	1:AC:82:LEU:HD23	2.18	0.43
2:AF:72:MET:HE3	2:AF:81:CYS:HB3	2.00	0.43
1:AH:40:ALA:CB	1:AH:97:VAL:HG22	2.49	0.43
1:AN:113:ARG:NH2	1:AN:161:SER:O	2.52	0.43
1:AP:116:TYR:HD1	1:AP:121:THR:HB	1.84	0.43
1:AR:47:ARG:O	1:AR:51:VAL:HG23	2.19	0.43
2:AU:62:TYR:OH	1:AV:80:THR:HG21	2.19	0.43
1:AV:107:ILE:HG22	10:AV:200:CYC:HBB1	2.01	0.43
1:AX:68:PRO:HA	1:AX:73:TYR:CG	2.54	0.43
6:BD:161:ARG:HG2	6:BD:165:TYR:CZ	2.54	0.43
6:BD:208:ASP:OD1	6:BD:209:TYR:N	2.52	0.43
6:BD:524:VAL:O	6:BD:524:VAL:HG12	2.18	0.43
6:BD:889:THR:HG22	2:CZ:118:SER:O	2.19	0.43
1:BK:97:VAL:HG11	2:BL:19:LEU:CD1	2.49	0.43
3:BM:47:GLU:CG	3:BM:89:LEU:HD23	2.48	0.43
2:BN:2:GLN:HB3	2:BN:102:SER:CB	2.48	0.43

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:BP:23:LEU:HD13	2:BQ:42:SER:OG	2.18	0.43
1:BP:81:CYS:HA	10:BP:200:CYC:CHD	2.49	0.43
1:BR:67:SER:O	1:BR:73:TYR:HB2	2.18	0.43
4:BS:44:ILE:CG2	4:BS:138:ILE:HD12	2.43	0.43
1:BV:8:ILE:HD13	1:BV:18:LEU:HD21	1.99	0.43
1:BV:107:ILE:O	2:CA:75:THR:HG23	2.18	0.43
2:CA:51:ILE:HG23	2:CA:136:VAL:CG1	2.48	0.43
2:CA:147:LYS:HE3	7:CO:93:PRO:HD3	2.01	0.43
1:CE:46:SER:O	1:CE:50:ILE:HG12	2.19	0.43
6:CM:485:ILE:CD1	6:CM:665:LEU:HD13	2.47	0.43
6:CM:840:ASN:CG	5:DX:35:ASN:HD21	2.27	0.43
9:CQ:249:LYS:C	9:CQ:250:LEU:HD23	2.44	0.43
1:CR:122:PRO:HD2	10:CR:200:CYC:OC	2.19	0.43
1:CT:66:VAL:HG23	1:CT:78:THR:CG2	2.48	0.43
2:CU:1:MET:SD	2:CU:103:ILE:HB	2.58	0.43
1:CV:85:MET:HE1	1:CV:129:SER:HB2	2.00	0.43
1:CY:85:MET:HB3	1:CY:133:MET:HE2	2.01	0.43
1:DA:39:ILE:HD12	1:DA:145:ASP:HB3	2.01	0.43
8:DG:213:GLY:HA2	1:DQ:83:ARG:HE	1.84	0.43
8:DG:221:MET:HE3	1:DQ:52:LYS:CA	2.49	0.43
9:DI:193:VAL:HG23	9:DI:255:GLY:C	2.44	0.43
9:DI:260:ALA:HB3	9:DI:264:PHE:CB	2.49	0.43
2:DK:15:GLN:HG3	2:DK:17:LYS:HG2	2.01	0.43
2:DK:41:ALA:N	2:DK:96:MET:HE1	2.33	0.43
1:DL:59:PHE:HE1	1:DL:65:ILE:HD11	1.84	0.43
1:DL:101:VAL:HB	1:DL:155:PHE:CE2	2.54	0.43
2:DM:71:ASN:HD22	2:DM:122:PRO:CD	2.32	0.43
1:DN:124:GLU:HG2	1:DN:125:ALA:N	2.34	0.43
2:DO:44:ILE:HD13	2:DO:149:MET:HE2	2.01	0.43
2:DO:50:THR:O	2:DO:54:GLU:HG2	2.19	0.43
2:DO:116:TYR:HE2	2:DO:160:LEU:HD21	1.84	0.43
2:DR:123:ILE:O	2:DR:127:VAL:HG23	2.19	0.43
2:DT:41:ALA:N	2:DT:96:MET:HE1	2.34	0.43
9:EA:49:LYS:HE3	9:EA:49:LYS:HB3	1.80	0.43
9:EA:223:LEU:HA	9:EA:301:VAL:HG12	2.01	0.43
3:AE:126:MET:HG2	3:AE:160:MET:HE3	2.00	0.43
2:AI:44:ILE:HG21	2:AI:93:THR:HG21	2.00	0.43
2:AI:58:LYS:NZ	2:AI:135:GLU:OE1	2.52	0.43
4:AK:72:ASN:ND2	4:AK:122:VAL:HG13	2.33	0.43
1:AN:8:ILE:HD12	2:AO:94:TYR:HD1	1.83	0.43
2:AS:110:ASN:O	6:BD:465:GLU:HB2	2.18	0.43

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
6:BD:799:LYS:CB	6:BD:883:LEU:HD21	2.47	0.43
1:BI:3:ILE:HG21	1:BP:25:ARG:HD3	2.00	0.43
2:BJ:77:ARG:HB3	10:BJ:200:CYC:HMD1	2.01	0.43
2:BL:125:SER:OG	10:BL:200:CYC:HMC1	2.19	0.43
2:BN:126:THR:CG2	10:BN:200:CYC:HMC1	2.49	0.43
1:BP:26:ILE:O	1:BP:30:VAL:HG23	2.19	0.43
2:BW:116:TYR:HB3	2:BW:121:VAL:HB	2.01	0.43
2:BW:144:ASP:OD1	2:BW:145:ALA:N	2.52	0.43
1:BX:97:VAL:HG11	2:BY:27:LEU:HD13	2.00	0.43
2:CA:124:SER:O	2:CA:128:GLN:HG2	2.19	0.43
1:CC:109:LEU:HD21	1:CC:159:LYS:HB3	2.01	0.43
2:CD:72:MET:HG2	2:CD:78:TYR:HD2	1.84	0.43
2:CD:148:GLU:HG3	2:CD:152:TYR:HE2	1.82	0.43
5:CK:21:GLU:OE1	5:CK:24:ASN:ND2	2.41	0.43
6:CM:151:LYS:HD3	6:CM:154:ARG:HH21	1.84	0.43
6:CM:208:ASP:OD1	6:CM:209:TYR:N	2.52	0.43
6:CM:371:GLY:HA3	6:CM:375:ALA:CB	2.48	0.43
6:CM:725:LYS:HG2	6:CM:852:THR:OG1	2.19	0.43
9:CQ:76:PRO:CB	9:CQ:123:ALA:HB2	2.46	0.43
1:CT:17:TYR:CD1	2:CU:90:ARG:HD3	2.54	0.43
1:CT:81:CYS:HB3	10:CT:200:CYC:HBC1	1.88	0.43
1:CT:97:VAL:HG21	2:CU:19:LEU:HD11	2.00	0.43
1:CY:48:GLU:HG2	1:CY:49:THR:N	2.34	0.43
1:CY:65:ILE:HG23	10:CY:200:CYC:OC	2.19	0.43
1:DA:105:GLU:HA	1:DA:109:LEU:HB2	2.01	0.43
9:DI:248:LEU:HD11	9:DI:277:TRP:CE3	2.53	0.43
1:DL:4:VAL:CG2	2:DM:97:LEU:HD23	2.49	0.43
2:DM:71:ASN:HD22	2:DM:121:VAL:HA	1.82	0.43
2:DM:71:ASN:HD22	2:DM:122:PRO:HD2	1.83	0.43
2:DM:87:TYR:OH	5:DX:19:GLN:O	2.34	0.43
1:DU:102:THR:O	1:DU:106:GLU:HG2	2.19	0.43
5:DX:27:PHE:HE1	5:DX:29:LYS:HE3	1.84	0.43
10:AA:200:CYC:O2A	2:AF:62:TYR:N	2.41	0.42
1:AC:14:GLU:HB3	1:AC:16:ARG:CD	2.49	0.42
1:AC:124:GLU:HG2	1:AC:125:ALA:N	2.34	0.42
2:AD:15:GLN:HB3	2:AD:17:LYS:HE2	2.01	0.42
2:AD:112:LEU:HA	2:AD:115:THR:CG2	2.48	0.42
1:AN:27:LYS:O	1:AN:30:VAL:HG22	2.19	0.42
2:AY:23:ALA:O	2:AY:27:LEU:HG	2.18	0.42
6:BD:210:PHE:HZ	6:BD:216:ALA:HB1	1.84	0.42
6:BD:532:VAL:O	6:BD:532:VAL:HG12	2.18	0.42

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
6:BD:603:GLU:OE2	6:BD:607:ARG:NE	2.24	0.42
1:BK:131:ARG:O	1:BK:134:LYS:HB3	2.19	0.42
2:BL:112:LEU:CA	2:BL:115:THR:HG22	2.48	0.42
2:BN:36:LEU:HD11	2:BN:145:ALA:HB2	2.00	0.42
2:BQ:87:TYR:CD2	10:BQ:200:CYC:HAB1	2.54	0.42
1:BR:71:ASN:C	1:BR:71:ASN:HD22	2.27	0.42
2:BT:92:ALA:HB3	2:BT:149:MET:HE1	2.01	0.42
2:BW:102:SER:O	2:BW:106:GLU:HG3	2.19	0.42
1:BX:10:ASN:OD1	6:CM:511:LYS:HE3	2.19	0.42
1:BZ:95:GLY:HA3	1:BZ:104:ILE:HD11	2.01	0.42
2:CA:11:SER:O	2:CA:14:VAL:HG12	2.19	0.42
1:CC:68:PRO:HA	1:CC:73:TYR:CG	2.54	0.42
1:CE:57:ARG:O	1:CE:60:GLN:NE2	2.52	0.42
1:CE:85:MET:HG3	10:CE:200:CYC:HBC1	2.01	0.42
1:CE:142:SER:O	1:CE:146:ALA:N	2.51	0.42
2:CH:123:ILE:O	2:CH:127:VAL:HG23	2.19	0.42
5:CJ:15:ARG:HH12	5:CJ:23:GLN:NE2	2.16	0.42
6:CM:238:ASN:HB3	6:CM:253:LEU:O	2.19	0.42
6:CM:811:ARG:HG2	6:CM:812:ALA:O	2.19	0.42
6:CM:818:GLU:HG2	6:CM:822:TYR:CE2	2.53	0.42
1:CR:131:ARG:HA	1:CR:157:ILE:HD11	2.01	0.42
2:CS:57:ALA:HA	2:CS:61:LEU:CD1	2.44	0.42
2:CU:64:ASP:OD1	2:CU:67:ARG:NH2	2.33	0.42
2:CU:109:LEU:HD13	2:CU:159:GLY:HA3	2.01	0.42
1:CV:37:LEU:HD21	2:CW:27:LEU:CD1	2.49	0.42
1:CY:71:ASN:HD22	1:CY:122:PRO:HD2	1.84	0.42
2:CZ:126:THR:HG22	10:CZ:200:CYC:H3C	2.00	0.42
1:DA:97:VAL:O	2:DB:5:ILE:HD11	2.19	0.42
2:DB:2:GLN:CD	2:DB:7:ALA:HA	2.44	0.42
8:DG:242:ILE:HD11	1:DL:83:ARG:HB2	2.00	0.42
8:DH:219:LEU:HD21	8:DY:203:PHE:HZ	1.83	0.42
8:DH:240:ILE:CD1	1:DJ:79:ALA:HB2	2.42	0.42
9:DI:28:PHE:CE2	9:DI:39:LEU:HD23	2.53	0.42
9:DI:197:THR:HG21	9:DI:292:LEU:HD11	2.01	0.42
9:DI:201:TYR:HD1	9:DI:213:LEU:CD1	2.32	0.42
1:DN:116:TYR:HE2	1:DN:126:VAL:HG11	1.84	0.42
2:DR:134:LYS:HA	2:DR:153:LEU:CD1	2.48	0.42
1:DS:21:GLY:O	1:DS:25:ARG:HG2	2.18	0.42
10:DT:200:CYC:HHA	10:DT:200:CYC:O2A	2.19	0.42
1:DU:100:ASP:OD1	1:DU:101:VAL:N	2.50	0.42
1:DU:141:MET:CE	1:DU:145:ASP:HB3	2.49	0.42

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
9:EA:32:ASN:HA	9:EA:170:LYS:HZ3	1.83	0.42
2:AB:41:ALA:CB	2:AB:97:LEU:HD21	2.50	0.42
3:AE:17:TYR:CE1	2:AF:90:ARG:HA	2.54	0.42
1:AH:41:GLU:HA	1:AH:44:THR:CG2	2.47	0.42
1:AH:75:GLU:HG2	8:BG:194:ILE:HD11	2.01	0.42
1:AJ:46:SER:HB2	1:AJ:50:ILE:HD11	2.00	0.42
2:AL:33:SER:OG	2:AL:37:ARG:NH2	2.50	0.42
1:AN:105:GLU:HG3	1:AN:106:GLU:N	2.33	0.42
2:AO:96:MET:HA	2:AO:152:TYR:CE2	2.54	0.42
2:AO:104:LEU:HD23	2:AO:108:VAL:HG21	2.01	0.42
2:AQ:112:LEU:HG	2:AQ:160:LEU:CD1	2.49	0.42
1:AR:24:ASP:O	1:AR:27:LYS:HG2	2.20	0.42
1:AR:52:LYS:HD3	8:BG:221:MET:HG3	2.02	0.42
2:AS:11:SER:O	2:AS:14:VAL:HG12	2.19	0.42
2:AS:41:ALA:HB1	2:AS:97:LEU:HD11	2.01	0.42
2:AU:61:LEU:HD22	10:AV:200:CYC:CBA	2.49	0.42
2:AU:141:VAL:HG12	2:AU:145:ALA:HB3	2.02	0.42
1:AV:101:VAL:HB	1:AV:155:PHE:CE1	2.53	0.42
1:AX:75:GLU:OE2	8:BG:240:ILE:HG23	2.18	0.42
2:AY:112:LEU:HD11	10:AY:200:CYC:HMB3	2.01	0.42
5:BA:7:THR:O	5:BA:51:VAL:HG22	2.18	0.42
6:BD:287:ARG:HA	6:BD:292:ARG:O	2.19	0.42
6:BD:355:PRO:HD2	6:BD:430:PHE:CZ	2.54	0.42
2:BJ:75:THR:HG22	10:BK:200:CYC:HMB2	2.01	0.42
1:BK:73:TYR:H	1:BK:77:MET:CB	2.29	0.42
1:BK:124:GLU:HG2	1:BK:125:ALA:N	2.34	0.42
3:BM:53:GLN:O	3:BM:56:LYS:HG2	2.18	0.42
10:BN:200:CYC:HB	10:BN:200:CYC:CMA	2.32	0.42
1:BV:46:SER:OG	1:BV:140:LEU:HD23	2.19	0.42
1:BV:131:ARG:HG3	1:BV:157:ILE:HD13	2.00	0.42
2:BW:134:LYS:CA	2:BW:153:LEU:HD13	2.49	0.42
2:BW:134:LYS:CB	2:BW:153:LEU:HD13	2.48	0.42
2:BY:81:CYS:HA	10:BY:200:CYC:HHD	2.01	0.42
2:CA:2:GLN:HG3	2:CA:100:ASP:OD2	2.19	0.42
1:CC:24:ASP:HA	1:CC:27:LYS:NZ	2.33	0.42
2:CD:126:THR:O	2:CD:130:ILE:HG13	2.19	0.42
2:CD:131:GLN:O	2:CD:134:LYS:HB3	2.19	0.42
1:CE:103:PRO:O	1:CE:107:ILE:HD12	2.19	0.42
6:CM:344:GLU:CD	6:CM:430:PHE:HB2	2.44	0.42
10:CM:902:CYC:NB	10:CM:902:CYC:HMA1	2.34	0.42
1:CR:126:VAL:HG12	1:CR:160:MET:HE2	2.01	0.42

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:CS:40:ALA:HB2	2:CS:145:ALA:HB1	2.01	0.42
1:CT:98:SER:HA	2:CU:5:ILE:HG21	2.01	0.42
1:CY:57:ARG:CZ	1:CY:57:ARG:HB2	2.49	0.42
2:DB:28:LYS:HB2	2:DB:28:LYS:HE2	1.92	0.42
1:DC:17:TYR:OH	2:DD:89:LEU:HG	2.19	0.42
1:DC:102:THR:O	1:DC:106:GLU:HG2	2.19	0.42
8:DG:225:ILE:HG23	1:DQ:56:ASP:OD1	2.19	0.42
8:DH:227:PRO:HG3	1:DS:78:THR:HG21	2.00	0.42
9:DI:213:LEU:CD1	9:DI:216:LEU:HD22	2.49	0.42
1:DJ:122:PRO:HD2	10:DJ:200:CYC:OC	2.19	0.42
2:DO:62:TYR:N	2:DO:66:THR:HG21	2.35	0.42
2:DO:105:ASP:HA	2:DO:109:LEU:HD12	2.00	0.42
1:DU:39:ILE:HG23	1:DU:141:MET:CE	2.49	0.42
2:AB:129:ALA:O	2:AB:133:ILE:HG13	2.19	0.42
1:AC:71:ASN:O	1:AC:77:MET:HB3	2.20	0.42
1:AH:47:ARG:HA	1:AH:89:LEU:HD21	2.01	0.42
1:AJ:72:ALA:HA	1:AJ:77:MET:HB3	2.01	0.42
4:AK:110:LEU:HD13	4:AK:163:GLU:HG3	2.00	0.42
2:AQ:36:LEU:HD13	2:AQ:39:ARG:HH21	1.84	0.42
2:AS:112:LEU:HD21	2:AS:160:LEU:HD21	2.00	0.42
2:AY:24:MET:HE3	2:AY:28:LYS:HD3	2.01	0.42
5:BA:59:ARG:HB2	5:BA:62:THR:CG2	2.49	0.42
6:BD:192:ILE:O	6:BD:196:ILE:HG12	2.20	0.42
6:BD:562:SER:O	6:BD:566:ILE:HG23	2.19	0.42
6:BD:755:ILE:O	6:BD:755:ILE:HG13	2.19	0.42
1:BK:25:ARG:CG	6:CM:42:GLU:HG3	2.49	0.42
3:BM:58:LEU:HD13	3:BM:129:ALA:N	2.34	0.42
10:BN:200:CYC:HC	10:BN:200:CYC:CMD	2.32	0.42
2:BW:47:ASN:O	2:BW:51:ILE:HG13	2.20	0.42
2:BW:57:ALA:HA	2:BW:61:LEU:HD12	2.02	0.42
6:CM:85:LEU:HA	6:CM:153:LEU:CD1	2.48	0.42
6:CM:543:ILE:HG23	6:CM:578:PHE:CZ	2.54	0.42
2:CS:85:LEU:HD23	2:CS:88:TYR:HD2	1.83	0.42
1:CT:30:VAL:CG2	2:CU:31:PHE:HA	2.49	0.42
1:CT:144:ASP:OD1	1:CT:145:ASP:N	2.50	0.42
1:CY:47:ARG:O	1:CY:50:ILE:HG22	2.19	0.42
1:CY:81:CYS:HA	10:CY:200:CYC:CHD	2.49	0.42
2:CZ:62:TYR:CG	8:DG:203:PHE:HB2	2.55	0.42
2:CZ:76:ARG:NH2	10:CZ:200:CYC:O1D	2.46	0.42
1:DJ:101:VAL:CG1	1:DQ:20:PRO:HG2	2.49	0.42
1:DL:49:THR:O	1:DL:53:GLN:HG2	2.19	0.42

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:DM:14:VAL:HG23	2:DM:15:GLN:OE1	2.19	0.42
1:DN:121:THR:HG23	10:DN:200:CYC:C1C	2.49	0.42
1:DQ:33:GLY:HA3	2:DR:31:PHE:CE2	2.54	0.42
1:DQ:131:ARG:HH12	1:DQ:157:ILE:HD12	1.84	0.42
2:DR:126:THR:HG23	10:DR:200:CYC:CBC	2.48	0.42
2:DT:47:ASN:O	2:DT:51:ILE:HG13	2.19	0.42
2:DV:126:THR:CG2	10:DV:200:CYC:HMC1	2.49	0.42
9:EA:47:MET:HE3	9:EA:52:THR:O	2.19	0.42
1:AJ:71:ASN:C	1:AJ:71:ASN:HD22	2.27	0.42
4:AK:4:ALA:HB2	4:AK:100:ALA:O	2.20	0.42
2:AQ:77:ARG:HG2	10:AQ:200:CYC:HAD1	2.01	0.42
2:AQ:141:VAL:HG12	2:AQ:145:ALA:HB3	2.00	0.42
2:AS:39:ARG:HA	2:AS:39:ARG:HH11	1.85	0.42
1:AV:111:GLY:HA2	1:AV:114:GLU:OE1	2.19	0.42
2:AW:3:ASP:N	2:AW:6:THR:OG1	2.50	0.42
2:AW:126:THR:O	2:AW:130:ILE:HG13	2.20	0.42
1:AX:130:VAL:HG12	1:AX:157:ILE:HD11	2.01	0.42
9:BH:229:ARG:HD2	9:EA:315:PHE:CD2	2.54	0.42
1:BK:90:ARG:HD2	1:BK:94:TYR:OH	2.18	0.42
1:BK:141:MET:CE	1:BK:146:ALA:HA	2.48	0.42
3:BM:58:LEU:HD13	3:BM:129:ALA:HA	2.00	0.42
10:BN:200:CYC:HBA1	5:CJ:33:TYR:CE2	2.55	0.42
1:BR:27:LYS:NZ	4:BS:35:SER:HA	2.34	0.42
1:BR:130:VAL:HB	1:BR:157:ILE:HD13	2.01	0.42
1:BV:33:GLY:HA2	1:BV:36:ARG:CG	2.49	0.42
1:BV:127:ALA:HB1	1:BV:161:SER:HB3	2.02	0.42
2:CA:56:VAL:HG13	2:CA:60:LEU:HB2	2.00	0.42
1:CC:39:ILE:HD11	1:CC:145:ASP:OD1	2.18	0.42
1:CE:40:ALA:O	1:CE:44:THR:HG23	2.18	0.42
1:CE:126:VAL:CG1	1:CE:160:MET:HE3	2.50	0.42
1:CG:137:ALA:HB1	1:CG:141:MET:HE2	2.00	0.42
6:CM:866:PHE:HB3	6:CM:867:PRO:HD3	2.02	0.42
1:CR:90:ARG:HD2	2:CS:17:LYS:C	2.44	0.42
1:CR:90:ARG:HB2	2:CS:18:TYR:CZ	2.54	0.42
1:CT:27:LYS:HD2	2:CU:35:GLU:OE1	2.18	0.42
1:CT:58:LEU:HD22	1:CT:129:SER:HB3	2.00	0.42
1:CV:37:LEU:HD12	2:CW:28:LYS:HE3	2.01	0.42
1:CV:94:TYR:CG	2:CW:9:ILE:HG23	2.53	0.42
1:CY:39:ILE:CD1	1:CY:145:ASP:HA	2.49	0.42
2:CZ:43:VAL:CG1	2:CZ:141:VAL:HG12	2.50	0.42
1:DA:51:VAL:CG1	8:DZ:221:MET:HE1	2.48	0.42

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:DA:72:ALA:HA	1:DA:77:MET:CB	2.49	0.42
9:DI:131:LEU:HD23	9:DI:136:ASN:HA	2.01	0.42
9:DI:146:GLU:HB2	9:DI:149:GLN:CD	2.45	0.42
2:DK:81:CYS:HA	10:DK:200:CYC:HAC1	1.78	0.42
2:DR:71:ASN:O	2:DR:77:ARG:NE	2.50	0.42
2:DT:81:CYS:SG	10:DT:200:CYC:HMD3	2.59	0.42
1:DU:134:LYS:HB2	1:DU:153:PHE:HB3	2.00	0.42
9:EA:32:ASN:HB3	9:EA:171:ARG:NH2	2.35	0.42
9:EA:72:GLN:HG2	9:EA:116:LEU:HD13	2.01	0.42
9:EA:210:PHE:CB	9:EA:241:PHE:HB3	2.50	0.42
2:AB:52:VAL:O	2:AB:56:VAL:HG23	2.19	0.42
2:AB:54:GLU:O	2:AB:58:LYS:HG2	2.19	0.42
3:AE:156:ILE:O	3:AE:160:MET:HG2	2.19	0.42
2:AF:78:TYR:O	2:AF:82:ILE:HG12	2.19	0.42
2:AF:141:VAL:CG1	2:AF:145:ALA:HB3	2.49	0.42
1:AH:13:ALA:HB2	6:BD:330:ARG:CD	2.50	0.42
1:AJ:49:THR:HG21	2:BW:54:GLU:OE2	2.19	0.42
4:AK:47:ASN:O	4:AK:51:ILE:HG13	2.19	0.42
2:AL:36:LEU:HD11	7:BF:85:GLN:HB2	2.01	0.42
1:AP:122:PRO:O	1:AP:126:VAL:HG23	2.19	0.42
1:AR:101:VAL:HG11	1:AR:155:PHE:CZ	2.55	0.42
2:AS:105:ASP:O	2:AS:110:ASN:HB2	2.20	0.42
2:AS:123:ILE:CA	2:AS:126:THR:HG22	2.45	0.42
2:AU:105:ASP:OD2	2:AU:155:TYR:OH	2.26	0.42
1:AV:59:PHE:CE2	1:AV:78:THR:HG23	2.55	0.42
1:AZ:109:LEU:HA	1:AZ:112:VAL:HG21	2.02	0.42
6:BD:357:SER:O	6:BD:361:VAL:HG23	2.20	0.42
8:BG:197:GLN:OE1	8:BG:197:GLN:N	2.44	0.42
9:BH:289:ARG:CZ	9:EA:306:LEU:HD12	2.50	0.42
10:BL:200:CYC:HMB2	5:CJ:22:LEU:HD13	2.01	0.42
3:BM:129:ALA:O	3:BM:132:VAL:HG22	2.20	0.42
2:BN:77:ARG:HB3	10:BN:200:CYC:HMD1	2.01	0.42
2:BN:115:THR:HG23	5:CJ:53:VAL:HG21	2.02	0.42
1:BP:97:VAL:CG1	2:BQ:27:LEU:HD22	2.49	0.42
2:BQ:74:THR:HG23	2:BQ:77:ARG:HH12	1.85	0.42
2:BT:119:LEU:HD21	6:CM:424:PHE:HE1	1.84	0.42
2:BW:56:VAL:HG21	2:BW:82:ILE:HD11	2.01	0.42
1:BZ:64:ASP:HA	1:BZ:67:SER:CB	2.50	0.42
1:CE:6:LYS:O	1:CE:9:VAL:HG22	2.18	0.42
1:CE:39:ILE:CG1	1:CE:145:ASP:HB3	2.50	0.42
2:CF:57:ALA:CA	2:CF:61:LEU:HD12	2.34	0.42

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:CF:64:ASP:N	2:CF:64:ASP:OD1	2.50	0.42
2:CF:93:THR:HA	2:CF:149:MET:HE1	2.01	0.42
2:CF:99:GLY:HA2	2:CF:152:TYR:OH	2.19	0.42
1:CG:82:LEU:O	1:CG:85:MET:HB2	2.19	0.42
5:CJ:2:ARG:HH12	5:CJ:66:LEU:HD13	1.84	0.42
6:CM:455:GLU:OE2	6:CM:607:ARG:NH1	2.52	0.42
6:CM:524:VAL:O	6:CM:524:VAL:HG12	2.20	0.42
6:CM:809:LEU:HD23	6:CM:809:LEU:HA	1.85	0.42
6:CM:851:ASP:OD1	6:CM:851:ASP:N	2.52	0.42
2:CW:3:ASP:OD1	2:CW:6:THR:OG1	2.38	0.42
2:CW:133:ILE:O	2:CW:137:THR:HG22	2.20	0.42
2:CW:137:THR:O	2:CW:141:VAL:HG22	2.19	0.42
1:CY:17:TYR:CE2	2:CZ:90:ARG:HG2	2.55	0.42
1:CY:46:SER:O	1:CY:50:ILE:HG22	2.20	0.42
1:CY:76:GLU:HB2	8:DH:201:ARG:O	2.20	0.42
2:CZ:52:VAL:O	2:CZ:56:VAL:HG23	2.20	0.42
1:DA:51:VAL:HG22	1:DA:133:MET:CE	2.42	0.42
9:DI:27:ARG:NH1	9:DI:141:THR:HG21	2.34	0.42
9:DI:41:TRP:O	9:DI:45:LEU:HG	2.19	0.42
9:DI:107:LEU:HD13	9:DI:158:VAL:HG21	2.02	0.42
1:DJ:58:LEU:HD13	1:DJ:128:GLN:HB3	2.01	0.42
1:DL:4:VAL:O	1:DL:8:ILE:HG12	2.20	0.42
2:DM:112:LEU:HD11	2:DM:116:TYR:CE2	2.54	0.42
2:DO:59:SER:HB3	2:DO:132:ALA:HB2	2.00	0.42
1:DQ:100:ASP:O	1:DQ:103:PRO:HD2	2.19	0.42
1:DU:81:CYS:HB3	10:DU:200:CYC:HBC1	1.32	0.42
9:EA:214:ILE:CD1	9:EA:238:LEU:HB2	2.50	0.42
1:AA:101:VAL:HG13	1:AH:20:PRO:HG2	2.02	0.42
2:AB:88:TYR:CD2	2:AB:130:ILE:HD11	2.54	0.42
1:AH:88:TYR:O	1:AH:92:VAL:HG23	2.19	0.42
1:AH:109:LEU:HA	1:AH:112:VAL:HG21	2.01	0.42
2:AI:81:CYS:HA	10:AI:200:CYC:CHD	2.50	0.42
2:AI:160:LEU:HD23	2:AI:160:LEU:HA	1.92	0.42
4:AK:62:GLU:OE1	4:AK:133:ILE:HD11	2.20	0.42
2:AL:114:GLU:HA	6:BD:466:THR:O	2.18	0.42
1:AN:78:THR:O	1:AN:82:LEU:HG	2.19	0.42
1:AN:128:GLN:O	1:AN:132:GLU:HG2	2.20	0.42
1:AP:74:GLY:O	1:AP:78:THR:HG23	2.18	0.42
2:AQ:53:LYS:HD2	1:AR:118:SER:O	2.19	0.42
1:AX:85:MET:HE3	1:AX:133:MET:HE3	2.01	0.42
5:BA:6:ILE:HD13	5:BA:36:TRP:CZ3	2.54	0.42

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
6:BD:77:THR:O	6:BD:140:ASN:HA	2.19	0.42
6:BD:318:GLU:CD	6:BD:321:ARG:HE	2.27	0.42
9:BH:34:GLU:HG2	9:BH:174:GLU:HG2	2.02	0.42
9:BH:249:LYS:N	9:BH:275:THR:OG1	2.40	0.42
1:BI:12:ASP:OD1	2:BJ:91:TYR:OH	2.33	0.42
1:BK:113:ARG:HD3	1:BK:123:ILE:HG12	2.02	0.42
2:BN:106:GLU:O	2:BN:110:ASN:HB2	2.19	0.42
2:BQ:107:ARG:O	6:CM:338:ILE:HD12	2.18	0.42
2:BT:64:ASP:OD1	2:BT:64:ASP:N	2.52	0.42
2:BW:41:ALA:HB2	2:BW:96:MET:CE	2.49	0.42
6:CM:314:ILE:HG21	6:CM:322:ARG:HH11	1.85	0.42
1:CR:14:GLU:HB3	1:CR:16:ARG:HG3	2.02	0.42
1:CR:108:GLY:HA2	2:CW:75:THR:CG2	2.49	0.42
1:CR:144:ASP:OD1	1:CR:145:ASP:N	2.52	0.42
1:CT:77:MET:HE2	1:CT:77:MET:N	2.34	0.42
2:CU:103:ILE:HD11	2:CU:107:ARG:HG3	2.01	0.42
2:CW:67:ARG:O	2:CW:73:TYR:HB2	2.19	0.42
1:DA:140:LEU:HD12	1:DA:140:LEU:O	2.19	0.42
2:DB:36:LEU:HD21	2:DB:145:ALA:CA	2.49	0.42
2:DB:68:PRO:HA	2:DB:73:TYR:CG	2.55	0.42
1:DC:115:MET:HE1	10:DC:200:CYC:HMB3	2.00	0.42
8:DH:197:GLN:HG3	8:DH:199:GLU:OE2	2.19	0.42
2:DK:36:LEU:HA	2:DK:39:ARG:CG	2.47	0.42
2:DK:75:THR:HG23	1:DL:108:GLY:HA2	2.01	0.42
2:DM:76:ARG:HG3	1:DN:110:VAL:CG1	2.48	0.42
2:DR:35:GLU:CA	2:DR:38:VAL:HG12	2.48	0.42
1:DS:74:GLY:O	1:DS:78:THR:HG23	2.19	0.42
9:EA:41:TRP:NE1	9:EA:125:ILE:HD13	2.35	0.42
9:EA:154:LEU:O	9:EA:158:VAL:HG22	2.20	0.42
9:EA:222:ALA:O	9:EA:300:PHE:HA	2.20	0.42
4:AK:8:LEU:HD23	4:AK:27:LEU:HD21	2.02	0.42
4:AK:93:ALA:O	4:AK:97:LEU:HG	2.19	0.42
2:AS:130:ILE:HD13	2:AS:156:ILE:CG2	2.50	0.42
2:AU:65:VAL:CG1	2:AU:72:MET:HE2	2.44	0.42
1:AZ:62:ARG:HH12	1:AZ:124:GLU:HG3	1.85	0.42
5:BA:29:LYS:NZ	6:BD:318:GLU:H	2.08	0.42
6:BD:845:LEU:HG	5:DF:42:ARG:NH1	2.35	0.42
9:BH:51:LEU:HD23	9:BH:150:GLN:NE2	2.35	0.42
3:BM:15:LEU:HA	2:BN:90:ARG:NH1	2.35	0.42
2:BN:105:ASP:HB2	2:BN:155:TYR:OH	2.19	0.42
1:BR:62:ARG:O	1:BR:65:ILE:HG12	2.19	0.42

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:BR:90:ARG:O	1:BR:93:THR:OG1	2.27	0.42
4:BS:2:ARG:CB	4:BS:103:ASN:HD22	2.31	0.42
4:BS:66:LEU:HD11	4:BS:126:PRO:CB	2.49	0.42
4:BS:110:LEU:HA	4:BS:113:LEU:HD12	2.02	0.42
2:BT:134:LYS:CB	2:BT:153:LEU:HD13	2.47	0.42
1:BV:91:LEU:CD2	1:BV:107:ILE:HB	2.49	0.42
1:CC:50:ILE:HA	1:CC:136:VAL:HG11	2.01	0.42
1:CC:57:ARG:HD2	1:CC:132:GLU:OE1	2.20	0.42
2:CD:57:ALA:HA	2:CD:61:LEU:HD12	2.01	0.42
1:CE:54:ALA:HB1	1:CE:133:MET:HG2	2.00	0.42
2:CF:2:GLN:OE1	2:CF:7:ALA:HA	2.20	0.42
6:CM:144:TYR:HB3	6:CM:149:MET:HG2	2.01	0.42
6:CM:285:ALA:CB	6:CM:316:MET:HE1	2.49	0.42
6:CM:410:ASN:HA	6:CM:413:MET:CE	2.50	0.42
6:CM:795:TYR:CE1	6:CM:881:LYS:HE3	2.54	0.42
1:CR:134:LYS:HG3	1:CR:153:PHE:HD2	1.84	0.42
2:CS:83:ARG:HD2	2:CS:87:TYR:OH	2.19	0.42
1:CV:47:ARG:O	1:CV:51:VAL:HG23	2.19	0.42
2:CW:36:LEU:CD2	2:CW:145:ALA:HB2	2.50	0.42
1:CY:35:ALA:O	1:CY:39:ILE:HG13	2.19	0.42
1:CY:39:ILE:HG12	1:CY:145:ASP:CG	2.44	0.42
1:DA:62:ARG:HD3	9:DI:58:ALA:N	2.35	0.42
1:DC:85:MET:C	1:DC:133:MET:HE1	2.44	0.42
8:DH:221:MET:HE3	1:DS:52:LYS:CA	2.50	0.42
9:DI:191:GLU:H	9:DI:254:ARG:HA	1.83	0.42
2:DM:3:ASP:HB2	2:DM:98:ALA:O	2.20	0.42
1:DQ:50:ILE:CG2	1:DQ:89:LEU:HD11	2.49	0.42
1:DQ:61:LYS:HE2	1:DQ:61:LYS:HA	2.01	0.42
2:DR:90:ARG:HG2	2:DR:94:TYR:CE2	2.55	0.42
2:DR:111:GLY:O	2:DR:115:THR:HG22	2.19	0.42
2:DT:56:VAL:HG12	2:DT:61:LEU:HG	2.01	0.42
9:EA:44:TYR:O	9:EA:47:MET:HB2	2.20	0.42
9:EA:107:LEU:HG	11:EA:400:45D:C26	2.50	0.42
2:AB:63:SER:O	2:AB:67:ARG:HD3	2.19	0.42
2:AB:74:THR:HB	1:AC:107:ILE:O	2.19	0.42
2:AD:81:CYS:HA	10:AD:200:CYC:HAC1	1.90	0.42
2:AD:105:ASP:OD2	2:AD:155:TYR:OH	2.37	0.42
3:AE:113:LYS:NZ	3:AE:123:VAL:HG11	2.35	0.42
2:AF:93:THR:HA	2:AF:149:MET:HE1	2.02	0.42
2:AF:110:ASN:HB3	5:BA:44:MET:HE3	2.00	0.42
1:AH:108:GLY:HA2	2:AL:75:THR:OG1	2.19	0.42

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:AJ:4:VAL:HG12	1:AJ:8:ILE:CD1	2.50	0.42
1:AJ:18:LEU:HD11	4:AK:95:TYR:HD1	1.85	0.42
4:AK:129:ARG:NH2	4:AK:133:ILE:HG13	2.26	0.42
1:AP:142:SER:H	2:DT:39:ARG:NH2	2.18	0.42
2:AW:104:LEU:O	2:AW:108:VAL:HB	2.20	0.42
10:AW:200:CYC:OB	6:BD:566:ILE:HD11	2.20	0.42
1:AX:76:GLU:HB3	8:BG:247:PRO:HG3	2.01	0.42
2:AY:100:ASP:OD1	2:AY:101:ALA:N	2.51	0.42
2:AY:104:LEU:HD22	2:AY:156:ILE:HD11	2.02	0.42
6:BD:31:GLN:CB	6:BD:33:ARG:HE	2.33	0.42
6:BD:268:ALA:HB3	6:BD:270:LYS:HZ2	1.85	0.42
6:BD:808:PHE:HB2	6:BD:838:MET:SD	2.60	0.42
10:BD:901:CYC:HB	10:BD:901:CYC:CMA	2.33	0.42
8:BG:225:ILE:O	8:BG:227:PRO:HD3	2.20	0.42
9:BH:112:ARG:HD2	9:BH:112:ARG:HA	1.84	0.42
9:BH:190:ILE:HG23	9:BH:193:VAL:HB	2.01	0.42
1:BI:50:ILE:HD11	1:BI:137:ALA:N	2.34	0.42
2:BL:51:ILE:HA	2:BL:136:VAL:HG11	2.02	0.42
2:BL:134:LYS:NZ	2:BL:154:ASP:OD1	2.50	0.42
3:BM:101:LYS:HE3	1:BR:16:ARG:NE	2.34	0.42
4:BS:62:GLU:CD	4:BS:133:ILE:HD11	2.45	0.42
4:BS:124:ILE:HD11	4:BS:164:LEU:CD1	2.50	0.42
1:BV:25:ARG:NE	1:BV:25:ARG:HA	2.34	0.42
1:BV:44:THR:O	1:BV:47:ARG:HG2	2.20	0.42
2:BY:14:VAL:CG1	6:CM:683:VAL:HG11	2.44	0.42
1:CC:134:LYS:HD3	1:CC:153:PHE:CB	2.47	0.42
2:CD:52:VAL:HG11	2:CD:82:ILE:HG13	2.02	0.42
1:CE:48:GLU:HG2	1:CE:49:THR:N	2.35	0.42
2:CH:21:GLY:O	2:CH:24:MET:HB3	2.19	0.42
9:CQ:39:LEU:HD12	9:CQ:139:LEU:HD12	2.00	0.42
9:CQ:89:ARG:HD3	9:CQ:170:LYS:HE3	2.01	0.42
9:CQ:115:GLU:CG	9:CQ:119:GLN:HE22	2.33	0.42
9:CQ:131:LEU:HD23	9:CQ:136:ASN:HA	2.01	0.42
9:CQ:311:GLU:O	9:CQ:312:LEU:HB2	2.18	0.42
1:CR:101:VAL:HB	1:CR:155:PHE:CE2	2.54	0.42
1:DA:54:ALA:HB1	1:DA:133:MET:HG2	2.01	0.42
1:DA:90:ARG:NH2	2:DB:13:ASP:OD1	2.52	0.42
10:DA:200:CYC:O1D	8:DG:201:ARG:NH2	2.53	0.42
2:DB:111:GLY:O	2:DB:114:GLU:HG2	2.19	0.42
8:DG:238:GLN:CD	1:DL:52:LYS:HG2	2.45	0.42
9:DI:86:LEU:CD2	9:DI:95:CYS:HA	2.50	0.42

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
9:DI:314:ASN:C	9:DI:314:ASN:ND2	2.76	0.42
1:DN:92:VAL:O	1:DN:96:VAL:HG23	2.20	0.42
1:DQ:140:LEU:HD23	1:DQ:140:LEU:O	2.20	0.42
1:DU:88:TYR:O	1:DU:92:VAL:HG23	2.19	0.42
2:DV:53:LYS:O	2:DV:56:VAL:HG12	2.20	0.42
9:EA:39:LEU:HD12	9:EA:139:LEU:HD12	2.02	0.42
9:EA:148:GLY:HA2	9:EA:151:ILE:HG22	2.02	0.42
1:AC:39:ILE:HD11	1:AC:145:ASP:OD1	2.20	0.42
2:AD:129:ALA:O	2:AD:133:ILE:HG13	2.20	0.42
3:AE:8:ILE:HG13	2:AF:1:MET:HE2	2.01	0.42
3:AE:15:LEU:HA	2:AF:90:ARG:NH1	2.35	0.42
1:AH:115:MET:HE1	10:AH:200:CYC:HMB2	2.01	0.42
4:AK:82:CYS:HA	10:AK:200:CYC:CHD	2.49	0.42
1:AR:87:TYR:CD2	10:AR:200:CYC:HBB3	2.55	0.42
2:BJ:59:SER:HB3	2:BJ:132:ALA:CB	2.50	0.42
3:BM:8:ILE:HG13	2:BN:1:MET:CE	2.50	0.42
3:BM:26:ILE:HD13	2:BN:97:LEU:HD21	2.02	0.42
3:BM:105:GLU:HG3	3:BM:110:ILE:HD11	2.01	0.42
4:BS:48:SER:OG	4:BS:49:ALA:N	2.53	0.42
1:BV:130:VAL:HA	1:BV:133:MET:HE2	2.01	0.42
2:BW:89:LEU:HD12	2:BW:89:LEU:HA	1.94	0.42
1:BX:14:GLU:HA	6:CM:548:ARG:NH2	2.35	0.42
1:BZ:25:ARG:NH2	1:CE:21:GLY:HA3	2.35	0.42
2:CA:144:ASP:OD1	2:CA:145:ALA:N	2.53	0.42
2:CD:61:LEU:HD22	10:CE:200:CYC:CAA	2.50	0.42
2:CD:72:MET:HG2	2:CD:78:TYR:CD2	2.54	0.42
2:CD:88:TYR:CE1	10:CD:200:CYC:HMB1	2.54	0.42
2:CD:109:LEU:HD22	2:CD:159:GLY:HA3	2.02	0.42
1:CE:14:GLU:CB	1:CE:16:ARG:HG2	2.50	0.42
1:CE:30:VAL:HG22	2:CF:31:PHE:O	2.19	0.42
1:CG:134:LYS:HG3	1:CG:153:PHE:CD2	2.55	0.42
2:CH:41:ALA:CA	2:CH:96:MET:HE1	2.50	0.42
2:CH:100:ASP:OD1	2:CH:101:ALA:N	2.52	0.42
6:CM:152:SER:HA	10:CM:902:CYC:C4C	2.50	0.42
9:CQ:34:GLU:CG	9:CQ:174:GLU:HG2	2.49	0.42
1:CR:126:VAL:C	1:CR:160:MET:HE1	2.44	0.42
2:CS:71:ASN:HB3	2:CS:122:PRO:HD3	2.01	0.42
2:CS:75:THR:CG2	1:CT:108:GLY:HA2	2.50	0.42
1:DA:53:GLN:HE21	1:DA:136:VAL:CG2	2.33	0.42
1:DA:76:GLU:CD	8:DG:202:ARG:HA	2.44	0.42
1:DC:128:GLN:O	1:DC:132:GLU:HG2	2.20	0.42

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:DD:88:TYR:OH	2:DD:112:LEU:HD21	2.20	0.42
2:DK:50:THR:HG22	2:DK:54:GLU:OE1	2.20	0.42
2:DK:71:ASN:O	2:DK:77:ARG:HD3	2.20	0.42
1:DL:39:ILE:CD1	1:DL:145:ASP:HB3	2.50	0.42
1:DL:104:ILE:HG22	1:DL:109:LEU:HD12	2.01	0.42
1:DU:109:LEU:HD13	1:DU:159:LYS:CG	2.45	0.42
8:DY:210:LEU:HD13	8:DY:216:MET:C	2.45	0.42
2:AB:40:ALA:HA	2:AB:141:VAL:CG2	2.49	0.42
2:AB:106:GLU:HG3	2:AB:107:ARG:CG	2.49	0.42
1:AC:152:TYR:O	1:AC:156:VAL:HG23	2.19	0.42
10:AD:200:CYC:HC	10:AD:200:CYC:CMD	2.33	0.42
3:AE:128:ASP:O	3:AE:131:THR:OG1	2.30	0.42
2:AF:100:ASP:OD1	2:AF:101:ALA:N	2.52	0.42
2:AF:111:GLY:HA3	5:BA:50:ILE:CD1	2.50	0.42
1:AN:25:ARG:NE	1:AN:25:ARG:HA	2.35	0.42
2:AQ:132:ALA:O	2:AQ:136:VAL:HG23	2.19	0.42
2:AS:148:GLU:HG3	2:AS:152:TYR:HE2	1.85	0.42
2:AY:96:MET:HA	2:AY:152:TYR:HE2	1.85	0.42
6:BD:152:SER:O	6:BD:156:MET:HG3	2.20	0.42
6:BD:314:ILE:HG21	6:BD:322:ARG:NH1	2.35	0.42
6:BD:458:PHE:HB3	6:BD:552:GLY:CA	2.50	0.42
6:BD:806:LYS:HD2	6:BD:872:LEU:HB3	2.02	0.42
9:BH:216:LEU:O	9:BH:298:ILE:N	2.45	0.42
2:BJ:132:ALA:O	2:BJ:136:VAL:HG23	2.19	0.42
1:BK:121:THR:HG23	10:BK:200:CYC:CMC	2.50	0.42
10:BL:200:CYC:CGA	10:BL:200:CYC:HMA3	2.50	0.42
3:BM:8:ILE:HG13	2:BN:1:MET:HE2	2.02	0.42
3:BM:50:ILE:CG1	3:BM:140:LEU:HD13	2.48	0.42
3:BM:126:MET:O	3:BM:130:VAL:HG23	2.19	0.42
2:BN:1:MET:CE	2:BN:103:ILE:HD12	2.50	0.42
2:BN:62:TYR:H	2:BN:66:THR:HG21	1.85	0.42
1:BP:153:PHE:O	1:BP:157:ILE:HG13	2.20	0.42
2:BQ:63:SER:HB3	6:CM:682:ARG:HB2	2.02	0.42
1:BR:5:THR:O	1:BR:8:ILE:HG22	2.20	0.42
1:BV:36:ARG:CZ	1:BV:36:ARG:HB3	2.50	0.42
2:BW:74:THR:HG23	1:BX:107:ILE:HA	2.01	0.42
2:BW:92:ALA:CB	2:BW:149:MET:HE3	2.50	0.42
2:CD:41:ALA:CA	2:CD:96:MET:HE1	2.50	0.42
2:CF:106:GLU:OE1	2:CF:107:ARG:HG3	2.20	0.42
1:CG:50:ILE:HA	1:CG:136:VAL:HG11	2.00	0.42
2:CH:104:LEU:HD13	2:CH:156:ILE:HG13	2.02	0.42

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
5:CJ:40:GLN:HG2	5:CJ:44:MET:HE3	2.02	0.42
6:CM:133:PRO:HB3	6:CM:136:PHE:CD1	2.55	0.42
6:CM:806:LYS:HB2	6:CM:872:LEU:HD13	2.01	0.42
9:CQ:55:ALA:N	11:CQ:400:45D:H221	2.35	0.42
9:CQ:74:MET:CE	9:CQ:78:GLN:HB2	2.50	0.42
9:CQ:277:TRP:CD1	9:CQ:305:LEU:HD11	2.55	0.42
2:CS:51:ILE:HD11	2:CS:140:LEU:HD12	2.02	0.42
2:CS:92:ALA:HB3	2:CS:149:MET:HE3	2.01	0.42
1:CT:56:ASP:O	1:CT:60:GLN:HG2	2.19	0.42
2:CU:47:ASN:O	2:CU:51:ILE:HG13	2.20	0.42
1:CV:97:VAL:HG11	2:CW:19:LEU:HD12	2.01	0.42
2:CW:153:LEU:HD23	2:CW:156:ILE:HD12	2.01	0.42
1:DA:83:ARG:HD3	8:DZ:213:GLY:O	2.20	0.42
2:DB:144:ASP:O	2:DB:147:LYS:HB3	2.19	0.42
2:DD:81:CYS:HB2	10:DD:200:CYC:HMD3	2.02	0.42
5:DF:5:ARG:HB3	5:DF:30:LEU:CD2	2.49	0.42
8:DG:248:ARG:HH12	2:DK:67:ARG:HD3	1.83	0.42
9:DI:126:PRO:HG3	11:DI:400:45D:H092	2.01	0.42
9:DI:126:PRO:HG3	11:DI:400:45D:C17	2.50	0.42
9:DI:146:GLU:C	9:DI:150:GLN:HE21	2.27	0.42
9:DI:269:VAL:HG23	9:DI:288:TRP:CE3	2.46	0.42
1:DJ:55:GLY:HA2	1:DJ:85:MET:HE1	2.01	0.42
2:DK:113:LYS:HE2	2:DK:161:SER:C	2.45	0.42
1:DL:130:VAL:HB	1:DL:157:ILE:CD1	2.50	0.42
1:DQ:39:ILE:CD1	1:DQ:145:ASP:HB3	2.49	0.42
1:DS:68:PRO:HG3	1:DS:73:TYR:CE1	2.55	0.42
2:DV:60:LEU:HD13	2:DV:72:MET:HE1	2.00	0.42
2:DV:119:LEU:HD11	10:DV:200:CYC:CAA	2.50	0.42
1:AA:108:GLY:HA2	2:AF:75:THR:OG1	2.20	0.41
2:AD:54:GLU:O	2:AD:58:LYS:HE3	2.20	0.41
3:AE:22:GLU:O	3:AE:26:ILE:HG13	2.20	0.41
3:AE:40:ALA:CB	3:AE:97:LEU:HD11	2.49	0.41
4:AK:115:GLU:OE2	6:BD:6:SER:OG	2.21	0.41
2:AL:51:ILE:HD13	2:AL:137:THR:HG22	2.01	0.41
1:AP:95:GLY:HA3	1:AP:104:ILE:CD1	2.50	0.41
1:AP:128:GLN:OE1	1:AP:131:ARG:NH1	2.53	0.41
2:AU:23:ALA:O	2:AU:27:LEU:HG	2.19	0.41
1:AV:27:LYS:HD2	2:AW:35:GLU:CD	2.45	0.41
6:BD:317:LYS:HD3	6:BD:392:GLU:OE2	2.20	0.41
6:BD:399:ARG:H	6:BD:399:ARG:HG2	1.65	0.41
6:BD:586:GLU:OE1	6:BD:586:GLU:N	2.53	0.41

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
6:BD:813:PRO:HD2	6:BD:885:VAL:HG11	2.02	0.41
8:BG:241:ASN:O	8:BG:245:SER:OG	2.31	0.41
1:BI:2:SER:OG	1:BI:5:THR:HG22	2.20	0.41
1:BK:46:SER:O	1:BK:50:ILE:HG12	2.19	0.41
1:BK:134:LYS:NZ	1:BK:154:ASP:HB3	2.35	0.41
2:BL:62:TYR:CE2	3:BM:80:GLN:HG3	2.48	0.41
2:BW:101:ALA:CB	2:BW:104:LEU:HD12	2.48	0.41
2:BY:53:LYS:O	2:BY:56:VAL:HG22	2.19	0.41
2:BY:66:THR:HA	2:BY:72:MET:O	2.20	0.41
1:BZ:76:GLU:OE2	9:CQ:155:ARG:NH2	2.39	0.41
1:CC:14:GLU:OE1	1:CC:16:ARG:NH2	2.51	0.41
1:CE:94:TYR:O	1:CE:97:VAL:HG22	2.19	0.41
2:CF:21:GLY:C	2:CF:24:MET:HB3	2.45	0.41
2:CH:104:LEU:HD22	2:CH:156:ILE:HD11	2.02	0.41
6:CM:155:ASP:OD2	10:CM:902:CYC:ND	2.53	0.41
6:CM:536:GLU:O	6:CM:539:THR:HG22	2.20	0.41
6:CM:703:GLU:CA	6:CM:706:ILE:HD12	2.42	0.41
1:CR:3:ILE:HD11	1:CR:29:PHE:CB	2.49	0.41
1:CR:27:LYS:HZ2	2:CS:38:VAL:HB	1.85	0.41
2:CS:106:GLU:OE2	5:DF:57:THR:HG22	2.19	0.41
1:DA:51:VAL:HB	8:DZ:221:MET:HE1	2.00	0.41
1:DA:62:ARG:HD3	9:DI:58:ALA:HB2	2.02	0.41
2:DD:100:ASP:OD1	2:DD:101:ALA:N	2.53	0.41
8:DG:213:GLY:HA2	1:DQ:87:TYR:OH	2.20	0.41
8:DG:218:PHE:CE1	1:DQ:83:ARG:HD2	2.55	0.41
9:DI:244:GLU:OE1	9:DI:244:GLU:N	2.51	0.41
2:DK:133:ILE:O	2:DK:137:THR:HG23	2.20	0.41
1:DN:44:THR:O	1:DN:47:ARG:HG2	2.20	0.41
1:DN:131:ARG:HG2	1:DN:157:ILE:CD1	2.42	0.41
10:DN:200:CYC:CMA	10:DN:200:CYC:HB	2.33	0.41
2:DO:3:ASP:OD1	2:DO:6:THR:OG1	2.11	0.41
2:DO:93:THR:N	2:DO:149:MET:HE1	2.34	0.41
2:DO:100:ASP:OD1	2:DO:101:ALA:N	2.52	0.41
1:DQ:53:GLN:HG2	1:DQ:57:ARG:HH12	1.85	0.41
2:DR:85:LEU:HD12	2:DR:133:ILE:CD1	2.50	0.41
1:DS:18:LEU:HD22	2:DT:97:LEU:HD23	2.02	0.41
1:DS:109:LEU:HD12	1:DS:159:LYS:CB	2.49	0.41
2:DT:44:ILE:HD11	2:DT:149:MET:HG3	2.02	0.41
1:DU:41:GLU:CG	2:DV:24:MET:HE2	2.50	0.41
2:DV:63:SER:O	2:DV:66:THR:HG22	2.19	0.41
9:EA:81:GLN:HA	9:EA:176:VAL:HG21	2.02	0.41

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:AA:77:MET:SD	10:AA:200:CYC:HAD1	2.60	0.41
1:AA:109:LEU:HD11	1:AA:159:LYS:HB3	2.01	0.41
1:AC:39:ILE:CD1	1:AC:141:MET:HE1	2.46	0.41
2:AD:106:GLU:HG3	2:AD:107:ARG:CD	2.49	0.41
2:AI:81:CYS:HA	10:AI:200:CYC:HHD	2.02	0.41
4:AK:14:LEU:HD21	2:AO:157:CYS:O	2.20	0.41
2:AU:68:PRO:HA	2:AU:73:TYR:CG	2.54	0.41
2:AU:77:ARG:NH2	5:BB:63:ASN:OD1	2.53	0.41
2:AY:24:MET:HE3	2:AY:28:LYS:CD	2.50	0.41
2:AY:71:ASN:HD22	2:AY:121:VAL:HA	1.83	0.41
2:AY:116:TYR:CD2	2:AY:160:LEU:HD11	2.54	0.41
5:BB:6:ILE:O	5:BB:6:ILE:HG13	2.20	0.41
6:BD:267:TYR:HB3	6:BD:281:VAL:HG22	2.02	0.41
6:BD:576:ARG:NH2	6:BD:641:ASP:HA	2.35	0.41
6:BD:576:ARG:NH1	6:BD:646:SER:OG	2.45	0.41
9:BH:141:THR:O	9:BH:145:LEU:HG	2.20	0.41
1:BK:131:ARG:HE	1:BK:157:ILE:CG2	2.34	0.41
4:BS:14:LEU:HD13	2:BW:161:SER:CB	2.51	0.41
4:BS:41:ALA:HA	4:BS:44:ILE:HD11	2.02	0.41
1:BV:68:PRO:O	1:CE:63:PRO:HG2	2.20	0.41
1:BV:108:GLY:HA2	2:CA:75:THR:HG21	2.01	0.41
2:BW:66:THR:HG21	10:BX:200:CYC:O1A	2.20	0.41
1:CC:61:LYS:HG3	1:CC:62:ARG:HG2	2.02	0.41
1:CE:132:GLU:O	1:CE:136:VAL:HG23	2.20	0.41
2:CF:36:LEU:CD2	2:CF:39:ARG:HH12	2.33	0.41
5:CK:23:GLN:HG2	5:CK:24:ASN:N	2.34	0.41
6:CM:802:GLU:OE2	6:CM:885:VAL:N	2.43	0.41
6:CM:811:ARG:HB2	6:CM:869:THR:OG1	2.19	0.41
9:CQ:42:PHE:CE2	9:CQ:131:LEU:HB2	2.55	0.41
9:CQ:87:ALA:O	9:CQ:170:LYS:NZ	2.42	0.41
1:CR:62:ARG:NE	1:DA:69:GLY:O	2.53	0.41
2:CW:15:GLN:HG3	2:CW:17:LYS:HZ3	1.83	0.41
2:CW:108:VAL:O	2:CW:112:LEU:HD13	2.19	0.41
1:CY:87:TYR:OH	8:DY:213:GLY:HA3	2.19	0.41
1:CY:97:VAL:CG1	2:CZ:27:LEU:HD13	2.51	0.41
2:CZ:64:ASP:O	2:CZ:70:GLY:HA3	2.20	0.41
1:DA:8:ILE:HD12	2:DB:94:TYR:CD1	2.55	0.41
1:DA:143:SER:O	1:DA:146:ALA:HB3	2.20	0.41
2:DB:132:ALA:O	2:DB:136:VAL:HG23	2.20	0.41
1:DC:66:VAL:O	1:DC:73:TYR:HA	2.20	0.41
9:DI:20:VAL:HB	9:DI:145:LEU:CD2	2.50	0.41

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:DL:92:VAL:HG11	1:DL:153:PHE:CE2	2.55	0.41
2:DM:75:THR:HG23	1:DN:108:GLY:HA2	2.01	0.41
1:DN:50:ILE:HA	1:DN:136:VAL:HG11	2.02	0.41
1:DQ:6:LYS:HD2	1:DQ:100:ASP:CG	2.46	0.41
1:DS:37:LEU:HD12	2:DT:24:MET:HE3	2.01	0.41
1:DS:39:ILE:HG23	1:DS:141:MET:CE	2.50	0.41
2:DV:3:ASP:N	2:DV:6:THR:OG1	2.52	0.41
9:EA:37:LEU:HA	9:EA:40:ILE:HG12	2.01	0.41
9:EA:188:VAL:O	9:EA:199:LEU:HD22	2.19	0.41
1:AA:37:LEU:HD13	2:AB:28:LYS:NZ	2.35	0.41
1:AA:46:SER:O	1:AA:50:ILE:HG22	2.20	0.41
1:AC:6:LYS:HD2	1:AC:6:LYS:HA	1.86	0.41
3:AE:64:GLU:O	3:AE:70:GLY:HA3	2.20	0.41
2:AF:116:TYR:CD1	2:AF:121:VAL:HG21	2.54	0.41
2:AF:123:ILE:O	2:AF:126:THR:OG1	2.34	0.41
1:AH:90:ARG:O	1:AH:93:THR:OG1	2.30	0.41
1:AH:150:SER:HA	1:AH:153:PHE:HB2	2.02	0.41
2:AI:91:TYR:HB3	2:AI:104:LEU:CD2	2.49	0.41
1:AJ:3:ILE:HG13	1:AJ:25:ARG:NH2	2.35	0.41
4:AK:75:THR:HB	6:BD:178:ASN:ND2	2.34	0.41
4:AK:161:THR:O	4:AK:165:SER:HB2	2.19	0.41
4:AK:167:LEU:CD2	6:BD:477:ASN:HB2	2.50	0.41
2:AL:5:ILE:HD13	6:BD:46:TYR:OH	2.21	0.41
1:AP:126:VAL:HG22	10:AP:200:CYC:HMC1	2.02	0.41
2:AQ:68:PRO:HD2	8:BG:213:GLY:HA3	2.02	0.41
2:AU:73:TYR:HH	1:AV:90:ARG:HH22	1.64	0.41
1:AV:91:LEU:HD21	1:AV:107:ILE:HB	2.02	0.41
1:AX:77:MET:HE2	1:AX:77:MET:N	2.35	0.41
1:AZ:81:CYS:HA	10:AZ:200:CYC:CHD	2.45	0.41
6:BD:606:HIS:HB2	6:BD:639:MET:HE1	2.01	0.41
9:BH:283:GLY:O	9:BH:284:MET:HE2	2.19	0.41
9:BH:284:MET:SD	9:BH:307:ALA:HA	2.60	0.41
2:BJ:112:LEU:HG	2:BJ:160:LEU:HD21	2.02	0.41
2:BL:5:ILE:O	2:BL:8:VAL:HG12	2.21	0.41
3:BM:150:ALA:N	3:BM:151:PRO:HD2	2.36	0.41
2:BN:123:ILE:HD13	2:BN:123:ILE:HA	1.90	0.41
1:BP:112:VAL:HA	2:BT:75:THR:HG21	2.02	0.41
1:BR:134:LYS:HD3	1:BR:153:PHE:HB3	2.02	0.41
4:BS:125:GLY:O	4:BS:128:VAL:HG22	2.19	0.41
2:BY:2:GLN:O	2:BY:100:ASP:HB3	2.21	0.41
2:BY:71:ASN:ND2	2:BY:121:VAL:HA	2.35	0.41

