



Full wwPDB EM Validation Report ⓘ

Mar 27, 2025 – 12:28 pm GMT

PDB ID : 6ZTZ
EMDB ID : EMD-22166
Title : Assembly intermediates of orthoreovirus captured in the cell
Authors : Sutton, G.C.; Stuart, D.I.
Deposited on : 2020-07-20
Resolution : 6.50 Å(reported)

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

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A user guide is available at

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

with specific help available everywhere you see the ⓘ symbol.

The types of validation reports are described at

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

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

EMDB validation analysis : 0.0.1.dev117
MolProbity : 4.02b-467
Percentile statistics : 20231227.v01 (using entries in the PDB archive December 27th 2023)
MapQ : 1.9.13
Ideal geometry (proteins) : Engh & Huber (2001)
Ideal geometry (DNA, RNA) : Parkinson et al. (1996)
Validation Pipeline (wwPDB-VP) : 2.41.5

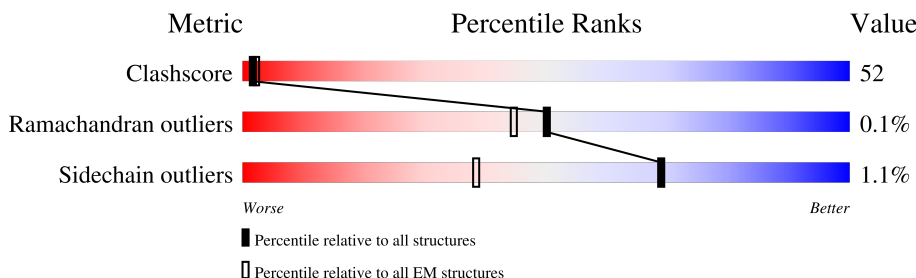
1 Overall quality at a glance

The following experimental techniques were used to determine the structure:

ELECTRON MICROSCOPY

The reported resolution of this entry is 6.50 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 ≥ 3 , 2, 1 and 0 types of geometric quality criteria respectively. A grey segment represents the fraction of residues that are not modelled. The numeric value for each fraction is indicated below the corresponding segment, with a dot representing fractions $\leq 5\%$. The upper red bar (where present) indicates the fraction of residues that have poor fit to the EM map (all-atom inclusion $< 40\%$). The numeric value is given above the bar.

Mol	Chain	Length	Quality of chain
1	B	1035	<div> <div>100%</div> <div> <div>30%</div> <div>69%</div> </div> </div>
2	C	1008	<div> <div>95%</div> <div> <div>26%</div> <div>73%</div> </div> </div>
3	D	417	<div> <div>96%</div> <div> <div>27%</div> <div>71%</div> </div> </div>
3	P	417	<div> <div>84%</div> <div> <div>25%</div> <div>73%</div> </div> </div>
4	K	641	<div> <div>89%</div> <div> <div>28%</div> <div>71%</div> </div> </div>
4	L	641	<div> <div>91%</div> <div> <div>32%</div> <div>68%</div> </div> </div>
4	M	641	<div> <div>92%</div> <div> <div>32%</div> <div>67%</div> </div> </div>
5	O	1284	<div> <div>91%</div> <div> <div>26%</div> <div>73%</div> </div> </div>

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Mol	Chain	Length	Quality of chain
6	X	365	<div><div></div><div>92%</div><div>25%</div><div>74%</div><div></div></div>
6	Y	365	<div><div></div><div>94%</div><div>30%</div><div>69%</div><div></div></div>
6	Z	365	<div><div></div><div>92%</div><div>27%</div><div>72%</div><div></div></div>

2 Entry composition [i](#)

There are 6 unique types of molecules in this entry. The entry contains 56150 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 Inner capsid protein lambda-1.

Mol	Chain	Residues	Atoms					AltConf	Trace
1	B	1035	Total	C	N	O	S	0	0
			8171	5222	1380	1519	50		

- Molecule 2 is a protein called Inner capsid protein lambda-1.

Mol	Chain	Residues	Atoms					AltConf	Trace
2	C	1008	Total	C	N	O	S	0	0
			7958	5091	1342	1475	50		

There are 8 discrepancies between the modelled and reference sequences:

Chain	Residue	Modelled	Actual	Comment	Reference
C	?	-	VAL	deletion	UNP Q9WAB2
C	?	-	SER	deletion	UNP Q9WAB2
C	?	-	GLU	deletion	UNP Q9WAB2
C	?	-	SER	deletion	UNP Q9WAB2
C	?	-	THR	deletion	UNP Q9WAB2
C	?	-	THR	deletion	UNP Q9WAB2
C	?	-	GLN	deletion	UNP Q9WAB2
C	?	-	THR	deletion	UNP Q9WAB2

- Molecule 3 is a protein called Inner capsid protein sigma-2.

Mol	Chain	Residues	Atoms					AltConf	Trace
3	D	417	Total	C	N	O	S	0	0
			3313	2092	600	604	17		
3	P	417	Total	C	N	O	S	0	0
			3313	2092	600	604	17		

- Molecule 4 is a protein called Outer capsid protein mu-1.

Mol	Chain	Residues	Atoms					AltConf	Trace
4	K	641	Total 4871	C 3091	N 807	O 954	S 19	0	0
4	L	641	Total 4871	C 3091	N 807	O 954	S 19	0	0
4	M	641	Total 4871	C 3091	N 807	O 954	S 19	0	0

There are 81 discrepancies between the modelled and reference sequences:

Chain	Residue	Modelled	Actual	Comment	Reference
K	?	-	PRO	deletion	UNP P11077
K	?	-	GLU	deletion	UNP P11077
K	?	-	THR	deletion	UNP P11077
K	?	-	ALA	deletion	UNP P11077
K	?	-	ILE	deletion	UNP P11077
K	?	-	ILE	deletion	UNP P11077
K	?	-	ASN	deletion	UNP P11077
K	?	-	THR	deletion	UNP P11077
K	?	-	ASP	deletion	UNP P11077
K	?	-	ASN	deletion	UNP P11077
K	?	-	SER	deletion	UNP P11077
K	?	-	SER	deletion	UNP P11077
K	?	-	GLY	deletion	UNP P11077
K	?	-	ALA	deletion	UNP P11077
K	?	-	VAL	deletion	UNP P11077
K	?	-	PRO	deletion	UNP P11077
K	?	-	SER	deletion	UNP P11077
K	?	-	GLU	deletion	UNP P11077
K	?	-	SER	deletion	UNP P11077
K	?	-	ALA	deletion	UNP P11077
K	?	-	LEU	deletion	UNP P11077
K	?	-	VAL	deletion	UNP P11077
K	?	-	PRO	deletion	UNP P11077
K	?	-	TYR	deletion	UNP P11077
K	?	-	ASN	deletion	UNP P11077
K	344	LEU	PRO	conflict	UNP P11077
K	359	PHE	LEU	conflict	UNP P11077
L	?	-	PRO	deletion	UNP P11077
L	?	-	GLU	deletion	UNP P11077
L	?	-	THR	deletion	UNP P11077
L	?	-	ALA	deletion	UNP P11077
L	?	-	ILE	deletion	UNP P11077
L	?	-	ILE	deletion	UNP P11077

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Chain	Residue	Modelled	Actual	Comment	Reference
L	?	-	ASN	deletion	UNP P11077
L	?	-	THR	deletion	UNP P11077
L	?	-	ASP	deletion	UNP P11077
L	?	-	ASN	deletion	UNP P11077
L	?	-	SER	deletion	UNP P11077
L	?	-	SER	deletion	UNP P11077
L	?	-	GLY	deletion	UNP P11077
L	?	-	ALA	deletion	UNP P11077
L	?	-	VAL	deletion	UNP P11077
L	?	-	PRO	deletion	UNP P11077
L	?	-	SER	deletion	UNP P11077
L	?	-	GLU	deletion	UNP P11077
L	?	-	SER	deletion	UNP P11077
L	?	-	ALA	deletion	UNP P11077
L	?	-	LEU	deletion	UNP P11077
L	?	-	VAL	deletion	UNP P11077
L	?	-	PRO	deletion	UNP P11077
L	?	-	TYR	deletion	UNP P11077
L	?	-	ASN	deletion	UNP P11077
L	344	LEU	PRO	conflict	UNP P11077
L	359	PHE	LEU	conflict	UNP P11077
M	?	-	PRO	deletion	UNP P11077
M	?	-	GLU	deletion	UNP P11077
M	?	-	THR	deletion	UNP P11077
M	?	-	ALA	deletion	UNP P11077
M	?	-	ILE	deletion	UNP P11077
M	?	-	ILE	deletion	UNP P11077
M	?	-	ASN	deletion	UNP P11077
M	?	-	THR	deletion	UNP P11077
M	?	-	ASP	deletion	UNP P11077
M	?	-	ASN	deletion	UNP P11077
M	?	-	SER	deletion	UNP P11077
M	?	-	SER	deletion	UNP P11077
M	?	-	GLY	deletion	UNP P11077
M	?	-	ALA	deletion	UNP P11077
M	?	-	VAL	deletion	UNP P11077
M	?	-	PRO	deletion	UNP P11077
M	?	-	SER	deletion	UNP P11077
M	?	-	GLU	deletion	UNP P11077
M	?	-	SER	deletion	UNP P11077
M	?	-	ALA	deletion	UNP P11077
M	?	-	LEU	deletion	UNP P11077

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Chain	Residue	Modelled	Actual	Comment	Reference
M	?	-	VAL	deletion	UNP P11077
M	?	-	PRO	deletion	UNP P11077
M	?	-	TYR	deletion	UNP P11077
M	?	-	ASN	deletion	UNP P11077
M	344	LEU	PRO	conflict	UNP P11077
M	359	PHE	LEU	conflict	UNP P11077

- Molecule 5 is a protein called Outer capsid protein lambda-2.

Mol	Chain	Residues	Atoms					AltConf	Trace
5	O	1284	Total	C	N	O	S	0	0
			10127	6468	1700	1917	42		

There are 4 discrepancies between the modelled and reference sequences:

Chain	Residue	Modelled	Actual	Comment	Reference
O	?	-	SER	deletion	UNP P11079
O	?	-	ALA	deletion	UNP P11079
O	?	-	SER	deletion	UNP P11079
O	?	-	GLY	deletion	UNP P11079

- Molecule 6 is a protein called Outer capsid protein sigma-3.

Mol	Chain	Residues	Atoms					AltConf	Trace
6	X	365	Total	C	N	O	S	0	0
			2885	1818	508	531	28		
6	Y	365	Total	C	N	O	S	0	0
			2885	1818	508	531	28		
6	Z	365	Total	C	N	O	S	0	0
			2885	1818	508	531	28		

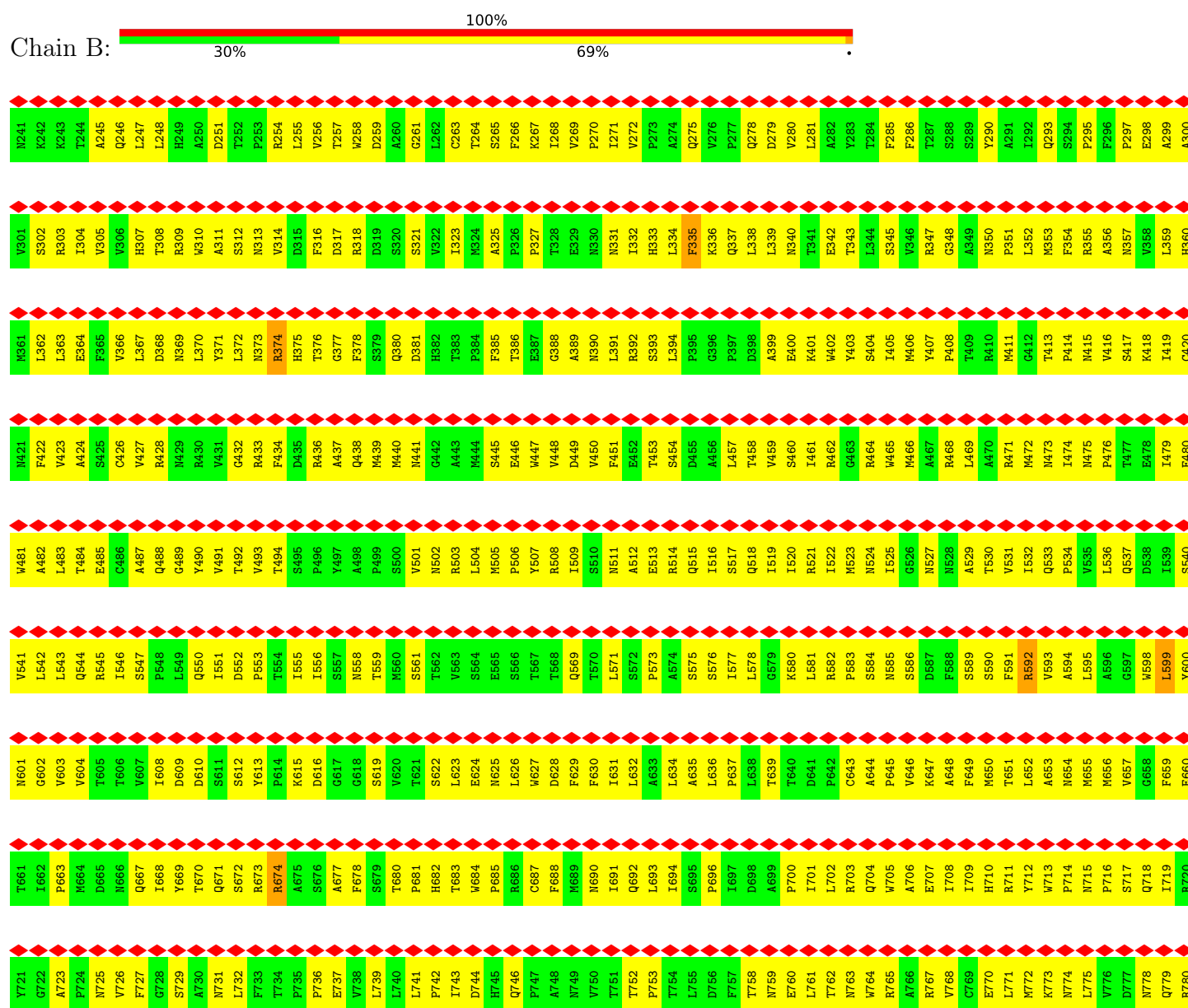
There are 6 discrepancies between the modelled and reference sequences:

Chain	Residue	Modelled	Actual	Comment	Reference
X	104	CYS	ALA	conflict	UNP P07939
X	325	ASN	ASP	conflict	UNP P07939
Y	104	CYS	ALA	conflict	UNP P07939
Y	325	ASN	ASP	conflict	UNP P07939
Z	104	CYS	ALA	conflict	UNP P07939
Z	325	ASN	ASP	conflict	UNP P07939

3 Residue-property plots

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

- Molecule 1: Inner capsid protein lambda-1



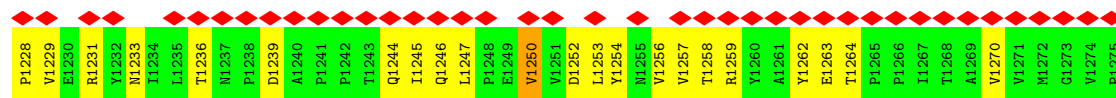
A1261	S1201	M1141	C1081	A1021	T961	Y901	D841	Y781
Y1262	T1202	G1142	R1082	D1022	A962	A902	R842	Q782
E1263	E1203	A1143	I1083	C1023	A963	A903	P843	P783
T1264	Y1204	Y1144	S1084	S1024	T964	A904	P844	G784
P1265	N1205	P1145	F1085	Q1025	F965	I905	T845	W785
P1266	D1206	Y1146	G1086	L1026	A966	Y906	M846	T786
T1267	R1207	M1147	M1087	T1027	E967	P907	V847	Q787
A1268	S1208	L1148	N1088	A1028	W968	M908	G848	S788
A1269	L1209	H1149	G1089	E1029	V969	Y909	V849	L789
V1270	F1210	Y1150	A1090	V1030	N970	A910	T850	V790
M1271	G1211	Y1151	A1091	F1031	T971	D911	R851	S791
M1272	T1212	D1152	P1092	M1032	S972	T912	Q852	S792
G1273	N1213	P1153	M1093	H1033	M973	E913	S853	M793
V1274	S1214	R1154	I1094	E1034	K974	V914	R854	R794
P1275	S1215	Q1155	R1095	Y1035	T975	F915	G795	G795
	S1216	Y1156	D1096	M1036	A976	S916	T856	T796
	P1217	A1157	E1097	L1037	F977	N917	I857	L797
	Q1218	N1158	T1098	F1038	D978	L918	T858	D798
	T1219	A1159	G1099	G1039	L979	Q919	Q859	K799
	I1220	W1160	M1100	I1040	S980	R920	P860	L800
	A1221	M1161	M1101	A1041	D981	D921	A861	K801
	G1222	L1162	V1102	R1042	M982	M922	L862	L802
	P1223	T1163	P1103	G1043	L983	I923	S863	I803
	D1224	S1164	F1104	D1044	L984	T924	L864	K804
	K1225	A1165	E1105	I1045	E985	C925	S865	S805
	H1226	W1166	G1106	I1046	P986	E926	T866	M806
	T1227	L1167	N1107	I1047	L987	A927	T867	T807
	P1228	E1168	W1108	G1048	L988	V928	N868	P808
	V1229	E1169	I1109	R1049	S989	Q929	M809	M809
	E1230	I1170	F1110	V1050	G990	T930	T870	Y810
	R1231	T1171	P1111	Q1051	D991	L931	L871	L811
	Y1232	P1172	L1112	S1052	P992	V932	Q872	Q812
	N1233	T1173	A1113	T1053	R993	T933	V873	Q813
	I1234	S1174	L1114	H1054	M994	L934	P874	L914
	L1235	I1175	W1115	L1055	T995	V935	L875	A815
	T1236	P1176	Q1116	V1056	Q996	A936	A876	P816
	N1237	S1177	M1117	S1057	L997	Q937	L877	V817
	P1238	V1178	N1118	P1058	A998	I938	D878	E918
	D1239	P1179	T1119	L1059	T999	S939	A879	L819
	A1240	F1180	R1120	A1060	Q1000	E940	R880	A820
	P1241	M1181	Y1121	P1061	Y1001	T941	A881	V821
	P1242	V1182	F1122	P1062	Q1002	Q942	I882	I822
	T1243	P1183	M1123	P1063	Y1003	T943	T883	A823
	Q1244	I1184	Q1124	D1064	Y1004	P944	V884	P824
	I1245	S1185	Q1125	L1065	M1005	V845	A885	M825
	Q1246	S1186	F1126	V1066	G1006	D946	L886	L826
	L1247	D1187	D1127	F1067	T1007	R947	L887	P827
	P1248	H1188	A1128	D1068	T1008	Y948	S988	F828
	E1249	D1189	W1129	R1069	F1009	L949	G889	P829
	V1250	I1190	T1130	D1070	M1010	D950	K890	P830
	V1251	S1191	K1131	T1071	V1011	W951	Y891	F831
	D1252	S1192	T1132	P1072	I1012	I952	P892	Q832
	L1253	A1193	G1133	G1073	P1013	P953	P893	V833
	Y1254	P1194	E1134	V1074	E1014	S954	D894	P834
	N1255	A1195	L1135	H1075	M1015	L955	L895	Y835
	V1256	V1196	R1136	I1076	P1016	R956	V896	V836
	V1257	Q1197	T1137	F1077	G1017	A957	T897	R837
	T1258	Y1198	R1138	G1078	S1018	S958	N898	L838
	R1259	T1199	I1139	R1079	V1019	A959	V899	D839
	V1260	T1200	F1140	D1080	T1020	A960	W900	R840

• Molecule 2: Inner capsid protein lambda-1

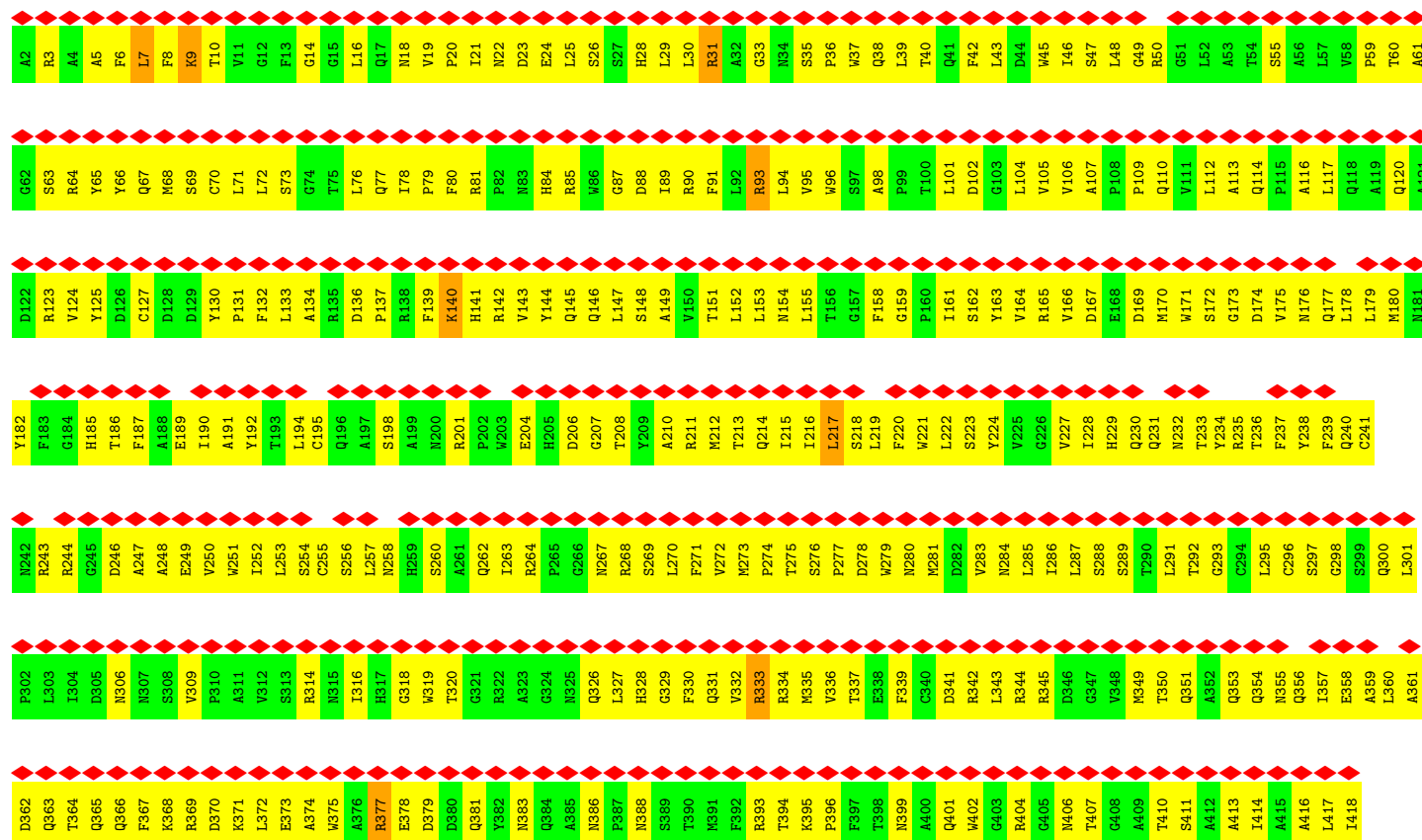


A260	G261	L262	C263	T264	S265	F266	K267	L268	A269	V270	V271	G272	M273	V274	P275	A276	Q277	F278	K279	L280	A281	G282	L283	T284	F285	F286	T287	S288	S289	V290	A291	L292	Q293	S294	P295	F296	P297	E298	A299	A300	V301	S302	R303	I304	V305	V306	K307	T308	R309	W310	A311	S312	N313	V314	D315	F316	D317	R318	D319
S320	S321	V322	I323	M324	A325	P326	P327	T328	E329	N330	N331	I332	H333	L334	F335	K336	Q337	L338	I339	N340	T341	E342	T343	L344	S345	V346	R347	G348	A349	N350	P351	L352	K353	F354	R355	A356	N357	V358	L359	R360	N361	L362	L363	E364	F365	V366	L367	D368	N369	L370	V371	L372	N373	R374	H375	T376	G377	F378	S379
Q380	D381	H382	T383	P384	F385	T386	E387	G388	A389	N390	L391	R392	S393	L394	F395	G396	P397	D398	A399	E400	K401	W402	Y403	L404	I405	M406	Y407	P408	T409	R410	M411	G412	T413	P414	M415	V416	S417	K418	I419	C420	M421	F422	V423	A424	S425	C426	V427	R428	M429	R430	V431	G432	R433	F434	D435	R436	A437	R438	A439

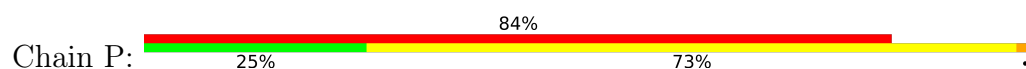
E1168	G1048	L988	Y928	N869	P808	A748	F888	D628	M560	S500	M440
E1169	R1049	S989	Q929	T869	M809	N749	M889	F829	S561	V501	N441
I1170	V1050	G990	T930	T870	Y810	V750	N690	F830	T562	N502	G442
T1171	Q1051	D991	L931	V871	L811	T751	I691	I831	L571	R503	A443
L1172	S1052	P992	V932	G872	Q812	T752	Q892	L632	S572	L504	M444
T1173	T1053	R993	T933	V873	Q813	P753	L693	A833	S573	M505	S445
S1174	H1054	M994	L934	P874	L814	T754	S895	L634	A574	P506	E446
I1175	L1055	T995	V935	L875	A815	L755	P896	A635	S575	Y507	E447
P1176	W1056	Q996	A936	A876	P816	D756	I897	P637	S576	L508	V448
S1177	S1057	L997	Q937	L877	V817	T757	D898	L638	I577	I509	D449
V1178	P1058	A998	T938	D878	B818	T758	A699	L639	L578	S510	V450
P1179	L1059	I999	S939	A879	L819	N759	P700	T639	G579	N511	F451
F1180	A1060	Q1000	E940	R880	A820	E760	I701	T640	K580	A512	E452
M1181	P1061	Y1001	T941	A881	V821	L761	L702	D641	L581	E513	T453
V1182	P1062	Q1002	Q942	T882	T822	T762	Q704	P642	R582	R514	S454
P1183	P1063	Q1003	Y943	T883	A823	N763	W705	C643	P583	Q515	D455
L1184	D1064	Y1004	P944	V884	P824	A764	A706	A644	S584	I516	A456
S1185	L1065	M1005	V945	A885	M825	R765	E707	P645	N585	S517	L457
L1186	V1066	G1006	D946	L886	L826	A766	I708	V646	S586	Q518	T458
D1187	F1067	R1007	R947	L887	P827	R767	I709	K647	D587	I519	V459
H1188	D1068	T1008	Y948	S888	P828	V768	I710	A648	F588	I520	S460
A1189	R1069	F1009	L949	G889	P829	C769	R711	F649	S589	R521	I461
I1190	D1070	N1010	D950	K890	P830	E770	Y712	M650	I522	I522	R462
I1191	T1071	V1011	D951	Y891	P831	L771	W713	M651	S590	I523	G463
K1192	P1072	I1012	P952	P892	Q832	M772	P714	L652	F591	N524	R464
S1193	G1073	P1013	P953	P893	V833	K773	W715	A653	R592	I525	W465
A1194	V1074	E1014	S954	D894	P834	N774	P716	N654	V593	G526	M466
H1075	H1075	M1015	L955	L895	Y835	L775	S717	M655	A594	G527	A467
I1076	I1076	P1016	L956	L896	V836	V776	Q718	M656	L595	N527	R468
F1077	G1077	G1017	R957	T897	V837	D777	I719	M657	A596	N528	L469
G1078	G1078	S1018	A957	V897	L838	N778	R720	G657	G597	A529	A470
R1079	R1079	N1019	S958	N898	D839	Y721	W598	G658	W598	T530	R471
I1139	D1080	I1020	A959	V899	D840	Q779	Y721	F659	L599	V531	M472
C1081	C1081	A1021	A960	Y900	R842	R780	A723	T661	Y600	I532	N473
T961	T961	T961	A962	Y901	V843	Q782	W725	I662	G602	Q533	I474
A962	A962	C1023	A963	D903	V844	P783	W726	P663	G603	P534	N475
S1084	S1084	V1024	T964	A904	T845	G784	F727	M664	V603	V535	P476
F1085	F1085	Q1025	F965	I905	M846	W785	G728	D665	L536	L536	T477
G1086	G1086	L1026	A966	Y906	V847	T786	S729	M666	T606	Q537	E478
M1087	M1087	T1027	E967	P907	G848	Q787	A730	Q667	T606	D538	I479
N1088	N1088	A1028	Y968	N908	V849	T788	N731	I668	V607	I539	E480
G1089	G1089	E1029	Y969	Y909	T850	L789	L732	Y669	I608	S540	W481
A1090	A1090	V1030	V969	T850	R851	V790	F733	T670	V541	L541	A482
A1091	A1091	F1031	N970	A910	Q852	S791	T734	Q671	D610	L542	L483
P1092	P1092	M1032	T971	D911	S853	S792	P735	S611	S612	L543	T484
C1093	C1093	H1033	S972	T912	R854	W793	E737	S672	S612	Q644	E485
M1094	M1094	E1034	N973	E913	T855	R854	E737	S673	S612	R545	C486
I1095	I1095	Y1035	K974	V914	D855	R794	V738	R674	R546	I546	A487
D1096	D1096	N1036	T975	F915	T856	G795	L739	A675	K615	S547	Q488
E1097	E1097	A976	A976	S916	T857	L796	L740	S676	D616	P548	G489
T1098	T1098	F1037	F977	N917	T858	L797	L741	S677	L549	Y490	Y490
G1099	G1099	L1038	D978	L918	Q859	D798	P742	G618	Q550	V491	V491
M1100	M1100	I1040	L979	Q919	P860	K799	I743	S679	S619	T492	T492
V1101	V1101	A1041	S980	R920	A861	L800	D744	T880	V620	D552	T494
M1102	M1102	R1042	D981	D921	L862	K801	H745	P681	P553	P553	S495
P1103	P1103	G1043	N982	N922	S863	L802	Q746	S822	T554	T554	P496
F1104	F1104	D1044	L983	T923	L864	R803	P747	T883	L623	I555	I555
E1105	E1105	I1045	L984	T924	S865	K804		W884	E624	I556	A498
G1106	G1106	I1046	E985	C925	T866	N806		P885	N625	S557	P499
N1107	N1107	L1047	P986	E926	T867	T807		G886	L626	N558	
			L987	A927				C887	W627		



• Molecule 3: Inner capsid protein sigma-2

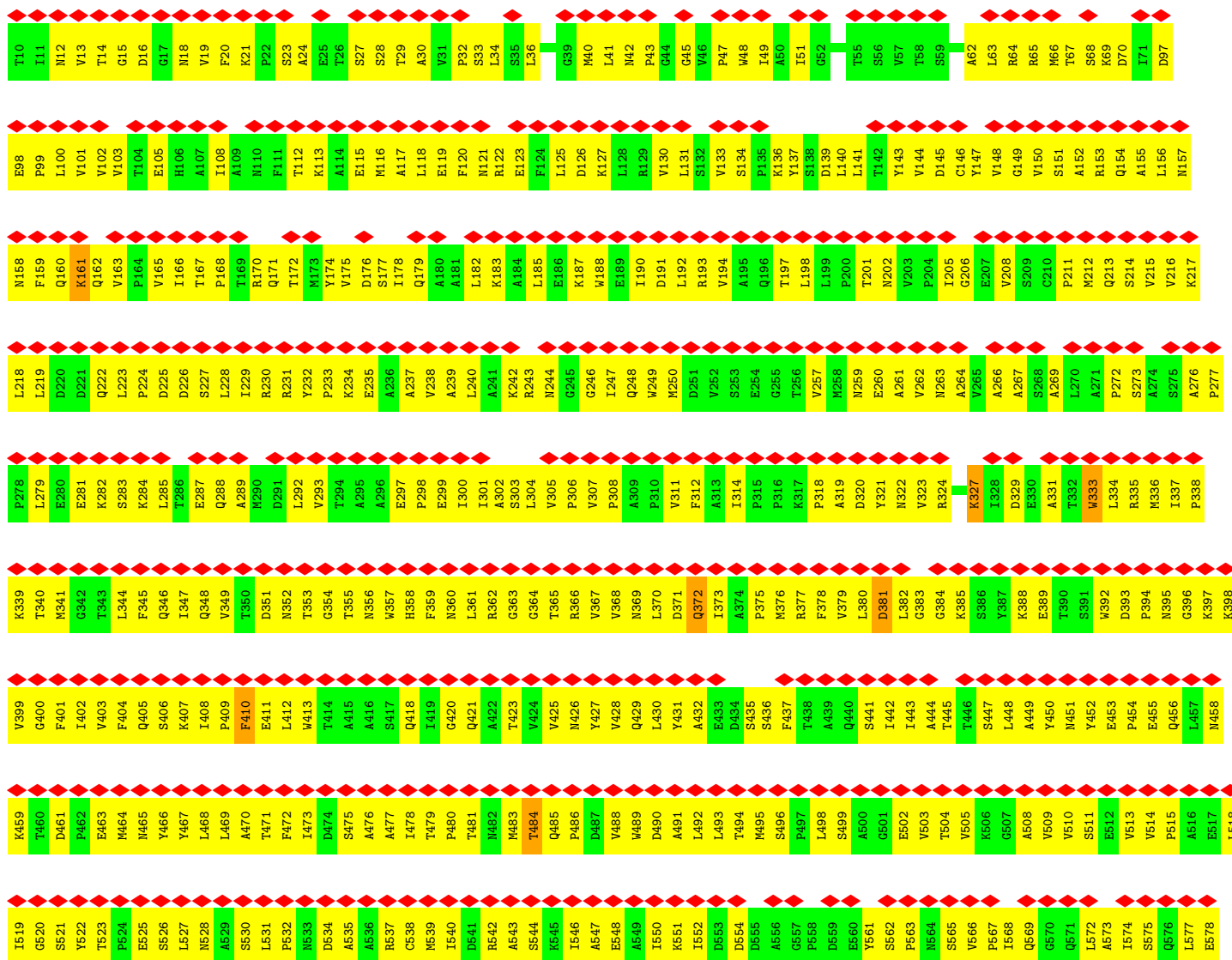
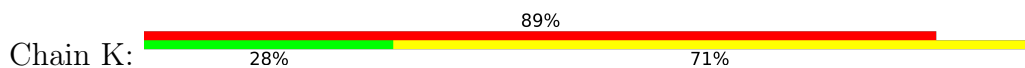


• Molecule 3: Inner capsid protein sigma-2



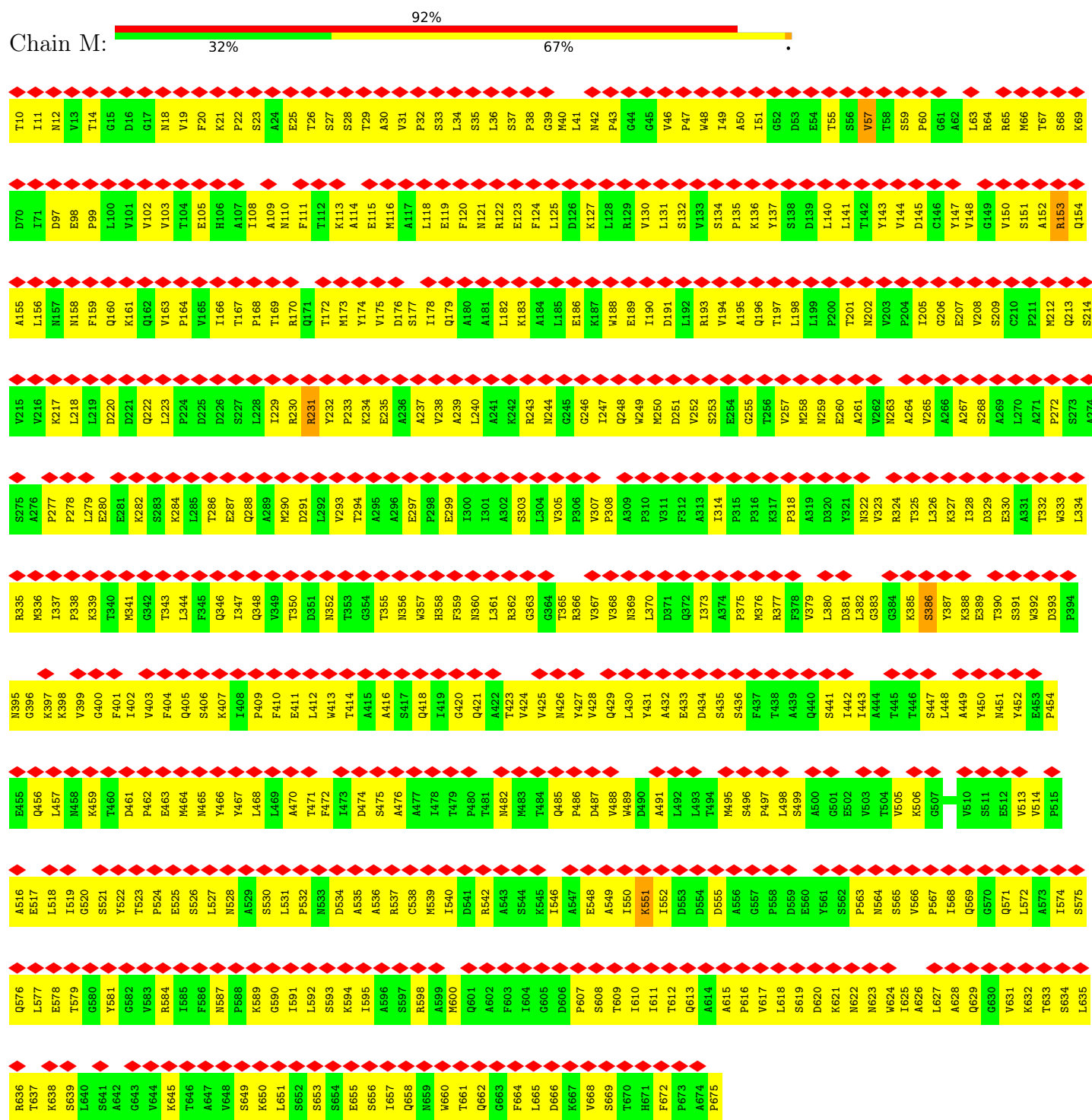


• Molecule 4: Outer capsid protein mu-1

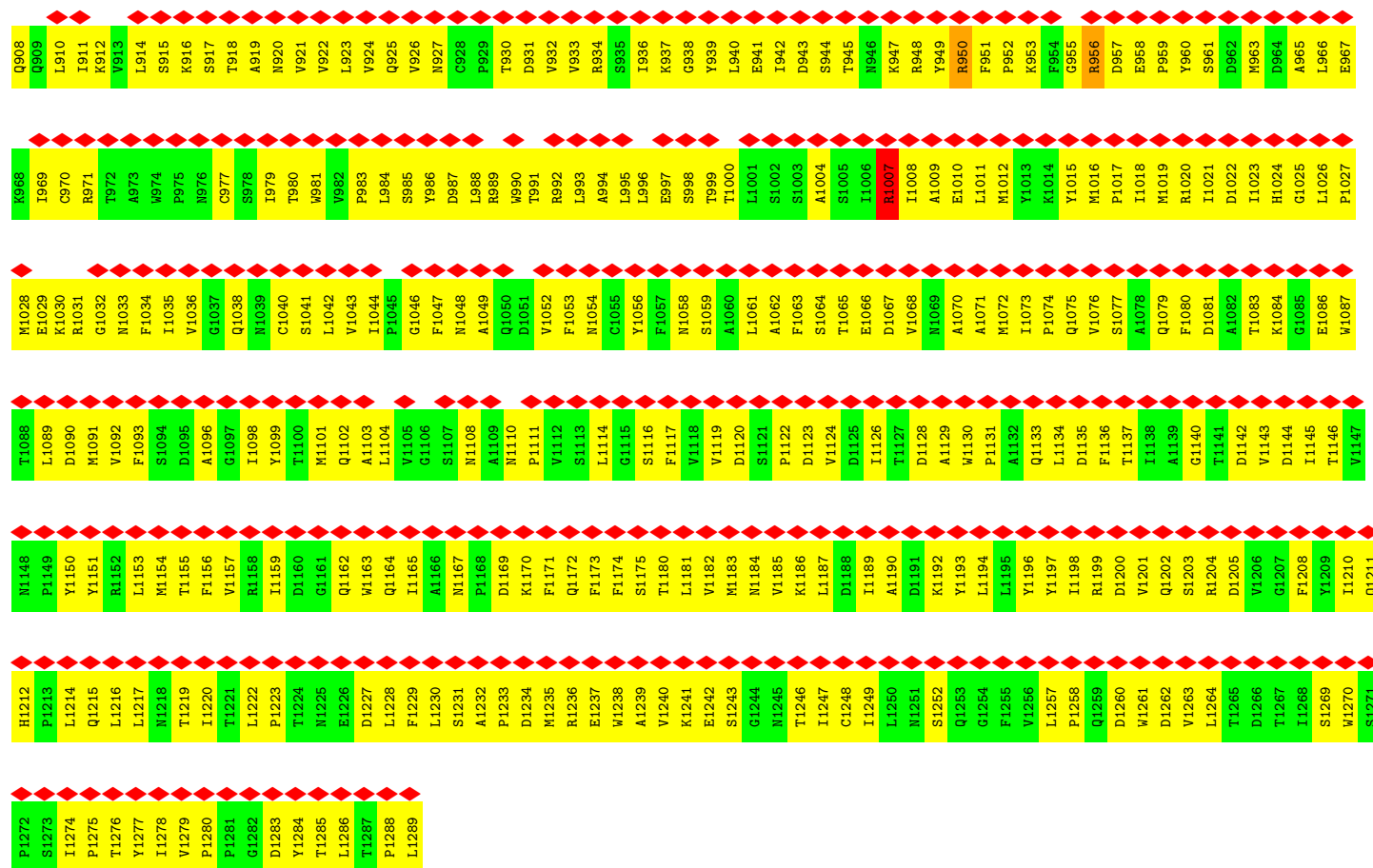




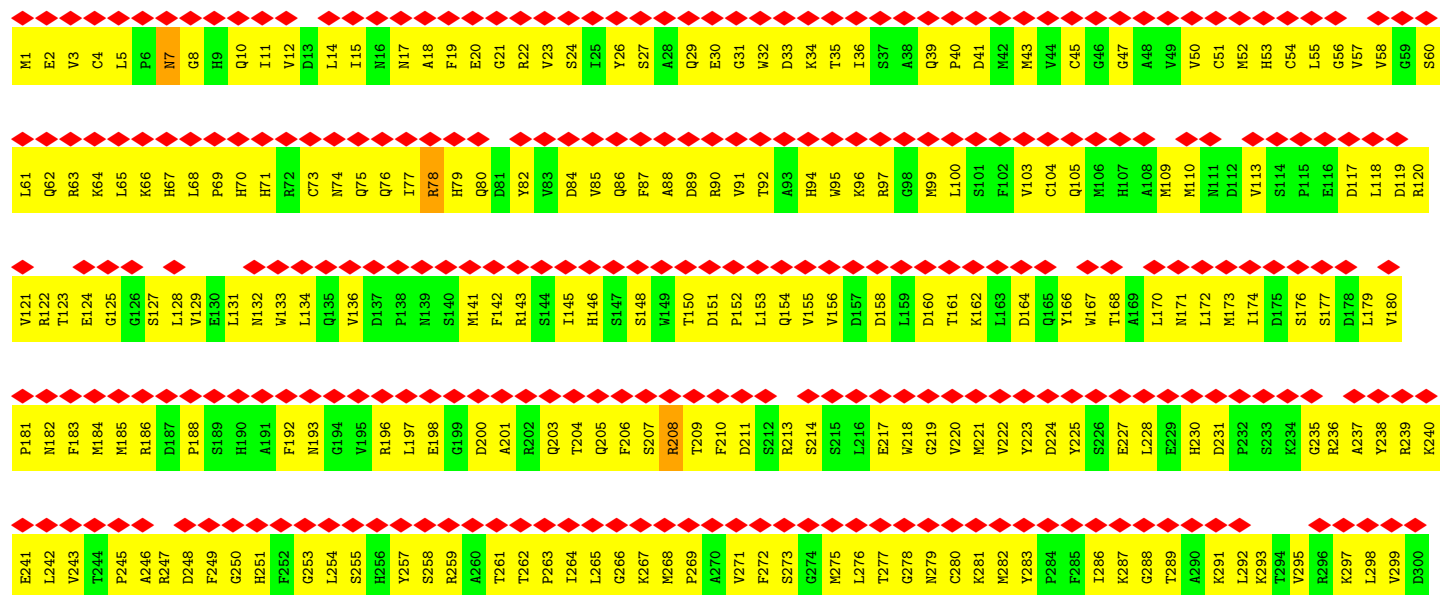
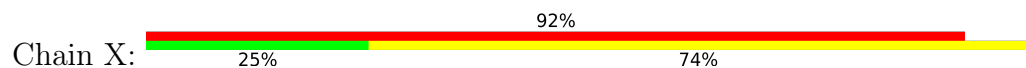
• Molecule 4: Outer capsid protein mu-1

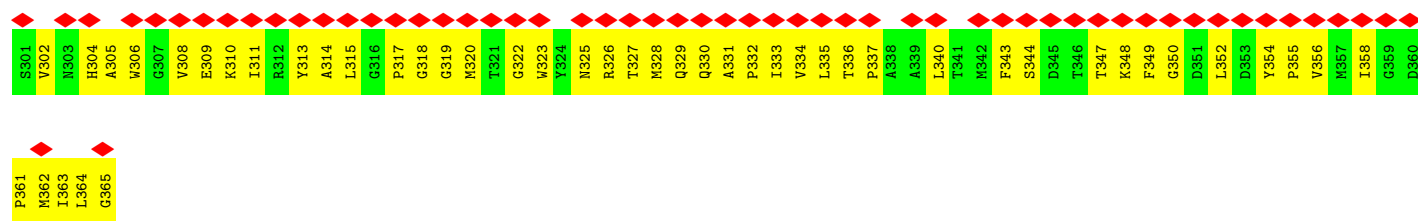


P845	V846	T847	C848	V849	D850	I851	R852	P853	T854	A855	L856	P857	S858	G859	C860	W861	N862	V863	R864	F867	L868	E869	L870	D871	Y872	L873	S874	C875	W876	L877	T878	S879	C880	V881	R882	G883	D884	I885	V886	T887	C888	W889	L890	S891	L892	G893	A894	G898	K899	S900	N901	T902	D904	A905	A906	F907						
A784	R785	T786	L787	L788	P789	A790	D791	P792	V793	L794	G795	E796	S858	G859	C860	W861	N862	V863	R864	F867	L868	E869	L870	D871	Y872	L873	S874	C875	W876	L877	T878	S879	C880	V881	R882	G883	D884	I885	V886	T887	C888	W889	L890	S891	L892	G893	A894	G898	K899	S900	N901	T902	D904	A905	A906	F907						
R724	I725	G726	I727	G728	G729	L730	C731	A732	N733	W734	G735	W736	A737	R738	W739	S740	R801	S802	P803	H804	W805	C806	L807	T808	M809	M810	Y811	L812	L813	T814	W815	S816	S817	A818	V819	D820	G821	G822	D823	W824	W825	L826	D827	L828	G829	T830	G831	P832	E833	A834	R835	L836	L837	F903	D904	A905	A906	F907				
L664	T665	W666	T667	S668	G669	W670	Y671	F672	F673	L674	W675	D676	H677	F678	F679	R680	Y681	E682	T683	L684	C685	T686	L687	S688	R689	Q690	L691	P692	T693	F694	G695	Y696	Y697	R698	D699	G700	S701	S702	W703	T704	G705	W706	E707	L708	I709	G710	I711	E712	N713	P714	G715	F716	S717	W718	L780	E781	T720	Q721	A722	A723		
H604	A605	T606	A607	P608	G609	G610	S611	F612	W613	W614	R615	L616	N617	F618	P619	T620	R621	P622	W623	W624	H625	Y626	L627	E628	G629	R630	L631	L632	P633	W634	L635	Y636	S637	Y638	Y639	Y640	Y641	Y642	Y643	Y644	Y645	Y646	Y647	Y648	Y649	Y650	Y651	Y652	Y653	Y654	Y655	Y656	Y657	Y658	Y659	Y660	Y661	Y662	Y663			
P544	S545	S546	V547	R548	Q549	P550	Q551	Y552	D553	W554	A555	R556	Q557	A558	I559	Y560	D561	L562	A563	R564	P565	F566	P567	S568	Y569	D570	Y571	Q572	F573	W574	Y575	S576	S577	D578	Y579	Q580	Y581	Y582	Y583	G584	H585	D586	D587	L588	S589	I590	S591	Y592	Y593	L594	Y595	E596	S597	L598	L599	S600	S601	C602	M603			
V484	K485	D486	T487	A488	V489	L490	K491	H492	A493	Y494	Q495	A496	T497	D498	P499	Y500	T501	G502	K503	E504	Y505	L506	R507	S508	R509	Q510	S511	Y512	A513	Y514	F515	G516	A517	Y518	A519	G520	H521	S522	G523	A524	D525	Q526	P527	L528	Y529	I530	E531	P532	Y533	L534	Q535	G536	K537	L538	S539	G540	V541	P542	P543			
G363	A364	T365	Q366	S367	G368	Y369	V370	R371	K372	N373	Q374	L375	R376	M379	R380	I381	S382	A383	L384	Q385	S386	L387	S388	D389	T390	P391	S392	P393	Y394	Q395	V396	L397	P398	Q399	Y400	T401	I402	D403	Q404	A405	A406	M407	D408	E409	G410	D411	L412	M413	V414	A415	R416	L417	T418	Q419	P421	L422	R423					
P424	D425	Y426	G427	N428	L429	W430	V431	G432	D433	A434	L435	S436	Y437	Y438	V439	D440	Y441	N442	R443	S444	H445	R446	Y447	V448	L449	S450	S451	E452	L453	P454	Q455	L456	P457	D458	T459	Y460	F461	D462	G463	D464	E465	Q466	Y467	G468	R469	S470	L471	F472	S473	L474	A475	R476	K477	L478	G479	D480	R481	S482	L483			
D182	D183	D184	P185	P186	L187	F188	A189	K190	D191	L192	S193	D194	Y195	K197	A198	F199	Y200	D202	D203	Q263	F264	S265	A266	N267	V268	R269	A270	F271	W272	T273	H274	A275	Q276	P275	V276	T277	A278	G279	V220	L221	Y222	H223	Y224	D225	K226	P227	T228	N229	G230	Q230	H231	Y232	Y233	L234	L235	G236	T237	L238	T239	Q240	M241	V242
A122	F123	L124	S125	N126	Q127	A128	F129	Y130	D131	L132	L133	P134	L135	L136	P137	I137	I138	N139	D140	T141	M142	I143	G144	D145	L146	L147	G148	T149	F150	A151	S152	L153	S154	Q155	F156	F157	Q158	S159	L160	G161	D162	V163	L164	E165	V166	A167	A168	G169	R170	K171	Y172	L173	Q174	M175	E176	N177	Y178	S179	N180	D181		
P62	L63	Q64	G65	L66	V67	L68	D69	T70	Q71	L72	Y73	G74	F75	P76	G77	A78	F79	D80	D81	Q82	E83	R84	F85	M86	R87	E88	K89	L90	R91	V92	L93	K94	Y95	E96	V97	L98	R99	I100	Y101	P102	I103	S104	N105	Y106	S107	N108	H109	H110	V111	N112	V113	F114	V115	A116	N117	A118	L119	V120	G121			

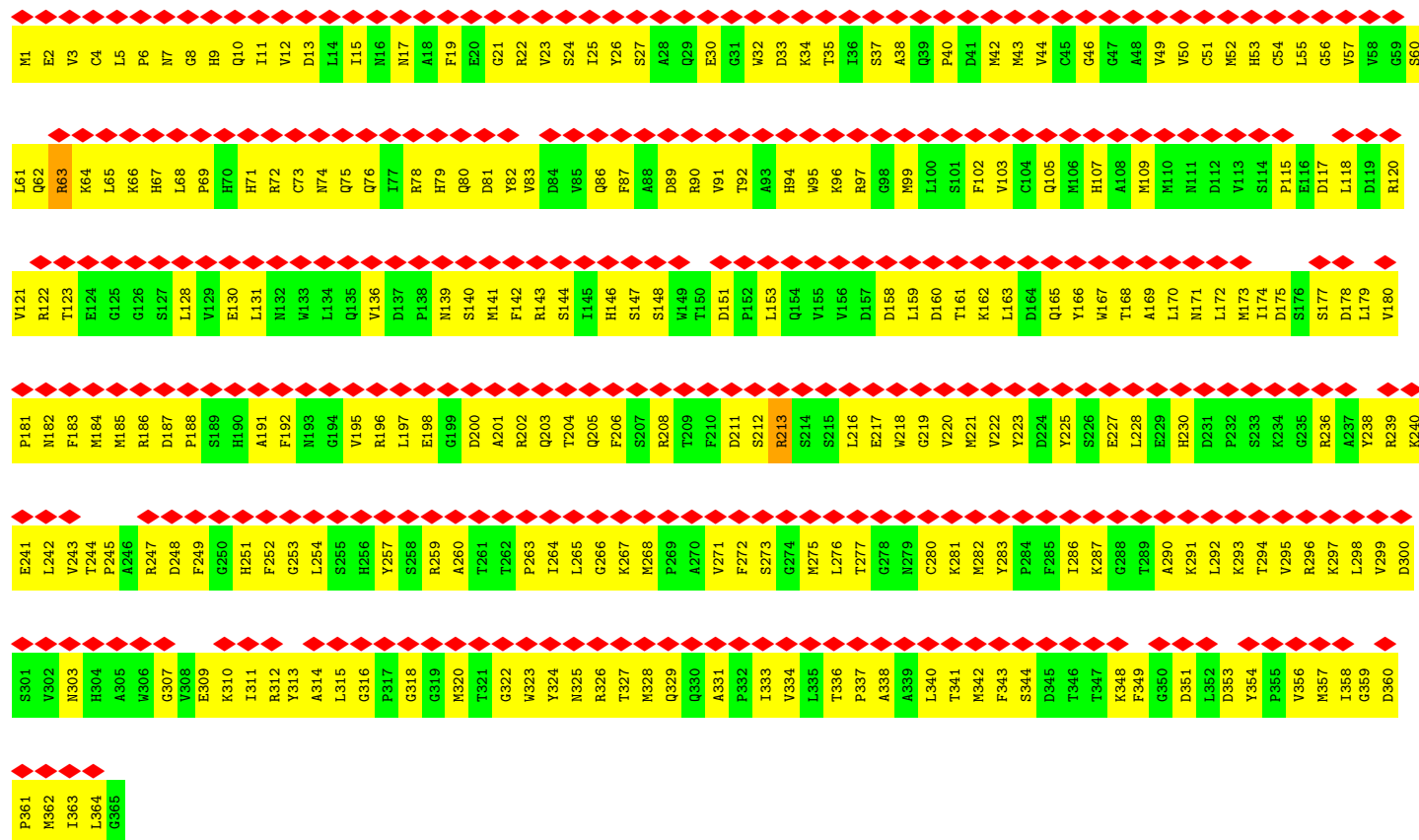


● Molecule 6: Outer capsid protein sigma-3

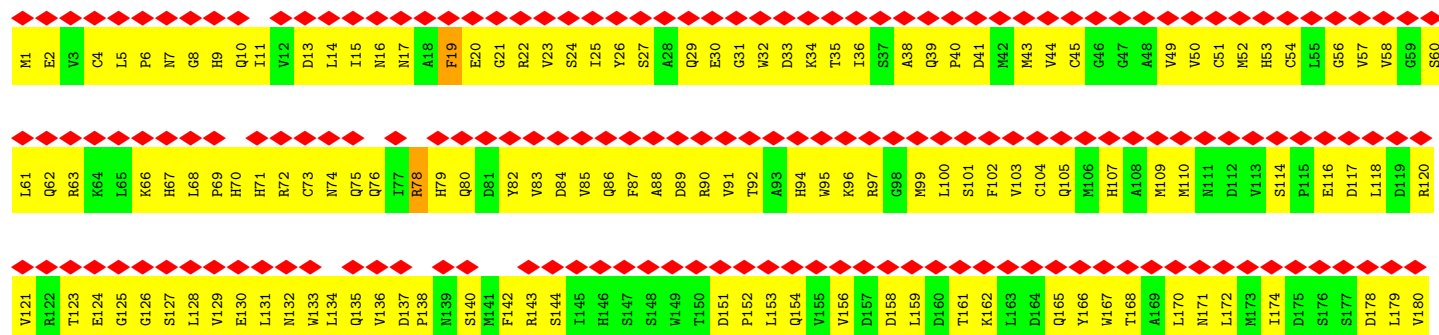
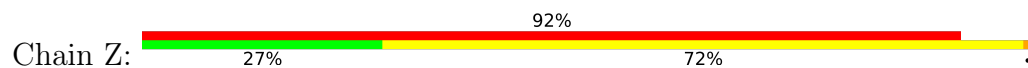




• Molecule 6: Outer capsid protein sigma-3



• Molecule 6: Outer capsid protein sigma-3





4 Experimental information

Property	Value	Source
EM reconstruction method	TOMOGRAPHY	Depositor
Imposed symmetry	POINT, Not provided	
Number of tilted images used	Not provided	
Resolution determination method	FSC 0.143 CUT-OFF	Depositor
CTF correction method	PHASE FLIPPING AND AMPLITUDE CORRECTION	Depositor
Microscope	FEI TITAN KRIOS	Depositor
Voltage (kV)	300	Depositor
Electron dose ($e^-/\text{\AA}^2$)	2	Depositor
Minimum defocus (nm)	Not provided	
Maximum defocus (nm)	Not provided	
Magnification	Not provided	
Image detector	GATAN K2 SUMMIT (4k x 4k)	Depositor
Maximum voxel value	8.955	Depositor
Minimum voxel value	-5.801	Depositor
Average voxel value	0.054	Depositor
Voxel value standard deviation	0.430	Depositor
Recommended contour level	2.7	Depositor
Tomogram size (\AA)	648.0, 648.0, 648.0	wwPDB
Tomogram dimensions	360, 360, 360	wwPDB
Tomogram angles ($^\circ$)	90.0, 90.0, 90.0	wwPDB
Grid spacing (\AA)	1.8, 1.8, 1.8	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	B	0.38	0/8391	0.56	1/11492 (0.0%)
2	C	0.40	0/8174	0.56	0/11194
3	D	0.39	0/3398	0.56	2/4626 (0.0%)
3	P	0.40	0/3398	0.56	1/4626 (0.0%)
4	K	0.36	0/4971	0.54	1/6787 (0.0%)
4	L	0.36	0/4971	0.56	0/6787
4	M	0.35	0/4971	0.54	0/6787
5	O	0.40	0/10385	0.56	1/14172 (0.0%)
6	X	0.39	0/2957	0.53	0/4005
6	Y	0.36	0/2957	0.53	0/4005
6	Z	0.37	0/2957	0.53	0/4005
All	All	0.38	0/57530	0.55	6/78486 (0.0%)

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

Mol	Chain	#Chirality outliers	#Planarity outliers
1	B	0	1
2	C	0	1
4	L	0	1
4	M	0	2
5	O	0	1
6	X	0	1
All	All	0	7

There are no bond length outliers.

All (6) bond angle outliers are listed below:

Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
5	O	1007	ARG	NE-CZ-NH1	-5.83	117.39	120.30

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
4	K	484	THR	C-N-CA	-5.68	107.50	121.70
3	D	7	LEU	CA-CB-CG	5.38	127.68	115.30
3	D	217	LEU	CA-CB-CG	-5.16	103.42	115.30
1	B	599	LEU	CA-CB-CG	-5.05	103.69	115.30
3	P	29	LEU	CA-CB-CG	-5.04	103.72	115.30

There are no chirality outliers.

All (7) planarity outliers are listed below:

Mol	Chain	Res	Type	Group
1	B	977	PHE	Peptide
2	C	383	THR	Peptide
4	L	615	ALA	Peptide
4	M	386	SER	Peptide
4	M	57	VAL	Peptide
5	O	200	TYR	Peptide
6	X	214	SER	Peptide

5.2 Too-close contacts

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

Mol	Chain	Non-H	H(model)	H(added)	Clashes	Symm-Clashes
1	B	8171	0	8083	851	0
2	C	7958	0	7871	867	0
3	D	3313	0	3215	361	0
3	P	3313	0	3215	415	0
4	K	4871	0	4900	587	0
4	L	4871	0	4900	541	0
4	M	4871	0	4900	524	0
5	O	10127	0	9910	1080	0
6	X	2885	0	2816	282	0
6	Y	2885	0	2816	296	0
6	Z	2885	0	2816	298	0
All	All	56150	0	55442	5837	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 52.

All (5837) 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)
5:O:1230:LEU:O	5:O:1276:THR:HA	1.45	1.14
1:B:760:GLU:O	1:B:764:TRP:HB2	1.46	1.14
5:O:704:THR:HA	5:O:758:ARG:O	1.47	1.12
4:K:68:SER:HG	4:K:97:ASP:N	1.47	1.11
4:L:142:THR:O	4:L:164:PRO:HA	1.50	1.10
6:X:319:GLY:O	6:X:323:TRP:HB2	1.54	1.08
1:B:1074:VAL:HA	1:B:1107:ASN:O	1.52	1.07
3:P:240:GLN:O	3:P:251:TRP:HA	1.57	1.04
4:L:368:VAL:HB	4:L:468:LEU:HB3	1.41	1.01
2:C:268:ILE:HA	2:C:304:ILE:HA	1.44	0.99
4:L:22:PRO:HD2	4:L:209:SER:HA	1.45	0.98
3:P:161:ILE:HA	3:P:272:VAL:O	1.63	0.98
3:P:316:ILE:H	3:P:320:THR:HB	1.29	0.97
2:C:385:PHE:HB2	2:C:410:ARG:HE	1.25	0.97
3:D:120:GLN:HB2	3:D:123:ARG:HE	1.30	0.96
4:L:143:TYR:HA	4:L:163:VAL:O	1.65	0.96
1:B:261:GLY:HA3	1:B:312:SER:HA	1.49	0.95
5:O:980:THR:HB	5:O:1020:ARG:O	1.67	0.95
5:O:940:LEU:HA	5:O:950:ARG:O	1.66	0.94
1:B:603:VAL:HG23	1:B:604:VAL:HG23	1.50	0.93
4:M:375:PRO:HA	4:M:452:TYR:O	1.69	0.93
3:P:41:GLN:O	3:P:45:TRP:HB2	1.70	0.92
2:C:607:VAL:O	2:C:876:ALA:HA	1.70	0.92
6:Y:271:VAL:O	6:Y:282:MET:HA	1.69	0.92
1:B:1231:ARG:NE	1:B:1251:VAL:O	2.02	0.92
4:K:358:HIS:ND1	4:K:475:SER:O	2.01	0.92
6:X:295:VAL:HA	6:X:298:LEU:HD13	1.50	0.91
4:M:434:ASP:HA	4:M:441:SER:HA	1.49	0.91
5:O:241:MET:HB3	5:O:249:ILE:HB	1.52	0.91
1:B:1036:ASN:HD21	1:B:1207:ARG:HE	1.18	0.91
1:B:293:GLN:HE22	1:B:401:LYS:HG2	1.35	0.91
2:C:398:ASP:HB3	2:C:401:LYS:HB3	1.50	0.91
5:O:230:GLY:O	5:O:232:HIS:ND1	2.04	0.91
4:K:384:GLY:HA2	4:K:442:ILE:HB	1.52	0.91
5:O:1153:LEU:HG	5:O:1200:ASP:HA	1.52	0.91
5:O:119:LEU:HD11	5:O:136:LEU:HD11	1.52	0.90
3:P:333:ARG:NH1	3:P:337:THR:OG1	2.04	0.90
5:O:1171:PHE:HA	5:O:1185:VAL:HA	1.52	0.90
4:M:67:THR:HA	4:M:99:PRO:HA	1.54	0.90
4:K:43:PRO:HD3	4:M:113:LYS:HG3	1.54	0.90

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
4:K:263:ASN:HD22	4:L:43:PRO:HD2	1.35	0.90
2:C:1007:ARG:NH1	2:C:1008:THR:O	2.06	0.89
4:K:48:TRP:HA	4:K:66:MET:H	1.35	0.89
4:M:400:GLY:O	4:M:470:ALA:HA	1.72	0.89
3:P:85:ARG:HE	3:P:86:TRP:H	1.21	0.89
4:K:147:TYR:HA	4:K:159:PHE:HA	1.54	0.88
4:K:320:ASP:HA	4:K:493:LEU:O	1.71	0.88
1:B:741:LEU:HD12	1:B:742:PRO:HD2	1.55	0.88
3:D:360:LEU:HA	3:D:363:GLN:HE21	1.38	0.88
4:K:357:TRP:HA	4:K:475:SER:HA	1.53	0.88
6:Z:249:PHE:HB2	6:Z:259:ARG:HH21	1.39	0.88
1:B:543:LEU:HD13	1:B:546:ILE:HD11	1.56	0.87
4:K:444:ALA:HB1	4:L:327:LYS:HG2	1.56	0.87
6:X:237:ALA:HA	6:X:240:LYS:HD3	1.55	0.87
3:P:165:ARG:HE	3:P:269:SER:HB2	1.38	0.87
6:X:223:TYR:HD2	6:X:354:TYR:HB2	1.39	0.87
3:P:67:GLN:NE2	3:P:392:PHE:O	2.07	0.86
1:B:920:ARG:NH2	1:B:981:ASP:OD2	2.09	0.86
5:O:988:LEU:O	5:O:991:THR:OG1	1.92	0.86
2:C:1041:ALA:O	2:C:1143:ALA:HA	1.76	0.86
6:Z:200:ASP:HA	6:Z:203:GLN:HB2	1.57	0.86
5:O:300:ARG:NH1	5:O:334:ILE:O	2.08	0.86
1:B:439:MET:HG2	2:C:862:LEU:HG	1.59	0.85
5:O:702:SER:HB3	5:O:758:ARG:HH21	1.38	0.85
2:C:1158:ASN:ND2	2:C:1198:TYR:OH	2.09	0.85
3:P:98:ALA:HB3	3:P:102:ASP:H	1.39	0.85
4:K:120:PHE:HA	4:K:242:LYS:HE2	1.57	0.85
5:O:526:GLN:HE22	5:O:528:LEU:HA	1.42	0.85
1:B:790:VAL:HG12	1:B:794:ARG:HE	1.40	0.85
2:C:468:ARG:NH2	2:C:922:MET:SD	2.49	0.85
2:C:741:LEU:HG	2:C:743:ILE:H	1.40	0.84
4:L:622:ASN:ND2	6:Z:2:GLU:OE2	2.10	0.84
6:Y:244:THR:HG23	6:Y:247:ARG:H	1.41	0.84
4:K:452:TYR:OH	4:K:459:LYS:NZ	2.09	0.84
6:Y:78:ARG:HE	6:Y:80:GLN:HE21	1.21	0.84
2:C:373:ASN:HB3	2:C:1259:ARG:HE	1.41	0.84
3:P:41:GLN:HE21	3:P:45:TRP:HD1	1.22	0.84
6:Y:174:ILE:O	6:Y:178:ASP:N	2.11	0.84
6:Y:259:ARG:HB2	6:Y:272:PHE:HB2	1.60	0.84
3:D:399:ASN:HD22	3:D:402:TRP:HD1	1.24	0.84
5:O:508:SER:OG	5:O:509:ARG:NH1	2.10	0.84

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:P:240:GLN:OE1	3:P:244:ARG:NH1	2.09	0.84
4:K:534:ASP:OD1	4:K:537:ARG:NE	2.10	0.84
5:O:851:ILE:HG22	5:O:871:ASP:HA	1.60	0.84
5:O:951:PHE:O	5:O:955:GLY:N	2.11	0.84
5:O:112:ASN:ND2	5:O:142:MET:SD	2.51	0.84
6:Z:182:ASN:HD21	6:Z:362:MET:H	1.24	0.84
1:B:892:PRO:HG2	1:B:895:LEU:HB2	1.59	0.83
4:L:122:ARG:NH1	4:L:133:VAL:O	2.10	0.83
4:M:19:VAL:HB	4:M:248:GLN:HB2	1.60	0.83
3:P:142:ARG:O	3:P:146:GLN:NE2	2.11	0.83
3:P:94:LEU:HD11	3:P:104:LEU:HD22	1.59	0.83
3:P:239:PHE:HA	3:P:252:ILE:O	1.77	0.83
4:K:306:PRO:HG2	5:O:880:GLY:HA2	1.60	0.83
2:C:392:ARG:O	2:C:403:TYR:OH	1.97	0.83
2:C:759:ASN:HB3	2:C:762:THR:HG23	1.61	0.83
5:O:419:GLN:HE21	5:O:709:ILE:HA	1.44	0.83
5:O:514:TYR:HA	5:O:575:TYR:HB3	1.61	0.83
1:B:600:TYR:OH	1:B:828:PHE:O	1.97	0.83
3:D:341:ASP:OD1	3:D:354:GLN:NE2	2.12	0.83
5:O:88:GLU:HA	5:O:91:ARG:HE	1.42	0.83
6:Z:72:ARG:NH2	6:Z:73:CYS:O	2.12	0.82
6:X:251:HIS:HD2	6:X:254:LEU:HG	1.43	0.82
2:C:579:GLY:O	2:C:582:ARG:NH1	2.10	0.82
2:C:1000:GLN:HG3	2:C:1010:ASN:HB2	1.60	0.82
5:O:318:ILE:HG23	5:O:322:LEU:HD21	1.58	0.82
6:Y:212:SER:OG	6:Y:213:ARG:NH1	2.12	0.82
6:Y:293:LYS:O	6:Y:297:LYS:NZ	2.12	0.82
5:O:610:GLY:O	5:O:658:VAL:N	2.10	0.82
3:P:9:LYS:HE2	3:P:122:ASP:HA	1.61	0.82
2:C:892:PRO:HD2	2:C:895:LEU:HD22	1.59	0.82
5:O:125:SER:O	5:O:127:GLN:NE2	2.11	0.82
5:O:635:ILE:O	5:O:662:SER:OG	1.96	0.82
1:B:483:LEU:HD22	1:B:493:VAL:HG22	1.59	0.82
1:B:859:GLN:NE2	3:P:96:TRP:O	2.13	0.82
2:C:920:ARG:HA	2:C:923:ILE:HD12	1.61	0.82
3:D:350:THR:HG22	3:D:353:GLN:HE21	1.43	0.82
4:K:115:GLU:HA	4:K:118:LEU:HD13	1.62	0.82
5:O:20:THR:H	5:O:277:THR:HA	1.45	0.82
5:O:226:LYS:O	5:O:278:ARG:NH1	2.13	0.82
6:X:182:ASN:H	6:X:363:ILE:HG12	1.42	0.82
2:C:920:ARG:NH1	2:C:924:THR:OG1	2.12	0.82

