



## Full wwPDB EM Validation Report ⓘ

Jun 18, 2025 – 02:43 PM JST

PDB ID : 9JYY / pdb\_00009jyy  
EMDB ID : EMD-61909  
Title : core proteins of mature T7  
Authors : Liu, H.R.; Chen, W.Y.  
Deposited on : 2024-10-13  
Resolution : 3.00 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](mailto: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>.

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

EMDB validation analysis : **FAILED**  
MolProbity : 4-5-2 with Phenix2.0rc1  
Percentile statistics : 20231227.v01 (using entries in the PDB archive December 27th 2023)  
MapQ : **FAILED**  
Ideal geometry (proteins) : Engh & Huber (2001)  
Ideal geometry (DNA, RNA) : Parkinson et al. (1996)  
Validation Pipeline (wwPDB-VP) : 2.44

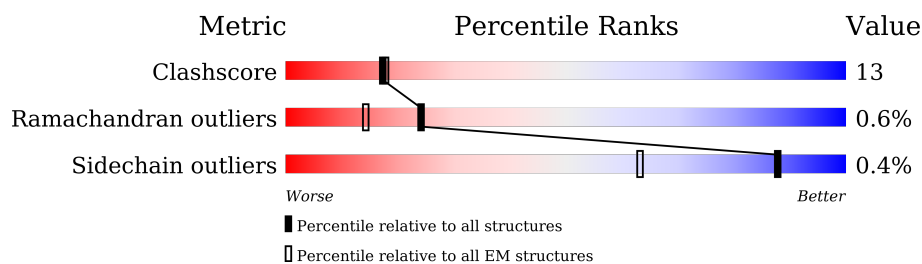
# 1 Overall quality at a glance

The following experimental techniques were used to determine the structure:

*ELECTRON MICROSCOPY*

The reported resolution of this entry is 3.00 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.



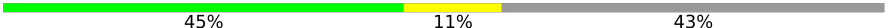
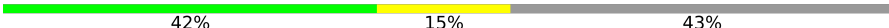
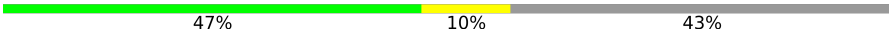
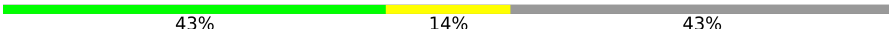
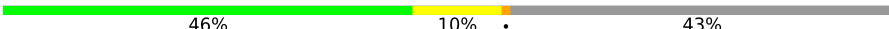
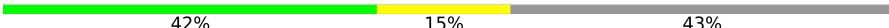
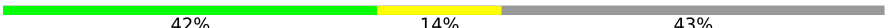












Metric	Whole archive (#Entries)	EM structures (#Entries)
Clashscore	210492	15764
Ramachandran outliers	207382	16835
Sidechain outliers	206894	16415

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  $\geq 3$ , 2, 1 and 0 types of geometric quality criteria respectively. A grey segment represents the fraction of residues that are not modelled. The numeric value for each fraction is indicated below the corresponding segment, with a dot representing fractions  $\leq 5\%$

Mol	Chain	Length	Quality of chain
1	A	88	15% 5% 81%
1	B	88	13% 7% 81%
1	C	88	18% . 81%
1	D	88	13% 6% 82%
1	E	88	13% 6% 82%
1	F	88	16% . 82%
1	a	88	17% . 81%
1	b	88	14% 5% 82%
2	S	196	41% 16% 43%

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Mol	Chain	Length	Quality of chain
2	T	196	
2	c	196	
2	d	196	
2	e	196	
2	f	196	
2	g	196	
2	h	196	
3	G	747	
3	H	747	
3	M	747	
3	N	747	
3	O	747	
3	P	747	
3	Q	747	
3	R	747	
4	I	1318	
4	J	1318	
4	K	1318	
4	L	1318	

## 2 Entry composition [i](#)

There are 4 unique types of molecules in this entry. The entry contains 87740 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 Protein 6.7.

Mol	Chain	Residues	Atoms				AltConf	Trace
1	a	17	Total	C	N	O	0	0
			116	66	17	33		
1	b	16	Total	C	N	O	0	0
			112	64	16	32		
1	A	17	Total	C	N	O	0	0
			116	66	17	33		
1	D	16	Total	C	N	O	0	0
			112	64	16	32		
1	B	17	Total	C	N	O	0	0
			116	66	17	33		
1	E	16	Total	C	N	O	0	0
			112	64	16	32		
1	C	17	Total	C	N	O	0	0
			116	66	17	33		
1	F	16	Total	C	N	O	0	0
			112	64	16	32		

- Molecule 2 is a protein called Internal virion protein gp14.

Mol	Chain	Residues	Atoms					AltConf	Trace
2	S	112	Total	C	N	O	S	0	0
			843	513	160	165	5		
2	T	111	Total	C	N	O	S	0	0
			835	507	159	164	5		
2	c	112	Total	C	N	O	S	0	0
			843	513	160	165	5		
2	d	111	Total	C	N	O	S	0	0
			835	507	159	164	5		
2	e	112	Total	C	N	O	S	0	0
			843	513	160	165	5		
2	f	111	Total	C	N	O	S	0	0
			835	507	159	164	5		
2	g	112	Total	C	N	O	S	0	0
			843	513	160	165	5		

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Mol	Chain	Residues	Atoms					AltConf	Trace
2	h	111	Total	C	N	O	S	0	0
			835	507	159	164	5		

- Molecule 3 is a protein called Internal virion protein gp15.

Mol	Chain	Residues	Atoms					AltConf	Trace
3	M	671	Total	C	N	O	S	0	0
			5335	3300	940	1065	30		
3	N	671	Total	C	N	O	S	0	0
			5335	3300	940	1065	30		
3	O	671	Total	C	N	O	S	0	0
			5335	3300	940	1065	30		
3	P	671	Total	C	N	O	S	0	0
			5335	3300	940	1065	30		
3	Q	671	Total	C	N	O	S	0	0
			5335	3300	940	1065	30		
3	R	671	Total	C	N	O	S	0	0
			5335	3300	940	1065	30		
3	G	671	Total	C	N	O	S	0	0
			5335	3300	940	1065	30		
3	H	671	Total	C	N	O	S	0	0
			5335	3300	940	1065	30		

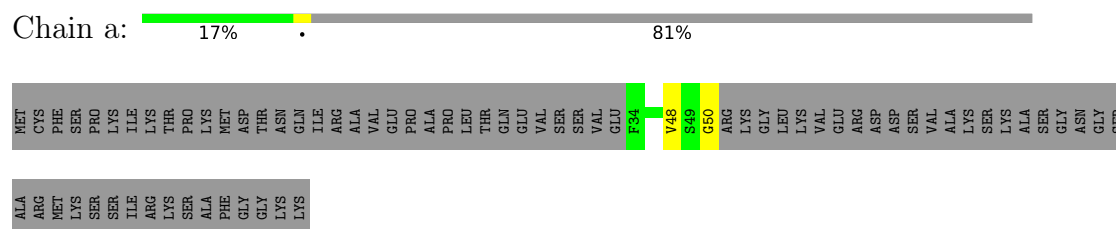
- Molecule 4 is a protein called Peptidoglycan transglycosylase gp16.

Mol	Chain	Residues	Atoms					AltConf	Trace
4	I	1223	Total	C	N	O	S	0	0
			9359	5853	1654	1805	47		
4	J	1223	Total	C	N	O	S	0	0
			9359	5853	1654	1805	47		
4	K	1223	Total	C	N	O	S	0	0
			9359	5853	1654	1805	47		
4	L	1223	Total	C	N	O	S	0	0
			9359	5853	1654	1805	47		

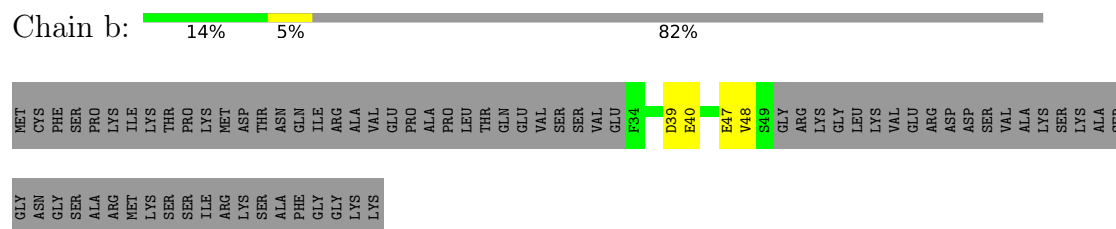
### 3 Residue-property plots [i](#)

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. 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. 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.

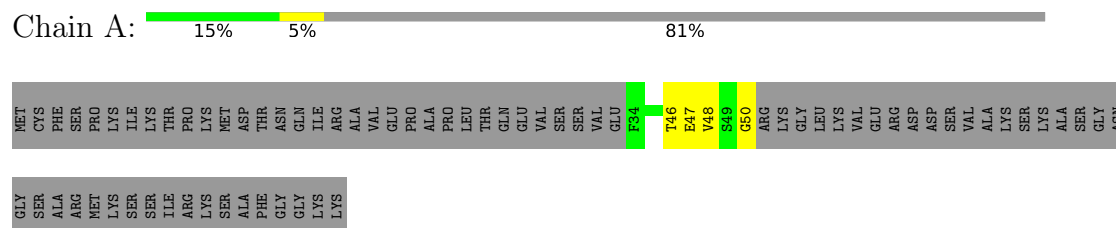
#### • Molecule 1: Protein 6.7



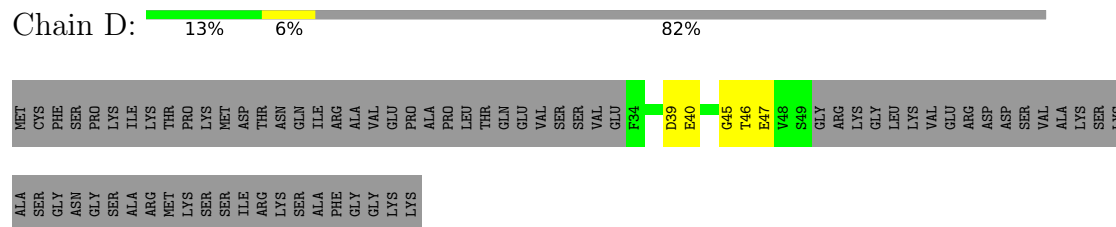
#### • Molecule 1: Protein 6.7



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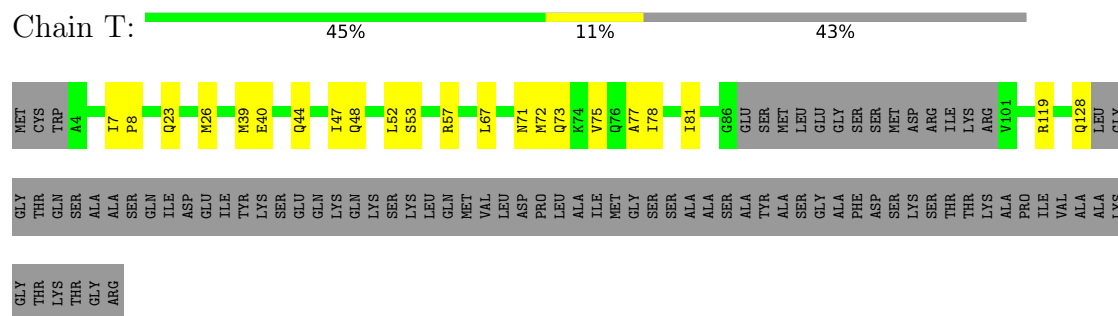
- Molecule 1: Protein 6.7

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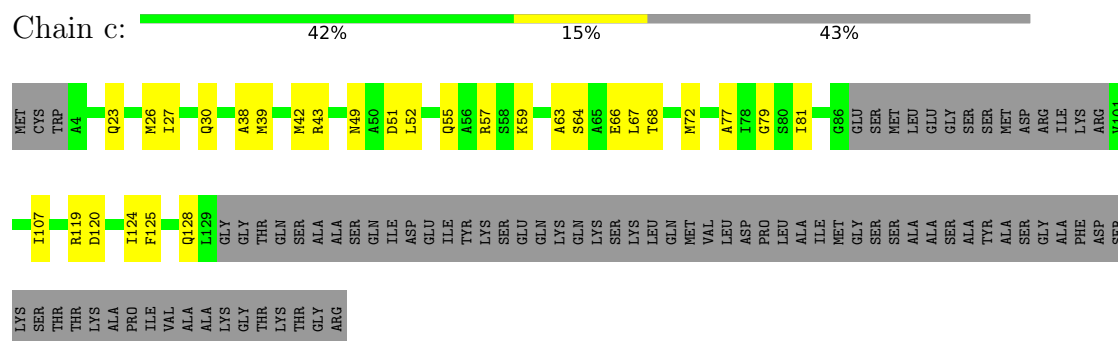
- Molecule 1: Protein 6.7

- Molecule 2: Internal virion protein gp14

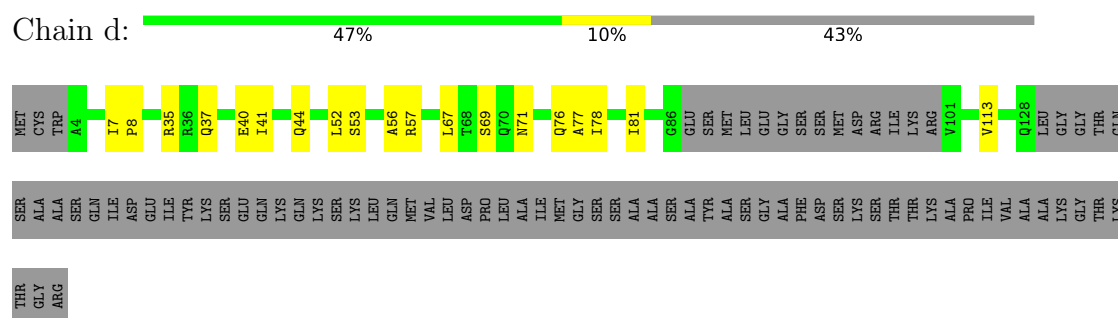
- Molecule 2: Internal virion protein gp14



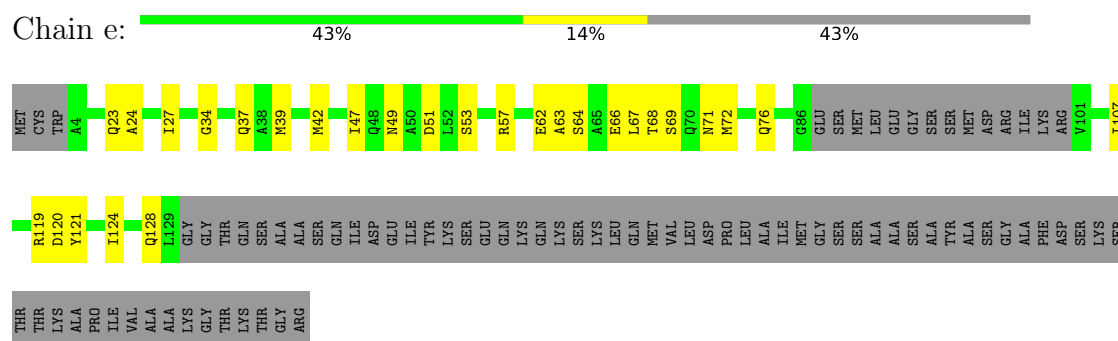
- Molecule 2: Internal virion protein gp14



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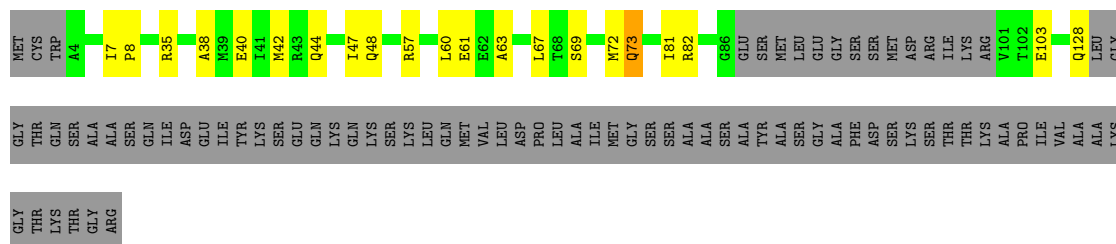


- Molecule 2: Internal virion protein gp14



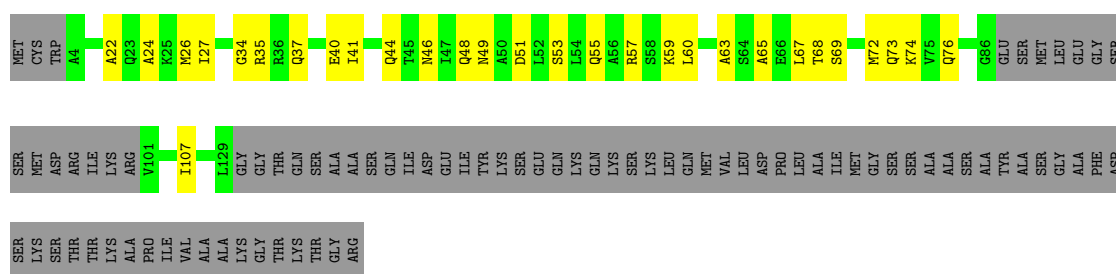
- Molecule 2: Internal virion protein gp14

Chain f: 



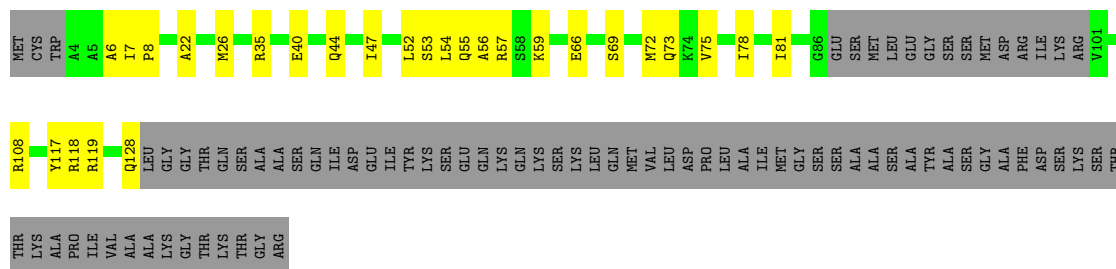
• Molecule 2: Internal virion protein gp14

Chain g: 



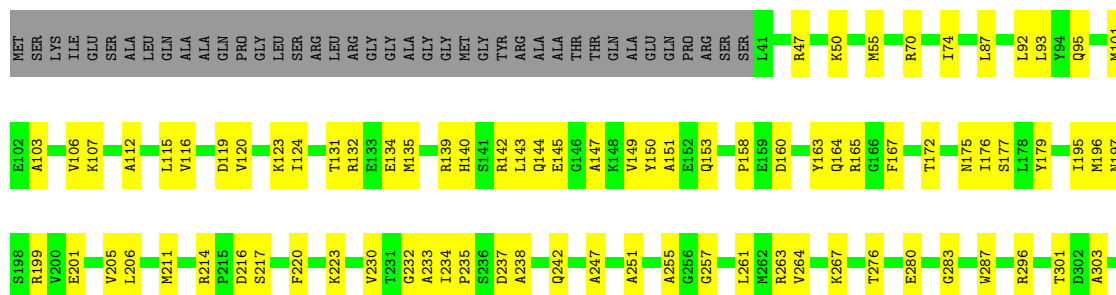
• Molecule 2: Internal virion protein gp14

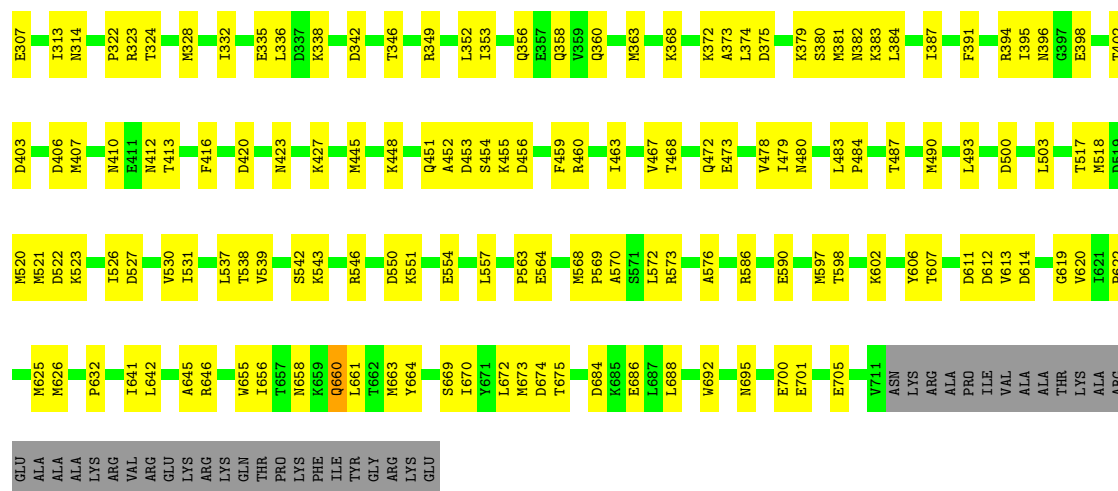
Chain h: 



• Molecule 3: Internal virion protein gp15

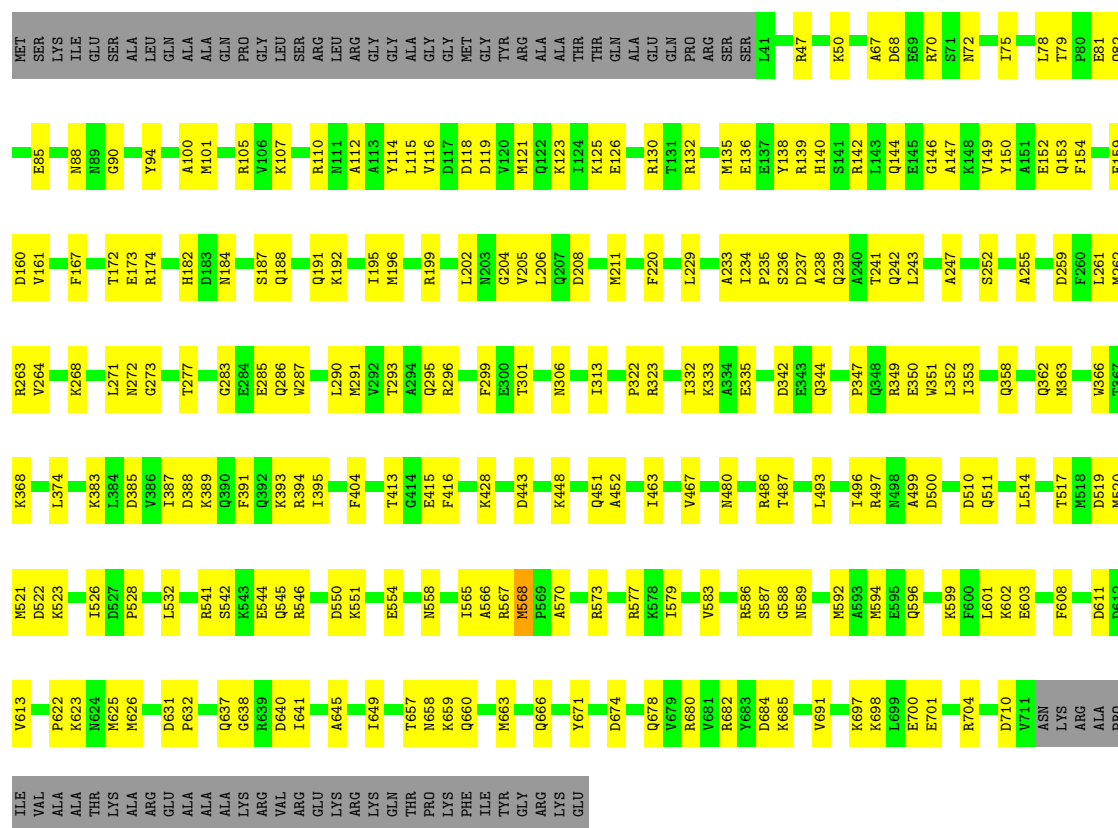
Chain M: 





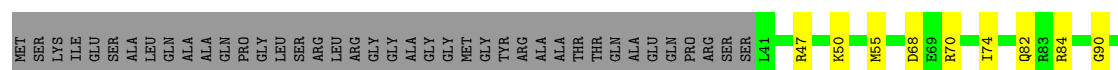
• Molecule 3: Internal virion protein gp15

Chain N: 59% 31% 10%



• Molecule 3: Internal virion protein gp15

Chain O: 60% 29% 10%







- Molecule 3: Internal virion protein gp15

[illegible]

- Molecule 3: Internal virion protein gp15

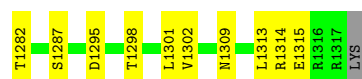
L450	L451	A452	D453	S454	K455	R460	I463	V467	E473	I479	T487	M490	L493	I496	R497	A498	A499	L503	L507	D510	Q511	F515	M518	D522	K523	I526	R536	R541	S542	E544	Q545	R546	D549	D560	K561	A562	F563												
E285	Q286	W287	M291	V292	T293	R296	E300	N318	E327	K338	Q344	M345	T346	Q356	V359	Q362	K368	L374	K383	F391	R394	I395	W399	F404	K405	D406	M407	T413	F416	K417	H418	S419	I437	D549	D560	K561	A562	F563											
D183	Q188	M196	K192	M196	M197	E201	L202	N203	G204	L206	Q207	M211	R214	F220	A233	I234	P235	D237	A238	Q239	A240	R254	L261	R262	R263	V264	K268	V269	F272	G273	T277	Y278	E173	N174	G283	F294	L178												
L93	Y94	A100	M101	E102	R105	G110	N111	A112	A113	Y114	L115	V116	D117	D118	D119	V120	M121	Q122	K125	E126	R130	E134	M135	V149	Y150	E159	D160	V161	D162	R165	G166	F167	D170	I171	I172	E173	N174	N175											
MET	SER	LVS	ILE	GLU	SER	ALA	LEU	GLN	ALA	ALA	GLN	PRO	GLY	LEU	ARG	SER	LEU	ARG	LEU	ARG	GLY	MET	GLY	TTR	GLY	ALA	ALA	THR	THR	GLN	GLN	GLU	PRO	SER	SER	SER	SER	L41	R47	K50	L66	A67	R70	N72	Y75	L87	N88	N89	L90

LYS	I649	A566
GLN	T657	A567
THR		P563
PRO		E564
LYS	Q660	I565
PHE		A566
ILE	M663	R567
TYR	Y664	M568
GLY		P569
ARG	S669	A570
LYS		S571
GLU	D674	L572
		R573
	Q678	
	V679	R577
	R680	
	V681	V583
	R682	K584
	Y683	Y585
	D684	R586
		S587
	V691	S588
		N589
	K697	
	K698	M592
	L699	A593
	E700	M594
	E701	E595
		Q596
	R704	M597
	E705	T598
	K706	K599
		F600
	D710	L601
	V711	K602
	ASN	
	LYS	F608
	ARG	
	ALA	D612
	PRO	
	ILE	G615
	VAL	D616
	ALA	T617
	ALA	
	THR	P622
	LYS	K623
	ALA	N624
	ARG	M625
	GLU	M626
	ALA	Q627
	ALA	V628
	ALA	
	LYS	S634
	ARG	
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	ARG	
	GLU	T641
	LYS	
	ARG	A645

● Molecule 4: Peptidoglycan transglycosylase gp16

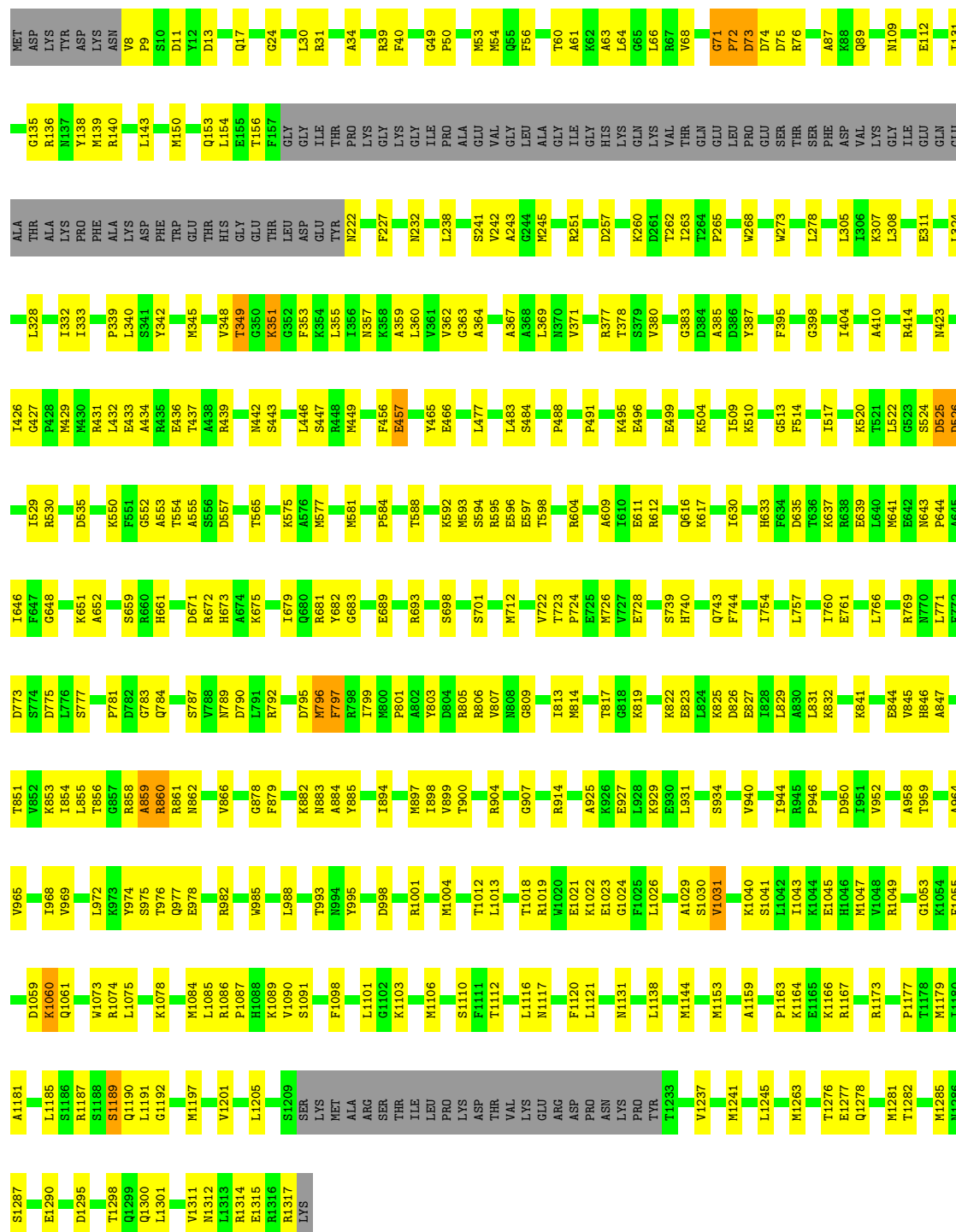
Chain I:  64% 28% 7%

MET	ASP	LYS	TRP	LYS	V8	S10	D11	Y12	D13	F16	Q17	K18	G24	L29	R31	F40	G49	P50	M63	M64	Q65	F66	T60	A61	K62	A63	L64	G65	L66	R67	V68	T69	P72	D73	R76	E80	K88	E101	K125	I131											
R136	M139	R140	L143	K147	L154	F157	GLY	GLY	GLY	ILE	THR	PRO	LYS	LYS	ILE	PRO	GLU	VAL	LEU	ALA	GLY	ILE	HIS	LYS	LYS	VAL	THR	GLM	GLU	LEU	PRO	GLU	SER	THR	PHE	ASP	VAL	LYS	ILE	GLU	GLN	GLU	ALA	THR	ALA	LYS	PRO				
PHE	ALA	LYS	ASP	PHE	TRP	GLU	THR	HIS	GLY	GLU	THR	LEU	ASP	GLU	TYR	N222	F227	N232	L238	S241	V242	G244	M245	R251	L252	D253	F256	T262	I263	T264	P265	R266	R667	W268	W273	L278	V293	L305	N310	E314	R318	L324									
L328	R439	M440	A441	I333	P339	L340	S341	Y342	M345	I356	N357	K358	L360	V361	R362	G363	A364	L369	N370	V371	L376	R377	G244	M245	R251	L252	D253	F395	G398	I404	V408	K504	I509	K510	F514	I517	K520	I426	G427	M429	D525	D526	I529	E433	A434						
R439	M440	A441	I333	P339	L340	S341	Y342	M345	I356	N357	K358	L360	V361	R362	G363	A364	L369	N370	V371	L376	R377	G244	M245	R251	L252	D253	F395	G398	I404	V408	K504	I509	K510	F514	I517	K520	I426	G427	M429	D525	D526	I529	E433	A434							
R538	K550	F551	G552	A553	T554	A555	S556	D557	E560	R561	L562	H563	T565	D572	M577	M581	P584	T588	M593	S594	R595	E596	E597	T598	R604	B611	R612	P613	E614	L615	Q616	M627	T630	H633	K637	M641	E642	N643	P644	A645	I646	L647	G648								
V653	F656	S659	R660	H661	V669	H673	A674	K675	I679	G680	R681	V682	G683	A684	L687	Q688	E689	R693	S701	M712	E715	V722	T723	P724	E725	M726	V727	H740	S748	E752	N753	I754	L757	T760	E761	L766	R769	H770	L771												
D775	L776	T777	T779	M780	P781	D782	S787	V788	M789	D790	L791	R792	F793	F794	D795	M796	R797	R798	M800	P801	A802	Y803	D804	R805	R806	V807	I811	M814	G815	S816	T817	G818	K819	T820	K822	E823	L824	K825	D826	E827	I828	L829	A830	L831	E844	A847	L848	M849	D850	R851	V852
K853	I854	L855	T856	A859	R860	R861	M862	Q863	V866	G878	F879	F880	A881	K882	T883	M884	Y885	A888	M889	Q889	T892	E893	L894	M897	L898	N900	T903	L906	G907	R914	K919	S920	K921	S934	L935	F936	G937	K938	F939	V940	D941	Q942	L943	I944	R945	P946	R947	R948	A949		
D950	E957	K1054	T959	D960	T961	A964	I968	V969	L972	K973	Y974	S975	T976	Q977	E978	L979	A980	A981	R982	Y985	L988	N989	N990	T993	L996	L997	D998	R1001	M1004	I1009	L1013	R1019	W1020	E1021	K1022	N1023	G1024	S1030	P1033	I1041	G1149	F1150	E1045								
R1049	G1053	F1055	D1059	K1060	Q1061	D1071	R1074	D1077	K1078	V1079	A1080	Q1081	E1082	L1083	M1084	A1085	R1086	P1087	V1090	F1098	G1099	A1100	L1101	G1102	L1103	K1109	S1110	F1111	T1112	L1116	K1119	F1120	L1121	K1129	M1241	G1242	S1243	M1244	L1245	Q1248	T1261	E1277	Q1278	M1153							
K1168	L1171	E1172	R1173	A1174	L1175	A1181	G1192	M1197	G1202	L1205	S1209	SER	LYS	MET	ALA	ARG	SER	THR	ILE	LEU	PRO	LYS	ASP	THR	VAL	LYS	GLU	ARG	ASP	PRO	ASN	LYS	PRO	TYR	T1233	V1237	M1241	G1242	S1243	M1244	L1245	Q1248	T1261	E1277	Q1278	M1153					



