



Full wwPDB X-ray Structure Validation Report ⓘ

May 14, 2020 – 01:46 pm BST

PDB ID : 4PKO
Title : Crystal structure of the Football-shaped GroEL-GroES2-(ADPBeFx)₁₄ complex
Authors : Fei, X.; Ye, X.; Laronde-Leblanc, N.; Lorimer, G.H.
Deposited on : 2014-05-15
Resolution : 3.84 Å(reported)

This is a Full wwPDB X-ray Structure 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/XrayValidationReportHelp>

with specific help available everywhere you see the ⓘ symbol.

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

MolProbity : 4.02b-467
Mogul : 1.8.5 (274361), CSD as541be (2020)
Xtriage (Phenix) : 1.13
EDS : 2.11
buster-report : 1.1.7 (2018)
Percentile statistics : 20191225.v01 (using entries in the PDB archive December 25th 2019)
Refmac : 5.8.0158
CCP4 : 7.0.044 (Gargrove)
Ideal geometry (proteins) : Engh & Huber (2001)
Ideal geometry (DNA, RNA) : Parkinson et al. (1996)
Validation Pipeline (wwPDB-VP) : 2.11

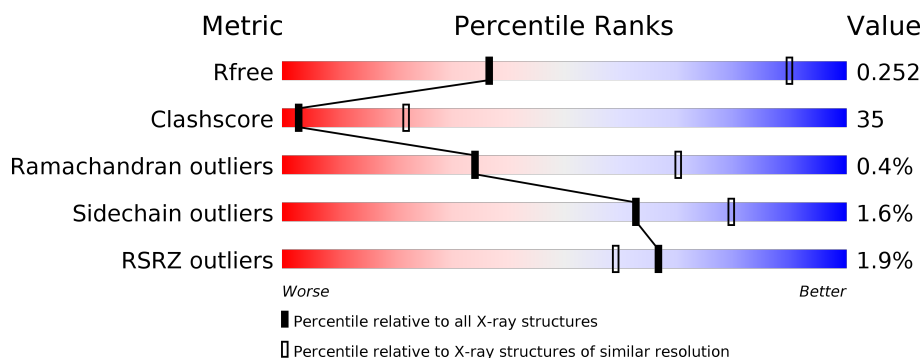
1 Overall quality at a glance

The following experimental techniques were used to determine the structure:

X-RAY DIFFRACTION

The reported resolution of this entry is 3.84 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.





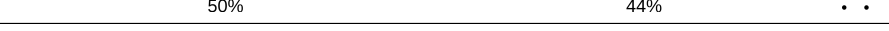

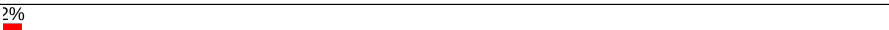





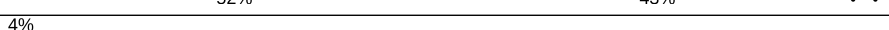
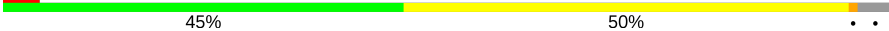










Metric	Whole archive (#Entries)	Similar resolution (#Entries, resolution range(Å))
R_{free}	130704	1242 (4.08-3.60)
Clashscore	141614	1004 (4.04-3.64)
Ramachandran outliers	138981	1003 (4.06-3.62)
Sidechain outliers	138945	1266 (4.08-3.60)
RSRZ outliers	127900	1149 (4.08-3.60)

The table below summarises the geometric issues observed across the polymeric chains and their fit to the electron density. The red, orange, yellow and green segments on the lower 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 electron density. The numeric value is given above the bar.

Mol	Chain	Length	Quality of chain
1	A	548	<div> <div>2%</div> <div> <div></div> <div>53%</div> <div>41%</div> <div>• •</div> </div> </div>
1	B	548	<div> <div>%</div> <div> <div></div> <div>50%</div> <div>45%</div> <div>• •</div> </div> </div>
1	C	548	<div> <div>%</div> <div> <div></div> <div>53%</div> <div>42%</div> <div>• •</div> </div> </div>
1	D	548	<div> <div></div> <div> <div></div> <div>57%</div> <div>38%</div> <div>• •</div> </div> </div>
1	E	548	<div> <div>2%</div> <div> <div></div> <div>52%</div> <div>42%</div> <div>• •</div> </div> </div>
1	F	548	<div> <div>3%</div> <div> <div></div> <div>51%</div> <div>44%</div> <div>• •</div> </div> </div>

Continued on next page...

Continued from previous page...

Mol	Chain	Length	Quality of chain
1	G	548	
1	H	548	
1	I	548	
1	J	548	
1	K	548	
1	L	548	
1	M	548	
1	N	548	
2	1	97	
2	2	97	
2	O	97	
2	P	97	
2	Q	97	
2	R	97	
2	S	97	
2	T	97	
2	U	97	
2	V	97	
2	W	97	
2	X	97	
2	Y	97	
2	Z	97	

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

Mol	Type	Chain	Res	Chirality	Geometry	Clashes	Electron density
4	BEF	A	602	-	-	X	-
4	BEF	B	602	-	-	X	-
4	BEF	C	602	-	-	X	-
4	BEF	H	602	-	-	X	-
4	BEF	I	602	-	-	X	-
4	BEF	J	602	-	-	X	-
4	BEF	K	602	-	-	X	-
4	BEF	L	602	-	-	X	-
4	BEF	N	602	-	-	X	-

2 Entry composition [i](#)

There are 6 unique types of molecules in this entry. The entry contains 64559 atoms, of which 0 are hydrogens and 0 are deuteriums.

In the tables below, the ZeroOcc column contains the number of atoms modelled with zero occupancy, 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.

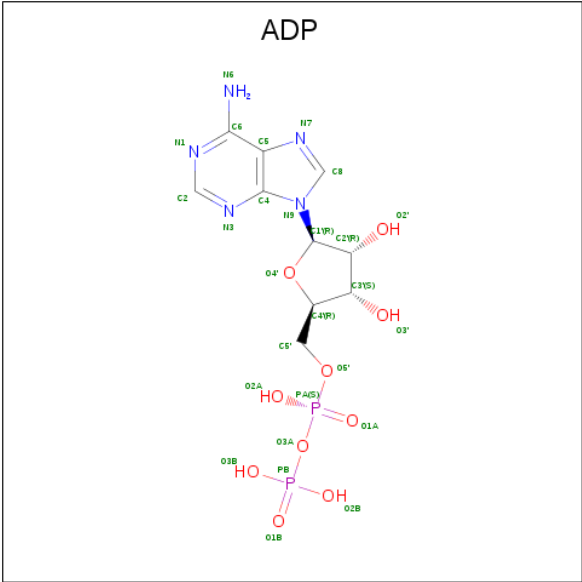
- Molecule 1 is a protein called 60 kDa chaperonin.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
1	A	524	Total	C	N	O	S	0	0	0
			3855	2397	665	773	20			
1	B	524	Total	C	N	O	S	0	0	0
			3855	2397	665	773	20			
1	C	524	Total	C	N	O	S	0	0	0
			3855	2397	665	773	20			
1	D	524	Total	C	N	O	S	0	0	0
			3855	2397	665	773	20			
1	E	524	Total	C	N	O	S	0	0	0
			3855	2397	665	773	20			
1	F	524	Total	C	N	O	S	0	0	0
			3855	2397	665	773	20			
1	G	524	Total	C	N	O	S	0	0	0
			3855	2397	665	773	20			
1	H	524	Total	C	N	O	S	0	0	0
			3855	2397	665	773	20			
1	I	524	Total	C	N	O	S	0	0	0
			3855	2397	665	773	20			
1	J	524	Total	C	N	O	S	0	0	0
			3855	2397	665	773	20			
1	K	524	Total	C	N	O	S	0	0	0
			3855	2397	665	773	20			
1	L	524	Total	C	N	O	S	0	0	0
			3855	2397	665	773	20			
1	M	524	Total	C	N	O	S	0	0	0
			3855	2397	665	773	20			
1	N	524	Total	C	N	O	S	0	0	0
			3855	2397	665	773	20			

- Molecule 2 is a protein called 10 kDa chaperonin.

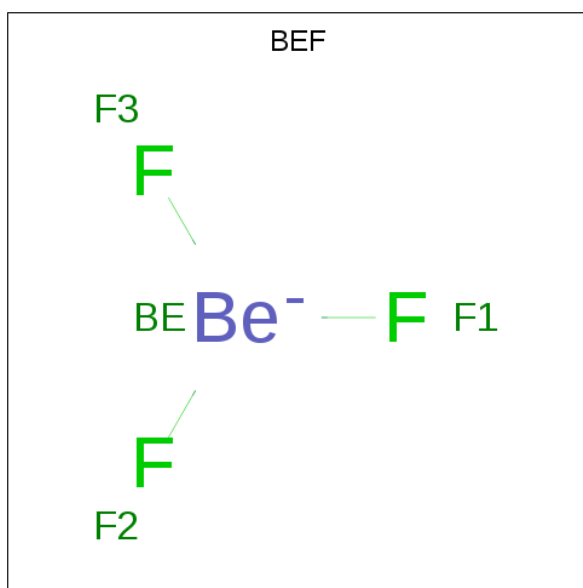
Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
2	O	95	Total	C	N	O	S	0	0	0
			711	445	124	141	1			
2	P	97	Total	C	N	O	S	0	0	0
			727	454	127	144	2			
2	Q	96	Total	C	N	O	S	0	0	0
			719	449	126	143	1			
2	R	97	Total	C	N	O	S	0	0	0
			727	454	127	144	2			
2	S	96	Total	C	N	O	S	0	0	0
			719	449	126	143	1			
2	T	97	Total	C	N	O	S	0	0	0
			727	454	127	144	2			
2	U	97	Total	C	N	O	S	0	0	0
			727	454	127	144	2			
2	V	97	Total	C	N	O	S	0	0	0
			727	454	127	144	2			
2	W	97	Total	C	N	O	S	0	0	0
			727	454	127	144	2			
2	X	97	Total	C	N	O	S	0	0	0
			727	454	127	144	2			
2	Y	96	Total	C	N	O	S	0	0	0
			722	451	126	143	2			
2	Z	95	Total	C	N	O	S	0	0	0
			713	446	125	140	2			
2	1	97	Total	C	N	O	S	0	0	0
			727	454	127	144	2			
2	2	97	Total	C	N	O	S	0	0	0
			727	454	127	144	2			

- Molecule 3 is ADENOSINE-5'-DIPHOSPHATE (three-letter code: ADP) (formula: $C_{10}H_{15}N_5O_{10}P_2$).



Mol	Chain	Residues	Atoms					ZeroOcc	AltConf
3	A	1	Total	C	N	O	P	0	0
			27	10	5	10	2		
3	B	1	Total	C	N	O	P	0	0
			27	10	5	10	2		
3	C	1	Total	C	N	O	P	0	0
			27	10	5	10	2		
3	D	1	Total	C	N	O	P	0	0
			27	10	5	10	2		
3	E	1	Total	C	N	O	P	0	0
			27	10	5	10	2		
3	F	1	Total	C	N	O	P	0	0
			27	10	5	10	2		
3	G	1	Total	C	N	O	P	0	0
			27	10	5	10	2		
3	H	1	Total	C	N	O	P	0	0
			27	10	5	10	2		
3	I	1	Total	C	N	O	P	0	0
			27	10	5	10	2		
3	J	1	Total	C	N	O	P	0	0
			27	10	5	10	2		
3	K	1	Total	C	N	O	P	0	0
			27	10	5	10	2		
3	L	1	Total	C	N	O	P	0	0
			27	10	5	10	2		
3	M	1	Total	C	N	O	P	0	0
			27	10	5	10	2		
3	N	1	Total	C	N	O	P	0	0
			27	10	5	10	2		

- Molecule 4 is BERYLLIUM TRIFLUORIDE ION (three-letter code: BEF) (formula: BeF₃).



Mol	Chain	Residues	Atoms			ZeroOcc	AltConf
4	A	1	Total 4	Be 1	F 3	0	0
4	B	1	Total 4	Be 1	F 3	0	0
4	C	1	Total 4	Be 1	F 3	0	0
4	D	1	Total 4	Be 1	F 3	0	0
4	E	1	Total 4	Be 1	F 3	0	0
4	F	1	Total 4	Be 1	F 3	0	0
4	G	1	Total 4	Be 1	F 3	0	0
4	H	1	Total 4	Be 1	F 3	0	0
4	I	1	Total 4	Be 1	F 3	0	0
4	J	1	Total 4	Be 1	F 3	0	0
4	K	1	Total 4	Be 1	F 3	0	0
4	L	1	Total 4	Be 1	F 3	0	0
4	M	1	Total 4	Be 1	F 3	0	0

Continued on next page...

Continued from previous page...

Mol	Chain	Residues	Atoms			ZeroOcc	AltConf
4	N	1	Total	Be	F	0	0
			4	1	3		

- Molecule 5 is MAGNESIUM ION (three-letter code: MG) (formula: Mg).

Mol	Chain	Residues	Atoms		ZeroOcc	AltConf
5	G	1	Total	Mg	0	0
			1	1		
5	J	1	Total	Mg	0	0
			1	1		
5	D	1	Total	Mg	0	0
			1	1		
5	K	1	Total	Mg	0	0
			1	1		
5	E	1	Total	Mg	0	0
			1	1		
5	H	1	Total	Mg	0	0
			1	1		
5	B	1	Total	Mg	0	0
			1	1		
5	I	1	Total	Mg	0	0
			1	1		
5	C	1	Total	Mg	0	0
			1	1		
5	A	1	Total	Mg	0	0
			1	1		
5	N	1	Total	Mg	0	0
			1	1		
5	L	1	Total	Mg	0	0
			1	1		
5	F	1	Total	Mg	0	0
			1	1		
5	M	1	Total	Mg	0	0
			1	1		

- Molecule 6 is POTASSIUM ION (three-letter code: K) (formula: K).

Mol	Chain	Residues	Atoms		ZeroOcc	AltConf
6	G	1	Total	K	0	0
			1	1		
6	J	1	Total	K	0	0
			1	1		

Continued on next page...

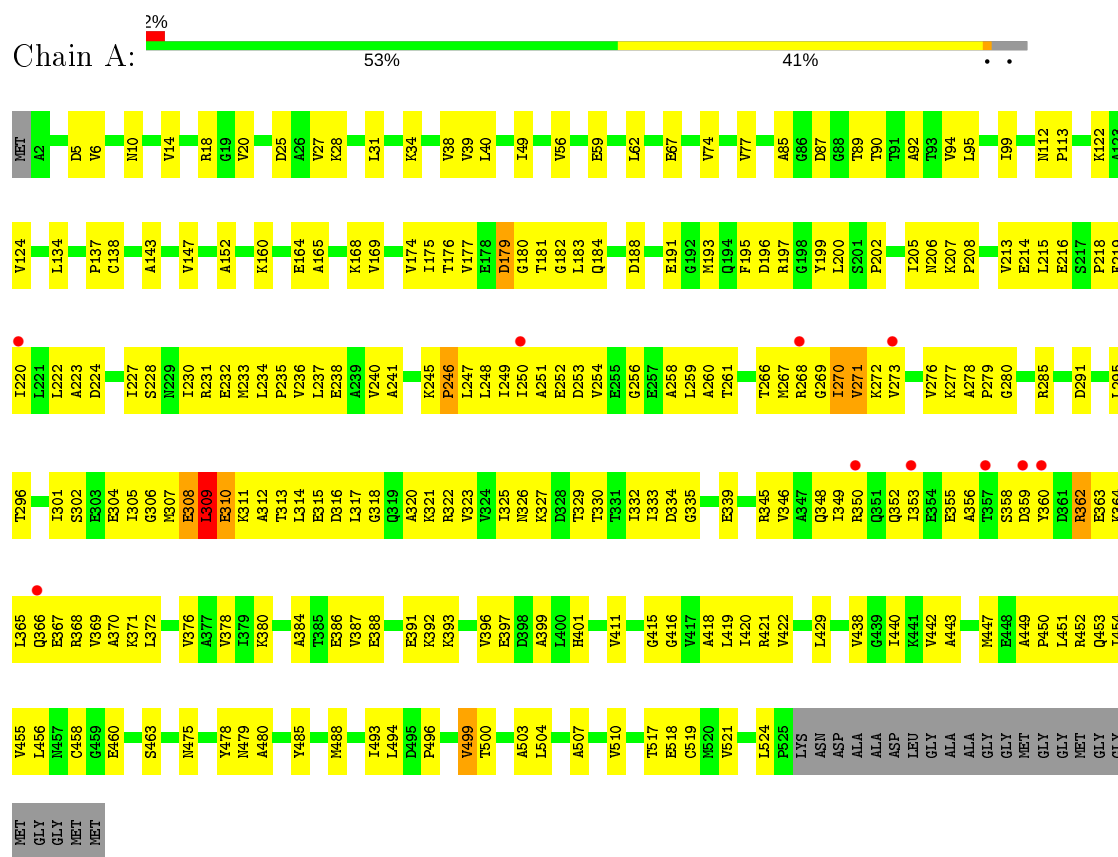
Continued from previous page...

Mol	Chain	Residues	Atoms		ZeroOcc	AltConf
6	D	1	Total 1	K 1	0	0
6	K	1	Total 1	K 1	0	0
6	E	1	Total 1	K 1	0	0
6	H	1	Total 1	K 1	0	0
6	B	1	Total 1	K 1	0	0
6	I	1	Total 1	K 1	0	0
6	C	1	Total 1	K 1	0	0
6	A	1	Total 1	K 1	0	0
6	N	1	Total 1	K 1	0	0
6	L	1	Total 1	K 1	0	0
6	F	1	Total 1	K 1	0	0
6	M	1	Total 1	K 1	0	0

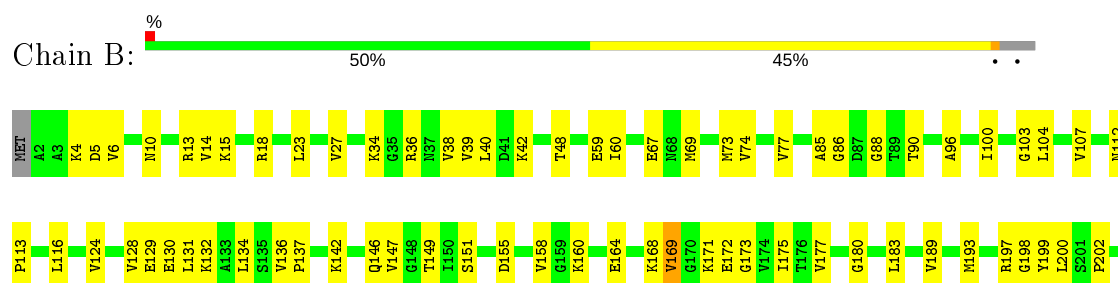
3 Residue-property plots [i](#)

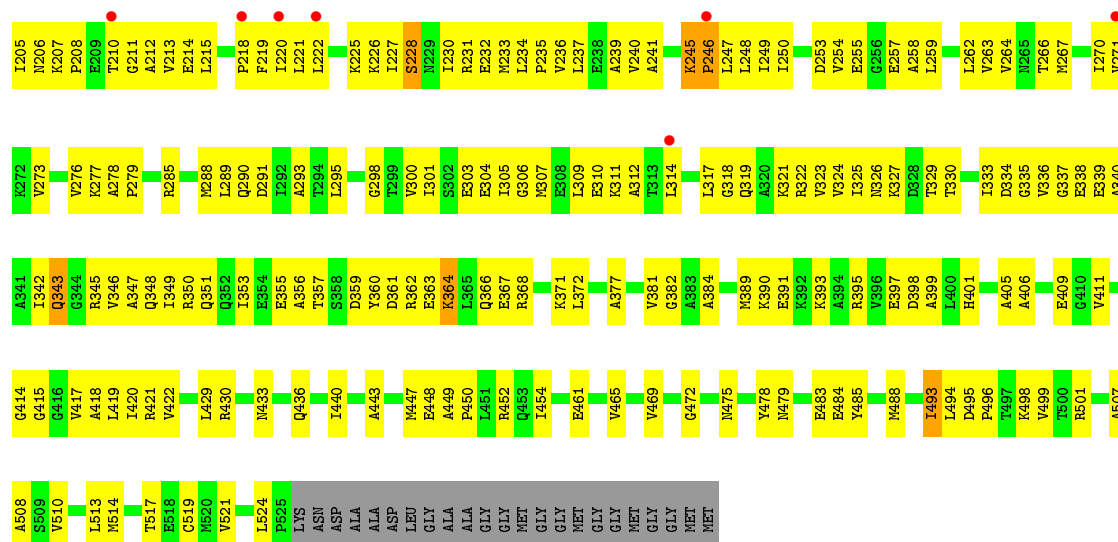
These plots are drawn for all protein, RNA and DNA chains in the entry. The first graphic for a chain summarises the proportions of the various outlier classes displayed in the second graphic. The second graphic shows the sequence view annotated by issues in geometry and electron density. Residues are 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 dot above a residue indicates a poor fit to the electron density ($RSRZ > 2$). 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: 60 kDa chaperonin

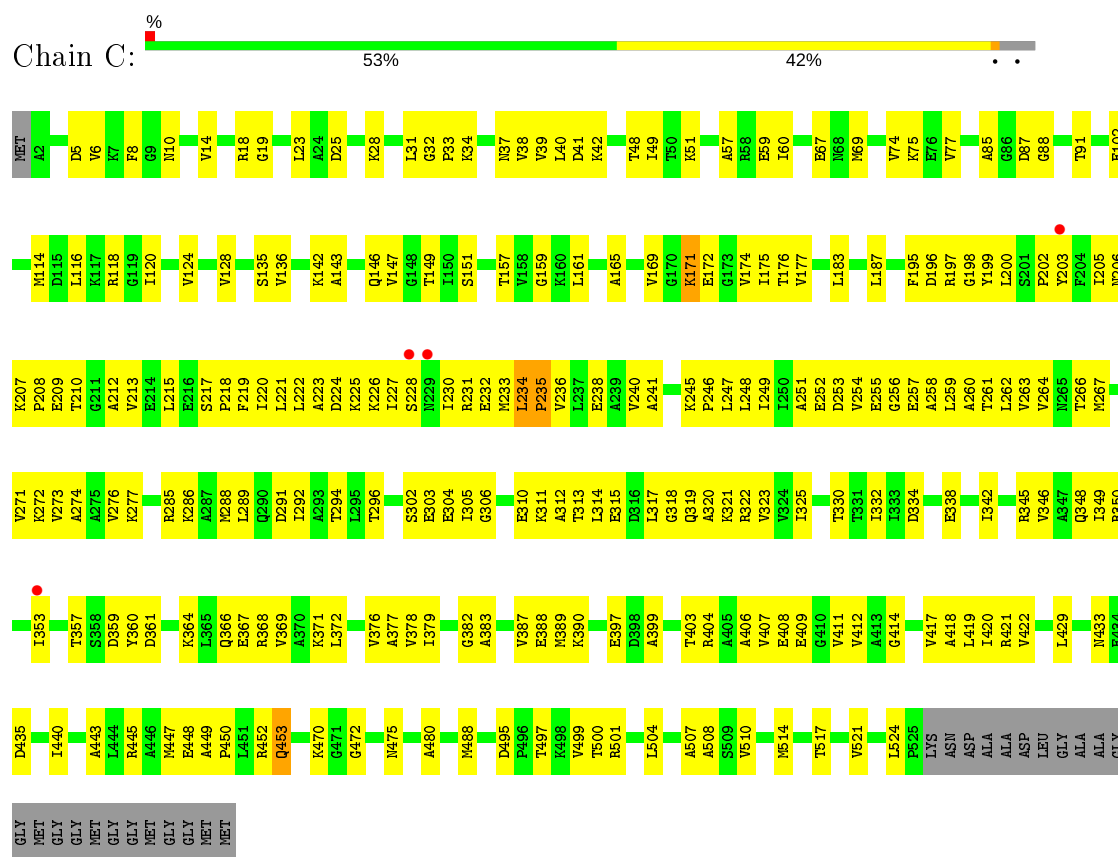


- Molecule 1: 60 kDa chaperonin

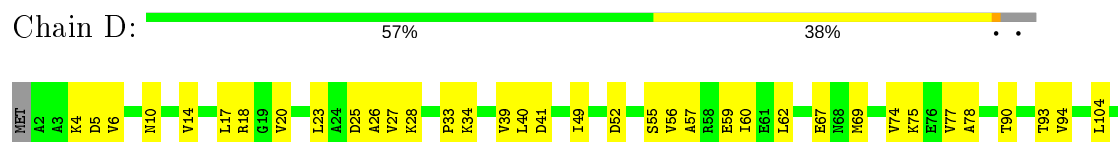


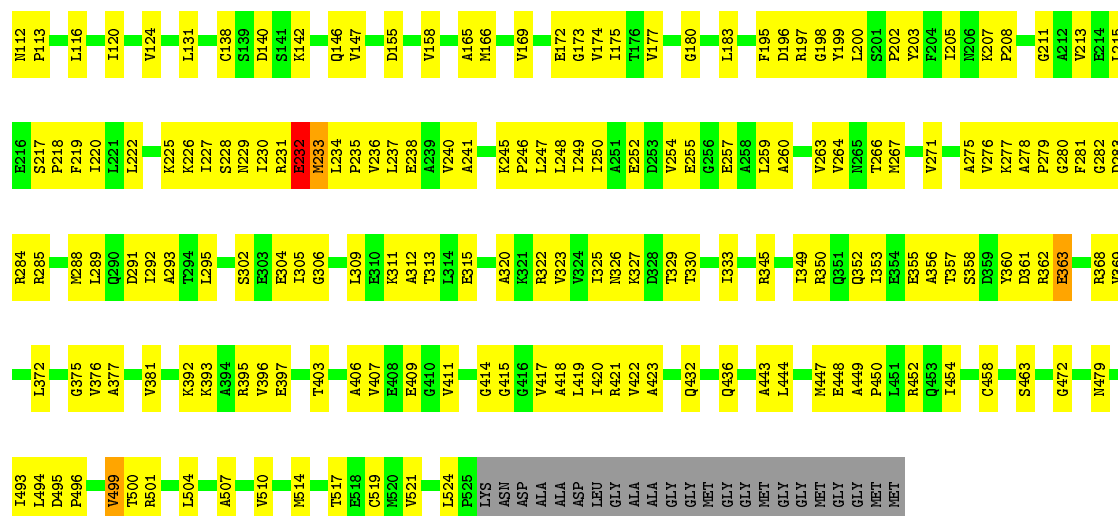


• Molecule 1: 60 kDa chaperonin

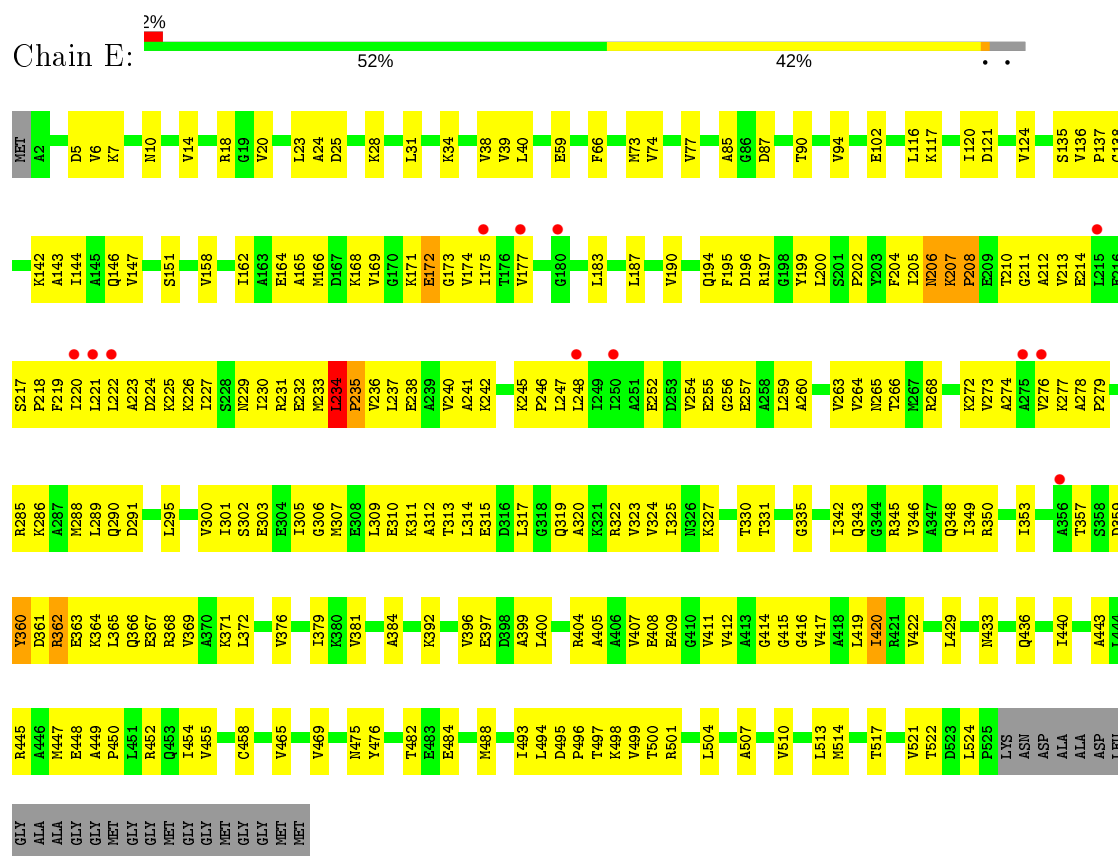


• Molecule 1: 60 kDa chaperonin

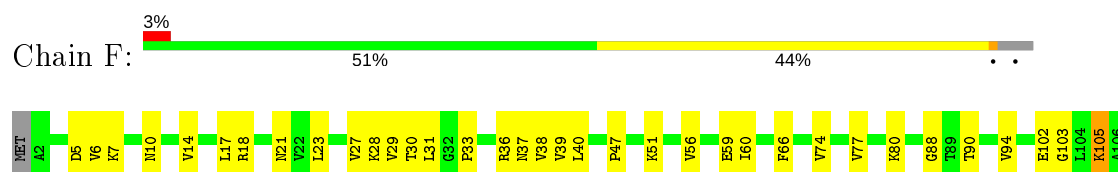


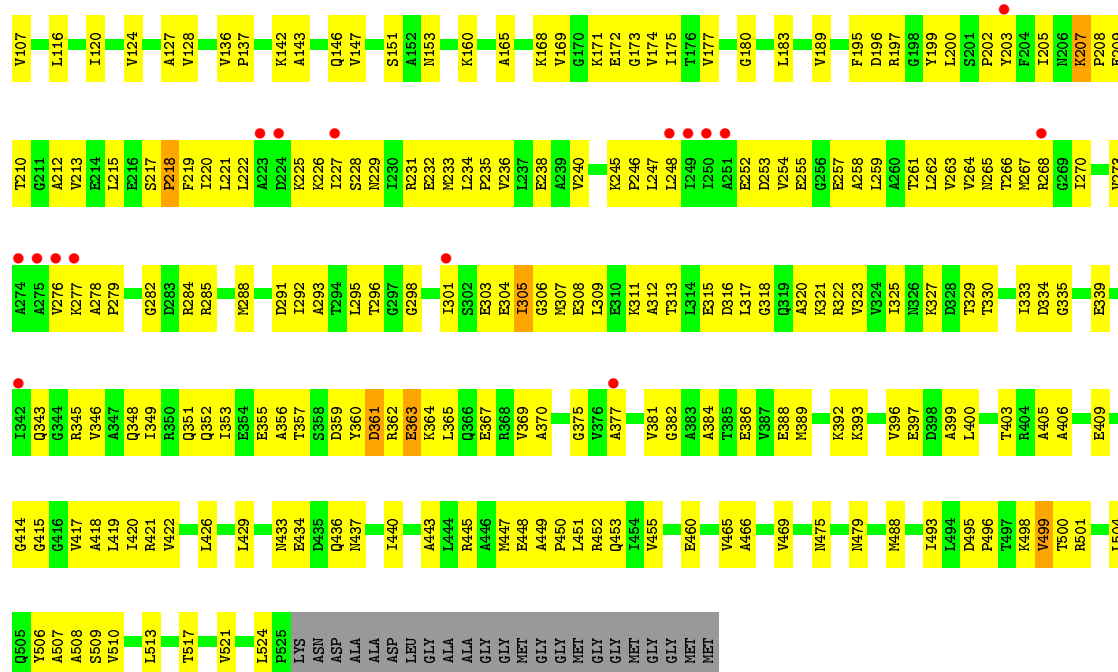


• Molecule 1: 60 kDa chaperonin

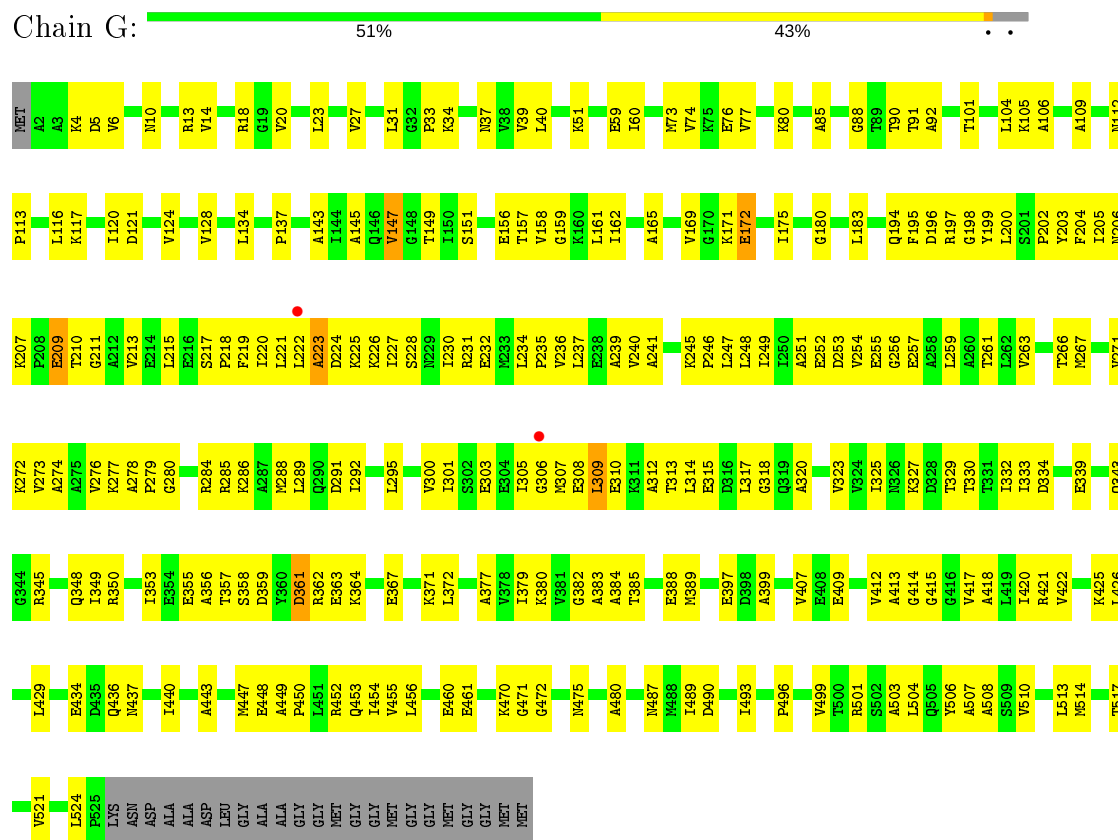


• Molecule 1: 60 kDa chaperonin



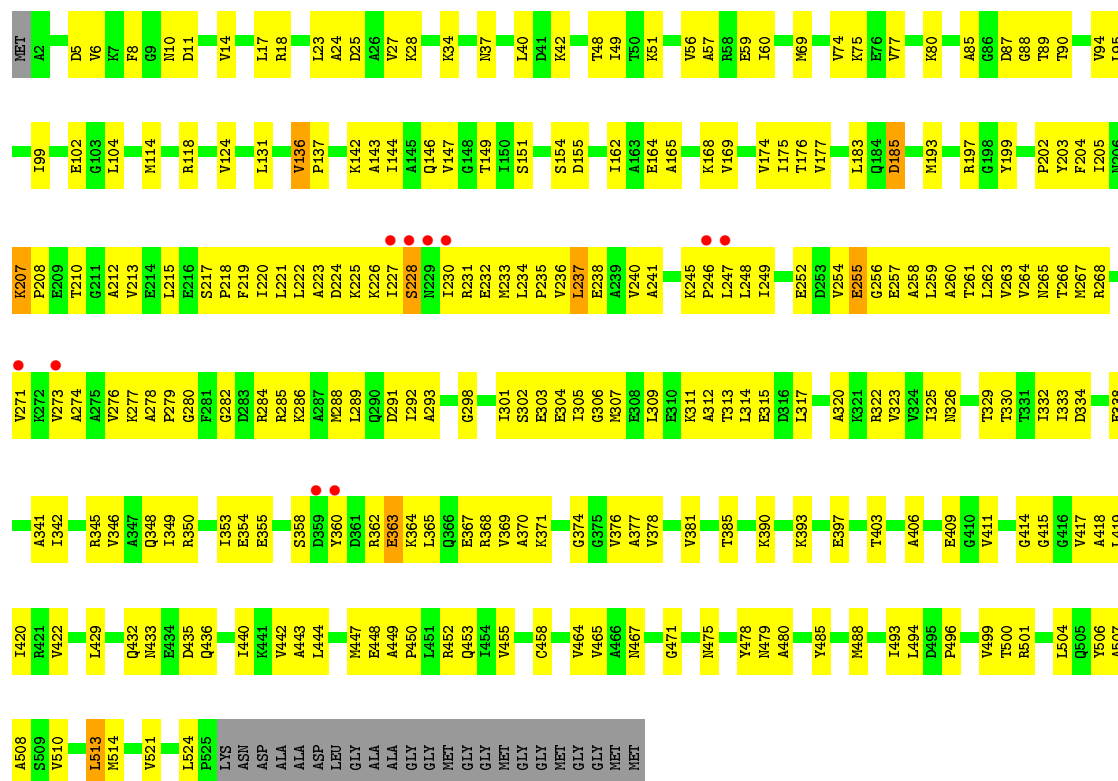


- Molecule 1: 60 kDa chaperonin



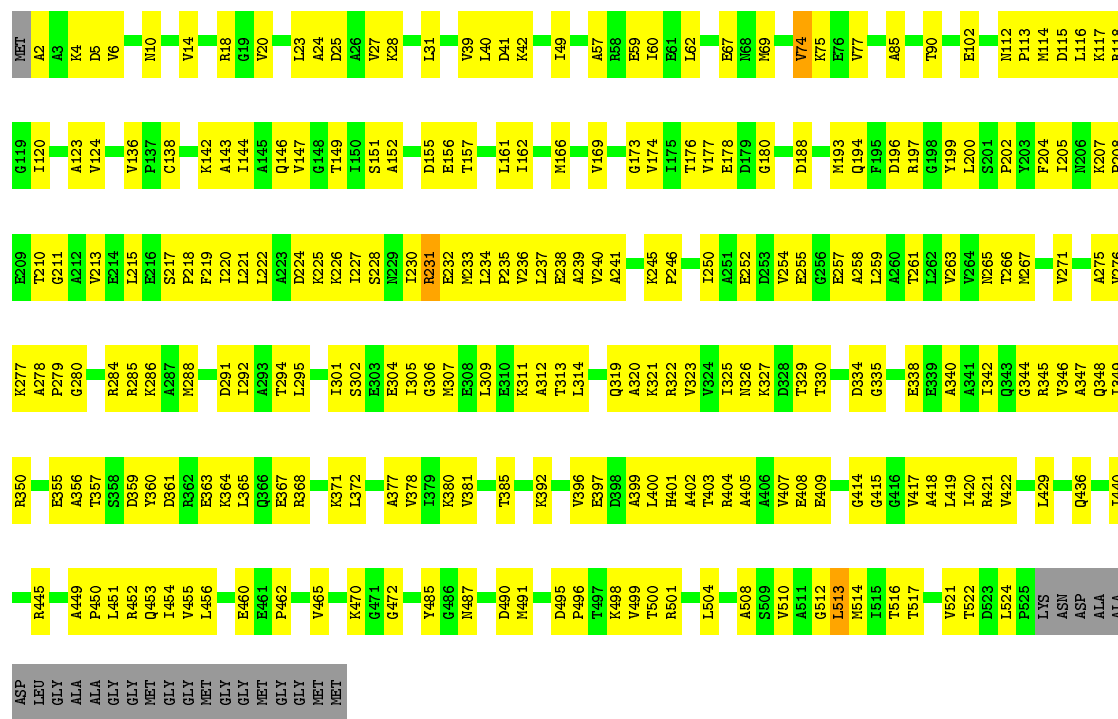
- Molecule 1: 60 kDa chaperonin



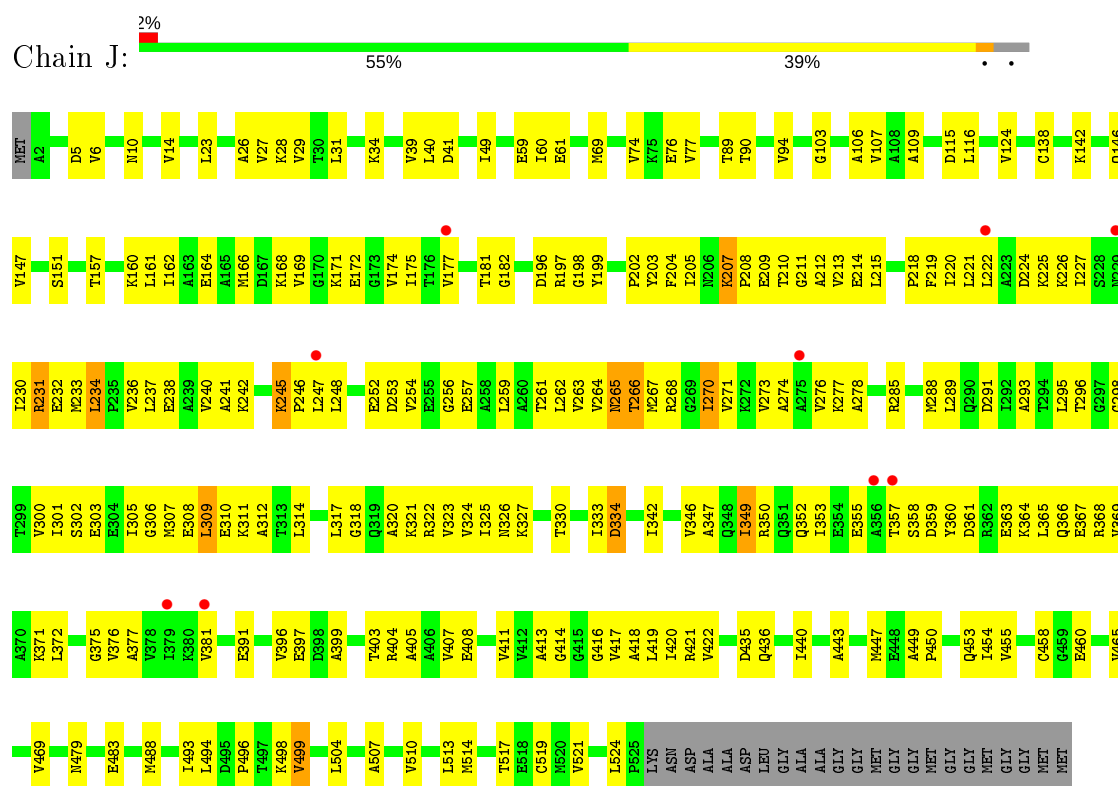


- Molecule 1: 60 kDa chaperonin

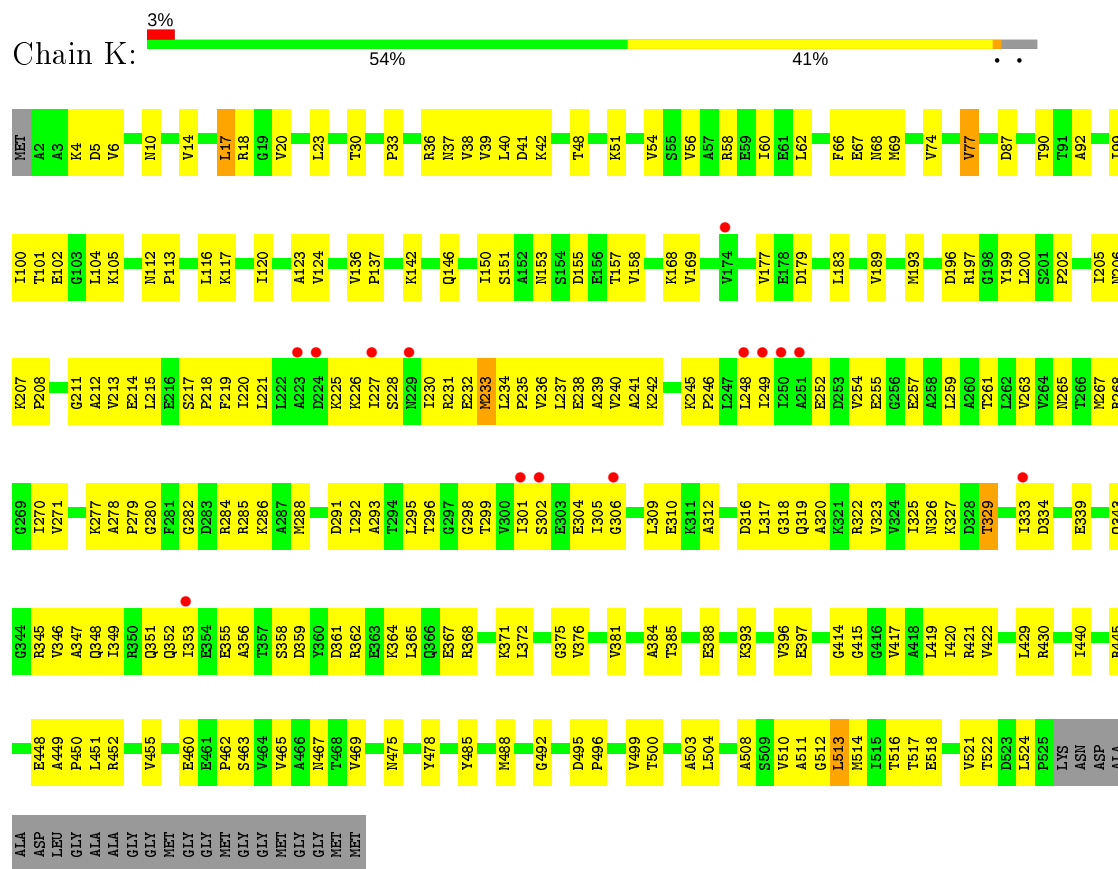
Chain I:



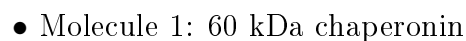
- Molecule 1: 60 kDa chaperonin



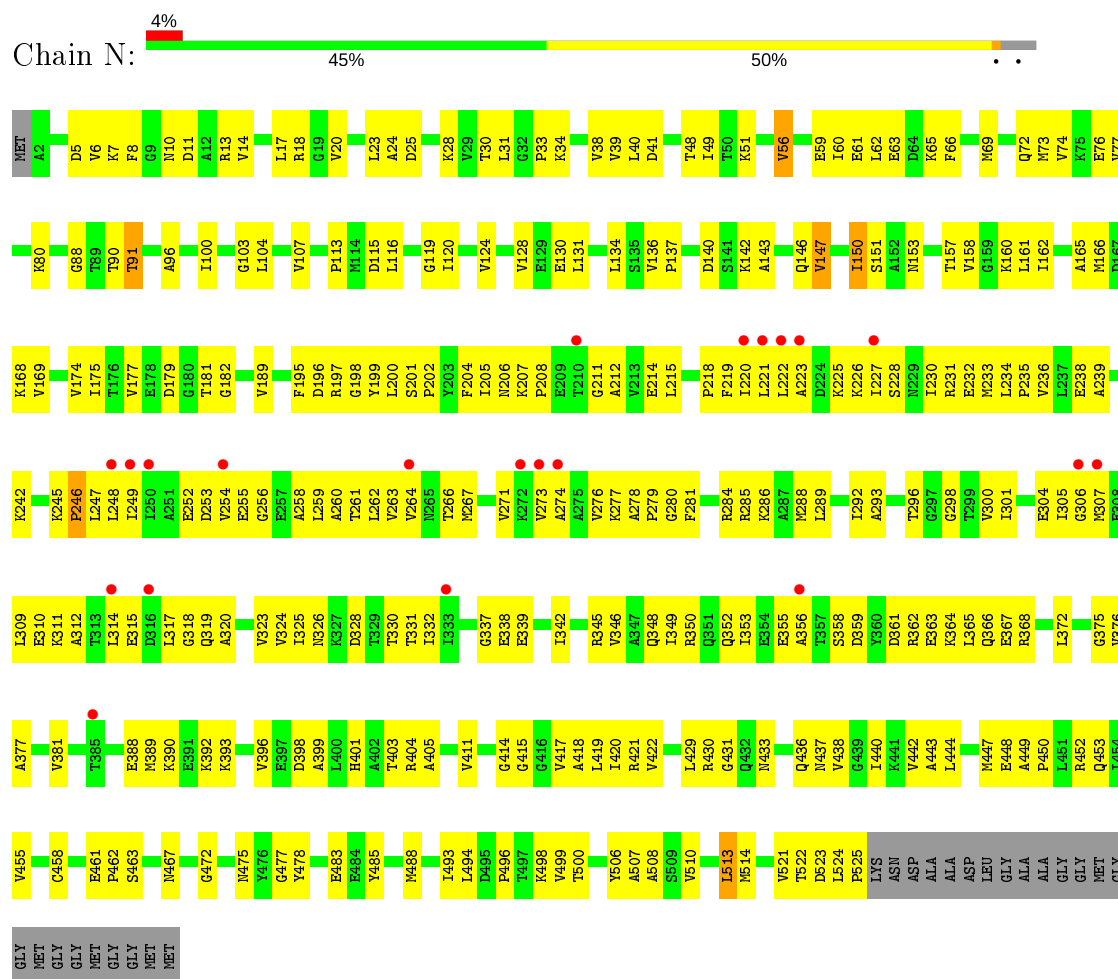
• Molecule 1: 60 kDa chaperonin



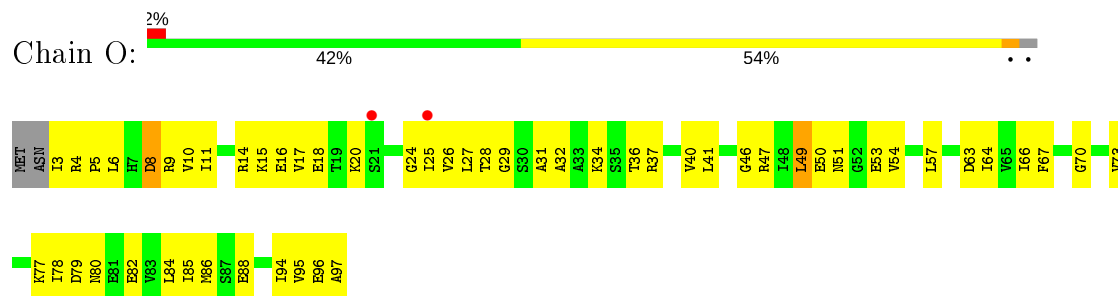
• Molecule 1: 60 kDa chaperonin



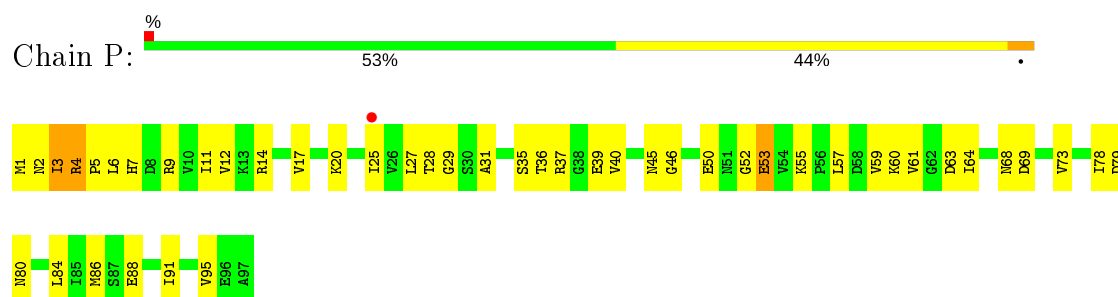
- Molecule 1: 60 kDa chaperonin



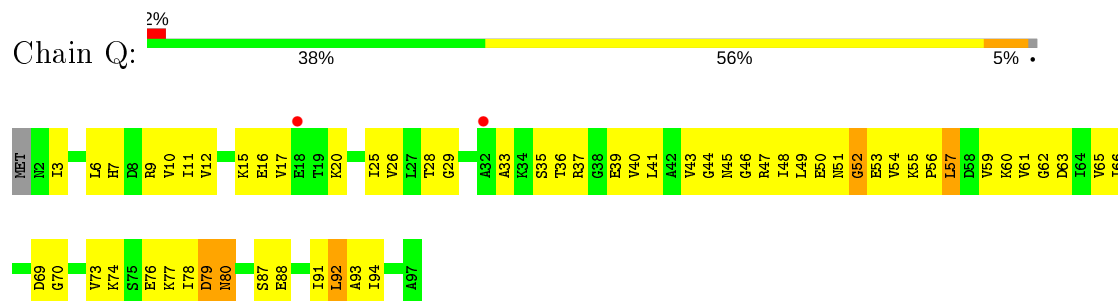
- Molecule 2: 10 kDa chaperonin



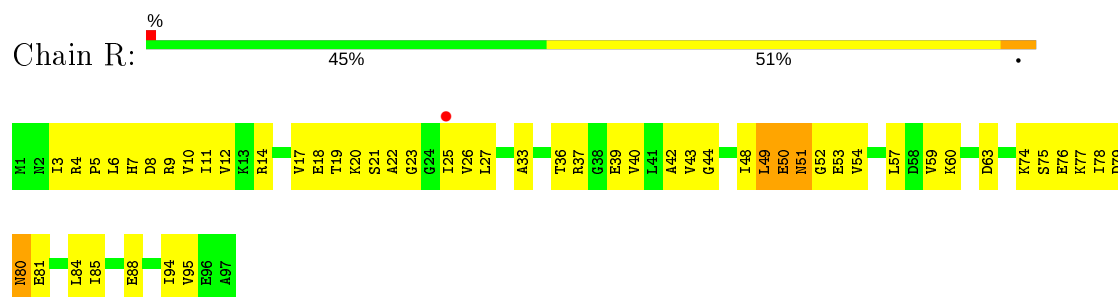
- Molecule 2: 10 kDa chaperonin



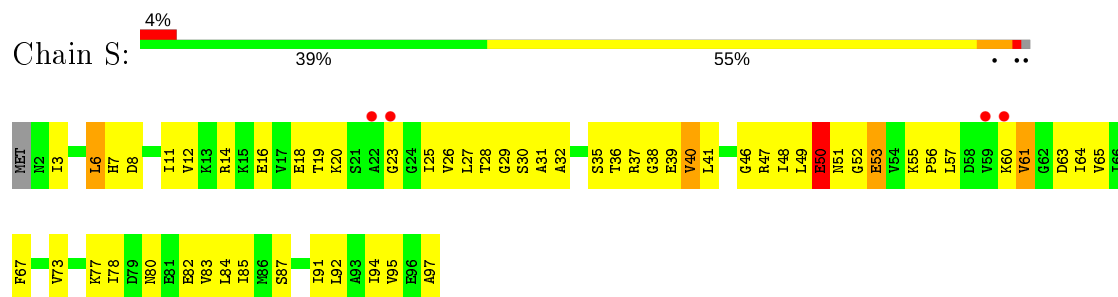
- Molecule 2: 10 kDa chaperonin



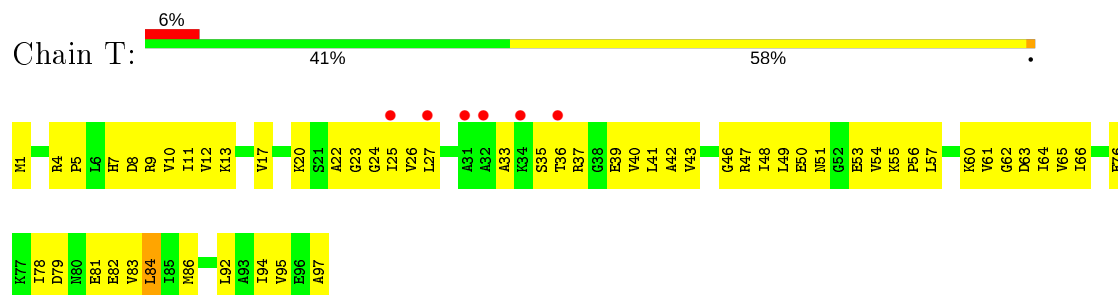
- Molecule 2: 10 kDa chaperonin



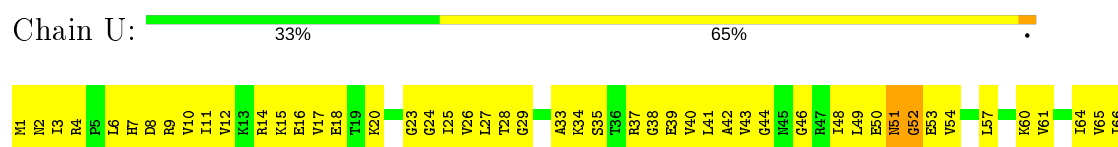
- Molecule 2: 10 kDa chaperonin

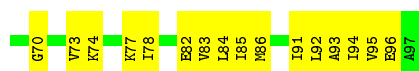


- Molecule 2: 10 kDa chaperonin

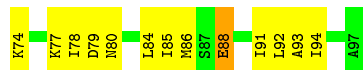
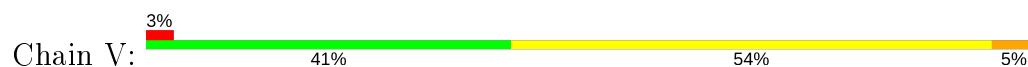


- Molecule 2: 10 kDa chaperonin





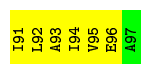
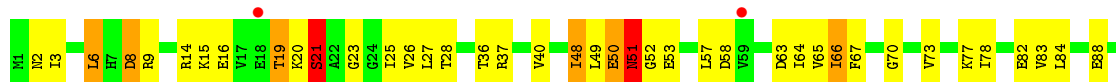
- Molecule 2: 10 kDa chaperonin



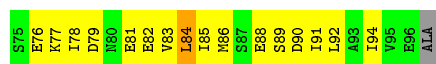
- Molecule 2: 10 kDa chaperonin



- Molecule 2: 10 kDa chaperonin

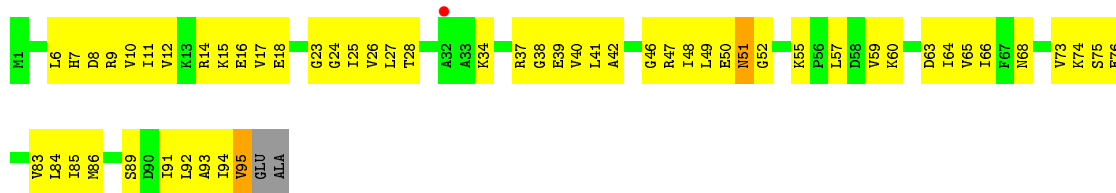


- Molecule 2: 10 kDa chaperonin

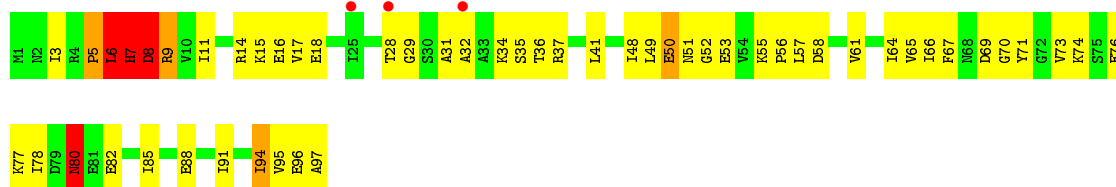


- Molecule 2: 10 kDa chaperonin





- Molecule 2: 10 kDa chaperonin



- Molecule 2: 10 kDa chaperonin



4 Data and refinement statistics

Property	Value	Source
Space group	P 21 21 21	Depositor
Cell constants a, b, c, α , β , γ	169.79Å 174.49Å 410.16Å 90.00° 90.00° 90.00°	Depositor
Resolution (Å)	90.90 – 3.84 90.90 – 3.84	Depositor EDS
% Data completeness (in resolution range)	99.9 (90.90-3.84) 99.9 (90.90-3.84)	Depositor EDS
R_{merge}	(Not available)	Depositor
R_{sym}	(Not available)	Depositor
$\langle I/\sigma(I) \rangle$ ¹	2.72 (at 3.89Å)	Xtriage
Refinement program	PHENIX (phenix.refine: 1.8.4_1496)	Depositor
R, R_{free}	0.183 , 0.248 0.187 , 0.252	Depositor DCC
R_{free} test set	1271 reflections (1.09%)	wwPDB-VP
Wilson B-factor (Å ²)	114.0	Xtriage
Anisotropy	0.331	Xtriage
Bulk solvent k_{sol} (e/Å ³), B_{sol} (Å ²)	0.27 , 110.2	EDS
L-test for twinning ²	$\langle L \rangle = 0.45$, $\langle L^2 \rangle = 0.27$	Xtriage
Estimated twinning fraction	0.118 for k,h,-l	Xtriage
F_o, F_c correlation	0.94	EDS
Total number of atoms	64559	wwPDB-VP
Average B, all atoms (Å ²)	160.0	wwPDB-VP

Xtriage's analysis on translational NCS is as follows: *The largest off-origin peak in the Patterson function is 2.24% of the height of the origin peak. No significant pseudotranslation is detected.*

¹Intensities estimated from amplitudes.

²Theoretical values of $\langle |L| \rangle$, $\langle L^2 \rangle$ for acentric reflections are 0.5, 0.333 respectively for untwinned datasets, and 0.375, 0.2 for perfectly twinned datasets.

5 Model quality ⓘ

5.1 Standard geometry ⓘ

Bond lengths and bond angles in the following residue types are not validated in this section: K, BEF, MG, ADP

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

Mol	Chain	Bond lengths		Bond angles	
		RMSZ	# $ Z > 5$	RMSZ	# $ Z > 5$
1	A	0.34	2/3883 (0.1%)	0.58	6/5243 (0.1%)
1	B	0.29	1/3883 (0.0%)	0.54	4/5243 (0.1%)
1	C	0.27	2/3883 (0.1%)	0.52	4/5243 (0.1%)
1	D	0.26	0/3883	0.50	1/5243 (0.0%)
1	E	0.31	2/3883 (0.1%)	0.59	11/5243 (0.2%)
1	F	0.25	0/3883	0.54	2/5243 (0.0%)
1	G	0.27	1/3883 (0.0%)	0.54	3/5243 (0.1%)
1	H	0.24	0/3883	0.48	0/5243
1	I	0.25	0/3883	0.51	0/5243
1	J	0.27	0/3883	0.56	4/5243 (0.1%)
1	K	0.25	1/3883 (0.0%)	0.51	2/5243 (0.0%)
1	L	0.25	0/3883	0.57	1/5243 (0.0%)
1	M	0.29	2/3883 (0.1%)	0.53	2/5243 (0.0%)
1	N	0.25	1/3883 (0.0%)	0.43	0/5243
2	1	0.35	1/731 (0.1%)	0.83	6/983 (0.6%)
2	2	0.29	0/731	0.53	0/983
2	O	0.26	0/715	0.54	0/962
2	P	0.28	0/731	0.64	1/983 (0.1%)
2	Q	0.25	0/723	0.57	0/973
2	R	0.26	0/731	0.54	1/983 (0.1%)
2	S	0.29	0/723	0.53	0/973
2	T	0.22	0/731	0.43	0/983
2	U	0.29	0/731	0.50	0/983
2	V	0.27	0/731	0.65	1/983 (0.1%)
2	W	0.28	0/731	0.64	2/983 (0.2%)
2	X	0.40	1/731 (0.1%)	0.99	9/983 (0.9%)
2	Y	0.26	0/726	0.59	1/976 (0.1%)
2	Z	0.25	0/717	0.47	0/964
All	All	0.27	14/64545 (0.0%)	0.55	61/87097 (0.1%)

Chiral center outliers are detected by calculating the chiral volume of a chiral center and verifying if

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

Mol	Chain	#Chirality outliers	#Planarity outliers
1	A	0	1
1	D	0	1
2	1	0	2
2	S	0	1
All	All	0	5

All (14) bond length outliers are listed below:

Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
1	A	246	PRO	N-CD	-10.73	1.32	1.47
1	E	208	PRO	N-CD	-9.79	1.34	1.47
1	B	246	PRO	N-CD	-9.12	1.35	1.47
1	A	137	PRO	N-CD	7.50	1.58	1.47
1	M	137	PRO	N-CD	6.88	1.57	1.47
1	M	246	PRO	N-CD	-6.41	1.38	1.47
1	E	171	LYS	CA-C	6.31	1.69	1.52
1	G	137	PRO	N-CD	5.92	1.56	1.47
1	C	171	LYS	CA-C	5.90	1.68	1.52
2	1	8	ASP	CA-C	5.61	1.67	1.52
2	X	51	ASN	CA-C	5.50	1.67	1.52
1	K	233	MET	CA-C	5.41	1.67	1.52
1	N	246	PRO	N-CD	-5.34	1.40	1.47
1	C	235	PRO	N-CD	-5.08	1.40	1.47

All (61) bond angle outliers are listed below:

Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
2	X	21	SER	N-CA-C	13.14	146.49	111.00
2	X	21	SER	N-CA-CB	-12.79	91.32	110.50
2	1	52	GLY	N-CA-C	11.15	140.97	113.10
2	V	93	ALA	N-CA-CB	-9.18	97.24	110.10
1	B	246	PRO	N-CA-CB	-9.11	92.36	103.30
1	J	266	THR	N-CA-CB	8.08	125.65	110.30
1	A	310	GLU	N-CA-C	7.78	132.00	111.00
1	A	310	GLU	N-CA-CB	-7.71	96.73	110.60
1	F	362	ARG	N-CA-CB	7.45	124.02	110.60
1	J	265	ASN	CB-CA-C	-7.40	95.60	110.40
1	B	246	PRO	N-CA-C	7.28	131.03	112.10
1	E	172	GLU	N-CA-C	7.02	129.96	111.00

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
1	F	361	ASP	CB-CA-C	-6.80	96.80	110.40
1	E	208	PRO	N-CD-CG	-6.60	93.30	103.20
1	A	246	PRO	CA-N-CD	6.51	120.81	111.70
2	R	51	ASN	N-CA-CB	6.51	122.31	110.60
1	E	208	PRO	N-CA-C	6.43	128.82	112.10
1	E	208	PRO	N-CA-CB	-6.38	95.58	102.60
1	B	364	LYS	N-CA-CB	-6.35	99.17	110.60
2	1	7	HIS	CB-CA-C	-6.27	97.86	110.40
2	W	71	TYR	CB-CA-C	-6.24	97.91	110.40
1	C	172	GLU	N-CA-C	6.23	127.82	111.00
2	P	79	ASP	CB-CA-C	6.21	122.83	110.40
2	X	8	ASP	CA-C-N	-6.20	103.56	117.20
2	1	80	ASN	N-CA-C	6.17	127.67	111.00
2	X	9	ARG	N-CA-C	6.11	127.51	111.00
1	J	210	THR	N-CA-CB	6.10	121.90	110.30
1	E	208	PRO	CA-N-CD	6.03	120.14	111.70
2	X	19	THR	CB-CA-C	5.94	127.64	111.60
1	M	310	GLU	N-CA-CB	-5.91	99.96	110.60
1	G	172	GLU	N-CA-C	5.89	126.91	111.00
2	X	50	GLU	CB-CA-C	-5.79	98.81	110.40
1	A	309	LEU	N-CA-C	5.75	126.53	111.00
2	X	51	ASN	CA-C-O	5.74	132.16	120.10
1	A	180	GLY	N-CA-C	5.71	127.39	113.10
1	A	246	PRO	N-CA-CB	-5.71	96.32	102.60
2	Y	84	LEU	N-CA-C	5.71	126.40	111.00
1	C	234	LEU	CA-CB-CG	5.70	128.41	115.30
1	E	360	TYR	N-CA-C	5.67	126.30	111.00
1	E	171	LYS	CA-C-N	-5.64	104.80	117.20
2	X	51	ASN	CA-C-N	-5.63	104.94	116.20
2	X	8	ASP	CA-C-O	5.62	131.90	120.10
2	1	8	ASP	CA-C-O	5.55	131.75	120.10
1	B	246	PRO	CA-N-CD	5.54	119.46	111.70
1	M	310	GLU	N-CA-C	5.48	125.79	111.00
1	L	217	SER	N-CA-C	5.47	125.78	111.00
2	1	9	ARG	N-CA-C	5.43	125.66	111.00
1	E	206	ASN	CB-CA-C	-5.38	99.64	110.40
1	D	232	GLU	N-CA-CB	5.33	120.20	110.60
1	E	171	LYS	CA-C-O	5.31	131.25	120.10
1	G	172	GLU	N-CA-CB	-5.30	101.06	110.60
1	E	172	GLU	N-CA-CB	-5.28	101.10	110.60
1	K	233	MET	CA-C-N	-5.27	105.60	117.20
1	C	171	LYS	CA-C-N	-5.24	105.68	117.20

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
1	E	362	ARG	N-CA-CB	5.21	119.97	110.60
2	1	8	ASP	CA-C-N	-5.17	105.83	117.20
1	G	382	GLY	N-CA-C	5.15	125.98	113.10
1	C	171	LYS	CA-C-O	5.11	130.83	120.10
1	K	233	MET	CA-C-O	5.11	130.84	120.10
2	W	72	GLY	N-CA-C	-5.11	100.33	113.10
1	J	265	ASN	N-CA-C	5.02	124.55	111.00

There are no chirality outliers.