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
6:Y:90:ARG:NH2	6:Y:151:ASP:O	2.13	0.82
5:O:212:TRP:HE1	3:P:137:PRO:HG2	1.44	0.81
1:B:764:TRP:HA	1:B:767:ARG:HD2	1.62	0.81
4:L:115:GLU:HA	4:L:118:LEU:HD12	1.59	0.81
4:L:505:VAL:O	6:Z:312:ARG:NH2	2.13	0.81
2:C:268:ILE:HG13	2:C:304:ILE:HG13	1.63	0.81
2:C:522:ILE:HA	2:C:525:ILE:HD12	1.62	0.81
4:M:581:TYR:N	6:Y:71:HIS:O	2.11	0.81
5:O:504:GLU:HG3	5:O:507:ARG:HH12	1.46	0.81
4:K:538:CYS:HB3	4:K:542:ARG:HH21	1.45	0.81
4:L:223:LEU:HD12	4:L:224:PRO:HD2	1.61	0.81
5:O:1239:ALA:HA	5:O:1249:ILE:HA	1.61	0.81
3:D:206:ASP:O	3:D:211:ARG:NH2	2.13	0.81
4:K:346:GLN:HA	4:K:360:ASN:HA	1.63	0.81
5:O:816:SER:HA	5:O:819:VAL:HB	1.63	0.81
3:D:78:ILE:HG23	3:D:81:ARG:HH21	1.44	0.81
5:O:932:VAL:O	5:O:1015:TYR:OH	1.99	0.81
6:X:143:ARG:NH1	6:X:218:TRP:O	2.13	0.81
2:C:385:PHE:HB2	2:C:410:ARG:NE	1.95	0.81
4:K:454:PRO:HA	4:K:459:LYS:HE2	1.62	0.81
2:C:604:VAL:HG22	2:C:873:VAL:HB	1.62	0.80
2:C:937:GLN:O	2:C:947:ARG:NH1	2.14	0.80
3:P:185:HIS:HB3	3:P:190:ILE:HD11	1.63	0.80
1:B:656:MET:O	1:B:660:GLU:N	2.12	0.80
4:K:379:VAL:HB	4:K:494:THR:HB	1.64	0.80
4:M:397:LYS:O	4:M:430:LEU:N	2.12	0.80
5:O:1184:ASN:O	5:O:1186:LYS:NZ	2.14	0.80
3:P:275:THR:N	3:P:330:PHE:O	2.15	0.80
1:B:680:THR:O	1:B:683:THR:OG1	1.99	0.80
4:L:391:SER:OG	6:Y:275:MET:O	1.98	0.80
4:M:140:LEU:HB3	4:M:166:ILE:HD12	1.63	0.80
4:L:306:PRO:HB2	4:L:512:GLU:HG2	1.64	0.80
5:O:49:GLN:HB2	5:O:183:ASP:HB3	1.63	0.80
6:Z:71:HIS:CE1	6:Z:73:CYS:HB2	2.16	0.80
2:C:451:PHE:HB3	2:C:1254:TYR:HE2	1.47	0.80
6:X:131:LEU:HB3	6:X:361:PRO:HB2	1.64	0.80
1:B:797:LEU:HA	1:B:800:LEU:HD12	1.64	0.80
4:L:140:LEU:O	4:L:170:ARG:NH1	2.14	0.80
4:L:324:ARG:NH2	4:L:487:ASP:O	2.15	0.80
1:B:582:ARG:HH12	1:B:880:ARG:HG3	1.46	0.80
6:X:4:CYS:SG	6:X:5:LEU:N	2.52	0.80

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:C:880:ARG:HG3	2:C:881:ALA:H	1.45	0.80
6:Z:263:PRO:HB3	6:Z:269:PRO:HD3	1.64	0.80
1:B:1083:ILE:HG13	1:B:1094:ILE:HD13	1.62	0.80
1:B:770:GLU:HA	1:B:773:LYS:HD2	1.63	0.79
5:O:768:ARG:NH1	5:O:840:ILE:O	2.15	0.79
3:P:52:LEU:HD21	3:P:281:MET:HA	1.63	0.79
6:Z:222:VAL:N	6:Z:357:MET:O	2.10	0.79
1:B:385:PHE:O	1:B:428:ARG:NH2	2.15	0.79
2:C:852:GLN:O	2:C:996:GLN:NE2	2.14	0.79
2:C:999:ILE:O	2:C:1010:ASN:HA	1.82	0.79
4:K:351:ASP:OD1	4:K:355:THR:N	2.14	0.79
5:O:112:ASN:OD1	5:O:115:VAL:N	2.13	0.79
6:Z:295:VAL:HA	6:Z:298:LEU:HD12	1.64	0.79
4:L:48:TRP:N	4:L:147:TYR:O	2.15	0.79
2:C:741:LEU:HD12	2:C:742:PRO:HD2	1.65	0.79
4:K:383:GLY:O	4:K:385:LYS:NZ	2.16	0.79
4:L:404:PHE:HB2	4:L:467:TYR:HB2	1.64	0.79
6:X:299:VAL:HG13	6:X:320:MET:HG3	1.65	0.79
2:C:694:ILE:O	2:C:703:ARG:NH1	2.16	0.79
4:K:380:LEU:O	4:K:447:SER:HA	1.83	0.79
5:O:1222:LEU:HD12	5:O:1223:PRO:HD2	1.65	0.79
2:C:365:PHE:O	2:C:369:ASN:ND2	2.14	0.79
4:M:18:ASN:HA	4:M:250:MET:H	1.45	0.79
6:Y:12:VAL:HA	6:Y:15:ILE:HD12	1.62	0.79
5:O:882:ARG:NH2	5:O:884:ASP:OD1	2.16	0.79
3:P:242:ASN:HD22	3:P:244:ARG:HH22	1.30	0.79
2:C:494:THR:HB	2:C:1270:VAL:HB	1.65	0.79
2:C:849:VAL:O	2:C:851:ARG:NH1	2.16	0.79
4:L:525:GLU:O	4:L:529:ALA:N	2.16	0.79
5:O:49:GLN:NE2	5:O:183:ASP:O	2.15	0.79
6:Y:236:ARG:HD2	6:Y:239:ARG:HE	1.47	0.79
1:B:309:ARG:HH12	1:B:321:SER:H	1.30	0.78
5:O:317:PRO:HA	5:O:387:LEU:HD13	1.65	0.78
3:P:183:PHE:HA	3:P:253:LEU:HD23	1.64	0.78
6:Y:171:ASN:HA	6:Y:174:ILE:HD12	1.65	0.78
5:O:55:VAL:HG12	5:O:177:ASN:HB2	1.64	0.78
1:B:602:GLY:HA2	1:B:833:VAL:HG13	1.65	0.78
4:L:198:LEU:HG	4:L:199:LEU:HG	1.63	0.78
5:O:585:HIS:ND1	5:O:586:ASP:OD2	2.17	0.78
5:O:418:THR:HA	5:O:709:ILE:HG22	1.65	0.78
5:O:525:ASP:HB3	5:O:554:VAL:HG23	1.65	0.78

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:C:309:ARG:NH2	2:C:368:ASP:O	2.17	0.78
3:D:8:PHE:CZ	3:D:125:TYR:HB2	2.19	0.78
3:D:35:SER:O	3:D:38:GLN:NE2	2.17	0.78
5:O:317:PRO:HG3	5:O:387:LEU:HD22	1.63	0.78
3:P:169:ASP:O	3:P:211:ARG:NH1	2.17	0.78
3:D:130:TYR:O	3:D:134:ALA:N	2.16	0.78
3:D:326:GLN:NE2	3:D:327:LEU:O	2.17	0.78
4:K:28:SER:HB2	4:K:244:ASN:HA	1.65	0.78
4:K:461:ASP:HB3	4:K:463:GLU:HG2	1.66	0.78
5:O:1129:ALA:HB3	5:O:1144:ASP:HB2	1.65	0.78
3:D:130:TYR:CE2	3:D:132:PHE:HB2	2.18	0.78
5:O:535:GLN:O	5:O:537:LYS:NZ	2.17	0.78
5:O:980:THR:CB	5:O:1020:ARG:O	2.30	0.78
1:B:1186:SER:H	1:B:1221:ALA:HB3	1.48	0.78
4:L:218:LEU:HD12	4:L:221:ASP:HB3	1.64	0.78
5:O:353:ILE:HB	5:O:371:ARG:HB2	1.65	0.78
5:O:743:ILE:H	5:O:787:ILE:HB	1.49	0.78
4:L:168:PRO:O	4:L:171:GLN:NE2	2.16	0.78
2:C:807:THR:HG21	2:C:886:LEU:HA	1.65	0.77
4:K:130:VAL:HG13	4:K:131:LEU:HG	1.66	0.77
4:K:344:LEU:HA	4:K:361:LEU:O	1.84	0.77
5:O:828:LEU:HD12	5:O:888:CYS:HB2	1.67	0.77
1:B:418:LYS:NZ	1:B:1224:ASP:OD1	2.18	0.77
2:C:708:ILE:HG23	2:C:712:TYR:HD1	1.49	0.77
2:C:797:LEU:HD23	2:C:800:LEU:HD12	1.67	0.77
4:M:375:PRO:HB2	4:M:498:LEU:HB2	1.67	0.77
6:Y:184:MET:O	6:Y:186:ARG:NH1	2.17	0.77
1:B:589:SER:OG	1:B:592:ARG:NH2	2.17	0.77
5:O:37:ASN:ND2	5:O:40:ARG:HE	1.81	0.77
5:O:253:ASP:HA	5:O:256:LEU:HD12	1.66	0.77
5:O:638:TYR:HB2	5:O:653:PHE:HE1	1.48	0.77
5:O:506:LEU:O	5:O:507:ARG:NH1	2.17	0.77
4:M:115:GLU:O	4:M:122:ARG:NH2	2.17	0.77
5:O:95:TYR:HA	5:O:98:LEU:HD12	1.67	0.77
5:O:532:PRO:HA	5:O:535:GLN:HB2	1.67	0.77
4:K:525:GLU:OE1	4:K:528:ASN:ND2	2.18	0.77
5:O:1171:PHE:HB3	5:O:1185:VAL:HG23	1.65	0.77
1:B:256:VAL:HA	1:B:372:LEU:HD12	1.67	0.77
2:C:270:PRO:HG3	2:C:297:PRO:HB2	1.65	0.77
2:C:684:TRP:O	2:C:840:ARG:NH2	2.18	0.77
1:B:746:GLN:OE1	5:O:158:GLN:NE2	2.17	0.77

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:D:123:ARG:NH1	3:D:125:TYR:OH	2.18	0.77
5:O:391:PRO:HA	5:O:788:LEU:HB2	1.65	0.77
3:P:139:PHE:HA	3:P:142:ARG:HE	1.49	0.77
6:X:246:ALA:HA	6:X:249:PHE:HB2	1.66	0.77
3:D:61:ALA:O	3:D:64:ARG:N	2.18	0.76
4:M:357:TRP:HA	4:M:475:SER:HA	1.68	0.76
2:C:718:GLN:HB2	2:C:738:VAL:HG13	1.67	0.76
2:C:1022:ASP:O	2:C:1025:GLN:NE2	2.18	0.76
4:M:34:LEU:HD21	4:M:110:ASN:HB3	1.68	0.76
6:Y:182:ASN:OD1	6:Y:362:MET:N	2.18	0.76
4:L:636:ARG:NH2	4:M:29:THR:OG1	2.17	0.76
1:B:366:VAL:HG21	1:B:1024:VAL:HG22	1.66	0.76
2:C:275:GLN:NE2	2:C:276:VAL:O	2.18	0.76
4:M:154:GLN:HG2	4:M:158:ASN:HD21	1.50	0.76
6:Z:30:GLU:OE2	6:Z:38:ALA:N	2.17	0.76
6:Z:180:VAL:N	6:Z:364:LEU:O	2.17	0.76
3:P:271:PHE:O	3:P:328:HIS:ND1	2.16	0.76
3:D:9:LYS:O	3:D:154:ASN:ND2	2.18	0.76
4:K:144:VAL:O	4:K:162:GLN:N	2.19	0.76
1:B:1187:ASP:OD1	1:B:1188:HIS:ND1	2.17	0.76
5:O:8:ARG:HD3	5:O:315:ALA:HB3	1.68	0.76
2:C:1146:TYR:HB2	2:C:1180:PHE:HE1	1.50	0.76
5:O:172:TYR:HB3	5:O:187:LEU:HD13	1.67	0.76
5:O:278:ARG:O	5:O:281:GLN:NE2	2.17	0.76
6:Z:95:TRP:HH2	6:Z:270:ALA:HB2	1.51	0.76
1:B:1034:GLU:HB3	1:B:1037:LEU:HD22	1.67	0.75
2:C:352:LEU:HA	2:C:954:SER:HA	1.67	0.75
4:K:304:LEU:HD21	4:K:621:LYS:HD2	1.68	0.75
3:P:90:ARG:HH21	3:P:115:PRO:HA	1.49	0.75
6:Z:347:THR:HG22	6:Z:349:PHE:H	1.51	0.75
2:C:1120:ARG:NH1	2:C:1170:ILE:O	2.18	0.75
5:O:16:PRO:HG3	5:O:270:ALA:HA	1.69	0.75
5:O:1053:PHE:N	5:O:1065:THR:OG1	2.19	0.75
4:L:614:ALA:O	6:Z:63:ARG:NH2	2.20	0.75
4:M:352:ASN:ND2	4:M:424:VAL:O	2.19	0.75
3:P:9:LYS:HA	3:P:124:VAL:HA	1.68	0.75
6:Y:5:LEU:HD12	6:Y:6:PRO:HD2	1.67	0.75
1:B:348:GLY:HA3	1:B:1175:ILE:H	1.51	0.75
3:D:5:ALA:O	3:D:144:TYR:OH	2.02	0.75
5:O:892:LEU:H	5:O:925:GLN:HE22	1.31	0.75
3:P:197:ALA:O	3:P:203:TRP:NE1	2.19	0.75

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:P:256:SER:OG	3:P:258:ASN:ND2	2.19	0.75
2:C:623:LEU:HA	2:C:626:LEU:HD13	1.68	0.75
5:O:282:CYS:SG	5:O:283:TYR:N	2.55	0.75
1:B:847:VAL:HG13	1:B:870:THR:HA	1.69	0.75
2:C:546:ILE:HD12	2:C:822:ILE:HD11	1.69	0.75
3:P:170:MET:HB3	3:P:215:ILE:HD11	1.69	0.75
1:B:375:HIS:HA	1:B:1259:ARG:HB3	1.68	0.75
1:B:439:MET:HG3	2:C:861:ALA:HA	1.67	0.75
4:K:298:PRO:O	4:K:302:ALA:N	2.19	0.75
4:M:151:SER:HG	4:M:154:GLN:H	1.34	0.75
4:L:250:MET:HB2	4:L:257:VAL:HG22	1.69	0.75
5:O:373:ASN:O	5:O:376:ARG:NH1	2.17	0.75
6:Y:257:TYR:HB3	6:Y:343:PHE:HA	1.67	0.75
6:Z:217:GLU:HA	6:Z:266:GLY:HA2	1.67	0.75
3:D:411:SER:HA	3:D:414:ILE:HD12	1.69	0.75
2:C:1121:TYR:O	2:C:1125:GLN:NE2	2.19	0.74
5:O:516:GLY:HA3	5:O:577:ASP:HB2	1.69	0.74
5:O:707:GLU:OE2	5:O:758:ARG:NH1	2.20	0.74
6:Y:259:ARG:N	6:Y:272:PHE:O	2.18	0.74
4:K:216:VAL:HA	4:K:219:LEU:HD12	1.68	0.74
5:O:353:ILE:HD11	5:O:369:TYR:HB3	1.69	0.74
5:O:722:ALA:HA	5:O:725:ILE:HD12	1.69	0.74
1:B:516:ILE:HG21	1:B:1013:PRO:HB3	1.69	0.74
3:D:250:VAL:HG12	3:D:251:TRP:H	1.52	0.74
4:L:582:GLY:O	6:Z:70:HIS:ND1	2.14	0.74
5:O:997:GLU:OE1	5:O:999:THR:OG1	2.04	0.74
3:P:164:VAL:O	3:P:269:SER:OG	2.00	0.74
6:Z:198:GLU:H	6:Z:201:ALA:HB3	1.51	0.74
2:C:1131:LYS:HA	2:C:1160:TRP:CH2	2.21	0.74
4:K:64:ARG:NH1	4:K:65:ARG:O	2.20	0.74
4:K:377:ARG:HB3	4:K:496:SER:HB2	1.69	0.74
4:K:566:VAL:O	4:K:569:GLN:NE2	2.19	0.74
4:L:250:MET:HG2	4:L:254:GLU:HB2	1.69	0.74
4:M:350:THR:HA	4:M:356:ASN:HA	1.67	0.74
1:B:415:ASN:O	2:C:1082:ARG:NE	2.19	0.74
2:C:262:LEU:N	2:C:311:ALA:O	2.20	0.74
2:C:1211:CYS:SG	2:C:1212:THR:N	2.59	0.74
4:K:393:ASP:O	4:K:397:LYS:NZ	2.13	0.74
5:O:1047:PHE:O	5:O:1087:TRP:NE1	2.19	0.74
1:B:374:ARG:NH1	1:B:1262:TYR:O	2.19	0.74
1:B:1166:TRP:CD1	1:B:1179:PRO:HD3	2.23	0.74

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:C:406:MET:HE1	2:C:453:THR:H	1.51	0.74
2:C:462:ARG:HA	2:C:465:TRP:HB3	1.70	0.74
2:C:527:ASN:ND2	2:C:867:THR:OG1	2.21	0.74
4:M:223:LEU:HB2	4:M:230:ARG:HH21	1.52	0.74
5:O:372:LYS:HB3	5:O:374:GLN:HG2	1.68	0.74
5:O:435:LEU:HB2	5:O:644:PHE:HD2	1.51	0.74
1:B:481:TRP:NE1	2:C:669:TYR:OH	2.19	0.74
2:C:479:ILE:O	2:C:483:LEU:HG	1.88	0.74
3:D:73:SER:O	3:D:77:GLN:NE2	2.21	0.74
4:L:46:VAL:HG11	4:L:65:ARG:HG3	1.67	0.74
4:L:435:SER:HA	4:L:442:ILE:HG23	1.68	0.74
5:O:1241:LYS:HA	5:O:1246:THR:HA	1.67	0.74
3:P:229:HIS:CE1	3:P:232:ASN:H	2.05	0.74
4:M:19:VAL:HG21	4:M:257:VAL:HG11	1.69	0.74
5:O:178:TYR:OH	5:O:180:ASN:ND2	2.20	0.74
5:O:409:GLU:OE1	5:O:409:GLU:N	2.21	0.74
6:Y:72:ARG:N	6:Y:72:ARG:HD2	2.01	0.74
2:C:536:LEU:HA	2:C:539:ILE:HD12	1.70	0.74
6:Z:20:GLU:HG3	6:Z:22:ARG:HB2	1.69	0.74
4:K:145:ASP:OD1	4:K:162:GLN:N	2.20	0.74
4:M:48:TRP:N	4:M:147:TYR:O	2.20	0.73
4:M:144:VAL:HG12	4:M:145:ASP:H	1.53	0.73
1:B:545:ARG:NH2	1:B:908:MET:SD	2.61	0.73
1:B:884:VAL:O	1:B:887:LEU:N	2.21	0.73
2:C:516:ILE:HA	2:C:519:ILE:HD12	1.67	0.73
2:C:1046:ILE:HB	2:C:1200:ILE:HB	1.70	0.73
4:K:426:ASN:HD22	6:Z:62:GLN:HG2	1.53	0.73
4:K:534:ASP:HB2	4:K:537:ARG:HG3	1.70	0.73
4:K:629:GLN:OE1	4:K:632:LYS:NZ	2.18	0.73
5:O:412:LEU:O	5:O:416:ARG:NH2	2.18	0.73
5:O:724:ARG:NE	5:O:775:ILE:O	2.21	0.73
6:Y:220:VAL:HG13	6:Y:264:ILE:HA	1.69	0.73
2:C:998:ALA:HB1	2:C:1010:ASN:HD21	1.53	0.73
4:K:250:MET:SD	4:K:257:VAL:N	2.61	0.73
3:P:66:TYR:O	3:P:69:SER:OG	2.03	0.73
6:X:221:MET:HA	6:X:358:ILE:HA	1.70	0.73
1:B:265:SER:O	1:B:307:HIS:N	2.20	0.73
1:B:623:LEU:HD12	1:B:626:LEU:HB2	1.70	0.73
1:B:883:THR:O	1:B:887:LEU:HG	1.89	0.73
5:O:1192:LYS:HE3	5:O:1219:THR:HA	1.70	0.73
3:P:170:MET:HA	3:P:211:ARG:HD2	1.70	0.73

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:P:227:VAL:O	3:P:232:ASN:ND2	2.22	0.73
1:B:433:ARG:HA	1:B:450:VAL:HA	1.69	0.73
1:B:944:PRO:HD3	3:P:105:VAL:HA	1.69	0.73
4:L:324:ARG:HE	4:L:489:TRP:HB2	1.53	0.73
6:Y:66:LYS:HB3	6:Y:67:HIS:CE1	2.23	0.73
1:B:338:LEU:HD23	1:B:339:LEU:HG	1.69	0.73
3:D:95:VAL:O	3:D:104:LEU:HA	1.88	0.73
4:K:510:VAL:HG21	4:K:513:VAL:HG22	1.69	0.73
4:M:69:LYS:HE2	4:M:97:ASP:HB2	1.70	0.73
1:B:458:THR:HA	1:B:461:ILE:HD12	1.71	0.73
3:D:3:ARG:HA	3:D:300:GLN:HB2	1.70	0.73
5:O:1173:PHE:HA	5:O:1183:MET:HG2	1.70	0.73
6:X:273:SER:OG	6:X:281:LYS:N	2.20	0.73
1:B:741:LEU:HG	1:B:743:ILE:H	1.52	0.73
4:K:385:LYS:H	4:K:443:ILE:HG22	1.53	0.73
1:B:1159:ALA:O	1:B:1163:THR:OG1	2.07	0.73
5:O:28:HIS:N	5:O:108:ASN:OD1	2.21	0.73
6:X:128:LEU:HG	6:X:129:VAL:HG13	1.71	0.73
1:B:359:LEU:HA	1:B:362:LEU:HD12	1.71	0.73
1:B:380:GLN:HE22	2:C:958:SER:H	1.35	0.72
2:C:466:MET:HA	2:C:469:LEU:HD12	1.71	0.72
4:M:190:ILE:HD12	4:M:193:ARG:HB2	1.69	0.72
5:O:480:ASP:OD1	5:O:481:ARG:NH1	2.22	0.72
1:B:1112:LEU:HD12	1:B:1115:TRP:HB3	1.71	0.72
4:M:624:TRP:HA	4:M:627:LEU:HD12	1.71	0.72
5:O:1054:ASN:OD1	5:O:1064:SER:OG	2.05	0.72
2:C:436:ARG:HB2	2:C:448:VAL:HG22	1.71	0.72
2:C:437:ALA:O	2:C:438:GLN:NE2	2.22	0.72
4:M:566:VAL:O	4:M:569:GLN:NE2	2.22	0.72
5:O:706:ILE:HD13	5:O:757:ARG:HG3	1.69	0.72
6:Z:236:ARG:HD3	6:Z:239:ARG:HH21	1.53	0.72
1:B:580:LYS:O	1:B:582:ARG:NH1	2.22	0.72
2:C:708:ILE:O	2:C:712:TYR:N	2.23	0.72
4:L:406:SER:HB2	4:L:420:GLY:HA3	1.72	0.72
6:Z:16:ASN:HD21	6:Z:335:LEU:HD22	1.54	0.72
6:Z:127:SER:OG	6:Z:130:GLU:OE1	2.08	0.72
1:B:275:GLN:NE2	1:B:1187:ASP:OD1	2.23	0.72
1:B:690:ASN:OD1	1:B:692:GLN:NE2	2.22	0.72
1:B:848:GLY:HA2	1:B:869:THR:H	1.52	0.72
4:L:50:ALA:HA	4:L:63:LEU:HA	1.71	0.72
5:O:836:ILE:HA	5:O:839:LEU:HD13	1.71	0.72

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:C:1130:ILE:HD13	2:C:1135:LEU:HB2	1.72	0.72
4:K:12:ASN:ND2	4:K:16:ASP:O	2.23	0.72
5:O:83:GLU:OE2	5:O:84:ARG:NH1	2.23	0.72
3:P:10:THR:HB	3:P:123:ARG:HB2	1.72	0.72
6:Z:258:SER:N	6:Z:342:MET:O	2.22	0.72
2:C:929:GLN:HE21	2:C:1017:GLY:HA2	1.54	0.72
4:M:251:ASP:O	4:M:255:GLY:N	2.23	0.72
4:M:362:ARG:HD2	4:M:486:PRO:HA	1.70	0.72
5:O:1145:ILE:HD13	5:O:1181:LEU:HG	1.71	0.72
6:Z:133:TRP:HD1	6:Z:134:LEU:HD23	1.53	0.72
4:K:401:PHE:HB3	4:K:468:LEU:HD11	1.70	0.72
4:K:595:ILE:HD13	4:K:598:ARG:HH11	1.55	0.72
4:L:34:LEU:HD11	4:L:113:LYS:HB2	1.71	0.72
5:O:768:ARG:NH2	5:O:841:PRO:O	2.23	0.72
6:X:96:LYS:HA	6:X:99:MET:HG2	1.71	0.72
6:X:236:ARG:HG3	6:X:240:LYS:HD2	1.70	0.72
2:C:844:PRO:HA	2:C:1003:GLN:HA	1.71	0.72
4:K:115:GLU:OE2	4:K:134:SER:OG	2.07	0.72
4:K:211:PRO:O	4:K:214:SER:OG	2.08	0.72
4:L:628:ALA:O	4:L:632:LYS:NZ	2.22	0.72
4:M:386:SER:HB3	4:M:441:SER:HB3	1.71	0.72
5:O:1172:GLN:N	5:O:1184:ASN:O	2.23	0.72
3:P:306:ASN:HB3	3:P:314:ARG:HD3	1.72	0.72
2:C:336:LYS:HG2	2:C:349:ALA:HB2	1.72	0.71
4:K:321:TYR:HH	4:K:537:ARG:HH22	1.38	0.71
4:L:357:TRP:HA	4:L:475:SER:HA	1.72	0.71
5:O:425:ASP:N	5:O:792:PRO:O	2.18	0.71
5:O:1004:ALA:O	5:O:1007:ARG:HD3	1.90	0.71
3:P:48:LEU:HD22	3:P:50:ARG:HD2	1.70	0.71
6:X:12:VAL:HA	6:X:15:ILE:HD12	1.71	0.71
3:D:24:GLU:OE2	3:D:28:HIS:ND1	2.22	0.71
4:K:260:GLU:OE1	4:L:65:ARG:NH1	2.23	0.71
4:L:601:GLN:HA	4:L:604:ILE:HD12	1.70	0.71
1:B:1031:PHE:HZ	1:B:1040:ILE:HG13	1.54	0.71
4:K:606:ASP:O	4:K:609:THR:OG1	2.08	0.71
4:M:505:VAL:O	6:Y:312:ARG:NH2	2.22	0.71
6:Y:121:VAL:HG21	6:Y:128:LEU:HD13	1.72	0.71
6:Y:325:ASN:O	6:Y:329:GLN:NE2	2.22	0.71
1:B:493:VAL:HG12	1:B:1273:GLY:HA2	1.71	0.71
2:C:1040:ILE:HA	2:C:1144:TYR:O	1.90	0.71
4:M:401:PHE:HB2	4:M:426:ASN:HB3	1.71	0.71

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
5:O:394:VAL:O	5:O:740:SER:HA	1.91	0.71
3:P:369:ARG:NH2	3:P:399:ASN:OD1	2.23	0.71
1:B:602:GLY:N	1:B:831:PHE:O	2.21	0.71
2:C:682:HIS:HB3	2:C:1004:TYR:HE2	1.55	0.71
2:C:1050:VAL:O	2:C:1196:VAL:HA	1.90	0.71
3:D:145:GLN:O	3:D:148:SER:OG	2.08	0.71
5:O:224:TYR:HD2	5:O:369:TYR:HH	1.37	0.71
5:O:314:LEU:HB2	5:O:318:ILE:HD11	1.72	0.71
6:X:318:GLY:HA3	6:X:322:GLY:HA3	1.72	0.71
2:C:708:ILE:HD13	2:C:711:ARG:HD2	1.73	0.71
2:C:1187:ASP:OD1	2:C:1188:HIS:ND1	2.23	0.71
2:C:1211:CYS:HA	2:C:1225:LYS:HG3	1.72	0.71
3:D:85:ARG:HH12	3:D:88:ASP:HB3	1.56	0.71
4:K:436:SER:HB3	4:L:324:ARG:HH12	1.54	0.71
5:O:227:PRO:HG3	5:O:232:HIS:HE1	1.56	0.71
5:O:829:GLY:HA3	5:O:891:SER:HB3	1.73	0.71
6:X:90:ARG:O	6:X:94:HIS:ND1	2.24	0.71
6:X:206:PHE:O	6:X:208:ARG:NH1	2.23	0.71
1:B:436:ARG:NH1	2:C:859:GLN:OE1	2.24	0.71
4:K:399:VAL:HG22	4:K:472:PHE:HA	1.72	0.71
4:M:459:LYS:HE2	4:M:462:PRO:HA	1.73	0.71
1:B:1144:TYR:HE2	1:B:1146:TYR:HB3	1.56	0.71
2:C:942:GLN:OE1	2:C:995:THR:OG1	2.08	0.71
3:P:47:SER:O	3:P:50:ARG:NH1	2.24	0.71
6:X:221:MET:HB3	6:X:356:VAL:HG12	1.71	0.71
2:C:380:GLN:HA	2:C:390:ASN:HA	1.73	0.71
4:K:362:ARG:HD2	4:K:486:PRO:HG3	1.73	0.71
5:O:212:TRP:CD2	3:P:138:ARG:HD2	2.26	0.71
3:D:373:GLU:O	3:D:377:ARG:NE	2.22	0.71
4:K:42:ASN:ND2	4:M:113:LYS:O	2.24	0.71
4:K:145:ASP:OD1	4:K:162:GLN:NE2	2.24	0.71
4:L:221:ASP:OD2	4:L:222:GLN:NE2	2.24	0.71
5:O:587:ASP:HA	5:O:590:ILE:HD13	1.72	0.71
3:D:271:PHE:O	3:D:328:HIS:ND1	2.24	0.70
3:D:344:ARG:HB3	3:D:349:MET:HE3	1.73	0.70
5:O:17:THR:HB	5:O:360:VAL:HG13	1.73	0.70
5:O:340:ASN:ND2	5:O:342:THR:O	2.23	0.70
1:B:544:GLN:OE1	1:B:592:ARG:NH2	2.23	0.70
2:C:376:THR:HG23	2:C:1259:ARG:HH12	1.56	0.70
4:M:264:ALA:O	4:M:268:SER:N	2.24	0.70
5:O:1156:PHE:HD2	5:O:1163:TRP:HB3	1.56	0.70

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:P:8:PHE:CE2	3:P:125:TYR:HB2	2.25	0.70
3:P:42:PHE:HA	3:P:64:ARG:HE	1.53	0.70
1:B:508:ARG:HH12	1:B:725:ASN:HB3	1.56	0.70
1:B:768:VAL:HA	1:B:771:LEU:HD12	1.72	0.70
1:B:924:THR:HG23	1:B:983:LEU:HD11	1.73	0.70
2:C:421:ASN:HD21	2:C:1217:PRO:HA	1.56	0.70
3:D:314:ARG:HA	3:D:320:THR:HG21	1.73	0.70
4:K:146:CYS:SG	4:K:147:TYR:N	2.62	0.70
4:K:636:ARG:O	4:L:29:THR:OG1	2.07	0.70
4:L:108:ILE:O	4:L:112:THR:OG1	2.09	0.70
4:M:115:GLU:OE2	4:M:174:TYR:OH	2.09	0.70
3:P:9:LYS:HZ3	3:P:11:VAL:HA	1.57	0.70
3:P:17:GLN:NE2	3:P:18:ASN:OD1	2.24	0.70
1:B:773:LYS:NZ	1:B:798:ASP:OD1	2.24	0.70
2:C:601:ASN:H	2:C:832:GLN:HG2	1.57	0.70
3:D:262:GLN:NE2	3:D:263:ILE:O	2.25	0.70
4:K:176:ASP:HA	4:K:179:GLN:HE21	1.56	0.70
4:M:218:LEU:O	4:M:222:GLN:HG2	1.91	0.70
3:P:200:ASN:OD1	3:P:201:ARG:N	2.23	0.70
6:Y:260:ALA:HA	6:Y:271:VAL:HA	1.74	0.70
2:C:580:LYS:O	2:C:582:ARG:NH1	2.25	0.70
2:C:1067:PHE:HD1	2:C:1136:ARG:HH21	1.37	0.70
3:D:79:PRO:HD3	3:D:96:TRP:HE1	1.57	0.70
4:L:370:LEU:N	4:L:466:TYR:O	2.25	0.70
5:O:27:LEU:HG	5:O:109:GLU:N	2.07	0.70
5:O:1248:CYS:HG	5:O:1261:TRP:HH2	1.37	0.70
4:K:32:PRO:HG3	4:L:38:PRO:HG2	1.71	0.70
4:L:277:PRO:O	4:L:282:LYS:NZ	2.24	0.70
5:O:202:ASP:O	5:O:231:HIS:ND1	2.25	0.70
6:Y:63:ARG:HD3	6:Y:63:ARG:H	1.54	0.70
6:Y:291:LYS:O	6:Y:294:THR:OG1	2.09	0.70
6:Z:109:MET:HG3	6:Z:131:LEU:HD22	1.73	0.70
2:C:375:HIS:HA	2:C:1259:ARG:HD3	1.72	0.70
2:C:1209:LEU:O	2:C:1225:LYS:NZ	2.24	0.70
4:K:396:GLY:O	4:K:429:GLN:NE2	2.19	0.70
5:O:223:HIS:CE1	5:O:281:GLN:HA	2.27	0.70
5:O:300:ARG:O	5:O:304:LEU:HG	1.92	0.70
5:O:830:THR:HG21	5:O:848:CYS:HB3	1.73	0.70
6:Z:22:ARG:HH22	6:Z:159:LEU:HD22	1.57	0.70
2:C:620:VAL:O	2:C:782:GLN:NE2	2.24	0.70
2:C:1224:ASP:OD2	2:C:1225:LYS:N	2.25	0.70

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
4:K:522:TYR:O	4:K:613:GLN:N	2.22	0.70
5:O:43:TRP:HD1	5:O:45:PRO:HD3	1.55	0.70
6:X:127:SER:HA	6:X:364:LEU:HA	1.74	0.70
2:C:759:ASN:HA	2:C:812:GLN:HE22	1.56	0.70
4:L:123:GLU:HG2	4:L:235:GLU:HG2	1.73	0.70
5:O:154:SER:O	5:O:158:GLN:NE2	2.24	0.70
5:O:352:GLN:OE1	5:O:372:LYS:N	2.18	0.70
6:X:64:LYS:O	6:X:66:LYS:NZ	2.23	0.70
6:Y:78:ARG:HE	6:Y:80:GLN:NE2	1.90	0.70
1:B:263:CYS:SG	1:B:462:ARG:NH1	2.65	0.70
2:C:707:GLU:O	2:C:711:ARG:CB	2.40	0.70
2:C:1120:ARG:O	2:C:1124:GLN:HG2	1.92	0.70
4:L:396:GLY:H	4:L:431:TYR:HA	1.56	0.70
2:C:731:ASN:OD1	2:C:734:THR:N	2.24	0.69
5:O:636:THR:N	5:O:656:PHE:O	2.24	0.69
5:O:1077:SER:O	5:O:1089:LEU:HA	1.92	0.69
2:C:401:LYS:O	2:C:404:SER:OG	2.09	0.69
2:C:543:LEU:HG	2:C:592:ARG:HD2	1.73	0.69
4:K:484:THR:OG1	4:K:485:GLN:NE2	2.25	0.69
5:O:472:PHE:HA	5:O:475:ALA:HB3	1.73	0.69
6:Y:79:HIS:HA	6:Y:82:TYR:CZ	2.26	0.69
1:B:1110:PHE:HE1	1:B:1115:TRP:HB2	1.55	0.69
2:C:267:LYS:O	2:C:305:VAL:N	2.19	0.69
3:D:333:ARG:NE	3:D:358:GLU:OE2	2.22	0.69
4:K:15:GLY:O	4:K:213:GLN:NE2	2.25	0.69
4:K:349:VAL:O	4:K:356:ASN:ND2	2.25	0.69
5:O:9:LEU:HD23	5:O:314:LEU:HA	1.72	0.69
5:O:849:VAL:HG22	5:O:868:LEU:HB2	1.73	0.69
3:P:25:LEU:HA	3:P:28:HIS:CD2	2.28	0.69
6:Y:73:CYS:SG	6:Y:74:ASN:N	2.65	0.69
6:Y:141:MET:HE2	6:Y:216:LEU:HD11	1.74	0.69
2:C:729:SER:HB3	2:C:736:PRO:HB2	1.73	0.69
2:C:929:GLN:NE2	2:C:1016:PRO:O	2.25	0.69
6:X:289:THR:O	6:X:293:LYS:NZ	2.26	0.69
1:B:483:LEU:O	1:B:487:ALA:N	2.24	0.69
2:C:500:SER:OG	2:C:1263:GLU:OE1	2.09	0.69
4:K:348:GLN:HB3	4:K:356:ASN:HD21	1.56	0.69
4:K:631:VAL:O	4:K:634:SER:OG	2.11	0.69
3:P:230:GLN:NE2	3:P:238:TYR:OH	2.24	0.69
1:B:760:GLU:O	1:B:764:TRP:CB	2.35	0.69
2:C:601:ASN:OD1	2:C:602:GLY:N	2.25	0.69

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:C:713:TRP:CD2	2:C:714:PRO:HD2	2.28	0.69
4:M:105:GLU:HA	4:M:108:ILE:HD12	1.74	0.69
5:O:10:ALA:N	5:O:313:GLN:O	2.25	0.69
5:O:403:ASP:HB3	5:O:774:LEU:HD12	1.75	0.69
5:O:491:LYS:NZ	5:O:539:SER:O	2.26	0.69
5:O:1123:ASP:OD1	5:O:1124:VAL:N	2.25	0.69
6:Z:295:VAL:O	6:Z:299:VAL:HG23	1.93	0.69
1:B:465:TRP:O	1:B:469:LEU:HG	1.93	0.69
1:B:374:ARG:O	1:B:1259:ARG:NH1	2.26	0.69
1:B:696:PRO:HB3	1:B:703:ARG:HG2	1.75	0.69
1:B:729:SER:OG	1:B:731:ASN:ND2	2.26	0.69
1:B:1044:ASP:OD1	1:B:1141:MET:N	2.22	0.69
2:C:325:ALA:HB3	2:C:331:ASN:HD21	1.58	0.69
2:C:517:SER:HA	2:C:520:ILE:HD12	1.73	0.69
2:C:577:ILE:HA	2:C:580:LYS:HD3	1.75	0.69
2:C:601:ASN:ND2	2:C:832:GLN:OE1	2.25	0.69
2:C:707:GLU:HB3	2:C:711:ARG:HH22	1.57	0.69
2:C:1161:ASN:O	2:C:1164:SER:OG	2.10	0.69
4:K:400:GLY:HA2	4:K:448:LEU:HD21	1.73	0.69
4:K:599:ALA:HB1	4:K:618:LEU:HD11	1.74	0.69
4:L:284:LYS:HA	4:L:287:GLU:HG3	1.74	0.69
4:M:60:PRO:HA	4:M:153:ARG:HD2	1.75	0.69
5:O:448:VAL:N	5:O:666:TRP:O	2.25	0.69
3:P:304:ILE:HA	3:P:324:GLY:HA3	1.75	0.69
6:Y:245:PRO:O	6:Y:249:PHE:N	2.21	0.69
6:Z:138:PRO:HA	6:Z:143:ARG:HB3	1.75	0.69
2:C:283:TYR:HB2	2:C:290:TYR:HB2	1.75	0.69
2:C:381:ASP:HB3	2:C:386:THR:HB	1.75	0.69
4:K:320:ASP:OD1	4:K:493:LEU:N	2.26	0.69
4:K:338:PRO:HG3	6:X:333:ILE:HG12	1.74	0.69
4:K:470:ALA:H	4:K:489:TRP:HE1	1.41	0.69
5:O:37:ASN:OD1	5:O:40:ARG:NH2	2.26	0.69
5:O:220:VAL:HG12	5:O:286:ARG:HB3	1.75	0.69
3:P:2:ALA:HB1	3:P:300:GLN:H	1.56	0.69
2:C:606:THR:H	2:C:640:THR:HG1	1.41	0.69
3:D:98:ALA:HB3	3:D:102:ASP:H	1.58	0.69
5:O:73:TYR:OH	5:O:170:ARG:NH2	2.26	0.69
5:O:172:TYR:HD2	5:O:190:LYS:HB2	1.58	0.69
5:O:509:ARG:HA	5:O:545:SER:N	2.08	0.69
6:Y:197:LEU:HA	6:Y:356:VAL:HB	1.75	0.69
2:C:541:VAL:HG12	2:C:545:ARG:HE	1.58	0.68

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
4:M:590:GLY:O	4:M:593:SER:OG	2.11	0.68
3:P:286:ILE:O	3:P:289:SER:OG	2.11	0.68
1:B:275:GLN:HG3	1:B:278:GLN:HE22	1.56	0.68
2:C:518:GLN:NE2	2:C:830:PRO:O	2.22	0.68
2:C:603:VAL:O	2:C:873:VAL:N	2.26	0.68
2:C:662:ILE:H	2:C:671:GLN:HE22	1.41	0.68
4:K:656:SER:HB3	4:M:305:VAL:HG13	1.74	0.68
4:L:455:GLU:O	4:L:458:ASN:ND2	2.26	0.68
4:M:569:GLN:HA	4:M:572:LEU:HD12	1.74	0.68
5:O:1197:TYR:HA	5:O:1212:HIS:O	1.93	0.68
5:O:1240:VAL:N	5:O:1248:CYS:O	2.22	0.68
5:O:1264:LEU:HD12	5:O:1276:THR:HG22	1.75	0.68
3:P:42:PHE:CZ	3:P:65:TYR:HB2	2.28	0.68
6:Y:198:GLU:OE1	6:Y:356:VAL:N	2.25	0.68
1:B:836:VAL:HB	1:B:839:ASP:HB3	1.76	0.68
1:B:858:THR:HG21	3:P:105:VAL:HG21	1.75	0.68
2:C:772:MET:HG2	2:C:775:LEU:HD23	1.75	0.68
4:L:402:ILE:HA	4:L:424:VAL:HA	1.73	0.68
4:L:631:VAL:O	4:L:635:LEU:HG	1.92	0.68
4:M:375:PRO:HG3	4:M:454:PRO:HD3	1.74	0.68
5:O:48:ASN:OD1	5:O:53:ASN:N	2.13	0.68
5:O:532:PRO:HB2	5:O:537:LYS:HB2	1.76	0.68
5:O:1063:PHE:HE1	5:O:1073:ILE:HG12	1.58	0.68
6:Y:257:TYR:HA	6:Y:342:MET:HG3	1.73	0.68
2:C:494:THR:HG22	2:C:496:PRO:HD3	1.75	0.68
4:K:639:SER:HG	4:L:120:PHE:HE2	1.42	0.68
4:M:151:SER:OG	4:M:154:GLN:N	2.18	0.68
4:M:233:PRO:HG2	4:M:234:LYS:HD3	1.75	0.68
5:O:532:PRO:O	5:O:536:GLY:N	2.26	0.68
6:Z:196:ARG:HE	6:Z:352:LEU:HB3	1.58	0.68
2:C:609:ASP:OD1	2:C:612:SER:N	2.27	0.68
4:L:280:GLU:HG3	4:L:284:LYS:HE2	1.76	0.68
4:L:346:GLN:HA	4:L:360:ASN:HA	1.76	0.68
4:M:472:PHE:HE1	4:M:485:GLN:HG3	1.58	0.68
4:M:522:TYR:O	4:M:612:THR:HA	1.94	0.68
4:M:634:SER:O	4:M:637:THR:OG1	2.09	0.68
5:O:1040:CYS:HB2	5:O:1093:PHE:HZ	1.58	0.68
1:B:438:GLN:O	2:C:859:GLN:NE2	2.27	0.68
1:B:1186:SER:N	1:B:1221:ALA:HB3	2.07	0.68
2:C:486:CYS:O	2:C:488:GLN:NE2	2.27	0.68
2:C:1067:PHE:CD2	2:C:1107:ASN:HB3	2.29	0.68

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
4:L:56:SER:OG	4:L:59:SER:O	2.08	0.68
4:M:49:ILE:N	4:M:66:MET:SD	2.67	0.68
4:M:471:THR:HA	4:M:488:VAL:HB	1.75	0.68
3:P:341:ASP:HA	3:P:354:GLN:HE22	1.59	0.68
6:X:196:ARG:NH2	6:X:352:LEU:O	2.26	0.68
1:B:464:ARG:HB3	1:B:1019:VAL:HG21	1.76	0.68
2:C:355:ARG:CZ	2:C:950:ASP:HA	2.24	0.68
4:L:370:LEU:HD21	4:L:468:LEU:HB2	1.76	0.68
4:M:143:TYR:HA	4:M:163:VAL:O	1.93	0.68
5:O:30:LEU:O	5:O:34:LEU:HG	1.94	0.68
5:O:41:GLU:OE1	5:O:87:ARG:NH2	2.25	0.68
5:O:317:PRO:HB3	5:O:387:LEU:HB3	1.75	0.68
6:X:21:GLY:HA3	6:X:92:THR:HG21	1.74	0.68
1:B:247:LEU:HG	1:B:521:ARG:NH2	2.09	0.68
1:B:809:MET:N	1:B:809:MET:SD	2.66	0.68
2:C:1233:ASN:HB3	2:C:1244:GLN:HE22	1.58	0.68
4:K:48:TRP:HB3	4:K:63:LEU:HD11	1.76	0.68
4:K:334:LEU:HA	4:K:369:ASN:H	1.58	0.68
4:L:324:ARG:NH2	4:L:489:TRP:H	1.92	0.68
4:M:530:SER:OG	4:M:531:LEU:N	2.27	0.68
5:O:391:PRO:HB3	5:O:788:LEU:HD12	1.75	0.68
5:O:742:ALA:HA	5:O:787:ILE:HB	1.74	0.68
5:O:1198:ILE:O	5:O:1211:GLN:NE2	2.26	0.68
6:X:33:ASP:OD2	6:X:34:LYS:N	2.27	0.68
6:Y:60:SER:HB3	6:Y:63:ARG:HH12	1.58	0.68
6:Z:199:GLY:H	6:Z:202:ARG:HE	1.42	0.68
1:B:251:ASP:HB2	1:B:979:LEU:HD22	1.75	0.68
2:C:1003:GLN:HB2	2:C:1007:ARG:N	2.09	0.68
3:D:351:GLN:O	3:D:355:ASN:ND2	2.27	0.68
4:M:379:VAL:HA	4:M:448:LEU:O	1.94	0.68
5:O:423:ARG:HA	5:O:697:VAL:HA	1.75	0.68
5:O:711:ILE:O	5:O:751:VAL:HA	1.94	0.68
6:X:20:GLU:O	6:X:22:ARG:NH1	2.27	0.68
6:X:113:VAL:HG11	6:X:118:LEU:HD13	1.73	0.68
2:C:337:GLN:HG2	2:C:357:ASN:HD22	1.59	0.68
2:C:383:THR:HB	2:C:410:ARG:HH22	1.58	0.68
2:C:1233:ASN:OD1	2:C:1236:THR:OG1	2.12	0.68
4:K:660:TRP:NE1	4:M:622:ASN:OD1	2.26	0.68
4:L:375:PRO:HB3	4:L:453:GLU:HG2	1.76	0.68
4:M:43:PRO:HB2	4:M:102:VAL:HG13	1.74	0.68
5:O:1133:GLN:NE2	5:O:1135:ASP:OD1	2.27	0.68

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
5:O:1197:TYR:HB3	5:O:1211:GLN:NE2	2.08	0.68
6:Z:245:PRO:HB2	6:Z:259:ARG:HH22	1.59	0.68
1:B:1150:TYR:HA	1:B:1182:VAL:O	1.93	0.67
2:C:846:MET:HA	2:C:871:VAL:HG22	1.76	0.67
3:D:70:CYS:O	3:D:73:SER:OG	2.11	0.67
4:M:325:THR:H	4:M:326:LEU:HD12	1.57	0.67
3:P:21:ILE:HD13	3:P:70:CYS:HB2	1.74	0.67
6:Y:259:ARG:O	6:Y:272:PHE:N	2.26	0.67
6:Z:128:LEU:HG	6:Z:129:VAL:HG23	1.75	0.67
6:Z:225:TYR:HA	6:Z:228:LEU:HD13	1.76	0.67
2:C:543:LEU:HD21	2:C:595:LEU:HD12	1.75	0.67
3:D:67:GLN:HG3	3:D:393:ARG:HA	1.76	0.67
4:L:66:MET:HG3	4:L:100:LEU:HD12	1.76	0.67
4:L:636:ARG:NH2	4:M:27:SER:O	2.27	0.67
4:M:10:THR:OG1	4:M:11:ILE:N	2.25	0.67
5:O:203:THR:HA	5:O:231:HIS:HD1	1.59	0.67
3:P:219:LEU:HD21	3:P:238:TYR:HA	1.74	0.67
6:X:7:ASN:HB2	6:X:10:GLN:HG2	1.76	0.67
6:Y:91:VAL:HA	6:Y:94:HIS:CD2	2.29	0.67
1:B:532:ILE:O	1:B:536:LEU:HG	1.94	0.67
2:C:954:SER:O	2:C:956:ARG:NH1	2.26	0.67
3:D:55:SER:HB3	3:D:59:PRO:HB3	1.75	0.67
4:K:378:PHE:O	4:K:449:ALA:HA	1.95	0.67
5:O:80:ASP:HB2	5:O:84:ARG:HH12	1.57	0.67
6:Y:191:ALA:HA	6:Y:195:VAL:HG11	1.77	0.67
1:B:473:ASN:OD1	1:B:505:MET:N	2.22	0.67
1:B:501:VAL:HG12	1:B:503:ARG:HH21	1.58	0.67
1:B:517:SER:HA	1:B:520:ILE:HD12	1.76	0.67
1:B:556:ILE:HD12	1:B:556:ILE:H	1.60	0.67
1:B:591:PHE:HB3	1:B:592:ARG:HH11	1.60	0.67
2:C:339:LEU:HG	2:C:968:TRP:HE1	1.60	0.67
2:C:494:THR:N	2:C:1270:VAL:O	2.22	0.67
2:C:1171:THR:HG1	2:C:1173:THR:HG1	1.40	0.67
4:K:523:THR:O	4:K:526:SER:OG	2.09	0.67
5:O:45:PRO:HB2	5:O:54:ILE:HD13	1.77	0.67
5:O:825:VAL:HA	5:O:885:ILE:HG23	1.76	0.67
3:P:211:ARG:HA	3:P:214:GLN:NE2	2.09	0.67
6:Y:169:ALA:HA	6:Y:172:LEU:HD12	1.76	0.67
2:C:822:ILE:HA	2:C:825:MET:HB2	1.77	0.67
4:M:176:ASP:OD2	4:M:179:GLN:NE2	2.27	0.67
4:M:661:THR:HA	4:M:664:PHE:CD2	2.30	0.67

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
5:O:310:ASN:O	5:O:357:SER:OG	2.12	0.67
3:P:42:PHE:HZ	3:P:65:TYR:HB2	1.60	0.67
2:C:685:PRO:O	2:C:689:MET:N	2.28	0.67
5:O:1054:ASN:HB3	5:O:1061:LEU:HD11	1.77	0.67
3:P:219:LEU:HA	3:P:222:LEU:HD12	1.76	0.67
3:P:300:GLN:NE2	3:P:301:LEU:O	2.27	0.67
6:Y:236:ARG:HH11	6:Y:239:ARG:HE	1.42	0.67
1:B:914:VAL:O	1:B:918:LEU:HG	1.95	0.67
2:C:1059:LEU:HD13	2:C:1205:ASN:H	1.60	0.67
4:K:392:TRP:HE1	4:K:394:PRO:HA	1.60	0.67
4:L:664:PHE:HA	4:L:667:LYS:HD2	1.77	0.67
5:O:59:LEU:HD11	5:O:63:LEU:HB3	1.75	0.67
5:O:460:TYR:OH	5:O:469:ARG:NH2	2.28	0.67
5:O:685:SER:O	5:O:688:SER:OG	2.11	0.67
3:P:58:VAL:HG23	3:P:63:SER:HB2	1.76	0.67
6:Y:180:VAL:N	6:Y:364:LEU:O	2.26	0.67
1:B:575:SER:HA	1:B:578:LEU:HD12	1.77	0.67
3:D:273:MET:O	3:D:329:GLY:HA2	1.94	0.67
4:M:520:GLY:N	4:M:610:ILE:O	2.22	0.67
5:O:18:ILE:HG21	5:O:273:ALA:HA	1.76	0.67
5:O:826:LEU:HD21	5:O:886:VAL:HG22	1.77	0.67
5:O:828:LEU:HB2	5:O:888:CYS:HA	1.77	0.67
6:Z:134:LEU:HA	6:Z:143:ARG:HH22	1.60	0.67
2:C:1063:PRO:HA	2:C:1066:VAL:HG22	1.77	0.67
4:L:402:ILE:HB	4:L:469:LEU:HB2	1.76	0.67
5:O:504:GLU:O	5:O:507:ARG:NH2	2.27	0.67
6:Z:308:VAL:HA	6:Z:311:ILE:HD12	1.76	0.67
2:C:475:ASN:O	2:C:479:ILE:HG12	1.95	0.67
4:K:372:GLN:HE22	4:K:509:VAL:H	1.43	0.67
4:K:561:TYR:HE1	4:L:224:PRO:HG3	1.60	0.67
4:L:540:ILE:HD11	4:L:607:PRO:HB2	1.75	0.67
5:O:425:ASP:HB2	5:O:793:VAL:HA	1.75	0.67
5:O:768:ARG:NH1	5:O:837:LEU:O	2.28	0.67
6:X:74:ASN:HD21	6:X:77:ILE:HD11	1.58	0.67
1:B:1210:PHE:CZ	1:B:1227:ILE:HG23	2.30	0.66
2:C:436:ARG:HA	2:C:448:VAL:HA	1.76	0.66
2:C:572:SER:O	2:C:575:SER:OG	2.13	0.66
2:C:647:LYS:NZ	2:C:678:PHE:O	2.20	0.66
2:C:681:PRO:HB3	2:C:840:ARG:HG3	1.77	0.66
2:C:681:PRO:HA	2:C:684:TRP:CD2	2.30	0.66
2:C:699:ALA:HB1	2:C:702:LEU:HB2	1.76	0.66

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
4:L:51:ILE:HB	4:L:64:ARG:HB3	1.77	0.66
4:L:508:ALA:HB3	6:Z:312:ARG:HH21	1.60	0.66
5:O:490:LEU:HD12	5:O:493:ALA:HB3	1.77	0.66
6:Y:240:LYS:O	6:Y:247:ARG:NH1	2.27	0.66
6:Y:245:PRO:HA	6:Y:248:ASP:HB2	1.78	0.66
1:B:355:ARG:NH2	1:B:954:SER:O	2.28	0.66
1:B:518:GLN:HA	1:B:521:ARG:HD3	1.75	0.66
1:B:1026:LEU:O	1:B:1030:VAL:HG23	1.95	0.66
1:B:1031:PHE:CZ	1:B:1040:ILE:HG13	2.30	0.66
1:B:1135:LEU:HD12	1:B:1136:ARG:H	1.59	0.66
2:C:680:THR:OG1	2:C:1004:TYR:OH	2.13	0.66
3:D:222:LEU:HD22	3:D:418:ILE:HD11	1.77	0.66
3:D:277:PRO:HA	3:D:279:TRP:CE2	2.31	0.66
4:K:262:VAL:HB	4:L:45:GLY:HA2	1.76	0.66
5:O:323:THR:O	5:O:327:LEU:HG	1.95	0.66
6:Y:182:ASN:N	6:Y:362:MET:O	2.27	0.66
6:Z:91:VAL:HG11	6:Z:210:PHE:HB3	1.77	0.66
2:C:624:GLU:HA	2:C:627:TRP:CE3	2.30	0.66
4:K:346:GLN:NE2	4:K:478:ILE:O	2.27	0.66
4:L:47:PRO:HA	4:L:148:VAL:HG22	1.77	0.66
4:M:357:TRP:HE3	4:M:474:ASP:HA	1.60	0.66
5:O:181:ASP:OD1	5:O:182:ASP:N	2.29	0.66
5:O:182:ASP:HB3	5:O:325:ARG:CZ	2.25	0.66
5:O:995:LEU:HA	5:O:999:THR:H	1.59	0.66
6:Z:4:CYS:HB3	6:Z:57:VAL:HB	1.75	0.66
6:Z:54:CYS:SG	6:Z:71:HIS:NE2	2.65	0.66
1:B:348:GLY:CA	1:B:1175:ILE:H	2.09	0.66
2:C:607:VAL:HG12	2:C:641:ASP:HB2	1.77	0.66
3:D:3:ARG:NH2	3:D:318:GLY:O	2.29	0.66
4:K:154:GLN:O	4:K:158:ASN:N	2.29	0.66
4:L:41:LEU:HD23	4:L:106:HIS:HE1	1.61	0.66
6:Y:217:GLU:OE1	6:Y:217:GLU:N	2.28	0.66
6:Z:5:LEU:HD21	6:Z:310:LYS:HE3	1.76	0.66
1:B:307:HIS:HD1	1:B:310:TRP:HD1	1.43	0.66
2:C:355:ARG:NH2	2:C:950:ASP:HA	2.11	0.66
2:C:1032:ASN:O	2:C:1042:ARG:NH2	2.29	0.66
2:C:1231:ARG:HG2	2:C:1250:VAL:HG13	1.78	0.66
4:M:666:ASP:O	4:M:669:SER:OG	2.13	0.66
5:O:340:ASN:ND2	5:O:342:THR:OG1	2.26	0.66
5:O:1258:PRO:HB2	5:O:1260:ASP:OD1	1.95	0.66
6:X:120:ARG:O	6:X:124:GLU:N	2.24	0.66

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:B:317:ASP:OD1	1:B:371:TYR:OH	2.12	0.66
2:C:262:LEU:HB3	2:C:264:THR:HG22	1.78	0.66
2:C:350:ASN:HB3	2:C:353:MET:HG2	1.78	0.66
2:C:362:LEU:O	2:C:366:VAL:HG23	1.96	0.66
2:C:1185:SER:OG	2:C:1213:ASN:ND2	2.28	0.66
3:D:25:LEU:HA	3:D:28:HIS:HB2	1.76	0.66
3:D:81:ARG:NH1	3:D:88:ASP:OD1	2.29	0.66
4:K:284:LYS:HD3	4:M:294:THR:HG21	1.77	0.66
4:K:369:ASN:OD1	4:K:370:LEU:N	2.27	0.66
4:L:385:LYS:HE3	4:L:491:ALA:HA	1.77	0.66
5:O:59:LEU:HD12	5:O:60:PHE:H	1.61	0.66
5:O:1197:TYR:CD2	5:O:1211:GLN:HG3	2.30	0.66
1:B:327:PRO:HG2	1:B:1148:LEU:HD12	1.78	0.66
1:B:1001:TYR:CZ	1:B:1009:PHE:HB3	2.31	0.66
2:C:707:GLU:O	2:C:711:ARG:HB3	1.96	0.66
2:C:968:TRP:O	2:C:972:SER:OG	2.13	0.66
4:K:51:ILE:O	5:O:374:GLN:NE2	2.29	0.66
4:K:455:GLU:OE1	4:K:458:ASN:ND2	2.20	0.66
4:M:327:LYS:HZ3	4:M:329:ASP:H	1.41	0.66
5:O:20:THR:N	5:O:276:VAL:O	2.29	0.66
5:O:1145:ILE:HB	5:O:1181:LEU:HB3	1.78	0.66
6:X:171:ASN:HA	6:X:174:ILE:HG12	1.78	0.66
1:B:440:MET:H	2:C:862:LEU:H	1.44	0.66
2:C:1209:LEU:HG	2:C:1225:LYS:HG2	1.78	0.66
4:K:336:MET:HA	4:K:366:ARG:HA	1.76	0.66
4:L:273:SER:HB3	4:L:276:ALA:HB3	1.77	0.66
4:L:362:ARG:HD2	4:L:486:PRO:HA	1.77	0.66
5:O:1091:MET:N	5:O:1091:MET:SD	2.69	0.66
3:D:123:ARG:HH11	3:D:125:TYR:HH	1.43	0.66
4:K:246:GLY:O	4:K:248:GLN:NE2	2.26	0.66
4:K:583:VAL:HG23	6:X:68:LEU:HD23	1.77	0.66
4:L:360:ASN:OD1	4:L:361:LEU:N	2.29	0.66
5:O:611:SER:HG	5:O:657:GLY:H	1.43	0.66
5:O:1142:ASP:OD1	5:O:1184:ASN:ND2	2.28	0.66
3:P:183:PHE:O	3:P:185:HIS:ND1	2.28	0.66
6:Y:227:GLU:HA	6:Y:230:HIS:CG	2.31	0.66
1:B:1031:PHE:HE1	1:B:1038:PHE:CD2	2.14	0.66
1:B:1077:PHE:HZ	1:B:1108:TRP:HB3	1.60	0.66
1:B:1106:GLY:H	1:B:1135:LEU:HA	1.61	0.66
2:C:1115:TRP:HA	2:C:1122:PHE:CE1	2.31	0.66
3:D:238:TYR:HE1	3:D:240:GLN:HE21	1.43	0.66

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
4:M:272:PRO:HG3	4:M:278:PRO:HD3	1.77	0.66
4:M:625:ILE:O	4:M:629:GLN:NE2	2.28	0.66
5:O:1151:TYR:HD2	5:O:1202:GLN:HE22	1.43	0.66
6:Z:35:THR:HG22	6:Z:152:PRO:HB3	1.77	0.66
1:B:984:LEU:HA	1:B:987:LEU:HD13	1.78	0.65
1:B:1044:ASP:N	1:B:1202:THR:OG1	2.19	0.65
2:C:458:THR:HA	2:C:461:ILE:HD12	1.78	0.65
2:C:681:PRO:HA	2:C:684:TRP:CE2	2.31	0.65
2:C:1085:PHE:HA	2:C:1092:PRO:HA	1.78	0.65
4:M:258:MET:SD	4:M:258:MET:N	2.69	0.65
6:X:74:ASN:OD1	6:X:75:GLN:N	2.29	0.65
6:Y:86:GLN:CD	6:Y:86:GLN:H	1.99	0.65
2:C:868:ASN:OD1	2:C:870:THR:N	2.29	0.65
2:C:983:LEU:H	2:C:985:GLU:HG3	1.60	0.65
4:K:298:PRO:HA	4:K:301:ILE:HG12	1.78	0.65
4:M:154:GLN:O	4:M:158:ASN:ND2	2.30	0.65
6:X:227:GLU:O	6:X:231:ASP:N	2.29	0.65
6:Y:122:ARG:NH2	6:Y:236:ARG:HH21	1.93	0.65
1:B:347:ARG:NH2	1:B:1177:SER:OG	2.30	0.65
1:B:1106:GLY:O	1:B:1136:ARG:N	2.28	0.65
2:C:472:MET:O	2:C:507:TYR:N	2.27	0.65
2:C:851:ARG:O	2:C:995:THR:N	2.29	0.65
4:M:540:ILE:HD12	4:M:608:SER:HA	1.77	0.65
3:P:137:PRO:HD2	3:P:138:ARG:HH11	1.61	0.65
1:B:436:ARG:HA	1:B:448:VAL:HA	1.78	0.65
3:D:14:GLY:O	3:D:395:LYS:NZ	2.28	0.65
3:D:331:GLN:NE2	3:D:333:ARG:H	1.93	0.65
4:K:572:LEU:O	4:K:575:SER:OG	2.12	0.65
4:M:265:VAL:O	4:M:268:SER:OG	2.14	0.65
6:Z:182:ASN:HD21	6:Z:362:MET:N	1.94	0.65
1:B:1033:HIS:O	1:B:1207:ARG:NH2	2.29	0.65
1:B:1186:SER:OG	1:B:1187:ASP:N	2.26	0.65
2:C:839:ASP:HB3	2:C:842:ARG:HE	1.61	0.65
3:D:191:ALA:HA	3:D:194:LEU:HD12	1.77	0.65
3:D:235:ARG:N	3:D:260:SER:OG	2.29	0.65
3:D:275:THR:HG23	3:D:329:GLY:HA3	1.78	0.65
4:M:567:PRO:HB2	4:M:626:ALA:HB1	1.79	0.65
1:B:558:ASN:OD1	1:B:559:THR:N	2.30	0.65
1:B:717:SER:N	1:B:743:ILE:O	2.30	0.65
1:B:759:ASN:O	1:B:763:ASN:N	2.26	0.65
2:C:646:VAL:HG21	2:C:684:TRP:CD2	2.32	0.65

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:C:1245:ILE:HG13	2:C:1247:LEU:HG	1.78	0.65
3:D:386:ASN:OD1	3:D:388:ASN:ND2	2.30	0.65
4:K:582:GLY:O	6:X:70:HIS:ND1	2.28	0.65
4:M:506:LYS:HB2	6:Y:312:ARG:HH12	1.62	0.65
5:O:51:THR:OG1	5:O:53:ASN:OD1	2.14	0.65
5:O:814:GLU:HG2	5:O:984:LEU:HD12	1.78	0.65
5:O:1130:TRP:HA	5:O:1143:VAL:HG13	1.77	0.65
3:P:354:GLN:HA	3:P:357:ILE:HD12	1.77	0.65
4:M:513:VAL:HG12	4:M:514:VAL:H	1.61	0.65
5:O:85:PHE:O	5:O:89:LYS:HG3	1.96	0.65
5:O:611:SER:OG	5:O:657:GLY:N	2.30	0.65
5:O:1235:MET:SD	5:O:1236:ARG:NH1	2.66	0.65
6:Y:326:ARG:HA	6:Y:329:GLN:HG2	1.79	0.65
1:B:680:THR:HG1	1:B:683:THR:HG23	1.61	0.65
4:L:409:PRO:HB2	4:L:412:LEU:HG	1.79	0.65
4:L:426:ASN:HA	6:Y:61:LEU:HD11	1.79	0.65
4:M:145:ASP:HB3	4:M:159:PHE:CE1	2.32	0.65
4:M:567:PRO:HD2	4:M:568:ILE:HD12	1.77	0.65
5:O:223:HIS:HE1	5:O:278:ARG:HE	1.44	0.65
6:X:60:SER:HB3	6:X:63:ARG:HH21	1.62	0.65
1:B:649:PHE:CZ	1:B:702:LEU:HB2	2.32	0.65
1:B:807:THR:O	1:B:810:TYR:HB3	1.97	0.65
1:B:1036:ASN:OD1	1:B:1207:ARG:NH2	2.28	0.65
3:D:186:THR:HA	3:D:252:ILE:HA	1.78	0.65
4:L:350:THR:HA	4:L:356:ASN:HA	1.78	0.65
3:P:7:LEU:N	3:P:144:TYR:OH	2.28	0.65
3:P:25:LEU:HD11	3:P:391:MET:HE2	1.79	0.65
1:B:656:MET:HB3	1:B:659:PHE:HD1	1.62	0.65
1:B:1051:GLN:HG3	1:B:1196:VAL:HG22	1.77	0.65
2:C:601:ASN:N	2:C:832:GLN:HG2	2.12	0.65
4:L:168:PRO:O	4:L:172:THR:OG1	2.13	0.65
4:L:662:GLN:NE2	4:L:666:ASP:OD1	2.29	0.65
5:O:371:ARG:HH12	5:O:375:LEU:HB2	1.62	0.65
1:B:394:LEU:HB3	1:B:399:ALA:HB2	1.78	0.64
1:B:400:GLU:HA	1:B:403:TYR:CD1	2.32	0.64
1:B:1147:MET:N	1:B:1178:VAL:O	2.30	0.64
2:C:373:ASN:HB3	2:C:1259:ARG:NE	2.12	0.64
2:C:1020:ILE:O	2:C:1024:VAL:HG23	1.97	0.64
2:C:1035:TYR:HA	2:C:1038:PHE:CD2	2.32	0.64
4:K:144:VAL:HG12	4:K:145:ASP:H	1.63	0.64
4:K:629:GLN:HE21	4:L:200:PRO:HB2	1.61	0.64

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
4:M:55:THR:HG22	4:M:156:LEU:HD21	1.78	0.64
4:M:229:ILE:HA	4:M:232:TYR:HB2	1.78	0.64
5:O:13:LEU:HD12	5:O:358:PRO:HD2	1.79	0.64
6:Y:33:ASP:OD2	6:Y:34:LYS:N	2.30	0.64
4:K:321:TYR:HB2	4:K:493:LEU:HD11	1.78	0.64
4:K:397:LYS:O	4:K:430:LEU:N	2.28	0.64
5:O:192:LEU:HB2	5:O:195:TYR:HD2	1.63	0.64
5:O:993:LEU:O	5:O:997:GLU:N	2.26	0.64
3:P:172:SER:O	3:P:176:ASN:ND2	2.30	0.64
6:X:158:ASP:OD2	6:X:162:LYS:NZ	2.29	0.64
6:Y:8:GLY:HA2	6:Y:11:ILE:HD12	1.79	0.64
6:Y:179:LEU:HB2	6:Y:363:ILE:HG12	1.79	0.64
1:B:266:PHE:HB3	1:B:304:ILE:HD11	1.77	0.64
1:B:746:GLN:NE2	5:O:155:GLN:OE1	2.28	0.64
1:B:1045:ILE:HG22	1:B:1201:SER:HA	1.79	0.64
1:B:1260:TYR:HB3	1:B:1262:TYR:HD2	1.61	0.64
4:K:339:LYS:HA	4:K:364:GLY:HA3	1.79	0.64
4:L:25:GLU:OE1	4:L:25:GLU:N	2.31	0.64
4:L:143:TYR:O	4:L:162:GLN:NE2	2.30	0.64
4:L:664:PHE:CD2	4:L:665:LEU:HD22	2.32	0.64
6:Y:109:MET:HG2	6:Y:131:LEU:HD11	1.79	0.64
2:C:475:ASN:OD1	2:C:478:GLU:N	2.30	0.64
2:C:577:ILE:HD13	2:C:627:TRP:HB3	1.80	0.64
2:C:749:ASN:OD1	2:C:750:VAL:N	2.31	0.64
4:K:581:TYR:O	6:X:71:HIS:N	2.27	0.64
4:M:383:GLY:HA2	4:M:443:ILE:HB	1.80	0.64
5:O:884:ASP:O	5:O:920:ASN:N	2.29	0.64
6:Z:131:LEU:HB3	6:Z:361:PRO:HG2	1.78	0.64
1:B:1151:TYR:CE2	1:B:1181:MET:HG3	2.33	0.64
2:C:825:MET:HE3	2:C:908:MET:HG2	1.79	0.64
4:L:382:LEU:HD12	4:L:385:LYS:HE2	1.79	0.64
4:L:398:LYS:HE3	4:L:473:ILE:HD11	1.80	0.64
5:O:45:PRO:HA	5:O:57:VAL:HG22	1.77	0.64
5:O:432:GLY:HA3	5:O:478:ILE:HB	1.78	0.64
5:O:595:VAL:HA	5:O:598:LEU:HG	1.78	0.64
3:P:229:HIS:N	3:P:232:ASN:O	2.22	0.64
2:C:649:PHE:HE2	2:C:706:ALA:HB2	1.62	0.64
2:C:869:THR:HG23	2:C:870:THR:HG23	1.78	0.64
3:D:190:ILE:O	3:D:194:LEU:HG	1.97	0.64
4:K:218:LEU:HD22	4:K:668:VAL:HG21	1.79	0.64
4:L:523:THR:H	4:L:526:SER:HG	1.46	0.64