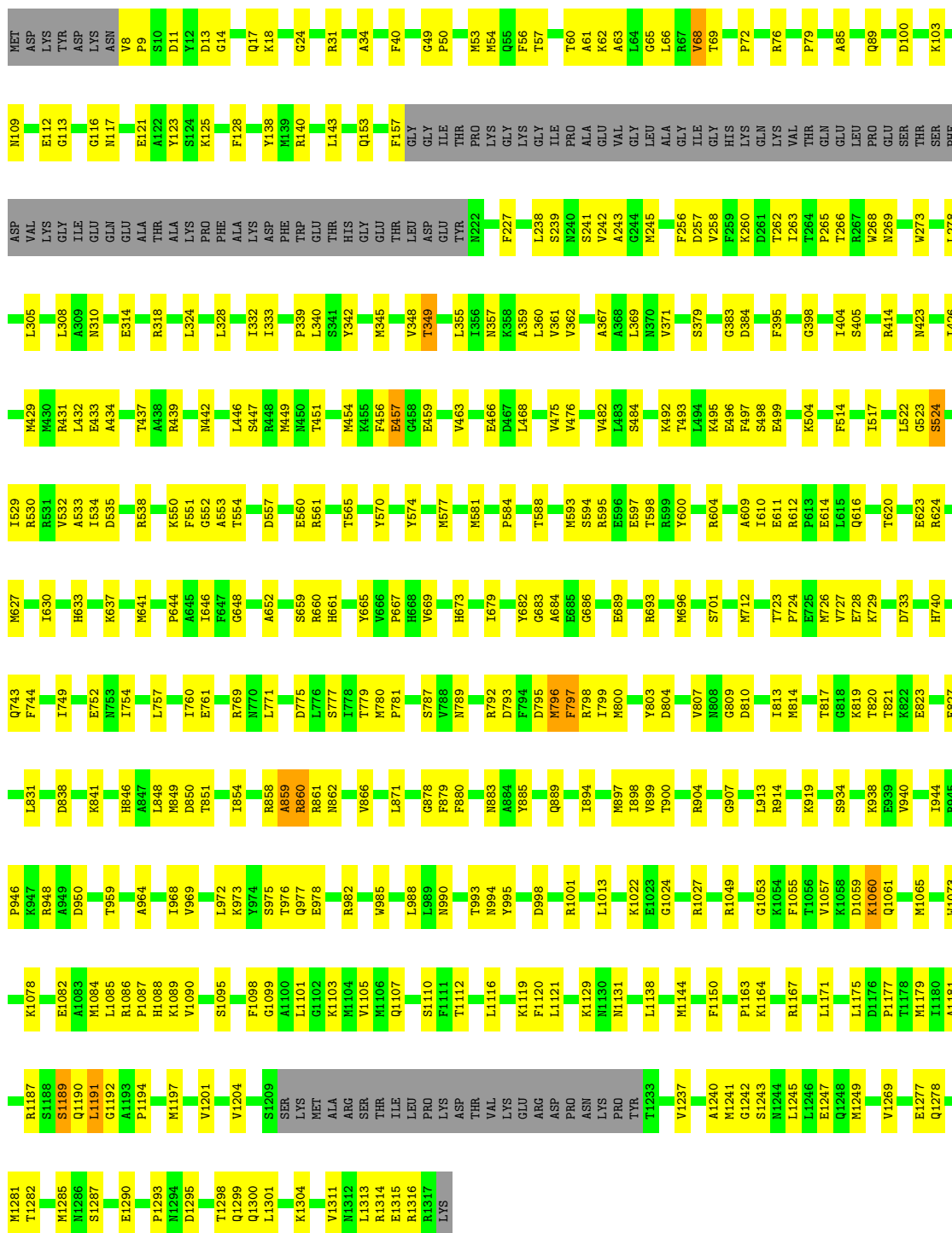
● Molecule 4: Peptidoglycan transglycosylase gp16

Chain J: 64% 28% 7%



● Molecule 4: Peptidoglycan transglycosylase gp16

Chain K:  63% 29% • 7%



- Molecule 4: Peptidoglycan transglycosylase gp16

Chain L:  64% 28% 7%





## 4 Experimental information

Property	Value	Source
EM reconstruction method	SINGLE PARTICLE	Depositor
Imposed symmetry	POINT, Not provided	
Number of particles used	45731	Depositor
Resolution determination method	FSC 0.143 CUT-OFF	Depositor
CTF correction method	PHASE FLIPPING AND AMPLITUDE CORRECTION	Depositor
Microscope	TFS KRIOS	Depositor
Voltage (kV)	300	Depositor
Electron dose ( $e^-/\text{\AA}^2$ )	35	Depositor
Minimum defocus (nm)	500	Depositor
Maximum defocus (nm)	4000	Depositor
Magnification	Not provided	
Image detector	GATAN K3 (6k x 4k)	Depositor

## 5 Model quality

### 5.1 Standard geometry

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

Mol	Chain	Bond lengths		Bond angles	
		RMSZ	# Z  >5	RMSZ	# Z  >5
1	A	0.14	0/116	0.33	0/155
1	B	0.18	0/116	0.38	0/155
1	C	0.14	0/116	0.34	0/155
1	D	0.21	0/112	0.42	0/150
1	E	0.21	0/112	0.43	0/150
1	F	0.20	0/112	0.46	0/150
1	a	0.18	0/116	0.31	0/155
1	b	0.20	0/112	0.41	0/150
2	S	0.21	0/846	0.42	0/1135
2	T	0.21	0/838	0.41	0/1124
2	c	0.19	0/846	0.42	0/1135
2	d	0.19	0/838	0.37	0/1124
2	e	0.18	0/846	0.39	0/1135
2	f	0.25	0/838	0.47	1/1124 (0.1%)
2	g	0.19	0/846	0.40	0/1135
2	h	0.18	0/838	0.38	0/1124
3	G	0.28	0/5415	0.52	5/7291 (0.1%)
3	H	0.27	0/5415	0.48	2/7291 (0.0%)
3	M	0.29	0/5415	0.53	6/7291 (0.1%)
3	N	0.28	0/5415	0.47	1/7291 (0.0%)
3	O	0.28	0/5415	0.50	5/7291 (0.1%)
3	P	0.27	0/5415	0.45	0/7291
3	Q	0.29	0/5415	0.52	5/7291 (0.1%)
3	R	0.27	0/5415	0.45	0/7291
4	I	0.29	0/9514	0.52	5/12824 (0.0%)
4	J	0.29	0/9514	0.54	11/12824 (0.1%)
4	K	0.29	0/9514	0.53	6/12824 (0.0%)
4	L	0.30	0/9514	0.55	11/12824 (0.1%)
All	All	0.28	0/89024	0.50	58/119880 (0.0%)

There are no bond length outliers.

All (58) bond angle outliers are listed below:

Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
3	Q	538	THR	N-CA-C	12.31	124.70	111.28
3	M	538	THR	N-CA-C	11.54	123.85	111.28
3	O	540	LYS	N-CA-C	10.10	122.29	111.28
3	G	540	LYS	N-CA-C	9.92	122.10	111.28
3	M	537	LEU	N-CA-C	9.13	121.00	111.14
4	L	72	PRO	N-CA-C	-9.10	102.83	114.03
4	L	524	SER	N-CA-C	8.81	120.89	111.28
3	M	542	SER	N-CA-C	8.40	123.12	108.75
4	I	525	ASP	N-CA-C	8.32	122.86	111.54
3	G	236	SER	N-CA-C	8.24	121.86	110.24
3	Q	542	SER	N-CA-C	8.16	122.70	108.75
3	G	539	VAL	N-CA-C	-7.80	103.66	111.77
4	I	524	SER	N-CA-C	7.55	120.95	108.49
4	K	348	VAL	N-CA-C	7.54	119.11	108.93
4	K	65	GLY	N-CA-C	7.50	121.73	112.73
3	Q	237	ASP	N-CA-C	-7.49	103.11	111.28
4	L	238	LEU	N-CA-C	7.44	119.47	111.36
4	L	797	PHE	N-CA-C	-7.36	104.27	113.55
3	O	538	THR	N-CA-C	7.34	121.06	109.39
4	J	72	PRO	N-CA-C	-7.33	105.02	114.03
4	I	73	ASP	N-CA-C	6.93	120.24	108.67
4	K	349	THR	N-CA-C	-6.86	98.21	109.40
3	N	451	GLN	N-CA-C	-6.83	101.52	110.53
3	G	538	THR	N-CA-C	6.74	119.62	108.49
4	I	68	VAL	N-CA-C	-6.68	106.62	113.10
3	O	539	VAL	N-CA-C	-6.67	104.84	111.77
4	J	525	ASP	N-CA-C	6.48	118.85	108.41
2	f	48	GLN	N-CA-C	6.42	118.84	110.43
4	J	353	PHE	N-CA-C	6.38	119.62	108.52
3	M	342	ASP	N-CA-C	6.37	118.30	111.36
4	L	1189	SER	N-CA-C	6.31	118.16	111.28
4	L	71	GLY	CA-C-N	-6.18	113.44	119.87
4	L	71	GLY	C-N-CA	-6.18	113.44	119.87
3	O	235	PRO	N-CA-C	6.18	121.03	111.21
4	K	1189	SER	N-CA-C	6.12	117.95	111.28
4	J	797	PHE	N-CA-C	-5.95	105.02	112.93
4	K	797	PHE	N-CA-C	-5.92	105.06	112.93
4	I	797	PHE	N-CA-C	-5.92	105.06	112.93
4	J	349	THR	N-CA-C	-5.88	100.32	109.72
4	J	71	GLY	CA-C-N	-5.85	113.78	119.87
4	J	71	GLY	C-N-CA	-5.85	113.78	119.87
4	L	553	ALA	CA-C-N	5.67	132.37	121.54
4	L	553	ALA	C-N-CA	5.67	132.37	121.54

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
3	Q	342	ASP	N-CA-C	5.60	117.38	111.28
3	H	235	PRO	N-CA-C	5.57	121.44	113.47
4	L	1164	LYS	N-CA-C	5.52	122.55	110.80
3	Q	230	VAL	N-CA-C	5.48	116.21	110.62
3	M	237	ASP	N-CA-C	-5.44	105.45	111.71
4	J	348	VAL	N-CA-C	5.39	115.95	108.84
3	O	236	SER	N-CA-C	5.34	123.97	113.29
4	K	1191	LEU	N-CA-C	5.28	120.08	111.37
3	M	230	VAL	N-CA-C	5.25	115.98	110.62
3	H	237	ASP	N-CA-C	-5.23	105.05	111.75
4	J	1189	SER	N-CA-C	5.21	116.95	111.28
3	G	238	ALA	N-CA-C	5.20	116.95	111.28
4	J	74	ASP	N-CA-C	-5.17	106.93	113.18
4	J	73	ASP	CB-CA-C	-5.14	110.67	116.63
4	L	73	ASP	CB-CA-C	-5.14	110.67	116.63

There are no chirality outliers.

There are no planarity outliers.

## 5.2 Too-close contacts ⓘ

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

Mol	Chain	Non-H	H(model)	H(added)	Clashes	Symm-Clashes
1	A	116	0	91	4	0
1	B	116	0	91	4	0
1	C	116	0	91	1	0
1	D	112	0	88	5	0
1	E	112	0	88	6	0
1	F	112	0	88	3	0
1	a	116	0	91	1	0
1	b	112	0	88	5	0
2	S	843	0	852	29	0
2	T	835	0	841	22	0
2	c	843	0	852	22	0
2	d	835	0	841	18	0
2	e	843	0	852	21	0
2	f	835	0	841	16	0

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Mol	Chain	Non-H	H(model)	H(added)	Clashes	Symm-Clashes
2	g	843	0	852	22	0
2	h	835	0	841	23	0
3	G	5335	0	5232	164	0
3	H	5335	0	5232	164	0
3	M	5335	0	5232	168	0
3	N	5335	0	5232	188	0
3	O	5335	0	5232	170	0
3	P	5335	0	5232	186	0
3	Q	5335	0	5232	143	0
3	R	5335	0	5232	183	0
4	I	9359	0	9352	280	0
4	J	9359	0	9352	275	0
4	K	9359	0	9352	274	0
4	L	9359	0	9352	283	0
All	All	87740	0	86752	2346	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 13.

All (2346) 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)
4:J:64:LEU:HD11	4:J:89:GLN:OE1	1.44	1.17
3:P:79:THR:H	3:P:82:GLN:HE21	1.14	0.95
4:K:669:VAL:HG11	4:K:799:ILE:HD12	1.49	0.95
4:I:669:VAL:HG11	4:I:799:ILE:HD12	1.51	0.91
2:f:42:MET:HG3	3:R:196:MET:HE3	1.54	0.89
3:N:678:GLN:NE2	4:J:449:MET:SD	2.47	0.88
3:O:590:GLU:H	3:P:541:ARG:HG3	1.36	0.88
3:R:678:GLN:HE21	4:L:449:MET:HE1	1.37	0.88
4:L:1314:ARG:HG3	4:L:1315:GLU:H	1.40	0.87
3:Q:590:GLU:H	3:R:541:ARG:HG3	1.39	0.87
3:P:678:GLN:NE2	4:K:449:MET:SD	2.48	0.87
4:K:993:THR:HG23	4:K:1085:LEU:HD21	1.56	0.87
3:M:590:GLU:H	3:N:541:ARG:HG3	1.40	0.86
4:K:398:GLY:HA3	4:K:429:MET:HE3	1.57	0.86
3:G:383:LYS:HB3	3:G:413:THR:HG23	1.61	0.83
4:K:904:ARG:HH22	4:K:1314:ARG:HD3	1.42	0.83
4:J:1314:ARG:HG3	4:J:1315:GLU:H	1.44	0.83
3:M:695:ASN:ND2	4:J:245:MET:SD	2.53	0.82
4:K:959:THR:HG21	4:K:1078:LYS:HE2	1.62	0.81

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:H:383:LYS:HB3	3:H:413:THR:HG23	1.63	0.81
3:R:404:PHE:HA	3:R:407:MET:HG3	1.63	0.80
4:L:1090:VAL:HG11	4:L:1110:SER:HB2	1.63	0.80
4:I:861:ARG:HH21	4:I:1205:LEU:HD11	1.46	0.80
3:Q:383:LYS:HB3	3:Q:413:THR:HG23	1.64	0.79
4:L:263:ILE:HD11	4:L:1116:LEU:HD22	1.63	0.79
3:P:404:PHE:HA	3:P:407:MET:HG3	1.62	0.79
3:R:383:LYS:HB3	3:R:413:THR:HG23	1.63	0.79
3:P:383:LYS:HB3	3:P:413:THR:HG23	1.65	0.78
3:G:590:GLU:H	3:H:541:ARG:HG3	1.47	0.78
3:H:563:PRO:HB3	3:H:617:THR:HG21	1.66	0.78
3:M:551:LYS:HD2	4:J:308:LEU:HD11	1.65	0.78
4:J:263:ILE:HD11	4:J:1116:LEU:HD22	1.66	0.78
4:J:976:THR:O	4:J:977:GLN:HG2	1.84	0.78
3:M:383:LYS:HB3	3:M:413:THR:HG23	1.65	0.78
3:R:125:LYS:HZ3	3:R:178:LEU:HD12	1.46	0.78
3:G:695:ASN:ND2	4:I:245:MET:SD	2.57	0.78
4:I:1090:VAL:HG11	4:I:1110:SER:HB2	1.64	0.77
3:O:383:LYS:HB3	3:O:413:THR:HG23	1.66	0.77
3:R:569:PRO:HG2	3:R:572:LEU:HD13	1.66	0.77
3:H:125:LYS:NZ	3:H:135:MET:SD	2.57	0.77
3:G:647:LYS:HD2	3:H:405:LYS:HG3	1.67	0.77
3:H:678:GLN:NE2	4:I:449:MET:SD	2.57	0.77
4:J:681:ARG:HH21	4:J:682:TYR:HE2	1.31	0.76
4:I:959:THR:HG21	4:I:1078:LYS:HE2	1.67	0.76
3:N:383:LYS:HB3	3:N:413:THR:HG23	1.66	0.76
4:J:968:ILE:HG13	4:J:969:VAL:HG23	1.69	0.75
4:K:241:SER:O	4:K:243:ALA:N	2.20	0.75
3:Q:483:LEU:HD12	3:Q:484:PRO:HD2	1.69	0.75
4:J:377:ARG:NH2	4:K:978:GLU:OE2	2.20	0.75
3:M:568:MET:HE3	3:M:573:ARG:HA	1.67	0.75
3:M:520:MET:HB3	3:M:526:ILE:HD11	1.69	0.75
3:G:360:GLN:NE2	3:H:285:GLU:OE2	2.20	0.74
3:R:563:PRO:HB3	3:R:617:THR:HG21	1.68	0.74
3:M:557:LEU:HD11	3:M:597:MET:HE1	1.69	0.74
4:L:959:THR:HG21	4:L:1078:LYS:HE2	1.70	0.73
4:K:595:ARG:HG2	4:K:1192:GLY:HA3	1.69	0.73
3:P:710:ASP:OD2	3:R:295:GLN:NE2	2.20	0.73
3:R:201:GLU:OE1	3:G:110:ARG:NH2	2.21	0.73
4:L:355:LEU:HD21	4:L:433:GLU:HG3	1.70	0.73
4:I:369:LEU:HD21	4:J:24:GLY:H	1.53	0.73

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
4:L:241:SER:O	4:L:243:ALA:N	2.21	0.73
4:L:456:PHE:O	4:L:457:GLU:HG3	1.89	0.73
4:L:38:SER:HB2	4:L:41:VAL:HG12	1.70	0.73
3:H:118:ASP:OD1	3:H:139:ARG:NH1	2.21	0.73
3:O:360:GLN:NE2	3:P:285:GLU:OE1	2.22	0.73
4:I:416:LYS:HZ2	4:I:418:GLU:H	1.37	0.72
3:O:520:MET:HB3	3:O:526:ILE:HD11	1.71	0.72
3:P:125:LYS:HZ1	3:P:178:LEU:HB3	1.53	0.72
4:L:1045:GLU:OE2	4:L:1068:ARG:NH2	2.21	0.72
3:R:214:ARG:O	3:R:263:ARG:NH2	2.23	0.72
4:I:1314:ARG:HG2	4:I:1315:GLU:H	1.54	0.72
3:G:214:ARG:HE	3:G:216:ASP:HB2	1.55	0.72
2:T:73:GLN:NE2	3:N:204:GLY:O	2.22	0.72
3:N:160:ASP:OD1	3:N:161:VAL:N	2.23	0.72
3:P:550:ASP:OD1	3:P:577:ARG:NH2	2.22	0.72
3:G:200:VAL:HG11	3:H:111:ASN:HD21	1.53	0.72
4:L:998:ASP:OD1	4:L:1001:ARG:NH2	2.23	0.72
4:K:611:GLU:OE1	4:K:792:ARG:NH1	2.20	0.72
4:J:241:SER:O	4:J:243:ALA:N	2.23	0.71
3:P:85:GLU:HA	3:P:88:ASN:HD21	1.55	0.71
2:d:78:ILE:HG12	3:P:211:MET:HE1	1.71	0.71
3:R:130:ARG:HH21	3:G:156:ILE:HG12	1.55	0.71
4:I:978:GLU:HA	4:L:371:VAL:HB	1.71	0.71
4:K:369:LEU:HD11	4:L:24:GLY:H	1.55	0.71
3:Q:661:LEU:HD23	3:Q:672:LEU:HD11	1.73	0.71
4:I:968:ILE:HG13	4:I:969:VAL:HG23	1.73	0.71
4:J:442:ASN:O	4:J:882:LYS:NZ	2.24	0.71
4:K:1247:GLU:HB3	4:K:1249:MET:HE3	1.72	0.71
3:Q:142:ARG:NH1	3:Q:145:GLU:OE2	2.23	0.71
3:O:551:LYS:HD2	4:K:308:LEU:HD11	1.72	0.71
3:O:589:ASN:HD22	3:O:592:MET:HG3	1.54	0.71
3:O:229:LEU:HD23	3:P:84:ARG:HH21	1.56	0.71
3:R:85:GLU:HA	3:R:88:ASN:HD21	1.56	0.71
3:R:542:SER:HB2	3:R:545:GLN:HB2	1.73	0.71
3:M:622:PRO:HB2	3:M:625:MET:HG3	1.73	0.70
3:Q:551:LYS:HD2	4:L:308:LEU:HD11	1.73	0.70
4:I:241:SER:O	4:I:243:ALA:N	2.23	0.70
4:J:611:GLU:OE1	4:J:792:ARG:NH1	2.24	0.70
4:J:1241:MET:O	4:J:1245:LEU:HG	1.92	0.70
4:K:257:ASP:HB3	4:K:260:LYS:HB2	1.73	0.70
4:I:611:GLU:OE1	4:I:792:ARG:NH1	2.22	0.70

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
4:L:553:ALA:O	4:L:554:THR:HG22	1.90	0.70
4:L:611:GLU:OE1	4:L:792:ARG:NH1	2.24	0.70
4:L:968:ILE:HG13	4:L:969:VAL:HG23	1.74	0.70
3:Q:520:MET:HB3	3:Q:526:ILE:HD11	1.72	0.70
3:P:682:ARG:HH22	3:P:684:ASP:HB2	1.57	0.69
4:I:456:PHE:O	4:I:457:GLU:HG3	1.92	0.69
4:J:819:LYS:HB3	4:J:823:GLU:HB2	1.74	0.69
2:S:49:ASN:OD1	3:H:338:LYS:NZ	2.25	0.69
3:P:671:TYR:CE1	3:P:682:ARG:HG3	2.27	0.69
4:K:1090:VAL:HG11	4:K:1110:SER:HB2	1.74	0.69
3:N:517:THR:OG1	3:N:586:ARG:NH2	2.25	0.69
3:Q:455:LYS:HD3	3:Q:460:ARG:HH11	1.58	0.69
1:D:39:ASP:OD2	3:O:323:ARG:NH2	2.25	0.69
3:H:262:MET:HE2	3:H:262:MET:HA	1.73	0.69
4:I:143:LEU:HD21	4:I:154:LEU:HD23	1.74	0.69
4:J:907:GLY:O	4:J:914:ARG:HB3	1.92	0.69
4:I:757:LEU:HD11	4:I:849:MET:HG3	1.74	0.69
4:L:552:GLY:HA2	4:L:648:GLY:HA3	1.74	0.69
1:C:47:GLU:OE1	3:R:362:GLN:NE2	2.26	0.69
2:f:57:ARG:HH22	3:Q:327:GLU:HB3	1.57	0.69
3:O:568:MET:HE2	3:O:601:LEU:HD11	1.74	0.69
4:L:1059:ASP:O	4:L:1061:GLN:N	2.26	0.69
3:G:661:LEU:HD23	3:G:672:LEU:HD11	1.74	0.68
4:I:1116:LEU:HA	4:I:1121:LEU:HD23	1.75	0.68
2:g:74:LYS:NZ	3:G:210:ASP:OD2	2.26	0.68
4:K:1059:ASP:O	4:K:1061:GLN:N	2.25	0.68
4:L:819:LYS:HB3	4:L:823:GLU:HB2	1.72	0.68
4:J:40:PHE:HA	4:J:53:MET:HE1	1.74	0.68
3:N:671:TYR:CE1	3:N:682:ARG:HG3	2.29	0.68
4:J:456:PHE:O	4:J:457:GLU:HG3	1.94	0.68
3:P:118:ASP:OD1	3:P:139:ARG:NH1	2.26	0.68
4:J:959:THR:HG21	4:J:1078:LYS:HE2	1.75	0.68
4:L:57:THR:HG23	4:L:74:ASP:HB3	1.76	0.68
4:K:968:ILE:HG13	4:K:969:VAL:HG23	1.75	0.68
4:K:1237:VAL:HG22	4:L:968:ILE:HG21	1.76	0.68
2:e:107:ILE:HD11	3:R:172:THR:HG23	1.76	0.67
2:h:73:GLN:NE2	3:H:204:GLY:O	2.27	0.67
3:M:661:LEU:HD23	3:M:672:LEU:HD11	1.77	0.67
4:K:612:ARG:NH2	4:K:614:GLU:OE2	2.26	0.67
3:N:79:THR:H	3:N:82:GLN:HE21	1.40	0.67
3:R:479:ILE:O	3:R:536:ARG:NH1	2.28	0.67

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:O:103:ALA:O	3:O:107:LYS:HG3	1.95	0.67
4:J:1116:LEU:HA	4:J:1121:LEU:HD23	1.77	0.67
3:Q:622:PRO:HB2	3:Q:625:MET:HG3	1.77	0.67
4:I:819:LYS:HB3	4:I:823:GLU:HB2	1.76	0.67
4:J:257:ASP:HB3	4:J:260:LYS:HB2	1.75	0.67
4:J:64:LEU:CD1	4:J:89:GLN:OE1	2.34	0.67
4:K:729:LYS:NZ	4:K:733:ASP:OD2	2.27	0.67
4:K:757:LEU:HD11	4:K:849:MET:HG2	1.77	0.67
3:N:85:GLU:HA	3:N:88:ASN:ND2	2.10	0.67
3:H:542:SER:HB2	3:H:545:GLN:HB2	1.77	0.67
4:J:131:ILE:O	4:J:136:ARG:NH2	2.28	0.67
3:N:710:ASP:OD2	3:P:295:GLN:NE2	2.28	0.66
4:I:1241:MET:O	4:I:1245:LEU:HG	1.94	0.66
4:L:463:VAL:HG22	4:L:875:ASN:HD22	1.60	0.66
3:G:569:PRO:HG2	3:G:572:LEU:HD12	1.76	0.66
4:J:681:ARG:NH2	4:J:784:GLN:OE1	2.28	0.66
2:d:71:ASN:ND2	3:Q:118:ASP:OD1	2.29	0.66
3:G:279:ARG:NH1	3:G:284:GLU:OE2	2.29	0.66
3:P:404:PHE:HB2	3:P:416:PHE:HB3	1.78	0.66
4:L:496:GLU:OE1	4:L:1173:ARG:NH1	2.28	0.66
3:R:541:ARG:HB3	3:R:546:ARG:HH11	1.60	0.66
4:I:442:ASN:O	4:I:882:LYS:NZ	2.28	0.66
3:R:413:THR:HG21	3:R:416:PHE:CE2	2.31	0.65
3:N:195:ILE:HD12	3:N:239:GLN:HG2	1.77	0.65
3:R:121:MET:HE2	3:R:139:ARG:HB2	1.79	0.65
3:H:394:ARG:HE	3:H:399:TRP:HZ3	1.44	0.65
3:H:404:PHE:HB2	3:H:416:PHE:HB3	1.78	0.65
4:I:550:LYS:HG2	4:I:552:GLY:H	1.60	0.65
3:N:85:GLU:HA	3:N:88:ASN:HD21	1.61	0.65
3:P:201:GLU:OE1	3:Q:110:ARG:NH2	2.30	0.65
4:I:371:VAL:HB	4:J:978:GLU:HA	1.79	0.65
4:L:819:LYS:NZ	4:L:827:GLU:OE1	2.29	0.65
4:I:318:ARG:HG2	4:I:318:ARG:HH11	1.61	0.65
4:K:1237:VAL:O	4:K:1241:MET:HG2	1.96	0.65
3:N:255:ALA:HB2	3:N:301:THR:HG21	1.78	0.65
4:I:310:ASN:O	4:I:314:GLU:HG2	1.96	0.65
4:K:265:PRO:HG3	4:K:988:LEU:HD11	1.79	0.65
4:L:222:ASN:N	4:L:232:ASN:OD1	2.30	0.65
3:R:447:LEU:HD12	3:R:503:LEU:HD23	1.78	0.64
4:K:1049:ARG:HH12	4:K:1053:GLY:HA2	1.62	0.64
4:L:560:GLU:HG3	4:L:1204:VAL:HG21	1.79	0.64