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:BZ:71:ASN:C	1:BZ:71:ASN:HD22	2.27	0.41
1:BZ:77:MET:SD	10:BZ:200:CYC:HAD1	2.60	0.41
2:CA:39:ARG:NH1	2:CA:42:SER:HB3	2.35	0.41
2:CD:119:LEU:HD21	5:CK:33:TYR:CE2	2.45	0.41
2:CF:1:MET:HB3	2:CF:103:ILE:HD13	2.01	0.41
2:CH:81:CYS:HA	10:CH:200:CYC:CHD	2.48	0.41
5:CK:2:ARG:HH21	5:CK:66:LEU:HD13	1.85	0.41
6:CM:341:ARG:HD2	6:CM:341:ARG:HA	1.90	0.41
1:CT:137:ALA:HB1	1:CT:141:MET:HE3	2.01	0.41
1:CY:102:THR:HB	1:CY:103:PRO:HD3	2.02	0.41
8:DH:245:SER:HB2	1:DJ:79:ALA:HB2	2.03	0.41
9:DI:74:MET:HE2	9:DI:74:MET:HB3	1.91	0.41
9:DI:213:LEU:HA	9:DI:216:LEU:CB	2.49	0.41
1:DJ:57:ARG:O	1:DJ:61:LYS:HD3	2.20	0.41
1:DJ:62:ARG:O	1:DJ:65:ILE:HG12	2.21	0.41
2:DK:131:GLN:O	2:DK:134:LYS:HB3	2.21	0.41
1:DL:143:SER:O	1:DL:146:ALA:HB3	2.19	0.41
2:DM:96:MET:HG3	2:DM:149:MET:HG2	2.02	0.41
2:DT:127:VAL:O	2:DT:131:GLN:HG2	2.20	0.41
5:DX:6:ILE:HD13	5:DX:36:TRP:HZ3	1.85	0.41
9:EA:109:PHE:CZ	9:EA:113:LEU:HD11	2.56	0.41
1:AA:90:ARG:HH22	2:AF:73:TYR:HH	1.67	0.41
1:AA:116:TYR:CE2	1:AA:126:VAL:HG11	2.56	0.41
1:AC:97:VAL:HG23	2:AD:9:ILE:HD11	2.03	0.41
2:AD:112:LEU:HD21	10:AD:200:CYC:C3B	2.50	0.41
2:AF:107:ARG:HA	5:BA:44:MET:HB3	2.02	0.41
1:AH:14:GLU:OE1	1:AH:16:ARG:NE	2.50	0.41
1:AH:112:VAL:HA	2:AL:75:THR:HG21	2.02	0.41
1:AJ:18:LEU:HD13	4:AK:98:ILE:HD12	2.02	0.41
4:AK:129:ARG:NH1	2:AO:15:GLN:OE1	2.53	0.41
4:AK:132:GLN:O	4:AK:136:GLU:HG3	2.21	0.41
4:AK:154:ALA:O	4:AK:158:ASP:HB2	2.21	0.41
1:AN:63:PRO:O	1:AN:67:SER:N	2.52	0.41
1:AR:96:VAL:HG22	1:AR:152:TYR:CD2	2.55	0.41
1:AR:128:GLN:NE2	1:AR:132:GLU:OE2	2.53	0.41
2:AS:37:ARG:NH1	2:AS:99:GLY:HA2	2.35	0.41
2:AS:39:ARG:HH12	2:AS:42:SER:HB3	1.85	0.41
2:AS:71:ASN:HD22	2:AS:121:VAL:HA	1.85	0.41
2:AS:81:CYS:HA	10:BD:901:CYC:HAC1	1.91	0.41
2:AU:10:ASN:O	2:AU:14:VAL:HG23	2.21	0.41
1:AV:121:THR:HG23	10:AV:200:CYC:C1C	2.50	0.41

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:AW:68:PRO:HA	2:AW:73:TYR:CD1	2.55	0.41
10:AW:200:CYC:HC	10:AW:200:CYC:CMD	2.33	0.41
1:AX:85:MET:SD	1:AX:129:SER:HB2	2.61	0.41
6:BD:193:ASP:O	6:BD:197:VAL:HG22	2.20	0.41
6:BD:889:THR:CG2	2:CZ:118:SER:HB2	2.50	0.41
9:BH:131:LEU:HD11	9:BH:135:ALA:HB1	2.02	0.41
9:BH:238:LEU:O	9:BH:242:ARG:HG2	2.20	0.41
2:BL:106:GLU:OE1	6:CM:273:LEU:HA	2.20	0.41
1:BR:130:VAL:HB	1:BR:157:ILE:CD1	2.51	0.41
2:BT:113:LYS:NZ	2:CA:155:TYR:OH	2.51	0.41
1:BV:25:ARG:HH11	1:BV:25:ARG:HG2	1.85	0.41
1:BV:102:THR:HB	1:BV:103:PRO:HD3	2.02	0.41
2:BY:58:LYS:HE2	2:BY:135:GLU:OE1	2.20	0.41
1:BZ:85:MET:HE3	1:BZ:129:SER:HB2	2.01	0.41
1:BZ:92:VAL:O	1:BZ:96:VAL:HG23	2.20	0.41
2:CD:15:GLN:HB2	2:CD:17:LYS:NZ	2.35	0.41
2:CH:148:GLU:OE2	2:CH:149:MET:HG2	2.21	0.41
5:CK:12:SER:HB3	5:CK:17:ARG:CZ	2.50	0.41
6:CM:353:ARG:HA	6:CM:418:PHE:CZ	2.56	0.41
6:CM:783:THR:HB	6:CM:832:LYS:HG2	2.02	0.41
9:CQ:98:TYR:O	9:CQ:106:LYS:HE3	2.20	0.41
9:CQ:280:GLY:HA3	9:CQ:284:MET:HG3	2.02	0.41
1:CR:27:LYS:O	1:CR:30:VAL:HG22	2.21	0.41
2:CW:46:ALA:HB2	1:DA:154:ASP:CG	2.45	0.41
2:CZ:130:ILE:HB	2:CZ:157:CYS:SG	2.60	0.41
1:DA:66:VAL:HG23	1:DA:78:THR:CG2	2.50	0.41
2:DB:25:ASP:HA	2:DB:28:LYS:HE2	2.03	0.41
9:DI:88:ASN:HB2	9:DI:170:LYS:HD2	2.02	0.41
1:DJ:108:GLY:C	1:DJ:109:LEU:HD22	2.45	0.41
1:DL:50:ILE:HD11	1:DL:137:ALA:CA	2.50	0.41
1:DL:123:ILE:HA	1:DL:126:VAL:HG23	2.02	0.41
2:DR:2:GLN:HB2	2:DR:6:THR:OG1	2.19	0.41
1:DS:50:ILE:HD11	1:DS:137:ALA:HA	2.03	0.41
1:DS:76:GLU:HG2	1:DS:77:MET:HE3	2.03	0.41
1:DU:17:TYR:CE2	2:DV:90:ARG:HA	2.56	0.41
9:EA:310:LYS:HZ1	9:EA:312:LEU:HD12	1.86	0.41
3:AE:14:GLN:CB	3:AE:16:ARG:HG2	2.50	0.41
2:AI:126:THR:HA	10:AI:200:CYC:HMC1	2.02	0.41
1:AJ:56:ASP:HA	1:BX:69:GLY:HA2	2.02	0.41
2:AL:134:LYS:HD2	2:AL:150:GLY:CA	2.50	0.41
1:AN:51:VAL:HG12	1:AN:133:MET:SD	2.60	0.41

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:AQ:68:PRO:HG3	2:AQ:73:TYR:CE1	2.56	0.41
2:AQ:81:CYS:HA	10:AQ:200:CYC:HHD	2.03	0.41
2:AS:57:ALA:HA	2:AS:61:LEU:CD1	2.51	0.41
1:AX:68:PRO:HG2	8:BG:230:ASN:CG	2.45	0.41
2:AY:106:GLU:HG3	2:AY:107:ARG:HG3	2.02	0.41
1:AZ:85:MET:SD	1:AZ:129:SER:HB2	2.59	0.41
1:AZ:89:LEU:HB2	1:AZ:133:MET:CE	2.50	0.41
6:BD:751:LEU:CD1	6:BD:755:ILE:HD13	2.51	0.41
6:BD:805:THR:HA	6:BD:838:MET:SD	2.61	0.41
6:BD:872:LEU:CD2	6:BD:884:VAL:HG11	2.50	0.41
9:BH:204:ASN:HD22	9:BH:213:LEU:HA	1.85	0.41
9:BH:267:ILE:HD12	9:BH:292:LEU:HD12	2.02	0.41
2:BQ:122:PRO:HG2	2:BQ:125:SER:OG	2.20	0.41
2:BY:96:MET:HA	2:BY:152:TYR:CE2	2.55	0.41
1:BZ:91:LEU:O	1:BZ:104:ILE:HG12	2.20	0.41
2:CA:81:CYS:HA	10:CM:901:CYC:HAC1	1.95	0.41
1:CC:109:LEU:HD12	1:CC:109:LEU:HA	1.89	0.41
2:CF:24:MET:HG3	2:CF:28:LYS:HZ3	1.85	0.41
2:CF:122:PRO:HB2	2:CF:125:SER:CB	2.51	0.41
1:CG:93:THR:HA	1:CG:96:VAL:HG22	2.02	0.41
1:CG:109:LEU:O	1:CG:112:VAL:HG12	2.21	0.41
5:CJ:6:ILE:HG13	5:CJ:6:ILE:O	2.20	0.41
5:CK:38:ARG:HE	6:CM:641:ASP:HB3	1.85	0.41
6:CM:186:ILE:HG13	6:CM:190:CYS:HB3	2.01	0.41
6:CM:811:ARG:NH2	6:CM:814:LEU:HG	2.35	0.41
1:CR:121:THR:HG23	10:CR:200:CYC:C1C	2.50	0.41
1:CV:47:ARG:HG3	1:CV:48:GLU:N	2.35	0.41
2:CZ:81:CYS:O	10:CZ:200:CYC:HBC1	2.21	0.41
1:DA:2:SER:HB3	2:DB:3:ASP:OD2	2.20	0.41
2:DB:11:SER:O	2:DB:14:VAL:HG22	2.20	0.41
2:DB:73:TYR:OH	1:DC:90:ARG:NH2	2.33	0.41
9:DI:86:LEU:HD21	9:DI:98:TYR:HB3	2.02	0.41
1:DJ:91:LEU:HD21	1:DJ:107:ILE:CG2	2.47	0.41
1:DL:101:VAL:HG11	1:DU:20:PRO:HG2	2.01	0.41
1:DU:39:ILE:HD11	1:DU:145:ASP:OD1	2.21	0.41
5:DX:6:ILE:HG21	5:DX:36:TRP:CZ3	2.56	0.41
2:AB:134:LYS:HB2	2:AB:153:LEU:HD13	2.02	0.41
2:AD:65:VAL:HB	2:AD:72:MET:HB2	2.02	0.41
2:AD:81:CYS:HA	10:AD:200:CYC:HHD	2.02	0.41
3:AE:2:SER:O	3:AE:5:SER:OG	2.36	0.41
1:AJ:114:GLU:OE1	1:AJ:114:GLU:N	2.51	0.41

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:AL:43:VAL:HG12	2:AL:140:LEU:HD21	2.02	0.41
1:AN:115:MET:HG2	2:AS:78:TYR:CD1	2.56	0.41
1:AN:159:LYS:HE3	1:AN:159:LYS:HB3	1.85	0.41
2:AO:75:THR:HG23	10:AP:200:CYC:HAB1	2.02	0.41
1:AR:153:PHE:O	1:AR:157:ILE:HG13	2.19	0.41
1:AV:104:ILE:HG22	1:AV:109:LEU:CD1	2.50	0.41
2:AW:110:ASN:HD21	5:BB:56:ALA:CB	2.32	0.41
1:AX:66:VAL:HG21	8:BG:235:VAL:HG21	2.03	0.41
1:AX:109:LEU:HD11	1:AX:159:LYS:HB2	2.02	0.41
2:AY:1:MET:HB3	2:AY:106:GLU:OE2	2.21	0.41
1:AZ:63:PRO:O	1:AZ:67:SER:N	2.54	0.41
5:BA:43:ILE:HG21	5:BA:50:ILE:CG2	2.50	0.41
6:BD:211:ARG:HD2	6:BD:211:ARG:O	2.20	0.41
6:BD:258:PHE:CE1	6:BD:263:LYS:HG3	2.56	0.41
2:BJ:68:PRO:HA	2:BJ:73:TYR:CD1	2.55	0.41
2:BJ:78:TYR:CE1	2:BJ:82:ILE:HD11	2.56	0.41
1:BK:17:TYR:CE1	2:BL:90:ARG:HA	2.56	0.41
2:BL:55:ALA:O	2:BL:59:SER:CB	2.68	0.41
1:BP:80:THR:CG2	1:BP:83:ARG:HH21	2.34	0.41
1:BP:134:LYS:HB2	1:BP:153:PHE:CG	2.56	0.41
4:BS:72:ASN:ND2	4:BS:122:VAL:HG22	2.36	0.41
4:BS:138:ILE:CG2	4:BS:153:ILE:HG21	2.51	0.41
1:BV:39:ILE:HG23	1:BV:141:MET:CE	2.51	0.41
1:BV:101:VAL:CG1	1:CC:20:PRO:HG2	2.50	0.41
2:BW:93:THR:CA	2:BW:149:MET:HE1	2.51	0.41
1:BX:108:GLY:O	1:BX:112:VAL:HG21	2.21	0.41
1:BZ:73:TYR:OH	9:CQ:147:SER:HB3	2.21	0.41
2:CA:41:ALA:N	2:CA:96:MET:HE1	2.35	0.41
1:CC:24:ASP:HA	1:CC:27:LYS:HZ3	1.86	0.41
1:CE:57:ARG:CA	1:CE:60:GLN:HE21	2.34	0.41
2:CF:41:ALA:CB	2:CF:96:MET:HE1	2.41	0.41
6:CM:720:GLN:NE2	2:DV:1:MET:O	2.54	0.41
9:CQ:208:ASN:HA	9:CQ:210:PHE:CE1	2.56	0.41
2:CS:43:VAL:HG13	2:CS:141:VAL:HG12	2.02	0.41
2:CS:73:TYR:O	2:CS:74:THR:OG1	2.21	0.41
1:CT:24:ASP:OD1	1:CT:25:ARG:N	2.53	0.41
1:CV:85:MET:CE	1:CV:129:SER:HB2	2.51	0.41
1:DA:90:ARG:HD2	2:DB:17:LYS:O	2.20	0.41
2:DB:112:LEU:CD2	2:DB:160:LEU:HD21	2.50	0.41
1:DJ:87:TYR:O	1:DJ:91:LEU:HD13	2.20	0.41
1:DL:72:ALA:HA	1:DL:77:MET:HB3	2.03	0.41

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:DN:90:ARG:HA	2:DO:18:TYR:CE2	2.56	0.41
1:DN:104:ILE:HG21	1:DN:156:VAL:HG22	2.03	0.41
1:DQ:6:LYS:HD2	1:DQ:100:ASP:OD1	2.21	0.41
1:DQ:134:LYS:HB2	1:DQ:153:PHE:HB3	2.01	0.41
1:DU:97:VAL:HG11	2:DV:27:LEU:CD2	2.51	0.41
1:DU:101:VAL:HB	1:DU:155:PHE:CE2	2.56	0.41
2:DV:25:ASP:OD1	2:DV:26:LYS:N	2.54	0.41
5:DX:8:ALA:HB2	5:DX:50:ILE:HA	2.03	0.41
8:DY:207:GLU:OE1	8:DY:207:GLU:N	2.39	0.41
9:EA:223:LEU:HD23	9:EA:224:GLN:N	2.35	0.41
1:AA:105:GLU:HG3	1:AA:110:VAL:CG2	2.48	0.41
2:AB:59:SER:HB3	2:AB:132:ALA:CB	2.51	0.41
1:AC:98:SER:OG	1:AC:103:PRO:HG2	2.21	0.41
2:AD:8:VAL:CG1	2:AD:27:LEU:HD21	2.49	0.41
1:AH:62:ARG:CG	1:AH:65:ILE:HG12	2.50	0.41
1:AJ:18:LEU:CD1	4:AK:98:ILE:HD12	2.51	0.41
2:AL:101:ALA:HB1	2:AL:104:LEU:HD12	2.03	0.41
1:AN:25:ARG:HG2	1:AN:25:ARG:HH11	1.85	0.41
1:AP:37:LEU:HD23	2:AQ:28:LYS:NZ	2.36	0.41
1:AP:83:ARG:NH1	10:AP:200:CYC:O2A	2.43	0.41
2:AQ:64:ASP:OD1	2:AQ:64:ASP:N	2.53	0.41
2:AU:1:MET:HE2	1:AZ:8:ILE:CG2	2.50	0.41
2:AU:47:ASN:OD1	2:AU:140:LEU:HD13	2.20	0.41
2:AU:111:GLY:HA2	2:AU:114:GLU:CD	2.46	0.41
2:AW:133:ILE:O	2:AW:137:THR:HG23	2.20	0.41
6:BD:176:VAL:HG22	6:BD:230:GLU:OE2	2.20	0.41
6:BD:268:ALA:HA	6:BD:393:GLU:O	2.20	0.41
6:BD:365:PHE:O	6:BD:369:SER:OG	2.32	0.41
6:BD:458:PHE:CB	6:BD:552:GLY:HA3	2.50	0.41
9:BH:226:PRO:HG3	9:BH:305:LEU:HG	2.03	0.41
2:BL:104:LEU:HB3	2:BL:109:LEU:HD13	2.01	0.41
2:BN:133:ILE:O	2:BN:137:THR:HG22	2.20	0.41
1:BR:107:ILE:HG22	10:BR:200:CYC:HBB1	2.01	0.41
4:BS:105:LEU:HD23	4:BS:159:HIS:CD2	2.55	0.41
2:BT:100:ASP:OD1	2:BT:101:ALA:N	2.54	0.41
2:BT:126:THR:O	2:BT:130:ILE:HG13	2.21	0.41
1:BX:76:GLU:HG2	1:BX:77:MET:HE2	2.03	0.41
1:BX:81:CYS:HA	10:BX:200:CYC:HHD	2.02	0.41
2:BY:39:ARG:O	2:BY:43:VAL:HG23	2.20	0.41
2:BY:160:LEU:HD12	2:BY:160:LEU:HA	1.91	0.41
1:BZ:3:ILE:HD11	1:BZ:25:ARG:HD3	2.02	0.41

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:BZ:47:ARG:O	1:BZ:51:VAL:HG23	2.20	0.41
1:BZ:66:VAL:HG21	8:CP:227:PRO:HB2	2.02	0.41
2:CA:108:VAL:O	2:CA:112:LEU:HB2	2.20	0.41
1:CE:30:VAL:CG2	2:CF:31:PHE:HA	2.50	0.41
1:CG:101:VAL:HG12	1:CG:152:TYR:CD1	2.55	0.41
2:CH:56:VAL:CG1	2:CH:85:LEU:HD12	2.50	0.41
6:CM:788:MET:HG2	6:CM:792:TYR:HB3	2.03	0.41
1:CR:77:MET:C	1:CR:80:THR:HG22	2.44	0.41
2:CU:124:SER:O	2:CU:128:GLN:HG3	2.21	0.41
1:CY:48:GLU:HG2	1:CY:49:THR:H	1.86	0.41
1:CY:50:ILE:HG21	1:CY:89:LEU:HD11	2.03	0.41
1:CY:100:ASP:O	1:CY:103:PRO:HD2	2.21	0.41
1:DC:30:VAL:HG21	2:DD:34:GLY:HA3	2.03	0.41
1:DC:113:ARG:NH2	1:DC:161:SER:OXT	2.53	0.41
8:DG:221:MET:HG3	1:DQ:52:LYS:HD3	2.02	0.41
2:DK:129:ALA:O	2:DK:133:ILE:HG13	2.20	0.41
1:DN:58:LEU:CD1	1:DN:128:GLN:HB3	2.51	0.41
1:DQ:5:THR:CA	1:DQ:8:ILE:HG22	2.50	0.41
1:DQ:23:LEU:HD22	2:DR:38:VAL:CG2	2.36	0.41
1:DQ:47:ARG:HA	1:DQ:50:ILE:HG22	2.03	0.41
2:DT:50:THR:O	2:DT:53:LYS:HB3	2.20	0.41
1:DU:37:LEU:HD22	2:DV:24:MET:CE	2.50	0.41
2:DV:71:ASN:CB	2:DV:122:PRO:HD3	2.50	0.41
5:DX:12:SER:HB2	5:DX:17:ARG:NH2	2.36	0.41
9:EA:98:TYR:CE1	9:EA:161:MET:HB2	2.55	0.41
2:AB:81:CYS:O	2:AB:85:LEU:HG	2.20	0.41
2:AB:144:ASP:N	2:AB:144:ASP:OD1	2.52	0.41
3:AE:58:LEU:CD2	3:AE:129:ALA:HB2	2.43	0.41
1:AJ:89:LEU:HB2	1:AJ:133:MET:HE1	2.01	0.41
4:AK:50:THR:O	4:AK:54:ARG:HG3	2.21	0.41
2:AO:19:LEU:O	2:AO:24:MET:HE1	2.21	0.41
2:AO:110:ASN:ND2	6:BD:481:LYS:O	2.52	0.41
1:AR:131:ARG:HG3	1:AR:157:ILE:HD13	2.01	0.41
2:AU:57:ALA:HA	2:AU:61:LEU:HD12	2.03	0.41
1:AV:40:ALA:O	1:AV:44:THR:HG23	2.20	0.41
2:AY:78:TYR:O	2:AY:82:ILE:HG12	2.21	0.41
6:BD:253:LEU:HD12	6:BD:254:PRO:CD	2.49	0.41
6:BD:358:ARG:HB2	6:BD:434:PHE:HB3	2.03	0.41
9:BH:64:ALA:HA	9:BH:101:TRP:CZ2	2.55	0.41
2:BL:58:LYS:HE2	2:BL:58:LYS:HB3	1.89	0.41
2:BL:72:MET:HE3	2:BL:78:TYR:CD1	2.56	0.41

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:BL:83:ARG:NE	5:CJ:22:LEU:HD11	2.36	0.41
3:BM:39:ILE:CG2	3:BM:96:VAL:HG11	2.43	0.41
3:BM:64:GLU:O	3:BM:70:GLY:HA3	2.20	0.41
3:BM:105:GLU:O	3:BM:110:ILE:HG13	2.21	0.41
2:BN:12:ALA:HB1	2:BN:17:LYS:O	2.21	0.41
2:BT:5:ILE:HG21	6:CM:169:ALA:HA	2.02	0.41
2:BT:24:MET:SD	6:CM:58:GLU:HB2	2.61	0.41
1:BX:62:ARG:O	1:BX:65:ILE:HG12	2.20	0.41
1:BX:126:VAL:O	1:BX:130:VAL:HG23	2.21	0.41
1:BZ:14:GLU:OE1	1:BZ:16:ARG:NE	2.35	0.41
1:BZ:25:ARG:HG3	1:CE:25:ARG:HD3	2.02	0.41
1:CC:101:VAL:HB	1:CC:155:PHE:CE2	2.55	0.41
2:CD:79:ALA:HA	2:CD:82:ILE:HG22	2.02	0.41
2:CF:100:ASP:OD1	2:CF:101:ALA:N	2.54	0.41
1:CG:107:ILE:CG2	10:CG:200:CYC:HBB1	2.47	0.41
6:CM:31:GLN:HG2	6:CM:267:TYR:OH	2.21	0.41
6:CM:52:LEU:HD11	6:CM:216:ALA:CB	2.47	0.41
9:CQ:131:LEU:HD11	9:CQ:135:ALA:HB1	2.01	0.41
9:CQ:235:GLU:HG3	1:DA:45:GLY:HA2	2.03	0.41
2:CS:123:ILE:O	2:CS:127:VAL:HG23	2.20	0.41
1:CT:67:SER:C	1:CT:73:TYR:HB2	2.45	0.41
1:CV:48:GLU:HG2	1:CV:49:THR:N	2.36	0.41
2:CZ:66:THR:CA	2:CZ:72:MET:HB3	2.33	0.41
2:DB:36:LEU:HD11	2:DB:144:ASP:CB	2.51	0.41
1:DC:50:ILE:CD1	1:DC:136:VAL:HG12	2.49	0.41
8:DH:203:PHE:HZ	8:DY:219:LEU:HD13	1.86	0.41
1:DL:131:ARG:N	1:DL:157:ILE:HD11	2.36	0.41
2:DO:71:ASN:ND2	10:DO:200:CYC:HMD2	2.36	0.41
2:DR:43:VAL:HG11	2:DR:141:VAL:HG12	2.03	0.41
10:DS:200:CYC:CMA	10:DS:200:CYC:HB	2.34	0.41
1:DU:140:LEU:HD12	1:DU:140:LEU:O	2.21	0.41
5:DX:27:PHE:CE1	5:DX:29:LYS:HE3	2.56	0.41
9:EA:32:ASN:OD1	9:EA:35:ASP:HB2	2.21	0.41
9:EA:151:ILE:O	9:EA:155:ARG:HG3	2.19	0.41
9:EA:269:VAL:HG23	9:EA:288:TRP:CE3	2.49	0.41
1:AA:50:ILE:HD12	1:AA:136:VAL:HG12	2.03	0.41
1:AA:71:ASN:OD1	10:AA:200:CYC:HBD2	2.20	0.41
1:AC:91:LEU:HD21	1:AC:107:ILE:CG2	2.51	0.41
1:AC:123:ILE:HG13	1:AC:160:MET:O	2.21	0.41
2:AD:66:THR:HG21	10:AE:200:CYC:CBA	2.49	0.41
1:AH:65:ILE:HG23	10:AH:200:CYC:OC	2.21	0.41

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:AL:119:LEU:HD23	2:AL:119:LEU:HA	1.95	0.41
2:AL:131:GLN:O	2:AL:135:GLU:HG2	2.20	0.41
1:AN:61:LYS:HG2	1:AN:62:ARG:HG2	2.03	0.41
2:AO:122:PRO:HB2	2:AO:125:SER:HB2	2.02	0.41
1:AP:89:LEU:HD12	1:AP:89:LEU:HA	1.91	0.41
1:AP:108:GLY:O	1:AP:112:VAL:HG21	2.21	0.41
1:AP:143:SER:HB2	1:DN:147:ALA:HB2	2.02	0.41
1:AR:47:ARG:HD2	2:AS:18:TYR:CZ	2.56	0.41
1:AR:109:LEU:O	1:AR:112:VAL:HG12	2.20	0.41
2:AS:91:TYR:HB3	2:AS:104:LEU:HD23	2.03	0.41
2:AU:60:LEU:HD23	2:AU:65:VAL:HG11	2.02	0.41
1:AV:48:GLU:HG2	1:AV:49:THR:N	2.36	0.41
1:AV:88:TYR:OH	1:AV:112:VAL:HG21	2.21	0.41
1:AV:98:SER:HB3	2:AW:9:ILE:HD11	2.03	0.41
2:AW:47:ASN:CG	2:AW:140:LEU:HD21	2.45	0.41
1:AX:2:SER:HB2	1:AX:5:THR:HB	2.01	0.41
1:AX:123:ILE:HD11	1:AX:160:MET:SD	2.61	0.41
1:AX:127:ALA:HB2	1:AX:160:MET:HB3	2.03	0.41
2:AY:62:TYR:N	2:AY:66:THR:HG21	2.36	0.41
5:BA:6:ILE:O	5:BA:6:ILE:HG13	2.20	0.41
6:BD:176:VAL:HG22	6:BD:230:GLU:CD	2.46	0.41
6:BD:755:ILE:HD12	6:BD:758:SER:N	2.36	0.41
10:BD:901:CYC:HBD1	10:BD:901:CYC:HHA	2.03	0.41
9:BH:88:ASN:HD21	9:BH:174:GLU:HB3	1.86	0.41
9:BH:112:ARG:NH2	9:BH:116:LEU:HD21	2.36	0.41
1:BI:37:LEU:O	1:BI:41:GLU:HG3	2.20	0.41
1:BI:109:LEU:HD12	1:BI:109:LEU:HA	1.94	0.41
1:BK:2:SER:OG	1:BK:4:VAL:HG22	2.20	0.41
1:BK:25:ARG:HG3	1:BK:26:ILE:N	2.36	0.41
2:BL:39:ARG:HH12	2:BL:42:SER:HB3	1.86	0.41
2:BL:116:TYR:HB3	2:BL:121:VAL:HB	2.03	0.41
3:BM:5:SER:HB3	2:BN:3:ASP:CG	2.45	0.41
3:BM:23:LEU:HD13	2:BN:42:SER:OG	2.21	0.41
2:BN:64:ASP:N	2:BN:64:ASP:OD1	2.54	0.41
2:BN:116:TYR:CD1	2:BN:121:VAL:HG21	2.56	0.41
1:BP:41:GLU:HA	1:BP:44:THR:CG2	2.51	0.41
1:BP:158:GLY:HA2	1:BP:161:SER:O	2.21	0.41
1:BR:91:LEU:O	1:BR:104:ILE:HG12	2.21	0.41
4:BS:123:PRO:HB3	2:CA:64:ASP:HB3	2.03	0.41
4:BS:129:ARG:HE	4:BS:133:ILE:HG13	1.86	0.41
2:BT:35:GLU:OE2	6:CM:48:GLN:HG3	2.21	0.41

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:BT:81:CYS:O	2:BT:85:LEU:HG	2.20	0.41
1:BV:71:ASN:OD1	10:BV:200:CYC:HMD2	2.21	0.41
1:BX:48:GLU:HG2	1:BX:49:THR:N	2.35	0.41
1:BX:102:THR:HB	1:BX:103:PRO:HD3	2.02	0.41
1:BX:113:ARG:NH1	1:BX:160:MET:O	2.51	0.41
1:BX:144:ASP:OD1	1:BX:144:ASP:N	2.52	0.41
2:BY:144:ASP:OD1	2:BY:144:ASP:N	2.51	0.41
1:CC:8:ILE:HG22	2:CD:1:MET:HE2	2.03	0.41
1:CC:74:GLY:O	1:CC:78:THR:HG23	2.20	0.41
2:CD:121:VAL:HG13	10:CD:200:CYC:NC	2.36	0.41
1:CE:57:ARG:O	1:CE:60:GLN:HG3	2.21	0.41
1:CG:147:ALA:HA	1:CG:150:SER:OG	2.20	0.41
2:CH:116:TYR:HB3	2:CH:123:ILE:HD11	2.02	0.41
5:CJ:17:ARG:NE	6:CM:240:VAL:HG21	2.36	0.41
6:CM:141:ILE:HG23	6:CM:149:MET:HE2	2.03	0.41
6:CM:701:ARG:HD2	6:CM:705:GLU:OE2	2.20	0.41
6:CM:760:PHE:O	6:CM:764:GLU:HG3	2.21	0.41
6:CM:827:ALA:HB2	2:DR:87:TYR:OH	2.20	0.41
9:CQ:284:MET:HE1	9:CQ:307:ALA:CB	2.50	0.41
2:CS:47:ASN:O	2:CS:50:THR:HB	2.21	0.41
1:CT:70:GLY:N	1:CY:64:ASP:HB3	2.35	0.41
2:CU:83:ARG:NH2	2:CU:84:ASP:OD1	2.44	0.41
2:CW:50:THR:HG21	9:DI:62:GLN:NE2	2.36	0.41
1:CY:43:LEU:HD11	1:CY:141:MET:HE1	2.03	0.41
1:CY:152:TYR:O	1:CY:156:VAL:HG12	2.21	0.41
2:CZ:35:GLU:CA	2:CZ:38:VAL:HG12	2.47	0.41
2:CZ:112:LEU:HA	2:CZ:115:THR:CG2	2.51	0.41
1:DC:116:TYR:CE2	1:DC:126:VAL:HG21	2.56	0.41
1:DC:124:GLU:OE1	1:DC:124:GLU:N	2.39	0.41
8:DG:227:PRO:HG3	1:DQ:59:PHE:CE2	2.55	0.41
8:DG:238:GLN:HA	1:DL:52:LYS:CD	2.50	0.41
9:DI:88:ASN:HB2	9:DI:170:LYS:CD	2.50	0.41
9:DI:286:ILE:HD13	9:DI:288:TRP:CZ2	2.55	0.41
1:DL:147:ALA:HA	1:DL:150:SER:HB3	2.02	0.41
1:DN:109:LEU:HD13	1:DN:159:LYS:HG2	2.03	0.41
1:DN:159:LYS:HD3	1:DN:159:LYS:HA	1.92	0.41
1:DU:92:VAL:HG11	1:DU:153:PHE:CZ	2.56	0.41
2:DV:33:SER:HB3	2:DV:37:ARG:HE	1.85	0.41
9:EA:69:LYS:HB3	9:EA:69:LYS:HE3	1.82	0.41
1:AA:156:VAL:HG13	1:AA:160:MET:HE3	2.02	0.41
2:AD:64:ASP:OD1	2:AD:67:ARG:NH1	2.48	0.41

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
4:AK:67:ILE:HG21	10:BD:902:CYC:HBA2	2.02	0.41
10:AL:200:CYC:CMD	10:AL:200:CYC:HC	2.34	0.41
2:AO:41:ALA:HB2	2:AO:96:MET:HE3	2.03	0.41
2:AO:56:VAL:HG12	2:AO:61:LEU:HG	2.02	0.41
1:AP:109:LEU:HA	1:AP:112:VAL:CG2	2.50	0.41
2:AQ:61:LEU:HA	2:AQ:66:THR:HG21	2.03	0.41
2:AS:141:VAL:HG22	2:AS:145:ALA:HB3	2.03	0.41
2:AU:73:TYR:CE1	10:AV:200:CYC:HBB3	2.56	0.41
2:AU:119:LEU:HD21	5:BB:4:PHE:HZ	1.85	0.41
1:AV:141:MET:HE1	1:AV:145:ASP:HB3	2.03	0.41
1:AX:36:ARG:NH1	1:AX:36:ARG:HB3	2.35	0.41
1:AX:126:VAL:HG22	10:AX:200:CYC:HMC2	2.02	0.41
2:AY:81:CYS:HA	10:AY:200:CYC:HAC1	1.88	0.41
6:BD:153:LEU:HA	6:BD:156:MET:SD	2.61	0.41
6:BD:458:PHE:HB3	6:BD:552:GLY:C	2.46	0.41
6:BD:489:PRO:HD2	6:BD:492:ASN:ND2	2.34	0.41
6:BD:816:GLN:HE22	6:BD:887:SER:HB3	1.86	0.41
9:BH:281:ASN:O	9:BH:282:VAL:HB	2.21	0.41
1:BI:5:THR:O	1:BI:9:VAL:HG23	2.21	0.41
1:BK:25:ARG:NH1	1:BK:26:ILE:HD11	2.36	0.41
3:BM:37:ILE:HA	3:BM:97:LEU:HD21	2.03	0.41
1:BP:131:ARG:O	1:BP:135:GLU:HG2	2.21	0.41
2:BQ:115:THR:CG2	6:CM:428:PRO:HG2	2.51	0.41
10:BQ:200:CYC:HMA3	10:BQ:200:CYC:NB	2.32	0.41
10:CC:200:CYC:O1A	2:CH:66:THR:HG21	2.21	0.41
2:CD:110:ASN:O	5:CK:44:MET:HE2	2.21	0.41
2:CF:116:TYR:CD2	2:CF:160:LEU:HD21	2.55	0.41
2:CF:144:ASP:N	2:CF:144:ASP:OD1	2.54	0.41
1:CG:144:ASP:OD1	1:CG:145:ASP:N	2.52	0.41
6:CM:182:LEU:HD12	6:CM:182:LEU:O	2.20	0.41
6:CM:748:GLU:O	6:CM:749:ARG:HD3	2.20	0.41
9:CQ:277:TRP:CD1	9:CQ:305:LEU:HD21	2.55	0.41
1:CR:116:TYR:CD2	1:CR:123:ILE:HG22	2.55	0.41
1:CR:141:MET:HE1	1:CR:145:ASP:O	2.21	0.41
10:CR:200:CYC:HAA1	2:CW:61:LEU:HD22	2.03	0.41
2:CS:77:ARG:CB	10:CS:200:CYC:HMD1	2.47	0.41
2:CW:134:LYS:HA	2:CW:137:THR:HG22	2.03	0.41
1:CY:90:ARG:HG2	2:CZ:18:TYR:CE1	2.56	0.41
2:CZ:71:ASN:ND2	2:CZ:121:VAL:HA	2.36	0.41
2:DD:112:LEU:HD13	10:DD:200:CYC:HMB2	2.02	0.41
8:DG:237:ALA:HA	8:DG:240:ILE:CD1	2.51	0.41

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
9:DI:248:LEU:HD11	9:DI:277:TRP:CZ3	2.56	0.41
1:DL:17:TYR:HB3	2:DM:45:SER:HB2	2.03	0.41
2:DM:134:LYS:HE2	2:DM:154:ASP:OD1	2.21	0.41
1:DQ:93:THR:O	1:DQ:97:VAL:HG23	2.21	0.41
2:DT:3:ASP:OD1	2:DT:6:THR:OG1	2.27	0.41
1:DU:115:MET:HE1	10:DU:200:CYC:C2B	2.51	0.41
9:EA:86:LEU:HD23	9:EA:95:CYS:SG	2.61	0.41
9:EA:223:LEU:HG	9:EA:301:VAL:CG1	2.50	0.41
1:AA:81:CYS:HA	10:AA:200:CYC:HAC1	1.98	0.40
1:AA:81:CYS:O	1:AA:85:MET:HG2	2.21	0.40
3:AE:17:TYR:CE1	2:AF:90:ARG:HG3	2.56	0.40
2:AF:114:GLU:CD	5:BA:53:VAL:HG12	2.45	0.40
4:AK:65:GLU:O	4:AK:71:GLY:HA3	2.21	0.40
2:AL:132:ALA:O	2:AL:136:VAL:HG23	2.21	0.40
1:AN:88:TYR:HD1	1:AN:91:LEU:HD12	1.86	0.40
2:AO:51:ILE:HG23	2:AO:136:VAL:HG12	2.03	0.40
1:AP:68:PRO:HA	1:AP:73:TYR:CG	2.55	0.40
1:AR:76:GLU:HG2	1:AR:77:MET:N	2.36	0.40
2:AS:36:LEU:HD11	2:AS:144:ASP:CG	2.45	0.40
1:AV:39:ILE:HG23	1:AV:141:MET:CE	2.51	0.40
1:AV:58:LEU:CD2	1:AV:129:SER:HB3	2.41	0.40
1:AZ:58:LEU:HD22	1:AZ:129:SER:HB2	2.04	0.40
1:AZ:61:LYS:HZ1	1:BR:60:GLN:HB3	1.86	0.40
5:BB:1:MET:CE	5:BB:32:PRO:HB3	2.51	0.40
6:BD:446:VAL:HB	6:BD:617:ARG:HG3	2.03	0.40
9:BH:92:THR:O	9:BH:96:ARG:HG2	2.21	0.40
9:BH:197:THR:HG21	9:BH:296:GLY:O	2.21	0.40
9:BH:232:VAL:CG1	9:EA:314:ASN:HD22	2.28	0.40
1:BK:63:PRO:HA	1:BK:66:VAL:HG12	2.02	0.40
1:BP:156:VAL:CG1	1:BP:160:MET:HE3	2.51	0.40
2:BQ:85:LEU:HD23	2:BQ:85:LEU:HA	1.84	0.40
1:BR:141:MET:HE2	1:BR:141:MET:HB3	1.89	0.40
4:BS:4:ALA:O	4:BS:7:THR:HG22	2.21	0.40
2:BT:3:ASP:OD1	2:BT:6:THR:HG22	2.20	0.40
2:BT:134:LYS:HD2	2:BT:150:GLY:HA3	2.03	0.40
1:CC:105:GLU:OE1	1:CC:159:LYS:NZ	2.54	0.40
2:CD:64:ASP:HA	2:CD:67:ARG:NH1	2.36	0.40
1:CE:27:LYS:HD2	2:CF:35:GLU:OE1	2.21	0.40
2:CF:93:THR:OG1	2:CF:149:MET:HE1	2.21	0.40
7:CO:83:PRO:O	7:CO:86:VAL:HG12	2.21	0.40
9:CQ:250:LEU:HD22	9:CQ:286:ILE:HD12	2.03	0.40

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
9:CQ:291:LEU:HD23	9:CQ:299:PHE:HE2	1.86	0.40
1:CR:36:ARG:CZ	1:CR:99:GLY:HA2	2.51	0.40
1:CV:25:ARG:HH12	1:DA:21:GLY:C	2.29	0.40
1:CV:46:SER:CB	1:CV:140:LEU:HD23	2.50	0.40
1:CY:47:ARG:CA	1:CY:50:ILE:HG22	2.52	0.40
1:CY:50:ILE:CD1	1:CY:136:VAL:HB	2.51	0.40
1:CY:68:PRO:HA	1:CY:73:TYR:CD1	2.56	0.40
2:CZ:71:ASN:ND2	2:CZ:122:PRO:HD3	2.31	0.40
1:DA:152:TYR:O	1:DA:156:VAL:HG23	2.20	0.40
1:DC:68:PRO:HG3	1:DC:73:TYR:CE1	2.56	0.40
9:DI:112:ARG:NH1	9:DI:115:GLU:OE1	2.54	0.40
1:DJ:27:LYS:O	1:DJ:30:VAL:HG22	2.22	0.40
1:DJ:50:ILE:HD11	1:DJ:140:LEU:CD1	2.36	0.40
1:DJ:123:ILE:HG12	1:DJ:160:MET:O	2.21	0.40
2:DK:39:ARG:NH1	2:DK:40:ALA:HB2	2.37	0.40
2:DK:100:ASP:OD2	2:DK:102:SER:HB3	2.21	0.40
1:DL:130:VAL:HG21	1:DL:160:MET:HE1	2.03	0.40
1:DN:27:LYS:HA	1:DN:30:VAL:HG22	2.02	0.40
2:DO:108:VAL:O	2:DO:112:LEU:HD22	2.21	0.40
2:DR:104:LEU:O	2:DR:108:VAL:N	2.43	0.40
1:DS:39:ILE:CD1	1:DS:145:ASP:HB3	2.51	0.40
1:DS:81:CYS:HB3	10:DS:200:CYC:HBC1	1.88	0.40
2:DV:71:ASN:HA	2:DV:77:ARG:HH11	1.85	0.40
8:DZ:241:ASN:O	8:DZ:245:SER:OG	2.25	0.40
2:AB:112:LEU:HD13	10:AB:200:CYC:CMB	2.46	0.40
3:AE:7:VAL:HG11	3:AE:26:ILE:HD11	2.02	0.40
1:AH:83:ARG:HD2	8:BG:206:GLN:NE2	2.37	0.40
2:AI:56:VAL:HG21	2:AI:82:ILE:CD1	2.51	0.40
2:AI:88:TYR:OH	2:AI:116:TYR:OH	2.28	0.40
4:AK:89:TYR:CB	4:AK:134:MET:HE1	2.51	0.40
4:AK:102:ASN:O	4:AK:105:LEU:HB3	2.21	0.40
1:AV:27:LYS:HE2	2:AW:38:VAL:HG11	2.02	0.40
1:AV:57:ARG:O	1:AV:60:GLN:HG3	2.20	0.40
1:AX:85:MET:SD	10:AX:200:CYC:HBC1	2.61	0.40
1:AX:131:ARG:N	1:AX:157:ILE:HD11	2.37	0.40
2:AY:10:ASN:ND2	6:BD:532:VAL:HB	2.36	0.40
2:AY:126:THR:CB	10:AY:200:CYC:HMC1	2.50	0.40
6:BD:447:TYR:HB2	6:BD:451:ASN:HD22	1.86	0.40
6:BD:756:ILE:HG22	6:BD:757:ASN:ND2	2.37	0.40
6:BD:785:GLU:O	6:BD:789:LYS:HG3	2.20	0.40
1:BI:79:ALA:HA	1:BI:82:LEU:HG	2.02	0.40

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
10:BI:200:CYC:O2A	2:BN:66:THR:HG21	2.20	0.40
2:BJ:123:ILE:H	2:BJ:123:ILE:HD12	1.86	0.40
1:BP:109:LEU:HA	1:BP:112:VAL:CG2	2.52	0.40
1:BR:7:SER:HA	1:BR:22:GLU:OE2	2.22	0.40
10:BW:200:CYC:CMD	10:BW:200:CYC:HC	2.35	0.40
1:BX:44:THR:O	1:BX:47:ARG:HG2	2.22	0.40
2:BY:71:ASN:ND2	2:BY:122:PRO:HD3	2.37	0.40
2:CA:6:THR:HG22	2:CA:10:ASN:OD1	2.21	0.40
1:CC:109:LEU:HD13	1:CC:160:MET:HE2	2.03	0.40
2:CD:47:ASN:OD1	2:CD:140:LEU:HD13	2.21	0.40
1:CG:44:THR:HG23	2:CH:18:TYR:HB3	2.03	0.40
6:CM:76:PHE:CE1	6:CM:139:ILE:HD12	2.56	0.40
6:CM:253:LEU:HD12	6:CM:254:PRO:CD	2.51	0.40
6:CM:293:ASP:OD1	6:CM:295:THR:HG22	2.22	0.40
6:CM:803:MET:HE1	6:CM:872:LEU:CD2	2.51	0.40
9:CQ:198:VAL:HG12	9:CQ:202:MET:CE	2.37	0.40
9:CQ:231:ILE:CG2	9:CQ:236:ASN:HB3	2.52	0.40
1:CR:8:ILE:HG21	2:CS:1:MET:CE	2.37	0.40
1:CR:62:ARG:O	1:CR:65:ILE:HG12	2.21	0.40
1:CR:66:VAL:HG22	1:CR:66:VAL:O	2.22	0.40
1:CR:76:GLU:HB3	8:DZ:247:PRO:CG	2.35	0.40
1:CR:115:MET:HG3	2:CW:78:TYR:CE1	2.55	0.40
2:CS:34:GLY:O	2:CS:38:VAL:HG23	2.21	0.40
2:CS:40:ALA:O	2:CS:44:ILE:HG12	2.20	0.40
1:CV:3:ILE:HD11	1:CV:25:ARG:HE	1.86	0.40
1:CV:124:GLU:HG2	1:CV:125:ALA:N	2.36	0.40
2:CW:56:VAL:CG1	2:CW:85:LEU:HD12	2.51	0.40
2:CZ:10:ASN:O	2:CZ:14:VAL:HG22	2.21	0.40
2:CZ:39:ARG:NH1	2:CZ:141:VAL:O	2.55	0.40
2:CZ:82:ILE:HD13	2:CZ:85:LEU:HD12	2.03	0.40
1:DC:116:TYR:HD1	1:DC:116:TYR:HA	1.76	0.40
2:DD:94:TYR:HB3	2:DD:103:ILE:HD13	2.03	0.40
9:DI:187:LYS:HA	9:DI:203:ASP:OD1	2.21	0.40
1:DJ:63:PRO:HG2	1:DS:68:PRO:O	2.20	0.40
1:DN:43:LEU:HD23	1:DN:43:LEU:HA	1.85	0.40
1:DN:87:TYR:O	1:DN:91:LEU:HG	2.22	0.40
2:DO:57:ALA:CA	2:DO:61:LEU:HD12	2.45	0.40
2:DT:84:ASP:OD2	10:DT:200:CYC:ND	2.54	0.40
1:DU:64:ASP:OD1	1:DU:64:ASP:N	2.50	0.40
9:EA:248:LEU:HD21	9:EA:277:TRP:CD2	2.56	0.40
1:AC:7:SER:CB	1:AC:25:ARG:HH22	2.33	0.40

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
4:AK:41:ALA:HA	4:AK:44:ILE:CD1	2.51	0.40
4:AK:114:ARG:HG2	4:AK:118:ASN:ND2	2.35	0.40
1:AN:104:ILE:HG21	1:AN:156:VAL:CG2	2.51	0.40
1:AP:39:ILE:HG12	1:AP:145:ASP:HB3	2.02	0.40
1:AP:47:ARG:HG3	1:AP:48:GLU:N	2.36	0.40
1:AR:56:ASP:OD1	1:AR:60:GLN:NE2	2.54	0.40
2:AU:104:LEU:O	2:AU:108:VAL:HG22	2.22	0.40
2:AU:108:VAL:HG23	2:AU:109:LEU:N	2.36	0.40
2:AW:25:ASP:HA	2:AW:28:LYS:NZ	2.36	0.40
2:AW:100:ASP:OD1	2:AW:101:ALA:N	2.54	0.40
2:AW:112:LEU:HD12	2:AW:112:LEU:HA	1.72	0.40
1:AZ:131:ARG:CA	1:AZ:157:ILE:HD11	2.50	0.40
6:BD:344:GLU:OE2	6:BD:428:PRO:HB3	2.22	0.40
6:BD:574:SER:OG	6:BD:577:GLU:OE1	2.40	0.40
6:BD:702:THR:CB	2:DT:151:VAL:HG22	2.48	0.40
6:BD:788:MET:HB2	6:BD:788:MET:HE2	2.01	0.40
7:BF:93:PRO:O	7:BF:95:PRO:HD3	2.22	0.40
1:BK:83:ARG:NH2	1:BK:84:ASP:OD1	2.36	0.40
3:BM:110:ILE:CG2	5:CJ:16:ILE:HD11	2.49	0.40
1:BP:63:PRO:O	1:BP:67:SER:N	2.54	0.40
1:BP:77:MET:SD	10:BP:200:CYC:HAD1	2.61	0.40
1:BR:61:LYS:HD2	1:BR:132:GLU:OE2	2.21	0.40
2:BT:5:ILE:HD13	6:CM:46:TYR:OH	2.21	0.40
1:BV:142:SER:O	1:BV:146:ALA:HB2	2.22	0.40
2:CF:20:ASP:OD1	2:CF:21:GLY:N	2.47	0.40
5:CJ:4:PHE:HE1	5:CJ:33:TYR:HD1	1.69	0.40
5:CJ:15:ARG:CD	5:CJ:22:LEU:HD23	2.52	0.40
5:CJ:18:THR:O	6:CM:239:LYS:HG2	2.21	0.40
6:CM:56:ILE:HD12	6:CM:220:VAL:HG21	2.03	0.40
6:CM:167:ILE:HD12	6:CM:219:ILE:HG22	2.03	0.40
6:CM:413:MET:SD	6:CM:433:THR:HB	2.61	0.40
6:CM:425:ARG:HD3	6:CM:429:GLN:HB2	2.03	0.40
9:CQ:290:PHE:CE1	9:CQ:301:VAL:HG23	2.57	0.40
10:CV:200:CYC:HBC2	10:CV:200:CYC:H2C	1.72	0.40
1:CY:47:ARG:C	1:CY:50:ILE:HG22	2.47	0.40
1:CY:64:ASP:HA	1:CY:67:SER:OG	2.21	0.40
1:CY:71:ASN:ND2	1:CY:121:THR:HA	2.36	0.40
1:CY:131:ARG:HA	1:CY:157:ILE:HD11	2.03	0.40
2:CZ:61:LEU:HD12	2:CZ:61:LEU:HA	1.88	0.40
2:CZ:111:GLY:O	2:CZ:114:GLU:HG3	2.21	0.40
1:DC:71:ASN:O	1:DC:71:ASN:ND2	2.51	0.40

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:DD:36:LEU:O	2:DD:39:ARG:HG2	2.21	0.40
9:DI:76:PRO:HB3	9:DI:123:ALA:CB	2.47	0.40
1:DJ:66:VAL:O	1:DJ:73:TYR:HD1	2.05	0.40
2:DK:64:ASP:N	2:DK:64:ASP:OD1	2.51	0.40
10:DN:200:CYC:HBC2	10:DN:200:CYC:H2C	1.76	0.40
2:DR:25:ASP:OD1	2:DR:25:ASP:N	2.53	0.40
2:DT:61:LEU:HD22	10:DU:200:CYC:CBA	2.49	0.40
1:DU:91:LEU:O	1:DU:104:ILE:HG12	2.21	0.40
2:DV:100:ASP:OD1	2:DV:101:ALA:N	2.55	0.40
2:DV:122:PRO:HG2	2:DV:125:SER:OG	2.21	0.40
5:DX:34:ASP:N	5:DX:34:ASP:OD1	2.53	0.40
1:AA:46:SER:CB	1:AA:140:LEU:HD21	2.52	0.40
1:AC:159:LYS:HD2	1:AC:159:LYS:HA	1.94	0.40
3:AE:40:ALA:HB2	3:AE:97:LEU:HD11	2.04	0.40
2:AF:37:ARG:NH1	2:AF:148:GLU:OE2	2.55	0.40
2:AI:24:MET:O	2:AI:28:LYS:HG3	2.21	0.40
2:AI:53:LYS:HE2	1:AJ:118:SER:O	2.21	0.40
1:AJ:92:VAL:O	1:AJ:96:VAL:HG23	2.22	0.40
2:AL:18:TYR:CE2	6:BD:161:ARG:HA	2.57	0.40
2:AO:78:TYR:CE1	1:AP:115:MET:HG3	2.56	0.40
6:BD:64:ALA:O	6:BD:68:VAL:HG22	2.20	0.40
6:BD:867:PRO:HG3	2:DB:115:THR:OG1	2.22	0.40
7:BF:113:MET:HE3	7:BF:113:MET:HB2	1.86	0.40
1:BI:108:GLY:HA2	2:BN:75:THR:OG1	2.22	0.40
1:BK:51:VAL:HG12	1:BK:82:LEU:HD12	2.02	0.40
2:BL:92:ALA:HB1	2:BL:149:MET:HE3	2.02	0.40
3:BM:112:VAL:HA	3:BM:115:MET:HB3	2.02	0.40
1:BP:37:LEU:HD22	2:BQ:24:MET:SD	2.61	0.40
1:BP:47:ARG:O	1:BP:51:VAL:HG23	2.21	0.40
2:BW:83:ARG:NH2	10:BW:200:CYC:O2A	2.54	0.40
1:BX:89:LEU:HD13	1:BX:133:MET:HE2	2.02	0.40
1:BX:137:ALA:O	1:BX:141:MET:HG2	2.21	0.40
10:BY:200:CYC:HHA	10:BY:200:CYC:HBD1	2.02	0.40
1:BZ:67:SER:HB3	1:BZ:68:PRO:HD2	2.03	0.40
1:CG:36:ARG:NH2	1:CG:152:TYR:OH	2.55	0.40
1:CG:82:LEU:HA	1:CG:85:MET:HG3	2.02	0.40
5:CK:31:VAL:HG11	5:CK:39:GLU:HG2	2.03	0.40
6:CM:373:LEU:HB3	6:CM:374:PRO:HD3	2.04	0.40
6:CM:504:PHE:HA	6:CM:505:PRO:HD3	1.92	0.40
6:CM:578:PHE:O	6:CM:582:LEU:HG	2.21	0.40
6:CM:622:GLN:O	6:CM:626:ILE:HG13	2.22	0.40

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
6:CM:657:GLU:OE2	6:CM:668:ARG:NH2	2.54	0.40
1:CR:92:VAL:HG21	1:CR:153:PHE:HE1	1.86	0.40
1:CT:67:SER:O	1:CT:73:TYR:HB2	2.21	0.40
1:CT:94:TYR:O	1:CT:97:VAL:HG22	2.22	0.40
2:CU:134:LYS:HG3	2:CU:150:GLY:HA2	2.03	0.40
10:CY:200:CYC:CAB	2:DD:74:THR:HA	2.51	0.40
1:DA:71:ASN:C	1:DA:71:ASN:HD22	2.28	0.40
9:DI:223:LEU:HA	9:DI:301:VAL:CG1	2.51	0.40
1:DJ:16:ARG:CA	2:DK:90:ARG:HD2	2.51	0.40
1:DQ:65:ILE:HD12	1:DQ:65:ILE:H	1.86	0.40
9:EA:39:LEU:HD12	9:EA:39:LEU:O	2.21	0.40
2:AB:59:SER:HB3	2:AB:132:ALA:HB2	2.02	0.40
2:AB:112:LEU:CD2	2:AB:160:LEU:HD21	2.51	0.40
1:AC:71:ASN:OD1	1:AC:121:THR:HA	2.21	0.40
1:AH:85:MET:HG2	1:AH:129:SER:OG	2.21	0.40
1:AH:153:PHE:O	1:AH:157:ILE:HG13	2.22	0.40
2:AL:106:GLU:HG3	2:AL:107:ARG:CG	2.47	0.40
1:AN:35:ALA:O	1:AN:39:ILE:HG13	2.21	0.40
2:AO:56:VAL:O	2:AO:61:LEU:HG	2.21	0.40
2:AO:85:LEU:HD22	2:AO:133:ILE:HD12	2.02	0.40
1:AV:6:LYS:HA	1:AV:9:VAL:HG22	2.03	0.40
1:AX:16:ARG:O	2:AY:94:TYR:OH	2.39	0.40
2:AY:101:ALA:HB1	2:AY:155:TYR:CE2	2.56	0.40
6:BD:210:PHE:CZ	6:BD:216:ALA:HB1	2.56	0.40
6:BD:341:ARG:HD2	6:BD:341:ARG:HA	1.89	0.40
6:BD:691:PHE:CZ	1:DC:51:VAL:HG21	2.56	0.40
6:BD:723:VAL:HG13	6:BD:852:THR:HG23	2.03	0.40
1:BI:90:ARG:HH22	2:BN:73:TYR:HH	1.63	0.40
1:BK:64:ASP:OD1	1:BK:64:ASP:N	2.55	0.40
3:BM:37:ILE:HG23	2:BN:24:MET:SD	2.62	0.40
2:BN:123:ILE:O	2:BN:127:VAL:HG23	2.22	0.40
10:BN:200:CYC:HAD2	5:CJ:33:TYR:OH	2.21	0.40
4:BS:54:ARG:HB2	4:BS:137:MET:CE	2.48	0.40
2:BT:36:LEU:HD11	7:CO:85:GLN:HB2	2.04	0.40
2:BY:47:ASN:CB	2:BY:140:LEU:HD21	2.51	0.40
2:CD:53:LYS:HG3	1:CE:118:SER:O	2.21	0.40
2:CD:160:LEU:HD23	2:CD:160:LEU:HA	1.96	0.40
2:CF:3:ASP:H	2:CF:6:THR:HG1	1.70	0.40
5:CK:31:VAL:HG21	5:CK:36:TRP:HE3	1.86	0.40
6:CM:288:GLN:HG2	6:CM:396:PRO:O	2.20	0.40
6:CM:691:PHE:CZ	1:DU:51:VAL:HG21	2.56	0.40

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
6:CM:792:TYR:CE2	1:DQ:13:ALA:HA	2.57	0.40
9:CQ:15:THR:HG23	9:CQ:16:LEU:N	2.37	0.40
9:CQ:253:GLU:OE2	9:CQ:272:LYS:HG3	2.20	0.40
1:CR:54:ALA:CB	1:CR:133:MET:HG3	2.52	0.40
1:CR:78:THR:O	1:CR:82:LEU:HG	2.22	0.40
1:CV:97:VAL:HG21	2:CW:19:LEU:HD11	2.04	0.40
1:CV:134:LYS:HE3	1:CV:150:SER:HA	2.03	0.40
1:CY:8:ILE:HD12	2:CZ:94:TYR:CD1	2.57	0.40
2:CZ:72:MET:HE1	2:CZ:81:CYS:SG	2.62	0.40
1:DA:130:VAL:HB	1:DA:157:ILE:HD11	2.03	0.40
2:DB:63:SER:OG	2:DB:65:VAL:HG22	2.20	0.40
1:DC:107:ILE:HD12	2:DD:13:ASP:OD2	2.20	0.40
8:DG:227:PRO:HG2	1:DQ:78:THR:HG21	2.03	0.40
9:DI:264:PHE:HB3	9:DI:291:LEU:CD1	2.51	0.40
1:DJ:30:VAL:HG11	2:DK:34:GLY:HA3	2.04	0.40
1:DJ:37:LEU:HD11	2:DK:24:MET:SD	2.61	0.40
1:DJ:68:PRO:HA	1:DJ:73:TYR:CG	2.57	0.40
1:DJ:94:TYR:O	1:DJ:97:VAL:HG22	2.22	0.40
1:DJ:141:MET:HE3	1:DJ:145:ASP:C	2.46	0.40
1:DL:142:SER:OG	1:DL:145:ASP:OD2	2.39	0.40
1:DN:47:ARG:HG3	1:DN:48:GLU:N	2.37	0.40
2:DO:56:VAL:CG2	2:DO:85:LEU:HD12	2.52	0.40
2:DO:73:TYR:O	2:DO:74:THR:OG1	2.23	0.40
1:DQ:101:VAL:HG11	1:DQ:155:PHE:CD2	2.57	0.40
2:DR:23:ALA:HA	2:DR:26:LYS:HG2	2.04	0.40
2:DR:85:LEU:HA	2:DR:88:TYR:CD2	2.57	0.40
9:EA:54:ALA:HB3	9:EA:111:TYR:CZ	2.55	0.40
9:EA:254:ARG:HG2	9:EA:270:THR:OG1	2.20	0.40

There are no symmetry-related clashes.

5.3 Torsion angles [i](#)

5.3.1 Protein backbone [i](#)

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

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

Mol	Chain	Analysed	Favoured	Allowed	Outliers	Percentiles	
1	AA	158/161 (98%)	157 (99%)	1 (1%)	0	100	100
1	AC	158/161 (98%)	153 (97%)	5 (3%)	0	100	100
1	AH	158/161 (98%)	157 (99%)	1 (1%)	0	100	100
1	AN	4/161 (2%)	4 (100%)	0	0	100	100
1	AP	21/161 (13%)	21 (100%)	0	0	100	100
1	AX	158/161 (98%)	158 (100%)	0	0	100	100
1	AZ	158/161 (98%)	157 (99%)	1 (1%)	0	100	100
1	BI	158/161 (98%)	158 (100%)	0	0	100	100
1	BK	158/161 (98%)	152 (96%)	6 (4%)	0	100	100
1	BP	158/161 (98%)	157 (99%)	1 (1%)	0	100	100
1	BR	158/161 (98%)	157 (99%)	1 (1%)	0	100	100
1	BV	158/161 (98%)	154 (98%)	4 (2%)	0	100	100
1	BX	109/161 (68%)	107 (98%)	2 (2%)	0	100	100
1	CG	56/161 (35%)	56 (100%)	0	0	100	100
1	CR	158/161 (98%)	157 (99%)	1 (1%)	0	100	100
1	CT	158/161 (98%)	153 (97%)	5 (3%)	0	100	100
1	CV	158/161 (98%)	156 (99%)	2 (1%)	0	100	100
1	CY	158/161 (98%)	156 (99%)	2 (1%)	0	100	100
1	DA	158/161 (98%)	154 (98%)	4 (2%)	0	100	100
1	DC	158/161 (98%)	156 (99%)	2 (1%)	0	100	100
1	DJ	129/161 (80%)	128 (99%)	1 (1%)	0	100	100
1	DL	142/161 (88%)	135 (95%)	6 (4%)	1 (1%)	18	38
2	AB	159/161 (99%)	159 (100%)	0	0	100	100
2	AD	159/161 (99%)	155 (98%)	4 (2%)	0	100	100
2	AF	159/161 (99%)	158 (99%)	1 (1%)	0	100	100
2	AI	151/161 (94%)	146 (97%)	5 (3%)	0	100	100
2	AO	10/161 (6%)	10 (100%)	0	0	100	100
2	AW	43/161 (27%)	42 (98%)	1 (2%)	0	100	100
2	AY	159/161 (99%)	159 (100%)	0	0	100	100
2	BJ	159/161 (99%)	159 (100%)	0	0	100	100
2	BL	159/161 (99%)	155 (98%)	4 (2%)	0	100	100
2	BN	159/161 (99%)	157 (99%)	2 (1%)	0	100	100

Continued on next page...

Continued from previous page...

Mol	Chain	Analysed	Favoured	Allowed	Outliers	Percentiles	
2	BQ	159/161 (99%)	156 (98%)	3 (2%)	0	100	100
2	BT	159/161 (99%)	155 (98%)	4 (2%)	0	100	100
2	BW	159/161 (99%)	157 (99%)	2 (1%)	0	100	100
2	CH	13/161 (8%)	12 (92%)	1 (8%)	0	100	100
2	CS	159/161 (99%)	156 (98%)	3 (2%)	0	100	100
2	CU	159/161 (99%)	154 (97%)	5 (3%)	0	100	100
2	CW	159/161 (99%)	153 (96%)	6 (4%)	0	100	100
2	CZ	159/161 (99%)	147 (92%)	12 (8%)	0	100	100
2	DB	159/161 (99%)	149 (94%)	9 (6%)	1 (1%)	21	42
2	DD	159/161 (99%)	154 (97%)	5 (3%)	0	100	100
2	DK	142/161 (88%)	139 (98%)	3 (2%)	0	100	100
2	DM	152/161 (94%)	149 (98%)	3 (2%)	0	100	100
3	AE	158/161 (98%)	157 (99%)	1 (1%)	0	100	100
3	BM	158/161 (98%)	157 (99%)	1 (1%)	0	100	100
4	BS	167/169 (99%)	161 (96%)	6 (4%)	0	100	100
5	BA	65/67 (97%)	64 (98%)	1 (2%)	0	100	100
5	BB	65/67 (97%)	65 (100%)	0	0	100	100
5	DF	40/67 (60%)	38 (95%)	2 (5%)	0	100	100
6	BD	846/896 (94%)	803 (95%)	43 (5%)	0	100	100
6	CM	753/896 (84%)	703 (93%)	50 (7%)	0	100	100
7	BF	34/121 (28%)	30 (88%)	4 (12%)	0	100	100
7	CO	34/121 (28%)	34 (100%)	0	0	100	100
8	BG	55/249 (22%)	52 (94%)	3 (6%)	0	100	100
8	CP	55/249 (22%)	51 (93%)	4 (7%)	0	100	100
8	DH	12/249 (5%)	9 (75%)	3 (25%)	0	100	100
9	BH	296/317 (93%)	283 (96%)	12 (4%)	1 (0%)	36	58
9	CQ	280/317 (88%)	266 (95%)	13 (5%)	1 (0%)	30	51
9	DI	265/317 (84%)	252 (95%)	12 (4%)	1 (0%)	30	51
9	EA	5/317 (2%)	4 (80%)	1 (20%)	0	100	100
All	All	9332/11825 (79%)	9053 (97%)	274 (3%)	5 (0%)	49	70

All (5) Ramachandran outliers are listed below:

Mol	Chain	Res	Type
9	BH	176	VAL
9	CQ	176	VAL
1	DL	143	SER
2	DB	75	THR
9	DI	298	ILE

5.3.2 Protein sidechains [i](#)

There are no protein residues with a non-rotameric sidechain to report in this entry.