All (5) planarity outliers are listed below:

Mol	Chain	Res	Type	Group
2	1	50	GLU	Peptide
2	1	6	LEU	Peptide
1	A	179	ASP	Peptide
1	D	232	GLU	Peptide
2	S	50	GLU	Peptide

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	A	3855	0	3974	266	0
1	B	3855	0	3975	282	5
1	C	3855	0	3974	260	0
1	D	3855	0	3975	221	0
1	E	3855	0	3976	268	0
1	F	3855	0	3975	257	0
1	G	3855	0	3975	288	2
1	H	3855	0	3975	286	0
1	I	3855	0	3976	251	0
1	J	3855	0	3976	242	0
1	K	3855	0	3976	235	12
1	L	3855	0	3975	272	1
1	M	3855	0	3974	254	7
1	N	3855	0	3976	317	0
2	1	727	0	762	69	1

Continued on next page...

Continued from previous page...

Mol	Chain	Non-H	H(model)	H(added)	Clashes	Symm-Clashes
2	2	727	0	762	74	1
2	O	711	0	744	85	0
2	P	727	0	762	71	2
2	Q	719	0	750	73	0
2	R	727	0	762	102	0
2	S	719	0	750	87	0
2	T	727	0	762	78	1
2	U	727	0	762	105	0
2	V	727	0	762	89	0
2	W	727	0	762	78	0
2	X	727	0	762	66	0
2	Y	722	0	757	85	0
2	Z	713	0	751	87	0
3	A	27	0	12	1	0
3	B	27	0	12	5	0
3	C	27	0	12	4	0
3	D	27	0	12	4	0
3	E	27	0	12	4	0
3	F	27	0	12	4	0
3	G	27	0	12	5	0
3	H	27	0	12	2	0
3	I	27	0	12	3	0
3	J	27	0	12	1	0
3	K	27	0	12	4	0
3	L	27	0	12	2	0
3	M	27	0	12	4	0
3	N	27	0	12	5	0
4	A	4	0	0	2	0
4	B	4	0	0	3	0
4	C	4	0	0	4	0
4	D	4	0	0	1	0
4	E	4	0	0	1	0
4	F	4	0	0	1	0
4	G	4	0	0	1	0
4	H	4	0	0	4	0
4	I	4	0	0	4	0
4	J	4	0	0	2	0
4	K	4	0	0	2	0
4	L	4	0	0	2	0
4	M	4	0	0	1	0
4	N	4	0	0	3	0
5	A	1	0	0	0	0

Continued on next page...

Continued from previous page...

Mol	Chain	Non-H	H(model)	H(added)	Clashes	Symm-Clashes
5	B	1	0	0	0	0
5	C	1	0	0	0	0
5	D	1	0	0	0	0
5	E	1	0	0	0	0
5	F	1	0	0	0	0
5	G	1	0	0	0	0
5	H	1	0	0	0	0
5	I	1	0	0	0	0
5	J	1	0	0	0	0
5	K	1	0	0	0	0
5	L	1	0	0	0	0
5	M	1	0	0	0	0
5	N	1	0	0	0	0
6	A	1	0	0	0	0
6	B	1	0	0	0	0
6	C	1	0	0	0	0
6	D	1	0	0	0	0
6	E	1	0	0	0	0
6	F	1	0	0	0	0
6	G	1	0	0	0	0
6	H	1	0	0	0	0
6	I	1	0	0	0	0
6	J	1	0	0	0	0
6	K	1	0	0	0	0
6	L	1	0	0	0	0
6	M	1	0	0	0	0
6	N	1	0	0	0	0
All	All	64559	0	66430	4540	16

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 35.

All (4540) 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)
2:O:95:VAL:HA	2:P:3:ILE:CG2	1.26	1.56
1:K:207:LYS:CG	1:K:208:PRO:HD3	1.46	1.42
2:O:95:VAL:CA	2:P:3:ILE:HG22	1.52	1.37
1:F:218:PRO:O	1:F:219:PHE:CD1	1.81	1.33
2:V:56:PRO:O	2:V:57:LEU:HG	1.26	1.32
2:R:6:LEU:HD21	2:R:7:HIS:CE1	1.65	1.30

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:R:49:LEU:HD12	2:S:50:GLU:O	1.28	1.26
1:B:363:GLU:OE1	1:B:364:LYS:HG3	1.34	1.26
1:L:208:PRO:O	1:L:209:GLU:CD	1.75	1.25
1:G:412:VAL:HG11	1:G:418:ALA:CB	1.64	1.24
2:U:48:ILE:O	2:U:49:LEU:HD12	1.12	1.24
2:O:96:GLU:O	2:P:1:MET:HA	1.22	1.23
2:Z:8:ASP:O	2:Z:57:LEU:HD21	1.39	1.22
1:G:412:VAL:CG1	1:G:418:ALA:HB2	1.70	1.20
2:R:6:LEU:CD2	2:R:7:HIS:ND1	2.06	1.19
1:G:205:ILE:HA	1:G:213:VAL:HG22	1.21	1.18
2:R:49:LEU:HD12	2:S:50:GLU:C	1.62	1.18
2:R:6:LEU:HD21	2:R:7:HIS:ND1	1.58	1.17
1:L:208:PRO:C	1:L:209:GLU:OE1	1.83	1.15
1:B:245:LYS:HB2	1:B:246:PRO:HD3	1.24	1.15
1:F:218:PRO:HB3	1:F:246:PRO:HG2	1.28	1.14
2:O:96:GLU:O	2:P:1:MET:CA	1.95	1.14
1:D:360:TYR:HA	1:D:363:GLU:HG3	1.27	1.14
1:G:412:VAL:HG11	1:G:418:ALA:HB1	1.26	1.13
2:T:20:LYS:HE2	2:T:22:ALA:HB3	1.23	1.12
1:F:233:MET:HG3	1:F:234:LEU:HD23	1.25	1.12
2:Y:12:VAL:HG12	2:Y:40:VAL:HA	1.32	1.12
1:C:197:ARG:HG2	1:C:277:LYS:HB3	1.29	1.12
1:K:207:LYS:HG2	1:K:208:PRO:HD3	1.29	1.11
2:U:48:ILE:O	2:U:49:LEU:CD1	1.98	1.11
1:C:6:VAL:HG12	1:C:521:VAL:HG22	1.33	1.11
1:K:207:LYS:HG3	1:K:208:PRO:CD	1.81	1.11
1:M:241:ALA:HA	1:M:271:VAL:HG11	1.30	1.11
2:W:20:LYS:HA	2:W:21:SER:HB3	1.14	1.11
2:O:95:VAL:CA	2:P:3:ILE:CG2	2.19	1.10
1:E:207:LYS:HB2	1:E:208:PRO:HD3	1.16	1.09
1:K:6:VAL:HG12	1:K:521:VAL:HG22	1.34	1.08
2:R:20:LYS:HA	2:R:21:SER:HB3	1.22	1.08
1:E:301:ILE:HG12	1:E:307:MET:HE1	1.18	1.08
1:H:262:LEU:HD11	1:H:273:VAL:HG11	1.30	1.08
1:C:177:VAL:HG11	1:C:397:GLU:HG2	1.30	1.08
2:U:51:ASN:O	2:U:53:GLU:HG2	1.54	1.08
1:K:207:LYS:HG3	1:K:208:PRO:HD3	1.10	1.07
1:N:77:VAL:HG21	1:N:507:ALA:HA	1.32	1.06
1:L:245:LYS:HB3	1:L:246:PRO:HD2	1.36	1.06
1:H:302:SER:HB2	1:H:305:ILE:HD13	1.35	1.06
1:H:238:GLU:HA	2:V:25:ILE:HD11	1.38	1.06

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:S:49:LEU:CD2	2:T:48:ILE:HD12	1.85	1.05
2:P:64:ILE:HG23	2:P:95:VAL:HB	1.38	1.05
1:F:6:VAL:HG12	1:F:521:VAL:HG22	1.35	1.05
1:H:197:ARG:HG2	1:H:277:LYS:HB3	1.38	1.05
1:J:205:ILE:HA	1:J:213:VAL:HG22	1.34	1.04
1:C:208:PRO:HB2	1:C:212:ALA:HB3	1.39	1.04
1:L:412:VAL:CG1	1:L:418:ALA:HB2	1.88	1.03
2:S:49:LEU:HD22	2:T:48:ILE:HD12	1.11	1.03
1:G:412:VAL:CG1	1:G:418:ALA:CB	2.33	1.03
2:S:8:ASP:HB3	2:S:47:ARG:HB3	1.38	1.03
1:A:236:VAL:HG21	1:A:312:ALA:HB3	1.40	1.03
1:G:412:VAL:HG12	1:G:418:ALA:HB2	1.36	1.02
2:R:50:GLU:N	2:R:50:GLU:OE1	1.90	1.02
1:D:232:GLU:HG2	1:D:309:LEU:HD23	1.40	1.02
1:E:205:ILE:HA	1:E:213:VAL:HG22	1.41	1.02
1:N:228:SER:HB3	1:N:255:GLU:HB2	1.38	1.02
1:H:221:LEU:HD13	1:H:317:LEU:HD21	1.37	1.02
1:L:412:VAL:HG11	1:L:418:ALA:CB	1.89	1.01
2:1:37:ARG:HH22	2:2:78:ILE:HG23	1.23	1.01
2:1:49:LEU:HD12	2:1:50:GLU:O	1.61	1.01
1:L:412:VAL:HG11	1:L:418:ALA:HB1	1.43	1.01
2:S:49:LEU:HD22	2:T:48:ILE:CD1	1.89	1.01
1:E:207:LYS:CB	1:E:208:PRO:HD3	1.81	1.00
1:L:230:ILE:HG21	1:L:309:LEU:HD21	1.43	1.00
2:T:12:VAL:HG12	2:T:40:VAL:HG23	1.44	1.00
1:L:367:GLU:HG2	1:L:371:LYS:HE2	1.43	1.00
2:2:49:LEU:H	2:2:52:GLY:HA2	1.26	1.00
1:H:233:MET:HE1	1:H:237:LEU:HD12	1.44	1.00
2:Z:95:VAL:HA	2:1:3:ILE:HG22	1.44	1.00
2:S:67:PHE:HA	2:S:92:LEU:HD13	1.40	1.00
2:T:97:ALA:HB2	2:U:1:MET:HA	1.43	1.00
1:L:412:VAL:HG12	1:L:418:ALA:HB2	1.40	1.00
1:B:222:LEU:HD22	1:B:300:VAL:HG22	1.44	0.99
1:M:291:ASP:HB3	1:M:372:LEU:HD21	1.43	0.99
1:M:325:ILE:HG13	1:M:330:THR:HG23	1.42	0.99
2:2:65:VAL:HG12	2:2:94:ILE:HG22	1.42	0.99
1:A:309:LEU:HD12	1:A:310:GLU:H	1.26	0.99
1:L:208:PRO:O	1:L:209:GLU:OE1	1.78	0.99
1:C:236:VAL:HG21	1:C:312:ALA:HB3	1.39	0.99
2:O:8:ASP:HB2	2:O:57:LEU:HD11	1.44	0.99
2:R:36:THR:HG23	2:R:37:ARG:HG3	1.45	0.99

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:C:226:LYS:HG2	1:C:252:GLU:HG2	1.39	0.99
1:H:322:ARG:HB3	1:H:333:ILE:HD12	1.41	0.99
2:O:50:GLU:O	2:U:50:GLU:HG3	1.63	0.99
1:K:208:PRO:HB2	1:K:212:ALA:HB3	1.45	0.99
1:H:350:ARG:HD3	1:H:353:ILE:HD12	1.44	0.99
1:K:207:LYS:CG	1:K:208:PRO:CD	2.38	0.99
1:G:221:LEU:HD13	1:G:249:ILE:HD13	1.45	0.99
1:K:234:LEU:O	1:K:237:LEU:HG	1.63	0.98
1:M:177:VAL:HG11	1:M:397:GLU:HG2	1.42	0.98
2:R:3:ILE:HD11	2:R:11:ILE:HD12	1.44	0.98
1:E:234:LEU:HD23	1:E:235:PRO:HD3	1.44	0.98
1:G:230:ILE:HG21	1:G:309:LEU:HD21	1.46	0.98
1:J:265:ASN:HB2	1:J:270:ILE:HB	1.45	0.97
2:R:48:ILE:HA	2:R:54:VAL:HG12	1.46	0.97
1:F:225:LYS:HG2	1:F:309:LEU:HD21	1.46	0.97
1:E:205:ILE:HD13	1:E:211:GLY:HA2	1.47	0.97
1:H:6:VAL:HG12	1:H:521:VAL:HG22	1.44	0.97
1:N:150:ILE:HD11	1:N:493:ILE:HG22	1.44	0.96
2:X:50:GLU:O	2:X:51:ASN:CG	2.04	0.96
2:R:8:ASP:O	2:R:57:LEU:HD11	1.63	0.96
2:S:64:ILE:HG23	2:S:95:VAL:HG13	1.46	0.96
1:B:230:ILE:HG21	1:B:309:LEU:HD13	1.48	0.96
2:X:37:ARG:HG2	2:X:66:ILE:HG23	1.48	0.96
1:B:363:GLU:OE1	1:B:364:LYS:CG	2.13	0.96
1:L:208:PRO:HB2	1:L:209:GLU:OE1	1.64	0.96
2:X:65:VAL:HG12	2:X:94:ILE:HG22	1.44	0.96
2:Z:93:ALA:HB2	2:1:5:PRO:HA	1.46	0.96
1:F:199:TYR:HA	1:F:276:VAL:HG12	1.46	0.95
2:O:95:VAL:HG22	2:P:3:ILE:HG21	1.48	0.95
1:A:309:LEU:HD12	1:A:310:GLU:N	1.82	0.95
1:J:322:ARG:HB3	1:J:333:ILE:HD12	1.47	0.95
1:K:177:VAL:HG11	1:K:397:GLU:HG2	1.48	0.95
2:R:49:LEU:HD11	2:S:50:GLU:HA	1.49	0.95
1:G:172:GLU:OE1	1:G:172:GLU:N	2.00	0.94
1:G:27:VAL:HG12	1:G:90:THR:HG23	1.48	0.94
2:Z:8:ASP:OD1	2:Z:9:ARG:N	2.00	0.94
1:F:357:THR:HG21	1:F:361:ASP:OD2	1.68	0.94
1:J:231:ARG:HD2	1:J:257:GLU:OE1	1.68	0.94
1:K:284:ARG:HD3	1:K:364:LYS:HE3	1.48	0.94
1:K:302:SER:HB2	1:K:305:ILE:HD13	1.50	0.94
2:U:49:LEU:HB3	2:U:50:GLU:OE1	1.67	0.94

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:D:230:ILE:O	1:D:231:ARG:HG2	1.68	0.93
2:O:64:ILE:HG23	2:O:95:VAL:HB	1.48	0.93
2:X:67:PHE:HA	2:X:92:LEU:HD13	1.48	0.93
1:H:367:GLU:HG2	1:H:371:LYS:HE2	1.48	0.93
1:J:367:GLU:HG2	1:J:371:LYS:HE2	1.51	0.93
1:N:247:LEU:HG	1:N:249:ILE:HD11	1.50	0.93
1:B:6:VAL:HG12	1:B:521:VAL:HG22	1.50	0.93
1:A:306:GLY:HA3	1:B:264:VAL:HG21	1.51	0.93
1:G:226:LYS:HB3	1:G:253:ASP:HB2	1.51	0.93
1:N:411:VAL:HG21	1:N:494:LEU:HD23	1.50	0.93
2:R:6:LEU:HD23	2:R:7:HIS:CG	2.03	0.93
1:B:245:LYS:CB	1:B:246:PRO:HD3	1.95	0.92
2:R:48:ILE:HG12	2:R:54:VAL:HG11	1.51	0.92
1:G:230:ILE:HD13	1:G:309:LEU:HD22	1.51	0.92
1:I:205:ILE:HA	1:I:213:VAL:HG22	1.50	0.92
1:I:199:TYR:HE2	1:I:202:PRO:HA	1.34	0.92
1:E:248:LEU:HD22	1:E:323:VAL:HG11	1.52	0.92
1:L:6:VAL:HG12	1:L:521:VAL:HG22	1.51	0.92
2:V:25:ILE:HG13	2:V:26:VAL:HG23	1.51	0.92
1:L:219:PHE:HA	1:L:317:LEU:HD21	1.48	0.92
2:Y:34:LYS:HE3	2:Z:76:GLU:HA	1.52	0.92
1:B:305:ILE:HG22	1:B:306:GLY:H	1.34	0.92
1:K:349:ILE:HD12	1:K:352:GLN:HB2	1.51	0.92
2:O:8:ASP:HA	2:O:57:LEU:HD21	1.50	0.91
1:N:477:GLY:HA3	1:N:488:MET:HE2	1.52	0.91
1:A:276:VAL:HG11	1:A:325:ILE:HG21	1.52	0.91
1:M:6:VAL:HG12	1:M:521:VAL:HG22	1.53	0.91
1:C:321:LYS:HB2	1:C:334:ASP:HB3	1.52	0.91
2:1:64:ILE:HG23	2:1:95:VAL:HB	1.51	0.91
1:B:325:ILE:HG13	1:B:330:THR:HG23	1.51	0.91
1:G:358:SER:HA	1:G:362:ARG:HE	1.35	0.91
2:R:49:LEU:CD1	2:S:50:GLU:O	2.18	0.91
2:R:49:LEU:CD1	2:S:50:GLU:HA	2.01	0.91
1:D:228:SER:HA	1:D:255:GLU:HB2	1.52	0.91
2:W:9:ARG:HG2	2:W:9:ARG:HH11	1.36	0.91
1:L:40:LEU:HD13	1:L:59:GLU:HG3	1.52	0.90
1:J:305:ILE:HG23	1:J:307:MET:H	1.34	0.90
2:R:6:LEU:CD2	2:R:7:HIS:CG	2.55	0.90
1:J:214:GLU:HG2	1:J:324:VAL:HG22	1.49	0.90
1:H:207:LYS:HB3	1:H:208:PRO:HD3	1.51	0.90
1:A:6:VAL:HG12	1:A:521:VAL:HG22	1.50	0.90

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:F:220:ILE:HG13	1:F:296:THR:HG21	1.51	0.90
1:C:420:ILE:HD11	1:C:470:LYS:HG3	1.53	0.90
2:U:15:LYS:HG2	2:U:38:GLY:HA2	1.51	0.90
1:N:225:LYS:HB3	1:N:231:ARG:HH21	1.37	0.90
1:K:221:LEU:HD13	1:K:317:LEU:HD21	1.53	0.90
1:G:222:LEU:HD22	1:G:300:VAL:HG22	1.52	0.89
2:V:11:ILE:HG22	2:V:85:ILE:HG12	1.55	0.89
1:L:412:VAL:CG1	1:L:418:ALA:CB	2.48	0.89
1:H:278:ALA:HB3	1:H:285:ARG:HD2	1.51	0.89
1:G:240:VAL:HG11	1:G:247:LEU:HD21	1.54	0.89
1:G:5:ASP:HB2	1:G:524:LEU:HD23	1.55	0.89
1:N:325:ILE:HG13	1:N:330:THR:HG23	1.53	0.89
2:U:49:LEU:C	2:U:50:GLU:OE1	2.10	0.88
2:Z:15:LYS:HG2	2:Z:38:GLY:HA2	1.54	0.88
1:N:6:VAL:HG12	1:N:521:VAL:HG22	1.54	0.88
1:M:405:ALA:HB1	1:M:498:LYS:HB3	1.56	0.88
2:1:17:VAL:HG23	2:1:34:LYS:HA	1.53	0.88
1:B:301:ILE:HG12	1:B:307:MET:HE1	1.55	0.88
2:1:65:VAL:HG12	2:1:94:ILE:HG22	1.55	0.88
2:V:57:LEU:HD13	2:V:88:GLU:HG3	1.56	0.88
1:E:221:LEU:HD21	1:E:301:ILE:HD12	1.56	0.88
1:A:177:VAL:HG21	1:A:397:GLU:HG2	1.53	0.87
1:A:40:LEU:HD13	1:A:59:GLU:HG3	1.56	0.87
1:F:205:ILE:HA	1:F:213:VAL:HG22	1.56	0.87
1:G:77:VAL:HG21	1:G:507:ALA:HA	1.54	0.87
2:Y:65:VAL:HG12	2:Y:94:ILE:HG22	1.56	0.87
2:R:49:LEU:CD1	2:S:50:GLU:C	2.42	0.87
1:H:302:SER:CB	1:H:305:ILE:HD13	2.03	0.87
1:K:518:GLU:HB3	1:L:29:VAL:HG11	1.55	0.87
2:W:20:LYS:HA	2:W:21:SER:CB	1.99	0.87
2:S:40:VAL:HG12	2:S:41:LEU:H	1.40	0.87
2:W:17:VAL:HG23	2:W:19:THR:H	1.39	0.87
1:C:171:LYS:HD2	1:C:407:VAL:HG11	1.56	0.87
2:W:1:MET:HE2	2:W:79:ASP:HB3	1.54	0.87
1:E:234:LEU:HD23	1:E:235:PRO:CD	2.04	0.86
1:E:183:LEU:HD22	1:E:384:ALA:HA	1.55	0.86
1:H:77:VAL:HG21	1:H:507:ALA:HA	1.56	0.86
2:Q:37:ARG:HG2	2:Q:66:ILE:HG12	1.58	0.86
2:1:95:VAL:HG22	2:2:3:ILE:HG21	1.57	0.86
1:C:197:ARG:HE	1:C:277:LYS:HD2	1.39	0.86
1:D:420:ILE:CD1	1:D:448:GLU:HG2	2.06	0.86

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:1:49:LEU:CD1	2:1:50:GLU:O	2.23	0.86
1:D:420:ILE:HD13	1:D:448:GLU:HG2	1.56	0.86
1:I:40:LEU:HD13	1:I:59:GLU:HG3	1.58	0.86
1:A:353:ILE:HG12	1:A:365:LEU:HD13	1.56	0.86
1:N:288:MET:HG2	1:N:368:ARG:HD3	1.57	0.86
1:N:77:VAL:HB	1:N:510:VAL:HG21	1.56	0.86
1:A:245:LYS:HB3	1:A:246:PRO:CD	2.05	0.86
1:A:325:ILE:HG13	1:A:330:THR:HG23	1.57	0.86
1:B:228:SER:HB3	1:B:255:GLU:HG2	1.56	0.86
1:C:232:GLU:HG3	1:C:310:GLU:HG3	1.56	0.85
1:F:236:VAL:HG11	1:F:312:ALA:HB3	1.56	0.85
1:F:284:ARG:HD3	1:F:364:LYS:HE2	1.57	0.85
1:E:5:ASP:HB2	1:E:524:LEU:HD23	1.58	0.85
1:F:305:ILE:HG22	1:F:306:GLY:H	1.40	0.85
1:B:230:ILE:CG2	1:B:309:LEU:HD13	2.06	0.85
1:M:305:ILE:HG22	1:M:306:GLY:H	1.38	0.85
2:R:78:ILE:HG13	2:R:79:ASP:H	1.38	0.85
2:T:17:VAL:HG23	2:T:33:ALA:HB2	1.58	0.85
1:M:197:ARG:HD2	1:M:277:LYS:HB2	1.58	0.85
1:M:429:LEU:HG	1:M:440:ILE:HD13	1.59	0.85
1:G:455:VAL:HG13	1:G:460:GLU:HB2	1.57	0.85
1:M:234:LEU:HB3	1:M:235:PRO:HD2	1.56	0.85
2:Y:11:ILE:HG22	2:Y:85:ILE:HB	1.59	0.85
1:D:305:ILE:HG22	1:D:306:GLY:H	1.41	0.85
1:H:197:ARG:CG	1:H:277:LYS:HB3	2.06	0.85
1:E:234:LEU:O	1:E:236:VAL:N	2.10	0.84
1:I:197:ARG:NH1	1:I:277:LYS:HD2	1.92	0.84
1:B:276:VAL:HG21	1:B:325:ILE:HG21	1.59	0.84
1:D:205:ILE:HA	1:D:213:VAL:HG22	1.59	0.84
1:E:229:ASN:OD1	1:E:230:ILE:HD12	1.77	0.84
2:W:55:LYS:HE3	2:X:53:GLU:HG2	1.57	0.84
1:B:77:VAL:HG21	1:B:507:ALA:HA	1.60	0.84
1:H:326:ASN:HB3	1:H:329:THR:HG22	1.57	0.84
1:I:252:GLU:HG3	1:I:285:ARG:NH1	1.92	0.84
1:J:278:ALA:HB1	1:J:285:ARG:HD2	1.60	0.84
1:K:455:VAL:HG13	1:K:460:GLU:HB2	1.59	0.84
1:M:279:PRO:HG2	1:M:285:ARG:HA	1.57	0.84
2:2:49:LEU:N	2:2:52:GLY:HA2	1.93	0.84
1:F:245:LYS:HB3	1:F:246:PRO:HD2	1.60	0.84
1:B:240:VAL:HG11	1:B:247:LEU:HD23	1.60	0.84
1:F:77:VAL:HB	1:F:510:VAL:HG21	1.59	0.83

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:G:40:LEU:HD13	1:G:59:GLU:HG3	1.60	0.83
1:J:357:THR:HG21	1:J:361:ASP:HB2	1.59	0.83
2:V:1:MET:N	2:2:97:ALA:O	2.11	0.83
1:B:351:GLN:HG2	1:C:210:THR:HG23	1.60	0.83
1:G:223:ALA:HB2	1:G:251:ALA:HA	1.58	0.83
1:J:314:LEU:HD23	1:J:314:LEU:O	1.77	0.83
1:A:181:THR:HG21	1:A:380:LYS:HD3	1.58	0.83
1:B:301:ILE:HG12	1:B:307:MET:CE	2.08	0.83
1:C:236:VAL:CG2	1:C:312:ALA:HB3	2.07	0.83
1:N:419:LEU:HD21	1:N:500:THR:HG23	1.60	0.83
2:2:78:ILE:HD12	2:2:78:ILE:H	1.44	0.83
1:L:205:ILE:HD13	1:L:211:GLY:HA2	1.60	0.83
2:V:50:GLU:HG3	2:W:50:GLU:HB3	1.61	0.83
1:C:225:LYS:HG3	1:C:303:GLU:HG3	1.60	0.83
1:D:278:ALA:HB1	1:D:285:ARG:HG3	1.60	0.83
1:F:221:LEU:HD13	1:F:317:LEU:HD21	1.61	0.82
1:N:245:LYS:HB3	1:N:246:PRO:HD2	1.60	0.82
2:1:50:GLU:O	2:1:51:ASN:OD1	1.96	0.82
1:J:162:ILE:HG21	1:J:403:THR:HG21	1.59	0.82
1:N:305:ILE:HG22	1:N:306:GLY:H	1.44	0.82
1:B:278:ALA:CB	1:B:285:ARG:HD2	2.10	0.82
1:F:27:VAL:HG12	1:F:90:THR:HG23	1.62	0.82
1:L:357:THR:HG21	1:L:361:ASP:HB2	1.59	0.82
1:N:228:SER:HA	1:N:258:ALA:HB2	1.59	0.82
2:Y:34:LYS:HD2	2:Z:76:GLU:HG2	1.62	0.82
1:M:234:LEU:HB3	1:M:235:PRO:CD	2.10	0.82
2:R:19:THR:HG21	2:R:33:ALA:HB1	1.61	0.81
2:W:78:ILE:HG13	2:W:79:ASP:H	1.44	0.81
1:H:367:GLU:CG	1:H:371:LYS:HE2	2.09	0.81
1:H:305:ILE:HG23	1:I:267:MET:HE2	1.62	0.81
2:P:17:VAL:HG12	2:P:35:SER:HB2	1.61	0.81
2:Y:40:VAL:HG13	2:Y:62:GLY:H	1.45	0.81
1:A:147:VAL:HG12	1:A:494:LEU:HB2	1.60	0.81
2:R:49:LEU:CD1	2:S:50:GLU:CA	2.58	0.81
2:V:50:GLU:O	2:V:51:ASN:ND2	2.12	0.81
1:L:218:PRO:HA	1:L:219:PHE:HB2	1.62	0.81
1:N:261:THR:HB	2:2:29:GLY:H	1.43	0.81
1:B:222:LEU:HD13	1:B:300:VAL:HA	1.61	0.81
2:R:12:VAL:HG12	2:R:40:VAL:HA	1.63	0.81
2:W:12:VAL:HG12	2:W:40:VAL:HA	1.63	0.81
1:F:233:MET:CG	1:F:234:LEU:HD23	2.06	0.81

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:R:59:VAL:HG22	2:R:94:ILE:HD11	1.62	0.81
1:H:174:VAL:HG23	1:H:376:VAL:HG13	1.62	0.81
1:J:204:PHE:CZ	1:J:262:LEU:HD22	2.16	0.81
1:L:27:VAL:HG12	1:L:90:THR:HG23	1.63	0.81
2:S:8:ASP:OD1	2:S:87:SER:OG	1.97	0.81
2:Z:93:ALA:CB	2:1:5:PRO:HA	2.10	0.81
1:F:321:LYS:HB2	1:F:334:ASP:HB3	1.63	0.81
1:K:205:ILE:HA	1:K:213:VAL:HG22	1.60	0.81
1:N:252:GLU:HA	1:N:285:ARG:HH12	1.46	0.81
1:H:305:ILE:HG22	1:H:306:GLY:H	1.45	0.81
2:O:6:LEU:CD1	2:U:93:ALA:HA	2.10	0.81
1:C:241:ALA:HA	1:C:271:VAL:HG11	1.62	0.81
1:K:353:ILE:HA	1:K:365:LEU:HD13	1.63	0.81
1:K:205:ILE:HD13	1:K:211:GLY:HA2	1.61	0.80
1:C:174:VAL:HG13	1:C:376:VAL:HG13	1.62	0.80
1:E:240:VAL:CG1	1:E:247:LEU:HD21	2.11	0.80
1:J:27:VAL:HG12	1:J:90:THR:HG23	1.62	0.80
1:L:218:PRO:HG2	1:L:220:ILE:HD11	1.63	0.80
2:X:92:LEU:HB3	2:Y:85:ILE:HD13	1.61	0.80
2:V:20:LYS:HG3	2:V:28:THR:HG23	1.63	0.80
1:G:230:ILE:CG2	1:G:309:LEU:HD11	2.11	0.80
1:F:308:GLU:HB2	1:F:311:LYS:HB3	1.63	0.80
1:G:254:VAL:HG12	1:G:259:LEU:HG	1.63	0.80
2:W:10:VAL:HG21	2:W:91:ILE:HD11	1.62	0.80
2:X:58:ASP:OD2	2:Y:7:HIS:ND1	2.14	0.80
1:D:360:TYR:CA	1:D:363:GLU:HG3	2.09	0.80
1:H:305:ILE:HG23	1:I:267:MET:CE	2.12	0.80
1:B:270:ILE:HG12	2:P:25:ILE:HD13	1.62	0.80
1:K:305:ILE:HG22	1:K:306:GLY:H	1.45	0.80
1:N:352:GLN:HB3	1:N:365:LEU:HD11	1.63	0.80
1:N:31:LEU:HD23	1:N:453:GLN:HB3	1.63	0.80
1:J:208:PRO:HB2	1:J:212:ALA:HB3	1.63	0.80
2:V:4:ARG:HD2	2:2:94:ILE:HD11	1.63	0.80
1:D:237:LEU:HD22	2:R:27:LEU:HD11	1.63	0.79
1:B:234:LEU:HA	1:B:237:LEU:HD13	1.64	0.79
2:O:97:ALA:HB2	2:P:1:MET:H1	1.47	0.79
2:R:20:LYS:HA	2:R:21:SER:CB	2.03	0.79
2:W:36:THR:HG23	2:W:37:ARG:HG3	1.65	0.79
1:N:228:SER:CB	1:N:255:GLU:HB2	2.13	0.79
1:G:230:ILE:HG21	1:G:309:LEU:HD11	1.64	0.79
1:B:199:TYR:CE1	1:B:202:PRO:HA	2.18	0.79

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:F:168:LYS:HG2	1:F:189:VAL:HG11	1.62	0.79
2:1:37:ARG:HH22	2:2:78:ILE:CG2	1.96	0.79
1:A:267:MET:HG3	1:G:305:ILE:HG12	1.65	0.79
1:G:6:VAL:HG12	1:G:521:VAL:HG22	1.65	0.79
1:H:278:ALA:CB	1:H:285:ARG:HD2	2.12	0.79
1:I:241:ALA:HB1	2:W:25:ILE:HD13	1.65	0.79
1:M:303:GLU:HG3	1:M:309:LEU:HD11	1.65	0.79
1:J:270:ILE:HG21	2:X:27:LEU:CD2	2.12	0.79
2:1:37:ARG:NH2	2:2:78:ILE:HG23	1.98	0.79
1:F:74:VAL:O	1:F:77:VAL:HG12	1.83	0.79
1:L:208:PRO:CB	1:L:209:GLU:OE1	2.30	0.79
1:M:233:MET:HB2	1:M:237:LEU:HD21	1.63	0.79
1:N:199:TYR:CE1	1:N:202:PRO:HA	2.18	0.79
1:J:325:ILE:HG13	1:J:330:THR:HG23	1.64	0.78
1:K:381:VAL:HG22	1:K:396:VAL:HG21	1.65	0.78
1:N:234:LEU:HD12	1:N:234:LEU:H	1.46	0.78
2:Q:50:GLU:CD	2:Q:51:ASN:H	1.85	0.78
1:B:305:ILE:HD12	1:C:263:VAL:HG11	1.65	0.78
1:E:183:LEU:CD2	1:E:384:ALA:HA	2.13	0.78
1:G:204:PHE:CD2	1:G:274:ALA:HA	2.17	0.78
1:J:230:ILE:HG13	1:J:231:ARG:H	1.46	0.78
1:J:349:ILE:HD12	1:J:369:VAL:HG22	1.64	0.78
1:L:306:GLY:O	1:L:307:MET:HG3	1.82	0.78
1:L:479:ASN:HD22	1:L:493:ILE:HD11	1.46	0.78
1:M:5:ASP:HB2	1:M:524:LEU:CD2	2.14	0.78
2:R:49:LEU:HD22	2:R:50:GLU:OE1	1.83	0.78
1:E:208:PRO:HB2	1:E:212:ALA:HB3	1.66	0.78
1:E:240:VAL:HG11	1:E:247:LEU:HD21	1.66	0.78
1:F:320:ALA:HA	1:F:335:GLY:HA2	1.65	0.78
1:F:325:ILE:HG13	1:F:330:THR:HG23	1.66	0.78
2:P:57:LEU:HD13	2:P:88:GLU:HB2	1.65	0.78
2:T:92:LEU:HD23	2:U:85:ILE:HD13	1.65	0.78
1:N:77:VAL:CG2	1:N:507:ALA:HA	2.12	0.78
2:T:12:VAL:HB	2:T:39:GLU:O	1.84	0.78
1:F:420:ILE:HD12	1:F:451:LEU:HD13	1.66	0.78
1:G:246:PRO:HA	1:G:272:LYS:O	1.84	0.78
1:H:40:LEU:HD13	1:H:59:GLU:HG3	1.66	0.78
1:I:74:VAL:O	1:I:77:VAL:HG12	1.83	0.78
1:K:215:LEU:HB2	1:K:323:VAL:HG22	1.65	0.78
1:C:177:VAL:HG11	1:C:397:GLU:CG	2.12	0.78
1:F:278:ALA:HB1	1:F:279:PRO:HD2	1.65	0.78

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:G:205:ILE:HD13	1:G:211:GLY:HA2	1.66	0.78
1:H:238:GLU:HA	2:V:25:ILE:CD1	2.14	0.77
1:L:207:LYS:HG2	1:L:208:PRO:HD2	1.66	0.77
1:E:214:GLU:HG2	1:E:324:VAL:HG22	1.63	0.77
1:E:5:ASP:HB2	1:E:524:LEU:CD2	2.14	0.77
1:G:200:LEU:HD11	1:G:277:LYS:HB2	1.66	0.77
1:J:322:ARG:HB3	1:J:333:ILE:CD1	2.15	0.77
1:N:175:ILE:HG12	1:N:377:ALA:HB3	1.66	0.77
1:F:218:PRO:HB3	1:F:246:PRO:CG	2.11	0.77
1:G:306:GLY:O	1:G:307:MET:HG3	1.82	0.77
1:H:349:ILE:HD12	1:H:369:VAL:HG22	1.65	0.77
1:N:40:LEU:HD21	1:N:56:VAL:HG22	1.66	0.77
1:A:188:ASP:HB2	1:A:378:VAL:HG23	1.64	0.77
1:C:231:ARG:HD3	1:C:258:ALA:HA	1.67	0.77
1:C:254:VAL:HG12	1:C:259:LEU:HG	1.67	0.77
1:F:5:ASP:HB2	1:F:524:LEU:CD2	2.13	0.77
1:H:245:LYS:HB3	1:H:246:PRO:HD2	1.64	0.77
1:M:199:TYR:CE1	1:M:202:PRO:HA	2.19	0.77
2:U:51:ASN:O	2:U:53:GLU:N	2.17	0.77
1:A:245:LYS:HB3	1:A:246:PRO:HD2	1.65	0.77
1:K:268:ARG:HG3	2:Y:27:LEU:HD21	1.64	0.77
1:B:233:MET:C	1:B:235:PRO:HD2	2.05	0.77
1:H:354:GLU:O	1:H:355:GLU:HG3	1.84	0.77
1:F:221:LEU:HD13	1:F:317:LEU:CD2	2.14	0.77
1:D:229:ASN:OD1	1:D:230:ILE:N	2.17	0.77
1:G:209:GLU:HG2	1:G:210:THR:H	1.49	0.77
1:E:174:VAL:HG22	1:E:376:VAL:HG13	1.67	0.77
1:G:232:GLU:HG3	1:G:310:GLU:OE2	1.85	0.77
2:R:8:ASP:O	2:R:57:LEU:CD1	2.32	0.76
1:B:236:VAL:HG21	1:B:312:ALA:HB3	1.67	0.76
1:H:74:VAL:O	1:H:77:VAL:HG12	1.85	0.76
1:L:479:ASN:ND2	1:L:493:ILE:HD11	2.00	0.76
1:H:385:THR:HG22	1:N:80:LYS:HG2	1.66	0.76
2:S:12:VAL:HG22	2:S:40:VAL:HG13	1.66	0.76
1:I:228:SER:HB3	1:I:255:GLU:HB2	1.67	0.76
1:L:355:GLU:HG3	1:L:356:ALA:H	1.51	0.76
2:R:17:VAL:HG23	2:R:19:THR:H	1.51	0.76
2:O:5:PRO:HA	2:U:93:ALA:CB	2.15	0.76
2:X:64:ILE:HG23	2:X:95:VAL:HG13	1.67	0.76
1:B:291:ASP:HB3	1:B:372:LEU:HD21	1.67	0.76
1:L:226:LYS:N	1:L:252:GLU:HB3	2.00	0.76

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:M:5:ASP:HB2	1:M:524:LEU:HD21	1.67	0.76
1:J:270:ILE:HG21	2:X:27:LEU:HD23	1.67	0.76
2:2:73:VAL:HG22	2:2:86:MET:HB3	1.68	0.76
1:F:218:PRO:CB	1:F:246:PRO:HG2	2.12	0.76
2:R:21:SER:N	2:R:27:LEU:O	2.15	0.76
2:R:49:LEU:O	2:R:51:ASN:N	2.18	0.76
1:A:77:VAL:HG21	1:A:507:ALA:HA	1.68	0.76
1:E:195:PHE:CE1	1:E:330:THR:HB	2.20	0.76
2:Q:6:LEU:HD13	2:Q:7:HIS:HD2	1.51	0.76
1:A:200:LEU:HD11	1:A:277:LYS:HB2	1.66	0.76
1:C:291:ASP:OD2	1:C:368:ARG:HD3	1.86	0.76
2:O:5:PRO:HA	2:U:93:ALA:HB2	1.66	0.76
2:W:20:LYS:CA	2:W:21:SER:HB3	2.07	0.76
1:K:291:ASP:HB2	1:K:372:LEU:HD21	1.66	0.76
2:Z:65:VAL:CG1	2:Z:94:ILE:HG22	2.16	0.76
1:N:77:VAL:HG21	1:N:507:ALA:CA	2.15	0.75
2:P:80:ASN:O	2:P:80:ASN:OD1	2.05	0.75
1:D:350:ARG:HD2	1:D:353:ILE:HD12	1.67	0.75
1:F:218:PRO:O	1:F:219:PHE:CG	2.38	0.75
2:V:6:LEU:HD21	2:2:91:ILE:HG22	1.67	0.75
1:C:23:LEU:HD23	1:C:60:ILE:HB	1.69	0.75
1:C:291:ASP:O	1:C:345:ARG:NH1	2.19	0.75
1:F:419:LEU:HD11	1:F:500:THR:HG23	1.68	0.75
1:G:230:ILE:HD13	1:G:309:LEU:CD2	2.17	0.75
2:S:12:VAL:HG13	2:S:40:VAL:HA	1.68	0.75
1:B:215:LEU:HD12	1:B:248:LEU:HD21	1.66	0.75
1:L:367:GLU:CG	1:L:371:LYS:HE2	2.16	0.75
1:L:381:VAL:HG11	1:L:392:LYS:HB3	1.67	0.75
2:U:48:ILE:C	2:U:49:LEU:HD12	2.05	0.75
1:B:307:MET:HG2	1:B:311:LYS:HZ2	1.52	0.75
1:C:367:GLU:CG	1:C:371:LYS:HE2	2.17	0.75
1:I:236:VAL:CG2	1:I:312:ALA:HB3	2.17	0.75
1:C:350:ARG:HH12	1:C:369:VAL:HG11	1.50	0.75
1:E:245:LYS:HE2	1:E:246:PRO:HD2	1.67	0.75
1:G:239:ALA:HB1	1:G:314:LEU:HD13	1.68	0.75
1:M:177:VAL:CG1	1:M:397:GLU:HG2	2.14	0.75
2:U:49:LEU:CB	2:U:50:GLU:OE1	2.35	0.75
1:E:276:VAL:HG11	1:E:325:ILE:HG21	1.67	0.75
1:F:220:ILE:N	1:F:318:GLY:O	2.19	0.75
1:J:245:LYS:HB3	1:J:246:PRO:HD2	1.68	0.75
1:E:200:LEU:HD11	1:E:277:LYS:HB2	1.67	0.75

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:M:302:SER:HB2	1:M:305:ILE:HD13	1.66	0.75
1:F:360:TYR:HA	1:F:363:GLU:HG3	1.68	0.74
1:J:357:THR:CG2	1:J:361:ASP:HB2	2.16	0.74
2:Z:65:VAL:HG12	2:Z:94:ILE:HG22	1.68	0.74
1:H:241:ALA:HB3	2:V:25:ILE:HD12	1.67	0.74
1:H:175:ILE:HG12	1:H:377:ALA:HB3	1.67	0.74
1:M:91:THR:OG1	3:M:601:ADP:O3B	2.05	0.74
1:I:151:SER:HB3	1:I:399:ALA:HA	1.67	0.74
1:L:234:LEU:HD23	2:Z:23:GLY:HA3	1.69	0.74
1:N:219:PHE:HB3	1:N:317:LEU:HB3	1.70	0.74
2:O:95:VAL:HA	2:P:3:ILE:HG21	1.58	0.74
2:S:50:GLU:CG	2:S:51:ASN:H	2.01	0.74
1:F:420:ILE:HD11	1:F:451:LEU:HB3	1.69	0.74
1:F:77:VAL:HG21	1:F:507:ALA:HA	1.66	0.74
1:I:199:TYR:CE2	1:I:202:PRO:HA	2.19	0.74
1:K:302:SER:CB	1:K:305:ILE:HD13	2.15	0.74
1:E:229:ASN:OD1	1:E:230:ILE:CD1	2.35	0.74
2:T:49:LEU:HD13	2:U:50:GLU:H	1.51	0.74
1:D:417:VAL:HA	1:D:420:ILE:HG22	1.70	0.74
1:G:230:ILE:HG21	1:G:309:LEU:CD2	2.17	0.74
1:L:230:ILE:HG12	1:L:309:LEU:HD21	1.68	0.74
1:L:175:ILE:HG12	1:L:377:ALA:HB3	1.70	0.74
1:M:241:ALA:CA	1:M:271:VAL:HG11	2.16	0.74
2:V:69:ASP:CB	2:V:73:VAL:HG11	2.18	0.74
1:B:231:ARG:HH22	2:P:31:ALA:HB2	1.52	0.74
1:J:266:THR:HG22	1:J:273:VAL:H	1.52	0.74
1:C:195:PHE:CE1	1:C:330:THR:HB	2.23	0.74
1:C:345:ARG:O	1:C:348:GLN:HG3	1.86	0.74
1:E:232:GLU:O	1:E:234:LEU:N	2.20	0.74
1:F:247:LEU:HB2	1:F:273:VAL:HG13	1.69	0.74
1:J:6:VAL:HG12	1:J:521:VAL:HG22	1.70	0.74
1:E:151:SER:HB3	1:E:399:ALA:HA	1.69	0.74
1:C:207:LYS:HB3	1:C:208:PRO:HD3	1.68	0.74
1:F:305:ILE:HG22	1:F:306:GLY:N	2.02	0.74
1:B:231:ARG:HH12	2:P:31:ALA:HB2	1.52	0.74
1:G:350:ARG:HD3	1:G:353:ILE:HD12	1.70	0.73
1:G:5:ASP:HB2	1:G:524:LEU:CD2	2.18	0.73
1:H:220:ILE:HG22	1:H:222:LEU:HD11	1.69	0.73
1:H:236:VAL:CG2	1:H:312:ALA:HB3	2.18	0.73
1:L:234:LEU:HD12	1:L:234:LEU:H	1.52	0.73
1:M:338:GLU:HG2	1:M:341:ALA:H	1.53	0.73

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:N:222:LEU:HD22	1:N:300:VAL:HG22	1.70	0.73
1:B:307:MET:HG2	1:B:311:LYS:NZ	2.03	0.73
1:E:199:TYR:HA	1:E:276:VAL:HG12	1.69	0.73
2:V:56:PRO:O	2:V:57:LEU:CG	2.22	0.73
1:D:215:LEU:HB2	1:D:323:VAL:CG2	2.19	0.73
1:B:230:ILE:HD11	1:B:232:GLU:HB3	1.71	0.73
1:K:124:VAL:HG21	1:K:508:ALA:HB1	1.68	0.73
1:N:381:VAL:HG22	1:N:396:VAL:HG21	1.71	0.73
2:U:49:LEU:O	2:U:50:GLU:CD	2.26	0.73
2:Y:60:LYS:HE3	2:Y:61:VAL:HG22	1.70	0.73
1:C:313:THR:HB	1:C:315:GLU:HG3	1.71	0.73
1:C:6:VAL:CG1	1:C:521:VAL:HG22	2.15	0.73
1:I:305:ILE:HG22	1:I:306:GLY:H	1.52	0.73
1:B:278:ALA:HB1	1:B:285:ARG:HD2	1.69	0.73
1:C:294:THR:OG1	1:C:345:ARG:NH1	2.22	0.73
1:C:367:GLU:HG2	1:C:371:LYS:HE2	1.70	0.73
1:H:276:VAL:CG1	1:H:325:ILE:HD13	2.19	0.73
1:H:305:ILE:HG22	1:H:306:GLY:N	2.04	0.73
1:M:519:CYS:HB3	1:N:38:VAL:HG13	1.70	0.73
1:N:5:ASP:HB2	1:N:524:LEU:HD21	1.70	0.73
1:K:309:LEU:HD12	1:K:310:GLU:N	2.04	0.73
1:N:307:MET:HG2	1:N:311:LYS:HZ2	1.54	0.73
2:W:9:ARG:HG2	2:W:9:ARG:NH1	2.02	0.73
1:E:194:GLN:HG2	1:E:331:THR:HG23	1.71	0.73
1:C:305:ILE:HG22	1:C:306:GLY:H	1.52	0.72
1:G:240:VAL:CG1	1:G:247:LEU:HD21	2.19	0.72
1:K:228:SER:HB3	1:K:255:GLU:HB2	1.71	0.72
2:W:55:LYS:HE3	2:X:53:GLU:CG	2.19	0.72
2:Z:55:LYS:HE2	2:1:48:ILE:HD11	1.71	0.72
2:Q:17:VAL:HG12	2:Q:35:SER:H	1.52	0.72
2:Q:69:ASP:HB2	2:Q:73:VAL:HG11	1.70	0.72
1:I:196:ASP:O	1:I:196:ASP:OD1	2.06	0.72
2:T:51:ASN:O	2:T:51:ASN:OD1	2.07	0.72
1:F:225:LYS:CG	1:F:309:LEU:HD21	2.19	0.72
1:J:74:VAL:O	1:J:77:VAL:HG12	1.89	0.72
1:K:215:LEU:HB2	1:K:323:VAL:CG2	2.18	0.72
2:O:95:VAL:CG2	2:P:3:ILE:HG21	2.19	0.72
2:T:20:LYS:HB3	2:T:26:VAL:O	1.88	0.72
1:A:181:THR:CG2	1:A:380:LYS:HD3	2.19	0.72
1:A:223:ALA:HB3	1:A:251:ALA:HB2	1.71	0.72
1:C:420:ILE:HD11	1:C:470:LYS:CG	2.19	0.72

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:D:205:ILE:HD13	1:D:211:GLY:HA2	1.72	0.72
1:E:219:PHE:HB3	1:E:317:LEU:HB3	1.70	0.72
1:L:31:LEU:HD23	1:L:453:GLN:HB3	1.70	0.72
2:W:49:LEU:HB2	2:W:52:GLY:HA3	1.70	0.72
1:A:232:GLU:HG3	1:A:234:LEU:HD23	1.71	0.72
1:B:488:MET:CE	1:B:493:ILE:HD12	2.19	0.72
1:D:349:ILE:HA	1:D:352:GLN:HG3	1.71	0.72
1:F:5:ASP:HB2	1:F:524:LEU:HD23	1.71	0.72
2:S:14:ARG:HE	2:S:35:SER:HB2	1.55	0.72
1:E:230:ILE:HG12	1:E:309:LEU:HD13	1.71	0.72
1:J:28:LYS:HG2	1:J:94:VAL:HG22	1.70	0.72
1:L:455:VAL:HG13	1:L:460:GLU:HB2	1.71	0.72
1:N:236:VAL:HG22	1:N:312:ALA:HB3	1.69	0.72
1:E:6:VAL:HG12	1:E:521:VAL:HG22	1.71	0.72
1:F:199:TYR:HA	1:F:276:VAL:CG1	2.18	0.72
1:I:429:LEU:HG	1:I:440:ILE:HD13	1.71	0.72
2:V:69:ASP:HB2	2:V:73:VAL:HG11	1.70	0.72
1:B:350:ARG:HD3	1:B:353:ILE:HD12	1.72	0.72
1:D:419:LEU:HD21	1:D:500:THR:HG23	1.71	0.72
1:F:177:VAL:HG11	1:F:397:GLU:HG2	1.71	0.72
2:W:14:ARG:HD3	2:W:35:SER:HB3	1.72	0.72
1:E:429:LEU:HG	1:E:440:ILE:HD13	1.72	0.71
1:G:417:VAL:HA	1:G:420:ILE:HG22	1.71	0.71
1:J:203:TYR:CD2	1:J:263:VAL:HG13	2.25	0.71
1:K:345:ARG:HD2	1:K:348:GLN:NE2	2.05	0.71
1:N:207:LYS:HB3	1:N:208:PRO:HD3	1.72	0.71
2:T:97:ALA:CB	2:U:1:MET:HA	2.19	0.71
1:A:362:ARG:HA	1:A:365:LEU:HD23	1.72	0.71
1:D:199:TYR:CE1	1:D:202:PRO:HA	2.25	0.71
1:D:215:LEU:HB2	1:D:323:VAL:HG22	1.72	0.71
1:N:227:ILE:HG23	1:N:231:ARG:HB2	1.69	0.71
1:N:225:LYS:HB3	1:N:231:ARG:NH2	2.04	0.71
2:U:64:ILE:HG23	2:U:95:VAL:CG2	2.20	0.71
1:B:321:LYS:HB2	1:B:334:ASP:HB2	1.72	0.71
1:D:415:GLY:N	3:D:601:ADP:O2'	2.20	0.71
1:L:221:LEU:HD11	1:L:249:ILE:HG23	1.71	0.71
1:N:284:ARG:HD3	1:N:364:LYS:HE2	1.71	0.71
2:U:10:VAL:HG11	2:U:91:ILE:HD11	1.72	0.71
2:1:17:VAL:CG2	2:1:34:LYS:HA	2.21	0.71
1:C:321:LYS:CB	1:C:334:ASP:HB3	2.20	0.71
1:E:197:ARG:HG2	1:E:279:PRO:HA	1.72	0.71

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:G:248:LEU:CD2	1:G:325:ILE:HD11	2.20	0.71
1:H:5:ASP:HB2	1:H:524:LEU:CD2	2.19	0.71
2:2:14:ARG:NH1	2:2:84:LEU:HD21	2.05	0.71
1:M:206:ASN:ND2	1:M:214:GLU:O	2.23	0.71
1:C:238:GLU:HG2	2:Q:25:ILE:HG12	1.71	0.71
1:K:23:LEU:HD23	1:K:60:ILE:HB	1.69	0.71
2:V:56:PRO:C	2:V:57:LEU:HG	2.10	0.71
2:2:78:ILE:N	2:2:78:ILE:HD12	2.03	0.71
1:A:74:VAL:O	1:A:77:VAL:HG12	1.90	0.71
1:M:305:ILE:HG22	1:M:306:GLY:N	2.05	0.71
1:N:136:VAL:HG13	1:N:137:PRO:HD2	1.72	0.71
2:S:20:LYS:NZ	2:S:27:LEU:O	2.23	0.71
2:X:36:THR:HG23	2:X:37:ARG:HG3	1.73	0.71
1:E:206:ASN:OD1	1:E:207:LYS:HG2	1.91	0.71
1:F:40:LEU:HD13	1:F:59:GLU:HG3	1.71	0.71
1:H:27:VAL:HG12	1:H:90:THR:HG23	1.73	0.71
1:K:305:ILE:HG22	1:K:306:GLY:N	2.06	0.71
1:M:200:LEU:HD22	1:M:254:VAL:HB	1.73	0.71
1:G:205:ILE:HA	1:G:213:VAL:CG2	2.12	0.71
1:I:252:GLU:HG3	1:I:285:ARG:HH11	1.53	0.71
1:I:449:ALA:HB3	1:I:450:PRO:HD3	1.71	0.71
1:K:193:MET:SD	1:K:371:LYS:HB3	2.31	0.71
1:N:420:ILE:HG22	1:N:448:GLU:HA	1.71	0.71
1:F:232:GLU:N	1:F:232:GLU:OE1	2.23	0.70
2:W:50:GLU:O	2:W:51:ASN:ND2	2.25	0.70
2:Z:46:GLY:HA2	2:Z:57:LEU:CD1	2.21	0.70
2:2:12:VAL:HG22	2:2:40:VAL:HA	1.72	0.70
1:H:204:PHE:CD2	1:H:274:ALA:HA	2.27	0.70
1:L:230:ILE:CG2	1:L:309:LEU:HD21	2.20	0.70
2:O:36:THR:HB	2:O:67:PHE:O	1.91	0.70
2:O:96:GLU:C	2:P:1:MET:HA	2.06	0.70
2:V:57:LEU:HB2	2:V:88:GLU:OE2	1.91	0.70
1:C:253:ASP:OD1	1:C:277:LYS:HE2	1.90	0.70
1:D:305:ILE:HG22	1:D:306:GLY:N	2.06	0.70
1:F:418:ALA:O	1:F:422:VAL:HG13	1.92	0.70
1:K:305:ILE:HG23	1:L:264:VAL:HG22	1.72	0.70
1:B:305:ILE:HG22	1:B:306:GLY:N	2.06	0.70
1:C:219:PHE:HE2	1:C:245:LYS:HB2	1.55	0.70
1:G:412:VAL:HG12	1:G:413:ALA:N	2.05	0.70
1:N:239:ALA:HB1	1:N:314:LEU:HD11	1.73	0.70
1:C:197:ARG:CG	1:C:277:LYS:HB3	2.15	0.70

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:F:301:ILE:HG23	1:F:307:MET:HB3	1.72	0.70
1:L:219:PHE:CA	1:L:317:LEU:HD21	2.22	0.70
2:Q:79:ASP:O	2:Q:80:ASN:ND2	2.24	0.70
1:C:305:ILE:HG22	1:C:306:GLY:N	2.06	0.70
1:F:282:GLY:HA2	1:F:285:ARG:HG2	1.73	0.70
1:H:225:LYS:CG	1:H:303:GLU:HG3	2.21	0.70
2:Q:12:VAL:HG23	2:Q:39:GLU:O	1.90	0.70
1:C:232:GLU:CG	1:C:310:GLU:HG3	2.20	0.70
1:E:230:ILE:HG21	1:E:309:LEU:HB3	1.72	0.70
1:E:219:PHE:HB3	1:E:317:LEU:HD13	1.73	0.70
1:I:302:SER:CB	1:I:305:ILE:HD13	2.22	0.70
1:N:197:ARG:HG2	1:N:277:LYS:HB3	1.73	0.70
1:A:27:VAL:HG12	1:A:90:THR:HG23	1.72	0.70
1:L:322:ARG:HB3	1:L:333:ILE:HD12	1.72	0.70
1:N:34:LYS:NZ	1:N:483:GLU:OE2	2.24	0.70
2:S:50:GLU:HG2	2:S:51:ASN:H	1.55	0.70
1:A:271:VAL:HG11	2:O:25:ILE:HD13	1.72	0.70
1:B:197:ARG:HE	1:B:277:LYS:HD3	1.55	0.70
2:Q:60:LYS:NZ	2:Q:63:ASP:HB3	2.06	0.70
2:W:17:VAL:HG23	2:W:19:THR:N	2.07	0.70
2:Y:74:LYS:NZ	2:Y:76:GLU:OE2	2.25	0.70
2:Z:55:LYS:HE2	2:1:48:ILE:CD1	2.22	0.70
1:E:25:ASP:HA	1:E:28:LYS:HE2	1.74	0.70
1:F:301:ILE:HG23	1:F:307:MET:CB	2.21	0.70
1:C:196:ASP:O	1:C:197:ARG:NH1	2.25	0.69
1:E:222:LEU:HD22	1:E:289:LEU:HD22	1.72	0.69
1:H:5:ASP:HB2	1:H:524:LEU:HD23	1.74	0.69
2:V:20:LYS:NZ	2:V:33:ALA:O	2.24	0.69
1:C:85:ALA:HB1	1:C:499:VAL:HG12	1.73	0.69
1:C:91:THR:N	4:C:602:BEF:F3	2.08	0.69
1:G:247:LEU:HB2	1:G:273:VAL:HG13	1.74	0.69
1:H:177:VAL:HG11	1:H:397:GLU:HG2	1.74	0.69
1:J:204:PHE:HE1	1:J:263:VAL:HA	1.58	0.69
2:O:6:LEU:HD12	2:U:93:ALA:HA	1.74	0.69
1:B:74:VAL:O	1:B:77:VAL:HG12	1.91	0.69
1:C:325:ILE:HG12	1:C:330:THR:HG23	1.73	0.69
1:J:213:VAL:O	1:J:324:VAL:HA	1.92	0.69
1:J:233:MET:SD	1:J:309:LEU:O	2.50	0.69
2:O:18:GLU:HB3	2:O:28:THR:HB	1.73	0.69
1:A:179:ASP:OD1	1:A:393:LYS:HD3	1.92	0.69
1:B:219:PHE:HB3	1:B:317:LEU:HD13	1.72	0.69

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:L:89:THR:N	4:L:602:BEF:F2	2.14	0.69
2:X:57:LEU:H	2:X:57:LEU:HD12	1.57	0.69
1:G:219:PHE:HA	1:G:318:GLY:O	1.92	0.69
1:H:205:ILE:HA	1:H:213:VAL:HG12	1.74	0.69
1:H:479:ASN:ND2	1:H:493:ILE:HD11	2.08	0.69
1:N:197:ARG:HH11	1:N:198:GLY:H	1.40	0.69
2:R:12:VAL:HG12	2:R:40:VAL:HG13	1.73	0.69
1:F:238:GLU:HG3	2:T:23:GLY:HA3	1.73	0.69
1:A:355:GLU:HG3	1:A:356:ALA:H	1.57	0.69
1:G:219:PHE:HE2	1:G:245:LYS:HB2	1.57	0.69
1:I:415:GLY:N	3:I:601:ADP:O2'	2.20	0.69
1:K:74:VAL:O	1:K:77:VAL:HG13	1.93	0.69
1:L:418:ALA:O	1:L:422:VAL:HG13	1.93	0.69
1:N:262:LEU:HD11	1:N:273:VAL:HB	1.74	0.69
2:Q:20:LYS:NZ	2:Q:33:ALA:O	2.25	0.69
2:X:70:GLY:O	2:X:73:VAL:HG12	1.93	0.69
2:1:95:VAL:HG13	2:2:3:ILE:HG22	1.74	0.69
2:1:95:VAL:HA	2:2:3:ILE:HG22	1.73	0.69
1:C:223:ALA:CB	1:C:227:ILE:HD11	2.22	0.69
1:F:124:VAL:HG13	1:F:504:LEU:CD1	2.22	0.69
1:J:496:PRO:HG2	1:J:499:VAL:CG1	2.23	0.69
1:M:405:ALA:CB	1:M:498:LYS:HD3	2.22	0.69
2:Q:92:LEU:H	2:Q:92:LEU:HD23	1.54	0.69
2:U:64:ILE:HG23	2:U:95:VAL:HG23	1.74	0.69
1:C:232:GLU:OE1	1:C:232:GLU:N	2.22	0.69
1:J:177:VAL:HG11	1:J:397:GLU:HG2	1.73	0.69
1:K:199:TYR:HE1	1:K:202:PRO:HA	1.57	0.69
1:L:208:PRO:O	1:L:209:GLU:CG	2.41	0.69
1:D:418:ALA:O	1:D:422:VAL:HG13	1.92	0.69
1:G:226:LYS:CB	1:G:253:ASP:HB2	2.23	0.69
1:H:233:MET:CE	1:H:237:LEU:HD12	2.22	0.69
1:H:350:ARG:HA	1:H:353:ILE:HD12	1.73	0.69
1:L:204:PHE:CD2	1:L:274:ALA:HA	2.28	0.69
2:R:49:LEU:HD11	2:S:50:GLU:CA	2.22	0.69
2:T:11:ILE:HD11	2:T:42:ALA:HB3	1.75	0.69
1:A:234:LEU:HA	1:A:237:LEU:HB3	1.75	0.69
1:E:513:LEU:HD11	1:F:388:GLU:HA	1.75	0.69
1:F:218:PRO:O	1:F:219:PHE:CE1	2.44	0.69
1:G:280:GLY:O	1:G:285:ARG:NE	2.22	0.69
1:I:238:GLU:HG2	2:W:27:LEU:CD1	2.23	0.69
1:K:234:LEU:HB3	1:K:235:PRO:HD3	1.73	0.69

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:L:291:ASP:HB2	1:L:372:LEU:HD21	1.74	0.69
1:M:278:ALA:HB1	1:M:279:PRO:HD2	1.74	0.69
2:R:6:LEU:HD21	2:R:7:HIS:CG	2.25	0.69
2:T:55:LYS:HG2	2:U:48:ILE:CD1	2.22	0.69
2:X:93:ALA:CB	2:Y:5:PRO:HA	2.23	0.69
1:D:289:LEU:O	1:D:293:ALA:N	2.26	0.68
1:F:320:ALA:HB1	1:F:334:ASP:O	1.93	0.68
1:F:172:GLU:O	1:F:370:ALA:HB2	1.93	0.68
1:G:358:SER:HA	1:G:362:ARG:NE	2.07	0.68
1:I:241:ALA:CB	2:W:25:ILE:HD13	2.23	0.68
2:1:17:VAL:HA	2:1:35:SER:H	1.57	0.68
1:A:254:VAL:HG12	1:A:259:LEU:HB2	1.74	0.68
1:A:236:VAL:CG2	1:A:312:ALA:HB3	2.21	0.68
1:D:226:LYS:HB3	1:D:252:GLU:HG2	1.75	0.68
1:G:350:ARG:HD3	1:G:353:ILE:CD1	2.22	0.68
1:I:5:ASP:HB2	1:I:524:LEU:HD23	1.74	0.68
1:K:449:ALA:HB3	1:K:450:PRO:HD3	1.74	0.68
1:L:74:VAL:O	1:L:77:VAL:HG12	1.94	0.68
1:M:231:ARG:HH21	1:M:257:GLU:HB3	1.58	0.68
1:N:429:LEU:O	1:N:430:ARG:NH1	2.25	0.68
1:F:238:GLU:HG3	2:T:23:GLY:CA	2.24	0.68
2:X:23:GLY:O	2:X:26:VAL:HG12	1.92	0.68
1:A:199:TYR:CE1	1:A:202:PRO:HA	2.28	0.68
1:C:74:VAL:O	1:C:77:VAL:HG12	1.92	0.68
1:D:226:LYS:HA	1:D:252:GLU:HB3	1.75	0.68
1:D:302:SER:OG	1:D:305:ILE:HD13	1.92	0.68
1:E:74:VAL:O	1:E:77:VAL:HG12	1.93	0.68
1:K:18:ARG:NH1	1:K:18:ARG:HB3	2.07	0.68
1:K:237:LEU:HD12	1:K:238:GLU:N	2.08	0.68
1:L:245:LYS:HB3	1:L:246:PRO:CD	2.20	0.68
1:N:254:VAL:HG12	1:N:259:LEU:HG	1.75	0.68
2:V:1:MET:HG2	2:V:1:MET:O	1.93	0.68
1:H:234:LEU:CD2	2:V:22:ALA:HB1	2.23	0.68
2:Z:64:ILE:HG23	2:Z:95:VAL:HG22	1.74	0.68
1:A:420:ILE:HD13	1:A:451:LEU:HD13	1.74	0.68
1:B:367:GLU:HG2	1:B:371:LYS:HE2	1.73	0.68
1:D:138:CYS:HB2	1:D:411:VAL:HG13	1.75	0.68
1:D:357:THR:CG2	1:D:361:ASP:HB2	2.23	0.68
1:M:23:LEU:HD22	1:M:74:VAL:HG23	1.75	0.68
1:B:175:ILE:CD1	1:B:377:ALA:HB3	2.23	0.68
1:E:411:VAL:HG21	1:E:494:LEU:HD23	1.75	0.68

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:G:241:ALA:HA	1:G:271:VAL:HG11	1.75	0.68
1:C:449:ALA:HB3	1:C:450:PRO:HD3	1.75	0.68
1:D:5:ASP:HB2	1:D:524:LEU:CD2	2.23	0.68
1:G:241:ALA:HA	1:G:271:VAL:CG1	2.23	0.68
1:H:25:ASP:HA	1:H:28:LYS:HE2	1.75	0.68
1:L:149:THR:HG23	1:L:159:GLY:HA3	1.76	0.68
1:N:25:ASP:HA	1:N:28:LYS:HE2	1.76	0.68
1:D:217:SER:HA	1:D:320:ALA:O	1.94	0.68
1:M:357:THR:HG21	1:M:361:ASP:HB3	1.75	0.68
2:U:40:VAL:HG12	2:U:41:LEU:H	1.58	0.68
1:B:288:MET:HG2	1:B:368:ARG:HG3	1.74	0.68
1:D:226:LYS:CA	1:D:252:GLU:HB3	2.23	0.68
1:F:304:GLU:HB3	1:F:305:ILE:HD12	1.75	0.68
2:U:3:ILE:HD13	2:U:78:ILE:HD13	1.76	0.68
2:W:3:ILE:HD11	2:W:11:ILE:HD12	1.76	0.68
1:F:127:ALA:CA	1:F:426:LEU:HD11	2.24	0.68
1:F:90:THR:N	3:F:601:ADP:O2B	2.26	0.68
1:I:174:VAL:HG21	1:I:367:GLU:HA	1.75	0.68
1:L:254:VAL:HG12	1:L:259:LEU:HG	1.76	0.68
1:K:268:ARG:HG3	2:Y:27:LEU:CD2	2.23	0.68
2:Z:46:GLY:HA2	2:Z:57:LEU:HD12	1.76	0.68
1:B:175:ILE:HD13	1:B:377:ALA:HB3	1.76	0.68
1:C:231:ARG:HH12	1:C:234:LEU:HD21	1.58	0.68
1:E:197:ARG:NH1	1:E:279:PRO:O	2.28	0.68
1:E:301:ILE:CG1	1:E:307:MET:HE1	2.10	0.68
1:H:193:MET:SD	1:H:371:LYS:HB3	2.34	0.68
1:H:124:VAL:HG13	1:H:504:LEU:HD12	1.76	0.68
1:J:479:ASN:ND2	1:J:493:ILE:HD11	2.09	0.68
1:M:177:VAL:HG11	1:M:397:GLU:CG	2.23	0.68
1:A:291:ASP:OD1	1:A:345:ARG:NH2	2.24	0.67
1:A:362:ARG:HA	1:A:365:LEU:CD2	2.24	0.67
1:C:226:LYS:HG2	1:C:252:GLU:CG	2.19	0.67
1:D:74:VAL:O	1:D:77:VAL:HG12	1.94	0.67
1:H:291:ASP:OD1	1:H:345:ARG:NH2	2.20	0.67
3:J:601:ADP:O3B	4:J:602:BEF:F1	2.02	0.67
1:N:488:MET:SD	1:N:493:ILE:HD11	2.35	0.67
2:U:37:ARG:HG2	2:U:66:ILE:HG12	1.76	0.67
2:Y:20:LYS:HB3	2:Y:26:VAL:O	1.94	0.67
1:E:207:LYS:HB2	1:E:208:PRO:CD	2.10	0.67
1:F:449:ALA:HB3	1:F:450:PRO:HD3	1.75	0.67
1:J:151:SER:HB3	1:J:399:ALA:HA	1.76	0.67

