Full wwPDB EM Validation Report ⓘ

Mar 30, 2026 – 08:52 AM UTC

PDB ID	:	5ZVT / pdb_00005zvt
EMDB ID	:	EMD-6969
Title	:	Structure of RNA polymerase complex and genome within a dsRNA virus provides insights into the mechanisms of transcription and assembly
Authors	:	Liu, H.; Fang, Q.; Cheng, L.
Deposited on	:	2018-05-12
Resolution	:	3.30 Å(reported)

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

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

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

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

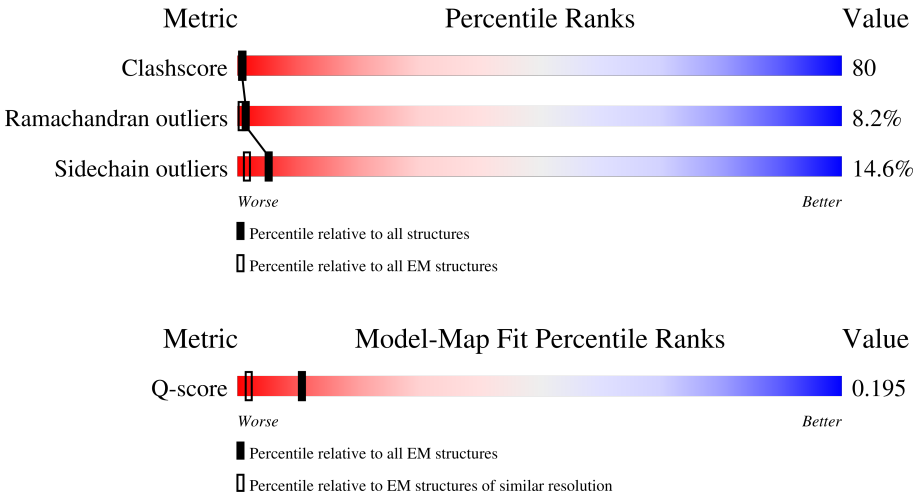
EMDB validation analysis	:	0.0.1.dev132
Mogul	:	2022.3.0, CSD as543be (2022)
MolProbity	:	4-5-2 with Phenix2.0
Percentile statistics	:	20250101.v01 (using entries in the PDB archive January 1st 2025)
EM percentile statistics	:	202505.v01 (Using data in the EMDB archive up until May 2025)
MapQ	:	1.9.13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

1 Overall quality at a glance

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

The reported resolution of this entry is 3.30 Å.

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



Metric	Whole archive (#Entries)	EM structures (#Entries)	Similar EM resolution (#Entries, resolution range(Å))
Clashscore	229148	23984	-
Ramachandran outliers	224038	23583	-
Sidechain outliers	223484	23102	-
Q-score	-	25397	15087 (2.80 - 3.80)

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

Mol	Chain	Length	Quality of chain
1	b	276	8% 14% 13% 69%
1	d	276	13% 15% 69%
1	f	276	5% 14% 13% 69%
1	h	276	6% 16% 12% 69%

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Mol	Chain	Length	Quality of chain
1	j	276	5% 14% 13% 69%
1	l	276	6% 14% 14% 69%
1	n	276	17% 12% 69%
1	p	276	8% 16% 11% 69%
1	r	276	7% 17% 11% 69%
1	t	276	12% 14% 14% 69%
2	A	42	38% 29% 36% 29% 5%
2	C	42	33% 19% 43% 31% 5%
2	E	42	31% 19% 40% 33% 5%
2	G	42	33% 33% 43% 17% 5%
2	I	42	36% 19% 43% 31% 5%
2	K	42	33% 19% 40% 33% 5%
2	M	42	36% 33% 43% 17% 5%
2	O	42	31% 19% 43% 31% 5%
2	Q	42	36% 17% 43% 33% 5%
2	S	42	26% 33% 43% 17% 5%
3	B	606	42% 24% 53% 18%
3	D	606	32% 20% 57% 18%
3	F	606	31% 20% 56% 19%
3	H	606	35% 20% 57% 19%
3	J	606	36% 20% 56% 19%
3	L	606	35% 21% 56% 18%
3	N	606	36% 19% 57% 19%
3	P	606	33% 20% 56% 19%
3	R	606	35% 21% 56% 19%

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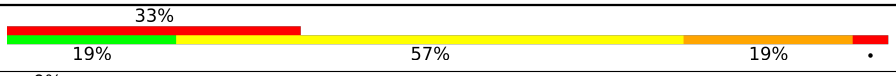
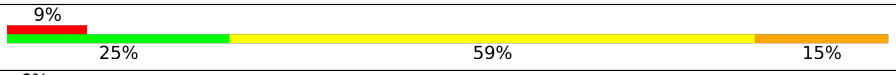
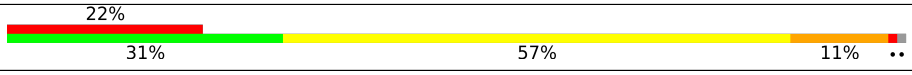
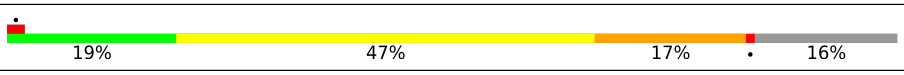
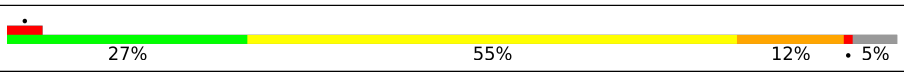
Mol	Chain	Length	Quality of chain
3	T	606	
4	U	412	
4	V	412	
5	W	1299	
6	X	1214	
6	Y	1214	

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
7	MYR	A	101	-	-	X	-
7	MYR	C	101	-	-	X	-
7	MYR	E	101	-	-	X	-
7	MYR	G	101	-	-	X	-
7	MYR	I	101	-	-	X	-
7	MYR	K	101	-	-	X	-
7	MYR	M	101	-	-	X	-
7	MYR	O	101	-	-	X	-
7	MYR	Q	101	-	-	X	-
7	MYR	S	101	-	-	X	-

2 Entry composition (i)

There are 7 unique types of molecules in this entry. The entry contains 87645 atoms, of which 0 are hydrogens and 0 are deuteriums.

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

- Molecule 1 is a protein called Outer capsid VP7.

Mol	Chain	Residues	Atoms					AltConf	Trace
1	l	86	Total	C	N	O	S	0	0
			666	411	125	124	6		
1	b	86	Total	C	N	O	S	0	0
			666	411	125	124	6		
1	f	86	Total	C	N	O	S	0	0
			666	411	125	124	6		
1	d	86	Total	C	N	O	S	0	0
			666	411	125	124	6		
1	h	86	Total	C	N	O	S	0	0
			666	411	125	124	6		
1	j	86	Total	C	N	O	S	0	0
			666	411	125	124	6		
1	n	86	Total	C	N	O	S	0	0
			666	411	125	124	6		
1	p	86	Total	C	N	O	S	0	0
			666	411	125	124	6		
1	r	86	Total	C	N	O	S	0	0
			666	411	125	124	6		
1	t	86	Total	C	N	O	S	0	0
			666	411	125	124	6		

- Molecule 2 is a protein called N-terminus of outer capsid protein VP5.

Mol	Chain	Residues	Atoms					AltConf	Trace
2	A	41	Total	C	N	O	S	0	0
			291	177	48	65	1		
2	C	41	Total	C	N	O	S	0	0
			291	177	48	65	1		
2	E	41	Total	C	N	O	S	0	0
			291	177	48	65	1		
2	G	41	Total	C	N	O	S	0	0
			284	174	46	63	1		
2	I	41	Total	C	N	O	S	0	0
			291	177	48	65	1		

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Mol	Chain	Residues	Atoms					AltConf	Trace
2	K	41	Total	C	N	O	S	0	0
			291	177	48	65	1		
2	M	41	Total	C	N	O	S	0	0
			284	174	46	63	1		
2	O	41	Total	C	N	O	S	0	0
			291	177	48	65	1		
2	Q	41	Total	C	N	O	S	0	0
			291	177	48	65	1		
2	S	41	Total	C	N	O	S	0	0
			284	174	46	63	1		

- Molecule 3 is a protein called C-terminus of outer capsid protein VP5.

Mol	Chain	Residues	Atoms					AltConf	Trace
3	B	604	Total	C	N	O	S	0	0
			4508	2858	761	872	17		
3	D	604	Total	C	N	O	S	0	0
			4508	2858	761	872	17		
3	F	604	Total	C	N	O	S	0	0
			4508	2858	761	872	17		
3	H	604	Total	C	N	O	S	0	0
			4508	2858	761	872	17		
3	J	604	Total	C	N	O	S	0	0
			4508	2858	761	872	17		
3	L	604	Total	C	N	O	S	0	0
			4508	2858	761	872	17		
3	N	604	Total	C	N	O	S	0	0
			4508	2858	761	872	17		
3	P	604	Total	C	N	O	S	0	0
			4508	2858	761	872	17		
3	R	604	Total	C	N	O	S	0	0
			4508	2858	761	872	17		
3	T	604	Total	C	N	O	S	0	0
			4508	2858	761	872	17		

- Molecule 4 is a protein called Core protein VP6.

Mol	Chain	Residues	Atoms					AltConf	Trace
4	U	411	Total	C	N	O	S	0	0
			3138	2008	544	571	15		
4	V	411	Total	C	N	O	S	0	0
			3138	2008	544	571	15		

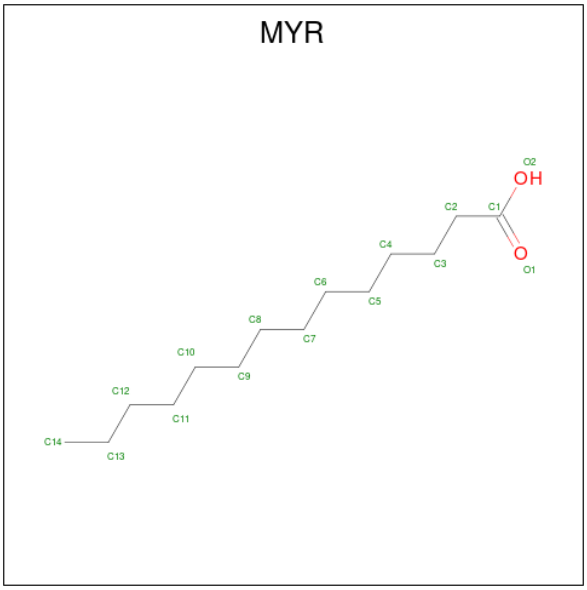
- Molecule 5 is a protein called VP1.

Mol	Chain	Residues	Atoms					AltConf	Trace
5	W	1284	Total	C	N	O	S	0	0
			9882	6335	1681	1839	27		

- Molecule 6 is a protein called VP3.

Mol	Chain	Residues	Atoms					AltConf	Trace
6	X	1018	Total	C	N	O	S	0	0
			7873	5033	1347	1447	46		
6	Y	1154	Total	C	N	O	S	0	0
			8835	5604	1525	1656	50		

- Molecule 7 is MYRISTIC ACID (CCD ID: MYR) (formula: $C_{14}H_{28}O_2$).



Mol	Chain	Residues	Atoms				AltConf
7	A	1	Total	C	O		0
			15	14	1		
7	C	1	Total	C	O		0
			15	14	1		
7	E	1	Total	C	O		0
			15	14	1		
7	G	1	Total	C	O		0
			15	14	1		
7	I	1	Total	C	O		0
			15	14	1		
7	K	1	Total	C	O		0
			15	14	1		

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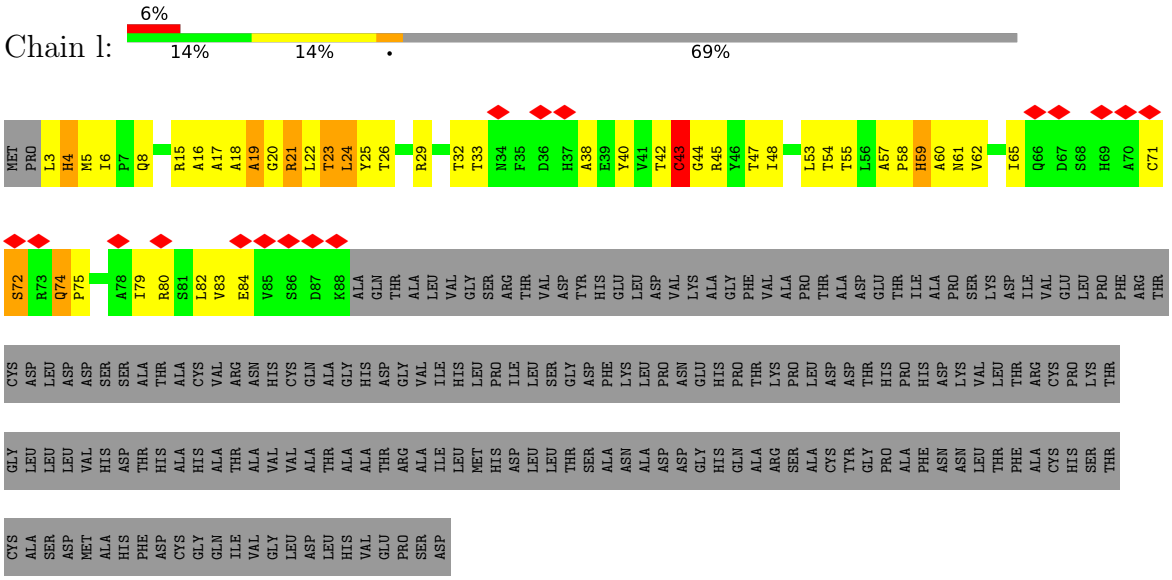
Mol	Chain	Residues	Atoms			AltConf
7	M	1	Total	C	O	0
			15	14	1	
7	O	1	Total	C	O	0
			15	14	1	
7	Q	1	Total	C	O	0
			15	14	1	
7	S	1	Total	C	O	0
			15	14	1	

3 Residue-property plots (i)

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

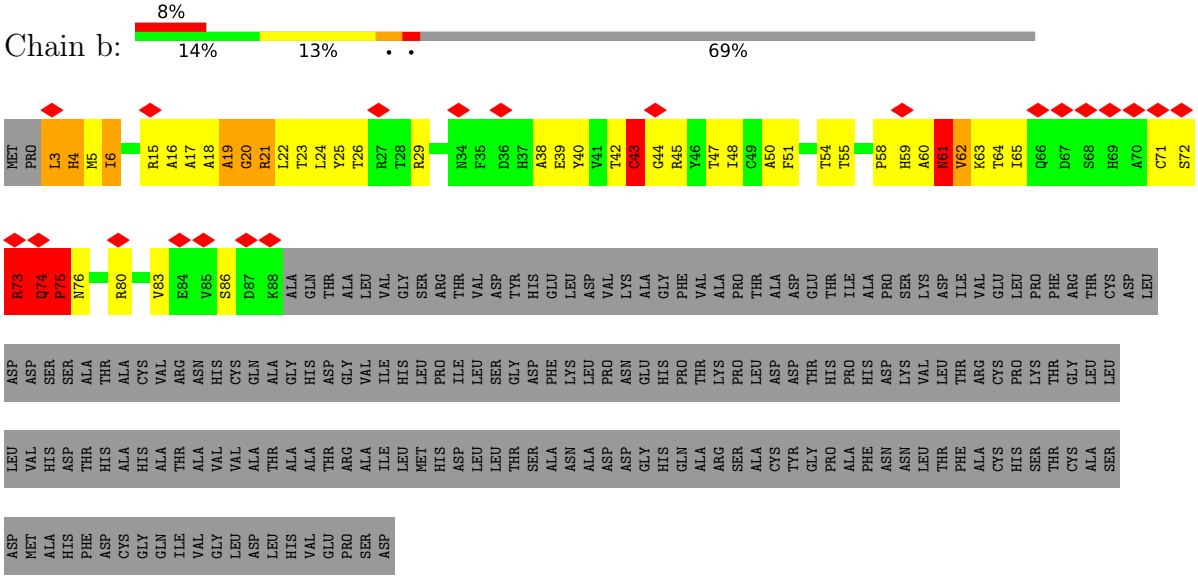
• Molecule 1: Outer capsid VP7

Chain l:



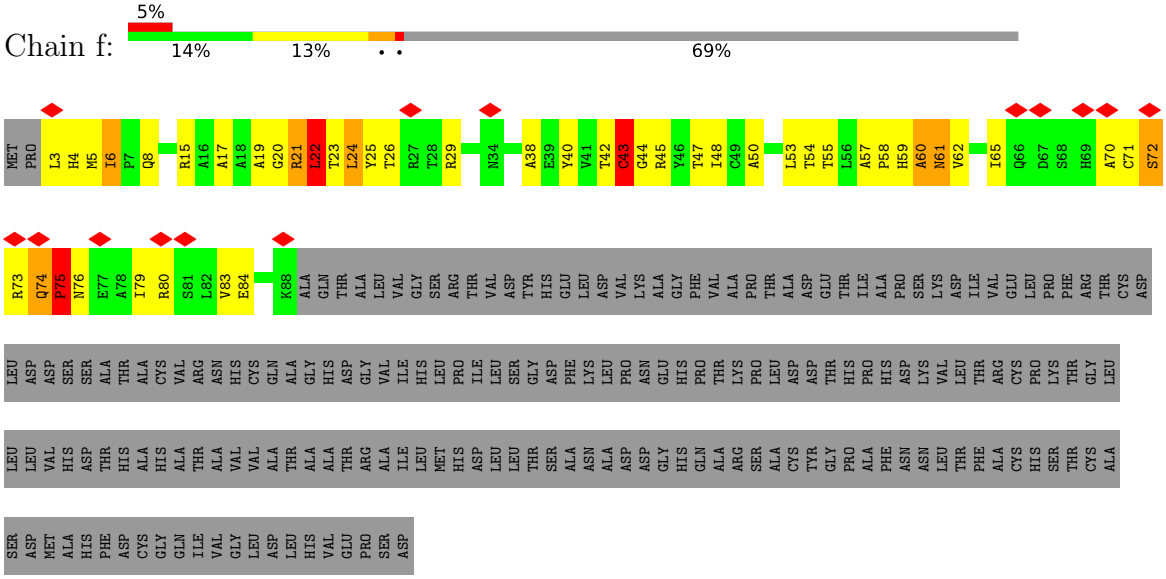
• Molecule 1: Outer capsid VP7

Chain b:



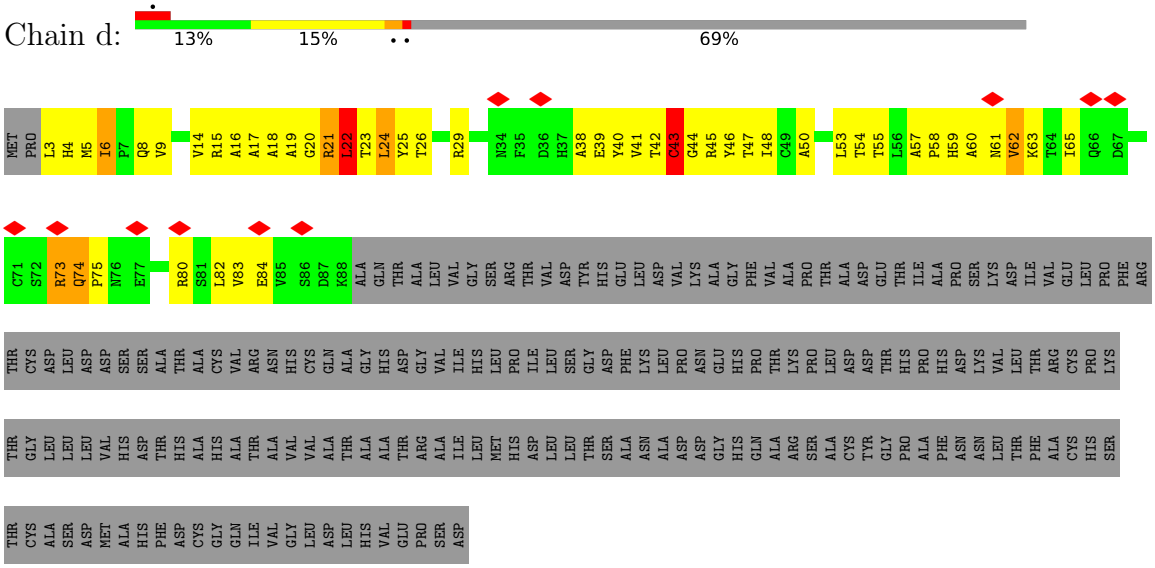
• Molecule 1: Outer capsid VP7

Chain f:



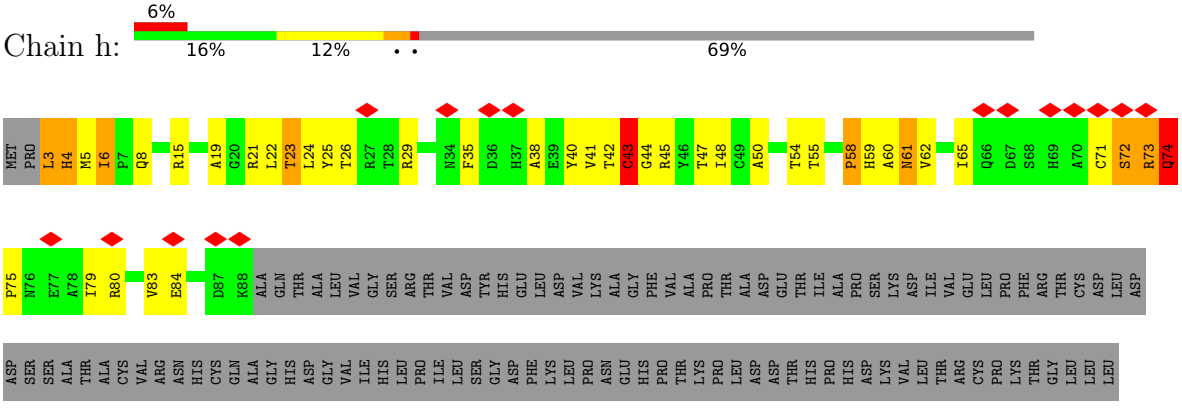
• Molecule 1: Outer capsid VP7

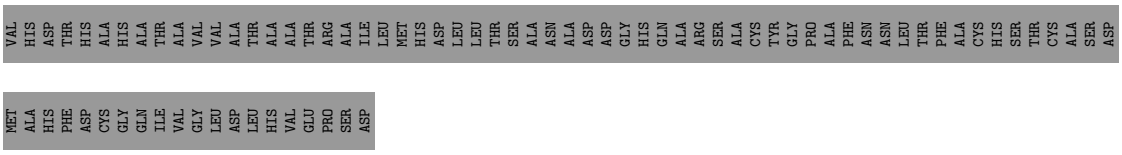
Chain d:



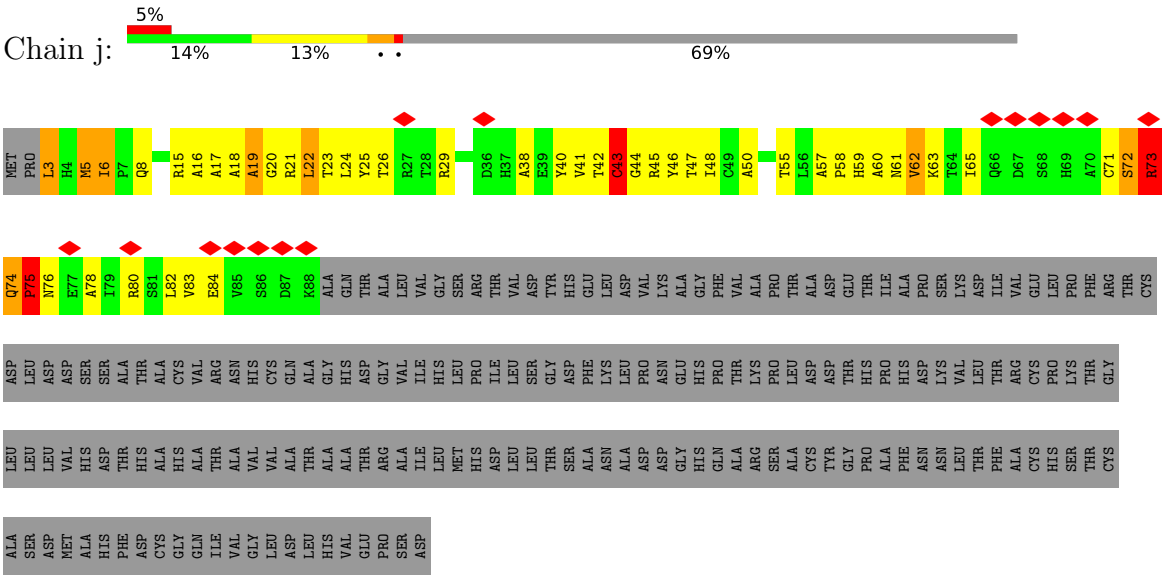
• Molecule 1: Outer capsid VP7

Chain h:

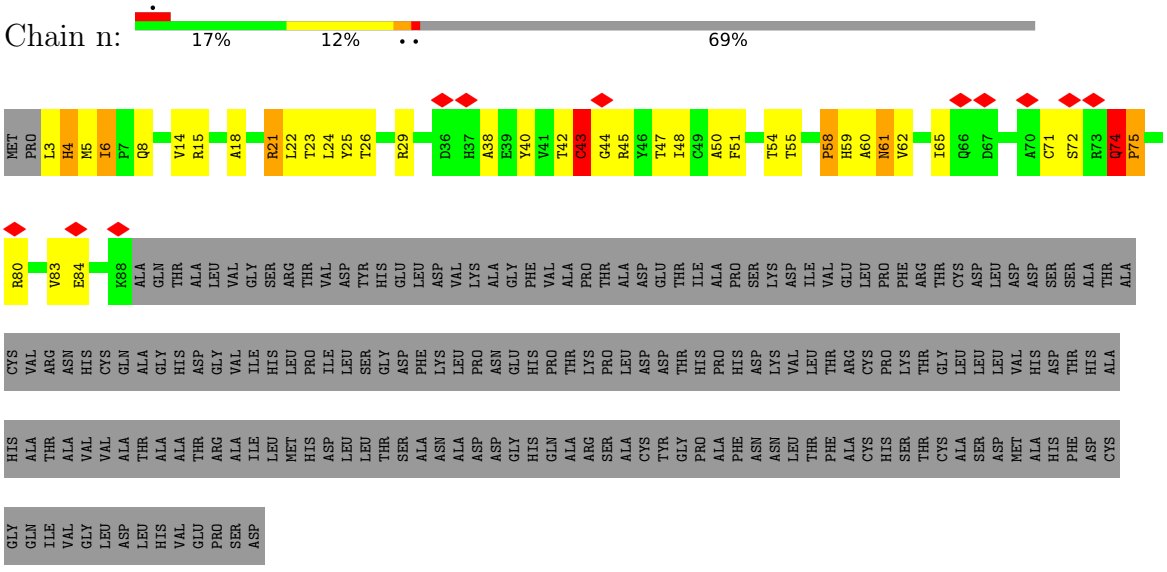




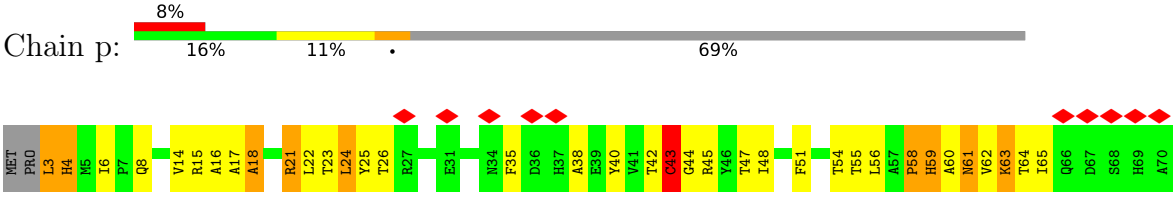
• Molecule 1: Outer capsid VP7

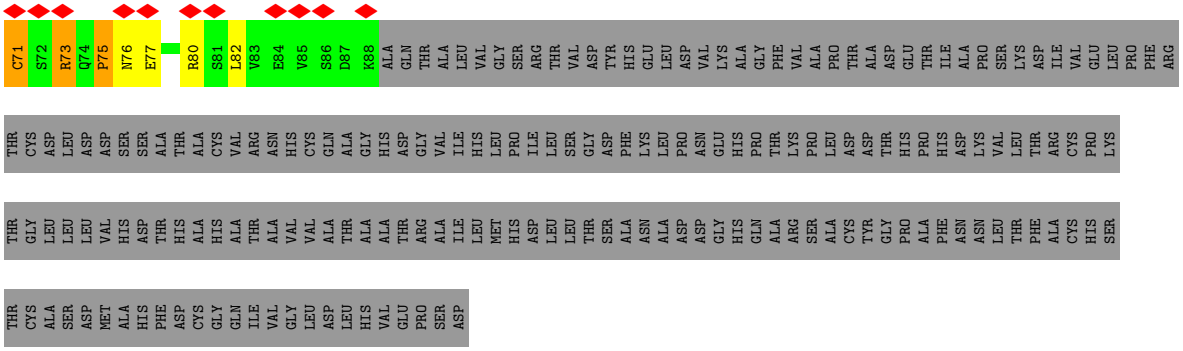


• Molecule 1: Outer capsid VP7



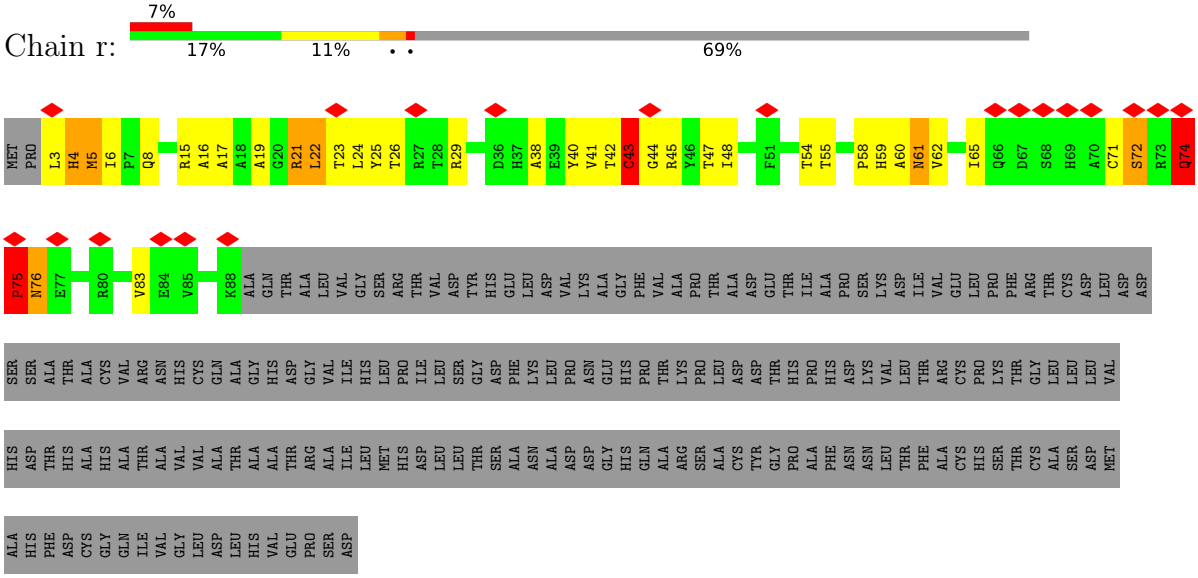
• Molecule 1: Outer capsid VP7





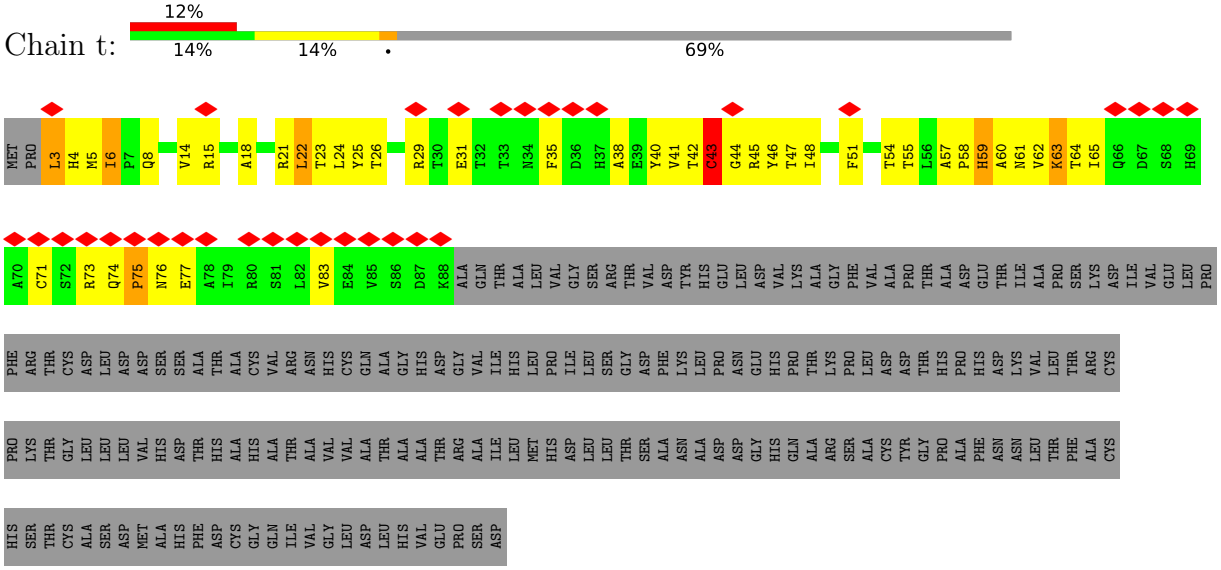
• Molecule 1: Outer capsid VP7

Chain r:



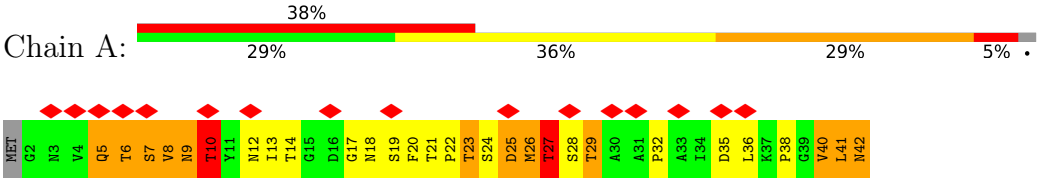
• Molecule 1: Outer capsid VP7

Chain t:



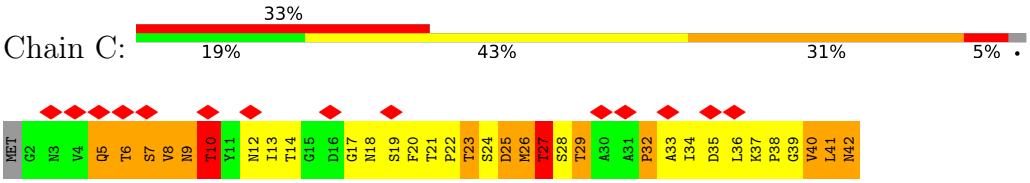
• Molecule 2: N-terminus of outer capsid protein VP5

Chain A:



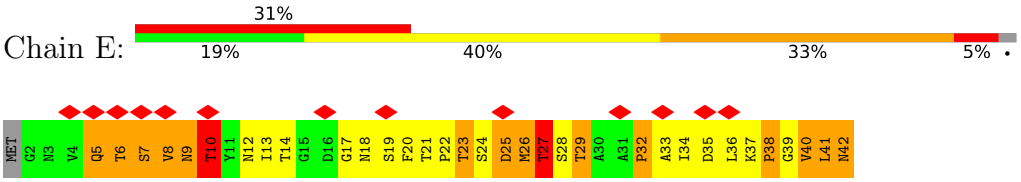
• Molecule 2: N-terminus of outer capsid protein VP5

Chain C:



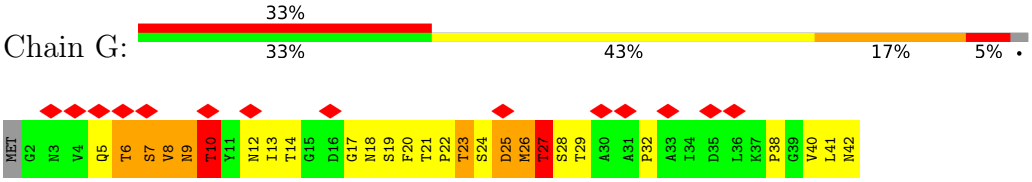
• Molecule 2: N-terminus of outer capsid protein VP5

Chain E:



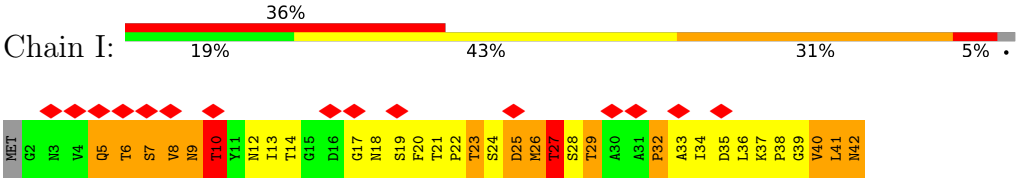
• Molecule 2: N-terminus of outer capsid protein VP5

Chain G:



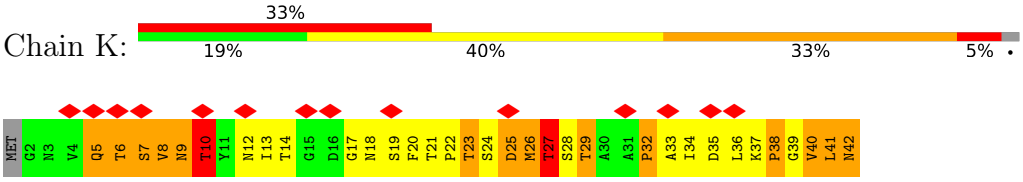
• Molecule 2: N-terminus of outer capsid protein VP5

Chain I:



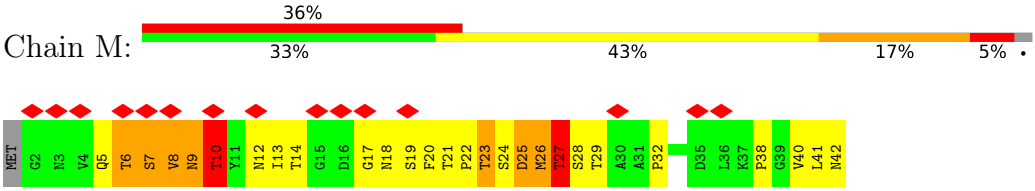
• Molecule 2: N-terminus of outer capsid protein VP5

Chain K:



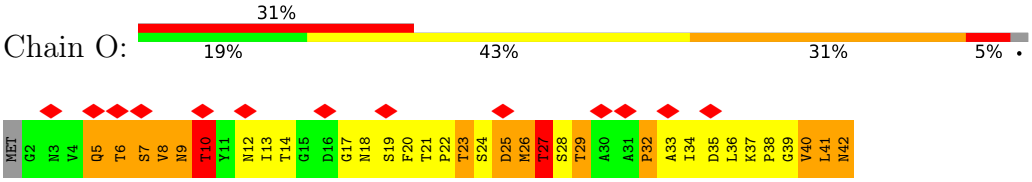
• Molecule 2: N-terminus of outer capsid protein VP5

Chain M:



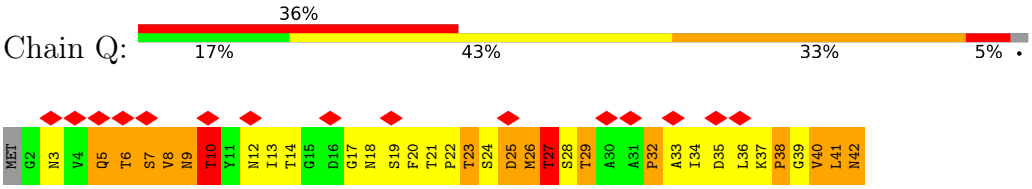
• Molecule 2: N-terminus of outer capsid protein VP5

Chain O:



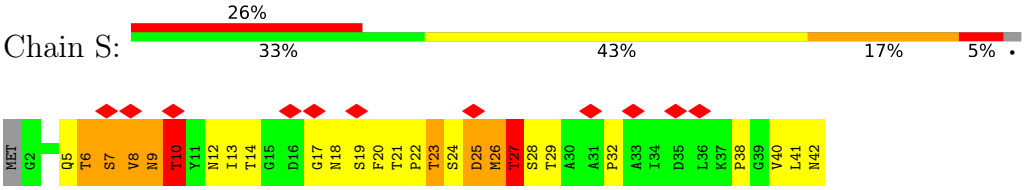
• Molecule 2: N-terminus of outer capsid protein VP5

Chain Q:



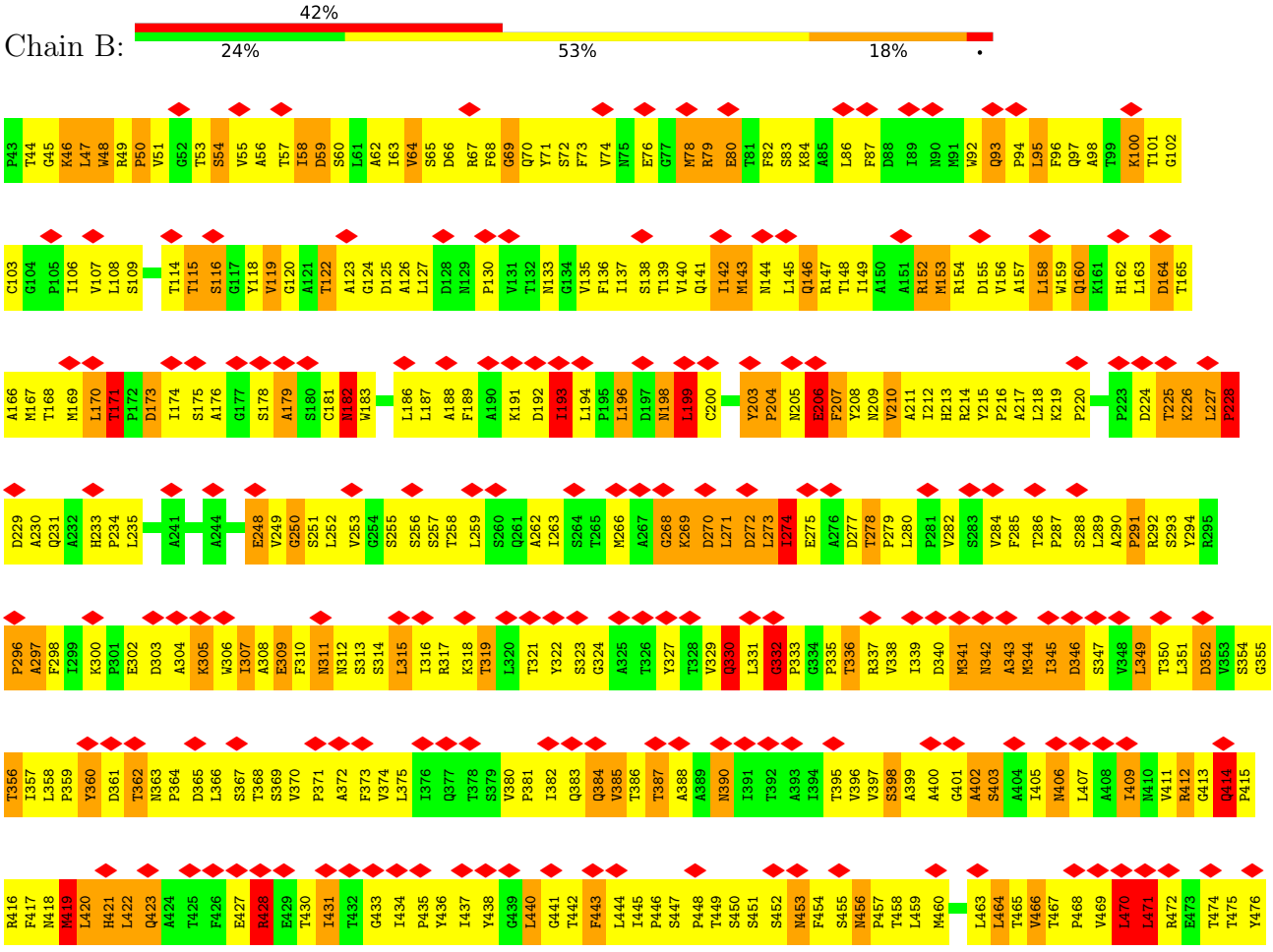
• Molecule 2: N-terminus of outer capsid protein VP5

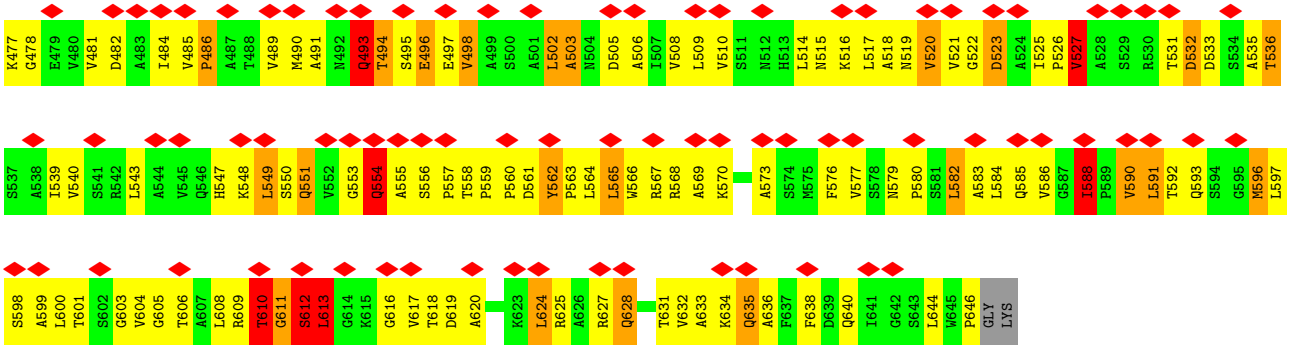
Chain S:



• Molecule 3: C-terminus of outer capsid protein VP5

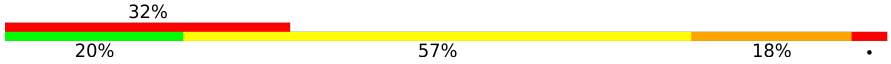
Chain B:

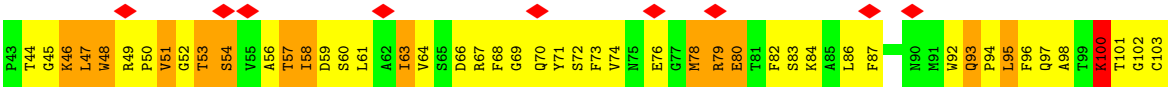


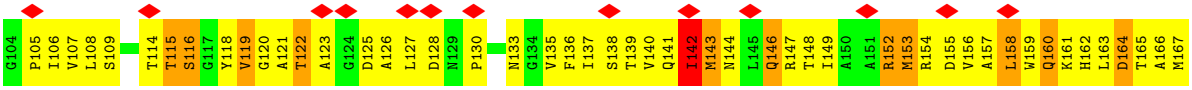


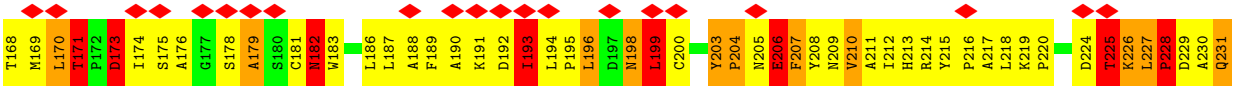
• Molecule 3: C-terminus of outer capsid protein VP5

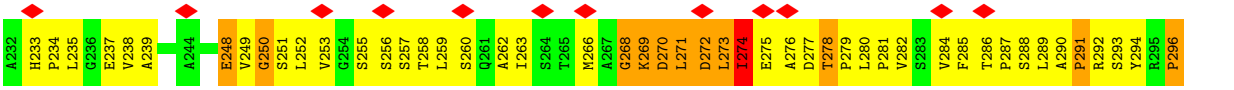
Chain D:

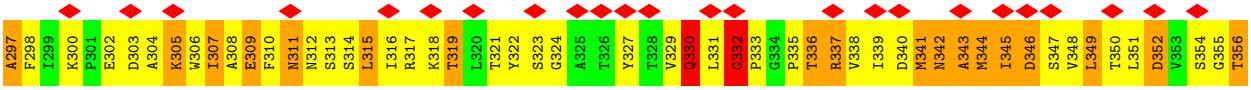


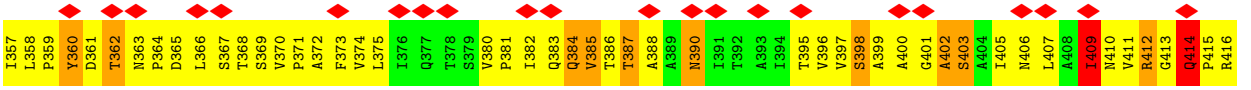


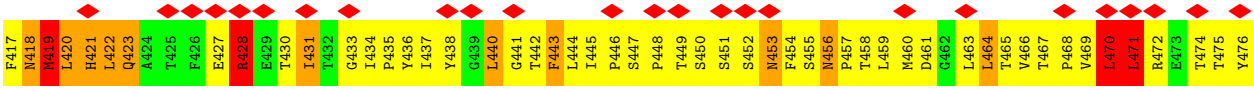


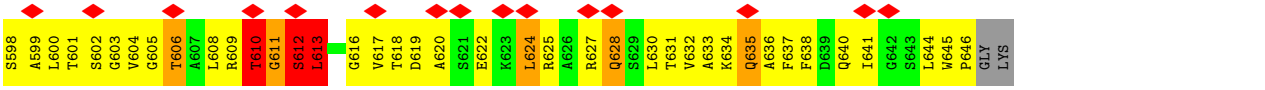






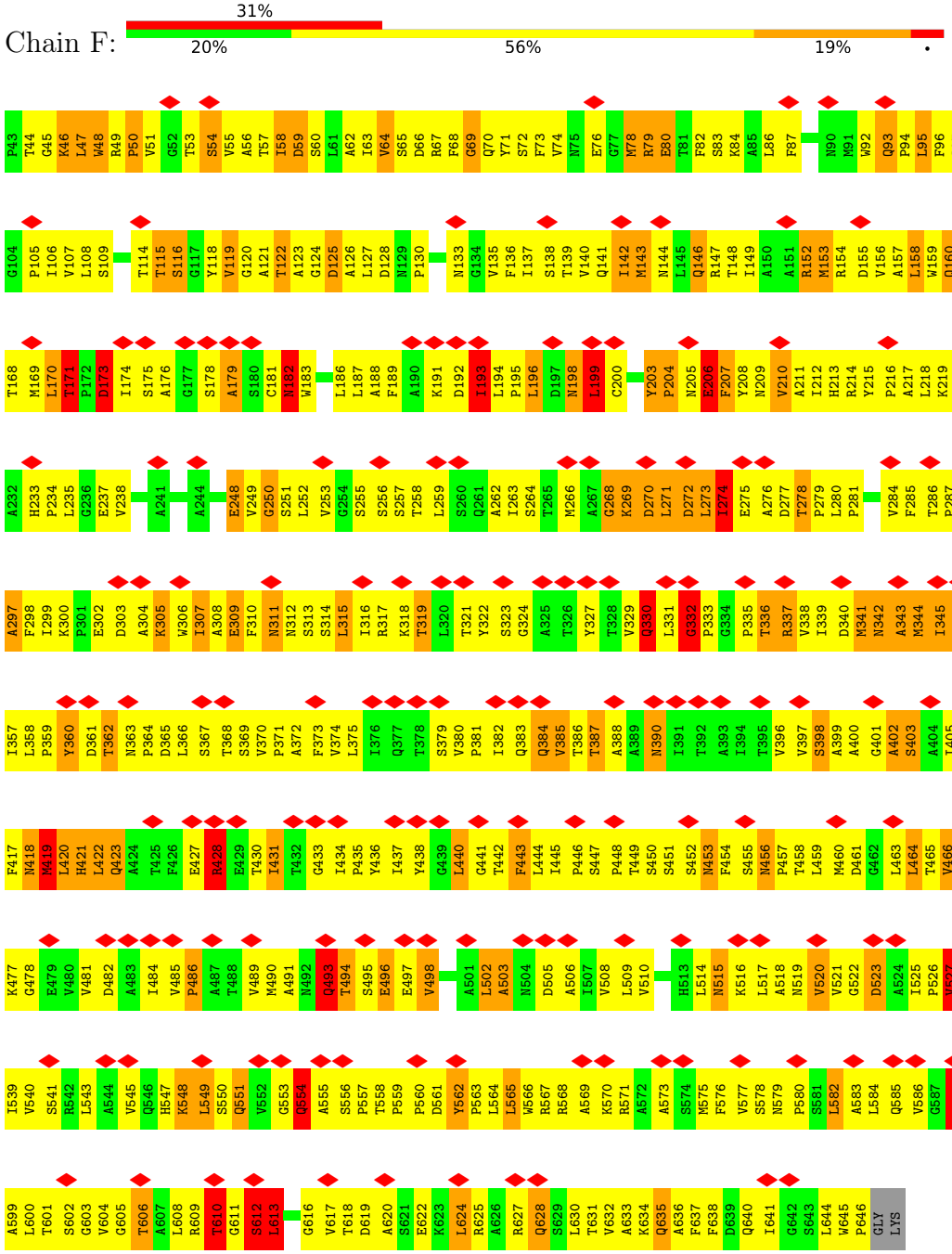






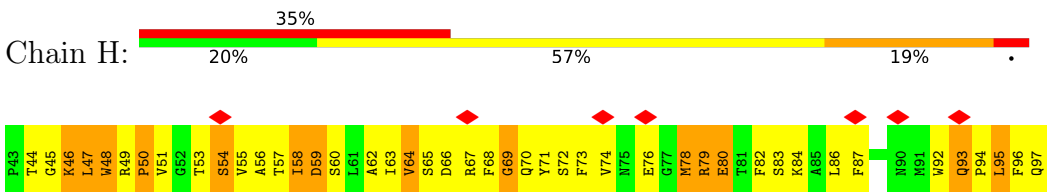
• Molecule 3: C-terminus of outer capsid protein VP5

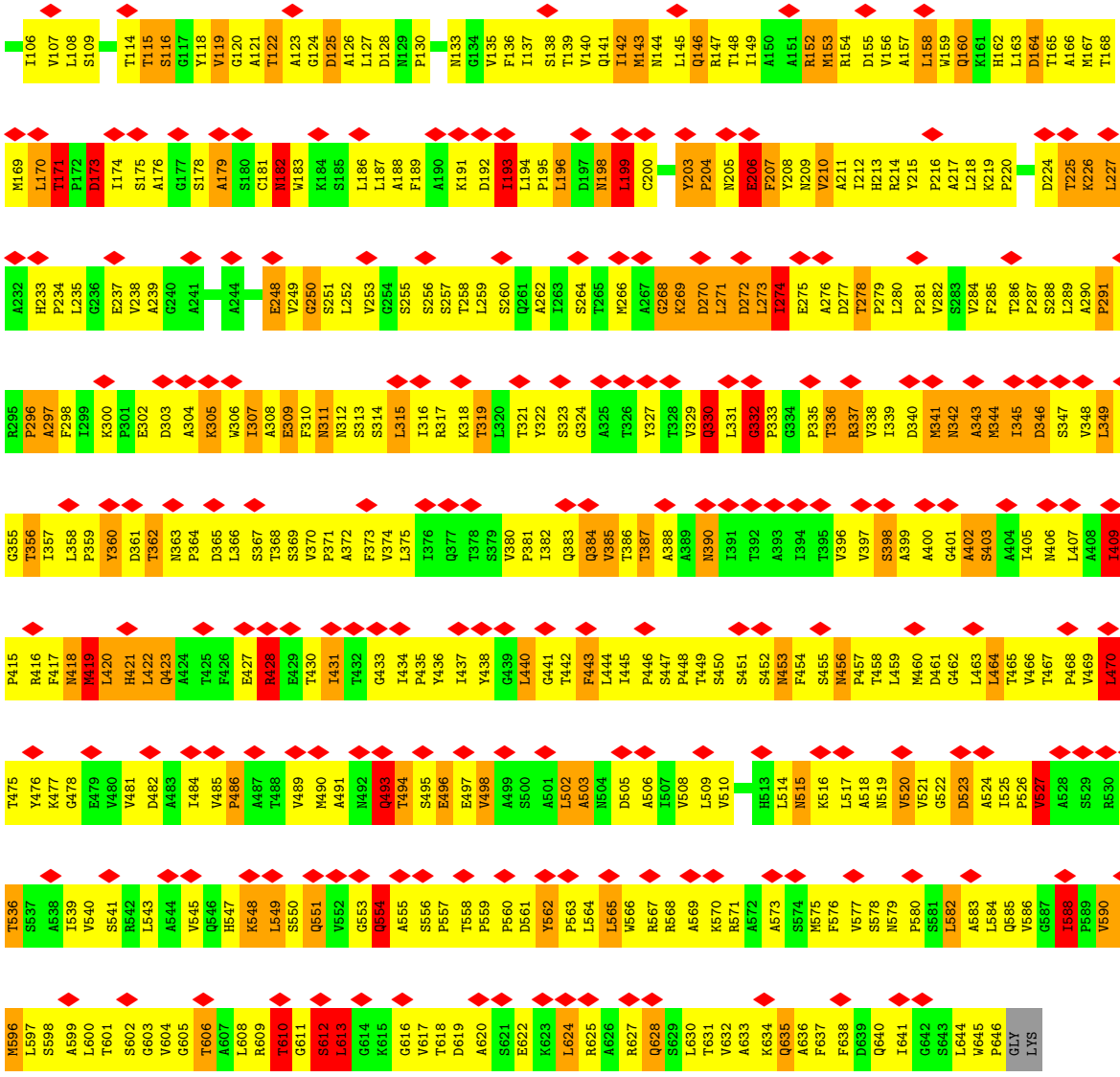
Chain F:



• Molecule 3: C-terminus of outer capsid protein VP5

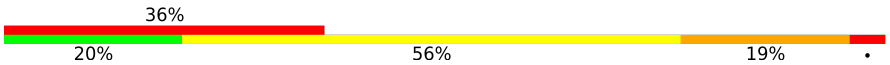
Chain H:

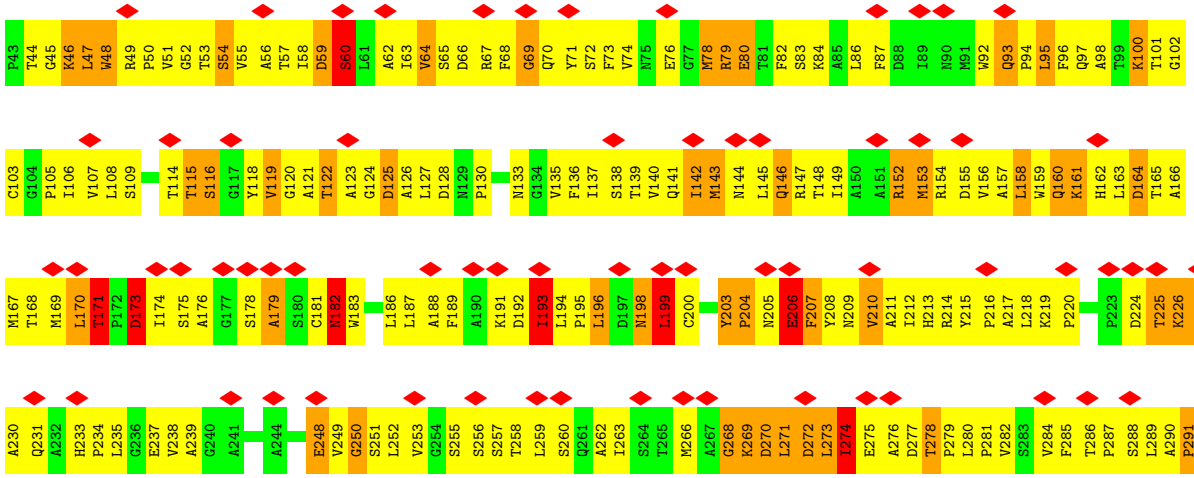


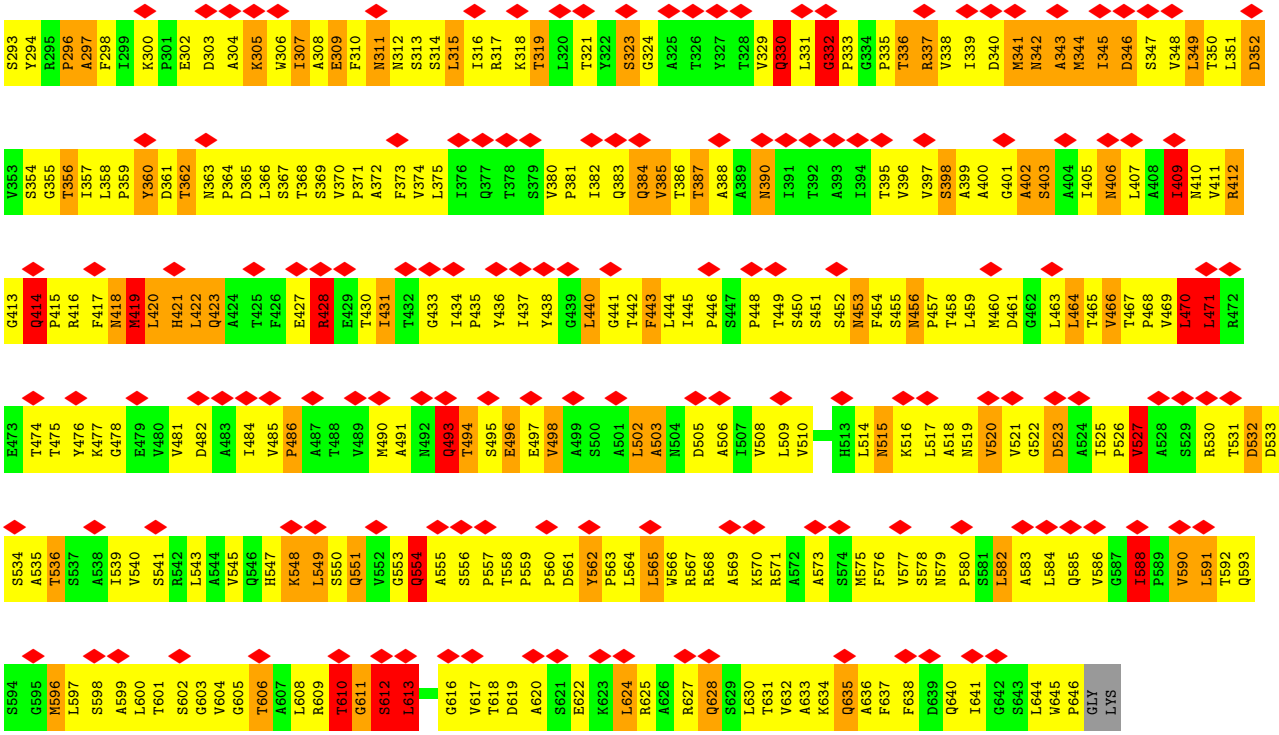


• Molecule 3: C-terminus of outer capsid protein VP5

Chain J:

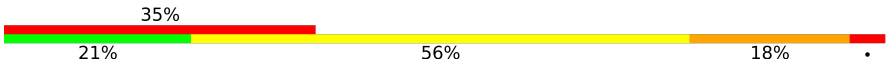


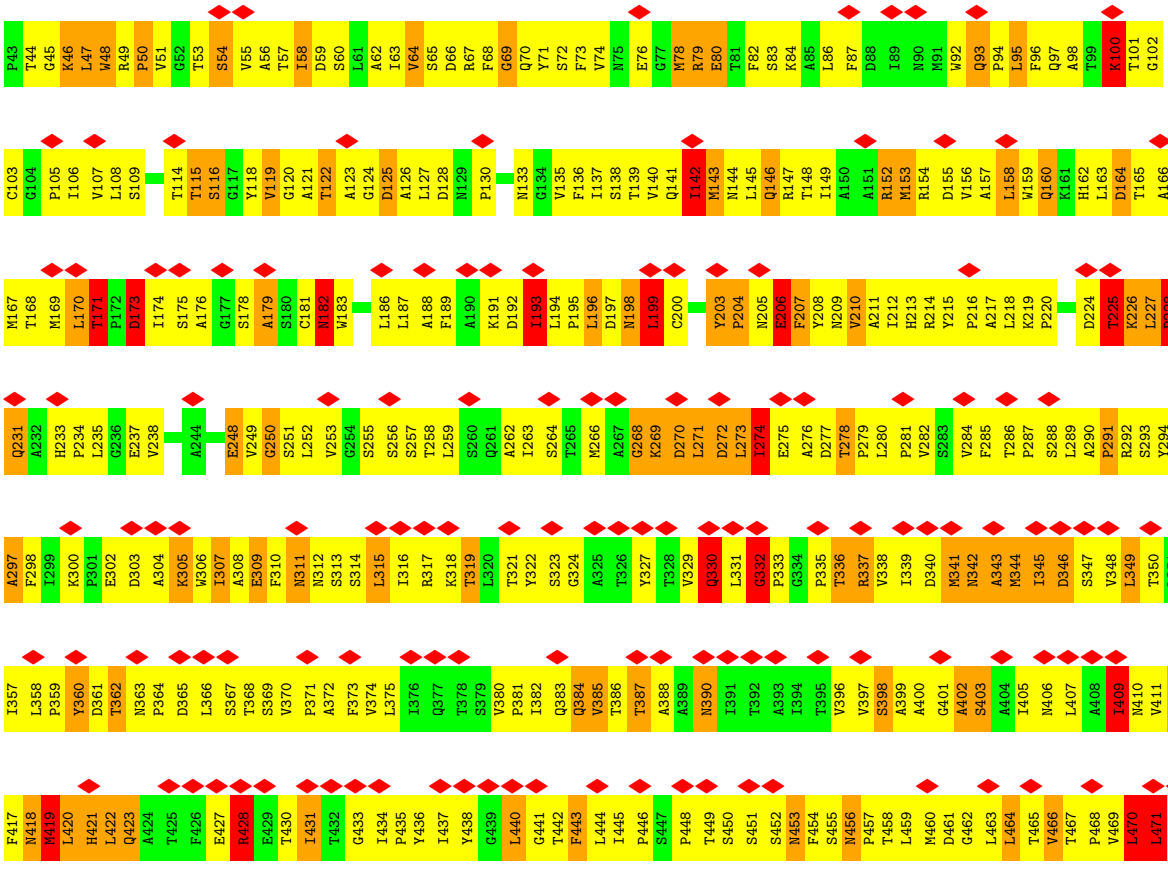


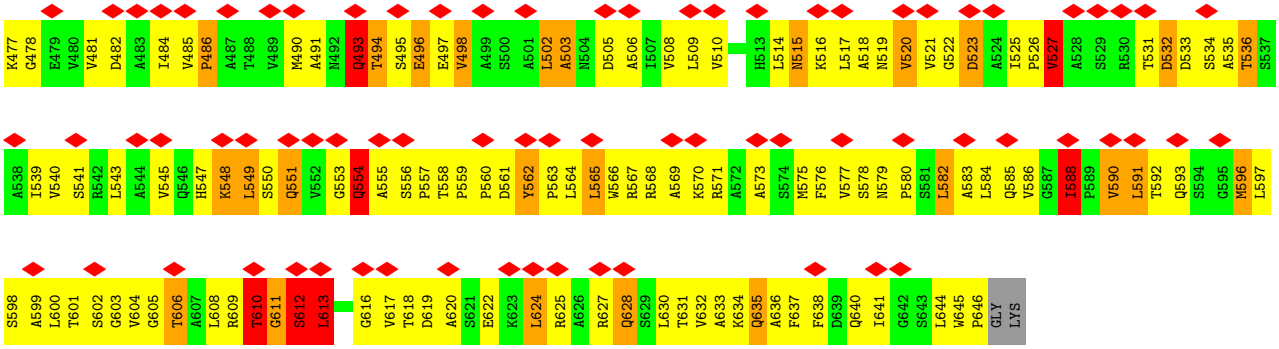


• Molecule 3: C-terminus of outer capsid protein VP5

Chain L:

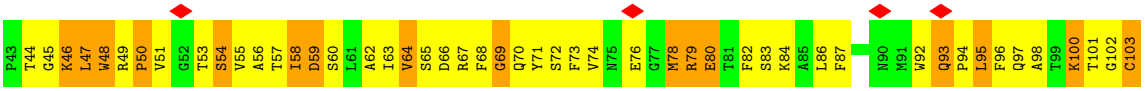


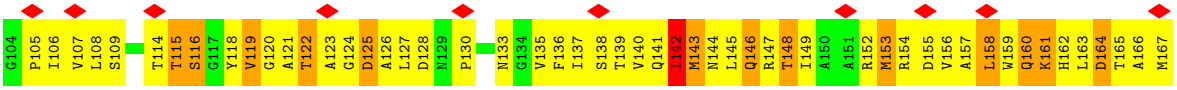


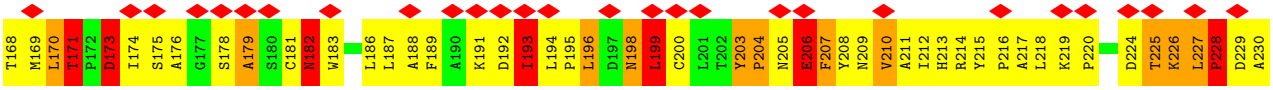


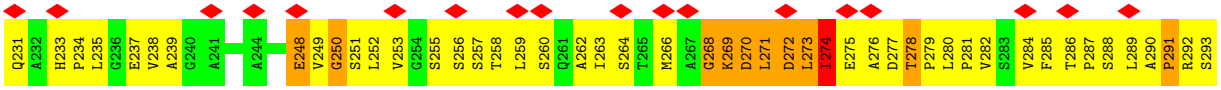
• Molecule 3: C-terminus of outer capsid protein VP5

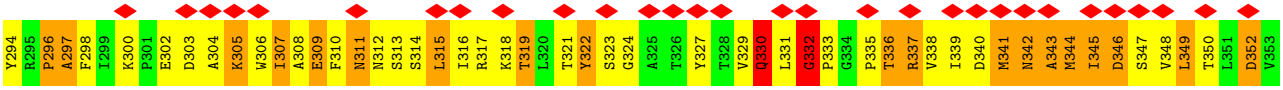
Chain N:

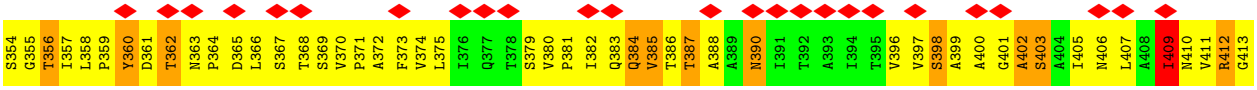


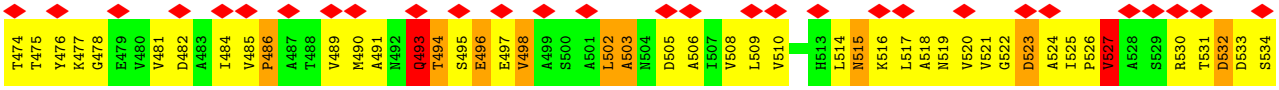


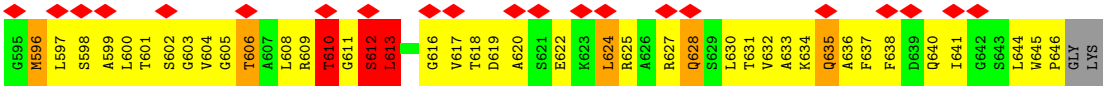






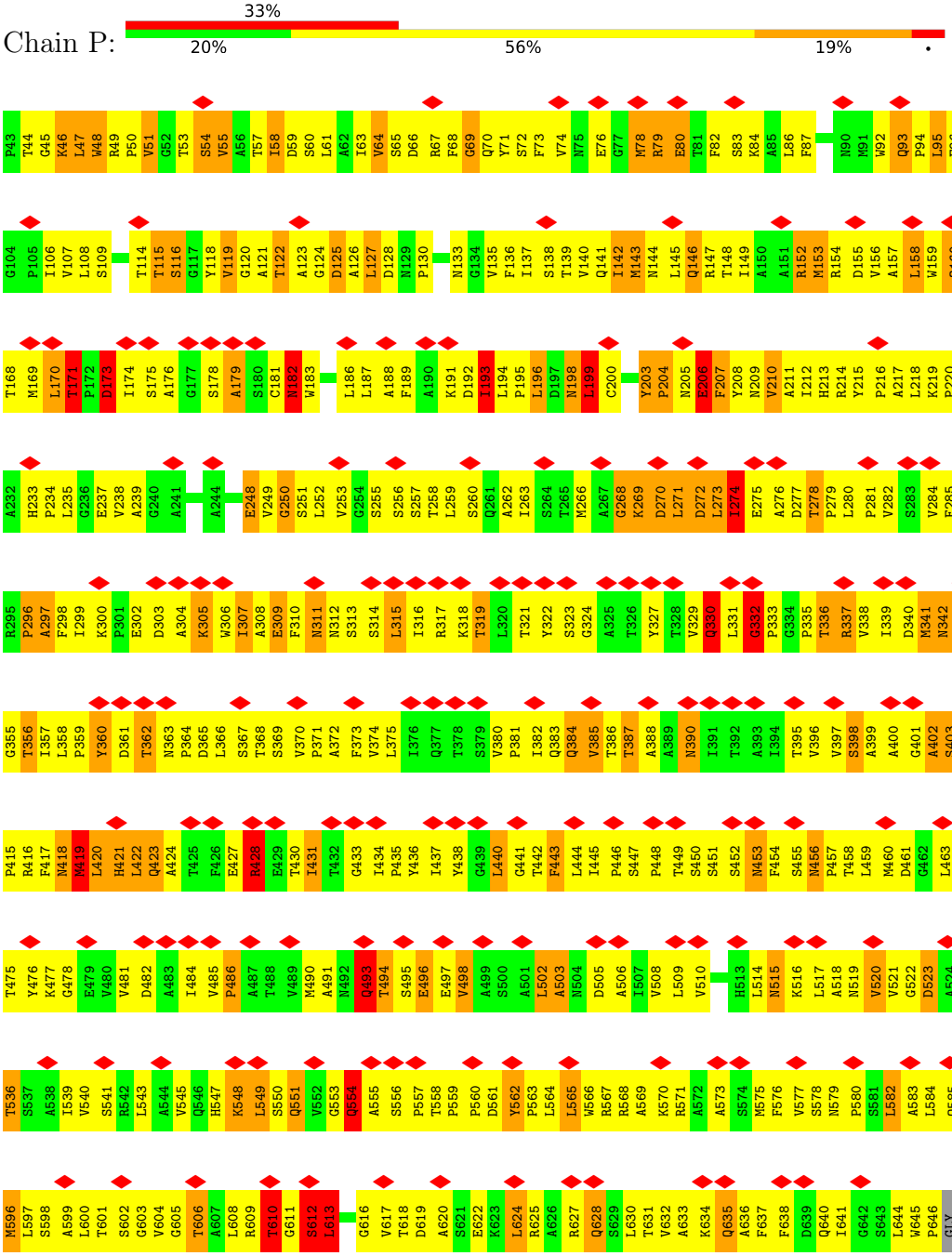






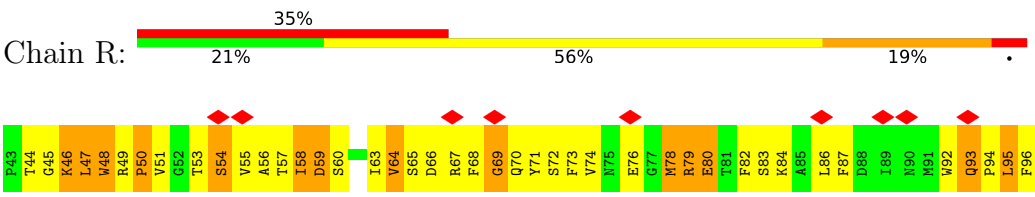
• Molecule 3: C-terminus of outer capsid protein VP5

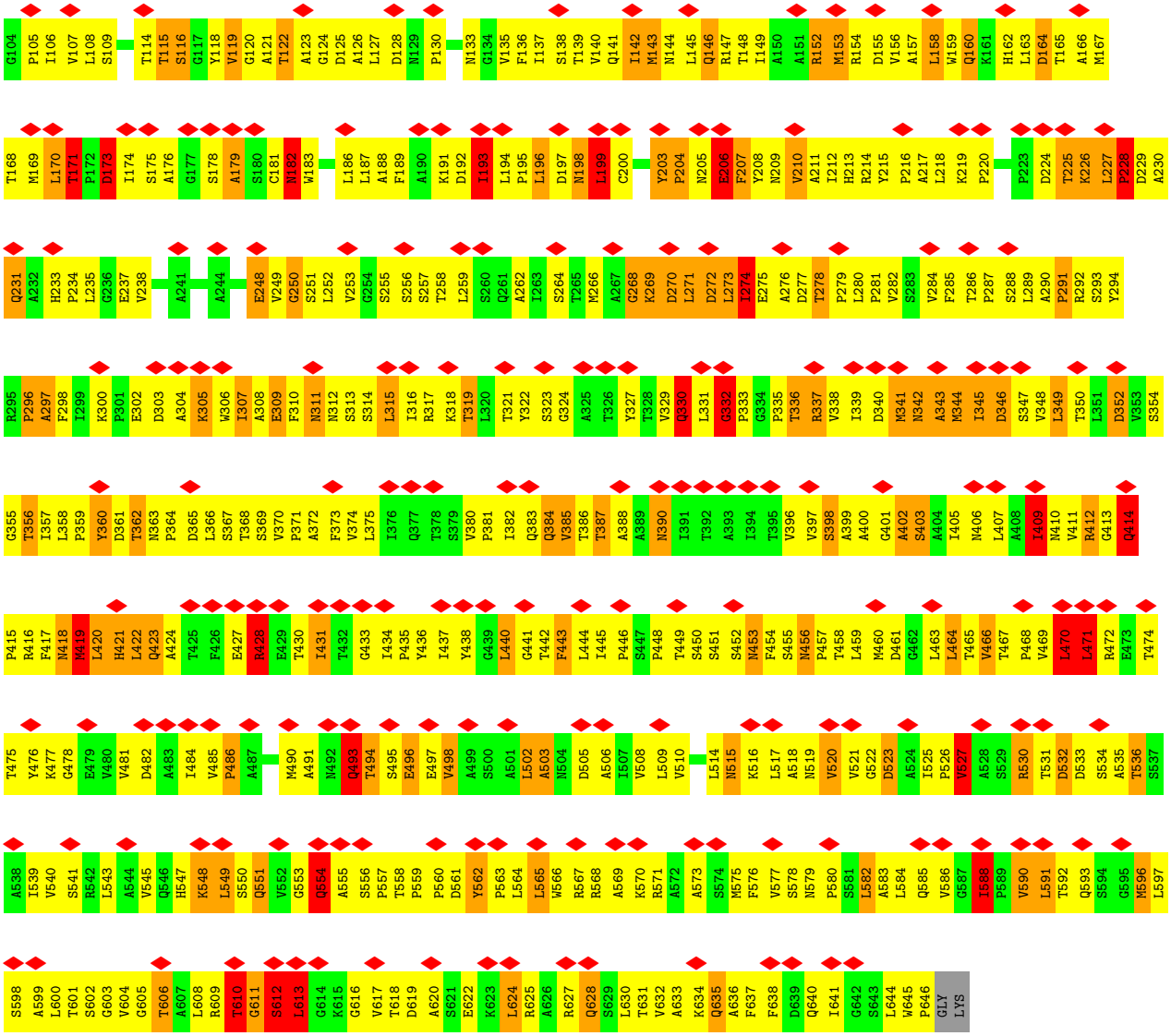
Chain P:



• Molecule 3: C-terminus of outer capsid protein VP5

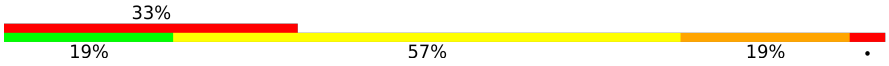
Chain R:

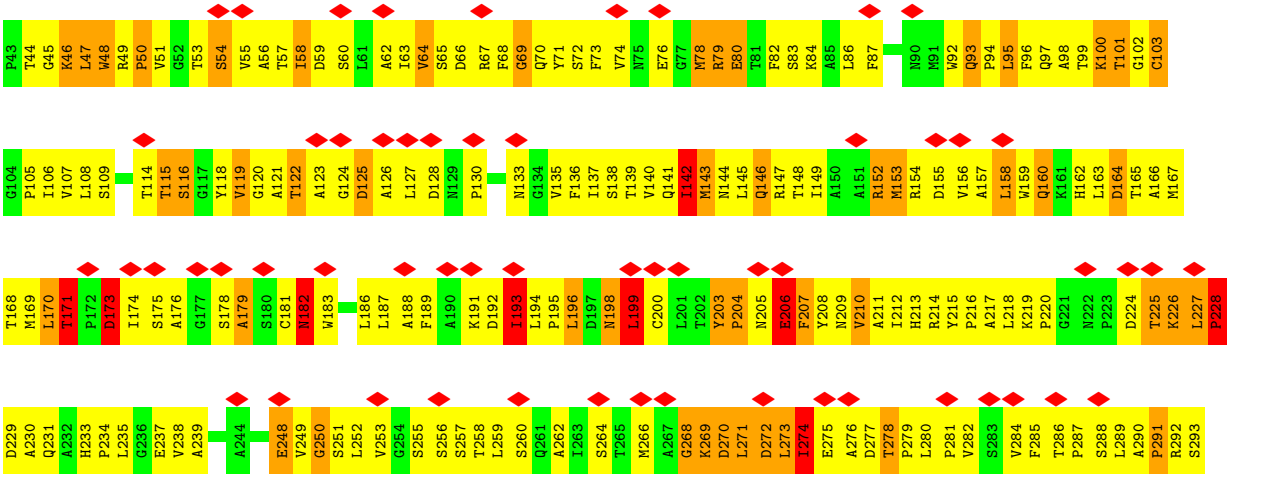


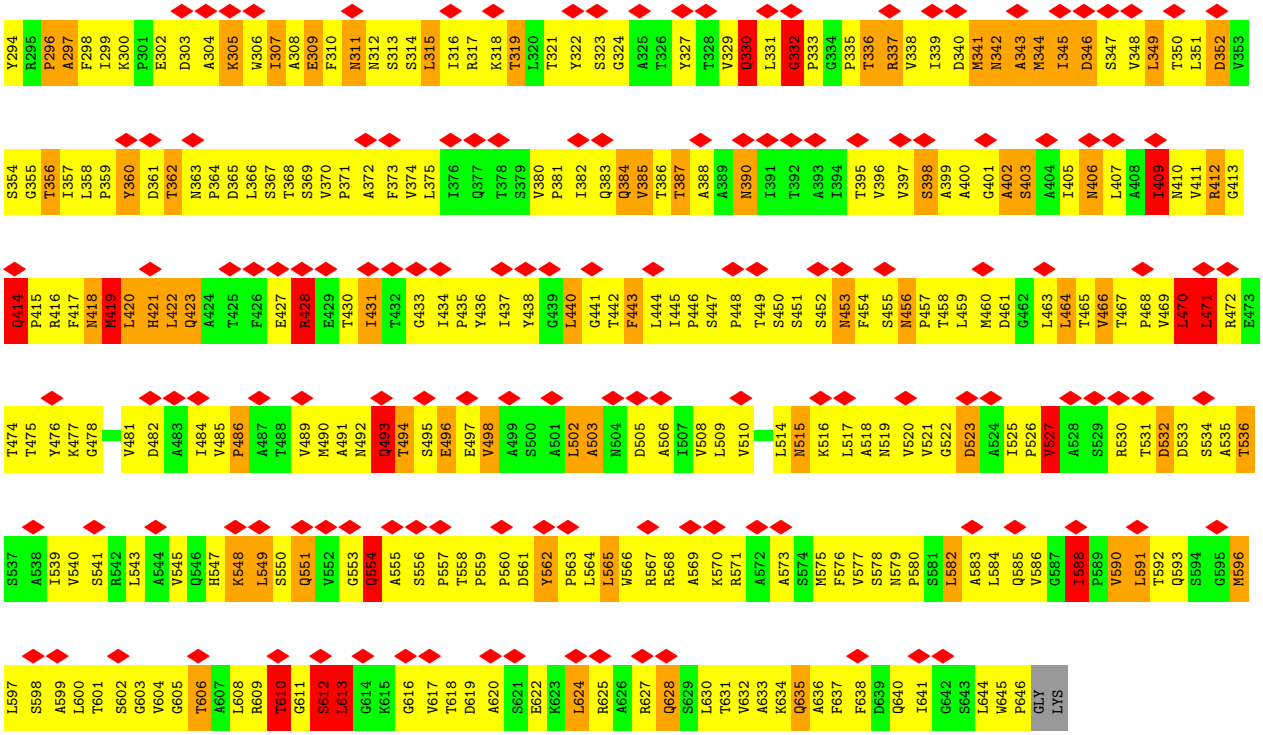


• Molecule 3: C-terminus of outer capsid protein VP5

Chain T:

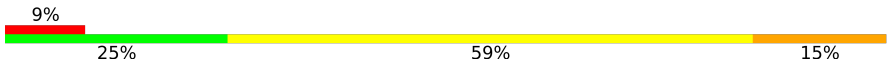


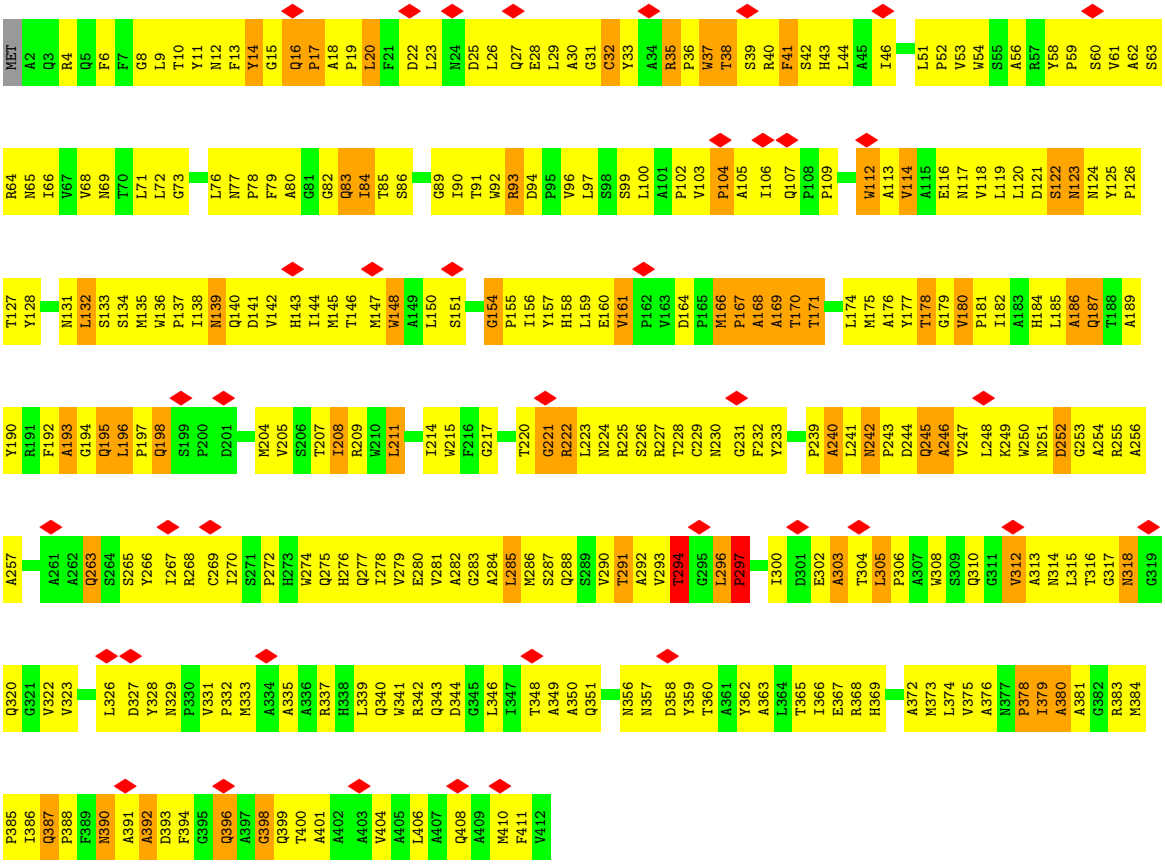




• Molecule 4: Core protein VP6

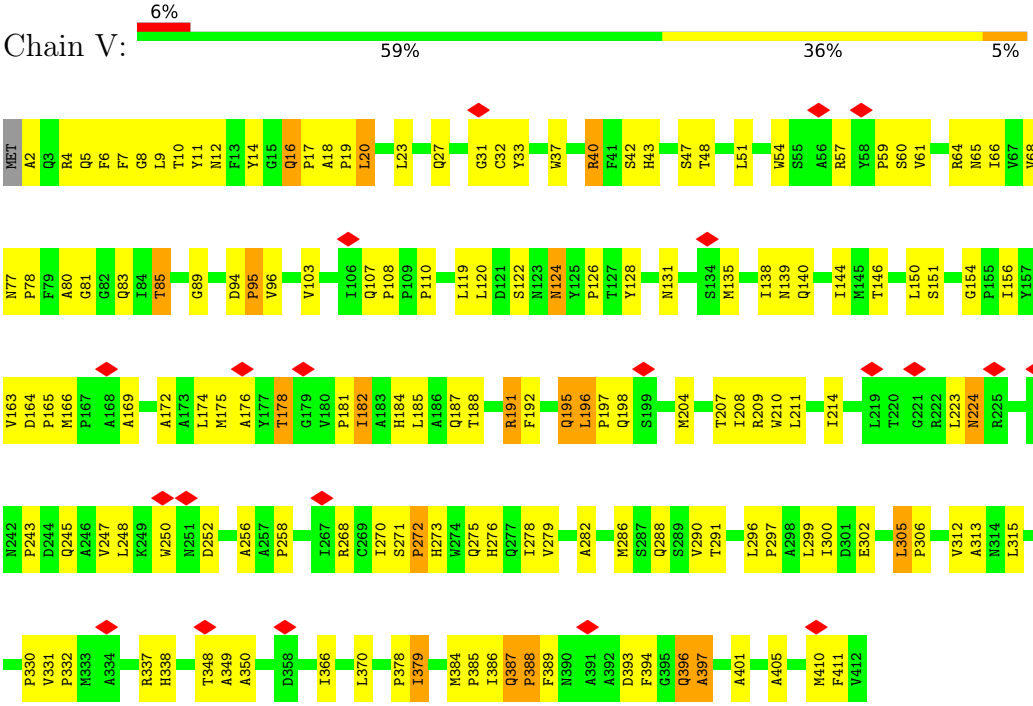
Chain U:





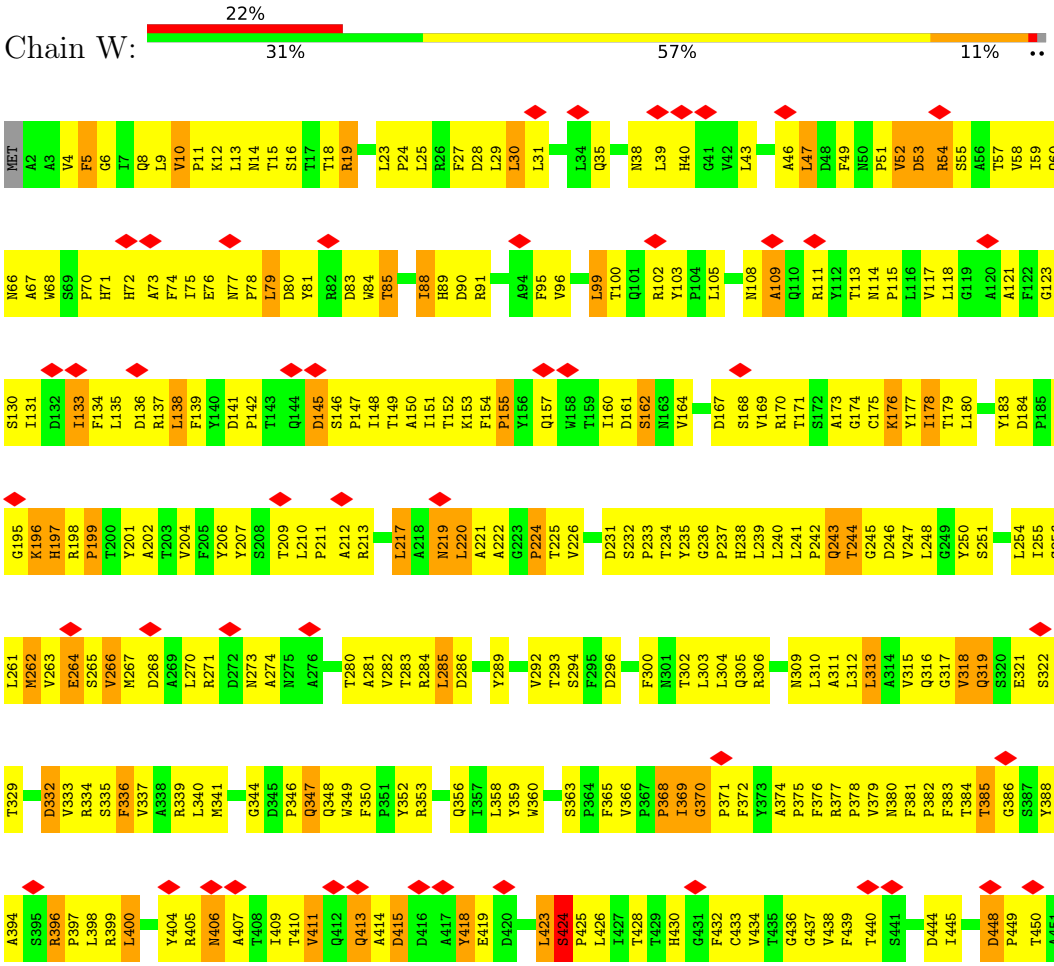
• Molecule 4: Core protein VP6

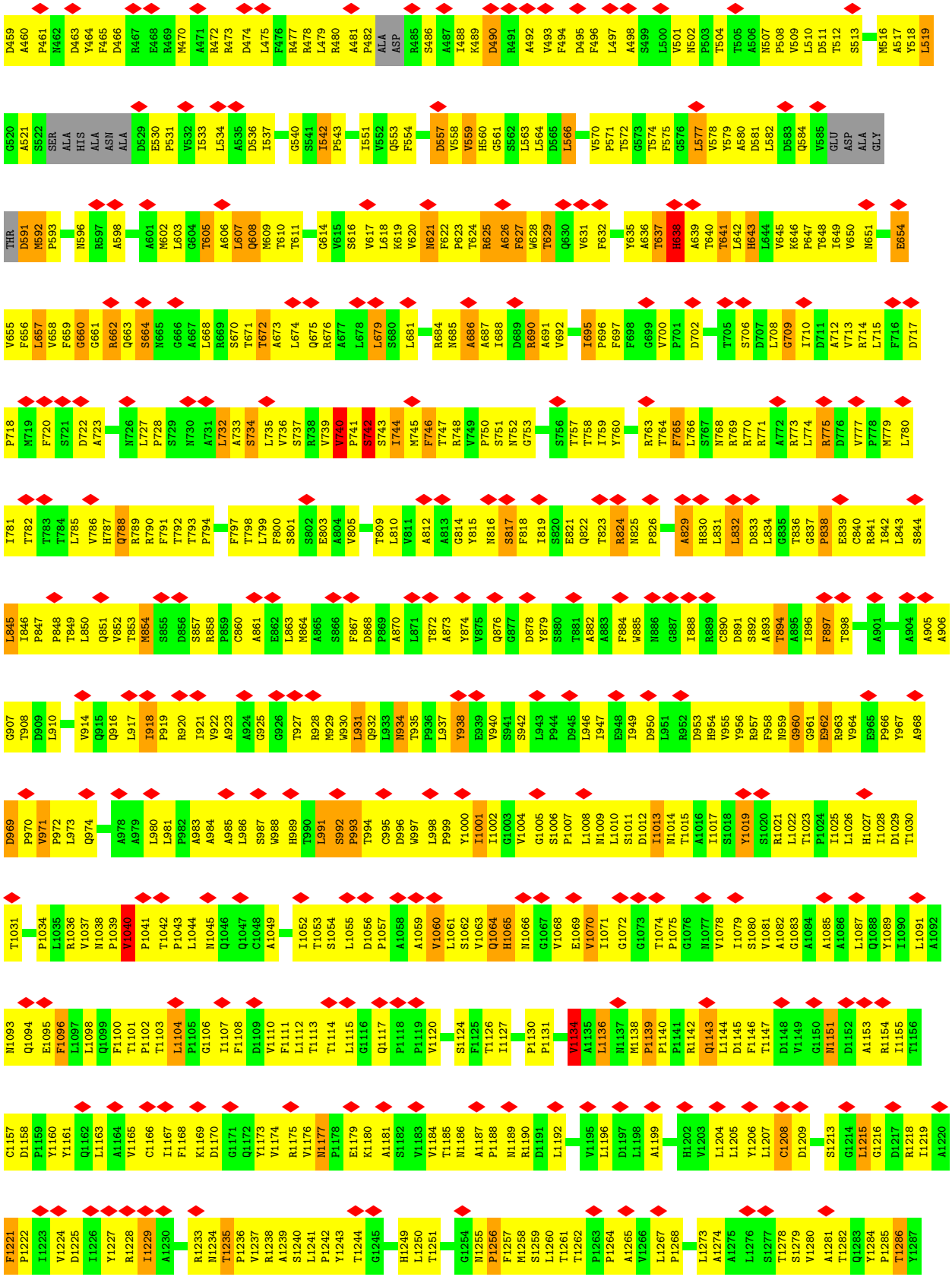
Chain V:

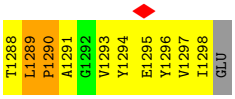


• Molecule 5: VP1

Chain W:

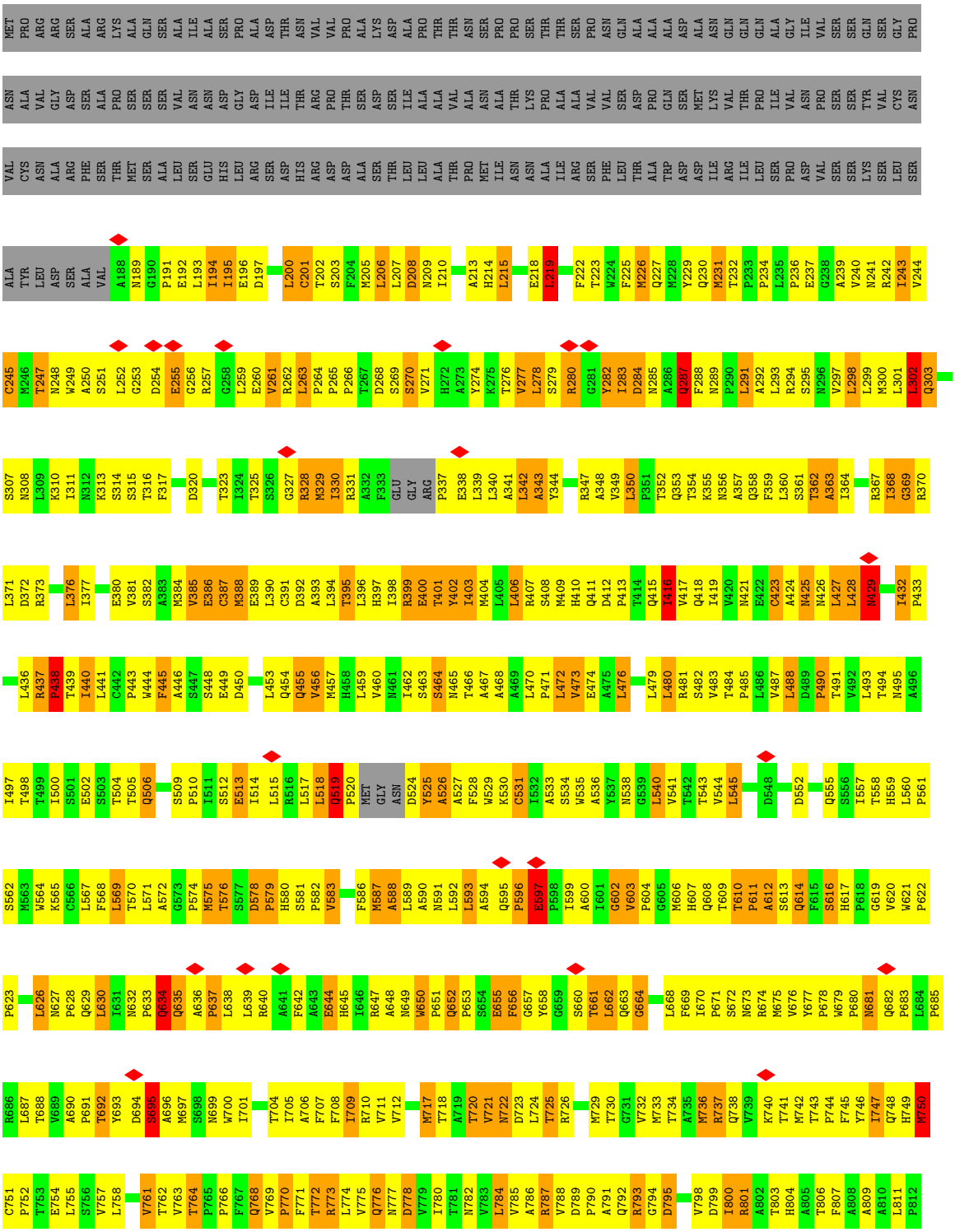


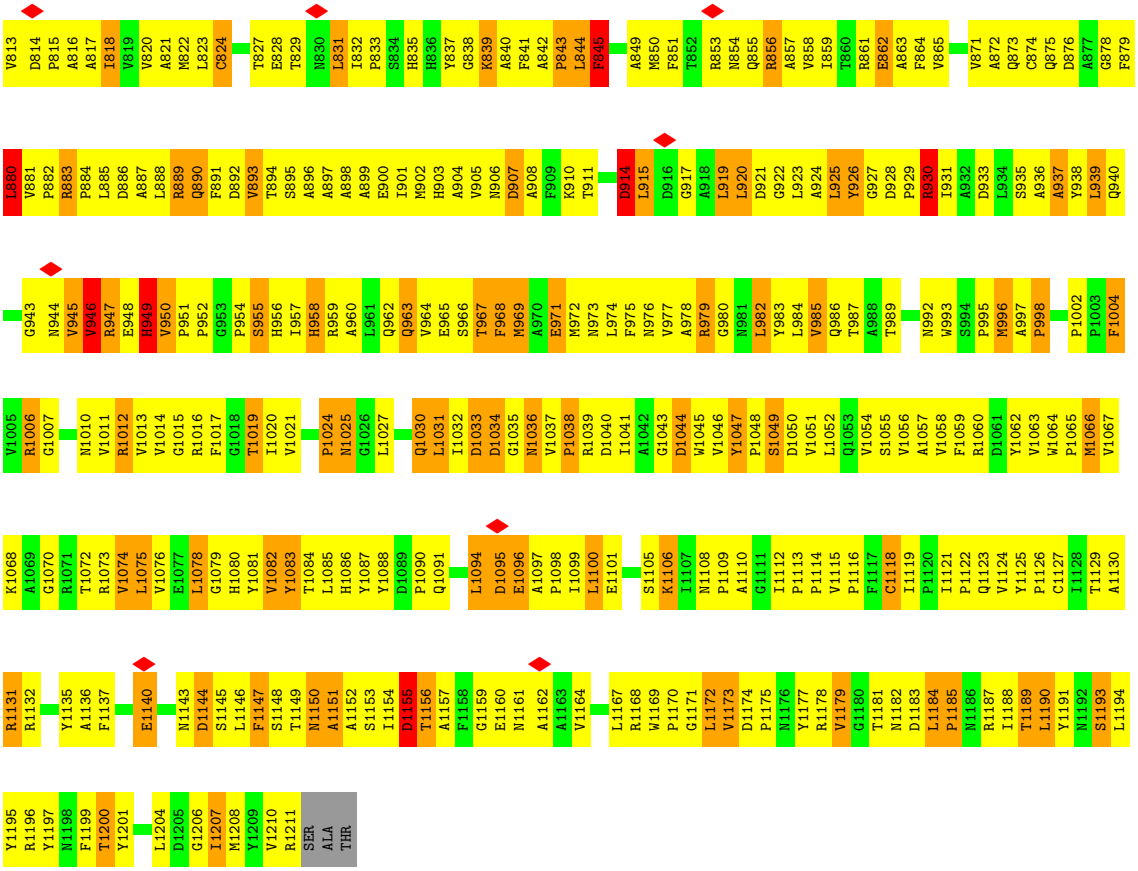




- Molecule 6: VP3

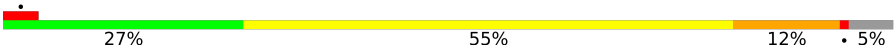
Chain X:

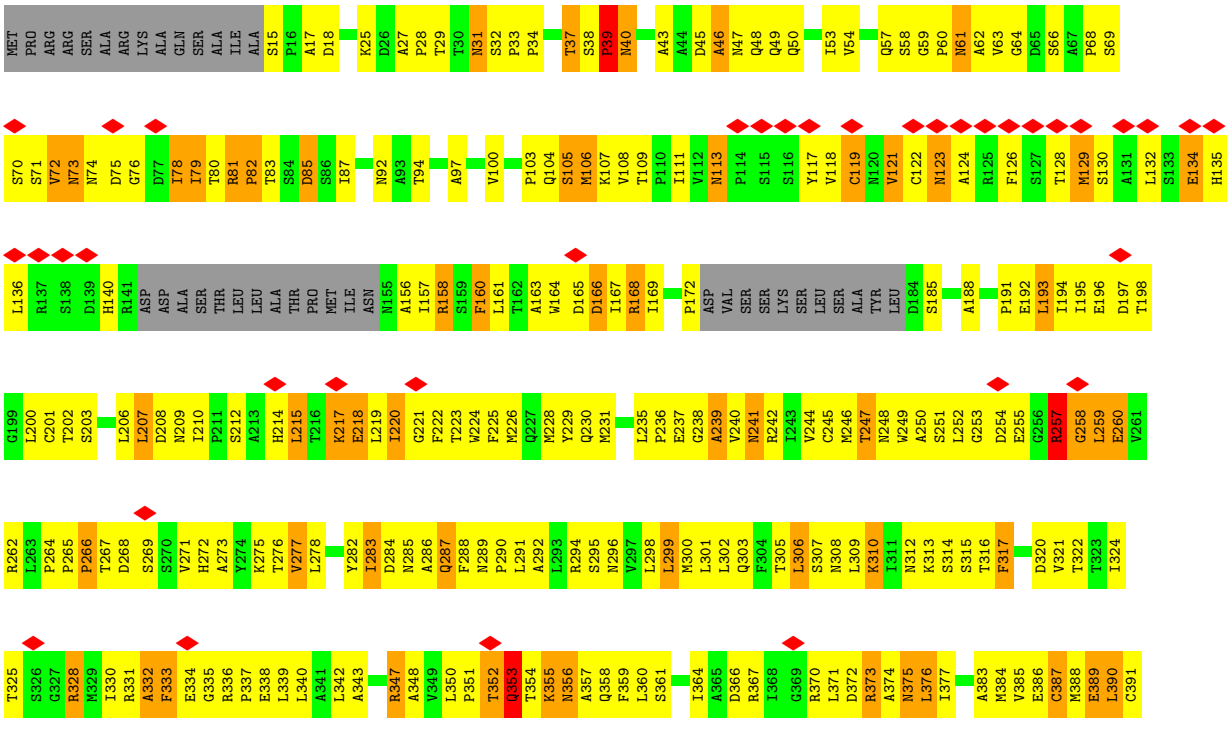


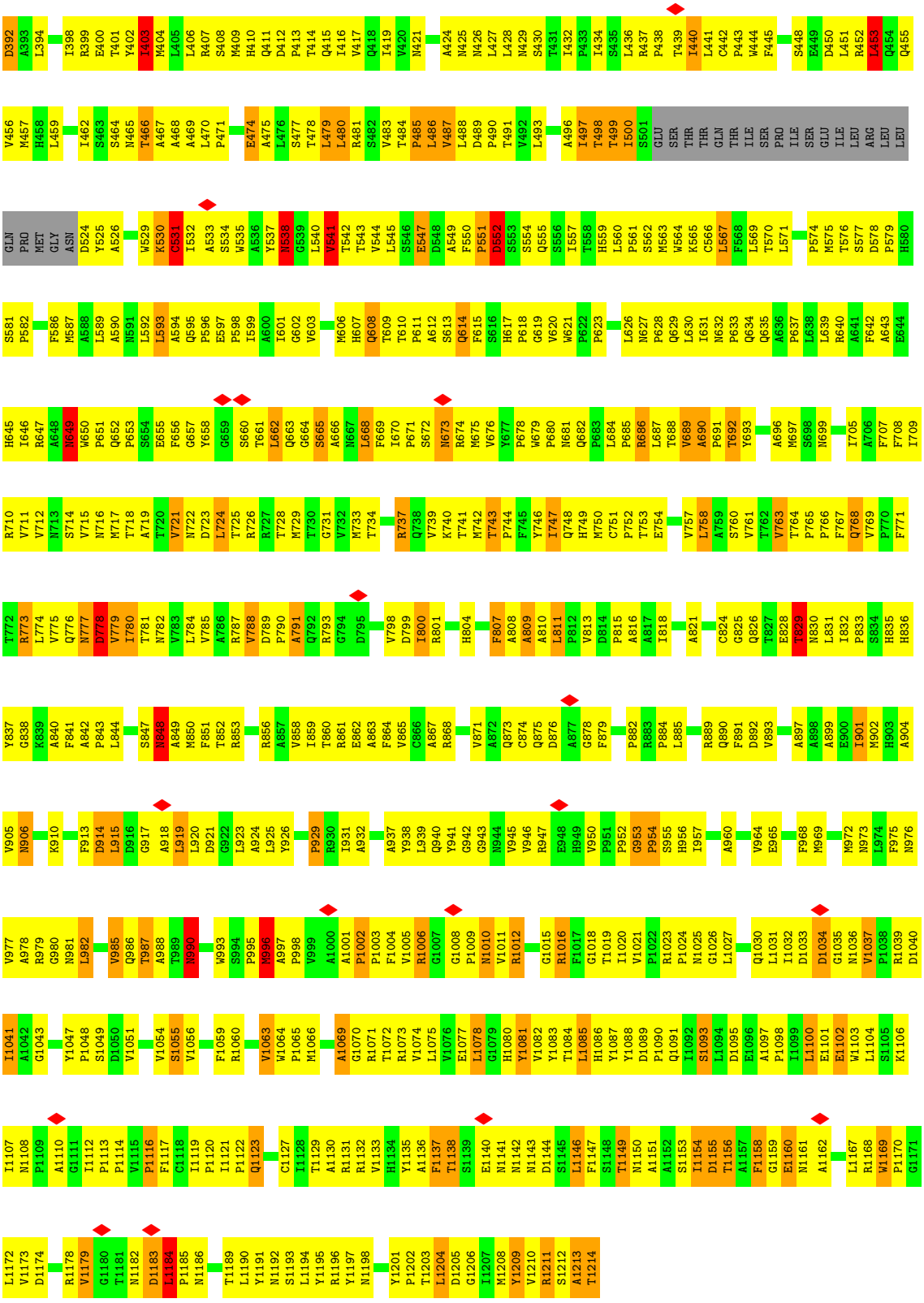


• Molecule 6: VP3

Chain Y:







4 Experimental information (i)

Property	Value	Source
EM reconstruction method	SINGLE PARTICLE	Depositor
Imposed symmetry	POINT, Not provided	
Number of particles used	41000	Depositor
Resolution determination method	FSC 0.143 CUT-OFF	Depositor
CTF correction method	PHASE FLIPPING AND AMPLITUDE CORRECTION	Depositor
Microscope	FEI TECNAI ARCTICA	Depositor
Voltage (kV)	200	Depositor
Electron dose ($e^-/Å^2$)	25	Depositor
Minimum defocus (nm)	Not provided	
Maximum defocus (nm)	Not provided	
Magnification	Not provided	
Image detector	FEI FALCON II (4k x 4k)	Depositor
Maximum map value	134.413	Depositor
Minimum map value	-88.945	Depositor
Average map value	0.680	Depositor
Map value standard deviation	8.761	Depositor
Recommended contour level	9	Depositor
Map size (Å)	838.8, 838.8, 838.8	wwPDB
Map dimensions	900, 900, 900	wwPDB
Map angles (°)	90.0, 90.0, 90.0	wwPDB
Pixel spacing (Å)	0.932, 0.932, 0.932	Depositor

5 Model quality (i)

5.1 Standard geometry (i)

Bond lengths and bond angles in the following residue types are not validated in this section: MYR

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	b	0.81	2/678 (0.3%)	1.50	8/922 (0.9%)
1	d	0.55	0/678	1.05	3/922 (0.3%)
1	f	0.55	0/678	1.39	16/922 (1.7%)
1	h	0.57	0/678	1.03	3/922 (0.3%)
1	j	0.56	0/678	1.08	11/922 (1.2%)
1	l	0.51	0/678	1.13	10/922 (1.1%)
1	n	0.53	0/678	1.01	3/922 (0.3%)
1	p	0.56	0/678	1.01	5/922 (0.5%)
1	r	0.51	0/678	1.10	7/922 (0.8%)
1	t	0.54	0/678	0.96	2/922 (0.2%)
2	A	0.90	2/295 (0.7%)	1.53	6/405 (1.5%)
2	C	0.90	2/295 (0.7%)	1.53	6/405 (1.5%)
2	E	0.90	2/295 (0.7%)	1.53	6/405 (1.5%)
2	G	0.89	2/286 (0.7%)	1.38	3/391 (0.8%)
2	I	0.90	2/295 (0.7%)	1.53	6/405 (1.5%)
2	K	0.90	2/295 (0.7%)	1.53	6/405 (1.5%)
2	M	0.89	2/286 (0.7%)	1.38	3/391 (0.8%)
2	O	0.90	2/295 (0.7%)	1.53	6/405 (1.5%)
2	Q	0.90	2/295 (0.7%)	1.53	6/405 (1.5%)
2	S	0.89	2/286 (0.7%)	1.38	3/391 (0.8%)
3	B	0.70	1/4601 (0.0%)	1.28	52/6295 (0.8%)
3	D	0.70	1/4601 (0.0%)	1.30	54/6295 (0.9%)
3	F	0.70	1/4601 (0.0%)	1.28	52/6295 (0.8%)
3	H	0.70	1/4601 (0.0%)	1.28	53/6295 (0.8%)
3	J	0.69	1/4601 (0.0%)	1.28	53/6295 (0.8%)
3	L	0.70	1/4601 (0.0%)	1.28	52/6295 (0.8%)
3	N	0.70	1/4601 (0.0%)	1.28	52/6295 (0.8%)
3	P	0.70	1/4601 (0.0%)	1.28	52/6295 (0.8%)
3	R	0.70	1/4601 (0.0%)	1.28	52/6295 (0.8%)
3	T	0.70	1/4601 (0.0%)	1.28	51/6295 (0.8%)
4	U	0.38	0/3233	0.93	16/4443 (0.4%)
4	V	0.41	0/3233	0.88	5/4443 (0.1%)

Mol	Chain	Bond lengths		Bond angles	
		RMSZ	#|Z| >5	RMSZ	#|Z| >5
5	W	0.38	0/10148	1.00	52/13935 (0.4%)
6	X	0.65	5/8078 (0.1%)	1.16	69/11071 (0.6%)
6	Y	0.45	1/9056 (0.0%)	1.04	52/12412 (0.4%)
All	All	0.62	38/89461 (0.0%)	1.19	836/122482 (0.7%)

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	b	0	4
1	d	0	3
1	f	0	2
1	h	0	2
1	j	0	2
1	l	0	2
1	n	0	3
1	p	0	3
1	r	0	2
1	t	0	2
3	B	0	1
3	D	0	1
3	F	0	1
3	H	0	1
3	J	0	1
3	L	0	1
3	N	0	1
3	P	0	1
3	R	0	1
3	T	0	1
6	X	0	3
6	Y	0	2
All	All	0	40

All (38) bond length outliers are listed below:

Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
1	b	74	GLN	C-N	14.77	1.68	1.33
3	N	227	LEU	N-CA	7.64	1.52	1.45
3	H	227	LEU	N-CA	7.64	1.52	1.45
3	R	227	LEU	N-CA	7.63	1.52	1.45

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
3	F	227	LEU	N-CA	7.60	1.52	1.45
3	D	227	LEU	N-CA	7.60	1.52	1.45
3	J	227	LEU	N-CA	7.58	1.52	1.45
3	L	227	LEU	N-CA	7.58	1.52	1.45
3	P	227	LEU	N-CA	7.54	1.52	1.45
3	B	227	LEU	N-CA	7.54	1.52	1.45
3	T	227	LEU	N-CA	7.51	1.51	1.45
6	X	969	MET	SD-CE	6.55	1.96	1.79
2	Q	10	THR	N-CA	5.96	1.53	1.46
2	O	10	THR	N-CA	5.96	1.53	1.46
2	K	10	THR	N-CA	5.94	1.53	1.46
2	A	10	THR	N-CA	5.92	1.53	1.46
2	I	10	THR	N-CA	5.91	1.53	1.46
2	C	10	THR	N-CA	5.90	1.53	1.46
2	E	10	THR	N-CA	5.89	1.53	1.46
6	X	231	MET	SD-CE	5.88	1.94	1.79
2	S	10	THR	N-CA	5.86	1.53	1.46
2	M	10	THR	N-CA	5.86	1.53	1.46
2	G	10	THR	N-CA	5.85	1.53	1.46
2	C	9	ASN	CA-C	5.84	1.59	1.52
6	X	717	MET	SD-CE	5.83	1.94	1.79
1	b	73	ARG	C-N	5.82	1.45	1.33
2	O	9	ASN	CA-C	5.80	1.59	1.52
2	A	9	ASN	CA-C	5.79	1.59	1.52
2	I	9	ASN	CA-C	5.79	1.59	1.52
2	Q	9	ASN	CA-C	5.79	1.59	1.52
2	G	9	ASN	CA-C	5.76	1.59	1.52
2	S	9	ASN	CA-C	5.75	1.59	1.52
2	E	9	ASN	CA-C	5.75	1.59	1.52
2	M	9	ASN	CA-C	5.75	1.59	1.52
6	Y	996	MET	SD-CE	-5.74	1.65	1.79
2	K	9	ASN	CA-C	5.71	1.59	1.52
6	X	226	MET	SD-CE	5.54	1.93	1.79
6	X	736	MET	SD-CE	5.14	1.92	1.79

All (836) bond angle outliers are listed below:

Mol	Chain	Res	Type	Atoms	Z	Observed(^{o})	Ideal(^{o})
1	b	73	ARG	O-C-N	-31.64	90.78	121.79
3	D	51	VAL	N-CA-C	17.02	126.53	110.53
1	f	73	ARG	N-CA-C	13.44	125.93	111.28
6	X	526	ALA	N-CA-C	-11.80	98.50	111.36

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Mol	Chain	Res	Type	Atoms	Z	Observed(^{o})	Ideal(^{o})
5	W	63	PRO	N-CA-C	11.78	125.07	110.70
1	b	74	GLN	O-C-N	-10.88	108.81	121.32
3	F	551	GLN	OE1-CD-NE2	-10.57	112.03	122.60
3	D	551	GLN	OE1-CD-NE2	-10.56	112.04	122.60
3	J	551	GLN	OE1-CD-NE2	-10.56	112.04	122.60
3	L	551	GLN	OE1-CD-NE2	-10.55	112.05	122.60
3	P	551	GLN	OE1-CD-NE2	-10.54	112.06	122.60
3	B	551	GLN	OE1-CD-NE2	-10.52	112.08	122.60
3	R	551	GLN	OE1-CD-NE2	-10.52	112.08	122.60
3	T	551	GLN	OE1-CD-NE2	-10.51	112.09	122.60
3	N	551	GLN	OE1-CD-NE2	-10.48	112.12	122.60
3	H	551	GLN	OE1-CD-NE2	-10.44	112.16	122.60
3	F	493	GLN	OE1-CD-NE2	-10.22	112.38	122.60
3	L	493	GLN	OE1-CD-NE2	-10.19	112.41	122.60
3	H	493	GLN	OE1-CD-NE2	-10.19	112.41	122.60
3	N	160	GLN	OE1-CD-NE2	-10.18	112.42	122.60
3	N	493	GLN	OE1-CD-NE2	-10.18	112.42	122.60
3	R	493	GLN	OE1-CD-NE2	-10.18	112.42	122.60
3	P	493	GLN	OE1-CD-NE2	-10.17	112.43	122.60
3	T	493	GLN	OE1-CD-NE2	-10.16	112.44	122.60
3	H	160	GLN	OE1-CD-NE2	-10.15	112.44	122.60
3	P	160	GLN	OE1-CD-NE2	-10.15	112.44	122.60
3	B	493	GLN	OE1-CD-NE2	-10.13	112.47	122.60
3	J	493	GLN	OE1-CD-NE2	-10.13	112.47	122.60
3	D	160	GLN	OE1-CD-NE2	-10.12	112.48	122.60
3	J	160	GLN	OE1-CD-NE2	-10.12	112.48	122.60
3	R	160	GLN	OE1-CD-NE2	-10.11	112.49	122.60
3	B	160	GLN	OE1-CD-NE2	-10.11	112.49	122.60
3	F	160	GLN	OE1-CD-NE2	-10.09	112.51	122.60
3	L	160	GLN	OE1-CD-NE2	-10.09	112.51	122.60
3	D	493	GLN	OE1-CD-NE2	-10.09	112.51	122.60
3	T	160	GLN	OE1-CD-NE2	-10.09	112.51	122.60
3	B	97	GLN	OE1-CD-NE2	-9.97	112.63	122.60
3	L	97	GLN	OE1-CD-NE2	-9.97	112.63	122.60
3	T	97	GLN	OE1-CD-NE2	-9.97	112.63	122.60
3	N	97	GLN	OE1-CD-NE2	-9.97	112.63	122.60
3	R	97	GLN	OE1-CD-NE2	-9.97	112.63	122.60
3	P	97	GLN	OE1-CD-NE2	-9.96	112.64	122.60
3	D	97	GLN	OE1-CD-NE2	-9.95	112.65	122.60
3	J	97	GLN	OE1-CD-NE2	-9.95	112.65	122.60
3	F	640	GLN	OE1-CD-NE2	-9.95	112.65	122.60
3	L	640	GLN	OE1-CD-NE2	-9.95	112.65	122.60

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Mol	Chain	Res	Type	Atoms	Z	Observed(^{o})	Ideal(^{o})
3	P	640	GLN	OE1-CD-NE2	-9.92	112.68	122.60
3	H	97	GLN	OE1-CD-NE2	-9.91	112.69	122.60
3	F	97	GLN	OE1-CD-NE2	-9.90	112.70	122.60
3	R	640	GLN	OE1-CD-NE2	-9.89	112.71	122.60
3	R	423	GLN	OE1-CD-NE2	-9.88	112.72	122.60
3	B	640	GLN	OE1-CD-NE2	-9.88	112.72	122.60
3	H	423	GLN	OE1-CD-NE2	-9.87	112.73	122.60
3	H	640	GLN	OE1-CD-NE2	-9.87	112.73	122.60
3	L	423	GLN	OE1-CD-NE2	-9.87	112.73	122.60
3	J	554	GLN	OE1-CD-NE2	-9.87	112.73	122.60
3	N	554	GLN	OE1-CD-NE2	-9.87	112.73	122.60
3	L	554	GLN	OE1-CD-NE2	-9.87	112.73	122.60
3	P	231	GLN	OE1-CD-NE2	-9.86	112.74	122.60
3	T	231	GLN	OE1-CD-NE2	-9.86	112.74	122.60
3	D	554	GLN	OE1-CD-NE2	-9.86	112.74	122.60
3	N	640	GLN	OE1-CD-NE2	-9.86	112.74	122.60
3	J	640	GLN	OE1-CD-NE2	-9.86	112.74	122.60
3	P	554	GLN	OE1-CD-NE2	-9.86	112.75	122.60
3	T	554	GLN	OE1-CD-NE2	-9.85	112.75	122.60
3	B	554	GLN	OE1-CD-NE2	-9.85	112.75	122.60
3	P	423	GLN	OE1-CD-NE2	-9.85	112.75	122.60
3	N	423	GLN	OE1-CD-NE2	-9.85	112.75	122.60
3	B	231	GLN	OE1-CD-NE2	-9.84	112.76	122.60
2	K	5	GLN	OE1-CD-NE2	-9.84	112.76	122.60
3	T	423	GLN	OE1-CD-NE2	-9.84	112.76	122.60
3	B	423	GLN	OE1-CD-NE2	-9.84	112.76	122.60
3	F	423	GLN	OE1-CD-NE2	-9.83	112.77	122.60
3	H	554	GLN	OE1-CD-NE2	-9.83	112.77	122.60
3	L	231	GLN	OE1-CD-NE2	-9.83	112.77	122.60
3	T	640	GLN	OE1-CD-NE2	-9.83	112.77	122.60
3	D	640	GLN	OE1-CD-NE2	-9.82	112.78	122.60
3	F	554	GLN	OE1-CD-NE2	-9.82	112.78	122.60
2	E	5	GLN	OE1-CD-NE2	-9.82	112.78	122.60
3	F	231	GLN	OE1-CD-NE2	-9.82	112.78	122.60
3	J	231	GLN	OE1-CD-NE2	-9.82	112.78	122.60
3	N	231	GLN	OE1-CD-NE2	-9.81	112.78	122.60
3	R	554	GLN	OE1-CD-NE2	-9.81	112.79	122.60
3	P	330	GLN	OE1-CD-NE2	-9.81	112.79	122.60
3	D	330	GLN	OE1-CD-NE2	-9.81	112.79	122.60
3	J	423	GLN	OE1-CD-NE2	-9.81	112.79	122.60
2	Q	5	GLN	OE1-CD-NE2	-9.81	112.79	122.60
3	R	231	GLN	OE1-CD-NE2	-9.80	112.80	122.60

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Mol	Chain	Res	Type	Atoms	Z	Observed(^{o})	Ideal(^{o})
3	J	330	GLN	OE1-CD-NE2	-9.80	112.80	122.60
3	R	330	GLN	OE1-CD-NE2	-9.80	112.80	122.60
6	X	635	GLN	OE1-CD-NE2	-9.80	112.80	122.60
3	D	231	GLN	OE1-CD-NE2	-9.79	112.81	122.60
3	H	231	GLN	OE1-CD-NE2	-9.79	112.81	122.60
3	B	330	GLN	OE1-CD-NE2	-9.78	112.82	122.60
3	D	423	GLN	OE1-CD-NE2	-9.78	112.82	122.60
6	X	1109	PRO	N-CA-C	-9.77	102.48	114.20
2	C	5	GLN	OE1-CD-NE2	-9.76	112.84	122.60
2	I	5	GLN	OE1-CD-NE2	-9.75	112.85	122.60
2	O	5	GLN	OE1-CD-NE2	-9.75	112.85	122.60
3	L	330	GLN	OE1-CD-NE2	-9.74	112.86	122.60
2	A	5	GLN	OE1-CD-NE2	-9.74	112.86	122.60
3	F	330	GLN	OE1-CD-NE2	-9.73	112.87	122.60
3	H	330	GLN	OE1-CD-NE2	-9.73	112.87	122.60
3	N	330	GLN	OE1-CD-NE2	-9.71	112.89	122.60
6	X	634	GLN	OE1-CD-NE2	-9.71	112.89	122.60
3	T	330	GLN	OE1-CD-NE2	-9.69	112.91	122.60
6	X	505	THR	N-CA-C	-9.65	101.67	112.57
3	F	414	GLN	OE1-CD-NE2	-9.63	112.97	122.60
3	T	628	GLN	OE1-CD-NE2	-9.63	112.97	122.60
6	X	401	THR	N-CA-C	-9.61	101.49	113.02
3	N	414	GLN	OE1-CD-NE2	-9.60	113.00	122.60
3	N	628	GLN	OE1-CD-NE2	-9.60	113.00	122.60
3	P	414	GLN	OE1-CD-NE2	-9.60	113.00	122.60
3	L	414	GLN	OE1-CD-NE2	-9.59	113.01	122.60
3	B	628	GLN	OE1-CD-NE2	-9.59	113.01	122.60
3	H	628	GLN	OE1-CD-NE2	-9.59	113.01	122.60
3	T	414	GLN	OE1-CD-NE2	-9.59	113.01	122.60
3	D	628	GLN	OE1-CD-NE2	-9.59	113.02	122.60
3	P	628	GLN	OE1-CD-NE2	-9.59	113.02	122.60
3	H	414	GLN	OE1-CD-NE2	-9.58	113.02	122.60
3	J	628	GLN	OE1-CD-NE2	-9.57	113.03	122.60
3	R	628	GLN	OE1-CD-NE2	-9.57	113.03	122.60
3	R	414	GLN	OE1-CD-NE2	-9.56	113.04	122.60
3	B	414	GLN	OE1-CD-NE2	-9.55	113.05	122.60
3	D	414	GLN	OE1-CD-NE2	-9.51	113.09	122.60
3	J	414	GLN	OE1-CD-NE2	-9.51	113.09	122.60
3	L	628	GLN	OE1-CD-NE2	-9.50	113.10	122.60
3	F	628	GLN	OE1-CD-NE2	-9.46	113.14	122.60
6	X	302	LEU	N-CA-C	-9.32	101.12	111.28
3	D	93	GLN	OE1-CD-NE2	-9.31	113.29	122.60

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Mol	Chain	Res	Type	Atoms	Z	Observed(^{o})	Ideal(^{o})
3	R	93	GLN	OE1-CD-NE2	-9.31	113.29	122.60
3	L	93	GLN	OE1-CD-NE2	-9.30	113.30	122.60
3	J	93	GLN	OE1-CD-NE2	-9.27	113.33	122.60
3	F	93	GLN	OE1-CD-NE2	-9.27	113.33	122.60
3	B	93	GLN	OE1-CD-NE2	-9.27	113.33	122.60
3	H	93	GLN	OE1-CD-NE2	-9.27	113.33	122.60
3	P	93	GLN	OE1-CD-NE2	-9.26	113.34	122.60
3	N	93	GLN	OE1-CD-NE2	-9.25	113.35	122.60
3	T	93	GLN	OE1-CD-NE2	-9.23	113.37	122.60
3	H	206	GLU	N-CA-C	-9.17	103.20	114.75
3	L	206	GLU	N-CA-C	-9.15	103.22	114.75
3	F	206	GLU	N-CA-C	-9.14	103.23	114.75
3	N	206	GLU	N-CA-C	-9.14	103.24	114.75
3	T	206	GLU	N-CA-C	-9.13	103.24	114.75
3	P	206	GLU	N-CA-C	-9.13	103.25	114.75
3	R	206	GLU	N-CA-C	-9.13	103.25	114.75
3	B	206	GLU	N-CA-C	-9.12	103.27	114.75
3	J	206	GLU	N-CA-C	-9.10	103.29	114.75
3	D	206	GLU	N-CA-C	-9.09	103.30	114.75
6	Y	1041	ILE	N-CA-C	8.73	118.81	110.42
1	f	76	ASN	N-CA-C	8.63	122.05	108.67
1	r	76	ASN	N-CA-C	8.63	122.05	108.67
6	X	616	SER	N-CA-C	8.57	121.76	111.82
1	l	18	ALA	CA-C-N	8.40	137.58	121.54
1	l	18	ALA	C-N-CA	8.40	137.58	121.54
5	W	62	TYR	CA-C-N	8.30	128.93	120.38
5	W	62	TYR	C-N-CA	8.30	128.93	120.38
1	r	74	GLN	N-CA-C	8.28	121.40	109.48
4	U	171	THR	N-CA-C	-8.04	103.61	113.50
6	X	793	ARG	N-CA-C	-7.92	97.69	110.20
1	f	72	SER	CA-C-N	7.75	130.67	120.28
1	f	72	SER	C-N-CA	7.75	130.67	120.28
6	Y	1186	ASN	N-CA-C	7.72	122.77	113.12
6	Y	1102	GLU	N-CA-C	-7.71	102.00	111.33
1	f	75	PRO	N-CA-CB	-7.64	95.23	103.25
3	F	332	GLY	CA-C-N	7.55	129.28	119.84
3	F	332	GLY	C-N-CA	7.55	129.28	119.84
3	L	332	GLY	CA-C-N	7.54	129.27	119.84
3	L	332	GLY	C-N-CA	7.54	129.27	119.84
3	D	332	GLY	CA-C-N	7.54	129.27	119.84
3	D	332	GLY	C-N-CA	7.54	129.27	119.84
6	X	649	ASN	N-CA-C	7.54	122.21	112.41

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Mol	Chain	Res	Type	Atoms	Z	Observed(^{o})	Ideal(^{o})
3	P	332	GLY	CA-C-N	7.53	129.25	119.84
3	P	332	GLY	C-N-CA	7.53	129.25	119.84
3	B	332	GLY	CA-C-N	7.51	129.23	119.84
3	B	332	GLY	C-N-CA	7.51	129.23	119.84
3	R	332	GLY	CA-C-N	7.51	129.22	119.84
3	R	332	GLY	C-N-CA	7.51	129.22	119.84
3	J	332	GLY	CA-C-N	7.50	129.22	119.84
3	J	332	GLY	C-N-CA	7.50	129.22	119.84
3	T	332	GLY	CA-C-N	7.50	129.21	119.84
3	T	332	GLY	C-N-CA	7.50	129.21	119.84
6	X	768	GLN	N-CA-C	7.50	121.76	110.14
3	H	332	GLY	CA-C-N	7.49	129.21	119.84
3	H	332	GLY	C-N-CA	7.49	129.21	119.84
3	N	332	GLY	CA-C-N	7.47	129.18	119.84
3	N	332	GLY	C-N-CA	7.47	129.18	119.84
6	Y	119	CYS	N-CA-C	-7.45	97.91	109.76
6	X	445	PHE	N-CA-C	7.45	121.58	109.59
6	Y	669	PHE	N-CA-C	7.32	119.04	111.14
4	U	380	ALA	N-CA-C	-7.30	104.25	113.01
1	j	74	GLN	N-CA-C	7.26	120.16	109.42
3	T	554	GLN	CG-CD-NE2	7.25	127.28	116.40
3	H	554	GLN	CG-CD-NE2	7.25	127.27	116.40
3	N	554	GLN	CG-CD-NE2	7.25	127.27	116.40
3	B	554	GLN	CG-CD-NE2	7.23	127.25	116.40
3	P	554	GLN	CG-CD-NE2	7.22	127.23	116.40
3	J	554	GLN	CG-CD-NE2	7.21	127.22	116.40
3	D	554	GLN	CG-CD-NE2	7.21	127.21	116.40
3	L	554	GLN	CG-CD-NE2	7.20	127.20	116.40
6	X	456	VAL	N-CA-C	-7.20	103.45	110.72
3	F	554	GLN	CG-CD-NE2	7.19	127.18	116.40
3	R	554	GLN	CG-CD-NE2	7.19	127.18	116.40
6	Y	1002	PRO	N-CA-CB	7.16	110.02	103.08
6	X	681	ASN	N-CA-C	7.11	121.03	109.59
6	X	958	HIS	N-CA-C	-7.08	103.60	113.37
6	X	427	LEU	N-CA-C	-7.07	100.53	110.50
6	X	849	ALA	N-CA-C	-7.06	103.34	112.23
3	F	551	GLN	CG-CD-NE2	7.06	126.98	116.40
6	X	1172	LEU	N-CA-C	7.04	119.98	111.82
3	L	640	GLN	CG-CD-NE2	7.04	126.96	116.40
3	R	640	GLN	CG-CD-NE2	7.03	126.95	116.40
3	L	551	GLN	CG-CD-NE2	7.03	126.95	116.40
6	X	1041	ILE	N-CA-C	7.03	118.63	110.62

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Mol	Chain	Res	Type	Atoms	Z	Observed(^{o})	Ideal(^{o})
6	Y	530	LYS	N-CA-C	-7.03	104.63	112.57
3	J	551	GLN	CG-CD-NE2	7.02	126.92	116.40
3	P	640	GLN	CG-CD-NE2	7.02	126.93	116.40
3	N	640	GLN	CG-CD-NE2	7.02	126.92	116.40
3	T	551	GLN	CG-CD-NE2	7.02	126.92	116.40
3	T	640	GLN	CG-CD-NE2	7.02	126.92	116.40
3	B	551	GLN	CG-CD-NE2	7.01	126.92	116.40
3	F	640	GLN	CG-CD-NE2	7.01	126.92	116.40
3	B	640	GLN	CG-CD-NE2	7.01	126.92	116.40
3	H	551	GLN	CG-CD-NE2	7.01	126.91	116.40
6	Y	533	ALA	N-CA-C	-7.01	104.60	113.01
3	D	551	GLN	CG-CD-NE2	7.01	126.91	116.40
3	H	640	GLN	CG-CD-NE2	7.00	126.90	116.40
3	N	551	GLN	CG-CD-NE2	7.00	126.90	116.40
3	J	640	GLN	CG-CD-NE2	7.00	126.90	116.40
6	Y	1035	GLY	N-CA-C	-7.00	106.64	115.31
5	W	540	GLY	N-CA-C	-7.00	105.23	115.30
3	P	551	GLN	CG-CD-NE2	6.99	126.89	116.40
3	D	640	GLN	CG-CD-NE2	6.99	126.89	116.40
3	R	551	GLN	CG-CD-NE2	6.99	126.88	116.40
6	X	709	ILE	N-CA-C	-6.93	102.84	113.16
3	P	207	PHE	N-CA-C	-6.86	104.55	113.12
1	b	19	ALA	N-CA-C	-6.86	101.15	110.68
3	L	207	PHE	N-CA-C	-6.86	104.55	113.12
3	F	207	PHE	N-CA-C	-6.85	104.55	113.12
1	f	22	LEU	CA-C-N	6.85	134.63	121.54
1	f	22	LEU	C-N-CA	6.85	134.63	121.54
3	J	207	PHE	N-CA-C	-6.85	104.56	113.12
6	X	777	ASN	N-CA-C	6.85	119.76	111.82
5	W	1001	ILE	N-CA-C	-6.84	106.21	111.62
3	N	207	PHE	N-CA-C	-6.84	104.57	113.12
3	N	231	GLN	CG-CD-NE2	6.83	126.65	116.40
3	R	207	PHE	N-CA-C	-6.83	104.58	113.12
3	H	231	GLN	CG-CD-NE2	6.83	126.65	116.40
3	B	207	PHE	N-CA-C	-6.83	104.58	113.12
3	H	207	PHE	N-CA-C	-6.83	104.59	113.12
3	R	231	GLN	CG-CD-NE2	6.82	126.63	116.40
3	D	231	GLN	CG-CD-NE2	6.82	126.63	116.40
3	D	207	PHE	N-CA-C	-6.81	104.60	113.12
3	N	160	GLN	CG-CD-NE2	6.81	126.62	116.40
3	L	231	GLN	CG-CD-NE2	6.81	126.62	116.40
3	B	231	GLN	CG-CD-NE2	6.81	126.61	116.40

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Mol	Chain	Res	Type	Atoms	Z	Observed(^{o})	Ideal(^{o})
3	P	231	GLN	CG-CD-NE2	6.81	126.61	116.40
3	J	231	GLN	CG-CD-NE2	6.81	126.61	116.40
3	H	160	GLN	CG-CD-NE2	6.80	126.60	116.40
3	T	231	GLN	CG-CD-NE2	6.80	126.60	116.40
6	Y	475	ALA	N-CA-C	-6.80	103.87	111.28
3	F	231	GLN	CG-CD-NE2	6.80	126.60	116.40
3	R	160	GLN	CG-CD-NE2	6.78	126.58	116.40
3	T	160	GLN	CG-CD-NE2	6.78	126.58	116.40
3	D	160	GLN	CG-CD-NE2	6.78	126.57	116.40
3	T	207	PHE	N-CA-C	-6.77	104.65	113.12
3	J	160	GLN	CG-CD-NE2	6.77	126.55	116.40
3	B	160	GLN	CG-CD-NE2	6.76	126.54	116.40
3	R	97	GLN	CG-CD-NE2	6.76	126.54	116.40
6	X	214	HIS	N-CA-C	-6.75	99.67	109.59
3	L	97	GLN	CG-CD-NE2	6.75	126.52	116.40
3	N	97	GLN	CG-CD-NE2	6.75	126.52	116.40
3	P	160	GLN	CG-CD-NE2	6.75	126.52	116.40
3	F	160	GLN	CG-CD-NE2	6.75	126.52	116.40
3	L	160	GLN	CG-CD-NE2	6.74	126.52	116.40
3	D	498	VAL	CB-CA-C	-6.74	105.98	111.71
3	T	97	GLN	CG-CD-NE2	6.74	126.50	116.40
3	B	97	GLN	CG-CD-NE2	6.73	126.50	116.40
6	X	955	SER	N-CA-C	6.73	120.44	110.48
3	J	97	GLN	CG-CD-NE2	6.73	126.49	116.40
3	T	423	GLN	CG-CD-NE2	6.73	126.49	116.40
3	F	97	GLN	CG-CD-NE2	6.72	126.48	116.40
6	X	403	ILE	N-CA-C	-6.72	103.76	110.62
3	N	423	GLN	CG-CD-NE2	6.71	126.47	116.40
3	P	97	GLN	CG-CD-NE2	6.71	126.47	116.40
3	R	423	GLN	CG-CD-NE2	6.71	126.47	116.40
3	H	97	GLN	CG-CD-NE2	6.71	126.47	116.40
3	D	97	GLN	CG-CD-NE2	6.71	126.46	116.40
3	H	498	VAL	CB-CA-C	-6.71	106.01	111.71
3	H	423	GLN	CG-CD-NE2	6.70	126.46	116.40
3	P	423	GLN	CG-CD-NE2	6.70	126.45	116.40
3	N	498	VAL	CB-CA-C	-6.70	106.02	111.71
3	B	423	GLN	CG-CD-NE2	6.69	126.44	116.40
3	L	423	GLN	CG-CD-NE2	6.69	126.44	116.40
3	J	498	VAL	CB-CA-C	-6.69	106.03	111.71
3	P	498	VAL	CB-CA-C	-6.69	106.03	111.71
3	F	423	GLN	CG-CD-NE2	6.67	126.41	116.40
6	X	635	GLN	CG-CD-NE2	6.67	126.41	116.40

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Mol	Chain	Res	Type	Atoms	Z	Observed(^{o})	Ideal(^{o})
3	J	330	GLN	CG-CD-NE2	6.67	126.41	116.40
3	T	498	VAL	CB-CA-C	-6.67	106.04	111.71
3	D	423	GLN	CG-CD-NE2	6.67	126.40	116.40
3	P	330	GLN	CG-CD-NE2	6.67	126.40	116.40
3	J	423	GLN	CG-CD-NE2	6.67	126.40	116.40
3	R	498	VAL	CB-CA-C	-6.66	106.05	111.71
3	B	330	GLN	CG-CD-NE2	6.66	126.38	116.40
3	B	498	VAL	CB-CA-C	-6.65	106.06	111.71
3	D	330	GLN	CG-CD-NE2	6.65	126.37	116.40
3	R	330	GLN	CG-CD-NE2	6.64	126.36	116.40
2	I	5	GLN	CG-CD-NE2	6.64	126.36	116.40
3	H	330	GLN	CG-CD-NE2	6.64	126.36	116.40
3	L	493	GLN	CG-CD-NE2	6.64	126.36	116.40
3	L	498	VAL	CB-CA-C	-6.64	106.07	111.71
6	Y	532	ILE	N-CA-C	-6.64	104.55	111.58
3	F	493	GLN	CG-CD-NE2	6.63	126.35	116.40
3	T	330	GLN	CG-CD-NE2	6.63	126.34	116.40
3	P	493	GLN	CG-CD-NE2	6.62	126.34	116.40
2	K	5	GLN	CG-CD-NE2	6.62	126.33	116.40
2	A	5	GLN	CG-CD-NE2	6.62	126.33	116.40
2	C	5	GLN	CG-CD-NE2	6.62	126.33	116.40
3	R	493	GLN	CG-CD-NE2	6.62	126.33	116.40
5	W	404	TYR	N-CA-C	6.62	119.49	110.55
3	F	498	VAL	CB-CA-C	-6.62	106.08	111.71
2	O	5	GLN	CG-CD-NE2	6.62	126.33	116.40
3	N	330	GLN	CG-CD-NE2	6.62	126.33	116.40
2	E	5	GLN	CG-CD-NE2	6.62	126.32	116.40
3	F	330	GLN	CG-CD-NE2	6.61	126.32	116.40
3	L	330	GLN	CG-CD-NE2	6.61	126.31	116.40
3	J	493	GLN	CG-CD-NE2	6.60	126.30	116.40
3	T	493	GLN	CG-CD-NE2	6.60	126.30	116.40
3	B	493	GLN	CG-CD-NE2	6.60	126.30	116.40
3	H	493	GLN	CG-CD-NE2	6.60	126.30	116.40
3	N	493	GLN	CG-CD-NE2	6.60	126.30	116.40
2	Q	5	GLN	CG-CD-NE2	6.60	126.30	116.40
6	X	634	GLN	CG-CD-NE2	6.59	126.28	116.40
3	D	493	GLN	CG-CD-NE2	6.58	126.27	116.40
6	Y	540	LEU	N-CA-C	-6.56	105.85	114.31
5	W	133	ILE	N-CA-C	6.55	117.30	110.62
3	F	414	GLN	CG-CD-NE2	6.54	126.21	116.40
3	P	414	GLN	CG-CD-NE2	6.53	126.19	116.40
6	X	630	LEU	N-CA-C	6.53	118.05	111.07

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Mol	Chain	Res	Type	Atoms	Z	Observed(^{o})	Ideal(^{o})
3	L	414	GLN	CG-CD-NE2	6.51	126.17	116.40
1	l	19	ALA	N-CA-C	-6.49	96.98	110.80
3	T	414	GLN	CG-CD-NE2	6.48	126.12	116.40
3	H	414	GLN	CG-CD-NE2	6.48	126.12	116.40
5	W	1040	VAL	CA-C-N	-6.48	113.17	121.91
5	W	1040	VAL	C-N-CA	-6.48	113.17	121.91
3	N	414	GLN	CG-CD-NE2	6.48	126.11	116.40
3	B	414	GLN	CG-CD-NE2	6.47	126.11	116.40
3	R	414	GLN	CG-CD-NE2	6.47	126.11	116.40
1	r	74	GLN	CA-C-N	6.47	127.92	119.84
1	r	74	GLN	C-N-CA	6.47	127.92	119.84
3	D	414	GLN	CG-CD-NE2	6.47	126.10	116.40
4	U	186	ALA	N-CA-C	-6.47	104.27	111.71
3	J	414	GLN	CG-CD-NE2	6.46	126.09	116.40
5	W	626	ALA	N-CA-C	-6.45	105.95	113.88
3	D	93	GLN	CG-CD-NE2	6.44	126.06	116.40
2	O	25	ASP	N-CA-C	6.44	119.29	111.82
5	W	607	LEU	N-CA-C	-6.42	105.74	112.93
6	X	437	ARG	CA-C-N	6.42	127.86	119.84
6	X	437	ARG	C-N-CA	6.42	127.86	119.84
2	A	25	ASP	N-CA-C	6.41	119.26	111.82
2	S	25	ASP	N-CA-C	6.41	119.26	111.82
3	J	93	GLN	CG-CD-NE2	6.41	126.01	116.40
3	J	628	GLN	CG-CD-NE2	6.41	126.01	116.40
3	T	203	TYR	CA-C-N	6.40	127.84	119.84
3	T	203	TYR	C-N-CA	6.40	127.84	119.84
3	T	628	GLN	CG-CD-NE2	6.40	126.00	116.40
3	T	93	GLN	CG-CD-NE2	6.40	126.00	116.40
6	X	966	SER	N-CA-C	-6.40	104.23	111.14
3	H	93	GLN	CG-CD-NE2	6.40	126.00	116.40
2	M	25	ASP	N-CA-C	6.40	119.24	111.82
3	N	203	TYR	CA-C-N	6.40	127.84	119.84
3	N	203	TYR	C-N-CA	6.40	127.84	119.84
3	B	628	GLN	CG-CD-NE2	6.39	125.99	116.40
3	R	203	TYR	CA-C-N	6.39	127.83	119.84
3	R	203	TYR	C-N-CA	6.39	127.83	119.84
3	D	628	GLN	CG-CD-NE2	6.39	125.99	116.40
2	E	25	ASP	N-CA-C	6.39	119.23	111.82
6	Y	260	GLU	N-CA-C	6.39	116.89	108.07
3	L	93	GLN	CG-CD-NE2	6.39	125.99	116.40
3	P	628	GLN	CG-CD-NE2	6.39	125.99	116.40
3	H	628	GLN	CG-CD-NE2	6.39	125.98	116.40

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Mol	Chain	Res	Type	Atoms	Z	Observed(^{o})	Ideal(^{o})
3	B	93	GLN	CG-CD-NE2	6.39	125.98	116.40
2	G	25	ASP	N-CA-C	6.39	119.23	111.82
3	F	93	GLN	CG-CD-NE2	6.39	125.98	116.40
2	I	25	ASP	N-CA-C	6.39	119.23	111.82
3	F	203	TYR	CA-C-N	6.38	127.82	119.84
3	F	203	TYR	C-N-CA	6.38	127.82	119.84
3	N	93	GLN	CG-CD-NE2	6.38	125.98	116.40
2	C	25	ASP	N-CA-C	6.38	119.22	111.82
3	L	203	TYR	CA-C-N	6.38	127.81	119.84
3	L	203	TYR	C-N-CA	6.38	127.81	119.84
3	N	628	GLN	CG-CD-NE2	6.38	125.96	116.40
3	P	93	GLN	CG-CD-NE2	6.38	125.97	116.40
3	R	93	GLN	CG-CD-NE2	6.38	125.96	116.40
3	B	203	TYR	CA-C-N	6.37	127.81	119.84
3	B	203	TYR	C-N-CA	6.37	127.81	119.84
3	D	203	TYR	CA-C-N	6.37	127.81	119.84
3	D	203	TYR	C-N-CA	6.37	127.81	119.84
2	K	25	ASP	N-CA-C	6.37	119.21	111.82
3	R	628	GLN	CG-CD-NE2	6.37	125.95	116.40
3	J	203	TYR	CA-C-N	6.37	127.80	119.84
3	J	203	TYR	C-N-CA	6.37	127.80	119.84
3	H	203	TYR	CA-C-N	6.36	127.79	119.84
3	H	203	TYR	C-N-CA	6.36	127.79	119.84
2	Q	25	ASP	N-CA-C	6.36	119.19	111.82
6	Y	1003	PRO	N-CA-CB	6.36	109.93	103.25
6	Y	1158	PHE	N-CA-C	6.35	124.33	110.80
6	X	429	ASN	N-CA-C	6.35	118.29	109.15
3	L	628	GLN	CG-CD-NE2	6.32	125.88	116.40
3	P	203	TYR	CA-C-N	6.32	127.73	119.84
3	P	203	TYR	C-N-CA	6.32	127.73	119.84
4	U	148	TRP	N-CA-C	-6.30	106.14	113.88
3	F	628	GLN	CG-CD-NE2	6.29	125.84	116.40
6	X	372	ASP	N-CA-C	6.29	119.65	107.71
6	Y	852	THR	N-CA-C	-6.28	104.44	111.28
6	Y	121	VAL	N-CA-C	-6.25	104.54	110.42
6	Y	668	LEU	N-CA-C	6.23	120.70	111.04
5	W	664	SER	N-CA-C	6.23	118.86	111.33
5	W	829	ALA	CB-CA-C	-6.16	109.48	116.63
1	f	74	GLN	CA-C-N	6.14	127.52	119.84
1	f	74	GLN	C-N-CA	6.14	127.52	119.84
5	W	197	HIS	N-CA-C	-6.12	105.08	112.54
5	W	686	ALA	N-CA-C	-6.10	104.19	111.69

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Mol	Chain	Res	Type	Atoms	Z	Observed(^{o})	Ideal(^{o})
6	X	402	TYR	N-CA-C	-6.09	105.88	113.55
1	f	21	ARG	N-CA-C	-6.08	103.97	112.12
5	W	10	VAL	CA-C-N	6.05	127.41	119.84
5	W	10	VAL	C-N-CA	6.05	127.41	119.84
4	V	158	HIS	N-CA-C	6.03	118.10	107.61
6	X	329	MET	N-CA-C	6.01	119.37	112.57
2	Q	17	GLY	N-CA-C	6.00	122.24	113.48
2	E	17	GLY	N-CA-C	5.99	122.23	113.48
2	M	17	GLY	N-CA-C	5.99	122.22	113.48
2	A	17	GLY	N-CA-C	5.98	122.22	113.48
4	U	166	MET	CA-C-N	5.98	127.32	119.84
4	U	166	MET	C-N-CA	5.98	127.32	119.84
2	K	17	GLY	N-CA-C	5.98	122.21	113.48
2	C	17	GLY	N-CA-C	5.97	122.19	113.48
2	G	17	GLY	N-CA-C	5.97	122.19	113.48
2	I	17	GLY	N-CA-C	5.96	122.18	113.48
2	O	17	GLY	N-CA-C	5.96	122.18	113.48
2	S	17	GLY	N-CA-C	5.96	122.18	113.48
5	W	1134	VAL	N-CA-C	5.95	117.47	107.18
1	p	58	PRO	CA-C-N	5.94	132.89	121.54
1	p	58	PRO	C-N-CA	5.94	132.89	121.54
6	Y	953	GLY	CA-C-N	5.94	127.27	119.84
6	Y	953	GLY	C-N-CA	5.94	127.27	119.84
6	X	558	THR	N-CA-C	-5.93	105.54	112.89
6	X	597	GLU	N-CA-C	5.92	120.18	108.59
3	P	142	ILE	N-CA-C	-5.91	105.26	112.76
3	N	142	ILE	N-CA-C	-5.90	105.27	112.76
3	B	142	ILE	N-CA-C	-5.89	105.28	112.76
3	T	142	ILE	N-CA-C	-5.89	105.28	112.76
6	X	1193	SER	N-CA-C	-5.89	101.29	110.42
3	D	142	ILE	N-CA-C	-5.89	105.28	112.76
3	L	142	ILE	N-CA-C	-5.89	105.28	112.76
6	X	387	CYS	N-CA-C	5.89	118.06	108.76
3	J	142	ILE	N-CA-C	-5.88	105.29	112.76
6	Y	1069	ALA	N-CA-C	-5.88	100.41	109.76
3	H	142	ILE	N-CA-C	-5.88	105.30	112.76
1	f	73	ARG	CA-C-O	-5.87	114.33	120.55
6	X	287	GLN	N-CA-C	5.87	119.46	110.36
3	H	69	GLY	N-CA-C	-5.87	104.48	112.52
6	Y	46	ALA	N-CA-C	-5.87	104.07	111.11
6	Y	113	ASN	CA-C-N	5.87	125.54	119.56
6	Y	113	ASN	C-N-CA	5.87	125.54	119.56

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Mol	Chain	Res	Type	Atoms	Z	Observed(^{o})	Ideal(^{o})
3	F	471	LEU	N-CA-C	5.86	118.15	111.11
3	R	142	ILE	N-CA-C	-5.86	105.32	112.76
5	W	629	THR	N-CA-C	-5.85	105.80	113.12
3	F	142	ILE	N-CA-C	-5.84	105.34	112.76
3	L	471	LEU	N-CA-C	5.84	118.12	111.11
5	W	495	ASP	N-CA-C	-5.84	104.92	111.28
3	N	69	GLY	N-CA-C	-5.84	104.52	112.52
3	R	471	LEU	N-CA-C	5.83	118.10	111.11
6	Y	445	PHE	N-CA-C	5.83	118.45	109.07
3	R	69	GLY	N-CA-C	-5.82	104.55	112.52
6	Y	1008	GLY	CA-C-N	5.81	126.00	119.90
6	Y	1008	GLY	C-N-CA	5.81	126.00	119.90
3	B	69	GLY	N-CA-C	-5.81	104.56	112.52
3	F	297	ALA	N-CA-C	-5.81	106.58	113.38
3	L	69	GLY	N-CA-C	-5.81	104.56	112.52
5	W	817	SER	N-CA-C	-5.81	106.32	113.41
3	L	297	ALA	N-CA-C	-5.80	106.59	113.38
3	T	69	GLY	N-CA-C	-5.80	104.58	112.52
3	T	471	LEU	N-CA-C	5.79	118.06	111.11
6	Y	768	GLN	N-CA-C	5.79	117.69	107.61
3	J	69	GLY	N-CA-C	-5.79	104.58	112.52
5	W	838	PRO	N-CA-C	-5.79	105.83	113.65
3	T	297	ALA	N-CA-C	-5.79	106.61	113.38
3	F	69	GLY	N-CA-C	-5.79	104.59	112.52
3	N	471	LEU	N-CA-C	5.79	118.06	111.11
3	R	297	ALA	N-CA-C	-5.79	106.61	113.38
3	B	471	LEU	N-CA-C	5.78	118.05	111.11
3	J	471	LEU	N-CA-C	5.77	118.04	111.11
3	P	69	GLY	N-CA-C	-5.77	104.61	112.52
6	X	569	LEU	N-CA-C	-5.77	104.36	111.40
3	D	471	LEU	N-CA-C	5.77	118.03	111.11
3	H	471	LEU	N-CA-C	5.76	118.03	111.11
3	P	471	LEU	N-CA-C	5.76	118.03	111.11
3	B	297	ALA	N-CA-C	-5.76	106.64	113.38
3	D	297	ALA	N-CA-C	-5.75	106.65	113.38
1	f	60	ALA	CA-C-N	5.75	132.53	121.54
1	f	60	ALA	C-N-CA	5.75	132.53	121.54
6	X	856	ARG	N-CA-C	-5.75	104.39	111.75
5	W	5	PHE	CB-CA-C	-5.75	109.96	116.63
1	h	58	PRO	N-CA-C	-5.75	105.70	113.40
3	P	635	GLN	N-CA-C	-5.75	105.10	111.36
3	J	297	ALA	N-CA-C	-5.74	106.66	113.38

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Mol	Chain	Res	Type	Atoms	Z	Observed(^{o})	Ideal(^{o})
3	P	297	ALA	N-CA-C	-5.74	106.67	113.38
3	L	635	GLN	N-CA-C	-5.74	105.11	111.36
3	R	635	GLN	N-CA-C	-5.73	105.11	111.36
1	f	75	PRO	CA-N-CD	-5.73	103.98	112.00
3	F	635	GLN	N-CA-C	-5.73	105.12	111.36
3	N	297	ALA	N-CA-C	-5.73	106.68	113.38
3	H	297	ALA	N-CA-C	-5.72	106.68	113.38
5	W	336	PHE	N-CA-C	-5.72	105.97	113.12
6	Y	1184	LEU	N-CA-C	5.72	122.46	109.81
5	W	236	GLY	N-CA-C	-5.72	100.67	112.34
3	B	635	GLN	N-CA-C	-5.71	105.14	111.36
4	U	230	ASN	N-CA-C	-5.71	106.32	113.28
3	J	635	GLN	N-CA-C	-5.70	105.14	111.36
6	Y	193	LEU	N-CA-C	5.70	119.85	113.01
3	H	173	ASP	N-CA-C	5.67	118.12	108.02
6	X	362	THR	N-CA-C	5.67	117.14	111.07
3	D	635	GLN	N-CA-C	-5.67	105.18	111.36
4	V	393	ASP	N-CA-C	5.67	113.43	108.78
6	X	650	TRP	CA-C-N	5.67	126.93	119.84
6	X	650	TRP	C-N-CA	5.67	126.93	119.84
3	N	173	ASP	N-CA-C	5.67	118.10	108.02
1	r	43	CYS	CA-C-N	5.66	131.89	121.70
1	r	43	CYS	C-N-CA	5.66	131.89	121.70
3	R	173	ASP	N-CA-C	5.66	118.09	108.02
6	X	1082	VAL	N-CA-C	-5.66	100.18	109.12
1	t	43	CYS	CA-C-N	5.65	131.87	121.70
1	t	43	CYS	C-N-CA	5.65	131.87	121.70
3	T	387	THR	N-CA-C	5.65	118.28	109.52
6	X	578	ASP	CA-C-N	5.65	126.90	119.84
6	X	578	ASP	C-N-CA	5.65	126.90	119.84
1	l	43	CYS	CA-C-N	5.65	131.87	121.70
1	l	43	CYS	C-N-CA	5.65	131.87	121.70
6	X	949	HIS	N-CA-C	5.65	117.95	108.23
3	T	635	GLN	N-CA-C	-5.65	105.21	111.36
1	j	43	CYS	CA-C-N	5.64	131.85	121.70
1	j	43	CYS	C-N-CA	5.64	131.85	121.70
1	n	43	CYS	CA-C-N	5.64	131.86	121.70
1	n	43	CYS	C-N-CA	5.64	131.86	121.70
3	H	635	GLN	N-CA-C	-5.64	105.21	111.36
3	L	173	ASP	N-CA-C	5.64	118.06	108.02
6	Y	474	GLU	N-CA-C	-5.64	106.53	113.41
1	d	43	CYS	CA-C-N	5.64	131.85	121.70

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Mol	Chain	Res	Type	Atoms	Z	Observed(^{o})	Ideal(^{o})
1	d	43	CYS	C-N-CA	5.64	131.85	121.70
1	p	43	CYS	CA-C-N	5.64	131.85	121.70
1	p	43	CYS	C-N-CA	5.64	131.85	121.70
4	V	154	GLY	CA-C-N	5.64	126.52	119.98
4	V	154	GLY	C-N-CA	5.64	126.52	119.98
6	Y	403	ILE	N-CA-C	-5.64	104.87	110.62
1	h	43	CYS	CA-C-N	5.64	131.85	121.70
1	h	43	CYS	C-N-CA	5.64	131.85	121.70
3	P	387	THR	N-CA-C	5.64	118.26	109.52
3	B	173	ASP	N-CA-C	5.64	118.05	108.02
3	J	387	THR	N-CA-C	5.64	118.26	109.52
3	T	173	ASP	N-CA-C	5.63	118.05	108.02
1	b	43	CYS	CA-C-N	5.63	131.84	121.70
1	b	43	CYS	C-N-CA	5.63	131.84	121.70
3	N	635	GLN	N-CA-C	-5.63	105.22	111.36
3	P	173	ASP	N-CA-C	5.63	118.03	108.02
6	X	243	ILE	CB-CA-C	-5.63	105.47	112.45
3	B	387	THR	N-CA-C	5.62	118.24	109.52
3	F	173	ASP	N-CA-C	5.62	118.03	108.02
1	f	43	CYS	CA-C-N	5.62	131.82	121.70
1	f	43	CYS	C-N-CA	5.62	131.82	121.70
6	X	395	THR	N-CA-C	-5.62	105.06	111.07
3	D	173	ASP	N-CA-C	5.62	118.02	108.02
5	W	1019	TYR	N-CA-C	-5.62	105.54	112.90
3	N	387	THR	N-CA-C	5.62	118.23	109.52
6	X	1083	TYR	N-CA-C	5.62	118.02	108.02
3	J	173	ASP	N-CA-C	5.61	118.01	108.02
3	R	387	THR	N-CA-C	5.61	118.22	109.52
3	H	387	THR	N-CA-C	5.61	118.22	109.52
3	D	598	SER	N-CA-C	-5.61	105.17	111.28
3	D	387	THR	N-CA-C	5.60	118.20	109.52
5	W	85	THR	N-CA-C	-5.59	105.18	111.28
4	U	154	GLY	CA-C-N	5.59	125.59	119.89
4	U	154	GLY	C-N-CA	5.59	125.59	119.89
3	F	387	THR	N-CA-C	5.58	118.17	109.52
6	Y	1174	ASP	CA-C-N	5.58	125.67	119.87
6	Y	1174	ASP	C-N-CA	5.58	125.67	119.87
5	W	740	VAL	CA-C-N	-5.58	112.87	119.84
5	W	740	VAL	C-N-CA	-5.58	112.87	119.84
3	L	387	THR	N-CA-C	5.58	118.16	109.52
3	F	233	HIS	CA-CB-CG	-5.57	108.23	113.80
5	W	384	THR	N-CA-C	-5.57	101.67	109.69

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Mol	Chain	Res	Type	Atoms	Z	Observed(^{o})	Ideal(^{o})
6	Y	929	PRO	N-CA-C	-5.57	106.90	113.86
3	R	598	SER	N-CA-C	-5.56	105.22	111.28
3	J	598	SER	N-CA-C	-5.55	105.23	111.28
1	n	58	PRO	N-CA-C	-5.55	105.97	113.40
3	L	233	HIS	CA-CB-CG	-5.55	108.25	113.80
3	L	598	SER	N-CA-C	-5.55	105.23	111.28
3	R	119	VAL	N-CA-C	5.55	116.10	107.28
5	W	775	ARG	N-CA-C	-5.54	106.64	113.41
6	Y	1006	ARG	N-CA-C	5.54	117.02	110.97
1	b	21	ARG	N-CA-C	-5.54	99.00	110.80
3	R	233	HIS	CA-CB-CG	-5.54	108.26	113.80
3	T	233	HIS	CA-CB-CG	-5.53	108.27	113.80
3	N	119	VAL	N-CA-C	5.53	116.07	107.28
3	T	598	SER	N-CA-C	-5.52	105.26	111.28
3	B	598	SER	N-CA-C	-5.51	105.27	111.28
3	H	233	HIS	CA-CB-CG	-5.51	108.29	113.80
3	J	119	VAL	N-CA-C	5.51	116.05	107.28
6	X	385	VAL	N-CA-C	-5.51	101.79	109.45
3	N	233	HIS	CA-CB-CG	-5.51	108.29	113.80
3	B	233	HIS	CA-CB-CG	-5.51	108.29	113.80
3	F	598	SER	N-CA-C	-5.50	105.28	111.28
3	H	598	SER	N-CA-C	-5.50	105.28	111.28
3	H	119	VAL	N-CA-C	5.50	116.03	107.28
4	V	60	SER	N-CA-C	5.50	115.81	108.38
3	D	119	VAL	N-CA-C	5.50	116.02	107.28
3	B	119	VAL	N-CA-C	5.50	116.02	107.28
3	D	57	THR	N-CA-C	-5.50	101.21	109.95
3	F	119	VAL	N-CA-C	5.50	116.02	107.28
3	L	119	VAL	N-CA-C	5.49	116.01	107.28
3	N	598	SER	N-CA-C	-5.49	105.29	111.28
3	P	233	HIS	CA-CB-CG	-5.49	108.31	113.80
4	U	257	ALA	CA-C-N	5.49	123.67	119.66
4	U	257	ALA	C-N-CA	5.49	123.67	119.66
1	l	18	ALA	O-C-N	5.49	129.42	122.28
3	T	119	VAL	N-CA-C	5.49	116.01	107.28
6	X	1034	ASP	N-CA-C	5.49	116.95	110.97
3	P	119	VAL	N-CA-C	5.48	115.99	107.28
3	D	233	HIS	CA-CB-CG	-5.48	108.32	113.80
3	J	233	HIS	CA-CB-CG	-5.47	108.33	113.80
3	P	598	SER	N-CA-C	-5.47	105.31	111.28
5	W	1143	GLN	N-CA-C	-5.47	105.20	113.51
6	X	629	GLN	N-CA-C	-5.47	104.72	111.33

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Mol	Chain	Res	Type	Atoms	Z	Observed(^{o})	Ideal(^{o})
4	U	180	VAL	CA-C-N	5.45	125.92	120.14
4	U	180	VAL	C-N-CA	5.45	125.92	120.14
5	W	1139	PRO	CA-C-N	5.45	123.58	119.66
5	W	1139	PRO	C-N-CA	5.45	123.58	119.66
3	T	342	ASN	N-CA-C	-5.44	103.35	110.53
3	F	342	ASN	N-CA-C	-5.43	103.36	110.53
6	Y	453	LEU	N-CA-C	-5.43	106.61	113.18
3	R	342	ASN	N-CA-C	-5.42	103.37	110.53
3	L	342	ASN	N-CA-C	-5.42	103.38	110.53
6	X	1173	VAL	N-CA-C	5.42	118.40	113.20
3	B	342	ASN	N-CA-C	-5.41	103.39	110.53
3	D	210	VAL	N-CA-C	-5.41	105.45	110.53
3	D	342	ASN	N-CA-C	-5.40	103.40	110.53
5	W	244	THR	N-CA-C	5.40	117.25	111.36
1	j	73	ARG	N-CA-C	5.40	117.25	111.36
3	N	342	ASN	N-CA-C	-5.39	103.41	110.53
3	P	342	ASN	N-CA-C	-5.39	103.41	110.53
1	r	21	ARG	N-CA-C	-5.39	99.32	110.80
1	l	21	ARG	N-CA-C	-5.39	106.72	113.72
3	J	342	ASN	N-CA-C	-5.39	103.42	110.53
1	j	19	ALA	N-CA-C	-5.38	99.75	108.52
6	X	472	LEU	N-CA-C	-5.38	107.37	114.04
2	C	29	THR	N-CA-C	5.38	117.14	108.32
3	H	342	ASN	N-CA-C	-5.37	103.45	110.53
2	O	29	THR	N-CA-C	5.36	117.11	108.32
3	D	54	SER	CB-CA-C	-5.36	110.41	116.63
3	J	210	VAL	N-CA-C	-5.36	105.49	110.53
1	j	74	GLN	CA-C-N	5.36	126.54	119.84
1	j	74	GLN	C-N-CA	5.36	126.54	119.84
6	Y	163	ALA	N-CA-C	5.36	116.80	111.07
2	I	29	THR	N-CA-C	5.36	117.10	108.32
2	A	29	THR	N-CA-C	5.35	117.10	108.32
3	L	210	VAL	N-CA-C	-5.35	105.50	110.53
6	X	956	HIS	N-CA-C	5.35	119.65	113.18
3	B	210	VAL	N-CA-C	-5.34	105.50	110.53
1	l	72	SER	CA-C-N	5.34	128.29	120.71
1	l	72	SER	C-N-CA	5.34	128.29	120.71
2	Q	29	THR	N-CA-C	5.34	117.08	108.32
2	K	29	THR	N-CA-C	5.33	117.07	108.32
2	E	29	THR	N-CA-C	5.33	117.07	108.32
3	P	210	VAL	N-CA-C	-5.33	105.52	110.53
3	D	164	ASP	N-CA-C	5.33	116.95	111.03

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Mol	Chain	Res	Type	Atoms	Z	Observed(^{o})	Ideal(^{o})
6	Y	17	ALA	N-CA-C	-5.33	105.42	112.24
6	Y	901	ILE	N-CA-C	-5.33	104.58	112.04
3	R	210	VAL	N-CA-C	-5.32	105.53	110.53
3	T	210	VAL	N-CA-C	-5.32	105.53	110.53
5	W	801	SER	N-CA-C	-5.31	103.02	110.35
6	X	968	PHE	N-CA-C	-5.31	105.18	110.97
3	H	210	VAL	N-CA-C	-5.30	105.54	110.53
5	W	695	ILE	CA-C-N	5.30	125.21	119.85
5	W	695	ILE	C-N-CA	5.30	125.21	119.85
3	F	210	VAL	N-CA-C	-5.30	105.55	110.53
3	J	164	ASP	N-CA-C	5.30	116.91	111.03
6	Y	807	PHE	N-CA-C	5.30	117.80	111.71
4	U	84	ILE	N-CA-C	5.29	116.77	111.00
3	N	210	VAL	N-CA-C	-5.29	105.56	110.53
6	X	772	THR	N-CA-C	5.29	117.83	107.57
6	X	610	THR	CA-C-N	5.29	126.45	119.84
6	X	610	THR	C-N-CA	5.29	126.45	119.84
1	b	76	ASN	N-CA-C	5.28	122.05	110.80
3	L	164	ASP	N-CA-C	5.28	116.89	111.03
6	Y	1047	TYR	CA-C-N	-5.27	113.87	119.98
6	Y	1047	TYR	C-N-CA	-5.27	113.87	119.98
3	B	164	ASP	N-CA-C	5.26	116.87	111.03
3	F	164	ASP	N-CA-C	5.26	116.87	111.03
6	X	1184	LEU	CA-C-N	-5.26	113.26	119.84
6	X	1184	LEU	C-N-CA	-5.26	113.26	119.84
1	l	18	ALA	N-CA-C	-5.25	102.34	109.54
3	R	164	ASP	N-CA-C	5.25	116.86	111.03
6	X	1206	GLY	N-CA-C	-5.25	105.99	113.86
2	C	26	MET	N-CA-C	5.25	119.23	112.41
5	W	638	HIS	N-CA-C	5.24	121.96	110.80
2	E	26	MET	N-CA-C	5.24	119.22	112.41
3	P	164	ASP	N-CA-C	5.24	116.84	111.03
2	Q	26	MET	N-CA-C	5.24	119.22	112.41
4	U	170	THR	N-CA-C	-5.24	106.80	114.39
2	I	26	MET	N-CA-C	5.23	119.21	112.41
6	Y	1138	THR	N-CA-C	5.23	117.28	110.53
2	K	26	MET	N-CA-C	5.23	119.21	112.41
2	S	26	MET	N-CA-C	5.23	119.21	112.41
6	X	602	GLY	N-CA-C	5.23	125.57	113.18
3	H	274	ILE	CB-CA-C	-5.22	103.80	112.26
2	O	26	MET	N-CA-C	5.22	119.20	112.41
2	G	26	MET	N-CA-C	5.22	119.19	112.41

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Mol	Chain	Res	Type	Atoms	Z	Observed(^{o})	Ideal(^{o})
5	W	502	ASN	CA-C-N	5.22	124.91	119.64
5	W	502	ASN	C-N-CA	5.22	124.91	119.64
5	W	1208	CYS	N-CA-C	5.22	117.95	109.24
6	Y	40	ASN	N-CA-C	5.21	116.65	111.07
6	Y	15	SER	CA-C-N	5.21	126.36	119.84
6	Y	15	SER	C-N-CA	5.21	126.36	119.84
2	A	26	MET	N-CA-C	5.21	119.19	112.41
2	M	26	MET	N-CA-C	5.21	119.18	112.41
1	j	18	ALA	N-CA-C	-5.21	105.50	111.07
3	F	274	ILE	CB-CA-C	-5.21	103.82	112.26
3	N	274	ILE	CB-CA-C	-5.21	103.82	112.26
3	T	164	ASP	N-CA-C	5.21	116.81	111.03
6	X	578	ASP	N-CA-C	5.21	117.10	109.71
3	L	274	ILE	CB-CA-C	-5.20	103.83	112.26
6	X	750	MET	N-CA-C	5.20	117.85	111.82
3	P	274	ILE	CB-CA-C	-5.20	103.84	112.26
3	H	588	ILE	CA-C-N	5.19	125.48	119.93
3	H	588	ILE	C-N-CA	5.19	125.48	119.93
3	R	274	ILE	CB-CA-C	-5.19	103.85	112.26
3	B	274	ILE	CB-CA-C	-5.19	103.86	112.26
3	P	421	HIS	N-CA-C	5.19	117.17	107.99
3	T	274	ILE	CB-CA-C	-5.19	103.86	112.26
3	N	164	ASP	N-CA-C	5.18	116.78	111.03
3	D	588	ILE	CA-C-N	5.18	125.47	119.93
3	D	588	ILE	C-N-CA	5.18	125.47	119.93
6	X	519	GLN	N-CA-C	5.18	121.25	109.81
5	W	1120	VAL	N-CA-C	5.17	113.53	107.89
3	R	421	HIS	N-CA-C	5.17	117.14	107.99
3	F	421	HIS	N-CA-C	5.17	117.14	107.99
3	J	274	ILE	CB-CA-C	-5.17	103.89	112.26
3	L	421	HIS	N-CA-C	5.17	117.14	107.99
6	Y	990	ASN	N-CA-C	-5.17	106.21	112.88
3	B	421	HIS	N-CA-C	5.17	117.13	107.99
3	T	421	HIS	N-CA-C	5.16	117.06	108.02
1	p	18	ALA	N-CA-C	-5.16	105.55	111.07
3	D	274	ILE	CB-CA-C	-5.16	103.90	112.26
3	D	421	HIS	N-CA-C	5.16	117.05	108.02
3	L	290	ALA	N-CA-C	5.16	118.36	108.97
1	d	19	ALA	N-CA-C	-5.16	100.11	108.52
3	B	290	ALA	N-CA-C	5.16	118.36	108.97
3	H	421	HIS	N-CA-C	5.16	117.05	108.02
3	R	290	ALA	N-CA-C	5.16	118.36	108.97

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Mol	Chain	Res	Type	Atoms	Z	Observed(^{o})	Ideal(^{o})
3	H	164	ASP	N-CA-C	5.16	116.75	111.03
3	T	290	ALA	N-CA-C	5.16	118.35	108.97
3	P	588	ILE	CA-C-N	5.15	125.44	119.93
3	P	588	ILE	C-N-CA	5.15	125.44	119.93
3	N	588	ILE	CA-C-N	5.15	125.44	119.93
3	N	588	ILE	C-N-CA	5.15	125.44	119.93
3	R	588	ILE	CA-C-N	5.15	125.44	119.93
3	R	588	ILE	C-N-CA	5.15	125.44	119.93
3	F	290	ALA	N-CA-C	5.15	118.33	108.97
3	J	290	ALA	N-CA-C	5.15	118.34	108.97
3	D	290	ALA	N-CA-C	5.14	118.33	108.97
3	H	290	ALA	N-CA-C	5.14	118.33	108.97
3	J	421	HIS	N-CA-C	5.14	117.09	107.99
3	J	588	ILE	CA-C-N	5.14	125.43	119.93
3	J	588	ILE	C-N-CA	5.14	125.43	119.93
3	N	290	ALA	N-CA-C	5.14	118.33	108.97
1	j	72	SER	CA-C-N	5.14	127.59	120.29
1	j	72	SER	C-N-CA	5.14	127.59	120.29
3	P	290	ALA	N-CA-C	5.14	118.32	108.97
1	b	75	PRO	N-CA-CB	-5.14	97.86	103.25
3	N	421	HIS	N-CA-C	5.14	117.08	107.99
3	B	588	ILE	CA-C-N	5.13	125.42	119.93
3	B	588	ILE	C-N-CA	5.13	125.42	119.93
3	T	343	ALA	N-CA-C	-5.13	104.36	111.28
3	F	588	ILE	CA-C-N	5.12	125.41	119.93
3	F	588	ILE	C-N-CA	5.12	125.41	119.93
3	J	60	SER	N-CA-C	-5.12	105.78	111.36
3	P	343	ALA	N-CA-C	-5.12	104.37	111.28
3	N	343	ALA	N-CA-C	-5.12	104.37	111.28
5	W	109	ALA	N-CA-C	-5.12	107.16	113.41
1	j	75	PRO	N-CA-CB	-5.12	97.88	103.25
3	L	588	ILE	CA-C-N	5.12	125.40	119.93
3	L	588	ILE	C-N-CA	5.12	125.40	119.93
3	H	224	ASP	N-CA-C	5.11	117.09	110.65
3	B	343	ALA	N-CA-C	-5.11	104.39	111.28
3	J	224	ASP	N-CA-C	5.11	117.08	110.65
3	H	343	ALA	N-CA-C	-5.10	104.39	111.28
5	W	992	SER	CA-C-N	5.10	126.22	119.84
5	W	992	SER	C-N-CA	5.10	126.22	119.84
3	N	224	ASP	N-CA-C	5.10	117.07	110.65
3	R	343	ALA	N-CA-C	-5.10	104.40	111.28
6	Y	565	LYS	N-CA-C	-5.09	106.76	112.87

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Mol	Chain	Res	Type	Atoms	Z	Observed(^{o})	Ideal(^{o})
3	H	524	ALA	N-CA-C	5.09	116.52	110.97
3	F	343	ALA	N-CA-C	-5.09	104.41	111.28
3	P	224	ASP	N-CA-C	5.09	117.07	110.65
6	Y	485	PRO	N-CA-C	-5.09	107.11	114.18
3	T	588	ILE	CA-C-N	5.09	125.37	119.93
3	T	588	ILE	C-N-CA	5.09	125.37	119.93
3	D	224	ASP	N-CA-C	5.08	117.05	110.65
3	L	343	ALA	N-CA-C	-5.08	104.42	111.28
6	X	946	VAL	N-CA-C	5.08	119.91	109.34
3	T	224	ASP	N-CA-C	5.08	117.05	110.65
3	B	224	ASP	N-CA-C	5.08	117.05	110.65
3	J	343	ALA	N-CA-C	-5.08	104.42	111.28
6	Y	643	ALA	N-CA-C	-5.07	105.83	111.36
3	D	343	ALA	N-CA-C	-5.07	104.43	111.28
3	R	224	ASP	N-CA-C	5.07	117.04	110.65
3	F	520	VAL	N-CA-C	-5.07	105.91	110.82
3	F	224	ASP	N-CA-C	5.06	117.03	110.65
3	L	224	ASP	N-CA-C	5.06	117.03	110.65
3	L	520	VAL	N-CA-C	-5.06	105.91	110.82
5	W	542	ILE	CA-C-N	5.06	125.30	119.83
5	W	542	ILE	C-N-CA	5.06	125.30	119.83
5	W	1013	ILE	N-CA-C	-5.05	104.87	110.62
4	U	368	ARG	N-CA-C	-5.04	107.08	113.18
5	W	709	GLY	N-CA-C	5.04	117.26	111.36
3	N	524	ALA	N-CA-C	5.03	116.46	110.97
3	H	520	VAL	N-CA-C	-5.03	105.94	110.82
3	J	520	VAL	N-CA-C	-5.03	105.94	110.82
6	Y	531	CYS	N-CA-C	-5.03	108.17	114.56
5	W	918	ILE	CB-CA-C	-5.02	108.93	113.70
5	W	1186	ASN	N-CA-C	-5.02	102.08	110.17
3	B	520	VAL	N-CA-C	-5.02	105.95	110.82
3	R	520	VAL	N-CA-C	-5.02	105.95	110.82
3	D	520	VAL	N-CA-C	-5.02	105.95	110.82
3	P	520	VAL	N-CA-C	-5.01	105.96	110.82

There are no chirality outliers.

All (40) planarity outliers are listed below:

Mol	Chain	Res	Type	Group
3	B	337	ARG	Sidechain
3	D	337	ARG	Sidechain
3	F	337	ARG	Sidechain

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Mol	Chain	Res	Type	Group
3	H	337	ARG	Sidechain
3	J	337	ARG	Sidechain
3	L	337	ARG	Sidechain
3	N	337	ARG	Sidechain
3	P	337	ARG	Sidechain
3	R	337	ARG	Sidechain
3	T	337	ARG	Sidechain
6	X	344	TYR	Sidechain
6	X	525	TYR	Sidechain
6	X	930	ARG	Sidechain
6	Y	347	ARG	Sidechain
6	Y	737	ARG	Sidechain
1	b	43	CYS	Peptide
1	b	73	ARG	Mainchain,Peptide
1	b	75	PRO	Peptide
1	d	21	ARG	Sidechain
1	d	43	CYS	Peptide
1	d	75	PRO	Peptide
1	f	43	CYS	Peptide
1	f	75	PRO	Peptide
1	h	43	CYS	Peptide
1	h	75	PRO	Peptide
1	j	43	CYS	Peptide
1	j	75	PRO	Peptide
1	l	43	CYS	Peptide
1	l	75	PRO	Peptide
1	n	21	ARG	Sidechain
1	n	43	CYS	Peptide
1	n	75	PRO	Peptide
1	p	21	ARG	Sidechain
1	p	43	CYS	Peptide
1	p	75	PRO	Peptide
1	r	43	CYS	Peptide
1	r	75	PRO	Peptide
1	t	43	CYS	Peptide
1	t	75	PRO	Peptide

5.2 Too-close contacts (i)

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

the asymmetric unit, whereas Symm-Clashes lists symmetry-related clashes.