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:Q:606:TYR:HB3	3:Q:626:MET:HE1	1.79	0.64
3:G:622:PRO:HB2	3:G:625:MET:HG3	1.80	0.64
3:G:680:ARG:HD2	4:I:256:PHE:HB2	1.78	0.64
4:J:675:LYS:O	4:J:679:ILE:HG12	1.98	0.64
3:M:606:TYR:HB3	3:M:626:MET:HE1	1.78	0.64
4:I:795:ASP:O	4:I:797:PHE:N	2.31	0.64
4:J:931:LEU:HD22	4:J:1153:MET:HG2	1.79	0.64
4:I:29:LEU:HD22	4:I:101:GLU:OE1	1.98	0.64
4:L:795:ASP:O	4:L:797:PHE:N	2.31	0.64
4:K:795:ASP:O	4:K:797:PHE:N	2.31	0.64
3:M:197:ASN:HD22	3:N:107:LYS:HB2	1.63	0.64
3:H:568:MET:HE2	3:H:601:LEU:HD11	1.80	0.64
4:K:310:ASN:O	4:K:314:GLU:HG2	1.98	0.64
3:O:344:GLN:HG3	3:O:345:MET:HG3	1.80	0.63
4:I:757:LEU:HD22	4:I:760:ILE:HD13	1.80	0.63
3:M:522:ASP:OD1	3:M:523:LYS:N	2.30	0.63
3:R:387:ILE:HA	3:R:407:MET:HE1	1.80	0.63
4:I:934:SER:HB2	4:I:1101:LEU:HB2	1.80	0.63
4:K:40:PHE:HA	4:K:53:MET:HE1	1.79	0.63
4:L:456:PHE:HA	4:L:466:GLU:HG2	1.79	0.63
3:N:413:THR:HG21	3:N:416:PHE:CE2	2.33	0.63
4:J:364:ALA:HA	4:J:443:SER:HB3	1.81	0.63
2:S:22:ALA:O	2:S:26:MET:HG2	1.98	0.63
4:K:263:ILE:HD11	4:K:1116:LEU:HD22	1.79	0.63
4:L:13:ASP:OD1	4:L:31:ARG:NH1	2.31	0.63
4:I:398:GLY:HA3	4:I:429:MET:SD	2.39	0.63
4:I:1059:ASP:O	4:I:1061:GLN:N	2.32	0.63
4:J:555:ALA:HB1	4:J:855:LEU:HD21	1.79	0.63
4:J:1059:ASP:O	4:J:1061:GLN:N	2.31	0.63
3:H:233:ALA:O	3:H:234:ILE:C	2.42	0.63
3:H:522:ASP:OD1	3:H:523:LYS:N	2.32	0.63
4:J:371:VAL:HB	4:K:978:GLU:HA	1.79	0.63
4:K:345:MET:HG2	4:K:1300:GLN:OE1	1.98	0.63
3:N:550:ASP:OD1	3:N:577:ARG:NH2	2.23	0.63
3:Q:214:ARG:HE	3:Q:216:ASP:HB2	1.63	0.63
4:I:357:ASN:ND2	4:I:433:GLU:OE1	2.32	0.63
4:J:1049:ARG:HH12	4:J:1053:GLY:HA2	1.64	0.63
3:P:323:ARG:HA	3:P:363:MET:HE1	1.80	0.63
3:Q:233:ALA:O	3:Q:234:ILE:C	2.41	0.63
4:L:995:TYR:HE1	4:L:1189:SER:HB2	1.63	0.63
4:L:1049:ARG:HH12	4:L:1053:GLY:HA2	1.64	0.63

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:N:385:ASP:OD1	3:N:389:LYS:HE3	1.99	0.63
3:R:700:GLU:HG2	3:G:368:LYS:HB3	1.80	0.63
4:K:60:THR:HA	4:K:63:ALA:HB2	1.81	0.63
4:L:988:LEU:HD22	4:L:1131:ASN:HD22	1.63	0.63
3:N:520:MET:HG2	3:N:526:ILE:HD12	1.81	0.62
3:N:700:GLU:HG2	3:O:368:LYS:HB3	1.80	0.62
4:K:550:LYS:HG2	4:K:552:GLY:H	1.64	0.62
3:N:388:ASP:OD1	3:N:428:LYS:NZ	2.32	0.62
3:N:657:THR:HB	3:N:660:GLN:HG3	1.81	0.62
3:G:234:ILE:O	3:G:235:PRO:C	2.41	0.62
3:G:568:MET:HE2	3:G:601:LEU:HD11	1.80	0.62
4:J:1237:VAL:O	4:J:1241:MET:HG2	1.99	0.62
3:G:70:ARG:O	3:G:74:ILE:HD12	1.99	0.62
4:J:13:ASP:OD1	4:J:31:ARG:NH1	2.32	0.62
4:K:371:VAL:HB	4:L:978:GLU:HA	1.82	0.62
4:L:577:MET:O	4:L:581:MET:HG3	1.97	0.62
3:Q:199:ARG:NH1	3:Q:246:GLN:OE1	2.27	0.62
4:I:64:LEU:HG	4:L:479:ASP:HA	1.80	0.62
4:J:222:ASN:N	4:J:232:ASN:OD1	2.31	0.62
4:L:550:LYS:HG2	4:L:552:GLY:H	1.64	0.62
3:N:599:LYS:HE3	3:O:472:GLN:OE1	1.99	0.62
3:O:569:PRO:HG2	3:O:572:LEU:HD12	1.80	0.62
3:Q:605:THR:HG22	3:Q:622:PRO:HA	1.82	0.62
4:I:988:LEU:HD23	4:I:1131:ASN:HD22	1.64	0.62
4:J:550:LYS:HG2	4:J:552:GLY:H	1.63	0.62
4:L:976:THR:O	4:L:977:GLN:HG2	1.99	0.62
4:J:988:LEU:HD22	4:J:1131:ASN:HD22	1.65	0.62
4:J:993:THR:HG23	4:J:1085:LEU:HD21	1.80	0.62
4:K:754:ILE:HA	4:K:858:ARG:HH11	1.64	0.62
3:N:295:GLN:NE2	3:H:710:ASP:OD2	2.33	0.62
3:P:541:ARG:HB3	3:P:546:ARG:HH11	1.63	0.62
3:Q:517:THR:HG23	3:Q:531:ILE:HG21	1.80	0.62
3:R:517:THR:OG1	3:R:586:ARG:NH2	2.32	0.62
4:K:907:GLY:O	4:K:914:ARG:HB3	1.98	0.62
3:M:517:THR:HA	3:M:520:MET:HE2	1.80	0.62
3:O:203:ASN:O	3:O:207:GLN:HB2	1.99	0.62
3:R:636:GLU:OE1	3:R:639:ARG:NH2	2.32	0.62
4:I:324:LEU:HD23	4:I:345:MET:HE3	1.82	0.62
3:P:93:LEU:HD13	3:P:95:GLN:HE21	1.64	0.62
4:I:449:MET:HE2	4:I:468:LEU:HD21	1.82	0.62
4:J:739:SER:OG	4:J:740:HIS:ND1	2.32	0.62

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:O:372:LYS:HE3	3:O:376:ASP:OD1	2.00	0.62
3:P:657:THR:HB	3:P:660:GLN:HG3	1.81	0.62
3:G:79:THR:H	3:G:82:GLN:HE21	1.45	0.62
4:I:40:PHE:HA	4:I:53:MET:HE1	1.82	0.62
4:I:555:ALA:HB1	4:I:855:LEU:HD21	1.82	0.62
3:P:522:ASP:OD1	3:P:523:LYS:N	2.33	0.61
3:H:188:GLN:O	3:H:192:LYS:HG3	1.99	0.61
4:J:262:THR:HG22	4:J:263:ILE:HG13	1.82	0.61
4:I:1049:ARG:HH12	4:I:1053:GLY:HA2	1.65	0.61
4:L:675:LYS:O	4:L:679:ILE:HG12	2.00	0.61
3:O:55:MET:HA	3:O:55:MET:HE3	1.82	0.61
3:P:388:ASP:OD1	3:P:428:LYS:NZ	2.33	0.61
3:R:85:GLU:HA	3:R:88:ASN:ND2	2.15	0.61
3:H:443:ASP:HB3	3:H:499:ALA:HB2	1.82	0.61
4:I:456:PHE:HA	4:I:466:GLU:HG2	1.81	0.61
4:J:307:LYS:NZ	4:J:311:GLU:OE1	2.33	0.61
3:R:568:MET:HE3	3:R:601:LEU:HD11	1.82	0.61
4:J:593:MET:HE3	4:J:598:THR:HG22	1.83	0.61
3:P:162:ASP:OD1	3:P:165:ARG:NH2	2.33	0.61
4:K:976:THR:O	4:K:977:GLN:HG2	2.00	0.61
3:Q:517:THR:HA	3:Q:520:MET:HE2	1.80	0.61
4:L:131:ILE:O	4:L:136:ARG:NH2	2.33	0.61
3:O:646:ARG:HG3	3:O:661:LEU:HD12	1.82	0.61
3:P:112:ALA:HB2	3:P:150:TYR:HD2	1.65	0.61
3:P:255:ALA:HB1	3:P:301:THR:HG21	1.83	0.61
4:L:1163:PRO:HB2	4:L:1166:LYS:HZ1	1.66	0.61
3:M:641:ILE:HD11	4:J:245:MET:HE2	1.83	0.61
3:Q:568:MET:HE2	3:Q:601:LEU:HD11	1.83	0.61
3:G:606:TYR:HB3	3:G:626:MET:HE1	1.81	0.61
4:J:998:ASP:OD1	4:J:1001:ARG:NH2	2.33	0.61
2:e:119:ARG:NH1	3:G:90:GLY:O	2.31	0.61
3:O:526:ILE:HA	3:P:480:ASN:HD21	1.66	0.61
3:G:692:TRP:CD1	4:I:245:MET:HE1	2.35	0.61
3:M:543:LYS:HE3	3:H:551:LYS:HZ3	1.66	0.61
3:G:187:SER:O	3:G:191:GLN:HG3	2.01	0.61
4:J:265:PRO:HG3	4:J:988:LEU:HD11	1.83	0.61
4:J:904:ARG:HB3	4:J:1311:VAL:HG13	1.83	0.61
4:L:265:PRO:HG3	4:L:988:LEU:HD11	1.83	0.61
4:L:328:LEU:HD13	4:L:332:ILE:HD13	1.82	0.61
3:P:85:GLU:HA	3:P:88:ASN:ND2	2.16	0.60
4:I:526:ASP:HB3	4:I:529:ILE:HB	1.83	0.60

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
4:J:328:LEU:HD13	4:J:332:ILE:HD13	1.82	0.60
4:K:360:LEU:HA	4:K:446:LEU:HD12	1.82	0.60
1:F:47:GLU:OE1	3:G:362:GLN:NE2	2.24	0.60
3:M:701:GLU:O	3:M:705:GLU:HG2	2.01	0.60
3:N:500:ASP:OD1	3:N:632:PRO:HB3	2.01	0.60
3:G:692:TRP:HD1	4:I:245:MET:HE1	1.65	0.60
4:I:681:ARG:HH12	4:L:1243:SER:HB2	1.65	0.60
4:J:552:GLY:HA2	4:J:648:GLY:HA3	1.83	0.60
2:f:35:ARG:HH21	3:G:115:LEU:HD22	1.66	0.60
3:M:214:ARG:HE	3:M:216:ASP:HB2	1.66	0.60
3:O:605:THR:HG22	3:O:622:PRO:HA	1.83	0.60
3:Q:138:TYR:O	3:Q:142:ARG:HG2	2.00	0.60
3:Q:526:ILE:HA	3:R:480:ASN:HD21	1.66	0.60
3:R:490:MET:HE1	3:R:521:MET:HE1	1.82	0.60
3:R:671:TYR:CE1	3:R:682:ARG:HG3	2.36	0.60
3:G:172:THR:O	3:G:176:ILE:HG13	2.01	0.60
4:J:795:ASP:O	4:J:797:PHE:N	2.34	0.60
4:K:995:TYR:HE1	4:K:1189:SER:HB2	1.66	0.60
3:P:542:SER:HB2	3:P:545:GLN:HB2	1.81	0.60
3:G:599:LYS:NZ	3:G:603:GLU:OE2	2.34	0.60
3:Q:522:ASP:OD1	3:Q:523:LYS:N	2.33	0.60
4:I:509:ILE:HD11	4:I:854:ILE:HG12	1.82	0.60
4:J:596:GLU:HA	4:J:1192:GLY:HA2	1.82	0.60
4:J:1089:LYS:NZ	4:J:1091:SER:OG	2.23	0.60
4:L:934:SER:HB2	4:L:1101:LEU:HB2	1.82	0.60
3:R:522:ASP:OD1	3:R:523:LYS:N	2.32	0.60
3:R:678:GLN:NE2	4:L:449:MET:HE1	2.12	0.60
4:K:1190:GLN:HG2	4:K:1290:GLU:HB3	1.82	0.60
3:M:468:THR:HG21	3:H:602:LYS:HE3	1.84	0.60
3:N:542:SER:HB2	3:N:545:GLN:HB2	1.82	0.60
3:O:233:ALA:O	3:O:234:ILE:C	2.45	0.60
4:I:612:ARG:NH2	4:I:614:GLU:OE2	2.34	0.60
4:I:723:THR:HG23	4:I:726:MET:H	1.66	0.60
4:J:641:MET:HE1	4:J:766:LEU:HD21	1.83	0.60
4:J:934:SER:HB2	4:J:1101:LEU:HB2	1.82	0.60
4:J:1163:PRO:HG2	4:J:1166:LYS:HE2	1.82	0.60
2:S:52:LEU:O	2:S:57:ARG:NH1	2.33	0.60
3:Q:569:PRO:HG2	3:Q:572:LEU:HD12	1.83	0.60
3:N:541:ARG:HB3	3:N:546:ARG:HH11	1.67	0.60
3:P:262:MET:HE2	3:P:262:MET:HA	1.84	0.60
4:I:673:HIS:CD2	4:I:946:PRO:HA	2.37	0.60

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
4:J:1041:SER:O	4:J:1045:GLU:HG3	2.00	0.60
2:g:73:GLN:NE2	3:G:208:ASP:OD2	2.30	0.60
2:h:72:MET:HE3	2:h:75:VAL:HG11	1.83	0.60
3:M:352:LEU:O	3:M:356:GLN:HG2	2.02	0.60
3:H:119:ASP:OD1	3:H:120:VAL:N	2.35	0.60
3:O:522:ASP:OD1	3:O:523:LYS:N	2.34	0.59
4:I:131:ILE:O	4:I:136:ARG:NH2	2.35	0.59
4:J:754:ILE:HA	4:J:858:ARG:HH11	1.67	0.59
3:M:692:TRP:CD1	4:J:245:MET:HE1	2.37	0.59
3:M:692:TRP:HD1	4:J:245:MET:HE1	1.67	0.59
3:N:404:PHE:HB2	3:N:416:PHE:HB3	1.84	0.59
4:L:907:GLY:O	4:L:914:ARG:HB3	2.00	0.59
3:M:242:GLN:NE2	3:H:327:GLU:OE2	2.35	0.59
3:R:541:ARG:HD2	3:R:546:ARG:HD3	1.83	0.59
3:H:467:VAL:HG12	3:H:511:GLN:HG3	1.83	0.59
3:O:642:LEU:HD21	3:O:670:ILE:HD13	1.85	0.59
3:H:622:PRO:HG2	3:H:625:MET:HE3	1.84	0.59
4:I:577:MET:O	4:I:581:MET:HG2	2.02	0.59
3:H:268:LYS:HA	3:H:277:THR:HA	1.83	0.59
4:I:524:SER:HB2	4:I:844:GLU:OE1	2.02	0.59
4:I:1137:ALA:O	4:I:1141:ILE:HG12	2.03	0.59
4:J:517:ILE:HG21	4:J:851:THR:OG1	2.03	0.59
4:K:13:ASP:OD1	4:K:31:ARG:NH1	2.32	0.59
2:c:72:MET:SD	3:P:118:ASP:HB3	2.43	0.59
3:M:384:LEU:HD21	3:M:427:LYS:HE2	1.84	0.59
3:M:473:GLU:OE2	3:M:487:THR:HA	2.03	0.59
3:P:669:SER:HB3	3:P:682:ARG:HH11	1.66	0.59
3:R:261:LEU:HA	3:R:264:VAL:HG22	1.83	0.59
3:H:87:LEU:HD11	3:H:94:TYR:HD1	1.67	0.59
4:I:222:ASN:N	4:I:232:ASN:OD1	2.36	0.59
3:P:417:LYS:NZ	3:P:419:SER:HB2	2.18	0.59
4:I:998:ASP:OD1	4:I:1001:ARG:NH2	2.33	0.59
4:L:1013:LEU:HD23	4:L:1049:ARG:HB2	1.83	0.59
3:M:642:LEU:HD21	3:M:670:ILE:HD13	1.85	0.59
3:O:599:LYS:NZ	3:P:472:GLN:OE1	2.36	0.59
3:R:50:LYS:NZ	3:R:159:GLU:OE2	2.35	0.59
3:R:255:ALA:HB2	3:R:301:THR:HG21	1.85	0.59
3:H:479:ILE:O	3:H:536:ARG:NH1	2.35	0.59
3:O:662:THR:HG23	4:K:1119:LYS:HE3	1.85	0.59
3:P:421:MET:HA	3:P:421:MET:HE3	1.85	0.59
3:P:522:ASP:OD1	3:P:523:LYS:HG2	2.03	0.59

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:P:622:PRO:HG2	3:P:625:MET:HE3	1.83	0.59
3:R:404:PHE:HB2	3:R:416:PHE:HB3	1.85	0.59
3:R:517:THR:HG22	3:R:521:MET:HE2	1.83	0.59
3:R:657:THR:HB	3:R:660:GLN:OE1	2.03	0.59
4:K:998:ASP:OD1	4:K:1001:ARG:NH2	2.35	0.59
3:O:611:ASP:N	3:O:658:ASN:HD21	2.00	0.59
4:L:757:LEU:HD22	4:L:760:ILE:HD13	1.85	0.59
4:L:1116:LEU:HA	4:L:1121:LEU:HD23	1.85	0.59
4:J:1317:ARG:H	4:J:1317:ARG:HD3	1.68	0.58
4:K:560:GLU:HG3	4:K:1204:VAL:HG21	1.84	0.58
4:L:574:TYR:OH	4:L:1290:GLU:OE2	2.20	0.58
3:M:387:ILE:HG12	3:M:407:MET:SD	2.43	0.58
3:M:472:GLN:OE1	3:H:599:LYS:HE3	2.02	0.58
3:N:306:ASN:HD21	3:N:351:TRP:HZ3	1.51	0.58
3:N:522:ASP:OD1	3:N:523:LYS:N	2.36	0.58
3:P:544:GLU:OE2	4:K:318:ARG:NH2	2.32	0.58
4:I:60:THR:HA	4:I:63:ALA:HB2	1.84	0.58
2:e:39:MET:HA	2:e:39:MET:HE2	1.84	0.58
3:R:214:ARG:HG2	3:R:216:ASP:H	1.67	0.58
3:H:130:ARG:N	3:H:134:GLU:OE2	2.35	0.58
4:I:416:LYS:NZ	4:I:418:GLU:H	1.99	0.58
4:I:825:LYS:O	4:I:829:LEU:HG	2.04	0.58
4:L:1237:VAL:O	4:L:1241:MET:HG2	2.03	0.58
3:N:184:ASN:O	3:N:188:GLN:HG2	2.03	0.58
3:O:380:SER:HA	3:O:383:LYS:HG2	1.85	0.58
3:O:622:PRO:HB2	3:O:625:MET:HG3	1.86	0.58
3:R:363:MET:HG2	3:G:286:GLN:HA	1.85	0.58
4:K:551:PHE:CD1	4:L:781:PRO:HD3	2.39	0.58
2:S:107:ILE:HD11	3:N:172:THR:HG23	1.86	0.58
3:H:196:MET:HE2	3:H:196:MET:HA	1.86	0.58
4:L:757:LEU:HD11	4:L:849:MET:HG3	1.85	0.58
2:g:107:ILE:HD11	3:H:172:THR:HG23	1.85	0.58
3:O:473:GLU:OE2	3:O:487:THR:HA	2.04	0.58
3:O:633:LYS:O	3:O:633:LYS:HD3	2.03	0.58
3:P:387:ILE:HA	3:P:407:MET:HE1	1.84	0.58
3:G:131:THR:HG22	3:G:134:GLU:HG3	1.85	0.58
3:G:255:ALA:HB2	3:G:301:THR:HG21	1.86	0.58
3:G:605:THR:HG22	3:G:622:PRO:HA	1.86	0.58
3:H:417:LYS:NZ	3:H:419:SER:HB2	2.19	0.58
3:H:549:ASP:OD2	3:H:585:TYR:OH	2.20	0.58
3:P:684:ASP:OD1	3:P:685:LYS:N	2.37	0.58

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
4:I:364:ALA:HA	4:I:443:SER:HB3	1.85	0.58
4:K:819:LYS:HB3	4:K:823:GLU:HB2	1.86	0.58
2:T:47:ILE:HA	2:T:52:LEU:HD21	1.85	0.58
3:M:455:LYS:HA	3:M:460:ARG:HG3	1.86	0.58
3:Q:387:ILE:HG12	3:Q:407:MET:SD	2.43	0.58
3:R:453:ASP:HB3	3:R:457:GLY:HA3	1.85	0.58
3:H:487:THR:HB	3:H:490:MET:HB2	1.86	0.58
4:K:333:ILE:HD11	4:K:426:ILE:HG13	1.86	0.58
4:K:463:VAL:HG11	4:K:871:LEU:HD21	1.86	0.58
1:E:40:GLU:OE1	3:Q:373:ALA:HB2	2.04	0.58
3:M:526:ILE:HA	3:N:480:ASN:HD21	1.68	0.58
3:P:413:THR:HG21	3:P:416:PHE:CE2	2.38	0.58
3:R:262:MET:HE2	3:R:262:MET:HA	1.85	0.58
4:J:592:LYS:HG2	4:J:593:MET:HG3	1.84	0.58
4:J:595:ARG:HG2	4:J:1192:GLY:HA3	1.85	0.58
4:J:681:ARG:NH1	4:J:784:GLN:HE22	2.02	0.58
4:K:439:ARG:NH1	4:K:883:ASN:OD1	2.36	0.58
3:N:622:PRO:HG2	3:N:625:MET:HE3	1.86	0.58
3:Q:259:ASP:HA	3:Q:262:MET:HE3	1.85	0.58
4:I:616:GLN:HG2	4:I:627:MET:HE2	1.86	0.58
4:I:988:LEU:CD2	4:I:1131:ASN:HD22	2.17	0.58
4:K:495:LYS:O	4:K:499:GLU:HG2	2.04	0.58
4:L:682:TYR:CE1	4:L:781:PRO:HD2	2.39	0.58
4:J:866:VAL:HG23	4:J:1201:VAL:HG23	1.86	0.57
4:J:899:VAL:HG13	4:J:1181:ALA:HB1	1.86	0.57
4:K:328:LEU:HD13	4:K:332:ILE:HD13	1.86	0.57
2:c:107:ILE:HD11	3:P:172:THR:HG23	1.85	0.57
2:d:35:ARG:HH21	3:Q:115:LEU:HD22	1.69	0.57
3:M:692:TRP:HA	4:J:245:MET:HE3	1.85	0.57
3:H:550:ASP:OD1	3:H:577:ARG:NH2	2.31	0.57
4:I:24:GLY:H	4:L:369:LEU:HD21	1.66	0.57
4:J:495:LYS:O	4:J:499:GLU:HG2	2.04	0.57
3:N:443:ASP:HB3	3:N:499:ALA:HB2	1.87	0.57
3:O:628:VAL:O	3:O:634:SER:OG	2.21	0.57
3:P:598:THR:HG22	3:P:602:LYS:HE2	1.85	0.57
3:Q:428:LYS:HE2	3:Q:449:TYR:CZ	2.39	0.57
3:G:363:MET:SD	3:H:286:GLN:HG3	2.44	0.57
3:G:692:TRP:HA	4:I:245:MET:HE3	1.86	0.57
3:H:125:LYS:HE3	3:H:178:LEU:HB3	1.87	0.57
3:H:413:THR:HG21	3:H:416:PHE:CE2	2.40	0.57
4:I:754:ILE:HD12	4:I:853:LYS:HD3	1.86	0.57

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
4:K:332:ILE:HG21	4:K:395:PHE:CD1	2.40	0.57
4:L:62:LYS:O	4:L:63:ALA:C	2.48	0.57
4:L:1190:GLN:HG2	4:L:1290:GLU:HB3	1.86	0.57
2:T:47:ILE:HD13	3:N:192:LYS:HA	1.87	0.57
4:L:1295:ASP:HB3	4:L:1298:THR:HG22	1.85	0.57
3:R:149:VAL:O	3:R:153:GLN:HG2	2.05	0.57
3:R:587:SER:HB3	3:G:479:ILE:HG12	1.87	0.57
4:K:985:TRP:O	4:K:993:THR:OG1	2.22	0.57
3:N:285:GLU:H	3:N:285:GLU:CD	2.13	0.57
3:O:363:MET:HG2	3:P:286:GLN:HA	1.87	0.57
4:I:682:TYR:OH	4:I:782:ASP:N	2.38	0.57
4:I:888:ALA:O	4:I:892:THR:HG23	2.05	0.57
4:I:1241:MET:SD	4:J:958:ALA:HB1	2.44	0.57
4:J:1190:GLN:HG2	4:J:1290:GLU:HB3	1.86	0.57
3:M:149:VAL:O	3:M:153:GLN:HG2	2.05	0.57
3:P:285:GLU:CD	3:P:285:GLU:H	2.12	0.57
3:R:261:LEU:HB3	3:R:291:MET:HE1	1.86	0.57
4:I:779:THR:HG21	4:L:538:ARG:HH12	1.70	0.57
4:K:593:MET:HE3	4:K:598:THR:HG22	1.85	0.57
4:L:555:ALA:HB1	4:L:855:LEU:HD21	1.87	0.57
1:A:48:VAL:O	3:N:358:GLN:NE2	2.38	0.57
3:M:493:LEU:HG	3:M:518:MET:HE1	1.87	0.57
3:P:374:LEU:HD13	3:Q:296:ARG:HA	1.86	0.57
4:L:723:THR:HG23	4:L:726:MET:H	1.70	0.57
3:M:201:GLU:OE1	3:N:110:ARG:NH1	2.37	0.56
3:O:358:GLN:O	3:O:362:GLN:HG3	2.05	0.56
3:O:556:ALA:O	3:O:560:SER:OG	2.22	0.56
4:I:1071:ASP:OD1	4:I:1074:ARG:NH2	2.38	0.56
4:L:538:ARG:NH1	4:L:557:ASP:OD2	2.38	0.56
3:H:88:ASN:OD1	3:H:89:ASN:N	2.38	0.56
2:h:55:GLN:O	2:h:59:LYS:HG3	2.05	0.56
3:M:478:VAL:HG21	3:H:592:MET:HE3	1.86	0.56
3:N:187:SER:O	3:N:191:GLN:HG3	2.05	0.56
3:P:160:ASP:O	3:P:164:GLN:HG2	2.06	0.56
4:I:332:ILE:HG21	4:I:395:PHE:CD1	2.40	0.56
4:K:689:GLU:OE2	4:K:693:ARG:NH2	2.38	0.56
4:K:1243:SER:OG	4:L:784:GLN:NE2	2.37	0.56
4:L:121:GLU:HG2	4:L:125:LYS:HE3	1.87	0.56
4:L:140:ARG:HA	4:L:143:LEU:HG	1.87	0.56
4:L:524:SER:OG	4:L:530:ARG:HG3	2.05	0.56
2:T:77:ALA:O	2:T:81:ILE:HG13	2.06	0.56

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:H:72:ASN:HA	3:H:75:ILE:HG22	1.88	0.56
2:e:64:SER:O	2:e:68:THR:HG23	2.05	0.56
3:O:520:MET:HE3	3:O:526:ILE:HD11	1.86	0.56
2:c:119:ARG:NH1	3:Q:90:GLY:O	2.31	0.56
3:P:79:THR:N	3:P:82:GLN:HE21	1.95	0.56
3:P:333:LYS:HG3	3:P:352:LEU:HD21	1.87	0.56
3:G:645:ALA:HA	4:I:227:PHE:HE2	1.70	0.56
3:H:542:SER:O	3:H:544:GLU:N	2.39	0.56
4:I:495:LYS:O	4:I:499:GLU:HG2	2.05	0.56
3:M:517:THR:HG23	3:M:531:ILE:HG21	1.88	0.56
3:M:569:PRO:HG2	3:M:572:LEU:HD12	1.87	0.56
3:M:645:ALA:HA	4:J:227:PHE:HE2	1.70	0.56
3:N:592:MET:HE1	3:O:474:TRP:CD1	2.40	0.56
3:G:47:ARG:HA	3:G:50:LYS:HE2	1.86	0.56
4:I:899:VAL:HG13	4:I:1181:ALA:HB1	1.88	0.56
4:L:1061:GLN:HE22	4:L:1065:MET:HE3	1.71	0.56
3:M:394:ARG:NH1	3:M:402:THR:OG1	2.38	0.56
3:M:598:THR:O	3:M:602:LYS:HG3	2.06	0.56
3:Q:151:ALA:HB2	3:Q:163:TYR:CZ	2.41	0.56
3:R:285:GLU:CD	3:R:285:GLU:H	2.14	0.56
4:I:968:ILE:HG21	4:L:1237:VAL:HG22	1.88	0.56
4:L:593:MET:HB2	4:L:597:GLU:HB3	1.86	0.56
4:L:1241:MET:O	4:L:1245:LEU:HG	2.06	0.56
3:Q:358:GLN:O	3:Q:362:GLN:HG3	2.05	0.56
3:G:214:ARG:O	3:G:263:ARG:NH2	2.39	0.56
4:I:1004:MET:HA	4:I:1004:MET:HE3	1.87	0.56
3:O:613:VAL:HG23	3:O:614:ASP:H	1.71	0.56
3:H:570:ALA:HA	3:H:573:ARG:HG2	1.87	0.56
3:M:500:ASP:OD1	3:M:632:PRO:HB3	2.05	0.55
3:O:313:ILE:HD13	3:O:332:ILE:HD13	1.89	0.55
3:P:117:ASP:O	3:P:121:MET:HG2	2.05	0.55
3:P:549:ASP:OD2	3:P:585:TYR:OH	2.24	0.55
3:R:589:ASN:HA	3:G:539:VAL:CG2	2.36	0.55
4:I:1243:SER:CB	4:J:784:GLN:HE21	2.19	0.55
4:J:333:ILE:HD11	4:J:426:ILE:HG13	1.88	0.55
3:P:541:ARG:HD2	3:P:546:ARG:HD3	1.87	0.55
3:R:123:LYS:HA	3:R:123:LYS:HE2	1.88	0.55
3:G:372:LYS:NZ	3:G:376:ASP:OD1	2.20	0.55
4:I:328:LEU:HD13	4:I:332:ILE:HD13	1.88	0.55
4:L:1120:PHE:HE2	4:L:1301:LEU:HD23	1.71	0.55
3:H:359:VAL:HA	3:H:362:GLN:HG3	1.88	0.55

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
4:I:1120:PHE:HE2	4:I:1301:LEU:HD23	1.71	0.55
4:J:1031:VAL:HG11	4:J:1075:LEU:HD13	1.89	0.55
4:K:723:THR:HG22	4:K:726:MET:HG3	1.87	0.55
4:K:899:VAL:HG13	4:K:1181:ALA:HB1	1.88	0.55
3:O:435:MET:HE1	3:P:318:ASN:HD22	1.71	0.55
3:G:387:ILE:HG12	3:G:407:MET:SD	2.47	0.55
4:I:1019:ARG:HA	4:I:1022:LYS:NZ	2.22	0.55
4:K:1277:GLU:HG2	4:L:974:TYR:CD1	2.42	0.55
4:I:976:THR:O	4:I:977:GLN:HG2	2.06	0.55
4:J:398:GLY:HA3	4:J:429:MET:SD	2.47	0.55
3:M:381:MET:HE2	3:N:299:PHE:HZ	1.71	0.55
3:P:264:VAL:HA	3:P:267:LYS:HG3	1.87	0.55
3:P:510:ASP:OD1	3:P:511:GLN:HG2	2.07	0.55
3:P:542:SER:O	3:P:544:GLU:N	2.38	0.55
4:J:822:LYS:NZ	4:J:826:ASP:OD2	2.39	0.55
4:J:1295:ASP:HB3	4:J:1298:THR:HG22	1.88	0.55
4:L:434:ALA:HB3	4:L:447:SER:HB3	1.89	0.55
4:L:604:ARG:NH1	4:L:950:ASP:OD2	2.39	0.55
3:M:296:ARG:HA	3:H:374:LEU:HD13	1.87	0.55
3:N:599:LYS:HZ3	3:N:603:GLU:CD	2.15	0.55
3:O:120:VAL:HG21	3:O:139:ARG:HG3	1.88	0.55
3:G:149:VAL:O	3:G:153:GLN:HG2	2.06	0.55
4:I:360:LEU:HA	4:I:446:LEU:HD12	1.88	0.55
4:J:761:GLU:OE1	4:J:825:LYS:NZ	2.40	0.55
4:J:925:ALA:O	4:J:929:LYS:HG2	2.07	0.55
1:D:47:GLU:OE1	3:O:362:GLN:NE2	2.25	0.55
3:Q:172:THR:O	3:Q:176:ILE:HG12	2.06	0.55
3:R:391:PHE:O	3:R:395:ILE:HG12	2.07	0.55
4:J:150:MET:SD	4:J:153:GLN:NE2	2.80	0.55
4:J:1023:GLU:HA	4:J:1026:LEU:HD12	1.89	0.55
4:K:577:MET:O	4:K:581:MET:HG2	2.06	0.55
4:K:1197:MET:HG2	4:K:1287:SER:HB2	1.89	0.55
2:e:69:SER:OG	3:R:119:ASP:OD1	2.25	0.55
3:M:356:GLN:O	3:M:360:GLN:HG2	2.05	0.55
3:P:170:ASP:OD1	3:P:173:GLU:HG3	2.06	0.55
3:R:160:ASP:OD1	3:R:161:VAL:N	2.40	0.55
3:G:132:ARG:HD2	3:G:179:TYR:CD1	2.41	0.55
4:I:1242:GLY:HA2	4:I:1245:LEU:HD12	1.87	0.55
4:J:604:ARG:NH1	4:J:950:ASP:OD2	2.40	0.55
4:J:952:VAL:HG23	4:J:1073:TRP:HH2	1.72	0.55
4:J:995:TYR:HE1	4:J:1189:SER:HB2	1.72	0.55