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:K:208:PRO:HB2	1:K:212:ALA:CB	2.23	0.67
1:K:355:GLU:HG3	1:K:356:ALA:H	1.58	0.67
2:W:11:ILE:CG2	2:W:42:ALA:HB3	2.24	0.67
2:Z:93:ALA:HA	2:1:5:PRO:O	1.95	0.67
1:B:349:ILE:CD1	1:B:368:ARG:HD2	2.25	0.67
1:I:90:THR:N	4:I:602:BEF:F2	2.17	0.67
1:M:124:VAL:HG13	1:M:504:LEU:CD1	2.24	0.67
1:N:247:LEU:HG	1:N:249:ILE:CD1	2.23	0.67
2:V:12:VAL:HG23	2:V:39:GLU:O	1.94	0.67
2:V:49:LEU:CD1	2:W:50:GLU:HG3	2.24	0.67
2:Z:50:GLU:O	2:Z:50:GLU:HG2	1.92	0.67
2:2:17:VAL:HG23	2:2:18:GLU:HG3	1.77	0.67
1:F:6:VAL:CG1	1:F:521:VAL:HG22	2.19	0.67
1:M:305:ILE:CG2	1:M:306:GLY:H	2.08	0.67
2:X:16:GLU:HG2	2:X:19:THR:HB	1.75	0.67
1:A:234:LEU:HG	1:A:235:PRO:HD3	1.77	0.67
1:E:222:LEU:HD21	1:E:289:LEU:HB3	1.76	0.67
1:F:233:MET:HG3	1:F:234:LEU:CD2	2.14	0.67
1:L:449:ALA:HB3	1:L:450:PRO:HD3	1.77	0.67
1:N:278:ALA:HB3	1:N:285:ARG:CZ	2.24	0.67
2:Q:43:VAL:HG21	2:Q:57:LEU:HD11	1.77	0.67
2:X:91:ILE:HB	2:Y:6:LEU:HD22	1.75	0.67
1:A:165:ALA:O	1:A:169:VAL:HG22	1.95	0.67
1:A:349:ILE:HD13	1:A:368:ARG:HD2	1.76	0.67
1:I:227:ILE:HG23	1:I:230:ILE:HB	1.76	0.67
1:M:418:ALA:O	1:M:422:VAL:HG13	1.95	0.67
2:U:50:GLU:O	2:U:50:GLU:HG2	1.94	0.67
1:B:357:THR:HG21	1:B:361:ASP:HB2	1.77	0.67
1:D:357:THR:HB	1:D:361:ASP:CB	2.25	0.67
1:F:254:VAL:HG12	1:F:259:LEU:HG	1.75	0.67
1:G:277:LYS:HG3	1:G:278:ALA:N	2.10	0.67
1:H:338:GLU:O	1:H:342:ILE:HD13	1.94	0.67
1:M:246:PRO:HA	1:M:272:LYS:O	1.95	0.67
1:M:239:ALA:HB1	1:M:314:LEU:HD23	1.77	0.67
2:S:12:VAL:HG13	2:S:40:VAL:H	1.60	0.67
2:X:20:LYS:HB3	2:X:28:THR:OG1	1.94	0.67
2:2:79:ASP:O	2:2:80:ASN:C	2.32	0.67
1:J:357:THR:CB	1:J:361:ASP:HB2	2.25	0.67
1:K:419:LEU:HD11	1:K:500:THR:HG23	1.77	0.67
1:L:412:VAL:HG12	1:L:413:ALA:N	2.10	0.67
1:N:147:VAL:CG2	1:N:403:THR:HG22	2.24	0.67

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:H:307:MET:HG2	1:H:311:LYS:HZ3	1.59	0.67
1:K:5:ASP:HB2	1:K:524:LEU:HD23	1.76	0.67
1:N:279:PRO:HB2	1:N:288:MET:HE3	1.77	0.67
1:C:219:PHE:HA	1:C:318:GLY:O	1.95	0.67
1:C:323:VAL:HG12	1:C:332:ILE:HG12	1.77	0.67
1:C:429:LEU:HG	1:C:440:ILE:HD13	1.76	0.67
1:E:124:VAL:HG13	1:E:504:LEU:CD1	2.24	0.67
1:H:234:LEU:N	1:H:235:PRO:HD2	2.10	0.67
1:L:415:GLY:N	3:L:601:ADP:O2'	2.27	0.67
1:N:218:PRO:HG3	1:N:323:VAL:CG2	2.25	0.67
1:B:411:VAL:HG21	1:B:494:LEU:HD23	1.76	0.66
1:C:135:SER:HB2	1:C:497:THR:HG21	1.78	0.66
1:I:149:THR:HG21	1:I:156:GLU:HG2	1.76	0.66
1:M:231:ARG:HD3	1:M:261:THR:HG21	1.76	0.66
1:M:513:LEU:HD22	1:N:49:ILE:HG21	1.77	0.66
1:D:234:LEU:N	1:D:235:PRO:HD2	2.10	0.66
1:F:339:GLU:O	1:F:343:GLN:HB2	1.95	0.66
1:I:367:GLU:HG2	1:I:371:LYS:HE2	1.75	0.66
1:J:264:VAL:O	1:J:268:ARG:HG2	1.95	0.66
1:K:168:LYS:HG2	1:K:189:VAL:HG11	1.77	0.66
2:O:6:LEU:HD12	2:O:6:LEU:N	2.10	0.66
2:Q:15:LYS:HB3	2:Q:16:GLU:CD	2.16	0.66
1:A:216:GLU:O	1:A:245:LYS:HE2	1.96	0.66
1:F:220:ILE:HG13	1:F:296:THR:CG2	2.25	0.66
1:G:449:ALA:HB3	1:G:450:PRO:HD3	1.76	0.66
1:I:231:ARG:HD3	1:I:257:GLU:OE2	1.95	0.66
1:K:270:ILE:HD12	1:K:270:ILE:H	1.58	0.66
2:R:20:LYS:HB2	2:R:22:ALA:N	2.10	0.66
1:B:13:ARG:HD2	1:B:104:LEU:HD22	1.75	0.66
1:B:449:ALA:HB3	1:B:450:PRO:HD3	1.76	0.66
1:G:203:TYR:HB3	1:G:267:MET:CE	2.26	0.66
1:K:146:GLN:OE1	1:K:492:GLY:HA2	1.95	0.66
1:M:443:ALA:O	1:M:447:MET:HG3	1.96	0.66
2:T:53:GLU:CG	2:T:54:VAL:H	2.08	0.66
1:D:449:ALA:HB3	1:D:450:PRO:HD3	1.76	0.66
2:X:77:LYS:HD3	2:X:82:GLU:HA	1.77	0.66
2:Z:57:LEU:H	2:Z:57:LEU:HD12	1.59	0.66
1:A:291:ASP:HB3	1:A:372:LEU:HD21	1.78	0.66
1:A:388:GLU:HA	1:G:513:LEU:CD1	2.26	0.66
1:K:197:ARG:NH1	1:K:277:LYS:HD2	2.11	0.66
1:M:309:LEU:HB2	1:M:310:GLU:OE1	1.95	0.66

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:C:234:LEU:N	1:C:235:PRO:HD2	2.10	0.66
1:C:302:SER:OG	1:C:305:ILE:HD13	1.95	0.66
1:C:305:ILE:CG2	1:C:306:GLY:H	2.08	0.66
1:H:226:LYS:HG2	1:H:252:GLU:HG2	1.78	0.66
1:I:305:ILE:HG22	1:I:306:GLY:N	2.11	0.66
1:M:199:TYR:HE1	1:M:202:PRO:HA	1.59	0.66
2:W:26:VAL:O	2:W:27:LEU:HB2	1.96	0.66
1:B:147:VAL:HG12	1:B:494:LEU:HB2	1.76	0.66
1:G:359:ASP:O	1:G:363:GLU:HG2	1.95	0.66
1:J:254:VAL:HG12	1:J:259:LEU:HG	1.78	0.66
1:J:419:LEU:O	1:J:422:VAL:HG22	1.96	0.66
1:N:5:ASP:HB2	1:N:524:LEU:CD2	2.25	0.66
1:E:208:PRO:HG2	1:E:214:GLU:HG3	1.78	0.66
1:F:203:TYR:HD2	1:F:263:VAL:HG11	1.61	0.66
1:H:236:VAL:HG21	1:H:312:ALA:HB3	1.78	0.66
1:I:200:LEU:HD13	1:I:275:ALA:O	1.95	0.66
1:J:253:ASP:OD1	1:J:277:LYS:NZ	2.23	0.66
2:Q:50:GLU:CG	2:Q:51:ASN:H	2.07	0.66
2:S:36:THR:HG23	2:S:37:ARG:HG3	1.77	0.66
2:V:49:LEU:HD22	2:V:50:GLU:H	1.59	0.66
1:A:305:ILE:HG12	1:A:306:GLY:H	1.60	0.66
1:C:517:THR:HG23	1:D:39:VAL:HG23	1.77	0.66
1:E:102:GLU:OE1	1:E:445:ARG:NH1	2.28	0.66
1:J:349:ILE:O	1:J:353:ILE:HG13	1.96	0.66
1:J:31:LEU:HD23	1:J:453:GLN:HB3	1.77	0.66
1:M:231:ARG:NH2	1:M:257:GLU:HB3	2.10	0.66
1:F:226:LYS:HA	1:F:252:GLU:HB3	1.77	0.65
1:G:195:PHE:CE1	1:G:330:THR:HB	2.30	0.65
1:G:417:VAL:HA	1:G:420:ILE:CG2	2.26	0.65
1:H:225:LYS:HG3	1:H:303:GLU:HG3	1.78	0.65
1:J:278:ALA:CB	1:J:285:ARG:HD2	2.25	0.65
1:J:220:ILE:HG22	1:J:318:GLY:CA	2.26	0.65
1:M:268:ARG:HB2	1:M:269:GLY:HA2	1.78	0.65
1:H:241:ALA:HB3	2:V:25:ILE:CD1	2.26	0.65
1:D:254:VAL:HG12	1:D:259:LEU:HG	1.77	0.65
1:F:18:ARG:NH1	1:F:18:ARG:HB3	2.12	0.65
1:H:305:ILE:CG2	1:H:306:GLY:H	2.09	0.65
1:K:230:ILE:HG22	1:K:233:MET:H	1.61	0.65
2:U:57:LEU:H	2:U:57:LEU:HD12	1.61	0.65
2:Y:15:LYS:CD	2:Y:37:ARG:HB3	2.25	0.65
1:A:202:PRO:O	1:A:205:ILE:HG12	1.96	0.65

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:479:ASN:ND2	1:A:493:ILE:HD11	2.11	0.65
1:B:254:VAL:HG12	1:B:259:LEU:HG	1.79	0.65
1:C:357:THR:CG2	1:C:361:ASP:HB2	2.27	0.65
1:G:209:GLU:CG	1:G:210:THR:H	2.09	0.65
1:G:222:LEU:HD23	1:G:289:LEU:HD22	1.77	0.65
1:G:240:VAL:HG11	1:G:247:LEU:CD2	2.26	0.65
1:L:204:PHE:CE2	1:L:274:ALA:HA	2.31	0.65
1:N:233:MET:O	1:N:236:VAL:N	2.28	0.65
1:N:218:PRO:HG3	1:N:323:VAL:HG22	1.77	0.65
1:B:351:GLN:CG	1:C:210:THR:HG23	2.24	0.65
1:F:219:PHE:CE2	1:F:240:VAL:HG13	2.31	0.65
1:L:496:PRO:HG2	1:L:499:VAL:CG2	2.26	0.65
1:B:40:LEU:HD13	1:B:59:GLU:HG3	1.77	0.65
1:C:345:ARG:HA	1:C:348:GLN:HG2	1.78	0.65
1:G:339:GLU:O	1:G:343:GLN:HB2	1.96	0.65
1:H:197:ARG:HG2	1:H:277:LYS:CB	2.22	0.65
1:K:254:VAL:HG12	1:K:259:LEU:HG	1.77	0.65
1:M:40:LEU:HD13	1:M:59:GLU:HG3	1.78	0.65
2:T:20:LYS:HE2	2:T:22:ALA:CB	2.14	0.65
1:A:276:VAL:CG1	1:A:325:ILE:HG21	2.26	0.65
1:D:358:SER:O	1:D:362:ARG:HG3	1.96	0.65
1:I:31:LEU:HD23	1:I:453:GLN:HB3	1.77	0.65
1:L:367:GLU:O	1:L:371:LYS:HG3	1.97	0.65
1:M:228:SER:O	1:M:231:ARG:HB3	1.96	0.65
2:X:65:VAL:CG1	2:X:94:ILE:HG22	2.21	0.65
1:A:199:TYR:HA	1:A:276:VAL:HG12	1.79	0.65
1:A:39:VAL:HG23	1:G:517:THR:HG23	1.79	0.65
1:C:448:GLU:O	1:C:452:ARG:HG2	1.95	0.65
1:D:443:ALA:O	1:D:447:MET:HG3	1.97	0.65
1:F:207:LYS:HB3	1:F:208:PRO:HD3	1.79	0.65
1:L:218:PRO:HG2	1:L:220:ILE:CD1	2.26	0.65
1:M:291:ASP:OD2	1:M:368:ARG:HD3	1.96	0.65
1:N:231:ARG:HD3	1:N:309:LEU:HB3	1.77	0.65
2:Z:95:VAL:CA	2:1:3:ILE:HG22	2.23	0.65
1:A:247:LEU:HG	1:A:273:VAL:CG1	2.26	0.65
1:A:252:GLU:HA	1:A:285:ARG:HH11	1.61	0.65
1:C:224:ASP:OD1	1:C:286:LYS:HD3	1.97	0.65
1:E:278:ALA:CB	1:E:285:ARG:HD2	2.26	0.65
1:F:228:SER:CB	1:F:255:GLU:HB2	2.27	0.65
1:K:252:GLU:HG3	1:K:285:ARG:NH1	2.10	0.65
2:R:12:VAL:HG11	2:R:40:VAL:HG22	1.79	0.65

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:245:LYS:CB	1:A:246:PRO:HD3	2.26	0.65
1:C:302:SER:CB	1:C:305:ILE:HD13	2.27	0.65
1:D:218:PRO:HA	1:D:246:PRO:HG2	1.77	0.65
1:G:325:ILE:HG12	1:G:330:THR:HG23	1.79	0.65
1:I:197:ARG:HH11	1:I:277:LYS:HD2	1.62	0.65
1:I:228:SER:HB3	1:I:255:GLU:CB	2.27	0.65
1:I:27:VAL:HG12	1:I:90:THR:HG23	1.79	0.65
1:M:233:MET:HB3	1:M:236:VAL:CG2	2.27	0.65
2:U:16:GLU:N	2:U:16:GLU:OE1	2.30	0.65
2:W:40:VAL:HB	2:W:63:ASP:H	1.62	0.65
1:B:136:VAL:CG2	1:B:137:PRO:HD2	2.26	0.65
1:B:207:LYS:HB3	1:B:208:PRO:HD3	1.79	0.65
1:B:219:PHE:HA	1:B:318:GLY:O	1.96	0.65
1:C:304:GLU:C	1:C:305:ILE:HD12	2.16	0.65
1:K:305:ILE:CG2	1:K:306:GLY:H	2.10	0.65
2:T:20:LYS:CE	2:T:22:ALA:HB3	2.15	0.65
1:A:320:ALA:HA	1:A:335:GLY:HA2	1.78	0.64
1:G:355:GLU:HG3	1:G:356:ALA:H	1.62	0.64
1:J:196:ASP:O	1:J:197:ARG:HG2	1.97	0.64
1:J:420:ILE:HD11	1:J:469:VAL:HG12	1.79	0.64
1:K:219:PHE:CZ	1:K:245:LYS:HG3	2.31	0.64
1:N:420:ILE:CG2	1:N:448:GLU:HA	2.27	0.64
2:V:86:MET:HG3	2:V:91:ILE:HD11	1.77	0.64
1:A:241:ALA:CB	1:A:271:VAL:HG12	2.28	0.64
1:D:200:LEU:CD1	1:D:276:VAL:HA	2.26	0.64
1:F:288:MET:HA	1:F:291:ASP:OD2	1.97	0.64
1:G:124:VAL:HG13	1:G:504:LEU:CD1	2.27	0.64
1:H:350:ARG:HD3	1:H:353:ILE:CD1	2.23	0.64
2:O:73:VAL:HG12	2:O:86:MET:HB3	1.78	0.64
2:R:20:LYS:CA	2:R:21:SER:HB3	2.15	0.64
2:W:64:ILE:CG2	2:W:95:VAL:HB	2.27	0.64
1:A:416:GLY:O	1:A:420:ILE:HD12	1.97	0.64
1:B:305:ILE:CG2	1:B:306:GLY:H	2.09	0.64
1:C:5:ASP:HB2	1:C:524:LEU:HD23	1.80	0.64
1:G:161:LEU:HD23	1:G:379:ILE:HD11	1.79	0.64
1:I:349:ILE:HG23	1:I:365:LEU:HD11	1.79	0.64
1:I:102:GLU:OE1	1:I:445:ARG:NH1	2.31	0.64
1:L:124:VAL:HG21	1:L:508:ALA:CB	2.27	0.64
1:L:364:LYS:O	1:L:367:GLU:HB3	1.97	0.64
2:R:3:ILE:HD11	2:R:11:ILE:CD1	2.23	0.64
2:W:8:ASP:O	2:W:57:LEU:HD11	1.97	0.64

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:310:GLU:CD	1:A:310:GLU:H	2.01	0.64
1:J:231:ARG:CD	1:J:257:GLU:OE1	2.43	0.64
1:B:27:VAL:HG12	1:B:90:THR:HG23	1.79	0.64
2:S:67:PHE:CA	2:S:92:LEU:HD13	2.24	0.64
1:C:217:SER:HA	1:C:320:ALA:O	1.98	0.64
1:G:230:ILE:HG22	1:G:232:GLU:HG2	1.77	0.64
1:G:240:VAL:HG11	1:G:247:LEU:HD11	1.78	0.64
1:J:301:ILE:HG12	1:J:307:MET:CE	2.28	0.64
1:K:199:TYR:CE1	1:K:202:PRO:HA	2.33	0.64
1:M:74:VAL:O	1:M:77:VAL:HG12	1.97	0.64
2:Q:78:ILE:HG13	2:Q:79:ASP:OD2	1.97	0.64
2:R:78:ILE:HG13	2:R:79:ASP:N	2.12	0.64
1:B:34:LYS:NZ	1:B:483:GLU:OE2	2.30	0.64
1:E:302:SER:O	1:E:305:ILE:HG22	1.97	0.64
1:F:23:LEU:HD23	1:F:60:ILE:HB	1.80	0.64
1:F:169:VAL:HB	1:F:375:GLY:O	1.97	0.64
1:H:177:VAL:HG11	1:H:397:GLU:CG	2.28	0.64
1:H:208:PRO:HB2	1:H:212:ALA:HB3	1.80	0.64
1:I:450:PRO:O	1:I:454:ILE:HG13	1.98	0.64
1:L:510:VAL:O	1:L:514:MET:HB2	1.97	0.64
1:N:201:SER:HB3	1:N:204:PHE:CZ	2.32	0.64
2:2:17:VAL:HA	2:2:35:SER:HB2	1.79	0.64
1:C:69:MET:SD	1:D:41:ASP:HB2	2.38	0.64
1:E:77:VAL:HG21	1:E:507:ALA:HA	1.80	0.64
1:F:349:ILE:HG21	1:F:369:VAL:HG22	1.80	0.64
1:G:284:ARG:HD3	1:G:364:LYS:HE2	1.79	0.64
1:I:311:LYS:HB3	1:I:311:LYS:NZ	2.12	0.64
1:J:230:ILE:HG21	1:J:309:LEU:HD13	1.80	0.64
1:L:282:GLY:HA2	1:L:285:ARG:HG2	1.80	0.64
2:T:64:ILE:HD12	2:T:64:ILE:H	1.63	0.64
2:X:50:GLU:O	2:X:50:GLU:CD	2.36	0.64
1:B:77:VAL:HB	1:B:510:VAL:HG21	1.80	0.64
1:D:177:VAL:CG2	1:D:393:LYS:HG3	2.27	0.64
1:E:196:ASP:O	1:E:197:ARG:HG2	1.98	0.64
1:H:77:VAL:HB	1:H:510:VAL:HG21	1.79	0.64
1:K:124:VAL:HG21	1:K:508:ALA:CB	2.26	0.64
1:M:197:ARG:CD	1:M:277:LYS:HB2	2.28	0.64
1:M:455:VAL:HG13	1:M:460:GLU:HB2	1.79	0.64
2:T:53:GLU:CD	2:T:54:VAL:H	2.00	0.64
2:T:92:LEU:CD2	2:U:85:ILE:HD13	2.27	0.64
1:A:147:VAL:HG12	1:A:494:LEU:CB	2.28	0.64

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:B:263:VAL:O	1:B:267:MET:HG2	1.97	0.64
1:I:291:ASP:OD1	1:I:345:ARG:NH2	2.28	0.64
1:L:320:ALA:HA	1:L:335:GLY:HA2	1.80	0.64
1:M:293:ALA:O	1:M:298:GLY:N	2.29	0.64
2:U:51:ASN:C	2:U:53:GLU:H	2.01	0.64
1:A:268:ARG:N	1:A:269:GLY:HA2	2.13	0.63
1:C:225:LYS:CG	1:C:303:GLU:HG3	2.28	0.63
1:C:174:VAL:CG1	1:C:376:VAL:HG13	2.27	0.63
1:E:136:VAL:CG2	1:E:137:PRO:HD2	2.28	0.63
1:I:417:VAL:HA	1:I:420:ILE:HG22	1.79	0.63
1:J:124:VAL:HG13	1:J:504:LEU:CD1	2.28	0.63
1:K:420:ILE:HD12	1:K:451:LEU:HD13	1.79	0.63
1:L:232:GLU:HB2	1:L:234:LEU:HD13	1.80	0.63
2:Q:40:VAL:HG11	2:Q:61:VAL:HA	1.79	0.63
1:H:406:ALA:HB1	1:H:411:VAL:HG12	1.79	0.63
1:J:449:ALA:HB3	1:J:450:PRO:HD3	1.80	0.63
1:J:496:PRO:HG2	1:J:499:VAL:HG13	1.80	0.63
1:K:221:LEU:HD13	1:K:317:LEU:CD2	2.27	0.63
1:M:225:LYS:HD3	1:M:303:GLU:OE1	1.97	0.63
2:T:7:HIS:HB2	2:T:46:GLY:O	1.98	0.63
1:E:205:ILE:HA	1:E:213:VAL:CG2	2.23	0.63
1:E:219:PHE:CB	1:E:317:LEU:HD13	2.29	0.63
1:I:397:GLU:O	1:I:401:HIS:ND1	2.30	0.63
1:J:321:LYS:HD2	1:J:334:ASP:HB2	1.78	0.63
2:R:6:LEU:HD23	2:R:7:HIS:ND1	1.97	0.63
1:A:276:VAL:HG11	1:A:325:ILE:CG2	2.27	0.63
1:B:245:LYS:HB2	1:B:246:PRO:CD	2.15	0.63
1:D:313:THR:CG2	1:D:315:GLU:HG2	2.28	0.63
3:G:601:ADP:O3B	4:G:602:BEF:F1	2.06	0.63
1:J:202:PRO:O	1:J:205:ILE:HG13	1.98	0.63
1:K:168:LYS:HG2	1:K:189:VAL:HG21	1.80	0.63
1:L:350:ARG:HD3	1:L:353:ILE:CD1	2.28	0.63
1:L:85:ALA:HB1	1:L:499:VAL:HG22	1.79	0.63
1:N:168:LYS:HD3	1:N:189:VAL:HG11	1.80	0.63
2:V:11:ILE:O	2:V:11:ILE:HD12	1.98	0.63
1:B:349:ILE:HD11	1:B:368:ARG:HD2	1.78	0.63
1:D:177:VAL:HG22	1:D:393:LYS:HG3	1.80	0.63
1:E:234:LEU:HD23	1:E:235:PRO:N	2.12	0.63
1:E:365:LEU:O	1:E:369:VAL:HG23	1.99	0.63
1:K:218:PRO:HA	1:K:246:PRO:HG2	1.81	0.63
1:L:299:THR:N	1:L:316:ASP:O	2.23	0.63

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:N:124:VAL:O	1:N:128:VAL:HG23	1.99	0.63
1:N:405:ALA:HB1	1:N:498:LYS:HB3	1.81	0.63
2:U:51:ASN:C	2:U:53:GLU:N	2.49	0.63
2:Z:47:ARG:CD	2:Z:49:LEU:HD12	2.28	0.63
1:A:308:GLU:HG2	1:A:311:LYS:HB2	1.81	0.63
1:I:205:ILE:HG21	1:I:211:GLY:HA2	1.80	0.63
1:J:301:ILE:HA	1:J:307:MET:CE	2.29	0.63
1:L:455:VAL:CG1	1:L:460:GLU:HB2	2.29	0.63
1:M:279:PRO:HG2	1:M:285:ARG:CA	2.27	0.63
2:U:48:ILE:HG13	2:U:53:GLU:HA	1.81	0.63
1:B:128:VAL:HG12	1:B:132:LYS:HE2	1.81	0.63
1:E:234:LEU:O	1:E:235:PRO:C	2.37	0.63
1:H:234:LEU:H	1:H:234:LEU:HD12	1.64	0.63
1:I:288:MET:HG3	1:I:368:ARG:CZ	2.28	0.63
1:C:37:ASN:OD1	1:C:51:LYS:HB2	1.99	0.63
1:C:77:VAL:HG21	1:C:507:ALA:HA	1.81	0.63
1:D:357:THR:HB	1:D:361:ASP:HB2	1.81	0.63
1:H:89:THR:N	4:H:602:BEF:F3	2.22	0.63
1:I:18:ARG:HB3	1:I:18:ARG:NH1	2.14	0.63
1:A:271:VAL:O	1:A:273:VAL:HG23	1.98	0.63
1:A:308:GLU:CD	1:A:311:LYS:HB2	2.20	0.63
1:E:230:ILE:HG23	1:E:310:GLU:OE2	1.99	0.63
1:E:226:LYS:HG2	1:E:252:GLU:OE1	1.99	0.63
1:F:209:GLU:HG3	1:F:210:THR:H	1.63	0.63
1:G:350:ARG:HH11	1:G:353:ILE:HD11	1.63	0.63
2:O:94:ILE:HD11	2:P:4:ARG:HB2	1.81	0.63
2:U:48:ILE:HA	2:U:53:GLU:O	1.99	0.63
1:H:349:ILE:O	1:H:353:ILE:HG13	1.98	0.62
1:I:392:LYS:O	1:I:396:VAL:HG23	1.99	0.62
1:N:6:VAL:CG1	1:N:521:VAL:HG22	2.29	0.62
1:B:293:ALA:O	1:B:298:GLY:N	2.32	0.62
1:C:383:ALA:HB3	1:C:389:MET:SD	2.39	0.62
1:D:496:PRO:HG2	1:D:499:VAL:CG1	2.27	0.62
1:K:228:SER:OG	1:K:255:GLU:OE2	2.17	0.62
1:L:247:LEU:HD13	1:L:248:LEU:N	2.14	0.62
2:Q:69:ASP:CB	2:Q:73:VAL:HG11	2.29	0.62
2:R:8:ASP:OD1	2:R:9:ARG:HG3	1.98	0.62
2:W:95:VAL:HA	2:X:3:ILE:HG22	1.80	0.62
2:Y:15:LYS:HD3	2:Y:37:ARG:HB3	1.81	0.62
2:Y:58:ASP:O	2:Y:58:ASP:OD1	2.17	0.62
1:A:496:PRO:HG2	1:A:499:VAL:CG1	2.29	0.62

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:B:322:ARG:HB3	1:B:333:ILE:HD12	1.81	0.62
1:D:237:LEU:HD21	2:R:27:LEU:HD21	1.80	0.62
3:F:601:ADP:O3B	4:F:602:BEF:F2	2.08	0.62
1:I:200:LEU:CD1	1:I:276:VAL:HA	2.30	0.62
1:I:302:SER:HB2	1:I:305:ILE:HD13	1.82	0.62
1:J:5:ASP:HB2	1:J:524:LEU:CD2	2.29	0.62
1:L:77:VAL:HB	1:L:510:VAL:HG21	1.81	0.62
2:Y:20:LYS:HE2	2:Y:22:ALA:HB3	1.82	0.62
1:C:124:VAL:HG21	1:C:508:ALA:CB	2.29	0.62
1:G:10:ASN:O	1:G:14:VAL:HG23	1.99	0.62
1:G:77:VAL:HB	1:G:510:VAL:HG21	1.79	0.62
1:G:88:GLY:N	3:G:601:ADP:O1B	2.32	0.62
1:K:448:GLU:O	1:K:452:ARG:HG2	1.99	0.62
1:A:5:ASP:HB2	1:A:524:LEU:HD21	1.82	0.62
1:B:304:GLU:O	1:B:305:ILE:HD13	2.00	0.62
1:F:215:LEU:HB2	1:F:323:VAL:CG2	2.30	0.62
1:G:345:ARG:O	1:G:349:ILE:HG13	2.00	0.62
1:I:280:GLY:O	1:I:285:ARG:NE	2.24	0.62
1:I:194:GLN:NE2	1:I:329:THR:HG21	2.14	0.62
1:L:433:ASN:HD21	1:L:435:ASP:HB2	1.64	0.62
2:O:53:GLU:HG3	2:O:54:VAL:H	1.62	0.62
2:1:50:GLU:O	2:1:51:ASN:CG	2.37	0.62
1:A:418:ALA:O	1:A:422:VAL:HG13	2.00	0.62
1:D:175:ILE:HA	1:D:377:ALA:O	2.00	0.62
1:E:285:ARG:O	1:E:289:LEU:HG	1.99	0.62
1:H:304:GLU:C	1:H:305:ILE:HD12	2.20	0.62
1:L:6:VAL:CG1	1:L:521:VAL:HG22	2.25	0.62
1:N:284:ARG:HG2	1:N:288:MET:CE	2.29	0.62
1:N:443:ALA:O	1:N:447:MET:HG3	1.99	0.62
1:F:270:ILE:HD13	2:T:26:VAL:HA	1.82	0.62
1:A:245:LYS:CB	1:A:246:PRO:CD	2.67	0.62
1:D:177:VAL:HG11	1:D:397:GLU:HG2	1.81	0.62
1:E:302:SER:HB2	1:E:305:ILE:CG2	2.30	0.62
1:E:419:LEU:HD11	1:E:500:THR:HG23	1.81	0.62
1:G:367:GLU:HG2	1:G:371:LYS:HE2	1.80	0.62
1:H:276:VAL:HG11	1:H:325:ILE:HD13	1.82	0.62
1:M:409:GLU:OE2	1:M:501:ARG:NH2	2.32	0.62
1:A:365:LEU:O	1:A:369:VAL:HG23	2.00	0.62
1:F:443:ALA:O	1:F:447:MET:HG3	2.00	0.62
1:F:31:LEU:HD23	1:F:453:GLN:HB3	1.82	0.62
1:I:69:MET:SD	1:J:41:ASP:HB2	2.40	0.62

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:K:280:GLY:O	1:K:285:ARG:HB3	1.99	0.62
1:K:62:LEU:HD23	1:K:67:GLU:HB3	1.81	0.62
1:B:198:GLY:O	1:B:276:VAL:HG13	1.99	0.62
1:C:231:ARG:NH1	1:C:234:LEU:HD21	2.14	0.62
1:E:136:VAL:HG22	1:E:137:PRO:HD2	1.82	0.62
1:E:225:LYS:HG2	1:E:226:LYS:N	2.15	0.62
1:E:291:ASP:OD2	1:E:368:ARG:HD3	2.00	0.62
1:F:513:LEU:HD11	1:G:388:GLU:HA	1.80	0.62
1:J:89:THR:N	4:J:602:BEF:F1	2.19	0.62
1:E:222:LEU:HD22	1:E:289:LEU:HD13	1.81	0.62
1:H:232:GLU:HG3	1:H:233:MET:N	2.14	0.62
1:I:224:ASP:OD1	1:I:286:LYS:HE3	1.99	0.62
2:S:14:ARG:HG2	2:S:35:SER:HB3	1.80	0.62
1:B:429:LEU:HG	1:B:440:ILE:HD13	1.81	0.61
1:C:142:LYS:O	1:C:146:GLN:HG3	2.00	0.61
1:D:479:ASN:ND2	1:D:493:ILE:HD11	2.15	0.61
1:E:247:LEU:HB2	1:E:273:VAL:HG13	1.81	0.61
1:G:231:ARG:HG2	1:G:257:GLU:OE1	2.00	0.61
1:I:363:GLU:OE1	1:I:364:LYS:HG2	2.00	0.61
1:K:296:THR:OG1	1:K:318:GLY:HA3	2.00	0.61
1:K:381:VAL:CG2	1:K:396:VAL:HG21	2.30	0.61
1:M:233:MET:HB2	1:M:237:LEU:CD2	2.30	0.61
1:M:302:SER:CB	1:M:305:ILE:HD13	2.30	0.61
1:N:222:LEU:HD22	1:N:293:ALA:HB2	1.82	0.61
2:T:55:LYS:HD2	2:T:56:PRO:HD2	1.80	0.61
1:H:234:LEU:HD23	2:V:22:ALA:HB1	1.82	0.61
2:W:70:GLY:O	2:W:73:VAL:HG12	2.00	0.61
2:1:66:ILE:HG21	2:2:76:GLU:HG3	1.82	0.61
1:A:388:GLU:HA	1:G:513:LEU:HD11	1.81	0.61
1:B:142:LYS:O	1:B:146:GLN:HG3	2.00	0.61
1:B:488:MET:HE1	1:B:493:ILE:HD12	1.81	0.61
1:D:142:LYS:O	1:D:146:GLN:HG3	1.99	0.61
1:D:241:ALA:HB2	1:D:271:VAL:HG11	1.83	0.61
1:E:234:LEU:O	1:E:237:LEU:N	2.33	0.61
1:G:172:GLU:CD	1:G:172:GLU:N	2.53	0.61
1:H:223:ALA:HA	1:H:301:ILE:HB	1.81	0.61
1:I:225:LYS:HG3	1:I:226:LYS:H	1.65	0.61
1:K:228:SER:CB	1:K:255:GLU:HB2	2.30	0.61
1:N:346:VAL:HG22	1:N:372:LEU:CD1	2.31	0.61
2:T:94:ILE:HD11	2:U:4:ARG:HD2	1.81	0.61
2:V:74:LYS:NZ	2:2:68:ASN:HB3	2.15	0.61

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:193:MET:HB3	1:A:332:ILE:HB	1.82	0.61
1:B:357:THR:CG2	1:B:361:ASP:HB2	2.30	0.61
1:C:48:THR:HG22	1:C:390:LYS:NZ	2.14	0.61
1:E:364:LYS:HD2	1:E:367:GLU:HB3	1.82	0.61
1:G:364:LYS:O	1:G:367:GLU:HB3	2.01	0.61
1:G:74:VAL:O	1:G:77:VAL:HG12	1.99	0.61
1:N:381:VAL:CG2	1:N:396:VAL:HG21	2.30	0.61
2:Q:3:ILE:HD11	2:Q:78:ILE:HD13	1.83	0.61
2:U:12:VAL:HG23	2:U:40:VAL:HA	1.80	0.61
1:B:443:ALA:O	1:B:447:MET:HG3	1.99	0.61
1:C:248:LEU:HD23	1:C:249:ILE:N	2.16	0.61
1:J:215:LEU:HB2	1:J:323:VAL:HG22	1.83	0.61
2:P:5:PRO:HD3	2:P:11:ILE:HD12	1.82	0.61
2:W:49:LEU:HB2	2:W:52:GLY:CA	2.29	0.61
1:C:218:PRO:HA	1:C:246:PRO:O	1.99	0.61
1:E:172:GLU:HB3	1:E:366:GLN:OE1	2.01	0.61
1:F:225:LYS:HG2	1:F:309:LEU:CD2	2.26	0.61
1:I:280:GLY:H	1:I:285:ARG:HB3	1.65	0.61
1:I:193:MET:CE	1:I:292:ILE:HG12	2.31	0.61
1:J:218:PRO:HD2	1:J:320:ALA:O	2.00	0.61
1:K:518:GLU:CB	1:L:29:VAL:HG11	2.28	0.61
1:M:34:LYS:HD2	1:M:458:CYS:SG	2.40	0.61
2:O:49:LEU:HD13	2:O:50:GLU:H	1.63	0.61
2:O:97:ALA:HB2	2:P:1:MET:N	2.15	0.61
1:B:234:LEU:N	1:B:235:PRO:HD2	2.15	0.61
1:E:165:ALA:O	1:E:169:VAL:HG22	2.01	0.61
1:H:358:SER:HA	1:H:362:ARG:CZ	2.30	0.61
1:H:48:THR:HG22	1:H:390:LYS:NZ	2.15	0.61
1:L:218:PRO:HB2	1:L:219:PHE:C	2.19	0.61
1:N:252:GLU:HA	1:N:285:ARG:NH1	2.15	0.61
1:A:195:PHE:O	1:A:329:THR:HG23	2.01	0.61
1:B:475:ASN:O	1:B:488:MET:HG2	2.00	0.61
1:C:345:ARG:O	1:C:349:ILE:HG13	2.00	0.61
1:D:218:PRO:HD2	1:D:320:ALA:O	2.00	0.61
1:E:225:LYS:NZ	1:E:230:ILE:HD11	2.16	0.61
1:D:517:THR:HG23	1:E:39:VAL:HG23	1.83	0.61
1:F:80:LYS:HD2	1:F:506:TYR:CZ	2.36	0.61
1:H:227:ILE:HA	1:H:230:ILE:CG2	2.30	0.61
1:I:225:LYS:HD2	1:I:252:GLU:OE2	2.01	0.61
1:I:20:VAL:HG13	1:I:74:VAL:HG11	1.82	0.61
1:I:517:THR:HG23	1:J:39:VAL:HG23	1.81	0.61

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:J:416:GLY:O	1:J:420:ILE:HG23	2.00	0.61
1:M:354:GLU:N	1:M:354:GLU:OE2	2.33	0.61
2:Y:40:VAL:HG13	2:Y:62:GLY:N	2.12	0.61
2:1:58:ASP:O	2:1:58:ASP:OD1	2.18	0.61
1:D:381:VAL:HG11	1:D:392:LYS:HB3	1.82	0.61
1:F:175:ILE:HA	1:F:377:ALA:O	2.00	0.61
1:H:258:ALA:O	1:H:262:LEU:HB2	2.00	0.61
1:H:218:PRO:HD2	1:H:320:ALA:O	2.01	0.61
1:I:227:ILE:HA	1:I:230:ILE:HD13	1.83	0.61
1:K:231:ARG:HD2	1:K:261:THR:HG21	1.82	0.61
1:L:443:ALA:O	1:L:447:MET:HG3	2.00	0.61
1:N:325:ILE:CG1	1:N:330:THR:HG23	2.29	0.61
2:R:12:VAL:CG1	2:R:40:VAL:HG22	2.30	0.61
2:V:57:LEU:HD12	2:V:59:VAL:HG13	1.83	0.61
1:G:215:LEU:HB3	1:G:246:PRO:HB2	1.83	0.61
1:H:143:ALA:O	1:H:147:VAL:HG13	2.00	0.61
1:J:207:LYS:HB3	1:J:208:PRO:HD3	1.83	0.61
1:K:92:ALA:HB2	1:K:503:ALA:HB1	1.82	0.61
2:T:49:LEU:HD23	2:T:50:GLU:HG3	1.83	0.61
1:B:278:ALA:HB3	1:B:285:ARG:HD2	1.80	0.61
1:G:234:LEU:HD12	1:G:234:LEU:H	1.64	0.61
1:H:136:VAL:HG23	1:H:137:PRO:HD2	1.81	0.61
1:J:350:ARG:HD3	1:J:353:ILE:CD1	2.31	0.61
1:L:218:PRO:CA	1:L:219:PHE:HB2	2.31	0.61
1:L:280:GLY:H	1:L:285:ARG:HB3	1.66	0.61
1:L:131:LEU:HD23	1:L:422:VAL:HG11	1.83	0.61
1:M:13:ARG:HD2	1:M:104:LEU:HD22	1.83	0.61
2:Y:7:HIS:O	2:Y:7:HIS:HD2	1.84	0.61
1:A:40:LEU:HD21	1:A:56:VAL:HG22	1.83	0.60
1:I:238:GLU:HG2	2:W:27:LEU:HD11	1.82	0.60
1:J:225:LYS:HG2	1:J:226:LYS:N	2.16	0.60
1:L:240:VAL:HG11	1:L:247:LEU:HB2	1.81	0.60
2:R:11:ILE:HG23	2:R:42:ALA:H	1.64	0.60
2:Y:20:LYS:HB2	2:Y:28:THR:HG23	1.83	0.60
1:B:222:LEU:CD2	1:B:300:VAL:HG22	2.24	0.60
1:C:233:MET:C	1:C:235:PRO:HD2	2.20	0.60
1:E:405:ALA:HB1	1:E:498:LYS:HB3	1.82	0.60
1:J:411:VAL:HG21	1:J:494:LEU:HD23	1.81	0.60
1:K:293:ALA:O	1:K:298:GLY:N	2.34	0.60
2:S:73:VAL:HA	2:S:85:ILE:O	2.01	0.60
1:D:237:LEU:CD2	2:R:27:LEU:HD21	2.31	0.60

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:D:360:TYR:HA	1:D:363:GLU:CG	2.17	0.60
1:G:418:ALA:O	1:G:422:VAL:HG13	2.01	0.60
1:G:134:LEU:HD11	1:G:425:LYS:NZ	2.16	0.60
1:H:313:THR:CG2	1:H:315:GLU:HG3	2.32	0.60
1:J:204:PHE:CD2	1:J:274:ALA:HA	2.36	0.60
1:B:448:GLU:O	1:B:452:ARG:HG2	2.01	0.60
1:E:10:ASN:O	1:E:14:VAL:HG23	2.01	0.60
1:F:200:LEU:HD22	1:F:254:VAL:HB	1.83	0.60
1:H:307:MET:HG2	1:H:311:LYS:NZ	2.17	0.60
1:L:205:ILE:CD1	1:L:211:GLY:HA2	2.30	0.60
1:N:284:ARG:HG2	1:N:288:MET:HE2	1.83	0.60
2:R:25:ILE:O	2:R:25:ILE:HD12	2.01	0.60
2:R:25:ILE:HG22	2:R:26:VAL:HG22	1.83	0.60
1:B:321:LYS:CB	1:B:334:ASP:HB2	2.31	0.60
1:C:183:LEU:HA	1:C:382:GLY:O	2.00	0.60
1:E:314:LEU:HD12	1:E:315:GLU:N	2.17	0.60
1:G:230:ILE:HG21	1:G:309:LEU:CD1	2.31	0.60
1:J:363:GLU:HG3	1:J:364:LYS:H	1.65	0.60
1:K:33:PRO:HA	1:K:153:ASN:OD1	2.01	0.60
1:L:513:LEU:HD22	1:M:49:ILE:HG21	1.83	0.60
1:N:247:LEU:CG	1:N:249:ILE:HD11	2.28	0.60
2:O:5:PRO:C	2:O:6:LEU:HD12	2.22	0.60
1:A:18:ARG:NH1	1:A:18:ARG:HB3	2.17	0.60
1:A:308:GLU:CG	1:A:311:LYS:HB2	2.31	0.60
1:E:177:VAL:HG11	1:E:397:GLU:CG	2.31	0.60
1:F:361:ASP:O	1:F:365:LEU:HB2	2.01	0.60
1:G:77:VAL:CG2	1:G:507:ALA:HA	2.29	0.60
1:H:232:GLU:C	1:H:234:LEU:HD12	2.22	0.60
1:J:157:THR:O	1:J:161:LEU:HD13	2.01	0.60
1:L:230:ILE:HG21	1:L:309:LEU:CD2	2.27	0.60
2:O:97:ALA:HB2	2:P:1:MET:HB2	1.82	0.60
2:U:40:VAL:HG12	2:U:41:LEU:N	2.16	0.60
2:Y:14:ARG:HD2	2:Y:84:LEU:HD11	1.84	0.60
2:Y:16:GLU:OE1	2:Y:16:GLU:N	2.28	0.60
1:B:305:ILE:HD11	1:C:203:TYR:HE2	1.66	0.60
1:C:200:LEU:HB3	1:C:259:LEU:HD11	1.83	0.60
1:C:245:LYS:HE2	1:C:246:PRO:HD2	1.83	0.60
1:C:40:LEU:HD13	1:C:59:GLU:HG3	1.82	0.60
1:C:5:ASP:HB2	1:C:524:LEU:CD2	2.32	0.60
1:D:18:ARG:HB3	1:D:18:ARG:NH1	2.17	0.60
1:G:124:VAL:HG21	1:G:508:ALA:CB	2.32	0.60

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:H:199:TYR:CE1	1:H:202:PRO:HA	2.37	0.60
1:I:207:LYS:HB3	1:I:208:PRO:HD3	1.84	0.60
1:K:168:LYS:CG	1:K:189:VAL:HG11	2.32	0.60
2:S:12:VAL:HG13	2:S:40:VAL:CA	2.32	0.60
2:U:14:ARG:CZ	2:U:34:LYS:HD3	2.31	0.60
2:2:69:ASP:OD1	2:2:69:ASP:N	2.34	0.60
1:B:218:PRO:HA	1:B:246:PRO:O	2.02	0.60
1:C:314:LEU:HD12	1:C:315:GLU:N	2.17	0.60
1:D:305:ILE:CG2	1:D:306:GLY:H	2.12	0.60
1:D:5:ASP:HB2	1:D:524:LEU:HD23	1.83	0.60
1:E:361:ASP:OD1	1:E:362:ARG:N	2.35	0.60
1:H:345:ARG:O	1:H:348:GLN:HG3	2.00	0.60
1:J:443:ALA:O	1:J:447:MET:HG3	2.01	0.60
1:N:346:VAL:HG12	1:N:350:ARG:HH12	1.66	0.60
2:O:96:GLU:O	2:P:1:MET:CB	2.49	0.60
2:Z:15:LYS:HB2	2:Z:37:ARG:O	2.02	0.60
1:A:455:VAL:HG13	1:A:460:GLU:HB2	1.83	0.60
1:B:168:LYS:HD3	1:B:189:VAL:HG21	1.84	0.60
1:G:236:VAL:HG22	1:G:312:ALA:HB3	1.82	0.60
1:N:236:VAL:HG22	1:N:312:ALA:CB	2.32	0.60
1:N:228:SER:HA	1:N:258:ALA:CB	2.31	0.60
2:Z:10:VAL:HG23	2:Z:42:ALA:O	2.02	0.60
2:1:95:VAL:HG22	2:2:3:ILE:CG2	2.30	0.60
1:C:18:ARG:HB3	1:C:18:ARG:NH1	2.17	0.60
1:D:77:VAL:HG21	1:D:507:ALA:HA	1.84	0.60
1:F:355:GLU:HG3	1:F:356:ALA:H	1.65	0.60
1:G:165:ALA:O	1:G:169:VAL:HG22	2.01	0.60
1:J:164:GLU:O	1:J:168:LYS:HG2	2.01	0.60
1:J:247:LEU:HD13	1:J:248:LEU:N	2.17	0.60
1:J:174:VAL:HG13	1:J:376:VAL:HG13	1.84	0.60
1:J:405:ALA:HB1	1:J:498:LYS:HB3	1.83	0.60
2:Q:25:ILE:HG13	2:Q:26:VAL:HG23	1.83	0.60
2:Q:77:LYS:NZ	2:Q:80:ASN:HA	2.17	0.60
1:D:198:GLY:O	1:D:276:VAL:HG23	2.02	0.59
1:I:205:ILE:HA	1:I:213:VAL:CG2	2.28	0.59
1:H:69:MET:SD	1:I:41:ASP:HB2	2.42	0.59
1:J:40:LEU:HD13	1:J:59:GLU:HG3	1.82	0.59
1:J:77:VAL:HG21	1:J:507:ALA:HA	1.84	0.59
1:L:208:PRO:CA	1:L:209:GLU:OE1	2.50	0.59
2:Y:5:PRO:HD3	2:Y:42:ALA:CB	2.32	0.59
1:C:199:TYR:CE1	1:C:202:PRO:HA	2.37	0.59

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:E:205:ILE:CD1	1:E:211:GLY:HA2	2.28	0.59
1:E:305:ILE:HG12	1:E:306:GLY:H	1.67	0.59
1:G:171:LYS:HB2	1:G:172:GLU:OE1	2.02	0.59
1:G:219:PHE:CZ	1:G:245:LYS:HD2	2.37	0.59
1:G:291:ASP:OD1	1:G:292:ILE:N	2.35	0.59
1:I:124:VAL:HG22	1:I:504:LEU:HD11	1.84	0.59
1:I:57:ALA:O	1:I:75:LYS:HE2	2.03	0.59
1:M:302:SER:OG	1:M:305:ILE:HB	2.02	0.59
1:N:349:ILE:O	1:N:353:ILE:HG13	2.02	0.59
2:W:48:ILE:HD13	2:W:54:VAL:HG12	1.84	0.59
2:Y:11:ILE:CG1	2:Y:42:ALA:HB3	2.32	0.59
1:B:42:LYS:HE2	1:B:48:THR:CG2	2.32	0.59
1:F:27:VAL:O	1:F:30:THR:HG22	2.02	0.59
1:F:284:ARG:CD	1:F:364:LYS:HE2	2.32	0.59
1:F:127:ALA:HA	1:F:426:LEU:HD11	1.83	0.59
1:J:381:VAL:HG22	1:J:396:VAL:HG21	1.85	0.59
1:L:131:LEU:CD2	1:L:422:VAL:HG11	2.32	0.59
1:M:355:GLU:HG2	1:M:356:ALA:H	1.67	0.59
2:W:10:VAL:CG2	2:W:91:ILE:HD11	2.33	0.59
1:B:180:GLY:HA3	1:B:382:GLY:HA2	1.83	0.59
1:E:450:PRO:O	1:E:454:ILE:HG13	2.03	0.59
1:F:177:VAL:HG11	1:F:397:GLU:CG	2.32	0.59
1:H:226:LYS:HE2	1:H:252:GLU:HG2	1.84	0.59
1:J:303:GLU:OE2	1:J:309:LEU:HD11	2.03	0.59
1:K:6:VAL:CG1	1:K:521:VAL:HG22	2.20	0.59
1:L:219:PHE:HA	1:L:317:LEU:HD11	1.83	0.59
1:N:368:ARG:O	1:N:372:LEU:HG	2.02	0.59
1:A:236:VAL:HG22	1:A:312:ALA:O	2.01	0.59
1:B:351:GLN:HG2	1:C:210:THR:CG2	2.29	0.59
1:E:236:VAL:CG2	1:E:312:ALA:HB3	2.32	0.59
1:M:239:ALA:HB1	1:M:314:LEU:HB3	1.83	0.59
1:N:143:ALA:O	1:N:147:VAL:HG13	2.03	0.59
2:X:20:LYS:CB	2:X:28:THR:OG1	2.50	0.59
2:Y:10:VAL:HG13	2:Y:42:ALA:O	2.02	0.59
1:C:32:GLY:N	3:C:601:ADP:O2A	2.35	0.59
1:E:266:THR:HG22	1:E:273:VAL:H	1.67	0.59
1:N:288:MET:HG2	1:N:368:ARG:CD	2.32	0.59
2:Y:10:VAL:HG22	2:Y:43:VAL:HA	1.83	0.59
2:Y:12:VAL:HG12	2:Y:40:VAL:CA	2.20	0.59
2:Z:85:ILE:HD12	2:Z:85:ILE:O	2.02	0.59
2:1:78:ILE:O	2:1:78:ILE:HG13	2.01	0.59

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:216:GLU:O	1:A:245:LYS:CE	2.51	0.59
1:D:304:GLU:C	1:D:305:ILE:HD12	2.22	0.59
1:D:325:ILE:HG13	1:D:330:THR:HG23	1.85	0.59
1:G:305:ILE:HG22	1:G:306:GLY:N	2.18	0.59
1:H:34:LYS:HD2	1:H:458:CYS:SG	2.42	0.59
1:K:124:VAL:HG13	1:K:504:LEU:CD1	2.32	0.59
1:K:5:ASP:N	1:K:522:THR:O	2.36	0.59
1:L:88:GLY:N	3:L:601:ADP:O1B	2.33	0.59
1:N:346:VAL:HG22	1:N:372:LEU:HD12	1.84	0.59
1:M:517:THR:HG23	1:N:39:VAL:HG23	1.85	0.59
2:P:17:VAL:HG12	2:P:35:SER:CB	2.31	0.59
2:R:51:ASN:OD1	2:R:52:GLY:N	2.36	0.59
2:X:50:GLU:CG	2:X:50:GLU:O	2.46	0.59
2:Y:11:ILE:HG22	2:Y:85:ILE:CB	2.31	0.59
1:H:124:VAL:HG13	1:H:504:LEU:CD1	2.32	0.59
1:I:258:ALA:HA	1:I:261:THR:HG22	1.84	0.59
1:J:224:ASP:OD2	1:J:302:SER:HB3	2.02	0.59
2:R:19:THR:CG2	2:R:33:ALA:HB1	2.30	0.59
2:T:12:VAL:HG12	2:T:40:VAL:CG2	2.25	0.59
2:W:20:LYS:HB3	2:W:22:ALA:N	2.17	0.59
1:G:455:VAL:CG1	1:G:460:GLU:HB2	2.29	0.59
1:H:217:SER:HA	1:H:320:ALA:O	2.02	0.59
1:H:418:ALA:O	1:H:422:VAL:HG13	2.03	0.59
1:I:124:VAL:HG13	1:I:504:LEU:HD12	1.84	0.59
1:N:199:TYR:HE2	1:N:326:ASN:HA	1.68	0.59
2:P:1:MET:O	2:P:1:MET:HG2	2.02	0.59
1:B:236:VAL:CG2	1:B:312:ALA:HB3	2.33	0.59
1:D:291:ASP:HB3	1:D:372:LEU:HD21	1.84	0.59
1:M:310:GLU:CD	1:M:310:GLU:N	2.57	0.59
1:N:418:ALA:O	1:N:422:VAL:HG13	2.03	0.59
1:A:320:ALA:HB1	1:A:334:ASP:O	2.02	0.58
1:B:222:LEU:HD23	1:B:289:LEU:HB3	1.84	0.58
1:C:276:VAL:HG21	1:C:330:THR:OG1	2.03	0.58
1:C:419:LEU:HD21	1:C:500:THR:HG23	1.85	0.58
1:F:205:ILE:HA	1:F:213:VAL:CG2	2.31	0.58
1:I:5:ASP:HB2	1:I:524:LEU:CD2	2.32	0.58
1:M:142:LYS:HE2	1:M:146:GLN:HE22	1.68	0.58
1:N:288:MET:CG	1:N:368:ARG:HD3	2.29	0.58
1:N:449:ALA:HB3	1:N:450:PRO:HD3	1.83	0.58
2:V:40:VAL:CG1	2:V:62:GLY:H	2.16	0.58
2:X:92:LEU:HB3	2:Y:85:ILE:CD1	2.31	0.58

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:358:SER:O	1:A:362:ARG:HB2	2.02	0.58
1:A:449:ALA:HB3	1:A:450:PRO:HD3	1.85	0.58
1:C:219:PHE:HB3	1:C:317:LEU:HB3	1.84	0.58
1:C:350:ARG:HA	1:C:353:ILE:HD12	1.86	0.58
1:I:10:ASN:O	1:I:14:VAL:HG23	2.02	0.58
1:I:285:ARG:HG3	1:I:286:LYS:N	2.18	0.58
2:O:11:ILE:HG12	2:O:85:ILE:HG12	1.85	0.58
2:S:50:GLU:CG	2:S:51:ASN:N	2.65	0.58
2:W:1:MET:CE	2:W:79:ASP:HB3	2.28	0.58
1:A:250:ILE:HG22	1:A:250:ILE:O	2.03	0.58
1:B:205:ILE:HA	1:B:213:VAL:HG22	1.85	0.58
1:B:218:PRO:HG3	1:B:323:VAL:CG1	2.34	0.58
1:E:225:LYS:HG2	1:E:226:LYS:H	1.68	0.58
1:F:351:GLN:HG2	1:G:210:THR:HA	1.85	0.58
1:H:257:GLU:O	1:H:261:THR:HG22	2.03	0.58
1:I:236:VAL:HG21	1:I:312:ALA:HB3	1.83	0.58
1:J:305:ILE:HG12	1:J:306:GLY:H	1.67	0.58
1:K:339:GLU:O	1:K:343:GLN:HB2	2.04	0.58
1:N:285:ARG:HG3	1:N:286:LYS:N	2.19	0.58
2:V:57:LEU:CD1	2:V:59:VAL:HG13	2.33	0.58
1:A:134:LEU:O	1:A:134:LEU:HD23	2.03	0.58
1:A:304:GLU:N	1:A:304:GLU:OE1	2.36	0.58
1:A:429:LEU:HG	1:A:440:ILE:HD13	1.85	0.58
1:E:359:ASP:OD1	1:E:362:ARG:NH2	2.36	0.58
1:G:510:VAL:O	1:G:514:MET:HB2	2.03	0.58
1:H:293:ALA:O	1:H:298:GLY:N	2.35	0.58
1:H:162:ILE:HG21	1:H:403:THR:HG21	1.86	0.58
1:L:291:ASP:OD1	1:L:292:ILE:HG13	2.04	0.58
2:O:46:GLY:HA2	2:O:57:LEU:HD22	1.86	0.58
1:B:231:ARG:NH1	2:P:31:ALA:HB2	2.18	0.58
2:T:48:ILE:HA	2:T:53:GLU:O	2.02	0.58
1:A:306:GLY:HA3	1:B:264:VAL:CG2	2.31	0.58
1:B:172:GLU:N	1:B:172:GLU:OE1	2.36	0.58
1:B:270:ILE:HG23	1:B:271:VAL:H	1.66	0.58
1:C:215:LEU:HD11	1:C:274:ALA:HB3	1.84	0.58
1:D:280:GLY:O	1:D:285:ARG:HB2	2.02	0.58
1:D:349:ILE:HA	1:D:352:GLN:CG	2.33	0.58
1:E:449:ALA:HB3	1:E:450:PRO:HD3	1.85	0.58
1:G:205:ILE:CA	1:G:213:VAL:HG22	2.14	0.58
1:J:285:ARG:O	1:J:289:LEU:HG	2.04	0.58
1:L:124:VAL:HG21	1:L:508:ALA:HB1	1.85	0.58

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:N:201:SER:HB3	1:N:204:PHE:CE2	2.38	0.58
2:Q:57:LEU:HD13	2:Q:59:VAL:HG22	1.84	0.58
2:R:36:THR:HG23	2:R:37:ARG:CG	2.26	0.58
2:S:23:GLY:O	2:S:26:VAL:HG12	2.04	0.58
1:B:234:LEU:HD23	1:B:235:PRO:HD3	1.84	0.58
1:F:232:GLU:O	1:F:235:PRO:HD2	2.04	0.58
1:G:248:LEU:HD23	1:G:325:ILE:HD11	1.85	0.58
1:G:412:VAL:CG1	1:G:413:ALA:N	2.66	0.58
1:J:513:LEU:HD11	1:K:388:GLU:HA	1.86	0.58
1:N:352:GLN:CB	1:N:365:LEU:HD11	2.33	0.58
1:N:496:PRO:HG2	1:N:499:VAL:CG1	2.34	0.58
2:S:55:LYS:HD2	2:S:56:PRO:HD2	1.84	0.58
2:U:43:VAL:HG23	2:U:57:LEU:HD22	1.84	0.58
1:K:238:GLU:HG3	2:Y:23:GLY:HA3	1.85	0.58
2:X:93:ALA:HB2	2:Y:5:PRO:HA	1.86	0.58
2:2:64:ILE:HG23	2:2:95:VAL:HB	1.86	0.58
1:H:252:GLU:OE1	1:H:285:ARG:NH2	2.36	0.58
1:H:220:ILE:O	1:H:317:LEU:HD22	2.03	0.58
1:I:322:ARG:HG2	1:I:323:VAL:N	2.19	0.58
2:Q:60:LYS:HZ2	2:Q:63:ASP:HB3	1.69	0.58
2:V:50:GLU:CG	2:W:50:GLU:HB3	2.29	0.58
1:A:199:TYR:CZ	1:A:205:ILE:HD11	2.39	0.58
1:A:517:THR:HG23	1:B:39:VAL:HG23	1.85	0.58
1:E:278:ALA:HB3	1:E:285:ARG:HD2	1.86	0.58
1:F:322:ARG:HG2	1:F:323:VAL:N	2.19	0.58
1:I:357:THR:HB	1:I:361:ASP:HB2	1.85	0.58
1:I:417:VAL:HA	1:I:420:ILE:CG2	2.33	0.58
1:L:392:LYS:HD2	1:L:395:ARG:HH21	1.69	0.58
1:M:169:VAL:HG11	1:M:377:ALA:HB2	1.86	0.58
2:Q:94:ILE:HD11	2:R:4:ARG:NH2	2.19	0.58
1:B:228:SER:CB	1:B:255:GLU:HG2	2.30	0.58
1:B:496:PRO:HG2	1:B:499:VAL:CG1	2.33	0.58
1:E:234:LEU:CG	1:E:235:PRO:N	2.67	0.58
1:E:342:ILE:O	1:E:346:VAL:HG23	2.03	0.58
1:F:226:LYS:HB3	1:F:252:GLU:HG2	1.85	0.58
1:F:414:GLY:HA2	1:F:495:ASP:OD2	2.04	0.58
1:H:207:LYS:CB	1:H:208:PRO:HD3	2.28	0.58
1:H:370:ALA:O	1:H:374:GLY:N	2.37	0.58
1:K:231:ARG:HD3	1:K:257:GLU:OE2	2.03	0.58
1:J:517:THR:HG23	1:K:39:VAL:HG23	1.84	0.58
1:N:30:THR:HB	1:N:51:LYS:HG3	1.86	0.58

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:B:231:ARG:NH2	2:P:31:ALA:HB2	2.18	0.58
2:S:14:ARG:HG2	2:S:35:SER:CB	2.33	0.58
2:T:17:VAL:HG23	2:T:33:ALA:CB	2.30	0.58
1:A:206:ASN:ND2	1:A:214:GLU:O	2.36	0.58
1:B:193:MET:SD	1:B:371:LYS:HB3	2.43	0.58
1:C:349:ILE:O	1:C:353:ILE:HG13	2.04	0.58
1:G:415:GLY:N	3:G:601:ADP:O2'	2.30	0.58
1:H:230:ILE:HB	1:H:309:LEU:HD13	1.85	0.58
1:H:338:GLU:HG2	1:H:341:ALA:H	1.69	0.58
1:I:124:VAL:HG13	1:I:504:LEU:CD1	2.34	0.58
1:K:510:VAL:O	1:K:514:MET:HB2	2.03	0.58
1:K:4:LYS:HG3	1:L:59:GLU:O	2.04	0.58
1:N:276:VAL:HG11	1:N:325:ILE:HG21	1.85	0.58
2:X:94:ILE:CD1	2:X:96:GLU:HG3	2.34	0.58
2:2:52:GLY:HA3	2:2:53:GLU:HB3	1.85	0.57
1:E:174:VAL:HG22	1:E:376:VAL:CG1	2.33	0.57
1:F:180:GLY:HA3	1:F:382:GLY:HA2	1.86	0.57
1:J:199:TYR:HD2	1:J:325:ILE:HG22	1.69	0.57
1:L:291:ASP:OD1	1:L:292:ILE:N	2.37	0.57
1:M:345:ARG:O	1:M:348:GLN:HG3	2.03	0.57
1:N:355:GLU:HG3	1:N:356:ALA:H	1.69	0.57
2:X:15:LYS:HG3	2:X:16:GLU:OE1	2.03	0.57
1:E:245:LYS:HA	1:E:245:LYS:HE2	1.85	0.57
1:E:290:GLN:HG2	1:E:300:VAL:HG21	1.85	0.57
1:F:419:LEU:HD12	1:F:419:LEU:H	1.68	0.57
1:G:313:THR:CG2	1:G:315:GLU:HG2	2.34	0.57
1:K:205:ILE:HA	1:K:213:VAL:CG2	2.33	0.57
1:K:322:ARG:HB3	1:K:333:ILE:CG2	2.34	0.57
1:L:237:LEU:HD22	2:Z:26:VAL:HG21	1.85	0.57
2:Q:92:LEU:HD23	2:Q:92:LEU:N	2.18	0.57
2:S:12:VAL:HG13	2:S:40:VAL:N	2.19	0.57
2:U:46:GLY:HA2	2:U:57:LEU:HD12	1.85	0.57
1:E:219:PHE:HB2	1:E:247:LEU:CD1	2.34	0.57
1:I:142:LYS:HE2	1:I:146:GLN:NE2	2.18	0.57
1:I:169:VAL:HG23	1:I:173:GLY:HA3	1.85	0.57
1:J:311:LYS:O	1:J:311:LYS:HD3	2.04	0.57
1:K:455:VAL:HG11	1:K:462:PRO:HA	1.85	0.57
1:L:368:ARG:O	1:L:372:LEU:HG	2.05	0.57
1:N:477:GLY:HA3	1:N:488:MET:CE	2.32	0.57
1:B:88:GLY:HA2	3:B:601:ADP:O2B	2.03	0.57
3:B:601:ADP:O3B	4:B:602:BEF:F2	2.13	0.57

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:H:304:GLU:OE1	1:H:304:GLU:N	2.37	0.57
1:H:360:TYR:HA	1:H:363:GLU:HG3	1.85	0.57
1:H:448:GLU:O	1:H:452:ARG:HG2	2.05	0.57
1:I:166:MET:CE	1:I:407:VAL:HG21	2.34	0.57
1:J:225:LYS:HD2	1:J:309:LEU:HD21	1.85	0.57
1:J:76:GLU:OE2	1:K:385:THR:HB	2.04	0.57
1:N:221:LEU:HD13	1:N:317:LEU:CD2	2.35	0.57
2:T:13:LYS:HD2	2:T:81:GLU:OE1	2.04	0.57
1:D:230:ILE:O	1:D:231:ARG:CG	2.48	0.57
1:D:166:MET:CE	1:D:407:VAL:HG21	2.35	0.57
1:H:210:THR:O	1:H:326:ASN:ND2	2.37	0.57
1:I:241:ALA:HB2	1:I:271:VAL:CG1	2.35	0.57
1:L:228:SER:O	1:L:231:ARG:HG3	2.03	0.57
1:M:254:VAL:HG21	1:M:275:ALA:HB1	1.86	0.57
2:Z:25:ILE:HD12	2:Z:25:ILE:H	1.69	0.57
2:Z:40:VAL:HG21	2:Z:59:VAL:CG1	2.35	0.57
1:A:25:ASP:HA	1:A:28:LYS:HE2	1.87	0.57
1:A:5:ASP:HB2	1:A:524:LEU:CD2	2.34	0.57
1:B:221:LEU:HD23	1:B:222:LEU:N	2.20	0.57
1:D:231:ARG:O	1:D:233:MET:HG2	2.04	0.57
1:D:40:LEU:HD13	1:D:59:GLU:HG3	1.86	0.57
1:E:350:ARG:HD3	1:E:353:ILE:HD12	1.86	0.57
1:F:199:TYR:CE1	1:F:202:PRO:HA	2.40	0.57
1:F:409:GLU:OE2	1:F:501:ARG:NH2	2.29	0.57
1:F:406:ALA:HB2	1:F:496:PRO:HB3	1.85	0.57
1:H:124:VAL:HG21	1:H:508:ALA:CB	2.35	0.57
1:M:308:GLU:HB2	1:M:311:LYS:HB3	1.84	0.57
1:N:147:VAL:HG23	1:N:403:THR:HG22	1.87	0.57
1:N:165:ALA:O	1:N:169:VAL:HG22	2.03	0.57
2:U:10:VAL:HG23	2:U:42:ALA:O	2.04	0.57
2:U:49:LEU:C	2:U:50:GLU:CD	2.62	0.57
2:X:40:VAL:HG12	2:X:63:ASP:O	2.04	0.57
2:X:50:GLU:O	2:X:51:ASN:ND2	2.38	0.57
2:2:49:LEU:HB2	2:2:52:GLY:CA	2.35	0.57
1:C:227:ILE:HG12	1:C:251:ALA:HB1	1.85	0.57
1:C:240:VAL:HG11	1:C:247:LEU:HD23	1.86	0.57
1:C:345:ARG:HA	1:C:348:GLN:CG	2.34	0.57
1:C:350:ARG:NH1	1:C:369:VAL:HG11	2.20	0.57
1:E:222:LEU:CD2	1:E:289:LEU:HD22	2.35	0.57
1:G:218:PRO:HG3	1:G:323:VAL:HG13	1.86	0.57
1:K:236:VAL:HG21	1:K:310:GLU:O	2.05	0.57