Mol	Chain	Non-H	H(model)	H(added)	Clashes	Symm-Clashes
1	b	666	0	660	204	0
1	d	666	0	660	172	0
1	f	666	0	657	151	0
1	h	666	0	655	167	0
1	j	666	0	659	188	0
1	l	666	0	656	194	0
1	n	666	0	658	170	0
1	p	666	0	659	224	0
1	r	666	0	657	171	0
1	t	666	0	659	230	0
2	A	291	0	277	57	0
2	C	291	0	277	81	0
2	E	291	0	277	80	0
2	G	284	0	267	58	0
2	I	291	0	277	84	0
2	K	291	0	277	81	0
2	M	284	0	267	60	0
2	O	291	0	277	78	0
2	Q	291	0	277	88	0
2	S	284	0	267	60	0
3	B	4508	0	4555	785	0
3	D	4508	0	4554	1045	0
3	F	4508	0	4554	1002	0
3	H	4508	0	4554	1005	0
3	J	4508	0	4553	1049	0
3	L	4508	0	4555	1031	0
3	N	4508	0	4555	1023	0
3	P	4508	0	4555	1083	0
3	R	4508	0	4554	1006	0
3	T	4508	0	4555	1053	0
4	U	3138	0	3061	471	0
4	V	3138	0	3061	181	0
5	W	9882	0	9821	1088	0
6	X	7873	0	7851	1319	0
6	Y	8835	0	8748	1111	0
7	A	15	0	27	27	0
7	C	15	0	27	31	0
7	E	15	0	27	27	0
7	G	15	0	27	28	0
7	I	15	0	27	29	0
7	K	15	0	27	27	0
7	M	15	0	27	26	0

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Mol	Chain	Non-H	H(model)	H(added)	Clashes	Symm-Clashes
7	O	15	0	27	28	0
7	Q	15	0	27	27	0
7	S	15	0	27	30	0
All	All	87645	0	87676	14002	0

The all-atom clashscore is defined as the number of clashes found per 1000 atoms (including hydrogen atoms). The all-atom clashscore for this structure is 80.

All (14002) 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)
1:p:8:GLN:HG2	3:R:302:GLU:CG	1.26	1.59
1:h:23:THR:CB	3:H:323:SER:HA	1.11	1.58
1:b:45:ARG:CG	3:B:400:ALA:HA	1.27	1.57
1:t:59:HIS:HB3	3:T:398:SER:CB	1.32	1.56
1:p:59:HIS:CB	3:P:398:SER:HA	1.35	1.55
1:p:45:ARG:CG	3:P:400:ALA:HA	1.19	1.54
1:j:45:ARG:CG	3:J:400:ALA:HA	1.37	1.53
1:d:4:HIS:CD2	1:d:54:THR:HG22	1.44	1.53
1:t:45:ARG:CG	3:T:400:ALA:HA	1.35	1.53
1:p:59:HIS:CB	3:P:398:SER:CA	1.84	1.52
1:j:8:GLN:HG2	3:L:302:GLU:CG	1.37	1.51
1:n:45:ARG:CG	3:N:400:ALA:HA	1.40	1.50
1:h:59:HIS:HB2	3:H:398:SER:CB	1.35	1.50
1:j:59:HIS:CB	3:J:398:SER:HB2	1.35	1.49
1:h:23:THR:HB	3:H:323:SER:CA	1.41	1.48
1:j:59:HIS:HB2	3:J:398:SER:CB	1.44	1.48
1:p:24:LEU:HD21	1:p:80:ARG:CD	1.41	1.48
1:j:24:LEU:CD1	1:j:41:VAL:HG22	1.45	1.47
1:p:59:HIS:HB3	3:P:398:SER:CB	1.43	1.47
1:j:23:THR:HG21	3:J:323:SER:C	1.33	1.46
1:f:44:GLY:CA	3:F:322:TYR:OH	1.64	1.45
1:d:45:ARG:CG	3:D:400:ALA:HA	1.43	1.45
1:j:71:CYS:SG	1:j:73:ARG:HG2	1.55	1.45
1:p:59:HIS:HB3	3:P:398:SER:CA	0.98	1.44
1:b:74:GLN:C	1:b:75:PRO:N	1.68	1.43
1:j:59:HIS:CB	3:J:398:SER:CB	1.96	1.42
1:r:44:GLY:CA	3:R:322:TYR:OH	1.67	1.42
1:d:22:LEU:CD1	1:d:83:VAL:HG21	1.51	1.41
1:r:6:ILE:CD1	3:T:586:VAL:HG21	1.51	1.41
1:h:6:ILE:CD1	3:D:586:VAL:HG21	1.51	1.40

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:l:45:ARG:CG	3:L:400:ALA:HA	1.48	1.40
1:t:59:HIS:CB	3:T:398:SER:CB	1.96	1.40
1:l:59:HIS:HB2	3:L:398:SER:CB	1.49	1.39
1:h:45:ARG:CG	3:H:400:ALA:HA	1.49	1.39
1:d:8:GLN:HG2	3:F:302:GLU:CG	1.48	1.39
1:t:61:ASN:ND2	3:T:395:THR:HG21	1.37	1.38
4:U:32:CYS:HB3	6:X:958:HIS:CE1	1.55	1.38
1:b:44:GLY:HA3	3:B:322:TYR:OH	1.24	1.38
1:f:45:ARG:CG	3:F:400:ALA:HA	1.54	1.38
6:X:464:SER:HB2	6:Y:500:ILE:CG2	1.52	1.38
1:l:24:LEU:CD1	1:l:32:THR:O	1.72	1.38
1:f:24:LEU:HD21	1:f:80:ARG:CD	1.51	1.38
1:l:21:ARG:O	3:L:323:SER:HB2	1.20	1.37
1:r:15:ARG:CZ	3:R:406:ASN:OD1	1.72	1.37
1:h:23:THR:CB	3:H:323:SER:CA	1.98	1.37
1:j:23:THR:CG2	3:J:323:SER:C	1.98	1.37
1:l:59:HIS:CB	3:L:398:SER:HB2	1.53	1.36
1:t:59:HIS:HB2	3:T:398:SER:CA	1.55	1.36
1:f:6:ILE:CD1	3:H:586:VAL:HG21	1.53	1.36
1:t:21:ARG:O	3:T:323:SER:CB	1.73	1.36
1:d:57:ALA:HB1	1:d:59:HIS:CD2	1.58	1.36
1:j:8:GLN:CG	3:L:302:GLU:HG3	1.54	1.35
1:p:8:GLN:CG	3:R:302:GLU:HG3	1.52	1.35
1:l:6:ILE:CD1	3:N:586:VAL:HG21	1.55	1.35
1:l:8:GLN:NE2	3:N:302:GLU:O	1.58	1.35
1:h:59:HIS:CB	3:H:398:SER:HB2	1.53	1.35
1:t:6:ILE:CD1	3:P:586:VAL:HG21	1.55	1.35
3:R:628:GLN:NE2	5:W:870:ALA:HA	1.41	1.35
1:l:15:ARG:NE	3:L:406:ASN:OD1	1.59	1.34
1:f:8:GLN:HG2	3:H:302:GLU:CG	1.57	1.34
1:n:23:THR:HG22	1:n:42:THR:OG1	1.17	1.34
1:t:8:GLN:NE2	3:P:302:GLU:O	1.60	1.34
1:b:50:ALA:O	1:b:74:GLN:HB2	1.19	1.33
1:j:8:GLN:NE2	3:L:302:GLU:HG2	1.41	1.33
1:t:8:GLN:HG2	3:P:302:GLU:CG	1.55	1.33
1:r:15:ARG:NH1	3:R:406:ASN:OD1	1.58	1.33
1:f:15:ARG:NE	3:F:406:ASN:OD1	1.59	1.33
1:p:45:ARG:CG	3:P:400:ALA:CA	2.06	1.33
3:D:469:VAL:HG21	3:F:575:MET:CE	1.59	1.33
1:p:23:THR:HB	3:P:323:SER:CA	1.59	1.33
3:F:469:VAL:HG21	3:H:575:MET:CE	1.59	1.33

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:h:23:THR:OG1	3:H:323:SER:HA	1.27	1.32
3:D:575:MET:CE	3:H:469:VAL:HG21	1.59	1.32
1:p:59:HIS:CD2	3:P:398:SER:HB2	1.65	1.32
1:n:8:GLN:NE2	3:J:302:GLU:O	1.61	1.32
3:P:469:VAL:HG21	3:R:575:MET:CE	1.59	1.32
1:b:4:HIS:NE2	1:b:54:THR:HG22	1.42	1.31
3:J:575:MET:CE	3:N:469:VAL:HG21	1.59	1.31
1:f:15:ARG:CZ	3:F:406:ASN:OD1	1.76	1.31
1:n:8:GLN:HG2	3:J:302:GLU:CG	1.57	1.31
1:p:23:THR:CB	3:P:323:SER:HA	1.59	1.31
1:t:21:ARG:O	3:T:323:SER:HB2	1.18	1.31
3:J:469:VAL:HG21	3:L:575:MET:CE	1.59	1.31
3:R:469:VAL:HG21	3:T:575:MET:CE	1.59	1.31
1:l:8:GLN:HG2	3:N:302:GLU:CG	1.61	1.31
1:b:45:ARG:CG	3:B:400:ALA:CA	2.09	1.31
1:f:44:GLY:N	3:F:322:TYR:OH	1.64	1.31
1:h:15:ARG:NE	3:H:406:ASN:OD1	1.61	1.31
1:r:45:ARG:CG	3:R:400:ALA:HA	1.60	1.31
1:h:8:GLN:HG2	3:D:302:GLU:CG	1.58	1.30
1:f:59:HIS:CB	3:F:398:SER:HB2	1.58	1.30
3:L:469:VAL:HG21	3:N:575:MET:CE	1.59	1.30
3:T:291:PRO:CB	3:T:467:THR:HG22	1.62	1.30
1:b:59:HIS:CB	3:B:397:VAL:O	1.79	1.29
3:L:291:PRO:CB	3:L:467:THR:HG22	1.62	1.29
3:P:575:MET:CE	3:T:469:VAL:HG21	1.59	1.29
1:n:6:ILE:CD1	3:J:586:VAL:HG21	1.60	1.29
3:P:291:PRO:CB	3:P:467:THR:HG22	1.62	1.29
3:R:291:PRO:CB	3:R:467:THR:HG22	1.62	1.29
3:B:291:PRO:CB	3:B:467:THR:HG22	1.62	1.29
3:F:291:PRO:CB	3:F:467:THR:HG22	1.62	1.29
1:r:23:THR:OG1	3:R:323:SER:CB	1.78	1.28
1:j:59:HIS:ND1	3:J:397:VAL:O	1.65	1.28
1:p:59:HIS:CA	3:P:398:SER:HA	1.62	1.28
3:H:291:PRO:CB	3:H:467:THR:HG22	1.62	1.28
1:p:6:ILE:CD1	3:R:586:VAL:HG21	1.62	1.28
1:t:23:THR:OG1	3:T:323:SER:HA	1.34	1.28
3:P:575:MET:HE3	3:T:469:VAL:CG2	1.64	1.28
1:t:44:GLY:HA3	3:T:322:TYR:OH	1.13	1.28
3:N:291:PRO:CB	3:N:467:THR:HG22	1.62	1.28
6:X:606:MET:CE	6:Y:718:THR:HG23	1.62	1.28
1:h:59:HIS:CG	3:H:397:VAL:O	1.88	1.27

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:r:8:GLN:HG2	3:T:302:GLU:CG	1.64	1.27
1:r:23:THR:HB	1:r:42:THR:OG1	1.19	1.27
3:D:469:VAL:CG2	3:F:575:MET:HE3	1.64	1.27
1:r:15:ARG:NE	3:R:406:ASN:OD1	1.66	1.27
1:r:44:GLY:HA3	3:R:322:TYR:OH	1.11	1.27
3:F:435:PRO:HG2	3:N:436:TYR:OH	1.29	1.27
3:P:469:VAL:CG2	3:R:575:MET:HE3	1.64	1.27
1:d:22:LEU:HD11	1:d:83:VAL:CG2	1.65	1.27
3:D:291:PRO:CB	3:D:467:THR:HG22	1.62	1.27
1:b:45:ARG:HA	3:B:399:ALA:O	1.35	1.26
1:h:15:ARG:CZ	3:H:406:ASN:OD1	1.83	1.26
1:f:59:HIS:HB2	3:F:398:SER:CB	1.63	1.26
1:p:15:ARG:HH11	3:P:406:ASN:CB	1.45	1.26
1:t:24:LEU:CD2	1:t:35:PHE:HA	1.65	1.26
3:B:274:ILE:HD11	3:B:514:LEU:CD2	1.66	1.26
3:F:274:ILE:HD11	3:F:514:LEU:CD2	1.66	1.26
3:J:575:MET:HE3	3:N:469:VAL:CG2	1.64	1.26
1:j:23:THR:OG1	3:J:323:SER:CA	1.83	1.26
1:t:59:HIS:HB3	3:T:398:SER:OG	1.15	1.26
3:J:291:PRO:CB	3:J:467:THR:HG22	1.62	1.26
3:L:469:VAL:CG2	3:N:575:MET:HE3	1.64	1.26
3:R:469:VAL:CG2	3:T:575:MET:HE3	1.64	1.26
3:T:274:ILE:HD11	3:T:514:LEU:CD2	1.66	1.26
1:p:15:ARG:NH1	3:P:406:ASN:HB3	1.47	1.26
3:D:274:ILE:HD11	3:D:514:LEU:CD2	1.66	1.26
1:b:59:HIS:HB2	3:B:397:VAL:O	1.22	1.26
1:p:45:ARG:HA	3:P:399:ALA:O	1.29	1.26
1:f:21:ARG:HD3	3:F:327:TYR:OH	1.36	1.25
1:d:8:GLN:CG	3:F:302:GLU:HG3	1.65	1.25
1:j:15:ARG:HH11	3:J:406:ASN:CB	1.48	1.25
1:j:22:LEU:HD23	1:j:42:THR:O	1.31	1.25
3:F:469:VAL:CG2	3:H:575:MET:HE3	1.64	1.25
3:J:274:ILE:HD11	3:J:514:LEU:CD2	1.66	1.25
3:J:469:VAL:CG2	3:L:575:MET:HE3	1.64	1.25
1:p:62:VAL:O	1:p:64:THR:N	1.69	1.25
3:H:274:ILE:HD11	3:H:514:LEU:CD2	1.66	1.25
1:r:23:THR:OG1	3:R:323:SER:HB3	1.28	1.25
3:D:575:MET:HE3	3:H:469:VAL:CG2	1.64	1.25
3:J:78:MET:CE	3:N:238:VAL:HG21	1.67	1.25
3:R:274:ILE:HD11	3:R:514:LEU:CD2	1.66	1.25
1:p:8:GLN:CG	3:R:302:GLU:CG	2.08	1.24

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:N:274:ILE:HD11	3:N:514:LEU:CD2	1.66	1.24
3:P:78:MET:CE	3:T:238:VAL:HG21	1.67	1.24
3:L:274:ILE:HD11	3:L:514:LEU:CD2	1.66	1.24
3:P:238:VAL:HG21	3:R:78:MET:CE	1.67	1.24
3:D:78:MET:CE	3:H:238:VAL:HG21	1.67	1.24
3:J:238:VAL:HG21	3:L:78:MET:CE	1.67	1.24
1:r:16:ALA:O	1:r:22:LEU:CD2	1.84	1.24
7:A:101:MYR:C14	7:A:101:MYR:H101	1.67	1.24
1:n:59:HIS:HB2	3:N:397:VAL:O	1.33	1.24
3:P:274:ILE:HD11	3:P:514:LEU:CD2	1.66	1.24
7:E:101:MYR:C14	7:E:101:MYR:H101	1.67	1.23
1:f:58:PRO:CB	3:F:423:GLN:HE21	1.52	1.23
1:f:59:HIS:ND1	3:F:397:VAL:O	1.72	1.23
3:R:238:VAL:HG21	3:T:78:MET:CE	1.67	1.23
7:S:101:MYR:C14	7:S:101:MYR:H101	1.67	1.23
3:D:238:VAL:HG21	3:F:78:MET:CE	1.67	1.23
7:M:101:MYR:C14	7:M:101:MYR:H101	1.67	1.23
7:Q:101:MYR:C14	7:Q:101:MYR:H101	1.67	1.23
3:F:238:VAL:HG21	3:H:78:MET:CE	1.67	1.22
7:I:101:MYR:C14	7:I:101:MYR:H101	1.67	1.22
3:L:238:VAL:HG21	3:N:78:MET:CE	1.67	1.22
3:N:161:LYS:HD3	3:N:162:HIS:N	1.55	1.22
1:b:23:THR:HG22	1:b:42:THR:OG1	1.38	1.22
1:l:24:LEU:HD12	1:l:32:THR:O	1.08	1.22
1:b:59:HIS:O	3:B:397:VAL:O	1.57	1.22
1:j:45:ARG:HG3	3:J:401:GLY:N	1.54	1.22
7:C:101:MYR:C14	7:C:101:MYR:H101	1.67	1.22
7:G:101:MYR:C14	7:G:101:MYR:H101	1.67	1.22
1:h:58:PRO:CB	3:H:423:GLN:HE21	1.54	1.21
1:n:23:THR:CB	3:N:323:SER:HA	1.71	1.21
1:f:15:ARG:NH1	3:F:406:ASN:OD1	1.72	1.21
1:f:44:GLY:HA3	3:F:322:TYR:OH	1.14	1.21
7:O:101:MYR:C14	7:O:101:MYR:H101	1.67	1.21
6:X:787:ARG:CG	6:X:787:ARG:HH21	1.54	1.21
1:l:15:ARG:CZ	3:L:406:ASN:OD1	1.89	1.20
1:n:8:GLN:CG	3:J:302:GLU:HG3	1.71	1.20
4:U:60:SER:HB3	4:U:63:SER:HB2	1.24	1.20
1:d:15:ARG:HE	3:D:406:ASN:ND2	1.40	1.20
1:p:61:ASN:CG	3:P:395:THR:CG2	2.15	1.20
1:t:59:HIS:CB	3:T:398:SER:CA	2.17	1.20
7:K:101:MYR:C14	7:K:101:MYR:H101	1.67	1.20

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:b:59:HIS:O	3:B:397:VAL:C	1.85	1.19
1:r:44:GLY:N	3:R:322:TYR:OH	1.74	1.19
1:p:8:GLN:CD	3:R:302:GLU:HG2	1.68	1.19
1:b:45:ARG:HG3	3:B:401:GLY:N	1.56	1.19
3:J:338:VAL:HG12	3:J:440:LEU:HG	1.20	1.19
5:W:840:CYS:HB3	5:W:854:MET:SD	1.81	1.19
6:X:616:SER:CB	6:Y:723:ASP:HB2	1.73	1.19
1:b:23:THR:CB	3:B:323:SER:HA	1.70	1.19
1:h:23:THR:HB	3:H:323:SER:CB	1.71	1.19
1:j:58:PRO:CG	3:J:423:GLN:HE21	1.55	1.19
1:n:23:THR:OG1	3:N:323:SER:HA	1.43	1.19
1:p:44:GLY:HA3	3:P:322:TYR:OH	1.38	1.19
1:t:59:HIS:CB	3:T:398:SER:HA	1.72	1.19
1:j:23:THR:OG1	3:J:323:SER:HA	1.00	1.18
1:p:45:ARG:HG3	3:P:401:GLY:N	1.58	1.18
1:t:8:GLN:CG	3:P:302:GLU:HG3	1.73	1.18
1:d:4:HIS:NE2	1:d:54:THR:HG22	1.57	1.18
1:j:71:CYS:SG	1:j:73:ARG:CG	2.31	1.18
5:W:198:ARG:NH2	6:X:648:ALA:HB1	1.58	1.18
4:U:178:THR:CG2	6:X:376:LEU:HD23	1.72	1.18
1:h:8:GLN:CG	3:D:302:GLU:HG3	1.74	1.18
1:b:15:ARG:NH1	3:B:402:ALA:HB1	1.59	1.17
1:t:8:GLN:NE2	3:P:302:GLU:C	2.02	1.17
3:P:472:ARG:NH1	3:R:530:ARG:HG3	1.58	1.17
4:U:32:CYS:CB	6:X:958:HIS:CE1	2.26	1.17
1:n:8:GLN:NE2	3:J:302:GLU:C	2.02	1.17
1:p:56:LEU:HD13	1:p:65:ILE:HG21	1.26	1.17
1:p:61:ASN:CG	3:P:395:THR:HG21	1.70	1.17
1:t:22:LEU:HD11	1:t:83:VAL:CG2	1.75	1.17
1:t:61:ASN:ND2	3:T:395:THR:CG2	2.06	1.17
3:D:238:VAL:HA	3:F:142:ILE:HD11	1.18	1.17
1:p:6:ILE:HD11	3:R:586:VAL:CG2	1.75	1.17
1:p:58:PRO:HG3	3:P:423:GLN:NE2	1.59	1.17
1:r:45:ARG:HG3	3:R:401:GLY:N	1.58	1.17
4:V:192:PHE:HB2	6:Y:376:LEU:HD21	1.26	1.17
1:h:59:HIS:CB	3:H:398:SER:CB	2.14	1.16
1:j:8:GLN:CD	3:L:302:GLU:HG2	1.70	1.16
1:r:58:PRO:CG	3:R:423:GLN:HE21	1.58	1.16
1:f:8:GLN:CG	3:H:302:GLU:HG3	1.74	1.16
1:d:45:ARG:HG3	3:D:401:GLY:N	1.59	1.16
1:h:59:HIS:ND1	1:h:60:ALA:N	1.91	1.16

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:t:21:ARG:NH1	3:T:327:TYR:OH	1.78	1.16
3:J:274:ILE:HD11	3:J:514:LEU:HD22	1.27	1.16
3:R:609:ARG:HB3	2:S:29:THR:HG22	1.27	1.16
1:j:45:ARG:CG	3:J:400:ALA:CA	2.23	1.16
1:p:59:HIS:CG	3:P:398:SER:HB2	1.80	1.16
1:t:59:HIS:CD2	3:T:397:VAL:O	1.98	1.16
1:r:45:ARG:HG3	3:R:401:GLY:H	1.02	1.16
1:t:61:ASN:CG	3:T:395:THR:HG21	1.70	1.16
1:d:23:THR:CG2	3:D:323:SER:C	2.18	1.15
1:l:8:GLN:CG	3:N:302:GLU:HG3	1.77	1.15
1:j:59:HIS:HB3	3:J:398:SER:HB2	1.28	1.15
3:R:338:VAL:HG12	3:R:440:LEU:HG	1.20	1.15
1:p:8:GLN:NE2	3:R:302:GLU:HG2	1.58	1.15
1:t:45:ARG:CG	3:T:400:ALA:CA	2.22	1.15
3:H:338:VAL:HG12	3:H:440:LEU:HG	1.20	1.15
1:f:59:HIS:CB	3:F:398:SER:CB	2.20	1.15
3:F:358:LEU:HD13	3:F:417:PHE:HE1	1.12	1.15
3:R:274:ILE:HD11	3:R:514:LEU:HD22	1.27	1.15
4:U:178:THR:CG2	6:X:376:LEU:CD2	2.25	1.15
3:B:291:PRO:HB2	3:B:467:THR:HG22	1.29	1.15
3:H:358:LEU:HD13	3:H:417:PHE:HE1	1.12	1.15
3:J:609:ARG:HB3	2:K:29:THR:HG22	1.27	1.15
3:P:338:VAL:HG12	3:P:440:LEU:HG	1.20	1.15
1:d:4:HIS:NE2	1:d:54:THR:CG2	2.10	1.14
1:d:15:ARG:HH11	3:D:406:ASN:CB	1.59	1.14
1:t:45:ARG:HA	3:T:399:ALA:O	1.47	1.14
3:L:358:LEU:HD13	3:L:417:PHE:HE1	1.12	1.14
4:U:178:THR:HG22	6:X:376:LEU:CD2	1.75	1.14
3:N:291:PRO:HB2	3:N:467:THR:HG22	1.29	1.14
3:B:358:LEU:HD13	3:B:417:PHE:HE1	1.12	1.14
4:U:27:GLN:HG2	6:X:880:LEU:HB3	1.15	1.14
3:P:609:ARG:HB3	2:Q:29:THR:HG22	1.27	1.14
1:f:58:PRO:HB3	3:F:423:GLN:HE21	1.03	1.14
3:T:274:ILE:HD11	3:T:514:LEU:HD22	1.28	1.14
1:f:24:LEU:CD2	1:f:80:ARG:HD2	1.77	1.13
3:F:274:ILE:HD11	3:F:514:LEU:HD22	1.28	1.13
2:O:29:THR:HG22	3:T:609:ARG:HB3	1.27	1.13
7:O:101:MYR:O1	3:P:204:PRO:HG3	1.49	1.13
3:R:341:MET:HE2	3:R:341:MET:HA	1.31	1.13
3:T:338:VAL:HG12	3:T:440:LEU:HG	1.20	1.13
6:X:787:ARG:HH21	6:X:787:ARG:HG3	0.99	1.13

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:d:23:THR:HG23	3:D:323:SER:C	1.69	1.13
1:j:45:ARG:HA	3:J:399:ALA:O	1.44	1.13
1:r:59:HIS:CB	3:R:398:SER:HB2	1.76	1.13
3:D:609:ARG:HB3	2:E:29:THR:HG22	1.27	1.13
3:T:375:LEU:HD23	3:T:437:ILE:HD11	1.31	1.13
6:X:1149:THR:HG21	6:X:1161:ASN:HA	1.29	1.13
1:n:15:ARG:NH1	3:N:402:ALA:HB1	1.62	1.13
1:r:8:GLN:CG	3:T:302:GLU:HG3	1.79	1.13
3:F:338:VAL:HG12	3:F:440:LEU:HG	1.20	1.13
3:L:375:LEU:HD23	3:L:437:ILE:HD11	1.31	1.13
1:l:8:GLN:NE2	3:N:302:GLU:C	2.04	1.13
1:n:8:GLN:NE2	3:J:302:GLU:HG2	1.63	1.13
1:b:15:ARG:CZ	3:B:402:ALA:CB	2.27	1.12
1:f:45:ARG:HG3	3:F:401:GLY:N	1.63	1.12
1:p:45:ARG:CA	3:P:399:ALA:O	1.96	1.12
1:t:6:ILE:HD11	3:P:586:VAL:CG2	1.78	1.12
3:B:431:ILE:HG22	3:B:431:ILE:O	1.49	1.12
1:b:24:LEU:HD21	1:b:80:ARG:HD2	1.13	1.12
1:b:59:HIS:HA	3:B:398:SER:CB	1.78	1.12
1:h:15:ARG:NH1	3:H:406:ASN:OD1	1.82	1.12
1:n:45:ARG:CG	3:N:400:ALA:CA	2.26	1.12
1:r:5:MET:HE2	1:r:5:MET:HA	1.26	1.12
3:B:375:LEU:HD23	3:B:437:ILE:HD11	1.31	1.12
3:N:358:LEU:HD13	3:N:417:PHE:CE1	1.84	1.12
3:P:358:LEU:HD13	3:P:417:PHE:HE1	1.12	1.12
1:j:15:ARG:NH1	3:J:406:ASN:HB3	1.65	1.12
3:H:358:LEU:HD13	3:H:417:PHE:CE1	1.84	1.12
3:L:358:LEU:HD13	3:L:417:PHE:CE1	1.84	1.12
6:Y:355:LYS:NZ	6:Y:1156:THR:HG22	1.64	1.12
1:b:16:ALA:O	1:b:22:LEU:CD2	1.98	1.12
1:p:24:LEU:CD2	1:p:80:ARG:CD	2.25	1.12
1:p:58:PRO:HG3	3:P:423:GLN:HE21	1.10	1.12
7:C:101:MYR:O1	3:D:204:PRO:HG3	1.49	1.12
7:K:101:MYR:H22	3:L:191:LYS:HZ3	1.09	1.12
3:N:431:ILE:HG22	3:N:431:ILE:O	1.49	1.12
3:R:628:GLN:HE21	5:W:870:ALA:CA	1.63	1.12
3:B:338:VAL:HG12	3:B:440:LEU:HG	1.20	1.12
3:R:237:GLU:HB2	3:T:46:LYS:HZ1	1.15	1.12
1:j:8:GLN:CG	3:L:302:GLU:CG	2.16	1.11
1:t:60:ALA:HB2	3:T:322:TYR:OH	1.50	1.11
7:A:101:MYR:O1	3:B:204:PRO:HG3	1.49	1.11

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:D:375:LEU:HD23	3:D:437:ILE:HD11	1.31	1.11
3:F:358:LEU:HD13	3:F:417:PHE:CE1	1.84	1.11
3:R:358:LEU:HD13	3:R:417:PHE:CE1	1.84	1.11
4:U:32:CYS:SG	6:X:958:HIS:NE2	2.23	1.11
3:J:46:LYS:HZ1	3:N:237:GLU:HB2	1.15	1.11
3:J:375:LEU:HD23	3:J:437:ILE:HD11	1.31	1.11
5:W:969:ASP:HB2	5:W:972:PRO:HD2	1.31	1.11
3:D:63:ILE:H	3:D:63:ILE:HD13	0.95	1.11
3:R:358:LEU:HD13	3:R:417:PHE:HE1	1.12	1.11
1:l:24:LEU:HD11	1:l:33:THR:HA	1.26	1.11
1:p:44:GLY:O	3:P:399:ALA:N	1.84	1.11
7:I:101:MYR:O1	3:J:204:PRO:HG3	1.49	1.11
3:P:116:SER:HB3	3:P:133:ASN:H	1.16	1.11
3:P:358:LEU:HD13	3:P:417:PHE:CE1	1.85	1.11
3:T:358:LEU:HD13	3:T:417:PHE:CE1	1.84	1.11
1:d:45:ARG:HA	3:D:399:ALA:O	1.50	1.11
1:n:15:ARG:CZ	3:N:402:ALA:CB	2.29	1.11
3:B:358:LEU:HD13	3:B:417:PHE:CE1	1.84	1.11
3:F:375:LEU:HD23	3:F:437:ILE:HD11	1.31	1.11
3:H:274:ILE:HD11	3:H:514:LEU:HD22	1.27	1.11
3:P:341:MET:HE2	3:P:341:MET:HA	1.30	1.11
1:b:15:ARG:HH11	3:B:406:ASN:CB	1.64	1.10
1:h:6:ILE:HD11	3:D:586:VAL:CG2	1.79	1.10
1:j:23:THR:HG21	3:J:324:GLY:N	1.65	1.10
2:C:29:THR:HG22	3:H:609:ARG:HB3	1.27	1.10
3:D:237:GLU:HB2	3:F:46:LYS:HZ1	1.16	1.10
3:J:238:VAL:HG21	3:L:78:MET:HE1	1.11	1.10
5:W:62:TYR:HB3	5:W:63:PRO:HD2	1.32	1.10
1:h:45:ARG:HG2	3:H:400:ALA:HA	1.11	1.10
1:h:59:HIS:CE1	1:h:60:ALA:O	2.03	1.10
3:F:237:GLU:HB2	3:H:46:LYS:HZ1	1.16	1.10
3:J:358:LEU:HD13	3:J:417:PHE:CE1	1.84	1.10
3:R:431:ILE:HG22	3:R:431:ILE:O	1.49	1.10
3:T:277:ASP:O	3:T:278:THR:HG22	1.52	1.10
6:X:827:THR:HG22	6:X:828:GLU:H	1.15	1.10
3:D:358:LEU:HD13	3:D:417:PHE:CE1	1.84	1.10
3:F:291:PRO:HB2	3:F:467:THR:HG22	1.29	1.10
3:H:277:ASP:O	3:H:278:THR:HG22	1.52	1.10
3:N:277:ASP:O	3:N:278:THR:HG22	1.52	1.10
3:P:274:ILE:HD11	3:P:514:LEU:HD22	1.28	1.10
1:j:24:LEU:HD11	1:j:41:VAL:HG22	1.20	1.10

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:r:45:ARG:HG2	3:R:400:ALA:HA	1.12	1.10
3:J:341:MET:HE2	3:J:341:MET:HA	1.30	1.10
3:L:237:GLU:HB2	3:N:46:LYS:HZ1	1.16	1.10
3:N:375:LEU:HD23	3:N:437:ILE:HD11	1.31	1.10
6:X:1012:ARG:HD3	6:X:1012:ARG:H	1.07	1.10
1:b:23:THR:HB	3:B:323:SER:CA	1.80	1.10
1:r:59:HIS:HA	3:R:398:SER:OG	1.50	1.10
3:B:274:ILE:HD11	3:B:514:LEU:HD22	1.28	1.10
2:I:29:THR:HG22	3:N:609:ARG:HB3	1.27	1.10
3:J:51:VAL:HG12	3:J:60:SER:O	1.52	1.10
3:L:341:MET:HE2	3:L:341:MET:HA	1.30	1.10
1:f:24:LEU:CD2	1:f:80:ARG:CD	2.27	1.09
1:f:59:HIS:HB2	3:F:398:SER:HB2	1.15	1.09
1:d:45:ARG:CG	3:D:400:ALA:CA	2.29	1.09
1:p:45:ARG:HG3	3:P:400:ALA:CA	1.72	1.09
3:D:341:MET:HE2	3:D:341:MET:HA	1.31	1.09
2:E:41:LEU:O	2:E:42:ASN:HB3	1.49	1.09
3:F:116:SER:HB3	3:F:133:ASN:H	1.16	1.09
3:L:291:PRO:HB3	3:L:467:THR:HG22	1.35	1.09
3:P:161:LYS:NZ	3:T:531:THR:HB	1.65	1.09
3:P:238:VAL:HG21	3:R:78:MET:HE1	1.11	1.09
3:T:116:SER:HB3	3:T:133:ASN:H	1.16	1.09
5:W:971:VAL:HG13	5:W:972:PRO:HD3	1.20	1.09
6:X:801:ARG:HH11	6:X:801:ARG:HB2	1.09	1.09
1:l:15:ARG:NH1	3:L:402:ALA:HB1	1.67	1.09
7:G:101:MYR:O1	3:H:204:PRO:HG3	1.53	1.09
3:L:291:PRO:HB2	3:L:467:THR:HG22	1.29	1.09
3:L:609:ARG:HB3	2:M:29:THR:HG22	1.27	1.09
3:R:277:ASP:O	3:R:278:THR:HG22	1.52	1.09
1:d:15:ARG:HH11	3:D:406:ASN:CG	1.60	1.09
3:F:277:ASP:O	3:F:278:THR:HG22	1.52	1.09
3:J:358:LEU:HD13	3:J:417:PHE:HE1	1.12	1.09
3:L:338:VAL:HG12	3:L:440:LEU:HG	1.20	1.09
1:d:59:HIS:HA	3:D:398:SER:CB	1.81	1.09
1:h:58:PRO:HB3	3:H:423:GLN:NE2	1.68	1.09
1:j:24:LEU:HD12	1:j:41:VAL:HG22	1.27	1.09
1:t:59:HIS:CB	3:T:398:SER:HB2	1.74	1.09
7:A:101:MYR:H101	7:A:101:MYR:H142	1.34	1.09
3:B:386:THR:HG22	3:B:387:THR:HG23	1.35	1.09
7:K:101:MYR:H101	7:K:101:MYR:H142	1.34	1.09
6:Y:374:ALA:HB1	6:Y:377:ILE:HD11	1.34	1.09

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:l:21:ARG:O	3:L:323:SER:CB	2.00	1.09
1:j:59:HIS:HB2	3:J:398:SER:CA	1.81	1.09
1:n:23:THR:HG23	1:n:25:TYR:CE2	1.88	1.09
3:B:341:MET:HE2	3:B:341:MET:HA	1.30	1.09
3:D:274:ILE:HD11	3:D:514:LEU:HD22	1.28	1.09
3:F:431:ILE:HG22	3:F:431:ILE:O	1.49	1.09
3:F:609:ARG:HB3	2:G:29:THR:HG22	1.27	1.09
3:N:338:VAL:HG12	3:N:440:LEU:HG	1.20	1.09
3:P:78:MET:HE1	3:T:238:VAL:HG21	1.11	1.09
3:P:277:ASP:O	3:P:278:THR:HG22	1.52	1.09
3:T:358:LEU:HD13	3:T:417:PHE:HE1	1.12	1.09
1:l:21:ARG:HD3	3:L:327:TYR:OH	1.53	1.08
1:l:59:HIS:HD2	3:L:397:VAL:HG22	1.11	1.08
1:d:45:ARG:O	1:d:58:PRO:O	1.69	1.08
1:r:16:ALA:C	1:r:22:LEU:HD21	1.77	1.08
1:r:23:THR:HG21	1:r:25:TYR:CZ	1.87	1.08
3:B:277:ASP:O	3:B:278:THR:HG22	1.52	1.08
3:D:338:VAL:HG12	3:D:440:LEU:HG	1.20	1.08
7:M:101:MYR:O1	3:N:204:PRO:HG3	1.53	1.08
3:N:116:SER:HB3	3:N:133:ASN:H	1.16	1.08
3:N:274:ILE:HD11	3:N:514:LEU:HD22	1.27	1.08
1:j:15:ARG:HH11	3:J:406:ASN:CG	1.59	1.08
1:t:44:GLY:CA	3:T:322:TYR:OH	2.00	1.08
3:B:274:ILE:HG21	3:B:601:THR:HG21	1.08	1.08
3:D:277:ASP:O	3:D:278:THR:HG22	1.52	1.08
3:J:291:PRO:HB2	3:J:467:THR:HG22	1.29	1.08
3:P:291:PRO:HB2	3:P:467:THR:HG22	1.30	1.08
3:R:375:LEU:HD23	3:R:437:ILE:HD11	1.31	1.08
7:S:101:MYR:H101	7:S:101:MYR:H142	1.34	1.08
3:T:358:LEU:HD22	3:T:417:PHE:CE1	1.89	1.08
3:T:386:THR:HG22	3:T:387:THR:HG23	1.35	1.08
6:Y:996:MET:HA	6:Y:996:MET:HE2	1.28	1.08
1:d:58:PRO:HG3	3:D:423:GLN:HE21	0.99	1.08
1:h:45:ARG:HG3	3:H:401:GLY:N	1.68	1.08
1:p:61:ASN:HB3	3:P:397:VAL:HA	1.31	1.08
2:C:41:LEU:O	2:C:42:ASN:HB3	1.49	1.08
3:D:291:PRO:HB2	3:D:467:THR:HG22	1.29	1.08
3:L:358:LEU:HD22	3:L:417:PHE:CE1	1.89	1.08
3:L:386:THR:HG22	3:L:387:THR:HG23	1.35	1.08
3:P:358:LEU:HD22	3:P:417:PHE:CE1	1.89	1.08
3:R:274:ILE:HG21	3:R:601:THR:HG21	1.08	1.08

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:p:59:HIS:HA	3:P:398:SER:HA	1.33	1.08
1:r:59:HIS:HB2	3:R:398:SER:HB2	1.13	1.08
1:t:8:GLN:NE2	3:P:302:GLU:HG2	1.69	1.08
1:t:24:LEU:HD23	1:t:35:PHE:HA	1.14	1.08
3:H:375:LEU:HD23	3:H:437:ILE:HD11	1.31	1.08
3:J:78:MET:HE1	3:N:238:VAL:HG21	1.11	1.08
3:J:274:ILE:HG21	3:J:601:THR:HG21	1.08	1.08
3:T:274:ILE:HG21	3:T:601:THR:HG21	1.08	1.08
1:j:58:PRO:HG2	3:J:423:GLN:HE21	1.08	1.08
1:n:45:ARG:HA	3:N:399:ALA:O	1.53	1.08
1:r:23:THR:HG21	1:r:25:TYR:OH	1.52	1.08
1:t:15:ARG:HH11	3:T:406:ASN:HB3	1.19	1.08
1:t:59:HIS:HB2	3:T:398:SER:HA	1.12	1.08
2:A:41:LEU:O	2:A:42:ASN:HB3	1.49	1.08
3:D:274:ILE:HG21	3:D:601:THR:HG21	1.08	1.08
3:D:358:LEU:HD22	3:D:417:PHE:CE1	1.89	1.08
7:E:101:MYR:H101	7:E:101:MYR:H142	1.34	1.08
2:K:41:LEU:O	2:K:42:ASN:HB3	1.49	1.08
3:L:238:VAL:HG21	3:N:78:MET:HE1	1.11	1.08
3:R:291:PRO:HB2	3:R:467:THR:HG22	1.29	1.08
1:d:4:HIS:CD2	1:d:54:THR:CG2	2.37	1.07
1:t:15:ARG:HH11	3:T:406:ASN:CB	1.67	1.07
7:C:101:MYR:H101	7:C:101:MYR:H142	1.34	1.07
3:D:386:THR:HG22	3:D:387:THR:HG23	1.35	1.07
3:D:431:ILE:HG22	3:D:431:ILE:O	1.49	1.07
3:F:358:LEU:HD22	3:F:417:PHE:CE1	1.89	1.07
3:P:431:ILE:HG22	3:P:431:ILE:O	1.49	1.07
3:R:358:LEU:HD22	3:R:417:PHE:CE1	1.89	1.07
3:T:291:PRO:HB2	3:T:467:THR:HG22	1.29	1.07
1:l:6:ILE:HD11	3:N:586:VAL:CG2	1.84	1.07
1:r:6:ILE:HD11	3:T:586:VAL:CG2	1.84	1.07
1:r:60:ALA:N	3:R:398:SER:HA	1.69	1.07
1:t:45:ARG:O	1:t:59:HIS:N	1.87	1.07
3:F:341:MET:HE2	3:F:341:MET:HA	1.30	1.07
2:O:41:LEU:O	2:O:42:ASN:HB3	1.49	1.07
7:S:101:MYR:O1	3:T:204:PRO:HG3	1.53	1.07
3:T:291:PRO:HB3	3:T:467:THR:HG22	1.35	1.07
1:f:45:ARG:HG2	3:F:400:ALA:HA	1.11	1.07
1:f:59:HIS:CG	3:F:397:VAL:O	2.08	1.07
7:E:101:MYR:H22	3:F:191:LYS:HZ3	1.12	1.07
3:H:358:LEU:HD22	3:H:417:PHE:CE1	1.89	1.07

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:L:274:ILE:HD11	3:L:514:LEU:HD22	1.28	1.07
3:L:277:ASP:O	3:L:278:THR:HG22	1.52	1.07
3:P:375:LEU:HD23	3:P:437:ILE:HD11	1.31	1.07
3:P:386:THR:HG22	3:P:387:THR:HG23	1.35	1.07
1:h:59:HIS:ND1	3:H:397:VAL:O	1.87	1.07
1:p:15:ARG:HH11	3:P:406:ASN:HB3	0.95	1.07
1:p:60:ALA:O	1:p:62:VAL:N	1.87	1.07
1:t:45:ARG:HG3	3:T:401:GLY:N	1.69	1.07
3:B:116:SER:HB3	3:B:133:ASN:H	1.16	1.07
3:D:46:LYS:HZ1	3:H:237:GLU:HB2	1.20	1.07
3:D:291:PRO:HB3	3:D:467:THR:HG22	1.35	1.07
3:F:291:PRO:HB3	3:F:467:THR:HG22	1.35	1.07
3:H:341:MET:HE2	3:H:341:MET:HA	1.30	1.07
3:J:116:SER:HB3	3:J:133:ASN:H	1.16	1.07
7:M:101:MYR:H102	3:N:200:CYS:SG	1.95	1.07
3:N:358:LEU:HD22	3:N:417:PHE:CE1	1.89	1.07
1:l:59:HIS:CD2	3:L:397:VAL:HG22	1.90	1.06
1:b:15:ARG:NH1	3:B:406:ASN:HB3	1.68	1.06
1:b:23:THR:HB	3:B:323:SER:HA	1.07	1.06
1:p:35:PHE:CE2	1:p:73:ARG:CG	2.38	1.06
3:D:238:VAL:HG21	3:F:78:MET:HE1	1.11	1.06
2:Q:41:LEU:O	2:Q:42:ASN:HB3	1.49	1.06
3:B:358:LEU:HD22	3:B:417:PHE:CE1	1.89	1.06
3:L:116:SER:HB3	3:L:133:ASN:H	1.16	1.06
6:X:606:MET:CE	6:Y:718:THR:CG2	2.33	1.06
1:b:15:ARG:HH11	3:B:406:ASN:HB3	1.16	1.06
1:r:8:GLN:NE2	3:T:302:GLU:HG2	1.69	1.06
7:G:101:MYR:H102	3:H:200:CYS:SG	1.95	1.06
3:L:431:ILE:HG22	3:L:431:ILE:O	1.49	1.06
3:N:161:LYS:HD3	3:N:161:LYS:C	1.79	1.06
3:N:386:THR:HG22	3:N:387:THR:HG23	1.35	1.06
1:f:45:ARG:HG3	3:F:401:GLY:H	1.11	1.06
1:d:8:GLN:CG	3:F:302:GLU:CG	2.27	1.06
1:d:57:ALA:HB1	1:d:59:HIS:NE2	1.71	1.06
3:B:443:PHE:HB2	3:B:457:PRO:HB2	1.38	1.06
3:D:530:ARG:HE	3:H:472:ARG:NH2	1.53	1.06
3:J:277:ASP:O	3:J:278:THR:HG22	1.52	1.06
3:J:358:LEU:HD22	3:J:417:PHE:CE1	1.89	1.06
3:N:274:ILE:HG21	3:N:601:THR:HG21	1.08	1.06
3:R:545:VAL:HG21	3:T:637:PHE:HE1	1.21	1.06
7:S:101:MYR:H102	3:T:200:CYS:SG	1.95	1.06

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:l:23:THR:CG2	1:l:25:TYR:CE2	2.39	1.06
1:d:59:HIS:HB2	3:D:397:VAL:O	1.54	1.06
1:j:23:THR:HG23	3:J:323:SER:O	1.54	1.06
3:D:472:ARG:NH2	3:F:530:ARG:HE	1.54	1.06
2:I:41:LEU:O	2:I:42:ASN:HB3	1.49	1.06
7:I:101:MYR:H101	7:I:101:MYR:H142	1.34	1.06
3:N:341:MET:HE2	3:N:341:MET:HA	1.31	1.06
3:P:46:LYS:HZ1	3:T:237:GLU:HB2	1.15	1.06
3:P:291:PRO:HB3	3:P:467:THR:HG22	1.35	1.06
3:R:116:SER:HB3	3:R:133:ASN:H	1.16	1.06
1:l:45:ARG:HG2	3:L:400:ALA:HA	1.13	1.05
1:f:6:ILE:HD11	3:H:586:VAL:CG2	1.84	1.05
3:D:116:SER:HB3	3:D:133:ASN:H	1.16	1.05
3:D:298:PHE:CZ	3:H:355:GLY:HA2	1.91	1.05
3:H:291:PRO:HB3	3:H:467:THR:HG22	1.35	1.05
7:M:101:MYR:H101	7:M:101:MYR:H142	1.34	1.05
3:P:443:PHE:HB2	3:P:457:PRO:HB2	1.38	1.05
7:Q:101:MYR:O1	3:R:204:PRO:HG3	1.56	1.05
3:T:341:MET:HE2	3:T:341:MET:HA	1.31	1.05
1:d:8:GLN:CD	3:F:302:GLU:HG2	1.80	1.05
3:D:355:GLY:HA2	3:F:298:PHE:CZ	1.91	1.05
3:F:274:ILE:HG21	3:F:601:THR:HG21	1.08	1.05
3:J:237:GLU:HB2	3:L:46:LYS:HZ1	1.15	1.05
3:L:274:ILE:HG21	3:L:601:THR:HG21	1.08	1.05
3:P:637:PHE:HE1	3:T:545:VAL:HG21	1.21	1.05
3:T:431:ILE:HG22	3:T:431:ILE:O	1.49	1.05
1:l:45:ARG:CG	3:L:400:ALA:CA	2.34	1.05
3:F:87:PHE:CD1	3:H:121:ALA:HB2	1.92	1.05
7:G:101:MYR:H101	7:G:101:MYR:H142	1.34	1.05
3:J:298:PHE:CZ	3:N:355:GLY:HA2	1.91	1.05
3:J:530:ARG:HE	3:N:472:ARG:NH2	1.54	1.05
3:N:443:PHE:HB2	3:N:457:PRO:HB2	1.38	1.05
3:P:298:PHE:CZ	3:T:355:GLY:HA2	1.91	1.05
1:l:45:ARG:HG3	3:L:401:GLY:N	1.72	1.05
1:b:15:ARG:NH2	3:B:402:ALA:CB	2.20	1.05
1:j:20:GLY:HA3	1:j:83:VAL:HG13	1.38	1.05
3:B:291:PRO:HB3	3:B:467:THR:HG22	1.35	1.05
3:D:78:MET:HE1	3:H:238:VAL:HG21	1.11	1.05
3:F:238:VAL:HG21	3:H:78:MET:HE1	1.11	1.05
3:H:431:ILE:HG22	3:H:431:ILE:O	1.49	1.05
3:L:355:GLY:HA2	3:N:298:PHE:CZ	1.91	1.05

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:N:322:TYR:HD2	3:N:327:TYR:CE2	1.74	1.05
4:U:198:GLN:HE22	6:X:661:THR:HG22	1.16	1.05
1:l:23:THR:OG1	3:L:323:SER:HA	1.57	1.05
1:l:60:ALA:H	3:L:398:SER:HA	1.12	1.05
1:b:24:LEU:HD21	1:b:80:ARG:CD	1.86	1.05
6:X:437:ARG:HH22	6:X:1207:ILE:HA	1.17	1.05
1:h:8:GLN:NE2	3:D:302:GLU:HG2	1.72	1.04
1:r:58:PRO:HG2	3:R:423:GLN:HE21	1.13	1.04
3:D:53:THR:HG22	3:D:54:SER:H	1.17	1.04
3:D:358:LEU:HD13	3:D:417:PHE:HE1	1.12	1.04
3:H:116:SER:HB3	3:H:133:ASN:H	1.15	1.04
3:N:358:LEU:HD13	3:N:417:PHE:HE1	1.12	1.04
3:R:355:GLY:HA2	3:T:298:PHE:CZ	1.91	1.04
3:R:386:THR:HG22	3:R:387:THR:HG23	1.35	1.04
4:U:32:CYS:SG	6:X:958:HIS:CE1	2.50	1.04
6:X:464:SER:HB2	6:Y:500:ILE:HG21	1.39	1.04
6:X:616:SER:HB3	6:Y:723:ASP:CB	1.86	1.04
6:Y:690:ALA:HB3	6:Y:691:PRO:HD3	1.38	1.04
1:h:45:ARG:CG	3:H:400:ALA:CA	2.35	1.04
1:t:15:ARG:NH1	3:T:402:ALA:HB1	1.71	1.04
7:E:101:MYR:O1	3:F:204:PRO:HG3	1.56	1.04
3:J:121:ALA:HB2	3:N:87:PHE:CD1	1.92	1.04
7:K:101:MYR:O1	3:L:204:PRO:HG3	1.56	1.04
3:P:121:ALA:HB2	3:T:87:PHE:CD1	1.92	1.04
3:R:238:VAL:HG21	3:T:78:MET:HE1	1.11	1.04
3:T:443:PHE:HB2	3:T:457:PRO:HB2	1.38	1.04
1:b:45:ARG:CA	3:B:399:ALA:O	2.04	1.04
1:n:15:ARG:HH11	3:N:406:ASN:HB3	1.22	1.04
1:n:45:ARG:HG2	3:N:400:ALA:HA	1.09	1.04
1:p:59:HIS:CB	3:P:398:SER:CB	2.20	1.04
3:D:142:ILE:HD11	3:H:238:VAL:CG2	1.86	1.04
3:J:87:PHE:CD1	3:L:121:ALA:HB2	1.92	1.04
3:J:291:PRO:HB3	3:J:467:THR:HG22	1.35	1.04
3:L:87:PHE:CD1	3:N:121:ALA:HB2	1.92	1.04
3:L:443:PHE:HB2	3:L:457:PRO:HB2	1.38	1.04
3:N:322:TYR:CD2	3:N:327:TYR:CE2	2.44	1.04
3:P:87:PHE:CD1	3:R:121:ALA:HB2	1.92	1.04
3:P:274:ILE:HG21	3:P:601:THR:HG21	1.08	1.04
1:h:59:HIS:HB2	3:H:398:SER:CA	1.87	1.04
1:p:45:ARG:HG3	3:P:400:ALA:C	1.81	1.04
3:D:87:PHE:CD1	3:F:121:ALA:HB2	1.92	1.04

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:D:121:ALA:HB2	3:H:87:PHE:CD1	1.92	1.04
3:F:355:GLY:HA2	3:H:298:PHE:CZ	1.91	1.04
3:F:443:PHE:HB2	3:F:457:PRO:HB2	1.38	1.04
3:P:237:GLU:HB2	3:R:46:LYS:HZ1	1.20	1.04
3:R:291:PRO:HB3	3:R:467:THR:HG22	1.35	1.04
6:X:571:LEU:HD11	6:X:705:ILE:HD11	1.39	1.04
6:Y:557:ILE:HD12	6:Y:593:LEU:HD22	1.36	1.04
1:b:45:ARG:CG	3:B:401:GLY:H	1.69	1.04
1:h:15:ARG:NH1	3:H:402:ALA:HB1	1.73	1.04
1:j:45:ARG:HG3	3:J:400:ALA:CA	1.88	1.04
1:n:6:ILE:HD11	3:J:586:VAL:CG2	1.86	1.04
2:C:26:MET:HG3	3:D:216:PRO:HG3	1.40	1.04
3:D:443:PHE:HB2	3:D:457:PRO:HB2	1.38	1.04
3:J:355:GLY:HA2	3:L:298:PHE:CZ	1.91	1.04
7:O:101:MYR:H101	7:O:101:MYR:H142	1.34	1.04
3:P:355:GLY:HA2	3:R:298:PHE:CZ	1.91	1.04
2:Q:26:MET:HG3	3:R:216:PRO:HG3	1.40	1.04
4:V:192:PHE:HB2	6:Y:376:LEU:CD2	1.86	1.04
1:l:8:GLN:NE2	3:N:302:GLU:HG2	1.71	1.03
1:b:45:ARG:HG3	3:B:400:ALA:CA	1.84	1.03
7:G:101:MYR:H22	3:H:191:LYS:HZ3	1.21	1.03
3:H:274:ILE:HG21	3:H:601:THR:HG21	1.08	1.03
3:H:386:THR:HG22	3:H:387:THR:HG23	1.35	1.03
7:Q:101:MYR:H101	7:Q:101:MYR:H142	1.34	1.03
1:f:24:LEU:HD21	1:f:80:ARG:HD2	1.05	1.03
1:n:23:THR:HB	3:N:323:SER:CB	1.89	1.03
1:r:45:ARG:CG	3:R:401:GLY:H	1.72	1.03
1:t:22:LEU:CD1	1:t:83:VAL:HG21	1.87	1.03
7:C:101:MYR:H102	3:D:200:CYS:SG	1.98	1.03
3:D:142:ILE:CD1	3:H:238:VAL:HG22	1.86	1.03
3:F:386:THR:HG22	3:F:387:THR:HG23	1.35	1.03
3:J:431:ILE:HG22	3:J:431:ILE:O	1.49	1.03
7:O:101:MYR:H102	3:P:200:CYS:SG	1.98	1.03
1:p:24:LEU:HD21	1:p:80:ARG:HD2	1.04	1.03
7:A:101:MYR:H102	3:B:200:CYS:SG	1.98	1.03
3:R:87:PHE:CD1	3:T:121:ALA:HB2	1.92	1.03
1:d:8:GLN:NE2	3:F:302:GLU:HG2	1.73	1.03
1:j:15:ARG:NH1	3:J:406:ASN:CG	2.15	1.03
1:n:45:ARG:HG3	3:N:401:GLY:N	1.73	1.03
1:t:23:THR:CG2	1:t:31:GLU:HB2	1.88	1.03
3:D:431:ILE:O	3:D:431:ILE:CG2	2.06	1.03

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:J:443:PHE:HB2	3:J:457:PRO:HB2	1.38	1.03
3:N:291:PRO:HB3	3:N:467:THR:HG22	1.35	1.03
6:X:528:PHE:HE1	6:X:575:MET:HE2	1.23	1.03
6:X:925:LEU:HD21	6:Y:496:ALA:HA	1.40	1.03
1:l:15:ARG:CZ	3:L:402:ALA:CB	2.37	1.03
1:t:44:GLY:O	3:T:399:ALA:N	1.91	1.03
7:C:101:MYR:H22	3:D:191:LYS:HZ3	1.20	1.03
3:H:291:PRO:HB2	3:H:467:THR:HG22	1.29	1.03
7:I:101:MYR:H102	3:J:200:CYS:SG	1.98	1.03
3:J:386:THR:HG22	3:J:387:THR:HG23	1.35	1.03
1:j:23:THR:CG2	3:J:323:SER:O	2.03	1.02
1:p:8:GLN:HE22	3:R:303:ASP:HA	1.21	1.02
1:t:15:ARG:CZ	3:T:402:ALA:CB	2.37	1.02
3:F:431:ILE:O	3:F:431:ILE:CG2	2.07	1.02
3:H:443:PHE:HB2	3:H:457:PRO:HB2	1.38	1.02
2:K:26:MET:HG3	3:L:216:PRO:HG3	1.40	1.02
1:h:45:ARG:HG3	3:H:401:GLY:H	1.19	1.02
2:I:26:MET:HG3	3:J:216:PRO:HG3	1.40	1.02
3:N:431:ILE:O	3:N:431:ILE:CG2	2.07	1.02
6:Y:277:VAL:HG22	6:Y:299:LEU:HD22	1.41	1.02
1:l:23:THR:CB	3:L:323:SER:HA	1.90	1.02
1:d:45:ARG:HG3	3:D:401:GLY:H	1.11	1.02
1:h:58:PRO:HB3	3:H:423:GLN:HE21	0.87	1.02
1:j:45:ARG:CG	3:J:401:GLY:H	1.73	1.02
1:p:4:HIS:CD2	1:p:54:THR:HG22	1.93	1.02
1:t:75:PRO:HB3	1:t:77:GLU:HG3	1.38	1.02
3:D:545:VAL:HG21	3:F:637:PHE:HE1	1.21	1.02
6:X:787:ARG:HG3	6:X:787:ARG:NH2	1.68	1.02
1:l:15:ARG:NH1	3:L:406:ASN:OD1	1.92	1.02
1:n:44:GLY:O	3:N:399:ALA:N	1.92	1.02
1:p:45:ARG:HG2	3:P:400:ALA:CA	1.78	1.02
3:R:431:ILE:O	3:R:431:ILE:CG2	2.06	1.02
3:R:443:PHE:HB2	3:R:457:PRO:HB2	1.38	1.02
6:X:603:VAL:HG11	6:Y:715:VAL:CG1	1.90	1.02
6:Y:235:LEU:HB2	6:Y:356:ASN:HD21	1.24	1.02
1:d:45:ARG:HG3	3:D:400:ALA:HA	1.42	1.02
1:r:23:THR:CG2	1:r:25:TYR:CZ	2.43	1.02
3:B:431:ILE:HD13	3:B:437:ILE:CG2	1.90	1.02
3:B:431:ILE:CD1	3:B:437:ILE:HG21	1.90	1.02
3:D:431:ILE:HD13	3:D:437:ILE:CG2	1.90	1.02
3:R:472:ARG:NH2	3:T:530:ARG:HE	1.58	1.02

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
4:U:178:THR:HG21	6:X:376:LEU:HD23	1.39	1.02
6:Y:355:LYS:HZ1	6:Y:1156:THR:HG22	1.18	1.02
1:b:45:ARG:HG2	3:B:400:ALA:CA	1.78	1.01
3:F:431:ILE:HD13	3:F:437:ILE:CG2	1.90	1.01
3:F:545:VAL:HG21	3:H:637:PHE:HE1	1.21	1.01
7:G:101:MYR:H101	7:G:101:MYR:H143	1.42	1.01
3:R:431:ILE:CD1	3:R:437:ILE:HG21	1.90	1.01
3:T:431:ILE:O	3:T:431:ILE:CG2	2.07	1.01
3:T:431:ILE:HD13	3:T:437:ILE:CG2	1.90	1.01
1:b:44:GLY:CA	3:B:322:TYR:OH	2.08	1.01
1:b:45:ARG:HG3	3:B:400:ALA:C	1.85	1.01
1:t:23:THR:HG21	1:t:31:GLU:HB2	1.41	1.01
7:E:101:MYR:H102	3:F:200:CYS:SG	2.00	1.01
3:F:431:ILE:HD13	3:F:437:ILE:HG21	1.43	1.01
3:F:431:ILE:CD1	3:F:437:ILE:HG21	1.90	1.01
3:L:431:ILE:HD13	3:L:437:ILE:CG2	1.90	1.01
4:V:175:MET:SD	6:Y:400:GLU:HG2	2.00	1.01
1:f:15:ARG:HH11	3:F:406:ASN:CB	1.72	1.01
1:f:59:HIS:HB3	3:F:398:SER:HB2	1.39	1.01
3:B:358:LEU:CD1	3:B:417:PHE:HE1	1.74	1.01
3:B:431:ILE:O	3:B:431:ILE:CG2	2.06	1.01
3:D:431:ILE:HD13	3:D:437:ILE:HG21	1.43	1.01
2:E:26:MET:HG3	3:F:216:PRO:HG3	1.40	1.01
3:H:431:ILE:O	3:H:431:ILE:CG2	2.06	1.01
3:J:431:ILE:HD13	3:J:437:ILE:HG21	1.43	1.01
3:J:637:PHE:HE1	3:N:545:VAL:HG21	1.21	1.01
3:L:431:ILE:CD1	3:L:437:ILE:HG21	1.90	1.01
3:L:545:VAL:HG21	3:N:637:PHE:HE1	1.21	1.01
3:P:431:ILE:O	3:P:431:ILE:CG2	2.06	1.01
3:P:431:ILE:HD13	3:P:437:ILE:HG21	1.43	1.01
3:R:431:ILE:HD13	3:R:437:ILE:CG2	1.90	1.01
1:d:15:ARG:HH11	3:D:406:ASN:HB3	1.26	1.01
3:J:431:ILE:O	3:J:431:ILE:CG2	2.06	1.01
7:K:101:MYR:H101	7:K:101:MYR:H143	1.42	1.01
3:N:274:ILE:CD1	3:N:514:LEU:CD2	2.39	1.01
3:N:431:ILE:CD1	3:N:437:ILE:HG21	1.90	1.01
5:W:1176:VAL:HG23	5:W:1280:VAL:HG13	1.42	1.01
1:f:15:ARG:HH11	3:F:406:ASN:CG	1.68	1.01
1:f:59:HIS:HB2	3:F:398:SER:CA	1.89	1.01
1:j:15:ARG:HH11	3:J:406:ASN:HB3	1.13	1.01
7:M:101:MYR:H101	7:M:101:MYR:H143	1.42	1.01

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:P:338:VAL:CG1	3:P:440:LEU:HG	1.91	1.01
3:P:431:ILE:HD13	3:P:437:ILE:CG2	1.90	1.01
3:P:545:VAL:HG21	3:R:637:PHE:HE1	1.21	1.01
7:Q:101:MYR:H102	3:R:200:CYS:SG	1.99	1.01
3:R:338:VAL:CG1	3:R:440:LEU:HG	1.91	1.01
6:X:718:THR:HG23	6:X:726:ARG:HH12	1.24	1.01
1:b:21:ARG:HD2	3:B:327:TYR:OH	1.61	1.00
1:p:21:ARG:O	3:P:323:SER:HB2	1.61	1.00
3:D:431:ILE:CD1	3:D:437:ILE:HG21	1.90	1.00
3:H:358:LEU:CD1	3:H:417:PHE:HE1	1.74	1.00
3:J:545:VAL:HG21	3:L:637:PHE:HE1	1.21	1.00
7:K:101:MYR:H102	3:L:200:CYS:SG	1.99	1.00
2:O:26:MET:HG3	3:P:216:PRO:HG3	1.40	1.00
3:T:431:ILE:CD1	3:T:437:ILE:HG21	1.90	1.00
4:U:178:THR:HG22	6:X:376:LEU:HD21	1.37	1.00
6:Y:557:ILE:HG23	6:Y:717:MET:HE1	1.42	1.00
1:d:15:ARG:NH1	3:D:406:ASN:CG	2.18	1.00
1:d:15:ARG:NH1	3:D:406:ASN:HB3	1.76	1.00
1:h:24:LEU:HD11	1:h:80:ARG:HD2	1.43	1.00
1:j:21:ARG:O	3:J:323:SER:HB3	1.62	1.00
1:n:8:GLN:CG	3:J:302:GLU:CG	2.36	1.00
1:p:58:PRO:CG	3:P:423:GLN:CG	2.39	1.00
1:r:58:PRO:HG2	3:R:423:GLN:NE2	1.77	1.00
3:B:274:ILE:CD1	3:B:514:LEU:CD2	2.39	1.00
3:D:142:ILE:HD11	3:H:238:VAL:HG22	1.00	1.00
3:H:431:ILE:CD1	3:H:437:ILE:HG21	1.90	1.00
3:J:431:ILE:HD13	3:J:437:ILE:CG2	1.90	1.00
2:M:26:MET:HG3	3:N:216:PRO:HG3	1.40	1.00
3:N:338:VAL:CG1	3:N:440:LEU:HG	1.91	1.00
3:P:431:ILE:CD1	3:P:437:ILE:HG21	1.90	1.00
3:R:358:LEU:CD1	3:R:417:PHE:HE1	1.74	1.00
1:b:24:LEU:CD2	1:b:80:ARG:HD2	1.89	1.00
1:p:56:LEU:HD13	1:p:65:ILE:CG2	1.91	1.00
7:A:101:MYR:H22	3:B:191:LYS:HZ3	1.25	1.00
3:R:274:ILE:CD1	3:R:514:LEU:CD2	2.39	1.00
3:T:358:LEU:CD1	3:T:417:PHE:HE1	1.74	1.00
5:W:1074:THR:HG22	5:W:1075:PRO:HD2	1.44	1.00
6:Y:542:THR:HG23	6:Y:808:ALA:HB3	1.44	1.00
1:j:15:ARG:NH1	3:J:406:ASN:CB	2.22	1.00
1:n:15:ARG:HH11	3:N:406:ASN:CB	1.73	1.00
3:B:338:VAL:CG1	3:B:440:LEU:HG	1.91	1.00

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:D:274:ILE:CD1	3:D:514:LEU:CD2	2.39	1.00
3:D:545:VAL:HG21	3:F:637:PHE:CE1	1.97	1.00
3:H:431:ILE:HD13	3:H:437:ILE:CG2	1.90	1.00
3:J:431:ILE:CD1	3:J:437:ILE:HG21	1.90	1.00
3:L:431:ILE:O	3:L:431:ILE:CG2	2.07	1.00
1:t:45:ARG:HG3	3:T:400:ALA:HA	1.40	1.00
2:A:26:MET:HG3	3:B:216:PRO:HG3	1.40	1.00
1:p:61:ASN:ND2	3:P:395:THR:CG2	2.24	1.00
1:r:15:ARG:NH1	3:R:406:ASN:CG	2.19	1.00
3:D:637:PHE:HE1	3:H:545:VAL:HG21	1.21	1.00
3:F:545:VAL:HG21	3:H:637:PHE:CE1	1.97	1.00
3:H:338:VAL:CG1	3:H:440:LEU:HG	1.91	1.00
3:J:274:ILE:CD1	3:J:514:LEU:CD2	2.39	1.00
3:L:338:VAL:CG1	3:L:440:LEU:HG	1.91	1.00
3:D:637:PHE:CE1	3:H:545:VAL:HG21	1.97	1.00
3:F:274:ILE:CD1	3:F:514:LEU:CD2	2.39	1.00
3:F:274:ILE:CG2	3:F:601:THR:HG21	1.92	1.00
2:G:26:MET:HG3	3:H:216:PRO:HG3	1.40	1.00
3:H:274:ILE:CG2	3:H:601:THR:HG21	1.92	1.00
3:N:431:ILE:HD13	3:N:437:ILE:CG2	1.90	1.00
3:P:274:ILE:CG2	3:P:601:THR:HG21	1.92	1.00
3:P:274:ILE:CD1	3:P:514:LEU:CD2	2.39	1.00
3:P:637:PHE:CE1	3:T:545:VAL:HG21	1.97	1.00
1:f:45:ARG:HA	3:F:399:ALA:O	1.62	0.99
3:F:338:VAL:CG1	3:F:440:LEU:HG	1.91	0.99
3:H:274:ILE:HG21	3:H:601:THR:CG2	1.92	0.99
3:H:274:ILE:CD1	3:H:514:LEU:CD2	2.39	0.99
3:J:338:VAL:CG1	3:J:440:LEU:HG	1.91	0.99
3:J:637:PHE:CE1	3:N:545:VAL:HG21	1.97	0.99
3:T:338:VAL:CG1	3:T:440:LEU:HG	1.91	0.99
5:W:748:ARG:HH12	5:W:789:ARG:HA	1.26	0.99
1:t:45:ARG:CA	3:T:399:ALA:O	2.10	0.99
3:L:274:ILE:CD1	3:L:514:LEU:CD2	2.39	0.99
3:N:431:ILE:HD13	3:N:437:ILE:HG21	1.43	0.99
3:N:579:ASN:HD22	3:N:582:LEU:HG	1.27	0.99
7:O:101:MYR:H101	7:O:101:MYR:H143	1.42	0.99
3:P:545:VAL:HG21	3:R:637:PHE:CE1	1.97	0.99
6:Y:594:ALA:HA	6:Y:608:GLN:NE2	1.76	0.99
1:h:45:ARG:HA	3:H:399:ALA:O	1.59	0.99
1:t:45:ARG:HG2	3:T:400:ALA:HA	1.04	0.99
7:C:101:MYR:H101	7:C:101:MYR:H143	1.42	0.99

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:J:237:GLU:HG3	3:L:142:ILE:HD13	1.44	0.99
3:J:274:ILE:HG21	3:J:601:THR:CG2	1.92	0.99
3:P:358:LEU:CD1	3:P:417:PHE:HE1	1.74	0.99
6:Y:632:ASN:HB2	6:Y:635:GLN:HB3	1.42	0.99
1:b:44:GLY:O	3:B:399:ALA:N	1.94	0.99
3:D:338:VAL:CG1	3:D:440:LEU:HG	1.91	0.99
7:E:101:MYR:H22	3:F:191:LYS:NZ	1.77	0.99
2:I:27:THR:HG23	3:N:609:ARG:NH1	1.78	0.99
3:N:358:LEU:CD1	3:N:417:PHE:HE1	1.74	0.99
7:O:101:MYR:H22	3:P:191:LYS:HZ3	1.28	0.99
5:W:313:LEU:HD13	5:W:318:VAL:HG11	1.44	0.99
1:l:45:ARG:HG3	3:L:401:GLY:H	1.25	0.99
1:f:45:ARG:CG	3:F:400:ALA:CA	2.39	0.99
1:d:61:ASN:O	1:d:63:LYS:N	1.95	0.99
1:p:61:ASN:ND2	3:P:395:THR:HG21	1.76	0.99
2:C:27:THR:HG23	3:H:609:ARG:NH1	1.78	0.99
3:D:274:ILE:HG21	3:D:601:THR:CG2	1.92	0.99
3:F:358:LEU:CD1	3:F:417:PHE:HE1	1.74	0.99
3:L:358:LEU:CD1	3:L:417:PHE:HE1	1.74	0.99
3:L:545:VAL:HG21	3:N:637:PHE:CE1	1.97	0.99
3:N:322:TYR:CD2	3:N:327:TYR:HE2	1.80	0.99
7:Q:101:MYR:H101	7:Q:101:MYR:H143	1.42	0.99
2:S:26:MET:HG3	3:T:216:PRO:HG3	1.40	0.99
1:l:45:ARG:HA	3:L:399:ALA:O	1.60	0.99
3:D:358:LEU:CD1	3:D:417:PHE:HE1	1.74	0.99
3:J:274:ILE:CG2	3:J:601:THR:HG21	1.92	0.99
4:U:84:ILE:HD11	4:U:92:TRP:HB2	1.39	0.99
1:b:59:HIS:CA	3:B:397:VAL:O	2.10	0.99
1:j:58:PRO:HG2	3:J:423:GLN:NE2	1.76	0.99
3:T:274:ILE:CD1	3:T:514:LEU:CD2	2.39	0.99
1:l:59:HIS:HA	3:L:398:SER:OG	1.62	0.99
3:F:609:ARG:NH1	2:G:27:THR:HG23	1.78	0.99
3:L:274:ILE:HG21	3:L:601:THR:CG2	1.92	0.99
6:X:282:TYR:CD2	6:Y:830:ASN:ND2	2.30	0.99
6:X:603:VAL:HG11	6:Y:715:VAL:HG12	1.42	0.99
1:h:15:ARG:HH11	3:H:406:ASN:CB	1.75	0.99
1:n:8:GLN:HE21	3:J:302:GLU:C	1.65	0.99
1:p:8:GLN:CG	3:R:302:GLU:HG2	1.89	0.99
3:F:274:ILE:HG21	3:F:601:THR:CG2	1.92	0.99
3:J:358:LEU:CD1	3:J:417:PHE:HE1	1.74	0.99
3:J:545:VAL:HG21	3:L:637:PHE:CE1	1.97	0.99

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
7:M:101:MYR:H22	3:N:191:LYS:HZ3	1.28	0.99
5:W:645:VAL:O	5:W:647:PRO:HD3	1.60	0.99
1:r:60:ALA:H	3:R:398:SER:HA	0.84	0.99
7:C:101:MYR:H72	3:D:200:CYS:SG	2.03	0.99
7:S:101:MYR:H101	7:S:101:MYR:H143	1.42	0.99
1:t:45:ARG:HG3	3:T:400:ALA:CA	1.88	0.98
3:R:274:ILE:HG21	3:R:601:THR:CG2	1.92	0.98
3:R:545:VAL:HG21	3:T:637:PHE:CE1	1.97	0.98
3:T:579:ASN:HD22	3:T:582:LEU:HG	1.28	0.98
5:W:1038:ASN:HD22	5:W:1049:ALA:HB3	1.26	0.98
1:f:8:GLN:NE2	3:H:302:GLU:HG2	1.75	0.98
1:t:8:GLN:CG	3:P:302:GLU:CG	2.36	0.98
3:H:341:MET:HB3	3:H:431:ILE:HD12	1.46	0.98
3:B:431:ILE:HD13	3:B:437:ILE:HG21	1.43	0.98
1:j:59:HIS:CG	3:J:397:VAL:O	2.17	0.98
3:N:274:ILE:CG2	3:N:601:THR:HG21	1.92	0.98
3:P:579:ASN:HD22	3:P:582:LEU:HG	1.28	0.98
7:S:101:MYR:H72	3:T:200:CYS:SG	2.04	0.98
5:W:1040:VAL:HB	5:W:1041:PRO:HD3	1.43	0.98
7:A:101:MYR:H101	7:A:101:MYR:H143	1.42	0.98
3:B:341:MET:HB3	3:B:431:ILE:CD1	1.94	0.98
3:J:341:MET:HB3	3:J:431:ILE:HD12	1.45	0.98
3:L:341:MET:HB3	3:L:431:ILE:CD1	1.93	0.98
3:R:341:MET:HB3	3:R:431:ILE:CD1	1.93	0.98
3:T:341:MET:HB3	3:T:431:ILE:CD1	1.94	0.98
1:b:59:HIS:HA	3:B:398:SER:HB2	1.45	0.98
1:r:23:THR:CB	1:r:42:THR:OG1	2.10	0.98
1:t:15:ARG:NH1	3:T:406:ASN:HB3	1.77	0.98
3:L:579:ASN:HD22	3:L:582:LEU:HG	1.27	0.98
3:P:274:ILE:HG21	3:P:601:THR:CG2	1.92	0.98
3:R:609:ARG:NH1	2:S:27:THR:HG23	1.78	0.98
5:W:818:PHE:HE2	5:W:989:HIS:HD1	1.05	0.98
6:Y:1027:LEU:H	6:Y:1027:LEU:HD12	1.27	0.98
1:d:45:ARG:HG2	3:D:400:ALA:HA	1.01	0.98
3:D:63:ILE:H	3:D:63:ILE:CD1	1.73	0.98
3:J:579:ASN:HD22	3:J:582:LEU:HG	1.27	0.98
3:R:274:ILE:CG2	3:R:601:THR:HG21	1.92	0.98
6:X:242:ARG:HH21	6:X:1085:LEU:CD1	1.75	0.98
1:l:24:LEU:CD2	1:l:80:ARG:CD	2.41	0.98
1:n:8:GLN:CD	3:J:302:GLU:HG2	1.88	0.98
1:r:60:ALA:H	3:R:398:SER:CA	1.74	0.98

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:B:274:ILE:CG2	3:B:601:THR:HG21	1.92	0.98
7:E:101:MYR:H101	7:E:101:MYR:H143	1.42	0.98
3:H:341:MET:HB3	3:H:431:ILE:CD1	1.94	0.98
3:N:341:MET:HB3	3:N:431:ILE:CD1	1.94	0.98
3:P:341:MET:HB3	3:P:431:ILE:CD1	1.94	0.98
3:T:274:ILE:CG2	3:T:601:THR:HG21	1.92	0.98
6:Y:432:ILE:HG21	6:Y:1204:LEU:HB3	1.44	0.98
7:A:101:MYR:H72	3:B:200:CYS:SG	2.03	0.98
3:H:380:VAL:CG1	3:H:384:GLN:HG2	1.94	0.98
3:J:341:MET:HB3	3:J:431:ILE:CD1	1.94	0.98
3:J:609:ARG:NH1	2:K:27:THR:HG23	1.78	0.98
7:M:101:MYR:H72	3:N:200:CYS:SG	2.04	0.98
6:Y:295:SER:HB2	6:Y:890:GLN:NE2	1.79	0.98
1:d:45:ARG:CG	3:D:401:GLY:H	1.76	0.98
1:h:22:LEU:CD1	1:h:83:VAL:HG21	1.94	0.98
1:t:59:HIS:HD2	3:T:397:VAL:O	1.36	0.98
3:D:274:ILE:CG2	3:D:601:THR:HG21	1.92	0.98
7:O:101:MYR:H72	3:P:200:CYS:SG	2.03	0.98
6:X:464:SER:CB	6:Y:500:ILE:CG2	2.40	0.98
1:f:58:PRO:CB	3:F:423:GLN:NE2	2.27	0.97
1:n:24:LEU:HD21	1:n:80:ARG:HD2	1.46	0.97
1:r:45:ARG:HA	3:R:399:ALA:O	1.64	0.97
3:D:380:VAL:CG1	3:D:384:GLN:HG2	1.94	0.97
3:F:380:VAL:CG1	3:F:384:GLN:HG2	1.94	0.97
4:U:27:GLN:CG	6:X:880:LEU:HB3	1.93	0.97
1:n:6:ILE:HD11	3:J:586:VAL:HG21	0.98	0.97
3:L:431:ILE:HD13	3:L:437:ILE:HG21	1.43	0.97
3:R:341:MET:HB3	3:R:431:ILE:HD12	1.45	0.97
3:T:380:VAL:CG1	3:T:384:GLN:HG2	1.94	0.97
1:n:58:PRO:HG3	3:N:423:GLN:HE21	1.29	0.97
3:B:341:MET:HB3	3:B:431:ILE:HD12	1.46	0.97
3:D:341:MET:HB3	3:D:431:ILE:HD12	1.45	0.97
3:D:579:ASN:HD22	3:D:582:LEU:HG	1.27	0.97
3:J:380:VAL:CG1	3:J:384:GLN:HG2	1.94	0.97
3:L:274:ILE:CG2	3:L:601:THR:HG21	1.92	0.97
7:Q:101:MYR:H22	3:R:191:LYS:NZ	1.77	0.97
3:T:341:MET:HB3	3:T:431:ILE:HD12	1.46	0.97
5:W:696:PRO:HG3	5:W:706:SER:HB2	1.46	0.97
2:C:27:THR:HG22	2:C:27:THR:O	1.64	0.97
3:N:380:VAL:CG1	3:N:384:GLN:HG2	1.94	0.97
2:O:27:THR:HG23	3:T:609:ARG:NH1	1.78	0.97

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:T:274:ILE:HG21	3:T:601:THR:CG2	1.92	0.97
5:W:584:GLN:HG2	5:W:621:ASN:HB2	1.47	0.97
3:D:341:MET:HB3	3:D:431:ILE:CD1	1.94	0.97
3:D:609:ARG:NH1	2:E:27:THR:HG23	1.78	0.97
7:I:101:MYR:H101	7:I:101:MYR:H143	1.42	0.97
7:I:101:MYR:H72	3:J:200:CYS:SG	2.03	0.97
3:P:380:VAL:CG1	3:P:384:GLN:HG2	1.94	0.97
6:X:606:MET:HE2	6:Y:718:THR:CG2	1.92	0.97
1:l:15:ARG:HH11	3:L:406:ASN:CB	1.76	0.97
1:f:45:ARG:CG	3:F:401:GLY:H	1.76	0.97
1:p:58:PRO:HG3	3:P:423:GLN:CD	1.89	0.97
3:L:609:ARG:NH1	2:M:27:THR:HG23	1.78	0.97
3:N:274:ILE:HG21	3:N:601:THR:CG2	1.92	0.97
6:Y:1150:ASN:HD22	6:Y:1156:THR:HB	1.28	0.97
1:h:24:LEU:HD22	1:h:41:VAL:HG22	1.46	0.97
3:B:369:SER:HG	3:B:443:PHE:HD1	1.12	0.97
7:I:101:MYR:C14	7:I:101:MYR:C10	2.42	0.97
1:b:73:ARG:NH1	1:b:74:GLN:OE1	1.97	0.97
1:j:24:LEU:CD1	1:j:41:VAL:CG2	2.41	0.97
3:B:274:ILE:HG21	3:B:601:THR:CG2	1.92	0.97
3:H:431:ILE:HD13	3:H:437:ILE:HG21	1.43	0.97
3:L:341:MET:HB3	3:L:431:ILE:HD12	1.45	0.97
1:d:23:THR:CG2	3:D:323:SER:HA	1.95	0.97
7:K:101:MYR:H22	3:L:191:LYS:NZ	1.77	0.97
6:X:606:MET:HE2	6:Y:718:THR:HG23	0.99	0.97
3:B:380:VAL:CG1	3:B:384:GLN:HG2	1.94	0.97
3:P:609:ARG:NH1	2:Q:27:THR:HG23	1.78	0.97
1:b:45:ARG:CD	3:B:400:ALA:HA	1.93	0.96
3:T:431:ILE:HD13	3:T:437:ILE:HG21	1.43	0.96
6:X:1012:ARG:HH21	6:X:1043:GLY:HA3	1.29	0.96
1:f:8:GLN:CG	3:H:302:GLU:CG	2.37	0.96
1:t:45:ARG:O	1:t:59:HIS:CA	2.13	0.96
3:L:380:VAL:CG1	3:L:384:GLN:HG2	1.94	0.96
3:D:50:PRO:HG3	3:D:56:ALA:HB2	1.47	0.96
3:H:297:ALA:HB2	3:H:460:MET:HB2	1.47	0.96
6:Y:410:HIS:HD2	6:Y:411:GLN:N	1.62	0.96
1:l:23:THR:HG21	1:l:25:TYR:CE2	1.98	0.96
1:f:8:GLN:CD	3:H:302:GLU:HG2	1.89	0.96
2:A:27:THR:HG22	2:A:27:THR:O	1.64	0.96
3:D:63:ILE:HD13	3:D:63:ILE:N	1.77	0.96
3:F:341:MET:HB3	3:F:431:ILE:CD1	1.94	0.96

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:R:380:VAL:CG1	3:R:384:GLN:HG2	1.94	0.96
3:T:297:ALA:HB2	3:T:460:MET:HB2	1.47	0.96
6:Y:705:ILE:HG21	6:Y:737:ARG:HB2	1.47	0.96
3:H:579:ASN:HD22	3:H:582:LEU:HG	1.28	0.96
6:X:464:SER:HB2	6:Y:500:ILE:HG23	1.47	0.96
6:Y:122:CYS:HB2	6:Y:168:ARG:HH22	1.31	0.96
1:b:58:PRO:HG3	3:B:423:GLN:HE21	1.31	0.96
1:j:45:ARG:CA	3:J:399:ALA:O	2.14	0.96
1:p:45:ARG:CG	3:P:401:GLY:H	1.79	0.96
1:p:45:ARG:HG3	3:P:401:GLY:H	1.23	0.96
7:G:101:MYR:H72	3:H:200:CYS:SG	2.04	0.96
3:R:297:ALA:HB2	3:R:460:MET:HB2	1.47	0.96
1:t:8:GLN:CD	3:P:302:GLU:HG2	1.91	0.96
7:G:101:MYR:H22	3:H:191:LYS:NZ	1.81	0.96
1:b:59:HIS:CE1	1:b:61:ASN:HD21	1.83	0.96
3:F:579:ASN:HD22	3:F:582:LEU:HG	1.28	0.96
3:R:418:ASN:HD21	3:T:337:ARG:HH21	1.13	0.96
6:Y:259:LEU:HD11	6:Y:303:GLN:HG2	1.45	0.96
1:l:21:ARG:CD	3:L:327:TYR:OH	2.13	0.96
3:F:341:MET:HB3	3:F:431:ILE:HD12	1.45	0.96
3:J:297:ALA:HB2	3:J:460:MET:HB2	1.47	0.96
7:M:101:MYR:H22	3:N:191:LYS:NZ	1.80	0.96
1:p:24:LEU:CD2	1:p:80:ARG:HD3	1.94	0.96
7:M:101:MYR:C14	7:M:101:MYR:C10	2.42	0.96
3:P:297:ALA:HB2	3:P:460:MET:HB2	1.47	0.96
6:X:368:ILE:HG12	6:X:369:GLY:H	1.29	0.96
1:j:23:THR:CB	3:J:323:SER:HA	1.94	0.95
3:L:297:ALA:HB2	3:L:460:MET:HB2	1.47	0.95
4:U:35:ARG:HD3	6:X:449:GLU:HG2	1.46	0.95
1:l:16:ALA:O	1:l:22:LEU:HD23	1.65	0.95
1:b:15:ARG:CZ	3:B:402:ALA:HB1	1.94	0.95
1:b:45:ARG:CG	3:B:401:GLY:N	2.27	0.95
1:h:8:GLN:CD	3:D:302:GLU:HG2	1.90	0.95
1:j:45:ARG:HG3	3:J:400:ALA:HA	1.39	0.95
7:K:101:MYR:H72	3:L:200:CYS:SG	2.06	0.95
1:l:58:PRO:CG	3:L:423:GLN:HE21	1.79	0.95
3:J:50:PRO:O	4:V:401:ALA:HB2	1.64	0.95
2:K:27:THR:HG22	2:K:27:THR:O	1.64	0.95
3:B:579:ASN:HD22	3:B:582:LEU:HG	1.27	0.95
3:F:297:ALA:HB2	3:F:460:MET:HB2	1.47	0.95
3:P:58:ILE:HG23	3:P:124:GLY:HA2	1.46	0.95

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
6:Y:376:LEU:HD13	6:Y:438:PRO:HB3	1.48	0.95
1:d:59:HIS:HA	3:D:398:SER:HB2	1.49	0.95
1:j:59:HIS:HB2	3:J:397:VAL:O	1.66	0.95
1:p:24:LEU:HD21	1:p:80:ARG:HD3	1.47	0.95
1:l:24:LEU:HD21	1:l:80:ARG:NE	1.81	0.95
1:d:45:ARG:HG3	3:D:400:ALA:CA	1.93	0.95
1:j:45:ARG:HG3	3:J:400:ALA:C	1.90	0.95
1:n:45:ARG:O	1:n:58:PRO:O	1.83	0.95
1:p:15:ARG:CZ	3:P:402:ALA:CB	2.45	0.95
3:D:341:MET:HB2	3:D:437:ILE:HG23	1.49	0.95
7:Q:101:MYR:H72	3:R:200:CYS:SG	2.06	0.95
3:T:103:CYS:HB3	5:W:407:ALA:HB2	1.44	0.95
5:W:840:CYS:CB	5:W:854:MET:SD	2.54	0.95
6:X:606:MET:HE3	6:Y:718:THR:CG2	1.96	0.95
1:h:6:ILE:CD1	3:D:586:VAL:CG2	2.41	0.95
3:N:297:ALA:HB2	3:N:460:MET:HB2	1.47	0.95
2:O:27:THR:HG22	2:O:27:THR:O	1.64	0.95
3:P:55:VAL:HG11	5:W:348:GLN:NE2	1.81	0.95
3:P:341:MET:HB3	3:P:431:ILE:HD12	1.46	0.95
1:f:15:ARG:NH1	3:F:402:ALA:HB1	1.81	0.95
3:P:337:ARG:HH21	3:T:418:ASN:HD21	1.13	0.95
1:d:21:ARG:NH1	3:D:327:TYR:CE1	2.33	0.95
1:n:45:ARG:HG3	3:N:400:ALA:HA	1.44	0.95
1:n:58:PRO:CG	3:N:423:GLN:HE21	1.80	0.95
3:N:341:MET:HB3	3:N:431:ILE:HD12	1.46	0.95
7:Q:101:MYR:C14	7:Q:101:MYR:C10	2.42	0.95
6:X:938:TYR:CZ	6:X:946:VAL:HG21	2.01	0.95
6:Y:1160:GLU:HG2	6:Y:1160:GLU:O	1.67	0.95
1:j:8:GLN:NE2	3:L:302:GLU:CG	2.29	0.95
1:j:59:HIS:HB3	3:J:398:SER:CB	1.81	0.95
1:p:35:PHE:CE2	1:p:73:ARG:HG2	2.01	0.95
3:J:341:MET:HB2	3:J:437:ILE:HG23	1.49	0.95
5:W:358:LEU:HB3	5:W:376:PHE:HB3	1.45	0.95
1:l:17:ALA:O	1:l:82:LEU:HD12	1.66	0.94
1:r:6:ILE:HD11	3:T:586:VAL:HG21	0.95	0.94
3:N:92:TRP:CE3	3:N:210:VAL:HG11	2.03	0.94
2:Q:27:THR:HG22	2:Q:27:THR:O	1.64	0.94
1:b:23:THR:CG2	1:b:25:TYR:CE2	2.49	0.94
1:j:8:GLN:HE21	3:L:302:GLU:HG2	1.27	0.94
1:n:45:ARG:CA	3:N:399:ALA:O	2.15	0.94
3:R:579:ASN:HD22	3:R:582:LEU:HG	1.27	0.94

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:b:59:HIS:CE1	1:b:61:ASN:ND2	2.34	0.94
1:n:15:ARG:NH1	3:N:402:ALA:CB	2.28	0.94
4:U:245:GLN:HE21	4:U:246:ALA:H	1.10	0.94
1:l:15:ARG:HH11	3:L:406:ASN:HB3	1.30	0.94
1:h:21:ARG:NH1	3:H:327:TYR:OH	2.01	0.94
1:j:45:ARG:HG3	3:J:401:GLY:H	1.09	0.94
1:t:61:ASN:CG	3:T:395:THR:CG2	2.39	0.94
3:H:92:TRP:CE3	3:H:210:VAL:HG11	2.02	0.94
7:I:101:MYR:H22	3:J:191:LYS:NZ	1.82	0.94
3:P:92:TRP:CE3	3:P:210:VAL:HG11	2.03	0.94
3:P:341:MET:HB2	3:P:437:ILE:HG23	1.49	0.94
4:U:26:LEU:HA	4:U:71:LEU:HD11	1.49	0.94
6:X:195:ILE:HG22	6:X:195:ILE:O	1.67	0.94
1:f:5:MET:HE2	1:f:5:MET:HA	1.49	0.94
1:r:59:HIS:HB3	3:R:423:GLN:O	1.66	0.94
7:C:101:MYR:H22	3:D:191:LYS:NZ	1.82	0.94
2:E:27:THR:HG22	2:E:27:THR:O	1.64	0.94
7:O:101:MYR:H22	3:P:191:LYS:NZ	1.82	0.94
3:R:431:ILE:HD13	3:R:437:ILE:HG21	1.43	0.94
2:S:27:THR:HG22	2:S:27:THR:O	1.64	0.94
6:Y:122:CYS:HB2	6:Y:168:ARG:NH2	1.81	0.94
1:d:44:GLY:O	3:D:399:ALA:N	2.00	0.94
1:j:45:ARG:HG2	3:J:400:ALA:HA	0.95	0.94
1:n:50:ALA:O	1:n:74:GLN:OE1	1.83	0.94
1:n:58:PRO:HG3	3:N:423:GLN:NE2	1.82	0.94
7:E:101:MYR:H21	3:F:191:LYS:HE2	1.49	0.94
3:L:178:SER:CB	2:Q:3:ASN:ND2	2.30	0.94
1:d:57:ALA:CB	1:d:59:HIS:CD2	2.49	0.94
3:B:297:ALA:HB2	3:B:460:MET:HB2	1.47	0.94
4:U:198:GLN:HE22	6:X:661:THR:CG2	1.79	0.94
1:l:6:ILE:HD11	3:N:586:VAL:HG21	0.96	0.94
1:b:59:HIS:C	3:B:397:VAL:O	2.10	0.94
1:f:6:ILE:HD11	3:H:586:VAL:HG21	0.94	0.94
1:d:23:THR:CG2	3:D:323:SER:CA	2.44	0.94
1:h:15:ARG:CZ	3:H:402:ALA:CB	2.46	0.94
1:j:45:ARG:HG2	3:J:400:ALA:CA	1.90	0.94
1:n:15:ARG:NH1	3:N:406:ASN:HB3	1.81	0.94
1:r:60:ALA:O	3:R:397:VAL:C	2.11	0.94
3:F:341:MET:HB2	3:F:437:ILE:HG23	1.49	0.94
2:I:27:THR:HG22	2:I:27:THR:O	1.64	0.94
3:J:369:SER:HG	3:J:443:PHE:HD1	1.11	0.94

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:N:341:MET:HB2	3:N:437:ILE:HG23	1.49	0.94
3:P:418:ASN:HD21	3:R:337:ARG:HH21	1.13	0.94
1:l:15:ARG:NH1	3:L:402:ALA:CB	2.31	0.94
1:p:59:HIS:CB	3:P:398:SER:HB2	1.92	0.94
7:E:101:MYR:H72	3:F:200:CYS:SG	2.06	0.94
5:W:1138:MET:HE3	5:W:1225:ASP:HB2	1.48	0.94
3:J:92:TRP:CE3	3:J:210:VAL:HG11	2.03	0.94
2:M:27:THR:HG22	2:M:27:THR:O	1.64	0.94
1:b:23:THR:HG22	1:b:42:THR:HG1	1.24	0.93
7:A:101:MYR:H22	3:B:191:LYS:NZ	1.82	0.93
3:B:92:TRP:CE3	3:B:210:VAL:HG11	2.03	0.93
2:G:27:THR:HG22	2:G:27:THR:O	1.64	0.93
5:W:162:SER:HA	5:W:171:THR:HG22	1.50	0.93
1:l:16:ALA:O	1:l:22:LEU:CD2	2.17	0.93
1:j:44:GLY:O	3:J:399:ALA:N	2.00	0.93
1:t:6:ILE:CD1	3:P:586:VAL:CG2	2.42	0.93
3:H:341:MET:HB2	3:H:437:ILE:HG23	1.49	0.93
3:J:549:LEU:HB3	3:L:644:LEU:HD11	1.51	0.93
3:P:58:ILE:CG2	3:P:124:GLY:HA2	1.98	0.93
3:R:628:GLN:HE21	5:W:870:ALA:HA	0.77	0.93
7:S:101:MYR:H22	3:T:191:LYS:HZ3	1.28	0.93
5:W:1037:VAL:HG12	5:W:1039:PRO:HD2	1.47	0.93
1:h:45:ARG:CG	3:H:401:GLY:H	1.80	0.93
3:D:237:GLU:HB2	3:F:46:LYS:NZ	1.83	0.93
3:F:92:TRP:CE3	3:F:210:VAL:HG11	2.03	0.93
3:T:92:TRP:CE3	3:T:210:VAL:HG11	2.02	0.93
4:U:30:ALA:HA	4:U:36:PRO:HB3	1.50	0.93
1:h:22:LEU:CG	1:h:83:VAL:HG21	1.97	0.93
1:n:15:ARG:NH2	3:N:402:ALA:CB	2.32	0.93
1:r:45:ARG:CG	3:R:400:ALA:CA	2.45	0.93
7:S:101:MYR:C14	7:S:101:MYR:C10	2.42	0.93
7:S:101:MYR:H22	3:T:191:LYS:NZ	1.80	0.93
1:j:22:LEU:HD21	1:j:41:VAL:HG13	1.51	0.93
1:h:8:GLN:CG	3:D:302:GLU:CG	2.38	0.93
3:D:46:LYS:NZ	3:H:237:GLU:HB2	1.84	0.93
3:L:92:TRP:CE3	3:L:210:VAL:HG11	2.03	0.93
7:I:101:MYR:H22	3:J:191:LYS:HZ3	1.33	0.93
3:J:418:ASN:HD21	3:L:337:ARG:HH21	1.13	0.93
3:P:369:SER:HG	3:P:443:PHE:HD1	1.13	0.93
7:Q:101:MYR:H21	3:R:191:LYS:HE2	1.49	0.93
3:R:92:TRP:CE3	3:R:210:VAL:HG11	2.03	0.93

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:l:8:GLN:CD	3:N:302:GLU:HG2	1.94	0.93
1:h:24:LEU:CD2	1:h:41:VAL:HG22	1.98	0.93
1:n:59:HIS:HA	3:N:398:SER:CB	1.98	0.93
3:J:46:LYS:NZ	3:N:237:GLU:HB2	1.84	0.93
2:K:9:ASN:ND2	3:L:206:GLU:HA	1.84	0.93
2:Q:9:ASN:ND2	3:R:206:GLU:HA	1.84	0.93
6:Y:355:LYS:NZ	6:Y:1156:THR:CG2	2.30	0.93
1:b:16:ALA:C	1:b:22:LEU:HD21	1.93	0.93
1:d:22:LEU:HD21	1:d:83:VAL:HG22	1.48	0.93
3:D:92:TRP:CE3	3:D:210:VAL:HG11	2.03	0.93
2:I:9:ASN:ND2	3:J:206:GLU:HA	1.84	0.93
3:T:341:MET:HB2	3:T:437:ILE:HG23	1.49	0.93
6:Y:196:GLU:OE1	6:Y:313:LYS:HE2	1.67	0.93
1:r:8:GLN:CG	3:T:302:GLU:CG	2.42	0.93
3:L:237:GLU:HB2	3:N:46:LYS:NZ	1.84	0.93
3:L:341:MET:CB	3:L:431:ILE:CD1	2.47	0.93
3:N:341:MET:CB	3:N:431:ILE:CD1	2.47	0.93
3:P:530:ARG:HE	3:T:472:ARG:NH2	1.67	0.93
3:P:549:LEU:HB3	3:R:644:LEU:HD11	1.51	0.93
5:W:1039:PRO:HG2	5:W:1042:THR:HG22	1.49	0.93
6:X:1167:LEU:HD22	6:X:1187:ARG:HH12	1.29	0.93
6:X:1170:PRO:HD2	6:X:1189:THR:HG23	1.46	0.93
3:F:341:MET:CB	3:F:431:ILE:CD1	2.47	0.92
3:R:341:MET:HB2	3:R:437:ILE:HG23	1.49	0.92
1:d:45:ARG:CA	3:D:399:ALA:O	2.17	0.92
1:n:23:THR:HB	3:N:323:SER:CA	2.00	0.92
1:r:8:GLN:CD	3:T:302:GLU:HG2	1.93	0.92
3:B:341:MET:HB2	3:B:437:ILE:HG23	1.49	0.92
3:D:337:ARG:HH21	3:H:418:ASN:HD21	1.13	0.92
7:G:101:MYR:C14	7:G:101:MYR:C10	2.42	0.92
3:H:341:MET:CB	3:H:431:ILE:CD1	2.47	0.92
3:J:58:ILE:HG23	3:J:124:GLY:HA2	1.50	0.92
3:J:237:GLU:HB2	3:L:46:LYS:NZ	1.83	0.92
3:R:237:GLU:HB2	3:T:46:LYS:NZ	1.84	0.92
1:f:59:HIS:HB2	3:F:397:VAL:O	1.69	0.92
3:L:341:MET:HB2	3:L:437:ILE:HG23	1.49	0.92
3:R:341:MET:CB	3:R:431:ILE:CD1	2.47	0.92
1:p:45:ARG:HG3	3:P:400:ALA:HA	1.28	0.92
1:t:21:ARG:O	3:T:323:SER:OG	1.85	0.92
7:A:101:MYR:C14	7:A:101:MYR:C10	2.42	0.92
2:E:9:ASN:ND2	3:F:206:GLU:HA	1.84	0.92

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
7:O:101:MYR:C14	7:O:101:MYR:C10	2.42	0.92
3:P:341:MET:CB	3:P:431:ILE:CD1	2.47	0.92
6:Y:209:ASN:HD22	6:Y:242:ARG:HG3	1.33	0.92
1:b:50:ALA:O	1:b:74:GLN:CB	2.14	0.92
3:J:49:ARG:HH12	3:J:115:THR:HG22	1.30	0.92
3:J:337:ARG:HH21	3:N:418:ASN:HD21	1.13	0.92
1:l:44:GLY:O	3:L:399:ALA:N	2.02	0.92
1:h:44:GLY:O	3:H:399:ALA:N	2.02	0.92
1:n:45:ARG:HG3	3:N:401:GLY:H	1.30	0.92
1:p:15:ARG:NH1	3:P:402:ALA:HB1	1.83	0.92
3:J:49:ARG:HH12	3:J:115:THR:CG2	1.81	0.92
3:T:101:THR:HA	5:W:406:ASN:HB2	1.48	0.92
4:U:182:ILE:HG21	4:U:279:VAL:HG23	1.50	0.92
1:l:6:ILE:CD1	3:N:586:VAL:CG2	2.44	0.92
1:h:15:ARG:HH11	3:H:406:ASN:HB3	1.34	0.92
3:B:341:MET:CB	3:B:431:ILE:CD1	2.47	0.92
3:R:346:ASP:HB3	3:R:428:ARG:H	1.35	0.92
3:T:341:MET:CB	3:T:431:ILE:CD1	2.47	0.92
6:X:1078:LEU:HD23	6:X:1078:LEU:H	1.35	0.92
3:J:428:ARG:HH11	3:J:428:ARG:HG2	1.35	0.92
3:L:418:ASN:HD21	3:N:337:ARG:HH21	1.13	0.92
3:L:428:ARG:HH11	3:L:428:ARG:HG2	1.35	0.92
3:P:46:LYS:NZ	3:T:237:GLU:HB2	1.84	0.92
3:P:644:LEU:HD11	3:T:549:LEU:HB3	1.51	0.92
4:U:93:ARG:HH11	4:U:93:ARG:HB3	1.34	0.92
1:f:58:PRO:HB2	3:F:423:GLN:CG	2.00	0.92
1:p:58:PRO:O	1:p:59:HIS:CG	2.23	0.92
1:t:45:ARG:CG	3:T:401:GLY:H	1.83	0.92
3:D:549:LEU:HB3	3:F:644:LEU:HD11	1.50	0.92
3:F:237:GLU:HB2	3:H:46:LYS:NZ	1.83	0.92
3:F:428:ARG:HH11	3:F:428:ARG:HG2	1.35	0.92
3:P:237:GLU:HB2	3:R:46:LYS:NZ	1.84	0.92
1:f:22:LEU:HG	1:f:83:VAL:HG21	1.51	0.92
3:D:341:MET:CB	3:D:431:ILE:CD1	2.47	0.92
3:J:341:MET:CB	3:J:431:ILE:CD1	2.47	0.92
6:X:594:ALA:HA	6:X:608:GLN:HE22	1.33	0.92
6:X:656:PHE:HE1	6:X:675:MET:HB2	1.35	0.92
1:d:58:PRO:HG3	3:D:423:GLN:NE2	1.85	0.91
1:p:58:PRO:HG3	3:P:423:GLN:CG	2.00	0.91
3:D:644:LEU:HD11	3:H:549:LEU:HB3	1.51	0.91
6:X:437:ARG:NH2	6:X:1207:ILE:HA	1.85	0.91

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:h:45:ARG:HG3	3:H:400:ALA:HA	1.50	0.91
1:r:23:THR:HG1	3:R:323:SER:HB3	1.12	0.91
3:H:346:ASP:HB3	3:H:428:ARG:H	1.35	0.91
7:Q:101:MYR:H22	3:R:191:LYS:HZ3	1.29	0.91
1:p:59:HIS:HD2	3:P:398:SER:HB2	1.32	0.91
1:r:5:MET:HA	1:r:5:MET:CE	1.99	0.91
2:A:9:ASN:ND2	3:B:206:GLU:HA	1.84	0.91
3:D:297:ALA:HB2	3:D:460:MET:HB2	1.47	0.91
7:K:101:MYR:H21	3:L:191:LYS:HE2	1.49	0.91
1:l:45:ARG:CG	3:L:401:GLY:H	1.83	0.91
1:f:58:PRO:HB3	3:F:423:GLN:NE2	1.84	0.91
1:t:22:LEU:HD11	1:t:83:VAL:HG21	0.92	0.91
3:T:271:LEU:HG	3:T:517:LEU:HB2	1.52	0.91
5:W:839:GLU:HB3	5:W:861:ALA:HB3	1.50	0.91
1:l:58:PRO:HG3	3:L:423:GLN:HE21	1.32	0.91
1:h:45:ARG:O	1:h:59:HIS:O	1.89	0.91
3:D:346:ASP:HB3	3:D:428:ARG:H	1.35	0.91
2:O:9:ASN:ND2	3:P:206:GLU:HA	1.84	0.91
1:n:45:ARG:HG3	3:N:400:ALA:CA	1.93	0.91
2:C:9:ASN:ND2	3:D:206:GLU:HA	1.84	0.91
3:D:271:LEU:HG	3:D:517:LEU:HB2	1.53	0.91
3:J:644:LEU:HD11	3:N:549:LEU:HB3	1.51	0.91
3:T:103:CYS:HB3	5:W:407:ALA:CB	2.01	0.91
1:l:21:ARG:NH1	3:L:370:VAL:HG13	1.86	0.91
1:b:15:ARG:NH1	3:B:402:ALA:CB	2.32	0.91
1:p:45:ARG:CD	3:P:400:ALA:HA	2.01	0.91
1:r:6:ILE:CD1	3:T:586:VAL:CG2	2.43	0.91
1:t:45:ARG:HG3	3:T:401:GLY:H	1.28	0.91
3:L:271:LEU:HG	3:L:517:LEU:HB2	1.53	0.91
3:R:428:ARG:HH11	3:R:428:ARG:HG2	1.35	0.91
6:Y:122:CYS:SG	6:Y:140:HIS:HE1	1.92	0.91
1:l:58:PRO:CB	3:L:423:GLN:HE21	1.83	0.91
1:n:23:THR:CB	3:N:323:SER:CA	2.49	0.91
1:t:5:MET:HE2	1:t:5:MET:HA	1.51	0.91
5:W:436:GLY:HA3	5:W:1009:ASN:HA	1.53	0.91
1:l:24:LEU:CD1	1:l:33:THR:HA	2.01	0.91
1:l:45:ARG:HG3	3:L:400:ALA:CA	2.01	0.91
1:h:15:ARG:HH11	3:H:406:ASN:CG	1.77	0.91
1:h:58:PRO:CB	3:H:423:GLN:NE2	2.31	0.91
1:h:59:HIS:HE1	1:h:60:ALA:O	1.54	0.91
1:p:62:VAL:HG12	1:p:65:ILE:HD12	1.51	0.91

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:R:271:LEU:HG	3:R:517:LEU:HB2	1.53	0.91
3:B:346:ASP:HB3	3:B:428:ARG:H	1.35	0.90
3:H:271:LEU:HG	3:H:517:LEU:HB2	1.52	0.90
3:J:346:ASP:HB3	3:J:428:ARG:H	1.35	0.90
3:P:428:ARG:HH11	3:P:428:ARG:HG2	1.35	0.90
5:W:1154:ARG:HD2	5:W:1189:ASN:HD22	1.36	0.90
1:f:15:ARG:NH1	3:F:406:ASN:CG	2.27	0.90
1:d:58:PRO:CG	3:D:423:GLN:HE21	1.84	0.90
1:p:58:PRO:CG	3:P:423:GLN:HG2	2.01	0.90
1:t:24:LEU:HD21	1:t:35:PHE:CD1	2.05	0.90
1:t:45:ARG:HG2	3:T:400:ALA:CA	1.95	0.90
3:B:428:ARG:HH11	3:B:428:ARG:HG2	1.35	0.90
3:T:346:ASP:HB3	3:T:428:ARG:H	1.35	0.90
1:h:59:HIS:HB2	3:H:398:SER:HB2	0.91	0.90
1:j:61:ASN:CG	3:J:395:THR:HG21	1.96	0.90
3:F:549:LEU:HB3	3:H:644:LEU:HD11	1.51	0.90
3:N:346:ASP:HB3	3:N:428:ARG:H	1.35	0.90
1:l:53:LEU:HD13	1:l:74:GLN:OE1	1.71	0.90
1:p:21:ARG:HH12	3:P:327:TYR:HE1	1.19	0.90
7:E:101:MYR:C14	7:E:101:MYR:C10	2.42	0.90
3:F:346:ASP:HB3	3:F:428:ARG:H	1.35	0.90
3:L:346:ASP:HB3	3:L:428:ARG:H	1.35	0.90
3:L:549:LEU:HB3	3:N:644:LEU:HD11	1.51	0.90
1:b:22:LEU:HG	1:b:83:VAL:CG2	2.00	0.90
7:A:101:MYR:C7	3:B:200:CYS:SG	2.60	0.90
3:R:472:ARG:HH21	3:T:530:ARG:HE	0.93	0.90
3:B:450:SER:HA	3:B:453:ASN:HD22	1.37	0.90
7:C:101:MYR:C14	7:C:101:MYR:C10	2.42	0.90
3:T:450:SER:HA	3:T:453:ASN:HD22	1.37	0.90
3:F:418:ASN:HD21	3:H:337:ARG:HH21	1.13	0.90
3:J:121:ALA:HB2	3:N:87:PHE:CE1	2.07	0.90
3:N:428:ARG:HH11	3:N:428:ARG:HG2	1.35	0.90
5:W:740:VAL:HG21	5:W:770:ARG:CZ	2.02	0.90
1:p:44:GLY:CA	3:P:322:TYR:OH	2.20	0.90
1:t:45:ARG:CD	3:T:400:ALA:HA	2.01	0.90
5:W:1238:ARG:HG2	5:W:1288:THR:HG22	1.54	0.90
6:X:416:ILE:HG22	6:X:445:PHE:CZ	2.07	0.90
1:n:45:ARG:CD	3:N:400:ALA:HA	2.01	0.90
1:t:6:ILE:HD11	3:P:586:VAL:HG21	0.91	0.90
3:D:238:VAL:HA	3:F:142:ILE:CD1	2.00	0.90
5:W:167:ASP:HB2	5:W:212:ALA:HB1	1.52	0.90

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:p:45:ARG:CG	3:P:401:GLY:N	2.35	0.90
7:Q:101:MYR:H31	3:R:191:LYS:HG3	1.54	0.90
7:S:101:MYR:H21	3:T:191:LYS:HE2	1.54	0.90
4:U:198:GLN:NE2	6:X:661:THR:HG22	1.87	0.90
6:X:1066:MET:HB3	6:X:1072:THR:HG21	1.54	0.90
1:n:59:HIS:CB	3:N:398:SER:HB2	2.01	0.89
1:p:44:GLY:O	3:P:399:ALA:CA	2.20	0.89
3:D:87:PHE:CE1	3:F:121:ALA:HB2	2.07	0.89
3:L:450:SER:HA	3:L:453:ASN:HD22	1.37	0.89
7:O:101:MYR:C7	3:P:200:CYS:SG	2.60	0.89
6:X:416:ILE:HG22	6:X:445:PHE:CE2	2.08	0.89
7:K:101:MYR:C14	7:K:101:MYR:C10	2.42	0.89
3:P:346:ASP:HB3	3:P:428:ARG:H	1.35	0.89
3:R:549:LEU:HB3	3:T:644:LEU:HD11	1.51	0.89
6:Y:40:ASN:HB3	6:Y:66:SER:HB3	1.54	0.89
6:Y:996:MET:HA	6:Y:996:MET:CE	2.02	0.89
1:d:15:ARG:NE	3:D:406:ASN:ND2	2.19	0.89
3:J:271:LEU:HG	3:J:517:LEU:HB2	1.53	0.89
3:L:87:PHE:CE1	3:N:121:ALA:HB2	2.07	0.89
7:M:101:MYR:H21	3:N:191:LYS:HE2	1.54	0.89
1:d:23:THR:HG21	3:D:323:SER:C	1.94	0.89
3:D:428:ARG:HH11	3:D:428:ARG:HG2	1.35	0.89
3:H:428:ARG:HH11	3:H:428:ARG:HG2	1.35	0.89
2:I:27:THR:HG23	3:N:609:ARG:HH11	1.37	0.89
3:N:271:LEU:HG	3:N:517:LEU:HB2	1.52	0.89
3:P:412:ARG:HH22	3:R:356:THR:CG2	1.86	0.89
1:d:57:ALA:CB	1:d:59:HIS:NE2	2.35	0.89
1:p:47:THR:HG23	1:p:59:HIS:O	1.72	0.89
3:D:152:ARG:HA	3:D:152:ARG:HH11	1.35	0.89
3:H:450:SER:HA	3:H:453:ASN:HD22	1.37	0.89
7:K:101:MYR:H31	3:L:191:LYS:HG3	1.54	0.89
3:R:609:ARG:HH11	2:S:27:THR:HG23	1.37	0.89
5:W:687:ALA:HB1	5:W:996:ASP:HA	1.54	0.89
6:Y:117:TYR:HB3	6:Y:132:LEU:HD22	1.51	0.89
6:Y:979:ARG:HG3	6:Y:1140:GLU:HG3	1.53	0.89
3:J:87:PHE:CE1	3:L:121:ALA:HB2	2.07	0.89
1:p:15:ARG:NH2	3:P:402:ALA:CB	2.36	0.89
3:B:271:LEU:HG	3:B:517:LEU:HB2	1.53	0.89
3:P:60:SER:O	5:W:379:VAL:HG21	1.72	0.89
6:X:786:ALA:HB2	6:X:804:HIS:CD2	2.08	0.89
1:l:17:ALA:O	1:l:82:LEU:CD1	2.21	0.89

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:b:44:GLY:HA3	3:B:322:TYR:CZ	2.06	0.89
1:h:45:ARG:HG3	3:H:400:ALA:CA	2.01	0.89
7:I:101:MYR:C7	3:J:200:CYS:SG	2.60	0.89
6:X:453:LEU:HD23	6:X:861:ARG:HA	1.54	0.89
6:Y:988:ALA:HA	6:Y:1133:VAL:HG12	1.55	0.89
6:Y:1170:PRO:HD2	6:Y:1189:THR:HG23	1.52	0.89
3:F:87:PHE:CE1	3:H:121:ALA:HB2	2.07	0.89
3:F:271:LEU:HG	3:F:517:LEU:HB2	1.53	0.89
3:R:87:PHE:CE1	3:T:121:ALA:HB2	2.07	0.89
3:R:412:ARG:HH22	3:T:356:THR:CG2	1.86	0.89
3:R:450:SER:HA	3:R:453:ASN:HD22	1.37	0.89
5:W:818:PHE:HE2	5:W:989:HIS:ND1	1.71	0.89
6:X:1130:ALA:HB2	6:Y:104:GLN:CB	2.02	0.89
6:Y:902:MET:CG	6:Y:924:ALA:HB2	2.02	0.89
1:l:60:ALA:CB	3:L:322:TYR:OH	2.21	0.89
3:P:530:ARG:NE	3:T:472:ARG:HH21	1.71	0.89
5:W:840:CYS:SG	5:W:854:MET:HG3	2.13	0.89
6:Y:198:THR:HG22	6:Y:313:LYS:HB3	1.53	0.89
6:Y:743:THR:HG22	6:Y:825:GLY:HA3	1.55	0.89
1:b:15:ARG:NH2	3:B:402:ALA:HB3	1.84	0.88
7:C:101:MYR:C7	3:D:200:CYS:SG	2.60	0.88
3:D:356:THR:CG2	3:H:412:ARG:HH22	1.86	0.88
3:D:450:SER:HA	3:D:453:ASN:HD22	1.37	0.88
3:L:412:ARG:HH22	3:N:356:THR:CG2	1.86	0.88
3:P:87:PHE:CE1	3:R:121:ALA:HB2	2.07	0.88
3:P:161:LYS:HZ1	3:T:531:THR:HB	1.34	0.88
3:T:428:ARG:HH11	3:T:428:ARG:HG2	1.35	0.88
5:W:114:ASN:HB3	5:W:117:VAL:HG12	1.52	0.88
6:X:237:GLU:HA	6:X:1151:ALA:CB	2.03	0.88
1:d:45:ARG:HG3	3:D:400:ALA:C	1.98	0.88
1:h:15:ARG:NH1	3:H:406:ASN:HB3	1.88	0.88
1:p:58:PRO:HG2	3:P:423:GLN:CG	2.04	0.88
1:r:4:HIS:CE1	1:r:54:THR:HG22	2.09	0.88
1:r:16:ALA:O	1:r:22:LEU:HD21	1.69	0.88
3:P:356:THR:CG2	3:T:412:ARG:HH22	1.86	0.88
1:j:22:LEU:CD2	1:j:42:THR:O	2.19	0.88
3:D:418:ASN:HD21	3:F:337:ARG:HH21	1.13	0.88
3:F:412:ARG:HH22	3:H:356:THR:CG2	1.86	0.88
3:F:450:SER:HA	3:F:453:ASN:HD22	1.37	0.88
6:X:1012:ARG:HD3	6:X:1012:ARG:N	1.87	0.88
1:n:23:THR:HB	3:N:323:SER:HA	1.56	0.88

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:n:45:ARG:CG	3:N:401:GLY:H	1.85	0.88
3:R:548:LYS:HG3	3:T:645:TRP:CZ2	2.08	0.88
6:X:206:LEU:HD22	6:X:206:LEU:H	1.36	0.88
1:d:8:GLN:NE2	3:F:303:ASP:OD1	2.06	0.88
1:d:15:ARG:NH1	3:D:406:ASN:CB	2.32	0.88
1:r:23:THR:HB	1:r:42:THR:HG1	1.35	0.88
1:t:24:LEU:HD12	1:t:41:VAL:HG22	1.52	0.88
3:R:472:ARG:HH21	3:T:530:ARG:NE	1.70	0.88
5:W:819:ILE:HD11	5:W:896:ILE:HD12	1.54	0.88
5:W:1139:PRO:HG2	5:W:1144:LEU:HD12	1.52	0.88
6:X:801:ARG:HB2	6:X:801:ARG:NH1	1.88	0.88
1:l:15:ARG:NH1	3:L:406:ASN:HB3	1.87	0.88
3:D:548:LYS:HG3	3:F:645:TRP:CZ2	2.08	0.88
7:S:101:MYR:C7	3:T:200:CYS:SG	2.62	0.88
5:W:1074:THR:CG2	5:W:1075:PRO:HD2	2.03	0.88
6:Y:902:MET:HG2	6:Y:924:ALA:HB2	1.54	0.88
1:f:5:MET:O	1:f:6:ILE:C	2.16	0.88
1:h:6:ILE:HD11	3:D:586:VAL:HG21	0.90	0.88
1:h:22:LEU:HG	1:h:83:VAL:HG21	1.54	0.88
1:n:58:PRO:CB	3:N:423:GLN:HE21	1.85	0.88
7:A:101:MYR:H21	3:B:191:LYS:HE2	1.55	0.88
3:D:53:THR:HG22	3:D:54:SER:N	1.86	0.88
3:J:645:TRP:CZ2	3:N:548:LYS:HG3	2.08	0.88
3:P:121:ALA:HB2	3:T:87:PHE:CE1	2.07	0.88
3:P:548:LYS:HG3	3:R:645:TRP:CZ2	2.08	0.88
1:r:60:ALA:O	3:R:397:VAL:O	1.92	0.88
1:t:62:VAL:O	1:t:64:THR:N	2.07	0.88
3:D:412:ARG:HH22	3:F:356:THR:CG2	1.86	0.88
3:F:435:PRO:CG	3:N:436:TYR:OH	2.21	0.88
3:J:548:LYS:HG3	3:L:645:TRP:CZ2	2.09	0.88
3:N:450:SER:HA	3:N:453:ASN:HD22	1.37	0.88
3:P:271:LEU:HG	3:P:517:LEU:HB2	1.53	0.88
3:P:645:TRP:CZ2	3:T:548:LYS:HG3	2.08	0.88
3:R:287:PRO:HB3	3:R:509:LEU:HD21	1.56	0.88
6:Y:842:ALA:HB3	6:Y:843:PRO:HD3	1.54	0.88
1:b:59:HIS:CE1	1:b:64:THR:OG1	2.27	0.88
1:h:74:GLN:O	1:h:74:GLN:NE2	2.07	0.88
1:r:23:THR:OG1	3:R:323:SER:CA	2.21	0.88
3:J:412:ARG:HH22	3:L:356:THR:CG2	1.86	0.88
3:J:450:SER:HA	3:J:453:ASN:HD22	1.37	0.88
3:L:548:LYS:HG3	3:N:645:TRP:CZ2	2.08	0.88

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:N:294:TYR:OH	3:N:502:LEU:HA	1.74	0.88
5:W:592:MET:H	5:W:593:PRO:HD2	1.39	0.88
6:X:655:GLU:CD	6:X:655:GLU:H	1.81	0.88
1:d:21:ARG:NH1	3:D:327:TYR:CZ	2.41	0.88
1:p:59:HIS:HB3	3:P:398:SER:HB2	1.47	0.88
3:F:548:LYS:HG3	3:H:645:TRP:CZ2	2.08	0.88
5:W:1177:ASN:HB2	5:W:1282:THR:HB	1.55	0.88
6:X:937:ALA:HB2	6:X:947:ARG:HA	1.54	0.88
3:D:645:TRP:CZ2	3:H:548:LYS:HG3	2.08	0.87
3:F:294:TYR:OH	3:F:502:LEU:HA	1.74	0.87
7:G:101:MYR:H21	3:H:191:LYS:HE2	1.54	0.87
3:T:167:MET:HA	3:T:167:MET:HE3	1.56	0.87
5:W:458:VAL:HG21	5:W:679:LEU:HG	1.54	0.87
1:f:15:ARG:NH1	3:F:406:ASN:HB3	1.89	0.87
1:h:22:LEU:HD12	1:h:83:VAL:HG21	1.53	0.87
3:D:121:ALA:HB2	3:H:87:PHE:CE1	2.07	0.87
4:U:27:GLN:HG2	6:X:880:LEU:CB	2.01	0.87
1:n:23:THR:HG23	1:n:23:THR:O	1.71	0.87
1:p:35:PHE:CE2	1:p:73:ARG:HG3	2.09	0.87
1:p:58:PRO:CG	3:P:423:GLN:HE21	1.86	0.87
1:t:8:GLN:HE22	3:P:303:ASP:HA	1.39	0.87
1:t:45:ARG:HG3	3:T:400:ALA:C	1.99	0.87
3:D:294:TYR:OH	3:D:502:LEU:HA	1.74	0.87
7:E:101:MYR:H31	3:F:191:LYS:HG3	1.54	0.87
3:H:294:TYR:OH	3:H:502:LEU:HA	1.74	0.87
3:J:356:THR:CG2	3:N:412:ARG:HH22	1.86	0.87
7:M:101:MYR:C7	3:N:200:CYS:SG	2.62	0.87
3:P:294:TYR:OH	3:P:502:LEU:HA	1.74	0.87
3:P:450:SER:HA	3:P:453:ASN:HD22	1.37	0.87
1:l:45:ARG:HG3	3:L:400:ALA:HA	1.51	0.87
1:b:45:ARG:HG3	3:B:401:GLY:H	1.22	0.87
1:b:45:ARG:HG2	3:B:400:ALA:HA	0.88	0.87
1:n:6:ILE:CD1	3:J:586:VAL:CG2	2.49	0.87
3:B:294:TYR:OH	3:B:502:LEU:HA	1.74	0.87
7:C:101:MYR:H21	3:D:191:LYS:HE2	1.55	0.87
3:H:287:PRO:HB3	3:H:509:LEU:HD21	1.57	0.87
3:J:287:PRO:HB3	3:J:509:LEU:HD21	1.56	0.87
1:l:45:ARG:O	1:l:58:PRO:O	1.92	0.87
1:b:4:HIS:NE2	1:b:54:THR:CG2	2.34	0.87
1:f:59:HIS:CB	3:F:397:VAL:O	2.22	0.87
1:d:23:THR:OG1	3:D:323:SER:HA	1.75	0.87

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:J:337:ARG:HH21	3:N:418:ASN:ND2	1.73	0.87
3:J:418:ASN:ND2	3:L:337:ARG:HH21	1.73	0.87
3:P:354:SER:CB	3:P:419:MET:HA	2.05	0.87
6:X:746:TYR:HA	6:X:750:MET:CE	2.05	0.87
6:Y:158:ARG:HG3	6:Y:1205:ASP:HB3	1.55	0.87
1:d:45:ARG:HG2	3:D:400:ALA:CA	1.97	0.87
1:p:60:ALA:O	1:p:62:VAL:HG13	1.75	0.87
1:t:15:ARG:NH1	3:T:402:ALA:CB	2.37	0.87
7:G:101:MYR:C7	3:H:200:CYS:SG	2.62	0.87
3:J:298:PHE:CE2	3:N:355:GLY:HA2	2.10	0.87
3:N:354:SER:CB	3:N:419:MET:HA	2.05	0.87
3:R:418:ASN:ND2	3:T:337:ARG:HH21	1.73	0.87
6:X:189:ASN:HD21	6:X:915:LEU:HA	1.38	0.87
1:l:8:GLN:CG	3:N:302:GLU:CG	2.41	0.87
1:b:16:ALA:O	1:b:22:LEU:HD23	1.73	0.87
1:f:24:LEU:HD21	1:f:80:ARG:NE	1.90	0.87
3:J:354:SER:CB	3:J:419:MET:HA	2.05	0.87
3:P:161:LYS:HZ2	3:T:531:THR:HB	1.40	0.87
3:R:354:SER:CB	3:R:419:MET:HA	2.05	0.87
5:W:766:LEU:HD13	5:W:841:ARG:HH22	1.40	0.87
1:h:15:ARG:NH1	3:H:402:ALA:CB	2.37	0.87
1:n:21:ARG:HH11	1:n:21:ARG:HG2	1.39	0.87
3:B:287:PRO:HB3	3:B:509:LEU:HD21	1.56	0.87
3:D:418:ASN:ND2	3:F:337:ARG:HH21	1.73	0.87
3:F:355:GLY:HA2	3:H:298:PHE:CE2	2.10	0.87
3:F:409:ILE:CD1	3:H:297:ALA:O	2.23	0.87
3:J:609:ARG:HH11	2:K:27:THR:HG23	1.37	0.87
3:L:167:MET:HA	3:L:167:MET:HE3	1.56	0.87
3:T:354:SER:CB	3:T:419:MET:HA	2.05	0.87
4:V:192:PHE:CB	6:Y:376:LEU:HD21	2.05	0.87
6:X:682:GLN:HG3	6:X:683:PRO:HD2	1.57	0.87
1:t:59:HIS:CB	3:T:398:SER:OG	2.10	0.87
1:t:59:HIS:CG	3:T:398:SER:HB2	2.10	0.87
3:D:287:PRO:HB3	3:D:509:LEU:HD21	1.56	0.87
6:Y:277:VAL:HG12	6:Y:278:LEU:HG	1.56	0.87
6:Y:1103:TRP:CD1	6:Y:1113:PRO:HG2	2.10	0.87
1:h:23:THR:HB	3:H:323:SER:HB2	1.56	0.86
3:D:309:GLU:OE1	3:D:335:PRO:HB3	1.75	0.86
3:D:354:SER:CB	3:D:419:MET:HA	2.05	0.86
1:b:51:PHE:O	1:b:74:GLN:HG3	1.74	0.86
1:d:15:ARG:NH1	3:D:402:ALA:HB1	1.90	0.86

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:P:409:ILE:CD1	3:R:297:ALA:O	2.23	0.86
1:l:58:PRO:HG3	3:L:423:GLN:NE2	1.91	0.86
1:b:23:THR:HG23	1:b:25:TYR:CE2	2.09	0.86
1:n:21:ARG:CZ	3:N:327:TYR:OH	2.23	0.86
1:r:58:PRO:CB	3:R:423:GLN:HE21	1.88	0.86
7:I:101:MYR:H21	3:J:191:LYS:HE2	1.55	0.86
3:J:167:MET:HE3	3:J:167:MET:HA	1.56	0.86
3:L:309:GLU:OE1	3:L:335:PRO:HB3	1.76	0.86
3:L:609:ARG:CB	2:M:29:THR:HG22	2.05	0.86
3:R:344:MET:HE2	3:R:502:LEU:HD11	1.58	0.86
6:X:594:ALA:HA	6:X:608:GLN:NE2	1.90	0.86
6:Y:737:ARG:HG2	6:Y:737:ARG:HH21	1.41	0.86
1:b:22:LEU:HG	1:b:83:VAL:HG21	1.54	0.86
1:f:6:ILE:CD1	3:H:586:VAL:CG2	2.45	0.86
3:D:344:MET:HE2	3:D:502:LEU:HD11	1.57	0.86
3:D:409:ILE:CD1	3:F:297:ALA:O	2.23	0.86
3:F:344:MET:HE2	3:F:502:LEU:HD11	1.58	0.86
3:F:418:ASN:ND2	3:H:337:ARG:HH21	1.73	0.86
3:F:609:ARG:HH11	2:G:27:THR:HG23	1.37	0.86
3:L:418:ASN:ND2	3:N:337:ARG:HH21	1.73	0.86
3:N:167:MET:HA	3:N:167:MET:HE3	1.56	0.86
3:R:355:GLY:HA2	3:T:298:PHE:CE2	2.10	0.86
5:W:836:THR:HG21	5:W:854:MET:HB2	1.57	0.86
1:l:45:ARG:CA	3:L:399:ALA:O	2.24	0.86
1:l:74:GLN:O	1:l:74:GLN:NE2	2.08	0.86
1:b:23:THR:HG21	1:b:25:TYR:CE2	2.10	0.86
1:j:59:HIS:HB2	3:J:398:SER:HB2	0.92	0.86
1:t:5:MET:O	1:t:6:ILE:C	2.14	0.86
3:J:491:ALA:HB2	3:J:592:THR:HB	1.58	0.86
3:P:355:GLY:HA2	3:R:298:PHE:CE2	2.10	0.86
6:X:250:ALA:HB1	6:X:339:LEU:HD11	1.55	0.86
2:C:27:THR:HG23	3:H:609:ARG:HH11	1.37	0.86
3:D:297:ALA:O	3:H:409:ILE:CD1	2.23	0.86
2:I:29:THR:HG22	3:N:609:ARG:CB	2.06	0.86
3:L:354:SER:CB	3:L:419:MET:HA	2.05	0.86
3:L:355:GLY:HA2	3:N:298:PHE:CE2	2.10	0.86
3:N:287:PRO:HB3	3:N:509:LEU:HD21	1.56	0.86
3:N:322:TYR:HD2	3:N:327:TYR:CD2	1.93	0.86
3:P:309:GLU:OE1	3:P:335:PRO:HB3	1.76	0.86
1:b:45:ARG:CD	3:B:401:GLY:H	1.88	0.86
1:n:59:HIS:HA	3:N:398:SER:OG	1.75	0.86

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:B:354:SER:CB	3:B:419:MET:HA	2.05	0.86
3:F:287:PRO:HB3	3:F:509:LEU:HD21	1.56	0.86
3:F:309:GLU:OE1	3:F:335:PRO:HB3	1.76	0.86
7:G:101:MYR:H31	3:H:191:LYS:HG3	1.58	0.86
3:H:344:MET:HE2	3:H:502:LEU:HD11	1.57	0.86
3:J:297:ALA:O	3:N:409:ILE:CD1	2.23	0.86
3:P:337:ARG:HH21	3:T:418:ASN:ND2	1.73	0.86
4:U:22:ASP:HB2	4:U:383:ARG:HD3	1.55	0.86
1:t:15:ARG:NH2	3:T:402:ALA:CB	2.37	0.86
2:C:29:THR:HG22	3:H:609:ARG:CB	2.05	0.86
3:D:68:PHE:HB3	3:D:140:VAL:CG1	2.05	0.86
3:D:298:PHE:CE2	3:H:355:GLY:HA2	2.10	0.86
1:b:71:CYS:SG	1:b:73:ARG:HG2	2.15	0.86
1:p:45:ARG:HG2	3:P:400:ALA:HA	0.86	0.86
1:r:15:ARG:NH1	3:R:402:ALA:HB1	1.91	0.86
3:D:337:ARG:HH21	3:H:418:ASN:ND2	1.73	0.86
3:F:167:MET:HE3	3:F:167:MET:HA	1.56	0.86
3:L:409:ILE:CD1	3:N:297:ALA:O	2.23	0.86
3:P:49:ARG:HH12	3:P:115:THR:HG22	1.41	0.86
3:P:287:PRO:HB3	3:P:509:LEU:HD21	1.56	0.86
3:T:294:TYR:OH	3:T:502:LEU:HA	1.74	0.86
6:X:620:VAL:HG11	6:Y:718:THR:HG21	1.57	0.86
6:Y:419:ILE:HD11	6:Y:662:LEU:HB2	1.58	0.86
1:n:23:THR:HB	3:N:323:SER:HB3	1.58	0.86
1:p:58:PRO:HG2	3:P:423:GLN:HG3	1.56	0.86
1:t:23:THR:HG21	1:t:31:GLU:CB	2.05	0.86
1:t:24:LEU:CD2	1:t:35:PHE:CA	2.53	0.86
1:t:44:GLY:HA3	3:T:322:TYR:CZ	2.10	0.86
3:B:491:ALA:HB2	3:B:592:THR:HB	1.58	0.86
3:P:491:ALA:HB2	3:P:592:THR:HB	1.58	0.86
7:S:101:MYR:H31	3:T:191:LYS:HG3	1.58	0.86
5:W:618:LEU:HD13	5:W:619:LYS:N	1.90	0.86
6:Y:1154:ILE:O	6:Y:1154:ILE:HD12	1.76	0.86
1:h:59:HIS:CE1	1:h:60:ALA:C	2.54	0.85
1:j:8:GLN:HE21	3:L:302:GLU:CG	1.88	0.85
1:p:15:ARG:HH11	3:P:406:ASN:CG	1.84	0.85
1:p:35:PHE:HE2	1:p:73:ARG:CG	1.88	0.85
3:D:355:GLY:HA2	3:F:298:PHE:CE2	2.10	0.85
3:D:491:ALA:HB2	3:D:592:THR:HB	1.58	0.85
3:H:309:GLU:OE1	3:H:335:PRO:HB3	1.76	0.85
3:H:494:THR:H	3:H:497:GLU:HB3	1.41	0.85

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:J:294:TYR:OH	3:J:502:LEU:HA	1.74	0.85
3:J:309:GLU:OE1	3:J:335:PRO:HB3	1.75	0.85
3:J:409:ILE:CD1	3:L:297:ALA:O	2.23	0.85
3:L:625:ARG:HG3	3:R:197:ASP:OD1	1.76	0.85
3:P:167:MET:HA	3:P:167:MET:HE3	1.56	0.85
3:T:309:GLU:OE1	3:T:335:PRO:HB3	1.75	0.85
3:T:494:THR:H	3:T:497:GLU:HB3	1.41	0.85
6:Y:367:ARG:O	6:Y:388:MET:HG2	1.76	0.85
6:Y:1093:SER:OG	6:Y:1130:ALA:HB1	1.76	0.85
1:f:45:ARG:HG3	3:F:400:ALA:HA	1.54	0.85
1:t:23:THR:OG1	3:T:323:SER:CA	2.21	0.85
3:H:58:ILE:HG23	3:H:124:GLY:HA2	1.59	0.85
3:J:435:PRO:HG2	3:T:436:TYR:OH	1.75	0.85
3:L:609:ARG:HH11	2:M:27:THR:HG23	1.37	0.85
3:P:57:THR:HG22	3:P:57:THR:O	1.76	0.85
3:R:628:GLN:NE2	5:W:870:ALA:CA	2.32	0.85
6:X:1012:ARG:H	6:X:1012:ARG:CD	1.88	0.85
6:X:1149:THR:CG2	6:X:1162:ALA:H	1.89	0.85
3:D:609:ARG:HH11	2:E:27:THR:HG23	1.37	0.85
7:E:101:MYR:C7	3:F:200:CYS:SG	2.65	0.85
3:F:436:TYR:OH	3:N:435:PRO:HG2	1.76	0.85
3:J:344:MET:HE2	3:J:502:LEU:HD11	1.57	0.85
3:P:298:PHE:CE2	3:T:355:GLY:HA2	2.10	0.85
3:P:472:ARG:NH1	3:R:530:ARG:CG	2.38	0.85
3:R:609:ARG:CB	2:S:29:THR:HG22	2.05	0.85
5:W:103:TYR:HB3	5:W:108:ASN:HD21	1.39	0.85
1:b:15:ARG:HH22	3:B:402:ALA:HB3	1.38	0.85
3:F:609:ARG:CB	2:G:29:THR:HG22	2.06	0.85
3:J:355:GLY:HA2	3:L:298:PHE:CE2	2.10	0.85
3:L:294:TYR:OH	3:L:502:LEU:HA	1.74	0.85
3:R:494:THR:H	3:R:497:GLU:HB3	1.42	0.85
1:b:21:ARG:NH1	3:B:327:TYR:OH	2.10	0.85
1:d:21:ARG:HH12	3:D:327:TYR:HE1	1.23	0.85
3:D:46:LYS:HZ3	3:H:237:GLU:CD	1.85	0.85
3:H:167:MET:HA	3:H:167:MET:HE3	1.56	0.85
3:H:354:SER:CB	3:H:419:MET:HA	2.05	0.85
3:R:49:ARG:HH12	3:R:115:THR:HG22	1.41	0.85
3:R:294:TYR:OH	3:R:502:LEU:HA	1.74	0.85
3:R:409:ILE:CD1	3:T:297:ALA:O	2.23	0.85
6:X:1130:ALA:HB2	6:Y:104:GLN:HG3	1.57	0.85
1:t:24:LEU:HG	1:t:38:ALA:CB	2.07	0.85

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
7:K:101:MYR:C7	3:L:200:CYS:SG	2.65	0.85
3:R:309:GLU:OE1	3:R:335:PRO:HB3	1.76	0.85
3:T:287:PRO:HB3	3:T:509:LEU:HD21	1.56	0.85
5:W:246:ASP:HB3	5:W:256:SER:HA	1.56	0.85
5:W:735:LEU:HD21	5:W:759:ILE:O	1.75	0.85
6:Y:599:ILE:HD12	6:Y:631:ILE:HG23	1.59	0.85
1:h:45:ARG:CA	3:H:399:ALA:O	2.24	0.85
1:j:59:HIS:CB	3:J:397:VAL:O	2.24	0.85
1:t:15:ARG:HH11	3:T:406:ASN:CG	1.83	0.85
3:F:354:SER:CB	3:F:419:MET:HA	2.05	0.85
3:J:84:LYS:HD3	2:K:39:GLY:O	1.77	0.85
3:N:344:MET:HE2	3:N:502:LEU:HD11	1.57	0.85
2:O:29:THR:HG22	3:T:609:ARG:CB	2.06	0.85
7:O:101:MYR:H21	3:P:191:LYS:HE2	1.55	0.85
3:P:84:LYS:HD3	2:Q:39:GLY:O	1.77	0.85
3:R:491:ALA:HB2	3:R:592:THR:HB	1.58	0.85
5:W:810:LEU:HD23	5:W:1001:ILE:HD13	1.57	0.85
1:n:15:ARG:NH2	3:N:402:ALA:HB3	1.91	0.85
3:L:58:ILE:HG23	3:L:124:GLY:HA2	1.59	0.85
3:N:309:GLU:OE1	3:N:335:PRO:HB3	1.76	0.85
3:P:154:ARG:NH1	3:P:154:ARG:HB3	1.92	0.85
3:P:297:ALA:O	3:T:409:ILE:CD1	2.23	0.85
7:Q:101:MYR:C7	3:R:200:CYS:SG	2.64	0.85
3:R:167:MET:HA	3:R:167:MET:HE3	1.56	0.85
1:b:59:HIS:CA	3:B:398:SER:HB2	2.07	0.85
1:p:59:HIS:HB3	3:P:398:SER:N	1.65	0.85
1:r:45:ARG:O	1:r:58:PRO:O	1.94	0.85
3:B:167:MET:HE3	3:B:167:MET:HA	1.56	0.85
2:C:39:GLY:O	3:H:84:LYS:HD3	1.77	0.85
2:I:21:THR:HG23	3:J:219:LYS:HE2	1.59	0.85
3:J:92:TRP:CD1	3:J:156:VAL:HG21	2.12	0.85
3:J:609:ARG:CB	2:K:29:THR:HG22	2.06	0.85
3:L:494:THR:H	3:L:497:GLU:HB3	1.42	0.85
2:Q:21:THR:HG23	3:R:219:LYS:HE2	1.59	0.85
3:T:344:MET:HE2	3:T:502:LEU:HD11	1.57	0.85
6:X:1130:ALA:HB2	6:Y:104:GLN:CG	2.07	0.85
1:r:59:HIS:HB2	3:R:397:VAL:O	1.76	0.85
1:t:60:ALA:CB	3:T:322:TYR:HE1	1.89	0.85
3:F:49:ARG:HH12	3:F:115:THR:HG22	1.41	0.85
3:H:49:ARG:HH12	3:H:115:THR:HG22	1.41	0.85
3:H:154:ARG:HB3	3:H:154:ARG:NH1	1.92	0.85

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:L:287:PRO:HB3	3:L:509:LEU:HD21	1.56	0.85
3:T:49:ARG:HH12	3:T:115:THR:HG22	1.41	0.85
6:Y:410:HIS:CD2	6:Y:411:GLN:N	2.44	0.85
1:f:15:ARG:HH11	3:F:406:ASN:HB3	1.37	0.84
3:F:494:THR:H	3:F:497:GLU:HB3	1.42	0.84
2:K:28:SER:HB3	3:L:216:PRO:HD2	1.59	0.84
3:N:92:TRP:CD1	3:N:156:VAL:HG21	2.12	0.84
3:N:491:ALA:HB2	3:N:592:THR:HB	1.58	0.84
3:P:418:ASN:ND2	3:R:337:ARG:HH21	1.73	0.84
6:X:455:GLN:OE1	6:X:455:GLN:HA	1.75	0.84
1:l:6:ILE:HD13	3:N:586:VAL:HG21	1.60	0.84
1:j:15:ARG:NH1	3:J:402:ALA:HB1	1.92	0.84
3:F:58:ILE:HG23	3:F:124:GLY:HA2	1.59	0.84
2:K:21:THR:HG23	3:L:219:LYS:HE2	1.59	0.84
3:L:491:ALA:HB2	3:L:592:THR:HB	1.58	0.84
3:P:92:TRP:CD1	3:P:156:VAL:HG21	2.12	0.84
3:R:92:TRP:CD1	3:R:156:VAL:HG21	2.12	0.84
6:X:1090:PRO:O	6:X:1129:THR:HG23	1.75	0.84
1:d:15:ARG:HE	3:D:406:ASN:HD21	1.21	0.84
2:C:28:SER:HB3	3:D:216:PRO:HD2	1.59	0.84
3:H:491:ALA:HB2	3:H:592:THR:HB	1.58	0.84
3:J:154:ARG:HB3	3:J:154:ARG:NH1	1.92	0.84
3:P:472:ARG:HH12	3:R:530:ARG:CG	1.90	0.84
5:W:424:SER:HB3	5:W:425:PRO:HD3	1.57	0.84
6:Y:995:PRO:HD2	6:Y:1119:ILE:HG12	1.58	0.84
1:r:6:ILE:HD13	3:T:586:VAL:HG21	1.57	0.84
1:r:45:ARG:HG3	3:R:400:ALA:HA	1.60	0.84
1:t:60:ALA:CB	3:T:322:TYR:CE1	2.60	0.84
3:B:344:MET:HE2	3:B:502:LEU:HD11	1.57	0.84
3:D:494:THR:H	3:D:497:GLU:HB3	1.42	0.84
3:F:154:ARG:HB3	3:F:154:ARG:NH1	1.92	0.84
2:O:39:GLY:O	3:T:84:LYS:HD3	1.77	0.84
3:P:472:ARG:HH12	3:R:530:ARG:HG3	1.41	0.84
6:X:528:PHE:CE1	6:X:575:MET:HE2	2.12	0.84
6:X:656:PHE:CE1	6:X:675:MET:HB2	2.13	0.84
1:d:21:ARG:O	3:D:323:SER:HB2	1.78	0.84
1:h:15:ARG:NH1	3:H:406:ASN:CG	2.36	0.84
1:p:21:ARG:NH1	3:P:327:TYR:CE1	2.44	0.84
1:p:45:ARG:O	1:p:59:HIS:N	2.09	0.84
1:p:60:ALA:CB	3:P:322:TYR:OH	2.25	0.84
3:B:494:THR:H	3:B:497:GLU:HB3	1.41	0.84

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:D:154:ARG:HB3	3:D:154:ARG:NH1	1.92	0.84
3:L:92:TRP:CD1	3:L:156:VAL:HG21	2.12	0.84
3:N:494:THR:H	3:N:497:GLU:HB3	1.42	0.84
5:W:625:ARG:HH11	5:W:625:ARG:HG2	1.42	0.84
1:b:45:ARG:CD	3:B:401:GLY:N	2.41	0.84
3:D:92:TRP:CD1	3:D:156:VAL:HG21	2.12	0.84
3:F:491:ALA:HB2	3:F:592:THR:HB	1.58	0.84
7:I:101:MYR:H31	3:J:191:LYS:HG3	1.60	0.84
3:L:344:MET:HE2	3:L:502:LEU:HD11	1.57	0.84
3:N:49:ARG:HH12	3:N:115:THR:HG22	1.41	0.84
3:P:203:TYR:HB3	3:P:206:GLU:HB3	1.60	0.84
3:P:609:ARG:CB	2:Q:29:THR:HG22	2.06	0.84
3:R:58:ILE:HG23	3:R:124:GLY:HA2	1.59	0.84
3:R:154:ARG:HB3	3:R:154:ARG:NH1	1.92	0.84
5:W:832:LEU:HB3	5:W:890:CYS:SG	2.17	0.84
6:X:827:THR:HG22	6:X:828:GLU:N	1.92	0.84
1:l:44:GLY:HA3	3:L:322:TYR:OH	1.78	0.84
1:f:45:ARG:HG3	3:F:400:ALA:CA	2.06	0.84
1:j:45:ARG:CG	3:J:401:GLY:N	2.35	0.84
1:j:59:HIS:CB	3:J:398:SER:OG	2.24	0.84
2:C:21:THR:HG23	3:D:219:LYS:HE2	1.59	0.84
3:P:609:ARG:HH11	2:Q:27:THR:HG23	1.37	0.84
5:W:497:LEU:HD11	5:W:617:VAL:HG21	1.60	0.84
1:b:15:ARG:NH1	3:B:406:ASN:CB	2.34	0.84
3:D:58:ILE:H	3:D:58:ILE:HD12	1.39	0.84
3:D:358:LEU:HD22	3:D:417:PHE:CD1	2.13	0.84
3:D:609:ARG:CB	2:E:29:THR:HG22	2.06	0.84
2:E:28:SER:HB3	3:F:216:PRO:HD2	1.59	0.84
3:N:154:ARG:HB3	3:N:154:ARG:NH1	1.92	0.84
4:U:198:GLN:NE2	6:X:661:THR:CG2	2.40	0.84
6:X:616:SER:HB3	6:Y:723:ASP:HB2	0.90	0.84
6:X:787:ARG:HE	6:X:925:LEU:HD13	1.43	0.84
6:Y:1122:PRO:HA	6:Y:1159:GLY:HA3	1.60	0.84
1:r:16:ALA:O	1:r:22:LEU:HD23	1.77	0.84
3:B:203:TYR:HB3	3:B:206:GLU:HB3	1.60	0.84
3:B:309:GLU:OE1	3:B:335:PRO:HB3	1.75	0.84
3:B:358:LEU:HD22	3:B:417:PHE:CD1	2.13	0.84
2:E:21:THR:HG23	3:F:219:LYS:HE2	1.59	0.84
3:L:154:ARG:HB3	3:L:154:ARG:NH1	1.92	0.84
3:T:58:ILE:HG23	3:T:124:GLY:HA2	1.58	0.84
1:n:23:THR:CG2	1:n:42:THR:OG1	2.14	0.84

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:p:24:LEU:CD2	1:p:80:ARG:HD2	1.98	0.84
3:F:92:TRP:CD1	3:F:156:VAL:HG21	2.12	0.84
3:F:358:LEU:HD22	3:F:417:PHE:CD1	2.13	0.84
7:M:101:MYR:H31	3:N:191:LYS:HG3	1.58	0.84
3:T:491:ALA:HB2	3:T:592:THR:HB	1.58	0.84
6:X:1106:LYS:HB3	6:X:1113:PRO:HG2	1.60	0.84
1:l:15:ARG:NH2	3:L:402:ALA:CB	2.41	0.83
1:l:21:ARG:HH11	3:L:370:VAL:HG13	1.40	0.83
1:f:58:PRO:HB2	3:F:423:GLN:HE21	1.42	0.83
1:t:58:PRO:HG3	3:T:423:GLN:HE21	1.41	0.83
3:B:58:ILE:HG23	3:B:124:GLY:HA2	1.59	0.83
3:J:203:TYR:HB3	3:J:206:GLU:HB3	1.60	0.83
3:N:358:LEU:HD22	3:N:417:PHE:CD1	2.13	0.83
3:P:494:THR:H	3:P:497:GLU:HB3	1.41	0.83
2:S:28:SER:HB3	3:T:216:PRO:HD2	1.59	0.83
6:X:705:ILE:HG21	6:X:737:ARG:HB3	1.57	0.83
1:p:21:ARG:O	3:P:323:SER:CB	2.26	0.83
7:A:101:MYR:H31	3:B:191:LYS:HG3	1.60	0.83
3:D:50:PRO:HA	3:D:61:LEU:HA	1.59	0.83
3:D:167:MET:HA	3:D:167:MET:HE3	1.56	0.83
2:O:21:THR:HG23	3:P:219:LYS:HE2	1.59	0.83
3:P:344:MET:HE2	3:P:502:LEU:HD11	1.57	0.83
2:S:21:THR:HG23	3:T:219:LYS:HE2	1.59	0.83
1:f:44:GLY:O	3:F:399:ALA:N	2.11	0.83
1:d:21:ARG:NH1	3:D:327:TYR:OH	2.10	0.83
1:r:4:HIS:CE1	1:r:54:THR:CG2	2.60	0.83
2:A:21:THR:HG23	3:B:219:LYS:HE2	1.59	0.83
3:D:57:THR:O	3:D:60:SER:HB3	1.78	0.83
3:N:58:ILE:HG23	3:N:124:GLY:HA2	1.58	0.83
2:O:28:SER:HB3	3:P:216:PRO:HD2	1.60	0.83
5:W:5:PHE:HA	5:W:344:GLY:HA3	1.60	0.83
6:X:277:VAL:HG13	6:X:278:LEU:HD13	1.60	0.83
6:X:782:ASN:OD1	6:X:806:THR:HG21	1.77	0.83
6:X:1032:ILE:HD13	6:X:1038:PRO:HB3	1.60	0.83
1:t:21:ARG:NH1	3:T:327:TYR:CE1	2.46	0.83
3:D:360:TYR:HB2	3:D:367:SER:HB2	1.61	0.83
3:L:358:LEU:HD22	3:L:417:PHE:CD1	2.13	0.83
3:N:360:TYR:HB2	3:N:367:SER:HB2	1.61	0.83
1:f:15:ARG:CZ	3:F:402:ALA:CB	2.56	0.83
1:n:15:ARG:CZ	3:N:402:ALA:HB1	2.03	0.83
1:t:21:ARG:NH1	3:T:327:TYR:CZ	2.44	0.83

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:B:92:TRP:CD1	3:B:156:VAL:HG21	2.12	0.83
3:B:154:ARG:HB3	3:B:154:ARG:NH1	1.92	0.83
3:H:92:TRP:CD1	3:H:156:VAL:HG21	2.12	0.83
3:T:92:TRP:CD1	3:T:156:VAL:HG21	2.12	0.83
1:h:24:LEU:CD1	1:h:80:ARG:HD2	2.08	0.83
3:J:237:GLU:HG3	3:L:142:ILE:CD1	2.08	0.83
3:L:322:TYR:CE1	3:L:396:VAL:HB	2.14	0.83
3:P:530:ARG:HE	3:T:472:ARG:HH21	0.86	0.83
6:X:1182:ASN:ND2	6:X:1184:LEU:HB2	1.94	0.83
1:l:15:ARG:HH11	3:L:406:ASN:CG	1.86	0.83
1:f:8:GLN:NE2	3:H:303:ASP:OD1	2.11	0.83
1:p:56:LEU:CD1	1:p:65:ILE:HG21	2.07	0.83
7:C:101:MYR:H31	3:D:191:LYS:HG3	1.60	0.83
3:F:291:PRO:HB2	3:F:467:THR:CG2	2.09	0.83
6:Y:210:ILE:HB	6:Y:239:ALA:HB3	1.60	0.83
1:l:45:ARG:CD	3:L:400:ALA:HA	2.09	0.83
1:p:4:HIS:HD2	1:p:54:THR:HG22	1.43	0.83
3:H:203:TYR:HB3	3:H:206:GLU:HB3	1.60	0.83
3:H:360:TYR:HB2	3:H:367:SER:HB2	1.61	0.83
3:J:358:LEU:HD22	3:J:417:PHE:CD1	2.13	0.83
3:P:59:ASP:O	5:W:379:VAL:CG1	2.26	0.83
3:R:203:TYR:HB3	3:R:206:GLU:HB3	1.60	0.83
5:W:39:LEU:H	5:W:89:HIS:CE1	1.96	0.83
5:W:748:ARG:HG2	5:W:757:THR:HG22	1.61	0.83
1:d:59:HIS:HA	3:D:398:SER:OG	1.78	0.83
1:n:23:THR:CG2	1:n:25:TYR:CE2	2.62	0.83
2:I:39:GLY:O	3:N:84:LYS:HD3	1.77	0.83
3:T:154:ARG:NH1	3:T:154:ARG:HB3	1.92	0.83
6:X:987:THR:HG22	6:X:989:THR:HG23	1.61	0.83
3:H:431:ILE:CD1	3:H:437:ILE:CG2	2.54	0.83
2:O:27:THR:HG23	3:T:609:ARG:HH11	1.37	0.83
4:U:18:ALA:HB1	4:U:19:PRO:HD2	1.59	0.83
1:l:59:HIS:CD2	3:L:397:VAL:O	2.32	0.82
1:b:21:ARG:CD	3:B:327:TYR:OH	2.27	0.82
1:r:8:GLN:HG2	3:T:302:GLU:HG3	0.87	0.82
3:B:380:VAL:HG11	3:B:384:GLN:HG2	1.60	0.82
3:D:380:VAL:HG11	3:D:384:GLN:HG2	1.60	0.82
3:L:203:TYR:HB3	3:L:206:GLU:HB3	1.60	0.82
2:M:21:THR:HG23	3:N:219:LYS:HE2	1.59	0.82
7:O:101:MYR:H31	3:P:191:LYS:HG3	1.60	0.82
3:P:360:TYR:HB2	3:P:367:SER:HB2	1.60	0.82

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:R:416:ARG:HB2	3:T:298:PHE:HE1	1.44	0.82
5:W:1281:ALA:O	5:W:1282:THR:HG22	1.78	0.82
6:X:787:ARG:CG	6:X:787:ARG:NH2	2.27	0.82
6:X:931:ILE:HD11	6:Y:689:VAL:CG1	2.08	0.82
1:h:59:HIS:HB3	3:H:398:SER:HB2	1.57	0.82
1:h:73:ARG:HD3	1:h:73:ARG:O	1.79	0.82
3:B:360:TYR:HB2	3:B:367:SER:HB2	1.61	0.82
7:E:101:MYR:C2	3:F:191:LYS:NZ	2.42	0.82
2:G:21:THR:HG23	3:H:219:LYS:HE2	1.59	0.82
3:J:360:TYR:HB2	3:J:367:SER:HB2	1.61	0.82
3:L:49:ARG:HH12	3:L:115:THR:HG22	1.41	0.82
3:N:291:PRO:HB2	3:N:467:THR:CG2	2.09	0.82
3:R:358:LEU:HD22	3:R:417:PHE:CD1	2.13	0.82
3:T:358:LEU:HD22	3:T:417:PHE:CD1	2.13	0.82
3:F:203:TYR:HB3	3:F:206:GLU:HB3	1.60	0.82
2:G:28:SER:HB3	3:H:216:PRO:HD2	1.59	0.82
3:P:380:VAL:HG11	3:P:384:GLN:HG2	1.60	0.82
5:W:516:MET:HG3	5:W:577:LEU:HD22	1.59	0.82
6:Y:742:MET:HB3	6:Y:825:GLY:O	1.80	0.82
1:l:59:HIS:CB	3:L:398:SER:CB	2.28	0.82
1:d:22:LEU:HD21	1:d:83:VAL:CG2	2.08	0.82
1:h:23:THR:CG2	3:H:323:SER:HA	2.09	0.82
3:B:49:ARG:HH12	3:B:115:THR:HG22	1.41	0.82
3:D:84:LYS:HD3	2:E:39:GLY:O	1.77	0.82
2:I:28:SER:HB3	3:J:216:PRO:HD2	1.59	0.82
3:P:358:LEU:HD22	3:P:417:PHE:CD1	2.13	0.82
3:R:176:ALA:HB1	3:R:628:GLN:HG3	1.62	0.82
6:X:317:PHE:HB2	6:X:373:ARG:NH2	1.94	0.82
1:p:59:HIS:CD2	3:P:398:SER:CB	2.58	0.82
3:J:380:VAL:HG11	3:J:384:GLN:HG2	1.60	0.82
7:K:101:MYR:C2	3:L:191:LYS:NZ	2.42	0.82
2:M:28:SER:HB3	3:N:216:PRO:HD2	1.59	0.82
3:P:322:TYR:O	3:P:323:SER:OG	1.96	0.82
5:W:198:ARG:HH21	6:X:648:ALA:HB1	1.43	0.82
1:r:17:ALA:N	1:r:22:LEU:HD21	1.95	0.82
3:D:317:ARG:NH1	3:D:448:PRO:HB2	1.95	0.82
3:P:317:ARG:NH1	3:P:448:PRO:HB2	1.95	0.82
2:Q:28:SER:HB3	3:R:216:PRO:HD2	1.59	0.82
4:V:192:PHE:HA	6:Y:376:LEU:HD23	1.60	0.82
6:X:237:GLU:HA	6:X:1151:ALA:HB3	1.62	0.82
6:X:425:ASN:N	6:X:425:ASN:HD22	1.77	0.82

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:j:8:GLN:NE2	3:L:303:ASP:OD1	2.12	0.82
3:D:298:PHE:HE1	3:H:416:ARG:HB2	1.45	0.82
3:J:176:ALA:HB1	3:J:628:GLN:HG3	1.62	0.82
3:J:494:THR:H	3:J:497:GLU:HB3	1.42	0.82
3:P:59:ASP:O	5:W:379:VAL:HG12	1.79	0.82
3:T:176:ALA:HB1	3:T:628:GLN:HG3	1.62	0.82
3:T:380:VAL:HG11	3:T:384:GLN:HG2	1.60	0.82
3:T:431:ILE:CD1	3:T:437:ILE:CG2	2.54	0.82
6:X:247:THR:CG2	6:X:308:ASN:OD1	2.28	0.82
6:X:697:MET:HE3	6:X:822:MET:HG3	1.62	0.82
6:Y:244:VAL:HG21	6:Y:265:PRO:HA	1.62	0.82
1:f:6:ILE:HD13	3:H:586:VAL:HG21	1.62	0.82
1:p:22:LEU:HD13	1:p:42:THR:O	1.79	0.82
1:r:59:HIS:HA	3:R:398:SER:CB	2.09	0.82
1:t:60:ALA:HB2	3:T:322:TYR:HH	1.44	0.82
3:H:358:LEU:HD22	3:H:417:PHE:CD1	2.13	0.82
3:T:317:ARG:NH1	3:T:448:PRO:HB2	1.95	0.82
6:X:409:MET:HB2	6:X:445:PHE:O	1.79	0.82
6:X:875:GLN:NE2	6:X:930:ARG:HG2	1.94	0.82
6:Y:707:PHE:O	6:Y:710:ARG:HG2	1.78	0.82
1:d:23:THR:HG23	3:D:323:SER:CA	2.08	0.82
1:d:57:ALA:HB1	1:d:59:HIS:HD2	1.35	0.82
1:p:15:ARG:CZ	3:P:402:ALA:HB1	2.09	0.82
1:r:58:PRO:CG	3:R:423:GLN:NE2	2.40	0.82
3:F:372:ALA:HB3	3:F:444:LEU:HD12	1.62	0.82
3:F:602:SER:OG	3:H:171:THR:HG21	1.80	0.82
3:J:416:ARG:HB2	3:L:298:PHE:HE1	1.44	0.82
3:L:602:SER:OG	3:N:171:THR:HG21	1.80	0.82
3:P:237:GLU:CB	3:R:46:LYS:HZ1	1.92	0.82
3:T:381:PRO:HG3	3:T:436:TYR:CZ	2.15	0.82
5:W:764:THR:HG21	5:W:1004:VAL:HA	1.62	0.82
1:d:23:THR:HG21	3:D:324:GLY:N	1.93	0.82
3:F:360:TYR:HB2	3:F:367:SER:HB2	1.61	0.82
3:H:317:ARG:NH1	3:H:448:PRO:HB2	1.95	0.82
3:L:149:ILE:O	3:L:153:MET:HB2	1.80	0.82
3:L:360:TYR:HB2	3:L:367:SER:HB2	1.61	0.82
3:L:372:ALA:HB3	3:L:444:LEU:HD12	1.62	0.82
3:L:416:ARG:HB2	3:N:298:PHE:HE1	1.45	0.82
3:N:149:ILE:O	3:N:153:MET:HB2	1.80	0.82
3:P:602:SER:OG	3:R:171:THR:HG21	1.80	0.82
5:W:409:ILE:HG12	5:W:779:MET:HE3	1.61	0.82

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:b:24:LEU:CD2	1:b:80:ARG:CD	2.53	0.81
3:D:171:THR:HG21	3:H:602:SER:OG	1.80	0.81
3:D:281:PRO:HD3	3:F:622:GLU:OE1	1.80	0.81
3:D:372:ALA:HB3	3:D:444:LEU:HD12	1.62	0.81
3:D:381:PRO:HG3	3:D:436:TYR:CZ	2.15	0.81
3:D:602:SER:OG	3:F:171:THR:HG21	1.80	0.81
3:J:372:ALA:HB3	3:J:444:LEU:HD12	1.62	0.81
3:N:431:ILE:CD1	3:N:437:ILE:CG2	2.54	0.81
3:P:298:PHE:HE1	3:T:416:ARG:HB2	1.45	0.81
3:R:602:SER:OG	3:T:171:THR:HG21	1.80	0.81
5:W:398:LEU:H	5:W:398:LEU:HD23	1.44	0.81
5:W:748:ARG:NH1	5:W:789:ARG:HA	1.94	0.81
6:X:996:MET:HE2	6:X:996:MET:HA	1.61	0.81
1:f:53:LEU:HD13	1:f:74:GLN:NE2	1.95	0.81
1:f:58:PRO:HB2	3:F:423:GLN:NE2	1.93	0.81
1:t:24:LEU:HD23	1:t:35:PHE:CA	2.04	0.81
3:B:381:PRO:HG3	3:B:436:TYR:CZ	2.15	0.81
3:D:176:ALA:HB1	3:D:628:GLN:HG3	1.62	0.81
3:D:291:PRO:HB2	3:D:467:THR:CG2	2.09	0.81
3:F:381:PRO:HG3	3:F:436:TYR:CZ	2.15	0.81
3:L:291:PRO:HB2	3:L:467:THR:CG2	2.09	0.81
3:N:380:VAL:HG11	3:N:384:GLN:HG2	1.60	0.81
3:T:203:TYR:HB3	3:T:206:GLU:HB3	1.60	0.81
6:X:958:HIS:CD2	6:X:959:ARG:HH21	1.97	0.81
6:Y:652:GLN:HA	6:Y:773:ARG:HD3	1.62	0.81
1:f:21:ARG:CD	3:F:327:TYR:OH	2.26	0.81
1:h:45:ARG:HG2	3:H:400:ALA:CA	2.04	0.81
1:j:23:THR:CG2	3:J:323:SER:CA	2.58	0.81
1:j:24:LEU:HD12	1:j:41:VAL:CG2	2.06	0.81
1:t:59:HIS:HB3	3:T:398:SER:HB2	1.42	0.81
2:A:28:SER:HB3	3:B:216:PRO:HD2	1.59	0.81
3:B:372:ALA:HB3	3:B:444:LEU:HD12	1.62	0.81
3:D:203:TYR:HB3	3:D:206:GLU:HB3	1.60	0.81
3:D:237:GLU:CD	3:F:46:LYS:HZ3	1.88	0.81
3:D:416:ARG:HB2	3:F:298:PHE:HE1	1.44	0.81
3:F:317:ARG:NH1	3:F:448:PRO:HB2	1.95	0.81
3:H:149:ILE:O	3:H:153:MET:HB2	1.80	0.81
3:H:170:LEU:HD12	3:H:170:LEU:O	1.81	0.81
3:H:277:ASP:C	3:H:278:THR:HG22	2.06	0.81
3:H:372:ALA:HB3	3:H:444:LEU:HD12	1.62	0.81
3:H:381:PRO:HG3	3:H:436:TYR:CZ	2.15	0.81

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:J:317:ARG:NH1	3:J:448:PRO:HB2	1.95	0.81
3:L:170:LEU:HD12	3:L:170:LEU:O	1.81	0.81
3:N:203:TYR:HB3	3:N:206:GLU:HB3	1.60	0.81
3:P:149:ILE:O	3:P:153:MET:HB2	1.80	0.81
3:R:277:ASP:C	3:R:278:THR:HG22	2.05	0.81
3:R:317:ARG:NH1	3:R:448:PRO:HB2	1.95	0.81
3:T:360:TYR:HB2	3:T:367:SER:HB2	1.61	0.81
4:U:27:GLN:O	6:X:882:PRO:HD3	1.80	0.81
5:W:823:THR:HG21	5:W:846:ILE:HG12	1.60	0.81
5:W:1096:PHE:H	5:W:1096:PHE:HD1	1.25	0.81
6:X:222:PHE:O	6:X:1154:ILE:HG23	1.79	0.81
6:Y:196:GLU:OE1	6:Y:313:LYS:HB2	1.80	0.81
6:Y:355:LYS:HZ3	6:Y:1156:THR:CG2	1.92	0.81
6:Y:773:ARG:HB2	6:Y:773:ARG:HH11	1.45	0.81
3:D:431:ILE:CD1	3:D:437:ILE:CG2	2.54	0.81
3:J:171:THR:HG21	3:N:602:SER:OG	1.80	0.81
3:J:281:PRO:HD3	3:L:622:GLU:OE1	1.81	0.81
3:N:291:PRO:CB	3:N:467:THR:CG2	2.56	0.81
3:P:416:ARG:HB2	3:R:298:PHE:HE1	1.44	0.81
4:U:224:ASN:ND2	4:U:346:LEU:HD11	1.95	0.81
6:Y:778:ASP:HA	6:Y:940:GLN:HB3	1.60	0.81
1:j:74:GLN:OE1	1:j:75:PRO:N	2.13	0.81
1:r:15:ARG:NH1	3:R:406:ASN:HB3	1.95	0.81
3:B:317:ARG:NH1	3:B:448:PRO:HB2	1.95	0.81
3:F:237:GLU:CD	3:H:46:LYS:HZ3	1.88	0.81
3:F:431:ILE:CD1	3:F:437:ILE:CG2	2.54	0.81
3:J:530:ARG:NE	3:N:472:ARG:NH2	2.28	0.81
3:L:317:ARG:NH1	3:L:448:PRO:HB2	1.95	0.81
3:N:277:ASP:C	3:N:278:THR:HG22	2.06	0.81
3:P:176:ALA:HB1	3:P:628:GLN:HG3	1.62	0.81
3:P:622:GLU:OE1	3:T:281:PRO:HD3	1.81	0.81
3:R:291:PRO:HB2	3:R:467:THR:CG2	2.09	0.81
1:t:45:ARG:CG	3:T:401:GLY:N	2.44	0.81
3:D:530:ARG:NE	3:H:472:ARG:NH2	2.28	0.81
3:F:277:ASP:C	3:F:278:THR:HG22	2.06	0.81
3:N:381:PRO:HG3	3:N:436:TYR:CZ	2.15	0.81
7:Q:101:MYR:C2	3:R:191:LYS:NZ	2.42	0.81
3:R:149:ILE:O	3:R:153:MET:HB2	1.81	0.81
3:R:381:PRO:HG3	3:R:436:TYR:CZ	2.15	0.81
6:Y:220:ILE:HG22	6:Y:221:GLY:H	1.45	0.81
1:r:16:ALA:C	1:r:22:LEU:CD2	2.41	0.81

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:B:149:ILE:O	3:B:153:MET:HB2	1.80	0.81
3:J:149:ILE:O	3:J:153:MET:HB2	1.80	0.81
3:J:170:LEU:O	3:J:170:LEU:HD12	1.81	0.81
6:Y:543:THR:HG22	6:Y:577:SER:H	1.46	0.81
1:b:61:ASN:HB2	3:B:395:THR:CG2	2.11	0.81
3:F:380:VAL:HG11	3:F:384:GLN:HG2	1.60	0.81
3:H:176:ALA:HB1	3:H:628:GLN:HG3	1.62	0.81
3:H:358:LEU:CD2	3:H:417:PHE:CE1	2.64	0.81
3:L:380:VAL:HG11	3:L:384:GLN:HG2	1.60	0.81
3:N:170:LEU:O	3:N:170:LEU:HD12	1.81	0.81
7:S:101:MYR:C2	3:T:191:LYS:NZ	2.44	0.81
3:T:372:ALA:HB3	3:T:444:LEU:HD12	1.62	0.81
6:X:795:ASP:CG	6:Y:737:ARG:HH12	1.87	0.81
1:l:24:LEU:CD2	1:l:80:ARG:NE	2.43	0.81
1:l:74:GLN:HE21	1:l:74:GLN:C	1.87	0.81
3:F:416:ARG:HB2	3:H:298:PHE:HE1	1.45	0.81
3:J:277:ASP:C	3:J:278:THR:HG22	2.06	0.81
3:L:358:LEU:CD2	3:L:417:PHE:CE1	2.64	0.81
3:L:381:PRO:HG3	3:L:436:TYR:CZ	2.15	0.81
3:R:360:TYR:HB2	3:R:367:SER:HB2	1.61	0.81
3:R:372:ALA:HB3	3:R:444:LEU:HD12	1.62	0.81
5:W:736:VAL:HG21	5:W:780:LEU:HD21	1.63	0.81
6:Y:627:ASN:HD22	6:Y:630:LEU:HG	1.43	0.81
6:Y:692:THR:HG23	6:Y:693:TYR:CD1	2.15	0.81
1:l:23:THR:HB	3:L:323:SER:HA	1.61	0.81
1:t:4:HIS:ND1	1:t:54:THR:HG22	1.96	0.81
3:D:277:ASP:C	3:D:278:THR:HG22	2.06	0.81
3:D:358:LEU:CD2	3:D:417:PHE:CE1	2.64	0.81
3:J:291:PRO:HB2	3:J:467:THR:CG2	2.09	0.81
3:J:381:PRO:HG3	3:J:436:TYR:CZ	2.15	0.81
3:J:602:SER:OG	3:L:171:THR:HG21	1.80	0.81
3:P:277:ASP:C	3:P:278:THR:HG22	2.06	0.81
3:P:381:PRO:HG3	3:P:436:TYR:CZ	2.15	0.81
3:P:575:MET:HB2	3:T:469:VAL:HG23	1.63	0.81
3:T:149:ILE:O	3:T:153:MET:HB2	1.80	0.81
6:X:242:ARG:HH21	6:X:1085:LEU:HD13	1.43	0.81
1:b:59:HIS:ND1	1:b:61:ASN:ND2	2.29	0.80
1:h:23:THR:OG1	3:H:323:SER:CA	2.13	0.80
3:D:469:VAL:HG23	3:F:575:MET:HB2	1.63	0.80
3:F:170:LEU:O	3:F:170:LEU:HD12	1.81	0.80
3:H:291:PRO:HB2	3:H:467:THR:CG2	2.09	0.80

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:J:298:PHE:HE1	3:N:416:ARG:HB2	1.45	0.80
3:N:317:ARG:NH1	3:N:448:PRO:HB2	1.95	0.80
4:U:242:ASN:HD21	4:U:244:ASP:HB3	1.46	0.80
6:X:875:GLN:HG3	6:X:876:ASP:H	1.46	0.80
6:Y:575:MET:HE1	6:Y:747:ILE:HD11	1.63	0.80
1:d:61:ASN:CG	3:D:395:THR:HG21	2.05	0.80
1:j:15:ARG:CZ	3:J:402:ALA:CB	2.59	0.80
1:j:21:ARG:O	3:J:323:SER:CB	2.29	0.80
3:H:380:VAL:HG11	3:H:384:GLN:HG2	1.60	0.80
3:R:281:PRO:HD3	3:T:622:GLU:OE1	1.81	0.80
3:T:277:ASP:C	3:T:278:THR:HG22	2.05	0.80
4:U:16:GLN:HG2	4:U:20:LEU:HD21	1.62	0.80
5:W:1134:VAL:HG22	5:W:1219:ILE:HD11	1.63	0.80
6:X:697:MET:HG2	6:X:744:PRO:HA	1.64	0.80
1:l:53:LEU:CD1	1:l:74:GLN:OE1	2.28	0.80
1:f:45:ARG:CA	3:F:399:ALA:O	2.30	0.80
1:p:15:ARG:NH1	3:P:406:ASN:CB	2.20	0.80
1:p:15:ARG:NH2	3:P:402:ALA:HB3	1.95	0.80
3:B:481:VAL:HG12	3:B:482:ASP:H	1.47	0.80
3:F:149:ILE:O	3:F:153:MET:HB2	1.80	0.80
3:F:358:LEU:CD2	3:F:417:PHE:CE1	2.64	0.80
3:H:291:PRO:CB	3:H:467:THR:CG2	2.56	0.80
3:P:481:VAL:HG12	3:P:482:ASP:H	1.47	0.80
3:R:380:VAL:HG11	3:R:384:GLN:HG2	1.60	0.80
3:R:431:ILE:CD1	3:R:437:ILE:CG2	2.54	0.80
5:W:9:LEU:HD13	5:W:318:VAL:HG21	1.63	0.80
5:W:823:THR:HG23	5:W:850:LEU:HD13	1.63	0.80
1:j:6:ILE:HD11	3:L:586:VAL:HG11	1.61	0.80
1:j:15:ARG:NH1	3:J:406:ASN:ND2	2.29	0.80
3:D:149:ILE:O	3:D:153:MET:HB2	1.80	0.80
3:J:575:MET:CB	3:N:469:VAL:HG23	2.12	0.80
3:N:176:ALA:HB1	3:N:628:GLN:HG3	1.62	0.80
3:P:372:ALA:HB3	3:P:444:LEU:HD12	1.62	0.80
4:U:135:MET:O	4:U:138:ILE:HG22	1.80	0.80
1:l:8:GLN:HG2	3:N:302:GLU:HG3	0.82	0.80
1:n:8:GLN:HE22	3:J:303:ASP:HA	1.47	0.80
1:n:45:ARG:HG3	3:N:400:ALA:C	2.05	0.80
1:t:24:LEU:HG	1:t:38:ALA:HB1	1.63	0.80
3:B:277:ASP:C	3:B:278:THR:HG22	2.06	0.80
3:B:431:ILE:CD1	3:B:437:ILE:CG2	2.54	0.80
3:D:68:PHE:HB3	3:D:140:VAL:HG12	1.62	0.80

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:D:469:VAL:HG23	3:F:575:MET:CB	2.12	0.80
3:F:176:ALA:HB1	3:F:628:GLN:HG3	1.62	0.80
3:F:281:PRO:HD3	3:H:622:GLU:OE1	1.81	0.80
3:F:355:GLY:CA	3:H:298:PHE:CE2	2.65	0.80
3:F:469:VAL:HG23	3:H:575:MET:CB	2.12	0.80
3:L:469:VAL:HG23	3:N:575:MET:CB	2.12	0.80
3:P:431:ILE:CD1	3:P:437:ILE:CG2	2.54	0.80
3:P:575:MET:CB	3:T:469:VAL:HG23	2.12	0.80
3:R:469:VAL:HG23	3:T:575:MET:HB2	1.63	0.80
4:U:171:THR:O	4:U:175:MET:HG3	1.81	0.80
6:X:587:MET:HE1	6:X:599:ILE:HD11	1.64	0.80
1:f:15:ARG:NH1	3:F:406:ASN:CB	2.44	0.80
1:d:15:ARG:CZ	3:D:402:ALA:CB	2.60	0.80
1:h:6:ILE:HD13	3:D:586:VAL:HG21	1.61	0.80
3:B:291:PRO:CB	3:B:467:THR:CG2	2.56	0.80
3:D:53:THR:CG2	3:D:54:SER:H	1.92	0.80
3:F:274:ILE:CD1	3:F:514:LEU:HD22	2.10	0.80
3:H:92:TRP:CZ3	3:H:210:VAL:HG11	2.17	0.80
3:H:274:ILE:CD1	3:H:514:LEU:HD22	2.10	0.80
3:L:237:GLU:CB	3:N:46:LYS:HZ1	1.93	0.80
3:L:469:VAL:HG23	3:N:575:MET:HB2	1.63	0.80
3:N:274:ILE:CD1	3:N:514:LEU:HD22	2.10	0.80
3:P:171:THR:HG21	3:T:602:SER:OG	1.80	0.80
3:P:298:PHE:CE2	3:T:355:GLY:CA	2.65	0.80
3:P:355:GLY:CA	3:R:298:PHE:CE2	2.65	0.80
3:T:92:TRP:CZ3	3:T:210:VAL:HG11	2.17	0.80
5:W:1130:PRO:HB2	5:W:1158:ASP:OD2	1.80	0.80
6:X:754:GLU:O	6:X:758:LEU:HB2	1.82	0.80
1:t:6:ILE:HD13	3:P:586:VAL:HG21	1.64	0.80
3:D:472:ARG:NH2	3:F:530:ARG:NE	2.28	0.80
3:L:176:ALA:HB1	3:L:628:GLN:HG3	1.62	0.80
3:N:481:VAL:HG12	3:N:482:ASP:H	1.47	0.80
3:T:291:PRO:HB2	3:T:467:THR:CG2	2.09	0.80
4:U:245:GLN:HE21	4:U:246:ALA:N	1.78	0.80
4:U:341:TRP:HA	4:U:346:LEU:HD12	1.64	0.80
5:W:248:LEU:HD23	5:W:254:LEU:HD22	1.61	0.80
6:X:827:THR:HG21	6:X:831:LEU:CD2	2.11	0.80
1:p:73:ARG:NE	1:p:77:GLU:OE2	2.14	0.80
3:B:176:ALA:HB1	3:B:628:GLN:HG3	1.62	0.80
3:D:575:MET:HB2	3:H:469:VAL:HG23	1.63	0.80
3:F:92:TRP:CZ3	3:F:210:VAL:HG11	2.17	0.80

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:L:355:GLY:CA	3:N:298:PHE:CE2	2.65	0.80
3:L:625:ARG:CG	3:R:197:ASP:OD1	2.28	0.80
3:P:116:SER:HB3	3:P:133:ASN:N	1.97	0.80
3:P:291:PRO:HB2	3:P:467:THR:CG2	2.09	0.80
3:P:358:LEU:CD2	3:P:417:PHE:CE1	2.64	0.80
3:R:358:LEU:CD2	3:R:417:PHE:CE1	2.64	0.80
3:R:469:VAL:HG23	3:T:575:MET:CB	2.12	0.80
3:T:358:LEU:CD2	3:T:417:PHE:CE1	2.64	0.80
1:r:15:ARG:NH1	3:R:406:ASN:CB	2.45	0.80
3:D:92:TRP:CZ3	3:D:210:VAL:HG11	2.17	0.80
3:D:354:SER:HB3	3:D:419:MET:HA	1.64	0.80
3:D:622:GLU:OE1	3:H:281:PRO:HD3	1.81	0.80
7:G:101:MYR:C2	3:H:191:LYS:NZ	2.44	0.80
3:J:51:VAL:CG1	3:J:60:SER:O	2.28	0.80
3:J:431:ILE:CD1	3:J:437:ILE:CG2	2.54	0.80
3:N:358:LEU:CD2	3:N:417:PHE:CE1	2.64	0.80
3:P:472:ARG:HH12	3:R:530:ARG:HD3	1.45	0.80
3:R:237:GLU:CB	3:T:46:LYS:HZ1	1.95	0.80
3:T:481:VAL:HG12	3:T:482:ASP:H	1.47	0.80
6:X:958:HIS:HD2	6:X:959:ARG:HH21	1.28	0.80
6:Y:593:LEU:HB2	6:Y:596:PRO:HG2	1.64	0.80
1:p:62:VAL:C	1:p:64:THR:N	2.40	0.80
1:t:45:ARG:O	1:t:59:HIS:HA	1.80	0.80
3:B:170:LEU:O	3:B:170:LEU:HD12	1.81	0.80
3:J:355:GLY:CA	3:L:298:PHE:CE2	2.65	0.80
3:J:358:LEU:CD2	3:J:417:PHE:CE1	2.64	0.80
3:P:170:LEU:HD12	3:P:170:LEU:O	1.80	0.80
3:R:355:GLY:CA	3:T:298:PHE:CE2	2.65	0.80
3:T:170:LEU:HD12	3:T:170:LEU:O	1.81	0.80
3:T:291:PRO:CB	3:T:467:THR:CG2	2.55	0.80
4:V:181:PRO:HG2	6:Y:412:ASP:OD2	1.82	0.80
5:W:71:HIS:CE1	5:W:73:ALA:HB3	2.17	0.80
6:X:195:ILE:O	6:X:195:ILE:CG2	2.28	0.80
1:j:59:HIS:HB3	3:J:398:SER:OG	1.81	0.79
1:j:61:ASN:CG	3:J:395:THR:CG2	2.54	0.79
1:j:74:GLN:OE1	1:j:75:PRO:C	2.25	0.79
1:n:44:GLY:N	3:N:322:TYR:OH	2.15	0.79
3:B:358:LEU:CD2	3:B:417:PHE:CE1	2.64	0.79
3:F:409:ILE:HD13	3:H:297:ALA:O	1.83	0.79
3:J:469:VAL:HG23	3:L:575:MET:CB	2.12	0.79
3:P:469:VAL:HG23	3:R:575:MET:CB	2.12	0.79