5.3.3 RNA [i](#)

There are no RNA molecules in this entry.

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

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

5.5 Carbohydrates [i](#)

There are no oligosaccharides in this entry.

5.6 Ligand geometry [i](#)

76 ligands are modelled in this entry.

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

Mol	Type	Chain	Res	Link	Bond lengths			Bond angles		
					Counts	RMSZ	$\# Z > 2$	Counts	RMSZ	$\# Z > 2$
10	CYC	BX	200	1	46,46,46	0.23	0	63,67,67	0.70	3 (4%)
10	CYC	AH	200	1	46,46,46	0.23	0	63,67,67	0.66	3 (4%)
10	CYC	BD	902	6	46,46,46	0.24	0	63,67,67	0.71	2 (3%)
10	CYC	AL	200	2	46,46,46	0.23	0	63,67,67	0.64	2 (3%)

Mol	Type	Chain	Res	Link	Bond lengths			Bond angles		
					Counts	RMSZ	# Z > 2	Counts	RMSZ	# Z > 2
10	CYC	CC	200	1	46,46,46	0.19	0	63,67,67	0.54	2 (3%)
10	CYC	DN	200	1	46,46,46	0.28	0	63,67,67	0.71	2 (3%)
10	CYC	BV	200	1	46,46,46	0.20	0	63,67,67	0.67	2 (3%)
10	CYC	BP	200	1	46,46,46	0.24	0	63,67,67	0.70	3 (4%)
10	CYC	AP	200	1	46,46,46	0.21	0	63,67,67	0.66	3 (4%)
10	CYC	BK	200	1	46,46,46	0.21	0	63,67,67	0.42	1 (1%)
10	CYC	BL	200	2	46,46,46	0.24	0	63,67,67	0.82	3 (4%)
10	CYC	AX	200	1	46,46,46	0.22	0	63,67,67	0.70	3 (4%)
10	CYC	BI	200	1	46,46,46	0.24	0	63,67,67	0.47	1 (1%)
10	CYC	DS	200	1	46,46,46	0.26	0	63,67,67	0.75	2 (3%)
10	CYC	DT	200	2	46,46,46	0.27	0	63,67,67	0.78	4 (6%)
10	CYC	DC	200	1	46,46,46	0.31	0	63,67,67	0.73	2 (3%)
10	CYC	CD	200	2	46,46,46	0.29	0	63,67,67	1.09	3 (4%)
10	CYC	AF	200	2	46,46,46	0.26	0	63,67,67	0.87	3 (4%)
10	CYC	AR	200	-	46,46,46	0.21	0	63,67,67	0.58	1 (1%)
10	CYC	CE	200	1	46,46,46	0.22	0	63,67,67	0.72	3 (4%)
10	CYC	AE	200	3	46,46,46	0.21	0	63,67,67	0.62	2 (3%)
10	CYC	AW	200	2	46,46,46	0.24	0	63,67,67	0.85	2 (3%)
10	CYC	BW	200	2	46,46,46	0.21	0	63,67,67	0.75	2 (3%)
10	CYC	BS	200	4	46,46,46	0.28	0	63,67,67	0.87	3 (4%)
10	CYC	AD	200	2	46,46,46	0.23	0	63,67,67	0.63	2 (3%)
10	CYC	CS	200	2	46,46,46	0.22	0	63,67,67	0.67	2 (3%)
10	CYC	DK	200	2	46,46,46	0.19	0	63,67,67	0.57	2 (3%)
10	CYC	DM	200	2	46,46,46	0.24	0	63,67,67	0.73	3 (4%)
10	CYC	BY	200	2	46,46,46	0.27	0	63,67,67	0.86	3 (4%)
10	CYC	AI	200	2	46,46,46	0.27	0	63,67,67	0.81	3 (4%)
10	CYC	AV	200	1	46,46,46	0.22	0	63,67,67	0.67	2 (3%)
10	CYC	BR	200	1	46,46,46	0.22	0	63,67,67	0.64	3 (4%)
10	CYC	BQ	200	2	46,46,46	0.24	0	63,67,67	0.91	3 (4%)
10	CYC	AA	200	1	46,46,46	0.21	0	63,67,67	0.61	3 (4%)
10	CYC	AQ	200	2	46,46,46	0.26	0	63,67,67	0.84	3 (4%)
10	CYC	CG	200	1	46,46,46	0.22	0	63,67,67	0.68	3 (4%)
10	CYC	CM	901	2	46,46,46	0.29	0	63,67,67	0.97	3 (4%)
10	CYC	CZ	200	2	46,46,46	0.23	0	63,67,67	0.79	2 (3%)
11	45D	BH	400	-	43,43,43	0.97	2 (4%)	54,60,60	1.57	12 (22%)

Mol	Type	Chain	Res	Link	Bond lengths			Bond angles		
					Counts	RMSZ	# Z > 2	Counts	RMSZ	# Z > 2
11	45D	CQ	400	-	43,43,43	0.97	1 (2%)	54,60,60	1.50	14 (25%)
10	CYC	DJ	200	1	46,46,46	0.23	0	63,67,67	0.46	1 (1%)
10	CYC	DL	200	1	46,46,46	0.22	0	63,67,67	0.72	2 (3%)
10	CYC	DQ	200	1	46,46,46	0.23	0	63,67,67	0.48	1 (1%)
10	CYC	CH	200	2	46,46,46	0.21	0	63,67,67	0.64	2 (3%)
10	CYC	AC	200	1	46,46,46	0.20	0	63,67,67	0.55	1 (1%)
10	CYC	CT	200	1	46,46,46	0.21	0	63,67,67	0.76	3 (4%)
10	CYC	CY	200	1	46,46,46	0.20	0	63,67,67	0.65	2 (3%)
10	CYC	AU	200	2	46,46,46	0.27	0	63,67,67	1.08	3 (4%)
10	CYC	AO	200	2	46,46,46	0.23	0	63,67,67	0.82	3 (4%)
10	CYC	AZ	200	1	46,46,46	0.20	0	63,67,67	0.52	2 (3%)
10	CYC	BT	200	2	46,46,46	0.26	0	63,67,67	0.78	2 (3%)
10	CYC	AB	200	2	46,46,46	0.20	0	63,67,67	0.65	2 (3%)
10	CYC	AY	200	2	46,46,46	0.20	0	63,67,67	0.58	2 (3%)
10	CYC	BZ	200	1	46,46,46	0.22	0	63,67,67	0.70	3 (4%)
10	CYC	AJ	200	1	46,46,46	0.21	0	63,67,67	0.66	3 (4%)
10	CYC	CV	200	1	46,46,46	0.25	0	63,67,67	0.61	2 (3%)
10	CYC	CR	200	1	46,46,46	0.28	0	63,67,67	0.73	3 (4%)
10	CYC	DD	200	2	46,46,46	0.20	0	63,67,67	0.56	2 (3%)
10	CYC	DU	200	1	46,46,46	0.31	0	63,67,67	0.74	2 (3%)
10	CYC	DO	200	2	46,46,46	0.25	0	63,67,67	0.85	3 (4%)
10	CYC	AN	200	1	46,46,46	0.20	0	63,67,67	0.72	3 (4%)
10	CYC	AK	200	4	46,46,46	0.22	0	63,67,67	0.79	3 (4%)
10	CYC	DV	200	2	46,46,46	0.21	0	63,67,67	0.55	2 (3%)
10	CYC	DB	200	2	46,46,46	0.26	0	63,67,67	0.84	4 (6%)
11	45D	DI	400	-	43,43,43	0.96	2 (4%)	54,60,60	1.50	12 (22%)
10	CYC	CU	200	2	46,46,46	0.26	0	63,67,67	0.79	3 (4%)
10	CYC	DR	200	2	46,46,46	0.28	0	63,67,67	0.93	3 (4%)
10	CYC	CW	200	2	46,46,46	0.28	0	63,67,67	0.93	3 (4%)
10	CYC	BJ	200	2	46,46,46	0.20	0	63,67,67	0.61	2 (3%)
10	CYC	DA	200	1	46,46,46	0.22	0	63,67,67	0.75	2 (3%)
10	CYC	BN	200	2	46,46,46	0.26	0	63,67,67	0.88	3 (4%)
11	45D	EA	400	-	43,43,43	1.00	3 (6%)	54,60,60	1.49	10 (18%)
10	CYC	CM	902	6	46,46,46	0.25	0	63,67,67	0.76	2 (3%)
10	CYC	BD	901	2	46,46,46	0.25	0	63,67,67	0.87	3 (4%)

Mol	Type	Chain	Res	Link	Bond lengths			Bond angles		
					Counts	RMSZ	# Z > 2	Counts	RMSZ	# Z > 2
10	CYC	BM	200	3	46,46,46	0.23	0	63,67,67	0.73	3 (4%)
10	CYC	CF	200	2	46,46,46	0.25	0	63,67,67	0.85	4 (6%)

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
10	CYC	BX	200	1	-	9/26/74/74	0/4/4/4
10	CYC	AH	200	1	-	13/26/74/74	0/4/4/4
10	CYC	BD	902	6	-	8/26/74/74	0/4/4/4
10	CYC	AL	200	2	-	13/26/74/74	0/4/4/4
10	CYC	CC	200	1	-	8/26/74/74	0/4/4/4
10	CYC	DN	200	1	-	9/26/74/74	0/4/4/4
10	CYC	BV	200	1	-	13/26/74/74	0/4/4/4
10	CYC	BP	200	1	-	10/26/74/74	0/4/4/4
10	CYC	AP	200	1	-	9/26/74/74	0/4/4/4
10	CYC	BK	200	1	-	8/26/74/74	0/4/4/4
10	CYC	BL	200	2	-	14/26/74/74	0/4/4/4
10	CYC	AX	200	1	-	10/26/74/74	0/4/4/4
10	CYC	BI	200	1	-	12/26/74/74	0/4/4/4
10	CYC	DS	200	1	-	7/26/74/74	0/4/4/4
10	CYC	DT	200	2	-	12/26/74/74	0/4/4/4
10	CYC	DC	200	1	-	12/26/74/74	0/4/4/4
10	CYC	CD	200	2	-	10/26/74/74	0/4/4/4
10	CYC	AF	200	2	-	7/26/74/74	0/4/4/4
10	CYC	AR	200	-	-	13/26/74/74	0/4/4/4
10	CYC	CE	200	1	-	11/26/74/74	0/4/4/4
10	CYC	AE	200	3	-	12/26/74/74	0/4/4/4
10	CYC	AW	200	2	-	11/26/74/74	0/4/4/4
10	CYC	BW	200	2	-	9/26/74/74	0/4/4/4
10	CYC	BS	200	4	-	9/26/74/74	0/4/4/4
10	CYC	AD	200	2	-	12/26/74/74	0/4/4/4
10	CYC	CS	200	2	-	9/26/74/74	0/4/4/4
10	CYC	DK	200	2	-	9/26/74/74	0/4/4/4

Continued on next page...

Continued from previous page...

Mol	Type	Chain	Res	Link	Chirals	Torsions	Rings
10	CYC	DM	200	2	-	9/26/74/74	0/4/4/4
10	CYC	BY	200	2	-	11/26/74/74	0/4/4/4
10	CYC	AI	200	2	-	7/26/74/74	0/4/4/4
10	CYC	AV	200	1	-	8/26/74/74	0/4/4/4
10	CYC	BR	200	1	-	10/26/74/74	0/4/4/4
10	CYC	BQ	200	2	-	9/26/74/74	0/4/4/4
10	CYC	AA	200	1	-	10/26/74/74	0/4/4/4
10	CYC	AQ	200	2	-	13/26/74/74	0/4/4/4
10	CYC	CG	200	1	-	10/26/74/74	0/4/4/4
10	CYC	CM	901	2	-	13/26/74/74	0/4/4/4
10	CYC	CZ	200	2	-	13/26/74/74	0/4/4/4
11	45D	BH	400	-	-	16/29/69/69	0/2/2/2
11	45D	CQ	400	-	-	15/29/69/69	0/2/2/2
10	CYC	DJ	200	1	-	11/26/74/74	0/4/4/4
10	CYC	DL	200	1	-	11/26/74/74	0/4/4/4
10	CYC	DQ	200	1	-	16/26/74/74	0/4/4/4
10	CYC	CH	200	2	-	11/26/74/74	0/4/4/4
10	CYC	AC	200	1	-	10/26/74/74	0/4/4/4
10	CYC	CT	200	1	-	13/26/74/74	0/4/4/4
10	CYC	CY	200	1	-	13/26/74/74	0/4/4/4
10	CYC	AU	200	2	-	7/26/74/74	0/4/4/4
10	CYC	AO	200	2	-	8/26/74/74	0/4/4/4
10	CYC	AZ	200	1	-	12/26/74/74	0/4/4/4
10	CYC	BT	200	2	-	15/26/74/74	0/4/4/4
10	CYC	AB	200	2	-	10/26/74/74	0/4/4/4
10	CYC	AY	200	2	-	12/26/74/74	0/4/4/4
10	CYC	BZ	200	1	-	10/26/74/74	0/4/4/4
10	CYC	AJ	200	1	-	10/26/74/74	0/4/4/4
10	CYC	CV	200	1	-	5/26/74/74	0/4/4/4
10	CYC	CR	200	1	-	10/26/74/74	0/4/4/4
10	CYC	DD	200	2	-	7/26/74/74	0/4/4/4
10	CYC	DU	200	1	-	10/26/74/74	0/4/4/4
10	CYC	DO	200	2	-	10/26/74/74	0/4/4/4
10	CYC	AN	200	1	-	11/26/74/74	0/4/4/4

Continued on next page...

Continued from previous page...

Mol	Type	Chain	Res	Link	Chirals	Torsions	Rings
10	CYC	AK	200	4	-	10/26/74/74	0/4/4/4
10	CYC	DV	200	2	-	11/26/74/74	0/4/4/4
10	CYC	DB	200	2	-	11/26/74/74	0/4/4/4
11	45D	DI	400	-	-	15/29/69/69	0/2/2/2
10	CYC	CU	200	2	-	10/26/74/74	0/4/4/4
10	CYC	DR	200	2	-	11/26/74/74	0/4/4/4
10	CYC	CW	200	2	-	12/26/74/74	0/4/4/4
10	CYC	BJ	200	2	-	8/26/74/74	0/4/4/4
10	CYC	DA	200	1	-	8/26/74/74	0/4/4/4
10	CYC	BN	200	2	-	8/26/74/74	0/4/4/4
11	45D	EA	400	-	-	16/29/69/69	0/2/2/2
10	CYC	CM	902	6	-	7/26/74/74	0/4/4/4
10	CYC	BD	901	2	-	10/26/74/74	0/4/4/4
10	CYC	BM	200	3	-	8/26/74/74	0/4/4/4
10	CYC	CF	200	2	-	10/26/74/74	0/4/4/4

All (8) bond length outliers are listed below:

Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
11	EA	400	45D	C34-C36	3.69	1.53	1.46
11	DI	400	45D	C34-C36	3.51	1.53	1.46
11	BH	400	45D	C34-C36	3.50	1.53	1.46
11	CQ	400	45D	C34-C36	3.47	1.53	1.46
11	BH	400	45D	C09-C17	2.17	1.53	1.50
11	EA	400	45D	C27-C25	2.06	1.55	1.50
11	DI	400	45D	C09-C17	2.05	1.53	1.50
11	EA	400	45D	C24-C20	2.00	1.39	1.33

All (226) bond angle outliers are listed below:

Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
10	AU	200	CYC	C1D-CHD-C4C	5.73	137.54	127.76
10	CD	200	CYC	C1D-CHD-C4C	5.72	137.52	127.76
10	DR	200	CYC	C1D-CHD-C4C	5.66	137.42	127.76
10	CW	200	CYC	C1B-CHB-C4A	5.34	141.17	128.06
10	CM	901	CYC	C1B-CHB-C4A	5.23	140.90	128.06
10	AW	200	CYC	C1D-CHD-C4C	5.19	136.63	127.76
10	CD	200	CYC	C1B-CHB-C4A	5.18	140.78	128.06
10	BL	200	CYC	C1D-CHD-C4C	5.10	136.46	127.76

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
10	AU	200	CYC	C1B-CHB-C4A	5.03	140.41	128.06
10	BQ	200	CYC	C1B-CHB-C4A	4.94	140.18	128.06
10	CZ	200	CYC	C1D-CHD-C4C	4.87	136.07	127.76
10	BD	901	CYC	C1B-CHB-C4A	4.83	139.93	128.06
10	BT	200	CYC	C1D-CHD-C4C	4.83	136.02	127.76
10	DO	200	CYC	C1B-CHB-C4A	4.83	139.90	128.06
10	BS	200	CYC	C1D-CHD-C4C	4.79	135.94	127.76
10	AO	200	CYC	C1D-CHD-C4C	4.72	135.82	127.76
10	CM	902	CYC	C1B-CHB-C4A	4.67	139.51	128.06
10	AK	200	CYC	C1D-CHD-C4C	4.55	135.53	127.76
10	BN	200	CYC	C1D-CHD-C4C	4.49	135.42	127.76
10	BW	200	CYC	C1D-CHD-C4C	4.43	135.33	127.76
10	CU	200	CYC	C1D-CHD-C4C	4.39	135.25	127.76
10	DU	200	CYC	C1B-CHB-C4A	4.34	138.72	128.06
10	AF	200	CYC	C1D-CHD-C4C	4.33	135.16	127.76
10	DC	200	CYC	C1B-CHB-C4A	4.25	138.48	128.06
10	BY	200	CYC	C1B-CHB-C4A	4.24	138.47	128.06
10	DN	200	CYC	C1B-CHB-C4A	4.20	138.36	128.06
10	CF	200	CYC	C1D-CHD-C4C	4.16	134.86	127.76
10	AQ	200	CYC	C1B-CHB-C4A	4.13	138.20	128.06
10	CM	901	CYC	C1D-CHD-C4C	4.10	134.77	127.76
10	AI	200	CYC	C1B-CHB-C4A	4.03	137.94	128.06
10	CS	200	CYC	C1D-CHD-C4C	4.01	134.61	127.76
10	DA	200	CYC	C1D-CHD-C4C	3.99	134.58	127.76
10	CY	200	CYC	C1D-CHD-C4C	3.99	134.57	127.76
10	BD	902	CYC	C1B-CHB-C4A	3.94	137.73	128.06
10	DB	200	CYC	C1D-CHD-C4C	3.92	134.45	127.76
10	DS	200	CYC	C1B-CHB-C4A	3.90	137.63	128.06
10	AF	200	CYC	C1B-CHB-C4A	3.85	137.50	128.06
10	AN	200	CYC	C1D-CHD-C4C	3.84	134.31	127.76
10	BY	200	CYC	C1D-CHD-C4C	3.83	134.30	127.76
10	BV	200	CYC	C1D-CHD-C4C	3.78	134.22	127.76
10	AV	200	CYC	C1D-CHD-C4C	3.77	134.21	127.76
10	BN	200	CYC	C1B-CHB-C4A	3.75	137.26	128.06
10	AB	200	CYC	C1D-CHD-C4C	3.74	134.15	127.76
10	BM	200	CYC	C1D-CHD-C4C	3.72	134.11	127.76
11	EA	400	45D	C05-C03-C07	3.71	115.83	110.44
10	AQ	200	CYC	C1D-CHD-C4C	3.70	134.08	127.76
10	DM	200	CYC	C1D-CHD-C4C	3.68	134.04	127.76
11	BH	400	45D	C32-C30-C26	-3.66	122.14	127.28
10	AI	200	CYC	C1D-CHD-C4C	3.60	133.91	127.76
10	AL	200	CYC	C1D-CHD-C4C	3.56	133.85	127.76

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
10	AX	200	CYC	C1D-CHD-C4C	3.54	133.81	127.76
10	CG	200	CYC	C1D-CHD-C4C	3.53	133.78	127.76
10	CR	200	CYC	C1B-CHB-C4A	3.52	136.70	128.06
10	CH	200	CYC	C1D-CHD-C4C	3.51	133.75	127.76
11	DI	400	45D	C19-C23-C25	-3.48	121.08	126.23
10	BJ	200	CYC	C1D-CHD-C4C	3.45	133.65	127.76
10	BP	200	CYC	C1D-CHD-C4C	3.44	133.63	127.76
10	BQ	200	CYC	C1D-CHD-C4C	3.41	133.59	127.76
10	BZ	200	CYC	C1D-CHD-C4C	3.40	133.57	127.76
10	BS	200	CYC	C1B-CHB-C4A	3.39	136.38	128.06
11	BH	400	45D	C19-C23-C25	-3.39	121.22	126.23
10	CE	200	CYC	C1D-CHD-C4C	3.38	133.54	127.76
10	BD	901	CYC	C1D-CHD-C4C	3.38	133.54	127.76
10	AD	200	CYC	C1D-CHD-C4C	3.37	133.51	127.76
10	AH	200	CYC	C1D-CHD-C4C	3.36	133.50	127.76
10	DK	200	CYC	C1D-CHD-C4C	3.30	133.40	127.76
10	CW	200	CYC	C1D-CHD-C4C	3.30	133.40	127.76
10	BX	200	CYC	C1D-CHD-C4C	3.28	133.36	127.76
10	CV	200	CYC	C1B-CHB-C4A	3.25	136.04	128.06
10	BR	200	CYC	C1D-CHD-C4C	3.23	133.28	127.76
10	DT	200	CYC	C1B-CHB-C4A	3.21	135.92	128.06
11	EA	400	45D	C19-C23-C25	-3.14	121.59	126.23
10	AJ	200	CYC	C1D-CHD-C4C	3.13	133.10	127.76
11	BH	400	45D	C32-C34-C36	-3.11	117.84	126.36
10	AP	200	CYC	C1D-CHD-C4C	3.08	133.02	127.76
10	DD	200	CYC	C1D-CHD-C4C	3.07	133.01	127.76
11	CQ	400	45D	C32-C30-C26	-3.07	122.97	127.28
11	CQ	400	45D	C23-C19-C07	-3.01	118.96	127.00
11	EA	400	45D	C32-C30-C26	-3.00	123.07	127.28
10	DV	200	CYC	C1D-CHD-C4C	2.99	132.87	127.76
10	AE	200	CYC	C1D-CHD-C4C	2.99	132.86	127.76
11	CQ	400	45D	C32-C34-C36	-2.98	118.18	126.36
10	DB	200	CYC	CAC-C3C-C4C	-2.98	105.02	112.67
10	CE	200	CYC	C1B-CHB-C4A	2.97	135.35	128.06
10	AY	200	CYC	C1D-CHD-C4C	2.97	132.83	127.76
11	CQ	400	45D	C19-C23-C25	-2.96	121.85	126.23
10	AA	200	CYC	C1D-CHD-C4C	2.96	132.81	127.76
10	CT	200	CYC	C1B-CHB-C4A	2.90	135.19	128.06
10	DO	200	CYC	C1D-CHD-C4C	2.90	132.72	127.76
10	AC	200	CYC	C1D-CHD-C4C	2.90	132.71	127.76
11	EA	400	45D	C32-C34-C36	-2.88	118.48	126.36
10	BM	200	CYC	C1B-CHB-C4A	2.87	135.11	128.06

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
11	BH	400	45D	C09-C17-C15	2.86	121.21	118.64
10	BX	200	CYC	C1B-CHB-C4A	2.84	135.03	128.06
11	BH	400	45D	C23-C19-C07	-2.83	119.43	127.00
11	DI	400	45D	C22-C16-C18	2.82	119.77	115.49
10	DC	200	CYC	C2C-C3C-C4C	2.80	105.53	101.34
11	DI	400	45D	C23-C19-C07	-2.79	119.54	127.00
10	DB	200	CYC	C2C-C3C-C4C	2.79	105.52	101.34
10	DT	200	CYC	C2C-C3C-C4C	2.79	105.51	101.34
10	DA	200	CYC	C1B-CHB-C4A	2.77	134.87	128.06
10	DU	200	CYC	C2C-C3C-C4C	2.76	105.47	101.34
10	CU	200	CYC	C2C-C3C-C4C	2.73	105.43	101.34
10	CR	200	CYC	C2C-C3C-C4C	2.71	105.40	101.34
10	DR	200	CYC	C2C-C3C-C4C	2.71	105.40	101.34
10	DT	200	CYC	C1D-CHD-C4C	2.70	132.38	127.76
11	EA	400	45D	C22-C16-C08	-2.67	119.81	124.11
11	DI	400	45D	C22-C16-C08	-2.67	119.82	124.11
11	DI	400	45D	C21-C15-C07	-2.66	119.83	124.11
10	AX	200	CYC	C1B-CHB-C4A	2.65	134.56	128.06
10	DS	200	CYC	C2C-C3C-C4C	2.64	105.30	101.34
10	DM	200	CYC	C2C-C3C-C4C	2.64	105.29	101.34
10	BP	200	CYC	C1B-CHB-C4A	2.63	134.52	128.06
10	CT	200	CYC	C2C-C3C-C4C	2.63	105.28	101.34
10	BZ	200	CYC	C1B-CHB-C4A	2.61	134.47	128.06
11	EA	400	45D	C23-C19-C07	-2.61	120.03	127.00
11	DI	400	45D	C32-C30-C26	-2.60	123.63	127.28
11	DI	400	45D	C32-C34-C36	-2.59	119.26	126.36
10	AR	200	CYC	C1D-CHD-C4C	2.54	132.10	127.76
10	DR	200	CYC	C1B-CHB-C4A	2.53	134.28	128.06
10	CG	200	CYC	C1B-CHB-C4A	2.53	134.27	128.06
10	CC	200	CYC	C1D-CHD-C4C	2.52	132.06	127.76
11	CQ	400	45D	C20-C24-C26	-2.51	122.53	126.23
10	DL	200	CYC	C4D-CHA-C1A	2.49	133.67	128.22
11	BH	400	45D	C21-C15-C07	-2.49	120.11	124.11
10	DL	200	CYC	C2C-C3C-C4C	2.49	105.06	101.34
10	AO	200	CYC	C1B-CHB-C4A	2.49	134.16	128.06
10	CV	200	CYC	C2C-C3C-C4C	2.44	105.00	101.34
10	BP	200	CYC	C2C-C3C-C4C	2.43	104.97	101.34
10	AP	200	CYC	C1B-CHB-C4A	2.42	133.99	128.06
10	CU	200	CYC	CAC-C3C-C4C	-2.41	106.48	112.67
10	AJ	200	CYC	C1B-CHB-C4A	2.40	133.96	128.06
10	BT	200	CYC	C2C-C3C-C4C	2.40	104.93	101.34
11	CQ	400	45D	C05-C03-C07	2.40	113.92	110.44

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
10	AH	200	CYC	C2C-C3C-C4C	2.39	104.92	101.34
10	CT	200	CYC	C4D-CHA-C1A	2.38	133.43	128.22
10	CZ	200	CYC	C2C-C3C-C4C	2.37	104.89	101.34
10	BR	200	CYC	C2C-C3C-C4C	2.36	104.88	101.34
10	CF	200	CYC	CAA-C2A-C1A	2.36	129.16	125.02
10	DM	200	CYC	CAC-C3C-C4C	-2.36	106.62	112.67
11	BH	400	45D	C28-C26-C24	2.35	121.68	118.09
11	BH	400	45D	C22-C16-C08	-2.35	120.33	124.11
10	AO	200	CYC	C2C-C3C-C4C	2.35	104.86	101.34
11	BH	400	45D	C24-C20-C08	-2.34	120.74	127.00
10	BD	902	CYC	C2C-C3C-C4C	2.34	104.84	101.34
10	DQ	200	CYC	C2C-C3C-C4C	2.33	104.83	101.34
10	CM	902	CYC	C2C-C3C-C4C	2.31	104.80	101.34
11	DI	400	45D	C24-C20-C08	-2.31	120.83	127.00
10	DN	200	CYC	C2C-C3C-C4C	2.31	104.80	101.34
10	BS	200	CYC	C2C-C3C-C4C	2.31	104.80	101.34
10	BW	200	CYC	C2C-C3C-C4C	2.30	104.79	101.34
10	BZ	200	CYC	C2C-C3C-C4C	2.30	104.79	101.34
10	BQ	200	CYC	C2C-C3C-C4C	2.30	104.78	101.34
10	AL	200	CYC	C2C-C3C-C4C	2.29	104.77	101.34
10	AZ	200	CYC	C1D-CHD-C4C	2.29	131.67	127.76
11	CQ	400	45D	C28-C26-C24	2.29	121.58	118.09
10	AK	200	CYC	C1B-CHB-C4A	2.28	133.65	128.06
10	AH	200	CYC	C1B-CHB-C4A	2.27	133.64	128.06
10	AK	200	CYC	C2C-C3C-C4C	2.27	104.74	101.34
11	DI	400	45D	C21-C15-C17	2.27	118.93	115.49
11	DI	400	45D	C20-C24-C26	-2.25	122.90	126.23
10	BK	200	CYC	C2C-C3C-C4C	2.25	104.71	101.34
10	AJ	200	CYC	C2C-C3C-C4C	2.25	104.70	101.34
10	AW	200	CYC	C2C-C3C-C4C	2.24	104.69	101.34
11	EA	400	45D	C28-C26-C24	2.24	121.51	118.09
10	AF	200	CYC	C2C-C3C-C4C	2.23	104.68	101.34
10	AP	200	CYC	C2C-C3C-C4C	2.22	104.67	101.34
10	CH	200	CYC	C2C-C3C-C4C	2.22	104.66	101.34
10	AQ	200	CYC	C2C-C3C-C4C	2.21	104.65	101.34
10	BY	200	CYC	C2C-C3C-C4C	2.21	104.65	101.34
10	DT	200	CYC	CAC-C3C-C4C	-2.21	107.00	112.67
10	BX	200	CYC	C2C-C3C-C4C	2.21	104.65	101.34
10	CD	200	CYC	C2C-C3C-C4C	2.20	104.63	101.34
10	CW	200	CYC	C2C-C3C-C4C	2.20	104.63	101.34
11	CQ	400	45D	C21-C15-C17	2.20	118.82	115.49
11	BH	400	45D	C05-C03-C07	2.20	113.63	110.44

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
10	BN	200	CYC	C2C-C3C-C4C	2.19	104.62	101.34
11	CQ	400	45D	C22-C16-C08	-2.19	120.59	124.11
10	CS	200	CYC	C2C-C3C-C4C	2.18	104.61	101.34
10	BI	200	CYC	C2C-C3C-C4C	2.18	104.61	101.34
10	DJ	200	CYC	C2C-C3C-C4C	2.18	104.60	101.34
10	AX	200	CYC	C2C-C3C-C4C	2.18	104.60	101.34
10	AY	200	CYC	C2C-C3C-C4C	2.18	104.60	101.34
10	AD	200	CYC	C2C-C3C-C4C	2.17	104.59	101.34
10	CF	200	CYC	C2C-C3C-C4C	2.17	104.59	101.34
11	EA	400	45D	C21-C15-C07	-2.17	120.62	124.11
10	AZ	200	CYC	C2C-C3C-C4C	2.17	104.59	101.34
10	BL	200	CYC	C2C-C3C-C4C	2.17	104.58	101.34
10	CR	200	CYC	C1D-CHD-C4C	2.16	131.46	127.76
11	CQ	400	45D	C06-C04-C08	2.16	113.58	110.44
10	AE	200	CYC	C2C-C3C-C4C	2.16	104.58	101.34
11	CQ	400	45D	C40-C36-C34	2.15	121.37	118.09
10	CE	200	CYC	C2C-C3C-C4C	2.15	104.55	101.34
11	CQ	400	45D	C24-C20-C08	-2.14	121.28	127.00
10	DD	200	CYC	C2C-C3C-C4C	2.14	104.54	101.34
10	CM	901	CYC	C2C-C3C-C4C	2.13	104.53	101.34
10	AB	200	CYC	C2C-C3C-C4C	2.13	104.53	101.34
11	BH	400	45D	C21-C15-C17	2.13	118.72	115.49
10	CG	200	CYC	C2C-C3C-C4C	2.13	104.53	101.34
10	AA	200	CYC	C1B-CHB-C4A	2.12	133.27	128.06
10	BJ	200	CYC	C2C-C3C-C4C	2.12	104.52	101.34
10	DB	200	CYC	C1B-CHB-C4A	2.12	133.25	128.06
10	BL	200	CYC	C1B-CHB-C4A	2.11	133.24	128.06
10	CC	200	CYC	C2C-C3C-C4C	2.11	104.49	101.34
10	DO	200	CYC	C2C-C3C-C4C	2.09	104.47	101.34
10	CF	200	CYC	C1B-CHB-C4A	2.09	133.18	128.06
10	AU	200	CYC	C2C-C3C-C4C	2.08	104.46	101.34
10	CY	200	CYC	C2C-C3C-C4C	2.08	104.46	101.34
11	EA	400	45D	C22-C16-C18	2.08	118.64	115.49
10	DV	200	CYC	C2C-C3C-C4C	2.07	104.44	101.34
10	AV	200	CYC	C2C-C3C-C4C	2.07	104.44	101.34
10	BM	200	CYC	C2C-C3C-C4C	2.06	104.43	101.34
10	AN	200	CYC	C1B-CHB-C4A	2.06	133.12	128.06
11	CQ	400	45D	C41-C42-C38	-2.05	119.32	123.52
11	DI	400	45D	C28-C26-C24	2.04	121.20	118.09
10	BR	200	CYC	C1B-CHB-C4A	2.04	133.06	128.06
10	AA	200	CYC	C2C-C3C-C4C	2.04	104.39	101.34
11	BH	400	45D	C42-C41-C37	-2.04	119.35	123.52

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
11	CQ	400	45D	C22-C16-C18	2.04	118.58	115.49
10	AN	200	CYC	C2C-C3C-C4C	2.03	104.38	101.34
10	BD	901	CYC	C2C-C3C-C4C	2.02	104.37	101.34
11	EA	400	45D	C24-C20-C08	-2.02	121.60	127.00
11	DI	400	45D	C41-C42-C38	-2.02	119.39	123.52
10	AI	200	CYC	C2C-C3C-C4C	2.01	104.34	101.34
10	DK	200	CYC	C2C-C3C-C4C	2.01	104.34	101.34
10	BV	200	CYC	C1B-CHB-C4A	2.00	132.97	128.06

There are no chirality outliers.

All (800) torsion outliers are listed below:

Mol	Chain	Res	Type	Atoms
10	AA	200	CYC	NA-C4A-CHB-C1B
10	AA	200	CYC	C3A-C4A-CHB-C1B
10	AA	200	CYC	NC-C4C-CHD-C1D
10	AA	200	CYC	C3C-C4C-CHD-C1D
10	AB	200	CYC	NA-C4A-CHB-C1B
10	AB	200	CYC	C3A-C4A-CHB-C1B
10	AB	200	CYC	NC-C4C-CHD-C1D
10	AB	200	CYC	C3C-C4C-CHD-C1D
10	AC	200	CYC	NA-C4A-CHB-C1B
10	AC	200	CYC	C3A-C4A-CHB-C1B
10	AC	200	CYC	C4B-C3B-CAB-CBB
10	AC	200	CYC	NC-C4C-CHD-C1D
10	AD	200	CYC	NA-C4A-CHB-C1B
10	AD	200	CYC	C3A-C4A-CHB-C1B
10	AD	200	CYC	NC-C4C-CHD-C1D
10	AD	200	CYC	C3C-C4C-CHD-C1D
10	AE	200	CYC	NA-C4A-CHB-C1B
10	AE	200	CYC	C3A-C4A-CHB-C1B
10	AE	200	CYC	NC-C4C-CHD-C1D
10	AE	200	CYC	C3C-C4C-CHD-C1D
10	AF	200	CYC	NA-C4A-CHB-C1B
10	AH	200	CYC	NA-C4A-CHB-C1B
10	AH	200	CYC	C3A-C4A-CHB-C1B
10	AH	200	CYC	NC-C4C-CHD-C1D
10	AH	200	CYC	C3C-C4C-CHD-C1D
10	AI	200	CYC	NA-C4A-CHB-C1B
10	AJ	200	CYC	NA-C4A-CHB-C1B
10	AJ	200	CYC	C3A-C4A-CHB-C1B
10	AK	200	CYC	NA-C4A-CHB-C1B

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type	Atoms
10	AK	200	CYC	C3A-C4A-CHB-C1B
10	AL	200	CYC	NA-C4A-CHB-C1B
10	AL	200	CYC	C3A-C4A-CHB-C1B
10	AN	200	CYC	NA-C4A-CHB-C1B
10	AN	200	CYC	C3A-C4A-CHB-C1B
10	AN	200	CYC	NC-C4C-CHD-C1D
10	AN	200	CYC	C3C-C4C-CHD-C1D
10	AO	200	CYC	NA-C4A-CHB-C1B
10	AO	200	CYC	C3A-C4A-CHB-C1B
10	AP	200	CYC	NA-C4A-CHB-C1B
10	AP	200	CYC	C3A-C4A-CHB-C1B
10	AQ	200	CYC	NA-C4A-CHB-C1B
10	AR	200	CYC	NA-C4A-CHB-C1B
10	AR	200	CYC	C3A-C4A-CHB-C1B
10	AR	200	CYC	C2C-C3C-CAC-CBC
10	AR	200	CYC	NC-C4C-CHD-C1D
10	AR	200	CYC	C3C-C4C-CHD-C1D
10	AV	200	CYC	NA-C4A-CHB-C1B
10	AV	200	CYC	C3A-C4A-CHB-C1B
10	AW	200	CYC	NA-C4A-CHB-C1B
10	AW	200	CYC	C3A-C4A-CHB-C1B
10	AX	200	CYC	NA-C4A-CHB-C1B
10	AX	200	CYC	C3A-C4A-CHB-C1B
10	AX	200	CYC	NC-C4C-CHD-C1D
10	AX	200	CYC	C3C-C4C-CHD-C1D
10	AY	200	CYC	NA-C4A-CHB-C1B
10	AY	200	CYC	C3A-C4A-CHB-C1B
10	AY	200	CYC	NC-C4C-CHD-C1D
10	AY	200	CYC	C3C-C4C-CHD-C1D
10	AZ	200	CYC	NA-C4A-CHB-C1B
10	AZ	200	CYC	C3A-C4A-CHB-C1B
10	AZ	200	CYC	C4B-C3B-CAB-CBB
10	BD	901	CYC	C2B-C3B-CAB-CBB
10	BD	901	CYC	NC-C4C-CHD-C1D
10	BD	901	CYC	C3C-C4C-CHD-C1D
10	BD	902	CYC	NA-C4A-CHB-C1B
10	BI	200	CYC	NA-C4A-CHB-C1B
10	BI	200	CYC	C3A-C4A-CHB-C1B
10	BI	200	CYC	C4B-C3B-CAB-CBB
10	BI	200	CYC	C2C-C3C-CAC-CBC
10	BI	200	CYC	C4C-C3C-CAC-CBC
10	BJ	200	CYC	NA-C4A-CHB-C1B

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type	Atoms
10	BJ	200	CYC	C3A-C4A-CHB-C1B
10	BJ	200	CYC	NC-C4C-CHD-C1D
10	BJ	200	CYC	C3C-C4C-CHD-C1D
10	BK	200	CYC	NA-C4A-CHB-C1B
10	BK	200	CYC	C3A-C4A-CHB-C1B
10	BK	200	CYC	C4B-C3B-CAB-CBB
10	BL	200	CYC	NA-C4A-CHB-C1B
10	BL	200	CYC	C3A-C4A-CHB-C1B
10	BL	200	CYC	C4B-C3B-CAB-CBB
10	BL	200	CYC	C2C-C3C-CAC-CBC
10	BL	200	CYC	C4C-C3C-CAC-CBC
10	BL	200	CYC	NC-C4C-CHD-C1D
10	BL	200	CYC	C3C-C4C-CHD-C1D
10	BM	200	CYC	NA-C4A-CHB-C1B
10	BM	200	CYC	C3A-C4A-CHB-C1B
10	BM	200	CYC	NC-C4C-CHD-C1D
10	BM	200	CYC	C3C-C4C-CHD-C1D
10	BN	200	CYC	NA-C4A-CHB-C1B
10	BN	200	CYC	NC-C4C-CHD-C1D
10	BN	200	CYC	C3C-C4C-CHD-C1D
10	BP	200	CYC	NA-C4A-CHB-C1B
10	BP	200	CYC	C3A-C4A-CHB-C1B
10	BQ	200	CYC	C2B-C3B-CAB-CBB
10	BR	200	CYC	NA-C4A-CHB-C1B
10	BR	200	CYC	C3A-C4A-CHB-C1B
10	BS	200	CYC	NA-C4A-CHB-C1B
10	BS	200	CYC	C3A-C4A-CHB-C1B
10	BS	200	CYC	C4C-C3C-CAC-CBC
10	BT	200	CYC	NA-C4A-CHB-C1B
10	BT	200	CYC	C3A-C4A-CHB-C1B
10	BT	200	CYC	NC-C4C-CHD-C1D
10	BT	200	CYC	C3C-C4C-CHD-C1D
10	BV	200	CYC	NA-C4A-CHB-C1B
10	BV	200	CYC	C3A-C4A-CHB-C1B
10	BW	200	CYC	NA-C4A-CHB-C1B
10	BW	200	CYC	C3A-C4A-CHB-C1B
10	BX	200	CYC	NA-C4A-CHB-C1B
10	BX	200	CYC	C3A-C4A-CHB-C1B
10	BZ	200	CYC	NA-C4A-CHB-C1B
10	BZ	200	CYC	C3A-C4A-CHB-C1B
10	BZ	200	CYC	NC-C4C-CHD-C1D
10	BZ	200	CYC	C3C-C4C-CHD-C1D

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type	Atoms
10	CC	200	CYC	NA-C4A-CHB-C1B
10	CC	200	CYC	C3A-C4A-CHB-C1B
10	CC	200	CYC	C4B-C3B-CAB-CBB
10	CD	200	CYC	C4C-C3C-CAC-CBC
10	CE	200	CYC	NA-C4A-CHB-C1B
10	CE	200	CYC	C3A-C4A-CHB-C1B
10	CE	200	CYC	NC-C4C-CHD-C1D
10	CE	200	CYC	C3C-C4C-CHD-C1D
10	CF	200	CYC	C1A-C2A-CAA-CBA
10	CF	200	CYC	C3A-C2A-CAA-CBA
10	CF	200	CYC	NA-C4A-CHB-C1B
10	CF	200	CYC	C3A-C4A-CHB-C1B
10	CG	200	CYC	NA-C4A-CHB-C1B
10	CG	200	CYC	C3A-C4A-CHB-C1B
10	CH	200	CYC	NA-C4A-CHB-C1B
10	CH	200	CYC	C3A-C4A-CHB-C1B
10	CH	200	CYC	NC-C4C-CHD-C1D
10	CH	200	CYC	C3C-C4C-CHD-C1D
10	CM	901	CYC	NA-C4A-CHB-C1B
10	CM	901	CYC	C2B-C3B-CAB-CBB
10	CM	901	CYC	C4C-C3C-CAC-CBC
10	CM	901	CYC	NC-C4C-CHD-C1D
10	CM	901	CYC	C3C-C4C-CHD-C1D
10	CR	200	CYC	NA-C4A-CHB-C1B
10	CS	200	CYC	NA-C4A-CHB-C1B
10	CS	200	CYC	C3A-C4A-CHB-C1B
10	CT	200	CYC	C3A-C2A-CAA-CBA
10	CT	200	CYC	C4B-C3B-CAB-CBB
10	CT	200	CYC	NC-C4C-CHD-C1D
10	CT	200	CYC	C3C-C4C-CHD-C1D
10	CU	200	CYC	NA-C4A-CHB-C1B
10	CU	200	CYC	C3A-C4A-CHB-C1B
10	CV	200	CYC	NA-C4A-CHB-C1B
10	CV	200	CYC	C3A-C4A-CHB-C1B
10	CW	200	CYC	C3A-C4A-CHB-C1B
10	CW	200	CYC	NC-C4C-CHD-C1D
10	CW	200	CYC	C3C-C4C-CHD-C1D
10	CY	200	CYC	C3A-C2A-CAA-CBA
10	CY	200	CYC	NA-C4A-CHB-C1B
10	CY	200	CYC	C3A-C4A-CHB-C1B
10	CZ	200	CYC	NA-C4A-CHB-C1B
10	CZ	200	CYC	C3A-C4A-CHB-C1B

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type	Atoms
10	CZ	200	CYC	C2C-C3C-CAC-CBC
10	CZ	200	CYC	C4C-C3C-CAC-CBC
10	CZ	200	CYC	NC-C4C-CHD-C1D
10	CZ	200	CYC	C3C-C4C-CHD-C1D
10	DA	200	CYC	NA-C4A-CHB-C1B
10	DA	200	CYC	C3A-C4A-CHB-C1B
10	DB	200	CYC	NA-C4A-CHB-C1B
10	DB	200	CYC	C3A-C4A-CHB-C1B
10	DB	200	CYC	NC-C4C-CHD-C1D
10	DB	200	CYC	C3C-C4C-CHD-C1D
10	DC	200	CYC	NA-C4A-CHB-C1B
10	DC	200	CYC	C4C-C3C-CAC-CBC
10	DD	200	CYC	NA-C4A-CHB-C1B
10	DD	200	CYC	C3A-C4A-CHB-C1B
10	DD	200	CYC	C4B-C3B-CAB-CBB
10	DJ	200	CYC	NA-C4A-CHB-C1B
10	DJ	200	CYC	C3A-C4A-CHB-C1B
10	DJ	200	CYC	C4B-C3B-CAB-CBB
10	DK	200	CYC	NA-C4A-CHB-C1B
10	DK	200	CYC	C3A-C4A-CHB-C1B
10	DL	200	CYC	C1A-C2A-CAA-CBA
10	DL	200	CYC	C3A-C2A-CAA-CBA
10	DL	200	CYC	NA-C4A-CHB-C1B
10	DL	200	CYC	C4B-C3B-CAB-CBB
10	DL	200	CYC	NC-C4C-CHD-C1D
10	DL	200	CYC	C3C-C4C-CHD-C1D
10	DM	200	CYC	NA-C4A-CHB-C1B
10	DM	200	CYC	C3A-C4A-CHB-C1B
10	DO	200	CYC	NC-C4C-CHD-C1D
10	DO	200	CYC	C3C-C4C-CHD-C1D
10	DQ	200	CYC	C1A-C2A-CAA-CBA
10	DQ	200	CYC	C3A-C2A-CAA-CBA
10	DQ	200	CYC	NA-C4A-CHB-C1B
10	DQ	200	CYC	C3A-C4A-CHB-C1B
10	DQ	200	CYC	C2C-C3C-CAC-CBC
10	DQ	200	CYC	C4C-C3C-CAC-CBC
10	DQ	200	CYC	NC-C4C-CHD-C1D
10	DQ	200	CYC	C3C-C4C-CHD-C1D
10	DR	200	CYC	NA-C4A-CHB-C1B
10	DR	200	CYC	C3A-C4A-CHB-C1B
10	DR	200	CYC	C4C-C3C-CAC-CBC
10	DR	200	CYC	NC-C4C-CHD-C1D

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type	Atoms
10	DR	200	CYC	C3C-C4C-CHD-C1D
10	DS	200	CYC	NA-C4A-CHB-C1B
10	DS	200	CYC	C3A-C4A-CHB-C1B
10	DT	200	CYC	NA-C4A-CHB-C1B
10	DT	200	CYC	C3A-C4A-CHB-C1B
10	DT	200	CYC	NC-C4C-CHD-C1D
10	DT	200	CYC	C3C-C4C-CHD-C1D
10	DU	200	CYC	C4C-C3C-CAC-CBC
10	DV	200	CYC	NA-C4A-CHB-C1B
10	DV	200	CYC	C3A-C4A-CHB-C1B
10	DV	200	CYC	C4B-C3B-CAB-CBB
10	DV	200	CYC	NC-C4C-CHD-C1D
10	DV	200	CYC	C3C-C4C-CHD-C1D
11	BH	400	45D	C16-C08-C20-C24
11	BH	400	45D	C20-C24-C26-C28
11	BH	400	45D	C20-C24-C26-C30
11	BH	400	45D	C26-C30-C32-C34
11	BH	400	45D	C31-C33-C35-C37
11	BH	400	45D	C31-C33-C35-C39
11	CQ	400	45D	C16-C08-C20-C24
11	CQ	400	45D	C19-C23-C25-C29
11	CQ	400	45D	C20-C24-C26-C28
11	CQ	400	45D	C26-C30-C32-C34
11	CQ	400	45D	C31-C33-C35-C37
11	CQ	400	45D	C31-C33-C35-C39
11	DI	400	45D	C15-C07-C19-C23
11	DI	400	45D	C16-C08-C20-C24
11	DI	400	45D	C26-C30-C32-C34
11	EA	400	45D	C16-C08-C20-C24
11	EA	400	45D	C19-C23-C25-C27
11	EA	400	45D	C20-C24-C26-C28
11	EA	400	45D	C20-C24-C26-C30
11	EA	400	45D	C26-C30-C32-C34
10	AB	200	CYC	C2B-C3B-CAB-CBB
10	AW	200	CYC	C2B-C3B-CAB-CBB
10	CF	200	CYC	C2B-C3B-CAB-CBB
10	CU	200	CYC	C2B-C3B-CAB-CBB
10	DC	200	CYC	C2B-C3B-CAB-CBB
10	DK	200	CYC	C2B-C3B-CAB-CBB
10	DM	200	CYC	C2B-C3B-CAB-CBB
10	AX	200	CYC	C2B-C3B-CAB-CBB
10	CG	200	CYC	C2B-C3B-CAB-CBB

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type	Atoms
10	AO	200	CYC	C2B-C3B-CAB-CBB
10	BX	200	CYC	C2B-C3B-CAB-CBB
10	CW	200	CYC	C2B-C3B-CAB-CBB
10	DQ	200	CYC	C2B-C3B-CAB-CBB
10	DO	200	CYC	C2B-C3B-CAB-CBB
10	DT	200	CYC	C2B-C3B-CAB-CBB
10	BN	200	CYC	C2B-C3B-CAB-CBB
10	DB	200	CYC	C2B-C3B-CAB-CBB
10	DS	200	CYC	C2B-C3B-CAB-CBB
10	DU	200	CYC	C2B-C3B-CAB-CBB
10	BY	200	CYC	C2B-C3B-CAB-CBB
10	CV	200	CYC	C2B-C3B-CAB-CBB
10	DC	200	CYC	C3A-C2A-CAA-CBA
10	DU	200	CYC	C3A-C2A-CAA-CBA
10	AI	200	CYC	ND-C1D-CHD-C4C
10	AL	200	CYC	ND-C1D-CHD-C4C
10	AO	200	CYC	ND-C1D-CHD-C4C
10	AV	200	CYC	ND-C1D-CHD-C4C
10	AW	200	CYC	ND-C1D-CHD-C4C
10	BL	200	CYC	ND-C1D-CHD-C4C
10	BS	200	CYC	ND-C1D-CHD-C4C
10	BT	200	CYC	ND-C1D-CHD-C4C
10	BW	200	CYC	ND-C1D-CHD-C4C
10	CF	200	CYC	ND-C1D-CHD-C4C
10	CR	200	CYC	ND-C1D-CHD-C4C
10	CT	200	CYC	ND-C1D-CHD-C4C
10	CU	200	CYC	ND-C1D-CHD-C4C
10	CY	200	CYC	ND-C1D-CHD-C4C
10	DJ	200	CYC	ND-C1D-CHD-C4C
10	DM	200	CYC	ND-C1D-CHD-C4C
10	DR	200	CYC	ND-C1D-CHD-C4C
10	DS	200	CYC	ND-C1D-CHD-C4C
10	DT	200	CYC	ND-C1D-CHD-C4C
10	DU	200	CYC	ND-C1D-CHD-C4C
10	AD	200	CYC	C2D-C1D-CHD-C4C
10	AF	200	CYC	C2D-C1D-CHD-C4C
10	AH	200	CYC	C2D-C1D-CHD-C4C
10	AI	200	CYC	C2D-C1D-CHD-C4C
10	AJ	200	CYC	C2D-C1D-CHD-C4C
10	AK	200	CYC	C2D-C1D-CHD-C4C
10	AO	200	CYC	C2D-C1D-CHD-C4C
10	AQ	200	CYC	C2D-C1D-CHD-C4C

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type	Atoms
10	AV	200	CYC	C2D-C1D-CHD-C4C
10	AX	200	CYC	C2D-C1D-CHD-C4C
10	BI	200	CYC	C2D-C1D-CHD-C4C
10	BL	200	CYC	C2D-C1D-CHD-C4C
10	BN	200	CYC	C2D-C1D-CHD-C4C
10	BQ	200	CYC	C2D-C1D-CHD-C4C
10	BR	200	CYC	C2D-C1D-CHD-C4C
10	BT	200	CYC	C2D-C1D-CHD-C4C
10	BV	200	CYC	C2D-C1D-CHD-C4C
10	BW	200	CYC	C2D-C1D-CHD-C4C
10	BX	200	CYC	C2D-C1D-CHD-C4C
10	BY	200	CYC	C2D-C1D-CHD-C4C
10	CD	200	CYC	C2D-C1D-CHD-C4C
10	CF	200	CYC	C2D-C1D-CHD-C4C
10	CG	200	CYC	C2D-C1D-CHD-C4C
10	CH	200	CYC	C2D-C1D-CHD-C4C
10	CM	901	CYC	C2D-C1D-CHD-C4C
10	CR	200	CYC	C2D-C1D-CHD-C4C
10	CS	200	CYC	C2D-C1D-CHD-C4C
10	CV	200	CYC	C2D-C1D-CHD-C4C
10	CY	200	CYC	C2D-C1D-CHD-C4C
10	DA	200	CYC	C2D-C1D-CHD-C4C
10	DC	200	CYC	C2D-C1D-CHD-C4C
10	DD	200	CYC	C2D-C1D-CHD-C4C
10	DJ	200	CYC	C2D-C1D-CHD-C4C
10	DQ	200	CYC	C2D-C1D-CHD-C4C
10	DU	200	CYC	C2D-C1D-CHD-C4C
10	CT	200	CYC	C1A-C2A-CAA-CBA
10	CY	200	CYC	C1A-C2A-CAA-CBA
10	DC	200	CYC	C1A-C2A-CAA-CBA
10	DU	200	CYC	C1A-C2A-CAA-CBA
10	DR	200	CYC	C2B-C3B-CAB-CBB
10	BY	200	CYC	NA-C4A-CHB-C1B
10	AY	200	CYC	C2B-C3B-CAB-CBB
10	CZ	200	CYC	C2B-C3B-CAB-CBB
10	BD	902	CYC	C3A-C4A-CHB-C1B
10	BN	200	CYC	C3A-C4A-CHB-C1B
10	CR	200	CYC	C3A-C4A-CHB-C1B
10	DL	200	CYC	C3A-C4A-CHB-C1B
10	AA	200	CYC	ND-C1D-CHD-C4C
10	AB	200	CYC	ND-C1D-CHD-C4C
10	AC	200	CYC	ND-C1D-CHD-C4C

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type	Atoms
10	AD	200	CYC	ND-C1D-CHD-C4C
10	AE	200	CYC	ND-C1D-CHD-C4C
10	AF	200	CYC	ND-C1D-CHD-C4C
10	AH	200	CYC	ND-C1D-CHD-C4C
10	AJ	200	CYC	ND-C1D-CHD-C4C
10	AK	200	CYC	ND-C1D-CHD-C4C
10	AN	200	CYC	ND-C1D-CHD-C4C
10	AP	200	CYC	ND-C1D-CHD-C4C
10	AQ	200	CYC	ND-C1D-CHD-C4C
10	AR	200	CYC	ND-C1D-CHD-C4C
10	AU	200	CYC	ND-C1D-CHD-C4C
10	AX	200	CYC	ND-C1D-CHD-C4C
10	AY	200	CYC	ND-C1D-CHD-C4C
10	AZ	200	CYC	ND-C1D-CHD-C4C
10	BD	901	CYC	ND-C1D-CHD-C4C
10	BI	200	CYC	ND-C1D-CHD-C4C
10	BJ	200	CYC	ND-C1D-CHD-C4C
10	BK	200	CYC	ND-C1D-CHD-C4C
10	BM	200	CYC	ND-C1D-CHD-C4C
10	BN	200	CYC	ND-C1D-CHD-C4C
10	BP	200	CYC	ND-C1D-CHD-C4C
10	BQ	200	CYC	ND-C1D-CHD-C4C
10	BR	200	CYC	ND-C1D-CHD-C4C
10	BV	200	CYC	ND-C1D-CHD-C4C
10	BX	200	CYC	ND-C1D-CHD-C4C
10	BY	200	CYC	ND-C1D-CHD-C4C
10	BZ	200	CYC	ND-C1D-CHD-C4C
10	CC	200	CYC	ND-C1D-CHD-C4C
10	CD	200	CYC	ND-C1D-CHD-C4C
10	CE	200	CYC	ND-C1D-CHD-C4C
10	CG	200	CYC	ND-C1D-CHD-C4C
10	CH	200	CYC	ND-C1D-CHD-C4C
10	CM	901	CYC	ND-C1D-CHD-C4C
10	CS	200	CYC	ND-C1D-CHD-C4C
10	CV	200	CYC	ND-C1D-CHD-C4C
10	CW	200	CYC	ND-C1D-CHD-C4C
10	CZ	200	CYC	ND-C1D-CHD-C4C
10	DA	200	CYC	ND-C1D-CHD-C4C
10	DC	200	CYC	ND-C1D-CHD-C4C
10	DD	200	CYC	ND-C1D-CHD-C4C
10	DK	200	CYC	ND-C1D-CHD-C4C
10	DL	200	CYC	ND-C1D-CHD-C4C

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type	Atoms
10	DN	200	CYC	ND-C1D-CHD-C4C
10	DO	200	CYC	ND-C1D-CHD-C4C
10	DQ	200	CYC	ND-C1D-CHD-C4C
10	DV	200	CYC	ND-C1D-CHD-C4C
10	AK	200	CYC	C2B-C3B-CAB-CBB
10	AA	200	CYC	C2D-C1D-CHD-C4C
10	AB	200	CYC	C2D-C1D-CHD-C4C
10	AC	200	CYC	C2D-C1D-CHD-C4C
10	AE	200	CYC	C2D-C1D-CHD-C4C
10	AL	200	CYC	C2D-C1D-CHD-C4C
10	AN	200	CYC	C2D-C1D-CHD-C4C
10	AP	200	CYC	C2D-C1D-CHD-C4C
10	AR	200	CYC	C2D-C1D-CHD-C4C
10	AU	200	CYC	C2D-C1D-CHD-C4C
10	AW	200	CYC	C2D-C1D-CHD-C4C
10	AY	200	CYC	C2D-C1D-CHD-C4C
10	AZ	200	CYC	C2D-C1D-CHD-C4C
10	BJ	200	CYC	C2D-C1D-CHD-C4C
10	BK	200	CYC	C2D-C1D-CHD-C4C
10	BM	200	CYC	C2D-C1D-CHD-C4C
10	BP	200	CYC	C2D-C1D-CHD-C4C
10	BS	200	CYC	C2D-C1D-CHD-C4C
10	BZ	200	CYC	C2D-C1D-CHD-C4C
10	CC	200	CYC	C2D-C1D-CHD-C4C
10	CE	200	CYC	C2D-C1D-CHD-C4C
10	CT	200	CYC	C2D-C1D-CHD-C4C
10	CU	200	CYC	C2D-C1D-CHD-C4C
10	CW	200	CYC	C2D-C1D-CHD-C4C
10	CZ	200	CYC	C2D-C1D-CHD-C4C
10	DB	200	CYC	C2D-C1D-CHD-C4C
10	DK	200	CYC	C2D-C1D-CHD-C4C
10	DM	200	CYC	C2D-C1D-CHD-C4C
10	DN	200	CYC	C2D-C1D-CHD-C4C
10	DO	200	CYC	C2D-C1D-CHD-C4C
10	DR	200	CYC	C2D-C1D-CHD-C4C
10	DS	200	CYC	C2D-C1D-CHD-C4C
10	DT	200	CYC	C2D-C1D-CHD-C4C
10	DV	200	CYC	C2D-C1D-CHD-C4C
10	CS	200	CYC	C2B-C3B-CAB-CBB
11	BH	400	45D	C19-C23-C25-C27
11	BH	400	45D	C32-C34-C36-C40
11	CQ	400	45D	C19-C23-C25-C27

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type	Atoms
11	CQ	400	45D	C32-C34-C36-C40
11	DI	400	45D	C20-C24-C26-C28
11	DI	400	45D	C31-C33-C35-C39
11	DI	400	45D	C32-C34-C36-C40
11	EA	400	45D	C31-C33-C35-C39
11	EA	400	45D	C32-C34-C36-C40
11	BH	400	45D	C19-C23-C25-C29
11	BH	400	45D	C32-C34-C36-C38
11	CQ	400	45D	C20-C24-C26-C30
11	CQ	400	45D	C32-C34-C36-C38
11	DI	400	45D	C20-C24-C26-C30
11	DI	400	45D	C31-C33-C35-C37
11	DI	400	45D	C32-C34-C36-C38
11	EA	400	45D	C19-C23-C25-C29
11	EA	400	45D	C31-C33-C35-C37
11	EA	400	45D	C32-C34-C36-C38
10	DB	200	CYC	ND-C1D-CHD-C4C
10	BD	901	CYC	C2D-C1D-CHD-C4C
10	DL	200	CYC	C2D-C1D-CHD-C4C
10	AF	200	CYC	C2B-C3B-CAB-CBB
10	AA	200	CYC	C4B-C3B-CAB-CBB
10	AD	200	CYC	C4B-C3B-CAB-CBB
10	AH	200	CYC	C4B-C3B-CAB-CBB
10	AI	200	CYC	C4B-C3B-CAB-CBB
10	AJ	200	CYC	C4B-C3B-CAB-CBB
10	AK	200	CYC	C4B-C3B-CAB-CBB
10	AL	200	CYC	C4B-C3B-CAB-CBB
10	AN	200	CYC	C4B-C3B-CAB-CBB
10	AR	200	CYC	C4B-C3B-CAB-CBB
10	BD	902	CYC	C4B-C3B-CAB-CBB
10	BQ	200	CYC	C4B-C3B-CAB-CBB
10	BS	200	CYC	C4B-C3B-CAB-CBB
10	BT	200	CYC	C4B-C3B-CAB-CBB
10	BV	200	CYC	C4B-C3B-CAB-CBB
10	BZ	200	CYC	C4B-C3B-CAB-CBB
10	CD	200	CYC	C4B-C3B-CAB-CBB
10	CR	200	CYC	C4B-C3B-CAB-CBB
10	DR	200	CYC	C4B-C3B-CAB-CBB
10	CE	200	CYC	C4D-C3D-CAD-CBD
10	AK	200	CYC	C3A-C2A-CAA-CBA
10	AW	200	CYC	C2A-CAA-CBA-CGA
10	BS	200	CYC	C2A-CAA-CBA-CGA

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type	Atoms
10	AL	200	CYC	C2B-C3B-CAB-CBB
10	CM	901	CYC	C3A-C4A-CHB-C1B
10	CR	200	CYC	C2B-C3B-CAB-CBB
10	AR	200	CYC	C2B-C3B-CAB-CBB
10	BM	200	CYC	C3A-C2A-CAA-CBA
10	BT	200	CYC	C2B-C3B-CAB-CBB
10	BR	200	CYC	C2B-C3B-CAB-CBB
10	BJ	200	CYC	C4D-C3D-CAD-CBD
10	CH	200	CYC	C4D-C3D-CAD-CBD
10	DT	200	CYC	C4D-C3D-CAD-CBD
10	AU	200	CYC	NA-C4A-CHB-C1B
10	BD	901	CYC	NA-C4A-CHB-C1B
10	BQ	200	CYC	NA-C4A-CHB-C1B
10	CD	200	CYC	NA-C4A-CHB-C1B
10	CM	902	CYC	NA-C4A-CHB-C1B
10	CW	200	CYC	NA-C4A-CHB-C1B
10	DN	200	CYC	NA-C4A-CHB-C1B
10	DO	200	CYC	NA-C4A-CHB-C1B
10	DU	200	CYC	NA-C4A-CHB-C1B
11	DI	400	45D	C19-C23-C25-C27
10	CE	200	CYC	C2B-C3B-CAB-CBB
10	AW	200	CYC	C3A-C2A-CAA-CBA
11	DI	400	45D	C19-C23-C25-C29
10	BV	200	CYC	C2A-CAA-CBA-CGA
10	BM	200	CYC	C1A-C2A-CAA-CBA
10	AF	200	CYC	C3A-C4A-CHB-C1B
10	AI	200	CYC	C3A-C4A-CHB-C1B
10	AQ	200	CYC	C3A-C4A-CHB-C1B
10	AU	200	CYC	C3A-C4A-CHB-C1B
10	BD	901	CYC	C3A-C4A-CHB-C1B
10	BQ	200	CYC	C3A-C4A-CHB-C1B
10	BY	200	CYC	C3A-C4A-CHB-C1B
10	CD	200	CYC	C3A-C4A-CHB-C1B
10	CM	902	CYC	C3A-C4A-CHB-C1B
10	DC	200	CYC	C3A-C4A-CHB-C1B
10	DN	200	CYC	C3A-C4A-CHB-C1B
10	DO	200	CYC	C3A-C4A-CHB-C1B
10	DU	200	CYC	C3A-C4A-CHB-C1B
10	AV	200	CYC	C4B-C3B-CAB-CBB
10	CM	901	CYC	C4B-C3B-CAB-CBB
10	CZ	200	CYC	C4B-C3B-CAB-CBB
10	DA	200	CYC	C4B-C3B-CAB-CBB

