



Full wwPDB EM Validation Report ⓘ

Apr 29, 2025 – 07:00 pm BST

PDB ID : 9GDY / pdb_00009gdy
EMDB ID : EMD-51280
Title : SARS-CoV-2 Spike protein Beta Variant at 37C structural flexibility / heterogeneity analyses
Authors : Herreros, D.; Mata, C.P.; Noddings, C.; Irene, D.; Agard, D.A.; Tsai, M.-D.; Sorzano, C.O.S.; Carazo, J.M.
Deposited on : 2024-08-06
Resolution : 2.80 Å (reported)
Based on initial models : 7VX1, 7WEV

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

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

A user guide is available at

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

with specific help available everywhere you see the ⓘ symbol.

The types of validation reports are described at

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

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

EMDB validation analysis : 0.0.1.dev118
MolProbity : 4-5-2 with Phenix2.0rc1
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) : 2.43.1

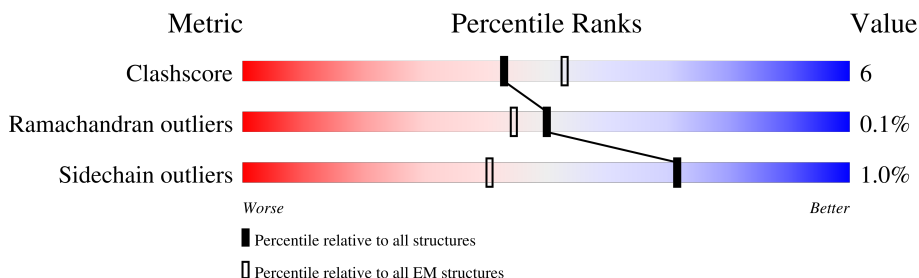
1 Overall quality at a glance

The following experimental techniques were used to determine the structure:

ELECTRON MICROSCOPY

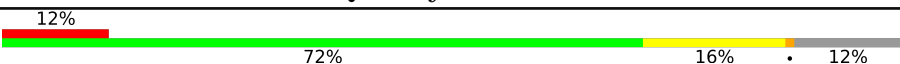
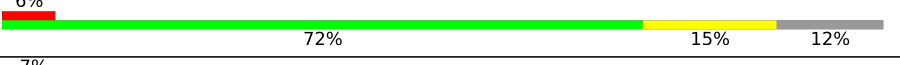

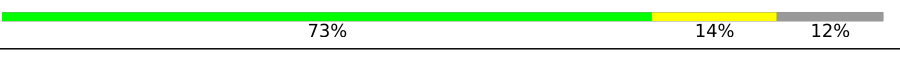
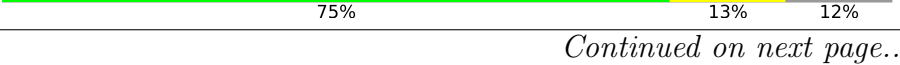

The reported resolution of this entry is 2.80 Å.

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	1-A	1230	
1	1-B	1230	
1	1-C	1230	
1	10-A	1230	
1	10-B	1230	
1	10-C	1230	
1	11-A	1230	
1	11-B	1230	


























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Mol	Chain	Length	Quality of chain
1	11-C	1230	
1	12-A	1230	
1	12-B	1230	
1	12-C	1230	
1	13-A	1230	
1	13-B	1230	
1	13-C	1230	
1	14-A	1230	
1	14-B	1230	
1	14-C	1230	
1	15-A	1230	
1	15-B	1230	
1	15-C	1230	
1	16-A	1230	
1	16-B	1230	
1	16-C	1230	
1	17-A	1230	
1	17-B	1230	
1	17-C	1230	
1	18-A	1230	
1	18-B	1230	
1	18-C	1230	
1	19-A	1230	
1	19-B	1230	
1	19-C	1230	



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Mol	Chain	Length	Quality of chain
1	2-A	1230	
1	2-B	1230	
1	2-C	1230	
1	20-A	1230	
1	20-B	1230	
1	20-C	1230	
1	3-A	1230	
1	3-B	1230	
1	3-C	1230	
1	4-A	1230	
1	4-B	1230	
1	4-C	1230	
1	5-A	1230	
1	5-B	1230	
1	5-C	1230	
1	6-A	1230	
1	6-B	1230	
1	6-C	1230	
1	7-A	1230	
1	7-B	1230	
1	7-C	1230	
1	8-A	1230	
1	8-B	1230	
1	8-C	1230	
1	9-A	1230	

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Mol	Chain	Length	Quality of chain
1	9-B	1230	 72% 16% 12%
1	9-C	1230	 71% 16% 12%

2 Entry composition

There is only 1 type of molecule in this entry. The entry contains 509160 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 Spike glycoprotein,Fibritin.

Mol	Chain	Residues	Atoms					AltConf	Trace
1	1-A	1079	Total	C	N	O	S	0	0
			8454	5399	1410	1606	39		
1	2-A	1079	Total	C	N	O	S	0	0
			8454	5399	1410	1606	39		
1	3-A	1085	Total	C	N	O	S	0	0
			8494	5425	1416	1615	38		
1	4-A	1085	Total	C	N	O	S	0	0
			8494	5425	1416	1615	38		
1	5-A	1085	Total	C	N	O	S	0	0
			8494	5425	1416	1615	38		
1	6-A	1085	Total	C	N	O	S	0	0
			8494	5425	1416	1615	38		
1	7-A	1085	Total	C	N	O	S	0	0
			8494	5425	1416	1615	38		
1	8-A	1085	Total	C	N	O	S	0	0
			8494	5425	1416	1615	38		
1	9-A	1079	Total	C	N	O	S	0	0
			8454	5399	1410	1606	39		
1	10-A	1085	Total	C	N	O	S	0	0
			8494	5425	1416	1615	38		
1	11-A	1079	Total	C	N	O	S	0	0
			8454	5399	1410	1606	39		
1	12-A	1085	Total	C	N	O	S	0	0
			8494	5425	1416	1615	38		
1	13-A	1085	Total	C	N	O	S	0	0
			8494	5425	1416	1615	38		
1	14-A	1085	Total	C	N	O	S	0	0
			8494	5425	1416	1615	38		
1	15-A	1085	Total	C	N	O	S	0	0
			8494	5425	1416	1615	38		
1	16-A	1085	Total	C	N	O	S	0	0
			8494	5425	1416	1615	38		
1	17-A	1085	Total	C	N	O	S	0	0
			8494	5425	1416	1615	38		

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Mol	Chain	Residues	Atoms					AltConf	Trace
1	18-A	1085	Total	C	N	O	S	0	0
			8494	5425	1416	1615	38		
1	19-A	1085	Total	C	N	O	S	0	0
			8494	5425	1416	1615	38		
1	20-A	1085	Total	C	N	O	S	0	0
			8494	5425	1416	1615	38		
1	1-B	1079	Total	C	N	O	S	0	0
			8454	5399	1410	1606	39		
1	2-B	1079	Total	C	N	O	S	0	0
			8454	5399	1410	1606	39		
1	3-B	1085	Total	C	N	O	S	0	0
			8494	5425	1416	1615	38		
1	4-B	1085	Total	C	N	O	S	0	0
			8494	5425	1416	1615	38		
1	5-B	1085	Total	C	N	O	S	0	0
			8494	5425	1416	1615	38		
1	6-B	1085	Total	C	N	O	S	0	0
			8494	5425	1416	1615	38		
1	7-B	1085	Total	C	N	O	S	0	0
			8494	5425	1416	1615	38		
1	8-B	1085	Total	C	N	O	S	0	0
			8494	5425	1416	1615	38		
1	9-B	1079	Total	C	N	O	S	0	0
			8454	5399	1410	1606	39		
1	10-B	1085	Total	C	N	O	S	0	0
			8494	5425	1416	1615	38		
1	11-B	1079	Total	C	N	O	S	0	0
			8454	5399	1410	1606	39		
1	12-B	1085	Total	C	N	O	S	0	0
			8494	5425	1416	1615	38		
1	13-B	1085	Total	C	N	O	S	0	0
			8494	5425	1416	1615	38		
1	14-B	1085	Total	C	N	O	S	0	0
			8494	5425	1416	1615	38		
1	15-B	1085	Total	C	N	O	S	0	0
			8494	5425	1416	1615	38		
1	16-B	1085	Total	C	N	O	S	0	0
			8494	5425	1416	1615	38		
1	17-B	1085	Total	C	N	O	S	0	0
			8494	5425	1416	1615	38		
1	18-B	1085	Total	C	N	O	S	0	0
			8494	5425	1416	1615	38		

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Mol	Chain	Residues	Atoms					AltConf	Trace
1	19-B	1085	Total	C	N	O	S	0	0
			8494	5425	1416	1615	38		
1	20-B	1085	Total	C	N	O	S	0	0
			8494	5425	1416	1615	38		
1	1-C	1079	Total	C	N	O	S	0	0
			8454	5399	1410	1606	39		
1	2-C	1079	Total	C	N	O	S	0	0
			8454	5399	1410	1606	39		
1	3-C	1085	Total	C	N	O	S	0	0
			8494	5425	1416	1615	38		
1	4-C	1085	Total	C	N	O	S	0	0
			8494	5425	1416	1615	38		
1	5-C	1085	Total	C	N	O	S	0	0
			8494	5425	1416	1615	38		
1	6-C	1085	Total	C	N	O	S	0	0
			8494	5425	1416	1615	38		
1	7-C	1085	Total	C	N	O	S	0	0
			8494	5425	1416	1615	38		
1	8-C	1085	Total	C	N	O	S	0	0
			8494	5425	1416	1615	38		
1	9-C	1079	Total	C	N	O	S	0	0
			8454	5399	1410	1606	39		
1	10-C	1085	Total	C	N	O	S	0	0
			8494	5425	1416	1615	38		
1	11-C	1079	Total	C	N	O	S	0	0
			8454	5399	1410	1606	39		
1	12-C	1085	Total	C	N	O	S	0	0
			8494	5425	1416	1615	38		
1	13-C	1085	Total	C	N	O	S	0	0
			8494	5425	1416	1615	38		
1	14-C	1085	Total	C	N	O	S	0	0
			8494	5425	1416	1615	38		
1	15-C	1085	Total	C	N	O	S	0	0
			8494	5425	1416	1615	38		
1	16-C	1085	Total	C	N	O	S	0	0
			8494	5425	1416	1615	38		
1	17-C	1085	Total	C	N	O	S	0	0
			8494	5425	1416	1615	38		
1	18-C	1085	Total	C	N	O	S	0	0
			8494	5425	1416	1615	38		
1	19-C	1085	Total	C	N	O	S	0	0
			8494	5425	1416	1615	38		

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Mol	Chain	Residues	Atoms					AltConf	Trace
1	20-C	1085	Total	C	N	O	S	0	0
			8494	5425	1416	1615	38		

There are 87 discrepancies between the modelled and reference sequences:

Chain	Residue	Modelled	Actual	Comment	Reference
A	18	PHE	LEU	variant	UNP P0DTC2
A	80	ALA	ASP	variant	UNP P0DTC2
A	215	GLY	ASP	variant	UNP P0DTC2
A	?	-	LEU	deletion	UNP P0DTC2
A	?	-	ALA	deletion	UNP P0DTC2
A	?	-	LEU	deletion	UNP P0DTC2
A	246	ILE	ARG	conflict	UNP P0DTC2
A	417	ASN	LYS	variant	UNP P0DTC2
A	484	LYS	GLU	variant	UNP P0DTC2
A	501	TYR	ASN	variant	UNP P0DTC2
A	614	GLY	ASP	variant	UNP P0DTC2
A	682	GLY	ARG	engineered mutation	UNP P0DTC2
A	683	SER	ARG	engineered mutation	UNP P0DTC2
A	685	SER	ARG	engineered mutation	UNP P0DTC2
A	701	VAL	ALA	variant	UNP P0DTC2
A	986	PRO	LYS	engineered mutation	UNP P0DTC2
A	987	PRO	VAL	engineered mutation	UNP P0DTC2
A	1209	GLY	-	linker	UNP P0DTC2
A	1210	SER	-	linker	UNP P0DTC2
A	1232	LEU	PHE	engineered mutation	UNP P10104
A	1238	GLY	-	expression tag	UNP P10104
A	1239	ARG	-	expression tag	UNP P10104
A	1240	SER	-	expression tag	UNP P10104
A	1241	LEU	-	expression tag	UNP P10104
A	1242	GLU	-	expression tag	UNP P10104
A	1243	VAL	-	expression tag	UNP P10104
A	1244	LEU	-	expression tag	UNP P10104
A	1245	PHE	-	expression tag	UNP P10104
A	1246	GLN	-	expression tag	UNP P10104
B	18	PHE	LEU	variant	UNP P0DTC2
B	80	ALA	ASP	variant	UNP P0DTC2
B	215	GLY	ASP	variant	UNP P0DTC2
B	?	-	LEU	deletion	UNP P0DTC2
B	?	-	ALA	deletion	UNP P0DTC2
B	?	-	LEU	deletion	UNP P0DTC2
B	246	ILE	ARG	conflict	UNP P0DTC2

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Chain	Residue	Modelled	Actual	Comment	Reference
B	417	ASN	LYS	variant	UNP P0DTC2
B	484	LYS	GLU	variant	UNP P0DTC2
B	501	TYR	ASN	variant	UNP P0DTC2
B	614	GLY	ASP	variant	UNP P0DTC2
B	682	GLY	ARG	engineered mutation	UNP P0DTC2
B	683	SER	ARG	engineered mutation	UNP P0DTC2
B	685	SER	ARG	engineered mutation	UNP P0DTC2
B	701	VAL	ALA	variant	UNP P0DTC2
B	986	PRO	LYS	engineered mutation	UNP P0DTC2
B	987	PRO	VAL	engineered mutation	UNP P0DTC2
B	1209	GLY	-	linker	UNP P0DTC2
B	1210	SER	-	linker	UNP P0DTC2
B	1232	LEU	PHE	engineered mutation	UNP P10104
B	1238	GLY	-	expression tag	UNP P10104
B	1239	ARG	-	expression tag	UNP P10104
B	1240	SER	-	expression tag	UNP P10104
B	1241	LEU	-	expression tag	UNP P10104
B	1242	GLU	-	expression tag	UNP P10104
B	1243	VAL	-	expression tag	UNP P10104
B	1244	LEU	-	expression tag	UNP P10104
B	1245	PHE	-	expression tag	UNP P10104
B	1246	GLN	-	expression tag	UNP P10104
C	18	PHE	LEU	variant	UNP P0DTC2
C	80	ALA	ASP	variant	UNP P0DTC2
C	215	GLY	ASP	variant	UNP P0DTC2
C	?	-	LEU	deletion	UNP P0DTC2
C	?	-	ALA	deletion	UNP P0DTC2
C	?	-	LEU	deletion	UNP P0DTC2
C	246	ILE	ARG	conflict	UNP P0DTC2
C	417	ASN	LYS	variant	UNP P0DTC2
C	484	LYS	GLU	variant	UNP P0DTC2
C	501	TYR	ASN	variant	UNP P0DTC2
C	614	GLY	ASP	variant	UNP P0DTC2
C	682	GLY	ARG	engineered mutation	UNP P0DTC2
C	683	SER	ARG	engineered mutation	UNP P0DTC2
C	685	SER	ARG	engineered mutation	UNP P0DTC2
C	701	VAL	ALA	variant	UNP P0DTC2
C	986	PRO	LYS	engineered mutation	UNP P0DTC2
C	987	PRO	VAL	engineered mutation	UNP P0DTC2
C	1209	GLY	-	linker	UNP P0DTC2
C	1210	SER	-	linker	UNP P0DTC2
C	1232	LEU	PHE	engineered mutation	UNP P10104

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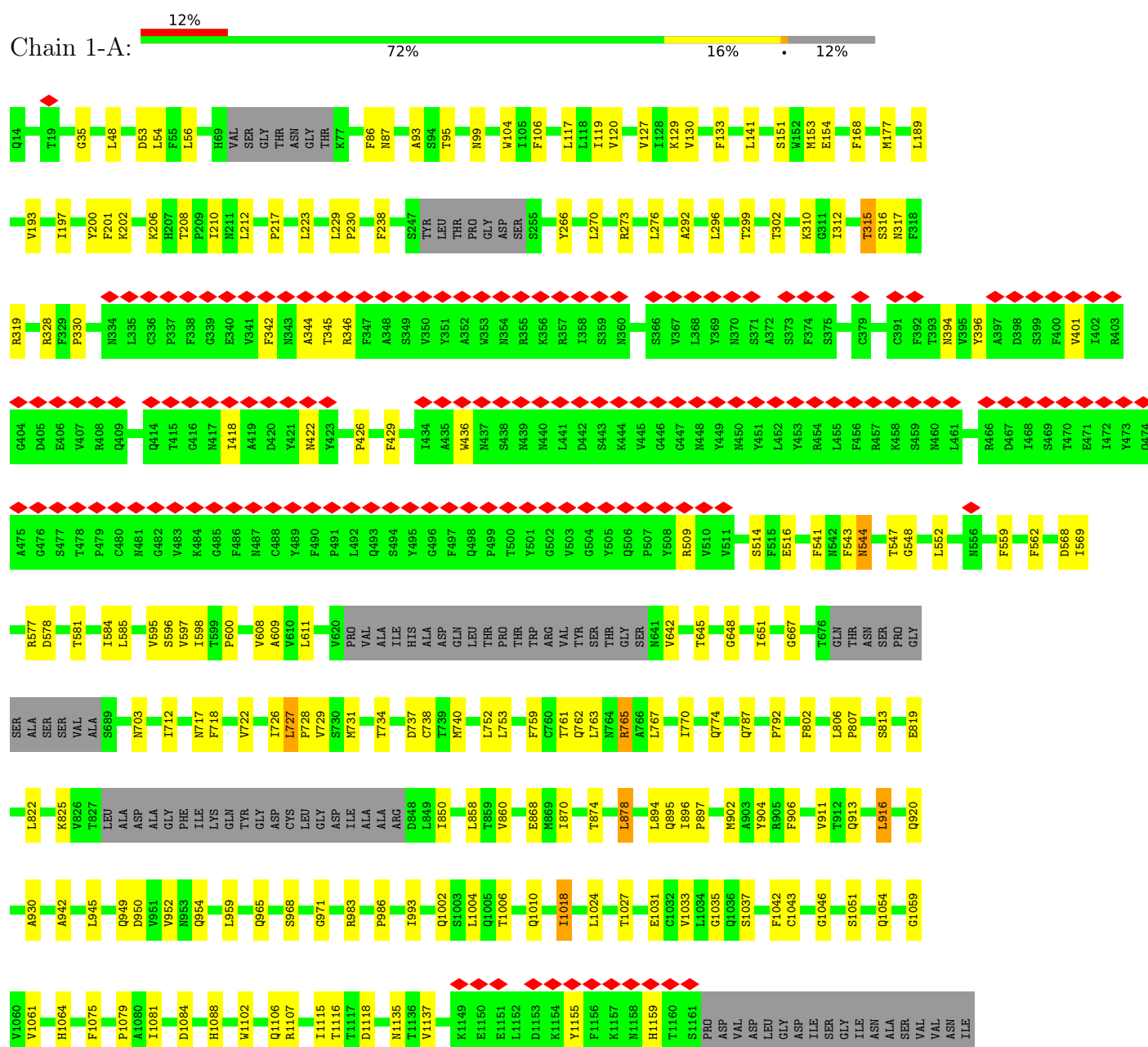
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Chain	Residue	Modelled	Actual	Comment	Reference
C	1238	GLY	-	expression tag	UNP P10104
C	1239	ARG	-	expression tag	UNP P10104
C	1240	SER	-	expression tag	UNP P10104
C	1241	LEU	-	expression tag	UNP P10104
C	1242	GLU	-	expression tag	UNP P10104
C	1243	VAL	-	expression tag	UNP P10104
C	1244	LEU	-	expression tag	UNP P10104
C	1245	PHE	-	expression tag	UNP P10104
C	1246	GLN	-	expression tag	UNP P10104

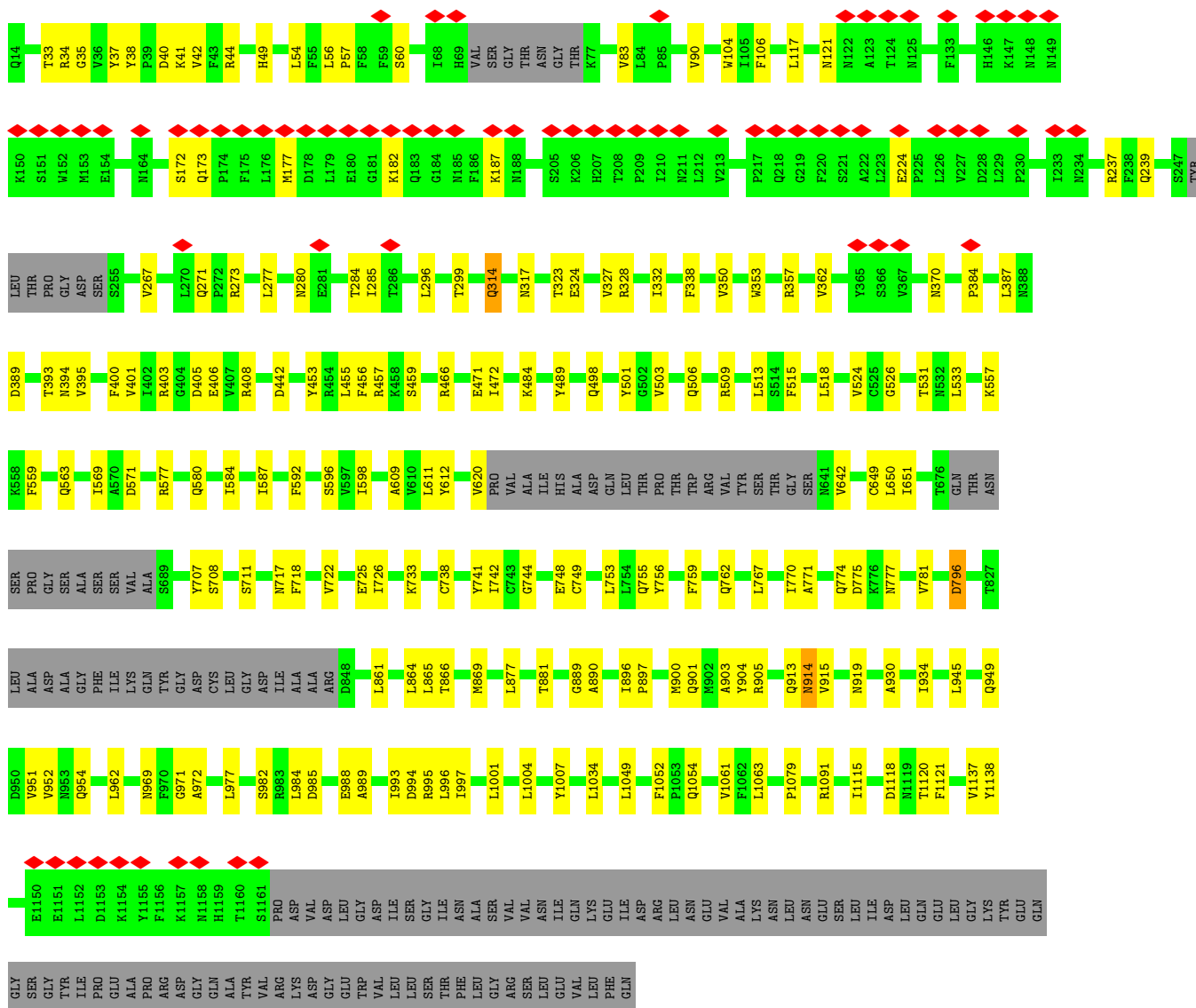
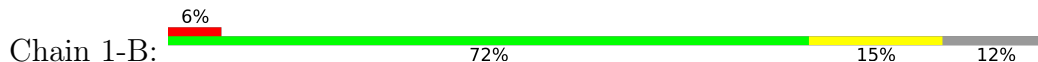
3 Residue-property plots

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.

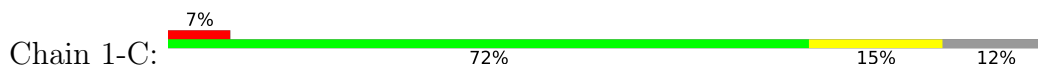
- Molecule 1: Spike glycoprotein,Fibritin

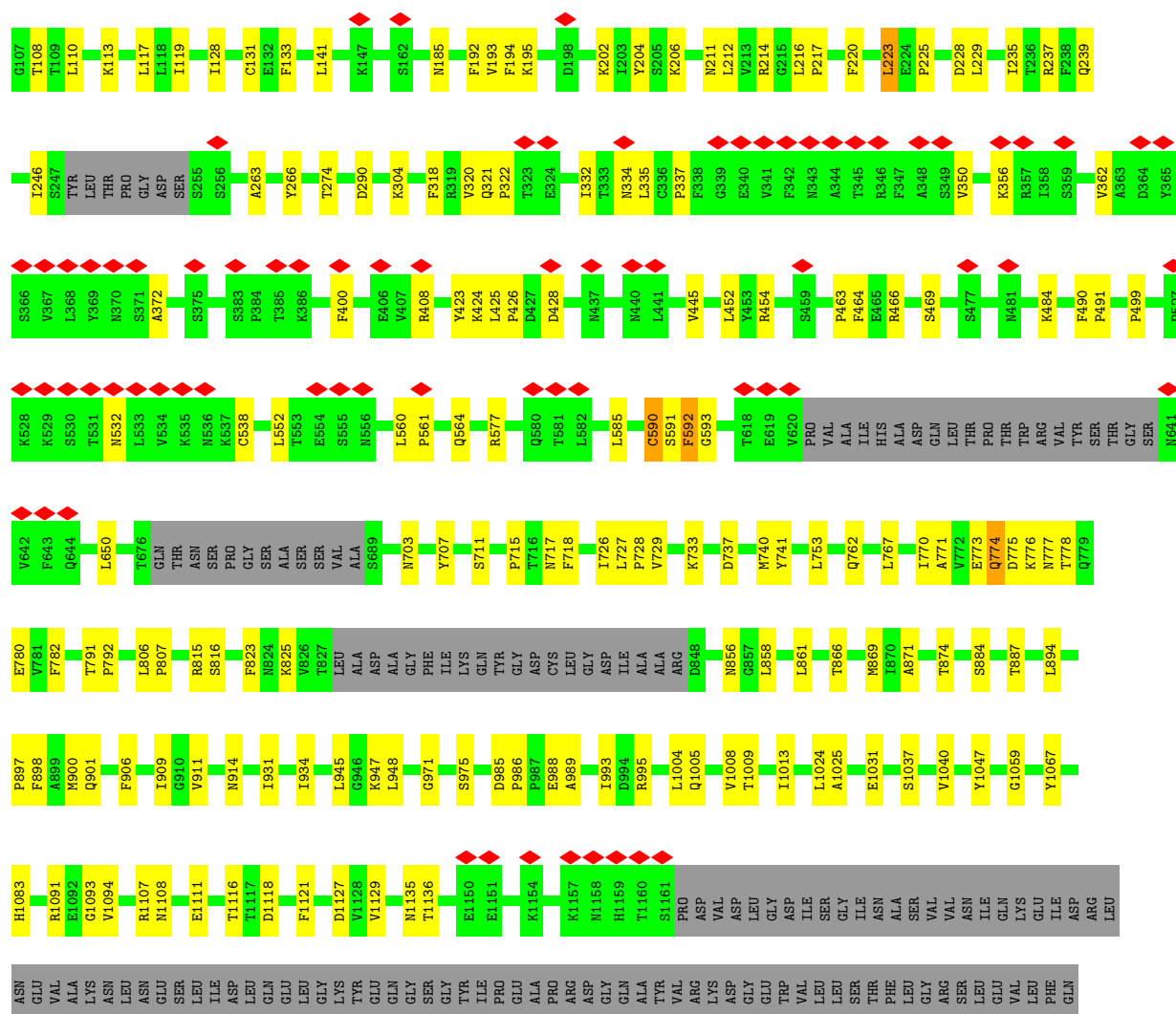


- Molecule 1: Spike glycoprotein, Fibrin



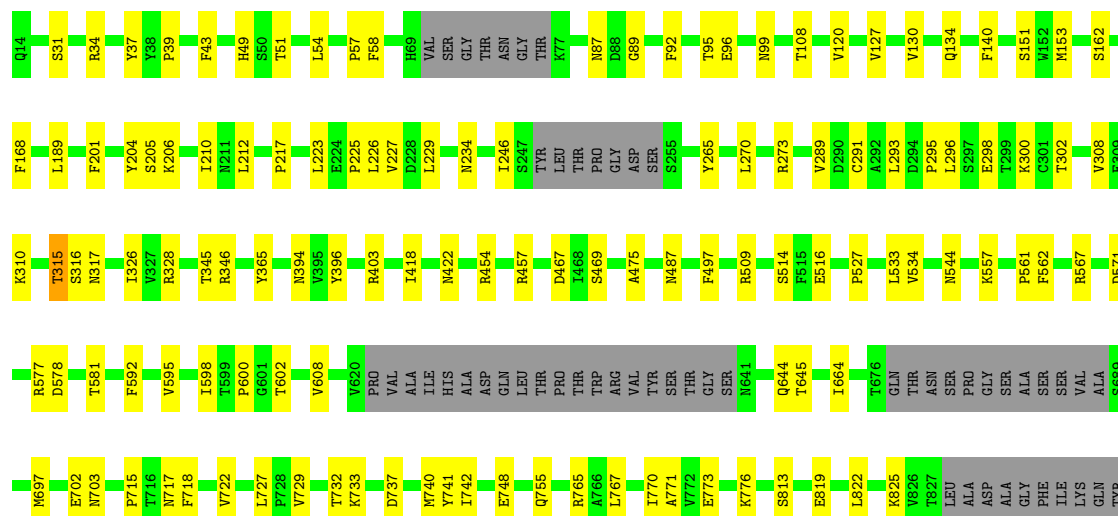
- Molecule 1: Spike glycoprotein, Fibrin