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:L:10:ASN:O	1:L:14:VAL:HG23	2.04	0.57
1:L:194:GLN:O	1:L:371:LYS:NZ	2.30	0.57
1:L:296:THR:O	1:L:319:GLN:N	2.36	0.57
1:L:517:THR:HG23	1:M:39:VAL:HG23	1.86	0.57
1:N:381:VAL:HG11	1:N:392:LYS:HB3	1.87	0.57
2:Q:40:VAL:CG1	2:Q:61:VAL:HA	2.35	0.57
1:B:418:ALA:O	1:B:422:VAL:HG13	2.05	0.57
1:C:42:LYS:HE2	1:C:48:THR:CG2	2.35	0.57
1:E:311:LYS:HD3	1:E:311:LYS:O	2.04	0.57
1:G:453:GLN:NE2	1:G:456:LEU:HD23	2.20	0.57
1:I:305:ILE:CG2	1:I:306:GLY:H	2.17	0.57
1:L:225:LYS:HG3	1:L:226:LYS:N	2.20	0.57
1:N:363:GLU:HG3	1:N:364:LYS:H	1.68	0.57
2:Y:36:THR:O	2:Y:66:ILE:HG12	2.05	0.57
2:2:65:VAL:HG12	2:2:94:ILE:CG2	2.25	0.57
1:A:393:LYS:O	1:A:397:GLU:HG3	2.05	0.57
1:B:169:VAL:CG1	1:B:173:GLY:HA3	2.35	0.57
1:C:220:ILE:HG22	1:C:222:LEU:HD11	1.87	0.57
1:E:24:ALA:O	1:E:28:LYS:HG3	2.05	0.57
1:G:305:ILE:HG22	1:G:306:GLY:H	1.70	0.57
1:H:204:PHE:CE2	1:H:274:ALA:HA	2.39	0.57
1:H:230:ILE:HG12	1:H:309:LEU:HB3	1.87	0.57
1:J:414:GLY:O	1:J:417:VAL:HG12	2.05	0.57
1:M:227:ILE:HD12	1:M:227:ILE:N	2.20	0.57
2:R:48:ILE:CA	2:R:54:VAL:HG12	2.28	0.57
2:T:20:LYS:HB3	2:T:27:LEU:HA	1.87	0.57
1:A:245:LYS:HE3	1:A:246:PRO:HD3	1.87	0.57
1:F:200:LEU:HD13	1:F:276:VAL:HA	1.86	0.57
1:H:18:ARG:NH1	1:H:18:ARG:HB3	2.20	0.57
1:H:231:ARG:HE	1:H:257:GLU:HG3	1.70	0.57
1:I:152:ALA:HB2	1:I:399:ALA:HB2	1.86	0.57
1:M:305:ILE:HD12	1:M:305:ILE:N	2.20	0.57
1:N:136:VAL:CG1	1:N:137:PRO:HD2	2.35	0.57
1:A:177:VAL:HG11	1:A:396:VAL:HG12	1.87	0.56
1:B:208:PRO:HB2	1:B:212:ALA:HB3	1.86	0.56
1:B:415:GLY:N	3:B:601:ADP:O2'	2.26	0.56
1:G:149:THR:HG23	1:G:159:GLY:HA3	1.86	0.56
1:G:219:PHE:HB2	1:G:247:LEU:CD1	2.34	0.56
1:G:256:GLY:HA2	1:G:259:LEU:HB2	1.86	0.56
1:I:169:VAL:CG2	1:I:173:GLY:HA3	2.34	0.56
1:J:363:GLU:HG3	1:J:364:LYS:N	2.20	0.56

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:K:215:LEU:HD12	1:K:323:VAL:HG21	1.86	0.56
2:T:49:LEU:HD22	2:U:50:GLU:HA	1.86	0.56
2:T:65:VAL:HG12	2:T:94:ILE:HG22	1.87	0.56
1:A:270:ILE:HG22	1:A:271:VAL:N	2.20	0.56
1:B:276:VAL:CG2	1:B:325:ILE:HG21	2.34	0.56
1:C:165:ALA:O	1:C:169:VAL:HG22	2.05	0.56
1:C:443:ALA:O	1:C:447:MET:HG3	2.05	0.56
1:E:302:SER:HB2	1:E:305:ILE:HG22	1.86	0.56
1:G:77:VAL:HG21	1:G:507:ALA:CA	2.33	0.56
1:J:181:THR:HG23	1:J:182:GLY:H	1.70	0.56
1:J:39:VAL:HG22	1:J:49:ILE:HG12	1.87	0.56
1:K:230:ILE:HG21	1:K:233:MET:HG3	1.86	0.56
1:K:252:GLU:HG3	1:K:285:ARG:HH12	1.71	0.56
1:L:355:GLU:HG3	1:L:356:ALA:N	2.18	0.56
1:M:304:GLU:C	1:M:305:ILE:HD12	2.26	0.56
1:N:168:LYS:HD3	1:N:189:VAL:HG21	1.87	0.56
1:N:301:ILE:HG12	1:N:307:MET:HE2	1.86	0.56
1:N:345:ARG:O	1:N:348:GLN:HG3	2.05	0.56
1:N:346:VAL:O	1:N:349:ILE:HG22	2.06	0.56
2:O:57:LEU:H	2:O:57:LEU:HD22	1.70	0.56
2:O:37:ARG:HG2	2:O:66:ILE:HG12	1.86	0.56
2:W:17:VAL:HG23	2:W:19:THR:HA	1.87	0.56
1:B:177:VAL:HG21	1:B:397:GLU:HG2	1.85	0.56
1:B:231:ARG:HD2	1:B:258:ALA:HA	1.87	0.56
1:C:349:ILE:HD13	1:C:368:ARG:HD2	1.88	0.56
1:E:222:LEU:HD13	1:E:289:LEU:HD13	1.88	0.56
1:E:248:LEU:HD13	1:E:323:VAL:HG21	1.88	0.56
1:G:225:LYS:HG3	1:G:226:LYS:H	1.70	0.56
1:H:263:VAL:O	1:H:267:MET:HG2	2.05	0.56
1:I:418:ALA:O	1:I:422:VAL:HG13	2.06	0.56
1:I:85:ALA:HB1	1:I:499:VAL:HG12	1.86	0.56
1:A:314:LEU:HD12	1:A:315:GLU:N	2.20	0.56
1:B:510:VAL:O	1:B:514:MET:HB2	2.05	0.56
1:D:236:VAL:O	1:D:240:VAL:HG23	2.05	0.56
1:F:219:PHE:HE2	1:F:240:VAL:HG13	1.69	0.56
1:G:18:ARG:HB3	1:G:18:ARG:NH1	2.21	0.56
1:G:203:TYR:HB3	1:G:267:MET:HE1	1.88	0.56
1:G:224:ASP:OD1	1:G:286:LYS:HG2	2.05	0.56
1:M:233:MET:HE3	1:M:237:LEU:HD23	1.88	0.56
1:M:23:LEU:HD12	1:M:60:ILE:HB	1.87	0.56
1:N:225:LYS:HG3	1:N:226:LYS:N	2.20	0.56

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:X:48:ILE:H	2:X:48:ILE:HD12	1.69	0.56
1:B:305:ILE:HD11	1:C:203:TYR:CE2	2.41	0.56
1:E:359:ASP:O	1:E:363:GLU:HG3	2.05	0.56
1:E:177:VAL:HG11	1:E:397:GLU:HG2	1.86	0.56
1:E:77:VAL:HB	1:E:510:VAL:HG21	1.88	0.56
1:H:302:SER:HB2	1:H:305:ILE:CD1	2.24	0.56
1:H:276:VAL:HG13	1:H:325:ILE:HD13	1.87	0.56
1:J:5:ASP:HB2	1:J:524:LEU:HD23	1.87	0.56
1:L:27:VAL:CG1	1:L:90:THR:HG23	2.35	0.56
2:V:49:LEU:HD13	2:V:50:GLU:N	2.20	0.56
2:X:77:LYS:HE3	2:X:82:GLU:HB2	1.87	0.56
2:V:74:LYS:HZ3	2:2:68:ASN:HB3	1.69	0.56
2:2:59:VAL:HG11	2:2:91:ILE:HD12	1.88	0.56
1:A:353:ILE:HA	1:A:365:LEU:HD22	1.87	0.56
1:A:174:VAL:HG13	1:A:370:ALA:CB	2.34	0.56
1:A:420:ILE:HD11	1:A:451:LEU:HB2	1.88	0.56
1:B:355:GLU:HG3	1:B:356:ALA:H	1.70	0.56
1:C:102:GLU:OE1	1:C:445:ARG:NH1	2.36	0.56
1:C:368:ARG:O	1:C:372:LEU:HG	2.06	0.56
1:D:23:LEU:HD23	1:D:60:ILE:HB	1.88	0.56
1:E:234:LEU:CD2	1:E:235:PRO:HD3	2.29	0.56
1:F:258:ALA:O	1:F:262:LEU:HB2	2.05	0.56
1:G:225:LYS:HD3	1:G:303:GLU:OE1	2.06	0.56
1:H:236:VAL:HG22	1:H:312:ALA:HB3	1.86	0.56
1:I:230:ILE:H	1:I:230:ILE:HD12	1.69	0.56
1:J:307:MET:HG2	1:J:311:LYS:NZ	2.20	0.56
1:L:18:ARG:HB3	1:L:18:ARG:NH1	2.21	0.56
1:N:130:GLU:O	1:N:134:LEU:HD13	2.06	0.56
1:N:284:ARG:CD	1:N:364:LYS:HE2	2.34	0.56
2:O:8:ASP:CA	2:O:57:LEU:HD21	2.31	0.56
2:U:23:GLY:O	2:U:26:VAL:HG12	2.05	0.56
2:V:57:LEU:HD11	2:V:59:VAL:HG22	1.87	0.56
2:Y:92:LEU:HD23	2:Z:85:ILE:CD1	2.35	0.56
2:1:95:VAL:HA	2:2:3:ILE:CG2	2.36	0.56
2:2:53:GLU:OE2	2:2:54:VAL:HG23	2.06	0.56
1:A:234:LEU:HG	1:A:235:PRO:CD	2.35	0.56
1:A:241:ALA:HB1	1:A:271:VAL:HG12	1.88	0.56
1:B:250:ILE:HG22	1:B:278:ALA:HB2	1.87	0.56
1:B:48:THR:HG22	1:B:390:LYS:NZ	2.19	0.56
1:H:165:ALA:O	1:H:169:VAL:HG22	2.06	0.56
1:H:225:LYS:HG2	1:H:303:GLU:HG3	1.87	0.56

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:H:323:VAL:HG12	1:H:332:ILE:HG12	1.88	0.56
1:I:326:ASN:HD21	1:I:329:THR:HB	1.71	0.56
1:M:18:ARG:NH1	1:M:18:ARG:HB3	2.20	0.56
1:M:448:GLU:O	1:M:452:ARG:HG3	2.06	0.56
1:N:231:ARG:HB3	1:N:233:MET:HG3	1.88	0.56
1:N:279:PRO:CB	1:N:288:MET:HE3	2.35	0.56
1:N:222:LEU:HB2	1:N:300:VAL:HA	1.85	0.56
1:N:214:GLU:HG2	1:N:324:VAL:HG22	1.87	0.56
2:R:6:LEU:CD2	2:R:7:HIS:CE1	2.55	0.56
2:Y:20:LYS:HE2	2:Y:22:ALA:CB	2.36	0.56
1:C:25:ASP:HA	1:C:28:LYS:HE2	1.86	0.56
1:E:231:ARG:HD3	1:E:257:GLU:OE2	2.06	0.56
1:E:345:ARG:O	1:E:348:GLN:HG3	2.06	0.56
1:H:260:ALA:O	1:H:264:VAL:HG23	2.06	0.56
1:J:230:ILE:O	1:J:232:GLU:N	2.39	0.56
1:J:175:ILE:HG12	1:J:377:ALA:HB3	1.87	0.56
1:K:225:LYS:HG3	1:K:226:LYS:H	1.71	0.56
1:N:77:VAL:HB	1:N:510:VAL:CG2	2.32	0.56
2:1:64:ILE:CG2	2:1:95:VAL:HB	2.31	0.56
1:A:266:THR:O	1:A:269:GLY:HA3	2.05	0.56
1:A:199:TYR:HB3	1:A:276:VAL:HG12	1.87	0.56
1:C:203:TYR:HB3	1:C:267:MET:CE	2.36	0.56
1:C:87:ASP:OD2	1:C:151:SER:OG	2.16	0.56
1:H:231:ARG:HE	1:H:257:GLU:CG	2.19	0.56
1:K:230:ILE:CG2	1:K:233:MET:HG3	2.36	0.56
1:K:237:LEU:C	1:K:237:LEU:HD12	2.26	0.56
1:M:208:PRO:HG3	1:M:214:GLU:CD	2.26	0.56
2:O:96:GLU:HB3	2:P:2:ASN:H	1.70	0.56
2:R:6:LEU:HD23	2:R:7:HIS:CB	2.35	0.56
2:R:95:VAL:HG22	2:S:3:ILE:HG21	1.88	0.56
1:F:127:ALA:N	1:F:426:LEU:HD11	2.21	0.56
1:G:215:LEU:HD22	1:G:246:PRO:CB	2.35	0.56
1:G:333:ILE:O	1:G:334:ASP:HB2	2.06	0.56
1:H:230:ILE:HD13	1:H:309:LEU:HB2	1.87	0.56
1:J:227:ILE:HD12	1:J:227:ILE:N	2.20	0.56
1:K:77:VAL:HG12	1:K:510:VAL:HG21	1.88	0.56
1:M:263:VAL:O	1:M:267:MET:HG2	2.06	0.56
1:M:151:SER:HB3	1:M:399:ALA:HA	1.86	0.56
2:P:73:VAL:HG22	2:P:86:MET:HB3	1.88	0.56
2:T:36:THR:O	2:T:66:ILE:HG12	2.06	0.56
2:Y:36:THR:HG22	2:Y:37:ARG:NH1	2.21	0.56

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:Y:7:HIS:O	2:Y:8:ASP:HB3	2.04	0.56
2:1:66:ILE:HD12	2:2:3:ILE:CD1	2.35	0.56
1:A:236:VAL:O	1:A:240:VAL:HG23	2.06	0.56
1:A:280:GLY:O	1:A:285:ARG:NE	2.39	0.56
1:B:128:VAL:CG1	1:B:132:LYS:HE2	2.34	0.56
1:J:308:GLU:HB3	1:J:311:LYS:HB3	1.87	0.56
1:K:90:THR:OG1	3:K:601:ADP:O2B	2.21	0.56
2:R:8:ASP:OD1	2:R:9:ARG:N	2.38	0.56
2:S:57:LEU:H	2:S:57:LEU:HD12	1.69	0.56
2:Z:47:ARG:HD3	2:Z:49:LEU:HD12	1.88	0.56
2:2:49:LEU:C	2:2:52:GLY:H	2.09	0.55
2:2:57:LEU:HD13	2:2:88:GLU:HB2	1.88	0.55
1:A:233:MET:C	1:A:235:PRO:HD2	2.27	0.55
1:A:6:VAL:CG1	1:A:521:VAL:HG22	2.31	0.55
1:D:10:ASN:O	1:D:14:VAL:HG23	2.05	0.55
1:D:230:ILE:O	1:D:230:ILE:HG23	2.06	0.55
1:F:345:ARG:O	1:F:348:GLN:HG3	2.06	0.55
1:G:183:LEU:HD12	1:G:384:ALA:HB2	1.89	0.55
1:G:280:GLY:H	1:G:285:ARG:HB3	1.71	0.55
1:L:417:VAL:HG21	1:L:477:GLY:HA3	1.87	0.55
1:N:231:ARG:C	1:N:232:GLU:HG2	2.26	0.55
2:S:50:GLU:HG2	2:S:51:ASN:N	2.20	0.55
2:U:15:LYS:HB2	2:U:37:ARG:O	2.06	0.55
2:W:77:LYS:HG3	2:W:81:GLU:O	2.06	0.55
1:L:237:LEU:HD22	2:Z:26:VAL:CG2	2.36	0.55
1:A:176:THR:OG1	1:A:378:VAL:HG12	2.06	0.55
1:A:301:ILE:CD1	1:A:312:ALA:HB2	2.35	0.55
1:B:359:ASP:O	1:B:360:TYR:HB2	2.05	0.55
1:C:57:ALA:O	1:C:75:LYS:HE2	2.06	0.55
1:D:496:PRO:HG2	1:D:499:VAL:HG11	1.88	0.55
1:E:368:ARG:O	1:E:372:LEU:HG	2.07	0.55
1:F:311:LYS:HZ1	1:F:313:THR:HG1	1.53	0.55
1:G:134:LEU:HD11	1:G:425:LYS:HZ1	1.69	0.55
1:G:291:ASP:HB2	1:G:372:LEU:HD21	1.87	0.55
1:G:461:GLU:OE2	1:I:452:ARG:NH2	2.38	0.55
1:G:85:ALA:HB1	1:G:499:VAL:HG12	1.88	0.55
1:I:381:VAL:HG11	1:I:392:LYS:HB3	1.86	0.55
1:N:227:ILE:HG23	1:N:231:ARG:CB	2.37	0.55
1:N:475:ASN:O	1:N:488:MET:HG2	2.06	0.55
2:O:95:VAL:HA	2:P:3:ILE:HG22	0.59	0.55
1:E:260:ALA:HB3	2:S:30:SER:HB3	1.87	0.55

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:V:49:LEU:HD13	2:V:50:GLU:CB	2.37	0.55
2:Y:40:VAL:HG12	2:Y:63:ASP:O	2.07	0.55
1:B:510:VAL:HG12	1:C:387:VAL:HG21	1.88	0.55
1:D:90:THR:O	1:D:94:VAL:HG23	2.07	0.55
1:E:325:ILE:N	1:E:325:ILE:HD12	2.21	0.55
1:F:33:PRO:HA	1:F:153:ASN:OD1	2.07	0.55
1:G:228:SER:O	1:G:231:ARG:HG3	2.07	0.55
1:L:349:ILE:O	1:L:353:ILE:HG13	2.06	0.55
1:L:479:ASN:HD22	1:L:493:ILE:CD1	2.15	0.55
1:N:158:VAL:O	1:N:162:ILE:HG13	2.06	0.55
1:C:10:ASN:O	1:C:14:VAL:HG23	2.06	0.55
1:C:245:LYS:HE2	1:C:245:LYS:HA	1.87	0.55
1:D:27:VAL:HG12	1:D:90:THR:HG23	1.88	0.55
1:F:31:LEU:CD2	1:F:453:GLN:HB3	2.36	0.55
1:H:449:ALA:HB3	1:H:450:PRO:HD3	1.87	0.55
1:I:147:VAL:HG22	1:I:403:THR:HG22	1.87	0.55
1:J:420:ILE:HD11	1:J:469:VAL:CG1	2.35	0.55
1:M:131:LEU:CD2	1:M:422:VAL:HG11	2.36	0.55
1:M:197:ARG:HD2	1:M:277:LYS:CB	2.34	0.55
2:O:10:VAL:N	2:O:86:MET:O	2.36	0.55
2:T:40:VAL:CG1	2:T:62:GLY:H	2.19	0.55
2:T:64:ILE:N	2:T:64:ILE:HD12	2.21	0.55
1:A:199:TYR:HE1	1:A:202:PRO:HA	1.71	0.55
1:A:31:LEU:HD23	1:A:453:GLN:HB3	1.88	0.55
1:G:223:ALA:HB2	1:G:251:ALA:CA	2.32	0.55
1:L:230:ILE:HG12	1:L:309:LEU:CD2	2.36	0.55
1:C:238:GLU:HG2	2:Q:25:ILE:CG1	2.35	0.55
2:R:17:VAL:HG23	2:R:19:THR:N	2.19	0.55
2:R:20:LYS:HB2	2:R:22:ALA:CB	2.36	0.55
2:Z:16:GLU:OE1	2:Z:16:GLU:N	2.39	0.55
1:B:232:GLU:O	1:B:234:LEU:HD23	2.06	0.55
1:C:231:ARG:CD	1:C:261:THR:HG21	2.36	0.55
1:D:463:SER:HB2	1:L:463:SER:HB2	1.89	0.55
1:E:420:ILE:HD11	1:E:469:VAL:HG12	1.87	0.55
1:F:124:VAL:HG13	1:F:504:LEU:HD12	1.88	0.55
1:I:404:ARG:O	1:I:408:GLU:HG2	2.06	0.55
1:J:124:VAL:HG13	1:J:504:LEU:HD12	1.88	0.55
1:J:174:VAL:O	1:J:376:VAL:HG13	2.07	0.55
1:J:367:GLU:O	1:J:371:LYS:HG3	2.07	0.55
1:L:219:PHE:HA	1:L:317:LEU:CD2	2.29	0.55
1:L:5:ASP:HB2	1:L:524:LEU:HD23	1.88	0.55

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:M:165:ALA:O	1:M:169:VAL:HG22	2.06	0.55
1:N:222:LEU:CD2	1:N:300:VAL:HG22	2.36	0.55
2:Y:17:VAL:HG23	2:Y:33:ALA:CB	2.37	0.55
2:1:80:ASN:OD1	2:1:80:ASN:N	2.39	0.55
1:A:245:LYS:HE3	1:A:246:PRO:CD	2.36	0.55
1:A:365:LEU:HD12	1:A:366:GLN:N	2.21	0.55
1:B:180:GLY:N	1:B:381:VAL:O	2.39	0.55
3:B:601:ADP:O3A	4:B:602:BEF:F3	2.15	0.55
1:C:203:TYR:HD2	1:C:263:VAL:HG11	1.72	0.55
1:C:176:THR:OG1	1:C:378:VAL:HG22	2.05	0.55
1:D:355:GLU:HG3	1:D:356:ALA:H	1.71	0.55
1:D:5:ASP:HB2	1:D:524:LEU:HD21	1.88	0.55
1:G:101:THR:O	1:G:105:LYS:HG3	2.05	0.55
1:I:265:ASN:OD1	2:W:27:LEU:HA	2.06	0.55
1:J:205:ILE:HD13	1:J:211:GLY:HA2	1.88	0.55
1:J:236:VAL:CG1	1:J:312:ALA:HB3	2.37	0.55
1:J:218:PRO:HA	1:J:246:PRO:HG2	1.88	0.55
1:K:219:PHE:HD2	1:K:240:VAL:HG13	1.71	0.55
1:K:325:ILE:N	1:K:325:ILE:HD12	2.21	0.55
1:L:225:LYS:C	1:L:252:GLU:HB3	2.27	0.55
2:T:36:THR:C	2:T:37:ARG:HD2	2.27	0.55
2:T:48:ILE:HG22	2:T:53:GLU:O	2.07	0.55
1:A:309:LEU:CD1	1:A:310:GLU:H	2.09	0.55
1:A:420:ILE:HD11	1:A:451:LEU:CB	2.36	0.55
1:A:455:VAL:CG1	1:A:460:GLU:HB2	2.37	0.55
1:B:346:VAL:O	1:B:349:ILE:HG22	2.07	0.55
1:B:517:THR:HG23	1:C:39:VAL:HG23	1.88	0.55
1:C:383:ALA:CB	1:C:389:MET:HA	2.36	0.55
1:D:199:TYR:HE1	1:D:202:PRO:HA	1.70	0.55
1:E:219:PHE:HE1	1:E:319:GLN:HG2	1.71	0.55
1:E:238:GLU:HA	1:E:241:ALA:HB3	1.88	0.55
1:F:349:ILE:O	1:F:353:ILE:N	2.40	0.55
1:G:220:ILE:C	1:G:221:LEU:HD12	2.27	0.55
1:H:203:TYR:CE2	1:N:305:ILE:HD11	2.41	0.55
1:H:230:ILE:HG23	1:H:231:ARG:N	2.22	0.55
1:M:233:MET:CE	1:M:237:LEU:HD23	2.37	0.55
1:A:199:TYR:CA	1:A:276:VAL:HG12	2.37	0.55
1:F:171:LYS:O	1:F:171:LYS:HD2	2.07	0.55
1:F:226:LYS:CA	1:F:252:GLU:HB3	2.36	0.55
1:H:314:LEU:HD12	1:H:315:GLU:N	2.22	0.55
1:M:280:GLY:O	1:M:285:ARG:HB3	2.07	0.55

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:N:231:ARG:O	1:N:232:GLU:HG2	2.07	0.55
2:S:40:VAL:HG12	2:S:41:LEU:N	2.18	0.55
2:V:17:VAL:HA	2:V:35:SER:HB2	1.87	0.55
2:V:48:ILE:HG13	2:V:54:VAL:HG12	1.87	0.55
2:Z:17:VAL:HG13	2:Z:18:GLU:CD	2.27	0.55
1:A:223:ALA:HB3	1:A:251:ALA:CB	2.36	0.55
1:B:208:PRO:HG2	1:B:214:GLU:HG3	1.89	0.55
1:C:226:LYS:CG	1:C:252:GLU:HG2	2.25	0.55
1:C:364:LYS:HA	1:C:367:GLU:HB3	1.88	0.55
1:C:418:ALA:O	1:C:422:VAL:HG13	2.06	0.55
1:G:252:GLU:HG3	1:G:285:ARG:NH1	2.21	0.55
1:G:313:THR:HB	1:G:315:GLU:HG2	1.89	0.55
1:H:142:LYS:O	1:H:146:GLN:HG3	2.07	0.55
1:H:305:ILE:CG2	1:H:306:GLY:N	2.69	0.55
1:J:270:ILE:HG21	2:X:27:LEU:HD22	1.89	0.55
1:J:23:LEU:HD23	1:J:60:ILE:CB	2.37	0.55
1:K:304:GLU:C	1:K:305:ILE:HD12	2.27	0.55
1:L:234:LEU:CD1	1:L:234:LEU:H	2.18	0.55
1:L:412:VAL:CG1	1:L:413:ALA:N	2.69	0.55
1:M:25:ASP:HA	1:M:28:LYS:HE2	1.88	0.55
2:P:3:ILE:H	2:P:3:ILE:HD12	1.71	0.55
2:R:10:VAL:HG13	2:R:57:LEU:HD13	1.88	0.55
1:B:398:ASP:OD2	4:B:602:BEF:F1	2.14	0.54
1:E:302:SER:HB2	1:E:305:ILE:HB	1.89	0.54
1:E:40:LEU:HD13	1:E:59:GLU:HG3	1.88	0.54
1:H:232:GLU:HG3	1:H:233:MET:H	1.72	0.54
1:K:219:PHE:CD1	1:K:319:GLN:HG3	2.41	0.54
1:K:219:PHE:HD1	1:K:319:GLN:HG3	1.72	0.54
1:L:227:ILE:N	1:L:227:ILE:HD12	2.22	0.54
1:L:215:LEU:O	1:L:322:ARG:HG3	2.07	0.54
1:N:293:ALA:HB1	1:N:298:GLY:O	2.08	0.54
2:R:8:ASP:C	2:R:57:LEU:HD11	2.27	0.54
2:S:16:GLU:HG3	2:S:19:THR:CG2	2.37	0.54
2:V:15:LYS:HB3	2:V:16:GLU:OE2	2.06	0.54
2:Z:40:VAL:HG21	2:Z:59:VAL:HG11	1.88	0.54
1:B:241:ALA:HA	1:B:271:VAL:HG11	1.89	0.54
1:D:245:LYS:HB3	1:D:246:PRO:HD2	1.88	0.54
1:E:18:ARG:NH1	1:E:18:ARG:HB3	2.22	0.54
1:E:325:ILE:HG13	1:E:330:THR:HG23	1.90	0.54
1:E:320:ALA:HA	1:E:335:GLY:HA2	1.89	0.54
1:E:443:ALA:O	1:E:447:MET:HG3	2.06	0.54

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:G:206:ASN:OD1	1:G:207:LYS:HG3	2.06	0.54
1:J:357:THR:HB	1:J:361:ASP:CB	2.38	0.54
1:K:217:SER:O	1:K:245:LYS:HD2	2.07	0.54
1:C:266:THR:HG22	1:C:273:VAL:H	1.73	0.54
1:E:236:VAL:HG21	1:E:312:ALA:HB3	1.90	0.54
1:E:237:LEU:O	1:E:241:ALA:N	2.37	0.54
1:F:349:ILE:O	1:F:353:ILE:HG13	2.07	0.54
1:L:204:PHE:CE1	1:L:266:THR:HG21	2.42	0.54
1:L:414:GLY:HA3	1:L:493:ILE:HG22	1.89	0.54
1:M:142:LYS:HE2	1:M:146:GLN:NE2	2.22	0.54
1:N:151:SER:HB3	1:N:399:ALA:HA	1.89	0.54
2:U:25:ILE:HD12	2:U:25:ILE:N	2.22	0.54
2:U:73:VAL:HG12	2:U:86:MET:CB	2.37	0.54
2:V:15:LYS:HB3	2:V:16:GLU:CD	2.27	0.54
2:V:43:VAL:HG22	2:V:44:GLY:N	2.22	0.54
2:V:78:ILE:HG13	2:V:79:ASP:OD2	2.07	0.54
1:A:122:LYS:HE2	1:A:429:LEU:HD11	1.89	0.54
1:D:169:VAL:HG23	1:D:173:GLY:HA3	1.88	0.54
1:D:172:GLU:N	1:D:172:GLU:OE1	2.30	0.54
1:G:448:GLU:O	1:G:452:ARG:HG2	2.07	0.54
1:L:429:LEU:HD12	1:L:430:ARG:H	1.72	0.54
1:N:363:GLU:O	1:N:367:GLU:HG3	2.08	0.54
1:N:8:PHE:O	1:N:11:ASP:HB3	2.07	0.54
2:S:20:LYS:HE3	2:S:29:GLY:N	2.22	0.54
2:U:65:VAL:HG12	2:U:94:ILE:HB	1.88	0.54
1:A:248:LEU:HD13	1:A:249:ILE:N	2.22	0.54
1:B:351:GLN:HG2	1:C:210:THR:HA	1.89	0.54
1:C:232:GLU:H	1:C:232:GLU:CD	2.09	0.54
1:D:230:ILE:HB	1:D:257:GLU:OE1	2.08	0.54
1:H:443:ALA:O	1:H:447:MET:HG3	2.07	0.54
1:L:279:PRO:HD2	1:L:285:ARG:HB2	1.89	0.54
1:M:201:SER:HB2	1:M:259:LEU:HD11	1.89	0.54
1:N:234:LEU:N	1:N:235:PRO:HD2	2.22	0.54
1:N:174:VAL:HB	1:N:376:VAL:HG13	1.89	0.54
2:Q:57:LEU:CD1	2:Q:59:VAL:HG22	2.37	0.54
2:S:48:ILE:HG13	2:S:52:GLY:H	1.73	0.54
2:U:50:GLU:C	2:U:52:GLY:H	2.11	0.54
2:V:78:ILE:HD13	2:2:37:ARG:HH12	1.72	0.54
1:J:234:LEU:HD11	2:X:26:VAL:HA	1.88	0.54
2:Y:20:LYS:CE	2:Y:22:ALA:HB3	2.38	0.54
1:C:218:PRO:HD2	1:C:320:ALA:O	2.07	0.54

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:D:279:PRO:HB2	1:D:288:MET:HE3	1.89	0.54
1:E:404:ARG:O	1:E:408:GLU:HG2	2.08	0.54
1:F:305:ILE:HD12	1:F:305:ILE:N	2.22	0.54
1:G:228:SER:HB3	1:G:255:GLU:HG2	1.89	0.54
1:G:23:LEU:HD23	1:G:60:ILE:HB	1.89	0.54
1:L:350:ARG:HD3	1:L:353:ILE:HD12	1.87	0.54
1:L:366:GLN:HA	1:L:369:VAL:HB	1.88	0.54
1:N:248:LEU:HD13	1:N:249:ILE:N	2.22	0.54
1:N:259:LEU:CD2	1:N:262:LEU:HD23	2.38	0.54
2:W:11:ILE:HB	2:W:85:ILE:CD1	2.38	0.54
2:Y:35:SER:OG	2:Y:37:ARG:O	2.25	0.54
1:A:183:LEU:HD22	1:A:384:ALA:HA	1.89	0.54
1:B:136:VAL:CG2	1:B:137:PRO:CD	2.86	0.54
1:B:311:LYS:O	1:B:311:LYS:HD3	2.08	0.54
1:D:236:VAL:CG2	1:D:312:ALA:HB3	2.37	0.54
1:E:227:ILE:HD12	1:E:227:ILE:N	2.23	0.54
1:H:215:LEU:HB2	1:H:323:VAL:HG22	1.89	0.54
1:I:284:ARG:HD2	1:I:364:LYS:NZ	2.22	0.54
1:I:6:VAL:HG22	1:I:521:VAL:HG22	1.89	0.54
1:N:254:VAL:CG1	1:N:259:LEU:HG	2.38	0.54
1:N:438:VAL:O	1:N:442:VAL:HG23	2.08	0.54
1:B:301:ILE:HG12	1:B:307:MET:HE2	1.88	0.54
1:B:218:PRO:HG3	1:B:323:VAL:CG2	2.37	0.54
1:D:305:ILE:HD12	1:D:305:ILE:N	2.23	0.54
1:D:326:ASN:OD1	1:D:329:THR:HB	2.07	0.54
1:E:183:LEU:HD22	1:E:384:ALA:CA	2.33	0.54
1:F:77:VAL:CG2	1:F:507:ALA:HA	2.38	0.54
1:G:227:ILE:N	1:G:227:ILE:HD12	2.22	0.54
1:H:237:LEU:HD21	1:H:271:VAL:HG21	1.90	0.54
1:J:222:LEU:HD22	1:J:300:VAL:HG22	1.90	0.54
1:J:357:THR:HB	1:J:361:ASP:HB2	1.89	0.54
1:K:368:ARG:O	1:K:372:LEU:HG	2.08	0.54
1:L:302:SER:CB	1:L:305:ILE:HB	2.38	0.54
1:L:309:LEU:HD23	1:L:309:LEU:H	1.73	0.54
1:K:517:THR:HG23	1:L:39:VAL:HG23	1.90	0.54
1:N:177:VAL:HG22	1:N:393:LYS:HG3	1.89	0.54
2:Y:7:HIS:O	2:Y:7:HIS:CD2	2.61	0.54
2:Z:25:ILE:N	2:Z:25:ILE:HD12	2.23	0.54
1:A:232:GLU:CG	1:A:234:LEU:HD23	2.37	0.54
1:A:306:GLY:CA	1:B:264:VAL:HG21	2.32	0.54
1:F:293:ALA:O	1:F:298:GLY:N	2.40	0.54

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:H:221:LEU:CD1	1:H:317:LEU:HD21	2.25	0.54
1:J:199:TYR:HE2	1:J:326:ASN:HA	1.73	0.54
1:L:488:MET:HE3	1:L:493:ILE:HG21	1.88	0.54
1:M:496:PRO:HG2	1:M:499:VAL:CG1	2.38	0.54
1:N:448:GLU:O	1:N:452:ARG:HG3	2.07	0.54
1:N:34:LYS:HD2	1:N:458:CYS:SG	2.48	0.54
2:P:73:VAL:HG11	2:P:84:LEU:HD23	1.90	0.54
2:Z:14:ARG:NH2	2:Z:34:LYS:HD3	2.23	0.54
2:2:17:VAL:CA	2:2:35:SER:HB2	2.38	0.54
1:A:227:ILE:HD12	1:A:227:ILE:N	2.23	0.54
1:A:247:LEU:HG	1:A:273:VAL:HG13	1.89	0.54
1:B:136:VAL:HG23	1:B:137:PRO:HD2	1.90	0.54
1:C:227:ILE:O	1:C:227:ILE:HG22	2.08	0.54
1:D:219:PHE:O	1:D:247:LEU:HD12	2.08	0.54
1:D:417:VAL:HA	1:D:420:ILE:CG2	2.37	0.54
1:K:196:ASP:O	1:K:197:ARG:HG2	2.07	0.54
1:K:225:LYS:HD2	1:K:252:GLU:OE1	2.08	0.54
1:K:322:ARG:HB3	1:K:333:ILE:HG22	1.89	0.54
1:L:230:ILE:CG1	1:L:309:LEU:HD21	2.36	0.54
1:M:151:SER:CB	1:M:399:ALA:HA	2.38	0.54
2:Q:36:THR:HG23	2:Q:37:ARG:HG3	1.90	0.54
2:S:6:LEU:HD12	2:S:6:LEU:O	2.08	0.54
2:U:14:ARG:NH1	2:U:84:LEU:HD21	2.22	0.54
1:A:252:GLU:HA	1:A:285:ARG:NH1	2.23	0.53
1:B:215:LEU:HB3	1:B:246:PRO:HB2	1.90	0.53
1:C:311:LYS:O	1:C:311:LYS:HD3	2.07	0.53
1:D:266:THR:HG22	1:D:271:VAL:O	2.08	0.53
1:D:199:TYR:OH	1:D:327:LYS:HG3	2.08	0.53
1:F:225:LYS:HB2	1:F:303:GLU:CD	2.29	0.53
1:F:386:GLU:O	1:F:389:MET:HB3	2.08	0.53
1:G:224:ASP:O	1:G:252:GLU:HB2	2.08	0.53
1:H:302:SER:OG	1:H:305:ILE:HB	2.08	0.53
1:H:406:ALA:HB1	1:H:411:VAL:CG1	2.37	0.53
1:J:496:PRO:HG2	1:J:499:VAL:HG11	1.90	0.53
1:K:18:ARG:HH11	1:K:18:ARG:HB3	1.72	0.53
1:M:34:LYS:HE3	1:M:480:ALA:O	2.08	0.53
1:M:5:ASP:HB2	1:M:524:LEU:HD23	1.87	0.53
1:N:40:LEU:HD13	1:N:59:GLU:HG3	1.90	0.53
2:O:94:ILE:O	2:P:3:ILE:HG22	2.08	0.53
2:Y:48:ILE:N	2:Y:48:ILE:HD12	2.23	0.53
1:A:92:ALA:HB2	1:A:503:ALA:HB1	1.90	0.53

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:B:136:VAL:HG22	1:B:137:PRO:HD2	1.88	0.53
1:H:220:ILE:HG22	1:H:222:LEU:CD1	2.38	0.53
1:K:261:THR:O	1:K:265:ASN:ND2	2.40	0.53
1:K:465:VAL:O	1:K:469:VAL:HG23	2.09	0.53
1:L:208:PRO:C	1:L:209:GLU:CD	2.46	0.53
1:M:409:GLU:CD	1:M:501:ARG:HH21	2.10	0.53
2:U:51:ASN:O	2:U:52:GLY:C	2.46	0.53
2:Y:7:HIS:HA	2:Y:45:ASN:H	1.73	0.53
2:1:97:ALA:HA	2:2:1:MET:SD	2.48	0.53
1:A:301:ILE:HD11	1:A:312:ALA:HB2	1.89	0.53
1:C:230:ILE:HB	1:C:232:GLU:OE1	2.08	0.53
1:C:232:GLU:HG3	1:C:310:GLU:CG	2.34	0.53
1:E:227:ILE:HG23	1:E:230:ILE:HB	1.89	0.53
1:I:233:MET:C	1:I:235:PRO:HD2	2.29	0.53
1:I:49:ILE:N	1:I:49:ILE:HD12	2.22	0.53
1:J:256:GLY:HA2	1:J:259:LEU:HB2	1.89	0.53
1:K:268:ARG:HB2	1:K:270:ILE:CD1	2.39	0.53
1:K:302:SER:OG	1:K:305:ILE:HB	2.08	0.53
1:L:406:ALA:HB2	1:L:496:PRO:HB3	1.90	0.53
1:M:333:ILE:O	1:M:334:ASP:HB2	2.08	0.53
1:M:357:THR:CG2	1:M:361:ASP:HB3	2.38	0.53
1:M:77:VAL:HG21	1:M:507:ALA:HA	1.90	0.53
1:N:219:PHE:HA	1:N:318:GLY:O	2.08	0.53
2:Q:17:VAL:HA	2:Q:35:SER:HB2	1.91	0.53
1:C:157:THR:O	1:C:161:LEU:HB2	2.08	0.53
1:D:4:LYS:HG3	1:E:59:GLU:O	2.08	0.53
1:F:208:PRO:HB2	1:F:212:ALA:HB3	1.89	0.53
1:I:90:THR:OG1	4:I:602:BEF:F1	2.10	0.53
1:J:23:LEU:HD23	1:J:60:ILE:HB	1.90	0.53
1:L:225:LYS:HB2	1:L:303:GLU:CD	2.28	0.53
1:L:417:VAL:HA	1:L:420:ILE:HG22	1.89	0.53
1:M:142:LYS:O	1:M:146:GLN:HG3	2.07	0.53
1:M:450:PRO:O	1:M:454:ILE:HG13	2.07	0.53
2:X:83:VAL:C	2:X:84:LEU:HD12	2.27	0.53
2:Y:5:PRO:CD	2:Y:42:ALA:HB1	2.38	0.53
1:A:182:GLY:C	1:A:184:GLN:H	2.11	0.53
1:A:321:LYS:HB2	1:A:334:ASP:HB3	1.89	0.53
1:A:496:PRO:HG2	1:A:499:VAL:HG13	1.90	0.53
1:B:177:VAL:HG21	1:B:397:GLU:CG	2.38	0.53
1:C:305:ILE:CG2	1:C:306:GLY:N	2.68	0.53
1:D:222:LEU:N	1:D:222:LEU:HD12	2.24	0.53

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:D:232:GLU:HG3	1:D:233:MET:H	1.73	0.53
1:G:202:PRO:O	1:G:205:ILE:HG13	2.08	0.53
1:H:440:ILE:O	1:H:444:LEU:HG	2.08	0.53
1:I:177:VAL:HG11	1:I:397:GLU:CG	2.39	0.53
1:J:305:ILE:HD11	1:K:267:MET:HE3	1.91	0.53
1:J:350:ARG:HD3	1:J:353:ILE:HD12	1.91	0.53
1:K:208:PRO:HG2	1:K:214:GLU:HG3	1.89	0.53
1:K:236:VAL:O	1:K:239:ALA:HB3	2.09	0.53
1:K:102:GLU:HG3	1:K:445:ARG:HD2	1.90	0.53
3:N:601:ADP:O3B	4:N:602:BEF:F3	2.17	0.53
2:Z:46:GLY:HA2	2:Z:57:LEU:HD11	1.91	0.53
2:Z:74:LYS:HE2	2:Z:75:SER:O	2.07	0.53
1:A:230:ILE:HG23	1:A:309:LEU:HD13	1.90	0.53
1:A:362:ARG:O	1:A:365:LEU:HG	2.08	0.53
1:B:270:ILE:HG23	1:B:271:VAL:N	2.23	0.53
1:F:233:MET:CG	1:F:234:LEU:CD2	2.80	0.53
1:F:360:TYR:HA	1:F:363:GLU:CG	2.37	0.53
1:G:225:LYS:HG3	1:G:226:LYS:N	2.23	0.53
1:G:199:TYR:CE2	1:G:327:LYS:HA	2.44	0.53
1:H:175:ILE:HA	1:H:377:ALA:O	2.08	0.53
1:M:69:MET:SD	1:N:41:ASP:HB2	2.48	0.53
2:R:12:VAL:HG12	2:R:40:VAL:CA	2.34	0.53
2:U:48:ILE:C	2:U:49:LEU:CD1	2.71	0.53
2:T:92:LEU:HA	2:U:9:ARG:HD2	1.90	0.53
2:W:17:VAL:HG23	2:W:19:THR:CA	2.39	0.53
2:W:9:ARG:HA	2:W:87:SER:HA	1.91	0.53
2:X:37:ARG:CG	2:X:66:ILE:HG23	2.30	0.53
1:D:25:ASP:HA	1:D:28:LYS:HE2	1.91	0.53
1:E:208:PRO:HB2	1:E:212:ALA:CB	2.36	0.53
1:F:142:LYS:O	1:F:146:GLN:HG3	2.08	0.53
1:F:455:VAL:HG13	1:F:460:GLU:HB2	1.90	0.53
1:I:304:GLU:C	1:I:305:ILE:HD12	2.28	0.53
1:I:456:LEU:HD13	1:I:462:PRO:HG3	1.90	0.53
1:J:138:CYS:HB2	1:J:411:VAL:HG13	1.90	0.53
1:K:280:GLY:O	1:K:285:ARG:NE	2.41	0.53
1:L:232:GLU:HB2	1:L:234:LEU:CD1	2.38	0.53
2:O:28:THR:HG22	2:O:29:GLY:O	2.09	0.53
2:U:11:ILE:CG2	2:U:42:ALA:HB3	2.39	0.53
2:W:19:THR:HG22	2:W:20:LYS:H	1.73	0.53
1:A:10:ASN:O	1:A:14:VAL:HG23	2.09	0.53
1:E:223:ALA:O	1:E:224:ASP:HB2	2.09	0.53

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:F:124:VAL:HG22	1:F:504:LEU:HD11	1.91	0.53
1:F:496:PRO:HG2	1:F:499:VAL:CG1	2.38	0.53
1:G:117:LYS:NZ	1:G:121:ASP:OD2	2.36	0.53
1:G:218:PRO:HA	1:G:246:PRO:O	2.09	0.53
1:G:291:ASP:OD1	1:G:292:ILE:HG13	2.08	0.53
1:H:305:ILE:N	1:H:305:ILE:HD12	2.23	0.53
1:H:42:LYS:HE2	1:H:48:THR:HG21	1.90	0.53
1:I:166:MET:HE1	1:I:407:VAL:HG21	1.91	0.53
1:J:368:ARG:O	1:J:372:LEU:HG	2.08	0.53
2:V:77:LYS:NZ	2:V:80:ASN:HA	2.24	0.53
1:D:392:LYS:O	1:D:396:VAL:HG23	2.09	0.53
1:E:124:VAL:HG13	1:E:504:LEU:HD12	1.89	0.53
1:E:234:LEU:HD11	1:E:238:GLU:OE1	2.08	0.53
1:H:346:VAL:HG13	1:H:369:VAL:HG13	1.91	0.53
1:I:355:GLU:HG3	1:I:356:ALA:H	1.74	0.53
1:J:174:VAL:HG12	1:J:376:VAL:HG22	1.91	0.53
1:J:181:THR:HG23	1:J:182:GLY:N	2.24	0.53
1:J:352:GLN:HA	1:J:355:GLU:HG2	1.90	0.53
1:K:227:ILE:HD12	1:K:233:MET:SD	2.49	0.53
1:L:95:LEU:O	1:L:99:ILE:HG13	2.09	0.53
1:N:230:ILE:O	1:N:309:LEU:HD22	2.09	0.53
1:N:74:VAL:O	1:N:77:VAL:HG12	2.09	0.53
2:P:3:ILE:O	2:P:3:ILE:HD12	2.08	0.53
2:R:48:ILE:HG12	2:R:54:VAL:CG1	2.32	0.53
1:A:220:ILE:HG13	1:A:296:THR:HG21	1.89	0.53
1:D:357:THR:CB	1:D:361:ASP:HB2	2.38	0.53
1:D:175:ILE:HG12	1:D:377:ALA:HB3	1.90	0.53
1:F:183:LEU:CD2	1:F:384:ALA:HA	2.39	0.53
1:F:479:ASN:ND2	1:F:493:ILE:HD11	2.24	0.53
1:H:215:LEU:O	1:H:322:ARG:HA	2.09	0.53
1:I:217:SER:N	1:I:218:PRO:HD3	2.24	0.53
1:J:203:TYR:OH	1:J:267:MET:HB2	2.08	0.53
1:N:227:ILE:HD12	1:N:227:ILE:N	2.24	0.53
2:P:37:ARG:NH2	2:Q:77:LYS:O	2.42	0.53
1:A:256:GLY:O	1:A:260:ALA:N	2.40	0.52
1:A:309:LEU:HD12	1:A:310:GLU:HB3	1.91	0.52
1:C:124:VAL:HG13	1:C:504:LEU:CD1	2.38	0.52
3:C:601:ADP:O2B	4:C:602:BEF:F1	2.17	0.52
1:D:368:ARG:O	1:D:372:LEU:HG	2.09	0.52
1:E:208:PRO:HG2	1:E:214:GLU:CG	2.38	0.52
1:F:218:PRO:C	1:F:219:PHE:CD1	2.75	0.52

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:G:13:ARG:HD2	1:G:104:LEU:HD22	1.90	0.52
1:H:232:GLU:CD	1:H:234:LEU:HD13	2.29	0.52
1:J:220:ILE:HG22	1:J:318:GLY:C	2.30	0.52
1:K:10:ASN:O	1:K:14:VAL:HG23	2.09	0.52
1:L:381:VAL:CG1	1:L:392:LYS:HB3	2.37	0.52
1:M:241:ALA:HA	1:M:271:VAL:CG1	2.21	0.52
1:N:124:VAL:HG21	1:N:508:ALA:CB	2.39	0.52
2:1:96:GLU:O	2:2:1:MET:HB2	2.09	0.52
1:A:219:PHE:HB3	1:A:317:LEU:HD13	1.92	0.52
1:C:510:VAL:O	1:C:514:MET:HB2	2.09	0.52
1:D:417:VAL:CA	1:D:420:ILE:HG22	2.37	0.52
1:G:349:ILE:O	1:G:353:ILE:HG13	2.10	0.52
1:H:221:LEU:HD13	1:H:317:LEU:CD2	2.26	0.52
1:I:218:PRO:HB3	1:I:246:PRO:HG2	1.90	0.52
1:L:305:ILE:HG22	1:L:306:GLY:H	1.75	0.52
1:N:496:PRO:HG2	1:N:499:VAL:HG11	1.92	0.52
2:1:17:VAL:HA	2:1:35:SER:N	2.24	0.52
1:B:218:PRO:HG3	1:B:323:VAL:HG13	1.91	0.52
1:B:5:ASP:HB2	1:B:524:LEU:HD23	1.92	0.52
1:E:448:GLU:O	1:E:452:ARG:HG2	2.10	0.52
1:G:422:VAL:O	1:G:426:LEU:HG	2.09	0.52
1:H:252:GLU:O	1:H:277:LYS:HG3	2.09	0.52
3:H:601:ADP:O3B	4:H:602:BEF:F3	2.18	0.52
1:M:219:PHE:O	1:M:247:LEU:HD12	2.10	0.52
1:M:25:ASP:OD1	1:M:97:GLN:NE2	2.43	0.52
2:T:40:VAL:HG22	2:T:41:LEU:N	2.24	0.52
2:Z:57:LEU:N	2:Z:57:LEU:HD12	2.24	0.52
2:1:36:THR:HB	2:1:67:PHE:O	2.10	0.52
1:A:346:VAL:O	1:A:350:ARG:HG2	2.08	0.52
1:A:364:LYS:HA	1:A:367:GLU:HB3	1.90	0.52
1:C:124:VAL:O	1:C:128:VAL:HG23	2.09	0.52
1:D:421:ARG:NH1	1:D:472:GLY:O	2.42	0.52
1:D:40:LEU:HD21	1:D:56:VAL:HG22	1.92	0.52
1:E:166:MET:CE	1:E:407:VAL:HG21	2.39	0.52
1:E:169:VAL:CG2	1:E:173:GLY:HA3	2.40	0.52
1:F:266:THR:CG2	1:F:273:VAL:H	2.23	0.52
1:F:313:THR:HG22	1:F:315:GLU:H	1.74	0.52
1:G:218:PRO:HG3	1:G:323:VAL:CG1	2.40	0.52
1:G:37:ASN:OD1	1:G:51:LYS:HB2	2.09	0.52
1:J:225:LYS:HG2	1:J:226:LYS:H	1.73	0.52
1:L:305:ILE:HG22	1:L:306:GLY:N	2.24	0.52

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:L:40:LEU:HD21	1:L:56:VAL:HG22	1.90	0.52
1:N:305:ILE:HG22	1:N:306:GLY:N	2.21	0.52
2:T:35:SER:OG	2:T:37:ARG:O	2.26	0.52
2:1:70:GLY:O	2:1:73:VAL:HG22	2.10	0.52
2:2:68:ASN:N	2:2:68:ASN:OD1	2.42	0.52
1:A:34:LYS:HE3	1:A:480:ALA:O	2.09	0.52
1:A:419:LEU:HD21	1:A:500:THR:HG23	1.91	0.52
1:D:260:ALA:O	1:D:264:VAL:HG23	2.09	0.52
1:E:365:LEU:HD23	1:E:369:VAL:HG23	1.89	0.52
1:E:90:THR:HB	3:E:601:ADP:O2B	2.10	0.52
1:F:225:LYS:CB	1:F:309:LEU:HD21	2.39	0.52
1:F:349:ILE:HD11	1:F:365:LEU:HD23	1.90	0.52
1:F:517:THR:HG23	1:G:39:VAL:HG23	1.91	0.52
1:I:419:LEU:HD21	1:I:500:THR:HG23	1.91	0.52
1:I:124:VAL:HG21	1:I:508:ALA:CB	2.39	0.52
1:J:245:LYS:CB	1:J:246:PRO:HD2	2.38	0.52
1:J:455:VAL:HG21	1:J:465:VAL:HG11	1.90	0.52
1:L:363:GLU:HA	1:L:366:GLN:OE1	2.08	0.52
1:L:40:LEU:HD13	1:L:59:GLU:CG	2.33	0.52
1:M:157:THR:O	1:M:161:LEU:HB2	2.09	0.52
1:L:305:ILE:HA	1:M:264:VAL:HG12	1.92	0.52
1:M:252:GLU:HA	1:M:285:ARG:NH1	2.25	0.52
1:M:519:CYS:CB	1:N:38:VAL:HG13	2.39	0.52
1:N:510:VAL:O	1:N:514:MET:HB2	2.09	0.52
1:E:219:PHE:HB2	1:E:247:LEU:HD12	1.92	0.52
1:G:228:SER:HB3	1:G:255:GLU:CG	2.39	0.52
1:H:230:ILE:HG23	1:H:231:ARG:H	1.75	0.52
1:I:225:LYS:HG3	1:I:226:LYS:N	2.25	0.52
1:I:239:ALA:HB1	1:I:314:LEU:HD21	1.92	0.52
1:N:266:THR:HA	1:N:271:VAL:O	2.10	0.52
2:S:50:GLU:CD	2:S:51:ASN:H	2.12	0.52
2:Y:15:LYS:HB2	2:Y:38:GLY:HA2	1.91	0.52
1:C:149:THR:OG1	1:C:159:GLY:HA3	2.08	0.52
1:C:412:VAL:HG13	1:C:497:THR:OG1	2.08	0.52
1:G:358:SER:HA	1:G:362:ARG:HH21	1.74	0.52
1:A:387:VAL:HG13	1:G:76:GLU:OE1	2.09	0.52
1:G:27:VAL:CG1	1:G:90:THR:HG23	2.29	0.52
1:H:325:ILE:HG12	1:H:330:THR:HG23	1.91	0.52
1:K:218:PRO:HD2	1:K:320:ALA:O	2.09	0.52
1:L:519:CYS:HB3	1:M:38:VAL:HG22	1.91	0.52
1:M:449:ALA:HB3	1:M:450:PRO:HD3	1.91	0.52

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:N:256:GLY:O	1:N:260:ALA:N	2.43	0.52
2:X:94:ILE:HD12	2:X:96:GLU:HG3	1.91	0.52
2:Z:93:ALA:CA	2:1:5:PRO:HA	2.39	0.52
1:B:77:VAL:CG2	1:B:507:ALA:HA	2.35	0.52
1:A:463:SER:HB3	1:H:464:VAL:HG23	1.91	0.52
1:M:261:THR:O	1:M:264:VAL:HG22	2.10	0.52
1:N:363:GLU:HG3	1:N:364:LYS:N	2.25	0.52
1:N:48:THR:HG22	1:N:390:LYS:NZ	2.24	0.52
2:R:20:LYS:HB2	2:R:22:ALA:H	1.75	0.52
2:S:12:VAL:HA	2:S:40:VAL:HA	1.92	0.52
2:S:46:GLY:HA2	2:S:57:LEU:CD1	2.40	0.52
2:V:88:GLU:HA	2:V:91:ILE:HD13	1.91	0.52
1:B:351:GLN:HG2	1:C:210:THR:CB	2.40	0.52
1:C:124:VAL:HG21	1:C:508:ALA:HB1	1.92	0.52
1:D:350:ARG:HD2	1:D:353:ILE:CD1	2.38	0.52
1:E:214:GLU:HA	1:E:323:VAL:O	2.10	0.52
1:F:28:LYS:HG2	1:F:94:VAL:HG22	1.92	0.52
1:G:199:TYR:HE1	1:G:202:PRO:HB3	1.75	0.52
1:I:396:VAL:O	1:I:400:LEU:HB2	2.10	0.52
1:J:236:VAL:HG11	1:J:312:ALA:HB3	1.92	0.52
1:K:116:LEU:O	1:K:120:ILE:HG13	2.10	0.52
1:L:39:VAL:HG22	1:L:49:ILE:HG12	1.92	0.52
1:M:325:ILE:CG1	1:M:330:THR:HG23	2.29	0.52
2:Q:94:ILE:O	2:Q:94:ILE:HG13	2.10	0.52
2:Z:60:LYS:HB3	2:Z:63:ASP:CG	2.30	0.52
1:E:415:GLY:N	3:E:601:ADP:O2'	2.34	0.52
1:F:509:SER:OG	1:G:385:THR:HG22	2.10	0.52
1:G:221:LEU:N	1:G:221:LEU:HD12	2.25	0.52
1:L:221:LEU:O	1:L:221:LEU:HD12	2.10	0.52
1:L:77:VAL:HG21	1:L:507:ALA:HA	1.92	0.52
1:M:147:VAL:HG12	1:M:494:LEU:HD12	1.91	0.52
1:M:201:SER:HB3	1:M:204:PHE:CZ	2.44	0.52
1:M:510:VAL:O	1:M:514:MET:HB2	2.10	0.52
2:U:39:GLU:HA	2:U:64:ILE:HA	1.91	0.52
1:A:254:VAL:HG11	1:A:259:LEU:HD13	1.92	0.51
1:D:219:PHE:HD2	1:D:240:VAL:HG22	1.75	0.51
1:D:200:LEU:HD13	1:D:276:VAL:HA	1.91	0.51
1:E:222:LEU:HD12	1:E:222:LEU:O	2.10	0.51
1:F:291:ASP:OD1	1:F:292:ILE:N	2.43	0.51
1:H:177:VAL:CG1	1:H:397:GLU:HG2	2.39	0.51
1:K:197:ARG:HH12	1:K:277:LYS:HD2	1.75	0.51

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:M:116:LEU:O	1:M:120:ILE:HG13	2.09	0.51
1:M:415:GLY:N	3:M:601:ADP:O2'	2.41	0.51
2:O:49:LEU:CD1	2:O:50:GLU:H	2.22	0.51
2:Q:51:ASN:O	2:Q:53:GLU:N	2.43	0.51
2:U:85:ILE:HD12	2:U:85:ILE:O	2.10	0.51
1:B:270:ILE:HG12	2:P:25:ILE:CD1	2.38	0.51
1:B:85:ALA:O	1:B:401:HIS:HB3	2.11	0.51
1:C:255:GLU:N	1:C:255:GLU:OE1	2.42	0.51
1:D:34:LYS:HD2	1:D:458:CYS:SG	2.50	0.51
1:H:220:ILE:HD12	1:H:220:ILE:N	2.25	0.51
1:H:227:ILE:HA	1:H:230:ILE:HG22	1.93	0.51
1:H:301:ILE:HD13	1:H:307:MET:HE1	1.91	0.51
1:I:115:ASP:O	1:I:436:GLN:HG2	2.10	0.51
1:J:103:GLY:O	1:J:107:VAL:HG23	2.10	0.51
1:J:265:ASN:OD1	1:J:266:THR:N	2.44	0.51
1:J:419:LEU:H	1:J:419:LEU:HD12	1.74	0.51
1:K:333:ILE:O	1:K:334:ASP:HB2	2.11	0.51
1:K:5:ASP:HB2	1:K:524:LEU:CD2	2.38	0.51
1:L:136:VAL:CG2	1:L:137:PRO:HD2	2.40	0.51
1:L:229:ASN:OD1	1:L:230:ILE:HD12	2.09	0.51
1:M:325:ILE:N	1:M:325:ILE:HD12	2.24	0.51
2:O:64:ILE:CG2	2:O:95:VAL:HB	2.30	0.51
2:Q:45:ASN:OD1	2:Q:46:GLY:N	2.42	0.51
2:R:11:ILE:CG2	2:R:42:ALA:HB3	2.40	0.51
2:U:10:VAL:HG11	2:U:91:ILE:CD1	2.39	0.51
2:Y:40:VAL:HG11	2:Y:60:LYS:O	2.11	0.51
2:2:49:LEU:HB2	2:2:52:GLY:HA2	1.92	0.51
1:A:450:PRO:O	1:A:454:ILE:HG13	2.10	0.51
1:B:255:GLU:HB3	1:B:257:GLU:OE1	2.09	0.51
1:C:302:SER:HB2	1:C:305:ILE:HD13	1.92	0.51
1:E:199:TYR:CA	1:E:276:VAL:HG12	2.37	0.51
1:F:245:LYS:CB	1:F:246:PRO:HD2	2.35	0.51
1:G:221:LEU:HD13	1:G:249:ILE:CD1	2.30	0.51
1:I:228:SER:CB	1:I:255:GLU:HB2	2.37	0.51
1:J:199:TYR:HA	1:J:276:VAL:HG12	1.91	0.51
1:J:305:ILE:HG23	1:J:306:GLY:N	2.25	0.51
1:J:350:ARG:HA	1:J:353:ILE:CD1	2.40	0.51
1:J:169:VAL:HB	1:J:375:GLY:O	2.10	0.51
1:L:247:LEU:HD22	1:L:248:LEU:H	1.74	0.51
1:L:4:LYS:HG3	1:M:59:GLU:O	2.10	0.51
1:M:199:TYR:HD2	1:M:325:ILE:HG22	1.75	0.51

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:M:88:GLY:HA2	3:M:601:ADP:PB	2.49	0.51
1:N:10:ASN:O	1:N:14:VAL:HG23	2.11	0.51
1:N:197:ARG:HG2	1:N:277:LYS:CB	2.39	0.51
2:Q:48:ILE:HD12	2:Q:48:ILE:N	2.25	0.51
2:T:13:LYS:HG3	2:T:83:VAL:HG12	1.91	0.51
2:V:60:LYS:HG3	2:V:63:ASP:CG	2.30	0.51
2:Z:50:GLU:O	2:Z:51:ASN:OD1	2.29	0.51
2:1:6:LEU:HB3	2:1:7:HIS:HB2	1.91	0.51
1:A:291:ASP:HB3	1:A:372:LEU:CD2	2.40	0.51
1:B:285:ARG:O	1:B:289:LEU:HG	2.10	0.51
1:C:285:ARG:HG3	1:C:286:LYS:N	2.25	0.51
1:C:350:ARG:HH12	1:C:369:VAL:CG1	2.19	0.51
1:D:279:PRO:HB2	1:D:288:MET:CE	2.41	0.51
1:D:381:VAL:HG13	1:D:392:LYS:HG2	1.90	0.51
1:E:256:GLY:HA2	1:E:259:LEU:HD12	1.91	0.51
1:E:419:LEU:O	1:E:422:VAL:HG22	2.11	0.51
1:G:215:LEU:HB2	1:G:323:VAL:CG2	2.40	0.51
1:I:414:GLY:HA2	1:I:495:ASP:OD2	2.11	0.51
1:M:10:ASN:O	1:M:14:VAL:HG23	2.11	0.51
1:M:279:PRO:HB2	1:M:288:MET:HE3	1.92	0.51
1:N:199:TYR:HE1	1:N:202:PRO:HA	1.69	0.51
1:N:222:LEU:N	1:N:222:LEU:HD12	2.26	0.51
2:S:20:LYS:HE3	2:S:29:GLY:H	1.75	0.51
2:V:11:ILE:O	2:V:41:LEU:HB2	2.10	0.51
1:A:199:TYR:OH	1:A:205:ILE:HD11	2.10	0.51
1:A:216:GLU:HB2	1:A:245:LYS:NZ	2.26	0.51
1:B:319:GLN:O	1:B:336:VAL:HG23	2.11	0.51
1:D:392:LYS:HD2	1:D:395:ARG:NH2	2.25	0.51
1:D:450:PRO:O	1:D:454:ILE:HG13	2.11	0.51
1:E:199:TYR:OH	1:E:327:LYS:HG3	2.11	0.51
1:E:85:ALA:HB1	1:E:499:VAL:HG12	1.92	0.51
1:K:305:ILE:HD12	1:K:305:ILE:N	2.25	0.51
1:J:519:CYS:HB3	1:K:38:VAL:HG22	1.90	0.51
1:L:233:MET:HE2	1:L:236:VAL:HB	1.91	0.51
1:M:414:GLY:HA2	1:M:495:ASP:OD2	2.09	0.51
1:M:405:ALA:HB1	1:M:498:LYS:HD3	1.92	0.51
1:M:37:ASN:OD1	1:M:51:LYS:HB2	2.09	0.51
2:P:12:VAL:HG13	2:P:39:GLU:O	2.10	0.51
2:R:48:ILE:HG23	2:R:54:VAL:CG1	2.41	0.51
2:S:25:ILE:N	2:S:25:ILE:HD12	2.25	0.51
2:W:20:LYS:NZ	2:X:77:LYS:HB2	2.26	0.51

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:Y:17:VAL:HG23	2:Y:33:ALA:HB3	1.92	0.51
2:Y:15:LYS:HD2	2:Y:37:ARG:HB3	1.91	0.51
1:A:231:ARG:HB3	1:A:258:ALA:HB1	1.92	0.51
1:B:230:ILE:O	1:B:230:ILE:HG23	2.11	0.51
1:B:342:ILE:O	1:B:346:VAL:HG23	2.11	0.51
1:B:393:LYS:O	1:B:397:GLU:HG3	2.11	0.51
1:B:90:THR:HB	3:B:601:ADP:O1B	2.11	0.51
1:E:278:ALA:HB1	1:E:285:ARG:HD2	1.91	0.51
1:F:245:LYS:HB3	1:F:246:PRO:CD	2.36	0.51
1:G:223:ALA:HB3	1:G:224:ASP:CA	2.40	0.51
1:H:176:THR:OG1	1:H:378:VAL:HG22	2.10	0.51
1:H:496:PRO:HG2	1:H:499:VAL:CG1	2.39	0.51
1:J:220:ILE:HG21	1:J:296:THR:OG1	2.10	0.51
1:M:175:ILE:HA	1:M:377:ALA:O	2.10	0.51
1:N:13:ARG:HD3	1:N:104:LEU:HD22	1.93	0.51
2:P:17:VAL:HG12	2:P:35:SER:CA	2.41	0.51
2:Q:11:ILE:O	2:Q:41:LEU:HB2	2.10	0.51
2:R:20:LYS:HB2	2:R:22:ALA:HB3	1.92	0.51
2:R:77:LYS:HG3	2:R:81:GLU:O	2.11	0.51
2:Z:14:ARG:NH1	2:Z:84:LEU:HD21	2.25	0.51
2:Z:92:LEU:HD22	2:1:9:ARG:HD3	1.93	0.51
2:1:57:LEU:H	2:1:57:LEU:HD22	1.76	0.51
1:B:228:SER:HA	1:B:255:GLU:HB2	1.92	0.51
1:C:205:ILE:HA	1:C:213:VAL:HG22	1.91	0.51
1:F:40:LEU:HD21	1:F:56:VAL:HG22	1.93	0.51
1:J:34:LYS:NZ	1:J:483:GLU:OE2	2.38	0.51
1:K:42:LYS:HE2	1:K:48:THR:OG1	2.11	0.51
1:L:357:THR:HG22	1:L:359:ASP:HB2	1.91	0.51
1:M:218:PRO:HD2	1:M:320:ALA:O	2.11	0.51
1:N:223:ALA:HA	1:N:301:ILE:O	2.11	0.51
2:U:83:VAL:HG12	2:U:84:LEU:N	2.25	0.51
2:V:49:LEU:HD12	2:W:50:GLU:HG3	1.92	0.51
2:V:57:LEU:C	2:V:58:ASP:CG	2.70	0.51
2:V:6:LEU:HD11	2:2:91:ILE:O	2.11	0.51
2:Y:36:THR:C	2:Y:37:ARG:HD2	2.30	0.51
1:A:291:ASP:OD2	1:A:368:ARG:HD3	2.11	0.51
1:A:355:GLU:HG3	1:A:356:ALA:N	2.26	0.51
1:A:77:VAL:HB	1:A:510:VAL:HG21	1.93	0.51
1:B:42:LYS:HE2	1:B:48:THR:HG21	1.92	0.51
1:F:17:LEU:HG	1:F:21:ASN:ND2	2.25	0.51
1:F:234:LEU:HB2	1:F:235:PRO:HD3	1.91	0.51

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:H:207:LYS:HB3	1:H:208:PRO:CD	2.33	0.51
1:H:241:ALA:CB	2:V:25:ILE:HD12	2.38	0.51
1:H:333:ILE:O	1:H:334:ASP:HB2	2.11	0.51
1:I:199:TYR:HA	1:I:276:VAL:HG12	1.93	0.51
1:I:220:ILE:HD12	1:I:220:ILE:N	2.25	0.51
1:I:230:ILE:C	1:I:232:GLU:H	2.12	0.51
1:I:234:LEU:N	1:I:235:PRO:HD2	2.25	0.51
1:N:227:ILE:HG23	1:N:231:ARG:CG	2.40	0.51
1:N:309:LEU:HD12	1:N:310:GLU:N	2.26	0.51
2:T:12:VAL:HA	2:T:40:VAL:HA	1.91	0.51
1:I:237:LEU:HD23	2:W:27:LEU:CD2	2.40	0.51
2:W:78:ILE:HG13	2:W:79:ASP:N	2.20	0.51
2:Z:7:HIS:C	2:Z:46:GLY:O	2.50	0.51
1:B:103:GLY:O	1:B:107:VAL:HG23	2.10	0.51
1:B:406:ALA:HB2	1:B:496:PRO:HB3	1.92	0.51
1:G:223:ALA:HB3	1:G:224:ASP:HA	1.93	0.51
1:H:88:GLY:N	4:H:602:BEF:F3	2.34	0.51
1:L:149:THR:HG22	1:L:155:ASP:O	2.10	0.51
1:M:309:LEU:CB	1:M:310:GLU:OE1	2.58	0.51
1:N:233:MET:HB3	1:N:236:VAL:HB	1.93	0.51
1:N:115:ASP:O	1:N:436:GLN:HG2	2.11	0.51
2:Q:43:VAL:HG22	2:Q:44:GLY:O	2.10	0.51
2:T:37:ARG:HH12	2:U:78:ILE:HG22	1.76	0.51
2:U:50:GLU:O	2:U:52:GLY:N	2.44	0.51
1:I:237:LEU:HD23	2:W:27:LEU:HD22	1.93	0.51
2:1:65:VAL:HG12	2:1:94:ILE:CG2	2.35	0.51
2:1:14:ARG:NH2	2:1:69:ASP:OD2	2.44	0.51
1:A:247:LEU:HG	1:A:273:VAL:HG11	1.92	0.51
1:A:278:ALA:HB3	1:A:285:ARG:NE	2.26	0.51
1:A:309:LEU:HD12	1:A:310:GLU:CB	2.37	0.51
1:A:174:VAL:HG13	1:A:370:ALA:HB1	1.92	0.51
1:B:348:GLN:HA	1:C:209:GLU:HG2	1.93	0.51
1:B:465:VAL:O	1:B:469:VAL:HG23	2.11	0.51
1:C:338:GLU:O	1:C:342:ILE:HD13	2.11	0.51
1:D:90:THR:HB	3:D:601:ADP:O2B	2.11	0.51
1:H:282:GLY:HA2	1:H:285:ARG:HH21	1.77	0.51
1:H:406:ALA:HB2	1:H:496:PRO:HB3	1.93	0.51
1:J:404:ARG:O	1:J:408:GLU:HG2	2.11	0.51
1:L:107:VAL:HG13	1:L:113:PRO:HG3	1.92	0.51
1:M:200:LEU:CD2	1:M:254:VAL:H	2.24	0.51
1:M:200:LEU:HD11	1:M:277:LYS:CG	2.41	0.51