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
4:U:32:CYS:HB3	6:X:958:HIS:HE1	1.06	0.79
5:W:35:GLN:HG3	5:W:96:VAL:HG21	1.64	0.79
6:Y:1168:ARG:HB3	6:Y:1190:LEU:HD23	1.64	0.79
1:j:24:LEU:HD11	1:j:41:VAL:CG2	2.08	0.79
1:r:24:LEU:HG	1:r:38:ALA:HB1	1.65	0.79
1:r:45:ARG:HG3	3:R:400:ALA:CA	2.12	0.79
1:t:47:THR:HG23	1:t:59:HIS:O	1.81	0.79
3:D:170:LEU:HD12	3:D:170:LEU:O	1.81	0.79
3:P:92:TRP:CZ3	3:P:210:VAL:HG11	2.17	0.79
6:Y:653:PRO:HA	6:Y:678:PRO:O	1.83	0.79
1:d:45:ARG:CG	3:D:401:GLY:N	2.40	0.79
1:n:23:THR:HG22	1:n:42:THR:HG1	1.45	0.79
1:p:58:PRO:O	1:p:59:HIS:CD2	2.36	0.79
7:A:101:MYR:H143	7:A:101:MYR:C10	2.11	0.79
3:B:291:PRO:HB2	3:B:467:THR:CG2	2.09	0.79
3:D:297:ALA:O	3:H:409:ILE:HD13	1.83	0.79
3:H:116:SER:HB3	3:H:133:ASN:N	1.97	0.79
3:J:291:PRO:CB	3:J:467:THR:CG2	2.56	0.79
3:J:622:GLU:OE1	3:N:281:PRO:HD3	1.81	0.79
7:M:101:MYR:C2	3:N:191:LYS:NZ	2.44	0.79
3:R:170:LEU:O	3:R:170:LEU:HD12	1.81	0.79
6:X:795:ASP:CG	6:X:798:VAL:HG21	2.07	0.79
6:Y:589:LEU:HG	6:Y:593:LEU:HD21	1.63	0.79
1:p:21:ARG:NH1	3:P:327:TYR:OH	2.14	0.79
1:t:15:ARG:NH2	3:T:402:ALA:HB3	1.96	0.79
3:D:343:ALA:HB2	3:D:478:GLY:O	1.83	0.79
3:J:358:LEU:CG	3:J:417:PHE:HE1	1.96	0.79
3:L:92:TRP:CZ3	3:L:210:VAL:HG11	2.17	0.79
3:L:431:ILE:CD1	3:L:437:ILE:CG2	2.54	0.79
3:P:281:PRO:HD3	3:R:622:GLU:OE1	1.81	0.79
3:R:92:TRP:CZ3	3:R:210:VAL:HG11	2.17	0.79
3:R:409:ILE:HD13	3:T:297:ALA:O	1.82	0.79
4:U:220:THR:HG21	4:U:404:VAL:HA	1.65	0.79
5:W:477:ARG:HA	5:W:650:VAL:HG11	1.64	0.79
6:X:299:LEU:HD11	6:X:904:ALA:HB3	1.63	0.79
6:X:889:ARG:HG3	6:X:890:GLN:H	1.47	0.79
6:Y:244:VAL:CG2	6:Y:265:PRO:HA	2.12	0.79
6:Y:593:LEU:HD23	6:Y:593:LEU:H	1.47	0.79
1:p:23:THR:HG23	1:p:25:TYR:CE2	2.17	0.79
7:E:101:MYR:H143	7:E:101:MYR:C10	2.12	0.79
3:J:343:ALA:HB2	3:J:478:GLY:O	1.83	0.79

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:J:609:ARG:HD2	2:K:29:THR:CG2	2.13	0.79
3:L:358:LEU:CG	3:L:417:PHE:HE1	1.96	0.79
3:T:287:PRO:O	3:T:471:LEU:HD12	1.83	0.79
6:X:242:ARG:HH21	6:X:1085:LEU:HD11	1.47	0.79
6:X:524:ASP:C	6:X:526:ALA:H	1.88	0.79
1:n:23:THR:CG2	1:n:25:TYR:HE2	1.96	0.79
3:D:575:MET:CB	3:H:469:VAL:HG23	2.12	0.79
3:D:609:ARG:HD2	2:E:29:THR:CG2	2.13	0.79
3:J:118:TYR:CE1	3:J:130:PRO:HB3	2.18	0.79
2:O:29:THR:CG2	3:T:609:ARG:HD2	2.13	0.79
3:R:238:VAL:HG11	3:T:78:MET:HE2	1.65	0.79
3:R:343:ALA:HB2	3:R:478:GLY:O	1.83	0.79
3:T:343:ALA:HB2	3:T:478:GLY:O	1.83	0.79
3:T:354:SER:HB3	3:T:419:MET:HA	1.64	0.79
4:U:9:LEU:HD23	4:U:119:LEU:HD23	1.63	0.79
6:X:1125:TYR:CE2	6:X:1127:CYS:HB3	2.17	0.79
1:b:15:ARG:NH1	3:B:406:ASN:CG	2.40	0.79
1:d:60:ALA:H	3:D:398:SER:HA	1.47	0.79
1:t:15:ARG:NH1	3:T:406:ASN:CG	2.40	0.79
3:B:92:TRP:CZ3	3:B:210:VAL:HG11	2.17	0.79
3:D:481:VAL:HG12	3:D:482:ASP:H	1.47	0.79
3:H:118:TYR:CE1	3:H:130:PRO:HB3	2.18	0.79
2:I:29:THR:CG2	3:N:609:ARG:HD2	2.13	0.79
3:J:297:ALA:O	3:N:409:ILE:HD13	1.82	0.79
3:L:281:PRO:HD3	3:N:622:GLU:OE1	1.81	0.79
3:L:287:PRO:O	3:L:471:LEU:HD12	1.83	0.79
7:O:101:MYR:H143	7:O:101:MYR:C10	2.11	0.79
7:Q:101:MYR:H143	7:Q:101:MYR:C10	2.11	0.79
4:V:195:GLN:NE2	6:Y:377:ILE:H	1.80	0.79
1:n:23:THR:HG23	1:n:25:TYR:HE2	1.47	0.79
3:D:50:PRO:CA	3:D:60:SER:O	2.30	0.79
3:D:78:MET:HE2	3:H:238:VAL:HG11	1.65	0.79
3:D:118:TYR:CE1	3:D:130:PRO:HB3	2.18	0.79
3:D:291:PRO:CB	3:D:467:THR:CG2	2.56	0.79
3:D:298:PHE:CE2	3:H:355:GLY:CA	2.65	0.79
3:F:238:VAL:HG11	3:H:78:MET:HE2	1.65	0.79
3:F:358:LEU:CG	3:F:417:PHE:HE1	1.96	0.79
3:J:298:PHE:CE2	3:N:355:GLY:CA	2.65	0.79
3:L:409:ILE:HD13	3:N:297:ALA:O	1.83	0.79
3:L:609:ARG:HD2	2:M:29:THR:CG2	2.13	0.79
3:P:564:LEU:HD21	3:P:568:ARG:NH2	1.98	0.79

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:R:287:PRO:O	3:R:471:LEU:HD12	1.83	0.79
6:Y:1025:ASN:CG	6:Y:1026:GLY:H	1.91	0.79
1:h:45:ARG:HG3	3:H:400:ALA:C	2.08	0.79
1:p:62:VAL:O	1:p:63:LYS:C	2.25	0.79
2:C:29:THR:CG2	3:H:609:ARG:HD2	2.13	0.79
3:F:139:THR:O	3:F:142:ILE:CG2	2.30	0.79
3:F:469:VAL:HG23	3:H:575:MET:HB2	1.63	0.79
3:J:469:VAL:HG23	3:L:575:MET:HB2	1.63	0.79
3:L:106:ILE:HD11	3:L:152:ARG:HG3	1.65	0.79
3:L:118:TYR:CE1	3:L:130:PRO:HB3	2.18	0.79
3:N:354:SER:HB3	3:N:419:MET:HA	1.64	0.79
3:N:372:ALA:HB3	3:N:444:LEU:HD12	1.62	0.79
3:P:78:MET:HE2	3:T:238:VAL:HG11	1.65	0.79
3:P:106:ILE:HD11	3:P:152:ARG:HG3	1.65	0.79
4:V:232:PHE:HB3	4:V:248:LEU:HD11	1.62	0.79
6:X:483:VAL:HB	6:X:840:ALA:HB1	1.63	0.79
6:Y:415:GLN:HE21	6:Y:662:LEU:HD23	1.48	0.79
1:l:16:ALA:C	1:l:22:LEU:CD2	2.56	0.79
3:B:358:LEU:CG	3:B:417:PHE:HE1	1.96	0.79
3:D:409:ILE:HD13	3:F:297:ALA:O	1.83	0.79
3:F:116:SER:HB3	3:F:133:ASN:N	1.97	0.79
3:F:118:TYR:CE1	3:F:130:PRO:HB3	2.18	0.79
3:J:92:TRP:CZ3	3:J:210:VAL:HG11	2.17	0.79
3:J:481:VAL:HG12	3:J:482:ASP:H	1.47	0.79
3:J:613:LEU:HD11	2:K:34:ILE:HB	1.65	0.79
3:L:343:ALA:HB2	3:L:478:GLY:O	1.83	0.79
3:P:358:LEU:CG	3:P:417:PHE:HE1	1.96	0.79
3:T:564:LEU:HD21	3:T:568:ARG:NH2	1.98	0.79
4:U:86:SER:HB3	4:U:91:THR:HB	1.65	0.79
4:U:224:ASN:HD21	4:U:226:SER:HB3	1.46	0.79
6:X:938:TYR:CE1	6:X:946:VAL:HG21	2.17	0.79
6:Y:409:MET:SD	6:Y:858:VAL:HG21	2.22	0.79
6:Y:862:GLU:HB3	6:Y:957:ILE:HD12	1.64	0.79
1:b:15:ARG:HH11	3:B:406:ASN:CG	1.91	0.78
1:h:22:LEU:CD1	1:h:83:VAL:CG2	2.61	0.78
3:D:238:VAL:HG11	3:F:78:MET:HE2	1.65	0.78
3:L:116:SER:HB3	3:L:133:ASN:N	1.97	0.78
3:L:277:ASP:C	3:L:278:THR:HG22	2.06	0.78
3:L:287:PRO:O	3:L:471:LEU:CD1	2.32	0.78
3:L:291:PRO:CB	3:L:467:THR:CG2	2.55	0.78
3:R:118:TYR:CE1	3:R:130:PRO:HB3	2.18	0.78

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:R:609:ARG:HD2	2:S:29:THR:CG2	2.13	0.78
5:W:733:ALA:HB2	5:W:779:MET:HE1	1.65	0.78
6:X:215:LEU:H	6:X:215:LEU:HD23	1.48	0.78
3:B:287:PRO:O	3:B:471:LEU:CD1	2.32	0.78
3:B:343:ALA:HB2	3:B:478:GLY:O	1.83	0.78
2:C:34:ILE:HB	3:H:613:LEU:HD11	1.65	0.78
3:D:355:GLY:CA	3:F:298:PHE:CE2	2.65	0.78
3:D:613:LEU:HD11	2:E:34:ILE:HB	1.65	0.78
3:F:139:THR:HA	3:F:142:ILE:HG22	1.65	0.78
3:J:46:LYS:HZ1	3:N:237:GLU:CB	1.95	0.78
3:J:287:PRO:O	3:J:471:LEU:CD1	2.31	0.78
3:J:575:MET:HB2	3:N:469:VAL:HG23	1.63	0.78
3:N:118:TYR:CE1	3:N:130:PRO:HB3	2.18	0.78
3:N:343:ALA:HB2	3:N:478:GLY:O	1.83	0.78
3:P:46:LYS:HZ1	3:T:237:GLU:CB	1.95	0.78
3:T:358:LEU:CG	3:T:417:PHE:HE1	1.96	0.78
6:Y:235:LEU:HB2	6:Y:356:ASN:ND2	1.99	0.78
1:b:59:HIS:HA	3:B:398:SER:CA	2.12	0.78
3:L:564:LEU:HD21	3:L:568:ARG:NH2	1.98	0.78
3:P:287:PRO:O	3:P:471:LEU:HD12	1.83	0.78
3:P:297:ALA:O	3:T:409:ILE:HD13	1.82	0.78
3:P:472:ARG:HH12	3:R:530:ARG:CD	1.96	0.78
3:P:609:ARG:HD2	2:Q:29:THR:CG2	2.13	0.78
3:R:116:SER:HB3	3:R:133:ASN:N	1.97	0.78
3:B:118:TYR:CE1	3:B:130:PRO:HB3	2.18	0.78
3:B:564:LEU:HD21	3:B:568:ARG:NH2	1.98	0.78
3:D:564:LEU:HD21	3:D:568:ARG:NH2	1.98	0.78
3:J:354:SER:HB3	3:J:419:MET:HA	1.64	0.78
3:R:354:SER:HB3	3:R:419:MET:HA	1.64	0.78
4:U:161:VAL:HG21	4:U:410:MET:HG3	1.66	0.78
5:W:618:LEU:HD13	5:W:619:LYS:H	1.46	0.78
6:X:243:ILE:HG13	6:X:1147:PHE:HB2	1.63	0.78
6:X:457:MET:HE1	6:X:923:LEU:HD11	1.64	0.78
6:X:940:GLN:HE22	6:X:944:ASN:HD22	1.31	0.78
1:f:8:GLN:HG2	3:H:302:GLU:HG3	0.80	0.78
1:f:45:ARG:HG3	3:F:400:ALA:C	2.09	0.78
1:n:45:ARG:HG2	3:N:400:ALA:CA	2.00	0.78
1:p:60:ALA:HB2	3:P:322:TYR:OH	1.84	0.78
3:B:354:SER:HB3	3:B:419:MET:HA	1.64	0.78
3:F:291:PRO:CB	3:F:467:THR:CG2	2.55	0.78
3:N:287:PRO:O	3:N:471:LEU:HD12	1.83	0.78

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:P:238:VAL:CG2	3:R:78:MET:CE	2.59	0.78
3:P:566:TRP:CE2	3:R:195:PRO:HD3	2.19	0.78
3:R:566:TRP:CE2	3:T:195:PRO:HD3	2.19	0.78
6:X:196:GLU:HB3	6:X:313:LYS:HB2	1.63	0.78
1:b:4:HIS:CE1	1:b:54:THR:HG22	2.18	0.78
3:H:294:TYR:HH	3:H:502:LEU:HA	1.47	0.78
3:H:564:LEU:HD21	3:H:568:ARG:NH2	1.98	0.78
3:N:116:SER:HB3	3:N:133:ASN:N	1.97	0.78
3:P:287:PRO:O	3:P:471:LEU:CD1	2.32	0.78
3:R:357:ILE:HG22	3:R:359:PRO:HD3	1.66	0.78
3:R:564:LEU:HD21	3:R:568:ARG:NH2	1.98	0.78
5:W:457:LEU:HB2	5:W:625:ARG:HB2	1.65	0.78
1:f:15:ARG:NH1	3:F:402:ALA:CB	2.45	0.78
3:D:195:PRO:HD3	3:H:566:TRP:CE2	2.19	0.78
3:D:287:PRO:O	3:D:471:LEU:CD1	2.31	0.78
3:F:287:PRO:O	3:F:471:LEU:HD12	1.83	0.78
3:F:343:ALA:HB2	3:F:478:GLY:O	1.83	0.78
3:F:354:SER:HB3	3:F:419:MET:HA	1.64	0.78
3:F:609:ARG:HD2	2:G:29:THR:CG2	2.13	0.78
3:H:481:VAL:HG12	3:H:482:ASP:H	1.47	0.78
3:J:287:PRO:O	3:J:471:LEU:HD12	1.83	0.78
3:P:343:ALA:HB2	3:P:478:GLY:O	1.83	0.78
5:W:341:MET:HE2	5:W:341:MET:HA	1.66	0.78
6:X:241:ASN:O	6:X:1147:PHE:HB3	1.83	0.78
6:X:773:ARG:O	6:X:774:LEU:HB2	1.83	0.78
6:X:780:ILE:HD12	6:X:938:TYR:CD2	2.18	0.78
1:r:44:GLY:O	3:R:399:ALA:N	2.17	0.78
3:D:58:ILE:HD12	3:D:58:ILE:N	1.98	0.78
3:D:358:LEU:CG	3:D:417:PHE:HE1	1.96	0.78
3:D:530:ARG:HE	3:H:472:ARG:HH21	1.29	0.78
3:H:287:PRO:O	3:H:471:LEU:HD12	1.83	0.78
3:H:358:LEU:CG	3:H:417:PHE:HE1	1.96	0.78
3:J:566:TRP:CE2	3:L:195:PRO:HD3	2.19	0.78
3:R:358:LEU:CG	3:R:417:PHE:HE1	1.96	0.78
6:X:247:THR:HG22	6:X:308:ASN:OD1	1.83	0.78
6:Y:765:PRO:HD3	6:Y:853:ARG:HD2	1.65	0.78
6:Y:1210:VAL:HG22	6:Y:1211:ARG:H	1.47	0.78
1:b:59:HIS:CB	3:B:398:SER:HB2	2.14	0.78
3:B:287:PRO:O	3:B:471:LEU:HD12	1.83	0.78
3:D:287:PRO:O	3:D:471:LEU:HD12	1.83	0.78
3:F:481:VAL:HG12	3:F:482:ASP:H	1.47	0.78

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:J:238:VAL:HG11	3:L:78:MET:HE2	1.65	0.78
2:K:28:SER:CB	3:L:216:PRO:HD2	2.14	0.78
3:R:189:PHE:CE1	3:R:193:ILE:HD11	2.19	0.78
5:W:398:LEU:HD11	5:W:788:GLN:NE2	1.99	0.78
5:W:1154:ARG:CD	5:W:1189:ASN:HD22	1.96	0.78
6:X:315:SER:HB2	6:X:331:ARG:HD2	1.66	0.78
6:X:1091:GLN:NE2	6:X:1127:CYS:HB2	1.99	0.78
6:Y:202:THR:CG2	6:Y:391:CYS:HA	2.14	0.78
1:l:59:HIS:HB3	3:L:423:GLN:O	1.84	0.78
1:t:15:ARG:NH1	3:T:406:ASN:CB	2.42	0.78
3:D:566:TRP:CE2	3:F:195:PRO:HD3	2.19	0.78
3:H:287:PRO:O	3:H:471:LEU:CD1	2.32	0.78
3:J:49:ARG:NH1	3:J:115:THR:HG22	1.98	0.78
3:L:189:PHE:CE1	3:L:193:ILE:HD11	2.19	0.78
3:N:358:LEU:CG	3:N:417:PHE:HE1	1.96	0.78
3:P:118:TYR:CE1	3:P:130:PRO:HB3	2.18	0.78
3:P:127:LEU:HD22	3:P:127:LEU:O	1.84	0.78
3:P:189:PHE:CE1	3:P:193:ILE:HD11	2.19	0.78
3:P:291:PRO:CB	3:P:467:THR:CG2	2.56	0.78
3:P:308:ALA:HB1	3:P:383:GLN:HE21	1.49	0.78
2:S:28:SER:CB	3:T:216:PRO:HD2	2.14	0.78
3:T:358:LEU:HB2	3:T:417:PHE:CD1	2.19	0.78
4:V:331:VAL:HB	4:V:332:PRO:HD3	1.66	0.78
5:W:1161:TYR:HE1	5:W:1218:ARG:HH11	1.32	0.78
1:h:59:HIS:CG	1:h:60:ALA:H	2.01	0.77
3:F:287:PRO:O	3:F:471:LEU:CD1	2.32	0.77
3:F:357:ILE:HG22	3:F:359:PRO:HD3	1.66	0.77
3:F:564:LEU:HD21	3:F:568:ARG:NH2	1.98	0.77
3:J:106:ILE:HD11	3:J:152:ARG:HG3	1.65	0.77
3:L:238:VAL:HG11	3:N:78:MET:HE2	1.65	0.77
3:L:274:ILE:CD1	3:L:514:LEU:HD22	2.10	0.77
2:M:28:SER:CB	3:N:216:PRO:HD2	2.14	0.77
3:N:308:ALA:HB1	3:N:383:GLN:HE21	1.49	0.77
3:N:358:LEU:HB2	3:N:417:PHE:CD1	2.20	0.77
4:V:184:HIS:NE2	6:Y:436:LEU:CD1	2.47	0.77
5:W:312:LEU:HD11	5:W:352:TYR:HB2	1.66	0.77
6:X:979:ARG:HG3	6:X:980:GLY:N	1.97	0.77
6:Y:353:GLN:OE1	6:Y:353:GLN:HA	1.83	0.77
3:B:106:ILE:HD11	3:B:152:ARG:HG3	1.65	0.77
3:B:137:ILE:HG12	3:B:142:ILE:HG13	1.66	0.77
3:D:358:LEU:HB2	3:D:417:PHE:CD1	2.20	0.77

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:F:566:TRP:CE2	3:H:195:PRO:HD3	2.19	0.77
2:I:28:SER:CB	3:J:216:PRO:HD2	2.14	0.77
3:L:354:SER:HB3	3:L:419:MET:HA	1.64	0.77
3:N:92:TRP:CZ3	3:N:210:VAL:HG11	2.17	0.77
3:P:469:VAL:HG23	3:R:575:MET:HB2	1.63	0.77
4:V:184:HIS:NE2	6:Y:436:LEU:HD13	1.98	0.77
5:W:949:ILE:HD11	5:W:1022:LEU:HD11	1.66	0.77
6:X:370:ARG:C	6:X:371:LEU:HD12	2.09	0.77
1:p:6:ILE:CD1	3:R:586:VAL:CG2	2.47	0.77
1:r:45:ARG:HG3	3:R:400:ALA:C	2.09	0.77
1:t:5:MET:HE2	1:t:5:MET:CA	2.14	0.77
3:B:49:ARG:HB2	3:B:64:VAL:HG22	1.66	0.77
7:C:101:MYR:C2	3:D:191:LYS:NZ	2.48	0.77
3:F:106:ILE:HD11	3:F:152:ARG:HG3	1.65	0.77
3:H:343:ALA:HB2	3:H:478:GLY:O	1.83	0.77
7:I:101:MYR:H143	7:I:101:MYR:C10	2.11	0.77
7:I:101:MYR:C2	3:J:191:LYS:NZ	2.48	0.77
3:J:277:ASP:C	3:J:278:THR:CG2	2.58	0.77
3:J:308:ALA:HB1	3:J:383:GLN:HE21	1.49	0.77
3:L:566:TRP:CE2	3:N:195:PRO:HD3	2.19	0.77
3:N:189:PHE:CE1	3:N:193:ILE:HD11	2.19	0.77
3:N:287:PRO:O	3:N:471:LEU:CD1	2.32	0.77
2:O:34:ILE:HB	3:T:613:LEU:HD11	1.65	0.77
3:P:137:ILE:HG12	3:P:142:ILE:HG13	1.66	0.77
3:T:357:ILE:HG22	3:T:359:PRO:HD3	1.66	0.77
4:U:222:ARG:HD3	4:U:408:GLN:HG2	1.65	0.77
6:X:1149:THR:HG21	6:X:1161:ASN:CA	2.11	0.77
1:l:45:ARG:HG2	3:L:400:ALA:CA	2.04	0.77
1:f:53:LEU:CD1	1:f:74:GLN:NE2	2.48	0.77
3:D:443:PHE:CE2	3:D:459:LEU:HD21	2.20	0.77
3:F:443:PHE:CE2	3:F:459:LEU:HD21	2.20	0.77
3:J:189:PHE:CE1	3:J:193:ILE:HD11	2.19	0.77
3:J:237:GLU:CB	3:L:46:LYS:HZ1	1.94	0.77
3:L:443:PHE:CE2	3:L:459:LEU:HD21	2.20	0.77
3:P:238:VAL:HG11	3:R:78:MET:HE2	1.65	0.77
1:l:16:ALA:C	1:l:22:LEU:HD21	2.09	0.77
1:p:6:ILE:HD11	3:R:586:VAL:HG21	0.82	0.77
3:B:57:THR:HG22	3:B:58:ILE:H	1.50	0.77
3:B:443:PHE:CE2	3:B:459:LEU:HD21	2.20	0.77
3:D:298:PHE:CE1	3:H:416:ARG:HB2	2.20	0.77
3:F:308:ALA:HB1	3:F:383:GLN:HE21	1.49	0.77

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:H:354:SER:HB3	3:H:419:MET:HA	1.64	0.77
3:H:443:PHE:CE2	3:H:459:LEU:HD21	2.20	0.77
3:J:564:LEU:HD21	3:J:568:ARG:NH2	1.98	0.77
3:P:354:SER:HB3	3:P:419:MET:HA	1.64	0.77
3:P:409:ILE:HD13	3:R:297:ALA:O	1.82	0.77
3:R:287:PRO:O	3:R:471:LEU:CD1	2.32	0.77
3:T:277:ASP:C	3:T:278:THR:CG2	2.58	0.77
6:X:763:VAL:HA	6:X:853:ARG:NH2	1.98	0.77
6:Y:1150:ASN:ND2	6:Y:1156:THR:HB	1.99	0.77
7:C:101:MYR:H143	7:C:101:MYR:C10	2.11	0.77
3:D:189:PHE:CE1	3:D:193:ILE:HD11	2.19	0.77
2:G:28:SER:CB	3:H:216:PRO:HD2	2.14	0.77
3:L:416:ARG:HB2	3:N:298:PHE:CE1	2.20	0.77
3:T:287:PRO:O	3:T:471:LEU:CD1	2.32	0.77
6:X:757:VAL:HG13	6:X:845:PHE:HE2	1.48	0.77
1:l:22:LEU:HD13	1:l:42:THR:O	1.84	0.77
1:h:58:PRO:HB2	3:H:423:GLN:HG2	1.67	0.77
1:j:23:THR:OG1	3:J:323:SER:OG	2.02	0.77
2:A:28:SER:CB	3:B:216:PRO:HD2	2.14	0.77
3:B:277:ASP:C	3:B:278:THR:CG2	2.58	0.77
3:J:358:LEU:HB2	3:J:417:PHE:CD1	2.20	0.77
3:L:277:ASP:C	3:L:278:THR:CG2	2.58	0.77
3:L:358:LEU:HB2	3:L:417:PHE:CD1	2.20	0.77
3:L:481:VAL:HG12	3:L:482:ASP:H	1.47	0.77
3:N:106:ILE:HD11	3:N:152:ARG:HG3	1.65	0.77
3:N:443:PHE:CE2	3:N:459:LEU:HD21	2.20	0.77
3:N:564:LEU:HD21	3:N:568:ARG:NH2	1.98	0.77
2:O:28:SER:CB	3:P:216:PRO:HD2	2.14	0.77
3:P:49:ARG:HB2	3:P:64:VAL:HG22	1.66	0.77
3:P:87:PHE:HZ	3:P:235:LEU:HD12	1.50	0.77
3:P:195:PRO:HD3	3:T:566:TRP:CE2	2.19	0.77
3:R:137:ILE:HG12	3:R:142:ILE:HG13	1.67	0.77
3:R:308:ALA:HB1	3:R:383:GLN:HE21	1.50	0.77
3:R:443:PHE:CE2	3:R:459:LEU:HD21	2.20	0.77
3:R:481:VAL:HG12	3:R:482:ASP:H	1.47	0.77
3:T:118:TYR:CE1	3:T:130:PRO:HB3	2.18	0.77
4:U:198:GLN:HE21	6:X:662:LEU:HD13	1.49	0.77
1:l:24:LEU:HD21	1:l:80:ARG:CZ	2.14	0.77
1:l:45:ARG:CG	3:L:401:GLY:N	2.46	0.77
1:l:59:HIS:CG	3:L:397:VAL:O	2.38	0.77
1:t:59:HIS:HB2	3:T:398:SER:CB	1.89	0.77

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:D:308:ALA:HB1	3:D:383:GLN:HE21	1.49	0.77
3:D:357:ILE:HG22	3:D:359:PRO:HD3	1.66	0.77
3:H:106:ILE:HD11	3:H:152:ARG:HG3	1.65	0.77
3:J:357:ILE:HG22	3:J:359:PRO:HD3	1.66	0.77
3:L:178:SER:CB	2:Q:3:ASN:HD21	1.95	0.77
3:N:357:ILE:HG22	3:N:359:PRO:HD3	1.66	0.77
3:T:116:SER:HB3	3:T:133:ASN:N	1.97	0.77
6:X:433:PRO:HD2	6:X:437:ARG:HH11	1.48	0.77
6:Y:978:ALA:HB2	6:Y:1083:TYR:HD2	1.49	0.77
1:r:59:HIS:CB	3:R:398:SER:CB	2.60	0.77
3:B:358:LEU:HB2	3:B:417:PHE:CD1	2.20	0.77
3:D:472:ARG:HH21	3:F:530:ARG:HE	1.29	0.77
3:F:277:ASP:C	3:F:278:THR:CG2	2.58	0.77
3:H:358:LEU:HB2	3:H:417:PHE:CD1	2.20	0.77
3:J:195:PRO:HD3	3:N:566:TRP:CE2	2.19	0.77
3:P:358:LEU:HB2	3:P:417:PHE:CD1	2.19	0.77
2:Q:28:SER:CB	3:R:216:PRO:HD2	2.14	0.77
3:R:106:ILE:HD11	3:R:152:ARG:HG3	1.65	0.77
3:R:277:ASP:C	3:R:278:THR:CG2	2.58	0.77
4:V:195:GLN:HE22	6:Y:377:ILE:HB	1.48	0.77
1:t:22:LEU:HD22	1:t:22:LEU:O	1.85	0.77
1:t:61:ASN:CB	3:T:397:VAL:HA	2.15	0.77
3:B:87:PHE:HZ	3:B:235:LEU:HD12	1.50	0.77
3:B:189:PHE:CE1	3:B:193:ILE:HD11	2.19	0.77
3:D:317:ARG:HH11	3:D:448:PRO:HB2	1.50	0.77
3:F:358:LEU:HB2	3:F:417:PHE:CE1	2.20	0.77
3:H:49:ARG:HB2	3:H:64:VAL:HG22	1.67	0.77
3:J:116:SER:HB3	3:J:133:ASN:N	1.97	0.77
3:R:416:ARG:HB2	3:T:298:PHE:CE1	2.20	0.77
3:R:519:ASN:O	3:R:523:ASP:HB2	1.85	0.77
3:T:106:ILE:HD11	3:T:152:ARG:HG3	1.65	0.77
3:T:274:ILE:CD1	3:T:514:LEU:HD22	2.10	0.77
3:T:443:PHE:CE2	3:T:459:LEU:HD21	2.20	0.77
6:Y:419:ILE:CD1	6:Y:662:LEU:HB2	2.15	0.77
1:j:15:ARG:CZ	3:J:406:ASN:ND2	2.48	0.76
1:j:23:THR:OG1	3:J:323:SER:CB	2.32	0.76
3:B:519:ASN:O	3:B:523:ASP:HB2	1.85	0.76
3:D:116:SER:HB3	3:D:133:ASN:N	1.97	0.76
3:F:49:ARG:HB2	3:F:64:VAL:HG22	1.66	0.76
3:F:416:ARG:HB2	3:H:298:PHE:CE1	2.20	0.76
3:J:298:PHE:CE1	3:N:416:ARG:HB2	2.20	0.76

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:L:308:ALA:HB1	3:L:383:GLN:HE21	1.49	0.76
3:N:358:LEU:HB2	3:N:417:PHE:CE1	2.20	0.76
3:P:358:LEU:HB2	3:P:417:PHE:CE1	2.20	0.76
3:R:87:PHE:HZ	3:R:235:LEU:HD12	1.50	0.76
6:X:979:ARG:HH11	6:X:979:ARG:HB3	1.50	0.76
6:Y:355:LYS:HZ1	6:Y:1156:THR:CG2	1.95	0.76
1:b:21:ARG:C	3:B:323:SER:OG	2.28	0.76
1:f:45:ARG:HG2	3:F:400:ALA:CA	2.05	0.76
3:B:358:LEU:HB2	3:B:417:PHE:CE1	2.20	0.76
3:F:238:VAL:CG2	3:H:78:MET:CE	2.59	0.76
3:H:137:ILE:HG12	3:H:142:ILE:HG13	1.66	0.76
3:H:341:MET:HA	3:H:341:MET:CE	2.15	0.76
2:I:34:ILE:HB	3:N:613:LEU:HD11	1.65	0.76
3:J:358:LEU:HB2	3:J:417:PHE:CE1	2.20	0.76
7:K:101:MYR:H143	7:K:101:MYR:C10	2.11	0.76
3:L:358:LEU:HB2	3:L:417:PHE:CE1	2.20	0.76
3:N:317:ARG:HH11	3:N:448:PRO:HB2	1.50	0.76
3:P:298:PHE:CE1	3:T:416:ARG:HB2	2.20	0.76
3:P:539:ILE:HD12	3:P:573:ALA:HB2	1.67	0.76
3:R:539:ILE:HD12	3:R:573:ALA:HB2	1.67	0.76
6:Y:309:LEU:HD22	6:Y:402:TYR:HB2	1.68	0.76
6:Y:440:ILE:HG13	6:Y:1202:PRO:HG2	1.66	0.76
6:Y:776:GLN:O	6:Y:779:VAL:HG12	1.85	0.76
1:l:45:ARG:HG3	3:L:400:ALA:C	2.09	0.76
1:b:45:ARG:HG2	3:B:399:ALA:O	1.85	0.76
1:b:73:ARG:NH1	1:b:74:GLN:CD	2.43	0.76
1:d:61:ASN:C	1:d:63:LYS:H	1.94	0.76
1:p:62:VAL:C	1:p:64:THR:H	1.93	0.76
1:r:25:TYR:HE2	1:r:42:THR:HG21	1.50	0.76
1:t:44:GLY:HA3	3:T:322:TYR:HH	1.51	0.76
2:C:28:SER:CB	3:D:216:PRO:HD2	2.14	0.76
3:D:87:PHE:HZ	3:D:235:LEU:HD12	1.50	0.76
3:D:539:ILE:HD12	3:D:573:ALA:HB2	1.67	0.76
3:F:57:THR:HG22	3:F:58:ILE:H	1.50	0.76
3:F:189:PHE:CE1	3:F:193:ILE:HD11	2.19	0.76
3:H:118:TYR:HE1	3:H:130:PRO:HB3	1.51	0.76
3:H:519:ASN:O	3:H:523:ASP:HB2	1.85	0.76
3:J:409:ILE:HD13	3:L:297:ALA:O	1.83	0.76
3:R:358:LEU:HB2	3:R:417:PHE:CE1	2.20	0.76
4:U:106:ILE:HD12	4:U:107:GLN:N	2.00	0.76
6:Y:266:PRO:HB2	6:Y:1085:LEU:HD11	1.66	0.76

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
6:Y:634:GLN:CD	6:Y:634:GLN:H	1.92	0.76
1:n:45:ARG:CG	3:N:399:ALA:O	2.34	0.76
1:p:15:ARG:NH1	3:P:406:ASN:CG	2.41	0.76
3:D:519:ASN:O	3:D:523:ASP:HB2	1.85	0.76
2:G:8:VAL:HG11	3:H:206:GLU:OE2	1.86	0.76
3:H:189:PHE:CE1	3:H:193:ILE:HD11	2.19	0.76
3:J:78:MET:HE2	3:N:238:VAL:HG11	1.65	0.76
3:J:416:ARG:HB2	3:L:298:PHE:CE1	2.20	0.76
7:K:101:MYR:C2	3:L:191:LYS:HZ3	1.92	0.76
3:T:57:THR:HG22	3:T:58:ILE:H	1.50	0.76
3:T:189:PHE:CE1	3:T:193:ILE:HD11	2.19	0.76
3:T:358:LEU:HB2	3:T:417:PHE:CE1	2.20	0.76
4:U:26:LEU:HD11	4:U:90:ILE:HG21	1.66	0.76
4:V:195:GLN:HE22	6:Y:377:ILE:CB	1.98	0.76
6:X:902:MET:HE3	6:X:929:PRO:HG3	1.67	0.76
1:b:17:ALA:N	1:b:22:LEU:HD21	2.01	0.76
3:B:317:ARG:HH11	3:B:448:PRO:HB2	1.50	0.76
7:E:101:MYR:H21	3:F:191:LYS:CE	2.16	0.76
3:H:57:THR:HG22	3:H:58:ILE:H	1.50	0.76
3:J:137:ILE:HG12	3:J:142:ILE:HG13	1.67	0.76
3:J:179:ALA:HB1	3:J:635:GLN:NE2	2.01	0.76
3:L:57:THR:HG22	3:L:58:ILE:H	1.50	0.76
3:R:358:LEU:HB2	3:R:417:PHE:CD1	2.20	0.76
3:T:49:ARG:HB2	3:T:64:VAL:HG22	1.67	0.76
6:X:545:LEU:HD12	6:X:815:PRO:HG3	1.66	0.76
6:Y:207:LEU:HD23	6:Y:207:LEU:O	1.84	0.76
6:Y:385:VAL:HG22	6:Y:386:GLU:N	2.00	0.76
6:Y:626:LEU:HA	6:Y:647:ARG:HH21	1.49	0.76
1:t:59:HIS:HB2	3:T:397:VAL:O	1.86	0.76
3:D:106:ILE:HD11	3:D:152:ARG:HG3	1.66	0.76
3:D:179:ALA:HB1	3:D:635:GLN:HE21	1.51	0.76
3:F:519:ASN:O	3:F:523:ASP:HB2	1.85	0.76
3:J:179:ALA:HB1	3:J:635:GLN:HE21	1.51	0.76
3:J:317:ARG:HH11	3:J:448:PRO:HB2	1.50	0.76
3:N:179:ALA:HB1	3:N:635:GLN:NE2	2.01	0.76
3:N:519:ASN:O	3:N:523:ASP:HB2	1.85	0.76
6:X:1156:THR:HG22	6:X:1161:ASN:HB3	1.68	0.76
6:Y:1101:GLU:HA	6:Y:1104:LEU:HD23	1.67	0.76
1:l:59:HIS:CA	3:L:398:SER:OG	2.33	0.76
1:n:24:LEU:CD2	1:n:80:ARG:HD2	2.15	0.76
1:p:61:ASN:ND2	3:P:395:THR:HG23	1.99	0.76

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:B:118:TYR:HE1	3:B:130:PRO:HB3	1.51	0.76
3:B:179:ALA:HB1	3:B:635:GLN:NE2	2.01	0.76
3:F:139:THR:O	3:F:142:ILE:HG22	1.85	0.76
3:F:139:THR:HA	3:F:142:ILE:CG2	2.15	0.76
3:F:179:ALA:HB1	3:F:635:GLN:HE21	1.51	0.76
3:H:330:GLN:NE2	3:H:332:GLY:HA2	2.01	0.76
3:H:358:LEU:HB2	3:H:417:PHE:CE1	2.20	0.76
3:N:277:ASP:C	3:N:278:THR:CG2	2.58	0.76
3:N:330:GLN:NE2	3:N:332:GLY:HA2	2.01	0.76
7:O:101:MYR:C2	3:P:191:LYS:NZ	2.48	0.76
7:Q:101:MYR:H21	3:R:191:LYS:CE	2.16	0.76
4:U:41:PHE:CE1	6:X:404:MET:HE1	2.21	0.76
4:U:327:ASP:H	4:U:399:GLN:HG3	1.49	0.76
6:X:201:CYS:O	6:X:391:CYS:HB2	1.86	0.76
6:X:1124:VAL:HG12	6:X:1153:SER:OG	1.86	0.76
1:b:44:GLY:O	3:B:399:ALA:CA	2.34	0.76
2:C:8:VAL:HG11	3:D:206:GLU:OE2	1.86	0.76
3:D:58:ILE:H	3:D:58:ILE:CD1	1.99	0.76
3:H:357:ILE:HG22	3:H:359:PRO:HD3	1.66	0.76
3:N:49:ARG:HB2	3:N:64:VAL:HG22	1.67	0.76
3:P:443:PHE:CE2	3:P:459:LEU:HD21	2.20	0.76
5:W:1074:THR:HG22	5:W:1075:PRO:CD	2.16	0.76
6:X:271:VAL:HG11	6:X:1114:PRO:HD3	1.68	0.76
1:f:22:LEU:HG	1:f:83:VAL:CG2	2.16	0.76
1:h:15:ARG:NH2	3:H:402:ALA:CB	2.47	0.76
1:h:59:HIS:HA	3:H:398:SER:OG	1.86	0.76
1:n:45:ARG:HG2	3:N:399:ALA:O	1.85	0.76
1:p:8:GLN:HE22	3:R:303:ASP:CA	1.85	0.76
3:D:56:ALA:HB1	3:D:61:LEU:HD21	1.68	0.76
3:F:179:ALA:HB1	3:F:635:GLN:NE2	2.01	0.76
3:F:358:LEU:HB2	3:F:417:PHE:CD1	2.20	0.76
3:H:87:PHE:HZ	3:H:235:LEU:HD12	1.50	0.76
3:H:179:ALA:HB1	3:H:635:GLN:NE2	2.01	0.76
3:P:179:ALA:HB1	3:P:635:GLN:HE21	1.51	0.76
3:P:317:ARG:HH11	3:P:448:PRO:HB2	1.50	0.76
3:P:519:ASN:O	3:P:523:ASP:HB2	1.85	0.76
3:P:613:LEU:HD11	2:Q:34:ILE:HB	1.65	0.76
3:R:238:VAL:CG2	3:T:78:MET:CE	2.59	0.76
5:W:424:SER:CB	5:W:425:PRO:HD3	2.16	0.76
7:A:101:MYR:C2	3:B:191:LYS:NZ	2.48	0.76
3:B:179:ALA:HB1	3:B:635:GLN:HE21	1.51	0.76

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:B:449:THR:HG22	3:B:450:SER:N	2.01	0.76
3:D:238:VAL:CA	3:F:142:ILE:HD11	2.07	0.76
3:D:277:ASP:C	3:D:278:THR:CG2	2.58	0.76
3:D:416:ARG:HB2	3:F:298:PHE:CE1	2.20	0.76
3:H:317:ARG:HH11	3:H:448:PRO:HB2	1.50	0.76
2:I:8:VAL:HG11	3:J:206:GLU:OE2	1.86	0.76
3:J:87:PHE:HZ	3:J:235:LEU:HD12	1.50	0.76
3:J:443:PHE:CE2	3:J:459:LEU:HD21	2.20	0.76
3:J:530:ARG:HE	3:N:472:ARG:HH21	1.29	0.76
3:L:118:TYR:HE1	3:L:130:PRO:HB3	1.51	0.76
3:L:179:ALA:HB1	3:L:635:GLN:HE21	1.51	0.76
3:P:277:ASP:C	3:P:278:THR:CG2	2.58	0.76
3:R:238:VAL:HG22	3:T:142:ILE:HD11	1.68	0.76
6:X:364:ILE:HG13	6:X:367:ARG:HH21	1.51	0.76
1:n:23:THR:OG1	3:N:323:SER:CA	2.30	0.75
2:E:28:SER:CB	3:F:216:PRO:HD2	2.15	0.75
3:L:357:ILE:HG22	3:L:359:PRO:HD3	1.66	0.75
3:N:179:ALA:HB1	3:N:635:GLN:HE21	1.51	0.75
3:P:416:ARG:HB2	3:R:298:PHE:CE1	2.20	0.75
3:R:49:ARG:HB2	3:R:64:VAL:HG22	1.66	0.75
3:T:179:ALA:HB1	3:T:635:GLN:NE2	2.01	0.75
6:Y:124:ALA:HB2	6:Y:168:ARG:HE	1.51	0.75
6:Y:1211:ARG:HH11	6:Y:1211:ARG:HG2	1.50	0.75
1:p:15:ARG:CZ	3:P:406:ASN:OD1	2.33	0.75
3:D:358:LEU:HB2	3:D:417:PHE:CE1	2.20	0.75
3:F:330:GLN:NE2	3:F:332:GLY:HA2	2.01	0.75
3:H:308:ALA:HB1	3:H:383:GLN:HE21	1.49	0.75
2:K:8:VAL:HG11	3:L:206:GLU:OE2	1.86	0.75
3:L:540:VAL:HG21	3:L:566:TRP:CE3	2.22	0.75
2:M:8:VAL:HG11	3:N:206:GLU:OE2	1.86	0.75
3:R:609:ARG:HB3	2:S:29:THR:CG2	2.14	0.75
2:S:8:VAL:HG11	3:T:206:GLU:OE2	1.86	0.75
3:T:118:TYR:HE1	3:T:130:PRO:HB3	1.51	0.75
4:V:2:ALA:HB2	6:Y:373:ARG:HH12	1.51	0.75
4:V:18:ALA:HB1	4:V:19:PRO:HD2	1.67	0.75
4:V:184:HIS:CD2	6:Y:436:LEU:HD13	2.22	0.75
6:X:1087:TYR:HA	6:X:1119:ILE:O	1.86	0.75
1:d:59:HIS:CA	3:D:398:SER:HB2	2.15	0.75
1:n:15:ARG:HH22	3:N:402:ALA:HB3	1.52	0.75
3:F:317:ARG:HH11	3:F:448:PRO:HB2	1.50	0.75
3:F:540:VAL:HG21	3:F:566:TRP:CE3	2.22	0.75

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:J:525:ILE:HD12	3:J:604:VAL:HG13	1.69	0.75
3:L:519:ASN:O	3:L:523:ASP:HB2	1.85	0.75
3:N:525:ILE:HD12	3:N:604:VAL:HG13	1.69	0.75
3:P:357:ILE:HG22	3:P:359:PRO:HD3	1.66	0.75
3:T:317:ARG:HH11	3:T:448:PRO:HB2	1.51	0.75
4:U:37:TRP:CZ3	4:U:72:LEU:HD11	2.20	0.75
6:X:528:PHE:HE1	6:X:575:MET:CE	1.99	0.75
1:f:24:LEU:HD12	1:f:24:LEU:N	2.01	0.75
1:d:23:THR:HG23	3:D:323:SER:O	1.86	0.75
1:n:59:HIS:CB	3:N:397:VAL:O	2.27	0.75
1:p:58:PRO:O	1:p:59:HIS:CE1	2.39	0.75
1:r:5:MET:HE2	1:r:5:MET:CA	2.13	0.75
3:B:116:SER:HB3	3:B:133:ASN:N	1.97	0.75
3:D:52:GLY:O	3:D:53:THR:CB	2.35	0.75
3:D:179:ALA:HB1	3:D:635:GLN:NE2	2.01	0.75
3:F:449:THR:HG22	3:F:450:SER:N	2.01	0.75
3:H:277:ASP:C	3:H:278:THR:CG2	2.58	0.75
3:J:118:TYR:HE1	3:J:130:PRO:HB3	1.51	0.75
3:J:330:GLN:NE2	3:J:332:GLY:HA2	2.01	0.75
3:J:519:ASN:O	3:J:523:ASP:HB2	1.85	0.75
3:L:87:PHE:HZ	3:L:235:LEU:HD12	1.50	0.75
6:X:208:ASP:HB2	6:X:236:PRO:HB2	1.68	0.75
1:p:24:LEU:HD21	1:p:80:ARG:NE	2.02	0.75
3:B:357:ILE:HG22	3:B:359:PRO:HD3	1.66	0.75
3:D:152:ARG:HH11	3:D:152:ARG:CA	1.98	0.75
3:H:449:THR:HG22	3:H:450:SER:N	2.01	0.75
3:H:525:ILE:HD12	3:H:604:VAL:HG13	1.68	0.75
3:L:179:ALA:HB1	3:L:635:GLN:NE2	2.01	0.75
3:L:525:ILE:HD12	3:L:604:VAL:HG13	1.69	0.75
3:N:449:THR:HG22	3:N:450:SER:N	2.01	0.75
3:P:179:ALA:HB1	3:P:635:GLN:NE2	2.01	0.75
2:Q:8:VAL:HG11	3:R:206:GLU:OE2	1.86	0.75
3:R:330:GLN:NE2	3:R:332:GLY:HA2	2.01	0.75
4:V:175:MET:O	4:V:175:MET:HG3	1.86	0.75
4:V:250:TRP:CE3	6:Y:956:HIS:HE1	2.04	0.75
6:X:194:ILE:HG21	6:X:1201:TYR:CE1	2.21	0.75
6:X:330:ILE:HG21	6:X:347:ARG:HD2	1.69	0.75
6:X:409:MET:O	6:X:443:PRO:HA	1.86	0.75
1:h:23:THR:CB	3:H:323:SER:CB	2.51	0.75
1:h:45:ARG:CD	3:H:400:ALA:HA	2.16	0.75
1:j:15:ARG:NH2	3:J:402:ALA:CB	2.50	0.75

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:B:339:ILE:HD13	3:B:349:LEU:CD2	2.17	0.75
3:F:87:PHE:HZ	3:F:235:LEU:HD12	1.50	0.75
3:F:539:ILE:HD12	3:F:573:ALA:HB2	1.67	0.75
3:H:179:ALA:HB1	3:H:635:GLN:HE21	1.51	0.75
7:K:101:MYR:H21	3:L:191:LYS:CE	2.16	0.75
3:L:330:GLN:NE2	3:L:332:GLY:HA2	2.01	0.75
3:P:330:GLN:NE2	3:P:332:GLY:HA2	2.01	0.75
3:P:339:ILE:HD13	3:P:349:LEU:CD2	2.17	0.75
4:U:160:GLU:OE1	4:U:209:ARG:HD2	1.87	0.75
4:U:331:VAL:HB	4:U:332:PRO:HD3	1.68	0.75
5:W:19:ARG:HA	5:W:282:VAL:O	1.87	0.75
5:W:1040:VAL:CB	5:W:1041:PRO:HD3	2.17	0.75
6:X:780:ILE:HD12	6:X:938:TYR:HD2	1.52	0.75
6:Y:567:LEU:HD23	6:Y:708:PHE:CD1	2.21	0.75
1:b:45:ARG:CG	3:B:399:ALA:O	2.35	0.75
1:f:58:PRO:HB2	3:F:423:GLN:HG2	1.67	0.75
1:h:5:MET:O	1:h:6:ILE:C	2.25	0.75
1:n:6:ILE:HD13	3:J:586:VAL:HG21	1.66	0.75
1:n:23:THR:O	1:n:23:THR:CG2	2.34	0.75
1:r:45:ARG:CA	3:R:399:ALA:O	2.34	0.75
1:t:59:HIS:CG	3:T:397:VAL:O	2.39	0.75
3:B:330:GLN:NE2	3:B:332:GLY:HA2	2.01	0.75
3:B:375:LEU:HD23	3:B:437:ILE:CD1	2.15	0.75
3:B:525:ILE:HD12	3:B:604:VAL:HG13	1.69	0.75
3:D:330:GLN:NE2	3:D:332:GLY:HA2	2.01	0.75
2:E:8:VAL:HG11	3:F:206:GLU:OE2	1.86	0.75
3:J:540:VAL:HG21	3:J:566:TRP:CE3	2.22	0.75
3:L:291:PRO:HB3	3:L:467:THR:CG2	2.16	0.75
3:L:449:THR:HG22	3:L:450:SER:N	2.01	0.75
3:N:341:MET:HA	3:N:341:MET:CE	2.15	0.75
2:O:8:VAL:HG11	3:P:206:GLU:OE2	1.86	0.75
3:T:308:ALA:HB1	3:T:383:GLN:HE21	1.49	0.75
3:T:330:GLN:NE2	3:T:332:GLY:HA2	2.01	0.75
3:T:339:ILE:HD13	3:T:349:LEU:CD2	2.17	0.75
3:T:540:VAL:HG21	3:T:566:TRP:CE3	2.22	0.75
4:U:158:HIS:HE1	4:U:160:GLU:HG3	1.51	0.75
5:W:914:VAL:HG12	5:W:918:ILE:HG13	1.68	0.75
6:X:232:THR:HG21	6:X:338:GLU:OE2	1.85	0.75
6:Y:212:SER:HB2	6:Y:238:GLY:HA2	1.67	0.75
6:Y:978:ALA:HB2	6:Y:1083:TYR:CD2	2.21	0.75
1:l:15:ARG:NH2	3:L:402:ALA:HB3	2.02	0.75

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:l:60:ALA:N	3:L:398:SER:HA	1.97	0.75
1:b:23:THR:CG2	1:b:25:TYR:HE2	1.98	0.75
1:j:44:GLY:O	3:J:399:ALA:CA	2.34	0.75
3:B:308:ALA:HB1	3:B:383:GLN:HE21	1.49	0.75
3:D:53:THR:O	3:D:54:SER:OG	2.03	0.75
3:H:540:VAL:HG21	3:H:566:TRP:CE3	2.22	0.75
3:J:469:VAL:CG2	3:L:575:MET:CE	2.43	0.75
3:L:49:ARG:HB2	3:L:64:VAL:HG22	1.66	0.75
3:N:339:ILE:HD13	3:N:349:LEU:CD2	2.17	0.75
3:P:525:ILE:HD12	3:P:604:VAL:HG13	1.69	0.75
2:Q:13:ILE:HD11	7:Q:101:MYR:H111	1.69	0.75
3:R:274:ILE:CD1	3:R:514:LEU:HD22	2.10	0.75
3:T:411:VAL:O	3:T:412:ARG:HB2	1.87	0.75
4:V:192:PHE:CB	6:Y:376:LEU:CD2	2.64	0.75
6:Y:714:SER:OG	6:Y:717:MET:HG2	1.85	0.75
1:b:23:THR:HG21	1:b:25:TYR:HE2	1.52	0.75
1:d:4:HIS:HD2	1:d:54:THR:HG22	1.43	0.75
1:d:23:THR:HG21	3:D:323:SER:HA	1.69	0.75
2:A:8:VAL:HG11	3:B:206:GLU:OE2	1.86	0.75
3:D:525:ILE:HD12	3:D:604:VAL:HG13	1.69	0.75
3:H:539:ILE:HD12	3:H:573:ALA:HB2	1.67	0.75
3:J:238:VAL:CG2	3:L:78:MET:CE	2.59	0.75
3:J:271:LEU:HD11	3:J:516:LYS:HB2	1.69	0.75
3:L:317:ARG:HH11	3:L:448:PRO:HB2	1.50	0.75
3:L:339:ILE:HD13	3:L:349:LEU:CD2	2.17	0.75
3:T:102:GLY:O	5:W:407:ALA:HB2	1.87	0.75
5:W:676:ARG:HB2	5:W:1081:VAL:HG11	1.68	0.75
1:n:45:ARG:CD	3:N:401:GLY:H	2.01	0.74
1:p:62:VAL:HG12	1:p:65:ILE:CD1	2.15	0.74
1:t:15:ARG:CZ	3:T:402:ALA:HB1	2.10	0.74
1:t:21:ARG:HH12	3:T:327:TYR:HE1	1.34	0.74
2:C:33:ALA:CB	3:H:260:SER:HB3	2.17	0.74
3:F:291:PRO:HB3	3:F:467:THR:CG2	2.17	0.74
3:J:449:THR:HG22	3:J:450:SER:N	2.01	0.74
3:N:469:VAL:O	3:N:470:LEU:HB2	1.87	0.74
3:T:519:ASN:O	3:T:523:ASP:HB2	1.85	0.74
4:U:241:LEU:C	6:X:1179:VAL:CG1	2.61	0.74
5:W:1233:ARG:O	5:W:1291:ALA:HB1	1.86	0.74
6:X:709:ILE:HD11	6:X:730:THR:HG23	1.67	0.74
1:l:21:ARG:NE	3:L:327:TYR:OH	2.20	0.74
1:l:45:ARG:HG2	3:L:399:ALA:O	1.88	0.74

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:h:22:LEU:HG	1:h:83:VAL:CG2	2.16	0.74
3:B:539:ILE:HD12	3:B:573:ALA:HB2	1.67	0.74
3:J:341:MET:HA	3:J:341:MET:CE	2.15	0.74
3:N:118:TYR:HE1	3:N:130:PRO:HB3	1.51	0.74
3:N:161:LYS:C	3:N:161:LYS:CD	2.54	0.74
3:N:291:PRO:HB3	3:N:467:THR:CG2	2.17	0.74
3:R:271:LEU:HD11	3:R:516:LYS:HB2	1.69	0.74
3:R:525:ILE:HD12	3:R:604:VAL:HG13	1.69	0.74
5:W:517:ALA:HB2	5:W:575:PHE:CD2	2.22	0.74
5:W:1206:TYR:CE2	5:W:1222:PRO:HB3	2.21	0.74
6:X:497:ILE:HD13	6:X:514:ILE:HD12	1.68	0.74
6:X:1017:PHE:HD1	6:Y:1210:VAL:O	1.70	0.74
6:X:1052:LEU:HD13	6:X:1052:LEU:O	1.87	0.74
6:Y:300:MET:HE1	6:Y:975:PHE:CD2	2.22	0.74
6:Y:1033:ASP:HB2	6:Y:1039:ARG:HG2	1.68	0.74
1:h:45:ARG:CG	3:H:401:GLY:N	2.45	0.74
3:D:449:THR:HG22	3:D:450:SER:N	2.01	0.74
3:F:469:VAL:O	3:F:470:LEU:HB2	1.87	0.74
2:O:33:ALA:CB	3:T:260:SER:HB3	2.17	0.74
3:P:118:TYR:HE1	3:P:130:PRO:HB3	1.51	0.74
3:R:179:ALA:HB1	3:R:635:GLN:NE2	2.01	0.74
3:R:540:VAL:HG21	3:R:566:TRP:CE3	2.22	0.74
4:U:35:ARG:CD	6:X:449:GLU:HG2	2.18	0.74
5:W:145:ASP:O	5:W:148:ILE:HG22	1.87	0.74
5:W:400:LEU:O	5:W:400:LEU:HD23	1.87	0.74
6:X:559:HIS:HD2	6:X:562:SER:OG	1.69	0.74
6:X:655:GLU:CD	6:X:655:GLU:N	2.43	0.74
1:b:59:HIS:HE1	1:b:64:THR:OG1	1.69	0.74
1:r:23:THR:HG22	1:r:25:TYR:CZ	2.22	0.74
3:B:411:VAL:O	3:B:412:ARG:HB2	1.87	0.74
3:B:540:VAL:HG21	3:B:566:TRP:CE3	2.22	0.74
3:D:540:VAL:HG21	3:D:566:TRP:CE3	2.22	0.74
3:F:375:LEU:HD23	3:F:437:ILE:CD1	2.15	0.74
3:F:428:ARG:HG2	3:F:428:ARG:NH1	2.03	0.74
3:L:92:TRP:NE1	3:L:156:VAL:HG21	2.03	0.74
3:N:57:THR:HG22	3:N:58:ILE:H	1.50	0.74
3:R:57:THR:HG22	3:R:58:ILE:H	1.50	0.74
3:R:179:ALA:HB1	3:R:635:GLN:HE21	1.51	0.74
3:T:179:ALA:HB1	3:T:635:GLN:HE21	1.51	0.74
6:X:226:MET:HB3	6:X:231:MET:O	1.87	0.74
6:X:864:PHE:CE2	6:X:952:PRO:HG3	2.22	0.74

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:l:15:ARG:NH1	3:L:406:ASN:CG	2.43	0.74
1:p:44:GLY:HA3	3:P:322:TYR:CZ	2.21	0.74
3:D:50:PRO:CG	3:D:56:ALA:HB2	2.17	0.74
3:H:411:VAL:O	3:H:412:ARG:HB2	1.87	0.74
3:H:469:VAL:O	3:H:470:LEU:HB2	1.87	0.74
3:J:354:SER:HB2	3:J:419:MET:HA	1.70	0.74
3:L:539:ILE:HD12	3:L:573:ALA:HB2	1.67	0.74
3:P:46:LYS:HZ3	3:T:237:GLU:CD	1.96	0.74
3:P:260:SER:HB3	2:Q:33:ALA:CB	2.17	0.74
3:R:339:ILE:HD13	3:R:349:LEU:CD2	2.17	0.74
3:R:449:THR:HG22	3:R:450:SER:N	2.01	0.74
7:S:101:MYR:H143	7:S:101:MYR:C10	2.12	0.74
3:T:539:ILE:HD12	3:T:573:ALA:HB2	1.67	0.74
5:W:971:VAL:CG1	5:W:972:PRO:HD3	2.10	0.74
6:X:801:ARG:HD3	6:X:939:LEU:CD1	2.18	0.74
6:Y:231:MET:HE3	6:Y:347:ARG:HH12	1.51	0.74
1:b:16:ALA:O	1:b:22:LEU:HD21	1.79	0.74
3:D:339:ILE:HD13	3:D:349:LEU:CD2	2.17	0.74
3:F:435:PRO:HG2	3:N:436:TYR:CZ	2.22	0.74
7:G:101:MYR:H143	7:G:101:MYR:C10	2.12	0.74
2:I:33:ALA:CB	3:N:260:SER:HB3	2.17	0.74
3:J:260:SER:HB3	2:K:33:ALA:CB	2.17	0.74
3:J:411:VAL:O	3:J:412:ARG:HB2	1.87	0.74
3:T:271:LEU:HD11	3:T:516:LYS:HB2	1.69	0.74
5:W:394:ALA:HB3	5:W:792:THR:HG22	1.67	0.74
5:W:1274:ALA:HB3	5:W:1286:THR:HG23	1.69	0.74
6:X:462:ILE:HD11	6:X:468:ALA:O	1.87	0.74
6:X:827:THR:CG2	6:X:828:GLU:H	1.98	0.74
1:p:44:GLY:HA3	3:P:322:TYR:HH	1.51	0.74
3:D:92:TRP:NE1	3:D:156:VAL:HG21	2.03	0.74
2:K:13:ILE:HD11	7:K:101:MYR:H111	1.69	0.74
3:L:138:SER:O	3:L:142:ILE:HG23	1.87	0.74
3:N:92:TRP:NE1	3:N:156:VAL:HG21	2.03	0.74
3:R:469:VAL:O	3:R:470:LEU:HB2	1.87	0.74
5:W:313:LEU:CD1	5:W:318:VAL:HG11	2.17	0.74
6:X:513:GLU:OE2	6:X:560:LEU:HB3	1.87	0.74
6:Y:203:SER:HB2	6:Y:247:THR:HA	1.67	0.74
1:h:59:HIS:CB	3:H:398:SER:OG	2.34	0.74
1:p:59:HIS:CG	3:P:398:SER:CB	2.60	0.74
3:D:356:THR:HG21	3:H:412:ARG:HH22	1.53	0.74
3:F:525:ILE:HD12	3:F:604:VAL:HG13	1.69	0.74

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:H:354:SER:HB2	3:H:419:MET:HA	1.69	0.74
3:J:469:VAL:O	3:J:470:LEU:HB2	1.88	0.74
3:L:178:SER:HB2	2:Q:3:ASN:ND2	2.01	0.74
3:P:411:VAL:O	3:P:412:ARG:HB2	1.87	0.74
3:P:449:THR:HG22	3:P:450:SER:N	2.01	0.74
3:R:317:ARG:HH11	3:R:448:PRO:HB2	1.51	0.74
3:T:322:TYR:O	3:T:323:SER:OG	2.06	0.74
3:T:449:THR:HG22	3:T:450:SER:N	2.01	0.74
4:U:176:ALA:CB	6:X:413:PRO:HD2	2.17	0.74
4:U:232:PHE:HB3	4:U:248:LEU:HD11	1.69	0.74
6:X:356:ASN:HD21	6:X:1151:ALA:HB1	1.50	0.74
6:X:652:GLN:HE22	6:X:773:ARG:HE	1.34	0.74
1:f:5:MET:HE2	1:f:5:MET:CA	2.15	0.74
1:r:8:GLN:NE2	3:T:303:ASP:OD1	2.19	0.74
1:r:17:ALA:CA	1:r:22:LEU:HD21	2.17	0.74
3:D:260:SER:HB3	2:E:33:ALA:CB	2.17	0.74
3:D:372:ALA:CB	3:D:399:ALA:HA	2.18	0.74
3:D:411:VAL:O	3:D:412:ARG:HB2	1.87	0.74
3:F:609:ARG:HB3	2:G:29:THR:CG2	2.14	0.74
3:H:92:TRP:NE1	3:H:156:VAL:HG21	2.03	0.74
3:H:339:ILE:HD13	3:H:349:LEU:CD2	2.17	0.74
3:J:339:ILE:HD13	3:J:349:LEU:CD2	2.17	0.74
3:J:372:ALA:CB	3:J:399:ALA:HA	2.18	0.74
3:J:539:ILE:HD12	3:J:573:ALA:HB2	1.67	0.74
3:N:87:PHE:HZ	3:N:235:LEU:HD12	1.50	0.74
3:N:539:ILE:HD12	3:N:573:ALA:HB2	1.67	0.74
3:P:637:PHE:HE1	3:T:545:VAL:CG2	2.01	0.74
6:X:1130:ALA:CB	6:Y:104:GLN:HB3	2.17	0.74
6:Y:413:PRO:HB2	6:Y:434:ILE:HD13	1.68	0.74
1:l:4:HIS:CE1	1:l:54:THR:HG22	2.22	0.74
3:D:69:GLY:HA3	3:D:139:THR:HB	1.70	0.74
3:D:271:LEU:HD11	3:D:516:LYS:HB2	1.69	0.74
3:L:238:VAL:CG2	3:N:78:MET:CE	2.59	0.74
3:N:322:TYR:O	3:N:323:SER:OG	2.06	0.74
3:P:412:ARG:HH22	3:R:356:THR:HG21	1.53	0.74
3:P:540:VAL:HG21	3:P:566:TRP:CE3	2.22	0.74
3:R:372:ALA:CB	3:R:399:ALA:HA	2.18	0.74
3:R:472:ARG:NH2	3:T:530:ARG:NE	2.32	0.74
4:U:52:PRO:HB3	6:X:963:GLN:NE2	2.03	0.74
6:X:957:ILE:C	6:X:959:ARG:H	1.96	0.74
6:Y:322:THR:HG21	6:Y:348:ALA:HB2	1.70	0.74

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:l:45:ARG:CD	3:L:401:GLY:H	2.01	0.73
1:b:15:ARG:NH2	3:B:402:ALA:HB2	2.03	0.73
1:b:23:THR:HB	3:B:323:SER:CB	2.18	0.73
3:B:341:MET:HA	3:B:341:MET:CE	2.15	0.73
3:D:121:ALA:CB	3:H:87:PHE:CE1	2.71	0.73
3:D:469:VAL:O	3:D:470:LEU:HB2	1.87	0.73
3:F:118:TYR:HE1	3:F:130:PRO:HB3	1.51	0.73
3:F:382:ILE:HA	3:F:385:VAL:HG12	1.70	0.73
2:I:29:THR:CG2	3:N:609:ARG:HB3	2.14	0.73
3:J:87:PHE:CE1	3:L:121:ALA:CB	2.71	0.73
3:J:412:ARG:HH22	3:L:356:THR:HG21	1.53	0.73
3:L:372:ALA:CB	3:L:399:ALA:HA	2.18	0.73
3:L:469:VAL:O	3:L:470:LEU:HB2	1.87	0.73
2:M:13:ILE:HD11	7:M:101:MYR:H111	1.70	0.73
3:N:372:ALA:CB	3:N:399:ALA:HA	2.18	0.73
7:S:101:MYR:H142	7:S:101:MYR:C10	2.11	0.73
5:W:736:VAL:HG21	5:W:780:LEU:CD2	2.18	0.73
5:W:1038:ASN:HD22	5:W:1049:ALA:CB	1.99	0.73
6:X:606:MET:HE3	6:Y:718:THR:HG21	1.67	0.73
6:Y:985:VAL:HG12	6:Y:1073:ARG:HB3	1.68	0.73
1:r:16:ALA:O	1:r:22:LEU:HD22	1.86	0.73
3:D:87:PHE:CE1	3:F:121:ALA:CB	2.71	0.73
3:D:354:SER:HB2	3:D:419:MET:HA	1.70	0.73
3:J:46:LYS:HZ3	3:N:237:GLU:CD	1.96	0.73
3:J:58:ILE:CG2	3:J:124:GLY:HA2	2.18	0.73
3:J:121:ALA:CB	3:N:87:PHE:CE1	2.71	0.73
7:K:101:MYR:H142	7:K:101:MYR:C10	2.10	0.73
3:P:469:VAL:O	3:P:470:LEU:HB2	1.88	0.73
2:Q:9:ASN:HD21	3:R:206:GLU:HA	1.53	0.73
3:R:92:TRP:NE1	3:R:156:VAL:HG21	2.03	0.73
3:T:87:PHE:HZ	3:T:235:LEU:HD12	1.50	0.73
3:T:291:PRO:HB3	3:T:467:THR:CG2	2.16	0.73
4:U:251:ASN:CG	4:U:252:ASP:H	1.94	0.73
5:W:157:GLN:O	5:W:157:GLN:HG2	1.86	0.73
5:W:1078:VAL:HG21	5:W:1087:LEU:HB3	1.68	0.73
1:p:8:GLN:NE2	3:R:302:GLU:CG	2.47	0.73
3:B:372:ALA:CB	3:B:399:ALA:HA	2.18	0.73
3:D:118:TYR:HE1	3:D:130:PRO:HB3	1.51	0.73
3:D:382:ILE:HA	3:D:385:VAL:HG12	1.70	0.73
2:E:13:ILE:HD11	7:E:101:MYR:H111	1.69	0.73
3:F:372:ALA:CB	3:F:399:ALA:HA	2.18	0.73

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:F:545:VAL:CG2	3:H:637:PHE:HE1	2.01	0.73
3:H:382:ILE:HA	3:H:385:VAL:HG12	1.70	0.73
3:J:49:ARG:HG3	3:J:64:VAL:HG21	1.70	0.73
3:L:87:PHE:CE1	3:N:121:ALA:CB	2.71	0.73
3:N:540:VAL:HG21	3:N:566:TRP:CE3	2.22	0.73
3:R:341:MET:HA	3:R:341:MET:CE	2.15	0.73
6:Y:202:THR:HG22	6:Y:391:CYS:HA	1.71	0.73
6:Y:486:LEU:HD22	6:Y:488:LEU:HD12	1.70	0.73
1:b:44:GLY:HA3	3:B:322:TYR:HH	1.51	0.73
1:p:15:ARG:NE	3:P:406:ASN:OD1	2.20	0.73
1:p:23:THR:CG2	1:p:25:TYR:CE2	2.71	0.73
3:D:78:MET:CE	3:H:238:VAL:CG2	2.59	0.73
3:F:48:TRP:CH2	3:F:122:THR:HA	2.24	0.73
3:H:271:LEU:HD11	3:H:516:LYS:HB2	1.69	0.73
3:L:271:LEU:HD11	3:L:516:LYS:HB2	1.69	0.73
3:P:356:THR:HG21	3:T:412:ARG:HH22	1.53	0.73
3:R:118:TYR:HE1	3:R:130:PRO:HB3	1.51	0.73
3:T:525:ILE:HD12	3:T:604:VAL:HG13	1.69	0.73
1:f:24:LEU:CD2	1:f:80:ARG:HD3	2.15	0.73
1:t:45:ARG:HG2	3:T:399:ALA:O	1.88	0.73
2:A:13:ILE:HD11	7:A:101:MYR:H111	1.70	0.73
3:D:164:ASP:OD2	3:H:534:SER:HB3	1.89	0.73
3:F:339:ILE:HD13	3:F:349:LEU:CD2	2.17	0.73
3:L:375:LEU:HD23	3:L:437:ILE:CD1	2.15	0.73
3:L:469:VAL:CG2	3:N:575:MET:CE	2.43	0.73
3:R:48:TRP:CH2	3:R:122:THR:HA	2.24	0.73
5:W:989:HIS:CE1	5:W:1027:HIS:HD1	2.06	0.73
6:Y:374:ALA:HB1	6:Y:377:ILE:CD1	2.16	0.73
1:h:59:HIS:CB	3:H:397:VAL:O	2.36	0.73
1:j:20:GLY:CA	1:j:83:VAL:HG13	2.17	0.73
1:n:45:ARG:CG	3:N:401:GLY:N	2.47	0.73
3:B:271:LEU:HD11	3:B:516:LYS:HB2	1.69	0.73
3:D:534:SER:HB3	3:F:164:ASP:OD2	1.89	0.73
3:H:48:TRP:CH2	3:H:122:THR:HA	2.23	0.73
3:H:644:LEU:O	3:H:646:PRO:HD3	1.89	0.73
3:L:411:VAL:O	3:L:412:ARG:HB2	1.87	0.73
3:P:48:TRP:CH2	3:P:122:THR:HA	2.23	0.73
6:X:976:ASN:HA	6:X:1083:TYR:HE1	1.54	0.73
6:Y:122:CYS:SG	6:Y:140:HIS:CE1	2.79	0.73
1:n:8:GLN:CD	3:J:302:GLU:CG	2.60	0.73
1:p:60:ALA:HB3	3:P:322:TYR:CE1	2.24	0.73

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:B:354:SER:HB2	3:B:419:MET:HA	1.69	0.73
3:P:92:TRP:NE1	3:P:156:VAL:HG21	2.03	0.73
3:P:372:ALA:CB	3:P:399:ALA:HA	2.18	0.73
3:R:411:VAL:O	3:R:412:ARG:HB2	1.87	0.73
3:T:92:TRP:NE1	3:T:156:VAL:HG21	2.03	0.73
3:T:372:ALA:CB	3:T:399:ALA:HA	2.18	0.73
4:U:61:VAL:HG12	4:U:276:HIS:HA	1.71	0.73
1:f:3:LEU:HD13	1:f:4:HIS:O	1.88	0.73
1:j:22:LEU:HD13	1:j:22:LEU:C	2.13	0.73
3:B:469:VAL:O	3:B:470:LEU:HB2	1.87	0.73
3:F:644:LEU:O	3:F:646:PRO:HD3	1.89	0.73
3:J:92:TRP:NE1	3:J:156:VAL:HG21	2.03	0.73
3:J:534:SER:HB3	3:L:164:ASP:OD2	1.89	0.73
3:L:644:LEU:O	3:L:646:PRO:HD3	1.89	0.73
3:R:291:PRO:HB3	3:R:467:THR:CG2	2.16	0.73
6:X:996:MET:O	6:X:998:PRO:HD3	1.88	0.73
6:X:1170:PRO:HD2	6:X:1189:THR:CG2	2.19	0.73
1:h:22:LEU:HD12	1:h:83:VAL:CG2	2.18	0.73
3:B:48:TRP:CH2	3:B:122:THR:HA	2.24	0.73
3:J:48:TRP:CH2	3:J:122:THR:HA	2.24	0.73
3:P:87:PHE:CE1	3:R:121:ALA:CB	2.71	0.73
3:P:291:PRO:HB3	3:P:467:THR:CG2	2.16	0.73
3:R:322:TYR:O	3:R:323:SER:OG	2.06	0.73
3:R:534:SER:HB3	3:T:164:ASP:OD2	1.89	0.73
3:T:368:THR:HG22	3:T:446:PRO:HG3	1.71	0.73
3:T:382:ILE:HA	3:T:385:VAL:HG12	1.70	0.73
4:U:268:ARG:HG2	4:U:327:ASP:OD1	1.89	0.73
5:W:30:LEU:HG	5:W:118:LEU:HD11	1.70	0.73
5:W:558:VAL:HG22	5:W:560:HIS:H	1.54	0.73
6:X:899:ALA:O	6:X:903:HIS:HB2	1.89	0.73
6:X:980:GLY:HA3	6:X:1140:GLU:HB2	1.68	0.73
6:Y:331:ARG:HD3	6:Y:387:CYS:SG	2.29	0.73
6:Y:980:GLY:HA3	6:Y:1140:GLU:HG2	1.70	0.73
1:f:58:PRO:HB2	3:F:423:GLN:CD	2.13	0.73
1:h:59:HIS:ND1	1:h:60:ALA:C	2.47	0.73
1:n:58:PRO:HB3	3:N:423:GLN:HE21	1.53	0.73
1:n:60:ALA:H	3:N:398:SER:HA	1.54	0.73
1:t:45:ARG:CG	3:T:399:ALA:O	2.36	0.73
1:t:59:HIS:HB3	3:T:398:SER:HG	1.48	0.73
3:D:368:THR:HG22	3:D:446:PRO:HG3	1.71	0.73
3:D:644:LEU:O	3:D:646:PRO:HD3	1.89	0.73

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:I:9:ASN:HD21	3:J:206:GLU:HA	1.53	0.73
3:J:98:ALA:HB2	3:J:206:GLU:HG2	1.71	0.73
3:J:375:LEU:HD23	3:J:437:ILE:CD1	2.15	0.73
3:L:382:ILE:HA	3:L:385:VAL:HG12	1.71	0.73
3:L:412:ARG:HH22	3:N:356:THR:HG21	1.53	0.73
3:N:354:SER:HB2	3:N:419:MET:HA	1.69	0.73
3:P:609:ARG:HB3	2:Q:29:THR:CG2	2.14	0.73
3:T:375:LEU:HD23	3:T:437:ILE:CD1	2.15	0.73
4:U:239:PRO:O	4:U:358:ASP:HB3	1.89	0.73
5:W:1208:CYS:HB3	5:W:1216:GLY:CA	2.18	0.73
6:X:1122:PRO:CB	6:X:1159:GLY:HA3	2.19	0.73
6:Y:618:PRO:HB2	6:Y:941:TYR:CG	2.24	0.73
6:Y:632:ASN:HB3	6:Y:634:GLN:NE2	2.03	0.73
1:b:58:PRO:HG3	3:B:423:GLN:NE2	2.03	0.72
1:r:22:LEU:HG	1:r:83:VAL:CG2	2.18	0.72
3:F:322:TYR:O	3:F:323:SER:OG	2.06	0.72
3:F:368:THR:HG22	3:F:446:PRO:HG3	1.71	0.72
2:G:13:ILE:HD11	7:G:101:MYR:H111	1.70	0.72
3:J:609:ARG:HB3	2:K:29:THR:CG2	2.14	0.72
3:N:271:LEU:HD11	3:N:516:LYS:HB2	1.69	0.72
2:O:29:THR:CG2	3:T:609:ARG:HB3	2.14	0.72
3:P:121:ALA:CB	3:T:87:PHE:CE1	2.71	0.72
3:P:289:LEU:HD21	3:R:515:ASN:ND2	2.04	0.72
3:R:98:ALA:HB2	3:R:206:GLU:HG2	1.71	0.72
3:R:412:ARG:HH22	3:T:356:THR:HG21	1.53	0.72
3:T:48:TRP:CH2	3:T:122:THR:HA	2.24	0.72
5:W:243:GLN:HE22	5:W:257:GLN:HG3	1.52	0.72
1:t:8:GLN:HG2	3:P:302:GLU:HG3	0.77	0.72
3:B:87:PHE:CZ	3:B:235:LEU:HD12	2.24	0.72
3:B:368:THR:HG22	3:B:446:PRO:HG3	1.71	0.72
3:B:382:ILE:HA	3:B:385:VAL:HG12	1.70	0.72
3:F:87:PHE:CE1	3:H:121:ALA:CB	2.71	0.72
3:F:271:LEU:HD11	3:F:516:LYS:HB2	1.69	0.72
3:H:291:PRO:HB3	3:H:467:THR:CG2	2.17	0.72
3:H:368:THR:HG22	3:H:446:PRO:HG3	1.71	0.72
3:J:57:THR:HG22	3:J:58:ILE:N	2.04	0.72
3:J:368:THR:HG22	3:J:446:PRO:HG3	1.71	0.72
3:J:637:PHE:HE1	3:N:545:VAL:CG2	2.01	0.72
3:L:48:TRP:CH2	3:L:122:THR:HA	2.23	0.72
3:R:87:PHE:CE1	3:T:121:ALA:CB	2.71	0.72
4:U:170:THR:HG23	4:U:207:THR:HA	1.71	0.72

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
5:W:63:PRO:HB2	5:W:64:PRO:CD	2.19	0.72
5:W:853:THR:HG22	5:W:873:ALA:HB3	1.71	0.72
6:X:801:ARG:HH11	6:X:801:ARG:CB	1.94	0.72
6:X:937:ALA:CB	6:X:947:ARG:HA	2.18	0.72
1:h:23:THR:HB	3:H:323:SER:HA	1.00	0.72
1:p:44:GLY:CA	3:P:399:ALA:HB3	2.19	0.72
1:t:59:HIS:CA	3:T:398:SER:HA	2.20	0.72
2:A:9:ASN:HD21	3:B:206:GLU:HA	1.53	0.72
3:D:472:ARG:HH21	3:F:530:ARG:NE	1.87	0.72
3:F:92:TRP:NE1	3:F:156:VAL:HG21	2.03	0.72
3:J:363:ASN:HD21	3:J:366:LEU:HB2	1.55	0.72
3:L:354:SER:HB2	3:L:419:MET:HA	1.69	0.72
3:N:644:LEU:O	3:N:646:PRO:HD3	1.89	0.72
2:O:13:ILE:HD11	7:O:101:MYR:H111	1.70	0.72
3:R:238:VAL:HG22	3:T:142:ILE:CD1	2.19	0.72
3:R:363:ASN:HD21	3:R:366:LEU:HB2	1.55	0.72
3:T:87:PHE:CZ	3:T:235:LEU:HD12	2.25	0.72
3:T:469:VAL:O	3:T:470:LEU:HB2	1.87	0.72
5:W:137:ARG:HH22	5:W:164:VAL:HG21	1.54	0.72
5:W:574:THR:HA	5:W:609:MET:O	1.89	0.72
6:Y:485:PRO:HB3	6:Y:836:HIS:CD2	2.24	0.72
1:h:23:THR:CG2	3:H:323:SER:CB	2.67	0.72
1:p:35:PHE:CD2	1:p:73:ARG:HG3	2.25	0.72
1:t:24:LEU:CD2	1:t:35:PHE:CD1	2.72	0.72
1:t:45:ARG:CD	3:T:401:GLY:H	2.03	0.72
3:B:92:TRP:NE1	3:B:156:VAL:HG21	2.03	0.72
3:J:164:ASP:OD2	3:N:534:SER:HB3	1.89	0.72
3:J:382:ILE:HA	3:J:385:VAL:HG12	1.70	0.72
3:L:87:PHE:CZ	3:L:235:LEU:HD12	2.25	0.72
2:O:8:VAL:CG1	3:P:206:GLU:OE2	2.38	0.72
3:R:237:GLU:CD	3:T:46:LYS:HZ3	1.97	0.72
3:R:375:LEU:HD23	3:R:437:ILE:CD1	2.15	0.72
2:S:8:VAL:CG1	3:T:206:GLU:OE2	2.38	0.72
6:X:497:ILE:HD13	6:X:514:ILE:CD1	2.19	0.72
6:X:820:VAL:O	6:X:824:CYS:HB3	1.89	0.72
3:B:428:ARG:HG2	3:B:428:ARG:NH1	2.03	0.72
3:D:515:ASN:ND2	3:H:289:LEU:HD21	2.05	0.72
3:F:412:ARG:HH22	3:H:356:THR:HG21	1.53	0.72
3:H:372:ALA:CB	3:H:399:ALA:HA	2.18	0.72
3:J:53:THR:HA	4:V:332:PRO:HG3	1.72	0.72
3:P:271:LEU:HD11	3:P:516:LYS:HB2	1.69	0.72

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:P:274:ILE:CD1	3:P:514:LEU:HD22	2.10	0.72
3:P:534:SER:HB3	3:R:164:ASP:OD2	1.89	0.72
3:R:545:VAL:CG2	3:T:637:PHE:HE1	2.01	0.72
2:S:13:ILE:HD11	7:S:101:MYR:H111	1.70	0.72
5:W:263:VAL:CG2	5:W:303:LEU:HD11	2.19	0.72
5:W:303:LEU:HD23	5:W:341:MET:HE3	1.71	0.72
5:W:1014:ASN:ND2	5:W:1213:SER:HB2	2.05	0.72
6:X:291:LEU:HD12	6:X:891:PHE:HB2	1.72	0.72
6:X:793:ARG:NH1	6:Y:693:TYR:CE2	2.57	0.72
6:Y:761:VAL:HG21	6:Y:844:LEU:HD13	1.71	0.72
3:D:428:ARG:HG2	3:D:428:ARG:NH1	2.03	0.72
3:F:341:MET:HA	3:F:341:MET:CE	2.15	0.72
3:F:354:SER:HB2	3:F:419:MET:HA	1.69	0.72
3:F:534:SER:HB3	3:H:164:ASP:OD2	1.89	0.72
2:G:8:VAL:CG1	3:H:206:GLU:OE2	2.38	0.72
2:I:8:VAL:CG1	3:J:206:GLU:OE2	2.38	0.72
3:L:411:VAL:O	3:L:411:VAL:HG12	1.90	0.72
3:N:48:TRP:CH2	3:N:122:THR:HA	2.23	0.72
3:N:363:ASN:HD21	3:N:366:LEU:HB2	1.55	0.72
3:P:78:MET:CE	3:T:238:VAL:CG2	2.59	0.72
3:P:354:SER:HB2	3:P:419:MET:HA	1.69	0.72
3:P:515:ASN:ND2	3:T:289:LEU:HD21	2.04	0.72
2:Q:8:VAL:CG1	3:R:206:GLU:OE2	2.38	0.72
2:Q:27:THR:O	2:Q:27:THR:CG2	2.38	0.72
3:R:289:LEU:HD21	3:T:515:ASN:ND2	2.04	0.72
4:U:102:PRO:HD3	6:X:880:LEU:HD11	1.70	0.72
4:U:166:MET:HE2	4:U:170:THR:HG22	1.70	0.72
4:U:223:LEU:HD11	4:U:411:PHE:CE2	2.24	0.72
4:U:231:GLY:HA2	4:U:255:ARG:NH1	2.04	0.72
6:X:509:SER:HB3	6:X:512:SER:HB3	1.72	0.72
2:A:8:VAL:CG1	3:B:206:GLU:OE2	2.38	0.72
3:D:545:VAL:CG2	3:F:637:PHE:HE1	2.01	0.72
3:F:411:VAL:O	3:F:412:ARG:HB2	1.87	0.72
3:H:98:ALA:HB2	3:H:206:GLU:HG2	1.71	0.72
2:I:27:THR:O	2:I:27:THR:CG2	2.38	0.72
2:K:27:THR:O	2:K:27:THR:CG2	2.38	0.72
3:P:363:ASN:HD21	3:P:366:LEU:HB2	1.55	0.72
3:P:375:LEU:HD23	3:P:437:ILE:CD1	2.15	0.72
3:P:644:LEU:O	3:P:646:PRO:HD3	1.89	0.72
3:T:98:ALA:HB2	3:T:206:GLU:HG2	1.71	0.72
6:X:801:ARG:HD3	6:X:939:LEU:HD13	1.72	0.72

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
6:Y:981:ASN:ND2	6:Y:1077:GLU:HG2	2.04	0.72
1:h:8:GLN:HG2	3:D:302:GLU:HG3	0.79	0.72
1:p:61:ASN:OD1	3:P:395:THR:HG21	1.90	0.72
3:B:609:ARG:O	3:B:610:THR:HG23	1.90	0.72
3:B:644:LEU:O	3:B:646:PRO:HD3	1.89	0.72
2:C:9:ASN:HD21	3:D:206:GLU:HA	1.54	0.72
3:D:98:ALA:HB2	3:D:206:GLU:HG2	1.71	0.72
3:D:363:ASN:HD21	3:D:366:LEU:HB2	1.55	0.72
2:E:8:VAL:CG1	3:F:206:GLU:OE2	2.38	0.72
3:J:356:THR:HG21	3:N:412:ARG:HH22	1.53	0.72
3:J:609:ARG:O	3:J:610:THR:HG23	1.90	0.72
3:P:411:VAL:O	3:P:411:VAL:HG12	1.90	0.72
3:R:87:PHE:CZ	3:R:235:LEU:HD12	2.24	0.72
4:V:48:THR:O	6:Y:417:VAL:HG21	1.89	0.72
5:W:623:PRO:HB2	5:W:628:TRP:HE1	1.53	0.72
6:X:398:ILE:O	6:X:401:THR:HG22	1.88	0.72
6:X:700:TRP:O	6:X:704:THR:HG22	1.90	0.72
1:t:44:GLY:O	3:T:399:ALA:CA	2.37	0.72
3:B:363:ASN:HD21	3:B:366:LEU:HB2	1.55	0.72
3:D:637:PHE:HE1	3:H:545:VAL:CG2	2.01	0.72
3:F:363:ASN:HD21	3:F:366:LEU:HB2	1.55	0.72
3:L:609:ARG:HB3	2:M:29:THR:CG2	2.14	0.72
3:N:382:ILE:HA	3:N:385:VAL:HG12	1.70	0.72
3:P:287:PRO:HG2	3:P:471:LEU:HD12	1.72	0.72
3:P:609:ARG:O	3:P:610:THR:HG23	1.90	0.72
3:T:287:PRO:HG2	3:T:471:LEU:HD12	1.72	0.72
5:W:137:ARG:HB3	5:W:153:LYS:HG2	1.71	0.72
5:W:672:THR:HG21	5:W:1081:VAL:HA	1.71	0.72
1:p:21:ARG:NH1	3:P:327:TYR:CZ	2.58	0.72
3:B:287:PRO:HG2	3:B:471:LEU:HD12	1.72	0.72
3:D:238:VAL:CG2	3:F:78:MET:CE	2.59	0.72
3:L:289:LEU:HD21	3:N:515:ASN:ND2	2.04	0.72
7:M:101:MYR:H143	7:M:101:MYR:C10	2.12	0.72
3:N:368:THR:HG22	3:N:446:PRO:HG3	1.71	0.72
3:P:341:MET:HA	3:P:341:MET:CE	2.15	0.72
7:Q:101:MYR:H142	7:Q:101:MYR:C10	2.10	0.72
3:R:644:LEU:O	3:R:646:PRO:HD3	1.89	0.72
5:W:564:LEU:HD13	5:W:570:VAL:HG13	1.70	0.72
5:W:566:LEU:HD13	5:W:605:THR:OG1	1.89	0.72
6:X:827:THR:HG21	6:X:831:LEU:HD22	1.72	0.72
6:Y:54:VAL:HB	6:Y:57:GLN:HG3	1.72	0.72

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
6:Y:990:ASN:HB2	6:Y:1131:ARG:HA	1.70	0.72
1:d:44:GLY:O	3:D:399:ALA:CA	2.38	0.71
1:j:15:ARG:CZ	3:J:406:ASN:CG	2.62	0.71
2:C:8:VAL:CG1	3:D:206:GLU:OE2	2.38	0.71
2:C:13:ILE:HD11	7:C:101:MYR:H111	1.70	0.71
3:J:87:PHE:CZ	3:J:235:LEU:HD12	2.24	0.71
3:J:575:MET:CE	3:N:469:VAL:CG2	2.43	0.71
2:O:13:ILE:HB	3:P:205:ASN:HB3	1.72	0.71
3:P:59:ASP:HB2	5:W:380:ASN:HB2	1.70	0.71
3:P:87:PHE:CZ	3:P:235:LEU:HD12	2.24	0.71
3:T:354:SER:HB2	3:T:419:MET:HA	1.70	0.71
3:T:644:LEU:O	3:T:646:PRO:HD3	1.89	0.71
4:V:195:GLN:HE22	6:Y:377:ILE:H	1.35	0.71
5:W:675:GLN:O	5:W:679:LEU:HB2	1.90	0.71
6:X:453:LEU:CD2	6:X:861:ARG:HA	2.18	0.71
1:f:45:ARG:CG	3:F:401:GLY:N	2.42	0.71
1:p:15:ARG:HH22	3:P:402:ALA:HB3	1.54	0.71
3:F:289:LEU:HD21	3:H:515:ASN:ND2	2.04	0.71
3:H:322:TYR:O	3:H:323:SER:OG	2.06	0.71
3:H:363:ASN:HD21	3:H:366:LEU:HB2	1.55	0.71
3:J:644:LEU:O	3:J:646:PRO:HD3	1.89	0.71
3:L:322:TYR:O	3:L:323:SER:OG	2.07	0.71
3:L:564:LEU:HD21	3:L:568:ARG:HH21	1.55	0.71
3:N:609:ARG:O	3:N:610:THR:HG23	1.90	0.71
2:O:9:ASN:HD21	3:P:206:GLU:HA	1.53	0.71
5:W:708:LEU:HD23	5:W:764:THR:HG22	1.72	0.71
5:W:740:VAL:HG21	5:W:770:ARG:NH1	2.05	0.71
5:W:1234:ASN:O	5:W:1235:THR:HG23	1.90	0.71
6:X:317:PHE:HB2	6:X:373:ARG:HH22	1.55	0.71
6:X:746:TYR:HA	6:X:750:MET:HE3	1.70	0.71
6:Y:640:ARG:HH11	6:Y:640:ARG:HG3	1.54	0.71
6:Y:692:THR:HG23	6:Y:693:TYR:HD1	1.55	0.71
1:j:58:PRO:CG	3:J:423:GLN:NE2	2.40	0.71
3:D:291:PRO:HB3	3:D:467:THR:CG2	2.17	0.71
3:D:411:VAL:O	3:D:411:VAL:HG12	1.90	0.71
3:F:287:PRO:HG2	3:F:471:LEU:HD12	1.72	0.71
3:F:369:SER:HG	3:F:443:PHE:HD1	1.34	0.71
3:F:469:VAL:CG2	3:H:575:MET:CE	2.43	0.71
3:J:289:LEU:HD21	3:L:515:ASN:ND2	2.04	0.71
3:J:564:LEU:HD21	3:J:568:ARG:HH21	1.55	0.71
3:L:287:PRO:HG2	3:L:471:LEU:HD12	1.72	0.71

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:R:368:THR:HG22	3:R:446:PRO:HG3	1.71	0.71
3:T:428:ARG:HG2	3:T:428:ARG:NH1	2.03	0.71
4:U:41:PHE:HD1	6:X:404:MET:CE	2.03	0.71
4:U:177:TYR:CZ	6:X:412:ASP:HB3	2.25	0.71
5:W:769:ARG:HD3	5:W:844:SER:O	1.90	0.71
1:p:59:HIS:HA	3:P:398:SER:CA	2.16	0.71
1:r:15:ARG:CZ	3:R:402:ALA:CB	2.68	0.71
3:B:564:LEU:HD21	3:B:568:ARG:HH21	1.55	0.71
3:D:87:PHE:CZ	3:D:235:LEU:HD12	2.24	0.71
3:D:287:PRO:HG2	3:D:471:LEU:HD12	1.72	0.71
2:E:9:ASN:HD21	3:F:206:GLU:HA	1.53	0.71
3:F:274:ILE:CD1	3:F:514:LEU:HD23	2.20	0.71
3:F:609:ARG:O	3:F:610:THR:HG23	1.90	0.71
3:H:411:VAL:O	3:H:411:VAL:HG12	1.90	0.71
3:L:363:ASN:HD21	3:L:366:LEU:HB2	1.55	0.71
7:M:101:MYR:H21	3:N:191:LYS:CE	2.20	0.71
3:N:411:VAL:O	3:N:412:ARG:HB2	1.87	0.71
3:R:354:SER:HB2	3:R:419:MET:HA	1.70	0.71
6:Y:317:PHE:HZ	6:Y:385:VAL:HG11	1.56	0.71
6:Y:611:PRO:O	6:Y:614:GLN:HB3	1.90	0.71
1:t:24:LEU:CD1	1:t:41:VAL:HG22	2.21	0.71
3:D:237:GLU:CB	3:F:46:LYS:HZ1	2.01	0.71
3:D:412:ARG:HH22	3:F:356:THR:HG21	1.53	0.71
3:D:609:ARG:O	3:D:610:THR:HG23	1.90	0.71
2:E:13:ILE:HB	3:F:205:ASN:HB3	1.72	0.71
3:F:87:PHE:CZ	3:F:235:LEU:HD12	2.24	0.71
3:L:341:MET:HA	3:L:341:MET:CE	2.15	0.71
2:M:13:ILE:HB	3:N:205:ASN:HB3	1.72	0.71
3:N:87:PHE:CZ	3:N:235:LEU:HD12	2.24	0.71
3:N:98:ALA:HB2	3:N:206:GLU:HG2	1.71	0.71
3:P:164:ASP:OD2	3:T:534:SER:HB3	1.89	0.71
3:R:287:PRO:HG2	3:R:471:LEU:HD12	1.72	0.71
3:R:469:VAL:CG2	3:T:575:MET:CE	2.43	0.71
3:T:363:ASN:HD21	3:T:366:LEU:HB2	1.55	0.71
3:T:564:LEU:HD21	3:T:568:ARG:HH21	1.55	0.71
5:W:433:CYS:O	5:W:1007:PRO:HA	1.89	0.71
5:W:981:LEU:HD12	5:W:1028:ILE:HG21	1.71	0.71
6:X:301:LEU:HD13	6:X:964:VAL:HG12	1.71	0.71
6:Y:489:ASP:OD1	6:Y:491:THR:HG23	1.90	0.71
6:Y:603:VAL:HG13	6:Y:606:MET:HB2	1.70	0.71
2:G:13:ILE:HB	3:H:205:ASN:HB3	1.72	0.71

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:L:534:SER:HB3	3:N:164:ASP:OD2	1.89	0.71
3:L:606:THR:CG2	3:N:173:ASP:HB2	2.21	0.71
3:T:58:ILE:HG22	3:T:58:ILE:O	1.90	0.71
5:W:102:ARG:HH11	5:W:102:ARG:HG2	1.56	0.71
5:W:1087:LEU:HD21	5:W:1096:PHE:HB3	1.73	0.71
6:X:597:GLU:HG3	6:X:636:ALA:HA	1.72	0.71
1:h:59:HIS:CG	1:h:60:ALA:N	2.52	0.71
1:r:5:MET:O	1:r:6:ILE:C	2.31	0.71
3:B:96:PHE:HD1	3:B:101:THR:HG23	1.56	0.71
3:F:411:VAL:O	3:F:411:VAL:HG12	1.90	0.71
3:H:300:LYS:HE3	3:H:303:ASP:OD2	1.91	0.71
3:J:237:GLU:CD	3:L:46:LYS:HZ3	1.99	0.71
3:J:515:ASN:ND2	3:N:289:LEU:HD21	2.04	0.71
3:N:300:LYS:HE3	3:N:303:ASP:OD2	1.91	0.71
3:P:51:VAL:HB	5:W:379:VAL:HG21	1.73	0.71
3:P:98:ALA:HB2	3:P:206:GLU:HG2	1.71	0.71
3:P:173:ASP:HB2	3:T:606:THR:CG2	2.21	0.71
3:R:96:PHE:HD1	3:R:101:THR:HG23	1.56	0.71
3:R:382:ILE:HA	3:R:385:VAL:HG12	1.71	0.71
3:R:411:VAL:O	3:R:411:VAL:HG12	1.90	0.71
2:S:13:ILE:HB	3:T:205:ASN:HB3	1.72	0.71
5:W:409:ILE:HG12	5:W:779:MET:CE	2.21	0.71
5:W:1057:PRO:HG3	5:W:1089:TYR:CE1	2.26	0.71
6:X:995:PRO:HG2	6:X:1118:CYS:O	1.90	0.71
2:I:13:ILE:HD11	7:I:101:MYR:H111	1.70	0.71
3:N:287:PRO:HG2	3:N:471:LEU:HD12	1.72	0.71
7:O:101:MYR:H142	7:O:101:MYR:C10	2.11	0.71
3:R:274:ILE:CD1	3:R:514:LEU:HD23	2.20	0.71
4:U:93:ARG:HB3	4:U:93:ARG:NH1	2.06	0.71
5:W:480:ARG:NE	5:W:649:ILE:HG22	2.05	0.71
5:W:714:ARG:HB2	5:W:758:THR:HG22	1.72	0.71
6:X:528:PHE:CE1	6:X:575:MET:CE	2.72	0.71
6:X:673:ASN:O	6:X:674:ARG:HD2	1.91	0.71
1:j:22:LEU:HD13	1:j:22:LEU:O	1.91	0.71
1:n:15:ARG:CZ	3:N:402:ALA:HB2	2.21	0.71
1:r:15:ARG:NH1	3:R:402:ALA:CB	2.53	0.71
3:B:98:ALA:HB2	3:B:206:GLU:HG2	1.71	0.71
3:D:375:LEU:HD23	3:D:437:ILE:CD1	2.15	0.71
3:F:98:ALA:HB2	3:F:206:GLU:HG2	1.71	0.71
3:F:606:THR:CG2	3:H:173:ASP:HB2	2.21	0.71
3:H:87:PHE:CZ	3:H:235:LEU:HD12	2.25	0.71

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:J:274:ILE:CD1	3:J:514:LEU:HD23	2.20	0.71
3:J:530:ARG:NE	3:N:472:ARG:HH21	1.87	0.71
3:J:606:THR:CG2	3:L:173:ASP:HB2	2.21	0.71
3:R:564:LEU:HD21	3:R:568:ARG:HH21	1.55	0.71
2:S:5:GLN:CG	2:S:5:GLN:OE1	2.39	0.71
2:S:26:MET:HG3	3:T:216:PRO:CG	2.21	0.71
4:U:31:GLY:C	4:U:32:CYS:SG	2.73	0.71
5:W:38:ASN:HD21	5:W:40:HIS:HB3	1.54	0.71
6:Y:923:LEU:CD2	6:Y:931:ILE:HD13	2.21	0.71
1:b:74:GLN:CA	1:b:75:PRO:N	2.54	0.71
1:j:60:ALA:H	3:J:398:SER:HA	1.54	0.71
1:n:8:GLN:HE22	3:J:302:GLU:C	1.94	0.71
3:B:291:PRO:HB3	3:B:467:THR:CG2	2.17	0.71
3:D:57:THR:H	3:D:60:SER:HB3	1.55	0.71
3:D:96:PHE:HD1	3:D:101:THR:HG23	1.56	0.71
7:G:101:MYR:H52	7:G:101:MYR:H91	1.73	0.71
3:H:58:ILE:HG22	3:H:58:ILE:O	1.90	0.71
3:H:609:ARG:O	3:H:610:THR:HG23	1.90	0.71
3:L:96:PHE:HD1	3:L:101:THR:HG23	1.56	0.71
3:L:178:SER:HB3	2:Q:3:ASN:HD21	1.55	0.71
3:L:609:ARG:O	3:L:610:THR:HG23	1.90	0.71
3:P:606:THR:CG2	3:R:173:ASP:HB2	2.21	0.71
5:W:335:SER:HA	5:W:752:ASN:O	1.91	0.71
6:X:590:ALA:HA	6:X:639:LEU:HD11	1.72	0.71
6:Y:167:ILE:O	6:Y:167:ILE:HG13	1.91	0.71
6:Y:209:ASN:ND2	6:Y:242:ARG:HG3	2.05	0.71
1:p:60:ALA:HB3	3:P:322:TYR:OH	1.89	0.70
1:r:4:HIS:ND1	1:r:54:THR:HG22	2.06	0.70
1:t:24:LEU:CD2	1:t:38:ALA:HB3	2.21	0.70
3:D:469:VAL:CG2	3:F:575:MET:CE	2.43	0.70
3:D:579:ASN:ND2	3:D:582:LEU:HG	2.06	0.70
3:D:606:THR:CG2	3:F:173:ASP:HB2	2.21	0.70
3:D:609:ARG:HB3	2:E:29:THR:CG2	2.14	0.70
7:E:101:MYR:H91	7:E:101:MYR:H52	1.73	0.70
3:J:613:LEU:CD1	2:K:34:ILE:HB	2.21	0.70
2:K:8:VAL:CG1	3:L:206:GLU:OE2	2.38	0.70
3:N:58:ILE:HG22	3:N:58:ILE:O	1.90	0.70
2:O:34:ILE:HB	3:T:613:LEU:CD1	2.21	0.70
3:P:300:LYS:HE3	3:P:303:ASP:OD2	1.91	0.70
3:P:575:MET:CE	3:T:469:VAL:CG2	2.43	0.70
3:T:341:MET:HA	3:T:341:MET:CE	2.15	0.70

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
5:W:148:ILE:O	5:W:151:ILE:HG12	1.90	0.70
6:X:782:ASN:ND2	6:X:801:ARG:HD2	2.05	0.70
6:X:928:ASP:HB2	6:Y:740:LYS:HG2	1.72	0.70
6:X:1094:LEU:HD23	6:X:1094:LEU:O	1.90	0.70
6:Y:128:THR:HG22	6:Y:130:SER:H	1.55	0.70
6:Y:403:ILE:HG13	6:Y:1197:TYR:CE2	2.26	0.70
1:n:24:LEU:HD11	1:n:80:ARG:NE	2.05	0.70
1:p:35:PHE:HE2	1:p:73:ARG:CD	2.04	0.70
3:D:274:ILE:CD1	3:D:514:LEU:HD23	2.20	0.70
7:G:101:MYR:H21	3:H:191:LYS:CE	2.20	0.70
3:H:274:ILE:CD1	3:H:514:LEU:HD23	2.20	0.70
2:I:13:ILE:HB	3:J:205:ASN:HB3	1.72	0.70
3:J:173:ASP:HB2	3:N:606:THR:CG2	2.21	0.70
3:J:291:PRO:HB3	3:J:467:THR:CG2	2.17	0.70
7:M:101:MYR:H91	7:M:101:MYR:H52	1.73	0.70
3:N:96:PHE:HD1	3:N:101:THR:HG23	1.56	0.70
3:P:274:ILE:CD1	3:P:514:LEU:HD23	2.20	0.70
3:P:450:SER:HA	3:P:453:ASN:ND2	2.07	0.70
3:R:291:PRO:CB	3:R:467:THR:CG2	2.56	0.70
3:R:606:THR:CG2	3:T:173:ASP:HB2	2.21	0.70
6:X:671:PRO:HB2	6:X:674:ARG:HG3	1.73	0.70
1:d:15:ARG:NH2	3:D:402:ALA:CB	2.55	0.70
3:B:300:LYS:HE3	3:B:303:ASP:OD2	1.91	0.70
3:D:289:LEU:HD21	3:F:515:ASN:ND2	2.04	0.70
3:F:300:LYS:HE3	3:F:303:ASP:OD2	1.91	0.70
3:H:287:PRO:HG2	3:H:471:LEU:HD12	1.72	0.70
3:H:450:SER:HA	3:H:453:ASN:ND2	2.06	0.70
2:K:13:ILE:HB	3:L:205:ASN:HB3	1.72	0.70
3:L:300:LYS:HE3	3:L:303:ASP:OD2	1.91	0.70
3:N:517:LEU:HG	3:N:601:THR:HG22	1.74	0.70
3:N:564:LEU:HD21	3:N:568:ARG:HH21	1.55	0.70
3:P:382:ILE:HA	3:P:385:VAL:HG12	1.70	0.70
3:T:336:THR:CG2	3:T:456:ASN:HD21	2.04	0.70
4:U:41:PHE:CD1	6:X:404:MET:CE	2.73	0.70
5:W:75:ILE:HD11	5:W:199:PRO:HG2	1.74	0.70
5:W:1065:HIS:O	5:W:1066:ASN:HB2	1.90	0.70
6:X:193:LEU:HD22	6:X:855:GLN:HE21	1.56	0.70
6:Y:457:MET:HG2	6:Y:860:THR:HG21	1.72	0.70
6:Y:882:PRO:O	6:Y:884:PRO:HD3	1.92	0.70
1:l:45:ARG:CG	3:L:399:ALA:O	2.39	0.70
1:b:59:HIS:CE1	1:b:64:THR:HG1	2.07	0.70

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:t:24:LEU:CD2	1:t:35:PHE:HD1	2.04	0.70
3:B:411:VAL:O	3:B:411:VAL:HG12	1.90	0.70
7:C:101:MYR:H52	7:C:101:MYR:H91	1.73	0.70
3:H:564:LEU:HD21	3:H:568:ARG:HH21	1.56	0.70
7:I:101:MYR:H91	7:I:101:MYR:H52	1.73	0.70
3:N:450:SER:HA	3:N:453:ASN:ND2	2.06	0.70
3:R:609:ARG:O	3:R:610:THR:HG23	1.90	0.70
5:W:14:ASN:HB3	5:W:319:GLN:NE2	2.06	0.70
5:W:35:GLN:CD	5:W:96:VAL:HG11	2.16	0.70
6:X:583:VAL:HG21	6:X:621:TRP:CG	2.26	0.70
6:Y:1160:GLU:O	6:Y:1160:GLU:CG	2.39	0.70
1:j:61:ASN:O	1:j:63:LYS:N	2.25	0.70
2:C:13:ILE:HB	3:D:205:ASN:HB3	1.72	0.70
3:F:58:ILE:HG22	3:F:58:ILE:O	1.90	0.70
3:F:336:THR:CG2	3:F:456:ASN:HD21	2.04	0.70
2:G:5:GLN:CG	2:G:5:GLN:OE1	2.39	0.70
3:H:96:PHE:HD1	3:H:101:THR:HG23	1.56	0.70
3:H:336:THR:CG2	3:H:456:ASN:HD21	2.04	0.70
3:J:84:LYS:CD	2:K:39:GLY:O	2.40	0.70
3:J:336:THR:CG2	3:J:456:ASN:HD21	2.04	0.70
3:J:428:ARG:HG2	3:J:428:ARG:NH1	2.03	0.70
3:P:517:LEU:HG	3:P:601:THR:HG22	1.74	0.70
3:R:450:SER:HA	3:R:453:ASN:ND2	2.07	0.70
4:V:51:LEU:HD13	6:X:1038:PRO:HD3	1.73	0.70
6:X:792:GLN:HG2	6:X:931:ILE:HG23	1.73	0.70
1:f:15:ARG:NH2	3:F:402:ALA:CB	2.55	0.70
7:A:101:MYR:H91	7:A:101:MYR:H52	1.73	0.70
3:D:336:THR:CG2	3:D:456:ASN:HD21	2.04	0.70
3:J:78:MET:CE	3:N:238:VAL:CG2	2.59	0.70
3:L:336:THR:CG2	3:L:456:ASN:HD21	2.04	0.70
3:P:368:THR:HG22	3:P:446:PRO:HG3	1.71	0.70
3:P:428:ARG:HG2	3:P:428:ARG:NH1	2.03	0.70
2:Q:13:ILE:HB	3:R:205:ASN:HB3	1.72	0.70
7:S:101:MYR:H91	7:S:101:MYR:H52	1.73	0.70
5:W:438:VAL:HB	5:W:649:ILE:HD13	1.72	0.70
5:W:519:LEU:HD13	5:W:566:LEU:HG	1.73	0.70
6:Y:832:ILE:HD13	6:Y:1213:ALA:HB2	1.73	0.70
1:l:24:LEU:HD21	1:l:80:ARG:CD	2.20	0.70
1:f:19:ALA:O	1:f:20:GLY:C	2.34	0.70
1:d:23:THR:HG21	3:D:323:SER:CA	2.13	0.70
1:p:62:VAL:CG1	1:p:65:ILE:HD12	2.22	0.70

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:B:58:ILE:HG22	3:B:58:ILE:O	1.90	0.70
2:I:34:ILE:HB	3:N:613:LEU:CD1	2.21	0.70
3:L:450:SER:HA	3:L:453:ASN:ND2	2.07	0.70
2:M:8:VAL:CG1	3:N:206:GLU:OE2	2.38	0.70
3:N:274:ILE:CD1	3:N:514:LEU:HD23	2.20	0.70
3:P:84:LYS:CD	2:Q:39:GLY:O	2.40	0.70
3:P:96:PHE:HD1	3:P:101:THR:HG23	1.56	0.70
3:R:336:THR:CG2	3:R:456:ASN:HD21	2.04	0.70
3:T:411:VAL:O	3:T:411:VAL:HG12	1.90	0.70
5:W:1204:LEU:HB3	5:W:1227:TYR:CD2	2.27	0.70
6:X:317:PHE:CE2	6:X:370:ARG:HB3	2.25	0.70
6:X:795:ASP:OD2	6:X:803:THR:HG22	1.91	0.70
6:X:1012:ARG:NH2	6:X:1043:GLY:HA3	2.03	0.70
1:l:4:HIS:CE1	1:l:54:THR:CG2	2.75	0.70
1:d:59:HIS:CB	3:D:397:VAL:O	2.36	0.70
1:j:6:ILE:CD1	3:L:586:VAL:HG21	2.22	0.70
1:j:45:ARG:CD	3:J:400:ALA:HA	2.20	0.70
1:t:60:ALA:HB2	3:T:322:TYR:CZ	2.27	0.70
3:J:277:ASP:O	3:J:278:THR:CG2	2.37	0.70
2:K:26:MET:HG3	3:L:216:PRO:CG	2.21	0.70
3:L:609:ARG:HH11	2:M:27:THR:CG2	2.05	0.70
3:N:579:ASN:ND2	3:N:582:LEU:HG	2.06	0.70
3:P:613:LEU:CD1	2:Q:34:ILE:HB	2.21	0.70
7:S:101:MYR:H21	3:T:191:LYS:CE	2.20	0.70
4:V:271:SER:HB3	4:V:330:PRO:HD2	1.73	0.70
5:W:934:ASN:O	5:W:967:TYR:HB2	1.92	0.70
6:X:249:TRP:HZ3	6:X:342:LEU:HD23	1.57	0.70
6:Y:104:GLN:CD	6:Y:104:GLN:H	2.00	0.70
6:Y:241:ASN:O	6:Y:1147:PHE:HB3	1.92	0.70
1:d:4:HIS:NE2	1:d:54:THR:HG21	2.07	0.70
1:j:5:MET:HE2	1:j:5:MET:CA	2.22	0.70
1:n:15:ARG:NH1	3:N:406:ASN:CG	2.47	0.70
3:B:274:ILE:CD1	3:B:514:LEU:HD23	2.20	0.70
3:B:517:LEU:HG	3:B:601:THR:HG22	1.73	0.70
3:D:300:LYS:HE3	3:D:303:ASP:OD2	1.91	0.70
3:F:564:LEU:HD21	3:F:568:ARG:HH21	1.55	0.70
2:I:27:THR:CG2	3:N:609:ARG:HH11	2.05	0.70
2:K:9:ASN:HD21	3:L:206:GLU:HA	1.53	0.70
3:N:411:VAL:O	3:N:411:VAL:HG12	1.90	0.70
3:R:300:LYS:HE3	3:R:303:ASP:OD2	1.91	0.70
3:T:609:ARG:O	3:T:610:THR:HG23	1.90	0.70

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
5:W:625:ARG:HG2	5:W:625:ARG:NH1	2.02	0.70
5:W:676:ARG:HB2	5:W:1081:VAL:CG1	2.21	0.70
5:W:679:LEU:HD22	5:W:1068:VAL:HG13	1.74	0.70
5:W:957:ARG:HB3	5:W:960:GLY:HA2	1.74	0.70
5:W:1233:ARG:HD2	5:W:1293:VAL:HG12	1.72	0.70
6:X:543:THR:O	6:X:578:ASP:HB2	1.92	0.70
1:f:59:HIS:HB2	3:F:398:SER:HA	1.74	0.70
1:d:21:ARG:O	3:D:323:SER:CB	2.40	0.70
1:j:61:ASN:OD1	3:J:395:THR:HG21	1.91	0.70
3:B:336:THR:CG2	3:B:456:ASN:HD21	2.04	0.70
3:L:58:ILE:HG22	3:L:58:ILE:O	1.90	0.70
3:R:58:ILE:HG22	3:R:58:ILE:O	1.90	0.70
3:R:341:MET:HE2	3:R:341:MET:CA	2.18	0.70
3:T:300:LYS:HE3	3:T:303:ASP:OD2	1.91	0.70
4:U:60:SER:HB3	4:U:63:SER:CB	2.13	0.70
5:W:1064:GLN:HB3	5:W:1069:GLU:HA	1.72	0.70
6:X:368:ILE:HG12	6:X:369:GLY:N	2.05	0.70
6:X:380:GLU:OE1	6:X:438:PRO:HG2	1.92	0.70
6:X:784:LEU:HD12	6:X:784:LEU:N	2.06	0.70
6:X:1125:TYR:O	6:X:1157:ALA:HB3	1.92	0.70
1:l:15:ARG:NH1	3:L:406:ASN:CB	2.50	0.69
1:f:22:LEU:HD11	1:f:79:ILE:HG22	1.74	0.69
1:d:23:THR:CG2	3:D:324:GLY:N	2.54	0.69
3:D:173:ASP:HB2	3:H:606:THR:CG2	2.21	0.69
3:H:582:LEU:HD23	3:H:585:GLN:NE2	2.07	0.69
3:J:287:PRO:HG2	3:J:471:LEU:HD12	1.72	0.69
3:L:368:THR:HG22	3:L:446:PRO:HG3	1.71	0.69
3:L:517:LEU:HG	3:L:601:THR:HG22	1.74	0.69
3:L:582:LEU:HD23	3:L:585:GLN:NE2	2.07	0.69
2:M:27:THR:O	2:M:27:THR:CG2	2.38	0.69
3:N:277:ASP:O	3:N:278:THR:CG2	2.37	0.69
5:W:149:THR:HA	5:W:152:THR:HG22	1.72	0.69
1:l:23:THR:HG23	1:l:25:TYR:CE2	2.26	0.69
1:n:8:GLN:NE2	3:J:302:GLU:CG	2.49	0.69
1:p:6:ILE:HG21	3:R:494:THR:HG22	1.74	0.69
2:C:34:ILE:HB	3:H:613:LEU:CD1	2.21	0.69
3:D:341:MET:HA	3:D:341:MET:CE	2.15	0.69
3:D:530:ARG:NE	3:H:472:ARG:HH21	1.87	0.69
3:L:274:ILE:CD1	3:L:514:LEU:HD23	2.20	0.69
2:M:5:GLN:OE1	2:M:5:GLN:CG	2.39	0.69
3:N:336:THR:CG2	3:N:456:ASN:HD21	2.04	0.69

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:N:582:LEU:HD23	3:N:585:GLN:NE2	2.07	0.69
2:O:39:GLY:O	3:T:84:LYS:CD	2.40	0.69
3:P:564:LEU:HD21	3:P:568:ARG:HH21	1.55	0.69
3:T:274:ILE:CD1	3:T:514:LEU:HD23	2.20	0.69
3:T:470:LEU:HD11	3:T:472:ARG:HH11	1.56	0.69
4:U:94:ASP:CG	6:X:949:HIS:HE2	2.00	0.69
5:W:584:GLN:CG	5:W:621:ASN:HB2	2.19	0.69
5:W:1243:TYR:CE1	5:W:1298:ILE:HD11	2.27	0.69
6:Y:479:LEU:HB3	6:Y:758:LEU:HD11	1.74	0.69
1:f:44:GLY:HA3	3:F:322:TYR:CZ	2.23	0.69
1:h:23:THR:CG2	3:H:323:SER:HB3	2.21	0.69
1:p:3:LEU:HD13	1:p:3:LEU:O	1.93	0.69
2:A:13:ILE:HB	3:B:205:ASN:HB3	1.72	0.69
3:D:582:LEU:HD23	3:D:585:GLN:NE2	2.07	0.69
3:F:237:GLU:CB	3:H:46:LYS:HZ1	2.01	0.69
3:F:379:SER:HB3	3:N:380:VAL:HG22	1.73	0.69
3:F:582:LEU:HD23	3:F:585:GLN:NE2	2.07	0.69
3:J:609:ARG:HH11	2:K:27:THR:CG2	2.05	0.69
3:P:582:LEU:HD23	3:P:585:GLN:NE2	2.08	0.69
5:W:63:PRO:HA	5:W:66:ASN:OD1	1.92	0.69
6:X:252:LEU:HD12	6:X:253:GLY:N	2.07	0.69
6:X:600:ALA:O	6:X:623:PRO:HD2	1.91	0.69
6:Y:371:LEU:HD21	6:Y:388:MET:HE2	1.73	0.69
6:Y:717:MET:HE2	6:Y:717:MET:HA	1.75	0.69
3:B:339:ILE:HD11	3:B:464:LEU:HD22	1.75	0.69
2:C:26:MET:HG3	3:D:216:PRO:CG	2.21	0.69
3:J:582:LEU:HD23	3:J:585:GLN:NE2	2.08	0.69
3:L:98:ALA:HB2	3:L:206:GLU:HG2	1.71	0.69
3:L:274:ILE:HD11	3:L:514:LEU:HD21	1.73	0.69
3:N:339:ILE:HD11	3:N:464:LEU:HD22	1.75	0.69
7:O:101:MYR:H91	7:O:101:MYR:H52	1.73	0.69
3:T:339:ILE:HD11	3:T:464:LEU:HD22	1.75	0.69
4:U:51:LEU:HB3	4:U:53:VAL:HG13	1.74	0.69
6:Y:451:LEU:HD13	6:Y:666:ALA:HB2	1.73	0.69
6:Y:1156:THR:CG2	6:Y:1161:ASN:HB2	2.23	0.69
1:p:58:PRO:O	1:p:59:HIS:ND1	2.25	0.69
1:t:5:MET:HA	1:t:5:MET:CE	2.22	0.69
3:D:416:ARG:CB	3:F:298:PHE:CE1	2.76	0.69
3:F:96:PHE:HD1	3:F:101:THR:HG23	1.56	0.69
3:P:294:TYR:HH	3:P:502:LEU:HA	1.57	0.69
3:P:336:THR:CG2	3:P:456:ASN:HD21	2.04	0.69

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:P:416:ARG:CB	3:R:298:PHE:CE1	2.76	0.69
3:R:609:ARG:HH11	2:S:27:THR:CG2	2.05	0.69
3:T:517:LEU:HG	3:T:601:THR:HG22	1.74	0.69
4:U:179:GLY:N	6:X:376:LEU:HD21	2.06	0.69
6:X:603:VAL:HG11	6:Y:715:VAL:HG11	1.75	0.69
1:d:23:THR:CB	3:D:323:SER:HA	2.23	0.69
1:d:74:GLN:O	1:d:74:GLN:NE2	2.26	0.69
2:C:27:THR:CG2	3:H:609:ARG:HH11	2.05	0.69
3:D:50:PRO:CB	3:D:60:SER:O	2.41	0.69
3:D:450:SER:HA	3:D:453:ASN:ND2	2.07	0.69
3:D:613:LEU:CD1	2:E:34:ILE:HB	2.21	0.69
7:E:101:MYR:H31	3:F:191:LYS:CG	2.23	0.69
2:I:26:MET:HG3	3:J:216:PRO:CG	2.21	0.69
3:J:416:ARG:CB	3:L:298:PHE:CE1	2.76	0.69
3:J:545:VAL:CG2	3:L:637:PHE:HE1	2.01	0.69
7:K:101:MYR:H52	7:K:101:MYR:H91	1.73	0.69
3:P:469:VAL:HG23	3:R:575:MET:CG	2.23	0.69
3:R:339:ILE:HD11	3:R:464:LEU:HD22	1.75	0.69
4:U:245:GLN:NE2	4:U:246:ALA:H	1.85	0.69
4:U:251:ASN:ND2	4:U:252:ASP:H	1.89	0.69
5:W:840:CYS:SG	5:W:854:MET:SD	2.90	0.69
5:W:1042:THR:OG1	5:W:1127:ILE:HA	1.92	0.69
6:X:875:GLN:HE22	6:X:930:ARG:HG2	1.56	0.69
1:b:16:ALA:C	1:b:22:LEU:CD2	2.59	0.69
1:t:22:LEU:HD22	1:t:22:LEU:C	2.18	0.69
2:A:27:THR:O	2:A:27:THR:CG2	2.38	0.69
3:B:339:ILE:HD13	3:B:349:LEU:HD21	1.75	0.69
7:C:101:MYR:H21	3:D:191:LYS:CE	2.23	0.69
3:R:517:LEU:HG	3:R:601:THR:HG22	1.74	0.69
4:U:65:ASN:HA	4:U:68:VAL:HG12	1.74	0.69
5:W:57:THR:HG23	5:W:183:TYR:HA	1.74	0.69
6:X:427:LEU:O	6:X:428:LEU:HB3	1.92	0.69
6:X:662:LEU:HD22	6:X:662:LEU:N	2.07	0.69
6:X:864:PHE:HA	6:X:920:LEU:HD21	1.75	0.69
6:X:976:ASN:HA	6:X:1083:TYR:CE1	2.28	0.69
6:X:1167:LEU:HD22	6:X:1187:ARG:NH1	2.06	0.69
6:Y:413:PRO:HG3	6:Y:442:CYS:SG	2.32	0.69
6:Y:589:LEU:O	6:Y:593:LEU:HD23	1.92	0.69
1:d:45:ARG:CD	3:D:400:ALA:HA	2.23	0.69
1:p:51:PHE:CZ	1:p:71:CYS:O	2.46	0.69
2:C:39:GLY:O	3:H:84:LYS:CD	2.40	0.69

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:D:50:PRO:HA	3:D:60:SER:O	1.91	0.69
3:D:469:VAL:HG23	3:F:575:MET:CG	2.23	0.69
3:D:564:LEU:HD21	3:D:568:ARG:HH21	1.55	0.69
2:E:37:LYS:NZ	3:H:264:SER:OG	2.23	0.69
3:F:416:ARG:CB	3:H:298:PHE:CE1	2.76	0.69
3:F:517:LEU:HG	3:F:601:THR:HG22	1.74	0.69
2:G:26:MET:HG3	3:H:216:PRO:CG	2.21	0.69
3:J:411:VAL:O	3:J:411:VAL:HG12	1.90	0.69
3:P:298:PHE:CE1	3:T:416:ARG:CB	2.76	0.69
3:P:339:ILE:HD11	3:P:464:LEU:HD22	1.75	0.69
6:X:453:LEU:O	6:X:453:LEU:HG	1.92	0.69
6:X:517:LEU:O	6:X:520:PRO:HD3	1.93	0.69
6:X:832:ILE:HB	6:X:835:HIS:HB3	1.75	0.69
6:X:993:TRP:NE1	6:X:995:PRO:HG3	2.08	0.69
6:Y:427:LEU:HD13	6:Y:837:TYR:HB3	1.74	0.69
6:Y:875:GLN:NE2	6:Y:878:GLY:HA3	2.08	0.69
6:Y:990:ASN:HD22	6:Y:1131:ARG:N	1.90	0.69
1:l:24:LEU:CD2	1:l:80:ARG:HD2	2.22	0.69
3:F:139:THR:CA	3:F:142:ILE:HG22	2.23	0.69
3:F:609:ARG:HH11	2:G:27:THR:CG2	2.05	0.69
3:H:339:ILE:HD13	3:H:349:LEU:HD21	1.75	0.69
3:J:300:LYS:HE3	3:J:303:ASP:OD2	1.91	0.69
3:J:450:SER:HA	3:J:453:ASN:ND2	2.06	0.69
3:J:469:VAL:HG23	3:L:575:MET:CG	2.23	0.69
3:J:517:LEU:HG	3:J:601:THR:HG22	1.74	0.69
3:L:339:ILE:HD11	3:L:464:LEU:HD22	1.75	0.69
2:O:27:THR:CG2	3:T:609:ARG:HH11	2.05	0.69
3:R:582:LEU:HD23	3:R:585:GLN:NE2	2.07	0.69
3:T:360:TYR:HD1	3:T:361:ASP:N	1.91	0.69
4:U:244:ASP:OD2	4:U:278:ILE:HG22	1.93	0.69
5:W:779:MET:HE2	5:W:779:MET:HA	1.74	0.69
6:X:206:LEU:HD22	6:X:206:LEU:N	2.08	0.69
6:X:597:GLU:HG3	6:X:636:ALA:CA	2.22	0.69
1:r:45:ARG:HG2	3:R:400:ALA:CA	2.07	0.69
3:D:84:LYS:CD	2:E:39:GLY:O	2.40	0.69
3:F:450:SER:HA	3:F:453:ASN:ND2	2.07	0.69
3:H:375:LEU:HD23	3:H:437:ILE:CD1	2.15	0.69
3:J:360:TYR:HD1	3:J:361:ASP:N	1.91	0.69
3:N:375:LEU:HD23	3:N:437:ILE:CD1	2.15	0.69
4:U:221:GLY:H	4:U:337:ARG:HG2	1.57	0.69
6:X:790:PRO:HB3	6:X:795:ASP:HB3	1.74	0.69

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
6:X:1097:ALA:HB3	6:X:1098:PRO:HD3	1.73	0.69
6:Y:413:PRO:HG2	6:Y:440:ILE:HD11	1.74	0.69
1:r:8:GLN:CD	3:T:302:GLU:CG	2.65	0.68
1:t:15:ARG:HH22	3:T:402:ALA:HB3	1.57	0.68
3:B:277:ASP:O	3:B:278:THR:CG2	2.37	0.68
3:F:339:ILE:HD13	3:F:349:LEU:HD21	1.75	0.68
3:H:360:TYR:HD1	3:H:361:ASP:N	1.91	0.68
3:H:517:LEU:HG	3:H:601:THR:HG22	1.74	0.68
3:J:274:ILE:CD1	3:J:514:LEU:HD22	2.10	0.68
3:L:322:TYR:CD1	3:L:396:VAL:HB	2.27	0.68
3:L:360:TYR:HD1	3:L:361:ASP:N	1.92	0.68
3:R:416:ARG:CB	3:T:298:PHE:CE1	2.76	0.68
3:R:579:ASN:ND2	3:R:582:LEU:HG	2.06	0.68
3:T:582:LEU:HD23	3:T:585:GLN:NE2	2.07	0.68
5:W:577:LEU:HD23	5:W:579:TYR:H	1.58	0.68
5:W:1063:VAL:HG22	5:W:1110:VAL:HG22	1.74	0.68
6:X:429:ASN:N	6:X:429:ASN:HD22	1.91	0.68
6:X:718:THR:HA	6:X:726:ARG:NH1	2.07	0.68
6:X:940:GLN:NE2	6:X:944:ASN:HD22	1.90	0.68
1:f:17:ALA:O	1:f:20:GLY:CA	2.41	0.68
1:h:45:ARG:CD	3:H:401:GLY:H	2.05	0.68
7:A:101:MYR:H21	3:B:191:LYS:CE	2.23	0.68
3:D:274:ILE:CD1	3:D:514:LEU:HD22	2.10	0.68
3:J:469:VAL:CG2	3:L:575:MET:HB2	2.23	0.68
3:L:237:GLU:CD	3:N:46:LYS:HZ3	2.02	0.68
3:R:411:VAL:HG11	3:T:461:ASP:HB3	1.76	0.68
5:W:261:LEU:O	5:W:264:GLU:HG3	1.94	0.68
5:W:470:MET:HG2	5:W:473:ARG:NH2	2.09	0.68
6:X:355:LYS:O	6:X:358:GLN:HB3	1.93	0.68
6:X:972:MET:HE2	6:X:977:VAL:O	1.94	0.68
6:Y:300:MET:HE1	6:Y:975:PHE:HD2	1.56	0.68
6:Y:606:MET:HE3	6:Y:610:THR:HG21	1.76	0.68
1:f:5:MET:HA	1:f:5:MET:CE	2.10	0.68
1:d:16:ALA:O	1:d:22:LEU:HD23	1.93	0.68
1:n:22:LEU:HG	1:n:83:VAL:HG21	1.75	0.68
3:B:274:ILE:CD1	3:B:514:LEU:HD22	2.10	0.68
2:C:29:THR:CG2	3:H:609:ARG:HB3	2.14	0.68
3:D:48:TRP:CE2	3:D:63:ILE:HG23	2.28	0.68
3:J:339:ILE:HD11	3:J:464:LEU:HD22	1.75	0.68
3:J:575:MET:CG	3:N:469:VAL:HG23	2.23	0.68
3:P:341:MET:HE2	3:P:341:MET:CA	2.18	0.68

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:P:609:ARG:NH1	2:Q:27:THR:CG2	2.56	0.68
3:P:609:ARG:HH11	2:Q:27:THR:CG2	2.05	0.68
3:T:339:ILE:HD13	3:T:349:LEU:HD21	1.75	0.68
6:Y:1213:ALA:O	6:Y:1214:THR:HG23	1.92	0.68
1:b:19:ALA:O	1:b:20:GLY:C	2.35	0.68
1:b:22:LEU:CG	1:b:83:VAL:HG21	2.23	0.68
1:h:22:LEU:HD11	1:h:79:ILE:HG22	1.75	0.68
1:t:60:ALA:HB1	3:T:322:TYR:HE1	1.58	0.68
3:B:360:TYR:HD1	3:B:361:ASP:N	1.91	0.68
2:E:26:MET:HG3	3:F:216:PRO:CG	2.21	0.68
3:H:579:ASN:ND2	3:H:582:LEU:HG	2.06	0.68
3:J:411:VAL:HG11	3:L:461:ASP:HB3	1.76	0.68
3:L:339:ILE:HD13	3:L:349:LEU:HD21	1.75	0.68
7:Q:101:MYR:C2	3:R:191:LYS:HE2	2.23	0.68
4:U:37:TRP:HZ3	4:U:72:LEU:HD11	1.58	0.68
5:W:243:GLN:HB2	6:X:652:GLN:HG2	1.74	0.68
5:W:577:LEU:HD21	5:W:579:TYR:HB2	1.74	0.68
5:W:831:LEU:HD12	5:W:892:SER:O	1.94	0.68
6:Y:777:ASN:HB2	6:Y:942:GLY:HA2	1.74	0.68
6:Y:1060:ARG:NH2	6:Y:1101:GLU:HB3	2.09	0.68
1:f:4:HIS:ND1	1:f:54:THR:HG22	2.09	0.68
1:t:21:ARG:C	3:T:323:SER:HB2	2.12	0.68
1:t:24:LEU:HD21	1:t:35:PHE:HA	1.71	0.68
2:I:39:GLY:O	3:N:84:LYS:CD	2.40	0.68
3:J:298:PHE:CE1	3:N:416:ARG:CB	2.76	0.68
3:L:469:VAL:HG23	3:N:575:MET:CG	2.23	0.68
3:P:575:MET:HE2	3:T:348:VAL:HG21	1.76	0.68
7:Q:101:MYR:H91	7:Q:101:MYR:H52	1.73	0.68
5:W:114:ASN:HB3	5:W:117:VAL:CG1	2.23	0.68
5:W:448:ASP:CG	5:W:673:ALA:HB2	2.18	0.68
5:W:617:VAL:HG13	5:W:658:VAL:HG12	1.76	0.68
3:D:298:PHE:CE1	3:H:416:ARG:CB	2.76	0.68
3:D:339:ILE:HD11	3:D:464:LEU:HD22	1.75	0.68
3:D:517:LEU:HG	3:D:601:THR:HG22	1.74	0.68
3:D:575:MET:HB2	3:H:469:VAL:CG2	2.24	0.68
3:D:575:MET:CG	3:H:469:VAL:HG23	2.23	0.68
3:D:609:ARG:HH11	2:E:27:THR:CG2	2.05	0.68
7:E:101:MYR:C2	3:F:191:LYS:HE2	2.23	0.68
3:J:549:LEU:C	3:J:549:LEU:HD12	2.19	0.68
3:J:575:MET:HB2	3:N:469:VAL:CG2	2.24	0.68
7:K:101:MYR:H31	3:L:191:LYS:CG	2.23	0.68

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:L:416:ARG:CB	3:N:298:PHE:CE1	2.76	0.68
3:N:339:ILE:HD13	3:N:349:LEU:HD21	1.75	0.68
4:U:16:GLN:OE1	4:U:380:ALA:HA	1.93	0.68
5:W:248:LEU:HD13	6:X:628:PRO:HD2	1.76	0.68
5:W:650:VAL:HG12	5:W:651:ASN:H	1.59	0.68
6:X:541:VAL:HG23	6:X:809:ALA:O	1.93	0.68
6:X:792:GLN:HB2	6:X:931:ILE:HD13	1.74	0.68
6:X:1019:THR:HG21	6:Y:429:ASN:HD21	1.58	0.68
1:b:45:ARG:O	1:b:58:PRO:O	2.11	0.68
1:r:59:HIS:CA	3:R:398:SER:CB	2.71	0.68
2:A:26:MET:HG3	3:B:216:PRO:CG	2.21	0.68
3:B:582:LEU:HD23	3:B:585:GLN:NE2	2.08	0.68
3:D:137:ILE:HD13	3:D:142:ILE:HG13	1.76	0.68
3:D:575:MET:HE2	3:H:348:VAL:HG21	1.76	0.68
3:H:413:GLY:O	3:H:415:PRO:HD3	1.94	0.68
3:J:98:ALA:CB	3:J:206:GLU:HG2	2.24	0.68
6:X:1122:PRO:HA	6:X:1159:GLY:HA3	1.74	0.68
1:l:23:THR:HG22	1:l:42:THR:OG1	1.92	0.68
1:d:15:ARG:NH1	3:D:402:ALA:CB	2.56	0.68
1:j:44:GLY:O	3:J:399:ALA:O	2.12	0.68
1:p:15:ARG:NH1	3:P:402:ALA:CB	2.54	0.68
1:t:8:GLN:CG	3:P:302:GLU:HG2	2.17	0.68
3:F:360:TYR:HD1	3:F:361:ASP:N	1.91	0.68
3:L:277:ASP:O	3:L:278:THR:CG2	2.37	0.68
3:N:360:TYR:HD1	3:N:361:ASP:N	1.91	0.68
3:P:545:VAL:CG2	3:R:637:PHE:HE1	2.01	0.68
3:P:549:LEU:C	3:P:549:LEU:HD12	2.19	0.68
5:W:934:ASN:HD22	5:W:934:ASN:H	1.42	0.68
6:X:464:SER:HB2	6:Y:500:ILE:HG22	1.65	0.68
6:X:535:TRP:CE2	6:X:678:PRO:HG3	2.28	0.68
6:Y:294:ARG:HD3	6:Y:885:LEU:O	1.93	0.68
6:Y:996:MET:HB3	6:Y:1143:ASN:HB3	1.75	0.68
1:d:22:LEU:HD11	1:d:83:VAL:HG21	0.74	0.68
1:t:8:GLN:HE22	3:P:302:GLU:C	1.98	0.68
3:D:98:ALA:CB	3:D:206:GLU:HG2	2.24	0.68
3:D:461:ASP:HB3	3:H:411:VAL:HG11	1.76	0.68
3:J:49:ARG:HB2	3:J:64:VAL:HG22	1.76	0.68
3:J:96:PHE:HD1	3:J:101:THR:HG23	1.56	0.68
3:N:294:TYR:HH	3:N:502:LEU:HA	1.57	0.68
3:P:92:TRP:CD1	3:P:156:VAL:CG2	2.77	0.68
3:P:461:ASP:HB3	3:T:411:VAL:HG11	1.76	0.68

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:P:469:VAL:CG2	3:R:575:MET:HB2	2.24	0.68
3:R:469:VAL:HG23	3:T:575:MET:CG	2.23	0.68
3:T:294:TYR:HH	3:T:502:LEU:HA	1.55	0.68
5:W:39:LEU:H	5:W:89:HIS:HE1	1.41	0.68
5:W:929:MET:HE1	5:W:981:LEU:HD11	1.73	0.68
6:X:718:THR:HA	6:X:726:ARG:HH11	1.59	0.68
6:X:850:MET:HE3	6:X:851:PHE:CE1	2.28	0.68
1:n:59:HIS:HB2	3:N:398:SER:HB2	1.75	0.68
1:r:45:ARG:CG	3:R:401:GLY:N	2.40	0.68
2:C:27:THR:O	2:C:27:THR:CG2	2.38	0.68
3:D:469:VAL:CG2	3:F:575:MET:HB2	2.23	0.68
3:F:339:ILE:HD11	3:F:464:LEU:HD22	1.75	0.68
3:P:55:VAL:HG11	5:W:348:GLN:CD	2.19	0.68
3:T:98:ALA:CB	3:T:206:GLU:HG2	2.24	0.68
4:U:211:LEU:HD13	4:U:211:LEU:O	1.93	0.68
5:W:1044:LEU:HD11	5:W:1103:THR:C	2.19	0.68
6:X:633:PRO:HG3	6:X:640:ARG:HH11	1.58	0.68
6:X:1122:PRO:CA	6:X:1159:GLY:HA3	2.24	0.68
6:Y:215:LEU:HD23	6:Y:215:LEU:N	2.09	0.68
1:l:3:LEU:HD13	1:l:4:HIS:O	1.94	0.67
1:d:8:GLN:HG2	3:F:302:GLU:HG3	0.71	0.67
3:D:549:LEU:C	3:D:549:LEU:HD12	2.19	0.67
3:F:413:GLY:O	3:F:415:PRO:HD3	1.94	0.67
3:L:549:LEU:C	3:L:549:LEU:HD12	2.19	0.67
3:P:98:ALA:CB	3:P:206:GLU:HG2	2.24	0.67
3:P:358:LEU:CD1	3:P:417:PHE:CE1	2.62	0.67
3:R:549:LEU:C	3:R:549:LEU:HD12	2.19	0.67
3:T:101:THR:OG1	3:T:102:GLY:N	2.26	0.67
6:X:444:TRP:O	6:X:854:ASN:HB3	1.94	0.67
6:Y:218:GLU:HG3	6:Y:219:LEU:N	2.07	0.67
6:Y:464:SER:HB3	6:Y:804:HIS:NE2	2.08	0.67
1:f:70:ALA:O	1:f:71:CYS:SG	2.51	0.67
1:j:6:ILE:HD11	3:L:586:VAL:HG21	1.76	0.67
1:r:61:ASN:HB2	3:R:396:VAL:O	1.94	0.67
3:B:274:ILE:HD11	3:B:514:LEU:HD21	1.73	0.67
3:B:294:TYR:HH	3:B:502:LEU:HA	1.56	0.67
3:B:413:GLY:O	3:B:415:PRO:HD3	1.94	0.67
3:B:450:SER:HA	3:B:453:ASN:ND2	2.06	0.67
3:D:360:TYR:HD1	3:D:361:ASP:N	1.91	0.67
3:D:411:VAL:HG11	3:F:461:ASP:HB3	1.76	0.67
3:F:469:VAL:HG23	3:H:575:MET:CG	2.23	0.67

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:H:92:TRP:CD1	3:H:156:VAL:CG2	2.77	0.67
3:P:339:ILE:HD13	3:P:349:LEU:HD21	1.75	0.67
3:P:360:TYR:HD1	3:P:361:ASP:N	1.91	0.67
2:Q:26:MET:HG3	3:R:216:PRO:CG	2.21	0.67
3:R:469:VAL:CG2	3:T:575:MET:HB2	2.24	0.67
3:T:450:SER:HA	3:T:453:ASN:ND2	2.06	0.67
6:X:432:ILE:HD13	6:X:432:ILE:O	1.93	0.67
1:f:60:ALA:H	3:F:398:SER:HA	1.59	0.67
1:p:47:THR:CG2	1:p:59:HIS:O	2.41	0.67
3:F:505:ASP:OD1	3:F:508:VAL:HG23	1.95	0.67
3:L:609:ARG:NH1	2:M:27:THR:CG2	2.56	0.67
3:P:575:MET:CG	3:T:469:VAL:HG23	2.23	0.67
3:R:339:ILE:HD13	3:R:349:LEU:HD21	1.75	0.67
3:R:348:VAL:HG21	3:T:575:MET:HE2	1.76	0.67
3:T:549:LEU:C	3:T:549:LEU:HD12	2.19	0.67
4:U:233:TYR:HE2	4:U:251:ASN:HB2	1.58	0.67
4:U:241:LEU:O	6:X:1179:VAL:CG1	2.43	0.67
6:X:347:ARG:HG3	6:X:368:ILE:CD1	2.24	0.67
6:Y:587:MET:HE1	6:Y:601:ILE:HD11	1.76	0.67
6:Y:645:HIS:O	6:Y:649:ASN:HB2	1.93	0.67
6:Y:784:LEU:HD23	6:Y:799:ASP:HB3	1.75	0.67
1:b:61:ASN:HB2	3:B:395:THR:HG23	1.76	0.67
1:h:15:ARG:CZ	3:H:402:ALA:HB1	2.21	0.67
3:F:549:LEU:C	3:F:549:LEU:HD12	2.19	0.67
3:H:505:ASP:OD1	3:H:508:VAL:HG23	1.95	0.67
2:I:27:THR:CG2	3:N:609:ARG:NH1	2.56	0.67
3:L:348:VAL:HG21	3:N:575:MET:HE2	1.76	0.67
3:L:449:THR:HG22	3:L:450:SER:H	1.60	0.67
3:L:525:ILE:CD1	3:L:604:VAL:HG13	2.25	0.67
3:N:525:ILE:CD1	3:N:604:VAL:HG13	2.25	0.67
2:O:27:THR:CG2	3:T:609:ARG:NH1	2.56	0.67
3:P:575:MET:HB2	3:T:469:VAL:CG2	2.24	0.67
3:R:92:TRP:CD1	3:R:156:VAL:CG2	2.77	0.67
3:R:360:TYR:HD1	3:R:361:ASP:N	1.91	0.67
3:T:521:VAL:CG2	3:T:577:VAL:HG22	2.25	0.67
4:U:143:HIS:CE1	4:U:291:THR:HB	2.29	0.67
6:X:269:SER:O	6:X:270:SER:HB3	1.94	0.67
6:X:416:ILE:CG2	6:X:445:PHE:CE2	2.77	0.67
6:Y:217:LYS:O	6:Y:218:GLU:HB3	1.93	0.67
6:Y:705:ILE:CG2	6:Y:737:ARG:HB2	2.24	0.67
1:b:44:GLY:C	3:B:399:ALA:HB3	2.19	0.67

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:f:22:LEU:CG	1:f:83:VAL:HG21	2.22	0.67
1:d:44:GLY:O	3:D:399:ALA:O	2.13	0.67
1:t:61:ASN:ND2	3:T:395:THR:HG22	2.06	0.67
3:B:549:LEU:C	3:B:549:LEU:HD12	2.19	0.67
3:D:525:ILE:CD1	3:D:604:VAL:HG13	2.25	0.67
3:H:339:ILE:HD11	3:H:464:LEU:HD22	1.75	0.67
3:N:413:GLY:O	3:N:415:PRO:HD3	1.94	0.67
3:R:147:ARG:NH1	3:R:147:ARG:HB2	2.10	0.67
3:T:413:GLY:O	3:T:415:PRO:HD3	1.94	0.67
3:T:505:ASP:OD1	3:T:508:VAL:HG23	1.95	0.67
5:W:998:LEU:HD13	5:W:1002:ILE:HD12	1.76	0.67
1:b:21:ARG:H	1:b:22:LEU:HD23	1.58	0.67
1:f:17:ALA:O	1:f:20:GLY:HA2	1.94	0.67
3:B:92:TRP:NE1	3:B:156:VAL:CG2	2.58	0.67
3:B:147:ARG:HB2	3:B:147:ARG:NH1	2.10	0.67
3:B:521:VAL:CG2	3:B:577:VAL:HG22	2.25	0.67
3:H:274:ILE:HD11	3:H:514:LEU:HD21	1.73	0.67
3:H:505:ASP:O	3:H:506:ALA:C	2.38	0.67
3:H:525:ILE:CD1	3:H:604:VAL:HG13	2.25	0.67
3:J:58:ILE:CG1	3:J:127:LEU:HD23	2.24	0.67
3:J:92:TRP:NE1	3:J:156:VAL:CG2	2.58	0.67
3:N:92:TRP:CD1	3:N:156:VAL:CG2	2.77	0.67
3:N:92:TRP:NE1	3:N:156:VAL:CG2	2.58	0.67
3:N:147:ARG:HB2	3:N:147:ARG:NH1	2.10	0.67
3:P:59:ASP:HB2	5:W:380:ASN:CB	2.24	0.67
3:P:274:ILE:HD11	3:P:514:LEU:HD21	1.73	0.67
3:R:92:TRP:NE1	3:R:156:VAL:CG2	2.58	0.67
3:T:92:TRP:CD1	3:T:156:VAL:CG2	2.77	0.67
3:T:92:TRP:NE1	3:T:156:VAL:CG2	2.58	0.67
4:U:31:GLY:O	4:U:32:CYS:SG	2.53	0.67
6:X:675:MET:HE2	6:X:768:GLN:NE2	2.10	0.67
6:X:982:LEU:HD23	6:X:982:LEU:C	2.19	0.67
6:Y:126:PHE:CE2	6:Y:135:HIS:HB2	2.29	0.67
6:Y:549:ALA:HB1	6:Y:815:PRO:HG3	1.75	0.67
1:f:58:PRO:CG	3:F:423:GLN:HG3	2.25	0.67
1:n:8:GLN:HE21	3:J:302:GLU:HG2	1.56	0.67
1:r:24:LEU:HG	1:r:38:ALA:CB	2.25	0.67
3:D:147:ARG:NH1	3:D:147:ARG:HB2	2.10	0.67
3:D:348:VAL:HG21	3:F:575:MET:HE2	1.76	0.67
3:F:92:TRP:NE1	3:F:156:VAL:CG2	2.58	0.67
3:H:521:VAL:CG2	3:H:577:VAL:HG22	2.25	0.67

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:H:549:LEU:C	3:H:549:LEU:HD12	2.19	0.67
3:L:545:VAL:CG2	3:N:637:PHE:HE1	2.01	0.67
3:P:521:VAL:CG2	3:P:577:VAL:HG22	2.25	0.67
3:R:413:GLY:O	3:R:415:PRO:HD3	1.94	0.67
3:T:174:ILE:HG22	3:T:634:LYS:HD2	1.77	0.67
5:W:145:ASP:OD2	5:W:148:ILE:HB	1.95	0.67
5:W:695:ILE:HD11	5:W:1002:ILE:HG22	1.77	0.67
6:Y:483:VAL:HG11	6:Y:841:PHE:CE1	2.29	0.67
6:Y:594:ALA:HA	6:Y:608:GLN:HE22	1.57	0.67
6:Y:1030:GLN:HB2	6:Y:1040:ASP:HA	1.76	0.67
3:B:92:TRP:CD1	3:B:156:VAL:CG2	2.77	0.67
2:C:27:THR:CG2	3:H:609:ARG:NH1	2.56	0.67
3:D:92:TRP:NE1	3:D:156:VAL:CG2	2.58	0.67
3:D:339:ILE:HD13	3:D:349:LEU:HD21	1.75	0.67
3:D:413:GLY:O	3:D:415:PRO:HD3	1.94	0.67
3:D:571:ARG:NH2	3:H:427:GLU:OE1	2.27	0.67
2:E:21:THR:CG2	3:F:219:LYS:HE2	2.25	0.67
3:J:521:VAL:CG2	3:J:577:VAL:HG22	2.25	0.67
3:N:98:ALA:CB	3:N:206:GLU:HG2	2.24	0.67
7:O:101:MYR:H21	3:P:191:LYS:CE	2.23	0.67
3:R:521:VAL:CG2	3:R:577:VAL:HG22	2.25	0.67
2:S:21:THR:CG2	3:T:219:LYS:HE2	2.25	0.67
4:U:78:PRO:HG3	4:U:84:ILE:HA	1.75	0.67
5:W:712:ALA:HB2	5:W:760:TYR:HB3	1.76	0.67
6:X:257:ARG:HB2	6:X:310:LYS:HZ3	1.59	0.67
6:X:583:VAL:HG21	6:X:621:TRP:CD1	2.30	0.67
6:X:1013:VAL:HG12	6:X:1014:VAL:N	2.09	0.67
6:X:1094:LEU:HD23	6:X:1132:ARG:HB3	1.77	0.67
6:Y:164:TRP:NE1	6:Y:165:ASP:HB2	2.09	0.67
1:r:44:GLY:HA3	3:R:322:TYR:CZ	2.25	0.67
1:t:61:ASN:HB2	3:T:397:VAL:HA	1.75	0.67
3:B:333:PRO:HG3	3:B:454:PHE:HA	1.77	0.67
3:F:427:GLU:OE1	3:H:571:ARG:NH2	2.27	0.67
3:F:469:VAL:CG2	3:H:575:MET:HB2	2.24	0.67
3:H:98:ALA:CB	3:H:206:GLU:HG2	2.24	0.67
3:J:92:TRP:CD1	3:J:156:VAL:CG2	2.77	0.67
3:J:147:ARG:HB2	3:J:147:ARG:NH1	2.10	0.67
3:L:147:ARG:NH1	3:L:147:ARG:HB2	2.10	0.67
3:L:505:ASP:O	3:L:506:ALA:C	2.38	0.67
3:L:521:VAL:CG2	3:L:577:VAL:HG22	2.25	0.67
2:M:26:MET:HG3	3:N:216:PRO:CG	2.21	0.67

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:O:21:THR:CG2	3:P:219:LYS:HE2	2.25	0.67
3:R:98:ALA:CB	3:R:206:GLU:HG2	2.24	0.67
3:R:428:ARG:HG2	3:R:428:ARG:NH1	2.03	0.67
4:U:198:GLN:HA	4:U:198:GLN:OE1	1.95	0.67
5:W:891:ASP:HA	5:W:925:GLY:O	1.95	0.67
6:X:387:CYS:O	6:X:1194:LEU:HD23	1.93	0.67
6:Y:464:SER:OG	6:Y:800:ILE:HG23	1.94	0.67
6:Y:1108:ASN:HD21	6:Y:1110:ALA:HB3	1.60	0.67
1:b:45:ARG:CG	3:B:400:ALA:C	2.55	0.67
1:h:23:THR:CG2	3:H:323:SER:CA	2.69	0.67
1:p:44:GLY:C	3:P:399:ALA:HB3	2.20	0.67
3:D:92:TRP:CD1	3:D:156:VAL:CG2	2.77	0.67
3:D:505:ASP:OD1	3:D:508:VAL:HG23	1.95	0.67
3:F:98:ALA:CB	3:F:206:GLU:HG2	2.24	0.67
3:H:92:TRP:NE1	3:H:156:VAL:CG2	2.58	0.67
3:J:418:ASN:OD1	3:L:337:ARG:NH2	2.28	0.67
3:R:505:ASP:OD1	3:R:508:VAL:HG23	1.95	0.67
3:T:277:ASP:O	3:T:278:THR:CG2	2.37	0.67
4:U:40:ARG:NE	4:U:144:ILE:HG21	2.09	0.67
4:V:192:PHE:CA	6:Y:376:LEU:HD23	2.25	0.67
5:W:38:ASN:HA	5:W:89:HIS:HE1	1.59	0.67
5:W:74:PHE:CE1	5:W:79:LEU:HD21	2.30	0.67
5:W:934:ASN:HB2	5:W:956:TYR:CE2	2.30	0.67
6:X:653:PRO:HD3	6:X:773:ARG:HG3	1.76	0.67
6:X:864:PHE:HE2	6:X:952:PRO:HG3	1.57	0.67
6:X:872:ALA:HB3	6:X:881:VAL:HG21	1.75	0.67
1:l:24:LEU:CD2	1:l:80:ARG:HD3	2.24	0.66
3:B:174:ILE:HG22	3:B:634:LYS:HD2	1.77	0.66
3:D:52:GLY:O	3:D:53:THR:HB	1.95	0.66
3:D:418:ASN:OD1	3:F:337:ARG:NH2	2.28	0.66
3:F:49:ARG:HB2	3:F:64:VAL:CG2	2.25	0.66
3:F:92:TRP:CD1	3:F:156:VAL:CG2	2.77	0.66
3:F:348:VAL:HG21	3:H:575:MET:HE2	1.76	0.66
3:F:521:VAL:CG2	3:F:577:VAL:HG22	2.25	0.66
3:H:174:ILE:HG22	3:H:634:LYS:HD2	1.77	0.66
7:I:101:MYR:H21	3:J:191:LYS:CE	2.23	0.66
3:N:181:CYS:HB2	3:N:638:PHE:HB3	1.77	0.66
3:P:411:VAL:HG11	3:R:461:ASP:HB3	1.76	0.66
3:P:413:GLY:O	3:P:415:PRO:HD3	1.94	0.66
7:Q:101:MYR:H31	3:R:191:LYS:CG	2.23	0.66
3:R:449:THR:HG22	3:R:450:SER:H	1.60	0.66

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
4:U:160:GLU:O	4:U:161:VAL:HG13	1.94	0.66
5:W:695:ILE:HG12	5:W:708:LEU:HD12	1.75	0.66
6:X:295:SER:HA	6:X:874:CYS:SG	2.34	0.66
6:X:958:HIS:CD2	6:X:958:HIS:O	2.48	0.66
6:X:1006:ARG:HG3	6:X:1007:GLY:N	2.07	0.66
6:Y:317:PHE:CZ	6:Y:385:VAL:HG11	2.30	0.66
6:Y:486:LEU:HD12	6:Y:746:TYR:CD2	2.30	0.66
1:l:15:ARG:CZ	3:L:402:ALA:HB2	2.26	0.66
1:l:15:ARG:HH22	3:L:402:ALA:HB3	1.60	0.66
1:h:21:ARG:O	3:H:323:SER:HB2	1.95	0.66
1:p:44:GLY:O	3:P:399:ALA:C	2.38	0.66
1:r:22:LEU:HG	1:r:83:VAL:HG21	1.76	0.66
3:D:56:ALA:HB1	3:D:61:LEU:CD2	2.24	0.66
3:F:147:ARG:HB2	3:F:147:ARG:NH1	2.10	0.66
2:G:21:THR:CG2	3:H:219:LYS:HE2	2.25	0.66
3:J:337:ARG:NH2	3:N:418:ASN:OD1	2.28	0.66
3:J:413:GLY:O	3:J:415:PRO:HD3	1.94	0.66
3:J:575:MET:HE2	3:N:348:VAL:HG21	1.76	0.66
3:J:630:LEU:HD21	3:N:276:ALA:HA	1.78	0.66
3:L:49:ARG:HB2	3:L:64:VAL:CG2	2.26	0.66
3:L:92:TRP:CD1	3:L:156:VAL:CG2	2.77	0.66
3:L:469:VAL:CG2	3:N:575:MET:HB2	2.24	0.66
3:N:174:ILE:HG22	3:N:634:LYS:HD2	1.77	0.66
3:N:428:ARG:HG2	3:N:428:ARG:NH1	2.03	0.66
3:P:49:ARG:HB2	3:P:64:VAL:CG2	2.26	0.66
4:U:150:LEU:HD11	4:U:315:LEU:HB3	1.77	0.66
6:Y:384:MET:O	6:Y:384:MET:HG3	1.95	0.66
1:b:45:ARG:CD	3:B:400:ALA:CA	2.62	0.66
1:b:51:PHE:O	1:b:74:GLN:CG	2.42	0.66
1:f:45:ARG:CD	3:F:400:ALA:HA	2.24	0.66
3:F:609:ARG:NH1	2:G:27:THR:CG2	2.56	0.66
3:L:92:TRP:NE1	3:L:156:VAL:CG2	2.58	0.66
3:L:174:ILE:HG22	3:L:634:LYS:HD2	1.77	0.66
3:L:309:GLU:OE1	3:L:335:PRO:CB	2.44	0.66
3:L:413:GLY:O	3:L:415:PRO:HD3	1.94	0.66
3:N:322:TYR:CE2	3:N:327:TYR:CE2	2.83	0.66
5:W:6:GLY:HA3	5:W:339:ARG:O	1.95	0.66
5:W:25:LEU:HD11	5:W:29:LEU:HB3	1.77	0.66
5:W:224:PRO:HG2	5:W:244:THR:HG22	1.76	0.66
5:W:709:GLY:HA3	5:W:763:ARG:NH2	2.11	0.66
6:X:282:TYR:CE2	6:Y:830:ASN:ND2	2.62	0.66

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
6:Y:54:VAL:HB	6:Y:57:GLN:HE21	1.61	0.66
6:Y:750:MET:HB3	6:Y:837:TYR:OH	1.96	0.66
1:f:24:LEU:HD12	1:f:24:LEU:H	1.58	0.66
1:f:45:ARG:CD	3:F:401:GLY:H	2.07	0.66
1:d:61:ASN:HB2	3:D:395:THR:CG2	2.26	0.66
1:h:8:GLN:NE2	3:D:302:GLU:CG	2.57	0.66
1:n:8:GLN:HG2	3:J:302:GLU:HG3	0.76	0.66
1:p:45:ARG:CD	3:P:401:GLY:N	2.59	0.66
1:t:61:ASN:HB3	3:T:397:VAL:HA	1.76	0.66
3:D:593:GLN:HB2	3:D:596:MET:HE2	1.78	0.66
3:F:411:VAL:HG11	3:H:461:ASP:HB3	1.76	0.66
3:H:181:CYS:HB2	3:H:638:PHE:HB3	1.78	0.66
3:J:333:PRO:HG3	3:J:454:PHE:HA	1.77	0.66
3:L:344:MET:CE	3:L:502:LEU:HD11	2.26	0.66
3:L:411:VAL:HG11	3:N:461:ASP:HB3	1.76	0.66
3:P:505:ASP:O	3:P:506:ALA:C	2.38	0.66
3:P:525:ILE:CD1	3:P:604:VAL:HG13	2.25	0.66
3:R:525:ILE:CD1	3:R:604:VAL:HG13	2.25	0.66
3:R:593:GLN:HB2	3:R:596:MET:HE2	1.78	0.66
4:U:378:PRO:O	4:U:379:ILE:HB	1.95	0.66
4:V:169:ALA:HB2	6:Y:1179:VAL:O	1.95	0.66
5:W:969:ASP:HB2	5:W:972:PRO:CD	2.18	0.66
6:Y:231:MET:HE3	6:Y:347:ARG:NH1	2.08	0.66
6:Y:709:ILE:HA	6:Y:733:MET:HE1	1.77	0.66
1:b:15:ARG:CZ	3:B:402:ALA:HB2	2.21	0.66
1:f:59:HIS:HA	3:F:398:SER:OG	1.95	0.66
1:n:45:ARG:CD	3:N:401:GLY:N	2.58	0.66
3:B:98:ALA:CB	3:B:206:GLU:HG2	2.24	0.66
3:B:449:THR:HG22	3:B:450:SER:H	1.60	0.66
3:D:337:ARG:NH2	3:H:418:ASN:OD1	2.28	0.66
3:D:521:VAL:CG2	3:D:577:VAL:HG22	2.25	0.66
3:F:276:ALA:HA	3:H:630:LEU:HD21	1.78	0.66
3:J:525:ILE:CD1	3:J:604:VAL:HG13	2.25	0.66
3:L:505:ASP:OD1	3:L:508:VAL:HG23	1.95	0.66
3:N:549:LEU:C	3:N:549:LEU:HD12	2.19	0.66
3:P:92:TRP:NE1	3:P:156:VAL:CG2	2.58	0.66
3:P:147:ARG:NH1	3:P:147:ARG:HB2	2.10	0.66
3:R:174:ILE:HG22	3:R:634:LYS:HD2	1.77	0.66
3:R:277:ASP:O	3:R:278:THR:CG2	2.37	0.66
3:T:147:ARG:NH1	3:T:147:ARG:HB2	2.09	0.66
3:T:333:PRO:HG3	3:T:454:PHE:HA	1.77	0.66

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:T:525:ILE:CD1	3:T:604:VAL:HG13	2.25	0.66
4:U:150:LEU:HD13	4:U:300:ILE:HD11	1.77	0.66
4:U:227:ARG:HG2	4:U:256:ALA:HB2	1.76	0.66
6:X:726:ARG:HA	6:X:729:MET:HE3	1.77	0.66
6:X:792:GLN:CG	6:X:931:ILE:HG23	2.25	0.66
6:X:979:ARG:HG3	6:X:980:GLY:H	1.58	0.66
6:Y:632:ASN:HB3	6:Y:634:GLN:HE22	1.59	0.66
6:Y:678:PRO:HG3	6:Y:771:PHE:CZ	2.30	0.66
6:Y:690:ALA:HB3	6:Y:691:PRO:CD	2.23	0.66
1:d:22:LEU:CG	1:d:83:VAL:HG21	2.24	0.66
2:A:35:ASP:OD1	2:A:40:VAL:HB	1.96	0.66
2:C:35:ASP:OD1	2:C:40:VAL:HB	1.96	0.66
3:D:630:LEU:HD21	3:H:276:ALA:HA	1.78	0.66
3:F:277:ASP:O	3:F:278:THR:CG2	2.37	0.66
3:H:49:ARG:HB2	3:H:64:VAL:CG2	2.26	0.66
2:I:21:THR:CG2	3:J:219:LYS:HE2	2.25	0.66
3:J:609:ARG:HD2	2:K:29:THR:HG21	1.78	0.66
2:K:35:ASP:OD1	2:K:40:VAL:HB	1.96	0.66
3:P:276:ALA:HA	3:R:630:LEU:HD21	1.78	0.66
3:P:348:VAL:HG21	3:R:575:MET:HE2	1.76	0.66
4:V:191:ARG:HH12	6:Y:438:PRO:CG	2.09	0.66
5:W:1196:LEU:HD11	5:W:1229:ILE:HD12	1.76	0.66
6:X:205:MET:HB3	6:X:249:TRP:CD2	2.30	0.66
1:l:24:LEU:CD1	1:l:32:THR:C	2.64	0.66
1:b:45:ARG:HD2	3:B:400:ALA:CA	2.25	0.66
1:j:15:ARG:CZ	3:J:402:ALA:HB1	2.25	0.66
1:r:44:GLY:O	1:r:60:ALA:HB2	1.96	0.66
1:t:45:ARG:CD	3:T:401:GLY:N	2.59	0.66
3:B:50:PRO:HA	3:B:60:SER:O	1.96	0.66
3:B:63:ILE:HG21	3:B:70:GLN:OE1	1.96	0.66
3:D:280:LEU:HD12	3:D:486:PRO:HG2	1.78	0.66
3:D:449:THR:HG22	3:D:450:SER:H	1.60	0.66
3:F:48:TRP:N	3:F:48:TRP:CD1	2.64	0.66
3:F:418:ASN:OD1	3:H:337:ARG:NH2	2.29	0.66
2:I:35:ASP:OD1	2:I:40:VAL:HB	1.96	0.66
3:J:461:ASP:HB3	3:N:411:VAL:HG11	1.76	0.66
3:J:571:ARG:NH2	3:N:427:GLU:OE1	2.27	0.66
3:L:50:PRO:HA	3:L:60:SER:O	1.96	0.66
3:L:98:ALA:CB	3:L:206:GLU:HG2	2.24	0.66
3:L:181:CYS:HB2	3:L:638:PHE:HB3	1.78	0.66
3:L:609:ARG:HD2	2:M:29:THR:HG21	1.78	0.66

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:M:21:THR:CG2	3:N:219:LYS:HE2	2.25	0.66
3:N:63:ILE:HG21	3:N:70:GLN:OE1	1.96	0.66
3:N:141:GLN:HA	3:N:141:GLN:NE2	2.11	0.66
3:N:521:VAL:CG2	3:N:577:VAL:HG22	2.25	0.66
3:R:309:GLU:OE1	3:R:335:PRO:CB	2.44	0.66
3:R:380:VAL:HG13	3:R:384:GLN:HG2	1.78	0.66
3:T:50:PRO:HA	3:T:60:SER:O	1.96	0.66
5:W:840:CYS:SG	5:W:854:MET:CG	2.83	0.66
5:W:1215:LEU:C	5:W:1215:LEU:HD12	2.21	0.66
6:X:884:PRO:HD3	6:X:962:GLN:HE22	1.61	0.66
6:X:1172:LEU:HB2	6:X:1191:TYR:CD2	2.30	0.66
6:Y:385:VAL:HG22	6:Y:386:GLU:H	1.60	0.66
6:Y:848:ASN:N	6:Y:848:ASN:HD22	1.92	0.66
1:p:16:ALA:O	1:p:22:LEU:CD2	2.44	0.66
3:D:609:ARG:NH1	2:E:27:THR:CG2	2.56	0.66
2:E:35:ASP:OD1	2:E:40:VAL:HB	1.96	0.66
3:H:147:ARG:NH1	3:H:147:ARG:HB2	2.10	0.66
3:H:280:LEU:HD12	3:H:486:PRO:HG2	1.78	0.66
3:J:505:ASP:OD1	3:J:508:VAL:HG23	1.95	0.66
3:L:48:TRP:N	3:L:48:TRP:CD1	2.64	0.66
3:L:216:PRO:HB3	3:L:234:PRO:HG3	1.78	0.66
3:L:428:ARG:HG2	3:L:428:ARG:NH1	2.03	0.66
2:O:26:MET:HG3	3:P:216:PRO:CG	2.21	0.66
3:P:309:GLU:OE1	3:P:335:PRO:CB	2.44	0.66
3:P:344:MET:CE	3:P:502:LEU:HD11	2.26	0.66
3:P:449:THR:HG22	3:P:450:SER:H	1.60	0.66
7:Q:101:MYR:C2	3:R:191:LYS:CE	2.73	0.66
3:T:48:TRP:N	3:T:48:TRP:CD1	2.64	0.66
4:U:198:GLN:NE2	6:X:662:LEU:HD13	2.11	0.66
6:X:856:ARG:HH22	6:X:917:GLY:C	2.04	0.66
6:X:1182:ASN:HD21	6:X:1184:LEU:HB2	1.59	0.66
6:Y:415:GLN:NE2	6:Y:662:LEU:HD23	2.10	0.66
6:Y:981:ASN:HD21	6:Y:1077:GLU:HG2	1.60	0.66
1:j:59:HIS:ND1	3:J:397:VAL:HG22	2.11	0.66
1:j:59:HIS:CA	3:J:398:SER:OG	2.44	0.66
1:p:51:PHE:CE2	1:p:71:CYS:O	2.49	0.66
3:B:181:CYS:HB2	3:B:638:PHE:HB3	1.78	0.66
3:D:68:PHE:HB3	3:D:140:VAL:HG11	1.76	0.66
3:F:181:CYS:HB2	3:F:638:PHE:HB3	1.78	0.66
3:F:380:VAL:HG13	3:F:384:GLN:HG2	1.78	0.66
2:I:29:THR:HG21	3:N:609:ARG:HD2	1.78	0.66

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:P:337:ARG:NH2	3:T:418:ASN:OD1	2.28	0.66
3:P:505:ASP:OD1	3:P:508:VAL:HG23	1.95	0.66
3:P:593:GLN:HB2	3:P:596:MET:HE2	1.78	0.66
3:R:427:GLU:OE1	3:T:571:ARG:NH2	2.27	0.66
3:T:593:GLN:HB2	3:T:596:MET:HE2	1.78	0.66
4:U:40:ARG:HE	4:U:144:ILE:HG21	1.60	0.66
4:V:184:HIS:CE1	6:Y:436:LEU:HD13	2.30	0.66
5:W:436:GLY:HA3	5:W:1009:ASN:CA	2.25	0.66
6:X:330:ILE:HD12	6:X:340:LEU:HD11	1.78	0.66
1:n:44:GLY:O	3:N:399:ALA:CA	2.43	0.66
3:B:505:ASP:O	3:B:506:ALA:C	2.38	0.66
3:B:593:GLN:HB2	3:B:596:MET:HE2	1.78	0.66
2:C:21:THR:CG2	3:D:219:LYS:HE2	2.25	0.66
3:D:56:ALA:CB	3:D:61:LEU:HD21	2.26	0.66
3:D:181:CYS:HB2	3:D:638:PHE:HB3	1.78	0.66
7:E:101:MYR:C2	3:F:191:LYS:CE	2.74	0.66
3:F:311:ASN:ND2	3:F:311:ASN:H	1.94	0.66
3:F:341:MET:HB2	3:F:431:ILE:CD1	2.26	0.66
3:J:280:LEU:HD12	3:J:486:PRO:HG2	1.78	0.66
7:K:101:MYR:C2	3:L:191:LYS:HE2	2.23	0.66
3:N:71:TYR:CE1	3:N:138:SER:HB3	2.31	0.66
3:N:505:ASP:OD1	3:N:508:VAL:HG23	1.95	0.66
3:P:216:PRO:HB3	3:P:234:PRO:HG3	1.78	0.66
3:P:427:GLU:OE1	3:R:571:ARG:NH2	2.27	0.66
3:P:571:ARG:NH2	3:T:427:GLU:OE1	2.27	0.66
2:Q:21:THR:CG2	3:R:219:LYS:HE2	2.25	0.66
3:R:71:TYR:CE1	3:R:138:SER:HB3	2.31	0.66
3:R:609:ARG:NH1	2:S:27:THR:CG2	2.56	0.66
3:T:344:MET:CE	3:T:502:LEU:HD11	2.26	0.66
3:T:449:THR:HG22	3:T:450:SER:H	1.60	0.66
4:U:178:THR:CG2	6:X:376:LEU:HD21	2.12	0.66
5:W:440:THR:HG21	5:W:492:ALA:HB1	1.77	0.66
5:W:1040:VAL:HB	5:W:1041:PRO:CD	2.20	0.66
6:X:424:ALA:O	6:X:427:LEU:HG	1.96	0.66
6:X:1030:GLN:HB3	6:X:1038:PRO:HB2	1.77	0.66
6:Y:394:LEU:HD11	6:Y:964:VAL:HG22	1.77	0.66
1:j:5:MET:HE2	1:j:5:MET:HA	1.77	0.65
1:n:51:PHE:CD1	1:n:71:CYS:O	2.49	0.65
1:r:23:THR:OG1	3:R:323:SER:HA	1.94	0.65
1:t:46:TYR:HD1	1:t:58:PRO:HA	1.61	0.65
2:A:21:THR:CG2	3:B:219:LYS:HE2	2.25	0.65

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:B:525:ILE:CD1	3:B:604:VAL:HG13	2.25	0.65
3:F:344:MET:CE	3:F:502:LEU:HD11	2.26	0.65
3:F:358:LEU:CG	3:F:417:PHE:CE1	2.79	0.65
3:F:525:ILE:CD1	3:F:604:VAL:HG13	2.25	0.65
3:H:63:ILE:HG21	3:H:70:GLN:OE1	1.96	0.65
3:H:449:THR:HG22	3:H:450:SER:H	1.60	0.65
3:J:63:ILE:HG21	3:J:70:GLN:OE1	1.96	0.65
3:L:63:ILE:HG21	3:L:70:GLN:OE1	1.96	0.65
3:P:418:ASN:OD1	3:R:337:ARG:NH2	2.28	0.65
3:R:358:LEU:CG	3:R:417:PHE:CE1	2.79	0.65
3:T:216:PRO:HB3	3:T:234:PRO:HG3	1.78	0.65
4:U:284:ALA:O	4:U:288:GLN:HG3	1.95	0.65
4:V:210:TRP:O	4:V:214:ILE:HB	1.96	0.65
6:X:627:ASN:HD22	6:X:630:LEU:HG	1.60	0.65
6:X:633:PRO:O	6:X:637:PRO:HD3	1.96	0.65
6:Y:627:ASN:HD21	6:Y:629:GLN:HB2	1.61	0.65
6:Y:923:LEU:HD23	6:Y:931:ILE:HD13	1.77	0.65
1:h:15:ARG:NH2	3:H:402:ALA:HB3	2.10	0.65
1:n:59:HIS:HA	3:N:398:SER:HB2	1.78	0.65
3:B:505:ASP:OD1	3:B:508:VAL:HG23	1.95	0.65
3:D:216:PRO:HB3	3:D:234:PRO:HG3	1.78	0.65
3:D:333:PRO:HG3	3:D:454:PHE:HA	1.77	0.65
3:H:358:LEU:CG	3:H:417:PHE:CE1	2.79	0.65
3:H:428:ARG:HG2	3:H:428:ARG:NH1	2.03	0.65
3:J:339:ILE:HD13	3:J:349:LEU:HD21	1.75	0.65
3:J:449:THR:HG22	3:J:450:SER:H	1.60	0.65
3:J:593:GLN:HB2	3:J:596:MET:HE2	1.78	0.65
2:K:21:THR:CG2	3:L:219:LYS:HE2	2.25	0.65
2:K:37:LYS:NZ	3:N:264:SER:OG	2.23	0.65
3:P:57:THR:O	3:P:57:THR:CG2	2.44	0.65
3:P:280:LEU:HD12	3:P:486:PRO:HG2	1.78	0.65
3:P:333:PRO:HG3	3:P:454:PHE:HA	1.77	0.65
2:Q:37:LYS:NZ	3:T:264:SER:OG	2.23	0.65
3:R:418:ASN:OD1	3:T:337:ARG:NH2	2.28	0.65
3:T:141:GLN:NE2	3:T:141:GLN:HA	2.11	0.65
4:U:211:LEU:HD21	4:U:248:LEU:HD22	1.78	0.65
4:V:160:GLU:O	4:V:161:VAL:HG22	1.95	0.65
6:Y:626:LEU:HD11	6:Y:776:GLN:HE22	1.60	0.65
1:t:58:PRO:CG	3:T:423:GLN:HE21	2.09	0.65
3:B:49:ARG:HB2	3:B:64:VAL:CG2	2.26	0.65
3:B:280:LEU:HD12	3:B:486:PRO:HG2	1.78	0.65

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:H:344:MET:CE	3:H:502:LEU:HD11	2.26	0.65
3:J:276:ALA:HA	3:L:630:LEU:HD21	1.77	0.65
3:L:333:PRO:HG3	3:L:454:PHE:HA	1.77	0.65
3:N:311:ASN:ND2	3:N:311:ASN:H	1.94	0.65
3:P:181:CYS:HB2	3:P:638:PHE:HB3	1.78	0.65
4:U:96:VAL:HG13	4:U:135:MET:HE2	1.78	0.65
4:U:140:GLN:HE22	4:U:190:TYR:HE1	1.44	0.65
4:U:168:ALA:C	4:U:170:THR:H	2.01	0.65
5:W:934:ASN:HB2	5:W:956:TYR:HE2	1.60	0.65
1:p:23:THR:HG22	1:p:42:THR:OG1	1.96	0.65
1:t:24:LEU:HD23	1:t:38:ALA:HB3	1.78	0.65
3:B:341:MET:HE2	3:B:341:MET:CA	2.18	0.65
3:B:344:MET:CE	3:B:502:LEU:HD11	2.26	0.65
3:F:593:GLN:HB2	3:F:596:MET:HE2	1.78	0.65
3:J:174:ILE:HG22	3:J:634:LYS:HD2	1.77	0.65
3:J:344:MET:CE	3:J:502:LEU:HD11	2.26	0.65
3:J:348:VAL:HG21	3:L:575:MET:HE2	1.76	0.65
3:J:358:LEU:CG	3:J:417:PHE:CE1	2.79	0.65
3:P:63:ILE:HG21	3:P:70:GLN:OE1	1.96	0.65
3:P:609:ARG:HD2	2:Q:29:THR:HG21	1.78	0.65
3:R:274:ILE:HD11	3:R:514:LEU:HD21	1.73	0.65
3:R:333:PRO:HG3	3:R:454:PHE:HA	1.77	0.65
3:T:63:ILE:HG21	3:T:70:GLN:OE1	1.96	0.65
6:X:600:ALA:C	6:X:623:PRO:HD2	2.20	0.65
1:j:61:ASN:OD1	3:J:395:THR:CG2	2.45	0.65
1:p:15:ARG:HE	3:P:406:ASN:CG	2.05	0.65
3:D:341:MET:HB2	3:D:431:ILE:CD1	2.26	0.65
3:F:280:LEU:HD12	3:F:486:PRO:HG2	1.78	0.65
3:F:609:ARG:HD2	2:G:29:THR:HG21	1.78	0.65
3:H:50:PRO:HA	3:H:60:SER:O	1.96	0.65
3:J:341:MET:HB2	3:J:431:ILE:CD1	2.26	0.65
3:L:579:ASN:ND2	3:L:582:LEU:HG	2.06	0.65
3:N:49:ARG:HB2	3:N:64:VAL:CG2	2.26	0.65
3:N:344:MET:CE	3:N:502:LEU:HD11	2.26	0.65
3:P:51:VAL:HB	5:W:379:VAL:CG2	2.27	0.65
3:P:174:ILE:HG22	3:P:634:LYS:HD2	1.77	0.65
3:P:277:ASP:O	3:P:278:THR:CG2	2.37	0.65
3:P:311:ASN:ND2	3:P:311:ASN:H	1.94	0.65
3:R:311:ASN:ND2	3:R:311:ASN:H	1.94	0.65
2:S:28:SER:CA	3:T:216:PRO:HD2	2.27	0.65
4:U:121:ASP:C	4:U:123:ASN:H	2.01	0.65

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
4:U:198:GLN:HE21	6:X:662:LEU:CD1	2.09	0.65
4:V:224:ASN:HD22	4:V:224:ASN:H	1.44	0.65
6:X:397:HIS:HB2	6:X:1184:LEU:HD21	1.79	0.65
6:X:697:MET:CG	6:X:744:PRO:HA	2.25	0.65
6:X:718:THR:CG2	6:X:726:ARG:HH12	2.03	0.65
6:Y:246:MET:HE2	6:Y:262:ARG:HD2	1.79	0.65
6:Y:586:PHE:HA	6:Y:642:PHE:HE2	1.62	0.65
6:Y:633:PRO:HG2	6:Y:634:GLN:OE1	1.95	0.65
1:d:60:ALA:N	3:D:398:SER:HA	2.11	0.65
1:j:23:THR:N	3:J:323:SER:OG	2.30	0.65
1:p:45:ARG:CD	3:P:400:ALA:CA	2.69	0.65
3:B:311:ASN:ND2	3:B:311:ASN:H	1.94	0.65
3:B:579:ASN:ND2	3:B:582:LEU:HG	2.06	0.65
3:D:344:MET:CE	3:D:502:LEU:HD11	2.26	0.65
3:D:380:VAL:HG13	3:D:384:GLN:HG2	1.78	0.65
3:D:575:MET:CE	3:H:469:VAL:CG2	2.43	0.65
2:E:28:SER:CA	3:F:216:PRO:HD2	2.27	0.65
3:F:579:ASN:ND2	3:F:582:LEU:HG	2.06	0.65
2:I:26:MET:HG3	2:I:26:MET:O	1.97	0.65
7:K:101:MYR:C2	3:L:191:LYS:CE	2.73	0.65
2:M:28:SER:CA	3:N:216:PRO:HD2	2.27	0.65
3:N:216:PRO:HB3	3:N:234:PRO:HG3	1.78	0.65
3:N:309:GLU:OE1	3:N:335:PRO:CB	2.44	0.65
3:P:71:TYR:CE1	3:P:138:SER:HB3	2.31	0.65
2:Q:35:ASP:OD1	2:Q:40:VAL:HB	1.96	0.65
3:T:67:ARG:CZ	3:T:67:ARG:HB2	2.27	0.65
3:T:280:LEU:HD12	3:T:486:PRO:HG2	1.78	0.65
3:T:358:LEU:CG	3:T:417:PHE:CE1	2.79	0.65
4:U:46:ILE:HD12	6:X:1197:TYR:HE2	1.62	0.65
4:U:136:TRP:HB3	4:U:137:PRO:HD3	1.78	0.65
4:U:150:LEU:HD13	4:U:300:ILE:CD1	2.27	0.65
5:W:650:VAL:HG12	5:W:651:ASN:N	2.11	0.65
6:Y:1015:GLY:CA	6:Y:1034:ASP:HA	2.26	0.65
1:l:45:ARG:CD	3:L:401:GLY:N	2.60	0.65
1:b:21:ARG:HD2	3:B:327:TYR:HH	1.60	0.65
1:d:24:LEU:HD21	1:d:80:ARG:HD2	1.79	0.65
1:d:50:ALA:O	1:d:74:GLN:HG2	1.97	0.65
1:h:45:ARG:HG2	3:H:399:ALA:O	1.97	0.65
1:j:23:THR:HG23	3:J:323:SER:C	1.96	0.65
3:D:505:ASP:O	3:D:506:ALA:C	2.38	0.65
3:F:174:ILE:HG22	3:F:634:LYS:HD2	1.77	0.65