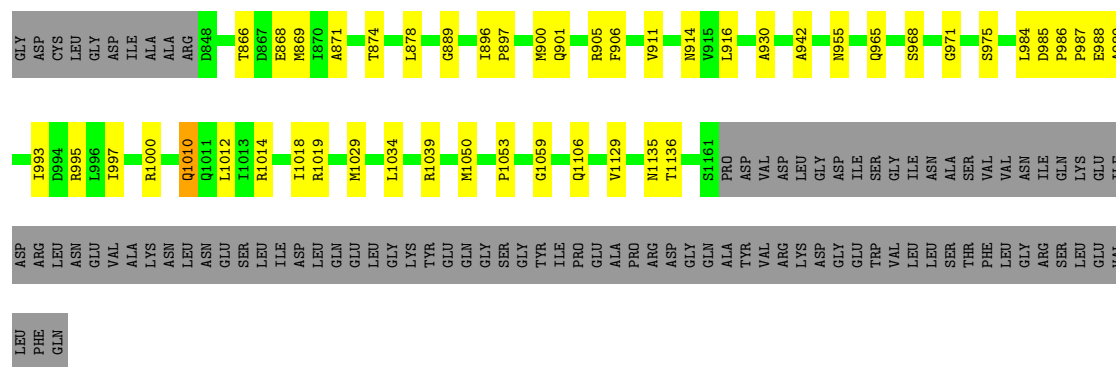




• Molecule 1: Spike glycoprotein,Fibrin

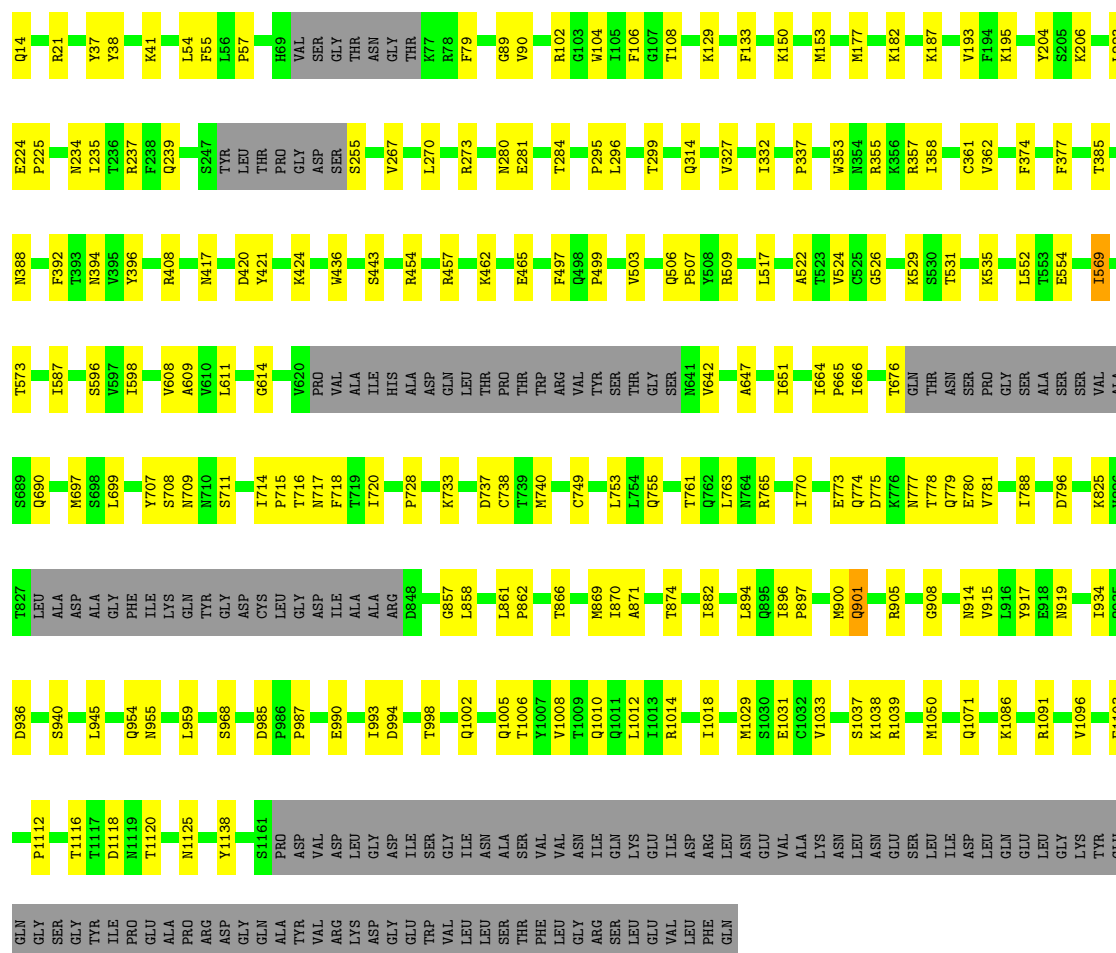
Chain 2-A: 74% 14% 12%





• Molecule 1: Spike glycoprotein,Fibrin

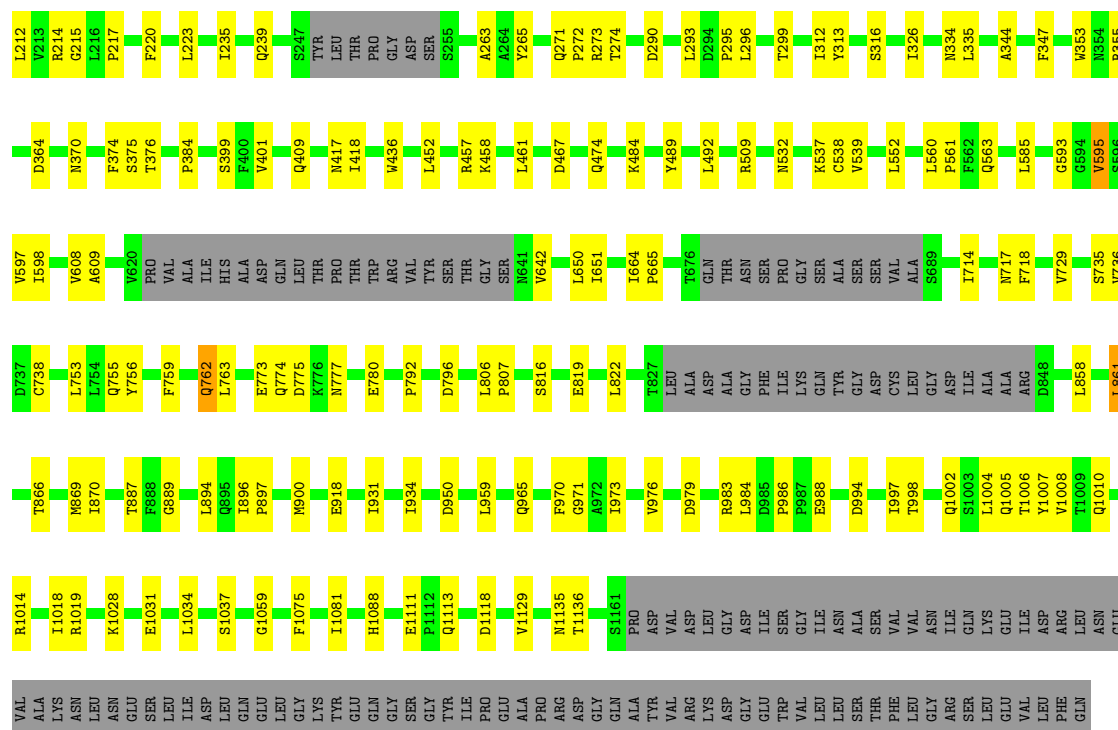
Chain 2-B: 72% 16% 12%



• Molecule 1: Spike glycoprotein,Fibrin

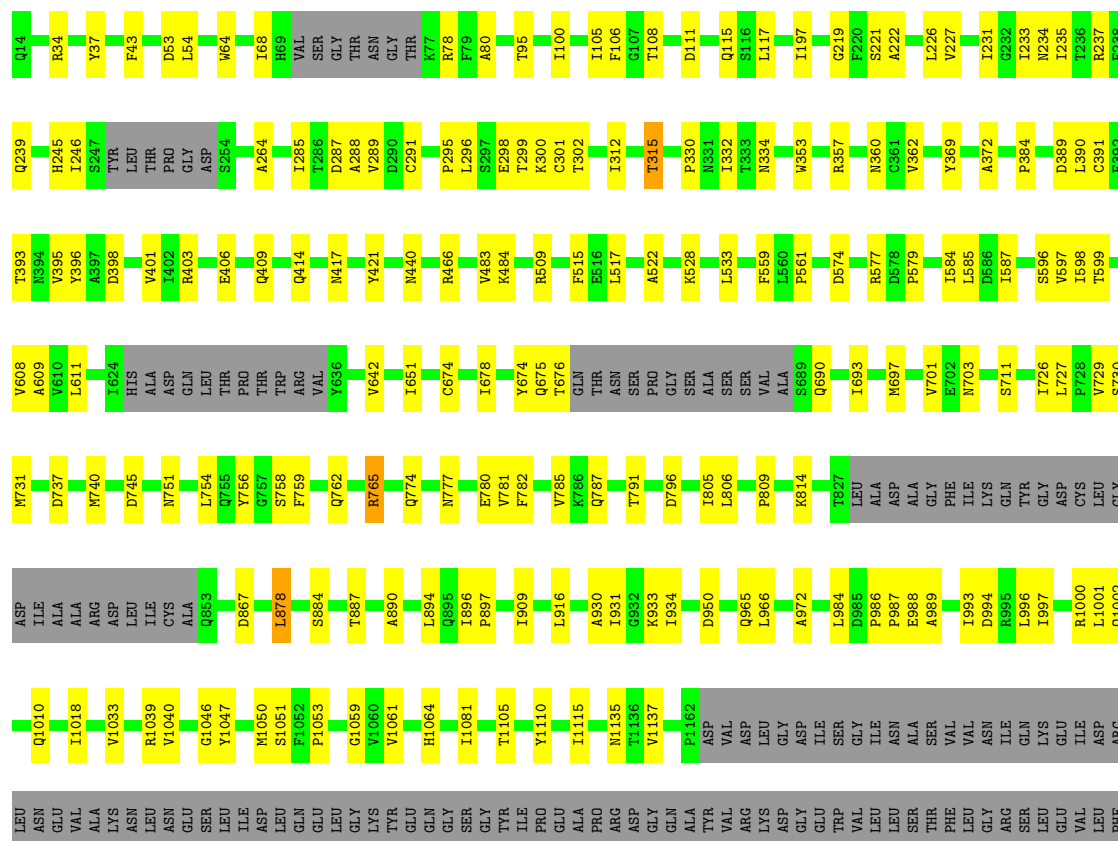
Chain 2-C: 73% 14% 12%

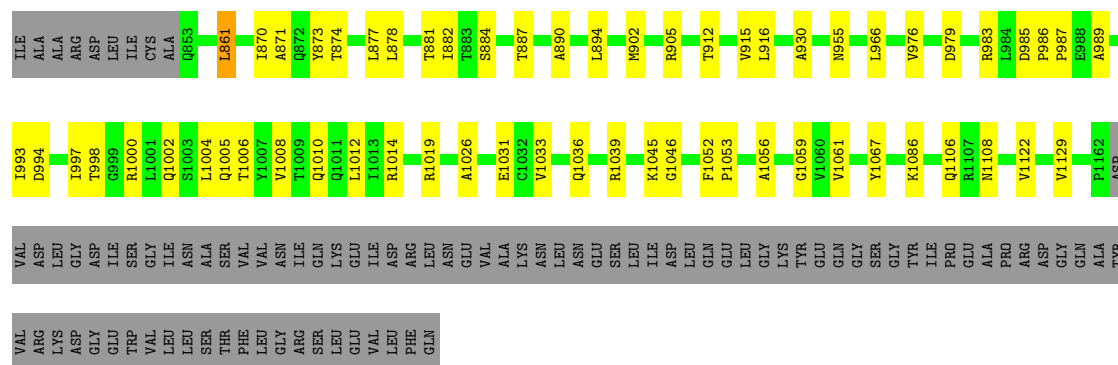




• Molecule 1: Spike glycoprotein,Fibritin

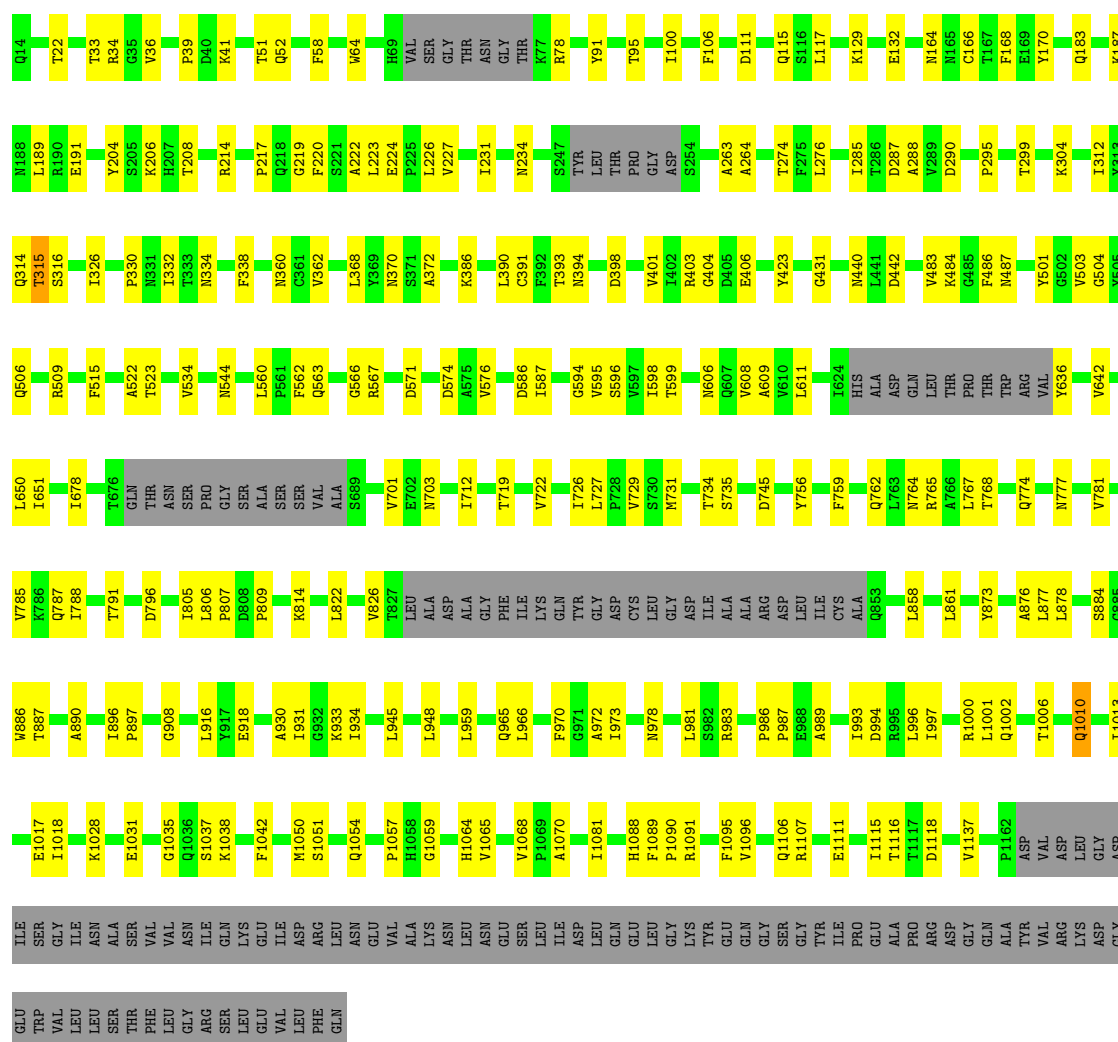
Chain 3-A: 73% 15% 12%





• Molecule 1: Spike glycoprotein,Fibrin

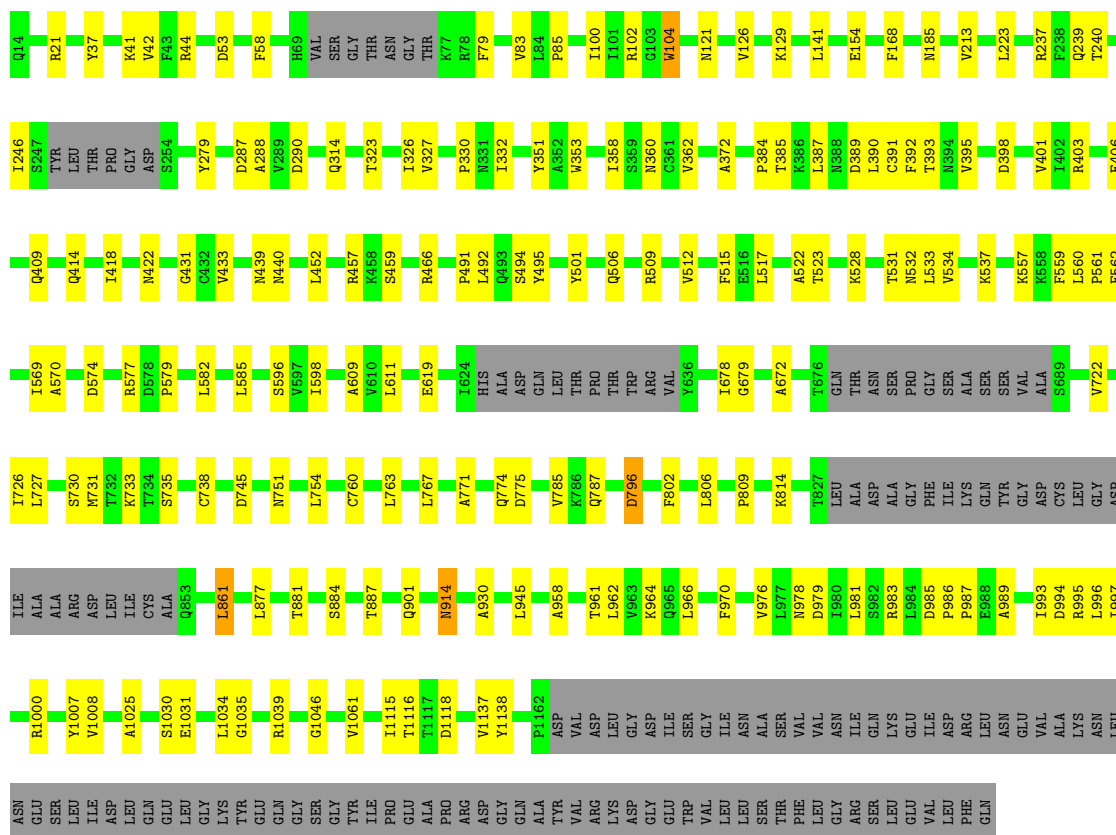
Chain 4-A: 70% 18% 12%



• Molecule 1: Spike glycoprotein,Fibrin

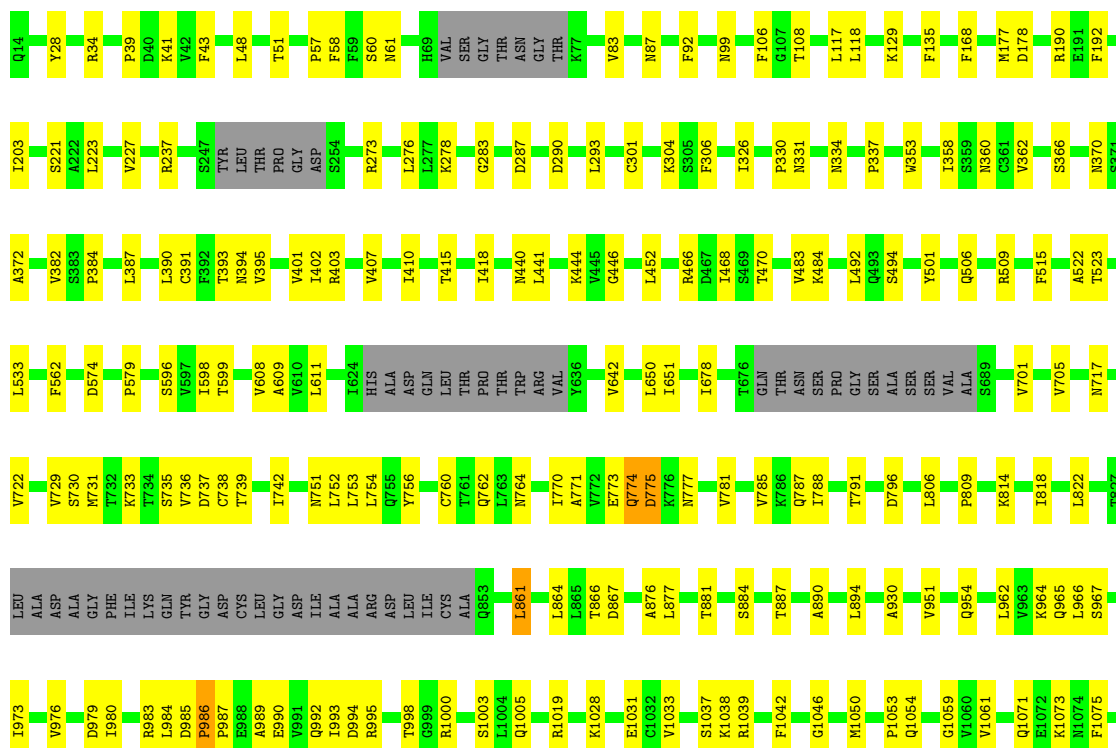
Chain 4-B: 70% 18% 12%

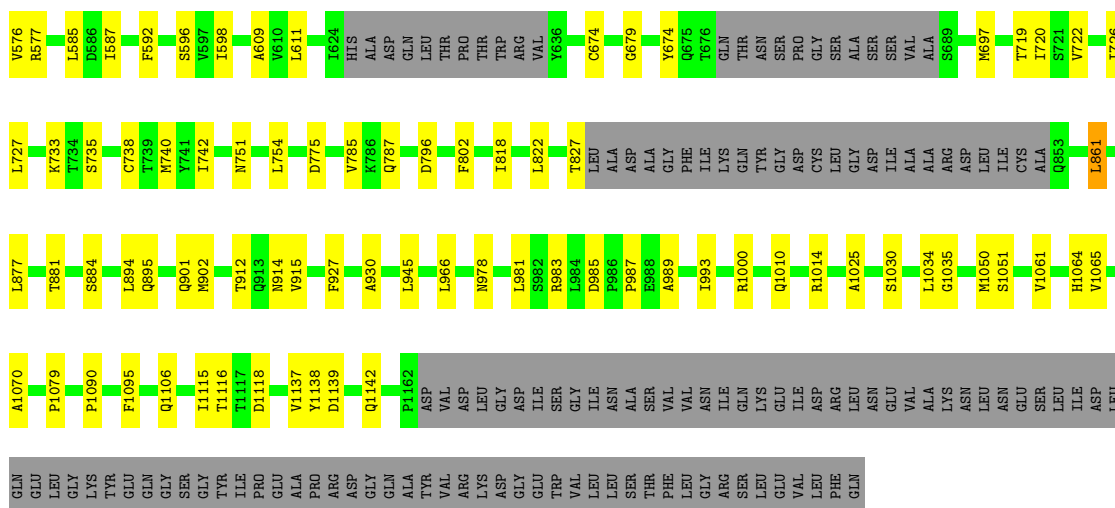




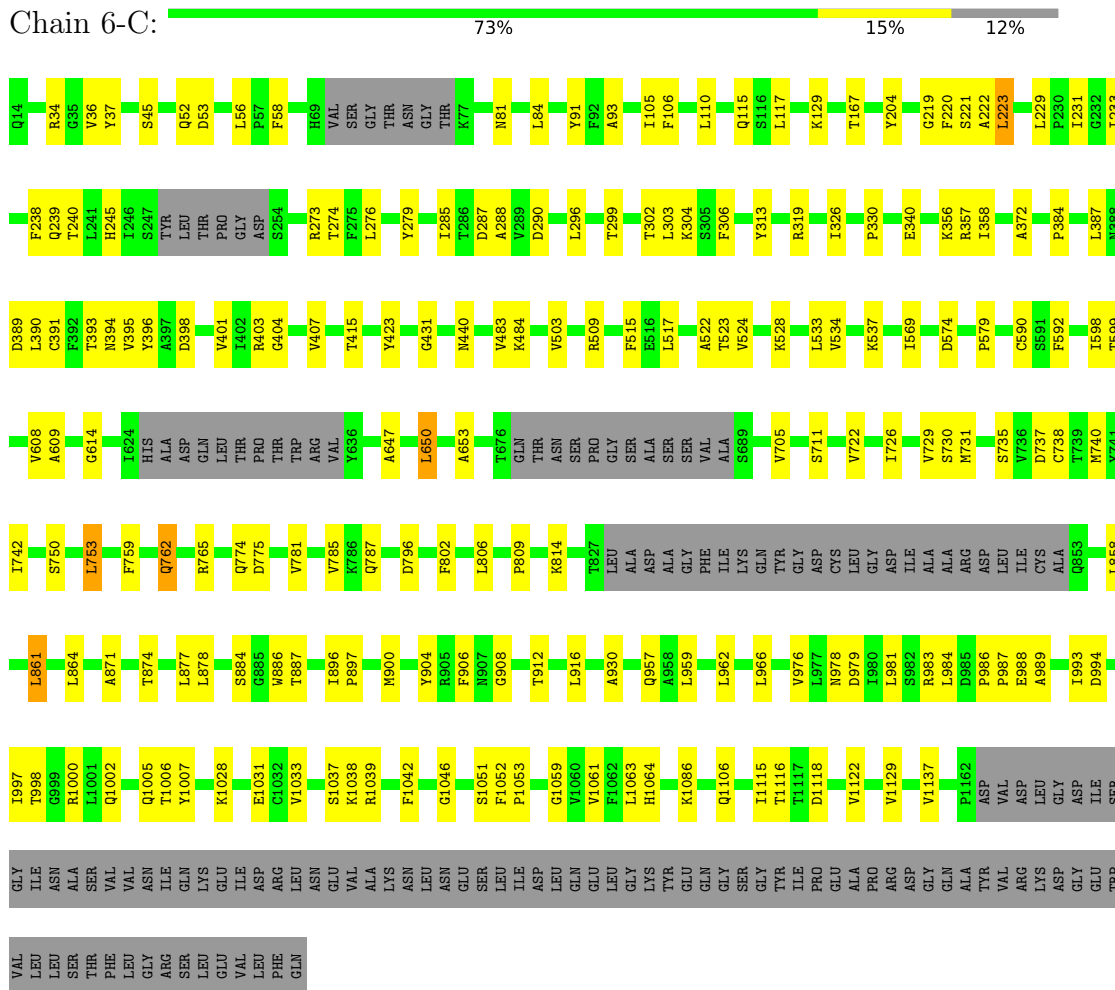
• Molecule 1: Spike glycoprotein,Fibrin

Chain 5-C: 72% 16% 12%



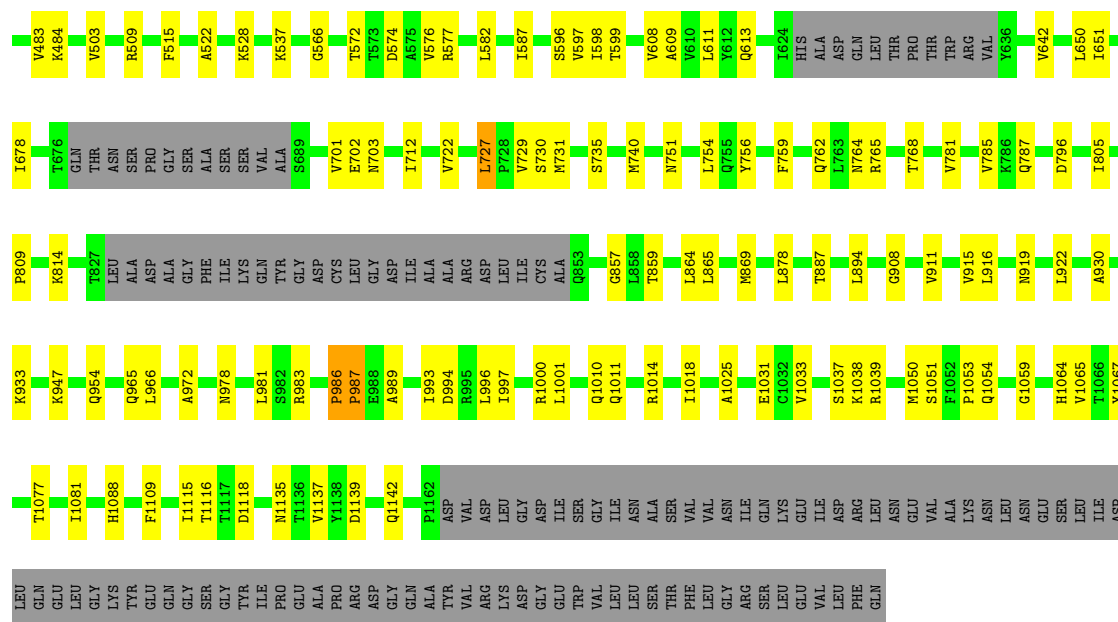


- Molecule 1: Spike glycoprotein, Fibrin



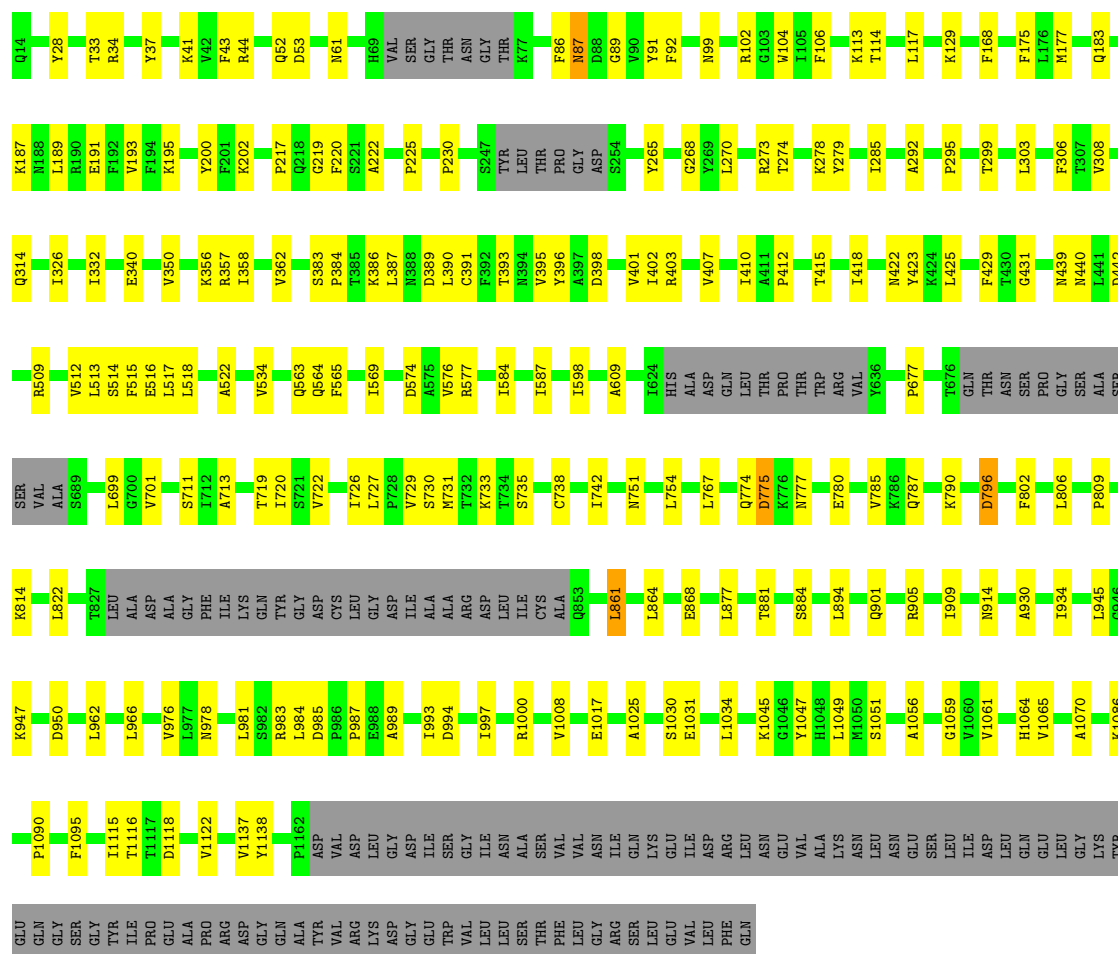
- Molecule 1: Spike glycoprotein, Fibrin