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
5:O:617:ASN:HA	5:O:651:LEU:HD13	1.79	0.64
6:X:47:GLY:HA3	6:X:61:LEU:HD23	1.80	0.64
1:B:432:GLY:O	1:B:451:PHE:N	2.31	0.64
1:B:472:MET:O	1:B:507:TYR:N	2.30	0.64
1:B:482:ALA:HB2	1:B:726:VAL:HG11	1.78	0.64
1:B:584:SER:OG	1:B:585:ASN:N	2.31	0.64
2:C:598:TRP:O	2:C:832:GLN:NE2	2.31	0.64
3:D:8:PHE:CE1	3:D:125:TYR:HB2	2.33	0.64
4:K:279:LEU:HD23	4:K:279:LEU:H	1.63	0.64
4:L:104:THR:HG23	4:L:107:ALA:H	1.63	0.64
4:L:404:PHE:HE2	4:L:419:ILE:HB	1.61	0.64
4:M:137:TYR:HA	4:M:140:LEU:HD13	1.78	0.64
4:M:147:TYR:CE1	4:M:159:PHE:HB2	2.33	0.64
5:O:240:GLN:OE1	5:O:250:ASN:ND2	2.31	0.64
5:O:515:PHE:CZ	5:O:550:PHE:HB2	2.32	0.64
5:O:736:ASN:ND2	5:O:838:GLU:OE2	2.30	0.64
5:O:768:ARG:HH22	5:O:840:ILE:HG13	1.62	0.64
6:X:119:ASP:O	6:X:123:THR:OG1	2.14	0.64
6:Z:4:CYS:SG	6:Z:5:LEU:N	2.67	0.64
6:Z:259:ARG:N	6:Z:272:PHE:O	2.31	0.64
1:B:1044:ASP:OD1	1:B:1142:GLY:N	2.26	0.64
1:B:1171:THR:HG23	1:B:1174:SER:HB3	1.80	0.64
2:C:533:GLN:O	2:C:537:GLN:HG2	1.97	0.64
4:K:404:PHE:O	4:K:466:TYR:HA	1.97	0.64
5:O:1154:MET:HB2	5:O:1201:VAL:HG21	1.79	0.64
5:O:1264:LEU:HD21	5:O:1278:ILE:HG13	1.80	0.64
6:Z:11:ILE:O	6:Z:15:ILE:HG12	1.98	0.64
1:B:860:PRO:HD3	3:P:97:SER:HB3	1.79	0.64
2:C:318:ARG:NH2	2:C:370:LEU:H	1.96	0.64
2:C:603:VAL:HG23	2:C:604:VAL:HG23	1.80	0.64
4:L:368:VAL:HG21	4:L:468:LEU:HD23	1.79	0.64
4:L:631:VAL:HG13	4:L:632:LYS:H	1.63	0.64
4:M:220:ASP:OD1	4:M:230:ARG:NH1	2.31	0.64
5:O:271:ARG:NE	5:O:274:GLN:OE1	2.31	0.64
3:P:399:ASN:HA	3:P:402:TRP:CD1	2.33	0.64
6:X:302:VAL:HA	6:X:305:ALA:HB3	1.80	0.64
6:Y:17:ASN:O	6:Y:21:GLY:N	2.31	0.64
6:Z:99:MET:O	6:Z:103:VAL:HG23	1.97	0.64
1:B:290:TYR:CZ	2:C:1120:ARG:HB3	2.32	0.64
1:B:558:ASN:O	1:B:561:SER:OG	2.15	0.64
2:C:318:ARG:HH21	2:C:370:LEU:H	1.46	0.64

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:C:473:ASN:ND2	2:C:503:ARG:O	2.30	0.64
3:D:16:LEU:HB2	3:D:20:PRO:HG3	1.80	0.64
3:D:316:ILE:H	3:D:320:THR:HB	1.61	0.64
3:D:360:LEU:O	3:D:364:THR:HG23	1.98	0.64
4:K:64:ARG:NH2	4:K:70:ASP:OD2	2.31	0.64
4:K:370:LEU:HG	4:K:467:TYR:HA	1.79	0.64
4:M:592:LEU:HA	4:M:595:ILE:HD12	1.79	0.64
1:B:381:ASP:HB2	1:B:388:GLY:HA2	1.80	0.63
1:B:931:LEU:HD13	1:B:934:LEU:HD12	1.80	0.63
1:B:1084:SER:HB2	1:B:1093:MET:H	1.63	0.63
2:C:473:ASN:OD1	2:C:505:MET:N	2.26	0.63
2:C:502:ASN:N	2:C:1263:GLU:OE1	2.31	0.63
2:C:707:GLU:HB3	2:C:711:ARG:NH2	2.12	0.63
3:D:130:TYR:HE2	3:D:132:PHE:HB2	1.59	0.63
4:M:167:THR:HG22	4:M:170:ARG:H	1.63	0.63
4:M:179:GLN:O	4:M:183:LYS:HG3	1.99	0.63
4:M:581:TYR:CG	6:Y:73:CYS:HB3	2.33	0.63
3:P:240:GLN:H	3:P:252:ILE:H	1.44	0.63
6:X:97:ARG:NH2	6:X:154:GLN:O	2.30	0.63
1:B:434:PHE:N	1:B:449:ASP:O	2.30	0.63
2:C:589:SER:OG	2:C:590:SER:N	2.31	0.63
2:C:767:ARG:O	2:C:771:LEU:HG	1.98	0.63
2:C:852:GLN:HA	2:C:995:THR:H	1.64	0.63
3:D:230:GLN:HE21	3:D:240:GLN:HE21	1.45	0.63
4:M:198:LEU:HD21	4:M:223:LEU:HD21	1.79	0.63
4:M:389:GLU:HB3	4:M:392:TRP:HB3	1.80	0.63
5:O:271:ARG:HG2	5:O:273:ALA:H	1.63	0.63
5:O:707:GLU:OE2	5:O:758:ARG:HG2	1.97	0.63
6:X:207:SER:OG	6:X:209:THR:O	2.16	0.63
2:C:261:GLY:HA3	2:C:312:SER:HA	1.80	0.63
2:C:307:HIS:HD2	2:C:310:TRP:HB3	1.62	0.63
2:C:451:PHE:HB3	2:C:1254:TYR:CE2	2.33	0.63
2:C:1067:PHE:CE1	2:C:1136:ARG:HB3	2.32	0.63
4:K:133:VAL:HG22	4:L:153:ARG:HH11	1.62	0.63
4:K:243:ARG:HH22	4:K:261:ALA:HA	1.63	0.63
4:L:49:ILE:O	4:L:64:ARG:N	2.30	0.63
5:O:227:PRO:HG2	5:O:230:GLY:H	1.63	0.63
6:Y:295:VAL:HG23	6:Y:324:TYR:HE1	1.64	0.63
1:B:999:ILE:N	1:B:1010:ASN:OD1	2.31	0.63
1:B:1101:MET:SD	1:B:1101:MET:N	2.71	0.63
5:O:622:PRO:HA	5:O:625:HIS:HB3	1.80	0.63

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:P:222:LEU:O	3:P:227:VAL:N	2.28	0.63
1:B:705:TRP:CZ3	1:B:771:LEU:HD13	2.34	0.63
2:C:574:ALA:O	2:C:578:LEU:HG	1.97	0.63
2:C:906:TYR:HB3	2:C:907:PRO:HD3	1.79	0.63
2:C:1158:ASN:HD21	2:C:1160:TRP:HB2	1.63	0.63
3:D:210:ALA:O	3:D:213:THR:N	2.30	0.63
4:K:358:HIS:CD2	4:K:477:ALA:HA	2.33	0.63
5:O:323:THR:O	5:O:326:TRP:HB3	1.99	0.63
5:O:886:VAL:HB	5:O:922:VAL:HA	1.79	0.63
5:O:1074:PRO:HB2	5:O:1092:VAL:HG13	1.80	0.63
3:P:282:ASP:OD1	3:P:283:VAL:N	2.31	0.63
2:C:282:ALA:HA	2:C:290:TYR:O	1.98	0.63
4:K:319:ALA:HB3	4:K:532:PRO:HG3	1.80	0.63
3:P:85:ARG:NE	3:P:86:TRP:H	1.94	0.63
6:X:27:SER:N	6:X:31:GLY:O	2.30	0.63
6:X:327:THR:HG23	6:X:330:GLN:HE21	1.63	0.63
6:Z:27:SER:N	6:Z:31:GLY:O	2.32	0.63
1:B:844:PRO:HA	1:B:1003:GLN:HA	1.80	0.63
2:C:902:ALA:HA	2:C:905:ILE:HG22	1.79	0.63
4:L:101:VAL:HG13	4:L:166:ILE:HG13	1.80	0.63
4:L:654:SER:HA	4:L:657:ILE:HD12	1.79	0.63
5:O:130:TYR:HA	5:O:133:LEU:HD21	1.81	0.63
5:O:686:THR:HA	5:O:689:ARG:HG3	1.80	0.63
5:O:906:ALA:O	5:O:910:LEU:HG	1.99	0.63
5:O:988:LEU:HD13	5:O:990:TRP:HE1	1.63	0.63
3:P:353:GLN:O	3:P:357:ILE:HG13	1.98	0.63
6:X:205:GLN:H	6:X:267:LYS:HZ3	1.45	0.63
1:B:537:GLN:O	1:B:541:VAL:HG23	1.99	0.63
1:B:849:VAL:HG13	1:B:999:ILE:HG13	1.81	0.63
2:C:850:THR:O	2:C:997:LEU:HA	1.99	0.63
2:C:1062:PRO:HB2	2:C:1065:LEU:HG	1.80	0.63
4:K:122:ARG:HH11	4:L:151:SER:HB2	1.63	0.63
4:K:398:LYS:HB3	4:K:473:ILE:HG13	1.81	0.63
4:L:623:ASN:O	4:L:627:LEU:HG	1.99	0.63
4:M:571:GLN:HA	4:M:574:ILE:HG12	1.80	0.63
1:B:427:VAL:HG22	1:B:428:ARG:H	1.64	0.63
2:C:324:MET:HA	2:C:334:LEU:HD11	1.81	0.63
2:C:418:LYS:NZ	2:C:1225:LYS:O	2.32	0.63
4:K:284:LYS:HZ3	4:M:291:ASP:HA	1.63	0.63
4:K:645:LYS:HE2	4:M:308:PRO:HG2	1.80	0.63
4:L:248:GLN:HB3	4:L:257:VAL:HG12	1.81	0.63

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:B:468:ARG:HA	1:B:471:ARG:HG2	1.80	0.62
1:B:599:LEU:O	1:B:832:GLN:NE2	2.32	0.62
1:B:681:PRO:HA	1:B:684:TRP:CD2	2.33	0.62
2:C:591:PHE:CE2	2:C:595:LEU:HD11	2.34	0.62
4:K:288:GLN:HE22	4:K:552:ILE:HG12	1.64	0.62
4:K:375:PRO:HA	4:K:452:TYR:CD2	2.33	0.62
5:O:206:VAL:HA	5:O:209:ARG:HD2	1.81	0.62
5:O:422:LEU:O	5:O:698:ASP:N	2.31	0.62
5:O:637:SER:H	5:O:656:PHE:HB2	1.64	0.62
5:O:1073:ILE:HB	5:O:1076:VAL:HG22	1.80	0.62
3:P:242:ASN:HD22	3:P:244:ARG:NH2	1.97	0.62
6:Z:6:PRO:HD3	6:Z:56:GLY:HA2	1.81	0.62
4:M:405:GLN:O	4:M:407:LYS:NZ	2.28	0.62
5:O:637:SER:OG	5:O:638:TYR:N	2.31	0.62
6:Y:167:TRP:HZ2	6:Y:259:ARG:HB3	1.62	0.62
1:B:787:GLN:OE1	1:B:787:GLN:N	2.32	0.62
1:B:930:THR:O	1:B:933:THR:OG1	2.17	0.62
2:C:883:THR:HA	2:C:886:LEU:HD12	1.82	0.62
3:D:331:GLN:HE22	3:D:333:ARG:H	1.47	0.62
4:L:292:LEU:HD11	4:L:545:LYS:HB2	1.81	0.62
4:L:565:SER:HA	4:L:568:ILE:HD13	1.82	0.62
4:L:584:ARG:HB3	6:Z:70:HIS:CG	2.34	0.62
5:O:15:SER:HB3	5:O:311:GLY:HA3	1.82	0.62
5:O:303:VAL:O	5:O:307:LEU:HG	1.99	0.62
6:X:82:TYR:HD2	6:X:86:GLN:HE22	1.47	0.62
6:X:251:HIS:CD2	6:X:254:LEU:HG	2.31	0.62
6:Y:165:GLN:O	6:Y:165:GLN:NE2	2.32	0.62
6:Y:331:ALA:O	6:Y:333:ILE:N	2.32	0.62
1:B:1230:GLU:OE1	1:B:1230:GLU:N	2.32	0.62
2:C:520:ILE:O	2:C:524:ASN:HB2	2.00	0.62
2:C:791:SER:OG	2:C:794:ARG:NH2	2.32	0.62
2:C:1204:TYR:HB2	2:C:1206:ASP:CG	2.20	0.62
3:D:9:LYS:NZ	3:D:154:ASN:O	2.24	0.62
3:D:244:ARG:HB3	3:D:357:ILE:HG21	1.80	0.62
4:K:266:ALA:HA	4:K:269:ALA:HB3	1.82	0.62
4:K:567:PRO:HG2	4:K:568:ILE:HD12	1.81	0.62
4:L:137:TYR:HD1	4:L:140:LEU:HD12	1.64	0.62
4:L:405:GLN:HE22	4:L:422:ALA:HA	1.62	0.62
4:M:362:ARG:O	4:M:365:THR:OG1	2.16	0.62
6:Y:175:ASP:OD2	6:Y:251:HIS:ND1	2.21	0.62
1:B:440:MET:HA	2:C:862:LEU:HB2	1.81	0.62

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:B:1109:ILE:HG23	1:B:1138:ARG:HE	1.64	0.62
1:B:1158:ASN:OD1	1:B:1159:ALA:N	2.30	0.62
2:C:847:VAL:HG13	2:C:871:VAL:HG13	1.80	0.62
2:C:1080:ASP:OD1	2:C:1082:ARG:NH1	2.31	0.62
3:D:29:LEU:HB2	3:D:30:LEU:HD12	1.80	0.62
4:K:405:GLN:HB2	4:K:421:GLN:HG2	1.80	0.62
4:M:19:VAL:O	4:M:248:GLN:N	2.21	0.62
4:M:369:ASN:HA	4:M:467:TYR:HD2	1.64	0.62
5:O:1137:THR:OG1	5:O:1140:GLY:N	2.33	0.62
3:P:22:ASN:ND2	3:P:389:SER:OG	2.33	0.62
6:Y:94:HIS:HA	6:Y:97:ARG:HH11	1.63	0.62
6:Y:243:VAL:HA	6:Y:349:PHE:HB2	1.80	0.62
1:B:1045:ILE:HG13	1:B:1139:ILE:HG23	1.82	0.62
2:C:589:SER:O	2:C:592:ARG:N	2.33	0.62
2:C:649:PHE:CE2	2:C:705:TRP:HD1	2.17	0.62
2:C:935:VAL:O	2:C:939:SER:OG	2.14	0.62
4:L:569:GLN:HA	4:L:572:LEU:HD12	1.82	0.62
4:M:653:SER:O	4:M:656:SER:OG	2.14	0.62
6:X:142:PHE:HE2	6:X:266:GLY:HA2	1.65	0.62
6:Y:281:LYS:HG2	6:Y:283:TYR:CZ	2.34	0.62
1:B:1188:HIS:HA	1:B:1220:ILE:HD12	1.82	0.62
2:C:1149:HIS:O	2:C:1182:VAL:N	2.28	0.62
3:D:7:LEU:HD12	3:D:8:PHE:H	1.65	0.62
3:D:273:MET:N	3:D:328:HIS:O	2.32	0.62
4:K:445:THR:OG1	4:L:327:LYS:NZ	2.32	0.62
4:K:631:VAL:O	4:K:635:LEU:HG	1.99	0.62
4:L:426:ASN:ND2	4:L:449:ALA:O	2.31	0.62
4:M:22:PRO:HB3	4:M:202:ASN:HA	1.81	0.62
4:M:23:SER:N	4:M:26:THR:OG1	2.32	0.62
5:O:329:SER:O	5:O:332:SER:OG	2.10	0.62
6:X:120:ARG:NH1	6:X:125:GLY:O	2.28	0.62
6:X:295:VAL:O	6:X:299:VAL:HG23	2.00	0.62
6:Y:182:ASN:OD1	6:Y:183:PHE:HD1	1.83	0.62
1:B:592:ARG:HA	1:B:595:LEU:HD12	1.82	0.62
1:B:1127:ASP:N	1:B:1127:ASP:OD1	2.32	0.62
2:C:269:VAL:N	2:C:303:ARG:O	2.25	0.62
2:C:764:TRP:O	2:C:768:VAL:HG23	2.00	0.62
3:D:66:TYR:O	3:D:69:SER:OG	2.14	0.62
4:K:34:LEU:HD13	4:M:279:LEU:HD21	1.82	0.62
4:L:428:VAL:HG12	4:L:448:LEU:HD22	1.82	0.62
4:M:409:PRO:HB2	4:M:412:LEU:HG	1.81	0.62

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
5:O:220:VAL:HB	5:O:287:TRP:O	2.00	0.62
3:P:93:ARG:O	3:P:107:ALA:N	2.32	0.62
6:Y:71:HIS:C	6:Y:72:ARG:HD2	2.20	0.62
6:Y:172:LEU:HG	6:Y:252:PHE:HE2	1.65	0.62
6:Y:318:GLY:HA3	6:Y:322:GLY:HA3	1.82	0.62
1:B:476:PRO:HA	1:B:479:ILE:HD12	1.81	0.62
1:B:799:LYS:HD2	1:B:802:LEU:HD13	1.81	0.62
1:B:1109:ILE:HD12	1:B:1138:ARG:HE	1.63	0.62
1:B:1201:SER:OG	1:B:1202:THR:N	2.33	0.62
4:L:428:VAL:HB	4:L:448:LEU:HD13	1.82	0.62
4:M:240:LEU:HD12	4:M:244:ASN:HB3	1.82	0.62
5:O:2:ALA:HB1	5:O:341:GLU:HG3	1.81	0.62
5:O:561:ASP:OD1	5:O:562:LEU:N	2.32	0.62
3:P:123:ARG:HB3	3:P:125:TYR:CZ	2.35	0.62
6:X:241:GLU:O	6:X:247:ARG:NE	2.33	0.62
6:X:262:THR:O	6:X:269:PRO:HB3	2.00	0.62
1:B:245:ALA:HA	1:B:248:LEU:HB2	1.82	0.62
1:B:1094:ILE:N	1:B:1102:VAL:O	2.33	0.62
2:C:407:TYR:CE1	2:C:431:VAL:HG22	2.34	0.62
2:C:557:SER:O	2:C:561:SER:N	2.33	0.62
2:C:602:GLY:H	2:C:832:GLN:HA	1.65	0.62
2:C:1110:PHE:CE2	2:C:1115:TRP:HB2	2.34	0.62
4:K:537:ARG:HA	4:K:540:ILE:HD12	1.81	0.62
4:K:565:SER:HA	4:K:568:ILE:HD13	1.81	0.62
4:M:154:GLN:HG2	4:M:158:ASN:ND2	2.15	0.62
5:O:1067:ASP:OD2	5:O:1070:ALA:N	2.31	0.62
6:X:52:MET:HA	6:X:55:LEU:HD22	1.82	0.62
6:Y:19:PHE:CG	6:Y:291:LYS:HE3	2.35	0.62
6:Y:99:MET:O	6:Y:103:VAL:HG23	2.00	0.62
1:B:464:ARG:NH2	1:B:1022:ASP:OD1	2.33	0.61
1:B:480:GLU:O	1:B:484:THR:HG23	1.99	0.61
1:B:984:LEU:O	1:B:988:LEU:HG	2.00	0.61
2:C:1111:PRO:HD2	2:C:1114:LEU:HD12	1.82	0.61
2:C:1150:TYR:HA	2:C:1182:VAL:O	1.99	0.61
4:K:36:LEU:HD11	4:M:280:GLU:HB3	1.81	0.61
4:K:321:TYR:OH	4:K:537:ARG:NH2	2.26	0.61
4:K:389:GLU:OE2	4:K:484:THR:OG1	2.18	0.61
4:M:405:GLN:HA	4:M:465:ASN:O	2.00	0.61
4:M:431:TYR:OH	4:M:433:GLU:OE1	2.16	0.61
5:O:372:LYS:HB2	5:O:375:LEU:HG	1.81	0.61
3:P:211:ARG:O	3:P:215:ILE:HG12	2.00	0.61

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
6:Z:199:GLY:O	6:Z:203:GLN:N	2.32	0.61
1:B:355:ARG:HG3	1:B:952:ILE:HB	1.81	0.61
1:B:878:ASP:OD1	1:B:880:ARG:N	2.32	0.61
3:D:110:GLN:O	3:D:114:GLN:NE2	2.34	0.61
4:K:435:SER:OG	4:K:437:PHE:O	2.19	0.61
4:L:505:VAL:HG12	4:L:506:LYS:HD2	1.83	0.61
4:M:22:PRO:O	4:M:207:GLU:HA	1.99	0.61
4:M:34:LEU:HD23	4:M:35:SER:N	2.15	0.61
5:O:371:ARG:HH22	5:O:375:LEU:HB2	1.65	0.61
6:X:18:ALA:O	6:X:21:GLY:N	2.32	0.61
2:C:801:LYS:HG3	2:C:802:LEU:HG	1.83	0.61
2:C:974:LYS:HD3	2:C:979:LEU:HD23	1.82	0.61
3:D:211:ARG:O	3:D:215:ILE:HG13	1.99	0.61
4:L:336:MET:HB2	4:L:366:ARG:HD3	1.80	0.61
4:M:327:LYS:HZ3	4:M:328:ILE:HG13	1.65	0.61
5:O:423:ARG:HB3	5:O:697:VAL:HG22	1.81	0.61
5:O:760:PRO:HB2	5:O:764:ARG:HH12	1.65	0.61
5:O:770:ARG:HE	5:O:860:CYS:HA	1.65	0.61
5:O:858:SER:HA	5:O:861:TRP:HE1	1.64	0.61
5:O:948:ARG:HG2	5:O:959:PRO:HA	1.82	0.61
5:O:1032:GLY:HA3	5:O:1040:CYS:HA	1.82	0.61
3:P:161:ILE:HG13	3:P:273:MET:HB3	1.81	0.61
6:X:39:GLN:HE21	6:X:41:ASP:HA	1.64	0.61
2:C:465:TRP:CE2	2:C:469:LEU:HD11	2.36	0.61
4:K:607:PRO:O	4:K:610:ILE:N	2.31	0.61
4:L:615:ALA:HB1	4:L:618:LEU:HB2	1.82	0.61
5:O:1167:ASN:HB2	5:O:1269:SER:O	2.00	0.61
3:P:238:TYR:O	3:P:254:SER:OG	2.14	0.61
6:Y:216:LEU:N	6:Y:217:GLU:OE1	2.33	0.61
6:Z:158:ASP:OD2	6:Z:161:THR:HB	1.99	0.61
1:B:938:ILE:O	1:B:956:ARG:NH1	2.34	0.61
1:B:942:GLN:O	3:P:105:VAL:HB	2.00	0.61
2:C:376:THR:HG23	2:C:1259:ARG:NH1	2.16	0.61
3:D:21:ILE:HG22	3:D:22:ASN:H	1.65	0.61
3:D:277:PRO:HA	3:D:279:TRP:NE1	2.15	0.61
4:K:647:ALA:O	4:K:651:LEU:HG	2.00	0.61
4:L:428:VAL:HG13	4:L:430:LEU:HD21	1.81	0.61
4:M:47:PRO:HA	4:M:148:VAL:HG22	1.82	0.61
1:B:491:VAL:HG13	1:B:1275:PRO:HA	1.83	0.61
1:B:1077:PHE:CZ	1:B:1108:TRP:HB3	2.35	0.61
4:K:327:LYS:HB3	4:K:329:ASP:OD1	1.99	0.61

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
4:L:410:PHE:HA	4:L:413:TRP:CE2	2.36	0.61
4:L:547:ALA:HA	4:L:550:ILE:HD12	1.81	0.61
4:M:68:SER:OG	4:M:97:ASP:OD1	2.19	0.61
5:O:27:LEU:HB2	5:O:108:ASN:HA	1.83	0.61
5:O:691:LEU:HD12	5:O:692:PRO:HD2	1.82	0.61
5:O:708:THR:HB	5:O:755:THR:HG23	1.81	0.61
5:O:940:LEU:HD12	5:O:950:ARG:H	1.65	0.61
5:O:1096:ALA:O	5:O:1099:TYR:OH	2.12	0.61
6:Y:142:PHE:HB2	6:Y:265:LEU:HD12	1.80	0.61
6:Y:325:ASN:OD1	6:Y:329:GLN:NE2	2.32	0.61
1:B:401:LYS:O	1:B:404:SER:OG	2.14	0.61
1:B:773:LYS:HZ3	1:B:797:LEU:HB2	1.66	0.61
2:C:400:GLU:N	2:C:400:GLU:OE1	2.34	0.61
2:C:714:PRO:O	2:C:837:ARG:NH2	2.34	0.61
2:C:845:THR:N	2:C:1002:GLN:O	2.34	0.61
4:K:160:GLN:HG3	4:K:161:LYS:H	1.64	0.61
4:L:124:PHE:CE1	4:L:128:LEU:HD21	2.36	0.61
4:L:666:ASP:O	4:L:669:SER:OG	2.18	0.61
4:M:12:ASN:ND2	4:M:14:THR:O	2.33	0.61
5:O:203:THR:HA	5:O:231:HIS:ND1	2.15	0.61
3:P:231:GLN:O	3:P:259:HIS:ND1	2.22	0.61
6:Z:20:GLU:HG3	6:Z:22:ARG:H	1.66	0.61
1:B:513:GLU:HG2	1:B:1016:PRO:HG3	1.82	0.61
1:B:589:SER:HG	1:B:592:ARG:NH2	1.99	0.61
3:D:93:ARG:HG3	3:D:112:LEU:HD22	1.82	0.61
3:D:237:PHE:HB3	3:D:253:LEU:HD11	1.83	0.61
4:L:174:TYR:O	4:L:177:SER:OG	2.17	0.61
4:L:324:ARG:NH2	4:L:487:ASP:OD1	2.33	0.61
4:L:445:THR:N	4:M:325:THR:O	2.33	0.61
5:O:494:TYR:HE2	5:O:541:VAL:HG22	1.64	0.61
6:Z:133:TRP:CD1	6:Z:134:LEU:HD23	2.35	0.61
1:B:731:ASN:HD21	1:B:737:GLU:H	1.48	0.61
1:B:985:GLU:O	1:B:989:SER:OG	2.17	0.61
1:B:1107:ASN:C	1:B:1108:TRP:HD1	2.03	0.61
1:B:1112:LEU:HG	1:B:1116:GLN:HE21	1.66	0.61
3:D:327:LEU:HD12	3:D:328:HIS:H	1.66	0.61
4:K:408:ILE:O	4:K:413:TRP:NE1	2.32	0.61
4:K:513:VAL:HG12	4:K:514:VAL:H	1.66	0.61
4:M:51:ILE:H	4:M:63:LEU:HD22	1.64	0.61
4:M:174:TYR:O	4:M:178:ILE:HG12	2.01	0.61
5:O:88:GLU:O	5:O:92:VAL:HG23	2.00	0.61

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
5:O:594:LEU:O	5:O:597:SER:OG	2.08	0.61
5:O:1193:TYR:CE2	5:O:1286:LEU:HB2	2.36	0.61
6:X:185:MET:SD	6:X:188:PRO:HB3	2.41	0.61
6:Y:160:ASP:OD1	6:Y:161:THR:N	2.34	0.61
2:C:1059:LEU:HB3	2:C:1204:TYR:HA	1.83	0.61
2:C:1125:GLN:HB2	2:C:1126:PHE:CE1	2.36	0.61
3:D:76:LEU:O	3:D:79:PRO:HD2	2.01	0.61
4:K:238:VAL:O	4:K:242:LYS:N	2.32	0.61
4:M:405:GLN:HE21	4:M:421:GLN:HB2	1.65	0.61
3:P:170:MET:O	3:P:175:VAL:HG11	2.00	0.61
6:Y:200:ASP:O	6:Y:203:GLN:HB3	2.00	0.61
1:B:490:TYR:O	2:C:674:ARG:NH1	2.34	0.60
1:B:790:VAL:O	1:B:794:ARG:HG3	2.01	0.60
1:B:1217:PRO:HG2	2:C:1087:MET:HE2	1.82	0.60
3:D:143:VAL:O	3:D:147:LEU:HG	2.01	0.60
3:D:229:HIS:HD2	3:D:232:ASN:HB2	1.66	0.60
3:D:285:LEU:O	3:D:288:SER:OG	2.17	0.60
4:K:33:SER:OG	4:K:34:LEU:N	2.34	0.60
4:K:349:VAL:O	4:K:356:ASN:HA	2.01	0.60
4:L:105:GLU:HG2	4:L:108:ILE:HD12	1.82	0.60
4:L:189:GLU:HA	4:L:192:LEU:HG	1.82	0.60
4:M:230:ARG:HB2	4:M:231:ARG:NH1	2.16	0.60
4:M:318:PRO:HB3	4:M:496:SER:HA	1.83	0.60
5:O:6:GLY:HA3	5:O:333:GLN:O	2.00	0.60
6:X:36:ILE:HG12	6:X:152:PRO:HA	1.83	0.60
1:B:941:THR:OG1	1:B:994:MET:SD	2.59	0.60
1:B:1090:ALA:O	1:B:1092:PRO:HD3	1.99	0.60
3:D:23:ASP:OD1	3:D:24:GLU:N	2.33	0.60
5:O:223:HIS:CE1	5:O:225:ASP:HB2	2.36	0.60
3:P:137:PRO:HA	3:P:140:LYS:HE2	1.83	0.60
6:X:297:LYS:HG3	6:X:298:LEU:HD12	1.83	0.60
1:B:373:ASN:HB2	1:B:1259:ARG:HD3	1.83	0.60
1:B:822:ILE:O	1:B:826:LEU:N	2.28	0.60
2:C:293:GLN:HE21	2:C:411:MET:HA	1.66	0.60
2:C:684:TRP:HB2	2:C:840:ARG:HH21	1.66	0.60
2:C:763:ASN:O	2:C:767:ARG:HG2	2.02	0.60
3:D:204:GLU:HG2	3:D:206:ASP:H	1.66	0.60
4:L:230:ARG:HB2	4:L:231:ARG:NH1	2.16	0.60
4:M:337:ILE:HG13	4:M:361:LEU:HD21	1.83	0.60
4:M:426:ASN:ND2	4:M:449:ALA:H	1.99	0.60
4:M:429:GLN:NE2	4:M:430:LEU:O	2.34	0.60

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
4:M:609:THR:OG1	4:M:612:THR:OG1	2.19	0.60
5:O:276:VAL:HG21	5:O:279:LEU:HD23	1.83	0.60
5:O:963:MET:O	5:O:966:LEU:HB3	2.01	0.60
5:O:1144:ASP:OD1	5:O:1182:VAL:HG13	2.01	0.60
6:X:164:ASP:O	6:X:168:THR:OG1	2.13	0.60
6:Y:179:LEU:CB	6:Y:363:ILE:HG12	2.31	0.60
1:B:408:PRO:HB2	1:B:423:VAL:HG21	1.83	0.60
1:B:946:ASP:HB3	3:P:31:ARG:HH22	1.65	0.60
1:B:1119:THR:HA	1:B:1122:PHE:HB2	1.84	0.60
3:D:8:PHE:HD2	3:D:127:CYS:HG	1.48	0.60
4:K:190:ILE:O	4:K:194:VAL:HG23	2.00	0.60
4:K:348:GLN:HA	4:K:357:TRP:O	2.00	0.60
4:L:32:PRO:HG3	4:M:38:PRO:HB2	1.82	0.60
4:L:459:LYS:HG3	4:L:464:MET:HB2	1.83	0.60
4:M:50:ALA:HA	4:M:63:LEU:HD22	1.82	0.60
5:O:1150:TYR:HB3	5:O:1151:TYR:CE1	2.36	0.60
3:P:50:ARG:HH11	3:P:51:GLY:H	1.50	0.60
2:C:418:LYS:NZ	2:C:1212:THR:O	2.31	0.60
2:C:995:THR:HG22	2:C:996:GLN:H	1.67	0.60
4:K:113:LYS:NZ	4:L:40:MET:O	2.34	0.60
4:K:188:TRP:HE1	4:K:192:LEU:HD22	1.66	0.60
4:K:671:HIS:HE1	4:M:578:GLU:HB2	1.66	0.60
3:P:90:ARG:NE	3:P:114:GLN:O	2.34	0.60
3:P:123:ARG:HH21	3:P:125:TYR:HE2	1.50	0.60
6:Y:92:THR:O	6:Y:96:LYS:HG2	2.00	0.60
6:Z:185:MET:H	6:Z:261:THR:HA	1.66	0.60
1:B:332:ILE:HD11	1:B:347:ARG:HD2	1.84	0.60
1:B:445:SER:OG	1:B:447:TRP:O	2.19	0.60
1:B:858:THR:OG1	1:B:942:GLN:OE1	2.19	0.60
1:B:1111:PRO:HB2	1:B:1114:LEU:HG	1.84	0.60
3:D:219:LEU:HD23	3:D:239:PHE:HD2	1.66	0.60
4:L:137:TYR:CD1	4:L:140:LEU:HD12	2.35	0.60
4:L:334:LEU:HB3	4:L:366:ARG:HH12	1.67	0.60
5:O:393:PRO:HB2	5:O:740:SER:HB2	1.83	0.60
5:O:680:ARG:HG2	5:O:684:LEU:HD11	1.84	0.60
6:X:211:ASP:HB2	6:X:213:ARG:HE	1.67	0.60
1:B:257:THR:OG1	1:B:372:LEU:N	2.31	0.60
1:B:356:ALA:O	1:B:359:LEU:HB3	2.02	0.60
1:B:520:ILE:HA	1:B:523:MET:SD	2.41	0.60
2:C:1056:TRP:HZ3	2:C:1058:PRO:HA	1.67	0.60
5:O:632:LEU:HB3	5:O:664:LEU:HB2	1.83	0.60

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
5:O:1156:PHE:HA	5:O:1165:ILE:HA	1.83	0.60
6:Z:21:GLY:O	6:Z:34:LYS:NZ	2.35	0.60
6:Z:39:GLN:HE22	6:Z:79:HIS:HD2	1.49	0.60
1:B:547:SER:OG	1:B:810:TYR:OH	2.12	0.60
3:D:23:ASP:O	3:D:26:SER:OG	2.13	0.60
4:L:602:ALA:O	4:L:606:ASP:N	2.31	0.60
4:M:103:VAL:HB	4:M:164:PRO:HG2	1.83	0.60
5:O:68:LEU:O	5:O:73:TYR:OH	2.10	0.60
5:O:80:ASP:N	5:O:80:ASP:OD1	2.32	0.60
3:P:230:GLN:HE21	3:P:258:ASN:HD21	1.50	0.60
6:Y:318:GLY:O	6:Y:322:GLY:N	2.28	0.60
6:Z:236:ARG:HA	6:Z:239:ARG:HB2	1.83	0.60
1:B:406:MET:HG3	1:B:453:THR:OG1	2.01	0.60
2:C:748:ALA:HA	2:C:813:GLN:HG2	1.84	0.60
2:C:809:MET:SD	2:C:810:TYR:N	2.75	0.60
2:C:897:THR:OG1	2:C:898:ASN:N	2.35	0.60
2:C:1180:PHE:HE2	2:C:1199:ILE:HG21	1.64	0.60
4:K:243:ARG:HH12	4:K:262:VAL:H	1.49	0.60
4:K:521:SER:N	4:K:611:ILE:O	2.35	0.60
4:M:20:PHE:HD1	4:M:247:ILE:HG12	1.66	0.60
5:O:421:PRO:HB3	5:O:699:ASP:HB3	1.84	0.60
5:O:680:ARG:O	5:O:684:LEU:HG	2.01	0.60
5:O:931:ASP:HB3	5:O:1015:TYR:CZ	2.37	0.60
5:O:1052:VAL:HG11	5:O:1066:GLU:OE1	2.02	0.60
5:O:1126:ILE:HG13	5:O:1146:THR:O	2.02	0.60
3:P:110:GLN:HG3	3:P:111:VAL:N	2.16	0.60
2:C:836:VAL:HG11	2:C:843:VAL:HG13	1.83	0.60
2:C:1074:VAL:HA	2:C:1107:ASN:O	2.01	0.60
4:K:396:GLY:H	4:K:430:LEU:HD23	1.66	0.60
4:K:426:ASN:ND2	6:Z:62:GLN:HG2	2.17	0.60
4:L:38:PRO:HA	4:L:41:LEU:HB2	1.84	0.60
5:O:981:TRP:HD1	5:O:1116:SER:O	1.85	0.60
5:O:1068:VAL:HA	5:O:1071:ALA:HB2	1.84	0.60
5:O:1122:PRO:HG2	5:O:1150:TYR:CG	2.37	0.60
5:O:1238:TRP:CE3	5:O:1286:LEU:HB3	2.37	0.60
6:Z:209:THR:OG1	6:Z:211:ASP:OD2	2.18	0.60
1:B:362:LEU:O	1:B:366:VAL:HG23	2.01	0.59
1:B:636:LEU:HD22	1:B:644:ALA:HB1	1.83	0.59
2:C:438:GLN:HB3	3:D:201:ARG:HE	1.65	0.59
2:C:600:TYR:CE1	2:C:830:PRO:HA	2.37	0.59
3:D:19:VAL:HG22	3:D:21:ILE:HD11	1.84	0.59

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
4:K:471:THR:HG22	4:K:488:VAL:HG12	1.84	0.59
4:K:522:TYR:CG	4:K:527:LEU:HG	2.36	0.59
4:M:27:SER:HA	4:M:202:ASN:HB3	1.84	0.59
5:O:389:ASP:OD1	5:O:390:THR:N	2.35	0.59
5:O:442:ASN:OD1	5:O:445:HIS:N	2.33	0.59
5:O:1155:THR:O	5:O:1165:ILE:HD12	2.02	0.59
1:B:542:LEU:HD13	1:B:545:ARG:HD3	1.83	0.59
2:C:699:ALA:O	2:C:703:ARG:N	2.31	0.59
3:D:42:PHE:HA	3:D:64:ARG:CZ	2.32	0.59
4:K:398:LYS:HG2	4:K:427:TYR:HE2	1.66	0.59
4:L:398:LYS:HB3	4:L:473:ILE:HD11	1.83	0.59
4:M:303:SER:OG	4:M:305:VAL:O	2.20	0.59
4:M:348:GLN:HB2	4:M:356:ASN:OD1	2.02	0.59
4:M:485:GLN:NE2	4:M:487:ASP:O	2.36	0.59
5:O:547:VAL:O	5:O:549:GLN:NE2	2.36	0.59
5:O:739:LYS:HB3	5:O:754:ILE:HD11	1.84	0.59
3:P:98:ALA:HB3	3:P:102:ASP:N	2.16	0.59
6:X:53:HIS:HD2	6:X:75:GLN:HE22	1.49	0.59
1:B:542:LEU:HA	1:B:545:ARG:HD3	1.85	0.59
1:B:1096:ASP:O	1:B:1099:GLY:N	2.34	0.59
2:C:646:VAL:HG11	2:C:684:TRP:CD1	2.38	0.59
3:D:48:LEU:O	3:D:50:ARG:N	2.36	0.59
3:D:165:ARG:HA	3:D:269:SER:HA	1.84	0.59
4:K:381:ASP:O	4:K:492:LEU:HB2	2.02	0.59
4:K:389:GLU:HB2	4:K:392:TRP:HB3	1.85	0.59
4:M:18:ASN:HA	4:M:250:MET:N	2.17	0.59
4:M:435:SER:HB3	4:M:442:ILE:HB	1.84	0.59
5:O:121:GLY:HA3	5:O:156:PHE:HZ	1.67	0.59
5:O:143:ILE:HG13	5:O:144:GLY:H	1.66	0.59
5:O:426:TYR:HB2	5:O:695:GLY:H	1.67	0.59
5:O:783:GLN:CD	5:O:785:ARG:HE	2.05	0.59
3:P:229:HIS:CD2	3:P:231:GLN:H	2.20	0.59
6:Y:324:TYR:O	6:Y:328:MET:HG3	2.01	0.59
6:Z:236:ARG:CZ	6:Z:240:LYS:HE3	2.32	0.59
1:B:992:PRO:HD2	1:B:993:ARG:NH1	2.18	0.59
4:M:367:VAL:HG21	4:M:467:TYR:HB3	1.83	0.59
4:M:393:ASP:O	4:M:397:LYS:NZ	2.24	0.59
3:P:21:ILE:HD11	3:P:71:LEU:HG	1.84	0.59
3:P:79:PRO:HG2	3:P:80:PHE:CE1	2.37	0.59
6:Y:44:VAL:HG22	6:Y:49:VAL:HG13	1.84	0.59
6:Y:168:THR:HA	6:Y:252:PHE:HZ	1.67	0.59

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
6:Y:263:PRO:HB3	6:Y:267:LYS:C	2.23	0.59
1:B:338:LEU:O	1:B:968:TRP:NE1	2.35	0.59
1:B:454:SER:H	1:B:1254:TYR:HA	1.67	0.59
1:B:501:VAL:HG13	1:B:502:ASN:H	1.68	0.59
1:B:550:GLN:HE22	1:B:552:ASP:HB2	1.65	0.59
1:B:653:ALA:O	1:B:671:GLN:NE2	2.28	0.59
1:B:1144:TYR:HD2	1:B:1145:PRO:O	1.86	0.59
2:C:332:ILE:HD11	2:C:336:LYS:HE2	1.85	0.59
2:C:470:ALA:HB3	3:D:180:MET:HG3	1.84	0.59
2:C:528:ASN:HB2	2:C:531:VAL:HG23	1.83	0.59
2:C:662:ILE:O	2:C:671:GLN:NE2	2.35	0.59
2:C:1186:SER:OG	2:C:1187:ASP:N	2.34	0.59
4:K:591:ILE:O	4:K:595:ILE:HG12	2.03	0.59
5:O:326:TRP:CE3	5:O:327:LEU:HD23	2.37	0.59
5:O:1007:ARG:O	5:O:1011:LEU:HG	2.01	0.59
3:P:96:TRP:HA	3:P:103:GLY:O	2.03	0.59
6:X:315:LEU:HB3	6:X:323:TRP:CD1	2.38	0.59
2:C:445:SER:OG	2:C:446:GLU:N	2.36	0.59
2:C:880:ARG:HG3	2:C:881:ALA:N	2.17	0.59
3:D:77:GLN:HA	3:D:80:PHE:HD1	1.68	0.59
4:K:153:ARG:HD3	4:M:134:SER:HB2	1.84	0.59
4:L:28:SER:HB2	4:L:244:ASN:HA	1.85	0.59
4:L:100:LEU:HB3	4:L:165:VAL:HG11	1.84	0.59
4:L:258:MET:SD	4:M:99:PRO:HD3	2.43	0.59
4:L:290:MET:O	4:L:294:THR:HG23	2.01	0.59
4:L:375:PRO:HG3	4:L:454:PRO:HD2	1.83	0.59
5:O:212:TRP:CE2	3:P:138:ARG:HD2	2.38	0.59
5:O:223:HIS:CE1	5:O:278:ARG:HH21	2.21	0.59
5:O:912:LYS:O	5:O:915:SER:OG	2.15	0.59
3:P:240:GLN:OE1	3:P:241:CYS:N	2.36	0.59
6:Z:30:GLU:HG2	6:Z:36:ILE:HB	1.84	0.59
1:B:372:LEU:HD21	1:B:465:TRP:HE1	1.67	0.59
1:B:401:LYS:O	1:B:405:ILE:HG12	2.03	0.59
1:B:1078:GLY:N	1:B:1097:GLU:OE1	2.36	0.59
4:L:506:LYS:HZ1	6:Z:313:TYR:C	2.05	0.59
4:M:334:LEU:HD21	4:M:366:ARG:HB3	1.83	0.59
5:O:297:LEU:O	5:O:301:LEU:HG	2.01	0.59
6:X:79:HIS:HA	6:X:82:TYR:CE1	2.37	0.59
6:X:196:ARG:HB2	6:X:355:PRO:CA	2.33	0.59
6:X:288:GLY:HA3	6:X:335:LEU:HD23	1.83	0.59
1:B:553:PRO:HA	1:B:556:ILE:HD13	1.83	0.59

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:B:1150:TYR:HB3	1:B:1184:ILE:HG12	1.85	0.59
2:C:394:LEU:HD22	2:C:403:TYR:CE1	2.38	0.59
2:C:685:PRO:HG2	2:C:688:PHE:HB2	1.84	0.59
2:C:747:PRO:O	2:C:813:GLN:NE2	2.36	0.59
2:C:961:THR:HG22	2:C:965:PHE:CZ	2.37	0.59
4:L:227:SER:OG	4:L:229:ILE:HG22	2.03	0.59
5:O:384:LEU:HD12	5:O:387:LEU:HD11	1.85	0.59
5:O:642:LYS:NZ	5:O:650:GLU:OE2	2.33	0.59
6:Z:78:ARG:HD3	6:Z:78:ARG:H	1.68	0.59
6:Z:211:ASP:H	6:Z:216:LEU:HD11	1.67	0.59
1:B:880:ARG:O	1:B:884:VAL:HG23	2.03	0.59
2:C:745:HIS:HA	2:C:812:GLN:O	2.03	0.59
2:C:880:ARG:O	2:C:884:VAL:HG23	2.02	0.59
3:D:407:THR:O	3:D:410:THR:OG1	2.18	0.59
4:L:299:GLU:HB2	4:M:650:LYS:HD2	1.84	0.59
5:O:62:PRO:HB3	5:O:113:VAL:HG13	1.85	0.59
5:O:113:VAL:HG12	5:O:117:ASN:HD21	1.66	0.59
5:O:743:ILE:HB	5:O:786:THR:HA	1.84	0.59
5:O:804:HIS:O	5:O:808:THR:OG1	2.16	0.59
3:P:279:TRP:O	3:P:282:ASP:N	2.36	0.59
6:X:332:PRO:O	6:X:336:THR:OG1	2.21	0.59
6:Z:196:ARG:HH21	6:Z:352:LEU:HB3	1.68	0.59
1:B:353:MET:SD	1:B:353:MET:N	2.73	0.59
1:B:1096:ASP:OD1	1:B:1097:GLU:N	2.36	0.59
2:C:880:ARG:O	2:C:883:THR:OG1	2.16	0.59
3:D:95:VAL:H	3:D:105:VAL:H	1.51	0.59
4:K:334:LEU:HD23	4:K:368:VAL:HG22	1.85	0.59
1:B:372:LEU:HD21	1:B:465:TRP:NE1	2.17	0.58
1:B:475:ASN:ND2	3:P:182:TYR:OH	2.36	0.58
1:B:628:ASP:O	1:B:632:LEU:HG	2.02	0.58
1:B:969:VAL:O	1:B:973:MET:HG3	2.02	0.58
2:C:1107:ASN:HA	2:C:1136:ARG:O	2.03	0.58
4:K:578:GLU:O	4:K:584:ARG:NH2	2.36	0.58
4:L:282:LYS:O	4:L:286:THR:HG23	2.02	0.58
4:M:392:TRP:HB2	4:M:482:ASN:HB3	1.84	0.58
4:M:514:VAL:HG13	4:M:517:GLU:HB2	1.84	0.58
5:O:777:PRO:O	5:O:781:GLU:N	2.27	0.58
6:X:22:ARG:HH22	6:X:96:LYS:HD3	1.68	0.58
6:X:204:THR:HA	6:X:267:LYS:HZ3	1.68	0.58
6:X:223:TYR:HB2	6:X:225:TYR:CZ	2.37	0.58
1:B:454:SER:N	1:B:1253:LEU:O	2.35	0.58

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:B:540:SER:OG	1:B:592:ARG:NH2	2.28	0.58
1:B:843:VAL:O	1:B:1004:TYR:N	2.35	0.58
1:B:965:PHE:O	1:B:969:VAL:HG23	2.03	0.58
2:C:580:LYS:HD2	2:C:627:TRP:CE3	2.39	0.58
2:C:622:SER:N	2:C:625:ASN:OD1	2.34	0.58
4:K:30:ALA:HB1	4:M:639:SER:HB2	1.84	0.58
4:M:546:ILE:O	4:M:550:ILE:HG12	2.03	0.58
4:M:664:PHE:O	4:M:668:VAL:HG23	2.03	0.58
5:O:200:TYR:CG	5:O:232:HIS:HB3	2.37	0.58
5:O:1063:PHE:HB3	5:O:1072:MET:HG2	1.85	0.58
6:X:181:PRO:HA	6:X:363:ILE:HG23	1.86	0.58
1:B:441:ASN:HD21	2:C:864:LEU:HG	1.68	0.58
2:C:296:PHE:HB3	2:C:419:ILE:HD12	1.85	0.58
2:C:448:VAL:HG12	2:C:450:VAL:HG13	1.85	0.58
2:C:550:GLN:N	2:C:890:LYS:O	2.34	0.58
3:D:9:LYS:HD2	3:D:10:THR:N	2.18	0.58
3:D:95:VAL:N	3:D:104:LEU:HB3	2.18	0.58
3:D:367:PHE:O	3:D:371:LYS:HG2	2.03	0.58
4:K:428:VAL:HG23	4:K:448:LEU:HD13	1.85	0.58
4:M:12:ASN:ND2	4:M:14:THR:OG1	2.26	0.58
4:M:581:TYR:CD1	6:Y:73:CYS:HB3	2.38	0.58
5:O:1040:CYS:HB2	5:O:1093:PHE:CZ	2.37	0.58
6:X:86:GLN:H	6:X:86:GLN:CD	2.07	0.58
6:X:222:VAL:O	6:X:356:VAL:HA	2.03	0.58
6:X:224:ASP:HB3	6:X:354:TYR:CD1	2.39	0.58
6:Z:29:GLN:HE22	6:Z:39:GLN:HB3	1.68	0.58
1:B:488:GLN:NE2	1:B:905:ILE:HD13	2.18	0.58
2:C:548:PRO:O	2:C:892:PRO:HD3	2.03	0.58
2:C:595:LEU:O	2:C:598:TRP:HB2	2.03	0.58
2:C:674:ARG:HB3	2:C:676:SER:OG	2.03	0.58
2:C:1041:ALA:HB3	2:C:1144:TYR:CE2	2.38	0.58
4:K:102:VAL:HA	4:K:165:VAL:HG22	1.84	0.58
4:K:635:LEU:HA	4:K:638:LYS:HD2	1.85	0.58
5:O:253:ASP:O	5:O:257:LEU:HG	2.03	0.58
5:O:1193:TYR:CZ	5:O:1286:LEU:HB2	2.38	0.58
5:O:1236:ARG:CZ	5:O:1236:ARG:HA	2.32	0.58
6:X:288:GLY:HA2	6:X:291:LYS:HG3	1.85	0.58
1:B:505:MET:HB2	1:B:507:TYR:HE1	1.69	0.58
2:C:457:LEU:O	2:C:460:SER:OG	2.18	0.58
3:D:80:PHE:HA	3:D:84:HIS:CD2	2.39	0.58
3:D:212:MET:O	3:D:216:ILE:HG12	2.03	0.58

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
4:K:282:LYS:HE2	4:K:640:LEU:HA	1.86	0.58
4:L:320:ASP:HA	4:L:493:LEU:O	2.03	0.58
4:L:410:PHE:HB2	4:L:467:TYR:CZ	2.39	0.58
4:L:581:TYR:O	6:Z:71:HIS:N	2.29	0.58
4:M:522:TYR:O	4:M:613:GLN:NE2	2.37	0.58
5:O:509:ARG:NH1	5:O:543:PRO:O	2.37	0.58
5:O:616:ILE:O	5:O:651:LEU:HB2	2.03	0.58
5:O:947:LYS:HB3	5:O:960:TYR:CE1	2.39	0.58
3:P:340:CYS:HA	3:P:343:LEU:HD12	1.85	0.58
6:X:87:PHE:O	6:X:91:VAL:HG23	2.04	0.58
6:Y:273:SER:OG	6:Y:281:LYS:N	2.37	0.58
6:Z:218:TRP:HB2	6:Z:267:LYS:HZ3	1.68	0.58
1:B:426:CYS:O	2:C:1079:ARG:NH2	2.31	0.58
1:B:473:ASN:HA	1:B:506:PRO:HA	1.86	0.58
1:B:1041:ALA:O	1:B:1143:ALA:HA	2.02	0.58
2:C:372:LEU:HD22	2:C:374:ARG:NH2	2.18	0.58
2:C:1114:LEU:O	2:C:1117:MET:HB2	2.03	0.58
3:D:155:LEU:HB2	3:D:161:ILE:HD11	1.86	0.58
4:K:351:ASP:OD2	4:K:355:THR:OG1	2.21	0.58
4:K:650:LYS:HD2	4:M:299:GLU:OE2	2.03	0.58
4:L:144:VAL:O	4:L:163:VAL:HG12	2.03	0.58
4:L:631:VAL:O	4:L:634:SER:OG	2.16	0.58
3:P:110:GLN:HG3	3:P:111:VAL:HG23	1.85	0.58
6:Y:130:GLU:HG2	6:Y:360:ASP:HB3	1.86	0.58
6:Y:191:ALA:O	6:Y:195:VAL:HG21	2.03	0.58
3:D:233:THR:O	3:D:260:SER:HA	2.04	0.58
4:K:284:LYS:NZ	4:M:291:ASP:HA	2.18	0.58
4:K:582:GLY:HA2	6:X:68:LEU:HD21	1.85	0.58
4:L:621:LYS:O	4:L:625:ILE:HG13	2.03	0.58
5:O:286:ARG:NH1	5:O:286:ARG:O	2.36	0.58
6:Z:184:MET:HA	6:Z:261:THR:HA	1.85	0.58
1:B:1094:ILE:HG22	1:B:1102:VAL:O	2.04	0.58
2:C:696:PRO:O	2:C:700:PRO:HB3	2.03	0.58
3:D:244:ARG:NH1	3:D:249:GLU:OE2	2.35	0.58
4:K:442:ILE:HD12	4:L:325:THR:HA	1.85	0.58
4:M:370:LEU:HD12	4:M:452:TYR:CD1	2.39	0.58
3:P:26:SER:O	3:P:30:LEU:N	2.20	0.58
6:X:50:VAL:HG22	6:X:57:VAL:HA	1.86	0.58
6:Y:323:TRP:O	6:Y:327:THR:OG1	2.16	0.58
1:B:257:THR:O	1:B:372:LEU:HB2	2.03	0.58
1:B:624:GLU:HA	1:B:627:TRP:HD1	1.69	0.58

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
4:K:49:ILE:HB	4:K:66:MET:HB2	1.85	0.58
4:K:228:LEU:HG	4:K:231:ARG:HD3	1.85	0.58
4:K:261:ALA:H	4:L:65:ARG:HH22	1.52	0.58
4:K:371:ASP:O	4:K:372:GLN:HB2	2.02	0.58
4:L:20:PHE:HB3	4:L:210:CYS:SG	2.44	0.58
4:L:154:GLN:OE1	4:L:158:ASN:N	2.37	0.58
5:O:430:TRP:HA	5:O:1000:THR:HG23	1.86	0.58
5:O:456:LEU:HG	5:O:457:PRO:HD2	1.86	0.58
5:O:948:ARG:HA	5:O:959:PRO:HA	1.86	0.58
5:O:971:ARG:HH22	5:O:1028:MET:N	2.01	0.58
5:O:981:TRP:O	5:O:983:PRO:HD3	2.04	0.58
3:P:25:LEU:HA	3:P:28:HIS:NE2	2.19	0.58
6:X:220:VAL:H	6:X:358:ILE:HG23	1.68	0.58
6:Y:74:ASN:ND2	6:Y:76:GLN:HB2	2.19	0.58
6:Z:128:LEU:HB3	6:Z:363:ILE:HG23	1.85	0.58
3:D:123:ARG:NH1	3:D:125:TYR:HH	2.01	0.58
3:D:195:CYS:O	3:D:198:SER:OG	2.17	0.58
3:D:243:ARG:NH2	3:D:246:ASP:OD2	2.37	0.58
4:K:378:PHE:HZ	4:K:452:TYR:HB3	1.69	0.58
5:O:553:ASP:HA	5:O:559:ILE:HD13	1.84	0.58
5:O:611:SER:HG	5:O:657:GLY:N	2.01	0.58
5:O:620:THR:OG1	5:O:623:VAL:HG23	2.04	0.58
6:X:196:ARG:HB2	6:X:355:PRO:HA	1.85	0.58
6:X:309:GLU:HB3	6:X:313:TYR:CZ	2.38	0.58
2:C:1084:SER:N	2:C:1093:MET:O	2.24	0.57
4:K:224:PRO:O	4:K:227:SER:OG	2.19	0.57
4:L:143:TYR:CE1	4:L:164:PRO:HB3	2.38	0.57
4:L:194:VAL:O	4:L:197:THR:OG1	2.18	0.57
4:L:597:SER:OG	4:L:598:ARG:N	2.36	0.57
4:M:347:ILE:HD13	4:M:416:ALA:HB1	1.85	0.57
5:O:397:LEU:O	5:O:399:GLN:NE2	2.36	0.57
5:O:590:ILE:H	5:O:590:ILE:HD12	1.69	0.57
5:O:1012:MET:HB3	5:O:1016:MET:HE2	1.85	0.57
3:P:78:ILE:HD11	3:P:91:PHE:HB2	1.86	0.57
6:X:277:THR:OG1	6:X:279:ASN:O	2.22	0.57
6:Z:15:ILE:HG22	6:Z:19:PHE:CZ	2.39	0.57
2:C:553:PRO:HB2	2:C:583:PRO:HB2	1.85	0.57
2:C:949:LEU:HD12	2:C:952:ILE:HD11	1.84	0.57
4:K:13:VAL:HG23	4:K:14:THR:HG23	1.86	0.57
4:K:378:PHE:CZ	4:K:452:TYR:HB3	2.40	0.57
4:M:174:TYR:O	4:M:177:SER:OG	2.16	0.57

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
5:O:8:ARG:HD3	5:O:315:ALA:CB	2.33	0.57
5:O:641:ILE:HG23	5:O:652:PHE:HB2	1.86	0.57
5:O:933:VAL:HG21	5:O:942:ILE:HG21	1.86	0.57
5:O:1180:THR:OG1	5:O:1181:LEU:N	2.32	0.57
3:P:283:VAL:O	3:P:286:ILE:N	2.38	0.57
3:P:314:ARG:HA	3:P:320:THR:HG21	1.86	0.57
6:Z:127:SER:HA	6:Z:364:LEU:HG	1.86	0.57
6:Z:199:GLY:HA2	6:Z:202:ARG:HH21	1.69	0.57
1:B:604:VAL:HG13	1:B:873:VAL:HG13	1.84	0.57
1:B:882:ILE:O	1:B:886:LEU:HG	2.04	0.57
1:B:1166:TRP:CZ2	1:B:1178:VAL:HG22	2.39	0.57
3:D:9:LYS:HA	3:D:124:VAL:HG12	1.86	0.57
4:K:348:GLN:HB3	4:K:356:ASN:ND2	2.18	0.57
4:M:282:LYS:O	4:M:286:THR:HG23	2.03	0.57
4:M:370:LEU:HA	4:M:373:ILE:HD11	1.86	0.57
5:O:80:ASP:HA	5:O:83:GLU:HG3	1.87	0.57
5:O:426:TYR:CG	5:O:694:PHE:HA	2.38	0.57
5:O:990:TRP:CE2	5:O:991:THR:HG23	2.39	0.57
5:O:1098:ILE:HG13	5:O:1117:PHE:O	2.04	0.57
5:O:1099:TYR:HB2	5:O:1117:PHE:CZ	2.39	0.57
6:X:253:GLY:HA2	6:X:275:MET:SD	2.45	0.57
1:B:381:ASP:N	1:B:389:ALA:O	2.30	0.57
1:B:1023:CYS:O	1:B:1027:THR:HG23	2.04	0.57
1:B:1110:PHE:CE1	1:B:1115:TRP:HB2	2.38	0.57
2:C:451:PHE:CE1	2:C:1256:VAL:HB	2.39	0.57
2:C:465:TRP:CH2	2:C:469:LEU:HD21	2.39	0.57
2:C:689:MET:HE2	2:C:840:ARG:HE	1.68	0.57
2:C:987:LEU:HG	2:C:992:PRO:HG3	1.86	0.57
4:K:297:GLU:HG3	4:K:300:ILE:HG22	1.85	0.57
4:K:540:ILE:HG12	4:K:608:SER:HA	1.87	0.57
4:M:212:MET:SD	4:M:212:MET:N	2.77	0.57
5:O:208:ASP:HA	5:O:211:PHE:CD2	2.39	0.57
5:O:1187:LEU:HD21	5:O:1217:LEU:HD21	1.86	0.57
1:B:946:ASP:OD2	1:B:947:ARG:N	2.38	0.57
1:B:1093:MET:HB3	1:B:1102:VAL:N	2.19	0.57
2:C:268:ILE:HG21	2:C:1210:PHE:CD2	2.38	0.57
2:C:810:TYR:CD1	2:C:814:LEU:HB2	2.39	0.57
4:K:653:SER:HB2	4:M:299:GLU:HG2	1.86	0.57
4:L:187:LYS:NZ	4:L:191:ASP:OD2	2.36	0.57
4:M:18:ASN:HB3	4:M:249:TRP:HE3	1.69	0.57
4:M:263:ASN:O	4:M:267:ALA:N	2.22	0.57

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
5:O:261:LEU:HA	5:O:264:PHE:CD2	2.39	0.57
5:O:435:LEU:HB2	5:O:644:PHE:CD2	2.38	0.57
5:O:807:LEU:HD12	5:O:810:MET:HB2	1.86	0.57
5:O:1238:TRP:CE2	5:O:1288:PRO:HB3	2.39	0.57
3:P:167:ASP:OD2	3:P:267:ASN:ND2	2.38	0.57
6:X:289:THR:O	6:X:292:LEU:HB3	2.03	0.57
6:Y:328:MET:HB2	6:Y:329:GLN:NE2	2.20	0.57
1:B:272:VAL:O	1:B:300:ALA:N	2.37	0.57
2:C:338:LEU:HD23	2:C:339:LEU:HB2	1.87	0.57
3:D:316:ILE:HB	3:D:320:THR:N	2.20	0.57
3:D:353:GLN:HA	3:D:356:GLN:CD	2.24	0.57
4:M:519:ILE:O	4:M:521:SER:OG	2.21	0.57
5:O:826:LEU:HD11	5:O:886:VAL:HG13	1.87	0.57
5:O:833:GLU:HG2	5:O:855:ALA:HB3	1.86	0.57
3:P:238:TYR:CE1	3:P:256:SER:HB2	2.39	0.57
3:P:273:MET:O	3:P:329:GLY:HA2	2.04	0.57
3:P:353:GLN:HA	3:P:356:GLN:HE21	1.69	0.57
6:Z:288:GLY:HA2	6:Z:291:LYS:HD2	1.86	0.57
1:B:969:VAL:O	1:B:972:SER:OG	2.17	0.57
3:D:6:PHE:HB3	3:D:127:CYS:HB2	1.87	0.57
4:K:193:ARG:O	4:K:197:THR:HG23	2.04	0.57
4:K:515:PRO:O	4:K:518:LEU:N	2.37	0.57
4:L:404:PHE:CE2	4:L:419:ILE:HB	2.40	0.57
4:M:49:ILE:HG23	4:M:64:ARG:HB3	1.85	0.57
5:O:192:LEU:HB2	5:O:195:TYR:CD2	2.39	0.57
5:O:528:LEU:O	5:O:532:PRO:HD2	2.04	0.57
3:P:3:ARG:NH2	3:P:299:SER:O	2.34	0.57
3:P:143:VAL:O	3:P:147:LEU:HG	2.04	0.57
1:B:357:ASN:HA	1:B:360:HIS:CD2	2.40	0.57
1:B:1115:TRP:O	1:B:1119:THR:OG1	2.20	0.57
2:C:649:PHE:CE2	2:C:706:ALA:HB2	2.40	0.57
3:D:31:ARG:HD2	3:D:45:TRP:HZ2	1.69	0.57
3:D:238:TYR:CE1	3:D:240:GLN:HB2	2.40	0.57
3:D:404:ARG:HH22	3:D:406:ASN:HA	1.69	0.57
4:M:546:ILE:HD12	4:M:549:ALA:HB3	1.86	0.57
5:O:632:LEU:HD23	5:O:635:ILE:HD11	1.86	0.57
3:P:178:LEU:HB3	3:P:212:MET:CE	2.35	0.57
6:X:310:LYS:HD3	6:X:311:ILE:HD13	1.87	0.57
6:Y:309:GLU:OE1	6:Y:309:GLU:N	2.35	0.57
1:B:713:TRP:O	1:B:837:ARG:NE	2.31	0.57
1:B:809:MET:HA	1:B:812:GLN:HB2	1.87	0.57

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:B:1052:SER:OG	1:B:1053:THR:N	2.38	0.57
3:D:172:SER:O	3:D:175:VAL:HB	2.05	0.57
3:D:246:ASP:N	3:D:249:GLU:OE1	2.30	0.57
3:D:330:PHE:HB2	3:D:407:THR:HG23	1.86	0.57
4:L:182:LEU:HA	4:L:185:LEU:HD21	1.87	0.57
4:M:35:SER:OG	4:M:110:ASN:ND2	2.38	0.57
4:M:338:PRO:HG3	4:M:411:GLU:HB3	1.85	0.57
4:M:572:LEU:O	4:M:575:SER:OG	2.20	0.57
5:O:172:TYR:CG	5:O:187:LEU:HB3	2.40	0.57
3:P:273:MET:N	3:P:328:HIS:O	2.33	0.57
6:Y:182:ASN:CG	6:Y:362:MET:H	2.08	0.57
6:Z:168:THR:HA	6:Z:171:ASN:ND2	2.20	0.57
1:B:355:ARG:NE	1:B:952:ILE:O	2.36	0.57
2:C:504:LEU:HB3	2:C:1264:THR:OG1	2.05	0.57
2:C:896:VAL:HB	2:C:899:VAL:HB	1.86	0.57
2:C:1005:ASN:O	3:P:235:ARG:NH1	2.38	0.57
2:C:1122:PHE:HB3	2:C:1126:PHE:CE2	2.40	0.57
4:K:436:SER:HB3	4:L:324:ARG:NH1	2.19	0.57
5:O:9:LEU:CD2	5:O:314:LEU:HA	2.34	0.57
5:O:112:ASN:ND2	5:O:146:LEU:HD11	2.19	0.57
5:O:425:ASP:O	5:O:794:LEU:N	2.31	0.57
5:O:426:TYR:OH	5:O:692:PRO:O	2.22	0.57
5:O:506:LEU:C	5:O:507:ARG:HH11	2.08	0.57
5:O:584:GLY:O	5:O:587:ASP:N	2.37	0.57
5:O:638:TYR:HD2	5:O:664:LEU:HD13	1.69	0.57
5:O:752:LEU:HD12	5:O:753:THR:H	1.70	0.57
3:P:47:SER:OG	3:P:51:GLY:O	2.23	0.57
3:P:151:THR:O	3:P:155:LEU:HG	2.04	0.57
6:Y:221:MET:N	6:Y:221:MET:SD	2.78	0.57
6:Z:217:GLU:O	6:Z:267:LYS:NZ	2.37	0.57
4:K:119:GLU:OE2	4:L:151:SER:N	2.37	0.56
4:K:397:LYS:HB2	4:K:472:PHE:HE2	1.69	0.56
4:K:450:TYR:CG	4:K:451:ASN:N	2.73	0.56
4:L:403:VAL:HG13	4:L:405:GLN:NE2	2.19	0.56
4:L:457:LEU:HD13	4:L:466:TYR:CD2	2.40	0.56
4:L:588:PRO:O	4:L:592:LEU:HG	2.05	0.56
4:M:441:SER:OG	4:M:442:ILE:N	2.38	0.56
4:M:660:TRP:CE2	4:M:664:PHE:HZ	2.23	0.56
5:O:464:ASP:OD1	5:O:464:ASP:N	2.38	0.56
5:O:815:VAL:O	5:O:819:VAL:N	2.38	0.56
5:O:949:TYR:CE1	5:O:960:TYR:HB3	2.40	0.56

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
6:Z:17:ASN:HB3	6:Z:23:VAL:HB	1.87	0.56
2:C:317:ASP:OD1	2:C:318:ARG:N	2.33	0.56
2:C:418:LYS:HZ3	2:C:1226:HIS:HA	1.70	0.56
2:C:488:GLN:HE21	2:C:821:VAL:HG12	1.70	0.56
3:D:94:LEU:C	3:D:104:LEU:HB3	2.26	0.56
4:L:159:PHE:CE2	4:L:161:LYS:HD2	2.40	0.56
4:L:218:LEU:O	4:L:222:GLN:HG2	2.05	0.56
4:M:339:LYS:O	4:M:341:MET:HG2	2.05	0.56
5:O:430:TRP:O	5:O:477:LYS:N	2.38	0.56
5:O:587:ASP:O	5:O:591:SER:N	2.38	0.56
5:O:778:ARG:O	5:O:782:VAL:HG23	2.05	0.56
6:X:96:LYS:O	6:X:100:LEU:HG	2.05	0.56
6:Y:181:PRO:N	6:Y:363:ILE:HG13	2.20	0.56
6:Z:94:HIS:CE1	6:Z:153:LEU:HB3	2.40	0.56
1:B:416:VAL:HA	2:C:1082:ARG:HH21	1.69	0.56
1:B:593:VAL:HG23	1:B:594:ALA:H	1.71	0.56
1:B:635:ALA:O	1:B:639:THR:HG23	2.06	0.56
1:B:878:ASP:OD1	1:B:879:ALA:N	2.38	0.56
1:B:1003:GLN:NE2	1:B:1004:TYR:HB3	2.21	0.56
1:B:1247:LEU:N	1:B:1249:GLU:OE1	2.39	0.56
2:C:298:GLU:OE2	2:C:1214:SER:N	2.38	0.56
2:C:675:ALA:HA	2:C:678:PHE:CE2	2.39	0.56
2:C:1051:GLN:HA	2:C:1196:VAL:HG22	1.87	0.56
2:C:1149:HIS:N	2:C:1180:PHE:O	2.38	0.56
3:D:231:GLN:OE1	3:D:258:ASN:ND2	2.38	0.56
3:D:331:GLN:N	3:D:407:THR:OG1	2.37	0.56
4:K:212:MET:O	4:K:215:VAL:HB	2.06	0.56
4:K:234:LYS:O	4:K:238:VAL:HG23	2.05	0.56
4:K:244:ASN:ND2	4:K:247:ILE:HB	2.20	0.56
4:K:427:TYR:CD2	4:K:473:ILE:HD11	2.39	0.56
4:K:502:GLU:N	4:K:502:GLU:OE1	2.38	0.56
4:M:217:LYS:HZ2	4:M:675:PRO:HD2	1.70	0.56
5:O:200:TYR:CE1	5:O:229:ASN:HB3	2.39	0.56
5:O:206:VAL:HA	5:O:209:ARG:HH11	1.68	0.56
5:O:271:ARG:NE	5:O:273:ALA:O	2.38	0.56
5:O:316:ARG:NH1	5:O:785:ARG:O	2.38	0.56
5:O:424:PRO:HD2	5:O:696:TYR:O	2.06	0.56
5:O:1101:MET:HB2	5:O:1114:LEU:HB2	1.86	0.56
5:O:1153:LEU:HD23	5:O:1198:ILE:HG22	1.86	0.56
5:O:1154:MET:O	5:O:1199:ARG:N	2.25	0.56
6:X:174:ILE:HA	6:X:179:LEU:HD13	1.87	0.56