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
4:J:1090:VAL:HG11	4:J:1110:SER:HA	1.89	0.55
4:K:359:ALA:O	4:K:360:LEU:HB2	2.07	0.55
4:K:1120:PHE:HE2	4:K:1301:LEU:HD23	1.71	0.55
4:K:1281:MET:O	4:K:1285:MET:HG2	2.07	0.55
4:I:333:ILE:HD11	4:I:426:ILE:HG13	1.89	0.55
4:J:30:LEU:HD11	4:J:87:ALA:HB2	1.87	0.55
4:K:517:ILE:HG21	4:K:851:THR:OG1	2.07	0.55
4:L:360:LEU:HA	4:L:446:LEU:HD12	1.87	0.55
4:L:509:ILE:HD11	4:L:854:ILE:HG12	1.89	0.55
3:N:261:LEU:HA	3:N:264:VAL:HG22	1.88	0.54
3:N:541:ARG:HD2	3:N:546:ARG:HD3	1.88	0.54
3:P:261:LEU:HA	3:P:264:VAL:HG22	1.89	0.54
3:G:550:ASP:O	3:G:554:GLU:HG2	2.07	0.54
3:G:642:LEU:HD21	3:G:670:ILE:HD13	1.89	0.54
3:H:285:GLU:CD	3:H:285:GLU:H	2.15	0.54
4:I:529:ILE:HD13	4:I:831:LEU:HD13	1.88	0.54
4:I:907:GLY:O	4:I:914:ARG:HB3	2.07	0.54
3:M:313:ILE:HD13	3:M:332:ILE:HD13	1.89	0.54
3:N:542:SER:O	3:N:544:GLU:N	2.40	0.54
3:G:662:THR:HG23	4:I:1119:LYS:HE3	1.88	0.54
3:H:162:ASP:OD1	3:H:165:ARG:NH1	2.41	0.54
4:J:359:ALA:O	4:J:360:LEU:HB2	2.08	0.54
4:K:1295:ASP:HB3	4:K:1298:THR:HG22	1.88	0.54
3:N:666:GLN:HB2	3:N:671:TYR:HE2	1.72	0.54
3:Q:363:MET:HE3	3:R:286:GLN:HB2	1.90	0.54
4:J:1021:GLU:OE2	4:J:1040:LYS:HE2	2.07	0.54
4:L:332:ILE:HG21	4:L:395:PHE:CD1	2.42	0.54
1:E:46:THR:O	3:P:392:GLN:NE2	2.40	0.54
3:M:235:PRO:HB3	3:N:94:TYR:CE1	2.43	0.54
3:G:451:GLN:HG2	3:G:503:LEU:HD11	1.88	0.54
4:L:357:ASN:ND2	4:L:433:GLU:OE2	2.41	0.54
3:P:82:GLN:HA	3:P:85:GLU:HG3	1.89	0.54
3:P:467:VAL:HG12	3:P:511:GLN:HG3	1.90	0.54
3:R:374:LEU:HD13	3:G:296:ARG:HA	1.90	0.54
3:M:70:ARG:O	3:M:74:ILE:HG12	2.08	0.54
3:O:323:ARG:O	3:O:327:GLU:HG3	2.07	0.54
3:O:538:THR:HA	3:O:540:LYS:HE3	1.90	0.54
3:P:47:ARG:HA	3:P:50:LYS:HE2	1.89	0.54
3:R:233:ALA:O	3:R:234:ILE:C	2.50	0.54
3:G:556:ALA:O	3:G:560:SER:OG	2.24	0.54
4:I:1197:MET:HG2	4:I:1287:SER:HB2	1.88	0.54

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
4:J:360:LEU:HA	4:J:446:LEU:HD12	1.88	0.54
4:J:535:ASP:HB3	4:J:817:THR:HG22	1.90	0.54
4:J:1281:MET:O	4:J:1285:MET:HG2	2.08	0.54
4:K:757:LEU:HD22	4:K:760:ILE:HD13	1.90	0.54
2:T:72:MET:HE3	3:O:121:MET:HB2	1.89	0.54
3:N:349:ARG:O	3:N:353:ILE:HD12	2.08	0.54
3:Q:71:SER:O	3:Q:75:ILE:HG12	2.08	0.54
4:J:345:MET:HE2	4:J:1300:GLN:HB2	1.90	0.54
4:J:577:MET:O	4:J:581:MET:HG2	2.06	0.54
4:K:451:THR:HA	4:K:454:MET:HE3	1.88	0.54
4:K:850:ASP:O	4:K:854:ILE:HG13	2.08	0.54
4:L:224:ARG:HG3	4:L:252:LEU:HD11	1.89	0.54
4:L:456:PHE:N	4:L:466:GLU:OE2	2.41	0.54
4:L:866:VAL:HG13	4:L:1201:VAL:HG23	1.90	0.54
1:b:39:ASP:OD2	3:M:323:ARG:NH2	2.40	0.54
3:N:323:ARG:HA	3:N:363:MET:HE1	1.88	0.54
3:H:47:ARG:HA	3:H:50:LYS:HE2	1.90	0.54
4:I:775:ASP:OD1	4:I:789:ASN:ND2	2.41	0.54
4:K:514:PHE:HE1	4:K:851:THR:HG23	1.72	0.54
4:L:581:MET:HG2	4:L:602:ILE:HD12	1.90	0.54
3:O:346:THR:HG23	3:O:349:ARG:H	1.73	0.54
4:I:637:LYS:O	4:I:641:MET:HG3	2.08	0.54
4:K:11:ASP:O	4:K:1024:GLY:HA3	2.08	0.54
4:K:495:LYS:O	4:K:498:SER:OG	2.20	0.54
4:L:989:LEU:HA	4:L:1135:ASP:HB2	1.89	0.54
2:h:22:ALA:O	2:h:26:MET:HG2	2.08	0.54
3:R:520:MET:HB3	3:R:526:ILE:HD12	1.90	0.54
4:J:11:ASP:O	4:J:1024:GLY:HA3	2.07	0.54
4:J:682:TYR:CE1	4:J:781:PRO:HD2	2.43	0.54
3:N:283:GLY:O	3:N:287:TRP:N	2.36	0.53
3:R:89:ASN:OD1	3:R:90:GLY:N	2.40	0.53
3:H:211:MET:O	3:H:214:ARG:HG3	2.08	0.53
3:H:344:GLN:HG2	3:H:346:THR:HG23	1.91	0.53
4:L:866:VAL:HG11	4:L:1197:MET:HE3	1.89	0.53
4:K:434:ALA:HB3	4:K:447:SER:HB3	1.89	0.53
3:M:55:MET:HA	3:M:55:MET:HE3	1.90	0.53
3:M:322:PRO:HB2	3:M:363:MET:HE2	1.90	0.53
3:O:201:GLU:OE1	3:P:110:ARG:NH1	2.41	0.53
3:O:701:GLU:O	3:O:705:GLU:HG2	2.08	0.53
3:R:252:SER:HB2	3:R:293:THR:HG23	1.90	0.53
3:G:435:MET:HE1	3:H:318:ASN:HD22	1.74	0.53

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:H:261:LEU:HA	3:H:264:VAL:HG22	1.90	0.53
4:I:644:PRO:HG2	4:I:814:MET:HG2	1.89	0.53
4:J:524:SER:HB3	4:J:844:GLU:OE1	2.09	0.53
4:J:898:ILE:HG22	4:J:1144:MET:HE1	1.89	0.53
4:K:266:THR:HG23	4:K:269:ASN:H	1.73	0.53
1:A:48:VAL:HG12	1:A:50:GLY:H	1.72	0.53
3:M:611:ASP:N	3:M:658:ASN:HD21	2.05	0.53
3:N:322:PRO:HG2	3:N:366:TRP:HB2	1.90	0.53
3:P:417:LYS:HZ3	3:P:419:SER:HB2	1.72	0.53
3:P:435:MET:HE1	3:Q:318:ASN:HD22	1.73	0.53
3:R:79:THR:OG1	3:R:82:GLN:OE1	2.26	0.53
4:K:964:ALA:O	4:K:968:ILE:HG12	2.08	0.53
3:P:344:GLN:HG2	3:P:346:THR:HG23	1.90	0.53
3:H:493:LEU:HA	3:H:496:ILE:HG22	1.91	0.53
4:I:496:GLU:OE2	4:I:1173:ARG:NH1	2.41	0.53
4:I:1084:MET:O	4:I:1109:LYS:NZ	2.33	0.53
4:J:514:PHE:HE1	4:J:851:THR:HG23	1.74	0.53
4:L:885:TYR:O	4:L:889:GLN:HG3	2.08	0.53
4:L:1281:MET:O	4:L:1285:MET:HG2	2.08	0.53
3:O:255:ALA:HB2	3:O:301:THR:HG21	1.91	0.53
3:O:349:ARG:O	3:O:353:ILE:HG12	2.08	0.53
3:P:551:LYS:HZ3	3:Q:543:LYS:HE3	1.74	0.53
3:H:447:LEU:HA	3:H:450:LEU:HG	1.91	0.53
4:K:258:VAL:HG12	4:K:1088:HIS:HB3	1.91	0.53
2:T:40:GLU:O	2:T:44:GLN:HG2	2.07	0.53
2:g:37:GLN:O	2:g:41:ILE:HG13	2.08	0.53
3:M:276:THR:HB	3:M:280:GLU:HB2	1.91	0.53
3:N:125:LYS:HZ1	3:N:135:MET:HG3	1.73	0.53
3:N:568:MET:HE2	3:N:601:LEU:HD11	1.91	0.53
3:P:682:ARG:NH2	3:P:684:ASP:HB2	2.24	0.53
4:I:641:MET:HE1	4:I:766:LEU:HD21	1.89	0.53
4:K:79:PRO:HD2	4:K:1027:ARG:HH22	1.73	0.53
4:L:899:VAL:HG13	4:L:1181:ALA:HB1	1.90	0.53
3:M:255:ALA:HB2	3:M:301:THR:HG21	1.90	0.53
3:R:542:SER:O	3:R:544:GLU:N	2.41	0.53
4:I:456:PHE:N	4:I:466:GLU:OE2	2.42	0.53
4:J:369:LEU:HD21	4:K:24:GLY:H	1.74	0.53
4:L:364:ALA:HA	4:L:443:SER:HB3	1.90	0.53
4:L:643:ASN:O	4:L:646:ILE:HG13	2.09	0.53
3:M:214:ARG:O	3:M:263:ARG:NH2	2.42	0.53
3:Q:483:LEU:HD11	3:Q:490:MET:HE1	1.90	0.53

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:G:346:THR:HG23	3:G:349:ARG:H	1.72	0.53
4:I:712:MET:HE2	4:I:771:LEU:HD12	1.91	0.53
4:J:801:PRO:O	4:J:805:ARG:HG3	2.09	0.53
3:O:597:MET:HE3	3:O:601:LEU:HD11	1.91	0.53
3:O:606:TYR:HB3	3:O:626:MET:HE1	1.90	0.53
3:P:135:MET:HE1	3:P:179:TYR:CE1	2.44	0.53
3:P:569:PRO:HG2	3:P:572:LEU:HD13	1.91	0.53
3:P:666:GLN:HB2	3:P:671:TYR:HE2	1.74	0.53
4:I:359:ALA:O	4:I:360:LEU:HB2	2.09	0.53
4:L:380:VAL:HB	4:L:575:LYS:HD2	1.91	0.53
4:L:529:ILE:HD13	4:L:831:LEU:HD23	1.91	0.53
2:c:77:ALA:O	2:c:81:ILE:HG13	2.10	0.52
3:N:123:LYS:O	3:N:126:GLU:HG3	2.09	0.52
3:P:659:LYS:HG2	4:K:362:VAL:HG22	1.91	0.52
3:Q:598:THR:O	3:Q:602:LYS:HG3	2.09	0.52
3:R:684:ASP:OD1	3:R:685:LYS:N	2.42	0.52
3:G:84:ARG:O	3:G:88:ASN:ND2	2.41	0.52
3:G:112:ALA:HB2	3:G:150:TYR:CD2	2.44	0.52
4:I:641:MET:HG2	4:I:811:ILE:HD11	1.91	0.52
4:J:897:MET:HA	4:J:900:THR:HG22	1.91	0.52
4:L:439:ARG:NH1	4:L:883:ASN:OD1	2.42	0.52
4:L:661:HIS:HB3	4:L:769:ARG:HH21	1.73	0.52
3:N:583:VAL:HG11	3:N:596:GLN:HG3	1.90	0.52
3:O:138:TYR:O	3:O:142:ARG:HG2	2.08	0.52
3:Q:211:MET:O	3:Q:217:SER:HB2	2.09	0.52
3:Q:684:ASP:OD1	3:Q:684:ASP:N	2.38	0.52
4:I:514:PHE:HE1	4:I:851:THR:HG23	1.74	0.52
4:I:563:HIS:NE2	4:I:1202:GLY:O	2.33	0.52
4:I:1295:ASP:HB3	4:I:1298:THR:HG22	1.91	0.52
4:J:434:ALA:HB3	4:J:447:SER:HB3	1.92	0.52
4:K:456:PHE:N	4:K:466:GLU:OE2	2.41	0.52
4:K:682:TYR:HB3	4:K:686:GLY:HA3	1.91	0.52
4:L:914:ARG:O	4:L:919:LYS:HG3	2.09	0.52
2:T:71:ASN:HB2	3:O:118:ASP:OD1	2.09	0.52
2:g:68:THR:HG22	3:G:200:VAL:HG13	1.90	0.52
3:Q:229:LEU:HD23	3:R:84:ARG:HH21	1.74	0.52
3:R:72:ASN:HA	3:R:75:ILE:HG22	1.92	0.52
4:I:827:GLU:O	4:I:831:LEU:HG	2.08	0.52
4:J:496:GLU:HG2	4:J:1179:MET:HE3	1.92	0.52
4:L:428:PRO:HB2	4:L:897:MET:HE2	1.90	0.52
4:L:975:SER:HB3	4:L:978:GLU:HG3	1.91	0.52

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:e:53:SER:O	2:e:57:ARG:HG3	2.10	0.52
3:M:448:LYS:HB3	3:M:448:LYS:HZ2	1.75	0.52
3:M:479:ILE:HG12	3:H:587:SER:HB3	1.90	0.52
3:P:583:VAL:HG11	3:P:596:GLN:HG3	1.90	0.52
3:H:139:ARG:NH2	3:H:175:ASN:OD1	2.43	0.52
4:I:669:VAL:HG13	4:I:793:ASP:HB3	1.91	0.52
4:K:1240:ALA:O	4:K:1243:SER:OG	2.24	0.52
4:L:8:VAL:HG12	4:L:31:ARG:HH21	1.75	0.52
4:L:359:ALA:O	4:L:360:LEU:HB2	2.09	0.52
2:T:128:GLN:HG2	3:P:82:GLN:OE1	2.10	0.52
2:h:53:SER:O	2:h:57:ARG:HG2	2.10	0.52
3:M:132:ARG:HD2	3:M:179:TYR:CD1	2.45	0.52
3:M:211:MET:O	3:M:217:SER:HB2	2.09	0.52
3:M:641:ILE:HG21	3:M:688:LEU:HD11	1.92	0.52
3:N:206:LEU:HD12	3:N:247:ALA:HB1	1.92	0.52
3:N:510:ASP:OD1	3:N:511:GLN:HG2	2.09	0.52
3:G:614:ASP:OD1	3:H:455:LYS:NZ	2.42	0.52
4:I:140:ARG:HA	4:I:143:LEU:HG	1.90	0.52
4:I:434:ALA:HB3	4:I:447:SER:HB3	1.92	0.52
4:K:34:ALA:HB2	4:K:54:MET:HE1	1.90	0.52
4:K:496:GLU:HG2	4:K:1179:MET:HE3	1.92	0.52
4:K:661:HIS:HB3	4:K:769:ARG:HH21	1.75	0.52
4:K:810:ASP:OD1	4:K:821:THR:OG1	2.26	0.52
4:L:80:GLU:HG3	4:L:1027:ARG:HE	1.74	0.52
2:S:119:ARG:NH1	3:O:90:GLY:O	2.37	0.52
3:P:587:SER:O	3:P:589:ASN:N	2.42	0.52
3:R:197:ASN:O	3:R:201:GLU:HG2	2.10	0.52
3:G:132:ARG:HD2	3:G:179:TYR:HD1	1.75	0.52
3:G:201:GLU:OE1	3:H:110:ARG:NH1	2.40	0.52
3:H:121:MET:HE2	3:H:139:ARG:HB2	1.92	0.52
3:H:697:LYS:O	3:H:701:GLU:HG2	2.09	0.52
4:I:985:TRP:O	4:I:993:THR:OG1	2.23	0.52
4:I:1248:GLN:HG3	4:J:790:ASP:HA	1.92	0.52
4:J:661:HIS:HB3	4:J:769:ARG:HH21	1.74	0.52
4:L:405:SER:OG	4:L:423:ASN:OD1	2.24	0.52
3:N:592:MET:HE3	3:O:478:VAL:HG21	1.89	0.52
3:O:514:LEU:O	3:O:518:MET:HG3	2.10	0.52
3:P:322:PRO:HG2	3:P:366:TRP:HB2	1.92	0.52
3:R:338:LYS:NZ	3:H:88:ASN:O	2.38	0.52
3:R:622:PRO:HG2	3:R:625:MET:HE3	1.90	0.52
3:G:673:MET:HB3	3:G:680:ARG:HG2	1.91	0.52

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:H:669:SER:HB3	3:H:682:ARG:HH11	1.74	0.52
4:J:66:LEU:C	4:J:68:VAL:H	2.18	0.52
4:K:604:ARG:NH1	4:K:950:ASP:OD2	2.43	0.52
4:K:1116:LEU:HA	4:K:1121:LEU:HD23	1.92	0.52
4:K:1293:PRO:HD2	4:K:1299:GLN:NE2	2.25	0.52
3:Q:214:ARG:O	3:Q:263:ARG:NH2	2.43	0.52
3:Q:439:ASP:OD2	3:Q:495:ARG:NH1	2.43	0.52
3:Q:473:GLU:HG2	3:Q:490:MET:HE2	1.92	0.52
3:G:358:GLN:O	3:G:362:GLN:HG3	2.09	0.52
3:H:556:ALA:HB1	3:H:594:MET:HE3	1.91	0.52
4:K:777:SER:HB3	4:K:787:SER:HB3	1.91	0.52
4:L:594:SER:O	4:L:598:THR:HG23	2.10	0.52
2:S:69:SER:HB2	3:N:115:LEU:HA	1.92	0.52
2:T:47:ILE:HG12	2:T:48:GLN:HE21	1.74	0.52
2:c:64:SER:O	2:c:68:THR:HG23	2.10	0.52
3:M:206:LEU:HD22	3:M:247:ALA:HB1	1.92	0.52
3:M:391:PHE:O	3:M:395:ILE:HG13	2.09	0.52
3:O:336:LEU:HD12	3:O:352:LEU:HD22	1.91	0.52
3:R:47:ARG:HA	3:R:50:LYS:HE2	1.92	0.52
3:R:383:LYS:O	3:R:387:ILE:HG13	2.10	0.52
4:J:827:GLU:O	4:J:831:LEU:HG	2.10	0.52
4:K:535:ASP:HB3	4:K:817:THR:HG22	1.92	0.52
4:K:552:GLY:HA2	4:K:648:GLY:HA3	1.91	0.52
2:d:53:SER:O	2:d:57:ARG:HG2	2.10	0.52
3:M:112:ALA:HB2	3:M:150:TYR:CD1	2.45	0.52
3:G:251:ALA:O	3:G:257:GLY:HA3	2.09	0.52
3:G:598:THR:O	3:G:602:LYS:HG3	2.09	0.52
4:I:801:PRO:O	4:I:805:ARG:HG3	2.09	0.52
4:J:825:LYS:O	4:J:829:LEU:HG	2.10	0.52
4:J:904:ARG:NH2	4:J:1177:PRO:HG2	2.25	0.52
2:S:39:MET:HE1	3:N:154:PHE:CE1	2.45	0.51
2:g:57:ARG:NH2	3:R:327:GLU:OE2	2.42	0.51
3:M:646:ARG:HG3	3:M:661:LEU:HD22	1.89	0.51
3:R:680:ARG:NH1	4:L:423:ASN:HD22	2.07	0.51
3:G:700:GLU:HB3	3:H:368:LYS:HB3	1.91	0.51
4:K:669:VAL:HG13	4:K:793:ASP:HB3	1.92	0.51
4:L:1197:MET:SD	4:L:1287:SER:HB2	2.50	0.51
3:P:519:ASP:OD1	3:P:523:LYS:HG3	2.10	0.51
3:Q:160:ASP:O	3:Q:164:GLN:HG2	2.10	0.51
4:I:905:ALA:O	4:I:1309:ASN:ND2	2.42	0.51
4:J:689:GLU:OE2	4:J:693:ARG:NE	2.42	0.51

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
4:J:722:VAL:HG13	4:J:726:MET:HE2	1.92	0.51
4:L:1074:ARG:O	4:L:1078:LYS:HG2	2.10	0.51
3:M:232:GLY:HA3	3:N:100:ALA:HB2	1.92	0.51
3:Q:645:ALA:HA	4:L:227:PHE:HE2	1.76	0.51
3:R:206:LEU:HD12	3:R:247:ALA:HB1	1.92	0.51
3:H:682:ARG:HH12	3:H:684:ASP:HB2	1.75	0.51
4:I:701:SER:HB2	4:L:646:ILE:HG12	1.93	0.51
4:I:978:GLU:OE1	4:L:377:ARG:NH2	2.43	0.51
4:K:846:HIS:HA	4:K:849:MET:HE3	1.92	0.51
2:T:81:ILE:HA	3:N:220:PHE:CD1	2.46	0.51
2:c:55:GLN:HG3	2:c:59:LYS:HE2	1.91	0.51
3:N:146:GLY:HA2	3:N:149:VAL:HG12	1.93	0.51
3:N:233:ALA:O	3:N:234:ILE:C	2.53	0.51
3:G:680:ARG:HB2	4:I:253:ASP:HB3	1.93	0.51
4:I:451:THR:HA	4:I:454:MET:HE3	1.90	0.51
4:J:643:ASN:O	4:J:646:ILE:HG13	2.11	0.51
4:K:262:THR:HG22	4:K:263:ILE:HG13	1.92	0.51
4:K:894:ILE:O	4:K:898:ILE:HG13	2.10	0.51
2:d:113:VAL:HG11	3:Q:176:ILE:HD12	1.92	0.51
2:h:35:ARG:NH1	2:h:66:GLU:O	2.41	0.51
4:I:604:ARG:NH1	4:I:950:ASP:OD2	2.43	0.51
4:I:863:GLN:O	4:I:866:VAL:HG22	2.10	0.51
4:K:405:SER:OG	4:K:423:ASN:OD1	2.24	0.51
1:A:46:THR:HG23	3:M:445:MET:HE3	1.91	0.51
2:g:65:ALA:O	2:g:68:THR:OG1	2.28	0.51
3:P:160:ASP:OD1	3:P:161:VAL:N	2.42	0.51
3:P:262:MET:HE3	3:P:291:MET:HE2	1.92	0.51
3:H:510:ASP:OD1	3:H:511:GLN:N	2.43	0.51
4:I:894:ILE:O	4:I:898:ILE:HG13	2.11	0.51
4:I:1099:GLY:O	4:I:1103:LYS:HG2	2.11	0.51
4:K:476:VAL:HA	4:K:482:VAL:HA	1.91	0.51
4:L:898:ILE:HG22	4:L:1144:MET:HE1	1.92	0.51
4:L:1163:PRO:HB2	4:L:1166:LYS:NZ	2.24	0.51
3:M:47:ARG:HA	3:M:50:LYS:HE2	1.93	0.51
3:M:380:SER:HA	3:M:383:LYS:HE2	1.93	0.51
3:O:661:LEU:HD13	3:O:672:LEU:HD11	1.92	0.51
3:Q:112:ALA:HB2	3:Q:150:TYR:CD2	2.45	0.51
4:K:432:LEU:HD22	4:K:897:MET:HE1	1.93	0.51
4:L:143:LEU:HD21	4:L:154:LEU:HD23	1.91	0.51
3:O:132:ARG:HD2	3:O:179:TYR:CD1	2.46	0.51
3:R:321:ASP:OD1	3:R:321:ASP:N	2.44	0.51

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
4:K:538:ARG:NH1	4:L:783:GLY:HA2	2.26	0.51
4:K:827:GLU:O	4:K:831:LEU:HG	2.10	0.51
4:K:897:MET:HA	4:K:900:THR:HG22	1.92	0.51
4:L:431:ARG:NH2	4:L:484:SER:HB3	2.25	0.51
2:T:52:LEU:HB2	2:T:57:ARG:HD3	1.93	0.51
3:M:420:ASP:OD1	3:M:420:ASP:N	2.41	0.51
3:N:608:PHE:CG	3:N:663:MET:HE1	2.46	0.51
3:N:680:ARG:NH1	4:J:423:ASN:HD22	2.08	0.51
3:O:563:PRO:HG2	3:O:564:GLU:OE2	2.11	0.51
3:Q:374:LEU:HD13	3:R:296:ARG:HA	1.93	0.51
3:Q:646:ARG:HG3	3:Q:661:LEU:HD22	1.92	0.51
3:R:101:MET:O	3:R:105:ARG:HG3	2.10	0.51
3:H:587:SER:O	3:H:589:ASN:N	2.42	0.51
4:I:553:ALA:O	4:I:557:ASP:HB2	2.11	0.51
4:I:897:MET:HA	4:I:900:THR:HG22	1.93	0.51
4:I:1112:THR:HG23	4:I:1138:LEU:HD13	1.93	0.51
4:L:1112:THR:HG23	4:L:1138:LEU:HD13	1.93	0.51
3:N:130:ARG:NH2	3:O:152:GLU:OE2	2.44	0.51
3:P:699:LEU:HB3	3:Q:372:LYS:HE2	1.92	0.51
4:K:814:MET:HE1	4:K:820:THR:HG22	1.91	0.51
2:S:39:MET:HE1	3:N:154:PHE:HE1	1.75	0.50
2:c:26:MET:O	2:c:30:GLN:HG2	2.11	0.50
2:c:52:LEU:O	2:c:57:ARG:NH1	2.42	0.50
3:M:103:ALA:O	3:M:107:LYS:HG3	2.10	0.50
3:M:526:ILE:HA	3:N:480:ASN:ND2	2.26	0.50
3:O:453:ASP:OD2	3:O:459:PHE:HB2	2.10	0.50
3:Q:197:ASN:O	3:Q:201:GLU:HG2	2.11	0.50
3:H:238:ALA:O	3:H:242:GLN:HG2	2.11	0.50
4:I:431:ARG:NH2	4:I:484:SER:HB3	2.27	0.50
4:K:100:ASP:OD2	4:K:123:TYR:OH	2.23	0.50
3:M:233:ALA:O	3:M:234:ILE:C	2.53	0.50
3:H:565:ILE:HD12	3:H:597:MET:HE2	1.92	0.50
4:J:581:MET:HE1	4:J:595:ARG:HG3	1.93	0.50
4:J:775:ASP:OD1	4:J:789:ASN:ND2	2.44	0.50
4:J:1164:LYS:HA	4:J:1167:ARG:HD2	1.93	0.50
4:L:473:GLY:HA3	4:L:485:ALA:HB2	1.93	0.50
1:E:45:GLY:HA2	3:P:448:LYS:NZ	2.27	0.50
3:N:118:ASP:OD2	3:N:174:ARG:HD2	2.11	0.50
3:Q:139:ARG:O	3:Q:143:LEU:HG	2.12	0.50
3:G:527:ASP:HB3	3:G:530:VAL:HG22	1.93	0.50
3:H:391:PHE:O	3:H:395:ILE:HG12	2.11	0.50

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
4:I:1030:SER:HB2	4:I:1074:ARG:NH1	2.26	0.50
4:I:1098:PHE:O	4:I:1103:LYS:NZ	2.44	0.50
4:J:34:ALA:HB2	4:J:54:MET:HE1	1.93	0.50
3:M:324:THR:O	3:M:328:MET:HG3	2.11	0.50
3:Q:121:MET:HG3	3:Q:125:LYS:HE2	1.92	0.50
3:Q:252:SER:HB2	3:Q:293:THR:HG23	1.93	0.50
3:G:303:ALA:O	3:G:307:GLU:HG2	2.12	0.50
4:K:679:ILE:HG12	4:K:683:GLY:O	2.11	0.50
2:d:71:ASN:ND2	3:Q:122:GLN:HB2	2.26	0.50
2:f:81:ILE:HA	3:R:220:PHE:CD1	2.47	0.50
2:h:73:GLN:HE21	3:H:204:GLY:HA2	1.77	0.50
3:P:500:ASP:OD1	3:P:632:PRO:HB3	2.12	0.50
3:P:556:ALA:HB1	3:P:594:MET:HE3	1.93	0.50
3:R:467:VAL:HG12	3:R:511:GLN:HG3	1.92	0.50
4:I:595:ARG:HG2	4:I:1192:GLY:HA3	1.94	0.50
4:I:1278:GLN:O	4:I:1282:THR:HG22	2.12	0.50
4:K:1084:MET:HG2	4:K:1098:PHE:CZ	2.46	0.50
2:g:53:SER:O	2:g:57:ARG:HG3	2.12	0.50
3:P:125:LYS:NZ	3:P:178:LEU:O	2.40	0.50
4:K:456:PHE:HA	4:K:466:GLU:HG2	1.93	0.50
4:K:712:MET:HE2	4:K:771:LEU:HD12	1.92	0.50
4:K:988:LEU:HD22	4:K:1131:ASN:HD22	1.75	0.50
4:L:964:ALA:O	4:L:968:ILE:HG12	2.11	0.50
3:N:684:ASP:OD1	3:N:685:LYS:N	2.44	0.50
3:P:136:GLU:OE1	3:Q:56:TYR:OH	2.30	0.50
3:P:233:ALA:O	3:P:234:ILE:C	2.53	0.50
3:R:67:ALA:O	3:R:70:ARG:HG2	2.11	0.50
3:R:388:ASP:OD1	3:R:428:LYS:NZ	2.45	0.50
3:G:117:ASP:OD2	3:G:174:ARG:NH1	2.33	0.50
3:G:646:ARG:HG3	3:G:661:LEU:HD22	1.93	0.50
3:H:66:LEU:HG	3:H:70:ARG:NH1	2.26	0.50
4:J:985:TRP:O	4:J:993:THR:OG1	2.25	0.50
4:L:637:LYS:O	4:L:641:MET:HE3	2.11	0.50
2:T:67:LEU:HD12	3:N:196:MET:SD	2.52	0.50
3:M:563:PRO:HG2	3:M:564:GLU:OE1	2.12	0.50
3:R:191:GLN:O	3:R:195:ILE:HG12	2.11	0.50
3:H:118:ASP:OD2	3:H:174:ARG:HD2	2.12	0.50
4:I:1277:GLU:HG2	4:J:974:TYR:CD1	2.47	0.50
4:J:1012:THR:HG22	4:J:1040:LYS:HD3	1.94	0.50
4:L:11:ASP:O	4:L:1024:GLY:HA3	2.12	0.50
4:L:825:LYS:O	4:L:829:LEU:HG	2.12	0.50