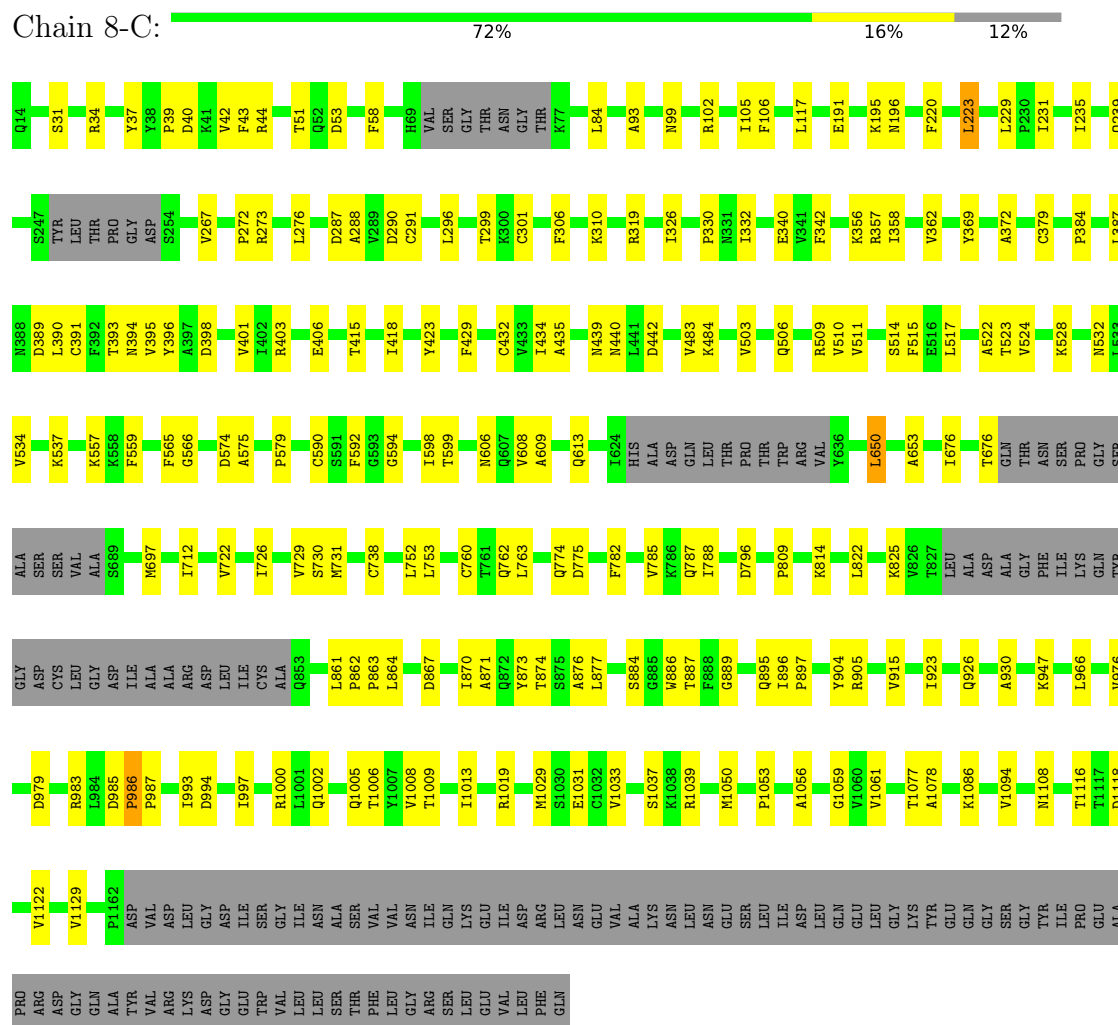


● Molecule 1: Spike glycoprotein,Fibrin

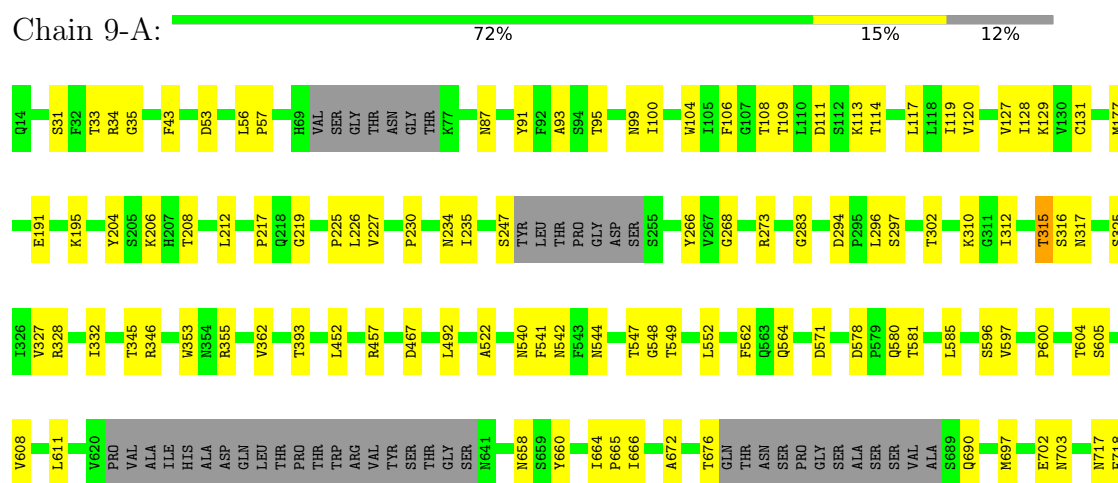
Chain 8-B: 72% 16% 12%

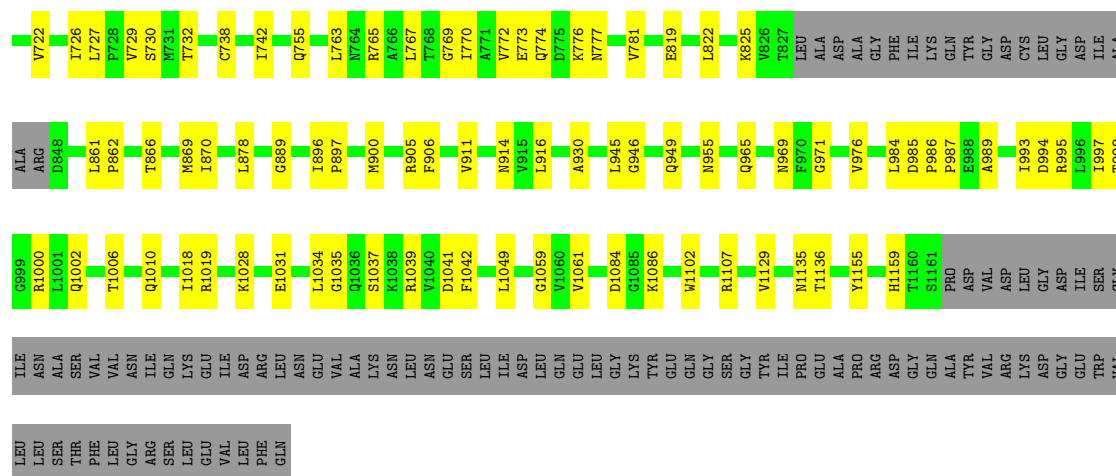


• Molecule 1: Spike glycoprotein,Fibrinin



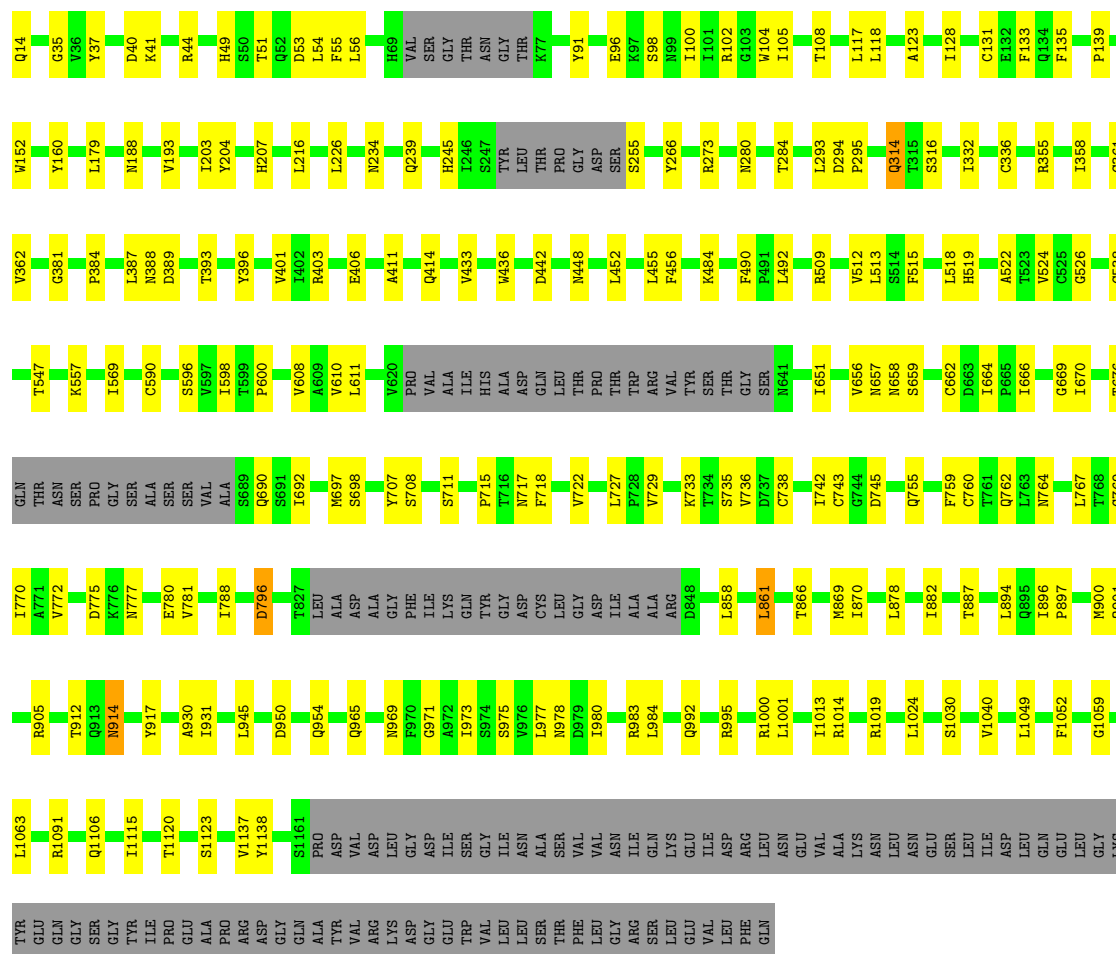
• Molecule 1: Spike glycoprotein,Fibrinin





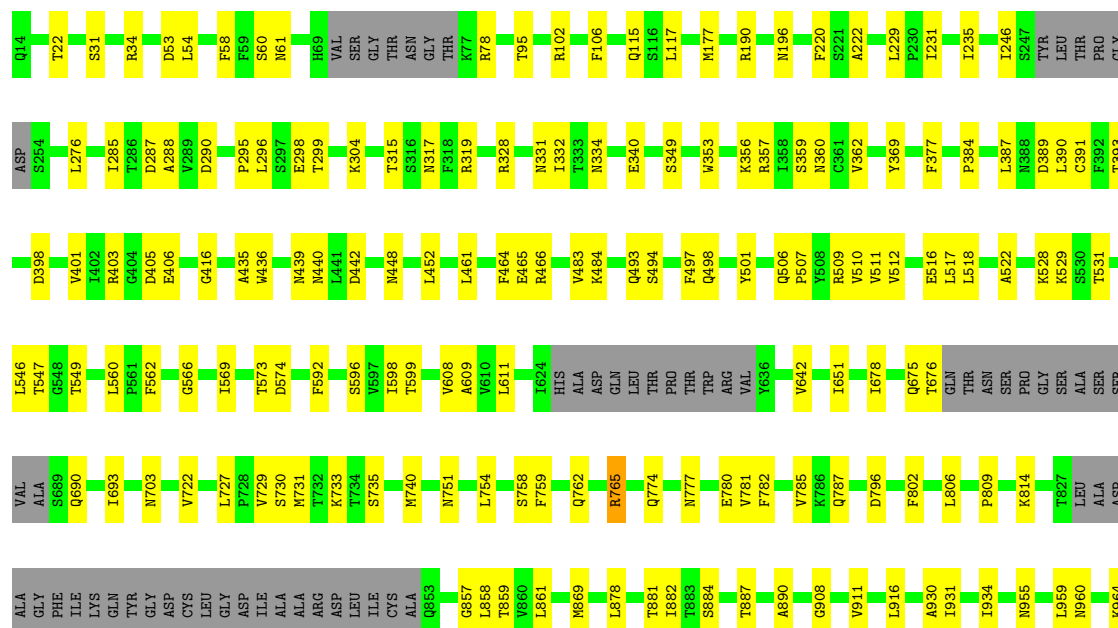
• Molecule 1: Spike glycoprotein,Fibrin

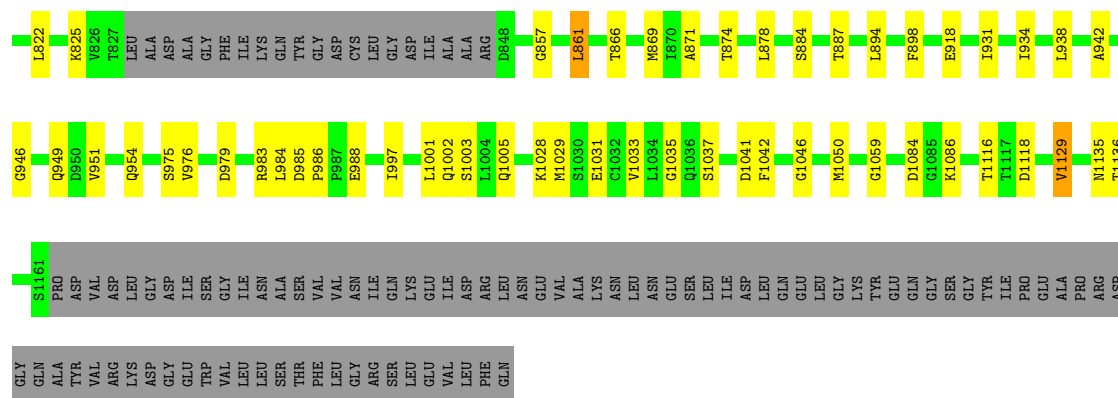
Chain 9-B: 72% 16% 12%



• Molecule 1: Spike glycoprotein,Fibrin

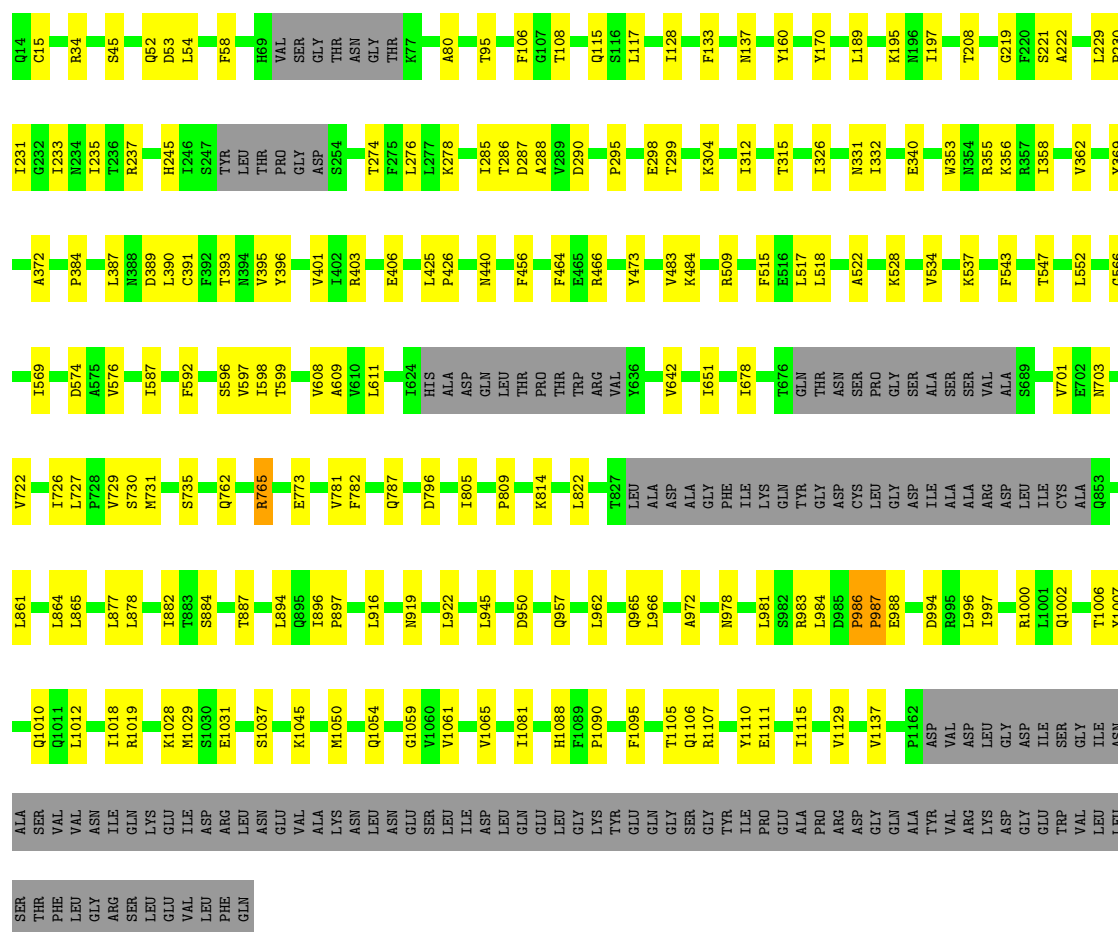
Chain 9-C: 71% 16% 12%





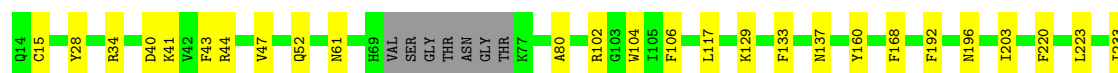
• Molecule 1: Spike glycoprotein, Fibrin

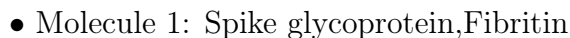
Chain 12-A: 73% 15% 12%



• Molecule 1: Spike glycoprotein, Fibrin

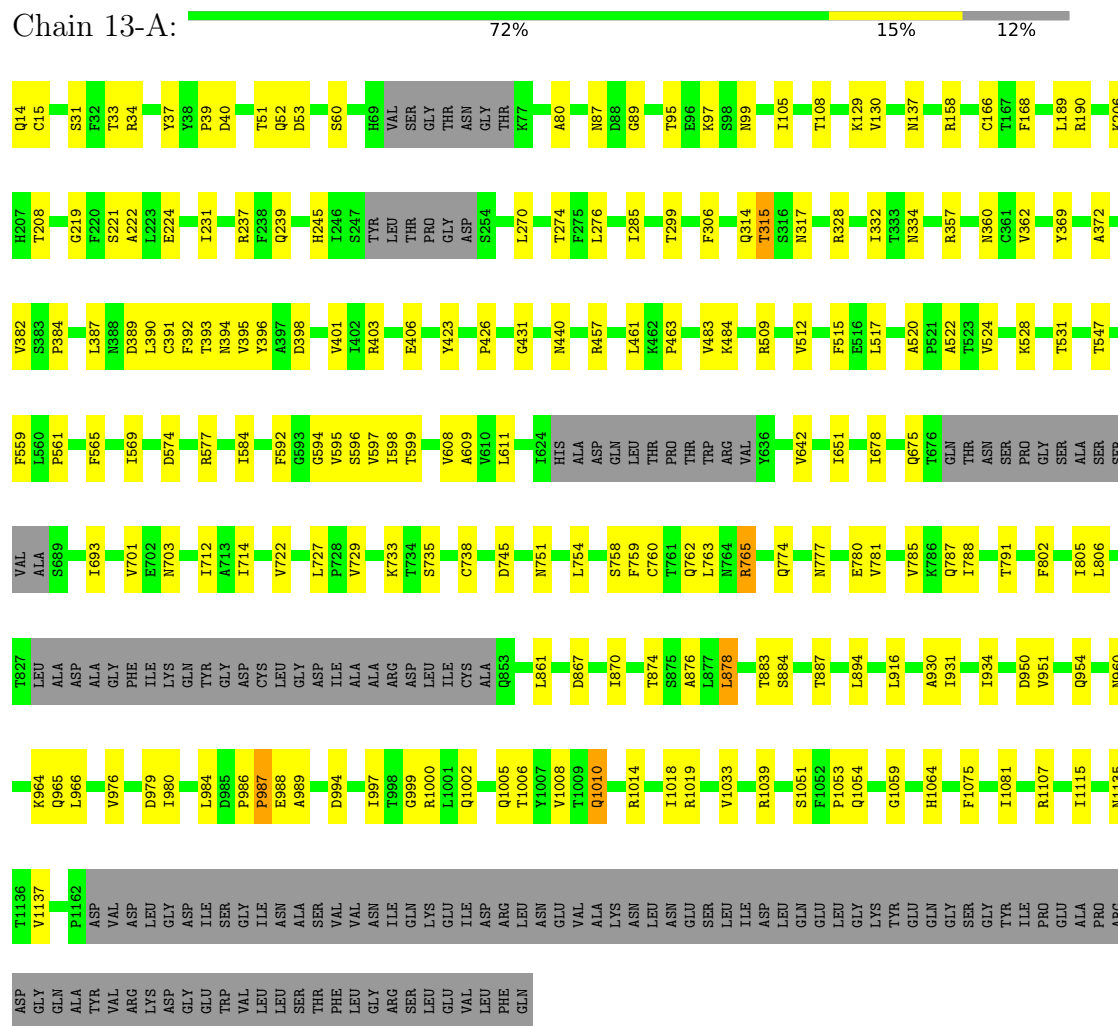
Chain 12-B: 72% 16% 12%



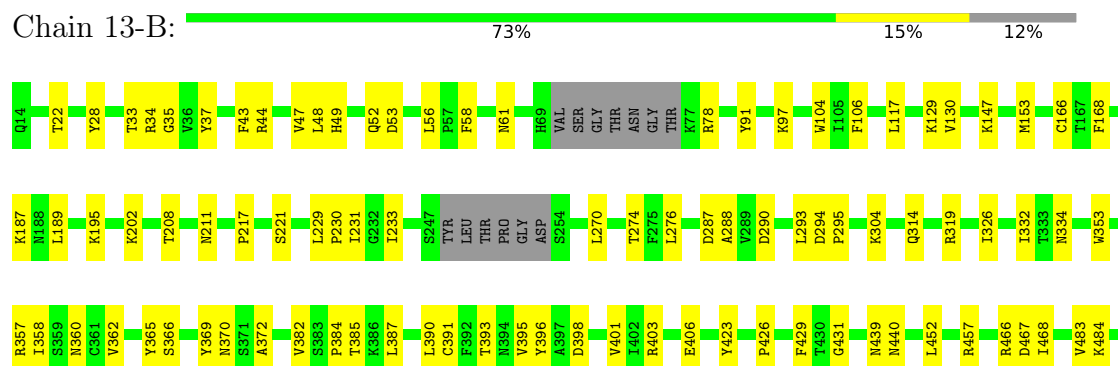


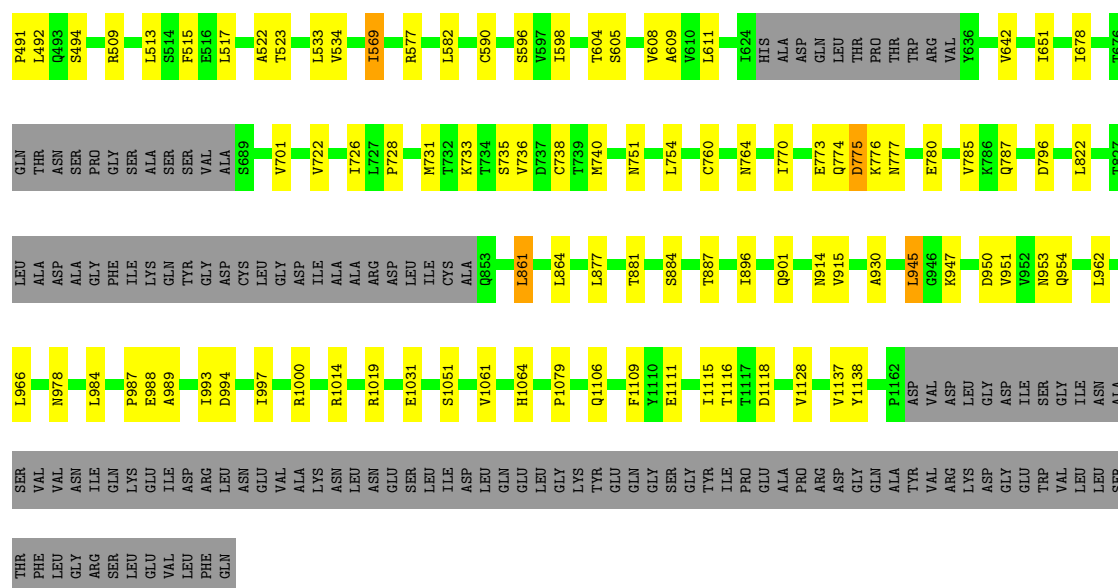
Q872	Q873	Q874	Q875	Q876	Q877	Q878	Q879	Q880	Q881	Q882	Q883	Q884	Q885	Q886	Q887	Q888	Q889	Q890	Q891	Q892	Q893	Q894	Q895	Q896	Q897	Q898	Q899	Q900	Q901	Q902	Q903	Q904	Q905	Q906	Q907	Q908	Q909	Q910	Q911	Q912	Q913	Q914	Q915	Q916	Q917	Q918	Q919	Q920	Q921	Q922	Q923	Q924	Q925	Q926	Q927	Q928	Q929	Q930	Q931	Q932	Q933	Q934	Q935	Q936	Q937	Q938	Q939	Q940	Q941	Q942	Q943	Q944	Q945	Q946	Q947	Q948	Q949	Q950	Q951	Q952	Q953	Q954	Q955	Q956	Q957	Q958	Q959	Q960	Q961	Q962	Q963	Q964	Q965	Q966	Q967	Q968	Q969	Q970	Q971	Q972	Q973	Q974	Q975	Q976	Q977	Q978	Q979	Q980	Q981	Q982	Q983	Q984	Q985	Q986	Q987	Q988	Q989	Q990	Q991	Q992	Q993	Q994	Q995	Q996	Q997	Q998	Q999	Q1000	Q1001	Q1002	Q1003	Q1004	Q1005	Q1006	Q1007	Q1008	Q1009	Q1010	Q1011	Q1012	Q1013	Q1014	Q1015	Q1016	Q1017	Q1018	Q1019	Q1020	Q1021	Q1022	Q1023	Q1024	Q1025	Q1026	Q1027	Q1028	Q1029	Q1030	Q1031	Q1032	Q1033	Q1034	Q1035	Q1036	Q1037	Q1038	Q1039	Q1040	Q1041	Q1042	Q1043	Q1044	Q1045	Q1046	Q1047	Q1048	Q1049	Q1050	Q1051	Q1052	Q1053	Q1054	Q1055	Q1056	Q1057	Q1058	Q1059	Q1060	Q1061	Q1062	Q1063	Q1064	Q1065	Q1066	Q1067	Q1068	Q1069	Q1070	Q1071	Q1072	Q1073	Q1074	Q1075	Q1076	Q1077	Q1078	Q1079	Q1080	Q1081	Q1082	Q1083	Q1084	Q1085	Q1086	Q1087	Q1088	Q1089	Q1090	Q1091	Q1092	Q1093	Q1094	Q1095	Q1096	Q1097	Q1098	Q1099	Q1100	Q1101	Q1102	Q1103	Q1104	Q1105	Q1106	Q1107	Q1108	Q1109	Q1110	Q1111	Q1112	Q1113	Q1114	Q1115	Q1116	Q1117	Q1118	Q1119	Q1120	Q1121	Q1122	Q1123	Q1124	Q1125	Q1126	Q1127	Q1128	Q1129	Q1130	Q1131	Q1132	Q1133	Q1134	Q1135	Q1136	Q1137	Q1138	Q1139	Q1140	Q1141	Q1142	Q1143	Q1144	Q1145	Q1146	Q1147	Q1148	Q1149	Q1150	Q1151	Q1152	Q1153	Q1154	Q1155	Q1156	Q1157	Q1158	Q1159	Q1160	Q1161	Q1162	Q1163	Q1164	Q1165	Q1166	Q1167	Q1168	Q1169	Q1170	Q1171	Q1172	Q1173	Q1174	Q1175	Q1176	Q1177	Q1178	Q1179	Q1180	Q1181	Q1182	Q1183	Q1184	Q1185	Q1186	Q1187	Q1188	Q1189	Q1190	Q1191	Q1192	Q1193	Q1194	Q1195	Q1196	Q1197	Q1198	Q1199	Q1200	Q1201	Q1202	Q1203	Q1204	Q1205	Q1206	Q1207	Q1208	Q1209	Q1210	Q1211	Q1212	Q1213	Q1214	Q1215	Q1216	Q1217	Q1218	Q1219	Q1220	Q1221	Q1222	Q1223	Q1224	Q1225	Q1226	Q1227	Q1228	Q1229	Q1230	Q1231	Q1232	Q1233	Q1234	Q1235	Q1236	Q1237	Q1238	Q1239	Q1240	Q1241	Q1242	Q1243	Q1244	Q1245	Q1246	Q1247	Q1248	Q1249	Q1250	Q1251	Q1252	Q1253	Q1254	Q1255	Q1256	Q1257	Q1258	Q1259	Q1260	Q1261	Q1262	Q1263	Q1264	Q1265	Q1266	Q1267	Q1268	Q1269	Q1270	Q1271	Q1272	Q1273	Q1274	Q1275	Q1276	Q1277	Q1278	Q1279	Q1280	Q1281	Q1282	Q1283	Q1284	Q1285	Q1286	Q1287	Q1288	Q1289	Q1290	Q1291	Q1292	Q1293
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- Molecule 1: Spike glycoprotein, Fibrin