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
6:Z:41:ASP:HA	6:Z:53:HIS:CE1	2.40	0.56
6:Z:223:TYR:HA	6:Z:356:VAL:HA	1.86	0.56
1:B:257:THR:OG1	1:B:318:ARG:NH2	2.36	0.56
2:C:473:ASN:O	2:C:508:ARG:NH2	2.38	0.56
2:C:511:ASN:HB3	2:C:730:ALA:HB2	1.88	0.56
2:C:622:SER:OG	2:C:624:GLU:OE1	2.23	0.56
2:C:1044:ASP:HA	2:C:1141:MET:HG3	1.88	0.56
4:K:29:THR:HG23	4:M:636:ARG:HB3	1.88	0.56
4:K:337:ILE:N	4:K:365:THR:O	2.25	0.56
5:O:208:ASP:HA	5:O:211:PHE:CE2	2.40	0.56
5:O:687:ILE:HG23	5:O:992:ARG:HD2	1.87	0.56
5:O:992:ARG:O	5:O:996:LEU:N	2.24	0.56
5:O:1227:ASP:HA	5:O:1279:VAL:O	2.06	0.56
6:X:295:VAL:HG12	6:X:298:LEU:HD22	1.87	0.56
1:B:303:ARG:HB3	1:B:1208:SER:O	2.06	0.56
1:B:580:LYS:HG3	1:B:631:ILE:HD11	1.87	0.56
1:B:987:LEU:HD12	1:B:987:LEU:H	1.71	0.56
1:B:1042:ARG:HG2	1:B:1043:GLY:H	1.71	0.56
1:B:1069:ARG:HA	1:B:1074:VAL:HG21	1.88	0.56
2:C:340:ASN:O	2:C:343:THR:OG1	2.16	0.56
2:C:446:GLU:HG3	2:C:447:TRP:HD1	1.70	0.56
3:D:151:THR:O	3:D:155:LEU:HG	2.06	0.56
4:K:370:LEU:HA	4:K:373:ILE:HD13	1.87	0.56
4:L:225:ASP:HA	4:L:230:ARG:HE	1.71	0.56
4:L:429:GLN:NE2	4:L:430:LEU:O	2.39	0.56
4:M:67:THR:OG1	4:M:97:ASP:OD1	2.22	0.56
5:O:66:LEU:HD13	5:O:170:ARG:HB3	1.88	0.56
5:O:1181:LEU:HD12	5:O:1182:VAL:H	1.70	0.56
3:P:78:ILE:HD13	3:P:117:LEU:HD23	1.87	0.56
3:P:123:ARG:HB3	3:P:125:TYR:CE1	2.41	0.56
6:Z:17:ASN:O	6:Z:21:GLY:N	2.34	0.56
1:B:880:ARG:O	1:B:883:THR:OG1	2.23	0.56
2:C:691:ILE:H	2:C:691:ILE:HD12	1.71	0.56
2:C:727:PHE:HD1	2:C:914:VAL:HG11	1.71	0.56
4:K:48:TRP:HE3	4:K:63:LEU:HG	1.70	0.56
4:K:69:LYS:HG2	4:K:97:ASP:HB2	1.86	0.56
4:K:182:LEU:HA	4:K:185:LEU:HD23	1.88	0.56
4:K:388:LYS:NZ	4:K:393:ASP:OD1	2.38	0.56
4:L:239:ALA:O	4:L:243:ARG:HG2	2.06	0.56
4:M:175:VAL:O	4:M:179:GLN:HG3	2.06	0.56
4:M:615:ALA:HB1	4:M:618:LEU:HB2	1.88	0.56

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
5:O:690:GLN:HE21	5:O:992:ARG:HE	1.52	0.56
5:O:1004:ALA:O	5:O:1007:ARG:NH1	2.38	0.56
5:O:1204:ARG:NH1	5:O:1205:ASP:OD1	2.39	0.56
3:P:277:PRO:HA	3:P:279:TRP:CD1	2.41	0.56
3:P:341:ASP:O	3:P:344:ARG:HG3	2.05	0.56
6:Z:326:ARG:HA	6:Z:329:GLN:NE2	2.20	0.56
1:B:255:LEU:HG	1:B:256:VAL:HG23	1.86	0.56
1:B:465:TRP:NE1	1:B:469:LEU:HD11	2.20	0.56
1:B:520:ILE:HA	1:B:523:MET:HG3	1.87	0.56
2:C:312:SER:OG	2:C:313:ASN:O	2.23	0.56
2:C:850:THR:O	2:C:997:LEU:HD13	2.04	0.56
3:D:367:PHE:CD1	3:D:371:LYS:HE2	2.41	0.56
4:K:369:ASN:HA	4:K:467:TYR:CD2	2.40	0.56
4:L:40:MET:HB3	4:L:106:HIS:CE1	2.40	0.56
4:L:64:ARG:NH2	4:L:70:ASP:OD2	2.37	0.56
4:M:137:TYR:CZ	4:M:141:LEU:HD21	2.41	0.56
4:M:152:ALA:HB3	4:M:153:ARG:NH1	2.21	0.56
5:O:212:TRP:NE1	3:P:137:PRO:HG2	2.18	0.56
5:O:1192:LYS:HZ1	5:O:1220:ILE:H	1.54	0.56
3:P:165:ARG:HD3	3:P:166:VAL:N	2.21	0.56
3:P:186:THR:O	3:P:189:GLU:HG2	2.06	0.56
6:Y:180:VAL:HG12	6:Y:225:TYR:HE1	1.69	0.56
2:C:416:VAL:HG23	2:C:1217:PRO:HG3	1.87	0.56
2:C:851:ARG:O	2:C:994:MET:HB2	2.05	0.56
2:C:951:TRP:CZ3	2:C:1042:ARG:HG3	2.40	0.56
2:C:1075:HIS:H	2:C:1108:TRP:HA	1.69	0.56
4:K:215:VAL:O	4:K:219:LEU:HG	2.05	0.56
4:L:398:LYS:HG2	4:L:429:GLN:HA	1.86	0.56
4:M:135:PRO:HB2	4:M:174:TYR:OH	2.06	0.56
5:O:186:PRO:HG2	5:O:188:PHE:HE1	1.71	0.56
5:O:408:ASP:O	5:O:412:LEU:HG	2.05	0.56
5:O:614:VAL:HG13	5:O:616:ILE:HD11	1.86	0.56
5:O:642:LYS:HZ1	5:O:644:PHE:HA	1.71	0.56
5:O:778:ARG:NH1	5:O:781:GLU:OE1	2.30	0.56
5:O:1231:SER:HA	5:O:1275:PRO:O	2.06	0.56
6:Y:208:ARG:HH12	6:Y:287:LYS:H	1.54	0.56
6:Y:247:ARG:O	6:Y:348:LYS:NZ	2.38	0.56
1:B:372:LEU:HD21	1:B:465:TRP:CD1	2.41	0.56
1:B:488:GLN:HE21	1:B:905:ILE:HD13	1.71	0.56
2:C:461:ILE:HG23	2:C:1023:CYS:SG	2.46	0.56
3:D:237:PHE:CD1	3:D:255:CYS:HA	2.41	0.56

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:D:361:ALA:O	3:D:365:GLN:HG3	2.05	0.56
4:L:250:MET:CG	4:L:254:GLU:HB2	2.34	0.56
4:M:314:ILE:HD13	4:M:538:CYS:HB2	1.88	0.56
5:O:47:ARG:HB3	5:O:53:ASN:C	2.26	0.56
5:O:888:CYS:SG	5:O:891:SER:OG	2.63	0.56
5:O:1170:LYS:O	5:O:1186:LYS:N	2.35	0.56
6:X:248:ASP:OD1	6:X:348:LYS:NZ	2.38	0.56
6:Y:54:CYS:HB3	6:Y:73:CYS:SG	2.46	0.56
6:Y:198:GLU:HA	6:Y:202:ARG:HE	1.70	0.56
2:C:464:ARG:HH22	2:C:1026:LEU:HD13	1.71	0.56
2:C:719:ILE:HG12	2:C:720:ARG:H	1.71	0.56
4:L:301:ILE:HG12	4:L:624:TRP:CZ2	2.41	0.56
4:M:28:SER:OG	4:M:243:ARG:O	2.17	0.56
4:M:122:ARG:O	4:M:125:LEU:HG	2.06	0.56
4:M:548:GLU:O	4:M:552:ILE:HG13	2.06	0.56
5:O:1203:SER:OG	5:O:1204:ARG:N	2.39	0.56
3:P:107:ALA:HB3	3:P:112:LEU:HD23	1.88	0.56
6:Y:60:SER:OG	6:Y:61:LEU:N	2.39	0.56
6:Y:299:VAL:HG23	6:Y:320:MET:HG3	1.87	0.56
6:Z:90:ARG:NH2	6:Z:151:ASP:O	2.39	0.56
1:B:318:ARG:N	1:B:371:TYR:OH	2.39	0.55
1:B:446:GLU:C	1:B:1261:ALA:HB3	2.26	0.55
1:B:576:SER:OG	1:B:577:ILE:N	2.36	0.55
1:B:931:LEU:O	1:B:935:VAL:HG23	2.05	0.55
2:C:394:LEU:HB2	2:C:403:TYR:OH	2.06	0.55
2:C:494:THR:O	2:C:1270:VAL:N	2.22	0.55
2:C:856:THR:HG22	2:C:857:ILE:H	1.71	0.55
3:D:213:THR:O	3:D:217:LEU:HG	2.06	0.55
4:K:48:TRP:CD1	4:K:65:ARG:HG3	2.41	0.55
4:K:479:THR:OG1	4:K:481:THR:OG1	2.14	0.55
4:K:617:VAL:HG22	4:K:618:LEU:HD22	1.88	0.55
4:L:199:LEU:HD13	4:L:665:LEU:HD11	1.87	0.55
4:M:169:THR:O	4:M:172:THR:OG1	2.18	0.55
4:M:396:GLY:O	4:M:429:GLN:NE2	2.36	0.55
5:O:394:VAL:O	5:O:741:ILE:HG12	2.06	0.55
5:O:617:ASN:ND2	5:O:649:VAL:O	2.39	0.55
6:Z:7:ASN:O	6:Z:11:ILE:HG13	2.06	0.55
1:B:259:ASP:OD2	1:B:312:SER:OG	2.24	0.55
2:C:313:ASN:OD1	2:C:314:VAL:N	2.39	0.55
2:C:476:PRO:O	2:C:479:ILE:HB	2.06	0.55
2:C:682:HIS:HB3	2:C:1004:TYR:CE2	2.41	0.55

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:D:334:ARG:HH11	3:D:337:THR:HG21	1.71	0.55
4:K:42:ASN:HB3	4:M:113:LYS:HG3	1.88	0.55
4:K:68:SER:OG	4:K:97:ASP:N	2.27	0.55
4:K:383:GLY:HA2	4:K:443:ILE:HG23	1.89	0.55
4:M:392:TRP:CE3	6:X:278:GLY:HA3	2.41	0.55
4:M:633:THR:O	4:M:637:THR:HG23	2.06	0.55
5:O:832:PRO:O	5:O:856:GLN:HG2	2.06	0.55
3:P:262:GLN:HE22	3:P:264:ARG:HB3	1.71	0.55
3:P:373:GLU:O	3:P:377:ARG:NH1	2.38	0.55
6:X:251:HIS:CE1	6:X:253:GLY:H	2.24	0.55
6:Y:312:ARG:HH11	6:Y:316:GLY:HA2	1.70	0.55
6:Z:300:ASP:N	6:Z:300:ASP:OD1	2.35	0.55
1:B:732:LEU:HD23	1:B:732:LEU:H	1.71	0.55
1:B:860:PRO:HA	1:B:863:SER:HB2	1.89	0.55
3:D:374:ALA:N	3:D:377:ARG:HH21	2.03	0.55
4:M:284:LYS:HZ3	4:M:287:GLU:HB2	1.70	0.55
5:O:575:TYR:HA	5:O:613:VAL:HG12	1.87	0.55
5:O:707:GLU:HB2	5:O:756:SER:OG	2.06	0.55
5:O:875:ASP:OD2	5:O:876:GLY:N	2.37	0.55
1:B:589:SER:HG	1:B:592:ARG:HH22	1.53	0.55
1:B:946:ASP:HB3	3:P:31:ARG:HH12	1.72	0.55
1:B:1092:PRO:O	1:B:1093:MET:HG3	2.07	0.55
1:B:1166:TRP:NE1	1:B:1176:PRO:O	2.39	0.55
2:C:300:ALA:O	2:C:1185:SER:OG	2.22	0.55
2:C:1035:TYR:CE2	2:C:1042:ARG:HB2	2.40	0.55
5:O:96:GLU:O	5:O:100:ILE:HD12	2.07	0.55
5:O:727:ILE:HD12	5:O:730:LEU:HD23	1.88	0.55
3:P:79:PRO:HG2	3:P:80:PHE:CZ	2.41	0.55
4:L:370:LEU:HB2	4:L:466:TYR:HB2	1.87	0.55
5:O:112:ASN:HD21	5:O:146:LEU:HD11	1.71	0.55
5:O:186:PRO:HG2	5:O:188:PHE:CE1	2.42	0.55
5:O:445:HIS:HB3	5:O:667:THR:HB	1.88	0.55
5:O:851:ILE:HG23	5:O:872:TYR:CE1	2.42	0.55
5:O:889:MET:HG2	5:O:890:LEU:HG	1.89	0.55
3:P:241:CYS:SG	3:P:243:ARG:HG2	2.46	0.55
6:Y:185:MET:HE3	6:Y:187:ASP:H	1.71	0.55
6:Z:79:HIS:HE1	6:Z:83:VAL:HA	1.71	0.55
1:B:600:TYR:CD2	1:B:830:PRO:HA	2.42	0.55
1:B:630:PHE:O	1:B:634:LEU:HG	2.07	0.55
1:B:701:ILE:HA	1:B:704:GLN:HE21	1.72	0.55
1:B:759:ASN:HB2	1:B:762:THR:H	1.71	0.55

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:B:764:TRP:HE3	1:B:767:ARG:HH11	1.53	0.55
2:C:265:SER:OG	2:C:309:ARG:N	2.35	0.55
2:C:506:PRO:HD2	2:C:507:TYR:CZ	2.41	0.55
2:C:555:ILE:O	2:C:559:THR:HG23	2.06	0.55
4:K:27:SER:OG	4:K:28:SER:N	2.40	0.55
4:K:64:ARG:NH1	4:K:67:THR:OG1	2.36	0.55
4:K:324:ARG:HA	4:K:366:ARG:NH1	2.22	0.55
4:K:639:SER:HA	4:L:31:VAL:O	2.07	0.55
4:L:169:THR:O	4:L:173:MET:HG2	2.07	0.55
4:L:284:LYS:HZ3	4:M:36:LEU:HD12	1.72	0.55
4:L:369:ASN:HA	4:L:467:TYR:CD2	2.41	0.55
4:M:206:GLY:HA3	4:M:658:GLN:HE22	1.72	0.55
4:M:264:ALA:HA	4:M:267:ALA:HB3	1.87	0.55
4:M:653:SER:O	4:M:657:ILE:HG12	2.06	0.55
5:O:327:LEU:O	5:O:331:VAL:HG23	2.07	0.55
5:O:491:LYS:HE2	5:O:538:ILE:HG13	1.87	0.55
5:O:904:ASP:OD1	5:O:961:SER:OG	2.18	0.55
5:O:908:GLN:HA	5:O:911:ILE:HG12	1.89	0.55
3:P:129:ASP:N	3:P:129:ASP:OD1	2.36	0.55
3:P:146:GLN:O	3:P:150:VAL:HG23	2.06	0.55
3:P:213:THR:HA	3:P:216:ILE:HG12	1.88	0.55
6:X:238:TYR:CZ	6:X:242:LEU:HD21	2.41	0.55
6:Y:238:TYR:HA	6:Y:241:GLU:CD	2.26	0.55
1:B:569:GLN:OE1	1:B:571:LEU:N	2.29	0.55
1:B:791:SER:OG	1:B:794:ARG:NH2	2.38	0.55
2:C:1166:TRP:CH2	2:C:1175:ILE:HD11	2.41	0.55
2:C:1233:ASN:HA	2:C:1236:THR:OG1	2.07	0.55
3:D:222:LEU:HB2	3:D:228:ILE:HG13	1.88	0.55
4:K:136:LYS:NZ	4:L:158:ASN:HA	2.22	0.55
4:L:371:ASP:OD1	4:L:459:LYS:NZ	2.36	0.55
4:M:327:LYS:HD2	4:M:328:ILE:H	1.71	0.55
4:M:373:ILE:HG13	4:M:452:TYR:CE2	2.41	0.55
5:O:115:VAL:O	5:O:119:LEU:HG	2.06	0.55
5:O:262:GLU:O	5:O:265:SER:OG	2.23	0.55
5:O:397:LEU:HD12	5:O:398:PRO:HD2	1.88	0.55
5:O:576:SER:OG	5:O:613:VAL:O	2.24	0.55
5:O:730:LEU:O	5:O:734:VAL:HG22	2.07	0.55
3:P:34:ASN:HD22	3:P:103:GLY:HA2	1.71	0.55
3:P:55:SER:H	3:P:59:PRO:HA	1.72	0.55
3:P:270:LEU:HB2	3:P:328:HIS:CE1	2.42	0.55
3:P:270:LEU:HD21	3:P:414:ILE:HG12	1.89	0.55

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
6:Z:7:ASN:HB3	6:Z:10:GLN:NE2	2.22	0.55
6:Z:206:PHE:HB3	6:Z:208:ARG:HG3	1.89	0.55
1:B:601:ASN:ND2	1:B:835:TYR:OH	2.39	0.55
1:B:815:ALA:O	1:B:819:LEU:HG	2.06	0.55
1:B:1029:GLU:HA	1:B:1032:ASN:HB2	1.89	0.55
1:B:1095:ARG:NH1	1:B:1099:GLY:O	2.40	0.55
2:C:340:ASN:HD21	2:C:353:MET:HG3	1.71	0.55
2:C:682:HIS:HB2	3:P:173:GLY:HA2	1.89	0.55
2:C:850:THR:OG1	2:C:998:ALA:HB3	2.06	0.55
2:C:1047:ILE:HD11	2:C:1137:ILE:HD12	1.89	0.55
2:C:1110:PHE:HE2	2:C:1115:TRP:HB2	1.70	0.55
4:K:12:ASN:HD21	4:K:16:ASP:N	2.04	0.55
4:K:153:ARG:O	4:K:157:ASN:ND2	2.39	0.55
4:M:108:ILE:HG12	4:M:137:TYR:HE2	1.72	0.55
4:M:426:ASN:HA	6:X:61:LEU:HD13	1.89	0.55
4:M:564:ASN:HB2	4:M:600:MET:SD	2.47	0.55
5:O:531:GLU:O	5:O:535:GLN:N	2.28	0.55
6:X:236:ARG:O	6:X:240:LYS:N	2.40	0.55
1:B:524:ASN:HD21	1:B:986:PRO:HG3	1.72	0.55
1:B:802:LEU:HG	1:B:804:LYS:NZ	2.21	0.55
1:B:943:TYR:HA	3:P:105:VAL:HG12	1.89	0.55
2:C:707:GLU:O	2:C:711:ARG:HB2	2.06	0.55
4:M:259:ASN:OD1	4:M:260:GLU:N	2.40	0.55
4:M:336:MET:HB2	4:M:366:ARG:CZ	2.37	0.55
4:M:347:ILE:HG23	4:M:359:PHE:CE1	2.41	0.55
4:M:592:LEU:HD12	4:M:595:ILE:HB	1.89	0.55
5:O:18:ILE:O	5:O:276:VAL:HG22	2.06	0.55
5:O:574:VAL:HB	5:O:612:PHE:HB3	1.89	0.55
3:P:392:PHE:CD1	3:P:396:PRO:HG3	2.42	0.55
6:X:8:GLY:O	6:X:11:ILE:N	2.38	0.55
6:X:132:ASN:OD1	6:X:133:TRP:N	2.40	0.55
6:Z:54:CYS:HB3	6:Z:73:CYS:HA	1.89	0.55
1:B:267:LYS:HD2	1:B:310:TRP:CE2	2.42	0.55
1:B:378:PHE:HA	1:B:391:LEU:O	2.06	0.55
1:B:489:GLY:O	2:C:672:SER:OG	2.22	0.55
1:B:670:THR:OG1	1:B:671:GLN:N	2.39	0.55
1:B:708:ILE:HA	1:B:711:ARG:HH21	1.72	0.55
3:D:229:HIS:CD2	3:D:232:ASN:HB2	2.42	0.55
3:D:336:VAL:HA	3:D:339:PHE:HB3	1.89	0.55
4:K:105:GLU:HA	4:K:108:ILE:HD12	1.89	0.55
4:K:174:TYR:O	4:K:178:ILE:HG12	2.07	0.55

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
4:L:621:LYS:HE3	6:Z:2:GLU:HG2	1.89	0.55
5:O:340:ASN:HD21	5:O:342:THR:HG1	1.53	0.55
5:O:687:ILE:HG12	5:O:992:ARG:HD2	1.88	0.55
5:O:926:VAL:HG21	5:O:1019:MET:HG2	1.89	0.55
5:O:971:ARG:HH22	5:O:1028:MET:H	1.54	0.55
5:O:1029:GLU:HB2	5:O:1043:VAL:O	2.07	0.55
5:O:1034:PHE:O	5:O:1120:ASP:N	2.37	0.55
3:P:233:THR:HB	3:P:236:THR:HA	1.89	0.55
6:X:74:ASN:HB2	6:X:76:GLN:HG3	1.88	0.55
6:X:166:TYR:O	6:X:170:LEU:HG	2.07	0.55
1:B:295:PRO:HA	1:B:405:ILE:HG23	1.88	0.54
1:B:468:ARG:HH12	1:B:1017:GLY:HA3	1.72	0.54
1:B:531:VAL:O	1:B:534:PRO:HD2	2.07	0.54
1:B:602:GLY:N	1:B:832:GLN:HA	2.22	0.54
1:B:646:VAL:HG22	1:B:688:PHE:HZ	1.73	0.54
1:B:882:ILE:HG13	1:B:883:THR:N	2.22	0.54
2:C:333:HIS:CE1	2:C:334:LEU:HG	2.42	0.54
2:C:550:GLN:O	2:C:890:LYS:N	2.39	0.54
2:C:807:THR:O	2:C:811:LEU:HG	2.06	0.54
4:K:18:ASN:ND2	4:K:249:TRP:HA	2.22	0.54
4:M:568:ILE:HD12	4:M:568:ILE:H	1.72	0.54
4:M:616:PRO:HA	6:Y:1:MET:HG3	1.89	0.54
5:O:166:VAL:HG12	5:O:167:ALA:N	2.22	0.54
5:O:352:GLN:NE2	5:O:370:VAL:O	2.40	0.54
5:O:1197:TYR:HD2	5:O:1211:GLN:HG3	1.69	0.54
3:P:71:LEU:O	3:P:75:THR:HG23	2.07	0.54
3:P:303:LEU:HB3	3:P:321:GLY:H	1.72	0.54
6:X:145:ILE:HG13	6:X:146:HIS:NE2	2.21	0.54
6:Z:118:LEU:HD13	6:Z:128:LEU:HD22	1.87	0.54
6:Z:131:LEU:O	6:Z:360:ASP:HB3	2.07	0.54
1:B:637:PRO:HG2	1:B:764:TRP:NE1	2.21	0.54
1:B:752:THR:HA	1:B:896:VAL:HG23	1.88	0.54
1:B:1001:TYR:O	1:B:1008:THR:HA	2.07	0.54
2:C:602:GLY:HA3	2:C:831:PHE:CE2	2.42	0.54
2:C:708:ILE:HD13	2:C:711:ARG:HH11	1.72	0.54
2:C:839:ASP:HB3	2:C:842:ARG:NE	2.21	0.54
2:C:843:VAL:O	2:C:1004:TYR:N	2.39	0.54
2:C:1064:ASP:N	2:C:1064:ASP:OD1	2.36	0.54
3:D:96:TRP:CE2	3:D:104:LEU:HD21	2.41	0.54
4:K:21:LYS:H	4:K:248:GLN:HE22	1.54	0.54
4:K:620:ASP:HB3	4:K:623:ASN:CG	2.28	0.54

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
5:O:192:LEU:HD12	5:O:195:TYR:HB3	1.89	0.54
5:O:394:VAL:HG12	5:O:395:GLN:H	1.70	0.54
5:O:778:ARG:O	5:O:781:GLU:HB3	2.06	0.54
5:O:892:LEU:H	5:O:925:GLN:NE2	2.03	0.54
5:O:941:GLU:HB2	5:O:950:ARG:HD2	1.89	0.54
5:O:1196:TYR:HB3	5:O:1214:LEU:HD12	1.89	0.54
6:X:58:VAL:O	6:X:63:ARG:NH1	2.39	0.54
6:X:265:LEU:N	6:X:268:MET:O	2.39	0.54
1:B:347:ARG:HH21	1:B:1177:SER:N	2.04	0.54
1:B:593:VAL:HG21	1:B:877:LEU:HD22	1.89	0.54
1:B:706:ALA:HA	1:B:709:ILE:HD12	1.89	0.54
2:C:483:LEU:O	2:C:487:ALA:N	2.24	0.54
2:C:1130:ILE:HG23	2:C:1160:TRP:HH2	1.71	0.54
3:D:165:ARG:HB2	3:D:267:ASN:HD21	1.72	0.54
3:D:246:ASP:OD1	3:D:247:ALA:N	2.40	0.54
3:D:362:ASP:HA	3:D:365:GLN:HE21	1.71	0.54
3:D:369:ARG:NH2	3:D:373:GLU:OE1	2.40	0.54
4:K:161:LYS:HD2	4:K:162:GLN:HB2	1.90	0.54
4:K:174:TYR:O	4:K:177:SER:OG	2.18	0.54
4:K:408:ILE:HD13	4:K:418:GLN:HG2	1.90	0.54
4:K:525:GLU:OE1	4:M:447:SER:OG	2.24	0.54
4:L:402:ILE:HG13	4:L:424:VAL:HG23	1.90	0.54
5:O:347:GLN:HB2	5:O:350:TYR:H	1.72	0.54
5:O:494:TYR:CE2	5:O:541:VAL:HG22	2.42	0.54
5:O:646:THR:HG22	5:O:647:ASN:H	1.72	0.54
5:O:712:GLU:HG2	5:O:750:ARG:O	2.08	0.54
5:O:988:LEU:H	5:O:989:ARG:CZ	2.20	0.54
3:P:230:GLN:HG3	3:P:258:ASN:ND2	2.23	0.54
6:X:261:THR:OG1	6:X:262:THR:O	2.23	0.54
1:B:717:SER:OG	1:B:744:ASP:HA	2.08	0.54
2:C:418:LYS:HD2	2:C:1214:SER:HB2	1.90	0.54
2:C:514:ARG:O	2:C:517:SER:OG	2.20	0.54
2:C:682:HIS:NE2	3:P:177:GLN:HB2	2.22	0.54
2:C:708:ILE:HG23	2:C:712:TYR:CD1	2.37	0.54
2:C:898:ASN:HA	2:C:901:TYR:CD1	2.43	0.54
2:C:983:LEU:O	2:C:984:LEU:HG	2.07	0.54
2:C:1005:ASN:OD1	2:C:1006:GLY:N	2.40	0.54
2:C:1151:TYR:CD2	2:C:1181:MET:HG2	2.42	0.54
3:D:316:ILE:O	3:D:320:THR:N	2.32	0.54
4:K:225:ASP:OD1	4:K:226:ASP:N	2.40	0.54
4:L:102:VAL:HG21	4:L:148:VAL:HG11	1.89	0.54

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
4:L:559:ASP:H	4:L:601:GLN:HE22	1.54	0.54
5:O:207:LEU:HA	5:O:210:PHE:HB3	1.90	0.54
5:O:743:ILE:N	5:O:787:ILE:HB	2.21	0.54
5:O:1192:LYS:HD3	5:O:1220:ILE:HG12	1.88	0.54
5:O:1222:LEU:HD12	5:O:1223:PRO:CD	2.36	0.54
6:X:221:MET:HG2	6:X:358:ILE:N	2.22	0.54
6:X:302:VAL:O	6:X:306:TRP:N	2.41	0.54
6:Z:17:ASN:O	6:Z:20:GLU:HG2	2.07	0.54
1:B:518:GLN:HE22	1:B:828:PHE:HD1	1.55	0.54
1:B:1165:ALA:O	1:B:1168:GLU:HG2	2.07	0.54
2:C:675:ALA:HA	2:C:678:PHE:CD2	2.42	0.54
2:C:854:ARG:HH22	2:C:1010:ASN:ND2	2.06	0.54
3:D:291:LEU:O	3:D:295:LEU:HG	2.08	0.54
3:D:344:ARG:HE	3:D:354:GLN:HE22	1.54	0.54
4:K:349:VAL:HB	4:K:357:TRP:HD1	1.72	0.54
4:L:124:PHE:O	4:L:128:LEU:HG	2.06	0.54
4:M:465:ASN:HB3	4:M:467:TYR:CE1	2.43	0.54
4:M:616:PRO:HD2	4:M:618:LEU:HD13	1.89	0.54
5:O:142:MET:O	5:O:146:LEU:N	2.26	0.54
5:O:157:PHE:HA	5:O:166:VAL:HG22	1.90	0.54
5:O:470:SER:O	5:O:474:LEU:HG	2.08	0.54
5:O:483:LEU:HD21	5:O:528:LEU:HB3	1.89	0.54
5:O:939:TYR:C	5:O:952:PRO:HD2	2.27	0.54
5:O:948:ARG:HE	5:O:957:ASP:CG	2.11	0.54
5:O:1169:ASP:OD2	5:O:1169:ASP:N	2.41	0.54
6:Y:65:LEU:HG	6:Y:67:HIS:H	1.70	0.54
6:Y:208:ARG:CZ	6:Y:287:LYS:HG3	2.38	0.54
6:Y:247:ARG:HB3	6:Y:348:LYS:HD2	1.88	0.54
6:Z:79:HIS:HA	6:Z:82:TYR:CE2	2.43	0.54
1:B:897:THR:OG1	1:B:898:ASN:N	2.40	0.54
1:B:1218:GLN:HE22	2:C:1087:MET:HG3	1.73	0.54
2:C:741:LEU:C	2:C:835:TYR:HE2	2.11	0.54
4:K:153:ARG:HH22	4:M:132:SER:HA	1.73	0.54
4:K:380:LEU:HD22	4:K:448:LEU:HB3	1.89	0.54
4:K:499:SER:HB2	4:K:502:GLU:CD	2.28	0.54
4:L:145:ASP:HB3	4:L:161:LYS:HG3	1.89	0.54
4:L:636:ARG:NE	4:M:28:SER:O	2.41	0.54
4:M:170:ARG:HA	4:M:173:MET:HE2	1.88	0.54
4:M:193:ARG:HA	4:M:196:GLN:NE2	2.23	0.54
5:O:26:THR:HB	5:O:108:ASN:HB3	1.89	0.54
5:O:527:PRO:O	5:O:530:ILE:HG12	2.08	0.54

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
5:O:600:SER:O	5:O:603:MET:HB3	2.07	0.54
5:O:671:TYR:O	5:O:675:VAL:HG23	2.07	0.54
3:P:144:TYR:CD1	3:P:147:LEU:HD12	2.43	0.54
6:Z:17:ASN:HA	6:Z:20:GLU:HG2	1.89	0.54
6:Z:88:ALA:O	6:Z:92:THR:HG23	2.08	0.54
1:B:516:ILE:HA	1:B:519:ILE:HD12	1.89	0.54
1:B:1110:PHE:CZ	1:B:1114:LEU:HB2	2.42	0.54
1:B:1115:TRP:CE2	1:B:1122:PHE:HB3	2.43	0.54
2:C:336:LYS:HD3	2:C:348:GLY:O	2.08	0.54
2:C:1148:LEU:O	2:C:1149:HIS:ND1	2.40	0.54
3:D:9:LYS:NZ	3:D:154:ASN:HB3	2.22	0.54
4:K:369:ASN:HA	4:K:467:TYR:HD2	1.72	0.54
4:L:322:ASN:HD21	4:L:324:ARG:HB2	1.72	0.54
4:M:565:SER:HA	4:M:568:ILE:HD13	1.90	0.54
5:O:948:ARG:HB3	5:O:957:ASP:OD1	2.07	0.54
5:O:1261:TRP:CD1	5:O:1279:VAL:HG13	2.43	0.54
3:P:161:ILE:HG23	3:P:272:VAL:N	2.21	0.54
6:Z:324:TYR:O	6:Z:328:MET:HG3	2.08	0.54
1:B:424:ALA:O	2:C:1079:ARG:NH2	2.41	0.54
1:B:1160:TRP:O	1:B:1163:THR:HB	2.07	0.54
2:C:690:ASN:HB3	2:C:693:LEU:HD11	1.90	0.54
2:C:1246:GLN:HB2	2:C:1250:VAL:O	2.08	0.54
3:D:166:VAL:O	3:D:267:ASN:HA	2.08	0.54
4:K:301:ILE:HG13	4:K:624:TRP:HE1	1.73	0.54
4:K:314:ILE:HG23	4:K:542:ARG:HH22	1.72	0.54
4:L:39:GLY:HA2	4:L:42:ASN:HD22	1.73	0.54
4:L:426:ASN:OD1	4:L:427:TYR:N	2.41	0.54
4:L:546:ILE:O	4:L:550:ILE:HG13	2.08	0.54
4:M:109:ALA:HB1	4:M:113:LYS:NZ	2.23	0.54
4:M:337:ILE:HB	4:M:365:THR:H	1.72	0.54
4:M:655:GLU:HA	4:M:658:GLN:HB2	1.90	0.54
5:O:27:LEU:H	5:O:109:GLU:H	1.55	0.54
5:O:210:PHE:HD2	5:O:211:PHE:CD1	2.26	0.54
5:O:307:LEU:HD22	5:O:342:THR:HG21	1.88	0.54
5:O:642:LYS:HD2	5:O:650:GLU:HG2	1.90	0.54
5:O:1236:ARG:NH1	5:O:1236:ARG:HA	2.23	0.54
3:P:306:ASN:HD21	3:P:322:ARG:HD2	1.73	0.54
6:X:180:VAL:H	6:X:365:GLY:HA2	1.73	0.54
6:Y:122:ARG:HH21	6:Y:236:ARG:HH21	1.53	0.54
6:Y:206:PHE:CD2	6:Y:267:LYS:HD2	2.43	0.54
6:Z:258:SER:HB2	6:Z:343:PHE:CE2	2.43	0.54

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:B:573:PRO:O	1:B:576:SER:OG	2.26	0.54
1:B:941:THR:CG2	1:B:943:TYR:HB2	2.38	0.54
1:B:983:LEU:HB3	1:B:984:LEU:HD12	1.89	0.54
2:C:398:ASP:O	2:C:401:LYS:N	2.40	0.54
4:K:281:GLU:HA	4:K:284:LYS:HG2	1.90	0.54
4:K:297:GLU:OE2	4:K:299:GLU:HB2	2.08	0.54
4:K:522:TYR:CD2	4:K:527:LEU:HG	2.43	0.54
4:K:610:ILE:HD12	4:K:624:TRP:HZ3	1.73	0.54
4:K:620:ASP:OD1	4:K:622:ASN:N	2.29	0.54
4:K:658:GLN:NE2	4:K:659:ASN:OD1	2.40	0.54
4:L:478:ILE:HG23	4:L:480:PRO:HD3	1.88	0.54
4:L:595:ILE:HG23	4:L:598:ARG:NH2	2.23	0.54
4:M:37:SER:O	4:M:37:SER:OG	2.23	0.54
5:O:59:LEU:HD23	5:O:64:GLN:HG2	1.90	0.54
5:O:690:GLN:HE21	5:O:992:ARG:NE	2.05	0.54
5:O:1248:CYS:SG	5:O:1261:TRP:HH2	2.30	0.54
3:P:239:PHE:HB2	3:P:251:TRP:HE3	1.71	0.54
6:Y:51:CYS:O	6:Y:55:LEU:HD13	2.08	0.54
6:Y:107:HIS:HE1	6:Y:169:ALA:HB3	1.73	0.54
6:Z:132:ASN:OD1	6:Z:133:TRP:N	2.41	0.54
6:Z:251:HIS:CB	6:Z:254:LEU:HB2	2.37	0.54
1:B:628:ASP:HA	1:B:631:ILE:HD12	1.89	0.54
2:C:719:ILE:HD12	2:C:745:HIS:CE1	2.43	0.54
2:C:919:GLN:O	2:C:923:ILE:HG13	2.08	0.54
2:C:1056:TRP:CE2	2:C:1062:PRO:HD3	2.43	0.54
4:K:123:GLU:OE2	4:K:238:VAL:HB	2.08	0.54
4:K:172:THR:O	4:K:175:VAL:HG22	2.08	0.54
4:K:308:PRO:HG2	4:K:311:VAL:HB	1.90	0.54
4:L:353:THR:HA	6:Y:32:TRP:CZ3	2.43	0.54
4:M:555:ASP:OD2	4:M:555:ASP:N	2.41	0.54
5:O:88:GLU:O	5:O:91:ARG:HG2	2.08	0.54
5:O:575:TYR:HA	5:O:613:VAL:CG1	2.38	0.54
5:O:716:PHE:CD2	5:O:781:GLU:HA	2.43	0.54
5:O:993:LEU:HA	5:O:996:LEU:HB2	1.90	0.54
6:Y:186:ARG:HB2	6:Y:351:ASP:OD2	2.07	0.54
6:Y:227:GLU:HB2	6:Y:354:TYR:CZ	2.43	0.54
6:Z:90:ARG:HH22	6:Z:151:ASP:C	2.12	0.54
1:B:636:LEU:CD1	1:B:648:ALA:HB2	2.38	0.53
1:B:670:THR:HG23	1:B:672:SER:H	1.73	0.53
2:C:325:ALA:HB1	2:C:329:GLU:OE1	2.08	0.53
2:C:474:ILE:HD11	2:C:507:TYR:HB2	1.90	0.53

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:C:904:ALA:O	2:C:907:PRO:HD2	2.07	0.53
4:L:405:GLN:HG2	4:L:457:LEU:HD22	1.89	0.53
4:M:402:ILE:HA	4:M:424:VAL:HA	1.90	0.53
5:O:458:ASP:N	5:O:458:ASP:OD1	2.40	0.53
5:O:490:LEU:HA	5:O:493:ALA:HB3	1.91	0.53
5:O:927:ASN:HB3	5:O:949:TYR:CE1	2.43	0.53
5:O:1046:GLY:H	5:O:1086:GLU:HG2	1.72	0.53
6:Z:51:CYS:HB3	6:Z:54:CYS:SG	2.49	0.53
6:Z:95:TRP:CZ3	6:Z:284:PRO:HG3	2.43	0.53
6:Z:222:VAL:O	6:Z:357:MET:N	2.26	0.53
2:C:298:GLU:HG2	2:C:1214:SER:HB3	1.88	0.53
2:C:417:SER:OG	2:C:1214:SER:O	2.26	0.53
3:D:96:TRP:HA	3:D:104:LEU:HG	1.89	0.53
3:D:171:TRP:NE1	3:D:234:TYR:OH	2.34	0.53
4:K:305:VAL:O	4:K:307:VAL:HG13	2.08	0.53
4:L:31:VAL:HG23	4:L:120:PHE:HZ	1.73	0.53
4:L:240:LEU:HB3	4:L:247:ILE:HD13	1.90	0.53
4:L:407:LYS:HB3	4:L:463:GLU:HB2	1.90	0.53
4:L:566:VAL:HG11	4:M:197:THR:HG23	1.89	0.53
4:M:355:THR:HG1	4:M:357:TRP:HE1	1.54	0.53
4:M:645:LYS:HZ1	4:M:649:SER:HB3	1.71	0.53
3:P:70:CYS:O	3:P:73:SER:OG	2.20	0.53
3:P:342:ARG:O	3:P:345:ARG:HB2	2.07	0.53
6:Y:131:LEU:O	6:Y:361:PRO:HD2	2.09	0.53
1:B:264:THR:HG21	1:B:1253:LEU:HD11	1.91	0.53
1:B:518:GLN:NE2	1:B:829:PRO:HD2	2.22	0.53
1:B:518:GLN:O	1:B:522:ILE:HG12	2.07	0.53
1:B:616:ASP:OD1	1:B:616:ASP:N	2.42	0.53
1:B:636:LEU:HD12	1:B:648:ALA:HB2	1.91	0.53
3:D:85:ARG:NH1	3:D:87:GLY:O	2.41	0.53
3:D:286:ILE:HD12	3:D:289:SER:OG	2.09	0.53
4:K:338:PRO:O	4:K:341:MET:HG2	2.08	0.53
4:K:393:ASP:H	6:Z:277:THR:HG22	1.73	0.53
4:K:607:PRO:O	4:K:610:ILE:HG12	2.08	0.53
4:M:358:HIS:N	4:M:474:ASP:O	2.29	0.53
5:O:533:TRP:CE2	5:O:544:PRO:HD3	2.44	0.53
5:O:760:PRO:HB2	5:O:764:ARG:NH1	2.22	0.53
5:O:924:VAL:O	5:O:926:VAL:HG13	2.07	0.53
3:P:242:ASN:ND2	3:P:244:ARG:HH22	2.04	0.53
6:X:336:THR:HG23	6:X:340:LEU:HD13	1.90	0.53
6:Y:307:GLY:O	6:Y:311:ILE:HG13	2.09	0.53

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
6:Z:10:GLN:HA	6:Z:13:ASP:OD1	2.08	0.53
1:B:268:ILE:HG13	1:B:303:ARG:C	2.28	0.53
2:C:352:LEU:O	2:C:954:SER:OG	2.15	0.53
2:C:374:ARG:O	2:C:1259:ARG:NH1	2.42	0.53
2:C:628:ASP:O	2:C:632:LEU:HG	2.07	0.53
4:K:229:ILE:HG13	4:K:233:PRO:HA	1.89	0.53
4:K:398:LYS:HA	4:K:429:GLN:HA	1.90	0.53
4:L:568:ILE:H	4:L:568:ILE:HD12	1.74	0.53
4:M:336:MET:SD	4:M:339:LYS:NZ	2.82	0.53
4:M:381:ASP:O	4:M:491:ALA:HB1	2.08	0.53
5:O:70:THR:HA	5:O:73:TYR:CD1	2.44	0.53
3:P:144:TYR:HD1	3:P:147:LEU:HD12	1.73	0.53
6:X:17:ASN:HA	6:X:20:GLU:OE1	2.08	0.53
6:X:186:ARG:HH12	6:X:245:PRO:HB3	1.72	0.53
1:B:474:ILE:H	1:B:505:MET:HG3	1.73	0.53
1:B:1171:THR:OG1	1:B:1174:SER:N	2.40	0.53
2:C:982:MET:SD	2:C:984:LEU:N	2.81	0.53
2:C:1001:TYR:CE1	2:C:1009:PHE:HB2	2.44	0.53
3:D:370:ASP:HB2	3:D:371:LYS:HZ2	1.72	0.53
4:K:385:LYS:N	4:K:443:ILE:HG22	2.21	0.53
4:L:133:VAL:HG22	4:L:134:SER:H	1.73	0.53
4:L:451:ASN:H	6:Y:62:GLN:NE2	2.06	0.53
4:L:631:VAL:HG13	4:L:632:LYS:N	2.23	0.53
4:M:214:SER:HA	4:M:217:LYS:HE2	1.90	0.53
4:M:457:LEU:HD13	4:M:466:TYR:CD2	2.43	0.53
5:O:48:ASN:HD21	5:O:51:THR:H	1.56	0.53
5:O:61:ARG:HH22	5:O:168:ALA:HB1	1.72	0.53
5:O:206:VAL:O	5:O:209:ARG:HB2	2.08	0.53
5:O:609:GLY:HA2	5:O:660:GLN:OE1	2.08	0.53
3:P:48:LEU:O	3:P:48:LEU:HD23	2.09	0.53
1:B:246:GLN:HE21	1:B:981:ASP:HB2	1.72	0.53
1:B:484:THR:O	1:B:487:ALA:N	2.42	0.53
1:B:511:ASN:HD21	1:B:729:SER:N	2.06	0.53
1:B:1107:ASN:C	1:B:1108:TRP:CD1	2.81	0.53
2:C:653:ALA:O	2:C:657:VAL:HG23	2.08	0.53
4:K:41:LEU:HD23	4:M:263:ASN:HA	1.90	0.53
4:K:554:ASP:OD2	4:K:554:ASP:N	2.41	0.53
4:L:600:MET:O	4:L:604:ILE:HG13	2.09	0.53
4:M:206:GLY:HA3	4:M:658:GLN:NE2	2.24	0.53
5:O:278:ARG:HB2	5:O:281:GLN:NE2	2.23	0.53
5:O:349:ARG:NH1	3:P:129:ASP:HA	2.24	0.53

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
6:X:121:VAL:HG13	6:X:125:GLY:HA2	1.90	0.53
6:Y:140:SER:OG	6:Y:144:SER:N	2.42	0.53
6:Z:117:ASP:O	6:Z:121:VAL:HG23	2.09	0.53
1:B:518:GLN:HE21	1:B:829:PRO:HD2	1.74	0.53
1:B:713:TRP:HA	1:B:764:TRP:CH2	2.44	0.53
1:B:719:ILE:O	1:B:739:LEU:HB3	2.09	0.53
1:B:1102:VAL:HG12	1:B:1103:PRO:O	2.08	0.53
2:C:279:ASP:OD2	2:C:280:VAL:HG23	2.09	0.53
2:C:336:LYS:NZ	2:C:346:VAL:O	2.30	0.53
2:C:337:GLN:N	2:C:337:GLN:OE1	2.42	0.53
2:C:619:SER:OG	2:C:620:VAL:N	2.42	0.53
4:M:335:ARG:HG2	4:M:369:ASN:HD21	1.73	0.53
4:M:367:VAL:CG2	4:M:467:TYR:HB3	2.39	0.53
4:M:405:GLN:CG	4:M:421:GLN:HB2	2.39	0.53
5:O:7:VAL:HG21	5:O:330:PHE:HE1	1.73	0.53
5:O:879:THR:HG23	5:O:916:LYS:HD3	1.91	0.53
5:O:1194:LEU:HD21	5:O:1233:PRO:HB2	1.90	0.53
3:P:189:GLU:O	3:P:193:THR:HG23	2.08	0.53
3:P:370:ASP:HB2	3:P:371:LYS:NZ	2.24	0.53
6:X:315:LEU:HD12	6:X:323:TRP:NE1	2.23	0.53
6:X:331:ALA:HA	6:X:334:VAL:HG22	1.90	0.53
2:C:316:PHE:HD2	2:C:371:TYR:HE2	1.56	0.53
4:L:526:SER:O	4:L:530:SER:N	2.41	0.53
4:M:410:PHE:HA	4:M:413:TRP:CE2	2.44	0.53
5:O:93:LEU:HD12	5:O:96:GLU:HB2	1.91	0.53
5:O:581:VAL:HG23	5:O:582:VAL:H	1.74	0.53
5:O:768:ARG:HA	5:O:862:ASN:HD21	1.73	0.53
5:O:813:PHE:O	5:O:817:SER:HB3	2.09	0.53
1:B:258:TRP:HB3	1:B:374:ARG:HB2	1.89	0.53
1:B:298:GLU:HG3	1:B:1214:SER:N	2.24	0.53
1:B:842:ARG:HA	1:B:1003:GLN:HE21	1.74	0.53
2:C:721:TYR:N	2:C:737:GLU:O	2.42	0.53
4:K:228:LEU:HA	4:K:231:ARG:HG3	1.91	0.53
4:K:373:ILE:HG13	4:K:452:TYR:CE2	2.43	0.53
4:K:629:GLN:NE2	4:L:202:ASN:O	2.36	0.53
4:L:37:SER:O	4:L:41:LEU:HG	2.09	0.53
4:L:393:ASP:H	4:L:397:LYS:NZ	2.07	0.53
4:M:399:VAL:HG22	4:M:430:LEU:HD21	1.90	0.53
4:M:523:THR:O	4:M:526:SER:N	2.42	0.53
5:O:1145:ILE:N	5:O:1181:LEU:O	2.41	0.53
5:O:1198:ILE:HG13	5:O:1214:LEU:HD11	1.91	0.53

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
5:O:1240:VAL:HB	5:O:1248:CYS:H	1.73	0.53
6:X:120:ARG:HG2	6:X:124:GLU:OE1	2.09	0.53
6:Y:50:VAL:HB	6:Y:52:MET:HE1	1.91	0.53
6:Y:168:THR:HA	6:Y:252:PHE:CZ	2.43	0.53
6:Z:168:THR:O	6:Z:172:LEU:HG	2.08	0.53
2:C:268:ILE:CG1	2:C:304:ILE:HG13	2.38	0.53
2:C:345:SER:OG	2:C:346:VAL:N	2.43	0.53
2:C:518:GLN:N	2:C:521:ARG:HH21	2.06	0.53
2:C:846:MET:HE3	2:C:847:VAL:H	1.74	0.53
2:C:1158:ASN:H	2:C:1181:MET:HE1	1.73	0.53
4:K:187:LYS:HD3	4:K:190:ILE:HD11	1.91	0.53
4:K:321:TYR:HH	4:K:537:ARG:NH2	2.06	0.53
4:K:378:PHE:HD1	4:K:495:MET:SD	2.31	0.53
4:K:430:LEU:HD13	4:K:472:PHE:CZ	2.44	0.53
4:L:304:LEU:HA	4:L:514:VAL:HG13	1.90	0.53
4:L:514:VAL:HG12	4:L:517:GLU:H	1.74	0.53
4:M:193:ARG:O	4:M:196:GLN:HG2	2.09	0.53
4:M:327:LYS:HE2	6:Y:334:VAL:HG11	1.90	0.53
4:M:399:VAL:HG12	4:M:472:PHE:HA	1.91	0.53
5:O:153:LEU:HD23	5:O:157:PHE:HE1	1.74	0.53
5:O:243:SER:OG	5:O:244:ALA:N	2.42	0.53
6:Y:102:PHE:HA	6:Y:105:GLN:NE2	2.24	0.53
6:Y:337:PRO:HD2	6:Y:340:LEU:HD11	1.91	0.53
1:B:293:GLN:HB3	1:B:405:ILE:HD11	1.92	0.52
1:B:377:GLY:HA3	1:B:393:SER:O	2.09	0.52
2:C:264:THR:O	2:C:309:ARG:HA	2.09	0.52
3:D:60:THR:HB	3:D:280:ASN:O	2.08	0.52
3:D:63:SER:HA	3:D:393:ARG:O	2.09	0.52
3:D:268:ARG:HB3	3:D:417:LEU:HD23	1.91	0.52
4:K:425:VAL:HG12	6:Z:61:LEU:O	2.09	0.52
4:M:169:THR:O	4:M:173:MET:HG3	2.09	0.52
4:M:346:GLN:HG3	4:M:360:ASN:HB3	1.91	0.52
5:O:415:SER:OG	5:O:416:ARG:NE	2.42	0.52
3:P:109:PRO:HA	3:P:112:LEU:HD12	1.91	0.52
3:P:139:PHE:HA	3:P:142:ARG:NE	2.22	0.52
6:X:74:ASN:ND2	6:X:77:ILE:HD11	2.23	0.52
6:X:248:ASP:OD2	6:X:347:THR:HA	2.09	0.52
6:Y:10:GLN:NE2	6:Y:46:GLY:O	2.42	0.52
6:Y:325:ASN:C	6:Y:329:GLN:HE21	2.13	0.52
1:B:1041:ALA:HB2	1:B:1146:TYR:CE1	2.44	0.52
1:B:1118:ASN:H	1:B:1122:PHE:HE2	1.56	0.52

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:C:371:TYR:H	2:C:462:ARG:NH2	2.07	0.52
2:C:517:SER:C	2:C:521:ARG:HE	2.12	0.52
2:C:650:MET:HG3	2:C:654:ASN:HD21	1.75	0.52
2:C:810:TYR:CD2	2:C:811:LEU:HD23	2.45	0.52
2:C:1197:GLN:NE2	2:C:1199:ILE:HB	2.24	0.52
3:D:81:ARG:NE	3:D:87:GLY:HA2	2.23	0.52
4:K:250:MET:HA	4:K:257:VAL:HB	1.92	0.52
4:K:395:ASN:HB2	4:K:432:ALA:HB3	1.92	0.52
4:K:573:ALA:O	4:K:577:LEU:HG	2.09	0.52
4:K:622:ASN:HD22	4:K:625:ILE:HD12	1.74	0.52
4:L:381:ASP:OD1	4:L:447:SER:HA	2.10	0.52
4:L:570:GLY:O	4:L:574:ILE:HG12	2.08	0.52
5:O:156:PHE:O	5:O:166:VAL:HG13	2.09	0.52
6:Z:244:THR:HB	6:Z:247:ARG:HG2	1.91	0.52
1:B:543:LEU:HD11	1:B:822:ILE:HD13	1.91	0.52
1:B:947:ARG:NH1	1:B:956:ARG:HH22	2.06	0.52
2:C:360:HIS:CG	2:C:363:LEU:HD23	2.44	0.52
2:C:949:LEU:HB3	2:C:952:ILE:HD12	1.91	0.52
2:C:1104:PHE:HE1	2:C:1126:PHE:HD1	1.58	0.52
4:K:123:GLU:HA	4:K:235:GLU:OE1	2.10	0.52
4:K:282:LYS:HG3	4:K:640:LEU:HD22	1.90	0.52
4:K:450:TYR:HD1	6:Z:62:GLN:HE22	1.57	0.52
5:O:426:TYR:HE2	5:O:429:ILE:HD11	1.73	0.52
5:O:511:SER:OG	5:O:572:GLN:N	2.26	0.52
3:P:137:PRO:HD2	3:P:138:ARG:NH1	2.23	0.52
3:P:350:THR:H	3:P:353:GLN:CD	2.13	0.52
6:X:172:LEU:HD23	6:X:173:MET:HG3	1.90	0.52
6:X:223:TYR:CD2	6:X:354:TYR:HB2	2.32	0.52
6:Y:26:TYR:HE2	6:Y:68:LEU:HB2	1.72	0.52
1:B:446:GLU:OE1	1:B:446:GLU:N	2.43	0.52
1:B:521:ARG:HH22	1:B:828:PHE:HA	1.75	0.52
1:B:811:LEU:HA	1:B:815:ALA:HB3	1.90	0.52
2:C:398:ASP:O	2:C:402:TRP:HD1	1.93	0.52
2:C:548:PRO:HD3	2:C:900:TRP:NE1	2.24	0.52
2:C:632:LEU:O	2:C:636:LEU:HG	2.10	0.52
2:C:851:ARG:HG3	2:C:997:LEU:HD11	1.90	0.52
2:C:1043:GLY:O	2:C:1144:TYR:HE1	1.91	0.52
4:K:19:VAL:HG12	4:K:248:GLN:HE21	1.75	0.52
4:K:34:LEU:HD22	4:M:279:LEU:HD21	1.92	0.52
4:K:345:PHE:N	4:K:360:ASN:OD1	2.42	0.52
4:K:639:SER:OG	4:L:31:VAL:N	2.42	0.52

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
4:L:377:ARG:HB3	4:L:496:SER:OG	2.10	0.52
4:L:410:PHE:HA	4:L:413:TRP:NE1	2.24	0.52
5:O:69:ASP:OD2	5:O:70:THR:N	2.42	0.52
5:O:80:ASP:O	5:O:83:GLU:HG3	2.09	0.52
5:O:181:ASP:N	5:O:184:ASP:OD2	2.35	0.52
5:O:624:TRP:HA	5:O:627:ILE:HD12	1.92	0.52
5:O:1075:GLN:O	5:O:1091:MET:HB2	2.08	0.52
5:O:1192:LYS:HE3	5:O:1220:ILE:H	1.75	0.52
3:P:108:PRO:O	3:P:112:LEU:N	2.38	0.52
6:X:317:PRO:HG2	6:X:326:ARG:CZ	2.39	0.52
6:Y:51:CYS:SG	6:Y:53:HIS:N	2.74	0.52
2:C:604:VAL:HA	2:C:873:VAL:H	1.74	0.52
2:C:740:LEU:HD13	2:C:835:TYR:CD1	2.45	0.52
4:K:334:LEU:HB3	4:K:367:VAL:O	2.09	0.52
4:K:384:GLY:N	4:K:443:ILE:O	2.35	0.52
4:K:488:VAL:O	4:K:490:ASP:N	2.38	0.52
4:M:525:GLU:HA	4:M:528:ASN:ND2	2.25	0.52
5:O:824:VAL:HA	5:O:845:PRO:O	2.10	0.52
3:P:106:VAL:HG12	3:P:112:LEU:HD21	1.90	0.52
3:P:408:GLY:O	3:P:411:SER:OG	2.12	0.52
6:Y:22:ARG:HH12	6:Y:96:LYS:HG3	1.74	0.52
6:Y:27:SER:HB2	6:Y:30:GLU:HB2	1.91	0.52
1:B:257:THR:HG1	1:B:372:LEU:H	1.52	0.52
1:B:351:PRO:HA	1:B:354:PHE:CZ	2.44	0.52
1:B:527:ASN:ND2	1:B:871:VAL:HG13	2.24	0.52
1:B:647:LYS:O	1:B:651:THR:HG23	2.09	0.52
1:B:930:THR:O	1:B:934:LEU:HG	2.10	0.52
1:B:1210:PHE:HZ	1:B:1227:ILE:HG23	1.74	0.52
2:C:308:THR:OG1	2:C:322:VAL:N	2.33	0.52
2:C:351:PRO:HA	2:C:354:PHE:CZ	2.45	0.52
2:C:1124:GLN:HB2	2:C:1125:GLN:NE2	2.25	0.52
4:K:300:ILE:HG13	4:K:312:PHE:CE1	2.44	0.52
4:K:349:VAL:HB	4:K:357:TRP:CD1	2.44	0.52
4:L:65:ARG:HE	4:L:99:PRO:HB3	1.73	0.52
4:L:356:ASN:O	4:L:475:SER:OG	2.18	0.52
4:L:376:MET:HB3	4:L:378:PHE:CZ	2.44	0.52
5:O:188:PHE:CD2	5:O:258:GLU:HG3	2.45	0.52
5:O:256:LEU:O	5:O:259:SER:OG	2.18	0.52
5:O:587:ASP:O	5:O:591:SER:OG	2.18	0.52
3:P:16:LEU:HG	3:P:18:ASN:H	1.75	0.52
3:P:152:LEU:HD21	3:P:319:TRP:CD2	2.44	0.52

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
6:X:208:ARG:HG2	6:X:287:LYS:HE2	1.90	0.52
6:X:315:LEU:HD12	6:X:323:TRP:HE1	1.74	0.52
6:Y:5:LEU:O	6:Y:314:ALA:HB1	2.10	0.52
6:Z:225:TYR:O	6:Z:228:LEU:HB2	2.09	0.52
1:B:325:ALA:HB2	1:B:331:ASN:HD21	1.75	0.52
1:B:649:PHE:HZ	1:B:702:LEU:HB2	1.72	0.52
1:B:673:ARG:HB3	1:B:678:PHE:CZ	2.44	0.52
1:B:1158:ASN:ND2	1:B:1160:TRP:HB2	2.25	0.52
2:C:607:VAL:O	2:C:608:ILE:HD13	2.09	0.52
2:C:942:GLN:HB2	2:C:995:THR:OG1	2.10	0.52
3:D:276:SER:O	3:D:279:TRP:NE1	2.42	0.52
4:K:498:LEU:HD23	4:L:606:ASP:OD2	2.09	0.52
4:M:67:THR:OG1	4:M:98:GLU:O	2.28	0.52
5:O:23:ARG:NH2	5:O:29:ASP:OD1	2.42	0.52
5:O:1044:ILE:HB	5:O:1087:TRP:HB2	1.91	0.52
3:P:64:ARG:O	3:P:68:MET:HG3	2.10	0.52
3:P:127:CYS:SG	3:P:144:TYR:HE1	2.33	0.52
3:P:230:GLN:HG3	3:P:258:ASN:HD22	1.74	0.52
3:P:292:THR:HA	3:P:295:LEU:HG	1.92	0.52
6:X:121:VAL:HA	6:X:125:GLY:H	1.74	0.52
1:B:1062:PRO:HD2	1:B:1066:VAL:HG22	1.92	0.52
3:D:63:SER:C	3:D:67:GLN:HE21	2.13	0.52
3:D:208:THR:HA	3:D:211:ARG:HE	1.75	0.52
3:D:263:ILE:HG13	3:D:264:ARG:H	1.74	0.52
3:D:274:PRO:HG2	3:D:279:TRP:CD1	2.45	0.52
4:K:214:SER:HA	4:K:217:LYS:HD2	1.91	0.52
4:K:583:VAL:C	6:X:70:HIS:HB2	2.28	0.52
4:L:400:GLY:HA2	4:L:427:TYR:HA	1.92	0.52
4:M:375:PRO:HG3	4:M:454:PRO:CD	2.40	0.52
4:M:390:THR:OG1	4:M:391:SER:N	2.43	0.52
5:O:63:LEU:O	5:O:120:VAL:HG11	2.09	0.52
5:O:70:THR:HA	5:O:73:TYR:CG	2.45	0.52
5:O:219:GLY:O	5:O:235:LEU:HA	2.09	0.52
5:O:472:PHE:HD1	5:O:475:ALA:HB3	1.75	0.52
5:O:670:VAL:O	5:O:674:LEU:HG	2.10	0.52
3:P:81:ARG:H	3:P:84:HIS:CD2	2.27	0.52
6:X:85:VAL:HB	6:X:86:GLN:NE2	2.24	0.52
6:X:255:SER:H	6:X:276:LEU:HG	1.74	0.52
6:Z:121:VAL:HG13	6:Z:125:GLY:HA2	1.91	0.52
1:B:550:GLN:OE1	1:B:890:LYS:HB3	2.10	0.52
1:B:649:PHE:HA	1:B:652:LEU:HD12	1.92	0.52

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:B:1068:ASP:CG	1:B:1069:ARG:H	2.14	0.52
2:C:601:ASN:OD1	2:C:833:VAL:N	2.33	0.52
2:C:928:VAL:HG12	2:C:984:LEU:HD23	1.92	0.52
4:K:65:ARG:NH2	4:M:258:MET:HG3	2.25	0.52
4:M:237:ALA:HA	4:M:249:TRP:HE1	1.74	0.52
5:O:28:HIS:HA	5:O:31:CYS:SG	2.49	0.52
5:O:142:MET:O	5:O:146:LEU:HG	2.09	0.52
5:O:1131:PRO:HD2	5:O:1143:VAL:HG13	1.92	0.52
3:P:156:THR:HG22	3:P:158:PHE:H	1.75	0.52
6:Y:65:LEU:HD12	6:Y:66:LYS:H	1.75	0.52
6:Z:44:VAL:HA	6:Z:49:VAL:HG12	1.92	0.52
1:B:465:TRP:CE2	1:B:469:LEU:HD11	2.44	0.52
1:B:753:PRO:HD3	1:B:896:VAL:HA	1.91	0.52
1:B:1108:TRP:HB2	1:B:1137:ILE:CD1	2.40	0.52
2:C:491:VAL:HG22	2:C:492:THR:H	1.74	0.52
2:C:810:TYR:HD2	2:C:811:LEU:HD23	1.75	0.52
2:C:905:ILE:HG13	2:C:909:TYR:CD2	2.45	0.52
3:D:370:ASP:HB2	3:D:371:LYS:NZ	2.25	0.52
4:L:56:SER:OG	4:L:59:SER:OG	2.25	0.52
4:L:244:ASN:OD1	4:L:247:ILE:N	2.24	0.52
4:L:405:GLN:N	4:L:421:GLN:O	2.38	0.52
4:M:47:PRO:HB2	4:M:66:MET:HG3	1.91	0.52
4:M:137:TYR:HB2	4:M:174:TYR:HE2	1.75	0.52
4:M:622:ASN:O	4:M:625:ILE:HB	2.10	0.52
4:M:661:THR:O	4:M:664:PHE:HB2	2.09	0.52
5:O:850:ASP:HA	5:O:872:TYR:CD1	2.45	0.52
3:P:75:THR:HB	3:P:96:TRP:CH2	2.44	0.52
3:P:86:TRP:HD1	3:P:130:TYR:OH	1.93	0.52
6:X:186:ARG:NH2	6:X:245:PRO:HA	2.25	0.52
6:X:271:VAL:O	6:X:282:MET:HA	2.10	0.52
6:Z:9:HIS:CE1	6:Z:10:GLN:HG3	2.45	0.52
1:B:914:VAL:HA	1:B:917:ASN:ND2	2.25	0.51
1:B:1034:GLU:OE1	1:B:1207:ARG:NH1	2.43	0.51
2:C:275:GLN:NE2	2:C:278:GLN:OE1	2.43	0.51
2:C:374:ARG:H	2:C:1259:ARG:NH2	2.08	0.51
2:C:606:THR:OG1	2:C:640:THR:N	2.34	0.51
2:C:635:ALA:HA	2:C:638:LEU:HD13	1.92	0.51
2:C:997:LEU:O	2:C:1012:ILE:HG23	2.10	0.51
3:D:253:LEU:HD12	3:D:254:SER:H	1.75	0.51
4:K:179:GLN:HA	4:K:182:LEU:HG	1.92	0.51
4:M:506:LYS:HD2	6:Y:316:GLY:CA	2.39	0.51

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
5:O:584:GLY:N	5:O:587:ASP:OD2	2.40	0.51
5:O:1026:LEU:HD12	5:O:1027:PRO:HD2	1.91	0.51
5:O:1034:PHE:HB3	5:O:1119:VAL:HA	1.92	0.51
5:O:1064:SER:HB3	5:O:1066:GLU:HG2	1.92	0.51
5:O:1258:PRO:HD2	5:O:1261:TRP:CE3	2.45	0.51
3:P:172:SER:HB2	3:P:175:VAL:HG23	1.92	0.51
6:X:315:LEU:HB3	6:X:323:TRP:HD1	1.75	0.51
6:Z:66:LYS:HG3	6:Z:67:HIS:ND1	2.25	0.51
6:Z:261:THR:OG1	6:Z:262:THR:N	2.42	0.51
1:B:258:TRP:CE3	1:B:374:ARG:HD3	2.45	0.51
1:B:295:PRO:HG2	1:B:408:PRO:HB3	1.91	0.51
1:B:746:GLN:NE2	5:O:154:SER:OG	2.43	0.51
1:B:1106:GLY:O	1:B:1135:LEU:HD12	2.10	0.51
1:B:1233:ASN:HA	1:B:1236:THR:OG1	2.11	0.51
2:C:276:VAL:O	2:C:278:GLN:NE2	2.43	0.51
2:C:288:SER:OG	2:C:289:SER:N	2.43	0.51
2:C:643:CYS:SG	2:C:684:TRP:NE1	2.83	0.51
2:C:772:MET:O	2:C:775:LEU:HB3	2.11	0.51
4:K:19:VAL:HB	4:K:248:GLN:HB2	1.92	0.51
4:L:620:ASP:OD2	4:L:623:ASN:N	2.40	0.51
5:O:252:THR:OG1	5:O:253:ASP:N	2.43	0.51
5:O:268:VAL:HG12	5:O:269:ARG:HD3	1.92	0.51
5:O:271:ARG:HB3	5:O:274:GLN:OE1	2.10	0.51
5:O:625:HIS:O	5:O:629:GLN:HG2	2.10	0.51
5:O:647:ASN:HD21	5:O:684:LEU:HB2	1.75	0.51
5:O:756:SER:C	5:O:757:ARG:HD3	2.30	0.51
5:O:809:MET:HA	5:O:812:ASN:ND2	2.25	0.51
5:O:1077:SER:O	5:O:1077:SER:OG	2.28	0.51
6:Y:208:ARG:HH22	6:Y:286:ILE:C	2.14	0.51
6:Y:236:ARG:HD2	6:Y:239:ARG:NE	2.23	0.51
6:Z:251:HIS:CG	6:Z:254:LEU:HD12	2.46	0.51
6:Z:340:LEU:C	6:Z:343:PHE:HE1	2.14	0.51
1:B:1213:ASN:HB2	1:B:1219:THR:OG1	2.10	0.51
3:D:123:ARG:HB3	3:D:125:TYR:CZ	2.44	0.51
3:D:212:MET:O	3:D:215:ILE:HB	2.11	0.51
4:K:288:GLN:O	4:K:292:LEU:HG	2.08	0.51
4:K:563:PRO:HB3	4:L:194:VAL:HG12	1.92	0.51
4:L:303:SER:O	4:L:515:PRO:HD2	2.11	0.51
4:L:405:GLN:NE2	4:L:422:ALA:HA	2.26	0.51
4:L:621:LYS:O	4:L:624:TRP:HB3	2.11	0.51
4:M:293:VAL:HG11	4:M:635:LEU:HD12	1.91	0.51

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
5:O:61:ARG:HE	5:O:64:GLN:CD	2.13	0.51
5:O:316:ARG:HH22	5:O:784:ALA:HA	1.74	0.51
6:X:188:PRO:HG2	6:X:343:PHE:CZ	2.46	0.51
6:Z:74:ASN:OD1	6:Z:75:GLN:N	2.43	0.51
1:B:1047:ILE:HA	1:B:1199:ILE:HG12	1.92	0.51
2:C:641:ASP:OD2	2:C:644:ALA:N	2.42	0.51
2:C:680:THR:O	2:C:684:TRP:NE1	2.43	0.51
2:C:818:GLU:O	2:C:822:ILE:HG13	2.11	0.51
2:C:1076:ILE:HD13	2:C:1109:ILE:HG22	1.93	0.51
2:C:1077:PHE:HA	2:C:1097:GLU:OE2	2.10	0.51
2:C:1146:TYR:HB2	2:C:1180:PHE:CE1	2.39	0.51
3:D:98:ALA:O	3:D:101:LEU:HD23	2.09	0.51
3:D:342:ARG:NH1	3:D:345:ARG:HD3	2.25	0.51
4:L:615:ALA:CB	4:L:618:LEU:HB2	2.39	0.51
4:M:382:LEU:HG	4:M:443:ILE:HD13	1.93	0.51
5:O:68:LEU:O	5:O:193:SER:OG	2.25	0.51
5:O:352:GLN:NE2	5:O:372:LYS:HG2	2.24	0.51
5:O:468:GLY:O	5:O:471:LEU:HB3	2.09	0.51
5:O:550:PHE:HA	5:O:554:VAL:HG11	1.91	0.51
5:O:1080:PHE:HB2	5:O:1087:TRP:CH2	2.45	0.51
6:X:84:ASP:OD1	6:X:85:VAL:N	2.43	0.51
6:Y:25:ILE:HD12	6:Y:40:PRO:HG3	1.93	0.51
1:B:334:LEU:HB2	1:B:335:PHE:CD1	2.45	0.51
1:B:723:ALA:HB3	1:B:726:VAL:HG22	1.93	0.51
2:C:459:VAL:HG11	2:C:1257:VAL:HG22	1.92	0.51
2:C:656:MET:HG3	2:C:660:GLU:HB2	1.91	0.51
2:C:656:MET:SD	2:C:659:PHE:HB2	2.51	0.51
2:C:796:THR:O	2:C:800:LEU:HG	2.11	0.51
2:C:931:LEU:O	2:C:935:VAL:HG23	2.11	0.51
2:C:1037:LEU:HB2	2:C:1038:PHE:CE1	2.44	0.51
4:K:178:ILE:O	4:K:182:LEU:HG	2.11	0.51
4:K:388:LYS:HZ3	4:K:394:PRO:HD2	1.73	0.51
4:K:522:TYR:N	4:K:611:ILE:O	2.44	0.51
4:L:13:VAL:HG23	4:L:249:TRP:HH2	1.74	0.51
4:L:37:SER:C	4:L:41:LEU:HG	2.31	0.51
4:L:64:ARG:NH1	4:L:65:ARG:O	2.44	0.51
4:L:339:LYS:HD3	4:L:340:THR:OG1	2.10	0.51
4:L:618:LEU:H	4:L:618:LEU:HD12	1.76	0.51
5:O:716:PHE:CE2	5:O:781:GLU:HA	2.46	0.51
5:O:1033:ASN:HD21	5:O:1038:GLN:HB3	1.76	0.51
3:P:55:SER:N	3:P:58:VAL:O	2.42	0.51