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:M:131:THR:HG22	3:M:134:GLU:HG3	1.94	0.50
3:M:349:ARG:O	3:M:353:ILE:HG12	2.12	0.50
3:R:125:LYS:NZ	3:R:181:ALA:HB3	2.27	0.50
3:H:583:VAL:HG11	3:H:596:GLN:HG3	1.94	0.50
3:H:608:PHE:CG	3:H:663:MET:HE1	2.47	0.50
4:I:643:ASN:O	4:I:646:ILE:HG13	2.12	0.50
4:K:273:TRP:HB2	4:K:278:LEU:HG	1.94	0.50
4:L:1278:GLN:O	4:L:1282:THR:HG22	2.11	0.50
1:D:40:GLU:OE1	3:O:373:ALA:HB2	2.11	0.49
3:P:503:LEU:O	3:P:507:LEU:HG	2.12	0.49
4:K:1112:THR:HG23	4:K:1138:LEU:HD13	1.94	0.49
4:L:8:VAL:HB	4:L:9:PRO:HD3	1.94	0.49
2:e:128:GLN:HB3	3:G:82:GLN:OE1	2.12	0.49
3:M:346:THR:HG23	3:M:349:ARG:H	1.77	0.49
3:N:112:ALA:HB2	3:N:150:TYR:HD2	1.76	0.49
3:N:517:THR:HG22	3:N:521:MET:HE2	1.94	0.49
3:O:132:ARG:HD2	3:O:179:TYR:HD1	1.77	0.49
3:O:401:SER:OG	3:O:406:ASP:OD2	2.26	0.49
3:R:262:MET:HE3	3:R:291:MET:SD	2.51	0.49
3:R:271:LEU:HG	3:R:272:ASN:H	1.77	0.49
3:R:574:GLU:O	3:R:578:LYS:HG3	2.12	0.49
3:R:626:MET:O	3:R:638:GLY:HA3	2.12	0.49
3:H:404:PHE:HA	3:H:407:MET:HG2	1.94	0.49
4:I:561:ARG:O	4:I:565:THR:HG23	2.12	0.49
4:I:968:ILE:CG2	4:L:1237:VAL:HG22	2.43	0.49
4:K:898:ILE:HG22	4:K:1144:MET:HE1	1.94	0.49
4:L:273:TRP:HB2	4:L:278:LEU:HG	1.94	0.49
2:S:57:ARG:HH21	3:M:242:GLN:NE2	2.10	0.49
2:d:40:GLU:O	2:d:44:GLN:HG2	2.12	0.49
3:M:151:ALA:HB2	3:M:163:TYR:CZ	2.47	0.49
3:N:132:ARG:O	3:N:136:GLU:HG2	2.11	0.49
3:Q:556:ALA:O	3:Q:560:SER:OG	2.26	0.49
3:R:272:ASN:OD1	3:R:273:GLY:N	2.45	0.49
3:G:232:GLY:HA3	3:H:100:ALA:HB2	1.94	0.49
3:H:112:ALA:HB2	3:H:150:TYR:CD2	2.47	0.49
3:H:160:ASP:OD1	3:H:161:VAL:N	2.45	0.49
4:I:1013:LEU:HD23	4:I:1049:ARG:HB2	1.94	0.49
3:N:626:MET:O	3:N:638:GLY:HA3	2.12	0.49
3:P:72:ASN:HB3	3:P:76:ARG:HH21	1.78	0.49
3:H:680:ARG:NH1	4:I:423:ASN:HD22	2.10	0.49
4:J:1197:MET:SD	4:J:1287:SER:HB2	2.52	0.49

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
4:K:633:HIS:CD2	4:K:800:MET:HE2	2.46	0.49
4:K:866:VAL:HG13	4:K:1201:VAL:HG23	1.95	0.49
4:K:904:ARG:HB3	4:K:1311:VAL:HG13	1.94	0.49
4:K:940:VAL:O	4:K:944:ILE:HG12	2.11	0.49
1:b:40:GLU:OE1	3:M:373:ALA:HB2	2.12	0.49
2:e:63:ALA:O	2:e:67:LEU:HD23	2.12	0.49
3:M:454:SER:C	3:M:456:ASP:H	2.20	0.49
3:O:526:ILE:HA	3:P:480:ASN:ND2	2.27	0.49
3:O:564:GLU:H	3:O:564:GLU:CD	2.19	0.49
3:P:130:ARG:HB3	3:Q:156:ILE:HD11	1.94	0.49
3:Q:336:LEU:HD12	3:Q:352:LEU:HD22	1.93	0.49
3:R:643:GLU:O	3:R:647:LYS:HG2	2.12	0.49
3:H:645:ALA:O	3:H:649:ILE:HG12	2.12	0.49
3:H:674:ASP:OD1	3:H:674:ASP:N	2.45	0.49
4:I:140:ARG:NH2	4:I:157:PHE:O	2.37	0.49
4:J:612:ARG:HD3	4:J:775:ASP:OD2	2.12	0.49
4:K:431:ARG:NH2	4:K:484:SER:HB3	2.27	0.49
4:K:1278:GLN:O	4:K:1282:THR:HG22	2.12	0.49
4:L:633:HIS:O	4:L:637:LYS:HG2	2.13	0.49
4:L:809:GLY:HA3	4:L:856:THR:HA	1.93	0.49
3:M:570:ALA:O	3:M:573:ARG:HB2	2.12	0.49
3:O:47:ARG:HA	3:O:50:LYS:HE2	1.94	0.49
3:O:550:ASP:O	3:O:554:GLU:HG2	2.12	0.49
3:R:589:ASN:HA	3:G:539:VAL:HG22	1.92	0.49
3:R:602:LYS:HE3	3:G:468:THR:HG21	1.95	0.49
3:G:639:ARG:HH12	3:H:418:HIS:HB3	1.77	0.49
4:I:416:LYS:HZ2	4:I:418:GLU:N	2.07	0.49
4:I:972:LEU:HD13	4:L:1277:GLU:HA	1.93	0.49
4:J:673:HIS:ND1	4:J:946:PRO:HA	2.28	0.49
4:L:17:GLN:HG3	4:L:1086:ARG:HG2	1.94	0.49
2:c:77:ALA:HB3	3:O:211:MET:HE3	1.95	0.49
2:h:72:MET:HE1	3:M:177:SER:OG	2.13	0.49
3:M:95:GLN:HA	3:M:101:MET:HE2	1.94	0.49
3:M:368:LYS:HZ2	3:H:704:ARG:HD3	1.78	0.49
3:M:453:ASP:OD2	3:M:459:PHE:HB2	2.13	0.49
3:N:112:ALA:HB2	3:N:150:TYR:CD2	2.47	0.49
3:H:205:VAL:HG12	3:H:206:LEU:HD23	1.94	0.49
4:I:101:GLU:HG3	4:I:147:LYS:H	1.78	0.49
4:I:593:MET:HE3	4:I:598:THR:HG22	1.94	0.49
4:J:866:VAL:HG21	4:J:1197:MET:HE3	1.94	0.49
4:J:1112:THR:HG23	4:J:1138:LEU:HD13	1.95	0.49

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
4:L:712:MET:HE2	4:L:771:LEU:HD12	1.95	0.49
4:L:1021:GLU:HG2	4:L:1036:MET:HE3	1.95	0.49
2:c:39:MET:HE1	3:P:154:PHE:HE1	1.77	0.49
3:O:383:LYS:HA	3:O:386:VAL:HG12	1.93	0.49
3:R:224:TYR:OH	3:G:107:LYS:NZ	2.46	0.49
3:G:139:ARG:O	3:G:143:LEU:HG	2.13	0.49
3:G:270:THR:HG23	3:G:275:THR:HG22	1.94	0.49
3:G:380:SER:HA	3:G:383:LYS:HG2	1.93	0.49
4:I:54:MET:HB2	4:I:56:PHE:CE1	2.48	0.49
4:I:416:LYS:NZ	4:I:418:GLU:OE1	2.46	0.49
4:I:556:SER:O	4:I:560:GLU:HG3	2.13	0.49
4:J:584:PRO:HA	4:J:588:THR:HG23	1.94	0.49
1:E:39:ASP:OD2	3:Q:323:ARG:NH2	2.46	0.49
3:N:238:ALA:O	3:N:242:GLN:HG2	2.13	0.49
3:O:645:ALA:HA	4:K:227:PHE:HE2	1.78	0.49
3:P:121:MET:O	3:P:125:LYS:HG3	2.13	0.49
3:P:674:ASP:OD1	3:P:674:ASP:N	2.44	0.49
4:I:1004:MET:HE1	4:I:1020:TRP:NE1	2.28	0.49
4:J:594:SER:O	4:J:598:THR:HG23	2.12	0.49
2:e:62:GLU:O	2:e:66:GLU:HG3	2.13	0.49
2:f:128:GLN:OE1	2:f:128:GLN:N	2.44	0.49
3:G:383:LYS:HB2	3:G:416:PHE:CE1	2.47	0.49
4:K:561:ARG:HH22	4:L:779:THR:HG21	1.77	0.49
4:K:633:HIS:O	4:K:637:LYS:HG2	2.13	0.49
4:L:584:PRO:HA	4:L:588:THR:HG23	1.95	0.49
4:L:1171:LEU:HD22	4:L:1175:LEU:HD12	1.95	0.49
3:M:374:LEU:HD13	3:N:296:ARG:HA	1.94	0.48
3:M:483:LEU:HD12	3:M:484:PRO:HD2	1.93	0.48
3:M:673:MET:HG2	3:M:674:ASP:O	2.13	0.48
3:N:47:ARG:HA	3:N:50:LYS:HE2	1.94	0.48
3:N:394:ARG:CG	3:N:452:ALA:HB1	2.43	0.48
3:O:568:MET:SD	3:O:573:ARG:HA	2.53	0.48
3:Q:563:PRO:HG2	3:Q:564:GLU:OE2	2.13	0.48
3:Q:611:ASP:N	3:Q:658:ASN:HD21	2.12	0.48
3:R:587:SER:O	3:R:589:ASN:N	2.43	0.48
3:R:700:GLU:HG3	3:G:372:LYS:HB2	1.95	0.48
3:H:455:LYS:HG2	3:H:455:LYS:O	2.12	0.48
3:H:541:ARG:HD2	3:H:546:ARG:HE	1.78	0.48
4:I:54:MET:HE2	4:I:54:MET:HA	1.95	0.48
4:L:442:ASN:OD1	4:L:442:ASN:O	2.31	0.48
4:L:904:ARG:HB3	4:L:1311:VAL:HG13	1.95	0.48

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
4:L:1084:MET:HE1	4:L:1105:VAL:HG13	1.95	0.48
2:h:54:LEU:HD11	3:G:319:GLN:OE1	2.13	0.48
2:h:69:SER:HB3	3:M:115:LEU:HD23	1.95	0.48
3:G:413:THR:HG21	3:G:416:PHE:CE2	2.48	0.48
3:H:453:ASP:HB2	3:H:460:ARG:HD3	1.94	0.48
4:K:414:ARG:HG2	4:K:414:ARG:HH11	1.78	0.48
4:K:497:PHE:HZ	4:K:1187:ARG:HH12	1.60	0.48
1:D:46:THR:HG23	3:N:448:LYS:HE3	1.95	0.48
2:S:37:GLN:HE22	2:S:41:ILE:HD11	1.78	0.48
3:M:500:ASP:OD2	3:M:503:LEU:HB3	2.13	0.48
3:N:125:LYS:NZ	3:N:182:HIS:HB2	2.28	0.48
3:N:510:ASP:OD1	3:N:511:GLN:N	2.46	0.48
3:O:132:ARG:O	3:O:136:GLU:HG3	2.13	0.48
3:O:147:ALA:HB2	3:O:167:PHE:CZ	2.48	0.48
3:G:116:VAL:O	3:G:120:VAL:HG23	2.13	0.48
3:H:541:ARG:HB3	3:H:546:ARG:HE	1.77	0.48
4:J:442:ASN:O	4:J:442:ASN:OD1	2.31	0.48
4:J:832:LYS:HE3	4:J:846:HIS:CE1	2.48	0.48
4:L:795:ASP:O	4:L:796:MET:C	2.56	0.48
2:f:82:ARG:NH2	2:f:103:GLU:OE2	2.47	0.48
2:g:60:LEU:HD12	3:G:196:MET:HE1	1.96	0.48
3:P:146:GLY:HA2	3:P:149:VAL:HG12	1.94	0.48
3:P:455:LYS:HG2	3:P:455:LYS:O	2.12	0.48
3:Q:323:ARG:NH1	3:R:293:THR:HG21	2.29	0.48
3:G:211:MET:O	3:G:217:SER:HB3	2.12	0.48
3:H:205:VAL:HG21	3:H:220:PHE:CZ	2.48	0.48
3:H:283:GLY:O	3:H:287:TRP:N	2.37	0.48
4:J:712:MET:HE3	4:J:712:MET:HB3	1.73	0.48
4:J:894:ILE:O	4:J:898:ILE:HG13	2.14	0.48
4:K:637:LYS:O	4:K:641:MET:HE3	2.14	0.48
4:K:914:ARG:O	4:K:919:LYS:HG3	2.13	0.48
4:K:1241:MET:O	4:K:1245:LEU:HG	2.12	0.48
3:N:645:ALA:O	3:N:649:ILE:HG12	2.13	0.48
3:O:380:SER:HA	3:O:383:LYS:HE2	1.95	0.48
3:P:283:GLY:O	3:P:287:TRP:N	2.42	0.48
4:I:379:SER:H	4:I:1282:THR:HB	1.79	0.48
4:J:140:ARG:HA	4:J:143:LEU:HG	1.95	0.48
4:K:553:ALA:O	4:K:557:ASP:HB2	2.13	0.48
4:L:398:GLY:HA3	4:L:429:MET:SD	2.53	0.48
4:L:806:ARG:NH1	4:L:856:THR:OG1	2.46	0.48
2:e:72:MET:SD	3:R:118:ASP:HB3	2.54	0.48

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:g:55:GLN:HG3	2:g:59:LYS:HE2	1.95	0.48
2:h:81:ILE:HA	3:H:220:PHE:CD1	2.49	0.48
3:M:172:THR:O	3:M:176:ILE:HG12	2.13	0.48
3:O:251:ALA:O	3:O:257:GLY:HA3	2.14	0.48
3:P:432:ILE:HG23	3:P:445:MET:HE3	1.96	0.48
3:G:382:ASN:ND2	3:G:412:ASN:HB3	2.29	0.48
4:J:273:TRP:HB2	4:J:278:LEU:HG	1.94	0.48
4:L:266:THR:HG23	4:L:269:ASN:H	1.78	0.48
3:M:197:ASN:O	3:M:201:GLU:HG2	2.13	0.48
3:M:518:MET:HA	3:M:521:MET:HG2	1.96	0.48
3:M:660:GLN:HA	3:M:675:THR:HG23	1.95	0.48
3:P:669:SER:HB3	3:P:682:ARG:NH1	2.28	0.48
3:Q:380:SER:HA	3:Q:383:LYS:HG2	1.95	0.48
3:R:146:GLY:HA2	3:R:149:VAL:HG12	1.95	0.48
3:G:132:ARG:O	3:G:136:GLU:HG3	2.13	0.48
4:I:681:ARG:HH22	4:L:1243:SER:HA	1.77	0.48
4:J:380:VAL:CG2	4:J:575:LYS:HG3	2.43	0.48
4:J:679:ILE:O	4:J:683:GLY:HA2	2.12	0.48
3:M:211:MET:O	3:M:214:ARG:HG2	2.14	0.48
3:M:480:ASN:HD21	3:H:526:ILE:HA	1.77	0.48
3:N:82:GLN:HA	3:N:85:GLU:HG3	1.96	0.48
3:P:238:ALA:O	3:P:242:GLN:HG2	2.14	0.48
3:P:252:SER:HB2	3:P:293:THR:HG23	1.96	0.48
3:P:349:ARG:O	3:P:353:ILE:HD12	2.13	0.48
3:Q:70:ARG:O	3:Q:74:ILE:HG12	2.14	0.48
3:G:200:VAL:HG11	3:H:111:ASN:ND2	2.24	0.48
3:H:170:ASP:OD1	3:H:173:GLU:HG3	2.13	0.48
4:I:975:SER:HB3	4:I:978:GLU:OE2	2.14	0.48
4:J:414:ARG:HG2	4:J:414:ARG:HH11	1.79	0.48
4:K:11:ASP:CG	4:K:1022:LYS:HB3	2.38	0.48
4:K:112:GLU:HA	4:K:116:GLY:HA3	1.94	0.48
4:K:355:LEU:HD21	4:K:433:GLU:HG3	1.95	0.48
4:K:594:SER:O	4:K:598:THR:HG23	2.14	0.48
4:L:1314:ARG:HG3	4:L:1315:GLU:N	2.19	0.48
3:M:487:THR:HG23	3:M:490:MET:HE3	1.96	0.48
3:N:587:SER:O	3:N:589:ASN:N	2.46	0.48
3:Q:413:THR:HG21	3:Q:416:PHE:CE1	2.48	0.48
3:R:674:ASP:OD1	3:R:674:ASP:N	2.45	0.48
3:H:497:ARG:HE	3:H:518:MET:HE3	1.77	0.48
4:I:11:ASP:CG	4:I:1022:LYS:HB3	2.39	0.48
4:J:1019:ARG:HA	4:J:1022:LYS:NZ	2.29	0.48

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
4:L:386:ASP:N	4:L:386:ASP:OD1	2.45	0.48
4:L:1090:VAL:HG22	4:L:1109:LYS:HD3	1.95	0.48
2:h:119:ARG:NH1	3:N:90:GLY:O	2.27	0.48
3:M:160:ASP:O	3:M:164:GLN:HG2	2.13	0.48
3:N:657:THR:HG22	3:N:659:LYS:H	1.79	0.48
3:R:132:ARG:HD2	3:R:179:TYR:CD2	2.49	0.48
3:G:568:MET:SD	3:G:573:ARG:HA	2.54	0.48
3:H:641:ILE:HD13	3:H:691:VAL:HG23	1.96	0.48
4:I:581:MET:HE1	4:I:595:ARG:HG3	1.96	0.48
4:J:513:GLY:O	4:J:517:ILE:HG12	2.14	0.48
4:J:520:LYS:HD2	4:J:847:ALA:HB2	1.96	0.48
4:L:806:ARG:HA	4:L:806:ARG:HD2	1.73	0.48
1:B:42:ASP:OD1	1:B:43:THR:N	2.46	0.47
3:N:599:LYS:HE3	3:O:472:GLN:CD	2.39	0.47
3:P:50:LYS:NZ	3:P:159:GLU:OE2	2.48	0.47
3:Q:453:ASP:OD2	3:Q:459:PHE:HB2	2.13	0.47
3:H:417:LYS:HZ3	3:H:419:SER:HB2	1.79	0.47
4:J:1277:GLU:HA	4:K:972:LEU:HD13	1.95	0.47
4:K:743:GLN:HG3	4:K:744:PHE:CD1	2.49	0.47
2:e:42:MET:HE1	3:R:154:PHE:CE1	2.49	0.47
3:M:382:ASN:ND2	3:M:412:ASN:HB3	2.29	0.47
3:M:613:VAL:HG23	3:M:614:ASP:H	1.78	0.47
3:N:374:LEU:HD13	3:O:296:ARG:HA	1.96	0.47
3:O:116:VAL:HG11	3:O:143:LEU:HA	1.95	0.47
3:P:467:VAL:HG22	3:P:493:LEU:HD13	1.96	0.47
3:Q:132:ARG:NH2	3:Q:183:ASP:OD1	2.41	0.47
3:Q:701:GLU:O	3:Q:705:GLU:HG2	2.13	0.47
3:R:250:ASP:O	3:R:254:ARG:HG2	2.14	0.47
3:G:483:LEU:HD11	3:G:490:MET:HE1	1.95	0.47
4:I:333:ILE:HG22	4:I:433:GLU:OE2	2.12	0.47
3:M:303:ALA:O	3:M:307:GLU:HG2	2.14	0.47
3:N:467:VAL:HG22	3:N:493:LEU:HD13	1.95	0.47
3:N:497:ARG:NH2	3:N:519:ASP:OD1	2.44	0.47
3:N:631:ASP:OD1	3:N:632:PRO:HD2	2.14	0.47
3:R:98:PRO:O	3:R:102:GLU:HG3	2.14	0.47
3:R:697:LYS:O	3:R:701:GLU:HG2	2.14	0.47
3:G:160:ASP:O	3:G:164:GLN:HG2	2.14	0.47
4:I:633:HIS:O	4:I:637:LYS:HG2	2.15	0.47
4:I:679:ILE:HG12	4:I:683:GLY:O	2.13	0.47
2:S:35:ARG:NH1	3:N:116:VAL:HG13	2.30	0.47
3:M:517:THR:OG1	3:M:586:ARG:NH2	2.44	0.47

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:O:172:THR:O	3:O:176:ILE:HG12	2.15	0.47
3:H:197:ASN:O	3:H:201:GLU:HG2	2.14	0.47
3:H:522:ASP:OD1	3:H:523:LYS:HG2	2.14	0.47
4:I:376:LEU:HG	4:I:378:THR:HG23	1.95	0.47
4:I:974:TYR:CD1	4:L:1277:GLU:HG2	2.49	0.47
4:J:777:SER:HB3	4:J:787:SER:HB3	1.95	0.47
4:J:1030:SER:HB2	4:J:1074:ARG:NH1	2.29	0.47
4:K:593:MET:HB3	4:K:597:GLU:HB3	1.95	0.47
4:K:1237:VAL:HG22	4:L:968:ILE:CG2	2.43	0.47
3:M:93:LEU:HD21	3:H:183:ASP:OD1	2.13	0.47
3:N:79:THR:H	3:N:82:GLN:NE2	2.10	0.47
3:O:394:ARG:HD3	3:O:452:ALA:O	2.14	0.47
3:G:191:GLN:O	3:G:195:ILE:HG13	2.15	0.47
4:J:363:GLY:O	4:J:443:SER:HB3	2.13	0.47
4:J:550:LYS:HD3	4:J:554:THR:HG22	1.97	0.47
4:J:1084:MET:HG2	4:J:1098:PHE:CZ	2.49	0.47
4:J:1278:GLN:O	4:J:1282:THR:HG22	2.14	0.47
4:K:990:ASN:O	4:K:994:ASN:ND2	2.48	0.47
3:M:283:GLY:O	3:M:287:TRP:N	2.41	0.47
3:N:394:ARG:HG3	3:N:452:ALA:HB1	1.97	0.47
3:O:70:ARG:O	3:O:74:ILE:HG12	2.15	0.47
3:R:576:ALA:HB1	3:R:597:MET:HE3	1.95	0.47
3:H:463:ILE:O	3:H:467:VAL:HG23	2.15	0.47
4:I:806:ARG:HA	4:I:806:ARG:HD2	1.74	0.47
4:I:1149:GLY:O	4:I:1153:MET:HG3	2.15	0.47
4:J:630:ILE:HD13	4:J:796:MET:HE3	1.96	0.47
4:K:866:VAL:HG11	4:K:1197:MET:CE	2.45	0.47
4:K:988:LEU:CD2	4:K:1131:ASN:HD22	2.28	0.47
4:L:777:SER:HB3	4:L:787:SER:HB2	1.96	0.47
4:L:779:THR:HG22	4:L:785:GLN:HG2	1.95	0.47
2:f:60:LEU:HD12	3:R:196:MET:HE1	1.95	0.47
3:M:238:ALA:O	3:M:242:GLN:HG2	2.14	0.47
3:M:655:TRP:CZ3	3:M:656:ILE:HG12	2.50	0.47
3:N:125:LYS:NZ	3:N:135:MET:HG3	2.29	0.47
3:N:199:ARG:HH11	3:N:243:LEU:HD13	1.78	0.47
3:N:641:ILE:HD13	3:N:691:VAL:HG23	1.96	0.47
3:O:386:VAL:HG11	3:O:416:PHE:HZ	1.79	0.47
3:O:541:ARG:HH22	3:O:544:GLU:HB3	1.79	0.47
3:P:641:ILE:HD13	3:P:691:VAL:HG23	1.96	0.47
3:R:551:LYS:HZ3	3:G:543:LYS:CG	2.28	0.47
3:R:598:THR:HG22	3:R:602:LYS:NZ	2.29	0.47

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:G:547:PHE:O	3:G:551:LYS:HG2	2.15	0.47
3:H:612:ASP:OD1	3:H:615:GLY:HA2	2.14	0.47
4:I:383:GLY:O	4:I:384:ASP:HB2	2.15	0.47
4:I:661:HIS:HB3	4:I:769:ARG:HH21	1.80	0.47
4:J:8:VAL:HB	4:J:9:PRO:HD3	1.96	0.47
4:K:609:ALA:HB1	4:K:616:GLN:HG2	1.97	0.47
4:K:724:PRO:HA	4:K:727:VAL:HG12	1.97	0.47
2:f:61:GLU:HA	3:R:199:ARG:NH2	2.30	0.47
2:h:52:LEU:HB3	2:h:56:ALA:HB3	1.97	0.47
3:N:78:LEU:HA	3:N:82:GLN:HE21	1.80	0.47
3:N:493:LEU:HA	3:N:496:ILE:HG22	1.96	0.47
3:O:102:GLU:HG3	3:O:162:ASP:CG	2.40	0.47
3:Q:628:VAL:O	3:Q:629:ASN:OD1	2.33	0.47
3:R:115:LEU:O	3:R:119:ASP:HB2	2.15	0.47
3:R:323:ARG:HH11	3:G:290:LEU:HD23	1.79	0.47
3:G:203:ASN:O	3:G:207:GLN:HB2	2.14	0.47
3:G:517:THR:HA	3:G:520:MET:HE2	1.95	0.47
3:G:543:LYS:HA	3:G:546:ARG:HB2	1.97	0.47
4:I:13:ASP:OD1	4:I:31:ARG:NH1	2.47	0.47
4:I:1277:GLU:HA	4:J:972:LEU:HD13	1.96	0.47
4:L:63:ALA:O	4:L:64:LEU:HB2	2.15	0.47
2:T:53:SER:O	2:T:57:ARG:HG2	2.14	0.47
3:M:261:LEU:HA	3:M:264:VAL:HG12	1.97	0.47
3:N:238:ALA:O	3:N:241:THR:OG1	2.33	0.47
3:N:526:ILE:HA	3:O:480:ASN:HD21	1.80	0.47
3:Q:255:ALA:HB2	3:Q:301:THR:HG21	1.97	0.47
3:Q:607:THR:HG23	3:Q:620:VAL:HG22	1.96	0.47
3:R:526:ILE:HA	3:G:480:ASN:HD21	1.80	0.47
4:I:679:ILE:O	4:I:683:GLY:HA2	2.13	0.47
4:I:964:ALA:O	4:I:968:ILE:HG12	2.15	0.47
4:J:806:ARG:NH1	4:J:856:THR:OG1	2.48	0.47
2:S:26:MET:HE1	2:h:6:ALA:N	2.30	0.47
2:e:39:MET:HE3	2:e:67:LEU:HD21	1.98	0.47
3:P:72:ASN:HA	3:P:75:ILE:HG22	1.95	0.47
3:P:437:ILE:CD1	3:P:445:MET:HE2	2.45	0.47
3:Q:303:ALA:O	3:Q:307:GLU:HG2	2.15	0.47
3:R:269:VAL:HG11	3:R:281:LEU:HD22	1.97	0.47
3:H:657:THR:HB	3:H:660:GLN:HG3	1.97	0.47
4:I:408:VAL:HB	4:I:1313:LEU:HD21	1.96	0.47
4:I:616:GLN:HG2	4:I:627:MET:CE	2.45	0.47
4:K:442:ASN:O	4:K:442:ASN:OD1	2.32	0.47

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
4:K:1084:MET:HE2	4:K:1084:MET:HB3	1.61	0.47
3:M:87:LEU:HB2	3:M:92:LEU:HD23	1.96	0.46
3:M:139:ARG:NH2	3:M:175:ASN:OD1	2.48	0.46
3:M:147:ALA:HB2	3:M:167:PHE:CZ	2.50	0.46
3:M:336:LEU:HD12	3:M:352:LEU:HD22	1.98	0.46
3:M:413:THR:HG21	3:M:416:PHE:CE2	2.49	0.46
3:M:520:MET:HB3	3:M:526:ILE:CD1	2.43	0.46
3:N:383:LYS:O	3:N:387:ILE:HG13	2.14	0.46
3:P:276:THR:OG1	3:P:280:GLU:OE1	2.33	0.46
3:R:238:ALA:O	3:R:242:GLN:HG2	2.15	0.46
3:G:151:ALA:HB2	3:G:163:TYR:CZ	2.50	0.46
3:H:66:LEU:HG	3:H:70:ARG:HH12	1.79	0.46
3:H:626:MET:O	3:H:638:GLY:HA3	2.16	0.46
4:I:572:ASP:OD2	4:I:1261:THR:HG22	2.15	0.46
4:J:496:GLU:OE1	4:J:1173:ARG:HG2	2.15	0.46
4:K:880:PHE:HB2	4:L:974:TYR:CZ	2.50	0.46
4:L:630:ILE:HD13	4:L:796:MET:HE3	1.97	0.46
2:T:39:MET:HE2	2:T:67:LEU:HD21	1.98	0.46
3:O:570:ALA:O	3:O:573:ARG:HB2	2.15	0.46
3:P:121:MET:HE1	3:P:138:TYR:CD2	2.50	0.46
3:Q:473:GLU:OE2	3:Q:487:THR:HA	2.16	0.46
3:R:584:LYS:HD3	3:R:590:GLU:CG	2.46	0.46
3:R:641:ILE:HD13	3:R:691:VAL:HG23	1.96	0.46
3:G:674:ASP:OD1	3:G:674:ASP:N	2.49	0.46
4:I:1171:LEU:HD22	4:I:1175:LEU:HD12	1.98	0.46
4:J:60:THR:HA	4:J:63:ALA:HB3	1.98	0.46
4:L:517:ILE:HG21	4:L:851:THR:OG1	2.16	0.46
4:L:1013:LEU:HD22	4:L:1055:PHE:CE1	2.50	0.46
2:S:72:MET:O	2:S:76:GLN:HG2	2.14	0.46
3:M:375:ASP:OD2	3:M:379:LYS:HE3	2.15	0.46
3:N:674:ASP:OD1	3:N:674:ASP:N	2.44	0.46
3:O:139:ARG:O	3:O:143:LEU:HG	2.15	0.46
3:O:151:ALA:HB2	3:O:163:TYR:CZ	2.51	0.46
3:Q:349:ARG:O	3:Q:353:ILE:HG12	2.14	0.46
3:R:162:ASP:OD1	3:R:165:ARG:NH2	2.49	0.46
3:R:556:ALA:HB1	3:R:594:MET:HE3	1.97	0.46
3:G:117:ASP:OD2	3:G:174:ARG:HD2	2.16	0.46
4:I:724:PRO:HA	4:I:727:VAL:HG12	1.97	0.46
4:J:525:ASP:HA	4:J:530:ARG:HH12	1.80	0.46
4:J:1314:ARG:HG3	4:J:1315:GLU:N	2.23	0.46
4:K:109:ASN:HB2	4:K:138:TYR:CE2	2.49	0.46