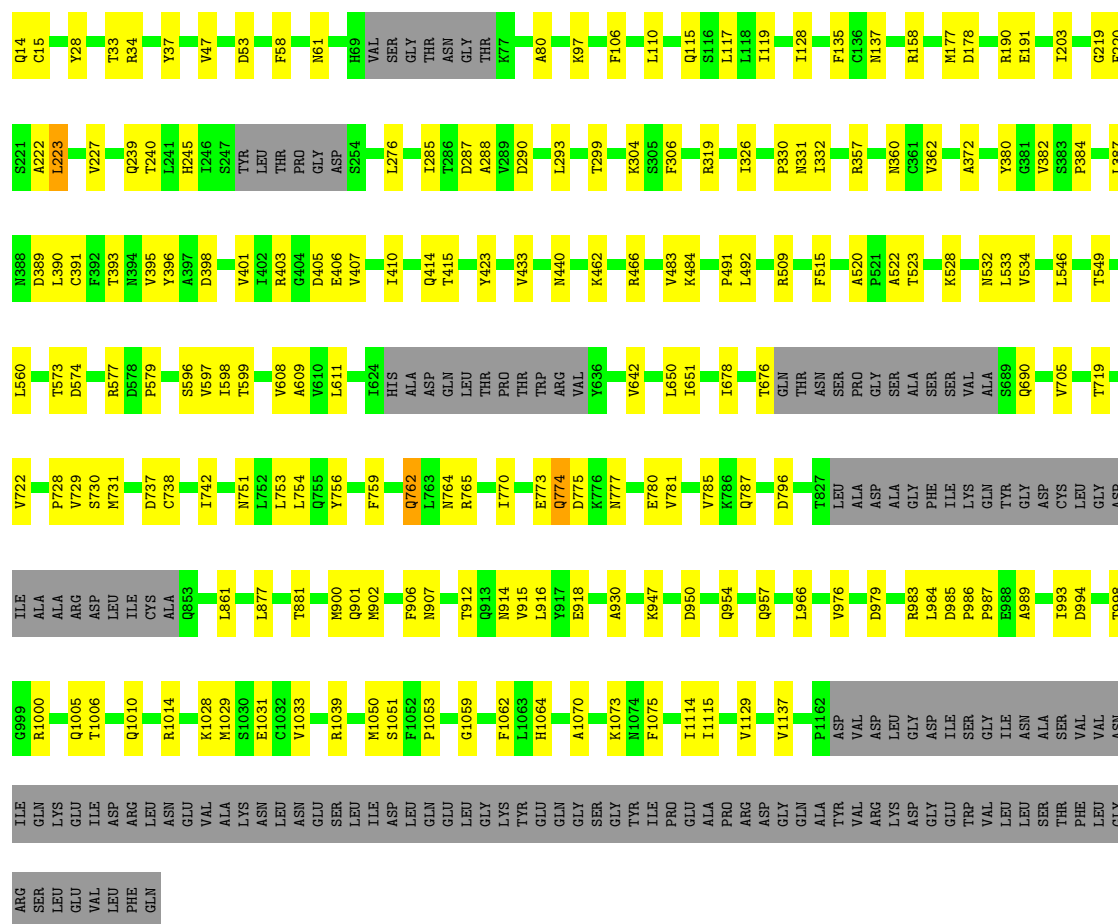
- Molecule 1: Spike glycoprotein, Fibrin





• Molecule 1: Spike glycoprotein,Fibritin

Chain 13-C: 73% 15% 12%



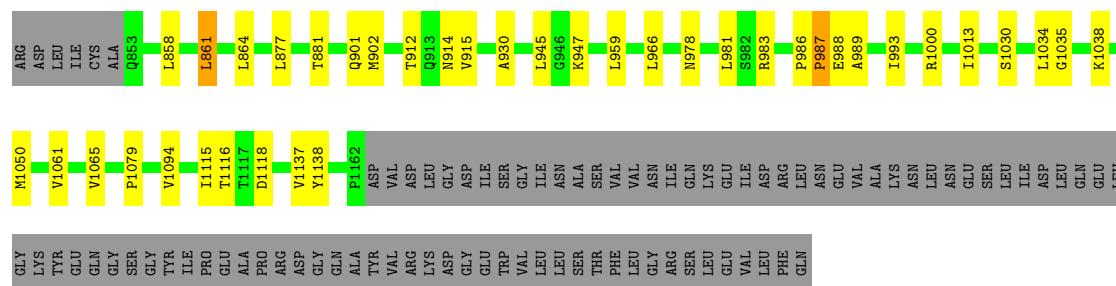
• Molecule 1: Spike glycoprotein,Fibritin

Category	Percentage
Very bad	73%
Bad	15%
Good	12%



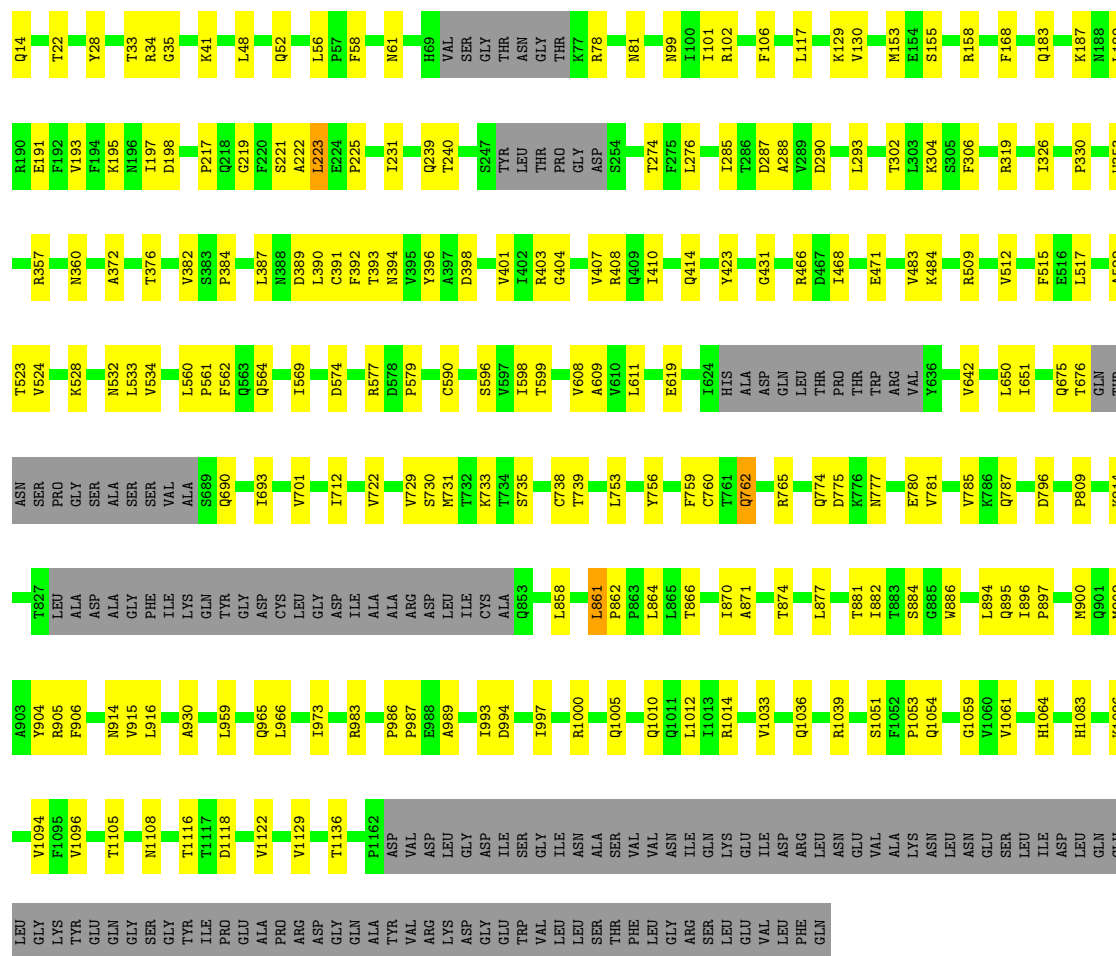
Response	Percentage
Yes	75%
No	13%
Not sure	12%





• Molecule 1: Spike glycoprotein,Fibritin

Chain 14-C: 71% 17% 12%



• Molecule 1: Spike glycoprotein,Fibritin

Chain 15-A: 73% 15% 12%



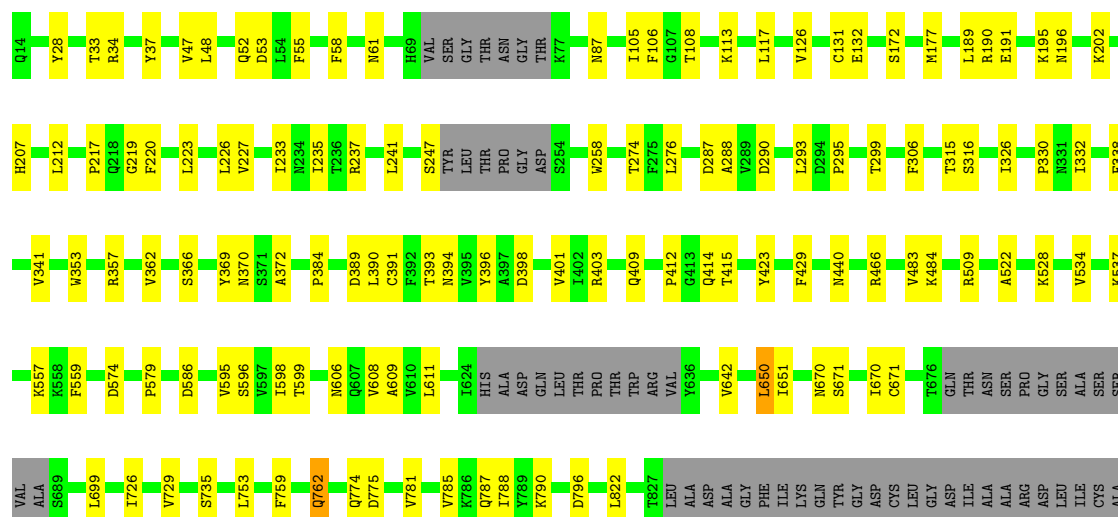
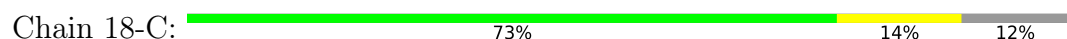
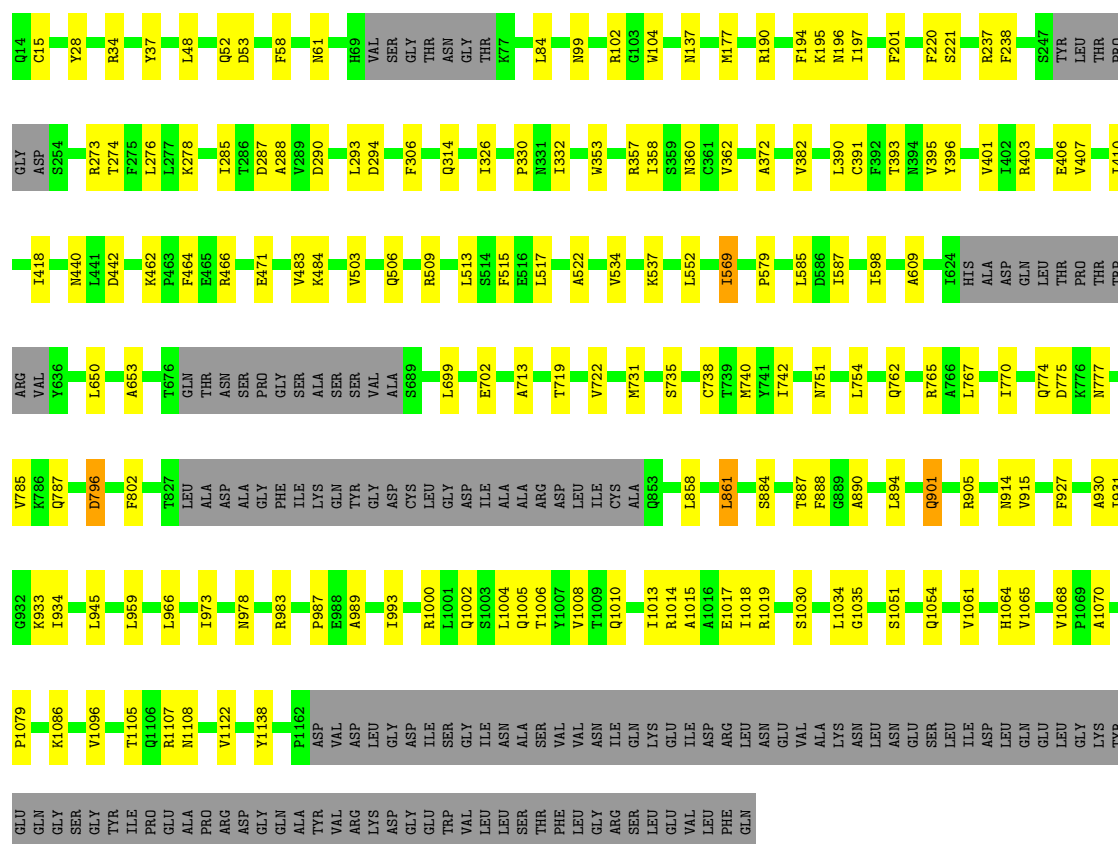









H1064	Y917	V781	V608	D389	I231	Q14
A1070	E918	V785	V610	L390	R237	T22
I1081	K947	K786	L611	F392	Y37	Y37
P1090	Q965	D796	L624	Y396	T240	Q52
F1095	L966	T805	H1S	A401	H245	D53
T1105	A972	P809	ASP	GLN	T247	F68
Q1106	V976	P809	LEU	R403	LEU	H69
Y1110	D979	K814	THR	D442	THR	VAL
E1111	L984	T827	PRO	R454	PRO	GLY
I1115	P985	LEU	THR	R457	GLY	THR
N1135	P986	ALA	ARG	F464	ASP	GLY
T1136	P987	ASP	VAL	R465	THR	THR
V1137	E988	ALA	V636	R466	S254	GLY
P1162	E993	ILE	V642	F465	T274	THR
VAL	P995	LYS	L651	V483	I285	K77
ASP	E996	GLN	L678	K484	T286	R78
LEU	L996	TYR	T676	R509	D287	F79
LEU	P997	GLY	GLN	L517	A288	A80
GLY	R1000	CYS	THR	D290	Y289	T95
ASP	L1001	LEU	ASN	T299	I101	I101
ILE	Q1002	GLY	SER	A522	R102	R102
SER	T1006	ASP	PRO	K528	F306	T108
ILE	Q1010	ALA	ALA	L533	E309	L117
ALA	R1014	ARG	ALA	V534	Q314	C131
SER	I1018	ASP	SER	K535	T315	F132
VAL	T1028	LEU	VAL	T547	S316	F133
ASN	K1028	ILE	A689	V159	Y160	Y159
GLN	E1031	ALA	O690	G550	M177	Y160
LYS	S1037	P862	V701	D574	P330	M177
ILE	K1038	P863	M703	P579	K331	R190
ASP	V1040	Y873	T719	L587	I332	F194
LEU	D1041	L878	L727	T588	N334	K195
ASN	F1042	S884	F728	P589	V353	N196
VAL	G1046	ALA	V729	F592	R357	I197
LYS	L1049	T887	L752	G593	N360	I203
LEU	H1050	A890	L763	V595	G361	K206
ASN	S1051	M902	F759	S596	V362	A222
SER	Q1054	A903	Q762	V597	Y369	L223
LEU	G1059	Y904	R765	T599	A372	E224
ILE	T1016	L916	802	M606	P384	P225
				802	V227	V227

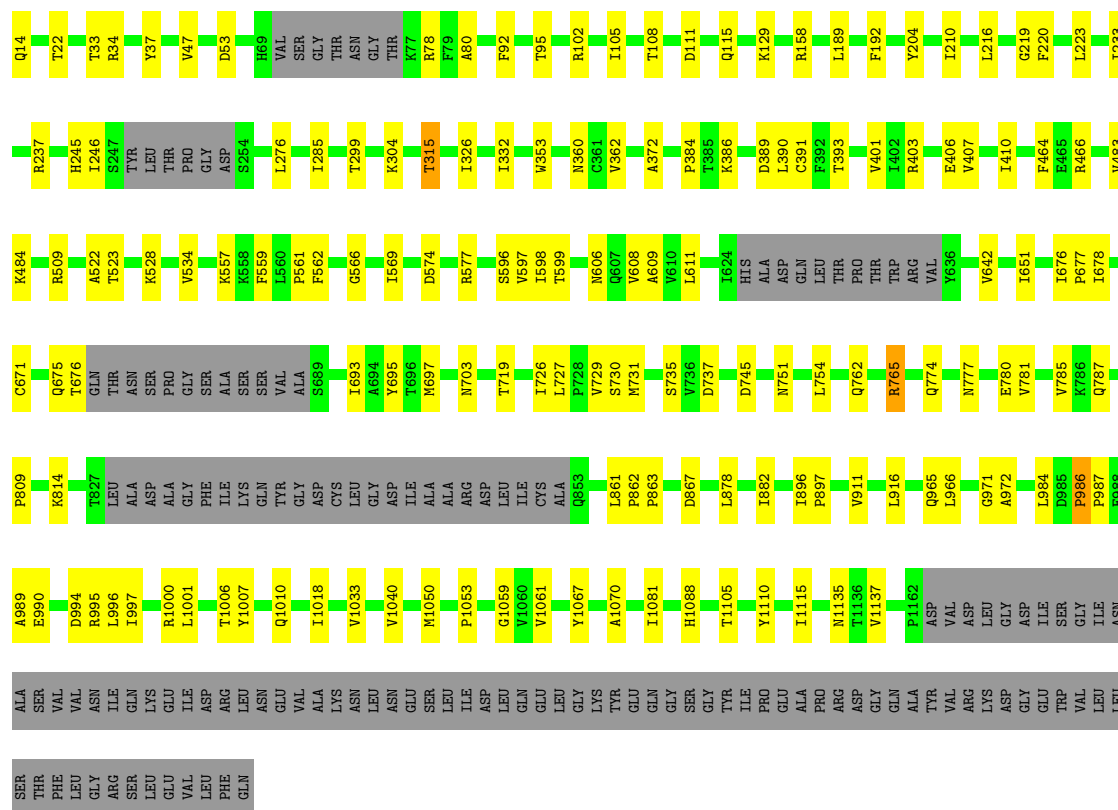





GLY
GLN
ALA
TYR
VAL
ARG
LYS
ASP
GLY
GLU
TRP
VAL
LEU
LEU
SER
THR
PHE
GLY
LEU
GLY
ARG
SER
SER
GLU
VAL
PHE
GLN

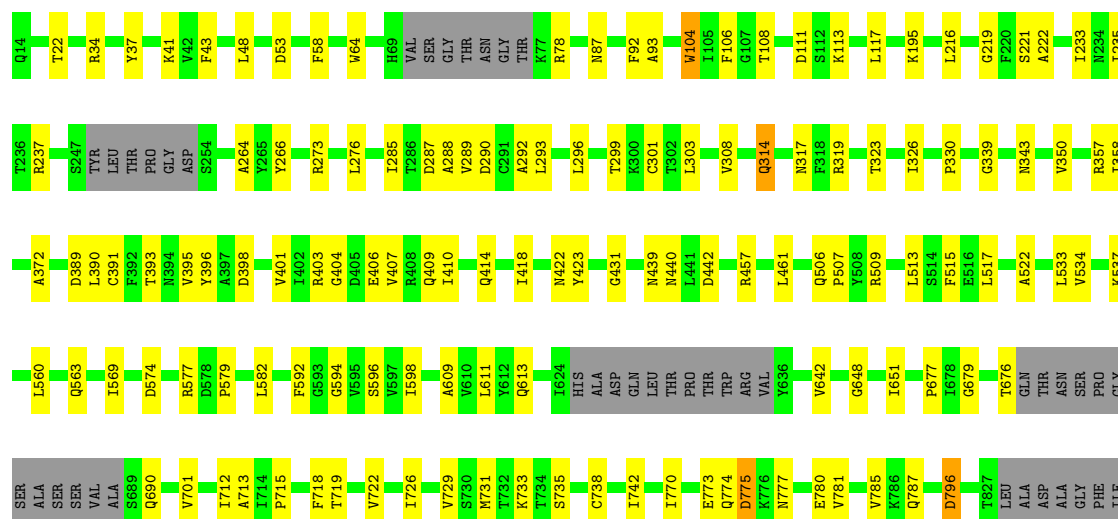
• Molecule 1: Spike glycoprotein,Fibritin

Chain 20-A:  76% 12% 12%



• Molecule 1: Spike glycoprotein,Fibritin

Chain 20-B:  73% 14% 12%



4 Experimental information

Property	Value	Source
EM reconstruction method	SINGLE PARTICLE	Depositor
Imposed symmetry	POINT, C1	Depositor
Number of particles used	309062	Depositor
Resolution determination method	FSC 0.143 CUT-OFF	Depositor
CTF correction method	NONE	Depositor
Microscope	TFS KRIOS	Depositor
Voltage (kV)	300	Depositor
Electron dose ($e^-/\text{\AA}^2$)	50	Depositor
Minimum defocus (nm)	1500	Depositor
Maximum defocus (nm)	2200	Depositor
Magnification	81000	Depositor
Image detector	GATAN K3 BIOQUANTUM (6k x 4k)	Depositor
Maximum map value	1.384	Depositor
Minimum map value	-0.335	Depositor
Average map value	0.001	Depositor
Map value standard deviation	0.057	Depositor
Recommended contour level	0.15	Depositor
Map size (\AA)	313.6, 313.6, 313.6	wwPDB
Map dimensions	224, 224, 224	wwPDB
Map angles ($^\circ$)	90.0, 90.0, 90.0	wwPDB
Pixel spacing (\AA)	1.4, 1.4, 1.4	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	1-A	0.08	0/8652	0.25	1/11768 (0.0%)
1	1-B	0.09	0/8652	0.26	0/11768
1	1-C	0.08	0/8652	0.25	1/11768 (0.0%)
1	2-A	0.08	0/8652	0.25	1/11768 (0.0%)
1	2-B	0.09	0/8652	0.26	0/11768
1	2-C	0.08	0/8652	0.25	1/11768 (0.0%)
1	3-A	0.09	0/8695	0.25	1/11829 (0.0%)
1	3-B	0.09	0/8695	0.27	0/11829
1	3-C	0.09	0/8695	0.26	1/11829 (0.0%)
1	4-A	0.09	0/8695	0.25	1/11829 (0.0%)
1	4-B	0.09	0/8695	0.27	0/11829
1	4-C	0.09	0/8695	0.26	1/11829 (0.0%)
1	5-A	0.09	0/8695	0.25	1/11829 (0.0%)
1	5-B	0.09	0/8695	0.27	0/11829
1	5-C	0.09	0/8695	0.26	1/11829 (0.0%)
1	6-A	0.09	0/8695	0.25	1/11829 (0.0%)
1	6-B	0.09	0/8695	0.27	0/11829
1	6-C	0.09	0/8695	0.26	1/11829 (0.0%)
1	7-A	0.09	0/8695	0.25	1/11829 (0.0%)
1	7-B	0.09	0/8695	0.27	0/11829
1	7-C	0.09	0/8695	0.26	1/11829 (0.0%)
1	8-A	0.09	0/8695	0.25	1/11829 (0.0%)
1	8-B	0.09	0/8695	0.27	0/11829
1	8-C	0.09	0/8695	0.26	1/11829 (0.0%)
1	9-A	0.08	0/8652	0.25	1/11768 (0.0%)
1	9-B	0.09	0/8652	0.26	0/11768
1	9-C	0.08	0/8652	0.25	1/11768 (0.0%)
1	10-A	0.09	0/8695	0.25	1/11829 (0.0%)
1	10-B	0.09	0/8695	0.27	0/11829
1	10-C	0.09	0/8695	0.26	1/11829 (0.0%)
1	11-A	0.08	0/8652	0.25	1/11768 (0.0%)
1	11-B	0.09	0/8652	0.26	0/11768
1	11-C	0.08	0/8652	0.25	1/11768 (0.0%)
1	12-A	0.09	0/8695	0.25	1/11829 (0.0%)

Mol	Chain	Bond lengths		Bond angles	
		RMSZ	# Z >5	RMSZ	# Z >5
1	12-B	0.09	0/8695	0.27	0/11829
1	12-C	0.09	0/8695	0.26	1/11829 (0.0%)
1	13-A	0.09	0/8695	0.25	1/11829 (0.0%)
1	13-B	0.09	0/8695	0.27	0/11829
1	13-C	0.09	0/8695	0.26	1/11829 (0.0%)
1	14-A	0.09	0/8695	0.25	1/11829 (0.0%)
1	14-B	0.09	0/8695	0.27	0/11829
1	14-C	0.09	0/8695	0.26	1/11829 (0.0%)
1	15-A	0.09	0/8695	0.25	1/11829 (0.0%)
1	15-B	0.09	0/8695	0.27	0/11829
1	15-C	0.09	0/8695	0.26	1/11829 (0.0%)
1	16-A	0.09	0/8695	0.25	1/11829 (0.0%)
1	16-B	0.09	0/8695	0.27	0/11829
1	16-C	0.09	0/8695	0.26	1/11829 (0.0%)
1	17-A	0.09	0/8695	0.25	1/11829 (0.0%)
1	17-B	0.09	0/8695	0.27	0/11829
1	17-C	0.09	0/8695	0.26	1/11829 (0.0%)
1	18-A	0.09	0/8695	0.25	1/11829 (0.0%)
1	18-B	0.09	0/8695	0.27	0/11829
1	18-C	0.09	0/8695	0.26	1/11829 (0.0%)
1	19-A	0.09	0/8695	0.25	1/11829 (0.0%)
1	19-B	0.09	0/8695	0.27	0/11829
1	19-C	0.09	0/8695	0.26	1/11829 (0.0%)
1	20-A	0.09	0/8695	0.25	1/11829 (0.0%)
1	20-B	0.09	0/8695	0.27	0/11829
1	20-C	0.09	0/8695	0.26	1/11829 (0.0%)
All	All	0.09	0/521184	0.26	40/709008 (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	1-B	0	1
1	1-C	0	2
1	2-C	0	1
1	8-B	0	1
1	10-B	0	1
1	11-C	0	1
1	17-C	0	1
1	20-B	0	1
All	All	0	9

There are no bond length outliers.