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
7:G:101:MYR:H31	3:H:191:LYS:CG	2.26	0.65
3:L:67:ARG:CZ	3:L:67:ARG:HB2	2.27	0.65
3:N:50:PRO:HA	3:N:60:SER:O	1.96	0.65
3:N:333:PRO:HG3	3:N:454:PHE:HA	1.77	0.65
3:P:566:TRP:CZ2	3:R:195:PRO:HD3	2.32	0.65
3:P:630:LEU:HD21	3:T:276:ALA:HA	1.78	0.65
2:Q:26:MET:HG3	2:Q:26:MET:O	1.97	0.65
3:R:50:PRO:HA	3:R:60:SER:O	1.96	0.65
3:T:49:ARG:HB2	3:T:64:VAL:CG2	2.26	0.65
3:T:341:MET:HB2	3:T:431:ILE:CD1	2.26	0.65
4:U:224:ASN:ND2	4:U:226:SER:HB3	2.12	0.65
5:W:439:PHE:HD2	5:W:684:ARG:NH1	1.95	0.65
6:X:708:PHE:O	6:X:733:MET:HE1	1.97	0.65
6:X:1050:ASP:OD2	6:Y:1213:ALA:HB3	1.97	0.65
1:d:8:GLN:NE2	3:F:302:GLU:CG	2.57	0.65
1:n:21:ARG:HE	3:N:370:VAL:HG11	1.62	0.65
1:p:8:GLN:NE2	3:R:303:ASP:HA	2.04	0.65
1:r:3:LEU:HD11	3:T:492:ASN:HD21	1.60	0.65
1:r:8:GLN:HE21	3:T:302:GLU:HG2	1.62	0.65
2:C:29:THR:HG21	3:H:609:ARG:HD2	1.78	0.65
3:D:92:TRP:CH2	3:D:160:GLN:HA	2.32	0.65
3:D:294:TYR:HH	3:D:502:LEU:HA	1.61	0.65
3:D:510:VAL:HG12	3:D:510:VAL:O	1.97	0.65
3:F:50:PRO:HA	3:F:60:SER:O	1.96	0.65
3:F:92:TRP:CH2	3:F:160:GLN:HA	2.32	0.65
3:F:216:PRO:HB3	3:F:234:PRO:HG3	1.78	0.65
3:F:333:PRO:HG3	3:F:454:PHE:HA	1.77	0.65
3:F:510:VAL:HG12	3:F:510:VAL:O	1.97	0.65
3:H:380:VAL:HG13	3:H:384:GLN:HG2	1.78	0.65
3:J:53:THR:O	3:J:54:SER:CB	2.43	0.65
3:J:92:TRP:CH2	3:J:160:GLN:HA	2.32	0.65
2:K:28:SER:CA	3:L:216:PRO:HD2	2.27	0.65
3:L:418:ASN:OD1	3:N:337:ARG:NH2	2.29	0.65
3:L:566:TRP:CZ2	3:N:195:PRO:HD3	2.32	0.65
3:R:49:ARG:HB2	3:R:64:VAL:CG2	2.26	0.65
3:T:510:VAL:HG12	3:T:510:VAL:O	1.97	0.65
4:U:46:ILE:HD11	6:X:384:MET:SD	2.36	0.65
4:U:241:LEU:O	4:U:242:ASN:HB3	1.94	0.65
5:W:222:ALA:HB3	6:X:655:GLU:HG3	1.78	0.65
5:W:727:LEU:HB3	5:W:728:PRO:HD3	1.77	0.65
6:X:931:ILE:HD11	6:Y:689:VAL:HG12	1.78	0.65

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
6:Y:544:VAL:O	6:Y:813:VAL:HG12	1.97	0.65
6:Y:582:PRO:HG3	6:Y:650:TRP:CE2	2.32	0.65
1:j:15:ARG:NH1	3:J:402:ALA:CB	2.60	0.65
1:j:16:ALA:O	1:j:19:ALA:O	2.15	0.65
1:r:8:GLN:NE2	3:T:302:GLU:CG	2.53	0.65
3:B:48:TRP:N	3:B:48:TRP:CD1	2.64	0.65
3:B:92:TRP:CH2	3:B:160:GLN:HA	2.32	0.65
3:D:427:GLU:OE1	3:F:571:ARG:NH2	2.27	0.65
3:H:49:ARG:HH12	3:H:115:THR:CG2	2.10	0.65
3:J:195:PRO:HD3	3:N:566:TRP:CZ2	2.32	0.65
3:L:92:TRP:CH2	3:L:160:GLN:HA	2.32	0.65
3:L:311:ASN:ND2	3:L:311:ASN:H	1.94	0.65
3:N:49:ARG:HH12	3:N:115:THR:CG2	2.10	0.65
3:P:510:VAL:HG12	3:P:510:VAL:O	1.97	0.65
2:Q:28:SER:CA	3:R:216:PRO:HD2	2.27	0.65
5:W:833:ASP:OD2	5:W:842:ILE:HD12	1.97	0.65
6:X:939:LEU:HG	6:X:945:VAL:HG23	1.79	0.65
6:Y:453:LEU:HD13	6:Y:952:PRO:HB3	1.79	0.65
1:l:8:GLN:NE2	3:N:302:GLU:CG	2.56	0.65
1:b:61:ASN:ND2	1:b:61:ASN:O	2.30	0.65
1:j:73:ARG:O	1:j:75:PRO:HD3	1.97	0.65
1:r:59:HIS:HA	3:R:398:SER:HG	1.59	0.65
3:B:309:GLU:OE1	3:B:335:PRO:CB	2.44	0.65
3:D:238:VAL:HG22	3:F:142:ILE:HD12	1.79	0.65
3:D:566:TRP:CZ2	3:F:195:PRO:HD3	2.32	0.65
3:H:92:TRP:CH2	3:H:160:GLN:HA	2.32	0.65
3:H:141:GLN:NE2	3:H:141:GLN:HA	2.11	0.65
3:J:341:MET:HE2	3:J:341:MET:CA	2.18	0.65
3:L:49:ARG:HH12	3:L:115:THR:CG2	2.10	0.65
3:L:294:TYR:HH	3:L:502:LEU:HA	1.60	0.65
3:N:449:THR:HG22	3:N:450:SER:H	1.60	0.65
3:P:579:ASN:ND2	3:P:582:LEU:HG	2.06	0.65
3:R:63:ILE:HG21	3:R:70:GLN:OE1	1.96	0.65
6:X:317:PHE:CD2	6:X:370:ARG:HB3	2.30	0.65
6:X:373:ARG:HG3	6:X:385:VAL:HG12	1.79	0.65
6:Y:468:ALA:C	6:Y:471:PRO:HD2	2.21	0.65
1:b:15:ARG:CZ	3:B:406:ASN:ND2	2.60	0.64
1:j:71:CYS:SG	1:j:73:ARG:HG3	2.34	0.64
2:A:28:SER:CA	3:B:216:PRO:HD2	2.27	0.64
3:B:141:GLN:HA	3:B:141:GLN:NE2	2.11	0.64
3:B:216:PRO:HB3	3:B:234:PRO:HG3	1.78	0.64

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
7:C:101:MYR:C2	3:D:191:LYS:HE2	2.28	0.64
3:D:277:ASP:O	3:D:278:THR:CG2	2.37	0.64
3:F:309:GLU:OE1	3:F:335:PRO:CB	2.44	0.64
3:F:341:MET:HE2	3:F:341:MET:CA	2.18	0.64
3:H:71:TYR:CE1	3:H:138:SER:HB3	2.32	0.64
3:J:48:TRP:N	3:J:48:TRP:CD1	2.64	0.64
3:J:181:CYS:HB2	3:J:638:PHE:HB3	1.78	0.64
3:J:609:ARG:NH1	2:K:27:THR:CG2	2.56	0.64
3:L:548:LYS:HG3	3:N:645:TRP:CH2	2.32	0.64
3:N:48:TRP:N	3:N:48:TRP:CD1	2.64	0.64
3:N:341:MET:HE2	3:N:341:MET:CA	2.18	0.64
3:P:358:LEU:CG	3:P:417:PHE:CE1	2.79	0.64
3:R:341:MET:HB2	3:R:431:ILE:CD1	2.26	0.64
3:R:505:ASP:O	3:R:506:ALA:C	2.38	0.64
7:S:101:MYR:H31	3:T:191:LYS:CG	2.26	0.64
3:T:71:TYR:CE1	3:T:138:SER:HB3	2.31	0.64
6:X:660:SER:C	6:X:673:ASN:HD21	2.05	0.64
6:X:678:PRO:HD3	6:X:771:PHE:CE2	2.32	0.64
1:l:15:ARG:CZ	3:L:402:ALA:HB1	2.13	0.64
1:n:59:HIS:CA	3:N:398:SER:HB2	2.27	0.64
1:r:25:TYR:CD2	1:r:62:VAL:HG21	2.33	0.64
1:r:45:ARG:CD	3:R:401:GLY:H	2.10	0.64
3:D:57:THR:O	3:D:60:SER:CB	2.45	0.64
3:D:71:TYR:CE1	3:D:138:SER:HB3	2.31	0.64
3:D:174:ILE:HG22	3:D:634:LYS:HD2	1.77	0.64
3:D:645:TRP:CH2	3:H:548:LYS:HG3	2.32	0.64
2:G:28:SER:CA	3:H:216:PRO:HD2	2.27	0.64
3:H:216:PRO:HB3	3:H:234:PRO:HG3	1.78	0.64
3:H:309:GLU:OE1	3:H:335:PRO:CB	2.44	0.64
3:H:311:ASN:ND2	3:H:311:ASN:H	1.94	0.64
3:J:71:TYR:CE1	3:J:138:SER:HB3	2.31	0.64
3:J:579:ASN:ND2	3:J:582:LEU:HG	2.06	0.64
3:L:71:TYR:CE1	3:L:138:SER:HB3	2.31	0.64
3:L:141:GLN:NE2	3:L:141:GLN:HA	2.11	0.64
2:M:26:MET:HG3	2:M:26:MET:O	1.97	0.64
3:R:216:PRO:HB3	3:R:234:PRO:HG3	1.78	0.64
3:R:344:MET:CE	3:R:502:LEU:HD11	2.26	0.64
3:R:609:ARG:HD2	2:S:29:THR:HG21	1.78	0.64
4:U:46:ILE:HD12	6:X:1197:TYR:CE2	2.31	0.64
5:W:219:ASN:ND2	6:X:674:ARG:NH1	2.45	0.64
5:W:450:THR:HB	5:W:672:THR:OG1	1.97	0.64

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
6:Y:988:ALA:CA	6:Y:1133:VAL:HG12	2.25	0.64
6:Y:1060:ARG:HH21	6:Y:1101:GLU:HB3	1.60	0.64
1:h:47:THR:CG2	1:h:60:ALA:HA	2.27	0.64
1:h:58:PRO:HB2	3:H:423:GLN:CG	2.27	0.64
1:r:59:HIS:ND1	3:R:424:ALA:HA	2.12	0.64
2:C:28:SER:CA	3:D:216:PRO:HD2	2.27	0.64
3:D:276:ALA:HA	3:F:630:LEU:HD21	1.78	0.64
3:F:63:ILE:HG21	3:F:70:GLN:OE1	1.96	0.64
3:L:358:LEU:CG	3:L:417:PHE:CE1	2.79	0.64
3:L:593:GLN:HB2	3:L:596:MET:HE2	1.78	0.64
3:N:380:VAL:HG13	3:N:384:GLN:HG2	1.78	0.64
3:N:593:GLN:HB2	3:N:596:MET:HE2	1.78	0.64
2:O:26:MET:HG3	2:O:26:MET:O	1.97	0.64
2:O:35:ASP:OD1	2:O:40:VAL:HB	1.96	0.64
3:P:147:ARG:NH1	3:P:147:ARG:CB	2.61	0.64
3:T:181:CYS:HB2	3:T:638:PHE:HB3	1.78	0.64
3:T:309:GLU:OE1	3:T:335:PRO:CB	2.44	0.64
3:T:505:ASP:O	3:T:506:ALA:C	2.38	0.64
3:T:536:THR:HB	3:T:570:LYS:HA	1.80	0.64
4:V:195:GLN:NE2	6:Y:377:ILE:HB	2.12	0.64
5:W:1163:LEU:HG	5:W:1190:ARG:NH2	2.12	0.64
6:X:787:ARG:HH21	6:X:787:ARG:HG2	1.54	0.64
6:X:1031:LEU:HD12	6:X:1045:TRP:CZ3	2.32	0.64
6:X:1167:LEU:HB3	6:X:1187:ARG:NH1	2.12	0.64
6:Y:354:THR:O	6:Y:355:LYS:C	2.40	0.64
6:Y:1030:GLN:HB2	6:Y:1039:ARG:O	1.96	0.64
1:t:24:LEU:HG	1:t:38:ALA:HB3	1.78	0.64
3:B:71:TYR:CE1	3:B:138:SER:HB3	2.31	0.64
3:F:548:LYS:HG3	3:H:645:TRP:CH2	2.32	0.64
3:H:147:ARG:NH1	3:H:147:ARG:CB	2.61	0.64
3:H:333:PRO:HG3	3:H:454:PHE:HA	1.77	0.64
3:J:93:GLN:N	3:J:94:PRO:HD2	2.13	0.64
3:J:297:ALA:HB2	3:J:460:MET:CB	2.26	0.64
3:J:311:ASN:ND2	3:J:311:ASN:H	1.94	0.64
3:J:645:TRP:CH2	3:N:548:LYS:HG3	2.32	0.64
3:P:141:GLN:NE2	3:P:141:GLN:HA	2.11	0.64
3:R:548:LYS:HG3	3:T:645:TRP:CH2	2.32	0.64
5:W:246:ASP:CB	5:W:256:SER:HA	2.26	0.64
5:W:641:THR:HG23	5:W:660:GLY:HA3	1.78	0.64
6:X:203:SER:OG	6:X:247:THR:HA	1.97	0.64
6:X:925:LEU:HD21	6:Y:496:ALA:CA	2.24	0.64

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
6:X:1014:VAL:HG13	6:X:1031:LEU:HD11	1.80	0.64
6:X:1106:LYS:HB3	6:X:1113:PRO:CG	2.27	0.64
1:n:58:PRO:CG	3:N:423:GLN:NE2	2.50	0.64
3:B:67:ARG:CZ	3:B:67:ARG:HB2	2.27	0.64
3:B:341:MET:HB2	3:B:431:ILE:CD1	2.26	0.64
3:B:358:LEU:CG	3:B:417:PHE:CE1	2.79	0.64
3:D:147:ARG:NH1	3:D:147:ARG:CB	2.61	0.64
3:D:195:PRO:HD3	3:H:566:TRP:CZ2	2.32	0.64
3:D:363:ASN:ND2	3:D:366:LEU:HB2	2.13	0.64
3:F:49:ARG:HH12	3:F:115:THR:CG2	2.10	0.64
3:F:71:TYR:CE1	3:F:138:SER:HB3	2.31	0.64
3:F:141:GLN:HA	3:F:141:GLN:NE2	2.11	0.64
3:F:363:ASN:ND2	3:F:366:LEU:HB2	2.13	0.64
3:H:341:MET:HE2	3:H:341:MET:CA	2.18	0.64
3:H:593:GLN:HB2	3:H:596:MET:HE2	1.78	0.64
3:J:566:TRP:CZ2	3:L:195:PRO:HD3	2.32	0.64
7:M:101:MYR:H31	3:N:191:LYS:CG	2.26	0.64
3:N:280:LEU:HD12	3:N:486:PRO:HG2	1.78	0.64
3:N:358:LEU:CG	3:N:417:PHE:CE1	2.79	0.64
3:N:510:VAL:HG12	3:N:510:VAL:O	1.97	0.64
3:P:93:GLN:N	3:P:94:PRO:HD2	2.13	0.64
3:P:121:ALA:HB2	3:T:87:PHE:HD1	1.61	0.64
3:P:195:PRO:HD3	3:T:566:TRP:CZ2	2.32	0.64
3:P:469:VAL:CG2	3:R:575:MET:CE	2.43	0.64
3:P:520:VAL:CG1	3:P:608:LEU:HD11	2.28	0.64
3:R:181:CYS:HB2	3:R:638:PHE:HB3	1.78	0.64
3:R:276:ALA:HA	3:T:630:LEU:HD21	1.78	0.64
3:T:147:ARG:NH1	3:T:147:ARG:CB	2.61	0.64
4:U:379:ILE:HD12	4:U:388:PRO:HD3	1.79	0.64
4:V:184:HIS:CE1	6:Y:436:LEU:HD22	2.32	0.64
4:V:195:GLN:HE22	6:Y:377:ILE:N	1.94	0.64
5:W:709:GLY:HA3	5:W:763:ARG:HH21	1.62	0.64
6:X:493:LEU:CD1	6:X:497:ILE:HD12	2.27	0.64
6:X:940:GLN:NE2	6:X:944:ASN:HB3	2.12	0.64
6:X:1015:GLY:HA2	6:X:1034:ASP:HA	1.77	0.64
6:X:1150:ASN:HD22	6:X:1153:SER:HB2	1.61	0.64
6:Y:487:VAL:HG13	6:Y:826:GLN:O	1.97	0.64
6:Y:660:SER:HB3	6:Y:663:GLN:HG2	1.79	0.64
6:Y:848:ASN:N	6:Y:848:ASN:ND2	2.44	0.64
1:b:71:CYS:SG	1:b:73:ARG:CG	2.85	0.64
1:j:59:HIS:CA	3:J:398:SER:CB	2.76	0.64

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:p:45:ARG:N	3:P:399:ALA:O	2.29	0.64
3:D:358:LEU:CD1	3:D:417:PHE:CE1	2.62	0.64
3:F:566:TRP:CZ2	3:H:195:PRO:HD3	2.32	0.64
3:J:141:GLN:HA	3:J:141:GLN:NE2	2.11	0.64
3:J:147:ARG:NH1	3:J:147:ARG:CB	2.61	0.64
3:J:309:GLU:OE1	3:J:335:PRO:CB	2.44	0.64
3:J:548:LYS:HG3	3:L:645:TRP:CH2	2.32	0.64
3:L:147:ARG:NH1	3:L:147:ARG:CB	2.61	0.64
3:L:308:ALA:HB1	3:L:383:GLN:NE2	2.13	0.64
3:N:363:ASN:ND2	3:N:366:LEU:HB2	2.13	0.64
2:O:29:THR:HG21	3:T:609:ARG:HD2	1.78	0.64
3:P:92:TRP:CH2	3:P:160:GLN:HA	2.32	0.64
3:R:93:GLN:N	3:R:94:PRO:HD2	2.13	0.64
3:R:238:VAL:CG2	3:T:78:MET:HE1	2.07	0.64
3:R:297:ALA:HB2	3:R:460:MET:CB	2.26	0.64
5:W:99:LEU:C	5:W:99:LEU:HD23	2.23	0.64
5:W:303:LEU:CD2	5:W:341:MET:HE3	2.27	0.64
5:W:775:ARG:HD2	5:W:868:ASP:HB2	1.79	0.64
5:W:843:LEU:CD2	5:W:854:MET:HE1	2.27	0.64
6:X:201:CYS:O	6:X:201:CYS:SG	2.56	0.64
6:X:464:SER:CB	6:Y:500:ILE:HG21	2.15	0.64
6:X:789:ASP:HB3	6:X:933:ASP:OD2	1.97	0.64
1:p:45:ARG:CD	3:P:401:GLY:H	2.09	0.64
7:A:101:MYR:C2	3:B:191:LYS:HE2	2.27	0.64
3:B:363:ASN:ND2	3:B:366:LEU:HB2	2.13	0.64
3:D:58:ILE:HD11	3:D:127:LEU:HD23	1.79	0.64
3:D:308:ALA:HB1	3:D:383:GLN:NE2	2.13	0.64
3:D:609:ARG:HD2	2:E:29:THR:HG21	1.78	0.64
2:E:26:MET:HG3	2:E:26:MET:O	1.97	0.64
3:F:449:THR:HG22	3:F:450:SER:H	1.60	0.64
3:H:341:MET:HB2	3:H:431:ILE:CD1	2.26	0.64
3:L:276:ALA:HA	3:N:630:LEU:HD21	1.78	0.64
3:L:280:LEU:HD12	3:L:486:PRO:HG2	1.78	0.64
3:L:520:VAL:CG1	3:L:608:LEU:HD11	2.28	0.64
3:P:48:TRP:N	3:P:48:TRP:CD1	2.64	0.64
3:P:297:ALA:HB2	3:P:460:MET:CB	2.26	0.64
3:R:280:LEU:HD12	3:R:486:PRO:HG2	1.78	0.64
3:T:363:ASN:ND2	3:T:366:LEU:HB2	2.13	0.64
4:V:175:MET:SD	6:Y:400:GLU:O	2.55	0.64
5:W:434:VAL:HG22	5:W:1008:LEU:HD11	1.78	0.64
5:W:598:ALA:O	5:W:602:MET:HG3	1.97	0.64

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
5:W:672:THR:HG22	5:W:1081:VAL:HG13	1.80	0.64
5:W:672:THR:CG2	5:W:1081:VAL:HA	2.28	0.64
6:X:541:VAL:HG21	6:X:811:LEU:HG	1.80	0.64
6:X:865:VAL:CG1	6:X:958:HIS:HB2	2.28	0.64
1:p:43:CYS:HB2	1:p:45:ARG:H	1.63	0.64
1:p:45:ARG:CG	3:P:399:ALA:O	2.46	0.64
1:t:43:CYS:HB2	1:t:45:ARG:H	1.63	0.64
3:B:93:GLN:N	3:B:94:PRO:HD2	2.13	0.64
3:B:147:ARG:NH1	3:B:147:ARG:CB	2.61	0.64
3:B:554:GLN:O	3:B:556:SER:N	2.31	0.64
3:D:66:ASP:OD1	3:D:67:ARG:N	2.30	0.64
3:D:554:GLN:O	3:D:556:SER:N	2.31	0.64
3:F:330:GLN:HE22	3:F:332:GLY:HA2	1.63	0.64
3:J:216:PRO:HB3	3:J:234:PRO:HG3	1.78	0.64
3:J:412:ARG:HA	3:L:455:SER:O	1.98	0.64
3:J:510:VAL:HG12	3:J:510:VAL:O	1.97	0.64
3:J:520:VAL:CG1	3:J:608:LEU:HD11	2.28	0.64
3:L:554:GLN:O	3:L:556:SER:N	2.31	0.64
3:N:92:TRP:CH2	3:N:160:GLN:HA	2.32	0.64
3:P:341:MET:HB2	3:P:431:ILE:CD1	2.26	0.64
3:R:141:GLN:HA	3:R:141:GLN:NE2	2.11	0.64
3:T:93:GLN:N	3:T:94:PRO:HD2	2.13	0.64
3:T:311:ASN:ND2	3:T:311:ASN:H	1.94	0.64
4:U:89:GLY:C	4:U:103:VAL:HG23	2.23	0.64
4:U:166:MET:HE1	4:U:174:LEU:HD12	1.79	0.64
4:U:305:LEU:HG	4:U:306:PRO:HD2	1.80	0.64
4:V:68:VAL:O	4:V:72:LEU:HB2	1.97	0.64
4:V:208:ILE:HG23	4:V:290:VAL:HG22	1.80	0.64
5:W:603:LEU:O	5:W:603:LEU:HD23	1.98	0.64
6:X:392:ASP:HB2	6:X:1188:ILE:CG2	2.27	0.64
6:Y:996:MET:CB	6:Y:1143:ASN:HB3	2.28	0.64
1:b:23:THR:N	3:B:323:SER:OG	2.31	0.64
2:A:26:MET:HG3	2:A:26:MET:O	1.97	0.64
3:B:490:MET:HE3	3:B:597:LEU:HD13	1.80	0.64
3:B:520:VAL:CG1	3:B:608:LEU:HD11	2.28	0.64
3:D:309:GLU:OE1	3:D:335:PRO:CB	2.44	0.64
3:F:139:THR:O	3:F:142:ILE:HG23	1.98	0.64
3:F:308:ALA:HB1	3:F:383:GLN:NE2	2.13	0.64
3:F:358:LEU:CD1	3:F:417:PHE:CE1	2.62	0.64
3:L:363:ASN:ND2	3:L:366:LEU:HB2	2.13	0.64
3:N:93:GLN:N	3:N:94:PRO:HD2	2.13	0.64

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:N:308:ALA:HB1	3:N:383:GLN:NE2	2.13	0.64
3:N:536:THR:HB	3:N:570:LYS:HA	1.80	0.64
3:P:67:ARG:CZ	3:P:67:ARG:HB2	2.27	0.64
3:P:363:ASN:ND2	3:P:366:LEU:HB2	2.13	0.64
3:P:455:SER:O	3:T:412:ARG:HA	1.98	0.64
3:R:48:TRP:N	3:R:48:TRP:CD1	2.64	0.64
3:R:92:TRP:CH2	3:R:160:GLN:HA	2.32	0.64
3:R:167:MET:HG2	3:R:215:TYR:CZ	2.33	0.64
4:U:44:LEU:HA	4:U:51:LEU:HD11	1.80	0.64
4:U:185:LEU:HD13	4:U:211:LEU:HD11	1.80	0.64
5:W:789:ARG:O	5:W:789:ARG:HG3	1.97	0.64
6:X:427:LEU:O	6:X:428:LEU:CB	2.45	0.64
6:X:691:PRO:HA	6:X:742:MET:HE3	1.78	0.64
3:D:49:ARG:NH2	3:D:115:THR:HG21	2.14	0.64
3:D:93:GLN:N	3:D:94:PRO:HD2	2.13	0.64
3:D:341:MET:HE2	3:D:341:MET:CA	2.18	0.64
2:G:26:MET:HG3	2:G:26:MET:O	1.97	0.64
3:H:67:ARG:CZ	3:H:67:ARG:HB2	2.27	0.64
3:J:330:GLN:HE22	3:J:332:GLY:HA2	1.63	0.64
3:L:167:MET:HG2	3:L:215:TYR:CZ	2.33	0.64
3:N:341:MET:HB2	3:N:431:ILE:CD1	2.26	0.64
3:N:358:LEU:CD1	3:N:417:PHE:CE1	2.62	0.64
3:P:167:MET:HG2	3:P:215:TYR:CZ	2.33	0.64
3:P:490:MET:HE3	3:P:597:LEU:HD13	1.80	0.64
3:R:490:MET:HE3	3:R:597:LEU:HD13	1.80	0.64
7:S:101:MYR:C2	3:T:191:LYS:HE2	2.28	0.64
3:T:341:MET:CB	3:T:431:ILE:HD11	2.28	0.64
4:U:178:THR:CB	6:X:376:LEU:HD23	2.27	0.64
5:W:906:ALA:O	5:W:908:THR:HG23	1.97	0.64
6:X:464:SER:CB	6:Y:500:ILE:HG22	2.26	0.64
6:X:597:GLU:HG3	6:X:636:ALA:CB	2.28	0.64
6:Y:848:ASN:ND2	6:Y:848:ASN:H	1.95	0.64
1:f:59:HIS:HB3	3:F:398:SER:CB	2.07	0.63
1:r:23:THR:OG1	3:R:323:SER:HB2	1.93	0.63
1:r:43:CYS:HB2	1:r:45:ARG:H	1.63	0.63
3:B:510:VAL:HG12	3:B:510:VAL:O	1.97	0.63
2:C:37:LYS:NZ	3:F:264:SER:OG	2.23	0.63
3:H:491:ALA:CB	3:H:592:THR:HB	2.28	0.63
3:L:93:GLN:N	3:L:94:PRO:HD2	2.13	0.63
3:N:147:ARG:NH1	3:N:147:ARG:CB	2.61	0.63
2:O:28:SER:CA	3:P:216:PRO:HD2	2.27	0.63

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:T:308:ALA:HB1	3:T:383:GLN:NE2	2.13	0.63
3:T:490:MET:HE3	3:T:597:LEU:HD13	1.80	0.63
3:T:579:ASN:ND2	3:T:582:LEU:HG	2.06	0.63
4:U:41:PHE:HE1	6:X:404:MET:HE1	1.61	0.63
5:W:733:ALA:HB2	5:W:779:MET:CE	2.28	0.63
5:W:934:ASN:H	5:W:934:ASN:ND2	1.95	0.63
6:X:242:ARG:NH2	6:X:1085:LEU:CD1	2.56	0.63
6:X:793:ARG:NH1	6:Y:693:TYR:HE2	1.95	0.63
6:Y:483:VAL:HG13	6:Y:840:ALA:HB1	1.78	0.63
1:j:61:ASN:C	1:j:63:LYS:H	2.06	0.63
1:n:43:CYS:HB2	1:n:45:ARG:H	1.63	0.63
3:B:167:MET:HG2	3:B:215:TYR:CZ	2.33	0.63
3:B:292:ARG:HH11	3:B:292:ARG:HG2	1.64	0.63
3:D:311:ASN:ND2	3:D:311:ASN:H	1.94	0.63
3:F:93:GLN:N	3:F:94:PRO:HD2	2.13	0.63
3:F:147:ARG:NH1	3:F:147:ARG:CB	2.61	0.63
3:H:308:ALA:HB1	3:H:383:GLN:NE2	2.13	0.63
3:J:67:ARG:CZ	3:J:67:ARG:HB2	2.27	0.63
3:J:308:ALA:HB1	3:J:383:GLN:NE2	2.13	0.63
3:J:380:VAL:HG13	3:J:384:GLN:HG2	1.78	0.63
3:N:67:ARG:CZ	3:N:67:ARG:HB2	2.27	0.63
3:N:274:ILE:HD11	3:N:514:LEU:HD21	1.73	0.63
3:P:173:ASP:HB2	3:T:606:THR:HG21	1.80	0.63
3:P:491:ALA:CB	3:P:592:THR:HB	2.28	0.63
3:P:554:GLN:O	3:P:556:SER:N	2.31	0.63
3:P:575:MET:HE2	3:T:348:VAL:CG2	2.29	0.63
3:R:341:MET:CB	3:R:431:ILE:HD11	2.28	0.63
3:R:491:ALA:CB	3:R:592:THR:HB	2.28	0.63
3:T:92:TRP:CH2	3:T:160:GLN:HA	2.32	0.63
5:W:242:PRO:HG3	5:W:261:LEU:HD12	1.79	0.63
5:W:651:ASN:HD21	5:W:685:ASN:CG	2.06	0.63
6:X:380:GLU:CD	6:X:438:PRO:HG2	2.24	0.63
6:X:391:CYS:SG	6:X:1194:LEU:HD13	2.38	0.63
6:X:697:MET:CE	6:X:822:MET:HG3	2.27	0.63
6:Y:390:LEU:HD12	6:Y:390:LEU:O	1.98	0.63
1:d:59:HIS:CB	3:D:398:SER:HB2	2.28	0.63
1:h:59:HIS:CA	3:H:398:SER:OG	2.45	0.63
1:j:43:CYS:HB2	1:j:45:ARG:H	1.63	0.63
1:t:60:ALA:H	3:T:398:SER:HA	1.63	0.63
3:F:67:ARG:CZ	3:F:67:ARG:HB2	2.27	0.63
3:F:341:MET:CB	3:F:431:ILE:HD11	2.28	0.63

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:H:311:ASN:H	3:H:311:ASN:HD22	1.47	0.63
3:H:341:MET:CB	3:H:431:ILE:HD11	2.28	0.63
3:J:154:ARG:HB3	3:J:154:ARG:HH11	1.63	0.63
3:J:173:ASP:HB2	3:N:606:THR:HG21	1.80	0.63
3:J:436:TYR:OH	3:T:435:PRO:HG2	1.98	0.63
3:J:505:ASP:O	3:J:506:ALA:C	2.38	0.63
3:L:165:THR:O	3:L:169:MET:HB2	1.99	0.63
3:N:330:GLN:HE22	3:N:332:GLY:HA2	1.63	0.63
3:R:67:ARG:CZ	3:R:67:ARG:HB2	2.27	0.63
3:R:520:VAL:CG1	3:R:608:LEU:HD11	2.28	0.63
4:U:196:LEU:HD21	6:X:448:SER:HB2	1.81	0.63
5:W:81:TYR:HA	5:W:180:LEU:HD23	1.80	0.63
6:X:283:ILE:HD11	6:X:287:GLN:C	2.23	0.63
6:Y:73:ASN:OD1	6:Y:74:ASN:N	2.31	0.63
6:Y:709:ILE:HG12	6:Y:733:MET:HE3	1.80	0.63
6:Y:1021:VAL:O	6:Y:1021:VAL:HG23	1.96	0.63
1:h:43:CYS:HB2	1:h:45:ARG:H	1.63	0.63
1:j:15:ARG:NE	3:J:406:ASN:CG	2.56	0.63
1:n:15:ARG:NH2	3:N:402:ALA:HB2	2.13	0.63
1:p:44:GLY:O	3:P:399:ALA:O	2.17	0.63
3:D:292:ARG:HH11	3:D:292:ARG:HG2	1.64	0.63
3:D:298:PHE:CE2	3:H:355:GLY:HA3	2.33	0.63
3:D:322:TYR:O	3:D:323:SER:OG	2.14	0.63
3:D:455:SER:O	3:H:412:ARG:HA	1.98	0.63
3:F:165:THR:O	3:F:169:MET:HB2	1.99	0.63
3:F:238:VAL:HG11	3:H:82:PHE:HZ	1.64	0.63
3:H:48:TRP:N	3:H:48:TRP:CD1	2.64	0.63
3:H:490:MET:HE3	3:H:597:LEU:HD13	1.80	0.63
3:J:341:MET:CB	3:J:431:ILE:HD11	2.28	0.63
3:J:427:GLU:OE1	3:L:571:ARG:NH2	2.27	0.63
3:J:554:GLN:O	3:J:556:SER:N	2.31	0.63
2:K:26:MET:HG3	2:K:26:MET:O	1.97	0.63
3:L:348:VAL:CG2	3:N:575:MET:HE2	2.29	0.63
3:P:176:ALA:O	3:P:631:THR:HG21	1.99	0.63
3:P:311:ASN:H	3:P:311:ASN:HD22	1.46	0.63
3:P:645:TRP:CH2	3:T:548:LYS:HG3	2.32	0.63
3:R:147:ARG:NH1	3:R:147:ARG:CB	2.61	0.63
3:R:176:ALA:O	3:R:631:THR:HG21	1.99	0.63
3:R:363:ASN:ND2	3:R:366:LEU:HB2	2.13	0.63
3:R:606:THR:HG21	3:T:173:ASP:HB2	1.80	0.63
2:S:26:MET:HG3	2:S:26:MET:O	1.97	0.63

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:T:165:THR:O	3:T:169:MET:HB2	1.99	0.63
4:V:156:ILE:HG22	4:V:268:ARG:HB2	1.81	0.63
4:V:166:MET:HE1	4:V:174:LEU:HD12	1.80	0.63
5:W:853:THR:CG2	5:W:873:ALA:HB3	2.28	0.63
5:W:1064:GLN:CB	5:W:1069:GLU:HA	2.28	0.63
6:Y:663:GLN:HE21	6:Y:663:GLN:HA	1.61	0.63
6:Y:1104:LEU:HD22	6:Y:1104:LEU:H	1.64	0.63
1:b:74:GLN:C	1:b:75:PRO:CA	2.69	0.63
3:B:165:THR:O	3:B:169:MET:HB2	1.99	0.63
3:B:536:THR:HB	3:B:570:LYS:HA	1.80	0.63
3:D:141:GLN:NE2	3:D:141:GLN:HA	2.11	0.63
3:D:173:ASP:HB2	3:H:606:THR:HG21	1.80	0.63
3:F:292:ARG:HH11	3:F:292:ARG:HG2	1.64	0.63
3:F:311:ASN:H	3:F:311:ASN:HD22	1.47	0.63
3:F:520:VAL:CG1	3:F:608:LEU:HD11	2.28	0.63
3:H:520:VAL:CG1	3:H:608:LEU:HD11	2.28	0.63
3:H:536:THR:HB	3:H:570:LYS:HA	1.80	0.63
2:I:28:SER:CA	3:J:216:PRO:HD2	2.27	0.63
3:J:491:ALA:CB	3:J:592:THR:HB	2.28	0.63
3:P:348:VAL:CG2	3:R:575:MET:HE2	2.29	0.63
3:R:554:GLN:O	3:R:556:SER:N	2.31	0.63
3:T:49:ARG:HH12	3:T:115:THR:CG2	2.10	0.63
6:Y:545:LEU:HD13	6:Y:569:LEU:CD1	2.29	0.63
1:l:59:HIS:CA	3:L:398:SER:CB	2.77	0.63
3:B:311:ASN:H	3:B:311:ASN:HD22	1.47	0.63
3:D:154:ARG:HB3	3:D:154:ARG:HH11	1.63	0.63
3:D:418:ASN:HD21	3:F:337:ARG:NH2	1.93	0.63
3:D:548:LYS:HG3	3:F:645:TRP:CH2	2.32	0.63
3:F:536:THR:HB	3:F:570:LYS:HA	1.80	0.63
3:H:358:LEU:CD1	3:H:417:PHE:CE1	2.61	0.63
3:J:348:VAL:CG2	3:L:575:MET:HE2	2.29	0.63
3:J:355:GLY:HA3	3:L:298:PHE:CE2	2.33	0.63
3:L:606:THR:HG21	3:N:173:ASP:HB2	1.80	0.63
3:N:341:MET:CB	3:N:431:ILE:HD11	2.28	0.63
3:P:238:VAL:HG11	3:R:82:PHE:HZ	1.64	0.63
3:P:412:ARG:HA	3:R:455:SER:O	1.98	0.63
3:P:536:THR:HB	3:P:570:LYS:HA	1.80	0.63
3:R:330:GLN:HE22	3:R:332:GLY:HA2	1.63	0.63
3:R:510:VAL:HG12	3:R:510:VAL:O	1.97	0.63
4:U:22:ASP:CB	4:U:383:ARG:HD3	2.28	0.63
4:U:59:PRO:HG2	4:U:386:ILE:HD11	1.78	0.63

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
4:U:184:HIS:O	4:U:187:GLN:HB3	1.99	0.63
4:V:178:THR:HG21	6:Y:408:SER:HA	1.79	0.63
5:W:12:LYS:HB3	5:W:386:GLY:O	1.98	0.63
5:W:624:THR:C	5:W:626:ALA:H	2.05	0.63
6:X:599:ILE:CG2	6:X:639:LEU:HD23	2.29	0.63
6:X:1091:GLN:NE2	6:X:1091:GLN:HA	2.14	0.63
6:X:1179:VAL:O	6:X:1179:VAL:HG13	1.98	0.63
1:l:43:CYS:HB2	1:l:45:ARG:H	1.63	0.63
1:b:61:ASN:O	1:b:63:LYS:N	2.32	0.63
1:f:53:LEU:HD13	1:f:74:GLN:HE21	1.62	0.63
1:j:15:ARG:NH2	3:J:402:ALA:HB3	2.13	0.63
1:j:59:HIS:HB2	3:J:397:VAL:C	2.24	0.63
1:p:42:THR:HG21	1:p:60:ALA:HB1	1.80	0.63
3:B:49:ARG:HH12	3:B:115:THR:CG2	2.10	0.63
3:D:348:VAL:CG2	3:F:575:MET:HE2	2.29	0.63
3:F:154:ARG:HB3	3:F:154:ARG:HH11	1.63	0.63
3:F:167:MET:HG2	3:F:215:TYR:CZ	2.33	0.63
3:H:167:MET:HG2	3:H:215:TYR:CZ	2.34	0.63
3:H:510:VAL:HG12	3:H:510:VAL:O	1.97	0.63
3:J:167:MET:HG2	3:J:215:TYR:CZ	2.33	0.63
3:J:199:LEU:O	3:J:207:PHE:CE2	2.52	0.63
3:L:238:VAL:HG11	3:N:82:PHE:HZ	1.64	0.63
3:L:341:MET:HE2	3:L:341:MET:CA	2.18	0.63
3:R:292:ARG:HH11	3:R:292:ARG:HG2	1.63	0.63
3:R:311:ASN:H	3:R:311:ASN:HD22	1.47	0.63
3:R:566:TRP:CZ2	3:T:195:PRO:HD3	2.32	0.63
3:T:520:VAL:CG1	3:T:608:LEU:HD11	2.28	0.63
5:W:57:THR:HG22	5:W:183:TYR:HD1	1.63	0.63
5:W:1038:ASN:ND2	5:W:1049:ALA:HB3	2.07	0.63
5:W:1096:PHE:CD1	5:W:1096:PHE:N	2.66	0.63
6:X:257:ARG:HB2	6:X:310:LYS:NZ	2.14	0.63
6:X:1091:GLN:HE22	6:X:1127:CYS:HB2	1.62	0.63
6:Y:560:LEU:HD21	6:Y:721:VAL:HG23	1.80	0.63
6:Y:993:TRP:NE1	6:Y:995:PRO:HG3	2.13	0.63
3:B:308:ALA:HB1	3:B:383:GLN:NE2	2.13	0.63
3:D:167:MET:HG2	3:D:215:TYR:CZ	2.33	0.63
3:F:98:ALA:HB2	3:F:206:GLU:CG	2.29	0.63
3:F:505:ASP:O	3:F:506:ALA:C	2.38	0.63
3:F:554:GLN:O	3:F:556:SER:N	2.31	0.63
3:H:330:GLN:HE22	3:H:332:GLY:HA2	1.63	0.63
3:H:554:GLN:O	3:H:556:SER:N	2.31	0.63

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:H:625:ARG:HG3	3:L:197:ASP:OD1	1.99	0.63
3:L:491:ALA:CB	3:L:592:THR:HB	2.28	0.63
3:L:510:VAL:HG12	3:L:510:VAL:O	1.97	0.63
3:L:635:GLN:HE21	3:L:635:GLN:HA	1.64	0.63
3:N:176:ALA:O	3:N:631:THR:HG21	1.99	0.63
3:N:520:VAL:CG1	3:N:608:LEU:HD11	2.28	0.63
3:N:554:GLN:O	3:N:556:SER:N	2.31	0.63
3:P:165:THR:O	3:P:169:MET:HB2	1.99	0.63
3:R:165:THR:O	3:R:169:MET:HB2	1.99	0.63
5:W:68:TRP:CZ2	5:W:70:PRO:HG3	2.34	0.63
5:W:454:PRO:HA	5:W:457:LEU:HD21	1.81	0.63
6:X:520:PRO:HD2	6:X:816:ALA:HB1	1.79	0.63
6:X:653:PRO:HD3	6:X:773:ARG:CG	2.28	0.63
6:X:875:GLN:HA	6:X:875:GLN:OE1	1.98	0.63
1:f:59:HIS:CB	3:F:398:SER:OG	2.47	0.63
1:n:58:PRO:HB2	3:N:423:GLN:HG2	1.80	0.63
1:n:59:HIS:CA	3:N:398:SER:CB	2.76	0.63
1:p:60:ALA:HB3	3:P:322:TYR:CZ	2.34	0.63
1:t:8:GLN:NE2	3:P:302:GLU:CG	2.56	0.63
3:D:165:THR:O	3:D:169:MET:HB2	1.99	0.63
3:D:311:ASN:H	3:D:311:ASN:HD22	1.46	0.63
3:D:520:VAL:CG1	3:D:608:LEU:HD11	2.28	0.63
3:F:635:GLN:HE21	3:F:635:GLN:HA	1.64	0.63
3:H:363:ASN:ND2	3:H:366:LEU:HB2	2.13	0.63
3:J:98:ALA:HB2	3:J:206:GLU:CG	2.29	0.63
3:L:292:ARG:HH11	3:L:292:ARG:HG2	1.63	0.63
3:N:154:ARG:HB3	3:N:154:ARG:HH11	1.63	0.63
3:P:548:LYS:HG3	3:R:645:TRP:CH2	2.32	0.63
3:R:308:ALA:HB1	3:R:383:GLN:NE2	2.13	0.63
2:S:27:THR:O	2:S:27:THR:CG2	2.38	0.63
3:T:292:ARG:HH11	3:T:292:ARG:HG2	1.64	0.63
4:U:158:HIS:CE1	4:U:160:GLU:HG3	2.32	0.63
4:U:223:LEU:HD11	4:U:411:PHE:CZ	2.34	0.63
4:V:124:ASN:O	4:V:126:PRO:HD3	1.99	0.63
5:W:834:LEU:HD11	5:W:921:ILE:HD11	1.79	0.63
5:W:879:TYR:HA	5:W:884:PHE:CD2	2.34	0.63
5:W:1057:PRO:HG3	5:W:1089:TYR:CZ	2.34	0.63
6:X:439:THR:O	6:X:1200:THR:HG23	1.99	0.63
6:X:467:ALA:O	6:X:471:PRO:HD3	1.99	0.63
6:X:528:PHE:HZ	6:X:747:ILE:HD11	1.64	0.63
6:X:813:VAL:HG11	6:X:818:ILE:HD11	1.81	0.63

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
6:Y:262:ARG:HH11	6:Y:262:ARG:HG3	1.62	0.63
6:Y:298:LEU:HD12	6:Y:885:LEU:HD12	1.81	0.63
7:C:101:MYR:H31	3:D:191:LYS:CG	2.30	0.62
3:F:490:MET:HE3	3:F:597:LEU:HD13	1.80	0.62
3:H:292:ARG:HH11	3:H:292:ARG:HG2	1.63	0.62
3:J:82:PHE:HZ	3:N:238:VAL:HG11	1.64	0.62
3:J:363:ASN:ND2	3:J:366:LEU:HB2	2.13	0.62
3:J:490:MET:HE3	3:J:597:LEU:HD13	1.80	0.62
3:L:311:ASN:H	3:L:311:ASN:HD22	1.47	0.62
3:P:98:ALA:HB2	3:P:206:GLU:CG	2.29	0.62
3:T:99:THR:OG1	3:T:101:THR:HG22	1.97	0.62
3:T:137:ILE:HD13	3:T:142:ILE:HG13	1.79	0.62
5:W:31:LEU:CD1	5:W:96:VAL:HG13	2.28	0.62
5:W:910:LEU:HD12	5:W:968:ALA:HB2	1.80	0.62
5:W:970:PRO:O	5:W:974:GLN:HB2	1.98	0.62
5:W:983:ALA:O	5:W:1034:PRO:HD3	1.98	0.62
5:W:1052:ILE:HG21	5:W:1112:LEU:HD22	1.81	0.62
6:Y:432:ILE:CG2	6:Y:1204:LEU:HB3	2.27	0.62
1:l:21:ARG:HH11	3:L:370:VAL:CG1	2.11	0.62
3:D:330:GLN:HE22	3:D:332:GLY:HA2	1.63	0.62
3:F:412:ARG:HA	3:H:455:SER:O	1.98	0.62
3:H:199:LEU:O	3:H:207:PHE:CE2	2.52	0.62
3:J:68:PHE:HD1	3:J:140:VAL:HG11	1.65	0.62
3:L:412:ARG:HA	3:N:455:SER:O	1.98	0.62
3:L:536:THR:HB	3:L:570:LYS:HA	1.80	0.62
3:N:490:MET:HE3	3:N:597:LEU:HD13	1.80	0.62
3:P:606:THR:HG21	3:R:173:ASP:HB2	1.80	0.62
3:R:98:ALA:HB2	3:R:206:GLU:CG	2.29	0.62
4:V:80:ALA:HB2	4:V:120:LEU:HD23	1.79	0.62
6:X:201:CYS:HB2	6:X:399:ARG:NH2	2.15	0.62
6:X:483:VAL:HG23	6:X:483:VAL:O	1.98	0.62
6:X:519:GLN:H	6:X:520:PRO:CD	2.12	0.62
6:X:1004:PHE:O	6:X:1004:PHE:CG	2.52	0.62
6:Y:169:ILE:HG21	6:Y:843:PRO:HA	1.80	0.62
6:Y:195:ILE:HD12	6:Y:257:ARG:NE	2.14	0.62
6:Y:412:ASP:CG	6:Y:414:THR:HG22	2.24	0.62
1:l:23:THR:HB	3:L:323:SER:CA	2.29	0.62
1:d:43:CYS:HB2	1:d:45:ARG:H	1.63	0.62
1:h:60:ALA:O	1:h:61:ASN:C	2.41	0.62
1:j:44:GLY:O	3:J:399:ALA:C	2.41	0.62
1:j:59:HIS:CA	3:J:398:SER:HA	2.30	0.62

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:B:176:ALA:O	3:B:631:THR:HG21	1.99	0.62
3:B:341:MET:CB	3:B:431:ILE:HD11	2.28	0.62
3:D:341:MET:HG3	3:D:437:ILE:HG12	1.82	0.62
3:D:412:ARG:HA	3:F:455:SER:O	1.98	0.62
3:D:575:MET:HE2	3:H:348:VAL:CG2	2.29	0.62
3:J:341:MET:HG3	3:J:437:ILE:HG12	1.82	0.62
3:J:358:LEU:CD1	3:J:417:PHE:CE1	2.62	0.62
3:J:575:MET:HE2	3:N:348:VAL:CG2	2.29	0.62
3:L:490:MET:HE3	3:L:597:LEU:HD13	1.80	0.62
3:R:68:PHE:HD1	3:R:140:VAL:HG11	1.64	0.62
3:T:341:MET:HE2	3:T:341:MET:CA	2.18	0.62
3:T:443:PHE:HB2	3:T:457:PRO:CB	2.24	0.62
4:V:175:MET:SD	6:Y:400:GLU:CG	2.83	0.62
5:W:814:GLY:HA3	5:W:991:LEU:HD13	1.80	0.62
6:X:567:LEU:HD11	6:X:705:ILE:HD13	1.81	0.62
6:X:637:PRO:HG2	6:X:638:LEU:H	1.64	0.62
6:X:712:VAL:HA	6:X:717:MET:HG2	1.81	0.62
6:X:1150:ASN:ND2	6:X:1153:SER:HB2	2.14	0.62
6:Y:601:ILE:CG2	6:Y:603:VAL:HG12	2.29	0.62
6:Y:769:VAL:HB	6:Y:780:ILE:CD1	2.30	0.62
6:Y:813:VAL:HG21	6:Y:818:ILE:HD11	1.81	0.62
1:j:45:ARG:CD	3:J:401:GLY:H	2.12	0.62
1:t:8:GLN:HE22	3:P:303:ASP:CA	2.11	0.62
2:C:26:MET:HG3	2:C:26:MET:O	1.97	0.62
3:F:294:TYR:HH	3:F:502:LEU:HA	1.62	0.62
3:H:165:THR:O	3:H:169:MET:HB2	1.99	0.62
3:H:277:ASP:O	3:H:278:THR:CG2	2.37	0.62
3:H:297:ALA:HB2	3:H:460:MET:CB	2.26	0.62
3:L:44:THR:HG22	3:L:45:GLY:N	2.14	0.62
3:L:176:ALA:O	3:L:631:THR:HG21	1.99	0.62
3:L:380:VAL:HG13	3:L:384:GLN:HG2	1.78	0.62
2:M:5:GLN:O	2:M:6:THR:HB	2.00	0.62
3:N:165:THR:O	3:N:169:MET:HB2	1.99	0.62
3:P:418:ASN:HD21	3:R:337:ARG:NH2	1.93	0.62
3:P:443:PHE:HB2	3:P:457:PRO:CB	2.24	0.62
3:R:536:THR:HB	3:R:570:LYS:HA	1.80	0.62
3:T:297:ALA:HB2	3:T:460:MET:CB	2.26	0.62
3:T:635:GLN:HE21	3:T:635:GLN:HA	1.64	0.62
5:W:47:LEU:HD11	5:W:327:ILE:CG2	2.29	0.62
5:W:710:ILE:HD11	5:W:798:THR:C	2.24	0.62
6:X:282:TYR:HD2	6:Y:830:ASN:ND2	1.94	0.62

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
6:X:350:LEU:HD23	6:X:357:ALA:HB1	1.81	0.62
6:X:454:GLN:HA	6:X:457:MET:HB3	1.82	0.62
6:X:488:LEU:HD11	6:X:821:ALA:HA	1.81	0.62
6:X:974:LEU:HD11	6:X:1168:ARG:HH22	1.63	0.62
6:Y:1023:ARG:HH21	6:Y:1030:GLN:HB3	1.64	0.62
1:b:59:HIS:HA	3:B:398:SER:HA	1.80	0.62
1:f:21:ARG:HH11	3:F:327:TYR:HH	1.44	0.62
3:B:98:ALA:HB2	3:B:206:GLU:CG	2.29	0.62
3:B:199:LEU:O	3:B:207:PHE:CE2	2.52	0.62
3:F:491:ALA:CB	3:F:592:THR:HB	2.28	0.62
2:G:5:GLN:O	2:G:6:THR:HB	2.00	0.62
3:H:341:MET:HG3	3:H:437:ILE:HG12	1.82	0.62
3:N:167:MET:HG2	3:N:215:TYR:CZ	2.34	0.62
3:N:292:ARG:HH11	3:N:292:ARG:HG2	1.64	0.62
3:P:55:VAL:HG11	5:W:348:GLN:HE22	1.62	0.62
3:P:355:GLY:HA3	3:R:298:PHE:CE2	2.34	0.62
7:S:101:MYR:C2	3:T:191:LYS:CE	2.78	0.62
3:T:98:ALA:HB2	3:T:206:GLU:CG	2.29	0.62
5:W:632:PHE:CD2	5:W:670:SER:HB3	2.35	0.62
1:l:23:THR:HG21	1:l:25:TYR:HE2	1.61	0.62
1:l:24:LEU:HD23	1:l:80:ARG:HD2	1.81	0.62
1:l:58:PRO:HB3	3:L:423:GLN:HE21	1.64	0.62
1:d:15:ARG:CZ	3:D:406:ASN:CG	2.71	0.62
3:D:199:LEU:O	3:D:207:PHE:CE2	2.52	0.62
3:D:356:THR:HG23	3:H:412:ARG:HH22	1.65	0.62
3:D:490:MET:HE3	3:D:597:LEU:HD13	1.80	0.62
3:D:491:ALA:CB	3:D:592:THR:HB	2.28	0.62
3:F:176:ALA:O	3:F:631:THR:HG21	1.99	0.62
3:F:199:LEU:O	3:F:207:PHE:CE2	2.52	0.62
3:F:330:GLN:NE2	3:F:332:GLY:CA	2.62	0.62
3:F:355:GLY:HA3	3:H:298:PHE:CE2	2.34	0.62
3:H:68:PHE:HD1	3:H:140:VAL:HG11	1.65	0.62
3:H:93:GLN:N	3:H:94:PRO:HD2	2.13	0.62
3:H:154:ARG:HB3	3:H:154:ARG:HH11	1.63	0.62
3:H:635:GLN:HE21	3:H:635:GLN:HA	1.64	0.62
2:I:9:ASN:HB2	3:J:209:ASN:ND2	2.15	0.62
3:J:176:ALA:O	3:J:631:THR:HG21	1.99	0.62
3:J:330:GLN:NE2	3:J:332:GLY:CA	2.62	0.62
3:L:199:LEU:O	3:L:207:PHE:CE2	2.52	0.62
3:L:273:LEU:HD23	3:L:273:LEU:C	2.25	0.62
3:P:199:LEU:O	3:P:207:PHE:CE2	2.52	0.62

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:P:292:ARG:HH11	3:P:292:ARG:HG2	1.64	0.62
3:R:44:THR:HG22	3:R:45:GLY:N	2.14	0.62
3:T:554:GLN:O	3:T:556:SER:N	2.31	0.62
5:W:79:LEU:HD23	5:W:79:LEU:H	1.64	0.62
5:W:577:LEU:HD23	5:W:577:LEU:C	2.24	0.62
5:W:874:TYR:HE2	5:W:876:GLN:HG3	1.63	0.62
6:X:350:LEU:HD12	6:X:350:LEU:N	2.14	0.62
6:X:590:ALA:HA	6:X:639:LEU:CD1	2.29	0.62
6:X:1156:THR:HG21	6:X:1161:ASN:CA	2.30	0.62
6:Y:401:THR:HG21	6:Y:960:ALA:HB2	1.82	0.62
6:Y:403:ILE:O	6:Y:403:ILE:HD13	2.00	0.62
7:A:101:MYR:H31	3:B:191:LYS:CG	2.30	0.62
3:B:154:ARG:HB3	3:B:154:ARG:HH11	1.63	0.62
2:E:5:GLN:O	2:E:6:THR:HB	1.99	0.62
3:F:348:VAL:CG2	3:H:575:MET:HE2	2.29	0.62
7:G:101:MYR:C2	3:H:191:LYS:CE	2.78	0.62
3:J:418:ASN:HD21	3:L:337:ARG:NH2	1.93	0.62
3:N:505:ASP:O	3:N:506:ALA:C	2.38	0.62
3:P:44:THR:HG22	3:P:45:GLY:N	2.14	0.62
3:P:154:ARG:HB3	3:P:154:ARG:HH11	1.63	0.62
3:R:330:GLN:NE2	3:R:332:GLY:CA	2.62	0.62
3:R:348:VAL:CG2	3:T:575:MET:HE2	2.29	0.62
3:R:635:GLN:NE2	3:R:635:GLN:HA	2.15	0.62
3:T:44:THR:HG22	3:T:45:GLY:N	2.14	0.62
3:T:273:LEU:HD23	3:T:273:LEU:C	2.25	0.62
3:T:330:GLN:NE2	3:T:332:GLY:CA	2.62	0.62
5:W:59:ILE:O	5:W:178:ILE:HD13	2.00	0.62
5:W:486:SER:O	5:W:490:ASP:HB2	2.00	0.62
6:X:192:GLU:O	6:X:192:GLU:HG3	1.99	0.62
6:X:898:ALA:HB3	6:X:929:PRO:HD2	1.81	0.62
6:X:978:ALA:HB3	6:X:1081:TYR:CE1	2.34	0.62
6:X:1095:ASP:O	6:X:1095:ASP:CG	2.42	0.62
6:Y:266:PRO:HA	6:Y:976:ASN:ND2	2.13	0.62
1:d:61:ASN:OD1	3:D:395:THR:HG21	1.99	0.62
1:r:23:THR:HG21	1:r:25:TYR:HH	1.58	0.62
2:C:9:ASN:HB2	3:D:209:ASN:ND2	2.15	0.62
3:D:82:PHE:HZ	3:H:238:VAL:HG11	1.64	0.62
3:D:635:GLN:HE21	3:D:635:GLN:HA	1.65	0.62
3:F:606:THR:HG21	3:H:173:ASP:HB2	1.80	0.62
3:H:98:ALA:HB2	3:H:206:GLU:CG	2.29	0.62
3:J:237:GLU:CG	3:L:142:ILE:HD13	2.25	0.62

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:J:606:THR:HG21	3:L:173:ASP:HB2	1.80	0.62
3:L:297:ALA:HB2	3:L:460:MET:CB	2.26	0.62
3:L:330:GLN:HE22	3:L:332:GLY:HA2	1.63	0.62
3:N:635:GLN:NE2	3:N:635:GLN:HA	2.15	0.62
3:R:49:ARG:HH12	3:R:115:THR:CG2	2.10	0.62
3:R:341:MET:HG3	3:R:437:ILE:HG12	1.82	0.62
3:R:412:ARG:HA	3:T:455:SER:O	1.98	0.62
3:T:199:LEU:O	3:T:207:PHE:CE2	2.52	0.62
3:T:311:ASN:H	3:T:311:ASN:HD22	1.47	0.62
4:U:224:ASN:ND2	4:U:226:SER:H	1.98	0.62
4:V:239:PRO:HA	4:V:243:PRO:HB3	1.81	0.62
5:W:131:ILE:O	5:W:131:ILE:HG12	1.99	0.62
5:W:739:VAL:HG12	5:W:742:SER:CB	2.30	0.62
6:X:453:LEU:HA	6:X:950:VAL:HG21	1.80	0.62
6:X:575:MET:HE1	6:X:747:ILE:HG12	1.81	0.62
6:Y:440:ILE:HG13	6:Y:1202:PRO:CG	2.29	0.62
1:b:17:ALA:CA	1:b:22:LEU:HD21	2.30	0.62
1:p:44:GLY:HA3	3:P:399:ALA:HB3	1.81	0.62
3:B:330:GLN:NE2	3:B:332:GLY:CA	2.62	0.62
3:B:380:VAL:HG13	3:B:384:GLN:HG2	1.78	0.62
3:D:336:THR:HG22	3:D:456:ASN:HD21	1.65	0.62
2:E:9:ASN:HB2	3:F:209:ASN:ND2	2.15	0.62
3:H:44:THR:HG22	3:H:45:GLY:N	2.14	0.62
3:H:330:GLN:NE2	3:H:332:GLY:CA	2.62	0.62
3:H:358:LEU:HD13	3:H:417:PHE:CZ	2.35	0.62
3:H:635:GLN:NE2	3:H:635:GLN:HA	2.15	0.62
3:J:58:ILE:O	3:J:59:ASP:HB2	1.99	0.62
3:J:455:SER:O	3:N:412:ARG:HA	1.98	0.62
3:L:98:ALA:HB2	3:L:206:GLU:CG	2.29	0.62
3:L:330:GLN:NE2	3:L:332:GLY:CA	2.62	0.62
3:N:330:GLN:NE2	3:N:332:GLY:CA	2.62	0.62
3:P:68:PHE:HD1	3:P:140:VAL:HG11	1.65	0.62
3:P:330:GLN:NE2	3:P:332:GLY:CA	2.62	0.62
3:P:330:GLN:HE22	3:P:332:GLY:HA2	1.63	0.62
3:P:341:MET:HG3	3:P:437:ILE:HG12	1.82	0.62
3:T:635:GLN:NE2	3:T:635:GLN:HA	2.15	0.62
4:U:93:ARG:HH11	4:U:93:ARG:CB	2.11	0.62
4:U:318:ASN:H	4:U:322:VAL:HG11	1.64	0.62
4:V:276:HIS:HB2	4:V:370:LEU:HD21	1.80	0.62
5:W:426:LEU:HD21	5:W:797:PHE:CD1	2.35	0.62
6:X:593:LEU:O	6:X:593:LEU:HG	1.99	0.62

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
6:Y:537:TYR:CE1	6:Y:766:PRO:HA	2.34	0.62
6:Y:586:PHE:HA	6:Y:642:PHE:CE2	2.34	0.62
1:h:45:ARG:CG	3:H:399:ALA:O	2.48	0.62
1:h:50:ALA:O	1:h:74:GLN:HG2	1.99	0.62
2:A:5:GLN:O	2:A:6:THR:HB	1.99	0.62
3:D:44:THR:HG22	3:D:45:GLY:N	2.14	0.62
3:D:330:GLN:NE2	3:D:332:GLY:CA	2.62	0.62
3:D:355:GLY:HA3	3:F:298:PHE:CE2	2.33	0.62
3:F:273:LEU:HD23	3:F:273:LEU:C	2.25	0.62
3:F:341:MET:HG3	3:F:437:ILE:HG12	1.81	0.62
3:F:549:LEU:HD12	3:F:550:SER:N	2.15	0.62
3:H:176:ALA:O	3:H:631:THR:HG21	1.99	0.62
3:J:44:THR:HG22	3:J:45:GLY:N	2.14	0.62
3:L:355:GLY:HA3	3:N:298:PHE:CE2	2.34	0.62
3:L:427:GLU:OE1	3:N:571:ARG:NH2	2.27	0.62
3:L:635:GLN:NE2	3:L:635:GLN:HA	2.15	0.62
7:M:101:MYR:C2	3:N:191:LYS:HE2	2.27	0.62
3:N:44:THR:HG22	3:N:45:GLY:N	2.14	0.62
3:N:57:THR:HG22	3:N:58:ILE:N	2.15	0.62
3:N:336:THR:HG22	3:N:456:ASN:HD21	1.65	0.62
3:P:341:MET:CB	3:P:431:ILE:HD11	2.28	0.62
3:T:167:MET:HG2	3:T:215:TYR:CZ	2.33	0.62
5:W:95:PHE:HB2	5:W:125:PHE:CE2	2.35	0.62
5:W:1134:VAL:CG2	5:W:1219:ILE:HD11	2.30	0.62
1:b:43:CYS:HB2	1:b:45:ARG:H	1.63	0.61
1:f:58:PRO:CB	3:F:423:GLN:CG	2.76	0.61
3:B:57:THR:HG22	3:B:58:ILE:N	2.15	0.61
3:B:635:GLN:HE21	3:B:635:GLN:HA	1.64	0.61
3:D:606:THR:HG21	3:F:173:ASP:HB2	1.80	0.61
3:D:635:GLN:NE2	3:D:635:GLN:HA	2.15	0.61
3:F:418:ASN:HD21	3:H:337:ARG:NH2	1.93	0.61
3:J:298:PHE:CE2	3:N:355:GLY:HA3	2.34	0.61
3:J:356:THR:HG23	3:N:412:ARG:HH22	1.65	0.61
3:L:341:MET:HB2	3:L:431:ILE:CD1	2.26	0.61
3:N:199:LEU:O	3:N:207:PHE:CE2	2.52	0.61
3:N:273:LEU:C	3:N:273:LEU:HD23	2.25	0.61
3:P:308:ALA:HB1	3:P:383:GLN:NE2	2.13	0.61
3:P:635:GLN:HE21	3:P:635:GLN:HA	1.64	0.61
3:R:199:LEU:O	3:R:207:PHE:CE2	2.52	0.61
3:T:176:ALA:O	3:T:631:THR:HG21	1.99	0.61
6:X:662:LEU:HD22	6:X:662:LEU:H	1.65	0.61

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
6:Y:631:ILE:O	6:Y:640:ARG:HD2	2.00	0.61
6:Y:722:ASN:ND2	6:Y:724:LEU:HB3	2.15	0.61
6:Y:731:GLY:O	6:Y:734:THR:HG22	2.00	0.61
6:Y:1012:ARG:HH12	6:Y:1043:GLY:HA3	1.65	0.61
1:d:20:GLY:HA3	1:d:83:VAL:HG13	1.82	0.61
1:d:45:ARG:CD	3:D:401:GLY:H	2.13	0.61
1:t:59:HIS:CB	3:T:397:VAL:O	2.48	0.61
3:B:329:VAL:HG13	3:B:442:THR:HG21	1.82	0.61
7:C:101:MYR:H71	3:D:200:CYS:SG	2.40	0.61
3:D:536:THR:HB	3:D:570:LYS:HA	1.80	0.61
3:F:297:ALA:HB2	3:F:460:MET:CB	2.26	0.61
3:J:87:PHE:HD1	3:L:121:ALA:HB2	1.61	0.61
3:J:329:VAL:HG13	3:J:442:THR:HG21	1.82	0.61
3:J:358:LEU:HD13	3:J:417:PHE:CZ	2.35	0.61
3:L:329:VAL:HG13	3:L:442:THR:HG21	1.82	0.61
3:L:549:LEU:HD12	3:L:550:SER:N	2.15	0.61
3:N:329:VAL:HG13	3:N:442:THR:HG21	1.82	0.61
3:P:49:ARG:HH12	3:P:115:THR:CG2	2.10	0.61
3:P:171:THR:CG2	3:T:602:SER:OG	2.48	0.61
3:T:491:ALA:CB	3:T:592:THR:HB	2.28	0.61
4:U:263:GLN:HE21	4:U:263:GLN:HA	1.64	0.61
5:W:81:TYR:O	5:W:85:THR:HG23	2.00	0.61
5:W:424:SER:HB3	5:W:425:PRO:CD	2.28	0.61
6:X:218:GLU:C	6:X:219:LEU:HD23	2.25	0.61
6:Y:428:LEU:HD21	6:Y:1208:MET:HE3	1.81	0.61
1:l:24:LEU:HD23	1:l:80:ARG:CD	2.30	0.61
1:f:4:HIS:CE1	1:f:54:THR:HG22	2.35	0.61
1:h:23:THR:HG22	1:h:42:THR:OG1	2.01	0.61
1:t:24:LEU:CG	1:t:38:ALA:CB	2.76	0.61
3:B:549:LEU:HD12	3:B:550:SER:N	2.15	0.61
2:I:5:GLN:O	2:I:6:THR:HB	1.99	0.61
2:K:9:ASN:HB2	3:L:209:ASN:ND2	2.15	0.61
3:N:98:ALA:HB2	3:N:206:GLU:CG	2.29	0.61
3:N:341:MET:O	3:N:344:MET:HB2	2.01	0.61
3:P:82:PHE:HZ	3:T:238:VAL:HG11	1.64	0.61
3:P:298:PHE:CE2	3:T:355:GLY:HA3	2.33	0.61
3:P:329:VAL:HG13	3:P:442:THR:HG21	1.82	0.61
2:Q:9:ASN:HB2	3:R:209:ASN:ND2	2.15	0.61
3:R:273:LEU:C	3:R:273:LEU:HD23	2.25	0.61
3:R:602:SER:OG	3:T:171:THR:CG2	2.48	0.61
3:R:635:GLN:HE21	3:R:635:GLN:HA	1.64	0.61

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:T:330:GLN:HE22	3:T:332:GLY:HA2	1.63	0.61
4:U:140:GLN:O	4:U:144:ILE:HG12	2.00	0.61
5:W:592:MET:H	5:W:593:PRO:CD	2.11	0.61
5:W:930:TRP:C	5:W:931:LEU:HD23	2.26	0.61
6:Y:722:ASN:HB3	6:Y:725:THR:HG23	1.81	0.61
6:Y:778:ASP:CA	6:Y:940:GLN:HB3	2.27	0.61
6:Y:990:ASN:HB2	6:Y:1131:ARG:CA	2.30	0.61
6:Y:1034:ASP:C	6:Y:1036:ASN:H	2.08	0.61
1:l:44:GLY:HA3	3:L:322:TYR:HH	1.65	0.61
1:d:61:ASN:CG	3:D:395:THR:CG2	2.73	0.61
1:p:61:ASN:CG	3:P:395:THR:HG22	2.21	0.61
3:B:491:ALA:CB	3:B:592:THR:HB	2.28	0.61
3:D:237:GLU:HG3	3:F:142:ILE:HD13	1.82	0.61
3:F:68:PHE:HD1	3:F:140:VAL:HG11	1.65	0.61
3:F:139:THR:C	3:F:142:ILE:HG22	2.24	0.61
3:F:238:VAL:CG2	3:H:78:MET:HE1	2.07	0.61
3:F:635:GLN:NE2	3:F:635:GLN:HA	2.15	0.61
7:I:101:MYR:H31	3:J:191:LYS:CG	2.30	0.61
3:J:165:THR:O	3:J:169:MET:HB2	1.99	0.61
3:J:238:VAL:HG11	3:L:82:PHE:HZ	1.64	0.61
3:J:273:LEU:C	3:J:273:LEU:HD23	2.25	0.61
3:J:311:ASN:H	3:J:311:ASN:HD22	1.47	0.61
3:N:311:ASN:H	3:N:311:ASN:HD22	1.46	0.61
3:N:443:PHE:HB2	3:N:457:PRO:CB	2.24	0.61
3:P:215:TYR:C	3:P:217:ALA:H	2.09	0.61
3:P:336:THR:HG22	3:P:456:ASN:HD21	1.65	0.61
3:P:549:LEU:HD12	3:P:550:SER:N	2.15	0.61
3:R:355:GLY:HA3	3:T:298:PHE:CE2	2.33	0.61
4:U:91:THR:HG21	4:U:109:PRO:HG2	1.82	0.61
4:V:163:VAL:O	4:V:258:PRO:HB2	1.99	0.61
5:W:204:VAL:HG13	5:W:238:HIS:HB2	1.81	0.61
5:W:217:LEU:HA	5:W:220:LEU:CD2	2.30	0.61
5:W:489:LYS:HG3	5:W:649:ILE:CG1	2.30	0.61
6:X:915:LEU:N	6:X:915:LEU:HD23	2.16	0.61
6:Y:317:PHE:CZ	6:Y:385:VAL:HG21	2.35	0.61
1:l:4:HIS:NE2	1:l:54:THR:HG22	2.15	0.61
1:h:59:HIS:HB2	3:H:397:VAL:O	2.00	0.61
1:j:45:ARG:O	1:j:58:PRO:O	2.18	0.61
1:r:4:HIS:HE1	1:r:54:THR:CG2	2.13	0.61
3:B:215:TYR:C	3:B:217:ALA:H	2.09	0.61
3:B:372:ALA:HB1	3:B:399:ALA:HA	1.83	0.61

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:D:98:ALA:HB2	3:D:206:GLU:CG	2.29	0.61
3:D:176:ALA:O	3:D:631:THR:HG21	1.99	0.61
3:J:171:THR:CG2	3:N:602:SER:OG	2.48	0.61
3:J:602:SER:OG	3:L:171:THR:CG2	2.48	0.61
3:L:341:MET:O	3:L:344:MET:HB2	2.01	0.61
3:N:491:ALA:CB	3:N:592:THR:HB	2.28	0.61
2:O:9:ASN:HB2	3:P:209:ASN:ND2	2.15	0.61
7:O:101:MYR:H31	3:P:191:LYS:CG	2.30	0.61
3:P:273:LEU:HD23	3:P:273:LEU:C	2.25	0.61
3:T:329:VAL:HG13	3:T:442:THR:HG21	1.82	0.61
3:T:549:LEU:HD12	3:T:550:SER:N	2.15	0.61
4:U:339:LEU:HG	4:U:343:GLN:HE21	1.66	0.61
4:V:268:ARG:HH12	4:V:305:LEU:HA	1.65	0.61
5:W:695:ILE:HD11	5:W:1002:ILE:CG2	2.31	0.61
6:X:416:ILE:CG2	6:X:445:PHE:HE2	2.12	0.61
6:X:518:LEU:HG	6:X:820:VAL:CG2	2.31	0.61
6:X:1078:LEU:HD23	6:X:1078:LEU:N	2.08	0.61
6:Y:210:ILE:HB	6:Y:239:ALA:CB	2.30	0.61
6:Y:354:THR:O	6:Y:356:ASN:N	2.34	0.61
6:Y:982:LEU:C	6:Y:982:LEU:HD23	2.25	0.61
6:Y:1170:PRO:HD2	6:Y:1189:THR:CG2	2.30	0.61
1:l:4:HIS:NE2	1:l:54:THR:CG2	2.63	0.61
1:d:22:LEU:CD1	1:d:83:VAL:CG2	2.48	0.61
1:r:24:LEU:HD12	1:r:40:TYR:O	2.01	0.61
1:t:15:ARG:CZ	3:T:402:ALA:HB2	2.30	0.61
2:A:9:ASN:HB2	3:B:209:ASN:ND2	2.15	0.61
3:B:44:THR:HG22	3:B:45:GLY:N	2.14	0.61
3:D:50:PRO:N	3:D:61:LEU:HD23	2.16	0.61
3:D:549:LEU:HD12	3:D:550:SER:N	2.15	0.61
3:F:57:THR:HG22	3:F:58:ILE:N	2.15	0.61
3:F:602:SER:OG	3:H:171:THR:CG2	2.48	0.61
3:J:635:GLN:HE21	3:J:635:GLN:HA	1.64	0.61
3:L:336:THR:HG22	3:L:456:ASN:HD21	1.65	0.61
7:M:101:MYR:C2	3:N:191:LYS:CE	2.78	0.61
3:N:549:LEU:HD12	3:N:550:SER:N	2.15	0.61
2:O:37:LYS:NZ	3:R:264:SER:OG	2.23	0.61
3:P:635:GLN:NE2	3:P:635:GLN:HA	2.15	0.61
3:R:57:THR:HG22	3:R:58:ILE:N	2.15	0.61
3:R:372:ALA:HB1	3:R:399:ALA:HA	1.83	0.61
3:T:215:TYR:C	3:T:217:ALA:H	2.09	0.61
3:T:341:MET:O	3:T:344:MET:HB2	2.01	0.61

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
4:U:175:MET:HB3	6:X:436:LEU:HA	1.82	0.61
4:V:250:TRP:CE3	6:Y:956:HIS:CE1	2.87	0.61
5:W:838:PRO:HB3	5:W:858:ARG:CZ	2.30	0.61
6:X:595:GLN:HB3	6:X:596:PRO:HD3	1.82	0.61
6:X:1032:ILE:HA	6:X:1038:PRO:HA	1.82	0.61
6:Y:367:ARG:HG3	6:Y:367:ARG:HH11	1.66	0.61
6:Y:777:ASN:HB2	6:Y:942:GLY:CA	2.30	0.61
6:Y:941:TYR:C	6:Y:943:GLY:H	2.09	0.61
1:b:22:LEU:CB	1:b:83:VAL:HG21	2.30	0.61
1:f:43:CYS:HB2	1:f:45:ARG:H	1.63	0.61
1:p:61:ASN:CB	3:P:395:THR:CG2	2.78	0.61
1:r:59:HIS:CA	3:R:398:SER:HB2	2.30	0.61
3:B:330:GLN:HE22	3:B:332:GLY:HA2	1.63	0.61
2:C:5:GLN:O	2:C:6:THR:HB	1.99	0.61
3:D:215:TYR:C	3:D:217:ALA:H	2.09	0.61
3:D:238:VAL:HG11	3:F:82:PHE:HZ	1.64	0.61
3:F:358:LEU:HD13	3:F:417:PHE:CZ	2.35	0.61
3:H:372:ALA:HB1	3:H:399:ALA:HA	1.83	0.61
3:J:57:THR:HG22	3:J:58:ILE:H	1.66	0.61
3:J:536:THR:HB	3:J:570:LYS:HA	1.80	0.61
3:J:549:LEU:HD12	3:J:550:SER:N	2.15	0.61
3:L:456:ASN:C	3:L:456:ASN:HD22	2.09	0.61
3:P:356:THR:HG23	3:T:412:ARG:HH22	1.65	0.61
3:P:380:VAL:HG13	3:P:384:GLN:HG2	1.78	0.61
3:R:540:VAL:HG23	3:R:569:ALA:HB3	1.83	0.61
4:U:52:PRO:HB3	6:X:963:GLN:HE22	1.66	0.61
4:U:215:TRP:CH2	4:U:279:VAL:HG22	2.36	0.61
5:W:149:THR:HA	5:W:152:THR:CG2	2.30	0.61
6:X:957:ILE:C	6:X:959:ARG:N	2.58	0.61
6:X:1122:PRO:HB3	6:X:1159:GLY:HA3	1.82	0.61
6:Y:777:ASN:O	6:Y:778:ASP:HB3	1.99	0.61
6:Y:840:ALA:O	6:Y:843:PRO:HD2	2.00	0.61
1:b:24:LEU:HD11	1:b:80:ARG:NE	2.16	0.61
1:j:23:THR:CB	3:J:323:SER:CA	2.66	0.61
1:n:43:CYS:O	3:N:322:TYR:OH	2.18	0.61
1:t:62:VAL:C	1:t:64:THR:N	2.58	0.61
3:D:341:MET:O	3:D:344:MET:HB2	2.01	0.61
3:F:329:VAL:HG13	3:F:442:THR:HG21	1.82	0.61
3:H:215:TYR:C	3:H:217:ALA:H	2.09	0.61
3:J:292:ARG:HH11	3:J:292:ARG:HG2	1.64	0.61
3:L:358:LEU:HD13	3:L:417:PHE:CZ	2.35	0.61

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:Q:5:GLN:O	2:Q:6:THR:HB	2.00	0.61
3:R:336:THR:HG22	3:R:456:ASN:HD21	1.65	0.61
3:T:358:LEU:HD13	3:T:417:PHE:CZ	2.35	0.61
5:W:639:ALA:CB	5:W:659:PHE:HB3	2.31	0.61
5:W:732:LEU:HD12	5:W:780:LEU:HD13	1.83	0.61
5:W:998:LEU:C	5:W:1000:TYR:H	2.08	0.61
6:X:827:THR:CG2	6:X:831:LEU:HD22	2.30	0.61
6:Y:244:VAL:HG21	6:Y:265:PRO:CA	2.30	0.61
6:Y:332:ALA:O	6:Y:333:PHE:HB2	1.99	0.61
1:b:23:THR:OG1	3:B:323:SER:HA	2.00	0.61
1:p:15:ARG:NE	3:P:406:ASN:CG	2.59	0.61
3:F:341:MET:O	3:F:344:MET:HB2	2.01	0.61
3:N:456:ASN:C	3:N:456:ASN:HD22	2.09	0.61
3:P:337:ARG:NH2	3:T:418:ASN:HD21	1.93	0.61
3:R:154:ARG:HB3	3:R:154:ARG:HH11	1.63	0.61
3:R:412:ARG:HH22	3:T:356:THR:HG23	1.65	0.61
4:U:318:ASN:N	4:U:322:VAL:HG11	2.16	0.61
5:W:266:VAL:CG2	5:W:311:ALA:HB2	2.30	0.61
5:W:518:TYR:CE1	5:W:531:PRO:HG2	2.35	0.61
5:W:823:THR:CG2	5:W:850:LEU:HD13	2.30	0.61
6:X:603:VAL:CG1	6:Y:715:VAL:HG11	2.31	0.61
6:X:628:PRO:HG2	6:X:647:ARG:HD3	1.83	0.61
6:Y:387:CYS:SG	6:Y:389:GLU:HG2	2.40	0.61
6:Y:709:ILE:HA	6:Y:733:MET:CE	2.30	0.61
1:b:44:GLY:HA3	3:B:322:TYR:CE2	2.35	0.61
1:h:59:HIS:HB2	3:H:398:SER:HA	1.81	0.61
3:B:273:LEU:HD23	3:B:273:LEU:C	2.25	0.61
3:D:297:ALA:HB2	3:D:460:MET:CB	2.26	0.61
3:D:431:ILE:HD13	3:D:437:ILE:HG23	1.82	0.61
3:D:456:ASN:C	3:D:456:ASN:HD22	2.09	0.61
3:H:273:LEU:HD23	3:H:273:LEU:C	2.25	0.61
3:J:372:ALA:HB1	3:J:399:ALA:HA	1.83	0.61
3:L:137:ILE:HG12	3:L:142:ILE:HG22	1.83	0.61
3:L:602:SER:OG	3:N:171:THR:CG2	2.48	0.61
3:P:48:TRP:CZ2	3:P:63:ILE:HD11	2.36	0.61
3:P:418:ASN:CG	3:R:337:ARG:HH21	2.09	0.61
3:P:540:VAL:HG23	3:P:569:ALA:HB3	1.83	0.61
3:T:372:ALA:HB1	3:T:399:ALA:HA	1.83	0.61
3:T:380:VAL:HG13	3:T:384:GLN:HG2	1.78	0.61
4:U:33:TYR:CD2	6:X:879:PHE:HD2	2.19	0.61
4:U:339:LEU:HG	4:U:343:GLN:NE2	2.14	0.61

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
4:U:348:THR:HB	4:U:351:GLN:HG3	1.81	0.61
5:W:12:LYS:HD2	5:W:385:THR:O	2.01	0.61
6:X:437:ARG:HH22	6:X:1207:ILE:CA	2.05	0.61
6:Y:272:HIS:HA	6:Y:275:LYS:HE2	1.83	0.61
6:Y:355:LYS:HZ3	6:Y:1156:THR:HG21	1.63	0.61
1:d:61:ASN:HB3	3:D:397:VAL:HA	1.83	0.60
1:n:74:GLN:O	1:n:74:GLN:NE2	2.27	0.60
3:B:341:MET:O	3:B:344:MET:HB2	2.01	0.60
3:B:456:ASN:HD22	3:B:456:ASN:C	2.09	0.60
3:F:274:ILE:HD11	3:F:514:LEU:HD21	1.73	0.60
3:H:154:ARG:HH11	3:H:154:ARG:CB	2.14	0.60
3:H:336:THR:HG22	3:H:456:ASN:HD21	1.65	0.60
3:H:341:MET:O	3:H:344:MET:HB2	2.01	0.60
3:J:337:ARG:NH2	3:N:418:ASN:HD21	1.93	0.60
3:L:48:TRP:CZ2	3:L:63:ILE:HD11	2.36	0.60
3:N:341:MET:HG3	3:N:437:ILE:HG12	1.82	0.60
3:R:178:SER:O	3:R:179:ALA:HB2	2.01	0.60
3:R:341:MET:O	3:R:344:MET:HB2	2.01	0.60
3:T:336:THR:HG22	3:T:456:ASN:HD21	1.65	0.60
4:V:176:ALA:HA	6:Y:407:ARG:HG3	1.82	0.60
5:W:193:THR:HG23	5:W:195:GLY:H	1.66	0.60
5:W:810:LEU:CD2	5:W:1001:ILE:HD13	2.31	0.60
5:W:938:TYR:OH	5:W:1106:GLY:HA2	2.00	0.60
5:W:988:TRP:CZ3	5:W:1026:LEU:HB2	2.36	0.60
5:W:1089:TYR:HA	5:W:1096:PHE:HA	1.83	0.60
5:W:1177:ASN:CB	5:W:1282:THR:HB	2.28	0.60
6:X:466:THR:HA	6:X:809:ALA:CB	2.31	0.60
6:X:534:SER:OG	6:X:543:THR:HG23	2.01	0.60
6:Y:195:ILE:HD12	6:Y:257:ARG:CZ	2.31	0.60
6:Y:1027:LEU:H	6:Y:1027:LEU:CD1	2.07	0.60
1:l:58:PRO:CG	3:L:423:GLN:NE2	2.54	0.60
3:B:635:GLN:NE2	3:B:635:GLN:HA	2.15	0.60
3:D:178:SER:O	3:D:179:ALA:HB2	2.01	0.60
3:D:341:MET:CB	3:D:431:ILE:HD11	2.28	0.60
2:I:37:LYS:NZ	3:L:264:SER:OG	2.23	0.60
3:J:341:MET:O	3:J:344:MET:HB2	2.01	0.60
2:K:5:GLN:O	2:K:6:THR:HB	1.99	0.60
2:O:5:GLN:O	2:O:6:THR:HB	2.00	0.60
3:P:372:ALA:HB1	3:P:399:ALA:HA	1.83	0.60
3:P:412:ARG:HH22	3:R:356:THR:HG23	1.65	0.60
3:R:238:VAL:HG11	3:T:82:PHE:HZ	1.64	0.60

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:R:358:LEU:CD1	3:R:417:PHE:CE1	2.62	0.60
5:W:885:TRP:HA	5:W:888:ILE:HD11	1.83	0.60
5:W:959:ASN:HB3	5:W:962:GLU:OE2	2.00	0.60
6:Y:490:PRO:HD3	6:Y:525:TYR:OH	2.00	0.60
6:Y:798:VAL:HG12	6:Y:799:ASP:N	2.17	0.60
6:Y:985:VAL:HG22	6:Y:1137:PHE:HE2	1.66	0.60
6:Y:1043:GLY:O	6:Y:1072:THR:HA	2.00	0.60
1:l:59:HIS:HB2	3:L:397:VAL:O	2.01	0.60
1:d:22:LEU:CD2	1:d:83:VAL:CG2	2.79	0.60
1:h:44:GLY:O	3:H:399:ALA:CA	2.49	0.60
1:n:58:PRO:CB	3:N:423:GLN:NE2	2.61	0.60
1:p:14:VAL:O	1:p:18:ALA:N	2.35	0.60
3:B:154:ARG:HH11	3:B:154:ARG:CB	2.15	0.60
3:B:336:THR:HG22	3:B:456:ASN:HD21	1.65	0.60
3:D:47:LEU:HD21	3:D:64:VAL:HG11	1.84	0.60
3:D:273:LEU:HD23	3:D:273:LEU:C	2.25	0.60
3:F:44:THR:HG22	3:F:45:GLY:N	2.14	0.60
3:F:418:ASN:CG	3:H:337:ARG:HH21	2.09	0.60
3:H:625:ARG:CG	3:L:197:ASP:OD1	2.49	0.60
3:J:154:ARG:HH11	3:J:154:ARG:CB	2.15	0.60
3:P:341:MET:O	3:P:344:MET:HB2	2.01	0.60
3:R:215:TYR:C	3:R:217:ALA:H	2.09	0.60
3:R:358:LEU:HD13	3:R:417:PHE:CZ	2.35	0.60
3:R:549:LEU:HD12	3:R:550:SER:N	2.15	0.60
2:S:5:GLN:O	2:S:6:THR:HB	2.00	0.60
4:U:143:HIS:NE2	4:U:291:THR:HB	2.17	0.60
4:U:241:LEU:O	6:X:1179:VAL:HG13	2.00	0.60
4:U:251:ASN:CG	4:U:252:ASP:N	2.59	0.60
5:W:398:LEU:CD2	5:W:792:THR:HG21	2.31	0.60
5:W:746:PHE:CZ	5:W:748:ARG:HG3	2.36	0.60
6:X:795:ASP:OD2	6:Y:737:ARG:NH1	2.25	0.60
6:Y:78:ILE:HD12	6:Y:78:ILE:N	2.16	0.60
6:Y:385:VAL:CG2	6:Y:386:GLU:H	2.15	0.60
6:Y:832:ILE:HD12	6:Y:835:HIS:CD2	2.35	0.60
6:Y:858:VAL:O	6:Y:862:GLU:HG2	2.01	0.60
1:b:44:GLY:O	3:B:399:ALA:HB3	2.00	0.60
1:d:73:ARG:C	1:d:73:ARG:HD2	2.27	0.60
1:j:59:HIS:HD1	3:J:397:VAL:HG22	1.66	0.60
1:t:24:LEU:CG	1:t:38:ALA:HB3	2.31	0.60
3:B:68:PHE:HD1	3:B:140:VAL:HG11	1.65	0.60
3:B:341:MET:HG3	3:B:437:ILE:HG12	1.82	0.60

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:D:358:LEU:CG	3:D:417:PHE:CE1	2.79	0.60
3:D:369:SER:OG	3:D:443:PHE:HD1	1.85	0.60
3:D:370:VAL:O	3:D:370:VAL:HG23	2.02	0.60
3:F:48:TRP:CZ2	3:F:63:ILE:HD11	2.36	0.60
3:H:48:TRP:CZ2	3:H:63:ILE:HD11	2.36	0.60
3:H:431:ILE:HD13	3:H:437:ILE:HG23	1.82	0.60
3:J:215:TYR:C	3:J:217:ALA:H	2.09	0.60
3:L:154:ARG:HB3	3:L:154:ARG:HH11	1.63	0.60
3:L:341:MET:CB	3:L:431:ILE:HD11	2.28	0.60
3:N:48:TRP:CZ2	3:N:63:ILE:HD11	2.36	0.60
3:N:369:SER:OG	3:N:443:PHE:HD1	1.85	0.60
3:N:370:VAL:HG23	3:N:370:VAL:O	2.02	0.60
3:N:635:GLN:HE21	3:N:635:GLN:HA	1.64	0.60
3:P:602:SER:OG	3:R:171:THR:CG2	2.48	0.60
3:T:540:VAL:HG23	3:T:569:ALA:HB3	1.83	0.60
4:U:151:SER:HB2	4:U:156:ILE:HG13	1.82	0.60
5:W:398:LEU:H	5:W:398:LEU:CD2	2.13	0.60
5:W:398:LEU:HD11	5:W:788:GLN:HE22	1.64	0.60
6:X:560:LEU:HD11	6:X:729:MET:SD	2.41	0.60
6:Y:385:VAL:HG13	6:Y:1196:ARG:HB2	1.84	0.60
6:Y:437:ARG:HB2	6:Y:439:THR:O	2.02	0.60
6:Y:680:PRO:O	6:Y:748:GLN:HG2	2.00	0.60
1:l:59:HIS:CB	3:L:398:SER:OG	2.48	0.60
1:t:44:GLY:CA	3:T:399:ALA:HB3	2.32	0.60
3:B:297:ALA:HB2	3:B:460:MET:CB	2.26	0.60
3:D:602:SER:OG	3:F:171:THR:CG2	2.48	0.60
3:F:154:ARG:HH11	3:F:154:ARG:CB	2.15	0.60
3:L:57:THR:HG22	3:L:58:ILE:N	2.15	0.60
7:O:101:MYR:H71	3:P:200:CYS:SG	2.40	0.60
3:R:48:TRP:CZ2	3:R:63:ILE:HD11	2.36	0.60
4:V:291:THR:HG21	4:V:296:LEU:HD23	1.83	0.60
5:W:113:THR:HB	5:W:118:LEU:HD12	1.83	0.60
5:W:918:ILE:N	5:W:919:PRO:HD2	2.16	0.60
6:X:1046:VAL:O	6:X:1048:PRO:HD3	2.01	0.60
6:Y:364:ILE:HD12	6:Y:367:ARG:NH1	2.15	0.60
1:l:60:ALA:HB3	3:L:322:TYR:OH	2.00	0.60
1:h:45:ARG:CD	3:H:401:GLY:N	2.64	0.60
3:F:87:PHE:HD1	3:H:121:ALA:HB2	1.61	0.60
3:H:329:VAL:HG13	3:H:442:THR:HG21	1.82	0.60
3:H:549:LEU:HD12	3:H:550:SER:N	2.15	0.60
3:J:370:VAL:HG23	3:J:370:VAL:O	2.02	0.60

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:J:456:ASN:C	3:J:456:ASN:HD22	2.09	0.60
3:L:341:MET:HG3	3:L:437:ILE:HG12	1.82	0.60
3:N:215:TYR:C	3:N:217:ALA:H	2.09	0.60
3:P:167:MET:HA	3:P:167:MET:CE	2.31	0.60
3:P:337:ARG:HH21	3:T:418:ASN:CG	2.09	0.60
3:T:68:PHE:HD1	3:T:140:VAL:HG11	1.65	0.60
3:T:456:ASN:HD22	3:T:456:ASN:C	2.09	0.60
4:U:224:ASN:HD22	4:U:346:LEU:HD11	1.66	0.60
5:W:15:THR:HG22	5:W:273:ASN:OD1	2.01	0.60
5:W:151:ILE:HG13	5:W:152:THR:N	2.16	0.60
5:W:1093:ASN:O	5:W:1094:GLN:HB2	2.00	0.60
6:X:480:LEU:HD22	6:X:758:LEU:HD13	1.84	0.60
6:X:862:GLU:OE1	6:X:955:SER:HB2	2.00	0.60
6:Y:544:VAL:HG12	6:Y:578:ASP:OD2	2.01	0.60
1:b:15:ARG:NH1	3:B:406:ASN:ND2	2.50	0.60
1:h:15:ARG:HH22	3:H:402:ALA:HB3	1.66	0.60
1:j:15:ARG:HE	3:J:406:ASN:CG	2.10	0.60
1:t:59:HIS:HB2	3:T:397:VAL:C	2.27	0.60
3:F:167:MET:HA	3:F:167:MET:CE	2.31	0.60
3:F:540:VAL:HG23	3:F:569:ALA:HB3	1.83	0.60
3:H:370:VAL:O	3:H:370:VAL:HG23	2.02	0.60
3:H:456:ASN:C	3:H:456:ASN:HD22	2.09	0.60
3:J:48:TRP:CZ2	3:J:63:ILE:HD11	2.36	0.60
3:J:53:THR:O	3:J:54:SER:HB3	2.01	0.60
3:J:58:ILE:HG13	3:J:127:LEU:HD23	1.84	0.60
3:J:635:GLN:NE2	3:J:635:GLN:HA	2.15	0.60
3:L:154:ARG:HH11	3:L:154:ARG:CB	2.14	0.60
3:L:370:VAL:O	3:L:370:VAL:HG23	2.02	0.60
3:P:189:PHE:CZ	3:P:193:ILE:HD11	2.37	0.60
3:R:329:VAL:HG13	3:R:442:THR:HG21	1.82	0.60
3:T:46:LYS:HA	3:T:72:SER:HA	1.84	0.60
3:T:341:MET:HG3	3:T:437:ILE:HG12	1.82	0.60
4:U:26:LEU:HA	4:U:71:LEU:CD1	2.28	0.60
4:U:40:ARG:HH22	4:U:69:ASN:ND2	1.98	0.60
4:U:225:ARG:HG3	4:U:225:ARG:HH11	1.67	0.60
5:W:521:ALA:HB1	5:W:531:PRO:HD3	1.84	0.60
6:X:301:LEU:HD11	6:X:968:PHE:HB2	1.84	0.60
6:X:590:ALA:CA	6:X:639:LEU:HD11	2.31	0.60
6:X:925:LEU:HD22	6:X:926:TYR:CD1	2.36	0.60
6:X:1130:ALA:CB	6:Y:104:GLN:CB	2.73	0.60
6:Y:392:ASP:HA	6:Y:1189:THR:O	2.02	0.60

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
6:Y:979:ARG:HH11	6:Y:979:ARG:HG2	1.67	0.60
6:Y:1082:VAL:O	6:Y:1114:PRO:HA	2.02	0.60
1:j:3:LEU:HD13	1:j:3:LEU:O	2.02	0.60
7:A:101:MYR:C2	3:B:191:LYS:CE	2.79	0.60
3:D:329:VAL:HG13	3:D:442:THR:HG21	1.82	0.60
3:D:337:ARG:HH21	3:H:418:ASN:CG	2.09	0.60
3:J:418:ASN:CG	3:L:337:ARG:HH21	2.09	0.60
3:J:540:VAL:HG23	3:J:569:ALA:HB3	1.83	0.60
3:N:540:VAL:HG23	3:N:569:ALA:HB3	1.83	0.60
3:R:370:VAL:HG23	3:R:370:VAL:O	2.02	0.60
4:U:12:ASN:HD22	4:U:13:PHE:N	2.00	0.60
6:X:377:ILE:HB	6:X:380:GLU:HB3	1.83	0.60
6:X:459:LEU:O	6:X:462:ILE:HG22	2.02	0.60
1:l:23:THR:CG2	1:l:23:THR:O	2.48	0.60
1:r:17:ALA:HA	1:r:22:LEU:HD21	1.83	0.60
1:t:22:LEU:CD1	1:t:83:VAL:CG2	2.64	0.60
3:B:48:TRP:CZ2	3:B:63:ILE:HD11	2.36	0.60
3:B:189:PHE:CZ	3:B:193:ILE:HD11	2.37	0.60
3:F:46:LYS:HA	3:F:72:SER:HA	1.84	0.60
3:F:372:ALA:HB1	3:F:399:ALA:HA	1.83	0.60
3:L:68:PHE:HD1	3:L:140:VAL:HG11	1.65	0.60
3:L:189:PHE:CZ	3:L:193:ILE:HD11	2.37	0.60
3:L:418:ASN:CG	3:N:337:ARG:HH21	2.09	0.60
2:O:13:ILE:CG2	2:O:14:THR:N	2.65	0.60
3:P:369:SER:CB	3:P:445:ILE:HA	2.32	0.60
3:R:46:LYS:HA	3:R:72:SER:HA	1.84	0.60
3:T:154:ARG:HB3	3:T:154:ARG:HH11	1.63	0.60
5:W:397:PRO:HB3	5:W:747:THR:CG2	2.32	0.60
5:W:805:VAL:HB	5:W:809:THR:OG1	2.02	0.60
5:W:1102:PRO:HG3	5:W:1127:ILE:HG13	1.84	0.60
6:X:789:ASP:HB3	6:X:933:ASP:HB3	1.82	0.60
6:Y:383:ALA:HB1	6:Y:439:THR:HG23	1.83	0.60
6:Y:687:LEU:O	6:Y:688:THR:HG23	2.01	0.60
6:Y:765:PRO:HD3	6:Y:853:ARG:CD	2.32	0.60
1:b:61:ASN:CG	3:B:397:VAL:HA	2.27	0.60
1:j:59:HIS:HB2	3:J:398:SER:HA	1.79	0.60
3:H:63:ILE:HG21	3:H:70:GLN:CD	2.27	0.60
3:H:369:SER:CB	3:H:445:ILE:HA	2.32	0.60
3:H:540:VAL:HG23	3:H:569:ALA:HB3	1.83	0.60
3:L:540:VAL:HG23	3:L:569:ALA:HB3	1.83	0.60
2:M:13:ILE:CG2	2:M:14:THR:N	2.65	0.60

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:N:68:PHE:HD1	3:N:140:VAL:HG11	1.65	0.60
3:N:154:ARG:HH11	3:N:154:ARG:CB	2.14	0.60
7:O:101:MYR:C2	3:P:191:LYS:CE	2.79	0.60
3:R:154:ARG:HH11	3:R:154:ARG:CB	2.15	0.60
3:R:189:PHE:CZ	3:R:193:ILE:HD11	2.37	0.60
3:T:48:TRP:CZ2	3:T:63:ILE:HD11	2.36	0.60
3:T:101:THR:HA	5:W:406:ASN:CB	2.27	0.60
3:T:369:SER:OG	3:T:443:PHE:HD1	1.85	0.60
4:U:32:CYS:CB	6:X:958:HIS:HE1	1.90	0.60
4:U:296:LEU:O	4:U:297:PRO:C	2.44	0.60
5:W:224:PRO:O	5:W:294:SER:HB2	2.01	0.60
5:W:968:ALA:HB1	5:W:973:LEU:HB2	1.83	0.60
6:X:261:VAL:HG22	6:X:261:VAL:O	2.01	0.60
6:X:384:MET:O	6:X:384:MET:HG3	2.02	0.60
6:Y:188:ALA:HB3	6:Y:763:VAL:HG11	1.83	0.60
6:Y:385:VAL:CG2	6:Y:386:GLU:N	2.64	0.60
6:Y:610:THR:CG2	6:Y:614:GLN:HG2	2.32	0.60
1:b:17:ALA:HA	1:b:22:LEU:HD21	1.84	0.59
1:d:44:GLY:O	3:D:399:ALA:C	2.45	0.59
1:j:59:HIS:CB	3:J:398:SER:CA	2.61	0.59
3:B:369:SER:CB	3:B:445:ILE:HA	2.32	0.59
7:C:101:MYR:C2	3:D:191:LYS:CE	2.79	0.59
3:F:178:SER:O	3:F:179:ALA:HB2	2.01	0.59
3:F:401:GLY:O	3:F:402:ALA:HB3	2.02	0.59
3:F:456:ASN:C	3:F:456:ASN:HD22	2.09	0.59
3:J:336:THR:HG22	3:J:456:ASN:HD21	1.65	0.59
3:J:401:GLY:O	3:J:402:ALA:HB3	2.02	0.59
3:L:147:ARG:HB2	3:L:147:ARG:CZ	2.32	0.59
3:L:358:LEU:CD1	3:L:417:PHE:CE1	2.62	0.59
3:L:369:SER:CB	3:L:445:ILE:HA	2.32	0.59
3:N:63:ILE:HG21	3:N:70:GLN:CD	2.27	0.59
3:N:358:LEU:HD13	3:N:417:PHE:CZ	2.35	0.59
3:T:369:SER:CB	3:T:445:ILE:HA	2.32	0.59
4:U:89:GLY:O	4:U:102:PRO:HA	2.02	0.59
4:U:161:VAL:CG2	4:U:263:GLN:HB2	2.31	0.59
4:U:167:PRO:O	4:U:169:ALA:N	2.35	0.59
4:U:267:ILE:HA	4:U:326:LEU:O	2.01	0.59
4:V:178:THR:OG1	6:Y:407:ARG:HB3	2.02	0.59
5:W:409:ILE:HG23	5:W:413:GLN:CD	2.27	0.59
5:W:477:ARG:HA	5:W:650:VAL:CG1	2.31	0.59
6:X:202:THR:HG23	6:X:249:TRP:NE1	2.17	0.59

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
6:X:677:TYR:CD1	6:X:678:PRO:HD2	2.38	0.59
6:X:1019:THR:HB	6:Y:1209:TYR:CD1	2.37	0.59
1:b:58:PRO:O	3:B:398:SER:OG	2.20	0.59
1:t:61:ASN:HD22	3:T:395:THR:CG2	2.10	0.59
3:B:147:ARG:HB2	3:B:147:ARG:CZ	2.33	0.59
3:D:189:PHE:CZ	3:D:193:ILE:HD11	2.37	0.59
3:D:418:ASN:CG	3:F:337:ARG:HH21	2.09	0.59
3:D:469:VAL:HG23	3:F:575:MET:HG3	1.84	0.59
3:F:336:THR:HG22	3:F:456:ASN:HD21	1.65	0.59
7:I:101:MYR:C2	3:J:191:LYS:CE	2.79	0.59
3:J:431:ILE:HD13	3:J:437:ILE:HG23	1.82	0.59
3:J:443:PHE:HB2	3:J:457:PRO:CB	2.24	0.59
3:L:63:ILE:HG21	3:L:70:GLN:CD	2.27	0.59
3:N:178:SER:O	3:N:179:ALA:HB2	2.01	0.59
3:P:287:PRO:HB3	3:P:509:LEU:CD2	2.32	0.59
3:P:456:ASN:HD22	3:P:456:ASN:C	2.09	0.59
3:T:147:ARG:HB2	3:T:147:ARG:CZ	2.32	0.59
4:U:241:LEU:CA	6:X:1179:VAL:CG1	2.80	0.59
5:W:79:LEU:HD23	5:W:79:LEU:N	2.16	0.59
5:W:220:LEU:HD12	5:W:241:LEU:HD12	1.84	0.59
6:X:959:ARG:O	6:X:962:GLN:HB3	2.02	0.59
6:X:1156:THR:CG2	6:X:1161:ASN:HB3	2.32	0.59
6:Y:410:HIS:HD2	6:Y:411:GLN:H	1.46	0.59
6:Y:899:ALA:HB2	6:Y:929:PRO:HG2	1.83	0.59
1:h:15:ARG:NH1	3:H:406:ASN:CB	2.48	0.59
3:B:287:PRO:HB3	3:B:509:LEU:CD2	2.32	0.59
2:E:13:ILE:CG2	2:E:14:THR:N	2.65	0.59
3:F:369:SER:CB	3:F:445:ILE:HA	2.32	0.59
7:I:101:MYR:H71	3:J:200:CYS:SG	2.40	0.59
3:N:372:ALA:HB1	3:N:399:ALA:HA	1.83	0.59
3:P:401:GLY:O	3:P:402:ALA:HB3	2.02	0.59
3:P:472:ARG:HH11	3:R:530:ARG:HG3	1.61	0.59
3:R:456:ASN:C	3:R:456:ASN:HD22	2.09	0.59
3:T:178:SER:O	3:T:179:ALA:HB2	2.01	0.59
4:U:117:ASN:HD21	4:U:308:TRP:CD1	2.21	0.59
5:W:713:VAL:HG23	5:W:735:LEU:HD13	1.83	0.59
5:W:1054:SER:HB3	5:W:1096:PHE:HE1	1.67	0.59
5:W:1236:PRO:HA	5:W:1290:PRO:O	2.02	0.59
6:X:302:LEU:O	6:X:302:LEU:HD23	2.02	0.59
6:X:315:SER:CB	6:X:331:ARG:HD2	2.31	0.59
6:X:653:PRO:CD	6:X:773:ARG:HB2	2.32	0.59

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
6:X:669:PHE:O	6:X:670:ILE:HD13	2.03	0.59
6:X:1087:TYR:HB3	6:X:1121:ILE:HG12	1.83	0.59
6:X:1149:THR:HG21	6:X:1162:ALA:H	1.66	0.59
6:Y:172:PRO:HD2	6:Y:847:SER:HA	1.83	0.59
1:h:23:THR:HG21	3:H:323:SER:HB3	1.82	0.59
1:j:45:ARG:CD	3:J:401:GLY:N	2.65	0.59
3:D:46:LYS:HA	3:D:72:SER:HA	1.84	0.59
3:H:178:SER:O	3:H:179:ALA:HB2	2.01	0.59
3:H:481:VAL:HG12	3:H:482:ASP:N	2.17	0.59
3:J:274:ILE:HD11	3:J:514:LEU:HD21	1.73	0.59
3:J:337:ARG:HH21	3:N:418:ASN:CG	2.09	0.59
3:J:356:THR:CG2	3:N:412:ARG:NH2	2.63	0.59
3:L:412:ARG:HH22	3:N:356:THR:HG23	1.65	0.59
3:N:199:LEU:HD12	3:N:203:TYR:CD2	2.38	0.59
7:O:101:MYR:C2	3:P:191:LYS:HE2	2.27	0.59
3:P:63:ILE:HG21	3:P:70:GLN:CD	2.27	0.59
3:P:147:ARG:HB2	3:P:147:ARG:CZ	2.32	0.59
3:R:443:PHE:HB2	3:R:457:PRO:CB	2.24	0.59
3:R:521:VAL:HG23	3:R:577:VAL:HG22	1.85	0.59
4:V:272:PRO:HB2	4:V:394:PHE:CE1	2.38	0.59
5:W:643:HIS:N	5:W:643:HIS:CD2	2.70	0.59
5:W:836:THR:CG2	5:W:854:MET:HB2	2.31	0.59
6:X:506:GLN:NE2	6:X:724:LEU:HD21	2.18	0.59
6:Y:124:ALA:HB3	6:Y:135:HIS:CD2	2.37	0.59
1:b:60:ALA:O	1:b:61:ASN:C	2.45	0.59
1:f:17:ALA:O	1:f:20:GLY:N	2.35	0.59
1:f:24:LEU:HD22	1:f:80:ARG:HD3	1.85	0.59
1:j:5:MET:HA	1:j:5:MET:CE	2.23	0.59
1:t:45:ARG:O	1:t:58:PRO:C	2.46	0.59
3:B:401:GLY:O	3:B:402:ALA:HB3	2.02	0.59
3:B:431:ILE:HD13	3:B:437:ILE:HG23	1.82	0.59
3:D:147:ARG:HB2	3:D:147:ARG:CZ	2.32	0.59
3:D:171:THR:CG2	3:H:602:SER:OG	2.48	0.59
3:D:337:ARG:NH2	3:H:418:ASN:HD21	1.93	0.59
3:F:73:PHE:HB2	3:F:119:VAL:HG21	1.85	0.59
3:F:189:PHE:CZ	3:F:193:ILE:HD11	2.37	0.59
3:H:46:LYS:HA	3:H:72:SER:HA	1.84	0.59
3:H:369:SER:OG	3:H:443:PHE:HD1	1.84	0.59
3:J:178:SER:O	3:J:179:ALA:HB2	2.01	0.59
3:J:521:VAL:HG23	3:J:577:VAL:HG22	1.85	0.59
3:L:178:SER:O	3:L:179:ALA:HB2	2.01	0.59

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:N:449:THR:CG2	3:N:450:SER:H	2.16	0.59
3:P:178:SER:O	3:P:179:ALA:HB2	2.01	0.59
3:R:481:VAL:HG12	3:R:482:ASP:N	2.17	0.59
2:S:13:ILE:CG2	2:S:14:THR:N	2.65	0.59
5:W:43:LEU:CD1	5:W:64:PRO:HG2	2.33	0.59
5:W:177:TYR:O	5:W:197:HIS:HA	2.02	0.59
5:W:564:LEU:HD13	5:W:570:VAL:CG1	2.33	0.59
5:W:991:LEU:HG	5:W:997:TRP:CZ3	2.37	0.59
5:W:1251:THR:HG22	5:W:1256:PRO:HA	1.84	0.59
6:X:294:ARG:HD3	6:X:885:LEU:O	2.02	0.59
6:Y:1203:THR:O	6:Y:1203:THR:HG23	2.02	0.59
1:j:8:GLN:HE21	3:L:302:GLU:CD	2.10	0.59
1:n:22:LEU:CB	1:n:83:VAL:HG21	2.33	0.59
1:p:59:HIS:CD2	3:P:423:GLN:O	2.56	0.59
1:p:59:HIS:NE2	3:P:424:ALA:HA	2.18	0.59
2:A:13:ILE:CG2	2:A:14:THR:N	2.65	0.59
7:A:101:MYR:H71	3:B:200:CYS:SG	2.40	0.59
2:C:13:ILE:CG2	2:C:14:THR:N	2.65	0.59
3:D:271:LEU:HG	3:D:517:LEU:CB	2.30	0.59
3:D:356:THR:CG2	3:H:412:ARG:NH2	2.63	0.59
3:F:189:PHE:CZ	3:F:193:ILE:CD1	2.86	0.59
3:F:199:LEU:HD12	3:F:203:TYR:CD2	2.38	0.59
3:F:287:PRO:HB3	3:F:509:LEU:CD2	2.32	0.59
3:F:341:MET:CB	3:F:437:ILE:HG23	2.29	0.59
3:F:449:THR:CG2	3:F:450:SER:N	2.66	0.59
3:H:73:PHE:HB2	3:H:119:VAL:HG21	1.85	0.59
3:H:189:PHE:CZ	3:H:193:ILE:CD1	2.86	0.59
3:H:199:LEU:HD12	3:H:203:TYR:CD2	2.38	0.59
3:H:199:LEU:HD12	3:H:203:TYR:HD2	1.68	0.59
3:H:449:THR:CG2	3:H:450:SER:H	2.16	0.59
3:J:189:PHE:CZ	3:J:193:ILE:CD1	2.86	0.59
3:N:199:LEU:HD12	3:N:203:TYR:HD2	1.68	0.59
3:N:369:SER:CB	3:N:445:ILE:HA	2.32	0.59
3:N:449:THR:CG2	3:N:450:SER:N	2.66	0.59
4:U:37:TRP:HA	4:U:37:TRP:CE3	2.37	0.59
4:U:59:PRO:HB2	4:U:373:MET:HG3	1.84	0.59
4:U:272:PRO:HB2	4:U:394:PHE:CE1	2.38	0.59
4:V:184:HIS:NE2	6:Y:436:LEU:HD11	2.17	0.59
5:W:843:LEU:HD23	5:W:854:MET:HE1	1.83	0.59
5:W:1209:ASP:HB2	5:W:1219:ILE:HG22	1.84	0.59
6:X:219:LEU:HB2	6:X:222:PHE:CD2	2.38	0.59

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
6:X:1059:PHE:O	6:X:1063:VAL:HG22	2.03	0.59
6:Y:249:TRP:HH2	6:Y:342:LEU:HD13	1.67	0.59
6:Y:277:VAL:HG22	6:Y:299:LEU:CD2	2.26	0.59
6:Y:355:LYS:NZ	6:Y:1161:ASN:OD1	2.31	0.59
6:Y:672:SER:O	6:Y:673:ASN:C	2.46	0.59
1:b:3:LEU:HD13	1:b:3:LEU:O	2.03	0.59
1:n:58:PRO:HG2	3:N:423:GLN:HG3	1.85	0.59
3:B:73:PHE:HB2	3:B:119:VAL:HG21	1.85	0.59
3:B:199:LEU:HD12	3:B:203:TYR:HD2	1.68	0.59
3:D:199:LEU:HD12	3:D:203:TYR:CD2	2.38	0.59
3:D:260:SER:HB3	2:E:33:ALA:HB2	1.85	0.59
3:D:372:ALA:HB1	3:D:399:ALA:HA	1.83	0.59
3:F:369:SER:OG	3:F:443:PHE:HD1	1.85	0.59
2:G:13:ILE:CG2	2:G:14:THR:N	2.65	0.59
3:H:92:TRP:CE2	3:H:156:VAL:HG23	2.38	0.59
2:I:13:ILE:CG2	2:I:14:THR:N	2.65	0.59
2:I:33:ALA:HB2	3:N:260:SER:HB3	1.85	0.59
3:J:73:PHE:HB2	3:J:119:VAL:HG21	1.85	0.59
3:L:369:SER:OG	3:L:443:PHE:HD1	1.85	0.59
3:L:449:THR:CG2	3:L:450:SER:H	2.16	0.59
3:N:103:CYS:SG	3:N:152:ARG:NH2	2.61	0.59
3:N:401:GLY:O	3:N:402:ALA:HB3	2.02	0.59
3:P:199:LEU:HD12	3:P:203:TYR:CD2	2.38	0.59
3:R:147:ARG:HB2	3:R:147:ARG:CZ	2.32	0.59
3:R:199:LEU:HD12	3:R:203:TYR:HD2	1.68	0.59
3:R:372:ALA:HB2	3:R:399:ALA:HA	1.85	0.59
3:T:57:THR:HG22	3:T:58:ILE:N	2.15	0.59
3:T:92:TRP:CE2	3:T:156:VAL:HG23	2.38	0.59
3:T:199:LEU:HD12	3:T:203:TYR:HD2	1.68	0.59
3:T:274:ILE:HD11	3:T:514:LEU:HD21	1.73	0.59
3:T:401:GLY:O	3:T:402:ALA:HB3	2.02	0.59
3:T:628:GLN:HA	3:T:631:THR:HG22	1.85	0.59
4:V:156:ILE:HD12	4:V:300:ILE:HD13	1.83	0.59
6:X:277:VAL:HG13	6:X:278:LEU:N	2.18	0.59
6:X:865:VAL:HG11	6:X:958:HIS:HB2	1.85	0.59
6:Y:302:LEU:HD22	6:Y:905:VAL:HG22	1.84	0.59
6:Y:626:LEU:HD11	6:Y:776:GLN:NE2	2.18	0.59
6:Y:655:GLU:OE1	6:Y:674:ARG:HD3	2.02	0.59
6:Y:1069:ALA:O	6:Y:1071:ARG:N	2.35	0.59
6:Y:1156:THR:HG23	6:Y:1161:ASN:HB2	1.85	0.59
1:h:58:PRO:CG	3:H:423:GLN:HG3	2.32	0.59

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:r:22:LEU:HD22	1:r:22:LEU:N	2.17	0.59
3:B:63:ILE:HG21	3:B:70:GLN:CD	2.27	0.59
3:B:370:VAL:HG23	3:B:370:VAL:O	2.02	0.59
3:D:189:PHE:CZ	3:D:193:ILE:CD1	2.86	0.59
3:D:252:LEU:O	3:D:256:SER:HB3	2.03	0.59
3:D:412:ARG:NH2	3:F:356:THR:CG2	2.63	0.59
3:F:147:ARG:HB2	3:F:147:ARG:CZ	2.32	0.59
3:F:215:TYR:C	3:F:217:ALA:H	2.09	0.59
3:F:350:THR:O	3:F:350:THR:HG23	2.03	0.59
3:H:401:GLY:O	3:H:402:ALA:HB3	2.02	0.59
3:H:449:THR:CG2	3:H:450:SER:N	2.66	0.59
3:L:46:LYS:HA	3:L:72:SER:HA	1.84	0.59
3:L:252:LEU:O	3:L:256:SER:HB3	2.03	0.59
2:M:28:SER:HA	3:N:216:PRO:HD2	1.85	0.59
3:N:46:LYS:HA	3:N:72:SER:HA	1.84	0.59
3:N:73:PHE:HB2	3:N:119:VAL:HG21	1.85	0.59
3:P:412:ARG:NH2	3:R:356:THR:CG2	2.63	0.59
2:Q:13:ILE:CG2	2:Q:14:THR:N	2.65	0.59
3:R:252:LEU:O	3:R:256:SER:HB3	2.03	0.59
3:R:271:LEU:HG	3:R:517:LEU:CB	2.30	0.59
3:R:431:ILE:HD13	3:R:437:ILE:HG23	1.82	0.59
3:T:73:PHE:HB2	3:T:119:VAL:HG21	1.85	0.59
3:T:562:TYR:HB2	3:T:563:PRO:HD3	1.85	0.59
4:U:214:ILE:HD13	4:U:232:PHE:HB2	1.85	0.59
4:U:275:GLN:NE2	4:U:387:GLN:HB3	2.17	0.59
5:W:4:VAL:HG22	5:W:347:GLN:HB2	1.85	0.59
5:W:175:CYS:C	5:W:177:TYR:H	2.10	0.59
5:W:426:LEU:HD23	5:W:710:ILE:O	2.02	0.59
5:W:474:ASP:O	5:W:477:ARG:HB3	2.03	0.59
5:W:498:ALA:O	5:W:501:VAL:HG12	2.03	0.59
6:X:248:ASN:HB3	6:X:251:SER:HB3	1.84	0.59
6:Y:412:ASP:OD1	6:Y:414:THR:HG22	2.03	0.59
6:Y:760:SER:OG	6:Y:850:MET:HE2	2.03	0.59
6:Y:902:MET:HG3	6:Y:924:ALA:HB2	1.85	0.59
6:Y:987:THR:HG21	6:Y:993:TRP:CH2	2.37	0.59
1:f:4:HIS:CE1	1:f:54:THR:CG2	2.85	0.59
1:f:57:ALA:HB1	1:f:58:PRO:HD2	1.85	0.59
1:d:60:ALA:N	3:D:397:VAL:O	2.35	0.59
1:n:74:GLN:NE2	1:n:74:GLN:HA	2.18	0.59
1:r:59:HIS:HB2	3:R:398:SER:CB	2.08	0.59
3:B:212:ILE:HD11	3:B:220:PRO:HB3	1.85	0.59

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:C:33:ALA:HB2	3:H:260:SER:HB3	1.85	0.59
3:D:540:VAL:HG23	3:D:569:ALA:HB3	1.83	0.59
2:E:28:SER:HA	3:F:216:PRO:HD2	1.85	0.59
2:G:28:SER:HA	3:H:216:PRO:HD2	1.85	0.59
3:H:294:TYR:OH	3:H:502:LEU:HD23	2.03	0.59
3:J:147:ARG:HB2	3:J:147:ARG:CZ	2.32	0.59
3:J:356:THR:HG23	3:N:412:ARG:NH2	2.18	0.59
3:L:199:LEU:HD12	3:L:203:TYR:CD2	2.38	0.59
3:L:412:ARG:NH2	3:N:356:THR:HG23	2.18	0.59
3:L:449:THR:CG2	3:L:450:SER:N	2.66	0.59
3:N:92:TRP:CE2	3:N:156:VAL:HG23	2.38	0.59
3:N:562:TYR:HB2	3:N:563:PRO:HD3	1.85	0.59
3:P:154:ARG:HH11	3:P:154:ARG:CB	2.14	0.59
3:P:252:LEU:O	3:P:256:SER:HB3	2.03	0.59
3:P:358:LEU:HD13	3:P:417:PHE:CZ	2.35	0.59
3:P:370:VAL:O	3:P:370:VAL:HG23	2.02	0.59
3:R:212:ILE:HD11	3:R:220:PRO:HB3	1.85	0.59
3:R:412:ARG:NH2	3:T:356:THR:HG23	2.18	0.59
3:T:449:THR:CG2	3:T:450:SER:H	2.16	0.59
3:T:521:VAL:HG23	3:T:577:VAL:HG22	1.85	0.59
5:W:150:ALA:O	5:W:164:VAL:HG11	2.02	0.59
5:W:831:LEU:O	5:W:852:VAL:HA	2.02	0.59
5:W:987:SER:HA	5:W:1124:SER:HB3	1.85	0.59
6:X:297:VAL:O	6:X:301:LEU:HG	2.03	0.59
6:X:597:GLU:HG3	6:X:636:ALA:HB1	1.84	0.59
6:X:931:ILE:HG12	6:Y:689:VAL:HG11	1.85	0.59
6:Y:119:CYS:HB3	6:Y:123:ASN:N	2.18	0.59
6:Y:387:CYS:HB3	6:Y:1194:LEU:HD21	1.84	0.59
1:f:8:GLN:CD	3:H:302:GLU:CG	2.62	0.59
1:n:15:ARG:NH1	3:N:406:ASN:CB	2.47	0.59
1:n:45:ARG:CD	3:N:400:ALA:CA	2.75	0.59
3:B:178:SER:O	3:B:179:ALA:HB2	2.01	0.59
3:B:252:LEU:O	3:B:256:SER:HB3	2.03	0.59
3:D:294:TYR:OH	3:D:502:LEU:HD23	2.03	0.59
3:D:449:THR:CG2	3:D:450:SER:H	2.16	0.59
3:F:92:TRP:CE2	3:F:156:VAL:HG23	2.38	0.59
3:F:412:ARG:NH2	3:H:356:THR:HG23	2.18	0.59
3:H:189:PHE:CZ	3:H:193:ILE:HD11	2.37	0.59
3:H:628:GLN:HA	3:H:631:THR:HG22	1.85	0.59
3:J:152:ARG:HH11	3:J:152:ARG:HA	1.68	0.59
3:J:294:TYR:OH	3:J:502:LEU:HD23	2.03	0.59

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:J:369:SER:CB	3:J:445:ILE:HA	2.32	0.59
3:L:372:ALA:HB2	3:L:399:ALA:HA	1.85	0.59
3:L:562:TYR:HB2	3:L:563:PRO:HD3	1.85	0.59
3:N:49:ARG:NH2	3:N:64:VAL:HG11	2.18	0.59
3:N:147:ARG:HB2	3:N:147:ARG:CZ	2.33	0.59
3:N:628:GLN:HA	3:N:631:THR:HG22	1.85	0.59
3:R:189:PHE:CZ	3:R:193:ILE:CD1	2.86	0.59
3:R:205:ASN:HA	3:R:208:TYR:HB3	1.85	0.59
3:R:294:TYR:OH	3:R:502:LEU:HD23	2.03	0.59
3:R:369:SER:CB	3:R:445:ILE:HA	2.32	0.59
3:T:154:ARG:HH11	3:T:154:ARG:CB	2.14	0.59
4:U:276:HIS:HB3	4:U:366:ILE:CG2	2.33	0.59
5:W:63:PRO:HB2	5:W:64:PRO:HD3	1.84	0.59
5:W:1053:THR:HG22	5:W:1095:GLU:HB3	1.85	0.59
6:X:473:VAL:HG12	6:X:474:GLU:N	2.18	0.59
6:X:924:ALA:O	6:X:926:TYR:N	2.36	0.59
6:Y:106:MET:HE3	6:Y:108:VAL:HG21	1.85	0.59
6:Y:210:ILE:HD11	6:Y:1087:TYR:CD1	2.38	0.59
1:b:44:GLY:CA	3:B:399:ALA:HB3	2.32	0.58
1:h:15:ARG:CZ	3:H:402:ALA:HB2	2.32	0.58
1:j:44:GLY:CA	3:J:399:ALA:HB3	2.33	0.58
1:j:46:TYR:HD1	1:j:58:PRO:HA	1.67	0.58
1:r:15:ARG:NH2	3:R:402:ALA:CB	2.66	0.58
1:r:21:ARG:H	1:r:22:LEU:HD23	1.67	0.58
3:B:189:PHE:CZ	3:B:193:ILE:CD1	2.86	0.58
3:B:562:TYR:HB2	3:B:563:PRO:HD3	1.85	0.58
3:D:48:TRP:N	3:D:48:TRP:CD1	2.70	0.58
3:D:372:ALA:HB3	3:D:444:LEU:CD1	2.33	0.58
3:D:412:ARG:NH2	3:F:356:THR:HG23	2.18	0.58
3:F:370:VAL:HG23	3:F:370:VAL:O	2.02	0.58
3:F:412:ARG:HH22	3:H:356:THR:HG23	1.65	0.58
3:H:49:ARG:NH2	3:H:64:VAL:HG11	2.18	0.58
3:H:484:ILE:HG22	3:H:485:VAL:N	2.19	0.58
7:I:101:MYR:C2	3:J:191:LYS:HE2	2.27	0.58
3:J:372:ALA:HB3	3:J:444:LEU:CD1	2.33	0.58
2:K:28:SER:HA	3:L:216:PRO:HD2	1.85	0.58
3:L:92:TRP:CE2	3:L:156:VAL:HG23	2.38	0.58
3:N:350:THR:O	3:N:350:THR:HG23	2.03	0.58
3:N:484:ILE:HG22	3:N:485:VAL:N	2.18	0.58
3:P:46:LYS:HA	3:P:72:SER:HA	1.84	0.58
3:P:205:ASN:HA	3:P:208:TYR:HB3	1.85	0.58

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:P:237:GLU:CD	3:R:46:LYS:HZ3	2.11	0.58
3:P:521:VAL:HG23	3:P:577:VAL:HG22	1.85	0.58
3:R:63:ILE:HG21	3:R:70:GLN:CD	2.27	0.58
3:R:369:SER:OG	3:R:443:PHE:HD1	1.85	0.58
3:T:189:PHE:CZ	3:T:193:ILE:HD11	2.37	0.58
4:U:225:ARG:HA	4:U:233:TYR:CZ	2.38	0.58
5:W:23:LEU:HD23	5:W:23:LEU:C	2.28	0.58
5:W:426:LEU:O	5:W:702:ASP:HB3	2.02	0.58
6:X:291:LEU:HD12	6:X:891:PHE:CB	2.32	0.58
6:Y:467:ALA:O	6:Y:471:PRO:HD3	2.03	0.58
1:l:44:GLY:O	3:L:399:ALA:CA	2.51	0.58
1:j:59:HIS:HA	3:J:398:SER:OG	2.02	0.58
1:t:23:THR:HG1	3:T:323:SER:HA	1.62	0.58
1:t:24:LEU:HD21	1:t:35:PHE:HD1	1.60	0.58
1:t:45:ARG:CD	3:T:400:ALA:CA	2.74	0.58
3:B:46:LYS:HA	3:B:72:SER:HA	1.84	0.58
3:B:92:TRP:CE2	3:B:156:VAL:HG23	2.38	0.58
3:B:199:LEU:HD12	3:B:203:TYR:CD2	2.38	0.58
3:B:358:LEU:HD13	3:B:417:PHE:CZ	2.35	0.58
3:D:199:LEU:HD12	3:D:203:TYR:HD2	1.68	0.58
3:D:369:SER:CB	3:D:445:ILE:HA	2.32	0.58
3:D:575:MET:HG3	3:H:469:VAL:HG23	1.85	0.58
3:F:49:ARG:NH2	3:F:64:VAL:HG11	2.18	0.58
3:F:63:ILE:HG21	3:F:70:GLN:CD	2.27	0.58
3:F:199:LEU:HD12	3:F:203:TYR:HD2	1.68	0.58
3:H:252:LEU:O	3:H:256:SER:HB3	2.03	0.58
3:J:63:ILE:HG21	3:J:70:GLN:CD	2.27	0.58
3:J:252:LEU:O	3:J:256:SER:HB3	2.03	0.58
3:J:369:SER:HB3	3:J:445:ILE:HA	1.86	0.58
3:N:189:PHE:CZ	3:N:193:ILE:HD11	2.37	0.58
3:P:87:PHE:HD1	3:R:121:ALA:HB2	1.61	0.58
3:T:63:ILE:HG21	3:T:70:GLN:CD	2.27	0.58
3:T:252:LEU:O	3:T:256:SER:HB3	2.03	0.58
4:U:327:ASP:N	4:U:399:GLN:HG3	2.18	0.58
5:W:204:VAL:CG1	5:W:238:HIS:HB2	2.34	0.58
5:W:816:ASN:ND2	5:W:845:LEU:HD21	2.18	0.58
5:W:992:SER:O	5:W:997:TRP:HZ3	1.86	0.58
6:X:215:LEU:H	6:X:215:LEU:CD2	2.16	0.58
6:Y:571:LEU:HD11	6:Y:705:ILE:HD11	1.85	0.58
6:Y:829:THR:HG23	6:Y:830:ASN:H	1.68	0.58
1:b:59:HIS:CG	3:B:397:VAL:O	2.53	0.58

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:f:15:ARG:CZ	3:F:402:ALA:HB1	2.30	0.58
1:p:75:PRO:HA	1:p:76:ASN:C	2.28	0.58
1:r:19:ALA:HB1	1:r:21:ARG:HH12	1.68	0.58
3:B:49:ARG:NH2	3:B:64:VAL:HG11	2.18	0.58
3:B:294:TYR:OH	3:B:502:LEU:HD23	2.03	0.58
3:B:350:THR:O	3:B:350:THR:HG23	2.03	0.58
3:B:484:ILE:HG22	3:B:485:VAL:N	2.19	0.58
3:D:350:THR:O	3:D:350:THR:HG23	2.03	0.58
3:F:294:TYR:OH	3:F:502:LEU:HD23	2.03	0.58
3:F:380:VAL:HG22	3:N:379:SER:HB3	1.85	0.58
3:F:449:THR:CG2	3:F:450:SER:H	2.16	0.58
3:J:189:PHE:CZ	3:J:193:ILE:HD11	2.37	0.58
3:J:469:VAL:HG23	3:L:575:MET:HG3	1.84	0.58
3:L:49:ARG:NH2	3:L:64:VAL:HG11	2.19	0.58
3:L:401:GLY:O	3:L:402:ALA:HB3	2.02	0.58
3:L:628:GLN:HA	3:L:631:THR:HG22	1.85	0.58
3:N:287:PRO:HB3	3:N:509:LEU:CD2	2.32	0.58
3:P:57:THR:O	3:P:58:ILE:HB	2.02	0.58
3:R:199:LEU:HD12	3:R:203:TYR:CD2	2.38	0.58
3:R:412:ARG:NH2	3:T:356:THR:CG2	2.63	0.58
3:R:418:ASN:CG	3:T:337:ARG:HH21	2.09	0.58
3:T:199:LEU:HD12	3:T:203:TYR:CD2	2.38	0.58
3:T:484:ILE:HG22	3:T:485:VAL:N	2.19	0.58
4:U:79:PHE:HE2	4:U:138:ILE:HD11	1.69	0.58
4:U:86:SER:CB	4:U:91:THR:HB	2.33	0.58
4:V:178:THR:CB	6:Y:407:ARG:HB3	2.33	0.58
5:W:15:THR:HG23	5:W:317:GLY:HA2	1.85	0.58
6:X:627:ASN:ND2	6:X:630:LEU:HG	2.17	0.58
1:r:44:GLY:O	3:R:399:ALA:O	2.22	0.58
1:t:75:PRO:HA	1:t:76:ASN:C	2.28	0.58
3:B:152:ARG:HH11	3:B:152:ARG:HA	1.69	0.58
3:B:551:GLN:HA	3:B:551:GLN:OE1	2.04	0.58
3:D:154:ARG:HH11	3:D:154:ARG:CB	2.15	0.58
3:D:380:VAL:HG13	3:D:381:PRO:HD2	1.85	0.58
3:D:521:VAL:HG23	3:D:577:VAL:HG22	1.85	0.58
3:F:369:SER:HB3	3:F:445:ILE:HA	1.86	0.58
3:F:484:ILE:HG22	3:F:485:VAL:N	2.19	0.58
3:F:551:GLN:HA	3:F:551:GLN:OE1	2.04	0.58
3:H:271:LEU:HG	3:H:517:LEU:CB	2.30	0.58
3:H:372:ALA:HB3	3:H:444:LEU:CD1	2.33	0.58
3:J:271:LEU:HG	3:J:517:LEU:CB	2.30	0.58

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:J:346:ASP:HB3	3:J:428:ARG:N	2.15	0.58
3:J:412:ARG:NH2	3:L:356:THR:HG23	2.18	0.58
3:J:575:MET:HG3	3:N:469:VAL:HG23	1.85	0.58
3:L:152:ARG:HA	3:L:152:ARG:HH11	1.69	0.58
3:L:625:ARG:HG2	3:R:197:ASP:OD1	2.03	0.58
3:N:297:ALA:HB2	3:N:460:MET:CB	2.26	0.58
3:P:189:PHE:CZ	3:P:193:ILE:CD1	2.86	0.58
3:P:369:SER:HB3	3:P:445:ILE:HA	1.86	0.58
3:R:401:GLY:O	3:R:402:ALA:HB3	2.02	0.58
3:R:628:GLN:HA	3:R:631:THR:HG22	1.85	0.58
2:S:28:SER:HA	3:T:216:PRO:HD2	1.85	0.58
4:U:198:GLN:OE1	6:X:661:THR:HG21	2.03	0.58
4:V:270:ILE:HD11	4:V:327:ASP:HB2	1.84	0.58
5:W:823:THR:HG22	5:W:847:PRO:HD2	1.85	0.58
6:X:718:THR:HG23	6:X:726:ARG:NH1	2.08	0.58
6:X:874:CYS:HB2	6:X:901:ILE:HD13	1.84	0.58
6:Y:415:GLN:HE21	6:Y:662:LEU:CD2	2.16	0.58
1:f:61:ASN:HB2	3:F:396:VAL:O	2.03	0.58
1:r:24:LEU:CD1	1:r:41:VAL:HG22	2.33	0.58
3:B:628:GLN:HA	3:B:631:THR:HG22	1.85	0.58
3:D:73:PHE:HB2	3:D:119:VAL:HG21	1.85	0.58
3:D:401:GLY:O	3:D:402:ALA:HB3	2.02	0.58
3:D:484:ILE:HG22	3:D:485:VAL:N	2.18	0.58
3:H:57:THR:HG22	3:H:58:ILE:N	2.15	0.58
3:H:205:ASN:HA	3:H:208:TYR:HB3	1.85	0.58
3:H:350:THR:O	3:H:350:THR:HG23	2.03	0.58
3:H:369:SER:HB3	3:H:445:ILE:HA	1.86	0.58
3:J:341:MET:CB	3:J:437:ILE:HG23	2.29	0.58
3:J:484:ILE:HG22	3:J:485:VAL:N	2.19	0.58
3:J:541:SER:HB3	3:L:169:MET:HG2	1.86	0.58
2:K:13:ILE:CG2	2:K:14:THR:N	2.65	0.58
3:L:73:PHE:HB2	3:L:119:VAL:HG21	1.85	0.58
3:L:199:LEU:HD12	3:L:203:TYR:HD2	1.68	0.58
3:P:199:LEU:HD12	3:P:203:TYR:HD2	1.68	0.58
3:P:369:SER:OG	3:P:443:PHE:HD1	1.85	0.58
3:P:380:VAL:HG13	3:P:381:PRO:HD2	1.86	0.58
3:T:49:ARG:NH2	3:T:64:VAL:HG11	2.18	0.58
3:T:287:PRO:HB3	3:T:509:LEU:CD2	2.32	0.58
5:W:81:TYR:CA	5:W:180:LEU:HD23	2.33	0.58
5:W:439:PHE:HD2	5:W:684:ARG:HH12	1.49	0.58
5:W:563:LEU:N	5:W:563:LEU:HD12	2.18	0.58

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
5:W:891:ASP:O	5:W:927:THR:N	2.33	0.58
6:X:259:LEU:HD23	6:X:911:THR:CG2	2.34	0.58
6:X:527:ALA:HB3	6:X:817:ALA:CB	2.34	0.58
6:X:831:LEU:HD12	6:X:832:ILE:N	2.18	0.58
6:X:936:ALA:O	6:X:937:ALA:HB2	2.03	0.58
6:X:940:GLN:HE21	6:X:944:ASN:HB3	1.69	0.58
6:Y:235:LEU:HB3	6:Y:236:PRO:HD2	1.85	0.58
3:B:540:VAL:HG23	3:B:569:ALA:HB3	1.83	0.58
3:D:47:LEU:C	3:D:48:TRP:CD1	2.82	0.58
3:D:57:THR:O	3:D:60:SER:N	2.36	0.58
3:D:205:ASN:HA	3:D:208:TYR:HB3	1.85	0.58
3:D:449:THR:CG2	3:D:450:SER:N	2.66	0.58
3:F:469:VAL:HG23	3:H:575:MET:HG3	1.84	0.58
3:H:380:VAL:HG13	3:H:381:PRO:HD2	1.85	0.58
3:J:294:TYR:HH	3:J:502:LEU:HA	1.68	0.58
3:J:449:THR:CG2	3:J:450:SER:N	2.66	0.58
3:L:541:SER:HB3	3:N:169:MET:HG2	1.86	0.58
3:P:469:VAL:HG23	3:R:575:MET:HG3	1.84	0.58
3:R:449:THR:CG2	3:R:450:SER:N	2.66	0.58
3:R:541:SER:HB3	3:T:169:MET:HG2	1.86	0.58
7:S:101:MYR:H71	3:T:200:CYS:SG	2.44	0.58
4:U:189:ALA:HA	4:U:208:ILE:HG12	1.85	0.58
4:V:211:LEU:O	4:V:214:ILE:HG22	2.04	0.58
5:W:438:VAL:HG22	5:W:439:PHE:N	2.18	0.58
5:W:918:ILE:HG23	5:W:929:MET:HE1	1.84	0.58
6:X:213:ALA:O	6:X:215:LEU:HD22	2.03	0.58
6:X:300:MET:O	6:X:303:GLN:HB3	2.03	0.58
6:X:742:MET:HG2	6:X:827:THR:OG1	2.04	0.58
6:Y:194:ILE:HD13	6:Y:1201:TYR:CE2	2.39	0.58
6:Y:200:LEU:HD21	6:Y:343:ALA:HB2	1.85	0.58
6:Y:200:LEU:HG	6:Y:339:LEU:HD22	1.86	0.58
6:Y:633:PRO:N	6:Y:640:ARG:HD3	2.19	0.58
1:d:60:ALA:H	3:D:398:SER:CA	2.15	0.58
1:h:45:ARG:O	1:h:59:HIS:C	2.47	0.58
1:h:72:SER:O	1:h:72:SER:OG	2.17	0.58
3:B:74:VAL:O	3:B:74:VAL:HG12	2.04	0.58
3:B:449:THR:CG2	3:B:450:SER:H	2.16	0.58
7:C:101:MYR:H142	7:C:101:MYR:C10	2.10	0.58
3:D:78:MET:HE1	3:H:238:VAL:CG2	2.07	0.58
3:D:356:THR:HG23	3:H:412:ARG:NH2	2.18	0.58
3:F:252:LEU:O	3:F:256:SER:HB3	2.03	0.58

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
7:G:101:MYR:C2	3:H:191:LYS:HE2	2.27	0.58
3:J:51:VAL:HG11	3:J:62:ALA:HB2	1.85	0.58
3:J:92:TRP:CE2	3:J:156:VAL:HG23	2.38	0.58
3:L:215:TYR:C	3:L:217:ALA:H	2.09	0.58
3:L:350:THR:O	3:L:350:THR:HG23	2.03	0.58
3:L:372:ALA:HB1	3:L:399:ALA:HA	1.83	0.58
3:L:380:VAL:HG13	3:L:381:PRO:HD2	1.86	0.58
3:N:189:PHE:CZ	3:N:193:ILE:CD1	2.86	0.58
3:P:49:ARG:NH2	3:P:64:VAL:HG11	2.18	0.58
3:P:78:MET:HE1	3:T:238:VAL:CG2	2.07	0.58
2:Q:28:SER:HA	3:R:216:PRO:HD2	1.85	0.58
6:X:189:ASN:ND2	6:X:915:LEU:HA	2.13	0.58
6:X:368:ILE:HG23	6:X:369:GLY:N	2.19	0.58
6:Y:499:THR:OG1	6:Y:500:ILE:N	2.37	0.58
6:Y:740:LYS:NZ	6:Y:742:MET:HE2	2.18	0.58
6:Y:1156:THR:HG21	6:Y:1161:ASN:HB2	1.86	0.58
1:h:58:PRO:CG	3:H:423:GLN:HE21	2.16	0.58
1:j:60:ALA:N	3:J:398:SER:HA	2.19	0.58
1:n:45:ARG:HD2	3:N:400:ALA:CA	2.33	0.58
3:B:380:VAL:HG13	3:B:381:PRO:HD2	1.85	0.58
3:D:212:ILE:HD11	3:D:220:PRO:HB3	1.85	0.58
3:D:274:ILE:HD11	3:D:514:LEU:HD21	1.73	0.58
7:E:101:MYR:H142	7:E:101:MYR:C10	2.10	0.58
3:F:205:ASN:HA	3:F:208:TYR:HB3	1.85	0.58
3:F:212:ILE:HD11	3:F:220:PRO:HB3	1.85	0.58
3:F:372:ALA:HB3	3:F:444:LEU:CD1	2.33	0.58
3:F:481:VAL:HG12	3:F:482:ASP:N	2.17	0.58
7:G:101:MYR:C2	3:H:191:LYS:HZ3	2.04	0.58
3:H:212:ILE:HD11	3:H:220:PRO:HB3	1.85	0.58
3:J:199:LEU:HD12	3:J:203:TYR:CD2	2.38	0.58
3:J:380:VAL:HG13	3:J:381:PRO:HD2	1.85	0.58
3:L:189:PHE:CZ	3:L:193:ILE:CD1	2.86	0.58
3:L:484:ILE:HG22	3:L:485:VAL:N	2.19	0.58
3:P:73:PHE:HB2	3:P:119:VAL:HG21	1.85	0.58
3:P:92:TRP:CE2	3:P:156:VAL:HG23	2.38	0.58
3:P:449:THR:CG2	3:P:450:SER:H	2.16	0.58
4:U:223:LEU:HD21	4:U:411:PHE:CD2	2.38	0.58
5:W:27:PHE:CE1	5:W:99:LEU:HD21	2.39	0.58
5:W:929:MET:CE	5:W:981:LEU:HD11	2.33	0.58
6:X:801:ARG:NH1	6:X:801:ARG:CB	2.60	0.58
6:Y:191:PRO:HG2	6:Y:193:LEU:HG	1.85	0.58

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:A:28:SER:HA	3:B:216:PRO:HD2	1.85	0.58
3:B:382:ILE:HA	3:B:385:VAL:CG1	2.34	0.58
2:C:28:SER:HA	3:D:216:PRO:HD2	1.85	0.58
3:D:561:ASP:O	3:D:562:TYR:HB2	2.04	0.58
3:F:380:VAL:HG13	3:F:381:PRO:HD2	1.85	0.58
3:H:147:ARG:HB2	3:H:147:ARG:CZ	2.33	0.58
3:J:46:LYS:HA	3:J:72:SER:HA	1.84	0.58
3:J:169:MET:HG2	3:N:541:SER:HB3	1.86	0.58
3:L:369:SER:HB3	3:L:445:ILE:HA	1.86	0.58
3:P:356:THR:HG23	3:T:412:ARG:NH2	2.18	0.58
3:P:372:ALA:HB2	3:P:399:ALA:HA	1.85	0.58
3:R:287:PRO:HB3	3:R:509:LEU:CD2	2.32	0.58
3:R:380:VAL:HG13	3:R:381:PRO:HD2	1.85	0.58
3:T:189:PHE:CZ	3:T:193:ILE:CD1	2.86	0.58
3:T:350:THR:HG23	3:T:350:THR:O	2.03	0.58
3:T:370:VAL:O	3:T:370:VAL:HG23	2.02	0.58
3:T:380:VAL:HG13	3:T:381:PRO:HD2	1.85	0.58
4:U:16:GLN:HG3	4:U:17:PRO:HD2	1.85	0.58
4:V:312:VAL:HG11	4:V:315:LEU:HD12	1.84	0.58
5:W:248:LEU:HD23	5:W:254:LEU:CD2	2.31	0.58
5:W:1208:CYS:HB3	5:W:1216:GLY:HA2	1.86	0.58
6:Y:215:LEU:N	6:Y:215:LEU:CD2	2.66	0.58
6:Y:294:ARG:HE	6:Y:890:GLN:HB3	1.68	0.58
6:Y:299:LEU:HD21	6:Y:904:ALA:HB3	1.85	0.58
6:Y:328:ARG:NH1	6:Y:366:ASP:O	2.37	0.58
6:Y:579:PRO:HB2	6:Y:779:VAL:HG23	1.86	0.58
6:Y:874:CYS:HB3	6:Y:893:VAL:HG21	1.86	0.58
1:h:24:LEU:CD1	1:h:80:ARG:CD	2.80	0.58
3:B:205:ASN:HA	3:B:208:TYR:HB3	1.85	0.58
3:D:169:MET:HG2	3:H:541:SER:HB3	1.85	0.58
3:H:175:SER:OG	3:H:627:ARG:HD3	2.04	0.58
3:H:272:ASP:O	3:H:273:LEU:HD22	2.04	0.58
3:H:341:MET:CB	3:H:437:ILE:HG23	2.29	0.58
3:H:369:SER:HB3	3:H:446:PRO:HD3	1.86	0.58
3:H:562:TYR:HB2	3:H:563:PRO:HD3	1.85	0.58
3:J:175:SER:OG	3:J:627:ARG:HD3	2.04	0.58
3:J:350:THR:O	3:J:350:THR:HG23	2.03	0.58
3:J:382:ILE:HA	3:J:385:VAL:CG1	2.34	0.58
3:J:449:THR:CG2	3:J:450:SER:H	2.16	0.58
3:L:294:TYR:OH	3:L:502:LEU:HD23	2.03	0.58
3:N:294:TYR:OH	3:N:502:LEU:HD23	2.03	0.58

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:N:369:SER:HB3	3:N:445:ILE:HA	1.86	0.58
2:O:28:SER:HA	3:P:216:PRO:HD2	1.85	0.58
3:P:74:VAL:HG12	3:P:74:VAL:O	2.04	0.58
3:P:350:THR:HG23	3:P:350:THR:O	2.03	0.58
3:P:423:GLN:HG3	3:P:423:GLN:O	2.04	0.58
3:P:481:VAL:HG12	3:P:482:ASP:N	2.17	0.58
3:R:73:PHE:HB2	3:R:119:VAL:HG21	1.85	0.58
3:R:92:TRP:CE2	3:R:156:VAL:HG23	2.38	0.58
3:T:449:THR:CG2	3:T:450:SER:N	2.66	0.58
3:T:481:VAL:HG12	3:T:482:ASP:N	2.17	0.58
5:W:146:SER:N	5:W:147:PRO:HD2	2.19	0.58
5:W:850:LEU:C	5:W:850:LEU:HD23	2.28	0.58
6:X:424:ALA:HA	6:X:757:VAL:HG22	1.86	0.58
6:X:608:GLN:HG2	6:X:608:GLN:O	2.03	0.58
6:X:899:ALA:HA	6:X:929:PRO:HG2	1.85	0.58
6:X:983:TYR:CE2	6:X:1002:PRO:HB2	2.39	0.58
6:Y:376:LEU:CD1	6:Y:438:PRO:HB3	2.27	0.58
6:Y:627:ASN:ND2	6:Y:629:GLN:HB2	2.18	0.58
6:Y:640:ARG:HG3	6:Y:640:ARG:NH1	2.19	0.58
6:Y:740:LYS:HZ1	6:Y:742:MET:HE2	1.69	0.58
3:B:272:ASP:O	3:B:273:LEU:HD22	2.04	0.57
3:B:369:SER:HB3	3:B:446:PRO:HD3	1.86	0.57
3:B:372:ALA:HB2	3:B:399:ALA:HA	1.85	0.57
3:D:175:SER:OG	3:D:627:ARG:HD3	2.04	0.57
3:D:369:SER:HB3	3:D:446:PRO:HD3	1.86	0.57
3:F:347:SER:HA	3:F:469:VAL:HG12	1.86	0.57
3:H:525:ILE:HG23	3:H:526:PRO:HD2	1.86	0.57
3:H:561:ASP:O	3:H:562:TYR:HB2	2.04	0.57
3:J:74:VAL:HG12	3:J:74:VAL:O	2.04	0.57
3:L:139:THR:HA	3:L:142:ILE:HG23	1.86	0.57
3:L:205:ASN:HA	3:L:208:TYR:HB3	1.85	0.57
3:L:341:MET:CB	3:L:437:ILE:HG23	2.29	0.57
3:L:481:VAL:HG12	3:L:482:ASP:N	2.17	0.57
3:N:74:VAL:HG12	3:N:74:VAL:O	2.04	0.57
3:N:175:SER:OG	3:N:627:ARG:HD3	2.04	0.57
3:N:252:LEU:O	3:N:256:SER:HB3	2.03	0.57
3:N:322:TYR:CE2	3:N:327:TYR:HE2	2.21	0.57
3:N:341:MET:CB	3:N:437:ILE:HG23	2.29	0.57
3:N:369:SER:HB3	3:N:446:PRO:HD3	1.86	0.57
3:N:521:VAL:HG23	3:N:577:VAL:HG22	1.85	0.57
3:N:561:ASP:O	3:N:562:TYR:HB2	2.04	0.57

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:P:58:ILE:HG12	3:P:127:LEU:CG	2.34	0.57
3:P:272:ASP:O	3:P:273:LEU:HD22	2.04	0.57
3:R:341:MET:CB	3:R:437:ILE:HG23	2.29	0.57
3:R:418:ASN:HD21	3:T:337:ARG:NH2	1.93	0.57
3:R:469:VAL:HG23	3:T:575:MET:HG3	1.85	0.57
3:T:272:ASP:O	3:T:273:LEU:HD22	2.04	0.57
3:T:294:TYR:OH	3:T:502:LEU:HD23	2.03	0.57
3:T:369:SER:HB3	3:T:445:ILE:HA	1.86	0.57
3:T:551:GLN:OE1	3:T:551:GLN:HA	2.04	0.57
4:U:399:GLN:O	4:U:401:ALA:N	2.37	0.57
5:W:396:ARG:HA	5:W:793:THR:OG1	2.04	0.57
5:W:937:LEU:HA	5:W:970:PRO:HB3	1.86	0.57
5:W:997:TRP:C	5:W:999:PRO:HD2	2.29	0.57
6:X:597:GLU:CG	6:X:636:ALA:HA	2.34	0.57
6:X:692:THR:HG23	6:X:745:PHE:HD1	1.69	0.57
1:l:71:CYS:SG	1:l:72:SER:N	2.77	0.57
1:t:21:ARG:O	3:T:323:SER:O	2.22	0.57
1:t:44:GLY:C	3:T:399:ALA:HB3	2.29	0.57
1:t:45:ARG:HD2	3:T:400:ALA:CA	2.34	0.57
1:t:73:ARG:HG3	1:t:75:PRO:HD3	1.86	0.57
3:B:491:ALA:HB2	3:B:592:THR:CB	2.32	0.57
3:D:107:VAL:HG21	3:F:128:ASP:CG	2.30	0.57
3:D:347:SER:HA	3:D:469:VAL:HG12	1.86	0.57
3:F:272:ASP:O	3:F:273:LEU:HD22	2.04	0.57
3:F:628:GLN:HA	3:F:631:THR:HG22	1.85	0.57
3:H:152:ARG:HA	3:H:152:ARG:HH11	1.69	0.57
3:H:491:ALA:HB2	3:H:592:THR:CB	2.32	0.57
3:J:128:ASP:CG	3:N:107:VAL:HG21	2.29	0.57
3:J:212:ILE:HD11	3:J:220:PRO:HB3	1.85	0.57
3:L:280:LEU:CD1	3:L:486:PRO:HG2	2.35	0.57
3:L:521:VAL:HG23	3:L:577:VAL:HG22	1.85	0.57
7:M:101:MYR:H71	3:N:200:CYS:SG	2.44	0.57
3:P:128:ASP:CG	3:T:107:VAL:HG21	2.30	0.57
3:P:260:SER:HB3	2:Q:33:ALA:HB2	1.85	0.57
3:R:369:SER:HB3	3:R:446:PRO:HD3	1.86	0.57
3:T:205:ASN:HA	3:T:208:TYR:HB3	1.85	0.57
3:T:280:LEU:CD1	3:T:486:PRO:HG2	2.34	0.57
3:T:431:ILE:HD13	3:T:437:ILE:HG23	1.82	0.57
4:U:374:LEU:HD21	4:U:388:PRO:HA	1.86	0.57
5:W:76:GLU:O	5:W:77:ASN:HB3	2.05	0.57
5:W:219:ASN:ND2	6:X:674:ARG:HH12	2.02	0.57

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
5:W:321:GLU:HG3	5:W:322:SER:N	2.18	0.57
5:W:559:VAL:O	5:W:559:VAL:HG12	2.04	0.57
5:W:864:MET:HE2	5:W:874:TYR:HB2	1.85	0.57
6:X:425:ASN:O	6:X:427:LEU:N	2.34	0.57
6:X:488:LEU:CD1	6:X:821:ALA:HA	2.34	0.57
6:X:597:GLU:CD	6:X:636:ALA:HA	2.28	0.57
6:X:782:ASN:CB	6:X:784:LEU:HD11	2.33	0.57
6:X:1017:PHE:CD1	6:Y:1210:VAL:O	2.56	0.57
6:X:1054:VAL:HG21	6:Y:1212:SER:CB	2.33	0.57
6:Y:192:GLU:HB2	6:Y:914:ASP:OD1	2.05	0.57
6:Y:997:ALA:CB	6:Y:1141:ASN:HB2	2.34	0.57
1:f:45:ARG:CD	3:F:401:GLY:N	2.66	0.57
1:d:15:ARG:NH2	3:D:402:ALA:HB3	2.18	0.57
1:h:23:THR:O	1:h:23:THR:HG23	2.03	0.57
3:B:521:VAL:HG23	3:B:577:VAL:HG22	1.85	0.57
3:D:272:ASP:O	3:D:273:LEU:HD22	2.04	0.57
3:D:628:GLN:HA	3:D:631:THR:HG22	1.85	0.57
2:E:13:ILE:HG23	2:E:14:THR:N	2.20	0.57
3:F:382:ILE:HA	3:F:385:VAL:CG1	2.34	0.57
3:F:541:SER:HB3	3:H:169:MET:HG2	1.86	0.57
3:F:562:TYR:HB2	3:F:563:PRO:HD3	1.85	0.57
3:H:517:LEU:HD22	3:H:517:LEU:O	2.05	0.57
3:J:107:VAL:HG21	3:L:128:ASP:CG	2.30	0.57
3:L:418:ASN:HD21	3:N:337:ARG:NH2	1.93	0.57
3:L:469:VAL:HG23	3:N:575:MET:HG3	1.84	0.57
3:R:449:THR:CG2	3:R:450:SER:H	2.16	0.57
3:T:147:ARG:HH11	3:T:147:ARG:HB3	1.70	0.57
3:T:175:SER:OG	3:T:627:ARG:HD3	2.04	0.57
3:T:212:ILE:HD11	3:T:220:PRO:HB3	1.85	0.57
4:U:182:ILE:HB	4:U:246:ALA:HB3	1.86	0.57
4:V:94:ASP:HB2	4:V:95:PRO:HD2	1.86	0.57
5:W:221:ALA:HA	5:W:293:THR:HG21	1.85	0.57
5:W:1107:ILE:HG23	5:W:1126:THR:HG22	1.86	0.57
6:X:1054:VAL:CG1	6:Y:835:HIS:CE1	2.87	0.57
6:X:1078:LEU:H	6:X:1078:LEU:CD2	2.15	0.57
6:X:1150:ASN:HB3	6:X:1153:SER:HB3	1.86	0.57
6:Y:545:LEU:HD13	6:Y:569:LEU:HD11	1.86	0.57
1:f:15:ARG:NH2	3:F:402:ALA:HB3	2.19	0.57
1:n:21:ARG:NH1	3:N:327:TYR:OH	2.38	0.57
1:n:22:LEU:HB2	1:n:83:VAL:HG21	1.86	0.57
1:t:59:HIS:HB2	3:T:398:SER:N	2.18	0.57

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:D:369:SER:HB3	3:D:445:ILE:HA	1.86	0.57
3:D:372:ALA:HB2	3:D:399:ALA:HA	1.85	0.57
3:D:562:TYR:HB2	3:D:563:PRO:HD3	1.85	0.57
3:F:372:ALA:HB2	3:F:399:ALA:HA	1.85	0.57
2:I:28:SER:HA	3:J:216:PRO:HD2	1.85	0.57
3:J:78:MET:HE1	3:N:238:VAL:CG2	2.07	0.57
3:J:272:ASP:O	3:J:273:LEU:HD22	2.04	0.57
3:J:372:ALA:HB2	3:J:399:ALA:HA	1.85	0.57
3:L:87:PHE:HD1	3:N:121:ALA:HB2	1.61	0.57
3:L:347:SER:HA	3:L:469:VAL:HG12	1.86	0.57
3:L:423:GLN:O	3:L:423:GLN:HG3	2.04	0.57
2:M:13:ILE:HG23	2:M:14:THR:N	2.20	0.57
3:N:205:ASN:HA	3:N:208:TYR:HB3	1.85	0.57
3:N:380:VAL:HG13	3:N:381:PRO:HD2	1.85	0.57
3:N:517:LEU:HD22	3:N:517:LEU:O	2.05	0.57
3:P:212:ILE:HD11	3:P:220:PRO:HB3	1.85	0.57
3:P:294:TYR:OH	3:P:502:LEU:HD23	2.03	0.57
3:P:369:SER:HB3	3:P:446:PRO:HD3	1.86	0.57
3:P:484:ILE:HG22	3:P:485:VAL:N	2.19	0.57
3:R:350:THR:O	3:R:350:THR:HG23	2.03	0.57
3:T:561:ASP:O	3:T:562:TYR:HB2	2.04	0.57
4:U:138:ILE:O	4:U:142:VAL:HG23	2.03	0.57
5:W:313:LEU:HD23	5:W:347:GLN:HG3	1.85	0.57
5:W:459:ASP:HB2	5:W:625:ARG:HD3	1.87	0.57
5:W:1036:ARG:NH2	5:W:1053:THR:HG21	2.19	0.57
6:X:680:PRO:HG3	6:X:696:ALA:HB2	1.86	0.57
6:Y:541:VAL:N	6:Y:808:ALA:HB1	2.19	0.57
6:Y:697:MET:HE2	6:Y:744:PRO:HA	1.84	0.57
1:d:6:ILE:CD1	3:F:586:VAL:HG21	2.34	0.57
1:p:15:ARG:NH1	3:P:406:ASN:OD1	2.38	0.57
3:B:368:THR:CA	3:B:403:SER:HB2	2.35	0.57
3:B:369:SER:HB3	3:B:445:ILE:HA	1.86	0.57
3:D:341:MET:CB	3:D:437:ILE:HG23	2.29	0.57
3:D:368:THR:CA	3:D:403:SER:HB2	2.35	0.57
3:F:369:SER:HB3	3:F:446:PRO:HD3	1.86	0.57
3:F:431:ILE:HD13	3:F:437:ILE:HG23	1.82	0.57
3:H:347:SER:HA	3:H:469:VAL:HG12	1.86	0.57
3:J:94:PRO:HG2	3:J:210:VAL:HG22	1.87	0.57
3:J:205:ASN:HA	3:J:208:TYR:HB3	1.85	0.57
3:J:354:SER:CB	3:J:419:MET:HG2	2.35	0.57
3:J:423:GLN:O	3:J:423:GLN:HG3	2.04	0.57

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:J:525:ILE:HG23	3:J:526:PRO:HD2	1.86	0.57
3:L:107:VAL:HG21	3:N:128:ASP:CG	2.30	0.57
3:N:347:SER:HA	3:N:469:VAL:HG12	1.86	0.57
3:N:354:SER:CB	3:N:419:MET:HG2	2.35	0.57
3:N:372:ALA:HB2	3:N:399:ALA:HA	1.85	0.57
3:N:551:GLN:HA	3:N:551:GLN:OE1	2.04	0.57
2:O:33:ALA:HB2	3:T:260:SER:HB3	1.85	0.57
3:P:449:THR:CG2	3:P:450:SER:N	2.66	0.57
3:P:561:ASP:O	3:P:562:TYR:HB2	2.04	0.57
3:P:628:GLN:HA	3:P:631:THR:HG22	1.85	0.57
3:R:382:ILE:HA	3:R:385:VAL:CG1	2.34	0.57
3:R:484:ILE:HG22	3:R:485:VAL:N	2.19	0.57
4:U:241:LEU:CA	6:X:1179:VAL:HG11	2.33	0.57
5:W:570:VAL:HG12	5:W:572:THR:H	1.70	0.57
5:W:627:PHE:C	5:W:627:PHE:CD1	2.82	0.57
5:W:831:LEU:HD11	5:W:894:THR:HG22	1.85	0.57
5:W:1138:MET:SD	5:W:1139:PRO:HD2	2.45	0.57
6:X:249:TRP:CZ3	6:X:342:LEU:HD23	2.38	0.57
6:X:271:VAL:CG1	6:X:1114:PRO:HD3	2.33	0.57
6:X:417:VAL:CG1	6:X:418:GLN:N	2.67	0.57
6:X:427:LEU:HD11	6:X:841:PHE:HD2	1.67	0.57
6:X:455:GLN:OE1	6:X:455:GLN:CA	2.50	0.57
6:X:971:GLU:HG2	6:X:974:LEU:HD13	1.86	0.57
6:Y:747:ILE:O	6:Y:747:ILE:HG23	2.04	0.57
6:Y:1033:ASP:HB3	6:Y:1037:VAL:O	2.03	0.57
1:l:24:LEU:HD13	1:l:32:THR:O	1.94	0.57
1:f:45:ARG:HG2	3:F:399:ALA:O	2.04	0.57
1:t:47:THR:CG2	1:t:59:HIS:O	2.52	0.57
3:B:94:PRO:HG2	3:B:210:VAL:HG22	1.87	0.57
3:D:51:VAL:N	3:D:60:SER:O	2.36	0.57
3:D:92:TRP:CE2	3:D:156:VAL:HG23	2.38	0.57
3:D:147:ARG:HH11	3:D:147:ARG:HB3	1.70	0.57
3:D:287:PRO:HB3	3:D:509:LEU:CD2	2.32	0.57
3:D:354:SER:CB	3:D:419:MET:HG2	2.35	0.57
3:D:382:ILE:HA	3:D:385:VAL:CG1	2.34	0.57
3:F:175:SER:OG	3:F:627:ARG:HD3	2.04	0.57
3:F:280:LEU:CD1	3:F:486:PRO:HG2	2.35	0.57
3:F:491:ALA:HB2	3:F:592:THR:CB	2.32	0.57
3:H:423:GLN:HG3	3:H:423:GLN:O	2.04	0.57
3:J:369:SER:HB3	3:J:446:PRO:HD3	1.86	0.57
3:J:450:SER:O	3:J:453:ASN:HB2	2.05	0.57

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:J:481:VAL:HG12	3:J:482:ASP:N	2.17	0.57
3:L:372:ALA:HB3	3:L:444:LEU:CD1	2.33	0.57
3:L:517:LEU:HD22	3:L:517:LEU:O	2.05	0.57
3:N:450:SER:O	3:N:453:ASN:HB2	2.05	0.57
3:P:356:THR:CG2	3:T:412:ARG:NH2	2.63	0.57
3:P:427:GLU:HB2	3:P:430:THR:HG23	1.87	0.57
3:P:551:GLN:OE1	3:P:551:GLN:HA	2.04	0.57
3:P:562:TYR:HB2	3:P:563:PRO:HD3	1.85	0.57
3:R:49:ARG:NH2	3:R:64:VAL:HG11	2.19	0.57
3:T:152:ARG:HA	3:T:152:ARG:HH11	1.68	0.57
3:T:517:LEU:O	3:T:517:LEU:HD22	2.05	0.57
4:U:128:TYR:OH	4:U:139:ASN:HB2	2.05	0.57
5:W:59:ILE:HB	5:W:178:ILE:HD11	1.86	0.57
5:W:267:MET:HE2	5:W:336:PHE:CG	2.40	0.57
6:X:983:TYR:HE1	6:X:1075:LEU:HG	1.70	0.57
6:X:1169:TRP:HB3	6:X:1172:LEU:HB3	1.87	0.57
6:Y:549:ALA:CB	6:Y:815:PRO:HG3	2.33	0.57
1:d:24:LEU:CD1	1:d:41:VAL:HG22	2.35	0.57
3:D:280:LEU:CD1	3:D:486:PRO:HG2	2.35	0.57
3:D:481:VAL:HG12	3:D:482:ASP:N	2.17	0.57
3:F:107:VAL:HG21	3:H:128:ASP:CG	2.30	0.57
3:F:368:THR:CA	3:F:403:SER:HB2	2.35	0.57
3:F:412:ARG:NH2	3:H:356:THR:CG2	2.63	0.57
2:G:13:ILE:HG23	2:G:14:THR:N	2.20	0.57
3:J:199:LEU:HD12	3:J:203:TYR:HD2	1.68	0.57
3:J:628:GLN:HA	3:J:631:THR:HG22	1.85	0.57
3:L:368:THR:CA	3:L:403:SER:HB2	2.35	0.57
3:L:382:ILE:HA	3:L:385:VAL:CG1	2.34	0.57
3:N:94:PRO:HG2	3:N:210:VAL:HG22	1.87	0.57
3:N:272:ASP:O	3:N:273:LEU:HD22	2.04	0.57
3:P:152:ARG:HA	3:P:152:ARG:HH11	1.69	0.57
3:P:271:LEU:HG	3:P:517:LEU:CB	2.30	0.57
3:R:427:GLU:HB2	3:R:430:THR:HG23	1.87	0.57
2:S:13:ILE:HG23	2:S:14:THR:N	2.20	0.57
4:U:327:ASP:H	4:U:399:GLN:CG	2.16	0.57
5:W:245:GLY:O	5:W:257:GLN:HB2	2.05	0.57
5:W:369:ILE:O	5:W:370:GLY:O	2.23	0.57
5:W:845:LEU:N	5:W:845:LEU:CD1	2.67	0.57
5:W:942:SER:HB3	5:W:949:ILE:HG13	1.87	0.57
6:X:229:TYR:O	6:X:230:GLN:C	2.47	0.57
6:X:242:ARG:NH2	6:X:1085:LEU:HD13	2.15	0.57

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
6:X:1075:LEU:HD23	6:X:1075:LEU:C	2.29	0.57
1:b:59:HIS:C	3:B:398:SER:HA	2.29	0.57
1:j:15:ARG:HH22	3:J:402:ALA:HB3	1.69	0.57
3:B:517:LEU:HD22	3:B:517:LEU:O	2.05	0.57
3:D:128:ASP:CG	3:H:107:VAL:HG21	2.30	0.57
3:H:94:PRO:HG2	3:H:210:VAL:HG22	1.86	0.57
3:H:443:PHE:HB2	3:H:457:PRO:CB	2.24	0.57
3:J:562:TYR:HB2	3:J:563:PRO:HD3	1.85	0.57
3:L:147:ARG:HH11	3:L:147:ARG:HB3	1.70	0.57
3:L:175:SER:OG	3:L:627:ARG:HD3	2.04	0.57
3:L:551:GLN:OE1	3:L:551:GLN:HA	2.04	0.57
2:M:7:SER:O	2:M:8:VAL:C	2.48	0.57
3:N:372:ALA:HB3	3:N:444:LEU:CD1	2.33	0.57
3:P:107:VAL:HG21	3:R:128:ASP:CG	2.30	0.57
3:P:147:ARG:HH11	3:P:147:ARG:HB3	1.70	0.57
3:P:368:THR:CA	3:P:403:SER:HB2	2.35	0.57
3:R:491:ALA:HB2	3:R:592:THR:CB	2.32	0.57
3:T:354:SER:CB	3:T:419:MET:HG2	2.35	0.57
3:T:372:ALA:HB2	3:T:399:ALA:HA	1.85	0.57
4:U:117:ASN:HD21	4:U:308:TRP:HD1	1.51	0.57
5:W:302:THR:HG22	5:W:304:LEU:H	1.70	0.57
5:W:607:LEU:HD22	5:W:638:HIS:HB3	1.85	0.57
6:X:249:TRP:CE3	6:X:250:ALA:HB2	2.40	0.57
6:X:675:MET:HE3	6:X:766:PRO:CB	2.35	0.57
6:X:861:ARG:O	6:X:865:VAL:HG23	2.04	0.57
6:X:889:ARG:O	6:X:890:GLN:HB2	2.05	0.57
6:X:1064:TRP:NE1	6:X:1068:LYS:HE3	2.19	0.57
6:X:1125:TYR:H	6:X:1125:TYR:HD1	1.47	0.57
6:Y:58:SER:OG	6:Y:61:ASN:HB2	2.04	0.57
6:Y:530:LYS:CD	6:Y:811:LEU:HD21	2.35	0.57
6:Y:535:TRP:CE2	6:Y:678:PRO:HG2	2.40	0.57
6:Y:557:ILE:HG23	6:Y:717:MET:CE	2.26	0.57
6:Y:969:MET:HA	6:Y:972:MET:HE3	1.86	0.57
6:Y:990:ASN:HB2	6:Y:1131:ARG:CB	2.35	0.57
1:b:59:HIS:HB2	3:B:398:SER:HB2	1.86	0.57
2:A:13:ILE:HG23	2:A:14:THR:N	2.20	0.57
3:B:347:SER:HA	3:B:469:VAL:HG12	1.86	0.57
3:B:354:SER:CB	3:B:419:MET:HG2	2.35	0.57
3:D:491:ALA:HB2	3:D:592:THR:CB	2.32	0.57
3:F:354:SER:CB	3:F:419:MET:HG2	2.35	0.57
3:F:423:GLN:HG3	3:F:423:GLN:O	2.04	0.57

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:H:372:ALA:HB2	3:H:399:ALA:HA	1.86	0.57
3:J:289:LEU:CD2	3:L:515:ASN:ND2	2.68	0.57
3:J:561:ASP:O	3:J:562:TYR:HB2	2.04	0.57
3:L:212:ILE:HD11	3:L:220:PRO:HB3	1.85	0.57
3:L:213:HIS:CG	3:L:235:LEU:HD23	2.40	0.57
3:L:272:ASP:O	3:L:273:LEU:HD22	2.04	0.57
3:L:287:PRO:HB3	3:L:509:LEU:CD2	2.32	0.57
3:N:427:GLU:HB2	3:N:430:THR:HG23	1.87	0.57
3:P:94:PRO:HG2	3:P:210:VAL:HG22	1.87	0.57
3:P:169:MET:HG2	3:T:541:SER:HB3	1.86	0.57
3:P:175:SER:OG	3:P:627:ARG:HD3	2.04	0.57
3:P:213:HIS:CG	3:P:235:LEU:HD23	2.40	0.57
3:P:412:ARG:NH2	3:R:356:THR:HG23	2.18	0.57
3:P:431:ILE:HD13	3:P:437:ILE:HG23	1.82	0.57
3:P:456:ASN:HB3	3:T:410:ASN:HD22	1.70	0.57
3:R:272:ASP:O	3:R:273:LEU:HD22	2.04	0.57
3:R:289:LEU:CD2	3:T:515:ASN:ND2	2.68	0.57
3:R:354:SER:CB	3:R:419:MET:HG2	2.35	0.57
3:R:423:GLN:O	3:R:423:GLN:HG3	2.04	0.57
3:R:562:TYR:HB2	3:R:563:PRO:HD3	1.85	0.57
4:U:27:GLN:CD	6:X:880:LEU:HB3	2.30	0.57
6:X:493:LEU:HD13	6:X:497:ILE:HD12	1.87	0.57
6:X:750:MET:HB3	6:X:837:TYR:OH	2.05	0.57
6:X:785:VAL:O	6:X:804:HIS:HB3	2.05	0.57
6:Y:249:TRP:CH2	6:Y:342:LEU:HD13	2.40	0.57
6:Y:252:LEU:HD13	6:Y:336:ARG:HG2	1.86	0.57
6:Y:356:ASN:HB2	6:Y:1151:ALA:HB1	1.87	0.57
6:Y:596:PRO:HB3	6:Y:716:ASN:HD22	1.70	0.57
1:d:3:LEU:HD13	1:d:4:HIS:O	2.05	0.57
1:d:46:TYR:HD1	1:d:58:PRO:HA	1.70	0.57
1:t:58:PRO:HD2	1:t:59:HIS:CE1	2.40	0.57
3:B:175:SER:OG	3:B:627:ARG:HD3	2.04	0.57
3:D:94:PRO:HG2	3:D:210:VAL:HG22	1.87	0.57
3:F:147:ARG:CB	3:F:147:ARG:HH11	2.18	0.57
3:F:152:ARG:HH11	3:F:152:ARG:HA	1.69	0.57
3:H:360:TYR:C	3:H:360:TYR:CD1	2.83	0.57
3:J:50:PRO:O	4:V:401:ALA:CB	2.45	0.57
3:J:354:SER:HB2	3:J:419:MET:CA	2.35	0.57
3:J:617:VAL:HG21	3:L:158:LEU:HD21	1.87	0.57
3:L:271:LEU:HG	3:L:517:LEU:CB	2.30	0.57
3:N:212:ILE:HD11	3:N:220:PRO:HB3	1.85	0.57

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:P:372:ALA:HB3	3:P:444:LEU:CD1	2.33	0.57
3:P:450:SER:O	3:P:453:ASN:HB2	2.05	0.57
3:R:152:ARG:HH11	3:R:152:ARG:HA	1.69	0.57
3:T:94:PRO:HG2	3:T:210:VAL:HG22	1.87	0.57
3:T:347:SER:HA	3:T:469:VAL:HG12	1.86	0.57
4:U:320:GLN:C	4:U:322:VAL:H	2.13	0.57
5:W:27:PHE:O	5:W:31:LEU:HD23	2.05	0.57
5:W:247:VAL:HG11	5:W:250:TYR:HE2	1.70	0.57
6:X:371:LEU:HD11	6:X:388:MET:CG	2.34	0.57
6:X:466:THR:HA	6:X:809:ALA:HB1	1.86	0.57
6:X:538:ASN:ND2	6:X:768:GLN:HG2	2.20	0.57
6:X:1210:VAL:O	6:X:1211:ARG:CB	2.53	0.57
6:Y:69:SER:HB3	6:Y:72:VAL:HG23	1.87	0.57
1:d:25:TYR:HB2	1:d:40:TYR:HB2	1.87	0.56
1:p:44:GLY:O	3:P:399:ALA:CB	2.53	0.56
1:t:62:VAL:O	1:t:63:LYS:C	2.47	0.56
3:D:46:LYS:O	3:D:48:TRP:CD1	2.58	0.56
3:D:412:ARG:HH22	3:F:356:THR:HG23	1.65	0.56
3:D:515:ASN:ND2	3:H:289:LEU:CD2	2.68	0.56
3:D:541:SER:HB3	3:F:169:MET:HG2	1.86	0.56
7:E:101:MYR:H71	3:F:200:CYS:SG	2.45	0.56
3:F:368:THR:HA	3:F:403:SER:HB2	1.87	0.56
3:F:517:LEU:HD22	3:F:517:LEU:O	2.05	0.56
3:F:617:VAL:HG21	3:H:158:LEU:HD21	1.87	0.56
2:G:26:MET:O	3:H:216:PRO:HG3	2.05	0.56
7:G:101:MYR:H71	3:H:200:CYS:SG	2.44	0.56
3:H:521:VAL:HG23	3:H:577:VAL:HG22	1.85	0.56
3:J:158:LEU:HD21	3:N:617:VAL:HG21	1.87	0.56
3:J:287:PRO:HB3	3:J:509:LEU:CD2	2.32	0.56
3:J:369:SER:OG	3:J:443:PHE:HD1	1.85	0.56
3:J:596:MET:O	3:J:597:LEU:HB2	2.05	0.56
3:L:412:ARG:NH2	3:N:356:THR:CG2	2.63	0.56
3:N:368:THR:HA	3:N:403:SER:HB2	1.87	0.56
3:N:525:ILE:HG23	3:N:526:PRO:HD2	1.86	0.56
3:P:515:ASN:ND2	3:T:289:LEU:CD2	2.68	0.56
3:P:541:SER:HB3	3:R:169:MET:HG2	1.86	0.56
3:P:575:MET:HG3	3:T:469:VAL:HG23	1.85	0.56
3:R:147:ARG:HH11	3:R:147:ARG:HB3	1.70	0.56
3:R:410:ASN:HD22	3:T:456:ASN:HB3	1.70	0.56
3:R:517:LEU:HD22	3:R:517:LEU:O	2.05	0.56
3:R:525:ILE:HG23	3:R:526:PRO:HD2	1.86	0.56

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:T:147:ARG:CB	3:T:147:ARG:HH11	2.18	0.56
5:W:62:TYR:HB3	5:W:63:PRO:CD	2.20	0.56
5:W:433:CYS:SG	5:W:479:LEU:HB3	2.45	0.56
5:W:437:GLY:O	5:W:1010:LEU:HB2	2.05	0.56
5:W:1144:LEU:C	5:W:1144:LEU:HD23	2.30	0.56
6:X:370:ARG:HA	6:X:387:CYS:CB	2.35	0.56
6:X:406:LEU:HB3	6:X:441:LEU:HD21	1.87	0.56
1:l:25:TYR:HB2	1:l:40:TYR:HB2	1.87	0.56
1:f:8:GLN:NE2	3:H:302:GLU:CG	2.59	0.56
1:p:25:TYR:HB2	1:p:40:TYR:HB2	1.88	0.56
1:p:44:GLY:O	3:P:399:ALA:HB3	2.05	0.56
1:p:45:ARG:HG2	3:P:399:ALA:O	2.02	0.56
3:B:341:MET:CB	3:B:437:ILE:HG23	2.29	0.56
2:C:26:MET:O	3:D:216:PRO:HG3	2.05	0.56
3:D:358:LEU:HD13	3:D:417:PHE:CZ	2.35	0.56
3:D:368:THR:HA	3:D:403:SER:HB2	1.87	0.56
3:D:596:MET:O	3:D:597:LEU:HB2	2.05	0.56
3:F:74:VAL:HG12	3:F:74:VAL:O	2.04	0.56
3:F:213:HIS:CG	3:F:235:LEU:HD23	2.40	0.56
3:H:596:MET:O	3:H:597:LEU:HB2	2.05	0.56
2:I:13:ILE:HG23	2:I:14:THR:N	2.20	0.56
3:J:410:ASN:HD22	3:L:456:ASN:HB3	1.70	0.56
3:L:238:VAL:HG22	3:N:142:ILE:CD1	2.35	0.56
3:L:617:VAL:HG21	3:N:158:LEU:HD21	1.87	0.56
3:N:147:ARG:CB	3:N:147:ARG:HH11	2.18	0.56
3:P:517:LEU:HD22	3:P:517:LEU:O	2.05	0.56
3:P:525:ILE:HG23	3:P:526:PRO:HD2	1.86	0.56
3:R:280:LEU:CD1	3:R:486:PRO:HG2	2.35	0.56
2:S:26:MET:O	3:T:216:PRO:HG3	2.05	0.56
3:T:450:SER:O	3:T:453:ASN:HB2	2.05	0.56
5:W:566:LEU:HD11	5:W:609:MET:HE2	1.86	0.56
5:W:637:THR:HB	5:W:664:SER:O	2.05	0.56
6:X:274:TYR:CE2	6:X:297:VAL:HG22	2.40	0.56
6:X:895:SER:O	6:X:898:ALA:N	2.38	0.56
1:l:23:THR:CG2	1:l:25:TYR:HE2	2.07	0.56
1:b:23:THR:CG2	1:b:42:THR:HG1	2.10	0.56
1:f:42:THR:HG22	1:f:47:THR:HG22	1.88	0.56
1:f:44:GLY:O	3:F:399:ALA:CA	2.53	0.56
1:n:15:ARG:CZ	3:N:402:ALA:HB3	2.27	0.56
1:n:74:GLN:HE21	1:n:74:GLN:C	2.12	0.56
1:r:42:THR:HG22	1:r:47:THR:HG22	1.88	0.56

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:B:147:ARG:HH11	3:B:147:ARG:HB3	1.70	0.56
3:B:358:LEU:CD1	3:B:417:PHE:CE1	2.62	0.56
3:B:596:MET:O	3:B:597:LEU:HB2	2.05	0.56
3:D:213:HIS:CG	3:D:235:LEU:HD23	2.40	0.56
3:D:427:GLU:HB2	3:D:430:THR:HG23	1.87	0.56
3:D:456:ASN:HB3	3:H:410:ASN:HD22	1.70	0.56
3:D:517:LEU:HD22	3:D:517:LEU:O	2.05	0.56
3:D:551:GLN:OE1	3:D:551:GLN:HA	2.04	0.56
3:F:271:LEU:HG	3:F:517:LEU:CB	2.30	0.56
3:F:360:TYR:CD1	3:F:360:TYR:C	2.83	0.56
3:F:427:GLU:HB2	3:F:430:THR:HG23	1.87	0.56
3:F:521:VAL:HG23	3:F:577:VAL:HG22	1.85	0.56
3:F:525:ILE:HG23	3:F:526:PRO:HD2	1.86	0.56
2:G:9:ASN:CB	3:H:209:ASN:CG	2.79	0.56
3:H:74:VAL:HG12	3:H:74:VAL:O	2.04	0.56
3:H:368:THR:CA	3:H:403:SER:HB2	2.35	0.56
3:H:368:THR:HA	3:H:403:SER:HB2	1.87	0.56
3:J:58:ILE:HG12	3:J:127:LEU:HD23	1.85	0.56
3:J:121:ALA:HB2	3:N:87:PHE:HD1	1.61	0.56
3:J:427:GLU:HB2	3:J:430:THR:HG23	1.87	0.56
3:J:491:ALA:HB2	3:J:592:THR:CB	2.32	0.56
3:L:74:VAL:HG12	3:L:74:VAL:O	2.04	0.56
3:L:94:PRO:HG2	3:L:210:VAL:HG22	1.87	0.56
3:L:268:GLY:O	3:L:269:LYS:C	2.49	0.56
3:L:450:SER:O	3:L:453:ASN:HB2	2.05	0.56
3:L:525:ILE:HG23	3:L:526:PRO:HD2	1.86	0.56
2:M:26:MET:O	3:N:216:PRO:HG3	2.05	0.56
3:N:213:HIS:CG	3:N:235:LEU:HD23	2.40	0.56
3:N:271:LEU:HG	3:N:517:LEU:CB	2.30	0.56
3:N:354:SER:HB2	3:N:419:MET:CA	2.35	0.56
3:N:382:ILE:HA	3:N:385:VAL:CG1	2.34	0.56
3:N:423:GLN:HG3	3:N:423:GLN:O	2.04	0.56
3:P:354:SER:CB	3:P:419:MET:HG2	2.35	0.56
3:P:354:SER:HB2	3:P:419:MET:CA	2.35	0.56
3:P:368:THR:HA	3:P:403:SER:HB2	1.87	0.56
3:R:74:VAL:HG12	3:R:74:VAL:O	2.04	0.56
3:R:94:PRO:HG2	3:R:210:VAL:HG22	1.87	0.56
3:R:450:SER:O	3:R:453:ASN:HB2	2.05	0.56
3:R:561:ASP:O	3:R:562:TYR:HB2	2.04	0.56
2:S:7:SER:O	2:S:8:VAL:C	2.49	0.56
3:T:213:HIS:CG	3:T:235:LEU:HD23	2.40	0.56