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type	Atoms
10	CE	200	CYC	C2D-C3D-CAD-CBD
11	BH	400	45D	C03-C07-C19-C23
11	BH	400	45D	C15-C07-C19-C23
11	BH	400	45D	C04-C08-C20-C24
11	CQ	400	45D	C03-C07-C19-C23
11	DI	400	45D	C03-C07-C19-C23
11	DI	400	45D	C04-C08-C20-C24
11	EA	400	45D	C03-C07-C19-C23
11	EA	400	45D	C15-C07-C19-C23
10	BV	200	CYC	C2B-C3B-CAB-CBB
10	BZ	200	CYC	C2B-C3B-CAB-CBB
10	DA	200	CYC	C2B-C3B-CAB-CBB
10	AE	200	CYC	C1A-C2A-CAA-CBA
10	CT	200	CYC	C2B-C3B-CAB-CBB
10	DL	200	CYC	C2B-C3B-CAB-CBB
10	AN	200	CYC	C2B-C3B-CAB-CBB
10	AP	200	CYC	C2B-C3B-CAB-CBB
10	CH	200	CYC	C2D-C3D-CAD-CBD
10	DT	200	CYC	C2D-C3D-CAD-CBD
10	AI	200	CYC	C2B-C3B-CAB-CBB
10	CY	200	CYC	C4D-C3D-CAD-CBD
10	AW	200	CYC	C1A-C2A-CAA-CBA
10	AV	200	CYC	C2B-C3B-CAB-CBB
10	BS	200	CYC	C2C-C3C-CAC-CBC
10	DC	200	CYC	C2C-C3C-CAC-CBC
10	DU	200	CYC	C2C-C3C-CAC-CBC
10	AK	200	CYC	C1A-C2A-CAA-CBA
10	BT	200	CYC	C1A-C2A-CAA-CBA
10	AE	200	CYC	C3A-C2A-CAA-CBA
10	BT	200	CYC	C3A-C2A-CAA-CBA
10	AQ	200	CYC	C1A-C2A-CAA-CBA
10	BY	200	CYC	C1A-C2A-CAA-CBA
10	BS	200	CYC	C2B-C3B-CAB-CBB
10	AJ	200	CYC	C2B-C3B-CAB-CBB
10	AL	200	CYC	C1A-C2A-CAA-CBA
10	AQ	200	CYC	C4D-C3D-CAD-CBD
10	AY	200	CYC	C4D-C3D-CAD-CBD
10	BN	200	CYC	C2A-CAA-CBA-CGA
10	DQ	200	CYC	C2A-CAA-CBA-CGA
10	DV	200	CYC	C2A-CAA-CBA-CGA
10	BD	901	CYC	C4B-C3B-CAB-CBB
10	BP	200	CYC	C4B-C3B-CAB-CBB

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type	Atoms
11	BH	400	45D	C25-C29-C31-C33
11	CQ	400	45D	C25-C29-C31-C33
10	AL	200	CYC	C3A-C2A-CAA-CBA
10	AA	200	CYC	C2B-C3B-CAB-CBB
10	AB	200	CYC	C4D-C3D-CAD-CBD
11	CQ	400	45D	C04-C08-C20-C24
11	EA	400	45D	C04-C08-C20-C24
10	DL	200	CYC	C2A-CAA-CBA-CGA
10	BL	200	CYC	C2B-C3B-CAB-CBB
10	AR	200	CYC	C4C-C3C-CAC-CBC
10	AU	200	CYC	C4C-C3C-CAC-CBC
10	DN	200	CYC	C2B-C3B-CAB-CBB
10	BV	200	CYC	C4D-C3D-CAD-CBD
10	DB	200	CYC	C4D-C3D-CAD-CBD
10	BJ	200	CYC	C2D-C3D-CAD-CBD
10	AQ	200	CYC	C4B-C3B-CAB-CBB
10	CW	200	CYC	C4B-C3B-CAB-CBB
10	AH	200	CYC	C2A-CAA-CBA-CGA
10	BY	200	CYC	C3A-C2A-CAA-CBA
10	CH	200	CYC	C2B-C3B-CAB-CBB
10	DJ	200	CYC	C2B-C3B-CAB-CBB
10	AC	200	CYC	C2B-C3B-CAB-CBB
10	AW	200	CYC	C4B-C3B-CAB-CBB
10	CF	200	CYC	C4B-C3B-CAB-CBB
10	DC	200	CYC	C4B-C3B-CAB-CBB
10	CC	200	CYC	C2B-C3B-CAB-CBB
10	CM	901	CYC	C2C-C3C-CAC-CBC
10	AH	200	CYC	C2B-C3B-CAB-CBB
10	BI	200	CYC	C2B-C3B-CAB-CBB
10	BW	200	CYC	C2B-C3B-CAB-CBB
10	AY	200	CYC	C2D-C3D-CAD-CBD
11	DI	400	45D	C25-C29-C31-C33
11	EA	400	45D	C25-C29-C31-C33
10	AQ	200	CYC	C3A-C2A-CAA-CBA
10	AK	200	CYC	NC-C4C-CHD-C1D
10	AL	200	CYC	NC-C4C-CHD-C1D
10	AZ	200	CYC	NC-C4C-CHD-C1D
10	BR	200	CYC	NC-C4C-CHD-C1D
10	CM	902	CYC	NC-C4C-CHD-C1D
11	BH	400	45D	C36-C38-C42-C41
10	BK	200	CYC	C2B-C3B-CAB-CBB
10	CU	200	CYC	C4B-C3B-CAB-CBB

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type	Atoms
10	DB	200	CYC	C2D-C3D-CAD-CBD
10	BD	902	CYC	ND-C1D-CHD-C4C
10	DO	200	CYC	C3A-C2A-CAA-CBA
11	EA	400	45D	C36-C38-C42-C41
10	CY	200	CYC	C2A-CAA-CBA-CGA
10	AQ	200	CYC	C2D-C3D-CAD-CBD
10	BV	200	CYC	C2D-C3D-CAD-CBD
10	CU	200	CYC	C1A-C2A-CAA-CBA
10	AQ	200	CYC	CAA-CBA-CGA-O2A
11	CQ	400	45D	C36-C38-C42-C41
11	DI	400	45D	C35-C37-C41-C42
10	AO	200	CYC	CAA-CBA-CGA-O2A
10	AY	200	CYC	CAA-CBA-CGA-O2A
10	BD	902	CYC	CAA-CBA-CGA-O2A
10	CU	200	CYC	C3A-C2A-CAA-CBA
10	BV	200	CYC	C1A-C2A-CAA-CBA
10	CT	200	CYC	C3A-C4A-CHB-C1B
10	AQ	200	CYC	CAA-CBA-CGA-O1A
10	BI	200	CYC	CAD-CBD-CGD-O1D
10	BP	200	CYC	CAA-CBA-CGA-O1A
10	DV	200	CYC	CAA-CBA-CGA-O1A
10	DV	200	CYC	CAA-CBA-CGA-O2A
10	CY	200	CYC	C2D-C3D-CAD-CBD
10	AC	200	CYC	CAA-CBA-CGA-O2A
10	BI	200	CYC	CAD-CBD-CGD-O2D
10	BX	200	CYC	CAA-CBA-CGA-O1A
10	BZ	200	CYC	CAA-CBA-CGA-O1A
10	AL	200	CYC	CAA-CBA-CGA-O1A
10	AR	200	CYC	CAA-CBA-CGA-O2A
10	BR	200	CYC	CAD-CBD-CGD-O2D
10	AP	200	CYC	CAD-CBD-CGD-O1D
10	BY	200	CYC	CAA-CBA-CGA-O2A
10	CC	200	CYC	CAA-CBA-CGA-O1A
10	CC	200	CYC	CAA-CBA-CGA-O2A
10	CM	902	CYC	CAA-CBA-CGA-O2A
10	DJ	200	CYC	CAA-CBA-CGA-O2A
10	AC	200	CYC	CAA-CBA-CGA-O1A
10	AP	200	CYC	CAA-CBA-CGA-O1A
10	AZ	200	CYC	CAA-CBA-CGA-O1A
10	BP	200	CYC	CAA-CBA-CGA-O2A
10	CR	200	CYC	CAD-CBD-CGD-O1D
10	DC	200	CYC	CAD-CBD-CGD-O1D

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type	Atoms
10	AX	200	CYC	CAA-CBA-CGA-O1A
10	BR	200	CYC	CAA-CBA-CGA-O2A
10	CH	200	CYC	CAA-CBA-CGA-O1A
10	DJ	200	CYC	C4D-C3D-CAD-CBD
10	BW	200	CYC	CAD-CBD-CGD-O2D
10	AJ	200	CYC	CAA-CBA-CGA-O1A
10	AQ	200	CYC	CAD-CBD-CGD-O1D
10	AR	200	CYC	CAA-CBA-CGA-O1A
10	CR	200	CYC	CAA-CBA-CGA-O1A
10	CS	200	CYC	CAD-CBD-CGD-O1D
10	DT	200	CYC	CAA-CBA-CGA-O1A
10	AP	200	CYC	CAA-CBA-CGA-O2A
10	BP	200	CYC	CAD-CBD-CGD-O1D
10	BY	200	CYC	CAD-CBD-CGD-O1D
10	DV	200	CYC	C2B-C3B-CAB-CBB
10	DK	200	CYC	C4B-C3B-CAB-CBB
10	AX	200	CYC	CAA-CBA-CGA-O2A
10	AZ	200	CYC	CAA-CBA-CGA-O2A
10	BK	200	CYC	CAA-CBA-CGA-O1A
10	BV	200	CYC	CAA-CBA-CGA-O1A
10	BW	200	CYC	CAA-CBA-CGA-O2A
10	DN	200	CYC	CAA-CBA-CGA-O1A
10	DN	200	CYC	CAA-CBA-CGA-O2A
10	DN	200	CYC	CAD-CBD-CGD-O1D
10	AB	200	CYC	C2D-C3D-CAD-CBD
10	AH	200	CYC	CAD-CBD-CGD-O1D
10	AJ	200	CYC	CAA-CBA-CGA-O2A
10	AL	200	CYC	CAA-CBA-CGA-O2A
10	AN	200	CYC	CAD-CBD-CGD-O1D
10	CG	200	CYC	CAA-CBA-CGA-O1A
10	CM	901	CYC	CAA-CBA-CGA-O1A
10	DR	200	CYC	CAA-CBA-CGA-O2A
10	BR	200	CYC	CAA-CBA-CGA-O1A
10	BT	200	CYC	CAD-CBD-CGD-O2D
10	CW	200	CYC	CAA-CBA-CGA-O1A
10	DJ	200	CYC	CAD-CBD-CGD-O1D
10	AJ	200	CYC	CAD-CBD-CGD-O1D
10	BX	200	CYC	CAD-CBD-CGD-O1D
10	BZ	200	CYC	CAA-CBA-CGA-O2A
10	CH	200	CYC	CAA-CBA-CGA-O2A
10	AD	200	CYC	CAD-CBD-CGD-O2D
10	AY	200	CYC	CAA-CBA-CGA-O1A

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type	Atoms
10	BD	902	CYC	CAA-CBA-CGA-O1A
10	BR	200	CYC	CAD-CBD-CGD-O1D
10	BX	200	CYC	CAA-CBA-CGA-O2A
10	DJ	200	CYC	CAA-CBA-CGA-O1A
10	BW	200	CYC	CAA-CBA-CGA-O1A
10	BW	200	CYC	CAD-CBD-CGD-O1D
10	BY	200	CYC	CAA-CBA-CGA-O1A
10	CM	902	CYC	CAA-CBA-CGA-O1A
10	CW	200	CYC	CAD-CBD-CGD-O2D
10	CZ	200	CYC	CAA-CBA-CGA-O2A
10	CT	200	CYC	NA-C4A-CHB-C1B
10	AN	200	CYC	CAD-CBD-CGD-O2D
10	BP	200	CYC	CAD-CBD-CGD-O2D
10	CW	200	CYC	CAA-CBA-CGA-O2A
10	BY	200	CYC	CAD-CBD-CGD-O2D
10	CT	200	CYC	C2A-CAA-CBA-CGA
10	AJ	200	CYC	CAD-CBD-CGD-O2D
10	AQ	200	CYC	CAD-CBD-CGD-O2D
10	DJ	200	CYC	CAD-CBD-CGD-O2D
10	BD	901	CYC	C4C-C3C-CAC-CBC
10	AD	200	CYC	CAA-CBA-CGA-O2A
10	AP	200	CYC	CAD-CBD-CGD-O2D
10	AW	200	CYC	CAA-CBA-CGA-O2A
10	BL	200	CYC	CAD-CBD-CGD-O2D
10	BV	200	CYC	CAA-CBA-CGA-O2A
10	BX	200	CYC	CAD-CBD-CGD-O2D
10	CF	200	CYC	CAA-CBA-CGA-O2A
10	CM	901	CYC	CAA-CBA-CGA-O2A
10	CR	200	CYC	CAD-CBD-CGD-O2D
10	CS	200	CYC	CAD-CBD-CGD-O2D
10	AA	200	CYC	CAA-CBA-CGA-O2A
10	AO	200	CYC	CAA-CBA-CGA-O1A
10	BK	200	CYC	CAA-CBA-CGA-O2A
10	CU	200	CYC	CAD-CBD-CGD-O2D
10	DN	200	CYC	CAD-CBD-CGD-O2D
10	CS	200	CYC	CAA-CBA-CGA-O2A
10	DQ	200	CYC	CAD-CBD-CGD-O1D
10	AF	200	CYC	CAD-CBD-CGD-O2D
10	BL	200	CYC	CAA-CBA-CGA-O2A
10	BT	200	CYC	CAA-CBA-CGA-O2A
10	BT	200	CYC	CAD-CBD-CGD-O1D
10	CS	200	CYC	CAA-CBA-CGA-O1A

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type	Atoms
10	DK	200	CYC	CAA-CBA-CGA-O2A
10	DT	200	CYC	CAA-CBA-CGA-O2A
10	AH	200	CYC	CAA-CBA-CGA-O1A
10	BQ	200	CYC	CAA-CBA-CGA-O2A
10	AD	200	CYC	CAD-CBD-CGD-O1D
10	AE	200	CYC	CAA-CBA-CGA-O2A
10	AH	200	CYC	CAA-CBA-CGA-O2A
10	AV	200	CYC	CAA-CBA-CGA-O2A
10	CR	200	CYC	CAA-CBA-CGA-O2A
10	DC	200	CYC	CAD-CBD-CGD-O2D
10	DU	200	CYC	C4B-C3B-CAB-CBB
10	BQ	200	CYC	CAA-CBA-CGA-O1A
10	CG	200	CYC	CAA-CBA-CGA-O2A
10	CU	200	CYC	CAD-CBD-CGD-O1D
10	DK	200	CYC	CAA-CBA-CGA-O1A
10	AH	200	CYC	CAD-CBD-CGD-O2D
10	BL	200	CYC	CAA-CBA-CGA-O1A
10	CW	200	CYC	CAD-CBD-CGD-O1D
10	CZ	200	CYC	CAA-CBA-CGA-O1A
10	BI	200	CYC	CAA-CBA-CGA-O2A
10	BT	200	CYC	CAA-CBA-CGA-O1A
10	AD	200	CYC	CAA-CBA-CGA-O1A
10	AW	200	CYC	CAA-CBA-CGA-O1A
10	BL	200	CYC	CAD-CBD-CGD-O1D
10	CY	200	CYC	CAD-CBD-CGD-O2D
10	DA	200	CYC	CAA-CBA-CGA-O2A
10	DR	200	CYC	CAA-CBA-CGA-O1A
10	AA	200	CYC	CAA-CBA-CGA-O1A
10	CF	200	CYC	CAA-CBA-CGA-O1A
10	DS	200	CYC	CAA-CBA-CGA-O2A
10	BD	902	CYC	C2D-C1D-CHD-C4C
10	AF	200	CYC	CAD-CBD-CGD-O1D
10	AX	200	CYC	C1A-C2A-CAA-CBA
10	BI	200	CYC	CAA-CBA-CGA-O1A
10	DA	200	CYC	CAA-CBA-CGA-O1A
10	DS	200	CYC	CAA-CBA-CGA-O1A
10	CM	901	CYC	C2D-C3D-CAD-CBD
10	CT	200	CYC	CAD-CBD-CGD-O1D
10	CY	200	CYC	CAD-CBD-CGD-O1D
10	DM	200	CYC	CAD-CBD-CGD-O1D
11	EA	400	45D	C35-C37-C41-C42
10	AE	200	CYC	CAA-CBA-CGA-O1A

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type	Atoms
10	AV	200	CYC	CAA-CBA-CGA-O1A
10	DM	200	CYC	C3A-C2A-CAA-CBA
10	DO	200	CYC	C4B-C3B-CAB-CBB
11	CQ	400	45D	C15-C07-C19-C23
10	DQ	200	CYC	CAD-CBD-CGD-O2D
10	AD	200	CYC	C2B-C3B-CAB-CBB
10	CE	200	CYC	CAA-CBA-CGA-O2A
10	CY	200	CYC	CAA-CBA-CGA-O2A
10	CD	200	CYC	CAA-CBA-CGA-O1A
10	CY	200	CYC	CAA-CBA-CGA-O1A
10	CG	200	CYC	CAD-CBD-CGD-O1D
10	DB	200	CYC	CAD-CBD-CGD-O1D
10	DQ	200	CYC	CAA-CBA-CGA-O1A
10	CD	200	CYC	CAD-CBD-CGD-O2D
10	DQ	200	CYC	CAA-CBA-CGA-O2A
10	AZ	200	CYC	CAD-CBD-CGD-O1D
10	CD	200	CYC	CAD-CBD-CGD-O1D
11	BH	400	45D	C35-C37-C41-C42
10	CG	200	CYC	CAD-CBD-CGD-O2D
10	CT	200	CYC	CAD-CBD-CGD-O2D
10	DM	200	CYC	CAD-CBD-CGD-O2D
10	AZ	200	CYC	C2B-C3B-CAB-CBB
10	CD	200	CYC	CAA-CBA-CGA-O2A
10	CE	200	CYC	CAA-CBA-CGA-O1A
10	BD	901	CYC	C2D-C3D-CAD-CBD
10	DM	200	CYC	C4B-C3B-CAB-CBB
10	DD	200	CYC	C2B-C3B-CAB-CBB
10	DB	200	CYC	CAD-CBD-CGD-O2D
10	AL	200	CYC	CAD-CBD-CGD-O2D
10	AE	200	CYC	C2B-C3B-CAB-CBB
10	AZ	200	CYC	CAD-CBD-CGD-O2D
10	DD	200	CYC	CAD-CBD-CGD-O2D
10	AN	200	CYC	CAA-CBA-CGA-O1A
10	BQ	200	CYC	C1A-C2A-CAA-CBA
10	AO	200	CYC	CAD-CBD-CGD-O2D
10	AU	200	CYC	CAA-CBA-CGA-O2A
10	CM	902	CYC	CAD-CBD-CGD-O2D
10	BD	902	CYC	CAD-CBD-CGD-O2D
10	DK	200	CYC	CAD-CBD-CGD-O2D
10	AC	200	CYC	C3C-C4C-CHD-C1D
10	AL	200	CYC	C3C-C4C-CHD-C1D
10	AZ	200	CYC	C3C-C4C-CHD-C1D

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type	Atoms
10	BP	200	CYC	C3C-C4C-CHD-C1D
10	CM	902	CYC	C3C-C4C-CHD-C1D
10	CZ	200	CYC	C3A-C2A-CAA-CBA
10	BT	200	CYC	C2A-CAA-CBA-CGA
10	DO	200	CYC	C1A-C2A-CAA-CBA
10	AI	200	CYC	CAA-CBA-CGA-O2A
10	AY	200	CYC	CAD-CBD-CGD-O2D
10	AB	200	CYC	C4B-C3B-CAB-CBB
10	AU	200	CYC	C4B-C3B-CAB-CBB
10	CG	200	CYC	C4B-C3B-CAB-CBB
10	AE	200	CYC	CAD-CBD-CGD-O2D
10	DT	200	CYC	CAD-CBD-CGD-O2D
10	AK	200	CYC	CAD-CBD-CGD-O2D
10	AR	200	CYC	CAD-CBD-CGD-O2D
10	BV	200	CYC	CAD-CBD-CGD-O2D

There are no ring outliers.

76 monomers are involved in 861 short contacts:

Mol	Chain	Res	Type	Clashes	Symm-Clashes
10	BX	200	CYC	10	0
10	AH	200	CYC	14	0
10	BD	902	CYC	12	0
10	AL	200	CYC	10	0
10	CC	200	CYC	7	0
10	DN	200	CYC	15	0
10	BV	200	CYC	14	0
10	BP	200	CYC	14	0
10	AP	200	CYC	10	0
10	BK	200	CYC	16	0
10	BL	200	CYC	15	0
10	AX	200	CYC	9	0
10	BI	200	CYC	15	0
10	DS	200	CYC	10	0
10	DT	200	CYC	13	0
10	DC	200	CYC	14	0
10	CD	200	CYC	10	0
10	AF	200	CYC	10	0
10	AR	200	CYC	9	0
10	CE	200	CYC	12	0
10	AE	200	CYC	9	0
10	AW	200	CYC	10	0

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type	Clashes	Symm-Clashes
10	BW	200	CYC	8	0
10	BS	200	CYC	11	0
10	AD	200	CYC	14	0
10	CS	200	CYC	7	0
10	DK	200	CYC	8	0
10	DM	200	CYC	8	0
10	BY	200	CYC	9	0
10	AI	200	CYC	11	0
10	AV	200	CYC	17	0
10	BR	200	CYC	6	0
10	BQ	200	CYC	15	0
10	AA	200	CYC	15	0
10	AQ	200	CYC	8	0
10	CG	200	CYC	11	0
10	CM	901	CYC	11	0
10	CZ	200	CYC	15	0
11	BH	400	45D	4	0
11	CQ	400	45D	6	0
10	DJ	200	CYC	13	0
10	DL	200	CYC	11	0
10	DQ	200	CYC	12	0
10	CH	200	CYC	13	0
10	AC	200	CYC	10	0
10	CT	200	CYC	13	0
10	CY	200	CYC	12	0
10	AU	200	CYC	14	0
10	AO	200	CYC	7	0
10	AZ	200	CYC	8	0
10	BT	200	CYC	10	0
10	AB	200	CYC	14	0
10	AY	200	CYC	15	0
10	BZ	200	CYC	12	0
10	AJ	200	CYC	7	0
10	CV	200	CYC	11	0
10	CR	200	CYC	7	0
10	DD	200	CYC	19	0
10	DU	200	CYC	12	0
10	DO	200	CYC	9	0
10	AN	200	CYC	11	0
10	AK	200	CYC	14	0
10	DV	200	CYC	12	0
10	DB	200	CYC	12	0

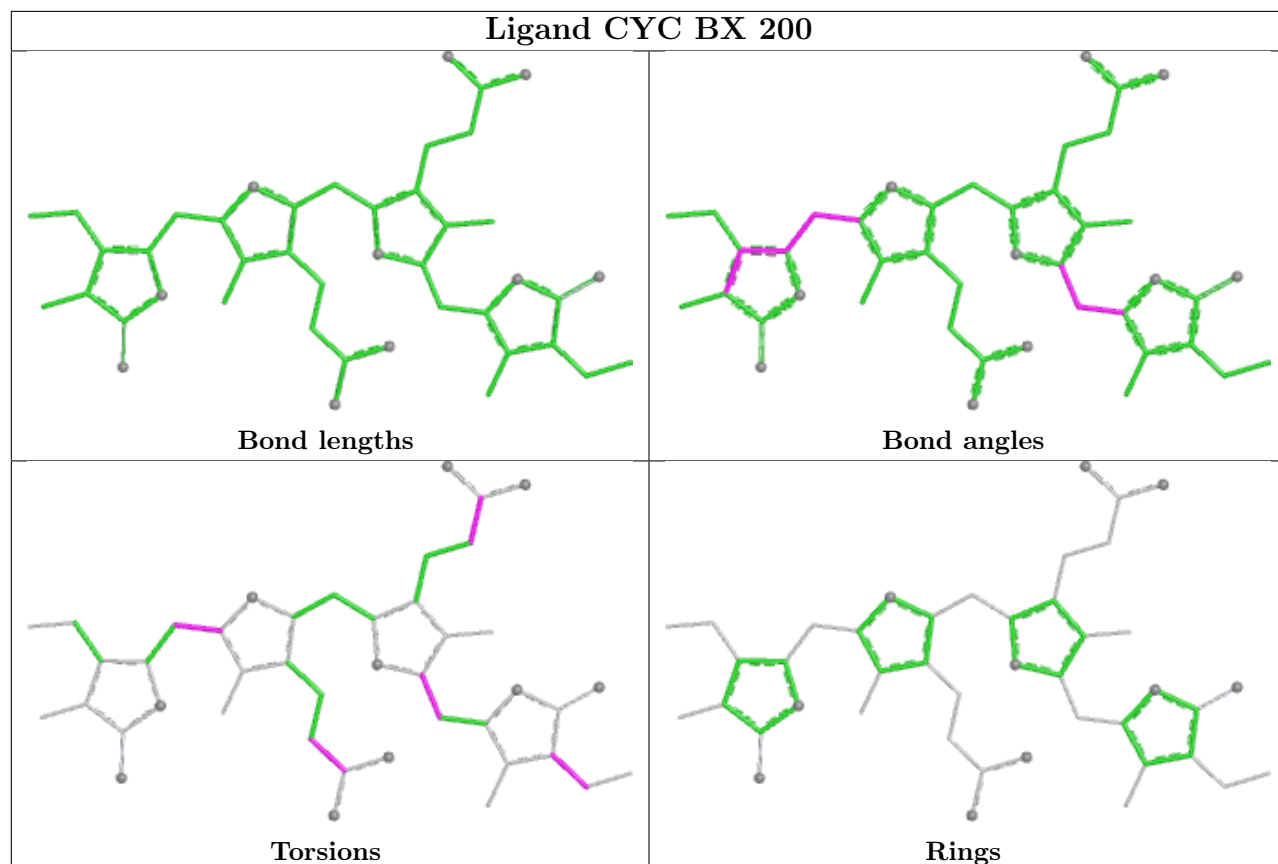
Continued on next page...

Continued from previous page...

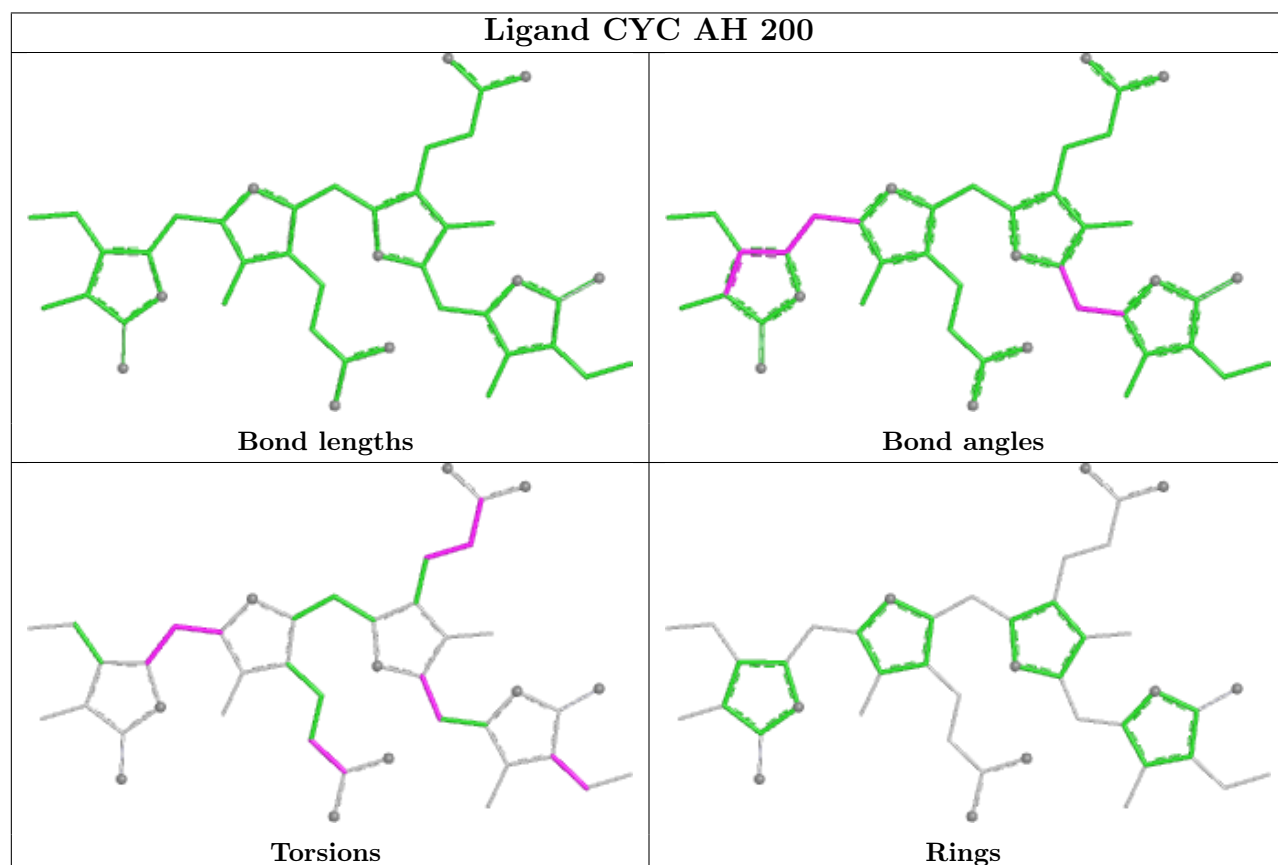
Mol	Chain	Res	Type	Clashes	Symm-Clashes
11	DI	400	45D	14	0
10	CU	200	CYC	8	0
10	DR	200	CYC	14	0
10	CW	200	CYC	12	0
10	BJ	200	CYC	11	0
10	DA	200	CYC	10	0
10	BN	200	CYC	16	0
11	EA	400	45D	12	0
10	CM	902	CYC	12	0
10	BD	901	CYC	11	0
10	BM	200	CYC	11	0
10	CF	200	CYC	10	0

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

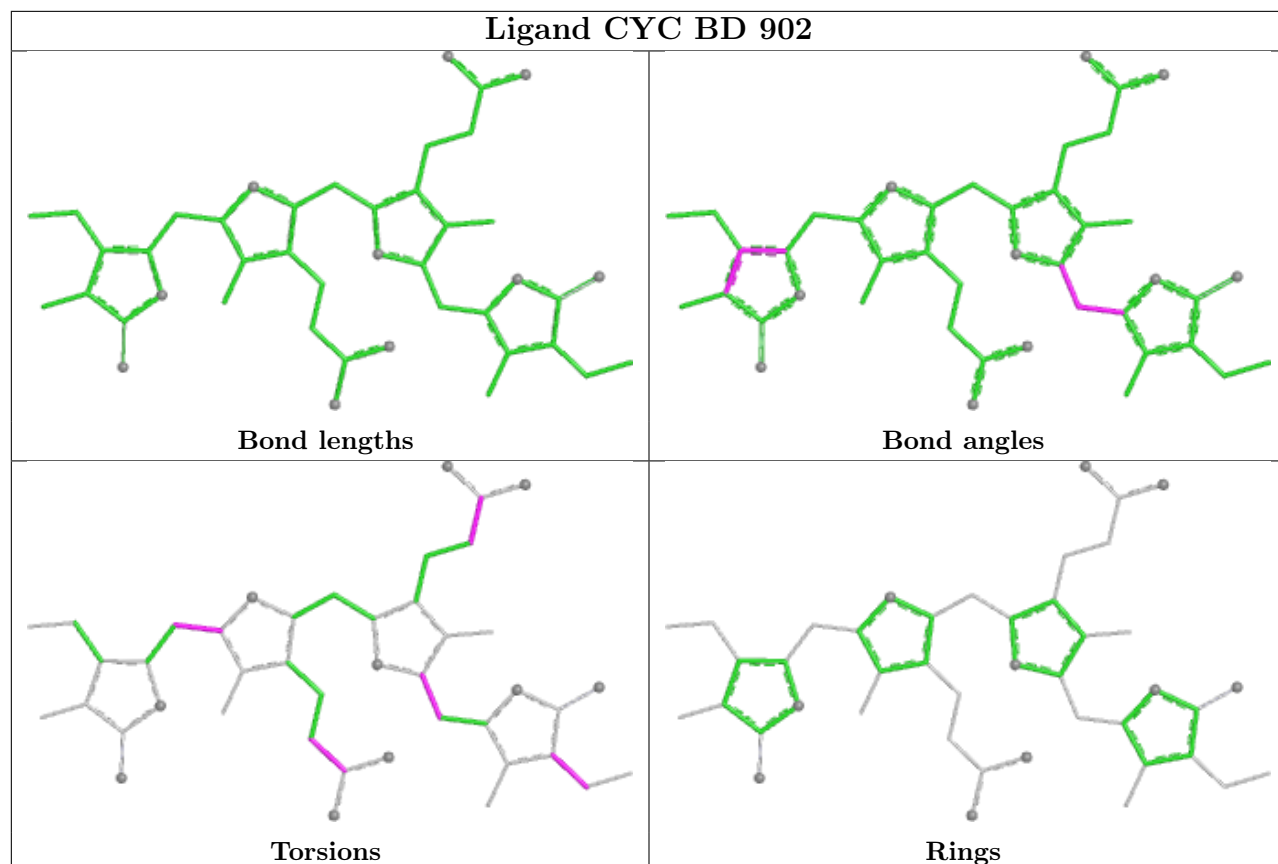
Ligand CYC BX 200



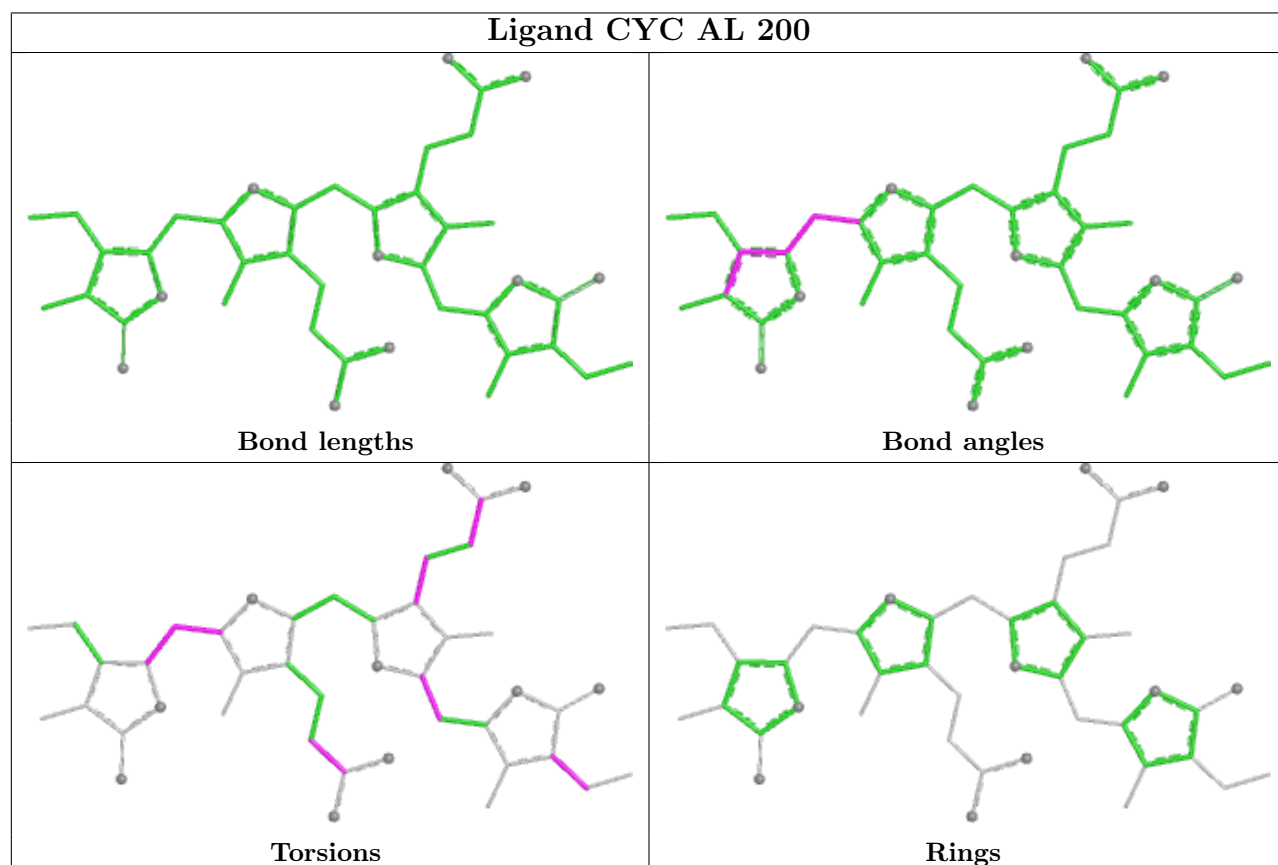
Ligand CYC AH 200

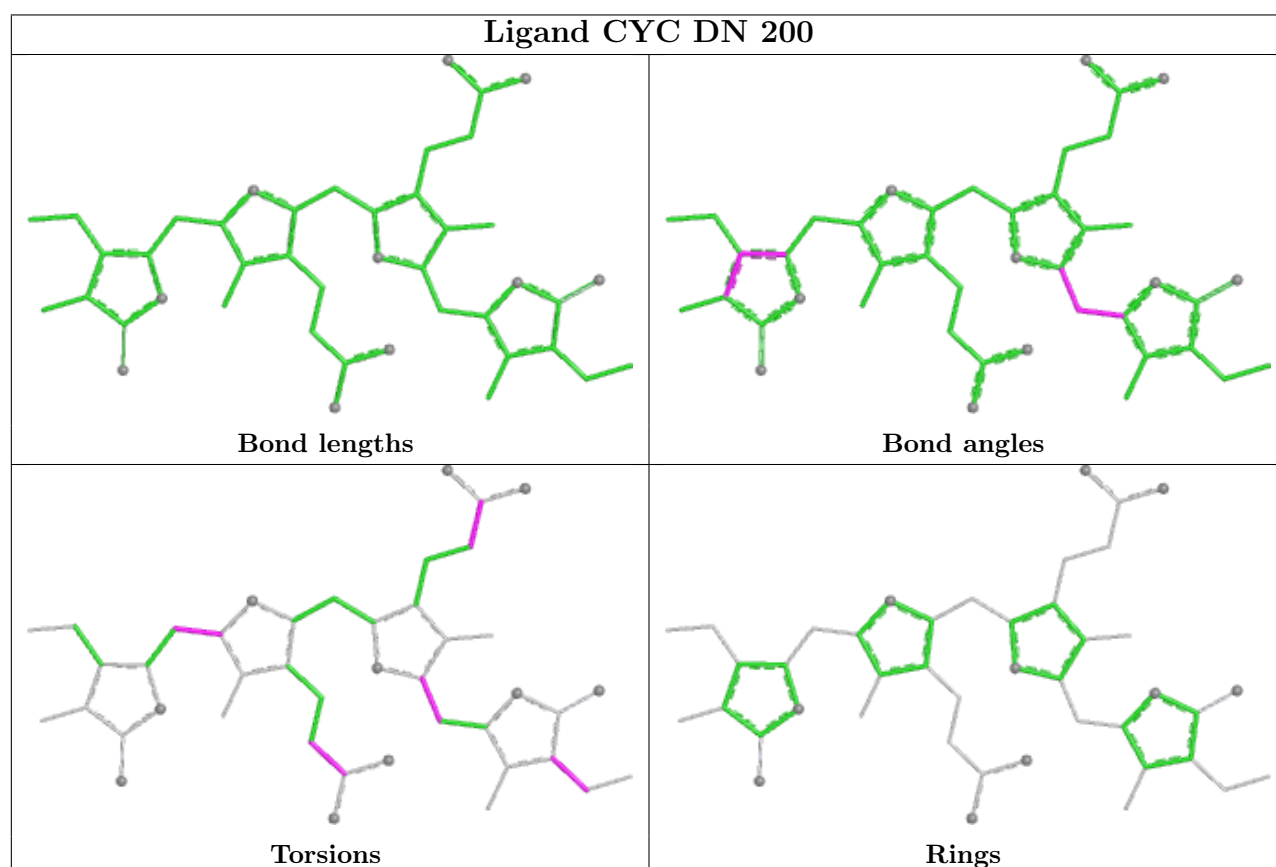
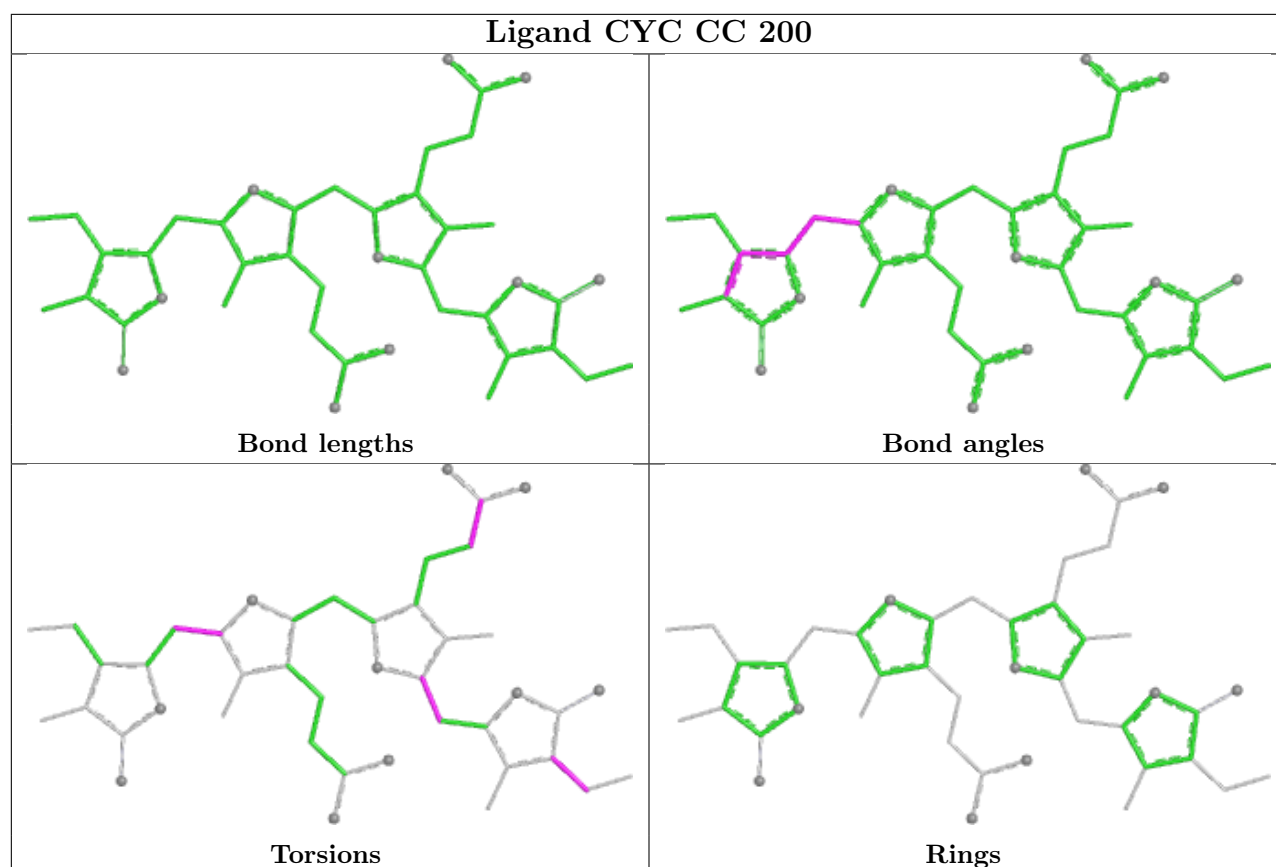