All (40) bond angle outliers are listed below:

Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
1	1-A	986	PRO	N-CA-C	5.63	117.57	110.70
1	2-A	986	PRO	N-CA-C	5.63	117.57	110.70
1	3-A	986	PRO	N-CA-C	5.63	117.57	110.70
1	4-A	986	PRO	N-CA-C	5.63	117.57	110.70
1	5-A	986	PRO	N-CA-C	5.63	117.57	110.70
1	6-A	986	PRO	N-CA-C	5.63	117.57	110.70
1	7-A	986	PRO	N-CA-C	5.63	117.57	110.70
1	8-A	986	PRO	N-CA-C	5.63	117.57	110.70
1	9-A	986	PRO	N-CA-C	5.63	117.57	110.70
1	10-A	986	PRO	N-CA-C	5.63	117.57	110.70
1	11-A	986	PRO	N-CA-C	5.63	117.57	110.70
1	12-A	986	PRO	N-CA-C	5.63	117.57	110.70
1	13-A	986	PRO	N-CA-C	5.63	117.57	110.70
1	14-A	986	PRO	N-CA-C	5.63	117.57	110.70
1	15-A	986	PRO	N-CA-C	5.63	117.57	110.70
1	16-A	986	PRO	N-CA-C	5.63	117.57	110.70
1	17-A	986	PRO	N-CA-C	5.63	117.57	110.70
1	18-A	986	PRO	N-CA-C	5.63	117.57	110.70
1	19-A	986	PRO	N-CA-C	5.63	117.57	110.70
1	20-A	986	PRO	N-CA-C	5.63	117.57	110.70
1	1-C	986	PRO	N-CA-C	5.04	116.85	110.70
1	2-C	986	PRO	N-CA-C	5.04	116.85	110.70
1	3-C	986	PRO	N-CA-C	5.04	116.85	110.70
1	4-C	986	PRO	N-CA-C	5.04	116.85	110.70
1	5-C	986	PRO	N-CA-C	5.04	116.85	110.70
1	6-C	986	PRO	N-CA-C	5.04	116.85	110.70
1	7-C	986	PRO	N-CA-C	5.04	116.85	110.70
1	8-C	986	PRO	N-CA-C	5.04	116.85	110.70
1	9-C	986	PRO	N-CA-C	5.04	116.85	110.70
1	10-C	986	PRO	N-CA-C	5.04	116.85	110.70
1	11-C	986	PRO	N-CA-C	5.04	116.85	110.70
1	12-C	986	PRO	N-CA-C	5.04	116.85	110.70
1	13-C	986	PRO	N-CA-C	5.04	116.85	110.70
1	14-C	986	PRO	N-CA-C	5.04	116.85	110.70
1	15-C	986	PRO	N-CA-C	5.04	116.85	110.70
1	16-C	986	PRO	N-CA-C	5.04	116.85	110.70
1	17-C	986	PRO	N-CA-C	5.04	116.85	110.70
1	18-C	986	PRO	N-CA-C	5.04	116.85	110.70
1	19-C	986	PRO	N-CA-C	5.04	116.85	110.70
1	20-C	986	PRO	N-CA-C	5.04	116.85	110.70

There are no chirality outliers.

All (9) planarity outliers are listed below:

Mol	Chain	Res	Type	Group
1	1-B	338	PHE	Sidechain
1	1-C	590	CYS	Peptide
1	1-C	592	PHE	Peptide
1	10-B	985	ASP	Sidechain
1	11-C	318	PHE	Peptide
1	17-C	1052	PHE	Sidechain
1	2-C	595	VAL	Peptide
1	20-B	1052	PHE	Sidechain
1	8-B	87	ASN	Mainchain

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	1-A	8454	0	8246	125	0
1	1-B	8454	0	8246	118	0
1	1-C	8454	0	8248	121	0
1	2-A	8454	0	8246	104	0
1	2-B	8454	0	8246	122	0
1	2-C	8454	0	8248	110	0
1	3-A	8494	0	8289	113	0
1	3-B	8494	0	8289	100	0
1	3-C	8494	0	8289	117	0
1	4-A	8494	0	8289	144	0
1	4-B	8494	0	8289	142	0
1	4-C	8494	0	8289	127	0
1	5-A	8494	0	8289	142	0
1	5-B	8494	0	8289	113	0
1	5-C	8494	0	8289	128	0
1	6-A	8494	0	8289	120	0
1	6-B	8494	0	8289	106	0
1	6-C	8494	0	8289	121	0
1	7-A	8494	0	8289	120	0
1	7-B	8494	0	8289	115	0
1	7-C	8494	0	8289	116	0

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Mol	Chain	Non-H	H(model)	H(added)	Clashes	Symm-Clashes
1	8-A	8494	0	8289	105	0
1	8-B	8494	0	8289	125	0
1	8-C	8494	0	8289	128	0
1	9-A	8454	0	8246	120	0
1	9-B	8454	0	8246	129	0
1	9-C	8454	0	8248	130	0
1	10-A	8494	0	8289	118	0
1	10-B	8494	0	8289	93	0
1	10-C	8494	0	8289	109	0
1	11-A	8454	0	8246	104	0
1	11-B	8454	0	8246	98	0
1	11-C	8454	0	8248	116	0
1	12-A	8494	0	8289	120	0
1	12-B	8494	0	8289	120	0
1	12-C	8494	0	8289	116	0
1	13-A	8494	0	8289	124	0
1	13-B	8494	0	8289	114	0
1	13-C	8494	0	8289	123	0
1	14-A	8494	0	8289	116	0
1	14-B	8494	0	8289	96	0
1	14-C	8494	0	8289	126	0
1	15-A	8494	0	8289	113	0
1	15-B	8494	0	8289	112	0
1	15-C	8494	0	8289	125	0
1	16-A	8494	0	8289	124	0
1	16-B	8494	0	8289	102	0
1	16-C	8494	0	8289	117	0
1	17-A	8494	0	8289	129	0
1	17-B	8494	0	8289	117	0
1	17-C	8494	0	8289	130	0
1	18-A	8494	0	8289	105	0
1	18-B	8494	0	8289	103	0
1	18-C	8494	0	8289	119	0
1	19-A	8494	0	8289	121	0
1	19-B	8494	0	8289	110	0
1	19-C	8494	0	8289	129	0
1	20-A	8494	0	8289	90	0
1	20-B	8494	0	8289	113	0
1	20-C	8494	0	8289	96	0
All	All	509160	0	496832	6472	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 6.

All (6472) close contacts within the same asymmetric unit are listed below, sorted by their clash magnitude.

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:C:320:VAL:C	1:C:321:GLN:N	1.70	1.50
1:A:14:GLN:HE21	1:A:14:GLN:N	1.41	1.18
1:C:853:GLN:N	1:C:853:GLN:HE21	1.62	0.97
1:C:353:TRP:HE1	1:C:466:ARG:HB2	1.41	0.84
1:A:353:TRP:HE1	1:A:466:ARG:HB2	1.44	0.81
1:B:237:ARG:HH12	1:B:239:GLN:HG3	1.46	0.81
1:A:552:LEU:HB3	1:A:585:LEU:HD12	1.63	0.81
1:A:353:TRP:HE1	1:A:466:ARG:HB2	1.46	0.80
1:B:353:TRP:HE1	1:B:466:ARG:HB2	1.44	0.80
1:C:353:TRP:HE1	1:C:466:ARG:HB2	1.47	0.80
1:C:730:SER:HB2	1:C:774:GLN:HE22	1.47	0.80
1:B:749:CYS:O	1:B:753:LEU:HB2	1.83	0.79
1:A:955:ASN:HD22	1:A:955:ASN:C	1.89	0.79
1:A:14:GLN:N	1:A:14:GLN:NE2	2.27	0.77
1:C:214:ARG:HE	1:C:215:GLY:H	1.31	0.77
1:C:778:THR:O	1:C:782:PHE:HB2	1.85	0.76
1:C:714:ILE:HD12	1:C:715:PRO:HD2	1.68	0.76
1:A:687:GLN:HG2	1:A:693:ILE:HD13	1.68	0.75
1:A:687:GLN:HG2	1:A:693:ILE:HD13	1.67	0.75
1:B:353:TRP:HE1	1:B:466:ARG:HB2	1.51	0.75
1:C:984:LEU:HD11	1:C:988:GLU:HB2	1.68	0.74
1:C:762:GLN:HA	1:C:765:ARG:HE	1.52	0.74
1:A:312:ILE:HD11	1:A:596:SER:HB3	1.70	0.74
1:A:312:ILE:HD11	1:A:596:SER:HB3	1.69	0.74
1:B:384:PRO:HA	1:B:387:LEU:HD22	1.70	0.74
1:C:853:GLN:N	1:C:853:GLN:NE2	2.34	0.74
1:A:353:TRP:HE1	1:A:466:ARG:HB2	1.53	0.74
1:B:714:ILE:HD12	1:B:715:PRO:HD2	1.70	0.74
1:A:596:SER:HB2	1:A:611:LEU:HB3	1.70	0.74
1:A:562:PHE:O	1:B:41:LYS:NZ	2.21	0.73
1:A:353:TRP:HE1	1:A:466:ARG:HB2	1.53	0.73
1:A:1128:VAL:HG21	1:B:918:GLU:HG3	1.70	0.73
1:B:384:PRO:HA	1:B:387:LEU:HG	1.71	0.73
1:C:611:LEU:HD22	1:C:678:ILE:HD11	1.69	0.73
1:B:408:ARG:HH22	1:B:409:GLN:HE21	1.36	0.73
1:A:966:LEU:HA	1:A:1000:ARG:HH12	1.54	0.72
1:C:1030:SER:HA	1:C:1034:LEU:HD23	1.71	0.72
1:B:611:LEU:HD22	1:B:678:ILE:HD11	1.70	0.72
1:A:596:SER:HB2	1:A:611:LEU:HB3	1.71	0.72
1:A:353:TRP:HE1	1:A:466:ARG:HB2	1.53	0.72

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:295:PRO:HA	1:A:298:GLU:HG2	1.71	0.72
1:B:312:ILE:HD11	1:B:596:SER:HB3	1.71	0.72
1:C:778:THR:O	1:C:782:PHE:HB2	1.90	0.72
1:C:353:TRP:HE1	1:C:355:ARG:HD3	1.55	0.72
1:A:353:TRP:HE1	1:A:466:ARG:HB2	1.54	0.72
1:A:43:PHE:O	1:C:567:ARG:NH2	2.23	0.72
1:C:984:LEU:HD11	1:C:988:GLU:HB2	1.72	0.72
1:A:353:TRP:HE1	1:A:466:ARG:HB2	1.54	0.72
1:A:353:TRP:HE1	1:A:466:ARG:HB2	1.53	0.72
1:A:177:MET:HE1	1:A:190:ARG:HH12	1.55	0.71
1:B:722:VAL:HG12	1:B:1065:VAL:HG12	1.72	0.71
1:B:966:LEU:HA	1:B:1000:ARG:HH12	1.54	0.71
1:C:983:ARG:HG3	1:C:984:LEU:HD22	1.72	0.71
1:A:611:LEU:HD22	1:A:678:ILE:HD11	1.72	0.71
1:A:687:GLN:HG2	1:A:693:ILE:HD13	1.71	0.71
1:B:517:LEU:HD23	1:B:518:LEU:HB2	1.73	0.71
1:C:611:LEU:HD22	1:C:678:ILE:HD11	1.72	0.71
1:B:439:ASN:ND2	1:B:506:GLN:OE1	2.24	0.71
1:A:735:SER:HB3	1:A:861:LEU:HD21	1.73	0.71
1:A:596:SER:HB2	1:A:611:LEU:HB3	1.73	0.71
1:A:353:TRP:HE1	1:A:466:ARG:HB2	1.56	0.71
1:B:117:LEU:HD21	1:B:231:ILE:HG21	1.73	0.71
1:A:745:ASP:OD1	1:C:319:ARG:NH1	2.24	0.71
1:C:735:SER:HB3	1:C:861:LEU:HD21	1.73	0.71
1:A:984:LEU:HD11	1:A:988:GLU:HB2	1.73	0.71
1:A:727:LEU:HD11	1:A:1025:ALA:HB2	1.72	0.71
1:A:611:LEU:HD22	1:A:678:ILE:HD11	1.73	0.71
1:A:562:PHE:O	1:B:41:LYS:NZ	2.25	0.70
1:A:904:TYR:HB3	1:C:1107:ARG:HH22	1.55	0.70
1:B:727:LEU:HD11	1:B:1025:ALA:HB2	1.74	0.70
1:A:596:SER:HB2	1:A:611:LEU:HB3	1.72	0.70
1:A:762:GLN:HA	1:A:765:ARG:HE	1.56	0.70
1:B:332:ILE:HG12	1:B:524:VAL:HG13	1.73	0.70
1:C:611:LEU:HD22	1:C:678:ILE:HD11	1.71	0.70
1:C:611:LEU:HD22	1:C:678:ILE:HD11	1.74	0.70
1:B:596:SER:HB2	1:B:611:LEU:HB3	1.73	0.70
1:A:312:ILE:HD11	1:A:596:SER:HB3	1.74	0.70
1:A:312:ILE:HD11	1:A:596:SER:HB3	1.72	0.70
1:A:596:SER:HB2	1:A:611:LEU:HB3	1.73	0.70
1:A:687:GLN:HG2	1:A:693:ILE:HD13	1.74	0.70
1:C:560:LEU:HD12	1:C:561:PRO:HD2	1.74	0.70

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:B:439:ASN:HD21	1:B:506:GLN:HB3	1.56	0.70
1:A:611:LEU:HD22	1:A:678:ILE:HD11	1.74	0.70
1:B:517:LEU:HD11	1:C:983:ARG:HH11	1.56	0.70
1:B:735:SER:HB3	1:B:861:LEU:HD21	1.73	0.70
1:B:395:VAL:HG13	1:B:513:LEU:HD11	1.73	0.69
1:C:353:TRP:HE1	1:C:466:ARG:HB2	1.57	0.69
1:C:523:THR:HG23	1:C:524:VAL:HG23	1.74	0.69
1:C:393:THR:HA	1:C:522:ALA:HA	1.73	0.69
1:A:353:TRP:HE1	1:A:466:ARG:HB2	1.55	0.69
1:A:517:LEU:HD23	1:A:518:LEU:HB2	1.73	0.69
1:A:384:PRO:HA	1:A:387:LEU:HD22	1.72	0.69
1:B:596:SER:HB2	1:B:611:LEU:HB3	1.74	0.69
1:A:517:LEU:HD23	1:A:518:LEU:HB2	1.73	0.69
1:A:596:SER:HB2	1:A:611:LEU:HB3	1.73	0.69
1:A:117:LEU:HD21	1:A:231:ILE:HG21	1.74	0.69
1:B:353:TRP:HE1	1:B:466:ARG:HB2	1.56	0.69
1:A:1123:SER:OG	1:B:914:ASN:ND2	2.26	0.69
1:A:611:LEU:HD22	1:A:678:ILE:HD11	1.73	0.69
1:B:611:LEU:HD22	1:B:678:ILE:HD11	1.73	0.69
1:B:598:ILE:HD11	1:B:678:ILE:HD13	1.75	0.69
1:A:966:LEU:HA	1:A:1000:ARG:HH12	1.57	0.69
1:A:598:ILE:HD11	1:A:664:ILE:HD12	1.75	0.69
1:B:353:TRP:HE1	1:B:466:ARG:HB2	1.57	0.69
1:C:106:PHE:HB2	1:C:117:LEU:HB3	1.75	0.69
1:A:61:ASN:HD22	1:A:61:ASN:C	2.01	0.69
1:B:1052:PHE:HB2	1:B:1063:LEU:HB2	1.75	0.69
1:C:290:ASP:HB3	1:C:293:LEU:HD23	1.74	0.69
1:B:735:SER:HB3	1:B:861:LEU:HD21	1.75	0.69
1:B:662:CYS:HB2	1:B:697:MET:HG2	1.75	0.69
1:A:61:ASN:HD22	1:A:61:ASN:C	2.01	0.69
1:A:966:LEU:HA	1:A:1000:ARG:HH12	1.57	0.69
1:A:596:SER:HB2	1:A:611:LEU:HB3	1.74	0.69
1:A:984:LEU:HD12	1:A:988:GLU:HG2	1.74	0.69
1:C:276:LEU:HD11	1:C:304:LYS:HA	1.76	0.68
1:A:755:GLN:NE2	1:C:969:ASN:OD1	2.22	0.68
1:C:401:VAL:HG22	1:C:509:ARG:HG2	1.73	0.68
1:C:312:ILE:HD12	1:C:598:ILE:HD12	1.75	0.68
1:A:966:LEU:HA	1:A:1000:ARG:HH12	1.58	0.68
1:C:596:SER:HB2	1:C:611:LEU:HB3	1.74	0.68
1:C:384:PRO:HA	1:C:387:LEU:HD23	1.75	0.68
1:A:969:ASN:HD21	1:B:755:GLN:HB2	1.59	0.68

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:C:884:SER:HB3	1:C:887:THR:HB	1.75	0.68
1:A:906:PHE:HB3	1:A:911:VAL:HB	1.76	0.68
1:C:770:ILE:O	1:C:774:GLN:HB3	1.94	0.68
1:B:237:ARG:HD2	1:B:238:PHE:H	1.59	0.68
1:A:596:SER:HB2	1:A:611:LEU:HB3	1.76	0.68
1:A:688:THR:HA	1:A:690:GLN:HG2	1.76	0.68
1:A:113:LYS:HD3	1:C:483:VAL:HA	1.76	0.68
1:C:806:LEU:HD12	1:C:807:PRO:HD2	1.73	0.68
1:C:319:ARG:NH2	1:C:590:CYS:SG	2.66	0.68
1:C:714:ILE:HD12	1:C:715:PRO:HD2	1.75	0.68
1:B:713:ALA:HB2	1:C:895:GLN:HG2	1.76	0.68
1:B:403:ARG:HG3	1:B:406:GLU:HG2	1.75	0.68
1:B:611:LEU:HD22	1:B:678:ILE:HD11	1.75	0.68
1:B:14:GLN:N	1:B:255:SER:HG	1.91	0.68
1:B:562:PHE:O	1:C:41:LYS:NZ	2.28	0.67
1:B:897:PRO:HG2	1:B:900:MET:HG2	1.75	0.67
1:A:43:PHE:H	1:C:567:ARG:HH12	1.41	0.67
1:B:353:TRP:HE1	1:B:466:ARG:HB2	1.58	0.67
1:C:100:ILE:HD12	1:C:263:ALA:HB2	1.76	0.67
1:A:360:ASN:H	1:A:523:THR:HB	1.60	0.67
1:C:983:ARG:HG3	1:C:984:LEU:HD22	1.74	0.67
1:C:1010:GLN:HE22	1:C:1014:ARG:HH22	1.42	0.67
1:C:735:SER:HB3	1:C:861:LEU:HD21	1.74	0.67
1:B:670:ASN:C	1:B:670:ASN:HD22	2.01	0.67
1:A:984:LEU:HD13	1:A:988:GLU:HG2	1.77	0.67
1:A:102:ARG:HE	1:A:246:ILE:HD11	1.59	0.67
1:C:196:ASN:HD22	1:C:235:ILE:HG12	1.59	0.67
1:B:1052:PHE:HB2	1:B:1063:LEU:HB2	1.77	0.67
1:C:320:VAL:C	1:C:321:GLN:CA	2.67	0.67
1:C:736:VAL:O	1:C:764:ASN:ND2	2.27	0.67
1:C:735:SER:HB3	1:C:861:LEU:HD21	1.76	0.67
1:A:564:GLN:H	1:B:41:LYS:HZ2	1.40	0.67
1:A:889:GLY:HA3	1:A:1034:LEU:HD21	1.76	0.67
1:B:727:LEU:HD11	1:B:1025:ALA:HB2	1.75	0.67
1:A:226:LEU:HG	1:A:227:VAL:HG23	1.77	0.67
1:A:418:ILE:HA	1:A:422:ASN:HD22	1.60	0.67
1:A:984:LEU:HB2	1:A:989:ALA:HB2	1.75	0.67
1:C:195:LYS:HE3	1:C:202:LYS:HD3	1.77	0.67
1:C:58:PHE:HD2	1:C:290:ASP:HB2	1.59	0.67
1:B:1040:VAL:HG11	1:C:1035:GLY:HA3	1.76	0.67
1:C:193:VAL:HG22	1:C:223:LEU:HD23	1.77	0.67

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:B:767:LEU:HD13	1:B:770:ILE:HD11	1.77	0.67
1:C:393:THR:HA	1:C:522:ALA:HA	1.77	0.67
1:C:596:SER:HB2	1:C:611:LEU:HB3	1.77	0.67
1:B:520:ALA:HA	1:C:41:LYS:HE2	1.76	0.67
1:B:731:MET:H	1:B:774:GLN:HE21	1.43	0.67
1:C:670:ASN:C	1:C:670:ASN:HD22	2.01	0.67
1:B:901:GLN:NE2	1:B:1050:MET:SD	2.68	0.67
1:C:907:ASN:C	1:C:907:ASN:HD22	2.02	0.67
1:B:393:THR:HA	1:B:522:ALA:HA	1.77	0.67
1:B:563:GLN:HE21	1:B:563:GLN:HA	1.60	0.67
1:A:762:GLN:HA	1:A:765:ARG:HE	1.59	0.67
1:A:206:LYS:HD3	1:A:223:LEU:HA	1.77	0.67
1:A:966:LEU:HA	1:A:1000:ARG:HH12	1.59	0.67
1:C:319:ARG:HH22	1:C:590:CYS:HB2	1.60	0.67
1:A:611:LEU:HD22	1:A:678:ILE:HD11	1.77	0.67
1:C:965:GLN:HE21	1:C:965:GLN:HA	1.59	0.66
1:A:729:VAL:HG12	1:A:1059:GLY:HA2	1.77	0.66
1:A:735:SER:HB3	1:A:861:LEU:HD21	1.76	0.66
1:A:369:TYR:HH	1:C:415:THR:HG1	1.41	0.66
1:C:126:VAL:HG22	1:C:172:SER:HB3	1.77	0.66
1:A:755:GLN:HE22	1:C:971:GLY:HA2	1.59	0.66
1:B:729:VAL:HG12	1:B:1059:GLY:HA2	1.78	0.66
1:C:729:VAL:HG12	1:C:1059:GLY:HA2	1.78	0.66
1:A:729:VAL:HG12	1:A:1059:GLY:HA2	1.77	0.66
1:A:1002:GLN:HE22	1:B:759:PHE:HE2	1.44	0.66
1:B:1079:PRO:HB3	1:C:900:MET:HE1	1.76	0.66
1:C:740:MET:HE1	1:C:857:GLY:HA3	1.78	0.66
1:C:35:GLY:HA3	1:C:56:LEU:HD13	1.78	0.66
1:C:82:PRO:O	1:C:239:GLN:NE2	2.28	0.66
1:A:596:SER:HB2	1:A:611:LEU:HB3	1.78	0.66
1:A:552:LEU:HB3	1:A:585:LEU:HD23	1.78	0.66
1:A:1123:SER:OG	1:B:914:ASN:ND2	2.29	0.66
1:C:733:LYS:HD3	1:C:771:ALA:HB1	1.78	0.66
1:C:401:VAL:HG22	1:C:509:ARG:HG2	1.78	0.66
1:A:596:SER:HB2	1:A:611:LEU:HB3	1.76	0.66
1:A:596:SER:HB2	1:A:611:LEU:HB3	1.78	0.66
1:C:106:PHE:HB2	1:C:117:LEU:HB3	1.78	0.66
1:A:57:PRO:HG3	1:A:273:ARG:HG3	1.76	0.66
1:A:688:THR:HA	1:A:690:GLN:HG2	1.78	0.66
1:B:393:THR:HA	1:B:522:ALA:HA	1.78	0.66
1:A:729:VAL:HG12	1:A:1059:GLY:HA2	1.78	0.66

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:B:1030:SER:HA	1:B:1034:LEU:HD23	1.77	0.66
1:C:714:ILE:HD12	1:C:715:PRO:HD2	1.77	0.66
1:B:722:VAL:HG22	1:B:1065:VAL:HG12	1.78	0.66
1:B:727:LEU:HD11	1:B:1025:ALA:HB2	1.77	0.65
1:B:596:SER:HB2	1:B:611:LEU:HB3	1.78	0.65
1:A:369:TYR:HH	1:C:415:THR:HG1	1.41	0.65
1:A:596:SER:HB2	1:A:611:LEU:HB3	1.77	0.65
1:B:393:THR:HA	1:B:522:ALA:HA	1.78	0.65
1:A:372:ALA:O	1:C:403:ARG:NH2	2.28	0.65
1:B:735:SER:HB3	1:B:861:LEU:HD21	1.78	0.65
1:C:729:VAL:HG12	1:C:1059:GLY:HA2	1.78	0.65
1:A:129:LYS:NZ	1:A:168:PHE:O	2.26	0.65
1:A:195:LYS:HE2	1:A:197:ILE:HD11	1.79	0.65
1:A:596:SER:HB2	1:A:611:LEU:HB3	1.77	0.65
1:B:560:LEU:HD12	1:B:561:PRO:HD2	1.76	0.65
1:B:729:VAL:HG12	1:B:1059:GLY:HA2	1.79	0.65
1:C:353:TRP:HE1	1:C:466:ARG:HB2	1.61	0.65
1:C:106:PHE:HB3	1:C:235:ILE:HG12	1.77	0.65
1:A:992:GLN:OE1	1:A:995:ARG:NH2	2.29	0.65
1:A:1002:GLN:NE2	1:B:1005:GLN:OE1	2.28	0.65
1:A:611:LEU:HD22	1:A:678:ILE:HD11	1.79	0.65
1:B:606:ASN:C	1:B:606:ASN:HD22	2.04	0.65
1:C:596:SER:HB2	1:C:611:LEU:HB3	1.77	0.65
1:B:729:VAL:HG12	1:B:1059:GLY:HA2	1.79	0.65
1:B:770:ILE:O	1:B:774:GLN:HB2	1.97	0.65
1:C:119:ILE:HG22	1:C:128:ILE:HG13	1.77	0.65
1:A:966:LEU:HA	1:A:1000:ARG:HH12	1.60	0.65
1:A:1028:LYS:NZ	1:A:1042:PHE:O	2.30	0.65
1:A:31:SER:HB3	1:A:34:ARG:HB2	1.79	0.65
1:A:596:SER:HB2	1:A:611:LEU:HB3	1.79	0.65
1:A:521:PRO:HG2	1:B:200:TYR:HE1	1.62	0.65
1:A:129:LYS:NZ	1:A:166:CYS:SG	2.70	0.65
1:C:239:GLN:NE2	1:C:240:THR:O	2.30	0.65
1:A:1046:GLY:HA2	1:B:890:ALA:HA	1.79	0.65
1:C:1028:LYS:O	1:C:1032:CYS:HB2	1.96	0.65
1:A:129:LYS:NZ	1:A:166:CYS:SG	2.69	0.65
1:A:729:VAL:HG12	1:A:1059:GLY:HA2	1.78	0.65
1:A:735:SER:HB3	1:A:861:LEU:HD21	1.79	0.65
1:B:725:GLU:OE2	1:B:1064:HIS:NE2	2.30	0.65
1:B:971:GLY:HA3	1:B:995:ARG:HH12	1.62	0.65
1:B:971:GLY:HA3	1:B:995:ARG:HH22	1.62	0.65

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:B:106:PHE:HB2	1:B:117:LEU:HB3	1.79	0.65
1:C:437:ASN:OD1	1:C:439:ASN:ND2	2.30	0.65
1:C:177:MET:SD	1:C:190:ARG:NH2	2.70	0.65
1:A:372:ALA:O	1:C:403:ARG:NH2	2.29	0.65
1:B:403:ARG:NH2	1:C:372:ALA:O	2.30	0.65
1:B:562:PHE:O	1:C:41:LYS:NZ	2.30	0.65
1:A:774:GLN:HE22	1:A:1018:ILE:HG21	1.62	0.65
1:A:715:PRO:HD3	1:B:894:LEU:HD21	1.77	0.64
1:C:384:PRO:HA	1:C:387:LEU:HD23	1.79	0.64
1:C:353:TRP:HE1	1:C:466:ARG:HB2	1.61	0.64
1:B:611:LEU:HD22	1:B:666:ILE:HD11	1.78	0.64
1:C:742:ILE:HD11	1:C:997:ILE:HA	1.79	0.64
1:C:393:THR:HA	1:C:522:ALA:HA	1.78	0.64
1:A:117:LEU:HD11	1:A:231:ILE:HG21	1.79	0.64
1:A:117:LEU:HD11	1:A:231:ILE:HG12	1.80	0.64
1:B:393:THR:HA	1:B:522:ALA:HA	1.79	0.64
1:A:729:VAL:HG12	1:A:1059:GLY:HA2	1.80	0.64
1:C:884:SER:HB3	1:C:887:THR:HB	1.79	0.64
1:A:312:ILE:HD11	1:A:596:SER:HB3	1.79	0.64
1:C:106:PHE:HB2	1:C:117:LEU:HB3	1.79	0.64
1:B:106:PHE:HB2	1:B:117:LEU:HB3	1.77	0.64
1:A:53:ASP:OD1	1:A:54:LEU:N	2.30	0.64
1:B:564:GLN:HG2	1:B:577:ARG:HD2	1.78	0.64
1:A:729:VAL:HG12	1:A:1059:GLY:HA2	1.79	0.64
1:A:762:GLN:HA	1:A:765:ARG:HE	1.62	0.64
1:A:200:TYR:O	1:A:202:LYS:NZ	2.31	0.64
1:B:353:TRP:O	1:B:466:ARG:NH2	2.31	0.64
1:A:729:VAL:HG12	1:A:1059:GLY:HA2	1.78	0.64
1:A:493:GLN:NE2	1:A:494:SER:O	2.30	0.64
1:A:984:LEU:HB3	1:A:989:ALA:HB2	1.77	0.64
1:C:729:VAL:HG12	1:C:1059:GLY:HA2	1.80	0.64
1:C:34:ARG:NH1	1:C:191:GLU:OE1	2.30	0.64
1:B:55:PHE:HB2	1:B:273:ARG:HB2	1.80	0.64
1:B:403:ARG:NH2	1:C:372:ALA:O	2.29	0.64
1:A:726:ILE:HD13	1:A:1061:VAL:HG22	1.78	0.64
1:A:384:PRO:HG3	1:C:415:THR:HG21	1.79	0.64
1:B:64:TRP:HE1	1:B:264:ALA:HB1	1.62	0.64
1:A:1081:ILE:HG23	1:A:1088:HIS:HB2	1.78	0.64
1:C:560:LEU:HD12	1:C:561:PRO:HD2	1.80	0.64
1:B:130:VAL:HB	1:B:168:PHE:HB2	1.78	0.64
1:C:598:ILE:HD11	1:C:678:ILE:HD13	1.80	0.64