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:P:168:GLU:OE1	3:P:171:TRP:HB2	2.10	0.51
3:P:276:SER:HB3	3:P:331:GLN:HG2	1.93	0.51
6:X:117:ASP:O	6:X:121:VAL:HG23	2.10	0.51
6:X:318:GLY:O	6:X:322:GLY:N	2.43	0.51
6:Y:204:THR:HG23	6:Y:218:TRP:CD2	2.46	0.51
1:B:433:ARG:HA	1:B:450:VAL:HG22	1.91	0.51
1:B:505:MET:HB2	1:B:507:TYR:CE1	2.45	0.51
1:B:629:PHE:CZ	1:B:652:LEU:HD21	2.46	0.51
1:B:631:ILE:HG23	1:B:883:THR:HG21	1.93	0.51
1:B:1031:PHE:CZ	1:B:1035:TYR:HA	2.46	0.51
1:B:1044:ASP:OD2	1:B:1140:GLU:HG2	2.10	0.51
2:C:405:ILE:O	2:C:408:PRO:HD3	2.11	0.51
2:C:549:LEU:HD12	2:C:890:LYS:C	2.31	0.51
2:C:765:ARG:NE	2:C:769:CYS:SG	2.84	0.51
2:C:1085:PHE:HA	2:C:1092:PRO:CA	2.41	0.51
3:D:169:ASP:OD1	3:D:169:ASP:N	2.38	0.51
4:K:657:ILE:O	4:K:661:THR:HG23	2.10	0.51
4:L:51:ILE:HG23	4:L:52:GLY:H	1.74	0.51
4:L:160:GLN:C	4:L:161:LYS:HD3	2.30	0.51
4:M:534:ASP:HB2	4:M:537:ARG:NH1	2.25	0.51
5:O:340:ASN:CG	5:O:342:THR:H	2.13	0.51
5:O:482:SER:HA	5:O:485:LYS:HD3	1.93	0.51
5:O:511:SER:OG	5:O:572:GLN:NE2	2.42	0.51
5:O:927:ASN:HB3	5:O:949:TYR:HE1	1.76	0.51
3:P:187:PHE:HE2	3:P:284:ASN:HD22	1.59	0.51
6:Y:280:CYS:SG	6:Y:281:LYS:N	2.84	0.51
6:Z:7:ASN:OD1	6:Z:8:GLY:N	2.41	0.51
6:Z:50:VAL:HG22	6:Z:57:VAL:HA	1.93	0.51
6:Z:137:ASP:O	6:Z:140:SER:OG	2.23	0.51
6:Z:257:TYR:HA	6:Z:342:MET:HG3	1.92	0.51
3:D:80:PHE:HA	3:D:84:HIS:NE2	2.26	0.51
3:D:90:ARG:O	3:D:117:LEU:HD12	2.11	0.51
3:D:173:GLY:HA2	3:D:176:ASN:OD1	2.11	0.51
3:D:210:ALA:HB1	3:D:214:GLN:HE22	1.75	0.51
3:D:410:THR:O	3:D:414:ILE:HG13	2.10	0.51
4:K:145:ASP:HA	4:K:161:LYS:HA	1.92	0.51
4:K:406:SER:HB2	4:K:420:GLY:HA3	1.92	0.51
4:K:515:PRO:HA	4:K:518:LEU:HG	1.92	0.51
4:L:159:PHE:CZ	4:L:161:LYS:HD2	2.46	0.51
4:L:468:LEU:HD12	4:L:469:LEU:H	1.75	0.51
4:L:615:ALA:O	4:L:619:SER:N	2.44	0.51

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
4:M:38:PRO:HA	4:M:41:LEU:HD12	1.92	0.51
4:M:334:LEU:HD11	4:M:489:TRP:HZ3	1.76	0.51
5:O:200:TYR:HE1	5:O:229:ASN:HB3	1.76	0.51
5:O:1211:GLN:HE21	5:O:1212:HIS:H	1.58	0.51
6:X:104:CYS:HA	6:X:166:TYR:CZ	2.46	0.51
6:Y:158:ASP:OD2	6:Y:162:LYS:HG3	2.11	0.51
6:Y:180:VAL:O	6:Y:363:ILE:HA	2.11	0.51
6:Z:114:SER:OG	6:Z:116:GLU:HG2	2.11	0.51
6:Z:244:THR:HG22	6:Z:246:ALA:H	1.76	0.51
1:B:1210:PHE:CE1	1:B:1228:PRO:HD2	2.46	0.51
2:C:480:GLU:HA	2:C:483:LEU:HD12	1.93	0.51
2:C:670:THR:H	2:C:673:ARG:HD3	1.75	0.51
3:D:66:TYR:CD1	3:D:394:THR:HA	2.45	0.51
3:D:238:TYR:HE1	3:D:240:GLN:NE2	2.09	0.51
4:K:153:ARG:O	4:K:156:LEU:HB3	2.11	0.51
4:K:348:GLN:HG2	4:K:358:HIS:CD2	2.46	0.51
4:K:437:PHE:HD2	4:L:388:LYS:HZ3	1.58	0.51
4:L:506:LYS:HZ3	6:Z:315:LEU:C	2.14	0.51
4:M:376:MET:HB3	4:M:495:MET:SD	2.51	0.51
5:O:457:PRO:HB2	5:O:459:THR:HG23	1.92	0.51
5:O:1042:LEU:N	5:O:1090:ASP:OD1	2.43	0.51
3:P:230:GLN:O	3:P:256:SER:OG	2.13	0.51
3:P:316:ILE:O	3:P:320:THR:N	2.43	0.51
6:X:54:CYS:HB3	6:X:73:CYS:HB3	1.93	0.51
6:X:105:GLN:O	6:X:109:MET:HG2	2.11	0.51
1:B:303:ARG:HA	1:B:1210:PHE:HB3	1.93	0.51
1:B:847:VAL:O	1:B:870:THR:OG1	2.22	0.51
2:C:798:ASP:HA	2:C:801:LYS:HD3	1.92	0.51
3:D:208:THR:N	3:D:211:ARG:HH21	2.09	0.51
4:K:150:VAL:HG23	4:M:119:GLU:HG2	1.93	0.51
4:K:632:LYS:HE3	4:L:202:ASN:HD22	1.75	0.51
4:K:639:SER:OG	4:L:30:ALA:HB1	2.10	0.51
4:L:233:PRO:O	4:L:236:ALA:HB3	2.11	0.51
4:L:350:THR:HG22	4:L:351:ASP:H	1.75	0.51
4:M:392:TRP:HE3	6:X:278:GLY:HA3	1.76	0.51
4:M:406:SER:OG	4:M:407:LYS:N	2.44	0.51
5:O:303:VAL:HG12	5:O:307:LEU:HD11	1.93	0.51
5:O:616:ILE:HD11	5:O:653:PHE:HB3	1.92	0.51
5:O:739:LYS:HZ3	5:O:756:SER:HA	1.75	0.51
5:O:1257:LEU:HD13	5:O:1277:TYR:CE1	2.46	0.51
6:Y:337:PRO:O	6:Y:340:LEU:HG	2.10	0.51

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
6:Z:79:HIS:CE1	6:Z:83:VAL:HA	2.45	0.51
1:B:256:VAL:HA	1:B:372:LEU:CD1	2.39	0.51
1:B:581:LEU:HD23	1:B:583:PRO:HG3	1.92	0.51
1:B:708:ILE:HD13	1:B:712:TYR:HE1	1.76	0.51
1:B:970:ASN:HA	1:B:973:MET:CE	2.40	0.51
2:C:480:GLU:HB3	2:C:495:SER:HB2	1.92	0.51
2:C:513:GLU:CD	2:C:513:GLU:H	2.14	0.51
2:C:718:GLN:CB	2:C:738:VAL:HG13	2.39	0.51
2:C:721:TYR:CG	2:C:722:GLY:N	2.78	0.51
2:C:999:ILE:N	2:C:1010:ASN:OD1	2.44	0.51
2:C:1212:THR:HG23	2:C:1225:LYS:HB3	1.93	0.51
3:D:206:ASP:OD1	3:D:206:ASP:N	2.43	0.51
3:D:239:PHE:CZ	3:D:253:LEU:HB2	2.46	0.51
4:K:151:SER:H	4:K:154:GLN:HE21	1.58	0.51
4:K:321:TYR:HE1	4:K:532:PRO:HG3	1.76	0.51
4:K:532:PRO:HA	4:K:537:ARG:NH1	2.25	0.51
4:L:265:VAL:O	4:L:268:SER:OG	2.17	0.51
4:L:351:ASP:OD1	4:L:355:THR:N	2.30	0.51
4:L:401:PHE:HB2	4:L:426:ASN:HB3	1.93	0.51
4:L:522:TYR:N	4:L:612:THR:HA	2.24	0.51
4:L:527:LEU:HA	4:L:530:SER:OG	2.11	0.51
4:M:168:PRO:O	4:M:172:THR:HG23	2.11	0.51
4:M:402:ILE:HG13	4:M:423:THR:C	2.31	0.51
4:M:594:LYS:C	4:M:598:ARG:HE	2.14	0.51
5:O:430:TRP:HB3	5:O:476:ARG:HG3	1.93	0.51
5:O:836:ILE:HG23	5:O:837:LEU:HD22	1.93	0.51
6:X:182:ASN:HB3	6:X:362:MET:O	2.11	0.51
6:X:227:GLU:HA	6:X:230:HIS:CG	2.46	0.51
6:X:298:LEU:O	6:X:302:VAL:HG22	2.10	0.51
6:Y:238:TYR:CZ	6:Y:242:LEU:HB2	2.46	0.51
1:B:355:ARG:NH2	1:B:952:ILE:O	2.44	0.50
1:B:437:ALA:HB2	1:B:447:TRP:NE1	2.27	0.50
1:B:714:PRO:HA	1:B:760:GLU:CD	2.32	0.50
1:B:942:GLN:HE21	1:B:994:MET:HE2	1.75	0.50
1:B:1112:LEU:HD11	1:B:1170:ILE:HG12	1.93	0.50
2:C:1067:PHE:CE2	2:C:1107:ASN:HB3	2.46	0.50
2:C:1148:LEU:HA	2:C:1180:PHE:HB2	1.92	0.50
3:D:369:ARG:NH1	3:D:370:ASP:OD1	2.44	0.50
4:K:609:THR:O	4:K:612:THR:N	2.44	0.50
4:L:15:GLY:O	4:L:18:ASN:ND2	2.44	0.50
4:L:105:GLU:HA	4:L:108:ILE:HG13	1.93	0.50

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
4:L:300:ILE:H	4:M:650:LYS:HZ3	1.60	0.50
4:M:152:ALA:HB3	4:M:153:ARG:HH11	1.77	0.50
4:M:233:PRO:HD2	4:M:234:LYS:NZ	2.26	0.50
4:M:402:ILE:O	4:M:468:LEU:HD12	2.11	0.50
5:O:197:LYS:HB3	5:O:199:PHE:CZ	2.46	0.50
5:O:405:ALA:HB3	5:O:772:LEU:HB2	1.93	0.50
5:O:485:LYS:O	5:O:489:VAL:HG13	2.10	0.50
5:O:573:PHE:CE2	5:O:575:TYR:HB2	2.46	0.50
5:O:634:ASN:HA	5:O:661:HIS:HA	1.93	0.50
5:O:693:SER:OG	5:O:696:TYR:HB3	2.11	0.50
5:O:1242:GLU:OE1	5:O:1243:SER:OG	2.22	0.50
6:Z:39:GLN:NE2	6:Z:79:HIS:HD2	2.09	0.50
6:Z:205:GLN:OE1	6:Z:205:GLN:N	2.45	0.50
1:B:473:ASN:HA	1:B:505:MET:O	2.10	0.50
1:B:650:MET:O	1:B:654:ASN:ND2	2.45	0.50
2:C:435:ASP:OD2	2:C:436:ARG:N	2.39	0.50
2:C:549:LEU:HD13	2:C:891:TYR:CZ	2.47	0.50
2:C:715:ASN:HB2	2:C:744:ASP:OD1	2.11	0.50
2:C:726:VAL:HG13	2:C:727:PHE:H	1.75	0.50
2:C:842:ARG:O	2:C:1003:GLN:HG2	2.11	0.50
2:C:1228:PRO:HB3	2:C:1231:ARG:HH11	1.75	0.50
2:C:1233:ASN:HB3	2:C:1244:GLN:NE2	2.23	0.50
3:D:274:PRO:HB2	3:D:278:ASP:OD2	2.11	0.50
3:D:360:LEU:CA	3:D:363:GLN:HE21	2.16	0.50
4:K:351:ASP:OD1	4:K:354:GLY:N	2.45	0.50
4:K:403:VAL:HG12	4:K:423:THR:O	2.10	0.50
4:K:543:ALA:HA	4:K:546:ILE:HG22	1.93	0.50
4:K:655:GLU:O	4:K:659:ASN:ND2	2.45	0.50
4:L:634:SER:HB2	4:M:193:ARG:NH2	2.26	0.50
4:M:344:LEU:HD11	4:M:360:ASN:HD22	1.75	0.50
5:O:172:TYR:CD1	5:O:187:LEU:HD22	2.46	0.50
5:O:491:LYS:O	5:O:494:TYR:HB2	2.11	0.50
5:O:783:GLN:HE22	5:O:785:ARG:HH21	1.59	0.50
3:P:55:SER:OG	3:P:58:VAL:N	2.35	0.50
3:P:174:ASP:O	3:P:177:GLN:HB3	2.11	0.50
3:P:197:ALA:HB3	3:P:203:TRP:HZ2	1.76	0.50
3:P:291:LEU:O	3:P:295:LEU:HG	2.12	0.50
6:X:78:ARG:HB3	6:X:80:GLN:HG2	1.93	0.50
6:X:193:ASN:HD22	6:X:197:LEU:HD13	1.76	0.50
6:X:211:ASP:HB3	6:X:213:ARG:HH21	1.77	0.50
6:Y:203:GLN:O	6:Y:205:GLN:NE2	2.39	0.50

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
6:Z:86:GLN:HA	6:Z:89:ASP:HB2	1.93	0.50
6:Z:270:ALA:HA	6:Z:284:PRO:HA	1.93	0.50
1:B:512:ALA:HA	1:B:515:GLN:NE2	2.25	0.50
1:B:520:ILE:HA	1:B:523:MET:CG	2.41	0.50
2:C:836:VAL:HG11	2:C:844:PRO:HD2	1.93	0.50
2:C:841:ASP:HB3	2:C:1005:ASN:HB3	1.94	0.50
3:D:6:PHE:HA	3:D:144:TYR:CE1	2.47	0.50
4:K:20:PHE:HD1	4:K:247:ILE:HD13	1.77	0.50
4:K:632:LYS:O	4:K:636:ARG:HD2	2.12	0.50
4:L:367:VAL:HG21	4:L:467:TYR:HB3	1.93	0.50
4:M:37:SER:O	4:M:41:LEU:HG	2.12	0.50
4:M:546:ILE:HD11	4:M:631:VAL:HG21	1.94	0.50
5:O:1153:LEU:HA	5:O:1201:VAL:H	1.76	0.50
5:O:1242:GLU:HB2	5:O:1284:TYR:CD1	2.46	0.50
3:P:3:ARG:HH21	3:P:300:GLN:HA	1.76	0.50
3:P:282:ASP:OD1	3:P:284:ASN:N	2.43	0.50
3:P:342:ARG:HA	3:P:345:ARG:HE	1.77	0.50
3:P:353:GLN:O	3:P:356:GLN:HG2	2.11	0.50
1:B:307:HIS:ND1	1:B:310:TRP:CD1	2.79	0.50
1:B:335:PHE:CD1	1:B:335:PHE:N	2.78	0.50
1:B:457:LEU:O	1:B:460:SER:OG	2.22	0.50
1:B:646:VAL:HG22	1:B:688:PHE:CZ	2.45	0.50
1:B:852:GLN:HA	1:B:996:GLN:HG2	1.93	0.50
1:B:1035:TYR:HB2	1:B:1040:ILE:HB	1.93	0.50
2:C:1034:GLU:HB3	2:C:1037:LEU:HD12	1.94	0.50
2:C:1158:ASN:HD21	2:C:1160:TRP:CB	2.24	0.50
3:D:152:LEU:HA	3:D:155:LEU:HD12	1.93	0.50
4:K:237:ALA:HA	4:K:240:LEU:HD12	1.93	0.50
4:K:465:ASN:HB3	4:K:467:TYR:CE1	2.46	0.50
4:L:297:GLU:HB2	4:L:542:ARG:HH21	1.76	0.50
4:M:579:THR:HG22	4:M:584:ARG:HB3	1.93	0.50
3:P:272:VAL:HA	3:P:328:HIS:HB2	1.93	0.50
3:P:275:THR:OG1	3:P:331:GLN:HG3	2.11	0.50
6:X:167:TRP:CE3	6:X:170:LEU:HD12	2.46	0.50
6:Y:10:GLN:HA	6:Y:46:GLY:HA3	1.92	0.50
6:Y:281:LYS:HG2	6:Y:283:TYR:CE1	2.47	0.50
6:Z:43:MET:HG2	6:Z:52:MET:SD	2.51	0.50
6:Z:101:SER:HA	6:Z:104:CYS:SG	2.50	0.50
6:Z:315:LEU:HG	6:Z:320:MET:SD	2.51	0.50
1:B:591:PHE:H	1:B:592:ARG:NH1	2.10	0.50
1:B:706:ALA:HB1	1:B:710:HIS:CE1	2.46	0.50

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:C:354:PHE:O	2:C:358:VAL:HG23	2.11	0.50
2:C:402:TRP:HA	2:C:405:ILE:HD12	1.93	0.50
2:C:465:TRP:CZ2	2:C:469:LEU:HD11	2.46	0.50
3:D:133:LEU:HD12	3:D:140:LYS:HB2	1.92	0.50
4:K:202:ASN:HD22	4:M:633:THR:HB	1.76	0.50
4:K:407:LYS:NZ	4:K:463:GLU:OE2	2.43	0.50
4:L:404:PHE:O	4:L:466:TYR:HA	2.10	0.50
4:L:468:LEU:HG	4:L:489:TRP:CH2	2.47	0.50
4:L:622:ASN:HA	4:L:625:ILE:HD12	1.94	0.50
4:M:595:ILE:N	4:M:598:ARG:HH21	2.10	0.50
5:O:153:LEU:HD23	5:O:157:PHE:CE1	2.46	0.50
5:O:302:GLY:O	5:O:305:SER:HB3	2.11	0.50
5:O:316:ARG:NH2	5:O:784:ALA:HA	2.27	0.50
5:O:329:SER:O	5:O:333:GLN:HG2	2.12	0.50
5:O:518:SER:OG	5:O:526:GLN:N	2.44	0.50
5:O:640:LEU:HG	5:O:653:PHE:HB2	1.94	0.50
5:O:943:ASP:O	5:O:947:LYS:HA	2.11	0.50
5:O:1034:PHE:HB3	5:O:1119:VAL:HG22	1.92	0.50
3:P:350:THR:N	3:P:353:GLN:OE1	2.44	0.50
3:P:367:PHE:O	3:P:371:LYS:NZ	2.33	0.50
6:Y:8:GLY:HA3	6:Y:323:TRP:HE1	1.76	0.50
6:Y:211:ASP:OD2	6:Y:211:ASP:N	2.44	0.50
6:Y:290:ALA:O	6:Y:294:THR:HG23	2.11	0.50
1:B:898:ASN:OD1	1:B:899:VAL:N	2.43	0.50
2:C:355:ARG:HH12	2:C:947:ARG:HH21	1.59	0.50
2:C:790:VAL:HG23	2:C:791:SER:H	1.76	0.50
3:D:241:CYS:HB2	3:D:251:TRP:CE3	2.47	0.50
4:K:358:HIS:HB2	4:K:476:ALA:O	2.12	0.50
4:K:409:PRO:HB2	4:K:412:LEU:HG	1.94	0.50
4:K:426:ASN:ND2	6:Z:61:LEU:HB2	2.27	0.50
4:K:621:LYS:O	4:K:625:ILE:HG13	2.12	0.50
4:L:299:GLU:H	4:L:299:GLU:CD	2.14	0.50
5:O:200:TYR:CD1	5:O:232:HIS:HB3	2.46	0.50
5:O:743:ILE:O	5:O:787:ILE:N	2.27	0.50
5:O:1007:ARG:HA	5:O:1010:GLU:HB2	1.94	0.50
5:O:1029:GLU:OE1	5:O:1029:GLU:N	2.44	0.50
5:O:1181:LEU:HD12	5:O:1182:VAL:N	2.26	0.50
5:O:1215:GLN:HE22	5:O:1216:LEU:HG	1.75	0.50
3:P:46:ILE:HG13	3:P:48:LEU:H	1.75	0.50
3:P:75:THR:HB	3:P:96:TRP:CZ2	2.47	0.50
3:P:397:PHE:HE2	3:P:402:TRP:HE1	1.60	0.50

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
6:X:24:SER:HA	6:X:34:LYS:HE3	1.93	0.50
6:Y:71:HIS:CG	6:Y:72:ARG:N	2.78	0.50
6:Y:105:GLN:HE22	6:Y:136:VAL:HG13	1.76	0.50
1:B:270:PRO:HG3	1:B:297:PRO:HG2	1.92	0.50
1:B:318:ARG:HB2	1:B:371:TYR:CZ	2.47	0.50
1:B:1107:ASN:OD1	1:B:1136:ARG:HD3	2.12	0.50
1:B:1115:TRP:HA	1:B:1122:PHE:CE1	2.46	0.50
2:C:525:ILE:HG12	2:C:532:ILE:HD11	1.94	0.50
2:C:759:ASN:O	2:C:762:THR:OG1	2.20	0.50
2:C:1119:THR:HA	2:C:1122:PHE:HD1	1.76	0.50
3:D:295:LEU:O	3:D:298:GLY:N	2.45	0.50
3:D:372:LEU:HD13	3:D:375:TRP:CE3	2.47	0.50
4:K:136:LYS:HB3	4:K:139:ASP:OD2	2.10	0.50
4:K:563:PRO:HB2	4:L:197:THR:HG21	1.93	0.50
4:L:334:LEU:HB3	4:L:366:ARG:HH22	1.76	0.50
4:M:190:ILE:HA	4:M:193:ARG:HG3	1.94	0.50
4:M:539:MET:HE1	4:M:611:ILE:HD11	1.94	0.50
5:O:246:PRO:HD2	5:O:247:TYR:CE1	2.46	0.50
5:O:708:THR:HA	5:O:754:ILE:O	2.11	0.50
5:O:1052:VAL:HG12	5:O:1053:PHE:N	2.27	0.50
5:O:1198:ILE:HG13	5:O:1214:LEU:HD21	1.93	0.50
3:P:176:ASN:O	3:P:179:LEU:HB3	2.12	0.50
6:X:40:PRO:HG2	6:X:82:TYR:HE2	1.76	0.50
6:X:54:CYS:HB3	6:X:73:CYS:CB	2.42	0.50
6:Z:116:GLU:HG3	6:Z:117:ASP:N	2.26	0.50
6:Z:154:GLN:O	6:Z:156:VAL:HG13	2.12	0.50
6:Z:249:PHE:HB2	6:Z:259:ARG:NH2	2.18	0.50
1:B:350:ASN:HB3	1:B:353:MET:SD	2.51	0.50
1:B:439:MET:HG2	2:C:862:LEU:H	1.77	0.50
1:B:518:GLN:O	1:B:521:ARG:HB2	2.11	0.50
2:C:392:ARG:HG2	2:C:393:SER:O	2.12	0.50
2:C:780:ARG:HG3	2:C:780:ARG:O	2.12	0.50
2:C:1015:MET:HG3	2:C:1016:PRO:O	2.12	0.50
3:D:365:GLN:HA	3:D:368:LYS:HE2	1.93	0.50
4:K:569:GLN:HB2	4:K:592:LEU:HB3	1.92	0.50
4:L:664:PHE:O	4:L:668:VAL:HG23	2.12	0.50
4:M:213:GLN:O	4:M:217:LYS:HG3	2.11	0.50
4:M:425:VAL:HG21	4:M:450:TYR:CD1	2.47	0.50
4:M:581:TYR:HE1	6:Y:75:GLN:HE21	1.60	0.50
5:O:43:TRP:CD1	5:O:45:PRO:HD3	2.43	0.50
5:O:212:TRP:CG	3:P:138:ARG:HD2	2.47	0.50

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
5:O:396:TRP:CH2	5:O:398:PRO:HA	2.47	0.50
5:O:481:ARG:HD3	5:O:481:ARG:N	2.27	0.50
5:O:602:CYS:O	5:O:606:THR:OG1	2.18	0.50
3:P:48:LEU:HB3	3:P:50:ARG:HD2	1.94	0.50
6:X:200:ASP:O	6:X:203:GLN:HG2	2.11	0.50
6:Z:118:LEU:HD12	6:Z:121:VAL:HB	1.93	0.50
6:Z:245:PRO:HA	6:Z:248:ASP:HB2	1.93	0.50
1:B:492:THR:HG23	1:B:1274:VAL:HB	1.94	0.50
1:B:529:ALA:O	1:B:532:ILE:HG22	2.12	0.50
1:B:637:PRO:HG2	1:B:764:TRP:HE1	1.76	0.50
1:B:1110:PHE:CE2	1:B:1114:LEU:HB2	2.46	0.50
2:C:1067:PHE:HE1	2:C:1136:ARG:HB3	1.76	0.50
3:D:7:LEU:HB3	3:D:319:TRP:HE1	1.77	0.50
3:D:85:ARG:NH1	3:D:88:ASP:HB3	2.24	0.50
3:D:283:VAL:O	3:D:287:LEU:HG	2.12	0.50
3:D:341:ASP:HA	3:D:344:ARG:NE	2.27	0.50
4:K:117:ALA:O	4:K:121:ASN:ND2	2.45	0.50
4:K:399:VAL:HG13	4:K:471:THR:C	2.32	0.50
4:K:406:SER:OG	4:K:407:LYS:N	2.44	0.50
4:L:121:ASN:O	4:L:125:LEU:HG	2.12	0.50
4:L:229:ILE:HG13	4:L:233:PRO:HA	1.94	0.50
4:L:427:TYR:HB2	4:L:473:ILE:HD13	1.92	0.50
4:M:457:LEU:HB3	4:M:466:TYR:CE1	2.46	0.50
4:M:631:VAL:O	4:M:635:LEU:HG	2.12	0.50
5:O:487:THR:HG23	5:O:538:ILE:HD12	1.94	0.50
5:O:988:LEU:H	5:O:989:ARG:NH2	2.10	0.50
6:X:272:PHE:CE2	6:X:282:MET:HB2	2.47	0.50
6:X:331:ALA:O	6:X:335:LEU:N	2.26	0.50
6:Z:258:SER:HA	6:Z:273:SER:HA	1.94	0.50
1:B:700:PRO:O	1:B:704:GLN:HG3	2.12	0.49
1:B:806:MET:H	1:B:889:GLY:HA3	1.77	0.49
1:B:985:GLU:CD	1:B:985:GLU:H	2.15	0.49
1:B:1075:HIS:ND1	1:B:1108:TRP:CD1	2.75	0.49
1:B:1086:GLY:CA	1:B:1092:PRO:HG3	2.42	0.49
2:C:622:SER:OG	2:C:623:LEU:N	2.45	0.49
2:C:880:ARG:NH1	2:C:881:ALA:HB2	2.27	0.49
2:C:1084:SER:O	2:C:1093:MET:N	2.31	0.49
2:C:1115:TRP:CD1	2:C:1119:THR:HA	2.47	0.49
4:K:141:LEU:HD12	4:K:165:VAL:N	2.27	0.49
4:K:337:ILE:HG23	4:K:341:MET:HG3	1.93	0.49
4:K:572:LEU:HB3	4:K:592:LEU:HD23	1.94	0.49

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
4:L:362:ARG:NH2	4:L:483:MET:O	2.26	0.49
4:L:435:SER:N	4:L:440:GLN:O	2.36	0.49
4:M:144:VAL:HG12	4:M:145:ASP:N	2.24	0.49
4:M:290:MET:HA	4:M:293:VAL:HG12	1.94	0.49
4:M:581:TYR:HB2	6:Y:73:CYS:H	1.77	0.49
5:O:20:THR:O	5:O:277:THR:HG23	2.12	0.49
5:O:462:ASP:HB2	5:O:465:GLU:OE1	2.12	0.49
5:O:1193:TYR:HH	5:O:1285:THR:HG1	1.59	0.49
3:P:28:HIS:HA	3:P:31:ARG:HG2	1.94	0.49
3:P:133:LEU:O	3:P:136:ASP:N	2.46	0.49
3:P:229:HIS:ND1	3:P:232:ASN:HB3	2.26	0.49
3:P:229:HIS:O	3:P:233:THR:HG22	2.12	0.49
3:P:239:PHE:HB3	3:P:251:TRP:HB3	1.94	0.49
6:Y:54:CYS:HG	6:Y:71:HIS:CE1	2.26	0.49
6:Y:95:TRP:O	6:Y:99:MET:HG2	2.11	0.49
1:B:439:MET:HA	2:C:861:ALA:HA	1.93	0.49
1:B:1115:TRP:HE1	1:B:1119:THR:HG23	1.77	0.49
1:B:1214:SER:O	1:B:1216:SER:N	2.45	0.49
2:C:453:THR:HA	2:C:1253:LEU:O	2.12	0.49
2:C:623:LEU:HD23	2:C:626:LEU:HD13	1.94	0.49
2:C:646:VAL:HG21	2:C:684:TRP:CE2	2.46	0.49
4:K:20:PHE:HB2	4:K:212:MET:SD	2.52	0.49
4:K:563:PRO:CB	4:L:197:THR:HG21	2.42	0.49
4:M:532:PRO:HA	4:M:537:ARG:HE	1.78	0.49
5:O:307:LEU:HD21	5:O:344:LEU:HD11	1.94	0.49
5:O:390:THR:O	5:O:787:ILE:HA	2.12	0.49
5:O:471:LEU:O	5:O:474:LEU:N	2.43	0.49
5:O:762:SER:O	5:O:765:ARG:HB3	2.12	0.49
6:X:51:CYS:O	6:X:55:LEU:N	2.46	0.49
6:X:333:ILE:HD12	6:X:333:ILE:H	1.75	0.49
1:B:272:VAL:HG12	1:B:300:ALA:HB3	1.94	0.49
1:B:309:ARG:NH1	1:B:321:SER:H	2.05	0.49
1:B:518:GLN:OE1	1:B:521:ARG:NE	2.46	0.49
2:C:432:GLY:N	2:C:451:PHE:HB2	2.27	0.49
2:C:542:LEU:HA	2:C:545:ARG:HG2	1.94	0.49
2:C:1104:PHE:HE1	2:C:1126:PHE:CD1	2.30	0.49
2:C:1162:LEU:HD12	2:C:1165:ALA:HB3	1.95	0.49
3:D:42:PHE:HA	3:D:64:ARG:NH2	2.27	0.49
3:D:171:TRP:CH2	3:D:235:ARG:HG3	2.47	0.49
4:K:222:GLN:HA	4:M:589:LYS:HZ3	1.78	0.49
4:K:380:LEU:HB2	4:K:448:LEU:HB3	1.94	0.49

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
4:K:408:ILE:C	4:K:413:TRP:HE1	2.15	0.49
4:L:624:TRP:HA	4:L:627:LEU:HD12	1.94	0.49
4:M:145:ASP:OD2	4:M:161:LYS:HA	2.12	0.49
4:M:272:PRO:HG3	4:M:277:PRO:HA	1.94	0.49
5:O:407:MET:HA	5:O:772:LEU:HD11	1.93	0.49
5:O:419:GLN:HB2	5:O:708:THR:HG23	1.94	0.49
5:O:555:ALA:HB3	5:O:556:ARG:NH1	2.28	0.49
5:O:627:ILE:HA	5:O:631:ILE:HD13	1.94	0.49
5:O:917:SER:OG	5:O:918:THR:N	2.46	0.49
3:P:3:ARG:NH1	3:P:296:CYS:HB3	2.27	0.49
3:P:42:PHE:HA	3:P:64:ARG:NE	2.25	0.49
6:Y:8:GLY:O	6:Y:12:VAL:HG23	2.12	0.49
6:Y:35:THR:HG23	6:Y:153:LEU:HB2	1.94	0.49
6:Y:130:GLU:OE2	6:Y:362:MET:HG2	2.12	0.49
6:Z:255:SER:N	6:Z:276:LEU:HD21	2.28	0.49
1:B:267:LYS:HB3	1:B:310:TRP:CZ2	2.47	0.49
1:B:476:PRO:O	1:B:480:GLU:HG2	2.13	0.49
1:B:1125:GLN:HB2	1:B:1126:PHE:CE2	2.48	0.49
1:B:1144:TYR:HD2	1:B:1145:PRO:C	2.15	0.49
2:C:622:SER:OG	2:C:624:GLU:N	2.43	0.49
2:C:837:ARG:H	2:C:842:ARG:HH12	1.60	0.49
2:C:856:THR:HA	3:P:348:VAL:O	2.13	0.49
2:C:1138:ARG:HD3	2:C:1139:ILE:O	2.12	0.49
4:K:213:GLN:O	4:K:217:LYS:HG3	2.11	0.49
4:K:324:ARG:HE	4:K:490:ASP:HB2	1.77	0.49
4:K:469:LEU:HD12	4:K:489:TRP:NE1	2.27	0.49
4:L:655:GLU:O	4:L:658:GLN:NE2	2.45	0.49
4:M:622:ASN:HA	4:M:625:ILE:HD12	1.93	0.49
5:O:37:ASN:ND2	5:O:40:ARG:HB2	2.27	0.49
5:O:371:ARG:HH12	5:O:375:LEU:HD12	1.76	0.49
5:O:705:GLY:N	5:O:758:ARG:HG3	2.27	0.49
5:O:744:TYR:CD1	5:O:789:PRO:HG3	2.47	0.49
5:O:939:TYR:O	5:O:951:PHE:HA	2.12	0.49
5:O:1190:ALA:O	5:O:1194:LEU:HG	2.12	0.49
5:O:1196:TYR:O	5:O:1214:LEU:N	2.34	0.49
3:P:23:ASP:O	3:P:26:SER:OG	2.26	0.49
3:P:48:LEU:HB3	3:P:50:ARG:CD	2.41	0.49
6:Y:314:ALA:O	6:Y:315:LEU:HD22	2.11	0.49
6:Z:302:VAL:HG22	6:Z:306:TRP:CD1	2.48	0.49
1:B:782:GLN:N	1:B:783:PRO:HD3	2.27	0.49
2:C:307:HIS:CD2	2:C:310:TRP:HB3	2.47	0.49

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:C:351:PRO:HA	2:C:354:PHE:CE1	2.47	0.49
2:C:456:ALA:HA	2:C:459:VAL:HG23	1.94	0.49
2:C:505:MET:O	2:C:507:TYR:N	2.45	0.49
2:C:946:ASP:OD2	2:C:947:ARG:N	2.45	0.49
4:K:348:GLN:HB3	4:K:356:ASN:OD1	2.12	0.49
4:K:353:THR:HG22	6:Z:24:SER:OG	2.13	0.49
4:L:188:TRP:O	4:L:192:LEU:HG	2.13	0.49
4:L:189:GLU:CD	4:L:192:LEU:HD12	2.32	0.49
4:L:322:ASN:ND2	4:L:324:ARG:HB2	2.28	0.49
4:L:468:LEU:HG	4:L:489:TRP:HH2	1.77	0.49
4:L:514:VAL:HB	4:L:517:GLU:HB2	1.95	0.49
4:M:662:GLN:HE22	4:M:665:LEU:HD12	1.78	0.49
5:O:212:TRP:CZ2	3:P:141:HIS:HE1	2.31	0.49
5:O:531:GLU:HB3	5:O:535:GLN:HE22	1.78	0.49
5:O:657:GLY:O	5:O:660:GLN:HB2	2.13	0.49
5:O:662:SER:O	5:O:664:LEU:HG	2.13	0.49
3:P:198:SER:HA	3:P:201:ARG:O	2.13	0.49
6:X:123:THR:HB	6:X:124:GLU:OE1	2.13	0.49
6:Y:196:ARG:HH21	6:Y:353:ASP:C	2.15	0.49
6:Y:243:VAL:HG13	6:Y:349:PHE:CD1	2.47	0.49
6:Z:218:TRP:HB3	6:Z:267:LYS:HB3	1.94	0.49
1:B:1112:LEU:HB2	1:B:1139:ILE:HD11	1.94	0.49
1:B:1141:MET:SD	1:B:1144:TYR:HB3	2.53	0.49
1:B:1166:TRP:CD1	1:B:1176:PRO:HB2	2.48	0.49
2:C:267:LYS:N	2:C:305:VAL:O	2.45	0.49
2:C:518:GLN:O	2:C:522:ILE:HG12	2.11	0.49
2:C:1049:ARG:HG3	2:C:1196:VAL:CG1	2.42	0.49
2:C:1125:GLN:HB2	2:C:1126:PHE:CD1	2.48	0.49
2:C:1128:ALA:O	2:C:1131:LYS:N	2.45	0.49
3:D:213:THR:HG21	3:D:295:LEU:HD21	1.95	0.49
3:D:244:ARG:HE	3:D:336:VAL:HG13	1.77	0.49
3:D:288:SER:O	3:D:292:THR:HG23	2.12	0.49
4:K:219:LEU:O	4:K:223:LEU:HG	2.12	0.49
4:K:527:LEU:O	4:K:530:SER:OG	2.18	0.49
5:O:460:TYR:HB3	5:O:461:PHE:CE1	2.47	0.49
5:O:653:PHE:CE2	5:O:655:ALA:HB2	2.47	0.49
5:O:926:VAL:HG11	5:O:1019:MET:HG2	1.95	0.49
5:O:1048:ASN:HA	5:O:1080:PHE:CE2	2.48	0.49
5:O:1241:LYS:O	5:O:1284:TYR:HA	2.11	0.49
6:X:264:ILE:N	6:X:268:MET:O	2.42	0.49
6:Y:86:GLN:OE1	6:Y:86:GLN:N	2.26	0.49

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
6:Z:199:GLY:O	6:Z:202:ARG:HB2	2.13	0.49
1:B:374:ARG:HH22	1:B:1263:GLU:HG2	1.78	0.49
1:B:651:THR:OG1	1:B:652:LEU:N	2.45	0.49
1:B:762:THR:HA	1:B:765:ARG:NH1	2.27	0.49
1:B:786:THR:O	1:B:790:VAL:HG23	2.13	0.49
1:B:1046:ILE:O	1:B:1199:ILE:HA	2.12	0.49
2:C:272:VAL:H	2:C:299:ALA:HB1	1.78	0.49
2:C:925:CYS:O	2:C:928:VAL:HG22	2.13	0.49
3:D:96:TRP:CD2	3:D:101:LEU:HD22	2.47	0.49
4:K:223:LEU:HB2	4:K:230:ARG:NH2	2.28	0.49
4:K:520:GLY:N	4:K:610:ILE:O	2.20	0.49
4:L:280:GLU:O	4:L:284:LYS:HG2	2.11	0.49
4:L:370:LEU:HD21	4:L:468:LEU:CB	2.42	0.49
4:L:506:LYS:HD2	6:Z:312:ARG:HH12	1.77	0.49
5:O:352:GLN:HA	5:O:370:VAL:O	2.13	0.49
5:O:524:ALA:O	5:O:556:ARG:NH2	2.46	0.49
5:O:995:LEU:HD12	5:O:998:SER:HA	1.95	0.49
3:P:84:HIS:HB3	3:P:130:TYR:CE1	2.48	0.49
3:P:171:TRP:CH2	3:P:235:ARG:HD2	2.48	0.49
3:P:213:THR:HG21	3:P:295:LEU:HD21	1.93	0.49
6:X:97:ARG:HA	6:X:100:LEU:HD12	1.94	0.49
6:Y:7:ASN:OD1	6:Y:9:HIS:N	2.22	0.49
6:Y:227:GLU:HB2	6:Y:354:TYR:CE1	2.48	0.49
6:Z:218:TRP:HB2	6:Z:267:LYS:NZ	2.28	0.49
1:B:340:ASN:ND2	1:B:343:THR:HB	2.27	0.49
1:B:687:CYS:HA	1:B:690:ASN:HB3	1.95	0.49
1:B:708:ILE:HD13	1:B:712:TYR:CE1	2.47	0.49
1:B:1031:PHE:CE1	1:B:1035:TYR:HA	2.48	0.49
1:B:1036:ASN:OD1	1:B:1036:ASN:N	2.46	0.49
1:B:1260:TYR:HB3	1:B:1262:TYR:CD2	2.47	0.49
3:D:50:ARG:HH11	3:D:50:ARG:HA	1.77	0.49
3:D:63:SER:HA	3:D:393:ARG:HB2	1.95	0.49
4:K:232:TYR:HA	4:K:234:LYS:HZ3	1.78	0.49
4:K:259:ASN:OD1	4:K:260:GLU:N	2.46	0.49
4:K:352:ASN:HB3	6:Z:32:TRP:CH2	2.47	0.49
4:K:505:VAL:HG22	4:K:510:VAL:HB	1.93	0.49
4:L:236:ALA:O	4:L:240:LEU:HG	2.13	0.49
4:M:145:ASP:HB3	4:M:159:PHE:HE1	1.78	0.49
5:O:102:PRO:HD2	5:O:105:ASN:HB2	1.94	0.49
5:O:200:TYR:OH	5:O:229:ASN:HB3	2.13	0.49
5:O:207:LEU:HB3	5:O:211:PHE:CZ	2.48	0.49

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
5:O:326:TRP:HE3	5:O:327:LEU:HD23	1.77	0.49
5:O:340:ASN:OD1	5:O:341:GLU:N	2.45	0.49
5:O:506:LEU:C	5:O:507:ARG:HD2	2.32	0.49
5:O:621:ARG:HD2	5:O:625:HIS:HB2	1.95	0.49
3:P:68:MET:HA	3:P:71:LEU:HD12	1.93	0.49
1:B:599:LEU:HA	1:B:832:GLN:NE2	2.28	0.49
1:B:837:ARG:HB3	5:O:238:LEU:HB2	1.95	0.49
2:C:438:GLN:HB3	3:D:201:ARG:NE	2.27	0.49
2:C:468:ARG:NH1	2:C:1018:SER:HA	2.28	0.49
2:C:1212:THR:OG1	2:C:1225:LYS:N	2.40	0.49
3:D:90:ARG:O	3:D:116:ALA:HA	2.12	0.49
3:D:153:LEU:C	3:D:155:LEU:H	2.16	0.49
3:D:284:ASN:HA	3:D:287:LEU:HD12	1.95	0.49
4:K:392:TRP:NE1	4:K:394:PRO:HA	2.28	0.49
4:L:297:GLU:HB2	4:L:542:ARG:NH2	2.28	0.49
4:L:346:GLN:HG3	4:L:360:ASN:HB2	1.94	0.49
4:L:379:VAL:HB	4:L:494:THR:HB	1.95	0.49
4:M:166:ILE:HG12	4:M:167:THR:O	2.13	0.49
4:M:333:TRP:CG	4:M:531:LEU:HD11	2.48	0.49
4:M:401:PHE:CD1	4:M:448:LEU:HD23	2.48	0.49
5:O:642:LYS:NZ	5:O:644:PHE:HA	2.27	0.49
3:P:253:LEU:HD12	3:P:254:SER:H	1.78	0.49
6:Y:139:ASN:HA	6:Y:146:HIS:HB2	1.94	0.49
6:Z:116:GLU:HG3	6:Z:117:ASP:H	1.78	0.49
6:Z:223:TYR:CG	6:Z:356:VAL:HG22	2.47	0.49
6:Z:236:ARG:HD3	6:Z:239:ARG:NH2	2.26	0.49
1:B:414:PRO:HD3	2:C:1118:ASN:ND2	2.28	0.49
1:B:973:MET:O	1:B:976:ALA:N	2.46	0.49
1:B:985:GLU:OE1	1:B:985:GLU:N	2.40	0.49
2:C:338:LEU:HA	2:C:968:TRP:CZ2	2.47	0.49
2:C:474:ILE:CD1	2:C:507:TYR:HB2	2.43	0.49
3:D:22:ASN:HD22	3:D:25:LEU:HD23	1.77	0.49
3:D:94:LEU:HA	3:D:105:VAL:O	2.12	0.49
3:D:250:VAL:HG12	3:D:251:TRP:N	2.25	0.49
4:K:112:THR:O	4:K:116:MET:HG2	2.12	0.49
4:K:335:ARG:HB3	4:K:411:GLU:CD	2.34	0.49
4:L:559:ASP:HA	4:L:597:SER:OG	2.12	0.49
4:M:516:ALA:O	4:M:519:ILE:HG12	2.13	0.49
5:O:448:VAL:HG13	5:O:452:GLU:CD	2.33	0.49
3:P:398:THR:HG23	3:P:401:GLN:HG3	1.95	0.49
6:Z:133:TRP:O	6:Z:136:VAL:HG23	2.13	0.49

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:B:275:GLN:NE2	1:B:1186:SER:OG	2.44	0.48
1:B:684:TRP:HB3	1:B:688:PHE:CD2	2.48	0.48
2:C:304:ILE:H	2:C:304:ILE:HD12	1.78	0.48
2:C:1044:ASP:OD1	2:C:1142:GLY:N	2.26	0.48
2:C:1165:ALA:HA	2:C:1168:GLU:CD	2.33	0.48
4:K:435:SER:N	4:K:441:SER:HA	2.28	0.48
4:L:435:SER:HB3	4:L:441:SER:C	2.34	0.48
4:L:639:SER:OG	4:M:30:ALA:HB1	2.13	0.48
4:M:193:ARG:HA	4:M:196:GLN:HE21	1.78	0.48
4:M:337:ILE:HG13	4:M:361:LEU:HD11	1.95	0.48
4:M:457:LEU:HB3	4:M:466:TYR:CZ	2.48	0.48
5:O:38:PRO:O	5:O:40:ARG:NH2	2.45	0.48
5:O:680:ARG:O	5:O:683:THR:OG1	2.17	0.48
5:O:911:ILE:O	5:O:914:LEU:HG	2.13	0.48
5:O:979:ILE:HG12	5:O:980:THR:N	2.28	0.48
5:O:1248:CYS:SG	5:O:1249:ILE:N	2.85	0.48
3:P:241:CYS:O	3:P:244:ARG:NH1	2.46	0.48
3:P:243:ARG:NE	3:P:248:ALA:O	2.46	0.48
6:Y:60:SER:H	6:Y:63:ARG:NH2	2.11	0.48
6:Z:244:THR:HB	6:Z:247:ARG:CG	2.43	0.48
1:B:773:LYS:NZ	1:B:797:LEU:HB2	2.28	0.48
1:B:778:ASN:HB2	1:B:780:ARG:HG2	1.96	0.48
1:B:1126:PHE:O	1:B:1130:ILE:HG22	2.13	0.48
2:C:263:CYS:HA	2:C:309:ARG:HG2	1.95	0.48
2:C:769:CYS:O	2:C:773:LYS:HG3	2.13	0.48
2:C:836:VAL:HB	2:C:843:VAL:HG22	1.94	0.48
2:C:935:VAL:HG11	2:C:992:PRO:O	2.13	0.48
2:C:1112:LEU:HA	2:C:1139:ILE:HD11	1.94	0.48
3:D:158:PHE:HB3	3:D:275:THR:O	2.13	0.48
3:D:222:LEU:HB3	3:D:227:VAL:HG23	1.95	0.48
3:D:241:CYS:HB2	3:D:251:TRP:CZ3	2.47	0.48
3:D:244:ARG:HA	3:D:244:ARG:HD2	1.51	0.48
4:K:197:THR:HG21	4:M:563:PRO:HB3	1.94	0.48
4:K:224:PRO:HG2	4:K:227:SER:HB3	1.94	0.48
4:K:396:GLY:HA2	4:K:431:TYR:CD1	2.48	0.48
4:L:410:PHE:CD2	4:L:411:GLU:HG3	2.48	0.48
4:L:582:GLY:HA3	6:Z:69:PRO:O	2.12	0.48
4:M:426:ASN:OD1	4:M:427:TYR:N	2.46	0.48
5:O:347:GLN:OE1	5:O:349:ARG:HD3	2.13	0.48
5:O:940:LEU:CA	5:O:950:ARG:O	2.51	0.48
5:O:1079:GLN:NE2	5:O:1080:PHE:O	2.46	0.48

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:P:243:ARG:NH2	3:P:250:VAL:HG23	2.27	0.48
6:X:22:ARG:HH22	6:X:96:LYS:CD	2.26	0.48
6:X:95:TRP:HE1	6:X:266:GLY:N	2.11	0.48
6:X:257:TYR:CD2	6:X:344:SER:HA	2.48	0.48
6:Y:37:SER:HB2	6:Y:90:ARG:HH22	1.77	0.48
6:Y:185:MET:CE	6:Y:187:ASP:H	2.26	0.48
6:Y:253:GLY:HA2	6:Y:275:MET:SD	2.53	0.48
6:Z:161:THR:O	6:Z:165:GLN:NE2	2.30	0.48
6:Z:312:ARG:NH1	6:Z:312:ARG:O	2.45	0.48
1:B:955:LEU:HG	1:B:956:ARG:O	2.13	0.48
2:C:360:HIS:HA	2:C:363:LEU:HB3	1.95	0.48
2:C:839:ASP:OD1	2:C:840:ARG:N	2.45	0.48
2:C:1157:ALA:HB1	2:C:1181:MET:HE2	1.95	0.48
3:D:230:GLN:HG2	3:D:238:TYR:CZ	2.48	0.48
4:K:212:MET:O	4:K:216:VAL:HG23	2.13	0.48
4:K:544:SER:O	4:K:548:GLU:HG2	2.13	0.48
4:L:39:GLY:HA2	4:L:42:ASN:ND2	2.27	0.48
4:L:404:PHE:CE1	4:L:413:TRP:HZ3	2.31	0.48
4:L:474:ASP:CG	4:L:476:ALA:H	2.15	0.48
4:M:151:SER:HB2	4:M:153:ARG:NH2	2.29	0.48
4:M:284:LYS:HZ1	4:M:288:GLN:HB2	1.79	0.48
5:O:15:SER:HB3	5:O:311:GLY:CA	2.43	0.48
5:O:30:LEU:HD11	5:O:34:LEU:HD11	1.94	0.48
5:O:91:ARG:O	5:O:94:LYS:HB2	2.13	0.48
5:O:371:ARG:NH1	5:O:375:LEU:HB2	2.27	0.48
5:O:478:ILE:HG23	5:O:479:GLY:O	2.13	0.48
5:O:609:GLY:N	5:O:658:VAL:O	2.40	0.48
5:O:649:VAL:HG12	5:O:651:LEU:HD11	1.94	0.48
5:O:1063:PHE:CE1	5:O:1073:ILE:HG12	2.45	0.48
3:P:3:ARG:NH2	3:P:301:LEU:HG	2.29	0.48
3:P:50:ARG:HH11	3:P:51:GLY:N	2.12	0.48
3:P:161:ILE:HG12	3:P:327:LEU:CD2	2.43	0.48
6:X:238:TYR:CE2	6:X:242:LEU:HD21	2.48	0.48
6:Y:182:ASN:HA	6:Y:225:TYR:CE2	2.48	0.48
6:Y:238:TYR:HE1	6:Y:242:LEU:HD12	1.78	0.48
6:Z:224:ASP:N	6:Z:355:PRO:O	2.47	0.48
1:B:508:ARG:HG3	3:P:196:GLN:OE1	2.13	0.48
1:B:715:ASN:ND2	1:B:760:GLU:OE2	2.46	0.48
2:C:810:TYR:HA	2:C:814:LEU:HD23	1.94	0.48
2:C:846:MET:HE3	2:C:847:VAL:HG22	1.94	0.48
3:D:88:ASP:OD1	3:D:89:ILE:HG13	2.13	0.48

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:D:218:SER:O	3:D:222:LEU:HG	2.14	0.48
4:K:121:ASN:O	4:K:125:LEU:HG	2.13	0.48
4:K:377:ARG:HH21	4:L:524:PRO:HG3	1.79	0.48
4:K:397:LYS:HB2	4:K:472:PHE:CE2	2.49	0.48
4:L:578:GLU:OE2	6:Z:1:MET:HG3	2.14	0.48
4:M:190:ILE:O	4:M:194:VAL:HG22	2.13	0.48
4:M:410:PHE:HA	4:M:413:TRP:NE1	2.27	0.48
4:M:517:GLU:HB3	6:Y:310:LYS:NZ	2.29	0.48
5:O:305:SER:O	5:O:309:THR:OG1	2.19	0.48
5:O:669:GLY:HA2	5:O:672:PHE:HD1	1.77	0.48
5:O:1157:VAL:HG22	5:O:1196:TYR:HE2	1.79	0.48
3:P:238:TYR:CZ	3:P:256:SER:HB2	2.49	0.48
6:X:136:VAL:O	6:X:143:ARG:HG3	2.13	0.48
6:X:327:THR:HA	6:X:330:GLN:HG2	1.95	0.48
6:Y:79:HIS:CG	6:Y:80:GLN:N	2.81	0.48
6:Y:107:HIS:HA	6:Y:170:LEU:HD21	1.96	0.48
1:B:414:PRO:HA	2:C:1082:ARG:HD2	1.94	0.48
1:B:460:SER:O	1:B:464:ARG:HG3	2.13	0.48
1:B:516:ILE:HA	1:B:519:ILE:HB	1.95	0.48
1:B:629:PHE:CE1	1:B:652:LEU:HD21	2.49	0.48
1:B:646:VAL:HG21	1:B:684:TRP:CH2	2.49	0.48
1:B:646:VAL:HG11	1:B:684:TRP:CE2	2.48	0.48
1:B:1167:LEU:O	1:B:1170:ILE:HG22	2.14	0.48
2:C:501:VAL:HG22	2:C:1263:GLU:OE2	2.14	0.48
2:C:623:LEU:HD22	2:C:626:LEU:HD22	1.95	0.48
2:C:1112:LEU:HD22	2:C:1140:GLU:O	2.14	0.48
3:D:333:ARG:HH12	3:D:334:ARG:CZ	2.26	0.48
4:K:548:GLU:HA	4:K:551:LYS:HE2	1.94	0.48
4:K:583:VAL:HB	6:X:68:LEU:O	2.13	0.48
4:L:625:ILE:HB	4:M:660:TRP:CZ2	2.47	0.48
4:M:632:LYS:NZ	4:M:636:ARG:HD3	2.29	0.48
5:O:296:SER:HG	5:O:299:TYR:H	1.61	0.48
5:O:992:ARG:NH1	5:O:996:LEU:HD11	2.28	0.48
5:O:1238:TRP:CZ3	5:O:1286:LEU:HB3	2.48	0.48
1:B:254:ARG:HG2	1:B:316:PHE:CE2	2.48	0.48
2:C:308:THR:OG1	2:C:322:VAL:HG12	2.14	0.48
2:C:371:TYR:O	2:C:462:ARG:NH1	2.46	0.48
2:C:650:MET:HG2	2:C:675:ALA:HB2	1.96	0.48
2:C:665:ASP:CG	2:C:687:CYS:HB3	2.34	0.48
2:C:803:ILE:HG21	2:C:887:LEU:HG	1.94	0.48
2:C:878:ASP:OD1	2:C:880:ARG:HG2	2.13	0.48

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:D:20:PRO:O	3:D:21:ILE:HD13	2.14	0.48
4:K:227:SER:HB2	4:K:229:ILE:HG22	1.94	0.48
4:M:51:ILE:N	4:M:63:LEU:HD22	2.28	0.48
5:O:172:TYR:CD2	5:O:187:LEU:HB3	2.49	0.48
5:O:649:VAL:HG13	5:O:677:HIS:CE1	2.48	0.48
5:O:771:TYR:HB2	5:O:860:CYS:SG	2.53	0.48
5:O:1049:ALA:HB2	5:O:1080:PHE:HD2	1.79	0.48
5:O:1102:GLN:HE21	5:O:1111:PRO:HB2	1.78	0.48
5:O:1235:MET:SD	5:O:1236:ARG:HD2	2.53	0.48
6:Y:7:ASN:O	6:Y:11:ILE:HG13	2.13	0.48
6:Y:338:ALA:O	6:Y:341:THR:OG1	2.23	0.48
6:Z:94:HIS:HE1	6:Z:153:LEU:HB3	1.77	0.48
2:C:497:TYR:OH	3:D:189:GLU:OE1	2.28	0.48
2:C:522:ILE:HG13	2:C:523:MET:N	2.29	0.48
2:C:773:LYS:O	2:C:776:VAL:HG12	2.13	0.48
2:C:868:ASN:OD1	2:C:869:THR:N	2.46	0.48
2:C:1159:ALA:O	2:C:1163:THR:OG1	2.29	0.48
3:D:96:TRP:CG	3:D:101:LEU:HD22	2.49	0.48
3:D:288:SER:O	3:D:291:LEU:HG	2.13	0.48
3:D:309:VAL:H	3:D:314:ARG:HH21	1.61	0.48
3:D:362:ASP:O	3:D:365:GLN:NE2	2.47	0.48
4:K:49:ILE:O	4:K:64:ARG:HG3	2.13	0.48
4:K:217:LYS:HE2	4:K:675:PRO:HG3	1.95	0.48
4:K:432:ALA:HB1	4:K:442:ILE:O	2.13	0.48
4:L:115:GLU:O	4:L:118:LEU:HB2	2.14	0.48
4:L:234:LYS:O	4:L:238:VAL:HG23	2.13	0.48
4:L:523:THR:N	4:L:526:SER:OG	2.25	0.48
4:M:51:ILE:HB	4:M:63:LEU:HA	1.95	0.48
4:M:426:ASN:ND2	4:M:449:ALA:O	2.36	0.48
5:O:673:PHE:HA	5:O:676:ASP:OD2	2.13	0.48
6:X:128:LEU:HD23	6:X:363:ILE:O	2.13	0.48
6:X:158:ASP:OD2	6:X:161:THR:HB	2.14	0.48
6:X:273:SER:O	6:X:280:CYS:HA	2.14	0.48
6:Y:182:ASN:HA	6:Y:225:TYR:CZ	2.49	0.48
6:Y:223:TYR:HB2	6:Y:225:TYR:HD2	1.78	0.48
6:Z:78:ARG:CG	6:Z:80:GLN:HB3	2.42	0.48
6:Z:158:ASP:CG	6:Z:162:LYS:HG3	2.34	0.48
6:Z:256:HIS:HB2	6:Z:342:MET:HE2	1.95	0.48
1:B:267:LYS:HD2	1:B:310:TRP:CD2	2.48	0.48
1:B:293:GLN:HB2	1:B:404:SER:HB2	1.95	0.48
1:B:492:THR:OG1	2:C:677:ALA:HB2	2.14	0.48

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:B:1121:TYR:O	1:B:1125:GLN:HG2	2.14	0.48
1:B:1147:MET:HG2	1:B:1177:SER:HG	1.78	0.48
2:C:350:ASN:ND2	2:C:1173:THR:O	2.46	0.48
2:C:516:ILE:O	2:C:519:ILE:HB	2.13	0.48
2:C:708:ILE:HA	2:C:711:ARG:HB3	1.96	0.48
2:C:759:ASN:HA	2:C:812:GLN:NE2	2.27	0.48
2:C:878:ASP:O	2:C:882:ILE:HG12	2.14	0.48
2:C:914:VAL:HA	2:C:917:ASN:ND2	2.29	0.48
2:C:989:SER:O	2:C:989:SER:OG	2.25	0.48
2:C:1116:GLN:HB3	2:C:1172:PRO:HA	1.96	0.48
3:D:316:ILE:HB	3:D:320:THR:H	1.78	0.48
4:L:98:GLU:OE1	4:L:169:THR:HG23	2.13	0.48
4:L:369:ASN:HA	4:L:467:TYR:HD2	1.78	0.48
4:L:432:ALA:O	4:L:433:GLU:HG3	2.13	0.48
4:M:47:PRO:C	4:M:66:MET:HG2	2.34	0.48
5:O:67:VAL:HG23	5:O:68:LEU:HG	1.95	0.48
5:O:199:PHE:O	5:O:232:HIS:HA	2.14	0.48
5:O:271:ARG:HB3	5:O:274:GLN:HG2	1.96	0.48
5:O:687:ILE:HD11	5:O:989:ARG:HD3	1.95	0.48
3:P:233:THR:HG21	3:P:256:SER:HB3	1.95	0.48
3:P:241:CYS:SG	3:P:250:VAL:N	2.65	0.48
1:B:468:ARG:HH22	1:B:1017:GLY:N	2.11	0.48
1:B:924:THR:O	1:B:928:VAL:HG23	2.14	0.48
1:B:1105:GLU:HA	1:B:1135:LEU:HB2	1.96	0.48
2:C:293:GLN:NE2	2:C:411:MET:HA	2.29	0.48
2:C:383:THR:HB	2:C:410:ARG:NH2	2.27	0.48
2:C:543:LEU:O	2:C:592:ARG:NH1	2.46	0.48
2:C:614:PRO:HG2	2:C:617:GLY:O	2.13	0.48
2:C:837:ARG:O	2:C:842:ARG:NH2	2.47	0.48
3:D:137:PRO:HA	3:D:140:LYS:HE3	1.96	0.48
3:D:281:MET:SD	3:D:368:LYS:HD2	2.54	0.48
3:D:357:ILE:HD13	3:D:360:LEU:HD12	1.96	0.48
4:L:18:ASN:HA	4:L:249:TRP:HA	1.95	0.48
5:O:58:GLN:HG3	5:O:171:LYS:HG3	1.95	0.48
5:O:299:TYR:OH	5:O:346:PRO:HD2	2.14	0.48
5:O:817:SER:O	5:O:820:TYR:OH	2.31	0.48
5:O:907:PHE:HA	5:O:910:LEU:HD12	1.96	0.48
5:O:924:VAL:C	5:O:1018:ILE:HG23	2.34	0.48
5:O:1172:GLN:HB3	5:O:1184:ASN:HB3	1.95	0.48
5:O:1222:LEU:HD11	5:O:1228:LEU:HD11	1.95	0.48
3:P:84:HIS:HE1	3:P:129:ASP:HB2	1.78	0.48

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
6:X:286:ILE:O	6:X:287:LYS:HD3	2.13	0.48
6:Z:35:THR:O	6:Z:90:ARG:NH1	2.47	0.48
6:Z:161:THR:O	6:Z:165:GLN:HG2	2.13	0.48
6:Z:309:GLU:HB2	6:Z:313:TYR:CE1	2.49	0.48
1:B:375:HIS:ND1	1:B:1261:ALA:HA	2.28	0.48
1:B:501:VAL:HG12	1:B:503:ARG:NH2	2.28	0.48
1:B:714:PRO:HD3	1:B:764:TRP:CH2	2.49	0.48
1:B:847:VAL:HG23	1:B:1000:GLN:O	2.14	0.48
1:B:1033:HIS:O	1:B:1207:ARG:NH1	2.46	0.48
2:C:372:LEU:HB3	2:C:1262:TYR:OH	2.14	0.48
3:D:141:HIS:O	3:D:145:GLN:HG2	2.14	0.48
3:D:175:VAL:O	3:D:178:LEU:HG	2.14	0.48
4:K:584:ARG:O	4:K:585:ILE:HD13	2.14	0.48
4:K:655:GLU:O	4:K:658:GLN:NE2	2.47	0.48
4:L:351:ASP:OD2	4:L:357:TRP:NE1	2.34	0.48
4:M:42:ASN:ND2	4:M:43:PRO:HD2	2.29	0.48
4:M:620:ASP:HB3	4:M:623:ASN:OD1	2.14	0.48
5:O:252:THR:O	5:O:255:MET:HG3	2.14	0.48
5:O:653:PHE:CG	5:O:654:VAL:N	2.82	0.48
5:O:885:ILE:HD12	5:O:921:VAL:O	2.14	0.48
5:O:1054:ASN:HB2	5:O:1056:TYR:HE1	1.79	0.48
3:P:361:ALA:O	3:P:364:THR:OG1	2.27	0.48
1:B:826:LEU:HG	1:B:827:PRO:O	2.14	0.47
2:C:359:LEU:HD13	2:C:934:LEU:HD12	1.95	0.47
4:K:324:ARG:NE	4:K:490:ASP:HB2	2.29	0.47
4:K:666:ASP:O	4:K:669:SER:OG	2.31	0.47
4:L:53:ASP:OD1	4:L:53:ASP:N	2.46	0.47
4:M:26:THR:HB	4:M:246:GLY:HA2	1.96	0.47
4:M:403:VAL:HG12	4:M:423:THR:O	2.13	0.47
4:M:628:ALA:HB3	4:M:629:GLN:NE2	2.29	0.47
5:O:59:LEU:HD21	5:O:63:LEU:HB3	1.96	0.47
5:O:531:GLU:HB3	5:O:535:GLN:NE2	2.28	0.47
5:O:1211:GLN:HE21	5:O:1212:HIS:N	2.11	0.47
6:X:64:LYS:HG3	6:X:66:LYS:HE3	1.96	0.47
6:X:192:PHE:O	6:X:197:LEU:HD11	2.13	0.47
6:Y:331:ALA:C	6:Y:333:ILE:H	2.16	0.47
6:Z:29:GLN:OE1	6:Z:38:ALA:HB3	2.13	0.47
6:Z:167:TRP:CZ3	6:Z:170:LEU:HD23	2.49	0.47
6:Z:220:VAL:HB	6:Z:264:ILE:HG23	1.96	0.47
1:B:949:LEU:HD13	1:B:951:TRP:HE1	1.79	0.47
1:B:1107:ASN:O	1:B:1108:TRP:HD1	1.98	0.47

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:C:337:GLN:HE21	2:C:361:MET:HB2	1.78	0.47
2:C:991:ASP:OD2	2:C:993:ARG:HB2	2.14	0.47
3:D:207:GLY:O	3:D:211:ARG:NE	2.46	0.47
4:K:285:LEU:HA	4:K:288:GLN:HG3	1.96	0.47
4:K:443:ILE:HG13	4:K:444:ALA:O	2.15	0.47
4:L:124:PHE:CE2	4:L:128:LEU:HD11	2.49	0.47
4:L:227:SER:O	4:L:231:ARG:NH2	2.47	0.47
4:L:330:GLU:OE2	6:Z:9:HIS:N	2.35	0.47
4:L:396:GLY:HA2	4:L:429:GLN:HE22	1.79	0.47
4:L:522:TYR:HB2	4:L:611:ILE:HG13	1.96	0.47
4:M:328:ILE:O	6:Y:326:ARG:NH1	2.34	0.47
4:M:524:PRO:O	4:M:527:LEU:HB3	2.15	0.47
5:O:85:PHE:CE2	5:O:89:LYS:HD2	2.49	0.47
5:O:411:ASP:O	5:O:414:VAL:HG22	2.15	0.47
5:O:831:GLY:H	5:O:834:ALA:HB2	1.79	0.47
5:O:1136:PHE:HE2	5:O:1229:PHE:H	1.62	0.47
5:O:1215:GLN:NE2	5:O:1216:LEU:HG	2.28	0.47
5:O:1236:ARG:O	5:O:1238:TRP:HD1	1.97	0.47
3:P:145:GLN:HE21	3:P:297:SER:CB	2.27	0.47
3:P:225:VAL:HB	3:P:227:VAL:HG23	1.95	0.47
6:X:110:MET:O	6:X:113:VAL:HG23	2.13	0.47
6:X:309:GLU:HB3	6:X:313:TYR:CE2	2.49	0.47
6:X:358:ILE:H	6:X:358:ILE:HD12	1.79	0.47
6:Y:71:HIS:CE1	6:Y:73:CYS:HB2	2.48	0.47
6:Z:94:HIS:CE1	6:Z:153:LEU:HD23	2.49	0.47
6:Z:199:GLY:HA2	6:Z:202:ARG:NH2	2.29	0.47
1:B:318:ARG:CZ	1:B:371:TYR:HA	2.45	0.47
1:B:589:SER:CA	1:B:592:ARG:HH22	2.27	0.47
1:B:609:ASP:HB2	1:B:612:SER:OG	2.14	0.47
1:B:691:ILE:O	1:B:703:ARG:NH1	2.47	0.47
1:B:779:GLN:HA	1:B:782:GLN:HE21	1.79	0.47
1:B:825:MET:HA	1:B:911:ASP:OD1	2.15	0.47
1:B:1048:GLY:O	1:B:1050:VAL:HG23	2.14	0.47
1:B:1150:TYR:HD2	1:B:1184:ILE:HG12	1.78	0.47
2:C:477:THR:HA	2:C:480:GLU:CD	2.35	0.47
2:C:1122:PHE:HB3	2:C:1126:PHE:HE2	1.79	0.47
2:C:1150:TYR:HB3	2:C:1184:ILE:HG12	1.97	0.47
3:D:30:LEU:HD23	3:D:33:GLY:C	2.35	0.47
4:K:98:GLU:HG3	4:K:99:PRO:HD2	1.96	0.47
4:K:322:ASN:OD1	4:K:323:VAL:N	2.47	0.47
4:K:382:LEU:HD21	4:K:491:ALA:HA	1.96	0.47

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
4:L:300:ILE:HG12	4:M:650:LYS:HZ1	1.79	0.47
4:M:176:ASP:HA	4:M:179:GLN:NE2	2.29	0.47
4:M:257:VAL:HG22	4:M:258:MET:H	1.80	0.47
4:M:398:LYS:NZ	4:M:474:ASP:OD1	2.38	0.47
5:O:94:LYS:O	5:O:98:LEU:HG	2.15	0.47
5:O:815:VAL:O	5:O:819:VAL:HG23	2.15	0.47
5:O:1054:ASN:HB3	5:O:1061:LEU:CD1	2.43	0.47
5:O:1232:ALA:HB3	5:O:1274:ILE:HG23	1.96	0.47
6:X:148:SER:N	6:X:151:ASP:OD2	2.41	0.47
6:X:286:ILE:HG22	6:X:287:LYS:H	1.79	0.47
6:Z:10:GLN:H	6:Z:10:GLN:CD	2.10	0.47
1:B:281:LEU:HD21	1:B:286:PHE:CE2	2.49	0.47
1:B:304:ILE:HD12	1:B:305:VAL:H	1.79	0.47
1:B:347:ARG:HB3	1:B:1176:PRO:HB3	1.96	0.47
1:B:593:VAL:HG23	1:B:594:ALA:N	2.29	0.47
1:B:707:GLU:HG2	1:B:711:ARG:CZ	2.43	0.47
1:B:708:ILE:HG23	1:B:712:TYR:HD1	1.79	0.47
1:B:762:THR:HA	1:B:765:ARG:CZ	2.44	0.47
3:D:123:ARG:HB3	3:D:125:TYR:CE1	2.50	0.47
3:D:267:ASN:ND2	3:D:269:SER:OG	2.47	0.47
3:D:283:VAL:O	3:D:286:ILE:HG22	2.15	0.47
4:K:148:VAL:O	4:K:158:ASN:HB3	2.14	0.47
4:K:194:VAL:O	4:K:197:THR:OG1	2.25	0.47
4:K:273:SER:OG	4:L:179:GLN:OE1	2.31	0.47
4:K:338:PRO:HB2	4:K:341:MET:HG2	1.96	0.47
4:L:273:SER:HA	4:M:179:GLN:NE2	2.30	0.47
4:L:378:PHE:HE2	4:L:452:TYR:HB3	1.80	0.47
4:M:160:GLN:CG	4:M:161:LYS:H	2.27	0.47
4:M:327:LYS:HD2	4:M:328:ILE:N	2.29	0.47
4:M:395:ASN:N	4:M:397:LYS:HD3	2.29	0.47
4:M:568:ILE:O	4:M:572:LEU:HG	2.14	0.47
5:O:616:ILE:C	5:O:651:LEU:HB2	2.35	0.47
5:O:641:ILE:HG13	5:O:643:PRO:HD3	1.96	0.47
5:O:769:LEU:HA	5:O:861:TRP:HA	1.96	0.47
3:P:307:ASN:HA	3:P:314:ARG:CZ	2.44	0.47
6:X:56:GLY:O	6:X:58:VAL:HG13	2.14	0.47
6:X:337:PRO:O	6:X:340:LEU:N	2.39	0.47
6:Y:227:GLU:HA	6:Y:230:HIS:ND1	2.30	0.47
1:B:336:LYS:HZ2	1:B:345:SER:C	2.18	0.47
1:B:406:MET:HB3	1:B:407:TYR:CE1	2.50	0.47
1:B:518:GLN:NE2	1:B:828:PHE:HD1	2.11	0.47