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:T:360:TYR:C	3:T:360:TYR:CD1	2.83	0.56
3:T:368:THR:CA	3:T:403:SER:HB2	2.35	0.56
3:T:423:GLN:O	3:T:423:GLN:HG3	2.04	0.56
4:U:269:CYS:SG	4:U:274:TRP:HB3	2.45	0.56
4:U:362:TYR:HB2	6:X:1178:ARG:NH2	2.21	0.56
5:W:465:PHE:CE2	5:W:592:MET:HE2	2.39	0.56
5:W:603:LEU:HD21	5:W:607:LEU:HD12	1.87	0.56
5:W:812:ALA:C	5:W:814:GLY:H	2.12	0.56
6:X:219:LEU:HB2	6:X:222:PHE:HD2	1.71	0.56
6:X:279:SER:HA	6:X:283:ILE:HG22	1.86	0.56
6:X:593:LEU:O	6:X:593:LEU:CG	2.53	0.56
6:X:793:ARG:NH1	6:Y:693:TYR:CD2	2.73	0.56
6:X:935:SER:OG	6:X:949:HIS:HB3	2.05	0.56
6:X:983:TYR:CE1	6:X:1075:LEU:HG	2.40	0.56
6:X:986:GLN:HG3	6:X:1135:TYR:CE1	2.40	0.56
6:X:1086:HIS:CD2	6:X:1086:HIS:C	2.84	0.56
6:X:1095:ASP:HB2	6:Y:107:LYS:O	2.06	0.56
6:Y:776:GLN:O	6:Y:778:ASP:N	2.38	0.56
6:Y:1122:PRO:HA	6:Y:1159:GLY:CA	2.31	0.56
6:Y:1129:THR:HG21	6:Y:1158:PHE:CZ	2.40	0.56
1:n:15:ARG:HH11	3:N:406:ASN:CG	1.92	0.56
1:n:51:PHE:CG	1:n:71:CYS:O	2.58	0.56
1:r:25:TYR:CE2	1:r:42:THR:HG21	2.35	0.56
1:r:45:ARG:CD	3:R:401:GLY:N	2.69	0.56
1:t:15:ARG:NH2	3:T:402:ALA:HB2	2.20	0.56
1:t:21:ARG:HD3	3:T:327:TYR:OH	2.05	0.56
2:A:26:MET:O	3:B:216:PRO:HG3	2.05	0.56
7:A:101:MYR:H142	7:A:101:MYR:C10	2.10	0.56
3:B:369:SER:OG	3:B:443:PHE:HD1	1.85	0.56
3:D:268:GLY:O	3:D:269:LYS:C	2.49	0.56
3:D:287:PRO:O	3:D:471:LEU:HD11	2.05	0.56
3:D:357:ILE:CG2	3:D:359:PRO:HD3	2.35	0.56
3:D:360:TYR:C	3:D:360:TYR:CD1	2.83	0.56
3:D:525:ILE:HG23	3:D:526:PRO:HD2	1.86	0.56
3:F:147:ARG:HH11	3:F:147:ARG:HB3	1.70	0.56
3:F:450:SER:O	3:F:453:ASN:HB2	2.05	0.56
3:H:147:ARG:HH11	3:H:147:ARG:HB3	1.70	0.56
3:H:287:PRO:O	3:H:471:LEU:HD11	2.05	0.56
3:H:450:SER:O	3:H:453:ASN:HB2	2.05	0.56
3:J:147:ARG:HH11	3:J:147:ARG:HB3	1.70	0.56
3:J:287:PRO:O	3:J:471:LEU:HD11	2.05	0.56

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:J:551:GLN:HA	3:J:551:GLN:OE1	2.04	0.56
3:L:561:ASP:O	3:L:562:TYR:HB2	2.04	0.56
3:P:347:SER:HA	3:P:469:VAL:HG12	1.86	0.56
3:P:491:ALA:HB2	3:P:592:THR:CB	2.32	0.56
2:Q:26:MET:O	3:R:216:PRO:HG3	2.05	0.56
3:R:372:ALA:HB3	3:R:444:LEU:CD1	2.33	0.56
5:W:398:LEU:HD13	5:W:790:ARG:HH21	1.71	0.56
6:Y:207:LEU:HD22	6:Y:244:VAL:CG2	2.36	0.56
6:Y:424:ALA:HA	6:Y:757:VAL:HG22	1.88	0.56
6:Y:485:PRO:HD3	6:Y:837:TYR:CE1	2.41	0.56
1:j:46:TYR:CD1	1:j:58:PRO:HA	2.40	0.56
1:t:25:TYR:HB2	1:t:40:TYR:HB2	1.87	0.56
3:B:280:LEU:CD1	3:B:486:PRO:HG2	2.34	0.56
3:B:427:GLU:HB2	3:B:430:THR:HG23	1.87	0.56
3:B:481:VAL:HG12	3:B:482:ASP:N	2.17	0.56
3:D:248:GLU:N	3:D:248:GLU:CD	2.64	0.56
3:D:617:VAL:HG21	3:F:158:LEU:HD21	1.88	0.56
3:F:154:ARG:NH1	3:F:154:ARG:CB	2.68	0.56
3:F:268:GLY:O	3:F:269:LYS:C	2.49	0.56
3:J:260:SER:HB3	2:K:33:ALA:HB2	1.85	0.56
3:J:280:LEU:CD1	3:J:486:PRO:HG2	2.35	0.56
3:J:456:ASN:HB3	3:N:410:ASN:HD22	1.70	0.56
3:N:481:VAL:HG12	3:N:482:ASP:N	2.17	0.56
2:O:9:ASN:CB	3:P:209:ASN:CG	2.79	0.56
2:O:26:MET:O	3:P:216:PRO:HG3	2.05	0.56
3:P:158:LEU:HD21	3:T:617:VAL:HG21	1.87	0.56
3:P:237:GLU:OE1	3:R:46:LYS:NZ	2.38	0.56
3:P:360:TYR:C	3:P:360:TYR:CD1	2.83	0.56
3:R:551:GLN:HA	3:R:551:GLN:OE1	2.03	0.56
2:S:28:SER:CB	3:T:215:TYR:HA	2.36	0.56
3:T:74:VAL:HG12	3:T:74:VAL:O	2.04	0.56
3:T:596:MET:O	3:T:597:LEU:HB2	2.05	0.56
5:W:71:HIS:HE1	5:W:73:ALA:HB3	1.67	0.56
5:W:727:LEU:HD13	5:W:727:LEU:C	2.30	0.56
5:W:732:LEU:O	5:W:732:LEU:HD13	2.06	0.56
5:W:1136:LEU:HD22	5:W:1155:ILE:HG22	1.87	0.56
6:X:874:CYS:O	6:X:890:GLN:HG2	2.06	0.56
6:X:1182:ASN:HD21	6:X:1184:LEU:HD22	1.70	0.56
6:Y:315:SER:OG	6:Y:316:THR:N	2.38	0.56
6:Y:579:PRO:HB2	6:Y:779:VAL:CG2	2.35	0.56
1:b:59:HIS:HE1	1:b:61:ASN:HD21	1.47	0.56

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:n:8:GLN:HE22	3:J:303:ASP:CA	2.17	0.56
1:t:62:VAL:C	1:t:64:THR:H	2.14	0.56
2:A:9:ASN:CB	3:B:209:ASN:CG	2.79	0.56
3:B:168:THR:C	3:B:170:LEU:H	2.14	0.56
2:C:9:ASN:CB	3:D:209:ASN:CG	2.79	0.56
2:C:13:ILE:HG23	2:C:14:THR:N	2.20	0.56
3:D:154:ARG:NH1	3:D:154:ARG:CB	2.68	0.56
3:D:336:THR:HG23	3:D:456:ASN:OD1	2.06	0.56
3:D:354:SER:HB2	3:D:419:MET:CA	2.36	0.56
3:F:289:LEU:CD2	3:H:515:ASN:ND2	2.68	0.56
3:H:147:ARG:CB	3:H:147:ARG:HH11	2.18	0.56
3:H:427:GLU:HB2	3:H:430:THR:HG23	1.87	0.56
3:J:347:SER:HA	3:J:469:VAL:HG12	1.86	0.56
3:J:360:TYR:CD1	3:J:360:TYR:C	2.83	0.56
3:J:517:LEU:HD22	3:J:517:LEU:O	2.05	0.56
3:N:147:ARG:HH11	3:N:147:ARG:HB3	1.70	0.56
3:N:368:THR:CA	3:N:403:SER:HB2	2.35	0.56
3:N:596:MET:O	3:N:597:LEU:HB2	2.05	0.56
3:P:287:PRO:O	3:P:471:LEU:HD11	2.05	0.56
3:R:107:VAL:HG21	3:T:128:ASP:CG	2.30	0.56
3:R:357:ILE:CG2	3:R:359:PRO:HD3	2.35	0.56
3:T:248:GLU:N	3:T:248:GLU:CD	2.64	0.56
3:T:382:ILE:HA	3:T:385:VAL:CG1	2.34	0.56
4:U:10:THR:HG22	4:U:11:TYR:N	2.21	0.56
4:U:390:ASN:O	4:U:392:ALA:N	2.38	0.56
5:W:953:ASP:O	5:W:955:VAL:HG12	2.06	0.56
6:X:208:ASP:CB	6:X:236:PRO:HB2	2.35	0.56
6:X:242:ARG:NH1	6:X:1146:LEU:HA	2.20	0.56
6:X:283:ILE:HD12	6:X:289:ASN:N	2.20	0.56
6:X:450:ASP:OD1	6:X:861:ARG:NH1	2.38	0.56
6:X:599:ILE:HG21	6:X:639:LEU:HD23	1.87	0.56
6:X:725:THR:CG2	6:X:726:ARG:N	2.69	0.56
6:X:757:VAL:HG13	6:X:845:PHE:CE2	2.37	0.56
6:Y:352:THR:HG23	6:Y:352:THR:O	2.04	0.56
6:Y:448:SER:OG	6:Y:665:SER:HB3	2.06	0.56
1:d:42:THR:HG22	1:d:47:THR:HG22	1.88	0.56
1:t:60:ALA:HB3	3:T:396:VAL:O	2.05	0.56
3:B:450:SER:O	3:B:453:ASN:HB2	2.05	0.56
3:D:147:ARG:CB	3:D:147:ARG:HH11	2.18	0.56
3:D:357:ILE:HD11	3:D:414:GLN:CG	2.36	0.56
3:D:610:THR:O	3:D:612:SER:N	2.39	0.56

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:H:280:LEU:CD1	3:H:486:PRO:HG2	2.35	0.56
3:H:357:ILE:CG2	3:H:359:PRO:HD3	2.35	0.56
3:H:551:GLN:OE1	3:H:551:GLN:HA	2.04	0.56
3:J:49:ARG:NH2	3:J:64:VAL:HG11	2.20	0.56
3:J:357:ILE:CG2	3:J:359:PRO:HD3	2.36	0.56
3:J:515:ASN:ND2	3:N:289:LEU:CD2	2.68	0.56
2:K:9:ASN:CB	3:L:209:ASN:CG	2.79	0.56
3:L:168:THR:C	3:L:170:LEU:H	2.14	0.56
3:L:178:SER:OG	2:Q:3:ASN:ND2	2.39	0.56
3:L:369:SER:HB3	3:L:446:PRO:HD3	1.86	0.56
3:L:410:ASN:HD22	3:N:456:ASN:HB3	1.70	0.56
3:L:491:ALA:HB2	3:L:592:THR:CB	2.32	0.56
2:M:28:SER:CB	3:N:215:TYR:HA	2.36	0.56
3:N:280:LEU:CD1	3:N:486:PRO:HG2	2.34	0.56
3:N:336:THR:HG23	3:N:456:ASN:OD1	2.06	0.56
2:O:13:ILE:HG23	2:O:14:THR:N	2.20	0.56
3:P:322:TYR:CD1	3:P:396:VAL:HB	2.41	0.56
3:P:596:MET:O	3:P:597:LEU:HB2	2.05	0.56
2:Q:9:ASN:CB	3:R:209:ASN:CG	2.79	0.56
3:R:168:THR:C	3:R:170:LEU:H	2.14	0.56
3:R:175:SER:OG	3:R:627:ARG:HD3	2.04	0.56
3:R:213:HIS:CG	3:R:235:LEU:HD23	2.40	0.56
3:R:368:THR:CA	3:R:403:SER:HB2	2.35	0.56
3:R:617:VAL:HG21	3:T:158:LEU:HD21	1.87	0.56
3:T:368:THR:HA	3:T:403:SER:HB2	1.87	0.56
3:T:369:SER:HB3	3:T:446:PRO:HD3	1.86	0.56
5:W:47:LEU:HD23	5:W:60:GLN:HB2	1.88	0.56
5:W:800:PHE:CE1	5:W:1005:GLY:HA2	2.40	0.56
5:W:1017:ILE:O	5:W:1021:ARG:HG3	2.05	0.56
6:X:514:ILE:HG13	6:X:515:LEU:N	2.20	0.56
6:X:557:ILE:HB	6:X:592:LEU:CD2	2.35	0.56
6:X:971:GLU:OE1	6:X:1188:ILE:HG13	2.06	0.56
6:X:1014:VAL:HG21	6:X:1047:TYR:CD1	2.41	0.56
6:Y:737:ARG:HG2	6:Y:737:ARG:NH2	2.11	0.56
1:p:58:PRO:C	1:p:59:HIS:CG	2.83	0.56
1:r:24:LEU:HD13	1:r:41:VAL:HG22	1.86	0.56
2:C:28:SER:CB	3:D:215:TYR:HA	2.36	0.56
3:D:121:ALA:HB2	3:H:87:PHE:HD1	1.61	0.56
3:D:158:LEU:HD21	3:H:617:VAL:HG21	1.87	0.56
3:D:168:THR:C	3:D:170:LEU:H	2.14	0.56
3:F:561:ASP:O	3:F:562:TYR:HB2	2.04	0.56

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:H:248:GLU:N	3:H:248:GLU:CD	2.64	0.56
2:I:9:ASN:CB	3:J:209:ASN:CG	2.79	0.56
3:J:238:VAL:CG1	3:L:82:PHE:HZ	2.19	0.56
3:J:248:GLU:CD	3:J:248:GLU:N	2.64	0.56
3:J:336:THR:HG23	3:J:456:ASN:OD1	2.06	0.56
3:J:368:THR:CA	3:J:403:SER:HB2	2.35	0.56
3:L:248:GLU:N	3:L:248:GLU:CD	2.64	0.56
3:N:248:GLU:CD	3:N:248:GLU:N	2.64	0.56
3:P:248:GLU:N	3:P:248:GLU:CD	2.64	0.56
3:P:280:LEU:CD1	3:P:486:PRO:HG2	2.34	0.56
3:P:410:ASN:HD22	3:R:456:ASN:HB3	1.70	0.56
3:R:248:GLU:CD	3:R:248:GLU:N	2.64	0.56
3:R:336:THR:HG23	3:R:456:ASN:OD1	2.06	0.56
3:R:357:ILE:HD11	3:R:414:GLN:CG	2.36	0.56
3:R:368:THR:HA	3:R:403:SER:HB2	1.87	0.56
2:S:9:ASN:CB	3:T:209:ASN:CG	2.79	0.56
3:T:336:THR:HG23	3:T:456:ASN:OD1	2.06	0.56
4:U:241:LEU:HB3	6:X:1179:VAL:HG11	1.87	0.56
4:V:164:ASP:HB2	4:V:165:PRO:HD2	1.87	0.56
5:W:838:PRO:O	5:W:860:CYS:HA	2.06	0.56
5:W:1258:MET:HG2	5:W:1259:SER:N	2.19	0.56
6:X:271:VAL:HG12	6:X:288:PHE:HB3	1.87	0.56
6:X:407:ARG:C	6:X:409:MET:H	2.12	0.56
6:X:514:ILE:HD13	6:X:732:VAL:CG1	2.36	0.56
6:X:640:ARG:O	6:X:644:GLU:HB2	2.06	0.56
6:X:872:ALA:CB	6:X:881:VAL:HG21	2.36	0.56
6:Y:81:ARG:HB2	6:Y:85:ASP:OD2	2.05	0.56
6:Y:662:LEU:HD12	6:Y:662:LEU:N	2.21	0.56
6:Y:1178:ARG:O	6:Y:1179:VAL:C	2.48	0.56
1:b:21:ARG:CD	3:B:327:TYR:HH	2.17	0.56
1:b:25:TYR:HB2	1:b:40:TYR:HB2	1.87	0.56
1:n:44:GLY:CA	3:N:399:ALA:HB3	2.36	0.56
1:t:42:THR:HG22	1:t:47:THR:HG22	1.88	0.56
2:A:7:SER:O	2:A:8:VAL:C	2.49	0.56
2:A:35:ASP:O	2:A:36:LEU:HD23	2.06	0.56
3:B:213:HIS:CG	3:B:235:LEU:HD23	2.40	0.56
3:B:372:ALA:HB3	3:B:444:LEU:CD1	2.33	0.56
3:B:423:GLN:O	3:B:423:GLN:HG3	2.04	0.56
3:D:280:LEU:HD12	3:D:486:PRO:CG	2.36	0.56
2:E:9:ASN:CB	3:F:209:ASN:CG	2.79	0.56
3:F:357:ILE:HD11	3:F:414:GLN:CG	2.36	0.56

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:F:410:ASN:HD22	3:H:456:ASN:HB3	1.70	0.56
3:F:596:MET:O	3:F:597:LEU:HB2	2.05	0.56
3:H:213:HIS:CG	3:H:235:LEU:HD23	2.40	0.56
3:H:354:SER:CB	3:H:419:MET:HG2	2.35	0.56
2:I:35:ASP:O	2:I:36:LEU:HD23	2.06	0.56
3:J:147:ARG:CB	3:J:147:ARG:HH11	2.18	0.56
3:J:213:HIS:CG	3:J:235:LEU:HD23	2.40	0.56
3:L:138:SER:O	3:L:142:ILE:CG2	2.54	0.56
3:L:280:LEU:HD12	3:L:486:PRO:CG	2.36	0.56
3:L:287:PRO:O	3:L:471:LEU:HD11	2.06	0.56
3:L:360:TYR:C	3:L:360:TYR:CD1	2.83	0.56
3:N:168:THR:C	3:N:170:LEU:H	2.14	0.56
3:P:336:THR:HG23	3:P:456:ASN:OD1	2.06	0.56
4:U:266:TYR:CE1	4:U:300:ILE:HG22	2.40	0.56
5:W:265:SER:O	5:W:268:ASP:HB2	2.04	0.56
5:W:1062:SER:HA	5:W:1071:ILE:O	2.05	0.56
5:W:1138:MET:HE3	5:W:1225:ASP:CB	2.29	0.56
6:X:350:LEU:N	6:X:350:LEU:CD1	2.69	0.56
6:X:360:LEU:O	6:X:361:SER:C	2.49	0.56
6:X:982:LEU:HD12	6:X:1115:VAL:HG11	1.88	0.56
6:X:1076:VAL:HG12	6:X:1078:LEU:HD22	1.88	0.56
6:X:1143:ASN:HD21	6:X:1146:LEU:HD22	1.70	0.56
6:Y:29:THR:HG22	6:Y:68:PRO:O	2.06	0.56
6:Y:398:ILE:O	6:Y:401:THR:HB	2.05	0.56
1:j:25:TYR:HB2	1:j:40:TYR:HB2	1.87	0.56
3:B:525:ILE:HG23	3:B:526:PRO:HD2	1.86	0.56
3:B:561:ASP:O	3:B:562:TYR:HB2	2.04	0.56
3:D:47:LEU:CD2	3:D:64:VAL:HG11	2.36	0.56
3:D:289:LEU:CD2	3:F:515:ASN:ND2	2.68	0.56
3:D:423:GLN:O	3:D:423:GLN:HG3	2.04	0.56
2:E:26:MET:O	3:F:216:PRO:HG3	2.05	0.56
2:E:28:SER:CB	3:F:215:TYR:HA	2.36	0.56
7:E:101:MYR:C2	3:F:191:LYS:HZ3	1.96	0.56
3:F:94:PRO:HG2	3:F:210:VAL:HG22	1.87	0.56
3:F:280:LEU:HD12	3:F:486:PRO:CG	2.36	0.56
3:F:341:MET:HB2	3:F:431:ILE:HD13	1.88	0.56
3:F:443:PHE:HB2	3:F:457:PRO:CB	2.24	0.56
2:G:28:SER:CB	3:H:215:TYR:HA	2.36	0.56
3:H:336:THR:HG23	3:H:456:ASN:OD1	2.06	0.56
3:H:357:ILE:HD11	3:H:414:GLN:CG	2.36	0.56
3:H:382:ILE:HA	3:H:385:VAL:CG1	2.34	0.56

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:J:212:ILE:CD1	3:J:220:PRO:HB3	2.36	0.56
3:L:354:SER:CB	3:L:419:MET:HG2	2.35	0.56
2:M:9:ASN:CB	3:N:209:ASN:CG	2.79	0.56
3:P:212:ILE:CD1	3:P:220:PRO:HB3	2.36	0.56
3:T:427:GLU:HB2	3:T:430:THR:HG23	1.87	0.56
4:V:187:GLN:HE22	6:Y:436:LEU:HD12	1.71	0.56
5:W:1013:ILE:O	5:W:1017:ILE:HG13	2.06	0.56
6:X:329:MET:HG3	6:X:370:ARG:CZ	2.36	0.56
6:X:459:LEU:HB3	6:X:807:PHE:HZ	1.71	0.56
6:X:706:ALA:O	6:X:709:ILE:HG22	2.06	0.56
6:Y:37:THR:HG22	6:Y:38:SER:N	2.20	0.56
1:b:23:THR:HG23	1:b:25:TYR:CD2	2.40	0.55
1:b:42:THR:HG22	1:b:47:THR:HG22	1.88	0.55
1:h:42:THR:HG22	1:h:47:THR:HG22	1.88	0.55
3:B:354:SER:HB2	3:B:419:MET:CA	2.35	0.55
3:B:357:ILE:HD11	3:B:414:GLN:CG	2.36	0.55
3:D:450:SER:O	3:D:453:ASN:HB2	2.05	0.55
3:F:287:PRO:O	3:F:471:LEU:HD11	2.06	0.55
3:F:357:ILE:CG2	3:F:359:PRO:HD3	2.36	0.55
3:F:610:THR:O	3:F:612:SER:N	2.39	0.55
3:H:268:GLY:O	3:H:269:LYS:C	2.49	0.55
3:H:610:THR:O	3:H:612:SER:N	2.39	0.55
2:I:7:SER:O	2:I:8:VAL:C	2.49	0.55
3:J:412:ARG:NH2	3:L:356:THR:CG2	2.63	0.55
3:N:167:MET:HG2	3:N:215:TYR:CE1	2.42	0.55
3:N:280:LEU:HD12	3:N:486:PRO:CG	2.36	0.55
2:O:28:SER:CB	3:P:215:TYR:HA	2.36	0.55
3:P:82:PHE:HZ	3:T:238:VAL:CG1	2.19	0.55
3:P:147:ARG:CB	3:P:147:ARG:HH11	2.18	0.55
3:P:382:ILE:HA	3:P:385:VAL:CG1	2.34	0.55
3:T:287:PRO:O	3:T:471:LEU:HD11	2.05	0.55
4:U:26:LEU:O	4:U:26:LEU:HD23	2.06	0.55
5:W:934:ASN:CB	5:W:1023:THR:HG22	2.36	0.55
5:W:1281:ALA:HB3	5:W:1284:TYR:HD2	1.70	0.55
6:X:411:GLN:HB2	6:X:416:ILE:HG23	1.88	0.55
6:X:721:VAL:HG12	6:X:722:ASN:O	2.05	0.55
6:X:763:VAL:HG22	6:X:763:VAL:O	2.06	0.55
6:X:1078:LEU:N	6:X:1078:LEU:CD2	2.69	0.55
6:X:1149:THR:HB	6:X:1162:ALA:HB3	1.88	0.55
6:Y:1037:VAL:HG11	6:Y:1039:ARG:NH2	2.20	0.55
3:B:271:LEU:HG	3:B:517:LEU:CB	2.30	0.55

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:D:298:PHE:HE1	3:H:416:ARG:CB	2.16	0.55
2:G:7:SER:O	2:G:8:VAL:C	2.49	0.55
3:H:212:ILE:CD1	3:H:220:PRO:HB3	2.37	0.55
3:H:287:PRO:HB3	3:H:509:LEU:CD2	2.32	0.55
2:I:28:SER:CB	3:J:215:TYR:HA	2.36	0.55
2:K:26:MET:O	3:L:216:PRO:HG3	2.05	0.55
3:L:238:VAL:CG1	3:N:82:PHE:HZ	2.19	0.55
3:L:357:ILE:HD11	3:L:414:GLN:CG	2.36	0.55
3:L:443:PHE:HB2	3:L:457:PRO:CB	2.24	0.55
3:L:596:MET:O	3:L:597:LEU:HB2	2.05	0.55
3:N:610:THR:O	3:N:612:SER:N	2.39	0.55
3:R:147:ARG:CB	3:R:147:ARG:HH11	2.18	0.55
3:R:369:SER:HB3	3:R:445:ILE:HA	1.86	0.55
3:T:354:SER:HB2	3:T:419:MET:CA	2.35	0.55
4:U:365:THR:C	4:U:367:GLU:H	2.15	0.55
4:V:42:SER:OG	6:X:1025:ASN:ND2	2.39	0.55
5:W:5:PHE:CZ	5:W:346:PRO:HB3	2.42	0.55
5:W:817:SER:HB3	5:W:1000:TYR:CD2	2.41	0.55
6:X:278:LEU:HD12	6:X:900:GLU:OE1	2.06	0.55
6:X:788:VAL:HG22	6:X:789:ASP:N	2.22	0.55
6:Y:207:LEU:HD22	6:Y:244:VAL:HB	1.88	0.55
6:Y:474:GLU:HA	6:Y:477:SER:HB3	1.88	0.55
6:Y:1136:ALA:O	6:Y:1137:PHE:HB3	2.06	0.55
1:l:42:THR:HG22	1:l:47:THR:HG22	1.88	0.55
1:f:25:TYR:HB2	1:f:40:TYR:HB2	1.87	0.55
1:d:44:GLY:CA	3:D:399:ALA:HB3	2.36	0.55
1:d:45:ARG:CD	3:D:401:GLY:N	2.68	0.55
1:j:42:THR:HG22	1:j:47:THR:HG22	1.87	0.55
3:B:262:ALA:HB1	3:B:520:VAL:HG13	1.89	0.55
3:B:322:TYR:CD1	3:B:396:VAL:HB	2.41	0.55
3:D:238:VAL:CG1	3:F:82:PHE:HZ	2.19	0.55
3:F:44:THR:HG22	3:F:45:GLY:H	1.72	0.55
3:F:358:LEU:CB	3:F:417:PHE:CE1	2.90	0.55
3:H:168:THR:C	3:H:170:LEU:H	2.14	0.55
2:I:26:MET:O	3:J:216:PRO:HG3	2.05	0.55
3:J:167:MET:HG2	3:J:215:TYR:CE1	2.41	0.55
3:J:168:THR:C	3:J:170:LEU:H	2.14	0.55
3:J:368:THR:HA	3:J:403:SER:HB2	1.87	0.55
2:K:28:SER:CB	3:L:215:TYR:HA	2.36	0.55
3:L:67:ARG:HG3	3:L:67:ARG:HH11	1.72	0.55
3:L:336:THR:HG23	3:L:456:ASN:OD1	2.06	0.55

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:L:358:LEU:CB	3:L:417:PHE:CE1	2.90	0.55
3:P:167:MET:HG2	3:P:215:TYR:CE1	2.42	0.55
3:R:347:SER:HA	3:R:469:VAL:HG12	1.86	0.55
3:R:360:TYR:CD1	3:R:360:TYR:C	2.83	0.55
3:T:357:ILE:HD11	3:T:414:GLN:CG	2.36	0.55
3:T:372:ALA:HB3	3:T:444:LEU:CD1	2.33	0.55
4:U:40:ARG:HH22	4:U:69:ASN:HD21	1.53	0.55
4:U:64:ARG:HH11	4:U:64:ARG:HG2	1.71	0.55
4:V:43:HIS:CB	6:X:1027:LEU:HD11	2.35	0.55
5:W:250:TYR:O	5:W:334:ARG:HG3	2.06	0.55
5:W:1096:PHE:HD1	5:W:1096:PHE:N	2.00	0.55
6:X:371:LEU:HD11	6:X:388:MET:HG3	1.88	0.55
6:X:396:LEU:HD11	6:X:1194:LEU:HB2	1.87	0.55
6:X:889:ARG:HG3	6:X:890:GLN:N	2.19	0.55
6:X:1015:GLY:CA	6:X:1034:ASP:HA	2.36	0.55
6:X:1019:THR:HG21	6:Y:429:ASN:ND2	2.20	0.55
6:Y:893:VAL:HG13	6:Y:897:ALA:HB3	1.88	0.55
6:Y:1154:ILE:O	6:Y:1155:ASP:HB3	2.06	0.55
1:n:43:CYS:C	3:N:322:TYR:OH	2.50	0.55
3:B:147:ARG:CB	3:B:147:ARG:HH11	2.18	0.55
3:B:248:GLU:CD	3:B:248:GLU:N	2.64	0.55
3:B:304:ALA:O	3:B:307:ILE:HG22	2.07	0.55
2:C:7:SER:O	2:C:8:VAL:C	2.49	0.55
3:D:212:ILE:CD1	3:D:220:PRO:HB3	2.36	0.55
3:F:212:ILE:CD1	3:F:220:PRO:HB3	2.37	0.55
3:F:248:GLU:CD	3:F:248:GLU:N	2.64	0.55
3:F:304:ALA:O	3:F:307:ILE:HG22	2.07	0.55
3:F:336:THR:HG23	3:F:456:ASN:OD1	2.06	0.55
3:F:354:SER:HB2	3:F:419:MET:CA	2.35	0.55
3:J:610:THR:O	3:J:612:SER:N	2.39	0.55
3:N:268:GLY:O	3:N:269:LYS:C	2.49	0.55
2:O:35:ASP:O	2:O:36:LEU:HD23	2.06	0.55
3:P:44:THR:HG22	3:P:45:GLY:H	1.72	0.55
3:P:341:MET:CB	3:P:437:ILE:HG23	2.29	0.55
2:Q:7:SER:O	2:Q:8:VAL:C	2.49	0.55
3:R:167:MET:HG2	3:R:215:TYR:CE1	2.42	0.55
3:R:238:VAL:CG1	3:T:82:PHE:HZ	2.19	0.55
3:T:525:ILE:HG23	3:T:526:PRO:HD2	1.86	0.55
4:U:79:PHE:HB2	4:U:120:LEU:HD22	1.88	0.55
4:U:198:GLN:CD	6:X:661:THR:CG2	2.78	0.55
4:V:140:GLN:HE22	4:V:144:ILE:HD11	1.70	0.55

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
5:W:49:PHE:CE1	5:W:54:ARG:HA	2.41	0.55
5:W:68:TRP:HB2	5:W:123:GLY:HA2	1.88	0.55
6:X:347:ARG:HG3	6:X:368:ILE:HD11	1.87	0.55
6:X:481:ARG:HD3	6:X:529:TRP:CD1	2.41	0.55
6:X:580:HIS:O	6:X:581:SER:C	2.47	0.55
6:X:785:VAL:HG12	6:X:786:ALA:N	2.22	0.55
6:X:931:ILE:CG1	6:Y:689:VAL:HG11	2.36	0.55
6:X:1054:VAL:HG21	6:Y:1212:SER:HB3	1.87	0.55
6:Y:390:LEU:HA	6:Y:1191:TYR:HA	1.88	0.55
6:Y:815:PRO:HG2	6:Y:816:ALA:H	1.72	0.55
1:l:24:LEU:HD22	1:l:80:ARG:CD	2.35	0.55
1:b:15:ARG:CZ	3:B:406:ASN:CG	2.80	0.55
1:h:25:TYR:HB2	1:h:40:TYR:HB2	1.87	0.55
1:j:8:GLN:HG2	3:L:302:GLU:HG3	0.59	0.55
1:n:42:THR:HG22	1:n:47:THR:HG22	1.88	0.55
1:p:44:GLY:C	3:P:399:ALA:O	2.49	0.55
1:r:59:HIS:CA	3:R:398:SER:OG	2.40	0.55
3:D:346:ASP:HB3	3:D:428:ARG:N	2.15	0.55
3:H:49:ARG:NH1	3:H:115:THR:HG22	2.18	0.55
3:H:262:ALA:HB1	3:H:520:VAL:HG13	1.89	0.55
3:J:80:GLU:O	3:J:84:LYS:HB2	2.07	0.55
3:J:280:LEU:HD12	3:J:486:PRO:CG	2.36	0.55
2:K:13:ILE:HG23	2:K:14:THR:N	2.20	0.55
3:N:341:MET:HB2	3:N:431:ILE:HD13	1.89	0.55
3:N:358:LEU:CB	3:N:417:PHE:CE1	2.90	0.55
3:P:357:ILE:HD11	3:P:414:GLN:CG	2.36	0.55
3:P:545:VAL:CG2	3:R:641:ILE:CD1	2.85	0.55
3:R:304:ALA:O	3:R:307:ILE:HG22	2.07	0.55
3:R:341:MET:HB2	3:R:431:ILE:HD13	1.88	0.55
3:R:471:LEU:HD23	3:R:471:LEU:H	1.72	0.55
3:T:341:MET:HB2	3:T:431:ILE:HD13	1.88	0.55
4:U:41:PHE:CD1	6:X:404:MET:HE1	2.40	0.55
4:U:178:THR:HG21	6:X:376:LEU:CD2	2.16	0.55
4:U:337:ARG:HG3	4:U:337:ARG:HH11	1.70	0.55
5:W:580:ALA:HB3	5:W:618:LEU:HA	1.87	0.55
5:W:708:LEU:CD2	5:W:764:THR:HG22	2.37	0.55
6:X:325:THR:CG2	6:X:347:ARG:HG2	2.36	0.55
6:X:1020:ILE:HD12	6:X:1062:TYR:CD2	2.42	0.55
6:Y:29:THR:HG22	6:Y:68:PRO:HB2	1.87	0.55
6:Y:122:CYS:SG	6:Y:135:HIS:CE1	3.00	0.55
6:Y:1085:LEU:HD12	6:Y:1085:LEU:H	1.72	0.55

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:n:21:ARG:HH11	1:n:21:ARG:CG	2.17	0.55
3:B:610:THR:O	3:B:612:SER:N	2.39	0.55
3:F:67:ARG:HH11	3:F:67:ARG:HG3	1.72	0.55
3:F:80:GLU:O	3:F:84:LYS:HB2	2.07	0.55
3:F:543:LEU:HD21	3:F:590:VAL:HG13	1.89	0.55
3:J:357:ILE:HD11	3:J:414:GLN:CG	2.36	0.55
3:J:612:SER:O	2:K:32:PRO:HA	2.07	0.55
3:L:289:LEU:CD2	3:N:515:ASN:ND2	2.68	0.55
3:L:368:THR:HA	3:L:403:SER:HB2	1.87	0.55
3:L:545:VAL:CG2	3:N:641:ILE:CD1	2.85	0.55
3:N:357:ILE:CG2	3:N:359:PRO:HD3	2.36	0.55
3:P:617:VAL:HG21	3:R:158:LEU:HD21	1.87	0.55
3:R:44:THR:HG22	3:R:45:GLY:H	1.72	0.55
3:R:212:ILE:CD1	3:R:220:PRO:HB3	2.37	0.55
3:R:280:LEU:HD12	3:R:486:PRO:CG	2.36	0.55
3:R:354:SER:HB2	3:R:419:MET:CA	2.35	0.55
3:T:168:THR:C	3:T:170:LEU:H	2.14	0.55
3:T:212:ILE:CD1	3:T:220:PRO:HB3	2.36	0.55
5:W:1037:VAL:HG12	5:W:1039:PRO:CD	2.28	0.55
6:X:428:LEU:HD21	6:X:842:ALA:HA	1.89	0.55
6:X:580:HIS:NE2	6:X:616:SER:O	2.40	0.55
6:X:743:THR:HG21	6:X:822:MET:HA	1.89	0.55
1:l:21:ARG:CZ	3:L:327:TYR:CE1	2.90	0.55
1:b:59:HIS:HE1	1:b:61:ASN:ND2	1.97	0.55
1:j:6:ILE:HD11	3:L:586:VAL:CG1	2.32	0.55
3:B:341:MET:HB2	3:B:431:ILE:HD13	1.88	0.55
3:B:368:THR:HA	3:B:403:SER:HB2	1.87	0.55
3:B:543:LEU:HD21	3:B:590:VAL:HG13	1.89	0.55
2:C:41:LEU:O	2:C:42:ASN:CB	2.37	0.55
3:D:74:VAL:HG12	3:D:74:VAL:O	2.04	0.55
3:D:167:MET:HG2	3:D:215:TYR:CE1	2.42	0.55
3:F:168:THR:C	3:F:170:LEU:H	2.14	0.55
3:H:167:MET:HG2	3:H:215:TYR:CE1	2.42	0.55
3:H:526:PRO:HG2	3:H:533:ASP:HB3	1.89	0.55
2:I:20:PHE:CE2	2:I:22:PRO:HD3	2.42	0.55
7:I:101:MYR:H142	7:I:101:MYR:C10	2.10	0.55
3:J:82:PHE:HZ	3:N:238:VAL:CG1	2.19	0.55
3:J:154:ARG:NH1	3:J:154:ARG:CB	2.68	0.55
3:J:304:ALA:O	3:J:307:ILE:HG22	2.07	0.55
3:J:545:VAL:CG2	3:L:641:ILE:CD1	2.85	0.55
2:K:41:LEU:O	2:K:42:ASN:CB	2.37	0.55

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:L:610:THR:O	3:L:612:SER:N	2.39	0.55
3:P:238:VAL:CG2	3:R:78:MET:HE1	2.07	0.55
3:P:289:LEU:CD2	3:R:515:ASN:ND2	2.68	0.55
3:P:641:ILE:CD1	3:T:545:VAL:CG2	2.85	0.55
3:R:610:THR:O	3:R:612:SER:N	2.39	0.55
3:T:44:THR:HG22	3:T:45:GLY:H	1.72	0.55
3:T:271:LEU:HG	3:T:517:LEU:CB	2.30	0.55
3:T:543:LEU:HD21	3:T:590:VAL:HG13	1.89	0.55
4:U:214:ILE:HA	4:U:223:LEU:CD1	2.37	0.55
4:U:239:PRO:O	4:U:240:ALA:HB2	2.07	0.55
4:V:192:PHE:CA	6:Y:376:LEU:CD2	2.85	0.55
6:X:429:ASN:ND2	6:X:429:ASN:H	2.05	0.55
6:X:603:VAL:CG1	6:Y:715:VAL:CG1	2.73	0.55
6:X:650:TRP:HB3	6:X:773:ARG:HD3	1.89	0.55
6:X:1096:GLU:HG3	6:X:1118:CYS:HB2	1.88	0.55
6:Y:266:PRO:HA	6:Y:976:ASN:HD21	1.72	0.55
6:Y:410:HIS:CD2	6:Y:411:GLN:H	2.21	0.55
6:Y:835:HIS:HE1	6:Y:1212:SER:HA	1.72	0.55
1:l:15:ARG:NH2	3:L:402:ALA:HB2	2.19	0.55
1:b:44:GLY:O	3:B:399:ALA:CB	2.54	0.55
3:B:80:GLU:O	3:B:84:LYS:HB2	2.07	0.55
3:B:268:GLY:O	3:B:269:LYS:C	2.49	0.55
3:B:336:THR:HG23	3:B:456:ASN:OD1	2.06	0.55
2:C:32:PRO:HA	3:H:612:SER:O	2.07	0.55
3:D:58:ILE:HD11	3:D:127:LEU:CD2	2.37	0.55
3:D:358:LEU:CB	3:D:417:PHE:CE1	2.90	0.55
3:D:545:VAL:CG2	3:F:641:ILE:CD1	2.85	0.55
2:E:7:SER:O	2:E:8:VAL:C	2.49	0.55
2:E:35:ASP:O	2:E:36:LEU:HD23	2.06	0.55
3:F:49:ARG:NH1	3:F:115:THR:HG22	2.18	0.55
3:H:531:THR:O	3:H:532:ASP:C	2.50	0.55
3:J:237:GLU:OE1	3:L:46:LYS:NZ	2.38	0.55
3:J:412:ARG:HH22	3:L:356:THR:HG23	1.65	0.55
3:L:427:GLU:HB2	3:L:430:THR:HG23	1.87	0.55
2:M:20:PHE:CE2	2:M:22:PRO:HD3	2.42	0.55
3:N:49:ARG:NH1	3:N:115:THR:HG22	2.18	0.55
3:P:67:ARG:HG3	3:P:67:ARG:HH11	1.72	0.55
3:P:262:ALA:HB1	3:P:520:VAL:HG13	1.89	0.55
3:R:87:PHE:HD1	3:T:121:ALA:HB2	1.61	0.55
3:R:416:ARG:CB	3:T:298:PHE:HE1	2.16	0.55
5:W:79:LEU:HB2	5:W:83:ASP:OD1	2.07	0.55

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
5:W:263:VAL:HG12	5:W:267:MET:HE3	1.88	0.55
5:W:409:ILE:CG2	5:W:414:ALA:HB2	2.36	0.55
5:W:418:TYR:C	5:W:418:TYR:CD1	2.85	0.55
5:W:521:ALA:CB	5:W:531:PRO:HD3	2.37	0.55
5:W:620:VAL:HG21	5:W:627:PHE:CZ	2.42	0.55
5:W:645:VAL:O	5:W:647:PRO:CD	2.47	0.55
5:W:679:LEU:HD22	5:W:1068:VAL:CG1	2.36	0.55
5:W:993:PRO:HA	5:W:1017:ILE:CD1	2.36	0.55
6:X:325:THR:OG1	6:X:328:ARG:HB2	2.07	0.55
6:X:417:VAL:HG13	6:X:418:GLN:N	2.19	0.55
6:X:524:ASP:O	6:X:526:ALA:N	2.39	0.55
6:X:580:HIS:C	6:X:582:PRO:HD2	2.31	0.55
6:X:789:ASP:HB3	6:X:933:ASP:CB	2.37	0.55
6:X:1054:VAL:HG11	6:Y:835:HIS:CE1	2.42	0.55
6:Y:126:PHE:CD2	6:Y:132:LEU:HA	2.41	0.55
6:Y:207:LEU:HD21	6:Y:265:PRO:HB3	1.87	0.55
1:l:21:ARG:CZ	3:L:327:TYR:OH	2.55	0.55
1:j:45:ARG:O	1:j:58:PRO:C	2.49	0.55
1:j:71:CYS:SG	1:j:72:SER:N	2.80	0.55
1:r:71:CYS:SG	1:r:72:SER:N	2.80	0.55
2:A:28:SER:CB	3:B:215:TYR:HA	2.36	0.55
3:B:471:LEU:H	3:B:471:LEU:HD23	1.72	0.55
3:D:410:ASN:HD22	3:F:456:ASN:HB3	1.70	0.55
3:F:526:PRO:HG2	3:F:533:ASP:HB3	1.89	0.55
2:G:20:PHE:CE2	2:G:22:PRO:HD3	2.42	0.55
3:H:44:THR:HG22	3:H:45:GLY:H	1.72	0.55
3:H:358:LEU:CB	3:H:417:PHE:CE1	2.90	0.55
3:J:358:LEU:CB	3:J:417:PHE:CE1	2.90	0.55
3:L:147:ARG:CB	3:L:147:ARG:HH11	2.18	0.55
3:L:212:ILE:CD1	3:L:220:PRO:HB3	2.37	0.55
3:L:304:ALA:O	3:L:307:ILE:HG22	2.07	0.55
3:N:526:PRO:HG2	3:N:533:ASP:HB3	1.89	0.55
3:P:268:GLY:O	3:P:269:LYS:C	2.49	0.55
3:P:304:ALA:O	3:P:307:ILE:HG22	2.07	0.55
3:P:357:ILE:CG2	3:P:359:PRO:HD3	2.36	0.55
3:P:543:LEU:HD21	3:P:590:VAL:HG13	1.89	0.55
3:P:612:SER:O	2:Q:32:PRO:HA	2.07	0.55
2:Q:28:SER:CB	3:R:215:TYR:HA	2.36	0.55
3:R:545:VAL:CG2	3:T:641:ILE:CD1	2.85	0.55
4:U:342:ARG:O	4:U:342:ARG:HD2	2.07	0.55
5:W:241:LEU:HD23	5:W:241:LEU:C	2.31	0.55

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
5:W:641:THR:H	5:W:660:GLY:HA3	1.72	0.55
5:W:918:ILE:O	5:W:922:VAL:HG23	2.06	0.55
6:X:705:ILE:CG2	6:X:737:ARG:HB3	2.32	0.55
6:X:827:THR:HG21	6:X:831:LEU:HD23	1.87	0.55
6:X:1052:LEU:O	6:X:1056:VAL:HG23	2.06	0.55
6:Y:71:SER:O	6:Y:79:ILE:HG22	2.06	0.55
6:Y:259:LEU:HD11	6:Y:303:GLN:CG	2.28	0.55
6:Y:400:GLU:HG2	6:Y:400:GLU:O	2.05	0.55
6:Y:668:LEU:HD12	6:Y:767:PHE:CE1	2.42	0.55
6:Y:1021:VAL:HG22	6:Y:1030:GLN:O	2.07	0.55
6:Y:1210:VAL:HG22	6:Y:1211:ARG:N	2.18	0.55
1:b:17:ALA:HA	1:b:22:LEU:CD2	2.37	0.55
2:A:20:PHE:CE2	2:A:22:PRO:HD3	2.42	0.55
3:D:49:ARG:NH2	3:D:115:THR:CG2	2.70	0.55
3:D:141:GLN:C	3:D:143:MET:N	2.63	0.55
3:D:262:ALA:HB1	3:D:520:VAL:HG13	1.89	0.55
3:F:262:ALA:HB1	3:F:520:VAL:HG13	1.89	0.55
3:F:531:THR:O	3:F:532:ASP:C	2.50	0.55
3:L:167:MET:HG2	3:L:215:TYR:CE1	2.42	0.55
3:N:357:ILE:HD11	3:N:414:GLN:CG	2.36	0.55
3:N:471:LEU:HD23	3:N:471:LEU:H	1.72	0.55
3:N:543:LEU:HD21	3:N:590:VAL:HG13	1.89	0.55
2:O:26:MET:O	2:O:27:THR:C	2.50	0.55
3:P:168:THR:C	3:P:170:LEU:H	2.14	0.55
3:P:341:MET:HB2	3:P:431:ILE:HD13	1.89	0.55
3:P:358:LEU:CB	3:P:417:PHE:CE1	2.90	0.55
3:R:268:GLY:O	3:R:269:LYS:C	2.49	0.55
5:W:81:TYR:HA	5:W:180:LEU:CD2	2.36	0.55
5:W:259:ALA:HB2	5:W:304:LEU:HB2	1.89	0.55
5:W:303:LEU:HD13	5:W:303:LEU:C	2.32	0.55
5:W:554:PHE:CZ	5:W:572:THR:HG21	2.41	0.55
5:W:805:VAL:CG1	5:W:1006:SER:HB3	2.36	0.55
5:W:916:GLN:HG2	5:W:916:GLN:O	2.06	0.55
6:X:196:GLU:CD	6:X:313:LYS:HD2	2.31	0.55
6:X:201:CYS:HB2	6:X:399:ARG:CZ	2.36	0.55
6:X:441:LEU:HD23	6:X:443:PRO:HD3	1.88	0.55
6:X:795:ASP:OD1	6:X:798:VAL:HG21	2.07	0.55
6:Y:1015:GLY:HA3	6:Y:1034:ASP:HA	1.88	0.55
1:l:59:HIS:HA	3:L:398:SER:CB	2.37	0.54
1:f:53:LEU:CD1	1:f:74:GLN:CD	2.80	0.54
1:p:42:THR:HG22	1:p:47:THR:HG22	1.88	0.54

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:B:280:LEU:HD12	3:B:486:PRO:CG	2.36	0.54
3:D:87:PHE:HD1	3:F:121:ALA:HB2	1.61	0.54
2:E:20:PHE:CE2	2:E:22:PRO:HD3	2.42	0.54
3:F:238:VAL:CG1	3:H:82:PHE:HZ	2.19	0.54
3:F:272:ASP:O	3:F:273:LEU:CD2	2.56	0.54
3:J:471:LEU:HD23	3:J:471:LEU:H	1.72	0.54
3:L:262:ALA:HB1	3:L:520:VAL:HG13	1.89	0.54
3:N:80:GLU:O	3:N:84:LYS:HB2	2.07	0.54
3:N:531:THR:O	3:N:532:ASP:C	2.50	0.54
3:P:238:VAL:CG1	3:R:82:PHE:HZ	2.19	0.54
7:Q:101:MYR:H71	3:R:200:CYS:SG	2.45	0.54
3:R:531:THR:O	3:R:532:ASP:C	2.50	0.54
3:R:596:MET:O	3:R:597:LEU:HB2	2.05	0.54
3:T:167:MET:HG2	3:T:215:TYR:CE1	2.41	0.54
3:T:341:MET:CB	3:T:437:ILE:HG23	2.29	0.54
4:U:263:GLN:HA	4:U:263:GLN:NE2	2.22	0.54
5:W:224:PRO:CG	5:W:244:THR:HG22	2.37	0.54
5:W:423:LEU:HD23	5:W:426:LEU:HD13	1.88	0.54
6:X:206:LEU:H	6:X:206:LEU:CD2	2.15	0.54
6:X:843:PRO:HG2	6:X:844:LEU:H	1.71	0.54
6:X:892:ASP:OD2	6:X:897:ALA:HB2	2.06	0.54
6:X:1011:VAL:HG12	6:X:1011:VAL:O	2.07	0.54
6:X:1125:TYR:HB2	6:X:1126:PRO:HD2	1.89	0.54
6:X:1156:THR:HG21	6:X:1161:ASN:N	2.22	0.54
6:X:1182:ASN:O	6:X:1182:ASN:CG	2.50	0.54
6:Y:40:ASN:HB3	6:Y:66:SER:CB	2.33	0.54
6:Y:294:ARG:HB3	6:Y:890:GLN:HB2	1.89	0.54
1:n:25:TYR:HB2	1:n:40:TYR:HB2	1.87	0.54
2:A:26:MET:O	2:A:27:THR:C	2.50	0.54
3:B:287:PRO:O	3:B:471:LEU:HD11	2.05	0.54
3:D:272:ASP:O	3:D:273:LEU:CD2	2.56	0.54
3:D:304:ALA:O	3:D:307:ILE:HG22	2.07	0.54
3:D:368:THR:C	3:D:403:SER:HB2	2.33	0.54
3:D:612:SER:O	2:E:32:PRO:HA	2.07	0.54
3:F:167:MET:HG2	3:F:215:TYR:CE1	2.42	0.54
3:H:280:LEU:HD12	3:H:486:PRO:CG	2.36	0.54
3:N:304:ALA:O	3:N:307:ILE:HG22	2.07	0.54
2:O:20:PHE:CE2	2:O:22:PRO:HD3	2.42	0.54
3:P:80:GLU:O	3:P:84:LYS:HB2	2.07	0.54
3:R:272:ASP:O	3:R:273:LEU:CD2	2.56	0.54
3:T:67:ARG:HG3	3:T:67:ARG:HH11	1.72	0.54

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:T:350:THR:HG22	3:T:465:THR:O	2.07	0.54
3:T:526:PRO:HG2	3:T:533:ASP:HB3	1.89	0.54
4:U:178:THR:CB	6:X:376:LEU:CD2	2.84	0.54
4:U:248:LEU:HD12	4:U:249:LYS:H	1.72	0.54
4:V:64:ARG:NH2	6:X:1027:LEU:HD13	2.23	0.54
4:V:182:ILE:HD13	4:V:182:ILE:O	2.07	0.54
5:W:466:ASP:H	5:W:470:MET:HG3	1.71	0.54
5:W:603:LEU:HD23	5:W:603:LEU:C	2.32	0.54
5:W:1060:VAL:HG23	5:W:1113:THR:O	2.06	0.54
6:X:202:THR:HA	6:X:392:ASP:OD1	2.08	0.54
6:X:476:LEU:HD13	6:X:480:LEU:HD23	1.89	0.54
6:Y:655:GLU:HB2	6:Y:675:MET:O	2.07	0.54
1:b:22:LEU:N	1:b:22:LEU:HD22	2.21	0.54
1:d:22:LEU:CD2	1:d:83:VAL:HG21	2.37	0.54
1:j:44:GLY:HA3	3:J:399:ALA:HB3	1.89	0.54
3:B:67:ARG:HH11	3:B:67:ARG:HG3	1.72	0.54
3:B:360:TYR:CD1	3:B:360:TYR:C	2.83	0.54
3:D:58:ILE:O	3:D:59:ASP:CB	2.56	0.54
3:D:82:PHE:HZ	3:H:238:VAL:CG1	2.19	0.54
3:F:545:VAL:CG2	3:H:641:ILE:CD1	2.85	0.54
3:H:272:ASP:O	3:H:273:LEU:CD2	2.56	0.54
3:H:368:THR:C	3:H:403:SER:HB2	2.33	0.54
3:J:371:PRO:HB3	3:J:459:LEU:HD11	1.90	0.54
3:L:178:SER:CB	2:Q:3:ASN:HD22	2.17	0.54
3:N:212:ILE:CD1	3:N:220:PRO:HB3	2.36	0.54
3:N:368:THR:C	3:N:403:SER:HB2	2.33	0.54
2:O:7:SER:O	2:O:8:VAL:C	2.49	0.54
3:P:141:GLN:C	3:P:143:MET:N	2.63	0.54
3:P:272:ASP:O	3:P:273:LEU:CD2	2.56	0.54
3:P:526:PRO:HG2	3:P:533:ASP:HB3	1.89	0.54
3:P:610:THR:O	3:P:612:SER:N	2.39	0.54
2:Q:13:ILE:HG23	2:Q:14:THR:N	2.20	0.54
3:R:80:GLU:O	3:R:84:LYS:HB2	2.07	0.54
2:S:20:PHE:CE2	2:S:22:PRO:HD3	2.42	0.54
3:T:141:GLN:C	3:T:143:MET:N	2.63	0.54
3:T:371:PRO:HB3	3:T:459:LEU:HD11	1.90	0.54
3:T:531:THR:O	3:T:532:ASP:C	2.50	0.54
4:U:26:LEU:HD21	4:U:100:LEU:HD12	1.89	0.54
4:V:175:MET:CE	6:Y:400:GLU:OE2	2.56	0.54
5:W:66:ASN:O	5:W:174:GLY:HA3	2.07	0.54
5:W:995:CYS:HA	5:W:997:TRP:CZ3	2.42	0.54

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
5:W:1101:THR:O	5:W:1101:THR:HG23	2.08	0.54
6:X:425:ASN:C	6:X:427:LEU:H	2.16	0.54
6:X:681:ASN:HD22	6:X:752:PRO:HG2	1.72	0.54
6:X:985:VAL:HG22	6:X:1137:PHE:HE2	1.72	0.54
6:Y:202:THR:HG21	6:Y:390:LEU:CD1	2.37	0.54
6:Y:231:MET:CE	6:Y:347:ARG:NH1	2.69	0.54
6:Y:312:ASN:HB3	6:Y:1196:ARG:NH1	2.22	0.54
6:Y:775:VAL:HG12	6:Y:775:VAL:O	2.06	0.54
1:b:4:HIS:CD2	1:b:54:THR:HG22	2.33	0.54
1:b:61:ASN:HB3	3:B:396:VAL:O	2.06	0.54
3:B:272:ASP:O	3:B:273:LEU:CD2	2.56	0.54
3:B:368:THR:C	3:B:403:SER:HB2	2.33	0.54
3:D:80:GLU:O	3:D:84:LYS:HB2	2.07	0.54
3:D:641:ILE:CD1	3:H:545:VAL:CG2	2.85	0.54
3:F:152:ARG:HH11	3:F:152:ARG:CA	2.21	0.54
2:I:32:PRO:HA	3:N:612:SER:O	2.07	0.54
3:J:67:ARG:HH11	3:J:67:ARG:HG3	1.72	0.54
3:J:268:GLY:O	3:J:269:LYS:C	2.49	0.54
3:J:272:ASP:O	3:J:273:LEU:CD2	2.56	0.54
3:L:154:ARG:NH1	3:L:154:ARG:CB	2.68	0.54
3:N:262:ALA:HB1	3:N:520:VAL:HG13	1.89	0.54
3:N:287:PRO:O	3:N:471:LEU:HD11	2.05	0.54
2:O:32:PRO:HA	3:T:612:SER:O	2.07	0.54
3:P:199:LEU:HD11	3:P:203:TYR:HE2	1.73	0.54
3:P:371:PRO:HB3	3:P:459:LEU:HD11	1.90	0.54
2:Q:20:PHE:CE2	2:Q:22:PRO:HD3	2.42	0.54
3:R:346:ASP:HB3	3:R:428:ARG:N	2.15	0.54
3:R:543:LEU:HD21	3:R:590:VAL:HG13	1.89	0.54
3:T:304:ALA:O	3:T:307:ILE:HG22	2.07	0.54
3:T:358:LEU:CB	3:T:417:PHE:CE1	2.90	0.54
3:T:491:ALA:HB2	3:T:592:THR:CB	2.32	0.54
4:U:241:LEU:HA	6:X:1179:VAL:CG1	2.37	0.54
4:V:9:LEU:HD23	4:V:119:LEU:HD23	1.88	0.54
5:W:432:PHE:CG	5:W:481:ALA:HB2	2.43	0.54
5:W:558:VAL:HG22	5:W:561:GLY:H	1.71	0.54
5:W:651:ASN:HB2	5:W:688:ILE:HG21	1.88	0.54
5:W:896:ILE:O	5:W:897:PHE:HB2	2.07	0.54
6:X:377:ILE:HD11	6:X:436:LEU:HG	1.88	0.54
6:X:476:LEU:CD1	6:X:533:ALA:HA	2.37	0.54
6:X:837:TYR:C	6:X:839:LYS:N	2.62	0.54
6:Y:83:THR:O	6:Y:87:ILE:HG13	2.07	0.54

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
6:Y:197:ASP:HA	6:Y:253:GLY:HA3	1.88	0.54
6:Y:479:LEU:HB3	6:Y:758:LEU:CD1	2.35	0.54
3:B:44:THR:HG22	3:B:45:GLY:H	1.72	0.54
3:B:167:MET:HG2	3:B:215:TYR:CE1	2.42	0.54
3:B:300:LYS:HB2	3:B:303:ASP:OD2	2.08	0.54
3:D:56:ALA:HB1	3:D:61:LEU:CG	2.38	0.54
3:D:199:LEU:HD11	3:D:203:TYR:HE2	1.73	0.54
3:F:373:PHE:HD2	3:F:397:VAL:CG1	2.21	0.54
3:H:80:GLU:O	3:H:84:LYS:HB2	2.07	0.54
3:H:354:SER:HB2	3:H:419:MET:CA	2.35	0.54
3:H:373:PHE:HD2	3:H:397:VAL:CG1	2.21	0.54
3:L:416:ARG:CB	3:N:298:PHE:HE1	2.16	0.54
3:N:272:ASP:O	3:N:273:LEU:CD2	2.56	0.54
3:N:346:ASP:HB3	3:N:428:ARG:N	2.15	0.54
3:N:360:TYR:CD1	3:N:360:TYR:C	2.83	0.54
3:P:273:LEU:HA	3:R:630:LEU:HD11	1.90	0.54
3:P:350:THR:HG22	3:P:465:THR:O	2.07	0.54
2:Q:35:ASP:O	2:Q:36:LEU:HD23	2.06	0.54
3:T:272:ASP:O	3:T:273:LEU:CD2	2.56	0.54
5:W:329:THR:HG23	5:W:332:ASP:H	1.73	0.54
5:W:470:MET:HG2	5:W:473:ARG:HH21	1.71	0.54
5:W:625:ARG:HH11	5:W:625:ARG:CG	2.17	0.54
5:W:799:LEU:HD22	5:W:799:LEU:H	1.73	0.54
5:W:1038:ASN:HB2	5:W:1049:ALA:HB3	1.88	0.54
6:X:392:ASP:HB2	6:X:1188:ILE:HG22	1.90	0.54
6:X:622:PRO:O	6:X:626:LEU:HD22	2.07	0.54
6:X:751:CYS:HB3	6:X:752:PRO:HD3	1.88	0.54
6:X:784:LEU:HD21	6:X:801:ARG:HA	1.88	0.54
6:Y:126:PHE:CZ	6:Y:135:HIS:HB2	2.43	0.54
6:Y:208:ASP:HA	6:Y:241:ASN:HA	1.89	0.54
6:Y:390:LEU:HD12	6:Y:390:LEU:C	2.32	0.54
6:Y:455:GLN:HA	6:Y:455:GLN:OE1	2.08	0.54
6:Y:799:ASP:OD1	6:Y:801:ARG:HD2	2.08	0.54
6:Y:1064:TRP:CH2	6:Y:1100:LEU:HD13	2.42	0.54
1:b:15:ARG:HH12	3:B:402:ALA:HB1	1.64	0.54
1:b:71:CYS:SG	1:b:72:SER:N	2.78	0.54
1:j:58:PRO:HG2	3:J:423:GLN:CD	2.32	0.54
1:t:57:ALA:HB3	1:t:65:ILE:HD11	1.89	0.54
3:B:350:THR:HG22	3:B:465:THR:O	2.07	0.54
3:J:57:THR:CG2	3:J:58:ILE:N	2.70	0.54
3:J:300:LYS:HB2	3:J:303:ASP:OD2	2.08	0.54

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:J:543:LEU:HD21	3:J:590:VAL:HG13	1.89	0.54
2:K:26:MET:O	2:K:27:THR:C	2.50	0.54
3:L:354:SER:HB2	3:L:419:MET:CA	2.35	0.54
3:L:471:LEU:HD23	3:L:471:LEU:H	1.72	0.54
3:P:300:LYS:HB2	3:P:303:ASP:OD2	2.08	0.54
3:R:287:PRO:O	3:R:471:LEU:HD11	2.05	0.54
3:R:371:PRO:HB3	3:R:459:LEU:HD11	1.90	0.54
3:T:280:LEU:HD12	3:T:486:PRO:CG	2.36	0.54
4:U:224:ASN:HD22	4:U:226:SER:H	1.53	0.54
4:U:266:TYR:CD1	4:U:300:ILE:HG22	2.42	0.54
5:W:63:PRO:O	5:W:65:LEU:N	2.41	0.54
5:W:75:ILE:CD1	5:W:199:PRO:HG2	2.37	0.54
5:W:164:VAL:HG13	5:W:169:VAL:HG22	1.89	0.54
5:W:454:PRO:HA	5:W:457:LEU:CD2	2.37	0.54
5:W:818:PHE:CE2	5:W:989:HIS:ND1	2.61	0.54
6:X:527:ALA:HB3	6:X:817:ALA:HB3	1.89	0.54
6:X:616:SER:CB	6:Y:723:ASP:CB	2.64	0.54
6:X:926:TYR:O	6:Y:740:LYS:HB3	2.07	0.54
6:X:1017:PHE:CZ	6:Y:426:ASN:O	2.60	0.54
6:X:1040:ASP:OD1	6:X:1040:ASP:N	2.39	0.54
6:Y:117:TYR:HB3	6:Y:132:LEU:CD2	2.30	0.54
1:b:47:THR:CG2	1:b:60:ALA:HA	2.37	0.54
1:r:21:ARG:HB3	1:r:43:CYS:O	2.08	0.54
3:B:346:ASP:HB3	3:B:428:ARG:N	2.15	0.54
3:D:48:TRP:CZ2	3:D:63:ILE:HG23	2.42	0.54
3:D:471:LEU:HD23	3:D:471:LEU:H	1.72	0.54
3:D:543:LEU:HD21	3:D:590:VAL:HG13	1.89	0.54
3:F:368:THR:C	3:F:403:SER:HB2	2.33	0.54
3:J:262:ALA:HB1	3:J:520:VAL:HG13	1.89	0.54
3:J:350:THR:HG22	3:J:465:THR:O	2.07	0.54
3:L:273:LEU:HA	3:N:630:LEU:HD11	1.90	0.54
3:L:350:THR:HG22	3:L:465:THR:O	2.07	0.54
3:L:431:ILE:HD13	3:L:437:ILE:HG23	1.82	0.54
3:N:44:THR:HG22	3:N:45:GLY:H	1.72	0.54
3:N:373:PHE:HD2	3:N:397:VAL:CG1	2.21	0.54
3:P:410:ASN:ND2	3:R:455:SER:O	2.41	0.54
3:P:472:ARG:NH1	3:R:530:ARG:HD3	2.19	0.54
3:R:350:THR:HG22	3:R:465:THR:O	2.07	0.54
3:T:152:ARG:HH11	3:T:152:ARG:CA	2.21	0.54
4:U:186:ALA:HB2	4:U:282:ALA:O	2.08	0.54
4:U:209:ARG:HA	4:U:290:VAL:HG21	1.90	0.54

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
4:U:317:GLY:O	4:U:318:ASN:HB3	2.07	0.54
5:W:58:VAL:HG22	5:W:179:THR:HG22	1.90	0.54
5:W:1168:PHE:HD1	5:W:1173:TYR:CE1	2.25	0.54
6:X:528:PHE:HZ	6:X:747:ILE:CD1	2.21	0.54
6:X:528:PHE:CZ	6:X:747:ILE:HD11	2.42	0.54
6:X:984:LEU:HB2	6:X:1074:VAL:HG22	1.88	0.54
6:X:1066:MET:HB3	6:X:1072:THR:CG2	2.34	0.54
6:Y:224:TRP:HB3	6:Y:229:TYR:CE1	2.43	0.54
6:Y:298:LEU:CD1	6:Y:885:LEU:HD12	2.38	0.54
6:Y:410:HIS:CD2	6:Y:410:HIS:C	2.86	0.54
6:Y:784:LEU:HB3	6:Y:799:ASP:HB3	1.90	0.54
1:j:23:THR:HG21	3:J:323:SER:CA	2.25	0.54
1:j:61:ASN:HB3	3:J:397:VAL:HA	1.90	0.54
1:p:45:ARG:HD2	3:P:400:ALA:CA	2.37	0.54
1:p:62:VAL:O	1:p:65:ILE:N	2.40	0.54
3:B:357:ILE:CG2	3:B:359:PRO:HD3	2.36	0.54
2:C:26:MET:O	2:C:27:THR:C	2.50	0.54
3:H:304:ALA:O	3:H:307:ILE:HG22	2.07	0.54
3:J:368:THR:C	3:J:403:SER:HB2	2.33	0.54
3:J:373:PHE:HD2	3:J:397:VAL:CG1	2.21	0.54
3:J:641:ILE:CD1	3:N:545:VAL:CG2	2.85	0.54
2:K:20:PHE:CE2	2:K:22:PRO:HD3	2.42	0.54
2:K:35:ASP:O	2:K:36:LEU:HD23	2.06	0.54
3:L:152:ARG:HH11	3:L:152:ARG:CA	2.21	0.54
3:L:543:LEU:HD21	3:L:590:VAL:HG13	1.89	0.54
2:M:26:MET:O	2:M:27:THR:C	2.50	0.54
3:N:371:PRO:HB3	3:N:459:LEU:HD11	1.90	0.54
3:P:152:ARG:HH11	3:P:152:ARG:CA	2.21	0.54
3:P:280:LEU:HD12	3:P:486:PRO:CG	2.36	0.54
3:R:273:LEU:HA	3:T:630:LEU:HD11	1.90	0.54
3:R:358:LEU:CB	3:R:417:PHE:CE1	2.90	0.54
3:T:346:ASP:HB3	3:T:428:ARG:N	2.15	0.54
3:T:368:THR:CG2	3:T:446:PRO:HG3	2.38	0.54
3:T:610:THR:O	3:T:612:SER:N	2.39	0.54
4:U:270:ILE:HD13	4:U:398:GLY:HA3	1.89	0.54
4:V:366:ILE:O	4:V:370:LEU:HD23	2.07	0.54
5:W:714:ARG:HE	5:W:758:THR:CG2	2.21	0.54
5:W:815:TYR:CD1	5:W:1025:ILE:HD11	2.42	0.54
6:X:557:ILE:CA	6:X:592:LEU:HD21	2.37	0.54
6:X:945:VAL:O	6:X:945:VAL:HG13	2.08	0.54
6:X:948:GLU:O	6:X:948:GLU:HG3	2.07	0.54

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
6:X:1168:ARG:C	6:X:1170:PRO:HD3	2.33	0.54
6:Y:299:LEU:HD21	6:Y:904:ALA:CB	2.37	0.54
6:Y:448:SER:HB2	6:Y:664:GLY:HA2	1.88	0.54
6:Y:459:LEU:HD23	6:Y:785:VAL:HG22	1.89	0.54
6:Y:1020:ILE:CD1	6:Y:1041:ILE:HD11	2.38	0.54
1:l:58:PRO:CB	3:L:423:GLN:NE2	2.64	0.54
1:h:61:ASN:CB	3:H:397:VAL:HA	2.38	0.54
3:B:199:LEU:HD11	3:B:203:TYR:HE2	1.73	0.54
3:D:44:THR:HG22	3:D:45:GLY:H	1.72	0.54
2:E:26:MET:O	2:E:27:THR:C	2.50	0.54
3:F:196:LEU:HD22	3:F:196:LEU:H	1.73	0.54
2:G:26:MET:O	2:G:27:THR:C	2.50	0.54
3:H:199:LEU:HD11	3:H:203:TYR:HE2	1.73	0.54
3:H:341:MET:HB2	3:H:431:ILE:HD13	1.88	0.54
3:J:152:ARG:HH11	3:J:152:ARG:CA	2.21	0.54
3:J:158:LEU:N	3:J:158:LEU:HD23	2.23	0.54
3:J:624:LEU:HD22	3:J:628:GLN:OE1	2.08	0.54
2:K:7:SER:O	2:K:8:VAL:C	2.49	0.54
3:L:80:GLU:O	3:L:84:LYS:HB2	2.07	0.54
3:L:341:MET:HB2	3:L:431:ILE:HD13	1.88	0.54
3:N:161:LYS:HE2	3:N:162:HIS:ND1	2.23	0.54
3:N:300:LYS:HB2	3:N:303:ASP:OD2	2.08	0.54
3:P:368:THR:C	3:P:403:SER:HB2	2.33	0.54
3:P:412:ARG:NH1	3:R:459:LEU:CD2	2.71	0.54
3:P:630:LEU:HD11	3:T:273:LEU:HA	1.90	0.54
3:R:422:LEU:O	3:R:423:GLN:HB3	2.08	0.54
3:T:80:GLU:O	3:T:84:LYS:HB2	2.07	0.54
3:T:373:PHE:HD2	3:T:397:VAL:CG1	2.21	0.54
5:W:10:VAL:HG22	5:W:319:GLN:O	2.08	0.54
5:W:766:LEU:CD1	5:W:841:ARG:HH22	2.17	0.54
5:W:1091:LEU:HD13	5:W:1091:LEU:C	2.33	0.54
6:X:415:GLN:O	6:X:417:VAL:N	2.41	0.54
6:X:524:ASP:C	6:X:526:ALA:N	2.55	0.54
6:X:676:VAL:HG22	6:X:768:GLN:OE1	2.08	0.54
6:X:1130:ALA:HB2	6:Y:104:GLN:HB3	1.77	0.54
6:Y:1025:ASN:CG	6:Y:1026:GLY:N	2.57	0.54
6:Y:1037:VAL:CG1	6:Y:1039:ARG:NH2	2.71	0.54
1:l:23:THR:HB	3:L:323:SER:CB	2.38	0.54
1:h:58:PRO:HG2	3:H:423:GLN:HG3	1.90	0.54
1:p:60:ALA:C	1:p:62:VAL:N	2.65	0.54
1:r:44:GLY:O	3:R:399:ALA:CA	2.55	0.54

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:B:255:SER:HA	3:B:258:THR:CG2	2.38	0.54
3:D:296:PRO:O	3:D:297:ALA:HB3	2.08	0.54
3:D:412:ARG:NH1	3:F:459:LEU:CD2	2.71	0.54
3:F:484:ILE:HG22	3:F:485:VAL:H	1.73	0.54
3:F:494:THR:HG23	3:F:497:GLU:OE1	2.08	0.54
3:F:624:LEU:HD22	3:F:628:GLN:OE1	2.08	0.54
3:H:196:LEU:H	3:H:196:LEU:HD22	1.73	0.54
3:H:543:LEU:HD21	3:H:590:VAL:HG13	1.89	0.54
3:J:199:LEU:HD11	3:J:203:TYR:HE2	1.73	0.54
3:J:531:THR:O	3:J:532:ASP:C	2.50	0.54
3:L:373:PHE:HD2	3:L:397:VAL:CG1	2.21	0.54
3:L:412:ARG:NH1	3:N:459:LEU:CD2	2.71	0.54
3:N:137:ILE:HD13	3:N:142:ILE:HG13	1.89	0.54
3:N:161:LYS:HD3	3:N:162:HIS:CA	2.36	0.54
3:N:419:MET:HE3	3:N:419:MET:O	2.08	0.54
3:P:459:LEU:CD2	3:T:412:ARG:NH1	2.71	0.54
3:P:471:LEU:HD23	3:P:471:LEU:H	1.72	0.54
3:R:410:ASN:ND2	3:T:455:SER:O	2.41	0.54
2:S:26:MET:O	2:S:27:THR:C	2.50	0.54
3:T:158:LEU:HD23	3:T:158:LEU:N	2.23	0.54
3:T:199:LEU:HD11	3:T:203:TYR:HE2	1.73	0.54
3:T:368:THR:C	3:T:403:SER:HB2	2.33	0.54
4:V:64:ARG:NH2	6:X:1027:LEU:HD22	2.23	0.54
5:W:39:LEU:HD12	5:W:43:LEU:HB2	1.89	0.54
5:W:632:PHE:O	5:W:636:ALA:HB2	2.08	0.54
5:W:821:GLU:HA	5:W:824:ARG:HH21	1.73	0.54
5:W:932:GLN:HG2	5:W:932:GLN:O	2.08	0.54
6:X:259:LEU:O	6:X:307:SER:HB3	2.07	0.54
6:X:423:CYS:HB3	6:X:845:PHE:CE1	2.43	0.54
6:X:891:PHE:CD1	6:X:891:PHE:O	2.61	0.54
6:Y:29:THR:CG2	6:Y:68:PRO:HB2	2.38	0.54
6:Y:223:THR:HG22	6:Y:223:THR:O	2.06	0.54
6:Y:356:ASN:CG	6:Y:1151:ALA:HB1	2.33	0.54
6:Y:717:MET:HE2	6:Y:717:MET:CA	2.37	0.54
6:Y:863:ALA:O	6:Y:920:LEU:HD22	2.08	0.54
1:d:8:GLN:OE1	3:F:302:GLU:O	2.27	0.53
1:d:22:LEU:C	3:D:323:SER:HB2	2.33	0.53
1:j:61:ASN:C	1:j:63:LYS:N	2.66	0.53
3:B:212:ILE:CD1	3:B:220:PRO:HB3	2.36	0.53
3:B:373:PHE:HD2	3:B:397:VAL:CG1	2.21	0.53
2:C:35:ASP:O	2:C:36:LEU:HD23	2.06	0.53

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:D:526:PRO:HG2	3:D:533:ASP:HB3	1.89	0.53
3:F:350:THR:HG22	3:F:465:THR:O	2.07	0.53
3:F:431:ILE:CG1	3:F:437:ILE:HG21	2.38	0.53
3:H:350:THR:HG22	3:H:465:THR:O	2.07	0.53
2:I:26:MET:O	2:I:27:THR:C	2.50	0.53
3:J:455:SER:O	3:N:410:ASN:ND2	2.41	0.53
3:J:484:ILE:HG22	3:J:485:VAL:H	1.73	0.53
3:L:296:PRO:O	3:L:297:ALA:HB3	2.08	0.53
3:L:624:LEU:HD22	3:L:628:GLN:OE1	2.08	0.53
3:N:350:THR:HG22	3:N:465:THR:O	2.07	0.53
3:N:494:THR:HG23	3:N:497:GLU:OE1	2.08	0.53
3:R:141:GLN:C	3:R:143:MET:N	2.63	0.53
3:R:229:ASP:O	3:R:230:ALA:HB3	2.09	0.53
3:R:300:LYS:HB2	3:R:303:ASP:OD2	2.08	0.53
3:R:373:PHE:HD2	3:R:397:VAL:CG1	2.21	0.53
4:U:166:MET:HE1	4:U:174:LEU:CD1	2.38	0.53
4:U:316:THR:HG22	4:U:317:GLY:N	2.22	0.53
4:V:178:THR:HB	6:Y:407:ARG:HB3	1.89	0.53
6:X:215:LEU:CD2	6:X:215:LEU:N	2.72	0.53
6:X:356:ASN:ND2	6:X:1151:ALA:HB1	2.22	0.53
6:X:411:GLN:CG	6:X:445:PHE:CD2	2.91	0.53
6:X:652:GLN:NE2	6:X:773:ARG:HE	2.05	0.53
1:l:24:LEU:HD22	1:l:80:ARG:HD3	1.89	0.53
1:b:61:ASN:C	1:b:63:LYS:H	2.16	0.53
1:h:58:PRO:CB	3:H:423:GLN:CG	2.85	0.53
1:t:22:LEU:CD1	1:t:22:LEU:H	2.21	0.53
3:B:367:SER:O	3:B:368:THR:HB	2.08	0.53
3:D:410:ASN:ND2	3:F:455:SER:O	2.41	0.53
3:F:346:ASP:HB3	3:F:428:ARG:N	2.15	0.53
3:F:368:THR:CG2	3:F:446:PRO:HG3	2.38	0.53
3:F:412:ARG:NH1	3:H:459:LEU:CD2	2.71	0.53
3:F:471:LEU:H	3:F:471:LEU:HD23	1.72	0.53
3:H:141:GLN:C	3:H:143:MET:N	2.63	0.53
3:H:144:ASN:C	3:H:146:GLN:N	2.65	0.53
3:H:419:MET:HE3	3:H:419:MET:O	2.08	0.53
3:J:255:SER:HA	3:J:258:THR:CG2	2.38	0.53
3:J:630:LEU:HD11	3:N:273:LEU:HA	1.90	0.53
3:L:255:SER:HA	3:L:258:THR:CG2	2.39	0.53
3:N:67:ARG:HH11	3:N:67:ARG:HG3	1.72	0.53
3:N:229:ASP:O	3:N:230:ALA:HB3	2.09	0.53
3:P:229:ASP:O	3:P:230:ALA:HB3	2.09	0.53

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:P:312:ASN:CG	3:P:316:ILE:HD11	2.34	0.53
3:P:431:ILE:CG1	3:P:437:ILE:HG21	2.38	0.53
3:R:152:ARG:HH11	3:R:152:ARG:CA	2.21	0.53
3:T:300:LYS:HB2	3:T:303:ASP:OD2	2.08	0.53
3:T:422:LEU:O	3:T:423:GLN:HB3	2.08	0.53
4:U:228:THR:HG22	4:U:233:TYR:CD2	2.43	0.53
5:W:77:ASN:N	5:W:78:PRO:HD3	2.23	0.53
5:W:823:THR:CG2	5:W:846:ILE:HG23	2.38	0.53
6:X:519:GLN:H	6:X:520:PRO:HD3	1.72	0.53
6:X:617:HIS:CE1	6:X:619:GLY:H	2.25	0.53
6:X:1017:PHE:HZ	6:Y:426:ASN:O	1.91	0.53
6:X:1100:LEU:HD21	6:X:1136:ALA:HB2	1.90	0.53
6:X:1108:ASN:C	6:X:1110:ALA:H	2.15	0.53
6:Y:407:ARG:NH2	6:Y:410:HIS:ND1	2.56	0.53
1:l:59:HIS:CB	3:L:397:VAL:O	2.55	0.53
1:f:47:THR:HG23	1:f:59:HIS:O	2.08	0.53
1:d:15:ARG:HH22	3:D:402:ALA:HB3	1.73	0.53
1:r:23:THR:CG2	1:r:25:TYR:CE1	2.89	0.53
3:B:296:PRO:O	3:B:297:ALA:HB3	2.08	0.53
3:B:526:PRO:HG2	3:B:533:ASP:HB3	1.89	0.53
3:D:158:LEU:HD23	3:D:158:LEU:N	2.23	0.53
3:D:255:SER:HA	3:D:258:THR:CG2	2.38	0.53
3:D:373:PHE:HD2	3:D:397:VAL:CG1	2.21	0.53
3:D:549:LEU:HB3	3:F:644:LEU:CD1	2.33	0.53
3:D:624:LEU:HD22	3:D:628:GLN:OE1	2.08	0.53
3:F:300:LYS:HB2	3:F:303:ASP:OD2	2.08	0.53
3:F:312:ASN:CG	3:F:316:ILE:HD11	2.34	0.53
3:F:367:SER:O	3:F:368:THR:HB	2.08	0.53
3:H:367:SER:O	3:H:368:THR:HB	2.08	0.53
3:J:296:PRO:O	3:J:297:ALA:HB3	2.08	0.53
3:L:49:ARG:NH1	3:L:115:THR:HG22	2.18	0.53
3:L:196:LEU:H	3:L:196:LEU:HD22	1.73	0.53
3:L:300:LYS:HB2	3:L:303:ASP:OD2	2.08	0.53
3:L:431:ILE:CG1	3:L:437:ILE:HG21	2.38	0.53
3:P:367:SER:O	3:P:368:THR:HB	2.08	0.53
3:P:373:PHE:HD2	3:P:397:VAL:CG1	2.21	0.53
3:P:416:ARG:CB	3:R:298:PHE:HE1	2.16	0.53
3:P:624:LEU:HD22	3:P:628:GLN:OE1	2.08	0.53
3:T:196:LEU:H	3:T:196:LEU:HD22	1.73	0.53
4:U:37:TRP:HA	4:U:37:TRP:HE3	1.73	0.53
4:V:185:LEU:HD22	4:V:211:LEU:HD21	1.89	0.53

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
6:X:269:SER:O	6:X:270:SER:CB	2.56	0.53
1:l:22:LEU:CD2	1:l:22:LEU:H	2.21	0.53
1:f:59:HIS:CA	3:F:398:SER:OG	2.56	0.53
1:p:22:LEU:HD22	1:p:22:LEU:N	2.23	0.53
1:p:58:PRO:CB	3:P:423:GLN:HG2	2.38	0.53
1:r:25:TYR:CD2	1:r:62:VAL:CG2	2.91	0.53
1:t:44:GLY:CA	3:T:322:TYR:HH	2.09	0.53
2:C:20:PHE:CE2	2:C:22:PRO:HD3	2.42	0.53
3:D:300:LYS:HB2	3:D:303:ASP:OD2	2.08	0.53
3:D:494:THR:HG23	3:D:497:GLU:OE1	2.08	0.53
3:D:531:THR:O	3:D:532:ASP:C	2.50	0.53
2:E:27:THR:O	2:E:27:THR:CG2	2.38	0.53
3:F:199:LEU:HD11	3:F:203:TYR:HE2	1.73	0.53
3:F:273:LEU:HA	3:H:630:LEU:HD11	1.90	0.53
3:H:158:LEU:HD23	3:H:158:LEU:N	2.23	0.53
3:H:471:LEU:HD23	3:H:471:LEU:H	1.72	0.53
3:J:412:ARG:NH1	3:L:459:LEU:CD2	2.71	0.53
3:J:459:LEU:CD2	3:N:412:ARG:NH1	2.71	0.53
3:L:141:GLN:C	3:L:143:MET:N	2.63	0.53
3:L:238:VAL:CG2	3:N:78:MET:HE1	2.07	0.53
3:L:357:ILE:HD11	3:L:414:GLN:HB3	1.91	0.53
3:L:371:PRO:HB2	3:L:459:LEU:CD1	2.39	0.53
3:L:419:MET:HE3	3:L:419:MET:O	2.08	0.53
3:N:167:MET:HA	3:N:167:MET:CE	2.31	0.53
3:N:484:ILE:HG22	3:N:485:VAL:H	1.73	0.53
3:P:419:MET:HE3	3:P:419:MET:O	2.08	0.53
3:P:494:THR:HG23	3:P:497:GLU:OE1	2.09	0.53
3:R:262:ALA:HB1	3:R:520:VAL:HG13	1.89	0.53
3:R:368:THR:C	3:R:403:SER:HB2	2.33	0.53
3:R:371:PRO:CB	3:R:459:LEU:HD11	2.39	0.53
3:R:526:PRO:HG2	3:R:533:ASP:HB3	1.89	0.53
3:T:255:SER:HA	3:T:258:THR:CG2	2.39	0.53
3:T:262:ALA:HB1	3:T:520:VAL:HG13	1.89	0.53
3:T:268:GLY:O	3:T:269:LYS:C	2.49	0.53
3:T:357:ILE:HD11	3:T:414:GLN:HB3	1.91	0.53
3:T:371:PRO:CB	3:T:459:LEU:HD11	2.39	0.53
4:U:28:GLU:O	4:U:29:LEU:HD23	2.08	0.53
4:U:263:GLN:HE21	4:U:263:GLN:CA	2.20	0.53
5:W:578:VAL:O	5:W:578:VAL:HG12	2.07	0.53
6:X:495:ASN:O	6:X:498:THR:HG22	2.08	0.53
6:X:502:GLU:OE2	6:X:510:PRO:HD2	2.08	0.53

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
6:Y:277:VAL:CG2	6:Y:299:LEU:HD22	2.26	0.53
6:Y:973:ASN:HD21	6:Y:1142:ASN:HA	1.74	0.53
1:n:22:LEU:CG	1:n:83:VAL:HG21	2.37	0.53
1:n:59:HIS:HB3	3:N:423:GLN:O	2.09	0.53
1:r:61:ASN:HB3	3:R:397:VAL:HA	1.90	0.53
1:t:60:ALA:CB	3:T:322:TYR:CZ	2.88	0.53
3:B:158:LEU:HD23	3:B:158:LEU:N	2.23	0.53
3:B:484:ILE:HG22	3:B:485:VAL:H	1.73	0.53
3:B:531:THR:O	3:B:532:ASP:C	2.50	0.53
3:B:624:LEU:HD22	3:B:628:GLN:OE1	2.08	0.53
3:D:341:MET:HB2	3:D:431:ILE:HD13	1.89	0.53
3:D:350:THR:HG22	3:D:465:THR:O	2.07	0.53
3:D:419:MET:HE3	3:D:419:MET:O	2.08	0.53
3:F:141:GLN:C	3:F:143:MET:N	2.63	0.53
3:H:67:ARG:HG3	3:H:67:ARG:HH11	1.72	0.53
3:H:229:ASP:O	3:H:230:ALA:HB3	2.08	0.53
3:H:312:ASN:CG	3:H:316:ILE:HD11	2.34	0.53
3:H:371:PRO:HB3	3:H:459:LEU:HD11	1.90	0.53
3:H:484:ILE:HG22	3:H:485:VAL:H	1.73	0.53
3:J:141:GLN:C	3:J:143:MET:H	2.16	0.53
3:J:312:ASN:CG	3:J:316:ILE:HD11	2.34	0.53
3:J:341:MET:HB2	3:J:431:ILE:HD13	1.89	0.53
3:J:431:ILE:CG1	3:J:437:ILE:HG21	2.38	0.53
3:L:58:ILE:HG13	3:L:127:LEU:HD23	1.91	0.53
3:L:272:ASP:O	3:L:273:LEU:CD2	2.56	0.53
3:L:312:ASN:CG	3:L:316:ILE:HD11	2.34	0.53
3:L:526:PRO:HG2	3:L:533:ASP:HB3	1.89	0.53
3:N:158:LEU:N	3:N:158:LEU:HD23	2.23	0.53
3:N:266:MET:HE3	3:N:605:GLY:HA2	1.91	0.53
3:P:255:SER:HA	3:P:258:THR:CG2	2.38	0.53
3:P:296:PRO:O	3:P:297:ALA:HB3	2.08	0.53
3:P:371:PRO:HB2	3:P:459:LEU:CD1	2.39	0.53
3:R:67:ARG:HH11	3:R:67:ARG:HG3	1.72	0.53
3:R:312:ASN:CG	3:R:316:ILE:HD11	2.34	0.53
3:R:521:VAL:HG23	3:R:522:GLY:H	1.74	0.53
3:T:494:THR:HG23	3:T:497:GLU:OE1	2.08	0.53
4:U:277:GLN:OE1	4:U:280:GLU:HG3	2.08	0.53
4:V:204:MET:HE1	6:Y:384:MET:SD	2.48	0.53
5:W:213:ARG:NH1	5:W:372:PHE:CZ	2.77	0.53
6:X:971:GLU:HG2	6:X:974:LEU:HD22	1.91	0.53
6:X:1149:THR:CG2	6:X:1162:ALA:N	2.66	0.53

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:j:44:GLY:C	3:J:399:ALA:O	2.52	0.53
1:p:15:ARG:CZ	3:P:406:ASN:CG	2.80	0.53
3:B:58:ILE:HG13	3:B:127:LEU:HD23	1.91	0.53
3:B:229:ASP:O	3:B:230:ALA:HB3	2.08	0.53
3:B:357:ILE:HD11	3:B:414:GLN:HB3	1.91	0.53
3:B:371:PRO:HB3	3:B:459:LEU:HD11	1.90	0.53
3:D:95:LEU:HD21	3:D:207:PHE:CZ	2.44	0.53
3:D:229:ASP:O	3:D:230:ALA:HB3	2.08	0.53
3:D:455:SER:O	3:H:410:ASN:ND2	2.41	0.53
3:D:459:LEU:CD2	3:H:412:ARG:NH1	2.71	0.53
3:D:578:SER:OG	3:H:468:PRO:O	2.21	0.53
3:F:410:ASN:ND2	3:H:455:SER:O	2.41	0.53
3:H:95:LEU:HD21	3:H:207:PHE:CZ	2.44	0.53
3:H:296:PRO:O	3:H:297:ALA:HB3	2.08	0.53
3:H:300:LYS:HB2	3:H:303:ASP:OD2	2.08	0.53
3:J:411:VAL:O	3:J:412:ARG:CB	2.56	0.53
3:L:237:GLU:OE1	3:N:46:LYS:NZ	2.38	0.53
3:L:410:ASN:ND2	3:N:455:SER:O	2.41	0.53
3:P:158:LEU:HD23	3:P:158:LEU:N	2.23	0.53
3:P:371:PRO:CB	3:P:459:LEU:HD11	2.39	0.53
3:R:199:LEU:HD11	3:R:203:TYR:HE2	1.73	0.53
3:R:484:ILE:HG22	3:R:485:VAL:H	1.73	0.53
3:R:518:ALA:CB	3:R:580:PRO:HG3	2.39	0.53
3:T:494:THR:H	3:T:497:GLU:CB	2.19	0.53
4:U:241:LEU:HA	6:X:1179:VAL:HG12	1.89	0.53
5:W:213:ARG:HE	5:W:217:LEU:CD2	2.22	0.53
5:W:736:VAL:HG11	5:W:780:LEU:HD21	1.90	0.53
5:W:819:ILE:CD1	5:W:896:ILE:HD12	2.34	0.53
6:X:380:GLU:HG3	6:X:381:VAL:N	2.24	0.53
6:X:979:ARG:CG	6:X:980:GLY:N	2.67	0.53
6:Y:123:ASN:HB3	6:Y:161:LEU:HD13	1.90	0.53
6:Y:693:TYR:OH	6:Y:740:LYS:HD3	2.08	0.53
1:b:60:ALA:O	1:b:62:VAL:N	2.41	0.53
1:h:59:HIS:ND1	1:h:60:ALA:CA	2.71	0.53
1:p:60:ALA:HB3	3:P:322:TYR:HE1	1.72	0.53
3:B:494:THR:HG23	3:B:497:GLU:OE1	2.09	0.53
3:F:95:LEU:HD21	3:F:207:PHE:CZ	2.44	0.53
3:F:296:PRO:O	3:F:297:ALA:HB3	2.08	0.53
3:H:287:PRO:HG2	3:H:471:LEU:CD1	2.39	0.53
3:H:321:THR:HG23	3:H:324:GLY:O	2.09	0.53
3:H:624:LEU:HD22	3:H:628:GLN:OE1	2.08	0.53

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:J:229:ASP:O	3:J:230:ALA:HB3	2.08	0.53
3:J:368:THR:CG2	3:J:446:PRO:HG3	2.38	0.53
3:J:518:ALA:CB	3:J:580:PRO:HG3	2.39	0.53
3:J:526:PRO:HG2	3:J:533:ASP:HB3	1.89	0.53
3:L:266:MET:HE3	3:L:605:GLY:HA2	1.91	0.53
3:N:199:LEU:O	3:N:207:PHE:HE2	1.91	0.53
3:N:321:THR:HG23	3:N:324:GLY:O	2.09	0.53
3:N:494:THR:O	3:N:495:SER:C	2.52	0.53
3:P:266:MET:HE3	3:P:605:GLY:HA2	1.91	0.53
2:Q:24:SER:HB3	3:R:176:ALA:N	2.24	0.53
3:T:367:SER:O	3:T:368:THR:HB	2.08	0.53
4:U:121:ASP:CG	4:U:123:ASN:HB2	2.33	0.53
5:W:460:ALA:HB2	5:W:624:THR:HG22	1.89	0.53
5:W:614:GLY:H	5:W:662:ARG:HB3	1.72	0.53
5:W:839:GLU:HB3	5:W:861:ALA:CB	2.31	0.53
5:W:845:LEU:N	5:W:845:LEU:HD12	2.24	0.53
5:W:942:SER:HB2	5:W:947:ILE:O	2.09	0.53
6:X:347:ARG:HG3	6:X:368:ILE:HD12	1.91	0.53
6:X:355:LYS:HE3	6:X:1149:THR:HG23	1.89	0.53
6:X:701:ILE:O	6:X:705:ILE:HG12	2.09	0.53
6:X:1006:ARG:HH11	6:X:1013:VAL:HG21	1.74	0.53
6:Y:75:ASP:O	6:Y:76:GLY:C	2.52	0.53
6:Y:132:LEU:O	6:Y:136:LEU:HG	2.09	0.53
6:Y:547:GLU:OE1	6:Y:547:GLU:HA	2.09	0.53
6:Y:787:ARG:HB3	6:Y:931:ILE:HG22	1.91	0.53
6:Y:923:LEU:HD21	6:Y:931:ILE:HD13	1.91	0.53
6:Y:1023:ARG:NH2	6:Y:1030:GLN:HB3	2.23	0.53
6:Y:1100:LEU:HD12	6:Y:1135:TYR:HB2	1.89	0.53
1:j:3:LEU:HD13	1:j:3:LEU:C	2.33	0.53
1:t:6:ILE:HG21	3:P:494:THR:HG22	1.91	0.53
3:B:368:THR:CG2	3:B:446:PRO:HG3	2.38	0.53
3:B:431:ILE:CG1	3:B:437:ILE:HG21	2.38	0.53
3:D:367:SER:O	3:D:368:THR:HB	2.08	0.53
3:D:371:PRO:CB	3:D:459:LEU:HD11	2.39	0.53
3:D:630:LEU:HD11	3:H:273:LEU:HA	1.90	0.53
3:F:50:PRO:HG3	3:F:56:ALA:HB2	1.91	0.53
3:F:158:LEU:N	3:F:158:LEU:HD23	2.23	0.53
3:F:199:LEU:O	3:F:207:PHE:HE2	1.91	0.53
3:F:255:SER:HA	3:F:258:THR:CG2	2.39	0.53
3:H:255:SER:HA	3:H:258:THR:CG2	2.38	0.53
3:H:521:VAL:HG23	3:H:522:GLY:H	1.74	0.53

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:L:367:SER:O	3:L:368:THR:HB	2.08	0.53
3:L:368:THR:C	3:L:403:SER:HB2	2.33	0.53
3:L:411:VAL:O	3:L:412:ARG:CB	2.56	0.53
3:N:312:ASN:CG	3:N:316:ILE:HD11	2.34	0.53
3:P:518:ALA:CB	3:P:580:PRO:HG3	2.39	0.53
3:R:419:MET:HE3	3:R:419:MET:O	2.08	0.53
3:R:566:TRP:NE1	3:T:195:PRO:HD3	2.24	0.53
3:T:322:TYR:CD1	3:T:396:VAL:HB	2.44	0.53
3:T:411:VAL:O	3:T:412:ARG:CB	2.57	0.53
3:T:624:LEU:HD22	3:T:628:GLN:OE1	2.08	0.53
4:V:175:MET:HE2	6:Y:404:MET:CE	2.39	0.53
5:W:620:VAL:HG23	5:W:657:LEU:HD11	1.91	0.53
5:W:814:GLY:HA3	5:W:991:LEU:CD1	2.39	0.53
5:W:816:ASN:CG	5:W:845:LEU:HD21	2.33	0.53
5:W:888:ILE:O	5:W:888:ILE:HG13	2.09	0.53
6:X:545:LEU:HD22	6:X:576:THR:HG21	1.89	0.53
6:X:557:ILE:HA	6:X:592:LEU:HD21	1.88	0.53
6:X:633:PRO:C	6:X:635:GLN:H	2.17	0.53
6:X:925:LEU:HD22	6:X:926:TYR:HD1	1.74	0.53
6:X:1078:LEU:HG	6:X:1081:TYR:HD2	1.74	0.53
6:Y:599:ILE:CD1	6:Y:631:ILE:HG12	2.38	0.53
6:Y:634:GLN:CD	6:Y:634:GLN:N	2.66	0.53
1:d:15:ARG:CZ	3:D:402:ALA:HB1	2.30	0.53
1:t:22:LEU:N	1:t:22:LEU:HD13	2.24	0.53
3:B:199:LEU:O	3:B:207:PHE:HE2	1.91	0.53
3:B:443:PHE:HB2	3:B:457:PRO:CB	2.24	0.53
3:D:485:VAL:O	3:D:485:VAL:HG12	2.09	0.53
3:F:141:GLN:C	3:F:143:MET:H	2.16	0.53
3:H:494:THR:H	3:H:497:GLU:CB	2.19	0.53
3:H:494:THR:O	3:H:495:SER:C	2.52	0.53
3:J:95:LEU:HD21	3:J:207:PHE:CZ	2.44	0.53
3:J:199:LEU:O	3:J:207:PHE:HE2	1.91	0.53
3:J:266:MET:HE3	3:J:605:GLY:HA2	1.91	0.53
3:J:273:LEU:HA	3:L:630:LEU:HD11	1.90	0.53
3:J:357:ILE:HD11	3:J:414:GLN:HB3	1.91	0.53
3:J:419:MET:O	3:J:419:MET:HE3	2.08	0.53
3:J:494:THR:HG23	3:J:497:GLU:OE1	2.08	0.53
3:L:476:TYR:CD1	3:L:506:ALA:HB3	2.44	0.53
3:N:255:SER:HA	3:N:258:THR:CG2	2.38	0.53
3:N:371:PRO:HB2	3:N:459:LEU:CD1	2.39	0.53
3:P:287:PRO:HG2	3:P:471:LEU:CD1	2.39	0.53

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:P:321:THR:HG23	3:P:324:GLY:O	2.09	0.53
3:P:484:ILE:HG22	3:P:485:VAL:H	1.73	0.53
3:R:196:LEU:HD22	3:R:196:LEU:H	1.74	0.53
3:R:266:MET:HE3	3:R:605:GLY:HA2	1.91	0.53
3:R:368:THR:CG2	3:R:446:PRO:HG3	2.38	0.53
3:R:624:LEU:HD22	3:R:628:GLN:OE1	2.08	0.53
3:T:58:ILE:HG13	3:T:127:LEU:HD23	1.91	0.53
3:T:199:LEU:O	3:T:207:PHE:HE2	1.91	0.53
3:T:518:ALA:CB	3:T:580:PRO:HG3	2.39	0.53
4:U:211:LEU:HD12	4:U:286:MET:SD	2.49	0.53
4:U:357:ASN:O	4:U:360:THR:HG22	2.09	0.53
5:W:175:CYS:O	5:W:177:TYR:N	2.42	0.53
5:W:558:VAL:HG11	5:W:561:GLY:O	2.09	0.53
5:W:1145:ASP:HB2	5:W:1147:THR:HG23	1.90	0.53
5:W:1215:LEU:HD12	5:W:1215:LEU:O	2.08	0.53
6:X:302:LEU:HD23	6:X:302:LEU:C	2.34	0.53
6:X:308:ASN:HD22	6:X:395:THR:HG23	1.73	0.53
6:X:407:ARG:C	6:X:409:MET:N	2.67	0.53
6:X:450:ASP:CG	6:X:861:ARG:NH1	2.66	0.53
6:X:720:THR:O	6:X:721:VAL:C	2.51	0.53
6:Y:39:PRO:HB3	6:Y:72:VAL:HG23	1.91	0.53
6:Y:413:PRO:HG2	6:Y:440:ILE:CD1	2.39	0.53
6:Y:1075:LEU:O	6:Y:1075:LEU:HG	2.09	0.53
1:l:15:ARG:HH12	3:L:402:ALA:CB	2.20	0.53
1:r:58:PRO:CB	3:R:423:GLN:NE2	2.66	0.53
1:t:74:GLN:O	1:t:74:GLN:HG3	2.09	0.53
3:B:345:ILE:O	3:B:346:ASP:C	2.52	0.53
3:B:422:LEU:O	3:B:423:GLN:HB3	2.08	0.53
3:D:141:GLN:C	3:D:143:MET:H	2.16	0.53
3:D:199:LEU:O	3:D:207:PHE:HE2	1.92	0.53
3:D:312:ASN:CG	3:D:316:ILE:HD11	2.34	0.53
3:F:422:LEU:O	3:F:423:GLN:HB3	2.08	0.53
3:F:476:TYR:CD1	3:F:506:ALA:HB3	2.44	0.53
2:I:24:SER:HB3	3:J:176:ALA:N	2.24	0.53
3:J:44:THR:HG22	3:J:45:GLY:H	1.72	0.53
3:L:95:LEU:HD21	3:L:207:PHE:CZ	2.44	0.53
3:L:494:THR:O	3:L:495:SER:C	2.52	0.53
3:L:521:VAL:HG23	3:L:522:GLY:H	1.74	0.53
3:N:196:LEU:HD22	3:N:196:LEU:H	1.73	0.53
3:N:371:PRO:CB	3:N:459:LEU:HD11	2.39	0.53
3:N:521:VAL:HG23	3:N:522:GLY:H	1.74	0.53

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:P:195:PRO:HD3	3:T:566:TRP:NE1	2.24	0.53
3:P:422:LEU:O	3:P:423:GLN:HB3	2.08	0.53
3:P:494:THR:O	3:P:495:SER:C	2.52	0.53
3:P:521:VAL:HG23	3:P:522:GLY:H	1.74	0.53
3:P:531:THR:O	3:P:532:ASP:C	2.50	0.53
3:R:371:PRO:HB2	3:R:459:LEU:CD1	2.39	0.53
3:T:229:ASP:O	3:T:230:ALA:HB3	2.08	0.53
3:T:476:TYR:CD1	3:T:506:ALA:HB3	2.44	0.53
4:V:384:MET:N	4:V:385:PRO:HD3	2.24	0.53
5:W:770:ARG:NH1	5:W:774:LEU:HD11	2.24	0.53
5:W:1134:VAL:HG21	5:W:1221:PHE:CD1	2.44	0.53
6:X:356:ASN:O	6:X:359:PHE:N	2.41	0.53
6:X:364:ILE:HG13	6:X:367:ARG:NH2	2.21	0.53
6:X:429:ASN:N	6:X:429:ASN:ND2	2.56	0.53
6:X:520:PRO:HD2	6:X:816:ALA:CB	2.39	0.53
6:Y:427:LEU:HD13	6:Y:837:TYR:CB	2.38	0.53
6:Y:497:ILE:HD13	6:Y:497:ILE:C	2.34	0.53
6:Y:1209:TYR:CD1	6:Y:1209:TYR:N	2.77	0.53
6:Y:1212:SER:O	6:Y:1213:ALA:C	2.51	0.53
1:f:15:ARG:HH22	3:F:402:ALA:HB3	1.74	0.52
1:f:45:ARG:CG	3:F:399:ALA:O	2.56	0.52
1:h:61:ASN:HB2	3:H:396:VAL:O	2.08	0.52
1:r:3:LEU:HD12	1:r:3:LEU:N	2.23	0.52
3:F:345:ILE:O	3:F:346:ASP:C	2.52	0.52
3:F:494:THR:O	3:F:495:SER:C	2.52	0.52
3:H:422:LEU:O	3:H:423:GLN:HB3	2.08	0.52
3:H:494:THR:HG23	3:H:497:GLU:OE1	2.08	0.52
3:H:518:ALA:CB	3:H:580:PRO:HG3	2.39	0.52
3:J:49:ARG:NH1	3:J:115:THR:CG2	2.61	0.52
3:J:49:ARG:NH1	4:V:405:ALA:CB	2.72	0.52
3:L:199:LEU:HD11	3:L:203:TYR:HE2	1.73	0.52
3:L:229:ASP:O	3:L:230:ALA:HB3	2.09	0.52
3:L:357:ILE:CG2	3:L:359:PRO:HD3	2.35	0.52
2:M:24:SER:HB3	3:N:176:ALA:N	2.24	0.52
3:N:144:ASN:C	3:N:146:GLN:N	2.65	0.52
3:N:367:SER:O	3:N:368:THR:HB	2.08	0.52
2:Q:26:MET:O	2:Q:27:THR:C	2.50	0.52
3:T:144:ASN:C	3:T:146:GLN:N	2.65	0.52
4:U:112:TRP:CZ3	4:U:114:VAL:HA	2.44	0.52
4:U:198:GLN:OE1	6:X:661:THR:CG2	2.57	0.52
4:U:222:ARG:N	4:U:340:GLN:HG2	2.23	0.52

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
4:V:16:GLN:HG2	4:V:20:LEU:HD21	1.91	0.52
5:W:167:ASP:HB2	5:W:212:ALA:CB	2.31	0.52
5:W:248:LEU:HD21	6:X:647:ARG:CZ	2.39	0.52
5:W:750:PRO:HD3	5:W:791:PHE:HB3	1.90	0.52
6:X:331:ARG:HH22	6:X:1196:ARG:CZ	2.21	0.52
6:X:409:MET:CB	6:X:445:PHE:O	2.55	0.52
6:X:416:ILE:HG22	6:X:445:PHE:HZ	1.69	0.52
6:X:444:TRP:HB3	6:X:851:PHE:HB3	1.91	0.52
6:X:992:ASN:HD22	6:X:992:ASN:C	2.16	0.52
6:X:1085:LEU:HD22	6:X:1119:ILE:CD1	2.39	0.52
6:X:1183:ASP:OD2	6:X:1187:ARG:HB2	2.09	0.52
6:Y:534:SER:OG	6:Y:543:THR:HB	2.09	0.52
6:Y:764:THR:O	6:Y:764:THR:HG23	2.09	0.52
6:Y:985:VAL:HG22	6:Y:1137:PHE:CE2	2.44	0.52
6:Y:1190:LEU:N	6:Y:1190:LEU:HD22	2.23	0.52
1:l:45:ARG:HD2	3:L:400:ALA:CA	2.40	0.52
1:j:22:LEU:CD2	1:j:41:VAL:HG13	2.31	0.52
1:p:24:LEU:CD2	1:p:80:ARG:NE	2.68	0.52
3:B:167:MET:HA	3:B:167:MET:CE	2.31	0.52
3:B:312:ASN:CG	3:B:316:ILE:HD11	2.34	0.52
3:B:358:LEU:CB	3:B:417:PHE:CE1	2.90	0.52
3:B:371:PRO:HB2	3:B:459:LEU:CD1	2.39	0.52
3:D:196:LEU:H	3:D:196:LEU:HD22	1.74	0.52
3:F:371:PRO:CB	3:F:459:LEU:HD11	2.39	0.52
3:F:371:PRO:HB2	3:F:459:LEU:CD1	2.39	0.52
3:F:419:MET:O	3:F:419:MET:HE3	2.08	0.52
3:H:152:ARG:HH11	3:H:152:ARG:CA	2.21	0.52
3:H:371:PRO:CB	3:H:459:LEU:HD11	2.39	0.52
3:L:422:LEU:O	3:L:423:GLN:HB3	2.08	0.52
3:N:95:LEU:HD21	3:N:207:PHE:CZ	2.44	0.52
3:N:148:THR:HA	3:R:100:LYS:HG3	1.91	0.52
3:N:199:LEU:HD11	3:N:203:TYR:HE2	1.73	0.52
3:N:518:ALA:CB	3:N:580:PRO:HG3	2.39	0.52
3:P:238:VAL:HG11	3:R:82:PHE:CZ	2.44	0.52
3:R:249:VAL:O	3:R:250:GLY:C	2.53	0.52
3:R:412:ARG:NH1	3:T:459:LEU:CD2	2.71	0.52
3:R:468:PRO:O	3:T:578:SER:OG	2.21	0.52
3:R:494:THR:HG23	3:R:497:GLU:OE1	2.08	0.52
3:T:141:GLN:C	3:T:143:MET:H	2.16	0.52
3:T:296:PRO:O	3:T:297:ALA:HB3	2.08	0.52
3:T:406:ASN:C	3:T:406:ASN:HD22	2.17	0.52

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:T:419:MET:HE3	3:T:419:MET:O	2.08	0.52
5:W:47:LEU:HD11	5:W:327:ILE:HG23	1.90	0.52
5:W:111:ARG:HD2	5:W:139:PHE:CB	2.39	0.52
5:W:533:ILE:HG12	5:W:537:ILE:HD11	1.91	0.52
5:W:839:GLU:O	5:W:839:GLU:HG3	2.08	0.52
6:X:428:LEU:CD2	6:X:842:ALA:HA	2.39	0.52
6:X:448:SER:HB2	6:X:664:GLY:CA	2.39	0.52
6:Y:29:THR:HG22	6:Y:68:PRO:C	2.34	0.52
6:Y:871:VAL:O	6:Y:875:GLN:HG2	2.08	0.52
6:Y:1081:TYR:OH	6:Y:1138:THR:CG2	2.57	0.52
6:Y:1156:THR:CG2	6:Y:1161:ASN:OD1	2.57	0.52
1:f:44:GLY:O	3:F:399:ALA:O	2.26	0.52
1:n:22:LEU:HG	1:n:83:VAL:CG2	2.38	0.52
3:B:50:PRO:HG3	3:B:56:ALA:HB2	1.91	0.52
3:B:266:MET:HE3	3:B:605:GLY:HA2	1.91	0.52
3:B:371:PRO:CB	3:B:459:LEU:HD11	2.39	0.52
3:D:266:MET:HE3	3:D:605:GLY:HA2	1.91	0.52
3:D:422:LEU:O	3:D:423:GLN:HB3	2.08	0.52
3:D:644:LEU:CD1	3:H:549:LEU:HB3	2.33	0.52
3:F:229:ASP:O	3:F:230:ALA:HB3	2.09	0.52
3:F:249:VAL:O	3:F:250:GLY:C	2.53	0.52
3:H:58:ILE:HG13	3:H:127:LEU:HD23	1.91	0.52
3:H:141:GLN:C	3:H:143:MET:H	2.16	0.52
3:H:199:LEU:O	3:H:207:PHE:HE2	1.91	0.52
2:I:29:THR:HG22	3:N:609:ARG:HD2	1.91	0.52
3:J:141:GLN:C	3:J:143:MET:N	2.63	0.52
3:J:371:PRO:CB	3:J:459:LEU:HD11	2.39	0.52
3:J:410:ASN:ND2	3:L:455:SER:O	2.41	0.52
3:J:485:VAL:HG12	3:J:485:VAL:O	2.09	0.52
3:L:199:LEU:O	3:L:207:PHE:HE2	1.91	0.52
3:L:371:PRO:HB3	3:L:459:LEU:HD11	1.90	0.52
3:L:485:VAL:O	3:L:485:VAL:HG12	2.09	0.52
3:P:199:LEU:O	3:P:207:PHE:HE2	1.91	0.52
3:R:58:ILE:HG13	3:R:127:LEU:HD23	1.91	0.52
3:R:255:SER:HA	3:R:258:THR:CG2	2.38	0.52
3:R:296:PRO:O	3:R:297:ALA:HB3	2.08	0.52
3:R:342:ASN:OD1	3:R:431:ILE:HG21	2.10	0.52
3:R:357:ILE:HD11	3:R:414:GLN:HB3	1.91	0.52
3:T:471:LEU:HD23	3:T:471:LEU:H	1.72	0.52
3:T:521:VAL:HG23	3:T:522:GLY:H	1.74	0.52
4:U:23:LEU:HD11	4:U:27:GLN:HE21	1.74	0.52

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
4:U:136:TRP:CE3	4:U:197:PRO:HG3	2.44	0.52
4:U:240:ALA:O	4:U:241:LEU:HD23	2.08	0.52
5:W:251:SER:HB2	5:W:255:ILE:HD11	1.90	0.52
5:W:296:ASP:HB3	5:W:305:GLN:NE2	2.24	0.52
5:W:304:LEU:HD23	5:W:304:LEU:C	2.34	0.52
5:W:439:PHE:CD2	5:W:684:ARG:NH1	2.77	0.52
5:W:1187:ALA:HB3	5:W:1188:PRO:HD3	1.91	0.52
6:X:241:ASN:HB2	6:X:1148:SER:OG	2.08	0.52
6:X:599:ILE:HD13	6:X:639:LEU:HD21	1.90	0.52
6:Y:220:ILE:HG22	6:Y:221:GLY:N	2.22	0.52
6:Y:457:MET:HG2	6:Y:860:THR:CG2	2.38	0.52
6:Y:610:THR:HG21	6:Y:614:GLN:HG2	1.91	0.52
1:b:22:LEU:CD2	1:b:22:LEU:N	2.73	0.52
1:b:59:HIS:HA	3:B:398:SER:OG	2.08	0.52
2:A:24:SER:HB3	3:B:176:ALA:N	2.24	0.52
3:B:93:GLN:N	3:B:94:PRO:CD	2.73	0.52
3:B:152:ARG:HH11	3:B:152:ARG:CA	2.21	0.52
3:B:322:TYR:CE1	3:B:396:VAL:HB	2.45	0.52
3:D:249:VAL:O	3:D:250:GLY:C	2.53	0.52
3:F:86:LEU:HD12	3:F:108:LEU:HB2	1.92	0.52
3:F:357:ILE:HD11	3:F:414:GLN:HB3	1.91	0.52
3:F:485:VAL:HG12	3:F:485:VAL:O	2.09	0.52
3:F:518:ALA:CB	3:F:580:PRO:HG3	2.39	0.52
2:G:27:THR:O	2:G:27:THR:CG2	2.38	0.52
3:H:93:GLN:N	3:H:94:PRO:CD	2.73	0.52
3:H:496:GLU:C	3:H:498:VAL:H	2.17	0.52
3:J:494:THR:O	3:J:495:SER:C	2.52	0.52
3:J:644:LEU:CD1	3:N:549:LEU:HB3	2.34	0.52
7:K:101:MYR:H71	3:L:200:CYS:SG	2.45	0.52
3:L:158:LEU:HD23	3:L:158:LEU:N	2.23	0.52
3:L:371:PRO:CB	3:L:459:LEU:HD11	2.39	0.52
3:L:531:THR:O	3:L:532:ASP:C	2.50	0.52
3:L:566:TRP:NE1	3:N:195:PRO:HD3	2.24	0.52
3:N:141:GLN:C	3:N:143:MET:H	2.16	0.52
3:N:269:LYS:O	3:N:270:ASP:HB2	2.10	0.52
3:N:368:THR:CG2	3:N:446:PRO:HG3	2.38	0.52
3:P:58:ILE:HG12	3:P:127:LEU:HG	1.90	0.52
3:P:345:ILE:O	3:P:346:ASP:C	2.52	0.52
3:P:357:ILE:HD11	3:P:414:GLN:HB3	1.91	0.52
3:R:95:LEU:HD21	3:R:207:PHE:CZ	2.44	0.52
3:T:86:LEU:HD12	3:T:108:LEU:HB2	1.92	0.52

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:T:321:THR:HG23	3:T:324:GLY:O	2.09	0.52
3:T:357:ILE:CG2	3:T:359:PRO:HD3	2.36	0.52
4:V:61:VAL:HG11	4:V:278:ILE:HA	1.92	0.52
4:V:275:GLN:HE22	4:V:389:PHE:HD2	1.58	0.52
5:W:102:ARG:HH11	5:W:102:ARG:CG	2.21	0.52
5:W:210:LEU:N	5:W:211:PRO:HD2	2.25	0.52
5:W:874:TYR:CE2	5:W:876:GLN:HG3	2.42	0.52
6:X:722:ASN:CG	6:X:723:ASP:N	2.67	0.52
6:Y:197:ASP:OD1	6:Y:310:LYS:HD3	2.10	0.52
6:Y:252:LEU:CD1	6:Y:336:ARG:HG2	2.40	0.52
6:Y:1074:VAL:CG1	6:Y:1075:LEU:N	2.71	0.52
1:b:45:ARG:HD2	3:B:400:ALA:C	2.35	0.52
1:d:44:GLY:C	3:D:399:ALA:O	2.52	0.52
3:B:321:THR:HG23	3:B:324:GLY:O	2.09	0.52
3:B:476:TYR:CD1	3:B:506:ALA:HB3	2.44	0.52
3:B:485:VAL:HG12	3:B:485:VAL:O	2.09	0.52
3:D:93:GLN:N	3:D:94:PRO:CD	2.73	0.52
3:D:345:ILE:O	3:D:346:ASP:C	2.52	0.52
3:H:368:THR:CG2	3:H:446:PRO:HG3	2.38	0.52
3:J:269:LYS:O	3:J:270:ASP:HB2	2.09	0.52
3:J:342:ASN:OD1	3:J:431:ILE:HG21	2.10	0.52
3:J:371:PRO:HB2	3:J:459:LEU:CD1	2.39	0.52
3:J:545:VAL:CG2	3:L:641:ILE:HD11	2.40	0.52
2:K:24:SER:HB3	3:L:176:ALA:N	2.24	0.52
3:L:518:ALA:CB	3:L:580:PRO:HG3	2.39	0.52
3:N:422:LEU:O	3:N:423:GLN:HB3	2.08	0.52
3:N:431:ILE:HD13	3:N:437:ILE:HG23	1.83	0.52
3:N:624:LEU:HD22	3:N:628:GLN:OE1	2.08	0.52
3:P:342:ASN:OD1	3:P:431:ILE:HG21	2.10	0.52
3:R:287:PRO:HG2	3:R:471:LEU:CD1	2.39	0.52
3:T:93:GLN:N	3:T:94:PRO:CD	2.73	0.52
3:T:296:PRO:C	3:T:298:PHE:N	2.68	0.52
3:T:345:ILE:O	3:T:346:ASP:C	2.52	0.52
3:T:371:PRO:HB2	3:T:459:LEU:CD1	2.39	0.52
4:U:121:ASP:C	4:U:123:ASN:N	2.66	0.52
5:W:5:PHE:CD1	5:W:306:ARG:NH1	2.77	0.52
5:W:95:PHE:HD2	5:W:125:PHE:CG	2.28	0.52
5:W:167:ASP:OD1	5:W:167:ASP:N	2.43	0.52
5:W:282:VAL:HG12	5:W:283:THR:N	2.22	0.52
5:W:398:LEU:HD22	5:W:792:THR:HG21	1.92	0.52
5:W:424:SER:CB	5:W:425:PRO:CD	2.87	0.52

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
5:W:697:PHE:O	5:W:700:VAL:HG12	2.10	0.52
5:W:782:THR:O	5:W:786:VAL:HG23	2.09	0.52
5:W:823:THR:HG21	5:W:846:ILE:HG23	1.91	0.52
6:X:531:CYS:SG	6:X:818:ILE:HD11	2.49	0.52
6:Y:277:VAL:CG1	6:Y:278:LEU:HG	2.34	0.52
6:Y:782:ASN:HB3	6:Y:784:LEU:HD11	1.92	0.52
1:h:61:ASN:HB3	3:H:397:VAL:HA	1.91	0.52
1:n:74:GLN:NE2	1:n:74:GLN:CA	2.73	0.52
1:t:51:PHE:CZ	1:t:71:CYS:O	2.63	0.52
3:D:273:LEU:HA	3:F:630:LEU:HD11	1.90	0.52
3:D:371:PRO:HB3	3:D:459:LEU:HD11	1.90	0.52
3:D:476:TYR:CD1	3:D:506:ALA:HB3	2.45	0.52
3:D:518:ALA:CB	3:D:580:PRO:HG3	2.39	0.52
2:E:9:ASN:HB3	3:F:209:ASN:CG	2.35	0.52
3:F:58:ILE:HG13	3:F:127:LEU:HD23	1.91	0.52
3:F:96:PHE:CD1	3:F:159:TRP:HH2	2.28	0.52
3:F:609:ARG:HD2	2:G:29:THR:HG22	1.91	0.52
3:J:321:THR:HG23	3:J:324:GLY:O	2.09	0.52
3:J:476:TYR:CD1	3:J:506:ALA:HB3	2.44	0.52
2:K:9:ASN:HB3	3:L:209:ASN:CG	2.35	0.52
3:L:342:ASN:OD1	3:L:431:ILE:HG21	2.10	0.52
3:L:496:GLU:C	3:L:498:VAL:H	2.17	0.52
3:N:342:ASN:OD1	3:N:431:ILE:HG21	2.10	0.52
3:P:485:VAL:O	3:P:485:VAL:HG12	2.09	0.52
3:R:50:PRO:HG3	3:R:56:ALA:HB2	1.91	0.52
3:R:93:GLN:N	3:R:94:PRO:CD	2.73	0.52
3:R:269:LYS:O	3:R:270:ASP:HB2	2.09	0.52
3:R:411:VAL:O	3:R:412:ARG:CB	2.56	0.52
3:T:49:ARG:NH1	3:T:115:THR:HG22	2.18	0.52
3:T:249:VAL:O	3:T:250:GLY:C	2.53	0.52
3:T:342:ASN:OD1	3:T:431:ILE:HG21	2.10	0.52
5:W:591:ASP:O	5:W:591:ASP:CG	2.53	0.52
5:W:607:LEU:CD2	5:W:638:HIS:HB3	2.40	0.52
5:W:940:VAL:O	5:W:940:VAL:HG12	2.10	0.52
6:X:241:ASN:HB2	6:X:1148:SER:HG	1.75	0.52
6:X:249:TRP:CZ3	6:X:250:ALA:HB2	2.45	0.52
6:X:1123:GLN:HG3	6:X:1127:CYS:SG	2.49	0.52
6:Y:317:PHE:N	6:Y:317:PHE:CD1	2.77	0.52
6:Y:560:LEU:HD11	6:Y:721:VAL:HG23	1.91	0.52
6:Y:1019:THR:H	6:Y:1032:ILE:HG22	1.74	0.52
6:Y:1169:TRP:N	6:Y:1169:TRP:CD1	2.77	0.52

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:b:45:ARG:CD	3:B:400:ALA:C	2.83	0.52
1:t:45:ARG:N	3:T:399:ALA:O	2.42	0.52
3:B:141:GLN:C	3:B:143:MET:N	2.63	0.52
3:B:419:MET:O	3:B:419:MET:HE3	2.08	0.52
3:B:496:GLU:C	3:B:498:VAL:H	2.17	0.52
3:F:266:MET:HE3	3:F:605:GLY:HA2	1.91	0.52
3:F:321:THR:HG23	3:F:324:GLY:O	2.09	0.52
3:F:360:TYR:CD1	3:F:361:ASP:N	2.77	0.52
3:J:86:LEU:HD12	3:J:108:LEU:HB2	1.92	0.52
3:J:195:PRO:HD3	3:N:566:TRP:NE1	2.24	0.52
3:J:249:VAL:O	3:J:250:GLY:C	2.53	0.52
3:L:144:ASN:C	3:L:146:GLN:N	2.65	0.52
3:L:494:THR:HG23	3:L:497:GLU:OE1	2.08	0.52
3:L:549:LEU:HB3	3:N:644:LEU:CD1	2.34	0.52
3:N:58:ILE:HG13	3:N:127:LEU:HD23	1.91	0.52
3:N:86:LEU:HD12	3:N:108:LEU:HB2	1.92	0.52
3:N:93:GLN:N	3:N:94:PRO:CD	2.73	0.52
2:O:24:SER:HB3	3:P:176:ALA:N	2.24	0.52
3:P:141:GLN:C	3:P:143:MET:H	2.16	0.52
3:P:196:LEU:H	3:P:196:LEU:HD22	1.73	0.52
3:P:411:VAL:O	3:P:412:ARG:CB	2.56	0.52
3:R:238:VAL:HG11	3:T:82:PHE:CZ	2.45	0.52
3:R:496:GLU:C	3:R:498:VAL:H	2.17	0.52
3:R:545:VAL:CG2	3:T:641:ILE:HD11	2.40	0.52
3:T:431:ILE:CG1	3:T:437:ILE:HG21	2.38	0.52
3:T:484:ILE:HG22	3:T:485:VAL:H	1.73	0.52
4:U:71:LEU:O	4:U:71:LEU:HD23	2.10	0.52
5:W:226:VAL:CG1	5:W:239:LEU:HD22	2.40	0.52
5:W:1204:LEU:HB3	5:W:1227:TYR:HD2	1.70	0.52
6:X:259:LEU:O	6:X:259:LEU:HD12	2.10	0.52
6:X:349:VAL:C	6:X:350:LEU:HD12	2.34	0.52
6:X:930:ARG:HH22	6:Y:692:THR:HG21	1.73	0.52
6:X:1013:VAL:CG1	6:X:1014:VAL:N	2.72	0.52
6:X:1108:ASN:C	6:X:1110:ALA:N	2.65	0.52
6:Y:416:ILE:HG21	6:Y:1204:LEU:CD1	2.40	0.52
6:Y:448:SER:HB2	6:Y:664:GLY:CA	2.39	0.52
6:Y:497:ILE:HG23	6:Y:498:THR:N	2.25	0.52
1:l:5:MET:O	1:l:6:ILE:HB	2.09	0.52
1:b:73:ARG:HH12	1:b:74:GLN:CD	2.11	0.52
1:d:22:LEU:O	3:D:323:SER:HB2	2.10	0.52
1:j:5:MET:HE2	1:j:5:MET:N	2.25	0.52

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:j:45:ARG:N	3:J:399:ALA:O	2.42	0.52
3:B:95:LEU:HD21	3:B:207:PHE:CZ	2.44	0.52
3:B:249:VAL:O	3:B:250:GLY:C	2.53	0.52
3:B:518:ALA:CB	3:B:580:PRO:HG3	2.39	0.52
3:D:371:PRO:HB2	3:D:459:LEU:CD1	2.39	0.52
2:E:24:SER:HB3	3:F:176:ALA:N	2.24	0.52
3:F:566:TRP:NE1	3:H:195:PRO:HD3	2.24	0.52
2:G:28:SER:HB2	3:H:214:ARG:O	2.10	0.52
3:H:371:PRO:HB2	3:H:459:LEU:CD1	2.39	0.52
3:L:321:THR:HG23	3:L:324:GLY:O	2.09	0.52
3:L:345:ILE:O	3:L:346:ASP:C	2.52	0.52
3:N:491:ALA:HB2	3:N:592:THR:CB	2.32	0.52
2:O:9:ASN:HB3	3:P:209:ASN:CG	2.35	0.52
3:P:60:SER:HA	5:W:379:VAL:HB	1.92	0.52
3:P:86:LEU:HD12	3:P:108:LEU:HB2	1.92	0.52
3:P:93:GLN:N	3:P:94:PRO:CD	2.73	0.52
3:R:476:TYR:CD1	3:R:506:ALA:HB3	2.44	0.52
3:T:312:ASN:CG	3:T:316:ILE:HD11	2.34	0.52
4:U:16:GLN:HG3	4:U:17:PRO:CD	2.38	0.52
4:U:369:HIS:O	4:U:373:MET:HG2	2.09	0.52
4:V:54:TRP:HZ2	6:X:1037:VAL:HG11	1.75	0.52
5:W:13:LEU:H	5:W:385:THR:HA	1.75	0.52
5:W:31:LEU:HD11	5:W:96:VAL:HG13	1.91	0.52
5:W:113:THR:CB	5:W:118:LEU:HD12	2.39	0.52
5:W:631:VAL:HG13	5:W:632:PHE:N	2.25	0.52
5:W:649:ILE:O	5:W:1010:LEU:HD11	2.10	0.52
5:W:662:ARG:O	5:W:662:ARG:HG3	2.09	0.52
6:X:354:THR:CG2	6:X:1151:ALA:O	2.58	0.52
6:X:898:ALA:HB1	6:X:929:PRO:HB2	1.92	0.52
6:X:935:SER:HA	6:X:949:HIS:HA	1.91	0.52
6:Y:278:LEU:HD13	6:Y:891:PHE:HZ	1.75	0.52
6:Y:798:VAL:HG12	6:Y:799:ASP:H	1.75	0.52
6:Y:835:HIS:CE1	6:Y:1213:ALA:H	2.27	0.52
6:Y:1100:LEU:O	6:Y:1103:TRP:N	2.39	0.52
1:n:14:VAL:O	1:n:18:ALA:N	2.42	0.52
1:t:48:ILE:HG13	1:t:55:THR:HA	1.92	0.52
3:B:196:LEU:HD22	3:B:196:LEU:H	1.73	0.52
3:D:368:THR:CG2	3:D:446:PRO:HG3	2.38	0.52
3:F:371:PRO:HB3	3:F:459:LEU:HD11	1.90	0.52
3:H:96:PHE:CD1	3:H:159:TRP:HH2	2.28	0.52
3:H:266:MET:HE3	3:H:605:GLY:HA2	1.91	0.52

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:H:344:MET:HE2	3:H:502:LEU:CD1	2.37	0.52
3:H:411:VAL:O	3:H:412:ARG:CB	2.57	0.52
3:H:476:TYR:CD1	3:H:506:ALA:HB3	2.44	0.52
3:J:345:ILE:O	3:J:346:ASP:C	2.52	0.52
3:L:50:PRO:HG3	3:L:56:ALA:HB2	1.91	0.52
3:L:141:GLN:C	3:L:143:MET:H	2.16	0.52
3:N:249:VAL:O	3:N:250:GLY:C	2.53	0.52
2:Q:28:SER:HB2	3:R:214:ARG:O	2.10	0.52
3:R:321:THR:HG23	3:R:324:GLY:O	2.09	0.52
2:S:28:SER:HB2	3:T:214:ARG:O	2.10	0.52
3:T:95:LEU:HD21	3:T:207:PHE:CZ	2.44	0.52
3:T:96:PHE:CD1	3:T:159:TRP:HH2	2.28	0.52
3:T:266:MET:HE3	3:T:605:GLY:HA2	1.91	0.52
4:U:40:ARG:HD2	4:U:141:ASP:OD1	2.09	0.52
4:V:78:PRO:C	4:V:80:ALA:H	2.18	0.52
5:W:47:LEU:HD11	5:W:327:ILE:HG21	1.92	0.52
5:W:558:VAL:CG2	5:W:561:GLY:H	2.23	0.52
5:W:1273:LEU:N	5:W:1273:LEU:HD22	2.24	0.52
6:X:675:MET:HE2	6:X:768:GLN:HE22	1.75	0.52
6:X:792:GLN:HB2	6:X:931:ILE:HG21	1.91	0.52
6:X:828:GLU:HA	6:X:828:GLU:OE1	2.10	0.52
6:X:880:LEU:H	6:X:880:LEU:HD12	1.74	0.52
6:X:931:ILE:CD1	6:Y:689:VAL:CG1	2.85	0.52
6:X:1149:THR:HG21	6:X:1162:ALA:N	2.25	0.52
6:Y:160:PHE:HD1	6:Y:1206:GLY:O	1.93	0.52
6:Y:322:THR:CG2	6:Y:348:ALA:HB2	2.38	0.52
1:b:45:ARG:CB	3:B:399:ALA:O	2.58	0.52
1:f:48:ILE:HG13	1:f:55:THR:HA	1.92	0.52
1:j:58:PRO:HG3	3:J:423:GLN:HE21	1.61	0.52
1:t:60:ALA:CB	3:T:322:TYR:OH	2.41	0.52
2:A:9:ASN:HB3	3:B:209:ASN:CG	2.35	0.52
3:B:285:PHE:CE2	3:B:484:ILE:HD12	2.45	0.52
3:B:494:THR:O	3:B:495:SER:C	2.52	0.52
3:D:64:VAL:O	3:D:64:VAL:HG13	2.10	0.52
3:D:321:THR:HG23	3:D:324:GLY:O	2.09	0.52
3:D:484:ILE:HG22	3:D:485:VAL:H	1.73	0.52
3:F:285:PHE:CE2	3:F:484:ILE:HD12	2.45	0.52
3:H:86:LEU:HD12	3:H:108:LEU:HB2	1.92	0.52
3:H:582:LEU:C	3:H:584:LEU:H	2.18	0.52
3:L:296:PRO:C	3:L:298:PHE:N	2.68	0.52
2:M:28:SER:HB2	3:N:214:ARG:O	2.10	0.52

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:N:296:PRO:O	3:N:297:ALA:HB3	2.08	0.52
3:N:357:ILE:HD11	3:N:414:GLN:HB3	1.91	0.52
3:P:237:GLU:CD	3:R:46:LYS:NZ	2.68	0.52
3:P:322:TYR:CE1	3:P:396:VAL:HB	2.45	0.52
3:R:367:SER:O	3:R:368:THR:HB	2.09	0.52
3:T:99:THR:OG1	3:T:101:THR:CG2	2.58	0.52
4:U:27:GLN:O	6:X:882:PRO:HB3	2.08	0.52
4:V:196:LEU:HD12	4:V:197:PRO:HD2	1.92	0.52
5:W:148:ILE:HG13	5:W:151:ILE:HD11	1.92	0.52
5:W:213:ARG:HE	5:W:217:LEU:HD21	1.75	0.52
5:W:453:PRO:HB2	5:W:456:GLN:HG2	1.92	0.52
5:W:657:LEU:HD12	5:W:657:LEU:N	2.23	0.52
5:W:1087:LEU:HD21	5:W:1096:PHE:CB	2.40	0.52
5:W:1185:THR:HG22	5:W:1190:ARG:HG2	1.92	0.52
6:X:234:PRO:HG3	6:X:348:ALA:HB2	1.92	0.52
6:X:354:THR:HG21	6:X:1152:ALA:HA	1.91	0.52
6:X:1106:LYS:O	6:X:1113:PRO:HD3	2.10	0.52
6:Y:607:HIS:O	6:Y:609:THR:N	2.42	0.52
6:Y:969:MET:O	6:Y:979:ARG:NH2	2.42	0.52
1:h:15:ARG:NH2	3:H:402:ALA:HB2	2.24	0.51
1:j:8:GLN:HE22	3:L:303:ASP:HA	1.75	0.51
1:p:45:ARG:CG	3:P:400:ALA:C	2.59	0.51
1:p:48:ILE:HG13	1:p:55:THR:HA	1.92	0.51
1:t:47:THR:OG1	1:t:57:ALA:N	2.40	0.51
3:B:49:ARG:NH1	3:B:115:THR:HG22	2.18	0.51
3:B:521:VAL:HG23	3:B:522:GLY:H	1.74	0.51
3:B:582:LEU:C	3:B:584:LEU:H	2.19	0.51
3:D:287:PRO:HG2	3:D:471:LEU:CD1	2.39	0.51
3:D:443:PHE:HB2	3:D:457:PRO:CB	2.24	0.51
3:D:494:THR:O	3:D:495:SER:C	2.52	0.51
3:F:342:ASN:OD1	3:F:431:ILE:HG21	2.10	0.51
3:F:411:VAL:O	3:F:412:ARG:CB	2.56	0.51
3:H:269:LYS:O	3:H:270:ASP:HB2	2.10	0.51
3:H:431:ILE:CG1	3:H:437:ILE:HG21	2.38	0.51
3:H:485:VAL:O	3:H:485:VAL:HG12	2.10	0.51
3:J:367:SER:O	3:J:368:THR:HB	2.08	0.51
3:J:521:VAL:HG23	3:J:522:GLY:H	1.74	0.51
3:L:249:VAL:O	3:L:250:GLY:C	2.53	0.51
3:N:285:PHE:CE2	3:N:484:ILE:HD12	2.45	0.51
3:N:431:ILE:CG1	3:N:437:ILE:HG21	2.38	0.51
3:N:540:VAL:CG2	3:N:569:ALA:HB3	2.40	0.51

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:O:9:ASN:HB2	3:P:209:ASN:CG	2.35	0.51
3:P:95:LEU:HD21	3:P:207:PHE:CZ	2.44	0.51
3:P:476:TYR:CD1	3:P:506:ALA:HB3	2.45	0.51
3:R:86:LEU:HD12	3:R:108:LEU:HB2	1.92	0.51
3:R:141:GLN:C	3:R:143:MET:H	2.16	0.51
3:R:158:LEU:N	3:R:158:LEU:HD23	2.23	0.51
3:T:284:VAL:HG12	3:T:284:VAL:O	2.10	0.51
3:T:357:ILE:HG12	3:T:416:ARG:NH1	2.26	0.51
4:U:10:THR:HG23	4:U:73:GLY:HA3	1.92	0.51
4:V:66:ILE:HD11	4:V:387:GLN:NE2	2.25	0.51
5:W:184:ASP:OD2	5:W:186:SER:HB3	2.09	0.51
5:W:672:THR:HG22	5:W:1081:VAL:HG22	1.91	0.51
5:W:971:VAL:HG13	5:W:972:PRO:CD	2.14	0.51
6:X:626:LEU:O	6:X:628:PRO:HD3	2.11	0.51
6:X:644:GLU:HA	6:X:644:GLU:OE2	2.10	0.51
6:X:883:ARG:HH11	6:X:883:ARG:HB3	1.74	0.51
6:X:979:ARG:HG3	6:X:1140:GLU:HG3	1.91	0.51
6:Y:238:GLY:O	6:Y:239:ALA:C	2.53	0.51
6:Y:484:THR:HA	6:Y:754:GLU:OE2	2.10	0.51
6:Y:685:PRO:HD3	6:Y:749:HIS:ND1	2.25	0.51
6:Y:990:ASN:HB2	6:Y:1131:ARG:HB2	1.92	0.51
1:d:45:ARG:N	3:D:399:ALA:O	2.43	0.51
1:h:23:THR:CG2	1:h:42:THR:OG1	2.59	0.51
1:h:48:ILE:HG13	1:h:55:THR:HA	1.92	0.51
1:n:44:GLY:C	3:N:399:ALA:HB3	2.34	0.51
1:n:48:ILE:HG13	1:n:55:THR:HA	1.92	0.51
1:p:8:GLN:HG2	3:R:302:GLU:HG3	0.55	0.51
1:r:48:ILE:HG13	1:r:55:THR:HA	1.92	0.51
1:r:59:HIS:O	1:r:59:HIS:HD2	1.92	0.51
3:B:96:PHE:CD1	3:B:159:TRP:HH2	2.28	0.51
2:C:24:SER:HB3	3:D:176:ALA:N	2.24	0.51
2:C:28:SER:HB2	3:D:214:ARG:O	2.10	0.51
3:D:269:LYS:O	3:D:270:ASP:HB2	2.09	0.51
3:D:431:ILE:CG1	3:D:437:ILE:HG21	2.38	0.51
3:D:521:VAL:HG23	3:D:522:GLY:H	1.74	0.51
3:F:93:GLN:N	3:F:94:PRO:CD	2.73	0.51
3:F:237:GLU:CB	3:H:46:LYS:NZ	2.67	0.51
3:F:269:LYS:O	3:F:270:ASP:HB2	2.09	0.51
3:H:249:VAL:O	3:H:250:GLY:C	2.52	0.51
3:H:285:PHE:CE2	3:H:484:ILE:HD12	2.45	0.51
3:J:92:TRP:CZ2	3:J:160:GLN:HB2	2.46	0.51

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:J:94:PRO:HG2	3:J:210:VAL:CG2	2.41	0.51
3:J:287:PRO:HG2	3:J:471:LEU:CD1	2.39	0.51
3:J:641:ILE:HD11	3:N:545:VAL:CG2	2.40	0.51
3:L:86:LEU:HD12	3:L:108:LEU:HB2	1.92	0.51
3:N:183:TRP:CZ3	3:N:186:LEU:HD23	2.46	0.51
2:O:29:THR:HG22	3:T:609:ARG:HD2	1.91	0.51
3:P:59:ASP:OD1	3:P:60:SER:N	2.43	0.51
3:P:298:PHE:CZ	3:T:416:ARG:HG3	2.45	0.51
3:P:472:ARG:NH1	3:R:530:ARG:CD	2.68	0.51
2:Q:9:ASN:HB3	3:R:209:ASN:CG	2.35	0.51
3:R:285:PHE:CE2	3:R:484:ILE:HD12	2.45	0.51
2:S:24:SER:HB3	3:T:176:ALA:N	2.24	0.51
3:T:50:PRO:HG3	3:T:56:ALA:HB2	1.91	0.51
3:T:431:ILE:CD1	3:T:437:ILE:HG23	2.39	0.51
4:U:94:ASP:OD1	6:X:949:HIS:NE2	2.36	0.51
4:U:106:ILE:HD12	4:U:107:GLN:H	1.74	0.51
5:W:248:LEU:CD1	6:X:628:PRO:HD2	2.39	0.51
5:W:751:SER:C	5:W:753:GLY:H	2.17	0.51
5:W:830:HIS:HE1	5:W:890:CYS:SG	2.32	0.51
5:W:843:LEU:HG	5:W:854:MET:SD	2.49	0.51
6:X:1060:ARG:HH12	6:X:1105:SER:HA	1.76	0.51
6:X:1125:TYR:CZ	6:X:1127:CYS:HB3	2.45	0.51
6:Y:440:ILE:CG1	6:Y:1202:PRO:HG2	2.36	0.51
6:Y:493:LEU:O	6:Y:497:ILE:HG22	2.09	0.51
6:Y:718:THR:HG23	6:Y:719:ALA:N	2.24	0.51
6:Y:919:LEU:HD22	6:Y:920:LEU:HB2	1.92	0.51
1:b:45:ARG:HD2	3:B:401:GLY:N	2.25	0.51
1:j:73:ARG:CZ	1:j:73:ARG:HB2	2.40	0.51
2:A:28:SER:HB2	3:B:214:ARG:O	2.10	0.51
3:B:287:PRO:HG2	3:B:471:LEU:CD1	2.39	0.51
3:D:47:LEU:C	3:D:48:TRP:HD1	2.17	0.51
3:D:183:TRP:CZ3	3:D:186:LEU:HD23	2.45	0.51
3:D:298:PHE:CZ	3:H:416:ARG:HG3	2.46	0.51
3:F:357:ILE:HG12	3:F:416:ARG:NH1	2.26	0.51
3:F:582:LEU:C	3:F:584:LEU:H	2.18	0.51
3:H:357:ILE:HG12	3:H:416:ARG:NH1	2.26	0.51
2:I:32:PRO:HD2	2:K:38:PRO:HB2	1.93	0.51
3:J:93:GLN:N	3:J:94:PRO:CD	2.73	0.51
3:J:296:PRO:C	3:J:298:PHE:N	2.68	0.51
3:J:422:LEU:O	3:J:423:GLN:HB3	2.08	0.51
3:L:96:PHE:CD1	3:L:159:TRP:HH2	2.28	0.51

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:L:238:VAL:HG11	3:N:82:PHE:CZ	2.45	0.51
3:L:484:ILE:HG22	3:L:485:VAL:H	1.73	0.51
3:N:357:ILE:HG12	3:N:416:ARG:NH1	2.26	0.51
3:P:96:PHE:CD1	3:P:159:TRP:HH2	2.28	0.51
3:P:183:TRP:CZ3	3:P:186:LEU:HD23	2.45	0.51
3:P:249:VAL:O	3:P:250:GLY:C	2.53	0.51
3:P:566:TRP:NE1	3:R:195:PRO:HD3	2.24	0.51
3:P:641:ILE:HD11	3:T:545:VAL:CG2	2.40	0.51
2:Q:9:ASN:HB2	3:R:209:ASN:CG	2.35	0.51
3:R:431:ILE:CG1	3:R:437:ILE:HG21	2.38	0.51
3:T:496:GLU:C	3:T:498:VAL:H	2.17	0.51
4:U:96:VAL:HG12	4:U:96:VAL:O	2.11	0.51
4:V:184:HIS:CE1	6:Y:436:LEU:CD1	2.91	0.51
5:W:407:ALA:O	5:W:779:MET:HG2	2.10	0.51
5:W:425:PRO:HA	5:W:702:ASP:O	2.09	0.51
5:W:480:ARG:HE	5:W:649:ILE:HG22	1.73	0.51
5:W:489:LYS:HG3	5:W:649:ILE:HG13	1.91	0.51
5:W:1136:LEU:HD11	5:W:1221:PHE:HZ	1.75	0.51
6:X:403:ILE:HG12	6:X:1197:TYR:CE1	2.46	0.51
6:X:651:PRO:HG3	6:X:679:TRP:CZ3	2.45	0.51
6:X:1168:ARG:O	6:X:1189:THR:HA	2.11	0.51
6:Y:353:GLN:NE2	6:Y:358:GLN:HG2	2.25	0.51
1:l:48:ILE:HG13	1:l:55:THR:HA	1.92	0.51
1:b:15:ARG:NE	3:B:406:ASN:CG	2.69	0.51
1:b:21:ARG:HH11	3:B:327:TYR:HH	1.56	0.51
1:b:48:ILE:HG13	1:b:55:THR:HA	1.92	0.51
1:j:48:ILE:HG13	1:j:55:THR:HA	1.92	0.51
1:n:60:ALA:O	1:n:61:ASN:C	2.53	0.51
3:B:86:LEU:HD12	3:B:108:LEU:HB2	1.92	0.51
3:B:144:ASN:C	3:B:146:GLN:N	2.65	0.51
3:D:114:THR:HG22	3:D:114:THR:O	2.10	0.51
3:D:566:TRP:NE1	3:F:195:PRO:HD3	2.24	0.51
3:F:431:ILE:CD1	3:F:437:ILE:HG23	2.39	0.51
2:G:24:SER:HB3	3:H:176:ALA:N	2.24	0.51
3:H:296:PRO:C	3:H:298:PHE:H	2.19	0.51
2:I:28:SER:HB2	3:J:214:ARG:O	2.10	0.51
3:J:357:ILE:HG12	3:J:416:ARG:NH1	2.26	0.51
3:J:416:ARG:HG3	3:L:298:PHE:CZ	2.46	0.51
3:L:44:THR:HG22	3:L:45:GLY:H	1.72	0.51
3:L:92:TRP:CZ2	3:L:160:GLN:HB2	2.46	0.51
3:L:93:GLN:N	3:L:94:PRO:CD	2.73	0.51

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:L:284:VAL:HG12	3:L:284:VAL:O	2.11	0.51
3:L:582:LEU:C	3:L:584:LEU:H	2.18	0.51
3:N:345:ILE:O	3:N:346:ASP:C	2.52	0.51
3:N:443:PHE:HE2	3:N:459:LEU:HD21	1.75	0.51
3:P:92:TRP:CZ2	3:P:160:GLN:HB2	2.46	0.51
3:P:494:THR:H	3:P:497:GLU:CB	2.19	0.51
3:R:49:ARG:NH1	3:R:115:THR:HG22	2.18	0.51
3:R:238:VAL:HG21	3:T:78:MET:HE2	1.83	0.51
3:T:540:VAL:CG2	3:T:569:ALA:HB3	2.40	0.51
4:U:123:ASN:C	4:U:125:TYR:H	2.19	0.51
4:U:313:ALA:O	4:U:314:ASN:HB2	2.10	0.51
4:V:195:GLN:HE22	6:Y:377:ILE:CA	2.24	0.51
4:V:282:ALA:O	4:V:286:MET:HB2	2.09	0.51
5:W:160:ILE:O	5:W:161:ASP:HB2	2.10	0.51
5:W:173:ALA:CB	5:W:202:ALA:HB3	2.41	0.51
5:W:426:LEU:O	5:W:702:ASP:CB	2.58	0.51
5:W:448:ASP:OD1	5:W:673:ALA:HB2	2.10	0.51
5:W:557:ASP:O	5:W:559:VAL:HG23	2.11	0.51
5:W:829:ALA:O	5:W:850:LEU:HG	2.11	0.51
6:X:670:ILE:HG22	6:X:674:ARG:HB2	1.90	0.51
6:X:682:GLN:HB3	6:X:749:HIS:ND1	2.25	0.51
6:Y:203:SER:CB	6:Y:247:THR:HA	2.39	0.51
6:Y:354:THR:O	6:Y:357:ALA:N	2.43	0.51
6:Y:386:GLU:HG2	6:Y:1195:TYR:CE2	2.44	0.51
6:Y:1034:ASP:C	6:Y:1036:ASN:N	2.68	0.51
1:d:22:LEU:HD11	1:d:83:VAL:CB	2.37	0.51
1:h:8:GLN:CD	3:D:302:GLU:CG	2.63	0.51
1:h:23:THR:HB	3:H:323:SER:C	2.28	0.51
1:h:58:PRO:CG	3:H:423:GLN:CG	2.88	0.51
1:n:8:GLN:HE21	3:J:302:GLU:CG	2.18	0.51
1:r:22:LEU:CD2	1:r:22:LEU:N	2.73	0.51
2:C:32:PRO:HD2	2:E:38:PRO:HB2	1.92	0.51
3:D:195:PRO:HD3	3:H:566:TRP:NE1	2.24	0.51
3:D:360:TYR:CD1	3:D:361:ASP:N	2.77	0.51
3:D:540:VAL:CG2	3:D:569:ALA:HB3	2.41	0.51
3:F:416:ARG:HG3	3:H:298:PHE:CZ	2.46	0.51
3:H:50:PRO:HG3	3:H:56:ALA:HB2	1.91	0.51
3:H:92:TRP:CE2	3:H:156:VAL:CG2	2.94	0.51
3:H:94:PRO:HG2	3:H:210:VAL:CG2	2.41	0.51
3:J:237:GLU:CD	3:L:46:LYS:NZ	2.68	0.51
3:J:496:GLU:C	3:J:498:VAL:H	2.17	0.51

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:L:368:THR:CG2	3:L:446:PRO:HG3	2.38	0.51
3:N:92:TRP:CZ2	3:N:160:GLN:HB2	2.46	0.51
3:N:96:PHE:CD1	3:N:159:TRP:HH2	2.28	0.51
3:N:333:PRO:HG3	3:N:454:PHE:CA	2.41	0.51
3:N:476:TYR:CD1	3:N:506:ALA:HB3	2.45	0.51
3:R:431:ILE:CD1	3:R:437:ILE:HG23	2.39	0.51
3:T:92:TRP:CZ2	3:T:160:GLN:HB2	2.46	0.51
3:T:94:PRO:HG2	3:T:210:VAL:CG2	2.41	0.51
3:T:494:THR:O	3:T:495:SER:C	2.52	0.51
3:T:582:LEU:C	3:T:584:LEU:H	2.18	0.51
4:U:375:VAL:O	4:U:375:VAL:HG12	2.11	0.51
5:W:79:LEU:HD12	5:W:84:TRP:HA	1.93	0.51
5:W:1103:THR:C	5:W:1104:LEU:HD23	2.35	0.51
5:W:1219:ILE:HG23	5:W:1219:ILE:O	2.11	0.51
5:W:1242:PRO:HB2	5:W:1243:TYR:CD1	2.45	0.51
6:X:302:LEU:CD1	6:X:905:VAL:HG22	2.41	0.51
6:X:518:LEU:HB2	6:X:568:PHE:CE2	2.46	0.51
6:Y:248:ASN:HB3	6:Y:251:SER:HB2	1.93	0.51
6:Y:275:LYS:NZ	6:Y:285:ASN:HA	2.26	0.51
6:Y:480:LEU:HD22	6:Y:529:TRP:HZ3	1.76	0.51
6:Y:593:LEU:HD12	6:Y:597:GLU:CG	2.40	0.51
6:Y:601:ILE:HG22	6:Y:603:VAL:HG12	1.91	0.51
6:Y:977:VAL:HA	6:Y:1081:TYR:O	2.10	0.51
6:Y:986:GLN:HG3	6:Y:1135:TYR:CE1	2.45	0.51
1:l:22:LEU:CD2	1:l:22:LEU:N	2.72	0.51
1:d:15:ARG:NE	3:D:406:ASN:CG	2.68	0.51
1:d:48:ILE:HG13	1:d:55:THR:HA	1.92	0.51
1:j:44:GLY:C	3:J:399:ALA:HB3	2.36	0.51
3:B:92:TRP:CE2	3:B:156:VAL:CG2	2.94	0.51
3:B:166:ALA:HA	3:B:194:LEU:HD21	1.93	0.51
3:B:269:LYS:O	3:B:270:ASP:HB2	2.09	0.51
3:B:333:PRO:HG3	3:B:454:PHE:CA	2.41	0.51
3:B:342:ASN:OD1	3:B:431:ILE:HG21	2.10	0.51
3:B:357:ILE:HG12	3:B:416:ARG:NH1	2.26	0.51
2:C:9:ASN:HB3	3:D:209:ASN:CG	2.35	0.51
3:D:342:ASN:OD1	3:D:431:ILE:HG21	2.10	0.51
3:D:545:VAL:CG2	3:F:641:ILE:HD11	2.40	0.51
2:E:28:SER:HB2	3:F:214:ARG:O	2.10	0.51
3:F:92:TRP:CZ2	3:F:160:GLN:HB2	2.46	0.51
3:F:296:PRO:C	3:F:298:PHE:H	2.19	0.51
3:F:540:VAL:CG2	3:F:569:ALA:HB3	2.41	0.51

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:H:540:VAL:CG2	3:H:569:ALA:HB3	2.40	0.51
2:I:9:ASN:HB2	3:J:209:ASN:CG	2.35	0.51
3:J:82:PHE:CZ	3:N:238:VAL:HG11	2.45	0.51
3:J:183:TRP:NE1	3:J:187:LEU:HD12	2.26	0.51
3:J:566:TRP:NE1	3:L:195:PRO:HD3	2.24	0.51
2:K:28:SER:HB2	3:L:214:ARG:O	2.10	0.51
3:L:92:TRP:CE2	3:L:156:VAL:CG2	2.94	0.51
3:N:496:GLU:C	3:N:498:VAL:H	2.17	0.51
3:P:455:SER:O	3:T:410:ASN:ND2	2.41	0.51
3:P:496:GLU:C	3:P:498:VAL:H	2.18	0.51
3:R:416:ARG:HG3	3:T:298:PHE:CZ	2.46	0.51
3:R:494:THR:O	3:R:495:SER:C	2.52	0.51
4:U:128:TYR:HA	4:U:135:MET:SD	2.51	0.51
4:U:136:TRP:CE2	4:U:197:PRO:HA	2.46	0.51
5:W:49:PHE:CZ	5:W:54:ARG:HA	2.45	0.51
5:W:66:ASN:HD22	5:W:176:LYS:HE3	1.76	0.51
5:W:246:ASP:HB2	5:W:255:ILE:O	2.11	0.51
5:W:955:VAL:HG21	5:W:964:VAL:CG1	2.41	0.51
5:W:1196:LEU:H	5:W:1196:LEU:HD22	1.76	0.51
6:X:355:LYS:HE3	6:X:1149:THR:CG2	2.41	0.51
6:X:432:ILE:HD13	6:X:432:ILE:C	2.35	0.51
6:X:611:PRO:O	6:X:612:ALA:C	2.52	0.51
6:X:673:ASN:C	6:X:674:ARG:HD2	2.34	0.51
6:X:758:LEU:O	6:X:762:THR:HG23	2.11	0.51
6:X:782:ASN:HB2	6:X:784:LEU:HD11	1.93	0.51
6:X:817:ALA:O	6:X:818:ILE:C	2.53	0.51
6:X:1124:VAL:HG23	6:X:1125:TYR:N	2.26	0.51
6:Y:210:ILE:O	6:Y:239:ALA:HB3	2.10	0.51
6:Y:237:GLU:HA	6:Y:1151:ALA:HB3	1.91	0.51
6:Y:543:THR:HG22	6:Y:577:SER:N	2.21	0.51
6:Y:1054:VAL:O	6:Y:1055:SER:HB3	2.11	0.51
6:Y:1146:LEU:HD11	6:Y:1149:THR:HG22	1.91	0.51
1:b:61:ASN:CB	3:B:395:THR:CG2	2.85	0.51
3:D:96:PHE:CD1	3:D:159:TRP:HH2	2.28	0.51
3:D:285:PHE:CE2	3:D:484:ILE:HD12	2.45	0.51
3:D:416:ARG:HG3	3:F:298:PHE:CZ	2.46	0.51
3:F:494:THR:H	3:F:497:GLU:CB	2.19	0.51
3:F:521:VAL:HG23	3:F:522:GLY:H	1.74	0.51
3:H:342:ASN:OD1	3:H:431:ILE:HG21	2.10	0.51
3:J:196:LEU:HD22	3:J:196:LEU:H	1.73	0.51
3:J:238:VAL:HG11	3:L:82:PHE:CZ	2.45	0.51

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:J:475:THR:O	3:J:505:ASP:HB2	2.11	0.51
3:L:166:ALA:HA	3:L:194:LEU:HD21	1.93	0.51
3:L:238:VAL:HG21	3:N:78:MET:HE2	1.83	0.51
3:L:287:PRO:HG2	3:L:471:LEU:CD1	2.39	0.51
3:L:545:VAL:CG2	3:N:641:ILE:HD11	2.40	0.51
3:P:416:ARG:HG3	3:R:298:PHE:CZ	2.46	0.51
3:R:183:TRP:NE1	3:R:187:LEU:HD12	2.26	0.51
3:R:237:GLU:OE1	3:T:46:LYS:NZ	2.38	0.51
3:R:284:VAL:O	3:R:284:VAL:HG12	2.10	0.51
3:T:183:TRP:CZ3	3:T:186:LEU:HD23	2.45	0.51
3:T:287:PRO:HG2	3:T:471:LEU:CD1	2.39	0.51
4:U:4:ARG:HH12	5:W:217:LEU:HG	1.75	0.51
4:U:27:GLN:O	6:X:882:PRO:CD	2.55	0.51
4:U:89:GLY:O	4:U:90:ILE:HD13	2.10	0.51
4:V:85:THR:HA	4:V:110:PRO:O	2.09	0.51
4:V:184:HIS:CE1	6:Y:436:LEU:CD2	2.93	0.51
5:W:38:ASN:ND2	5:W:40:HIS:HB3	2.26	0.51
5:W:68:TRP:CD2	5:W:126:LEU:HD12	2.46	0.51
5:W:359:TYR:OH	5:W:375:PRO:HB3	2.10	0.51
5:W:800:PHE:CD1	5:W:1005:GLY:HA2	2.45	0.51
5:W:821:GLU:HG3	5:W:824:ARG:HH21	1.76	0.51
5:W:1161:TYR:O	5:W:1190:ARG:NH1	2.44	0.51
5:W:1180:LYS:HG2	5:W:1280:VAL:HG22	1.93	0.51
5:W:1196:LEU:HD22	5:W:1196:LEU:N	2.25	0.51
5:W:1243:TYR:HE1	5:W:1298:ILE:HD11	1.72	0.51
6:X:252:LEU:HD12	6:X:252:LEU:C	2.34	0.51
6:X:671:PRO:HB2	6:X:674:ARG:CG	2.39	0.51
6:X:746:TYR:OH	6:X:754:GLU:OE2	2.22	0.51
6:X:821:ALA:O	6:X:824:CYS:N	2.44	0.51
6:X:925:LEU:C	6:X:927:GLY:H	2.18	0.51
6:Y:219:LEU:HG	6:Y:350:LEU:HD21	1.91	0.51
6:Y:434:ILE:CG2	6:Y:1204:LEU:HD22	2.41	0.51
6:Y:602:GLY:HA3	6:Y:623:PRO:HG3	1.93	0.51
6:Y:876:ASP:HB2	6:Y:892:ASP:OD1	2.10	0.51
1:l:23:THR:HG22	1:l:23:THR:O	2.10	0.51
3:B:141:GLN:C	3:B:143:MET:H	2.16	0.51
3:B:360:TYR:CD1	3:B:361:ASP:N	2.77	0.51
3:D:63:ILE:CD1	3:D:63:ILE:N	2.50	0.51
3:D:92:TRP:CE2	3:D:156:VAL:CG2	2.94	0.51
3:D:496:GLU:C	3:D:498:VAL:H	2.17	0.51
3:F:306:TRP:CE3	3:F:306:TRP:O	2.64	0.51

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:F:545:VAL:CG2	3:H:641:ILE:HD11	2.40	0.51
3:H:357:ILE:HD11	3:H:414:GLN:HB3	1.91	0.51
2:I:9:ASN:HB3	3:J:209:ASN:CG	2.35	0.51
3:J:96:PHE:CD1	3:J:159:TRP:HH2	2.28	0.51
3:J:431:ILE:CD1	3:J:437:ILE:HG23	2.39	0.51
2:K:22:PRO:HD2	3:L:179:ALA:O	2.11	0.51
3:L:306:TRP:CE3	3:L:306:TRP:O	2.64	0.51
3:N:94:PRO:HG2	3:N:210:VAL:CG2	2.41	0.51
3:P:122:THR:O	3:P:123:ALA:C	2.54	0.51
3:R:122:THR:O	3:R:123:ALA:C	2.54	0.51
3:T:333:PRO:HG3	3:T:454:PHE:CA	2.41	0.51
3:T:357:ILE:HD11	3:T:414:GLN:CD	2.36	0.51
4:U:147:MET:HB3	4:U:288:GLN:NE2	2.26	0.51
4:U:168:ALA:C	4:U:170:THR:N	2.69	0.51
4:U:174:LEU:HD22	4:U:211:LEU:HD23	1.92	0.51
5:W:642:LEU:HA	5:W:658:VAL:O	2.10	0.51
5:W:766:LEU:HD13	5:W:841:ARG:NH2	2.18	0.51
5:W:914:VAL:HG12	5:W:918:ILE:CG1	2.40	0.51
6:X:315:SER:HB2	6:X:331:ARG:CD	2.40	0.51
6:X:331:ARG:NH2	6:X:1196:ARG:NE	2.58	0.51
6:X:493:LEU:HD11	6:X:497:ILE:HD12	1.91	0.51
6:X:1033:ASP:OD1	6:X:1033:ASP:C	2.53	0.51
6:Y:257:ARG:NH1	6:Y:309:LEU:O	2.44	0.51
6:Y:295:SER:HB2	6:Y:890:GLN:HE22	1.70	0.51
6:Y:403:ILE:HD13	6:Y:407:ARG:HG2	1.92	0.51
6:Y:586:PHE:HB2	6:Y:646:ILE:HD12	1.93	0.51
6:Y:1023:ARG:HD2	6:Y:1027:LEU:HD22	1.93	0.51
1:d:44:GLY:HA3	3:D:399:ALA:HB3	1.92	0.51
1:p:44:GLY:HA3	3:P:322:TYR:CE2	2.46	0.51
1:p:58:PRO:O	1:p:59:HIS:NE2	2.44	0.51
1:r:61:ASN:CB	3:R:397:VAL:HA	2.40	0.51
1:t:22:LEU:CD1	1:t:22:LEU:N	2.73	0.51
3:B:284:VAL:O	3:B:284:VAL:HG12	2.11	0.51
3:D:296:PRO:C	3:D:298:PHE:H	2.19	0.51
3:D:641:ILE:HD11	3:H:545:VAL:CG2	2.40	0.51
2:E:9:ASN:HB2	3:F:209:ASN:CG	2.35	0.51
3:F:475:THR:O	3:F:505:ASP:HB2	2.11	0.51
3:H:114:THR:HG22	3:H:114:THR:O	2.10	0.51
3:H:183:TRP:CZ3	3:H:186:LEU:HD23	2.46	0.51
3:H:306:TRP:CE3	3:H:306:TRP:O	2.64	0.51
3:H:333:PRO:HG3	3:H:454:PHE:CA	2.41	0.51

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:J:306:TRP:CE3	3:J:306:TRP:O	2.64	0.51
2:K:28:SER:HB3	3:L:215:TYR:HA	1.93	0.51
3:L:357:ILE:HG12	3:L:416:ARG:NH1	2.26	0.51
3:N:48:TRP:N	3:N:48:TRP:HD1	2.09	0.51
3:N:50:PRO:HG3	3:N:56:ALA:HB2	1.91	0.51
3:N:475:THR:O	3:N:505:ASP:HB2	2.11	0.51
3:P:46:LYS:NZ	3:T:237:GLU:CD	2.68	0.51
3:P:269:LYS:O	3:P:270:ASP:HB2	2.09	0.51
3:P:306:TRP:CE3	3:P:306:TRP:O	2.64	0.51
3:P:357:ILE:HD11	3:P:414:GLN:CD	2.36	0.51
3:P:368:THR:CG2	3:P:446:PRO:HG3	2.38	0.51
3:P:540:VAL:CG2	3:P:569:ALA:HB3	2.40	0.51
3:R:92:TRP:CE2	3:R:156:VAL:CG2	2.94	0.51
3:R:94:PRO:HG2	3:R:210:VAL:CG2	2.41	0.51
3:R:96:PHE:CD1	3:R:159:TRP:HH2	2.28	0.51
3:R:166:ALA:HA	3:R:194:LEU:HD21	1.93	0.51
3:R:296:PRO:C	3:R:298:PHE:N	2.67	0.51
3:R:345:ILE:O	3:R:346:ASP:C	2.52	0.51
3:R:357:ILE:HG12	3:R:416:ARG:NH1	2.26	0.51
3:T:369:SER:OG	3:T:443:PHE:CD1	2.64	0.51
3:T:475:THR:O	3:T:505:ASP:HB2	2.11	0.51
5:W:292:VAL:HG12	5:W:292:VAL:O	2.11	0.51
5:W:628:TRP:HA	5:W:631:VAL:HG12	1.93	0.51
5:W:748:ARG:HG2	5:W:757:THR:CG2	2.36	0.51
5:W:815:TYR:CG	5:W:1025:ILE:HD11	2.46	0.51
5:W:1075:PRO:HA	5:W:1087:LEU:CD1	2.40	0.51
6:X:483:VAL:CB	6:X:840:ALA:HB1	2.40	0.51
6:X:589:LEU:O	6:X:590:ALA:C	2.54	0.51
6:X:893:VAL:HG21	6:Y:687:LEU:O	2.11	0.51
6:X:997:ALA:HB2	6:X:1143:ASN:HB3	1.92	0.51
6:X:1017:PHE:CE1	6:Y:1211:ARG:HA	2.46	0.51
6:Y:334:GLU:O	6:Y:335:GLY:C	2.53	0.51
6:Y:665:SER:O	6:Y:666:ALA:HB3	2.11	0.51
1:t:5:MET:CA	1:t:5:MET:CE	2.86	0.51
2:A:28:SER:HB3	3:B:215:TYR:HA	1.93	0.51
3:B:114:THR:HG22	3:B:114:THR:O	2.11	0.51
3:B:183:TRP:NE1	3:B:187:LEU:HD12	2.26	0.51
3:B:306:TRP:CE3	3:B:306:TRP:O	2.64	0.51
3:B:431:ILE:CD1	3:B:437:ILE:HG23	2.39	0.51
3:D:333:PRO:HG3	3:D:454:PHE:CA	2.41	0.51
3:D:357:ILE:HD11	3:D:414:GLN:HB3	1.91	0.51

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:F:183:TRP:CZ3	3:F:186:LEU:HD23	2.46	0.51
3:F:203:TYR:O	3:F:206:GLU:N	2.44	0.51
3:F:496:GLU:C	3:F:498:VAL:H	2.17	0.51
3:J:114:THR:HG22	3:J:114:THR:O	2.11	0.51
3:J:137:ILE:HD11	3:J:142:ILE:HA	1.93	0.51
3:J:298:PHE:CZ	3:N:416:ARG:HG3	2.46	0.51
3:J:582:LEU:C	3:J:584:LEU:H	2.19	0.51
3:L:183:TRP:NE1	3:L:187:LEU:HD12	2.26	0.51
3:L:237:GLU:CD	3:N:46:LYS:NZ	2.68	0.51
3:N:141:GLN:C	3:N:143:MET:N	2.63	0.51
3:N:166:ALA:HA	3:N:194:LEU:HD21	1.93	0.51
3:P:333:PRO:HG3	3:P:454:PHE:CA	2.41	0.51
3:P:582:LEU:C	3:P:584:LEU:H	2.19	0.51
2:Q:22:PRO:HD2	3:R:179:ALA:O	2.11	0.51
7:Q:101:MYR:C2	3:R:191:LYS:HZ1	2.21	0.51
3:R:183:TRP:CZ3	3:R:186:LEU:HD23	2.46	0.51
3:R:273:LEU:HD23	3:R:273:LEU:O	2.12	0.51
3:T:48:TRP:N	3:T:48:TRP:HD1	2.09	0.51
3:T:137:ILE:CD1	3:T:142:ILE:HG13	2.41	0.51
4:V:181:PRO:HG2	6:Y:412:ASP:CG	2.35	0.51
5:W:1059:ALA:HB3	5:W:1096:PHE:HZ	1.76	0.51
6:X:448:SER:HB2	6:X:664:GLY:HA3	1.93	0.51
6:Y:560:LEU:HD21	6:Y:721:VAL:CG2	2.41	0.51
6:Y:945:VAL:O	6:Y:945:VAL:HG13	2.10	0.51
1:b:59:HIS:HB2	3:B:397:VAL:HG22	1.92	0.50
1:f:59:HIS:HB2	3:F:397:VAL:C	2.34	0.50
1:j:59:HIS:HA	3:J:398:SER:HA	1.93	0.50
1:j:74:GLN:OE1	1:j:75:PRO:CD	2.58	0.50
1:n:5:MET:O	1:n:6:ILE:HB	2.11	0.50
1:n:15:ARG:HH12	3:N:402:ALA:CB	2.21	0.50
1:n:61:ASN:CB	3:N:397:VAL:HA	2.40	0.50
1:p:59:HIS:CB	3:P:398:SER:N	2.29	0.50
1:t:15:ARG:CZ	3:T:406:ASN:CG	2.84	0.50
3:D:166:ALA:HA	3:D:194:LEU:HD21	1.93	0.50
3:D:575:MET:HE3	3:H:469:VAL:HG21	0.69	0.50
3:F:137:ILE:HD11	3:F:142:ILE:HA	1.93	0.50
3:F:144:ASN:C	3:F:146:GLN:N	2.65	0.50
3:H:92:TRP:CZ2	3:H:160:GLN:HB2	2.46	0.50
3:H:166:ALA:HA	3:H:194:LEU:HD21	1.93	0.50
3:H:345:ILE:O	3:H:346:ASP:C	2.52	0.50
3:J:166:ALA:HA	3:J:194:LEU:HD21	1.93	0.50

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:J:183:TRP:CZ3	3:J:186:LEU:HD23	2.46	0.50
3:J:285:PHE:CE2	3:J:484:ILE:HD12	2.45	0.50
3:J:357:ILE:HD11	3:J:414:GLN:CD	2.36	0.50
3:L:114:THR:HG22	3:L:114:THR:O	2.11	0.50
3:L:122:THR:O	3:L:123:ALA:C	2.54	0.50
3:L:285:PHE:CE2	3:L:484:ILE:HD12	2.45	0.50
3:L:475:THR:O	3:L:505:ASP:HB2	2.11	0.50
3:N:296:PRO:C	3:N:298:PHE:H	2.19	0.50
3:N:306:TRP:CE3	3:N:306:TRP:O	2.64	0.50
3:N:485:VAL:HG12	3:N:485:VAL:O	2.10	0.50
2:O:22:PRO:HD2	3:P:179:ALA:O	2.11	0.50
3:P:82:PHE:CZ	3:T:238:VAL:HG11	2.44	0.50
3:P:94:PRO:HG2	3:P:210:VAL:CG2	2.41	0.50
3:P:284:VAL:HG12	3:P:284:VAL:O	2.11	0.50
3:R:154:ARG:NH1	3:R:154:ARG:CB	2.68	0.50
3:R:296:PRO:C	3:R:298:PHE:H	2.19	0.50
3:R:306:TRP:CE3	3:R:306:TRP:O	2.64	0.50
3:R:609:ARG:HD2	2:S:29:THR:HG22	1.91	0.50
3:T:92:TRP:C	3:T:94:PRO:HD2	2.37	0.50
4:V:64:ARG:HH21	6:X:1027:LEU:CD2	2.23	0.50
4:V:78:PRO:HG3	4:V:83:GLN:O	2.11	0.50
5:W:773:ARG:HH11	5:W:872:THR:CG2	2.24	0.50
5:W:1059:ALA:HB3	5:W:1096:PHE:CZ	2.46	0.50
5:W:1208:CYS:HB3	5:W:1216:GLY:HA3	1.90	0.50
6:X:301:LEU:HD11	6:X:968:PHE:CB	2.41	0.50
6:X:462:ILE:CD1	6:X:468:ALA:HB1	2.40	0.50
6:X:562:SER:HB3	6:X:592:LEU:CD1	2.41	0.50
6:X:581:SER:N	6:X:582:PRO:HD2	2.26	0.50
6:X:898:ALA:HB2	6:X:930:ARG:HG3	1.93	0.50
6:Y:38:SER:O	6:Y:39:PRO:C	2.54	0.50
6:Y:254:ASP:CG	6:Y:255:GLU:N	2.68	0.50
6:Y:538:ASN:ND2	6:Y:769:VAL:O	2.44	0.50
6:Y:567:LEU:HD23	6:Y:708:PHE:CG	2.45	0.50
6:Y:718:THR:CG2	6:Y:719:ALA:N	2.74	0.50
6:Y:799:ASP:OD1	6:Y:801:ARG:HB2	2.10	0.50
1:l:45:ARG:CD	3:L:400:ALA:CA	2.83	0.50
1:f:58:PRO:HG2	3:F:423:GLN:HG3	1.93	0.50
1:d:6:ILE:HD11	3:F:586:VAL:HG21	1.93	0.50
1:d:6:ILE:O	1:d:8:GLN:N	2.45	0.50
1:d:21:ARG:O	3:D:323:SER:O	2.29	0.50
1:t:59:HIS:HD2	3:T:397:VAL:HG22	1.75	0.50

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:B:540:VAL:CG2	3:B:569:ALA:HB3	2.41	0.50
3:D:92:TRP:CZ2	3:D:160:GLN:HB2	2.46	0.50
3:D:122:THR:O	3:D:123:ALA:C	2.54	0.50
3:D:137:ILE:HD11	3:D:142:ILE:HA	1.93	0.50
3:D:203:TYR:O	3:D:206:GLU:N	2.44	0.50
3:F:92:TRP:CE2	3:F:156:VAL:CG2	2.94	0.50
3:F:114:THR:HG22	3:F:114:THR:O	2.10	0.50
3:F:287:PRO:HG2	3:F:471:LEU:CD1	2.39	0.50
3:F:358:LEU:CB	3:F:417:PHE:HE1	2.24	0.50
3:H:215:TYR:C	3:H:217:ALA:N	2.70	0.50
3:J:203:TYR:O	3:J:206:GLU:N	2.44	0.50
3:L:92:TRP:C	3:L:94:PRO:HD2	2.37	0.50
3:L:609:ARG:HD2	2:M:29:THR:HG22	1.91	0.50
2:M:28:SER:HB3	3:N:215:TYR:HA	1.93	0.50
3:P:114:THR:HG22	3:P:114:THR:O	2.11	0.50
3:P:285:PHE:CE2	3:P:484:ILE:HD12	2.45	0.50
3:R:92:TRP:C	3:R:94:PRO:HD2	2.37	0.50
3:R:92:TRP:CZ2	3:R:160:GLN:HB2	2.46	0.50
3:R:199:LEU:O	3:R:207:PHE:HE2	1.92	0.50
2:S:22:PRO:HD2	3:T:179:ALA:O	2.11	0.50
3:T:114:THR:HG22	3:T:114:THR:O	2.10	0.50
3:T:269:LYS:O	3:T:270:ASP:HB2	2.10	0.50
4:V:195:GLN:O	4:V:196:LEU:HD22	2.11	0.50
5:W:815:TYR:HA	5:W:818:PHE:HB3	1.91	0.50
5:W:946:LEU:N	5:W:946:LEU:HD22	2.26	0.50
5:W:1100:PHE:CE2	5:W:1102:PRO:HB3	2.46	0.50
5:W:1142:ARG:O	5:W:1143:GLN:HB3	2.10	0.50
5:W:1237:VAL:HG23	5:W:1289:LEU:HD23	1.93	0.50
6:X:209:ASN:O	6:X:210:ILE:C	2.53	0.50
6:X:242:ARG:HA	6:X:1145:SER:O	2.12	0.50
6:X:514:ILE:HD13	6:X:732:VAL:HG11	1.93	0.50
6:X:587:MET:O	6:X:588:ALA:C	2.54	0.50
6:X:895:SER:O	6:X:896:ALA:C	2.53	0.50
6:Y:355:LYS:HE2	6:Y:1161:ASN:OD1	2.12	0.50
6:Y:470:LEU:N	6:Y:471:PRO:CD	2.74	0.50
6:Y:868:ARG:CD	6:Y:932:ALA:O	2.60	0.50
1:h:24:LEU:HD12	1:h:80:ARG:CD	2.41	0.50
1:j:22:LEU:HD21	1:j:41:VAL:CG1	2.32	0.50
1:r:19:ALA:CB	1:r:21:ARG:HH12	2.24	0.50
2:A:22:PRO:HD2	3:B:179:ALA:O	2.11	0.50
2:C:9:ASN:HB2	3:D:209:ASN:CG	2.36	0.50

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:D:82:PHE:CZ	3:H:238:VAL:HG11	2.45	0.50
3:D:152:ARG:NH2	3:L:100:LYS:O	2.44	0.50
2:E:22:PRO:HD2	3:F:179:ALA:O	2.11	0.50
2:E:28:SER:HB3	3:F:215:TYR:HA	1.93	0.50
3:F:80:GLU:HA	3:F:80:GLU:OE2	2.12	0.50
3:F:296:PRO:C	3:F:298:PHE:N	2.68	0.50
3:L:203:TYR:O	3:L:206:GLU:N	2.44	0.50
3:L:249:VAL:HG12	3:L:617:VAL:HB	1.94	0.50
3:L:333:PRO:HG3	3:L:454:PHE:CA	2.41	0.50
3:L:431:ILE:CD1	3:L:437:ILE:HG23	2.39	0.50
3:L:606:THR:HG22	3:N:173:ASP:HB2	1.94	0.50
3:N:122:THR:O	3:N:123:ALA:C	2.54	0.50
2:O:26:MET:O	2:O:26:MET:CG	2.60	0.50
3:P:545:VAL:CG2	3:R:641:ILE:HD11	2.40	0.50
3:R:203:TYR:O	3:R:206:GLU:N	2.44	0.50
3:R:357:ILE:HD11	3:R:414:GLN:CD	2.36	0.50
3:R:369:SER:OG	3:R:443:PHE:CD1	2.64	0.50
3:R:540:VAL:CG2	3:R:569:ALA:HB3	2.41	0.50
2:S:28:SER:HB3	3:T:215:TYR:HA	1.93	0.50
3:T:92:TRP:CE2	3:T:156:VAL:CG2	2.94	0.50
3:T:166:ALA:HA	3:T:194:LEU:HD21	1.93	0.50
3:T:306:TRP:CE3	3:T:306:TRP:O	2.64	0.50
4:V:192:PHE:HB2	6:Y:376:LEU:HD23	1.84	0.50
5:W:504:THR:CG2	5:W:1184:VAL:HG21	2.41	0.50
5:W:619:LYS:HE3	5:W:654:GLU:CD	2.37	0.50
6:Y:531:CYS:SG	6:Y:818:ILE:HD11	2.51	0.50
6:Y:552:ASP:OD2	6:Y:555:GLN:HB2	2.12	0.50
6:Y:996:MET:HE3	6:Y:1117:PHE:CD2	2.46	0.50
6:Y:1019:THR:O	6:Y:1032:ILE:HB	2.11	0.50
1:n:21:ARG:NE	3:N:370:VAL:HG11	2.25	0.50
1:t:23:THR:CG2	1:t:31:GLU:CB	2.70	0.50
2:A:9:ASN:HB2	3:B:209:ASN:CG	2.35	0.50
3:B:92:TRP:C	3:B:94:PRO:HD2	2.37	0.50
3:B:203:TYR:O	3:B:206:GLU:N	2.44	0.50
3:B:344:MET:HE2	3:B:502:LEU:CD1	2.37	0.50
3:B:411:VAL:O	3:B:412:ARG:CB	2.56	0.50
3:B:443:PHE:HE2	3:B:459:LEU:HD21	1.75	0.50
2:C:22:PRO:HD2	3:D:179:ALA:O	2.11	0.50
3:D:94:PRO:HG2	3:D:210:VAL:CG2	2.41	0.50
3:D:357:ILE:HG12	3:D:416:ARG:NH1	2.26	0.50
3:F:536:THR:HG21	3:F:570:LYS:HD2	1.94	0.50

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:H:273:LEU:HD23	3:H:273:LEU:O	2.12	0.50
3:J:57:THR:CG2	3:J:58:ILE:H	2.25	0.50
3:J:92:TRP:CE2	3:J:156:VAL:CG2	2.94	0.50
3:L:416:ARG:HG3	3:N:298:PHE:CZ	2.46	0.50
3:N:80:GLU:HA	3:N:80:GLU:OE2	2.12	0.50
3:N:273:LEU:HD23	3:N:273:LEU:O	2.12	0.50
2:O:28:SER:HB3	3:P:215:TYR:HA	1.93	0.50
3:P:137:ILE:HD11	3:P:142:ILE:HA	1.93	0.50
3:P:249:VAL:HG12	3:P:617:VAL:HB	1.94	0.50
3:T:285:PHE:CE2	3:T:484:ILE:HD12	2.45	0.50
4:U:26:LEU:HD13	4:U:90:ILE:HG13	1.93	0.50
4:U:176:ALA:HB1	6:X:413:PRO:CD	2.41	0.50
4:U:348:THR:HG22	4:U:349:ALA:N	2.27	0.50
5:W:53:ASP:O	5:W:55:SER:N	2.44	0.50
5:W:135:LEU:HD23	5:W:138:LEU:HD12	1.92	0.50
5:W:327:ILE:O	5:W:327:ILE:HG22	2.11	0.50
5:W:579:TYR:CE1	5:W:581:ASP:HB2	2.46	0.50
6:X:293:LEU:HD22	6:X:1080:HIS:O	2.11	0.50
6:X:481:ARG:O	6:X:482:SER:C	2.54	0.50
6:X:557:ILE:HG12	6:X:557:ILE:O	2.12	0.50
6:X:902:MET:HG3	6:X:929:PRO:HB3	1.94	0.50
6:Y:111:ILE:HG22	6:Y:111:ILE:O	2.10	0.50
6:Y:157:ILE:CD1	6:Y:1204:LEU:HB2	2.42	0.50
6:Y:543:THR:O	6:Y:543:THR:HG23	2.11	0.50
6:Y:1211:ARG:HG2	6:Y:1211:ARG:NH1	2.20	0.50
1:n:4:HIS:CD2	1:n:54:THR:HG22	2.46	0.50
1:t:60:ALA:N	3:T:398:SER:HA	2.26	0.50
1:t:62:VAL:O	1:t:65:ILE:N	2.44	0.50
3:B:122:THR:O	3:B:123:ALA:C	2.54	0.50
3:B:296:PRO:C	3:B:298:PHE:N	2.68	0.50
3:B:475:THR:O	3:B:505:ASP:HB2	2.11	0.50
3:D:86:LEU:HD12	3:D:108:LEU:HB2	1.92	0.50
3:F:94:PRO:HG2	3:F:210:VAL:CG2	2.41	0.50
3:H:183:TRP:NE1	3:H:187:LEU:HD12	2.26	0.50
3:H:284:VAL:HG12	3:H:284:VAL:O	2.10	0.50
2:I:22:PRO:HD2	3:J:179:ALA:O	2.11	0.50
3:J:79:ARG:NH1	3:J:79:ARG:HB2	2.27	0.50
3:J:122:THR:O	3:J:123:ALA:C	2.54	0.50
3:L:269:LYS:O	3:L:270:ASP:HB2	2.09	0.50
3:L:431:ILE:HG12	3:L:437:ILE:HD13	1.94	0.50
3:N:92:TRP:CE2	3:N:156:VAL:CG2	2.94	0.50