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:B:596:SER:HB2	1:B:611:LEU:HB3	1.79	0.64
1:A:729:VAL:HG12	1:A:1059:GLY:HA2	1.79	0.64
1:B:1031:GLU:HB3	1:B:1037:SER:HB2	1.80	0.64
1:C:896:ILE:HD12	1:C:897:PRO:HD2	1.80	0.64
1:C:129:LYS:NZ	1:C:166:CYS:SG	2.63	0.64
1:B:360:ASN:H	1:B:523:THR:HB	1.62	0.64
1:A:611:LEU:HD22	1:A:678:ILE:HD11	1.79	0.64
1:A:104:TRP:H	1:A:119:ILE:HB	1.62	0.64
1:B:1030:SER:HA	1:B:1034:LEU:HD23	1.79	0.64
1:B:403:ARG:HB2	1:B:495:TYR:HE2	1.63	0.64
1:B:353:TRP:HE1	1:B:466:ARG:HB2	1.63	0.64
1:C:106:PHE:HB2	1:C:117:LEU:HB3	1.80	0.64
1:B:735:SER:HB3	1:B:861:LEU:HD21	1.79	0.64
1:C:384:PRO:HA	1:C:387:LEU:HD23	1.80	0.64
1:A:577:ARG:HG3	1:A:584:ILE:HG12	1.78	0.64
1:B:317:ASN:ND2	1:C:737:ASP:OD2	2.31	0.64
1:C:401:VAL:HG22	1:C:509:ARG:HG2	1.79	0.64
1:A:598:ILE:HD11	1:A:678:ILE:HD12	1.79	0.63
1:B:332:ILE:HG12	1:B:524:VAL:HG13	1.80	0.63
1:B:577:ARG:HG2	1:B:584:ILE:HG12	1.79	0.63
1:C:1106:GLN:HA	1:C:1106:GLN:HE21	1.62	0.63
1:A:966:LEU:HA	1:A:1000:ARG:HH12	1.61	0.63
1:C:726:ILE:HG22	1:C:1061:VAL:HG22	1.79	0.63
1:C:966:LEU:HA	1:C:1000:ARG:HH12	1.63	0.63
1:C:83:VAL:HG13	1:C:239:GLN:HE21	1.63	0.63
1:A:894:LEU:HD21	1:C:715:PRO:HD3	1.81	0.63
1:A:1054:GLN:HB2	1:A:1061:VAL:HG23	1.79	0.63
1:A:711:SER:OG	1:B:895:GLN:NE2	2.30	0.63
1:C:302:THR:O	1:C:304:LYS:NZ	2.31	0.63
1:A:908:GLY:O	1:A:1038:LYS:NZ	2.30	0.63
1:C:384:PRO:HA	1:C:387:LEU:HD23	1.80	0.63
1:A:966:LEU:HA	1:A:1000:ARG:HH12	1.63	0.63
1:C:384:PRO:HA	1:C:387:LEU:HD23	1.78	0.63
1:C:393:THR:HA	1:C:522:ALA:HA	1.79	0.63
1:C:726:ILE:HG22	1:C:1061:VAL:HG22	1.80	0.63
1:B:353:TRP:HE1	1:B:466:ARG:HB2	1.64	0.63
1:B:714:ILE:HD12	1:B:715:PRO:HD2	1.79	0.63
1:B:897:PRO:HG2	1:B:900:MET:HG2	1.80	0.63
1:A:353:TRP:HE1	1:A:466:ARG:HB2	1.64	0.63
1:C:596:SER:HB2	1:C:611:LEU:HB3	1.81	0.63
1:C:762:GLN:HA	1:C:765:ARG:HE	1.61	0.63

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:B:736:VAL:O	1:B:764:ASN:ND2	2.31	0.63
1:A:517:LEU:HD11	1:B:983:ARG:HD2	1.81	0.63
1:A:330:PRO:HD3	1:A:544:ASN:HD22	1.63	0.63
1:C:295:PRO:HG2	1:C:608:VAL:HG11	1.79	0.63
1:A:81:ASN:O	1:A:239:GLN:NE2	2.31	0.63
1:B:577:ARG:HH21	1:B:582:LEU:HD12	1.64	0.63
1:B:914:ASN:N	1:B:914:ASN:HD22	1.94	0.63
1:C:735:SER:HB3	1:C:861:LEU:HD21	1.79	0.63
1:C:726:ILE:HG22	1:C:1061:VAL:HG22	1.81	0.63
1:A:966:LEU:HA	1:A:1000:ARG:HH12	1.62	0.63
1:A:598:ILE:HD11	1:A:678:ILE:HD12	1.79	0.63
1:C:941:THR:HG23	1:C:944:ALA:HB2	1.79	0.63
1:B:966:LEU:HA	1:B:1000:ARG:HH12	1.62	0.63
1:A:394:ASN:ND2	1:B:200:TYR:OH	2.31	0.63
1:C:393:THR:HA	1:C:522:ALA:HA	1.81	0.63
1:B:722:VAL:HG12	1:B:1065:VAL:HG22	1.79	0.63
1:A:994:ASP:HA	1:A:997:ILE:HG12	1.81	0.63
1:A:729:VAL:HG12	1:A:1059:GLY:HA2	1.80	0.63
1:A:31:SER:HB2	1:A:34:ARG:HB2	1.80	0.63
1:C:598:ILE:HB	1:C:609:ALA:HB3	1.80	0.63
1:C:1028:LYS:NZ	1:C:1042:PHE:O	2.23	0.63
1:A:278:LYS:NZ	1:A:286:THR:OG1	2.30	0.63
1:B:353:TRP:HE1	1:B:466:ARG:HB2	1.64	0.63
1:C:1043:CYS:O	1:C:1064:HIS:ND1	2.31	0.63
1:B:966:LEU:HA	1:B:1000:ARG:HH12	1.63	0.63
1:A:729:VAL:HG12	1:A:1059:GLY:HA2	1.80	0.63
1:B:360:ASN:H	1:B:523:THR:HB	1.64	0.63
1:A:752:LEU:HD22	1:A:993:ILE:HD12	1.80	0.63
1:A:394:ASN:HB2	1:A:516:GLU:HB3	1.81	0.63
1:B:64:TRP:HE1	1:B:264:ALA:HB1	1.63	0.63
1:B:1079:PRO:HB3	1:C:900:MET:HE1	1.81	0.63
1:A:393:THR:HA	1:A:522:ALA:HA	1.80	0.63
1:A:719:THR:HG22	1:A:1070:ALA:HB2	1.81	0.63
1:A:896:ILE:HD12	1:A:897:PRO:HD2	1.80	0.62
1:B:707:TYR:HD2	1:C:792:PRO:HG3	1.64	0.62
1:C:994:ASP:HA	1:C:997:ILE:HG12	1.81	0.62
1:A:983:ARG:HH21	1:C:517:LEU:HD21	1.63	0.62
1:C:1030:SER:HA	1:C:1034:LEU:HD23	1.81	0.62
1:B:106:PHE:HB2	1:B:117:LEU:HB3	1.81	0.62
1:A:735:SER:HB3	1:A:861:LEU:HD21	1.80	0.62
1:B:517:LEU:HD11	1:C:983:ARG:HH11	1.63	0.62

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:B:102:ARG:HH21	1:B:246:ILE:HG12	1.64	0.62
1:B:984:LEU:HB2	1:B:989:ALA:HB2	1.80	0.62
1:A:1010:GLN:OE1	1:A:1014:ARG:NH1	2.30	0.62
1:C:606:ASN:HD22	1:C:606:ASN:C	2.06	0.62
1:C:966:LEU:HA	1:C:1000:ARG:HH12	1.62	0.62
1:A:234:ASN:ND2	1:C:465:GLU:OE1	2.32	0.62
1:B:729:VAL:HG12	1:B:1059:GLY:HA2	1.81	0.62
1:A:372:ALA:O	1:C:403:ARG:NH2	2.30	0.62
1:B:1030:SER:HA	1:B:1034:LEU:HD23	1.81	0.62
1:A:1081:ILE:HG23	1:A:1088:HIS:HB2	1.81	0.62
1:C:866:THR:H	1:C:869:MET:HE2	1.64	0.62
1:C:908:GLY:O	1:C:1038:LYS:NZ	2.31	0.62
1:A:702:GLU:OE2	1:B:790:LYS:NZ	2.32	0.62
1:C:53:ASP:OD1	1:C:54:LEU:N	2.32	0.62
1:B:517:LEU:HD11	1:C:983:ARG:HH11	1.65	0.62
1:C:1031:GLU:HG3	1:C:1039:ARG:HH21	1.64	0.62
1:C:887:THR:HG21	1:C:894:LEU:HG	1.80	0.62
1:C:598:ILE:HB	1:C:609:ALA:HB3	1.82	0.62
1:A:1051:SER:OG	1:A:1064:HIS:ND1	2.31	0.62
1:A:877:LEU:HD21	1:A:1029:MET:HE1	1.81	0.62
1:A:226:LEU:HG	1:A:227:VAL:HG23	1.81	0.62
1:C:53:ASP:OD2	1:C:54:LEU:N	2.31	0.62
1:A:984:LEU:HD13	1:A:988:GLU:HG2	1.81	0.62
1:A:403:ARG:NH2	1:B:372:ALA:O	2.31	0.62
1:B:598:ILE:HB	1:B:609:ALA:HB3	1.81	0.62
1:C:229:LEU:HD22	1:C:231:ILE:HD13	1.81	0.62
1:B:729:VAL:HG12	1:B:1059:GLY:HA2	1.80	0.62
1:C:99:ASN:O	1:C:102:ARG:NH2	2.32	0.62
1:A:763:LEU:HD23	1:A:1004:LEU:HD13	1.81	0.62
1:A:1102:TRP:HD1	1:A:1135:ASN:HD22	1.47	0.62
1:A:314:GLN:HE22	1:A:594:GLY:HA3	1.65	0.62
1:C:1031:GLU:HG3	1:C:1039:ARG:HH21	1.64	0.62
1:A:372:ALA:O	1:C:403:ARG:NH2	2.28	0.62
1:C:897:PRO:HG2	1:C:900:MET:HG2	1.80	0.62
1:C:106:PHE:HB2	1:C:117:LEU:HB3	1.81	0.62
1:A:825:LYS:HD3	1:A:945:LEU:HD13	1.81	0.62
1:B:353:TRP:HE1	1:B:466:ARG:HB2	1.63	0.62
1:A:521:PRO:HD3	1:B:41:LYS:HE2	1.82	0.62
1:A:369:TYR:HH	1:C:415:THR:HG1	1.46	0.62
1:A:571:ASP:OD1	1:B:975:SER:OG	2.18	0.62
1:A:99:ASN:HB3	1:A:102:ARG:HH12	1.64	0.62

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:598:ILE:HB	1:A:609:ALA:HB3	1.82	0.62
1:C:393:THR:HA	1:C:522:ALA:HA	1.81	0.62
1:C:596:SER:HB2	1:C:611:LEU:HB3	1.81	0.62
1:A:729:VAL:HG12	1:A:1059:GLY:HA2	1.82	0.62
1:C:726:ILE:HG22	1:C:1061:VAL:HG22	1.82	0.62
1:C:362:VAL:HG21	1:C:524:VAL:HB	1.81	0.62
1:C:338:PHE:HA	1:C:341:VAL:HG12	1.81	0.62
1:A:393:THR:HA	1:A:522:ALA:HA	1.81	0.62
1:B:596:SER:HB2	1:B:611:LEU:HB3	1.81	0.62
1:A:393:THR:HA	1:A:522:ALA:HA	1.81	0.62
1:B:969:ASN:OD1	1:C:755:GLN:NE2	2.26	0.62
1:A:1031:GLU:HB3	1:A:1037:SER:HB2	1.81	0.62
1:B:353:TRP:HE1	1:B:466:ARG:HB2	1.64	0.62
1:A:984:LEU:HB2	1:A:989:ALA:HB2	1.82	0.62
1:A:317:ASN:ND2	1:B:737:ASP:OD2	2.32	0.62
1:A:372:ALA:O	1:C:403:ARG:NH2	2.33	0.62
1:C:555:SER:HB2	1:C:586:ASP:HB2	1.82	0.62
1:C:130:VAL:HG21	1:C:231:ILE:HD11	1.81	0.62
1:B:735:SER:HB3	1:B:861:LEU:HD21	1.80	0.62
1:C:287:ASP:HB2	1:C:306:PHE:HE2	1.65	0.62
1:C:393:THR:HA	1:C:522:ALA:HA	1.82	0.62
1:B:360:ASN:HD22	1:B:360:ASN:C	2.07	0.61
1:A:994:ASP:HA	1:A:997:ILE:HG12	1.81	0.61
1:C:729:VAL:HG21	1:C:781:VAL:HG11	1.81	0.61
1:C:729:VAL:HG12	1:C:1059:GLY:HA2	1.82	0.61
1:A:117:LEU:HD11	1:A:231:ILE:HG12	1.81	0.61
1:C:606:ASN:HD22	1:C:606:ASN:C	2.08	0.61
1:B:393:THR:HA	1:B:522:ALA:HA	1.80	0.61
1:C:1106:GLN:NE2	1:C:1111:GLU:OE1	2.33	0.61
1:C:560:LEU:HD12	1:C:561:PRO:HD2	1.82	0.61
1:A:968:SER:OG	1:B:755:GLN:O	2.18	0.61
1:C:1010:GLN:OE1	1:C:1014:ARG:NH2	2.33	0.61
1:C:1019:ARG:NH2	1:C:1023:ASN:OD1	2.33	0.61
1:A:1081:ILE:HG23	1:A:1088:HIS:HB2	1.81	0.61
1:C:357:ARG:NH2	1:C:396:TYR:OH	2.33	0.61
1:A:774:GLN:O	1:A:777:ASN:ND2	2.34	0.61
1:B:735:SER:HB3	1:B:861:LEU:HD21	1.82	0.61
1:A:598:ILE:HD11	1:A:678:ILE:HD12	1.82	0.61
1:B:1079:PRO:HB3	1:C:900:MET:HE1	1.81	0.61
1:B:393:THR:HA	1:B:522:ALA:HA	1.81	0.61
1:B:393:THR:HA	1:B:522:ALA:HA	1.83	0.61

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:B:401:VAL:HG22	1:B:509:ARG:HG2	1.82	0.61
1:A:729:VAL:HG12	1:A:1059:GLY:HA2	1.81	0.61
1:A:729:VAL:HG12	1:A:1059:GLY:HA2	1.81	0.61
1:A:729:VAL:HG21	1:A:781:VAL:HG11	1.82	0.61
1:A:273:ARG:HD2	1:A:292:ALA:HB3	1.82	0.61
1:A:369:TYR:HH	1:C:415:THR:HG1	1.48	0.61
1:A:576:VAL:HG12	1:A:587:ILE:HD11	1.82	0.61
1:C:393:THR:HA	1:C:522:ALA:HA	1.82	0.61
1:B:317:ASN:ND2	1:C:737:ASP:OD2	2.34	0.61
1:C:606:ASN:HD22	1:C:606:ASN:C	2.08	0.61
1:A:730:SER:HA	1:A:774:GLN:HE22	1.64	0.61
1:C:802:PHE:HB3	1:C:806:LEU:HD23	1.81	0.61
1:A:547:THR:O	1:B:978:ASN:ND2	2.34	0.61
1:A:735:SER:HB3	1:A:861:LEU:HD21	1.81	0.61
1:B:319:ARG:HH22	1:C:745:ASP:HA	1.64	0.61
1:A:117:LEU:HD11	1:A:231:ILE:HG21	1.82	0.61
1:A:1051:SER:OG	1:A:1064:HIS:ND1	2.33	0.61
1:C:106:PHE:HD1	1:C:235:ILE:HD13	1.65	0.61
1:A:897:PRO:HG2	1:A:900:MET:HG2	1.82	0.61
1:A:992:GLN:OE1	1:A:995:ARG:NH2	2.33	0.61
1:C:393:THR:HA	1:C:522:ALA:HA	1.82	0.61
1:C:105:ILE:HG12	1:C:241:LEU:HD21	1.83	0.61
1:B:729:VAL:HG12	1:B:1059:GLY:HA2	1.83	0.61
1:C:401:VAL:HG22	1:C:509:ARG:HG2	1.81	0.61
1:B:403:ARG:NH2	1:C:372:ALA:O	2.31	0.61
1:B:729:VAL:HG12	1:B:1059:GLY:HA2	1.82	0.61
1:A:687:GLN:HG2	1:A:693:ILE:HD13	1.82	0.61
1:A:887:THR:HB	1:A:894:LEU:HD23	1.83	0.61
1:A:394:ASN:HB2	1:A:516:GLU:HB3	1.81	0.61
1:C:858:LEU:HD22	1:C:959:LEU:HD12	1.82	0.61
1:A:1081:ILE:HG23	1:A:1088:HIS:HB2	1.83	0.61
1:B:1051:SER:OG	1:B:1064:HIS:ND1	2.31	0.61
1:C:735:SER:HB3	1:C:861:LEU:HD21	1.82	0.61
1:C:379:CYS:HA	1:C:432:CYS:HB3	1.83	0.61
1:A:914:ASN:OD1	1:C:1123:SER:OG	2.18	0.61
1:C:99:ASN:O	1:C:102:ARG:NH1	2.33	0.61
1:A:1017:GLU:OE1	1:B:1019:ARG:NH1	2.31	0.61
1:A:598:ILE:HB	1:A:609:ALA:HB3	1.83	0.61
1:C:28:TYR:HB3	1:C:61:ASN:HB3	1.83	0.61
1:C:729:VAL:HG12	1:C:1059:GLY:HA2	1.83	0.61
1:A:278:LYS:NZ	1:A:286:THR:OG1	2.33	0.61

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:389:ASP:OD1	1:A:528:LYS:NZ	2.34	0.61
1:C:596:SER:HB2	1:C:611:LEU:HB3	1.83	0.61
1:C:1031:GLU:OE2	1:C:1039:ARG:NH2	2.34	0.61
1:A:393:THR:HA	1:A:522:ALA:HA	1.83	0.61
1:A:606:ASN:C	1:A:606:ASN:HD22	2.08	0.61
1:C:444:LYS:HD3	1:C:446:GLY:H	1.66	0.61
1:A:915:VAL:HG12	1:A:1109:PHE:HD2	1.65	0.61
1:A:517:LEU:HD23	1:A:518:LEU:HB2	1.83	0.61
1:B:1052:PHE:HB2	1:B:1063:LEU:HB2	1.82	0.61
1:A:596:SER:HB2	1:A:611:LEU:HB3	1.83	0.61
1:A:189:LEU:HD22	1:A:210:ILE:HD13	1.83	0.61
1:C:106:PHE:HB3	1:C:235:ILE:HD12	1.82	0.61
1:B:317:ASN:HD21	1:B:592:PHE:HD1	1.49	0.61
1:B:564:GLN:H	1:C:41:LYS:HZ3	1.49	0.61
1:B:577:ARG:HH21	1:B:582:LEU:HD12	1.66	0.61
1:B:731:MET:H	1:B:774:GLN:HE21	1.48	0.61
1:A:598:ILE:HB	1:A:609:ALA:HB3	1.83	0.61
1:B:729:VAL:HG12	1:B:1059:GLY:HA2	1.83	0.61
1:A:908:GLY:O	1:A:1038:LYS:NZ	2.30	0.61
1:C:858:LEU:HD22	1:C:959:LEU:HD12	1.83	0.60
1:C:598:ILE:HB	1:C:609:ALA:HB3	1.82	0.60
1:A:923:ILE:HA	1:A:926:GLN:HG3	1.82	0.60
1:B:326:ILE:HD11	1:B:534:VAL:HG12	1.81	0.60
1:B:106:PHE:HB2	1:B:117:LEU:HB3	1.83	0.60
1:B:707:TYR:HD2	1:C:792:PRO:HG3	1.66	0.60
1:A:606:ASN:C	1:A:606:ASN:HD22	2.08	0.60
1:A:93:ALA:HB3	1:A:266:TYR:HB2	1.82	0.60
1:C:452:LEU:HD22	1:C:492:LEU:HD11	1.83	0.60
1:C:908:GLY:O	1:C:1038:LYS:NZ	2.34	0.60
1:C:1030:SER:HA	1:C:1034:LEU:HD23	1.84	0.60
1:B:896:ILE:HD12	1:B:897:PRO:HD2	1.82	0.60
1:B:384:PRO:HA	1:B:387:LEU:HD23	1.84	0.60
1:A:664:ILE:HD12	1:A:665:PRO:HD2	1.81	0.60
1:A:418:ILE:HA	1:A:422:ASN:HD22	1.66	0.60
1:B:966:LEU:HA	1:B:1000:ARG:HH12	1.66	0.60
1:B:729:VAL:HG12	1:B:1059:GLY:HA2	1.82	0.60
1:A:562:PHE:O	1:B:41:LYS:NZ	2.35	0.60
1:A:983:ARG:NH2	1:C:428:ASP:OD1	2.35	0.60
1:C:1031:GLU:HB3	1:C:1037:SER:HB2	1.82	0.60
1:C:357:ARG:NH2	1:C:396:TYR:OH	2.34	0.60
1:C:720:ILE:HD11	1:C:1065:VAL:HG13	1.84	0.60

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:C:598:ILE:HB	1:C:609:ALA:HB3	1.84	0.60
1:A:596:SER:HB2	1:A:611:LEU:HB3	1.83	0.60
1:A:729:VAL:HG12	1:A:1059:GLY:HA2	1.82	0.60
1:B:398:ASP:OD2	1:B:423:TYR:OH	2.19	0.60
1:C:67:ALA:HB3	1:C:263:ALA:H	1.65	0.60
1:B:738:CYS:O	1:B:742:ILE:HB	2.00	0.60
1:A:973:ILE:HG21	1:A:983:ARG:HH22	1.66	0.60
1:C:389:ASP:OD1	1:C:528:LYS:NZ	2.35	0.60
1:A:372:ALA:O	1:C:403:ARG:NH2	2.34	0.60
1:C:109:THR:HG21	1:C:113:LYS:HD3	1.82	0.60
1:A:598:ILE:HB	1:A:609:ALA:HB3	1.84	0.60
1:C:735:SER:HB3	1:C:861:LEU:HD21	1.82	0.60
1:B:598:ILE:HB	1:B:609:ALA:HB3	1.83	0.60
1:A:598:ILE:HB	1:A:609:ALA:HB3	1.84	0.60
1:C:762:GLN:HA	1:C:765:ARG:HE	1.66	0.60
1:A:403:ARG:NH1	1:A:405:ASP:OD2	2.34	0.60
1:A:729:VAL:HG12	1:A:1059:GLY:HA2	1.82	0.60
1:A:729:VAL:HG21	1:A:781:VAL:HG11	1.84	0.60
1:A:981:LEU:O	1:C:386:LYS:NZ	2.34	0.60
1:B:966:LEU:HA	1:B:1000:ARG:HH12	1.65	0.60
1:B:401:VAL:HG22	1:B:509:ARG:HG2	1.83	0.60
1:B:738:CYS:HB3	1:B:763:LEU:HD11	1.82	0.60
1:C:806:LEU:HD12	1:C:807:PRO:HD2	1.81	0.60
1:B:733:LYS:HD3	1:B:771:ALA:HB1	1.82	0.60
1:A:357:ARG:NH2	1:A:394:ASN:OD1	2.34	0.60
1:B:560:LEU:HD23	1:B:563:GLN:HB2	1.83	0.60
1:A:372:ALA:O	1:C:403:ARG:NH2	2.31	0.60
1:A:557:LYS:NZ	1:B:281:GLU:O	2.35	0.60
1:B:598:ILE:HD11	1:B:678:ILE:HD13	1.83	0.60
1:B:726:ILE:HG22	1:B:1061:VAL:HG22	1.84	0.60
1:C:902:MET:HB3	1:C:916:LEU:HD11	1.84	0.60
1:C:299:THR:HG22	1:C:597:VAL:HG11	1.83	0.60
1:A:598:ILE:HB	1:A:609:ALA:HB3	1.82	0.60
1:A:984:LEU:HD13	1:A:988:GLU:HG2	1.83	0.60
1:C:729:VAL:HG12	1:C:1059:GLY:HA2	1.83	0.60
1:C:1033:VAL:HG21	1:C:1053:PRO:HG3	1.82	0.60
1:A:226:LEU:HG	1:A:227:VAL:HG23	1.82	0.60
1:C:452:LEU:HB3	1:C:492:LEU:HD12	1.82	0.60
1:C:378:LYS:NZ	1:C:407:VAL:O	2.35	0.60
1:A:729:VAL:HG12	1:A:1059:GLY:HA2	1.83	0.60
1:A:195:LYS:HD2	1:A:197:ILE:HD11	1.84	0.60

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:34:ARG:NH1	1:A:219:GLY:O	2.33	0.60
1:C:598:ILE:HB	1:C:609:ALA:HB3	1.83	0.60
1:A:403:ARG:NH2	1:B:372:ALA:O	2.28	0.60
1:C:598:ILE:HB	1:C:609:ALA:HB3	1.84	0.60
1:A:726:ILE:HG22	1:A:1061:VAL:HG22	1.83	0.60
1:C:53:ASP:OD1	1:C:54:LEU:N	2.34	0.60
1:A:119:ILE:HD12	1:A:128:ILE:HG12	1.84	0.60
1:A:889:GLY:HA3	1:A:1034:LEU:HD21	1.84	0.60
1:B:332:ILE:HG23	1:B:529:LYS:HE3	1.84	0.60
1:B:598:ILE:HB	1:B:609:ALA:HB3	1.84	0.60
1:B:729:VAL:HG21	1:B:781:VAL:HG11	1.83	0.60
1:C:407:VAL:HG21	1:C:508:TYR:HD2	1.65	0.60
1:B:58:PHE:HB3	1:B:293:LEU:HD12	1.84	0.60
1:B:1079:PRO:HB3	1:C:900:MET:HE1	1.84	0.60
1:A:1081:ILE:HG23	1:A:1088:HIS:HB2	1.84	0.60
1:C:598:ILE:HB	1:C:609:ALA:HB3	1.84	0.60
1:A:230:PRO:O	1:C:466:ARG:NH2	2.34	0.60
1:A:396:TYR:HB2	1:A:514:SER:HB3	1.84	0.60
1:B:498:GLN:HE21	1:B:501:TYR:HB2	1.66	0.60
1:A:598:ILE:HB	1:A:609:ALA:HB3	1.82	0.60
1:C:598:ILE:HD11	1:C:678:ILE:HD13	1.84	0.60
1:C:825:LYS:NZ	1:C:938:LEU:O	2.35	0.60
1:B:147:LYS:HB3	1:B:153:MET:HE1	1.83	0.60
1:A:117:LEU:HD11	1:A:231:ILE:HG12	1.84	0.60
1:B:1079:PRO:HB3	1:C:900:MET:HE1	1.82	0.60
1:C:756:TYR:HB3	1:C:759:PHE:HD2	1.66	0.60
1:A:315:THR:HG23	1:A:595:VAL:HG23	1.84	0.60
1:A:276:LEU:HD11	1:A:304:LYS:HE2	1.84	0.60
1:A:418:ILE:HA	1:A:422:ASN:HD22	1.66	0.59
1:B:393:THR:HA	1:B:522:ALA:HA	1.82	0.59
1:A:193:VAL:HG12	1:A:204:TYR:HB2	1.84	0.59
1:A:598:ILE:HD11	1:A:678:ILE:HD12	1.84	0.59
1:B:393:THR:HA	1:B:522:ALA:HA	1.84	0.59
1:A:34:ARG:HH22	1:A:220:PHE:HA	1.65	0.59
1:B:1030:SER:HA	1:B:1034:LEU:HD23	1.84	0.59
1:C:108:THR:O	1:C:237:ARG:NH2	2.35	0.59
1:C:731:MET:H	1:C:774:GLN:HE21	1.49	0.59
1:B:328:ARG:HH21	1:B:533:LEU:H	1.49	0.59
1:B:327:VAL:O	1:B:531:THR:OG1	2.21	0.59
1:C:777:ASN:HA	1:C:780:GLU:HG3	1.83	0.59
1:B:418:ILE:HD12	1:B:422:ASN:HD22	1.67	0.59