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:B:702:LEU:HA	1:B:705:TRP:HB2	1.96	0.47
1:B:1007:ARG:NH1	1:B:1007:ARG:HA	2.29	0.47
1:B:1094:ILE:HD12	1:B:1095:ARG:H	1.79	0.47
1:B:1146:TYR:HA	1:B:1178:VAL:N	2.29	0.47
1:B:1253:LEU:HD12	1:B:1253:LEU:HA	1.68	0.47
2:C:419:ILE:HG23	2:C:420:CYS:H	1.79	0.47
2:C:446:GLU:HG3	2:C:447:TRP:CD1	2.50	0.47
2:C:482:ALA:O	2:C:485:GLU:HB3	2.14	0.47
2:C:541:VAL:O	2:C:544:GLN:HB3	2.14	0.47
2:C:851:ARG:NH1	2:C:863:SER:OG	2.48	0.47
3:D:18:ASN:HD22	3:D:91:PHE:HZ	1.63	0.47
3:D:21:ILE:HG22	3:D:22:ASN:N	2.30	0.47
3:D:137:PRO:O	3:D:140:LYS:HB3	2.14	0.47
3:D:165:ARG:NH2	3:D:167:ASP:OD2	2.48	0.47
3:D:339:PHE:O	3:D:343:LEU:HD23	2.14	0.47
4:K:324:ARG:NH2	4:K:490:ASP:HB2	2.30	0.47
4:L:426:ASN:OD1	4:L:448:LEU:HD11	2.14	0.47
4:M:471:THR:HG22	4:M:472:PHE:N	2.30	0.47
5:O:13:LEU:HD22	5:O:311:GLY:C	2.34	0.47
5:O:57:VAL:HG23	5:O:175:MET:HE2	1.96	0.47
5:O:737:ALA:HB1	5:O:757:ARG:NH2	2.30	0.47
5:O:1199:ARG:HG2	5:O:1211:GLN:OE1	2.14	0.47
5:O:1263:VAL:HA	5:O:1277:TYR:CD1	2.50	0.47
3:P:361:ALA:O	3:P:365:GLN:HG2	2.15	0.47
6:Y:180:VAL:H	6:Y:364:LEU:H	1.63	0.47
1:B:846:MET:SD	1:B:870:THR:HG21	2.55	0.47
1:B:1041:ALA:HB2	1:B:1146:TYR:HE1	1.80	0.47
2:C:931:LEU:HD11	2:C:987:LEU:HD23	1.97	0.47
2:C:1003:GLN:CD	2:C:1007:ARG:HB3	2.35	0.47
2:C:1120:ARG:N	2:C:1120:ARG:HD2	2.29	0.47
3:D:137:PRO:O	3:D:141:HIS:ND1	2.47	0.47
3:D:174:ASP:HA	3:D:177:GLN:NE2	2.29	0.47
3:D:344:ARG:HE	3:D:354:GLN:NE2	2.13	0.47
4:K:24:ALA:HB1	4:K:205:ILE:HD12	1.95	0.47
4:K:40:MET:HE2	4:K:40:MET:HB2	1.65	0.47
4:K:229:ILE:HG23	4:K:230:ARG:HD3	1.96	0.47
4:K:284:LYS:HA	4:K:287:GLU:OE1	2.15	0.47
4:K:463:GLU:HG3	4:K:464:MET:HG2	1.96	0.47
4:L:216:VAL:HA	4:L:219:LEU:HG	1.96	0.47
4:L:242:LYS:HE3	4:L:243:ARG:HH21	1.80	0.47
4:L:368:VAL:O	4:L:467:TYR:HA	2.15	0.47

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
4:M:67:THR:OG1	4:M:68:SER:N	2.47	0.47
4:M:581:TYR:CD2	6:Y:71:HIS:CE1	3.02	0.47
5:O:290:GLN:OE1	3:P:135:ARG:HG2	2.15	0.47
5:O:635:ILE:HA	5:O:657:GLY:O	2.15	0.47
5:O:1192:LYS:CE	5:O:1220:ILE:H	2.28	0.47
3:P:46:ILE:HG13	3:P:48:LEU:N	2.29	0.47
6:X:183:PHE:CD1	6:X:261:THR:HB	2.49	0.47
6:Y:117:ASP:O	6:Y:121:VAL:HG22	2.13	0.47
6:Y:236:ARG:HH11	6:Y:239:ARG:NE	2.12	0.47
6:Z:347:THR:HG22	6:Z:349:PHE:N	2.27	0.47
1:B:267:LYS:HB3	1:B:310:TRP:CH2	2.50	0.47
1:B:433:ARG:HG3	1:B:449:ASP:O	2.15	0.47
1:B:622:SER:OG	1:B:624:GLU:OE1	2.20	0.47
1:B:1151:TYR:O	1:B:1184:ILE:HG13	2.15	0.47
1:B:1189:ASP:OD1	1:B:1190:ILE:N	2.48	0.47
1:B:1228:PRO:HB2	1:B:1230:GLU:HB2	1.96	0.47
2:C:381:ASP:HB2	2:C:391:LEU:HD21	1.96	0.47
2:C:418:LYS:HE3	2:C:1214:SER:HA	1.97	0.47
2:C:516:ILE:HG13	2:C:517:SER:N	2.30	0.47
2:C:599:LEU:C	2:C:832:GLN:HE21	2.17	0.47
2:C:810:TYR:HD1	2:C:814:LEU:HB2	1.80	0.47
2:C:913:GLU:HB3	2:C:917:ASN:OD1	2.14	0.47
2:C:998:ALA:HB1	2:C:1010:ASN:ND2	2.26	0.47
2:C:1077:PHE:HB3	2:C:1081:CYS:SG	2.55	0.47
2:C:1121:TYR:O	2:C:1124:GLN:HB2	2.15	0.47
2:C:1185:SER:CB	2:C:1213:ASN:HD21	2.27	0.47
2:C:1229:VAL:HG23	2:C:1236:THR:HG21	1.95	0.47
3:D:107:ALA:HB3	3:D:112:LEU:CD2	2.45	0.47
3:D:306:ASN:HD21	3:D:320:THR:HG22	1.80	0.47
3:D:336:VAL:O	3:D:339:PHE:HB3	2.15	0.47
4:K:16:ASP:N	4:K:16:ASP:OD1	2.46	0.47
4:K:120:PHE:CE2	4:K:242:LYS:HG2	2.50	0.47
4:K:401:PHE:HA	4:K:470:ALA:HA	1.95	0.47
4:K:405:GLN:HG3	4:K:421:GLN:HB2	1.96	0.47
4:L:10:THR:OG1	4:L:11:ILE:HD12	2.14	0.47
4:L:47:PRO:HD3	4:L:102:VAL:HG13	1.96	0.47
4:L:48:TRP:HD1	4:L:148:VAL:HA	1.80	0.47
4:L:563:PRO:HG3	4:M:190:ILE:HD11	1.96	0.47
4:M:205:ILE:HD12	4:M:651:LEU:HD12	1.95	0.47
4:M:229:ILE:HD13	4:M:232:TYR:HB2	1.96	0.47
4:M:382:LEU:HD23	4:M:382:LEU:H	1.79	0.47

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
4:M:388:LYS:NZ	4:M:389:GLU:O	2.40	0.47
4:M:395:ASN:H	4:M:397:LYS:HD3	1.80	0.47
4:M:576:GLN:NE2	4:M:587:ASN:H	2.13	0.47
4:M:581:TYR:CD2	6:Y:73:CYS:HB3	2.50	0.47
4:M:661:THR:O	4:M:665:LEU:HG	2.14	0.47
5:O:44:LYS:HB2	5:O:60:PHE:HZ	1.79	0.47
5:O:218:ALA:HB1	5:O:236:GLY:O	2.15	0.47
5:O:267:ASN:ND2	5:O:311:GLY:HA2	2.30	0.47
5:O:509:ARG:HH22	5:O:543:PRO:HD2	1.79	0.47
5:O:530:ILE:HG23	5:O:533:TRP:CE3	2.50	0.47
5:O:648:ASN:HA	5:O:681:TYR:CZ	2.50	0.47
5:O:650:GLU:C	5:O:651:LEU:HD12	2.35	0.47
5:O:944:SER:HA	5:O:947:LYS:HD3	1.96	0.47
5:O:1004:ALA:O	5:O:1008:ILE:HG12	2.15	0.47
3:P:305:ASP:OD1	3:P:306:ASN:N	2.47	0.47
3:P:373:GLU:O	3:P:377:ARG:HG2	2.14	0.47
6:X:14:LEU:HD13	6:X:45:CYS:HA	1.96	0.47
6:X:224:ASP:H	6:X:354:TYR:HB3	1.80	0.47
6:Y:22:ARG:HH22	6:Y:96:LYS:HG3	1.80	0.47
6:Y:51:CYS:O	6:Y:55:LEU:HA	2.13	0.47
6:Y:263:PRO:HA	6:Y:268:MET:C	2.35	0.47
6:Z:322:GLY:HA2	6:Z:325:ASN:ND2	2.30	0.47
1:B:1109:ILE:CD1	1:B:1138:ARG:HB3	2.44	0.47
2:C:283:TYR:HB2	2:C:290:TYR:CB	2.43	0.47
2:C:982:MET:HA	2:C:985:GLU:CD	2.35	0.47
2:C:1239:ASP:OD1	2:C:1239:ASP:N	2.48	0.47
3:D:36:PRO:HG2	3:D:37:TRP:CD1	2.50	0.47
3:D:207:GLY:C	3:D:211:ARG:HE	2.18	0.47
3:D:373:GLU:C	3:D:377:ARG:HE	2.14	0.47
4:K:137:TYR:CE1	4:K:140:LEU:HD13	2.50	0.47
4:K:143:TYR:HA	4:K:163:VAL:O	2.15	0.47
4:K:401:PHE:CE1	4:K:448:LEU:HG	2.49	0.47
4:K:565:SER:OG	4:K:600:MET:SD	2.73	0.47
4:L:48:TRP:HB3	4:L:63:LEU:HD23	1.95	0.47
4:L:571:GLN:HE22	4:L:572:LEU:HD23	1.80	0.47
4:M:123:GLU:OE2	4:M:127:LYS:HE2	2.14	0.47
4:M:414:THR:OG1	4:M:418:GLN:NE2	2.42	0.47
4:M:617:VAL:C	4:M:619:SER:H	2.18	0.47
5:O:1056:TYR:CE1	5:O:1061:LEU:HD13	2.50	0.47
6:X:176:SER:OG	6:X:177:SER:N	2.47	0.47
6:Y:79:HIS:HA	6:Y:82:TYR:CE2	2.50	0.47

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
6:Y:105:GLN:OE1	6:Y:136:VAL:HG13	2.15	0.47
6:Y:197:LEU:HB3	6:Y:201:ALA:HB1	1.96	0.47
6:Y:310:LYS:HZ2	6:Y:313:TYR:HD2	1.61	0.47
6:Z:102:PHE:HA	6:Z:105:GLN:OE1	2.15	0.47
6:Z:132:ASN:HB3	6:Z:135:GLN:CD	2.35	0.47
1:B:279:ASP:OD1	1:B:280:VAL:N	2.46	0.47
2:C:634:LEU:O	2:C:637:PRO:HD2	2.14	0.47
2:C:669:TYR:HB3	2:C:673:ARG:CZ	2.45	0.47
2:C:682:HIS:HB2	3:P:173:GLY:CA	2.45	0.47
3:D:96:TRP:CZ3	3:D:104:LEU:HD11	2.49	0.47
4:K:222:GLN:HE22	4:M:589:LYS:HB3	1.78	0.47
4:K:308:PRO:HA	4:K:511:SER:O	2.15	0.47
4:K:324:ARG:HH21	4:K:490:ASP:HB2	1.80	0.47
4:L:113:LYS:HE2	4:M:40:MET:HA	1.96	0.47
4:L:381:ASP:O	4:L:491:ALA:HB1	2.14	0.47
4:M:30:ALA:O	4:M:32:PRO:HD3	2.14	0.47
4:M:358:HIS:HB2	4:M:476:ALA:O	2.15	0.47
5:O:56:ALA:HA	5:O:175:MET:HG2	1.97	0.47
5:O:61:ARG:HG3	5:O:64:GLN:HG3	1.97	0.47
5:O:456:LEU:HD11	5:O:460:TYR:CG	2.50	0.47
5:O:1054:ASN:ND2	5:O:1104:LEU:HD21	2.29	0.47
5:O:1243:SER:OG	5:O:1283:ASP:HB3	2.14	0.47
3:P:80:PHE:HA	3:P:84:HIS:HD2	1.80	0.47
3:P:80:PHE:HA	3:P:84:HIS:CD2	2.50	0.47
3:P:335:MET:O	3:P:338:GLU:HG3	2.15	0.47
3:P:377:ARG:HA	3:P:380:ASP:OD1	2.15	0.47
6:X:10:GLN:HB3	6:X:45:CYS:SG	2.55	0.47
6:X:228:LEU:HD22	6:X:239:ARG:HD3	1.97	0.47
6:Y:192:PHE:HD1	6:Y:206:PHE:HE2	1.63	0.47
6:Y:236:ARG:HG2	6:Y:239:ARG:HH21	1.79	0.47
1:B:600:TYR:CE2	1:B:830:PRO:HA	2.50	0.47
1:B:623:LEU:HG	1:B:627:TRP:NE1	2.29	0.47
1:B:770:GLU:OE2	1:B:774:ASN:ND2	2.48	0.47
1:B:849:VAL:H	1:B:869:THR:N	2.13	0.47
1:B:851:ARG:NH1	1:B:986:PRO:HA	2.30	0.47
2:C:474:ILE:HG22	2:C:475:ASN:H	1.80	0.47
2:C:682:HIS:NE2	3:P:177:GLN:OE1	2.47	0.47
3:D:223:SER:N	3:D:228:ILE:HD12	2.30	0.47
4:L:193:ARG:CZ	4:L:196:GLN:HB2	2.45	0.47
4:L:207:GLU:HG3	4:L:208:VAL:N	2.29	0.47
4:M:380:LEU:HD11	4:M:491:ALA:HB3	1.97	0.47

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
5:O:438:TYR:HB3	5:O:641:ILE:HD13	1.97	0.47
5:O:923:LEU:HA	5:O:1019:MET:O	2.15	0.47
5:O:1074:PRO:HG2	5:O:1093:PHE:C	2.34	0.47
3:P:35:SER:OG	3:P:37:TRP:O	2.27	0.47
3:P:61:ALA:HA	3:P:64:ARG:CB	2.45	0.47
3:P:336:VAL:HA	3:P:339:PHE:HB3	1.96	0.47
6:X:26:TYR:CE1	6:X:31:GLY:HA2	2.50	0.47
6:Y:180:VAL:N	6:Y:364:LEU:H	2.13	0.47
1:B:1049:ARG:HH12	1:B:1132:THR:HG23	1.80	0.46
2:C:392:ARG:HH21	2:C:394:LEU:HD11	1.80	0.46
2:C:606:THR:N	2:C:640:THR:HG1	2.12	0.46
2:C:1104:PHE:HB2	2:C:1129:TRP:CE3	2.50	0.46
2:C:1107:ASN:O	2:C:1108:TRP:HD1	1.98	0.46
3:D:149:ALA:HA	3:D:152:LEU:HD23	1.97	0.46
3:D:174:ASP:N	3:D:174:ASP:OD2	2.48	0.46
4:K:272:PRO:HA	4:K:276:ALA:O	2.14	0.46
4:K:321:TYR:OH	4:K:530:SER:O	2.30	0.46
4:K:590:GLY:O	4:K:593:SER:OG	2.32	0.46
4:L:402:ILE:HG23	4:L:424:VAL:N	2.31	0.46
4:L:457:LEU:HD23	4:L:457:LEU:HA	1.68	0.46
4:L:569:GLN:NE2	4:L:592:LEU:HB2	2.29	0.46
4:M:229:ILE:C	4:M:232:TYR:H	2.19	0.46
4:M:650:LYS:O	4:M:653:SER:OG	2.26	0.46
5:O:728:SER:HA	5:O:731:CYS:SG	2.55	0.46
5:O:1019:MET:HB3	5:O:1021:ILE:HD11	1.97	0.46
5:O:1230:LEU:HB2	5:O:1279:VAL:HG21	1.97	0.46
3:P:152:LEU:HD21	3:P:319:TRP:CE2	2.51	0.46
6:Y:222:VAL:HG22	6:Y:359:GLY:H	1.79	0.46
6:Z:132:ASN:HD21	6:Z:134:LEU:HB2	1.80	0.46
6:Z:320:MET:HA	6:Z:323:TRP:HB3	1.96	0.46
1:B:441:ASN:ND2	2:C:864:LEU:HG	2.29	0.46
1:B:705:TRP:CE3	1:B:705:TRP:HA	2.49	0.46
1:B:746:GLN:HE22	5:O:154:SER:C	2.13	0.46
1:B:807:THR:HA	1:B:810:TYR:HB3	1.96	0.46
1:B:1119:THR:HG21	1:B:1170:ILE:HG23	1.96	0.46
2:C:432:GLY:H	2:C:451:PHE:HB2	1.79	0.46
2:C:591:PHE:O	2:C:595:LEU:HG	2.14	0.46
2:C:854:ARG:NE	3:P:258:ASN:HB3	2.30	0.46
2:C:969:VAL:O	2:C:973:MET:HG2	2.14	0.46
2:C:1189:ASP:OD1	2:C:1190:ILE:N	2.45	0.46
4:K:24:ALA:HB2	4:K:206:GLY:H	1.79	0.46

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
4:K:171:GLN:HA	4:K:174:TYR:HB3	1.97	0.46
4:K:175:VAL:HA	4:K:178:ILE:HG12	1.97	0.46
4:K:260:GLU:HB3	4:L:65:ARG:HH12	1.80	0.46
4:K:358:HIS:CG	4:K:477:ALA:HA	2.50	0.46
4:L:187:LYS:HA	4:L:190:ILE:HD12	1.97	0.46
4:L:629:GLN:HA	4:L:632:LYS:HZ3	1.78	0.46
4:M:263:ASN:ND2	4:M:267:ALA:HB2	2.30	0.46
4:M:337:ILE:HB	4:M:365:THR:N	2.30	0.46
4:M:448:LEU:HA	4:M:448:LEU:HD12	1.75	0.46
4:M:463:GLU:OE2	4:M:463:GLU:N	2.45	0.46
5:O:371:ARG:NH2	5:O:375:LEU:HB2	2.29	0.46
5:O:581:VAL:HG23	5:O:582:VAL:N	2.30	0.46
5:O:594:LEU:O	5:O:598:LEU:HG	2.16	0.46
5:O:1264:LEU:HG	5:O:1277:TYR:HA	1.97	0.46
3:P:221:TRP:O	3:P:224:TYR:HB3	2.16	0.46
6:X:177:SER:HB2	6:X:179:LEU:HD11	1.96	0.46
6:Y:141:MET:HA	6:Y:144:SER:O	2.16	0.46
6:Z:97:ARG:NH2	6:Z:156:VAL:HG22	2.29	0.46
6:Z:236:ARG:HA	6:Z:239:ARG:HE	1.80	0.46
1:B:338:LEU:HG	1:B:968:TRP:CD2	2.50	0.46
1:B:488:GLN:NE2	1:B:905:ILE:HG21	2.30	0.46
1:B:657:VAL:HG22	1:B:671:GLN:NE2	2.30	0.46
1:B:772:MET:O	1:B:775:LEU:HB3	2.14	0.46
1:B:1185:SER:OG	1:B:1186:SER:N	2.49	0.46
2:C:294:SER:OG	2:C:296:PHE:O	2.33	0.46
2:C:745:HIS:CG	2:C:816:PRO:HG3	2.51	0.46
2:C:807:THR:OG1	2:C:885:ALA:O	2.26	0.46
2:C:856:THR:HA	3:P:348:VAL:C	2.36	0.46
2:C:1147:MET:HG2	2:C:1177:SER:OG	2.16	0.46
3:D:169:ASP:OD1	3:D:170:MET:HE2	2.15	0.46
3:D:182:TYR:HA	3:D:185:HIS:CE1	2.51	0.46
3:D:350:THR:HG23	3:D:353:GLN:H	1.80	0.46
4:K:123:GLU:HG3	4:K:239:ALA:HB2	1.97	0.46
4:K:193:ARG:NH1	4:M:638:LYS:HE3	2.29	0.46
4:K:336:MET:SD	4:K:366:ARG:HB3	2.55	0.46
4:K:402:ILE:HD13	4:K:469:LEU:HG	1.97	0.46
4:L:122:ARG:HE	4:M:153:ARG:HB2	1.79	0.46
4:L:244:ASN:HB3	4:L:247:ILE:HD12	1.98	0.46
4:L:284:LYS:HA	4:L:287:GLU:CG	2.42	0.46
4:L:487:ASP:HB2	4:L:489:TRP:CD1	2.51	0.46
4:M:155:ALA:HA	4:M:158:ASN:HD22	1.78	0.46

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
4:M:386:SER:OG	4:M:441:SER:O	2.34	0.46
4:M:578:GLU:C	4:M:584:ARG:HH12	2.19	0.46
5:O:131:ASP:N	5:O:131:ASP:OD1	2.47	0.46
5:O:319:PRO:HD2	5:O:322:LEU:HD23	1.96	0.46
5:O:550:PHE:HE2	5:O:557:GLY:HA3	1.80	0.46
5:O:1092:VAL:HG22	5:O:1093:PHE:H	1.80	0.46
3:P:136:ASP:C	3:P:140:LYS:HZ1	2.19	0.46
6:X:52:MET:SD	6:X:55:LEU:HD13	2.56	0.46
6:X:196:ARG:O	6:X:198:GLU:HG3	2.15	0.46
6:Y:10:GLN:HA	6:Y:13:ASP:OD1	2.15	0.46
6:Y:172:LEU:HG	6:Y:252:PHE:CE2	2.48	0.46
1:B:247:LEU:HG	1:B:521:ARG:HH21	1.79	0.46
1:B:318:ARG:HH11	1:B:370:LEU:HB3	1.79	0.46
1:B:426:CYS:HB3	1:B:1235:LEU:HB3	1.98	0.46
1:B:731:ASN:ND2	1:B:737:GLU:H	2.14	0.46
1:B:782:GLN:HA	1:B:785:TRP:HB2	1.96	0.46
1:B:1187:ASP:CG	1:B:1188:HIS:HD1	2.16	0.46
2:C:606:THR:HG1	2:C:640:THR:H	1.54	0.46
2:C:930:THR:O	2:C:933:THR:OG1	2.33	0.46
3:D:377:ARG:O	3:D:381:GLN:HG3	2.15	0.46
4:K:140:LEU:HD21	4:K:166:ILE:HG21	1.97	0.46
4:K:224:PRO:O	4:K:230:ARG:NE	2.48	0.46
4:K:572:LEU:HD22	4:K:592:LEU:HD21	1.97	0.46
4:L:389:GLU:HB3	4:L:392:TRP:HB3	1.97	0.46
5:O:37:ASN:HD21	5:O:40:ARG:HE	1.61	0.46
5:O:508:SER:OG	5:O:543:PRO:O	2.32	0.46
5:O:512:VAL:N	5:O:548:ARG:HH22	2.12	0.46
5:O:574:VAL:HB	5:O:612:PHE:CB	2.45	0.46
5:O:730:LEU:O	5:O:733:ASN:HB2	2.15	0.46
6:X:201:ALA:HA	6:X:218:TRP:HZ2	1.80	0.46
6:X:224:ASP:HB3	6:X:354:TYR:HB3	1.96	0.46
6:X:266:GLY:O	6:X:268:MET:HG2	2.15	0.46
6:Y:33:ASP:OD2	6:Y:35:THR:N	2.47	0.46
1:B:501:VAL:HG13	1:B:502:ASN:N	2.29	0.46
1:B:864:LEU:HD12	1:B:867:THR:HB	1.96	0.46
2:C:949:LEU:HD13	2:C:951:TRP:NE1	2.31	0.46
2:C:974:LYS:HB3	2:C:979:LEU:O	2.16	0.46
2:C:985:GLU:N	2:C:986:PRO:HD2	2.30	0.46
2:C:1180:PHE:CE2	2:C:1199:ILE:HG21	2.48	0.46
2:C:1212:THR:HB	2:C:1213:ASN:HD22	1.79	0.46
3:D:350:THR:HG22	3:D:353:GLN:NE2	2.21	0.46

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
4:K:42:ASN:ND2	4:M:116:MET:HB2	2.30	0.46
4:K:208:VAL:HA	4:K:662:GLN:HE22	1.81	0.46
4:K:223:LEU:O	4:M:589:LYS:NZ	2.48	0.46
4:K:398:LYS:O	4:K:473:ILE:N	2.31	0.46
4:K:435:SER:H	4:K:441:SER:HA	1.81	0.46
4:K:539:MET:HE1	4:K:611:ILE:HG13	1.98	0.46
4:L:515:PRO:O	4:L:519:ILE:HG12	2.15	0.46
4:M:548:GLU:O	4:M:551:LYS:HG3	2.15	0.46
4:M:609:THR:HA	4:M:612:THR:HG23	1.98	0.46
5:O:441:TYR:OH	5:O:443:ARG:HD2	2.16	0.46
3:P:8:PHE:CZ	3:P:125:TYR:HD1	2.34	0.46
3:P:228:ILE:HG21	3:P:234:TYR:HB2	1.97	0.46
6:X:53:HIS:CD2	6:X:75:GLN:HE22	2.32	0.46
6:X:171:ASN:HD22	6:X:174:ILE:HD11	1.79	0.46
6:Y:19:PHE:CD1	6:Y:291:LYS:HE3	2.49	0.46
6:Y:292:LEU:O	6:Y:295:VAL:HG22	2.15	0.46
6:Z:15:ILE:HG22	6:Z:19:PHE:HZ	1.79	0.46
6:Z:235:GLY:C	6:Z:239:ARG:HG3	2.36	0.46
6:Z:308:VAL:HA	6:Z:311:ILE:CD1	2.45	0.46
6:Z:320:MET:HG3	6:Z:323:TRP:CE3	2.51	0.46
1:B:577:ILE:HD13	1:B:627:TRP:HB3	1.97	0.46
1:B:622:SER:HB3	1:B:625:ASN:ND2	2.30	0.46
1:B:650:MET:HG3	1:B:654:ASN:HD21	1.81	0.46
1:B:657:VAL:C	1:B:659:PHE:H	2.19	0.46
1:B:1047:ILE:HG23	1:B:1137:ILE:HB	1.96	0.46
2:C:481:TRP:O	2:C:484:THR:OG1	2.30	0.46
2:C:512:ALA:O	2:C:516:ILE:HG23	2.16	0.46
2:C:552:ASP:OD2	2:C:555:ILE:HG23	2.15	0.46
2:C:681:PRO:HD2	2:C:1004:TYR:CE1	2.51	0.46
2:C:856:THR:HG23	3:P:349:MET:HA	1.97	0.46
3:D:19:VAL:HG21	3:D:73:SER:OG	2.15	0.46
3:D:140:LYS:O	3:D:144:TYR:HB2	2.16	0.46
3:D:233:THR:OG1	3:D:258:ASN:O	2.33	0.46
4:L:18:ASN:OD1	4:L:250:MET:N	2.48	0.46
4:L:332:THR:N	4:L:506:LYS:O	2.43	0.46
4:L:344:LEU:HD13	4:L:362:ARG:HB2	1.97	0.46
4:L:559:ASP:N	4:L:601:GLN:HE22	2.13	0.46
4:M:505:VAL:HG12	6:Y:312:ARG:NH1	2.31	0.46
5:O:121:GLY:HA3	5:O:156:PHE:CZ	2.49	0.46
5:O:580:GLN:HE22	5:O:618:PHE:HZ	1.64	0.46
5:O:1044:ILE:O	5:O:1086:GLU:HB3	2.15	0.46

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:P:39:LEU:O	3:P:43:LEU:HD13	2.15	0.46
6:X:211:ASP:CB	6:X:213:ARG:HE	2.28	0.46
6:Y:208:ARG:HH12	6:Y:286:ILE:HA	1.81	0.46
1:B:674:ARG:O	1:B:677:ALA:HB3	2.16	0.46
1:B:1135:LEU:HD12	1:B:1136:ARG:N	2.29	0.46
1:B:1246:GLN:HB3	1:B:1249:GLU:HB2	1.97	0.46
2:C:265:SER:OG	2:C:309:ARG:HG3	2.16	0.46
2:C:692:GLN:HA	2:C:703:ARG:NH2	2.30	0.46
2:C:755:LEU:HD23	2:C:755:LEU:HA	1.76	0.46
4:K:19:VAL:O	4:K:248:GLN:HB2	2.16	0.46
4:K:118:LEU:H	4:K:118:LEU:HD12	1.81	0.46
4:K:133:VAL:HG22	4:L:153:ARG:NH1	2.29	0.46
4:K:298:PRO:O	4:K:301:ILE:HG12	2.15	0.46
4:L:152:ALA:HA	4:L:155:ALA:HB3	1.98	0.46
4:L:508:ALA:HB3	6:Z:312:ARG:NH2	2.28	0.46
4:M:119:GLU:HA	4:M:122:ARG:HG3	1.97	0.46
4:M:327:LYS:NZ	4:M:328:ILE:HG13	2.29	0.46
5:O:13:LEU:CD1	5:O:358:PRO:HD2	2.44	0.46
5:O:47:ARG:HB2	5:O:52:ASN:OD1	2.16	0.46
5:O:907:PHE:O	5:O:911:ILE:HG12	2.15	0.46
5:O:1012:MET:HB3	5:O:1016:MET:CE	2.46	0.46
5:O:1073:ILE:HG23	5:O:1092:VAL:O	2.16	0.46
5:O:1131:PRO:HD2	5:O:1143:VAL:HG22	1.98	0.46
5:O:1155:THR:HG23	5:O:1198:ILE:HA	1.98	0.46
5:O:1193:TYR:CZ	5:O:1286:LEU:HD13	2.51	0.46
3:P:186:THR:HG22	3:P:188:ALA:H	1.79	0.46
6:Y:128:LEU:HD23	6:Y:128:LEU:H	1.81	0.46
6:Y:336:THR:HB	6:Y:340:LEU:HD12	1.98	0.46
6:Z:328:MET:SD	6:Z:329:GLN:HG3	2.55	0.46
1:B:440:MET:H	2:C:862:LEU:N	2.11	0.46
1:B:474:ILE:HD11	1:B:507:TYR:HB2	1.97	0.46
1:B:521:ARG:NH2	1:B:828:PHE:HA	2.31	0.46
1:B:868:ASN:OD1	1:B:868:ASN:N	2.49	0.46
1:B:920:ARG:NH2	1:B:983:LEU:HD13	2.30	0.46
1:B:946:ASP:HB3	3:P:31:ARG:NH2	2.30	0.46
1:B:1104:PHE:HD1	1:B:1129:TRP:CD1	2.33	0.46
2:C:798:ASP:O	2:C:801:LYS:HG2	2.16	0.46
2:C:1178:VAL:HG22	2:C:1179:PRO:O	2.16	0.46
3:D:217:LEU:HD21	3:D:291:LEU:HD11	1.97	0.46
4:K:48:TRP:CG	4:K:65:ARG:HA	2.51	0.46
4:K:334:LEU:O	4:K:335:ARG:NE	2.49	0.46

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
4:K:632:LYS:CE	4:L:202:ASN:HD22	2.29	0.46
4:L:245:GLY:O	4:L:248:GLN:HG3	2.15	0.46
4:M:337:ILE:HG22	4:M:363:GLY:HA2	1.97	0.46
4:M:385:LYS:HD2	4:M:385:LYS:N	2.31	0.46
4:M:660:TRP:CE2	4:M:664:PHE:CZ	3.04	0.46
5:O:240:GLN:CD	5:O:248:ILE:HB	2.36	0.46
5:O:617:ASN:HB3	5:O:618:PHE:CE1	2.51	0.46
5:O:899:LYS:HB3	5:O:901:MET:SD	2.56	0.46
5:O:950:ARG:HD3	5:O:952:PRO:HD3	1.97	0.46
5:O:1194:LEU:HB2	5:O:1196:TYR:CE1	2.50	0.46
5:O:1227:ASP:OD2	5:O:1280:PRO:HA	2.16	0.46
3:P:47:SER:C	3:P:49:GLY:H	2.18	0.46
3:P:315:ASN:HD22	3:P:322:ARG:NE	2.14	0.46
3:P:344:ARG:HG2	3:P:354:GLN:NE2	2.30	0.46
3:P:354:GLN:HG2	3:P:358:GLU:OE2	2.16	0.46
6:Y:195:VAL:HG23	6:Y:197:LEU:HD21	1.96	0.46
6:Z:126:GLY:O	6:Z:364:LEU:HA	2.15	0.46
1:B:251:ASP:CB	1:B:979:LEU:HD22	2.45	0.46
1:B:414:PRO:HG2	2:C:1122:PHE:HE2	1.80	0.46
1:B:1125:GLN:HB2	1:B:1126:PHE:CD2	2.51	0.46
1:B:1149:HIS:N	1:B:1180:PHE:O	2.48	0.46
2:C:510:SER:OG	2:C:513:GLU:OE2	2.25	0.46
2:C:1082:ARG:O	2:C:1094:ILE:HG13	2.16	0.46
2:C:1108:TRP:HB2	2:C:1137:ILE:HG23	1.98	0.46
3:D:173:GLY:O	3:D:176:ASN:N	2.49	0.46
3:D:263:ILE:HG12	3:D:268:ARG:CZ	2.46	0.46
4:K:191:ASP:OD2	4:K:192:LEU:N	2.49	0.46
4:K:372:GLN:NE2	4:K:509:VAL:HG23	2.31	0.46
4:K:401:PHE:CD2	4:K:470:ALA:HB2	2.51	0.46
4:L:126:ASP:O	4:L:129:ARG:N	2.48	0.46
4:L:131:LEU:HG	4:L:133:VAL:O	2.16	0.46
4:L:535:ALA:O	4:L:539:MET:HG2	2.16	0.46
4:M:239:ALA:O	4:M:243:ARG:HG2	2.16	0.46
4:M:243:ARG:C	4:M:261:ALA:HB2	2.36	0.46
5:O:412:LEU:O	5:O:415:SER:OG	2.19	0.46
5:O:530:ILE:O	5:O:533:TRP:HB3	2.16	0.46
5:O:1035:ILE:HG23	5:O:1120:ASP:HB2	1.98	0.46
5:O:1053:PHE:HD2	5:O:1103:ALA:HA	1.80	0.46
3:P:333:ARG:NH1	3:P:337:THR:HG1	2.09	0.46
6:X:92:THR:HG22	6:X:96:LYS:HD2	1.97	0.46
6:Z:312:ARG:O	6:Z:316:GLY:HA2	2.16	0.46

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:B:368:ASP:N	1:B:368:ASP:OD1	2.48	0.46
1:B:556:ILE:O	1:B:559:THR:OG1	2.33	0.46
1:B:947:ARG:CZ	1:B:956:ARG:HH22	2.29	0.46
2:C:324:MET:HA	2:C:334:LEU:CD1	2.45	0.46
2:C:419:ILE:O	2:C:423:VAL:HG23	2.16	0.46
2:C:421:ASN:HD21	2:C:1217:PRO:CA	2.26	0.46
2:C:623:LEU:HD21	2:C:785:TRP:CH2	2.51	0.46
2:C:637:PRO:HA	2:C:713:TRP:CH2	2.50	0.46
2:C:677:ALA:HB3	2:C:678:PHE:CE1	2.51	0.46
2:C:754:THR:CG2	2:C:804:LYS:HB3	2.46	0.46
2:C:1093:MET:HB3	2:C:1101:MET:HB3	1.98	0.46
2:C:1166:TRP:CD1	2:C:1179:PRO:HD3	2.50	0.46
4:L:224:PRO:O	4:L:230:ARG:NH2	2.48	0.46
4:L:559:ASP:H	4:L:601:GLN:NE2	2.14	0.46
4:M:375:PRO:HD2	4:M:499:SER:OG	2.16	0.46
4:M:382:LEU:HG	4:M:443:ILE:HG21	1.98	0.46
5:O:371:ARG:HG3	5:O:372:LYS:N	2.31	0.46
5:O:436:SER:HG	5:O:438:TYR:HD2	1.62	0.46
3:P:161:ILE:CA	3:P:272:VAL:O	2.50	0.46
3:P:185:HIS:C	3:P:252:ILE:HG13	2.36	0.46
3:P:185:HIS:O	3:P:252:ILE:HG13	2.16	0.46
3:P:189:GLU:HG2	3:P:190:ILE:HD12	1.98	0.46
6:Y:219:GLY:HA3	6:Y:358:ILE:HD13	1.98	0.46
6:Y:293:LYS:HA	6:Y:296:ARG:HB2	1.97	0.46
6:Z:134:LEU:H	6:Z:134:LEU:HG	1.59	0.46
1:B:364:GLU:O	1:B:367:LEU:HB3	2.16	0.45
1:B:370:LEU:HD11	1:B:465:TRP:CD2	2.52	0.45
1:B:413:THR:O	2:C:1082:ARG:HD2	2.16	0.45
1:B:1003:GLN:HE21	1:B:1004:TYR:HB3	1.80	0.45
1:B:1047:ILE:CD1	1:B:1198:TYR:HB3	2.46	0.45
1:B:1113:ALA:HA	1:B:1116:GLN:CD	2.36	0.45
2:C:347:ARG:HD2	2:C:347:ARG:HA	1.81	0.45
2:C:451:PHE:CD1	2:C:1256:VAL:HB	2.51	0.45
2:C:600:TYR:O	2:C:603:VAL:HG22	2.16	0.45
2:C:1096:ASP:OD1	2:C:1096:ASP:N	2.48	0.45
3:D:9:LYS:HG2	3:D:124:VAL:HG12	1.98	0.45
3:D:283:VAL:HG22	3:D:287:LEU:HG	1.97	0.45
3:D:365:GLN:O	3:D:369:ARG:HG2	2.16	0.45
3:D:395:LYS:HD2	3:D:396:PRO:HD2	1.99	0.45
4:K:445:THR:HG23	4:L:325:THR:O	2.16	0.45
4:K:483:MET:HE2	4:K:485:GLN:H	1.81	0.45

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
4:L:193:ARG:HH21	4:L:197:THR:HA	1.81	0.45
4:L:297:GLU:HB3	4:M:650:LYS:HZ1	1.81	0.45
4:L:324:ARG:NH2	4:L:489:TRP:N	2.62	0.45
4:M:137:TYR:HB2	4:M:174:TYR:CE2	2.51	0.45
4:M:662:GLN:NE2	4:M:665:LEU:HD12	2.31	0.45
5:O:27:LEU:CB	5:O:108:ASN:HA	2.45	0.45
5:O:38:PRO:HG3	5:O:87:ARG:HH12	1.80	0.45
5:O:414:VAL:HA	5:O:417:LEU:HD12	1.97	0.45
5:O:723:ALA:O	5:O:727:ILE:HG22	2.16	0.45
5:O:765:ARG:HB2	5:O:838:GLU:OE2	2.16	0.45
5:O:1053:PHE:O	5:O:1065:THR:N	2.48	0.45
5:O:1133:GLN:HE21	5:O:1134:LEU:C	2.20	0.45
6:Y:171:ASN:HB2	6:Y:252:PHE:CE2	2.51	0.45
6:Z:120:ARG:NH2	6:Z:124:GLU:OE1	2.50	0.45
6:Z:195:VAL:HG13	6:Z:197:LEU:HG	1.98	0.45
1:B:608:ILE:HG23	1:B:877:LEU:O	2.16	0.45
1:B:737:GLU:HB3	5:O:213:THR:HG23	1.97	0.45
1:B:920:ARG:NH1	1:B:924:THR:OG1	2.49	0.45
2:C:268:ILE:HG12	2:C:1210:PHE:CG	2.51	0.45
2:C:740:LEU:HD13	2:C:835:TYR:CG	2.52	0.45
2:C:754:THR:HG23	2:C:804:LYS:HB3	1.97	0.45
2:C:1197:GLN:CD	2:C:1198:TYR:H	2.19	0.45
3:D:31:ARG:HD2	3:D:45:TRP:CZ2	2.49	0.45
3:D:66:TYR:HB3	3:D:393:ARG:O	2.16	0.45
4:K:282:LYS:O	4:K:285:LEU:HG	2.16	0.45
4:K:654:SER:O	4:K:658:GLN:HG3	2.16	0.45
4:L:636:ARG:HH12	4:M:27:SER:HB3	1.80	0.45
4:M:393:ASP:OD2	4:M:395:ASN:HB3	2.16	0.45
4:M:395:ASN:ND2	4:M:432:ALA:O	2.49	0.45
4:M:404:PHE:O	4:M:466:TYR:HA	2.16	0.45
5:O:160:HIS:O	5:O:163:VAL:HB	2.17	0.45
5:O:324:ASN:N	5:O:324:ASN:OD1	2.49	0.45
5:O:888:CYS:SG	5:O:924:VAL:HG23	2.56	0.45
5:O:967:GLU:HA	5:O:970:CYS:SG	2.56	0.45
3:P:380:ASP:O	3:P:384:GLN:HG2	2.17	0.45
6:X:167:TRP:CD1	6:X:280:CYS:HB2	2.51	0.45
6:X:218:TRP:HD1	6:X:219:GLY:O	1.98	0.45
6:X:347:THR:HG22	6:X:350:GLY:H	1.81	0.45
6:Y:95:TRP:HD1	6:Y:142:PHE:CE2	2.34	0.45
6:Z:307:GLY:H	6:Z:310:LYS:HE2	1.81	0.45
1:B:451:PHE:CD1	1:B:1256:VAL:HG22	2.52	0.45

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:B:504:LEU:HD21	1:B:1262:TYR:CD1	2.51	0.45
1:B:680:THR:HG21	1:B:682:HIS:NE2	2.30	0.45
1:B:690:ASN:CG	1:B:693:LEU:HG	2.37	0.45
2:C:685:PRO:O	2:C:688:PHE:N	2.46	0.45
2:C:851:ARG:HB2	2:C:852:GLN:OE1	2.15	0.45
2:C:1084:SER:O	2:C:1092:PRO:HA	2.16	0.45
3:D:285:LEU:H	3:D:285:LEU:HD12	1.82	0.45
4:K:118:LEU:HA	4:K:121:ASN:HD22	1.81	0.45
4:K:304:LEU:HD11	4:K:621:LYS:HZ3	1.79	0.45
4:L:193:ARG:O	4:L:197:THR:HG23	2.15	0.45
4:L:212:MET:HA	4:L:215:VAL:CG2	2.46	0.45
4:L:223:LEU:HG	4:L:224:PRO:O	2.15	0.45
4:L:435:SER:HA	4:L:442:ILE:CG2	2.45	0.45
4:L:640:LEU:HD23	4:M:182:LEU:HD11	1.97	0.45
4:M:33:SER:OG	4:M:34:LEU:N	2.48	0.45
4:M:127:LYS:HA	4:M:127:LYS:HD3	1.55	0.45
5:O:19:GLU:HB3	5:O:277:THR:O	2.16	0.45
5:O:221:LEU:HD13	5:O:285:LEU:HA	1.98	0.45
5:O:295:ASP:OD1	5:O:295:ASP:N	2.47	0.45
5:O:439:VAL:O	5:O:639:MET:HG3	2.16	0.45
5:O:583:ASP:C	5:O:587:ASP:HB2	2.37	0.45
5:O:891:SER:H	5:O:925:GLN:HE22	1.64	0.45
5:O:1084:LYS:O	5:O:1086:GLU:HG3	2.16	0.45
3:P:167:ASP:N	3:P:170:MET:HE3	2.31	0.45
3:P:176:ASN:HA	3:P:179:LEU:HB3	1.99	0.45
3:P:182:TYR:HA	3:P:185:HIS:CD2	2.51	0.45
3:P:225:VAL:C	3:P:339:PHE:HD1	2.19	0.45
6:Y:99:MET:N	6:Y:99:MET:SD	2.89	0.45
6:Z:86:GLN:H	6:Z:86:GLN:CD	2.18	0.45
1:B:474:ILE:HG22	1:B:475:ASN:N	2.31	0.45
1:B:731:ASN:OD1	1:B:736:PRO:HA	2.16	0.45
1:B:983:LEU:HA	1:B:983:LEU:HD12	1.74	0.45
2:C:308:THR:HG21	2:C:368:ASP:OD2	2.17	0.45
2:C:402:TRP:O	2:C:406:MET:HG2	2.16	0.45
2:C:537:GLN:O	2:C:541:VAL:HG23	2.16	0.45
2:C:556:ILE:O	2:C:560:MET:HG2	2.16	0.45
2:C:713:TRP:HA	2:C:713:TRP:CE3	2.50	0.45
2:C:974:LYS:O	2:C:978:ASP:N	2.50	0.45
4:K:123:GLU:HB3	4:K:188:TRP:CH2	2.52	0.45
4:K:147:TYR:HB3	4:K:155:ALA:O	2.17	0.45
4:K:272:PRO:HG3	4:K:277:PRO:HA	1.98	0.45

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
4:K:334:LEU:HD23	4:K:368:VAL:HA	1.99	0.45
4:K:600:MET:O	4:K:604:ILE:HG12	2.16	0.45
4:L:133:VAL:HG22	4:L:134:SER:N	2.31	0.45
4:L:476:ALA:HB2	6:Y:160:ASP:HB2	1.98	0.45
4:M:223:LEU:HA	4:M:223:LEU:HD23	1.71	0.45
4:M:571:GLN:NE2	4:M:574:ILE:HD11	2.31	0.45
5:O:248:ILE:C	5:O:249:ILE:HG13	2.37	0.45
5:O:404:GLN:O	5:O:774:LEU:HD11	2.17	0.45
5:O:511:SER:HG	5:O:572:GLN:HE21	1.63	0.45
5:O:617:ASN:CA	5:O:651:LEU:HD13	2.44	0.45
5:O:658:VAL:HG12	5:O:659:HIS:CD2	2.50	0.45
5:O:712:GLU:OE1	5:O:713:ASN:ND2	2.49	0.45
5:O:985:SER:O	5:O:985:SER:OG	2.28	0.45
5:O:1035:ILE:HG12	5:O:1120:ASP:OD2	2.17	0.45
5:O:1159:ILE:HD12	5:O:1164:GLN:HB2	1.98	0.45
3:P:218:SER:O	3:P:222:LEU:HG	2.16	0.45
6:Z:95:TRP:HZ3	6:Z:282:MET:HE1	1.81	0.45
1:B:259:ASP:N	1:B:372:LEU:O	2.48	0.45
1:B:299:ALA:HB3	1:B:302:SER:OG	2.16	0.45
1:B:685:PRO:HD2	1:B:688:PHE:HB2	1.98	0.45
1:B:764:TRP:HE3	1:B:767:ARG:NH1	2.14	0.45
1:B:848:GLY:O	1:B:864:LEU:HD21	2.17	0.45
2:C:265:SER:HB3	2:C:308:THR:HA	1.98	0.45
2:C:737:GLU:OE1	2:C:737:GLU:N	2.36	0.45
2:C:856:THR:HG22	2:C:857:ILE:N	2.32	0.45
2:C:1052:SER:OG	2:C:1053:THR:N	2.49	0.45
3:D:95:VAL:HG22	3:D:105:VAL:O	2.16	0.45
4:K:377:ARG:NH2	4:L:524:PRO:HG3	2.32	0.45
4:K:578:GLU:N	4:K:584:ARG:HH22	2.14	0.45
4:L:124:PHE:CD2	4:L:125:LEU:HD23	2.52	0.45
4:L:301:ILE:HG23	4:L:624:TRP:HZ2	1.81	0.45
4:L:639:SER:HB3	4:M:120:PHE:HE2	1.82	0.45
4:M:160:GLN:HG3	4:M:161:LYS:H	1.80	0.45
4:M:548:GLU:HG2	4:M:551:LYS:HE3	1.98	0.45
5:O:93:LEU:HA	5:O:96:GLU:CD	2.37	0.45
5:O:142:MET:HA	5:O:145:ASP:HB2	1.98	0.45
5:O:224:TYR:HD2	5:O:369:TYR:OH	1.95	0.45
5:O:403:ASP:CG	5:O:404:GLN:H	2.18	0.45
5:O:802:SER:O	5:O:805:VAL:HG22	2.16	0.45
5:O:1048:ASN:HA	5:O:1080:PHE:CZ	2.52	0.45
5:O:1056:TYR:HA	5:O:1062:ALA:H	1.82	0.45

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:P:20:PRO:HB2	3:P:391:MET:HA	1.99	0.45
3:P:132:PHE:HA	3:P:135:ARG:HH11	1.82	0.45
3:P:136:ASP:O	3:P:140:LYS:HG3	2.15	0.45
3:P:333:ARG:HH11	3:P:333:ARG:C	2.20	0.45
3:P:334:ARG:CZ	3:P:337:THR:HG21	2.46	0.45
3:P:334:ARG:HA	3:P:337:THR:OG1	2.16	0.45
3:P:355:ASN:HA	3:P:358:GLU:OE2	2.17	0.45
6:X:1:MET:O	6:X:3:VAL:HG23	2.16	0.45
6:X:258:SER:OG	6:X:259:ARG:N	2.49	0.45
6:Y:87:PHE:CZ	6:Y:212:SER:HB3	2.51	0.45
1:B:245:ALA:HA	1:B:248:LEU:HD12	1.98	0.45
1:B:313:ASN:OD1	1:B:314:VAL:N	2.50	0.45
1:B:770:GLU:HA	1:B:773:LYS:CD	2.41	0.45
1:B:981:ASP:OD2	1:B:983:LEU:HD13	2.17	0.45
2:C:905:ILE:HG13	2:C:909:TYR:CE2	2.51	0.45
2:C:1157:ALA:HB3	2:C:1195:ALA:HB1	1.98	0.45
3:D:136:ASP:OD2	3:D:139:PHE:N	2.45	0.45
3:D:182:TYR:HA	3:D:185:HIS:ND1	2.32	0.45
4:K:151:SER:OG	4:K:152:ALA:N	2.50	0.45
4:L:273:SER:O	4:L:273:SER:OG	2.32	0.45
4:L:410:PHE:HA	4:L:413:TRP:CD1	2.52	0.45
4:M:252:VAL:HG23	4:M:253:SER:H	1.80	0.45
5:O:219:GLY:O	5:O:235:LEU:HD12	2.15	0.45
5:O:349:ARG:CZ	3:P:129:ASP:HA	2.47	0.45
5:O:481:ARG:CZ	5:O:482:SER:HG	2.29	0.45
5:O:498:ASP:OD1	5:O:499:PRO:HD2	2.16	0.45
5:O:836:ILE:HD12	5:O:839:LEU:HB2	1.98	0.45
5:O:965:ALA:O	5:O:969:ILE:HG13	2.16	0.45
3:P:76:LEU:O	3:P:79:PRO:HD2	2.17	0.45
3:P:78:ILE:HB	3:P:79:PRO:HD3	1.99	0.45
3:P:186:THR:N	3:P:189:GLU:OE2	2.50	0.45
3:P:331:GLN:HE22	3:P:403:GLY:H	1.65	0.45
1:B:375:HIS:HA	1:B:1259:ARG:HD2	1.99	0.45
1:B:378:PHE:HB3	1:B:390:ASN:HB3	1.98	0.45
1:B:636:LEU:HA	1:B:636:LEU:HD23	1.59	0.45
1:B:681:PRO:HA	1:B:684:TRP:CE3	2.52	0.45
1:B:859:GLN:OE1	3:P:97:SER:HA	2.17	0.45
1:B:1144:TYR:CE2	1:B:1146:TYR:HB3	2.45	0.45
2:C:268:ILE:HG12	2:C:1210:PHE:CD1	2.52	0.45
2:C:471:ARG:NH1	3:D:257:LEU:HD21	2.32	0.45
2:C:713:TRP:CE3	2:C:714:PRO:HD2	2.52	0.45

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:C:818:GLU:HA	2:C:821:VAL:HG22	1.97	0.45
2:C:972:SER:HB2	2:C:973:MET:HE2	1.97	0.45
2:C:974:LYS:HG2	2:C:979:LEU:HB3	1.98	0.45
2:C:1045:ILE:O	2:C:1139:ILE:HG22	2.16	0.45
3:D:189:GLU:O	3:D:192:TYR:HB2	2.17	0.45
3:D:244:ARG:O	3:D:357:ILE:HD12	2.16	0.45
4:K:122:ARG:CZ	4:L:153:ARG:HB2	2.46	0.45
4:K:126:ASP:OD1	4:K:127:LYS:N	2.50	0.45
4:K:151:SER:HB3	4:K:154:GLN:HE22	1.82	0.45
4:L:28:SER:HA	4:L:263:ASN:ND2	2.31	0.45
4:L:280:GLU:HA	4:M:36:LEU:HD13	1.98	0.45
4:L:668:VAL:O	4:L:671:HIS:HB3	2.16	0.45
4:M:232:TYR:O	4:M:235:GLU:HB2	2.17	0.45
4:M:284:LYS:HA	4:M:287:GLU:OE1	2.16	0.45
4:M:591:ILE:HA	4:M:594:LYS:HE3	1.98	0.45
5:O:14:SER:HB3	5:O:313:GLN:OE1	2.16	0.45
5:O:511:SER:H	5:O:572:GLN:NE2	2.15	0.45
5:O:638:TYR:CD2	5:O:664:LEU:HD13	2.51	0.45
5:O:770:ARG:HG2	5:O:860:CYS:C	2.37	0.45
5:O:977:CYS:N	5:O:1025:GLY:O	2.50	0.45
5:O:1190:ALA:HB2	5:O:1231:SER:O	2.17	0.45
3:P:233:THR:OG1	3:P:260:SER:HA	2.17	0.45
6:X:184:MET:HE1	6:X:243:VAL:HG12	1.98	0.45
6:Y:122:ARG:CZ	6:Y:236:ARG:HH21	2.29	0.45
6:Y:192:PHE:HD1	6:Y:206:PHE:CE2	2.34	0.45
6:Z:78:ARG:HG3	6:Z:80:GLN:HB3	1.98	0.45
6:Z:168:THR:HG22	6:Z:172:LEU:HD11	1.98	0.45
6:Z:289:THR:O	6:Z:292:LEU:HB3	2.17	0.45
1:B:437:ALA:HB2	1:B:447:TRP:CE2	2.52	0.45
1:B:668:ILE:HG12	1:B:669:TYR:CD1	2.52	0.45
1:B:970:ASN:O	1:B:974:LYS:HE3	2.17	0.45
1:B:1056:TRP:HZ3	1:B:1060:ALA:O	1.99	0.45
2:C:714:PRO:HA	2:C:760:GLU:CD	2.36	0.45
2:C:1155:GLN:NE2	2:C:1156:TYR:HB3	2.32	0.45
3:D:263:ILE:HG13	3:D:264:ARG:N	2.31	0.45
3:D:375:TRP:HA	3:D:378:GLU:OE1	2.16	0.45
4:K:333:TRP:CE3	4:K:334:LEU:HD11	2.52	0.45
4:L:39:GLY:O	4:L:42:ASN:HB2	2.16	0.45
4:L:212:MET:HA	4:L:215:VAL:HB	1.98	0.45
4:L:323:VAL:HA	4:L:326:LEU:CD1	2.47	0.45
4:L:395:ASN:OD1	4:L:432:ALA:N	2.38	0.45

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
4:M:524:PRO:HB3	4:M:612:THR:HB	1.98	0.45
5:O:69:ASP:OD2	5:O:71:GLN:HG2	2.17	0.45
5:O:172:TYR:CD2	5:O:190:LYS:HB2	2.43	0.45
5:O:327:LEU:O	5:O:330:PHE:HB3	2.17	0.45
5:O:362:ASP:OD1	5:O:363:GLY:N	2.49	0.45
5:O:497:ILE:HA	5:O:504:GLU:HA	1.99	0.45
5:O:949:TYR:HB2	5:O:958:GLU:HG2	1.99	0.45
5:O:984:LEU:HD23	5:O:984:LEU:HA	1.84	0.45
5:O:1073:ILE:HD13	5:O:1073:ILE:HA	1.89	0.45
3:P:316:ILE:HB	3:P:320:THR:H	1.81	0.45
3:P:376:ALA:HB3	3:P:377:ARG:NH1	2.32	0.45
6:X:110:MET:HG2	6:X:173:MET:SD	2.57	0.45
6:Y:158:ASP:OD2	6:Y:161:THR:HB	2.17	0.45
6:Y:171:ASN:HB2	6:Y:252:PHE:CZ	2.51	0.45
6:Y:241:GLU:HB3	6:Y:242:LEU:HG	1.99	0.45
6:Z:36:ILE:HA	6:Z:152:PRO:HG3	1.98	0.45
6:Z:188:PRO:HB2	6:Z:285:PHE:CE2	2.51	0.45
1:B:408:PRO:O	1:B:411:MET:HB3	2.16	0.45
1:B:615:LYS:NZ	1:B:674:ARG:HB3	2.32	0.45
1:B:692:GLN:NE2	1:B:692:GLN:H	2.14	0.45
1:B:764:TRP:O	1:B:768:VAL:HG22	2.17	0.45
1:B:1014:GLU:HG3	3:P:34:ASN:OD1	2.17	0.45
1:B:1109:ILE:HG23	1:B:1138:ARG:NE	2.32	0.45
1:B:1115:TRP:CD2	1:B:1122:PHE:HB3	2.52	0.45
2:C:464:ARG:NH2	2:C:1026:LEU:HD13	2.32	0.45
2:C:979:LEU:HG	2:C:980:SER:N	2.31	0.45
3:D:47:SER:HA	3:D:192:TYR:OH	2.16	0.45
3:D:164:VAL:HB	3:D:270:LEU:HG	1.97	0.45
3:D:179:LEU:HD23	3:D:182:TYR:CD1	2.52	0.45
3:D:314:ARG:HA	3:D:320:THR:CG2	2.44	0.45
3:D:366:GLN:HA	3:D:369:ARG:HG2	1.98	0.45
3:D:404:ARG:NH2	3:D:406:ASN:HA	2.32	0.45
4:K:45:GLY:H	4:K:102:VAL:HB	1.82	0.45
4:K:243:ARG:NH1	4:K:262:VAL:HG22	2.31	0.45
4:K:398:LYS:HG2	4:K:427:TYR:CE2	2.51	0.45
4:K:490:ASP:OD2	4:M:436:SER:HB3	2.17	0.45
4:K:627:LEU:O	4:K:631:VAL:HG12	2.17	0.45
4:L:665:LEU:HD13	4:L:665:LEU:HA	1.79	0.45
4:M:121:ASN:OD1	4:M:188:TRP:NE1	2.49	0.45
4:M:323:VAL:O	4:M:324:ARG:HD2	2.17	0.45
4:M:396:GLY:HA2	4:M:430:LEU:O	2.17	0.45

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
5:O:154:SER:OG	5:O:155:GLN:OE1	2.22	0.45
5:O:456:LEU:HD21	5:O:460:TYR:HB3	1.97	0.45
5:O:515:PHE:CE1	5:O:550:PHE:HB2	2.52	0.45
5:O:606:THR:HG22	5:O:607:ALA:H	1.81	0.45
5:O:803:PRO:HA	5:O:806:CYS:SG	2.57	0.45
5:O:931:ASP:HB3	5:O:1015:TYR:OH	2.16	0.45
5:O:1011:LEU:HD22	5:O:1208:PHE:CZ	2.52	0.45
5:O:1049:ALA:HB2	5:O:1080:PHE:CD2	2.52	0.45
5:O:1236:ARG:HA	5:O:1236:ARG:NE	2.31	0.45
3:P:168:GLU:O	3:P:170:MET:N	2.50	0.45
6:X:7:ASN:CB	6:X:10:GLN:HG2	2.46	0.45
6:X:18:ALA:HA	6:X:23:VAL:HG23	1.98	0.45
6:Y:23:VAL:HG12	6:Y:24:SER:H	1.81	0.45
6:Z:68:LEU:HD12	6:Z:69:PRO:HD2	1.98	0.45
6:Z:290:ALA:HA	6:Z:293:LYS:NZ	2.32	0.45
1:B:380:GLN:NE2	2:C:958:SER:H	2.09	0.45
1:B:701:ILE:H	1:B:701:ILE:HG13	1.51	0.45
1:B:811:LEU:O	1:B:816:PRO:HD3	2.17	0.45
1:B:1054:HIS:CD2	1:B:1056:TRP:HB3	2.52	0.45
2:C:647:LYS:HB3	2:C:675:ALA:HB1	1.99	0.45
2:C:969:VAL:HG13	2:C:973:MET:HE3	1.99	0.45
4:K:47:PRO:HG2	4:K:100:LEU:HD11	1.99	0.45
4:K:145:ASP:CG	4:K:162:GLN:HE21	2.18	0.45
4:L:68:SER:N	4:L:100:LEU:HD21	2.32	0.45
4:L:375:PRO:HA	4:L:452:TYR:O	2.17	0.45
4:L:527:LEU:HA	4:L:527:LEU:HD12	1.77	0.45
5:O:114:PHE:CZ	5:O:229:ASN:HA	2.52	0.45
5:O:344:LEU:HD12	5:O:344:LEU:O	2.17	0.45
5:O:768:ARG:O	5:O:861:TRP:HE3	1.99	0.45
5:O:879:THR:HG23	5:O:916:LYS:HB3	1.99	0.45
5:O:1054:ASN:HB2	5:O:1056:TYR:CE1	2.51	0.45
3:P:36:PRO:O	3:P:72:LEU:HD21	2.17	0.45
6:X:36:ILE:HD11	6:X:153:LEU:H	1.82	0.45
6:Y:19:PHE:CZ	6:Y:291:LYS:HG2	2.51	0.45
6:Y:198:GLU:HA	6:Y:202:ARG:NE	2.32	0.45
6:Y:198:GLU:N	6:Y:201:ALA:HB3	2.32	0.45
6:Y:272:PHE:HB3	6:Y:280:CYS:SG	2.56	0.45
6:Z:297:LYS:O	6:Z:301:SER:OG	2.27	0.45
1:B:269:VAL:HB	1:B:271:ILE:HD11	1.99	0.44
1:B:417:SER:H	1:B:420:CYS:HB2	1.81	0.44
1:B:505:MET:HG3	1:B:505:MET:O	2.17	0.44

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:B:608:ILE:HG22	1:B:609:ASP:N	2.32	0.44
1:B:928:VAL:O	1:B:932:VAL:HG22	2.17	0.44
2:C:325:ALA:HB3	2:C:331:ASN:ND2	2.29	0.44
2:C:577:ILE:HB	2:C:627:TRP:CD1	2.51	0.44
2:C:743:ILE:HG22	2:C:744:ASP:H	1.82	0.44
2:C:1167:LEU:HD23	2:C:1167:LEU:HA	1.81	0.44
3:D:222:LEU:CB	3:D:228:ILE:HG13	2.47	0.44
3:D:399:ASN:HA	3:D:402:TRP:CD1	2.51	0.44
4:K:201:THR:HG23	4:K:202:ASN:OD1	2.17	0.44
4:K:298:PRO:HG2	4:K:299:GLU:OE2	2.17	0.44
4:K:401:PHE:HA	4:K:469:LEU:O	2.17	0.44
4:K:453:GLU:H	4:K:456:GLN:HG3	1.82	0.44
4:L:20:PHE:O	4:L:210:CYS:N	2.43	0.44
4:L:166:ILE:HA	4:L:170:ARG:HH11	1.81	0.44
4:L:195:ALA:O	4:L:198:LEU:HD22	2.17	0.44
4:L:299:GLU:HB2	4:M:650:LYS:CD	2.48	0.44
4:M:234:LYS:HB2	4:M:234:LYS:HE2	1.74	0.44
4:M:395:ASN:OD1	4:M:396:GLY:N	2.51	0.44
4:M:405:GLN:O	4:M:420:GLY:HA3	2.18	0.44
4:M:426:ASN:HD21	4:M:449:ALA:H	1.62	0.44
5:O:189:ALA:O	5:O:192:LEU:HG	2.17	0.44
5:O:448:VAL:HG12	5:O:666:TRP:HB2	1.99	0.44
5:O:654:VAL:HG22	5:O:656:PHE:HE1	1.82	0.44
5:O:990:TRP:CZ2	5:O:991:THR:HG23	2.52	0.44
5:O:1008:ILE:O	5:O:1012:MET:HG2	2.17	0.44
5:O:1193:TYR:CE1	5:O:1286:LEU:HD13	2.52	0.44
5:O:1246:THR:OG1	5:O:1247:ILE:N	2.50	0.44
3:P:42:PHE:HE1	3:P:64:ARG:HB3	1.82	0.44
3:P:242:ASN:HA	3:P:244:ARG:NH2	2.31	0.44
6:X:220:VAL:O	6:X:358:ILE:HA	2.17	0.44
6:X:299:VAL:HG13	6:X:320:MET:CG	2.43	0.44
6:Y:23:VAL:HG12	6:Y:24:SER:N	2.31	0.44
6:Y:34:LYS:O	6:Y:153:LEU:HD12	2.17	0.44
6:Z:96:LYS:NZ	6:Z:284:PRO:HD3	2.31	0.44
6:Z:162:LYS:HG2	6:Z:166:TYR:CE2	2.52	0.44
1:B:300:ALA:O	1:B:1212:THR:OG1	2.35	0.44
1:B:949:LEU:HB3	1:B:952:ILE:HD12	1.99	0.44
1:B:1036:ASN:HD22	1:B:1205:ASN:CA	2.30	0.44
2:C:474:ILE:HG22	2:C:475:ASN:N	2.32	0.44
2:C:733:PHE:CE1	2:C:1001:TYR:HB3	2.52	0.44
3:D:227:VAL:HB	3:D:418:ILE:HG13	2.00	0.44

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:D:332:VAL:O	3:D:335:MET:HB3	2.17	0.44
4:K:28:SER:HA	4:K:243:ARG:HG3	1.99	0.44
4:K:376:MET:SD	4:K:495:MET:HB3	2.58	0.44
4:K:469:LEU:HD12	4:K:489:TRP:HE1	1.81	0.44
4:L:213:GLN:HB3	4:L:675:PRO:HD2	1.99	0.44
4:L:583:VAL:O	6:Z:70:HIS:N	2.49	0.44
4:M:328:ILE:HG13	4:M:329:ASP:H	1.81	0.44
4:M:471:THR:HG23	4:M:488:VAL:HG23	1.99	0.44
5:O:178:TYR:HE1	5:O:184:ASP:HB3	1.81	0.44
5:O:429:ILE:HB	5:O:475:ALA:HB1	1.98	0.44
5:O:632:LEU:HD22	5:O:664:LEU:HB2	1.98	0.44
5:O:832:PRO:HB3	5:O:853:PRO:O	2.17	0.44
5:O:936:ILE:HA	5:O:937:LYS:HZ2	1.83	0.44
6:X:160:ASP:N	6:X:160:ASP:OD1	2.48	0.44
6:X:227:GLU:HA	6:X:230:HIS:HB2	2.00	0.44
6:Y:120:ARG:O	6:Y:123:THR:HB	2.16	0.44
6:Y:228:LEU:N	6:Y:354:TYR:OH	2.50	0.44
6:Z:120:ARG:O	6:Z:123:THR:HB	2.17	0.44
1:B:246:GLN:NE2	1:B:982:MET:H	2.15	0.44
1:B:278:GLN:O	1:B:281:LEU:HB3	2.17	0.44
1:B:342:GLU:OE1	1:B:343:THR:N	2.50	0.44
1:B:550:GLN:NE2	1:B:551:ILE:O	2.51	0.44
1:B:903:ASP:OD1	1:B:904:ALA:N	2.50	0.44
1:B:997:LEU:O	1:B:1012:ILE:HG23	2.18	0.44
4:K:13:VAL:HG12	4:K:249:TRP:HZ3	1.83	0.44
4:K:146:CYS:O	4:K:160:GLN:N	2.34	0.44
4:K:340:THR:N	4:K:363:GLY:O	2.49	0.44
4:K:347:ILE:O	4:K:358:HIS:HA	2.17	0.44
4:K:562:SER:OG	4:K:565:SER:OG	2.28	0.44
4:L:41:LEU:HD23	4:L:106:HIS:CE1	2.46	0.44
4:L:377:ARG:NH1	4:L:451:ASN:OD1	2.50	0.44
5:O:41:GLU:HB3	5:O:43:TRP:CH2	2.52	0.44
5:O:48:ASN:ND2	5:O:50:ARG:H	2.15	0.44
5:O:114:PHE:HA	5:O:117:ASN:ND2	2.32	0.44
5:O:496:ALA:O	5:O:504:GLU:HA	2.18	0.44
5:O:554:VAL:H	5:O:559:ILE:CD1	2.30	0.44
5:O:705:GLY:O	5:O:757:ARG:HA	2.17	0.44
5:O:919:ALA:O	5:O:1023:ILE:HD13	2.17	0.44
5:O:988:LEU:HB2	5:O:989:ARG:HH21	1.82	0.44
5:O:1098:ILE:HA	5:O:1117:PHE:O	2.17	0.44
3:P:161:ILE:HG12	3:P:327:LEU:HD22	2.00	0.44

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:P:168:GLU:OE2	3:P:171:TRP:HD1	2.00	0.44
6:X:205:GLN:N	6:X:267:LYS:HZ3	2.13	0.44
6:Y:249:PHE:HD1	6:Y:344:SER:OG	2.00	0.44
6:Z:179:LEU:HB3	6:Z:363:ILE:HD11	1.98	0.44
1:B:920:ARG:HH22	1:B:983:LEU:HD13	1.83	0.44
1:B:1116:GLN:OE1	1:B:1172:PRO:HA	2.17	0.44
2:C:271:ILE:HG23	2:C:272:VAL:HG22	2.00	0.44
2:C:337:GLN:H	2:C:357:ASN:ND2	2.14	0.44
2:C:716:PRO:HB3	2:C:742:PRO:HA	2.00	0.44
2:C:1112:LEU:CA	2:C:1139:ILE:HD11	2.48	0.44
4:K:103:VAL:HG11	4:K:141:LEU:HD11	1.98	0.44
4:K:149:GLY:HA3	4:K:154:GLN:HB2	1.99	0.44
4:K:334:LEU:CD2	4:K:368:VAL:HG22	2.48	0.44
4:L:18:ASN:HA	4:L:248:GLN:O	2.17	0.44
4:L:171:GLN:NE2	4:L:172:THR:OG1	2.50	0.44
4:M:111:PHE:O	4:M:114:ALA:HB3	2.17	0.44
4:M:329:ASP:OD1	4:M:330:GLU:HG2	2.18	0.44
4:M:376:MET:HE3	4:M:497:PRO:HA	1.98	0.44
4:M:459:LYS:HE3	4:M:464:MET:H	1.82	0.44
5:O:173:LEU:HB3	5:O:175:MET:HE1	1.99	0.44
5:O:432:GLY:N	5:O:477:LYS:O	2.36	0.44
5:O:740:SER:O	5:O:754:ILE:HD12	2.17	0.44
6:X:32:TRP:HZ2	6:X:65:LEU:HB3	1.82	0.44
6:X:243:VAL:HG22	6:X:349:PHE:CE2	2.53	0.44
6:Y:295:VAL:HG23	6:Y:324:TYR:CE1	2.48	0.44
6:Y:303:ASN:ND2	6:Y:320:MET:HB3	2.31	0.44
1:B:518:GLN:HE22	1:B:521:ARG:NH2	2.15	0.44
1:B:1094:ILE:O	1:B:1102:VAL:N	2.48	0.44
2:C:406:MET:HB3	2:C:1254:TYR:OH	2.17	0.44
2:C:662:ILE:HG23	2:C:688:PHE:CZ	2.52	0.44
2:C:753:PRO:HB2	2:C:756:ASP:OD2	2.18	0.44
3:D:159:GLY:HA3	3:D:274:PRO:HG2	2.00	0.44
3:D:306:ASN:ND2	3:D:320:THR:HG22	2.33	0.44
4:K:198:LEU:HA	4:M:566:VAL:HG12	2.00	0.44
4:K:322:ASN:ND2	4:K:490:ASP:OD1	2.50	0.44
4:K:520:GLY:O	4:K:613:GLN:HA	2.17	0.44
4:K:522:TYR:HD2	4:K:526:SER:OG	2.01	0.44
4:L:235:GLU:OE1	4:L:235:GLU:N	2.36	0.44
4:M:307:VAL:HG12	4:M:513:VAL:O	2.18	0.44
5:O:28:HIS:HB3	5:O:108:ASN:HD21	1.82	0.44
5:O:285:LEU:HD11	5:O:287:TRP:O	2.17	0.44