Ligand CYC BD 902

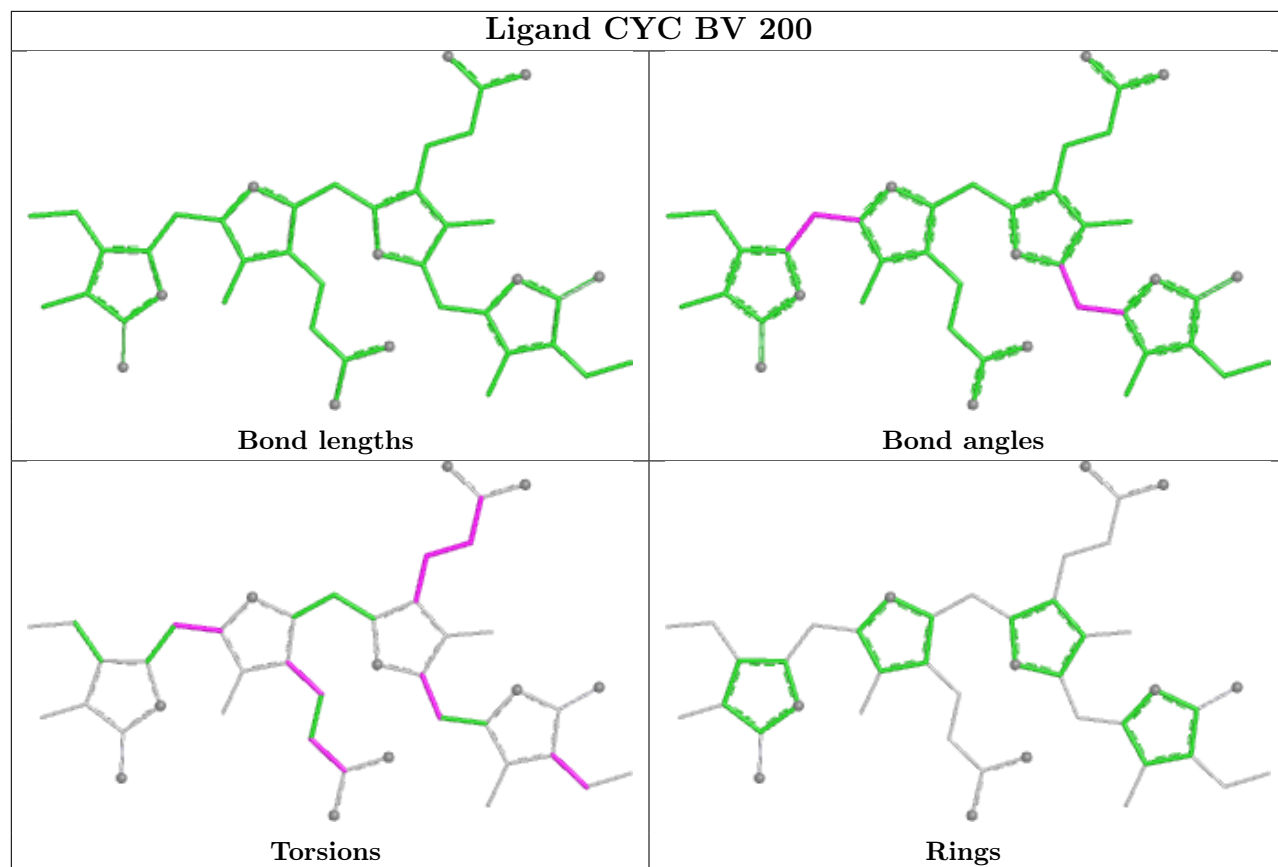


Ligand CYC AL 200

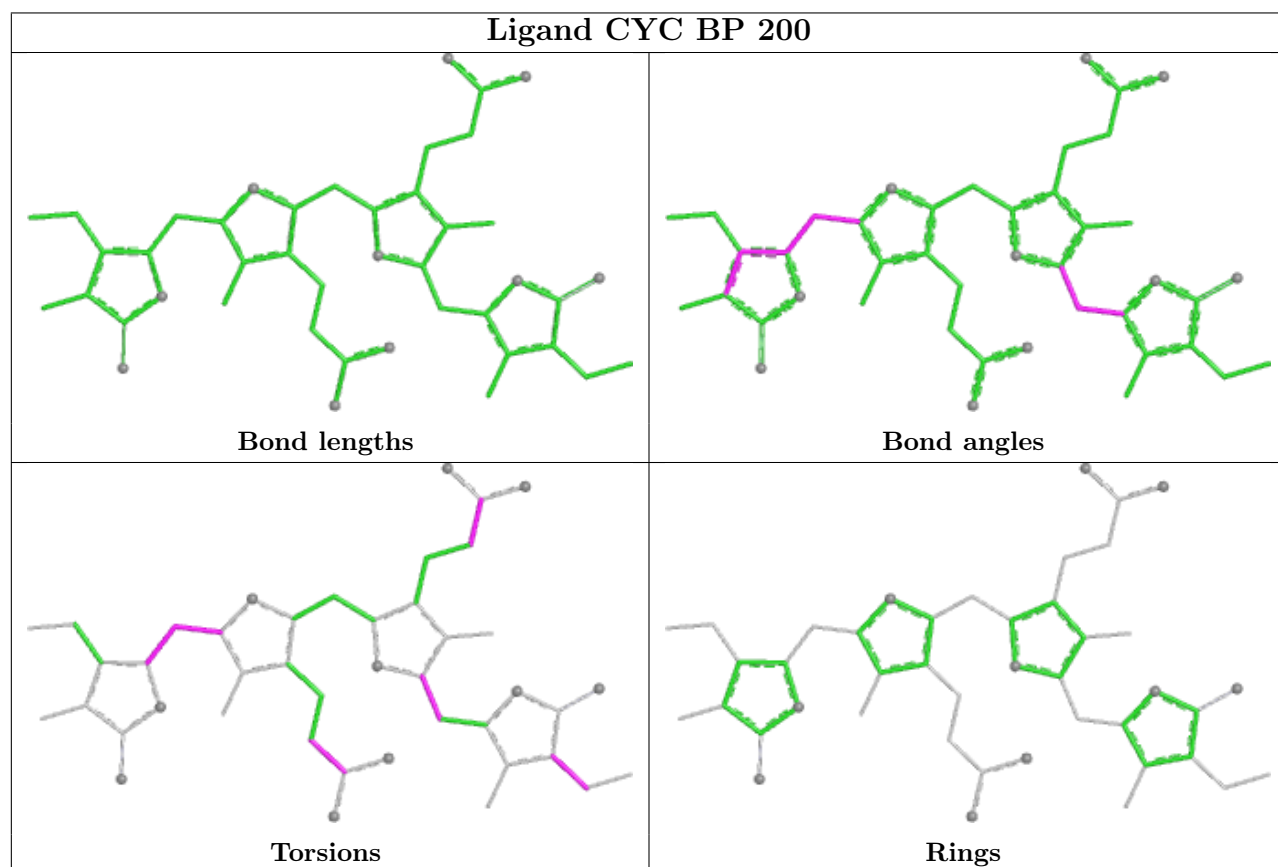




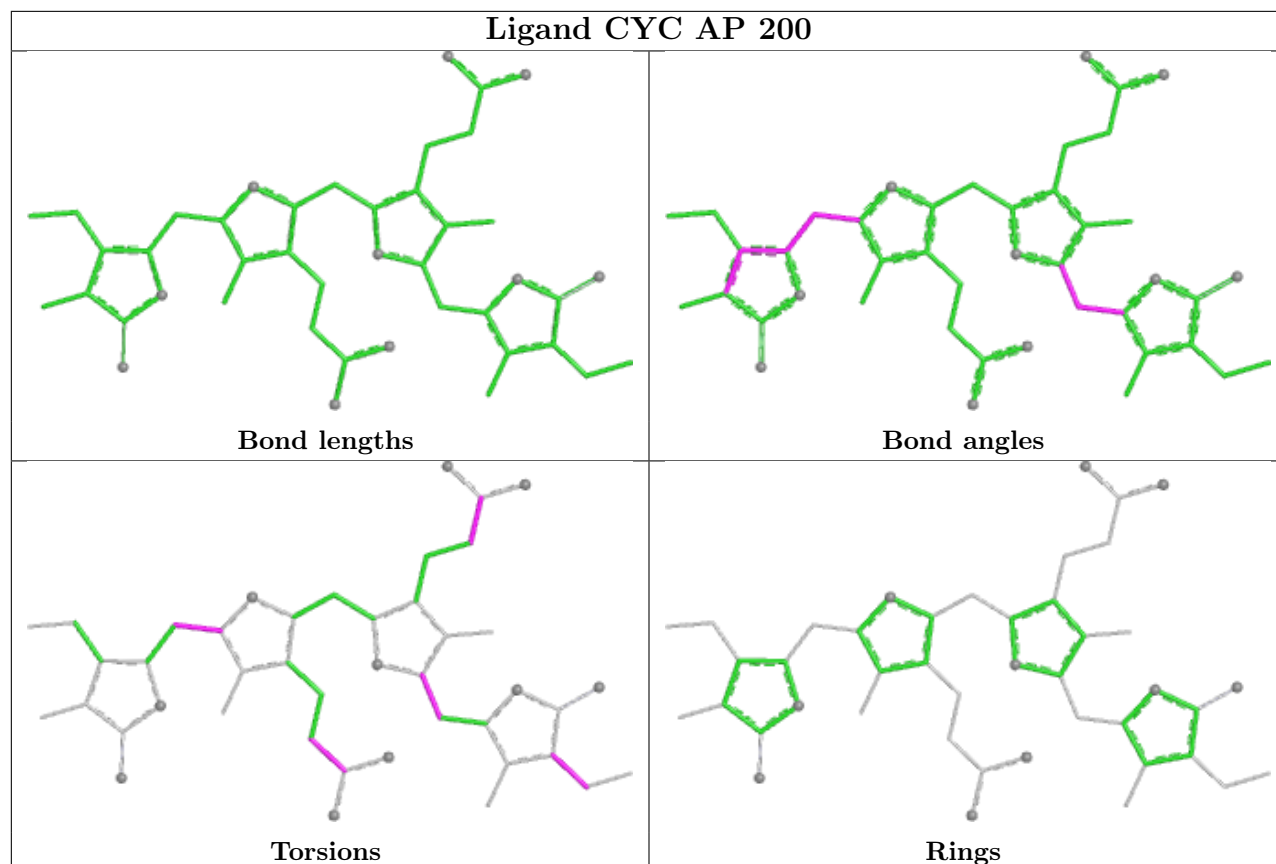
Ligand CYC BV 200



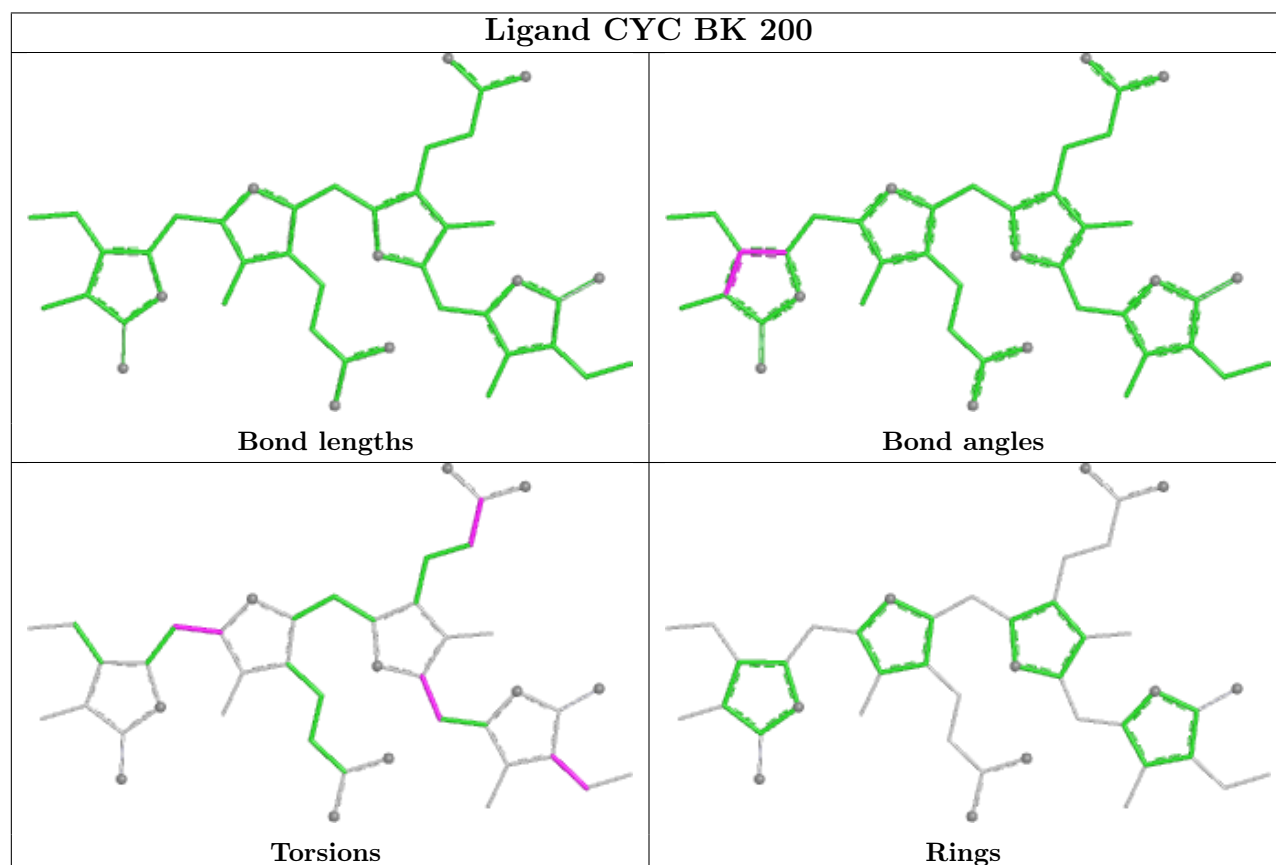
Ligand CYC BP 200



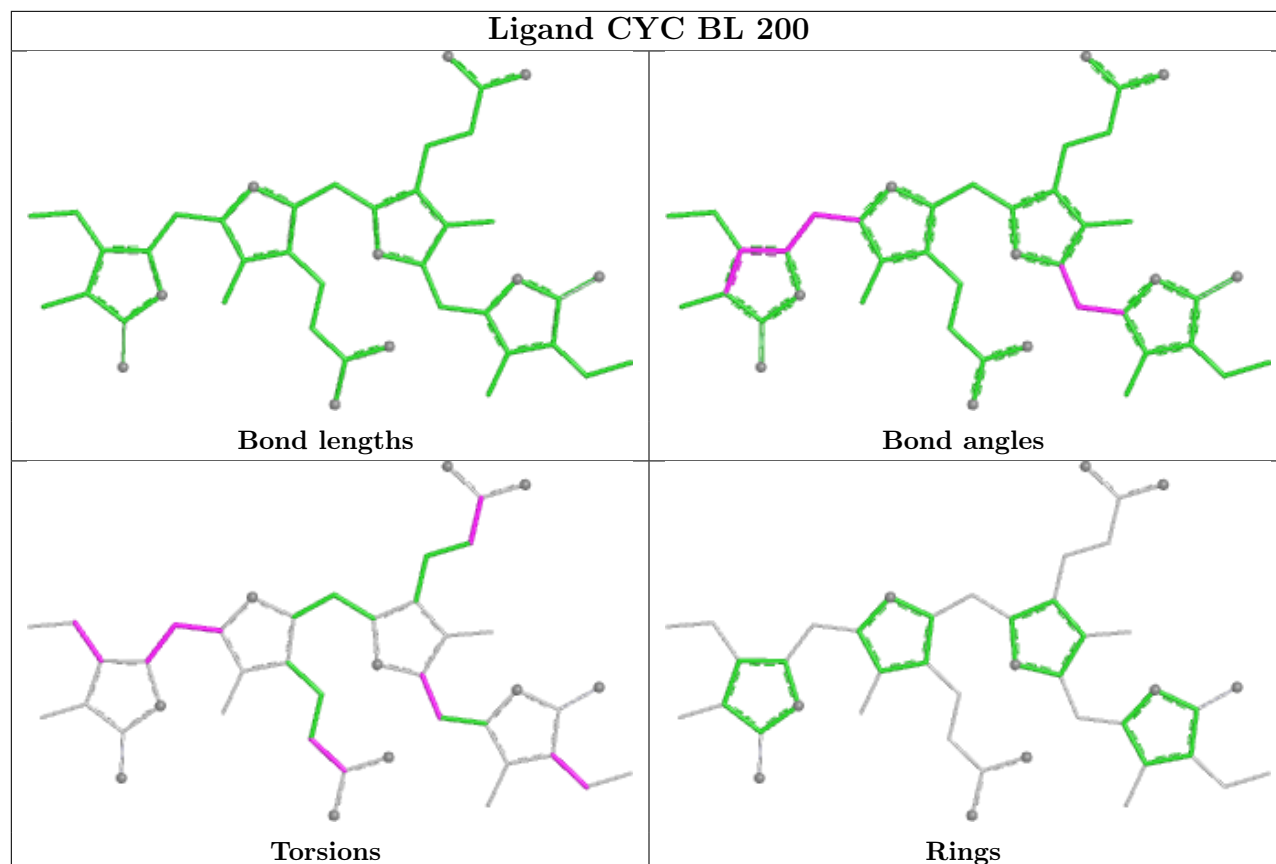
Ligand CYC AP 200



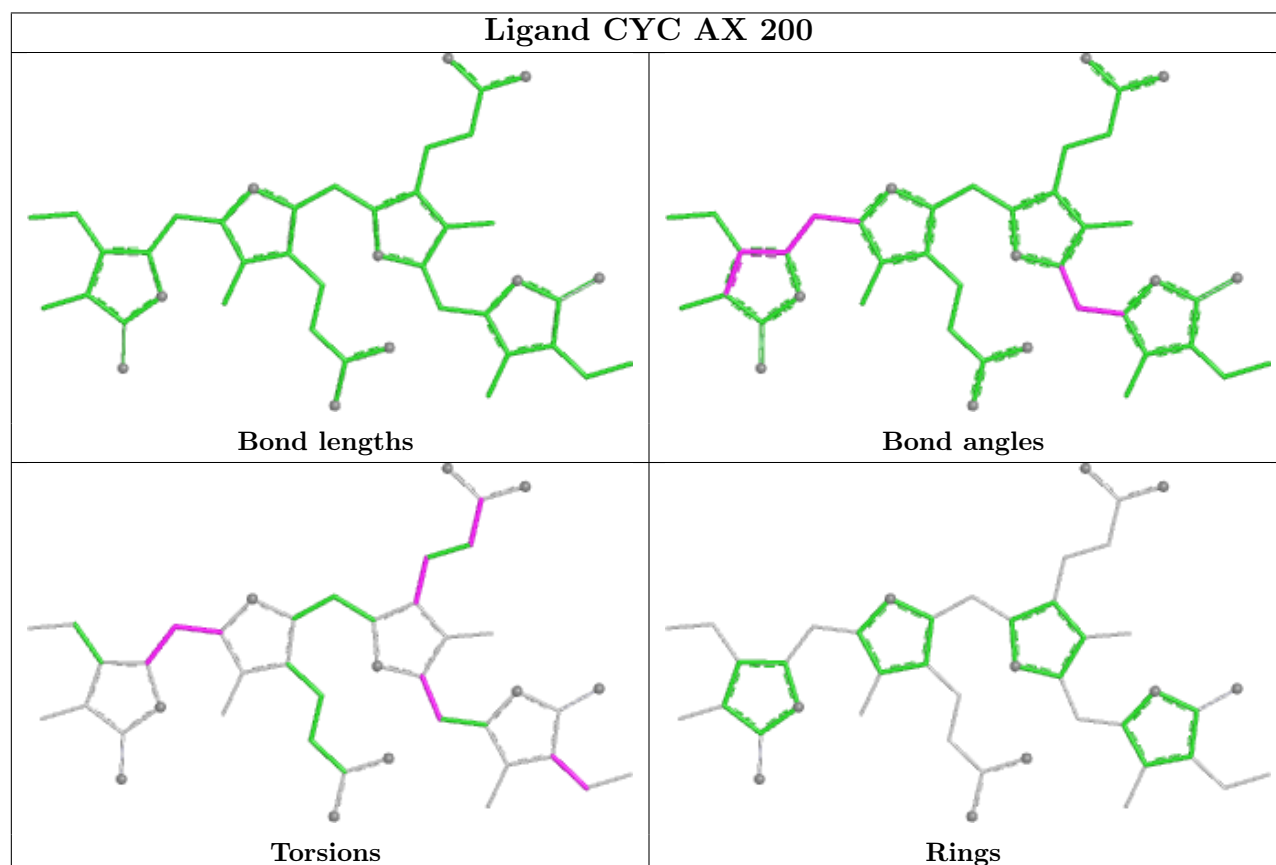
Ligand CYC BK 200



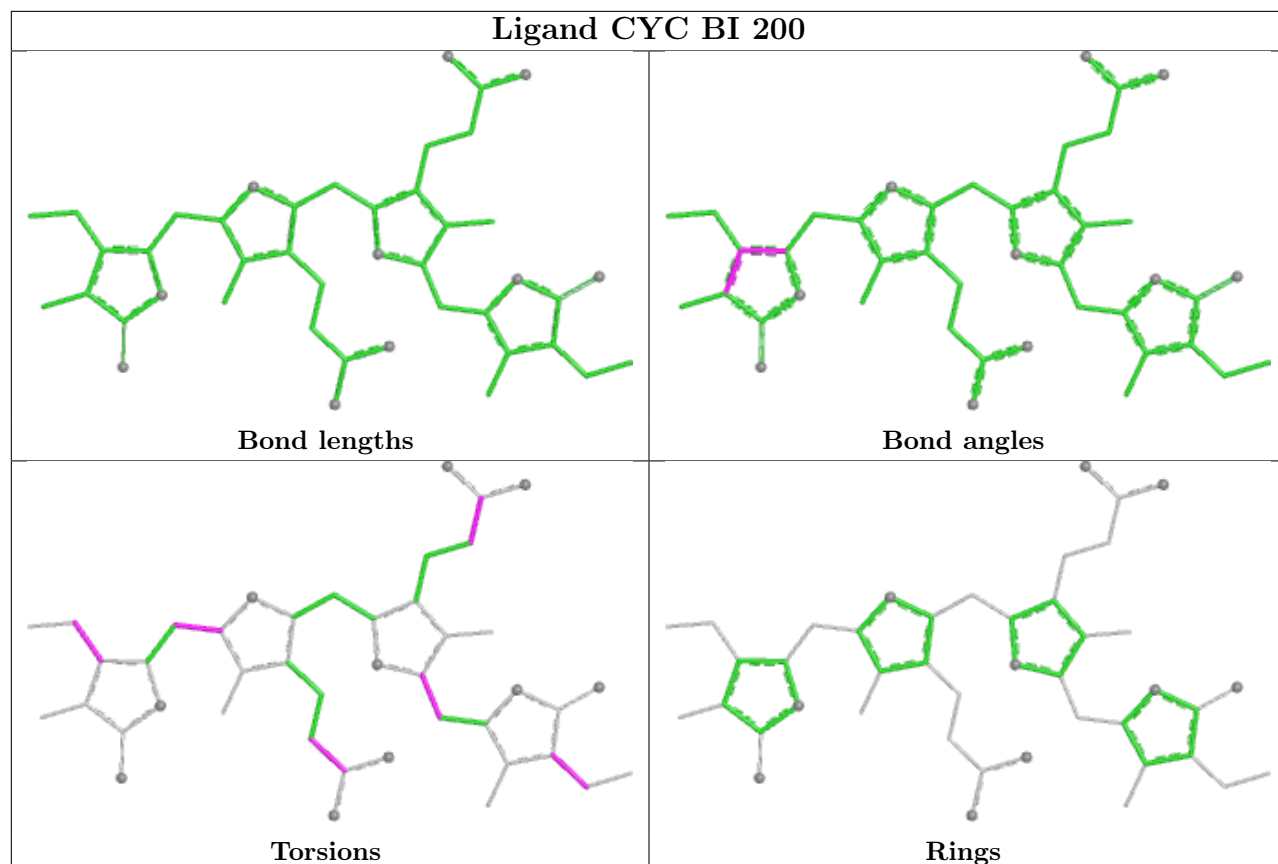
Ligand CYC BL 200



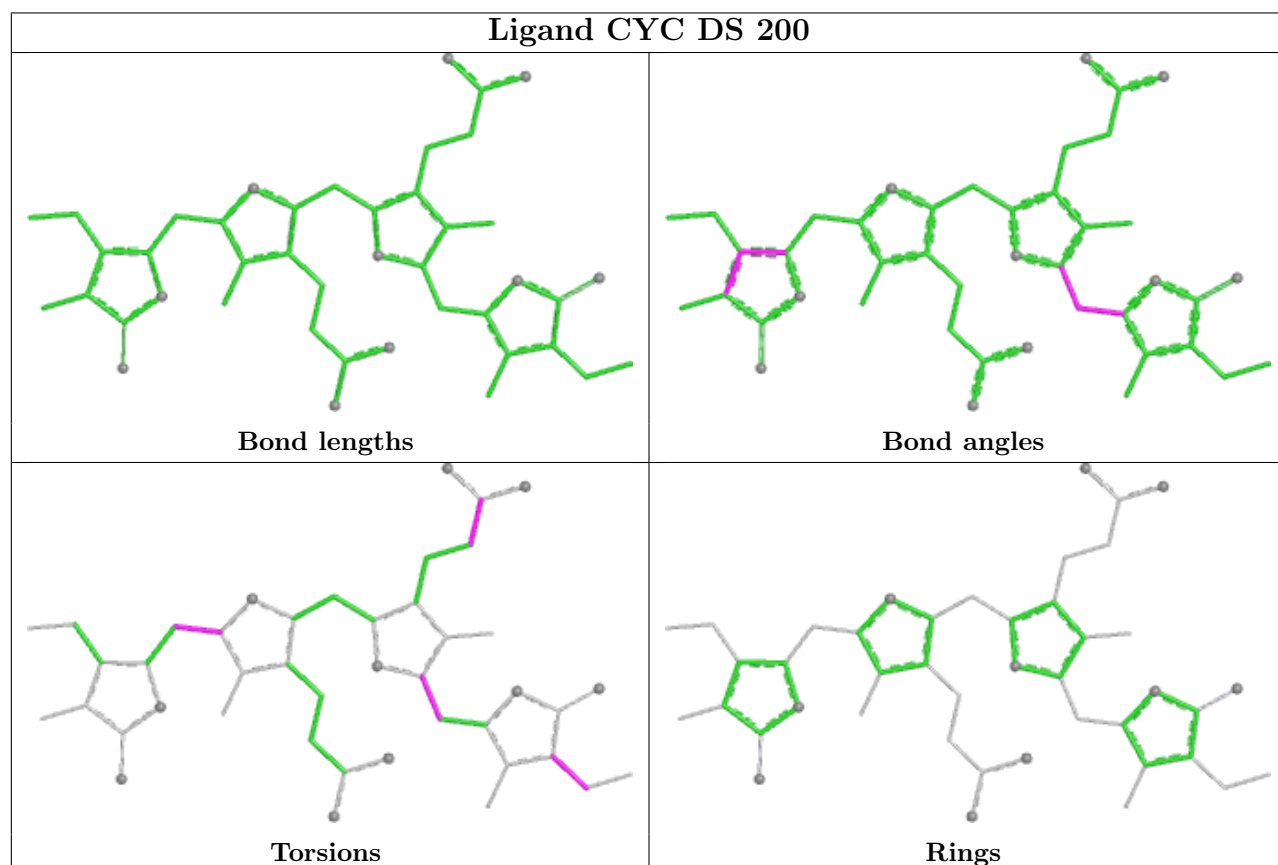
Ligand CYC AX 200

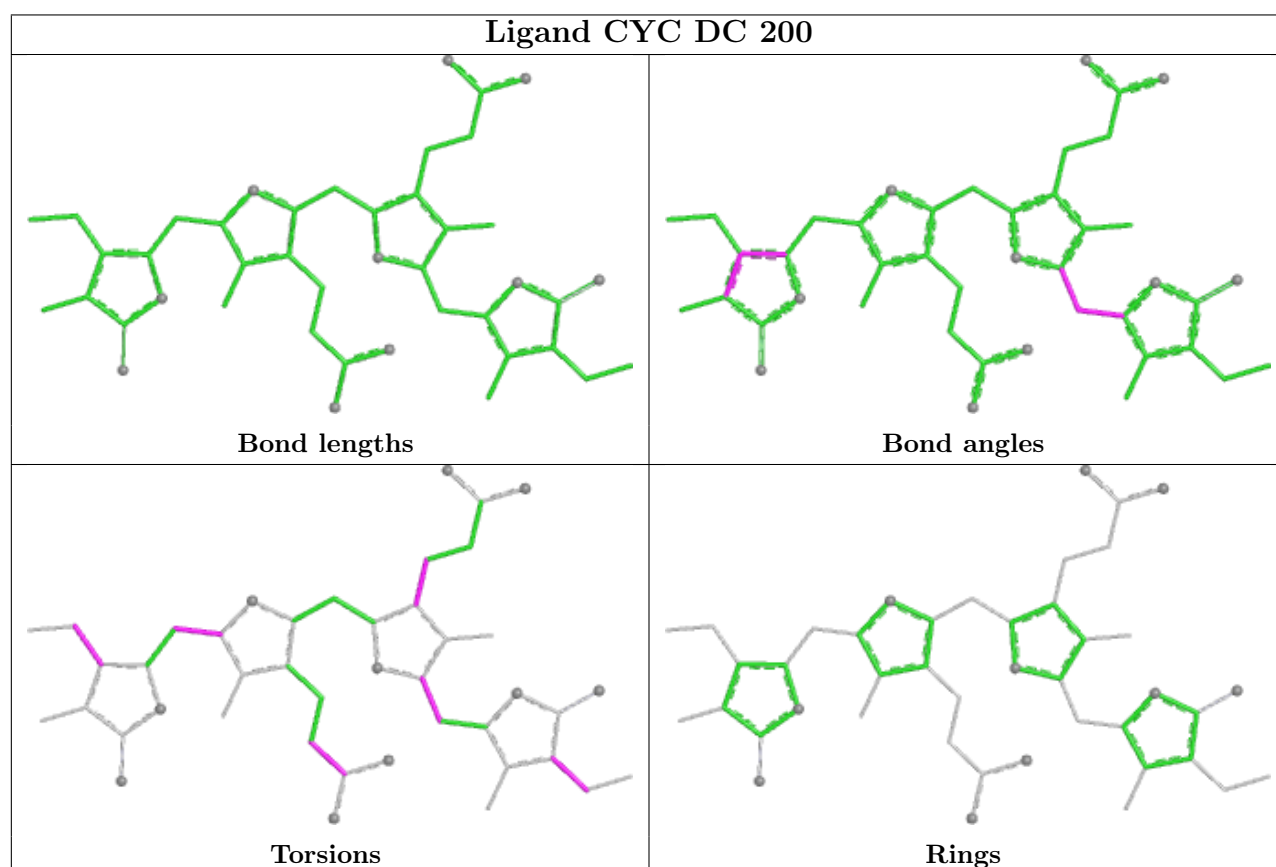
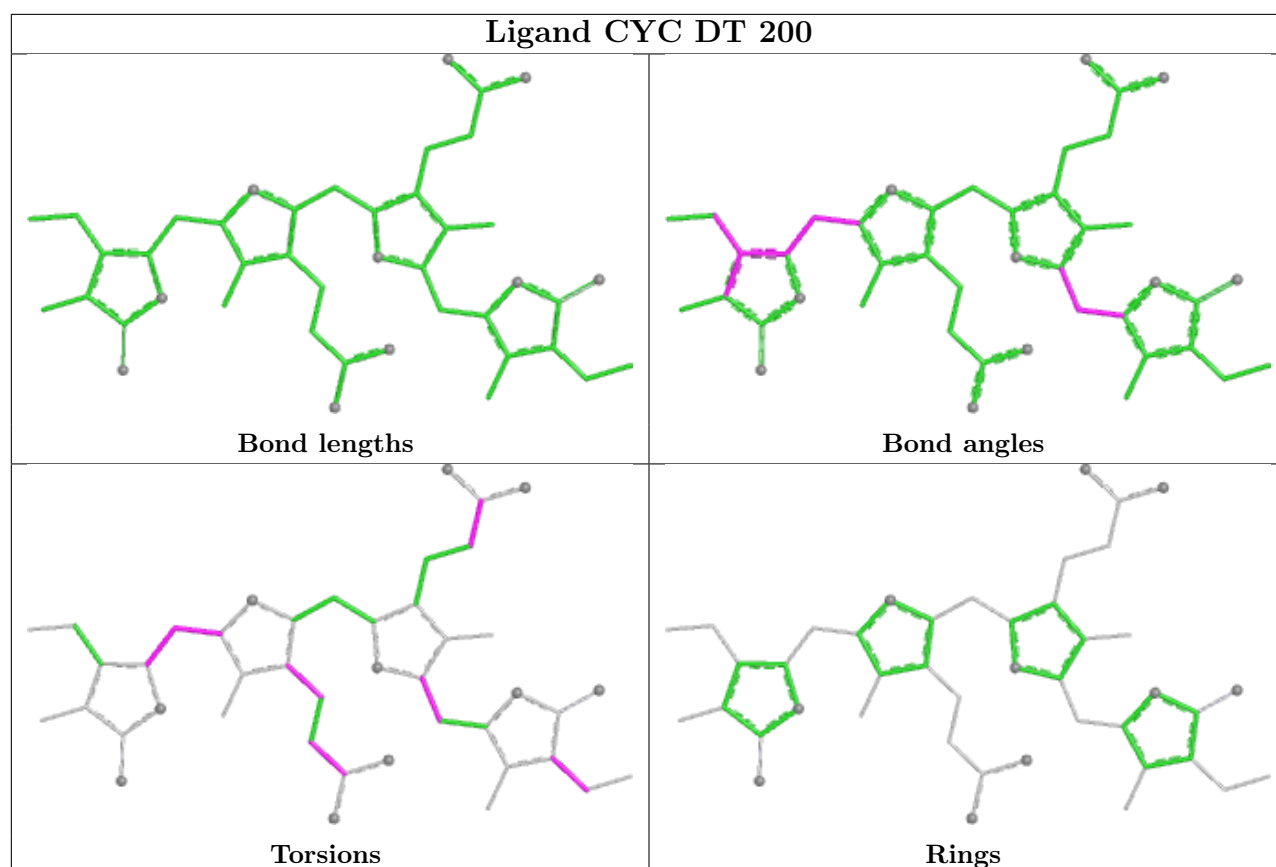