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:276:LEU:HG	1:A:306:PHE:HE1	1.66	0.59
1:C:598:ILE:HB	1:C:609:ALA:HB3	1.84	0.59
1:C:58:PHE:HD1	1:C:290:ASP:HB2	1.67	0.59
1:A:702:GLU:OE2	1:B:790:LYS:NZ	2.32	0.59
1:C:877:LEU:HD21	1:C:1029:MET:HE1	1.84	0.59
1:B:118:LEU:HD21	1:B:160:TYR:HE2	1.67	0.59
1:C:53:ASP:OD2	1:C:54:LEU:N	2.34	0.59
1:B:40:ASP:OD1	1:B:41:LYS:N	2.36	0.59
1:A:729:VAL:HG21	1:A:781:VAL:HG11	1.84	0.59
1:B:1079:PRO:HB3	1:C:900:MET:HE1	1.83	0.59
1:A:905:ARG:HH21	1:A:1050:MET:HB2	1.66	0.59
1:A:966:LEU:HA	1:A:1000:ARG:HH12	1.68	0.59
1:A:61:ASN:HD22	1:A:61:ASN:C	2.03	0.59
1:A:729:VAL:HG21	1:A:781:VAL:HG11	1.83	0.59
1:B:970:PHE:HA	1:C:756:TYR:HE2	1.67	0.59
1:A:108:THR:O	1:A:237:ARG:NH2	2.36	0.59
1:C:712:ILE:HG13	1:C:1094:VAL:HG11	1.84	0.59
1:B:401:VAL:HG22	1:B:509:ARG:HG2	1.83	0.59
1:A:788:ILE:HD12	1:A:876:ALA:HB2	1.83	0.59
1:C:53:ASP:OD1	1:C:54:LEU:N	2.35	0.59
1:A:904:TYR:OH	1:C:1093:GLY:O	2.20	0.59
1:A:729:VAL:HG12	1:A:1059:GLY:HA2	1.85	0.59
1:B:28:TYR:HB3	1:B:61:ASN:HB3	1.84	0.59
1:A:1116:THR:OG1	1:A:1118:ASP:OD1	2.20	0.59
1:C:229:LEU:HD22	1:C:231:ILE:HG22	1.84	0.59
1:C:393:THR:HA	1:C:522:ALA:HA	1.84	0.59
1:C:55:PHE:HB2	1:C:273:ARG:HB2	1.83	0.59
1:C:1033:VAL:HG21	1:C:1053:PRO:HG3	1.84	0.59
1:A:598:ILE:HB	1:A:609:ALA:HB3	1.84	0.59
1:A:598:ILE:HB	1:A:609:ALA:HB3	1.84	0.59
1:B:129:LYS:NZ	1:B:168:PHE:O	2.34	0.59
1:B:966:LEU:HA	1:B:1000:ARG:HH12	1.67	0.59
1:A:1129:VAL:HG22	1:B:917:TYR:HB3	1.83	0.59
1:B:560:LEU:HD12	1:B:562:PHE:H	1.68	0.59
1:C:384:PRO:HA	1:C:387:LEU:HD23	1.83	0.59
1:C:408:ARG:NH1	1:C:414:GLN:OE1	2.35	0.59
1:B:908:GLY:O	1:B:1038:LYS:NZ	2.34	0.59
1:B:915:VAL:HG11	1:B:1108:ASN:HB2	1.85	0.59
1:A:598:ILE:HB	1:A:609:ALA:HB3	1.85	0.59
1:A:598:ILE:HB	1:A:609:ALA:HB3	1.85	0.59
1:B:108:THR:O	1:B:237:ARG:NH2	2.35	0.59

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:B:563:GLN:HA	1:C:41:LYS:HZ1	1.68	0.59
1:A:201:PHE:HB2	1:A:229:LEU:HB2	1.83	0.59
1:C:598:ILE:HB	1:C:609:ALA:HB3	1.84	0.59
1:B:129:LYS:HZ3	1:B:168:PHE:H	1.49	0.59
1:C:729:VAL:HG12	1:C:1059:GLY:HA2	1.84	0.59
1:A:729:VAL:HG21	1:A:781:VAL:HG11	1.84	0.59
1:B:393:THR:HA	1:B:522:ALA:HA	1.84	0.59
1:A:102:ARG:NH1	1:A:121:ASN:O	2.36	0.59
1:A:299:THR:HG21	1:A:597:VAL:HG21	1.84	0.59
1:A:1051:SER:OG	1:A:1064:HIS:ND1	2.33	0.59
1:B:521:PRO:HD3	1:C:41:LYS:HE2	1.84	0.59
1:A:34:ARG:NH1	1:A:219:GLY:O	2.35	0.59
1:C:598:ILE:HB	1:C:609:ALA:HB3	1.85	0.59
1:B:954:GLN:OE1	1:B:1014:ARG:NH1	2.36	0.59
1:C:240:THR:HG1	1:C:245:HIS:HE2	1.47	0.59
1:A:598:ILE:HB	1:A:609:ALA:HB3	1.84	0.59
1:A:1043:CYS:O	1:A:1064:HIS:ND1	2.36	0.59
1:C:726:ILE:HG22	1:C:1061:VAL:HG22	1.85	0.59
1:A:729:VAL:HG12	1:A:1059:GLY:HA2	1.85	0.59
1:A:988:GLU:OE2	1:C:383:SER:OG	2.20	0.59
1:A:896:ILE:HD12	1:A:897:PRO:HD2	1.85	0.59
1:B:777:ASN:OD1	1:B:1019:ARG:NH2	2.35	0.59
1:A:1129:VAL:HG12	1:B:917:TYR:HB3	1.84	0.59
1:A:37:TYR:OH	1:A:53:ASP:OD1	2.20	0.59
1:B:596:SER:HB2	1:B:611:LEU:HB3	1.85	0.59
1:A:372:ALA:O	1:C:403:ARG:NH2	2.32	0.59
1:C:726:ILE:HG22	1:C:1061:VAL:HG22	1.85	0.59
1:C:546:LEU:HD21	1:C:573:THR:HG21	1.85	0.59
1:B:990:GLU:HA	1:B:993:ILE:HG22	1.85	0.59
1:B:408:ARG:HH22	1:B:409:GLN:HE21	1.49	0.59
1:C:276:LEU:HD11	1:C:304:LYS:HA	1.85	0.59
1:C:461:LEU:HD22	1:C:465:GLU:HG2	1.84	0.59
1:A:389:ASP:OD1	1:A:528:LYS:NZ	2.35	0.59
1:A:517:LEU:HD11	1:B:983:ARG:HH11	1.68	0.59
1:A:99:ASN:ND2	1:A:177:MET:SD	2.76	0.58
1:A:970:PHE:HD2	1:A:996:LEU:HA	1.67	0.58
1:C:1031:GLU:HB3	1:C:1037:SER:HB2	1.84	0.58
1:A:729:VAL:HG12	1:A:1059:GLY:HA2	1.85	0.58
1:B:398:ASP:OD2	1:B:423:TYR:OH	2.21	0.58
1:A:226:LEU:HG	1:A:227:VAL:HG23	1.83	0.58
1:B:130:VAL:HB	1:B:168:PHE:HB3	1.85	0.58

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:B:770:ILE:HA	1:B:773:GLU:HG3	1.85	0.58
1:A:372:ALA:O	1:C:403:ARG:NH2	2.32	0.58
1:C:177:MET:SD	1:C:190:ARG:NH2	2.77	0.58
1:C:81:ASN:O	1:C:239:GLN:NE2	2.36	0.58
1:B:1030:SER:HA	1:B:1034:LEU:HD23	1.85	0.58
1:C:1030:SER:OG	1:C:1031:GLU:OE1	2.20	0.58
1:A:398:ASP:HB2	1:A:512:VAL:HG12	1.85	0.58
1:A:596:SER:HB2	1:A:611:LEU:HB3	1.86	0.58
1:A:866:THR:OG1	1:A:869:MET:SD	2.57	0.58
1:C:729:VAL:HG21	1:C:781:VAL:HG11	1.85	0.58
1:A:384:PRO:HG3	1:C:415:THR:HG21	1.85	0.58
1:A:129:LYS:NZ	1:A:166:CYS:SG	2.68	0.58
1:B:357:ARG:NH2	1:B:396:TYR:OH	2.37	0.58
1:A:727:LEU:HD11	1:A:1025:ALA:HB2	1.84	0.58
1:A:296:LEU:HB2	1:A:608:VAL:HG21	1.85	0.58
1:A:726:ILE:HG12	1:A:1061:VAL:HG23	1.86	0.58
1:A:401:VAL:HG22	1:A:509:ARG:HG2	1.85	0.58
1:C:384:PRO:HA	1:C:387:LEU:HD23	1.85	0.58
1:A:984:LEU:HB2	1:A:989:ALA:HB2	1.84	0.58
1:C:393:THR:HA	1:C:522:ALA:HA	1.85	0.58
1:C:611:LEU:HD22	1:C:678:ILE:HD11	1.86	0.58
1:A:44:ARG:NH2	1:A:279:TYR:OH	2.36	0.58
1:C:222:ALA:HB2	1:C:285:ILE:HB	1.85	0.58
1:B:436:TRP:HZ3	1:B:509:ARG:HD2	1.67	0.58
1:C:983:ARG:HG3	1:C:984:LEU:HD22	1.85	0.58
1:A:598:ILE:HB	1:A:609:ALA:HB3	1.85	0.58
1:A:566:GLY:HA2	1:B:43:PHE:HB3	1.86	0.58
1:A:902:MET:HE1	1:A:1049:LEU:HD23	1.86	0.58
1:B:385:THR:O	1:B:388:ASN:ND2	2.36	0.58
1:C:738:CYS:O	1:C:742:ILE:HB	2.03	0.58
1:C:1031:GLU:HB3	1:C:1037:SER:HB2	1.84	0.58
1:A:372:ALA:O	1:C:403:ARG:NH2	2.36	0.58
1:C:96:GLU:OE1	1:C:188:ASN:ND2	2.36	0.58
1:A:64:TRP:HE1	1:A:264:ALA:HB1	1.68	0.58
1:B:1051:SER:OG	1:B:1064:HIS:ND1	2.35	0.58
1:A:1091:ARG:NH1	1:A:1118:ASP:O	2.36	0.58
1:C:129:LYS:HZ3	1:C:168:PHE:H	1.52	0.58
1:B:733:LYS:HE3	1:B:771:ALA:HB1	1.85	0.58
1:A:598:ILE:HB	1:A:609:ALA:HB3	1.86	0.58
1:B:403:ARG:NH2	1:C:372:ALA:O	2.32	0.58
1:B:403:ARG:NH2	1:C:372:ALA:O	2.32	0.58

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:B:896:ILE:HD12	1:B:897:PRO:HD2	1.85	0.58
1:A:106:PHE:HB2	1:A:117:LEU:HB2	1.85	0.58
1:C:896:ILE:HD12	1:C:897:PRO:HD2	1.86	0.58
1:A:561:PRO:O	1:A:577:ARG:NH1	2.36	0.58
1:C:357:ARG:NH2	1:C:396:TYR:OH	2.37	0.58
1:B:38:TYR:HE2	1:B:224:GLU:HG3	1.68	0.58
1:A:299:THR:HG21	1:A:597:VAL:HG21	1.85	0.58
1:A:972:ALA:HB2	1:A:996:LEU:HD21	1.86	0.58
1:C:777:ASN:HB3	1:C:1019:ARG:HH21	1.69	0.58
1:A:195:LYS:HD2	1:A:197:ILE:HD11	1.86	0.58
1:B:34:ARG:HH22	1:B:220:PHE:HA	1.69	0.58
1:B:398:ASP:OD2	1:B:423:TYR:OH	2.22	0.58
1:C:193:VAL:HG22	1:C:223:LEU:HD23	1.85	0.58
1:A:129:LYS:NZ	1:A:168:PHE:O	2.33	0.58
1:B:28:TYR:HB3	1:B:61:ASN:HB3	1.85	0.58
1:C:598:ILE:HB	1:C:609:ALA:HB3	1.86	0.58
1:C:1031:GLU:HB3	1:C:1037:SER:HB2	1.85	0.58
1:A:372:ALA:O	1:C:403:ARG:NH2	2.36	0.58
1:B:995:ARG:HE	1:B:996:LEU:HD22	1.67	0.58
1:B:106:PHE:HB2	1:B:117:LEU:HB2	1.85	0.58
1:A:729:VAL:HG21	1:A:781:VAL:HG11	1.84	0.58
1:A:805:ILE:HD12	1:A:878:LEU:HD11	1.86	0.58
1:A:983:ARG:HE	1:C:517:LEU:HD11	1.68	0.58
1:A:774:GLN:OE1	1:A:777:ASN:ND2	2.37	0.58
1:A:115:GLN:HE22	1:C:468:ILE:HG12	1.68	0.58
1:A:84:LEU:HD12	1:A:85:PRO:HD2	1.84	0.58
1:C:915:VAL:HG12	1:C:1109:PHE:HD2	1.69	0.58
1:A:767:LEU:HA	1:A:770:ILE:HG22	1.85	0.58
1:B:866:THR:HG22	1:B:869:MET:HE2	1.84	0.58
1:C:552:LEU:HB3	1:C:585:LEU:HD22	1.86	0.58
1:A:729:VAL:HG21	1:A:781:VAL:HG11	1.84	0.58
1:C:101:ILE:HD11	1:C:240:THR:HB	1.85	0.58
1:C:226:LEU:HG	1:C:227:VAL:HG23	1.86	0.58
1:A:566:GLY:HA2	1:B:43:PHE:HB3	1.85	0.58
1:B:598:ILE:HB	1:B:609:ALA:HB3	1.85	0.58
1:B:896:ILE:HD12	1:B:897:PRO:HD2	1.85	0.58
1:B:64:TRP:HE1	1:B:264:ALA:HB1	1.69	0.58
1:A:276:LEU:HD11	1:A:304:LYS:HE2	1.85	0.58
1:C:357:ARG:NH2	1:C:396:TYR:OH	2.36	0.58
1:A:276:LEU:HG	1:A:306:PHE:HE1	1.69	0.58
1:B:677:PRO:HB2	1:C:864:LEU:HD11	1.86	0.58

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:C:316:SER:OG	1:C:593:GLY:O	2.21	0.58
1:A:517:LEU:HD11	1:B:983:ARG:HD2	1.86	0.58
1:B:357:ARG:NH2	1:B:396:TYR:OH	2.37	0.58
1:C:389:ASP:OD1	1:C:528:LYS:NZ	2.37	0.58
1:A:742:ILE:O	1:A:1000:ARG:NH1	2.36	0.58
1:C:738:CYS:HB2	1:C:763:LEU:HD11	1.85	0.58
1:B:702:GLU:OE2	1:C:790:LYS:NZ	2.36	0.58
1:A:310:LYS:HG2	1:A:664:ILE:HG21	1.86	0.58
1:A:389:ASP:OD1	1:A:528:LYS:NZ	2.37	0.58
1:A:911:VAL:HG11	1:A:1067:TYR:HE1	1.69	0.58
1:C:326:ILE:HD11	1:C:534:VAL:HG22	1.86	0.58
1:C:273:ARG:NH1	1:C:290:ASP:OD2	2.37	0.58
1:B:503:VAL:HA	1:B:506:GLN:HE21	1.69	0.57
1:A:389:ASP:OD1	1:A:528:LYS:NZ	2.36	0.57
1:C:966:LEU:HB2	1:C:1000:ARG:HH12	1.68	0.57
1:A:806:LEU:HD22	1:A:878:LEU:HD21	1.86	0.57
1:B:912:THR:OG1	1:B:1106:GLN:NE2	2.37	0.57
1:A:805:ILE:HD12	1:A:1054:GLN:HE21	1.69	0.57
1:B:738:CYS:O	1:B:742:ILE:HB	2.04	0.57
1:B:559:PHE:HB2	1:B:584:ILE:HD13	1.84	0.57
1:C:809:PRO:O	1:C:814:LYS:NZ	2.36	0.57
1:B:326:ILE:HD11	1:B:534:VAL:HG22	1.85	0.57
1:A:372:ALA:O	1:C:403:ARG:NH2	2.32	0.57
1:B:34:ARG:HH22	1:B:220:PHE:HA	1.69	0.57
1:B:415:THR:OG1	1:C:369:TYR:OH	2.18	0.57
1:A:738:CYS:HB3	1:A:763:LEU:HD11	1.85	0.57
1:C:357:ARG:NH2	1:C:396:TYR:OH	2.36	0.57
1:A:774:GLN:OE1	1:A:777:ASN:ND2	2.36	0.57
1:B:961:THR:HA	1:B:964:LYS:HD3	1.86	0.57
1:A:106:PHE:HB2	1:A:117:LEU:HB3	1.85	0.57
1:A:896:ILE:HD12	1:A:897:PRO:HD2	1.86	0.57
1:C:340:GLU:OE2	1:C:356:LYS:NZ	2.37	0.57
1:A:966:LEU:HA	1:A:1000:ARG:HH12	1.70	0.57
1:C:726:ILE:HG22	1:C:1061:VAL:HG22	1.85	0.57
1:B:106:PHE:HB2	1:B:117:LEU:HB3	1.87	0.57
1:A:393:THR:HA	1:A:522:ALA:HA	1.86	0.57
1:A:130:VAL:HG21	1:A:231:ILE:HD11	1.86	0.57
1:C:726:ILE:HG22	1:C:1061:VAL:HG22	1.85	0.57
1:B:902:MET:HE1	1:B:1050:MET:HE2	1.85	0.57
1:C:48:LEU:HB3	1:C:276:LEU:HD11	1.85	0.57
1:A:552:LEU:HB3	1:A:585:LEU:HD23	1.86	0.57

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:17:ASN:HB2	1:A:21:ARG:HD3	1.87	0.57
1:B:403:ARG:NH2	1:C:372:ALA:O	2.36	0.57
1:A:276:LEU:HG	1:A:306:PHE:HE1	1.68	0.57
1:A:1051:SER:OG	1:A:1064:HIS:ND1	2.35	0.57
1:A:129:LYS:NZ	1:A:168:PHE:O	2.36	0.57
1:B:1079:PRO:HB3	1:C:900:MET:HE1	1.86	0.57
1:C:1106:GLN:NE2	1:C:1111:GLU:OE2	2.36	0.57
1:B:1079:PRO:HB3	1:C:900:MET:HE1	1.84	0.57
1:A:966:LEU:HA	1:A:1000:ARG:HH12	1.68	0.57
1:A:676:ILE:HD12	1:A:677:PRO:HD2	1.86	0.57
1:A:1002:GLN:OE1	1:C:1002:GLN:NE2	2.37	0.57
1:C:426:PRO:HB3	1:C:464:PHE:HB2	1.87	0.57
1:A:1051:SER:OG	1:A:1064:HIS:ND1	2.36	0.57
1:B:770:ILE:HA	1:B:773:GLU:HG3	1.87	0.57
1:B:401:VAL:HG22	1:B:509:ARG:HG2	1.86	0.57
1:C:357:ARG:NH2	1:C:396:TYR:OH	2.37	0.57
1:A:369:TYR:HH	1:C:415:THR:HG1	1.52	0.57
1:C:978:ASN:HA	1:C:981:LEU:HG	1.86	0.57
1:C:774:GLN:HE22	1:C:1018:ILE:HG21	1.69	0.57
1:B:332:ILE:HA	1:B:524:VAL:HG22	1.86	0.57
1:B:905:ARG:NH1	1:B:1049:LEU:O	2.37	0.57
1:B:984:LEU:HD11	1:B:988:GLU:HB2	1.85	0.57
1:C:884:SER:HB3	1:C:887:THR:HB	1.85	0.57
1:A:563:GLN:O	1:A:577:ARG:NH2	2.37	0.57
1:A:946:GLY:HA2	1:A:949:GLN:HB3	1.86	0.57
1:A:1093:GLY:O	1:B:904:TYR:OH	2.18	0.57
1:B:457:ARG:NH1	1:B:467:ASP:OD2	2.33	0.57
1:B:229:LEU:HD12	1:B:230:PRO:HD2	1.87	0.57
1:B:360:ASN:H	1:B:523:THR:HB	1.69	0.57
1:C:129:LYS:NZ	1:C:166:CYS:SG	2.76	0.57
1:A:105:ILE:HG22	1:A:239:GLN:HB3	1.87	0.57
1:C:14:GLN:O	1:C:158:ARG:NH1	2.35	0.57
1:C:389:ASP:OD1	1:C:528:LYS:NZ	2.38	0.57
1:B:317:ASN:ND2	1:C:737:ASP:OD1	2.37	0.57
1:A:762:GLN:HA	1:A:765:ARG:HE	1.70	0.57
1:A:805:ILE:HD12	1:A:1054:GLN:HE21	1.69	0.57
1:C:775:ASP:HB3	1:C:864:LEU:HD23	1.86	0.57
1:B:129:LYS:NZ	1:B:168:PHE:O	2.33	0.57
1:A:106:PHE:HB2	1:A:117:LEU:HB2	1.86	0.57
1:C:185:ASN:ND2	1:C:211:ASN:OD1	2.37	0.57
1:A:102:ARG:NH1	1:A:121:ASN:O	2.37	0.57

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:B:598:ILE:HB	1:B:609:ALA:HB3	1.86	0.57
1:B:1079:PRO:HB3	1:C:900:MET:HE1	1.85	0.57
1:C:316:SER:OG	1:C:317:ASN:N	2.37	0.57
1:A:501:TYR:O	1:A:506:GLN:NE2	2.37	0.57
1:C:781:VAL:HG13	1:C:782:PHE:HD2	1.70	0.57
1:A:567:ARG:NH1	1:A:573:THR:OG1	2.38	0.57
1:B:384:PRO:HA	1:B:387:LEU:HD23	1.87	0.57
1:C:108:THR:O	1:C:237:ARG:NH2	2.38	0.57
1:C:177:MET:SD	1:C:190:ARG:NH2	2.78	0.57
1:B:362:VAL:HG21	1:B:526:GLY:H	1.70	0.57
1:C:126:VAL:HB	1:C:172:SER:HB3	1.86	0.57
1:A:745:ASP:OD1	1:C:319:ARG:NH1	2.38	0.57
1:B:86:PHE:HE1	1:B:89:GLY:HA2	1.70	0.57
1:A:47:VAL:HG12	1:C:569:ILE:HA	1.87	0.57
1:A:177:MET:SD	1:A:190:ARG:NH2	2.78	0.57
1:C:129:LYS:HZ1	1:C:167:THR:H	1.52	0.57
1:B:1030:SER:HA	1:B:1034:LEU:HD23	1.86	0.57
1:B:28:TYR:HB3	1:B:61:ASN:HB3	1.86	0.57
1:B:569:ILE:HG13	1:C:47:VAL:HG12	1.86	0.57
1:A:360:ASN:H	1:A:523:THR:HB	1.69	0.57
1:A:973:ILE:HG21	1:A:983:ARG:HH22	1.70	0.57
1:C:560:LEU:HD22	1:C:562:PHE:HB3	1.87	0.57
1:C:99:ASN:HB3	1:C:102:ARG:HH12	1.70	0.57
1:C:401:VAL:HG22	1:C:509:ARG:HG2	1.87	0.57
1:C:994:ASP:HA	1:C:997:ILE:HG12	1.85	0.57
1:C:375:SER:HB3	1:C:436:TRP:HA	1.87	0.57
1:B:37:TYR:OH	1:B:53:ASP:OD1	2.22	0.57
1:B:978:ASN:HA	1:B:981:LEU:HG	1.86	0.57
1:A:299:THR:HG21	1:A:597:VAL:HG21	1.87	0.57
1:B:129:LYS:NZ	1:B:168:PHE:O	2.35	0.57
1:B:767:LEU:HA	1:B:770:ILE:HG12	1.86	0.57
1:B:28:TYR:HB3	1:B:61:ASN:HB3	1.86	0.57
1:A:393:THR:HA	1:A:522:ALA:HA	1.87	0.57
1:C:101:ILE:HD11	1:C:240:THR:HB	1.87	0.57
1:A:276:LEU:HD11	1:A:304:LYS:HE2	1.85	0.57
1:B:403:ARG:NH2	1:C:372:ALA:O	2.36	0.57
1:C:741:TYR:HE1	1:C:1004:LEU:HD13	1.70	0.57
1:C:1010:GLN:HB3	1:C:1014:ARG:HH21	1.70	0.57
1:A:574:ASP:OD1	1:A:574:ASP:N	2.38	0.57
1:C:822:LEU:HD22	1:C:945:LEU:HD21	1.85	0.57
1:A:299:THR:HG21	1:A:597:VAL:HG21	1.87	0.57

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:C:340:GLU:OE2	1:C:356:LYS:NZ	2.38	0.57
1:A:1051:SER:OG	1:A:1064:HIS:ND1	2.36	0.57
1:B:14:GLN:N	1:B:255:SER:HG	2.02	0.57
1:B:48:LEU:HB3	1:B:276:LEU:HD11	1.87	0.57
1:A:384:PRO:HA	1:A:387:LEU:HD23	1.86	0.57
1:B:401:VAL:HG22	1:B:509:ARG:HG2	1.87	0.57
1:A:809:PRO:O	1:A:814:LYS:NZ	2.37	0.57
1:C:984:LEU:HG	1:C:989:ALA:HB2	1.86	0.57
1:C:276:LEU:HG	1:C:306:PHE:HE1	1.70	0.57
1:B:984:LEU:HB2	1:B:989:ALA:HB2	1.86	0.57
1:B:741:TYR:HE2	1:B:1004:LEU:HB2	1.70	0.56
1:B:353:TRP:HE1	1:B:355:ARG:HH21	1.52	0.56
1:B:376:THR:OG1	1:B:378:LYS:NZ	2.38	0.56
1:B:966:LEU:HA	1:B:1000:ARG:HH12	1.69	0.56
1:A:687:GLN:HG2	1:A:693:ILE:HD13	1.87	0.56
1:B:382:VAL:HG22	1:C:983:ARG:HH12	1.70	0.56
1:B:984:LEU:HB2	1:B:989:ALA:HB2	1.87	0.56
1:C:564:GLN:HG2	1:C:577:ARG:HD2	1.86	0.56
1:A:384:PRO:HG3	1:C:415:THR:HG21	1.87	0.56
1:C:384:PRO:HA	1:C:387:LEU:HD23	1.87	0.56
1:A:726:ILE:HG22	1:A:1061:VAL:HG22	1.87	0.56
1:C:440:ASN:OD1	1:C:440:ASN:N	2.38	0.56
1:C:1033:VAL:HG21	1:C:1053:PRO:HG3	1.87	0.56
1:A:1040:VAL:HG11	1:B:1035:GLY:HA3	1.86	0.56
1:A:105:ILE:HG23	1:A:239:GLN:HB3	1.88	0.56
1:B:576:VAL:HG12	1:B:587:ILE:HD11	1.87	0.56
1:A:761:THR:HG22	1:A:765:ARG:HH21	1.68	0.56
1:B:390:LEU:HD21	1:C:983:ARG:HA	1.86	0.56
1:C:119:ILE:HG12	1:C:128:ILE:HG12	1.87	0.56
1:A:517:LEU:HD11	1:B:983:ARG:HH21	1.70	0.56
1:B:102:ARG:NH1	1:B:121:ASN:O	2.37	0.56
1:A:439:ASN:OD1	1:A:506:GLN:NE2	2.38	0.56
1:A:326:ILE:HD11	1:A:534:VAL:HG12	1.86	0.56
1:B:390:LEU:HD21	1:C:983:ARG:HG2	1.87	0.56
1:C:326:ILE:HD11	1:C:534:VAL:HG22	1.88	0.56
1:B:35:GLY:HA3	1:B:56:LEU:HB3	1.86	0.56
1:A:896:ILE:HD12	1:A:897:PRO:HD2	1.86	0.56
1:B:777:ASN:HA	1:B:780:GLU:HG3	1.85	0.56
1:A:108:THR:O	1:A:237:ARG:NH2	2.39	0.56
1:A:1039:ARG:HD2	1:B:1039:ARG:HH22	1.70	0.56
1:A:34:ARG:HH22	1:A:220:PHE:HA	1.70	0.56

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:132:GLU:OE1	1:A:164:ASN:ND2	2.37	0.56
1:A:809:PRO:O	1:A:814:LYS:NZ	2.38	0.56
1:B:763:LEU:HB2	1:B:1008:VAL:HG11	1.88	0.56
1:B:384:PRO:HA	1:B:387:LEU:HD23	1.86	0.56
1:C:357:ARG:NH2	1:C:396:TYR:OH	2.38	0.56
1:C:908:GLY:O	1:C:1038:LYS:NZ	2.38	0.56
1:C:99:ASN:HB3	1:C:102:ARG:HH12	1.70	0.56
1:C:374:PHE:CG	1:C:434:ILE:HD11	2.40	0.56
1:B:759:PHE:HA	1:B:762:GLN:OE1	2.05	0.56
1:C:360:ASN:HB3	1:C:523:THR:HG22	1.85	0.56
1:C:901:GLN:OE1	1:C:905:ARG:NH1	2.38	0.56
1:C:735:SER:HB3	1:C:861:LEU:HD21	1.85	0.56
1:C:809:PRO:O	1:C:814:LYS:NZ	2.36	0.56
1:A:890:ALA:HA	1:C:1046:GLY:HA2	1.87	0.56
1:A:340:GLU:OE2	1:A:356:LYS:NZ	2.38	0.56
1:B:409:GLN:HA	1:B:414:GLN:HG2	1.86	0.56
1:C:129:LYS:HZ3	1:C:168:PHE:H	1.52	0.56
1:C:731:MET:H	1:C:774:GLN:HE21	1.52	0.56
1:A:353:TRP:HE1	1:A:466:ARG:HB2	1.71	0.56
1:A:702:GLU:OE2	1:B:790:LYS:NZ	2.35	0.56
1:C:357:ARG:NH2	1:C:396:TYR:OH	2.38	0.56
1:C:973:ILE:HG21	1:C:983:ARG:HH12	1.70	0.56
1:C:866:THR:H	1:C:869:MET:HE2	1.69	0.56
1:B:896:ILE:HD12	1:B:897:PRO:HD2	1.88	0.56
1:B:1002:GLN:HA	1:B:1005:GLN:HE21	1.70	0.56
1:A:80:ALA:O	1:A:245:HIS:NE2	2.38	0.56
1:B:733:LYS:HB2	1:B:861:LEU:HB2	1.86	0.56
1:C:383:SER:HB3	1:C:386:LYS:HE2	1.87	0.56
1:C:498:GLN:HE22	1:C:505:TYR:HD2	1.53	0.56
1:C:393:THR:HA	1:C:522:ALA:HA	1.86	0.56
1:A:569:ILE:HG13	1:B:47:VAL:HG23	1.87	0.56
1:C:877:LEU:HD21	1:C:1029:MET:HE1	1.87	0.56
1:B:1049:LEU:HB3	1:B:1050:MET:HE3	1.87	0.56
1:A:1115:ILE:HG22	1:A:1137:VAL:HG23	1.87	0.56
1:A:384:PRO:HG3	1:C:415:THR:HG21	1.87	0.56
1:C:52:GLN:OE1	1:C:274:THR:OG1	2.21	0.56
1:B:966:LEU:HA	1:B:1000:ARG:HH12	1.70	0.56
1:C:106:PHE:HD1	1:C:235:ILE:HD13	1.71	0.56
1:A:439:ASN:ND2	1:A:506:GLN:OE1	2.34	0.56
1:A:394:ASN:HB2	1:A:516:GLU:HB3	1.87	0.56
1:A:1031:GLU:HB3	1:A:1037:SER:HB2	1.85	0.56