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
5:O:419:GLN:HG2	5:O:709:ILE:HA	1.98	0.44
5:O:481:ARG:HB2	5:O:645:VAL:HG11	2.00	0.44
5:O:547:VAL:HG12	5:O:549:GLN:HG3	2.00	0.44
3:P:111:VAL:HA	3:P:114:GLN:CD	2.38	0.44
3:P:174:ASP:O	3:P:178:LEU:HD23	2.18	0.44
3:P:209:TYR:N	3:P:209:TYR:CD1	2.83	0.44
3:P:292:THR:HG23	3:P:295:LEU:HD12	2.00	0.44
3:P:360:LEU:O	3:P:364:THR:HG23	2.17	0.44
6:X:26:TYR:HB2	6:X:32:TRP:CH2	2.52	0.44
6:X:158:ASP:O	6:X:162:LYS:HG2	2.16	0.44
6:X:218:TRP:CD1	6:X:219:GLY:N	2.85	0.44
6:X:330:GLN:CD	6:X:334:VAL:HG21	2.38	0.44
6:Y:2:GLU:HB2	6:Y:4:CYS:SG	2.58	0.44
6:Y:128:LEU:HD21	6:Y:179:LEU:HD13	2.00	0.44
6:Y:142:PHE:HA	6:Y:217:GLU:HA	1.98	0.44
6:Z:179:LEU:HD22	6:Z:363:ILE:HG13	1.99	0.44
6:Z:330:GLN:HG3	6:Z:334:VAL:HG13	1.99	0.44
1:B:285:PHE:CZ	1:B:1217:PRO:HD3	2.53	0.44
1:B:530:THR:HA	1:B:533:GLN:CD	2.38	0.44
1:B:589:SER:HA	1:B:592:ARG:HH22	1.83	0.44
1:B:963:ALA:HA	1:B:966:ALA:HB3	2.00	0.44
1:B:1147:MET:HG2	1:B:1177:SER:OG	2.18	0.44
2:C:347:ARG:HA	2:C:347:ARG:HH11	1.82	0.44
2:C:682:HIS:NE2	2:C:683:THR:HG23	2.33	0.44
2:C:770:GLU:HA	2:C:773:LYS:HD2	2.00	0.44
2:C:806:MET:SD	2:C:890:LYS:HA	2.57	0.44
2:C:1151:TYR:O	2:C:1183:PRO:HA	2.17	0.44
3:D:71:LEU:HD23	3:D:72:LEU:N	2.32	0.44
3:D:230:GLN:HE21	3:D:240:GLN:NE2	2.14	0.44
3:D:331:GLN:CD	3:D:333:ARG:H	2.20	0.44
4:L:103:VAL:HB	4:L:164:PRO:HG2	1.99	0.44
4:L:118:LEU:HD21	4:L:182:LEU:HG	2.00	0.44
5:O:21:ARG:CZ	5:O:22:THR:HG22	2.47	0.44
5:O:221:LEU:HD22	5:O:285:LEU:HD12	1.99	0.44
5:O:240:GLN:HB2	5:O:249:ILE:C	2.37	0.44
5:O:448:VAL:HB	5:O:666:TRP:C	2.38	0.44
5:O:632:LEU:HD11	5:O:666:TRP:CZ3	2.52	0.44
5:O:955:GLY:O	5:O:956:ARG:HG2	2.18	0.44
5:O:1242:GLU:HB2	5:O:1284:TYR:CE1	2.53	0.44
3:P:202:PRO:HB3	3:P:294:CYS:SG	2.58	0.44
3:P:231:GLN:HE22	3:P:258:ASN:HB2	1.82	0.44

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
6:X:86:GLN:HA	6:X:89:ASP:OD2	2.18	0.44
6:X:183:PHE:CE1	6:X:264:ILE:HD11	2.52	0.44
6:Y:303:ASN:HD21	6:Y:320:MET:HB3	1.83	0.44
6:Z:174:ILE:O	6:Z:178:ASP:N	2.50	0.44
6:Z:251:HIS:ND1	6:Z:254:LEU:HD12	2.32	0.44
6:Z:309:GLU:OE1	6:Z:309:GLU:N	2.42	0.44
1:B:351:PRO:HA	1:B:354:PHE:CE1	2.53	0.44
1:B:381:ASP:HB2	1:B:388:GLY:CA	2.48	0.44
1:B:543:LEU:HA	1:B:546:ILE:HG12	2.00	0.44
1:B:623:LEU:HA	1:B:626:LEU:HG	2.00	0.44
1:B:1036:ASN:ND2	1:B:1205:ASN:OD1	2.51	0.44
1:B:1110:PHE:HZ	1:B:1115:TRP:N	2.15	0.44
2:C:361:MET:SD	2:C:362:LEU:HD12	2.57	0.44
2:C:719:ILE:HG12	2:C:720:ARG:N	2.32	0.44
2:C:761:LEU:HD12	2:C:808:PRO:HB3	2.00	0.44
2:C:1118:ASN:HB2	2:C:1122:PHE:CZ	2.53	0.44
3:D:79:PRO:HD3	3:D:96:TRP:NE1	2.29	0.44
3:D:350:THR:HG22	3:D:353:GLN:HG3	2.00	0.44
4:K:144:VAL:HG12	4:K:145:ASP:N	2.29	0.44
4:K:530:SER:O	4:K:537:ARG:NH2	2.49	0.44
4:K:566:VAL:C	4:K:569:GLN:HE21	2.18	0.44
4:K:620:ASP:OD2	4:K:622:ASN:HB2	2.18	0.44
4:L:50:ALA:N	4:L:63:LEU:HG	2.32	0.44
4:L:296:ALA:O	4:L:542:ARG:NE	2.51	0.44
4:L:367:VAL:CG2	4:L:467:TYR:HB3	2.46	0.44
4:M:377:ARG:HD3	4:M:496:SER:OG	2.17	0.44
4:M:426:ASN:ND2	4:M:449:ALA:N	2.66	0.44
4:M:431:TYR:CE2	4:M:433:GLU:HG3	2.52	0.44
4:M:577:LEU:C	4:M:584:ARG:HH12	2.21	0.44
4:M:577:LEU:O	4:M:584:ARG:NH2	2.51	0.44
4:M:589:LYS:HA	4:M:592:LEU:HB3	1.99	0.44
5:O:260:CYS:HB3	5:O:264:PHE:CZ	2.53	0.44
5:O:316:ARG:NH1	5:O:784:ALA:O	2.51	0.44
5:O:472:PHE:CD1	5:O:475:ALA:HB3	2.52	0.44
5:O:768:ARG:HH12	5:O:840:ILE:HG13	1.82	0.44
5:O:940:LEU:HD13	5:O:951:PHE:CZ	2.53	0.44
5:O:1015:TYR:C	5:O:1017:PRO:HD3	2.37	0.44
5:O:1096:ALA:HA	5:O:1119:VAL:HB	1.99	0.44
5:O:1128:ASP:OD2	5:O:1130:TRP:HB2	2.17	0.44
3:P:243:ARG:HE	3:P:249:GLU:HA	1.81	0.44
6:X:160:ASP:OD1	6:X:161:THR:N	2.44	0.44

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
6:X:164:ASP:HA	6:X:280:CYS:SG	2.58	0.44
6:Y:140:SER:N	6:Y:144:SER:OG	2.51	0.44
6:Z:17:ASN:HA	6:Z:20:GLU:CD	2.38	0.44
6:Z:84:ASP:OD1	6:Z:85:VAL:N	2.51	0.44
6:Z:185:MET:H	6:Z:261:THR:CA	2.30	0.44
6:Z:196:ARG:NE	6:Z:352:LEU:HB3	2.27	0.44
6:Z:331:ALA:HB3	6:Z:332:PRO:HD3	1.99	0.44
1:B:376:THR:HB	1:B:392:ARG:CZ	2.48	0.44
1:B:414:PRO:HA	2:C:1082:ARG:HB3	2.00	0.44
1:B:419:ILE:O	1:B:423:VAL:HG23	2.18	0.44
1:B:619:SER:O	1:B:622:SER:N	2.43	0.44
1:B:741:LEU:HG	1:B:743:ILE:N	2.28	0.44
1:B:1115:TRP:HA	1:B:1122:PHE:CD1	2.52	0.44
2:C:355:ARG:HD3	2:C:949:LEU:HB3	1.98	0.44
2:C:464:ARG:CZ	2:C:1026:LEU:HD22	2.47	0.44
2:C:514:ARG:HH12	2:C:729:SER:N	2.16	0.44
2:C:580:LYS:HE2	2:C:627:TRP:HB2	2.00	0.44
2:C:991:ASP:O	2:C:993:ARG:N	2.50	0.44
2:C:1070:ASP:OD1	2:C:1071:THR:N	2.51	0.44
2:C:1168:GLU:HB3	2:C:1169:GLU:OE1	2.18	0.44
3:D:6:PHE:HA	3:D:144:TYR:HE1	1.82	0.44
3:D:350:THR:H	3:D:353:GLN:NE2	2.16	0.44
4:K:582:GLY:HA3	6:X:69:PRO:O	2.17	0.44
4:K:657:ILE:HD12	4:K:660:TRP:HB3	1.99	0.44
4:L:273:SER:OG	4:M:179:GLN:HB3	2.17	0.44
4:L:324:ARG:HH21	4:L:489:TRP:N	2.15	0.44
4:L:525:GLU:HA	4:L:528:ASN:HB3	2.00	0.44
4:L:608:SER:O	4:L:611:ILE:HG12	2.17	0.44
4:M:234:LYS:HD3	4:M:234:LYS:H	1.82	0.44
5:O:182:ASP:HB3	5:O:325:ARG:NE	2.33	0.44
5:O:507:ARG:HD2	5:O:507:ARG:N	2.33	0.44
5:O:638:TYR:HE2	5:O:664:LEU:HB3	1.83	0.44
5:O:1159:ILE:HB	5:O:1162:GLN:HB2	2.00	0.44
3:P:3:ARG:HB2	3:P:300:GLN:HB2	1.99	0.44
3:P:6:PHE:O	3:P:127:CYS:N	2.45	0.44
3:P:99:PRO:HB2	3:P:132:PHE:CZ	2.53	0.44
3:P:143:VAL:HG12	3:P:147:LEU:HD21	1.99	0.44
3:P:156:THR:HG22	3:P:158:PHE:N	2.33	0.44
3:P:333:ARG:HD3	3:P:334:ARG:N	2.32	0.44
6:X:32:TRP:CZ2	6:X:65:LEU:HB3	2.52	0.44
6:Y:5:LEU:HD23	6:Y:315:LEU:HD23	2.00	0.44

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
6:Y:186:ARG:NH2	6:Y:225:TYR:OH	2.51	0.44
6:Z:185:MET:HA	6:Z:223:TYR:OH	2.18	0.44
6:Z:238:TYR:CE2	6:Z:242:LEU:HD13	2.53	0.44
6:Z:288:GLY:HA2	6:Z:291:LYS:CD	2.48	0.44
1:B:303:ARG:HG2	1:B:1209:LEU:HD12	1.99	0.44
1:B:307:HIS:CE1	1:B:310:TRP:HB3	2.53	0.44
1:B:1044:ASP:O	1:B:1202:THR:N	2.50	0.44
1:B:1110:PHE:CE1	1:B:1112:LEU:HA	2.53	0.44
2:C:556:ILE:H	2:C:556:ILE:HG13	1.67	0.44
2:C:638:LEU:H	2:C:638:LEU:HD12	1.82	0.44
2:C:674:ARG:HA	2:C:674:ARG:HD3	1.75	0.44
2:C:790:VAL:HG23	2:C:791:SER:N	2.33	0.44
2:C:1042:ARG:O	2:C:1144:TYR:OH	2.18	0.44
2:C:1231:ARG:HA	2:C:1252:ASP:OD1	2.17	0.44
3:D:208:THR:HA	3:D:211:ARG:NE	2.33	0.44
4:K:19:VAL:O	4:K:248:GLN:NE2	2.51	0.44
4:L:583:VAL:O	6:Z:70:HIS:HB2	2.18	0.44
4:M:235:GLU:OE1	4:M:235:GLU:N	2.30	0.44
4:M:328:ILE:HG13	4:M:329:ASP:N	2.33	0.44
4:M:406:SER:C	4:M:407:LYS:HD2	2.38	0.44
5:O:111:VAL:HG22	5:O:137:ILE:HA	1.98	0.44
5:O:669:GLY:HA2	5:O:672:PHE:CD1	2.53	0.44
5:O:877:TRP:O	5:O:880:GLY:N	2.51	0.44
3:P:7:LEU:HG	3:P:126:ASP:HA	2.00	0.44
3:P:197:ALA:HA	3:P:200:ASN:HD21	1.83	0.44
6:X:35:THR:C	6:X:36:ILE:HG13	2.38	0.44
6:X:259:ARG:O	6:X:272:PHE:N	2.50	0.44
6:Z:142:PHE:C	6:Z:144:SER:H	2.21	0.44
1:B:311:ALA:HB1	1:B:402:TRP:CH2	2.53	0.43
1:B:451:PHE:HD1	1:B:1256:VAL:HG13	1.82	0.43
1:B:520:ILE:HG23	1:B:523:MET:SD	2.58	0.43
1:B:1115:TRP:CZ2	1:B:1167:LEU:HD11	2.53	0.43
1:B:1147:MET:O	1:B:1180:PHE:N	2.51	0.43
1:B:1158:ASN:HD21	1:B:1160:TRP:HB2	1.83	0.43
2:C:303:ARG:HE	2:C:1209:LEU:HD13	1.83	0.43
2:C:543:LEU:O	2:C:546:ILE:HG22	2.18	0.43
2:C:554:THR:O	2:C:558:ASN:ND2	2.51	0.43
2:C:658:GLY:H	2:C:661:THR:HG22	1.83	0.43
2:C:703:ARG:HG3	2:C:707:GLU:OE2	2.18	0.43
2:C:1154:ARG:HE	2:C:1190:ILE:HB	1.83	0.43
3:D:96:TRP:CZ3	3:D:101:LEU:HB3	2.53	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:D:107:ALA:H	3:D:112:LEU:HD11	1.83	0.43
3:D:296:CYS:SG	3:D:301:LEU:HD12	2.57	0.43
4:L:522:TYR:HE2	4:L:530:SER:OG	2.00	0.43
4:M:21:LYS:HA	4:M:208:VAL:O	2.18	0.43
5:O:17:THR:HG22	5:O:358:PRO:C	2.38	0.43
5:O:476:ARG:HH12	5:O:644:PHE:HZ	1.66	0.43
5:O:626:TYR:HA	5:O:629:GLN:HB2	1.99	0.43
5:O:628:GLU:OE1	5:O:665:THR:HA	2.19	0.43
5:O:696:TYR:CE2	5:O:698:ASP:HA	2.53	0.43
5:O:1192:LYS:NZ	5:O:1220:ILE:H	2.15	0.43
5:O:1240:VAL:HB	5:O:1247:ILE:HB	2.00	0.43
3:P:212:MET:O	3:P:216:ILE:HG12	2.17	0.43
6:X:287:LYS:O	6:X:291:LYS:HE3	2.17	0.43
6:Y:141:MET:HB3	6:Y:147:SER:O	2.18	0.43
6:Y:216:LEU:HD23	6:Y:216:LEU:H	1.83	0.43
6:Z:16:ASN:HA	6:Z:19:PHE:CD2	2.53	0.43
6:Z:252:PHE:HE1	6:Z:258:SER:C	2.21	0.43
1:B:303:ARG:HA	1:B:1210:PHE:CB	2.48	0.43
1:B:761:LEU:HA	1:B:764:TRP:HB3	2.00	0.43
1:B:970:ASN:HA	1:B:973:MET:HE2	2.00	0.43
2:C:477:THR:OG1	2:C:478:GLU:N	2.51	0.43
2:C:1151:TYR:N	2:C:1182:VAL:O	2.51	0.43
3:D:151:THR:HG23	3:D:154:ASN:HD22	1.82	0.43
3:D:342:ARG:CZ	3:D:345:ARG:HD3	2.48	0.43
3:D:359:ALA:O	3:D:363:GLN:HG3	2.18	0.43
4:K:233:PRO:HD2	4:K:234:LYS:HZ2	1.82	0.43
4:L:336:MET:N	4:L:411:GLU:OE2	2.49	0.43
4:L:367:VAL:HG22	4:L:368:VAL:H	1.82	0.43
4:L:573:ALA:O	4:L:577:LEU:HG	2.18	0.43
4:L:663:GLY:O	4:L:667:LYS:HG3	2.18	0.43
4:M:108:ILE:HG23	4:M:137:TYR:CE2	2.54	0.43
4:M:288:GLN:OE1	4:M:291:ASP:HB2	2.18	0.43
4:M:578:GLU:N	4:M:584:ARG:HH12	2.17	0.43
5:O:5:TRP:HB3	5:O:336:SER:O	2.18	0.43
5:O:47:ARG:HG2	5:O:54:ILE:HG12	1.99	0.43
5:O:83:GLU:OE2	5:O:84:ARG:HG3	2.18	0.43
5:O:118:ALA:HA	5:O:156:PHE:CE2	2.52	0.43
5:O:422:LEU:C	5:O:697:VAL:HG13	2.38	0.43
5:O:606:THR:HG22	5:O:607:ALA:N	2.33	0.43
5:O:654:VAL:HG13	5:O:656:PHE:CE1	2.53	0.43
5:O:671:TYR:HA	5:O:674:LEU:HG	1.99	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
5:O:995:LEU:HA	5:O:995:LEU:HD12	1.83	0.43
5:O:1145:ILE:HB	5:O:1181:LEU:CB	2.48	0.43
3:P:228:ILE:HA	3:P:232:ASN:ND2	2.33	0.43
3:P:375:TRP:HA	3:P:378:GLU:OE1	2.18	0.43
6:Z:10:GLN:OE1	6:Z:10:GLN:N	2.21	0.43
6:Z:87:PHE:O	6:Z:91:VAL:HG23	2.17	0.43
6:Z:231:ASP:OD2	6:Z:233:SER:OG	2.34	0.43
6:Z:312:ARG:NH2	6:Z:317:PRO:HG3	2.33	0.43
1:B:307:HIS:CD2	1:B:323:ILE:HD13	2.53	0.43
1:B:474:ILE:HD12	1:B:505:MET:O	2.18	0.43
1:B:613:TYR:HD2	1:B:615:LYS:HD3	1.83	0.43
1:B:1039:GLY:HA2	1:B:1146:TYR:OH	2.18	0.43
1:B:1109:ILE:HD12	1:B:1138:ARG:NE	2.31	0.43
2:C:1049:ARG:CZ	2:C:1132:THR:HG23	2.48	0.43
2:C:1075:HIS:HD1	2:C:1108:TRP:HD1	1.65	0.43
2:C:1123:ASN:N	2:C:1123:ASN:OD1	2.51	0.43
3:D:210:ALA:O	3:D:211:ARG:C	2.57	0.43
4:K:167:THR:HG23	4:K:170:ARG:H	1.83	0.43
4:K:367:VAL:HG21	4:K:467:TYR:HB3	1.99	0.43
4:K:645:LYS:NZ	4:K:649:SER:HB3	2.33	0.43
4:L:13:VAL:HA	4:L:249:TRP:CH2	2.52	0.43
4:L:103:VAL:HG21	4:L:141:LEU:HD22	2.01	0.43
4:L:352:ASN:OD1	6:Y:64:LYS:HD2	2.19	0.43
4:L:393:ASP:OD2	6:Y:277:THR:HB	2.19	0.43
4:L:610:ILE:HG13	4:L:611:ILE:HG23	2.01	0.43
4:L:636:ARG:NH1	4:M:27:SER:HB3	2.32	0.43
4:M:233:PRO:HD2	4:M:234:LYS:HZ3	1.84	0.43
5:O:8:ARG:HE	5:O:716:PHE:HD2	1.65	0.43
5:O:61:ARG:NE	5:O:64:GLN:OE1	2.44	0.43
5:O:616:ILE:HB	5:O:651:LEU:CB	2.48	0.43
5:O:994:ALA:O	5:O:999:THR:N	2.51	0.43
5:O:1054:ASN:HD21	5:O:1104:LEU:HD11	1.82	0.43
5:O:1230:LEU:HD22	5:O:1277:TYR:CD2	2.52	0.43
3:P:83:ASN:OD1	3:P:83:ASN:N	2.42	0.43
3:P:110:GLN:O	3:P:114:GLN:HG3	2.18	0.43
3:P:219:LEU:HD22	3:P:239:PHE:CE1	2.53	0.43
3:P:306:ASN:ND2	3:P:322:ARG:HA	2.33	0.43
6:X:14:LEU:CD1	6:X:45:CYS:HA	2.48	0.43
6:Y:24:SER:O	6:Y:43:MET:HA	2.19	0.43
6:Z:50:VAL:HG13	6:Z:56:GLY:O	2.18	0.43
1:B:1076:ILE:HG22	1:B:1077:PHE:H	1.83	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:C:317:ASP:CG	2:C:318:ARG:H	2.20	0.43
2:C:398:ASP:HB2	2:C:402:TRP:HE1	1.83	0.43
2:C:808:PRO:HA	2:C:811:LEU:HG	1.99	0.43
2:C:1095:ARG:N	2:C:1101:MET:HG3	2.33	0.43
3:D:179:LEU:O	3:D:182:TYR:N	2.49	0.43
3:D:349:MET:HE3	3:D:349:MET:HB3	1.88	0.43
4:L:299:GLU:OE2	4:M:650:LYS:NZ	2.48	0.43
4:L:362:ARG:CD	4:L:486:PRO:HA	2.45	0.43
4:L:658:GLN:NE2	4:L:659:ASN:HB2	2.33	0.43
4:M:57:VAL:HG13	4:M:59:SER:O	2.18	0.43
4:M:385:LYS:NZ	4:M:491:ALA:HA	2.34	0.43
4:M:620:ASP:OD1	4:M:621:LYS:HG2	2.19	0.43
4:M:631:VAL:HG13	4:M:632:LYS:H	1.84	0.43
5:O:10:ALA:HB2	5:O:313:GLN:CG	2.48	0.43
5:O:26:THR:H	5:O:29:ASP:HB2	1.81	0.43
5:O:114:PHE:HA	5:O:117:ASN:HD22	1.82	0.43
5:O:453:LEU:HD21	5:O:624:TRP:CD1	2.54	0.43
5:O:492:HIS:C	5:O:494:TYR:H	2.21	0.43
5:O:529:VAL:HG23	5:O:530:ILE:HD13	1.99	0.43
5:O:573:PHE:HE2	5:O:575:TYR:HB2	1.81	0.43
5:O:770:ARG:NH2	5:O:859:GLY:O	2.28	0.43
5:O:907:PHE:CZ	5:O:911:ILE:HD11	2.53	0.43
5:O:931:ASP:HB3	5:O:1015:TYR:CE1	2.52	0.43
5:O:1167:ASN:HB3	5:O:1170:LYS:HB2	1.99	0.43
5:O:1189:ILE:HG22	5:O:1193:TYR:HE1	1.83	0.43
5:O:1262:ASP:HB3	5:O:1278:ILE:HB	2.01	0.43
3:P:81:ARG:NE	3:P:83:ASN:O	2.52	0.43
3:P:241:CYS:O	3:P:244:ARG:HG3	2.19	0.43
3:P:267:ASN:OD1	3:P:268:ARG:N	2.51	0.43
3:P:333:ARG:HG2	3:P:362:ASP:OD1	2.19	0.43
6:Y:283:TYR:CE1	6:Y:337:PRO:HG2	2.53	0.43
6:Z:74:ASN:HB2	6:Z:76:GLN:OE1	2.18	0.43
1:B:768:VAL:O	1:B:772:MET:HG2	2.18	0.43
1:B:1036:ASN:ND2	1:B:1207:ARG:HE	2.00	0.43
2:C:268:ILE:H	2:C:268:ILE:HD12	1.84	0.43
2:C:1110:PHE:HZ	2:C:1115:TRP:N	2.16	0.43
2:C:1115:TRP:CD1	2:C:1119:THR:HG23	2.54	0.43
2:C:1160:TRP:HD1	2:C:1161:ASN:OD1	2.01	0.43
2:C:1253:LEU:HD23	2:C:1253:LEU:HA	1.77	0.43
3:D:246:ASP:OD1	3:D:248:ALA:N	2.39	0.43
4:K:193:ARG:NH2	4:M:634:SER:O	2.50	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
4:K:299:GLU:CD	4:K:299:GLU:H	2.21	0.43
4:K:377:ARG:HB2	4:K:498:LEU:HD21	2.00	0.43
4:K:606:ASP:OD1	4:K:608:SER:N	2.51	0.43
4:L:20:PHE:O	4:L:209:SER:OG	2.19	0.43
4:L:35:SER:O	4:L:35:SER:OG	2.36	0.43
4:L:126:ASP:OD2	4:L:132:SER:HA	2.17	0.43
4:L:228:LEU:O	4:L:232:TYR:N	2.51	0.43
4:L:347:ILE:HD12	4:L:348:GLN:H	1.83	0.43
4:L:392:TRP:CH2	4:L:397:LYS:HD2	2.54	0.43
4:L:654:SER:O	4:L:658:GLN:HG3	2.17	0.43
4:M:105:GLU:O	4:M:108:ILE:N	2.51	0.43
4:M:131:LEU:HD12	4:M:131:LEU:HA	1.82	0.43
4:M:402:ILE:HG13	4:M:423:THR:O	2.18	0.43
5:O:9:LEU:O	5:O:315:ALA:HB2	2.19	0.43
5:O:207:LEU:HD13	5:O:210:PHE:HB3	1.99	0.43
5:O:241:MET:N	5:O:249:ILE:O	2.40	0.43
5:O:768:ARG:HG2	5:O:842:ALA:HB2	2.00	0.43
3:P:84:HIS:CD2	3:P:130:TYR:HD1	2.36	0.43
3:P:283:VAL:HG22	3:P:287:LEU:HD23	1.99	0.43
3:P:327:LEU:HD23	3:P:327:LEU:HA	1.63	0.43
3:P:331:GLN:HE22	3:P:403:GLY:N	2.15	0.43
3:P:367:PHE:O	3:P:371:LYS:HG2	2.18	0.43
6:X:84:ASP:OD1	6:X:85:VAL:HG23	2.18	0.43
6:X:142:PHE:O	6:X:217:GLU:HA	2.18	0.43
6:Y:180:VAL:HB	6:Y:364:LEU:HD23	2.01	0.43
6:Y:222:VAL:HG23	6:Y:357:MET:SD	2.59	0.43
6:Z:142:PHE:CD1	6:Z:217:GLU:HB3	2.54	0.43
6:Z:212:SER:O	6:Z:213:ARG:NE	2.52	0.43
1:B:318:ARG:NE	1:B:371:TYR:HA	2.33	0.43
1:B:532:ILE:HG23	1:B:536:LEU:HD11	2.00	0.43
1:B:634:LEU:HB2	1:B:883:THR:HG22	2.00	0.43
1:B:718:GLN:OE1	1:B:718:GLN:N	2.50	0.43
1:B:931:LEU:HA	1:B:934:LEU:HG	2.00	0.43
1:B:1019:VAL:HG22	1:B:1023:CYS:SG	2.58	0.43
1:B:1166:TRP:NE1	1:B:1179:PRO:HD3	2.32	0.43
2:C:1146:TYR:HE1	2:C:1148:LEU:HD21	1.83	0.43
3:D:236:THR:O	3:D:237:PHE:HD1	2.01	0.43
4:K:353:THR:HA	6:Z:32:TRP:CE3	2.53	0.43
4:K:371:ASP:N	4:K:373:ILE:HG12	2.34	0.43
4:K:583:VAL:HG11	6:X:67:HIS:NE2	2.34	0.43
4:L:23:SER:HB2	4:L:25:GLU:OE1	2.19	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
4:L:283:SER:O	4:L:286:THR:OG1	2.26	0.43
4:L:393:ASP:H	4:L:397:LYS:HZ2	1.67	0.43
4:L:398:LYS:HE3	4:L:473:ILE:CD1	2.46	0.43
4:L:437:PHE:HB3	4:L:440:GLN:CG	2.49	0.43
4:M:27:SER:HA	4:M:202:ASN:CB	2.49	0.43
4:M:403:VAL:HG22	4:M:404:PHE:H	1.83	0.43
4:M:405:GLN:HE21	4:M:421:GLN:CB	2.29	0.43
4:M:472:PHE:CE1	4:M:485:GLN:HG3	2.46	0.43
5:O:96:GLU:O	5:O:99:ARG:N	2.51	0.43
5:O:587:ASP:O	5:O:590:ILE:HB	2.18	0.43
5:O:1257:LEU:HD11	5:O:1261:TRP:HB2	1.99	0.43
3:P:311:ALA:HA	3:P:314:ARG:CD	2.47	0.43
6:X:104:CYS:HA	6:X:166:TYR:CE1	2.53	0.43
6:X:170:LEU:O	6:X:174:ILE:HG23	2.19	0.43
6:Z:311:ILE:O	6:Z:316:GLY:N	2.45	0.43
6:Z:315:LEU:HD12	6:Z:323:TRP:CD1	2.53	0.43
1:B:643:CYS:SG	1:B:684:TRP:HZ2	2.42	0.43
1:B:760:GLU:O	1:B:767:ARG:NH1	2.51	0.43
1:B:819:LEU:HA	1:B:822:ILE:HD11	2.01	0.43
1:B:1093:MET:HA	1:B:1102:VAL:O	2.18	0.43
2:C:404:SER:O	2:C:408:PRO:HA	2.19	0.43
2:C:544:GLN:HA	2:C:592:ARG:NE	2.34	0.43
2:C:898:ASN:HA	2:C:901:TYR:HD1	1.83	0.43
2:C:1022:ASP:HA	2:C:1025:GLN:HG3	2.01	0.43
2:C:1048:GLY:O	2:C:1198:TYR:HA	2.18	0.43
2:C:1057:SER:O	2:C:1060:ALA:N	2.36	0.43
2:C:1101:MET:N	2:C:1101:MET:SD	2.92	0.43
2:C:1153:PRO:HG3	2:C:1183:PRO:HB3	2.01	0.43
3:D:139:PHE:O	3:D:143:VAL:HG22	2.19	0.43
3:D:185:HIS:O	3:D:252:ILE:HG13	2.19	0.43
4:K:24:ALA:HB2	4:K:206:GLY:N	2.33	0.43
4:K:297:GLU:O	4:K:300:ILE:HG22	2.19	0.43
4:K:413:TRP:HA	4:K:418:GLN:OE1	2.18	0.43
4:K:562:SER:HG	4:K:565:SER:HG	1.59	0.43
4:L:337:ILE:HG13	4:L:341:MET:HB2	2.01	0.43
4:L:650:LYS:HD3	4:L:650:LYS:C	2.39	0.43
4:M:195:ALA:O	4:M:198:LEU:HG	2.19	0.43
4:M:218:LEU:HD13	4:M:672:PHE:CD2	2.54	0.43
4:M:338:PRO:O	4:M:343:THR:HG21	2.18	0.43
5:O:140:ASP:OD2	5:O:141:THR:N	2.52	0.43
5:O:198:ALA:C	5:O:199:PHE:CG	2.92	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
5:O:322:LEU:N	5:O:322:LEU:HD12	2.34	0.43
5:O:424:PRO:HA	5:O:792:PRO:HG2	2.00	0.43
5:O:794:LEU:HG	5:O:795:PHE:CD1	2.54	0.43
5:O:804:HIS:H	5:O:804:HIS:CD2	2.36	0.43
5:O:828:LEU:HG	5:O:887:THR:O	2.19	0.43
5:O:977:CYS:HA	5:O:1022:ASP:O	2.18	0.43
3:P:41:GLN:HG3	3:P:64:ARG:HH22	1.83	0.43
6:X:242:LEU:HD23	6:X:242:LEU:H	1.84	0.43
6:Z:107:HIS:O	6:Z:110:MET:HB3	2.19	0.43
1:B:667:GLN:NE2	1:B:667:GLN:HA	2.34	0.43
1:B:702:LEU:O	1:B:705:TRP:HB2	2.19	0.43
1:B:781:TYR:O	1:B:785:TRP:HB2	2.19	0.43
1:B:928:VAL:HG13	1:B:987:LEU:HD21	2.00	0.43
1:B:1047:ILE:CG2	1:B:1137:ILE:HB	2.49	0.43
1:B:1148:LEU:HD13	1:B:1150:TYR:HE1	1.84	0.43
2:C:955:LEU:HD12	2:C:956:ARG:N	2.34	0.43
2:C:1062:PRO:HD2	2:C:1065:LEU:HD12	2.01	0.43
3:D:293:GLY:O	3:D:297:SER:OG	2.27	0.43
4:K:337:ILE:O	4:K:364:GLY:HA2	2.18	0.43
4:K:388:LYS:NZ	4:K:394:PRO:HD2	2.32	0.43
4:K:547:ALA:HA	4:K:550:ILE:HD12	2.00	0.43
4:L:300:ILE:HG12	4:M:650:LYS:NZ	2.34	0.43
4:L:407:LYS:HE2	4:L:463:GLU:H	1.84	0.43
4:L:648:VAL:O	4:L:651:LEU:HG	2.17	0.43
5:O:456:LEU:HD21	5:O:460:TYR:CB	2.49	0.43
5:O:509:ARG:HA	5:O:544:PRO:C	2.39	0.43
5:O:633:PRO:HA	5:O:662:SER:O	2.18	0.43
5:O:742:ALA:HA	5:O:787:ILE:CB	2.47	0.43
5:O:842:ALA:HB1	5:O:863:VAL:HG11	1.99	0.43
5:O:1030:LYS:O	5:O:1031:ARG:HD3	2.18	0.43
5:O:1081:ASP:OD1	5:O:1083:THR:OG1	2.33	0.43
5:O:1257:LEU:HD12	5:O:1258:PRO:HD2	2.01	0.43
3:P:231:GLN:O	3:P:233:THR:HG23	2.18	0.43
3:P:363:GLN:HB3	3:P:367:PHE:CZ	2.53	0.43
6:X:73:CYS:HB2	6:X:75:GLN:OE1	2.19	0.43
6:X:246:ALA:O	6:X:250:GLY:N	2.51	0.43
6:Y:206:PHE:CG	6:Y:267:LYS:HD2	2.53	0.43
6:Z:133:TRP:N	6:Z:360:ASP:OD2	2.41	0.43
1:B:340:ASN:ND2	1:B:342:GLU:O	2.51	0.43
1:B:459:VAL:HG21	1:B:1255:ASN:ND2	2.34	0.43
1:B:1034:GLU:O	1:B:1037:LEU:HB2	2.19	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:B:1115:TRP:NE1	1:B:1119:THR:HG23	2.33	0.43
2:C:604:VAL:HA	2:C:873:VAL:N	2.34	0.43
2:C:716:PRO:HB3	2:C:835:TYR:CD2	2.54	0.43
2:C:837:ARG:HD2	2:C:837:ARG:HA	1.77	0.43
2:C:1022:ASP:OD1	2:C:1023:CYS:N	2.52	0.43
2:C:1204:TYR:HD1	2:C:1206:ASP:H	1.65	0.43
3:D:16:LEU:HB2	3:D:20:PRO:CG	2.47	0.43
3:D:59:PRO:HG3	3:D:375:TRP:CH2	2.54	0.43
3:D:190:ILE:HG12	3:D:194:LEU:HD21	2.00	0.43
3:D:357:ILE:HD13	3:D:357:ILE:HA	1.86	0.43
4:K:147:TYR:CZ	4:K:159:PHE:HD1	2.36	0.43
4:K:574:ILE:HA	4:K:577:LEU:HB2	2.01	0.43
4:L:18:ASN:HB3	4:L:249:TRP:CD2	2.54	0.43
4:L:540:ILE:HD13	4:L:608:SER:HA	2.00	0.43
4:L:589:LYS:HA	4:L:592:LEU:HD12	2.01	0.43
4:M:327:LYS:HZ2	4:M:329:ASP:CG	2.21	0.43
4:M:456:GLN:NE2	4:M:456:GLN:O	2.51	0.43
4:M:517:GLU:OE1	6:Y:313:TYR:HE2	2.01	0.43
4:M:645:LYS:HA	4:M:645:LYS:HD2	1.89	0.43
5:O:34:LEU:O	5:O:90:LEU:HD11	2.19	0.43
5:O:186:PRO:HD3	5:O:324:ASN:HB3	2.01	0.43
5:O:426:TYR:CD2	5:O:694:PHE:HA	2.54	0.43
5:O:768:ARG:HA	5:O:862:ASN:ND2	2.34	0.43
5:O:811:TYR:CE2	5:O:1018:ILE:HD11	2.54	0.43
5:O:939:TYR:O	5:O:952:PRO:HD2	2.19	0.43
5:O:1133:GLN:NE2	5:O:1134:LEU:O	2.52	0.43
5:O:1150:TYR:HB3	5:O:1151:TYR:CD1	2.54	0.43
3:P:190:ILE:HD12	3:P:190:ILE:H	1.82	0.43
3:P:285:LEU:HG	3:P:286:ILE:HD13	2.01	0.43
3:P:353:GLN:HA	3:P:356:GLN:NE2	2.31	0.43
6:X:128:LEU:HD23	6:X:128:LEU:H	1.84	0.43
6:X:245:PRO:HB2	6:X:249:PHE:CD1	2.54	0.43
6:X:328:MET:SD	6:X:329:GLN:HG3	2.58	0.43
6:Y:21:GLY:C	6:Y:22:ARG:HD3	2.38	0.43
6:Y:86:GLN:CD	6:Y:86:GLN:N	2.69	0.43
6:Y:143:ARG:HG2	6:Y:265:LEU:HD11	2.01	0.43
6:Y:265:LEU:HD13	6:Y:265:LEU:HA	1.91	0.43
6:Z:190:HIS:HA	6:Z:193:ASN:O	2.18	0.43
6:Z:299:VAL:HG21	6:Z:324:TYR:CD1	2.54	0.43
1:B:293:GLN:HB2	1:B:404:SER:CB	2.48	0.43
1:B:369:ASN:HD22	1:B:458:THR:HG22	1.82	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:B:578:LEU:O	1:B:581:LEU:N	2.52	0.43
1:B:1035:TYR:HD2	1:B:1040:ILE:O	2.01	0.43
1:B:1086:GLY:HA3	1:B:1092:PRO:HG3	2.00	0.43
1:B:1093:MET:HB3	1:B:1101:MET:C	2.39	0.43
2:C:265:SER:CB	2:C:309:ARG:H	2.30	0.43
2:C:732:LEU:HG	2:C:1011:VAL:CG1	2.48	0.43
2:C:931:LEU:HD23	2:C:932:VAL:N	2.34	0.43
2:C:1117:MET:HE3	2:C:1117:MET:HB3	1.72	0.43
3:D:55:SER:HB3	3:D:59:PRO:CB	2.46	0.43
3:D:236:THR:HB	3:D:256:SER:O	2.18	0.43
4:K:13:VAL:HG12	4:K:249:TRP:CZ3	2.53	0.43
4:K:147:TYR:CE1	4:K:159:PHE:HD1	2.37	0.43
4:K:168:PRO:O	4:K:172:THR:OG1	2.30	0.43
4:K:227:SER:O	4:K:230:ARG:HG2	2.18	0.43
4:K:289:ALA:O	4:K:293:VAL:HG22	2.19	0.43
4:L:111:PHE:HD1	4:L:137:TYR:CE2	2.37	0.43
4:L:145:ASP:OD1	4:L:162:GLN:HG2	2.18	0.43
4:L:506:LYS:HD3	6:Z:316:GLY:CA	2.48	0.43
4:L:537:ARG:O	4:L:540:ILE:HG22	2.19	0.43
4:M:672:PHE:HD1	4:M:672:PHE:HA	1.67	0.43
5:O:4:VAL:HG13	5:O:341:GLU:HG2	2.00	0.43
5:O:67:VAL:O	5:O:68:LEU:HD23	2.18	0.43
5:O:93:LEU:O	5:O:97:VAL:HG13	2.18	0.43
5:O:608:PRO:HA	5:O:659:HIS:CD2	2.54	0.43
5:O:981:TRP:HB2	5:O:1116:SER:OG	2.19	0.43
5:O:1036:VAL:HG11	5:O:1150:TYR:CE1	2.54	0.43
3:P:65:TYR:HA	3:P:68:MET:SD	2.59	0.43
3:P:233:THR:CB	3:P:236:THR:HA	2.49	0.43
6:X:99:MET:O	6:X:103:VAL:HG22	2.19	0.43
6:X:262:THR:O	6:X:264:ILE:HG12	2.19	0.43
6:Y:159:LEU:H	6:Y:159:LEU:HD12	1.84	0.43
6:Y:180:VAL:HG12	6:Y:225:TYR:CE1	2.52	0.43
6:Z:40:PRO:HB2	6:Z:52:MET:SD	2.59	0.43
6:Z:50:VAL:HA	6:Z:58:VAL:HG22	2.01	0.43
1:B:333:HIS:HB2	1:B:337:GLN:NE2	2.34	0.42
1:B:484:THR:OG1	1:B:485:GLU:N	2.53	0.42
1:B:552:ASP:OD2	1:B:555:ILE:HG13	2.18	0.42
2:C:521:ARG:O	2:C:525:ILE:HG13	2.19	0.42
2:C:600:TYR:HA	2:C:831:PHE:O	2.18	0.42
2:C:949:LEU:HD13	2:C:949:LEU:HA	1.90	0.42
2:C:951:TRP:O	2:C:953:PRO:HD3	2.18	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:C:1110:PHE:O	2:C:1139:ILE:HD12	2.19	0.42
2:C:1209:LEU:HD12	2:C:1210:PHE:H	1.84	0.42
3:D:63:SER:CA	3:D:393:ARG:HB2	2.48	0.42
3:D:66:TYR:CE1	3:D:394:THR:HA	2.53	0.42
3:D:146:GLN:HG2	3:D:147:LEU:N	2.34	0.42
4:K:359:PHE:HA	4:K:478:ILE:HG21	2.01	0.42
4:L:45:GLY:HA3	4:L:149:GLY:O	2.18	0.42
4:L:254:GLU:HB3	4:L:256:THR:OG1	2.19	0.42
4:L:284:LYS:N	4:L:284:LYS:HD3	2.34	0.42
4:M:442:ILE:HD12	4:M:442:ILE:HA	1.83	0.42
4:M:450:TYR:CG	4:M:451:ASN:N	2.87	0.42
5:O:114:PHE:HZ	5:O:229:ASN:HA	1.84	0.42
5:O:122:ALA:O	5:O:125:SER:OG	2.29	0.42
5:O:188:PHE:CE2	5:O:258:GLU:HG3	2.54	0.42
5:O:372:LYS:HB3	5:O:374:GLN:CG	2.44	0.42
5:O:440:ASP:HB3	5:O:639:MET:SD	2.59	0.42
5:O:442:ASN:OD1	5:O:444:SER:N	2.52	0.42
5:O:486:ASP:HA	5:O:489:VAL:HG22	2.01	0.42
5:O:509:ARG:HB3	5:O:545:SER:HB3	1.99	0.42
5:O:707:GLU:CD	5:O:758:ARG:HH11	2.22	0.42
5:O:750:ARG:HD3	5:O:750:ARG:HA	1.82	0.42
5:O:986:TYR:OH	5:O:1059:SER:HB3	2.19	0.42
5:O:987:ASP:C	5:O:988:LEU:HD22	2.40	0.42
3:P:61:ALA:HA	3:P:64:ARG:HB2	2.00	0.42
3:P:165:ARG:NH1	3:P:267:ASN:HD21	2.17	0.42
3:P:223:SER:HB2	3:P:251:TRP:CZ3	2.54	0.42
3:P:370:ASP:HB2	3:P:371:LYS:HZ2	1.84	0.42
3:P:399:ASN:HA	3:P:402:TRP:HD1	1.79	0.42
6:X:304:HIS:O	6:X:306:TRP:N	2.52	0.42
6:Y:27:SER:N	6:Y:30:GLU:HB2	2.33	0.42
6:Y:42:MET:SD	6:Y:51:CYS:HA	2.59	0.42
6:Y:148:SER:N	6:Y:151:ASP:OD2	2.52	0.42
6:Z:54:CYS:HB3	6:Z:73:CYS:SG	2.58	0.42
6:Z:231:ASP:CG	6:Z:233:SER:HG	2.22	0.42
1:B:373:ASN:HB2	1:B:1259:ARG:CD	2.49	0.42
1:B:385:PHE:CZ	2:C:1117:MET:HA	2.55	0.42
1:B:502:ASN:C	1:B:503:ARG:HD2	2.40	0.42
1:B:644:ALA:HB3	1:B:645:PRO:HD3	2.01	0.42
2:C:573:PRO:HB3	2:C:623:LEU:HD13	2.01	0.42
2:C:685:PRO:CG	2:C:688:PHE:HB2	2.48	0.42
2:C:746:GLN:HB2	2:C:813:GLN:HG3	2.00	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:C:1070:ASP:OD1	2:C:1071:THR:HG23	2.19	0.42
3:D:39:LEU:HG	3:D:40:THR:N	2.34	0.42
3:D:63:SER:HB2	3:D:393:ARG:HD2	2.00	0.42
3:D:239:PHE:HB3	3:D:251:TRP:HB3	2.00	0.42
3:D:280:ASN:ND2	3:D:394:THR:OG1	2.51	0.42
4:K:262:VAL:O	4:L:43:PRO:HG2	2.17	0.42
4:K:344:LEU:HD11	4:K:480:PRO:HB3	2.01	0.42
4:K:615:ALA:O	4:K:619:SER:HB3	2.18	0.42
4:K:627:LEU:HD12	4:K:628:ALA:N	2.34	0.42
4:L:309:ALA:O	4:L:313:ALA:N	2.35	0.42
4:L:448:LEU:HD12	4:L:448:LEU:HA	1.90	0.42
4:L:515:PRO:HA	4:L:539:MET:HE1	2.01	0.42
4:M:357:TRP:CE3	4:M:474:ASP:HA	2.47	0.42
4:M:423:THR:HG22	4:M:424:VAL:O	2.18	0.42
4:M:520:GLY:N	4:M:611:ILE:HA	2.34	0.42
4:M:633:THR:OG1	4:M:634:SER:N	2.53	0.42
5:O:361:VAL:O	5:O:365:THR:HG22	2.19	0.42
5:O:617:ASN:HB3	5:O:618:PHE:CZ	2.54	0.42
5:O:743:ILE:HG12	5:O:752:LEU:HD13	2.00	0.42
5:O:1052:VAL:HG11	5:O:1066:GLU:CD	2.39	0.42
5:O:1052:VAL:O	5:O:1104:LEU:HB2	2.19	0.42
3:P:18:ASN:ND2	3:P:19:VAL:O	2.51	0.42
3:P:108:PRO:O	3:P:112:LEU:HG	2.20	0.42
3:P:205:HIS:CD2	3:P:206:ASP:HB2	2.54	0.42
3:P:375:TRP:HA	3:P:378:GLU:CD	2.39	0.42
6:X:29:GLN:HG3	6:X:30:GLU:OE2	2.18	0.42
6:X:211:ASP:N	6:X:211:ASP:OD1	2.51	0.42
6:Y:3:VAL:HA	6:Y:56:GLY:HA2	2.01	0.42
6:Z:158:ASP:OD1	6:Z:162:LYS:HG3	2.19	0.42
1:B:599:LEU:HA	1:B:832:GLN:CD	2.39	0.42
1:B:627:TRP:O	1:B:631:ILE:HG13	2.18	0.42
1:B:680:THR:OG1	1:B:683:THR:HG23	2.18	0.42
1:B:849:VAL:HA	1:B:998:ALA:O	2.18	0.42
1:B:1231:ARG:HH21	1:B:1250:VAL:HG13	1.83	0.42
2:C:437:ALA:HA	2:C:447:TRP:CE2	2.54	0.42
2:C:532:ILE:HA	2:C:535:VAL:HG22	1.99	0.42
2:C:866:THR:O	2:C:866:THR:HG22	2.18	0.42
2:C:982:MET:O	2:C:983:LEU:HG	2.19	0.42
2:C:1116:GLN:OE1	2:C:1116:GLN:HA	2.18	0.42
2:C:1257:VAL:HG12	2:C:1258:THR:O	2.19	0.42
3:D:162:SER:HB3	3:D:272:VAL:HG13	2.01	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:D:371:LYS:HA	3:D:371:LYS:HD3	1.95	0.42
4:K:346:GLN:CA	4:K:360:ASN:HA	2.43	0.42
4:K:360:ASN:OD1	4:K:361:LEU:N	2.40	0.42
4:K:519:ILE:O	4:K:521:SER:OG	2.29	0.42
4:K:523:THR:HG23	4:K:613:GLN:OE1	2.20	0.42
4:L:111:PHE:HD1	4:L:137:TYR:HE2	1.67	0.42
4:L:188:TRP:HD1	4:L:189:GLU:HG2	1.84	0.42
4:L:401:PHE:CZ	4:L:448:LEU:HB3	2.53	0.42
4:L:578:GLU:O	6:Z:70:HIS:NE2	2.51	0.42
4:M:68:SER:N	4:M:98:GLU:O	2.35	0.42
4:M:118:LEU:O	4:M:122:ARG:HG3	2.19	0.42
4:M:631:VAL:HG13	4:M:632:LYS:N	2.34	0.42
5:O:405:ALA:C	5:O:772:LEU:HD13	2.39	0.42
5:O:646:THR:HG22	5:O:647:ASN:N	2.33	0.42
5:O:760:PRO:O	5:O:763:ALA:HB3	2.20	0.42
5:O:1024:HIS:CD2	5:O:1024:HIS:N	2.86	0.42
5:O:1032:GLY:HA3	5:O:1041:SER:H	1.82	0.42
5:O:1080:PHE:HB2	5:O:1087:TRP:CZ3	2.55	0.42
3:P:86:TRP:CZ2	3:P:99:PRO:HB3	2.54	0.42
3:P:354:GLN:O	3:P:358:GLU:HG3	2.19	0.42
6:X:88:ALA:HB2	6:X:210:PHE:CE2	2.54	0.42
6:X:211:ASP:OD2	6:X:213:ARG:HB2	2.20	0.42
6:X:238:TYR:OH	6:X:242:LEU:HD11	2.20	0.42
6:Z:1:MET:SD	6:Z:71:HIS:HA	2.59	0.42
6:Z:192:PHE:HD2	6:Z:206:PHE:CG	2.37	0.42
1:B:267:LYS:O	1:B:304:ILE:HD12	2.20	0.42
1:B:331:ASN:O	1:B:334:LEU:HG	2.19	0.42
1:B:377:GLY:O	1:B:392:ARG:HA	2.19	0.42
1:B:713:TRP:CD2	1:B:714:PRO:HD2	2.55	0.42
1:B:793:MET:O	1:B:797:LEU:HG	2.18	0.42
1:B:803:ILE:C	1:B:804:LYS:HD2	2.40	0.42
1:B:1020:ILE:O	1:B:1024:VAL:HG23	2.18	0.42
1:B:1121:TYR:CE1	1:B:1125:GLN:HG3	2.54	0.42
1:B:1131:LYS:HE3	1:B:1160:TRP:CG	2.54	0.42
1:B:1231:ARG:HD3	1:B:1231:ARG:HA	1.74	0.42
2:C:460:SER:O	2:C:464:ARG:HG3	2.18	0.42
2:C:1077:PHE:CE2	2:C:1108:TRP:HB3	2.53	0.42
3:D:109:PRO:O	3:D:113:ALA:N	2.47	0.42
4:K:349:VAL:CG2	4:K:357:TRP:HB2	2.50	0.42
4:K:402:ILE:HB	4:K:469:LEU:HB3	2.01	0.42
4:K:504:THR:HA	4:K:508:ALA:O	2.19	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
4:L:34:LEU:O	4:L:36:LEU:HG	2.20	0.42
4:L:175:VAL:HA	4:L:178:ILE:HG12	2.01	0.42
4:L:250:MET:HG2	4:L:254:GLU:OE1	2.19	0.42
4:L:410:PHE:HB2	4:L:467:TYR:CE1	2.54	0.42
4:M:20:PHE:CD2	4:M:209:SER:HA	2.54	0.42
4:M:124:PHE:HE1	4:M:232:TYR:CZ	2.36	0.42
5:O:23:ARG:HD3	5:O:25:TYR:CE2	2.54	0.42
5:O:91:ARG:HA	5:O:94:LYS:HG3	2.00	0.42
5:O:93:LEU:HD22	5:O:123:PHE:CD1	2.55	0.42
5:O:241:MET:HG2	5:O:249:ILE:HD12	2.01	0.42
5:O:394:VAL:HG12	5:O:395:GLN:N	2.34	0.42
5:O:401:THR:HG21	5:O:724:ARG:HH21	1.84	0.42
5:O:826:LEU:HA	5:O:847:THR:HG23	2.01	0.42
5:O:1175:SER:HA	5:O:1181:LEU:HA	2.01	0.42
3:P:84:HIS:CG	3:P:130:TYR:HD1	2.36	0.42
3:P:108:PRO:HD2	3:P:111:VAL:HB	2.02	0.42
6:X:142:PHE:CE2	6:X:266:GLY:HA2	2.50	0.42
6:X:218:TRP:HE1	6:X:358:ILE:HD11	1.83	0.42
6:Y:38:ALA:O	6:Y:40:PRO:HD3	2.20	0.42
6:Z:33:ASP:OD1	6:Z:34:LYS:N	2.52	0.42
6:Z:107:HIS:ND1	6:Z:170:LEU:HB2	2.35	0.42
6:Z:251:HIS:CE1	6:Z:254:LEU:HD12	2.54	0.42
6:Z:253:GLY:HA2	6:Z:275:MET:HB2	2.01	0.42
6:Z:256:HIS:HB2	6:Z:342:MET:CE	2.49	0.42
6:Z:326:ARG:CZ	6:Z:329:GLN:HE22	2.33	0.42
1:B:427:VAL:H	1:B:1235:LEU:HD12	1.84	0.42
1:B:432:GLY:O	1:B:451:PHE:CG	2.72	0.42
1:B:512:ALA:O	1:B:516:ILE:HG12	2.19	0.42
1:B:522:ILE:HA	1:B:525:ILE:HD11	2.01	0.42
1:B:571:LEU:HD23	1:B:571:LEU:HA	1.83	0.42
1:B:630:PHE:CE1	1:B:768:VAL:HB	2.54	0.42
1:B:767:ARG:O	1:B:770:GLU:HG3	2.19	0.42
1:B:851:ARG:HH11	1:B:986:PRO:HA	1.83	0.42
1:B:1086:GLY:HA2	1:B:1092:PRO:HG3	2.02	0.42
2:C:678:PHE:N	2:C:678:PHE:CD1	2.83	0.42
2:C:816:PRO:HA	2:C:819:LEU:HD11	2.02	0.42
4:K:205:ILE:HD12	4:K:205:ILE:HA	1.88	0.42
4:K:376:MET:CA	4:K:498:LEU:HG	2.49	0.42
4:K:514:VAL:O	4:K:518:LEU:HG	2.19	0.42
4:L:292:LEU:HD21	4:L:545:LYS:O	2.19	0.42
4:M:176:ASP:HA	4:M:179:GLN:CG	2.50	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
4:M:535:ALA:O	4:M:539:MET:HG3	2.19	0.42
5:O:514:TYR:CD1	5:O:575:TYR:CG	3.08	0.42
5:O:1044:ILE:CD1	5:O:1089:LEU:HG	2.50	0.42
5:O:1058:ASN:O	5:O:1059:SER:OG	2.33	0.42
3:P:27:SER:O	3:P:31:ARG:N	2.53	0.42
3:P:130:TYR:HA	3:P:131:PRO:HD3	1.86	0.42
3:P:186:THR:O	3:P:190:ILE:HD12	2.19	0.42
6:Y:3:VAL:HA	6:Y:56:GLY:CA	2.50	0.42
6:Y:82:TYR:HB3	6:Y:298:LEU:HD11	2.01	0.42
6:Y:188:PRO:HA	6:Y:191:ALA:HB3	2.01	0.42
6:Y:266:GLY:O	6:Y:267:LYS:HG2	2.18	0.42
6:Z:71:HIS:CG	6:Z:72:ARG:N	2.87	0.42
1:B:800:LEU:O	1:B:803:ILE:HG12	2.20	0.42
1:B:1046:ILE:HG13	1:B:1138:ARG:HA	2.02	0.42
1:B:1204:TYR:CE2	1:B:1207:ARG:HG3	2.54	0.42
2:C:576:SER:HB2	2:C:627:TRP:CH2	2.55	0.42
2:C:602:GLY:CA	2:C:833:VAL:HG23	2.49	0.42
2:C:763:ASN:HA	2:C:766:ALA:HB3	2.01	0.42
4:K:136:LYS:HZ1	4:L:158:ASN:HA	1.85	0.42
4:K:426:ASN:CG	6:Z:61:LEU:HB2	2.40	0.42
4:L:178:ILE:O	4:L:181:ALA:HB3	2.19	0.42
4:L:198:LEU:O	4:L:199:LEU:HD23	2.20	0.42
4:L:375:PRO:HA	4:L:452:TYR:CD2	2.55	0.42
4:L:380:LEU:HB2	4:L:401:PHE:HZ	1.85	0.42
4:L:385:LYS:NZ	4:L:492:LEU:HG	2.35	0.42
4:L:621:LYS:HG3	4:L:622:ASN:N	2.35	0.42
4:M:147:TYR:CZ	4:M:159:PHE:HB2	2.54	0.42
4:M:348:GLN:HA	4:M:359:PHE:CE2	2.55	0.42
4:M:370:LEU:HB3	4:M:452:TYR:CE1	2.55	0.42
4:M:395:ASN:OD1	4:M:432:ALA:N	2.31	0.42
4:M:476:ALA:HB2	6:X:160:ASP:OD2	2.20	0.42
4:M:525:GLU:HA	4:M:528:ASN:HD22	1.84	0.42
5:O:73:TYR:HE1	5:O:170:ARG:HH12	1.67	0.42
5:O:223:HIS:NE2	5:O:281:GLN:HA	2.34	0.42
5:O:808:THR:O	5:O:811:TYR:HB2	2.19	0.42
5:O:810:MET:HG2	5:O:990:TRP:CE2	2.55	0.42
3:P:77:GLN:HG3	3:P:91:PHE:CE1	2.54	0.42
6:X:61:LEU:HD12	6:X:62:GLN:N	2.34	0.42
6:X:322:GLY:HA2	6:X:325:ASN:ND2	2.34	0.42
6:Y:65:LEU:HD12	6:Y:66:LYS:N	2.35	0.42
6:Y:86:GLN:HA	6:Y:89:ASP:OD2	2.20	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
6:Y:144:SER:HB2	6:Y:146:HIS:ND1	2.34	0.42
6:Y:169:ALA:O	6:Y:173:MET:HG3	2.20	0.42
6:Y:182:ASN:ND2	6:Y:362:MET:HG3	2.35	0.42
6:Z:39:GLN:HE22	6:Z:79:HIS:CD2	2.32	0.42
1:B:613:TYR:CD2	1:B:615:LYS:HD3	2.54	0.42
1:B:970:ASN:HA	1:B:973:MET:HE3	2.02	0.42
2:C:418:LYS:NZ	2:C:1211:CYS:SG	2.67	0.42
2:C:536:LEU:HD11	2:C:596:ALA:HB1	2.02	0.42
2:C:713:TRP:CD1	2:C:837:ARG:NH2	2.88	0.42
2:C:982:MET:SD	2:C:983:LEU:N	2.93	0.42
2:C:1003:GLN:CG	2:C:1007:ARG:HB3	2.49	0.42
3:D:91:PHE:C	3:D:93:ARG:H	2.21	0.42
3:D:93:ARG:O	3:D:106:VAL:HA	2.19	0.42
3:D:239:PHE:HA	3:D:252:ILE:O	2.19	0.42
4:K:283:SER:HA	4:K:640:LEU:HD21	2.02	0.42
4:K:337:ILE:HG13	4:K:410:PHE:CZ	2.55	0.42
4:K:620:ASP:HB3	4:K:623:ASN:ND2	2.35	0.42
4:K:620:ASP:HA	6:X:2:GLU:CD	2.40	0.42
4:L:334:LEU:HD22	4:L:366:ARG:NH2	2.35	0.42
4:L:381:ASP:HA	4:L:446:THR:OG1	2.19	0.42
4:L:429:GLN:C	4:L:430:LEU:HD22	2.40	0.42
4:M:46:VAL:O	4:M:48:TRP:NE1	2.52	0.42
4:M:150:VAL:N	4:M:154:GLN:OE1	2.36	0.42
4:M:186:GLU:O	4:M:189:GLU:HG3	2.20	0.42
5:O:78:ALA:HB3	5:O:81:ASP:CG	2.40	0.42
5:O:200:TYR:CD2	5:O:232:HIS:HB3	2.54	0.42
5:O:298:THR:HA	5:O:301:LEU:HG	2.01	0.42
5:O:1022:ASP:OD2	5:O:1024:HIS:HB2	2.19	0.42
5:O:1096:ALA:HB1	5:O:1119:VAL:O	2.20	0.42
5:O:1247:ILE:HD13	5:O:1247:ILE:HA	1.80	0.42
3:P:3:ARG:NH2	3:P:300:GLN:HA	2.35	0.42
3:P:304:ILE:HG23	3:P:305:ASP:O	2.20	0.42
6:X:119:ASP:O	6:X:122:ARG:HB2	2.20	0.42
6:X:314:ALA:C	6:X:315:LEU:HD22	2.39	0.42
6:X:348:LYS:HE2	6:X:348:LYS:HB2	1.72	0.42
6:Y:240:LYS:HG2	6:Y:247:ARG:HH12	1.84	0.42
1:B:268:ILE:HG13	1:B:303:ARG:O	2.20	0.42
1:B:772:MET:HB2	1:B:797:LEU:HD13	2.00	0.42
2:C:756:ASP:O	2:C:762:THR:HG21	2.19	0.42
2:C:924:THR:O	2:C:928:VAL:HG13	2.19	0.42
2:C:937:GLN:O	2:C:947:ARG:HD2	2.20	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:D:142:ARG:HA	3:D:145:GLN:OE1	2.19	0.42
3:D:236:THR:HG22	3:D:258:ASN:O	2.20	0.42
3:D:375:TRP:N	3:D:375:TRP:CD1	2.86	0.42
3:D:379:ASP:OD2	3:D:383:ASN:ND2	2.51	0.42
4:K:146:CYS:HA	4:K:159:PHE:CZ	2.55	0.42
4:K:621:LYS:HZ2	4:K:625:ILE:HD11	1.84	0.42
4:L:34:LEU:HD13	4:L:110:ASN:HA	2.01	0.42
4:L:587:ASN:HA	4:L:588:PRO:HD3	1.92	0.42
4:M:297:GLU:HB2	4:M:542:ARG:HH12	1.85	0.42
4:M:413:TRP:CH2	4:M:467:TYR:HD1	2.37	0.42
4:M:428:VAL:HG12	4:M:429:GLN:O	2.19	0.42
4:M:456:GLN:O	4:M:457:LEU:HD23	2.20	0.42
5:O:133:LEU:O	5:O:136:LEU:HD23	2.20	0.42
5:O:501:THR:HG23	5:O:503:LYS:HG2	2.00	0.42
5:O:930:THR:OG1	5:O:1015:TYR:HA	2.19	0.42
5:O:1154:MET:HB2	5:O:1201:VAL:CG2	2.47	0.42
6:Y:241:GLU:O	6:Y:242:LEU:HD23	2.20	0.42
6:Z:7:ASN:HB3	6:Z:10:GLN:HE22	1.84	0.42
1:B:352:LEU:HD13	1:B:955:LEU:HD23	2.01	0.42
1:B:386:THR:OG1	1:B:388:GLY:N	2.51	0.42
1:B:589:SER:CB	1:B:592:ARG:HH22	2.33	0.42
1:B:602:GLY:HA2	1:B:833:VAL:CG1	2.42	0.42
1:B:663:PRO:HG2	1:B:693:LEU:HB3	2.00	0.42
1:B:758:THR:O	1:B:763:ASN:ND2	2.42	0.42
1:B:1000:GLN:OE1	1:B:1008:THR:HB	2.19	0.42
2:C:801:LYS:CG	2:C:802:LEU:HG	2.47	0.42
2:C:1116:GLN:OE1	2:C:1119:THR:OG1	2.27	0.42
2:C:1158:ASN:CG	2:C:1160:TRP:H	2.23	0.42
4:K:333:TRP:CD2	4:K:531:LEU:HD21	2.53	0.42
4:L:21:LYS:HA	4:L:209:SER:HA	2.02	0.42
4:L:101:VAL:CG1	4:L:166:ILE:HG13	2.50	0.42
4:L:348:GLN:HA	4:L:357:TRP:O	2.19	0.42
4:M:130:VAL:HG12	4:M:131:LEU:HB2	2.01	0.42
4:M:332:THR:C	4:M:335:ARG:HH12	2.23	0.42
4:M:651:LEU:O	4:M:655:GLU:HG3	2.20	0.42
4:M:662:GLN:OE1	4:M:665:LEU:HB2	2.20	0.42
5:O:134:PRO:O	5:O:137:ILE:HG12	2.19	0.42
5:O:538:ILE:HG12	5:O:541:VAL:CG2	2.50	0.42
5:O:826:LEU:HD13	5:O:828:LEU:HD21	2.02	0.42
5:O:931:ASP:OD2	5:O:932:VAL:N	2.50	0.42
5:O:1136:PHE:CG	5:O:1228:LEU:HD22	2.55	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:P:134:ALA:HA	3:P:140:LYS:HD3	2.02	0.42
6:X:82:TYR:HD2	6:X:86:GLN:NE2	2.16	0.42
6:Z:60:SER:H	6:Z:63:ARG:HD2	1.85	0.42
1:B:378:PHE:CZ	1:B:433:ARG:HB2	2.54	0.42
1:B:694:ILE:O	1:B:703:ARG:NH1	2.53	0.42
1:B:793:MET:HA	1:B:796:THR:OG1	2.20	0.42
1:B:898:ASN:N	1:B:898:ASN:OD1	2.53	0.42
1:B:946:ASP:HB3	3:P:31:ARG:NH1	2.35	0.42
1:B:1116:GLN:HG2	1:B:1170:ILE:HG13	2.01	0.42
2:C:435:ASP:HB3	2:C:449:ASP:OD2	2.20	0.42
2:C:722:GLY:HA2	2:C:729:SER:OG	2.19	0.42
3:D:393:ARG:O	3:D:394:THR:OG1	2.29	0.42
4:K:64:ARG:HH22	4:K:67:THR:H	1.68	0.42
4:K:232:TYR:HB3	4:K:234:LYS:HE2	2.02	0.42
4:K:264:ALA:HA	4:K:267:ALA:HB3	2.00	0.42
4:K:660:TRP:CZ2	4:M:622:ASN:HB3	2.55	0.42
4:K:661:THR:O	4:K:664:PHE:HB3	2.20	0.42
4:L:193:ARG:HH21	4:L:197:THR:N	2.17	0.42
4:L:302:ALA:HA	4:L:624:TRP:HE1	1.85	0.42
4:L:380:LEU:HD22	4:L:401:PHE:CE1	2.55	0.42
4:M:290:MET:CE	4:M:636:ARG:HG2	2.50	0.42
5:O:396:TRP:CZ3	5:O:398:PRO:HA	2.54	0.42
5:O:940:LEU:HD13	5:O:951:PHE:CE2	2.54	0.42
5:O:1023:ILE:H	5:O:1023:ILE:HD12	1.84	0.42
3:P:90:ARG:HE	3:P:116:ALA:N	2.18	0.42
3:P:96:TRP:CE3	3:P:104:LEU:HD21	2.55	0.42
3:P:100:THR:OG1	3:P:133:LEU:HD11	2.19	0.42
3:P:161:ILE:HG21	3:P:271:PHE:HB3	2.02	0.42
3:P:171:TRP:HA	3:P:171:TRP:CE3	2.55	0.42
3:P:241:CYS:HB3	3:P:251:TRP:NE1	2.35	0.42
3:P:249:GLU:HB3	3:P:251:TRP:CD1	2.55	0.42
3:P:275:THR:HG1	3:P:330:PHE:C	2.23	0.42
6:X:5:LEU:HD13	6:X:310:LYS:NZ	2.34	0.42
6:X:221:MET:CB	6:X:356:VAL:HG12	2.44	0.42
6:X:227:GLU:HA	6:X:230:HIS:CD2	2.55	0.42
6:Y:102:PHE:HA	6:Y:105:GLN:CD	2.40	0.42
6:Y:105:GLN:NE2	6:Y:136:VAL:HG13	2.34	0.42
6:Y:163:LEU:HD21	6:Y:272:PHE:CE2	2.55	0.42
6:Z:24:SER:HB3	6:Z:44:VAL:HB	2.01	0.42
1:B:436:ARG:HB3	1:B:448:VAL:HG13	2.01	0.41
1:B:439:MET:HG2	1:B:440:MET:H	1.84	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:B:716:PRO:HB3	1:B:742:PRO:HA	2.02	0.41
1:B:1144:TYR:CE2	1:B:1146:TYR:HD1	2.37	0.41
2:C:936:ALA:HB2	2:C:941:THR:HB	2.02	0.41
3:D:96:TRP:CE3	3:D:104:LEU:HD11	2.55	0.41
4:K:48:TRP:CA	4:K:66:MET:H	2.20	0.41
4:K:118:LEU:HA	4:K:121:ASN:ND2	2.35	0.41
4:K:503:VAL:HG11	4:K:535:ALA:HB2	2.02	0.41
4:K:547:ALA:O	4:K:551:LYS:HG3	2.20	0.41
4:L:16:ASP:CG	4:L:17:GLY:H	2.24	0.41
4:M:42:ASN:CG	4:M:43:PRO:HD2	2.40	0.41
4:M:322:ASN:OD1	4:M:324:ARG:HG2	2.20	0.41
4:M:327:LYS:NZ	4:M:329:ASP:H	2.15	0.41
5:O:392:SER:O	5:O:394:VAL:HG23	2.20	0.41
5:O:570:ASP:HA	5:O:605:ALA:O	2.20	0.41
5:O:616:ILE:HD12	5:O:652:PHE:HA	2.01	0.41
5:O:679:TYR:O	5:O:683:THR:HG23	2.20	0.41
5:O:779:SER:OG	5:O:780:LEU:HD22	2.20	0.41
3:P:172:SER:O	3:P:175:VAL:HB	2.20	0.41
3:P:183:PHE:O	3:P:185:HIS:CE1	2.73	0.41
3:P:351:GLN:HG3	3:P:355:ASN:HD21	1.85	0.41
6:X:79:HIS:CD2	6:X:82:TYR:CZ	3.08	0.41
6:Y:107:HIS:ND1	6:Y:166:TYR:HB3	2.35	0.41
6:Z:312:ARG:HA	6:Z:316:GLY:O	2.20	0.41
1:B:527:ASN:H	1:B:871:VAL:HG11	1.84	0.41
1:B:691:ILE:HB	1:B:703:ARG:HD2	2.00	0.41
1:B:746:GLN:NE2	5:O:154:SER:O	2.28	0.41
1:B:1153:PRO:HG3	1:B:1184:ILE:O	2.20	0.41
1:B:1161:ASN:OD1	1:B:1162:LEU:N	2.45	0.41
2:C:338:LEU:CD2	2:C:339:LEU:HB2	2.50	0.41
2:C:371:TYR:HB3	2:C:372:LEU:H	1.69	0.41
2:C:453:THR:HG23	2:C:1253:LEU:O	2.20	0.41
2:C:518:GLN:HA	2:C:521:ARG:NE	2.36	0.41
2:C:549:LEU:HD13	2:C:891:TYR:CE2	2.55	0.41
2:C:581:LEU:O	2:C:582:ARG:HG2	2.19	0.41
2:C:702:LEU:O	2:C:705:TRP:HB3	2.20	0.41
2:C:815:ALA:N	2:C:816:PRO:HD2	2.35	0.41
2:C:1003:GLN:HB2	2:C:1006:GLY:C	2.40	0.41
3:D:43:LEU:HA	3:D:46:ILE:HG13	2.02	0.41
3:D:130:TYR:HA	3:D:131:PRO:HD3	1.94	0.41
3:D:163:TYR:HE1	3:D:269:SER:HB3	1.85	0.41
3:D:270:LEU:HD22	3:D:417:LEU:CD2	2.50	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:D:413:ALA:O	3:D:416:ALA:HB3	2.20	0.41
4:K:45:GLY:O	4:K:102:VAL:N	2.36	0.41
4:K:113:LYS:HA	4:K:116:MET:HG2	2.01	0.41
4:K:329:ASP:OD1	4:K:329:ASP:N	2.50	0.41
4:K:398:LYS:HE2	4:K:427:TYR:OH	2.20	0.41
4:K:469:LEU:HD12	4:K:470:ALA:H	1.85	0.41
4:K:522:TYR:O	4:K:612:THR:HA	2.20	0.41
4:L:13:VAL:HG13	4:L:14:THR:HG23	2.01	0.41
4:L:20:PHE:CD2	4:L:22:PRO:HD3	2.55	0.41
4:L:67:THR:HB	4:L:70:ASP:OD1	2.19	0.41
4:L:229:ILE:HG13	4:L:233:PRO:CA	2.50	0.41
4:L:280:GLU:CG	4:L:284:LYS:HE2	2.46	0.41
4:L:284:LYS:NZ	4:M:36:LEU:HB2	2.35	0.41
4:L:392:TRP:HD1	4:L:394:PRO:HD3	1.84	0.41
4:L:537:ARG:NH2	4:L:608:SER:OG	2.53	0.41
5:O:164:LEU:HD12	5:O:164:LEU:HA	1.75	0.41
5:O:417:LEU:O	5:O:709:ILE:HB	2.19	0.41
5:O:995:LEU:C	5:O:998:SER:H	2.22	0.41
3:P:3:ARG:NH2	3:P:296:CYS:HA	2.36	0.41
3:P:34:ASN:ND2	3:P:103:GLY:HA2	2.35	0.41
3:P:42:PHE:HD1	3:P:64:ARG:HE	1.66	0.41
6:X:118:LEU:HD12	6:X:118:LEU:HA	1.90	0.41
6:Z:26:TYR:CG	6:Z:27:SER:N	2.88	0.41
6:Z:34:LYS:H	6:Z:34:LYS:HG2	1.54	0.41
1:B:351:PRO:HA	1:B:354:PHE:CE2	2.56	0.41
1:B:370:LEU:HD21	1:B:465:TRP:CZ3	2.56	0.41
1:B:590:SER:H	1:B:592:ARG:NH2	2.19	0.41
1:B:610:ASP:HA	1:B:613:TYR:CZ	2.54	0.41
1:B:652:LEU:H	1:B:652:LEU:HG	1.41	0.41
1:B:685:PRO:HB2	1:B:687:CYS:SG	2.60	0.41
2:C:329:GLU:O	2:C:331:ASN:ND2	2.50	0.41
2:C:355:ARG:NH1	2:C:950:ASP:OD1	2.53	0.41
2:C:417:SER:O	2:C:420:CYS:N	2.53	0.41
2:C:509:ILE:HD12	2:C:513:GLU:CD	2.41	0.41
2:C:551:ILE:HA	2:C:889:GLY:HA2	2.03	0.41
2:C:731:ASN:N	2:C:736:PRO:HB3	2.35	0.41
2:C:740:LEU:HD23	2:C:740:LEU:HA	1.89	0.41
2:C:1039:GLY:O	2:C:1145:PRO:HA	2.19	0.41
4:K:171:GLN:HG2	4:K:174:TYR:CD2	2.55	0.41
4:K:303:SER:HB2	4:K:307:VAL:HG11	2.03	0.41
4:K:331:ALA:HA	4:K:333:TRP:NE1	2.35	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
4:K:493:LEU:O	4:K:493:LEU:HD12	2.19	0.41
4:K:566:VAL:N	4:K:567:PRO:HD2	2.35	0.41
4:K:651:LEU:O	4:K:655:GLU:HG2	2.20	0.41
4:K:656:SER:HA	4:K:659:ASN:HD22	1.84	0.41
4:L:188:TRP:CZ3	4:L:228:LEU:HD13	2.56	0.41
4:L:228:LEU:O	4:L:231:ARG:N	2.42	0.41
4:M:136:LYS:HD3	4:M:136:LYS:HA	1.80	0.41
4:M:397:LYS:N	4:M:430:LEU:HB2	2.35	0.41
5:O:61:ARG:HB2	5:O:64:GLN:NE2	2.35	0.41
5:O:96:GLU:O	5:O:99:ARG:HB3	2.20	0.41
5:O:480:ASP:OD1	5:O:481:ARG:HD3	2.21	0.41
5:O:504:GLU:HG3	5:O:507:ARG:NH1	2.25	0.41
5:O:537:LYS:HA	5:O:537:LYS:HD3	1.78	0.41
5:O:647:ASN:HB3	5:O:681:TYR:HE1	1.85	0.41
5:O:778:ARG:HD2	5:O:781:GLU:OE1	2.21	0.41
5:O:944:SER:OG	5:O:947:LYS:NZ	2.46	0.41
5:O:1004:ALA:HA	5:O:1007:ARG:CZ	2.50	0.41
5:O:1044:ILE:HD11	5:O:1089:LEU:HG	2.01	0.41
3:P:130:TYR:HE2	3:P:132:PHE:HD1	1.68	0.41
3:P:145:GLN:HG3	3:P:297:SER:HB3	2.02	0.41
3:P:211:ARG:HA	3:P:214:GLN:CD	2.40	0.41
6:Z:17:ASN:HA	6:Z:20:GLU:CG	2.50	0.41
6:Z:196:ARG:HE	6:Z:352:LEU:CB	2.29	0.41
1:B:285:PHE:HB3	1:B:286:PHE:CE2	2.56	0.41
1:B:338:LEU:HG	1:B:968:TRP:CE2	2.55	0.41
1:B:483:LEU:HD13	1:B:493:VAL:HG13	2.02	0.41
1:B:530:THR:HA	1:B:533:GLN:NE2	2.34	0.41
1:B:584:SER:O	1:B:586:SER:HA	2.21	0.41
1:B:623:LEU:HG	1:B:627:TRP:CE2	2.55	0.41
1:B:623:LEU:HD21	1:B:627:TRP:CZ2	2.56	0.41
1:B:904:ALA:O	1:B:907:PRO:HD2	2.21	0.41
1:B:1049:ARG:HH11	1:B:1134:GLU:CD	2.21	0.41
1:B:1232:TYR:O	1:B:1236:THR:HG23	2.20	0.41
2:C:337:GLN:NE2	2:C:357:ASN:HB3	2.36	0.41
2:C:857:ILE:HG12	3:P:348:VAL:O	2.20	0.41
2:C:1165:ALA:HA	2:C:1168:GLU:OE1	2.20	0.41
3:D:10:THR:HA	3:D:154:ASN:ND2	2.36	0.41
3:D:251:TRP:N	3:D:251:TRP:CD1	2.87	0.41
3:D:283:VAL:HG22	3:D:287:LEU:CG	2.50	0.41
4:K:101:VAL:O	4:K:165:VAL:HG13	2.21	0.41
4:K:193:ARG:HH21	4:M:634:SER:C	2.23	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
4:K:228:LEU:HD23	4:K:231:ARG:CZ	2.50	0.41
4:K:337:ILE:HG22	4:K:364:GLY:N	2.35	0.41
4:K:401:PHE:HE1	4:K:448:LEU:HG	1.85	0.41
4:K:667:LYS:HE3	4:K:667:LYS:HB2	1.82	0.41
4:L:332:THR:HB	4:L:507:GLY:HA3	2.02	0.41
4:M:584:ARG:HG3	4:M:584:ARG:HH11	1.84	0.41
5:O:806:CYS:O	5:O:810:MET:HG3	2.20	0.41
5:O:888:CYS:HB3	5:O:924:VAL:HG23	2.03	0.41
5:O:1172:GLN:HG2	5:O:1174:PHE:HD2	1.85	0.41
5:O:1269:SER:HB2	5:O:1274:ILE:HB	2.02	0.41
3:P:41:GLN:HG3	3:P:64:ARG:NH2	2.34	0.41
3:P:174:ASP:HA	3:P:177:GLN:OE1	2.19	0.41
3:P:266:GLY:O	3:P:268:ARG:NH2	2.53	0.41
6:X:19:PHE:CE2	6:X:291:LYS:HB3	2.56	0.41
6:Z:8:GLY:N	6:Z:315:LEU:HD13	2.35	0.41
6:Z:257:TYR:HB3	6:Z:343:PHE:HA	2.00	0.41
1:B:303:ARG:HH11	1:B:1208:SER:HB2	1.85	0.41
1:B:441:ASN:N	2:C:862:LEU:O	2.31	0.41
1:B:514:ARG:NH1	1:B:727:PHE:O	2.53	0.41
1:B:601:ASN:O	1:B:603:VAL:N	2.54	0.41
1:B:707:GLU:HG2	1:B:711:ARG:NE	2.35	0.41
1:B:708:ILE:HG13	1:B:771:LEU:HD21	2.02	0.41
1:B:1112:LEU:O	1:B:1115:TRP:HB3	2.21	0.41
1:B:1231:ARG:HD3	1:B:1252:ASP:HA	2.02	0.41
2:C:355:ARG:HE	2:C:952:ILE:HB	1.85	0.41
2:C:360:HIS:CD2	2:C:969:VAL:HG22	2.55	0.41
3:D:22:ASN:OD1	3:D:23:ASP:N	2.53	0.41
3:D:94:LEU:HA	3:D:105:VAL:C	2.40	0.41
3:D:306:ASN:O	3:D:314:ARG:NH1	2.54	0.41
4:K:23:SER:OG	4:K:24:ALA:N	2.54	0.41
4:K:51:ILE:HG12	4:K:62:ALA:O	2.21	0.41
4:K:240:LEU:HB3	4:K:244:ASN:ND2	2.36	0.41
4:K:461:ASP:C	4:K:463:GLU:H	2.24	0.41
4:K:574:ILE:HA	4:K:577:LEU:HD12	2.01	0.41
4:K:671:HIS:CE1	4:M:578:GLU:HB2	2.51	0.41
4:L:47:PRO:CG	4:L:100:LEU:HB2	2.51	0.41
4:L:290:MET:O	4:L:290:MET:HE3	2.21	0.41
4:L:471:THR:HG23	4:L:472:PHE:O	2.20	0.41
4:L:603:PHE:CE1	4:L:607:PRO:HB3	2.54	0.41
4:M:25:GLU:OE2	4:M:26:THR:HG23	2.20	0.41
4:M:176:ASP:HA	4:M:179:GLN:HG3	2.02	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
4:M:284:LYS:HD2	4:M:287:GLU:OE1	2.20	0.41
4:M:403:VAL:HG22	4:M:404:PHE:N	2.35	0.41
4:M:506:LYS:HG2	4:M:522:TYR:CE2	2.56	0.41
4:M:536:ALA:O	4:M:540:ILE:HG12	2.20	0.41
4:M:625:ILE:C	4:M:629:GLN:HE21	2.20	0.41
5:O:202:ASP:C	5:O:231:HIS:HB2	2.41	0.41
5:O:405:ALA:O	5:O:772:LEU:HD13	2.20	0.41
5:O:457:PRO:C	5:O:459:THR:H	2.24	0.41
5:O:469:ARG:HD2	5:O:647:ASN:O	2.21	0.41
5:O:809:MET:SD	5:O:812:ASN:HB2	2.61	0.41
3:P:3:ARG:HH22	3:P:301:LEU:HG	1.84	0.41
6:X:272:PHE:HA	6:X:281:LYS:O	2.20	0.41
6:Z:14:LEU:HD13	6:Z:45:CYS:SG	2.60	0.41
1:B:268:ILE:HA	1:B:304:ILE:HA	2.02	0.41
1:B:623:LEU:HD13	1:B:626:LEU:HD12	2.02	0.41
1:B:637:PRO:HB3	1:B:714:PRO:HG2	2.02	0.41
1:B:1083:ILE:HG22	1:B:1084:SER:O	2.20	0.41
2:C:528:ASN:HB3	2:C:530:THR:HG22	2.02	0.41
2:C:854:ARG:NH2	3:P:258:ASN:HA	2.36	0.41
2:C:928:VAL:HG12	2:C:984:LEU:CD2	2.50	0.41
2:C:1116:GLN:O	2:C:1172:PRO:HG3	2.21	0.41
4:K:49:ILE:HD12	4:K:49:ILE:HA	1.87	0.41
4:K:375:PRO:CG	4:K:454:PRO:HD3	2.49	0.41
4:L:24:ALA:N	4:L:203:VAL:HB	2.35	0.41
4:L:328:ILE:HD11	4:L:366:ARG:NH2	2.35	0.41
4:L:380:LEU:HB2	4:L:401:PHE:CZ	2.55	0.41
4:L:401:PHE:CE1	4:L:448:LEU:HD23	2.55	0.41
4:L:409:PRO:O	4:L:413:TRP:CD1	2.73	0.41
4:L:437:PHE:HB3	4:L:440:GLN:HG2	2.02	0.41
4:L:506:LYS:HD3	6:Z:316:GLY:HA2	2.03	0.41
4:M:201:THR:O	4:M:202:ASN:ND2	2.53	0.41
4:M:645:LYS:NZ	4:M:649:SER:HB3	2.35	0.41
5:O:44:LYS:HB2	5:O:60:PHE:CZ	2.55	0.41
5:O:61:ARG:NH2	5:O:169:GLY:O	2.54	0.41
5:O:215:ASP:OD2	5:O:235:LEU:HD21	2.20	0.41
5:O:347:GLN:OE1	5:O:349:ARG:HB3	2.20	0.41
5:O:449:LEU:HD12	5:O:450:SER:H	1.84	0.41
5:O:530:ILE:HG23	5:O:533:TRP:HE3	1.85	0.41
5:O:1234:ASP:OD1	5:O:1236:ARG:HB2	2.21	0.41
3:P:7:LEU:HA	3:P:127:CYS:H	1.85	0.41
3:P:43:LEU:O	3:P:46:ILE:HG23	2.20	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:P:132:PHE:HA	3:P:135:ARG:HD2	2.03	0.41
3:P:208:THR:O	3:P:212:MET:HG3	2.21	0.41
3:P:379:ASP:O	3:P:382:TYR:HB3	2.21	0.41
6:X:64:LYS:HB3	6:X:66:LYS:HZ1	1.85	0.41
6:Y:49:VAL:O	6:Y:57:VAL:HA	2.20	0.41
6:Y:163:LEU:HD21	6:Y:272:PHE:HE2	1.84	0.41
6:Z:25:ILE:HD13	6:Z:34:LYS:HE2	2.03	0.41
1:B:377:GLY:CA	1:B:393:SER:O	2.69	0.41
1:B:471:ARG:HE	1:B:471:ARG:HA	1.85	0.41
1:B:1058:PRO:O	1:B:1200:ILE:HG13	2.20	0.41
2:C:549:LEU:HD13	2:C:891:TYR:CE1	2.56	0.41
2:C:600:TYR:CZ	2:C:830:PRO:HA	2.55	0.41
2:C:610:ASP:HA	2:C:613:TYR:CD2	2.56	0.41
2:C:689:MET:HG3	2:C:840:ARG:NH2	2.36	0.41
2:C:791:SER:O	2:C:794:ARG:HB3	2.20	0.41
2:C:1154:ARG:HB3	2:C:1190:ILE:HD12	2.01	0.41
3:D:220:PHE:HE1	3:D:251:TRP:CD1	2.38	0.41
3:D:279:TRP:HB3	3:D:285:LEU:HD22	2.02	0.41
4:K:337:ILE:HG22	4:K:364:GLY:H	1.85	0.41
4:L:218:LEU:HD12	4:L:218:LEU:HA	1.81	0.41
4:L:525:GLU:O	4:L:528:ASN:HB3	2.20	0.41
4:L:621:LYS:HZ2	4:L:622:ASN:HD21	1.69	0.41
4:M:137:TYR:CD1	4:M:140:LEU:HD22	2.55	0.41
4:M:327:LYS:NZ	4:M:329:ASP:OD1	2.46	0.41
4:M:405:GLN:O	4:M:421:GLN:N	2.49	0.41
4:M:488:VAL:O	4:M:489:TRP:CD1	2.73	0.41
5:O:72:LEU:HD21	5:O:127:GLN:NE2	2.35	0.41
5:O:237:THR:OG1	5:O:238:LEU:N	2.53	0.41
5:O:624:TRP:HA	5:O:627:ILE:HB	2.02	0.41
5:O:804:HIS:CD2	5:O:804:HIS:N	2.87	0.41
5:O:943:ASP:OD1	5:O:945:THR:OG1	2.30	0.41
3:P:123:ARG:HB3	3:P:125:TYR:OH	2.20	0.41
3:P:139:PHE:O	3:P:143:VAL:HG23	2.20	0.41
1:B:308:THR:HG23	1:B:323:ILE:HG12	2.02	0.41
1:B:793:MET:O	1:B:796:THR:HB	2.20	0.41
1:B:1000:GLN:HB3	1:B:1010:ASN:HB2	2.03	0.41
1:B:1135:LEU:HD21	1:B:1137:ILE:HD11	2.02	0.41
2:C:352:LEU:HB3	2:C:955:LEU:HB2	2.02	0.41
2:C:355:ARG:NE	2:C:952:ILE:HB	2.36	0.41
2:C:371:TYR:N	2:C:462:ARG:CZ	2.84	0.41
2:C:650:MET:O	2:C:654:ASN:ND2	2.54	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:C:713:TRP:CD1	2:C:838:LEU:HD21	2.56	0.41
2:C:805:SER:O	2:C:808:PRO:HD2	2.21	0.41
3:D:130:TYR:CD1	3:D:131:PRO:HD2	2.55	0.41
3:D:300:GLN:NE2	3:D:301:LEU:O	2.54	0.41
3:D:316:ILE:O	3:D:319:TRP:N	2.53	0.41
4:K:409:PRO:HD2	4:K:412:LEU:HD12	2.03	0.41
4:K:471:THR:HG22	4:K:488:VAL:H	1.86	0.41
4:K:503:VAL:O	4:K:510:VAL:N	2.48	0.41
4:K:657:ILE:HD11	4:M:625:ILE:HG23	2.03	0.41
4:L:33:SER:O	4:M:39:GLY:HA3	2.20	0.41
4:L:47:PRO:HA	4:L:148:VAL:HA	2.03	0.41
4:L:47:PRO:HG2	4:L:66:MET:HB3	2.02	0.41
4:L:51:ILE:HG23	4:L:52:GLY:N	2.35	0.41
4:L:143:TYR:HE1	4:L:164:PRO:HB3	1.83	0.41
4:L:413:TRP:CD1	4:L:418:GLN:CD	2.93	0.41
4:L:503:VAL:HG13	4:L:510:VAL:HB	2.02	0.41
4:L:566:VAL:HB	4:L:567:PRO:HD3	2.03	0.41
4:L:607:PRO:O	4:L:610:ILE:HG23	2.20	0.41
4:L:613:GLN:HG2	6:Z:4:CYS:SG	2.60	0.41
4:M:368:VAL:HG23	4:M:370:LEU:HD21	2.03	0.41
4:M:576:GLN:O	4:M:584:ARG:NH1	2.53	0.41
5:O:6:GLY:HA3	5:O:333:GLN:C	2.40	0.41
5:O:6:GLY:O	5:O:333:GLN:HB3	2.20	0.41
5:O:342:THR:HA	5:O:343:PRO:HD3	1.89	0.41
5:O:459:THR:O	5:O:461:PHE:N	2.54	0.41
5:O:616:ILE:HB	5:O:651:LEU:HB2	2.02	0.41
5:O:807:LEU:HB2	5:O:1009:ALA:HB1	2.03	0.41
3:P:156:THR:C	3:P:158:PHE:H	2.24	0.41
3:P:352:ALA:HA	3:P:355:ASN:ND2	2.35	0.41
3:P:414:ILE:H	3:P:414:ILE:HG13	1.61	0.41
6:X:35:THR:HG21	6:X:155:VAL:HG22	2.03	0.41
6:X:99:MET:N	6:X:99:MET:SD	2.93	0.41
6:X:120:ARG:O	6:X:123:THR:N	2.54	0.41
6:X:204:THR:HA	6:X:267:LYS:NZ	2.34	0.41
6:Y:220:VAL:O	6:Y:358:ILE:HG23	2.20	0.41
6:Y:254:LEU:HA	6:Y:276:LEU:HD11	2.03	0.41
6:Z:120:ARG:C	6:Z:120:ARG:HD3	2.41	0.41
1:B:381:ASP:OD2	1:B:389:ALA:N	2.54	0.41
1:B:399:ALA:HA	1:B:402:TRP:CD1	2.56	0.41
1:B:422:PHE:CE1	1:B:1227:ILE:HB	2.56	0.41
1:B:434:PHE:HB2	1:B:451:PHE:CE1	2.55	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:B:653:ALA:C	1:B:671:GLN:HE21	2.19	0.41
1:B:847:VAL:HA	1:B:1001:TYR:HA	2.02	0.41
1:B:1000:GLN:HA	1:B:1009:PHE:O	2.20	0.41
1:B:1036:ASN:C	1:B:1038:PHE:N	2.74	0.41
1:B:1045:ILE:HA	1:B:1200:ILE:O	2.20	0.41
1:B:1156:TYR:CG	1:B:1194:PRO:HB2	2.56	0.41
2:C:347:ARG:NH2	2:C:1176:PRO:HB3	2.36	0.41
2:C:477:THR:HA	2:C:480:GLU:OE2	2.21	0.41
2:C:548:PRO:HD3	2:C:900:TRP:CD1	2.56	0.41
2:C:682:HIS:CE1	2:C:683:THR:HG23	2.56	0.41
2:C:701:ILE:HA	2:C:704:GLN:OE1	2.20	0.41
2:C:716:PRO:HD2	2:C:837:ARG:HD3	2.03	0.41
2:C:749:ASN:N	2:C:813:GLN:OE1	2.54	0.41
2:C:789:LEU:O	2:C:793:MET:HG2	2.20	0.41
2:C:1037:LEU:HB2	2:C:1038:PHE:CD1	2.56	0.41
2:C:1166:TRP:HE1	2:C:1177:SER:C	2.24	0.41
2:C:1170:ILE:HG12	2:C:1176:PRO:HD2	2.03	0.41
2:C:1199:ILE:C	2:C:1200:ILE:HG13	2.41	0.41
3:D:42:PHE:HA	3:D:64:ARG:NH1	2.35	0.41
3:D:95:VAL:HA	3:D:104:LEU:HD23	2.02	0.41
3:D:223:SER:HA	3:D:228:ILE:HB	2.03	0.41
3:D:229:HIS:HB2	3:D:232:ASN:H	1.86	0.41
3:D:381:GLN:HG3	3:D:381:GLN:H	1.67	0.41
3:D:395:LYS:HD2	3:D:395:LYS:HA	1.78	0.41
4:K:65:ARG:HH22	4:M:258:MET:HG3	1.85	0.41
4:K:122:ARG:HA	4:K:125:LEU:HD12	2.02	0.41
4:K:171:GLN:O	4:K:174:TYR:HB3	2.20	0.41
4:K:305:VAL:HG21	4:L:652:SER:OG	2.21	0.41
4:K:377:ARG:CZ	4:K:379:VAL:HG22	2.51	0.41
4:K:538:CYS:O	4:K:542:ARG:NE	2.54	0.41
4:K:638:LYS:C	4:L:30:ALA:HB3	2.41	0.41
4:L:145:ASP:HA	4:L:162:GLN:H	1.85	0.41
4:L:213:GLN:CB	4:L:675:PRO:HD2	2.51	0.41
4:L:327:LYS:HE2	4:L:327:LYS:HB2	1.93	0.41
4:L:435:SER:OG	4:L:440:GLN:HB2	2.21	0.41
4:L:470:ALA:N	4:L:489:TRP:CZ2	2.89	0.41
4:L:471:THR:OG1	4:L:472:PHE:N	2.54	0.41
4:L:537:ARG:HA	4:L:540:ILE:HG22	2.03	0.41
4:L:639:SER:HA	4:M:31:VAL:O	2.19	0.41
4:M:234:LYS:O	4:M:238:VAL:HG22	2.20	0.41
4:M:413:TRP:HA	4:M:418:GLN:OE1	2.21	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
4:M:433:GLU:O	4:M:441:SER:OG	2.25	0.41
4:M:607:PRO:O	4:M:610:ILE:HG12	2.21	0.41
5:O:81:ASP:N	5:O:81:ASP:OD2	2.53	0.41
5:O:323:THR:OG1	5:O:326:TRP:N	2.35	0.41
5:O:334:ILE:C	5:O:336:SER:H	2.24	0.41
5:O:371:ARG:HH12	5:O:375:LEU:CB	2.30	0.41
5:O:393:PRO:HA	5:O:741:ILE:O	2.21	0.41
5:O:408:ASP:O	5:O:411:ASP:HB3	2.20	0.41
5:O:410:GLY:O	5:O:414:VAL:HG13	2.21	0.41
5:O:426:TYR:CD1	5:O:694:PHE:HA	2.55	0.41
5:O:430:TRP:CZ3	5:O:1000:THR:HA	2.55	0.41
5:O:431:VAL:HA	5:O:477:LYS:H	1.86	0.41
5:O:476:ARG:NH1	5:O:646:THR:HA	2.35	0.41
5:O:510:GLN:HB2	5:O:572:GLN:CD	2.41	0.41
5:O:580:GLN:HB3	5:O:583:ASP:OD2	2.20	0.41
5:O:735:GLY:HA3	5:O:765:ARG:CZ	2.51	0.41
5:O:778:ARG:CZ	5:O:782:VAL:HG22	2.50	0.41
5:O:938:GLY:O	5:O:953:LYS:HE2	2.21	0.41
3:P:81:ARG:HB2	3:P:83:ASN:O	2.21	0.41
3:P:85:ARG:HG3	3:P:86:TRP:N	2.36	0.41
3:P:362:ASP:HA	3:P:365:GLN:CG	2.51	0.41
6:X:282:MET:HG2	6:X:283:TYR:N	2.35	0.41
6:Y:79:HIS:O	6:Y:81:ASP:N	2.54	0.41
6:Y:115:PRO:O	6:Y:118:LEU:HB3	2.21	0.41
6:Y:185:MET:HE2	6:Y:187:ASP:O	2.21	0.41
6:Y:297:LYS:HB2	6:Y:297:LYS:HE2	1.92	0.41
6:Y:309:GLU:HB2	6:Y:313:TYR:OH	2.21	0.41
6:Z:95:TRP:HE1	6:Z:268:MET:HB3	1.86	0.41
6:Z:121:VAL:HG21	6:Z:128:LEU:HD13	2.03	0.41
6:Z:130:GLU:HB3	6:Z:362:MET:HB3	2.03	0.41
6:Z:136:VAL:H	6:Z:143:ARG:HH12	1.69	0.41
6:Z:310:LYS:HE2	6:Z:310:LYS:HB2	1.88	0.41
1:B:437:ALA:HB2	1:B:447:TRP:CD1	2.56	0.41
1:B:585:ASN:N	1:B:585:ASN:OD1	2.44	0.41
1:B:702:LEU:HD12	1:B:703:ARG:N	2.36	0.41
1:B:807:THR:C	1:B:810:TYR:HB3	2.41	0.41
1:B:1116:GLN:HG2	1:B:1170:ILE:CG1	2.51	0.41
2:C:394:LEU:HD23	2:C:399:ALA:HA	2.03	0.41
2:C:534:PRO:HA	2:C:537:GLN:CG	2.51	0.41
2:C:767:ARG:O	2:C:768:VAL:C	2.59	0.41
2:C:1000:GLN:HG2	3:P:257:LEU:HB3	2.03	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:D:95:VAL:H	3:D:105:VAL:N	2.15	0.41
3:D:179:LEU:HD11	3:D:212:MET:HE1	2.02	0.41
3:D:213:THR:HA	3:D:216:ILE:HG12	2.03	0.41
3:D:221:TRP:O	3:D:224:TYR:HB3	2.21	0.41
4:K:335:ARG:HB3	4:K:411:GLU:OE2	2.21	0.41
4:K:503:VAL:O	4:K:510:VAL:HG12	2.21	0.41
4:K:631:VAL:HG13	4:K:632:LYS:H	1.86	0.41
4:L:20:PHE:HD1	4:L:247:ILE:N	2.19	0.41
4:L:188:TRP:NE1	4:L:192:LEU:HD11	2.35	0.41
4:L:336:MET:HG2	4:L:365:THR:O	2.21	0.41
4:L:385:LYS:HE2	4:L:385:LYS:HB2	1.95	0.41
4:L:407:LYS:HD3	4:L:463:GLU:OE1	2.21	0.41
4:L:586:PHE:O	4:L:587:ASN:ND2	2.54	0.41
4:M:461:ASP:HB2	4:M:464:MET:HE3	2.02	0.41
5:O:10:ALA:HB2	5:O:313:GLN:HG2	2.03	0.41
5:O:37:ASN:CG	5:O:40:ARG:HB2	2.42	0.41
5:O:131:ASP:OD1	5:O:132:LEU:HG	2.21	0.41
5:O:649:VAL:HG23	5:O:681:TYR:CE2	2.55	0.41
5:O:1108:ASN:HD21	5:O:1110:ASN:ND2	2.19	0.41
5:O:1136:PHE:CD2	5:O:1228:LEU:HD22	2.56	0.41
5:O:1237:GLU:CB	5:O:1289:LEU:HB2	2.51	0.41
3:P:11:VAL:HG22	3:P:12:GLY:O	2.21	0.41
3:P:171:TRP:CZ3	3:P:235:ARG:HD2	2.56	0.41
3:P:221:TRP:HA	3:P:224:TYR:HB3	2.03	0.41
3:P:243:ARG:CZ	3:P:250:VAL:HG23	2.51	0.41
3:P:273:MET:HG3	3:P:274:PRO:O	2.21	0.41
3:P:288:SER:O	3:P:291:LEU:HG	2.21	0.41
6:X:186:ARG:HG2	6:X:347:THR:HG23	2.03	0.41
6:X:235:GLY:O	6:X:239:ARG:HG3	2.21	0.41
6:X:263:PRO:HA	6:X:269:PRO:N	2.36	0.41
6:X:306:TRP:CE3	6:X:310:LYS:HD2	2.56	0.41
6:Y:51:CYS:SG	6:Y:52:MET:N	2.94	0.41
6:Y:54:CYS:HB3	6:Y:73:CYS:CB	2.51	0.41
6:Y:79:HIS:O	6:Y:83:VAL:HG13	2.21	0.41
6:Z:227:GLU:HA	6:Z:230:HIS:ND1	2.36	0.41
6:Z:238:TYR:CZ	6:Z:242:LEU:HD13	2.56	0.41
6:Z:258:SER:CB	6:Z:273:SER:HA	2.51	0.41
1:B:281:LEU:HD21	1:B:286:PHE:HE2	1.86	0.40
1:B:350:ASN:HA	1:B:1173:THR:HG22	2.02	0.40
1:B:466:MET:HA	1:B:469:LEU:HD12	2.03	0.40
1:B:685:PRO:O	1:B:688:PHE:HB3	2.21	0.40