Ligand CYC BI 200

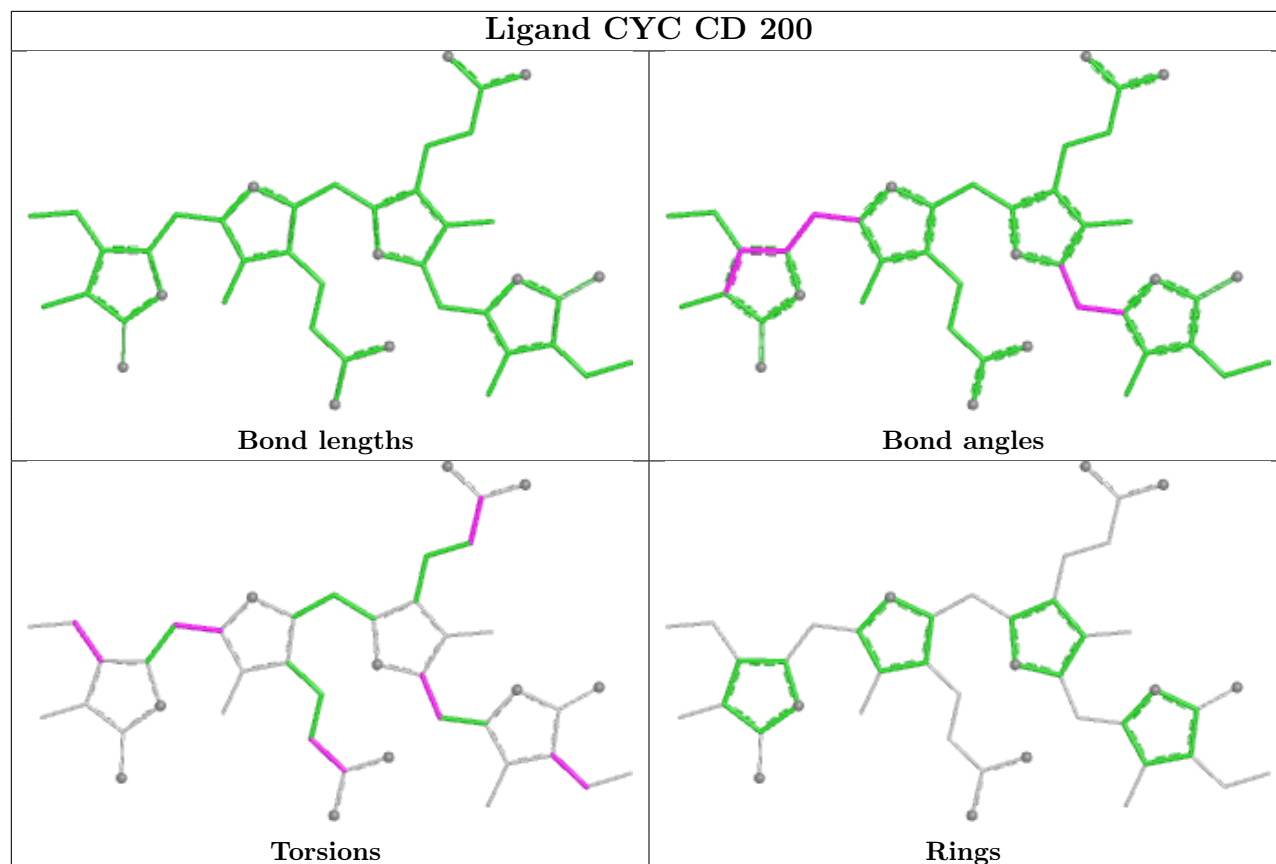


Ligand CYC DS 200

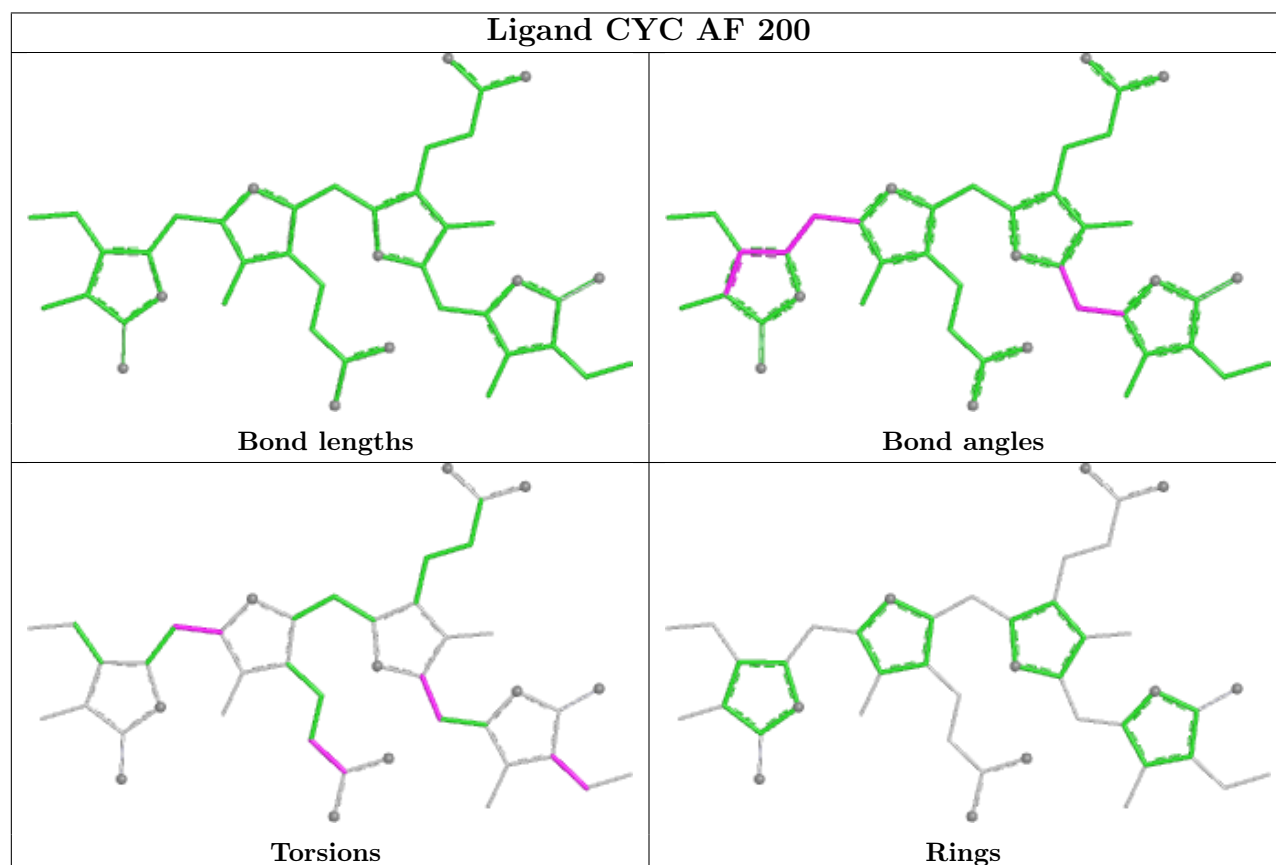




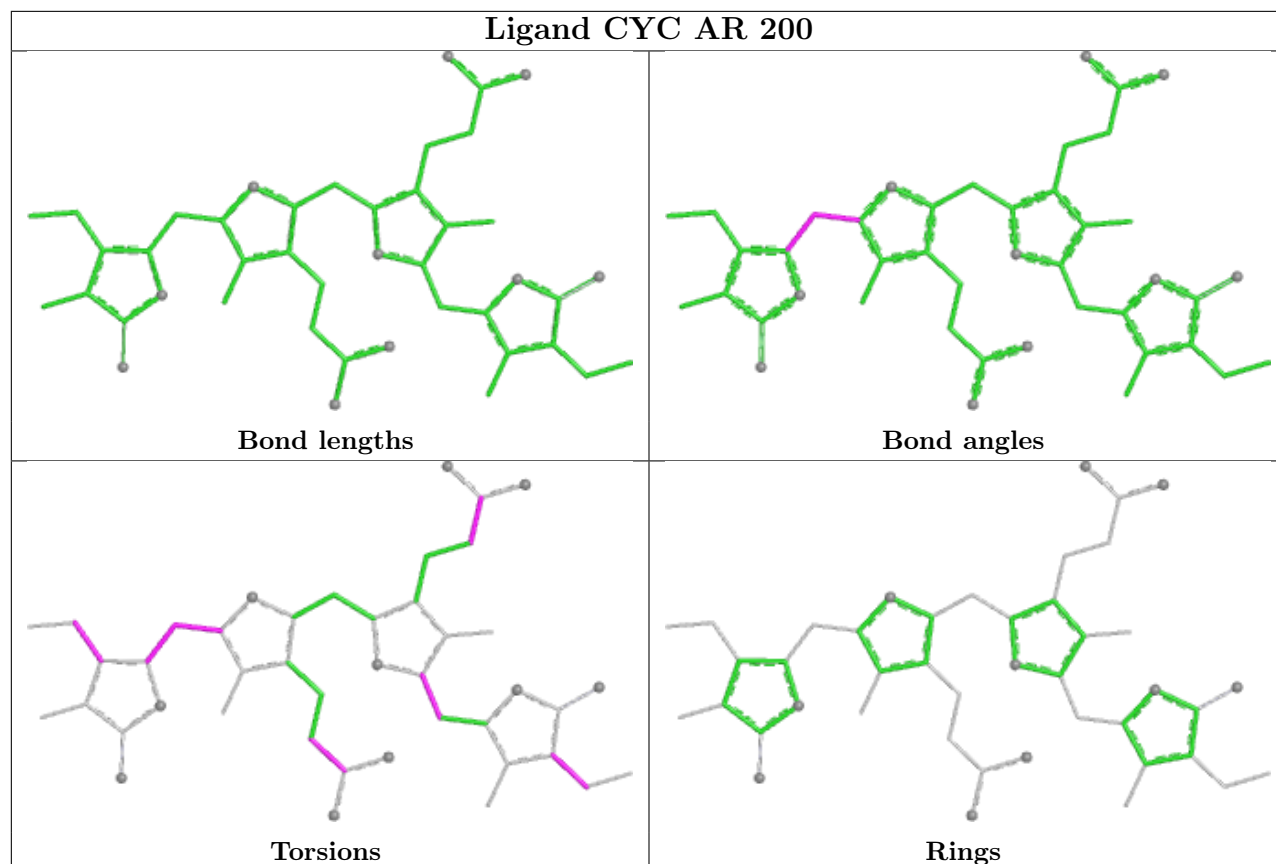
Ligand CYC CD 200



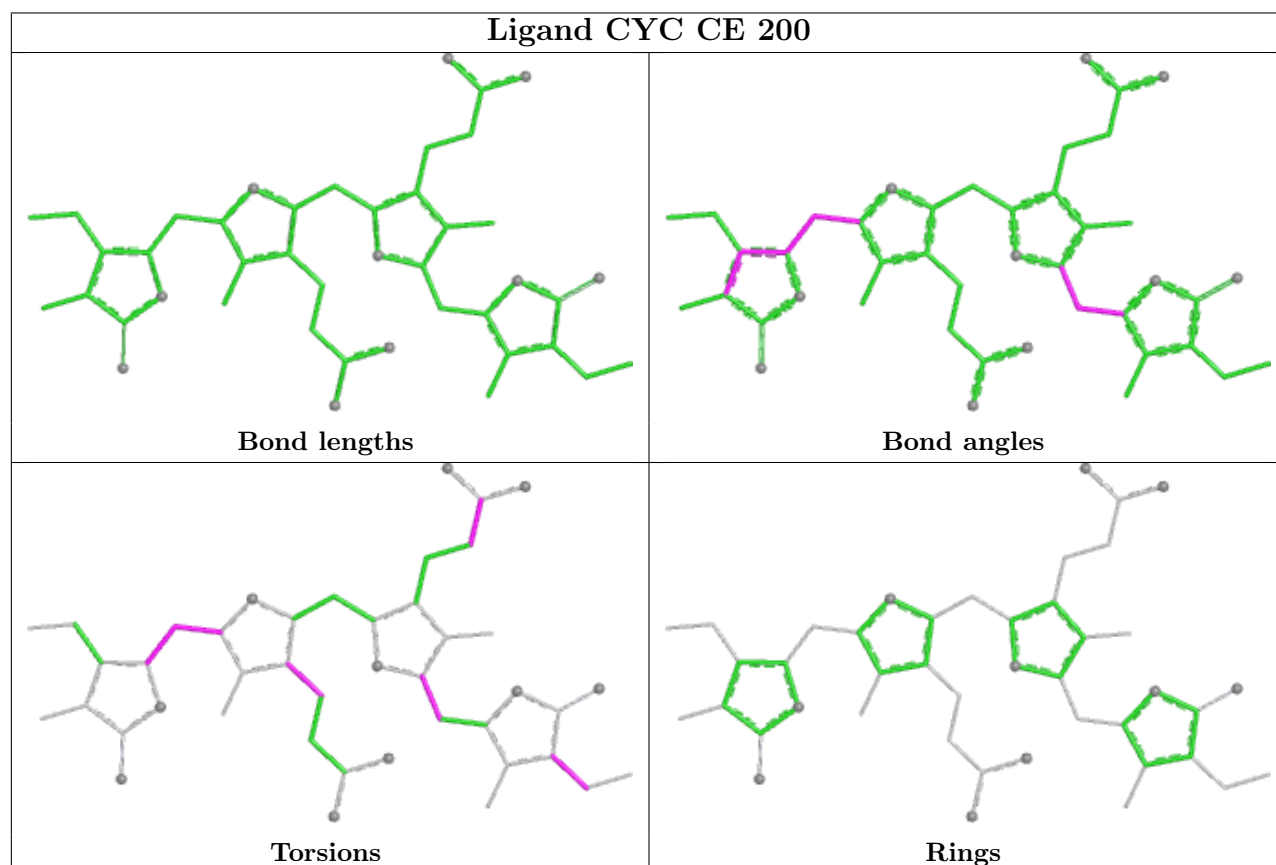
Ligand CYC AF 200



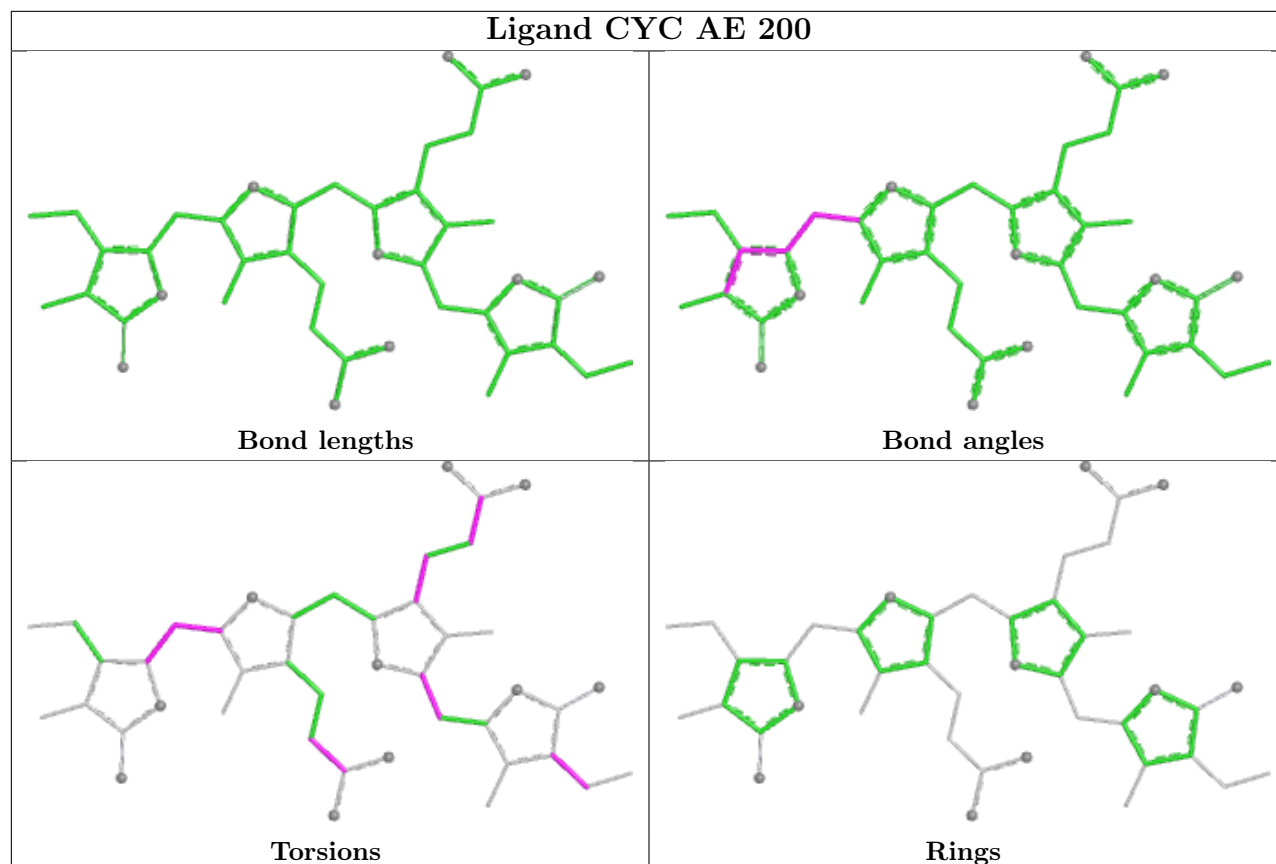
Ligand CYC AR 200



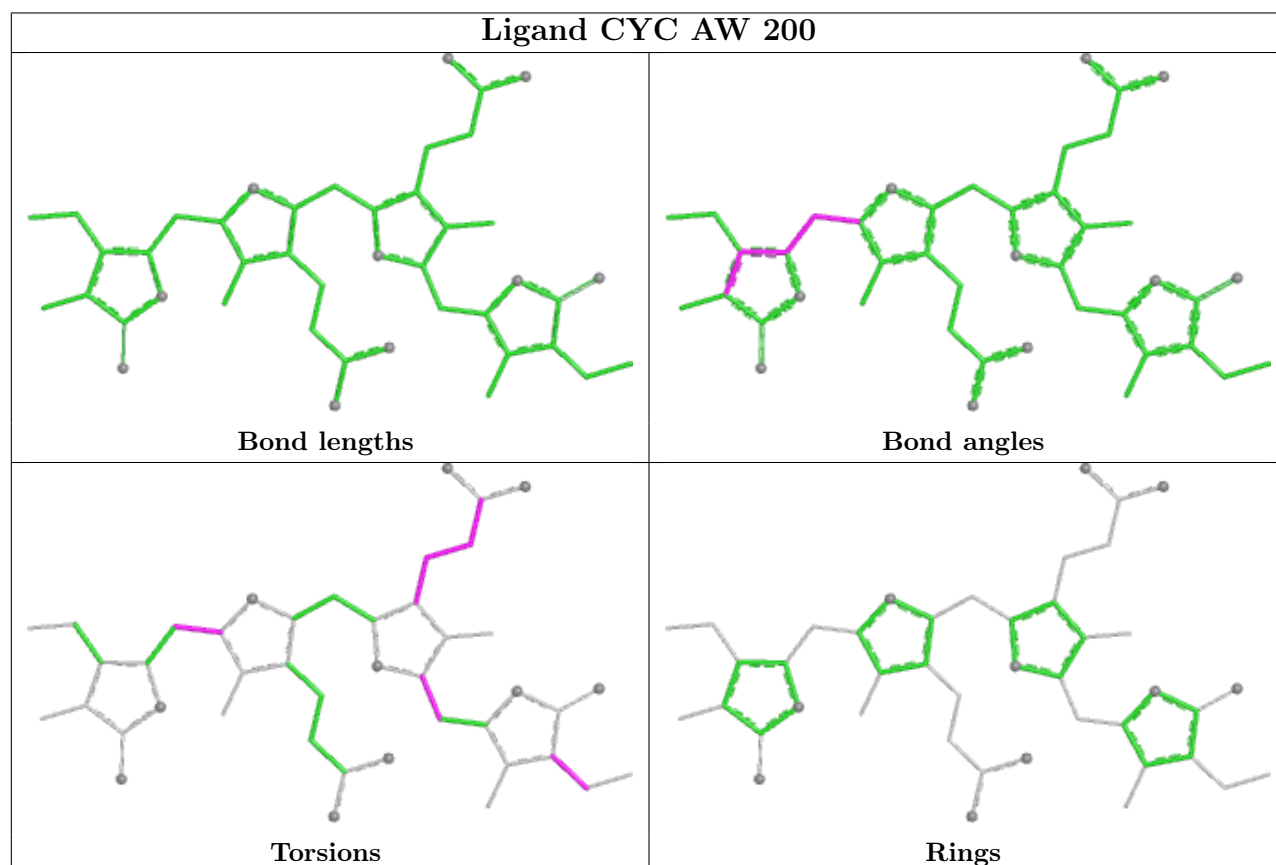
Ligand CYC CE 200



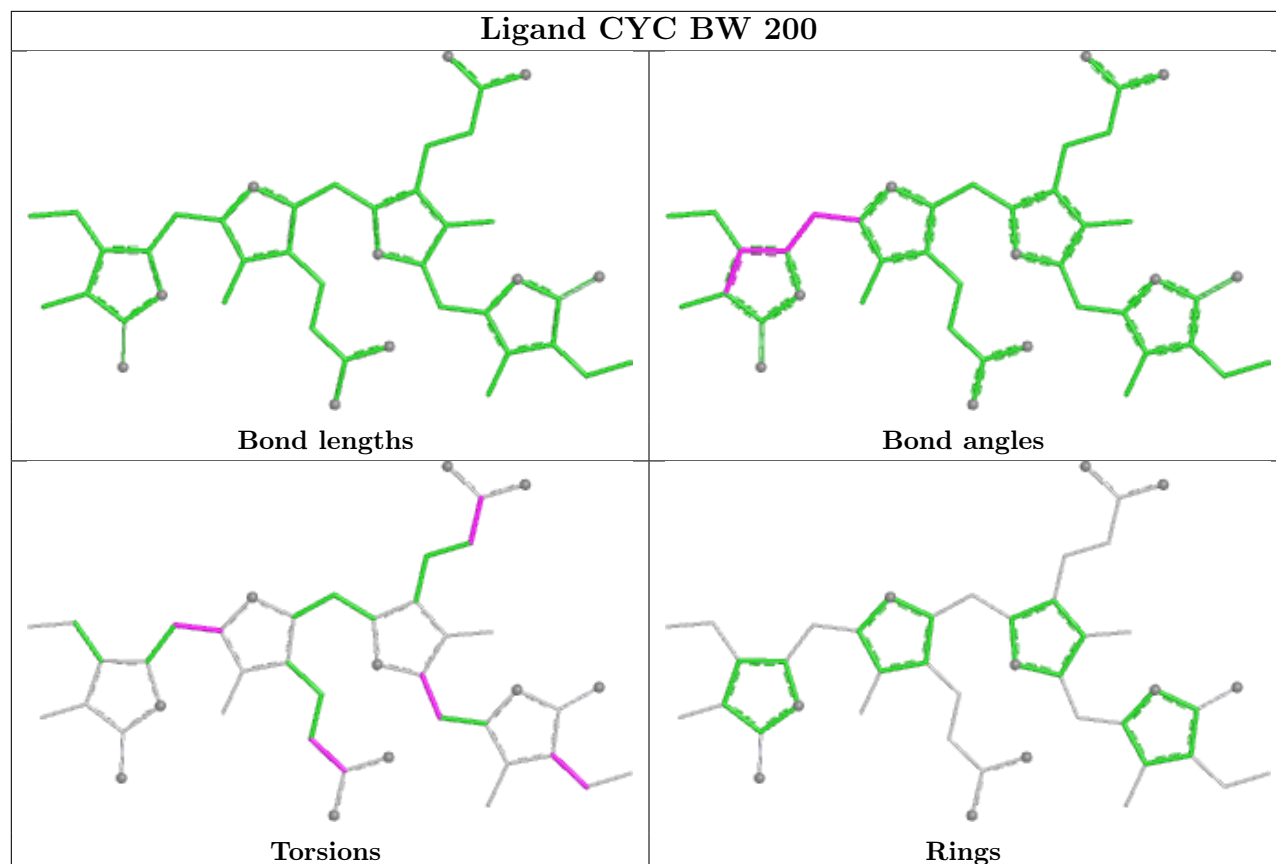
Ligand CYC AE 200



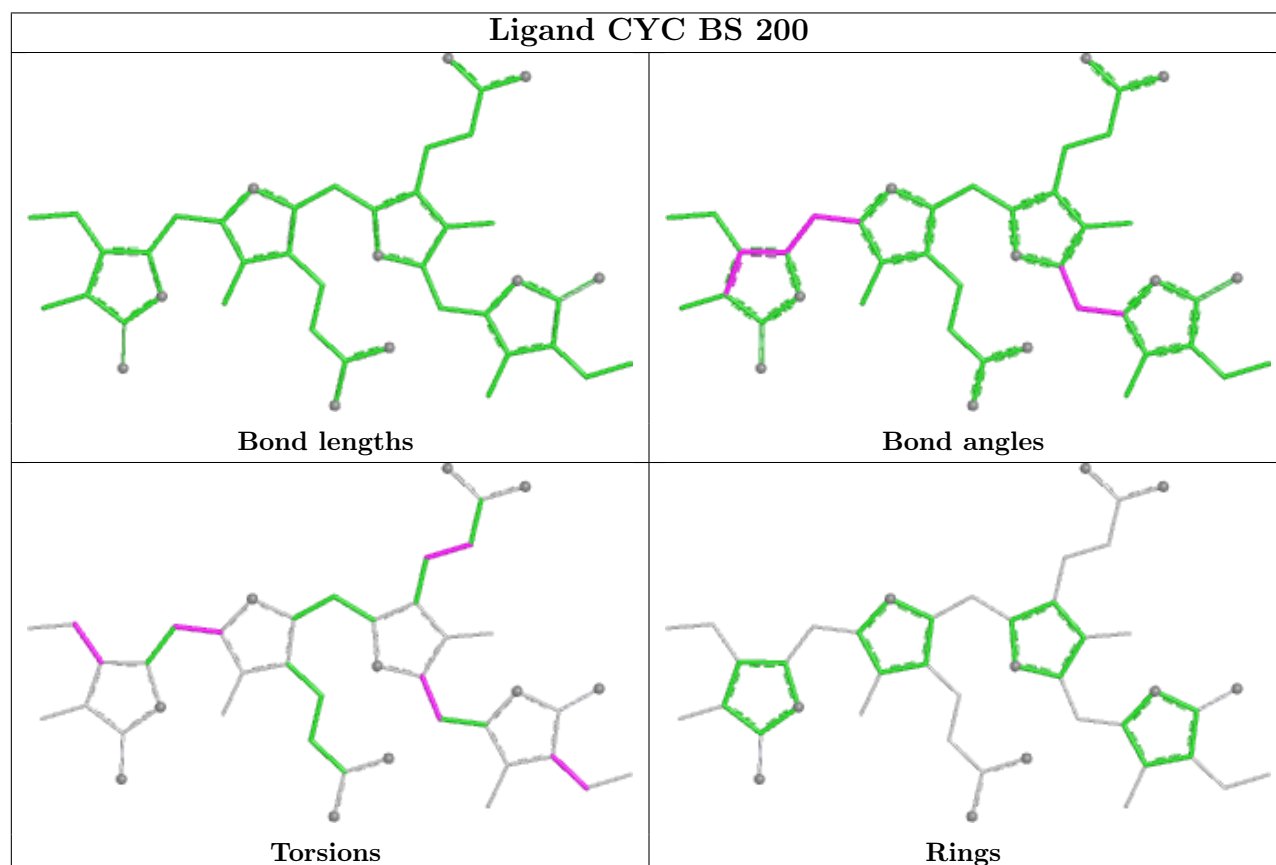
Ligand CYC AW 200



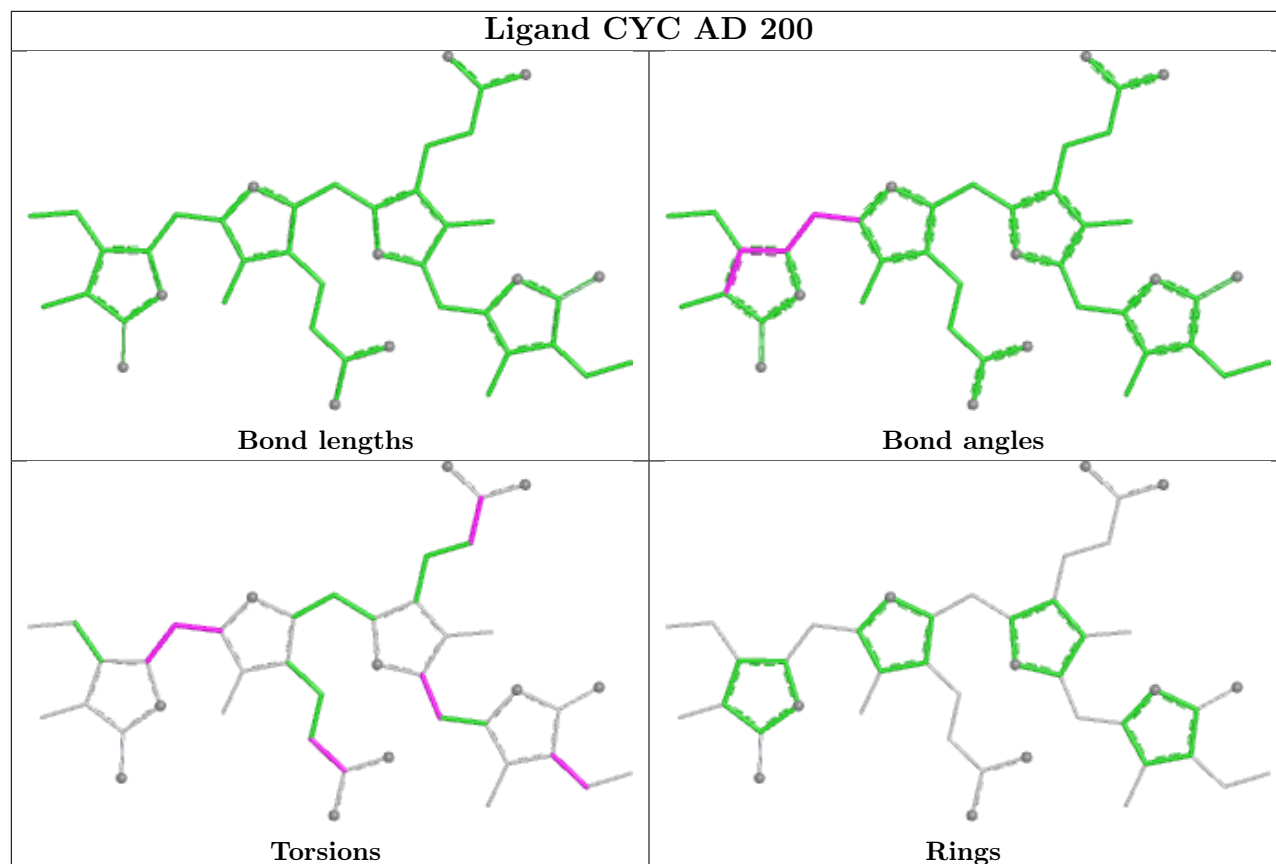
Ligand CYC BW 200



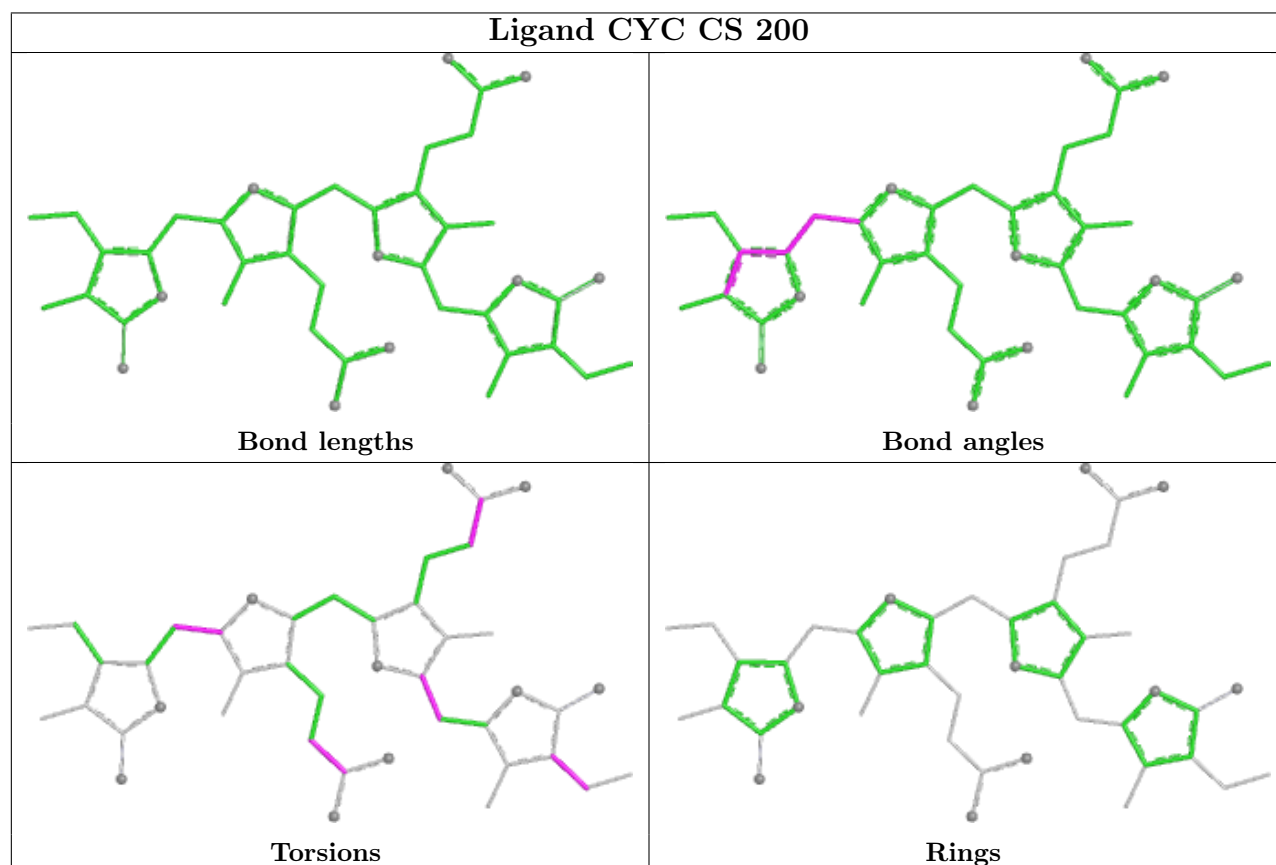
Ligand CYC BS 200



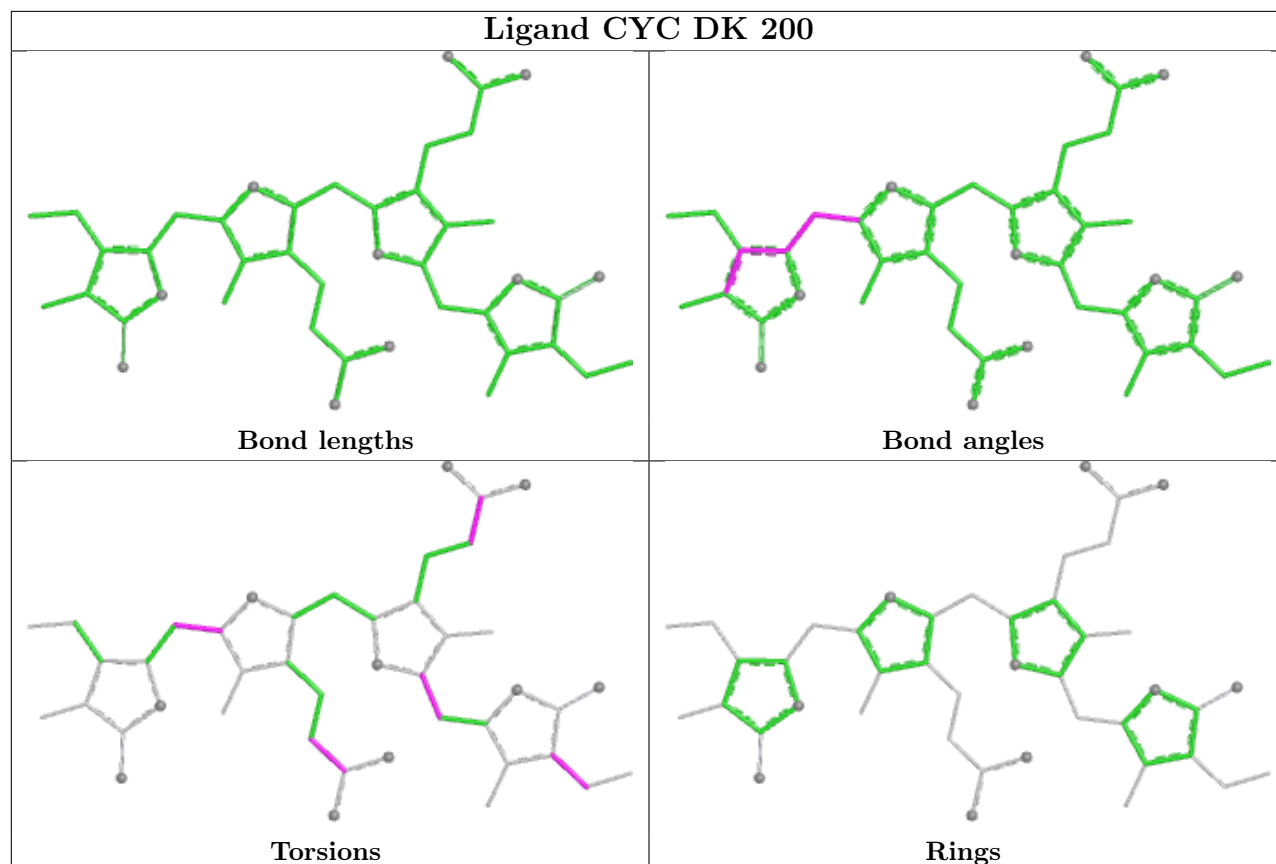
Ligand CYC AD 200



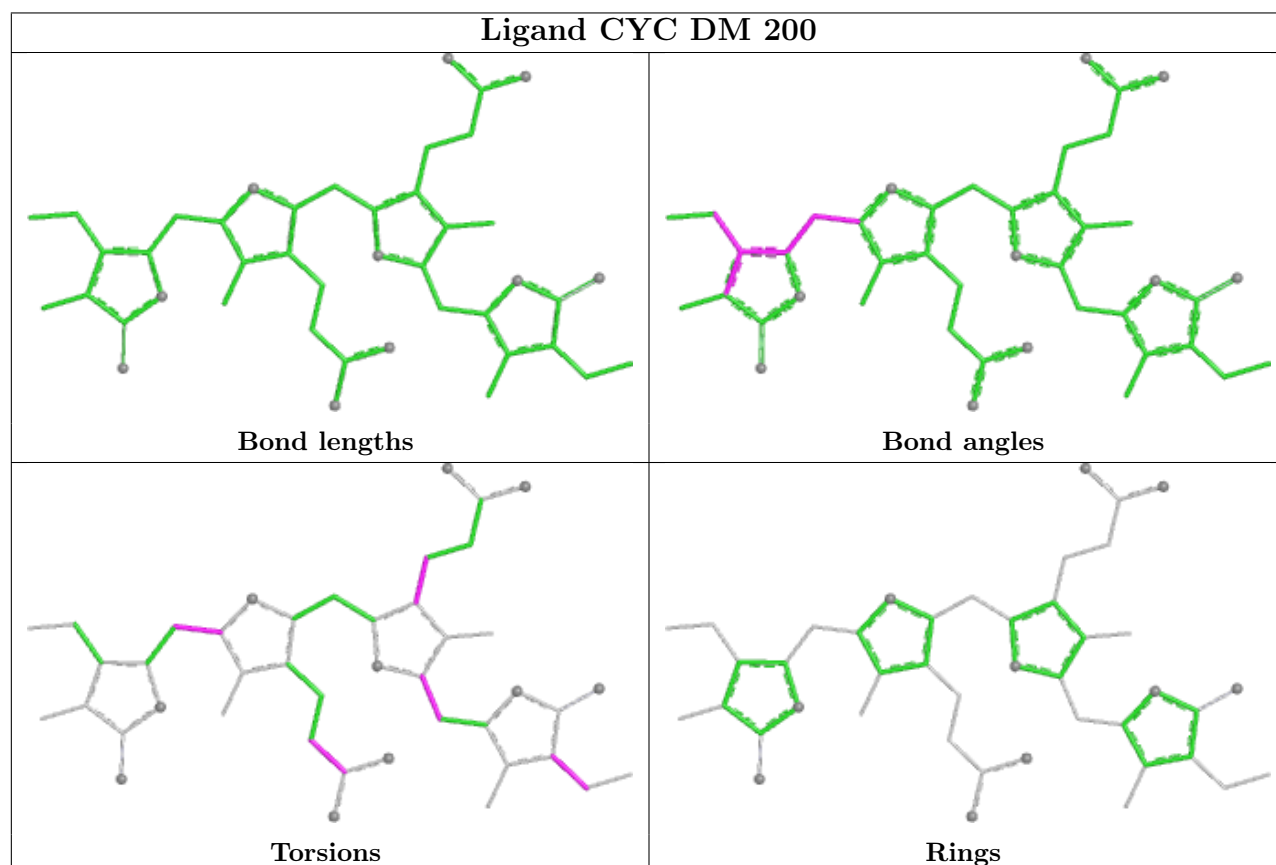
Ligand CYC CS 200



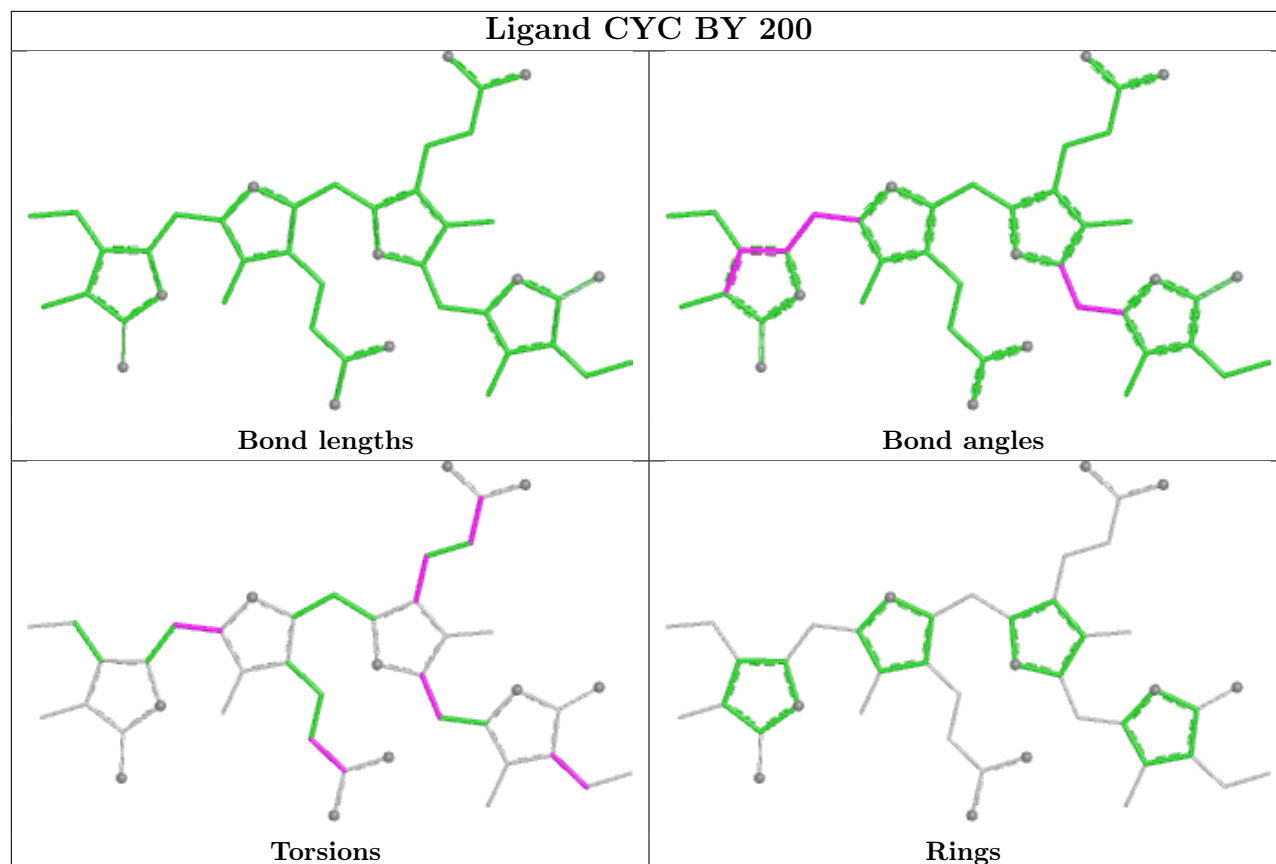
Ligand CYC DK 200



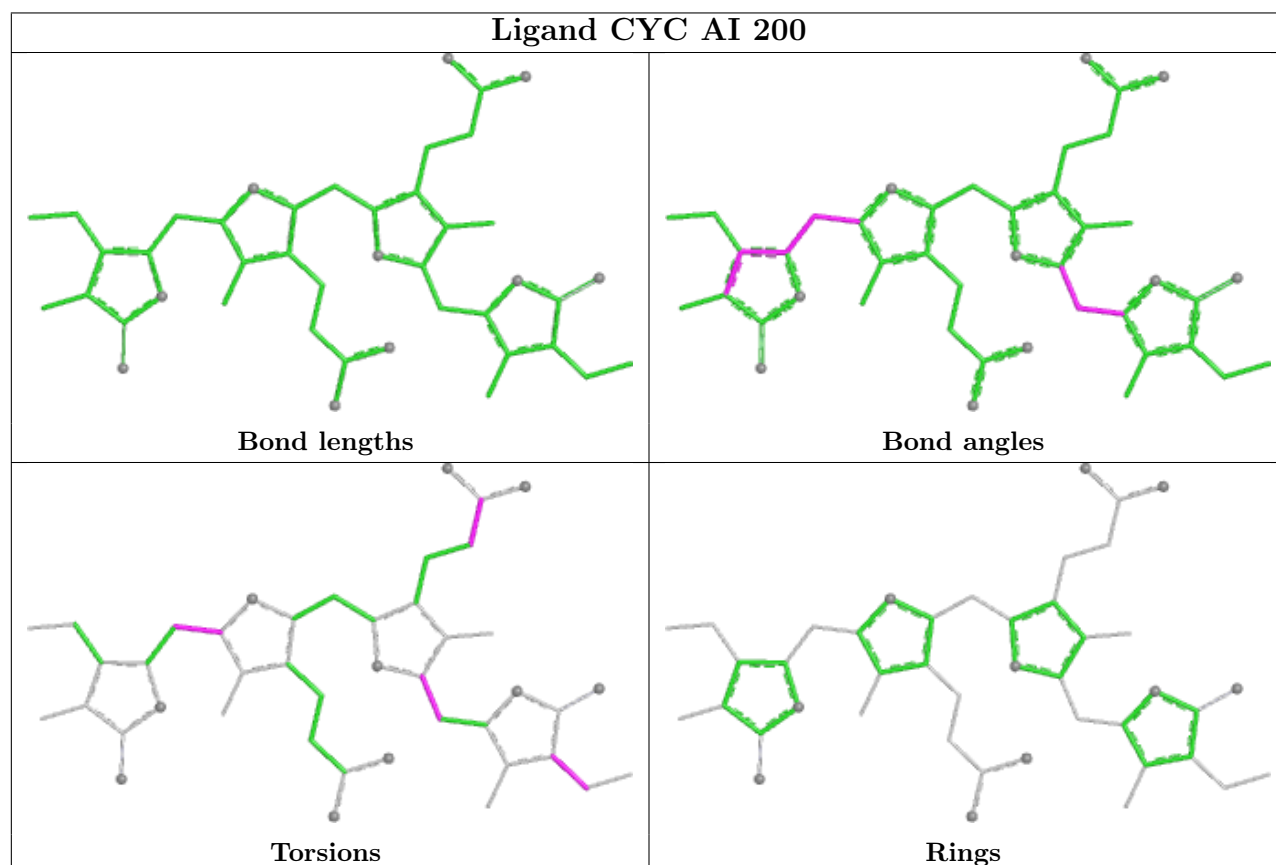
Ligand CYC DM 200



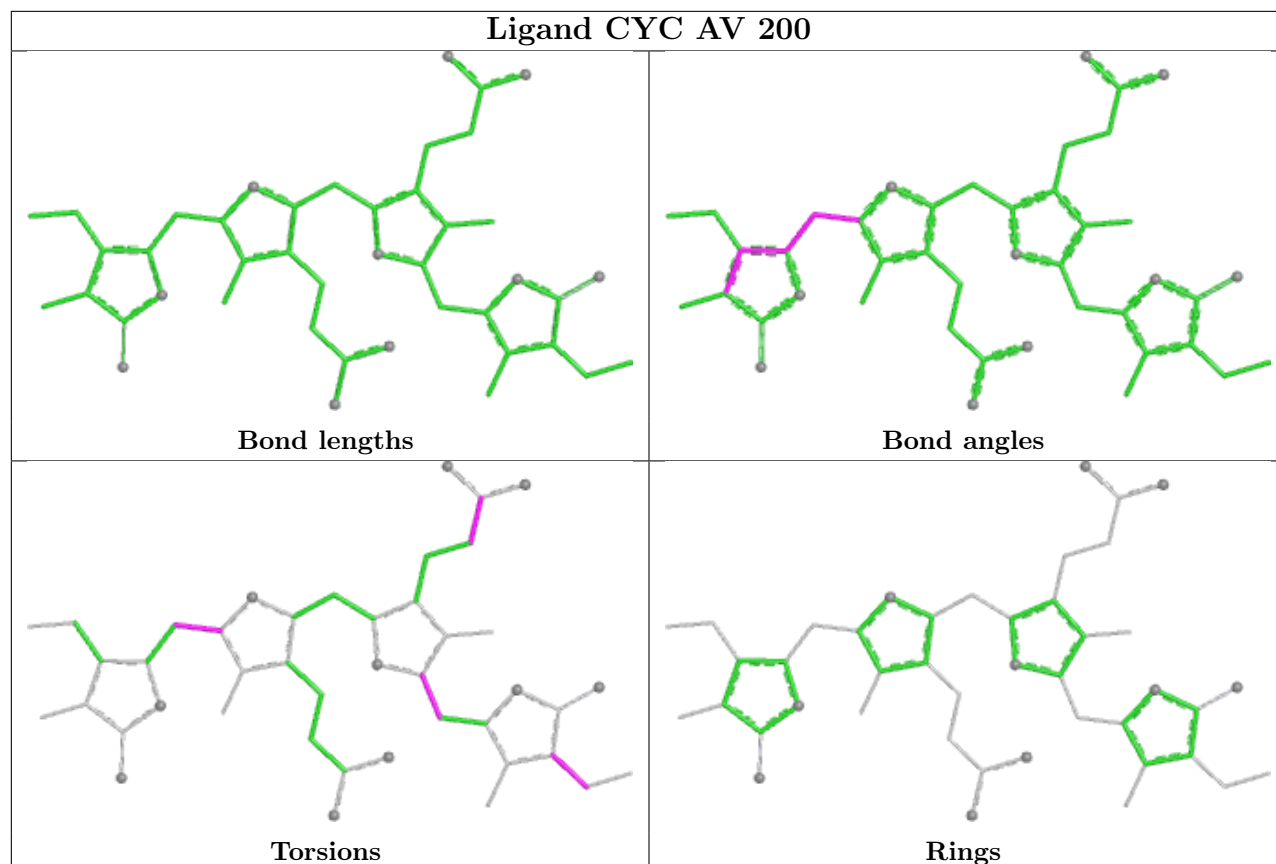
Ligand CYC BY 200



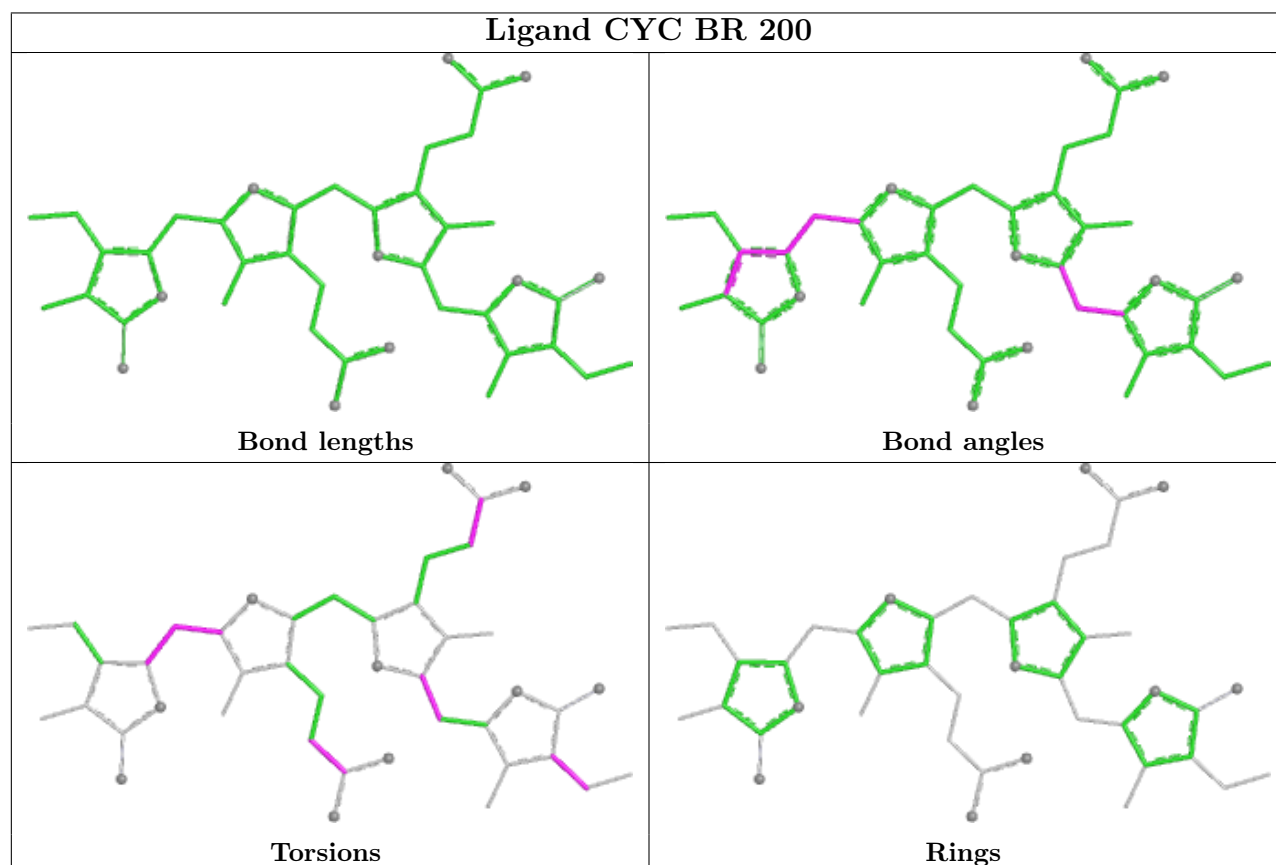
Ligand CYC AI 200

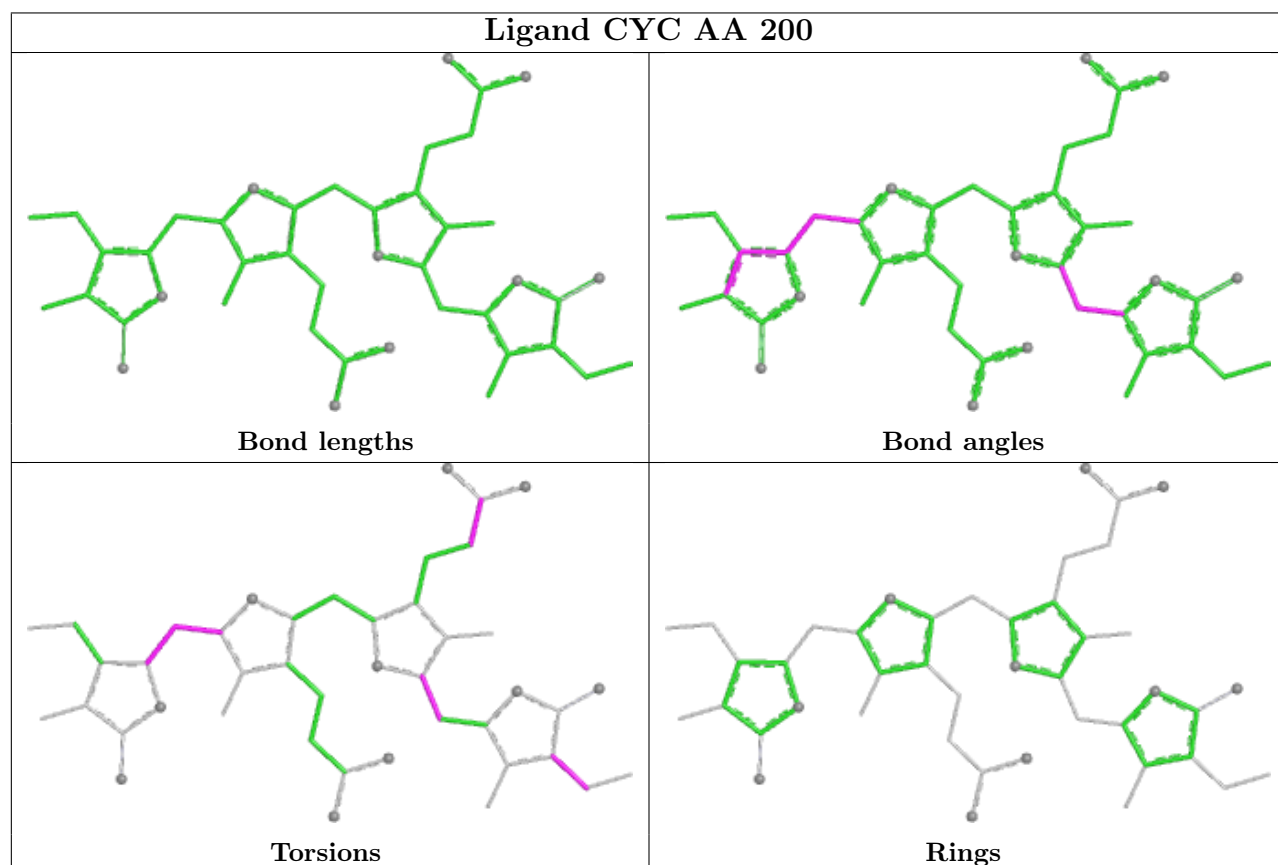
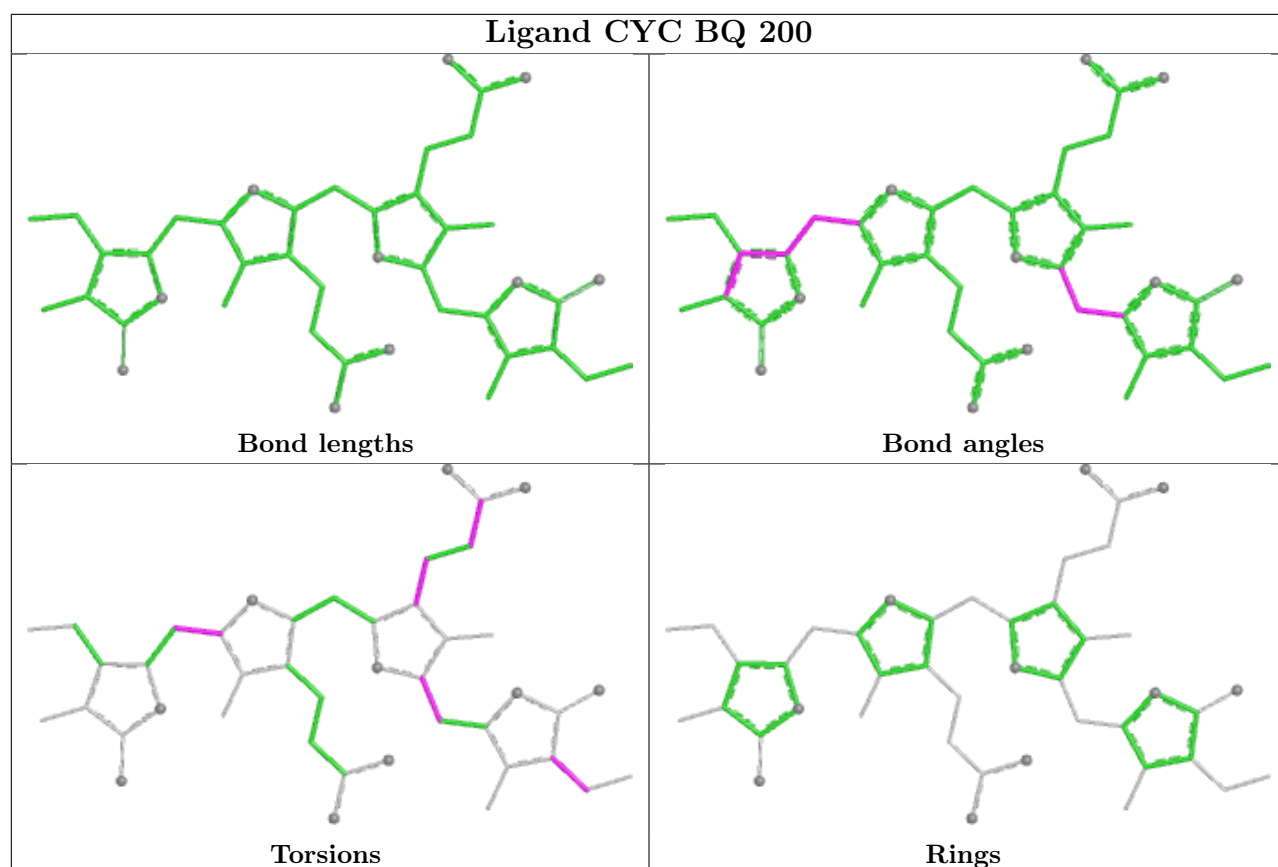


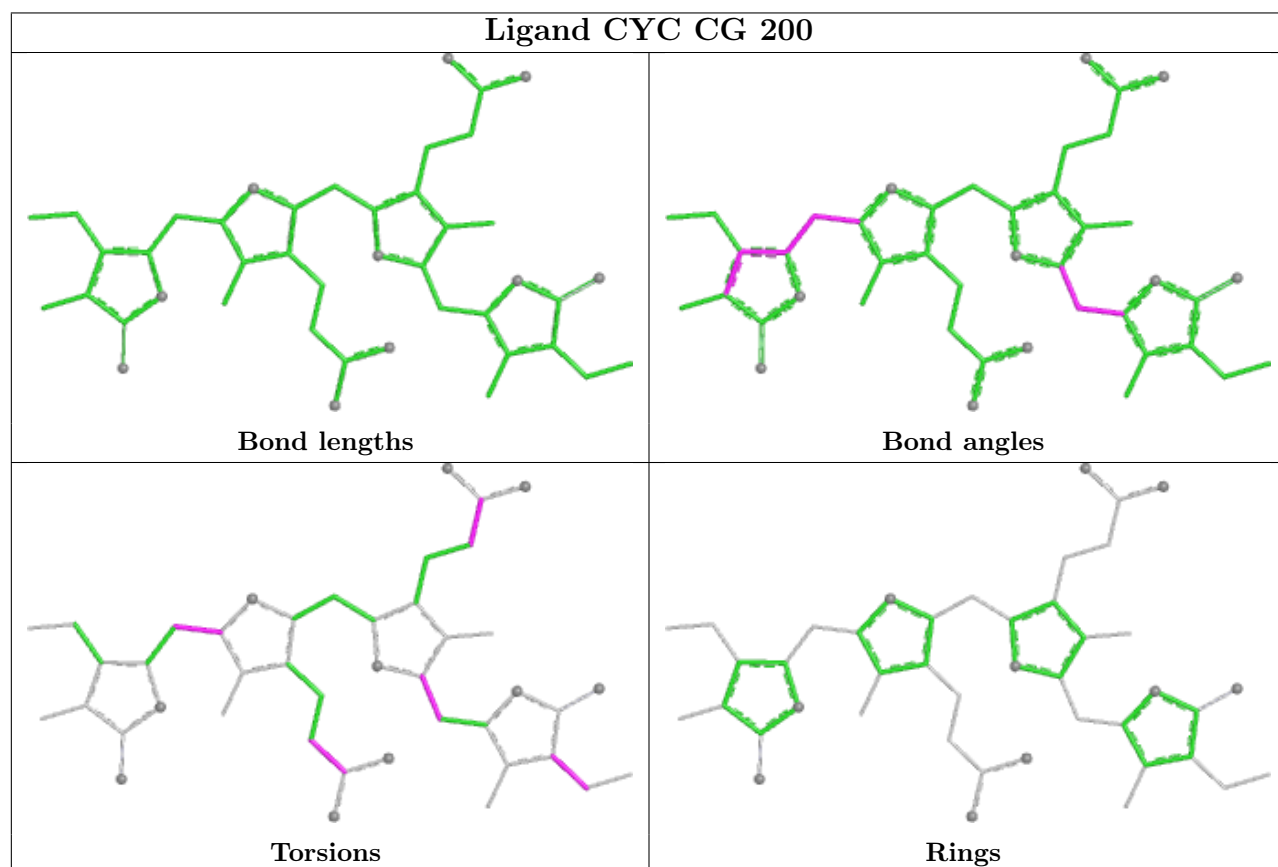
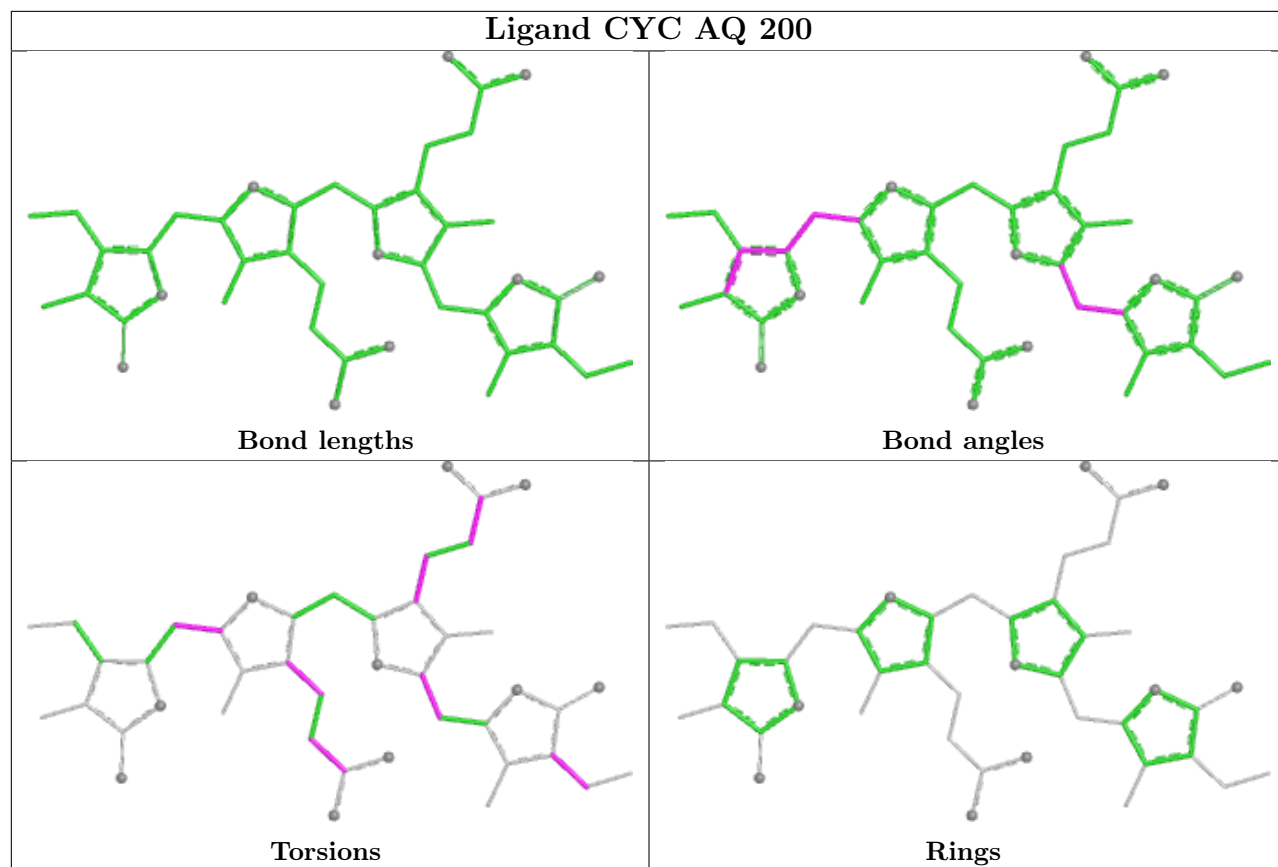
Ligand CYC AV 200



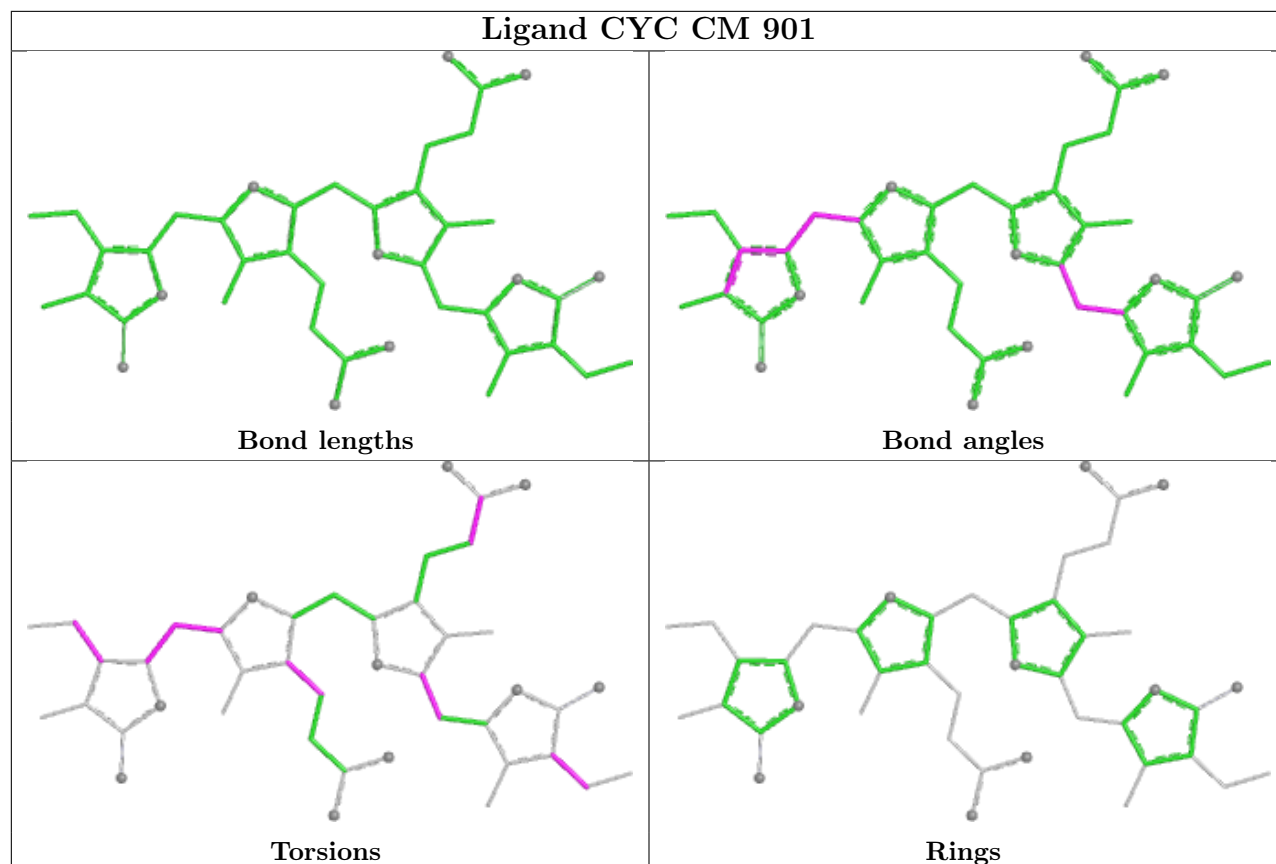
Ligand CYC BR 200



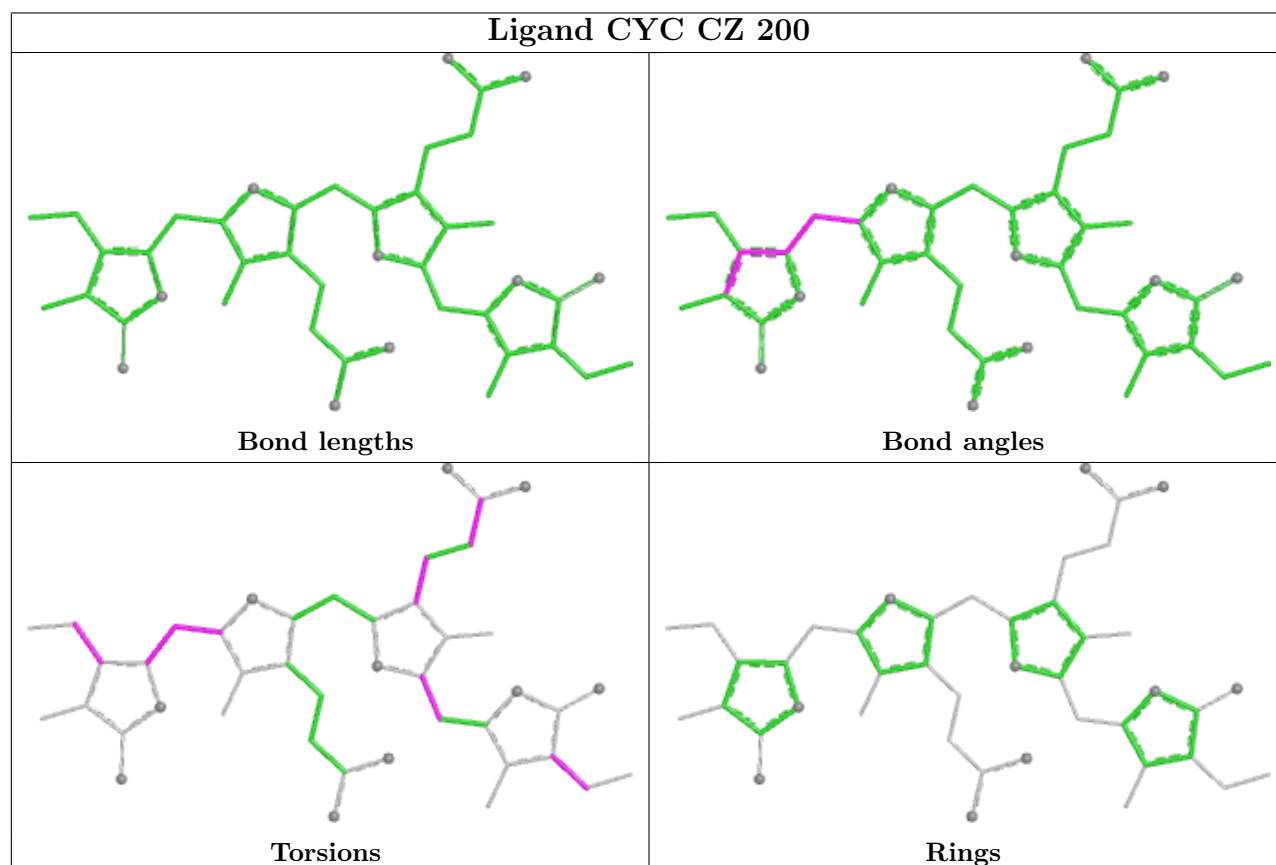


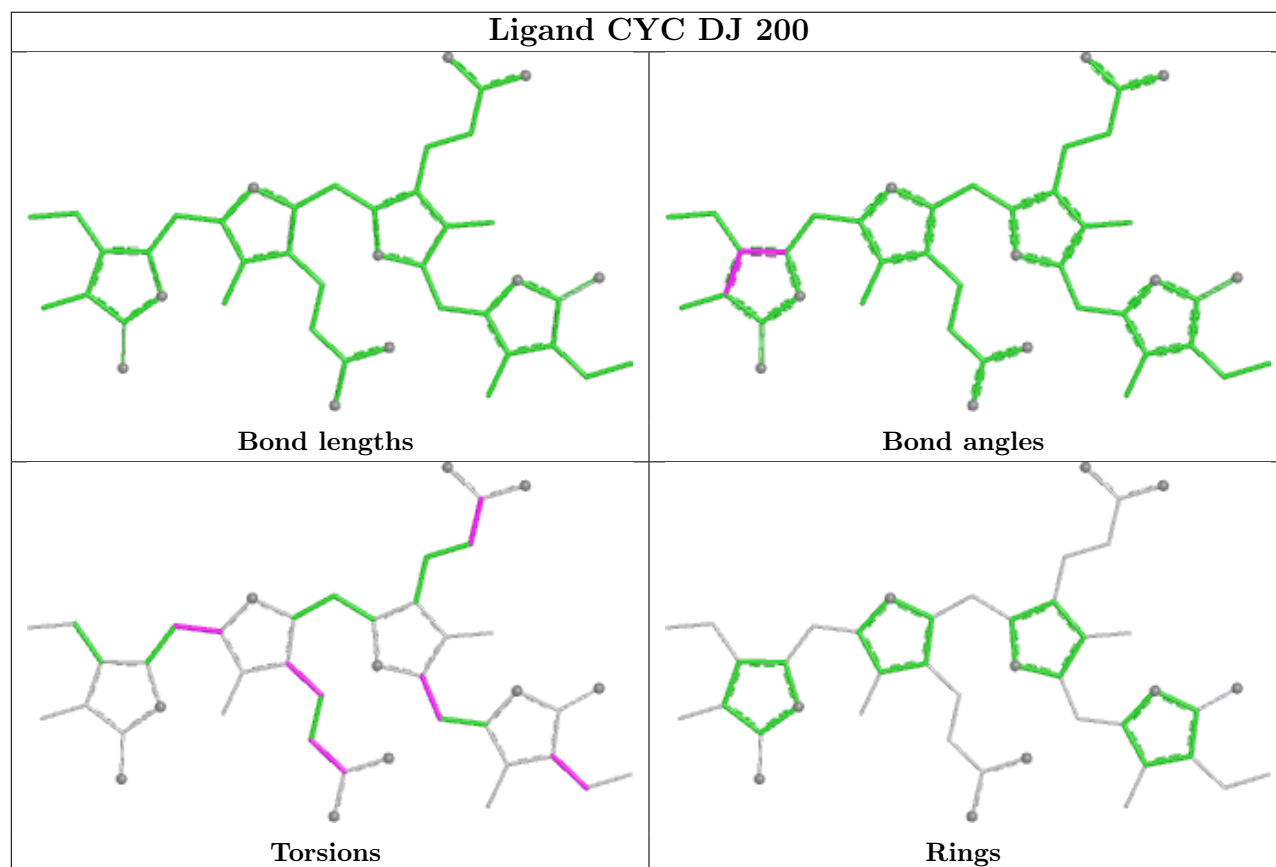
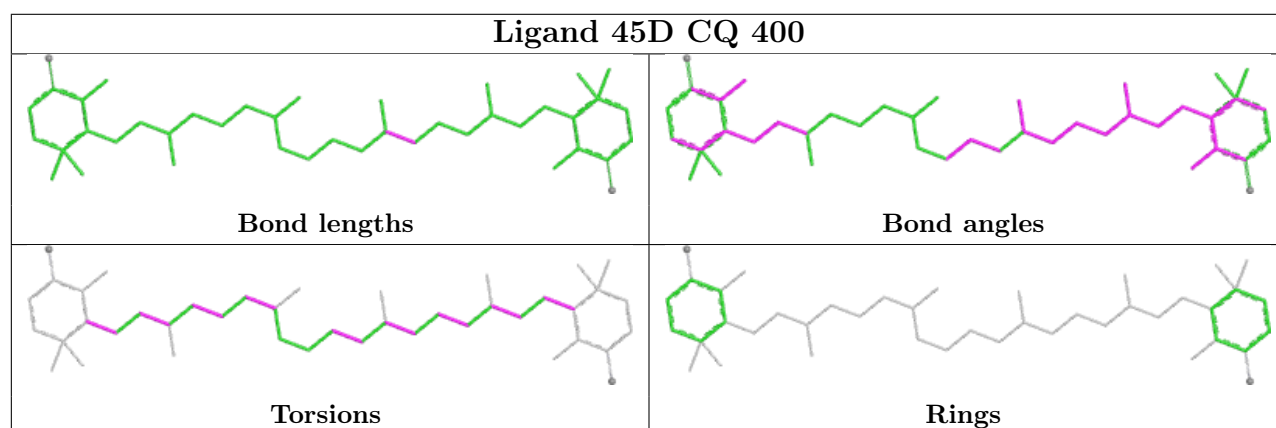
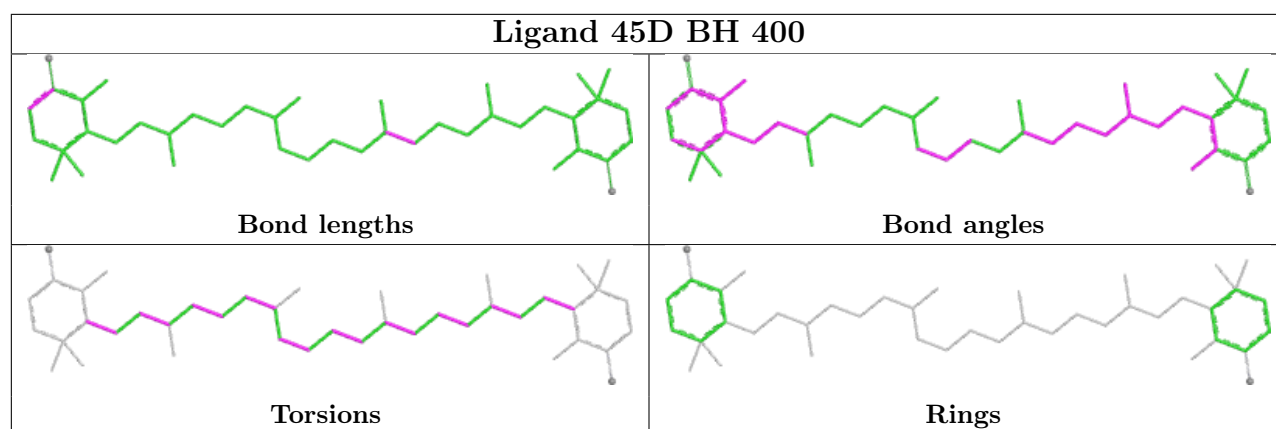


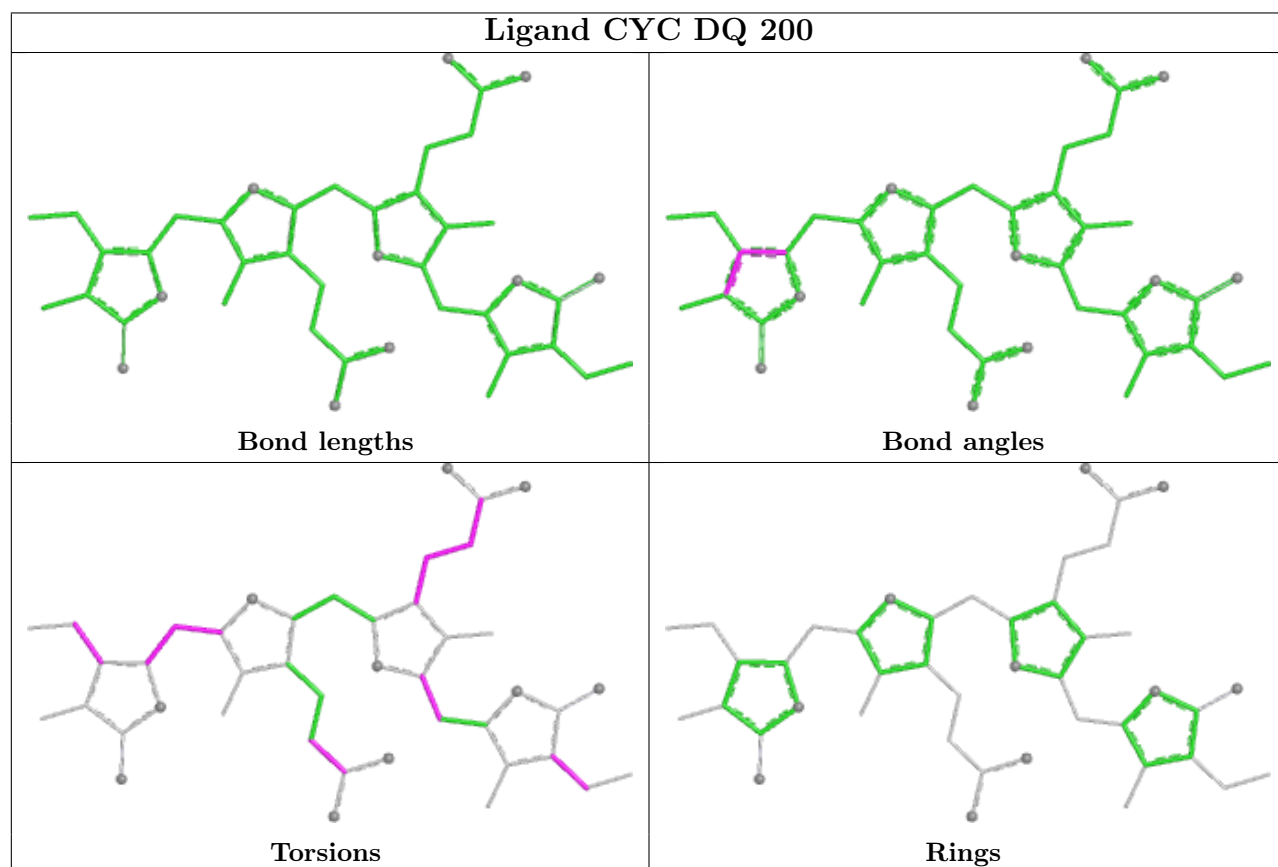
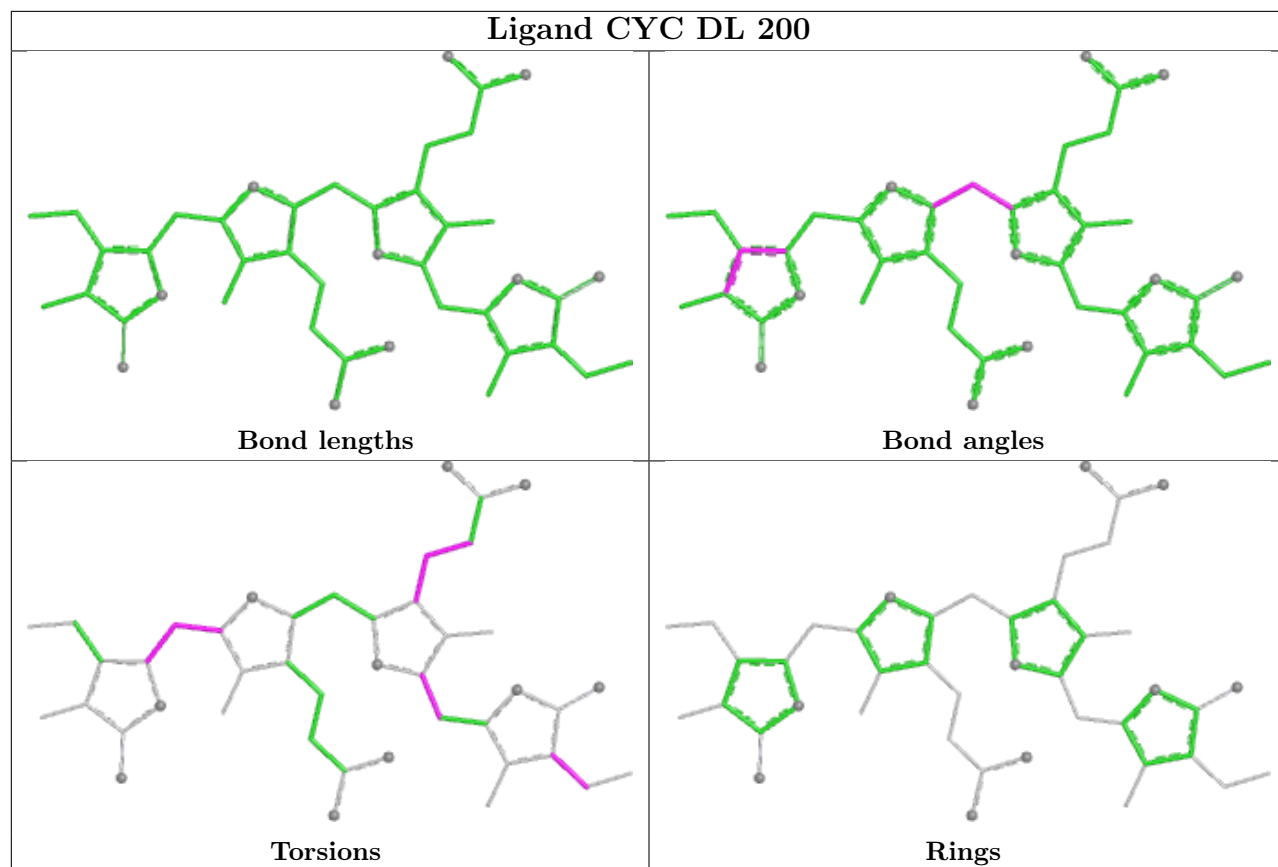
Ligand CYC CM 901