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:N:148:THR:OG1	3:R:100:LYS:HD2	2.11	0.50
3:N:215:TYR:C	3:N:217:ALA:N	2.70	0.50
3:N:357:ILE:HD11	3:N:414:GLN:CD	2.36	0.50
3:N:582:LEU:C	3:N:584:LEU:H	2.18	0.50
3:R:249:VAL:HG12	3:R:617:VAL:HB	1.94	0.50
3:R:294:TYR:HH	3:R:502:LEU:HD23	1.75	0.50
3:R:485:VAL:HG12	3:R:485:VAL:O	2.10	0.50
3:R:536:THR:HG21	3:R:570:LYS:HD2	1.94	0.50
3:T:215:TYR:C	3:T:217:ALA:N	2.70	0.50
3:T:296:PRO:C	3:T:298:PHE:H	2.19	0.50
4:U:171:THR:O	4:U:175:MET:CG	2.54	0.50
5:W:35:GLN:NE2	5:W:96:VAL:HG11	2.26	0.50
5:W:46:ALA:HA	5:W:59:ILE:HG12	1.93	0.50
5:W:47:LEU:CD1	5:W:327:ILE:HG21	2.42	0.50
5:W:316:GLN:HA	5:W:316:GLN:OE1	2.12	0.50
5:W:360:TRP:HB3	5:W:363:SER:HB2	1.92	0.50
5:W:472:ARG:HG2	5:W:692:VAL:HG12	1.93	0.50
5:W:475:LEU:HD23	5:W:475:LEU:C	2.36	0.50
5:W:816:ASN:HA	5:W:845:LEU:CD2	2.41	0.50
5:W:1107:ILE:HA	5:W:1126:THR:HG22	1.92	0.50
6:X:476:LEU:HD12	6:X:533:ALA:HA	1.94	0.50
6:X:1020:ILE:CD1	6:X:1062:TYR:HB2	2.41	0.50
6:X:1169:TRP:N	6:X:1169:TRP:CD1	2.79	0.50
6:Y:278:LEU:CD1	6:Y:891:PHE:HZ	2.25	0.50
6:Y:579:PRO:HG3	6:Y:807:PHE:HE2	1.74	0.50
6:Y:938:TYR:CE2	6:Y:946:VAL:HG21	2.46	0.50
1:b:59:HIS:CA	3:B:398:SER:CB	2.66	0.50
1:d:45:ARG:CG	3:D:399:ALA:O	2.60	0.50
1:n:45:ARG:N	3:N:399:ALA:O	2.44	0.50
1:r:4:HIS:HE1	1:r:54:THR:HG23	1.76	0.50
3:B:92:TRP:CZ2	3:B:160:GLN:HB2	2.46	0.50
3:B:183:TRP:CZ3	3:B:186:LEU:HD23	2.45	0.50
3:D:53:THR:CG2	3:D:54:SER:N	2.57	0.50
3:D:79:ARG:HB2	3:D:79:ARG:NH1	2.27	0.50
3:D:306:TRP:CE3	3:D:306:TRP:O	2.64	0.50
3:D:344:MET:HE2	3:D:502:LEU:CD1	2.37	0.50
3:D:582:LEU:C	3:D:584:LEU:H	2.19	0.50
3:F:284:VAL:O	3:F:284:VAL:HG12	2.11	0.50
2:G:22:PRO:HD2	3:H:179:ALA:O	2.11	0.50
3:J:92:TRP:C	3:J:94:PRO:HD2	2.37	0.50
3:J:360:TYR:CD1	3:J:361:ASP:N	2.77	0.50

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:J:540:VAL:CG2	3:J:569:ALA:HB3	2.41	0.50
2:K:9:ASN:HB2	3:L:209:ASN:CG	2.35	0.50
3:L:94:PRO:HG2	3:L:210:VAL:CG2	2.41	0.50
3:L:357:ILE:HD11	3:L:414:GLN:CD	2.36	0.50
3:L:536:THR:HG21	3:L:570:LYS:HD2	1.94	0.50
3:L:540:VAL:CG2	3:L:569:ALA:HB3	2.41	0.50
3:N:92:TRP:C	3:N:94:PRO:HD2	2.37	0.50
3:N:203:TYR:O	3:N:206:GLU:N	2.44	0.50
3:N:284:VAL:O	3:N:284:VAL:HG12	2.10	0.50
3:N:369:SER:OG	3:N:443:PHE:CD1	2.64	0.50
2:O:32:PRO:HD2	2:Q:38:PRO:HB2	1.93	0.50
3:P:92:TRP:CE2	3:P:156:VAL:CG2	2.94	0.50
3:P:166:ALA:HA	3:P:194:LEU:HD21	1.93	0.50
3:P:173:ASP:HB2	3:T:606:THR:HG22	1.94	0.50
3:P:431:ILE:HG12	3:P:437:ILE:HD13	1.94	0.50
3:P:609:ARG:HD2	2:Q:29:THR:HG22	1.91	0.50
3:R:114:THR:HG22	3:R:114:THR:O	2.10	0.50
3:R:144:ASN:C	3:R:146:GLN:N	2.65	0.50
3:R:360:TYR:CD1	3:R:361:ASP:N	2.77	0.50
2:S:26:MET:O	2:S:26:MET:CG	2.60	0.50
3:T:118:TYR:HB3	3:T:126:ALA:O	2.12	0.50
3:T:183:TRP:NE1	3:T:187:LEU:HD12	2.26	0.50
3:T:485:VAL:HG12	3:T:485:VAL:O	2.10	0.50
5:W:383:PHE:N	5:W:383:PHE:CD1	2.79	0.50
5:W:989:HIS:CE1	5:W:1027:HIS:ND1	2.79	0.50
6:X:370:ARG:HA	6:X:387:CYS:HB3	1.94	0.50
6:X:572:ALA:HA	6:X:818:ILE:HG21	1.94	0.50
6:X:841:PHE:HA	6:X:844:LEU:CD1	2.41	0.50
6:X:1149:THR:HG22	6:X:1162:ALA:H	1.72	0.50
6:Y:535:TRP:HD1	6:Y:577:SER:HB3	1.75	0.50
6:Y:595:GLN:O	6:Y:598:PRO:HD3	2.10	0.50
6:Y:690:ALA:CB	6:Y:691:PRO:HD3	2.24	0.50
6:Y:746:TYR:CD1	6:Y:750:MET:HB2	2.46	0.50
1:f:61:ASN:CB	3:F:397:VAL:HA	2.42	0.50
1:h:71:CYS:SG	1:h:72:SER:N	2.85	0.50
1:p:42:THR:HB	1:p:60:ALA:HB2	1.92	0.50
1:r:25:TYR:HB2	1:r:40:TYR:HB2	1.92	0.50
3:B:94:PRO:HG2	3:B:210:VAL:CG2	2.41	0.50
3:B:215:TYR:C	3:B:217:ALA:N	2.70	0.50
3:B:249:VAL:HG12	3:B:617:VAL:HB	1.94	0.50
3:D:64:VAL:O	3:D:64:VAL:HG22	2.11	0.50

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:D:475:THR:O	3:D:505:ASP:HB2	2.11	0.50
3:F:273:LEU:HD23	3:F:273:LEU:O	2.11	0.50
3:F:344:MET:O	3:F:345:ILE:C	2.55	0.50
3:F:431:ILE:HG12	3:F:437:ILE:HD13	1.94	0.50
3:H:137:ILE:HD11	3:H:142:ILE:HA	1.93	0.50
3:H:203:TYR:O	3:H:206:GLU:N	2.44	0.50
3:H:564:LEU:HA	3:H:567:ARG:NH2	2.27	0.50
3:J:118:TYR:HB3	3:J:126:ALA:O	2.12	0.50
3:L:183:TRP:CZ3	3:L:186:LEU:HD23	2.45	0.50
3:L:468:PRO:O	3:N:578:SER:OG	2.21	0.50
3:N:183:TRP:NE1	3:N:187:LEU:HD12	2.26	0.50
3:N:360:TYR:CD1	3:N:361:ASP:N	2.77	0.50
3:N:431:ILE:CD1	3:N:437:ILE:HG23	2.39	0.50
2:O:28:SER:HB2	3:P:214:ARG:O	2.10	0.50
3:P:92:TRP:C	3:P:94:PRO:HD2	2.37	0.50
3:P:203:TYR:O	3:P:206:GLU:N	2.44	0.50
3:P:273:LEU:HD23	3:P:273:LEU:O	2.12	0.50
3:P:357:ILE:HG12	3:P:416:ARG:NH1	2.26	0.50
3:R:606:THR:HG22	3:T:173:ASP:HB2	1.94	0.50
3:T:358:LEU:CB	3:T:417:PHE:HE1	2.24	0.50
5:W:356:GLN:OE1	5:W:374:ALA:HB2	2.12	0.50
5:W:400:LEU:HD13	5:W:785:LEU:HD21	1.93	0.50
5:W:917:LEU:HD23	5:W:917:LEU:C	2.37	0.50
5:W:1142:ARG:O	5:W:1143:GLN:CB	2.59	0.50
6:X:390:LEU:HA	6:X:1191:TYR:HD1	1.76	0.50
6:X:456:VAL:HG21	6:X:950:VAL:HG11	1.94	0.50
6:X:685:PRO:HG2	6:X:833:PRO:HG2	1.94	0.50
6:X:906:ASN:OD1	6:X:919:LEU:HD23	2.10	0.50
6:X:1004:PHE:CD2	6:X:1044:ASP:HB3	2.47	0.50
6:X:1130:ALA:HB2	6:Y:104:GLN:CA	2.41	0.50
6:X:1179:VAL:HG23	6:X:1193:SER:HB3	1.94	0.50
6:Y:54:VAL:CB	6:Y:57:GLN:HE21	2.23	0.50
6:Y:1182:ASN:O	6:Y:1183:ASP:HB2	2.11	0.50
1:d:16:ALA:O	1:d:22:LEU:CD2	2.59	0.50
1:p:22:LEU:CD2	1:p:22:LEU:N	2.75	0.50
1:r:3:LEU:N	1:r:3:LEU:CD1	2.73	0.50
1:r:17:ALA:HA	1:r:22:LEU:CD2	2.41	0.50
3:B:79:ARG:NH1	3:B:79:ARG:HB2	2.27	0.50
3:B:449:THR:CG2	3:B:450:SER:N	2.66	0.50
3:D:92:TRP:C	3:D:94:PRO:HD2	2.37	0.50
3:D:183:TRP:NE1	3:D:187:LEU:HD12	2.26	0.50

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:D:249:VAL:HG12	3:D:617:VAL:HB	1.93	0.50
3:D:357:ILE:HD11	3:D:414:GLN:CD	2.36	0.50
3:D:385:VAL:HG23	3:D:390:ASN:HB3	1.94	0.50
3:F:215:TYR:C	3:F:217:ALA:N	2.70	0.50
3:F:357:ILE:HD11	3:F:414:GLN:CD	2.36	0.50
3:H:122:THR:O	3:H:123:ALA:C	2.54	0.50
3:H:536:THR:HG21	3:H:570:LYS:HD2	1.94	0.50
3:J:144:ASN:C	3:J:146:GLN:N	2.65	0.50
3:J:249:VAL:HG12	3:J:617:VAL:HB	1.93	0.50
3:J:273:LEU:HD23	3:J:273:LEU:O	2.12	0.50
3:J:284:VAL:O	3:J:284:VAL:HG12	2.11	0.50
3:J:344:MET:O	3:J:345:ILE:C	2.55	0.50
3:J:361:ASP:CG	3:J:362:THR:N	2.70	0.50
3:L:137:ILE:HD11	3:L:142:ILE:HA	1.93	0.50
3:N:79:ARG:NH1	3:N:79:ARG:HB2	2.27	0.50
3:N:181:CYS:O	3:N:182:ASN:C	2.55	0.50
3:N:287:PRO:HG2	3:N:471:LEU:CD1	2.39	0.50
3:N:536:THR:HG21	3:N:570:LYS:HD2	1.94	0.50
3:P:361:ASP:CG	3:P:362:THR:N	2.70	0.50
2:Q:26:MET:O	2:Q:26:MET:CG	2.60	0.50
3:R:48:TRP:N	3:R:48:TRP:HD1	2.09	0.50
3:R:494:THR:H	3:R:497:GLU:CB	2.19	0.50
5:W:75:ILE:O	5:W:78:PRO:HG3	2.11	0.50
5:W:88:ILE:C	5:W:90:ASP:N	2.69	0.50
5:W:686:ALA:O	5:W:690:ARG:HB3	2.12	0.50
5:W:736:VAL:HG13	5:W:744:ILE:CD1	2.42	0.50
6:X:274:TYR:CE2	6:X:300:MET:HE3	2.47	0.50
6:X:441:LEU:CD2	6:X:443:PRO:HD3	2.41	0.50
6:X:459:LEU:C	6:X:462:ILE:HG22	2.37	0.50
6:X:640:ARG:O	6:X:640:ARG:HG2	2.11	0.50
6:X:747:ILE:O	6:X:747:ILE:HG23	2.12	0.50
6:X:875:GLN:HG3	6:X:876:ASP:N	2.21	0.50
6:X:899:ALA:N	6:X:929:PRO:HG2	2.27	0.50
6:X:972:MET:HE1	6:X:1080:HIS:HB3	1.93	0.50
6:X:974:LEU:HD11	6:X:1168:ARG:NH2	2.27	0.50
6:Y:292:ALA:O	6:Y:296:ASN:HB2	2.11	0.50
6:Y:355:LYS:HB2	6:Y:1151:ALA:HA	1.94	0.50
6:Y:1103:TRP:O	6:Y:1104:LEU:C	2.54	0.50
1:j:74:GLN:OE1	1:j:75:PRO:CA	2.60	0.50
3:B:296:PRO:C	3:B:298:PHE:H	2.19	0.50
3:D:273:LEU:HD23	3:D:273:LEU:O	2.12	0.50

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:D:296:PRO:C	3:D:298:PHE:N	2.67	0.50
3:D:298:PHE:CZ	3:H:355:GLY:CA	2.80	0.50
3:D:344:MET:O	3:D:345:ILE:C	2.55	0.50
3:D:369:SER:OG	3:D:443:PHE:CD1	2.64	0.50
3:F:361:ASP:CG	3:F:362:THR:N	2.70	0.50
3:H:79:ARG:HB2	3:H:79:ARG:NH1	2.27	0.50
3:H:346:ASP:HB3	3:H:428:ARG:N	2.15	0.50
3:H:431:ILE:HG12	3:H:437:ILE:HD13	1.94	0.50
3:J:385:VAL:HG23	3:J:390:ASN:HB3	1.94	0.50
3:J:606:THR:HG22	3:L:173:ASP:HB2	1.94	0.50
3:P:144:ASN:C	3:P:146:GLN:N	2.65	0.50
3:P:183:TRP:NE1	3:P:187:LEU:HD12	2.26	0.50
3:P:536:THR:HG21	3:P:570:LYS:HD2	1.94	0.50
3:P:564:LEU:HA	3:P:567:ARG:NH2	2.27	0.50
3:R:80:GLU:HA	3:R:80:GLU:OE2	2.12	0.50
3:R:118:TYR:HB3	3:R:126:ALA:O	2.12	0.50
3:R:137:ILE:HD11	3:R:142:ILE:HA	1.94	0.50
3:R:333:PRO:HG3	3:R:454:PHE:CA	2.41	0.50
3:R:564:LEU:HA	3:R:567:ARG:NH2	2.27	0.50
3:T:122:THR:O	3:T:123:ALA:C	2.54	0.50
5:W:821:GLU:HG3	5:W:824:ARG:NH2	2.27	0.50
5:W:1166:CYS:HB2	5:W:1174:VAL:O	2.12	0.50
6:X:371:LEU:HD13	6:X:387:CYS:HA	1.94	0.50
6:X:509:SER:CB	6:X:512:SER:HB3	2.40	0.50
6:X:769:VAL:CG1	6:X:770:PRO:HD2	2.42	0.50
6:X:1004:PHE:HB3	6:X:1073:ARG:NH2	2.27	0.50
6:Y:123:ASN:CB	6:Y:161:LEU:HD13	2.42	0.50
6:Y:336:ARG:HG3	6:Y:336:ARG:O	2.12	0.50
6:Y:535:TRP:CD2	6:Y:678:PRO:HG2	2.46	0.50
1:r:23:THR:CB	1:r:42:THR:HG1	2.10	0.49
3:B:361:ASP:CG	3:B:362:THR:N	2.70	0.49
3:D:167:MET:HA	3:D:167:MET:CE	2.31	0.49
7:G:101:MYR:H142	7:G:101:MYR:C10	2.10	0.49
3:H:47:LEU:HB2	3:H:136:PHE:CE2	2.47	0.49
3:H:274:ILE:HD12	3:H:514:LEU:HD23	1.94	0.49
3:H:475:THR:O	3:H:505:ASP:HB2	2.11	0.49
3:J:215:TYR:C	3:J:217:ALA:N	2.70	0.49
3:J:296:PRO:C	3:J:298:PHE:H	2.19	0.49
3:J:333:PRO:HG3	3:J:454:PHE:CA	2.41	0.49
3:N:47:LEU:HB2	3:N:136:PHE:CE2	2.47	0.49
3:P:181:CYS:O	3:P:182:ASN:C	2.55	0.49

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:P:215:TYR:C	3:P:217:ALA:N	2.70	0.49
3:R:107:VAL:HG12	3:R:109:SER:H	1.77	0.49
3:R:288:SER:HB3	3:T:522:GLY:HA3	1.94	0.49
3:R:443:PHE:HE2	3:R:459:LEU:HD21	1.74	0.49
3:T:181:CYS:O	3:T:182:ASN:C	2.55	0.49
3:T:203:TYR:O	3:T:206:GLU:N	2.44	0.49
3:T:361:ASP:CG	3:T:362:THR:N	2.70	0.49
3:T:536:THR:HG21	3:T:570:LYS:HD2	1.94	0.49
4:V:2:ALA:HB2	6:Y:373:ARG:NH1	2.22	0.49
4:V:161:VAL:HG21	4:V:410:MET:HG3	1.93	0.49
4:V:245:GLN:HG3	6:Y:415:GLN:NE2	2.26	0.49
5:W:445:ILE:HD12	5:W:674:LEU:HB2	1.92	0.49
5:W:646:LYS:HD3	5:W:655:VAL:HG12	1.94	0.49
5:W:649:ILE:O	5:W:1010:LEU:CD1	2.60	0.49
5:W:961:GLY:O	5:W:963:ARG:N	2.45	0.49
6:X:975:PHE:O	6:X:976:ASN:HB2	2.12	0.49
6:X:993:TRP:HE1	6:X:995:PRO:HG3	1.75	0.49
6:X:1056:VAL:O	6:X:1057:ALA:C	2.55	0.49
6:X:1067:VAL:HG22	6:X:1072:THR:HG23	1.94	0.49
6:Y:244:VAL:HG22	6:Y:265:PRO:HA	1.93	0.49
6:Y:681:ASN:O	6:Y:682:GLN:HG3	2.12	0.49
6:Y:1019:THR:H	6:Y:1032:ILE:CG2	2.25	0.49
1:l:15:ARG:HH12	3:L:402:ALA:HB1	1.71	0.49
1:h:44:GLY:O	3:H:399:ALA:O	2.29	0.49
1:p:23:THR:HB	3:P:323:SER:CB	2.35	0.49
3:B:107:VAL:HG12	3:B:109:SER:H	1.77	0.49
3:B:154:ARG:NH1	3:B:154:ARG:CB	2.68	0.49
3:B:273:LEU:HD23	3:B:273:LEU:O	2.12	0.49
2:C:28:SER:HB3	3:D:215:TYR:HA	1.93	0.49
3:D:284:VAL:HG12	3:D:284:VAL:O	2.11	0.49
3:D:431:ILE:CD1	3:D:437:ILE:HG23	2.39	0.49
3:D:564:LEU:HA	3:D:567:ARG:NH2	2.27	0.49
3:F:369:SER:OG	3:F:443:PHE:CD1	2.64	0.49
3:H:344:MET:O	3:H:345:ILE:C	2.55	0.49
3:H:357:ILE:HD11	3:H:414:GLN:CD	2.36	0.49
3:J:181:CYS:O	3:J:182:ASN:C	2.55	0.49
3:J:431:ILE:HG12	3:J:437:ILE:HD13	1.94	0.49
3:L:118:TYR:HB3	3:L:126:ALA:O	2.12	0.49
3:N:114:THR:HG22	3:N:114:THR:O	2.10	0.49
3:N:358:LEU:CB	3:N:417:PHE:HE1	2.25	0.49
3:P:469:VAL:HG21	3:R:575:MET:HE3	0.69	0.49

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:R:475:THR:O	3:R:505:ASP:HB2	2.11	0.49
3:R:582:LEU:C	3:R:584:LEU:H	2.18	0.49
4:U:37:TRP:O	4:U:39:SER:N	2.42	0.49
5:W:72:HIS:HA	5:W:75:ILE:HD12	1.93	0.49
5:W:681:LEU:HD23	5:W:681:LEU:C	2.37	0.49
6:X:574:PRO:HB3	6:X:651:PRO:HG2	1.93	0.49
6:X:653:PRO:HD3	6:X:773:ARG:HB2	1.92	0.49
6:X:795:ASP:CG	6:Y:737:ARG:NH1	2.64	0.49
6:X:928:ASP:OD1	6:X:930:ARG:HD3	2.11	0.49
6:Y:485:PRO:HB3	6:Y:836:HIS:HD2	1.73	0.49
6:Y:670:ILE:HD13	6:Y:768:GLN:HB3	1.93	0.49
6:Y:776:GLN:HG2	6:Y:777:ASN:N	2.28	0.49
6:Y:891:PHE:CD1	6:Y:891:PHE:C	2.90	0.49
1:p:59:HIS:HD2	3:P:398:SER:CB	2.13	0.49
1:r:45:ARG:CD	3:R:400:ALA:HA	2.34	0.49
1:t:24:LEU:O	1:t:38:ALA:CB	2.60	0.49
1:t:61:ASN:HD21	3:T:395:THR:HG21	1.58	0.49
3:B:357:ILE:HD11	3:B:414:GLN:CD	2.36	0.49
3:B:385:VAL:HG23	3:B:390:ASN:HB3	1.94	0.49
3:D:80:GLU:OE2	3:D:80:GLU:HA	2.12	0.49
3:D:361:ASP:CG	3:D:362:THR:N	2.70	0.49
3:D:469:VAL:O	3:D:469:VAL:HG13	2.12	0.49
3:F:118:TYR:HB3	3:F:126:ALA:O	2.12	0.49
3:F:183:TRP:NE1	3:F:187:LEU:HD12	2.26	0.49
2:G:28:SER:HB3	3:H:215:TYR:HA	1.93	0.49
3:H:92:TRP:C	3:H:94:PRO:HD2	2.37	0.49
3:H:296:PRO:C	3:H:298:PHE:N	2.68	0.49
3:L:344:MET:O	3:L:345:ILE:C	2.55	0.49
3:L:346:ASP:HB3	3:L:428:ARG:N	2.15	0.49
3:L:361:ASP:CG	3:L:362:THR:N	2.70	0.49
3:P:47:LEU:HB2	3:P:136:PHE:CE2	2.47	0.49
3:P:296:PRO:C	3:P:298:PHE:H	2.19	0.49
3:P:369:SER:OG	3:P:443:PHE:CD1	2.64	0.49
3:P:469:VAL:O	3:P:469:VAL:HG13	2.12	0.49
3:R:174:ILE:HD11	3:R:178:SER:HA	1.95	0.49
3:T:174:ILE:HD11	3:T:178:SER:HA	1.95	0.49
3:T:249:VAL:HG12	3:T:617:VAL:HB	1.94	0.49
4:U:4:ARG:HH12	5:W:217:LEU:CB	2.25	0.49
4:U:302:GLU:O	4:U:303:ALA:C	2.55	0.49
4:U:328:TYR:CB	4:U:333:MET:HE2	2.42	0.49
5:W:62:TYR:CB	5:W:63:PRO:HD2	2.10	0.49

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
5:W:365:PHE:CD1	5:W:365:PHE:C	2.90	0.49
5:W:418:TYR:HB2	5:W:734:SER:HB2	1.95	0.49
5:W:1102:PRO:HG3	5:W:1127:ILE:CG1	2.42	0.49
6:X:650:TRP:HB3	6:X:773:ARG:HH11	1.76	0.49
6:X:692:THR:C	6:X:693:TYR:HD1	2.19	0.49
6:X:742:MET:O	6:X:742:MET:HG3	2.11	0.49
6:X:772:THR:HG21	6:X:778:ASP:HB2	1.94	0.49
6:X:795:ASP:OD1	6:Y:737:ARG:NH1	2.45	0.49
6:X:879:PHE:O	6:X:881:VAL:N	2.44	0.49
6:Y:861:ARG:HD3	6:Y:953:GLY:O	2.13	0.49
6:Y:988:ALA:CB	6:Y:1133:VAL:HG12	2.41	0.49
1:t:61:ASN:CB	3:T:395:THR:CG2	2.91	0.49
3:B:137:ILE:HD11	3:B:142:ILE:HA	1.93	0.49
3:B:300:LYS:HB3	3:B:302:GLU:OE1	2.13	0.49
3:B:469:VAL:O	3:B:469:VAL:HG13	2.12	0.49
3:D:238:VAL:HG11	3:F:82:PHE:CZ	2.45	0.49
3:D:431:ILE:HG12	3:D:437:ILE:HD13	1.94	0.49
3:F:47:LEU:HB2	3:F:136:PHE:CE2	2.47	0.49
3:F:79:ARG:NH1	3:F:79:ARG:HB2	2.27	0.49
3:F:92:TRP:C	3:F:94:PRO:HD2	2.36	0.49
3:F:274:ILE:HD12	3:F:514:LEU:HD23	1.94	0.49
2:I:26:MET:O	2:I:26:MET:CG	2.60	0.49
3:J:143:MET:SD	2:M:25:ASP:OD2	2.71	0.49
3:J:238:VAL:CG2	3:L:78:MET:HE1	2.07	0.49
3:J:269:LYS:HG2	3:J:516:LYS:HZ3	1.77	0.49
3:J:358:LEU:CB	3:J:417:PHE:HE1	2.25	0.49
3:J:469:VAL:O	3:J:469:VAL:HG13	2.12	0.49
2:K:25:ASP:OD2	3:N:143:MET:SD	2.71	0.49
3:L:107:VAL:HG12	3:L:109:SER:H	1.77	0.49
3:L:373:PHE:HA	3:L:441:GLY:HA2	1.95	0.49
3:N:274:ILE:HD12	3:N:514:LEU:HD23	1.94	0.49
2:O:25:ASP:OD2	3:R:143:MET:SD	2.71	0.49
3:P:65:SER:O	3:P:66:ASP:HB3	2.12	0.49
3:P:107:VAL:HG12	3:P:109:SER:H	1.77	0.49
3:P:360:TYR:CD1	3:P:361:ASP:N	2.77	0.49
3:R:65:SER:O	3:R:66:ASP:HB3	2.12	0.49
3:R:79:ARG:NH1	3:R:79:ARG:HB2	2.27	0.49
5:W:81:TYR:N	5:W:180:LEU:HD23	2.27	0.49
5:W:830:HIS:CE1	5:W:890:CYS:SG	3.06	0.49
6:X:255:GLU:OE1	6:X:255:GLU:HA	2.12	0.49
6:X:782:ASN:HB3	6:X:784:LEU:HD11	1.93	0.49

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
6:X:821:ALA:O	6:X:822:MET:C	2.55	0.49
6:X:931:ILE:HG22	6:X:931:ILE:O	2.12	0.49
6:Y:46:ALA:O	6:Y:50:GLN:HB2	2.12	0.49
6:Y:466:THR:HB	6:Y:809:ALA:HB1	1.94	0.49
6:Y:545:LEU:HD13	6:Y:569:LEU:HD12	1.95	0.49
1:b:21:ARG:CZ	3:B:327:TYR:OH	2.60	0.49
1:f:58:PRO:HG2	3:F:423:GLN:O	2.12	0.49
1:t:8:GLN:NE2	3:P:303:ASP:N	2.59	0.49
3:B:344:MET:O	3:B:345:ILE:C	2.55	0.49
3:B:354:SER:C	3:B:356:THR:H	2.21	0.49
3:F:107:VAL:HG12	3:F:109:SER:H	1.77	0.49
3:F:166:ALA:HA	3:F:194:LEU:HD21	1.93	0.49
3:F:174:ILE:HD11	3:F:178:SER:HA	1.95	0.49
3:F:564:LEU:HA	3:F:567:ARG:NH2	2.27	0.49
3:H:65:SER:O	3:H:66:ASP:HB3	2.12	0.49
3:H:181:CYS:O	3:H:182:ASN:C	2.55	0.49
3:H:300:LYS:HB3	3:H:302:GLU:OE1	2.13	0.49
3:H:361:ASP:CG	3:H:362:THR:N	2.70	0.49
3:H:385:VAL:HG23	3:H:390:ASN:HB3	1.94	0.49
3:J:300:LYS:HB3	3:J:302:GLU:OE1	2.13	0.49
3:J:522:GLY:HA3	3:N:288:SER:HB3	1.94	0.49
3:L:369:SER:OG	3:L:443:PHE:CD1	2.64	0.49
3:L:427:GLU:CD	3:N:571:ARG:HH22	2.19	0.49
3:L:469:VAL:O	3:L:469:VAL:HG13	2.13	0.49
2:M:22:PRO:HD2	3:N:179:ALA:O	2.11	0.49
3:N:137:ILE:HD11	3:N:142:ILE:HA	1.93	0.49
3:N:344:MET:O	3:N:345:ILE:C	2.55	0.49
3:N:361:ASP:CG	3:N:362:THR:N	2.70	0.49
3:N:431:ILE:HG12	3:N:437:ILE:HD13	1.94	0.49
3:P:78:MET:HE2	3:T:238:VAL:HG21	1.83	0.49
3:P:143:MET:SD	2:S:25:ASP:OD2	2.71	0.49
3:P:274:ILE:HD12	3:P:514:LEU:HD23	1.94	0.49
3:P:344:MET:O	3:P:345:ILE:C	2.55	0.49
2:Q:25:ASP:OD2	3:T:143:MET:SD	2.71	0.49
3:R:47:LEU:HD23	3:R:71:TYR:HB2	1.95	0.49
3:R:181:CYS:O	3:R:182:ASN:C	2.55	0.49
3:T:137:ILE:HD11	3:T:142:ILE:HA	1.93	0.49
3:T:273:LEU:HD23	3:T:273:LEU:O	2.12	0.49
3:T:330:GLN:HE21	3:T:332:GLY:CA	2.26	0.49
4:U:44:LEU:O	6:X:404:MET:SD	2.71	0.49
4:U:46:ILE:HG13	4:U:181:PRO:HG2	1.93	0.49

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
4:U:52:PRO:HG2	4:U:64:ARG:HH22	1.77	0.49
4:U:220:THR:O	4:U:222:ARG:N	2.44	0.49
4:U:248:LEU:HD12	4:U:249:LYS:N	2.28	0.49
5:W:496:PHE:HE2	5:W:658:VAL:HG21	1.77	0.49
5:W:1028:ILE:HG22	5:W:1029:ASP:N	2.28	0.49
5:W:1060:VAL:HG23	5:W:1113:THR:HG23	1.95	0.49
6:X:297:VAL:HG11	6:X:968:PHE:CZ	2.48	0.49
6:X:393:ALA:HB1	6:X:1184:LEU:HD13	1.95	0.49
6:X:463:SER:O	6:X:465:ASN:N	2.45	0.49
6:X:560:LEU:N	6:X:561:PRO:CD	2.75	0.49
6:X:788:VAL:HG23	6:X:931:ILE:HB	1.93	0.49
6:Y:206:LEU:HB3	6:Y:236:PRO:HG3	1.95	0.49
6:Y:271:VAL:HG13	6:Y:272:HIS:N	2.28	0.49
6:Y:658:TYR:CD1	6:Y:658:TYR:C	2.91	0.49
6:Y:864:PHE:CZ	6:Y:931:ILE:HD12	2.47	0.49
1:d:45:ARG:HG2	3:D:399:ALA:O	2.12	0.49
1:p:58:PRO:CD	3:P:423:GLN:HE21	2.25	0.49
3:B:47:LEU:HD23	3:B:71:TYR:HB2	1.95	0.49
3:B:118:TYR:HB3	3:B:126:ALA:O	2.12	0.49
3:D:118:TYR:HB3	3:D:126:ALA:O	2.12	0.49
3:D:358:LEU:CB	3:D:417:PHE:HE1	2.25	0.49
3:F:181:CYS:O	3:F:182:ASN:C	2.55	0.49
3:H:107:VAL:HG12	3:H:109:SER:H	1.77	0.49
3:H:249:VAL:HG12	3:H:617:VAL:HB	1.94	0.49
3:H:330:GLN:HE21	3:H:332:GLY:CA	2.26	0.49
3:J:65:SER:O	3:J:66:ASP:HB3	2.13	0.49
3:J:167:MET:HA	3:J:167:MET:CE	2.31	0.49
3:J:564:LEU:HA	3:J:567:ARG:NH2	2.27	0.49
3:L:80:GLU:OE2	3:L:80:GLU:HA	2.12	0.49
3:L:330:GLN:HE21	3:L:332:GLY:CA	2.26	0.49
3:L:469:VAL:HG21	3:N:575:MET:HE3	0.69	0.49
3:L:494:THR:H	3:L:497:GLU:CB	2.19	0.49
3:L:564:LEU:HA	3:L:567:ARG:NH2	2.27	0.49
3:P:80:GLU:OE2	3:P:80:GLU:HA	2.12	0.49
3:P:164:ASP:C	3:P:166:ALA:N	2.70	0.49
3:P:192:ASP:O	3:P:193:ILE:C	2.56	0.49
3:P:330:GLN:HE21	3:P:332:GLY:CA	2.26	0.49
3:P:475:THR:O	3:P:505:ASP:HB2	2.11	0.49
3:R:344:MET:O	3:R:345:ILE:C	2.55	0.49
3:R:361:ASP:CG	3:R:362:THR:N	2.70	0.49
3:T:47:LEU:HD23	3:T:71:TYR:HB2	1.95	0.49

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:T:82:PHE:O	3:T:86:LEU:HG	2.13	0.49
3:T:192:ASP:O	3:T:193:ILE:C	2.56	0.49
3:T:344:MET:O	3:T:345:ILE:C	2.55	0.49
5:W:740:VAL:HG12	5:W:741:PRO:N	2.27	0.49
5:W:992:SER:O	5:W:997:TRP:CZ3	2.65	0.49
6:X:662:LEU:H	6:X:662:LEU:CD2	2.25	0.49
6:X:837:TYR:C	6:X:839:LYS:H	2.21	0.49
6:X:1125:TYR:C	6:X:1157:ALA:HB3	2.37	0.49
6:Y:603:VAL:HG13	6:Y:603:VAL:O	2.12	0.49
6:Y:975:PHE:O	6:Y:976:ASN:HB2	2.12	0.49
1:l:23:THR:HG21	1:l:25:TYR:CZ	2.45	0.49
1:b:61:ASN:C	1:b:61:ASN:HD22	2.20	0.49
1:h:26:THR:HA	1:h:38:ALA:HA	1.95	0.49
1:h:35:PHE:CZ	1:h:74:GLN:HB3	2.47	0.49
1:h:58:PRO:CB	3:H:423:GLN:CD	2.86	0.49
1:j:6:ILE:HD11	3:L:586:VAL:CG2	2.43	0.49
1:t:58:PRO:C	1:t:59:HIS:ND1	2.70	0.49
3:D:107:VAL:HG12	3:D:109:SER:H	1.77	0.49
3:D:330:GLN:HE21	3:D:332:GLY:CA	2.26	0.49
3:D:606:THR:HG22	3:F:173:ASP:HB2	1.94	0.49
3:F:65:SER:O	3:F:66:ASP:HB3	2.12	0.49
3:F:606:THR:HG22	3:H:173:ASP:HB2	1.94	0.49
3:H:118:TYR:HB3	3:H:126:ALA:O	2.12	0.49
3:H:297:ALA:CB	3:H:460:MET:HB2	2.33	0.49
3:L:47:LEU:HB2	3:L:136:PHE:CE2	2.47	0.49
3:L:65:SER:O	3:L:66:ASP:HB3	2.12	0.49
3:L:215:TYR:C	3:L:217:ALA:N	2.70	0.49
3:L:358:LEU:CB	3:L:417:PHE:HE1	2.25	0.49
3:N:137:ILE:CD1	3:N:142:ILE:HG13	2.42	0.49
3:R:297:ALA:CB	3:R:460:MET:HB2	2.33	0.49
4:U:131:ASN:O	4:U:133:SER:N	2.45	0.49
4:U:291:THR:HA	4:U:294:THR:HG22	1.95	0.49
4:V:312:VAL:HG12	4:V:313:ALA:N	2.27	0.49
5:W:75:ILE:HD11	5:W:199:PRO:CG	2.39	0.49
5:W:736:VAL:HG22	5:W:744:ILE:HD13	1.95	0.49
5:W:832:LEU:HD21	5:W:893:ALA:HB2	1.94	0.49
5:W:847:PRO:O	5:W:849:THR:N	2.45	0.49
5:W:1295:GLU:O	5:W:1297:VAL:HG23	2.13	0.49
6:X:191:PRO:CB	6:X:859:ILE:HG13	2.42	0.49
6:X:386:GLU:HG3	6:X:1195:TYR:CD2	2.48	0.49
6:X:441:LEU:HD12	6:X:1199:PHE:CE2	2.48	0.49

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
6:Y:441:LEU:HG	6:Y:443:PRO:HD3	1.95	0.49
6:Y:483:VAL:HG13	6:Y:840:ALA:CB	2.41	0.49
6:Y:862:GLU:CB	6:Y:957:ILE:HD12	2.39	0.49
1:l:44:GLY:HA3	3:L:322:TYR:CZ	2.48	0.49
1:f:61:ASN:HB3	3:F:397:VAL:HA	1.94	0.49
1:d:61:ASN:CB	3:D:395:THR:CG2	2.90	0.49
1:n:50:ALA:HB1	1:n:74:GLN:HG2	1.95	0.49
1:r:26:THR:HA	1:r:38:ALA:HA	1.95	0.49
1:t:61:ASN:HD22	3:T:395:THR:HG22	1.73	0.49
3:B:174:ILE:HD11	3:B:178:SER:HA	1.95	0.49
3:B:181:CYS:O	3:B:182:ASN:C	2.55	0.49
3:B:431:ILE:HG12	3:B:437:ILE:HD13	1.94	0.49
3:B:494:THR:H	3:B:497:GLU:CB	2.19	0.49
3:B:536:THR:HG21	3:B:570:LYS:HD2	1.94	0.49
3:D:373:PHE:HD2	3:D:397:VAL:HG11	1.78	0.49
3:D:427:GLU:O	3:D:428:ARG:C	2.56	0.49
3:F:122:THR:O	3:F:123:ALA:C	2.54	0.49
3:F:385:VAL:HG23	3:F:390:ASN:HB3	1.94	0.49
3:J:47:LEU:HB2	3:J:136:PHE:CE2	2.47	0.49
3:L:181:CYS:O	3:L:182:ASN:C	2.55	0.49
3:N:107:VAL:HG12	3:N:109:SER:H	1.77	0.49
3:N:300:LYS:HB3	3:N:302:GLU:OE1	2.13	0.49
3:N:564:LEU:HA	3:N:567:ARG:NH2	2.27	0.49
3:P:571:ARG:HH22	3:T:427:GLU:CD	2.19	0.49
3:R:527:VAL:HA	3:R:533:ASP:OD2	2.13	0.49
3:T:47:LEU:HB2	3:T:136:PHE:CE2	2.47	0.49
3:T:79:ARG:HB2	3:T:79:ARG:NH1	2.27	0.49
4:U:9:LEU:HD11	4:U:117:ASN:ND2	2.28	0.49
4:U:275:GLN:HE22	4:U:387:GLN:HB3	1.77	0.49
4:U:291:THR:HG23	4:U:297:PRO:HD3	1.95	0.49
4:U:359:TYR:O	4:U:362:TYR:HB3	2.12	0.49
4:V:65:ASN:HA	4:V:68:VAL:HG12	1.95	0.49
5:W:313:LEU:CD2	5:W:347:GLN:HG3	2.43	0.49
5:W:508:PRO:HB2	5:W:511:ASP:HB2	1.95	0.49
5:W:720:PHE:HB3	5:W:782:THR:HB	1.95	0.49
5:W:816:ASN:HA	5:W:845:LEU:HD23	1.94	0.49
5:W:843:LEU:HD21	5:W:854:MET:CE	2.43	0.49
6:X:367:ARG:HG3	6:X:388:MET:HE2	1.94	0.49
6:X:450:ASP:OD2	6:X:861:ARG:NH1	2.46	0.49
6:X:536:ALA:HA	6:X:768:GLN:HE21	1.78	0.49
6:X:538:ASN:HD21	6:X:768:GLN:HG2	1.78	0.49

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
6:X:899:ALA:CA	6:X:929:PRO:HG2	2.43	0.49
6:X:979:ARG:CG	6:X:1140:GLU:HG3	2.42	0.49
6:Y:224:TRP:HB3	6:Y:229:TYR:HE1	1.78	0.49
6:Y:920:LEU:O	6:Y:920:LEU:HG	2.12	0.49
6:Y:1100:LEU:O	6:Y:1103:TRP:HB3	2.12	0.49
1:d:59:HIS:HB3	3:D:423:GLN:O	2.12	0.49
1:h:8:GLN:NE2	3:D:303:ASP:HA	2.28	0.49
1:n:26:THR:HA	1:n:38:ALA:HA	1.95	0.49
1:p:23:THR:HG21	1:p:25:TYR:CE2	2.47	0.49
3:B:47:LEU:HB2	3:B:136:PHE:CE2	2.47	0.49
3:B:80:GLU:OE2	3:B:80:GLU:HA	2.12	0.49
3:B:82:PHE:O	3:B:86:LEU:HG	2.13	0.49
3:B:373:PHE:HA	3:B:441:GLY:HA2	1.95	0.49
3:B:564:LEU:HA	3:B:567:ARG:NH2	2.27	0.49
3:D:143:MET:SD	2:G:25:ASP:OD2	2.71	0.49
3:D:181:CYS:O	3:D:182:ASN:C	2.55	0.49
3:D:536:THR:HG21	3:D:570:LYS:HD2	1.94	0.49
3:F:82:PHE:O	3:F:86:LEU:HG	2.13	0.49
3:F:238:VAL:HG11	3:H:82:PHE:CZ	2.45	0.49
3:F:355:GLY:O	3:F:416:ARG:HD2	2.13	0.49
3:F:427:GLU:O	3:F:428:ARG:C	2.56	0.49
3:H:80:GLU:OE2	3:H:80:GLU:HA	2.12	0.49
3:H:120:GLY:HA3	3:H:126:ALA:HA	1.95	0.49
3:H:373:PHE:HD2	3:H:397:VAL:HG11	1.78	0.49
3:J:288:SER:HB3	3:L:522:GLY:HA3	1.95	0.49
3:L:79:ARG:HB2	3:L:79:ARG:NH1	2.27	0.49
3:L:300:LYS:HB3	3:L:302:GLU:OE1	2.13	0.49
3:N:82:PHE:O	3:N:86:LEU:HG	2.13	0.49
3:N:118:TYR:HB3	3:N:126:ALA:O	2.12	0.49
3:P:346:ASP:HB3	3:P:428:ARG:N	2.15	0.49
2:Q:28:SER:HB3	3:R:215:TYR:HA	1.93	0.49
3:R:469:VAL:HG21	3:T:575:MET:HE3	0.69	0.49
3:T:381:PRO:O	3:T:384:GLN:HB3	2.13	0.49
4:U:406:LEU:HD13	4:U:406:LEU:O	2.13	0.49
4:V:198:GLN:OE1	6:Y:373:ARG:HB2	2.12	0.49
5:W:271:ARG:O	5:W:274:ALA:HB3	2.12	0.49
5:W:438:VAL:CG2	5:W:439:PHE:N	2.76	0.49
6:X:200:LEU:HD23	6:X:200:LEU:N	2.27	0.49
6:X:298:LEU:HG	6:X:965:GLU:OE2	2.12	0.49
6:X:340:LEU:O	6:X:341:ALA:C	2.55	0.49
6:X:599:ILE:HG23	6:X:639:LEU:HD23	1.93	0.49

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
6:X:669:PHE:C	6:X:670:ILE:HD13	2.37	0.49
6:X:982:LEU:HD12	6:X:1115:VAL:CG1	2.43	0.49
6:X:1171:GLY:HA2	6:X:1177:TYR:CD2	2.48	0.49
6:Y:164:TRP:CD1	6:Y:165:ASP:N	2.81	0.49
6:Y:388:MET:O	6:Y:389:GLU:O	2.31	0.49
6:Y:465:ASN:OD1	6:Y:468:ALA:N	2.42	0.49
6:Y:832:ILE:HD12	6:Y:835:HIS:HD2	1.76	0.49
6:Y:1009:PRO:O	6:Y:1010:ASN:HB2	2.11	0.49
3:B:427:GLU:O	3:B:428:ARG:C	2.56	0.49
2:C:42:ASN:O	3:H:239:ALA:HB2	2.13	0.49
3:D:47:LEU:HB2	3:D:136:PHE:CE2	2.47	0.49
3:D:144:ASN:C	3:D:146:GLN:N	2.65	0.49
3:D:469:VAL:HG21	3:F:575:MET:HE3	0.69	0.49
3:F:288:SER:HB3	3:H:522:GLY:HA3	1.94	0.49
3:F:333:PRO:HG3	3:F:454:PHE:CA	2.41	0.49
3:F:469:VAL:O	3:F:469:VAL:HG13	2.13	0.49
3:H:162:HIS:CD2	3:H:199:LEU:HB2	2.48	0.49
2:I:28:SER:HB3	3:J:215:TYR:HA	1.93	0.49
3:J:47:LEU:HD23	3:J:71:TYR:HB2	1.95	0.49
3:J:80:GLU:HA	3:J:80:GLU:OE2	2.12	0.49
3:J:239:ALA:HB2	2:K:42:ASN:O	2.13	0.49
3:J:358:LEU:HB2	3:J:417:PHE:HD1	1.77	0.49
3:J:494:THR:H	3:J:497:GLU:CB	2.19	0.49
3:L:273:LEU:HD23	3:L:273:LEU:O	2.12	0.49
3:N:65:SER:O	3:N:66:ASP:HB3	2.12	0.49
3:N:385:VAL:HG23	3:N:390:ASN:HB3	1.94	0.49
3:P:354:SER:C	3:P:356:THR:H	2.21	0.49
3:R:82:PHE:O	3:R:86:LEU:HG	2.13	0.49
3:R:237:GLU:CD	3:T:46:LYS:NZ	2.68	0.49
3:R:385:VAL:HG23	3:R:390:ASN:HB3	1.94	0.49
3:T:120:GLY:HA3	3:T:126:ALA:HA	1.95	0.49
3:T:162:HIS:CD2	3:T:199:LEU:HB2	2.48	0.49
3:T:164:ASP:C	3:T:166:ALA:N	2.71	0.49
3:T:385:VAL:HG23	3:T:390:ASN:HB3	1.94	0.49
3:T:431:ILE:HG12	3:T:437:ILE:HD13	1.94	0.49
4:U:372:ALA:HB3	4:U:373:MET:HE2	1.95	0.49
5:W:99:LEU:HD23	5:W:100:THR:N	2.28	0.49
5:W:240:LEU:HD21	5:W:261:LEU:HD13	1.95	0.49
5:W:750:PRO:HB3	5:W:791:PHE:CE1	2.47	0.49
5:W:1215:LEU:C	5:W:1215:LEU:CD1	2.85	0.49
6:X:279:SER:O	6:X:280:ARG:C	2.56	0.49

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
6:X:621:TRP:HE3	6:X:626:LEU:CD1	2.26	0.49
6:X:658:TYR:C	6:X:658:TYR:CD1	2.91	0.49
6:X:837:TYR:O	6:X:839:LYS:N	2.46	0.49
6:X:1091:GLN:HA	6:X:1091:GLN:HE21	1.77	0.49
6:Y:453:LEU:O	6:Y:453:LEU:HG	2.12	0.49
6:Y:621:TRP:HE3	6:Y:626:LEU:HD21	1.77	0.49
6:Y:774:LEU:C	6:Y:776:GLN:H	2.21	0.49
6:Y:829:THR:HG23	6:Y:830:ASN:OD1	2.13	0.49
6:Y:906:ASN:ND2	6:Y:921:ASP:HB3	2.28	0.49
6:Y:1041:ILE:HG22	6:Y:1066:MET:HG2	1.95	0.49
1:b:21:ARG:NH1	3:B:327:TYR:CZ	2.81	0.48
1:p:45:ARG:HD2	3:P:400:ALA:HB1	1.95	0.48
3:B:420:LEU:HD12	3:B:422:LEU:HD23	1.95	0.48
3:B:521:VAL:HG21	3:B:577:VAL:HG22	1.95	0.48
3:D:549:LEU:C	3:D:549:LEU:CD1	2.86	0.48
3:F:527:VAL:HA	3:F:533:ASP:OD2	2.13	0.48
3:H:82:PHE:O	3:H:86:LEU:HG	2.13	0.48
3:H:354:SER:C	3:H:356:THR:H	2.21	0.48
3:J:173:ASP:HB2	3:N:606:THR:HG22	1.94	0.48
3:J:274:ILE:HD12	3:J:514:LEU:HD23	1.94	0.48
3:J:420:LEU:HD12	3:J:422:LEU:HD23	1.95	0.48
3:J:427:GLU:O	3:J:428:ARG:C	2.56	0.48
3:L:82:PHE:O	3:L:86:LEU:HG	2.13	0.48
3:L:297:ALA:CB	3:L:460:MET:HB2	2.33	0.48
3:L:373:PHE:HD2	3:L:397:VAL:HG11	1.78	0.48
3:N:120:GLY:HA3	3:N:126:ALA:HA	1.95	0.48
3:N:358:LEU:HB2	3:N:417:PHE:HD1	1.77	0.48
3:P:174:ILE:HD11	3:P:178:SER:HA	1.95	0.48
3:P:239:ALA:HB2	2:Q:42:ASN:O	2.13	0.48
3:P:315:LEU:HD23	3:P:315:LEU:N	2.28	0.48
3:P:385:VAL:HG23	3:P:390:ASN:HB3	1.94	0.48
3:R:47:LEU:HB2	3:R:136:PHE:CE2	2.47	0.48
3:R:203:TYR:HB3	3:R:206:GLU:CB	2.39	0.48
3:R:358:LEU:CB	3:R:417:PHE:HE1	2.24	0.48
3:T:65:SER:O	3:T:66:ASP:HB3	2.12	0.48
4:U:72:LEU:O	4:U:76:LEU:HG	2.12	0.48
4:U:193:ALA:HB1	4:U:293:VAL:HG13	1.94	0.48
4:V:64:ARG:HH21	6:X:1027:LEU:CD1	2.25	0.48
4:V:209:ARG:HB2	4:V:290:VAL:HG11	1.95	0.48
5:W:38:ASN:HA	5:W:89:HIS:CE1	2.45	0.48
5:W:58:VAL:HG12	5:W:59:ILE:N	2.27	0.48

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
5:W:405:ARG:NH1	5:W:779:MET:HG3	2.27	0.48
5:W:836:THR:HG23	5:W:837:GLY:O	2.13	0.48
5:W:893:ALA:HB2	5:W:921:ILE:HD12	1.95	0.48
5:W:1037:VAL:HG11	5:W:1042:THR:HG21	1.95	0.48
5:W:1139:PRO:HD3	5:W:1153:ALA:HB2	1.95	0.48
5:W:1281:ALA:HB3	5:W:1284:TYR:CD2	2.47	0.48
6:X:373:ARG:HG3	6:X:382:SER:OG	2.13	0.48
6:X:377:ILE:HD11	6:X:436:LEU:CG	2.43	0.48
6:X:569:LEU:HD23	6:X:569:LEU:HA	1.70	0.48
6:X:743:THR:HB	6:X:822:MET:O	2.13	0.48
6:X:1094:LEU:HD23	6:X:1094:LEU:C	2.37	0.48
6:Y:743:THR:N	6:Y:744:PRO:CD	2.76	0.48
1:h:5:MET:O	1:h:5:MET:HG3	2.13	0.48
1:j:26:THR:HA	1:j:38:ALA:HA	1.95	0.48
1:n:21:ARG:HG2	1:n:21:ARG:NH1	2.13	0.48
1:p:24:LEU:CD1	1:p:24:LEU:N	2.76	0.48
1:p:26:THR:HA	1:p:38:ALA:HA	1.95	0.48
1:r:45:ARG:HG2	3:R:399:ALA:O	2.13	0.48
3:B:65:SER:O	3:B:66:ASP:HB3	2.12	0.48
3:B:78:MET:HG2	3:B:78:MET:O	2.13	0.48
3:B:274:ILE:HD12	3:B:514:LEU:HD23	1.94	0.48
3:B:358:LEU:HB2	3:B:417:PHE:HD1	1.77	0.48
3:D:300:LYS:HB3	3:D:302:GLU:OE1	2.13	0.48
3:D:315:LEU:HD23	3:D:315:LEU:N	2.28	0.48
3:D:494:THR:H	3:D:497:GLU:CB	2.19	0.48
3:D:522:GLY:HA3	3:H:288:SER:HB3	1.94	0.48
2:E:25:ASP:OD2	3:H:143:MET:SD	2.71	0.48
3:F:249:VAL:HG12	3:F:617:VAL:HB	1.94	0.48
3:F:300:LYS:HB3	3:F:302:GLU:OE1	2.13	0.48
3:H:47:LEU:HD23	3:H:71:TYR:HB2	1.95	0.48
3:H:192:ASP:O	3:H:193:ILE:C	2.56	0.48
3:H:355:GLY:O	3:H:416:ARG:HD2	2.13	0.48
3:J:203:TYR:HB3	3:J:206:GLU:CB	2.39	0.48
3:L:192:ASP:O	3:L:193:ILE:C	2.56	0.48
3:N:494:THR:H	3:N:497:GLU:CB	2.19	0.48
3:P:47:LEU:HD23	3:P:71:TYR:HB2	1.95	0.48
3:P:82:PHE:O	3:P:86:LEU:HG	2.13	0.48
3:P:118:TYR:HB3	3:P:126:ALA:O	2.12	0.48
3:P:381:PRO:O	3:P:384:GLN:HB3	2.13	0.48
3:P:549:LEU:C	3:P:549:LEU:CD1	2.87	0.48
3:P:575:MET:HE3	3:T:469:VAL:HG21	0.69	0.48

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:R:315:LEU:N	3:R:315:LEU:HD23	2.28	0.48
3:R:420:LEU:HD12	3:R:422:LEU:HD23	1.95	0.48
3:R:431:ILE:HG12	3:R:437:ILE:HD13	1.94	0.48
3:T:373:PHE:HA	3:T:441:GLY:HA2	1.95	0.48
4:U:89:GLY:HA2	4:U:103:VAL:HG23	1.94	0.48
4:V:271:SER:C	4:V:273:HIS:H	2.20	0.48
5:W:30:LEU:HD22	5:W:30:LEU:O	2.13	0.48
5:W:175:CYS:C	5:W:177:TYR:N	2.68	0.48
5:W:480:ARG:O	5:W:482:PRO:HD3	2.13	0.48
5:W:603:LEU:HD22	5:W:635:TYR:CD2	2.48	0.48
5:W:829:ALA:HB3	5:W:891:ASP:OD2	2.13	0.48
5:W:994:THR:O	5:W:995:CYS:SG	2.69	0.48
6:X:718:THR:CB	6:X:726:ARG:NH1	2.75	0.48
6:X:982:LEU:HD23	6:X:982:LEU:O	2.13	0.48
6:X:1013:VAL:HG12	6:X:1014:VAL:H	1.76	0.48
6:Y:38:SER:O	6:Y:40:ASN:N	2.46	0.48
6:Y:353:GLN:OE1	6:Y:353:GLN:CA	2.59	0.48
6:Y:721:VAL:HG13	6:Y:726:ARG:HB2	1.93	0.48
6:Y:1087:TYR:HA	6:Y:1119:ILE:O	2.13	0.48
6:Y:1088:TYR:CZ	6:Y:1120:PRO:HB3	2.48	0.48
1:d:26:THR:HA	1:d:38:ALA:HA	1.95	0.48
1:t:6:ILE:CD1	3:P:586:VAL:CB	2.90	0.48
1:t:51:PHE:CE2	1:t:71:CYS:O	2.66	0.48
2:A:5:GLN:O	2:A:6:THR:CB	2.61	0.48
3:B:381:PRO:O	3:B:384:GLN:HB3	2.13	0.48
3:B:527:VAL:HA	3:B:533:ASP:OD2	2.13	0.48
3:D:239:ALA:HB2	2:E:42:ASN:O	2.13	0.48
3:F:162:HIS:CD2	3:F:199:LEU:HB2	2.48	0.48
3:F:315:LEU:N	3:F:315:LEU:HD23	2.28	0.48
3:J:355:GLY:O	3:J:416:ARG:HD2	2.13	0.48
3:J:381:PRO:O	3:J:384:GLN:HB3	2.13	0.48
3:L:162:HIS:CD2	3:L:199:LEU:HB2	2.48	0.48
3:L:381:PRO:O	3:L:384:GLN:HB3	2.13	0.48
3:L:521:VAL:HG21	3:L:577:VAL:HG22	1.95	0.48
3:N:373:PHE:HA	3:N:441:GLY:HA2	1.95	0.48
3:P:49:ARG:NH1	3:P:115:THR:HG22	2.18	0.48
3:P:79:ARG:HB2	3:P:79:ARG:NH1	2.27	0.48
3:P:120:GLY:HA3	3:P:126:ALA:HA	1.95	0.48
3:P:355:GLY:O	3:P:416:ARG:HD2	2.13	0.48
3:P:522:GLY:HA3	3:T:288:SER:HB3	1.94	0.48
3:R:300:LYS:HB3	3:R:302:GLU:OE1	2.13	0.48

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:T:471:LEU:N	3:T:471:LEU:CD2	2.76	0.48
4:U:79:PHE:HE2	4:U:138:ILE:CD1	2.25	0.48
4:U:241:LEU:O	4:U:242:ASN:CB	2.60	0.48
4:U:245:GLN:HE21	4:U:245:GLN:CA	2.25	0.48
4:V:379:ILE:HD13	4:V:388:PRO:HD3	1.94	0.48
5:W:398:LEU:HD23	5:W:398:LEU:N	2.22	0.48
5:W:624:THR:C	5:W:626:ALA:N	2.71	0.48
5:W:733:ALA:CB	5:W:779:MET:HE1	2.38	0.48
5:W:1014:ASN:HD21	5:W:1213:SER:HB2	1.77	0.48
5:W:1070:VAL:HG11	5:W:1082:ALA:HB2	1.95	0.48
6:X:656:PHE:CD2	6:X:752:PRO:HB3	2.48	0.48
6:X:687:LEU:O	6:X:832:ILE:HD13	2.14	0.48
6:X:718:THR:CA	6:X:726:ARG:NH1	2.75	0.48
6:X:792:GLN:CB	6:X:931:ILE:HD13	2.43	0.48
6:X:1129:THR:HG21	6:X:1132:ARG:NH1	2.28	0.48
6:Y:80:THR:HG22	6:Y:80:THR:O	2.13	0.48
6:Y:266:PRO:CB	6:Y:1085:LEU:HD11	2.41	0.48
6:Y:543:THR:O	6:Y:543:THR:CG2	2.60	0.48
6:Y:633:PRO:O	6:Y:637:PRO:HA	2.14	0.48
1:b:19:ALA:O	1:b:20:GLY:O	2.32	0.48
1:h:59:HIS:HB3	3:H:398:SER:CB	2.25	0.48
1:p:15:ARG:HH12	3:P:406:ASN:HB3	1.65	0.48
3:B:162:HIS:CD2	3:B:199:LEU:HB2	2.48	0.48
3:B:164:ASP:C	3:B:166:ALA:N	2.70	0.48
3:B:203:TYR:HB3	3:B:206:GLU:CB	2.39	0.48
3:D:78:MET:O	3:D:78:MET:HG2	2.13	0.48
3:D:355:GLY:O	3:D:416:ARG:HD2	2.13	0.48
3:D:381:PRO:O	3:D:384:GLN:HB3	2.13	0.48
3:F:48:TRP:CE2	3:F:63:ILE:HD13	2.49	0.48
3:F:382:ILE:O	3:F:383:GLN:HB2	2.14	0.48
3:H:527:VAL:HA	3:H:533:ASP:OD2	2.13	0.48
3:J:48:TRP:CE2	3:J:63:ILE:HD13	2.49	0.48
3:J:82:PHE:O	3:J:86:LEU:HG	2.13	0.48
3:J:107:VAL:HG12	3:J:109:SER:H	1.77	0.48
3:J:154:ARG:HB3	3:J:154:ARG:CZ	2.44	0.48
3:J:192:ASP:O	3:J:193:ILE:C	2.56	0.48
3:J:330:GLN:HE21	3:J:332:GLY:CA	2.26	0.48
3:J:344:MET:HE2	3:J:502:LEU:CD1	2.37	0.48
3:N:47:LEU:HD23	3:N:71:TYR:HB2	1.95	0.48
3:N:174:ILE:HD11	3:N:178:SER:HA	1.95	0.48
3:T:107:VAL:HG12	3:T:109:SER:H	1.77	0.48

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:T:154:ARG:HB3	3:T:154:ARG:CZ	2.44	0.48
3:T:355:GLY:O	3:T:416:ARG:HD2	2.13	0.48
4:U:136:TRP:HZ2	4:U:293:VAL:HG12	1.77	0.48
4:U:166:MET:O	4:U:168:ALA:N	2.46	0.48
4:U:195:GLN:O	4:U:196:LEU:HB2	2.14	0.48
4:U:342:ARG:HD3	4:U:349:ALA:HA	1.95	0.48
4:U:376:ALA:O	4:U:378:PRO:HD3	2.12	0.48
5:W:111:ARG:HD2	5:W:139:PHE:HB3	1.94	0.48
5:W:193:THR:O	5:W:194:TYR:HB2	2.13	0.48
5:W:1179:GLU:HA	5:W:1179:GLU:OE1	2.13	0.48
5:W:1233:ARG:NH1	5:W:1293:VAL:HG12	2.28	0.48
6:X:419:ILE:HD13	6:X:663:GLN:HB2	1.94	0.48
6:X:593:LEU:HD21	6:X:597:GLU:HB2	1.95	0.48
6:Y:938:TYR:CZ	6:Y:946:VAL:HG21	2.49	0.48
6:Y:1097:ALA:HB3	6:Y:1098:PRO:HD3	1.93	0.48
1:l:21:ARG:NH2	3:L:327:TYR:HE1	2.12	0.48
1:l:53:LEU:HD12	1:l:74:GLN:OE1	2.13	0.48
1:b:45:ARG:N	3:B:399:ALA:O	2.47	0.48
1:d:24:LEU:HB3	1:d:38:ALA:CB	2.44	0.48
1:d:60:ALA:H	3:D:397:VAL:C	2.21	0.48
1:j:45:ARG:CD	3:J:400:ALA:CA	2.88	0.48
2:C:25:ASP:OD2	3:F:143:MET:SD	2.71	0.48
3:D:82:PHE:O	3:D:86:LEU:HG	2.13	0.48
3:D:164:ASP:C	3:D:166:ALA:N	2.71	0.48
3:D:527:VAL:HA	3:D:533:ASP:OD2	2.13	0.48
3:F:373:PHE:HA	3:F:441:GLY:HA2	1.95	0.48
2:G:5:GLN:O	2:G:6:THR:CB	2.61	0.48
3:H:96:PHE:CE1	3:H:159:TRP:CH2	3.02	0.48
2:I:25:ASP:OD2	3:L:143:MET:SD	2.71	0.48
3:J:527:VAL:HA	3:J:533:ASP:OD2	2.13	0.48
3:L:98:ALA:HB2	3:L:206:GLU:CD	2.39	0.48
3:L:288:SER:HB3	3:N:522:GLY:HA3	1.94	0.48
3:L:296:PRO:C	3:L:298:PHE:H	2.19	0.48
3:L:385:VAL:HG23	3:L:390:ASN:HB3	1.94	0.48
3:L:427:GLU:O	3:L:428:ARG:C	2.56	0.48
3:N:354:SER:C	3:N:356:THR:H	2.21	0.48
3:N:381:PRO:O	3:N:384:GLN:HB3	2.13	0.48
2:O:42:ASN:O	3:T:239:ALA:HB2	2.13	0.48
3:P:260:SER:HB3	2:Q:33:ALA:HB1	1.96	0.48
3:P:427:GLU:O	3:P:428:ARG:C	2.56	0.48
3:P:443:PHE:HE2	3:P:459:LEU:HD21	1.74	0.48

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:R:98:ALA:HB2	3:R:206:GLU:CD	2.39	0.48
3:R:469:VAL:O	3:R:469:VAL:HG13	2.13	0.48
3:T:360:TYR:CD1	3:T:361:ASP:N	2.77	0.48
3:T:427:GLU:O	3:T:428:ARG:C	2.56	0.48
4:U:10:THR:CG2	4:U:73:GLY:HA3	2.43	0.48
4:U:294:THR:HG23	4:U:294:THR:O	2.13	0.48
4:U:384:MET:N	4:U:385:PRO:HD3	2.28	0.48
5:W:1187:ALA:N	5:W:1188:PRO:CD	2.77	0.48
5:W:1258:MET:HG3	5:W:1264:PRO:HG2	1.94	0.48
6:X:248:ASN:O	6:X:249:TRP:C	2.55	0.48
6:X:603:VAL:HG13	6:X:620:VAL:HG12	1.95	0.48
6:X:973:ASN:ND2	6:X:1144:ASP:HB2	2.29	0.48
6:X:1124:VAL:O	6:X:1157:ALA:HB2	2.13	0.48
6:Y:54:VAL:CG2	6:Y:57:GLN:HE21	2.26	0.48
6:Y:649:ASN:ND2	6:Y:699:ASN:HD21	2.12	0.48
6:Y:656:PHE:HD1	6:Y:657:GLY:O	1.96	0.48
1:j:6:ILE:HD11	3:L:586:VAL:CB	2.44	0.48
1:n:61:ASN:HB3	3:N:397:VAL:HA	1.94	0.48
1:p:4:HIS:CD2	1:p:54:THR:CG2	2.83	0.48
1:p:8:GLN:NE2	3:R:303:ASP:CA	2.62	0.48
1:t:26:THR:HA	1:t:38:ALA:HA	1.95	0.48
1:t:59:HIS:N	1:t:59:HIS:ND1	2.61	0.48
3:B:98:ALA:HB2	3:B:206:GLU:CD	2.39	0.48
3:B:154:ARG:HB3	3:B:154:ARG:CZ	2.44	0.48
3:D:52:GLY:O	3:D:53:THR:OG1	2.31	0.48
3:D:192:ASP:O	3:D:193:ILE:C	2.56	0.48
3:F:120:GLY:HA3	3:F:126:ALA:HA	1.95	0.48
3:F:420:LEU:HD12	3:F:422:LEU:HD23	1.95	0.48
3:H:96:PHE:CD1	3:H:101:THR:HG23	2.44	0.48
3:H:549:LEU:C	3:H:549:LEU:CD1	2.87	0.48
3:J:162:HIS:CD2	3:J:199:LEU:HB2	2.48	0.48
3:J:238:VAL:HG21	3:L:78:MET:HE2	1.83	0.48
3:J:315:LEU:N	3:J:315:LEU:HD23	2.28	0.48
3:L:48:TRP:N	3:L:48:TRP:HD1	2.09	0.48
3:L:48:TRP:CE2	3:L:63:ILE:HD13	2.49	0.48
3:L:315:LEU:HD23	3:L:315:LEU:N	2.28	0.48
3:L:344:MET:HE2	3:L:502:LEU:CD1	2.37	0.48
3:L:420:LEU:HD12	3:L:422:LEU:HD23	1.95	0.48
3:N:296:PRO:C	3:N:298:PHE:N	2.68	0.48
3:N:330:GLN:HE21	3:N:332:GLY:CA	2.26	0.48
3:N:527:VAL:HA	3:N:533:ASP:OD2	2.13	0.48

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:P:162:HIS:CD2	3:P:199:LEU:HB2	2.48	0.48
3:P:300:LYS:HB3	3:P:302:GLU:OE1	2.13	0.48
3:P:373:PHE:HA	3:P:441:GLY:HA2	1.95	0.48
3:R:78:MET:HG2	3:R:78:MET:O	2.13	0.48
3:R:120:GLY:HA3	3:R:126:ALA:HA	1.95	0.48
3:R:238:VAL:CG2	3:T:142:ILE:HD11	2.42	0.48
3:R:381:PRO:O	3:R:384:GLN:HB3	2.13	0.48
3:R:471:LEU:CD2	3:R:471:LEU:N	2.76	0.48
3:R:549:LEU:HB3	3:T:644:LEU:CD1	2.34	0.48
3:T:315:LEU:HD23	3:T:315:LEU:N	2.28	0.48
4:U:41:PHE:CZ	4:U:186:ALA:HB3	2.49	0.48
4:U:339:LEU:CG	4:U:343:GLN:HE21	2.26	0.48
4:V:337:ARG:HE	4:V:338:HIS:CD2	2.32	0.48
5:W:409:ILE:HB	5:W:414:ALA:HB2	1.95	0.48
5:W:519:LEU:HG	5:W:580:ALA:HA	1.94	0.48
5:W:582:LEU:CB	5:W:602:MET:HE1	2.44	0.48
5:W:610:THR:OG1	5:W:662:ARG:HG2	2.13	0.48
5:W:860:CYS:HB2	5:W:874:TYR:CE1	2.49	0.48
5:W:1040:VAL:CB	5:W:1041:PRO:CD	2.88	0.48
5:W:1157:CYS:HB3	5:W:1190:ARG:NH2	2.28	0.48
6:X:373:ARG:HA	6:X:385:VAL:HA	1.95	0.48
6:X:415:GLN:C	6:X:417:VAL:N	2.70	0.48
6:X:681:ASN:HD22	6:X:752:PRO:CG	2.27	0.48
6:X:921:ASP:O	6:X:923:LEU:N	2.44	0.48
6:Y:165:ASP:C	6:Y:166:ASP:CG	2.81	0.48
6:Y:559:HIS:HD2	6:Y:562:SER:CB	2.25	0.48
6:Y:587:MET:CE	6:Y:601:ILE:HD11	2.42	0.48
6:Y:1037:VAL:CG1	6:Y:1039:ARG:HH21	2.26	0.48
1:l:4:HIS:CD2	1:l:54:THR:HG22	2.49	0.48
1:f:15:ARG:CZ	3:F:402:ALA:HB2	2.40	0.48
1:f:50:ALA:O	1:f:74:GLN:HB3	2.14	0.48
1:j:45:ARG:CG	3:J:399:ALA:O	2.61	0.48
1:t:58:PRO:HD2	1:t:59:HIS:HE1	1.76	0.48
3:B:120:GLY:HA3	3:B:126:ALA:HA	1.95	0.48
3:B:315:LEU:N	3:B:315:LEU:HD23	2.28	0.48
3:B:382:ILE:O	3:B:383:GLN:HB2	2.14	0.48
7:C:101:MYR:H72	3:D:200:CYS:HG	1.79	0.48
3:D:47:LEU:HD23	3:D:71:TYR:HB2	1.95	0.48
3:D:58:ILE:O	3:D:59:ASP:HB3	2.13	0.48
3:D:120:GLY:HA3	3:D:126:ALA:HA	1.95	0.48
3:D:162:HIS:CD2	3:D:199:LEU:HB2	2.48	0.48

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:D:288:SER:HB3	3:F:522:GLY:HA3	1.95	0.48
3:D:354:SER:C	3:D:356:THR:H	2.21	0.48
3:D:373:PHE:HA	3:D:441:GLY:HA2	1.95	0.48
2:E:28:SER:HB2	3:F:215:TYR:HA	1.96	0.48
3:F:98:ALA:HB2	3:F:206:GLU:CD	2.39	0.48
3:F:381:PRO:O	3:F:384:GLN:HB3	2.13	0.48
3:F:471:LEU:CD2	3:F:471:LEU:N	2.76	0.48
3:H:164:ASP:C	3:H:166:ALA:N	2.71	0.48
3:H:174:ILE:HD11	3:H:178:SER:HA	1.95	0.48
3:H:204:PRO:O	3:H:208:TYR:HB2	2.14	0.48
3:H:360:TYR:CD1	3:H:361:ASP:N	2.77	0.48
3:H:471:LEU:N	3:H:471:LEU:CD2	2.76	0.48
2:I:5:GLN:O	2:I:6:THR:CB	2.61	0.48
3:J:120:GLY:HA3	3:J:126:ALA:HA	1.95	0.48
3:J:204:PRO:O	3:J:208:TYR:HB2	2.14	0.48
2:K:26:MET:O	2:K:26:MET:CG	2.60	0.48
3:L:96:PHE:CE1	3:L:159:TRP:CH2	3.02	0.48
3:L:354:SER:C	3:L:356:THR:H	2.21	0.48
3:L:355:GLY:O	3:L:416:ARG:HD2	2.13	0.48
3:N:249:VAL:HG12	3:N:617:VAL:HB	1.94	0.48
3:P:48:TRP:N	3:P:48:TRP:HD1	2.09	0.48
3:P:98:ALA:HB2	3:P:206:GLU:CD	2.39	0.48
3:P:288:SER:HB3	3:R:522:GLY:HA3	1.94	0.48
3:P:298:PHE:HE1	3:T:416:ARG:CB	2.16	0.48
3:R:330:GLN:HE21	3:R:332:GLY:CA	2.26	0.48
3:R:373:PHE:HA	3:R:441:GLY:HA2	1.95	0.48
3:T:98:ALA:HB2	3:T:206:GLU:CD	2.39	0.48
3:T:204:PRO:O	3:T:208:TYR:HB2	2.14	0.48
3:T:344:MET:HE2	3:T:502:LEU:CD1	2.37	0.48
3:T:420:LEU:HD12	3:T:422:LEU:HD23	1.96	0.48
3:T:527:VAL:HA	3:T:533:ASP:OD2	2.13	0.48
3:T:564:LEU:HA	3:T:567:ARG:NH2	2.27	0.48
5:W:19:ARG:NH1	5:W:366:VAL:O	2.47	0.48
5:W:63:PRO:C	5:W:65:LEU:H	2.20	0.48
5:W:740:VAL:O	5:W:742:SER:N	2.47	0.48
5:W:773:ARG:CD	5:W:843:LEU:HD22	2.43	0.48
5:W:893:ALA:CB	5:W:921:ILE:HD12	2.43	0.48
5:W:1080:SER:HB3	5:W:1085:ALA:HB3	1.94	0.48
5:W:1233:ARG:HH11	5:W:1293:VAL:HG12	1.78	0.48
6:X:410:HIS:CG	6:X:411:GLN:N	2.80	0.48
6:Y:321:VAL:O	6:Y:321:VAL:HG12	2.13	0.48

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
6:Y:549:ALA:HB1	6:Y:569:LEU:HD13	1.96	0.48
6:Y:996:MET:O	6:Y:998:PRO:HD3	2.14	0.48
1:l:8:GLN:NE2	3:N:303:ASP:HA	2.28	0.48
1:b:22:LEU:HD22	1:b:22:LEU:H	1.77	0.48
1:r:25:TYR:CG	1:r:62:VAL:HG21	2.49	0.48
2:A:26:MET:O	2:A:26:MET:CG	2.60	0.48
3:D:411:VAL:O	3:D:412:ARG:CB	2.56	0.48
3:F:204:PRO:O	3:F:208:TYR:HB2	2.14	0.48
3:H:322:TYR:C	3:H:323:SER:HG	2.15	0.48
3:H:382:ILE:O	3:H:383:GLN:HB2	2.14	0.48
3:J:59:ASP:OD1	3:J:59:ASP:O	2.32	0.48
3:L:527:VAL:HA	3:L:533:ASP:OD2	2.13	0.48
3:L:561:ASP:O	3:L:563:PRO:CD	2.62	0.48
2:M:28:SER:HB2	3:N:215:TYR:HA	1.96	0.48
3:N:192:ASP:O	3:N:193:ILE:C	2.56	0.48
3:P:373:PHE:HD2	3:P:397:VAL:HG11	1.78	0.48
3:P:521:VAL:HG21	3:P:577:VAL:HG22	1.95	0.48
3:P:527:VAL:HA	3:P:533:ASP:OD2	2.13	0.48
3:P:561:ASP:O	3:P:563:PRO:CD	2.62	0.48
3:P:644:LEU:CD1	3:T:549:LEU:HB3	2.33	0.48
3:R:48:TRP:CE2	3:R:63:ILE:HD13	2.49	0.48
3:R:192:ASP:O	3:R:193:ILE:C	2.56	0.48
3:R:493:GLN:O	3:R:586:VAL:HG23	2.14	0.48
3:T:48:TRP:CE2	3:T:63:ILE:HD13	2.49	0.48
3:T:96:PHE:CE1	3:T:159:TRP:CH2	3.02	0.48
3:T:443:PHE:HE2	3:T:459:LEU:HD21	1.75	0.48
4:U:58:TYR:HB2	4:U:64:ARG:NH1	2.27	0.48
4:V:77:ASN:OD1	4:V:83:GLN:HB2	2.14	0.48
5:W:300:PHE:CE1	5:W:309:ASN:ND2	2.82	0.48
5:W:821:GLU:CB	5:W:824:ARG:HH21	2.26	0.48
5:W:1081:VAL:HG12	5:W:1081:VAL:O	2.14	0.48
5:W:1244:THR:HA	5:W:1284:TYR:HE2	1.78	0.48
6:X:518:LEU:HG	6:X:820:VAL:HG23	1.96	0.48
6:X:1017:PHE:CD1	6:Y:1211:ARG:HA	2.49	0.48
6:Y:231:MET:SD	6:Y:348:ALA:HB3	2.53	0.48
6:Y:294:ARG:HB3	6:Y:890:GLN:CB	2.44	0.48
6:Y:439:THR:O	6:Y:440:ILE:HB	2.13	0.48
6:Y:676:VAL:HG12	6:Y:771:PHE:HD2	1.78	0.48
6:Y:707:PHE:O	6:Y:711:VAL:HG23	2.13	0.48
1:d:24:LEU:HD13	1:d:41:VAL:HG22	1.96	0.48
1:n:59:HIS:HB3	3:N:398:SER:HB2	1.92	0.48

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:t:6:ILE:HD11	3:P:586:VAL:CB	2.43	0.48
1:t:24:LEU:O	1:t:38:ALA:HB1	2.14	0.48
3:B:561:ASP:O	3:B:563:PRO:CD	2.62	0.48
3:D:96:PHE:CE1	3:D:159:TRP:CH2	3.02	0.48
3:D:561:ASP:O	3:D:563:PRO:CD	2.62	0.48
3:F:96:PHE:CE1	3:F:159:TRP:CH2	3.02	0.48
3:F:549:LEU:HB3	3:H:644:LEU:CD1	2.34	0.48
3:H:48:TRP:CE2	3:H:63:ILE:HD13	2.49	0.48
3:H:381:PRO:O	3:H:384:GLN:HB3	2.13	0.48
3:H:633:ALA:O	3:H:636:ALA:HB3	2.14	0.48
2:I:42:ASN:O	3:N:239:ALA:HB2	2.13	0.48
3:J:354:SER:C	3:J:356:THR:H	2.21	0.48
3:J:373:PHE:HA	3:J:441:GLY:HA2	1.95	0.48
3:J:521:VAL:HG21	3:J:577:VAL:HG22	1.95	0.48
3:J:536:THR:HG21	3:J:570:LYS:HD2	1.94	0.48
3:L:166:ALA:CB	3:L:194:LEU:HD11	2.44	0.48
2:M:5:GLN:O	2:M:6:THR:CB	2.61	0.48
3:N:427:GLU:O	3:N:428:ARG:C	2.56	0.48
3:N:549:LEU:C	3:N:549:LEU:CD1	2.87	0.48
3:P:48:TRP:CE2	3:P:63:ILE:HD13	2.49	0.48
2:S:28:SER:HB2	3:T:215:TYR:HA	1.96	0.48
4:U:80:ALA:HB1	4:U:118:VAL:HG21	1.95	0.48
4:U:121:ASP:O	4:U:123:ASN:N	2.47	0.48
4:V:94:ASP:HB2	4:V:95:PRO:CD	2.43	0.48
4:V:188:THR:HG23	6:Y:376:LEU:HD13	1.95	0.48
5:W:59:ILE:HB	5:W:178:ILE:CD1	2.43	0.48
5:W:1160:TYR:HE2	5:W:1218:ARG:HH12	1.62	0.48
6:X:411:GLN:HG3	6:X:445:PHE:CD2	2.48	0.48
6:X:789:ASP:HB3	6:X:933:ASP:CG	2.39	0.48
6:X:855:GLN:C	6:X:857:ALA:N	2.67	0.48
6:X:1167:LEU:HD13	6:X:1187:ARG:NH2	2.29	0.48
6:Y:374:ALA:O	6:Y:375:ASN:HB2	2.12	0.48
6:Y:590:ALA:O	6:Y:608:GLN:HG2	2.13	0.48
1:f:26:THR:HA	1:f:38:ALA:HA	1.95	0.48
1:p:61:ASN:CB	3:P:395:THR:HG22	2.43	0.48
1:t:15:ARG:HE	3:T:406:ASN:CG	2.21	0.48
3:B:96:PHE:CD1	3:B:101:THR:HG23	2.44	0.48
3:B:192:ASP:O	3:B:193:ILE:C	2.56	0.48
3:B:471:LEU:CD2	3:B:471:LEU:N	2.76	0.48
2:E:26:MET:O	2:E:26:MET:CG	2.60	0.48
3:F:352:ASP:HB2	3:F:421:HIS:CD2	2.49	0.48

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:F:561:ASP:O	3:F:563:PRO:CD	2.62	0.48
3:F:633:ALA:O	3:F:636:ALA:HB3	2.14	0.48
2:G:28:SER:HB2	3:H:215:TYR:HA	1.96	0.48
3:H:98:ALA:HB2	3:H:206:GLU:CD	2.39	0.48
3:H:154:ARG:HB3	3:H:154:ARG:CZ	2.44	0.48
3:H:431:ILE:CD1	3:H:437:ILE:HG23	2.39	0.48
3:H:493:GLN:O	3:H:586:VAL:HG23	2.14	0.48
3:J:98:ALA:HB2	3:J:206:GLU:CD	2.39	0.48
3:J:396:VAL:HG11	3:J:399:ALA:HB2	1.96	0.48
3:L:78:MET:O	3:L:78:MET:HG2	2.13	0.48
3:L:396:VAL:HG11	3:L:399:ALA:HB2	1.96	0.48
3:N:96:PHE:CE1	3:N:159:TRP:CH2	3.02	0.48
3:N:315:LEU:N	3:N:315:LEU:HD23	2.28	0.48
3:N:373:PHE:HD2	3:N:397:VAL:HG11	1.78	0.48
3:N:396:VAL:HG11	3:N:399:ALA:HB2	1.96	0.48
3:N:471:LEU:CD2	3:N:471:LEU:N	2.76	0.48
3:P:305:LYS:HG3	3:P:477:LYS:O	2.14	0.48
3:P:396:VAL:HG11	3:P:399:ALA:HB2	1.96	0.48
7:Q:101:MYR:H21	3:R:191:LYS:NZ	2.24	0.48
3:R:354:SER:C	3:R:356:THR:H	2.21	0.48
4:U:131:ASN:O	4:U:132:LEU:C	2.56	0.48
4:U:161:VAL:CG2	4:U:410:MET:HG3	2.38	0.48
4:U:174:LEU:C	4:U:176:ALA:H	2.22	0.48
4:V:20:LEU:HD13	4:V:384:MET:HB2	1.96	0.48
4:V:140:GLN:O	4:V:144:ILE:HG12	2.14	0.48
5:W:313:LEU:O	5:W:318:VAL:CG1	2.61	0.48
5:W:493:VAL:HG21	5:W:656:PHE:CD2	2.49	0.48
5:W:765:PHE:CE1	5:W:766:LEU:HG	2.49	0.48
5:W:847:PRO:C	5:W:849:THR:H	2.22	0.48
5:W:1154:ARG:HD2	5:W:1189:ASN:ND2	2.16	0.48
6:X:362:THR:O	6:X:363:ALA:C	2.56	0.48
6:Y:750:MET:HB3	6:Y:837:TYR:HH	1.79	0.48
6:Y:925:LEU:O	6:Y:926:TYR:HB3	2.13	0.48
6:Y:1080:HIS:O	6:Y:1081:TYR:HB3	2.14	0.48
1:b:24:LEU:HD22	1:b:80:ARG:CD	2.41	0.47
1:b:61:ASN:ND2	1:b:61:ASN:C	2.72	0.47
1:p:15:ARG:NH2	3:P:402:ALA:HB2	2.25	0.47
1:r:23:THR:HG22	1:r:25:TYR:CE2	2.49	0.47
3:B:48:TRP:CE2	3:B:63:ILE:HD13	2.49	0.47
3:B:204:PRO:O	3:B:208:TYR:HB2	2.14	0.47
3:D:423:GLN:O	3:D:423:GLN:CG	2.62	0.47

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:F:192:ASP:O	3:F:193:ILE:C	2.56	0.47
3:F:358:LEU:HB2	3:F:417:PHE:HD1	1.77	0.47
3:F:396:VAL:HG11	3:F:399:ALA:HB2	1.96	0.47
3:H:251:SER:OG	3:H:253:VAL:HG12	2.14	0.47
3:H:396:VAL:HG11	3:H:399:ALA:HB2	1.96	0.47
3:J:55:VAL:HG22	3:J:56:ALA:N	2.29	0.47
3:J:166:ALA:CB	3:J:194:LEU:HD11	2.44	0.47
3:J:352:ASP:HB2	3:J:421:HIS:CD2	2.49	0.47
3:J:423:GLN:O	3:J:423:GLN:CG	2.62	0.47
3:J:427:GLU:CD	3:L:571:ARG:HH22	2.20	0.47
3:L:47:LEU:HD23	3:L:71:TYR:HB2	1.95	0.47
3:L:443:PHE:HE2	3:L:459:LEU:HD21	1.74	0.47
3:L:471:LEU:N	3:L:471:LEU:CD2	2.76	0.47
3:L:549:LEU:C	3:L:549:LEU:CD1	2.87	0.47
2:M:26:MET:CG	3:N:216:PRO:HG3	2.29	0.47
3:N:78:MET:HG2	3:N:78:MET:O	2.13	0.47
3:P:51:VAL:CG1	5:W:379:VAL:CG2	2.92	0.47
3:P:57:THR:O	3:P:58:ILE:CB	2.62	0.47
3:P:96:PHE:CE1	3:P:159:TRP:CH2	3.02	0.47
3:P:493:GLN:O	3:P:586:VAL:HG23	2.14	0.47
2:Q:28:SER:HB2	3:R:215:TYR:HA	1.96	0.47
3:R:96:PHE:CE1	3:R:159:TRP:CH2	3.02	0.47
3:R:305:LYS:HG3	3:R:477:LYS:O	2.14	0.47
3:T:80:GLU:OE2	3:T:80:GLU:HA	2.12	0.47
3:T:166:ALA:CB	3:T:194:LEU:HD11	2.44	0.47
5:W:30:LEU:HD13	5:W:30:LEU:C	2.39	0.47
5:W:95:PHE:HB2	5:W:125:PHE:CD2	2.49	0.47
5:W:105:LEU:HD23	5:W:105:LEU:C	2.38	0.47
5:W:439:PHE:HB3	5:W:648:THR:OG1	2.12	0.47
5:W:1102:PRO:HG3	5:W:1127:ILE:CD1	2.43	0.47
5:W:1192:LEU:HD13	5:W:1192:LEU:C	2.38	0.47
6:X:389:GLU:HG2	6:X:1194:LEU:HD21	1.95	0.47
6:X:393:ALA:HB1	6:X:1182:ASN:HD22	1.78	0.47
6:X:784:LEU:N	6:X:784:LEU:CD1	2.74	0.47
6:X:935:SER:CB	6:X:949:HIS:HB3	2.44	0.47
6:X:976:ASN:CA	6:X:1083:TYR:HE1	2.26	0.47
6:Y:302:LEU:CD2	6:Y:905:VAL:HG22	2.43	0.47
6:Y:842:ALA:HB3	6:Y:843:PRO:CD	2.34	0.47
1:h:15:ARG:HH12	3:H:402:ALA:CB	2.23	0.47
1:p:44:GLY:C	3:P:399:ALA:C	2.82	0.47
3:B:166:ALA:CB	3:B:194:LEU:HD11	2.44	0.47

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:B:330:GLN:HE21	3:B:332:GLY:CA	2.26	0.47
3:B:373:PHE:HD2	3:B:397:VAL:HG11	1.78	0.47
2:C:5:GLN:O	2:C:6:THR:CB	2.61	0.47
3:D:433:GLY:O	3:D:435:PRO:HD3	2.14	0.47
2:E:5:GLN:O	2:E:6:THR:CB	2.61	0.47
3:F:373:PHE:HD2	3:F:397:VAL:HG11	1.78	0.47
3:F:427:GLU:CD	3:H:571:ARG:HH22	2.19	0.47
3:F:433:GLY:O	3:F:435:PRO:HD3	2.14	0.47
3:J:78:MET:HG2	3:J:78:MET:O	2.13	0.47
3:J:416:ARG:CB	3:L:298:PHE:HE1	2.16	0.47
3:J:443:PHE:HE2	3:J:459:LEU:HD21	1.75	0.47
3:J:549:LEU:HB3	3:L:644:LEU:CD1	2.33	0.47
3:L:305:LYS:HG3	3:L:477:LYS:O	2.14	0.47
3:N:98:ALA:HB2	3:N:206:GLU:CD	2.39	0.47
3:N:203:TYR:HB3	3:N:206:GLU:CB	2.39	0.47
3:N:469:VAL:O	3:N:469:VAL:HG13	2.13	0.47
2:O:28:SER:HA	3:P:216:PRO:CD	2.44	0.47
3:P:204:PRO:O	3:P:208:TYR:HB2	2.14	0.47
3:P:296:PRO:C	3:P:298:PHE:N	2.68	0.47
3:P:358:LEU:CB	3:P:417:PHE:HE1	2.25	0.47
3:P:471:LEU:N	3:P:471:LEU:CD2	2.76	0.47
3:R:352:ASP:HB2	3:R:421:HIS:CD2	2.49	0.47
3:R:373:PHE:HD2	3:R:397:VAL:HG11	1.78	0.47
3:R:423:GLN:O	3:R:423:GLN:CG	2.62	0.47
3:R:427:GLU:O	3:R:428:ARG:C	2.56	0.47
3:T:469:VAL:O	3:T:469:VAL:HG13	2.12	0.47
3:T:493:GLN:O	3:T:586:VAL:HG23	2.14	0.47
3:T:633:ALA:O	3:T:636:ALA:HB3	2.14	0.47
4:V:10:THR:HG23	4:V:73:GLY:HA3	1.95	0.47
4:V:12:ASN:ND2	4:V:14:TYR:H	2.12	0.47
5:W:192:ALA:CB	5:W:333:VAL:HG21	2.44	0.47
5:W:1052:ILE:CG2	5:W:1112:LEU:HD22	2.43	0.47
5:W:1054:SER:HB3	5:W:1096:PHE:CE1	2.49	0.47
5:W:1062:SER:O	5:W:1111:PHE:HB2	2.14	0.47
5:W:1206:TYR:CD2	5:W:1222:PRO:HB3	2.49	0.47
6:X:311:ILE:HD11	6:X:406:LEU:CD2	2.43	0.47
6:X:865:VAL:HG12	6:X:958:HIS:HB2	1.95	0.47
6:X:887:ALA:HB3	6:X:969:MET:HE1	1.95	0.47
6:Y:32:SER:OG	6:Y:70:SER:HB2	2.13	0.47
6:Y:617:HIS:CD2	6:Y:619:GLY:H	2.32	0.47
6:Y:1020:ILE:HD13	6:Y:1031:LEU:HA	1.96	0.47

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
6:Y:1049:SER:HB2	6:Y:1078:LEU:HG	1.96	0.47
1:l:16:ALA:CB	1:l:22:LEU:HD21	2.44	0.47
1:f:53:LEU:HD13	1:f:74:GLN:CD	2.38	0.47
1:j:21:ARG:O	3:J:323:SER:OG	2.31	0.47
1:t:15:ARG:NE	3:T:406:ASN:CG	2.72	0.47
1:t:45:ARG:CG	3:T:400:ALA:C	2.73	0.47
3:B:96:PHE:CE1	3:B:159:TRP:CH2	3.02	0.47
3:B:560:PRO:HB2	3:B:562:TYR:CE1	2.50	0.47
2:C:29:THR:HG22	3:H:609:ARG:HD2	1.91	0.47
3:D:203:TYR:HB3	3:D:206:GLU:CB	2.39	0.47
3:D:297:ALA:CB	3:D:460:MET:HB2	2.33	0.47
3:D:305:LYS:HG3	3:D:477:LYS:O	2.14	0.47
3:D:352:ASP:HB2	3:D:421:HIS:CD2	2.49	0.47
3:D:382:ILE:O	3:D:383:GLN:HB2	2.14	0.47
3:F:47:LEU:HD23	3:F:71:TYR:HB2	1.95	0.47
3:F:330:GLN:HE21	3:F:332:GLY:CA	2.26	0.47
3:F:564:LEU:CD2	3:F:568:ARG:NH2	2.75	0.47
3:H:305:LYS:HG3	3:H:477:LYS:O	2.14	0.47
3:H:433:GLY:O	3:H:435:PRO:HD3	2.14	0.47
3:H:560:PRO:HB2	3:H:562:TYR:CE1	2.50	0.47
3:H:561:ASP:O	3:H:562:TYR:CB	2.62	0.47
3:J:46:LYS:NZ	3:N:237:GLU:CD	2.68	0.47
3:J:373:PHE:HD2	3:J:397:VAL:HG11	1.78	0.47
3:J:560:PRO:HB2	3:J:562:TYR:CE1	2.50	0.47
3:L:154:ARG:HB3	3:L:154:ARG:CZ	2.44	0.47
3:N:365:ASP:OD2	3:N:452:SER:HB2	2.15	0.47
3:P:423:GLN:CG	3:P:423:GLN:O	2.62	0.47
3:T:354:SER:C	3:T:356:THR:H	2.21	0.47
3:T:406:ASN:O	3:T:406:ASN:ND2	2.43	0.47
4:U:367:GLU:OE2	4:U:394:PHE:HB3	2.14	0.47
4:V:7:PHE:O	4:V:315:LEU:HD21	2.14	0.47
5:W:8:GLN:HB2	5:W:321:GLU:HG2	1.95	0.47
5:W:226:VAL:HG11	5:W:239:LEU:HD22	1.95	0.47
5:W:910:LEU:HD13	5:W:910:LEU:C	2.40	0.47
5:W:993:PRO:HA	5:W:1017:ILE:HD13	1.96	0.47
6:X:474:GLU:HG2	6:X:474:GLU:O	2.14	0.47
6:X:633:PRO:CG	6:X:640:ARG:HH11	2.24	0.47
6:X:925:LEU:C	6:X:925:LEU:HD23	2.38	0.47
6:Y:46:ALA:O	6:Y:49:GLN:HG2	2.14	0.47
6:Y:105:SER:O	6:Y:106:MET:C	2.57	0.47
6:Y:202:THR:HG22	6:Y:392:ASP:H	1.79	0.47

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
6:Y:283:ILE:HG12	6:Y:287:GLN:O	2.15	0.47
6:Y:413:PRO:HB2	6:Y:434:ILE:CD1	2.39	0.47
6:Y:462:ILE:HG21	6:Y:469:ALA:HB2	1.95	0.47
6:Y:728:THR:C	6:Y:729:MET:HE2	2.40	0.47
6:Y:790:PRO:HG3	6:Y:793:ARG:NH2	2.30	0.47
1:b:58:PRO:HG2	3:B:423:GLN:CG	2.44	0.47
1:h:5:MET:O	1:h:6:ILE:O	2.32	0.47
1:r:8:GLN:HE21	3:T:302:GLU:CG	2.22	0.47
1:r:74:GLN:HE22	1:r:76:ASN:N	2.12	0.47
3:B:352:ASP:HB2	3:B:421:HIS:CD2	2.49	0.47
3:B:396:VAL:HG11	3:B:399:ALA:HB2	1.96	0.47
3:B:521:VAL:HG23	3:B:522:GLY:N	2.30	0.47
2:C:28:SER:HA	3:D:216:PRO:CD	2.44	0.47
3:D:174:ILE:HD11	3:D:178:SER:HA	1.95	0.47
3:D:204:PRO:O	3:D:208:TYR:HB2	2.14	0.47
3:D:471:LEU:N	3:D:471:LEU:CD2	2.76	0.47
3:D:560:PRO:HB2	3:D:562:TYR:CE1	2.50	0.47
3:F:78:MET:HG2	3:F:78:MET:O	2.13	0.47
3:H:166:ALA:CB	3:H:194:LEU:HD11	2.44	0.47
3:H:341:MET:CE	3:H:341:MET:CA	2.88	0.47
3:H:369:SER:OG	3:H:443:PHE:CD1	2.64	0.47
3:H:373:PHE:HA	3:H:441:GLY:HA2	1.95	0.47
3:H:427:GLU:O	3:H:428:ARG:C	2.56	0.47
3:H:469:VAL:O	3:H:469:VAL:HG13	2.13	0.47
3:J:96:PHE:CE1	3:J:159:TRP:CH2	3.02	0.47
3:J:174:ILE:HD11	3:J:178:SER:HA	1.95	0.47
3:J:365:ASP:OD2	3:J:452:SER:HB2	2.15	0.47
3:J:521:VAL:HG23	3:J:522:GLY:N	2.30	0.47
3:J:549:LEU:C	3:J:549:LEU:CD1	2.86	0.47
2:K:28:SER:HB2	3:L:215:TYR:HA	1.96	0.47
3:L:433:GLY:O	3:L:435:PRO:HD3	2.14	0.47
3:L:493:GLN:O	3:L:586:VAL:HG23	2.14	0.47
3:L:560:PRO:HB2	3:L:562:TYR:CE1	2.50	0.47
2:Q:28:SER:HA	3:R:216:PRO:CD	2.44	0.47
3:R:162:HIS:CD2	3:R:199:LEU:HB2	2.48	0.47
3:T:274:ILE:HD12	3:T:514:LEU:HD23	1.94	0.47
3:T:470:LEU:HD11	3:T:472:ARG:NH1	2.26	0.47
3:T:549:LEU:C	3:T:549:LEU:CD1	2.87	0.47
4:U:328:TYR:CG	4:U:333:MET:HE2	2.50	0.47
4:V:64:ARG:NH2	6:X:1027:LEU:CD1	2.78	0.47
4:V:250:TRP:CZ3	6:Y:956:HIS:CE1	3.02	0.47

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
4:V:299:LEU:HB2	4:V:322:VAL:HG12	1.97	0.47
4:V:348:THR:HG22	4:V:350:ALA:H	1.78	0.47
5:W:391:VAL:CG1	5:W:790:ARG:HH11	2.27	0.47
5:W:494:PHE:HA	5:W:497:LEU:HB2	1.96	0.47
5:W:998:LEU:C	5:W:1000:TYR:N	2.73	0.47
6:X:416:ILE:CD1	6:X:1207:ILE:HG21	2.45	0.47
6:Y:202:THR:HG21	6:Y:390:LEU:HD12	1.96	0.47
6:Y:317:PHE:H	6:Y:317:PHE:HD1	1.61	0.47
6:Y:325:THR:HG21	6:Y:347:ARG:HA	1.96	0.47
6:Y:589:LEU:O	6:Y:593:LEU:CD2	2.60	0.47
6:Y:618:PRO:CG	6:Y:779:VAL:HG13	2.44	0.47
6:Y:1020:ILE:HD11	6:Y:1041:ILE:HD11	1.95	0.47
6:Y:1078:LEU:HD12	6:Y:1078:LEU:N	2.29	0.47
6:Y:1088:TYR:CE1	6:Y:1132:ARG:HD3	2.50	0.47
1:l:26:THR:HA	1:l:38:ALA:HA	1.95	0.47
1:l:59:HIS:C	1:l:59:HIS:ND1	2.73	0.47
1:b:45:ARG:HD3	3:B:401:GLY:H	1.71	0.47
1:r:60:ALA:O	3:R:397:VAL:CA	2.63	0.47
1:t:44:GLY:O	3:T:399:ALA:O	2.32	0.47
2:A:13:ILE:HG13	3:B:208:TYR:CG	2.50	0.47
2:A:41:LEU:O	2:A:42:ASN:CB	2.37	0.47
3:B:633:ALA:O	3:B:636:ALA:HB3	2.14	0.47
3:D:274:ILE:HD12	3:D:514:LEU:HD23	1.94	0.47
3:F:521:VAL:HG23	3:F:522:GLY:N	2.30	0.47
3:H:315:LEU:HD23	3:H:315:LEU:N	2.28	0.47
3:H:420:LEU:HD12	3:H:422:LEU:HD23	1.95	0.47
3:J:251:SER:OG	3:J:253:VAL:HG12	2.14	0.47
3:J:369:SER:OG	3:J:443:PHE:CD1	2.64	0.47
3:J:471:LEU:CD2	3:J:471:LEU:N	2.76	0.47
3:J:493:GLN:O	3:J:586:VAL:HG23	2.14	0.47
3:J:633:ALA:O	3:J:636:ALA:HB3	2.14	0.47
2:K:13:ILE:HG13	3:L:208:TYR:CG	2.50	0.47
3:L:120:GLY:HA3	3:L:126:ALA:HA	1.95	0.47
3:L:164:ASP:C	3:L:166:ALA:N	2.71	0.47
3:L:174:ILE:HD11	3:L:178:SER:HA	1.95	0.47
3:L:251:SER:OG	3:L:253:VAL:HG12	2.14	0.47
3:N:305:LYS:HG3	3:N:477:LYS:O	2.14	0.47
3:N:355:GLY:O	3:N:416:ARG:HD2	2.13	0.47
3:N:521:VAL:HG23	3:N:522:GLY:N	2.30	0.47
7:O:101:MYR:H72	3:P:200:CYS:HG	1.75	0.47
3:P:78:MET:HG2	3:P:78:MET:O	2.13	0.47

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:P:352:ASP:HB2	3:P:421:HIS:CD2	2.49	0.47
3:P:382:ILE:O	3:P:383:GLN:HB2	2.14	0.47
3:P:433:GLY:O	3:P:435:PRO:HD3	2.14	0.47
3:R:166:ALA:CB	3:R:194:LEU:HD11	2.44	0.47
3:R:355:GLY:O	3:R:416:ARG:HD2	2.13	0.47
3:R:365:ASP:OD2	3:R:452:SER:HB2	2.15	0.47
3:R:633:ALA:O	3:R:636:ALA:HB3	2.14	0.47
2:S:28:SER:HA	3:T:216:PRO:CD	2.44	0.47
3:T:78:MET:HG2	3:T:78:MET:O	2.13	0.47
3:T:352:ASP:HB2	3:T:421:HIS:CD2	2.49	0.47
4:U:12:ASN:HD22	4:U:12:ASN:C	2.21	0.47
4:U:318:ASN:H	4:U:322:VAL:CG1	2.26	0.47
5:W:620:VAL:HG23	5:W:657:LEU:CD1	2.45	0.47
5:W:642:LEU:HD12	5:W:668:LEU:HD13	1.96	0.47
5:W:1163:LEU:HD13	5:W:1207:LEU:HB3	1.95	0.47
6:X:242:ARG:NH2	6:X:1085:LEU:HD11	2.23	0.47
6:X:263:LEU:HD22	6:X:263:LEU:N	2.29	0.47
6:X:298:LEU:O	6:X:298:LEU:HD23	2.13	0.47
6:X:419:ILE:HA	6:X:660:SER:CB	2.44	0.47
6:X:578:ASP:HA	6:X:579:PRO:HD3	1.75	0.47
6:Y:250:ALA:HA	6:Y:336:ARG:NH2	2.30	0.47
6:Y:355:LYS:CE	6:Y:1161:ASN:OD1	2.63	0.47
6:Y:632:ASN:HB2	6:Y:635:GLN:CB	2.30	0.47
6:Y:1048:PRO:HG2	6:Y:1051:VAL:HG23	1.95	0.47
6:Y:1088:TYR:O	6:Y:1121:ILE:HG13	2.14	0.47
1:l:23:THR:HG23	1:l:25:TYR:CD2	2.50	0.47
1:b:58:PRO:CG	3:B:423:GLN:CG	2.92	0.47
1:j:24:LEU:N	1:j:24:LEU:HD22	2.30	0.47
1:j:45:ARG:HG2	3:J:399:ALA:O	2.13	0.47
1:n:71:CYS:SG	1:n:72:SER:N	2.87	0.47
2:A:28:SER:HA	3:B:216:PRO:CD	2.44	0.47
3:B:382:ILE:HB	3:B:438:TYR:CZ	2.50	0.47
3:B:423:GLN:O	3:B:423:GLN:CG	2.62	0.47
3:D:98:ALA:HB2	3:D:206:GLU:CD	2.39	0.47
3:D:633:ALA:O	3:D:636:ALA:HB3	2.14	0.47
3:F:237:GLU:CD	3:H:46:LYS:NZ	2.68	0.47
3:F:354:SER:C	3:F:356:THR:H	2.21	0.47
3:J:49:ARG:NH1	4:V:405:ALA:HB1	2.29	0.47
3:J:227:LEU:O	3:J:228:PRO:O	2.33	0.47
3:J:298:PHE:HE1	3:N:416:ARG:CB	2.16	0.47
3:L:204:PRO:O	3:L:208:TYR:HB2	2.14	0.47

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:L:358:LEU:HB2	3:L:417:PHE:HD1	1.77	0.47
3:L:382:ILE:HB	3:L:438:TYR:CZ	2.50	0.47
3:L:382:ILE:O	3:L:383:GLN:HB2	2.14	0.47
3:N:369:SER:HB2	3:N:445:ILE:HA	1.97	0.47
3:N:420:LEU:HD12	3:N:422:LEU:HD23	1.96	0.47
3:N:493:GLN:O	3:N:586:VAL:HG23	2.14	0.47
3:N:561:ASP:O	3:N:562:TYR:CB	2.62	0.47
2:O:5:GLN:O	2:O:6:THR:CB	2.61	0.47
3:P:166:ALA:CB	3:P:194:LEU:HD11	2.44	0.47
3:P:365:ASP:OD2	3:P:452:SER:HB2	2.15	0.47
3:P:382:ILE:HB	3:P:438:TYR:CZ	2.50	0.47
3:R:433:GLY:O	3:R:435:PRO:HD3	2.15	0.47
3:T:300:LYS:HB3	3:T:302:GLU:OE1	2.13	0.47
3:T:382:ILE:HB	3:T:438:TYR:CZ	2.50	0.47
3:T:560:PRO:HB2	3:T:562:TYR:CE1	2.50	0.47
3:T:564:LEU:CD2	3:T:568:ARG:NH2	2.75	0.47
4:U:92:TRP:CD1	4:U:97:LEU:HD22	2.50	0.47
4:V:8:GLY:HA2	4:V:146:THR:OG1	2.15	0.47
5:W:398:LEU:CD1	5:W:790:ARG:HH21	2.27	0.47
5:W:453:PRO:HB2	5:W:456:GLN:CG	2.44	0.47
6:X:192:GLU:O	6:X:192:GLU:CG	2.62	0.47
6:X:429:ASN:HD22	6:X:429:ASN:H	1.57	0.47
6:X:470:LEU:N	6:X:471:PRO:CD	2.78	0.47
6:X:589:LEU:HD23	6:X:589:LEU:C	2.40	0.47
6:X:660:SER:C	6:X:673:ASN:ND2	2.73	0.47
6:X:704:THR:HG23	6:X:705:ILE:N	2.30	0.47
6:X:732:VAL:CG1	6:X:732:VAL:O	2.62	0.47
6:X:734:THR:HG22	6:X:738:GLN:OE1	2.15	0.47
6:X:1058:VAL:O	6:X:1059:PHE:C	2.58	0.47
6:Y:164:TRP:CG	6:Y:165:ASP:N	2.82	0.47
6:Y:288:PHE:O	6:Y:290:PRO:HD3	2.14	0.47
6:Y:778:ASP:C	6:Y:940:GLN:HB3	2.40	0.47
6:Y:998:PRO:HG3	6:Y:1138:THR:O	2.15	0.47
6:Y:1024:PRO:O	6:Y:1025:ASN:HB3	2.14	0.47
1:b:21:ARG:NH1	3:B:327:TYR:CE1	2.82	0.47
1:b:26:THR:HA	1:b:38:ALA:HA	1.95	0.47
1:d:74:GLN:NE2	1:d:74:GLN:C	2.72	0.47
1:n:47:THR:CG2	1:n:60:ALA:HA	2.45	0.47
1:n:58:PRO:CG	3:N:423:GLN:CG	2.92	0.47
1:p:6:ILE:HG21	3:R:494:THR:CG2	2.43	0.47
1:t:46:TYR:CD1	1:t:58:PRO:HA	2.46	0.47

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
7:A:101:MYR:C4	3:B:187:LEU:HD22	2.45	0.47
3:B:48:TRP:N	3:B:48:TRP:HD1	2.09	0.47
3:B:355:GLY:O	3:B:416:ARG:HD2	2.13	0.47
3:B:358:LEU:CB	3:B:417:PHE:HE1	2.25	0.47
3:B:369:SER:HB2	3:B:445:ILE:HA	1.97	0.47
2:C:28:SER:HB2	3:D:215:TYR:HA	1.96	0.47
3:D:166:ALA:CB	3:D:194:LEU:HD11	2.44	0.47
3:D:251:SER:OG	3:D:253:VAL:HG12	2.14	0.47
3:D:365:ASP:OD2	3:D:452:SER:HB2	2.15	0.47
3:D:561:ASP:O	3:D:562:TYR:CB	2.62	0.47
3:F:48:TRP:N	3:F:48:TRP:HD1	2.09	0.47
3:F:166:ALA:CB	3:F:194:LEU:HD11	2.44	0.47
3:F:365:ASP:OD2	3:F:452:SER:HB2	2.15	0.47
3:F:423:GLN:CG	3:F:423:GLN:O	2.62	0.47
3:F:436:TYR:HH	3:N:435:PRO:HG2	1.74	0.47
3:F:493:GLN:O	3:F:586:VAL:HG23	2.14	0.47
3:H:352:ASP:HB2	3:H:421:HIS:CD2	2.49	0.47
3:H:382:ILE:HB	3:H:438:TYR:CZ	2.49	0.47
3:H:423:GLN:CG	3:H:423:GLN:O	2.62	0.47
3:H:564:LEU:HD23	3:H:564:LEU:C	2.40	0.47
2:I:28:SER:HA	3:J:216:PRO:CD	2.44	0.47
7:I:101:MYR:C4	3:J:187:LEU:HD22	2.45	0.47
3:J:48:TRP:N	3:J:48:TRP:HD1	2.09	0.47
3:J:164:ASP:C	3:J:166:ALA:N	2.70	0.47
3:J:298:PHE:CE1	3:N:355:GLY:HA2	2.47	0.47
3:J:382:ILE:O	3:J:383:GLN:HB2	2.14	0.47
3:J:433:GLY:O	3:J:435:PRO:HD3	2.14	0.47
3:J:561:ASP:O	3:J:562:TYR:CB	2.62	0.47
3:J:561:ASP:O	3:J:563:PRO:CD	2.62	0.47
2:K:5:GLN:O	2:K:6:THR:CB	2.61	0.47
7:K:101:MYR:H102	3:L:200:CYS:HG	1.73	0.47
3:L:227:LEU:O	3:L:228:PRO:O	2.33	0.47
2:M:13:ILE:HG13	3:N:208:TYR:CG	2.50	0.47
3:N:154:ARG:HB3	3:N:154:ARG:CZ	2.44	0.47
3:N:164:ASP:C	3:N:166:ALA:N	2.71	0.47
3:N:352:ASP:HB2	3:N:421:HIS:CD2	2.49	0.47
3:N:521:VAL:HG21	3:N:577:VAL:HG22	1.95	0.47
3:P:46:LYS:NZ	3:T:237:GLU:OE1	2.38	0.47
3:P:431:ILE:CD1	3:P:437:ILE:HG23	2.39	0.47
3:P:633:ALA:O	3:P:636:ALA:HB3	2.14	0.47
3:R:204:PRO:O	3:R:208:TYR:HB2	2.14	0.47

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:R:560:PRO:HB2	3:R:562:TYR:CE1	2.50	0.47
2:S:5:GLN:O	2:S:6:THR:CB	2.61	0.47
3:T:365:ASP:OD2	3:T:452:SER:HB2	2.15	0.47
3:T:373:PHE:HD2	3:T:397:VAL:HG11	1.78	0.47
3:T:423:GLN:O	3:T:423:GLN:CG	2.62	0.47
3:T:521:VAL:HG21	3:T:577:VAL:HG22	1.95	0.47
3:T:564:LEU:HD23	3:T:564:LEU:C	2.40	0.47
4:U:250:TRP:CE3	6:X:436:LEU:HD13	2.50	0.47
4:U:253:GLY:O	4:U:254:ALA:C	2.58	0.47
4:V:268:ARG:HB3	4:V:327:ASP:OD1	2.13	0.47
5:W:57:THR:HG22	5:W:183:TYR:CD1	2.48	0.47
5:W:146:SER:N	5:W:147:PRO:CD	2.77	0.47
5:W:241:LEU:HD23	5:W:241:LEU:O	2.15	0.47
5:W:368:PRO:O	5:W:369:ILE:HG13	2.14	0.47
5:W:534:LEU:HD22	5:W:551:ILE:HG21	1.96	0.47
5:W:577:LEU:CD2	5:W:579:TYR:H	2.26	0.47
5:W:603:LEU:HD22	5:W:635:TYR:CE2	2.49	0.47
5:W:769:ARG:NH1	5:W:1000:TYR:OH	2.46	0.47
5:W:860:CYS:HB2	5:W:874:TYR:CZ	2.49	0.47
5:W:882:ALA:HA	5:W:920:ARG:CZ	2.44	0.47
5:W:955:VAL:HA	5:W:966:PRO:HA	1.97	0.47
5:W:1017:ILE:O	5:W:1017:ILE:HG22	2.13	0.47
5:W:1044:LEU:O	5:W:1045:ASN:HB2	2.13	0.47
5:W:1143:GLN:HA	5:W:1143:GLN:OE1	2.15	0.47
5:W:1180:LYS:HB2	5:W:1278:THR:O	2.14	0.47
5:W:1239:ALA:HB2	5:W:1289:LEU:HD21	1.96	0.47
6:X:327:GLY:O	6:X:328:ARG:O	2.32	0.47
6:X:679:TRP:CZ3	6:X:697:MET:HA	2.49	0.47
6:X:862:GLU:CG	6:X:957:ILE:HD12	2.44	0.47
6:X:1149:THR:HB	6:X:1162:ALA:N	2.29	0.47
6:Y:104:GLN:CD	6:Y:104:GLN:N	2.71	0.47
6:Y:260:GLU:OE1	6:Y:262:ARG:NH2	2.48	0.47
6:Y:294:ARG:HH21	6:Y:890:GLN:HB3	1.80	0.47
6:Y:324:ILE:HG23	6:Y:325:THR:HG23	1.97	0.47
6:Y:350:LEU:O	6:Y:352:THR:N	2.47	0.47
6:Y:459:LEU:HA	6:Y:459:LEU:HD12	1.78	0.47
6:Y:575:MET:HG2	6:Y:818:ILE:CG2	2.45	0.47
6:Y:662:LEU:N	6:Y:662:LEU:CD1	2.77	0.47
6:Y:684:LEU:HD11	6:Y:750:MET:HE3	1.97	0.47
6:Y:865:VAL:HB	6:Y:955:SER:CB	2.44	0.47
6:Y:1027:LEU:HD12	6:Y:1027:LEU:N	2.11	0.47

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:l:17:ALA:O	1:l:82:LEU:HD13	2.10	0.47
1:l:21:ARG:CZ	3:L:327:TYR:HE1	2.27	0.47
3:B:227:LEU:O	3:B:228:PRO:O	2.33	0.47
3:B:251:SER:OG	3:B:253:VAL:HG12	2.14	0.47
3:B:271:LEU:HD12	3:B:271:LEU:HA	1.79	0.47
3:D:396:VAL:HG11	3:D:399:ALA:HB2	1.96	0.47
3:D:420:LEU:HD12	3:D:422:LEU:HD23	1.95	0.47
3:D:521:VAL:HG23	3:D:522:GLY:N	2.30	0.47
3:F:355:GLY:CA	3:H:298:PHE:CZ	2.80	0.47
3:H:561:ASP:O	3:H:563:PRO:CD	2.62	0.47
3:H:564:LEU:CD2	3:H:568:ARG:NH2	2.75	0.47
3:L:564:LEU:HD23	3:L:564:LEU:C	2.40	0.47
3:N:166:ALA:CB	3:N:194:LEU:HD11	2.44	0.47
3:N:204:PRO:O	3:N:208:TYR:HB2	2.14	0.47
3:N:251:SER:OG	3:N:253:VAL:HG12	2.14	0.47
3:N:382:ILE:HB	3:N:438:TYR:CZ	2.50	0.47
3:P:420:LEU:HD12	3:P:422:LEU:HD23	1.96	0.47
3:P:428:ARG:NH1	3:P:428:ARG:CG	2.76	0.47
3:R:561:ASP:O	3:R:563:PRO:CD	2.62	0.47
3:T:561:ASP:O	3:T:563:PRO:CD	2.62	0.47
4:U:241:LEU:CB	6:X:1179:VAL:HG11	2.44	0.47
4:U:398:GLY:O	4:U:399:GLN:C	2.56	0.47
4:V:175:MET:HB2	6:Y:404:MET:HE2	1.97	0.47
5:W:209:THR:O	5:W:212:ALA:HB3	2.15	0.47
5:W:805:VAL:HG13	5:W:1006:SER:HB3	1.95	0.47
5:W:938:TYR:CG	5:W:1021:ARG:HD3	2.50	0.47
6:X:368:ILE:CG1	6:X:369:GLY:H	2.04	0.47
6:X:400:GLU:C	6:X:402:TYR:H	2.23	0.47
6:X:425:ASN:N	6:X:425:ASN:ND2	2.49	0.47
6:X:914:ASP:CG	6:X:914:ASP:O	2.58	0.47
6:X:1182:ASN:O	6:X:1184:LEU:N	2.48	0.47
6:Y:271:VAL:HG23	6:Y:288:PHE:HB3	1.96	0.47
6:Y:417:VAL:HG12	6:Y:421:ASN:ND2	2.30	0.47
6:Y:906:ASN:HD21	6:Y:921:ASP:HB3	1.78	0.47
6:Y:1087:TYR:HB3	6:Y:1121:ILE:HG12	1.96	0.47
1:j:17:ALA:O	1:j:82:LEU:CD1	2.62	0.47
1:p:4:HIS:CG	1:p:54:THR:HG22	2.46	0.47
3:B:493:GLN:O	3:B:586:VAL:HG23	2.14	0.47
3:B:561:ASP:O	3:B:562:TYR:CB	2.62	0.47
3:B:564:LEU:HD23	3:B:564:LEU:C	2.40	0.47
2:C:13:ILE:HG13	3:D:208:TYR:CG	2.50	0.47

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
7:C:101:MYR:C4	3:D:187:LEU:HD22	2.45	0.47
3:D:56:ALA:HB1	3:D:61:LEU:HG	1.96	0.47
3:D:564:LEU:HD23	3:D:564:LEU:C	2.40	0.47
7:E:101:MYR:H21	3:F:191:LYS:NZ	2.24	0.47
7:E:101:MYR:C4	3:F:187:LEU:HD22	2.45	0.47
3:F:164:ASP:C	3:F:166:ALA:N	2.71	0.47
2:G:28:SER:HA	3:H:216:PRO:CD	2.44	0.47
3:H:203:TYR:HB3	3:H:206:GLU:CB	2.39	0.47
3:J:96:PHE:CD1	3:J:101:THR:HG23	2.44	0.47
3:J:161:LYS:HB2	3:J:161:LYS:HE2	1.64	0.47
3:L:96:PHE:CD1	3:L:101:THR:HG23	2.44	0.47
3:N:382:ILE:O	3:N:383:GLN:HB2	2.14	0.47
3:N:423:GLN:O	3:N:423:GLN:CG	2.63	0.47
3:N:561:ASP:O	3:N:563:PRO:CD	2.62	0.47
3:N:633:ALA:O	3:N:636:ALA:HB3	2.14	0.47
3:P:322:TYR:C	3:P:323:SER:OG	2.58	0.47
3:P:365:ASP:HB3	3:P:452:SER:OG	2.15	0.47
3:R:251:SER:OG	3:R:253:VAL:HG12	2.14	0.47
3:T:305:LYS:HG3	3:T:477:LYS:O	2.14	0.47
3:T:382:ILE:O	3:T:383:GLN:HB2	2.14	0.47
4:U:329:ASN:CG	4:U:332:PRO:HD2	2.39	0.47
4:V:43:HIS:HB2	6:X:1027:LEU:HD11	1.97	0.47
5:W:47:LEU:HD23	5:W:60:GLN:CB	2.45	0.47
5:W:217:LEU:HA	5:W:220:LEU:HD22	1.97	0.47
5:W:464:TYR:CE2	5:W:473:ARG:NH1	2.83	0.47
5:W:691:ALA:HA	5:W:999:PRO:HB3	1.97	0.47
5:W:923:ALA:C	5:W:925:GLY:H	2.23	0.47
5:W:1008:LEU:C	5:W:1008:LEU:HD12	2.39	0.47
6:X:456:VAL:O	6:X:460:VAL:HG23	2.14	0.47
6:X:693:TYR:N	6:X:693:TYR:CD1	2.83	0.47
6:X:718:THR:OG1	6:X:726:ARG:NH1	2.48	0.47
6:X:920:LEU:HD13	6:X:920:LEU:O	2.15	0.47
6:X:1031:LEU:HD13	6:X:1031:LEU:C	2.40	0.47
6:X:1156:THR:HG21	6:X:1161:ASN:HA	1.97	0.47
6:Y:442:CYS:HB3	6:Y:444:TRP:CE2	2.50	0.47
6:Y:649:ASN:HD22	6:Y:649:ASN:HA	1.54	0.47
1:l:4:HIS:CE1	1:l:54:THR:HG21	2.47	0.47
1:d:43:CYS:H	1:d:44:GLY:HA2	1.80	0.47
1:h:3:LEU:HD12	1:h:6:ILE:H	1.80	0.47
1:r:22:LEU:CG	1:r:83:VAL:HG21	2.45	0.47
1:t:44:GLY:HA3	3:T:399:ALA:HB3	1.97	0.47

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:B:92:TRP:CD2	3:B:156:VAL:HG23	2.51	0.47
3:B:433:GLY:O	3:B:435:PRO:HD3	2.15	0.47
3:D:238:VAL:CG2	3:F:78:MET:HE1	2.07	0.47
2:E:13:ILE:HG13	3:F:208:TYR:CG	2.50	0.47
3:F:431:ILE:HD11	3:F:437:ILE:CG2	2.45	0.47
3:F:560:PRO:HB2	3:F:562:TYR:CE1	2.50	0.47
3:H:58:ILE:CG1	3:H:127:LEU:HD23	2.45	0.47
3:H:275:GLU:HG2	3:H:275:GLU:O	2.15	0.47
3:H:358:LEU:HB2	3:H:417:PHE:HD1	1.77	0.47
3:H:365:ASP:OD2	3:H:452:SER:HB2	2.15	0.47
3:J:564:LEU:HD23	3:J:564:LEU:C	2.40	0.47
3:J:578:SER:OG	3:N:468:PRO:O	2.21	0.47
3:L:352:ASP:HB2	3:L:421:HIS:CD2	2.49	0.47
3:L:588:ILE:HD13	3:L:588:ILE:HA	1.80	0.47
3:N:48:TRP:CE2	3:N:63:ILE:HD13	2.49	0.47
3:N:58:ILE:CG1	3:N:127:LEU:HD23	2.45	0.47
3:N:560:PRO:HB2	3:N:562:TYR:CE1	2.50	0.47
3:N:564:LEU:CD2	3:N:568:ARG:NH2	2.75	0.47
2:O:13:ILE:HG13	3:P:208:TYR:CG	2.50	0.47
7:O:101:MYR:C4	3:P:187:LEU:HD22	2.45	0.47
2:Q:5:GLN:O	2:Q:6:THR:CB	2.61	0.47
3:R:561:ASP:O	3:R:562:TYR:CB	2.62	0.47
3:R:564:LEU:HD23	3:R:564:LEU:C	2.40	0.47
3:T:58:ILE:O	3:T:58:ILE:CG2	2.62	0.47
3:T:169:MET:HG2	3:T:169:MET:O	2.15	0.47
5:W:934:ASN:HD22	5:W:934:ASN:N	2.04	0.47
5:W:986:LEU:HD23	5:W:1028:ILE:HG23	1.97	0.47
6:X:200:LEU:CD1	6:X:343:ALA:HA	2.45	0.47
6:X:386:GLU:HG3	6:X:1195:TYR:CE2	2.50	0.47
6:X:853:ARG:CZ	6:X:853:ARG:HB3	2.45	0.47
6:X:958:HIS:HD2	6:X:958:HIS:O	1.97	0.47
1:b:45:ARG:HG2	3:B:399:ALA:C	2.38	0.46
1:b:59:HIS:HB2	3:B:397:VAL:C	2.22	0.46
1:f:43:CYS:H	1:f:44:GLY:HA2	1.80	0.46
1:h:6:ILE:HG21	3:D:494:THR:HG22	1.98	0.46
1:h:44:GLY:CA	3:H:399:ALA:HB3	2.45	0.46
1:j:44:GLY:O	3:J:399:ALA:HB3	2.15	0.46
1:j:76:ASN:C	1:j:78:ALA:H	2.23	0.46
1:t:44:GLY:O	3:T:399:ALA:C	2.58	0.46
3:B:169:MET:HG2	3:B:169:MET:O	2.15	0.46
3:B:536:THR:HG22	3:B:573:ALA:HB3	1.97	0.46

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:B:564:LEU:CD2	3:B:568:ARG:NH2	2.75	0.46
2:C:26:MET:O	2:C:26:MET:CG	2.60	0.46
3:D:173:ASP:HB2	3:H:606:THR:HG22	1.94	0.46
3:D:427:GLU:CD	3:F:571:ARG:HH22	2.20	0.46
3:F:154:ARG:HB3	3:F:154:ARG:CZ	2.43	0.46
3:F:251:SER:OG	3:F:253:VAL:HG12	2.15	0.46
3:F:355:GLY:HA2	3:H:298:PHE:CE1	2.47	0.46
3:F:525:ILE:HD11	3:F:604:VAL:HA	1.98	0.46
3:H:48:TRP:N	3:H:48:TRP:HD1	2.09	0.46
2:I:28:SER:HB2	3:J:215:TYR:HA	1.96	0.46
3:L:92:TRP:CD2	3:L:156:VAL:HG23	2.51	0.46
3:L:365:ASP:OD2	3:L:452:SER:HB2	2.15	0.46
2:M:28:SER:HA	3:N:216:PRO:CD	2.44	0.46
2:O:28:SER:HB2	3:P:215:TYR:HA	1.96	0.46
3:P:128:ASP:CB	3:T:107:VAL:HG11	2.45	0.46
3:R:431:ILE:O	3:R:431:ILE:HG23	2.10	0.46
3:T:58:ILE:CG1	3:T:127:LEU:HD23	2.45	0.46
4:U:12:ASN:O	4:U:14:TYR:N	2.48	0.46
4:U:243:PRO:C	4:U:245:GLN:H	2.23	0.46
4:V:119:LEU:HD13	4:V:313:ALA:HB2	1.96	0.46
5:W:436:GLY:CA	5:W:1009:ASN:HA	2.34	0.46
5:W:610:THR:HG21	5:W:616:SER:HB3	1.96	0.46
5:W:850:LEU:HD23	5:W:851:GLN:N	2.30	0.46
5:W:1075:PRO:HA	5:W:1087:LEU:HD13	1.97	0.46
5:W:1180:LYS:HD3	5:W:1278:THR:O	2.16	0.46
5:W:1224:VAL:HG12	5:W:1224:VAL:O	2.15	0.46
6:X:364:ILE:CG1	6:X:367:ARG:HH21	2.25	0.46
6:X:491:THR:O	6:X:495:ASN:HB2	2.15	0.46
6:X:814:ASP:OD2	6:X:816:ALA:HB3	2.16	0.46
6:X:841:PHE:O	6:X:842:ALA:C	2.58	0.46
6:X:1064:TRP:HB3	6:X:1065:PRO:HD3	1.97	0.46
6:Y:383:ALA:HB1	6:Y:439:THR:CG2	2.45	0.46
6:Y:658:TYR:CD2	6:Y:675:MET:HG3	2.50	0.46
6:Y:990:ASN:O	6:Y:990:ASN:CG	2.58	0.46
1:l:21:ARG:O	3:L:323:SER:OG	2.33	0.46
1:l:60:ALA:CB	3:L:322:TYR:HH	2.27	0.46
1:d:15:ARG:HH11	3:D:406:ASN:ND2	2.07	0.46
1:d:44:GLY:C	3:D:399:ALA:HB3	2.40	0.46
1:n:58:PRO:HB3	3:N:423:GLN:NE2	2.24	0.46
1:p:61:ASN:HB2	3:P:396:VAL:O	2.15	0.46
1:r:60:ALA:O	1:r:61:ASN:CB	2.62	0.46

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:E:28:SER:HA	3:F:216:PRO:CD	2.44	0.46
3:F:365:ASP:HB3	3:F:452:SER:OG	2.15	0.46
2:G:26:MET:O	2:G:26:MET:CG	2.60	0.46
3:H:78:MET:O	3:H:78:MET:HG2	2.13	0.46
3:H:137:ILE:HD13	3:H:142:ILE:HG12	1.97	0.46
3:H:521:VAL:HG23	3:H:522:GLY:N	2.30	0.46
3:H:521:VAL:HG21	3:H:577:VAL:HG22	1.95	0.46
2:I:13:ILE:HG13	3:J:208:TYR:CG	2.50	0.46
3:J:128:ASP:CB	3:N:107:VAL:HG11	2.45	0.46
3:J:137:ILE:HD13	3:J:142:ILE:HG12	1.98	0.46
3:J:305:LYS:HG3	3:J:477:LYS:O	2.14	0.46
3:J:369:SER:HB2	3:J:445:ILE:HA	1.97	0.46
3:J:382:ILE:HB	3:J:438:TYR:CZ	2.50	0.46
2:K:28:SER:HA	3:L:216:PRO:CD	2.44	0.46
3:L:423:GLN:O	3:L:423:GLN:CG	2.62	0.46
3:N:227:LEU:O	3:N:228:PRO:O	2.33	0.46
3:N:433:GLY:O	3:N:435:PRO:HD3	2.14	0.46
3:P:251:SER:OG	3:P:253:VAL:HG12	2.14	0.46
3:P:525:ILE:HD11	3:P:604:VAL:HA	1.97	0.46
2:Q:24:SER:HB2	3:R:175:SER:C	2.41	0.46
3:R:227:LEU:O	3:R:228:PRO:O	2.33	0.46
3:R:354:SER:HB3	3:R:419:MET:HG2	1.97	0.46
3:R:382:ILE:O	3:R:383:GLN:HB2	2.14	0.46
3:R:382:ILE:HB	3:R:438:TYR:CZ	2.50	0.46
2:S:13:ILE:HG13	3:T:208:TYR:CG	2.50	0.46
3:T:251:SER:OG	3:T:253:VAL:HG12	2.14	0.46
3:T:536:THR:HG22	3:T:573:ALA:HB3	1.98	0.46
3:T:559:PRO:O	3:T:560:PRO:C	2.58	0.46
4:U:223:LEU:O	4:U:233:TYR:HB2	2.16	0.46
4:V:151:SER:OG	4:V:268:ARG:HD2	2.15	0.46
4:V:386:ILE:O	4:V:387:GLN:HB3	2.14	0.46
5:W:225:THR:HG22	5:W:226:VAL:N	2.30	0.46
5:W:233:PRO:HG3	5:W:237:PRO:HA	1.97	0.46
5:W:955:VAL:HG22	5:W:956:TYR:N	2.30	0.46
6:X:205:MET:HB3	6:X:249:TRP:CE2	2.50	0.46
6:X:417:VAL:HG21	6:X:432:ILE:HG22	1.97	0.46
6:X:696:ALA:O	6:X:697:MET:C	2.59	0.46
6:X:1064:TRP:CE3	6:X:1064:TRP:HA	2.51	0.46
6:X:1106:LYS:CB	6:X:1113:PRO:HG2	2.39	0.46
6:Y:602:GLY:HA3	6:Y:623:PRO:CD	2.45	0.46
1:d:73:ARG:C	1:d:73:ARG:CD	2.87	0.46

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:r:74:GLN:C	1:r:74:GLN:NE2	2.73	0.46
3:D:128:ASP:CB	3:H:107:VAL:HG11	2.45	0.46
3:D:365:ASP:HB3	3:D:452:SER:OG	2.15	0.46
3:D:369:SER:HB2	3:D:445:ILE:HA	1.97	0.46
3:H:358:LEU:CB	3:H:417:PHE:HE1	2.24	0.46
3:H:365:ASP:HB3	3:H:452:SER:OG	2.16	0.46
3:H:369:SER:HB2	3:H:445:ILE:HA	1.97	0.46
7:I:101:MYR:C2	3:J:191:LYS:HZ1	2.28	0.46
3:J:365:ASP:HB3	3:J:452:SER:OG	2.15	0.46
3:J:536:THR:HG22	3:J:573:ALA:HB3	1.97	0.46
3:J:571:ARG:HH22	3:N:427:GLU:CD	2.19	0.46
3:L:107:VAL:HG11	3:N:128:ASP:CB	2.45	0.46
7:O:101:MYR:H21	3:P:191:LYS:NZ	2.30	0.46
3:P:355:GLY:HA2	3:R:298:PHE:CE1	2.47	0.46
3:P:606:THR:HG22	3:R:173:ASP:HB2	1.94	0.46
3:R:525:ILE:HD11	3:R:604:VAL:HA	1.97	0.46
3:T:396:VAL:HG11	3:T:399:ALA:HB2	1.96	0.46
3:T:521:VAL:HG23	3:T:522:GLY:N	2.30	0.46
4:U:192:PHE:C	4:U:194:GLY:H	2.23	0.46
4:U:221:GLY:N	4:U:337:ARG:HG2	2.28	0.46
4:V:77:ASN:N	4:V:78:PRO:CD	2.78	0.46
4:V:302:GLU:HG2	4:V:302:GLU:O	2.14	0.46
5:W:381:PHE:CD1	5:W:382:PRO:HD2	2.51	0.46
5:W:969:ASP:OD2	5:W:972:PRO:HG2	2.15	0.46
5:W:1145:ASP:O	5:W:1196:LEU:HD23	2.16	0.46
5:W:1239:ALA:CB	5:W:1250:LEU:HD11	2.45	0.46
5:W:1240:SER:HB2	5:W:1279:SER:HA	1.97	0.46
6:X:472:LEU:HD11	6:X:764:THR:HG21	1.96	0.46
6:X:481:ARG:O	6:X:484:THR:HG22	2.15	0.46
6:X:497:ILE:HD13	6:X:514:ILE:HD11	1.96	0.46
6:Y:430:SER:HB3	6:Y:1208:MET:HG3	1.97	0.46
6:Y:453:LEU:O	6:Y:457:MET:HB2	2.16	0.46
6:Y:835:HIS:CE1	6:Y:1212:SER:HA	2.51	0.46
1:l:59:HIS:HB2	3:L:398:SER:HB2	0.62	0.46
1:h:35:PHE:HZ	1:h:74:GLN:HB3	1.80	0.46
1:j:45:ARG:CG	3:J:400:ALA:C	2.68	0.46
1:p:61:ASN:HB2	3:P:395:THR:CG2	2.45	0.46
1:t:44:GLY:O	3:T:399:ALA:HB3	2.16	0.46
3:B:58:ILE:CG1	3:B:127:LEU:HD23	2.45	0.46
3:B:305:LYS:HG3	3:B:477:LYS:O	2.14	0.46
3:D:340:ASP:O	3:D:344:MET:HG3	2.16	0.46

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:F:561:ASP:O	3:F:562:TYR:CB	2.62	0.46
3:H:525:ILE:HD11	3:H:604:VAL:HA	1.98	0.46
3:H:600:LEU:HG	3:H:601:THR:N	2.31	0.46
2:K:36:LEU:HD22	2:K:41:LEU:HD11	1.98	0.46
3:L:431:ILE:O	3:L:431:ILE:HG23	2.10	0.46
3:L:536:THR:HG22	3:L:573:ALA:HB3	1.98	0.46
3:L:633:ALA:O	3:L:636:ALA:HB3	2.14	0.46
2:M:24:SER:HB2	3:N:175:SER:C	2.41	0.46
3:N:92:TRP:CD2	3:N:156:VAL:HG23	2.51	0.46
3:N:169:MET:HG2	3:N:169:MET:O	2.15	0.46
3:N:369:SER:HB2	3:N:444:LEU:O	2.16	0.46
3:N:431:ILE:O	3:N:431:ILE:HG23	2.10	0.46
3:N:564:LEU:HD23	3:N:564:LEU:C	2.40	0.46
3:P:369:SER:HB2	3:P:445:ILE:HA	1.97	0.46
3:P:560:PRO:HB2	3:P:562:TYR:CE1	2.50	0.46
3:P:561:ASP:O	3:P:562:TYR:CB	2.62	0.46
3:P:564:LEU:HD23	3:P:564:LEU:C	2.40	0.46
2:Q:26:MET:CG	3:R:216:PRO:HG3	2.29	0.46
3:R:69:GLY:HA3	3:R:139:THR:HB	1.97	0.46
3:R:274:ILE:HD12	3:R:514:LEU:HD23	1.94	0.46
3:R:369:SER:HB2	3:R:444:LEU:O	2.16	0.46
3:R:396:VAL:HG11	3:R:399:ALA:HB2	1.96	0.46
2:S:24:SER:HB2	3:T:175:SER:C	2.41	0.46
3:T:92:TRP:CD2	3:T:156:VAL:HG23	2.51	0.46
3:T:369:SER:HB2	3:T:444:LEU:O	2.16	0.46
3:T:561:ASP:O	3:T:562:TYR:CB	2.62	0.46
4:U:341:TRP:O	4:U:346:LEU:HB2	2.14	0.46
5:W:247:VAL:CG2	5:W:257:GLN:HE21	2.28	0.46
5:W:736:VAL:CG2	5:W:780:LEU:HD21	2.41	0.46
5:W:793:THR:HB	5:W:794:PRO:HD2	1.98	0.46
6:X:278:LEU:HD23	6:X:292:ALA:HB2	1.96	0.46
6:X:377:ILE:HG13	6:X:438:PRO:HG3	1.97	0.46
6:X:593:LEU:CD2	6:X:597:GLU:HB2	2.46	0.46
6:X:983:TYR:HD2	6:X:1137:PHE:CZ	2.34	0.46
6:Y:63:VAL:HG12	6:Y:64:GLY:N	2.30	0.46
6:Y:763:VAL:O	6:Y:763:VAL:HG13	2.15	0.46
6:Y:976:ASN:O	6:Y:1083:TYR:CD2	2.69	0.46
1:l:44:GLY:CA	3:L:399:ALA:HB3	2.45	0.46
1:d:15:ARG:CZ	3:D:406:ASN:ND2	2.78	0.46
1:d:24:LEU:CD2	1:d:80:ARG:HD2	2.45	0.46
1:r:43:CYS:H	1:r:44:GLY:HA2	1.80	0.46

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:r:59:HIS:CB	3:R:397:VAL:O	2.57	0.46
1:t:43:CYS:H	1:t:44:GLY:HA2	1.80	0.46
3:B:600:LEU:HG	3:B:601:THR:N	2.30	0.46
3:D:238:VAL:CA	3:F:142:ILE:CD1	2.81	0.46
3:D:609:ARG:HD2	2:E:29:THR:HG22	1.91	0.46
3:F:305:LYS:HG3	3:F:477:LYS:O	2.14	0.46
3:F:345:ILE:HG23	3:F:475:THR:HG1	1.81	0.46
3:F:354:SER:HB3	3:F:419:MET:HG2	1.97	0.46
3:F:369:SER:HB2	3:F:444:LEU:O	2.16	0.46
3:H:69:GLY:HA3	3:H:139:THR:HB	1.97	0.46
3:H:431:ILE:HD11	3:H:437:ILE:CG2	2.45	0.46
2:I:24:SER:HB2	3:J:175:SER:C	2.41	0.46
3:J:67:ARG:HH11	3:J:67:ARG:CG	2.29	0.46
3:L:345:ILE:HG23	3:L:475:THR:OG1	2.16	0.46
3:P:369:SER:HB2	3:P:444:LEU:O	2.16	0.46
7:Q:101:MYR:H22	3:R:191:LYS:HZ1	1.72	0.46
3:R:107:VAL:HG11	3:T:128:ASP:CB	2.45	0.46
3:R:154:ARG:HB3	3:R:154:ARG:CZ	2.44	0.46
3:R:164:ASP:C	3:R:166:ALA:N	2.71	0.46
3:R:427:GLU:CD	3:T:571:ARG:HH22	2.19	0.46
3:T:275:GLU:HG2	3:T:275:GLU:O	2.15	0.46
4:V:305:LEU:HB3	4:V:306:PRO:HD2	1.98	0.46
5:W:5:PHE:CZ	5:W:306:ARG:HG2	2.50	0.46
5:W:261:LEU:HA	5:W:264:GLU:HG2	1.97	0.46
5:W:1166:CYS:HB3	5:W:1175:ARG:HA	1.98	0.46
6:X:823:LEU:HG	6:X:823:LEU:O	2.16	0.46
6:X:864:PHE:CD2	6:X:952:PRO:HG3	2.50	0.46
6:X:895:SER:HA	6:X:930:ARG:HD2	1.98	0.46
6:X:1094:LEU:CD2	6:X:1132:ARG:HB3	2.45	0.46
6:Y:262:ARG:HG3	6:Y:262:ARG:NH1	2.31	0.46
6:Y:428:LEU:HG	6:Y:1208:MET:HG2	1.97	0.46
6:Y:664:GLY:O	6:Y:665:SER:C	2.58	0.46
6:Y:668:LEU:HD12	6:Y:767:PHE:HE1	1.79	0.46
6:Y:868:ARG:HD3	6:Y:932:ALA:O	2.16	0.46
1:h:74:GLN:NE2	1:h:74:GLN:C	2.73	0.46
1:n:21:ARG:HE	3:N:370:VAL:CG1	2.27	0.46
1:t:60:ALA:HB3	3:T:322:TYR:CE1	2.48	0.46
3:B:365:ASP:HB3	3:B:452:SER:OG	2.15	0.46
3:B:369:SER:HB2	3:B:444:LEU:O	2.16	0.46
3:D:154:ARG:HB3	3:D:154:ARG:CZ	2.44	0.46
3:D:275:GLU:HG2	3:D:275:GLU:O	2.15	0.46

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:D:382:ILE:HB	3:D:438:TYR:CZ	2.50	0.46
3:D:443:PHE:HE2	3:D:459:LEU:HD21	1.75	0.46
3:D:559:PRO:O	3:D:560:PRO:C	2.58	0.46
3:F:227:LEU:O	3:F:228:PRO:O	2.33	0.46
3:F:345:ILE:HG23	3:F:475:THR:OG1	2.16	0.46
3:F:469:VAL:HG21	3:H:575:MET:HE3	0.69	0.46
2:G:13:ILE:HG13	3:H:208:TYR:CG	2.50	0.46
3:J:69:GLY:HA3	3:J:139:THR:HB	1.97	0.46
3:J:107:VAL:HG11	3:L:128:ASP:CB	2.45	0.46
3:J:369:SER:HB2	3:J:444:LEU:O	2.16	0.46
3:L:58:ILE:CG1	3:L:127:LEU:HD23	2.46	0.46
3:L:274:ILE:HD12	3:L:514:LEU:HD23	1.94	0.46
3:L:521:VAL:HG23	3:L:522:GLY:N	2.30	0.46
3:N:147:ARG:HH12	3:R:100:LYS:NZ	2.14	0.46
3:N:275:GLU:HG2	3:N:275:GLU:O	2.15	0.46
3:N:411:VAL:O	3:N:412:ARG:CB	2.57	0.46
3:N:536:THR:HG22	3:N:573:ALA:HB3	1.97	0.46
2:O:24:SER:HB2	3:P:175:SER:C	2.41	0.46
3:P:227:LEU:O	3:P:228:PRO:O	2.33	0.46
3:P:345:ILE:HG23	3:P:475:THR:OG1	2.16	0.46
3:P:367:SER:O	3:P:368:THR:CB	2.64	0.46
2:Q:13:ILE:HG13	3:R:208:TYR:CG	2.50	0.46
3:R:540:VAL:CG2	3:R:566:TRP:HA	2.46	0.46
4:U:140:GLN:HE21	4:U:144:ILE:CG1	2.28	0.46
4:U:180:VAL:CG1	4:U:184:HIS:HB2	2.45	0.46
4:V:195:GLN:NE2	6:Y:377:ILE:CB	2.73	0.46
4:V:348:THR:HG22	4:V:349:ALA:N	2.30	0.46
5:W:259:ALA:HB1	5:W:303:LEU:HD12	1.95	0.46
5:W:853:THR:HA	5:W:873:ALA:O	2.16	0.46
5:W:857:SER:HB3	5:W:878:ASP:OD1	2.15	0.46
5:W:1290:PRO:HD2	5:W:1294:TYR:OH	2.16	0.46
6:X:377:ILE:CD1	6:X:436:LEU:HD11	2.45	0.46
6:X:413:PRO:C	6:X:415:GLN:N	2.74	0.46
6:X:658:TYR:CD2	6:X:675:MET:HG3	2.50	0.46
6:X:732:VAL:O	6:X:732:VAL:HG12	2.16	0.46
6:Y:941:TYR:C	6:Y:943:GLY:N	2.74	0.46
6:Y:1153:SER:C	6:Y:1155:ASP:H	2.24	0.46
1:l:60:ALA:HB1	3:L:322:TYR:OH	2.11	0.46
1:b:44:GLY:O	3:B:399:ALA:C	2.58	0.46
1:p:15:ARG:HB3	3:P:402:ALA:HB2	1.97	0.46
3:D:178:SER:O	3:D:179:ALA:CB	2.64	0.46

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:D:564:LEU:CD2	3:D:568:ARG:NH2	2.75	0.46
3:F:96:PHE:CD1	3:F:101:THR:HG23	2.44	0.46
3:F:107:VAL:HG11	3:H:128:ASP:CB	2.45	0.46
3:F:275:GLU:HG2	3:F:275:GLU:O	2.15	0.46
3:F:382:ILE:HB	3:F:438:TYR:CZ	2.50	0.46
3:H:540:VAL:CG2	3:H:566:TRP:HA	2.46	0.46
3:J:178:SER:O	3:J:179:ALA:CB	2.64	0.46
2:K:24:SER:HB2	3:L:175:SER:C	2.41	0.46
3:L:365:ASP:HB3	3:L:452:SER:OG	2.15	0.46
3:L:561:ASP:O	3:L:562:TYR:CB	2.62	0.46
3:N:346:ASP:CB	3:N:427:GLU:HA	2.46	0.46
3:N:559:PRO:O	3:N:560:PRO:C	2.58	0.46
3:P:57:THR:HB	3:P:59:ASP:OD1	2.15	0.46
3:P:298:PHE:CE1	3:T:355:GLY:HA2	2.47	0.46
3:P:340:ASP:O	3:P:344:MET:HG3	2.16	0.46
3:P:536:THR:HG22	3:P:573:ALA:HB3	1.97	0.46
3:R:169:MET:HG2	3:R:169:MET:O	2.15	0.46
3:R:365:ASP:HB3	3:R:452:SER:OG	2.15	0.46
3:R:559:PRO:O	3:R:560:PRO:C	2.58	0.46
3:R:600:LEU:HG	3:R:601:THR:N	2.30	0.46
3:T:561:ASP:O	3:T:563:PRO:HD3	2.16	0.46
4:U:6:PHE:CD2	4:U:122:SER:HB3	2.51	0.46
4:U:136:TRP:O	4:U:139:ASN:HB3	2.16	0.46
4:U:268:ARG:O	4:U:327:ASP:HA	2.15	0.46
4:U:300:ILE:O	4:U:300:ILE:HG13	2.15	0.46
5:W:860:CYS:SG	5:W:864:MET:SD	3.14	0.46
5:W:973:LEU:HD13	5:W:973:LEU:O	2.16	0.46
5:W:1144:LEU:HA	5:W:1151:ASN:OD1	2.16	0.46
5:W:1155:ILE:HD12	5:W:1163:LEU:HD12	1.97	0.46
5:W:1241:LEU:HG	5:W:1242:PRO:HD2	1.98	0.46
6:X:356:ASN:C	6:X:358:GLN:N	2.73	0.46
6:X:555:GLN:HA	6:X:555:GLN:OE1	2.15	0.46
6:X:559:HIS:CG	6:X:561:PRO:HD2	2.51	0.46
6:X:656:PHE:CE2	6:X:752:PRO:HB3	2.51	0.46
6:X:889:ARG:CG	6:X:890:GLN:H	2.24	0.46
6:X:894:THR:O	6:X:897:ALA:HB3	2.15	0.46
6:Y:48:GLN:HE21	6:Y:53:ILE:HB	1.79	0.46
6:Y:202:THR:HB	6:Y:392:ASP:OD1	2.15	0.46
6:Y:663:GLN:HA	6:Y:663:GLN:NE2	2.30	0.46
6:Y:691:PRO:O	6:Y:692:THR:HB	2.16	0.46
6:Y:1122:PRO:CA	6:Y:1159:GLY:HA3	2.38	0.46

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:l:43:CYS:H	1:l:44:GLY:HA2	1.80	0.46
1:j:43:CYS:H	1:j:44:GLY:HA2	1.80	0.46
1:n:24:LEU:CD2	1:n:80:ARG:CD	2.90	0.46
1:n:74:GLN:O	1:n:75:PRO:C	2.56	0.46
1:p:61:ASN:CG	3:P:395:THR:HG23	2.25	0.46
1:t:59:HIS:CD2	3:T:398:SER:HB2	2.51	0.46
2:A:28:SER:HB2	3:B:215:TYR:HA	1.96	0.46
2:A:36:LEU:HD22	2:A:41:LEU:HD11	1.98	0.46
3:B:67:ARG:HH11	3:B:67:ARG:CG	2.29	0.46
3:D:260:SER:HB3	2:E:33:ALA:HB1	1.96	0.46
3:D:341:MET:CE	3:D:341:MET:CA	2.88	0.46
3:D:369:SER:HB2	3:D:444:LEU:O	2.16	0.46
3:D:493:GLN:O	3:D:586:VAL:HG23	2.14	0.46
2:E:6:THR:HG23	2:E:7:SER:N	2.31	0.46
2:E:24:SER:HB2	3:F:175:SER:C	2.41	0.46
3:F:178:SER:O	3:F:179:ALA:CB	2.64	0.46
3:F:344:MET:HE2	3:F:502:LEU:CD1	2.37	0.46
3:F:536:THR:HG22	3:F:573:ALA:HB3	1.98	0.46
3:F:540:VAL:CG2	3:F:566:TRP:HA	2.46	0.46
3:H:345:ILE:HG23	3:H:475:THR:HG1	1.81	0.46
3:H:369:SER:HB2	3:H:444:LEU:O	2.16	0.46
3:L:178:SER:O	3:L:179:ALA:CB	2.64	0.46
3:L:275:GLU:HG2	3:L:275:GLU:O	2.15	0.46
3:L:561:ASP:O	3:L:563:PRO:HD3	2.16	0.46
3:L:582:LEU:HD23	3:L:582:LEU:HA	1.69	0.46
3:P:358:LEU:HB2	3:P:417:PHE:HD1	1.77	0.46
3:P:521:VAL:HG23	3:P:522:GLY:N	2.30	0.46
3:P:600:LEU:HG	3:P:601:THR:N	2.31	0.46
2:Q:6:THR:HG23	2:Q:7:SER:N	2.31	0.46
3:R:275:GLU:O	3:R:275:GLU:HG2	2.15	0.46
3:R:521:VAL:HG23	3:R:522:GLY:N	2.30	0.46
3:T:178:SER:O	3:T:179:ALA:CB	2.64	0.46
3:T:345:ILE:HG23	3:T:475:THR:OG1	2.16	0.46
3:T:431:ILE:HD11	3:T:437:ILE:CG2	2.45	0.46
3:T:433:GLY:O	3:T:435:PRO:HD3	2.14	0.46
4:U:14:TYR:O	4:U:15:GLY:C	2.59	0.46
4:U:62:ALA:HB2	4:U:281:VAL:HG21	1.97	0.46
4:U:178:THR:HG22	6:X:376:LEU:CG	2.42	0.46
4:U:180:VAL:HA	4:U:181:PRO:HD3	1.76	0.46
4:U:248:LEU:C	4:U:249:LYS:HG2	2.41	0.46
4:V:96:VAL:HA	4:V:135:MET:HE2	1.97	0.46

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
5:W:271:ARG:HB2	5:W:324:ILE:HB	1.97	0.46
5:W:622:PHE:O	5:W:624:THR:N	2.46	0.46
5:W:687:ALA:HB1	5:W:996:ASP:CA	2.35	0.46
5:W:739:VAL:HG12	5:W:742:SER:HB2	1.97	0.46
5:W:879:TYR:OH	5:W:898:THR:HG22	2.16	0.46
6:X:839:LYS:HZ2	6:X:839:LYS:HB2	1.79	0.46
6:X:1098:PRO:C	6:X:1100:LEU:H	2.21	0.46
6:Y:188:ALA:CB	6:Y:763:VAL:HG11	2.45	0.46
6:Y:247:THR:HG22	6:Y:308:ASN:ND2	2.31	0.46
6:Y:356:ASN:CB	6:Y:1151:ALA:HB1	2.45	0.46
6:Y:634:GLN:H	6:Y:634:GLN:NE2	2.14	0.46
6:Y:996:MET:CE	6:Y:996:MET:CA	2.84	0.46
1:l:22:LEU:N	1:l:22:LEU:HD22	2.30	0.46
1:p:4:HIS:HD2	1:p:54:THR:CG2	2.22	0.46
3:B:178:SER:O	3:B:179:ALA:CB	2.64	0.46
3:B:345:ILE:HG23	3:B:475:THR:OG1	2.16	0.46
3:D:92:TRP:CD2	3:D:156:VAL:HG23	2.51	0.46
3:D:107:VAL:HG11	3:F:128:ASP:CB	2.45	0.46
3:D:354:SER:HB3	3:D:419:MET:HG2	1.97	0.46
3:D:367:SER:O	3:D:368:THR:CB	2.64	0.46
3:D:533:ASP:OD1	3:D:535:ALA:HB3	2.16	0.46
3:H:92:TRP:CD2	3:H:156:VAL:HG23	2.51	0.46
3:H:178:SER:O	3:H:179:ALA:CB	2.64	0.46
3:J:260:SER:HB3	2:K:33:ALA:HB1	1.96	0.46
3:J:525:ILE:HD11	3:J:604:VAL:HA	1.98	0.46
3:L:346:ASP:CB	3:L:427:GLU:HA	2.46	0.46
3:L:540:VAL:CG2	3:L:566:TRP:HA	2.46	0.46
3:L:600:LEU:HG	3:L:601:THR:N	2.30	0.46
3:N:345:ILE:HG23	3:N:475:THR:OG1	2.16	0.46
3:N:365:ASP:HB3	3:N:452:SER:OG	2.16	0.46
3:P:67:ARG:HH11	3:P:67:ARG:CG	2.29	0.46
3:P:92:TRP:CD2	3:P:156:VAL:HG23	2.50	0.46
3:P:154:ARG:HB3	3:P:154:ARG:CZ	2.44	0.46
3:P:346:ASP:CB	3:P:427:GLU:HA	2.46	0.46
3:P:411:VAL:O	3:P:411:VAL:CG1	2.61	0.46
3:P:564:LEU:CD2	3:P:568:ARG:NH2	2.75	0.46
3:R:345:ILE:HG23	3:R:475:THR:OG1	2.16	0.46
3:R:369:SER:HB2	3:R:445:ILE:HA	1.97	0.46
3:T:227:LEU:O	3:T:228:PRO:O	2.33	0.46
3:T:365:ASP:HB3	3:T:452:SER:OG	2.15	0.46
4:U:39:SER:HB2	6:X:954:PRO:CB	2.45	0.46

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
4:U:51:LEU:C	4:U:53:VAL:H	2.24	0.46
4:U:178:THR:HB	6:X:376:LEU:HD23	1.98	0.46
4:V:64:ARG:NH2	6:X:1027:LEU:CD2	2.79	0.46
4:V:224:ASN:HD22	4:V:224:ASN:N	2.12	0.46
5:W:1061:LEU:O	5:W:1072:GLY:HA2	2.16	0.46
5:W:1289:LEU:N	5:W:1289:LEU:HD22	2.31	0.46
6:X:243:ILE:CG1	6:X:1147:PHE:HB2	2.41	0.46
6:X:271:VAL:CG1	6:X:288:PHE:HB3	2.45	0.46
6:X:476:LEU:O	6:X:480:LEU:HD23	2.16	0.46
6:X:590:ALA:HB1	6:X:639:LEU:HD21	1.97	0.46
6:Y:197:ASP:CB	6:Y:253:GLY:HA3	2.46	0.46
6:Y:530:LYS:HD3	6:Y:811:LEU:HD21	1.98	0.46
6:Y:712:VAL:HG21	6:Y:729:MET:HB3	1.98	0.46
6:Y:978:ALA:HB3	6:Y:1081:TYR:CE1	2.51	0.46
1:b:61:ASN:C	1:b:63:LYS:N	2.74	0.46
1:n:43:CYS:H	1:n:44:GLY:HA2	1.80	0.46
1:t:15:ARG:CZ	3:T:402:ALA:HB3	2.34	0.46
2:A:24:SER:HB2	3:B:175:SER:C	2.41	0.46
3:B:69:GLY:HA3	3:B:139:THR:HB	1.97	0.46
3:B:275:GLU:HG2	3:B:275:GLU:O	2.15	0.46
3:B:340:ASP:O	3:B:344:MET:HG3	2.16	0.46
3:B:365:ASP:OD2	3:B:452:SER:HB2	2.15	0.46
3:B:367:SER:O	3:B:368:THR:CB	2.64	0.46
3:B:540:VAL:CG2	3:B:566:TRP:HA	2.46	0.46
3:D:215:TYR:C	3:D:217:ALA:N	2.70	0.46
3:D:269:LYS:HG2	3:D:516:LYS:HZ3	1.81	0.46
3:D:521:VAL:HG21	3:D:577:VAL:HG22	1.95	0.46
3:D:561:ASP:O	3:D:563:PRO:HD3	2.16	0.46
3:D:571:ARG:HH22	3:H:427:GLU:CD	2.19	0.46
3:F:533:ASP:OD1	3:F:535:ALA:HB3	2.16	0.46
3:H:547:HIS:CE1	3:H:560:PRO:HA	2.52	0.46
2:I:33:ALA:HB1	3:N:260:SER:HB3	1.96	0.46
3:J:46:LYS:NZ	3:N:237:GLU:OE1	2.38	0.46
3:J:561:ASP:O	3:J:563:PRO:HD3	2.16	0.46
3:L:58:ILE:O	3:L:58:ILE:CG2	2.62	0.46
3:L:203:TYR:HB3	3:L:206:GLU:CB	2.39	0.46
3:L:367:SER:O	3:L:368:THR:CB	2.64	0.46
3:L:385:VAL:HA	3:L:390:ASN:OD1	2.16	0.46
3:L:431:ILE:HD11	3:L:437:ILE:CG2	2.45	0.46
3:N:525:ILE:HD11	3:N:604:VAL:HA	1.97	0.46
3:P:137:ILE:HD13	3:P:142:ILE:HG12	1.97	0.46

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:P:178:SER:O	3:P:179:ALA:CB	2.64	0.46
3:P:354:SER:HB3	3:P:419:MET:HG2	1.97	0.46
3:P:547:HIS:CE1	3:P:560:PRO:HA	2.51	0.46
3:R:67:ARG:HH11	3:R:67:ARG:CG	2.29	0.46
3:R:341:MET:HG3	3:R:431:ILE:HD11	1.98	0.46
3:R:547:HIS:CE1	3:R:560:PRO:HA	2.51	0.46
3:R:549:LEU:C	3:R:549:LEU:CD1	2.87	0.46
3:T:346:ASP:CB	3:T:427:GLU:HA	2.46	0.46
3:T:525:ILE:HD11	3:T:604:VAL:HA	1.97	0.46
3:T:540:VAL:CG2	3:T:566:TRP:HA	2.46	0.46
4:U:132:LEU:O	4:U:133:SER:C	2.59	0.46
4:U:285:LEU:HD13	4:U:285:LEU:O	2.16	0.46
5:W:23:LEU:HD23	5:W:24:PRO:N	2.31	0.46
5:W:399:ARG:HA	5:W:745:MET:HA	1.98	0.46
5:W:463:ASP:OD1	5:W:464:TYR:N	2.49	0.46
5:W:574:THR:HB	5:W:611:THR:HA	1.97	0.46
5:W:1054:SER:HB2	5:W:1112:LEU:CD2	2.46	0.46
5:W:1139:PRO:HA	5:W:1140:PRO:HD3	1.82	0.46
5:W:1228:ARG:O	5:W:1229:ILE:C	2.59	0.46
6:X:406:LEU:HD12	6:X:409:MET:HE3	1.98	0.46
6:Y:559:HIS:HD2	6:Y:562:SER:HB2	1.81	0.46
6:Y:780:ILE:HG12	6:Y:938:TYR:CD1	2.51	0.46
1:b:29:ARG:NH2	1:b:65:ILE:O	2.49	0.45
1:d:5:MET:O	1:d:9:VAL:CG2	2.64	0.45
1:h:58:PRO:CG	3:H:423:GLN:NE2	2.79	0.45
1:j:57:ALA:HB1	1:j:59:HIS:CD2	2.51	0.45
1:j:59:HIS:CB	3:J:398:SER:HA	2.39	0.45
1:r:3:LEU:HD11	3:T:492:ASN:ND2	2.28	0.45
1:r:44:GLY:O	3:R:399:ALA:C	2.58	0.45
3:B:346:ASP:CB	3:B:427:GLU:HA	2.46	0.45
3:B:385:VAL:HA	3:B:390:ASN:OD1	2.16	0.45
3:B:561:ASP:O	3:B:563:PRO:HD3	2.16	0.45
2:C:24:SER:HB2	3:D:175:SER:C	2.41	0.45
3:D:346:ASP:CB	3:D:427:GLU:HA	2.46	0.45
3:D:418:ASN:CG	3:F:337:ARG:NH2	2.74	0.45
3:F:58:ILE:CG1	3:F:127:LEU:HD23	2.46	0.45
3:F:69:GLY:HA3	3:F:139:THR:HB	1.97	0.45
3:F:169:MET:HG2	3:F:169:MET:O	2.15	0.45
3:F:346:ASP:CB	3:F:427:GLU:HA	2.46	0.45
3:F:369:SER:HB2	3:F:445:ILE:HA	1.97	0.45
3:F:521:VAL:HG21	3:F:577:VAL:HG22	1.95	0.45

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:J:169:MET:HG2	3:J:169:MET:O	2.15	0.45
3:J:354:SER:HB3	3:J:419:MET:HG2	1.97	0.45
3:J:367:SER:O	3:J:368:THR:CB	2.64	0.45
3:J:547:HIS:CE1	3:J:560:PRO:HA	2.51	0.45
3:J:609:ARG:HD2	2:K:29:THR:HG22	1.91	0.45
3:L:599:ALA:O	3:L:600:LEU:C	2.60	0.45
3:N:533:ASP:OD1	3:N:535:ALA:HB3	2.16	0.45
3:N:561:ASP:O	3:N:563:PRO:HD3	2.16	0.45
3:P:169:MET:HG2	3:P:169:MET:O	2.15	0.45
3:R:53:THR:HG22	3:R:54:SER:N	2.31	0.45
4:U:41:PHE:HZ	4:U:186:ALA:HB3	1.81	0.45
4:U:89:GLY:CA	4:U:103:VAL:HG23	2.46	0.45
4:V:192:PHE:CB	6:Y:376:LEU:HD23	2.41	0.45
5:W:27:PHE:CZ	5:W:99:LEU:HD21	2.51	0.45
5:W:440:THR:HG22	5:W:647:PRO:HB3	1.97	0.45
5:W:448:ASP:O	5:W:671:THR:HG21	2.16	0.45
5:W:452:TYR:HD2	5:W:672:THR:HG23	1.81	0.45
5:W:934:ASN:HB3	5:W:1023:THR:HG22	1.98	0.45
5:W:958:PHE:O	5:W:959:ASN:HB2	2.16	0.45
6:X:388:MET:HA	6:X:1193:SER:HA	1.98	0.45
6:X:560:LEU:N	6:X:561:PRO:HD3	2.31	0.45
6:X:578:ASP:C	6:X:580:HIS:H	2.24	0.45
6:X:1131:ARG:O	6:Y:106:MET:HG3	2.16	0.45
6:Y:773:ARG:HG3	6:Y:773:ARG:O	2.16	0.45
6:Y:787:ARG:O	6:Y:788:VAL:HB	2.15	0.45
6:Y:1084:THR:CG2	6:Y:1114:PRO:HB2	2.46	0.45
6:Y:1169:TRP:HE3	6:Y:1191:TYR:HB2	1.80	0.45
1:b:22:LEU:HB2	1:b:83:VAL:HG21	1.97	0.45
1:b:60:ALA:C	1:b:62:VAL:N	2.74	0.45
1:p:43:CYS:H	1:p:44:GLY:HA2	1.80	0.45
1:r:29:ARG:NH2	1:r:65:ILE:O	2.49	0.45
1:t:24:LEU:HD12	1:t:41:VAL:CG2	2.35	0.45
1:t:44:GLY:HA3	3:T:322:TYR:CE2	2.51	0.45
3:B:137:ILE:HD13	3:B:142:ILE:HG12	1.97	0.45
3:D:547:HIS:CE1	3:D:560:PRO:HA	2.51	0.45
3:F:341:MET:CE	3:F:341:MET:CA	2.88	0.45
3:H:53:THR:HG22	3:H:54:SER:N	2.32	0.45
3:H:227:LEU:O	3:H:228:PRO:O	2.33	0.45
3:H:536:THR:HG22	3:H:573:ALA:HB3	1.97	0.45
3:J:58:ILE:HG23	3:J:124:GLY:CA	2.32	0.45
3:J:92:TRP:CD2	3:J:156:VAL:HG23	2.51	0.45

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:J:275:GLU:HG2	3:J:275:GLU:O	2.15	0.45
3:J:346:ASP:CB	3:J:427:GLU:HA	2.46	0.45
3:J:416:ARG:HG3	3:L:298:PHE:HZ	1.81	0.45
3:J:600:LEU:HG	3:J:601:THR:N	2.31	0.45
2:K:6:THR:HG23	2:K:7:SER:N	2.31	0.45
7:K:101:MYR:C4	3:L:187:LEU:HD22	2.45	0.45
3:L:255:SER:HA	3:L:258:THR:HG22	1.99	0.45
3:L:354:SER:HB3	3:L:419:MET:HG2	1.97	0.45
3:L:525:ILE:HD11	3:L:604:VAL:HA	1.97	0.45
3:N:67:ARG:HH11	3:N:67:ARG:CG	2.29	0.45
3:N:340:ASP:O	3:N:344:MET:HG3	2.16	0.45
3:P:275:GLU:HG2	3:P:275:GLU:O	2.15	0.45
3:R:521:VAL:HG21	3:R:577:VAL:HG22	1.95	0.45
3:R:599:ALA:O	3:R:600:LEU:C	2.60	0.45
3:T:44:THR:O	3:T:119:VAL:HG22	2.17	0.45
3:T:330:GLN:O	3:T:331:LEU:HB2	2.16	0.45
3:T:533:ASP:OD1	3:T:535:ALA:HB3	2.16	0.45
4:U:241:LEU:C	6:X:1179:VAL:HG11	2.41	0.45
5:W:133:ILE:HG13	5:W:134:PHE:N	2.31	0.45
5:W:751:SER:C	5:W:753:GLY:N	2.73	0.45
5:W:766:LEU:HD12	5:W:1004:VAL:CG2	2.46	0.45
5:W:967:TYR:HD1	5:W:968:ALA:O	1.99	0.45
5:W:997:TRP:C	5:W:997:TRP:CD1	2.94	0.45
6:X:362:THR:CG2	6:X:1173:VAL:HB	2.46	0.45
6:X:456:VAL:HG23	6:X:668:LEU:HD11	1.97	0.45
6:X:594:ALA:CA	6:X:608:GLN:NE2	2.70	0.45
6:X:694:ASP:O	6:X:695:SER:CB	2.64	0.45
6:X:1068:LYS:C	6:X:1070:GLY:N	2.73	0.45
6:Y:284:ASP:C	6:Y:286:ALA:N	2.74	0.45
6:Y:331:ARG:CD	6:Y:387:CYS:SG	3.01	0.45
6:Y:574:PRO:HB2	6:Y:679:TRP:CH2	2.52	0.45
1:l:61:ASN:HB2	3:L:396:VAL:O	2.16	0.45
1:p:61:ASN:CB	3:P:396:VAL:O	2.65	0.45
1:t:61:ASN:HB2	3:T:395:THR:HG22	1.98	0.45
3:B:354:SER:HB3	3:B:419:MET:HG2	1.97	0.45
3:B:390:ASN:HD22	3:B:390:ASN:HA	1.60	0.45
3:D:175:SER:CB	3:D:627:ARG:HD3	2.47	0.45
3:D:227:LEU:O	3:D:228:PRO:O	2.33	0.45
3:D:345:ILE:HG23	3:D:475:THR:OG1	2.16	0.45
3:F:468:PRO:O	3:H:578:SER:OG	2.21	0.45
3:F:564:LEU:HD23	3:F:564:LEU:C	2.40	0.45

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:G:6:THR:HG23	2:G:7:SER:N	2.30	0.45
2:G:24:SER:HB2	3:H:175:SER:C	2.41	0.45
3:H:44:THR:O	3:H:119:VAL:HG22	2.17	0.45
3:H:354:SER:HB3	3:H:419:MET:HG2	1.97	0.45
3:J:44:THR:O	3:J:119:VAL:HG22	2.17	0.45
3:J:255:SER:HA	3:J:258:THR:HG22	1.98	0.45
3:J:345:ILE:HG23	3:J:475:THR:OG1	2.16	0.45
3:J:469:VAL:HG21	3:L:575:MET:HE3	0.69	0.45
3:J:533:ASP:OD1	3:J:535:ALA:HB3	2.16	0.45
3:J:540:VAL:CG2	3:J:566:TRP:HA	2.46	0.45
3:J:562:TYR:HA	3:J:565:LEU:HB2	1.99	0.45
3:L:418:ASN:HD22	3:L:418:ASN:HA	1.51	0.45
3:N:69:GLY:HA3	3:N:139:THR:HB	1.97	0.45
3:N:255:SER:HA	3:N:258:THR:HG22	1.98	0.45
3:P:559:PRO:O	3:P:560:PRO:C	2.58	0.45
3:P:599:ALA:O	3:P:600:LEU:C	2.60	0.45
3:R:58:ILE:CG1	3:R:127:LEU:HD23	2.45	0.45
3:R:340:ASP:O	3:R:344:MET:HG3	2.16	0.45
3:R:418:ASN:HD22	3:R:418:ASN:HA	1.50	0.45
3:T:431:ILE:O	3:T:431:ILE:HG23	2.10	0.45
3:T:562:TYR:HA	3:T:565:LEU:HB2	1.99	0.45
4:U:52:PRO:HG2	4:U:64:ARG:NH2	2.31	0.45
5:W:152:THR:HG23	5:W:153:LYS:N	2.32	0.45
5:W:300:PHE:HE1	5:W:309:ASN:HD22	1.64	0.45
5:W:489:LYS:O	5:W:492:ALA:HB3	2.17	0.45
5:W:885:TRP:HA	5:W:888:ILE:CD1	2.46	0.45
6:X:283:ILE:HD11	6:X:288:PHE:N	2.31	0.45
6:X:308:ASN:ND2	6:X:395:THR:HG23	2.30	0.45
6:X:325:THR:HG22	6:X:347:ARG:HG2	1.99	0.45
6:X:337:PRO:HG2	6:X:338:GLU:H	1.82	0.45
6:X:389:GLU:O	6:X:390:LEU:C	2.59	0.45
6:X:651:PRO:CG	6:X:679:TRP:CZ3	2.99	0.45
6:X:736:MET:C	6:X:738:GLN:H	2.24	0.45
6:X:839:LYS:HB2	6:X:839:LYS:NZ	2.31	0.45
6:X:1013:VAL:CG1	6:X:1014:VAL:H	2.29	0.45
6:X:1122:PRO:HA	6:X:1159:GLY:CA	2.42	0.45
6:X:1130:ALA:CB	6:Y:104:GLN:HA	2.47	0.45
6:X:1140:GLU:OE2	6:X:1140:GLU:HA	2.16	0.45
6:X:1154:ILE:O	6:X:1155:ASP:HB2	2.16	0.45
6:X:1208:MET:HE2	6:X:1208:MET:HB3	1.87	0.45
6:Y:106:MET:HE3	6:Y:108:VAL:CG2	2.45	0.45

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
6:Y:283:ILE:HD11	6:Y:288:PHE:C	2.41	0.45
6:Y:312:ASN:O	6:Y:313:LYS:C	2.60	0.45
6:Y:450:ASP:C	6:Y:450:ASP:OD1	2.58	0.45
6:Y:486:LEU:HD21	6:Y:825:GLY:HA3	1.99	0.45
6:Y:751:CYS:N	6:Y:752:PRO:CD	2.80	0.45
1:l:44:GLY:HA3	3:L:322:TYR:CE2	2.51	0.45
1:b:59:HIS:CA	3:B:398:SER:HA	2.47	0.45
1:d:15:ARG:NE	3:D:406:ASN:HD21	1.97	0.45
1:d:61:ASN:HB2	3:D:395:THR:HG22	1.98	0.45
2:A:6:THR:HG23	2:A:7:SER:N	2.31	0.45
3:B:357:ILE:CG2	3:B:358:LEU:N	2.80	0.45
3:D:357:ILE:CG2	3:D:358:LEU:N	2.80	0.45
3:D:525:ILE:HD11	3:D:604:VAL:HA	1.98	0.45
3:D:536:THR:HG22	3:D:573:ALA:HB3	1.97	0.45
3:F:193:ILE:H	3:F:193:ILE:HG12	1.55	0.45
3:F:340:ASP:O	3:F:344:MET:HG3	2.16	0.45
3:F:547:HIS:CE1	3:F:560:PRO:HA	2.51	0.45
3:H:345:ILE:HG23	3:H:475:THR:OG1	2.16	0.45
3:H:346:ASP:CB	3:H:427:GLU:HA	2.46	0.45
3:H:357:ILE:CG2	3:H:358:LEU:N	2.80	0.45
3:H:385:VAL:HA	3:H:390:ASN:OD1	2.16	0.45
3:H:533:ASP:OD1	3:H:535:ALA:HB3	2.16	0.45
3:J:49:ARG:HB2	3:J:64:VAL:CG2	2.43	0.45
3:J:170:LEU:O	3:J:171:THR:C	2.60	0.45
3:L:69:GLY:HA3	3:L:139:THR:HB	1.97	0.45
3:L:178:SER:HB3	3:L:179:ALA:H	1.65	0.45
3:L:369:SER:HB2	3:L:445:ILE:HA	1.97	0.45
2:M:26:MET:O	2:M:26:MET:CG	2.60	0.45
3:N:367:SER:O	3:N:368:THR:CB	2.64	0.45
3:N:540:VAL:CG2	3:N:566:TRP:HA	2.46	0.45
2:O:36:LEU:HD22	2:O:41:LEU:HD11	1.98	0.45
3:P:69:GLY:HA3	3:P:139:THR:HB	1.97	0.45
3:R:92:TRP:CD2	3:R:156:VAL:HG23	2.51	0.45
3:R:533:ASP:OD1	3:R:535:ALA:HB3	2.16	0.45
3:T:471:LEU:HD23	3:T:471:LEU:N	2.32	0.45
4:U:225:ARG:HE	4:U:346:LEU:CD2	2.30	0.45
4:U:335:ALA:HA	4:U:356:ASN:OD1	2.16	0.45
5:W:282:VAL:CG1	5:W:283:THR:N	2.79	0.45
5:W:306:ARG:HD3	5:W:341:MET:HE2	1.99	0.45
5:W:341:MET:HA	5:W:341:MET:CE	2.42	0.45
5:W:415:ASP:O	5:W:419:GLU:HG3	2.16	0.45

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
5:W:438:VAL:HB	5:W:649:ILE:CD1	2.43	0.45
5:W:1087:LEU:HB2	5:W:1098:LEU:HD13	1.97	0.45
5:W:1261:THR:HG23	5:W:1262:THR:N	2.32	0.45
6:X:264:PRO:HB2	6:X:268:ASP:OD2	2.17	0.45
6:X:415:GLN:C	6:X:417:VAL:H	2.25	0.45
6:X:446:ALA:HB2	6:X:854:ASN:O	2.16	0.45
6:X:483:VAL:O	6:X:483:VAL:CG2	2.63	0.45
6:X:614:GLN:OE1	6:X:614:GLN:HA	2.15	0.45
6:X:780:ILE:HD13	6:X:940:GLN:HG3	1.98	0.45
6:X:914:ASP:C	6:X:915:LEU:HD23	2.42	0.45
6:X:1087:TYR:HB3	6:X:1121:ILE:CG1	2.45	0.45
6:Y:287:GLN:HB3	6:Y:1113:PRO:CA	2.47	0.45
6:Y:575:MET:HG2	6:Y:818:ILE:HG23	1.98	0.45
6:Y:1023:ARG:HD2	6:Y:1027:LEU:CD2	2.46	0.45
6:Y:1074:VAL:HG12	6:Y:1075:LEU:N	2.30	0.45
6:Y:1119:ILE:HG23	6:Y:1120:PRO:HD2	1.97	0.45
1:h:43:CYS:H	1:h:44:GLY:HA2	1.80	0.45
1:n:61:ASN:HB2	3:N:396:VAL:O	2.16	0.45
1:r:74:GLN:C	1:r:74:GLN:CD	2.85	0.45
3:D:333:PRO:HG3	3:D:454:PHE:C	2.42	0.45
3:D:562:TYR:HA	3:D:565:LEU:HB2	1.99	0.45
3:F:139:THR:CA	3:F:142:ILE:CG2	2.90	0.45
3:F:175:SER:CB	3:F:627:ARG:HD3	2.47	0.45
3:F:333:PRO:HG3	3:F:454:PHE:C	2.42	0.45
3:H:255:SER:HA	3:H:258:THR:HG22	1.98	0.45
2:I:6:THR:HG23	2:I:7:SER:N	2.31	0.45
2:I:29:THR:HG21	3:N:609:ARG:CD	2.47	0.45
3:J:235:LEU:HD13	3:J:235:LEU:HA	1.80	0.45
3:J:298:PHE:HZ	3:N:416:ARG:HG3	1.81	0.45
3:J:471:LEU:HD23	3:J:471:LEU:N	2.32	0.45
3:L:175:SER:CB	3:L:627:ARG:HD3	2.47	0.45
3:P:44:THR:O	3:P:119:VAL:HG22	2.17	0.45
3:P:61:LEU:HD21	3:P:118:TYR:CD2	2.51	0.45
3:P:357:ILE:CG2	3:P:358:LEU:N	2.80	0.45
3:P:540:VAL:CG2	3:P:566:TRP:HA	2.46	0.45
3:R:371:PRO:HG3	3:R:407:LEU:HD23	1.99	0.45
3:T:322:TYR:CE1	3:T:396:VAL:HB	2.51	0.45
3:T:371:PRO:HG3	3:T:407:LEU:HD23	1.99	0.45
3:T:599:ALA:O	3:T:600:LEU:C	2.60	0.45
4:U:121:ASP:OD1	4:U:123:ASN:HB2	2.16	0.45
4:U:176:ALA:HB1	6:X:413:PRO:HD2	1.94	0.45

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
4:U:184:HIS:C	4:U:187:GLN:HB3	2.42	0.45
4:U:217:GLY:C	4:U:223:LEU:HB2	2.42	0.45
4:U:344:ASP:C	4:U:346:LEU:H	2.23	0.45
4:V:214:ILE:CD1	4:V:232:PHE:HB2	2.46	0.45
5:W:518:TYR:O	5:W:554:PHE:N	2.50	0.45
5:W:554:PHE:O	5:W:558:VAL:HB	2.16	0.45
5:W:606:ALA:C	5:W:608:GLN:H	2.23	0.45
5:W:818:PHE:C	5:W:818:PHE:CD1	2.93	0.45
5:W:843:LEU:CD2	5:W:854:MET:CE	2.94	0.45
6:X:557:ILE:CB	6:X:592:LEU:HD21	2.46	0.45
6:X:1085:LEU:HD22	6:X:1119:ILE:HD11	1.98	0.45
6:Y:358:GLN:HE21	6:Y:1154:ILE:HD13	1.81	0.45
6:Y:567:LEU:HD22	6:Y:567:LEU:O	2.17	0.45
6:Y:990:ASN:CB	6:Y:1131:ARG:HA	2.45	0.45
6:Y:1083:TYR:CD1	6:Y:1083:TYR:C	2.95	0.45
6:Y:1117:PHE:CZ	6:Y:1138:THR:HG23	2.52	0.45
6:Y:1123:GLN:HG2	6:Y:1127:CYS:SG	2.57	0.45
1:b:15:ARG:HH12	3:B:402:ALA:CB	2.22	0.45
1:j:24:LEU:HB3	1:j:38:ALA:HB1	1.99	0.45
1:n:51:PHE:CE1	1:n:71:CYS:O	2.70	0.45
1:p:16:ALA:C	1:p:22:LEU:CD2	2.90	0.45
1:p:45:ARG:CB	3:P:399:ALA:O	2.61	0.45
2:C:6:THR:HG23	2:C:7:SER:N	2.31	0.45
3:D:199:LEU:HB3	3:D:207:PHE:CE2	2.52	0.45
3:D:385:VAL:HA	3:D:390:ASN:OD1	2.16	0.45
3:F:53:THR:HG22	3:F:54:SER:N	2.32	0.45
3:F:203:TYR:HB3	3:F:206:GLU:CB	2.39	0.45
3:H:367:SER:O	3:H:368:THR:CB	2.64	0.45
3:J:175:SER:CB	3:J:627:ARG:HD3	2.47	0.45
3:J:341:MET:HG3	3:J:431:ILE:HD11	1.98	0.45
3:L:53:THR:HG22	3:L:54:SER:N	2.32	0.45
3:L:360:TYR:CD1	3:L:361:ASP:N	2.77	0.45
3:L:369:SER:HB2	3:L:444:LEU:O	2.16	0.45
3:L:533:ASP:OD1	3:L:535:ALA:HB3	2.16	0.45
3:N:175:SER:CB	3:N:627:ARG:HD3	2.47	0.45
3:N:178:SER:O	3:N:179:ALA:CB	2.64	0.45
3:N:385:VAL:HA	3:N:390:ASN:OD1	2.16	0.45
3:N:419:MET:HE2	3:N:419:MET:HB3	1.80	0.45
2:O:6:THR:HG23	2:O:7:SER:N	2.30	0.45
3:P:96:PHE:CD1	3:P:101:THR:HG23	2.44	0.45
3:P:442:THR:OG1	3:P:443:PHE:N	2.50	0.45