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:M:205:ILE:HA	1:M:213:VAL:HG22	1.93	0.51
1:M:323:VAL:HG12	1:M:332:ILE:HG12	1.93	0.51
1:N:177:VAL:CG2	1:N:396:VAL:HG11	2.42	0.51
2:O:95:VAL:CB	2:P:3:ILE:CG2	2.87	0.51
2:R:95:VAL:HA	2:S:3:ILE:HG22	1.92	0.51
2:W:84:LEU:O	2:W:84:LEU:HD12	2.10	0.51
2:X:25:ILE:HD12	2:X:25:ILE:N	2.26	0.51
1:E:218:PRO:HA	1:E:246:PRO:O	2.11	0.50
1:E:248:LEU:HD22	1:E:323:VAL:CG1	2.34	0.50
1:F:102:GLU:OE1	1:F:445:ARG:NH1	2.39	0.50
1:F:419:LEU:HD11	1:F:500:THR:CG2	2.39	0.50
1:F:405:ALA:HB1	1:F:498:LYS:HD3	1.92	0.50
1:H:232:GLU:CG	1:H:233:MET:H	2.22	0.50
1:H:510:VAL:O	1:H:514:MET:HB2	2.11	0.50
1:I:496:PRO:O	1:I:499:VAL:HG22	2.11	0.50
1:J:231:ARG:HH21	1:J:261:THR:HG21	1.76	0.50
1:K:102:GLU:OE1	1:K:445:ARG:NH1	2.41	0.50
1:K:77:VAL:CG1	1:K:510:VAL:HG21	2.41	0.50
1:M:197:ARG:CG	1:M:277:LYS:HB2	2.41	0.50
1:M:215:LEU:HD22	1:M:246:PRO:HB3	1.93	0.50
1:N:247:LEU:HB3	1:N:273:VAL:HG13	1.93	0.50
2:Q:65:VAL:HG12	2:Q:94:ILE:HG22	1.93	0.50
2:Y:5:PRO:HD3	2:Y:42:ALA:HB1	1.93	0.50
1:C:208:PRO:HB2	1:C:212:ALA:CB	2.26	0.50
1:C:219:PHE:H	1:C:247:LEU:HA	1.75	0.50
1:C:197:ARG:NE	1:C:277:LYS:HD2	2.18	0.50
1:D:288:MET:HG3	1:D:368:ARG:HE	1.75	0.50
1:D:381:VAL:HG13	1:D:392:LYS:CG	2.42	0.50
1:D:381:VAL:CG1	1:D:392:LYS:HB3	2.41	0.50
1:E:245:LYS:HE2	1:E:246:PRO:CD	2.38	0.50
1:E:20:VAL:HG13	1:E:74:VAL:HG11	1.93	0.50
1:F:414:GLY:HA2	1:F:495:ASP:CG	2.32	0.50
1:G:199:TYR:HE2	1:G:327:LYS:HG3	1.74	0.50
1:J:233:MET:O	1:J:236:VAL:N	2.42	0.50
1:L:248:LEU:HD13	1:L:249:ILE:N	2.25	0.50
1:M:225:LYS:HG3	1:M:226:LYS:N	2.26	0.50
1:N:220:ILE:N	1:N:220:ILE:HD12	2.27	0.50
2:V:57:LEU:CD1	2:V:59:VAL:HG22	2.41	0.50
2:X:57:LEU:HD23	2:X:88:GLU:HB2	1.93	0.50
1:B:200:LEU:HD11	1:B:277:LYS:HB2	1.94	0.50
1:B:496:PRO:HG2	1:B:499:VAL:HG13	1.94	0.50

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:E:117:LYS:NZ	1:E:121:ASP:OD2	2.43	0.50
1:E:138:CYS:SG	1:E:144:ILE:HD13	2.51	0.50
1:E:219:PHE:CE1	1:E:319:GLN:HG2	2.46	0.50
1:E:455:VAL:HG21	1:E:465:VAL:HG11	1.93	0.50
1:F:225:LYS:HB2	1:F:303:GLU:OE2	2.10	0.50
1:F:353:ILE:HD11	1:F:369:VAL:HG21	1.93	0.50
1:H:87:ASP:OD2	1:H:151:SER:OG	2.27	0.50
1:I:177:VAL:HG11	1:I:397:GLU:HG3	1.93	0.50
1:I:305:ILE:N	1:I:305:ILE:HD12	2.27	0.50
1:I:405:ALA:HB1	1:I:498:LYS:HB3	1.92	0.50
1:L:350:ARG:HD3	1:L:353:ILE:HD11	1.92	0.50
1:L:429:LEU:HD12	1:L:430:ARG:N	2.26	0.50
1:L:436:GLN:O	1:L:440:ILE:HG13	2.11	0.50
1:M:267:MET:HA	1:M:267:MET:CE	2.42	0.50
1:M:367:GLU:HG2	1:M:371:LYS:NZ	2.25	0.50
1:N:415:GLY:N	3:N:601:ADP:O2'	2.29	0.50
2:O:17:VAL:HG23	2:O:34:LYS:HA	1.94	0.50
2:Q:60:LYS:HZ1	2:Q:63:ASP:HB3	1.75	0.50
2:W:1:MET:HE2	2:W:79:ASP:CB	2.33	0.50
1:A:267:MET:CE	1:A:267:MET:HA	2.41	0.50
1:A:267:MET:HG3	1:G:305:ILE:CG1	2.37	0.50
1:A:478:TYR:HD1	1:A:485:TYR:CE1	2.30	0.50
1:C:256:GLY:HA2	1:C:259:LEU:HB2	1.92	0.50
1:D:228:SER:OG	1:D:255:GLU:HG2	2.12	0.50
1:D:349:ILE:O	1:D:353:ILE:HG13	2.11	0.50
1:E:116:LEU:O	1:E:120:ILE:HG13	2.11	0.50
1:J:241:ALA:HA	1:J:271:VAL:CG1	2.42	0.50
1:J:301:ILE:HA	1:J:307:MET:HE2	1.94	0.50
1:J:172:GLU:HB3	1:J:366:GLN:OE1	2.12	0.50
1:J:450:PRO:O	1:J:454:ILE:HG13	2.10	0.50
1:K:220:ILE:HD12	1:K:220:ILE:N	2.27	0.50
1:M:268:ARG:N	1:M:269:GLY:HA2	2.26	0.50
1:N:90:THR:HB	3:N:601:ADP:O2B	2.12	0.50
2:1:18:GLU:HB3	2:1:28:THR:HB	1.93	0.50
1:B:218:PRO:CB	1:B:248:LEU:HD23	2.41	0.50
1:B:6:VAL:CG1	1:B:521:VAL:HG22	2.34	0.50
1:C:8:PHE:HE1	1:D:26:ALA:HA	1.77	0.50
3:D:601:ADP:O3B	4:D:602:BEF:F3	2.19	0.50
1:E:7:LYS:HG3	1:E:66:PHE:CE2	2.46	0.50
1:F:311:LYS:NZ	1:F:313:THR:OG1	2.35	0.50
1:G:247:LEU:HD23	1:G:273:VAL:HG22	1.93	0.50

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:H:367:GLU:O	1:H:371:LYS:HG3	2.12	0.50
1:K:343:GLN:HA	1:K:346:VAL:HG22	1.93	0.50
1:L:213:VAL:O	1:L:324:VAL:HA	2.12	0.50
1:L:352:GLN:HE21	1:M:209:GLU:CD	2.15	0.50
2:Q:50:GLU:CG	2:Q:51:ASN:N	2.74	0.50
2:T:53:GLU:CG	2:T:54:VAL:N	2.75	0.50
2:2:28:THR:HG22	2:2:29:GLY:O	2.12	0.50
1:A:220:ILE:HG22	1:A:222:LEU:HD11	1.93	0.50
1:A:234:LEU:O	1:A:238:GLU:N	2.34	0.50
1:C:367:GLU:O	1:C:371:LYS:HG3	2.12	0.50
1:D:166:MET:HE1	1:D:407:VAL:HG21	1.93	0.50
1:F:348:GLN:OE1	1:F:352:GLN:NE2	2.44	0.50
1:G:450:PRO:O	1:G:454:ILE:HG13	2.12	0.50
1:H:174:VAL:CG2	1:H:376:VAL:HG22	2.41	0.50
1:I:240:VAL:HG12	1:I:245:LYS:O	2.11	0.50
1:I:347:ALA:O	1:I:350:ARG:HB2	2.12	0.50
1:J:303:GLU:CD	1:J:309:LEU:HD21	2.32	0.50
1:J:381:VAL:CG2	1:J:396:VAL:HG21	2.42	0.50
1:L:179:ASP:OD1	1:L:393:LYS:HD2	2.11	0.50
1:N:398:ASP:OD2	4:N:602:BEF:F1	2.20	0.50
2:V:49:LEU:HD13	2:V:50:GLU:HB2	1.94	0.50
2:1:91:ILE:HG22	2:2:6:LEU:HD12	1.92	0.50
1:D:200:LEU:HD13	1:D:275:ALA:O	2.11	0.50
1:E:28:LYS:HG2	1:E:94:VAL:HG22	1.93	0.50
1:F:392:LYS:O	1:F:396:VAL:HG23	2.11	0.50
1:G:221:LEU:HD11	1:G:317:LEU:HD22	1.94	0.50
1:H:17:LEU:HD13	1:H:104:LEU:HD12	1.94	0.50
1:H:233:MET:HG2	1:H:233:MET:O	2.10	0.50
1:I:313:THR:HG22	1:I:314:LEU:N	2.27	0.50
1:K:207:LYS:HG2	1:K:208:PRO:CD	2.21	0.50
1:K:414:GLY:HA2	1:K:495:ASP:OD2	2.12	0.50
1:K:23:LEU:CD2	1:K:60:ILE:HB	2.41	0.50
1:M:124:VAL:HG13	1:M:504:LEU:HD12	1.92	0.50
1:N:197:ARG:CG	1:N:277:LYS:HB3	2.41	0.50
1:A:215:LEU:HB2	1:A:323:VAL:CG2	2.42	0.50
1:B:285:ARG:NH1	1:B:289:LEU:HD11	2.27	0.50
1:E:291:ASP:OD1	1:E:345:ARG:NH2	2.39	0.50
1:F:168:LYS:CG	1:F:189:VAL:HG11	2.36	0.50
1:G:215:LEU:HD22	1:G:246:PRO:HB2	1.93	0.50
1:H:14:VAL:O	1:H:18:ARG:HG3	2.11	0.50
1:H:183:LEU:N	1:H:183:LEU:HD12	2.27	0.50

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:H:415:GLY:N	3:H:601:ADP:O2'	2.39	0.50
1:I:157:THR:O	1:I:161:LEU:HD13	2.11	0.50
1:J:234:LEU:HA	1:J:237:LEU:HB3	1.94	0.50
1:K:17:LEU:HD13	1:K:104:LEU:HD12	1.94	0.50
1:M:175:ILE:HG12	1:M:377:ALA:HB3	1.93	0.50
1:M:409:GLU:OE1	1:M:498:LYS:HA	2.12	0.50
1:N:208:PRO:HB2	1:N:212:ALA:HB3	1.94	0.50
1:N:197:ARG:HG2	1:N:277:LYS:N	2.27	0.50
2:1:57:LEU:N	2:1:57:LEU:HD22	2.27	0.50
1:C:246:PRO:HA	1:C:272:LYS:O	2.12	0.50
1:D:313:THR:HG21	1:D:315:GLU:HG2	1.93	0.50
1:E:142:LYS:O	1:E:146:GLN:HG3	2.11	0.50
1:H:232:GLU:HA	1:H:234:LEU:CD1	2.42	0.50
1:H:248:LEU:HD13	1:H:249:ILE:N	2.26	0.50
1:I:359:ASP:O	1:I:360:TYR:HB3	2.11	0.50
1:J:360:TYR:O	1:J:363:GLU:HG2	2.12	0.50
1:K:155:ASP:OD1	1:K:157:THR:OG1	2.20	0.50
1:L:417:VAL:O	1:L:421:ARG:HB2	2.12	0.50
1:L:405:ALA:HB1	1:L:498:LYS:HD3	1.94	0.50
1:M:233:MET:O	1:M:234:LEU:HB2	2.12	0.50
1:N:261:THR:HB	2:2:29:GLY:N	2.21	0.50
1:N:342:ILE:O	1:N:346:VAL:HG23	2.12	0.50
2:O:95:VAL:CA	2:P:3:ILE:HG21	2.28	0.50
2:O:97:ALA:HB2	2:P:1:MET:CB	2.42	0.50
2:S:39:GLU:HG2	2:S:39:GLU:O	2.11	0.50
1:A:191:GLU:N	1:A:191:GLU:OE1	2.36	0.49
1:A:40:LEU:HD11	1:A:56:VAL:HA	1.93	0.49
1:B:234:LEU:O	1:B:237:LEU:N	2.40	0.49
1:B:479:ASN:HB3	1:B:484:GLU:O	2.12	0.49
1:D:6:VAL:HG22	1:D:521:VAL:HG22	1.94	0.49
1:E:254:VAL:HG12	1:E:259:LEU:HG	1.93	0.49
1:F:195:PHE:CE1	1:F:330:THR:HB	2.46	0.49
1:F:420:ILE:HD11	1:F:451:LEU:CB	2.39	0.49
1:G:409:GLU:OE2	1:G:501:ARG:NH2	2.27	0.49
1:G:496:PRO:HG2	1:G:499:VAL:CG1	2.42	0.49
1:G:31:LEU:HD12	3:G:601:ADP:O2A	2.11	0.49
1:H:241:ALA:HA	1:H:271:VAL:HG11	1.94	0.49
1:M:339:GLU:O	1:M:343:GLN:HB2	2.12	0.49
1:M:89:THR:OG1	4:M:602:BEF:F3	2.13	0.49
1:N:88:GLY:N	4:N:602:BEF:F3	2.34	0.49
2:O:17:VAL:HB	2:O:34:LYS:HE3	1.94	0.49

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:456:LEU:O	1:A:456:LEU:HD23	2.11	0.49
1:B:336:VAL:O	1:B:336:VAL:HG12	2.12	0.49
1:D:345:ARG:O	1:D:349:ILE:HG13	2.11	0.49
1:D:414:GLY:HA2	1:D:495:ASP:OD2	2.13	0.49
1:G:91:THR:OG1	3:G:601:ADP:O2B	2.29	0.49
1:H:149:THR:HG22	1:H:154:SER:HA	1.94	0.49
1:J:265:ASN:OD1	1:J:265:ASN:C	2.51	0.49
1:K:101:THR:O	1:K:105:LYS:HG2	2.11	0.49
1:K:232:GLU:O	1:K:235:PRO:HD2	2.12	0.49
2:Y:34:LYS:CE	2:Z:76:GLU:HA	2.34	0.49
1:A:231:ARG:HD2	1:A:261:THR:HG21	1.94	0.49
1:A:90:THR:O	1:A:94:VAL:HG23	2.12	0.49
1:B:414:GLY:HA2	1:B:495:ASP:OD2	2.12	0.49
1:C:232:GLU:C	1:C:234:LEU:H	2.16	0.49
1:E:345:ARG:O	1:E:349:ILE:HG13	2.12	0.49
1:F:228:SER:HG	1:F:229:ASN:H	1.60	0.49
1:H:5:ASP:HB2	1:H:524:LEU:HD21	1.92	0.49
1:J:350:ARG:HD3	1:J:353:ILE:HD11	1.93	0.49
1:K:205:ILE:CD1	1:K:211:GLY:HA2	2.37	0.49
1:K:415:GLY:N	3:K:601:ADP:O2'	2.32	0.49
1:K:20:VAL:HG13	1:K:74:VAL:HG11	1.93	0.49
1:N:524:LEU:CD2	1:N:525:PRO:HD2	2.42	0.49
2:P:36:THR:HG21	2:Q:76:GLU:OE2	2.12	0.49
2:Q:77:LYS:HZ1	2:Q:80:ASN:HA	1.75	0.49
2:S:64:ILE:HG23	2:S:95:VAL:CG1	2.32	0.49
1:B:197:ARG:NE	1:B:277:LYS:HD3	2.25	0.49
1:C:238:GLU:O	2:Q:25:ILE:HD11	2.12	0.49
1:C:218:PRO:HG3	1:C:323:VAL:HG13	1.94	0.49
1:D:409:GLU:OE2	1:D:501:ARG:NH2	2.28	0.49
3:E:601:ADP:O3B	4:E:602:BEF:F1	2.21	0.49
1:G:6:VAL:CG1	1:G:521:VAL:HG22	2.39	0.49
1:K:299:THR:N	1:K:316:ASP:O	2.41	0.49
1:M:268:ARG:CB	1:M:269:GLY:HA2	2.41	0.49
1:N:181:THR:OG1	1:N:182:GLY:N	2.46	0.49
2:O:4:ARG:CD	2:U:96:GLU:HG3	2.41	0.49
2:P:40:VAL:HG23	2:P:63:ASP:O	2.12	0.49
2:Q:56:PRO:O	2:Q:57:LEU:HG	2.12	0.49
2:W:8:ASP:N	2:W:8:ASP:OD2	2.42	0.49
1:B:10:ASN:O	1:B:14:VAL:HG23	2.12	0.49
1:B:227:ILE:HG22	1:B:227:ILE:O	2.13	0.49
1:C:116:LEU:O	1:C:120:ILE:HG13	2.13	0.49

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:C:276:VAL:HG11	1:C:330:THR:HG21	1.94	0.49
1:C:49:ILE:HD12	1:C:49:ILE:N	2.27	0.49
1:F:233:MET:HG3	1:F:234:LEU:N	2.27	0.49
1:I:234:LEU:HD12	1:I:234:LEU:H	1.77	0.49
1:I:420:ILE:HD11	1:I:470:LYS:HG3	1.93	0.49
1:J:171:LYS:HD3	1:J:407:VAL:CG1	2.43	0.49
1:K:415:GLY:O	1:K:451:LEU:HD12	2.12	0.49
1:L:200:LEU:HD11	1:L:277:LYS:HB2	1.94	0.49
1:L:29:VAL:HG12	1:L:29:VAL:O	2.12	0.49
1:L:34:LYS:HE3	1:L:480:ALA:O	2.13	0.49
1:N:285:ARG:O	1:N:289:LEU:N	2.45	0.49
2:O:70:GLY:O	2:O:73:VAL:HG22	2.12	0.49
2:Q:41:LEU:O	2:Q:61:VAL:HG13	2.12	0.49
2:1:3:ILE:HD12	2:1:3:ILE:O	2.12	0.49
2:1:77:LYS:HG2	2:1:82:GLU:HA	1.94	0.49
1:A:175:ILE:N	1:A:175:ILE:HD12	2.28	0.49
1:A:92:ALA:HB2	1:A:503:ALA:CB	2.42	0.49
1:C:342:ILE:HD12	1:C:342:ILE:N	2.28	0.49
1:F:305:ILE:CG2	1:F:306:GLY:H	2.14	0.49
1:G:355:GLU:HG3	1:G:356:ALA:N	2.27	0.49
1:G:417:VAL:CA	1:G:420:ILE:HG22	2.40	0.49
1:I:258:ALA:HA	1:I:261:THR:CG2	2.42	0.49
1:I:25:ASP:HA	1:I:28:LYS:HE2	1.95	0.49
1:I:338:GLU:O	1:I:342:ILE:HG13	2.13	0.49
1:J:240:VAL:HA	1:J:314:LEU:HD12	1.93	0.49
1:K:196:ASP:OD1	1:K:196:ASP:N	2.45	0.49
1:M:301:ILE:HD11	1:M:312:ALA:HB2	1.95	0.49
1:M:31:LEU:HD23	1:M:453:GLN:HB3	1.95	0.49
1:N:88:GLY:N	3:N:601:ADP:O3B	2.39	0.49
1:N:63:GLU:OE1	1:N:63:GLU:N	2.27	0.49
2:P:14:ARG:NH2	2:P:69:ASP:OD2	2.45	0.49
2:T:12:VAL:HG23	2:T:84:LEU:CD1	2.43	0.49
2:U:74:LYS:HB3	2:U:85:ILE:HD11	1.94	0.49
2:V:88:GLU:HA	2:V:91:ILE:CD1	2.43	0.49
1:A:182:GLY:O	1:A:183:LEU:HB2	2.11	0.49
1:B:220:ILE:HD12	1:B:220:ILE:N	2.27	0.49
1:D:169:VAL:HB	1:D:375:GLY:O	2.12	0.49
1:G:199:TYR:CE1	1:G:202:PRO:HA	2.47	0.49
1:G:203:TYR:HB3	1:G:267:MET:HE2	1.94	0.49
1:G:197:ARG:HG3	1:G:278:ALA:O	2.13	0.49
1:G:301:ILE:HG23	1:G:307:MET:HB3	1.94	0.49

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:H:293:ALA:HB1	1:H:298:GLY:O	2.12	0.49
1:H:174:VAL:HG22	1:H:376:VAL:HG22	1.94	0.49
1:I:326:ASN:ND2	1:I:329:THR:HB	2.27	0.49
1:L:325:ILE:HG12	1:L:330:THR:HG23	1.93	0.49
1:L:353:ILE:O	1:L:362:ARG:HD2	2.13	0.49
1:M:220:ILE:N	1:M:220:ILE:HD12	2.28	0.49
1:N:219:PHE:O	1:N:247:LEU:HD12	2.13	0.49
1:N:315:GLU:N	1:N:315:GLU:OE1	2.44	0.49
1:N:346:VAL:HG12	1:N:350:ARG:NH1	2.26	0.49
2:S:25:ILE:O	2:S:25:ILE:HG22	2.12	0.49
2:T:43:VAL:CG1	2:T:57:LEU:HD12	2.43	0.49
2:Y:78:ILE:HG13	2:Y:79:ASP:H	1.77	0.49
2:2:49:LEU:CA	2:2:52:GLY:HA2	2.43	0.49
1:F:253:ASP:HA	1:F:277:LYS:HB3	1.95	0.49
1:H:183:LEU:H	1:H:183:LEU:HD12	1.77	0.49
1:H:322:ARG:HB3	1:H:333:ILE:CD1	2.28	0.49
1:L:138:CYS:HB2	1:L:411:VAL:HG13	1.93	0.49
1:L:219:PHE:CE2	1:L:245:LYS:HB2	2.48	0.49
1:L:220:ILE:O	1:L:317:LEU:HG	2.13	0.49
1:M:320:ALA:HA	1:M:335:GLY:HA2	1.95	0.49
1:M:346:VAL:O	1:M:350:ARG:HG2	2.13	0.49
1:N:249:ILE:HD12	1:N:249:ILE:N	2.28	0.49
1:N:259:LEU:HD23	1:N:262:LEU:HD23	1.93	0.49
2:Q:59:VAL:O	2:Q:59:VAL:HG23	2.13	0.49
2:X:16:GLU:CG	2:X:19:THR:HB	2.43	0.49
2:2:84:LEU:N	2:2:84:LEU:HD12	2.28	0.49
1:B:218:PRO:HB2	1:B:248:LEU:HD23	1.94	0.49
1:C:34:LYS:HE3	1:C:480:ALA:O	2.13	0.49
1:E:496:PRO:HG2	1:E:499:VAL:CG2	2.43	0.49
1:F:220:ILE:HB	1:F:318:GLY:C	2.32	0.49
1:F:325:ILE:CG1	1:F:330:THR:HG23	2.40	0.49
1:G:223:ALA:CB	1:G:224:ASP:C	2.81	0.49
1:G:223:ALA:HB1	1:G:225:LYS:N	2.27	0.49
1:G:443:ALA:O	1:G:447:MET:HG3	2.12	0.49
1:H:147:VAL:HG12	1:H:494:LEU:HB2	1.94	0.49
1:H:285:ARG:O	1:H:289:LEU:HG	2.13	0.49
1:I:455:VAL:HG13	1:I:460:GLU:HB2	1.95	0.49
1:K:225:LYS:HG3	1:K:226:LYS:N	2.28	0.49
1:K:358:SER:HA	1:K:362:ARG:HH21	1.78	0.49
1:L:230:ILE:HD12	1:L:230:ILE:H	1.78	0.49
1:L:247:LEU:HD12	1:L:249:ILE:HD11	1.94	0.49

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:N:231:ARG:O	1:N:309:LEU:HD13	2.13	0.49
1:N:284:ARG:CZ	1:N:364:LYS:HE2	2.42	0.49
2:P:5:PRO:HD3	2:P:11:ILE:CD1	2.43	0.49
2:P:20:LYS:HA	2:P:27:LEU:HG	1.95	0.49
2:U:46:GLY:HA2	2:U:57:LEU:CD1	2.42	0.49
2:V:86:MET:CG	2:V:91:ILE:HD11	2.42	0.49
2:1:15:LYS:O	2:1:16:GLU:HB2	2.13	0.49
1:B:322:ARG:HG2	1:B:323:VAL:N	2.27	0.49
1:B:367:GLU:O	1:B:371:LYS:HG3	2.13	0.49
1:D:282:GLY:O	1:D:283:ASP:HB2	2.13	0.49
1:E:207:LYS:N	1:E:208:PRO:CD	2.67	0.49
1:F:40:LEU:HD11	1:F:56:VAL:HA	1.95	0.49
1:H:230:ILE:HB	1:H:309:LEU:CD1	2.43	0.49
1:H:349:ILE:CD1	1:H:369:VAL:HG22	2.40	0.49
1:I:41:ASP:O	1:I:42:LYS:HD2	2.13	0.49
1:L:302:SER:HB3	1:L:305:ILE:HB	1.95	0.49
1:M:269:GLY:C	1:M:271:VAL:H	2.16	0.49
1:M:20:VAL:HG13	1:M:74:VAL:HG11	1.95	0.49
1:N:337:GLY:O	1:N:338:GLU:HG2	2.12	0.49
1:N:365:LEU:HD23	1:N:365:LEU:O	2.12	0.49
1:N:80:LYS:HG3	1:N:506:TYR:CZ	2.48	0.49
2:Q:12:VAL:HG23	2:Q:39:GLU:C	2.34	0.49
2:T:11:ILE:HG13	2:T:42:ALA:H	1.77	0.49
1:D:226:LYS:N	1:D:252:GLU:HB3	2.27	0.48
1:E:227:ILE:HA	1:E:230:ILE:HD13	1.95	0.48
1:F:196:ASP:O	1:F:197:ARG:HG2	2.13	0.48
1:F:240:VAL:HG11	1:F:247:LEU:HD11	1.94	0.48
1:F:220:ILE:HB	1:F:318:GLY:CA	2.43	0.48
1:H:221:LEU:HA	1:H:317:LEU:HD22	1.95	0.48
1:H:350:ARG:O	1:H:353:ILE:HB	2.12	0.48
3:I:601:ADP:O3B	4:I:602:BEF:F3	2.20	0.48
1:K:291:ASP:HB2	1:K:372:LEU:CD2	2.41	0.48
1:L:116:LEU:O	1:L:120:ILE:HG13	2.12	0.48
1:L:219:PHE:CE1	1:L:240:VAL:HG22	2.47	0.48
1:L:398:ASP:OD2	4:L:602:BEF:F3	2.21	0.48
1:N:228:SER:CA	1:N:258:ALA:HB2	2.37	0.48
1:N:320:ALA:HB1	1:N:332:ILE:HG23	1.95	0.48
1:N:284:ARG:NE	1:N:364:LYS:HE2	2.28	0.48
2:R:6:LEU:C	2:R:6:LEU:HD23	2.33	0.48
2:2:12:VAL:HG13	2:2:39:GLU:O	2.13	0.48
1:A:164:GLU:O	1:A:168:LYS:HB2	2.13	0.48

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:452:ARG:NH2	1:A:463:SER:HA	2.28	0.48
1:C:42:LYS:HE2	1:C:48:THR:HG21	1.93	0.48
1:D:417:VAL:O	1:D:420:ILE:HG22	2.13	0.48
1:E:151:SER:CB	1:E:399:ALA:HA	2.39	0.48
1:F:174:VAL:HG11	1:F:367:GLU:HA	1.95	0.48
1:F:5:ASP:HB2	1:F:524:LEU:HD21	1.93	0.48
1:G:222:LEU:HD22	1:G:300:VAL:CG2	2.33	0.48
1:G:34:LYS:HE3	1:G:480:ALA:O	2.13	0.48
1:I:266:THR:HG22	1:I:271:VAL:O	2.13	0.48
1:I:24:ALA:O	1:I:28:LYS:HG3	2.12	0.48
1:J:166:MET:CE	1:J:171:LYS:HG2	2.43	0.48
1:K:221:LEU:HB2	1:K:317:LEU:HD22	1.95	0.48
1:K:255:GLU:OE2	1:K:255:GLU:N	2.46	0.48
1:K:355:GLU:HG3	1:K:356:ALA:N	2.28	0.48
1:L:143:ALA:O	1:L:147:VAL:HG12	2.13	0.48
2:S:16:GLU:HG3	2:S:19:THR:HG23	1.95	0.48
2:T:5:PRO:HB3	2:T:11:ILE:HG23	1.94	0.48
2:T:7:HIS:O	2:T:8:ASP:HB3	2.13	0.48
2:Z:89:SER:O	2:1:9:ARG:NH2	2.46	0.48
1:B:325:ILE:CG1	1:B:330:THR:HG23	2.34	0.48
1:B:346:VAL:O	1:B:350:ARG:HG2	2.12	0.48
1:D:180:GLY:N	1:D:381:VAL:O	2.46	0.48
1:D:448:GLU:O	1:D:452:ARG:HG3	2.13	0.48
1:E:136:VAL:CG2	1:E:137:PRO:CD	2.91	0.48
1:E:236:VAL:O	1:E:240:VAL:HG23	2.13	0.48
1:F:346:VAL:O	1:F:349:ILE:HG22	2.13	0.48
1:F:151:SER:HB3	1:F:399:ALA:HA	1.95	0.48
1:G:183:LEU:HD12	1:G:384:ALA:CB	2.43	0.48
1:G:414:GLY:HA3	1:G:493:ILE:HG22	1.96	0.48
1:H:57:ALA:O	1:H:75:LYS:HE2	2.13	0.48
1:M:419:LEU:HD21	1:M:500:THR:HG23	1.94	0.48
1:N:113:PRO:HA	1:N:116:LEU:HD12	1.95	0.48
2:2:7:HIS:O	2:2:44:GLY:HA3	2.13	0.48
1:C:232:GLU:HG3	1:C:310:GLU:OE2	2.13	0.48
1:C:305:ILE:HD12	1:C:305:ILE:N	2.28	0.48
1:D:174:VAL:HG23	1:D:376:VAL:HG13	1.96	0.48
1:F:147:VAL:CG2	1:F:403:THR:HG22	2.43	0.48
1:G:357:THR:HG22	1:G:359:ASP:HB2	1.96	0.48
1:H:226:LYS:HG2	1:H:252:GLU:HB3	1.94	0.48
1:H:85:ALA:HB1	1:H:499:VAL:HG12	1.95	0.48
1:I:138:CYS:SG	1:I:144:ILE:HD13	2.53	0.48

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:J:77:VAL:CG2	1:J:507:ALA:HA	2.43	0.48
1:K:87:ASP:OD2	1:K:151:SER:OG	2.24	0.48
1:L:234:LEU:N	1:L:235:PRO:HD2	2.28	0.48
1:N:293:ALA:O	1:N:298:GLY:N	2.46	0.48
2:Q:47:ARG:HB2	2:Q:47:ARG:NH1	2.28	0.48
2:V:47:ARG:O	2:V:54:VAL:HA	2.13	0.48
2:X:67:PHE:CA	2:X:92:LEU:HD13	2.33	0.48
2:Y:48:ILE:O	2:Y:48:ILE:HG22	2.14	0.48
1:A:218:PRO:HD2	1:A:320:ALA:O	2.13	0.48
1:A:496:PRO:HG2	1:A:499:VAL:HG11	1.95	0.48
1:E:234:LEU:CD2	1:E:235:PRO:N	2.77	0.48
1:E:197:ARG:HG3	1:E:278:ALA:O	2.13	0.48
1:F:136:VAL:HG12	1:F:137:PRO:HD2	1.96	0.48
1:F:217:SER:HA	1:F:320:ALA:O	2.13	0.48
1:F:88:GLY:N	3:F:601:ADP:O1B	2.38	0.48
1:G:223:ALA:HB3	1:G:224:ASP:C	2.33	0.48
1:G:345:ARG:O	1:G:348:GLN:HG3	2.13	0.48
1:H:227:ILE:O	1:H:227:ILE:HG22	2.14	0.48
1:I:197:ARG:NH1	1:I:277:LYS:CD	2.72	0.48
1:I:285:ARG:HG3	1:I:286:LYS:HG3	1.95	0.48
1:K:320:ALA:HB1	1:K:334:ASP:O	2.13	0.48
1:K:463:SER:O	1:K:467:ASN:HB2	2.14	0.48
1:K:478:TYR:HD1	1:K:485:TYR:CE1	2.32	0.48
1:M:131:LEU:HD23	1:M:422:VAL:HG11	1.94	0.48
1:N:147:VAL:HG22	1:N:403:THR:HG22	1.95	0.48
2:S:7:HIS:HA	2:S:46:GLY:O	2.14	0.48
1:G:237:LEU:HD22	2:U:26:VAL:CG2	2.43	0.48
2:O:5:PRO:CA	2:U:93:ALA:HB2	2.40	0.48
2:Y:92:LEU:HD23	2:Z:85:ILE:HD13	1.93	0.48
2:2:11:ILE:O	2:2:41:LEU:HB2	2.14	0.48
1:B:276:VAL:HG21	1:B:325:ILE:CG2	2.38	0.48
1:E:166:MET:HE1	1:E:407:VAL:HG21	1.93	0.48
1:E:219:PHE:HB3	1:E:317:LEU:CD1	2.43	0.48
1:G:240:VAL:HG11	1:G:247:LEU:CD1	2.43	0.48
1:G:333:ILE:HG22	1:G:333:ILE:O	2.14	0.48
1:I:228:SER:H	1:I:230:ILE:CD1	2.27	0.48
1:I:320:ALA:HA	1:I:335:GLY:HA2	1.94	0.48
1:J:230:ILE:HG13	1:J:231:ARG:N	2.22	0.48
1:K:231:ARG:O	1:K:234:LEU:HB3	2.13	0.48
1:K:414:GLY:O	1:K:417:VAL:HG12	2.14	0.48
1:L:396:VAL:O	1:L:400:LEU:HB2	2.13	0.48

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:M:405:ALA:HB2	1:M:498:LYS:HD3	1.94	0.48
1:N:150:ILE:HG22	1:N:151:SER:N	2.28	0.48
1:N:222:LEU:CD2	1:N:293:ALA:HB2	2.42	0.48
2:O:96:GLU:O	2:P:1:MET:N	2.44	0.48
2:P:1:MET:HE1	2:P:78:ILE:C	2.30	0.48
1:A:200:LEU:HD21	1:A:253:ASP:OD1	2.13	0.48
1:A:228:SER:O	1:A:258:ALA:HB2	2.14	0.48
1:A:307:MET:C	1:A:308:GLU:HG3	2.34	0.48
1:B:199:TYR:HE1	1:B:202:PRO:HA	1.74	0.48
1:E:202:PRO:O	1:E:205:ILE:HG13	2.14	0.48
1:F:231:ARG:HB2	1:F:234:LEU:HD12	1.94	0.48
1:F:448:GLU:O	1:F:452:ARG:HG2	2.14	0.48
1:F:475:ASN:O	1:F:488:MET:HG2	2.12	0.48
1:G:215:LEU:HD22	1:G:246:PRO:HB3	1.95	0.48
1:L:218:PRO:HB2	1:L:219:PHE:O	2.13	0.48
1:L:85:ALA:O	1:L:401:HIS:HB3	2.14	0.48
1:N:339:GLU:OE1	1:N:339:GLU:N	2.47	0.48
2:V:37:ARG:HG2	2:V:66:ILE:HG12	1.95	0.48
2:V:40:VAL:HG11	2:V:63:ASP:OD1	2.13	0.48
2:Z:23:GLY:O	2:Z:26:VAL:HG12	2.13	0.48
1:A:196:ASP:O	1:A:197:ARG:HG2	2.14	0.48
1:B:337:GLY:O	1:B:338:GLU:HG2	2.13	0.48
1:C:259:LEU:O	1:C:263:VAL:N	2.42	0.48
1:C:357:THR:HB	1:C:361:ASP:CB	2.44	0.48
1:D:228:SER:CA	1:D:255:GLU:HB2	2.34	0.48
1:H:124:VAL:HG22	1:H:504:LEU:HD11	1.95	0.48
1:H:80:LYS:HG2	1:I:385:THR:HG22	1.95	0.48
1:I:199:TYR:HB3	1:I:325:ILE:HG21	1.95	0.48
1:I:193:MET:HG3	1:I:371:LYS:HB3	1.96	0.48
1:I:4:LYS:HG3	1:J:59:GLU:O	2.13	0.48
1:A:197:ARG:HG3	1:A:278:ALA:C	2.34	0.48
1:B:305:ILE:HD12	1:C:263:VAL:CG1	2.41	0.48
1:C:215:LEU:HB2	1:C:323:VAL:CG2	2.43	0.48
1:C:364:LYS:O	1:C:367:GLU:HB3	2.14	0.48
1:C:417:VAL:HA	1:C:420:ILE:HG22	1.96	0.48
1:E:240:VAL:HG12	1:E:247:LEU:HD21	1.92	0.48
1:F:165:ALA:O	1:F:169:VAL:HG22	2.13	0.48
1:G:195:PHE:O	1:G:329:THR:HG23	2.12	0.48
1:I:513:LEU:O	1:I:516:THR:OG1	2.23	0.48
1:J:10:ASN:O	1:J:14:VAL:HG23	2.13	0.48
1:J:214:GLU:HA	1:J:323:VAL:O	2.13	0.48

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:K:345:ARG:O	1:K:348:GLN:HG2	2.14	0.48
1:K:419:LEU:O	1:K:422:VAL:HG22	2.14	0.48
1:L:136:VAL:HG22	1:L:137:PRO:HD2	1.95	0.48
1:M:160:LYS:O	1:M:164:GLU:HG3	2.14	0.48
1:N:175:ILE:HA	1:N:377:ALA:O	2.14	0.48
2:O:78:ILE:HD12	2:O:79:ASP:OD1	2.14	0.48
2:Q:56:PRO:C	2:Q:57:LEU:HG	2.33	0.48
2:V:15:LYS:HA	2:V:16:GLU:HA	1.59	0.48
2:V:80:ASN:ND2	2:V:80:ASN:O	2.47	0.48
2:X:15:LYS:HB2	2:X:37:ARG:O	2.14	0.48
2:1:7:HIS:O	2:1:8:ASP:HB3	2.14	0.48
1:A:234:LEU:HB2	2:O:26:VAL:HG21	1.95	0.48
1:A:234:LEU:N	1:A:235:PRO:HD2	2.28	0.48
1:A:345:ARG:O	1:A:348:GLN:HG3	2.13	0.48
1:A:367:GLU:HG3	1:A:371:LYS:HZ3	1.79	0.48
1:A:89:THR:N	4:A:602:BEF:F1	2.37	0.48
1:C:228:SER:HB3	1:C:255:GLU:OE1	2.14	0.48
1:D:291:ASP:OD1	1:D:345:ARG:NH2	2.36	0.48
1:D:62:LEU:HD23	1:D:67:GLU:HB3	1.95	0.48
1:E:66:PHE:CE1	1:E:522:THR:HG22	2.49	0.48
1:H:230:ILE:CD1	1:H:309:LEU:HB2	2.44	0.48
1:H:48:THR:HG22	1:H:390:LYS:HZ1	1.78	0.48
1:H:89:THR:OG1	4:H:602:BEF:F2	2.22	0.48
1:I:217:SER:O	1:I:245:LYS:HD3	2.14	0.48
1:I:199:TYR:CE1	1:I:327:LYS:HA	2.49	0.48
1:N:304:GLU:O	1:N:305:ILE:HD13	2.13	0.48
2:W:12:VAL:HA	2:W:41:LEU:HD13	1.96	0.48
1:A:224:ASP:OD2	1:A:302:SER:HB3	2.14	0.47
1:A:348:GLN:OE1	1:A:352:GLN:NE2	2.47	0.47
1:C:114:MET:HG3	1:C:118:ARG:NH1	2.29	0.47
1:E:433:ASN:OD1	1:E:436:GLN:HG3	2.13	0.47
1:E:417:VAL:HG21	1:E:476:TYR:O	2.13	0.47
1:F:227:ILE:O	1:F:254:VAL:HA	2.13	0.47
1:G:252:GLU:HA	1:G:285:ARG:NH1	2.29	0.47
1:G:284:ARG:HG2	1:G:288:MET:CE	2.43	0.47
1:K:280:GLY:C	1:K:285:ARG:HB3	2.33	0.47
1:K:429:LEU:HG	1:K:440:ILE:HD13	1.95	0.47
1:L:196:ASP:O	1:L:197:ARG:HG2	2.14	0.47
1:N:288:MET:O	1:N:292:ILE:HG13	2.14	0.47
1:N:199:TYR:CE2	1:N:326:ASN:HA	2.49	0.47
1:N:440:ILE:O	1:N:444:LEU:HG	2.14	0.47

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:O:96:GLU:O	2:O:97:ALA:CB	2.61	0.47
1:B:69:MET:SD	1:C:41:ASP:HB2	2.54	0.47
1:D:220:ILE:HD12	1:D:220:ILE:N	2.29	0.47
1:D:238:GLU:OE2	2:R:23:GLY:N	2.34	0.47
1:E:230:ILE:HG22	1:E:230:ILE:O	2.14	0.47
1:E:234:LEU:HG	1:E:235:PRO:N	2.28	0.47
1:E:302:SER:HB2	1:E:305:ILE:CB	2.44	0.47
1:E:175:ILE:HD12	1:E:400:LEU:HD21	1.96	0.47
1:F:263:VAL:O	1:F:267:MET:HG3	2.14	0.47
1:I:222:LEU:HD23	1:I:250:ILE:HB	1.96	0.47
1:I:278:ALA:HB1	1:I:279:PRO:CD	2.43	0.47
1:I:325:ILE:N	1:I:325:ILE:HD12	2.29	0.47
1:I:326:ASN:OD1	1:I:329:THR:HB	2.13	0.47
1:L:233:MET:C	1:L:235:PRO:HD2	2.35	0.47
1:L:433:ASN:ND2	1:L:435:ASP:HB2	2.29	0.47
1:M:305:ILE:CG2	1:M:306:GLY:N	2.72	0.47
1:M:126:ALA:HB1	1:M:426:LEU:HD22	1.96	0.47
1:N:307:MET:HG2	1:N:311:LYS:NZ	2.27	0.47
2:P:52:GLY:HA3	2:P:53:GLU:HA	1.60	0.47
2:2:17:VAL:HG23	2:2:18:GLU:CG	2.43	0.47
1:A:152:ALA:HB2	1:A:399:ALA:HB2	1.96	0.47
1:B:169:VAL:HG13	1:B:173:GLY:HA3	1.95	0.47
1:B:339:GLU:OE1	1:B:339:GLU:N	2.48	0.47
1:C:223:ALA:HB1	1:C:227:ILE:HD11	1.94	0.47
1:C:219:PHE:CE2	1:C:245:LYS:HB2	2.43	0.47
1:D:226:LYS:HB3	1:D:252:GLU:CG	2.43	0.47
1:H:311:LYS:HD3	1:H:311:LYS:O	2.14	0.47
1:H:102:GLU:HB2	1:H:442:VAL:HG13	1.96	0.47
1:H:49:ILE:N	1:H:49:ILE:HD12	2.29	0.47
1:J:268:ARG:HB2	1:J:270:ILE:HG12	1.96	0.47
1:L:278:ALA:HB1	1:L:285:ARG:HB2	1.96	0.47
1:N:177:VAL:HG21	1:N:396:VAL:HG11	1.96	0.47
2:P:45:ASN:OD1	2:P:46:GLY:N	2.47	0.47
2:V:6:LEU:O	2:V:6:LEU:HD12	2.14	0.47
2:W:49:LEU:HD12	2:W:52:GLY:HA3	1.96	0.47
2:Z:11:ILE:CG1	2:Z:83:VAL:HG11	2.44	0.47
2:2:12:VAL:HG13	2:2:39:GLU:N	2.29	0.47
1:D:169:VAL:CG2	1:D:173:GLY:HA3	2.44	0.47
1:D:147:VAL:CG2	1:D:403:THR:HG22	2.44	0.47
1:D:131:LEU:CD2	1:D:422:VAL:HG11	2.43	0.47
1:D:39:VAL:HG22	1:D:49:ILE:HG12	1.95	0.47

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:E:224:ASP:OD2	1:E:286:LYS:HE2	2.14	0.47
1:E:217:SER:HA	1:E:320:ALA:O	2.15	0.47
1:F:257:GLU:O	1:F:261:THR:HG22	2.14	0.47
1:G:225:LYS:HB2	1:G:303:GLU:CD	2.35	0.47
1:H:417:VAL:O	1:H:420:ILE:HG22	2.14	0.47
1:I:197:ARG:HA	1:I:197:ARG:HD3	1.73	0.47
1:I:255:GLU:HA	1:I:255:GLU:OE1	2.14	0.47
1:I:311:LYS:HB3	1:I:311:LYS:HZ3	1.79	0.47
1:K:305:ILE:CG2	1:K:306:GLY:N	2.72	0.47
1:L:365:LEU:O	1:L:369:VAL:HG23	2.14	0.47
1:M:123:ALA:HB2	1:M:440:ILE:HG23	1.96	0.47
1:N:169:VAL:HB	1:N:375:GLY:O	2.14	0.47
1:N:197:ARG:CD	1:N:277:LYS:HB3	2.45	0.47
1:N:242:LYS:HD3	1:N:242:LYS:C	2.35	0.47
1:N:276:VAL:CG1	1:N:325:ILE:HG21	2.45	0.47
1:N:325:ILE:HD12	1:N:325:ILE:N	2.28	0.47
1:N:392:LYS:O	1:N:396:VAL:HG23	2.13	0.47
1:N:177:VAL:CG2	1:N:393:LYS:HG3	2.44	0.47
2:T:60:LYS:HB3	2:T:63:ASP:OD1	2.13	0.47
2:Z:27:LEU:HD23	2:Z:28:THR:N	2.29	0.47
1:A:322:ARG:HG2	1:A:323:VAL:N	2.29	0.47
1:B:247:LEU:HD12	1:B:247:LEU:C	2.35	0.47
1:D:496:PRO:HG2	1:D:499:VAL:HG13	1.95	0.47
1:E:343:GLN:HA	1:E:346:VAL:HB	1.96	0.47
1:F:10:ASN:O	1:F:14:VAL:HG23	2.14	0.47
1:F:177:VAL:CG2	1:F:393:LYS:HG3	2.45	0.47
1:F:452:ARG:HH21	1:F:466:ALA:CB	2.27	0.47
1:F:124:VAL:HG21	1:F:508:ALA:CB	2.45	0.47
1:G:257:GLU:O	1:G:261:THR:HG22	2.13	0.47
1:H:280:GLY:C	1:H:285:ARG:HB2	2.35	0.47
1:H:364:LYS:HA	1:H:367:GLU:HB3	1.96	0.47
1:I:124:VAL:HG21	1:I:508:ALA:HB2	1.94	0.47
1:I:510:VAL:O	1:I:514:MET:HB2	2.14	0.47
1:M:288:MET:HG3	1:M:364:LYS:NZ	2.29	0.47
1:M:381:VAL:HG22	1:M:396:VAL:HG21	1.97	0.47
1:N:286:LYS:NZ	1:N:304:GLU:OE1	2.34	0.47
1:N:478:TYR:N	1:N:488:MET:HE1	2.30	0.47
1:N:493:ILE:C	1:N:494:LEU:HD12	2.35	0.47
2:R:60:LYS:HG3	2:R:63:ASP:OD1	2.14	0.47
2:V:37:ARG:HH21	2:W:76:GLU:HG3	1.79	0.47
2:X:50:GLU:O	2:X:51:ASN:CB	2.62	0.47

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:C:198:GLY:O	1:C:276:VAL:HG23	2.14	0.47
1:C:345:ARG:NH2	1:C:372:LEU:HD22	2.30	0.47
1:D:280:GLY:O	1:D:285:ARG:NH1	2.48	0.47
1:D:349:ILE:HD12	1:D:369:VAL:HG22	1.97	0.47
1:D:20:VAL:HG13	1:D:74:VAL:HG11	1.97	0.47
1:F:209:GLU:CG	1:F:210:THR:H	2.26	0.47
1:F:360:TYR:CA	1:F:363:GLU:HG3	2.41	0.47
1:F:455:VAL:CG1	1:F:460:GLU:HB2	2.45	0.47
1:H:236:VAL:O	1:H:240:VAL:HG23	2.14	0.47
1:I:355:GLU:HG3	1:I:356:ALA:N	2.29	0.47
1:J:219:PHE:HB3	1:J:317:LEU:HD13	1.95	0.47
1:K:326:ASN:HD21	1:K:329:THR:HG22	1.80	0.47
1:M:88:GLY:HA2	3:M:601:ADP:O3B	2.13	0.47
1:N:14:VAL:O	1:N:18:ARG:HG3	2.15	0.47
2:O:31:ALA:O	2:O:32:ALA:HB3	2.14	0.47
2:T:60:LYS:CG	2:T:61:VAL:H	2.28	0.47
1:A:188:ASP:HB2	1:A:378:VAL:CG2	2.41	0.47
1:A:313:THR:O	1:A:316:ASP:HB2	2.15	0.47
1:A:333:ILE:O	1:A:334:ASP:HB2	2.14	0.47
1:B:519:CYS:HB3	1:C:38:VAL:HG22	1.96	0.47
1:G:219:PHE:HB3	1:G:317:LEU:HB3	1.97	0.47
1:G:412:VAL:CG1	1:G:413:ALA:H	2.27	0.47
1:L:76:GLU:O	1:L:80:LYS:HB2	2.15	0.47
1:N:227:ILE:CG1	1:N:231:ARG:HG3	2.44	0.47
2:O:57:LEU:HD12	2:O:88:GLU:CB	2.44	0.47
2:V:50:GLU:HG2	2:2:50:GLU:OE1	2.14	0.47
2:W:71:TYR:C	2:W:71:TYR:CD1	2.88	0.47
1:A:270:ILE:HG22	1:A:271:VAL:HG13	1.97	0.47
1:A:197:ARG:HG3	1:A:278:ALA:O	2.15	0.47
1:B:310:GLU:N	1:B:310:GLU:OE1	2.47	0.47
1:B:417:VAL:HA	1:B:420:ILE:HG22	1.96	0.47
1:B:27:VAL:CG1	1:B:90:THR:HG23	2.44	0.47
1:C:48:THR:HG22	1:C:390:LYS:HZ2	1.78	0.47
1:D:165:ALA:O	1:D:169:VAL:HG22	2.14	0.47
1:E:194:GLN:CG	1:E:331:THR:HG23	2.41	0.47
1:E:350:ARG:HD3	1:E:353:ILE:CD1	2.45	0.47
1:F:169:VAL:CG2	1:F:173:GLY:HA3	2.45	0.47
1:F:496:PRO:HG2	1:F:499:VAL:HG13	1.97	0.47
1:G:196:ASP:O	1:G:197:ARG:HG2	2.15	0.47
1:I:346:VAL:O	1:I:350:ARG:HG2	2.14	0.47
1:J:399:ALA:O	1:J:403:THR:HG23	2.15	0.47

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:J:455:VAL:HG13	1:J:460:GLU:HB2	1.96	0.47
1:K:293:ALA:HB1	1:K:298:GLY:O	2.15	0.47
1:K:327:LYS:H	1:K:327:LYS:HD3	1.80	0.47
1:M:205:ILE:HG22	1:M:206:ASN:O	2.15	0.47
1:N:76:GLU:O	1:N:80:LYS:N	2.45	0.47
2:P:1:MET:HE3	2:P:78:ILE:HA	1.34	0.47
2:Q:40:VAL:CG1	2:Q:62:GLY:H	2.27	0.47
2:S:18:GLU:HG3	2:S:32:ALA:HB2	1.97	0.47
2:S:48:ILE:HG13	2:S:53:GLU:H	1.79	0.47
2:T:10:VAL:N	2:T:86:MET:O	2.40	0.47
2:U:6:LEU:HD12	2:U:7:HIS:HB2	1.97	0.47
2:Y:13:LYS:HD2	2:Y:81:GLU:OE1	2.15	0.47
1:A:62:LEU:HD23	1:A:67:GLU:HB3	1.96	0.47
1:B:221:LEU:HD23	1:B:221:LEU:C	2.34	0.47
1:D:200:LEU:HD11	1:D:277:LYS:N	2.30	0.47
1:D:327:LYS:H	1:D:327:LYS:HD3	1.80	0.47
1:F:220:ILE:HG22	1:F:222:LEU:HD11	1.97	0.47
1:G:234:LEU:HD23	2:U:23:GLY:HA3	1.96	0.47
1:J:364:LYS:NZ	1:J:367:GLU:OE2	2.42	0.47
1:K:100:ILE:O	1:K:104:LEU:HG	2.14	0.47
1:K:421:ARG:HA	1:K:421:ARG:HD3	1.77	0.47
1:L:458:CYS:SG	1:L:480:ALA:HB1	2.55	0.47
1:M:221:LEU:HD11	1:M:301:ILE:HD12	1.96	0.47
1:M:280:GLY:H	1:M:285:ARG:HB3	1.80	0.47
1:N:13:ARG:HG2	1:N:104:LEU:HD13	1.97	0.47
1:N:278:ALA:HB1	1:N:279:PRO:HD2	1.96	0.47
1:N:431:GLY:N	1:N:437:ASN:OD1	2.32	0.47
1:C:261:THR:HB	2:Q:29:GLY:O	2.14	0.47
2:R:12:VAL:HG12	2:R:40:VAL:CG1	2.42	0.47
2:R:51:ASN:C	2:R:53:GLU:H	2.17	0.47
2:Z:15:LYS:NZ	2:Z:64:ILE:HD12	2.29	0.47
2:Z:17:VAL:HG22	2:Z:18:GLU:N	2.30	0.47
2:Z:84:LEU:HD12	2:Z:84:LEU:N	2.29	0.47
1:B:493:ILE:C	1:B:494:LEU:HD12	2.35	0.47
1:C:175:ILE:HA	1:C:377:ALA:O	2.15	0.47
1:C:404:ARG:O	1:C:408:GLU:HG2	2.15	0.47
1:C:406:ALA:HB1	1:C:411:VAL:CG1	2.45	0.47
1:H:245:LYS:CB	1:H:246:PRO:HD2	2.33	0.47
1:I:124:VAL:HG22	1:I:504:LEU:CD1	2.44	0.47
1:I:5:ASP:N	1:I:522:THR:O	2.46	0.47
1:L:331:THR:HG22	1:L:332:ILE:N	2.30	0.47

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:N:414:GLY:O	1:N:417:VAL:HG12	2.15	0.47
1:N:429:LEU:HG	1:N:440:ILE:HD13	1.97	0.47
2:O:77:LYS:HE2	2:O:82:GLU:HB2	1.97	0.47
2:U:20:LYS:HB3	2:U:27:LEU:HD12	1.96	0.47
2:Y:73:VAL:HG22	2:Y:86:MET:HB3	1.96	0.47
2:Z:92:LEU:HD13	2:1:85:ILE:HG21	1.96	0.47
1:A:85:ALA:O	1:A:401:HIS:HB3	2.14	0.47
1:B:13:ARG:CD	1:B:104:LEU:HD22	2.42	0.47
1:F:220:ILE:HG13	1:F:296:THR:CB	2.44	0.47
1:F:231:ARG:O	1:F:234:LEU:HG	2.15	0.47
1:F:282:GLY:CA	1:F:285:ARG:HG2	2.42	0.47
1:F:291:ASP:OD1	1:F:292:ILE:HG13	2.15	0.47
1:F:419:LEU:N	1:F:419:LEU:HD12	2.30	0.47
1:G:436:GLN:O	1:G:440:ILE:HG13	2.14	0.47
1:I:77:VAL:HB	1:I:510:VAL:HG21	1.96	0.47
1:L:115:ASP:O	1:L:436:GLN:HG2	2.14	0.47
1:L:450:PRO:O	1:L:454:ILE:HG13	2.15	0.47
1:N:204:PHE:CE2	1:N:274:ALA:HA	2.50	0.47
2:V:49:LEU:HD22	2:V:50:GLU:N	2.28	0.47
2:X:27:LEU:C	2:X:27:LEU:HD12	2.35	0.47
1:A:143:ALA:O	1:A:147:VAL:HG13	2.14	0.46
1:A:266:THR:HG22	1:A:273:VAL:H	1.80	0.46
1:B:213:VAL:O	1:B:324:VAL:HA	2.15	0.46
1:F:301:ILE:HD11	1:F:316:ASP:OD2	2.15	0.46
1:G:306:GLY:O	1:G:307:MET:CG	2.60	0.46
1:G:367:GLU:O	1:G:371:LYS:HG3	2.15	0.46
1:H:429:LEU:HG	1:H:440:ILE:HD13	1.96	0.46
1:I:217:SER:N	1:I:218:PRO:CD	2.78	0.46
1:I:237:LEU:CD2	2:W:27:LEU:CD2	2.93	0.46
1:I:487:ASN:HB3	1:I:490:ASP:HB2	1.96	0.46
1:L:195:PHE:O	1:L:329:THR:OG1	2.29	0.46
1:M:207:LYS:N	1:M:208:PRO:HD2	2.30	0.46
1:M:219:PHE:HB3	1:M:317:LEU:HD13	1.97	0.46
1:N:478:TYR:HD1	1:N:485:TYR:CE1	2.32	0.46
2:O:80:ASN:ND2	2:U:24:GLY:H	2.14	0.46
1:H:234:LEU:HD21	2:V:22:ALA:HB1	1.96	0.46
2:V:36:THR:HG21	2:W:76:GLU:OE1	2.15	0.46
1:B:180:GLY:HA3	1:B:382:GLY:CA	2.45	0.46
1:E:200:LEU:HD11	1:E:277:LYS:CB	2.42	0.46
1:E:204:PHE:CD2	1:E:274:ALA:HA	2.51	0.46
1:H:278:ALA:HB1	1:H:279:PRO:HD2	1.97	0.46

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:H:455:VAL:HG21	1:H:465:VAL:HG11	1.97	0.46
1:I:18:ARG:HH11	1:I:18:ARG:HB3	1.80	0.46
1:J:226:LYS:HB2	1:J:252:GLU:CD	2.35	0.46
1:J:215:LEU:HB2	1:J:323:VAL:CG2	2.44	0.46
1:J:510:VAL:O	1:J:514:MET:HB2	2.15	0.46
1:K:359:ASP:C	1:K:361:ASP:H	2.19	0.46
1:M:230:ILE:H	1:M:230:ILE:HD12	1.80	0.46
1:M:293:ALA:HB1	1:M:298:GLY:O	2.15	0.46
1:N:222:LEU:HD22	1:N:300:VAL:CG2	2.44	0.46
1:N:226:LYS:HE3	1:N:253:ASP:HB3	1.98	0.46
1:N:221:LEU:HD13	1:N:317:LEU:HD21	1.97	0.46
1:N:455:VAL:HG11	1:N:462:PRO:HA	1.97	0.46
2:S:60:LYS:HB2	2:S:63:ASP:OD2	2.15	0.46
2:Z:23:GLY:H	2:1:80:ASN:HB3	1.80	0.46
1:B:218:PRO:HG3	1:B:323:VAL:HG22	1.96	0.46
1:C:357:THR:O	1:C:357:THR:HG22	2.16	0.46
1:H:232:GLU:CG	1:H:233:MET:N	2.76	0.46
1:H:34:LYS:HE3	1:H:480:ALA:O	2.15	0.46
1:I:347:ALA:HA	1:I:350:ARG:HG3	1.97	0.46
1:I:180:GLY:HA2	1:I:380:LYS:HB3	1.97	0.46
1:J:291:ASP:HB3	1:J:372:LEU:HD21	1.98	0.46
1:J:198:GLY:HA3	1:J:327:LYS:O	2.16	0.46
1:J:34:LYS:HD2	1:J:458:CYS:SG	2.55	0.46
1:J:23:LEU:CD2	1:J:60:ILE:HG13	2.46	0.46
1:K:248:LEU:HD13	1:K:249:ILE:N	2.30	0.46
1:K:513:LEU:CD2	1:L:49:ILE:HG21	2.45	0.46
1:L:193:MET:SD	1:L:371:LYS:HB3	2.56	0.46
1:L:285:ARG:HG3	1:L:286:LYS:N	2.30	0.46
1:M:2:ALA:O	1:N:61:GLU:HB3	2.15	0.46
1:M:151:SER:OG	1:M:399:ALA:HA	2.14	0.46
1:N:96:ALA:O	1:N:100:ILE:HG13	2.15	0.46
1:N:161:LEU:HD12	1:N:161:LEU:HA	1.84	0.46
1:N:66:PHE:CE1	1:N:522:THR:HG22	2.51	0.46
2:R:17:VAL:HG23	2:R:19:THR:HA	1.97	0.46
2:S:64:ILE:CG2	2:S:95:VAL:HG13	2.32	0.46
2:U:83:VAL:C	2:U:84:LEU:HD12	2.35	0.46
1:B:130:GLU:O	1:B:134:LEU:HD13	2.15	0.46
1:B:226:LYS:HE3	1:B:253:ASP:HB3	1.96	0.46
1:B:419:LEU:O	1:B:422:VAL:HG22	2.16	0.46
1:D:263:VAL:O	1:D:267:MET:HG3	2.16	0.46
1:E:177:VAL:HG23	1:E:381:VAL:CG2	2.46	0.46

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:F:327:LYS:HD3	1:F:327:LYS:H	1.80	0.46
1:G:143:ALA:O	1:G:147:VAL:HG13	2.16	0.46
1:G:210:THR:HG22	1:G:210:THR:O	2.14	0.46
1:H:419:LEU:O	1:H:422:VAL:HG22	2.16	0.46
1:I:278:ALA:HB1	1:I:279:PRO:HD2	1.96	0.46
1:J:350:ARG:HA	1:J:353:ILE:HD12	1.96	0.46
1:K:288:MET:O	1:K:292:ILE:HG13	2.15	0.46
1:L:218:PRO:O	1:L:319:GLN:HG2	2.14	0.46
1:M:349:ILE:HG21	1:M:369:VAL:HG22	1.97	0.46
1:N:150:ILE:HD11	1:N:493:ILE:CG2	2.31	0.46
1:N:273:VAL:HG12	1:N:274:ALA:N	2.31	0.46
1:N:61:GLU:C	1:N:62:LEU:HD12	2.35	0.46
2:P:59:VAL:HG11	2:P:91:ILE:HD12	1.97	0.46
2:W:6:LEU:C	2:W:6:LEU:HD23	2.36	0.46
2:Y:78:ILE:HG13	2:Y:79:ASP:OD1	2.15	0.46
2:2:40:VAL:HG23	2:2:63:ASP:O	2.16	0.46
1:A:193:MET:SD	1:A:371:LYS:HB3	2.56	0.46
1:A:327:LYS:HD3	1:A:327:LYS:H	1.81	0.46
1:C:23:LEU:CD2	1:C:60:ILE:HB	2.43	0.46
1:D:57:ALA:O	1:D:75:LYS:HE2	2.16	0.46
1:E:234:LEU:N	1:E:235:PRO:HD2	2.31	0.46
1:F:465:VAL:O	1:F:469:VAL:HG23	2.15	0.46
1:H:77:VAL:HG21	1:H:507:ALA:CA	2.36	0.46
1:H:95:LEU:O	1:H:99:ILE:HG13	2.15	0.46
1:I:215:LEU:HB2	1:I:323:VAL:CG2	2.46	0.46
1:I:233:MET:CB	1:I:309:LEU:HD21	2.45	0.46
1:L:234:LEU:HD23	2:Z:23:GLY:CA	2.44	0.46
1:L:333:ILE:O	1:L:334:ASP:HB2	2.15	0.46
1:M:232:GLU:OE1	1:M:232:GLU:N	2.39	0.46
1:M:147:VAL:HG12	1:M:494:LEU:CD1	2.46	0.46
1:N:296:THR:HG22	1:N:320:ALA:HB2	1.98	0.46
2:R:25:ILE:HA	2:R:26:VAL:HG22	1.96	0.46
2:X:20:LYS:HG3	2:X:21:SER:H	1.80	0.46
2:X:6:LEU:O	2:X:6:LEU:HD12	2.15	0.46
2:Y:11:ILE:CG2	2:Y:85:ILE:HB	2.39	0.46
1:B:169:VAL:O	1:B:169:VAL:HG13	2.15	0.46
1:E:223:ALA:HA	1:E:301:ILE:O	2.15	0.46
1:E:415:GLY:HA3	1:E:488:MET:HE2	1.98	0.46
1:F:215:LEU:HB2	1:F:323:VAL:HG22	1.97	0.46
1:G:116:LEU:O	1:G:120:ILE:HG13	2.16	0.46
1:H:185:ASP:OD1	1:H:185:ASP:N	2.49	0.46

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:H:234:LEU:H	1:H:234:LEU:CD1	2.28	0.46
1:H:256:GLY:O	1:H:260:ALA:N	2.40	0.46
1:H:278:ALA:HB1	1:H:279:PRO:CD	2.46	0.46
1:K:179:ASP:OD1	1:K:393:LYS:HD3	2.14	0.46
2:O:17:VAL:HB	2:O:34:LYS:CE	2.46	0.46
2:Q:40:VAL:HG11	2:Q:60:LYS:O	2.16	0.46
2:S:97:ALA:HA	2:T:1:MET:HG3	1.97	0.46
2:U:17:VAL:HG22	2:U:18:GLU:N	2.31	0.46
2:U:57:LEU:HD12	2:U:57:LEU:N	2.29	0.46
2:Y:24:GLY:HA2	2:Y:25:ILE:HA	1.43	0.46
1:A:197:ARG:HG2	1:A:279:PRO:HA	1.97	0.46
1:G:219:PHE:HB2	1:G:247:LEU:HD12	1.98	0.46
1:G:252:GLU:HA	1:G:285:ARG:HH11	1.81	0.46
1:G:228:SER:HB3	1:G:255:GLU:CD	2.35	0.46
1:I:513:LEU:HD21	1:J:391:GLU:HG2	1.98	0.46
1:J:465:VAL:O	1:J:469:VAL:HG23	2.16	0.46
1:M:233:MET:HB3	1:M:236:VAL:HG21	1.97	0.46
1:M:91:THR:HG23	1:M:450:PRO:HB3	1.98	0.46
1:N:301:ILE:HG23	1:N:307:MET:HE2	1.98	0.46
2:O:4:ARG:HD3	2:U:96:GLU:HG3	1.97	0.46
2:Q:9:ARG:HA	2:Q:87:SER:HA	1.98	0.46
2:U:83:VAL:HG12	2:U:84:LEU:H	1.80	0.46
1:A:415:GLY:N	3:A:601:ADP:O2'	2.44	0.46
1:B:4:LYS:HG3	1:C:59:GLU:O	2.16	0.46
1:C:220:ILE:HD12	1:C:220:ILE:N	2.31	0.46
1:A:59:GLU:O	1:G:4:LYS:HG3	2.16	0.46
1:I:259:LEU:O	1:I:263:VAL:HG23	2.16	0.46
1:I:381:VAL:CG1	1:I:392:LYS:HB3	2.45	0.46
1:L:200:LEU:CD2	1:L:254:VAL:HB	2.46	0.46
1:M:200:LEU:HD11	1:M:277:LYS:HG3	1.98	0.46
1:M:314:LEU:HD12	1:M:315:GLU:N	2.31	0.46
1:M:91:THR:CG2	1:M:450:PRO:HB3	2.46	0.46
1:N:10:ASN:HA	1:N:13:ARG:HB3	1.96	0.46
1:N:231:ARG:HA	1:N:231:ARG:HD3	1.74	0.46
2:O:15:LYS:O	2:O:16:GLU:HB2	2.15	0.46
2:O:14:ARG:HE	2:O:84:LEU:HD11	1.81	0.46
2:R:12:VAL:CG1	2:R:40:VAL:HA	2.40	0.46
2:T:40:VAL:HG13	2:T:62:GLY:H	1.80	0.46
2:U:14:ARG:HG2	2:U:35:SER:OG	2.15	0.46
2:V:94:ILE:HG13	2:V:94:ILE:O	2.16	0.46
2:2:17:VAL:HG11	2:2:34:LYS:HD3	1.97	0.46