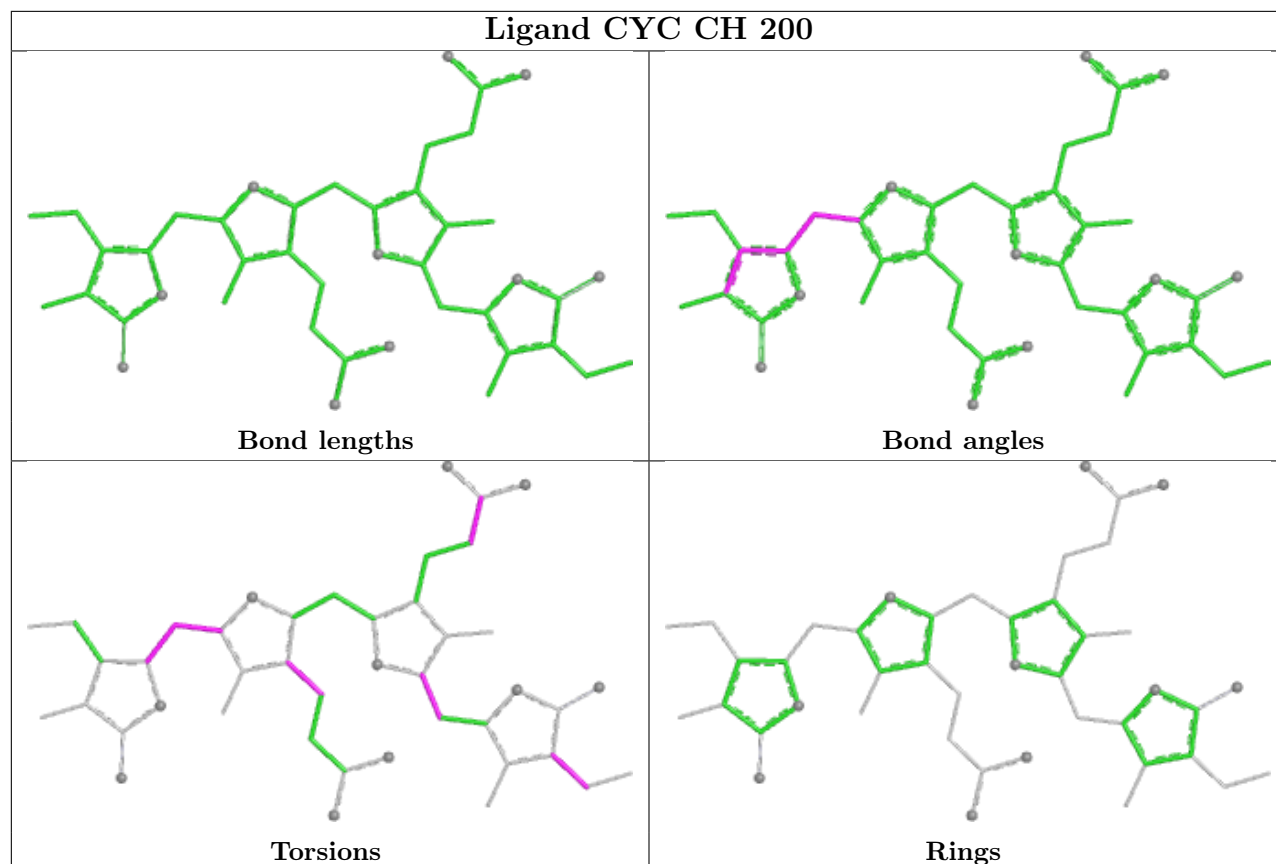
Ligand CYC CZ 200



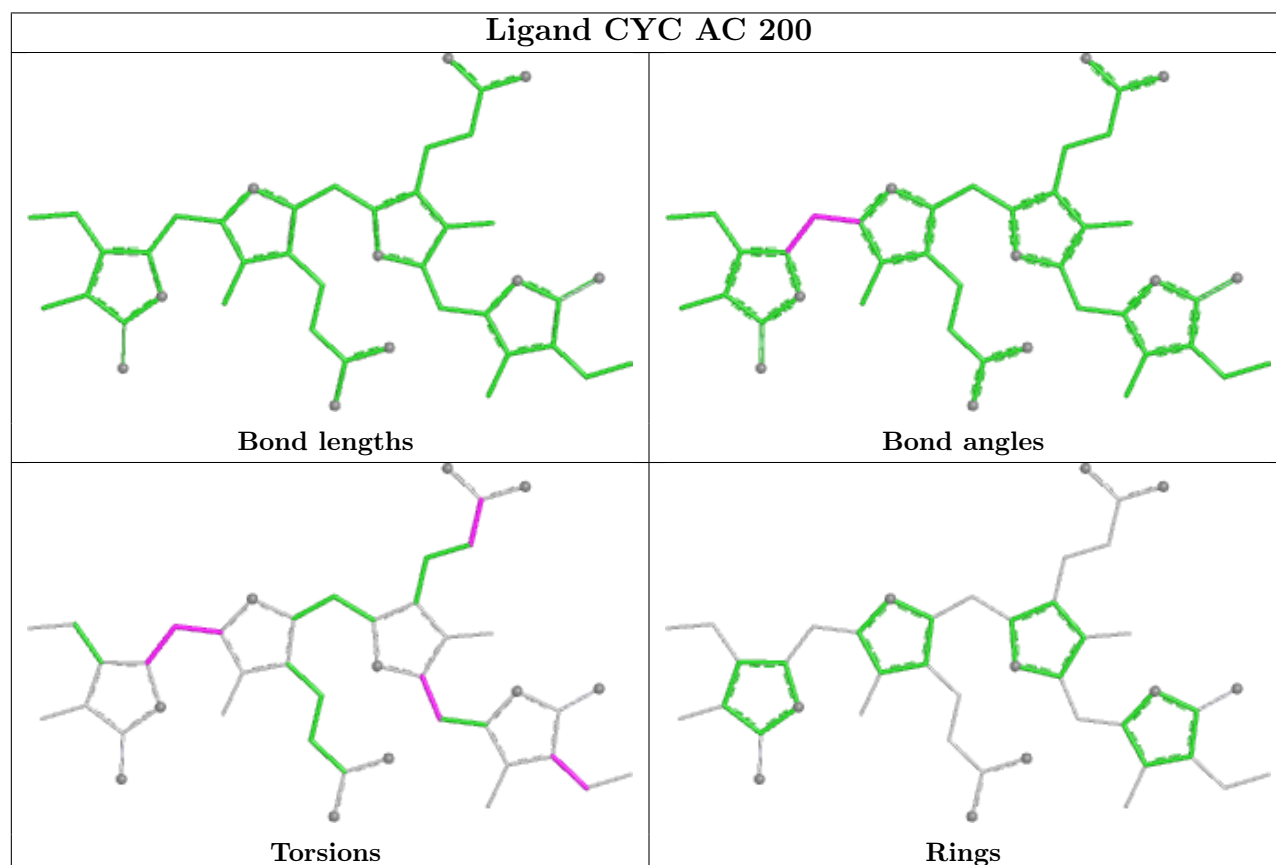




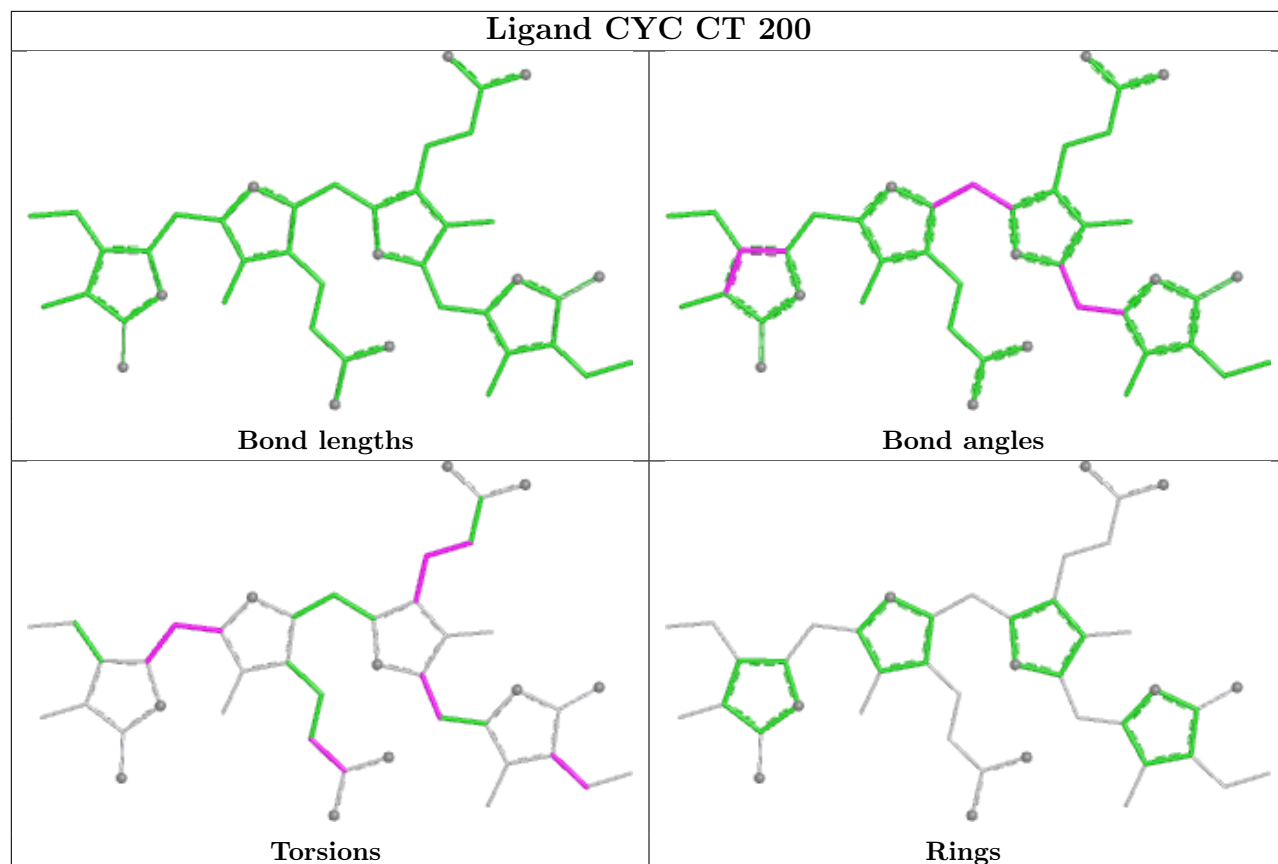
Ligand CYC CH 200



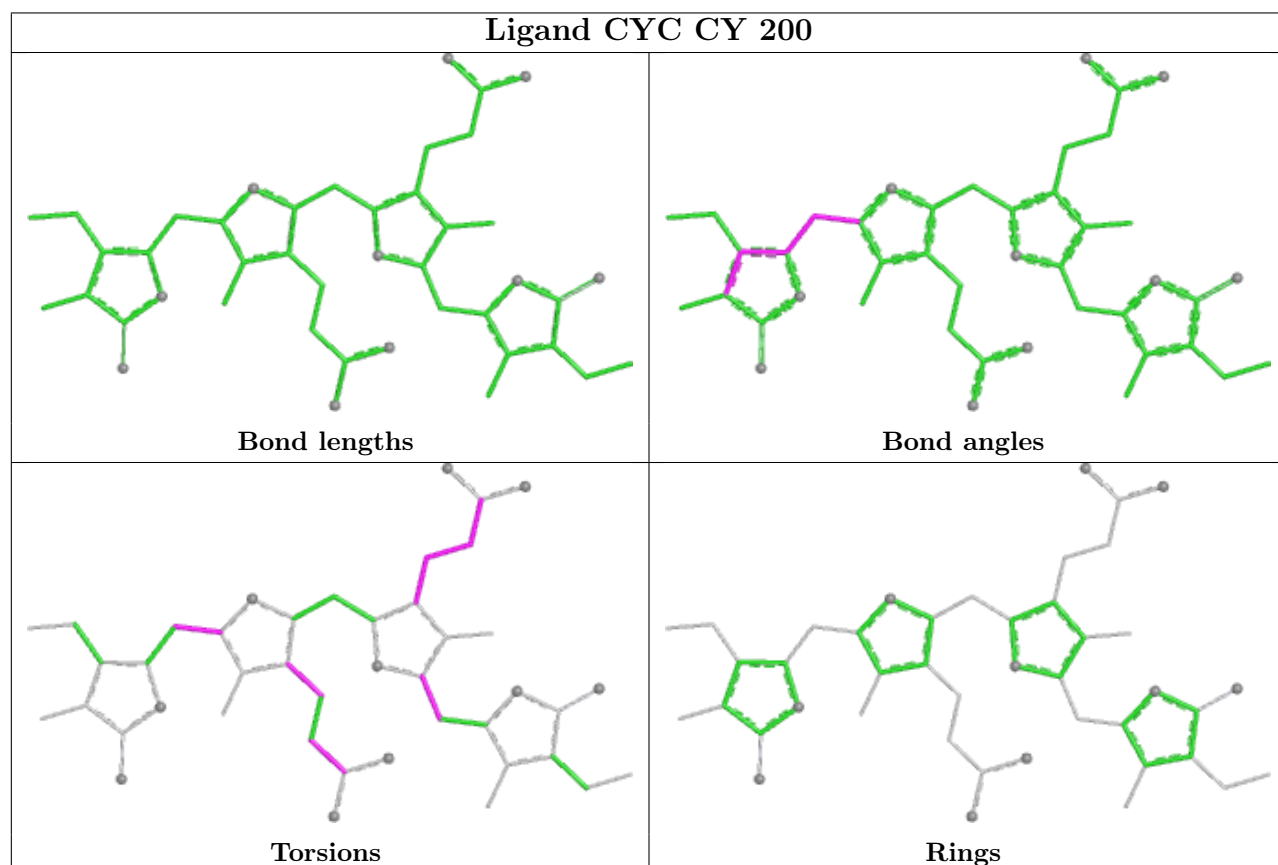
Ligand CYC AC 200



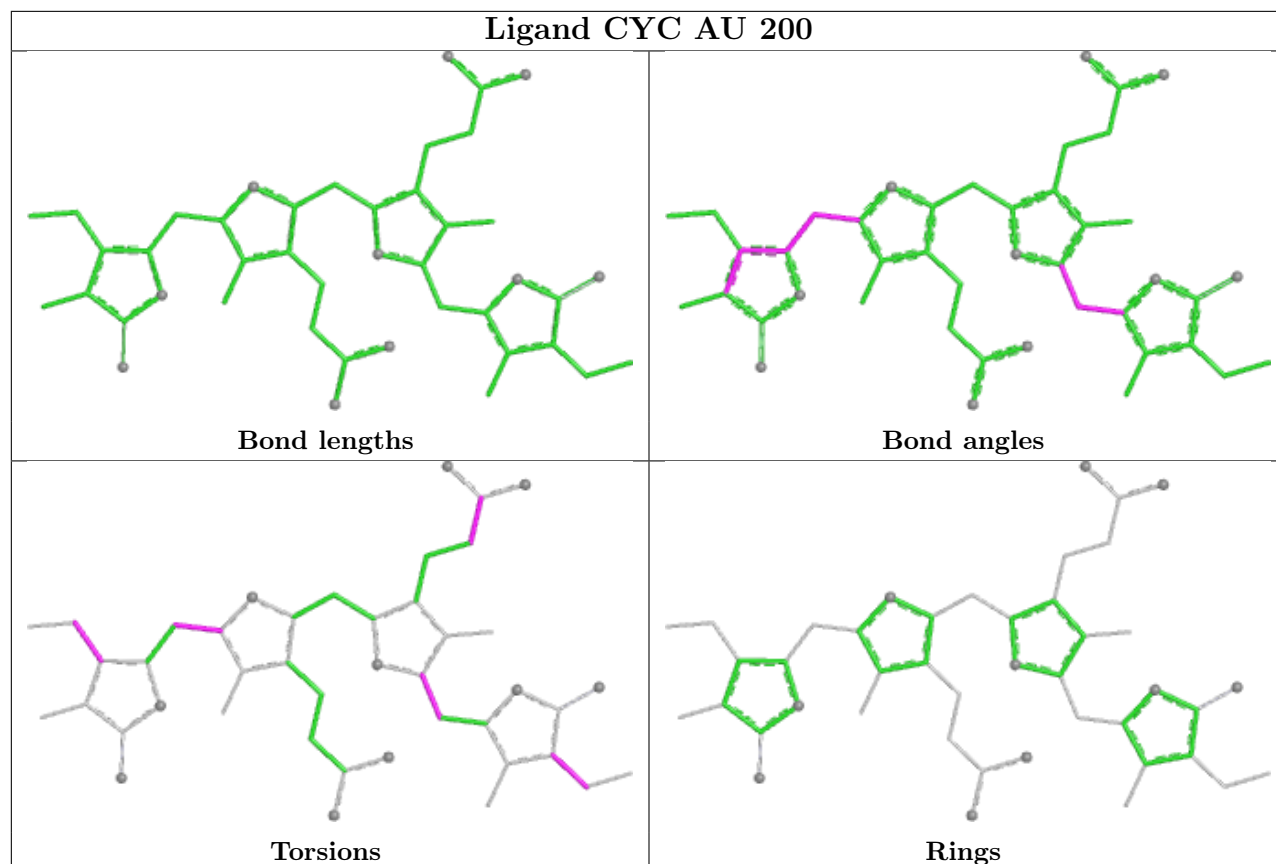
Ligand CYC CT 200



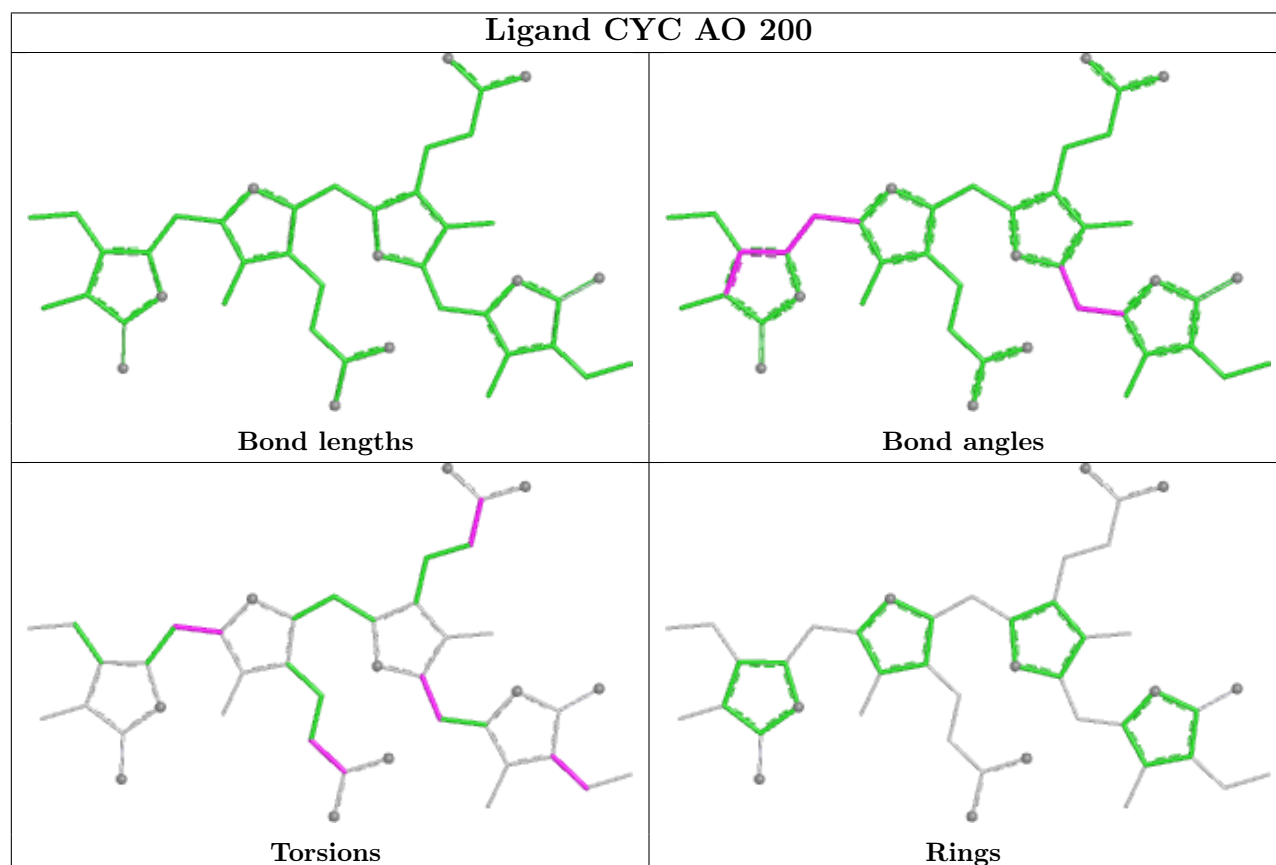
Ligand CYC CY 200



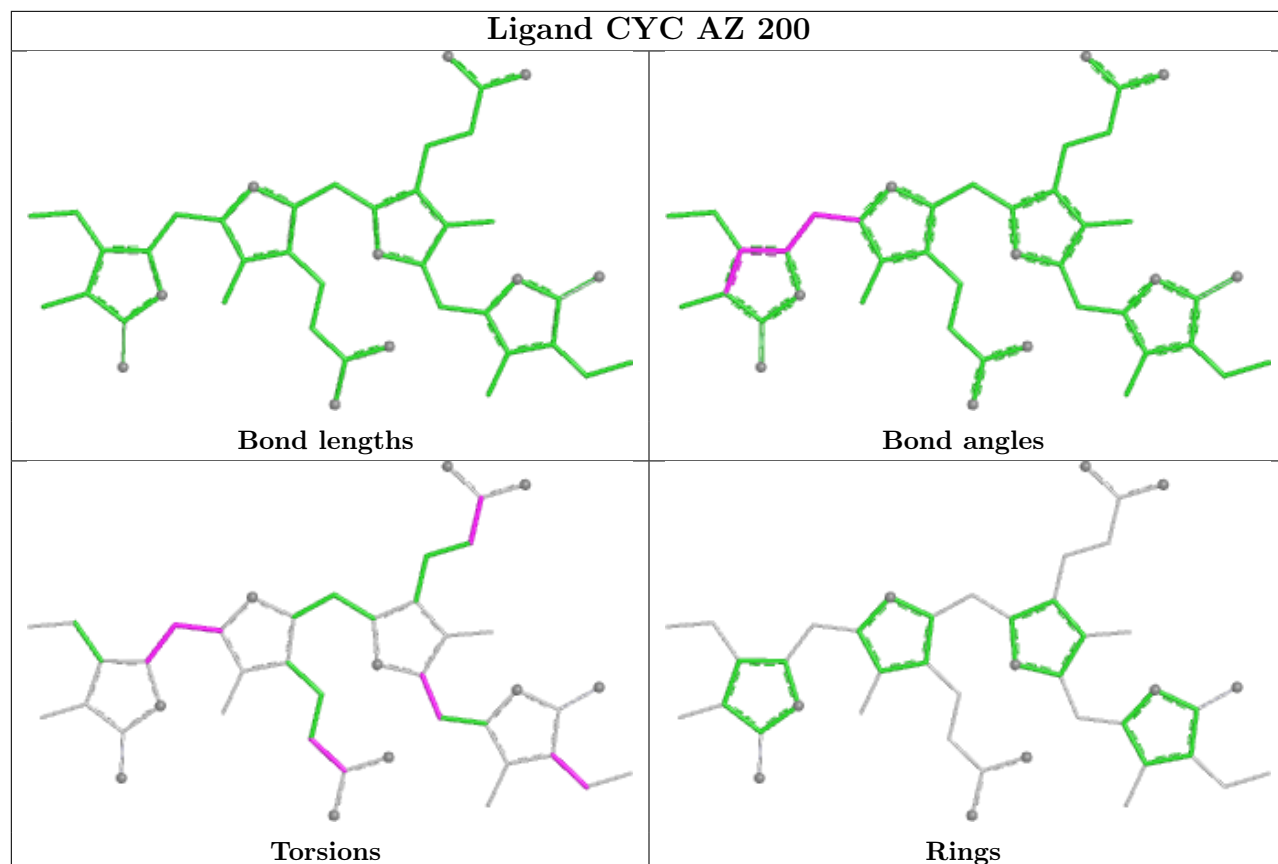
Ligand CYC AU 200



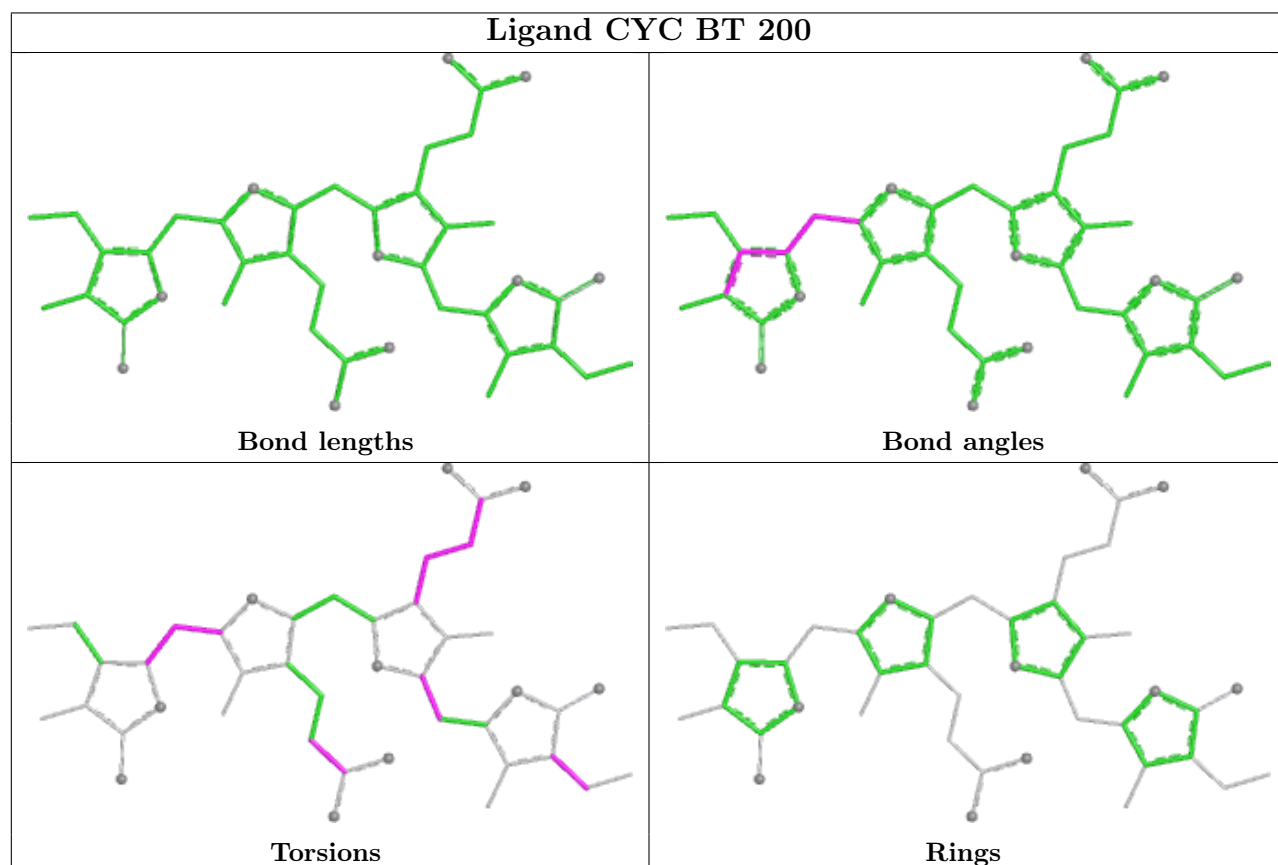
Ligand CYC AO 200



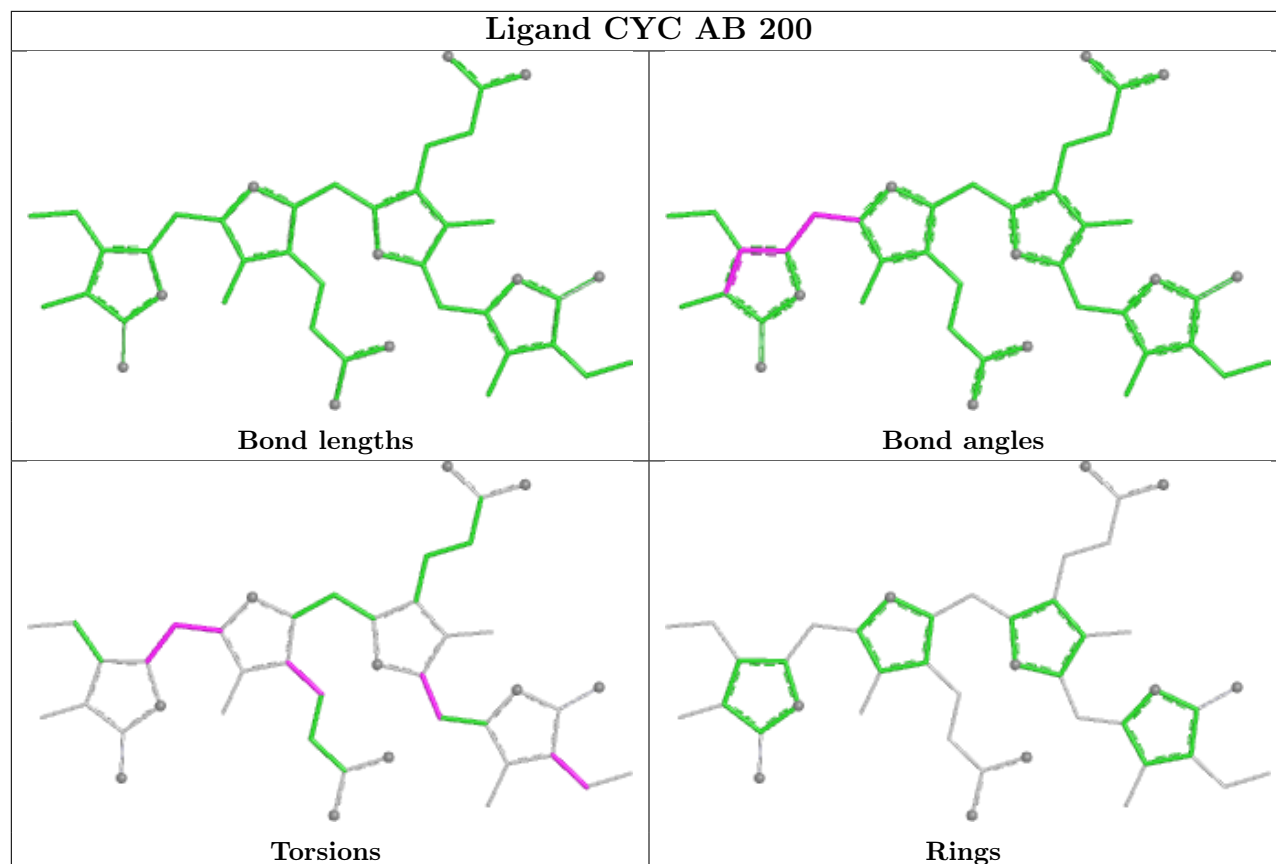
Ligand CYC AZ 200



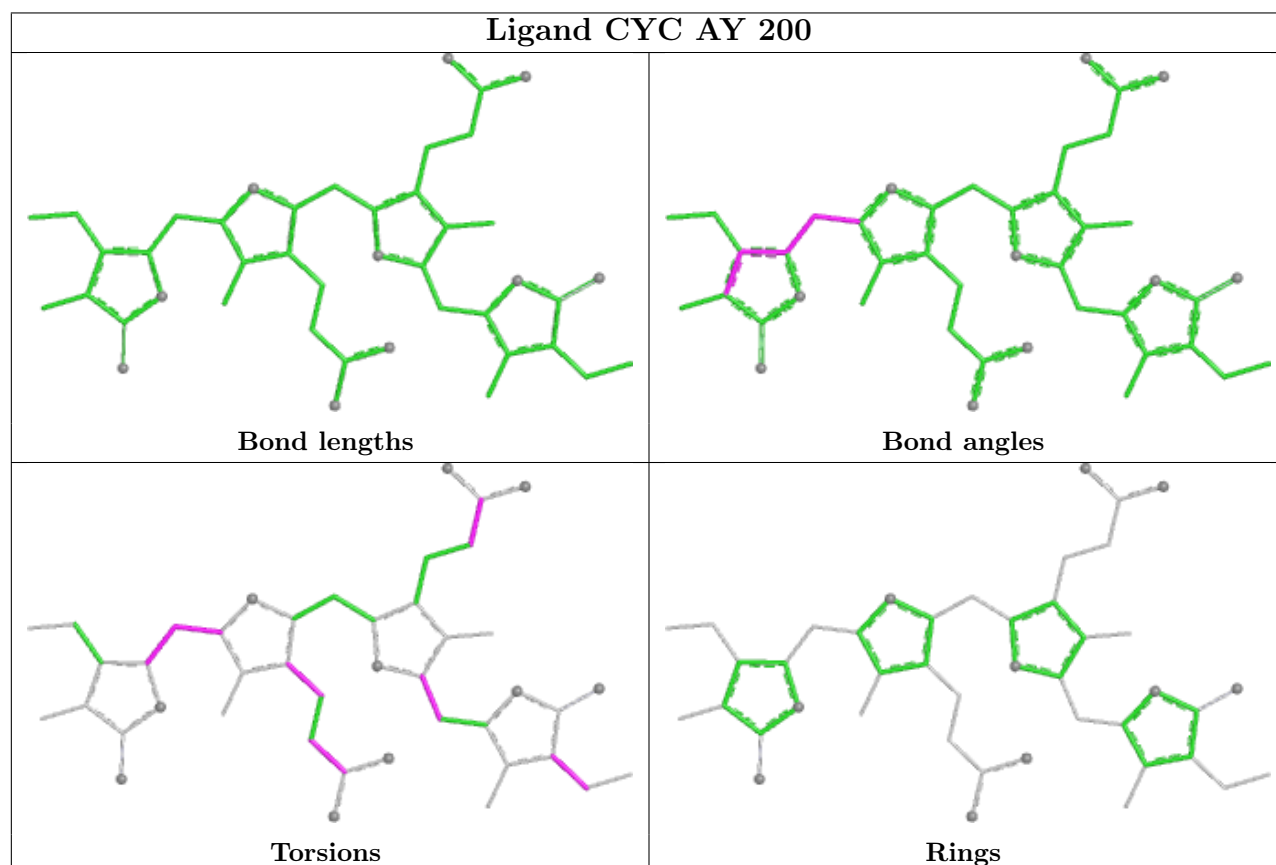
Ligand CYC BT 200



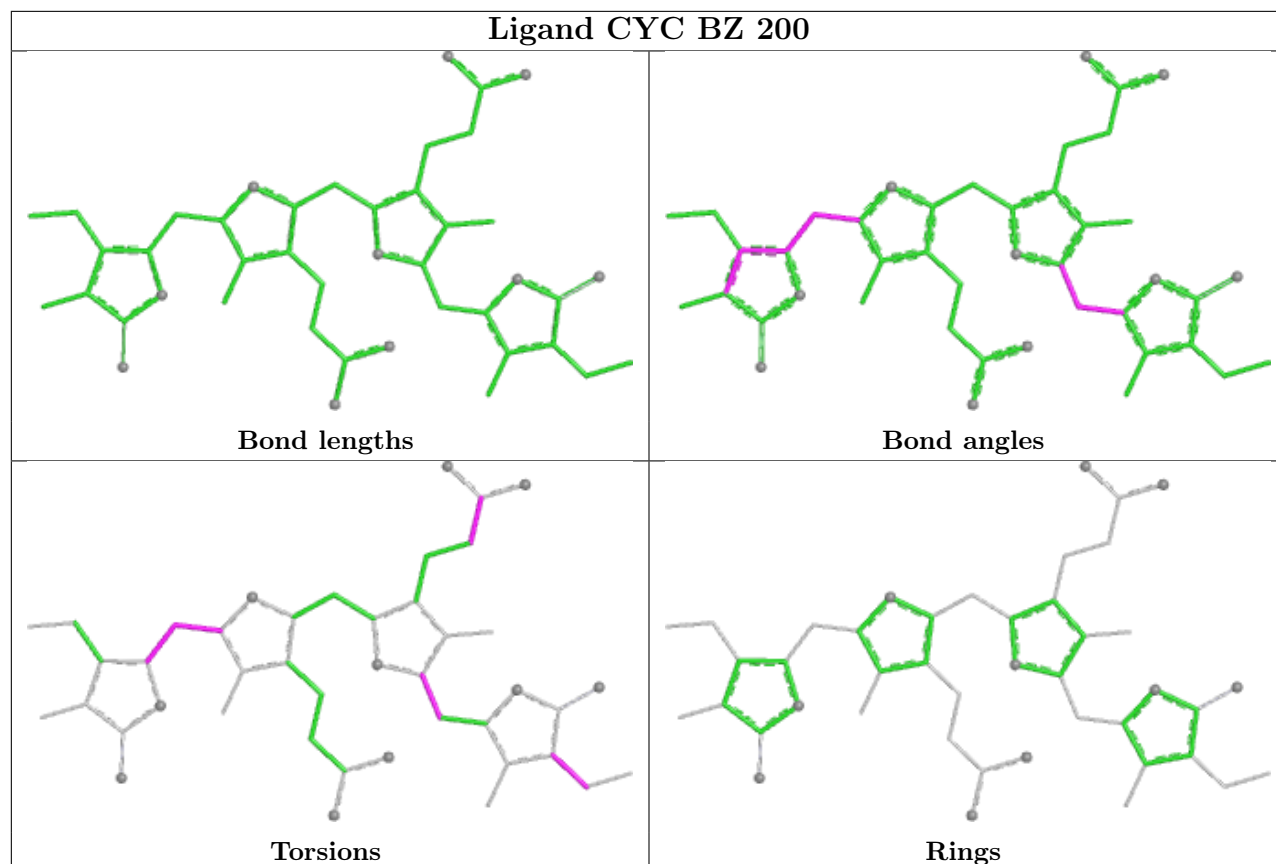
Ligand CYC AB 200



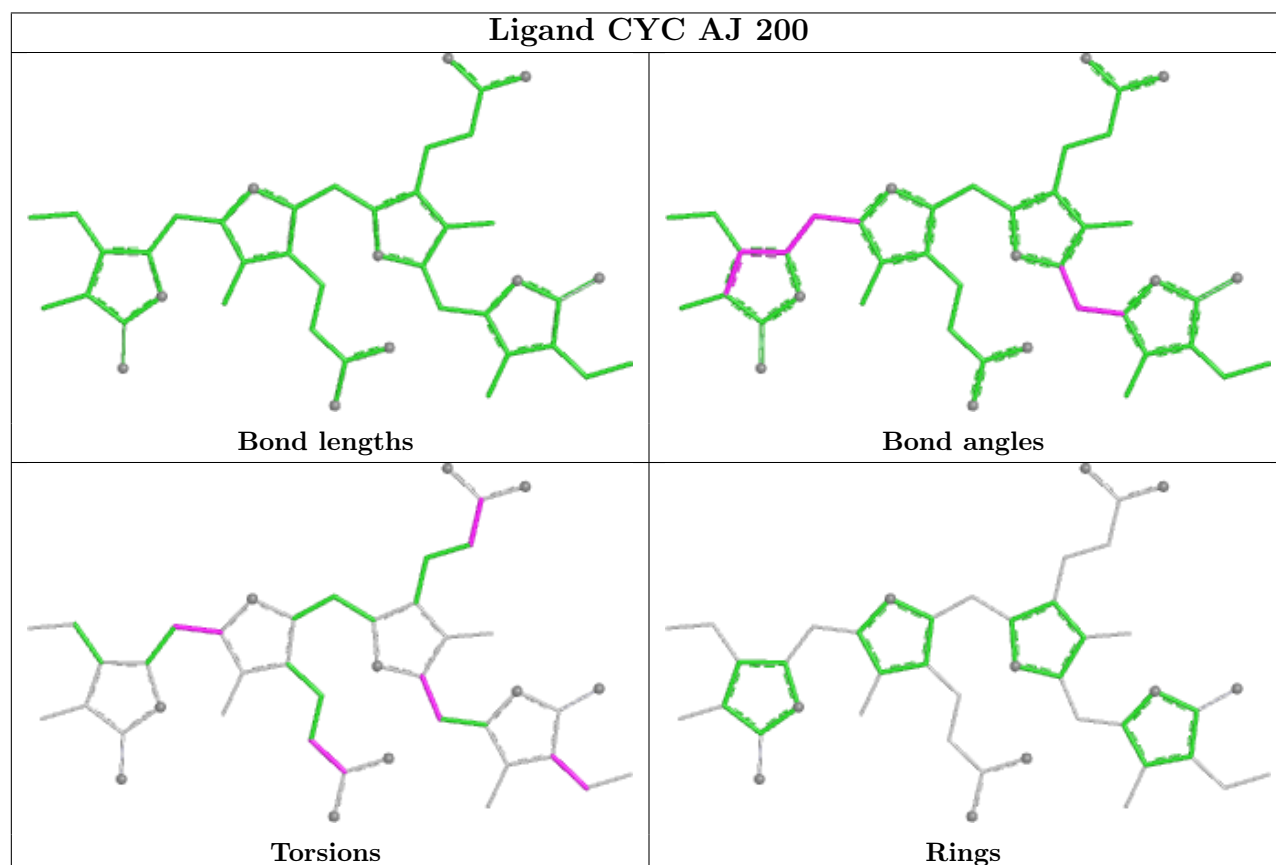
Ligand CYC AY 200



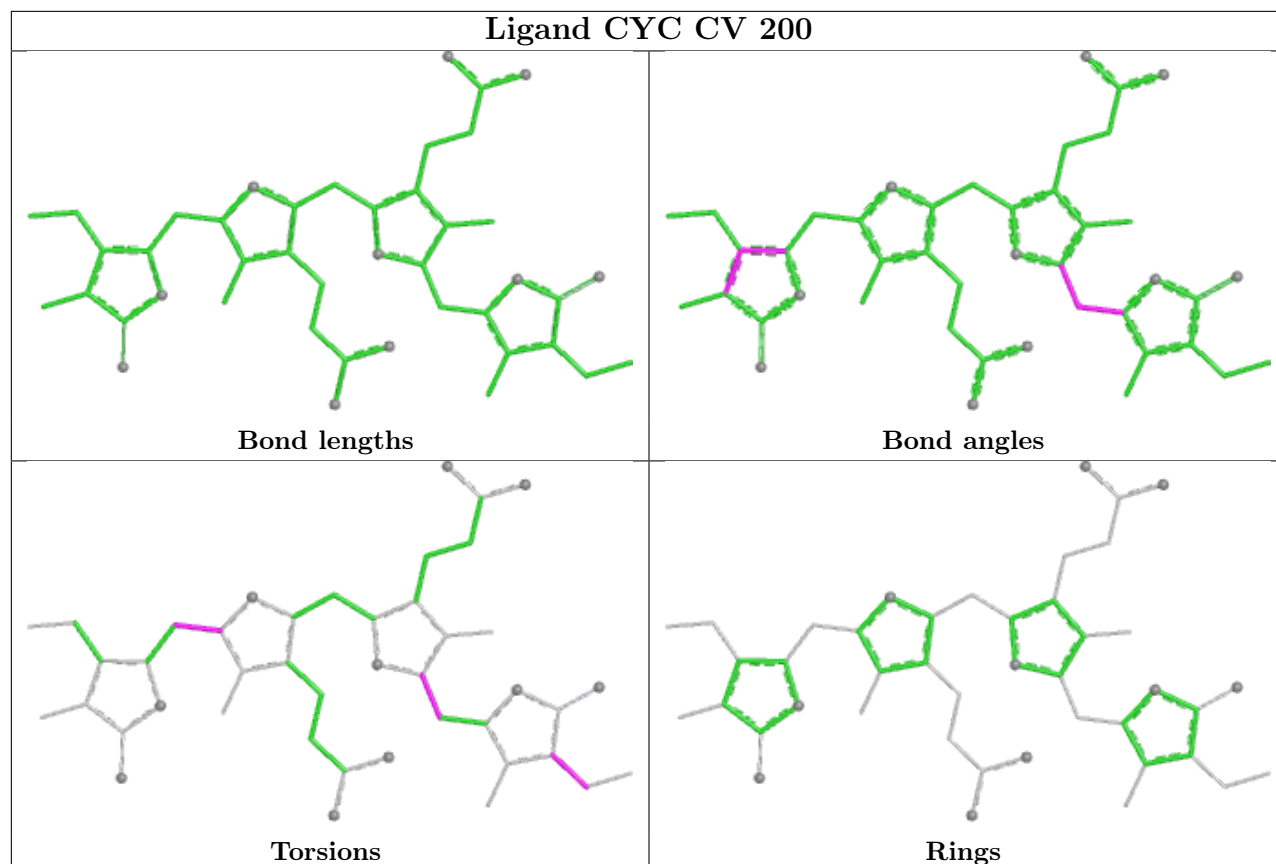
Ligand CYC BZ 200



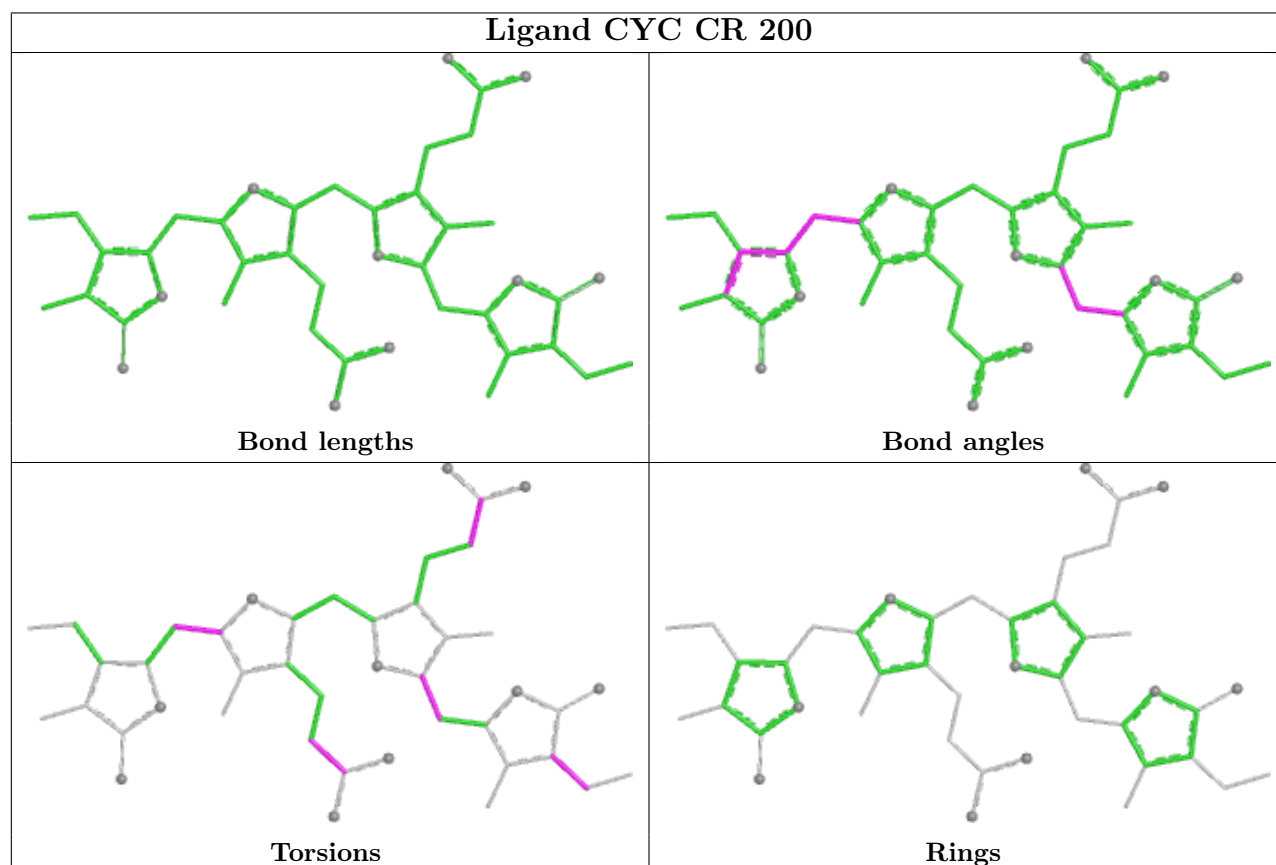
Ligand CYC AJ 200

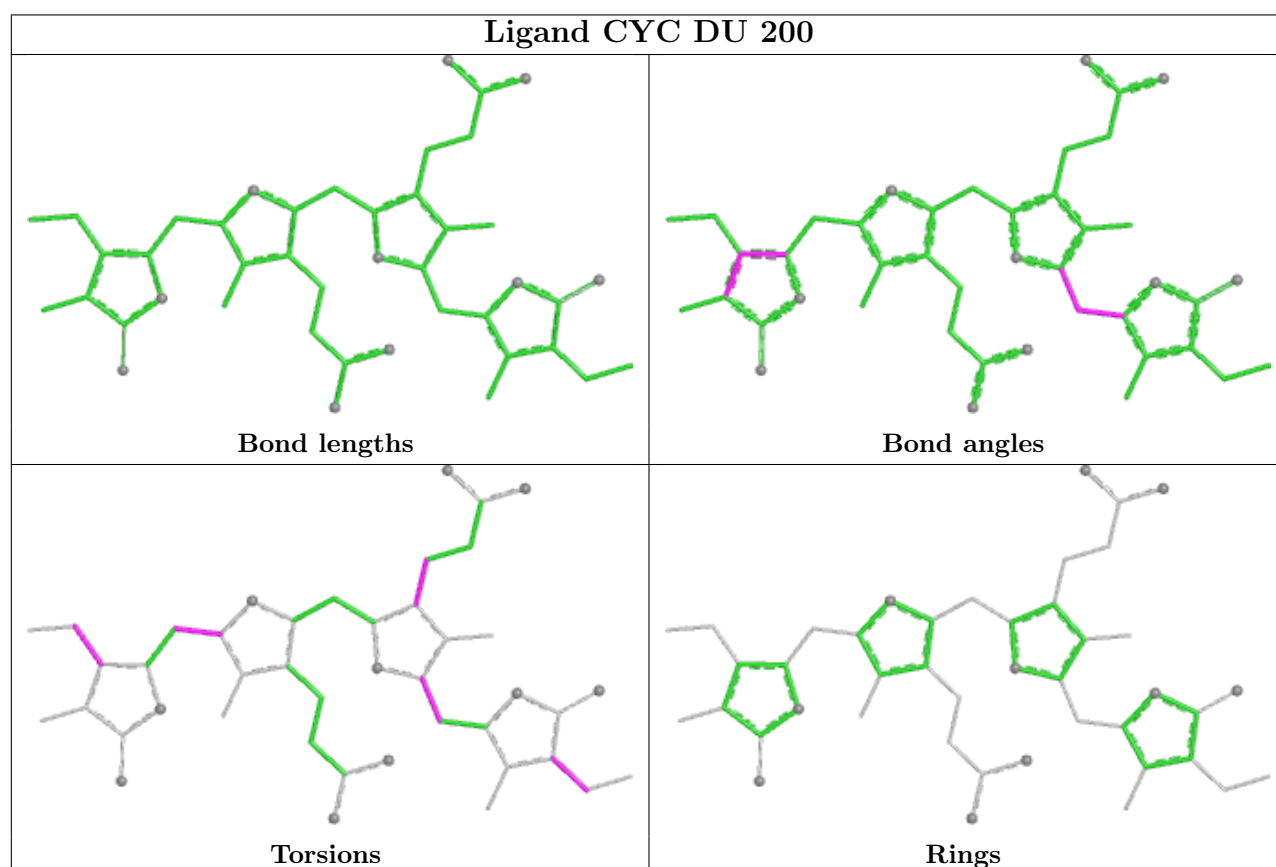
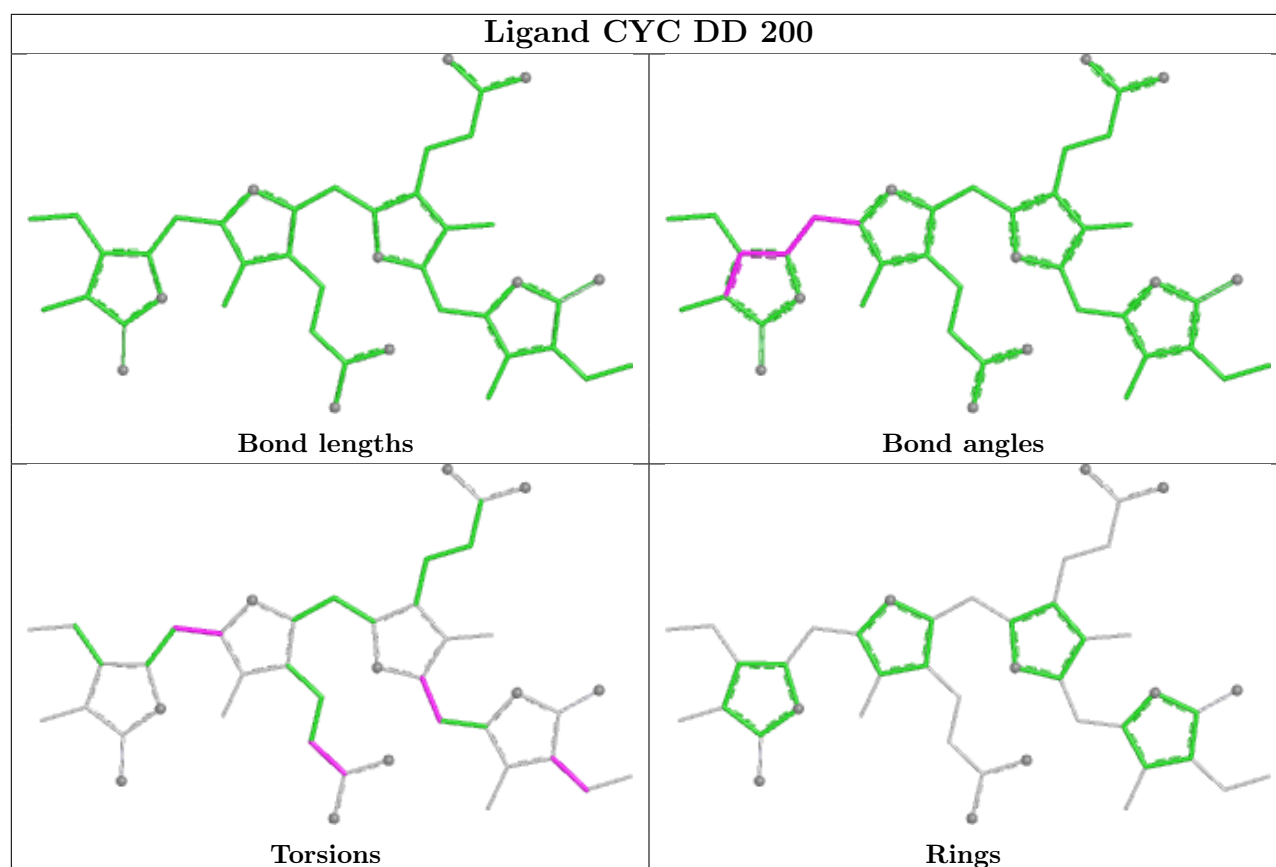


Ligand CYC CV 200

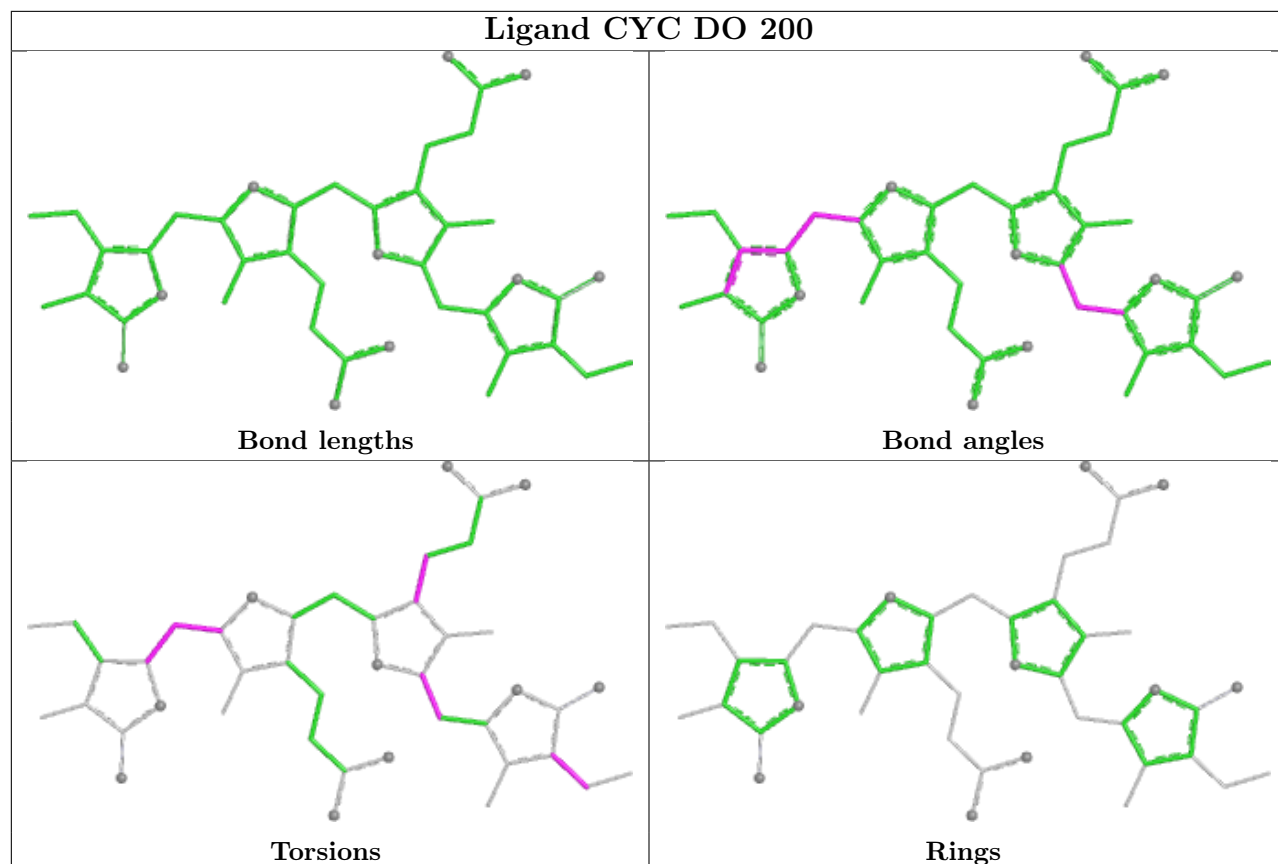


Ligand CYC CR 200

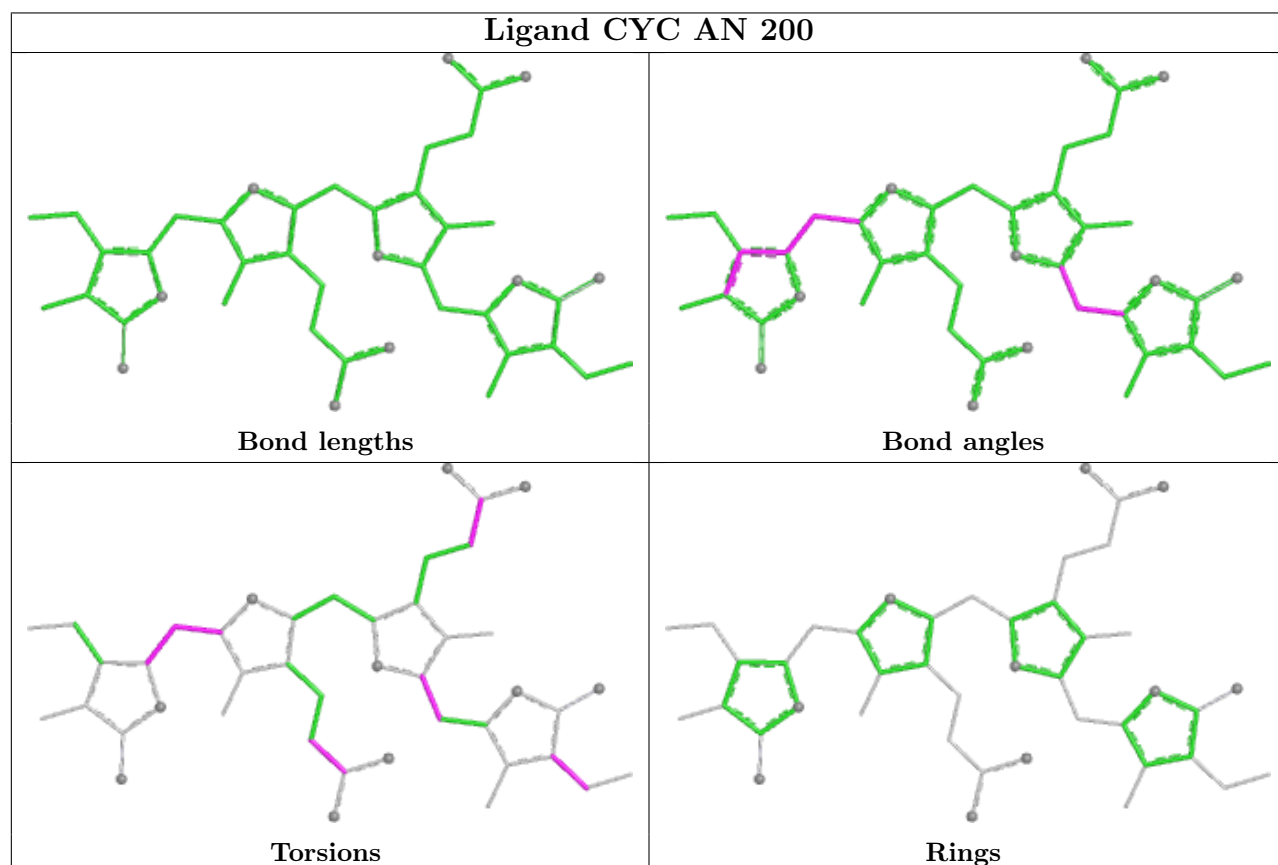




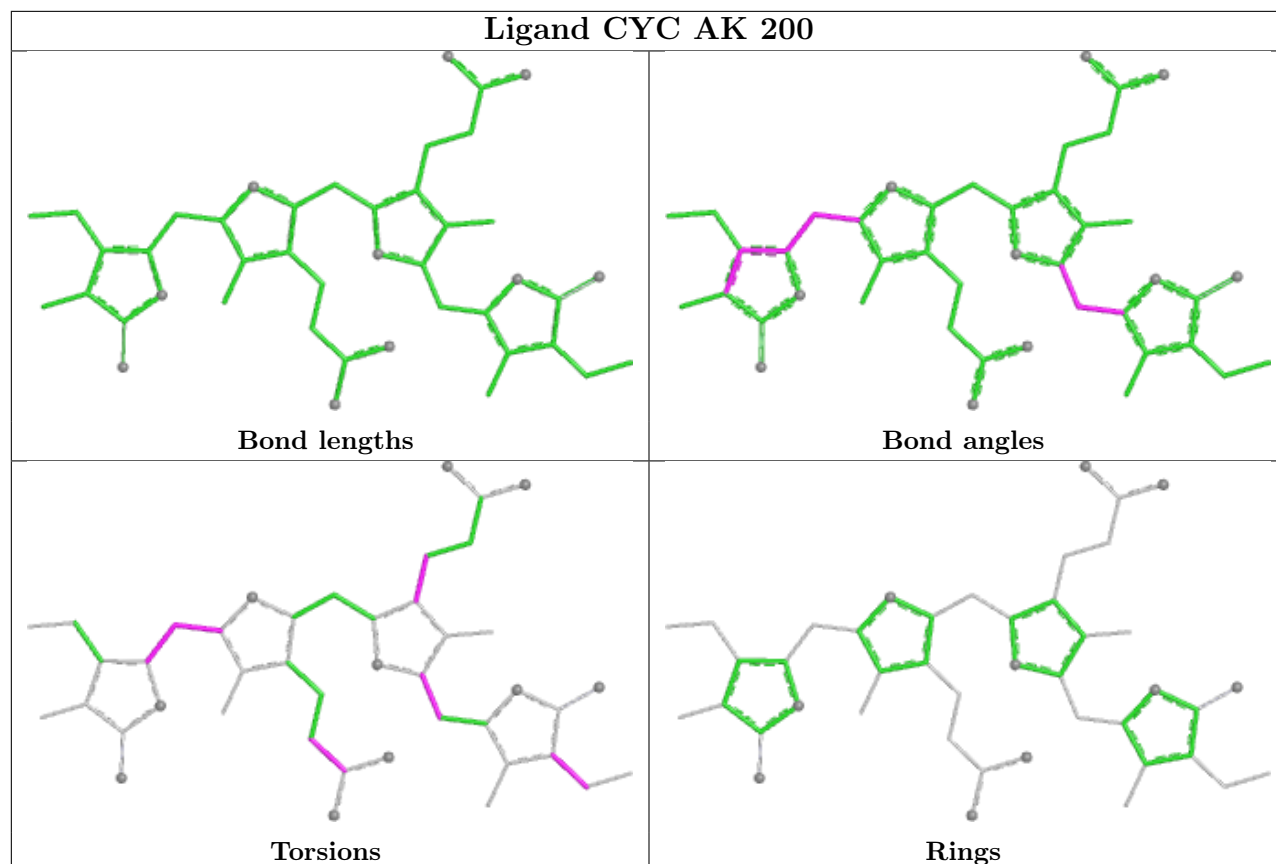
Ligand CYC DO 200



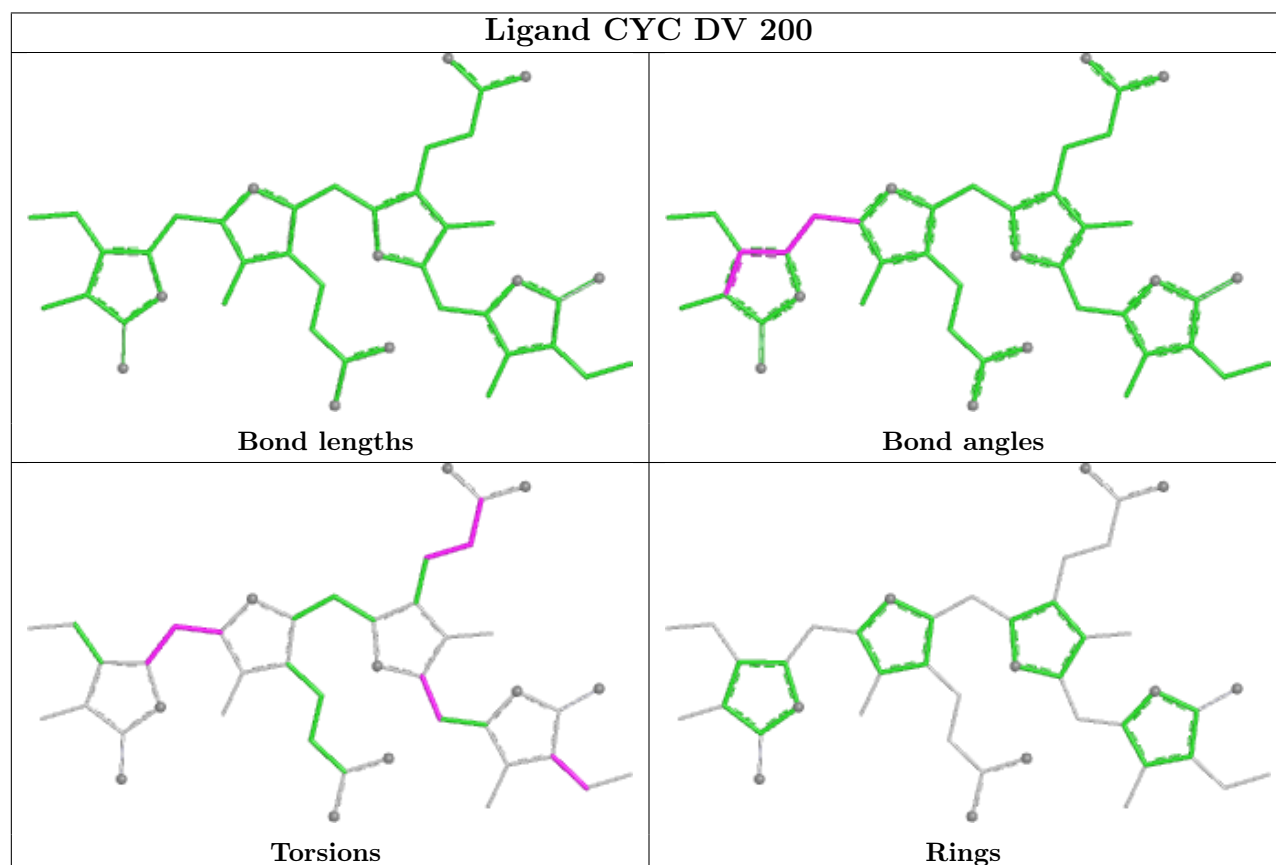
Ligand CYC AN 200

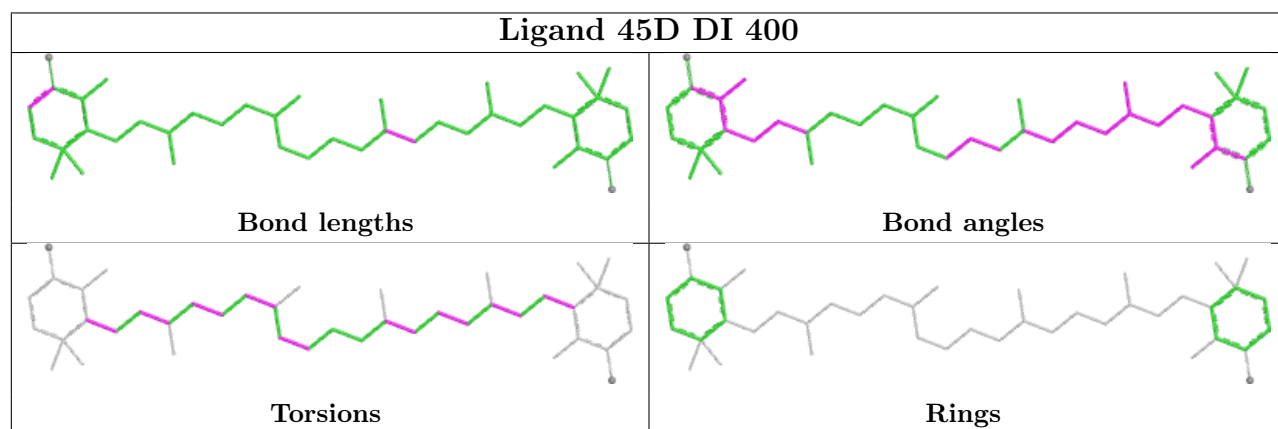
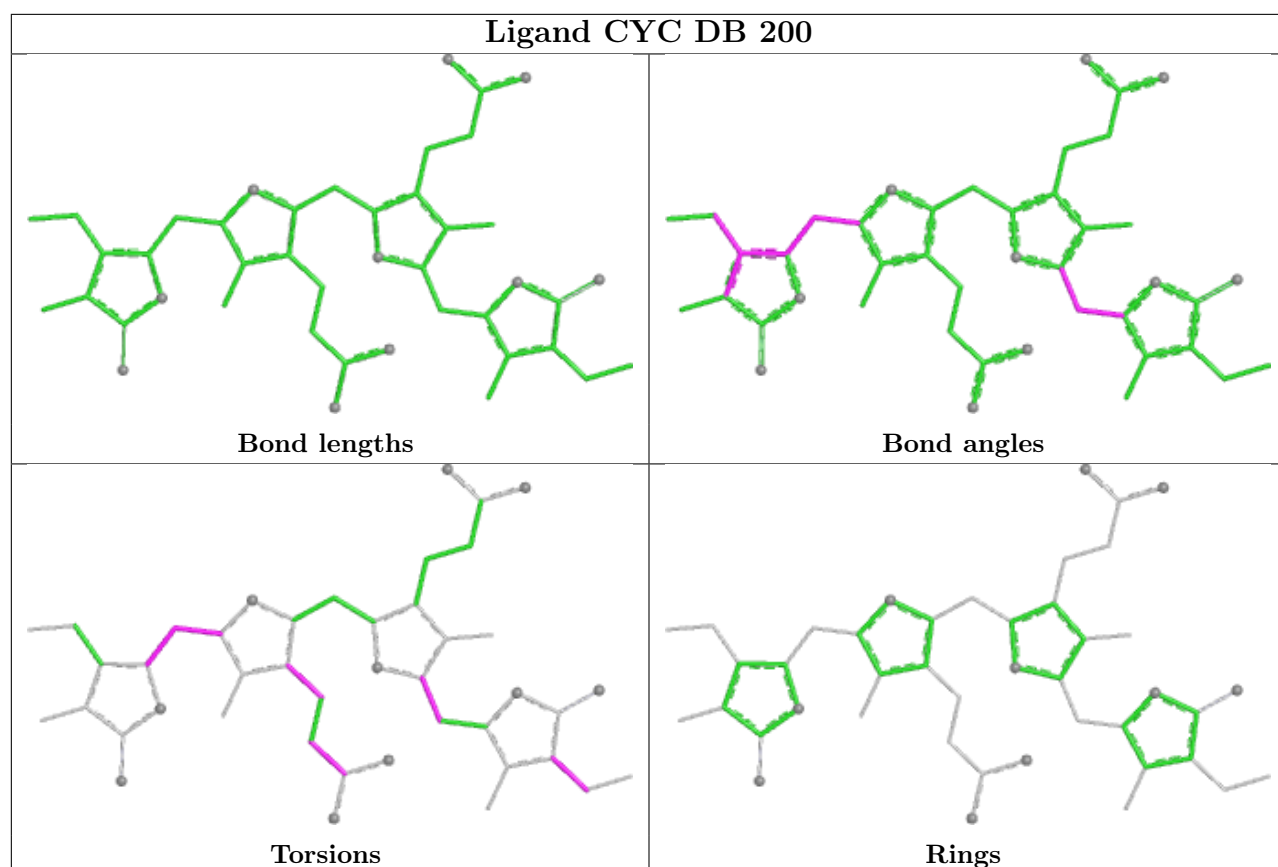


Ligand CYC AK 200

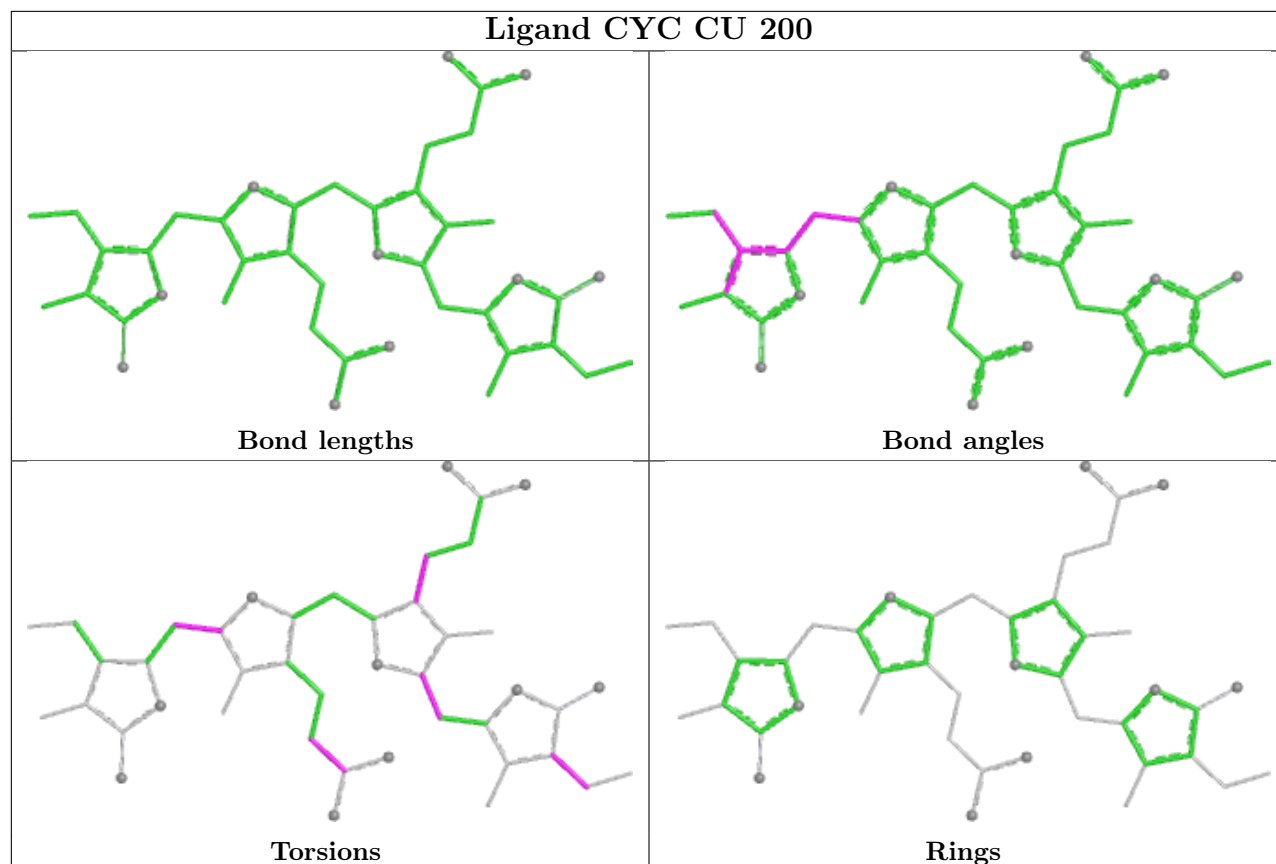


Ligand CYC DV 200

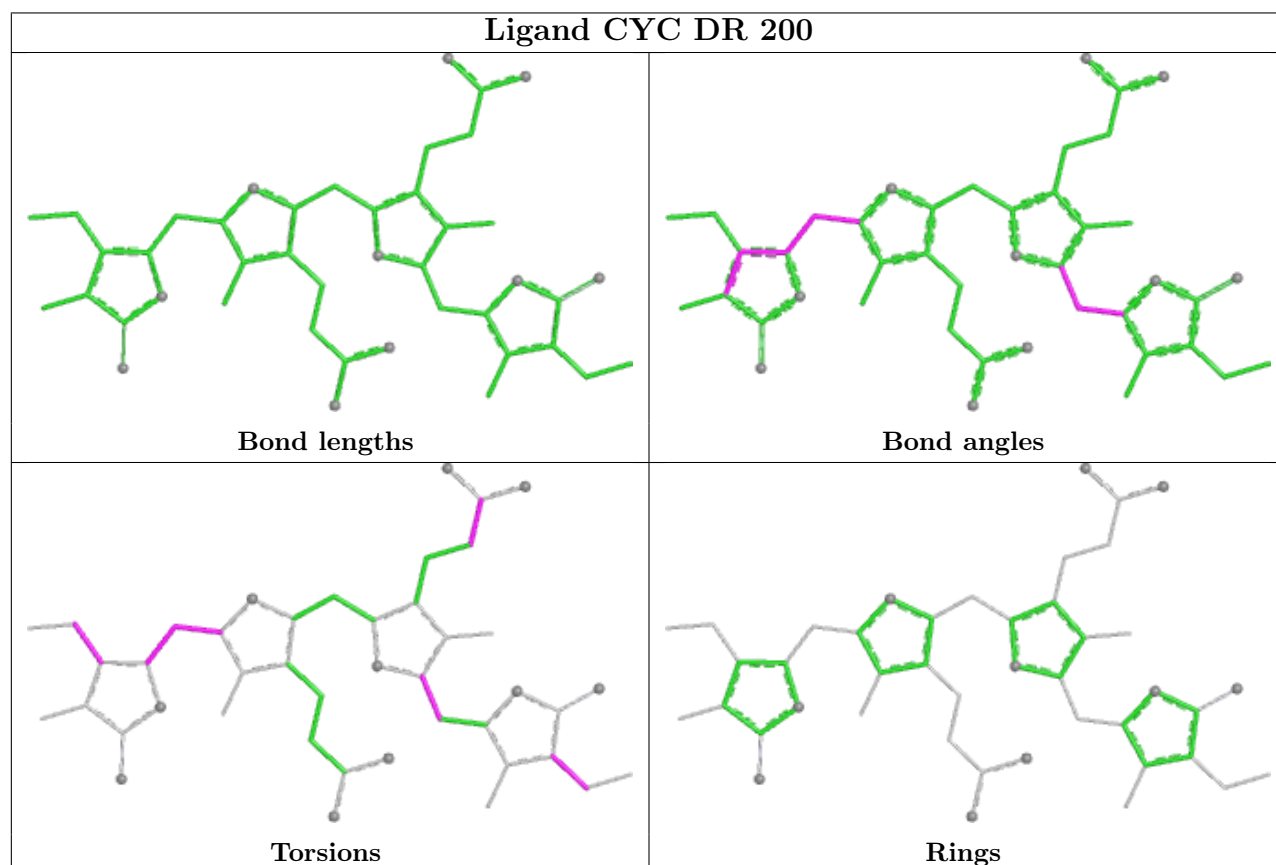




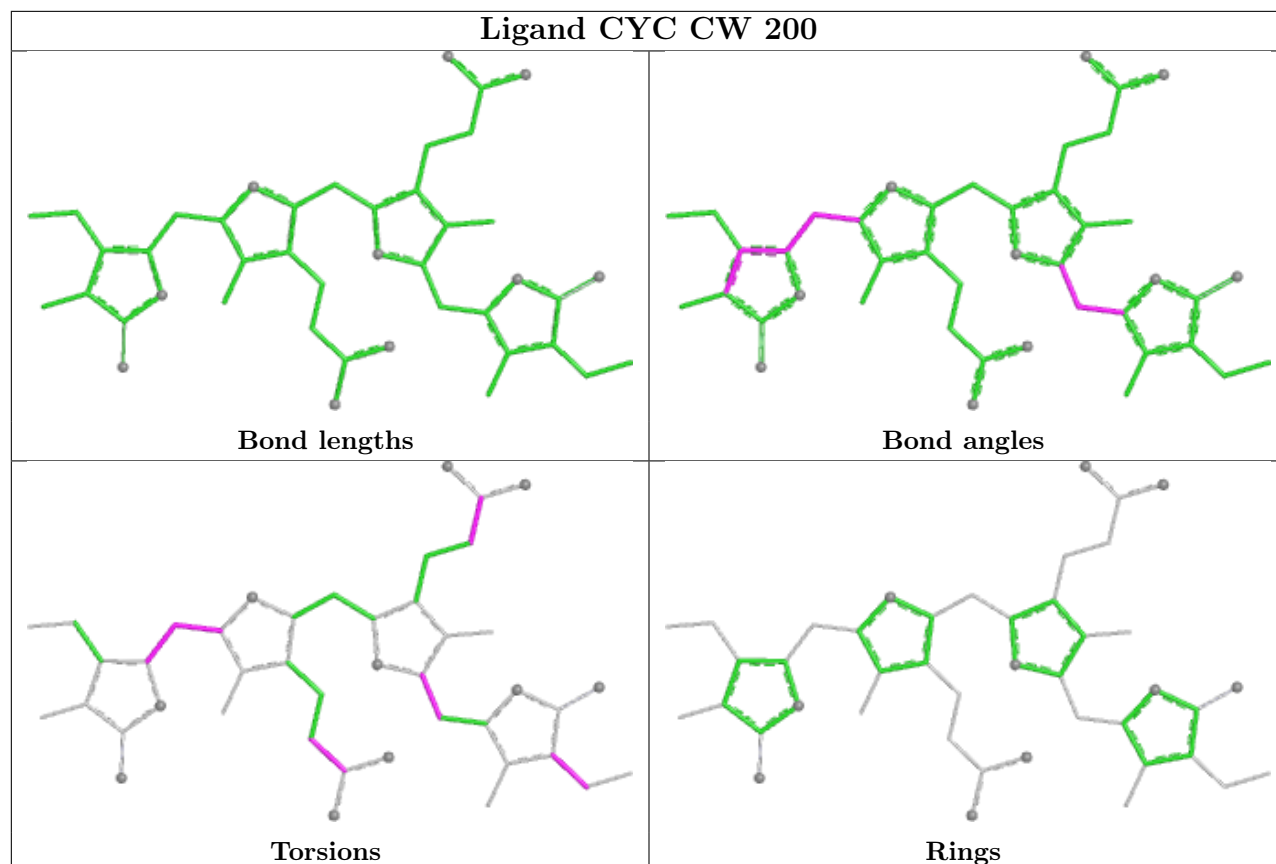
Ligand CYC CU 200



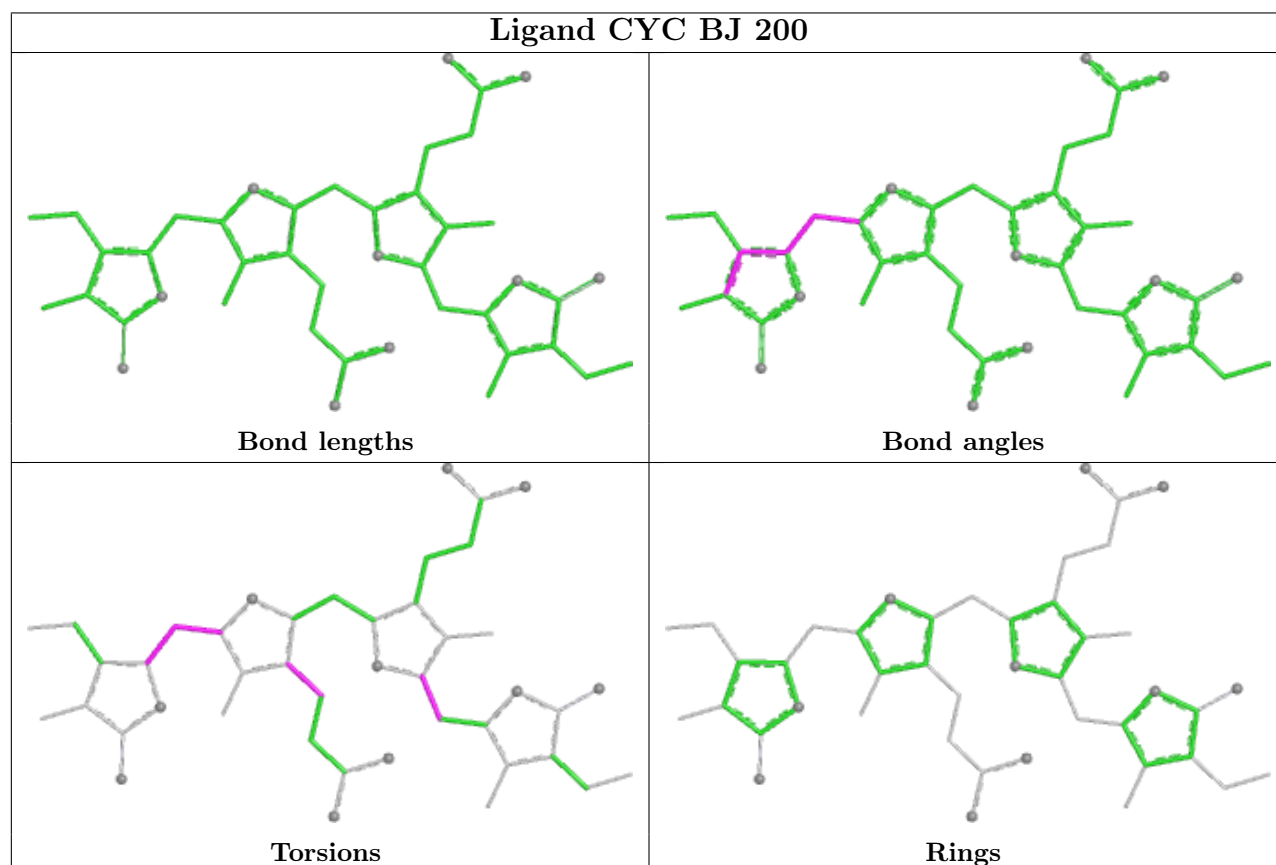
Ligand CYC DR 200



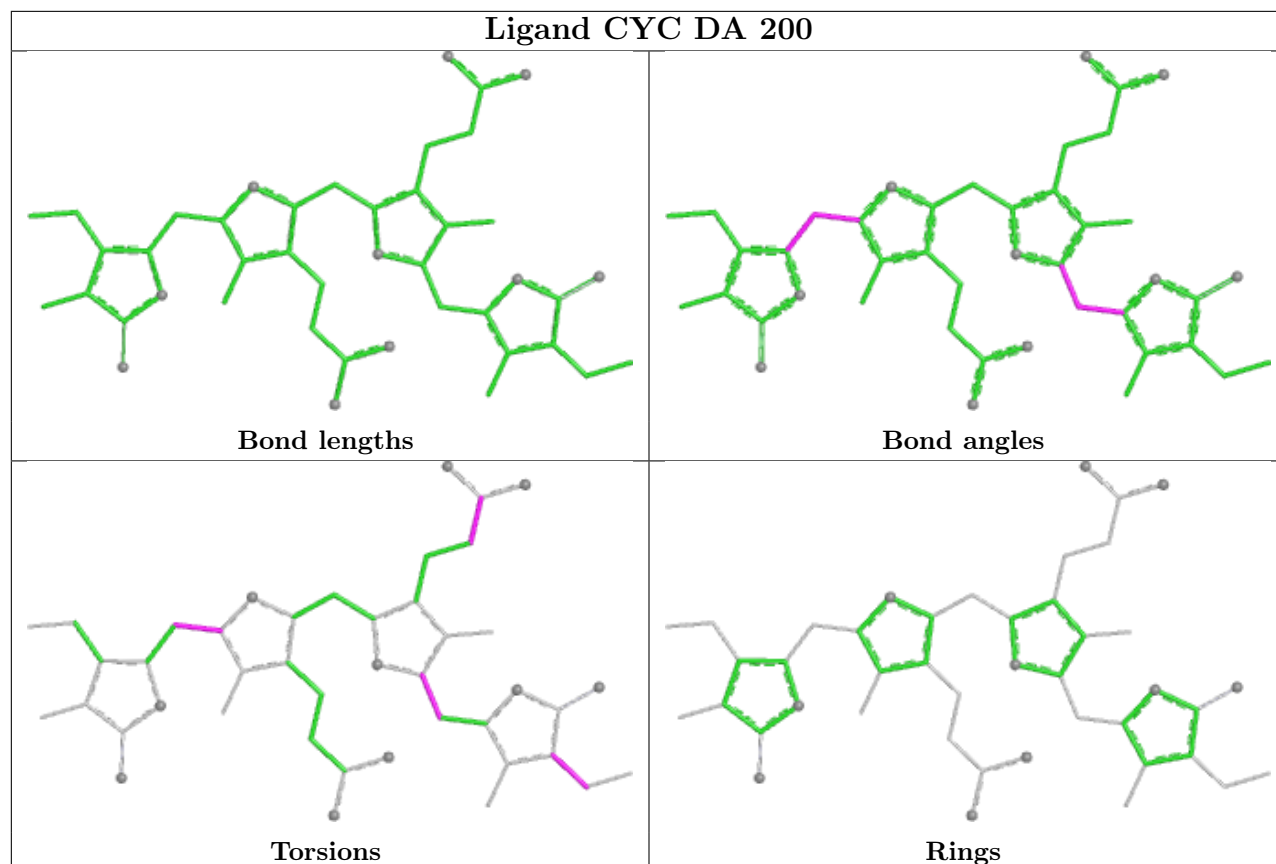
Ligand CYC CW 200



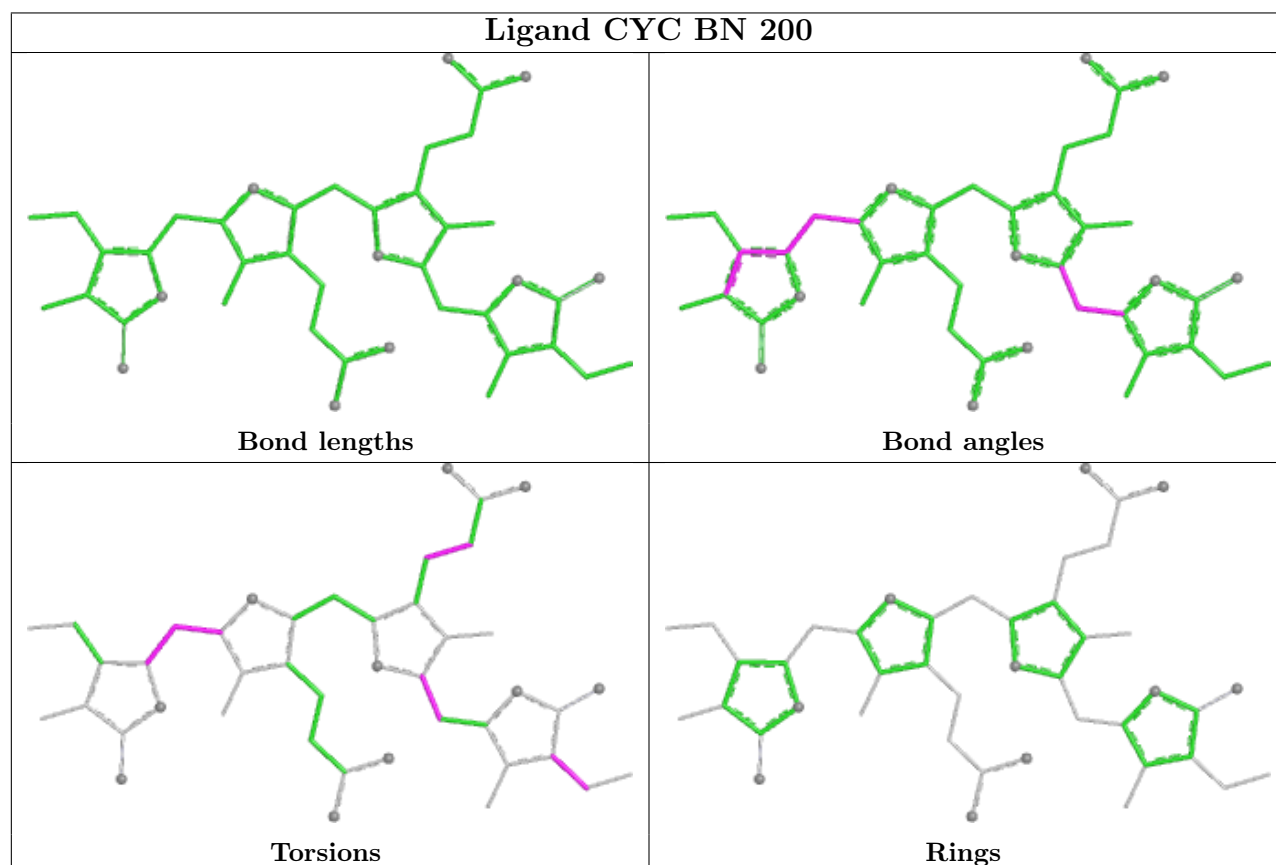
Ligand CYC BJ 200

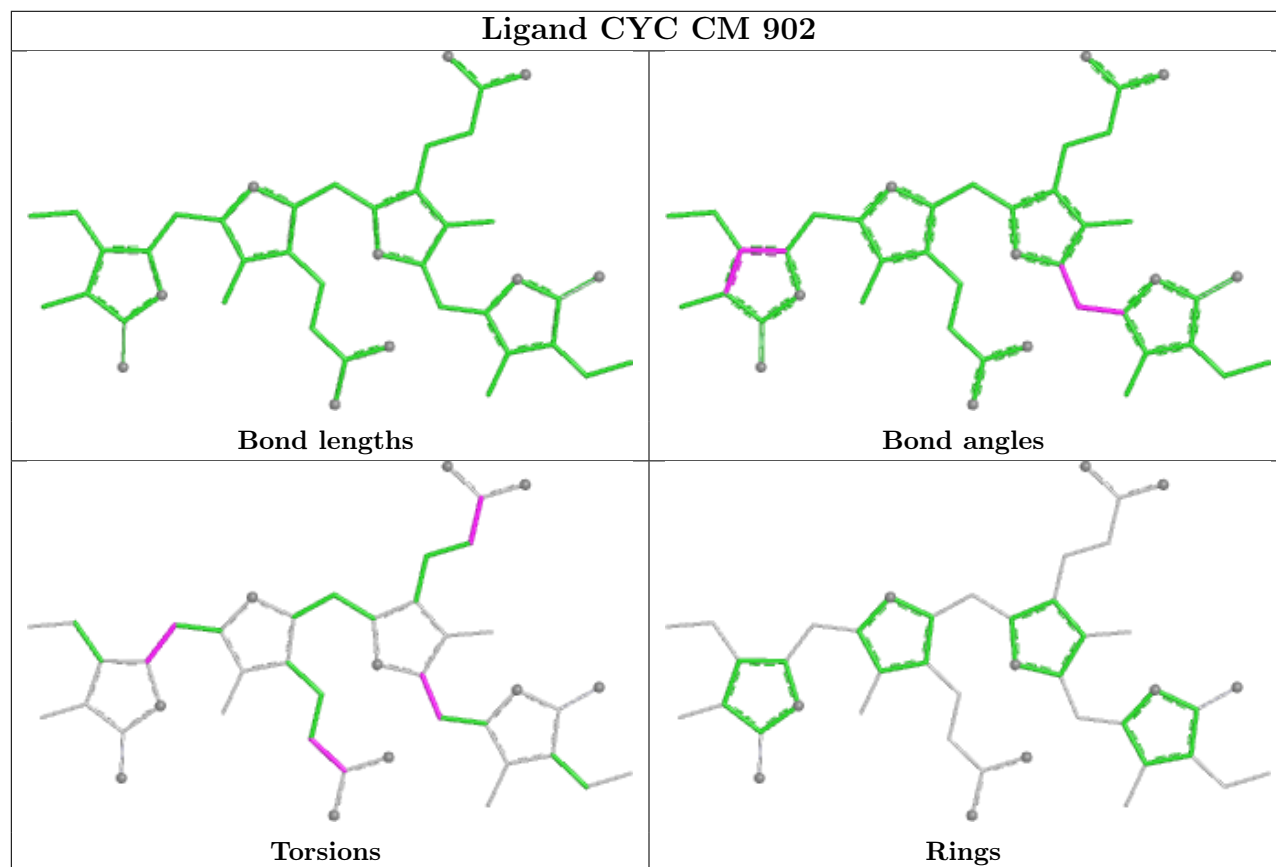
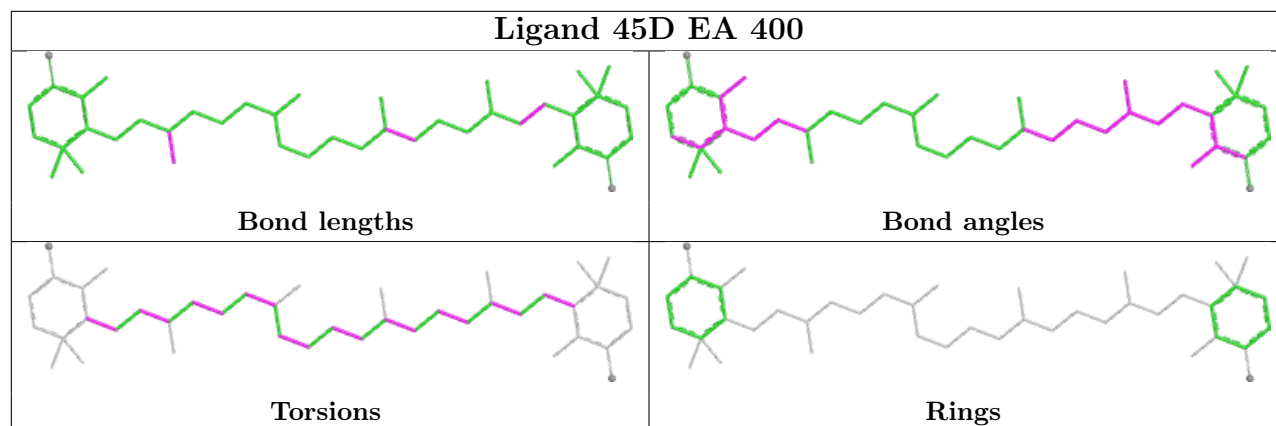