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:P:561:ASP:O	3:P:563:PRO:HD3	2.16	0.45
3:R:137:ILE:HD13	3:R:142:ILE:HG12	1.98	0.45
3:R:170:LEU:O	3:R:171:THR:C	2.60	0.45
3:R:175:SER:CB	3:R:627:ARG:HD3	2.47	0.45
3:R:257:SER:C	3:R:259:LEU:N	2.74	0.45
3:R:357:ILE:CG2	3:R:358:LEU:N	2.80	0.45
3:T:358:LEU:CD1	3:T:417:PHE:CE1	2.62	0.45
4:U:228:THR:HG22	4:U:233:TYR:HD2	1.82	0.45
5:W:63:PRO:CB	5:W:64:PRO:CD	2.92	0.45
5:W:625:ARG:CG	5:W:625:ARG:O	2.65	0.45
5:W:737:SER:HB2	5:W:777:VAL:HG13	1.98	0.45
5:W:998:LEU:O	5:W:1000:TYR:N	2.50	0.45
5:W:1225:ASP:C	5:W:1227:TYR:N	2.73	0.45
6:X:413:PRO:HA	6:X:416:ILE:HG12	1.99	0.45
6:X:562:SER:HB3	6:X:592:LEU:HD11	1.98	0.45
6:X:572:ALA:CB	6:X:818:ILE:HG21	2.47	0.45
6:X:610:THR:O	6:X:612:ALA:N	2.50	0.45
6:X:672:SER:O	6:X:673:ASN:HB2	2.16	0.45
6:X:910:LYS:O	6:X:914:ASP:HA	2.17	0.45
6:X:985:VAL:HG22	6:X:1137:PHE:CE2	2.52	0.45
6:X:1055:SER:O	6:X:1056:VAL:C	2.59	0.45
6:Y:595:GLN:H	6:Y:596:PRO:HD2	1.82	0.45
6:Y:628:PRO:CG	6:Y:647:ARG:HG3	2.47	0.45
6:Y:1167:LEU:O	6:Y:1170:PRO:HD3	2.17	0.45
1:l:44:GLY:C	3:L:399:ALA:HB3	2.42	0.45
1:f:29:ARG:NH2	1:f:65:ILE:O	2.49	0.45
1:h:45:ARG:CD	3:H:400:ALA:CA	2.89	0.45
3:B:175:SER:CB	3:B:627:ARG:HD3	2.47	0.45
3:B:431:ILE:HD11	3:B:437:ILE:CG2	2.45	0.45
3:B:599:ALA:O	3:B:600:LEU:C	2.60	0.45
2:C:33:ALA:HB1	3:H:260:SER:HB3	1.96	0.45
2:C:36:LEU:HD22	2:C:41:LEU:HD11	1.98	0.45
3:D:46:LYS:NZ	3:H:237:GLU:CB	2.68	0.45
3:D:255:SER:HA	3:D:258:THR:HG22	1.98	0.45
3:D:468:PRO:O	3:F:578:SER:OG	2.21	0.45
3:F:92:TRP:CD2	3:F:156:VAL:HG23	2.51	0.45
3:F:345:ILE:HD12	3:F:468:PRO:HB3	1.99	0.45
3:F:367:SER:O	3:F:368:THR:CB	2.64	0.45
3:F:471:LEU:HD23	3:F:471:LEU:N	2.32	0.45
3:F:561:ASP:O	3:F:563:PRO:HD3	2.16	0.45
2:G:23:THR:C	2:G:25:ASP:N	2.75	0.45

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:H:600:LEU:O	3:H:601:THR:C	2.60	0.45
3:H:644:LEU:N	3:H:644:LEU:HD23	2.32	0.45
2:I:36:LEU:HD22	2:I:41:LEU:HD11	1.98	0.45
3:J:371:PRO:HG3	3:J:407:LEU:HD23	1.99	0.45
3:J:385:VAL:HA	3:J:390:ASN:OD1	2.16	0.45
3:J:600:LEU:O	3:J:601:THR:C	2.60	0.45
3:L:257:SER:C	3:L:259:LEU:N	2.74	0.45
3:L:416:ARG:HG3	3:N:298:PHE:HZ	1.81	0.45
3:N:341:MET:HG3	3:N:431:ILE:HD11	1.98	0.45
3:N:345:ILE:HD12	3:N:468:PRO:HB3	1.99	0.45
3:P:107:VAL:HG11	3:R:128:ASP:CB	2.45	0.45
3:P:175:SER:CB	3:P:627:ARG:HD3	2.47	0.45
7:Q:101:MYR:C4	3:R:187:LEU:HD22	2.45	0.45
3:R:346:ASP:CB	3:R:427:GLU:HA	2.46	0.45
3:R:367:SER:O	3:R:368:THR:CB	2.64	0.45
3:R:536:THR:HG22	3:R:573:ALA:HB3	1.98	0.45
3:T:175:SER:CB	3:T:627:ARG:HD3	2.47	0.45
3:T:333:PRO:HG3	3:T:454:PHE:C	2.42	0.45
3:T:600:LEU:O	3:T:601:THR:C	2.60	0.45
3:T:644:LEU:N	3:T:644:LEU:HD23	2.32	0.45
4:U:56:ALA:HB2	6:X:1185:PRO:HG2	1.99	0.45
4:U:63:SER:C	4:U:66:ILE:HG22	2.42	0.45
5:W:39:LEU:N	5:W:89:HIS:HE1	2.12	0.45
5:W:67:ALA:O	5:W:174:GLY:HA2	2.16	0.45
5:W:246:ASP:HB2	5:W:255:ILE:C	2.42	0.45
5:W:382:PRO:HG2	5:W:383:PHE:CD1	2.52	0.45
5:W:654:GLU:O	5:W:654:GLU:HG3	2.17	0.45
5:W:981:LEU:CD1	5:W:1028:ILE:HG21	2.44	0.45
5:W:985:ALA:HA	5:W:1034:PRO:HB3	1.98	0.45
5:W:1249:HIS:HA	5:W:1259:SER:HA	1.98	0.45
6:Y:73:ASN:HB3	6:Y:79:ILE:HB	1.99	0.45
6:Y:200:LEU:HD21	6:Y:343:ALA:CB	2.47	0.45
6:Y:207:LEU:HD22	6:Y:244:VAL:CB	2.47	0.45
6:Y:602:GLY:HA3	6:Y:623:PRO:CG	2.47	0.45
6:Y:661:THR:OG1	6:Y:662:LEU:HD12	2.15	0.45
6:Y:769:VAL:HB	6:Y:780:ILE:HD13	1.98	0.45
6:Y:1081:TYR:OH	6:Y:1138:THR:HG22	2.17	0.45
1:b:15:ARG:HE	3:B:406:ASN:CG	2.24	0.45
1:j:6:ILE:HD13	3:L:586:VAL:HG21	1.99	0.45
1:p:16:ALA:O	1:p:22:LEU:HD23	2.16	0.45
1:p:62:VAL:CB	1:p:65:ILE:HD12	2.47	0.45

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:t:14:VAL:O	1:t:18:ALA:N	2.47	0.45
3:B:533:ASP:OD1	3:B:535:ALA:HB3	2.16	0.45
3:D:343:ALA:CB	3:D:478:GLY:O	2.62	0.45
3:D:600:LEU:O	3:D:601:THR:C	2.60	0.45
2:E:36:LEU:HD22	2:E:41:LEU:HD11	1.98	0.45
3:F:199:LEU:HB3	3:F:207:PHE:CE2	2.52	0.45
3:F:252:LEU:HB3	3:F:613:LEU:CD2	2.47	0.45
3:F:443:PHE:HE2	3:F:459:LEU:HD21	1.74	0.45
3:H:175:SER:CB	3:H:627:ARG:HD3	2.47	0.45
3:H:333:PRO:HG3	3:H:454:PHE:C	2.42	0.45
3:H:346:ASP:HB3	3:H:427:GLU:HA	1.99	0.45
2:I:26:MET:CG	3:J:216:PRO:HG3	2.29	0.45
3:J:199:LEU:HB3	3:J:207:PHE:CE2	2.52	0.45
3:J:262:ALA:CB	3:J:608:LEU:HD13	2.47	0.45
3:L:44:THR:O	3:L:119:VAL:HG22	2.17	0.45
3:L:139:THR:CA	3:L:142:ILE:HG23	2.45	0.45
3:L:169:MET:HG2	3:L:169:MET:O	2.15	0.45
3:L:199:LEU:HB3	3:L:207:PHE:CE2	2.52	0.45
3:L:330:GLN:O	3:L:331:LEU:HB2	2.16	0.45
3:L:341:MET:CE	3:L:341:MET:CA	2.88	0.45
3:N:95:LEU:HD12	3:N:95:LEU:HA	1.71	0.45
3:N:170:LEU:O	3:N:171:THR:C	2.60	0.45
3:N:257:SER:C	3:N:259:LEU:N	2.74	0.45
3:N:428:ARG:NH1	3:N:428:ARG:CG	2.76	0.45
3:P:345:ILE:HG23	3:P:475:THR:HG1	1.81	0.45
3:P:533:ASP:OD1	3:P:535:ALA:HB3	2.16	0.45
3:R:561:ASP:O	3:R:563:PRO:HD3	2.16	0.45
3:R:562:TYR:HA	3:R:565:LEU:HB2	1.99	0.45
3:T:101:THR:HG21	3:T:159:TRP:HZ2	1.82	0.45
3:T:367:SER:O	3:T:368:THR:CB	2.64	0.45
4:U:224:ASN:CG	4:U:346:LEU:HD11	2.42	0.45
5:W:19:ARG:NH1	5:W:285:LEU:HB2	2.31	0.45
5:W:247:VAL:HG12	5:W:248:LEU:N	2.30	0.45
5:W:717:ASP:N	5:W:718:PRO:CD	2.78	0.45
5:W:765:PHE:C	5:W:765:PHE:CD1	2.94	0.45
5:W:775:ARG:CD	5:W:868:ASP:HB2	2.46	0.45
5:W:852:VAL:HG22	5:W:853:THR:N	2.31	0.45
5:W:1063:VAL:CG1	5:W:1108:PHE:HB3	2.47	0.45
6:X:191:PRO:O	6:X:193:LEU:HD13	2.17	0.45
6:X:842:ALA:N	6:X:843:PRO:HD2	2.31	0.45
6:X:1099:ILE:HG22	6:X:1099:ILE:O	2.17	0.45

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
6:Y:276:THR:CG2	6:Y:299:LEU:HD23	2.47	0.45
6:Y:847:SER:O	6:Y:849:ALA:N	2.42	0.45
6:Y:1082:VAL:O	6:Y:1114:PRO:CA	2.65	0.45
1:n:6:ILE:HG21	3:J:494:THR:HG22	1.99	0.45
1:n:23:THR:O	1:n:25:TYR:CD2	2.70	0.45
3:B:252:LEU:HB3	3:B:613:LEU:CD2	2.47	0.45
3:B:297:ALA:CB	3:B:460:MET:HB2	2.33	0.45
3:B:330:GLN:O	3:B:331:LEU:HB2	2.17	0.45
3:B:341:MET:HG3	3:B:431:ILE:HD11	1.98	0.45
3:B:345:ILE:HD12	3:B:468:PRO:HB3	1.99	0.45
3:B:562:TYR:HA	3:B:565:LEU:HB2	1.99	0.45
3:B:600:LEU:O	3:B:601:THR:C	2.60	0.45
3:D:471:LEU:HD23	3:D:471:LEU:N	2.32	0.45
3:F:170:LEU:O	3:F:171:THR:C	2.60	0.45
3:F:255:SER:HA	3:F:258:THR:HG22	1.99	0.45
3:F:458:THR:HG23	3:F:458:THR:O	2.17	0.45
3:H:67:ARG:HH11	3:H:67:ARG:CG	2.29	0.45
3:H:199:LEU:HB3	3:H:207:PHE:CE2	2.52	0.45
3:H:345:ILE:HD12	3:H:468:PRO:HB3	1.99	0.45
3:H:599:ALA:O	3:H:600:LEU:C	2.60	0.45
3:J:330:GLN:O	3:J:331:LEU:HB2	2.17	0.45
3:J:340:ASP:O	3:J:344:MET:HG3	2.16	0.45
3:L:238:VAL:HG22	3:N:142:ILE:HD11	1.97	0.45
3:N:96:PHE:CD1	3:N:101:THR:HG23	2.44	0.45
2:O:33:ALA:HB1	3:T:260:SER:HB3	1.96	0.45
3:P:533:ASP:CG	3:P:536:THR:HG23	2.42	0.45
3:R:44:THR:O	3:R:119:VAL:HG22	2.17	0.45
3:R:96:PHE:CD1	3:R:101:THR:HG23	2.44	0.45
3:R:141:GLN:HE22	3:R:144:ASN:ND2	2.15	0.45
3:R:252:LEU:HB3	3:R:613:LEU:CD2	2.47	0.45
3:R:333:PRO:HG3	3:R:454:PHE:C	2.42	0.45
2:S:6:THR:HG23	2:S:7:SER:N	2.30	0.45
3:T:69:GLY:HA3	3:T:139:THR:HB	1.97	0.45
3:T:340:ASP:O	3:T:344:MET:HG3	2.16	0.45
3:T:354:SER:HB3	3:T:419:MET:HG2	1.97	0.45
3:T:385:VAL:HA	3:T:390:ASN:OD1	2.16	0.45
3:T:547:HIS:CE1	3:T:560:PRO:HA	2.52	0.45
4:U:9:LEU:O	4:U:145:MET:HE3	2.17	0.45
4:U:32:CYS:SG	6:X:958:HIS:CD2	3.04	0.45
4:U:54:TRP:CH2	6:X:1181:THR:HG22	2.52	0.45
4:U:140:GLN:HE21	4:U:144:ILE:HG12	1.82	0.45

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
4:U:223:LEU:HA	4:U:227:ARG:O	2.17	0.45
4:V:175:MET:SD	6:Y:404:MET:HE3	2.57	0.45
5:W:207:TYR:CZ	5:W:237:PRO:CB	2.99	0.45
5:W:234:THR:OG1	5:W:284:ARG:NH2	2.50	0.45
6:X:559:HIS:CE1	6:X:561:PRO:HD2	2.52	0.45
6:X:741:THR:HG23	6:X:744:PRO:HG2	1.98	0.45
6:Y:801:ARG:HH11	6:Y:801:ARG:HG2	1.82	0.45
6:Y:1032:ILE:HG23	6:Y:1032:ILE:O	2.17	0.45
1:l:6:ILE:CD1	3:N:586:VAL:CB	2.95	0.45
1:p:35:PHE:HE2	1:p:73:ARG:HD3	1.79	0.45
3:B:346:ASP:HB3	3:B:427:GLU:HA	1.99	0.45
2:C:23:THR:C	2:C:25:ASP:N	2.75	0.45
3:D:141:GLN:HE22	3:D:144:ASN:ND2	2.15	0.45
3:D:252:LEU:HB3	3:D:613:LEU:CD2	2.47	0.45
3:D:341:MET:HG3	3:D:431:ILE:HD11	1.99	0.45
2:E:24:SER:CB	3:F:176:ALA:N	2.80	0.45
3:F:341:MET:HG3	3:F:431:ILE:HD11	1.99	0.45
3:F:357:ILE:CG2	3:F:358:LEU:N	2.80	0.45
3:H:169:MET:HG2	3:H:169:MET:O	2.15	0.45
3:H:183:TRP:CE2	3:H:187:LEU:CD1	3.00	0.45
3:H:340:ASP:O	3:H:344:MET:HG3	2.16	0.45
3:H:431:ILE:O	3:H:431:ILE:HG23	2.10	0.45
3:H:471:LEU:HD23	3:H:471:LEU:N	2.32	0.45
3:H:559:PRO:O	3:H:560:PRO:C	2.58	0.45
3:J:369:SER:CB	3:J:446:PRO:HD3	2.47	0.45
3:L:340:ASP:O	3:L:344:MET:HG3	2.16	0.45
3:N:322:TYR:O	3:N:322:TYR:CG	2.66	0.45
3:N:333:PRO:HG3	3:N:454:PHE:C	2.42	0.45
3:N:354:SER:HB3	3:N:419:MET:HG2	1.97	0.45
3:P:199:LEU:HB3	3:P:207:PHE:CE2	2.52	0.45
3:P:255:SER:HA	3:P:258:THR:HG22	1.98	0.45
3:P:345:ILE:HD12	3:P:468:PRO:HB3	1.99	0.45
3:R:262:ALA:CB	3:R:608:LEU:HD13	2.47	0.45
3:T:458:THR:O	3:T:458:THR:HG23	2.17	0.45
4:U:10:THR:HG22	4:U:11:TYR:H	1.82	0.45
4:U:38:THR:HA	4:U:190:TYR:OH	2.16	0.45
4:U:54:TRP:CZ2	6:X:1181:THR:HA	2.52	0.45
4:U:123:ASN:C	4:U:125:TYR:N	2.73	0.45
4:U:251:ASN:ND2	4:U:253:GLY:H	2.15	0.45
4:U:386:ILE:O	4:U:387:GLN:HB3	2.17	0.45
5:W:102:ARG:CG	5:W:102:ARG:NH1	2.78	0.45

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
5:W:126:LEU:C	5:W:128:ALA:H	2.25	0.45
5:W:145:ASP:CG	5:W:148:ILE:HB	2.42	0.45
5:W:397:PRO:HB3	5:W:747:THR:HG21	1.99	0.45
5:W:923:ALA:C	5:W:925:GLY:N	2.73	0.45
5:W:1093:ASN:HB2	5:W:1095:GLU:HG2	1.99	0.45
6:X:197:ASP:OD1	6:X:253:GLY:HA3	2.17	0.45
6:X:518:LEU:CD1	6:X:820:VAL:HG23	2.47	0.45
6:X:534:SER:CB	6:X:543:THR:HG23	2.47	0.45
6:X:597:GLU:OE1	6:X:637:PRO:HD2	2.17	0.45
6:X:724:LEU:HD12	6:X:724:LEU:C	2.42	0.45
6:X:832:ILE:HB	6:X:835:HIS:CB	2.46	0.45
6:X:1021:VAL:HG23	6:X:1021:VAL:O	2.16	0.45
6:Y:218:GLU:HA	6:Y:218:GLU:OE1	2.17	0.45
6:Y:246:MET:CE	6:Y:262:ARG:HD2	2.46	0.45
6:Y:651:PRO:HB3	6:Y:679:TRP:CE3	2.51	0.45
1:f:21:ARG:NH1	3:F:327:TYR:OH	2.28	0.44
1:h:19:ALA:O	1:h:21:ARG:HG3	2.17	0.44
1:n:21:ARG:HH21	3:N:370:VAL:HG13	1.81	0.44
3:B:491:ALA:HB2	3:B:592:THR:CG2	2.48	0.44
3:D:330:GLN:HB2	3:D:457:PRO:HG3	1.99	0.44
3:D:431:ILE:HD11	3:D:437:ILE:CG2	2.45	0.44
3:F:44:THR:O	3:F:119:VAL:HG22	2.17	0.44
3:F:141:GLN:HE22	3:F:144:ASN:ND2	2.15	0.44
3:F:436:TYR:CZ	3:N:435:PRO:HG2	2.52	0.44
3:F:533:ASP:CG	3:F:536:THR:HG23	2.42	0.44
3:F:559:PRO:O	3:F:560:PRO:C	2.58	0.44
3:H:141:GLN:HE22	3:H:144:ASN:ND2	2.15	0.44
3:J:252:LEU:HB3	3:J:613:LEU:CD2	2.47	0.44
3:J:346:ASP:HB3	3:J:427:GLU:HA	1.99	0.44
3:J:442:THR:OG1	3:J:443:PHE:N	2.50	0.44
3:J:533:ASP:OD1	3:J:536:THR:HG23	2.17	0.44
3:L:170:LEU:O	3:L:171:THR:C	2.60	0.44
3:L:183:TRP:CE2	3:L:187:LEU:CD1	3.00	0.44
3:L:333:PRO:HG3	3:L:454:PHE:C	2.42	0.44
3:L:458:THR:O	3:L:458:THR:HG23	2.17	0.44
3:L:471:LEU:HD23	3:L:471:LEU:N	2.32	0.44
3:N:199:LEU:HB3	3:N:207:PHE:CE2	2.52	0.44
3:N:600:LEU:O	3:N:601:THR:C	2.60	0.44
3:P:330:GLN:O	3:P:331:LEU:HB2	2.17	0.44
3:P:333:PRO:HG3	3:P:454:PHE:C	2.42	0.44
3:P:346:ASP:HB3	3:P:427:GLU:HA	1.99	0.44

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:P:385:VAL:HA	3:P:390:ASN:OD1	2.16	0.44
3:P:562:TYR:HA	3:P:565:LEU:HB2	1.99	0.44
3:R:178:SER:O	3:R:179:ALA:CB	2.64	0.44
3:R:330:GLN:O	3:R:331:LEU:HB2	2.17	0.44
3:R:345:ILE:HD12	3:R:468:PRO:HB3	1.99	0.44
3:R:533:ASP:CG	3:R:536:THR:HG23	2.42	0.44
4:U:12:ASN:C	4:U:12:ASN:ND2	2.73	0.44
4:U:41:PHE:HD1	6:X:404:MET:HE2	1.80	0.44
4:U:393:ASP:O	4:U:396:GLN:HB3	2.17	0.44
4:V:196:LEU:HD13	4:V:196:LEU:HA	1.86	0.44
5:W:39:LEU:HD12	5:W:43:LEU:CB	2.46	0.44
5:W:391:VAL:HG11	5:W:790:ARG:HH11	1.82	0.44
5:W:448:ASP:HA	5:W:449:PRO:HD3	1.80	0.44
5:W:824:ARG:HA	5:W:847:PRO:HG2	1.98	0.44
6:X:247:THR:O	6:X:260:GLU:OE1	2.35	0.44
6:X:284:ASP:OD1	6:X:284:ASP:C	2.61	0.44
6:X:376:LEU:N	6:X:376:LEU:CD1	2.80	0.44
6:X:479:LEU:HD23	6:X:761:VAL:CG2	2.47	0.44
6:X:586:PHE:HA	6:X:642:PHE:CE2	2.52	0.44
6:X:697:MET:HE1	6:X:822:MET:HE2	2.00	0.44
6:X:1006:ARG:HH11	6:X:1013:VAL:CG2	2.29	0.44
6:X:1082:VAL:HB	6:X:1114:PRO:HB3	1.98	0.44
6:Y:282:TYR:OH	6:Y:289:ASN:ND2	2.51	0.44
6:Y:868:ARG:HD2	6:Y:932:ALA:O	2.17	0.44
1:b:59:HIS:CA	3:B:398:SER:CA	2.92	0.44
1:j:59:HIS:CA	3:J:398:SER:CA	2.93	0.44
3:B:333:PRO:HG3	3:B:454:PHE:C	2.42	0.44
3:B:525:ILE:HD11	3:B:604:VAL:HA	1.97	0.44
3:B:547:HIS:CE1	3:B:560:PRO:HA	2.51	0.44
3:D:416:ARG:HG3	3:F:298:PHE:HZ	1.82	0.44
3:D:458:THR:O	3:D:458:THR:HG23	2.17	0.44
3:D:540:VAL:CG2	3:D:566:TRP:HA	2.46	0.44
3:D:644:LEU:N	3:D:644:LEU:HD23	2.32	0.44
2:E:41:LEU:O	2:E:42:ASN:CB	2.37	0.44
3:F:330:GLN:HB2	3:F:457:PRO:HG3	1.99	0.44
3:F:343:ALA:CB	3:F:478:GLY:O	2.62	0.44
3:H:55:VAL:O	3:H:55:VAL:HG13	2.18	0.44
3:H:257:SER:C	3:H:259:LEU:N	2.74	0.44
3:H:458:THR:O	3:H:458:THR:HG23	2.17	0.44
3:H:533:ASP:CG	3:H:536:THR:HG23	2.42	0.44
3:H:561:ASP:O	3:H:563:PRO:HD3	2.16	0.44

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:J:333:PRO:HG3	3:J:454:PHE:C	2.42	0.44
3:L:262:ALA:CB	3:L:608:LEU:HD13	2.47	0.44
2:M:6:THR:HG23	2:M:7:SER:N	2.30	0.44
3:N:262:ALA:CB	3:N:608:LEU:HD13	2.47	0.44
3:N:599:ALA:O	3:N:600:LEU:C	2.60	0.44
3:P:208:TYR:CE2	3:P:212:ILE:HD11	2.53	0.44
3:P:533:ASP:OD1	3:P:536:THR:HG23	2.18	0.44
3:P:644:LEU:N	3:P:644:LEU:HD23	2.32	0.44
3:R:330:GLN:HB2	3:R:457:PRO:HG3	1.99	0.44
3:R:385:VAL:HA	3:R:390:ASN:OD1	2.16	0.44
3:R:416:ARG:HG3	3:T:298:PHE:HZ	1.81	0.44
3:R:458:THR:HG23	3:R:458:THR:O	2.17	0.44
3:R:644:LEU:HD23	3:R:644:LEU:N	2.32	0.44
2:S:26:MET:CG	3:T:216:PRO:HG3	2.29	0.44
3:T:183:TRP:CE2	3:T:187:LEU:CD1	3.01	0.44
3:T:199:LEU:HB3	3:T:207:PHE:CE2	2.52	0.44
3:T:330:GLN:HB2	3:T:457:PRO:HG3	1.99	0.44
3:T:369:SER:HB2	3:T:445:ILE:HA	1.97	0.44
3:T:491:ALA:HB2	3:T:592:THR:CG2	2.48	0.44
4:U:6:PHE:CD1	4:U:6:PHE:C	2.95	0.44
4:U:42:SER:C	4:U:43:HIS:CD2	2.95	0.44
4:U:150:LEU:HD13	4:U:300:ILE:HD13	1.99	0.44
4:U:283:GLY:HA2	4:U:286:MET:HE2	1.98	0.44
4:U:302:GLU:HG2	4:U:310:GLN:HE21	1.81	0.44
5:W:271:ARG:HH21	5:W:328:PRO:HG2	1.83	0.44
5:W:553:GLN:HB3	5:W:558:VAL:HG23	1.98	0.44
5:W:1108:PHE:CD1	5:W:1108:PHE:N	2.85	0.44
6:X:531:CYS:SG	6:X:813:VAL:HG11	2.58	0.44
6:X:757:VAL:HG11	6:X:841:PHE:CD2	2.52	0.44
6:X:865:VAL:HG11	6:X:955:SER:OG	2.16	0.44
6:X:973:ASN:ND2	6:X:1144:ASP:CB	2.81	0.44
6:X:1059:PHE:C	6:X:1059:PHE:CD1	2.96	0.44
6:X:1130:ALA:O	6:X:1132:ARG:HG3	2.17	0.44
6:Y:27:ALA:O	6:Y:28:PRO:C	2.59	0.44
6:Y:201:CYS:SG	6:Y:201:CYS:O	2.76	0.44
6:Y:202:THR:HG23	6:Y:391:CYS:HA	1.96	0.44
6:Y:231:MET:HE1	6:Y:347:ARG:NH2	2.32	0.44
6:Y:257:ARG:HE	6:Y:257:ARG:HB3	1.50	0.44
6:Y:358:GLN:NE2	6:Y:1154:ILE:HD13	2.32	0.44
1:l:24:LEU:CD1	1:l:33:THR:CA	2.86	0.44
1:d:17:ALA:O	1:d:82:LEU:CD1	2.65	0.44

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:h:45:ARG:N	3:H:399:ALA:O	2.50	0.44
1:j:20:GLY:HA2	1:j:83:VAL:HG22	1.99	0.44
1:n:15:ARG:HB3	3:N:402:ALA:HB2	1.99	0.44
1:n:29:ARG:NH2	1:n:65:ILE:O	2.49	0.44
1:p:23:THR:HB	3:P:323:SER:HA	0.67	0.44
3:B:471:LEU:HD23	3:B:471:LEU:N	2.32	0.44
3:B:549:LEU:C	3:B:549:LEU:CD1	2.86	0.44
3:D:169:MET:HG2	3:D:169:MET:O	2.15	0.44
3:D:262:ALA:CB	3:D:608:LEU:HD13	2.47	0.44
3:F:55:VAL:O	3:F:55:VAL:HG13	2.18	0.44
3:F:183:TRP:CE2	3:F:187:LEU:CD1	3.00	0.44
7:G:101:MYR:H21	3:H:191:LYS:NZ	2.25	0.44
3:H:442:THR:OG1	3:H:443:PHE:N	2.50	0.44
3:H:562:TYR:HA	3:H:565:LEU:HB2	1.99	0.44
2:I:24:SER:CB	3:J:176:ALA:N	2.80	0.44
3:J:208:TYR:CE2	3:J:212:ILE:HD11	2.53	0.44
3:J:298:PHE:CZ	3:N:355:GLY:CA	2.80	0.44
3:L:252:LEU:HB3	3:L:613:LEU:CD2	2.47	0.44
3:N:46:LYS:O	3:N:48:TRP:CD1	2.71	0.44
3:N:141:GLN:HE22	3:N:144:ASN:ND2	2.15	0.44
3:N:208:TYR:CE2	3:N:212:ILE:HD11	2.52	0.44
3:N:491:ALA:HB2	3:N:592:THR:CG2	2.48	0.44
3:P:183:TRP:CE2	3:P:187:LEU:CD1	3.00	0.44
3:P:298:PHE:HZ	3:T:416:ARG:HG3	1.81	0.44
2:Q:36:LEU:HD22	2:Q:41:LEU:HD11	1.98	0.44
3:R:442:THR:OG1	3:R:443:PHE:N	2.50	0.44
3:R:471:LEU:HD23	3:R:471:LEU:N	2.32	0.44
3:T:53:THR:HG22	3:T:54:SER:N	2.31	0.44
3:T:141:GLN:HE22	3:T:144:ASN:ND2	2.15	0.44
3:T:152:ARG:HA	3:T:155:ASP:OD2	2.18	0.44
3:T:257:SER:C	3:T:259:LEU:N	2.74	0.44
4:U:154:GLY:HA2	4:U:274:TRP:CE2	2.52	0.44
4:U:406:LEU:O	4:U:406:LEU:HD22	2.18	0.44
4:V:150:LEU:HB2	4:V:156:ILE:HD11	1.98	0.44
5:W:310:LEU:HD13	5:W:310:LEU:O	2.17	0.44
5:W:993:PRO:HA	5:W:1017:ILE:HD11	1.98	0.44
5:W:998:LEU:HD23	5:W:998:LEU:HA	1.72	0.44
5:W:1258:MET:HG2	5:W:1259:SER:H	1.82	0.44
6:X:206:LEU:HD12	6:X:243:ILE:HG12	1.99	0.44
6:X:225:PHE:C	6:X:227:GLN:N	2.74	0.44
6:X:243:ILE:HD11	6:X:1147:PHE:HD1	1.82	0.44

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
6:X:354:THR:O	6:X:355:LYS:C	2.60	0.44
6:X:531:CYS:SG	6:X:818:ILE:CD1	3.06	0.44
6:X:590:ALA:CB	6:X:639:LEU:HD21	2.47	0.44
6:X:883:ARG:HH11	6:X:883:ARG:CB	2.30	0.44
6:X:949:HIS:O	6:X:951:PRO:HD3	2.17	0.44
6:Y:191:PRO:HA	6:Y:915:LEU:CD1	2.47	0.44
6:Y:265:PRO:O	6:Y:267:THR:N	2.50	0.44
6:Y:737:ARG:HH21	6:Y:737:ARG:CG	2.20	0.44
6:Y:874:CYS:SG	6:Y:901:ILE:HD13	2.56	0.44
1:t:23:THR:HG21	1:t:31:GLU:OE1	2.17	0.44
2:A:24:SER:CB	3:B:176:ALA:N	2.80	0.44
3:B:44:THR:O	3:B:119:VAL:HG22	2.17	0.44
3:B:53:THR:HG22	3:B:54:SER:N	2.31	0.44
3:B:208:TYR:CE2	3:B:212:ILE:HD11	2.53	0.44
3:B:362:THR:O	3:B:364:PRO:HD3	2.18	0.44
3:B:533:ASP:CG	3:B:536:THR:HG23	2.42	0.44
3:D:455:SER:O	3:H:412:ARG:CA	2.66	0.44
3:D:567:ARG:NH1	3:D:567:ARG:HB2	2.33	0.44
3:F:46:LYS:O	3:F:48:TRP:CD1	2.71	0.44
3:F:152:ARG:HA	3:F:155:ASP:OD2	2.18	0.44
3:F:346:ASP:HB3	3:F:427:GLU:HA	1.99	0.44
3:F:416:ARG:HG3	3:H:298:PHE:HZ	1.82	0.44
3:F:600:LEU:HG	3:F:601:THR:N	2.30	0.44
3:J:345:ILE:HD12	3:J:468:PRO:HB3	1.99	0.44
3:J:455:SER:O	3:N:412:ARG:CA	2.66	0.44
3:J:559:PRO:O	3:J:560:PRO:C	2.58	0.44
3:J:644:LEU:HD23	3:J:644:LEU:N	2.32	0.44
3:L:346:ASP:HB3	3:L:427:GLU:HA	1.99	0.44
3:L:357:ILE:CG2	3:L:358:LEU:N	2.80	0.44
3:L:418:ASN:CG	3:N:337:ARG:NH2	2.74	0.44
3:L:491:ALA:HB2	3:L:592:THR:CG2	2.48	0.44
3:L:559:PRO:O	3:L:560:PRO:C	2.58	0.44
2:M:24:SER:CB	3:N:176:ALA:N	2.80	0.44
3:N:44:THR:O	3:N:119:VAL:HG22	2.17	0.44
3:P:46:LYS:O	3:P:48:TRP:CD1	2.71	0.44
3:P:51:VAL:HG12	5:W:379:VAL:HG23	2.00	0.44
3:P:53:THR:HG22	3:P:54:SER:N	2.31	0.44
3:P:341:MET:HG3	3:P:431:ILE:HD11	1.98	0.44
2:Q:24:SER:CB	3:R:176:ALA:N	2.80	0.44
3:R:533:ASP:OD1	3:R:536:THR:HG23	2.17	0.44
3:R:582:LEU:O	3:R:584:LEU:N	2.51	0.44

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:T:346:ASP:HB3	3:T:427:GLU:HA	1.99	0.44
3:T:360:TYR:HD1	3:T:360:TYR:C	2.24	0.44
4:U:312:VAL:O	4:U:313:ALA:C	2.60	0.44
4:U:406:LEU:HD13	4:U:406:LEU:C	2.42	0.44
5:W:63:PRO:C	5:W:65:LEU:N	2.76	0.44
5:W:244:THR:OG1	6:X:652:GLN:NE2	2.49	0.44
5:W:312:LEU:CD1	5:W:352:TYR:HB2	2.42	0.44
5:W:453:PRO:HA	5:W:454:PRO:HD2	1.87	0.44
5:W:566:LEU:O	5:W:602:MET:HA	2.18	0.44
5:W:955:VAL:CG2	5:W:964:VAL:CG1	2.95	0.44
5:W:1014:ASN:HD22	5:W:1213:SER:HB2	1.80	0.44
6:X:277:VAL:HG22	6:X:900:GLU:HB3	2.00	0.44
6:X:844:LEU:HD12	6:X:844:LEU:N	2.33	0.44
6:Y:206:LEU:O	6:Y:236:PRO:HB3	2.18	0.44
6:Y:231:MET:HE1	6:Y:347:ARG:CZ	2.48	0.44
6:Y:277:VAL:HG11	6:Y:295:SER:OG	2.17	0.44
6:Y:294:ARG:NE	6:Y:890:GLN:HB3	2.32	0.44
6:Y:406:LEU:HD23	6:Y:409:MET:HE3	2.00	0.44
1:l:60:ALA:H	3:L:398:SER:CA	2.04	0.44
1:d:74:GLN:C	1:d:74:GLN:CD	2.86	0.44
1:h:23:THR:HG21	3:H:323:SER:CA	2.46	0.44
1:r:60:ALA:O	1:r:61:ASN:HB3	2.18	0.44
3:B:170:LEU:O	3:B:171:THR:C	2.60	0.44
3:B:255:SER:HA	3:B:258:THR:HG22	1.99	0.44
3:D:57:THR:N	3:D:60:SER:HB3	2.26	0.44
3:F:362:THR:O	3:F:364:PRO:HD3	2.18	0.44
3:F:369:SER:CB	3:F:446:PRO:HD3	2.47	0.44
3:F:371:PRO:HG3	3:F:407:LEU:HD23	1.99	0.44
3:F:549:LEU:C	3:F:549:LEU:CD1	2.87	0.44
3:F:582:LEU:HD23	3:F:582:LEU:HA	1.69	0.44
3:F:582:LEU:O	3:F:584:LEU:N	2.51	0.44
3:H:252:LEU:HB3	3:H:613:LEU:CD2	2.47	0.44
3:H:262:ALA:CB	3:H:608:LEU:HD13	2.47	0.44
3:L:341:MET:HG3	3:L:431:ILE:HD11	1.98	0.44
3:L:547:HIS:CE1	3:L:560:PRO:HA	2.51	0.44
3:N:471:LEU:HD23	3:N:471:LEU:N	2.32	0.44
3:N:644:LEU:HD23	3:N:644:LEU:N	2.32	0.44
3:P:371:PRO:HG3	3:P:407:LEU:HD23	1.99	0.44
3:P:416:ARG:HG3	3:R:298:PHE:HZ	1.81	0.44
3:P:600:LEU:O	3:P:601:THR:C	2.60	0.44
3:T:46:LYS:O	3:T:48:TRP:CD1	2.71	0.44

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:T:55:VAL:O	3:T:55:VAL:HG13	2.18	0.44
3:T:178:SER:HB3	3:T:179:ALA:H	1.65	0.44
3:T:362:THR:O	3:T:364:PRO:HD3	2.18	0.44
4:U:156:ILE:HG12	4:U:268:ARG:HB3	2.00	0.44
4:U:178:THR:HB	6:X:376:LEU:CD2	2.47	0.44
4:U:291:THR:HG23	4:U:297:PRO:CD	2.47	0.44
4:U:304:THR:O	4:U:304:THR:HG23	2.18	0.44
5:W:440:THR:CG2	5:W:647:PRO:HB3	2.47	0.44
6:X:330:ILE:HA	6:X:369:GLY:HA2	2.00	0.44
6:X:504:THR:C	6:X:506:GLN:H	2.26	0.44
6:X:541:VAL:CG2	6:X:811:LEU:HG	2.45	0.44
6:Y:92:ASN:C	6:Y:94:THR:H	2.25	0.44
6:Y:359:PHE:CE1	6:Y:1173:VAL:HG21	2.53	0.44
6:Y:424:ALA:O	6:Y:425:ASN:HB2	2.18	0.44
6:Y:442:CYS:HB3	6:Y:444:TRP:CD2	2.52	0.44
6:Y:986:GLN:HB2	6:Y:1135:TYR:CE1	2.53	0.44
1:d:59:HIS:HB2	3:D:398:SER:HB2	2.00	0.44
1:r:59:HIS:O	1:r:59:HIS:CD2	2.70	0.44
1:t:24:LEU:HD22	1:t:35:PHE:HD1	1.80	0.44
3:B:79:ARG:NH1	3:B:79:ARG:CB	2.81	0.44
3:B:262:ALA:CB	3:B:608:LEU:HD13	2.47	0.44
3:B:345:ILE:HG23	3:B:475:THR:HG1	1.83	0.44
7:C:101:MYR:H52	7:C:101:MYR:C9	2.42	0.44
3:D:68:PHE:O	3:D:140:VAL:HB	2.18	0.44
3:D:235:LEU:HD13	3:D:235:LEU:HA	1.80	0.44
3:D:257:SER:C	3:D:259:LEU:N	2.74	0.44
3:F:193:ILE:O	3:F:194:LEU:C	2.61	0.44
3:F:330:GLN:O	3:F:331:LEU:HB2	2.17	0.44
3:F:385:VAL:HA	3:F:390:ASN:OD1	2.16	0.44
3:F:533:ASP:OD1	3:F:536:THR:HG23	2.18	0.44
3:F:599:ALA:O	3:F:600:LEU:C	2.60	0.44
3:H:170:LEU:O	3:H:171:THR:C	2.60	0.44
3:H:341:MET:HG3	3:H:431:ILE:HD11	1.99	0.44
3:J:79:ARG:NH1	3:J:79:ARG:CB	2.81	0.44
3:J:341:MET:CE	3:J:341:MET:CA	2.88	0.44
3:J:458:THR:HG23	3:J:458:THR:O	2.17	0.44
3:J:491:ALA:HB2	3:J:592:THR:CG2	2.48	0.44
2:K:9:ASN:HD21	3:L:206:GLU:HG3	1.83	0.44
3:L:369:SER:CB	3:L:446:PRO:HD3	2.47	0.44
3:L:401:GLY:O	3:L:402:ALA:CB	2.66	0.44
3:L:644:LEU:N	3:L:644:LEU:HD23	2.32	0.44

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:N:79:ARG:NH1	3:N:79:ARG:CB	2.81	0.44
3:N:183:TRP:CE2	3:N:187:LEU:CD1	3.00	0.44
3:N:256:SER:OG	3:N:257:SER:N	2.51	0.44
3:N:547:HIS:CE1	3:N:560:PRO:HA	2.52	0.44
2:O:18:ASN:O	2:O:19:SER:HB3	2.18	0.44
3:P:170:LEU:O	3:P:171:THR:C	2.60	0.44
3:P:256:SER:OG	3:P:257:SER:N	2.51	0.44
3:P:458:THR:O	3:P:458:THR:HG23	2.17	0.44
3:R:152:ARG:HA	3:R:155:ASP:OD2	2.18	0.44
3:R:199:LEU:HB3	3:R:207:PHE:CE2	2.52	0.44
3:R:255:SER:HA	3:R:258:THR:HG22	1.98	0.44
3:R:341:MET:HB2	3:R:437:ILE:CG2	2.35	0.44
2:S:24:SER:CB	3:T:176:ALA:N	2.81	0.44
3:T:208:TYR:CE2	3:T:212:ILE:HD11	2.53	0.44
3:T:256:SER:OG	3:T:257:SER:N	2.51	0.44
3:T:341:MET:HG3	3:T:431:ILE:HD11	1.98	0.44
3:T:567:ARG:NH1	3:T:567:ARG:HB2	2.33	0.44
4:U:148:TRP:CD1	4:U:155:PRO:HB3	2.53	0.44
4:U:148:TRP:HZ2	4:U:285:LEU:HD23	1.83	0.44
4:U:167:PRO:HB2	4:U:170:THR:OG1	2.17	0.44
4:V:223:LEU:HD11	4:V:411:PHE:CZ	2.52	0.44
5:W:234:THR:HG22	5:W:235:TYR:CD1	2.53	0.44
5:W:383:PHE:H	5:W:383:PHE:HD1	1.66	0.44
5:W:432:PHE:CB	5:W:481:ALA:HB2	2.48	0.44
6:X:411:GLN:HG2	6:X:445:PHE:CD2	2.52	0.44
6:X:453:LEU:O	6:X:457:MET:HB2	2.17	0.44
6:X:879:PHE:O	6:X:881:VAL:HG22	2.18	0.44
6:X:931:ILE:CD1	6:Y:689:VAL:HG11	2.47	0.44
6:X:1088:TYR:CE1	6:X:1132:ARG:HD2	2.51	0.44
6:Y:407:ARG:O	6:Y:408:SER:C	2.61	0.44
6:Y:647:ARG:HD3	6:Y:774:LEU:HD23	2.00	0.44
6:Y:651:PRO:O	6:Y:773:ARG:HD2	2.17	0.44
1:l:61:ASN:CB	3:L:397:VAL:HA	2.48	0.44
1:b:43:CYS:H	1:b:44:GLY:HA2	1.81	0.44
1:d:24:LEU:HD21	1:d:80:ARG:CD	2.47	0.44
1:j:8:GLN:HE22	3:L:303:ASP:CA	2.31	0.44
1:r:15:ARG:NH2	3:R:402:ALA:HB3	2.32	0.44
1:r:23:THR:CG2	1:r:25:TYR:CE2	3.00	0.44
1:r:45:ARG:CG	3:R:399:ALA:O	2.66	0.44
1:r:58:PRO:HB2	3:R:423:GLN:HG2	1.99	0.44
1:t:15:ARG:HB3	3:T:402:ALA:HB2	2.00	0.44

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:A:9:ASN:HD21	3:B:206:GLU:HG3	1.83	0.44
3:B:55:VAL:O	3:B:55:VAL:HG13	2.18	0.44
3:B:152:ARG:HA	3:B:155:ASP:OD2	2.18	0.44
3:B:360:TYR:HD1	3:B:360:TYR:C	2.24	0.44
3:B:458:THR:HG23	3:B:458:THR:O	2.17	0.44
3:B:533:ASP:OD1	3:B:536:THR:HG23	2.18	0.44
3:B:567:ARG:NH1	3:B:567:ARG:HB2	2.33	0.44
3:B:582:LEU:O	3:B:584:LEU:N	2.51	0.44
3:D:50:PRO:CD	3:D:61:LEU:HD23	2.47	0.44
3:D:588:ILE:HD13	3:D:588:ILE:HA	1.80	0.44
3:D:599:ALA:O	3:D:600:LEU:C	2.60	0.44
3:H:79:ARG:NH1	3:H:79:ARG:CB	2.81	0.44
3:H:533:ASP:OD1	3:H:536:THR:HG23	2.18	0.44
3:H:603:GLY:C	3:H:605:GLY:N	2.75	0.44
3:J:58:ILE:O	3:J:59:ASP:CB	2.63	0.44
3:J:337:ARG:NH2	3:N:418:ASN:CG	2.74	0.44
3:J:468:PRO:O	3:L:578:SER:OG	2.21	0.44
3:J:575:MET:HE3	3:N:469:VAL:HG21	0.69	0.44
3:L:141:GLN:HE22	3:L:144:ASN:ND2	2.15	0.44
3:L:412:ARG:CA	3:N:455:SER:O	2.66	0.44
3:N:55:VAL:O	3:N:55:VAL:HG13	2.18	0.44
3:N:458:THR:HG23	3:N:458:THR:O	2.17	0.44
3:N:533:ASP:OD1	3:N:536:THR:HG23	2.18	0.44
3:N:562:TYR:HA	3:N:565:LEU:HB2	1.99	0.44
2:O:9:ASN:HD21	3:P:206:GLU:HG3	1.83	0.44
2:O:23:THR:C	2:O:25:ASP:H	2.26	0.44
3:P:344:MET:HE2	3:P:502:LEU:CD1	2.37	0.44
3:R:401:GLY:O	3:R:402:ALA:CB	2.66	0.44
3:R:567:ARG:HB2	3:R:567:ARG:NH1	2.33	0.44
3:T:79:ARG:NH1	3:T:79:ARG:CB	2.81	0.44
3:T:341:MET:CE	3:T:341:MET:CA	2.88	0.44
3:T:369:SER:CB	3:T:446:PRO:HD3	2.47	0.44
4:U:22:ASP:O	4:U:25:ASP:HB2	2.18	0.44
4:U:228:THR:HG21	4:U:251:ASN:HB3	2.00	0.44
4:U:231:GLY:HA3	4:U:250:TRP:NE1	2.33	0.44
4:V:89:GLY:C	4:V:103:VAL:HG23	2.43	0.44
4:V:312:VAL:HG12	4:V:313:ALA:H	1.83	0.44
5:W:16:SER:O	5:W:285:LEU:HD11	2.17	0.44
5:W:148:ILE:CG2	5:W:149:THR:N	2.81	0.44
5:W:270:LEU:HA	5:W:270:LEU:HD23	1.82	0.44
5:W:424:SER:O	5:W:426:LEU:N	2.47	0.44

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
5:W:1289:LEU:HD22	5:W:1289:LEU:H	1.83	0.44
6:X:266:PRO:HB2	6:X:1085:LEU:HD12	1.99	0.44
6:X:314:SER:N	6:X:1197:TYR:O	2.43	0.44
6:X:354:THR:CG2	6:X:1152:ALA:HA	2.47	0.44
6:X:446:ALA:HB2	6:X:858:VAL:HG23	2.00	0.44
6:X:459:LEU:HA	6:X:462:ILE:HG22	2.00	0.44
6:X:517:LEU:O	6:X:519:GLN:N	2.49	0.44
6:X:517:LEU:HD12	6:X:565:LYS:HG3	2.00	0.44
6:X:718:THR:CA	6:X:726:ARG:HH11	2.29	0.44
6:X:844:LEU:H	6:X:844:LEU:HD12	1.83	0.44
6:X:996:MET:HB3	6:X:1143:ASN:HB2	2.00	0.44
6:Y:128:THR:HG22	6:Y:130:SER:N	2.26	0.44
6:Y:172:PRO:CD	6:Y:847:SER:HA	2.46	0.44
6:Y:370:ARG:NE	6:Y:385:VAL:HG23	2.32	0.44
6:Y:612:ALA:O	6:Y:613:SER:C	2.61	0.44
6:Y:1023:ARG:NH2	6:Y:1030:GLN:NE2	2.66	0.44
1:l:22:LEU:HD23	1:l:22:LEU:H	1.82	0.44
1:b:4:HIS:CE1	1:b:54:THR:CG2	2.92	0.44
1:f:5:MET:CA	1:f:5:MET:CE	2.85	0.44
1:h:6:ILE:CD1	3:D:586:VAL:CB	2.96	0.44
1:h:45:ARG:HD2	3:H:400:ALA:CA	2.48	0.44
1:r:22:LEU:HD22	1:r:22:LEU:H	1.79	0.44
3:B:343:ALA:CB	3:B:478:GLY:O	2.62	0.44
3:B:371:PRO:HG3	3:B:407:LEU:HD23	1.99	0.44
3:B:387:THR:O	3:B:390:ASN:HB2	2.18	0.44
3:B:559:PRO:O	3:B:560:PRO:C	2.58	0.44
3:D:330:GLN:O	3:D:331:LEU:HB2	2.17	0.44
3:D:491:ALA:HB2	3:D:592:THR:CG2	2.48	0.44
3:D:533:ASP:CG	3:D:536:THR:HG23	2.42	0.44
3:F:491:ALA:HB2	3:F:592:THR:CG2	2.47	0.44
3:F:644:LEU:HD23	3:F:644:LEU:N	2.32	0.44
3:H:167:MET:HA	3:H:167:MET:CE	2.31	0.44
3:H:208:TYR:CE2	3:H:212:ILE:HD11	2.52	0.44
3:H:567:ARG:NH1	3:H:567:ARG:HB2	2.33	0.44
3:J:152:ARG:HA	3:J:155:ASP:OD2	2.18	0.44
3:J:533:ASP:CG	3:J:536:THR:HG23	2.42	0.44
3:J:564:LEU:CD2	3:J:568:ARG:NH2	2.75	0.44
2:K:23:THR:C	2:K:25:ASP:N	2.75	0.44
3:L:107:VAL:HG21	3:N:128:ASP:OD2	2.18	0.44
3:L:152:ARG:HA	3:L:155:ASP:OD2	2.18	0.44
3:L:533:ASP:OD1	3:L:536:THR:HG23	2.18	0.44

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:L:540:VAL:HG21	3:L:566:TRP:HA	2.00	0.44
3:N:53:THR:HG22	3:N:54:SER:N	2.32	0.44
3:N:387:THR:O	3:N:390:ASN:HB2	2.18	0.44
3:N:431:ILE:HD11	3:N:437:ILE:CG2	2.45	0.44
3:N:533:ASP:CG	3:N:536:THR:HG23	2.42	0.44
3:N:567:ARG:HB2	3:N:567:ARG:NH1	2.33	0.44
3:P:193:ILE:O	3:P:194:LEU:C	2.61	0.44
3:P:582:LEU:O	3:P:584:LEU:N	2.51	0.44
3:T:92:TRP:CZ2	3:T:160:GLN:CB	3.01	0.44
3:T:255:SER:HA	3:T:258:THR:HG22	1.99	0.44
4:U:157:TYR:HB2	4:U:287:SER:HB2	2.00	0.44
5:W:15:THR:HG23	5:W:317:GLY:CA	2.47	0.44
5:W:353:ARG:HB2	5:W:356:GLN:O	2.18	0.44
5:W:773:ARG:HG2	5:W:867:PHE:HD2	1.83	0.44
5:W:773:ARG:HD3	5:W:872:THR:HG21	2.00	0.44
5:W:1264:PRO:O	5:W:1265:ALA:C	2.61	0.44
6:X:210:ILE:O	6:X:239:ALA:N	2.51	0.44
6:X:394:LEU:HD13	6:X:967:THR:HG21	1.99	0.44
6:X:513:GLU:O	6:X:514:ILE:C	2.61	0.44
6:X:700:TRP:O	6:X:704:THR:CG2	2.60	0.44
6:X:789:ASP:C	6:X:791:ALA:H	2.25	0.44
6:X:813:VAL:CG1	6:X:818:ILE:HD11	2.46	0.44
6:X:861:ARG:HE	6:X:861:ARG:HB3	1.66	0.44
6:X:987:THR:CG2	6:X:989:THR:HG23	2.40	0.44
6:Y:43:ALA:C	6:Y:45:ASP:H	2.24	0.44
6:Y:61:ASN:C	6:Y:63:VAL:H	2.25	0.44
6:Y:260:GLU:HA	6:Y:307:SER:HB2	2.00	0.44
6:Y:686:ARG:CG	6:Y:686:ARG:HH11	2.30	0.44
6:Y:718:THR:O	6:Y:719:ALA:C	2.61	0.44
6:Y:831:LEU:O	6:Y:833:PRO:HD3	2.17	0.44
6:Y:862:GLU:HA	6:Y:862:GLU:OE1	2.17	0.44
6:Y:910:LYS:O	6:Y:914:ASP:N	2.50	0.44
1:l:6:ILE:HG21	3:N:494:THR:HG22	2.00	0.44
1:f:44:GLY:O	3:F:399:ALA:C	2.61	0.44
1:d:24:LEU:CD2	1:d:80:ARG:CD	2.96	0.44
1:n:24:LEU:HD21	1:n:80:ARG:CD	2.31	0.44
1:p:17:ALA:O	1:p:82:LEU:CD1	2.66	0.44
2:C:24:SER:CB	3:D:176:ALA:N	2.81	0.44
3:D:256:SER:OG	3:D:257:SER:N	2.51	0.44
3:D:298:PHE:HZ	3:H:416:ARG:HG3	1.81	0.44
3:D:346:ASP:HB3	3:D:427:GLU:HA	1.99	0.44

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:D:533:ASP:OD1	3:D:536:THR:HG23	2.17	0.44
3:D:582:LEU:O	3:D:584:LEU:N	2.51	0.44
3:F:262:ALA:CB	3:F:608:LEU:HD13	2.47	0.44
3:F:387:THR:O	3:F:390:ASN:HB2	2.18	0.44
3:F:456:ASN:ND2	3:F:458:THR:HG22	2.33	0.44
2:G:23:THR:C	2:G:25:ASP:H	2.26	0.44
3:H:46:LYS:O	3:H:48:TRP:CD1	2.71	0.44
3:H:152:ARG:HA	3:H:155:ASP:OD2	2.18	0.44
3:H:369:SER:CB	3:H:446:PRO:HD3	2.47	0.44
3:H:371:PRO:HG3	3:H:407:LEU:HD23	1.99	0.44
2:I:9:ASN:HD21	3:J:206:GLU:HG3	1.83	0.44
3:J:141:GLN:HE22	3:J:144:ASN:ND2	2.15	0.44
3:J:330:GLN:HB2	3:J:457:PRO:HG3	1.99	0.44
3:J:582:LEU:O	3:J:584:LEU:N	2.51	0.44
3:J:609:ARG:CD	2:K:29:THR:HG21	2.47	0.44
2:K:23:THR:C	2:K:25:ASP:H	2.26	0.44
3:L:67:ARG:HH11	3:L:67:ARG:CG	2.29	0.44
3:L:208:TYR:CE2	3:L:212:ILE:HD11	2.53	0.44
3:L:330:GLN:HB2	3:L:457:PRO:HG3	1.99	0.44
3:L:387:THR:O	3:L:390:ASN:HB2	2.18	0.44
3:L:603:GLY:C	3:L:605:GLY:N	2.75	0.44
2:M:23:THR:C	2:M:25:ASP:H	2.26	0.44
7:M:101:MYR:C2	3:N:191:LYS:HZ1	2.27	0.44
3:N:152:ARG:HA	3:N:155:ASP:OD2	2.18	0.44
3:N:252:LEU:HB3	3:N:613:LEU:CD2	2.47	0.44
3:N:330:GLN:HB2	3:N:457:PRO:HG3	1.99	0.44
3:N:346:ASP:HB3	3:N:427:GLU:HA	1.99	0.44
3:N:582:LEU:O	3:N:584:LEU:N	2.51	0.44
2:O:29:THR:HG21	3:T:609:ARG:CD	2.47	0.44
3:P:141:GLN:HE22	3:P:144:ASN:ND2	2.15	0.44
3:P:455:SER:O	3:T:412:ARG:CA	2.66	0.44
2:Q:9:ASN:HD21	3:R:206:GLU:HG3	1.83	0.44
3:R:55:VAL:O	3:R:55:VAL:HG13	2.18	0.44
3:R:183:TRP:NE1	3:R:187:LEU:CD1	2.81	0.44
3:R:199:LEU:CD1	3:R:203:TYR:CE2	3.01	0.44
3:R:208:TYR:CE2	3:R:212:ILE:HD11	2.53	0.44
3:R:235:LEU:HD13	3:R:235:LEU:HA	1.80	0.44
3:R:369:SER:CB	3:R:446:PRO:HD3	2.47	0.44
3:T:67:ARG:HH11	3:T:67:ARG:CG	2.29	0.44
3:T:193:ILE:O	3:T:194:LEU:C	2.61	0.44
3:T:357:ILE:CG2	3:T:358:LEU:N	2.80	0.44

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:T:387:THR:O	3:T:390:ASN:HB2	2.18	0.44
3:T:588:ILE:HA	3:T:588:ILE:HD13	1.80	0.44
3:T:600:LEU:HG	3:T:601:THR:N	2.30	0.44
4:U:42:SER:O	4:U:43:HIS:HD2	2.01	0.44
4:U:63:SER:O	4:U:66:ILE:HG22	2.18	0.44
4:U:124:ASN:HA	5:W:353:ARG:NH1	2.33	0.44
4:V:191:ARG:HH12	6:Y:438:PRO:HG2	1.80	0.44
5:W:10:VAL:HB	5:W:388:TYR:CE2	2.52	0.44
5:W:632:PHE:CE2	5:W:670:SER:HB3	2.52	0.44
5:W:700:VAL:O	5:W:700:VAL:HG13	2.16	0.44
5:W:735:LEU:CD2	5:W:759:ILE:HB	2.48	0.44
5:W:765:PHE:CD1	5:W:766:LEU:N	2.86	0.44
5:W:1267:LEU:HA	5:W:1268:PRO:HD3	1.89	0.44
6:X:241:ASN:ND2	6:X:1148:SER:OG	2.37	0.44
6:X:244:VAL:HG22	6:X:245:CYS:N	2.33	0.44
6:X:297:VAL:O	6:X:297:VAL:HG12	2.17	0.44
6:X:514:ILE:CG1	6:X:515:LEU:N	2.81	0.44
6:X:983:TYR:HE1	6:X:1075:LEU:CG	2.30	0.44
6:X:1059:PHE:CD2	6:X:1063:VAL:HG21	2.53	0.44
6:Y:257:ARG:O	6:Y:259:LEU:N	2.50	0.44
6:Y:412:ASP:OD2	6:Y:414:THR:HG22	2.16	0.44
6:Y:621:TRP:HB2	6:Y:776:GLN:NE2	2.32	0.44
6:Y:660:SER:HB3	6:Y:663:GLN:CG	2.47	0.44
6:Y:874:CYS:HB3	6:Y:893:VAL:CG2	2.46	0.44
1:l:29:ARG:NH2	1:l:65:ILE:O	2.49	0.43
1:j:8:GLN:HE22	3:L:303:ASP:CG	2.23	0.43
1:p:61:ASN:CB	3:P:397:VAL:HA	2.22	0.43
1:p:73:ARG:CZ	1:p:77:GLU:OE2	2.65	0.43
2:C:9:ASN:HD21	3:D:206:GLU:HG3	1.83	0.43
3:D:44:THR:O	3:D:119:VAL:HG22	2.17	0.43
3:D:78:MET:HE2	3:H:238:VAL:HG21	1.83	0.43
3:D:170:LEU:O	3:D:171:THR:C	2.60	0.43
2:E:23:THR:C	2:E:25:ASP:H	2.26	0.43
3:F:540:VAL:HG21	3:F:566:TRP:HA	2.00	0.43
2:G:24:SER:CB	3:H:176:ALA:N	2.81	0.43
3:H:582:LEU:O	3:H:584:LEU:N	2.51	0.43
3:J:92:TRP:CZ2	3:J:160:GLN:CB	3.01	0.43
3:J:183:TRP:CE2	3:J:187:LEU:CD1	3.01	0.43
3:J:257:SER:C	3:J:259:LEU:N	2.74	0.43
3:J:387:THR:O	3:J:390:ASN:HB2	2.18	0.43
3:J:456:ASN:ND2	3:J:458:THR:HG22	2.33	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:L:46:LYS:O	3:L:48:TRP:CD1	2.71	0.43
3:L:55:VAL:O	3:L:55:VAL:HG13	2.18	0.43
3:L:345:ILE:HD12	3:L:468:PRO:HB3	1.99	0.43
3:L:371:PRO:HG3	3:L:407:LEU:HD23	1.99	0.43
3:N:330:GLN:O	3:N:331:LEU:HB2	2.17	0.43
3:N:362:THR:O	3:N:364:PRO:HD3	2.18	0.43
3:N:369:SER:CB	3:N:446:PRO:HD3	2.47	0.43
3:N:540:VAL:HG21	3:N:566:TRP:HA	2.00	0.43
3:P:152:ARG:HA	3:P:155:ASP:OD2	2.18	0.43
3:P:252:LEU:HB3	3:P:613:LEU:CD2	2.47	0.43
3:P:456:ASN:ND2	3:P:458:THR:HG22	2.33	0.43
3:R:46:LYS:O	3:R:48:TRP:CD1	2.71	0.43
3:R:92:TRP:CZ2	3:R:160:GLN:CB	3.01	0.43
3:R:256:SER:OG	3:R:257:SER:N	2.51	0.43
3:R:588:ILE:HG22	3:R:591:LEU:HB2	2.00	0.43
3:T:297:ALA:CB	3:T:460:MET:HB2	2.33	0.43
4:U:13:PHE:HD2	4:U:17:PRO:O	2.01	0.43
4:U:91:THR:O	4:U:92:TRP:C	2.60	0.43
4:U:125:TYR:CE1	4:U:127:THR:HG22	2.53	0.43
4:U:198:GLN:OE1	4:U:198:GLN:CA	2.65	0.43
4:U:241:LEU:O	6:X:1179:VAL:HG11	2.17	0.43
4:U:379:ILE:C	4:U:381:ALA:N	2.75	0.43
5:W:148:ILE:O	5:W:152:THR:HG22	2.18	0.43
5:W:315:VAL:O	5:W:315:VAL:HG22	2.18	0.43
5:W:635:TYR:HD1	5:W:638:HIS:CE1	2.36	0.43
5:W:1154:ARG:NE	5:W:1189:ASN:HB2	2.33	0.43
6:X:277:VAL:O	6:X:278:LEU:C	2.60	0.43
6:X:839:LYS:HZ2	6:X:839:LYS:CB	2.30	0.43
6:X:940:GLN:HE21	6:X:940:GLN:HB2	1.53	0.43
6:X:1064:TRP:HE1	6:X:1101:GLU:HG2	1.83	0.43
6:X:1064:TRP:HE1	6:X:1101:GLU:CG	2.31	0.43
6:X:1149:THR:HB	6:X:1162:ALA:CB	2.47	0.43
6:Y:202:THR:HG22	6:Y:392:ASP:OD1	2.18	0.43
6:Y:314:SER:OG	6:Y:1198:ASN:ND2	2.51	0.43
6:Y:452:ARG:C	6:Y:950:VAL:HG21	2.43	0.43
6:Y:574:PRO:HB2	6:Y:679:TRP:CZ2	2.53	0.43
6:Y:618:PRO:HB2	6:Y:941:TYR:CB	2.48	0.43
6:Y:705:ILE:HG21	6:Y:733:MET:O	2.17	0.43
6:Y:781:THR:O	6:Y:807:PHE:HB3	2.18	0.43
6:Y:790:PRO:O	6:Y:791:ALA:C	2.60	0.43
6:Y:1001:ALA:O	6:Y:1002:PRO:C	2.61	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
6:Y:1072:THR:CG2	6:Y:1073:ARG:N	2.81	0.43
6:Y:1149:THR:HG23	6:Y:1162:ALA:HB3	2.00	0.43
1:b:58:PRO:HG2	3:B:423:GLN:HG3	2.00	0.43
1:n:43:CYS:C	3:N:322:TYR:HH	2.17	0.43
1:t:3:LEU:HD22	1:t:3:LEU:HA	1.83	0.43
7:A:101:MYR:H42	3:B:187:LEU:HD22	2.00	0.43
3:B:46:LYS:O	3:B:48:TRP:CD1	2.71	0.43
3:B:193:ILE:O	3:B:194:LEU:C	2.61	0.43
3:B:322:TYR:O	3:B:323:SER:HB3	2.17	0.43
3:B:330:GLN:HB2	3:B:457:PRO:HG3	1.99	0.43
3:B:456:ASN:ND2	3:B:458:THR:HG22	2.33	0.43
2:C:23:THR:C	2:C:25:ASP:H	2.26	0.43
7:C:101:MYR:H42	3:D:187:LEU:HD22	2.00	0.43
3:D:79:ARG:NH1	3:D:79:ARG:CB	2.81	0.43
3:D:96:PHE:CE1	3:D:159:TRP:HH2	2.37	0.43
3:D:208:TYR:CE2	3:D:212:ILE:HD11	2.53	0.43
3:D:298:PHE:CE1	3:H:355:GLY:HA2	2.47	0.43
3:D:345:ILE:HG23	3:D:475:THR:HG1	1.82	0.43
3:D:369:SER:CB	3:D:446:PRO:HD3	2.47	0.43
3:D:371:PRO:HG3	3:D:407:LEU:HD23	1.99	0.43
3:F:183:TRP:NE1	3:F:187:LEU:CD1	2.81	0.43
3:H:397:VAL:O	3:H:398:SER:HB2	2.18	0.43
3:J:357:ILE:CG2	3:J:358:LEU:N	2.80	0.43
2:K:24:SER:CB	3:L:176:ALA:N	2.80	0.43
3:L:92:TRP:CZ2	3:L:160:GLN:CB	3.01	0.43
3:L:193:ILE:O	3:L:194:LEU:C	2.61	0.43
3:L:390:ASN:HD22	3:L:390:ASN:HA	1.60	0.43
3:L:533:ASP:CG	3:L:536:THR:HG23	2.42	0.43
3:L:562:TYR:HA	3:L:565:LEU:HB2	1.99	0.43
2:M:18:ASN:O	2:M:19:SER:HB3	2.18	0.43
2:M:23:THR:C	2:M:25:ASP:N	2.75	0.43
3:N:183:TRP:NE1	3:N:187:LEU:CD1	2.81	0.43
3:P:79:ARG:NH1	3:P:79:ARG:CB	2.81	0.43
3:P:101:THR:HG23	3:P:102:GLY:N	2.33	0.43
3:P:183:TRP:NE1	3:P:187:LEU:CD1	2.81	0.43
3:P:362:THR:O	3:P:364:PRO:HD3	2.18	0.43
3:P:530:ARG:HH21	3:T:472:ARG:HH22	1.66	0.43
3:P:582:LEU:HD23	3:P:585:GLN:HE22	1.83	0.43
3:P:603:GLY:C	3:P:605:GLY:N	2.75	0.43
3:R:79:ARG:NH1	3:R:79:ARG:CB	2.81	0.43
3:R:183:TRP:CE2	3:R:187:LEU:CD1	3.00	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:R:336:THR:CG2	3:R:456:ASN:ND2	2.79	0.43
3:R:345:ILE:HG23	3:R:475:THR:HG1	1.82	0.43
3:R:346:ASP:HB3	3:R:427:GLU:HA	1.99	0.43
3:T:183:TRP:NE1	3:T:187:LEU:CD1	2.81	0.43
3:T:345:ILE:HD12	3:T:468:PRO:HB3	1.99	0.43
3:T:397:VAL:O	3:T:398:SER:HB2	2.18	0.43
4:U:10:THR:HG23	4:U:73:GLY:CA	2.47	0.43
4:U:249:LYS:HD2	6:X:376:LEU:HG	1.99	0.43
4:V:31:GLY:H	6:X:1025:ASN:CB	2.31	0.43
4:V:131:ASN:O	4:V:135:MET:HG2	2.18	0.43
5:W:286:ASP:OD1	5:W:286:ASP:C	2.61	0.43
5:W:461:PRO:O	5:W:464:TYR:HB2	2.17	0.43
5:W:579:TYR:HA	5:W:617:VAL:HB	1.99	0.43
5:W:1040:VAL:C	5:W:1042:THR:H	2.27	0.43
5:W:1071:ILE:HD11	5:W:1100:PHE:CE1	2.54	0.43
6:X:200:LEU:HD13	6:X:343:ALA:CB	2.48	0.43
6:X:302:LEU:C	6:X:302:LEU:CD2	2.91	0.43
6:X:388:MET:HE3	6:X:388:MET:HB3	1.70	0.43
6:X:557:ILE:HB	6:X:592:LEU:HD21	2.00	0.43
6:X:902:MET:HE3	6:X:929:PRO:CG	2.45	0.43
6:X:1031:LEU:N	6:X:1039:ARG:O	2.48	0.43
6:Y:272:HIS:ND1	6:Y:273:ALA:N	2.66	0.43
6:Y:350:LEU:C	6:Y:352:THR:H	2.25	0.43
6:Y:358:GLN:O	6:Y:361:SER:HB3	2.18	0.43
6:Y:421:ASN:OD1	6:Y:429:ASN:HA	2.17	0.43
6:Y:606:MET:CE	6:Y:610:THR:HG21	2.44	0.43
6:Y:607:HIS:C	6:Y:609:THR:N	2.76	0.43
6:Y:828:GLU:O	6:Y:829:THR:C	2.61	0.43
6:Y:996:MET:HB2	6:Y:1143:ASN:HB3	2.01	0.43
6:Y:1095:ASP:OD1	6:Y:1098:PRO:HD3	2.18	0.43
1:t:44:GLY:O	3:T:399:ALA:CB	2.67	0.43
1:t:45:ARG:CB	3:T:399:ALA:O	2.64	0.43
2:A:23:THR:C	2:A:25:ASP:N	2.75	0.43
3:B:183:TRP:CE2	3:B:187:LEU:CD1	3.01	0.43
3:D:107:VAL:HG21	3:F:128:ASP:OD2	2.18	0.43
3:D:345:ILE:HD12	3:D:468:PRO:HB3	1.99	0.43
3:D:355:GLY:CA	3:F:298:PHE:CZ	2.80	0.43
3:D:431:ILE:O	3:D:431:ILE:HG23	2.10	0.43
3:F:79:ARG:NH1	3:F:79:ARG:CB	2.81	0.43
3:J:345:ILE:O	3:J:347:SER:N	2.52	0.43
3:L:79:ARG:NH1	3:L:79:ARG:CB	2.81	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:L:95:LEU:HD12	3:L:95:LEU:HA	1.71	0.43
3:L:183:TRP:NE1	3:L:187:LEU:CD1	2.81	0.43
3:N:92:TRP:CZ2	3:N:160:GLN:CB	3.01	0.43
3:N:336:THR:CG2	3:N:456:ASN:ND2	2.79	0.43
3:N:345:ILE:O	3:N:347:SER:N	2.52	0.43
3:N:357:ILE:CG2	3:N:358:LEU:N	2.80	0.43
3:N:401:GLY:O	3:N:402:ALA:CB	2.66	0.43
3:N:600:LEU:HG	3:N:601:THR:N	2.31	0.43
3:P:58:ILE:HG22	3:P:58:ILE:O	2.19	0.43
3:P:257:SER:C	3:P:259:LEU:N	2.74	0.43
3:P:412:ARG:CA	3:R:455:SER:O	2.66	0.43
3:P:418:ASN:CG	3:R:337:ARG:NH2	2.74	0.43
3:T:170:LEU:O	3:T:171:THR:C	2.60	0.43
3:T:203:TYR:HB3	3:T:206:GLU:CB	2.39	0.43
4:U:192:PHE:CZ	4:U:204:MET:HB3	2.53	0.43
4:V:204:MET:HA	4:V:207:THR:HG22	1.98	0.43
5:W:445:ILE:HB	5:W:642:LEU:HD22	2.00	0.43
5:W:512:THR:O	5:W:513:SER:HB2	2.18	0.43
5:W:577:LEU:C	5:W:577:LEU:CD2	2.91	0.43
5:W:722:ASP:O	5:W:723:ALA:HB3	2.16	0.43
5:W:765:PHE:CZ	5:W:999:PRO:O	2.71	0.43
6:X:263:LEU:HB3	6:X:976:ASN:ND2	2.34	0.43
6:X:463:SER:C	6:X:465:ASN:N	2.76	0.43
6:X:592:LEU:HD23	6:X:592:LEU:C	2.44	0.43
6:X:878:GLY:O	6:X:879:PHE:C	2.61	0.43
6:X:907:ASP:O	6:X:908:ALA:C	2.61	0.43
6:X:1072:THR:O	6:X:1074:VAL:HG12	2.18	0.43
6:Y:31:ASN:HB2	6:Y:47:ASN:OD1	2.17	0.43
6:Y:264:PRO:HB2	6:Y:268:ASP:HB2	1.99	0.43
6:Y:416:ILE:HD12	6:Y:851:PHE:CZ	2.53	0.43
6:Y:672:SER:O	6:Y:674:ARG:HB3	2.18	0.43
6:Y:741:THR:OG1	6:Y:743:THR:HG23	2.17	0.43
1:b:4:HIS:NE2	1:b:54:THR:HA	2.33	0.43
3:B:92:TRP:CZ2	3:B:160:GLN:CB	3.01	0.43
3:B:183:TRP:NE1	3:B:187:LEU:CD1	2.81	0.43
3:B:199:LEU:HB3	3:B:207:PHE:CE2	2.52	0.43
3:B:199:LEU:CD1	3:B:203:TYR:CE2	3.01	0.43
3:B:644:LEU:HD23	3:B:644:LEU:N	2.32	0.43
3:D:183:TRP:CE2	3:D:187:LEU:CD1	3.01	0.43
3:D:281:PRO:CD	3:F:622:GLU:OE1	2.62	0.43
3:D:310:PHE:CD1	3:D:382:ILE:HG13	2.54	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:E:9:ASN:HD21	3:F:206:GLU:HG3	1.83	0.43
3:F:562:TYR:HA	3:F:565:LEU:HB2	1.99	0.43
3:H:154:ARG:NH1	3:H:154:ARG:CB	2.68	0.43
3:H:199:LEU:CD1	3:H:203:TYR:CE2	3.01	0.43
3:H:345:ILE:O	3:H:347:SER:N	2.52	0.43
3:J:49:ARG:CZ	4:V:405:ALA:CB	2.97	0.43
3:J:183:TRP:NE1	3:J:187:LEU:CD1	2.81	0.43
3:L:135:VAL:O	3:L:135:VAL:HG13	2.19	0.43
3:L:427:GLU:HB2	3:L:430:THR:CG2	2.49	0.43
3:L:588:ILE:HG22	3:L:591:LEU:HB2	2.00	0.43
3:N:101:THR:HG23	3:N:102:GLY:N	2.34	0.43
3:N:136:PHE:CD1	3:N:136:PHE:N	2.87	0.43
3:N:371:PRO:HG3	3:N:407:LEU:HD23	1.99	0.43
3:P:262:ALA:CB	3:P:608:LEU:HD13	2.47	0.43
3:P:491:ALA:HB2	3:P:592:THR:CG2	2.48	0.43
3:P:567:ARG:NH1	3:P:567:ARG:HB2	2.33	0.43
3:R:491:ALA:HB2	3:R:592:THR:CG2	2.48	0.43
2:S:18:ASN:O	2:S:19:SER:HB3	2.18	0.43
3:T:540:VAL:HG21	3:T:566:TRP:HA	2.00	0.43
4:U:4:ARG:HH12	5:W:217:LEU:CG	2.32	0.43
4:U:54:TRP:O	4:U:59:PRO:HA	2.17	0.43
4:U:83:GLN:H	4:U:83:GLN:HG3	1.52	0.43
4:U:92:TRP:HA	4:U:99:SER:O	2.18	0.43
4:U:265:SER:HA	4:U:323:VAL:HG12	2.01	0.43
5:W:28:ASP:HB3	5:W:109:ALA:HB1	2.00	0.43
5:W:219:ASN:HD21	6:X:674:ARG:HH12	1.66	0.43
5:W:350:PHE:CE1	5:W:378:PRO:HD3	2.52	0.43
5:W:577:LEU:HD23	5:W:579:TYR:N	2.30	0.43
5:W:779:MET:HE2	5:W:779:MET:CA	2.47	0.43
5:W:1012:ASP:O	5:W:1013:ILE:C	2.61	0.43
6:X:234:PRO:HG2	6:X:360:LEU:HD11	2.00	0.43
6:X:263:LEU:N	6:X:263:LEU:CD2	2.81	0.43
6:X:265:PRO:O	6:X:268:ASP:HB2	2.18	0.43
6:X:393:ALA:CB	6:X:1184:LEU:HD13	2.48	0.43
6:X:401:THR:HG21	6:X:960:ALA:HB2	1.99	0.43
6:X:413:PRO:HA	6:X:416:ILE:CG1	2.49	0.43
6:X:603:VAL:HB	6:X:604:PRO:HD2	1.99	0.43
6:X:930:ARG:NH2	6:Y:692:THR:HG21	2.32	0.43
6:X:1072:THR:HG23	6:X:1072:THR:O	2.17	0.43
6:Y:462:ILE:O	6:Y:465:ASN:HB3	2.17	0.43
6:Y:742:MET:HG3	6:Y:742:MET:O	2.18	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
6:Y:1064:TRP:HB3	6:Y:1065:PRO:HD3	1.99	0.43
6:Y:1089:ASP:C	6:Y:1091:GLN:H	2.26	0.43
6:Y:1090:PRO:HB3	6:Y:1158:PHE:HE2	1.83	0.43
1:f:60:ALA:N	3:F:398:SER:HA	2.32	0.43
1:j:61:ASN:CB	3:J:397:VAL:HA	2.47	0.43
1:p:62:VAL:O	1:p:64:THR:CA	2.60	0.43
1:t:29:ARG:NH2	1:t:65:ILE:O	2.49	0.43
3:B:369:SER:CB	3:B:446:PRO:HD3	2.47	0.43
3:D:106:ILE:CD1	3:D:152:ARG:HG3	2.45	0.43
3:D:128:ASP:OD2	3:H:107:VAL:HG21	2.18	0.43
3:D:456:ASN:ND2	3:D:458:THR:HG22	2.33	0.43
3:D:582:LEU:HD23	3:D:585:GLN:HE22	1.83	0.43
3:F:257:SER:C	3:F:259:LEU:N	2.74	0.43
3:H:341:MET:HB2	3:H:437:ILE:CG2	2.35	0.43
2:I:23:THR:C	2:I:25:ASP:N	2.75	0.43
3:J:193:ILE:O	3:J:194:LEU:C	2.61	0.43
3:J:292:ARG:HG2	3:J:292:ARG:NH1	2.32	0.43
3:J:343:ALA:CB	3:J:478:GLY:O	2.62	0.43
3:J:540:VAL:HG21	3:J:566:TRP:HA	2.00	0.43
3:J:588:ILE:HG22	3:J:591:LEU:HB2	2.00	0.43
3:L:336:THR:CG2	3:L:456:ASN:ND2	2.79	0.43
3:L:442:THR:OG1	3:L:443:PHE:N	2.50	0.43
3:L:582:LEU:O	3:L:584:LEU:N	2.51	0.43
2:M:13:ILE:CD1	7:M:101:MYR:H111	2.46	0.43
3:N:193:ILE:O	3:N:194:LEU:C	2.61	0.43
3:N:343:ALA:CB	3:N:478:GLY:O	2.62	0.43
3:N:427:GLU:HB2	3:N:430:THR:CG2	2.49	0.43
3:N:603:GLY:C	3:N:605:GLY:N	2.75	0.43
3:P:96:PHE:CE1	3:P:159:TRP:HH2	2.36	0.43
3:P:107:VAL:HG21	3:R:128:ASP:OD2	2.18	0.43
3:P:237:GLU:CG	3:R:46:LYS:HZ1	2.31	0.43
3:R:565:LEU:HD23	3:R:565:LEU:N	2.34	0.43
2:S:23:THR:C	2:S:25:ASP:H	2.26	0.43
3:T:310:PHE:CD1	3:T:382:ILE:HG13	2.54	0.43
3:T:533:ASP:CG	3:T:536:THR:HG23	2.42	0.43
4:U:161:VAL:HG23	4:U:263:GLN:HB2	1.99	0.43
4:V:40:ARG:HH21	4:V:65:ASN:HB3	1.84	0.43
5:W:617:VAL:HG13	5:W:658:VAL:CG1	2.47	0.43
5:W:1241:LEU:HD13	5:W:1260:LEU:HB3	2.01	0.43
6:X:242:ARG:HE	6:X:1085:LEU:CD1	2.30	0.43
6:X:371:LEU:HD11	6:X:388:MET:HG2	2.01	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
6:X:653:PRO:HD3	6:X:773:ARG:CB	2.49	0.43
6:X:894:THR:O	6:X:895:SER:C	2.62	0.43
6:X:964:VAL:O	6:X:968:PHE:N	2.48	0.43
6:X:992:ASN:C	6:X:992:ASN:ND2	2.75	0.43
6:X:1035:GLY:O	6:X:1036:ASN:HB3	2.19	0.43
6:X:1060:ARG:HA	6:X:1064:TRP:HB2	1.99	0.43
6:X:1083:TYR:O	6:X:1083:TYR:CG	2.71	0.43
6:Y:106:MET:O	6:Y:106:MET:HG2	2.18	0.43
6:Y:188:ALA:HB1	6:Y:856:ARG:CD	2.48	0.43
6:Y:269:SER:HB2	6:Y:271:VAL:HG12	2.00	0.43
6:Y:913:PHE:O	6:Y:915:LEU:HD13	2.18	0.43
6:Y:1016:ARG:C	6:Y:1018:GLY:H	2.25	0.43
6:Y:1031:LEU:HD12	6:Y:1032:ILE:H	1.83	0.43
6:Y:1107:ILE:HG12	6:Y:1113:PRO:HD2	1.99	0.43
1:l:23:THR:CG2	1:l:25:TYR:CD2	2.99	0.43
1:f:15:ARG:HH12	3:F:402:ALA:CB	2.29	0.43
1:d:24:LEU:HD13	1:d:24:LEU:HA	1.89	0.43
1:n:5:MET:O	1:n:6:ILE:CB	2.67	0.43
1:n:5:MET:HE2	1:n:5:MET:HB2	1.80	0.43
1:p:60:ALA:O	1:p:62:VAL:CG1	2.58	0.43
2:A:23:THR:C	2:A:25:ASP:H	2.26	0.43
3:B:141:GLN:HE22	3:B:144:ASN:ND2	2.15	0.43
3:B:603:GLY:C	3:B:605:GLY:N	2.75	0.43
3:D:92:TRP:CZ2	3:D:160:GLN:CB	3.01	0.43
3:D:337:ARG:NH2	3:H:418:ASN:CG	2.74	0.43
3:D:362:THR:O	3:D:364:PRO:HD3	2.18	0.43
3:D:609:ARG:O	2:E:29:THR:HA	2.19	0.43
3:F:67:ARG:HH11	3:F:67:ARG:CG	2.29	0.43
3:F:92:TRP:CZ2	3:F:160:GLN:CB	3.01	0.43
3:F:199:LEU:CD1	3:F:203:TYR:CE2	3.01	0.43
3:F:208:TYR:CE2	3:F:212:ILE:HD11	2.53	0.43
3:F:293:SER:HB3	3:F:465:THR:HG23	2.01	0.43
3:F:588:ILE:HD13	3:F:588:ILE:HA	1.80	0.43
3:H:330:GLN:O	3:H:331:LEU:HB2	2.17	0.43
3:H:330:GLN:HB2	3:H:457:PRO:HG3	1.99	0.43
3:H:362:THR:O	3:H:364:PRO:HD3	2.18	0.43
3:H:401:GLY:O	3:H:402:ALA:CB	2.66	0.43
3:H:456:ASN:ND2	3:H:458:THR:HG22	2.33	0.43
3:H:551:GLN:NE2	3:H:558:THR:HB	2.34	0.43
3:J:203:TYR:N	3:J:204:PRO:HD2	2.34	0.43
3:J:551:GLN:NE2	3:J:558:THR:HB	2.34	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:J:609:ARG:O	2:K:29:THR:HA	2.19	0.43
3:L:96:PHE:CE1	3:L:159:TRP:HH2	2.37	0.43
3:L:345:ILE:O	3:L:347:SER:N	2.52	0.43
2:O:24:SER:CB	3:P:176:ALA:N	2.80	0.43
3:P:92:TRP:CZ2	3:P:160:GLN:CB	3.01	0.43
3:P:199:LEU:CD1	3:P:203:TYR:CE2	3.01	0.43
3:P:540:VAL:HG21	3:P:566:TRP:HA	2.00	0.43
2:Q:23:THR:C	2:Q:25:ASP:H	2.26	0.43
3:R:96:PHE:CE1	3:R:159:TRP:HH2	2.37	0.43
3:R:397:VAL:O	3:R:398:SER:HB2	2.18	0.43
3:T:262:ALA:CB	3:T:608:LEU:HD13	2.47	0.43
3:T:603:GLY:C	3:T:605:GLY:N	2.75	0.43
4:U:348:THR:HG22	4:U:350:ALA:H	1.83	0.43
5:W:173:ALA:HB3	5:W:202:ALA:HB3	2.00	0.43
5:W:194:TYR:HB2	5:W:196:LYS:HD3	2.01	0.43
5:W:335:SER:CA	5:W:752:ASN:O	2.64	0.43
5:W:405:ARG:HB3	5:W:779:MET:HB2	2.00	0.43
5:W:715:LEU:HB3	5:W:718:PRO:HB3	2.00	0.43
5:W:934:ASN:ND2	5:W:934:ASN:N	2.58	0.43
5:W:996:ASP:O	5:W:999:PRO:HD2	2.18	0.43
5:W:1052:ILE:HD11	5:W:1110:VAL:HG11	2.00	0.43
6:X:248:ASN:HD22	6:X:260:GLU:CD	2.26	0.43
6:X:747:ILE:O	6:X:747:ILE:CG2	2.62	0.43
6:X:1068:LYS:C	6:X:1070:GLY:H	2.27	0.43
6:X:1189:THR:C	6:X:1191:TYR:H	2.27	0.43
6:Y:134:GLU:CD	6:Y:134:GLU:N	2.76	0.43
6:Y:200:LEU:HB2	6:Y:250:ALA:CB	2.47	0.43
6:Y:207:LEU:HD23	6:Y:207:LEU:C	2.44	0.43
6:Y:607:HIS:O	6:Y:610:THR:N	2.49	0.43
6:Y:893:VAL:HG13	6:Y:897:ALA:CB	2.48	0.43
6:Y:1172:LEU:HB2	6:Y:1191:TYR:CD2	2.53	0.43
1:d:29:ARG:NH2	1:d:65:ILE:O	2.49	0.43
1:h:23:THR:HG21	3:H:323:SER:CB	2.44	0.43
2:A:9:ASN:HD21	3:B:206:GLU:CG	2.32	0.43
3:B:565:LEU:HD23	3:B:565:LEU:N	2.34	0.43
3:B:588:ILE:HG22	3:B:591:LEU:HB2	2.00	0.43
3:D:100:LYS:HZ3	3:L:147:ARG:HH12	1.67	0.43
3:D:101:THR:HG23	3:D:102:GLY:N	2.33	0.43
3:D:193:ILE:O	3:D:194:LEU:C	2.61	0.43
3:D:540:VAL:HG21	3:D:566:TRP:HA	2.00	0.43
3:F:107:VAL:HG21	3:H:128:ASP:OD2	2.18	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:F:135:VAL:O	3:F:135:VAL:HG13	2.19	0.43
3:F:256:SER:OG	3:F:257:SER:N	2.51	0.43
3:F:412:ARG:CA	3:H:455:SER:O	2.66	0.43
3:F:420:LEU:HD11	3:F:422:LEU:CD2	2.49	0.43
3:F:567:ARG:NH1	3:F:567:ARG:HB2	2.33	0.43
3:H:135:VAL:HG13	3:H:135:VAL:O	2.19	0.43
3:J:107:VAL:HG21	3:L:128:ASP:OD2	2.18	0.43
3:J:136:PHE:N	3:J:136:PHE:CD1	2.87	0.43
3:J:310:PHE:CD1	3:J:382:ILE:HG13	2.54	0.43
3:J:567:ARG:HB2	3:J:567:ARG:NH1	2.33	0.43
3:J:599:ALA:O	3:J:600:LEU:C	2.60	0.43
3:L:199:LEU:CD1	3:L:203:TYR:CE2	3.01	0.43
3:L:551:GLN:NE2	3:L:558:THR:HB	2.34	0.43
3:L:609:ARG:CD	2:M:29:THR:HG21	2.47	0.43
3:N:135:VAL:O	3:N:135:VAL:HG13	2.19	0.43
3:N:397:VAL:O	3:N:398:SER:HB2	2.18	0.43
3:N:463:LEU:HD23	3:N:463:LEU:C	2.44	0.43
3:P:203:TYR:HB3	3:P:206:GLU:CB	2.39	0.43
3:P:292:ARG:HG2	3:P:292:ARG:NH1	2.32	0.43
3:P:330:GLN:HB2	3:P:457:PRO:HG3	1.99	0.43
3:P:369:SER:CB	3:P:446:PRO:HD3	2.47	0.43
3:P:401:GLY:O	3:P:402:ALA:CB	2.66	0.43
3:P:471:LEU:HD23	3:P:471:LEU:N	2.32	0.43
2:Q:9:ASN:HD21	3:R:206:GLU:CG	2.32	0.43
2:Q:18:ASN:O	2:Q:19:SER:HB3	2.18	0.43
3:R:540:VAL:HG21	3:R:566:TRP:HA	2.00	0.43
3:R:564:LEU:CD2	3:R:568:ARG:NH2	2.75	0.43
3:T:96:PHE:CE1	3:T:159:TRP:HH2	2.37	0.43
3:T:456:ASN:ND2	3:T:458:THR:HG22	2.33	0.43
3:T:533:ASP:OD1	3:T:536:THR:HG23	2.18	0.43
3:T:588:ILE:HG22	3:T:591:LEU:HB2	2.00	0.43
4:U:26:LEU:CA	4:U:71:LEU:HD11	2.34	0.43
4:U:156:ILE:HD13	4:U:305:LEU:HD11	2.00	0.43
4:U:159:LEU:CD1	4:U:267:ILE:HD12	2.48	0.43
4:U:251:ASN:ND2	4:U:253:GLY:N	2.67	0.43
5:W:88:ILE:C	5:W:90:ASP:H	2.27	0.43
5:W:129:ARG:O	5:W:130:SER:C	2.62	0.43
5:W:213:ARG:HG2	5:W:217:LEU:CD2	2.49	0.43
5:W:316:GLN:OE1	5:W:316:GLN:CA	2.66	0.43
5:W:994:THR:HG23	5:W:1066:ASN:O	2.19	0.43
5:W:1063:VAL:CG2	5:W:1110:VAL:HG22	2.45	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
6:X:194:ILE:HG21	6:X:1201:TYR:CZ	2.53	0.43
6:X:617:HIS:ND1	6:X:619:GLY:N	2.60	0.43
6:X:694:ASP:O	6:X:695:SER:HB3	2.19	0.43
6:X:793:ARG:HH11	6:Y:693:TYR:HE2	1.57	0.43
6:X:862:GLU:O	6:X:863:ALA:C	2.62	0.43
6:Y:49:GLN:CG	6:Y:50:GLN:N	2.81	0.43
6:Y:404:MET:HG2	6:Y:956:HIS:CE1	2.53	0.43
6:Y:832:ILE:O	6:Y:833:PRO:C	2.60	0.43
6:Y:838:GLY:HA3	6:Y:1210:VAL:HG23	2.00	0.43
1:b:74:GLN:HA	1:b:75:PRO:CD	2.47	0.43
1:f:15:ARG:NH2	3:F:402:ALA:HB2	2.30	0.43
1:d:15:ARG:NH2	3:D:402:ALA:HB2	2.34	0.43
1:h:45:ARG:O	1:h:58:PRO:O	2.37	0.43
1:j:44:GLY:O	3:J:399:ALA:CB	2.66	0.43
1:r:5:MET:O	1:r:6:ILE:O	2.36	0.43
3:B:341:MET:CE	3:B:341:MET:CA	2.88	0.43
3:B:463:LEU:HD23	3:B:463:LEU:C	2.44	0.43
2:C:18:ASN:O	2:C:19:SER:HB3	2.18	0.43
3:D:154:ARG:O	3:D:155:ASP:C	2.62	0.43
3:F:48:TRP:CE2	3:F:63:ILE:CD1	3.02	0.43
3:H:96:PHE:CE1	3:H:159:TRP:HH2	2.37	0.43
3:H:101:THR:HG23	3:H:102:GLY:N	2.34	0.43
3:H:370:VAL:O	3:H:370:VAL:CG2	2.67	0.43
3:H:387:THR:O	3:H:390:ASN:HB2	2.18	0.43
3:H:420:LEU:HD11	3:H:422:LEU:CD2	2.49	0.43
3:H:463:LEU:HD23	3:H:463:LEU:C	2.44	0.43
2:I:23:THR:C	2:I:25:ASP:H	2.26	0.43
7:I:101:MYR:H42	3:J:187:LEU:HD22	2.00	0.43
3:J:46:LYS:O	3:J:48:TRP:CD1	2.71	0.43
3:J:128:ASP:OD2	3:N:107:VAL:HG21	2.18	0.43
3:L:203:TYR:N	3:L:204:PRO:HD2	2.34	0.43
3:L:352:ASP:OD1	3:L:352:ASP:C	2.62	0.43
3:L:567:ARG:NH1	3:L:567:ARG:HB2	2.33	0.43
3:N:203:TYR:N	3:N:204:PRO:HD2	2.34	0.43
3:N:420:LEU:HD11	3:N:422:LEU:CD2	2.49	0.43
3:N:456:ASN:C	3:N:456:ASN:ND2	2.76	0.43
3:P:345:ILE:O	3:P:347:SER:N	2.52	0.43
3:P:387:THR:O	3:P:390:ASN:HB2	2.18	0.43
3:P:463:LEU:HD23	3:P:463:LEU:C	2.44	0.43
3:P:551:GLN:NE2	3:P:558:THR:HB	2.34	0.43
3:P:565:LEU:N	3:P:565:LEU:HD23	2.34	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:R:101:THR:HG23	3:R:102:GLY:N	2.34	0.43
3:R:107:VAL:HG21	3:T:128:ASP:OD2	2.18	0.43
3:R:266:MET:HG2	3:R:520:VAL:HG11	2.01	0.43
3:R:332:GLY:HA3	3:R:456:ASN:HA	2.01	0.43
3:R:362:THR:O	3:R:364:PRO:HD3	2.18	0.43
3:R:456:ASN:ND2	3:R:458:THR:HG22	2.33	0.43
3:T:48:TRP:CE2	3:T:63:ILE:CD1	3.02	0.43
3:T:318:LYS:HG3	3:T:319:THR:H	1.84	0.43
3:T:332:GLY:HA3	3:T:456:ASN:HA	2.01	0.43
3:T:341:MET:CG	3:T:431:ILE:HD11	2.49	0.43
3:T:428:ARG:HH11	3:T:428:ARG:CG	2.16	0.43
4:U:225:ARG:HG3	4:U:225:ARG:NH1	2.32	0.43
4:U:225:ARG:HA	4:U:233:TYR:CE1	2.54	0.43
4:V:178:THR:OG1	6:Y:407:ARG:CB	2.66	0.43
5:W:1255:ASN:O	5:W:1257:PHE:N	2.51	0.43
6:X:359:PHE:O	6:X:360:LEU:C	2.60	0.43
6:X:670:ILE:HG22	6:X:671:PRO:HD2	2.00	0.43
6:X:707:PHE:O	6:X:710:ARG:HB3	2.19	0.43
6:Y:81:ARG:HD2	6:Y:85:ASP:OD1	2.19	0.43
6:Y:257:ARG:H	6:Y:257:ARG:HG2	1.61	0.43
6:Y:259:LEU:O	6:Y:307:SER:HB2	2.18	0.43
6:Y:497:ILE:HG23	6:Y:498:THR:H	1.83	0.43
6:Y:497:ILE:O	6:Y:500:ILE:HD12	2.19	0.43
6:Y:543:THR:O	6:Y:576:THR:HB	2.19	0.43
6:Y:547:GLU:OE1	6:Y:547:GLU:CA	2.66	0.43
6:Y:865:VAL:CG2	6:Y:952:PRO:HB2	2.49	0.43
6:Y:919:LEU:CD2	6:Y:920:LEU:N	2.82	0.43
6:Y:1095:ASP:OD1	6:Y:1097:ALA:N	2.51	0.43
6:Y:1156:THR:CG2	6:Y:1161:ASN:CB	2.96	0.43
1:p:51:PHE:CE1	1:p:71:CYS:O	2.71	0.43
1:r:3:LEU:HG	3:T:492:ASN:ND2	2.34	0.43
3:B:203:TYR:N	3:B:204:PRO:HD2	2.34	0.43
3:B:401:GLY:O	3:B:402:ALA:CB	2.66	0.43
3:B:433:GLY:O	3:B:435:PRO:CD	2.67	0.43
3:B:540:VAL:HG21	3:B:566:TRP:HA	2.00	0.43
7:C:101:MYR:H21	3:D:191:LYS:NZ	2.30	0.43
3:D:96:PHE:CD1	3:D:101:THR:HG23	2.44	0.43
3:D:588:ILE:HG22	3:D:591:LEU:HB2	2.00	0.43
3:D:603:GLY:C	3:D:605:GLY:N	2.75	0.43
3:F:62:ALA:O	3:F:63:ILE:C	2.62	0.43
3:F:588:ILE:HG22	3:F:591:LEU:HB2	2.00	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:H:136:PHE:CD1	3:H:136:PHE:N	2.87	0.43
3:H:411:VAL:O	3:H:411:VAL:CG1	2.61	0.43
3:H:427:GLU:HB2	3:H:430:THR:CG2	2.49	0.43
3:H:517:LEU:C	3:H:517:LEU:HD13	2.44	0.43
2:I:18:ASN:O	2:I:19:SER:HB3	2.18	0.43
3:J:199:LEU:CD1	3:J:203:TYR:CE2	3.01	0.43
3:J:256:SER:OG	3:J:257:SER:N	2.51	0.43
3:J:266:MET:HG2	3:J:520:VAL:HG11	2.01	0.43
3:J:431:ILE:O	3:J:431:ILE:HG23	2.10	0.43
2:K:9:ASN:HD21	3:L:206:GLU:CG	2.32	0.43
3:L:266:MET:HG2	3:L:520:VAL:HG11	2.01	0.43
3:L:362:THR:O	3:L:364:PRO:HD3	2.18	0.43
3:L:456:ASN:ND2	3:L:458:THR:HG22	2.33	0.43
3:N:318:LYS:HG3	3:N:319:THR:H	1.84	0.43
3:N:456:ASN:ND2	3:N:458:THR:HG22	2.33	0.43
3:P:271:LEU:HD12	3:P:271:LEU:HA	1.79	0.43
3:P:420:LEU:HD11	3:P:422:LEU:CD2	2.49	0.43
3:P:433:GLY:O	3:P:435:PRO:CD	2.67	0.43
3:P:590:VAL:O	3:P:592:THR:N	2.52	0.43
3:R:215:TYR:C	3:R:217:ALA:N	2.70	0.43
3:R:600:LEU:O	3:R:601:THR:C	2.60	0.43
3:T:144:ASN:O	3:T:145:LEU:C	2.62	0.43
3:T:203:TYR:N	3:T:204:PRO:HD2	2.34	0.43
3:T:428:ARG:NH1	3:T:428:ARG:CG	2.76	0.43
3:T:582:LEU:O	3:T:584:LEU:N	2.51	0.43
3:T:590:VAL:O	3:T:592:THR:N	2.52	0.43
4:U:8:GLY:HA2	4:U:146:THR:OG1	2.19	0.43
4:U:60:SER:OG	4:U:275:GLN:O	2.37	0.43
5:W:23:LEU:C	5:W:23:LEU:CD2	2.91	0.43
5:W:135:LEU:CD2	5:W:138:LEU:HD12	2.49	0.43
5:W:517:ALA:HB2	5:W:575:PHE:CE2	2.52	0.43
5:W:578:VAL:HG23	5:W:610:THR:HG22	2.01	0.43
5:W:661:GLY:O	5:W:663:GLN:N	2.45	0.43
5:W:695:ILE:HA	5:W:696:PRO:HD3	1.79	0.43
5:W:771:ARG:O	5:W:774:LEU:HB2	2.18	0.43
5:W:910:LEU:HD13	5:W:910:LEU:O	2.19	0.43
5:W:1165:VAL:HG11	5:W:1181:ALA:HB3	2.01	0.43
5:W:1204:LEU:HD23	5:W:1204:LEU:H	1.84	0.43
6:X:242:ARG:HH12	6:X:1146:LEU:CD1	2.32	0.43
6:X:252:LEU:O	6:X:253:GLY:C	2.61	0.43
6:X:390:LEU:C	6:X:390:LEU:HD23	2.44	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
6:X:509:SER:O	6:X:510:PRO:C	2.62	0.43
6:X:590:ALA:HB1	6:X:599:ILE:CD1	2.49	0.43
6:X:757:VAL:CG1	6:X:845:PHE:HE2	2.26	0.43
6:X:904:ALA:O	6:X:907:ASP:HB2	2.18	0.43
6:Y:33:PRO:HA	6:Y:34:PRO:HD3	1.86	0.43
6:Y:399:ARG:C	6:Y:401:THR:N	2.75	0.43
6:Y:403:ILE:CG2	6:Y:404:MET:N	2.82	0.43
6:Y:1089:ASP:C	6:Y:1091:GLN:N	2.75	0.43
6:Y:1182:ASN:HB3	6:Y:1183:ASP:H	1.41	0.43
1:l:45:ARG:N	3:L:399:ALA:O	2.52	0.43
1:l:60:ALA:O	3:L:397:VAL:O	2.37	0.43
1:d:5:MET:O	1:d:9:VAL:HG21	2.19	0.43
1:d:5:MET:HE1	1:d:53:LEU:HD23	2.01	0.43
2:A:18:ASN:O	2:A:19:SER:HB3	2.18	0.43
3:B:96:PHE:CE1	3:B:159:TRP:HH2	2.37	0.43
3:B:101:THR:HG23	3:B:102:GLY:N	2.34	0.43
3:B:256:SER:OG	3:B:257:SER:N	2.51	0.43
3:B:318:LYS:HG3	3:B:319:THR:H	1.84	0.43
3:B:332:GLY:HA3	3:B:456:ASN:HA	2.01	0.43
3:D:46:LYS:HD3	3:D:70:GLN:OE1	2.19	0.43
3:D:199:LEU:CD1	3:D:203:TYR:CE2	3.01	0.43
3:D:336:THR:CG2	3:D:456:ASN:ND2	2.79	0.43
3:D:412:ARG:CA	3:F:455:SER:O	2.66	0.43
3:D:556:SER:N	3:D:557:PRO:CD	2.82	0.43
3:F:310:PHE:CD1	3:F:382:ILE:HG13	2.54	0.43
3:F:418:ASN:HD22	3:F:418:ASN:HA	1.50	0.43
3:F:463:LEU:HD23	3:F:463:LEU:C	2.44	0.43
3:F:551:GLN:NE2	3:F:558:THR:HB	2.34	0.43
3:F:590:VAL:O	3:F:592:THR:N	2.52	0.43
3:F:603:GLY:C	3:F:605:GLY:N	2.75	0.43
3:H:183:TRP:NE1	3:H:187:LEU:CD1	2.81	0.43
3:H:203:TYR:N	3:H:204:PRO:HD2	2.34	0.43
3:H:256:SER:OG	3:H:257:SER:N	2.51	0.43
3:H:259:LEU:HD23	3:H:259:LEU:HA	1.88	0.43
3:H:491:ALA:HB2	3:H:592:THR:CG2	2.48	0.43
3:H:588:ILE:HD13	3:H:588:ILE:HA	1.80	0.43
3:H:590:VAL:O	3:H:592:THR:N	2.52	0.43
3:J:135:VAL:O	3:J:135:VAL:HG13	2.19	0.43
3:J:332:GLY:HA3	3:J:456:ASN:HA	2.01	0.43
3:J:341:MET:CG	3:J:431:ILE:HD11	2.49	0.43
3:L:136:PHE:CD1	3:L:136:PHE:N	2.87	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:L:293:SER:HB3	3:L:465:THR:HG23	2.01	0.43
3:L:310:PHE:CD1	3:L:382:ILE:HG13	2.54	0.43
3:L:463:LEU:HD23	3:L:463:LEU:C	2.44	0.43
3:L:625:ARG:HG3	3:R:197:ASP:CG	2.42	0.43
3:N:96:PHE:CE1	3:N:159:TRP:HH2	2.37	0.43
3:N:199:LEU:CD1	3:N:203:TYR:CE2	3.01	0.43
3:N:433:GLY:O	3:N:435:PRO:CD	2.67	0.43
3:P:203:TYR:N	3:P:204:PRO:HD2	2.34	0.43
3:P:427:GLU:CD	3:R:571:ARG:HH22	2.20	0.43
3:R:135:VAL:O	3:R:135:VAL:HG13	2.19	0.43
3:R:144:ASN:O	3:R:145:LEU:C	2.62	0.43
3:R:345:ILE:O	3:R:347:SER:N	2.52	0.43
3:R:463:LEU:HD23	3:R:463:LEU:C	2.44	0.43
3:R:590:VAL:O	3:R:592:THR:N	2.52	0.43
2:S:13:ILE:CD1	7:S:101:MYR:H111	2.46	0.43
3:T:62:ALA:O	3:T:63:ILE:C	2.62	0.43
3:T:136:PHE:N	3:T:136:PHE:CD1	2.87	0.43
3:T:143:MET:O	3:T:146:GLN:HB3	2.19	0.43
3:T:154:ARG:O	3:T:155:ASP:C	2.62	0.43
3:T:343:ALA:CB	3:T:478:GLY:O	2.62	0.43
3:T:420:LEU:HD11	3:T:422:LEU:CD2	2.49	0.43
4:U:208:ILE:O	4:U:208:ILE:HG23	2.18	0.43
4:U:329:ASN:HB3	4:U:332:PRO:HD2	2.00	0.43
4:V:76:LEU:HD13	4:V:120:LEU:HD11	2.01	0.43
4:V:188:THR:CG2	6:Y:376:LEU:HD11	2.49	0.43
5:W:51:PRO:O	5:W:52:VAL:C	2.60	0.43
5:W:170:ARG:HH21	6:X:682:GLN:NE2	2.17	0.43
5:W:225:THR:CG2	5:W:226:VAL:N	2.81	0.43
5:W:950:ASP:HB3	5:W:955:VAL:CG1	2.48	0.43
5:W:959:ASN:CB	5:W:962:GLU:CD	2.92	0.43
6:X:288:PHE:CE1	6:X:293:LEU:HD13	2.54	0.43
6:X:407:ARG:O	6:X:409:MET:N	2.51	0.43
6:X:544:VAL:HG12	6:X:545:LEU:N	2.34	0.43
6:X:628:PRO:HB2	6:X:644:GLU:OE2	2.19	0.43
6:X:663:GLN:O	6:X:664:GLY:C	2.60	0.43
6:X:769:VAL:HG13	6:X:770:PRO:HD2	2.01	0.43
6:X:841:PHE:O	6:X:844:LEU:HD12	2.18	0.43
6:X:1150:ASN:ND2	6:X:1153:SER:CB	2.82	0.43
6:Y:191:PRO:CA	6:Y:915:LEU:HD11	2.49	0.43
6:Y:1153:SER:C	6:Y:1155:ASP:N	2.75	0.43
6:Y:1179:VAL:HG13	6:Y:1193:SER:OG	2.18	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:d:14:VAL:O	1:d:18:ALA:N	2.51	0.42
1:n:44:GLY:O	3:N:399:ALA:O	2.37	0.42
1:p:59:HIS:HB2	3:P:398:SER:N	2.13	0.42
1:t:24:LEU:CD2	1:t:35:PHE:CB	2.97	0.42
3:B:143:MET:O	3:B:146:GLN:HB3	2.19	0.42
3:B:154:ARG:O	3:B:155:ASP:C	2.62	0.42
3:B:257:SER:C	3:B:259:LEU:N	2.74	0.42
3:B:345:ILE:O	3:B:347:SER:N	2.52	0.42
3:B:427:GLU:HB2	3:B:430:THR:CG2	2.49	0.42
2:C:29:THR:HA	3:H:609:ARG:O	2.19	0.42
3:D:237:GLU:CB	3:F:46:LYS:NZ	2.67	0.42
3:D:238:VAL:HG21	3:F:78:MET:HE2	1.83	0.42
3:D:420:LEU:HD11	3:D:422:LEU:CD2	2.49	0.42
3:D:433:GLY:O	3:D:435:PRO:CD	2.67	0.42
3:F:163:LEU:HD11	3:F:211:ALA:HB2	2.01	0.42
3:H:62:ALA:O	3:H:63:ILE:C	2.62	0.42
3:H:73:PHE:CB	3:H:119:VAL:HG21	2.49	0.42
3:H:92:TRP:CZ2	3:H:160:GLN:CB	3.01	0.42
3:J:46:LYS:HD3	3:J:70:GLN:OE1	2.19	0.42
3:J:49:ARG:HG3	3:J:64:VAL:CG2	2.45	0.42
3:J:51:VAL:N	3:J:60:SER:O	2.38	0.42
3:L:564:LEU:CD2	3:L:568:ARG:NH2	2.75	0.42
3:N:48:TRP:CE2	3:N:63:ILE:CD1	3.02	0.42
3:N:266:MET:HG2	3:N:520:VAL:HG11	2.01	0.42
3:N:551:GLN:NE2	3:N:558:THR:HB	2.34	0.42
3:N:588:ILE:HG22	3:N:591:LEU:HB2	2.00	0.42
3:N:625:ARG:HH11	3:N:625:ARG:HG3	1.84	0.42
2:O:23:THR:C	2:O:25:ASP:N	2.75	0.42
2:O:27:THR:HG22	2:O:29:THR:HG23	2.01	0.42
3:P:259:LEU:HD23	3:P:259:LEU:HA	1.88	0.42
3:P:266:MET:HG2	3:P:520:VAL:HG11	2.01	0.42
3:P:609:ARG:O	2:Q:29:THR:HA	2.19	0.42
3:R:163:LEU:HD11	3:R:211:ALA:HB2	2.01	0.42
3:T:135:VAL:O	3:T:135:VAL:HG13	2.19	0.42
3:T:266:MET:HG2	3:T:520:VAL:HG11	2.01	0.42
3:T:293:SER:HB3	3:T:465:THR:HG23	2.01	0.42
3:T:551:GLN:NE2	3:T:558:THR:HB	2.34	0.42
4:U:32:CYS:O	4:U:33:TYR:HB2	2.18	0.42
4:U:53:VAL:HB	4:U:60:SER:C	2.43	0.42
4:U:156:ILE:HG12	4:U:268:ARG:HD3	1.99	0.42
4:V:191:ARG:NH1	6:Y:438:PRO:CG	2.80	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
4:V:211:LEU:HD11	4:V:248:LEU:HD22	2.00	0.42
5:W:91:ARG:NH2	5:W:125:PHE:CZ	2.87	0.42
5:W:141:ASP:OD1	5:W:142:PRO:HD2	2.19	0.42
5:W:168:SER:HA	5:W:206:TYR:O	2.19	0.42
5:W:243:GLN:HE21	5:W:243:GLN:H	1.67	0.42
5:W:405:ARG:HG2	5:W:406:ASN:H	1.84	0.42
5:W:662:ARG:O	5:W:662:ARG:CG	2.67	0.42
5:W:733:ALA:HB1	5:W:777:VAL:HG12	2.00	0.42
5:W:927:THR:OG1	5:W:928:ARG:HG2	2.19	0.42
5:W:1242:PRO:HD3	5:W:1296:TYR:OH	2.18	0.42
6:X:294:ARG:HB2	6:X:888:LEU:HB3	2.01	0.42
6:X:377:ILE:HD11	6:X:436:LEU:HD11	2.00	0.42
6:X:557:ILE:O	6:X:557:ILE:CG1	2.66	0.42
6:X:972:MET:O	6:X:973:ASN:C	2.62	0.42
6:X:1130:ALA:CA	6:Y:104:GLN:HB3	2.49	0.42
6:X:1167:LEU:HD13	6:X:1187:ARG:CZ	2.49	0.42
6:Y:188:ALA:HB1	6:Y:856:ARG:HD2	2.01	0.42
6:Y:305:THR:O	6:Y:306:LEU:C	2.62	0.42
1:b:24:LEU:HD21	1:b:80:ARG:NE	2.31	0.42
1:f:45:ARG:CG	3:F:400:ALA:C	2.82	0.42
2:A:27:THR:HG22	2:A:29:THR:HG23	2.02	0.42
3:B:551:GLN:NE2	3:B:558:THR:HB	2.34	0.42
3:B:582:LEU:HD23	3:B:585:GLN:HE22	1.83	0.42
3:D:183:TRP:NE1	3:D:187:LEU:CD1	2.81	0.42
3:D:355:GLY:HA2	3:F:298:PHE:CE1	2.47	0.42
3:D:387:THR:O	3:D:390:ASN:HB2	2.18	0.42
3:D:463:LEU:HD23	3:D:463:LEU:C	2.44	0.42
3:D:582:LEU:HD23	3:D:582:LEU:HA	1.69	0.42
2:G:18:ASN:O	2:G:19:SER:HB3	2.18	0.42
3:H:163:LEU:HD11	3:H:211:ALA:HB2	2.01	0.42
3:H:433:GLY:O	3:H:435:PRO:CD	2.67	0.42
3:J:101:THR:HG23	3:J:102:GLY:N	2.33	0.42
3:J:154:ARG:O	3:J:155:ASP:C	2.62	0.42
3:J:166:ALA:HB2	3:J:194:LEU:HD11	2.02	0.42
3:J:450:SER:CA	3:J:453:ASN:HD22	2.21	0.42
3:L:332:GLY:HA3	3:L:456:ASN:HA	2.01	0.42
3:L:365:ASP:HB2	3:L:451:SER:OG	2.20	0.42
3:L:433:GLY:O	3:L:435:PRO:CD	2.67	0.42
3:L:517:LEU:C	3:L:517:LEU:HD13	2.44	0.42
3:L:625:ARG:HG3	3:L:625:ARG:HH11	1.84	0.42
3:N:310:PHE:CD1	3:N:382:ILE:HG13	2.54	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:N:352:ASP:OD1	3:N:352:ASP:C	2.62	0.42
3:N:365:ASP:HB2	3:N:451:SER:OG	2.20	0.42
3:N:590:VAL:O	3:N:592:THR:N	2.52	0.42
7:O:101:MYR:H42	3:P:187:LEU:HD22	2.00	0.42
3:R:154:ARG:O	3:R:155:ASP:C	2.62	0.42
3:R:171:THR:HG21	3:R:634:LYS:NZ	2.35	0.42
3:R:433:GLY:O	3:R:435:PRO:CD	2.67	0.42
3:T:556:SER:N	3:T:557:PRO:CD	2.82	0.42
4:U:35:ARG:H	4:U:35:ARG:HG2	1.70	0.42
4:U:136:TRP:N	4:U:137:PRO:CD	2.82	0.42
4:U:363:ALA:O	4:U:367:GLU:HB2	2.19	0.42
5:W:192:ALA:HB1	5:W:333:VAL:HG21	2.00	0.42
5:W:196:LYS:HG3	5:W:201:TYR:CZ	2.54	0.42
5:W:359:TYR:CZ	5:W:375:PRO:HB3	2.54	0.42
5:W:478:ARG:C	5:W:480:ARG:H	2.26	0.42
5:W:1009:ASN:C	5:W:1011:SER:N	2.77	0.42
6:X:593:LEU:HD22	6:X:639:LEU:HD13	2.01	0.42
6:X:1048:PRO:HB2	6:X:1051:VAL:HG23	2.01	0.42
6:X:1085:LEU:HD22	6:X:1119:ILE:HD12	2.01	0.42
6:X:1210:VAL:O	6:X:1211:ARG:HB2	2.19	0.42
6:Y:305:THR:HA	6:Y:398:ILE:HD13	2.01	0.42
6:Y:410:HIS:HD2	6:Y:411:GLN:C	2.27	0.42
6:Y:722:ASN:O	6:Y:726:ARG:HB2	2.20	0.42
6:Y:859:ILE:HD13	6:Y:859:ILE:HA	1.91	0.42
1:h:29:ARG:NH2	1:h:65:ILE:O	2.49	0.42
3:B:144:ASN:O	3:B:145:LEU:C	2.62	0.42
3:B:341:MET:CG	3:B:431:ILE:HD11	2.49	0.42
3:B:370:VAL:O	3:B:370:VAL:CG2	2.67	0.42
2:C:9:ASN:HD21	3:D:206:GLU:CG	2.32	0.42
2:C:13:ILE:CD1	7:C:101:MYR:H111	2.45	0.42
2:C:29:THR:CG2	3:H:609:ARG:CD	2.93	0.42
3:D:136:PHE:CD1	3:D:136:PHE:N	2.87	0.42
3:D:143:MET:O	3:D:146:GLN:HB3	2.19	0.42
3:D:266:MET:HG2	3:D:520:VAL:HG11	2.01	0.42
3:D:293:SER:HB3	3:D:465:THR:HG23	2.01	0.42
3:D:565:LEU:N	3:D:565:LEU:HD23	2.34	0.42
3:F:188:ALA:HA	3:F:191:LYS:HD2	2.01	0.42
3:F:365:ASP:HB2	3:F:451:SER:OG	2.20	0.42
3:F:416:ARG:CB	3:H:298:PHE:HE1	2.16	0.42
3:H:310:PHE:CD1	3:H:382:ILE:HG13	2.54	0.42
3:H:540:VAL:HG21	3:H:566:TRP:HA	2.00	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:H:625:ARG:HG2	3:L:197:ASP:OD1	2.18	0.42
3:J:427:GLU:HB2	3:J:430:THR:CG2	2.49	0.42
3:J:433:GLY:O	3:J:435:PRO:CD	2.67	0.42
3:J:565:LEU:N	3:J:565:LEU:HD23	2.34	0.42
3:J:590:VAL:O	3:J:592:THR:N	2.52	0.42
3:J:603:GLY:C	3:J:605:GLY:N	2.75	0.42
3:L:252:LEU:HB3	3:L:613:LEU:HD23	2.02	0.42
3:L:341:MET:CG	3:L:431:ILE:HD11	2.49	0.42
3:L:345:ILE:HG23	3:L:475:THR:HG1	1.84	0.42
7:M:101:MYR:C4	3:N:187:LEU:HD22	2.49	0.42
3:N:188:ALA:HA	3:N:191:LYS:HD2	2.02	0.42
3:N:634:LYS:HE3	3:N:634:LYS:HB2	1.89	0.42
2:O:29:THR:HA	3:T:609:ARG:O	2.19	0.42
3:P:143:MET:O	3:P:146:GLN:HB3	2.19	0.42
3:P:144:ASN:O	3:P:145:LEU:C	2.62	0.42
3:P:427:GLU:HB2	3:P:430:THR:CG2	2.49	0.42
3:P:517:LEU:C	3:P:517:LEU:HD13	2.44	0.42
3:R:252:LEU:HB3	3:R:613:LEU:HD23	2.01	0.42
3:R:279:PRO:HA	3:R:484:ILE:O	2.20	0.42
3:R:310:PHE:CD1	3:R:382:ILE:HG13	2.54	0.42
3:R:341:MET:CE	3:R:341:MET:CA	2.88	0.42
3:R:390:ASN:HD22	3:R:390:ASN:HA	1.60	0.42
3:R:427:GLU:HB2	3:R:430:THR:CG2	2.49	0.42
3:T:46:LYS:HD3	3:T:70:GLN:OE1	2.19	0.42
3:T:252:LEU:HB3	3:T:613:LEU:CD2	2.47	0.42
3:T:285:PHE:O	3:T:286:THR:C	2.62	0.42
3:T:456:ASN:C	3:T:456:ASN:ND2	2.76	0.42
4:U:175:MET:HG3	4:U:232:PHE:HZ	1.83	0.42
4:U:270:ILE:HG22	4:U:328:TYR:O	2.18	0.42
5:W:1107:ILE:HG23	5:W:1126:THR:CG2	2.48	0.42
5:W:1167:ILE:HG22	5:W:1205:LEU:HD23	2.00	0.42
6:X:485:PRO:O	6:X:827:THR:HG23	2.20	0.42
6:X:514:ILE:CG2	6:X:564:TRP:CD2	3.02	0.42
6:X:751:CYS:N	6:X:752:PRO:CD	2.82	0.42
6:X:780:ILE:CD1	6:X:940:GLN:HG3	2.50	0.42
6:X:1075:LEU:C	6:X:1075:LEU:CD2	2.93	0.42
6:Y:484:THR:HG21	6:Y:529:TRP:CZ2	2.55	0.42
6:Y:599:ILE:N	6:Y:608:GLN:OE1	2.52	0.42
6:Y:1048:PRO:HG2	6:Y:1051:VAL:CG2	2.50	0.42
1:l:61:ASN:HB3	3:L:397:VAL:HA	2.01	0.42
3:B:48:TRP:CE2	3:B:63:ILE:CD1	3.02	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:B:79:ARG:CB	3:B:79:ARG:HH11	2.33	0.42
3:B:135:VAL:O	3:B:135:VAL:HG13	2.19	0.42
3:B:310:PHE:CD1	3:B:382:ILE:HG13	2.54	0.42
3:B:616:GLY:O	3:B:620:ALA:HB2	2.20	0.42
3:D:171:THR:HG21	3:D:634:LYS:NZ	2.34	0.42
3:D:279:PRO:HA	3:D:484:ILE:O	2.20	0.42
2:E:9:ASN:HD21	3:F:206:GLU:CG	2.32	0.42
3:F:171:THR:HG21	3:F:634:LYS:NZ	2.34	0.42
3:F:266:MET:HG2	3:F:520:VAL:HG11	2.01	0.42
3:F:297:ALA:CB	3:F:460:MET:HB2	2.33	0.42
3:F:341:MET:CG	3:F:431:ILE:HD11	2.49	0.42
3:F:345:ILE:O	3:F:347:SER:N	2.52	0.42
3:F:397:VAL:O	3:F:398:SER:HB2	2.18	0.42
3:F:427:GLU:HB2	3:F:430:THR:CG2	2.49	0.42
3:H:318:LYS:HG3	3:H:319:THR:H	1.84	0.42
3:J:96:PHE:CE1	3:J:159:TRP:HH2	2.37	0.42
3:J:285:PHE:O	3:J:286:THR:C	2.62	0.42
3:J:288:SER:O	3:J:289:LEU:C	2.62	0.42
3:J:463:LEU:HD23	3:J:463:LEU:C	2.44	0.42
2:K:18:ASN:O	2:K:19:SER:HB3	2.18	0.42
3:L:355:GLY:HA2	3:N:298:PHE:CE1	2.47	0.42
3:L:616:GLY:O	3:L:620:ALA:HB2	2.20	0.42
3:L:618:THR:HG22	3:L:619:ASP:OD2	2.19	0.42
2:M:5:GLN:HB3	2:M:6:THR:H	1.74	0.42
2:M:27:THR:HG22	2:M:29:THR:HG23	2.02	0.42
3:N:517:LEU:HD13	3:N:517:LEU:C	2.44	0.42
3:N:556:SER:N	3:N:557:PRO:CD	2.82	0.42
3:N:618:THR:HG22	3:N:619:ASP:OD2	2.19	0.42
3:P:293:SER:HB3	3:P:465:THR:HG23	2.01	0.42
3:P:588:ILE:HG22	3:P:591:LEU:HB2	2.00	0.42
3:T:93:GLN:OE1	3:T:93:GLN:HA	2.19	0.42
3:T:616:GLY:O	3:T:620:ALA:HB2	2.20	0.42
4:U:194:GLY:O	4:U:195:GLN:HB3	2.19	0.42
4:V:174:LEU:HD21	4:V:211:LEU:HD13	2.00	0.42
5:W:360:TRP:CD1	5:W:378:PRO:HD2	2.55	0.42
5:W:632:PHE:HE2	5:W:674:LEU:HD22	1.84	0.42
5:W:769:ARG:CD	5:W:844:SER:O	2.65	0.42
5:W:916:GLN:O	5:W:920:ARG:HG3	2.19	0.42
5:W:1055:LEU:HD23	5:W:1056:ASP:H	1.84	0.42
5:W:1225:ASP:C	5:W:1227:TYR:H	2.25	0.42
6:X:299:LEU:HD11	6:X:904:ALA:CB	2.42	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
6:X:570:THR:O	6:X:571:LEU:C	2.63	0.42
6:X:632:ASN:HA	6:X:633:PRO:HD3	1.87	0.42
6:X:772:THR:CG2	6:X:775:VAL:HB	2.49	0.42
6:X:983:TYR:HE2	6:X:1002:PRO:HB2	1.83	0.42
6:X:1124:VAL:CG1	6:X:1153:SER:OG	2.61	0.42
6:X:1130:ALA:HB2	6:Y:104:GLN:HA	2.00	0.42
6:Y:97:ALA:HB3	6:Y:100:VAL:HG22	2.01	0.42
6:Y:593:LEU:H	6:Y:593:LEU:CD2	2.24	0.42
6:Y:865:VAL:HB	6:Y:955:SER:HB2	2.02	0.42
6:Y:867:ALA:HA	6:Y:905:VAL:HG11	2.00	0.42
6:Y:1021:VAL:O	6:Y:1021:VAL:CG2	2.64	0.42
1:l:20:GLY:HA3	1:l:83:VAL:HG13	2.00	0.42
1:l:80:ARG:NH1	1:l:84:GLU:OE2	2.49	0.42
1:d:21:ARG:HD3	3:D:327:TYR:OH	2.19	0.42
1:h:4:HIS:ND1	1:h:54:THR:HG22	2.35	0.42
1:h:44:GLY:O	3:H:399:ALA:C	2.62	0.42
1:r:15:ARG:HH22	3:R:402:ALA:HB3	1.84	0.42
1:t:6:ILE:HD13	3:P:586:VAL:CG2	2.37	0.42
1:t:21:ARG:NH1	3:T:327:TYR:HE1	1.96	0.42
3:B:136:PHE:CD1	3:B:136:PHE:N	2.87	0.42
3:B:352:ASP:OD1	3:B:352:ASP:C	2.62	0.42
3:D:237:GLU:CD	3:F:46:LYS:NZ	2.68	0.42
3:D:345:ILE:O	3:D:347:SER:N	2.52	0.42
2:E:18:ASN:O	2:E:19:SER:HB3	2.18	0.42
3:F:154:ARG:O	3:F:155:ASP:C	2.62	0.42
3:F:442:THR:OG1	3:F:443:PHE:N	2.50	0.42
3:F:556:SER:N	3:F:557:PRO:CD	2.82	0.42
3:H:235:LEU:HD13	3:H:235:LEU:HA	1.80	0.42
3:H:266:MET:HG2	3:H:520:VAL:HG11	2.01	0.42
3:H:508:VAL:O	3:H:508:VAL:HG12	2.20	0.42
3:H:565:LEU:N	3:H:565:LEU:HD23	2.34	0.42
3:H:588:ILE:HG22	3:H:591:LEU:HB2	2.00	0.42
2:I:27:THR:HG22	2:I:29:THR:HG23	2.02	0.42
3:J:171:THR:HG21	3:J:634:LYS:NZ	2.34	0.42
3:J:362:THR:O	3:J:364:PRO:HD3	2.18	0.42
3:J:397:VAL:O	3:J:398:SER:HB2	2.18	0.42
3:J:625:ARG:HH11	3:J:625:ARG:HG3	1.84	0.42
3:L:285:PHE:O	3:L:286:THR:C	2.62	0.42
3:L:318:LYS:HG3	3:L:319:THR:H	1.84	0.42
3:L:469:VAL:O	3:L:470:LEU:CB	2.64	0.42
3:L:590:VAL:O	3:L:592:THR:N	2.52	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:N:163:LEU:HD11	3:N:211:ALA:HB2	2.01	0.42
3:N:214:ARG:HB2	3:N:215:TYR:CE2	2.55	0.42
3:N:235:LEU:HD13	3:N:235:LEU:HA	1.80	0.42
3:N:341:MET:CE	3:N:341:MET:CA	2.88	0.42
3:P:79:ARG:CB	3:P:79:ARG:HH11	2.33	0.42
3:P:188:ALA:HA	3:P:191:LYS:HD2	2.01	0.42
3:P:288:SER:O	3:P:289:LEU:C	2.62	0.42
3:P:332:GLY:HA3	3:P:456:ASN:HA	2.01	0.42
3:P:341:MET:CE	3:P:341:MET:CA	2.88	0.42
3:P:341:MET:CG	3:P:431:ILE:HD11	2.49	0.42
3:P:469:VAL:O	3:P:469:VAL:HG22	2.20	0.42
3:P:618:THR:HG22	3:P:619:ASP:OD2	2.19	0.42
3:R:143:MET:O	3:R:146:GLN:HB3	2.19	0.42
3:R:178:SER:HB3	3:R:179:ALA:H	1.65	0.42
3:R:387:THR:O	3:R:390:ASN:HB2	2.18	0.42
3:R:508:VAL:O	3:R:508:VAL:HG12	2.20	0.42
3:R:551:GLN:NE2	3:R:558:THR:HB	2.34	0.42
3:T:199:LEU:CD1	3:T:203:TYR:CE2	3.01	0.42
3:T:365:ASP:HB2	3:T:451:SER:OG	2.20	0.42
3:T:433:GLY:O	3:T:435:PRO:CD	2.67	0.42
4:U:121:ASP:OD2	4:U:123:ASN:HB2	2.19	0.42
4:V:27:GLN:O	4:V:27:GLN:HG2	2.19	0.42
5:W:306:ARG:HD3	5:W:341:MET:CE	2.49	0.42
5:W:566:LEU:HD11	5:W:609:MET:CE	2.49	0.42
5:W:905:ALA:C	5:W:907:GLY:H	2.25	0.42
5:W:953:ASP:O	5:W:955:VAL:N	2.52	0.42
6:X:255:GLU:OE1	6:X:256:GLY:N	2.53	0.42
6:X:413:PRO:O	6:X:415:GLN:N	2.53	0.42
6:X:416:ILE:HD12	6:X:1207:ILE:HG13	2.00	0.42
6:X:662:LEU:N	6:X:662:LEU:CD2	2.75	0.42
6:X:750:MET:HE3	6:X:750:MET:HB2	1.63	0.42
6:X:786:ALA:HB2	6:X:804:HIS:NE2	2.34	0.42
6:X:1146:LEU:O	6:X:1148:SER:N	2.48	0.42
6:Y:633:PRO:CD	6:Y:640:ARG:HD3	2.48	0.42
6:Y:722:ASN:HD22	6:Y:724:LEU:HB3	1.84	0.42
1:b:59:HIS:O	3:B:398:SER:N	2.44	0.42
1:d:15:ARG:NH1	3:D:406:ASN:ND2	2.66	0.42
1:j:22:LEU:C	1:j:22:LEU:CD1	2.85	0.42
1:r:21:ARG:CB	1:r:43:CYS:O	2.67	0.42
3:B:163:LEU:HD11	3:B:211:ALA:HB2	2.01	0.42
3:D:46:LYS:HZ1	3:H:237:GLU:CB	2.08	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:D:96:PHE:CD1	3:D:159:TRP:CH2	3.08	0.42
3:D:397:VAL:O	3:D:398:SER:HB2	2.18	0.42
3:D:517:LEU:C	3:D:517:LEU:HD13	2.44	0.42
3:F:143:MET:O	3:F:146:GLN:HB3	2.20	0.42
3:F:352:ASP:OD1	3:F:352:ASP:C	2.62	0.42
3:F:618:THR:HG22	3:F:619:ASP:OD2	2.19	0.42
2:G:13:ILE:CD1	7:G:101:MYR:H111	2.46	0.42
3:H:46:LYS:HD3	3:H:70:GLN:OE1	2.19	0.42
3:H:143:MET:O	3:H:146:GLN:HB3	2.20	0.42
3:H:341:MET:CG	3:H:431:ILE:HD11	2.49	0.42
3:H:618:THR:HG22	3:H:619:ASP:OD2	2.19	0.42
3:H:625:ARG:HG3	3:H:625:ARG:HH11	1.84	0.42
2:I:26:MET:CG	3:J:216:PRO:CG	2.95	0.42
3:J:48:TRP:CE2	3:J:63:ILE:CD1	3.02	0.42
3:J:188:ALA:HA	3:J:191:LYS:HD2	2.01	0.42
3:J:279:PRO:HA	3:J:484:ILE:O	2.20	0.42
3:J:289:LEU:HA	3:L:578:SER:O	2.20	0.42
3:J:293:SER:HB3	3:J:465:THR:HG23	2.01	0.42
3:J:556:SER:N	3:J:557:PRO:CD	2.82	0.42
3:J:618:THR:HG22	3:J:619:ASP:OD2	2.19	0.42
3:L:171:THR:HG21	3:L:634:LYS:NZ	2.35	0.42
3:N:137:ILE:HD13	3:N:142:ILE:CG1	2.50	0.42
3:N:616:GLY:O	3:N:620:ALA:HB2	2.20	0.42
3:P:136:PHE:N	3:P:136:PHE:CD1	2.87	0.42
3:P:252:LEU:HB3	3:P:613:LEU:HD23	2.02	0.42
3:P:279:PRO:HA	3:P:484:ILE:O	2.20	0.42
3:P:616:GLY:O	3:P:620:ALA:HB2	2.19	0.42
3:P:622:GLU:OE1	3:T:281:PRO:CD	2.62	0.42
3:R:136:PHE:CD1	3:R:136:PHE:N	2.87	0.42
3:R:193:ILE:O	3:R:194:LEU:C	2.61	0.42
3:R:203:TYR:N	3:R:204:PRO:HD2	2.34	0.42
3:R:288:SER:O	3:R:289:LEU:C	2.62	0.42
3:R:292:ARG:HG2	3:R:292:ARG:NH1	2.32	0.42
3:R:344:MET:HE2	3:R:502:LEU:CD1	2.37	0.42
3:R:412:ARG:CA	3:T:455:SER:O	2.66	0.42
3:R:556:SER:N	3:R:557:PRO:CD	2.82	0.42
7:S:101:MYR:H21	3:T:191:LYS:NZ	2.25	0.42
3:T:336:THR:CG2	3:T:456:ASN:ND2	2.79	0.42
3:T:370:VAL:O	3:T:370:VAL:CG2	2.67	0.42
3:T:517:LEU:C	3:T:517:LEU:HD13	2.44	0.42
3:T:565:LEU:N	3:T:565:LEU:HD23	2.34	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:T:625:ARG:HG3	3:T:625:ARG:HH11	1.84	0.42
4:U:221:GLY:O	4:U:222:ARG:C	2.62	0.42
4:U:272:PRO:HD3	4:U:393:ASP:HB3	2.01	0.42
4:V:37:TRP:CZ3	4:V:72:LEU:HD21	2.55	0.42
4:V:396:GLN:H	4:V:396:GLN:NE2	2.17	0.42
5:W:619:LYS:O	5:W:619:LYS:HG2	2.19	0.42
6:X:310:LYS:O	6:X:399:ARG:HG3	2.19	0.42
6:X:377:ILE:HD11	6:X:436:LEU:CD1	2.50	0.42
6:X:656:PHE:HE2	6:X:752:PRO:HA	1.83	0.42
6:X:1024:PRO:O	6:X:1025:ASN:C	2.62	0.42
6:X:1031:LEU:C	6:X:1031:LEU:CD1	2.92	0.42
6:Y:218:GLU:HG3	6:Y:219:LEU:H	1.78	0.42
6:Y:284:ASP:O	6:Y:286:ALA:N	2.52	0.42
6:Y:484:THR:HG21	6:Y:529:TRP:CH2	2.55	0.42
1:f:57:ALA:HB1	1:f:58:PRO:CD	2.49	0.42
1:r:22:LEU:CB	1:r:83:VAL:HG21	2.50	0.42
1:t:23:THR:HG21	1:t:31:GLU:HB3	1.95	0.42
3:B:188:ALA:HA	3:B:191:LYS:HD2	2.01	0.42
3:B:365:ASP:HB2	3:B:451:SER:OG	2.20	0.42
3:B:556:SER:N	3:B:557:PRO:CD	2.82	0.42
3:D:73:PHE:CB	3:D:119:VAL:HG21	2.49	0.42
3:D:163:LEU:HD11	3:D:211:ALA:HB2	2.01	0.42
3:D:166:ALA:HB2	3:D:194:LEU:HD11	2.02	0.42
3:D:188:ALA:HA	3:D:191:LYS:HD2	2.01	0.42
3:D:288:SER:O	3:D:289:LEU:C	2.62	0.42
3:D:373:PHE:CD2	3:D:397:VAL:HG11	2.55	0.42
3:D:427:GLU:HB2	3:D:430:THR:CG2	2.49	0.42
3:D:590:VAL:O	3:D:592:THR:N	2.52	0.42
2:E:28:SER:C	2:E:29:THR:HG23	2.45	0.42
3:F:46:LYS:HD3	3:F:70:GLN:OE1	2.19	0.42
3:F:101:THR:HG23	3:F:102:GLY:N	2.34	0.42
3:F:152:ARG:NH1	3:F:155:ASP:OD2	2.53	0.42
3:F:517:LEU:HD13	3:F:517:LEU:C	2.44	0.42
7:G:101:MYR:C4	3:H:187:LEU:HD22	2.49	0.42
3:H:48:TRP:CE2	3:H:63:ILE:CD1	3.02	0.42
3:H:352:ASP:OD1	3:H:352:ASP:C	2.62	0.42
3:H:373:PHE:CD2	3:H:397:VAL:HG11	2.55	0.42
3:H:567:ARG:HB2	3:H:567:ARG:HH11	1.85	0.42
2:I:9:ASN:HD21	3:J:206:GLU:CG	2.32	0.42
3:J:252:LEU:HB3	3:J:613:LEU:HD23	2.02	0.42
3:J:412:ARG:CA	3:L:455:SER:O	2.66	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:J:418:ASN:CG	3:L:337:ARG:NH2	2.74	0.42
3:J:420:LEU:HD11	3:J:422:LEU:CD2	2.49	0.42
3:J:567:ARG:HB2	3:J:567:ARG:HH11	1.85	0.42
3:L:96:PHE:CD1	3:L:159:TRP:CH2	3.08	0.42
3:L:139:THR:HA	3:L:142:ILE:CG2	2.50	0.42
3:L:144:ASN:O	3:L:145:LEU:C	2.62	0.42
3:L:188:ALA:HA	3:L:191:LYS:HD2	2.01	0.42
3:L:565:LEU:N	3:L:565:LEU:HD23	2.34	0.42
2:M:28:SER:C	2:M:29:THR:HG23	2.45	0.42
3:N:154:ARG:O	3:N:155:ASP:C	2.62	0.42
3:N:293:SER:HB3	3:N:465:THR:HG23	2.01	0.42
3:N:341:MET:CG	3:N:431:ILE:HD11	2.49	0.42
3:P:128:ASP:OD2	3:T:107:VAL:HG21	2.18	0.42
3:P:152:ARG:NH1	3:P:155:ASP:OD2	2.53	0.42
3:P:171:THR:HG21	3:P:634:LYS:NZ	2.34	0.42
3:P:310:PHE:CD1	3:P:382:ILE:HG13	2.54	0.42
3:P:337:ARG:NH2	3:T:418:ASN:CG	2.74	0.42
3:P:357:ILE:CD1	3:P:414:GLN:HB3	2.50	0.42
3:P:365:ASP:HB2	3:P:451:SER:OG	2.20	0.42
3:P:578:SER:O	3:T:289:LEU:HA	2.20	0.42
2:Q:23:THR:C	2:Q:25:ASP:N	2.75	0.42
3:R:73:PHE:CB	3:R:119:VAL:HG21	2.49	0.42
3:R:293:SER:HB3	3:R:465:THR:HG23	2.01	0.42
3:R:370:VAL:O	3:R:370:VAL:CG2	2.67	0.42
3:R:469:VAL:O	3:R:469:VAL:HG22	2.20	0.42
3:R:517:LEU:HD13	3:R:517:LEU:C	2.44	0.42
3:R:618:THR:HG22	3:R:619:ASP:OD2	2.19	0.42
3:T:279:PRO:HA	3:T:484:ILE:O	2.20	0.42
4:U:53:VAL:HG11	4:U:61:VAL:HA	2.01	0.42
4:V:128:TYR:OH	4:V:139:ASN:HB2	2.20	0.42
4:V:175:MET:HE2	6:Y:404:MET:HE1	2.02	0.42
4:V:184:HIS:CD2	6:Y:436:LEU:CD1	2.95	0.42
4:V:196:LEU:HD23	6:Y:377:ILE:HD12	2.01	0.42
5:W:207:TYR:CZ	5:W:237:PRO:HB3	2.54	0.42
5:W:266:VAL:HG11	5:W:310:LEU:HD12	2.01	0.42
5:W:410:THR:O	5:W:411:VAL:C	2.62	0.42
5:W:530:GLU:O	5:W:531:PRO:C	2.61	0.42
5:W:651:ASN:HD21	5:W:685:ASN:ND2	2.17	0.42
5:W:918:ILE:N	5:W:919:PRO:CD	2.82	0.42
6:X:390:LEU:CA	6:X:1191:TYR:HD1	2.33	0.42
6:X:427:LEU:HD13	6:X:838:GLY:HA2	2.02	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
6:X:682:GLN:CG	6:X:683:PRO:HD2	2.40	0.42
6:X:792:GLN:HG2	6:X:931:ILE:CG2	2.46	0.42
6:X:1174:ASP:HA	6:X:1175:PRO:HD3	1.92	0.42
6:Y:156:ALA:HB3	6:Y:851:PHE:CD2	2.55	0.42
6:Y:551:PRO:O	6:Y:552:ASP:C	2.63	0.42
6:Y:564:TRP:C	6:Y:566:CYS:N	2.76	0.42
6:Y:684:LEU:HD11	6:Y:750:MET:CE	2.50	0.42
6:Y:864:PHE:CD1	6:Y:864:PHE:C	2.97	0.42
6:Y:882:PRO:C	6:Y:884:PRO:HD3	2.44	0.42
1:l:16:ALA:CB	1:l:22:LEU:CD2	2.98	0.42
1:b:24:LEU:HD11	1:b:80:ARG:CZ	2.49	0.42
1:f:5:MET:O	1:f:6:ILE:O	2.35	0.42
1:f:75:PRO:O	1:f:75:PRO:HD2	2.20	0.42
1:j:50:ALA:O	1:j:74:GLN:HB3	2.20	0.42
1:p:22:LEU:CD2	1:p:22:LEU:H	2.31	0.42
2:A:28:SER:C	2:A:29:THR:HG23	2.45	0.42
3:B:79:ARG:O	3:B:83:SER:HB2	2.20	0.42
3:B:252:LEU:HB3	3:B:613:LEU:HD23	2.02	0.42
3:B:285:PHE:O	3:B:286:THR:C	2.62	0.42
3:B:293:SER:HB3	3:B:465:THR:HG23	2.01	0.42
3:B:397:VAL:O	3:B:398:SER:HB2	2.18	0.42
7:C:101:MYR:H131	3:D:163:LEU:HD11	2.02	0.42
3:D:135:VAL:O	3:D:135:VAL:HG13	2.19	0.42
3:D:282:VAL:HG21	3:D:474:THR:HB	2.02	0.42
3:D:352:ASP:OD1	3:D:352:ASP:C	2.62	0.42
3:D:416:ARG:CB	3:F:298:PHE:HE1	2.16	0.42
3:D:625:ARG:HH11	3:D:625:ARG:HG3	1.85	0.42
3:F:166:ALA:HB2	3:F:194:LEU:HD11	2.02	0.42
3:F:199:LEU:HD11	3:F:203:TYR:CE2	2.54	0.42
3:F:433:GLY:O	3:F:435:PRO:CD	2.67	0.42
3:F:565:LEU:HD23	3:F:565:LEU:N	2.34	0.42
3:H:79:ARG:CB	3:H:79:ARG:HH11	2.33	0.42
3:H:193:ILE:O	3:H:194:LEU:C	2.61	0.42
3:J:79:ARG:CB	3:J:79:ARG:HH11	2.33	0.42
3:L:152:ARG:NH1	3:L:155:ASP:OD2	2.53	0.42
3:L:289:LEU:HA	3:N:578:SER:O	2.20	0.42
3:L:600:LEU:O	3:L:601:THR:C	2.60	0.42
3:N:46:LYS:HD3	3:N:70:GLN:OE1	2.19	0.42
3:N:79:ARG:O	3:N:83:SER:HB2	2.20	0.42
3:N:285:PHE:O	3:N:286:THR:C	2.62	0.42
3:P:48:TRP:CE2	3:P:63:ILE:CD1	3.02	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:P:58:ILE:HG12	3:P:127:LEU:CB	2.50	0.42
3:P:59:ASP:OD2	5:W:349:TRP:NE1	2.53	0.42
3:P:158:LEU:O	3:P:161:LYS:HB3	2.20	0.42
3:P:285:PHE:O	3:P:286:THR:C	2.62	0.42
3:R:48:TRP:CE2	3:R:63:ILE:CD1	3.02	0.42
3:R:214:ARG:HB2	3:R:215:TYR:CE2	2.55	0.42
3:R:470:LEU:O	3:R:470:LEU:HD13	2.20	0.42
3:R:502:LEU:O	3:R:503:ALA:C	2.63	0.42
3:R:616:GLY:O	3:R:620:ALA:HB2	2.20	0.42
7:S:101:MYR:C2	3:T:191:LYS:HZ1	2.27	0.42
3:T:152:ARG:NH1	3:T:155:ASP:OD2	2.53	0.42
3:T:156:VAL:HG13	3:T:157:ALA:N	2.35	0.42
3:T:199:LEU:HD11	3:T:203:TYR:CE2	2.55	0.42
3:T:252:LEU:HB3	3:T:613:LEU:HD23	2.02	0.42
3:T:288:SER:O	3:T:289:LEU:C	2.62	0.42
3:T:345:ILE:O	3:T:347:SER:N	2.52	0.42
4:V:172:ALA:HA	4:V:175:MET:HE3	2.02	0.42
5:W:157:GLN:O	6:X:683:PRO:HG3	2.19	0.42
5:W:582:LEU:HB2	5:W:602:MET:HE1	2.01	0.42
5:W:1169:LYS:O	5:W:1170:ASP:HB2	2.20	0.42
6:X:207:LEU:HB3	6:X:244:VAL:HG12	2.01	0.42
6:X:476:LEU:O	6:X:476:LEU:HD22	2.20	0.42
6:X:861:ARG:O	6:X:861:ARG:HG2	2.19	0.42
6:X:996:MET:HE2	6:X:996:MET:CA	2.41	0.42
6:X:1037:VAL:O	6:X:1038:PRO:C	2.63	0.42
6:Y:595:GLN:N	6:Y:596:PRO:HD2	2.35	0.42
6:Y:1156:THR:HG21	6:Y:1161:ASN:CB	2.50	0.42
6:Y:1183:ASP:O	6:Y:1185:PRO:HD2	2.20	0.42
6:Y:1192:ASN:HB3	6:Y:1193:SER:H	1.57	0.42
1:d:22:LEU:O	1:d:42:THR:OG1	2.36	0.42
3:B:106:ILE:CD1	3:B:152:ARG:HG3	2.44	0.42
3:B:152:ARG:NH1	3:B:155:ASP:OD2	2.53	0.42
3:B:266:MET:HG2	3:B:520:VAL:HG11	2.01	0.42
3:B:456:ASN:C	3:B:456:ASN:ND2	2.76	0.42
3:D:289:LEU:HA	3:F:578:SER:O	2.20	0.42
3:D:365:ASP:HB2	3:D:451:SER:OG	2.20	0.42
3:D:418:ASN:HD22	3:D:418:ASN:HA	1.51	0.42
3:D:551:GLN:NE2	3:D:558:THR:HB	2.34	0.42
3:D:600:LEU:HG	3:D:601:THR:N	2.30	0.42
3:D:618:THR:HG22	3:D:619:ASP:OD2	2.19	0.42
3:F:79:ARG:O	3:F:83:SER:HB2	2.20	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:F:171:THR:HG21	3:F:634:LYS:HZ1	1.84	0.42
3:F:252:LEU:HB3	3:F:613:LEU:HD23	2.02	0.42
3:F:288:SER:O	3:F:289:LEU:C	2.62	0.42
3:F:600:LEU:O	3:F:601:THR:C	2.60	0.42
3:F:609:ARG:CD	2:G:29:THR:HG21	2.47	0.42
3:F:625:ARG:HH11	3:F:625:ARG:HG3	1.84	0.42
3:H:95:LEU:HD12	3:H:95:LEU:HA	1.71	0.42
3:H:152:ARG:NH1	3:H:155:ASP:OD2	2.53	0.42
3:H:188:ALA:HA	3:H:191:LYS:HD2	2.02	0.42
3:H:282:VAL:HG21	3:H:474:THR:HB	2.02	0.42
3:H:292:ARG:HG2	3:H:292:ARG:NH1	2.32	0.42
3:H:443:PHE:HE2	3:H:459:LEU:HD21	1.75	0.42
2:I:29:THR:HA	3:N:609:ARG:O	2.19	0.42
3:J:214:ARG:HB2	3:J:215:TYR:CE2	2.55	0.42
3:J:318:LYS:HG3	3:J:319:THR:H	1.84	0.42
3:J:352:ASP:OD1	3:J:352:ASP:C	2.62	0.42
3:J:622:GLU:OE1	3:N:281:PRO:CD	2.62	0.42
2:K:27:THR:HG22	2:K:29:THR:HG23	2.02	0.42
3:L:48:TRP:CE2	3:L:63:ILE:CD1	3.02	0.42
3:L:166:ALA:HB2	3:L:194:LEU:HD11	2.02	0.42
3:L:288:SER:O	3:L:289:LEU:C	2.62	0.42
3:L:343:ALA:CB	3:L:478:GLY:O	2.62	0.42
3:L:431:ILE:CG1	3:L:437:ILE:HD13	2.50	0.42
3:L:502:LEU:O	3:L:503:ALA:C	2.63	0.42
3:N:79:ARG:CB	3:N:79:ARG:HH11	2.33	0.42
3:N:143:MET:O	3:N:146:GLN:HB3	2.19	0.42
3:N:567:ARG:HB2	3:N:567:ARG:HH11	1.85	0.42
3:P:135:VAL:O	3:P:135:VAL:HG13	2.19	0.42
3:P:166:ALA:HB2	3:P:194:LEU:HD11	2.02	0.42
3:P:409:ILE:HD11	3:R:297:ALA:O	2.17	0.42
3:P:556:SER:N	3:P:557:PRO:CD	2.82	0.42
3:P:567:ARG:HB2	3:P:567:ARG:HH11	1.85	0.42
3:R:341:MET:CG	3:R:431:ILE:HD11	2.49	0.42
3:R:365:ASP:HB2	3:R:451:SER:OG	2.20	0.42
3:R:431:ILE:CG1	3:R:437:ILE:HD13	2.50	0.42
3:T:357:ILE:CD1	3:T:414:GLN:HB3	2.50	0.42
4:U:12:ASN:ND2	4:U:116:GLU:OE2	2.52	0.42
4:U:77:ASN:N	4:U:78:PRO:CD	2.82	0.42
4:U:329:ASN:CB	4:U:332:PRO:HD2	2.50	0.42
4:V:162:PRO:HD2	4:V:210:TRP:CZ2	2.55	0.42
4:V:188:THR:HG23	6:Y:376:LEU:CD1	2.50	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
5:W:231:ASP:O	5:W:232:SER:C	2.62	0.42
5:W:300:PHE:CD2	5:W:346:PRO:HG3	2.55	0.42
5:W:957:ARG:HG2	5:W:964:VAL:HG22	2.01	0.42
6:X:205:MET:HB3	6:X:249:TRP:CG	2.54	0.42
6:X:249:TRP:CG	6:X:250:ALA:N	2.88	0.42
6:X:291:LEU:HD13	6:X:291:LEU:HA	1.85	0.42
6:X:389:GLU:O	6:X:389:GLU:HG3	2.20	0.42
6:X:589:LEU:C	6:X:591:ASN:N	2.76	0.42
6:X:633:PRO:O	6:X:635:GLN:N	2.53	0.42
6:X:645:HIS:CE1	6:X:700:TRP:CZ3	3.08	0.42
6:X:722:ASN:CG	6:X:723:ASP:H	2.27	0.42
6:X:780:ILE:HD13	6:X:940:GLN:CG	2.49	0.42
6:X:793:ARG:HB3	6:Y:693:TYR:CD2	2.55	0.42
6:X:1125:TYR:N	6:X:1125:TYR:CD1	2.76	0.42
6:Y:113:ASN:N	6:Y:113:ASN:HD22	2.18	0.42
6:Y:122:CYS:SG	6:Y:135:HIS:NE2	2.93	0.42
6:Y:367:ARG:HH11	6:Y:367:ARG:CG	2.32	0.42
6:Y:545:LEU:CD2	6:Y:576:THR:HG21	2.50	0.42
6:Y:590:ALA:HA	6:Y:639:LEU:HD11	2.01	0.42
6:Y:938:TYR:CD1	6:Y:938:TYR:C	2.98	0.42
6:Y:979:ARG:HG2	6:Y:979:ARG:NH1	2.32	0.42
6:Y:1056:VAL:HG21	6:Y:1107:ILE:HB	2.02	0.42
1:h:59:HIS:CE1	3:H:397:VAL:O	2.66	0.42
1:p:22:LEU:CD1	1:p:42:THR:O	2.62	0.42
1:p:59:HIS:CD2	3:P:424:ALA:HA	2.55	0.42
2:A:26:MET:CG	3:B:216:PRO:CG	2.95	0.42
2:A:28:SER:O	2:A:29:THR:CG2	2.68	0.42
3:B:156:VAL:HG13	3:B:157:ALA:N	2.35	0.42
3:B:279:PRO:HA	3:B:484:ILE:O	2.20	0.42
3:B:420:LEU:HD11	3:B:422:LEU:CD2	2.49	0.42
3:B:470:LEU:O	3:B:470:LEU:HD13	2.20	0.42
3:B:508:VAL:O	3:B:508:VAL:HG12	2.20	0.42
3:B:517:LEU:HD13	3:B:517:LEU:C	2.44	0.42
3:D:318:LYS:HG3	3:D:319:THR:H	1.84	0.42
3:D:502:LEU:O	3:D:503:ALA:C	2.63	0.42
3:D:578:SER:O	3:H:289:LEU:HA	2.20	0.42
3:D:609:ARG:CD	2:E:29:THR:HG21	2.47	0.42
2:E:27:THR:HG22	2:E:29:THR:HG23	2.02	0.42
3:F:136:PHE:CD1	3:F:136:PHE:N	2.87	0.42
3:F:156:VAL:HG13	3:F:157:ALA:N	2.35	0.42
3:F:203:TYR:N	3:F:204:PRO:HD2	2.34	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:F:214:ARG:HB2	3:F:215:TYR:CE2	2.55	0.42
3:F:274:ILE:O	3:F:274:ILE:HG13	2.20	0.42
3:F:502:LEU:O	3:F:503:ALA:C	2.63	0.42
3:F:616:GLY:O	3:F:620:ALA:HB2	2.20	0.42
2:G:9:ASN:CB	3:H:209:ASN:ND2	2.83	0.42
2:G:26:MET:CG	3:H:216:PRO:CG	2.95	0.42
2:G:28:SER:C	2:G:29:THR:HG23	2.45	0.42
3:H:154:ARG:O	3:H:155:ASP:C	2.62	0.42
3:H:166:ALA:HB2	3:H:194:LEU:HD11	2.02	0.42
3:H:365:ASP:HB2	3:H:451:SER:OG	2.20	0.42
3:H:470:LEU:HD13	3:H:470:LEU:O	2.20	0.42
3:J:143:MET:O	3:J:146:GLN:HB3	2.19	0.42
3:J:273:LEU:HA	3:L:630:LEU:CD1	2.50	0.42
3:J:355:GLY:HA2	3:L:298:PHE:CE1	2.47	0.42
3:L:93:GLN:OE1	3:L:93:GLN:HA	2.19	0.42
3:L:101:THR:HG23	3:L:102:GLY:N	2.34	0.42
3:L:167:MET:HA	3:L:167:MET:CE	2.31	0.42
3:L:561:ASP:C	3:L:563:PRO:HD2	2.45	0.42
2:M:9:ASN:CB	3:N:209:ASN:ND2	2.83	0.42
3:N:96:PHE:CD1	3:N:159:TRP:CH2	3.08	0.42
3:N:166:ALA:HB2	3:N:194:LEU:HD11	2.02	0.42
3:N:360:TYR:CE1	3:N:415:PRO:HG3	2.55	0.42
3:N:460:MET:C	3:N:462:GLY:N	2.78	0.42
7:O:101:MYR:H131	3:P:163:LEU:HD11	2.02	0.42
3:P:46:LYS:HD3	3:P:70:GLN:OE1	2.19	0.42
3:P:370:VAL:O	3:P:370:VAL:CG2	2.67	0.42
3:P:431:ILE:CG1	3:P:437:ILE:HD13	2.50	0.42
3:P:547:HIS:ND1	3:P:547:HIS:C	2.78	0.42
3:P:609:ARG:CD	2:Q:29:THR:CG2	2.93	0.42
3:P:625:ARG:HG3	3:P:625:ARG:HH11	1.84	0.42
3:R:79:ARG:CB	3:R:79:ARG:HH11	2.33	0.42
3:R:96:PHE:CD1	3:R:159:TRP:CH2	3.08	0.42
3:R:271:LEU:HD12	3:R:271:LEU:HA	1.79	0.42
3:R:318:LYS:HG3	3:R:319:THR:H	1.84	0.42
2:S:28:SER:O	2:S:29:THR:CG2	2.68	0.42
7:S:101:MYR:C4	3:T:187:LEU:HD22	2.49	0.42
3:T:292:ARG:HG2	3:T:292:ARG:NH1	2.32	0.42
3:T:427:GLU:HB2	3:T:430:THR:CG2	2.48	0.42
3:T:463:LEU:C	3:T:463:LEU:HD23	2.44	0.42
3:T:618:THR:HG22	3:T:619:ASP:OD2	2.19	0.42
4:U:31:GLY:C	4:U:33:TYR:H	2.28	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
4:U:64:ARG:HG2	4:U:64:ARG:NH1	2.33	0.42
4:U:146:THR:O	4:U:146:THR:HG22	2.20	0.42
4:U:182:ILE:CG2	4:U:246:ALA:HB3	2.50	0.42
4:U:239:PRO:CA	4:U:243:PRO:HB3	2.50	0.42
5:W:263:VAL:HG23	5:W:303:LEU:HD11	2.00	0.42
5:W:346:PRO:O	5:W:347:GLN:C	2.61	0.42
5:W:450:THR:O	5:W:671:THR:HA	2.20	0.42
5:W:533:ILE:CG1	5:W:537:ILE:HD11	2.49	0.42
5:W:574:THR:HB	5:W:611:THR:CA	2.50	0.42
5:W:1208:CYS:HA	5:W:1219:ILE:O	2.20	0.42
5:W:1250:LEU:HD12	5:W:1258:MET:HE2	2.00	0.42
6:X:657:GLY:HA2	6:X:673:ASN:O	2.19	0.42
6:X:687:LEU:HB3	6:X:688:THR:H	1.50	0.42
6:X:871:VAL:O	6:X:875:GLN:HB2	2.20	0.42
6:X:1182:ASN:ND2	6:X:1184:LEU:HD22	2.33	0.42
6:Y:277:VAL:HG12	6:Y:278:LEU:CG	2.40	0.42
6:Y:330:ILE:HD12	6:Y:340:LEU:HD11	2.01	0.42
6:Y:366:ASP:OD1	6:Y:366:ASP:C	2.63	0.42
6:Y:1032:ILE:HD12	6:Y:1032:ILE:HA	1.84	0.42
1:l:59:HIS:CD2	3:L:397:VAL:CG2	2.82	0.41
1:f:8:GLN:OE1	3:H:302:GLU:O	2.37	0.41
1:h:80:ARG:NH1	1:h:84:GLU:OE2	2.49	0.41
1:j:23:THR:HG21	3:J:324:GLY:CA	2.43	0.41
1:p:21:ARG:CD	3:P:370:VAL:HG11	2.50	0.41
3:B:46:LYS:HD3	3:B:70:GLN:OE1	2.19	0.41
3:B:288:SER:O	3:B:289:LEU:C	2.62	0.41
3:B:442:THR:OG1	3:B:443:PHE:N	2.50	0.41
3:B:618:THR:HG22	3:B:619:ASP:OD2	2.19	0.41
3:D:50:PRO:HB3	3:D:56:ALA:CB	2.50	0.41
3:D:79:ARG:CB	3:D:79:ARG:HH11	2.33	0.41
3:D:203:TYR:N	3:D:204:PRO:HD2	2.34	0.41
2:E:23:THR:C	2:E:25:ASP:N	2.75	0.41
3:F:292:ARG:HG2	3:F:292:ARG:NH1	2.32	0.41
3:F:312:ASN:ND2	3:F:314:SER:O	2.53	0.41
3:F:318:LYS:HG3	3:F:319:THR:H	1.84	0.41
3:F:561:ASP:C	3:F:563:PRO:HD2	2.45	0.41
3:H:156:VAL:HG13	3:H:157:ALA:N	2.35	0.41
3:H:332:GLY:HA3	3:H:456:ASN:HA	2.01	0.41
3:J:152:ARG:NH1	3:J:155:ASP:OD2	2.53	0.41
3:J:282:VAL:HG21	3:J:474:THR:HB	2.02	0.41
3:J:361:ASP:OD2	3:J:362:THR:HG23	2.20	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:J:370:VAL:O	3:J:370:VAL:CG2	2.67	0.41
3:J:517:LEU:HD13	3:J:517:LEU:C	2.44	0.41
2:K:9:ASN:CB	3:L:209:ASN:ND2	2.83	0.41
3:L:199:LEU:HD11	3:L:203:TYR:CE2	2.55	0.41
3:L:460:MET:C	3:L:462:GLY:N	2.78	0.41
2:M:28:SER:O	2:M:29:THR:CG2	2.68	0.41
3:N:279:PRO:HA	3:N:484:ILE:O	2.20	0.41
3:P:147:ARG:CB	3:P:147:ARG:CZ	2.98	0.41
3:P:360:TYR:CE1	3:P:415:PRO:HG3	2.55	0.41
3:P:468:PRO:O	3:R:578:SER:OG	2.21	0.41
3:R:188:ALA:HA	3:R:191:LYS:HD2	2.01	0.41
3:R:358:LEU:HB2	3:R:417:PHE:HD1	1.77	0.41
3:R:561:ASP:C	3:R:563:PRO:HD2	2.45	0.41
3:R:609:ARG:CD	2:S:29:THR:HG21	2.47	0.41
3:R:625:ARG:HH11	3:R:625:ARG:HG3	1.84	0.41
3:T:214:ARG:HB2	3:T:215:TYR:CE2	2.55	0.41
3:T:360:TYR:CE1	3:T:415:PRO:HG3	2.55	0.41
3:T:502:LEU:O	3:T:503:ALA:C	2.63	0.41
4:U:103:VAL:O	4:U:104:PRO:O	2.37	0.41
4:U:143:HIS:CD2	4:U:292:ALA:HA	2.55	0.41
4:U:379:ILE:C	4:U:381:ALA:H	2.27	0.41
5:W:121:ALA:HB2	5:W:134:PHE:HB3	2.02	0.41
5:W:248:LEU:HD11	6:X:647:ARG:HE	1.84	0.41
5:W:293:THR:CG2	5:W:294:SER:N	2.83	0.41
5:W:519:LEU:CD1	5:W:566:LEU:HG	2.48	0.41
5:W:832:LEU:HD23	5:W:893:ALA:HA	2.02	0.41
5:W:1249:HIS:CE1	5:W:1297:VAL:HB	2.55	0.41
6:X:493:LEU:O	6:X:494:THR:C	2.63	0.41
6:X:593:LEU:O	6:X:596:PRO:HD2	2.20	0.41
6:X:690:ALA:O	6:X:691:PRO:C	2.60	0.41
6:X:1131:ARG:O	6:Y:106:MET:HA	2.20	0.41
6:Y:986:GLN:HB2	6:Y:1135:TYR:HE1	1.84	0.41
6:Y:1005:VAL:HG22	6:Y:1006:ARG:N	2.35	0.41
6:Y:1112:ILE:HG23	6:Y:1113:PRO:HD2	2.02	0.41
6:Y:1154:ILE:HD13	6:Y:1154:ILE:HA	1.81	0.41
1:l:17:ALA:CB	1:l:79:ILE:HG23	2.51	0.41
1:b:5:MET:O	1:b:5:MET:SD	2.78	0.41
1:t:44:GLY:C	3:T:399:ALA:O	2.63	0.41
3:B:166:ALA:HB2	3:B:194:LEU:HD11	2.02	0.41
3:D:214:ARG:HB2	3:D:215:TYR:CE2	2.55	0.41
3:D:285:PHE:O	3:D:286:THR:C	2.62	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:D:357:ILE:HD13	3:D:357:ILE:HA	1.97	0.41
3:F:273:LEU:HA	3:H:630:LEU:CD1	2.50	0.41
3:F:285:PHE:O	3:F:286:THR:C	2.62	0.41
3:F:471:LEU:O	3:F:472:ARG:C	2.64	0.41
3:F:554:GLN:C	3:F:556:SER:H	2.28	0.41
3:H:79:ARG:O	3:H:83:SER:HB2	2.20	0.41
3:H:144:ASN:O	3:H:145:LEU:C	2.62	0.41
2:I:28:SER:O	2:I:29:THR:CG2	2.68	0.41
2:I:28:SER:C	2:I:29:THR:HG23	2.45	0.41
3:J:101:THR:HG21	3:J:159:TRP:CZ2	2.55	0.41
3:J:266:MET:HE2	3:J:604:VAL:HG12	2.03	0.41
3:J:289:LEU:HD13	3:L:580:PRO:CD	2.50	0.41
3:J:365:ASP:HB2	3:J:451:SER:OG	2.20	0.41
3:J:373:PHE:CD2	3:J:397:VAL:HG11	2.55	0.41
3:J:401:GLY:O	3:J:402:ALA:CB	2.66	0.41
3:L:79:ARG:O	3:L:83:SER:HB2	2.20	0.41
3:L:289:LEU:HD13	3:N:580:PRO:CD	2.50	0.41
3:L:360:TYR:HD1	3:L:360:TYR:C	2.24	0.41
3:N:62:ALA:O	3:N:63:ILE:C	2.62	0.41
3:N:147:ARG:CB	3:N:147:ARG:CZ	2.99	0.41
3:N:156:VAL:O	3:N:157:ALA:C	2.64	0.41
3:N:156:VAL:HG13	3:N:157:ALA:N	2.35	0.41
3:N:344:MET:HE2	3:N:502:LEU:CD1	2.37	0.41
3:N:357:ILE:CD1	3:N:414:GLN:HB3	2.50	0.41
3:N:370:VAL:O	3:N:370:VAL:CG2	2.67	0.41
3:N:469:VAL:O	3:N:469:VAL:HG22	2.20	0.41
3:N:471:LEU:O	3:N:472:ARG:C	2.63	0.41
3:N:502:LEU:O	3:N:503:ALA:C	2.63	0.41
3:N:508:VAL:O	3:N:508:VAL:HG12	2.20	0.41
2:O:9:ASN:HD21	3:P:206:GLU:CG	2.32	0.41
3:P:156:VAL:HG13	3:P:157:ALA:N	2.35	0.41
3:P:163:LEU:HD11	3:P:211:ALA:HB2	2.01	0.41
3:P:289:LEU:HD13	3:R:580:PRO:CD	2.50	0.41
3:P:549:LEU:HB3	3:R:644:LEU:CD1	2.34	0.41
3:R:46:LYS:HD3	3:R:70:GLN:OE1	2.19	0.41
3:R:93:GLN:OE1	3:R:93:GLN:HA	2.19	0.41
3:R:289:LEU:HD13	3:T:580:PRO:CD	2.50	0.41
3:R:361:ASP:OD2	3:R:362:THR:HG23	2.20	0.41
7:S:101:MYR:C2	3:T:191:LYS:HZ3	2.11	0.41
3:T:361:ASP:OD2	3:T:362:THR:HG23	2.20	0.41
3:T:469:VAL:O	3:T:469:VAL:HG22	2.20	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:T:518:ALA:HB1	3:T:580:PRO:HG3	2.03	0.41
4:U:159:LEU:HD12	4:U:267:ILE:HD12	2.01	0.41
4:U:184:HIS:CD2	4:U:184:HIS:N	2.86	0.41
4:U:205:VAL:O	4:U:209:ARG:HB2	2.19	0.41
4:U:272:PRO:HD3	4:U:393:ASP:CB	2.49	0.41
4:V:6:PHE:O	4:V:122:SER:HB2	2.20	0.41
4:V:291:THR:HG22	4:V:291:THR:O	2.20	0.41
5:W:47:LEU:HB3	5:W:191:PRO:HG3	2.03	0.41
5:W:280:THR:HG22	5:W:281:ALA:N	2.35	0.41
5:W:519:LEU:HD11	5:W:580:ALA:CB	2.50	0.41
5:W:640:THR:HB	5:W:661:GLY:H	1.85	0.41
5:W:714:ARG:HA	5:W:758:THR:HA	2.02	0.41
5:W:821:GLU:CA	5:W:824:ARG:HH21	2.32	0.41
5:W:825:ASN:OD1	5:W:825:ASN:C	2.63	0.41
5:W:961:GLY:O	5:W:962:GLU:C	2.63	0.41
5:W:1199:ALA:HB2	5:W:1296:TYR:CZ	2.56	0.41
5:W:1273:LEU:HD12	5:W:1285:PRO:HB2	2.02	0.41
6:X:200:LEU:CD2	6:X:200:LEU:H	2.33	0.41
6:X:240:VAL:HG11	6:X:1087:TYR:CZ	2.56	0.41
6:X:282:TYR:HE2	6:X:289:ASN:ND2	2.17	0.41
6:X:362:THR:HG21	6:X:1173:VAL:HB	2.01	0.41
6:X:377:ILE:CG1	6:X:438:PRO:HG3	2.50	0.41
6:X:448:SER:HB2	6:X:664:GLY:HA2	2.01	0.41
6:X:567:LEU:HD11	6:X:705:ILE:CD1	2.50	0.41
6:X:587:MET:HA	6:X:590:ALA:HB3	2.02	0.41
6:X:593:LEU:CD2	6:X:639:LEU:HD13	2.51	0.41
6:X:693:TYR:HD1	6:X:693:TYR:N	2.18	0.41
6:X:798:VAL:HG23	6:X:803:THR:HG21	2.01	0.41
6:Y:220:ILE:O	6:Y:350:LEU:HD11	2.20	0.41
6:Y:291:LEU:O	6:Y:890:GLN:HA	2.19	0.41
6:Y:376:LEU:HB2	6:Y:383:ALA:HB3	2.01	0.41
6:Y:526:ALA:O	6:Y:530:LYS:N	2.50	0.41
6:Y:576:THR:HG22	6:Y:813:VAL:HG11	2.02	0.41
6:Y:592:LEU:HD23	6:Y:592:LEU:HA	1.76	0.41
1:f:6:ILE:HG21	3:H:494:THR:HG22	2.03	0.41
7:A:101:MYR:H131	3:B:163:LEU:HD11	2.02	0.41
3:B:147:ARG:CB	3:B:147:ARG:CZ	2.98	0.41
3:B:373:PHE:CD2	3:B:397:VAL:HG11	2.55	0.41
3:B:431:ILE:CG1	3:B:437:ILE:HD13	2.50	0.41
3:B:469:VAL:O	3:B:469:VAL:HG22	2.20	0.41
3:B:554:GLN:C	3:B:556:SER:H	2.28	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:B:576:PHE:HB2	3:B:591:LEU:HD21	2.02	0.41
3:B:625:ARG:HH11	3:B:625:ARG:HG3	1.84	0.41
2:C:28:SER:C	2:C:29:THR:HG23	2.45	0.41
7:C:101:MYR:H62	3:D:200:CYS:O	2.21	0.41
3:D:79:ARG:O	3:D:83:SER:HB2	2.20	0.41
3:D:199:LEU:HD11	3:D:203:TYR:CE2	2.54	0.41
3:D:266:MET:HE2	3:D:604:VAL:HG12	2.03	0.41
3:D:357:ILE:CD1	3:D:414:GLN:HB3	2.50	0.41
3:D:428:ARG:NH1	3:D:428:ARG:CG	2.76	0.41
3:D:580:PRO:CD	3:H:289:LEU:HD13	2.51	0.41
2:E:28:SER:O	2:E:29:THR:CG2	2.68	0.41
3:F:96:PHE:CE1	3:F:159:TRP:HH2	2.36	0.41
2:G:28:SER:O	2:G:29:THR:CG2	2.68	0.41
3:H:171:THR:HG21	3:H:634:LYS:NZ	2.34	0.41
3:H:274:ILE:O	3:H:274:ILE:HG13	2.20	0.41
3:H:288:SER:O	3:H:289:LEU:C	2.63	0.41
3:H:381:PRO:C	3:H:382:ILE:O	2.63	0.41
3:H:556:SER:N	3:H:557:PRO:CD	2.82	0.41
2:I:36:LEU:HD11	3:N:613:LEU:HD21	2.02	0.41
7:I:101:MYR:H62	3:J:200:CYS:O	2.21	0.41
3:J:52:GLY:HA2	4:V:397:ALA:HB3	2.02	0.41
3:J:360:TYR:CE1	3:J:415:PRO:HG3	2.55	0.41
3:J:431:ILE:CG1	3:J:437:ILE:HD13	2.50	0.41
3:L:154:ARG:O	3:L:155:ASP:C	2.62	0.41
3:L:360:TYR:CE1	3:L:415:PRO:HG3	2.56	0.41
3:L:361:ASP:OD2	3:L:362:THR:HG23	2.20	0.41
3:L:397:VAL:O	3:L:398:SER:HB2	2.18	0.41
3:N:282:VAL:HG21	3:N:474:THR:HB	2.02	0.41
3:N:288:SER:O	3:N:289:LEU:C	2.62	0.41
3:P:79:ARG:O	3:P:83:SER:HB2	2.20	0.41
3:P:96:PHE:CD1	3:P:159:TRP:CH2	3.08	0.41
3:P:214:ARG:HB2	3:P:215:TYR:CE2	2.55	0.41
3:P:361:ASP:OD2	3:P:362:THR:HG23	2.20	0.41
3:P:470:LEU:HD13	3:P:470:LEU:O	2.20	0.41
3:P:554:GLN:C	3:P:556:SER:H	2.28	0.41
3:P:634:LYS:HE3	3:P:634:LYS:HB2	1.89	0.41
2:Q:27:THR:HG22	2:Q:29:THR:HG23	2.02	0.41
2:Q:28:SER:O	2:Q:29:THR:CG2	2.68	0.41
3:R:273:LEU:HA	3:T:630:LEU:CD1	2.50	0.41
3:R:285:PHE:O	3:R:286:THR:C	2.63	0.41
3:R:289:LEU:HA	3:T:578:SER:O	2.20	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:R:360:TYR:HD1	3:R:360:TYR:C	2.24	0.41
3:R:420:LEU:HD11	3:R:422:LEU:CD2	2.49	0.41
3:R:603:GLY:C	3:R:605:GLY:N	2.75	0.41
2:S:9:ASN:CB	3:T:209:ASN:ND2	2.83	0.41
3:T:48:TRP:CZ3	3:T:123:ALA:N	2.89	0.41
3:T:466:VAL:HG11	3:T:502:LEU:CD2	2.50	0.41
3:T:576:PHE:HB2	3:T:591:LEU:HD21	2.02	0.41
5:W:126:LEU:C	5:W:128:ALA:N	2.78	0.41
5:W:240:LEU:HD22	5:W:289:TYR:CZ	2.55	0.41
5:W:248:LEU:HD21	6:X:647:ARG:NH2	2.35	0.41
5:W:646:LYS:NZ	5:W:648:THR:HG22	2.35	0.41
5:W:993:PRO:HG3	5:W:1017:ILE:HG21	2.01	0.41
5:W:1042:THR:HA	5:W:1043:PRO:HD3	1.87	0.41
6:X:303:GLN:HE21	6:X:303:GLN:HB2	1.47	0.41
6:X:352:THR:O	6:X:354:THR:N	2.48	0.41
6:X:389:GLU:OE2	6:X:1194:LEU:HD21	2.21	0.41
6:X:490:PRO:O	6:X:491:THR:C	2.63	0.41
6:X:530:LYS:O	6:X:534:SER:N	2.48	0.41
6:X:755:LEU:HD13	6:X:755:LEU:HA	1.95	0.41
6:X:835:HIS:NE2	6:X:839:LYS:NZ	2.69	0.41
6:Y:219:LEU:CD2	6:Y:350:LEU:HG	2.50	0.41
6:Y:355:LYS:HD2	6:Y:1149:THR:O	2.20	0.41
6:Y:399:ARG:C	6:Y:401:THR:H	2.28	0.41
6:Y:414:THR:HG23	6:Y:415:GLN:N	2.35	0.41
6:Y:468:ALA:O	6:Y:471:PRO:HD2	2.20	0.41
6:Y:483:VAL:HG11	6:Y:841:PHE:HE1	1.82	0.41
6:Y:993:TRP:HE1	6:Y:995:PRO:HG3	1.85	0.41
6:Y:1211:ARG:NH1	6:Y:1211:ARG:CG	2.83	0.41
3:B:235:LEU:HA	3:B:235:LEU:HD13	1.80	0.41
3:B:466:VAL:HG11	3:B:502:LEU:CD2	2.51	0.41
3:B:561:ASP:C	3:B:563:PRO:HD2	2.45	0.41
3:B:590:VAL:O	3:B:592:THR:N	2.52	0.41
3:D:105:PRO:HB3	3:F:125:ASP:HB3	2.03	0.41
3:D:252:LEU:HB3	3:D:613:LEU:HD23	2.02	0.41
3:D:312:ASN:ND2	3:D:314:SER:O	2.54	0.41
3:D:470:LEU:HD13	3:D:470:LEU:O	2.20	0.41
3:D:508:VAL:O	3:D:508:VAL:HG12	2.20	0.41
3:D:567:ARG:HB2	3:D:567:ARG:HH11	1.85	0.41
3:F:147:ARG:CB	3:F:147:ARG:CZ	2.98	0.41
3:F:370:VAL:O	3:F:370:VAL:CG2	2.67	0.41
3:H:293:SER:HB3	3:H:465:THR:HG23	2.01	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:H:361:ASP:OD2	3:H:362:THR:HG23	2.20	0.41
3:H:485:VAL:O	3:H:486:PRO:C	2.64	0.41
3:H:554:GLN:C	3:H:556:SER:H	2.28	0.41
3:H:561:ASP:C	3:H:563:PRO:HD2	2.45	0.41
3:H:576:PHE:HB2	3:H:591:LEU:HD21	2.03	0.41
3:H:616:GLY:O	3:H:620:ALA:HB2	2.20	0.41
3:J:62:ALA:O	3:J:63:ILE:C	2.62	0.41
3:J:163:LEU:HD11	3:J:211:ALA:HB2	2.01	0.41
3:J:336:THR:CG2	3:J:456:ASN:ND2	2.79	0.41
3:J:508:VAL:O	3:J:508:VAL:HG12	2.20	0.41
3:J:554:GLN:C	3:J:556:SER:H	2.28	0.41
3:J:561:ASP:C	3:J:563:PRO:HD2	2.45	0.41
3:J:578:SER:O	3:N:289:LEU:HA	2.20	0.41
2:K:26:MET:CG	3:L:216:PRO:HG3	2.29	0.41
3:L:105:PRO:HB3	3:N:125:ASP:HB3	2.03	0.41
3:L:143:MET:O	3:L:146:GLN:HB3	2.20	0.41
3:L:256:SER:OG	3:L:257:SER:N	2.51	0.41
3:L:266:MET:HE2	3:L:604:VAL:HG12	2.03	0.41
3:L:373:PHE:CD2	3:L:397:VAL:HG11	2.55	0.41
3:N:93:GLN:OE1	3:N:93:GLN:HA	2.19	0.41
3:N:161:LYS:HE3	3:N:165:THR:CG2	2.50	0.41
3:N:162:HIS:CD2	3:N:199:LEU:HB2	2.54	0.41
3:N:361:ASP:OD2	3:N:362:THR:HG23	2.20	0.41
3:N:456:ASN:HA	3:N:457:PRO:HD2	1.93	0.41
3:P:518:ALA:HB1	3:P:580:PRO:HG3	2.03	0.41
3:R:44:THR:CG2	3:R:45:GLY:N	2.82	0.41
3:R:105:PRO:HB3	3:T:125:ASP:HB3	2.03	0.41
3:R:282:VAL:HG21	3:R:474:THR:HB	2.02	0.41
3:R:357:ILE:CD1	3:R:414:GLN:HB3	2.50	0.41
3:R:609:ARG:CD	2:S:29:THR:CG2	2.93	0.41
3:T:431:ILE:CG1	3:T:437:ILE:HD13	2.50	0.41
4:U:118:VAL:O	4:U:118:VAL:HG13	2.20	0.41
4:U:196:LEU:HD11	6:X:672:SER:CB	2.51	0.41
4:U:302:GLU:CG	4:U:310:GLN:HE21	2.32	0.41
4:U:329:ASN:OD1	4:U:332:PRO:HD2	2.20	0.41
4:U:374:LEU:HD23	4:U:386:ILE:HG22	2.02	0.41
4:V:296:LEU:HA	4:V:297:PRO:HD3	1.94	0.41
5:W:220:LEU:O	5:W:220:LEU:HG	2.21	0.41
5:W:220:LEU:CD1	5:W:241:LEU:HD12	2.50	0.41
5:W:489:LYS:HG3	5:W:649:ILE:HD11	2.02	0.41
5:W:542:ILE:HA	5:W:543:PRO:HD2	1.90	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
5:W:563:LEU:N	5:W:563:LEU:CD1	2.83	0.41
5:W:959:ASN:HB2	5:W:962:GLU:HB2	2.02	0.41
5:W:984:ALA:CB	5:W:1030:THR:HA	2.50	0.41
5:W:1238:ARG:CZ	5:W:1274:ALA:HB1	2.50	0.41
6:X:218:GLU:HB3	6:X:219:LEU:HD22	2.03	0.41
6:X:350:LEU:HD23	6:X:357:ALA:CB	2.50	0.41
6:X:425:ASN:C	6:X:427:LEU:N	2.79	0.41
6:X:433:PRO:HD2	6:X:437:ARG:NH1	2.26	0.41
6:X:518:LEU:HD13	6:X:568:PHE:CZ	2.55	0.41
6:X:597:GLU:CA	6:X:597:GLU:OE2	2.69	0.41
6:Y:200:LEU:HA	6:Y:389:GLU:OE1	2.20	0.41
1:f:29:ARG:NH1	1:f:62:VAL:O	2.54	0.41
1:d:44:GLY:O	3:D:399:ALA:HB3	2.20	0.41
1:n:8:GLN:NE2	3:J:303:ASP:N	2.62	0.41
3:B:44:THR:CG2	3:B:45:GLY:N	2.82	0.41
3:B:62:ALA:O	3:B:63:ILE:C	2.62	0.41
3:B:101:THR:HG21	3:B:159:TRP:CZ2	2.55	0.41
3:B:431:ILE:O	3:B:431:ILE:HG23	2.10	0.41
3:D:616:GLY:O	3:D:620:ALA:HB2	2.19	0.41
7:E:101:MYR:H52	3:F:187:LEU:CD2	2.51	0.41
3:F:73:PHE:CB	3:F:119:VAL:HG21	2.49	0.41
3:F:79:ARG:CB	3:F:79:ARG:HH11	2.33	0.41
3:F:292:ARG:HB2	3:F:503:ALA:HB2	2.03	0.41
3:F:357:ILE:CD1	3:F:414:GLN:HB3	2.50	0.41
3:F:360:TYR:CE1	3:F:415:PRO:HG3	2.56	0.41
3:F:455:SER:O	3:F:456:ASN:HB3	2.21	0.41
2:G:27:THR:HG22	2:G:29:THR:HG23	2.02	0.41
3:H:285:PHE:O	3:H:286:THR:C	2.62	0.41
3:H:360:TYR:CE1	3:H:415:PRO:HG3	2.55	0.41
3:H:502:LEU:O	3:H:503:ALA:C	2.63	0.41
3:J:259:LEU:O	3:J:263:ILE:HG13	2.21	0.41
3:J:455:SER:O	3:J:456:ASN:HB3	2.21	0.41
3:J:469:VAL:O	3:J:469:VAL:HG22	2.20	0.41
3:J:502:LEU:O	3:J:503:ALA:C	2.63	0.41
3:J:576:PHE:HB2	3:J:591:LEU:HD21	2.02	0.41
3:L:156:VAL:HG13	3:L:157:ALA:N	2.35	0.41
3:L:273:LEU:HA	3:N:630:LEU:CD1	2.50	0.41
3:L:279:PRO:HA	3:L:484:ILE:O	2.20	0.41
3:L:420:LEU:HD11	3:L:422:LEU:CD2	2.49	0.41
3:N:565:LEU:HD23	3:N:565:LEU:N	2.34	0.41
3:N:596:MET:SD	3:N:596:MET:N	2.94	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:P:455:SER:O	3:P:456:ASN:HB3	2.21	0.41
3:P:502:LEU:O	3:P:503:ALA:C	2.63	0.41
3:R:343:ALA:CB	3:R:478:GLY:O	2.62	0.41
3:T:79:ARG:O	3:T:83:SER:HB2	2.20	0.41
3:T:166:ALA:HB2	3:T:194:LEU:HD11	2.02	0.41
3:T:171:THR:HG21	3:T:634:LYS:NZ	2.34	0.41
3:T:456:ASN:HA	3:T:457:PRO:HD2	1.93	0.41
3:T:470:LEU:HD21	3:T:472:ARG:HD3	2.02	0.41
3:T:547:HIS:ND1	3:T:547:HIS:C	2.78	0.41
4:U:158:HIS:CG	4:U:297:PRO:HG2	2.55	0.41
4:U:179:GLY:O	4:U:247:VAL:HG13	2.20	0.41
4:U:196:LEU:HD21	6:X:448:SER:CB	2.50	0.41
4:U:240:ALA:C	4:U:241:LEU:HD23	2.46	0.41
4:V:175:MET:HE1	6:Y:400:GLU:OE2	2.19	0.41
4:V:211:LEU:CD1	4:V:248:LEU:HD22	2.51	0.41
5:W:43:LEU:HD11	5:W:64:PRO:HG2	2.01	0.41
5:W:162:SER:CA	5:W:171:THR:HG22	2.37	0.41
5:W:271:ARG:HH21	5:W:328:PRO:CG	2.32	0.41
5:W:310:LEU:HD13	5:W:310:LEU:C	2.46	0.41
5:W:377:ARG:HE	5:W:377:ARG:HB2	1.51	0.41
5:W:509:VAL:O	5:W:510:LEU:HB2	2.20	0.41
5:W:961:GLY:C	5:W:963:ARG:N	2.77	0.41
5:W:1160:TYR:HE2	5:W:1218:ARG:NH1	2.17	0.41
5:W:1204:LEU:HD12	5:W:1222:PRO:HB2	2.01	0.41
6:X:459:LEU:HD22	6:X:540:LEU:HD11	2.02	0.41
6:X:569:LEU:HD23	6:X:815:PRO:HG2	2.03	0.41
6:X:742:MET:HE2	6:X:745:PHE:CB	2.49	0.41
6:X:837:TYR:O	6:X:838:GLY:C	2.62	0.41
6:X:1075:LEU:HD23	6:X:1076:VAL:C	2.46	0.41
6:Y:247:THR:OG1	6:Y:260:GLU:HB3	2.21	0.41
6:Y:277:VAL:HG21	6:Y:295:SER:C	2.46	0.41
6:Y:301:LEU:HD11	6:Y:968:PHE:HB2	2.02	0.41
6:Y:478:THR:O	6:Y:479:LEU:C	2.63	0.41
6:Y:621:TRP:HB2	6:Y:776:GLN:HE21	1.85	0.41
6:Y:1033:ASP:CB	6:Y:1037:VAL:O	2.68	0.41
1:l:16:ALA:HB1	1:l:22:LEU:HD22	2.01	0.41
1:f:58:PRO:CB	3:F:423:GLN:CD	2.79	0.41
1:h:29:ARG:NH1	1:h:62:VAL:O	2.54	0.41
1:j:80:ARG:NH1	1:j:84:GLU:OE2	2.49	0.41
1:n:8:GLN:HE21	3:J:302:GLU:CB	2.34	0.41
3:B:156:VAL:O	3:B:157:ALA:C	2.64	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:B:351:LEU:HA	3:B:464:LEU:HA	2.03	0.41
3:B:360:TYR:CE1	3:B:415:PRO:HG3	2.55	0.41
3:B:369:SER:OG	3:B:443:PHE:CD1	2.64	0.41
3:B:455:SER:O	3:B:456:ASN:HB3	2.21	0.41
3:B:588:ILE:HD13	3:B:588:ILE:HA	1.80	0.41
3:D:360:TYR:CE1	3:D:415:PRO:HG3	2.55	0.41
2:E:26:MET:CG	3:F:216:PRO:CG	2.95	0.41
3:F:431:ILE:CG1	3:F:437:ILE:HD13	2.50	0.41
3:F:445:ILE:HG22	3:F:447:SER:H	1.86	0.41
3:F:469:VAL:O	3:F:469:VAL:HG22	2.20	0.41
3:F:470:LEU:HD13	3:F:470:LEU:O	2.20	0.41
3:F:485:VAL:O	3:F:486:PRO:C	2.64	0.41
3:H:164:ASP:C	3:H:166:ALA:H	2.29	0.41
3:H:214:ARG:HB2	3:H:215:TYR:CE2	2.55	0.41
3:H:279:PRO:HA	3:H:484:ILE:O	2.20	0.41
3:J:96:PHE:CD1	3:J:159:TRP:CH2	3.08	0.41
3:J:144:ASN:O	3:J:145:LEU:C	2.62	0.41
3:J:431:ILE:HD11	3:J:437:ILE:CG2	2.45	0.41
3:J:466:VAL:HG11	3:J:502:LEU:CD2	2.51	0.41
3:J:580:PRO:CD	3:N:289:LEU:HD13	2.50	0.41
3:J:596:MET:SD	3:J:596:MET:N	2.94	0.41
2:K:28:SER:O	2:K:29:THR:CG2	2.68	0.41
3:L:214:ARG:HB2	3:L:215:TYR:CE2	2.55	0.41
3:L:466:VAL:HG11	3:L:502:LEU:CD2	2.51	0.41
3:L:554:GLN:C	3:L:556:SER:H	2.28	0.41
3:L:567:ARG:HB2	3:L:567:ARG:HH11	1.85	0.41
3:N:137:ILE:CG1	3:N:142:ILE:HG13	2.50	0.41
3:N:292:ARG:HB2	3:N:503:ALA:HB2	2.03	0.41
3:N:312:ASN:ND2	3:N:314:SER:O	2.54	0.41
2:O:28:SER:C	2:O:29:THR:HG23	2.45	0.41
3:P:274:ILE:O	3:P:274:ILE:HG13	2.20	0.41
3:P:373:PHE:CD2	3:P:397:VAL:HG11	2.55	0.41
3:P:508:VAL:O	3:P:508:VAL:HG12	2.20	0.41
3:P:580:PRO:CD	3:T:289:LEU:HD13	2.50	0.41
3:P:613:LEU:HD21	2:Q:36:LEU:HD11	2.03	0.41
2:Q:28:SER:C	2:Q:29:THR:HG23	2.45	0.41
3:R:355:GLY:HA2	3:T:298:PHE:CE1	2.47	0.41
3:R:418:ASN:ND2	3:T:337:ARG:NH2	2.55	0.41
3:R:582:LEU:HD23	3:R:582:LEU:HA	1.70	0.41
3:T:106:ILE:CD1	3:T:152:ARG:HG3	2.44	0.41
3:T:164:ASP:C	3:T:166:ALA:H	2.29	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:T:445:ILE:HG22	3:T:447:SER:H	1.86	0.41
4:U:39:SER:HB2	6:X:954:PRO:HB2	2.02	0.41
4:U:112:TRP:CG	4:U:113:ALA:N	2.89	0.41
4:U:245:GLN:NE2	4:U:245:GLN:CA	2.83	0.41
4:U:267:ILE:HG12	4:U:326:LEU:HB2	2.01	0.41
5:W:19:ARG:NH2	5:W:368:PRO:HB3	2.35	0.41
5:W:405:ARG:HH12	5:W:779:MET:HG3	1.86	0.41
5:W:812:ALA:C	5:W:814:GLY:N	2.74	0.41
5:W:896:ILE:HD11	5:W:1025:ILE:HD13	2.02	0.41
5:W:956:TYR:O	5:W:957:ARG:HG3	2.21	0.41
6:X:205:MET:HE2	6:X:207:LEU:HB2	2.01	0.41
6:X:317:PHE:CZ	6:X:331:ARG:HD3	2.55	0.41
6:X:587:MET:HE3	6:X:590:ALA:HB3	2.03	0.41
6:X:633:PRO:C	6:X:635:GLN:N	2.79	0.41
6:X:743:THR:N	6:X:744:PRO:CD	2.83	0.41
6:X:864:PHE:HA	6:X:920:LEU:CD2	2.48	0.41
6:X:874:CYS:CB	6:X:901:ILE:HD13	2.50	0.41
6:X:1013:VAL:HB	6:X:1034:ASP:HB2	2.02	0.41
6:Y:276:THR:O	6:Y:277:VAL:C	2.63	0.41
6:Y:278:LEU:CD1	6:Y:891:PHE:CZ	3.03	0.41
6:Y:410:HIS:HE2	6:Y:413:PRO:HD3	1.85	0.41
6:Y:596:PRO:CB	6:Y:716:ASN:HD22	2.30	0.41
6:Y:606:MET:SD	6:Y:620:VAL:HG13	2.61	0.41
6:Y:686:ARG:O	6:Y:687:LEU:C	2.64	0.41
6:Y:915:LEU:HD12	6:Y:915:LEU:HA	1.79	0.41
6:Y:986:GLN:HG3	6:Y:1135:TYR:HE1	1.83	0.41
6:Y:1205:ASP:OD1	6:Y:1205:ASP:N	2.52	0.41
1:l:24:LEU:HD23	1:l:80:ARG:NE	2.33	0.41
1:f:59:HIS:CA	3:F:398:SER:CB	2.96	0.41
1:d:80:ARG:NH1	1:d:84:GLU:OE2	2.49	0.41
1:h:44:GLY:C	3:H:399:ALA:HB3	2.45	0.41
1:j:29:ARG:NH1	1:j:62:VAL:O	2.54	0.41
1:r:29:ARG:NH1	1:r:62:VAL:O	2.54	0.41
1:t:47:THR:OG1	1:t:57:ALA:HB3	2.21	0.41
2:A:13:ILE:CD1	7:A:101:MYR:H111	2.45	0.41
7:A:101:MYR:H62	3:B:200:CYS:O	2.21	0.41
3:B:48:TRP:CZ3	3:B:123:ALA:N	2.89	0.41
3:B:259:LEU:O	3:B:263:ILE:HG13	2.21	0.41
3:B:282:VAL:HG21	3:B:474:THR:HB	2.02	0.41
2:C:27:THR:HG22	2:C:29:THR:HG23	2.02	0.41
3:D:268:GLY:O	3:D:270:ASP:N	2.54	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:D:332:GLY:HA3	3:D:456:ASN:HA	2.01	0.41
3:D:341:MET:CG	3:D:431:ILE:HD11	2.49	0.41
3:D:576:PHE:HB2	3:D:591:LEU:HD21	2.02	0.41
3:F:48:TRP:CZ3	3:F:123:ALA:N	2.89	0.41
3:F:101:THR:HG21	3:F:159:TRP:CZ2	2.55	0.41
3:F:105:PRO:HB3	3:H:125:ASP:HB3	2.03	0.41
3:F:238:VAL:HG21	3:H:78:MET:HE2	1.83	0.41
3:F:289:LEU:HD13	3:H:580:PRO:CD	2.51	0.41
3:H:93:GLN:OE1	3:H:93:GLN:HA	2.19	0.41
3:H:262:ALA:HB1	3:H:608:LEU:HD13	2.03	0.41
3:H:582:LEU:HD23	3:H:582:LEU:HA	1.70	0.41
2:I:41:LEU:O	2:I:42:ASN:CB	2.37	0.41
3:J:79:ARG:O	3:J:83:SER:HB2	2.20	0.41
3:J:92:TRP:CD2	3:J:210:VAL:HG11	2.54	0.41
3:J:93:GLN:OE1	3:J:93:GLN:HA	2.19	0.41
3:J:105:PRO:HB3	3:L:125:ASP:HB3	2.03	0.41
3:L:101:THR:HG21	3:L:159:TRP:CZ2	2.55	0.41
3:L:312:ASN:ND2	3:L:314:SER:O	2.54	0.41
3:L:469:VAL:O	3:L:469:VAL:HG22	2.20	0.41
3:L:470:LEU:HD13	3:L:470:LEU:O	2.20	0.41
3:N:137:ILE:HG12	3:N:142:ILE:HG13	2.02	0.41
3:N:144:ASN:O	3:N:145:LEU:C	2.62	0.41
3:N:262:ALA:HB1	3:N:608:LEU:HD13	2.03	0.41
3:N:455:SER:O	3:N:456:ASN:HB3	2.21	0.41
3:P:44:THR:CG2	3:P:45:GLY:N	2.82	0.41
3:P:178:SER:HB3	3:P:179:ALA:H	1.65	0.41
3:P:318:LYS:HG3	3:P:319:THR:H	1.84	0.41
3:P:431:ILE:HD11	3:P:437:ILE:CG2	2.45	0.41
2:Q:9:ASN:CB	3:R:209:ASN:ND2	2.83	0.41
3:R:58:ILE:O	3:R:58:ILE:CG2	2.62	0.41
3:R:101:THR:HG21	3:R:159:TRP:CZ2	2.55	0.41
3:R:237:GLU:CB	3:T:46:LYS:NZ	2.68	0.41
3:R:352:ASP:OD1	3:R:352:ASP:C	2.62	0.41
3:R:576:PHE:HB2	3:R:591:LEU:HD21	2.02	0.41
3:T:470:LEU:HD13	3:T:470:LEU:O	2.20	0.41
3:T:485:VAL:O	3:T:486:PRO:C	2.64	0.41
3:T:561:ASP:C	3:T:563:PRO:HD2	2.45	0.41
4:U:30:ALA:CA	4:U:36:PRO:HB3	2.35	0.41
4:U:96:VAL:HA	4:U:135:MET:HE2	2.03	0.41
4:U:170:THR:HG22	4:U:170:THR:O	2.20	0.41
4:U:174:LEU:CD2	4:U:211:LEU:HD23	2.51	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
5:W:1080:SER:HB2	5:W:1083:GLY:O	2.21	0.41
5:W:1102:PRO:HG3	5:W:1127:ILE:HD12	2.01	0.41
5:W:1199:ALA:HB2	5:W:1296:TYR:CE2	2.55	0.41
6:X:311:ILE:HD11	6:X:406:LEU:HD23	2.03	0.41
6:X:638:LEU:HD12	6:X:638:LEU:HA	1.90	0.41
6:X:725:THR:HG23	6:X:726:ARG:N	2.34	0.41
6:X:882:PRO:O	6:X:884:PRO:HD3	2.20	0.41
6:X:1020:ILE:HD12	6:X:1062:TYR:HD2	1.84	0.41
6:X:1150:ASN:O	6:X:1151:ALA:C	2.63	0.41
6:X:1172:LEU:HG	6:X:1173:VAL:HG13	2.02	0.41
6:Y:78:ILE:N	6:Y:78:ILE:CD1	2.83	0.41
6:Y:453:LEU:CD1	6:Y:952:PRO:HD3	2.50	0.41
6:Y:596:PRO:CB	6:Y:716:ASN:ND2	2.83	0.41
6:Y:996:MET:HG2	6:Y:1119:ILE:HD11	2.02	0.41
6:Y:1100:LEU:O	6:Y:1103:TRP:CB	2.69	0.41
1:b:4:HIS:CE1	1:b:54:THR:CB	3.03	0.41
1:b:24:LEU:HD22	1:b:80:ARG:HD3	2.03	0.41
1:b:39:GLU:O	1:b:50:ALA:N	2.47	0.41
1:b:44:GLY:HA3	3:B:399:ALA:HB3	2.00	0.41
1:b:74:GLN:HA	1:b:75:PRO:N	2.34	0.41
1:n:29:ARG:NH1	1:n:62:VAL:O	2.54	0.41
1:n:80:ARG:NH1	1:n:84:GLU:OE2	2.49	0.41
1:r:6:ILE:HD13	3:T:586:VAL:CG2	2.36	0.41
1:t:15:ARG:NH1	3:T:406:ASN:ND2	2.68	0.41
3:B:292:ARG:HB2	3:B:503:ALA:HB2	2.03	0.41
3:B:471:LEU:O	3:B:472:ARG:C	2.64	0.41
2:C:9:ASN:CB	3:D:209:ASN:ND2	2.83	0.41
2:C:28:SER:O	2:C:29:THR:CG2	2.68	0.41
3:D:351:LEU:HA	3:D:464:LEU:HA	2.03	0.41
3:D:561:ASP:C	3:D:563:PRO:HD2	2.45	0.41
3:F:79:ARG:HH22	3:F:135:VAL:HG21	1.86	0.41
3:F:332:GLY:HA3	3:F:456:ASN:HA	2.01	0.41
7:G:101:MYR:H62	3:H:200:CYS:O	2.21	0.41
3:H:266:MET:HE2	3:H:604:VAL:HG12	2.03	0.41
3:H:268:GLY:O	3:H:270:ASP:N	2.54	0.41
3:H:312:ASN:ND2	3:H:314:SER:O	2.54	0.41
3:H:351:LEU:HA	3:H:464:LEU:HA	2.03	0.41
2:I:13:ILE:CD1	7:I:101:MYR:H111	2.45	0.41
3:J:199:LEU:HD11	3:J:203:TYR:CE2	2.54	0.41
3:J:351:LEU:HA	3:J:464:LEU:HA	2.03	0.41
3:J:470:LEU:O	3:J:470:LEU:HD13	2.20	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:J:588:ILE:HD13	3:J:588:ILE:HA	1.80	0.41
3:L:106:ILE:CD1	3:L:152:ARG:HG3	2.44	0.41
3:L:268:GLY:O	3:L:270:ASP:N	2.54	0.41
3:L:556:SER:N	3:L:557:PRO:CD	2.82	0.41
3:L:610:THR:O	3:L:611:GLY:C	2.64	0.41
3:N:470:LEU:HD13	3:N:470:LEU:O	2.20	0.41
3:N:518:ALA:HB1	3:N:580:PRO:HG3	2.03	0.41
3:P:73:PHE:CB	3:P:119:VAL:HG21	2.49	0.41
3:P:125:ASP:HB3	3:T:105:PRO:HB3	2.03	0.41
3:P:312:ASN:ND2	3:P:314:SER:O	2.54	0.41
3:P:336:THR:CG2	3:P:456:ASN:ND2	2.79	0.41
3:P:355:GLY:CA	3:R:298:PHE:CZ	2.80	0.41
3:R:152:ARG:NH1	3:R:155:ASP:OD2	2.53	0.41
3:R:156:VAL:O	3:R:157:ALA:C	2.64	0.41
3:R:156:VAL:HG13	3:R:157:ALA:N	2.35	0.41
3:R:262:ALA:HB1	3:R:608:LEU:HD13	2.03	0.41
3:R:381:PRO:C	3:R:382:ILE:O	2.63	0.41
3:R:455:SER:O	3:R:456:ASN:HB3	2.21	0.41
3:T:79:ARG:CB	3:T:79:ARG:HH11	2.33	0.41
3:T:188:ALA:HA	3:T:191:LYS:HD2	2.02	0.41
3:T:292:ARG:HB2	3:T:503:ALA:HB2	2.03	0.41
3:T:596:MET:SD	3:T:596:MET:N	2.94	0.41
4:U:40:ARG:NH2	4:U:69:ASN:HD21	2.16	0.41
4:U:54:TRP:CZ3	4:U:276:HIS:HE1	2.38	0.41
4:V:107:GLN:N	4:V:108:PRO:HD3	2.35	0.41
4:V:124:ASN:HD22	4:V:124:ASN:HA	1.66	0.41
5:W:178:ILE:HD13	5:W:178:ILE:H	1.86	0.41
5:W:225:THR:HA	5:W:293:THR:O	2.21	0.41
5:W:240:LEU:CD2	5:W:261:LEU:HD13	2.50	0.41
5:W:246:ASP:N	5:W:246:ASP:OD1	2.54	0.41
5:W:376:PHE:CD1	5:W:376:PHE:C	2.98	0.41
5:W:1012:ASP:O	5:W:1015:THR:N	2.53	0.41
6:X:371:LEU:HD12	6:X:371:LEU:N	2.34	0.41
6:X:440:ILE:HG13	6:X:440:ILE:O	2.20	0.41
6:X:607:HIS:C	6:X:609:THR:N	2.79	0.41
6:X:628:PRO:CG	6:X:647:ARG:HD3	2.49	0.41
6:X:651:PRO:O	6:X:773:ARG:CG	2.69	0.41
6:X:985:VAL:HG12	6:X:1073:ARG:HB3	2.01	0.41
6:X:1124:VAL:HG12	6:X:1153:SER:CB	2.51	0.41
6:Y:119:CYS:HB3	6:Y:123:ASN:CA	2.51	0.41
6:Y:425:ASN:C	6:Y:427:LEU:H	2.29	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
6:Y:485:PRO:HD3	6:Y:837:TYR:HE1	1.81	0.41
6:Y:688:THR:OG1	6:Y:691:PRO:HD2	2.21	0.41
6:Y:717:MET:HA	6:Y:717:MET:CE	2.44	0.41
6:Y:729:MET:HE2	6:Y:729:MET:N	2.35	0.41
6:Y:1015:GLY:HA2	6:Y:1034:ASP:HA	2.03	0.41
1:l:29:ARG:NH1	1:l:62:VAL:O	2.54	0.41
1:l:44:GLY:O	3:L:399:ALA:O	2.38	0.41
1:l:57:ALA:HB1	1:l:59:HIS:HD1	1.86	0.41
1:f:44:GLY:CA	3:F:399:ALA:HB3	2.51	0.41
1:n:24:LEU:HD11	1:n:80:ARG:CZ	2.51	0.41
1:n:45:ARG:CB	3:N:399:ALA:O	2.66	0.41
1:p:61:ASN:HB3	3:P:397:VAL:CA	2.23	0.41
1:t:4:HIS:CE1	1:t:54:THR:HG22	2.55	0.41
1:t:51:PHE:HB3	1:t:71:CYS:HB3	2.02	0.41
3:B:73:PHE:CB	3:B:119:VAL:HG21	2.49	0.41
3:B:96:PHE:CD1	3:B:159:TRP:CH2	3.08	0.41
3:B:214:ARG:HB2	3:B:215:TYR:CE2	2.55	0.41
3:B:489:VAL:O	3:B:489:VAL:HG12	2.21	0.41
2:C:36:LEU:HD11	3:H:613:LEU:HD21	2.03	0.41
3:D:101:THR:HG21	3:D:159:TRP:CZ2	2.55	0.41
3:D:156:VAL:HG13	3:D:157:ALA:N	2.35	0.41
3:D:289:LEU:HD13	3:F:580:PRO:CD	2.50	0.41
3:D:370:VAL:O	3:D:370:VAL:CG2	2.67	0.41
3:D:455:SER:O	3:D:456:ASN:HB3	2.21	0.41
3:F:93:GLN:OE1	3:F:93:GLN:HA	2.19	0.41
3:F:259:LEU:O	3:F:263:ILE:HG13	2.21	0.41
3:F:271:LEU:HD12	3:F:271:LEU:HA	1.79	0.41
3:F:279:PRO:HA	3:F:484:ILE:O	2.20	0.41
3:F:289:LEU:HA	3:H:578:SER:O	2.20	0.41
3:F:336:THR:CG2	3:F:456:ASN:ND2	2.79	0.41
3:F:390:ASN:HD22	3:F:390:ASN:HA	1.60	0.41
3:F:567:ARG:HB2	3:F:567:ARG:HH11	1.85	0.41
7:G:101:MYR:C7	3:H:200:CYS:O	2.69	0.41
3:H:47:LEU:C	3:H:48:TRP:CD1	2.99	0.41
3:H:471:LEU:O	3:H:472:ARG:C	2.63	0.41
7:I:101:MYR:C7	3:J:200:CYS:O	2.69	0.41
7:I:101:MYR:H131	3:J:163:LEU:HD11	2.02	0.41
3:J:48:TRP:CZ3	3:J:123:ALA:N	2.89	0.41
3:J:341:MET:HB2	3:J:437:ILE:CG2	2.35	0.41
3:J:418:ASN:HD22	3:J:418:ASN:HA	1.50	0.41
2:K:28:SER:C	2:K:29:THR:HG23	2.45	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
7:K:101:MYR:H52	3:L:187:LEU:CD2	2.51	0.41
3:L:46:LYS:HD3	3:L:70:GLN:OE1	2.19	0.41
3:L:48:TRP:CZ3	3:L:123:ALA:N	2.89	0.41
3:L:62:ALA:O	3:L:63:ILE:C	2.62	0.41
3:L:163:LEU:HD11	3:L:211:ALA:HB2	2.01	0.41
3:L:282:VAL:HG21	3:L:474:THR:HB	2.02	0.41
3:L:292:ARG:HG2	3:L:292:ARG:NH1	2.32	0.41
3:L:370:VAL:O	3:L:370:VAL:CG2	2.67	0.41
3:L:455:SER:O	3:L:456:ASN:HB3	2.21	0.41
3:L:485:VAL:O	3:L:486:PRO:C	2.64	0.41
3:L:596:MET:SD	3:L:596:MET:N	2.94	0.41
3:L:624:LEU:CD2	3:L:624:LEU:C	2.94	0.41
3:N:101:THR:HG21	3:N:159:TRP:CZ2	2.55	0.41
3:N:171:THR:HG21	3:N:634:LYS:NZ	2.34	0.41
3:N:252:LEU:HB3	3:N:613:LEU:HD23	2.02	0.41
3:N:274:ILE:O	3:N:274:ILE:HG13	2.20	0.41
3:N:373:PHE:CD2	3:N:397:VAL:HG11	2.55	0.41
3:N:445:ILE:HG22	3:N:447:SER:H	1.86	0.41
3:N:547:HIS:ND1	3:N:547:HIS:C	2.78	0.41
3:N:554:GLN:C	3:N:556:SER:H	2.28	0.41
3:N:561:ASP:C	3:N:563:PRO:HD2	2.45	0.41
3:N:576:PHE:HB2	3:N:591:LEU:HD21	2.03	0.41
2:O:28:SER:O	2:O:29:THR:CG2	2.68	0.41
2:O:36:LEU:HD11	3:T:613:LEU:HD21	2.03	0.41
3:P:101:THR:HG21	3:P:159:TRP:CZ2	2.55	0.41
3:P:154:ARG:O	3:P:155:ASP:C	2.62	0.41
3:P:268:GLY:O	3:P:270:ASP:N	2.54	0.41
3:P:289:LEU:HA	3:R:578:SER:O	2.20	0.41
3:P:297:ALA:O	3:T:409:ILE:HD11	2.17	0.41
3:P:456:ASN:HA	3:P:457:PRO:HD2	1.93	0.41
3:P:466:VAL:HG11	3:P:502:LEU:CD2	2.51	0.41
3:P:561:ASP:C	3:P:563:PRO:HD2	2.45	0.41
3:R:48:TRP:CZ3	3:R:123:ALA:N	2.89	0.41
3:R:57:THR:HB	3:R:59:ASP:OD1	2.21	0.41
3:R:147:ARG:CB	3:R:147:ARG:CZ	2.98	0.41
3:R:194:LEU:HA	3:R:195:PRO:HD2	1.98	0.41
3:R:274:ILE:O	3:R:274:ILE:HG13	2.20	0.41
3:R:332:GLY:HA3	3:R:333:PRO:HD2	1.90	0.41
3:R:518:ALA:HB1	3:R:580:PRO:HG3	2.03	0.41
3:R:547:HIS:ND1	3:R:547:HIS:C	2.78	0.41
3:R:567:ARG:HB2	3:R:567:ARG:HH11	1.85	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:S:27:THR:HG22	2:S:29:THR:HG23	2.02	0.41
3:T:163:LEU:HD11	3:T:211:ALA:HB2	2.01	0.41
3:T:235:LEU:HA	3:T:235:LEU:HD13	1.80	0.41
3:T:268:GLY:O	3:T:270:ASP:N	2.54	0.41
3:T:312:ASN:ND2	3:T:314:SER:O	2.54	0.41
3:T:401:GLY:O	3:T:402:ALA:CB	2.66	0.41
4:U:13:PHE:CD2	4:U:17:PRO:O	2.73	0.41
4:U:51:LEU:CB	4:U:53:VAL:HG13	2.48	0.41
4:U:150:LEU:HD23	4:U:150:LEU:HA	1.93	0.41
4:U:160:GLU:C	4:U:161:VAL:HG22	2.46	0.41
5:W:4:VAL:O	5:W:5:PHE:HB2	2.20	0.41
5:W:207:TYR:CZ	5:W:237:PRO:HB2	2.55	0.41
5:W:393:ASP:HB3	5:W:791:PHE:O	2.20	0.41
5:W:885:TRP:HH2	5:W:921:ILE:HD11	1.86	0.41
5:W:956:TYR:O	5:W:964:VAL:HG13	2.21	0.41
5:W:967:TYR:CD1	5:W:967:TYR:C	2.99	0.41
5:W:1071:ILE:HA	5:W:1079:ILE:O	2.21	0.41
5:W:1074:THR:O	5:W:1075:PRO:C	2.64	0.41
6:X:329:MET:O	6:X:330:ILE:C	2.64	0.41
6:X:390:LEU:HG	6:X:1190:LEU:O	2.20	0.41
6:X:406:LEU:HD12	6:X:406:LEU:HA	1.86	0.41
6:X:488:LEU:HA	6:X:488:LEU:HD13	1.72	0.41
6:X:510:PRO:HA	6:X:513:GLU:OE1	2.21	0.41
6:X:541:VAL:HA	6:X:809:ALA:O	2.21	0.41
6:X:620:VAL:HG13	6:Y:715:VAL:CG1	2.51	0.41
6:X:681:ASN:ND2	6:X:752:PRO:CB	2.84	0.41
6:X:722:ASN:OD1	6:X:723:ASP:N	2.53	0.41
6:X:795:ASP:O	6:X:798:VAL:HG22	2.21	0.41
6:X:855:GLN:O	6:X:856:ARG:C	2.63	0.41
6:X:1014:VAL:HG21	6:X:1047:TYR:CE1	2.55	0.41
6:X:1084:THR:O	6:X:1116:PRO:HA	2.21	0.41
6:Y:128:THR:O	6:Y:129:MET:C	2.63	0.41
6:Y:257:ARG:O	6:Y:258:GLY:C	2.62	0.41
6:Y:284:ASP:C	6:Y:286:ALA:H	2.27	0.41
6:Y:336:ARG:HA	6:Y:337:PRO:HD2	1.93	0.41
6:Y:579:PRO:HG3	6:Y:807:PHE:CE2	2.55	0.41
6:Y:593:LEU:HD12	6:Y:597:GLU:HG3	2.02	0.41
6:Y:633:PRO:HD3	6:Y:640:ARG:HD3	2.02	0.41
6:Y:715:VAL:O	6:Y:718:THR:HG22	2.20	0.41
6:Y:725:THR:O	6:Y:729:MET:HG2	2.21	0.41
6:Y:773:ARG:O	6:Y:774:LEU:HB2	2.20	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
6:Y:1059:PHE:HA	6:Y:1063:VAL:CG1	2.51	0.41
6:Y:1100:LEU:HD22	6:Y:1104:LEU:HD21	2.03	0.41
6:Y:1102:GLU:OE2	6:Y:1106:LYS:HE3	2.21	0.41
6:Y:1123:GLN:HB2	6:Y:1158:PHE:HB3	2.03	0.41
6:Y:1211:ARG:O	6:Y:1212:SER:CB	2.69	0.41
1:f:80:ARG:NH1	1:f:84:GLU:OE2	2.49	0.41
1:d:8:GLN:CD	3:F:302:GLU:CG	2.55	0.41
1:h:21:ARG:O	3:H:323:SER:CB	2.65	0.41
1:j:15:ARG:NE	3:J:406:ASN:ND2	2.67	0.41
1:r:15:ARG:HH12	3:R:402:ALA:CB	2.33	0.41
1:r:74:GLN:NE2	1:r:75:PRO:C	2.78	0.41
3:B:361:ASP:OD2	3:B:362:THR:HG23	2.20	0.41
3:B:518:ALA:HB1	3:B:580:PRO:HG3	2.03	0.41
3:B:596:MET:SD	3:B:596:MET:N	2.94	0.41
3:D:229:ASP:O	3:D:231:GLN:N	2.54	0.41
3:D:469:VAL:O	3:D:469:VAL:HG22	2.20	0.41
3:F:57:THR:HB	3:F:59:ASP:OD1	2.21	0.41
3:F:58:ILE:HG23	3:F:124:GLY:CA	2.42	0.41
3:F:229:ASP:O	3:F:231:GLN:N	2.54	0.41
3:F:489:VAL:O	3:F:489:VAL:HG12	2.21	0.41
3:H:229:ASP:O	3:H:231:GLN:N	2.54	0.41
3:H:252:LEU:HB3	3:H:613:LEU:HD23	2.02	0.41
3:H:582:LEU:HD23	3:H:585:GLN:HE22	1.83	0.41
3:H:624:LEU:CD2	3:H:624:LEU:C	2.94	0.41
3:J:268:GLY:O	3:J:270:ASP:N	2.54	0.41
7:K:101:MYR:H42	3:L:187:LEU:HD22	2.03	0.41
3:L:79:ARG:CB	3:L:79:ARG:HH11	2.33	0.41
3:L:381:PRO:C	3:L:382:ILE:O	2.63	0.41
3:L:547:HIS:ND1	3:L:547:HIS:C	2.78	0.41
3:N:73:PHE:CB	3:N:119:VAL:HG21	2.49	0.41
3:N:142:ILE:O	3:N:142:ILE:CG2	2.69	0.41
3:N:259:LEU:O	3:N:263:ILE:HG13	2.21	0.41
3:N:332:GLY:HA3	3:N:456:ASN:HA	2.01	0.41
3:N:485:VAL:O	3:N:486:PRO:C	2.64	0.41
2:O:26:MET:CG	3:P:216:PRO:CG	2.95	0.41
3:P:67:ARG:CG	3:P:67:ARG:NH1	2.84	0.41
3:P:93:GLN:OE1	3:P:93:GLN:HA	2.19	0.41
3:P:124:GLY:HA3	3:T:105:PRO:HG2	2.03	0.41
3:P:229:ASP:O	3:P:231:GLN:N	2.54	0.41
3:P:262:ALA:HB1	3:P:608:LEU:HD13	2.03	0.41
3:P:282:VAL:HG21	3:P:474:THR:HB	2.02	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:P:609:ARG:CD	2:Q:29:THR:HG21	2.47	0.41
3:R:292:ARG:HB2	3:R:503:ALA:HB2	2.03	0.41
3:R:466:VAL:HG11	3:R:502:LEU:CD2	2.51	0.41
2:S:23:THR:C	2:S:25:ASP:N	2.75	0.41
7:S:101:MYR:H62	3:T:200:CYS:O	2.21	0.41
3:T:351:LEU:HA	3:T:464:LEU:HA	2.03	0.41
3:T:455:SER:O	3:T:456:ASN:HB3	2.21	0.41
4:U:124:ASN:HA	5:W:353:ARG:HH12	1.86	0.41
4:U:181:PRO:HG3	6:X:384:MET:CE	2.50	0.41
4:U:328:TYR:HB2	4:U:333:MET:HE2	2.03	0.41
5:W:570:VAL:HA	5:W:571:PRO:HD3	1.90	0.41
5:W:650:VAL:CG1	5:W:651:ASN:N	2.81	0.41
5:W:732:LEU:CD1	5:W:780:LEU:HD22	2.50	0.41
5:W:1019:TYR:O	5:W:1019:TYR:CD1	2.74	0.41
5:W:1053:THR:HG23	5:W:1095:GLU:OE1	2.21	0.41
5:W:1258:MET:CG	5:W:1259:SER:N	2.82	0.41
6:X:638:LEU:HD11	6:X:711:VAL:HA	2.02	0.41
6:X:784:LEU:HD22	6:X:800:ILE:HG23	2.03	0.41
6:X:1004:PHE:CZ	6:X:1075:LEU:HB2	2.56	0.41
6:X:1178:ARG:O	6:X:1181:THR:HG23	2.21	0.41
6:Y:374:ALA:CB	6:Y:377:ILE:HD11	2.25	0.41
6:Y:550:PHE:CD2	6:Y:612:ALA:HB3	2.56	0.41
6:Y:615:PHE:CD2	6:Y:620:VAL:HG12	2.56	0.41
6:Y:873:GLN:NE2	6:Y:965:GLU:OE1	2.48	0.41
6:Y:1072:THR:HG22	6:Y:1073:ARG:N	2.35	0.41
6:Y:1090:PRO:HB2	6:Y:1129:THR:HG22	2.02	0.41
1:d:29:ARG:NH1	1:d:62:VAL:O	2.54	0.40
1:d:61:ASN:C	1:d:63:LYS:N	2.54	0.40
1:n:15:ARG:NH1	3:N:402:ALA:HB3	2.27	0.40
1:n:61:ASN:HB2	3:N:397:VAL:HA	2.03	0.40
1:n:74:GLN:NE2	1:n:74:GLN:C	2.77	0.40
1:r:21:ARG:CD	3:R:327:TYR:OH	2.69	0.40
3:B:47:LEU:C	3:B:48:TRP:CD1	2.99	0.40
3:B:93:GLN:OE1	3:B:93:GLN:HA	2.19	0.40
3:B:193:ILE:H	3:B:193:ILE:HG12	1.55	0.40
3:B:610:THR:O	3:B:611:GLY:C	2.64	0.40
3:D:463:LEU:HD23	3:D:463:LEU:O	2.22	0.40
7:E:101:MYR:H42	3:F:187:LEU:HD22	2.03	0.40
3:F:47:LEU:C	3:F:48:TRP:CD1	2.99	0.40
3:F:431:ILE:O	3:F:431:ILE:HG23	2.10	0.40
3:F:463:LEU:HD23	3:F:463:LEU:O	2.21	0.40