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:38:VAL:HB	1:A:56:VAL:HG21	1.98	0.46
1:C:219:PHE:HB3	1:C:317:LEU:HD13	1.97	0.46
1:E:164:GLU:O	1:E:168:LYS:HG2	2.16	0.46
1:H:197:ARG:CD	1:H:277:LYS:HB3	2.45	0.46
1:I:241:ALA:HB2	1:I:271:VAL:HG11	1.98	0.46
1:J:230:ILE:C	1:J:231:ARG:HG2	2.36	0.46
1:J:204:PHE:CE1	1:J:263:VAL:HA	2.44	0.46
1:J:346:VAL:O	1:J:350:ARG:HG2	2.16	0.46
1:J:27:VAL:CG1	1:J:90:THR:HG23	2.40	0.46
1:K:288:MET:HG3	1:K:368:ARG:NE	2.31	0.46
1:L:263:VAL:O	1:L:267:MET:HG2	2.16	0.46
1:L:249:ILE:HD12	1:L:274:ALA:O	2.15	0.46
1:L:295:LEU:C	1:L:295:LEU:HD23	2.36	0.46
1:N:358:SER:HA	1:N:362:ARG:NH2	2.31	0.46
1:N:74:VAL:HG13	1:N:514:MET:SD	2.56	0.46
2:O:16:GLU:HB3	2:O:17:VAL:H	1.61	0.46
2:R:12:VAL:HB	2:R:39:GLU:O	2.16	0.46
2:S:57:LEU:HD12	2:S:57:LEU:N	2.31	0.46
2:U:38:GLY:O	2:U:65:VAL:N	2.28	0.46
2:U:60:LYS:HG2	2:U:61:VAL:H	1.80	0.46
2:V:70:GLY:O	2:V:73:VAL:HG12	2.16	0.46
2:X:8:ASP:O	2:X:57:LEU:HD21	2.16	0.46
2:1:95:VAL:CA	2:2:3:ILE:HG22	2.46	0.46
1:A:199:TYR:CB	1:A:276:VAL:HG12	2.46	0.46
1:B:225:LYS:HB2	1:B:303:GLU:HG3	1.98	0.46
1:C:259:LEU:O	1:C:263:VAL:HG23	2.15	0.46
1:C:342:ILE:O	1:C:346:VAL:HG23	2.16	0.46
1:E:187:LEU:HD13	1:E:379:ILE:HG12	1.97	0.46
1:G:171:LYS:HD3	1:G:407:VAL:HG11	1.98	0.46
1:H:69:MET:CE	1:I:39:VAL:HG12	2.46	0.46
1:I:233:MET:HB3	1:I:309:LEU:CD2	2.46	0.46
1:I:252:GLU:O	1:I:277:LYS:HG3	2.15	0.46
1:I:409:GLU:CD	1:I:501:ARG:HH21	2.18	0.46
1:I:2:ALA:N	1:J:61:GLU:OE1	2.48	0.46
1:K:112:ASN:HA	1:K:113:PRO:HD3	1.85	0.46
1:N:421:ARG:NH1	1:N:472:GLY:O	2.44	0.46
1:N:77:VAL:CG2	1:N:507:ALA:CA	2.86	0.46
2:P:14:ARG:NH1	2:P:84:LEU:HD21	2.30	0.46
2:Q:41:LEU:O	2:Q:61:VAL:HG22	2.16	0.46
2:S:40:VAL:HG23	2:S:63:ASP:O	2.16	0.46
2:T:24:GLY:HA2	2:T:25:ILE:HA	1.37	0.46

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:X:49:LEU:O	2:X:52:GLY:N	2.48	0.46
2:Z:64:ILE:CG2	2:Z:95:VAL:HG22	2.42	0.46
1:C:350:ARG:HA	1:C:353:ILE:CD1	2.45	0.45
1:E:255:GLU:OE1	1:E:255:GLU:N	2.49	0.45
1:E:364:LYS:O	1:E:367:GLU:HB3	2.16	0.45
1:F:325:ILE:HD12	1:F:325:ILE:N	2.31	0.45
1:H:291:ASP:OD2	1:H:368:ARG:HD3	2.16	0.45
1:I:193:MET:HE1	1:I:292:ILE:CG1	2.46	0.45
1:J:245:LYS:HB3	1:J:245:LYS:HE2	1.60	0.45
1:L:488:MET:HE3	1:L:493:ILE:CG2	2.47	0.45
1:M:215:LEU:HD22	1:M:246:PRO:CB	2.46	0.45
1:M:234:LEU:CB	1:M:235:PRO:CD	2.84	0.45
1:N:116:LEU:O	1:N:120:ILE:HG13	2.16	0.45
1:N:221:LEU:HD13	1:N:317:LEU:HD23	1.97	0.45
1:N:91:THR:OG1	3:N:601:ADP:O2B	2.31	0.45
2:Q:55:LYS:HE3	2:R:52:GLY:HA2	1.98	0.45
2:R:7:HIS:O	2:R:44:GLY:HA3	2.16	0.45
2:X:66:ILE:HG21	2:Y:76:GLU:HG3	1.96	0.45
2:Z:48:ILE:HG13	2:Z:52:GLY:O	2.17	0.45
2:1:41:LEU:O	2:1:61:VAL:HG13	2.16	0.45
1:D:237:LEU:HD22	2:R:27:LEU:CD1	2.42	0.45
1:D:510:VAL:O	1:D:514:MET:HB2	2.17	0.45
1:E:219:PHE:HB3	1:E:317:LEU:CB	2.41	0.45
1:E:264:VAL:O	1:E:268:ARG:HG2	2.17	0.45
1:F:322:ARG:HG2	1:F:323:VAL:H	1.81	0.45
1:G:151:SER:HB3	1:G:399:ALA:HA	1.99	0.45
1:J:230:ILE:O	1:J:231:ARG:HG2	2.16	0.45
1:J:305:ILE:CG2	1:J:307:MET:H	2.18	0.45
1:J:325:ILE:CG1	1:J:330:THR:HG23	2.40	0.45
1:J:291:ASP:OD2	1:J:368:ARG:HD3	2.17	0.45
1:K:40:LEU:HD11	1:K:56:VAL:HA	1.97	0.45
1:M:231:ARG:HB3	1:M:258:ALA:HB2	1.98	0.45
1:E:265:ASN:OD1	2:S:27:LEU:HD22	2.16	0.45
2:T:78:ILE:HG13	2:T:79:ASP:N	2.31	0.45
2:U:11:ILE:HG23	2:U:42:ALA:H	1.80	0.45
2:Y:5:PRO:HD3	2:Y:42:ALA:HB3	1.98	0.45
1:A:205:ILE:HA	1:A:213:VAL:HG22	1.98	0.45
1:A:230:ILE:O	1:A:230:ILE:HG22	2.16	0.45
1:B:197:ARG:HH21	1:B:277:LYS:HD3	1.82	0.45
1:D:254:VAL:CG1	1:D:259:LEU:HG	2.46	0.45
1:D:357:THR:HB	1:D:361:ASP:CG	2.37	0.45

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:E:220:ILE:HD12	1:E:220:ILE:O	2.16	0.45
1:F:175:ILE:HB	1:F:400:LEU:HD11	1.98	0.45
1:F:228:SER:HB2	1:F:255:GLU:HB2	1.98	0.45
1:H:210:THR:HG22	1:H:210:THR:O	2.16	0.45
1:H:266:THR:HG22	1:H:273:VAL:H	1.82	0.45
1:H:40:LEU:HD11	1:H:56:VAL:HA	1.97	0.45
1:I:62:LEU:HD23	1:I:67:GLU:HB3	1.99	0.45
1:J:162:ILE:O	1:J:166:MET:HG3	2.16	0.45
1:J:208:PRO:HG2	1:J:214:GLU:HG3	1.97	0.45
1:L:225:LYS:CG	1:L:226:LYS:N	2.79	0.45
1:L:247:LEU:C	1:L:247:LEU:HD13	2.36	0.45
2:S:16:GLU:CG	2:S:19:THR:HG23	2.46	0.45
2:T:48:ILE:HG13	2:T:48:ILE:O	2.16	0.45
2:Y:13:LYS:HG3	2:Y:83:VAL:HG12	1.99	0.45
1:A:227:ILE:CG2	1:A:231:ARG:HA	2.47	0.45
1:B:513:LEU:HD11	1:C:388:GLU:HA	1.98	0.45
1:C:263:VAL:O	1:C:267:MET:HG2	2.16	0.45
1:C:88:GLY:N	4:C:602:BEF:F2	2.36	0.45
1:E:200:LEU:HD22	1:E:254:VAL:HB	1.99	0.45
1:E:226:LYS:HE2	1:E:255:GLU:OE2	2.15	0.45
1:E:409:GLU:CD	1:E:501:ARG:HH21	2.18	0.45
1:G:157:THR:O	1:G:161:LEU:HB2	2.15	0.45
1:G:358:SER:CA	1:G:362:ARG:HE	2.18	0.45
1:H:149:THR:HG23	1:H:155:ASP:C	2.37	0.45
1:I:207:LYS:N	1:I:208:PRO:CD	2.80	0.45
1:I:239:ALA:CB	1:I:314:LEU:HD21	2.47	0.45
1:K:169:VAL:HB	1:K:375:GLY:O	2.17	0.45
1:K:475:ASN:O	1:K:488:MET:HG2	2.16	0.45
1:L:221:LEU:HG	1:L:248:LEU:O	2.16	0.45
1:L:241:ALA:HA	1:L:271:VAL:CG1	2.46	0.45
1:L:433:ASN:OD1	1:L:436:GLN:HG3	2.16	0.45
1:M:496:PRO:O	1:M:499:VAL:HG22	2.17	0.45
1:N:358:SER:OG	1:N:359:ASP:N	2.49	0.45
2:O:15:LYS:HG2	2:O:37:ARG:O	2.17	0.45
2:U:73:VAL:HG12	2:U:86:MET:HB2	1.98	0.45
2:X:25:ILE:O	2:X:25:ILE:HG22	2.16	0.45
2:X:14:ARG:HG3	2:X:84:LEU:HD13	1.99	0.45
1:A:207:LYS:HB3	1:A:208:PRO:HD3	1.99	0.45
1:B:239:ALA:CB	1:B:314:LEU:HD11	2.46	0.45
1:B:305:ILE:CG2	1:B:306:GLY:N	2.75	0.45
1:D:203:TYR:HB3	1:D:267:MET:HE2	1.98	0.45

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:E:322:ARG:HG2	1:E:323:VAL:N	2.31	0.45
1:I:152:ALA:HB2	1:I:399:ALA:CB	2.44	0.45
1:J:142:LYS:O	1:J:146:GLN:HG3	2.16	0.45
1:J:227:ILE:HD12	1:J:227:ILE:H	1.80	0.45
1:K:217:SER:HA	1:K:320:ALA:O	2.17	0.45
1:K:54:VAL:HG22	1:K:58:ARG:HE	1.81	0.45
1:L:149:THR:CG2	1:L:156:GLU:HA	2.47	0.45
1:L:204:PHE:C	1:L:213:VAL:HG22	2.37	0.45
1:L:216:GLU:CD	1:L:217:SER:N	2.70	0.45
1:N:69:MET:O	1:N:73:MET:HG3	2.16	0.45
2:P:25:ILE:C	2:P:25:ILE:HD12	2.36	0.45
2:Q:88:GLU:HA	2:Q:91:ILE:HG22	1.98	0.45
2:V:57:LEU:O	2:V:59:VAL:HG13	2.17	0.45
2:Z:11:ILE:O	2:Z:41:LEU:HB2	2.17	0.45
1:A:419:LEU:H	1:A:419:LEU:HD22	1.82	0.45
1:B:338:GLU:HG3	1:B:340:ALA:H	1.82	0.45
1:D:423:ALA:HA	1:D:444:LEU:HD22	1.99	0.45
1:E:34:LYS:HD2	1:E:458:CYS:SG	2.57	0.45
1:H:224:ASP:OD1	1:H:286:LYS:HG3	2.17	0.45
1:J:293:ALA:O	1:J:298:GLY:N	2.49	0.45
1:K:361:ASP:O	1:K:365:LEU:HG	2.17	0.45
1:L:392:LYS:O	1:L:396:VAL:HG23	2.16	0.45
1:M:364:LYS:HD3	1:M:364:LYS:C	2.37	0.45
1:M:92:ALA:HB2	1:M:503:ALA:HB1	1.97	0.45
2:O:51:ASN:ND2	2:P:50:GLU:O	2.49	0.45
2:P:55:LYS:HA	2:P:55:LYS:HD2	1.75	0.45
2:U:28:THR:HG22	2:U:29:GLY:O	2.17	0.45
2:W:50:GLU:OE1	2:W:50:GLU:N	2.50	0.45
1:A:222:LEU:N	1:A:222:LEU:HD12	2.32	0.45
1:A:443:ALA:O	1:A:447:MET:HG3	2.17	0.45
1:B:23:LEU:HD23	1:B:60:ILE:HB	1.99	0.45
1:D:177:VAL:HG21	1:D:393:LYS:HG3	1.96	0.45
1:D:78:ALA:HB2	1:D:93:THR:OG1	2.17	0.45
1:E:360:TYR:O	1:E:360:TYR:CG	2.70	0.45
1:E:73:MET:HG3	1:F:47:PRO:HG2	1.98	0.45
1:F:295:LEU:C	1:F:295:LEU:HD23	2.37	0.45
1:H:268:ARG:HD2	1:H:268:ARG:HA	1.78	0.45
1:H:432:GLN:HB2	1:H:436:GLN:OE1	2.17	0.45
1:I:193:MET:CE	1:I:292:ILE:CG1	2.95	0.45
1:J:240:VAL:HA	1:J:314:LEU:CD1	2.47	0.45
1:K:208:PRO:CB	1:K:212:ALA:HB3	2.32	0.45

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:L:237:LEU:HD23	1:L:237:LEU:O	2.17	0.45
1:L:434:GLU:O	1:L:437:ASN:HB2	2.16	0.45
1:M:164:GLU:O	1:M:168:LYS:HB2	2.17	0.45
1:N:227:ILE:HG12	1:N:231:ARG:HG3	1.98	0.45
1:N:419:LEU:O	1:N:422:VAL:HG22	2.17	0.45
2:T:23:GLY:O	2:T:26:VAL:HG12	2.17	0.45
2:U:49:LEU:O	2:U:50:GLU:CG	2.64	0.45
2:X:16:GLU:O	2:X:16:GLU:HG2	2.16	0.45
1:A:278:ALA:HB3	1:A:285:ARG:HD3	1.99	0.45
1:A:518:GLU:HB2	1:B:36:ARG:HB2	1.99	0.45
1:B:222:LEU:N	1:B:222:LEU:HD12	2.32	0.45
1:D:124:VAL:HG13	1:D:504:LEU:CD1	2.47	0.45
1:E:414:GLY:HA2	1:E:495:ASP:OD2	2.17	0.45
1:E:31:LEU:HA	3:E:601:ADP:O2A	2.16	0.45
1:F:264:VAL:O	1:F:268:ARG:HG2	2.17	0.45
1:F:304:GLU:CB	1:F:305:ILE:HD12	2.43	0.45
1:F:301:ILE:HG12	1:F:307:MET:HG3	1.99	0.45
1:F:381:VAL:HG21	1:F:392:LYS:HG3	1.99	0.45
1:G:288:MET:O	1:G:292:ILE:HG13	2.17	0.45
1:G:313:THR:HB	1:G:315:GLU:CG	2.47	0.45
1:H:228:SER:HB2	1:H:255:GLU:OE1	2.17	0.45
1:H:414:GLY:O	1:H:417:VAL:HG12	2.17	0.45
1:H:419:LEU:HD22	1:H:419:LEU:H	1.81	0.45
1:I:307:MET:O	1:I:307:MET:HG3	2.17	0.45
1:K:248:LEU:HD13	1:K:248:LEU:C	2.37	0.45
1:L:488:MET:CE	1:L:493:ILE:HG21	2.47	0.45
1:L:28:LYS:HG2	1:L:94:VAL:HG22	1.99	0.45
1:M:200:LEU:CD2	1:M:254:VAL:HB	2.44	0.45
1:M:349:ILE:HG21	1:M:369:VAL:CG2	2.47	0.45
2:Q:15:LYS:HB3	2:Q:16:GLU:OE2	2.17	0.45
2:S:82:GLU:O	2:S:82:GLU:HG2	2.15	0.45
2:U:43:VAL:HG22	2:U:44:GLY:N	2.32	0.45
2:Y:57:LEU:HD13	2:Y:88:GLU:HB2	1.99	0.45
2:Z:39:GLU:HA	2:Z:64:ILE:HA	1.99	0.45
2:1:31:ALA:O	2:1:32:ALA:HB3	2.17	0.45
1:A:359:ASP:O	1:A:360:TYR:HB2	2.17	0.45
1:B:247:LEU:C	1:B:248:LEU:HD22	2.37	0.45
1:G:220:ILE:N	1:G:220:ILE:HD12	2.31	0.45
1:G:266:THR:CG2	1:G:273:VAL:HB	2.47	0.45
1:G:222:LEU:CD2	1:G:300:VAL:HG22	2.37	0.45
1:H:363:GLU:O	1:H:367:GLU:HB2	2.17	0.45

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:H:42:LYS:HE2	1:H:48:THR:CG2	2.47	0.45
1:K:259:LEU:O	1:K:263:VAL:HG23	2.17	0.45
1:K:285:ARG:HG3	1:K:286:LYS:N	2.32	0.45
1:L:230:ILE:O	1:L:232:GLU:HG2	2.16	0.45
1:L:25:ASP:OD1	1:L:28:LYS:HE2	2.16	0.45
1:L:320:ALA:HB1	1:L:334:ASP:O	2.17	0.45
1:L:361:ASP:O	1:L:365:LEU:HD13	2.17	0.45
1:H:203:TYR:HE2	1:N:305:ILE:HD11	1.81	0.45
2:Q:54:VAL:O	2:Q:54:VAL:HG23	2.17	0.45
2:U:78:ILE:HD12	2:U:78:ILE:C	2.37	0.45
2:X:94:ILE:HD11	2:Y:4:ARG:HD3	1.98	0.45
1:B:18:ARG:NH1	1:B:18:ARG:HB3	2.32	0.45
1:B:262:LEU:O	1:B:266:THR:HG23	2.16	0.45
1:B:48:THR:HG22	1:B:390:LYS:HZ1	1.80	0.45
1:C:226:LYS:HA	1:C:252:GLU:HB3	1.99	0.45
1:C:226:LYS:HE2	1:C:252:GLU:HG2	1.99	0.45
1:E:230:ILE:HD12	1:E:230:ILE:H	1.82	0.45
1:G:429:LEU:HG	1:G:440:ILE:HD13	1.98	0.45
1:H:10:ASN:O	1:H:14:VAL:HG23	2.16	0.45
1:I:321:LYS:HD2	1:I:334:ASP:OD2	2.16	0.45
1:J:365:LEU:O	1:J:369:VAL:HG23	2.17	0.45
3:K:601:ADP:O3B	4:K:602:BEF:F1	2.25	0.45
1:L:282:GLY:CA	1:L:285:ARG:HG2	2.45	0.45
1:M:228:SER:HB3	1:M:255:GLU:HB3	1.99	0.45
1:M:236:VAL:CG1	1:M:312:ALA:HB3	2.46	0.45
1:N:331:THR:HG22	1:N:332:ILE:N	2.32	0.45
2:R:12:VAL:HG23	2:R:84:LEU:HD11	1.99	0.45
2:R:11:ILE:HB	2:R:85:ILE:CD1	2.46	0.45
2:S:12:VAL:CG1	2:S:40:VAL:HA	2.43	0.45
1:A:174:VAL:HG23	1:A:376:VAL:HG13	1.99	0.44
1:B:232:GLU:O	1:B:232:GLU:HG2	2.17	0.44
1:D:278:ALA:CB	1:D:285:ARG:HG3	2.37	0.44
1:D:77:VAL:HB	1:D:510:VAL:HG21	1.98	0.44
1:E:496:PRO:O	1:E:499:VAL:HG22	2.17	0.44
1:E:510:VAL:O	1:E:514:MET:HB2	2.17	0.44
1:H:238:GLU:HG2	2:V:25:ILE:HG12	1.99	0.44
1:I:149:THR:HG23	1:I:155:ASP:C	2.38	0.44
1:I:202:PRO:O	1:I:205:ILE:HG13	2.17	0.44
1:I:294:THR:OG1	1:I:345:ARG:HG3	2.17	0.44
1:J:175:ILE:HA	1:J:377:ALA:O	2.17	0.44
1:J:197:ARG:HA	1:J:197:ARG:HD3	1.79	0.44

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:L:345:ARG:O	1:L:348:GLN:HG3	2.17	0.44
1:M:196:ASP:O	1:M:197:ARG:HG2	2.18	0.44
2:O:57:LEU:HD22	2:O:57:LEU:N	2.32	0.44
2:P:20:LYS:HG2	2:P:27:LEU:HD23	1.99	0.44
2:S:6:LEU:H	2:S:6:LEU:HD12	1.81	0.44
2:W:50:GLU:C	2:W:52:GLY:H	2.21	0.44
1:A:247:LEU:CD1	1:A:249:ILE:HD11	2.47	0.44
1:A:271:VAL:HG11	2:O:25:ILE:CD1	2.45	0.44
1:B:124:VAL:HG21	1:B:508:ALA:CB	2.47	0.44
1:B:86:GLY:HA3	1:B:401:HIS:ND1	2.31	0.44
1:C:196:ASP:OD1	1:C:196:ASP:N	2.48	0.44
1:E:493:ILE:C	1:E:494:LEU:HD12	2.38	0.44
1:F:245:LYS:CB	1:F:246:PRO:CD	2.94	0.44
1:F:259:LEU:O	1:F:263:VAL:HG23	2.17	0.44
1:G:196:ASP:OD1	1:G:196:ASP:N	2.50	0.44
1:G:358:SER:HA	1:G:362:ARG:NH2	2.32	0.44
1:H:226:LYS:HG2	1:H:252:GLU:CG	2.46	0.44
1:I:230:ILE:O	1:I:230:ILE:HG22	2.17	0.44
1:I:204:PHE:CE1	1:I:263:VAL:HG22	2.51	0.44
1:I:31:LEU:CD2	1:I:453:GLN:HB3	2.45	0.44
1:I:485:TYR:O	1:I:491:MET:HE1	2.17	0.44
1:K:37:ASN:OD1	1:K:51:LYS:HB2	2.17	0.44
1:M:259:LEU:O	1:M:262:LEU:HB3	2.17	0.44
1:N:196:ASP:OD1	1:N:196:ASP:N	2.48	0.44
2:S:38:GLY:O	2:S:64:ILE:HA	2.18	0.44
2:U:15:LYS:HA	2:U:15:LYS:HD3	1.69	0.44
2:U:2:ASN:C	2:U:3:ILE:HD12	2.37	0.44
2:Z:47:ARG:NE	2:Z:49:LEU:HD12	2.32	0.44
1:A:278:ALA:HB3	1:A:285:ARG:CD	2.47	0.44
1:A:329:THR:CG2	1:A:330:THR:N	2.80	0.44
1:B:245:LYS:N	1:B:246:PRO:HD2	2.32	0.44
1:C:221:LEU:HB2	1:C:317:LEU:CD2	2.47	0.44
1:G:284:ARG:HG2	1:G:288:MET:HE1	1.99	0.44
1:G:33:PRO:CG	1:G:480:ALA:HB3	2.47	0.44
1:I:210:THR:HG22	1:I:210:THR:O	2.16	0.44
1:J:238:GLU:HG3	1:J:242:LYS:HZ1	1.81	0.44
1:J:325:ILE:HD12	1:J:325:ILE:N	2.32	0.44
1:J:353:ILE:HG12	1:J:365:LEU:CD2	2.46	0.44
1:J:69:MET:SD	1:K:41:ASP:HB2	2.57	0.44
1:K:302:SER:HB2	1:K:305:ILE:CD1	2.36	0.44
1:L:217:SER:O	1:L:245:LYS:HE3	2.16	0.44

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:L:253:ASP:OD1	1:L:277:LYS:HE2	2.17	0.44
1:M:234:LEU:O	1:M:237:LEU:HG	2.17	0.44
1:M:236:VAL:HG11	1:M:312:ALA:HB3	1.99	0.44
1:M:147:VAL:HG22	1:M:403:THR:HG22	1.99	0.44
2:O:6:LEU:HD11	2:U:94:ILE:H	1.81	0.44
2:V:57:LEU:C	2:V:57:LEU:HD12	2.38	0.44
2:Z:6:LEU:O	2:Z:6:LEU:HD12	2.16	0.44
1:B:325:ILE:HD12	1:B:325:ILE:N	2.33	0.44
1:C:177:VAL:CG1	1:C:397:GLU:HG2	2.22	0.44
1:C:419:LEU:N	1:C:419:LEU:HD22	2.32	0.44
1:D:419:LEU:H	1:D:419:LEU:HD22	1.83	0.44
1:E:197:ARG:CG	1:E:279:PRO:HA	2.45	0.44
1:F:222:LEU:N	1:F:222:LEU:HD12	2.33	0.44
1:F:329:THR:CG2	1:F:330:THR:N	2.80	0.44
1:I:367:GLU:CG	1:I:371:LYS:HE2	2.46	0.44
1:I:176:THR:OG1	1:I:378:VAL:HG22	2.18	0.44
1:J:493:ILE:C	1:J:494:LEU:HD12	2.38	0.44
1:K:168:LYS:HG2	1:K:189:VAL:CG1	2.45	0.44
1:K:207:LYS:HG3	1:K:208:PRO:N	2.30	0.44
1:M:365:LEU:O	1:M:369:VAL:HG23	2.17	0.44
1:N:220:ILE:O	1:N:317:LEU:HA	2.17	0.44
2:O:6:LEU:N	2:O:6:LEU:CD1	2.80	0.44
2:V:14:ARG:O	2:V:14:ARG:HG3	2.18	0.44
2:V:40:VAL:HG11	2:V:60:LYS:O	2.17	0.44
2:Y:15:LYS:HD3	2:Y:37:ARG:CB	2.46	0.44
2:Z:17:VAL:HG13	2:Z:18:GLU:OE2	2.17	0.44
1:B:215:LEU:CD1	1:B:248:LEU:HD21	2.44	0.44
1:B:429:LEU:HD12	1:B:430:ARG:N	2.32	0.44
1:B:421:ARG:NH1	1:B:472:GLY:O	2.50	0.44
1:D:196:ASP:O	1:D:197:ARG:HG2	2.16	0.44
1:G:226:LYS:HE2	1:G:253:ASP:HB2	1.98	0.44
1:G:198:GLY:O	1:G:276:VAL:HG23	2.17	0.44
1:I:284:ARG:HD2	1:I:364:LYS:HZ3	1.82	0.44
1:K:155:ASP:CG	1:K:158:VAL:HG23	2.38	0.44
1:L:200:LEU:HD22	1:L:254:VAL:HB	2.00	0.44
1:N:311:LYS:O	1:N:311:LYS:HD3	2.17	0.44
2:V:14:ARG:NE	2:V:84:LEU:HD21	2.33	0.44
2:W:9:ARG:H	2:W:9:ARG:HG3	1.58	0.44
2:Z:15:LYS:HD2	2:Z:16:GLU:OE1	2.18	0.44
1:A:305:ILE:HG23	1:A:306:GLY:N	2.32	0.44
1:B:236:VAL:O	1:B:240:VAL:HG23	2.17	0.44

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:B:391:GLU:OE2	1:B:395:ARG:NE	2.43	0.44
1:C:260:ALA:O	1:C:264:VAL:HG23	2.18	0.44
1:D:230:ILE:C	1:D:231:ARG:HG2	2.33	0.44
1:D:236:VAL:HG22	1:D:312:ALA:HB3	1.99	0.44
1:D:240:VAL:HG12	1:D:245:LYS:O	2.17	0.44
1:E:143:ALA:O	1:E:147:VAL:HG13	2.17	0.44
1:E:367:GLU:O	1:E:371:LYS:HG3	2.17	0.44
1:E:517:THR:HG23	1:F:39:VAL:HG23	1.99	0.44
1:G:285:ARG:HG3	1:G:286:LYS:N	2.33	0.44
1:G:361:ASP:OD1	1:G:361:ASP:N	2.51	0.44
1:H:199:TYR:CD1	1:H:202:PRO:HA	2.53	0.44
1:I:280:GLY:O	1:I:285:ARG:HB3	2.17	0.44
1:J:222:LEU:HD12	1:J:222:LEU:N	2.32	0.44
1:M:221:LEU:HB2	1:M:317:LEU:CD2	2.47	0.44
1:N:24:ALA:O	1:N:28:LYS:HG3	2.17	0.44
2:O:40:VAL:HG23	2:O:63:ASP:O	2.17	0.44
2:Z:47:ARG:HG3	2:Z:49:LEU:HG	2.00	0.44
2:2:17:VAL:CG1	2:2:35:SER:H	2.31	0.44
1:A:34:LYS:HD2	1:A:458:CYS:SG	2.58	0.44
1:B:205:ILE:HD12	1:B:211:GLY:HA2	2.00	0.44
1:B:429:LEU:HD12	1:B:430:ARG:H	1.82	0.44
1:B:77:VAL:HG21	1:B:507:ALA:CA	2.39	0.44
1:C:23:LEU:HD13	1:C:23:LEU:C	2.38	0.44
1:F:414:GLY:O	1:F:417:VAL:HG12	2.17	0.44
1:H:222:LEU:HD12	1:H:222:LEU:N	2.32	0.44
1:I:417:VAL:CA	1:I:420:ILE:HG22	2.46	0.44
1:J:421:ARG:HD3	1:J:421:ARG:HA	1.68	0.44
1:K:353:ILE:HG12	1:K:365:LEU:HB3	2.00	0.44
1:L:412:VAL:CG1	1:L:413:ALA:H	2.31	0.44
1:M:200:LEU:HD22	1:M:254:VAL:CB	2.46	0.44
1:N:142:LYS:O	1:N:146:GLN:HG3	2.17	0.44
2:Z:6:LEU:O	2:Z:9:ARG:HG2	2.18	0.44
1:A:195:PHE:CE1	1:A:330:THR:HB	2.53	0.44
1:A:367:GLU:HG3	1:A:371:LYS:NZ	2.33	0.44
1:A:475:ASN:O	1:A:488:MET:HG2	2.17	0.44
1:B:478:TYR:HD1	1:B:485:TYR:CE1	2.35	0.44
1:B:73:MET:SD	1:C:49:ILE:HD11	2.57	0.44
1:C:143:ALA:O	1:C:147:VAL:HG13	2.18	0.44
1:B:305:ILE:CD1	1:C:203:TYR:HE2	2.31	0.44
1:C:215:LEU:HD22	1:C:246:PRO:HB2	2.00	0.44
1:B:305:ILE:CD1	1:C:263:VAL:HG11	2.44	0.44

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:D:155:ASP:CG	1:D:158:VAL:HG23	2.38	0.44
1:D:254:VAL:HB	1:D:259:LEU:HD21	1.99	0.44
1:D:329:THR:CG2	1:D:330:THR:N	2.81	0.44
1:D:326:ASN:ND2	1:D:329:THR:H	2.15	0.44
1:D:393:LYS:O	1:D:397:GLU:HG3	2.17	0.44
1:E:248:LEU:CD2	1:E:323:VAL:HG11	2.36	0.44
1:G:124:VAL:O	1:G:128:VAL:HG23	2.17	0.44
1:H:365:LEU:O	1:H:369:VAL:HG23	2.18	0.44
1:J:116:LEU:HD23	1:J:435:ASP:O	2.18	0.44
1:K:371:LYS:HE2	1:K:376:VAL:HG21	2.00	0.44
1:L:271:VAL:O	1:L:273:VAL:HG23	2.16	0.44
1:M:90:THR:O	1:M:94:VAL:HG23	2.17	0.44
1:N:248:LEU:C	1:N:249:ILE:HD12	2.38	0.44
2:O:94:ILE:HG13	2:P:4:ARG:O	2.17	0.44
2:R:80:ASN:OD1	2:R:81:GLU:HG3	2.18	0.44
2:R:57:LEU:HD22	2:R:88:GLU:HB2	2.00	0.44
1:A:392:LYS:O	1:A:396:VAL:HG23	2.17	0.44
1:B:136:VAL:HG22	1:B:137:PRO:CD	2.48	0.44
1:B:155:ASP:CG	1:B:158:VAL:HG23	2.38	0.44
1:B:450:PRO:O	1:B:454:ILE:HG13	2.18	0.44
1:C:200:LEU:HB3	1:C:259:LEU:CD1	2.48	0.44
1:D:234:LEU:N	1:D:235:PRO:CD	2.80	0.44
1:E:361:ASP:O	1:E:365:LEU:HB2	2.18	0.44
1:F:183:LEU:HD22	1:F:384:ALA:HA	2.00	0.44
1:H:114:MET:HG3	1:H:118:ARG:NH1	2.33	0.44
1:K:268:ARG:HB2	1:K:270:ILE:HD12	1.99	0.44
1:L:127:ALA:O	1:L:131:LEU:HG	2.17	0.44
1:L:282:GLY:HA2	1:L:285:ARG:CG	2.47	0.44
1:M:364:LYS:HZ3	1:M:368:ARG:HE	1.66	0.44
1:N:259:LEU:HA	1:N:262:LEU:HB3	2.00	0.44
2:R:25:ILE:HA	2:R:26:VAL:CG2	2.48	0.44
2:S:41:LEU:O	2:S:61:VAL:HG23	2.17	0.44
2:U:50:GLU:C	2:U:52:GLY:N	2.71	0.44
2:U:50:GLU:OE1	2:U:50:GLU:N	2.49	0.44
2:Y:7:HIS:O	2:Y:8:ASP:CB	2.66	0.44
1:A:205:ILE:HG22	1:A:206:ASN:O	2.18	0.43
1:A:213:VAL:HB	1:A:325:ILE:HB	2.00	0.43
1:A:147:VAL:CG1	1:A:494:LEU:HD12	2.48	0.43
1:B:496:PRO:HG2	1:B:499:VAL:HG11	1.99	0.43
1:C:218:PRO:O	1:C:319:GLN:HA	2.18	0.43
1:D:166:MET:HE2	1:D:407:VAL:HG21	2.00	0.43

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:F:177:VAL:HG22	1:F:393:LYS:HG3	1.99	0.43
1:G:323:VAL:HG12	1:G:332:ILE:HG12	1.99	0.43
1:I:295:LEU:O	1:I:295:LEU:HD23	2.18	0.43
1:J:265:ASN:CB	1:J:270:ILE:HB	2.32	0.43
1:J:350:ARG:HA	1:J:353:ILE:HG13	2.00	0.43
1:J:23:LEU:HD22	1:J:60:ILE:HG13	1.98	0.43
1:K:197:ARG:HA	1:K:197:ARG:HD3	1.82	0.43
1:L:230:ILE:O	1:L:231:ARG:HB2	2.17	0.43
1:L:225:LYS:HD3	1:L:303:GLU:OE1	2.18	0.43
1:M:200:LEU:HD11	1:M:277:LYS:HG2	2.00	0.43
1:N:157:THR:O	1:N:161:LEU:HB2	2.18	0.43
2:Q:40:VAL:HG21	2:Q:60:LYS:O	2.17	0.43
2:U:3:ILE:CD1	2:U:78:ILE:HD13	2.46	0.43
2:Y:91:ILE:HG22	2:Z:6:LEU:HD21	2.00	0.43
1:B:96:ALA:O	1:B:100:ILE:HG13	2.19	0.43
1:B:198:GLY:HA3	1:B:327:LYS:O	2.18	0.43
1:B:368:ARG:O	1:B:372:LEU:HG	2.18	0.43
1:C:231:ARG:HH12	1:C:234:LEU:CD2	2.29	0.43
1:C:364:LYS:HA	1:C:367:GLU:CB	2.48	0.43
1:G:158:VAL:O	1:G:162:ILE:HG13	2.18	0.43
1:G:313:THR:CB	1:G:315:GLU:HG2	2.49	0.43
1:H:207:LYS:N	1:H:208:PRO:CD	2.81	0.43
1:H:77:VAL:CG2	1:H:507:ALA:HA	2.39	0.43
1:I:367:GLU:O	1:I:371:LYS:HG3	2.17	0.43
1:J:358:SER:OG	1:J:359:ASP:N	2.50	0.43
1:K:213:VAL:HB	1:K:325:ILE:HB	1.99	0.43
1:K:232:GLU:N	1:K:232:GLU:OE1	2.50	0.43
1:K:99:ILE:CG2	1:K:511:ALA:HB1	2.47	0.43
1:M:217:SER:N	1:M:218:PRO:HD3	2.33	0.43
1:M:285:ARG:HG3	1:M:286:LYS:N	2.34	0.43
1:A:95:LEU:O	1:A:99:ILE:HG13	2.17	0.43
1:E:136:VAL:HG22	1:E:137:PRO:CD	2.48	0.43
1:F:393:LYS:O	1:F:397:GLU:HG3	2.18	0.43
1:F:421:ARG:HA	1:F:421:ARG:HD3	1.75	0.43
1:G:209:GLU:CG	1:G:210:THR:N	2.79	0.43
1:G:223:ALA:HB1	1:G:225:LYS:O	2.19	0.43
1:H:25:ASP:OD1	1:H:28:LYS:HE2	2.19	0.43
1:I:213:VAL:HB	1:I:325:ILE:HB	2.01	0.43
1:I:219:PHE:HD1	1:I:319:GLN:HG2	1.84	0.43
1:I:401:HIS:O	1:I:404:ARG:HB2	2.19	0.43
1:I:496:PRO:HG2	1:I:499:VAL:CG1	2.48	0.43

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:J:208:PRO:HG2	1:J:214:GLU:CG	2.49	0.43
1:J:238:GLU:HG3	1:J:242:LYS:NZ	2.34	0.43
1:K:218:PRO:CA	1:K:246:PRO:HG2	2.47	0.43
1:K:69:MET:HB3	1:K:69:MET:HE2	1.89	0.43
1:L:510:VAL:HG12	1:M:387:VAL:HG21	2.00	0.43
1:N:200:LEU:HD12	1:N:200:LEU:N	2.33	0.43
1:N:199:TYR:CD1	1:N:202:PRO:HA	2.52	0.43
1:N:197:ARG:HG2	1:N:277:LYS:CA	2.48	0.43
2:P:28:THR:HG22	2:P:29:GLY:O	2.18	0.43
2:Q:50:GLU:HB3	2:Q:52:GLY:H	1.83	0.43
2:Q:92:LEU:O	2:R:6:LEU:HB3	2.18	0.43
2:T:60:LYS:HG3	2:T:61:VAL:H	1.82	0.43
2:U:65:VAL:CG1	2:U:94:ILE:HG22	2.48	0.43
2:X:73:VAL:CG2	2:X:84:LEU:HB3	2.48	0.43
2:V:78:ILE:HD13	2:2:37:ARG:HH22	1.83	0.43
1:B:219:PHE:HB3	1:B:317:LEU:HB3	2.00	0.43
1:B:367:GLU:CG	1:B:371:LYS:HE2	2.43	0.43
1:C:421:ARG:NH1	1:C:472:GLY:O	2.50	0.43
1:H:220:ILE:CG2	1:H:222:LEU:HD11	2.44	0.43
1:H:234:LEU:N	1:H:235:PRO:CD	2.80	0.43
1:H:305:ILE:HG23	1:I:267:MET:HE1	1.96	0.43
1:H:147:VAL:HG23	1:H:403:THR:HG22	1.99	0.43
1:I:23:LEU:HD23	1:I:60:ILE:HB	2.00	0.43
1:J:232:GLU:O	1:J:310:GLU:HG3	2.18	0.43
1:K:168:LYS:HG2	1:K:189:VAL:CG2	2.48	0.43
1:K:496:PRO:HG2	1:K:499:VAL:CG2	2.49	0.43
1:L:496:PRO:HG2	1:L:499:VAL:HG21	1.97	0.43
1:M:177:VAL:HG23	1:M:381:VAL:HG23	2.00	0.43
1:M:65:LYS:HB3	1:M:522:THR:HG21	2.00	0.43
1:N:225:LYS:CG	1:N:226:LYS:N	2.81	0.43
2:Q:94:ILE:HD11	2:R:4:ARG:HH21	1.83	0.43
2:U:77:LYS:HG2	2:U:82:GLU:HA	2.00	0.43
2:V:40:VAL:HG13	2:V:41:LEU:N	2.33	0.43
2:Y:11:ILE:HG12	2:Y:42:ALA:HB3	2.01	0.43
2:1:66:ILE:HD12	2:2:3:ILE:HD12	2.00	0.43
2:2:27:LEU:HD12	2:2:27:LEU:N	2.34	0.43
1:B:278:ALA:HB1	1:B:279:PRO:HD2	1.99	0.43
1:C:359:ASP:O	1:C:360:TYR:HB3	2.18	0.43
1:D:233:MET:HB3	1:D:309:LEU:CD2	2.48	0.43
1:D:198:GLY:HA3	1:D:327:LYS:O	2.17	0.43
1:E:392:LYS:O	1:E:396:VAL:HG23	2.18	0.43

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:F:160:LYS:HD2	1:F:160:LYS:HA	1.82	0.43
1:G:106:ALA:O	1:G:109:ALA:HB3	2.19	0.43
1:H:228:SER:HA	1:H:255:GLU:HB2	2.00	0.43
1:H:267:MET:CE	1:H:267:MET:HA	2.48	0.43
1:H:381:VAL:HG21	1:H:393:LYS:HA	2.00	0.43
1:I:302:SER:OG	1:I:305:ILE:HD13	2.18	0.43
1:I:368:ARG:O	1:I:372:LEU:HG	2.19	0.43
1:K:23:LEU:HD13	1:K:23:LEU:C	2.39	0.43
1:M:115:ASP:O	1:M:436:GLN:HG2	2.18	0.43
1:M:143:ALA:O	1:M:147:VAL:HG13	2.18	0.43
1:M:419:LEU:N	1:M:419:LEU:HD22	2.34	0.43
1:N:301:ILE:HG12	1:N:307:MET:CE	2.49	0.43
2:R:76:GLU:HG3	2:R:77:LYS:N	2.33	0.43
2:Q:88:GLU:OE2	2:R:7:HIS:HE1	2.00	0.43
2:R:12:VAL:CG2	2:R:84:LEU:HD11	2.48	0.43
2:R:5:PRO:HB2	2:R:9:ARG:O	2.18	0.43
2:S:11:ILE:CG1	2:S:83:VAL:HG11	2.48	0.43
2:X:36:THR:OG1	2:Y:76:GLU:OE2	2.28	0.43
1:A:227:ILE:HD12	1:A:227:ILE:H	1.83	0.43
1:C:205:ILE:HG12	1:C:206:ASN:O	2.18	0.43
1:C:256:GLY:O	1:C:260:ALA:N	2.35	0.43
3:C:601:ADP:O3B	4:C:602:BEF:F2	2.26	0.43
1:D:112:ASN:HA	1:D:113:PRO:HD3	1.80	0.43
1:E:136:VAL:HG23	1:E:137:PRO:HD2	2.00	0.43
1:E:158:VAL:O	1:E:162:ILE:HG13	2.19	0.43
1:E:87:ASP:OD2	1:E:151:SER:OG	2.29	0.43
1:G:149:THR:HG22	1:G:156:GLU:HA	1.99	0.43
1:G:255:GLU:N	1:G:255:GLU:OE1	2.50	0.43
1:G:263:VAL:O	1:G:267:MET:HG2	2.19	0.43
1:I:200:LEU:CD2	1:I:254:VAL:HB	2.49	0.43
1:I:230:ILE:N	1:I:230:ILE:HD12	2.34	0.43
1:I:221:LEU:HD11	1:I:301:ILE:HD12	1.99	0.43
1:J:26:ALA:O	1:J:29:VAL:HG22	2.19	0.43
1:J:455:VAL:CG2	1:J:465:VAL:HG11	2.48	0.43
1:L:37:ASN:OD1	1:L:51:LYS:HB2	2.18	0.43
1:M:225:LYS:HD3	1:M:303:GLU:CD	2.38	0.43
1:M:455:VAL:HG21	1:M:465:VAL:HG11	2.01	0.43
2:O:96:GLU:HB2	2:P:2:ASN:O	2.19	0.43
2:R:74:LYS:HD2	2:R:75:SER:H	1.84	0.43
2:T:48:ILE:HG22	2:T:53:GLU:C	2.39	0.43
2:T:78:ILE:HG13	2:T:79:ASP:H	1.84	0.43

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:W:19:THR:HG22	2:W:20:LYS:N	2.33	0.43
2:W:12:VAL:HB	2:W:39:GLU:O	2.18	0.43
2:Z:8:ASP:OD1	2:Z:9:ARG:CA	2.66	0.43
2:Z:94:ILE:HG13	2:Z:94:ILE:O	2.19	0.43
1:B:171:LYS:HA	1:B:171:LYS:HD2	1.70	0.43
1:B:326:ASN:ND2	1:B:329:THR:O	2.51	0.43
1:C:215:LEU:HB2	1:C:323:VAL:HG22	2.00	0.43
1:C:220:ILE:HD13	1:C:296:THR:HB	2.00	0.43
1:D:17:LEU:HD13	1:D:104:LEU:HD12	2.00	0.43
1:F:124:VAL:O	1:F:128:VAL:HG23	2.19	0.43
1:G:112:ASN:HA	1:G:113:PRO:HD3	1.89	0.43
1:G:180:GLY:HA2	1:G:380:LYS:HG2	2.01	0.43
1:H:232:GLU:O	1:H:233:MET:HB3	2.18	0.43
1:H:40:LEU:HD13	1:H:59:GLU:CG	2.42	0.43
1:H:513:LEU:CD2	1:I:49:ILE:HG21	2.49	0.43
1:I:112:ASN:HA	1:I:113:PRO:HD3	1.84	0.43
1:I:415:GLY:O	1:I:451:LEU:HD23	2.17	0.43
1:K:117:LYS:HG3	1:K:512:GLY:C	2.39	0.43
1:K:282:GLY:HA2	1:K:285:ARG:CD	2.48	0.43
1:K:301:ILE:HD11	1:K:312:ALA:HB2	2.01	0.43
1:M:146:GLN:OE1	1:M:492:GLY:HA2	2.19	0.43
1:N:205:ILE:HD12	1:N:211:GLY:HA2	2.00	0.43
1:N:195:PHE:CZ	1:N:330:THR:HB	2.54	0.43
1:N:524:LEU:HD22	1:N:525:PRO:HD2	2.01	0.43
2:O:97:ALA:CB	2:P:1:MET:HB2	2.46	0.43
2:P:6:LEU:O	2:P:9:ARG:HD2	2.19	0.43
2:X:64:ILE:CG2	2:X:95:VAL:HG13	2.44	0.43
2:Z:95:VAL:HG12	2:1:3:ILE:HG21	1.99	0.43
2:2:49:LEU:H	2:2:52:GLY:CA	2.12	0.43
1:B:362:ARG:HB3	1:B:366:GLN:NE2	2.33	0.43
1:B:180:GLY:H	1:B:389:MET:HE1	1.84	0.43
1:C:230:ILE:HG22	1:C:232:GLU:HB3	2.01	0.43
1:C:247:LEU:CD1	1:C:249:ILE:HG12	2.49	0.43
1:C:203:TYR:HD2	1:C:263:VAL:CG1	2.31	0.43
1:C:285:ARG:O	1:C:289:LEU:N	2.52	0.43
1:D:417:VAL:C	1:D:420:ILE:HG22	2.39	0.43
1:E:190:VAL:HG11	1:E:194:GLN:CD	2.39	0.43
1:E:475:ASN:O	1:E:488:MET:HG2	2.18	0.43
1:I:420:ILE:HD11	1:I:470:LYS:CG	2.49	0.43
1:J:322:ARG:CB	1:J:333:ILE:HD12	2.34	0.43
1:J:342:ILE:O	1:J:346:VAL:HG23	2.19	0.43

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:K:200:LEU:HD22	1:K:254:VAL:HB	2.00	0.43
1:K:30:THR:HA	1:K:36:ARG:O	2.19	0.43
1:M:190:VAL:HG11	1:M:194:GLN:CD	2.39	0.43
1:M:30:THR:HB	1:M:51:LYS:HG3	2.01	0.43
1:N:103:GLY:O	1:N:107:VAL:HG23	2.19	0.43
1:A:322:ARG:HD2	1:A:333:ILE:HD12	2.01	0.43
1:A:138:CYS:HB2	1:A:411:VAL:HG13	2.01	0.43
1:A:438:VAL:O	1:A:442:VAL:HG23	2.19	0.43
1:B:205:ILE:HG12	1:B:206:ASN:O	2.19	0.43
1:A:519:CYS:HB3	1:B:38:VAL:HG22	2.01	0.43
1:C:257:GLU:O	1:C:261:THR:HG22	2.18	0.43
1:C:31:LEU:HD23	1:C:453:GLN:HB3	2.00	0.43
1:C:409:GLU:CD	1:C:501:ARG:HH21	2.21	0.43
1:D:116:LEU:O	1:D:120:ILE:HG13	2.19	0.43
1:D:421:ARG:HD3	1:D:421:ARG:HA	1.76	0.43
1:E:23:LEU:HD13	1:E:23:LEU:C	2.38	0.43
1:F:103:GLY:O	1:F:107:VAL:HG23	2.19	0.43
1:H:207:LYS:HE3	1:H:207:LYS:HB2	1.82	0.43
1:H:226:LYS:HG2	1:H:252:GLU:CB	2.49	0.43
1:J:488:MET:CE	1:J:493:ILE:HG21	2.49	0.43
1:L:100:ILE:O	1:L:104:LEU:HG	2.18	0.43
1:L:30:THR:HB	1:L:51:LYS:HG3	2.01	0.43
1:L:358:SER:HA	1:L:362:ARG:CZ	2.49	0.43
1:L:176:THR:OG1	1:L:378:VAL:HG22	2.19	0.43
1:M:197:ARG:HD3	1:M:197:ARG:HA	1.86	0.43
1:M:339:GLU:HA	1:M:342:ILE:HB	2.01	0.43
1:N:215:LEU:O	1:N:218:PRO:HD3	2.19	0.43
1:N:401:HIS:O	1:N:404:ARG:HB2	2.19	0.43
2:R:84:LEU:O	2:R:84:LEU:HD12	2.19	0.43
1:E:231:ARG:NH1	2:S:31:ALA:O	2.49	0.43
2:S:46:GLY:HA2	2:S:57:LEU:HD11	2.00	0.43
2:X:94:ILE:HD11	2:Y:4:ARG:CG	2.49	0.43
2:Z:8:ASP:O	2:Z:57:LEU:CD2	2.34	0.43
2:1:94:ILE:O	2:2:3:ILE:HB	2.18	0.43
1:A:199:TYR:HD2	1:A:325:ILE:HG22	1.83	0.43
1:A:247:LEU:O	1:A:273:VAL:HG13	2.19	0.43
1:A:268:ARG:N	1:A:269:GLY:CA	2.81	0.43
1:B:488:MET:O	1:B:493:ILE:HG13	2.19	0.43
1:B:405:ALA:HB1	1:B:498:LYS:HD3	2.01	0.43
1:C:14:VAL:O	1:C:18:ARG:HG3	2.19	0.43
1:D:140:ASP:OD2	1:D:142:LYS:HB3	2.18	0.43

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:D:225:LYS:HG3	1:D:226:LYS:N	2.33	0.43
1:F:359:ASP:O	1:F:360:TYR:HB3	2.19	0.43
1:G:195:PHE:CD1	1:G:195:PHE:C	2.93	0.43
1:G:234:LEU:N	1:G:235:PRO:HD2	2.34	0.43
1:H:232:GLU:O	1:H:234:LEU:HD12	2.18	0.43
1:J:23:LEU:HD13	1:J:23:LEU:C	2.39	0.43
1:J:271:VAL:O	1:J:273:VAL:HG23	2.18	0.43
1:K:349:ILE:HD12	1:K:352:GLN:CB	2.36	0.43
1:M:448:GLU:OE1	1:M:452:ARG:NH1	2.52	0.43
1:N:461:GLU:HA	1:N:462:PRO:HD3	1.77	0.43
2:P:3:ILE:N	2:P:3:ILE:HD12	2.29	0.43
2:Q:6:LEU:HD13	2:Q:7:HIS:CD2	2.41	0.43
2:S:97:ALA:O	2:T:1:MET:HA	2.18	0.43
2:U:84:LEU:HD12	2:U:84:LEU:N	2.34	0.43
1:A:295:LEU:C	1:A:295:LEU:HD23	2.40	0.42
1:A:310:GLU:HG2	1:A:311:LYS:HG3	2.01	0.42
1:B:149:THR:HG23	1:B:155:ASP:C	2.39	0.42
1:B:347:ALA:O	1:B:350:ARG:HB2	2.19	0.42
1:C:187:LEU:HD13	1:C:379:ILE:HG12	2.01	0.42
1:D:248:LEU:HD13	1:D:249:ILE:N	2.33	0.42
1:F:105:LYS:HD3	1:F:105:LYS:HA	1.66	0.42
1:F:228:SER:O	1:F:258:ALA:HB2	2.19	0.42
1:G:239:ALA:CB	1:G:314:LEU:HD13	2.45	0.42
1:H:254:VAL:HG12	1:H:259:LEU:HG	2.00	0.42
1:H:210:THR:HG22	1:H:326:ASN:HD21	1.84	0.42
1:I:117:LYS:HG3	1:I:512:GLY:C	2.39	0.42
1:I:228:SER:HB3	1:I:255:GLU:CG	2.48	0.42
1:I:455:VAL:HG21	1:I:465:VAL:HG11	2.01	0.42
1:K:199:TYR:OH	1:K:327:LYS:HG3	2.19	0.42
1:L:265:ASN:OD1	1:L:270:ILE:HD12	2.20	0.42
1:M:165:ALA:HB1	1:M:175:ILE:HD13	2.00	0.42
1:M:231:ARG:HH11	2:1:29:GLY:HA3	1.84	0.42
1:N:346:VAL:CG1	1:N:350:ARG:HH12	2.32	0.42
2:Q:40:VAL:HG13	2:Q:41:LEU:N	2.34	0.42
2:S:49:LEU:H	2:S:52:GLY:HA3	1.83	0.42
2:V:4:ARG:HB3	2:2:94:ILE:HG13	2.00	0.42
1:B:197:ARG:HH11	1:B:198:GLY:H	1.67	0.42
1:B:314:LEU:N	1:B:314:LEU:HD12	2.34	0.42
1:D:207:LYS:N	1:D:208:PRO:CD	2.82	0.42
1:E:234:LEU:CD1	1:E:238:GLU:OE1	2.67	0.42
1:E:225:LYS:HD2	1:E:303:GLU:OE1	2.19	0.42

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:F:433:ASN:OD1	1:F:436:GLN:HG3	2.19	0.42
1:G:209:GLU:HG2	1:G:210:THR:N	2.25	0.42
1:G:217:SER:HA	1:G:320:ALA:O	2.19	0.42
1:I:162:ILE:HG21	1:I:403:THR:HG21	2.01	0.42
1:J:106:ALA:O	1:J:109:ALA:HB3	2.19	0.42
1:J:417:VAL:HG11	1:J:488:MET:HG3	2.01	0.42
1:J:77:VAL:HB	1:J:510:VAL:HG21	1.99	0.42
1:K:136:VAL:HG13	1:K:137:PRO:HD2	2.01	0.42
1:K:347:ALA:O	1:K:351:GLN:NE2	2.39	0.42
1:L:357:THR:HG21	1:L:361:ASP:CB	2.40	0.42
1:M:488:MET:HE3	1:M:493:ILE:HG21	2.01	0.42
1:N:18:ARG:HB3	1:N:18:ARG:NH1	2.35	0.42
2:R:19:THR:HG21	2:R:33:ALA:CB	2.42	0.42
2:W:17:VAL:C	2:W:19:THR:H	2.21	0.42
2:Z:65:VAL:HA	2:Z:94:ILE:HA	2.02	0.42
1:A:325:ILE:N	1:A:325:ILE:HD12	2.34	0.42
1:E:238:GLU:O	1:E:242:LYS:HG3	2.18	0.42
1:E:416:GLY:O	1:E:420:ILE:HG23	2.18	0.42
1:E:412:VAL:HG13	1:E:497:THR:OG1	2.19	0.42
1:G:247:LEU:HD22	1:G:247:LEU:N	2.34	0.42
1:G:475:ASN:HB3	1:G:489:ILE:HG12	2.01	0.42
1:H:349:ILE:O	1:H:353:ILE:N	2.51	0.42
1:H:90:THR:O	1:H:94:VAL:HG23	2.19	0.42
1:K:228:SER:OG	1:K:255:GLU:HB2	2.19	0.42
1:M:357:THR:HB	1:M:361:ASP:CG	2.39	0.42
1:N:319:GLN:OE1	1:N:319:GLN:N	2.53	0.42
1:N:326:ASN:OD1	1:N:328:ASP:HB2	2.19	0.42
2:R:8:ASP:HA	2:R:57:LEU:HD21	2.01	0.42
2:S:14:ARG:CZ	2:S:84:LEU:HD21	2.49	0.42
2:S:36:THR:HG23	2:S:37:ARG:CG	2.44	0.42
2:Z:57:LEU:CD1	2:Z:57:LEU:H	2.29	0.42
2:2:54:VAL:HG12	2:2:56:PRO:HD3	2.01	0.42
1:A:18:ARG:HH11	1:A:18:ARG:HB3	1.83	0.42
1:B:221:LEU:HD22	1:B:249:ILE:HG23	2.01	0.42
1:B:461:GLU:OE1	1:N:463:SER:OG	2.24	0.42
1:C:253:ASP:OD1	1:C:277:LYS:CE	2.64	0.42
1:D:219:PHE:CD2	1:D:240:VAL:HG22	2.54	0.42
1:K:367:GLU:O	1:K:371:LYS:HG2	2.20	0.42
1:L:77:VAL:HG21	1:L:507:ALA:CA	2.49	0.42
1:M:193:MET:SD	1:M:371:LYS:HB3	2.59	0.42
2:Q:92:LEU:HG	2:Q:93:ALA:H	1.85	0.42

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:R:36:THR:C	2:R:37:ARG:HG3	2.40	0.42
2:V:40:VAL:HG13	2:V:62:GLY:H	1.84	0.42
2:X:48:ILE:N	2:X:48:ILE:HD12	2.34	0.42
2:Z:12:VAL:HG23	2:Z:39:GLU:O	2.19	0.42
2:Z:74:LYS:HG3	2:Z:76:GLU:HG3	2.01	0.42
1:A:177:VAL:HG21	1:A:397:GLU:CG	2.39	0.42
1:A:421:ARG:HA	1:A:421:ARG:HD3	1.77	0.42
1:B:421:ARG:HA	1:B:421:ARG:HD3	1.85	0.42
1:C:175:ILE:N	1:C:175:ILE:HD12	2.34	0.42
1:F:127:ALA:HB2	1:F:426:LEU:HD12	2.01	0.42
1:F:23:LEU:HD23	1:F:60:ILE:CB	2.48	0.42
1:F:240:VAL:HG11	1:F:247:LEU:CD1	2.49	0.42
1:F:266:THR:HG22	1:F:273:VAL:H	1.84	0.42
1:G:207:LYS:O	1:G:209:GLU:N	2.42	0.42
1:G:276:VAL:HG22	1:G:277:LYS:N	2.34	0.42
1:H:278:ALA:HB3	1:H:285:ARG:CD	2.36	0.42
1:K:105:LYS:HA	1:K:105:LYS:HD3	1.72	0.42
1:M:23:LEU:O	1:M:27:VAL:HG23	2.19	0.42
1:N:20:VAL:HG11	1:N:100:ILE:HD12	2.00	0.42
1:N:168:LYS:CD	1:N:189:VAL:HG21	2.49	0.42
1:N:151:SER:CB	1:N:399:ALA:HA	2.50	0.42
2:V:5:PRO:HG2	2:V:43:VAL:C	2.40	0.42
2:V:86:MET:SD	2:V:91:ILE:HD11	2.59	0.42
2:Y:86:MET:CE	2:Y:90:ASP:HB2	2.50	0.42
2:1:49:LEU:C	2:1:49:LEU:HD12	2.40	0.42
2:2:49:LEU:CB	2:2:52:GLY:HA2	2.49	0.42
1:A:339:GLU:OE1	1:A:339:GLU:N	2.53	0.42
1:D:305:ILE:HG23	1:E:263:VAL:HG11	2.01	0.42
1:F:18:ARG:HB3	1:F:18:ARG:HH11	1.80	0.42
1:F:127:ALA:HB2	1:F:426:LEU:CD1	2.49	0.42
1:H:475:ASN:O	1:H:488:MET:HG2	2.19	0.42
1:H:478:TYR:HD1	1:H:485:TYR:CD1	2.37	0.42
1:I:197:ARG:HH12	1:I:277:LYS:HD2	1.78	0.42
1:I:302:SER:OG	1:I:305:ILE:HB	2.20	0.42
1:J:175:ILE:HG23	1:J:377:ALA:O	2.19	0.42
1:L:232:GLU:HG3	1:L:310:GLU:OE2	2.19	0.42
1:N:263:VAL:O	1:N:267:MET:HG2	2.18	0.42
1:E:238:GLU:CD	2:S:25:ILE:HD13	2.39	0.42
2:T:53:GLU:CD	2:T:54:VAL:HG22	2.40	0.42
2:U:54:VAL:O	2:U:54:VAL:HG23	2.19	0.42
2:U:65:VAL:HB	2:U:94:ILE:HG22	2.01	0.42

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:Y:11:ILE:HD11	2:Y:42:ALA:HB3	2.01	0.42
1:A:230:ILE:HD12	1:A:230:ILE:H	1.85	0.42
1:A:20:VAL:HG13	1:A:74:VAL:HG11	2.01	0.42
1:B:266:THR:CG2	1:B:273:VAL:HB	2.50	0.42
1:B:290:GLN:HG3	1:B:345:ARG:NH2	2.34	0.42
1:C:234:LEU:N	1:C:235:PRO:CD	2.81	0.42
1:C:151:SER:HB3	1:C:399:ALA:HA	2.01	0.42
1:D:282:GLY:HA2	1:D:285:ARG:NH2	2.35	0.42
1:D:311:LYS:HE2	1:D:311:LYS:HB3	1.75	0.42
1:D:33:PRO:HG3	3:D:601:ADP:C6	2.54	0.42
1:E:246:PRO:HA	1:E:272:LYS:O	2.18	0.42
1:F:434:GLU:O	1:F:437:ASN:HB2	2.20	0.42
1:H:164:GLU:O	1:H:168:LYS:HG3	2.20	0.42
1:I:402:ALA:HB1	1:I:496:PRO:HG3	2.01	0.42
1:K:513:LEU:O	1:K:516:THR:OG1	2.24	0.42
1:L:157:THR:O	1:L:161:LEU:HB2	2.19	0.42
1:L:327:LYS:HG3	1:L:328:ASP:OD1	2.20	0.42
1:L:8:PHE:CE2	1:M:26:ALA:HA	2.55	0.42
1:M:310:GLU:OE1	1:M:310:GLU:N	2.52	0.42
1:M:363:GLU:HG3	1:M:364:LYS:N	2.34	0.42
1:N:150:ILE:CD1	1:N:493:ILE:HA	2.49	0.42
2:O:20:LYS:HB3	2:O:27:LEU:HD12	2.01	0.42
2:O:95:VAL:CB	2:P:3:ILE:HG21	2.49	0.42
2:P:60:LYS:HG2	2:P:61:VAL:H	1.84	0.42
2:R:12:VAL:HG12	2:R:40:VAL:CB	2.49	0.42
2:R:43:VAL:HG23	2:R:57:LEU:HD12	2.01	0.42
1:F:265:ASN:OD1	2:T:26:VAL:HG23	2.19	0.42
2:U:6:LEU:O	2:U:9:ARG:HG2	2.20	0.42
2:Y:20:LYS:HB2	2:Y:28:THR:CG2	2.47	0.42
2:Y:77:LYS:HE2	2:Y:82:GLU:OE1	2.20	0.42
1:C:226:LYS:HG2	1:C:252:GLU:CB	2.50	0.42
1:C:258:ALA:O	1:C:262:LEU:HB2	2.20	0.42
1:D:138:CYS:HB2	1:D:411:VAL:CG1	2.48	0.42
1:D:218:PRO:CA	1:D:246:PRO:HG2	2.47	0.42
1:E:365:LEU:HD23	1:E:369:VAL:CG2	2.48	0.42
1:D:519:CYS:HB3	1:E:38:VAL:HG22	2.01	0.42
1:G:175:ILE:HA	1:G:377:ALA:O	2.19	0.42
1:H:136:VAL:HG23	1:H:137:PRO:CD	2.49	0.42
1:I:142:LYS:HE2	1:I:146:GLN:HE22	1.81	0.42
1:K:17:LEU:CD1	1:K:104:LEU:HD12	2.49	0.42
1:K:38:VAL:HB	1:K:56:VAL:HG21	2.01	0.42

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:M:196:ASP:N	1:M:196:ASP:OD1	2.52	0.42
1:M:288:MET:HG3	1:M:364:LYS:HZ3	1.85	0.42
1:N:362:ARG:O	1:N:366:GLN:HB2	2.20	0.42
2:P:27:LEU:N	2:P:27:LEU:HD12	2.34	0.42
2:Q:49:LEU:C	2:Q:49:LEU:HD12	2.40	0.42
2:X:93:ALA:HB1	2:Y:5:PRO:HA	2.00	0.42
2:1:5:PRO:HB3	2:1:11:ILE:HD11	2.01	0.42
2:V:6:LEU:CD2	2:2:91:ILE:HG22	2.43	0.42
1:A:219:PHE:HA	1:A:318:GLY:O	2.19	0.42
1:B:231:ARG:CD	1:B:258:ALA:HA	2.48	0.42
1:C:288:MET:O	1:C:292:ILE:HG13	2.18	0.42
1:C:357:THR:HG22	1:C:361:ASP:HB2	2.00	0.42
1:D:147:VAL:HG21	1:D:403:THR:HA	2.01	0.42
1:E:238:GLU:O	1:E:242:LYS:N	2.47	0.42
1:E:349:ILE:CG2	1:E:365:LEU:HD21	2.50	0.42
1:F:174:VAL:CG1	1:F:367:GLU:HA	2.49	0.42
1:G:219:PHE:HE2	1:G:245:LYS:CB	2.30	0.42
1:G:383:ALA:HB3	1:G:389:MET:HA	2.02	0.42
1:G:496:PRO:HG2	1:G:499:VAL:CG2	2.50	0.42
1:H:37:ASN:OD1	1:H:51:LYS:HB2	2.20	0.42
1:I:143:ALA:O	1:I:147:VAL:HG13	2.20	0.42
1:J:488:MET:HE3	1:J:493:ILE:HG21	2.01	0.42
1:L:199:TYR:CE1	1:L:202:PRO:HA	2.54	0.42
1:L:421:ARG:HD3	1:L:421:ARG:HA	1.93	0.42
1:L:42:LYS:HE2	1:L:48:THR:HB	2.01	0.42
1:L:513:LEU:CD1	1:M:388:GLU:HA	2.50	0.42
1:M:228:SER:HB3	1:M:255:GLU:CB	2.50	0.42
1:N:496:PRO:HG2	1:N:499:VAL:HG13	2.00	0.42
2:O:5:PRO:HA	2:U:93:ALA:CA	2.49	0.42
2:S:78:ILE:CG2	2:S:83:VAL:HG21	2.50	0.42
2:Z:24:GLY:H	2:1:80:ASN:ND2	2.18	0.42
2:2:55:LYS:O	2:2:55:LYS:HG3	2.19	0.42
1:A:245:LYS:HB3	1:A:246:PRO:HD3	1.85	0.42
1:A:252:GLU:O	1:A:285:ARG:NH1	2.53	0.42
1:B:234:LEU:HD23	1:B:235:PRO:CD	2.47	0.42
1:B:414:GLY:O	1:B:417:VAL:HG12	2.19	0.42
1:C:419:LEU:HD22	1:C:419:LEU:H	1.84	0.42
1:C:433:ASN:OD1	1:C:435:ASP:HB2	2.20	0.42
1:F:210:THR:HG22	1:F:210:THR:O	2.19	0.42
1:F:270:ILE:HD13	2:T:25:ILE:O	2.20	0.42
1:H:183:LEU:CD1	1:H:183:LEU:H	2.32	0.42