Ligand CYC DA 200

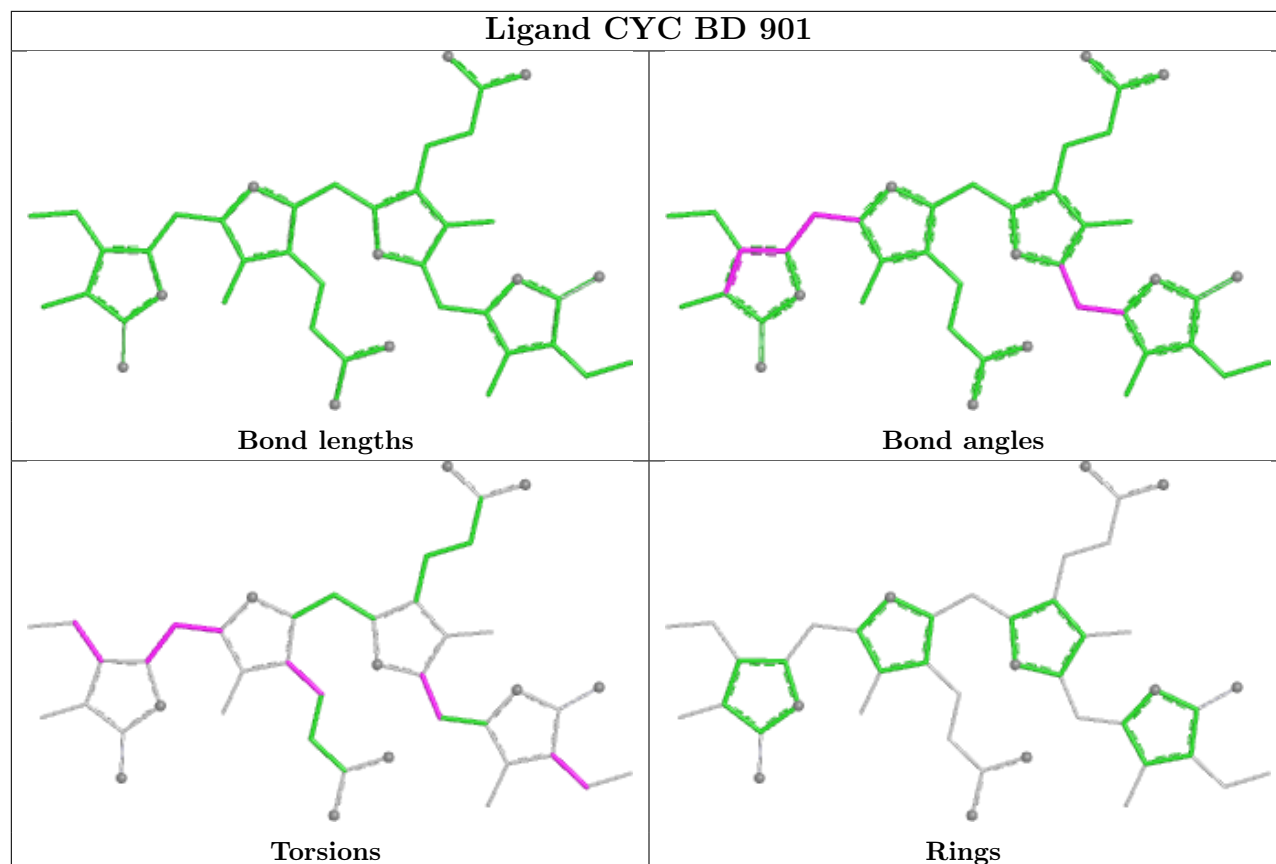


Ligand CYC BN 200

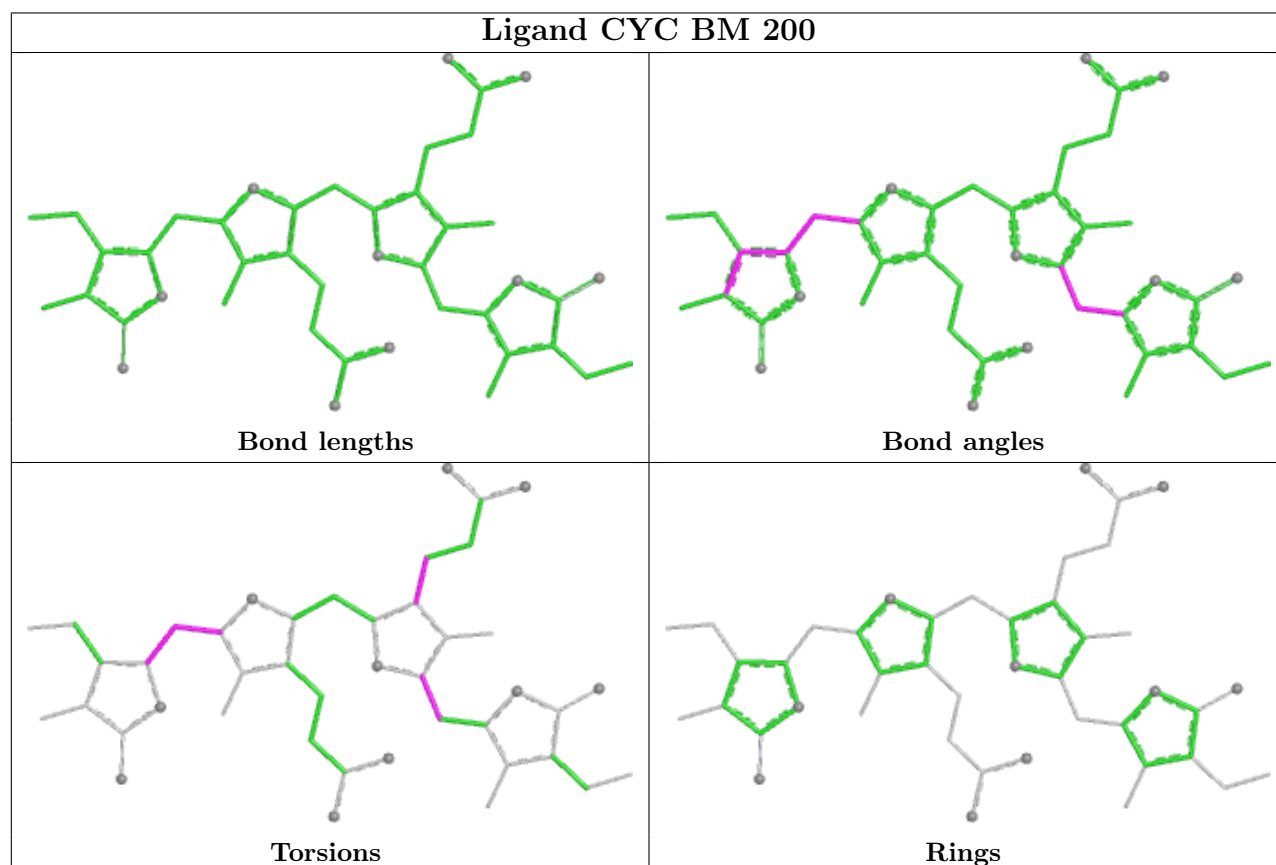


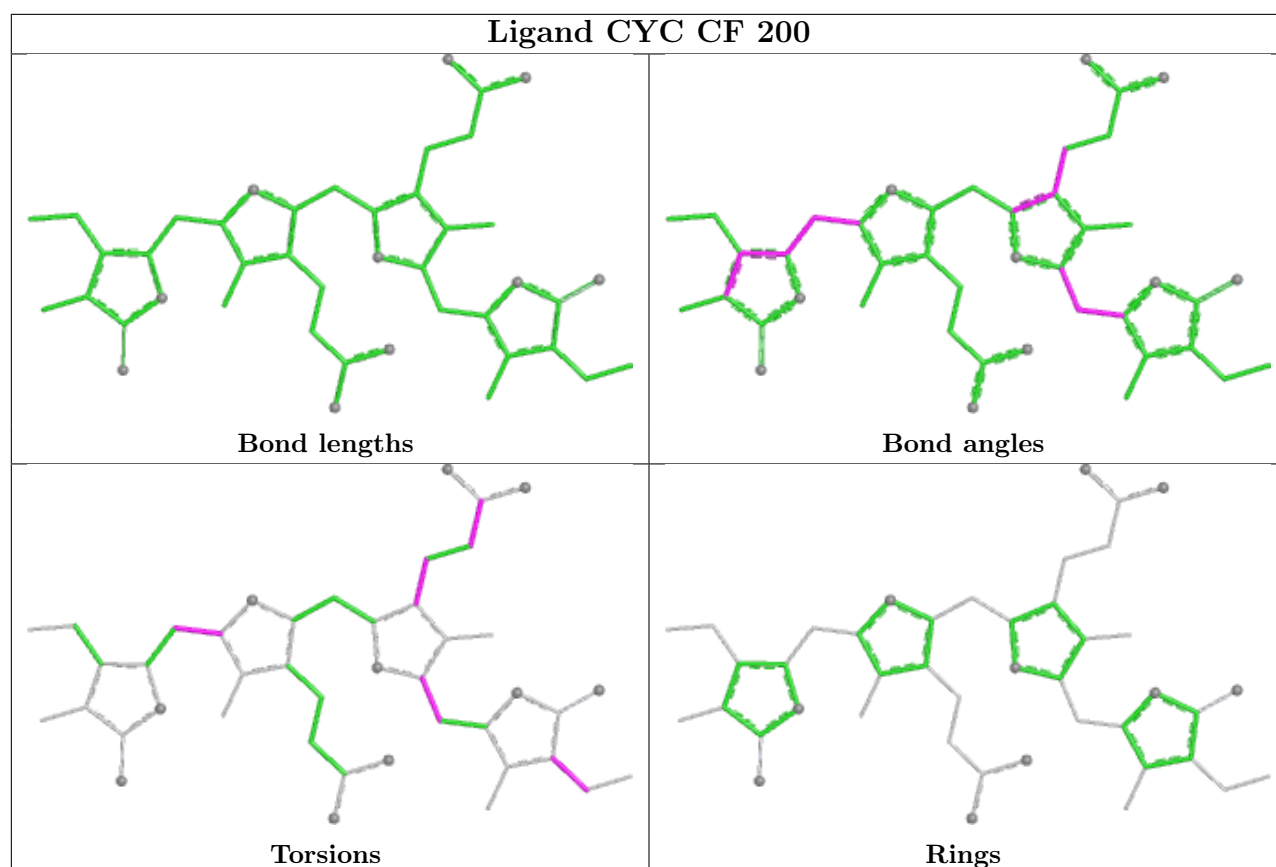


Ligand CYC BD 901



Ligand CYC BM 200





5.7 Other polymers [i](#)

There are no such residues in this entry.

5.8 Polymer linkage issues [i](#)

There are no chain breaks in this entry.

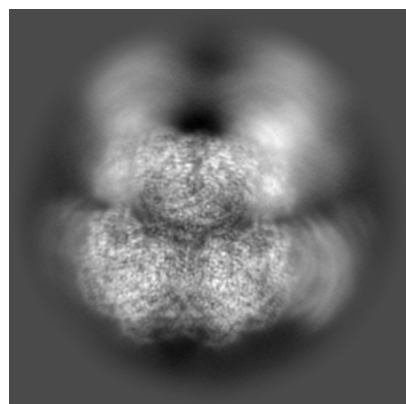
6 Map visualisation [i](#)

This section contains visualisations of the EMDB entry EMD-25030. These allow visual inspection of the internal detail of the map and identification of artifacts.

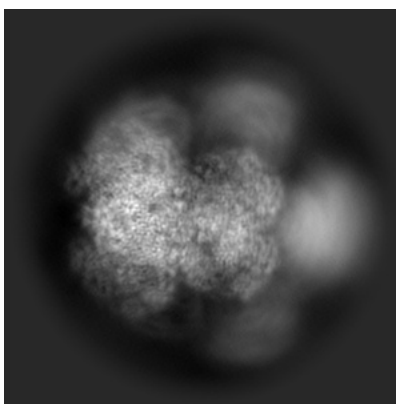
Images derived from a raw map, generated by summing the deposited half-maps, are presented below the corresponding image components of the primary map to allow further visual inspection and comparison with those of the primary map.

6.1 Orthogonal projections [i](#)

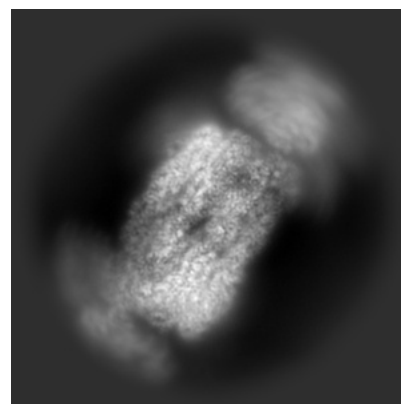
6.1.1 Primary map



X

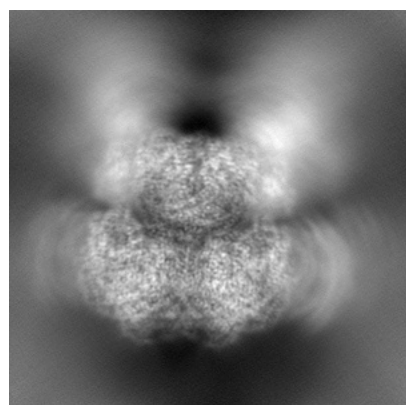


Y

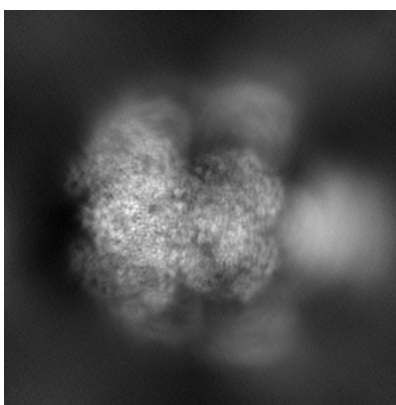


Z

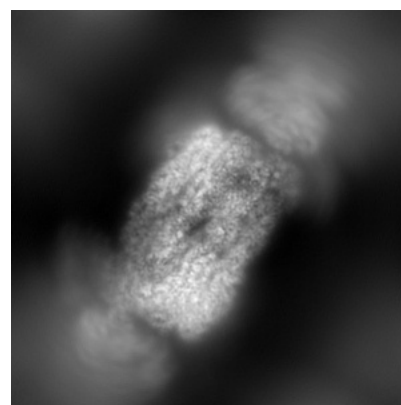
6.1.2 Raw map



X



Y

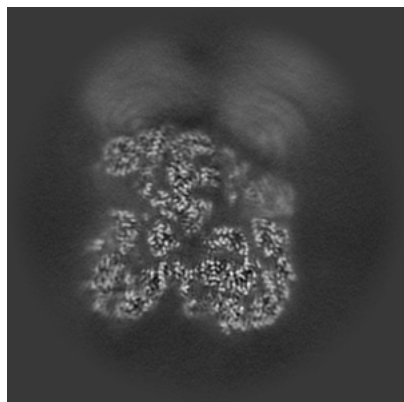


Z

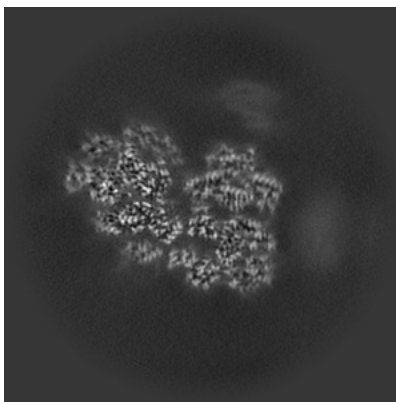
The images above show the map projected in three orthogonal directions.

6.2 Central slices [i](#)

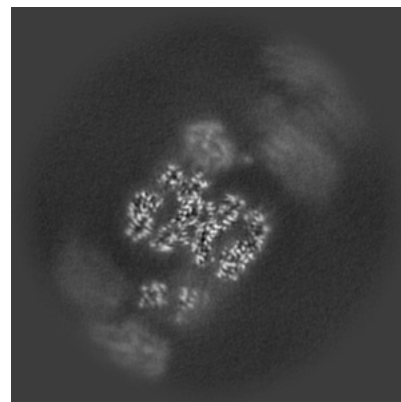
6.2.1 Primary map



X Index: 180

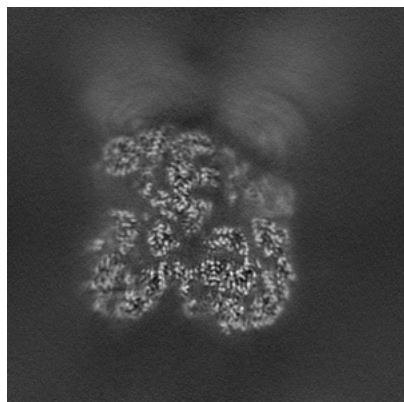


Y Index: 180

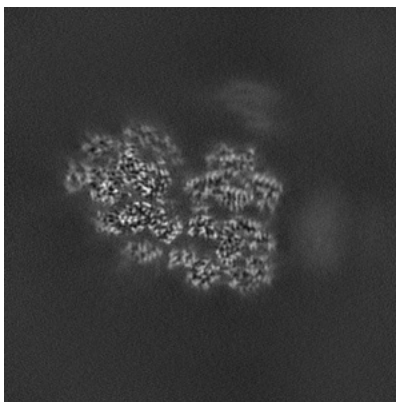


Z Index: 180

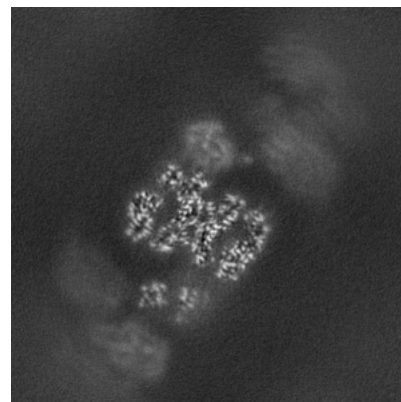
6.2.2 Raw map



X Index: 180



Y Index: 180

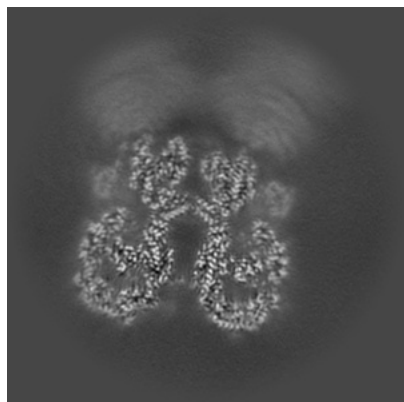


Z Index: 180

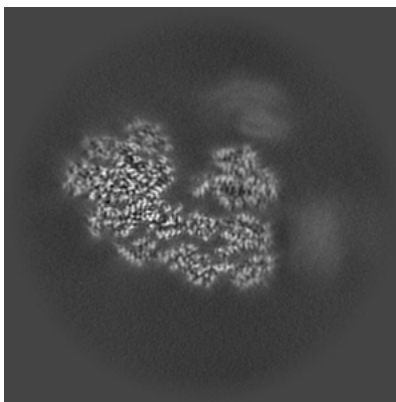
The images above show central slices of the map in three orthogonal directions.

6.3 Largest variance slices [i](#)

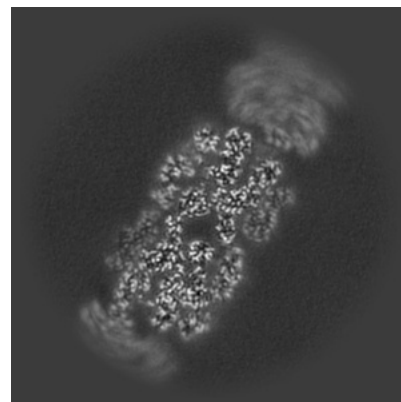
6.3.1 Primary map



X Index: 168

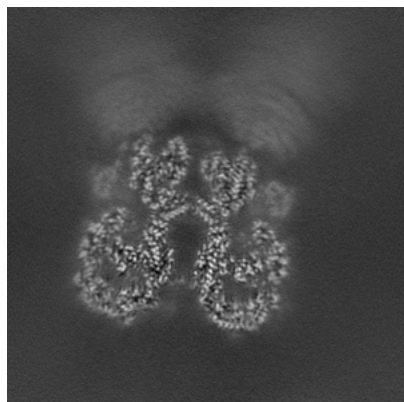


Y Index: 186

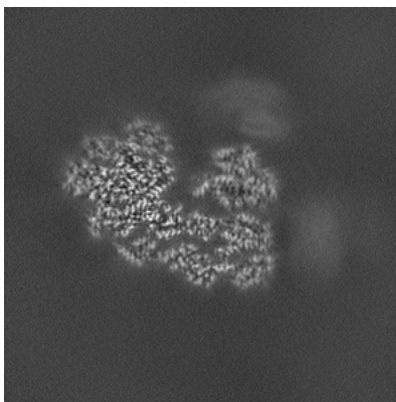


Z Index: 139

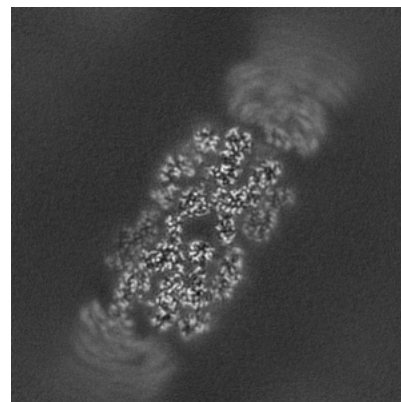
6.3.2 Raw map



X Index: 168



Y Index: 186

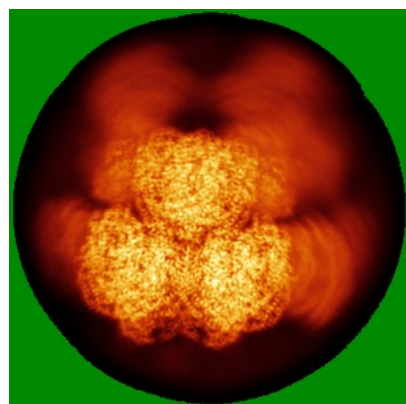


Z Index: 139

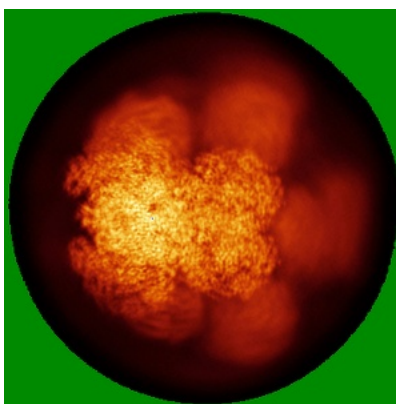
The images above show the largest variance slices of the map in three orthogonal directions.

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

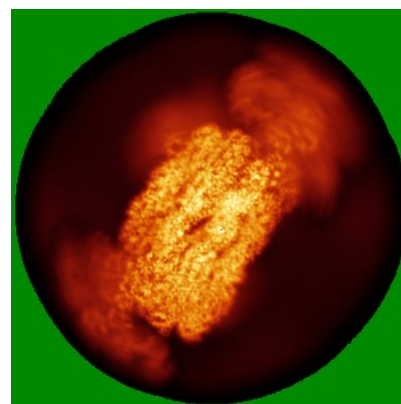
6.4.1 Primary map



X

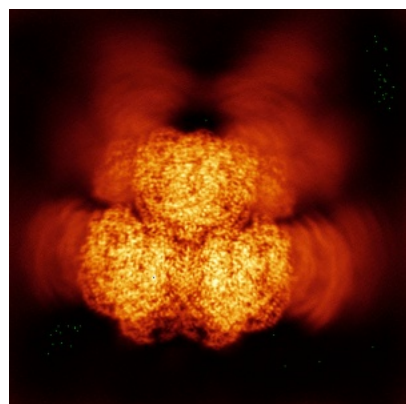


Y

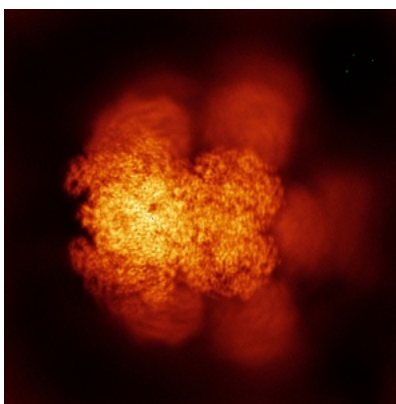


Z

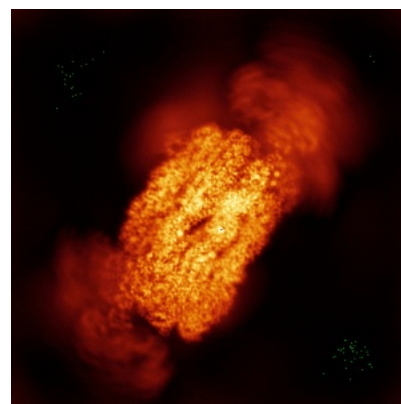
6.4.2 Raw map



X



Y

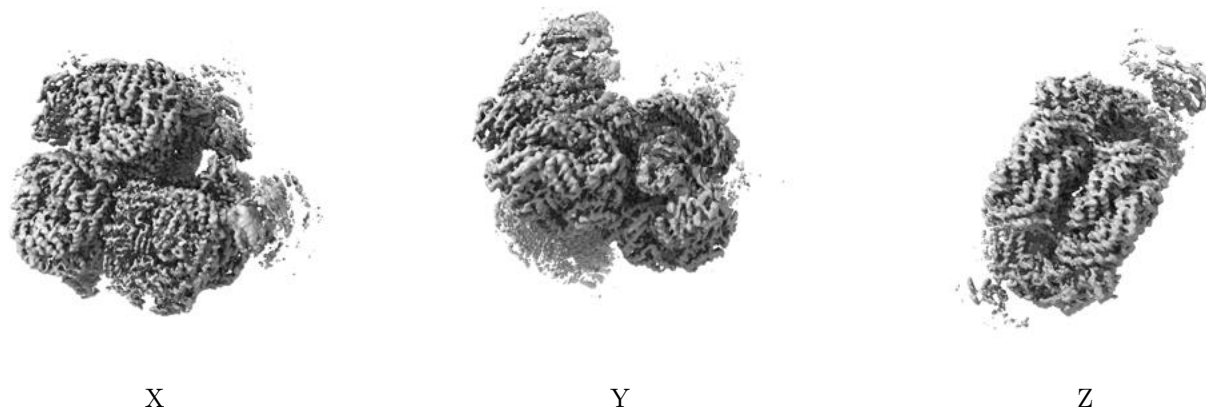


Z

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

6.5 Orthogonal surface views [i](#)

6.5.1 Primary map



The images above show the 3D surface view of the map at the recommended contour level 0.437. These images, in conjunction with the slice images, may facilitate assessment of whether an appropriate contour level has been provided.

6.5.2 Raw map



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

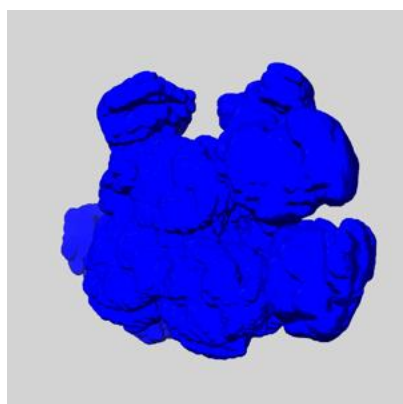
6.6 Mask visualisation [i](#)

This section shows the 3D surface view of the primary map at 50% transparency overlaid with the specified mask at 0% transparency

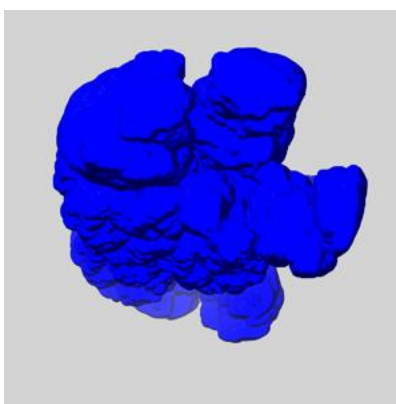
A mask typically either:

- Encompasses the whole structure
- Separates out a domain, a functional unit, a monomer or an area of interest from a larger structure

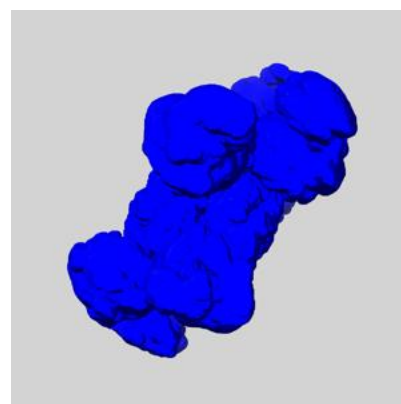
6.6.1 emd_25030_msk_1.map [i](#)



X



Y

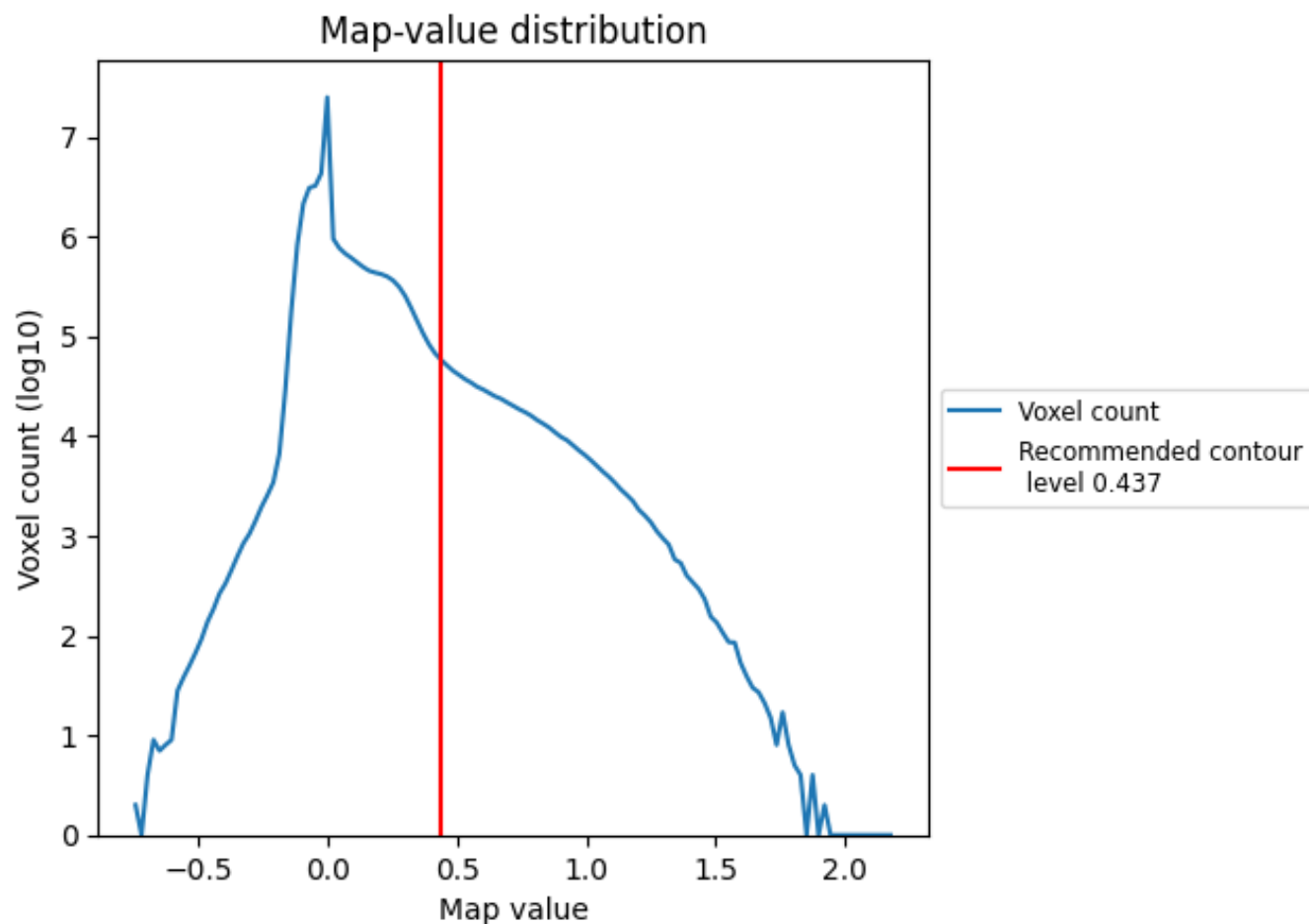


Z

7 Map analysis [i](#)

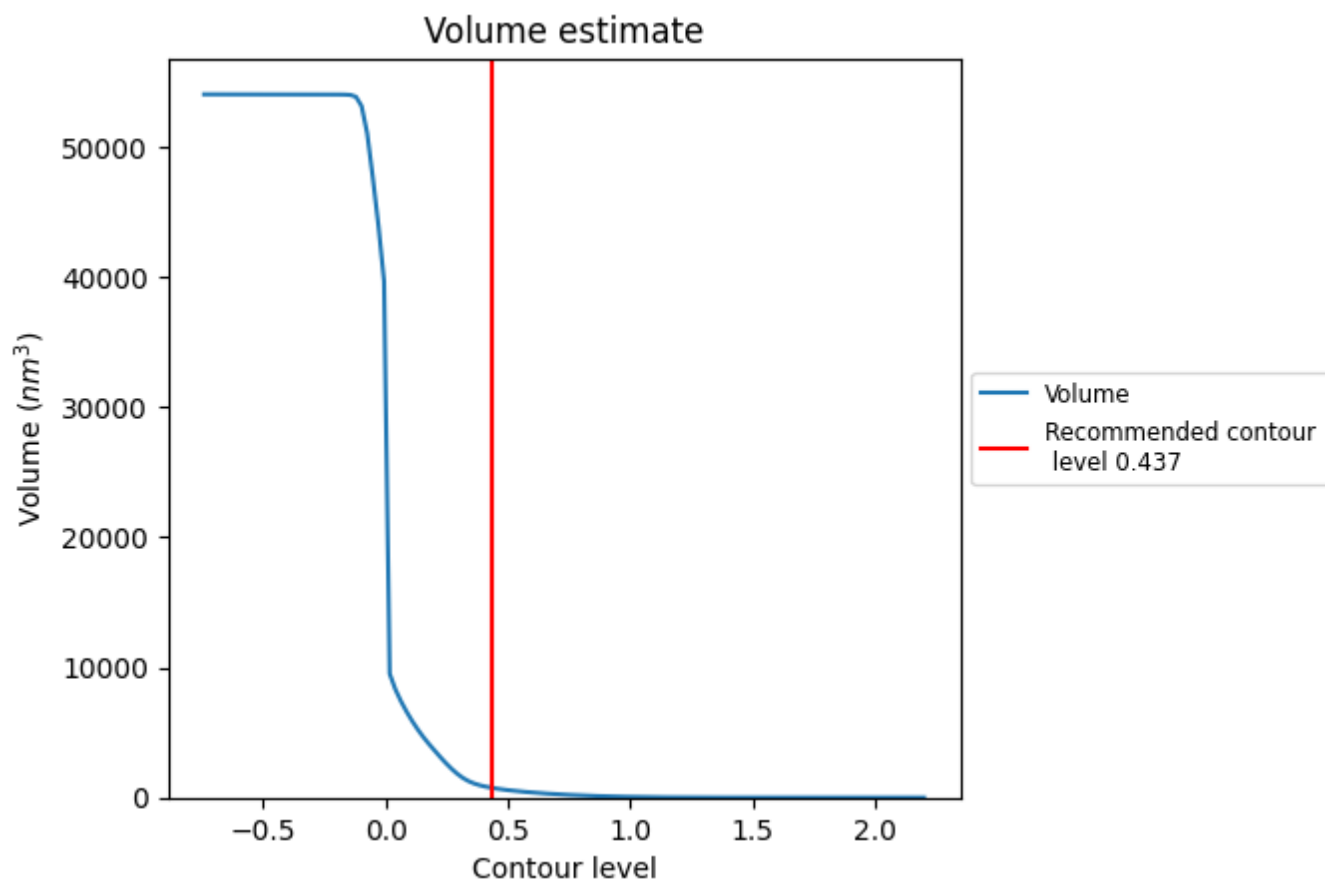
This section contains the results of statistical analysis of the map.

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.

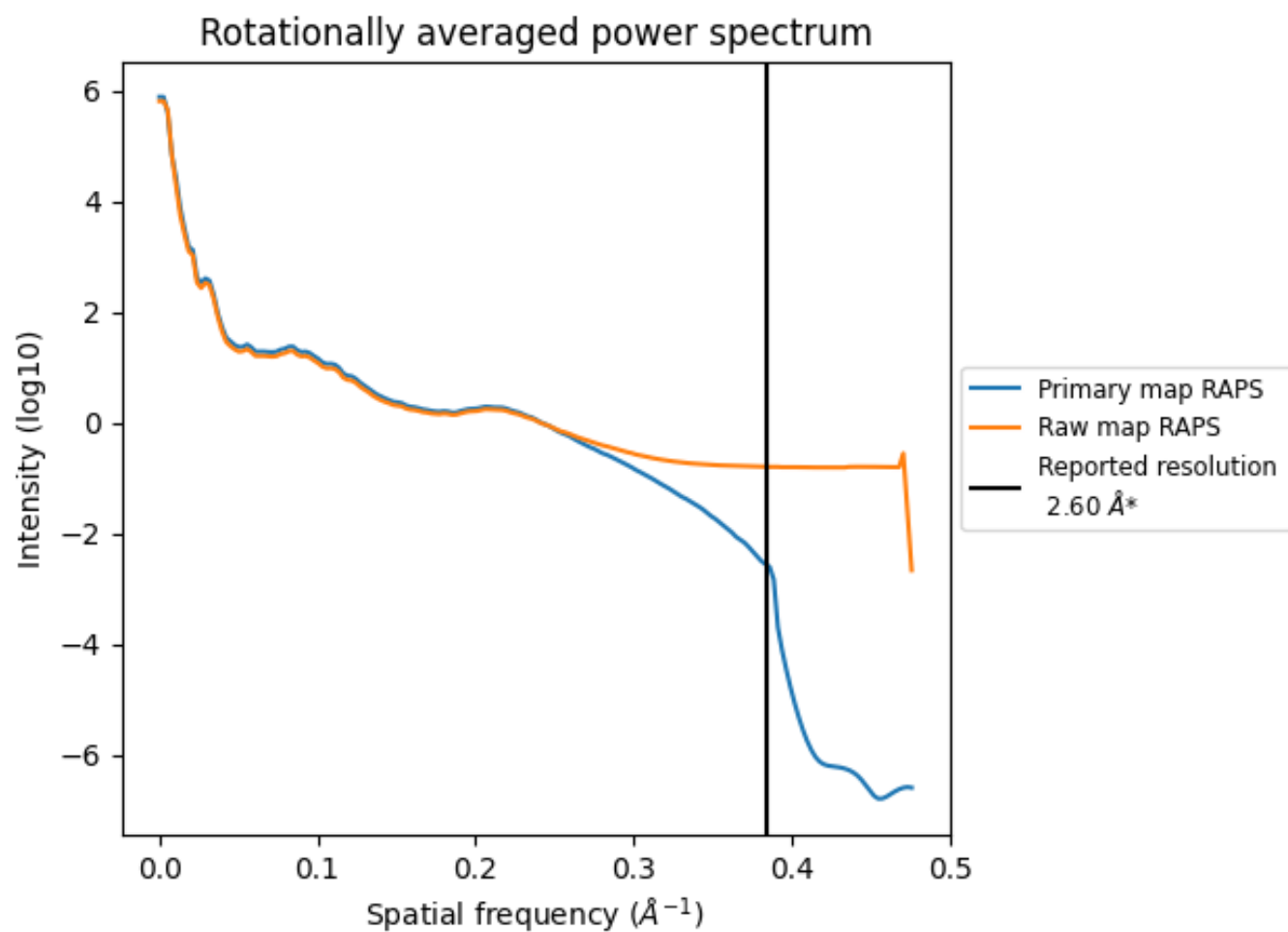
7.2 Volume estimate [i](#)



The volume at the recommended contour level is 737 nm³; this corresponds to an approximate mass of 666 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 ⓘ

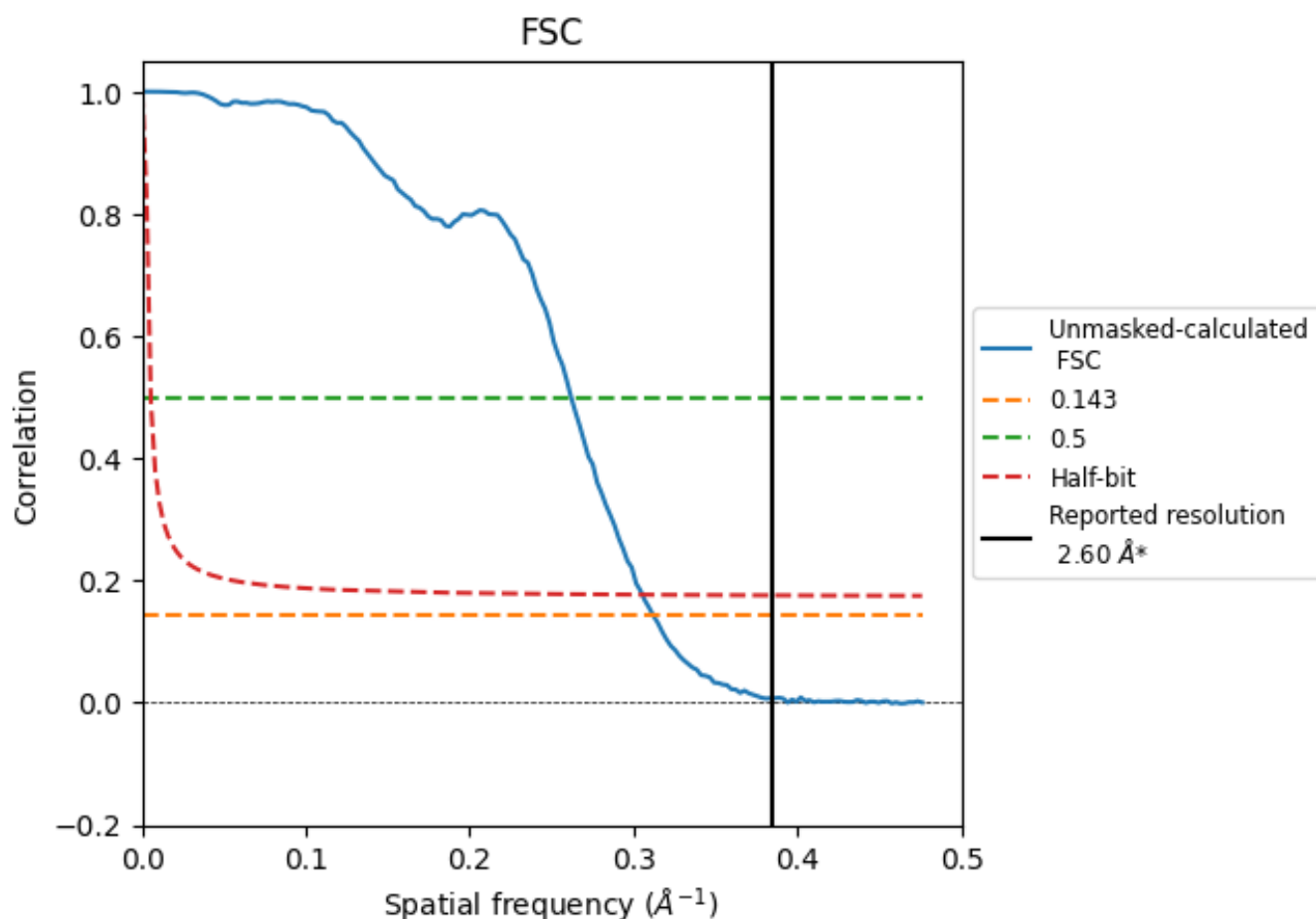


*Reported resolution corresponds to spatial frequency of 0.385 Å⁻¹

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 0.385 $\text{\AA}^{-1}$

8.2 Resolution estimates [i](#)

Resolution estimate (Å)	Estimation criterion (FSC cut-off)		
	0.143	0.5	Half-bit
Reported by author	2.60	-	-
Author-provided FSC curve	-	-	-
Unmasked-calculated*	3.21	3.82	3.28

*Resolution estimate based on FSC curve calculated by comparison of deposited half-maps. The value from deposited half-maps intersecting FSC 0.143 CUT-OFF 3.21 differs from the reported value 2.6 by more than 10 %

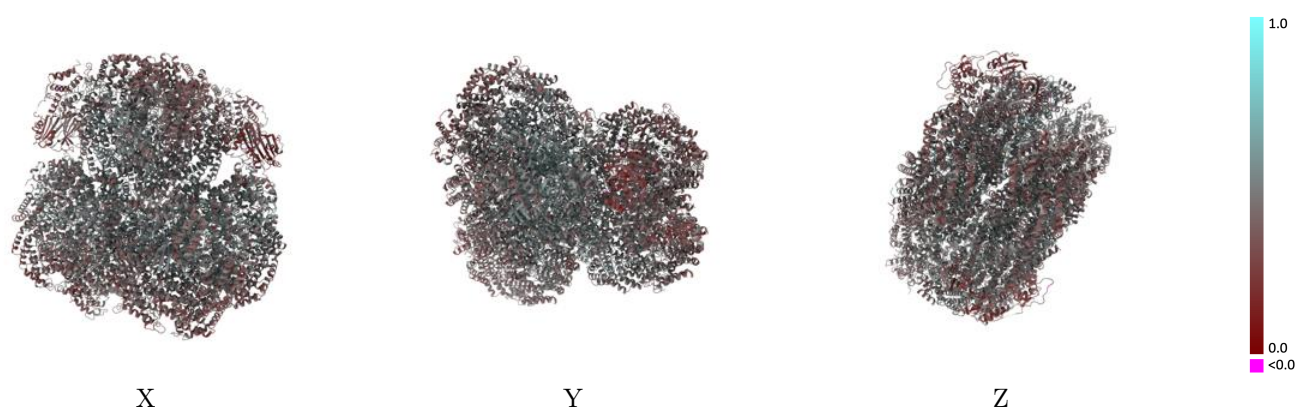
9 Map-model fit [i](#)

This section contains information regarding the fit between EMDB map EMD-25030 and PDB model 7SC9. Per-residue inclusion information can be found in section 3 on page 18.

9.1 Map-model overlay [i](#)

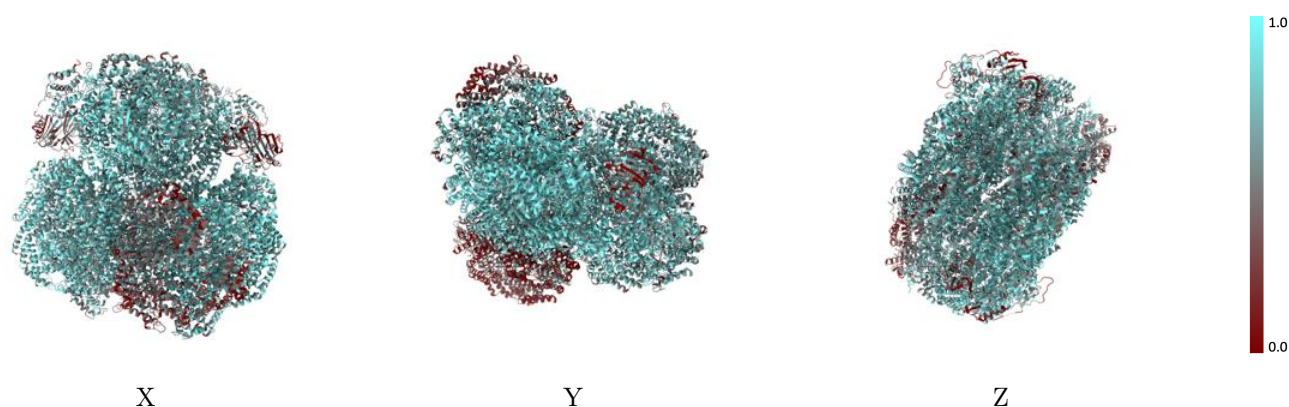
This section was not generated.

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



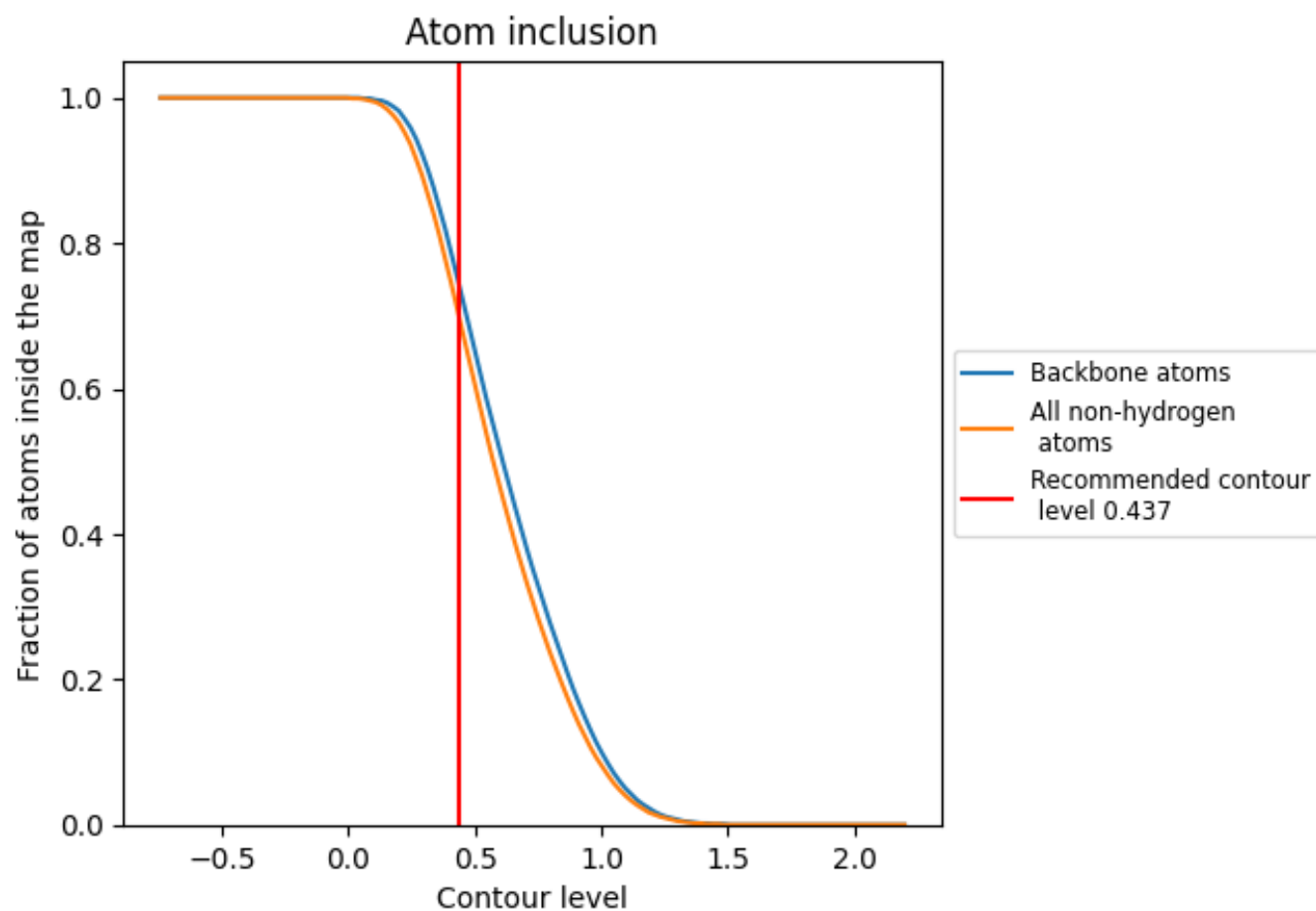
The images above show the model with each residue coloured according to its Q-score. This shows their resolvability in the map with higher Q-score values reflecting better resolvability. Please note: Q-score is calculating the resolvability of atoms, and thus high values are only expected at resolutions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

9.3 Atom inclusion mapped to coordinate model [i](#)



The images above show the model with each residue coloured according to its atom inclusion. This shows to what extent they are inside the map at the recommended contour level (0.437).

9.4 Atom inclusion [i](#)



At the recommended contour level, 74% of all backbone atoms, 70% of all non-hydrogen atoms, are inside the map.

9.5 Map-model fit summary ⓘ

The table lists the average atom inclusion at the recommended contour level (0.437) and Q-score for the entire model and for each chain.

Chain	Atom inclusion	Q-score
All	 0.7000	 0.4490
AA	 0.1520	 0.4250
AB	 0.1360	 0.4150
AC	 0.1510	 0.3650
AD	 0.1330	 0.3650
AE	 0.1950	 0.3820
AF	 0.1110	 0.4160
AH	 0.7940	 0.4590
AI	 0.8780	 0.5300
AJ	 0.8590	 0.5140
AK	 0.8170	 0.4500
AL	 0.8520	 0.4480
AN	 0.8150	 0.4280
AO	 0.8740	 0.5150
AP	 0.8640	 0.5240
AQ	 0.8760	 0.5290
AR	 0.8240	 0.4710
AS	 0.8470	 0.4560
AU	 0.7730	 0.4120
AV	 0.7770	 0.3790
AW	 0.8360	 0.4240
AX	 0.8320	 0.4630
AY	 0.8430	 0.4670
AZ	 0.7550	 0.4430
BA	 0.2260	 0.4480
BB	 0.6910	 0.4870
BD	 0.8150	 0.4770
BF	 0.7100	 0.4110
BG	 0.7790	 0.4650
BH	 0.4920	 0.3840
BI	 0.3540	 0.4400
BJ	 0.3880	 0.4340
BK	 0.3630	 0.3980
BL	 0.3660	 0.3950
BM	 0.4280	 0.4030



Continued on next page...

Continued from previous page...

Chain	Atom inclusion	Q-score
BN	 0.3660	 0.4290
BP	 0.8290	 0.4820
BQ	 0.8910	 0.5340
BR	 0.8610	 0.5150
BS	 0.8450	 0.4730
BT	 0.8600	 0.4790
BV	 0.8430	 0.4430
BW	 0.8770	 0.5250
BX	 0.8760	 0.5350
BY	 0.8810	 0.5450
BZ	 0.8640	 0.5050
CA	 0.8620	 0.4900
CC	 0.7390	 0.4680
CD	 0.7860	 0.4440
CE	 0.8190	 0.4310
CF	 0.8330	 0.4530
CG	 0.8620	 0.4790
CH	 0.8350	 0.4820
CJ	 0.4930	 0.4900
CK	 0.7020	 0.5160
CM	 0.8370	 0.4890
CO	 0.7360	 0.4290
CP	 0.7950	 0.4820
CQ	 0.5480	 0.3920
CR	 0.7290	 0.3970
CS	 0.7170	 0.4240
CT	 0.6720	 0.4280
CU	 0.7300	 0.4660
CV	 0.8490	 0.4910
CW	 0.8240	 0.4460
CY	 0.6870	 0.4010
CZ	 0.6930	 0.4120
DA	 0.8000	 0.4400
DB	 0.8350	 0.4930
DC	 0.8360	 0.5030
DD	 0.7940	 0.4570
DF	 0.8370	 0.5040
DG	 0.4650	 0.3670
DH	 0.6460	 0.4310
DI	 0.4850	 0.3140
DJ	 0.7160	 0.4080
DK	 0.6920	 0.4030

Continued on next page...

Continued from previous page...

Chain	Atom inclusion	Q-score
DL	 0.7080	 0.3920
DM	 0.7990	 0.4600
DN	 0.8510	 0.4840
DO	 0.8420	 0.4730
DQ	 0.6830	 0.3800
DR	 0.7130	 0.4220
DS	 0.8300	 0.4720
DT	 0.8240	 0.4920
DU	 0.8310	 0.4930
DV	 0.7550	 0.4100
DX	 0.8290	 0.5090
DY	 0.4460	 0.4080
DZ	 0.6390	 0.4060
EA	 0.5100	 0.3400