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:F:576:PHE:HB2	3:F:591:LEU:HD21	2.02	0.40
3:F:582:LEU:HD23	3:F:585:GLN:HE22	1.83	0.40
3:H:57:THR:HB	3:H:59:ASP:OD1	2.21	0.40
3:H:193:ILE:H	3:H:193:ILE:HG12	1.55	0.40
3:H:489:VAL:O	3:H:489:VAL:HG12	2.21	0.40
3:J:47:LEU:C	3:J:48:TRP:CD1	2.99	0.40
3:J:156:VAL:HG13	3:J:157:ALA:N	2.35	0.40
3:J:485:VAL:O	3:J:486:PRO:C	2.64	0.40
3:J:582:LEU:HD23	3:J:585:GLN:HE22	1.83	0.40
3:J:616:GLY:O	3:J:620:ALA:HB2	2.20	0.40
3:L:44:THR:CG2	3:L:45:GLY:N	2.82	0.40
3:L:259:LEU:O	3:L:263:ILE:HG13	2.21	0.40
3:L:508:VAL:O	3:L:508:VAL:HG12	2.20	0.40
3:L:582:LEU:C	3:L:584:LEU:N	2.79	0.40
3:N:47:LEU:C	3:N:48:TRP:CD1	2.99	0.40
3:N:199:LEU:HD11	3:N:203:TYR:CE2	2.54	0.40
3:N:588:ILE:HD13	3:N:588:ILE:HA	1.80	0.40
3:P:47:LEU:C	3:P:48:TRP:CD1	2.99	0.40
3:P:48:TRP:CZ3	3:P:123:ALA:N	2.89	0.40
3:P:78:MET:HE2	3:T:238:VAL:CG1	2.45	0.40
3:P:322:TYR:C	3:P:323:SER:HG	2.16	0.40
3:P:343:ALA:CB	3:P:478:GLY:O	2.62	0.40
3:P:373:PHE:O	3:P:397:VAL:HG12	2.21	0.40
3:P:445:ILE:HG22	3:P:447:SER:H	1.86	0.40
3:R:166:ALA:HB2	3:R:194:LEU:HD11	2.02	0.40
3:R:373:PHE:CD2	3:R:397:VAL:HG11	2.55	0.40
3:R:373:PHE:O	3:R:397:VAL:HG12	2.22	0.40
3:T:373:PHE:O	3:T:397:VAL:HG12	2.21	0.40
3:T:508:VAL:O	3:T:508:VAL:HG12	2.20	0.40
4:U:136:TRP:CD2	4:U:197:PRO:HG3	2.56	0.40
4:U:242:ASN:ND2	4:U:244:ASP:HB3	2.26	0.40
4:V:182:ILE:HG21	4:V:279:VAL:HG23	2.02	0.40
5:W:10:VAL:HA	5:W:11:PRO:HD3	1.88	0.40
5:W:14:ASN:HB3	5:W:319:GLN:HE21	1.81	0.40
5:W:250:TYR:HB2	5:W:255:ILE:HD12	2.02	0.40
5:W:369:ILE:HG22	5:W:370:GLY:N	2.35	0.40
5:W:396:ARG:NH1	5:W:396:ARG:HG2	2.36	0.40
5:W:486:SER:HB2	5:W:490:ASP:OD1	2.22	0.40
5:W:765:PHE:HA	5:W:768:ASN:HD22	1.86	0.40
5:W:823:THR:HG22	5:W:847:PRO:CD	2.49	0.40
5:W:879:TYR:HE2	5:W:898:THR:HG21	1.85	0.40

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
5:W:1066:ASN:HD22	5:W:1066:ASN:HA	1.61	0.40
5:W:1290:PRO:HB2	5:W:1291:ALA:H	1.59	0.40
6:X:282:TYR:HD2	6:Y:830:ASN:HD21	1.68	0.40
6:X:409:MET:O	6:X:409:MET:HG3	2.20	0.40
6:X:497:ILE:CD1	6:X:514:ILE:HD12	2.46	0.40
6:X:596:PRO:C	6:X:597:GLU:OE2	2.64	0.40
6:X:693:TYR:OH	6:X:740:LYS:HA	2.21	0.40
6:X:749:HIS:O	6:X:752:PRO:HD2	2.20	0.40
6:X:1020:ILE:HG12	6:X:1031:LEU:HD22	2.02	0.40
6:X:1067:VAL:CG2	6:X:1072:THR:HG23	2.52	0.40
6:Y:200:LEU:HB2	6:Y:250:ALA:HB3	2.03	0.40
6:Y:221:GLY:HA3	6:Y:230:GLN:HA	2.04	0.40
6:Y:685:PRO:HD3	6:Y:749:HIS:HD1	1.86	0.40
6:Y:832:ILE:CG2	6:Y:1213:ALA:HA	2.52	0.40
6:Y:864:PHE:CE2	6:Y:931:ILE:HD12	2.55	0.40
6:Y:1136:ALA:O	6:Y:1137:PHE:CB	2.68	0.40
1:b:18:ALA:O	1:b:86:SER:HB3	2.21	0.40
1:j:22:LEU:HD22	1:j:23:THR:N	2.36	0.40
3:B:79:ARG:HH22	3:B:135:VAL:HG21	1.86	0.40
3:B:171:THR:HG21	3:B:634:LYS:NZ	2.34	0.40
3:B:502:LEU:O	3:B:503:ALA:C	2.63	0.40
7:C:101:MYR:C7	3:D:200:CYS:O	2.69	0.40
3:D:259:LEU:O	3:D:263:ILE:HG13	2.21	0.40
3:D:361:ASP:OD2	3:D:362:THR:HG23	2.20	0.40
3:D:431:ILE:CG1	3:D:437:ILE:HD13	2.50	0.40
3:D:445:ILE:HG22	3:D:447:SER:H	1.86	0.40
3:D:547:HIS:ND1	3:D:547:HIS:C	2.78	0.40
3:F:268:GLY:O	3:F:270:ASP:N	2.54	0.40
3:F:466:VAL:HG11	3:F:502:LEU:CD2	2.51	0.40
3:H:106:ILE:CD1	3:H:152:ARG:HG3	2.45	0.40
3:H:431:ILE:CG1	3:H:437:ILE:HD13	2.50	0.40
3:H:460:MET:C	3:H:462:GLY:N	2.78	0.40
3:H:469:VAL:O	3:H:469:VAL:HG22	2.20	0.40
2:I:9:ASN:CB	3:J:209:ASN:ND2	2.83	0.40
3:J:73:PHE:CB	3:J:119:VAL:HG21	2.49	0.40
3:J:125:ASP:HB3	3:N:105:PRO:HB3	2.03	0.40
3:J:373:PHE:O	3:J:397:VAL:HG12	2.21	0.40
3:L:274:ILE:O	3:L:274:ILE:HG13	2.20	0.40
3:N:79:ARG:HH22	3:N:135:VAL:HG21	1.86	0.40
3:N:373:PHE:O	3:N:397:VAL:HG12	2.21	0.40
3:N:624:LEU:CD2	3:N:624:LEU:C	2.94	0.40

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
7:O:101:MYR:C7	3:P:200:CYS:O	2.69	0.40
3:P:235:LEU:HD13	3:P:235:LEU:HA	1.80	0.40
3:P:352:ASP:OD1	3:P:352:ASP:C	2.62	0.40
3:R:409:ILE:HD11	3:T:297:ALA:O	2.17	0.40
7:S:101:MYR:H131	3:T:163:LEU:HD11	2.04	0.40
3:T:489:VAL:O	3:T:489:VAL:HG12	2.21	0.40
3:T:582:LEU:C	3:T:584:LEU:N	2.79	0.40
4:U:26:LEU:HD21	4:U:100:LEU:CD1	2.51	0.40
4:U:140:GLN:NE2	4:U:190:TYR:CE1	2.87	0.40
4:U:209:ARG:HG3	4:U:290:VAL:HG11	2.03	0.40
4:U:344:ASP:C	4:U:346:LEU:N	2.79	0.40
5:W:145:ASP:HB2	5:W:148:ILE:HB	2.02	0.40
5:W:262:MET:HE2	5:W:262:MET:HB3	1.81	0.40
5:W:268:ASP:C	5:W:270:LEU:N	2.79	0.40
5:W:430:HIS:HA	5:W:799:LEU:HD23	2.03	0.40
5:W:480:ARG:HH21	5:W:488:ILE:CD1	2.34	0.40
5:W:766:LEU:HD21	5:W:1000:TYR:HE1	1.85	0.40
5:W:787:HIS:O	5:W:788:GLN:C	2.63	0.40
5:W:885:TRP:CZ3	5:W:921:ILE:HD13	2.55	0.40
5:W:989:HIS:N	5:W:989:HIS:CD2	2.88	0.40
5:W:1026:LEU:HD13	5:W:1028:ILE:HG13	2.02	0.40
5:W:1250:LEU:HG	5:W:1260:LEU:HD21	2.03	0.40
6:X:218:GLU:O	6:X:219:LEU:HD23	2.21	0.40
6:X:259:LEU:HD23	6:X:911:THR:HB	2.03	0.40
6:X:772:THR:HG22	6:X:775:VAL:H	1.85	0.40
6:X:795:ASP:CG	6:X:798:VAL:CG2	2.89	0.40
6:X:798:VAL:HG21	6:Y:737:ARG:NH1	2.36	0.40
6:X:798:VAL:HG23	6:X:803:THR:CG2	2.52	0.40
6:X:850:MET:HE3	6:X:851:PHE:HE1	1.82	0.40
6:X:973:ASN:C	6:X:975:PHE:H	2.29	0.40
6:X:1049:SER:O	6:X:1050:ASP:C	2.64	0.40
6:Y:560:LEU:HB2	6:Y:561:PRO:HD3	2.03	0.40
6:Y:680:PRO:CG	6:Y:696:ALA:HB2	2.51	0.40
6:Y:937:ALA:HB2	6:Y:947:ARG:HA	2.03	0.40
6:Y:976:ASN:O	6:Y:1083:TYR:CE2	2.74	0.40
6:Y:978:ALA:O	6:Y:1080:HIS:HA	2.22	0.40
6:Y:1085:LEU:HD22	6:Y:1119:ILE:HD12	2.03	0.40
1:b:29:ARG:NH1	1:b:62:VAL:O	2.54	0.40
1:b:44:GLY:O	3:B:399:ALA:O	2.39	0.40
1:j:15:ARG:NH2	3:J:402:ALA:HB2	2.33	0.40
1:j:29:ARG:NH2	1:j:65:ILE:O	2.49	0.40

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:n:24:LEU:HD11	1:n:80:ARG:HE	1.83	0.40
1:t:61:ASN:HB2	3:T:395:THR:CG2	2.52	0.40
2:A:9:ASN:CB	3:B:209:ASN:ND2	2.83	0.40
3:B:199:LEU:HD11	3:B:203:TYR:CE2	2.54	0.40
3:B:312:ASN:ND2	3:B:314:SER:O	2.54	0.40
3:B:357:ILE:CD1	3:B:414:GLN:HB3	2.50	0.40
3:B:445:ILE:HG22	3:B:447:SER:H	1.86	0.40
3:D:100:LYS:O	3:L:152:ARG:NH2	2.49	0.40
3:D:156:VAL:O	3:D:157:ALA:C	2.64	0.40
3:D:225:THR:O	3:D:225:THR:HG23	2.22	0.40
3:D:271:LEU:HD12	3:D:271:LEU:HA	1.79	0.40
3:D:273:LEU:HA	3:F:630:LEU:CD1	2.50	0.40
3:D:401:GLY:O	3:D:402:ALA:CB	2.66	0.40
3:D:456:ASN:C	3:D:456:ASN:ND2	2.76	0.40
3:D:489:VAL:HG12	3:D:489:VAL:O	2.21	0.40
3:D:582:LEU:C	3:D:584:LEU:N	2.79	0.40
3:F:67:ARG:CG	3:F:67:ARG:NH1	2.84	0.40
3:F:373:PHE:CD2	3:F:397:VAL:HG11	2.55	0.40
3:F:547:HIS:ND1	3:F:547:HIS:C	2.78	0.40
3:H:44:THR:CG2	3:H:45:GLY:N	2.82	0.40
3:H:455:SER:O	3:H:456:ASN:HB3	2.21	0.40
2:I:29:THR:CG2	3:N:609:ARG:CD	2.93	0.40
3:J:53:THR:HA	4:V:332:PRO:CG	2.47	0.40
3:J:312:ASN:ND2	3:J:314:SER:O	2.54	0.40
3:J:428:ARG:NH1	3:J:428:ARG:CG	2.76	0.40
3:J:463:LEU:HD23	3:J:463:LEU:O	2.21	0.40
3:L:229:ASP:O	3:L:231:GLN:N	2.54	0.40
3:L:471:LEU:O	3:L:472:ARG:C	2.64	0.40
2:M:13:ILE:CG2	2:M:14:THR:H	2.35	0.40
7:M:101:MYR:C7	3:N:200:CYS:O	2.69	0.40
7:M:101:MYR:H62	3:N:200:CYS:O	2.21	0.40
3:N:48:TRP:CZ3	3:N:123:ALA:N	2.89	0.40
3:N:57:THR:HB	3:N:59:ASP:OD1	2.21	0.40
3:N:167:MET:HG2	3:N:215:TYR:CE2	2.57	0.40
3:N:297:ALA:CB	3:N:460:MET:HB2	2.33	0.40
3:N:489:VAL:HG12	3:N:489:VAL:O	2.21	0.40
3:P:485:VAL:O	3:P:486:PRO:C	2.64	0.40
3:P:576:PHE:HB2	3:P:591:LEU:HD21	2.03	0.40
7:Q:101:MYR:H52	3:R:187:LEU:CD2	2.51	0.40
3:R:193:ILE:H	3:R:193:ILE:HG12	1.55	0.40
3:R:266:MET:HE2	3:R:604:VAL:HG12	2.03	0.40

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:R:292:ARG:HH11	3:R:292:ARG:CG	2.32	0.40
3:R:312:ASN:ND2	3:R:314:SER:O	2.54	0.40
3:R:360:TYR:CE1	3:R:415:PRO:HG3	2.56	0.40
3:R:485:VAL:O	3:R:486:PRO:C	2.64	0.40
3:T:182:ASN:O	3:T:182:ASN:CG	2.65	0.40
3:T:282:VAL:HG21	3:T:474:THR:HB	2.02	0.40
3:T:463:LEU:HD23	3:T:463:LEU:O	2.21	0.40
4:U:22:ASP:HB2	4:U:383:ARG:CD	2.39	0.40
4:U:61:VAL:HG21	4:U:278:ILE:HB	2.02	0.40
4:U:157:TYR:CD1	4:U:157:TYR:N	2.88	0.40
4:U:157:TYR:CB	4:U:287:SER:HB2	2.51	0.40
4:U:243:PRO:HD2	4:U:362:TYR:CD1	2.56	0.40
4:U:265:SER:HA	4:U:323:VAL:CG1	2.52	0.40
4:V:65:ASN:O	4:V:68:VAL:HG12	2.21	0.40
4:V:156:ILE:O	4:V:156:ILE:HG13	2.21	0.40
5:W:162:SER:HA	5:W:171:THR:CG2	2.36	0.40
5:W:459:ASP:O	5:W:460:ALA:C	2.63	0.40
5:W:722:ASP:OD1	5:W:782:THR:OG1	2.39	0.40
5:W:992:SER:HA	5:W:993:PRO:HD3	1.87	0.40
6:X:242:ARG:HH12	6:X:1146:LEU:HD12	1.86	0.40
6:X:298:LEU:C	6:X:298:LEU:CD2	2.94	0.40
6:X:470:LEU:C	6:X:472:LEU:H	2.29	0.40
6:X:606:MET:HE2	6:Y:719:ALA:N	2.35	0.40
6:X:788:VAL:HG22	6:X:789:ASP:H	1.86	0.40
6:X:945:VAL:HG13	6:X:947:ARG:HG2	2.03	0.40
6:Y:43:ALA:C	6:Y:45:ASP:N	2.79	0.40
6:Y:126:PHE:HD2	6:Y:132:LEU:HD13	1.87	0.40
6:Y:338:GLU:H	6:Y:338:GLU:HG3	1.63	0.40
6:Y:451:LEU:HB3	6:Y:666:ALA:CB	2.51	0.40
6:Y:488:LEU:CD2	6:Y:821:ALA:HB1	2.50	0.40
6:Y:581:SER:N	6:Y:582:PRO:CD	2.84	0.40
1:l:45:ARG:CG	3:L:400:ALA:C	2.81	0.40
1:b:21:ARG:HB2	1:b:43:CYS:O	2.22	0.40
1:d:39:GLU:O	1:d:50:ALA:N	2.47	0.40
1:n:6:ILE:CD1	3:J:586:VAL:CB	2.98	0.40
1:p:58:PRO:HG3	3:P:423:GLN:HG2	1.79	0.40
1:p:58:PRO:CG	3:P:423:GLN:NE2	2.50	0.40
3:B:567:ARG:HB2	3:B:567:ARG:HH11	1.85	0.40
3:B:591:LEU:HD23	3:B:591:LEU:HA	1.89	0.40
3:D:50:PRO:HD3	3:D:61:LEU:CD2	2.51	0.40
3:D:262:ALA:HB1	3:D:608:LEU:HD13	2.03	0.40

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:D:292:ARG:HG2	3:D:292:ARG:NH1	2.32	0.40
3:D:292:ARG:HB2	3:D:503:ALA:HB2	2.03	0.40
3:D:442:THR:OG1	3:D:443:PHE:N	2.50	0.40
3:D:624:LEU:CD2	3:D:624:LEU:C	2.94	0.40
2:E:19:SER:OG	3:F:228:PRO:HG3	2.22	0.40
3:H:79:ARG:HH22	3:H:135:VAL:HG21	1.86	0.40
3:H:156:VAL:O	3:H:157:ALA:C	2.64	0.40
3:J:357:ILE:CD1	3:J:414:GLN:HB3	2.50	0.40
3:J:547:HIS:ND1	3:J:547:HIS:C	2.78	0.40
3:J:630:LEU:CD1	3:N:273:LEU:HA	2.50	0.40
2:K:19:SER:OG	3:L:228:PRO:HG3	2.22	0.40
3:L:167:MET:HG2	3:L:215:TYR:CE2	2.57	0.40
3:L:576:PHE:HB2	3:L:591:LEU:HD21	2.03	0.40
3:N:431:ILE:CG1	3:N:437:ILE:HD13	2.50	0.40
3:N:582:LEU:HD23	3:N:585:GLN:HE22	1.83	0.40
3:P:156:VAL:O	3:P:157:ALA:C	2.64	0.40
3:P:259:LEU:O	3:P:263:ILE:HG13	2.21	0.40
3:P:266:MET:HE2	3:P:604:VAL:HG12	2.03	0.40
3:P:273:LEU:HA	3:R:630:LEU:CD1	2.50	0.40
2:Q:13:ILE:CG2	2:Q:14:THR:H	2.35	0.40
3:R:229:ASP:O	3:R:231:GLN:N	2.54	0.40
3:R:582:LEU:C	3:R:584:LEU:N	2.79	0.40
3:R:610:THR:O	3:R:611:GLY:C	2.64	0.40
2:S:28:SER:C	2:S:29:THR:HG23	2.45	0.40
3:T:266:MET:HE2	3:T:604:VAL:HG12	2.03	0.40
4:U:217:GLY:HA3	4:U:223:LEU:HD12	2.03	0.40
4:U:242:ASN:HB2	6:X:1179:VAL:O	2.21	0.40
4:V:20:LEU:HD13	4:V:384:MET:SD	2.61	0.40
4:V:57:ARG:O	4:V:59:PRO:HD3	2.21	0.40
5:W:154:PHE:HA	5:W:155:PRO:HD3	1.77	0.40
5:W:477:ARG:CA	5:W:650:VAL:HG11	2.41	0.40
5:W:534:LEU:HD22	5:W:551:ILE:HD13	2.03	0.40
5:W:780:LEU:N	5:W:780:LEU:HD12	2.35	0.40
5:W:815:TYR:C	5:W:817:SER:N	2.79	0.40
6:X:200:LEU:HD13	6:X:343:ALA:HB2	2.03	0.40
6:X:206:LEU:HB2	6:X:236:PRO:HD3	2.02	0.40
6:X:367:ARG:HG3	6:X:388:MET:CE	2.51	0.40
6:X:580:HIS:C	6:X:582:PRO:CD	2.94	0.40
6:X:593:LEU:HD21	6:X:597:GLU:CB	2.51	0.40
6:X:754:GLU:HG3	6:X:837:TYR:CZ	2.56	0.40
6:X:859:ILE:HA	6:X:859:ILE:HD13	1.82	0.40

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
6:X:929:PRO:O	6:X:931:ILE:N	2.54	0.40
6:X:1034:ASP:C	6:X:1036:ASN:H	2.29	0.40
6:X:1043:GLY:O	6:X:1072:THR:HA	2.22	0.40
6:Y:59:GLY:HA3	6:Y:60:PRO:HD3	1.98	0.40
6:Y:195:ILE:HD12	6:Y:257:ARG:NH2	2.36	0.40
6:Y:222:PHE:O	6:Y:228:MET:HA	2.22	0.40
6:Y:456:VAL:O	6:Y:456:VAL:HG12	2.22	0.40
6:Y:541:VAL:CA	6:Y:808:ALA:HB1	2.52	0.40
1:f:59:HIS:HB3	3:F:398:SER:OG	2.17	0.40
1:d:73:ARG:CD	1:d:73:ARG:O	2.69	0.40
1:j:76:ASN:C	1:j:78:ALA:N	2.79	0.40
1:t:4:HIS:CE1	1:t:54:THR:CG2	3.05	0.40
3:B:57:THR:HB	3:B:59:ASP:OD1	2.21	0.40
3:B:485:VAL:O	3:B:486:PRO:C	2.64	0.40
3:D:48:TRP:CH2	3:D:63:ILE:HD12	2.56	0.40
3:D:190:ALA:C	3:D:192:ASP:N	2.80	0.40
3:D:358:LEU:HB2	3:D:417:PHE:HD1	1.77	0.40
3:D:373:PHE:O	3:D:397:VAL:HG12	2.22	0.40
3:D:518:ALA:HB1	3:D:580:PRO:HG3	2.03	0.40
3:D:610:THR:O	3:D:611:GLY:C	2.64	0.40
3:F:158:LEU:O	3:F:161:LYS:HB3	2.21	0.40
3:F:182:ASN:O	3:F:182:ASN:CG	2.65	0.40
3:F:299:ILE:O	3:F:337:ARG:HD2	2.22	0.40
3:F:508:VAL:O	3:F:508:VAL:HG12	2.20	0.40
3:H:101:THR:HG21	3:H:159:TRP:CZ2	2.55	0.40
3:H:182:ASN:O	3:H:182:ASN:CG	2.65	0.40
3:H:373:PHE:O	3:H:397:VAL:HG12	2.21	0.40
3:H:420:LEU:CD1	3:H:422:LEU:HD23	2.52	0.40
3:H:445:ILE:HG22	3:H:447:SER:H	1.86	0.40
3:H:463:LEU:HD23	3:H:463:LEU:O	2.22	0.40
3:J:496:GLU:C	3:J:498:VAL:N	2.80	0.40
3:J:610:THR:O	3:J:611:GLY:C	2.64	0.40
3:J:624:LEU:C	3:J:624:LEU:CD2	2.94	0.40
3:L:225:THR:O	3:L:225:THR:HG23	2.22	0.40
3:L:262:ALA:HB1	3:L:608:LEU:HD13	2.03	0.40
3:N:411:VAL:O	3:N:411:VAL:CG1	2.61	0.40
3:N:466:VAL:HG11	3:N:502:LEU:CD2	2.51	0.40
3:P:299:ILE:O	3:P:337:ARG:HD2	2.22	0.40
3:P:431:ILE:O	3:P:431:ILE:HG23	2.10	0.40
3:P:450:SER:CA	3:P:453:ASN:HD22	2.21	0.40
3:P:582:LEU:C	3:P:584:LEU:N	2.79	0.40

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:R:79:ARG:O	3:R:83:SER:HB2	2.20	0.40
3:R:79:ARG:HH22	3:R:135:VAL:HG21	1.86	0.40
3:R:182:ASN:O	3:R:182:ASN:CG	2.64	0.40
3:R:385:VAL:O	3:R:385:VAL:HG22	2.22	0.40
3:R:554:GLN:C	3:R:556:SER:H	2.28	0.40
3:R:591:LEU:HD23	3:R:591:LEU:HA	1.89	0.40
2:S:19:SER:OG	3:T:228:PRO:HG3	2.22	0.40
7:S:101:MYR:C7	3:T:200:CYS:O	2.69	0.40
3:T:154:ARG:NH1	3:T:154:ARG:CB	2.68	0.40
3:T:299:ILE:O	3:T:337:ARG:HD2	2.22	0.40
3:T:554:GLN:C	3:T:556:SER:H	2.28	0.40
4:U:252:ASP:HA	4:U:255:ARG:NH1	2.36	0.40
4:U:362:TYR:CZ	4:U:366:ILE:HG13	2.57	0.40
4:U:367:GLU:HG3	4:U:394:PHE:CD1	2.55	0.40
4:U:374:LEU:HD13	4:U:374:LEU:C	2.46	0.40
4:V:20:LEU:HD13	4:V:379:ILE:HD11	2.03	0.40
4:V:72:LEU:O	4:V:76:LEU:HG	2.21	0.40
5:W:822:GLN:HA	5:W:822:GLN:OE1	2.22	0.40
5:W:934:ASN:HA	5:W:1023:THR:HA	2.03	0.40
5:W:1114:THR:HG22	5:W:1115:LEU:N	2.36	0.40
5:W:1166:CYS:HA	5:W:1176:VAL:H	1.87	0.40
6:X:294:ARG:HB2	6:X:888:LEU:CB	2.52	0.40
6:X:328:ARG:HA	6:X:328:ARG:HD2	1.57	0.40
6:X:450:ASP:OD2	6:X:861:ARG:CZ	2.69	0.40
6:X:607:HIS:C	6:X:609:THR:H	2.29	0.40
6:X:842:ALA:HB3	6:X:843:PRO:CD	2.52	0.40
6:X:879:PHE:O	6:X:880:LEU:C	2.65	0.40
6:X:921:ASP:C	6:X:923:LEU:H	2.29	0.40
6:X:1064:TRP:C	6:X:1066:MET:N	2.79	0.40
6:Y:541:VAL:O	6:Y:541:VAL:HG22	2.21	0.40
6:Y:563:MET:C	6:Y:563:MET:SD	3.04	0.40
6:Y:652:GLN:HA	6:Y:773:ARG:CD	2.44	0.40

There are no symmetry-related clashes.

5.3 Torsion angles (i)

5.3.1 Protein backbone (i)

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

entries.

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

Mol	Chain	Analysed	Favoured	Allowed	Outliers	Percentiles
1	b	84/276 (30%)	70 (83%)	9 (11%)	5 (6%)	1 9
1	d	84/276 (30%)	70 (83%)	11 (13%)	3 (4%)	2 17
1	f	84/276 (30%)	68 (81%)	12 (14%)	4 (5%)	2 12
1	h	84/276 (30%)	71 (84%)	10 (12%)	3 (4%)	2 17
1	j	84/276 (30%)	72 (86%)	9 (11%)	3 (4%)	2 17
1	l	84/276 (30%)	70 (83%)	13 (16%)	1 (1%)	10 37
1	n	84/276 (30%)	69 (82%)	12 (14%)	3 (4%)	2 17
1	p	84/276 (30%)	74 (88%)	7 (8%)	3 (4%)	2 17
1	r	84/276 (30%)	71 (84%)	11 (13%)	2 (2%)	4 24
1	t	84/276 (30%)	72 (86%)	10 (12%)	2 (2%)	4 24
2	A	39/42 (93%)	25 (64%)	9 (23%)	5 (13%)	0 1
2	C	39/42 (93%)	25 (64%)	9 (23%)	5 (13%)	0 1
2	E	39/42 (93%)	25 (64%)	9 (23%)	5 (13%)	0 1
2	G	37/42 (88%)	24 (65%)	8 (22%)	5 (14%)	0 1
2	I	39/42 (93%)	25 (64%)	9 (23%)	5 (13%)	0 1
2	K	39/42 (93%)	25 (64%)	9 (23%)	5 (13%)	0 1
2	M	37/42 (88%)	24 (65%)	8 (22%)	5 (14%)	0 1
2	O	39/42 (93%)	25 (64%)	9 (23%)	5 (13%)	0 1
2	Q	39/42 (93%)	25 (64%)	9 (23%)	5 (13%)	0 1
2	S	37/42 (88%)	24 (65%)	8 (22%)	5 (14%)	0 1
3	B	602/606 (99%)	407 (68%)	134 (22%)	61 (10%)	0 3
3	D	602/606 (99%)	416 (69%)	129 (21%)	57 (10%)	0 3
3	F	602/606 (99%)	408 (68%)	133 (22%)	61 (10%)	0 3
3	H	602/606 (99%)	409 (68%)	132 (22%)	61 (10%)	0 3
3	J	602/606 (99%)	411 (68%)	132 (22%)	59 (10%)	0 3
3	L	602/606 (99%)	409 (68%)	132 (22%)	61 (10%)	0 3
3	N	602/606 (99%)	408 (68%)	133 (22%)	61 (10%)	0 3
3	P	602/606 (99%)	410 (68%)	132 (22%)	60 (10%)	0 3
3	R	602/606 (99%)	406 (67%)	135 (22%)	61 (10%)	0 3

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Mol	Chain	Analysed	Favoured	Allowed	Outliers	Percentiles
3	T	602/606 (99%)	406 (67%)	135 (22%)	61 (10%)	0 3
4	U	409/412 (99%)	283 (69%)	81 (20%)	45 (11%)	0 2
4	V	409/412 (99%)	330 (81%)	60 (15%)	19 (5%)	2 13
5	W	1276/1299 (98%)	1063 (83%)	173 (14%)	40 (3%)	3 20
6	X	1012/1214 (83%)	714 (71%)	219 (22%)	79 (8%)	1 5
6	Y	1146/1214 (94%)	860 (75%)	204 (18%)	82 (7%)	1 6
All	All	11496/13791 (83%)	8294 (72%)	2255 (20%)	947 (8%)	1 5

All (947) Ramachandran outliers are listed below:

Mol	Chain	Res	Type
1	l	19	ALA
1	b	62	VAL
1	b	75	PRO
1	f	75	PRO
1	d	22	LEU
1	d	62	VAL
1	j	62	VAL
1	j	75	PRO
1	p	61	ASN
1	p	63	LYS
1	r	61	ASN
1	t	63	LYS
3	B	59	ASP
3	B	103	CYS
3	B	179	ALA
3	B	225	THR
3	B	228	PRO
3	B	250	GLY
3	B	270	ASP
3	B	291	PRO
3	B	345	ILE
3	B	388	ALA
3	B	405	ILE
3	B	409	ILE
3	B	412	ARG
3	B	470	LEU
3	B	486	PRO
3	B	555	ALA
3	B	562	TYR

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Mol	Chain	Res	Type
3	B	591	LEU
3	D	53	THR
3	D	103	CYS
3	D	179	ALA
3	D	225	THR
3	D	228	PRO
3	D	250	GLY
3	D	270	ASP
3	D	291	PRO
3	D	345	ILE
3	D	388	ALA
3	D	405	ILE
3	D	409	ILE
3	D	412	ARG
3	D	470	LEU
3	D	486	PRO
3	D	555	ALA
3	D	562	TYR
3	D	591	LEU
3	F	59	ASP
3	F	103	CYS
3	F	179	ALA
3	F	225	THR
3	F	228	PRO
3	F	250	GLY
3	F	270	ASP
3	F	291	PRO
3	F	345	ILE
3	F	388	ALA
3	F	405	ILE
3	F	409	ILE
3	F	412	ARG
3	F	470	LEU
3	F	486	PRO
3	F	555	ALA
3	F	562	TYR
3	F	591	LEU
3	H	59	ASP
3	H	103	CYS
3	H	179	ALA
3	H	225	THR
3	H	228	PRO

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Mol	Chain	Res	Type
3	H	250	GLY
3	H	270	ASP
3	H	291	PRO
3	H	345	ILE
3	H	388	ALA
3	H	405	ILE
3	H	409	ILE
3	H	412	ARG
3	H	470	LEU
3	H	486	PRO
3	H	555	ALA
3	H	562	TYR
3	H	591	LEU
3	J	54	SER
3	J	59	ASP
3	J	103	CYS
3	J	179	ALA
3	J	225	THR
3	J	228	PRO
3	J	250	GLY
3	J	270	ASP
3	J	291	PRO
3	J	345	ILE
3	J	388	ALA
3	J	405	ILE
3	J	409	ILE
3	J	412	ARG
3	J	470	LEU
3	J	486	PRO
3	J	555	ALA
3	J	562	TYR
3	J	591	LEU
3	L	59	ASP
3	L	103	CYS
3	L	179	ALA
3	L	225	THR
3	L	228	PRO
3	L	250	GLY
3	L	270	ASP
3	L	291	PRO
3	L	345	ILE
3	L	388	ALA

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Mol	Chain	Res	Type
3	L	405	ILE
3	L	409	ILE
3	L	412	ARG
3	L	470	LEU
3	L	486	PRO
3	L	555	ALA
3	L	562	TYR
3	L	591	LEU
3	N	59	ASP
3	N	103	CYS
3	N	179	ALA
3	N	225	THR
3	N	228	PRO
3	N	250	GLY
3	N	270	ASP
3	N	291	PRO
3	N	345	ILE
3	N	388	ALA
3	N	405	ILE
3	N	409	ILE
3	N	412	ARG
3	N	470	LEU
3	N	486	PRO
3	N	555	ALA
3	N	562	TYR
3	N	591	LEU
3	P	58	ILE
3	P	103	CYS
3	P	179	ALA
3	P	225	THR
3	P	228	PRO
3	P	250	GLY
3	P	270	ASP
3	P	291	PRO
3	P	345	ILE
3	P	388	ALA
3	P	405	ILE
3	P	409	ILE
3	P	412	ARG
3	P	470	LEU
3	P	486	PRO
3	P	555	ALA

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Mol	Chain	Res	Type
3	P	562	TYR
3	P	591	LEU
3	R	59	ASP
3	R	103	CYS
3	R	179	ALA
3	R	225	THR
3	R	228	PRO
3	R	250	GLY
3	R	270	ASP
3	R	291	PRO
3	R	345	ILE
3	R	388	ALA
3	R	405	ILE
3	R	409	ILE
3	R	412	ARG
3	R	470	LEU
3	R	486	PRO
3	R	555	ALA
3	R	562	TYR
3	R	591	LEU
3	T	59	ASP
3	T	103	CYS
3	T	179	ALA
3	T	225	THR
3	T	228	PRO
3	T	250	GLY
3	T	270	ASP
3	T	291	PRO
3	T	345	ILE
3	T	388	ALA
3	T	405	ILE
3	T	409	ILE
3	T	412	ARG
3	T	470	LEU
3	T	486	PRO
3	T	555	ALA
3	T	562	TYR
3	T	591	LEU
4	U	37	TRP
4	U	104	PRO
4	U	132	LEU
4	U	168	ALA

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Mol	Chain	Res	Type
4	U	221	GLY
4	U	240	ALA
4	U	252	ASP
4	U	303	ALA
4	U	318	ASN
4	U	378	PRO
4	U	379	ILE
4	U	390	ASN
4	U	391	ALA
4	U	400	THR
4	V	40	ARG
4	V	378	PRO
5	W	63	PRO
5	W	369	ILE
5	W	370	GLY
5	W	638	HIS
5	W	1040	VAL
5	W	1290	PRO
6	X	280	ARG
6	X	328	ARG
6	X	343	ALA
6	X	438	PRO
6	X	519	GLN
6	X	611	PRO
6	X	612	ALA
6	X	773	ARG
6	X	776	GLN
6	X	845	PHE
6	X	880	LEU
6	X	890	GLN
6	X	925	LEU
6	X	1025	ASN
6	X	1096	GLU
6	Y	39	PRO
6	Y	103	PRO
6	Y	105	SER
6	Y	106	MET
6	Y	123	ASN
6	Y	217	LYS
6	Y	239	ALA
6	Y	277	VAL
6	Y	355	LYS

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Mol	Chain	Res	Type
6	Y	372	ASP
6	Y	389	GLU
6	Y	440	ILE
6	Y	554	SER
6	Y	665	SER
6	Y	692	THR
6	Y	777	ASN
6	Y	778	ASP
6	Y	788	VAL
6	Y	879	PHE
6	Y	918	ALA
6	Y	954	PRO
6	Y	1055	SER
6	Y	1137	PHE
6	Y	1183	ASP
6	Y	1184	LEU
6	Y	1213	ALA
1	f	23	THR
1	f	61	ASN
2	A	6	THR
2	A	10	THR
2	A	27	THR
3	B	100	LYS
3	B	170	LEU
3	B	193	ILE
3	B	196	LEU
3	B	198	ASN
3	B	268	GLY
3	B	313	SER
3	B	332	GLY
3	B	346	ASP
3	B	384	GLN
3	B	428	ARG
3	B	582	LEU
3	B	583	ALA
3	B	596	MET
3	B	610	THR
3	B	611	GLY
3	B	612	SER
2	C	6	THR
2	C	10	THR
2	C	27	THR

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Mol	Chain	Res	Type
3	D	100	LYS
3	D	170	LEU
3	D	193	ILE
3	D	196	LEU
3	D	198	ASN
3	D	268	GLY
3	D	313	SER
3	D	332	GLY
3	D	346	ASP
3	D	384	GLN
3	D	428	ARG
3	D	582	LEU
3	D	583	ALA
3	D	596	MET
3	D	610	THR
3	D	611	GLY
3	D	612	SER
2	E	6	THR
2	E	10	THR
2	E	27	THR
3	F	100	LYS
3	F	170	LEU
3	F	193	ILE
3	F	196	LEU
3	F	198	ASN
3	F	268	GLY
3	F	313	SER
3	F	332	GLY
3	F	346	ASP
3	F	384	GLN
3	F	428	ARG
3	F	582	LEU
3	F	583	ALA
3	F	596	MET
3	F	610	THR
3	F	611	GLY
3	F	612	SER
2	G	6	THR
2	G	10	THR
2	G	27	THR
3	H	100	LYS
3	H	170	LEU

Continued on next page...

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Mol	Chain	Res	Type
3	H	193	ILE
3	H	196	LEU
3	H	198	ASN
3	H	268	GLY
3	H	313	SER
3	H	332	GLY
3	H	346	ASP
3	H	384	GLN
3	H	428	ARG
3	H	582	LEU
3	H	583	ALA
3	H	596	MET
3	H	610	THR
3	H	611	GLY
3	H	612	SER
2	I	6	THR
2	I	10	THR
2	I	27	THR
3	J	100	LYS
3	J	170	LEU
3	J	193	ILE
3	J	196	LEU
3	J	198	ASN
3	J	268	GLY
3	J	313	SER
3	J	332	GLY
3	J	346	ASP
3	J	384	GLN
3	J	428	ARG
3	J	582	LEU
3	J	583	ALA
3	J	596	MET
3	J	610	THR
3	J	611	GLY
3	J	612	SER
2	K	6	THR
2	K	10	THR
2	K	27	THR
3	L	100	LYS
3	L	170	LEU
3	L	193	ILE
3	L	196	LEU

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Mol	Chain	Res	Type
3	L	198	ASN
3	L	268	GLY
3	L	313	SER
3	L	332	GLY
3	L	346	ASP
3	L	384	GLN
3	L	428	ARG
3	L	582	LEU
3	L	583	ALA
3	L	596	MET
3	L	610	THR
3	L	611	GLY
3	L	612	SER
2	M	6	THR
2	M	10	THR
2	M	27	THR
3	N	100	LYS
3	N	170	LEU
3	N	193	ILE
3	N	196	LEU
3	N	198	ASN
3	N	268	GLY
3	N	313	SER
3	N	332	GLY
3	N	346	ASP
3	N	384	GLN
3	N	428	ARG
3	N	582	LEU
3	N	583	ALA
3	N	596	MET
3	N	610	THR
3	N	611	GLY
3	N	612	SER
2	O	6	THR
2	O	10	THR
2	O	27	THR
3	P	100	LYS
3	P	170	LEU
3	P	193	ILE
3	P	196	LEU
3	P	198	ASN
3	P	268	GLY

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Mol	Chain	Res	Type
3	P	313	SER
3	P	332	GLY
3	P	346	ASP
3	P	384	GLN
3	P	428	ARG
3	P	582	LEU
3	P	583	ALA
3	P	596	MET
3	P	610	THR
3	P	611	GLY
3	P	612	SER
2	Q	6	THR
2	Q	10	THR
2	Q	27	THR
3	R	100	LYS
3	R	170	LEU
3	R	193	ILE
3	R	196	LEU
3	R	198	ASN
3	R	268	GLY
3	R	313	SER
3	R	332	GLY
3	R	346	ASP
3	R	384	GLN
3	R	428	ARG
3	R	582	LEU
3	R	583	ALA
3	R	596	MET
3	R	610	THR
3	R	611	GLY
3	R	612	SER
2	S	6	THR
2	S	10	THR
2	S	27	THR
3	T	100	LYS
3	T	170	LEU
3	T	193	ILE
3	T	196	LEU
3	T	198	ASN
3	T	268	GLY
3	T	313	SER
3	T	332	GLY

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Mol	Chain	Res	Type
3	T	346	ASP
3	T	384	GLN
3	T	428	ARG
3	T	582	LEU
3	T	583	ALA
3	T	596	MET
3	T	610	THR
3	T	611	GLY
3	T	612	SER
4	U	17	PRO
4	U	105	ALA
4	U	122	SER
4	U	167	PRO
4	U	178	THR
4	U	195	GLN
4	U	222	ARG
4	U	229	CYS
4	U	294	THR
4	U	312	VAL
4	V	5	GLN
4	V	20	LEU
4	V	247	VAL
4	V	379	ILE
5	W	176	LYS
5	W	368	PRO
5	W	559	VAL
5	W	592	MET
5	W	660	GLY
5	W	788	GLN
5	W	954	HIS
5	W	962	GLU
6	X	270	SER
6	X	363	ALA
6	X	369	GLY
6	X	416	ILE
6	X	426	ASN
6	X	428	LEU
6	X	464	SER
6	X	518	LEU
6	X	602	GLY
6	X	634	GLN
6	X	721	VAL

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Mol	Chain	Res	Type
6	X	818	ILE
6	X	914	ASP
6	X	937	ALA
6	X	946	VAL
6	X	996	MET
6	X	1004	PHE
6	X	1016	ARG
6	X	1155	ASP
6	X	1190	LEU
6	Y	257	ARG
6	Y	258	GLY
6	Y	259	LEU
6	Y	332	ALA
6	Y	333	PHE
6	Y	351	PRO
6	Y	353	GLN
6	Y	375	ASN
6	Y	376	LEU
6	Y	541	VAL
6	Y	608	GLN
6	Y	673	ASN
6	Y	791	ALA
6	Y	848	ASN
6	Y	996	MET
6	Y	1070	GLY
6	Y	1155	ASP
6	Y	1179	VAL
1	b	61	ASN
1	h	61	ASN
1	r	75	PRO
2	A	32	PRO
3	B	204	PRO
3	B	226	LYS
3	B	307	ILE
3	B	503	ALA
3	B	532	ASP
3	B	553	GLY
3	B	554	GLN
2	C	32	PRO
3	D	204	PRO
3	D	226	LYS
3	D	307	ILE

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Mol	Chain	Res	Type
3	D	503	ALA
3	D	532	ASP
3	D	553	GLY
3	D	554	GLN
2	E	32	PRO
3	F	204	PRO
3	F	226	LYS
3	F	307	ILE
3	F	503	ALA
3	F	532	ASP
3	F	553	GLY
3	F	554	GLN
2	G	32	PRO
3	H	204	PRO
3	H	226	LYS
3	H	307	ILE
3	H	503	ALA
3	H	532	ASP
3	H	553	GLY
3	H	554	GLN
2	I	32	PRO
3	J	204	PRO
3	J	226	LYS
3	J	307	ILE
3	J	503	ALA
3	J	532	ASP
3	J	553	GLY
3	J	554	GLN
2	K	32	PRO
3	L	204	PRO
3	L	226	LYS
3	L	307	ILE
3	L	503	ALA
3	L	532	ASP
3	L	553	GLY
3	L	554	GLN
2	M	32	PRO
3	N	204	PRO
3	N	226	LYS
3	N	307	ILE
3	N	503	ALA
3	N	532	ASP

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Mol	Chain	Res	Type
3	N	553	GLY
3	N	554	GLN
2	O	32	PRO
3	P	204	PRO
3	P	226	LYS
3	P	307	ILE
3	P	503	ALA
3	P	532	ASP
3	P	553	GLY
3	P	554	GLN
2	Q	32	PRO
3	R	204	PRO
3	R	226	LYS
3	R	307	ILE
3	R	503	ALA
3	R	532	ASP
3	R	553	GLY
3	R	554	GLN
2	S	32	PRO
3	T	204	PRO
3	T	226	LYS
3	T	307	ILE
3	T	503	ALA
3	T	532	ASP
3	T	553	GLY
3	T	554	GLN
4	U	14	TYR
4	U	20	LEU
4	U	134	SER
4	U	169	ALA
4	U	193	ALA
4	U	297	PRO
4	U	396	GLN
4	U	398	GLY
4	V	17	PRO
4	V	81	GLY
4	V	85	THR
4	V	256	ALA
4	V	397	ALA
5	W	54	ARG
5	W	162	SER
5	W	662	ARG

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Mol	Chain	Res	Type
5	W	742	SER
5	W	848	PRO
5	W	863	LEU
5	W	991	LEU
5	W	1229	ILE
5	W	1256	PRO
6	X	278	LEU
6	X	353	GLN
6	X	525	TYR
6	X	588	ALA
6	X	613	SER
6	X	695	SER
6	X	720	THR
6	X	829	THR
6	X	930	ARG
6	X	1024	PRO
6	X	1131	ARG
6	X	1147	PHE
6	X	1151	ALA
6	X	1185	PRO
6	X	1189	THR
6	Y	62	ALA
6	Y	72	VAL
6	Y	82	PRO
6	Y	168	ARG
6	Y	214	HIS
6	Y	218	GLU
6	Y	225	PHE
6	Y	352	THR
6	Y	538	ASN
6	Y	552	ASP
6	Y	809	ALA
6	Y	824	CYS
6	Y	829	THR
6	Y	1016	ARG
6	Y	1081	TYR
1	h	74	GLN
1	p	59	HIS
3	B	182	ASN
3	B	296	PRO
3	B	398	SER
3	B	402	ALA

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Mol	Chain	Res	Type
3	B	414	GLN
3	B	590	VAL
3	B	613	LEU
3	D	182	ASN
3	D	296	PRO
3	D	398	SER
3	D	402	ALA
3	D	414	GLN
3	D	590	VAL
3	D	613	LEU
3	F	182	ASN
3	F	296	PRO
3	F	398	SER
3	F	402	ALA
3	F	414	GLN
3	F	590	VAL
3	F	613	LEU
3	H	182	ASN
3	H	296	PRO
3	H	398	SER
3	H	402	ALA
3	H	414	GLN
3	H	590	VAL
3	H	613	LEU
3	J	182	ASN
3	J	296	PRO
3	J	398	SER
3	J	402	ALA
3	J	414	GLN
3	J	590	VAL
3	J	613	LEU
3	L	182	ASN
3	L	296	PRO
3	L	398	SER
3	L	402	ALA
3	L	414	GLN
3	L	590	VAL
3	L	613	LEU
3	N	182	ASN
3	N	296	PRO
3	N	398	SER
3	N	402	ALA

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Mol	Chain	Res	Type
3	N	414	GLN
3	N	590	VAL
3	N	613	LEU
3	P	182	ASN
3	P	296	PRO
3	P	398	SER
3	P	402	ALA
3	P	414	GLN
3	P	590	VAL
3	P	613	LEU
3	R	182	ASN
3	R	296	PRO
3	R	398	SER
3	R	402	ALA
3	R	414	GLN
3	R	590	VAL
3	R	613	LEU
3	T	182	ASN
3	T	296	PRO
3	T	398	SER
3	T	402	ALA
3	T	414	GLN
3	T	590	VAL
3	T	613	LEU
4	U	38	THR
4	U	114	VAL
4	U	187	GLN
4	U	196	LEU
4	U	246	ALA
4	U	387	GLN
4	V	33	TYR
4	V	161	VAL
4	V	178	THR
4	V	387	GLN
5	W	743	SER
5	W	1117	GLN
6	X	219	LEU
6	X	285	ASN
6	X	408	SER
6	X	579	PRO
6	X	722	ASN
6	X	1036	ASN

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Mol	Chain	Res	Type
6	X	1079	GLY
6	Y	498	THR
6	Y	649	ASN
6	Y	780	ILE
6	Y	810	ALA
6	Y	914	ASP
1	d	6	ILE
1	n	61	ASN
3	B	54	SER
3	B	199	LEU
3	B	305	LYS
3	B	362	THR
3	B	403	SER
3	B	419	MET
3	B	434	ILE
3	D	199	LEU
3	D	305	LYS
3	D	362	THR
3	D	403	SER
3	D	419	MET
3	D	434	ILE
3	F	54	SER
3	F	199	LEU
3	F	305	LYS
3	F	362	THR
3	F	403	SER
3	F	419	MET
3	F	434	ILE
3	H	54	SER
3	H	199	LEU
3	H	305	LYS
3	H	362	THR
3	H	403	SER
3	H	419	MET
3	H	434	ILE
3	J	199	LEU
3	J	305	LYS
3	J	362	THR
3	J	403	SER
3	J	419	MET
3	J	434	ILE
3	L	54	SER

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Mol	Chain	Res	Type
3	L	199	LEU
3	L	305	LYS
3	L	362	THR
3	L	403	SER
3	L	419	MET
3	L	434	ILE
3	N	54	SER
3	N	199	LEU
3	N	305	LYS
3	N	362	THR
3	N	403	SER
3	N	419	MET
3	N	434	ILE
3	P	54	SER
3	P	199	LEU
3	P	305	LYS
3	P	362	THR
3	P	403	SER
3	P	419	MET
3	P	434	ILE
3	R	54	SER
3	R	199	LEU
3	R	305	LYS
3	R	362	THR
3	R	403	SER
3	R	419	MET
3	R	434	ILE
3	T	54	SER
3	T	199	LEU
3	T	305	LYS
3	T	362	THR
3	T	403	SER
3	T	419	MET
3	T	434	ILE
4	U	82	GLY
4	U	85	THR
4	U	242	ASN
4	U	392	ALA
4	V	195	GLN
4	V	305	LEU
5	W	52	VAL
5	W	199	PRO

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Mol	Chain	Res	Type
5	W	424	SER
5	W	453	PRO
5	W	740	VAL
5	W	1131	PRO
6	X	368	ILE
6	X	421	ASN
6	X	513	GLU
6	X	596	PRO
6	X	664	GLY
6	X	824	CYS
6	X	843	PRO
6	X	926	TYR
6	X	947	ARG
6	Y	18	ASP
6	Y	185	SER
6	Y	360	LEU
6	Y	373	ARG
6	Y	387	CYS
6	Y	614	GLN
6	Y	919	LEU
6	Y	1093	SER
1	t	6	ILE
4	U	112	TRP
4	V	388	PRO
5	W	64	PRO
5	W	960	GLY
6	X	255	GLU
6	X	794	GLY
6	X	943	GLY
6	X	1049	SER
6	Y	1004	PHE
6	Y	1116	PRO
1	b	20	GLY
3	B	50	PRO
3	B	58	ILE
3	B	171	THR
3	B	527	VAL
3	D	171	THR
3	D	527	VAL
3	F	50	PRO
3	F	58	ILE
3	F	171	THR

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Mol	Chain	Res	Type
3	F	527	VAL
3	H	50	PRO
3	H	58	ILE
3	H	171	THR
3	H	527	VAL
3	J	171	THR
3	J	527	VAL
3	L	50	PRO
3	L	58	ILE
3	L	171	THR
3	L	527	VAL
3	N	50	PRO
3	N	58	ILE
3	N	171	THR
3	N	527	VAL
3	P	50	PRO
3	P	171	THR
3	P	527	VAL
3	R	50	PRO
3	R	58	ILE
3	R	171	THR
3	R	527	VAL
3	T	50	PRO
3	T	58	ILE
3	T	171	THR
3	T	527	VAL
5	W	155	PRO
5	W	744	ILE
5	W	781	ILE
6	X	1164	VAL
6	X	1179	VAL
6	X	1207	ILE
6	Y	81	ARG
1	f	6	ILE
1	n	6	ILE
2	A	38	PRO
2	C	38	PRO
2	E	38	PRO
2	G	38	PRO
2	I	38	PRO
2	K	38	PRO
2	M	38	PRO

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Mol	Chain	Res	Type
2	O	38	PRO
2	Q	38	PRO
2	S	38	PRO
4	U	126	PRO
4	V	272	PRO
5	W	411	VAL
6	X	637	PRO
6	Y	266	PRO
6	Y	690	ALA
1	b	6	ILE
1	j	6	ILE
1	n	74	GLN
3	B	64	VAL
3	F	64	VAL
3	H	64	VAL
3	J	64	VAL
3	L	64	VAL
3	N	64	VAL
3	P	64	VAL
3	R	64	VAL
3	T	64	VAL
4	U	296	LEU
5	W	993	PRO
6	X	490	PRO
6	X	922	GLY
6	X	1038	PRO
6	Y	220	ILE
6	Y	671	PRO
5	W	371	PRO
6	X	770	PRO
6	Y	551	PRO
6	Y	917	GLY
1	h	6	ILE
5	W	826	PRO
6	X	440	ILE
6	Y	800	ILE

5.3.2 Protein sidechains (i)

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

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

Mol	Chain	Analysed	Rotameric	Outliers	Percentiles	
1	b	72/228 (32%)	66 (92%)	6 (8%)	10	34
1	d	72/228 (32%)	68 (94%)	4 (6%)	19	47
1	f	72/228 (32%)	69 (96%)	3 (4%)	26	55
1	h	72/228 (32%)	66 (92%)	6 (8%)	10	34
1	j	72/228 (32%)	68 (94%)	4 (6%)	19	47
1	l	72/228 (32%)	67 (93%)	5 (7%)	14	41
1	n	72/228 (32%)	69 (96%)	3 (4%)	26	55
1	p	72/228 (32%)	67 (93%)	5 (7%)	14	41
1	r	72/228 (32%)	67 (93%)	5 (7%)	14	41
1	t	72/228 (32%)	69 (96%)	3 (4%)	26	55
2	A	34/35 (97%)	25 (74%)	9 (26%)	0	2
2	C	34/35 (97%)	25 (74%)	9 (26%)	0	2
2	E	34/35 (97%)	25 (74%)	9 (26%)	0	2
2	G	32/35 (91%)	23 (72%)	9 (28%)	0	2
2	I	34/35 (97%)	25 (74%)	9 (26%)	0	2
2	K	34/35 (97%)	25 (74%)	9 (26%)	0	2
2	M	32/35 (91%)	23 (72%)	9 (28%)	0	2
2	O	34/35 (97%)	25 (74%)	9 (26%)	0	2
2	Q	34/35 (97%)	25 (74%)	9 (26%)	0	2
2	S	32/35 (91%)	23 (72%)	9 (28%)	0	2
3	B	499/501 (100%)	412 (83%)	87 (17%)	2	9
3	D	499/501 (100%)	410 (82%)	89 (18%)	2	8
3	F	499/501 (100%)	413 (83%)	86 (17%)	2	10
3	H	499/501 (100%)	413 (83%)	86 (17%)	2	10
3	J	499/501 (100%)	410 (82%)	89 (18%)	2	8
3	L	499/501 (100%)	412 (83%)	87 (17%)	2	9
3	N	499/501 (100%)	410 (82%)	89 (18%)	2	8
3	P	499/501 (100%)	410 (82%)	89 (18%)	2	8
3	R	499/501 (100%)	412 (83%)	87 (17%)	2	9
3	T	499/501 (100%)	410 (82%)	89 (18%)	2	8

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Mol	Chain	Analysed	Rotameric	Outliers	Percentiles
4	U	325/326 (100%)	305 (94%)	20 (6%)	16 45
4	V	325/326 (100%)	306 (94%)	19 (6%)	18 47
5	W	1082/1092 (99%)	979 (90%)	103 (10%)	8 29
6	X	869/1030 (84%)	726 (84%)	143 (16%)	2 11
6	Y	976/1030 (95%)	865 (89%)	111 (11%)	5 21
All	All	9621/11444 (84%)	8213 (85%)	1408 (15%)	5 14

All (1408) residues with a non-rotameric sidechain are listed below:

Mol	Chain	Res	Type
1	l	4	HIS
1	l	23	THR
1	l	24	LEU
1	l	59	HIS
1	l	74	GLN
1	b	3	LEU
1	b	4	HIS
1	b	6	ILE
1	b	61	ASN
1	b	73	ARG
1	b	74	GLN
1	f	22	LEU
1	f	24	LEU
1	f	72	SER
1	d	22	LEU
1	d	24	LEU
1	d	73	ARG
1	d	74	GLN
1	h	3	LEU
1	h	4	HIS
1	h	23	THR
1	h	72	SER
1	h	73	ARG
1	h	74	GLN
1	j	3	LEU
1	j	5	MET
1	j	22	LEU
1	j	73	ARG
1	n	3	LEU
1	n	4	HIS

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Mol	Chain	Res	Type
1	n	74	GLN
1	p	3	LEU
1	p	4	HIS
1	p	24	LEU
1	p	71	CYS
1	p	73	ARG
1	r	4	HIS
1	r	5	MET
1	r	22	LEU
1	r	72	SER
1	r	74	GLN
1	t	3	LEU
1	t	22	LEU
1	t	59	HIS
2	A	7	SER
2	A	8	VAL
2	A	10	THR
2	A	12	ASN
2	A	23	THR
2	A	27	THR
2	A	40	VAL
2	A	41	LEU
2	A	42	ASN
3	B	46	LYS
3	B	47	LEU
3	B	48	TRP
3	B	51	VAL
3	B	76	GLU
3	B	78	MET
3	B	79	ARG
3	B	80	GLU
3	B	95	LEU
3	B	100	LYS
3	B	115	THR
3	B	116	SER
3	B	122	THR
3	B	125	ASP
3	B	143	MET
3	B	146	GLN
3	B	148	THR
3	B	152	ARG
3	B	153	MET

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Mol	Chain	Res	Type
3	B	158	LEU
3	B	171	THR
3	B	173	ASP
3	B	182	ASN
3	B	193	ILE
3	B	198	ASN
3	B	199	LEU
3	B	206	GLU
3	B	218	LEU
3	B	225	THR
3	B	226	LYS
3	B	228	PRO
3	B	248	GLU
3	B	269	LYS
3	B	271	LEU
3	B	272	ASP
3	B	273	LEU
3	B	274	ILE
3	B	278	THR
3	B	309	GLU
3	B	311	ASN
3	B	315	LEU
3	B	319	THR
3	B	330	GLN
3	B	336	THR
3	B	341	MET
3	B	344	MET
3	B	349	LEU
3	B	352	ASP
3	B	356	THR
3	B	360	TYR
3	B	374	VAL
3	B	385	VAL
3	B	390	ASN
3	B	406	ASN
3	B	409	ILE
3	B	418	ASN
3	B	419	MET
3	B	420	LEU
3	B	422	LEU
3	B	428	ARG
3	B	431	ILE

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Mol	Chain	Res	Type
3	B	440	LEU
3	B	443	PHE
3	B	453	ASN
3	B	456	ASN
3	B	464	LEU
3	B	466	VAL
3	B	470	LEU
3	B	471	LEU
3	B	493	GLN
3	B	494	THR
3	B	496	GLU
3	B	502	LEU
3	B	515	ASN
3	B	523	ASP
3	B	527	VAL
3	B	536	THR
3	B	548	LYS
3	B	549	LEU
3	B	565	LEU
3	B	588	ILE
3	B	606	THR
3	B	610	THR
3	B	612	SER
3	B	613	LEU
3	B	624	LEU
3	B	632	VAL
2	C	7	SER
2	C	8	VAL
2	C	10	THR
2	C	12	ASN
2	C	23	THR
2	C	27	THR
2	C	40	VAL
2	C	41	LEU
2	C	42	ASN
3	D	46	LYS
3	D	47	LEU
3	D	48	TRP
3	D	58	ILE
3	D	63	ILE
3	D	76	GLU
3	D	78	MET

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Mol	Chain	Res	Type
3	D	79	ARG
3	D	80	GLU
3	D	95	LEU
3	D	100	LYS
3	D	115	THR
3	D	116	SER
3	D	122	THR
3	D	125	ASP
3	D	142	ILE
3	D	143	MET
3	D	146	GLN
3	D	148	THR
3	D	152	ARG
3	D	153	MET
3	D	158	LEU
3	D	161	LYS
3	D	171	THR
3	D	173	ASP
3	D	182	ASN
3	D	193	ILE
3	D	198	ASN
3	D	199	LEU
3	D	206	GLU
3	D	218	LEU
3	D	225	THR
3	D	226	LYS
3	D	228	PRO
3	D	248	GLU
3	D	269	LYS
3	D	271	LEU
3	D	272	ASP
3	D	273	LEU
3	D	274	ILE
3	D	278	THR
3	D	309	GLU
3	D	311	ASN
3	D	315	LEU
3	D	319	THR
3	D	330	GLN
3	D	336	THR
3	D	341	MET
3	D	344	MET

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Mol	Chain	Res	Type
3	D	349	LEU
3	D	352	ASP
3	D	356	THR
3	D	360	TYR
3	D	374	VAL
3	D	385	VAL
3	D	390	ASN
3	D	409	ILE
3	D	418	ASN
3	D	419	MET
3	D	420	LEU
3	D	422	LEU
3	D	428	ARG
3	D	431	ILE
3	D	440	LEU
3	D	443	PHE
3	D	453	ASN
3	D	456	ASN
3	D	464	LEU
3	D	466	VAL
3	D	470	LEU
3	D	471	LEU
3	D	493	GLN
3	D	494	THR
3	D	496	GLU
3	D	502	LEU
3	D	515	ASN
3	D	523	ASP
3	D	527	VAL
3	D	536	THR
3	D	548	LYS
3	D	549	LEU
3	D	565	LEU
3	D	588	ILE
3	D	606	THR
3	D	610	THR
3	D	612	SER
3	D	613	LEU
3	D	624	LEU
3	D	632	VAL
2	E	7	SER
2	E	8	VAL

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Mol	Chain	Res	Type
2	E	10	THR
2	E	12	ASN
2	E	23	THR
2	E	27	THR
2	E	40	VAL
2	E	41	LEU
2	E	42	ASN
3	F	46	LYS
3	F	47	LEU
3	F	48	TRP
3	F	51	VAL
3	F	76	GLU
3	F	78	MET
3	F	79	ARG
3	F	80	GLU
3	F	95	LEU
3	F	100	LYS
3	F	115	THR
3	F	116	SER
3	F	122	THR
3	F	125	ASP
3	F	143	MET
3	F	146	GLN
3	F	148	THR
3	F	152	ARG
3	F	153	MET
3	F	158	LEU
3	F	171	THR
3	F	173	ASP
3	F	182	ASN
3	F	193	ILE
3	F	198	ASN
3	F	199	LEU
3	F	206	GLU
3	F	218	LEU
3	F	225	THR
3	F	226	LYS
3	F	228	PRO
3	F	248	GLU
3	F	269	LYS
3	F	271	LEU
3	F	272	ASP

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Mol	Chain	Res	Type
3	F	273	LEU
3	F	274	ILE
3	F	278	THR
3	F	309	GLU
3	F	311	ASN
3	F	315	LEU
3	F	319	THR
3	F	330	GLN
3	F	336	THR
3	F	341	MET
3	F	344	MET
3	F	349	LEU
3	F	352	ASP
3	F	356	THR
3	F	360	TYR
3	F	374	VAL
3	F	385	VAL
3	F	390	ASN
3	F	409	ILE
3	F	418	ASN
3	F	419	MET
3	F	420	LEU
3	F	422	LEU
3	F	428	ARG
3	F	431	ILE
3	F	440	LEU
3	F	443	PHE
3	F	453	ASN
3	F	456	ASN
3	F	464	LEU
3	F	466	VAL
3	F	470	LEU
3	F	471	LEU
3	F	493	GLN
3	F	494	THR
3	F	496	GLU
3	F	502	LEU
3	F	515	ASN
3	F	523	ASP
3	F	527	VAL
3	F	536	THR
3	F	548	LYS

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Mol	Chain	Res	Type
3	F	549	LEU
3	F	565	LEU
3	F	588	ILE
3	F	606	THR
3	F	610	THR
3	F	612	SER
3	F	613	LEU
3	F	624	LEU
3	F	632	VAL
2	G	7	SER
2	G	8	VAL
2	G	10	THR
2	G	12	ASN
2	G	23	THR
2	G	27	THR
2	G	40	VAL
2	G	41	LEU
2	G	42	ASN
3	H	46	LYS
3	H	47	LEU
3	H	48	TRP
3	H	51	VAL
3	H	76	GLU
3	H	78	MET
3	H	79	ARG
3	H	80	GLU
3	H	95	LEU
3	H	100	LYS
3	H	115	THR
3	H	116	SER
3	H	122	THR
3	H	125	ASP
3	H	143	MET
3	H	146	GLN
3	H	148	THR
3	H	152	ARG
3	H	153	MET
3	H	158	LEU
3	H	171	THR
3	H	173	ASP
3	H	182	ASN
3	H	193	ILE

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Mol	Chain	Res	Type
3	H	198	ASN
3	H	199	LEU
3	H	206	GLU
3	H	218	LEU
3	H	225	THR
3	H	226	LYS
3	H	228	PRO
3	H	248	GLU
3	H	269	LYS
3	H	271	LEU
3	H	272	ASP
3	H	273	LEU
3	H	274	ILE
3	H	278	THR
3	H	309	GLU
3	H	311	ASN
3	H	315	LEU
3	H	319	THR
3	H	330	GLN
3	H	336	THR
3	H	341	MET
3	H	344	MET
3	H	349	LEU
3	H	352	ASP
3	H	356	THR
3	H	360	TYR
3	H	374	VAL
3	H	385	VAL
3	H	390	ASN
3	H	409	ILE
3	H	418	ASN
3	H	419	MET
3	H	420	LEU
3	H	422	LEU
3	H	428	ARG
3	H	431	ILE
3	H	440	LEU
3	H	443	PHE
3	H	453	ASN
3	H	456	ASN
3	H	464	LEU
3	H	466	VAL

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Mol	Chain	Res	Type
3	H	470	LEU
3	H	471	LEU
3	H	493	GLN
3	H	494	THR
3	H	496	GLU
3	H	502	LEU
3	H	515	ASN
3	H	523	ASP
3	H	527	VAL
3	H	536	THR
3	H	548	LYS
3	H	549	LEU
3	H	565	LEU
3	H	588	ILE
3	H	606	THR
3	H	610	THR
3	H	612	SER
3	H	613	LEU
3	H	624	LEU
3	H	632	VAL
2	I	7	SER
2	I	8	VAL
2	I	10	THR
2	I	12	ASN
2	I	23	THR
2	I	27	THR
2	I	40	VAL
2	I	41	LEU
2	I	42	ASN
3	J	46	LYS
3	J	47	LEU
3	J	48	TRP
3	J	60	SER
3	J	76	GLU
3	J	78	MET
3	J	79	ARG
3	J	80	GLU
3	J	95	LEU
3	J	100	LYS
3	J	115	THR
3	J	116	SER
3	J	122	THR

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Mol	Chain	Res	Type
3	J	125	ASP
3	J	143	MET
3	J	146	GLN
3	J	148	THR
3	J	152	ARG
3	J	153	MET
3	J	158	LEU
3	J	161	LYS
3	J	171	THR
3	J	173	ASP
3	J	182	ASN
3	J	193	ILE
3	J	198	ASN
3	J	199	LEU
3	J	206	GLU
3	J	218	LEU
3	J	225	THR
3	J	226	LYS
3	J	228	PRO
3	J	248	GLU
3	J	269	LYS
3	J	271	LEU
3	J	272	ASP
3	J	273	LEU
3	J	274	ILE
3	J	278	THR
3	J	309	GLU
3	J	311	ASN
3	J	315	LEU
3	J	319	THR
3	J	323	SER
3	J	330	GLN
3	J	336	THR
3	J	341	MET
3	J	344	MET
3	J	349	LEU
3	J	352	ASP
3	J	356	THR
3	J	360	TYR
3	J	374	VAL
3	J	385	VAL
3	J	390	ASN

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Mol	Chain	Res	Type
3	J	406	ASN
3	J	409	ILE
3	J	418	ASN
3	J	419	MET
3	J	420	LEU
3	J	422	LEU
3	J	428	ARG
3	J	431	ILE
3	J	440	LEU
3	J	443	PHE
3	J	453	ASN
3	J	456	ASN
3	J	464	LEU
3	J	466	VAL
3	J	470	LEU
3	J	471	LEU
3	J	493	GLN
3	J	494	THR
3	J	496	GLU
3	J	502	LEU
3	J	515	ASN
3	J	523	ASP
3	J	527	VAL
3	J	536	THR
3	J	548	LYS
3	J	549	LEU
3	J	565	LEU
3	J	588	ILE
3	J	606	THR
3	J	610	THR
3	J	612	SER
3	J	613	LEU
3	J	624	LEU
3	J	632	VAL
2	K	7	SER
2	K	8	VAL
2	K	10	THR
2	K	12	ASN
2	K	23	THR
2	K	27	THR
2	K	40	VAL
2	K	41	LEU

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Mol	Chain	Res	Type
2	K	42	ASN
3	L	46	LYS
3	L	47	LEU
3	L	48	TRP
3	L	51	VAL
3	L	76	GLU
3	L	78	MET
3	L	79	ARG
3	L	80	GLU
3	L	95	LEU
3	L	100	LYS
3	L	115	THR
3	L	116	SER
3	L	122	THR
3	L	125	ASP
3	L	142	ILE
3	L	143	MET
3	L	146	GLN
3	L	148	THR
3	L	152	ARG
3	L	153	MET
3	L	158	LEU
3	L	171	THR
3	L	173	ASP
3	L	182	ASN
3	L	193	ILE
3	L	198	ASN
3	L	199	LEU
3	L	206	GLU
3	L	218	LEU
3	L	225	THR
3	L	226	LYS
3	L	228	PRO
3	L	248	GLU
3	L	269	LYS
3	L	271	LEU
3	L	272	ASP
3	L	273	LEU
3	L	274	ILE
3	L	278	THR
3	L	309	GLU
3	L	311	ASN

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Mol	Chain	Res	Type
3	L	315	LEU
3	L	319	THR
3	L	330	GLN
3	L	336	THR
3	L	341	MET
3	L	344	MET
3	L	349	LEU
3	L	352	ASP
3	L	356	THR
3	L	360	TYR
3	L	374	VAL
3	L	385	VAL
3	L	390	ASN
3	L	409	ILE
3	L	418	ASN
3	L	419	MET
3	L	420	LEU
3	L	422	LEU
3	L	428	ARG
3	L	431	ILE
3	L	440	LEU
3	L	443	PHE
3	L	453	ASN
3	L	456	ASN
3	L	464	LEU
3	L	466	VAL
3	L	470	LEU
3	L	471	LEU
3	L	493	GLN
3	L	494	THR
3	L	496	GLU
3	L	502	LEU
3	L	515	ASN
3	L	523	ASP
3	L	527	VAL
3	L	536	THR
3	L	548	LYS
3	L	549	LEU
3	L	565	LEU
3	L	588	ILE
3	L	606	THR
3	L	610	THR

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Mol	Chain	Res	Type
3	L	612	SER
3	L	613	LEU
3	L	624	LEU
3	L	632	VAL
2	M	7	SER
2	M	8	VAL
2	M	10	THR
2	M	12	ASN
2	M	23	THR
2	M	27	THR
2	M	40	VAL
2	M	41	LEU
2	M	42	ASN
3	N	46	LYS
3	N	47	LEU
3	N	48	TRP
3	N	51	VAL
3	N	76	GLU
3	N	78	MET
3	N	79	ARG
3	N	80	GLU
3	N	95	LEU
3	N	100	LYS
3	N	115	THR
3	N	116	SER
3	N	122	THR
3	N	125	ASP
3	N	142	ILE
3	N	143	MET
3	N	146	GLN
3	N	148	THR
3	N	153	MET
3	N	158	LEU
3	N	161	LYS
3	N	171	THR
3	N	173	ASP
3	N	182	ASN
3	N	193	ILE
3	N	198	ASN
3	N	199	LEU
3	N	206	GLU
3	N	218	LEU

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Mol	Chain	Res	Type
3	N	225	THR
3	N	226	LYS
3	N	228	PRO
3	N	248	GLU
3	N	269	LYS
3	N	271	LEU
3	N	272	ASP
3	N	273	LEU
3	N	274	ILE
3	N	278	THR
3	N	309	GLU
3	N	311	ASN
3	N	315	LEU
3	N	319	THR
3	N	322	TYR
3	N	330	GLN
3	N	336	THR
3	N	341	MET
3	N	344	MET
3	N	349	LEU
3	N	352	ASP
3	N	356	THR
3	N	360	TYR
3	N	374	VAL
3	N	385	VAL
3	N	390	ASN
3	N	409	ILE
3	N	418	ASN
3	N	419	MET
3	N	420	LEU
3	N	422	LEU
3	N	428	ARG
3	N	431	ILE
3	N	440	LEU
3	N	443	PHE
3	N	453	ASN
3	N	456	ASN
3	N	464	LEU
3	N	466	VAL
3	N	470	LEU
3	N	471	LEU
3	N	493	GLN

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Mol	Chain	Res	Type
3	N	494	THR
3	N	496	GLU
3	N	502	LEU
3	N	515	ASN
3	N	523	ASP
3	N	527	VAL
3	N	530	ARG
3	N	536	THR
3	N	548	LYS
3	N	549	LEU
3	N	565	LEU
3	N	588	ILE
3	N	606	THR
3	N	610	THR
3	N	612	SER
3	N	613	LEU
3	N	624	LEU
3	N	632	VAL
2	O	7	SER
2	O	8	VAL
2	O	10	THR
2	O	12	ASN
2	O	23	THR
2	O	27	THR
2	O	40	VAL
2	O	41	LEU
2	O	42	ASN
3	P	46	LYS
3	P	47	LEU
3	P	48	TRP
3	P	51	VAL
3	P	55	VAL
3	P	76	GLU
3	P	78	MET
3	P	79	ARG
3	P	80	GLU
3	P	95	LEU
3	P	100	LYS
3	P	115	THR
3	P	116	SER
3	P	122	THR
3	P	125	ASP

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Mol	Chain	Res	Type
3	P	127	LEU
3	P	143	MET
3	P	146	GLN
3	P	148	THR
3	P	152	ARG
3	P	153	MET
3	P	158	LEU
3	P	171	THR
3	P	173	ASP
3	P	182	ASN
3	P	193	ILE
3	P	198	ASN
3	P	199	LEU
3	P	206	GLU
3	P	218	LEU
3	P	225	THR
3	P	226	LYS
3	P	228	PRO
3	P	248	GLU
3	P	269	LYS
3	P	271	LEU
3	P	272	ASP
3	P	273	LEU
3	P	274	ILE
3	P	278	THR
3	P	309	GLU
3	P	311	ASN
3	P	315	LEU
3	P	319	THR
3	P	330	GLN
3	P	336	THR
3	P	341	MET
3	P	344	MET
3	P	349	LEU
3	P	352	ASP
3	P	356	THR
3	P	360	TYR
3	P	374	VAL
3	P	385	VAL
3	P	390	ASN
3	P	409	ILE
3	P	418	ASN

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Mol	Chain	Res	Type
3	P	419	MET
3	P	420	LEU
3	P	422	LEU
3	P	428	ARG
3	P	431	ILE
3	P	440	LEU
3	P	443	PHE
3	P	453	ASN
3	P	456	ASN
3	P	464	LEU
3	P	466	VAL
3	P	470	LEU
3	P	471	LEU
3	P	472	ARG
3	P	493	GLN
3	P	494	THR
3	P	496	GLU
3	P	502	LEU
3	P	515	ASN
3	P	523	ASP
3	P	527	VAL
3	P	536	THR
3	P	548	LYS
3	P	549	LEU
3	P	565	LEU
3	P	588	ILE
3	P	606	THR
3	P	610	THR
3	P	612	SER
3	P	613	LEU
3	P	624	LEU
3	P	632	VAL
2	Q	7	SER
2	Q	8	VAL
2	Q	10	THR
2	Q	12	ASN
2	Q	23	THR
2	Q	27	THR
2	Q	40	VAL
2	Q	41	LEU
2	Q	42	ASN
3	R	46	LYS

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Mol	Chain	Res	Type
3	R	47	LEU
3	R	48	TRP
3	R	51	VAL
3	R	76	GLU
3	R	78	MET
3	R	79	ARG
3	R	80	GLU
3	R	95	LEU
3	R	100	LYS
3	R	115	THR
3	R	116	SER
3	R	122	THR
3	R	125	ASP
3	R	143	MET
3	R	146	GLN
3	R	148	THR
3	R	152	ARG
3	R	153	MET
3	R	158	LEU
3	R	171	THR
3	R	173	ASP
3	R	182	ASN
3	R	193	ILE
3	R	198	ASN
3	R	199	LEU
3	R	206	GLU
3	R	218	LEU
3	R	225	THR
3	R	226	LYS
3	R	228	PRO
3	R	248	GLU
3	R	269	LYS
3	R	271	LEU
3	R	272	ASP
3	R	273	LEU
3	R	274	ILE
3	R	278	THR
3	R	309	GLU
3	R	311	ASN
3	R	315	LEU
3	R	319	THR
3	R	330	GLN

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Mol	Chain	Res	Type
3	R	336	THR
3	R	341	MET
3	R	344	MET
3	R	349	LEU
3	R	352	ASP
3	R	356	THR
3	R	360	TYR
3	R	374	VAL
3	R	385	VAL
3	R	390	ASN
3	R	409	ILE
3	R	418	ASN
3	R	419	MET
3	R	420	LEU
3	R	422	LEU
3	R	428	ARG
3	R	431	ILE
3	R	440	LEU
3	R	443	PHE
3	R	453	ASN
3	R	456	ASN
3	R	464	LEU
3	R	466	VAL
3	R	470	LEU
3	R	471	LEU
3	R	493	GLN
3	R	494	THR
3	R	496	GLU
3	R	502	LEU
3	R	515	ASN
3	R	523	ASP
3	R	527	VAL
3	R	530	ARG
3	R	536	THR
3	R	548	LYS
3	R	549	LEU
3	R	565	LEU
3	R	588	ILE
3	R	606	THR
3	R	610	THR
3	R	612	SER
3	R	613	LEU

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Mol	Chain	Res	Type
3	R	624	LEU
3	R	632	VAL
2	S	7	SER
2	S	8	VAL
2	S	10	THR
2	S	12	ASN
2	S	23	THR
2	S	27	THR
2	S	40	VAL
2	S	41	LEU
2	S	42	ASN
3	T	46	LYS
3	T	47	LEU
3	T	48	TRP
3	T	51	VAL
3	T	76	GLU
3	T	78	MET
3	T	79	ARG
3	T	80	GLU
3	T	95	LEU
3	T	100	LYS
3	T	101	THR
3	T	115	THR
3	T	116	SER
3	T	122	THR
3	T	125	ASP
3	T	142	ILE
3	T	143	MET
3	T	146	GLN
3	T	148	THR
3	T	152	ARG
3	T	153	MET
3	T	158	LEU
3	T	171	THR
3	T	173	ASP
3	T	182	ASN
3	T	193	ILE
3	T	198	ASN
3	T	199	LEU
3	T	206	GLU
3	T	218	LEU
3	T	225	THR

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Mol	Chain	Res	Type
3	T	226	LYS
3	T	228	PRO
3	T	248	GLU
3	T	269	LYS
3	T	271	LEU
3	T	272	ASP
3	T	273	LEU
3	T	274	ILE
3	T	278	THR
3	T	309	GLU
3	T	311	ASN
3	T	315	LEU
3	T	319	THR
3	T	330	GLN
3	T	336	THR
3	T	341	MET
3	T	344	MET
3	T	349	LEU
3	T	352	ASP
3	T	356	THR
3	T	360	TYR
3	T	374	VAL
3	T	385	VAL
3	T	390	ASN
3	T	406	ASN
3	T	409	ILE
3	T	418	ASN
3	T	419	MET
3	T	420	LEU
3	T	422	LEU
3	T	428	ARG
3	T	431	ILE
3	T	440	LEU
3	T	443	PHE
3	T	453	ASN
3	T	456	ASN
3	T	464	LEU
3	T	466	VAL
3	T	470	LEU
3	T	471	LEU
3	T	493	GLN
3	T	494	THR

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Mol	Chain	Res	Type
3	T	496	GLU
3	T	502	LEU
3	T	515	ASN
3	T	523	ASP
3	T	527	VAL
3	T	536	THR
3	T	548	LYS
3	T	549	LEU
3	T	565	LEU
3	T	588	ILE
3	T	606	THR
3	T	610	THR
3	T	612	SER
3	T	613	LEU
3	T	624	LEU
3	T	632	VAL
4	U	16	GLN
4	U	32	CYS
4	U	35	ARG
4	U	41	PHE
4	U	83	GLN
4	U	93	ARG
4	U	123	ASN
4	U	139	ASN
4	U	161	VAL
4	U	164	ASP
4	U	198	GLN
4	U	208	ILE
4	U	211	LEU
4	U	245	GLN
4	U	263	GLN
4	U	285	LEU
4	U	291	THR
4	U	294	THR
4	U	297	PRO
4	U	305	LEU
4	V	4	ARG
4	V	11	TYR
4	V	16	GLN
4	V	23	LEU
4	V	32	CYS
4	V	47	SER

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Mol	Chain	Res	Type
4	V	72	LEU
4	V	95	PRO
4	V	124	ASN
4	V	138	ILE
4	V	161	VAL
4	V	182	ILE
4	V	191	ARG
4	V	196	LEU
4	V	224	ASN
4	V	238	LYS
4	V	252	ASP
4	V	288	GLN
4	V	396	GLN
5	W	18	THR
5	W	19	ARG
5	W	30	LEU
5	W	47	LEU
5	W	53	ASP
5	W	79	LEU
5	W	80	ASP
5	W	88	ILE
5	W	99	LEU
5	W	115	PRO
5	W	136	ASP
5	W	138	LEU
5	W	145	ASP
5	W	178	ILE
5	W	196	LYS
5	W	217	LEU
5	W	219	ASN
5	W	220	LEU
5	W	224	PRO
5	W	243	GLN
5	W	257	GLN
5	W	262	MET
5	W	264	GLU
5	W	266	VAL
5	W	285	LEU
5	W	313	LEU
5	W	318	VAL
5	W	319	GLN
5	W	332	ASP

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Mol	Chain	Res	Type
5	W	337	VAL
5	W	340	LEU
5	W	347	GLN
5	W	385	THR
5	W	396	ARG
5	W	400	LEU
5	W	406	ASN
5	W	413	GLN
5	W	415	ASP
5	W	418	TYR
5	W	423	LEU
5	W	424	SER
5	W	428	THR
5	W	444	ASP
5	W	448	ASP
5	W	490	ASP
5	W	507	ASN
5	W	519	LEU
5	W	536	ASP
5	W	557	ASP
5	W	566	LEU
5	W	577	LEU
5	W	591	ASP
5	W	596	ASN
5	W	605	THR
5	W	608	GLN
5	W	621	ASN
5	W	625	ARG
5	W	627	PHE
5	W	629	THR
5	W	637	THR
5	W	641	THR
5	W	643	HIS
5	W	654	GLU
5	W	657	LEU
5	W	672	THR
5	W	679	LEU
5	W	690	ARG
5	W	732	LEU
5	W	734	SER
5	W	742	SER
5	W	746	PHE

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Mol	Chain	Res	Type
5	W	765	PHE
5	W	803	GLU
5	W	824	ARG
5	W	832	LEU
5	W	845	LEU
5	W	854	MET
5	W	894	THR
5	W	897	PHE
5	W	931	LEU
5	W	934	ASN
5	W	935	THR
5	W	938	TYR
5	W	969	ASP
5	W	971	VAL
5	W	980	LEU
5	W	1031	THR
5	W	1060	VAL
5	W	1064	GLN
5	W	1065	HIS
5	W	1070	VAL
5	W	1096	PHE
5	W	1104	LEU
5	W	1134	VAL
5	W	1136	LEU
5	W	1146	PHE
5	W	1151	ASN
5	W	1177	ASN
5	W	1215	LEU
5	W	1221	PHE
5	W	1235	THR
5	W	1286	THR
5	W	1289	LEU
6	X	194	ILE
6	X	195	ILE
6	X	200	LEU
6	X	201	CYS
6	X	206	LEU
6	X	208	ASP
6	X	215	LEU
6	X	219	LEU
6	X	223	THR
6	X	245	CYS

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Mol	Chain	Res	Type
6	X	247	THR
6	X	254	ASP
6	X	261	VAL
6	X	262	ARG
6	X	263	LEU
6	X	276	THR
6	X	277	VAL
6	X	282	TYR
6	X	283	ILE
6	X	284	ASP
6	X	287	GLN
6	X	291	LEU
6	X	298	LEU
6	X	302	LEU
6	X	303	GLN
6	X	316	THR
6	X	320	ASP
6	X	323	THR
6	X	330	ILE
6	X	342	LEU
6	X	350	LEU
6	X	376	LEU
6	X	386	GLU
6	X	388	MET
6	X	399	ARG
6	X	400	GLU
6	X	406	LEU
6	X	416	ILE
6	X	423	CYS
6	X	425	ASN
6	X	429	ASN
6	X	432	ILE
6	X	438	PRO
6	X	455	GLN
6	X	473	VAL
6	X	476	LEU
6	X	480	LEU
6	X	487	VAL
6	X	488	LEU
6	X	500	ILE
6	X	506	GLN
6	X	531	CYS

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Mol	Chain	Res	Type
6	X	540	LEU
6	X	545	LEU
6	X	552	ASP
6	X	575	MET
6	X	576	THR
6	X	583	VAL
6	X	587	MET
6	X	593	LEU
6	X	597	GLU
6	X	603	VAL
6	X	614	GLN
6	X	626	LEU
6	X	634	GLN
6	X	644	GLU
6	X	652	GLN
6	X	655	GLU
6	X	656	PHE
6	X	661	THR
6	X	662	LEU
6	X	692	THR
6	X	695	SER
6	X	699	ASN
6	X	725	THR
6	X	737	ARG
6	X	747	ILE
6	X	748	GLN
6	X	750	MET
6	X	761	VAL
6	X	764	THR
6	X	776	GLN
6	X	778	ASP
6	X	784	LEU
6	X	787	ARG
6	X	795	ASP
6	X	799	ASP
6	X	800	ILE
6	X	801	ARG
6	X	831	LEU
6	X	839	LYS
6	X	844	LEU
6	X	845	PHE
6	X	862	GLU

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Mol	Chain	Res	Type
6	X	873	GLN
6	X	880	LEU
6	X	883	ARG
6	X	886	ASP
6	X	889	ARG
6	X	893	VAL
6	X	907	ASP
6	X	914	ASP
6	X	915	LEU
6	X	919	LEU
6	X	920	LEU
6	X	939	LEU
6	X	945	VAL
6	X	949	HIS
6	X	950	VAL
6	X	963	GLN
6	X	967	THR
6	X	971	GLU
6	X	979	ARG
6	X	982	LEU
6	X	985	VAL
6	X	998	PRO
6	X	1006	ARG
6	X	1010	ASN
6	X	1012	ARG
6	X	1019	THR
6	X	1030	GLN
6	X	1031	LEU
6	X	1033	ASP
6	X	1044	ASP
6	X	1047	TYR
6	X	1066	MET
6	X	1074	VAL
6	X	1075	LEU
6	X	1078	LEU
6	X	1094	LEU
6	X	1095	ASP
6	X	1100	LEU
6	X	1106	LYS
6	X	1112	ILE
6	X	1118	CYS
6	X	1140	GLU

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Mol	Chain	Res	Type
6	X	1144	ASP
6	X	1150	ASN
6	X	1155	ASP
6	X	1156	THR
6	X	1160	GLU
6	X	1200	THR
6	X	1204	LEU
6	Y	25	LYS
6	Y	31	ASN
6	Y	37	THR
6	Y	39	PRO
6	Y	61	ASN
6	Y	73	ASN
6	Y	78	ILE
6	Y	79	ILE
6	Y	82	PRO
6	Y	85	ASP
6	Y	109	THR
6	Y	118	VAL
6	Y	121	VAL
6	Y	129	MET
6	Y	134	GLU
6	Y	158	ARG
6	Y	160	PHE
6	Y	166	ASP
6	Y	207	LEU
6	Y	215	LEU
6	Y	226	MET
6	Y	240	VAL
6	Y	241	ASN
6	Y	245	CYS
6	Y	247	THR
6	Y	257	ARG
6	Y	283	ILE
6	Y	287	GLN
6	Y	299	LEU
6	Y	306	LEU
6	Y	310	LYS
6	Y	317	PHE
6	Y	320	ASP
6	Y	328	ARG
6	Y	353	GLN

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Mol	Chain	Res	Type
6	Y	356	ASN
6	Y	390	LEU
6	Y	392	ASP
6	Y	403	ILE
6	Y	453	LEU
6	Y	466	THR
6	Y	479	LEU
6	Y	480	LEU
6	Y	481	ARG
6	Y	486	LEU
6	Y	487	VAL
6	Y	497	ILE
6	Y	499	THR
6	Y	500	ILE
6	Y	524	ASP
6	Y	531	CYS
6	Y	538	ASN
6	Y	541	VAL
6	Y	547	GLU
6	Y	552	ASP
6	Y	567	LEU
6	Y	570	THR
6	Y	593	LEU
6	Y	649	ASN
6	Y	662	LEU
6	Y	686	ARG
6	Y	689	VAL
6	Y	721	VAL
6	Y	724	LEU
6	Y	739	VAL
6	Y	743	THR
6	Y	747	ILE
6	Y	753	THR
6	Y	758	LEU
6	Y	763	VAL
6	Y	773	ARG
6	Y	778	ASP
6	Y	779	VAL
6	Y	789	ASP
6	Y	811	LEU
6	Y	829	THR
6	Y	848	ASN

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Mol	Chain	Res	Type
6	Y	889	ARG
6	Y	906	ASN
6	Y	915	LEU
6	Y	939	LEU
6	Y	954	PRO
6	Y	982	LEU
6	Y	985	VAL
6	Y	987	THR
6	Y	990	ASN
6	Y	996	MET
6	Y	1010	ASN
6	Y	1011	VAL
6	Y	1012	ARG
6	Y	1034	ASP
6	Y	1037	VAL
6	Y	1063	VAL
6	Y	1078	LEU
6	Y	1085	LEU
6	Y	1086	HIS
6	Y	1100	LEU
6	Y	1116	PRO
6	Y	1123	GLN
6	Y	1144	ASP
6	Y	1146	LEU
6	Y	1149	THR
6	Y	1154	ILE
6	Y	1156	THR
6	Y	1160	GLU
6	Y	1169	TRP
6	Y	1184	LEU
6	Y	1204	LEU
6	Y	1209	TYR
6	Y	1211	ARG
6	Y	1214	THR

Sometimes sidechains can be flipped to improve hydrogen bonding and reduce clashes. All (433) such sidechains are listed below:

Mol	Chain	Res	Type
1	l	37	HIS
1	l	59	HIS
1	b	37	HIS
1	b	59	HIS

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Mol	Chain	Res	Type
1	b	61	ASN
1	f	37	HIS
1	d	37	HIS
1	d	59	HIS
1	h	37	HIS
1	j	8	GLN
1	j	37	HIS
1	n	8	GLN
1	n	37	HIS
1	n	74	GLN
1	p	37	HIS
1	p	61	ASN
1	p	74	GLN
1	r	4	HIS
1	r	37	HIS
1	r	59	HIS
1	r	74	GLN
1	t	37	HIS
1	t	59	HIS
1	t	61	ASN
2	A	9	ASN
2	A	42	ASN
3	B	144	ASN
3	B	160	GLN
3	B	162	HIS
3	B	209	ASN
3	B	233	HIS
3	B	311	ASN
3	B	312	ASN
3	B	330	GLN
3	B	383	GLN
3	B	390	ASN
3	B	406	ASN
3	B	410	ASN
3	B	418	ASN
3	B	421	HIS
3	B	423	GLN
3	B	453	ASN
3	B	456	ASN
3	B	492	ASN
3	B	504	ASN
3	B	512	ASN

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Mol	Chain	Res	Type
3	B	515	ASN
3	B	579	ASN
3	B	585	GLN
3	B	635	GLN
2	C	9	ASN
3	D	144	ASN
3	D	160	GLN
3	D	162	HIS
3	D	209	ASN
3	D	233	HIS
3	D	311	ASN
3	D	312	ASN
3	D	330	GLN
3	D	383	GLN
3	D	390	ASN
3	D	406	ASN
3	D	410	ASN
3	D	418	ASN
3	D	421	HIS
3	D	423	GLN
3	D	453	ASN
3	D	456	ASN
3	D	492	ASN
3	D	504	ASN
3	D	512	ASN
3	D	515	ASN
3	D	579	ASN
3	D	585	GLN
3	D	635	GLN
3	D	640	GLN
2	E	9	ASN
3	F	144	ASN
3	F	160	GLN
3	F	162	HIS
3	F	209	ASN
3	F	233	HIS
3	F	311	ASN
3	F	312	ASN
3	F	330	GLN
3	F	383	GLN
3	F	390	ASN
3	F	410	ASN

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Mol	Chain	Res	Type
3	F	418	ASN
3	F	421	HIS
3	F	423	GLN
3	F	453	ASN
3	F	456	ASN
3	F	492	ASN
3	F	504	ASN
3	F	512	ASN
3	F	515	ASN
3	F	579	ASN
3	F	585	GLN
3	F	635	GLN
3	F	640	GLN
3	H	144	ASN
3	H	160	GLN
3	H	162	HIS
3	H	209	ASN
3	H	233	HIS
3	H	311	ASN
3	H	312	ASN
3	H	330	GLN
3	H	383	GLN
3	H	390	ASN
3	H	410	ASN
3	H	418	ASN
3	H	423	GLN
3	H	453	ASN
3	H	456	ASN
3	H	492	ASN
3	H	504	ASN
3	H	512	ASN
3	H	515	ASN
3	H	579	ASN
3	H	585	GLN
3	H	635	GLN
3	H	640	GLN
2	I	9	ASN
3	J	144	ASN
3	J	160	GLN
3	J	162	HIS
3	J	209	ASN
3	J	233	HIS

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Mol	Chain	Res	Type
3	J	311	ASN
3	J	312	ASN
3	J	330	GLN
3	J	383	GLN
3	J	390	ASN
3	J	410	ASN
3	J	418	ASN
3	J	421	HIS
3	J	423	GLN
3	J	453	ASN
3	J	456	ASN
3	J	492	ASN
3	J	504	ASN
3	J	512	ASN
3	J	515	ASN
3	J	579	ASN
3	J	585	GLN
3	J	635	GLN
3	J	640	GLN
2	K	9	ASN
3	L	144	ASN
3	L	160	GLN
3	L	162	HIS
3	L	209	ASN
3	L	231	GLN
3	L	233	HIS
3	L	311	ASN
3	L	312	ASN
3	L	330	GLN
3	L	383	GLN
3	L	390	ASN
3	L	410	ASN
3	L	418	ASN
3	L	421	HIS
3	L	423	GLN
3	L	453	ASN
3	L	456	ASN
3	L	492	ASN
3	L	504	ASN
3	L	512	ASN
3	L	515	ASN
3	L	579	ASN

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Mol	Chain	Res	Type
3	L	585	GLN
3	L	635	GLN
3	L	640	GLN
3	N	144	ASN
3	N	160	GLN
3	N	209	ASN
3	N	233	HIS
3	N	311	ASN
3	N	312	ASN
3	N	330	GLN
3	N	383	GLN
3	N	390	ASN
3	N	410	ASN
3	N	418	ASN
3	N	421	HIS
3	N	423	GLN
3	N	453	ASN
3	N	456	ASN
3	N	492	ASN
3	N	504	ASN
3	N	512	ASN
3	N	515	ASN
3	N	579	ASN
3	N	585	GLN
3	N	635	GLN
3	N	640	GLN
2	O	9	ASN
3	P	144	ASN
3	P	160	GLN
3	P	162	HIS
3	P	209	ASN
3	P	233	HIS
3	P	311	ASN
3	P	312	ASN
3	P	330	GLN
3	P	383	GLN
3	P	390	ASN
3	P	410	ASN
3	P	418	ASN
3	P	421	HIS
3	P	423	GLN
3	P	453	ASN

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Mol	Chain	Res	Type
3	P	456	ASN
3	P	492	ASN
3	P	504	ASN
3	P	512	ASN
3	P	515	ASN
3	P	579	ASN
3	P	585	GLN
3	P	635	GLN
3	P	640	GLN
2	Q	3	ASN
2	Q	9	ASN
3	R	144	ASN
3	R	160	GLN
3	R	162	HIS
3	R	209	ASN
3	R	233	HIS
3	R	311	ASN
3	R	312	ASN
3	R	330	GLN
3	R	383	GLN
3	R	390	ASN
3	R	410	ASN
3	R	418	ASN
3	R	423	GLN
3	R	453	ASN
3	R	456	ASN
3	R	492	ASN
3	R	504	ASN
3	R	512	ASN
3	R	515	ASN
3	R	579	ASN
3	R	585	GLN
3	R	628	GLN
3	R	635	GLN
3	R	640	GLN
3	T	144	ASN
3	T	160	GLN
3	T	162	HIS
3	T	209	ASN
3	T	233	HIS
3	T	311	ASN
3	T	312	ASN

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Mol	Chain	Res	Type
3	T	330	GLN
3	T	383	GLN
3	T	390	ASN
3	T	410	ASN
3	T	418	ASN
3	T	423	GLN
3	T	453	ASN
3	T	456	ASN
3	T	492	ASN
3	T	504	ASN
3	T	512	ASN
3	T	515	ASN
3	T	579	ASN
3	T	585	GLN
3	T	635	GLN
3	T	640	GLN
4	U	3	GLN
4	U	12	ASN
4	U	24	ASN
4	U	27	GLN
4	U	43	HIS
4	U	69	ASN
4	U	75	HIS
4	U	124	ASN
4	U	140	GLN
4	U	158	HIS
4	U	198	GLN
4	U	224	ASN
4	U	242	ASN
4	U	245	GLN
4	U	251	ASN
4	U	263	GLN
4	U	275	GLN
4	U	276	HIS
4	U	310	GLN
4	U	314	ASN
4	U	343	GLN
4	U	357	ASN
4	U	377	ASN
4	U	387	GLN
4	V	3	GLN
4	V	12	ASN

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Mol	Chain	Res	Type
4	V	65	ASN
4	V	87	HIS
4	V	88	GLN
4	V	124	ASN
4	V	140	GLN
4	V	158	HIS
4	V	184	HIS
4	V	187	GLN
4	V	195	GLN
4	V	242	ASN
4	V	245	GLN
4	V	277	GLN
4	V	310	GLN
4	V	338	HIS
4	V	351	GLN
4	V	354	GLN
4	V	357	ASN
4	V	387	GLN
4	V	396	GLN
4	V	399	GLN
5	W	35	GLN
5	W	38	ASN
5	W	40	HIS
5	W	60	GLN
5	W	89	HIS
5	W	108	ASN
5	W	163	ASN
5	W	219	ASN
5	W	229	HIS
5	W	243	GLN
5	W	257	GLN
5	W	319	GLN
5	W	347	GLN
5	W	406	ASN
5	W	553	GLN
5	W	584	GLN
5	W	596	ASN
5	W	630	GLN
5	W	633	ASN
5	W	638	HIS
5	W	651	ASN
5	W	685	ASN

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Mol	Chain	Res	Type
5	W	768	ASN
5	W	788	GLN
5	W	816	ASN
5	W	830	HIS
5	W	934	ASN
5	W	1014	ASN
5	W	1038	ASN
5	W	1045	ASN
5	W	1047	GLN
5	W	1066	ASN
5	W	1151	ASN
5	W	1189	ASN
5	W	1249	HIS
5	W	1283	GLN
6	X	214	HIS
6	X	230	GLN
6	X	287	GLN
6	X	296	ASN
6	X	303	GLN
6	X	353	GLN
6	X	356	ASN
6	X	421	ASN
6	X	425	ASN
6	X	429	ASN
6	X	506	GLN
6	X	559	HIS
6	X	627	ASN
6	X	635	GLN
6	X	645	HIS
6	X	652	GLN
6	X	673	ASN
6	X	681	ASN
6	X	682	GLN
6	X	749	HIS
6	X	777	ASN
6	X	826	GLN
6	X	854	ASN
6	X	855	GLN
6	X	873	GLN
6	X	940	GLN
6	X	944	ASN
6	X	962	GLN

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Mol	Chain	Res	Type
6	X	973	ASN
6	X	986	GLN
6	X	990	ASN
6	X	1025	ASN
6	X	1053	GLN
6	X	1091	GLN
6	X	1134	HIS
6	X	1143	ASN
6	X	1150	ASN
6	X	1182	ASN
6	Y	48	GLN
6	Y	49	GLN
6	Y	57	GLN
6	Y	113	ASN
6	Y	120	ASN
6	Y	140	HIS
6	Y	308	ASN
6	Y	356	ASN
6	Y	358	GLN
6	Y	410	HIS
6	Y	415	GLN
6	Y	429	ASN
6	Y	559	HIS
6	Y	580	HIS
6	Y	614	GLN
6	Y	617	HIS
6	Y	627	ASN
6	Y	632	ASN
6	Y	652	GLN
6	Y	663	GLN
6	Y	699	ASN
6	Y	722	ASN
6	Y	748	GLN
6	Y	776	GLN
6	Y	835	HIS
6	Y	836	HIS
6	Y	848	ASN
6	Y	875	GLN
6	Y	906	ASN
6	Y	940	GLN
6	Y	944	ASN
6	Y	949	HIS

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Mol	Chain	Res	Type
6	Y	956	HIS
6	Y	973	ASN
6	Y	976	ASN
6	Y	981	ASN
6	Y	990	ASN
6	Y	1010	ASN
6	Y	1030	GLN
6	Y	1150	ASN
6	Y	1198	ASN

5.3.3 RNA (i)

There are no RNA molecules in this entry.

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

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

5.5 Carbohydrates (i)

There are no oligosaccharides in this entry.

5.6 Ligand geometry (i)

10 ligands are modelled in this entry.

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

Mol	Type	Chain	Res	Link	Bond lengths			Bond angles		
					Counts	RMSZ	$\#\|Z\| > 2$	Counts	RMSZ	$\#\|Z\| > 2$
7	MYR	A	101	2	13,14,15	0.44	0	12,13,15	0.49	0
7	MYR	C	101	2	13,14,15	0.44	0	12,13,15	0.49	0
7	MYR	E	101	2	13,14,15	0.44	0	12,13,15	0.49	0
7	MYR	O	101	2	13,14,15	0.44	0	12,13,15	0.49	0
7	MYR	S	101	2	13,14,15	0.44	0	12,13,15	0.49	0

Mol	Type	Chain	Res	Link	Bond lengths			Bond angles		
					Counts	RMSZ	$\#\|Z\|>2$	Counts	RMSZ	$\#\|Z\|>2$
7	MYR	I	101	2	13,14,15	0.44	0	12,13,15	0.49	0
7	MYR	M	101	2	13,14,15	0.44	0	12,13,15	0.49	0
7	MYR	G	101	2	13,14,15	0.44	0	12,13,15	0.49	0
7	MYR	K	101	2	13,14,15	0.44	0	12,13,15	0.49	0
7	MYR	Q	101	2	13,14,15	0.45	0	12,13,15	0.49	0

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
7	MYR	A	101	2	-	8/12/12/13	-
7	MYR	C	101	2	-	8/12/12/13	-
7	MYR	E	101	2	-	8/12/12/13	-
7	MYR	O	101	2	-	8/12/12/13	-
7	MYR	S	101	2	-	8/12/12/13	-
7	MYR	I	101	2	-	8/12/12/13	-
7	MYR	M	101	2	-	8/12/12/13	-
7	MYR	G	101	2	-	8/12/12/13	-
7	MYR	K	101	2	-	8/12/12/13	-
7	MYR	Q	101	2	-	8/12/12/13	-

There are no bond length outliers.

There are no bond angle outliers.

There are no chirality outliers.

All (80) torsion outliers are listed below:

Mol	Chain	Res	Type	Atoms
7	E	101	MYR	C3-C4-C5-C6
7	K	101	MYR	C3-C4-C5-C6
7	Q	101	MYR	C3-C4-C5-C6
7	A	101	MYR	C3-C4-C5-C6
7	C	101	MYR	C3-C4-C5-C6
7	G	101	MYR	C3-C4-C5-C6
7	I	101	MYR	C3-C4-C5-C6
7	M	101	MYR	C3-C4-C5-C6
7	O	101	MYR	C3-C4-C5-C6

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Mol	Chain	Res	Type	Atoms
7	S	101	MYR	C3-C4-C5-C6
7	A	101	MYR	C11-C12-C13-C14
7	C	101	MYR	C11-C12-C13-C14
7	E	101	MYR	C11-C12-C13-C14
7	G	101	MYR	C11-C12-C13-C14
7	I	101	MYR	C11-C12-C13-C14
7	K	101	MYR	C11-C12-C13-C14
7	M	101	MYR	C11-C12-C13-C14
7	O	101	MYR	C11-C12-C13-C14
7	Q	101	MYR	C11-C12-C13-C14
7	S	101	MYR	C11-C12-C13-C14
7	A	101	MYR	C6-C7-C8-C9
7	K	101	MYR	C6-C7-C8-C9
7	O	101	MYR	C6-C7-C8-C9
7	C	101	MYR	C6-C7-C8-C9
7	E	101	MYR	C6-C7-C8-C9
7	G	101	MYR	C6-C7-C8-C9
7	I	101	MYR	C6-C7-C8-C9
7	M	101	MYR	C6-C7-C8-C9
7	Q	101	MYR	C6-C7-C8-C9
7	S	101	MYR	C6-C7-C8-C9
7	C	101	MYR	C2-C3-C4-C5
7	Q	101	MYR	C2-C3-C4-C5
7	A	101	MYR	C2-C3-C4-C5
7	E	101	MYR	C2-C3-C4-C5
7	G	101	MYR	C2-C3-C4-C5
7	I	101	MYR	C2-C3-C4-C5
7	K	101	MYR	C2-C3-C4-C5
7	M	101	MYR	C2-C3-C4-C5
7	O	101	MYR	C2-C3-C4-C5
7	S	101	MYR	C2-C3-C4-C5
7	G	101	MYR	C11-C10-C9-C8
7	M	101	MYR	C11-C10-C9-C8
7	A	101	MYR	C11-C10-C9-C8
7	C	101	MYR	C11-C10-C9-C8
7	E	101	MYR	C11-C10-C9-C8
7	I	101	MYR	C11-C10-C9-C8
7	K	101	MYR	C11-C10-C9-C8
7	O	101	MYR	C11-C10-C9-C8
7	Q	101	MYR	C11-C10-C9-C8
7	S	101	MYR	C11-C10-C9-C8
7	C	101	MYR	C4-C5-C6-C7

Continued on next page...

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Mol	Chain	Res	Type	Atoms
7	I	101	MYR	C4-C5-C6-C7
7	A	101	MYR	C4-C5-C6-C7
7	E	101	MYR	C4-C5-C6-C7
7	G	101	MYR	C4-C5-C6-C7
7	K	101	MYR	C4-C5-C6-C7
7	O	101	MYR	C4-C5-C6-C7
7	S	101	MYR	C4-C5-C6-C7
7	M	101	MYR	C4-C5-C6-C7
7	Q	101	MYR	C4-C5-C6-C7
7	A	101	MYR	C1-C2-C3-C4
7	C	101	MYR	C1-C2-C3-C4
7	E	101	MYR	C1-C2-C3-C4
7	G	101	MYR	C1-C2-C3-C4
7	I	101	MYR	C1-C2-C3-C4
7	K	101	MYR	C1-C2-C3-C4
7	M	101	MYR	C1-C2-C3-C4
7	O	101	MYR	C1-C2-C3-C4
7	Q	101	MYR	C1-C2-C3-C4
7	S	101	MYR	C1-C2-C3-C4
7	I	101	MYR	C5-C6-C7-C8
7	C	101	MYR	C5-C6-C7-C8
7	G	101	MYR	C5-C6-C7-C8
7	S	101	MYR	C5-C6-C7-C8
7	M	101	MYR	C5-C6-C7-C8
7	A	101	MYR	C5-C6-C7-C8
7	E	101	MYR	C5-C6-C7-C8
7	K	101	MYR	C5-C6-C7-C8
7	O	101	MYR	C5-C6-C7-C8
7	Q	101	MYR	C5-C6-C7-C8

There are no ring outliers.

10 monomers are involved in 280 short contacts:

Mol	Chain	Res	Type	Clashes	Symm-Clashes
7	A	101	MYR	27	0
7	C	101	MYR	31	0
7	E	101	MYR	27	0
7	O	101	MYR	28	0
7	S	101	MYR	30	0
7	I	101	MYR	29	0
7	M	101	MYR	26	0
7	G	101	MYR	28	0

Continued on next page...

Continued from previous page...

Mol	Chain	Res	Type	Clashes	Symm-Clashes
7	K	101	MYR	27	0
7	Q	101	MYR	27	0

5.7 Other polymers (i)

There are no such residues in this entry.

5.8 Polymer linkage issues (i)

The following chains have linkage breaks:

Mol	Chain	Number of breaks
1	b	1

All chain breaks are listed below:

Model	Chain	Residue-1	Atom-1	Residue-2	Atom-2	Distance (Å)
1	b	74:GLN	C	75:PRO	N	1.68

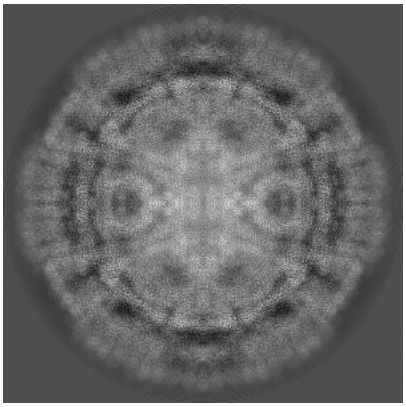
6 Map visualisation

This section contains visualisations of the EMDB entry EMD-6969. These allow visual inspection of the internal detail of the map and identification of artifacts.

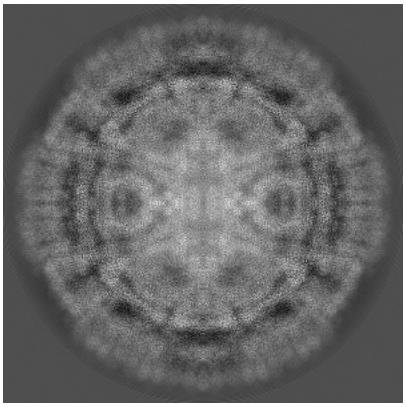
No raw map or half-maps were deposited for this entry and therefore no images, graphs, etc. pertaining to the raw map can be shown.

6.1 Orthogonal projections

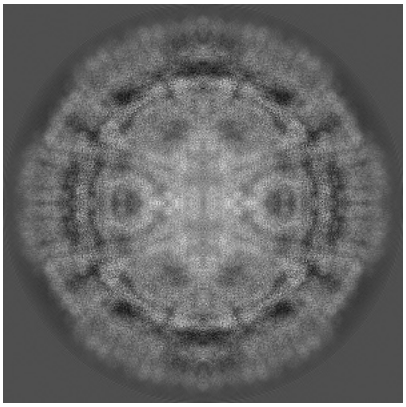
6.1.1 Primary map



X



Y

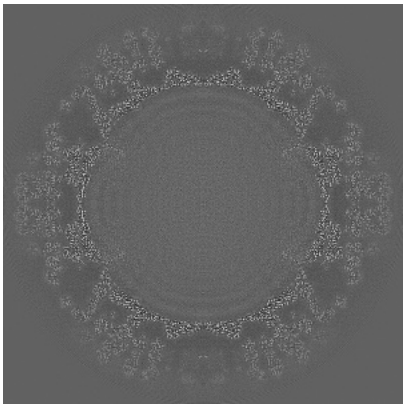


Z

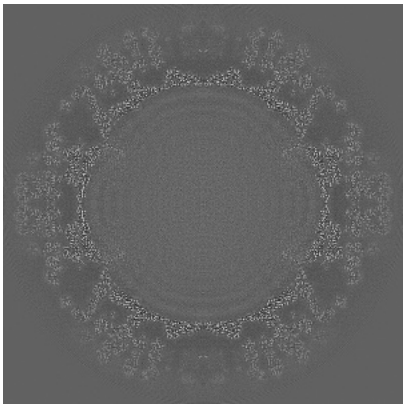
The images above show the map projected in three orthogonal directions.

6.2 Central slices

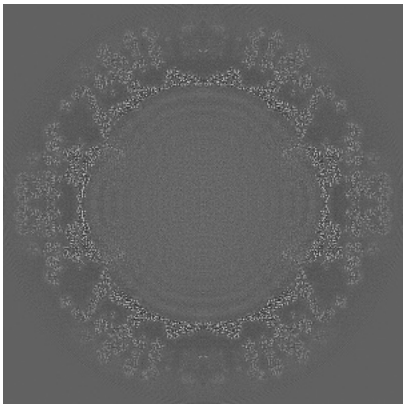
6.2.1 Primary map



X Index: 450



Y Index: 450

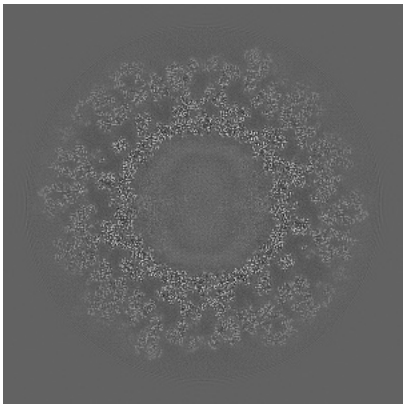


Z Index: 450

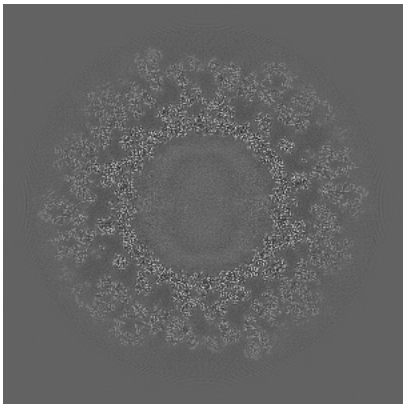
The images above show central slices of the map in three orthogonal directions.

6.3 Largest variance slices (i)

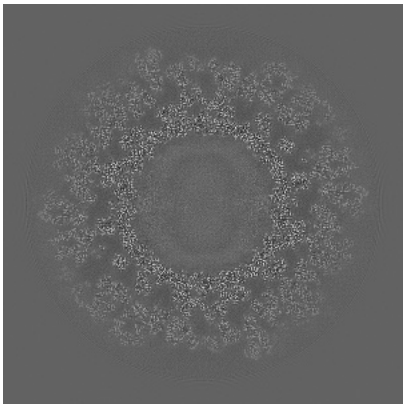
6.3.1 Primary map



X Index: 241



Y Index: 659

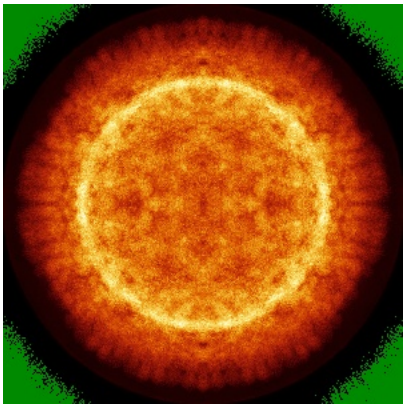


Z Index: 659

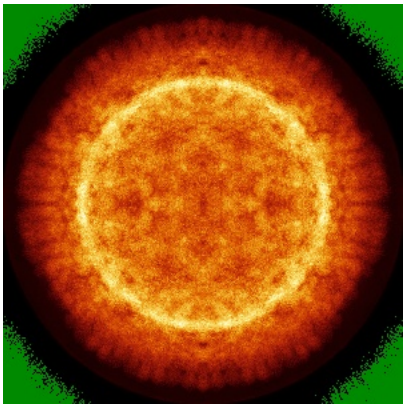
The images above show the largest variance slices of the map in three orthogonal directions.

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

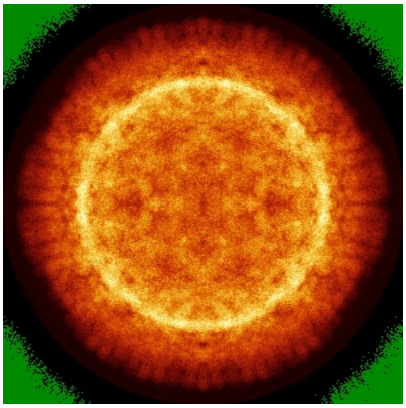
6.4.1 Primary map



X



Y

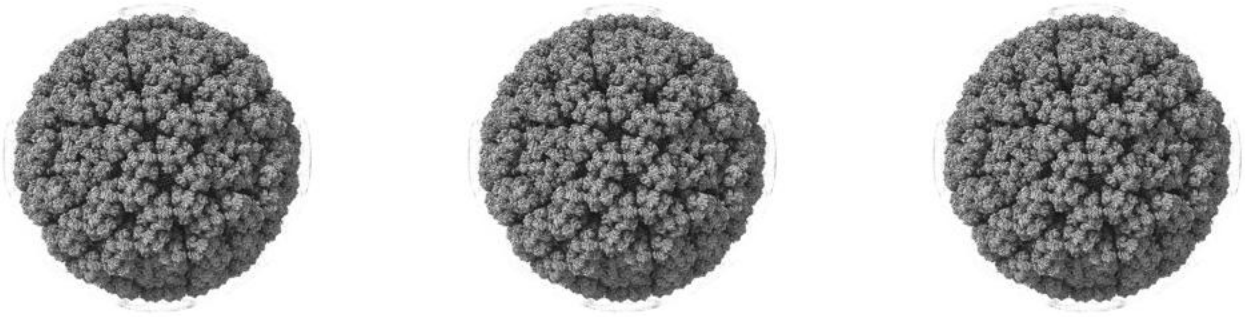


Z

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

6.5 Orthogonal surface views (i)

6.5.1 Primary map



X Y Z

The images above show the 3D surface view of the map at the recommended contour level 9.0. These images, in conjunction with the slice images, may facilitate assessment of whether an appropriate contour level has been provided.

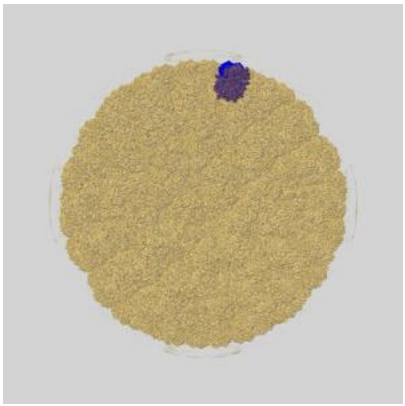
6.6 Mask visualisation (i)

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

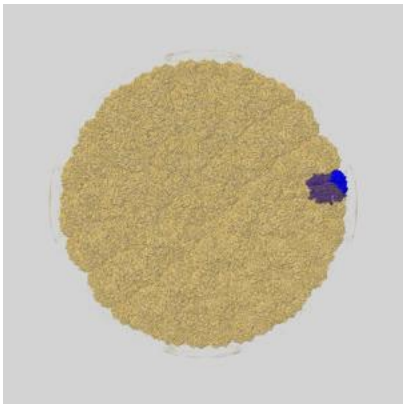
A mask typically either:

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

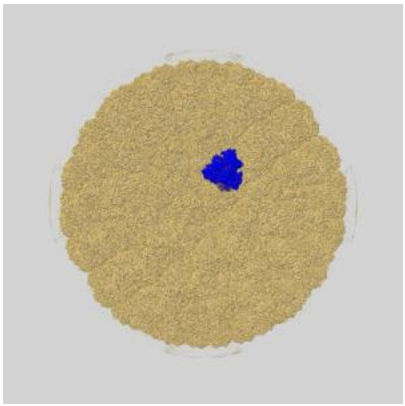
6.6.1 emd_6969_msk_1.map (i)



X



Y

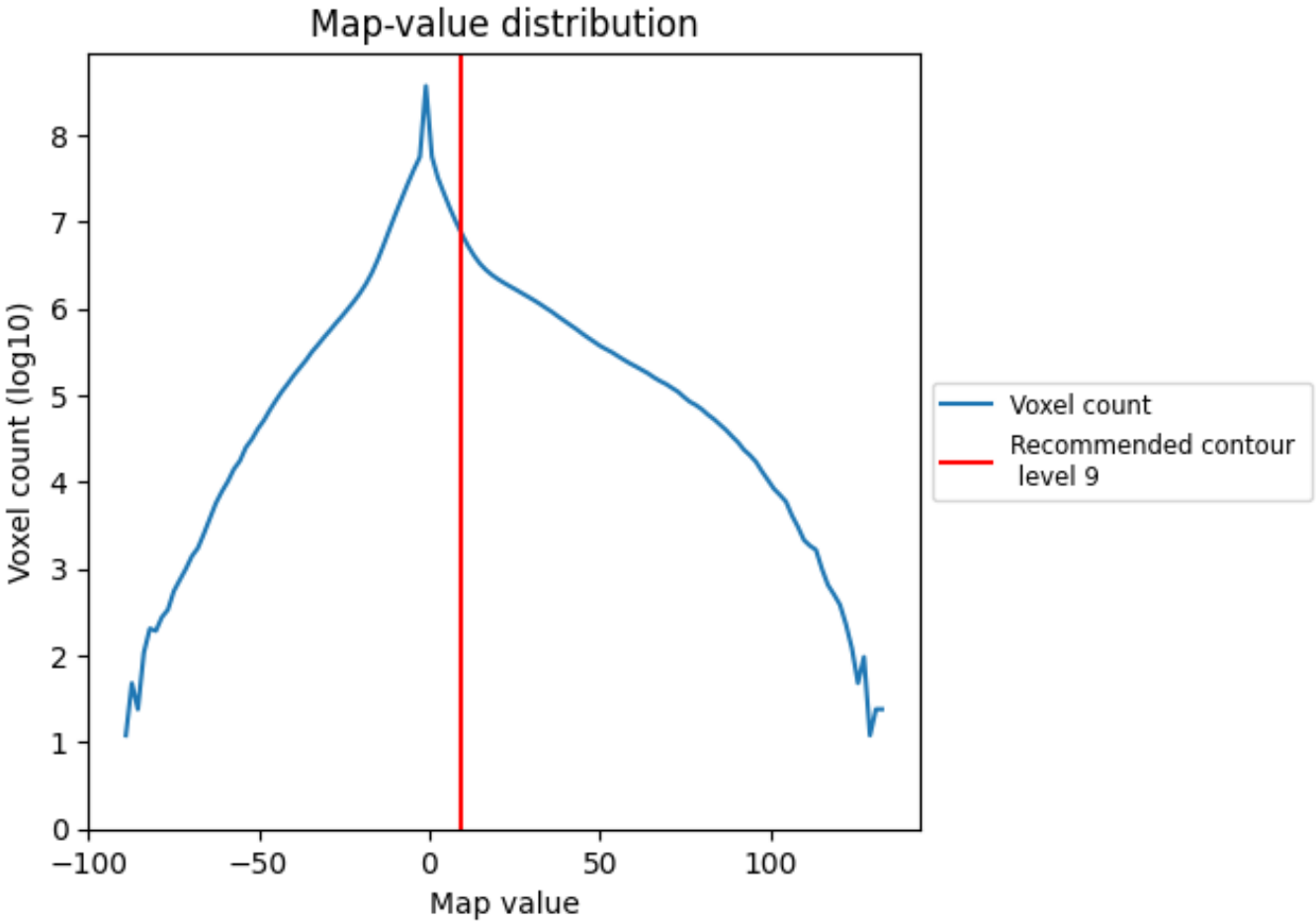


Z

7 Map analysis

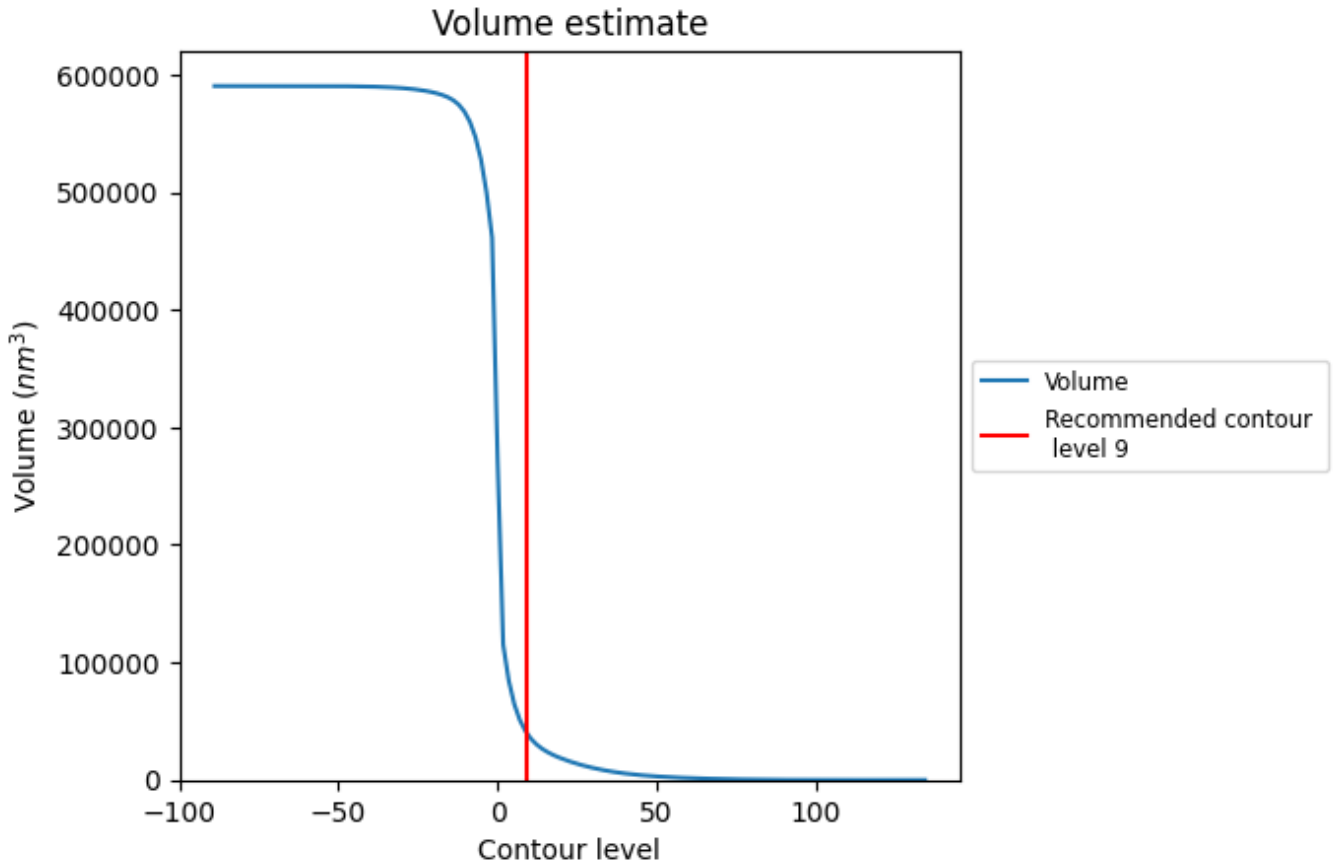
This section contains the results of statistical analysis of the map.

7.1 Map-value distribution



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

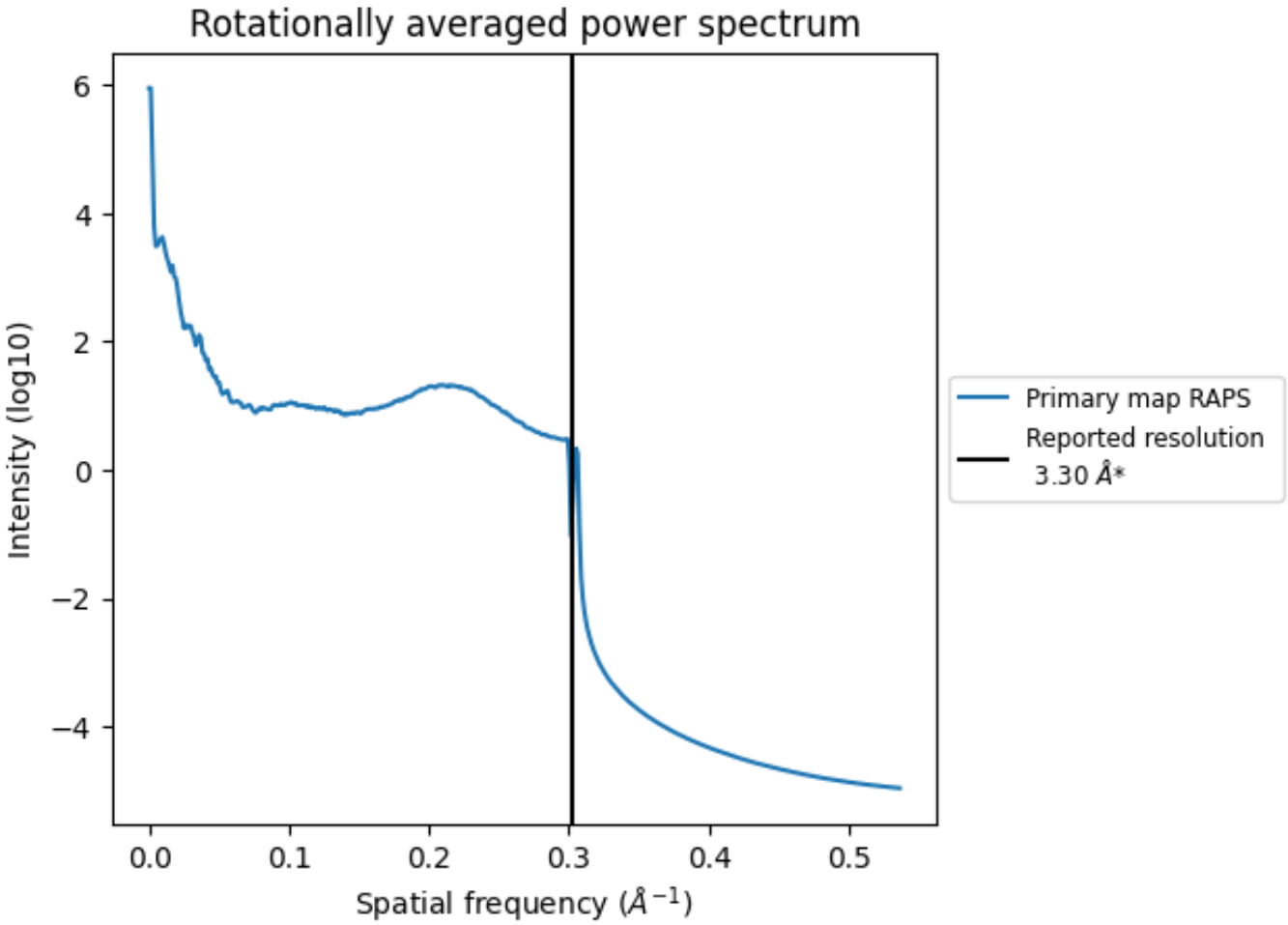
7.2 Volume estimate



The volume at the recommended contour level is 40818 nm^3; this corresponds to an approximate mass of 36872 kDa.

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

7.3 Rotationally averaged power spectrum (i)



*Reported resolution corresponds to spatial frequency of 0.303 Å^{-1}

8 Fourier-Shell correlation

This section was not generated. No FSC curve or half-maps provided.

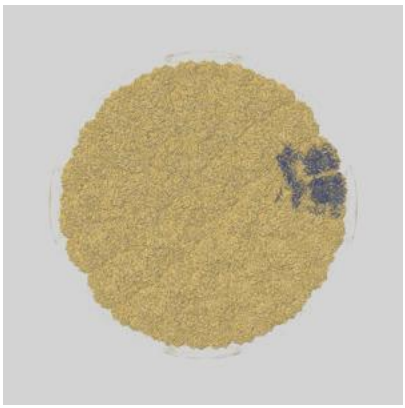
9 Map-model fit

This section contains information regarding the fit between EMDB map EMD-6969 and PDB model 5ZVT. Per-residue inclusion information can be found in section 3 on page 9.

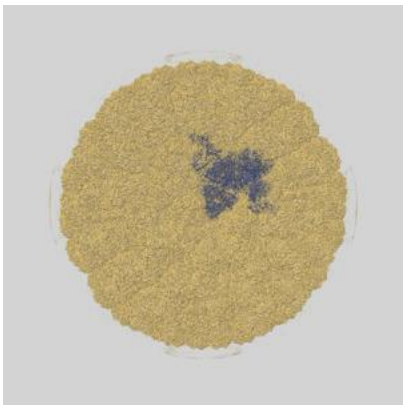
9.1 Map-model overlays

9.1.1 Map-model overlay

X

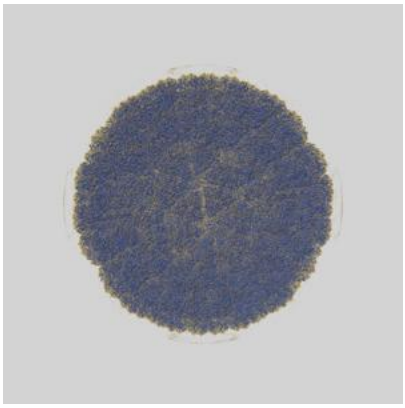


Y

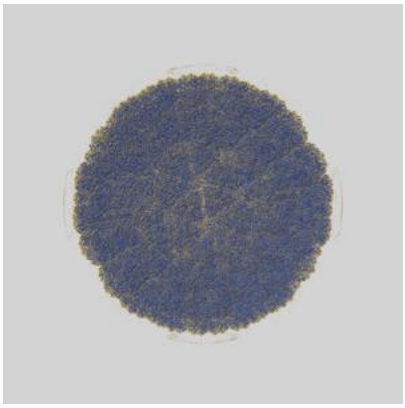


Z

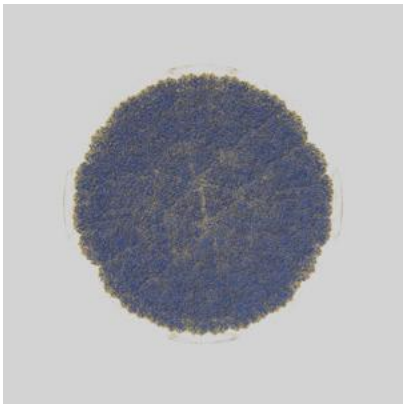
9.1.2 Map-model assembly overlay



X



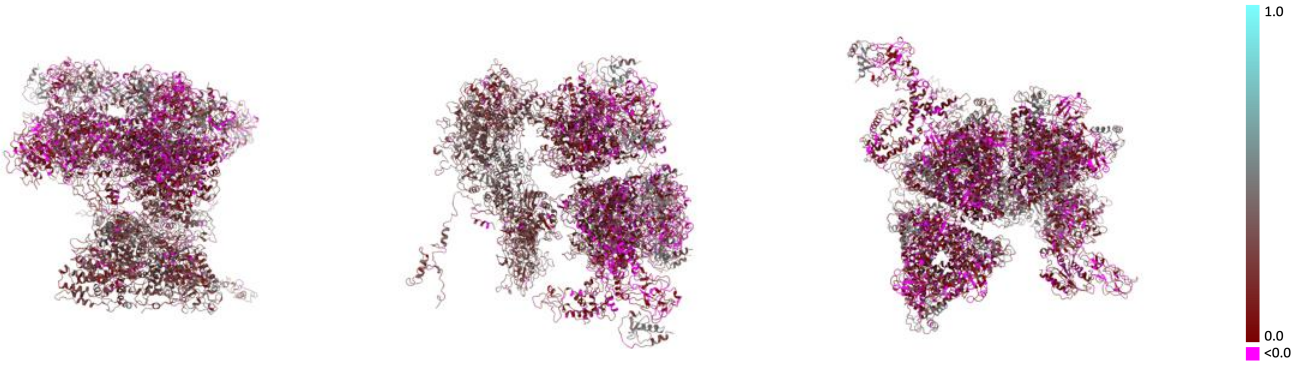
Y



Z

The images above show the 3D surface view of the map at the recommended contour level 9.0 at 50% transparency in yellow overlaid with a ribbon representation of the model coloured in blue. These images allow for the visual assessment of the quality of fit between the atomic model and the map.

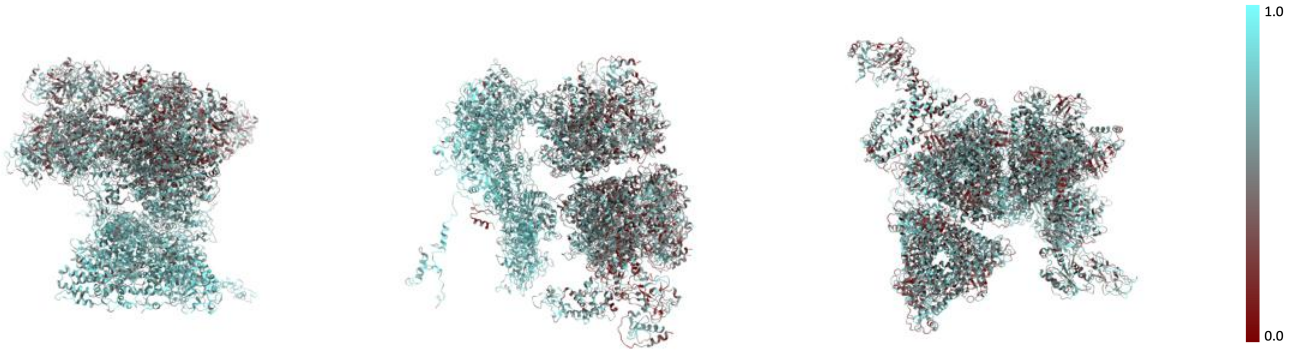
9.2 Q-score mapped to coordinate model



X Y Z

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

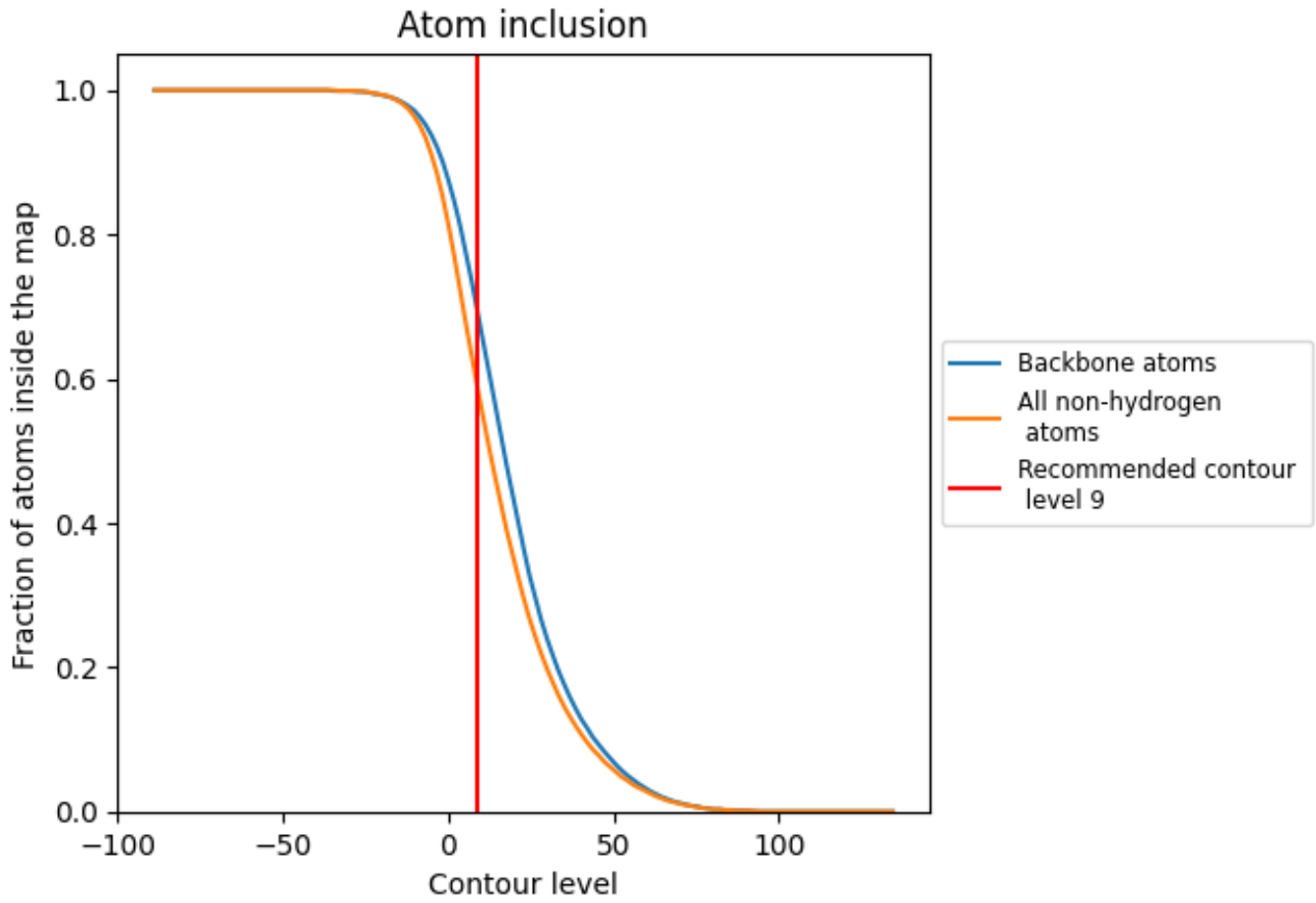
9.3 Atom inclusion mapped to coordinate model



X Y Z

The images above show the model with each residue coloured according to its atom inclusion. This shows to what extent they are inside the map at the recommended contour level (9).

9.4 Atom inclusion (i)



At the recommended contour level, 69% of all backbone atoms, 58% of all non-hydrogen atoms, are inside the map.

9.5 Map-model fit summary

The table lists the average atom inclusion at the recommended contour level (9) and Q-score for the entire model and for each chain.

Chain	Atom inclusion	Q-score
All	0.5820	0.1950
A	0.5020	0.1370
B	0.4820	0.1040
C	0.5180	0.1540
D	0.5200	0.1290
E	0.5180	0.1500
F	0.5270	0.1390
G	0.5300	0.1670
H	0.5220	0.1340
I	0.5050	0.1360
J	0.5110	0.1220
K	0.5280	0.1550
L	0.5070	0.1240
M	0.4800	0.1230
N	0.5110	0.1200
O	0.5340	0.1600
P	0.5200	0.1270
Q	0.5210	0.1220
R	0.5070	0.1210
S	0.5600	0.1760
T	0.5240	0.1310
U	0.6810	0.2370
V	0.7020	0.2490
W	0.5820	0.1600
X	0.7510	0.3170
Y	0.7450	0.3150
b	0.5450	0.3850
d	0.6160	0.3980
f	0.5960	0.4170
h	0.5810	0.4070
j	0.5810	0.4010
l	0.5640	0.3930
n	0.6380	0.4120
p	0.5380	0.4020
r	0.5670	0.4080
t	0.4410	0.3620