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
4:K:551:PHE:HD1	4:L:781:PRO:HD3	1.80	0.46
4:L:592:LYS:HG2	4:L:593:MET:HE2	1.96	0.46
2:S:57:ARG:HH21	3:M:242:GLN:HE22	1.64	0.46
2:d:67:LEU:HD22	3:P:196:MET:SD	2.55	0.46
3:O:276:THR:HB	3:O:280:GLU:HB2	1.97	0.46
3:O:500:ASP:HB2	3:O:632:PRO:HG3	1.97	0.46
3:Q:526:ILE:HA	3:R:480:ASN:ND2	2.28	0.46
3:R:520:MET:HE1	3:R:586:ARG:HE	1.81	0.46
3:G:635:TRP:HZ3	3:G:639:ARG:HD3	1.79	0.46
3:H:236:SER:HB3	3:H:239:GLN:H	1.80	0.46
4:I:439:ARG:NH1	4:I:883:ASN:OD1	2.48	0.46
4:J:712:MET:HE2	4:J:771:LEU:HD12	1.97	0.46
4:K:140:ARG:NH2	4:K:157:PHE:O	2.47	0.46
4:L:496:GLU:HG2	4:L:1179:MET:HE3	1.97	0.46
4:L:1013:LEU:HD22	4:L:1055:PHE:HE1	1.80	0.46
2:S:75:VAL:HG13	3:N:173:GLU:OE1	2.15	0.46
2:d:77:ALA:HB3	3:P:211:MET:HE3	1.98	0.46
2:f:7:ILE:HB	2:f:8:PRO:HD3	1.97	0.46
2:g:24:ALA:O	2:g:27:ILE:HG12	2.16	0.46
2:h:40:GLU:O	2:h:44:GLN:HG2	2.15	0.46
3:N:565:ILE:HD13	3:N:601:LEU:HD12	1.98	0.46
3:N:570:ALA:HA	3:N:573:ARG:HB2	1.97	0.46
3:O:394:ARG:NH2	3:O:400:VAL:O	2.49	0.46
3:P:700:GLU:HG2	3:Q:372:LYS:HD3	1.97	0.46
3:Q:653:ASN:HB2	3:Q:656:ILE:HD12	1.98	0.46
3:G:611:ASP:N	3:G:658:ASN:HD21	2.13	0.46
3:H:206:LEU:HD12	3:H:247:ALA:HB1	1.98	0.46
4:I:11:ASP:O	4:I:1024:GLY:HA3	2.15	0.46
4:I:517:ILE:HG21	4:I:851:THR:OG1	2.15	0.46
4:K:50:PRO:HB3	4:K:57:THR:H	1.81	0.46
4:L:801:PRO:O	4:L:805:ARG:HG3	2.15	0.46
2:S:65:ALA:O	2:S:68:THR:OG1	2.31	0.46
3:N:67:ALA:O	3:N:70:ARG:HG2	2.15	0.46
3:N:332:ILE:O	3:N:335:GLU:HG3	2.16	0.46
4:I:885:TYR:O	4:I:889:GLN:HG3	2.16	0.46
4:J:465:TYR:CD2	4:J:491:PRO:HB2	2.51	0.46
4:J:796:MET:HA	4:J:799:ILE:HB	1.96	0.46
4:J:1098:PHE:HE2	4:J:1106:MET:HE2	1.80	0.46
4:K:630:ILE:HD13	4:K:796:MET:HE3	1.97	0.46
2:S:75:VAL:HG12	3:N:114:TYR:CE1	2.51	0.46
3:N:237:ASP:HB3	3:N:271:LEU:HD12	1.98	0.46

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:N:541:ARG:HB3	3:N:546:ARG:NH1	2.31	0.46
3:P:115:LEU:O	3:P:119:ASP:HB2	2.15	0.46
3:P:272:ASN:OD1	3:P:273:GLY:N	2.48	0.46
3:G:260:PHE:O	3:G:264:VAL:HG12	2.16	0.46
3:G:467:VAL:HG22	3:G:493:LEU:HD13	1.98	0.46
3:G:698:LYS:O	3:G:702:LYS:HG2	2.15	0.46
4:I:49:GLY:N	4:I:50:PRO:HD2	2.31	0.46
4:I:878:GLY:HA2	4:I:885:TYR:CG	2.51	0.46
4:J:432:LEU:HD22	4:J:897:MET:HE1	1.98	0.46
4:J:477:LEU:HD11	4:J:483:LEU:HD11	1.97	0.46
4:J:553:ALA:O	4:J:557:ASP:HB2	2.16	0.46
4:J:832:LYS:HE3	4:J:846:HIS:HE1	1.81	0.46
4:L:383:GLY:O	4:L:384:ASP:HB2	2.15	0.46
4:L:451:THR:HA	4:L:454:MET:HE3	1.96	0.46
2:c:81:ILE:HG23	3:O:220:PHE:HB2	1.98	0.46
2:f:38:ALA:HB1	2:f:63:ALA:HA	1.97	0.46
3:N:697:LYS:O	3:N:701:GLU:HG2	2.16	0.46
3:P:129:PHE:HZ	3:P:138:TYR:CD2	2.34	0.46
3:P:680:ARG:CZ	4:K:423:ASN:HD22	2.29	0.46
3:R:549:ASP:OD2	3:R:585:TYR:OH	2.28	0.46
3:G:366:TRP:O	3:G:370:GLN:HG2	2.16	0.46
3:G:627:GLN:HA	3:G:635:TRP:HA	1.97	0.46
4:I:136:ARG:HA	4:I:139:MET:HE2	1.98	0.46
4:I:795:ASP:O	4:I:796:MET:C	2.58	0.46
4:J:633:HIS:O	4:J:637:LYS:HG2	2.15	0.46
4:L:449:MET:HE2	4:L:468:LEU:HD21	1.98	0.46
1:a:48:VAL:HG12	1:a:50:GLY:H	1.80	0.46
2:d:81:ILE:HA	3:P:220:PHE:CD1	2.51	0.46
2:g:35:ARG:NH1	3:H:116:VAL:HA	2.30	0.46
3:N:208:ASP:HB2	3:N:211:MET:HB2	1.97	0.46
3:N:323:ARG:HH11	3:O:290:LEU:HD23	1.80	0.46
3:O:352:LEU:O	3:O:356:GLN:HG2	2.16	0.46
3:Q:592:MET:SD	3:R:479:ILE:HD11	2.56	0.46
3:G:352:LEU:O	3:G:356:GLN:HG2	2.16	0.46
4:I:660:ARG:NH2	4:I:715:GLU:OE1	2.49	0.46
4:J:509:ILE:HD11	4:J:854:ILE:HG12	1.98	0.46
4:J:940:VAL:O	4:J:944:ILE:HG12	2.15	0.46
4:K:684:ALA:HB1	4:K:740:HIS:NE2	2.31	0.46
4:L:340:LEU:HD11	4:L:404:ILE:HG13	1.97	0.46
4:L:775:ASP:OD1	4:L:789:ASN:ND2	2.48	0.46
2:S:128:GLN:HG2	3:O:82:GLN:HG2	1.98	0.46

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:N:347:PRO:HA	3:N:350:GLU:HG2	1.98	0.46
3:N:587:SER:HB3	3:O:479:ILE:HG12	1.98	0.46
3:O:382:ASN:ND2	3:O:412:ASN:HB3	2.31	0.46
3:O:413:THR:HG21	3:O:416:PHE:CZ	2.51	0.46
3:R:238:ALA:O	3:R:241:THR:OG1	2.32	0.46
4:J:262:THR:OG1	4:J:1117:ASN:OD1	2.29	0.46
3:R:583:VAL:HG11	3:R:596:GLN:HG3	1.98	0.45
3:R:621:ILE:HD11	3:R:663:MET:HB3	1.98	0.45
3:G:607:THR:HG23	3:G:620:VAL:HG22	1.97	0.45
3:H:101:MET:O	3:H:105:ARG:HG3	2.16	0.45
3:H:680:ARG:CZ	4:I:423:ASN:HD22	2.30	0.45
4:I:550:LYS:HD3	4:I:554:THR:HG22	1.98	0.45
4:I:978:GLU:O	4:L:370:ASN:HB3	2.16	0.45
4:J:743:GLN:HG3	4:J:744:PHE:CD1	2.52	0.45
4:J:832:LYS:CE	4:J:846:HIS:HE1	2.27	0.45
4:K:140:ARG:HA	4:K:143:LEU:HG	1.98	0.45
4:K:838:ASP:HB2	4:K:841:LYS:HB2	1.98	0.45
4:L:54:MET:HE2	4:L:54:MET:HA	1.97	0.45
4:L:673:HIS:ND1	4:L:946:PRO:HA	2.31	0.45
3:M:106:VAL:HG21	3:M:165:ARG:HH21	1.80	0.45
3:N:50:LYS:NZ	3:N:159:GLU:OE2	2.49	0.45
3:O:112:ALA:HB2	3:O:150:TYR:CD2	2.51	0.45
3:O:356:GLN:O	3:O:360:GLN:HG2	2.17	0.45
3:P:197:ASN:O	3:P:201:GLU:HG2	2.16	0.45
3:R:306:ASN:OD1	3:R:351:TRP:HZ3	1.99	0.45
3:G:628:VAL:O	3:G:629:ASN:OD1	2.33	0.45
4:J:698:SER:OG	4:J:773:ASP:HB2	2.16	0.45
4:L:760:ILE:HG13	4:L:761:GLU:N	2.32	0.45
1:b:47:GLU:HB3	3:M:358:GLN:OE1	2.17	0.45
3:O:516:LEU:O	3:O:520:MET:HG3	2.16	0.45
3:P:626:MET:O	3:P:638:GLY:HA3	2.16	0.45
3:Q:662:THR:HG22	4:L:1119:LYS:NZ	2.30	0.45
3:H:473:GLU:OE1	3:H:487:THR:HA	2.15	0.45
4:I:820:THR:OG1	4:I:823:GLU:HG3	2.16	0.45
4:J:238:LEU:HD23	4:J:238:LEU:HA	1.78	0.45
4:J:644:PRO:HG2	4:J:814:MET:HG2	1.97	0.45
4:K:14:GLY:O	4:K:18:LYS:HG3	2.15	0.45
4:K:367:ALA:HB2	4:L:265:PRO:HB2	1.98	0.45
4:K:795:ASP:O	4:K:796:MET:C	2.58	0.45
4:L:1084:MET:HE2	4:L:1084:MET:HB3	1.69	0.45
2:f:40:GLU:O	2:f:44:GLN:HG2	2.15	0.45

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:M:700:GLU:HB3	3:N:368:LYS:HB3	1.99	0.45
3:P:493:LEU:HA	3:P:496:ILE:HG22	1.97	0.45
3:Q:550:ASP:O	3:Q:554:GLU:HG2	2.15	0.45
3:Q:564:GLU:H	3:Q:564:GLU:CD	2.24	0.45
3:R:121:MET:HE3	3:R:135:MET:HG2	1.99	0.45
3:R:125:LYS:HB3	3:R:125:LYS:HE2	1.45	0.45
3:R:664:TYR:OH	4:L:426:ILE:HD11	2.17	0.45
4:I:520:LYS:HD2	4:I:847:ALA:HB2	1.99	0.45
4:I:630:ILE:HD13	4:I:796:MET:HE3	1.98	0.45
4:I:641:MET:HE3	4:I:656:PHE:CD1	2.51	0.45
4:J:975:SER:O	4:J:978:GLU:HB2	2.16	0.45
4:J:1084:MET:HE2	4:J:1084:MET:HB3	1.74	0.45
4:J:1120:PHE:HE2	4:J:1301:LEU:HD23	1.80	0.45
4:K:1311:VAL:HG12	4:K:1313:LEU:HD22	1.97	0.45
4:L:476:VAL:HA	4:L:482:VAL:HA	1.99	0.45
4:L:600:TYR:CE2	4:L:948:ARG:HG2	2.51	0.45
2:S:118:ARG:NH2	3:O:68:ASP:HA	2.32	0.45
3:M:135:MET:HE1	3:M:179:TYR:CD1	2.52	0.45
3:N:188:GLN:O	3:N:192:LYS:HG3	2.17	0.45
3:Q:454:SER:C	3:Q:456:ASP:H	2.25	0.45
3:Q:520:MET:HB3	3:Q:526:ILE:CD1	2.44	0.45
3:R:205:VAL:HG12	3:R:206:LEU:HD23	1.99	0.45
3:R:669:SER:CB	3:R:682:ARG:HE	2.30	0.45
3:G:74:ILE:HA	3:G:77:LYS:HD2	1.98	0.45
3:G:454:SER:C	3:G:456:ASP:H	2.24	0.45
4:I:53:MET:HG3	4:I:54:MET:HE3	1.98	0.45
4:I:442:ASN:O	4:I:442:ASN:OD1	2.35	0.45
4:J:143:LEU:HD21	4:J:154:LEU:HD23	1.98	0.45
4:K:383:GLY:O	4:K:384:ASP:HB2	2.17	0.45
4:K:859:ALA:O	4:K:860:ARG:HB3	2.17	0.45
4:K:1099:GLY:O	4:K:1103:LYS:HG3	2.16	0.45
2:c:63:ALA:O	2:c:67:LEU:HD23	2.16	0.45
3:M:372:LYS:HB2	3:H:700:GLU:HG2	1.98	0.45
3:N:205:VAL:HG11	3:N:220:PHE:CE2	2.52	0.45
3:N:551:LYS:HZ3	3:O:543:LYS:HG3	1.81	0.45
3:O:700:GLU:HB3	3:P:368:LYS:HB3	1.97	0.45
3:P:205:VAL:HG21	3:P:220:PHE:CZ	2.51	0.45
3:P:645:ALA:O	3:P:649:ILE:HG12	2.16	0.45
3:R:261:LEU:CB	3:R:291:MET:HE1	2.47	0.45
3:R:628:VAL:HG22	3:R:634:SER:HB2	1.99	0.45
3:H:67:ALA:HA	3:H:70:ARG:CZ	2.47	0.45

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
4:I:476:VAL:HA	4:I:482:VAL:HA	1.98	0.45
4:I:594:SER:O	4:I:598:THR:HG23	2.15	0.45
4:J:54:MET:HB2	4:J:56:PHE:CE1	2.52	0.45
4:J:332:ILE:HG21	4:J:395:PHE:CD1	2.52	0.45
4:J:439:ARG:NH1	4:J:883:ASN:OD1	2.50	0.45
4:J:964:ALA:O	4:J:968:ILE:HG12	2.17	0.45
4:K:637:LYS:C	4:K:641:MET:HE3	2.41	0.45
4:K:1013:LEU:HD12	4:K:1055:PHE:CE1	2.51	0.45
4:L:940:VAL:O	4:L:944:ILE:HG12	2.17	0.45
2:S:60:LEU:HD12	3:M:196:MET:HE1	1.99	0.45
3:N:140:HIS:O	3:N:144:GLN:HG3	2.16	0.45
3:N:272:ASN:OD1	3:N:273:GLY:N	2.50	0.45
3:P:697:LYS:O	3:P:701:GLU:HG2	2.16	0.45
3:R:103:ALA:O	3:R:107:LYS:HG3	2.17	0.45
3:H:147:ALA:HB2	3:H:167:PHE:CZ	2.52	0.45
3:H:356:GLN:HA	3:H:359:VAL:HG12	1.97	0.45
4:I:17:GLN:HG3	4:I:1086:ARG:HG2	1.97	0.45
4:J:49:GLY:N	4:J:50:PRO:HD2	2.32	0.45
4:J:377:ARG:HG2	4:J:884:ALA:HB2	1.98	0.45
4:K:8:VAL:HG12	4:K:31:ARG:HH21	1.81	0.45
4:K:644:PRO:HG2	4:K:814:MET:HG2	1.98	0.45
4:L:609:ALA:HB1	4:L:616:GLN:HG2	1.99	0.45
4:L:953:GLN:OE1	4:L:954:ARG:NH2	2.49	0.45
1:A:47:GLU:CD	3:N:362:GLN:HE21	2.25	0.45
2:S:49:ASN:C	2:S:51:ASP:H	2.25	0.45
2:e:121:TYR:OH	3:G:96:ASP:OD2	2.27	0.45
3:N:366:TRP:HD1	3:O:289:ALA:HB1	1.81	0.45
3:O:557:LEU:HD22	3:O:573:ARG:HH21	1.81	0.45
3:R:55:MET:HE1	3:R:58:ALA:HB3	1.99	0.45
4:J:757:LEU:HD22	4:J:760:ILE:HD13	1.99	0.45
4:K:62:LYS:O	4:K:63:ALA:C	2.59	0.45
4:L:329:SER:OG	4:L:354:LYS:O	2.30	0.45
2:S:47:ILE:HD12	3:M:195:ILE:HD12	1.99	0.45
2:c:125:PHE:HD1	2:c:128:GLN:NE2	2.15	0.45
3:M:251:ALA:O	3:M:257:GLY:HA3	2.17	0.45
3:M:472:GLN:CD	3:H:599:LYS:HE3	2.41	0.45
3:M:543:LYS:HA	3:M:546:ARG:HB2	1.97	0.45
3:N:125:LYS:HD3	3:N:182:HIS:N	2.32	0.45
3:O:142:ARG:NE	3:O:145:GLU:OE2	2.50	0.45
3:O:420:ASP:OD1	3:O:420:ASP:N	2.46	0.45
3:Q:307:GLU:HA	3:Q:310:ARG:NH1	2.32	0.45

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:R:541:ARG:HB3	3:R:546:ARG:NH1	2.29	0.45
3:H:238:ALA:O	3:H:241:THR:HG22	2.17	0.45
3:H:252:SER:HB2	3:H:293:THR:HG23	1.97	0.45
4:I:1131:ASN:OD1	4:I:1132:ARG:N	2.50	0.45
4:J:11:ASP:CG	4:J:1022:LYS:HB3	2.42	0.45
4:K:449:MET:HE3	4:K:468:LEU:HD21	1.99	0.45
4:K:1103:LYS:O	4:K:1107:GLN:HG3	2.17	0.45
4:L:357:ASN:H	4:L:437:THR:HG23	1.81	0.45
4:L:698:SER:OG	4:L:773:ASP:HB2	2.17	0.45
4:L:1084:MET:HG2	4:L:1098:PHE:CZ	2.52	0.45
2:g:107:ILE:HD11	3:H:172:THR:CG2	2.47	0.45
3:O:451:GLN:HG2	3:O:503:LEU:HD21	1.99	0.45
3:H:272:ASN:OD1	3:H:273:GLY:N	2.49	0.45
3:H:503:LEU:O	3:H:507:LEU:HG	2.16	0.45
4:I:339:PRO:O	4:I:342:TYR:HB2	2.18	0.45
4:I:646:ILE:HG12	4:J:701:SER:HB2	1.98	0.45
4:J:339:PRO:O	4:J:342:TYR:HB2	2.17	0.45
4:J:878:GLY:HA2	4:J:885:TYR:CG	2.52	0.45
4:K:8:VAL:HB	4:K:9:PRO:HD3	1.98	0.45
4:L:463:VAL:HG22	4:L:875:ASN:ND2	2.29	0.45
4:L:530:ARG:O	4:L:534:ILE:HG13	2.17	0.45
4:L:679:ILE:O	4:L:683:GLY:HA2	2.17	0.45
3:P:587:SER:HB3	3:Q:479:ILE:HG12	1.99	0.44
4:I:54:MET:HB2	4:I:56:PHE:CD1	2.53	0.44
4:K:673:HIS:ND1	4:K:946:PRO:HA	2.32	0.44
4:K:938:LYS:HD2	4:K:1150:PHE:CE1	2.52	0.44
4:L:110:GLN:NE2	4:L:134:GLU:OE1	2.49	0.44
4:L:1242:GLY:HA2	4:L:1245:LEU:HD12	1.98	0.44
2:S:81:ILE:O	2:S:85:ILE:HG12	2.17	0.44
2:T:119:ARG:NH1	3:P:90:GLY:O	2.36	0.44
2:e:47:ILE:HD12	3:Q:195:ILE:HD12	2.00	0.44
3:M:550:ASP:O	3:M:554:GLU:HG2	2.17	0.44
3:O:159:GLU:N	3:O:159:GLU:OE2	2.51	0.44
3:O:197:ASN:O	3:O:201:GLU:HG2	2.17	0.44
3:O:463:ILE:O	3:O:467:VAL:HG23	2.17	0.44
3:P:92:LEU:HD12	3:P:93:LEU:N	2.32	0.44
3:P:101:MET:O	3:P:105:ARG:HG3	2.17	0.44
3:R:487:THR:HB	3:R:490:MET:HB2	1.98	0.44
3:H:394:ARG:HG3	3:H:452:ALA:HB1	1.99	0.44
4:J:410:ALA:O	4:J:414:ARG:HG3	2.18	0.44
4:J:431:ARG:NH2	4:J:484:SER:HB3	2.31	0.44

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
4:J:861:ARG:O	4:J:862:ASN:HB2	2.17	0.44
4:K:17:GLN:HG3	4:K:1086:ARG:HG2	2.00	0.44
4:K:1057:VAL:HG11	4:K:1060:LYS:HD3	1.99	0.44
4:L:581:MET:HE1	4:L:1194:PRO:HD3	1.99	0.44
4:L:1177:PRO:HD2	4:L:1314:ARG:HH22	1.81	0.44
1:b:48:VAL:HG13	3:M:314:ASN:HD21	1.81	0.44
2:c:57:ARG:HG3	2:c:57:ARG:HH11	1.82	0.44
2:g:34:GLY:O	2:g:37:GLN:HG3	2.17	0.44
2:g:69:SER:HB3	3:H:115:LEU:HA	1.99	0.44
3:M:120:VAL:O	3:M:124:ILE:HG13	2.18	0.44
3:O:187:SER:O	3:O:191:GLN:HG3	2.18	0.44
3:Q:251:ALA:O	3:Q:257:GLY:HA3	2.17	0.44
4:I:293:VAL:HG12	4:I:1130:ASN:HD22	1.82	0.44
4:I:940:VAL:O	4:I:944:ILE:HG12	2.18	0.44
4:J:339:PRO:HA	4:J:342:TYR:CD1	2.52	0.44
4:K:1277:GLU:HA	4:L:972:LEU:HD13	1.99	0.44
4:L:761:GLU:HG3	4:L:822:LYS:HG3	1.99	0.44
2:c:49:ASN:C	2:c:51:ASP:H	2.24	0.44
2:f:47:ILE:HD13	3:R:192:LYS:HA	1.99	0.44
3:O:673:MET:HG2	3:O:674:ASP:O	2.17	0.44
3:P:643:GLU:O	3:P:647:LYS:HG2	2.18	0.44
3:Q:123:LYS:HD3	3:Q:128:VAL:HG21	2.00	0.44
3:R:566:ALA:O	3:R:568:MET:N	2.50	0.44
3:H:122:GLN:O	3:H:126:GLU:HG2	2.18	0.44
3:H:553:PHE:CZ	3:H:557:LEU:HD11	2.53	0.44
4:I:416:LYS:HD2	4:I:416:LYS:HA	1.76	0.44
4:I:777:SER:HB3	4:I:787:SER:HB2	1.98	0.44
4:J:345:MET:HG2	4:J:1300:GLN:OE1	2.17	0.44
4:K:809:GLY:O	4:K:813:ILE:HG12	2.17	0.44
4:K:885:TYR:CZ	4:K:889:GLN:NE2	2.85	0.44
4:K:975:SER:HB3	4:K:978:GLU:HG3	1.98	0.44
4:L:49:GLY:N	4:L:50:PRO:HD2	2.33	0.44
4:L:514:PHE:HE1	4:L:851:THR:HG23	1.81	0.44
4:L:659:SER:O	4:L:660:ARG:HB2	2.17	0.44
2:S:38:ALA:HB2	2:S:66:GLU:OE1	2.18	0.44
2:e:76:GLN:HG2	3:R:114:TYR:CD2	2.52	0.44
2:e:120:ASP:O	2:e:124:ILE:HG13	2.18	0.44
3:M:576:ALA:HB1	3:M:597:MET:HE3	1.99	0.44
3:N:152:GLU:OE2	3:N:153:GLN:NE2	2.50	0.44
3:N:463:ILE:O	3:N:467:VAL:HG23	2.17	0.44
3:N:554:GLU:OE2	3:N:558:ASN:ND2	2.51	0.44

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:P:211:MET:O	3:P:214:ARG:HG3	2.17	0.44
3:Q:232:GLY:HA3	3:R:100:ALA:HB2	1.99	0.44
3:R:551:LYS:HZ3	3:G:543:LYS:HG3	1.82	0.44
3:G:211:MET:O	3:G:214:ARG:HG2	2.17	0.44
3:G:326:TRP:O	3:G:330:GLN:HG2	2.18	0.44
3:G:336:LEU:HB3	3:G:349:ARG:HH12	1.82	0.44
3:G:510:ASP:OD2	3:G:511:GLN:NE2	2.50	0.44
3:H:623:LYS:HE3	3:H:623:LYS:HB2	1.88	0.44
4:I:18:LYS:HD2	4:I:1082:GLU:OE2	2.18	0.44
4:I:362:VAL:HB	4:I:363:GLY:H	1.62	0.44
4:I:627:MET:HE3	4:I:627:MET:HB3	1.81	0.44
4:J:367:ALA:HB2	4:K:265:PRO:HB2	1.99	0.44
4:J:646:ILE:HG12	4:K:701:SER:HB2	1.99	0.44
4:J:795:ASP:O	4:J:796:MET:C	2.59	0.44
4:K:278:LEU:HD23	4:K:278:LEU:HA	1.84	0.44
4:K:339:PRO:HA	4:K:342:TYR:CD1	2.53	0.44
4:L:65:GLY:C	4:L:67:ARG:H	2.26	0.44
4:L:809:GLY:O	4:L:813:ILE:HG12	2.17	0.44
4:L:1042:LEU:HD13	4:L:1066:ASP:OD2	2.18	0.44
2:T:72:MET:SD	3:O:117:ASP:HB3	2.57	0.44
3:N:205:VAL:HG12	3:N:206:LEU:HD23	2.00	0.44
3:N:391:PHE:O	3:N:395:ILE:HG13	2.17	0.44
3:N:415:GLU:HG2	4:J:241:SER:HB2	1.99	0.44
3:O:131:THR:HG22	3:O:134:GLU:HG3	1.99	0.44
3:P:118:ASP:OD2	3:P:174:ARG:HD2	2.17	0.44
3:Q:673:MET:HG2	3:Q:674:ASP:O	2.17	0.44
3:G:545:GLN:HA	3:G:548:GLU:OE1	2.17	0.44
4:J:496:GLU:CG	4:J:1179:MET:HE3	2.47	0.44
4:L:477:LEU:HD11	4:L:483:LEU:HD11	2.00	0.44
4:L:1115:SER:HB3	4:L:1305:ILE:HG12	1.98	0.44
2:c:39:MET:O	2:c:43:ARG:HG3	2.18	0.44
2:e:34:GLY:O	2:e:37:GLN:HG3	2.18	0.44
3:M:383:LYS:O	3:M:387:ILE:HG13	2.18	0.44
3:M:463:ILE:O	3:M:467:VAL:HG23	2.18	0.44
3:N:202:LEU:HA	3:N:202:LEU:HD23	1.80	0.44
3:P:463:ILE:O	3:P:467:VAL:HG23	2.17	0.44
3:P:657:THR:HG22	3:P:659:LYS:H	1.83	0.44
3:Q:420:ASP:N	3:Q:420:ASP:OD1	2.43	0.44
3:H:698:LYS:HB2	3:H:698:LYS:HE2	1.72	0.44
4:I:1013:LEU:HB3	4:I:1049:ARG:HD3	2.00	0.44
4:K:49:GLY:N	4:K:50:PRO:HD2	2.33	0.44