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:H:322:ARG:HG2	1:H:323:VAL:N	2.35	0.42
1:H:49:ILE:HG21	1:N:513:LEU:HD22	2.02	0.42
1:I:236:VAL:O	1:I:240:VAL:HG23	2.20	0.42
1:J:115:ASP:O	1:J:436:GLN:HG2	2.20	0.42
1:J:295:LEU:C	1:J:295:LEU:HD23	2.41	0.42
1:K:343:GLN:HA	1:K:346:VAL:CG2	2.50	0.42
1:N:131:LEU:HD23	1:N:131:LEU:HA	1.93	0.42
1:N:40:LEU:HD21	1:N:56:VAL:CG2	2.42	0.42
2:O:17:VAL:CG2	2:O:18:GLU:HG3	2.50	0.42
2:R:18:GLU:O	2:R:19:THR:OG1	2.33	0.42
2:Y:29:GLY:HA2	2:Y:30:SER:HA	1.74	0.42
2:1:16:GLU:HB3	2:1:17:VAL:H	1.54	0.41
2:1:95:VAL:HG13	2:2:3:ILE:CG2	2.46	0.41
1:A:241:ALA:CB	1:A:271:VAL:CG1	2.97	0.41
1:B:510:VAL:HG12	1:C:387:VAL:CG2	2.50	0.41
1:C:322:ARG:HG2	1:C:323:VAL:N	2.35	0.41
1:E:124:VAL:HG22	1:E:504:LEU:HD11	2.02	0.41
1:E:206:ASN:OD1	1:E:207:LYS:N	2.43	0.41
1:E:295:LEU:HD23	1:E:295:LEU:O	2.20	0.41
1:F:322:ARG:HB3	1:F:333:ILE:CG2	2.50	0.41
1:G:237:LEU:C	1:G:237:LEU:HD23	2.41	0.41
1:H:219:PHE:O	1:H:247:LEU:HG	2.20	0.41
1:H:271:VAL:O	1:H:273:VAL:HG23	2.20	0.41
1:H:345:ARG:O	1:H:349:ILE:HG13	2.20	0.41
1:H:8:PHE:O	1:H:11:ASP:HB3	2.19	0.41
1:I:194:GLN:HG3	1:I:330:THR:O	2.20	0.41
1:I:419:LEU:O	1:I:422:VAL:HG22	2.20	0.41
1:J:230:ILE:O	1:J:231:ARG:CG	2.68	0.41
1:J:314:LEU:CD2	1:J:314:LEU:O	2.60	0.41
1:K:295:LEU:HD23	1:K:295:LEU:O	2.19	0.41
1:L:313:THR:HG22	1:L:314:LEU:N	2.35	0.41
1:M:303:GLU:HG2	1:M:303:GLU:O	2.20	0.41
1:N:225:LYS:HG3	1:N:226:LYS:H	1.83	0.41
2:S:16:GLU:CG	2:S:19:THR:CG2	2.98	0.41
2:S:20:LYS:NZ	2:S:26:VAL:HG22	2.34	0.41
2:S:73:VAL:CG2	2:S:84:LEU:HB3	2.49	0.41
2:O:9:ARG:HD3	2:U:92:LEU:HD22	2.02	0.41
2:V:13:LYS:HB2	2:V:41:LEU:HD11	2.02	0.41
2:W:36:THR:C	2:W:37:ARG:HG3	2.39	0.41
2:W:9:ARG:CG	2:W:9:ARG:NH1	2.73	0.41
2:W:37:ARG:CZ	2:X:78:ILE:HD11	2.50	0.41

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:Z:68:ASN:HD21	2:1:74:LYS:HE2	1.86	0.41
2:1:76:GLU:HG3	2:1:77:LYS:N	2.35	0.41
2:1:57:LEU:HD12	2:1:88:GLU:CD	2.40	0.41
1:A:147:VAL:HG12	1:A:494:LEU:HD12	2.02	0.41
1:C:323:VAL:CG1	1:C:332:ILE:HG12	2.48	0.41
1:D:432:GLN:HB2	1:D:436:GLN:OE1	2.19	0.41
1:E:195:PHE:CZ	1:E:330:THR:HB	2.55	0.41
1:F:151:SER:CB	1:F:399:ALA:HA	2.49	0.41
1:G:234:LEU:CD1	1:G:234:LEU:H	2.32	0.41
1:H:288:MET:HA	1:H:291:ASP:OD2	2.21	0.41
1:H:174:VAL:CG2	1:H:376:VAL:HG13	2.42	0.41
1:I:114:MET:HG3	1:I:118:ARG:NH1	2.35	0.41
1:I:178:GLU:OE1	1:I:378:VAL:HG11	2.20	0.41
1:I:230:ILE:O	1:I:232:GLU:N	2.52	0.41
1:J:209:GLU:C	1:J:211:GLY:N	2.73	0.41
1:J:413:ALA:O	1:J:418:ALA:HB2	2.20	0.41
1:K:90:THR:HG23	4:K:602:BEF:F1	2.10	0.41
1:L:224:ASP:OD2	1:L:286:LYS:HE2	2.19	0.41
1:L:506:TYR:O	1:L:510:VAL:HG22	2.20	0.41
1:M:190:VAL:HG11	1:M:194:GLN:NE2	2.35	0.41
1:M:333:ILE:HG22	1:M:333:ILE:O	2.19	0.41
1:N:239:ALA:HB1	1:N:314:LEU:CD1	2.46	0.41
2:W:24:GLY:O	2:W:25:ILE:HG22	2.20	0.41
2:1:55:LYS:HA	2:1:56:PRO:HD2	1.88	0.41
2:1:95:VAL:CG1	2:2:3:ILE:HG22	2.48	0.41
1:B:295:LEU:HD23	1:B:335:GLY:HA3	2.02	0.41
1:C:33:PRO:CG	1:C:480:ALA:HB3	2.50	0.41
1:C:33:PRO:HG3	3:C:601:ADP:C6	2.55	0.41
1:D:281:PHE:O	1:D:284:ARG:HB2	2.20	0.41
1:E:232:GLU:C	1:E:234:LEU:N	2.74	0.41
1:E:234:LEU:C	1:E:236:VAL:N	2.72	0.41
1:F:143:ALA:O	1:F:146:GLN:HB2	2.20	0.41
1:G:194:GLN:NE2	1:G:329:THR:HG21	2.34	0.41
1:G:223:ALA:CB	1:G:224:ASP:CA	2.98	0.41
1:A:386:GLU:HB3	1:G:76:GLU:OE2	2.20	0.41
1:H:225:LYS:HA	1:H:225:LYS:HD3	1.62	0.41
1:H:265:ASN:OD1	2:V:26:VAL:HG13	2.21	0.41
1:I:219:PHE:CD1	1:I:319:GLN:HG2	2.55	0.41
1:J:237:LEU:C	1:J:237:LEU:HD23	2.41	0.41
1:J:288:MET:O	1:J:288:MET:HG3	2.20	0.41
1:J:221:LEU:HD13	1:J:317: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:K:242:LYS:O	1:K:242:LYS:HD3	2.20	0.41
1:L:225:LYS:HD2	1:L:226:LYS:H	1.83	0.41
1:L:227:ILE:H	1:L:227:ILE:HD12	1.84	0.41
1:L:237:LEU:HD23	1:L:237:LEU:C	2.40	0.41
1:K:305:ILE:CG2	1:L:264:VAL:HG22	2.47	0.41
1:L:31:LEU:CD2	1:L:453:GLN:HB3	2.44	0.41
1:N:162:ILE:O	1:N:166:MET:N	2.49	0.41
1:N:7:LYS:HG3	1:N:66:PHE:CE2	2.55	0.41
2:O:96:GLU:OE1	2:P:1:MET:N	2.48	0.41
2:Z:59:VAL:HG21	2:Z:91:ILE:HG21	2.02	0.41
1:A:160:LYS:HD2	1:A:160:LYS:HA	1.89	0.41
1:A:181:THR:HG21	1:A:380:LYS:CD	2.41	0.41
1:F:219:PHE:O	1:F:248:LEU:HB3	2.20	0.41
1:F:7:LYS:HD2	1:F:66:PHE:CE2	2.55	0.41
1:G:266:THR:HG22	1:G:273:VAL:H	1.85	0.41
1:G:421:ARG:HD3	1:G:421:ARG:HA	1.90	0.41
1:I:174:VAL:HG21	1:I:367:GLU:CA	2.48	0.41
1:I:364:LYS:HB3	1:I:364:LYS:HE2	1.90	0.41
1:L:136:VAL:CG2	1:L:137:PRO:CD	2.98	0.41
1:L:155:ASP:CG	1:L:158:VAL:HG23	2.40	0.41
1:L:216:GLU:OE1	1:L:245:LYS:HD2	2.21	0.41
1:M:103:GLY:O	1:M:106:ALA:HB3	2.19	0.41
1:M:140:ASP:OD1	1:M:143:ALA:N	2.49	0.41
1:M:279:PRO:HB2	1:M:288:MET:CE	2.50	0.41
1:N:72:GLN:HB3	1:N:72:GLN:HE21	1.63	0.41
2:U:70:GLY:O	2:U:73:VAL:HG22	2.21	0.41
2:W:79:ASP:O	2:W:80:ASN:OD1	2.38	0.41
2:Y:36:THR:HG22	2:Y:36:THR:O	2.20	0.41
1:A:124:VAL:HG22	1:A:504:LEU:HD11	2.03	0.41
1:B:149:THR:HG23	1:B:155:ASP:O	2.20	0.41
1:B:183:LEU:CD2	1:B:384:ALA:HA	2.50	0.41
1:B:210:THR:O	1:B:210:THR:HG22	2.20	0.41
1:E:135:SER:HB2	1:E:497:THR:HG21	2.03	0.41
1:F:349:ILE:HG21	1:F:369:VAL:CG2	2.49	0.41
1:G:18:ARG:HB3	1:G:18:ARG:HH11	1.86	0.41
1:G:314:LEU:HA	1:G:314:LEU:HD12	1.95	0.41
1:G:487:ASN:HB3	1:G:490:ASP:HB2	2.03	0.41
1:A:49:ILE:HD11	1:G:73:MET:SD	2.60	0.41
1:J:151:SER:CB	1:J:399:ALA:HA	2.48	0.41
1:L:222:LEU:N	1:L:222:LEU:HD12	2.35	0.41
1:M:158:VAL:O	1:M:162:ILE:HG13	2.20	0.41

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:N:278:ALA:HB1	1:N:279:PRO:CD	2.50	0.41
2:R:14:ARG:HB3	2:R:84:LEU:HD21	2.03	0.41
2:R:14:ARG:HE	2:R:84:LEU:CD2	2.34	0.41
2:S:48:ILE:HG13	2:S:53:GLU:N	2.35	0.41
2:S:94:ILE:HD13	2:T:4:ARG:HH21	1.85	0.41
2:T:43:VAL:HG13	2:T:57:LEU:HD12	2.02	0.41
2:T:76:GLU:O	2:T:82:GLU:HB2	2.21	0.41
2:W:25:ILE:O	2:W:25:ILE:HG23	2.20	0.41
1:A:231:ARG:HB3	1:A:258:ALA:CB	2.51	0.41
1:A:307:MET:O	1:A:308:GLU:HG3	2.20	0.41
1:A:87:ASP:OD1	4:A:602:BEF:F2	2.29	0.41
1:B:250:ILE:HG23	1:B:278:ALA:HA	2.03	0.41
1:B:325:ILE:HG23	1:B:330:THR:OG1	2.19	0.41
1:C:399:ALA:O	1:C:403:THR:HG23	2.20	0.41
1:C:475:ASN:O	1:C:488:MET:HG2	2.21	0.41
1:D:313:THR:HG22	1:D:315:GLU:H	1.85	0.41
1:E:222:LEU:CD2	1:E:289:LEU:HD13	2.49	0.41
1:F:225:LYS:HB2	1:F:303:GLU:OE1	2.20	0.41
1:F:37:ASN:OD1	1:F:51:LYS:HB2	2.20	0.41
1:F:415:GLY:N	3:F:601:ADP:O2'	2.51	0.41
1:G:145:ALA:O	1:G:149:THR:HG23	2.21	0.41
1:G:278:ALA:HA	1:G:279:PRO:HD3	1.86	0.41
1:G:222:LEU:HD23	1:G:289:LEU:HB3	2.03	0.41
1:G:305:ILE:CG2	1:G:306:GLY:N	2.83	0.41
1:H:218:PRO:HA	1:H:246:PRO:HG2	2.03	0.41
1:H:222:LEU:HD21	1:H:292:ILE:CG2	2.51	0.41
1:I:305:ILE:CG2	1:I:306:GLY:N	2.76	0.41
1:L:142:LYS:O	1:L:146:GLN:HG3	2.21	0.41
1:M:233:MET:HE2	1:M:237:LEU:CD2	2.51	0.41
1:M:221:LEU:HB2	1:M:317:LEU:HD22	2.02	0.41
1:N:33:PRO:HA	1:N:153:ASN:OD1	2.21	0.41
1:N:160:LYS:HA	1:N:160:LYS:HD2	1.89	0.41
1:N:23:LEU:HD23	1:N:60:ILE:HB	2.01	0.41
1:N:285:ARG:O	1:N:289:LEU:HG	2.21	0.41
2:Q:3:ILE:HD11	2:Q:78:ILE:CD1	2.50	0.41
2:Q:88:GLU:O	2:Q:88:GLU:HG2	2.21	0.41
2:T:64:ILE:HB	2:T:95:VAL:HG13	2.02	0.41
2:W:65:VAL:HG12	2:W:94:ILE:HG12	2.03	0.41
1:A:363:GLU:O	1:A:367:GLU:HB2	2.21	0.41
1:C:210:THR:HG22	1:C:210:THR:O	2.21	0.41
1:D:305:ILE:CG2	1:D:306:GLY:N	2.74	0.41

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:D:52:ASP:O	1:D:55:SER:HB2	2.20	0.41
1:E:207:LYS:N	1:E:208:PRO:HD2	2.35	0.41
1:F:333:ILE:O	1:F:334:ASP:HB2	2.21	0.41
1:G:420:ILE:HD11	1:G:470:LYS:HG3	2.03	0.41
1:H:23:LEU:HD23	1:H:60:ILE:HB	2.02	0.41
1:I:340:ALA:O	1:I:344:GLY:N	2.54	0.41
1:J:291:ASP:HB3	1:J:372:LEU:CD2	2.50	0.41
1:J:347:ALA:HA	1:J:350:ARG:CG	2.51	0.41
1:J:419:LEU:N	1:J:419:LEU:HD12	2.34	0.41
1:M:183:LEU:HD22	1:M:384:ALA:HA	2.03	0.41
1:N:228:SER:O	1:N:258:ALA:HB2	2.21	0.41
1:N:355:GLU:HG3	1:N:356:ALA:N	2.32	0.41
1:N:284:ARG:NH2	1:N:364:LYS:HG3	2.35	0.41
2:T:5:PRO:HD3	2:T:11:ILE:CD1	2.51	0.41
2:Z:17:VAL:O	2:Z:18:GLU:HG2	2.21	0.41
1:N:261:THR:CB	2:2:29:GLY:H	2.24	0.41
1:A:278:ALA:HB3	1:A:285:ARG:CZ	2.50	0.41
1:A:326:ASN:ND2	1:A:329:THR:O	2.54	0.41
1:B:264:VAL:O	1:B:267:MET:HB2	2.21	0.41
1:B:409:GLU:CD	1:B:501:ARG:HH21	2.24	0.41
1:D:197:ARG:HA	1:D:197:ARG:HD3	1.88	0.41
1:D:259:LEU:O	1:D:263:VAL:HG23	2.21	0.41
1:D:322:ARG:HD2	1:D:333:ILE:HD12	2.03	0.41
1:D:77:VAL:HG21	1:D:507:ALA:CA	2.49	0.41
1:F:200:LEU:N	1:F:200:LEU:HD12	2.36	0.41
1:G:350:ARG:HD3	1:G:353:ILE:HD11	2.01	0.41
1:G:434:GLU:HA	1:G:437:ASN:HB2	2.02	0.41
1:H:144:ILE:O	1:H:147:VAL:HG22	2.21	0.41
1:H:197:ARG:HA	1:H:197:ARG:HD3	1.94	0.41
1:H:467:ASN:O	1:H:471:GLY:N	2.51	0.41
3:I:601:ADP:O2B	4:I:602:BEF:F2	2.28	0.41
1:J:205:ILE:HA	1:J:213:VAL:CG2	2.25	0.41
1:J:301:ILE:HA	1:J:307:MET:HE1	2.02	0.41
1:K:183:LEU:HD21	1:K:384:ALA:HA	2.03	0.41
1:K:66:PHE:CE1	1:K:522:THR:HG22	2.55	0.41
1:L:196:ASP:OD1	1:L:196:ASP:N	2.53	0.41
1:L:219:PHE:CD1	1:L:240:VAL:HG22	2.56	0.41
1:M:306:GLY:HA3	1:N:264:VAL:HG21	2.03	0.41
1:N:230:ILE:HD11	1:N:231:ARG:HG2	2.03	0.41
1:N:234:LEU:CD1	1:N:234:LEU:H	2.22	0.41
2:S:12:VAL:HG22	2:S:40:VAL:HA	2.03	0.41

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:S:65:VAL:HG21	2:S:91:ILE:HD12	2.02	0.41
2:U:17:VAL:HG23	2:U:33:ALA:O	2.20	0.41
2:O:3:ILE:N	2:U:95:VAL:HA	2.36	0.41
2:W:37:ARG:NH2	2:X:78:ILE:HD11	2.35	0.41
2:Y:13:LYS:HB2	2:Y:41:LEU:HD11	2.02	0.41
2:2:51:ASN:O	2:2:51:ASN:OD1	2.39	0.41
1:B:197:ARG:HG2	1:B:277:LYS:O	2.20	0.41
1:B:343:GLN:HA	1:B:346:VAL:HB	2.02	0.41
1:C:183:LEU:N	1:C:183:LEU:HD12	2.36	0.41
1:C:254:VAL:CG1	1:C:259:LEU:HG	2.44	0.41
1:D:229:ASN:CG	1:D:230:ILE:H	2.12	0.41
1:D:248:LEU:CD1	1:D:250:ILE:HG13	2.51	0.41
1:D:419:LEU:HB3	1:D:447:MET:HB3	2.03	0.41
1:G:221:LEU:HD22	1:G:249:ILE:HD12	2.02	0.41
1:G:279:PRO:HB2	1:G:288:MET:HE3	2.02	0.41
1:H:409:GLU:CD	1:H:501:ARG:HH21	2.24	0.41
1:I:295:LEU:C	1:I:295:LEU:HD23	2.41	0.41
1:J:349:ILE:HG22	1:J:365:LEU:HD21	2.01	0.41
1:J:171:LYS:HD3	1:J:407:VAL:HG11	2.02	0.41
1:K:123:ALA:HB2	1:K:440:ILE:HG23	2.02	0.41
1:K:215:LEU:HD12	1:K:323:VAL:CG2	2.50	0.41
1:K:296:THR:CB	1:K:318:GLY:HA3	2.51	0.41
1:K:455:VAL:CG1	1:K:460:GLU:HB2	2.40	0.41
1:L:350:ARG:HA	1:L:353:ILE:CD1	2.51	0.41
1:L:349:ILE:O	1:L:353:ILE:N	2.53	0.41
1:M:183:LEU:CD2	1:M:384:ALA:HA	2.50	0.41
1:M:210:THR:O	1:M:210:THR:HG22	2.20	0.41
1:M:363:GLU:HG3	1:M:364:LYS:H	1.86	0.41
1:M:406:ALA:HB2	1:M:496:PRO:HB3	2.02	0.41
1:M:62:LEU:HD23	1:M:67:GLU:HB3	2.02	0.41
1:N:388:GLU:O	1:N:392:LYS:HB2	2.20	0.41
1:N:61:GLU:O	1:N:62:LEU:HD12	2.21	0.41
2:O:77:LYS:HG2	2:O:82:GLU:HA	2.02	0.41
2:P:68:ASN:HB3	2:Q:74:LYS:NZ	2.36	0.41
2:Q:43:VAL:CG2	2:Q:57:LEU:HD11	2.48	0.41
2:Q:70:GLY:O	2:Q:73:VAL:HG12	2.19	0.41
2:T:5:PRO:HB2	2:T:9:ARG:O	2.21	0.41
2:U:49:LEU:CA	2:U:50:GLU:OE1	2.69	0.41
2:V:86:MET:SD	2:V:91:ILE:HG13	2.61	0.41
2:W:11:ILE:HG21	2:W:42:ALA:HB3	2.02	0.41
2:W:64:ILE:HG21	2:W:95:VAL:HB	2.01	0.41

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:X:58:ASP:OD2	2:Y:7:HIS:CE1	2.73	0.41
2:Y:78:ILE:HG13	2:Y:79:ASP:N	2.36	0.41
2:Z:47:ARG:HE	2:Z:49:LEU:HD12	1.86	0.41
2:Z:73:VAL:HG12	2:Z:86:MET:HB3	2.02	0.41
2:2:17:VAL:HG12	2:2:35:SER:H	1.86	0.41
1:B:219:PHE:CB	1:B:317:LEU:HD13	2.45	0.41
1:C:124:VAL:HG21	1:C:508:ALA:HB2	2.03	0.41
1:C:276:VAL:HG22	1:C:277:LYS:N	2.35	0.41
1:C:414:GLY:O	1:C:417:VAL:HG12	2.21	0.41
1:D:305:ILE:CG2	1:E:263:VAL:HG11	2.50	0.41
1:E:210:THR:O	1:E:210:THR:HG22	2.21	0.41
1:E:327:LYS:HD3	1:E:327:LYS:H	1.86	0.41
1:E:400:LEU:O	1:E:400:LEU:HD13	2.21	0.41
1:E:5:ASP:HB2	1:E:524:LEU:HD21	1.99	0.41
1:G:230:ILE:CG2	1:G:309:LEU:HD21	2.33	0.41
1:G:92:ALA:HB2	1:G:503:ALA:CB	2.50	0.41
1:H:284:ARG:HA	1:H:284:ARG:HD3	1.83	0.41
1:I:218:PRO:HD2	1:I:320:ALA:O	2.21	0.41
1:J:314:LEU:C	1:J:314:LEU:HD23	2.40	0.41
1:K:23:LEU:HD23	1:K:60:ILE:CB	2.44	0.41
1:K:278:ALA:HB1	1:K:279:PRO:HD2	2.02	0.41
1:M:229:ASN:C	1:M:231:ARG:H	2.22	0.41
1:N:215:LEU:HB2	1:N:323:VAL:CG2	2.50	0.41
1:N:325:ILE:HG22	1:N:326:ASN:N	2.35	0.41
2:S:3:ILE:HD12	2:S:3:ILE:C	2.40	0.41
2:V:2:ASN:C	2:V:2:ASN:ND2	2.73	0.41
2:Z:47:ARG:CG	2:Z:49:LEU:HG	2.51	0.41
1:A:321:LYS:HD2	1:A:334:ASP:OD2	2.20	0.41
1:A:27:VAL:CG1	1:A:90:THR:HG23	2.45	0.41
1:B:112:ASN:HA	1:B:113:PRO:HD3	1.85	0.41
1:B:180:GLY:H	1:B:389:MET:CE	2.33	0.41
1:B:222:LEU:HD21	1:B:293:ALA:HB2	2.02	0.41
1:B:151:SER:HB3	1:B:399:ALA:HA	2.03	0.41
1:B:494:LEU:N	1:B:494:LEU:HD12	2.36	0.41
1:C:406:ALA:HB1	1:C:411:VAL:HG12	2.03	0.41
1:D:263:VAL:CG1	1:D:267:MET:HE3	2.51	0.41
1:D:291:ASP:HB3	1:D:372:LEU:HD11	2.03	0.41
1:E:221:LEU:HB2	1:E:317:LEU:CD2	2.51	0.41
1:E:496:PRO:HG2	1:E:499:VAL:CG1	2.50	0.41
1:F:18:ARG:CZ	1:F:18:ARG:HB3	2.51	0.41
1:F:29:VAL:O	1:F:36:ARG:N	2.43	0.41

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:F:452:ARG:HH21	1:F:466:ALA:HB3	1.86	0.41
1:G:307:MET:O	1:G:308:GLU:HG2	2.21	0.41
1:F:513:LEU:CD1	1:G:388:GLU:HA	2.48	0.41
1:I:188:ASP:O	1:I:377:ALA:HA	2.21	0.41
1:J:147:VAL:HG21	1:J:403:THR:HA	2.03	0.41
1:K:150:ILE:HG22	1:K:151:SER:N	2.36	0.41
1:K:62:LEU:HB2	1:K:68:ASN:HA	2.03	0.41
1:L:419:LEU:HD11	1:L:500:THR:HG23	2.02	0.41
1:L:518:GLU:HB3	1:M:29:VAL:HG21	2.02	0.41
1:M:202:PRO:O	1:M:205:ILE:HG12	2.20	0.41
1:N:140:ASP:OD2	1:N:142:LYS:HB3	2.21	0.41
1:N:179:ASP:HA	1:N:389:MET:SD	2.61	0.41
1:N:65:LYS:O	1:N:69:MET:HG3	2.21	0.41
2:P:17:VAL:HG12	2:P:35:SER:N	2.36	0.41
2:T:12:VAL:HG23	2:T:84:LEU:HD12	2.02	0.41
2:V:2:ASN:HD22	2:V:2:ASN:H	1.69	0.41
2:W:11:ILE:HB	2:W:85:ILE:HD13	2.03	0.41
2:Y:92:LEU:O	2:Z:9:ARG:HD2	2.21	0.41
2:2:10:VAL:HG12	2:2:11:ILE:N	2.36	0.40
1:B:129:GLU:OE2	1:B:132:LYS:HD2	2.21	0.40
1:B:433:ASN:OD1	1:B:436:GLN:HG3	2.21	0.40
1:C:19:GLY:HA3	1:C:67:GLU:O	2.21	0.40
1:D:183:LEU:HD12	1:D:183:LEU:N	2.36	0.40
1:D:406:ALA:HB2	1:D:496:PRO:HB3	2.03	0.40
1:E:247:LEU:HD22	1:E:247:LEU:N	2.36	0.40
1:E:199:TYR:CB	1:E:276:VAL:HG12	2.51	0.40
1:E:494:LEU:HD12	1:E:494:LEU:N	2.35	0.40
1:F:236:VAL:HG11	1:F:312:ALA:CB	2.38	0.40
1:F:420:ILE:HG12	1:F:448:GLU:HA	2.03	0.40
1:G:420:ILE:HD11	1:G:470:LYS:CG	2.51	0.40
1:H:245:LYS:CB	1:H:246:PRO:CD	3.00	0.40
1:H:24:ALA:O	1:H:28:LYS:HG3	2.22	0.40
1:H:301:ILE:HD13	1:H:307:MET:CE	2.51	0.40
1:H:77:VAL:HG23	1:H:506:TYR:C	2.41	0.40
1:J:5:ASP:HB2	1:J:524:LEU:HD21	2.02	0.40
1:K:241:ALA:HB2	1:K:271:VAL:HG11	2.03	0.40
1:M:226:LYS:HG2	1:M:252:GLU:HG2	2.02	0.40
1:N:234:LEU:O	1:N:238:GLU:HG3	2.21	0.40
2:P:17:VAL:CG1	2:P:35:SER:HB2	2.43	0.40
2:Y:49:LEU:HD23	2:Y:49:LEU:C	2.42	0.40
1:A:272:LYS:HG3	1:A:272:LYS:O	2.21	0.40

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:25:ASP:OD1	1:A:28:LYS:HE2	2.21	0.40
1:A:77:VAL:HG21	1:A:507:ALA:CA	2.47	0.40
1:C:313:THR:HG22	1:C:314:LEU:N	2.36	0.40
1:E:346:VAL:O	1:E:350:ARG:HG2	2.21	0.40
1:E:482:THR:OG1	1:E:484:GLU:HG2	2.21	0.40
1:F:116:LEU:O	1:F:120:ILE:HG13	2.21	0.40
1:F:400:LEU:HD23	1:F:400:LEU:C	2.41	0.40
1:G:20:VAL:HG13	1:G:74:VAL:HG11	2.03	0.40
1:G:309:LEU:HD23	1:G:310:GLU:H	1.85	0.40
1:G:383:ALA:CB	1:G:389:MET:HA	2.51	0.40
1:G:417:VAL:O	1:G:421:ARG:HB2	2.21	0.40
1:G:80:LYS:HD2	1:G:506:TYR:CZ	2.56	0.40
1:H:358:SER:HA	1:H:362:ARG:NE	2.36	0.40
1:I:116:LEU:O	1:I:120:ILE:HG13	2.22	0.40
1:I:288:MET:HG3	1:I:368:ARG:NH1	2.35	0.40
1:J:160:LYS:HA	1:J:160:LYS:HD2	1.81	0.40
1:J:23:LEU:HD23	1:J:60:ILE:HG21	2.03	0.40
1:K:142:LYS:O	1:K:146:GLN:HG3	2.21	0.40
1:K:30:THR:OG1	1:K:90:THR:HG21	2.21	0.40
1:N:433:ASN:OD1	1:N:436:GLN:HG3	2.21	0.40
1:N:524:LEU:HA	1:N:524:LEU:HD23	1.91	0.40
2:Q:10:VAL:HG22	2:Q:11:ILE:N	2.36	0.40
2:Q:43:VAL:HG22	2:Q:44:GLY:N	2.36	0.40
2:S:77:LYS:HD3	2:S:82:GLU:HA	2.03	0.40
2:U:6:LEU:CD1	2:U:7:HIS:HB2	2.51	0.40
2:T:92:LEU:CA	2:U:9:ARG:HD2	2.50	0.40
2:1:6:LEU:HA	2:1:6:LEU:HD23	1.79	0.40
1:A:49:ILE:O	1:A:391:GLU:HB2	2.21	0.40
1:B:160:LYS:O	1:B:164:GLU:HG3	2.21	0.40
1:B:131:LEU:CD2	1:B:422:VAL:HG11	2.51	0.40
1:B:513:LEU:CD1	1:C:388:GLU:HA	2.51	0.40
1:D:195:PHE:CE1	1:D:330:THR:HB	2.56	0.40
1:D:278:ALA:HB3	1:D:285:ARG:CD	2.50	0.40
1:E:288:MET:HA	1:E:291:ASP:OD2	2.21	0.40
1:E:419:LEU:HB3	1:E:447:MET:HB3	2.02	0.40
1:F:282:GLY:HA2	1:F:285:ARG:CG	2.47	0.40
1:G:226:LYS:HD3	1:G:255:GLU:OE2	2.21	0.40
1:G:289:LEU:HD23	1:G:289:LEU:HA	1.83	0.40
1:G:295:LEU:C	1:G:295:LEU:HD23	2.41	0.40
1:I:233:MET:CB	1:I:309:LEU:CD2	3.00	0.40
1:I:241:ALA:HB1	2:W:25:ILE:CD1	2.43	0.40

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:L:250:ILE:HG22	1:L:251:ALA:N	2.35	0.40
1:L:307:MET:O	1:L:308:GLU:HG2	2.21	0.40
1:N:119:GLY:O	1:N:440:ILE:HG12	2.21	0.40
2:O:17:VAL:HG22	2:O:18:GLU:HG3	2.02	0.40
2:O:40:VAL:HG12	2:O:41:LEU:N	2.37	0.40
2:S:27:LEU:HD12	2:S:28:THR:O	2.21	0.40
2:T:8:ASP:HB2	2:T:47:ARG:HB3	2.03	0.40
2:V:11:ILE:HG13	2:V:42:ALA:O	2.22	0.40
2:Z:49:LEU:O	2:Z:52:GLY:N	2.49	0.40
1:B:231:ARG:HH22	2:P:31:ALA:CB	2.27	0.40
1:B:307:MET:HE3	1:B:311:LYS:HZ2	1.86	0.40
1:B:406:ALA:CA	1:B:496:PRO:HB3	2.51	0.40
1:C:236:VAL:O	1:C:240:VAL:HG23	2.21	0.40
1:D:146:GLN:HB2	1:D:494:LEU:HD12	2.04	0.40
1:E:247:LEU:HD23	1:E:273:VAL:HG22	2.02	0.40
1:E:357:THR:HG22	1:E:357:THR:O	2.21	0.40
1:D:69:MET:CE	1:E:39:VAL:HG12	2.52	0.40
1:F:38:VAL:HB	1:F:56:VAL:CG2	2.52	0.40
1:G:124:VAL:HG13	1:G:504:LEU:HD12	2.02	0.40
1:G:236:VAL:CG2	1:G:312:ALA:HB3	2.50	0.40
1:H:131:LEU:HD13	1:H:500:THR:HG22	2.04	0.40
1:H:433:ASN:OD1	1:H:435:ASP:HB2	2.22	0.40
1:I:123:ALA:HB2	1:I:440:ILE:HG23	2.04	0.40
1:I:263:VAL:O	1:I:267:MET:HG3	2.20	0.40
1:I:200:LEU:HD13	1:I:276:VAL:HA	2.03	0.40
1:K:322:ARG:HB3	1:K:333:ILE:HG21	2.01	0.40
1:L:388:GLU:O	1:L:392:LYS:HB2	2.22	0.40
1:L:30:THR:O	1:L:51:LYS:HE2	2.21	0.40
1:M:92:ALA:HB2	1:M:503:ALA:CB	2.51	0.40
1:N:13:ARG:CD	1:N:104:LEU:HD22	2.50	0.40
1:N:280:GLY:O	1:N:285:ARG:HB3	2.22	0.40
1:A:238:GLU:OE2	2:O:24:GLY:HA3	2.21	0.40
2:R:17:VAL:HG23	2:R:19:THR:CA	2.52	0.40
2:R:36:THR:CG2	2:R:37:ARG:HG3	2.34	0.40
1:E:238:GLU:OE1	2:S:25:ILE:HD13	2.20	0.40
2:U:6:LEU:C	2:U:6:LEU:HD12	2.41	0.40
1:J:238:GLU:OE1	2:X:25:ILE:HD13	2.21	0.40
2:X:3:ILE:HD12	2:X:3:ILE:O	2.22	0.40
2:Z:37:ARG:HG2	2:Z:66:ILE:HG12	2.02	0.40
2:Z:48:ILE:O	2:Z:48:ILE:HG23	2.21	0.40
1:A:112:ASN:HA	1:A:113:PRO:HD3	1.89	0.40

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:B:113:PRO:O	1:B:116:LEU:HB2	2.21	0.40
1:B:15:LYS:O	1:B:67:GLU:HG2	2.22	0.40
1:C:236:VAL:HG22	1:C:312:ALA:O	2.22	0.40
1:D:295:LEU:HD23	1:D:295:LEU:C	2.42	0.40
1:E:313:THR:HG22	1:E:314:LEU:N	2.37	0.40
1:F:429:LEU:HG	1:F:440:ILE:HD13	2.04	0.40
1:G:197:ARG:HD3	1:G:197:ARG:HA	1.93	0.40
1:G:305:ILE:CG2	1:G:306:GLY:H	2.35	0.40
1:I:295:LEU:HD12	1:I:372:LEU:HD23	2.02	0.40
1:I:421:ARG:NH1	1:I:472:GLY:O	2.54	0.40
1:J:436:GLN:O	1:J:440:ILE:HG13	2.21	0.40
1:K:51:LYS:NZ	3:K:601:ADP:O2A	2.47	0.40
1:L:381:VAL:HG22	1:L:396:VAL:HG21	2.02	0.40
1:L:87:ASP:O	1:L:499:VAL:HG13	2.21	0.40
1:M:295:LEU:HD23	1:M:295:LEU:O	2.22	0.40
1:N:143:ALA:O	1:N:146:GLN:HB2	2.21	0.40
1:N:20:VAL:HG13	1:N:74:VAL:HG21	2.04	0.40
1:N:281:PHE:O	1:N:284:ARG:N	2.55	0.40
2:O:47:ARG:O	2:O:54:VAL:HA	2.20	0.40
2:P:7:HIS:HA	2:P:45:ASN:OD1	2.22	0.40
2:T:92:LEU:HD12	2:T:92:LEU:N	2.36	0.40
2:U:65:VAL:HG12	2:U:94:ILE:CG2	2.52	0.40
2:V:37:ARG:HH21	2:W:76:GLU:CG	2.34	0.40
2:Y:92:LEU:HD23	2:Z:85:ILE:HD11	2.03	0.40

All (16) symmetry-related close contacts are listed below. The label for Atom-2 includes the symmetry operator and encoded unit-cell translations to be applied.

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:B:339:GLU:OE2	1:K:207:LYS:NZ[1_545]	0.50	1.70
1:K:430:ARG:CG	1:M:245:LYS:NZ[4_565]	0.86	1.34
1:B:339:GLU:CD	1:K:207:LYS:NZ[1_545]	1.06	1.14
1:K:430:ARG:CB	1:M:245:LYS:NZ[4_565]	1.09	1.11
1:B:339:GLU:OE2	1:K:207:LYS:CE[1_545]	1.58	0.62
1:K:430:ARG:CA	1:M:245:LYS:NZ[4_565]	1.76	0.44
1:K:430:ARG:CG	1:M:245:LYS:CE[4_565]	1.78	0.42
1:G:472:GLY:N	2:P:2:ASN:ND2[3_554]	1.79	0.41
1:G:471:GLY:O	2:P:2:ASN:ND2[3_554]	1.84	0.36
1:K:430:ARG:CD	1:M:245:LYS:NZ[4_565]	1.86	0.34
1:L:476:TYR:OH	2:2:79:ASP:OD2[4_565]	1.95	0.25

Continued on next page...

Continued from previous page...

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:K:430:ARG:CB	1:M:245:LYS:CE[4_565]	2.03	0.17
1:B:339:GLU:CG	1:K:207:LYS:NZ[1_545]	2.06	0.14
2:T:53:GLU:OE1	2:1:53:GLU:OE2[1_645]	2.09	0.11
1:B:339:GLU:OE1	1:K:207:LYS:NZ[1_545]	2.11	0.09
1:K:430:ARG:N	1:M:245:LYS:CE[4_565]	2.17	0.03

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 X-ray entries followed by that with respect to entries of similar resolution.

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

Mol	Chain	Analysed	Favoured	Allowed	Outliers	Percentiles	
1	A	522/548 (95%)	496 (95%)	23 (4%)	3 (1%)	25	62
1	B	522/548 (95%)	499 (96%)	21 (4%)	2 (0%)	34	70
1	C	522/548 (95%)	509 (98%)	13 (2%)	0	100	100
1	D	522/548 (95%)	502 (96%)	19 (4%)	1 (0%)	47	78
1	E	522/548 (95%)	500 (96%)	18 (3%)	4 (1%)	19	56
1	F	522/548 (95%)	504 (97%)	16 (3%)	2 (0%)	34	70
1	G	522/548 (95%)	508 (97%)	12 (2%)	2 (0%)	34	70
1	H	522/548 (95%)	502 (96%)	19 (4%)	1 (0%)	47	78
1	I	522/548 (95%)	507 (97%)	14 (3%)	1 (0%)	47	78
1	J	522/548 (95%)	503 (96%)	17 (3%)	2 (0%)	34	70
1	K	522/548 (95%)	507 (97%)	15 (3%)	0	100	100
1	L	522/548 (95%)	504 (97%)	18 (3%)	0	100	100
1	M	522/548 (95%)	503 (96%)	16 (3%)	3 (1%)	25	62
1	N	522/548 (95%)	503 (96%)	19 (4%)	0	100	100
2	1	95/97 (98%)	87 (92%)	4 (4%)	4 (4%)	3	26
2	2	95/97 (98%)	87 (92%)	7 (7%)	1 (1%)	14	50
2	O	93/97 (96%)	89 (96%)	4 (4%)	0	100	100

Continued on next page...

Continued from previous page...

Mol	Chain	Analysed	Favoured	Allowed	Outliers	Percentiles	
2	P	95/97 (98%)	86 (90%)	9 (10%)	0	100	100
2	Q	94/97 (97%)	84 (89%)	9 (10%)	1 (1%)	14	50
2	R	95/97 (98%)	88 (93%)	6 (6%)	1 (1%)	14	50
2	S	94/97 (97%)	83 (88%)	10 (11%)	1 (1%)	14	50
2	T	95/97 (98%)	90 (95%)	5 (5%)	0	100	100
2	U	95/97 (98%)	85 (90%)	8 (8%)	2 (2%)	7	38
2	V	95/97 (98%)	89 (94%)	5 (5%)	1 (1%)	14	50
2	W	95/97 (98%)	84 (88%)	10 (10%)	1 (1%)	14	50
2	X	95/97 (98%)	87 (92%)	6 (6%)	2 (2%)	7	38
2	Y	94/97 (97%)	90 (96%)	4 (4%)	0	100	100
2	Z	93/97 (96%)	89 (96%)	4 (4%)	0	100	100
All	All	8631/9030 (96%)	8265 (96%)	331 (4%)	35 (0%)	34	70

All (35) Ramachandran outliers are listed below:

Mol	Chain	Res	Type
1	D	233	MET
1	E	233	MET
1	E	234	LEU
1	M	270	ILE
2	R	50	GLU
2	S	40	VAL
1	A	270	ILE
1	B	228	SER
1	H	228	SER
1	I	231	ARG
2	X	21	SER
2	X	51	ASN
2	1	6	LEU
2	1	7	HIS
1	B	245	LYS
1	E	235	PRO
1	F	305	ILE
1	G	209	GLU
2	Q	52	GLY
2	U	51	ASN
1	A	271	VAL
1	A	309	LEU

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type
1	F	218	PRO
1	G	223	ALA
1	J	231	ARG
2	U	52	GLY
2	W	26	VAL
2	2	54	VAL
1	M	234	LEU
1	M	334	ASP
2	1	5	PRO
1	E	207	LYS
1	J	270	ILE
2	V	92	LEU
2	1	8	ASP

5.3.2 Protein sidechains ⓘ

In the following table, the Percentiles column shows the percent sidechain outliers of the chain as a percentile score with respect to all X-ray entries followed by that with respect to entries of similar resolution.

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

Mol	Chain	Analysed	Rotameric	Outliers	Percentiles	
1	A	404/415 (97%)	401 (99%)	3 (1%)	84	91
1	B	404/415 (97%)	401 (99%)	3 (1%)	84	91
1	C	404/415 (97%)	400 (99%)	4 (1%)	76	86
1	D	404/415 (97%)	400 (99%)	4 (1%)	76	86
1	E	404/415 (97%)	402 (100%)	2 (0%)	88	94
1	F	404/415 (97%)	400 (99%)	4 (1%)	76	86
1	G	404/415 (97%)	400 (99%)	4 (1%)	76	86
1	H	404/415 (97%)	396 (98%)	8 (2%)	55	74
1	I	404/415 (97%)	400 (99%)	4 (1%)	76	86
1	J	404/415 (97%)	397 (98%)	7 (2%)	60	78
1	K	404/415 (97%)	399 (99%)	5 (1%)	71	84
1	L	404/415 (97%)	401 (99%)	3 (1%)	84	91
1	M	404/415 (97%)	399 (99%)	5 (1%)	71	84
1	N	404/415 (97%)	394 (98%)	10 (2%)	47	70

Continued on next page...

Continued from previous page...

Mol	Chain	Analysed	Rotameric	Outliers	Percentiles	
2	1	80/80 (100%)	77 (96%)	3 (4%)	33	60
2	2	80/80 (100%)	76 (95%)	4 (5%)	24	54
2	O	78/80 (98%)	76 (97%)	2 (3%)	46	69
2	P	80/80 (100%)	77 (96%)	3 (4%)	33	60
2	Q	79/80 (99%)	74 (94%)	5 (6%)	18	48
2	R	80/80 (100%)	78 (98%)	2 (2%)	47	70
2	S	79/80 (99%)	74 (94%)	5 (6%)	18	48
2	T	80/80 (100%)	79 (99%)	1 (1%)	69	82
2	U	80/80 (100%)	79 (99%)	1 (1%)	69	82
2	V	80/80 (100%)	75 (94%)	5 (6%)	18	48
2	W	80/80 (100%)	77 (96%)	3 (4%)	33	60
2	X	80/80 (100%)	75 (94%)	5 (6%)	18	48
2	Y	80/80 (100%)	78 (98%)	2 (2%)	47	70
2	Z	79/80 (99%)	77 (98%)	2 (2%)	47	70
All	All	6771/6930 (98%)	6662 (98%)	109 (2%)	62	79

All (109) residues with a non-rotameric sidechain are listed below:

Mol	Chain	Res	Type
1	A	308	GLU
1	A	362	ARG
1	A	499	VAL
1	B	169	VAL
1	B	343	GLN
1	B	493	ILE
1	C	136	VAL
1	C	366	GLN
1	C	453	GLN
1	C	495	ASP
1	D	227	ILE
1	D	292	ILE
1	D	363	GLU
1	D	499	VAL
1	E	234	LEU
1	E	420	ILE
1	F	105	LYS
1	F	207	LYS

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type
1	F	363	GLU
1	F	499	VAL
1	G	147	VAL
1	G	309	LEU
1	G	361	ASP
1	G	397	GLU
1	H	136	VAL
1	H	185	ASP
1	H	207	LYS
1	H	237	LEU
1	H	255	GLU
1	H	363	GLU
1	H	453	GLN
1	H	513	LEU
1	I	74	VAL
1	I	136	VAL
1	I	348	GLN
1	I	513	LEU
1	J	207	LYS
1	J	234	LEU
1	J	245	LYS
1	J	309	LEU
1	J	334	ASP
1	J	349	ILE
1	J	499	VAL
1	K	17	LEU
1	K	77	VAL
1	K	206	ASN
1	K	329	THR
1	K	513	LEU
1	L	510	VAL
1	L	513	LEU
1	L	523	ASP
1	M	147	VAL
1	M	216	GLU
1	M	224	ASP
1	M	467	ASN
1	M	513	LEU
1	N	17	LEU
1	N	56	VAL
1	N	91	THR
1	N	147	VAL

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type
1	N	150	ILE
1	N	206	ASN
1	N	361	ASP
1	N	467	ASN
1	N	513	LEU
1	N	523	ASP
2	O	8	ASP
2	O	49	LEU
2	P	3	ILE
2	P	4	ARG
2	P	53	GLU
2	Q	28	THR
2	Q	57	LEU
2	Q	79	ASP
2	Q	80	ASN
2	Q	92	LEU
2	R	49	LEU
2	R	80	ASN
2	S	6	LEU
2	S	50	GLU
2	S	53	GLU
2	S	61	VAL
2	S	80	ASN
2	T	84	LEU
2	U	8	ASP
2	V	2	ASN
2	V	12	VAL
2	V	51	ASN
2	V	58	ASP
2	V	88	GLU
2	W	8	ASP
2	W	9	ARG
2	W	51	ASN
2	X	2	ASN
2	X	6	LEU
2	X	21	SER
2	X	48	ILE
2	X	66	ILE
2	Y	7	HIS
2	Y	89	SER
2	Z	51	ASN
2	Z	95	VAL

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type
2	1	71	TYR
2	1	80	ASN
2	1	94	ILE
2	2	43	VAL
2	2	45	ASN
2	2	78	ILE
2	2	88	GLU

Some sidechains can be flipped to improve hydrogen bonding and reduce clashes. All (12) such sidechains are listed below:

Mol	Chain	Res	Type
1	B	351	GLN
1	F	348	GLN
1	F	352	GLN
1	F	366	GLN
1	H	194	GLN
1	I	194	GLN
1	J	453	GLN
1	L	453	GLN
1	M	194	GLN
2	O	7	HIS
2	T	51	ASN
2	V	2	ASN

5.3.3 RNA ⓘ

There are no RNA molecules in this entry.

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

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

5.5 Carbohydrates ⓘ

There are no carbohydrates in this entry.

5.6 Ligand geometry ⓘ

Of 56 ligands modelled in this entry, 28 are monoatomic - leaving 28 for Mogul analysis.

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

Mol	Type	Chain	Res	Link	Bond lengths			Bond angles		
					Counts	RMSZ	# $ Z > 2$	Counts	RMSZ	# $ Z > 2$
4	BEF	D	602	-	0,3,3	0.00	-	-		
3	ADP	B	601	5,6	24,29,29	0.94	1 (4%)	29,45,45	1.45	4 (13%)
4	BEF	L	602	-	0,3,3	0.00	-	-		
3	ADP	H	601	5,6	24,29,29	0.95	1 (4%)	29,45,45	1.66	5 (17%)
3	ADP	J	601	5,6	24,29,29	0.93	1 (4%)	29,45,45	1.56	6 (20%)
4	BEF	B	602	-	0,3,3	0.00	-	-		
3	ADP	A	601	5,4,6	24,29,29	0.94	1 (4%)	29,45,45	1.48	4 (13%)
4	BEF	I	602	-	0,3,3	0.00	-	-		
4	BEF	K	602	-	0,3,3	0.00	-	-		
4	BEF	A	602	3	0,3,3	0.00	-	-		
4	BEF	J	602	-	0,3,3	0.00	-	-		
3	ADP	C	601	5,6	24,29,29	0.95	1 (4%)	29,45,45	1.46	4 (13%)
3	ADP	I	601	5,6	24,29,29	0.95	1 (4%)	29,45,45	1.53	4 (13%)
3	ADP	K	601	5,6	24,29,29	0.94	1 (4%)	29,45,45	1.50	4 (13%)
3	ADP	D	601	5,6	24,29,29	0.93	1 (4%)	29,45,45	1.58	5 (17%)
4	BEF	G	602	-	0,3,3	0.00	-	-		
4	BEF	N	602	-	0,3,3	0.00	-	-		
3	ADP	L	601	5,6	24,29,29	0.94	1 (4%)	29,45,45	1.56	6 (20%)
4	BEF	F	602	-	0,3,3	0.00	-	-		
4	BEF	E	602	-	0,3,3	0.00	-	-		
4	BEF	M	602	-	0,3,3	0.00	-	-		
3	ADP	E	601	5,6	24,29,29	0.92	1 (4%)	29,45,45	1.54	5 (17%)
3	ADP	M	601	5,6	24,29,29	0.96	1 (4%)	29,45,45	1.49	5 (17%)
3	ADP	F	601	5,6	24,29,29	0.96	1 (4%)	29,45,45	1.50	6 (20%)
3	ADP	N	601	5,6	24,29,29	0.94	1 (4%)	29,45,45	1.46	5 (17%)
4	BEF	H	602	-	0,3,3	0.00	-	-		
3	ADP	G	601	5,6	24,29,29	0.95	1 (4%)	29,45,45	1.37	3 (10%)
4	BEF	C	602	-	0,3,3	0.00	-	-		

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
3	ADP	A	601	5,4,6	-	1/12/32/32	0/3/3/3
3	ADP	C	601	5,6	-	4/12/32/32	0/3/3/3
3	ADP	I	601	5,6	-	6/12/32/32	0/3/3/3
3	ADP	K	601	5,6	-	5/12/32/32	0/3/3/3
3	ADP	E	601	5,6	-	4/12/32/32	0/3/3/3
3	ADP	D	601	5,6	-	6/12/32/32	0/3/3/3
3	ADP	F	601	5,6	-	3/12/32/32	0/3/3/3
3	ADP	M	601	5,6	-	7/12/32/32	0/3/3/3
3	ADP	B	601	5,6	-	2/12/32/32	0/3/3/3
3	ADP	L	601	5,6	-	1/12/32/32	0/3/3/3
3	ADP	N	601	5,6	-	8/12/32/32	0/3/3/3
3	ADP	H	601	5,6	-	6/12/32/32	0/3/3/3
3	ADP	J	601	5,6	-	6/12/32/32	0/3/3/3
3	ADP	G	601	5,6	-	4/12/32/32	0/3/3/3

All (14) bond length outliers are listed below:

Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
3	C	601	ADP	C5-C4	2.44	1.47	1.40
3	H	601	ADP	C5-C4	2.39	1.47	1.40
3	B	601	ADP	C5-C4	2.36	1.47	1.40
3	N	601	ADP	C5-C4	2.35	1.47	1.40
3	A	601	ADP	C5-C4	2.34	1.47	1.40
3	J	601	ADP	C5-C4	2.33	1.47	1.40
3	G	601	ADP	C5-C4	2.33	1.47	1.40
3	E	601	ADP	C5-C4	2.32	1.47	1.40
3	L	601	ADP	C5-C4	2.32	1.47	1.40
3	F	601	ADP	C5-C4	2.32	1.47	1.40
3	K	601	ADP	C5-C4	2.28	1.46	1.40
3	M	601	ADP	C5-C4	2.27	1.46	1.40
3	I	601	ADP	C5-C4	2.26	1.46	1.40
3	D	601	ADP	C5-C4	2.23	1.46	1.40

All (66) bond angle outliers are listed below:

Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
3	H	601	ADP	PA-O3A-PB	-4.16	118.55	132.83
3	I	601	ADP	PA-O3A-PB	-4.03	118.98	132.83
3	F	601	ADP	PA-O3A-PB	-3.91	119.42	132.83

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
3	F	601	ADP	N3-C2-N1	-3.84	122.68	128.68
3	E	601	ADP	N3-C2-N1	-3.83	122.70	128.68
3	G	601	ADP	N3-C2-N1	-3.78	122.77	128.68
3	L	601	ADP	N3-C2-N1	-3.77	122.78	128.68
3	D	601	ADP	N3-C2-N1	-3.71	122.88	128.68
3	H	601	ADP	N3-C2-N1	-3.70	122.89	128.68
3	C	601	ADP	PA-O3A-PB	-3.70	120.14	132.83
3	K	601	ADP	N3-C2-N1	-3.66	122.95	128.68
3	J	601	ADP	N3-C2-N1	-3.63	123.00	128.68
3	A	601	ADP	N3-C2-N1	-3.61	123.03	128.68
3	N	601	ADP	N3-C2-N1	-3.59	123.07	128.68
3	M	601	ADP	N3-C2-N1	-3.58	123.08	128.68
3	D	601	ADP	PA-O3A-PB	-3.56	120.61	132.83
3	I	601	ADP	N3-C2-N1	-3.49	123.22	128.68
3	B	601	ADP	N3-C2-N1	-3.46	123.26	128.68
3	C	601	ADP	N3-C2-N1	-3.44	123.31	128.68
3	K	601	ADP	PA-O3A-PB	-3.37	121.25	132.83
3	H	601	ADP	C3'-C2'-C1'	3.28	105.91	100.98
3	M	601	ADP	PA-O3A-PB	-3.25	121.66	132.83
3	E	601	ADP	C3'-C2'-C1'	3.24	105.86	100.98
3	A	601	ADP	C3'-C2'-C1'	3.15	105.72	100.98
3	J	601	ADP	C3'-C2'-C1'	3.14	105.70	100.98
3	J	601	ADP	C4-C5-N7	-3.12	106.15	109.40
3	B	601	ADP	C4-C5-N7	-3.10	106.17	109.40
3	K	601	ADP	C3'-C2'-C1'	2.99	105.48	100.98
3	L	601	ADP	C3'-C2'-C1'	2.97	105.45	100.98
3	N	601	ADP	C3'-C2'-C1'	2.97	105.44	100.98
3	L	601	ADP	PA-O3A-PB	-2.97	122.65	132.83
3	N	601	ADP	C4-C5-N7	-2.95	106.32	109.40
3	H	601	ADP	C4-C5-N7	-2.94	106.33	109.40
3	D	601	ADP	C3'-C2'-C1'	2.91	105.36	100.98
3	N	601	ADP	PA-O3A-PB	-2.90	122.86	132.83
3	M	601	ADP	C4-C5-N7	-2.87	106.41	109.40
3	D	601	ADP	C4-C5-N7	-2.76	106.53	109.40
3	A	601	ADP	C4-C5-N7	-2.74	106.55	109.40
3	E	601	ADP	PA-O3A-PB	-2.74	123.44	132.83
3	B	601	ADP	C3'-C2'-C1'	2.72	105.07	100.98
3	J	601	ADP	PA-O3A-PB	-2.71	123.53	132.83
3	A	601	ADP	PA-O3A-PB	-2.71	123.53	132.83
3	B	601	ADP	PA-O3A-PB	-2.67	123.66	132.83
3	K	601	ADP	C4-C5-N7	-2.63	106.66	109.40
3	C	601	ADP	C4-C5-N7	-2.62	106.67	109.40

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
3	G	601	ADP	C4-C5-N7	-2.59	106.69	109.40
3	E	601	ADP	C4-C5-N7	-2.59	106.70	109.40
3	F	601	ADP	C4-C5-N7	-2.54	106.75	109.40
3	L	601	ADP	C4-C5-N7	-2.51	106.78	109.40
3	C	601	ADP	C3'-C2'-C1'	2.48	104.72	100.98
3	I	601	ADP	C4-C5-N7	-2.45	106.84	109.40
3	G	601	ADP	PA-O3A-PB	-2.44	124.45	132.83
3	M	601	ADP	C3'-C2'-C1'	2.41	104.60	100.98
3	I	601	ADP	C3'-C2'-C1'	2.27	104.40	100.98
3	F	601	ADP	O3B-PB-O2B	2.24	116.20	107.64
3	L	601	ADP	O3B-PB-O2B	2.18	115.96	107.64
3	F	601	ADP	C3'-C2'-C1'	2.09	104.12	100.98
3	J	601	ADP	C2-N1-C6	2.08	122.31	118.75
3	H	601	ADP	C2-N1-C6	2.07	122.30	118.75
3	M	601	ADP	O3B-PB-O2B	2.07	115.56	107.64
3	D	601	ADP	C2-N1-C6	2.06	122.28	118.75
3	E	601	ADP	C2-N1-C6	2.06	122.28	118.75
3	F	601	ADP	C2-N1-C6	2.03	122.23	118.75
3	L	601	ADP	C2-N1-C6	2.03	122.23	118.75
3	J	601	ADP	C1'-N9-C4	-2.03	123.08	126.64
3	N	601	ADP	C2-N1-C6	2.00	122.18	118.75

There are no chirality outliers.

All (63) torsion outliers are listed below:

Mol	Chain	Res	Type	Atoms
3	H	601	ADP	C5'-O5'-PA-O1A
3	H	601	ADP	C5'-O5'-PA-O2A
3	J	601	ADP	PA-O3A-PB-O2B
3	J	601	ADP	C5'-O5'-PA-O1A
3	J	601	ADP	C5'-O5'-PA-O2A
3	C	601	ADP	C5'-O5'-PA-O2A
3	C	601	ADP	C5'-O5'-PA-O3A
3	I	601	ADP	C5'-O5'-PA-O2A
3	I	601	ADP	C3'-C4'-C5'-O5'
3	K	601	ADP	C5'-O5'-PA-O3A
3	K	601	ADP	C3'-C4'-C5'-O5'
3	D	601	ADP	O4'-C4'-C5'-O5'
3	F	601	ADP	C3'-C4'-C5'-O5'
3	M	601	ADP	PA-O3A-PB-O2B
3	M	601	ADP	PB-O3A-PA-O5'
3	M	601	ADP	C5'-O5'-PA-O3A

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type	Atoms
3	M	601	ADP	O4'-C4'-C5'-O5'
3	N	601	ADP	PB-O3A-PA-O5'
3	G	601	ADP	C5'-O5'-PA-O1A
3	G	601	ADP	C5'-O5'-PA-O3A
3	C	601	ADP	C3'-C4'-C5'-O5'
3	M	601	ADP	C3'-C4'-C5'-O5'
3	N	601	ADP	O4'-C4'-C5'-O5'
3	N	601	ADP	C3'-C4'-C5'-O5'
3	I	601	ADP	O4'-C4'-C5'-O5'
3	K	601	ADP	O4'-C4'-C5'-O5'
3	D	601	ADP	C3'-C4'-C5'-O5'
3	F	601	ADP	O4'-C4'-C5'-O5'
3	L	601	ADP	C3'-C4'-C5'-O5'
3	C	601	ADP	O4'-C4'-C5'-O5'
3	G	601	ADP	PA-O3A-PB-O1B
3	I	601	ADP	PB-O3A-PA-O5'
3	D	601	ADP	PB-O3A-PA-O5'
3	E	601	ADP	PB-O3A-PA-O5'
3	H	601	ADP	O4'-C4'-C5'-O5'
3	N	601	ADP	PA-O3A-PB-O1B
3	K	601	ADP	PA-O3A-PB-O2B
3	B	601	ADP	C5'-O5'-PA-O3A
3	H	601	ADP	C5'-O5'-PA-O3A
3	D	601	ADP	C5'-O5'-PA-O3A
3	N	601	ADP	C5'-O5'-PA-O3A
3	H	601	ADP	PB-O3A-PA-O2A
3	K	601	ADP	C5'-O5'-PA-O2A
3	D	601	ADP	C5'-O5'-PA-O2A
3	M	601	ADP	C5'-O5'-PA-O1A
3	N	601	ADP	C5'-O5'-PA-O2A
3	M	601	ADP	PA-O3A-PB-O1B
3	F	601	ADP	C4'-C5'-O5'-PA
3	G	601	ADP	PB-O3A-PA-O2A
3	E	601	ADP	O4'-C4'-C5'-O5'
3	D	601	ADP	PB-O3A-PA-O1A
3	E	601	ADP	PB-O3A-PA-O1A
3	J	601	ADP	O4'-C4'-C5'-O5'
3	J	601	ADP	PA-O3A-PB-O1B
3	N	601	ADP	PA-O3A-PB-O2B
3	N	601	ADP	PA-O3A-PB-O3B
3	J	601	ADP	C5'-O5'-PA-O3A
3	I	601	ADP	C5'-O5'-PA-O3A

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type	Atoms
3	E	601	ADP	C5'-O5'-PA-O3A
3	H	601	ADP	PB-O3A-PA-O1A
3	A	601	ADP	PB-O3A-PA-O1A
3	B	601	ADP	C5'-O5'-PA-O1A
3	I	601	ADP	C5'-O5'-PA-O1A

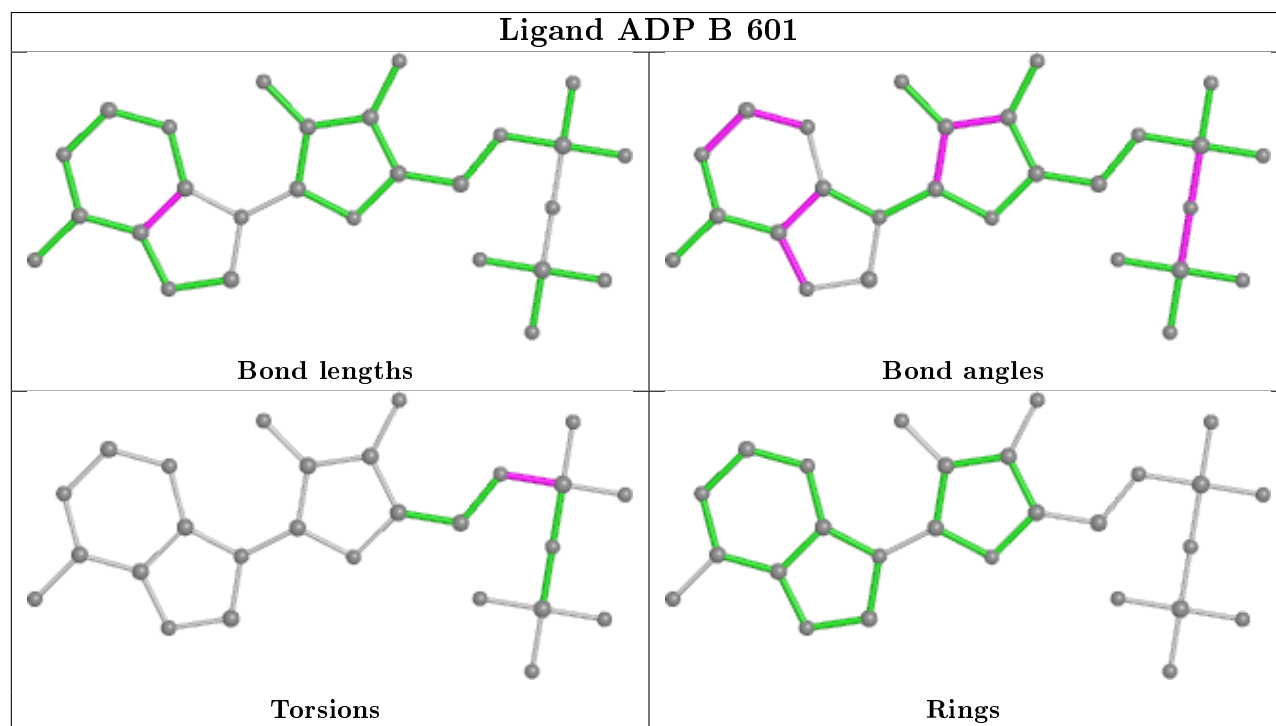
There are no ring outliers.