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:C:300:ALA:O	2:C:1213:ASN:ND2	2.53	0.40
2:C:528:ASN:HB2	2:C:531:VAL:CG2	2.50	0.40
2:C:854:ARG:CZ	3:P:258:ASN:HA	2.51	0.40
2:C:1037:LEU:H	2:C:1037:LEU:HG	1.45	0.40
2:C:1056:TRP:HE3	2:C:1057:SER:O	2.03	0.40
2:C:1074:VAL:HG13	2:C:1107:ASN:C	2.41	0.40
3:D:230:GLN:NE2	3:D:240:GLN:NE2	2.69	0.40
3:D:236:THR:O	3:D:256:SER:N	2.46	0.40
3:D:333:ARG:HH12	3:D:334:ARG:NH1	2.19	0.40
4:K:19:VAL:O	4:K:248:GLN:N	2.48	0.40
4:K:388:LYS:HD3	4:K:394:PRO:HD2	2.03	0.40
4:K:399:VAL:O	4:K:427:TYR:HB2	2.21	0.40
4:K:577:LEU:C	4:K:584:ARG:HH22	2.24	0.40
4:K:625:ILE:HG21	4:L:660:TRP:CZ3	2.56	0.40
4:L:32:PRO:CG	4:M:38:PRO:HB2	2.49	0.40
4:L:234:LYS:H	4:L:234:LYS:HG3	1.72	0.40
4:L:334:LEU:HD13	4:L:366:ARG:HH22	1.86	0.40
5:O:119:LEU:HD21	5:O:136:LEU:HD21	2.03	0.40
5:O:273:ALA:O	5:O:275:PRO:HD3	2.21	0.40
5:O:285:LEU:HD21	5:O:289:ALA:N	2.36	0.40
5:O:1056:TYR:CZ	5:O:1061:LEU:HD13	2.56	0.40
3:P:110:GLN:HG3	3:P:111:VAL:H	1.82	0.40
3:P:133:LEU:HB3	3:P:140:LYS:HG2	2.04	0.40
3:P:292:THR:HA	3:P:295:LEU:CG	2.51	0.40
3:P:302:PRO:O	3:P:303:LEU:HD23	2.22	0.40
6:X:183:PHE:CG	6:X:261:THR:HB	2.56	0.40
6:Y:68:LEU:HD12	6:Y:69:PRO:HD2	2.03	0.40
6:Z:159:LEU:H	6:Z:159:LEU:HD12	1.85	0.40
6:Z:295:VAL:O	6:Z:295:VAL:HG22	2.21	0.40
1:B:598:TRP:O	1:B:601:ASN:HB2	2.22	0.40
1:B:937:GLN:OE1	1:B:947:ARG:HG2	2.22	0.40
1:B:1217:PRO:HG2	2:C:1087:MET:HG2	2.02	0.40
2:C:797:LEU:HA	2:C:800:LEU:HB2	2.03	0.40
2:C:915:PHE:CG	2:C:916:SER:N	2.89	0.40
2:C:922:MET:HA	2:C:925:CYS:SG	2.61	0.40
2:C:979:LEU:HD23	2:C:983:LEU:HD11	2.03	0.40
3:D:110:GLN:H	3:D:110:GLN:CD	2.23	0.40
3:D:120:GLN:CG	3:D:123:ARG:HH21	2.34	0.40
3:D:187:PHE:N	3:D:239:PHE:HE1	2.20	0.40
3:D:333:ARG:NH1	3:D:334:ARG:HD3	2.35	0.40
4:K:661:THR:HA	4:K:664:PHE:HB3	2.04	0.40

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
4:L:315:PRO:HA	4:L:316:PRO:HD3	1.92	0.40
4:L:367:VAL:HG22	4:L:368:VAL:N	2.36	0.40
4:M:188:TRP:HH2	4:M:235:GLU:HB3	1.85	0.40
4:M:188:TRP:CH2	4:M:235:GLU:HB3	2.56	0.40
4:M:229:ILE:HG23	4:M:232:TYR:O	2.21	0.40
4:M:252:VAL:HG23	4:M:253:SER:N	2.35	0.40
5:O:27:LEU:N	5:O:109:GLU:H	2.17	0.40
5:O:691:LEU:HG	5:O:692:PRO:N	2.35	0.40
5:O:830:THR:CG2	5:O:848:CYS:HB3	2.48	0.40
5:O:885:ILE:HD13	5:O:921:VAL:HB	2.03	0.40
5:O:934:ARG:HH12	5:O:1210:ILE:HA	1.86	0.40
6:X:132:ASN:ND2	6:X:134:LEU:HB2	2.37	0.40
6:X:218:TRP:NE1	6:X:358:ILE:HD11	2.36	0.40
6:X:308:VAL:HG23	6:X:309:GLU:H	1.86	0.40
6:Y:177:SER:OG	6:Y:179:LEU:HG	2.21	0.40
6:Z:100:LEU:HD23	6:Z:101:SER:N	2.36	0.40
6:Z:137:ASP:OD2	6:Z:140:SER:HB3	2.21	0.40
6:Z:162:LYS:HG2	6:Z:166:TYR:HE2	1.87	0.40
6:Z:222:VAL:C	6:Z:356:VAL:HG13	2.42	0.40
6:Z:223:TYR:HA	6:Z:356:VAL:HG22	2.03	0.40
6:Z:227:GLU:HG3	6:Z:228:LEU:HD12	2.01	0.40
6:Z:300:ASP:O	6:Z:304:HIS:CD2	2.74	0.40
1:B:266:PHE:CB	1:B:1253:LEU:HD21	2.51	0.40
1:B:494:THR:HG22	1:B:1272:MET:HB2	2.03	0.40
1:B:602:GLY:H	1:B:832:GLN:HA	1.85	0.40
1:B:629:PHE:HA	1:B:632:LEU:HD12	2.04	0.40
1:B:837:ARG:HH12	1:B:838:LEU:HD11	1.85	0.40
1:B:852:GLN:OE1	1:B:991:ASP:HB3	2.22	0.40
1:B:914:VAL:HG13	1:B:915:PHE:HD1	1.87	0.40
1:B:935:VAL:O	1:B:938:ILE:N	2.51	0.40
1:B:961:THR:O	1:B:964:THR:HB	2.22	0.40
2:C:504:LEU:O	2:C:1264:THR:HG23	2.21	0.40
2:C:636:LEU:HA	2:C:639:THR:HG23	2.04	0.40
2:C:1042:ARG:HB3	2:C:1203:GLU:OE2	2.21	0.40
3:D:65:TYR:HA	3:D:68:MET:SD	2.62	0.40
3:D:81:ARG:CZ	3:D:87:GLY:HA2	2.52	0.40
4:K:20:PHE:CD1	4:K:247:ILE:HD13	2.56	0.40
4:K:175:VAL:O	4:K:179:GLN:HG2	2.22	0.40
4:K:218:LEU:HD12	4:K:218:LEU:HA	1.89	0.40
4:K:223:LEU:C	4:K:230:ARG:HH21	2.25	0.40
4:K:248:GLN:HB3	4:K:250:MET:CE	2.51	0.40

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
4:K:318:PRO:HB3	4:K:496:SER:HA	2.04	0.40
4:K:451:ASN:N	4:K:451:ASN:OD1	2.52	0.40
4:L:180:ALA:O	4:L:183:LYS:HB3	2.21	0.40
4:L:343:THR:O	4:L:363:GLY:N	2.54	0.40
4:L:379:VAL:HG22	4:L:449:ALA:HB2	2.03	0.40
4:M:188:TRP:HA	4:M:191:ASP:OD2	2.22	0.40
4:M:549:ALA:HA	4:M:552:ILE:CD1	2.51	0.40
5:O:207:LEU:HD13	5:O:210:PHE:CB	2.51	0.40
5:O:426:TYR:HB3	5:O:694:PHE:CE2	2.57	0.40
5:O:498:ASP:OD2	5:O:501:THR:HG22	2.21	0.40
5:O:988:LEU:HA	5:O:990:TRP:NE1	2.36	0.40
5:O:1019:MET:HB3	5:O:1021:ILE:CD1	2.52	0.40
3:P:101:LEU:HA	3:P:101:LEU:HD13	1.77	0.40
3:P:217:LEU:HG	3:P:221:TRP:CD1	2.57	0.40
3:P:309:VAL:CG2	3:P:314:ARG:HG2	2.51	0.40
6:X:155:VAL:HG12	6:X:156:VAL:N	2.36	0.40
6:X:308:VAL:HG23	6:X:309:GLU:N	2.36	0.40
6:Z:7:ASN:HA	6:Z:315:LEU:HD22	2.02	0.40
6:Z:210:PHE:CZ	6:Z:268:MET:HG2	2.57	0.40
1:B:332:ILE:O	1:B:334:LEU:N	2.54	0.40
1:B:363:LEU:O	1:B:366:VAL:HB	2.21	0.40
1:B:509:ILE:HD12	1:B:513:GLU:HB2	2.03	0.40
1:B:651:THR:O	1:B:655:MET:HG2	2.22	0.40
1:B:985:GLU:HB2	1:B:986:PRO:HD3	2.04	0.40
1:B:1062:PRO:HB2	1:B:1065:LEU:H	1.86	0.40
1:B:1080:ASP:OD1	1:B:1095:ARG:HG2	2.21	0.40
1:B:1168:GLU:HG3	1:B:1169:GLU:OE2	2.22	0.40
2:C:260:ALA:HB3	2:C:312:SER:OG	2.21	0.40
2:C:285:PHE:CG	2:C:286:PHE:N	2.89	0.40
2:C:301:VAL:HG22	2:C:1212:THR:HG22	2.03	0.40
2:C:624:GLU:HA	2:C:627:TRP:HE3	1.84	0.40
2:C:668:ILE:HB	2:C:669:TYR:CE1	2.57	0.40
2:C:853:SER:HB2	2:C:942:GLN:HB3	2.04	0.40
3:D:275:THR:HB	3:D:401:GLN:HE22	1.86	0.40
3:D:368:LYS:HG3	3:D:369:ARG:N	2.37	0.40
4:K:263:ASN:HA	4:L:43:PRO:CD	2.51	0.40
4:K:380:LEU:HD13	4:K:401:PHE:CZ	2.56	0.40
4:K:473:ILE:HG13	4:K:473:ILE:O	2.22	0.40
4:K:499:SER:HB2	4:K:502:GLU:OE1	2.22	0.40
4:K:645:LYS:HZ1	4:K:649:SER:HB3	1.86	0.40
4:L:34:LEU:HD23	4:L:34:LEU:HA	1.85	0.40

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
4:L:148:VAL:HG12	4:L:149:GLY:N	2.36	0.40
4:L:158:ASN:CG	4:L:160:GLN:HE21	2.23	0.40
4:L:468:LEU:HD12	4:L:469:LEU:N	2.35	0.40
4:L:625:ILE:HG21	4:M:657:ILE:HD12	2.02	0.40
4:M:46:VAL:CG1	4:M:65:ARG:HG3	2.51	0.40
4:M:518:LEU:O	4:M:611:ILE:HG23	2.21	0.40
5:O:483:LEU:O	5:O:529:VAL:HG11	2.22	0.40
5:O:653:PHE:HE2	5:O:655:ALA:HB2	1.87	0.40
5:O:1270:TRP:HE1	5:O:1274:ILE:HG12	1.87	0.40
3:P:392:PHE:CE1	3:P:396:PRO:HG3	2.57	0.40
6:X:8:GLY:HA2	6:X:11:ILE:HG12	2.03	0.40
6:X:43:MET:HE2	6:X:52:MET:HG3	2.03	0.40
6:X:68:LEU:HA	6:X:69:PRO:HD3	1.95	0.40
6:X:171:ASN:HA	6:X:174:ILE:CG1	2.49	0.40
6:X:223:TYR:HA	6:X:355:PRO:O	2.22	0.40
6:X:255:SER:HG	6:X:276:LEU:H	1.60	0.40
6:Y:103:VAL:O	6:Y:107:HIS:HB2	2.21	0.40
6:Y:221:MET:HE3	6:Y:263:PRO:HB2	2.03	0.40
6:Y:236:ARG:HA	6:Y:236:ARG:NH1	2.36	0.40
1:B:472:MET:HB3	1:B:507:TYR:C	2.41	0.40
1:B:522:ILE:HA	1:B:525:ILE:CD1	2.52	0.40
1:B:764:TRP:HA	1:B:767:ARG:HH11	1.87	0.40
2:C:541:VAL:HG13	2:C:545:ARG:HH21	1.87	0.40
2:C:623:LEU:O	2:C:626:LEU:HB2	2.21	0.40
2:C:837:ARG:C	2:C:842:ARG:HH22	2.24	0.40
2:C:948:TYR:OH	2:C:1029:GLU:N	2.55	0.40
2:C:1049:ARG:HG3	2:C:1196:VAL:HG11	2.04	0.40
3:D:230:GLN:NE2	3:D:240:GLN:HE21	2.15	0.40
4:K:113:LYS:O	4:K:116:MET:HB2	2.22	0.40
4:K:171:GLN:O	4:K:175:VAL:HG13	2.22	0.40
4:K:282:LYS:HD3	4:K:642:ALA:HB2	2.03	0.40
4:K:632:LYS:HZ2	4:L:202:ASN:HB3	1.86	0.40
4:K:655:GLU:HA	4:K:658:GLN:HE21	1.87	0.40
4:L:101:VAL:O	4:L:165:VAL:HA	2.22	0.40
4:L:216:VAL:O	4:L:219:LEU:HG	2.21	0.40
4:L:237:ALA:HA	4:L:240:LEU:HG	2.04	0.40
4:L:667:LYS:O	4:L:670:THR:HB	2.22	0.40
4:M:37:SER:HG	4:M:40:MET:HB2	1.87	0.40
4:M:109:ALA:HB1	4:M:113:LYS:HZ3	1.86	0.40
4:M:153:ARG:HD3	4:M:153:ARG:N	2.37	0.40
4:M:369:ASN:HA	4:M:467:TYR:CD2	2.51	0.40

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
5:O:17:THR:HA	5:O:279:LEU:HD11	2.03	0.40
5:O:280:ASP:OD2	5:O:309:THR:HG23	2.21	0.40
5:O:297:LEU:O	5:O:300:ARG:HB2	2.22	0.40
5:O:609:GLY:HA2	5:O:660:GLN:CD	2.42	0.40
5:O:908:GLN:NE2	5:O:911:ILE:HB	2.37	0.40
5:O:1249:ILE:O	5:O:1252:SER:OG	2.31	0.40
3:P:211:ARG:O	3:P:214:GLN:HB2	2.22	0.40
3:P:223:SER:C	3:P:226:GLY:H	2.24	0.40
3:P:229:HIS:CE1	3:P:231:GLN:HB2	2.57	0.40
6:X:20:GLU:O	6:X:22:ARG:HG2	2.22	0.40
6:Y:44:VAL:HG22	6:Y:49:VAL:HG22	2.03	0.40
6:Y:54:CYS:HB3	6:Y:73:CYS:HB2	2.03	0.40
6:Y:299:VAL:HG13	6:Y:300:ASP:H	1.86	0.40
6:Z:123:THR:HG22	6:Z:124:GLU:HG3	2.03	0.40
6:Z:197:LEU:HD22	6:Z:201:ALA:HB1	2.03	0.40
6:Z:199:GLY:N	6:Z:202:ARG:HE	2.13	0.40

There are no symmetry-related clashes.

5.3 Torsion angles [i](#)

5.3.1 Protein backbone [i](#)

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

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

Mol	Chain	Analysed	Favoured	Allowed	Outliers	Percentiles	
1	B	1031/1035 (100%)	841 (82%)	190 (18%)	0	100	100
2	C	1004/1008 (100%)	820 (82%)	179 (18%)	5 (0%)	25	64
3	D	415/417 (100%)	343 (83%)	71 (17%)	1 (0%)	44	78
3	P	415/417 (100%)	352 (85%)	62 (15%)	1 (0%)	44	78
4	K	637/641 (99%)	548 (86%)	88 (14%)	1 (0%)	44	78
4	L	637/641 (99%)	536 (84%)	100 (16%)	1 (0%)	44	78
4	M	637/641 (99%)	550 (86%)	86 (14%)	1 (0%)	44	78

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Mol	Chain	Analysed	Favoured	Allowed	Outliers	Percentiles	
5	O	1280/1284 (100%)	1069 (84%)	211 (16%)	0	100	100
6	X	363/365 (100%)	309 (85%)	54 (15%)	0	100	100
6	Y	363/365 (100%)	317 (87%)	46 (13%)	0	100	100
6	Z	363/365 (100%)	296 (82%)	67 (18%)	0	100	100
All	All	7145/7179 (100%)	5981 (84%)	1154 (16%)	10 (0%)	50	83

All (10) Ramachandran outliers are listed below:

Mol	Chain	Res	Type
2	C	1088	ASN
3	D	49	GLY
4	M	387	TYR
4	K	372	GLN
3	P	170	MET
2	C	996	GLN
2	C	992	PRO
2	C	1250	VAL
2	C	907	PRO
4	L	616	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	B	915/915 (100%)	908 (99%)	7 (1%)	79	85
2	C	890/890 (100%)	882 (99%)	8 (1%)	75	83
3	D	352/352 (100%)	346 (98%)	6 (2%)	56	72
3	P	352/352 (100%)	346 (98%)	6 (2%)	56	72
4	K	541/541 (100%)	534 (99%)	7 (1%)	65	77
4	L	541/541 (100%)	537 (99%)	4 (1%)	81	87
4	M	541/541 (100%)	538 (99%)	3 (1%)	84	88
5	O	1118/1118 (100%)	1101 (98%)	17 (2%)	60	75

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Mol	Chain	Analysed	Rotameric	Outliers	Percentiles	
6	X	317/317 (100%)	312 (98%)	5 (2%)	58	74
6	Y	317/317 (100%)	315 (99%)	2 (1%)	84	88
6	Z	317/317 (100%)	314 (99%)	3 (1%)	75	83
All	All	6201/6201 (100%)	6133 (99%)	68 (1%)	69	80

All (68) residues with a non-rotameric sidechain are listed below:

Mol	Chain	Res	Type
1	B	335	PHE
1	B	374	ARG
1	B	592	ARG
1	B	674	ARG
1	B	993	ARG
1	B	1101	MET
1	B	1138	ARG
2	C	521	ARG
2	C	720	ARG
2	C	781	TYR
2	C	851	ARG
2	C	920	ARG
2	C	956	ARG
2	C	995	THR
2	C	1069	ARG
3	D	9	LYS
3	D	31	ARG
3	D	93	ARG
3	D	140	LYS
3	D	333	ARG
3	D	377	ARG
4	K	161	LYS
4	K	183	LYS
4	K	327	LYS
4	K	333	TRP
4	K	381	ASP
4	K	410	PHE
4	K	666	ASP
4	L	183	LYS
4	L	231	ARG
4	L	362	ARG
4	L	664	PHE
4	M	153	ARG

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Mol	Chain	Res	Type
4	M	231	ARG
4	M	551	LYS
5	O	8	ARG
5	O	40	ARG
5	O	200	TYR
5	O	226	LYS
5	O	269	ARG
5	O	312	TYR
5	O	416	ARG
5	O	446	ARG
5	O	456	LEU
5	O	481	ARG
5	O	548	ARG
5	O	620	THR
5	O	673	PHE
5	O	757	ARG
5	O	950	ARG
5	O	956	ARG
5	O	1007	ARG
3	P	50	ARG
3	P	187	PHE
3	P	203	TRP
3	P	258	ASN
3	P	284	ASN
3	P	333	ARG
6	X	7	ASN
6	X	78	ARG
6	X	141	MET
6	X	150	THR
6	X	208	ARG
6	Y	63	ARG
6	Y	213	ARG
6	Z	19	PHE
6	Z	78	ARG
6	Z	353	ASP

Sometimes sidechains can be flipped to improve hydrogen bonding and reduce clashes. All (189) such sidechains are listed below:

Mol	Chain	Res	Type
1	B	246	GLN
1	B	278	GLN
1	B	333	HIS

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Mol	Chain	Res	Type
1	B	390	ASN
1	B	475	ASN
1	B	488	GLN
1	B	518	GLN
1	B	524	ASN
1	B	537	GLN
1	B	601	ASN
1	B	625	ASN
1	B	667	GLN
1	B	710	HIS
1	B	715	ASN
1	B	731	ASN
1	B	759	ASN
1	B	782	GLN
1	B	996	GLN
1	B	1125	GLN
1	B	1197	GLN
2	C	275	GLN
2	C	278	GLN
2	C	340	ASN
2	C	357	ASN
2	C	421	ASN
2	C	488	GLN
2	C	527	ASN
2	C	550	GLN
2	C	558	ASN
2	C	654	ASN
2	C	671	GLN
2	C	745	HIS
2	C	782	GLN
2	C	812	GLN
2	C	929	GLN
2	C	1002	GLN
2	C	1003	GLN
2	C	1010	ASN
2	C	1054	HIS
2	C	1118	ASN
2	C	1158	ASN
2	C	1213	ASN
2	C	1244	GLN
2	C	1246	GLN
2	C	1255	ASN

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Mol	Chain	Res	Type
3	D	38	GLN
3	D	67	GLN
3	D	118	GLN
3	D	181	ASN
3	D	200	ASN
3	D	214	GLN
3	D	229	HIS
3	D	230	GLN
3	D	231	GLN
3	D	232	ASN
3	D	240	GLN
3	D	258	ASN
3	D	262	GLN
3	D	280	ASN
3	D	353	GLN
3	D	363	GLN
3	D	365	GLN
3	D	381	GLN
3	D	388	ASN
3	D	399	ASN
4	K	12	ASN
4	K	42	ASN
4	K	110	ASN
4	K	121	ASN
4	K	154	GLN
4	K	157	ASN
4	K	162	GLN
4	K	179	GLN
4	K	196	GLN
4	K	213	GLN
4	K	263	ASN
4	K	288	GLN
4	K	372	GLN
4	K	426	ASN
4	K	485	GLN
4	K	571	GLN
4	K	601	GLN
4	K	622	ASN
4	K	671	HIS
4	L	42	ASN
4	L	106	HIS
4	L	157	ASN

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Mol	Chain	Res	Type
4	L	196	GLN
4	L	202	ASN
4	L	222	GLN
4	L	405	GLN
4	L	418	GLN
4	L	465	ASN
4	L	528	ASN
4	L	587	ASN
4	L	601	GLN
4	L	629	GLN
4	L	658	GLN
4	L	671	HIS
4	M	42	ASN
4	M	157	ASN
4	M	158	ASN
4	M	179	GLN
4	M	222	GLN
4	M	248	GLN
4	M	360	ASN
4	M	369	ASN
4	M	372	GLN
4	M	405	GLN
4	M	440	GLN
4	M	456	GLN
4	M	485	GLN
4	M	528	ASN
4	M	564	ASN
4	M	571	GLN
4	M	613	GLN
4	M	671	HIS
5	O	49	GLN
5	O	117	ASN
5	O	180	ASN
5	O	214	HIS
5	O	250	ASN
5	O	399	GLN
5	O	419	GLN
5	O	492	HIS
5	O	526	GLN
5	O	572	GLN
5	O	617	ASN
5	O	659	HIS

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Mol	Chain	Res	Type
5	O	736	ASN
5	O	804	HIS
5	O	812	ASN
5	O	908	GLN
5	O	909	GLN
5	O	925	GLN
5	O	1048	ASN
5	O	1079	GLN
5	O	1108	ASN
5	O	1110	ASN
5	O	1133	GLN
5	O	1172	GLN
5	O	1215	GLN
3	P	22	ASN
3	P	84	HIS
3	P	141	HIS
3	P	154	ASN
3	P	176	ASN
3	P	214	GLN
3	P	229	HIS
3	P	230	GLN
3	P	242	ASN
3	P	258	ASN
3	P	262	GLN
3	P	307	ASN
3	P	315	ASN
3	P	331	GLN
3	P	354	GLN
3	P	355	ASN
3	P	356	GLN
3	P	365	GLN
3	P	406	ASN
6	X	39	GLN
6	X	53	HIS
6	X	79	HIS
6	X	86	GLN
6	X	107	HIS
6	X	139	ASN
6	X	171	ASN
6	X	193	ASN
6	X	205	GLN
6	X	251	HIS

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Mol	Chain	Res	Type
6	Y	9	HIS
6	Y	39	GLN
6	Y	62	GLN
6	Y	80	GLN
6	Y	107	HIS
6	Y	139	ASN
6	Y	165	GLN
6	Y	171	ASN
6	Y	230	HIS
6	Y	329	GLN
6	Y	330	GLN
6	Z	39	GLN
6	Z	62	GLN
6	Z	79	HIS
6	Z	80	GLN
6	Z	135	GLN
6	Z	171	ASN
6	Z	304	HIS

5.3.3 RNA [i](#)

There are no RNA molecules in this entry.

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

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

5.5 Carbohydrates [i](#)

There are no oligosaccharides in this entry.

5.6 Ligand geometry [i](#)

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

The following chains have linkage breaks:

Mol	Chain	Number of breaks
2	C	1
4	L	1
4	M	1
4	K	1
1	B	1
5	O	1

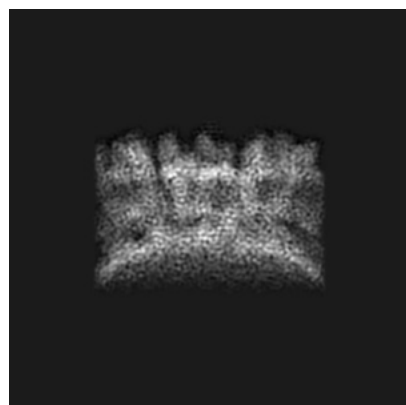
All chain breaks are listed below:

Model	Chain	Residue-1	Atom-1	Residue-2	Atom-2	Distance (Å)
1	C	562:THR	C	571:LEU	N	14.85
1	L	71:ILE	C	97:ASP	N	12.37
1	M	71:ILE	C	97:ASP	N	11.46
1	K	71:ILE	C	97:ASP	N	9.27
1	B	583:PRO	C	584:SER	N	8.38
1	O	1175:SER	C	1180:THR	N	6.54

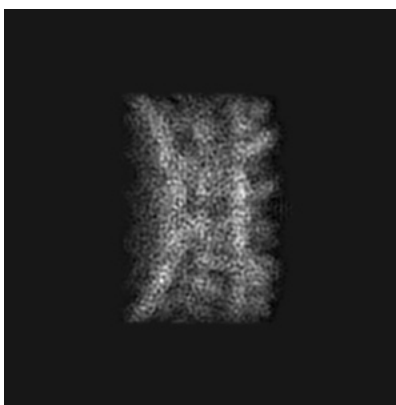
6 Tomogram visualisation [i](#)

This section contains visualisations of the EMDB entry EMD-22166. These allow visual inspection of the internal detail of the tomogram and identification of artifacts.

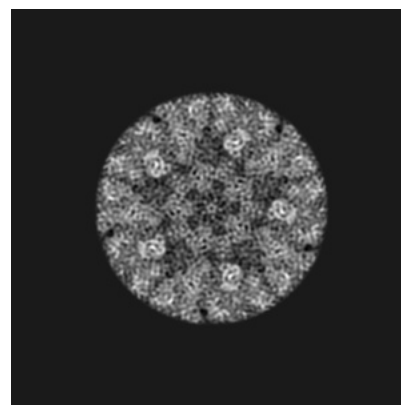
6.1 Orthogonal projections [i](#)



X



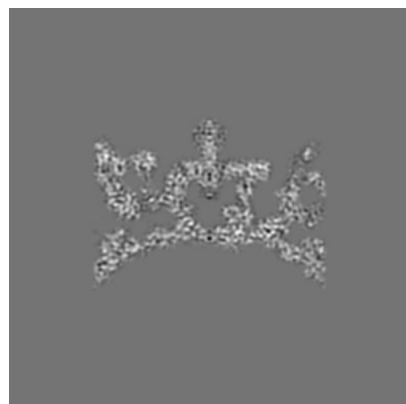
Y



Z

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

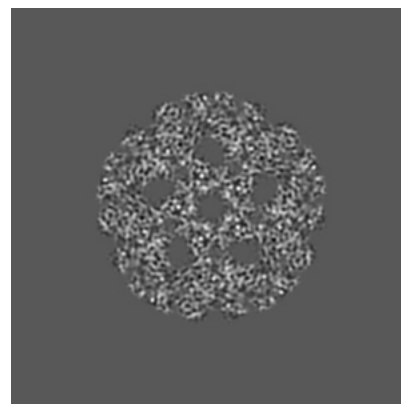
6.2 Central slices [i](#)



X Index: 180



Y Index: 180



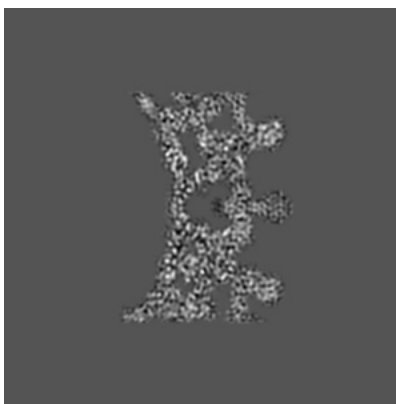
Z Index: 180

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

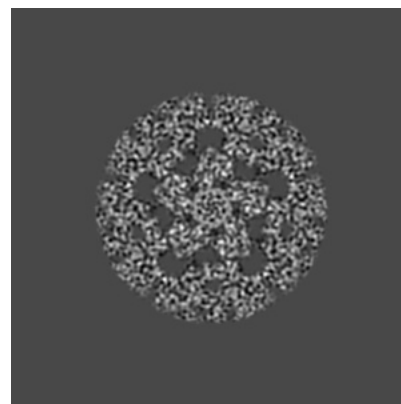
6.3 Largest variance slices [i](#)



X Index: 195



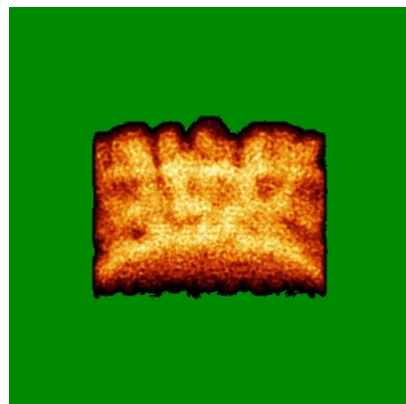
Y Index: 176



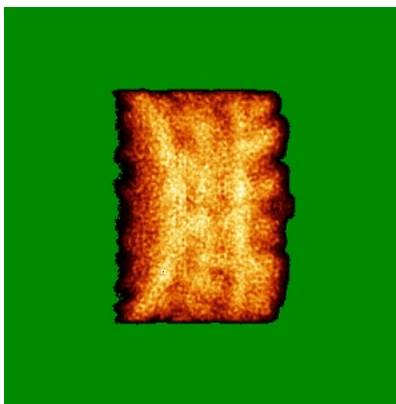
Z Index: 213

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

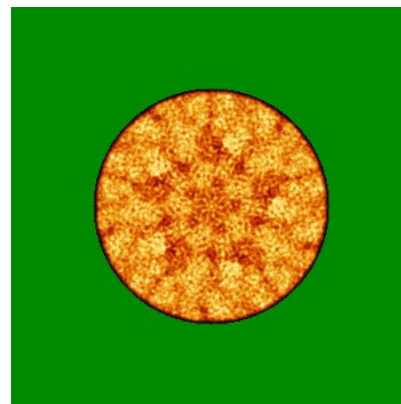
6.4 Orthogonal standard-deviation projections (False-color) [i](#)



X



Y



Z

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

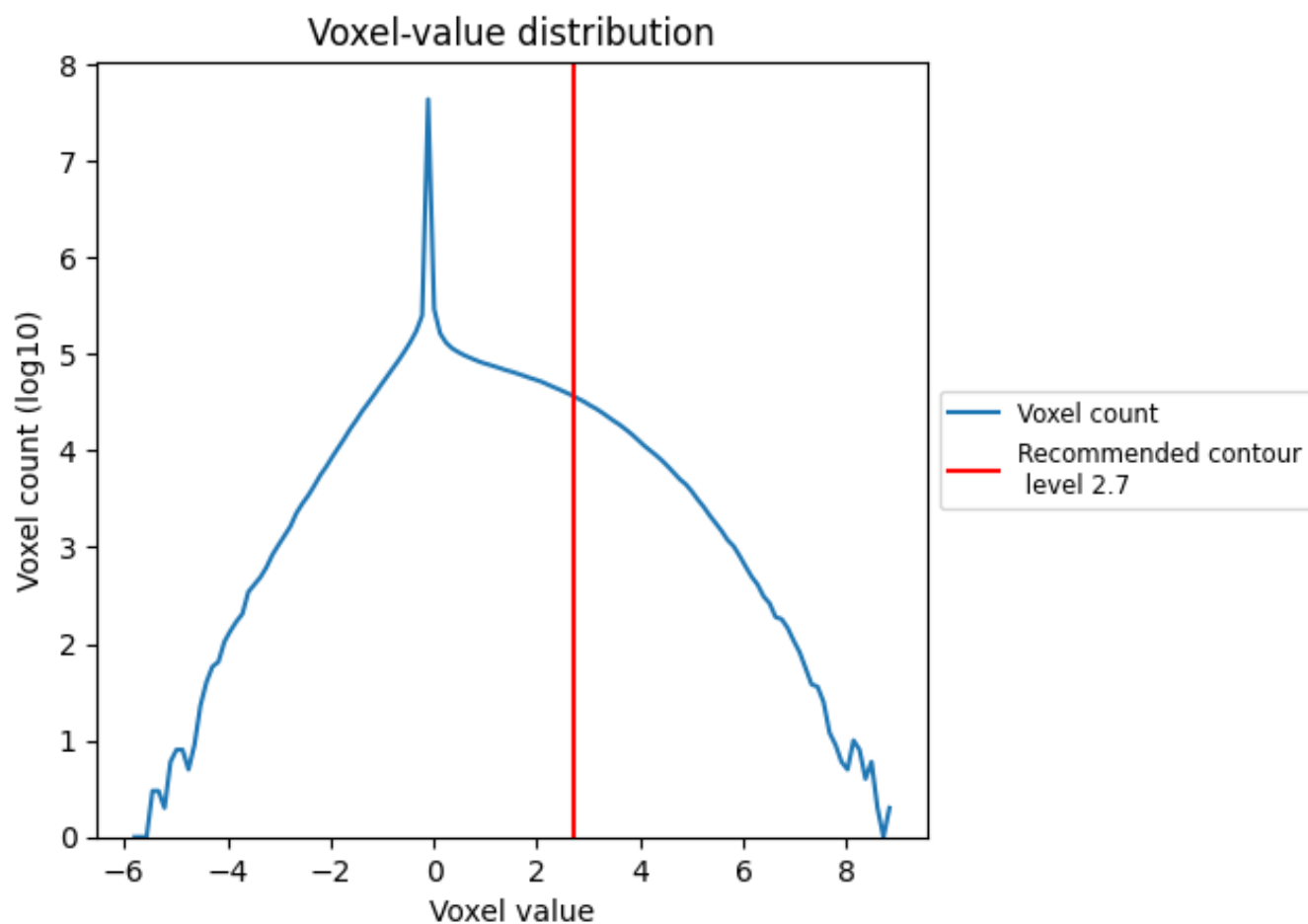
6.5 Mask visualisation [i](#)

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

7 Tomogram analysis [i](#)

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

7.1 Voxel-value distribution [i](#)



The voxel-value distribution is plotted in 128 intervals along the x-axis. The y-axis is logarithmic.

8 Map-model fit [i](#)

This section contains information regarding the fit between EMD map EMD-22166 and PDB model 6ZTZ. Per-residue inclusion information can be found in section 3 on page 8.

8.1 Map-model overlay [i](#)

This section was not generated.

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

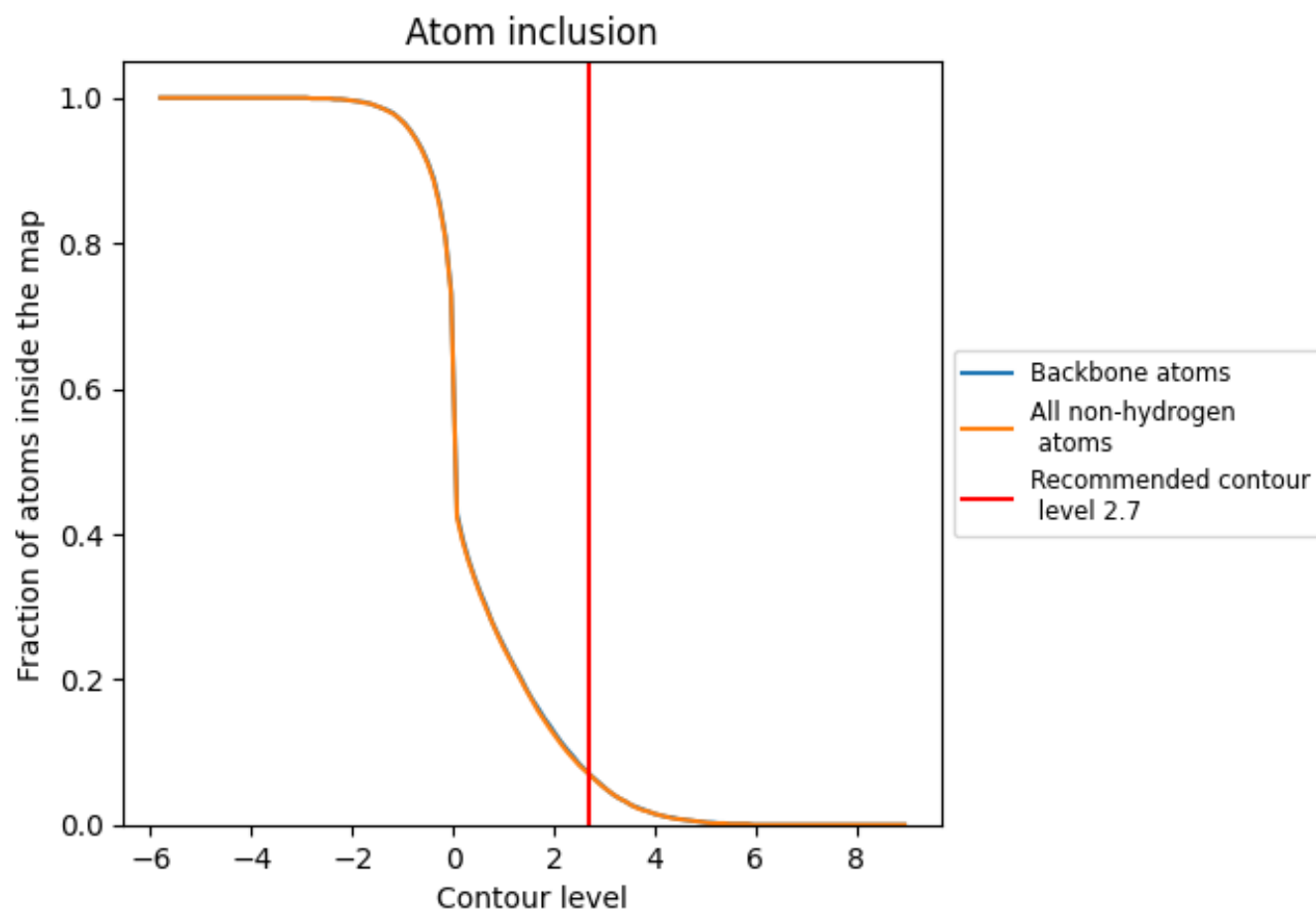


The images above show the model with each residue coloured according to its Q-score. This shows their resolvability in the map with higher Q-score values reflecting better resolvability. Please note: Q-score is calculating the resolvability of atoms, and thus high values are only expected at resolutions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

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

This section was not generated.

8.4 Atom inclusion [i](#)



At the recommended contour level, 7% of all backbone atoms, 7% of all non-hydrogen atoms, are inside the map.

8.5 Map-model fit summary ⓘ

The table lists the average atom inclusion at the recommended contour level (2.7) and Q-score for the entire model and for each chain.

Chain	Atom inclusion	Q-score
All	<div></div> 0.0690	<div></div> 0.0100
B	<div></div> 0.0010	<div></div> 0.0230
C	<div></div> 0.0530	<div></div> 0.0150
D	<div></div> 0.0450	<div></div> 0.0040
K	<div></div> 0.0870	<div></div> -0.0040
L	<div></div> 0.0870	<div></div> 0.0090
M	<div></div> 0.0830	<div></div> 0.0020
O	<div></div> 0.0910	<div></div> 0.0230
P	<div></div> 0.1510	<div></div> -0.0060
X	<div></div> 0.0720	<div></div> -0.0060
Y	<div></div> 0.0560	<div></div> -0.0050
Z	<div></div> 0.0830	<div></div> 0.0030

1.0
0.0
-0.0