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
4:L:577:MET:HE3	4:L:577:MET:HB3	1.76	0.44
1:E:45:GLY:HA2	3:P:448:LYS:HZ3	1.83	0.44
2:h:78:ILE:HD12	2:h:78:ILE:HA	1.87	0.44
3:M:448:LYS:HB3	3:M:448:LYS:NZ	2.33	0.44
3:N:637:GLN:O	3:N:641:ILE:HG13	2.17	0.44
3:P:526:ILE:HA	3:Q:480:ASN:HD21	1.82	0.44
3:Q:322:PRO:HB2	3:Q:363:MET:SD	2.58	0.44
3:Q:428:LYS:O	3:Q:428:LYS:HG3	2.17	0.44
3:Q:486:ARG:HG3	3:Q:488:PRO:HD3	2.00	0.44
3:R:269:VAL:HB	3:R:278:TYR:CZ	2.53	0.44
3:G:568:MET:HG2	3:G:573:ARG:HG2	2.00	0.44
3:G:613:VAL:CG1	4:I:414:ARG:HE	2.31	0.44
3:H:450:LEU:O	3:H:460:ARG:NE	2.51	0.44
4:J:724:PRO:O	4:J:728:GLU:HG3	2.17	0.44
4:J:761:GLU:CD	4:J:822:LYS:HG3	2.43	0.44
4:J:1276:THR:HG22	4:K:973:LYS:HB3	1.98	0.44
4:K:593:MET:HE3	4:K:593:MET:HB2	1.83	0.44
4:L:513:GLY:O	4:L:517:ILE:HG12	2.17	0.44
4:L:938:LYS:HD2	4:L:1150:PHE:CE1	2.53	0.44
4:L:1084:MET:HE1	4:L:1105:VAL:CG1	2.47	0.44
1:F:40:GLU:OE1	3:G:373:ALA:HB2	2.17	0.44
2:d:52:LEU:HB3	2:d:56:ALA:HB3	2.00	0.44
2:f:72:MET:SD	3:G:121:MET:HE2	2.57	0.44
2:h:78:ILE:HB	3:H:211:MET:HE1	1.99	0.44
3:O:102:GLU:HG3	3:O:162:ASP:OD2	2.17	0.44
3:O:616:ASP:OD1	3:O:616:ASP:N	2.51	0.44
3:Q:147:ALA:HB2	3:Q:167:PHE:CZ	2.53	0.44
3:G:410:ASN:C	3:G:412:ASN:H	2.24	0.44
3:G:541:ARG:HD2	3:G:541:ARG:HA	1.74	0.44
3:H:437:ILE:HD12	3:H:441:ALA:HB3	2.00	0.44
4:I:268:TRP:CD2	4:I:305:LEU:HD22	2.53	0.44
4:I:449:MET:HE3	4:I:449:MET:HB3	1.83	0.44
4:I:641:MET:SD	4:I:807:VAL:HB	2.57	0.44
4:J:109:ASN:HB2	4:J:138:TYR:CE2	2.53	0.44
4:J:278:LEU:HD23	4:J:278:LEU:HA	1.83	0.44
4:J:593:MET:HB3	4:J:597:GLU:OE1	2.17	0.44
4:J:1030:SER:HB2	4:J:1074:ARG:HH12	1.83	0.44
4:K:9:PRO:HB3	4:K:13:ASP:OD2	2.18	0.44
4:K:581:MET:HE1	4:K:595:ARG:HG3	2.00	0.44
4:L:395:PHE:CD1	4:L:429:MET:HE1	2.53	0.44
3:N:101:MET:O	3:N:105:ARG:HG3	2.18	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:O:543:LYS:H	3:O:546:ARG:HG3	1.83	0.43
3:H:146:GLY:HA2	3:H:149:VAL:HG22	1.99	0.43
4:I:880:PHE:HB2	4:J:974:TYR:CZ	2.53	0.43
4:I:936:PHE:HE2	4:I:1009:ILE:HG21	1.83	0.43
4:I:1013:LEU:HD22	4:I:1055:PHE:CE1	2.53	0.43
4:J:1029:ALA:HB3	4:J:1031:VAL:HG12	2.00	0.43
4:K:532:VAL:HG12	4:K:848:LEU:HD11	2.00	0.43
4:K:627:MET:HE3	4:K:627:MET:HB3	1.90	0.43
4:L:354:LYS:HB3	4:L:354:LYS:HE2	1.62	0.43
4:L:743:GLN:HG3	4:L:744:PHE:CD1	2.53	0.43
3:N:138:TYR:CE1	3:N:142:ARG:HG3	2.53	0.43
3:O:117:ASP:OD2	3:O:174:ARG:HD2	2.17	0.43
3:O:160:ASP:OD1	3:O:160:ASP:N	2.51	0.43
3:O:322:PRO:HB2	3:O:363:MET:SD	2.58	0.43
3:O:483:LEU:HD11	3:O:490:MET:HE1	2.00	0.43
3:O:655:TRP:CZ3	3:O:656:ILE:HG13	2.53	0.43
3:P:323:ARG:HH11	3:Q:290:LEU:HD23	1.83	0.43
3:R:149:VAL:O	3:R:152:GLU:HG3	2.18	0.43
4:I:278:LEU:HD23	4:I:278:LEU:HA	1.82	0.43
4:I:552:GLY:HA2	4:I:648:GLY:HA3	1.99	0.43
4:K:379:SER:H	4:K:1282:THR:HB	1.83	0.43
4:K:523:GLY:O	4:K:524:SER:HB3	2.17	0.43
4:K:878:GLY:HA2	4:K:885:TYR:CG	2.53	0.43
4:K:1061:GLN:OE1	4:K:1065:MET:HE3	2.18	0.43
4:L:339:PRO:HA	4:L:342:TYR:CD1	2.53	0.43
3:N:306:ASN:ND2	3:N:351:TRP:HZ3	2.13	0.43
3:O:631:ASP:O	3:O:634:SER:OG	2.36	0.43
3:P:202:LEU:HD23	3:P:202:LEU:HA	1.82	0.43
3:R:120:VAL:O	3:R:124:ILE:HG12	2.18	0.43
3:G:103:ALA:O	3:G:107:LYS:HG3	2.19	0.43
3:H:112:ALA:HB2	3:H:150:TYR:HD2	1.80	0.43
4:I:432:LEU:HD22	4:I:897:MET:HE1	2.01	0.43
4:I:659:SER:C	4:I:661:HIS:H	2.27	0.43
4:I:790:ASP:O	4:I:948:ARG:NH1	2.49	0.43
4:J:488:PRO:HB3	4:J:1185:LEU:HD13	2.00	0.43
4:J:526:ASP:HB3	4:J:529:ILE:HB	1.99	0.43
4:K:18:LYS:HD2	4:K:1082:GLU:OE2	2.18	0.43
4:K:414:ARG:HG2	4:K:414:ARG:NH1	2.33	0.43
4:K:553:ALA:O	4:K:554:THR:OG1	2.31	0.43
4:L:40:PHE:HA	4:L:53:MET:HE1	2.00	0.43
4:L:563:HIS:NE2	4:L:1202:GLY:O	2.39	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
4:L:859:ALA:O	4:L:860:ARG:HB3	2.17	0.43
2:e:23:GLN:O	2:e:27:ILE:HG23	2.18	0.43
2:f:67:LEU:O	3:G:115:LEU:HD21	2.18	0.43
3:M:116:VAL:HG11	3:M:143:LEU:HD23	2.00	0.43
3:O:120:VAL:O	3:O:124:ILE:HG13	2.18	0.43
3:O:410:ASN:C	3:O:412:ASN:H	2.25	0.43
3:P:580:TYR:HB2	3:P:597:MET:HE2	1.99	0.43
3:R:195:ILE:CD1	3:R:239:GLN:HG2	2.48	0.43
3:R:205:VAL:HG11	3:R:220:PHE:CE2	2.53	0.43
3:G:463:ILE:O	3:G:467:VAL:HG23	2.18	0.43
3:H:568:MET:SD	3:H:573:ARG:HD2	2.59	0.43
4:K:561:ARG:O	4:K:565:THR:HG23	2.19	0.43
4:L:39:ARG:HA	4:L:39:ARG:HD3	1.68	0.43
4:L:339:PRO:O	4:L:342:TYR:HB2	2.18	0.43
4:L:627:MET:HE3	4:L:627:MET:HB3	1.80	0.43
4:L:651:LYS:HG3	4:L:651:LYS:O	2.19	0.43
4:L:754:ILE:HB	4:L:853:LYS:HD3	2.00	0.43
4:L:1013:LEU:HD11	4:L:1047:MET:HB2	1.99	0.43
4:L:1090:VAL:CG2	4:L:1109:LYS:HD3	2.48	0.43
2:T:128:GLN:N	2:T:128:GLN:OE1	2.52	0.43
2:h:128:GLN:OE1	2:h:128:GLN:N	2.52	0.43
3:M:142:ARG:NE	3:M:145:GLU:OE2	2.51	0.43
3:M:381:MET:HE2	3:N:299:PHE:CZ	2.53	0.43
3:N:125:LYS:HZ2	3:N:182:HIS:HB2	1.82	0.43
3:P:435:MET:HE1	3:Q:318:ASN:ND2	2.33	0.43
3:P:510:ASP:OD1	3:P:511:GLN:N	2.51	0.43
3:Q:410:ASN:C	3:Q:412:ASN:H	2.27	0.43
3:Q:695:ASN:ND2	4:L:245:MET:HE2	2.34	0.43
3:Q:700:GLU:HB3	3:R:368:LYS:HB3	2.00	0.43
3:R:453:ASP:C	3:R:455:LYS:H	2.26	0.43
3:G:435:MET:HB3	3:G:437:ILE:HG23	2.01	0.43
4:I:468:LEU:HD12	4:I:474:ALA:O	2.18	0.43
4:K:610:ILE:HG13	4:K:627:MET:HG2	1.99	0.43
4:K:1300:GLN:O	4:K:1304:LYS:HG2	2.18	0.43
4:K:1315:GLU:HG3	4:K:1316:ARG:HG2	2.00	0.43
4:L:565:THR:HG22	4:L:1263:MET:HE1	2.00	0.43
4:L:712:MET:HE3	4:L:712:MET:HB3	1.73	0.43
3:M:394:ARG:NH2	3:M:452:ALA:O	2.52	0.43
3:M:686:GLU:HB3	4:J:156:THR:HG21	2.01	0.43
3:N:205:VAL:HG21	3:N:220:PHE:CZ	2.53	0.43
3:N:594:MET:HE2	3:N:594:MET:HA	1.99	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:P:437:ILE:HD11	3:P:445:MET:HE2	2.00	0.43
3:R:519:ASP:OD1	3:R:523:LYS:HE2	2.17	0.43
3:H:50:LYS:NZ	3:H:159:GLU:OE2	2.52	0.43
3:H:205:VAL:HG11	3:H:220:PHE:CE2	2.54	0.43
4:I:363:GLY:O	4:I:443:SER:HB3	2.19	0.43
4:I:921:LYS:HA	4:I:1168:LYS:HZ3	1.82	0.43
4:J:1018:THR:HG22	4:J:1021:GLU:HG2	1.99	0.43
4:K:339:PRO:O	4:K:342:TYR:HB2	2.18	0.43
4:K:459:GLU:HG2	4:K:463:VAL:C	2.43	0.43
4:L:553:ALA:O	4:L:557:ASP:HB2	2.18	0.43
3:M:140:HIS:O	3:M:144:GLN:HG2	2.19	0.43
3:M:363:MET:SD	3:N:286:GLN:HA	2.58	0.43
3:M:423:ASN:HB3	3:M:427:LYS:NZ	2.33	0.43
3:M:451:GLN:HE21	3:M:503:LEU:HD22	1.84	0.43
3:O:454:SER:C	3:O:456:ASP:H	2.25	0.43
3:Q:568:MET:SD	3:Q:573:ARG:HA	2.59	0.43
3:H:568:MET:HE2	3:H:601:LEU:CD1	2.45	0.43
4:I:659:SER:O	4:I:660:ARG:HB2	2.17	0.43
4:I:1080:ALA:O	4:I:1084:MET:HB2	2.19	0.43
4:J:340:LEU:HD11	4:J:404:ILE:HG13	2.01	0.43
4:K:861:ARG:O	4:K:862:ASN:HB2	2.18	0.43
4:K:1163:PRO:O	4:K:1167:ARG:HG3	2.18	0.43
4:L:155:GLU:OE1	4:L:251:ARG:HD2	2.18	0.43
4:L:266:THR:OG1	4:L:267:ARG:N	2.52	0.43
4:L:858:ARG:HD3	4:L:858:ARG:HA	1.75	0.43
3:M:403:ASP:HB3	3:M:406:ASP:OD2	2.18	0.43
3:O:486:ARG:HG3	3:O:488:PRO:HD3	2.00	0.43
3:O:695:ASN:ND2	4:K:245:MET:HE2	2.33	0.43
3:H:565:ILE:HD13	3:H:601:LEU:HD12	2.01	0.43
4:I:80:GLU:OE1	4:I:961:THR:HB	2.19	0.43
4:I:473:GLY:HA2	4:I:492:LYS:NZ	2.33	0.43
4:I:681:ARG:NH2	4:L:1243:SER:HA	2.34	0.43
4:J:609:ALA:HB1	4:J:616:GLN:HG2	1.99	0.43
4:K:128:PHE:HE2	4:K:153:GLN:HB3	1.82	0.43
4:L:395:PHE:HD1	4:L:429:MET:HE1	1.84	0.43
2:d:67:LEU:O	3:Q:115:LEU:HD21	2.19	0.43
2:h:118:ARG:NH2	3:N:68:ASP:OD1	2.52	0.43
3:N:467:VAL:HG12	3:N:511:GLN:HG3	2.01	0.43
3:N:566:ALA:O	3:N:568:MET:N	2.52	0.43
3:O:614:ASP:OD1	3:P:455:LYS:NZ	2.51	0.43
3:P:431:GLU:O	3:P:435:MET:HG3	2.19	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:Q:201:GLU:OE1	3:R:110:ARG:NH1	2.41	0.43
4:I:712:MET:HE3	4:I:712:MET:HB3	1.80	0.43
4:I:780:MET:HE3	4:I:780:MET:HB3	1.84	0.43
4:J:552:GLY:O	4:K:779:THR:OG1	2.36	0.43
4:L:344:PRO:HB3	4:L:1124:PHE:HD1	1.83	0.43
3:M:267:LYS:HZ2	3:M:267:LYS:HG3	1.66	0.43
3:N:229:LEU:HD22	3:O:84:ARG:NH2	2.34	0.43
3:O:518:MET:HE2	3:O:518:MET:HB3	1.94	0.43
3:O:653:ASN:HB2	3:O:656:ILE:HD12	2.01	0.43
3:O:660:GLN:HA	3:O:675:THR:HG23	1.99	0.43
3:P:205:VAL:HG12	3:P:206:LEU:HD23	2.01	0.43
3:P:706:LYS:HD2	3:P:706:LYS:HA	1.79	0.43
3:Q:261:LEU:HA	3:Q:264:VAL:HG12	2.00	0.43
3:Q:507:LEU:HD23	3:Q:507:LEU:HA	1.87	0.43
3:Q:627:GLN:O	3:Q:685:LYS:HE3	2.18	0.43
3:R:152:GLU:OE2	3:R:153:GLN:NE2	2.51	0.43
3:R:279:ARG:NH2	3:R:284:GLU:OE2	2.51	0.43
4:I:265:PRO:HG3	4:I:988:LEU:HD11	2.00	0.43
4:I:273:TRP:HB2	4:I:278:LEU:HG	2.00	0.43
4:I:465:TYR:CD2	4:I:491:PRO:HB2	2.54	0.43
4:I:555:ALA:N	4:I:816:SER:HB3	2.34	0.43
4:I:604:ARG:HH11	4:L:1248:GLN:HG2	1.82	0.43
4:I:996:LEU:HB2	4:I:1085:LEU:HD11	2.01	0.43
4:I:1129:LYS:HA	4:I:1129:LYS:HD3	1.86	0.43
4:I:1141:ILE:HD11	4:I:1302:VAL:HG22	2.01	0.43
4:J:1060:LYS:HE3	4:J:1060:LYS:HB2	1.84	0.43
4:K:796:MET:HA	4:K:800:MET:H	1.84	0.43
4:K:798:ARG:H	4:K:798:ARG:HG2	1.61	0.43
4:L:28:ASP:OD2	4:L:1089:LYS:HE3	2.19	0.43
4:L:362:VAL:HB	4:L:363:GLY:H	1.66	0.43
4:L:913:LEU:HD22	4:L:918:TYR:CE1	2.54	0.43
3:O:467:VAL:HG22	3:O:493:LEU:HD13	2.01	0.42
3:P:268:LYS:HA	3:P:277:THR:HA	2.01	0.42
3:P:541:ARG:HB3	3:P:546:ARG:NH1	2.31	0.42
3:P:566:ALA:O	3:P:568:MET:N	2.52	0.42
3:R:112:ALA:HB2	3:R:150:TYR:CD1	2.54	0.42
3:R:283:GLY:O	3:R:287:TRP:N	2.46	0.42
3:G:309:TYR:OH	3:G:336:LEU:HD11	2.19	0.42
3:G:491:ASP:OD1	3:G:494:ARG:NH1	2.52	0.42
3:G:627:GLN:O	3:G:685:LYS:HE3	2.18	0.42
4:I:72:PRO:CB	4:I:76:ARG:HB2	2.49	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
4:I:760:ILE:HG13	4:I:761:GLU:N	2.34	0.42
4:I:1001:ARG:NE	4:I:1077:ASP:OD2	2.49	0.42
4:I:1243:SER:HB3	4:J:784:GLN:NE2	2.34	0.42
4:L:96:LYS:HB3	4:L:96:LYS:HE2	1.82	0.42
2:T:75:VAL:HA	2:T:78:ILE:HG22	2.01	0.42
3:M:612:ASP:HB2	3:M:658:ASN:HD22	1.84	0.42
3:N:623:LYS:HE3	3:N:623:LYS:HB2	1.89	0.42
3:O:495:ARG:HE	3:O:495:ARG:HB3	1.69	0.42
3:O:623:LYS:HE3	3:O:623:LYS:HB2	1.88	0.42
3:Q:116:VAL:HG11	3:Q:143:LEU:HA	2.01	0.42
3:Q:211:MET:O	3:Q:214:ARG:HG2	2.18	0.42
3:Q:510:ASP:OD2	3:Q:511:GLN:NE2	2.44	0.42
3:R:363:MET:HE2	3:R:363:MET:HB2	1.93	0.42
3:R:493:LEU:HA	3:R:496:ILE:HG22	2.01	0.42
3:R:666:GLN:HB2	3:R:671:TYR:HE2	1.84	0.42
3:G:380:SER:HA	3:G:383:LYS:HE2	2.01	0.42
3:H:203:ASN:O	3:H:207:GLN:HG3	2.19	0.42
3:H:282:ILE:HG23	3:H:286:GLN:OE1	2.19	0.42
4:I:675:LYS:HG3	4:I:687:LEU:CD2	2.50	0.42
4:I:803:TYR:O	4:I:807:VAL:HG22	2.19	0.42
4:I:861:ARG:O	4:I:862:ASN:HB2	2.18	0.42
4:I:919:LYS:HG2	4:I:921:LYS:H	1.84	0.42
4:I:957:GLU:HG2	4:I:980:ALA:CB	2.49	0.42
4:J:809:GLY:O	4:J:813:ILE:HG12	2.19	0.42
4:J:1012:THR:OG1	4:J:1043:ILE:HG21	2.19	0.42
4:J:1159:ALA:O	4:J:1167:ARG:HG2	2.18	0.42
4:K:456:PHE:O	4:K:457:GLU:HB2	2.19	0.42
4:L:257:ASP:HB3	4:L:260:LYS:HB3	2.01	0.42
4:L:565:THR:HG21	4:L:1268:VAL:HG13	2.00	0.42
4:L:644:PRO:HG2	4:L:814:MET:HG2	2.01	0.42
4:L:760:ILE:HD11	4:L:825:LYS:HG3	2.02	0.42
2:S:64:SER:O	2:S:68:THR:HG23	2.18	0.42
2:c:23:GLN:O	2:c:27:ILE:HG23	2.19	0.42
2:g:22:ALA:O	2:g:26:MET:HG2	2.19	0.42
2:g:40:GLU:O	2:g:44:GLN:HG2	2.18	0.42
3:M:119:ASP:O	3:M:123:LYS:HG2	2.19	0.42
3:M:607:THR:HG23	3:M:620:VAL:HG22	2.02	0.42
3:N:72:ASN:HA	3:N:75:ILE:HG22	2.01	0.42
3:N:640:ASP:OD2	3:O:417:LYS:HB3	2.19	0.42
3:P:356:GLN:O	3:P:359:VAL:HG12	2.19	0.42
3:Q:131:THR:HG22	3:Q:134:GLU:HG3	2.01	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:Q:428:LYS:HE2	3:Q:449:TYR:CE2	2.54	0.42
3:Q:613:VAL:HG23	3:Q:614:ASP:H	1.84	0.42
3:G:197:ASN:O	3:G:201:GLU:HG2	2.19	0.42
3:G:680:ARG:NH1	4:I:253:ASP:OD2	2.53	0.42
3:H:269:VAL:HB	3:H:278:TYR:CZ	2.54	0.42
4:J:1163:PRO:O	4:J:1167:ARG:HG3	2.18	0.42
4:K:340:LEU:HD11	4:K:404:ILE:HG13	2.00	0.42
4:K:362:VAL:HG23	4:K:446:LEU:HD11	2.01	0.42
4:K:624:ARG:HA	4:K:624:ARG:HD2	1.87	0.42
4:L:333:ILE:HG22	4:L:433:GLU:OE1	2.18	0.42
4:L:350:GLY:HA2	4:L:392:LEU:HD23	2.01	0.42
4:L:659:SER:C	4:L:661:HIS:H	2.26	0.42
3:M:674:ASP:OD1	3:M:674:ASP:N	2.44	0.42
3:P:115:LEU:HD12	3:P:119:ASP:OD2	2.20	0.42
3:P:229:LEU:HG	3:P:234:ILE:HD11	2.02	0.42
3:P:232:GLY:HA3	3:Q:100:ALA:HB2	2.01	0.42
3:P:656:ILE:HD11	3:P:679:VAL:HG21	2.00	0.42
3:Q:326:TRP:O	3:Q:330:GLN:HG2	2.20	0.42
3:R:271:LEU:HG	3:R:272:ASN:N	2.34	0.42
3:R:394:ARG:HG2	3:R:452:ALA:HB1	2.00	0.42
4:I:879:PHE:CE2	4:J:68:VAL:HG11	2.54	0.42
4:J:268:TRP:CD2	4:J:305:LEU:HD22	2.54	0.42
4:J:526:ASP:H	4:J:530:ARG:NH1	2.16	0.42
4:J:927:GLU:O	4:J:931:LEU:HG	2.19	0.42
4:K:522:LEU:HG	4:K:533:ALA:HB1	2.01	0.42
4:K:819:LYS:HE3	4:K:827:GLU:OE2	2.19	0.42
4:K:1171:LEU:HD22	4:K:1175:LEU:HD12	2.00	0.42
4:L:529:ILE:HG21	4:L:844:GLU:OE2	2.19	0.42
4:L:641:MET:HG2	4:L:655:ILE:HG13	2.01	0.42
4:L:1289:LYS:HE2	4:L:1289:LYS:HB2	1.88	0.42
2:S:52:LEU:HB3	2:S:56:ALA:HB3	2.01	0.42
2:d:37:GLN:O	2:d:41:ILE:HG13	2.19	0.42
3:N:252:SER:HB2	3:N:293:THR:HG23	2.00	0.42
3:O:232:GLY:HA3	3:P:100:ALA:HB2	2.00	0.42
3:P:66:LEU:O	3:P:70:ARG:HG2	2.18	0.42
3:P:680:ARG:NH1	4:K:423:ASN:HD22	2.18	0.42
3:Q:660:GLN:HA	3:Q:675:THR:HG23	2.01	0.42
4:J:355:LEU:HD21	4:J:433:GLU:HG3	2.00	0.42
4:J:525:ASP:HA	4:J:530:ARG:NH1	2.34	0.42
4:J:652:ALA:HB1	4:J:814:MET:HG3	2.00	0.42
4:K:72:PRO:HB3	4:K:76:ARG:HE	1.83	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
4:K:324:LEU:CD1	4:K:342:TYR:HE2	2.32	0.42
4:K:357:ASN:H	4:K:437:THR:HG23	1.84	0.42
4:K:529:ILE:HD13	4:K:831:LEU:HD13	2.01	0.42
4:K:570:TYR:OH	4:K:804:ASP:OD2	2.30	0.42
4:K:749:ILE:HA	4:K:752:GLU:CD	2.44	0.42
4:L:704:GLU:OE1	4:L:704:GLU:N	2.31	0.42
4:L:803:TYR:O	4:L:807:VAL:HG22	2.20	0.42
4:L:861:ARG:HH21	4:L:1205:LEU:HD11	1.83	0.42
4:L:1042:LEU:HA	4:L:1045:GLU:HG2	2.02	0.42
1:B:47:GLU:CD	3:P:362:GLN:HE21	2.27	0.42
2:c:120:ASP:O	2:c:124:ILE:HG13	2.20	0.42
2:e:71:ASN:HB3	3:R:122:GLN:HG2	2.02	0.42
2:h:75:VAL:HA	2:h:78:ILE:HG22	2.02	0.42
3:M:205:VAL:HG11	3:M:220:PHE:CE2	2.53	0.42
3:M:614:ASP:N	3:M:614:ASP:OD1	2.52	0.42
3:N:121:MET:HE1	3:N:139:ARG:HA	2.01	0.42
3:N:195:ILE:HD11	3:N:235:PRO:HD2	2.00	0.42
3:P:574:GLU:O	3:P:578:LYS:HG3	2.19	0.42
3:P:577:ARG:O	3:P:577:ARG:HD2	2.20	0.42
3:P:628:VAL:HG22	3:P:634:SER:HB2	2.01	0.42
3:R:148:LYS:HA	3:R:158:PRO:HG3	2.01	0.42
3:R:394:ARG:CG	3:R:452:ALA:HB1	2.49	0.42
3:R:455:LYS:HA	3:R:460:ARG:HD2	2.00	0.42
3:G:244:ILE:HD13	3:G:278:TYR:HE1	1.85	0.42
4:I:8:VAL:HB	4:I:9:PRO:HD3	2.00	0.42
4:I:509:ILE:HD13	4:I:859:ALA:HB1	2.02	0.42
4:J:496:GLU:OE1	4:J:1173:ARG:NH1	2.51	0.42
4:J:641:MET:HE1	4:J:766:LEU:CD2	2.49	0.42
4:L:842:THR:HA	4:L:845:VAL:HG22	2.01	0.42
2:c:107:ILE:HD11	3:P:172:THR:CG2	2.50	0.42
3:N:389:LYS:O	3:N:393:LYS:HG3	2.20	0.42
3:O:135:MET:HE3	3:O:135:MET:HB3	1.85	0.42
3:P:415:GLU:HG2	4:K:241:SER:HB2	2.02	0.42
3:P:612:ASP:OD1	3:P:615:GLY:HA2	2.20	0.42
3:Q:117:ASP:OD2	3:Q:174:ARG:HD2	2.20	0.42
3:Q:403:ASP:HB3	3:Q:406:ASP:OD2	2.19	0.42
3:Q:455:LYS:HD3	3:Q:460:ARG:NH1	2.30	0.42
3:R:156:ILE:H	3:R:156:ILE:HD12	1.84	0.42
3:R:237:ASP:O	3:R:241:THR:HG23	2.18	0.42
3:R:616:ASP:OD1	3:R:616:ASP:N	2.51	0.42
3:H:443:ASP:O	3:H:447:LEU:HD23	2.20	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:H:553:PHE:CD2	3:H:577:ARG:HD3	2.55	0.42
4:I:340:LEU:HD11	4:I:404:ILE:HG13	2.02	0.42
4:J:11:ASP:OD2	4:J:1022:LYS:HB3	2.19	0.42
4:J:324:LEU:CD1	4:J:342:TYR:HE2	2.32	0.42
4:J:671:ASP:OD1	4:J:672:ARG:N	2.51	0.42
4:J:1013:LEU:HD12	4:J:1055:PHE:CE1	2.55	0.42
4:K:600:TYR:CE2	4:K:948:ARG:HG2	2.55	0.42
4:K:652:ALA:HB1	4:K:814:MET:HG3	2.02	0.42
4:K:1245:LEU:HD13	4:L:955:LEU:HG	2.01	0.42
4:L:681:ARG:NH2	4:L:782:ASP:OD2	2.52	0.42
4:L:817:THR:HB	4:L:819:LYS:HE3	2.01	0.42
1:E:47:GLU:HB3	3:Q:358:GLN:OE1	2.20	0.42
3:M:410:ASN:C	3:M:412:ASN:H	2.26	0.42
3:N:114:TYR:CD1	3:N:174:ARG:HD3	2.55	0.42
3:N:602:LYS:HE3	3:O:468:THR:HG21	2.02	0.42
3:P:623:LYS:HE3	3:P:623:LYS:HB2	1.91	0.42
3:Q:612:ASP:HB2	3:Q:658:ASN:HD22	1.85	0.42
3:H:236:SER:C	3:H:238:ALA:N	2.76	0.42
4:I:684:ALA:HB1	4:I:740:HIS:NE2	2.35	0.42
4:I:938:LYS:O	4:I:942:GLN:HG2	2.19	0.42
4:J:1013:LEU:HD11	4:J:1047:MET:HB2	2.01	0.42
4:K:682:TYR:CE1	4:K:781:PRO:HD2	2.55	0.42
4:L:749:ILE:HD11	4:L:1061:GLN:HG2	2.00	0.42
2:c:38:ALA:HB2	2:c:66:GLU:OE1	2.20	0.42
2:c:42:MET:HE1	3:P:154:PHE:CE1	2.55	0.42
2:d:76:GLN:HG2	3:Q:114:TYR:CG	2.55	0.42
2:g:46:ASN:OD1	2:g:48:GLN:N	2.40	0.42
3:O:319:GLN:NE2	3:O:328:MET:HE1	2.35	0.42
3:O:545:GLN:HG2	3:O:548:GLU:OE1	2.20	0.42
3:O:607:THR:HG23	3:O:620:VAL:HG22	2.02	0.42
3:R:623:LYS:HE3	3:R:623:LYS:HB2	1.91	0.42
4:I:510:LYS:HA	4:I:510:LYS:HD2	1.86	0.42
4:I:748:SER:O	4:I:752:GLU:HG3	2.20	0.42
4:K:803:TYR:O	4:K:807:VAL:HG22	2.20	0.42
4:K:934:SER:HB3	4:K:1101:LEU:HB2	2.01	0.42
4:K:1095:SER:O	4:K:1103:LYS:HG2	2.20	0.42
4:K:1121:LEU:HD12	4:K:1121:LEU:HA	1.92	0.42
4:K:1164:LYS:HA	4:K:1167:ARG:HD2	2.00	0.42
4:L:754:ILE:HA	4:L:858:ARG:HH11	1.84	0.42
2:d:76:GLN:HG2	3:Q:114:TYR:CD2	2.55	0.42
3:N:268:LYS:HA	3:N:277:THR:HA	2.01	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:N:358:GLN:O	3:N:362:GLN:HG3	2.19	0.42
3:N:486:ARG:O	3:N:487:THR:C	2.63	0.42
3:O:435:MET:HG3	3:P:311:LEU:HD11	2.01	0.42
3:O:613:VAL:HG23	3:O:614:ASP:N	2.32	0.42
3:P:114:TYR:CD1	3:P:174:ARG:HD3	2.55	0.42
3:P:120:VAL:O	3:P:124:ILE:HG12	2.20	0.42
3:P:188:GLN:O	3:P:192:LYS:HG3	2.19	0.42
3:P:589:ASN:ND2	3:P:592:MET:HG2	2.34	0.42
3:R:70:ARG:O	3:R:74:ILE:HG12	2.20	0.42
3:R:459:PHE:O	3:R:463:ILE:HG12	2.20	0.42
3:G:623:LYS:HE3	3:G:623:LYS:HB2	1.92	0.42
3:H:92:LEU:HD12	3:H:93:LEU:N	2.34	0.42
4:I:262:THR:HG22	4:I:263:ILE:HG12	2.01	0.42
4:J:71:GLY:C	4:J:73:ASP:N	2.74	0.42
4:J:378:THR:OG1	4:J:380:VAL:HG12	2.19	0.42
4:J:577:MET:HE3	4:J:577:MET:HB3	1.90	0.42
4:K:50:PRO:HA	4:K:57:THR:OG1	2.20	0.42
4:K:85:ALA:O	4:K:89:GLN:HG3	2.20	0.42
4:K:574:TYR:CD1	4:K:1194:PRO:HG2	2.55	0.42
4:K:659:SER:O	4:K:660:ARG:HB2	2.20	0.42
2:T:67:LEU:O	3:O:115:LEU:HD21	2.20	0.41
2:c:79:GLY:HA3	3:P:110:ARG:HH21	1.84	0.41
3:M:158:PRO:HG2	3:M:160:ASP:CG	2.44	0.41
3:M:323:ARG:HH11	3:N:290:LEU:HD23	1.85	0.41
3:N:147:ALA:HB2	3:N:167:PHE:CZ	2.55	0.41
3:N:528:PRO:O	3:N:532:LEU:HB2	2.20	0.41
3:O:259:ASP:HA	3:O:262:MET:HE3	2.02	0.41
3:O:303:ALA:O	3:O:307:GLU:HG2	2.20	0.41
3:O:413:THR:HG21	3:O:416:PHE:CE2	2.54	0.41
3:P:614:ASP:O	3:P:615:GLY:C	2.63	0.41
3:Q:244:ILE:HD13	3:Q:278:TYR:HE1	1.85	0.41
3:Q:592:MET:HE1	3:R:479:ILE:HD11	2.01	0.41
3:G:639:ARG:HG2	3:G:639:ARG:HH11	1.84	0.41
3:G:639:ARG:NH1	3:G:640:ASP:OD1	2.53	0.41
3:H:262:MET:HE3	3:H:291:MET:SD	2.59	0.41
4:I:974:TYR:CZ	4:L:880:PHE:HB2	2.55	0.41
4:K:268:TRP:CD2	4:K:305:LEU:HD22	2.55	0.41
4:K:641:MET:HE1	4:K:803:TYR:OH	2.19	0.41
4:K:646:ILE:HG12	4:K:1269:VAL:HG21	2.02	0.41
4:L:362:VAL:HG21	4:L:446:LEU:HD21	2.02	0.41
3:M:197:ASN:ND2	3:N:107:LYS:HB2	2.34	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:M:669:SER:OG	3:M:684:ASP:OD1	2.37	0.41
3:N:342:ASP:HB2	3:N:344:GLN:HE22	1.85	0.41
3:O:568:MET:SD	3:O:573:ARG:HG2	2.60	0.41
3:O:619:GLY:HA3	3:O:663:MET:HB2	2.02	0.41
3:P:87:LEU:HD11	3:P:94:TYR:HD1	1.86	0.41
3:Q:570:ALA:O	3:Q:573:ARG:HB2	2.20	0.41
3:R:118:ASP:OD2	3:R:174:ARG:HD2	2.19	0.41
3:R:680:ARG:CZ	4:L:423:ASN:HD22	2.32	0.41
4:I:596:GLU:HG2	4:I:949:ALA:HB2	2.01	0.41
4:J:357:ASN:H	4:J:437:THR:HG23	1.85	0.41
4:K:493:THR:HA	4:K:1179:MET:HE1	2.02	0.41
4:L:1029:ALA:HB2	4:L:1078:LYS:HB2	2.01	0.41
4:L:1061:GLN:NE2	4:L:1065:MET:HE3	2.33	0.41
4:L:1078:LYS:HB3	4:L:1078:LYS:HE3	1.91	0.41
2:d:69:SER:HB3	3:Q:115:LEU:HD23	2.02	0.41
3:M:413:THR:HG21	3:M:416:PHE:CZ	2.55	0.41
3:M:619:GLY:HA3	3:M:663:MET:HB2	2.02	0.41
3:M:673:MET:HG2	3:M:674:ASP:N	2.36	0.41
3:N:511:GLN:HB3	3:N:514:LEU:HB3	2.02	0.41
3:R:356:GLN:O	3:R:359:VAL:HG12	2.20	0.41
3:R:565:ILE:HD13	3:R:601:LEU:HD12	2.02	0.41
3:H:296:ARG:O	3:H:300:GLU:HG2	2.20	0.41
4:I:356:ILE:HG12	4:I:440:ASN:HB2	2.02	0.41
4:I:722:VAL:HG13	4:I:726:MET:HE2	2.03	0.41
4:J:251:ARG:HA	4:J:251:ARG:HD3	1.88	0.41
4:J:529:ILE:HD13	4:J:831:LEU:HD13	2.02	0.41
4:J:841:LYS:O	4:J:845:VAL:HG13	2.20	0.41
4:J:861:ARG:HH21	4:J:1205:LEU:HD11	1.85	0.41
4:K:944:ILE:HD12	4:K:1073:TRP:HB2	2.02	0.41
4:L:938:LYS:O	4:L:942:GLN:HG2	2.21	0.41
4:L:1241:MET:C	4:L:1243:SER:N	2.78	0.41
3:O:202:LEU:HA	3:O:202:LEU:HD12	1.78	0.41
3:O:444:ALA:O	3:O:448:LYS:HG3	2.21	0.41
3:Q:323:ARG:HH12	3:R:293:THR:HG21	1.84	0.41
3:R:336:LEU:HD23	3:R:349:ARG:NH1	2.36	0.41
3:H:493:LEU:HD21	3:H:515:PHE:HE1	1.85	0.41
3:H:566:ALA:O	3:H:568:MET:N	2.54	0.41
4:I:653:VAL:O	4:I:653:VAL:HG13	2.21	0.41
4:J:553:ALA:O	4:J:554:THR:OG1	2.34	0.41
4:J:565:THR:HG22	4:J:1263:MET:HE1	2.03	0.41
4:J:651:LYS:HE3	4:J:651:LYS:HB3	1.84	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
4:J:806:ARG:HD2	4:J:806:ARG:HA	1.82	0.41
4:J:1237:VAL:HG22	4:K:968:ILE:CG2	2.50	0.41
4:K:324:LEU:HD11	4:K:342:TYR:HE2	1.86	0.41
4:K:1084:MET:HE1	4:K:1105:VAL:CG2	2.51	0.41
4:K:1242:GLY:HA2	4:K:1245:LEU:HG	2.02	0.41
4:L:555:ALA:N	4:L:816:SER:HB3	2.35	0.41
4:L:822:LYS:HE3	4:L:822:LYS:HB3	1.83	0.41
3:N:259:ASP:HB3	3:N:263:ARG:HH12	1.86	0.41
3:O:185:PHE:CD2	3:O:186:LEU:HD12	2.56	0.41
3:Q:383:LYS:HB2	3:Q:416:PHE:CE2	2.56	0.41
3:R:188:GLN:O	3:R:192:LYS:HG3	2.21	0.41
3:H:598:THR:HG22	3:H:602:LYS:NZ	2.35	0.41
4:I:88:LYS:HB3	4:I:88:LYS:HE2	1.77	0.41
4:I:859:ALA:O	4:I:860:ARG:HB3	2.20	0.41
4:I:1019:ARG:HA	4:I:1022:LYS:HZ3	1.84	0.41
4:I:1237:VAL:HG22	4:J:968:ILE:CG2	2.50	0.41
4:J:387:TYR:HD1	4:J:1285:MET:HE1	1.86	0.41
4:J:456:PHE:HA	4:J:466:GLU:HB2	2.02	0.41
4:J:723:THR:HG23	4:J:726:MET:H	1.86	0.41
4:J:904:ARG:HD2	4:J:1312:ASN:HB2	2.02	0.41
4:J:1086:ARG:N	4:J:1087:PRO:HD2	2.35	0.41
4:K:550:LYS:HD3	4:K:554:THR:HG22	2.02	0.41
2:f:69:SER:HB3	3:G:115:LEU:HD23	2.02	0.41
2:f:69:SER:O	2:f:73:GLN:HB2	2.19	0.41
3:N:579:ILE:O	3:N:583:VAL:HG23	2.20	0.41
3:P:528:PRO:O	3:P:532:LEU:HB2	2.21	0.41
3:P:607:THR:HG23	3:P:617:THR:HG23	2.02	0.41
3:Q:527:ASP:O	3:Q:530:VAL:HG22	2.21	0.41
3:G:448:LYS:NZ	3:G:448:LYS:HB3	2.35	0.41
3:G:676:THR:OG1	3:G:678:GLN:HG3	2.20	0.41
4:I:938:LYS:HD2	4:I:1150:PHE:CE1	2.56	0.41
4:J:66:LEU:C	4:J:68:VAL:N	2.78	0.41
4:J:517:ILE:HG23	4:J:847:ALA:HB1	2.01	0.41
4:J:1098:PHE:O	4:J:1103:LYS:HE2	2.21	0.41
4:L:71:GLY:C	4:L:73:ASP:N	2.74	0.41
2:d:7:ILE:HB	2:d:8:PRO:HD3	2.01	0.41
3:N:79:THR:N	3:N:82:GLN:HE21	2.12	0.41
3:O:160:ASP:O	3:O:164:GLN:HG2	2.21	0.41
3:O:565:ILE:HG23	3:O:597:MET:HE1	2.02	0.41
3:P:178:LEU:HD23	3:P:178:LEU:HA	1.94	0.41
3:R:195:ILE:HD12	3:R:239:GLN:HG2	2.01	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:G:323:ARG:O	3:G:327:GLU:HG3	2.20	0.41
3:G:493:LEU:O	3:G:497:ARG:HB2	2.20	0.41
4:K:113:GLY:N	4:K:117:ASN:OD1	2.53	0.41
4:K:475:VAL:HG11	4:K:492:LYS:HB2	2.02	0.41
4:K:760:ILE:HG13	4:K:761:GLU:N	2.34	0.41
4:K:775:ASP:OD1	4:K:789:ASN:ND2	2.53	0.41
4:L:54:MET:HB2	4:L:56:PHE:CD1	2.55	0.41
4:L:941:ASP:OD2	4:L:1001:ARG:NH1	2.53	0.41
2:d:71:ASN:HD21	3:Q:122:GLN:HB2	1.86	0.41
3:N:149:VAL:O	3:N:152:GLU:HG3	2.21	0.41
3:N:568:MET:HE2	3:N:601:LEU:CD1	2.51	0.41
3:N:698:LYS:HB2	3:N:698:LYS:HE2	1.75	0.41
3:G:368:LYS:HZ2	3:G:368:LYS:HG3	1.67	0.41
4:I:251:ARG:HA	4:I:251:ARG:HD3	1.88	0.41
4:I:682:TYR:CE1	4:I:781:PRO:HD2	2.56	0.41
4:I:761:GLU:CD	4:I:822:LYS:HG3	2.46	0.41
4:I:1086:ARG:N	4:I:1087:PRO:HD2	2.35	0.41
4:J:1021:GLU:O	4:J:1026:LEU:HD11	2.21	0.41
4:K:121:GLU:HG2	4:K:125:LYS:HE3	2.02	0.41
4:K:679:ILE:O	4:K:683:GLY:HA2	2.20	0.41
4:L:344:PRO:HB3	4:L:1124:PHE:CD1	2.56	0.41
4:L:610:ILE:HG13	4:L:627:MET:HG2	2.03	0.41
1:B:48:VAL:HG12	1:B:50:GLY:H	1.86	0.41
2:T:23:GLN:HA	2:T:26:MET:HE2	2.02	0.41
2:h:108:ARG:NH2	2:h:117:TYR:OH	2.54	0.41
3:M:383:LYS:HB2	3:M:416:PHE:CE1	2.56	0.41
3:M:527:ASP:O	3:M:530:VAL:HG22	2.20	0.41
3:M:543:LYS:H	3:M:546:ARG:HG3	1.84	0.41
3:N:114:TYR:O	3:N:118:ASP:HB2	2.20	0.41
3:N:115:LEU:HG	3:N:116:VAL:N	2.34	0.41
3:N:118:ASP:OD1	3:N:139:ARG:NH1	2.53	0.41
3:N:195:ILE:CD1	3:N:239:GLN:HG2	2.45	0.41
3:N:262:MET:HE3	3:N:291:MET:HE2	2.03	0.41
3:N:682:ARG:HH12	3:N:684:ASP:HB2	1.86	0.41
3:O:209:PRO:HA	3:O:212:LEU:HD12	2.03	0.41
3:O:394:ARG:HD2	3:O:402:THR:HG21	2.03	0.41
3:P:329:LEU:HD23	3:P:329:LEU:HA	1.88	0.41
3:Q:205:VAL:HG11	3:Q:220:PHE:CE2	2.56	0.41
3:Q:307:GLU:HA	3:Q:310:ARG:HH12	1.86	0.41
3:R:92:LEU:HD12	3:R:93:LEU:N	2.35	0.41
3:R:614:ASP:O	3:R:615:GLY:C	2.64	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:G:142:ARG:HD2	3:G:142:ARG:N	2.36	0.41
3:G:210:ASP:OD1	3:G:211:MET:N	2.54	0.41
3:G:495:ARG:HE	3:G:495:ARG:HB3	1.58	0.41
3:G:513:GLU:O	3:G:517:THR:HG23	2.20	0.41
3:H:569:PRO:C	3:H:571:SER:H	2.29	0.41
3:H:664:TYR:OH	4:I:426:ILE:HD11	2.21	0.41
4:I:535:ASP:HB3	4:I:817:THR:HG22	2.03	0.41
4:I:806:ARG:NH1	4:I:856:THR:OG1	2.54	0.41
4:I:1060:LYS:HE3	4:I:1060:LYS:HB2	1.86	0.41
4:J:39:ARG:HA	4:J:39:ARG:HD3	1.71	0.41
4:J:72:PRO:HB2	4:J:76:ARG:O	2.20	0.41
4:J:635:ASP:O	4:J:639:GLU:HG3	2.20	0.41
4:J:754:ILE:HB	4:J:853:LYS:HD3	2.03	0.41
4:J:859:ALA:O	4:J:860:ARG:HB3	2.21	0.41
4:J:879:PHE:HE2	4:K:68:VAL:HG11	1.86	0.41
4:K:238:LEU:HA	4:K:238:LEU:HD23	1.73	0.41
4:K:584:PRO:HA	4:K:588:THR:HG23	2.03	0.41
4:K:780:MET:HE3	4:K:780:MET:HB3	1.93	0.41
4:K:904:ARG:NH2	4:K:1177:PRO:HG2	2.36	0.41
4:L:268:TRP:CD2	4:L:305:LEU:HD22	2.55	0.41
4:L:619:LEU:HD22	4:L:623:GLU:HB3	2.03	0.41
4:L:861:ARG:O	4:L:862:ASN:HB2	2.20	0.41
4:L:1084:MET:HE3	4:L:1106:MET:HG2	2.02	0.41
4:L:1103:LYS:O	4:L:1107:GLN:HG3	2.20	0.41
2:S:63:ALA:O	2:S:67:LEU:HD23	2.20	0.41
2:e:24:ALA:O	2:e:27:ILE:HG12	2.21	0.41
3:M:223:LYS:HE3	3:M:223:LYS:HB3	1.83	0.41
3:N:333:LYS:HB2	3:N:352:LEU:HD21	2.02	0.41
3:N:611:ASP:HB3	4:K:125:LYS:HE2	2.03	0.41
3:O:211:MET:O	3:O:217:SER:HB3	2.21	0.41
3:O:366:TRP:O	3:O:370:GLN:HG2	2.21	0.41
3:P:383:LYS:O	3:P:387:ILE:HG13	2.21	0.41
3:Q:463:ILE:O	3:Q:467:VAL:HG23	2.20	0.41
3:R:579:ILE:O	3:R:583:VAL:HG23	2.21	0.41
3:R:611:ASP:HB3	4:I:125:LYS:HE2	2.03	0.41
3:G:304:LYS:HE2	3:G:304:LYS:HB3	1.81	0.41
4:I:760:ILE:HD11	4:I:825:LYS:HG3	2.02	0.41
4:I:1004:MET:HE1	4:I:1020:TRP:HE1	1.85	0.41
4:I:1041:SER:O	4:I:1045:GLU:HG2	2.20	0.41
4:J:617:LYS:HB3	4:J:617:LYS:HE3	1.88	0.41
4:K:103:LYS:HD3	4:K:123:TYR:CD1	2.56	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
4:L:535:ASP:HB3	4:L:817:THR:HG22	2.02	0.41
4:L:1171:LEU:HA	4:L:1171:LEU:HD23	1.82	0.41
1:B:40:GLU:OE1	3:P:373:ALA:HB1	2.20	0.40
2:T:7:ILE:HB	2:T:8:PRO:HD3	2.02	0.40
2:e:49:ASN:C	2:e:51:ASP:H	2.29	0.40
2:g:76:GLN:HG2	3:H:114:TYR:CD1	2.55	0.40
3:M:557:LEU:HD22	3:M:573:ARG:HH21	1.86	0.40
3:M:664:TYR:HE1	3:M:673:MET:HE2	1.86	0.40
3:Q:565:ILE:HD13	3:Q:597:MET:HE1	2.03	0.40
3:G:75:ILE:O	3:G:78:LEU:HB3	2.21	0.40
3:G:356:GLN:O	3:G:360:GLN:HG2	2.21	0.40
4:I:266:THR:HG21	4:I:1129:LYS:HG3	2.03	0.40
4:J:510:LYS:HA	4:J:510:LYS:HD2	1.88	0.40
4:J:809:GLY:HA3	4:J:856:THR:HA	2.03	0.40
4:K:256:PHE:O	4:K:1089:LYS:NZ	2.48	0.40
4:K:665:TYR:CZ	4:K:667:PRO:HA	2.56	0.40
4:K:757:LEU:HD22	4:K:760:ILE:HG21	2.03	0.40
4:K:879:PHE:HE2	4:L:68:VAL:HG11	1.86	0.40
4:K:968:ILE:HG13	4:K:969:VAL:N	2.36	0.40
4:K:1086:ARG:N	4:K:1087:PRO:HD2	2.36	0.40
4:K:1277:GLU:HG2	4:L:974:TYR:CE1	2.56	0.40
4:L:814:MET:O	4:L:814:MET:HG3	2.21	0.40
2:S:118:ARG:HH22	3:O:68:ASP:HA	1.85	0.40
2:T:73:GLN:HE22	3:N:208:ASP:CG	2.29	0.40
2:g:49:ASN:C	2:g:51:ASP:H	2.29	0.40
3:M:335:GLU:HA	3:M:338:LYS:HE2	2.03	0.40
3:M:396:ASN:HB2	3:M:398:GLU:OE1	2.22	0.40
3:N:313:ILE:HD13	3:N:332:ILE:HD13	2.02	0.40
3:N:588:GLY:O	3:O:539:VAL:HG22	2.21	0.40
3:N:704:ARG:HD3	3:O:368:LYS:HG2	2.02	0.40
3:P:435:MET:SD	3:P:445:MET:HE1	2.61	0.40
3:P:628:VAL:HG23	3:P:629:ASN:HD22	1.86	0.40
3:P:704:ARG:HD3	3:Q:368:LYS:HG3	2.02	0.40
3:Q:120:VAL:O	3:Q:124:ILE:HG13	2.21	0.40
3:Q:133:GLU:OE2	3:R:64:ARG:NH2	2.38	0.40
3:Q:142:ARG:HH11	3:Q:142:ARG:HA	1.86	0.40
3:G:135:MET:HE1	3:G:179:TYR:CD1	2.56	0.40
3:G:197:ASN:N	3:G:197:ASN:HD22	2.19	0.40
3:H:628:VAL:HG22	3:H:634:SER:HB2	2.04	0.40
3:H:706:LYS:HA	3:H:706:LYS:HD2	1.85	0.40
4:I:238:LEU:HD23	4:I:238:LEU:HA	1.81	0.40