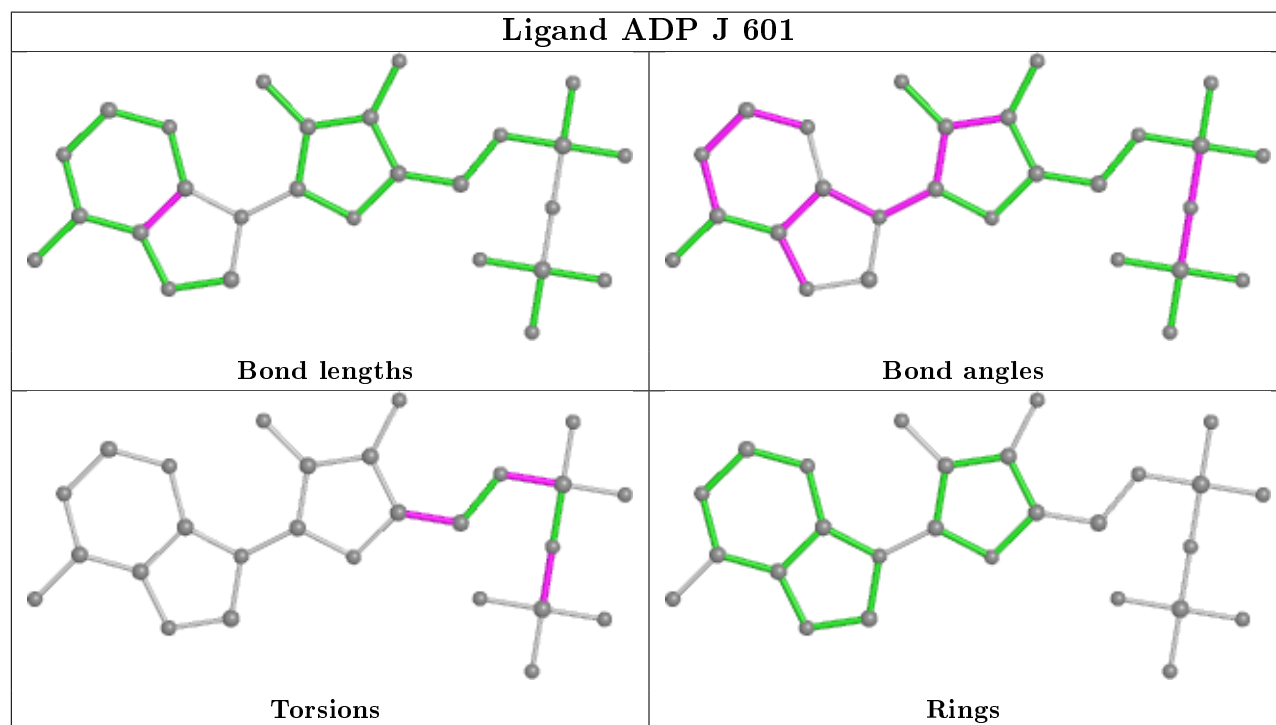
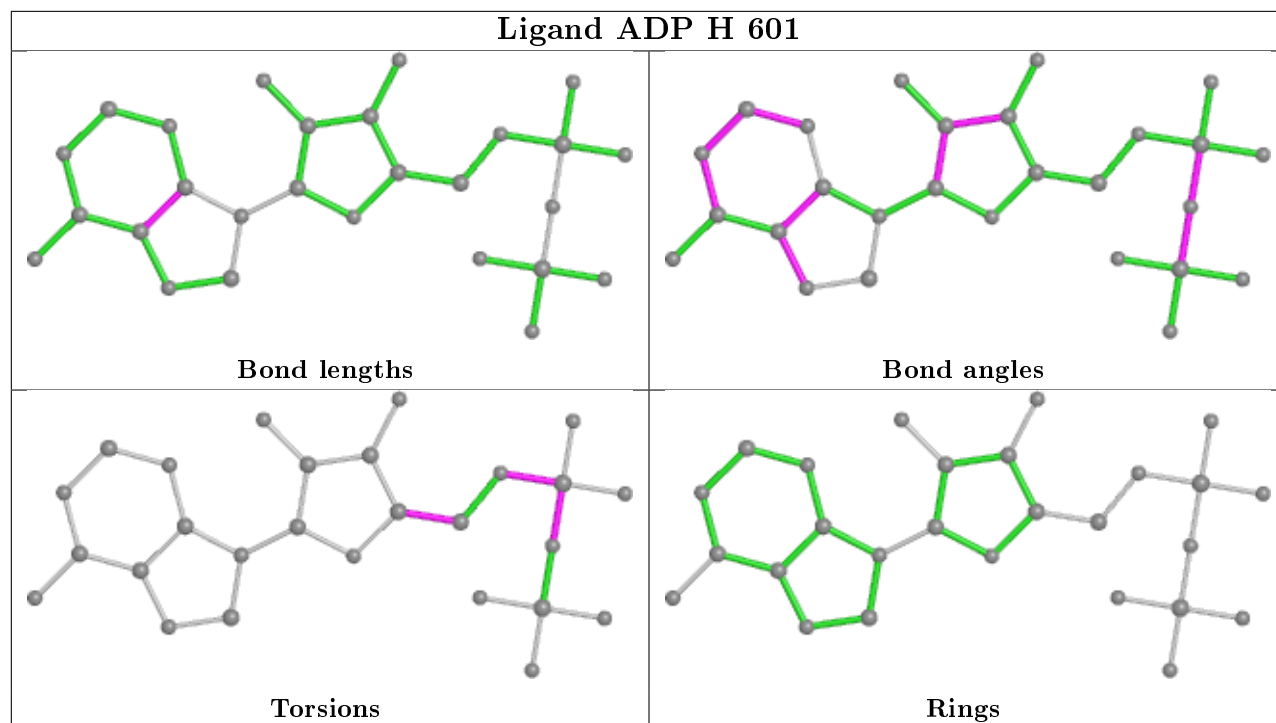
28 monomers are involved in 65 short contacts:

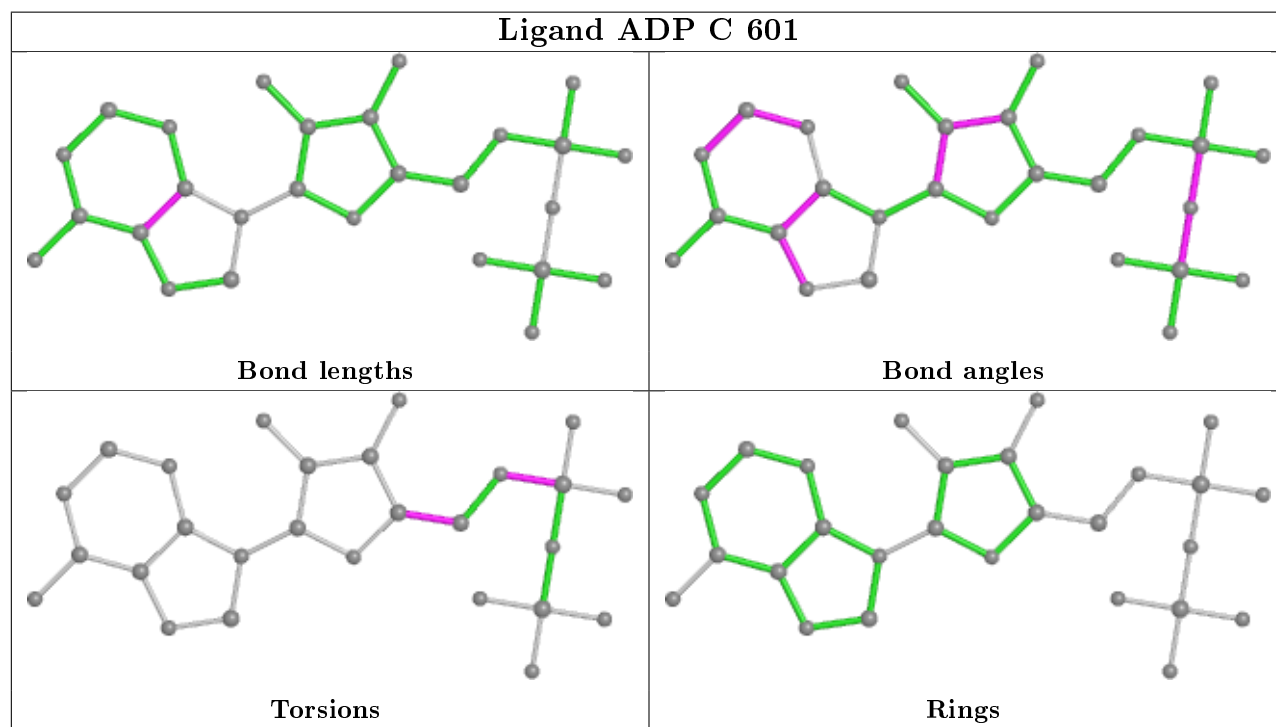
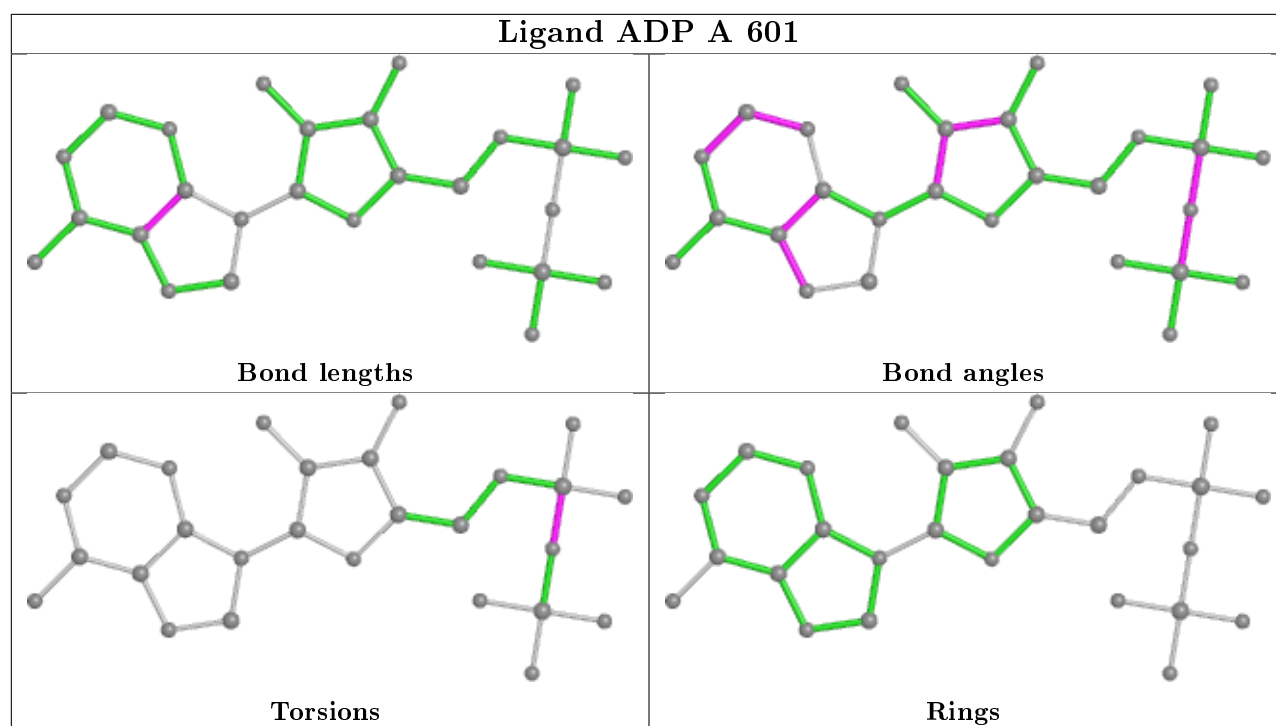
Mol	Chain	Res	Type	Clashes	Symm-Clashes
4	D	602	BEF	1	0
3	B	601	ADP	5	0
4	L	602	BEF	2	0
3	H	601	ADP	2	0
3	J	601	ADP	1	0
4	B	602	BEF	3	0
3	A	601	ADP	1	0
4	I	602	BEF	4	0
4	K	602	BEF	2	0
4	A	602	BEF	2	0
4	J	602	BEF	2	0
3	C	601	ADP	4	0
3	I	601	ADP	3	0
3	K	601	ADP	4	0
3	D	601	ADP	4	0
4	G	602	BEF	1	0
4	N	602	BEF	3	0
3	L	601	ADP	2	0
4	F	602	BEF	1	0
4	E	602	BEF	1	0
4	M	602	BEF	1	0
3	E	601	ADP	4	0
3	M	601	ADP	4	0
3	F	601	ADP	4	0
3	N	601	ADP	5	0
4	H	602	BEF	4	0
3	G	601	ADP	5	0
4	C	602	BEF	4	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

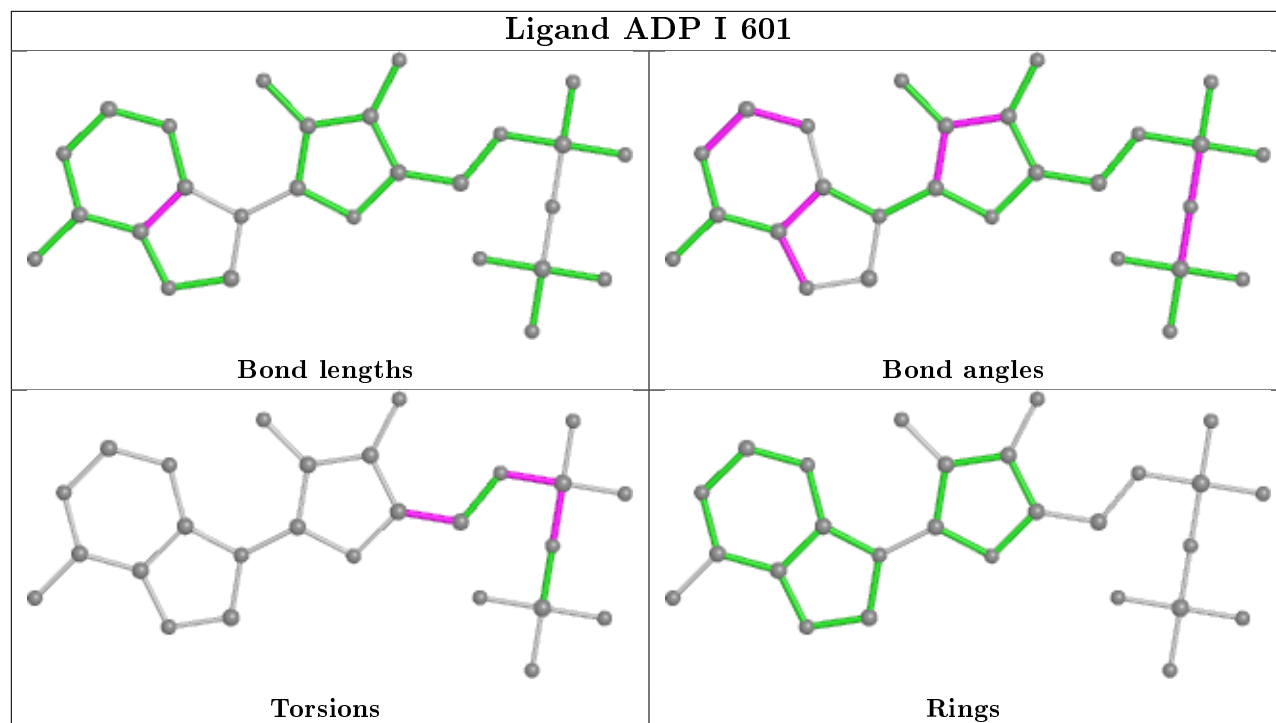
also be included. For torsion angles, if less than 5% of the Mogul distribution of torsion angles is within 10 degrees of the torsion angle in question, then that torsion angle is considered an outlier. Any bond that is central to one or more torsion angles identified as an outlier by Mogul will be highlighted 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.



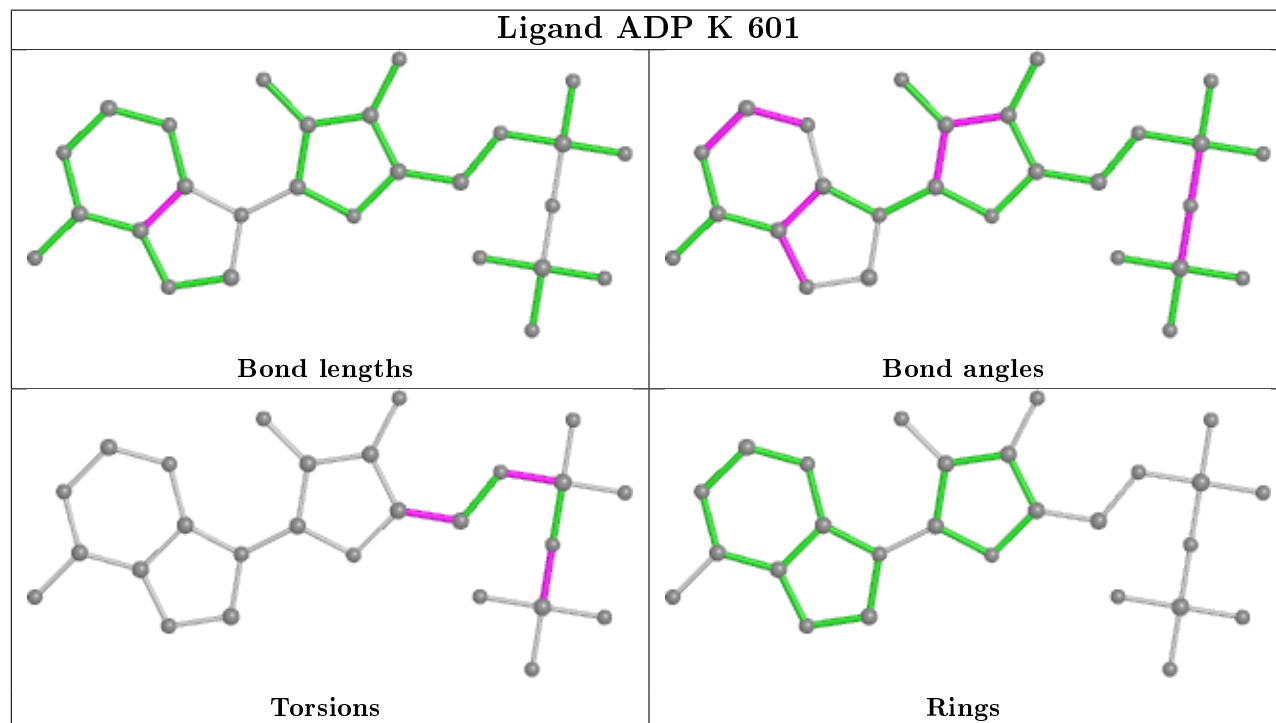


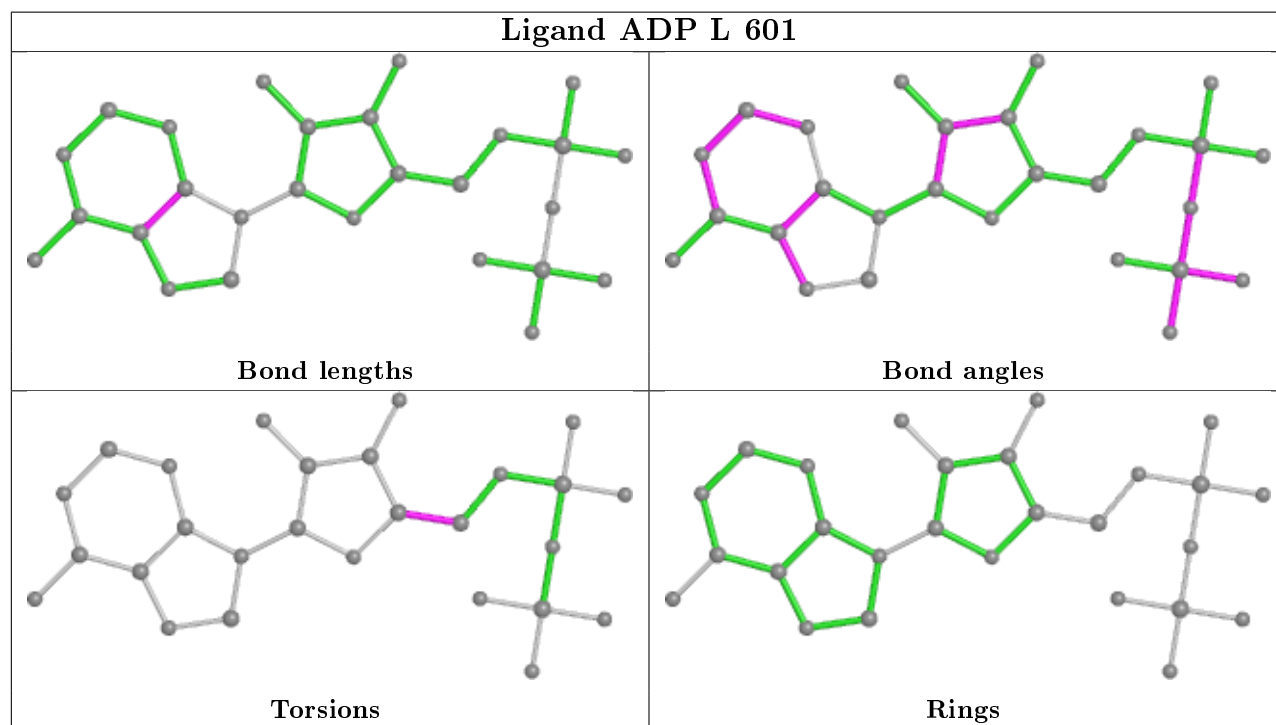
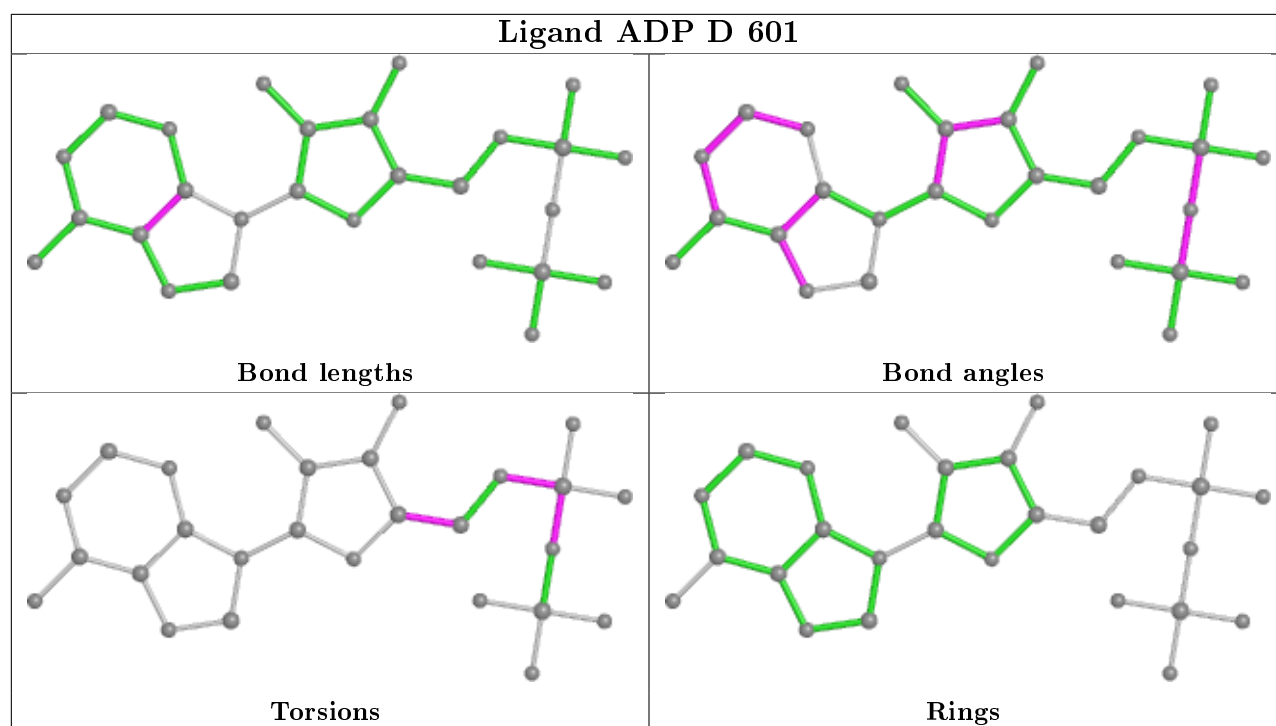


Ligand ADP I 601

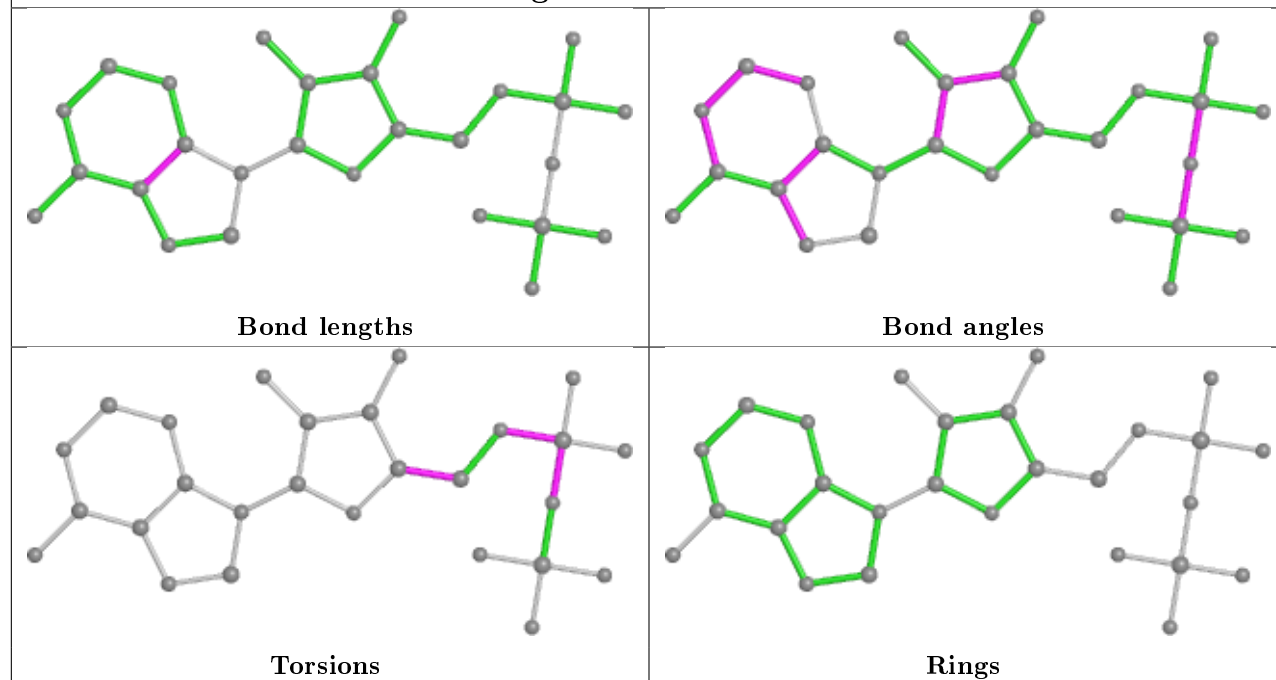


Ligand ADP K 601

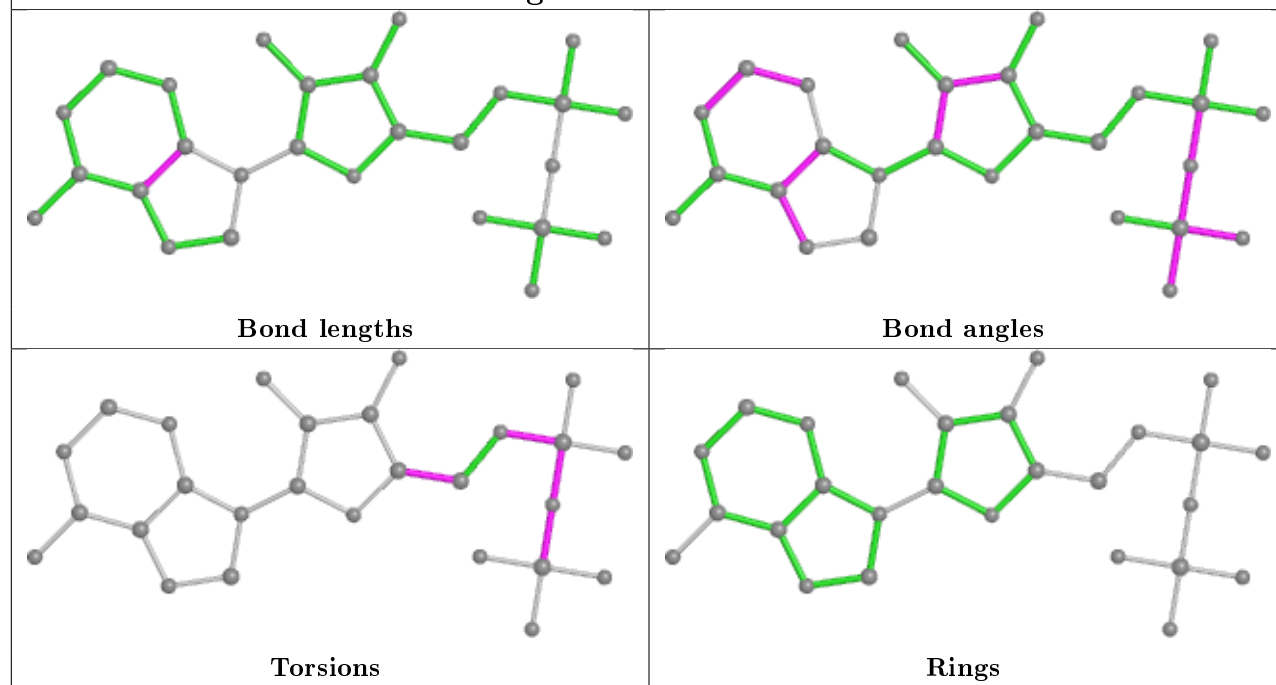


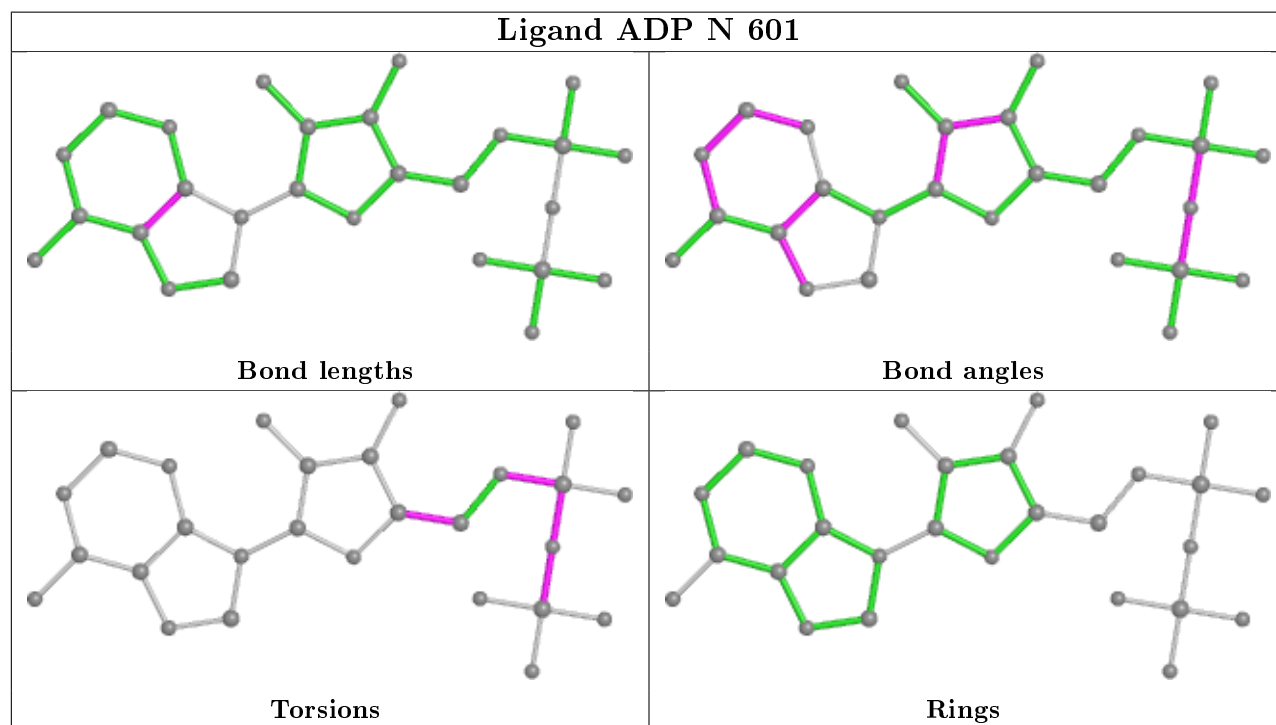
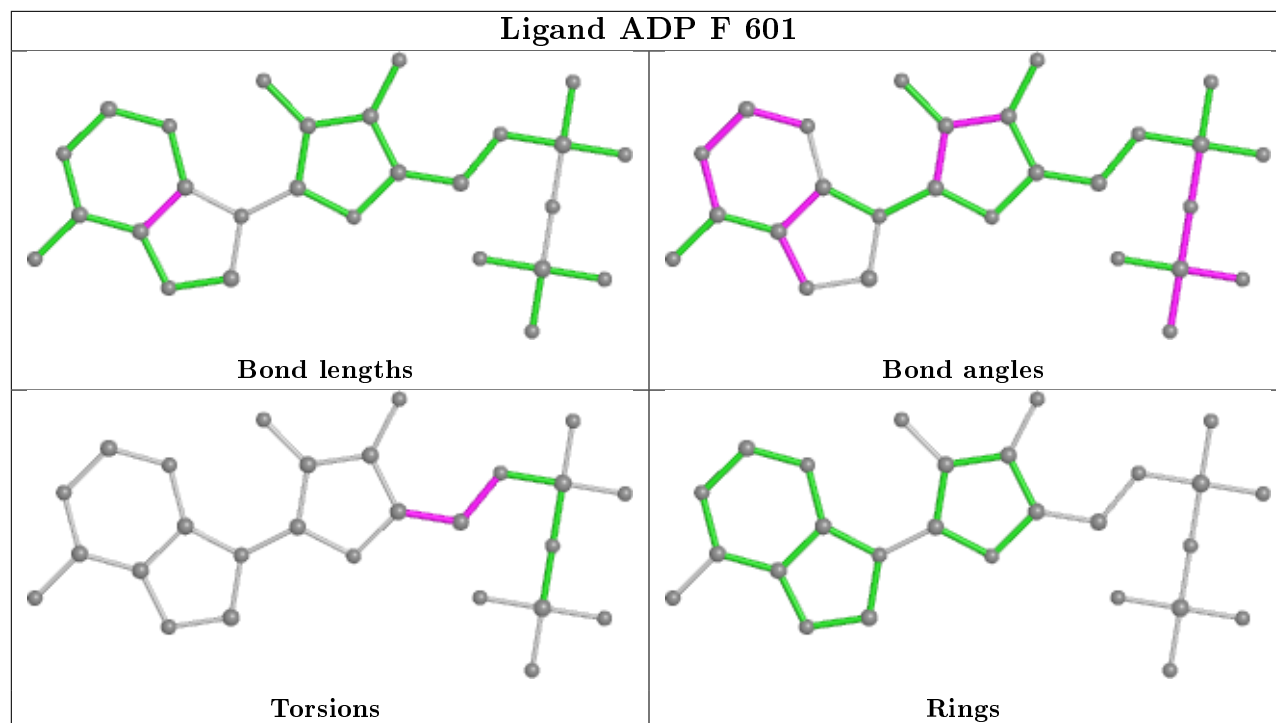


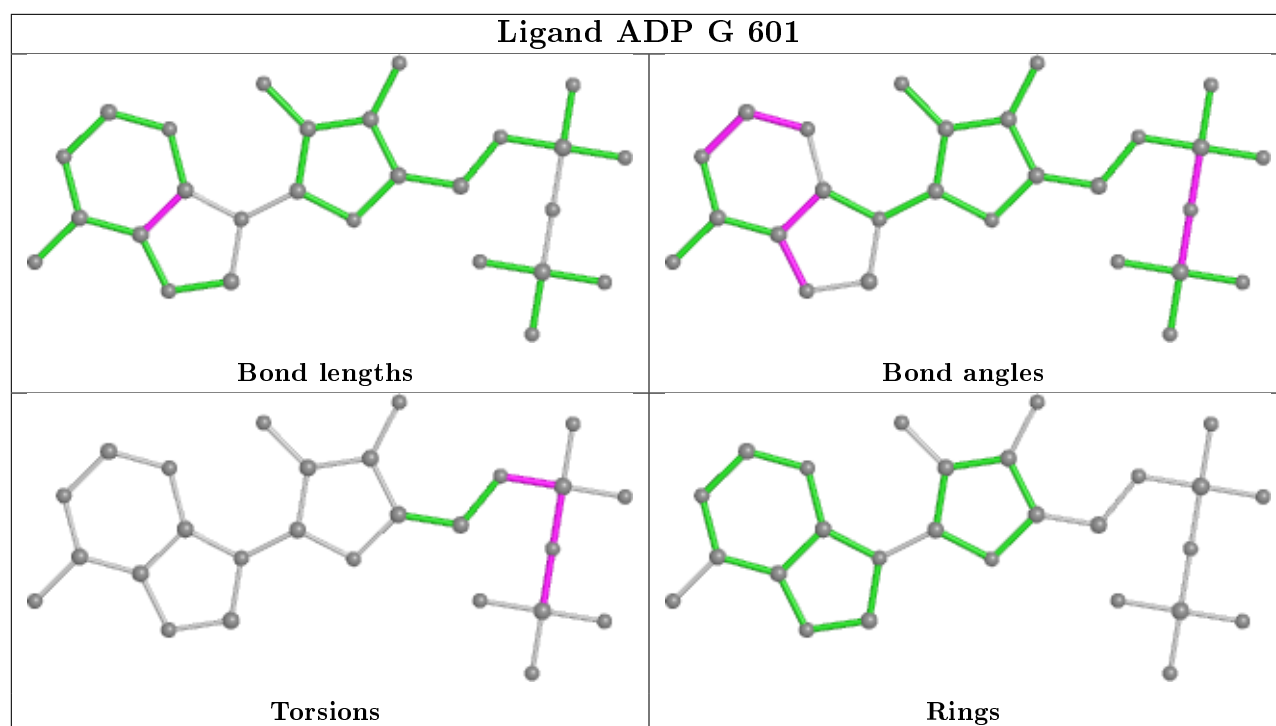
Ligand ADP E 601



Ligand ADP M 601







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 Fit of model and data ⓘ

6.1 Protein, DNA and RNA chains ⓘ

In the following table, the column labelled ‘#RSRZ> 2’ contains the number (and percentage) of RSRZ outliers, followed by percent RSRZ outliers for the chain as percentile scores relative to all X-ray entries and entries of similar resolution. The OWAB column contains the minimum, median, 95th percentile and maximum values of the occupancy-weighted average B-factor per residue. The column labelled ‘Q< 0.9’ lists the number of (and percentage) of residues with an average occupancy less than 0.9.

Mol	Chain	Analysed	<RSRZ>	#RSRZ>2	OWAB(Å ²)	Q<0.9
1	A	524/548 (95%)	-0.17	10 (1%) 66 59	54, 121, 305, 467	0
1	B	524/548 (95%)	-0.22	7 (1%) 77 70	57, 135, 293, 355	0
1	C	524/548 (95%)	-0.26	4 (0%) 86 80	54, 126, 274, 415	0
1	D	524/548 (95%)	-0.32	0 100 100	53, 113, 212, 321	0
1	E	524/548 (95%)	-0.16	12 (2%) 60 52	62, 139, 293, 376	0
1	F	524/548 (95%)	-0.16	16 (3%) 49 39	51, 122, 288, 363	0
1	G	524/548 (95%)	-0.26	2 (0%) 92 88	43, 111, 278, 398	0
1	H	524/548 (95%)	-0.16	10 (1%) 66 59	61, 150, 325, 459	0
1	I	524/548 (95%)	-0.33	0 100 100	49, 117, 213, 354	0
1	J	524/548 (95%)	-0.17	9 (1%) 70 62	63, 139, 287, 398	0
1	K	524/548 (95%)	-0.18	14 (2%) 54 45	59, 141, 293, 379	0
1	L	524/548 (95%)	-0.23	6 (1%) 80 74	53, 130, 298, 397	0
1	M	524/548 (95%)	-0.07	22 (4%) 36 30	71, 146, 330, 456	0
1	N	524/548 (95%)	-0.12	21 (4%) 38 31	70, 159, 306, 375	0
2	1	97/97 (100%)	-0.10	3 (3%) 49 39	93, 149, 321, 352	0
2	2	97/97 (100%)	-0.12	1 (1%) 82 76	98, 162, 314, 350	0
2	O	95/97 (97%)	-0.20	2 (2%) 63 55	88, 136, 315, 390	0
2	P	97/97 (100%)	-0.18	1 (1%) 82 76	74, 152, 323, 361	0
2	Q	96/97 (98%)	-0.03	2 (2%) 63 55	113, 182, 322, 375	0
2	R	97/97 (100%)	-0.15	1 (1%) 82 76	112, 168, 267, 319	0
2	S	96/97 (98%)	0.00	4 (4%) 36 30	106, 165, 341, 367	0
2	T	97/97 (100%)	0.10	6 (6%) 20 16	116, 178, 338, 434	0
2	U	97/97 (100%)	-0.28	0 100 100	88, 156, 354, 422	0
2	V	97/97 (100%)	0.01	3 (3%) 49 39	119, 183, 325, 378	0

Continued on next page...

Continued from previous page...

Mol	Chain	Analysed	<RSRZ>	#RSRZ>2		OWAB(Å ²)	Q<0.9
2	W	97/97 (100%)	-0.15	1 (1%)	82 76	106, 161, 248, 306	0
2	X	97/97 (100%)	-0.07	2 (2%)	63 55	101, 153, 297, 380	0
2	Y	96/97 (98%)	-0.00	5 (5%)	27 23	104, 155, 304, 366	0
2	Z	95/97 (97%)	-0.22	1 (1%)	80 74	108, 160, 312, 358	0
All	All	8687/9030 (96%)	-0.19	165 (1%)	66 59	43, 139, 302, 467	0

All (165) RSRZ outliers are listed below:

Mol	Chain	Res	Type	RSRZ
2	Y	33	ALA	6.3
1	H	229	ASN	5.7
1	M	273	VAL	5.5
1	C	229	ASN	5.3
1	H	359	ASP	5.2
1	K	223	ALA	5.1
1	K	250	ILE	5.0
1	F	275	ALA	4.8
1	N	307	MET	4.7
2	X	18	GLU	4.6
1	M	268	ARG	4.6
2	P	25	ILE	4.5
1	B	246	PRO	4.5
1	F	251	ALA	4.5
2	T	34	LYS	4.4
1	J	356	ALA	4.4
1	K	302	SER	4.3
1	K	251	ALA	4.3
1	N	210	THR	4.2
1	H	360	TYR	4.2
1	K	227	ILE	4.1
2	T	36	THR	4.0
2	1	32	ALA	4.0
1	C	353	ILE	4.0
1	M	243	ALA	3.9
1	N	250	ILE	3.9
1	F	250	ILE	3.8
1	N	248	LEU	3.7
1	B	210	THR	3.7
1	E	356	ALA	3.7
1	F	342	ILE	3.6
2	S	22	ALA	3.5

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type	RSRZ
1	M	192	GLY	3.5
2	V	22	ALA	3.5
1	J	229	ASN	3.4
1	M	242	LYS	3.4
1	E	276	VAL	3.4
2	Y	25	ILE	3.4
1	A	250	ILE	3.4
1	L	306	GLY	3.3
1	A	268	ARG	3.3
1	E	220	ILE	3.3
1	J	275	ALA	3.3
1	K	306	GLY	3.2
2	2	25	ILE	3.2
1	N	273	VAL	3.1
1	M	306	GLY	3.1
1	A	359	ASP	3.1
2	S	23	GLY	3.1
2	O	21	SER	3.1
1	A	350	ARG	3.0
1	F	249	ILE	3.0
2	Y	32	ALA	3.0
1	E	215	LEU	3.0
1	A	366	GLN	3.0
1	M	246	PRO	2.9
1	E	248	LEU	2.9
1	G	306	GLY	2.9
1	N	274	ALA	2.8
2	V	62	GLY	2.8
1	K	333	ILE	2.8
1	A	220	ILE	2.8
1	N	220	ILE	2.8
1	H	228	SER	2.8
1	F	301	ILE	2.8
1	E	221	LEU	2.8
1	M	264	VAL	2.7
1	B	271	VAL	2.7
1	M	238	GLU	2.7
1	H	246	PRO	2.7
1	N	249	ILE	2.7
1	K	353	ILE	2.7
1	N	227	ILE	2.7
1	A	353	ILE	2.7

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type	RSRZ
2	O	25	ILE	2.7
1	E	250	ILE	2.7
1	F	268	ARG	2.6
1	M	250	ILE	2.6
2	Q	18	GLU	2.6
1	N	272	LYS	2.6
1	M	274	ALA	2.6
2	T	25	ILE	2.6
1	J	222	LEU	2.6
2	Y	27	LEU	2.6
1	J	381	VAL	2.6
1	K	174	VAL	2.6
1	K	224	ASP	2.6
2	I	28	THR	2.5
1	H	271	VAL	2.5
1	K	249	ILE	2.5
1	E	275	ALA	2.5
2	T	32	ALA	2.5
1	M	307	MET	2.5
1	F	224	ASP	2.4
1	L	304	GLU	2.4
1	F	223	ALA	2.4
1	H	230	ILE	2.4
1	L	273	VAL	2.4
1	M	265	ASN	2.4
1	F	203	TYR	2.4
1	B	314	LEU	2.4
1	H	247	LEU	2.4
1	M	267	MET	2.4
1	M	353	ILE	2.4
2	X	59	VAL	2.4
1	E	180	GLY	2.4
1	G	222	LEU	2.3
1	M	271	VAL	2.3
2	T	31	ALA	2.3
1	L	333	ILE	2.3
1	L	271	VAL	2.3
1	B	220	ILE	2.3
1	J	177	VAL	2.3
1	N	254	VAL	2.3
2	W	97	ALA	2.3
1	N	221	LEU	2.3

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type	RSRZ
1	N	306	GLY	2.3
1	F	227	ILE	2.3
1	C	203	TYR	2.3
1	M	212	ALA	2.3
2	Z	32	ALA	2.3
1	F	377	ALA	2.3
1	N	385	THR	2.3
1	F	276	VAL	2.2
1	E	222	LEU	2.2
2	S	59	VAL	2.2
1	M	275	ALA	2.2
1	F	274	ALA	2.2
1	M	247	LEU	2.2
2	R	25	ILE	2.2
1	K	229	ASN	2.2
1	K	248	LEU	2.2
2	Y	30	SER	2.2
1	M	175	ILE	2.2
1	B	218	PRO	2.2
1	N	333	ILE	2.2
1	K	301	ILE	2.2
2	T	27	LEU	2.2
2	1	25	ILE	2.1
1	A	273	VAL	2.1
1	M	213	VAL	2.1
1	M	240	VAL	2.1
1	F	248	LEU	2.1
1	N	264	VAL	2.1
1	N	222	LEU	2.1
1	C	228	SER	2.1
1	N	223	ALA	2.1
1	N	314	LEU	2.1
1	E	177	VAL	2.1
2	S	60	LYS	2.1
1	J	247	LEU	2.1
1	J	357	THR	2.1
1	A	360	TYR	2.1
1	J	379	ILE	2.0
1	L	250	ILE	2.0
1	B	222	LEU	2.0
1	H	273	VAL	2.0
1	F	277	LYS	2.0

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type	RSRZ
1	E	175	ILE	2.0
1	N	356	ALA	2.0
2	Q	32	ALA	2.0
2	V	21	SER	2.0
1	H	227	ILE	2.0
1	N	316	ASP	2.0
1	A	357	THR	2.0

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

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

6.3 Carbohydrates [i](#)

There are no carbohydrates in this entry.

6.4 Ligands [i](#)

In the following table, the Atoms column lists the number of modelled atoms in the group and the number defined in the chemical component dictionary. The B-factors column lists the minimum, median, 95th percentile and maximum values of B factors of atoms in the group. The column labelled 'Q<0.9' lists the number of atoms with occupancy less than 0.9.

Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors(Å ²)	Q<0.9
6	K	K	604	1/1	0.94	0.10	110,110,110,110	0
4	BEF	F	602	4/4	0.94	0.16	53,80,165,206	0
6	K	N	604	1/1	0.94	0.10	107,107,107,107	0
4	BEF	I	602	4/4	0.95	0.18	59,65,81,85	0
4	BEF	A	602	4/4	0.95	0.18	62,131,196,204	0
3	ADP	K	601	27/27	0.95	0.24	80,97,118,125	0
3	ADP	H	601	27/27	0.95	0.24	72,93,142,191	0
6	K	A	604	1/1	0.95	0.16	84,84,84,84	0
3	ADP	J	601	27/27	0.95	0.24	35,90,137,139	0
3	ADP	F	601	27/27	0.95	0.24	44,79,128,140	0
3	ADP	A	601	27/27	0.96	0.24	34,80,131,147	0
4	BEF	J	602	4/4	0.96	0.15	40,132,142,204	0
5	MG	G	603	1/1	0.96	0.22	48,48,48,48	0
3	ADP	L	601	27/27	0.96	0.23	69,83,118,128	0
3	ADP	M	601	27/27	0.96	0.21	25,77,196,198	0
6	K	G	604	1/1	0.97	0.15	67,67,67,67	0

Continued on next page...

Continued from previous page...

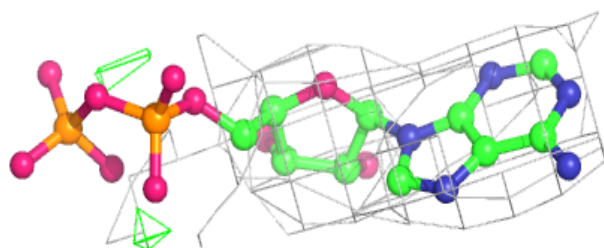
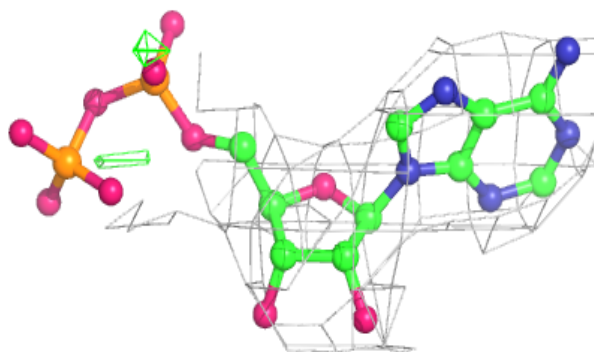
Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors(\AA^2)	Q<0.9
4	BEF	N	602	4/4	0.97	0.15	98,99,105,118	0
4	BEF	K	602	4/4	0.97	0.15	54,86,101,160	0
4	BEF	D	602	4/4	0.97	0.16	31,64,104,109	0
4	BEF	L	602	4/4	0.97	0.14	43,102,141,206	0
3	ADP	C	601	27/27	0.97	0.22	66,83,148,156	0
3	ADP	I	601	27/27	0.97	0.22	47,72,117,158	0
4	BEF	M	602	4/4	0.97	0.19	119,139,180,221	0
6	K	C	604	1/1	0.97	0.10	126,126,126,126	0
6	K	H	604	1/1	0.97	0.07	110,110,110,110	0
3	ADP	N	601	27/27	0.97	0.20	66,106,128,133	0
3	ADP	G	601	27/27	0.97	0.22	3,59,88,94	0
3	ADP	B	601	27/27	0.98	0.23	16,86,98,110	0
4	BEF	B	602	4/4	0.98	0.22	45,60,96,100	0
3	ADP	E	601	27/27	0.98	0.20	36,84,99,115	0
3	ADP	D	601	27/27	0.98	0.22	25,73,84,87	0
6	K	D	604	1/1	0.98	0.11	82,82,82,82	0
4	BEF	G	602	4/4	0.98	0.19	36,46,68,99	0
4	BEF	H	602	4/4	0.98	0.13	13,85,125,156	0
6	K	I	604	1/1	0.98	0.18	75,75,75,75	0
4	BEF	C	602	4/4	0.98	0.20	2,4,92,93	0
4	BEF	E	602	4/4	0.99	0.15	55,87,89,89	0
6	K	L	604	1/1	0.99	0.12	87,87,87,87	0
5	MG	F	603	1/1	0.99	0.23	51,51,51,51	0
5	MG	B	603	1/1	0.99	0.22	66,66,66,66	0
5	MG	I	603	1/1	0.99	0.19	81,81,81,81	0
6	K	M	604	1/1	0.99	0.12	94,94,94,94	0
5	MG	H	603	1/1	0.99	0.16	103,103,103,103	0
6	K	E	604	1/1	0.99	0.14	89,89,89,89	0
6	K	B	604	1/1	0.99	0.11	79,79,79,79	0
5	MG	A	603	1/1	0.99	0.23	78,78,78,78	0
5	MG	N	603	1/1	0.99	0.15	71,71,71,71	0
5	MG	D	603	1/1	0.99	0.20	44,44,44,44	0
6	K	F	604	1/1	0.99	0.16	103,103,103,103	0
6	K	J	604	1/1	0.99	0.17	122,122,122,122	0
5	MG	C	603	1/1	0.99	0.17	57,57,57,57	0
5	MG	M	603	1/1	0.99	0.17	73,73,73,73	0
5	MG	K	603	1/1	0.99	0.21	50,50,50,50	0
5	MG	E	603	1/1	1.00	0.16	45,45,45,45	0
5	MG	L	603	1/1	1.00	0.20	66,66,66,66	0
5	MG	J	603	1/1	1.00	0.18	56,56,56,56	0

The following is a graphical depiction of the model fit to experimental electron density of all instances of the Ligand of Interest. In addition, ligands with molecular weight > 250 and outliers

as shown on the geometry validation Tables will also be included. Each fit is shown from different orientation to approximate a three-dimensional view.

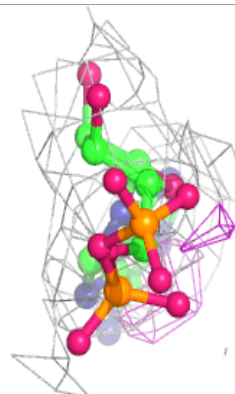
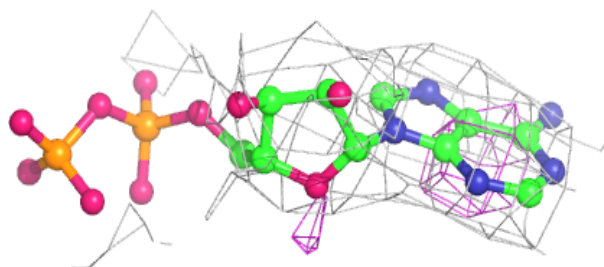
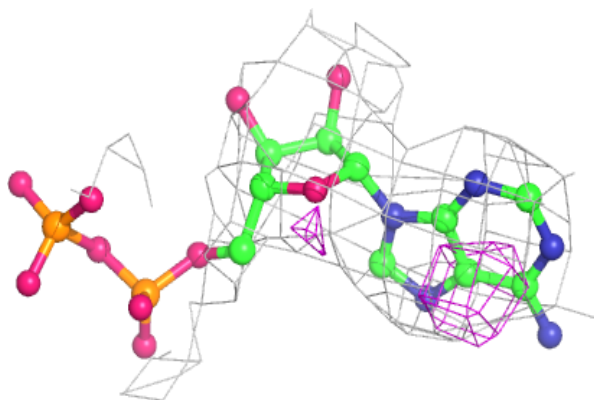
Electron density around ADP K 601:

$2mF_o-DF_c$ (at 0.7 rmsd) in gray
 mF_o-DF_c (at 3 rmsd) in purple (negative)
and green (positive)



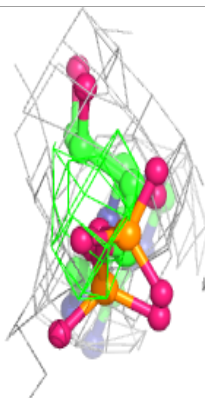
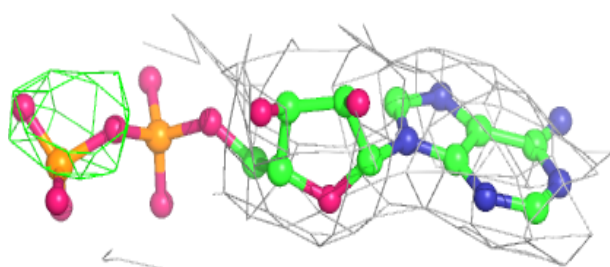
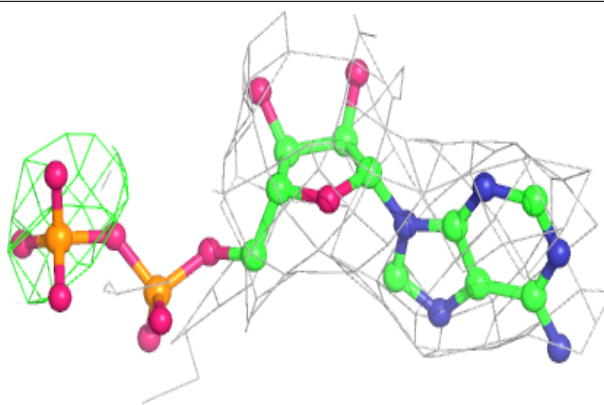
Electron density around ADP H 601:

$2mF_o-DF_c$ (at 0.7 rmsd) in gray
 mF_o-DF_c (at 3 rmsd) in purple (negative)
and green (positive)

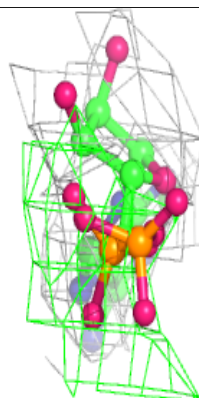
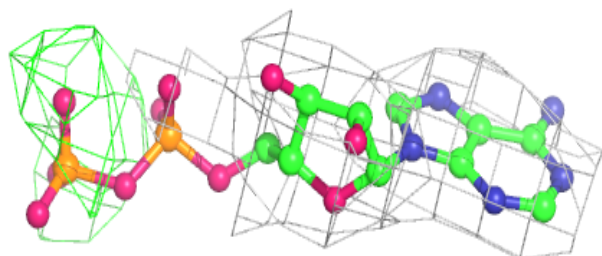
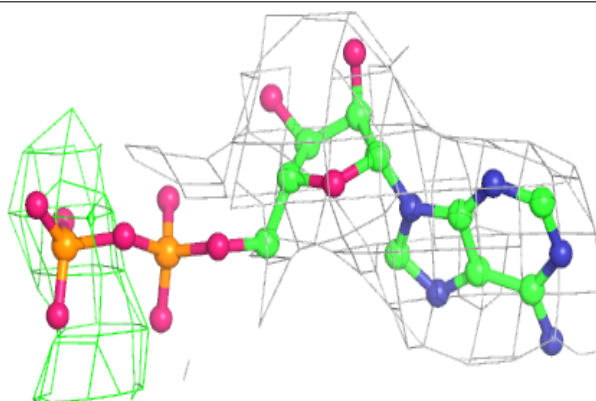


Electron density around ADP J 601:

$2mF_o-DF_c$ (at 0.7 rmsd) in gray
 mF_o-DF_c (at 3 rmsd) in purple (negative)
and green (positive)

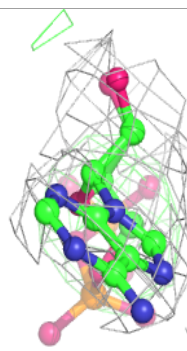
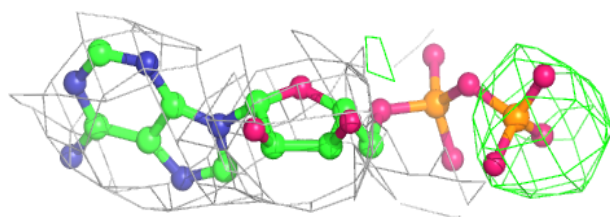
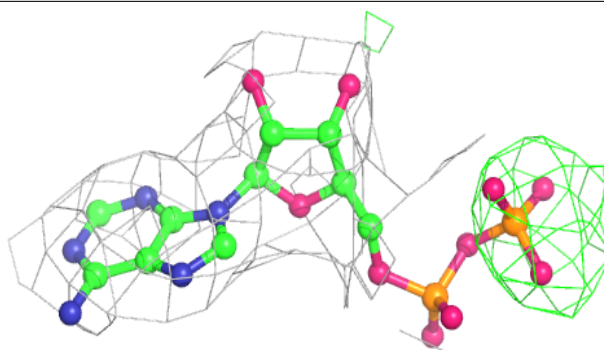
**Electron density around ADP F 601:**

$2mF_o-DF_c$ (at 0.7 rmsd) in gray
 mF_o-DF_c (at 3 rmsd) in purple (negative)
and green (positive)

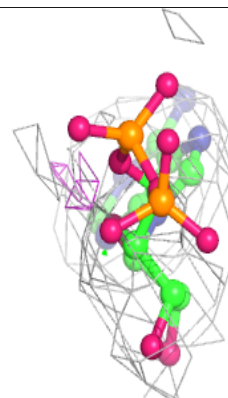
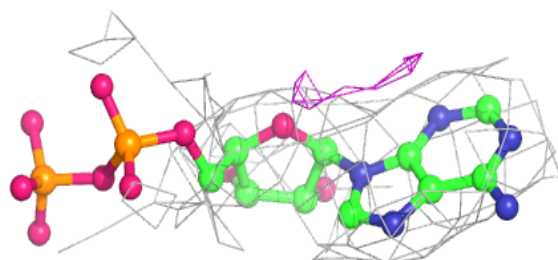
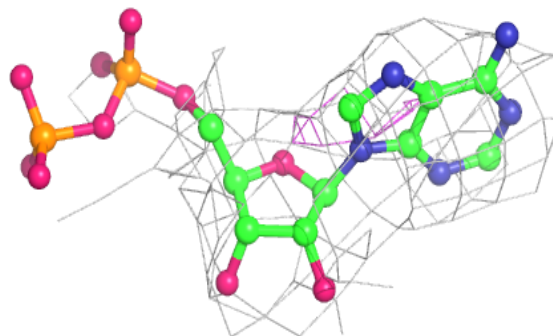


Electron density around ADP A 601:

$2mF_o-DF_c$ (at 0.7 rmsd) in gray
 mF_o-DF_c (at 3 rmsd) in purple (negative)
and green (positive)

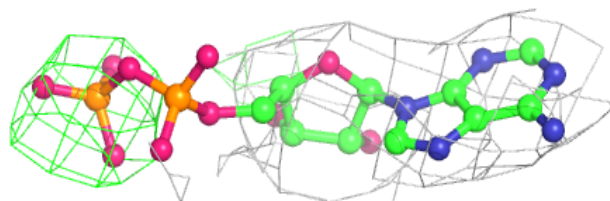
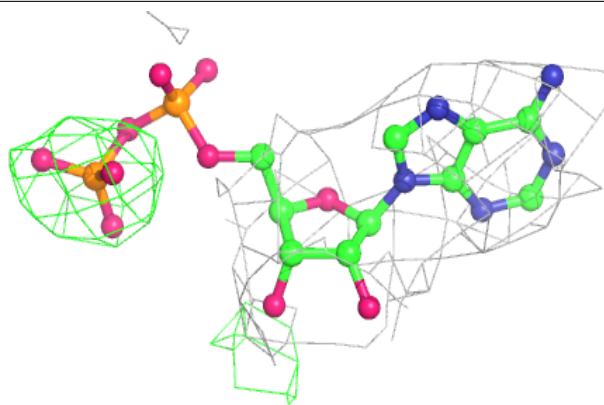
**Electron density around ADP L 601:**

$2mF_o-DF_c$ (at 0.7 rmsd) in gray
 mF_o-DF_c (at 3 rmsd) in purple (negative)
and green (positive)

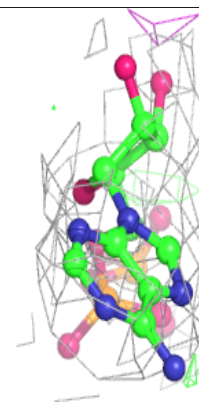
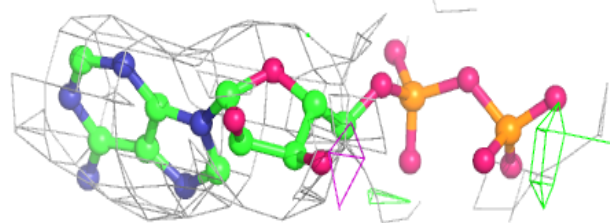
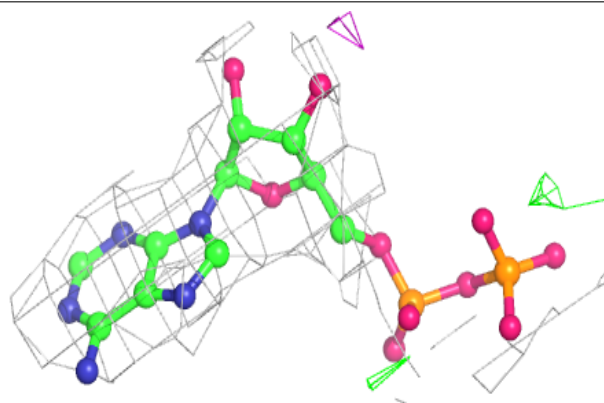


Electron density around ADP M 601:

$2mF_o - DF_c$ (at 0.7 rmsd) in gray
 $mF_o - DF_c$ (at 3 rmsd) in purple (negative)
and green (positive)

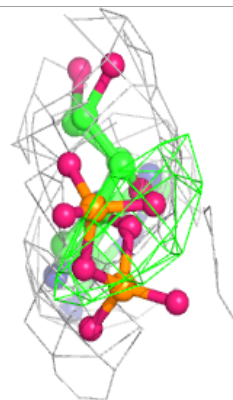
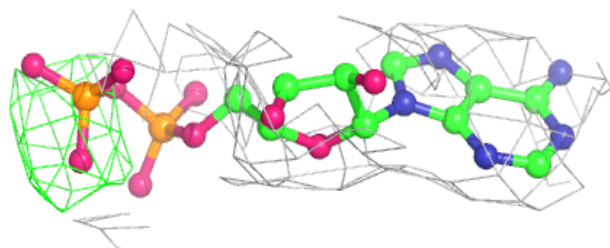
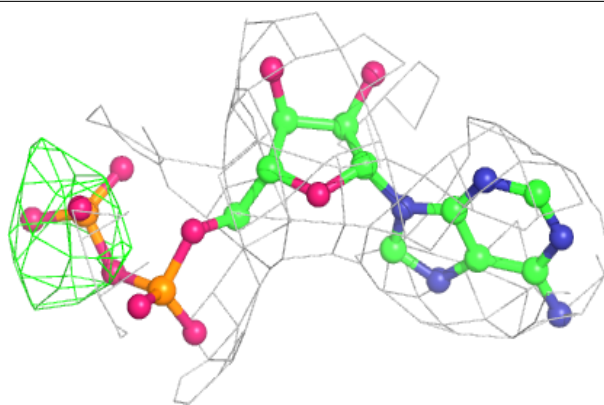
**Electron density around ADP C 601:**

$2mF_o - DF_c$ (at 0.7 rmsd) in gray
 $mF_o - DF_c$ (at 3 rmsd) in purple (negative)
and green (positive)

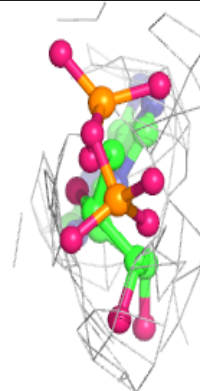
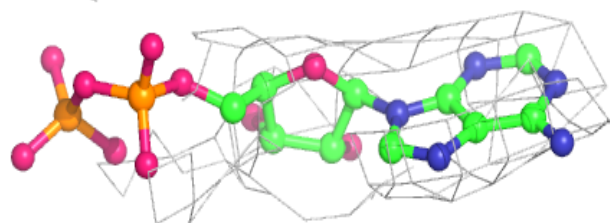
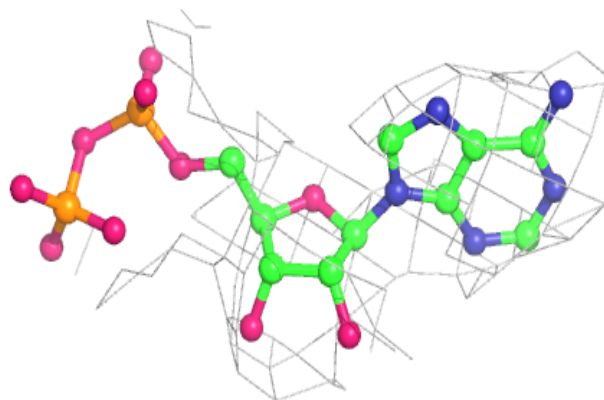


Electron density around ADP I 601:

$2mF_o-DF_c$ (at 0.7 rmsd) in gray
 mF_o-DF_c (at 3 rmsd) in purple (negative)
and green (positive)

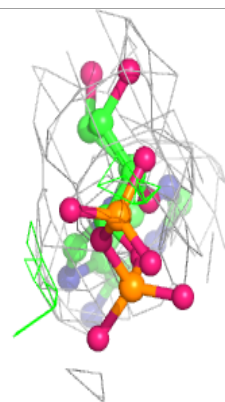
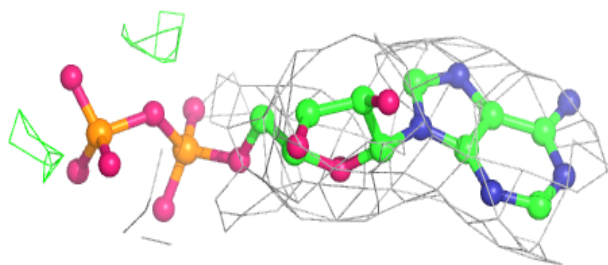
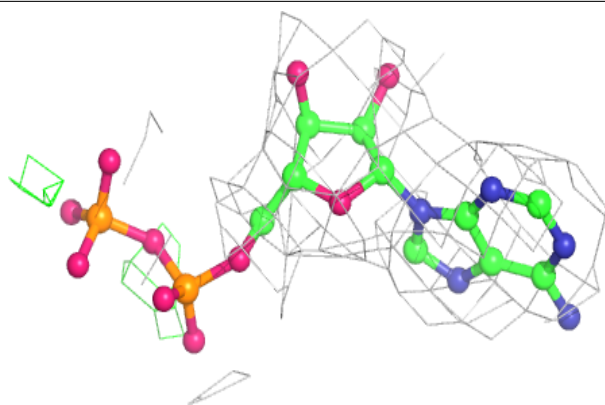
**Electron density around ADP N 601:**

$2mF_o-DF_c$ (at 0.7 rmsd) in gray
 mF_o-DF_c (at 3 rmsd) in purple (negative)
and green (positive)

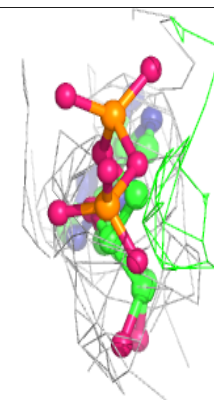
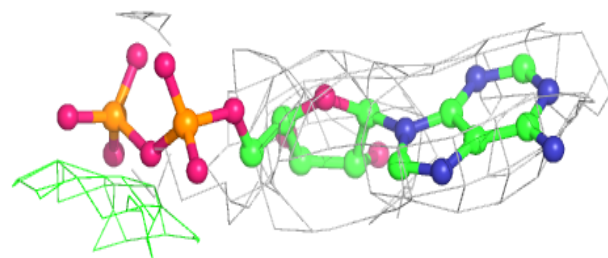
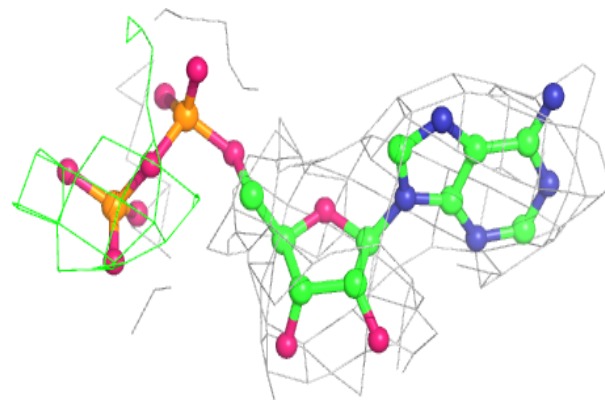


Electron density around ADP G 601:

$2mF_o-DF_c$ (at 0.7 rmsd) in gray
 mF_o-DF_c (at 3 rmsd) in purple (negative)
and green (positive)

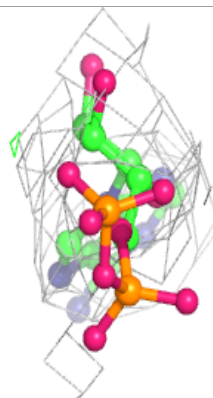
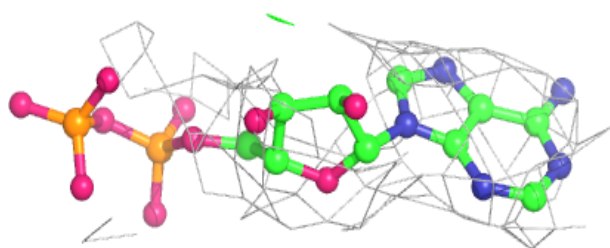
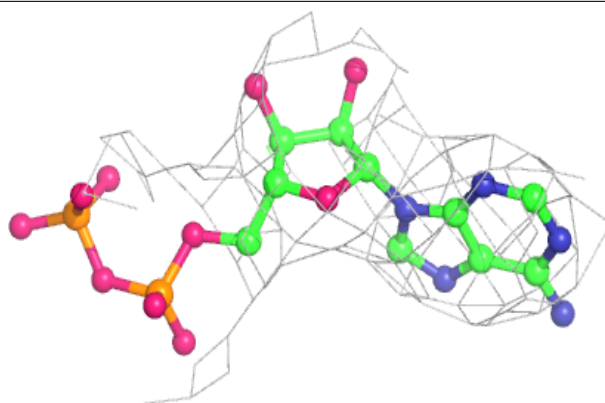
**Electron density around ADP B 601:**

$2mF_o-DF_c$ (at 0.7 rmsd) in gray
 mF_o-DF_c (at 3 rmsd) in purple (negative)
and green (positive)

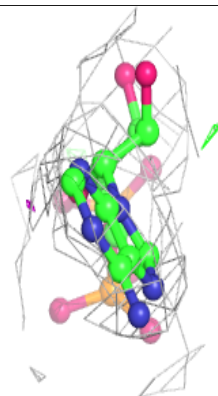
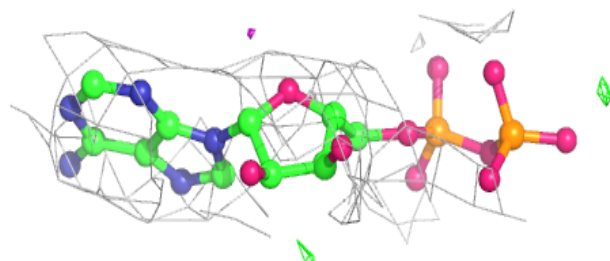
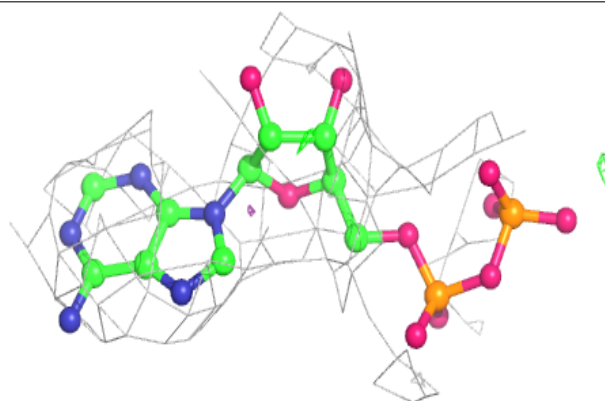


Electron density around ADP E 601:

$2mF_o-DF_c$ (at 0.7 rmsd) in gray
 mF_o-DF_c (at 3 rmsd) in purple (negative)
and green (positive)

**Electron density around ADP D 601:**

$2mF_o-DF_c$ (at 0.7 rmsd) in gray
 mF_o-DF_c (at 3 rmsd) in purple (negative)
and green (positive)



6.5 Other polymers [i](#)

There are no such residues in this entry.