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
4:I:450:ASN:HD21	4:J:112:GLU:HG2	1.86	0.40
4:I:584:PRO:HA	4:I:588:THR:HG23	2.03	0.40
4:J:17:GLN:HG3	4:J:1086:ARG:HG2	2.03	0.40
4:J:965:VAL:HA	4:J:968:ILE:HG12	2.03	0.40
4:L:497:PHE:O	4:L:501:ASP:HB2	2.21	0.40
1:b:48:VAL:HG13	3:M:314:ASN:ND2	2.36	0.40
2:g:63:ALA:O	2:g:67:LEU:HD23	2.21	0.40
2:h:7:ILE:HB	2:h:8:PRO:HD3	2.03	0.40
2:h:47:ILE:HA	2:h:52:LEU:HD21	2.03	0.40
3:M:467:VAL:HG22	3:M:493:LEU:HD13	2.04	0.40
3:N:81:GLU:HG2	3:N:82:GLN:N	2.35	0.40
3:N:682:ARG:HH22	3:N:684:ASP:HB2	1.86	0.40
3:O:673:MET:HE2	3:O:673:MET:HB3	2.01	0.40
3:G:610:GLY:C	3:G:658:ASN:HD21	2.29	0.40
3:G:701:GLU:O	3:G:705:GLU:HG2	2.22	0.40
3:H:250:ASP:O	3:H:254:ARG:HB2	2.21	0.40
3:H:594:MET:O	3:H:598:THR:OG1	2.32	0.40
4:I:68:VAL:HG11	4:L:879:PHE:HE2	1.86	0.40
4:I:427:GLY:O	4:I:431:ARG:HG3	2.22	0.40
4:I:689:GLU:OE1	4:I:693:ARG:NH2	2.54	0.40
4:I:996:LEU:CB	4:I:1085:LEU:HD11	2.50	0.40
4:J:355:LEU:HD23	4:J:436:GLU:HB3	2.01	0.40
4:J:659:SER:C	4:J:661:HIS:H	2.30	0.40
4:K:362:VAL:CG2	4:K:446:LEU:HD11	2.51	0.40
4:K:1129:LYS:HA	4:K:1129:LYS:HD3	1.87	0.40
1:D:45:GLY:O	1:D:47:GLU:HG3	2.21	0.40
1:F:40:GLU:CD	3:G:373:ALA:HB2	2.46	0.40
2:S:54:LEU:HD11	3:G:438:PRO:HD3	2.02	0.40
2:g:72:MET:HE3	2:g:72:MET:HB3	1.94	0.40
3:M:195:ILE:HG23	3:M:199:ARG:HH21	1.85	0.40
3:N:658:ASN:OD1	3:N:659:LYS:N	2.54	0.40
3:P:579:ILE:O	3:P:583:VAL:HG23	2.22	0.40
3:P:671:TYR:HE1	3:P:682:ARG:HG3	1.85	0.40
3:R:125:LYS:HZ1	3:R:181:ALA:HB3	1.86	0.40
3:R:463:ILE:O	3:R:467:VAL:HG23	2.21	0.40
3:G:606:TYR:CZ	3:G:639:ARG:HD2	2.56	0.40
3:G:625:MET:HE2	3:G:625:MET:HB3	2.01	0.40
4:I:477:LEU:HD11	4:I:483:LEU:HD11	2.04	0.40
4:I:641:MET:HE3	4:I:656:PHE:HD1	1.86	0.40
4:I:825:LYS:HD3	4:I:829:LEU:HD11	2.04	0.40
4:I:973:LYS:HA	4:L:1274:LYS:NZ	2.36	0.40

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
4:J:383:GLY:C	4:J:385:ALA:H	2.30	0.40
4:J:803:TYR:O	4:J:807:VAL:HG22	2.21	0.40
4:J:1004:MET:HE3	4:J:1084:MET:SD	2.61	0.40
3:M:322:PRO:HB2	3:M:363:MET:CE	2.51	0.40
3:O:601:LEU:O	3:O:605:THR:OG1	2.30	0.40
3:O:610:GLY:C	3:O:658:ASN:HD21	2.30	0.40
3:P:121:MET:HE1	3:P:138:TYR:CE2	2.57	0.40
3:P:206:LEU:HD12	3:P:247:ALA:HB1	2.03	0.40
3:R:467:VAL:CG1	3:R:511:GLN:HG3	2.52	0.40
3:G:78:LEU:CD1	3:G:82:GLN:HG3	2.51	0.40
3:G:639:ARG:NH1	3:H:418:HIS:HB3	2.36	0.40
3:H:102:GLU:O	3:H:106:VAL:HG23	2.21	0.40
3:H:121:MET:O	3:H:125:LYS:HG3	2.22	0.40
3:H:140:HIS:NE2	3:H:144:GLN:OE1	2.54	0.40
4:I:16:PHE:HE2	4:I:40:PHE:HE1	1.69	0.40
4:I:538:ARG:HH11	4:J:783:GLY:HA2	1.85	0.40
4:J:135:GLY:O	4:J:139:MET:HG3	2.21	0.40
4:J:427:GLY:O	4:J:431:ARG:HG3	2.22	0.40
4:K:54:MET:HB2	4:K:56:PHE:CE1	2.57	0.40
4:K:238:LEU:HB3	4:K:239:SER:H	1.78	0.40
4:K:357:ASN:O	4:K:361:VAL:HG23	2.21	0.40
4:K:530:ARG:O	4:K:534:ILE:HG13	2.22	0.40
4:K:620:THR:HG1	4:K:623:GLU:HG3	1.87	0.40
4:K:696:MET:HE2	4:K:728:GLU:HG2	2.04	0.40
4:K:820:THR:OG1	4:K:823:GLU:HG3	2.21	0.40
4:K:913:LEU:HD12	4:K:913:LEU:HA	1.90	0.40
4:L:363:GLY:O	4:L:443:SER:HB3	2.22	0.40
4:L:485:ALA:O	4:L:486:SER:OG	2.33	0.40
4:L:927:GLU:O	4:L:931:LEU:HG	2.21	0.40
4:L:1234:SER:O	4:L:1238:MET:HG2	2.21	0.40
4:L:1241:MET:O	4:L:1243:SER:N	2.54	0.40

There are no symmetry-related clashes.

## 5.3 Torsion angles ⓘ

### 5.3.1 Protein backbone ⓘ

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

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

Mol	Chain	Analysed	Favoured	Allowed	Outliers	Percentiles	
1	A	15/88 (17%)	15 (100%)	0	0	100	100
1	B	15/88 (17%)	15 (100%)	0	0	100	100
1	C	15/88 (17%)	15 (100%)	0	0	100	100
1	D	14/88 (16%)	14 (100%)	0	0	100	100
1	E	14/88 (16%)	14 (100%)	0	0	100	100
1	F	14/88 (16%)	14 (100%)	0	0	100	100
1	a	15/88 (17%)	15 (100%)	0	0	100	100
1	b	14/88 (16%)	14 (100%)	0	0	100	100
2	S	108/196 (55%)	106 (98%)	2 (2%)	0	100	100
2	T	107/196 (55%)	104 (97%)	3 (3%)	0	100	100
2	c	108/196 (55%)	104 (96%)	4 (4%)	0	100	100
2	d	107/196 (55%)	105 (98%)	2 (2%)	0	100	100
2	e	108/196 (55%)	105 (97%)	3 (3%)	0	100	100
2	f	107/196 (55%)	106 (99%)	1 (1%)	0	100	100
2	g	108/196 (55%)	106 (98%)	2 (2%)	0	100	100
2	h	107/196 (55%)	105 (98%)	2 (2%)	0	100	100
3	G	669/747 (90%)	642 (96%)	25 (4%)	2 (0%)	37	70
3	H	669/747 (90%)	636 (95%)	31 (5%)	2 (0%)	37	70
3	M	669/747 (90%)	642 (96%)	27 (4%)	0	100	100
3	N	669/747 (90%)	642 (96%)	24 (4%)	3 (0%)	30	66
3	O	669/747 (90%)	634 (95%)	34 (5%)	1 (0%)	48	81
3	P	669/747 (90%)	641 (96%)	25 (4%)	3 (0%)	30	66
3	Q	669/747 (90%)	640 (96%)	29 (4%)	0	100	100
3	R	669/747 (90%)	634 (95%)	31 (5%)	4 (1%)	22	57
4	I	1217/1318 (92%)	1118 (92%)	86 (7%)	13 (1%)	12	44
4	J	1217/1318 (92%)	1120 (92%)	85 (7%)	12 (1%)	13	46
4	K	1217/1318 (92%)	1112 (91%)	95 (8%)	10 (1%)	16	51
4	L	1217/1318 (92%)	1119 (92%)	84 (7%)	14 (1%)	11	41
All	All	11196/13520 (83%)	10537 (94%)	595 (5%)	64 (1%)	24	57

All (64) Ramachandran outliers are listed below:

Mol	Chain	Res	Type
4	I	61	ALA
4	I	63	ALA
4	I	504	LYS
4	I	1060	LYS
4	J	61	ALA
4	J	242	VAL
4	J	504	LYS
4	J	1060	LYS
4	K	61	ALA
4	K	242	VAL
4	K	504	LYS
4	K	524	SER
4	K	1060	LYS
4	L	61	ALA
4	L	242	VAL
4	L	504	LYS
4	L	1060	LYS
3	R	237	ASP
4	I	362	VAL
4	I	796	MET
4	I	860	ARG
4	J	362	VAL
4	J	526	ASP
4	J	796	MET
4	J	860	ARG
4	K	796	MET
4	K	860	ARG
4	L	362	VAL
4	L	796	MET
4	L	860	ARG
3	N	567	ARG
3	P	567	ARG
3	R	567	ARG
3	G	235	PRO
4	J	351	LYS
4	L	64	LEU
4	L	236	ALA
4	L	457	GLU
3	O	537	LEU
3	P	236	SER
3	R	236	SER
3	G	537	LEU
3	H	567	ARG

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Mol	Chain	Res	Type
4	I	457	GLU
4	I	982	ARG
4	J	457	GLU
4	J	982	ARG
4	K	982	ARG
4	L	65	GLY
4	L	982	ARG
3	N	236	SER
3	R	568	MET
4	I	66	LEU
4	I	859	ALA
4	J	859	ALA
4	K	457	GLU
4	L	63	ALA
4	L	859	ALA
3	P	568	MET
4	I	242	VAL
4	K	859	ALA
3	N	568	MET
3	H	568	MET
4	I	1033	PRO

### 5.3.2 Protein sidechains ⓘ

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

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

Mol	Chain	Analysed	Rotameric	Outliers	Percentiles	
1	A	13/73 (18%)	13 (100%)	0	100	100
1	B	13/73 (18%)	13 (100%)	0	100	100
1	C	13/73 (18%)	13 (100%)	0	100	100
1	D	13/73 (18%)	13 (100%)	0	100	100
1	E	13/73 (18%)	13 (100%)	0	100	100
1	F	13/73 (18%)	13 (100%)	0	100	100
1	a	13/73 (18%)	13 (100%)	0	100	100
1	b	13/73 (18%)	13 (100%)	0	100	100

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Mol	Chain	Analysed	Rotameric	Outliers	Percentiles	
2	S	83/149 (56%)	83 (100%)	0	100	100
2	T	82/149 (55%)	82 (100%)	0	100	100
2	c	83/149 (56%)	83 (100%)	0	100	100
2	d	82/149 (55%)	82 (100%)	0	100	100
2	e	83/149 (56%)	83 (100%)	0	100	100
2	f	82/149 (55%)	81 (99%)	1 (1%)	67	86
2	g	83/149 (56%)	83 (100%)	0	100	100
2	h	82/149 (55%)	82 (100%)	0	100	100
3	G	569/624 (91%)	565 (99%)	4 (1%)	81	91
3	H	569/624 (91%)	568 (100%)	1 (0%)	92	97
3	M	569/624 (91%)	567 (100%)	2 (0%)	89	95
3	N	569/624 (91%)	567 (100%)	2 (0%)	89	95
3	O	569/624 (91%)	569 (100%)	0	100	100
3	P	569/624 (91%)	568 (100%)	1 (0%)	92	97
3	Q	569/624 (91%)	567 (100%)	2 (0%)	89	95
3	R	569/624 (91%)	568 (100%)	1 (0%)	92	97
4	I	979/1059 (92%)	975 (100%)	4 (0%)	89	95
4	J	979/1059 (92%)	972 (99%)	7 (1%)	81	91
4	K	979/1059 (92%)	974 (100%)	5 (0%)	86	94
4	L	979/1059 (92%)	975 (100%)	4 (0%)	89	95
All	All	9232/11004 (84%)	9198 (100%)	34 (0%)	88	95

All (34) residues with a non-rotameric sidechain are listed below:

Mol	Chain	Res	Type
2	f	73	GLN
3	M	539	VAL
3	M	660	GLN
3	N	119	ASP
3	N	613	VAL
3	P	111	ASN
3	Q	539	VAL
3	Q	660	GLN
3	R	111	ASN
3	G	340	GLN

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Mol	Chain	Res	Type
3	G	537	LEU
3	G	539	VAL
3	G	660	GLN
3	H	239	GLN
4	I	68	VAL
4	I	69	THR
4	I	362	VAL
4	I	990	ASN
4	J	75	ASP
4	J	349	THR
4	J	351	LYS
4	J	522	LEU
4	J	1031	VAL
4	J	1187	ARG
4	J	1191	LEU
4	K	66	LEU
4	K	68	VAL
4	K	69	THR
4	K	349	THR
4	K	1191	LEU
4	L	240	ASN
4	L	362	VAL
4	L	522	LEU
4	L	1021	GLU

Sometimes sidechains can be flipped to improve hydrogen bonding and reduce clashes. All (134) such sidechains are listed below:

Mol	Chain	Res	Type
2	T	44	GLN
2	T	48	GLN
2	T	73	GLN
2	c	30	GLN
2	c	55	GLN
2	c	76	GLN
2	e	44	GLN
2	f	30	GLN
2	f	44	GLN
2	f	71	ASN
2	g	71	ASN
2	g	76	GLN
3	M	88	ASN
3	M	140	HIS

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Mol	Chain	Res	Type
3	M	164	GLN
3	M	191	GLN
3	M	197	ASN
3	M	242	GLN
3	M	314	ASN
3	M	364	ASN
3	M	451	GLN
3	M	529	GLN
3	M	658	ASN
3	N	82	GLN
3	N	144	GLN
3	N	164	GLN
3	N	191	GLN
3	N	295	GLN
3	N	306	ASN
3	N	319	GLN
3	N	344	GLN
3	N	370	GLN
3	N	524	GLN
3	N	589	ASN
3	N	637	GLN
3	O	88	ASN
3	O	111	ASN
3	O	140	HIS
3	O	182	HIS
3	O	286	GLN
3	O	314	ASN
3	O	318	ASN
3	O	319	GLN
3	O	382	ASN
3	O	529	GLN
3	O	589	ASN
3	O	658	ASN
3	O	695	ASN
3	P	82	GLN
3	P	182	HIS
3	P	295	GLN
3	P	318	ASN
3	P	356	GLN
3	P	358	GLN
3	P	370	GLN
3	P	382	ASN

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Mol	Chain	Res	Type
3	P	524	GLN
3	Q	140	HIS
3	Q	144	GLN
3	Q	182	HIS
3	Q	191	GLN
3	Q	314	ASN
3	Q	426	ASN
3	Q	529	GLN
3	Q	627	GLN
3	Q	637	GLN
3	Q	658	ASN
3	Q	695	ASN
3	R	182	HIS
3	R	382	ASN
3	R	423	ASN
3	R	545	GLN
3	R	678	GLN
3	G	82	GLN
3	G	88	ASN
3	G	144	GLN
3	G	197	ASN
3	G	314	ASN
3	G	340	GLN
3	G	529	GLN
3	G	627	GLN
3	G	637	GLN
3	G	658	ASN
3	H	111	ASN
3	H	182	HIS
3	H	239	GLN
3	H	308	GLN
3	H	330	GLN
3	H	358	GLN
3	H	392	GLN
3	H	423	ASN
3	H	451	GLN
3	H	511	GLN
3	H	524	GLN
4	I	84	ASN
4	I	269	ASN
4	I	287	ASN
4	I	292	ASN

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Mol	Chain	Res	Type
4	I	423	ASN
4	I	440	ASN
4	I	450	ASN
4	I	717	HIS
4	I	932	HIS
4	I	953	GLN
4	I	977	GLN
4	I	990	ASN
4	I	1294	ASN
4	J	17	GLN
4	J	109	ASN
4	J	110	GLN
4	J	292	ASN
4	J	357	ASN
4	J	442	ASN
4	J	460	HIS
4	J	567	GLN
4	J	846	HIS
4	K	17	GLN
4	K	292	ASN
4	K	442	ASN
4	K	567	GLN
4	K	746	ASN
4	K	953	GLN
4	K	1061	GLN
4	L	17	GLN
4	L	109	ASN
4	L	240	ASN
4	L	292	ASN
4	L	450	ASN
4	L	567	GLN
4	L	846	HIS
4	L	875	ASN
4	L	977	GLN
4	L	994	ASN
4	L	1061	GLN

### 5.3.3 RNA ⓘ

There are no RNA molecules in this entry.

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

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

## 5.5 Carbohydrates [i](#)

There are no oligosaccharides in this entry.

## 5.6 Ligand geometry [i](#)

There are no ligands in this entry.

## 5.7 Other polymers [i](#)

There are no such residues in this entry.

## 5.8 Polymer linkage issues [i](#)

There are no chain breaks in this entry.