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:B:562:PHE:O	1:C:41:LYS:NZ	2.39	0.56
1:C:1104:VAL:H	1:C:1115:ILE:HD11	1.71	0.56
1:B:360:ASN:H	1:B:523:THR:HB	1.69	0.56
1:C:777:ASN:HA	1:C:780:GLU:HG3	1.88	0.56
1:A:971:GLY:HA3	1:A:995:ARG:HH12	1.70	0.56
1:A:598:ILE:HB	1:A:609:ALA:HB3	1.87	0.56
1:A:562:PHE:O	1:B:41:LYS:NZ	2.38	0.56
1:A:802:PHE:HB3	1:A:806:LEU:HD23	1.86	0.56
1:A:130:VAL:HB	1:A:168:PHE:HB3	1.88	0.56
1:C:389:ASP:OD1	1:C:528:LYS:NZ	2.36	0.56
1:C:896:ILE:HD12	1:C:897:PRO:HD2	1.88	0.56
1:A:357:ARG:NH2	1:A:396:TYR:OH	2.38	0.56
1:A:954:GLN:OE1	1:A:1014:ARG:NH2	2.39	0.56
1:B:357:ARG:NH2	1:B:396:TYR:OH	2.38	0.56
1:A:697:MET:HB3	1:B:869:MET:HE1	1.88	0.56
1:A:869:MET:HG2	1:C:699:LEU:HD21	1.88	0.56
1:A:1035:GLY:HA3	1:C:1040:VAL:HG11	1.88	0.56
1:C:319:ARG:NH2	1:C:590:CYS:SG	2.78	0.56
1:C:401:VAL:HG22	1:C:509:ARG:HG2	1.88	0.56
1:A:310:LYS:HG3	1:A:600:PRO:HA	1.87	0.56
1:C:809:PRO:O	1:C:814:LYS:NZ	2.36	0.56
1:A:719:THR:HG22	1:A:1070:ALA:HB2	1.86	0.56
1:C:896:ILE:HD12	1:C:897:PRO:HD2	1.88	0.56
1:C:994:ASP:HA	1:C:997:ILE:HG12	1.86	0.56
1:A:988:GLU:OE1	1:C:383:SER:OG	2.24	0.56
1:A:106:PHE:HB2	1:A:117:LEU:HB2	1.88	0.56
1:B:216:LEU:HD12	1:B:217:PRO:HD2	1.86	0.56
1:A:334:ASN:ND2	1:A:360:ASN:O	2.34	0.56
1:A:199:GLY:HA2	1:A:232:GLY:HA2	1.86	0.56
1:A:1129:VAL:HG22	1:B:917:TYR:HB3	1.88	0.56
1:B:912:THR:OG1	1:B:1106:GLN:NE2	2.38	0.56
1:A:887:THR:HB	1:A:894:LEU:HD13	1.88	0.56
1:B:742:ILE:O	1:B:1000:ARG:NH1	2.39	0.56
1:B:598:ILE:HB	1:B:609:ALA:HB3	1.87	0.56
1:A:401:VAL:HG22	1:A:509:ARG:HG2	1.87	0.56
1:B:189:LEU:HD21	1:B:210:ILE:HD13	1.88	0.56
1:B:384:PRO:HA	1:B:387:LEU:HD23	1.88	0.56
1:C:106:PHE:HB2	1:C:117:LEU:HB2	1.88	0.56
1:B:398:ASP:OD2	1:B:423:TYR:OH	2.22	0.56
1:B:966:LEU:HA	1:B:1000:ARG:HH12	1.70	0.56
1:C:777:ASN:HB3	1:C:1019:ARG:HH21	1.70	0.56

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:C:195:LYS:HG3	1:C:197:ILE:HG12	1.88	0.56
1:A:914:ASN:HD21	1:C:1121:PHE:HE2	1.53	0.56
1:C:973:ILE:HG21	1:C:983:ARG:HH12	1.70	0.56
1:B:358:ILE:HB	1:B:395:VAL:HG23	1.86	0.56
1:C:395:VAL:HG12	1:C:515:PHE:HB3	1.86	0.56
1:A:1011:GLN:OE1	1:A:1014:ARG:NH1	2.37	0.56
1:A:384:PRO:HG3	1:C:415:THR:HG21	1.86	0.56
1:A:809:PRO:O	1:A:814:LYS:NZ	2.38	0.56
1:B:1051:SER:OG	1:B:1064:HIS:ND1	2.36	0.56
1:A:729:VAL:HG21	1:A:781:VAL:HG11	1.88	0.56
1:B:506:GLN:HG3	1:B:507:PRO:HD2	1.87	0.56
1:A:564:GLN:OE1	1:B:41:LYS:NZ	2.38	0.56
1:A:133:PHE:HE2	1:A:159:VAL:HG12	1.69	0.56
1:C:357:ARG:NH2	1:C:396:TYR:OH	2.38	0.56
1:A:369:TYR:HH	1:C:415:THR:HG1	1.51	0.56
1:C:353:TRP:HB2	1:C:399:SER:H	1.70	0.56
1:B:191:GLU:HG2	1:B:223:LEU:HD21	1.88	0.56
1:A:918:GLU:HG3	1:C:1128:VAL:HG21	1.87	0.56
1:C:34:ARG:HH22	1:C:220:PHE:HA	1.70	0.56
1:C:440:ASN:OD1	1:C:440:ASN:N	2.39	0.56
1:A:316:SER:OG	1:A:317:ASN:N	2.37	0.56
1:A:972:ALA:HB2	1:A:996:LEU:HD21	1.88	0.56
1:A:296:LEU:HB2	1:A:608:VAL:HG21	1.86	0.56
1:A:1013:ILE:HG21	1:B:1012:LEU:HB3	1.87	0.56
1:A:440:ASN:OD1	1:A:440:ASN:N	2.39	0.56
1:C:384:PRO:HA	1:C:387:LEU:HD23	1.88	0.56
1:C:393:THR:HA	1:C:522:ALA:HA	1.87	0.56
1:B:357:ARG:NH2	1:B:396:TYR:OH	2.39	0.56
1:C:401:VAL:HG22	1:C:509:ARG:HG2	1.87	0.56
1:B:670:ASN:C	1:B:670:ASN:ND2	2.64	0.56
1:A:34:ARG:HH22	1:A:220:PHE:HA	1.71	0.56
1:A:394:ASN:OD1	1:B:200:TYR:OH	2.23	0.56
1:A:1051:SER:OG	1:A:1064:HIS:ND1	2.33	0.56
1:A:57:PRO:HG3	1:A:273:ARG:HG3	1.88	0.56
1:B:357:ARG:NH2	1:B:396:TYR:OH	2.38	0.56
1:B:22:THR:OG1	1:B:78:ARG:NH1	2.38	0.56
1:A:954:GLN:HA	1:A:957:GLN:HG3	1.88	0.56
1:A:369:TYR:OH	1:C:415:THR:OG1	2.22	0.56
1:C:401:VAL:HG22	1:C:509:ARG:HG2	1.88	0.56
1:A:547:THR:O	1:B:978:ASN:ND2	2.34	0.56
1:A:300:LYS:NZ	1:A:306:PHE:O	2.39	0.56

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:B:966:LEU:HA	1:B:1000:ARG:HH12	1.70	0.56
1:B:355:ARG:NH1	1:C:230:PRO:O	2.38	0.56
1:B:519:HIS:HE1	1:C:42:VAL:HA	1.71	0.56
1:A:31:SER:OG	1:A:60:SER:O	2.24	0.56
1:A:461:LEU:HD12	1:A:465:GLU:HB3	1.86	0.56
1:A:316:SER:OG	1:A:317:ASN:N	2.39	0.56
1:C:425:LEU:HD22	1:C:512:VAL:HG11	1.87	0.56
1:B:457:ARG:NH1	1:B:467:ASP:OD1	2.36	0.56
1:C:984:LEU:HB2	1:C:989:ALA:HB2	1.87	0.56
1:B:966:LEU:HA	1:B:1000:ARG:HH12	1.71	0.56
1:C:712:ILE:HG12	1:C:1094:VAL:HG11	1.87	0.56
1:A:896:ILE:HD12	1:C:712:ILE:HD11	1.88	0.56
1:A:369:TYR:OH	1:C:415:THR:OG1	2.22	0.56
1:B:1017:GLU:OE2	1:C:1019:ARG:NH1	2.39	0.56
1:C:196:ASN:ND2	1:C:233:ILE:O	2.34	0.56
1:B:22:THR:O	1:B:78:ARG:NH1	2.39	0.56
1:C:454:ARG:NH1	1:C:469:SER:O	2.38	0.55
1:A:68:ILE:H	1:A:78:ARG:HB2	1.71	0.55
1:B:298:GLU:HB3	1:B:315:THR:HG21	1.89	0.55
1:A:332:ILE:HG13	1:A:362:VAL:HG22	1.87	0.55
1:B:317:ASN:HD21	1:B:592:PHE:HD2	1.55	0.55
1:A:401:VAL:HG22	1:A:509:ARG:HG2	1.88	0.55
1:A:401:VAL:HG22	1:A:509:ARG:HG2	1.88	0.55
1:B:726:ILE:HG22	1:B:1061:VAL:HG22	1.88	0.55
1:C:395:VAL:HG22	1:C:515:PHE:HB3	1.88	0.55
1:C:1010:GLN:OE1	1:C:1014:ARG:NH1	2.39	0.55
1:B:104:TRP:HD1	1:B:238:PHE:HZ	1.55	0.55
1:C:177:MET:SD	1:C:190:ARG:NH2	2.79	0.55
1:C:357:ARG:NH2	1:C:396:TYR:OH	2.39	0.55
1:A:296:LEU:HB3	1:A:608:VAL:HG21	1.86	0.55
1:B:726:ILE:HG22	1:B:1061:VAL:HG22	1.89	0.55
1:C:340:GLU:OE1	1:C:356:LYS:NZ	2.39	0.55
1:A:394:ASN:ND2	1:B:200:TYR:OH	2.38	0.55
1:C:1031:GLU:OE2	1:C:1039:ARG:NE	2.39	0.55
1:A:106:PHE:HB3	1:A:235:ILE:HD12	1.89	0.55
1:C:34:ARG:NH1	1:C:219:GLY:O	2.40	0.55
1:C:877:LEU:HG	1:C:1053:PRO:HG2	1.87	0.55
1:B:195:LYS:HZ3	1:B:202:LYS:HD3	1.71	0.55
1:C:994:ASP:HA	1:C:997:ILE:HG12	1.87	0.55
1:C:52:GLN:OE1	1:C:274:THR:OG1	2.25	0.55
1:B:357:ARG:NH2	1:B:396:TYR:OH	2.39	0.55

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:C:401:VAL:HG22	1:C:509:ARG:HG2	1.87	0.55
1:B:598:ILE:HD11	1:B:678:ILE:HD12	1.87	0.55
1:B:598:ILE:HB	1:B:609:ALA:HB3	1.86	0.55
1:B:1031:GLU:HB3	1:B:1037:SER:HB2	1.88	0.55
1:C:39:PRO:HG3	1:C:51:THR:HG21	1.87	0.55
1:A:372:ALA:O	1:C:403:ARG:NH2	2.39	0.55
1:B:994:ASP:HA	1:B:997:ILE:HG12	1.88	0.55
1:C:276:LEU:HG	1:C:306:PHE:HE1	1.71	0.55
1:A:753:LEU:HD21	1:A:997:ILE:HG21	1.88	0.55
1:B:1139:ASP:HB3	1:B:1142:GLN:HG2	1.87	0.55
1:A:34:ARG:NH1	1:A:219:GLY:O	2.39	0.55
1:A:338:PHE:HE1	1:A:368:LEU:HD23	1.71	0.55
1:A:611:LEU:HD22	1:A:678:ILE:HD11	1.88	0.55
1:B:336:CYS:HB3	1:B:358:ILE:HD12	1.89	0.55
1:B:442:ASP:O	1:B:448:ASN:ND2	2.39	0.55
1:C:394:ASN:ND2	1:C:516:GLU:OE1	2.35	0.55
1:C:1088:HIS:CE1	1:C:1122:VAL:HG22	2.41	0.55
1:C:384:PRO:HA	1:C:387:LEU:HD23	1.89	0.55
1:C:738:CYS:O	1:C:742:ILE:HB	2.06	0.55
1:A:57:PRO:O	1:A:60:SER:OG	2.24	0.55
1:A:108:THR:O	1:A:237:ARG:NH2	2.39	0.55
1:B:117:LEU:HD13	1:B:235:ILE:HD11	1.89	0.55
1:A:970:PHE:O	1:A:995:ARG:NH1	2.40	0.55
1:B:28:TYR:HB3	1:B:61:ASN:HB3	1.88	0.55
1:C:229:LEU:HD12	1:C:230:PRO:HD2	1.88	0.55
1:B:994:ASP:HA	1:B:997:ILE:HG12	1.88	0.55
1:A:606:ASN:O	1:A:606:ASN:ND2	2.39	0.55
1:C:64:TRP:HE1	1:C:264:ALA:HB1	1.72	0.55
1:C:440:ASN:OD1	1:C:440:ASN:N	2.40	0.55
1:A:566:GLY:HA2	1:B:43:PHE:HB3	1.88	0.55
1:C:729:VAL:HG12	1:C:1059:GLY:HA2	1.87	0.55
1:C:809:PRO:HA	1:C:814:LYS:HD2	1.86	0.55
1:B:384:PRO:HA	1:B:387:LEU:HD23	1.88	0.55
1:B:357:ARG:NH2	1:B:396:TYR:OH	2.40	0.55
1:A:717:ASN:HB3	1:A:1071:GLN:HB2	1.88	0.55
1:C:287:ASP:HB2	1:C:306:PHE:HE2	1.71	0.55
1:A:576:VAL:HG12	1:A:587:ILE:HD11	1.88	0.55
1:C:326:ILE:HD11	1:C:534:VAL:HG22	1.87	0.55
1:A:230:PRO:O	1:C:466:ARG:NH2	2.40	0.55
1:C:524:VAL:HG13	1:C:529:LYS:HD3	1.87	0.55
1:A:369:TYR:HH	1:C:415:THR:HG1	1.44	0.55

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:C:34:ARG:NH1	1:C:219:GLY:O	2.39	0.55
1:B:222:ALA:HB2	1:B:285:ILE:HB	1.87	0.55
1:B:358:ILE:HB	1:B:395:VAL:HG23	1.88	0.55
1:C:994:ASP:HA	1:C:997:ILE:HG12	1.88	0.55
1:A:1031:GLU:HB3	1:A:1037:SER:HB2	1.89	0.55
1:C:108:THR:O	1:C:237:ARG:NH2	2.40	0.55
1:A:729:VAL:HG21	1:A:781:VAL:HG11	1.89	0.55
1:A:887:THR:HB	1:A:894:LEU:HD23	1.89	0.55
1:B:978:ASN:HA	1:B:981:LEU:HG	1.88	0.55
1:A:972:ALA:HB2	1:A:996:LEU:HD21	1.87	0.55
1:C:966:LEU:HA	1:C:1000:ARG:HH12	1.71	0.55
1:A:312:ILE:HG22	1:A:664:ILE:HD11	1.88	0.55
1:A:328:ARG:NH2	1:A:580:GLN:OE1	2.40	0.55
1:B:357:ARG:NH2	1:B:396:TYR:OH	2.40	0.55
1:A:1129:VAL:HG22	1:B:917:TYR:HB3	1.88	0.55
1:B:950:ASP:OD1	1:B:950:ASP:N	2.38	0.55
1:B:596:SER:HB2	1:B:611:LEU:HB3	1.89	0.55
1:C:37:TYR:OH	1:C:53:ASP:OD1	2.24	0.55
1:C:809:PRO:O	1:C:814:LYS:NZ	2.38	0.55
1:B:276:LEU:HB3	1:B:289:VAL:HG22	1.88	0.55
1:C:398:ASP:HB2	1:C:512:VAL:HG12	1.88	0.55
1:C:58:PHE:HD2	1:C:290:ASP:HB2	1.72	0.55
1:C:815:ARG:HH11	1:C:823:PHE:HD1	1.55	0.55
1:B:358:ILE:HB	1:B:395:VAL:HG23	1.88	0.55
1:B:471:GLU:OE2	1:C:113:LYS:NZ	2.39	0.55
1:B:598:ILE:HB	1:B:609:ALA:HB3	1.89	0.55
1:B:735:SER:HB3	1:B:861:LEU:HD21	1.88	0.55
1:B:809:PRO:HA	1:B:814:LYS:HD2	1.87	0.55
1:B:102:ARG:HH21	1:B:246:ILE:HG12	1.72	0.55
1:C:729:VAL:HG12	1:C:1059:GLY:HA2	1.88	0.55
1:B:393:THR:HA	1:B:522:ALA:HA	1.89	0.55
1:C:99:ASN:O	1:C:102:ARG:NH1	2.40	0.55
1:C:553:THR:O	1:C:586:ASP:N	2.39	0.55
1:A:877:LEU:HD21	1:A:1029:MET:HE1	1.88	0.55
1:C:776:LYS:NZ	1:C:777:ASN:OD1	2.40	0.55
1:C:467:ASP:OD1	1:C:467:ASP:N	2.37	0.55
1:C:867:ASP:OD2	1:C:867:ASP:N	2.39	0.55
1:B:457:ARG:NH1	1:B:467:ASP:OD2	2.40	0.55
1:B:37:TYR:OH	1:B:53:ASP:OD1	2.24	0.55
1:C:140:PHE:HA	1:C:246:ILE:HG23	1.89	0.55
1:A:401:VAL:HG22	1:A:509:ARG:HG2	1.89	0.55

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:702:GLU:HA	1:B:788:ILE:HB	1.88	0.55
1:A:807:PRO:HA	1:A:816:SER:HA	1.88	0.55
1:B:662:CYS:HB2	1:B:697:MET:HG3	1.88	0.55
1:A:80:ALA:O	1:A:245:HIS:NE2	2.36	0.55
1:B:357:ARG:NH2	1:B:396:TYR:OH	2.38	0.55
1:C:896:ILE:HD12	1:C:897:PRO:HD2	1.88	0.55
1:A:115:GLN:HB2	1:A:233:ILE:HG12	1.89	0.55
1:B:702:GLU:OE2	1:C:790:LYS:NZ	2.35	0.55
1:B:888:PHE:HE2	1:B:1034:LEU:HA	1.71	0.55
1:C:37:TYR:OH	1:C:195:LYS:NZ	2.39	0.55
1:C:670:ASN:C	1:C:670:ASN:ND2	2.65	0.55
1:C:598:ILE:HD11	1:C:678:ILE:HD13	1.87	0.55
1:A:722:VAL:HG22	1:A:930:ALA:HB1	1.89	0.55
1:A:741:TYR:HD1	1:A:742:ILE:HD13	1.71	0.55
1:C:979:ASP:OD2	1:C:983:ARG:NH1	2.39	0.55
1:A:403:ARG:NH2	1:B:372:ALA:O	2.33	0.55
1:B:358:ILE:HB	1:B:395:VAL:HG23	1.88	0.55
1:A:806:LEU:HD12	1:A:807:PRO:HD2	1.87	0.55
1:C:276:LEU:HD13	1:C:301:CYS:HA	1.88	0.55
1:A:358:ILE:HB	1:A:395:VAL:HG23	1.87	0.55
1:A:719:THR:HG22	1:A:1070:ALA:HB2	1.89	0.55
1:B:200:TYR:HA	1:B:230:PRO:HA	1.89	0.55
1:C:357:ARG:NH2	1:C:396:TYR:OH	2.40	0.55
1:A:319:ARG:NH2	1:A:549:THR:OG1	2.40	0.55
1:A:403:ARG:NH2	1:B:372:ALA:O	2.32	0.55
1:C:360:ASN:H	1:C:523:THR:HB	1.72	0.55
1:C:994:ASP:HA	1:C:997:ILE:HG12	1.88	0.55
1:B:401:VAL:HG22	1:B:509:ARG:HG2	1.89	0.55
1:C:99:ASN:OD1	1:C:190:ARG:NH2	2.34	0.55
1:A:968:SER:OG	1:B:755:GLN:O	2.25	0.55
1:B:393:THR:HG21	1:B:518:LEU:HB2	1.88	0.55
1:B:296:LEU:O	1:B:299:THR:OG1	2.22	0.55
1:C:273:ARG:NH2	1:C:290:ASP:OD2	2.39	0.55
1:C:395:VAL:HG12	1:C:515:PHE:HB3	1.88	0.55
1:A:1091:ARG:NH1	1:A:1118:ASP:O	2.37	0.55
1:A:752:LEU:HD21	1:A:994:ASP:HB3	1.89	0.55
1:B:726:ILE:HG22	1:B:1061:VAL:HG22	1.88	0.55
1:B:37:TYR:OH	1:B:53:ASP:OD1	2.24	0.55
1:B:37:TYR:OH	1:B:53:ASP:OD1	2.25	0.55
1:C:340:GLU:OE2	1:C:356:LYS:NZ	2.39	0.55
1:C:809:PRO:HA	1:C:814:LYS:HD2	1.89	0.55

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:B:362:VAL:HG21	1:B:526:GLY:H	1.71	0.55
1:A:193:VAL:HG12	1:A:204:TYR:HB2	1.89	0.55
1:C:33:THR:OG1	1:C:219:GLY:O	2.25	0.55
1:C:729:VAL:HG21	1:C:781:VAL:HG11	1.89	0.55
1:C:1031:GLU:HB3	1:C:1037:SER:HB2	1.88	0.55
1:A:357:ARG:HD2	1:A:359:SER:HB2	1.88	0.55
1:C:52:GLN:OE1	1:C:274:THR:OG1	2.23	0.55
1:C:1031:GLU:HG3	1:C:1039:ARG:HH21	1.72	0.55
1:C:1033:VAL:HG21	1:C:1053:PRO:HG3	1.89	0.55
1:C:546:LEU:HD21	1:C:573:THR:HG21	1.88	0.55
1:B:738:CYS:O	1:B:742:ILE:HB	2.07	0.55
1:C:276:LEU:HD22	1:C:306:PHE:HE1	1.70	0.55
1:A:966:LEU:HA	1:A:1000:ARG:HH12	1.72	0.55
1:B:106:PHE:HB2	1:B:117:LEU:HB3	1.88	0.55
1:B:440:ASN:OD1	1:B:440:ASN:N	2.37	0.55
1:B:707:TYR:HD2	1:C:792:PRO:HG3	1.70	0.55
1:B:950:ASP:OD1	1:B:950:ASP:N	2.37	0.55
1:A:177:MET:SD	1:A:190:ARG:NH2	2.78	0.55
1:C:204:TYR:HB3	1:C:223:LEU:HB3	1.88	0.55
1:B:971:GLY:HA3	1:B:995:ARG:HH12	1.72	0.55
1:B:983:ARG:HG2	1:B:984:LEU:HD23	1.88	0.55
1:A:1005:GLN:HA	1:A:1008:VAL:HG22	1.88	0.55
1:C:276:LEU:HG	1:C:306:PHE:HE1	1.71	0.55
1:B:357:ARG:NH2	1:B:396:TYR:OH	2.39	0.55
1:A:1028:LYS:NZ	1:A:1042:PHE:O	2.38	0.55
1:C:276:LEU:HG	1:C:306:PHE:HE1	1.71	0.55
1:A:117:LEU:HD11	1:A:231:ILE:HG21	1.87	0.55
1:A:731:MET:H	1:A:774:GLN:HE21	1.55	0.55
1:C:735:SER:HB3	1:C:861:LEU:HD21	1.90	0.54
1:A:97:LYS:HE2	1:A:187:LYS:H	1.72	0.54
1:A:439:ASN:ND2	1:A:506:GLN:OE1	2.40	0.54
1:C:337:PRO:HD2	1:C:358:ILE:HD12	1.90	0.54
1:C:440:ASN:OD1	1:C:440:ASN:N	2.40	0.54
1:A:369:TYR:OH	1:C:415:THR:OG1	2.21	0.54
1:A:440:ASN:OD1	1:A:440:ASN:N	2.39	0.54
1:B:577:ARG:HH21	1:B:582:LEU:HD13	1.72	0.54
1:A:389:ASP:OD1	1:A:528:LYS:NZ	2.38	0.54
1:C:753:LEU:HD13	1:C:756:TYR:HD2	1.71	0.54
1:B:357:ARG:NH2	1:B:396:TYR:OH	2.39	0.54
1:A:973:ILE:HG21	1:A:983:ARG:HH22	1.72	0.54
1:B:358:ILE:HB	1:B:395:VAL:HG23	1.88	0.54

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:B:403:ARG:NH2	1:C:372:ALA:O	2.32	0.54
1:C:129:LYS:NZ	1:C:167:THR:OG1	2.40	0.54
1:A:278:LYS:HB2	1:A:306:PHE:HE1	1.71	0.54
1:B:269:TYR:O	1:B:271:GLN:NE2	2.39	0.54
1:A:596:SER:HB2	1:A:611:LEU:HB3	1.88	0.54
1:B:726:ILE:HG23	1:B:1061:VAL:HG22	1.90	0.54
1:C:99:ASN:OD1	1:C:190:ARG:NH2	2.40	0.54
1:B:269:TYR:O	1:B:271:GLN:NE2	2.40	0.54
1:C:389:ASP:OD1	1:C:528:LYS:NZ	2.41	0.54
1:B:501:TYR:O	1:B:506:GLN:NE2	2.40	0.54
1:C:1073:LYS:HE3	1:C:1075:PHE:HZ	1.71	0.54
1:B:777:ASN:HA	1:B:780:GLU:HG3	1.89	0.54
1:C:1031:GLU:HB3	1:C:1037:SER:HB2	1.89	0.54
1:C:319:ARG:NH2	1:C:590:CYS:SG	2.79	0.54
1:C:398:ASP:OD2	1:C:423:TYR:OH	2.18	0.54
1:A:113:LYS:NZ	1:C:471:GLU:OE2	2.34	0.54
1:C:393:THR:HA	1:C:522:ALA:HA	1.88	0.54
1:A:826:VAL:HB	1:A:1057:PRO:HG2	1.88	0.54
1:C:398:ASP:OD2	1:C:423:TYR:OH	2.24	0.54
1:C:606:ASN:O	1:C:606:ASN:ND2	2.40	0.54
1:A:340:GLU:OE2	1:A:356:LYS:NZ	2.40	0.54
1:B:598:ILE:HB	1:B:609:ALA:HB3	1.89	0.54
1:A:403:ARG:NH2	1:B:372:ALA:O	2.35	0.54
1:B:533:LEU:HD21	1:B:585:LEU:HD11	1.89	0.54
1:B:295:PRO:HA	1:B:298:GLU:HG3	1.89	0.54
1:B:403:ARG:NH2	1:C:372:ALA:O	2.32	0.54
1:A:819:GLU:HA	1:A:822:LEU:HB2	1.90	0.54
1:A:971:GLY:HA2	1:B:755:GLN:HE21	1.71	0.54
1:C:37:TYR:OH	1:C:195:LYS:NZ	2.38	0.54
1:C:108:THR:O	1:C:237:ARG:NH2	2.41	0.54
1:C:202:LYS:NZ	1:C:228:ASP:OD1	2.40	0.54
1:C:212:LEU:HD22	1:C:217:PRO:HD3	1.90	0.54
1:C:1081:ILE:HG13	1:C:1088:HIS:HB2	1.90	0.54
1:C:28:TYR:HB3	1:C:61:ASN:HB3	1.90	0.54
1:A:358:ILE:HB	1:A:395:VAL:HG23	1.89	0.54
1:C:994:ASP:HA	1:C:997:ILE:HG12	1.89	0.54
1:A:37:TYR:OH	1:A:53:ASP:OD1	2.22	0.54
1:A:1028:LYS:NZ	1:A:1042:PHE:O	2.40	0.54
1:B:1091:ARG:NH2	1:B:1120:THR:O	2.40	0.54
1:A:58:PHE:HD1	1:A:290:ASP:HB2	1.72	0.54
1:A:384:PRO:HA	1:A:387:LEU:HD23	1.88	0.54

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:440:ASN:OD1	1:A:440:ASN:N	2.40	0.54
1:C:287:ASP:HB2	1:C:306:PHE:HE2	1.71	0.54
1:B:206:LYS:NZ	1:B:222:ALA:O	2.39	0.54
1:B:316:SER:OG	1:B:317:ASN:N	2.40	0.54
1:B:949:GLN:HA	1:B:952:VAL:HG22	1.89	0.54
1:B:403:ARG:NH2	1:C:372:ALA:O	2.30	0.54
1:B:360:ASN:H	1:B:523:THR:HB	1.72	0.54
1:C:357:ARG:NH2	1:C:396:TYR:OH	2.40	0.54
1:C:440:ASN:OD1	1:C:440:ASN:N	2.40	0.54
1:A:299:THR:HG21	1:A:597:VAL:HG21	1.89	0.54
1:B:560:LEU:HD23	1:B:562:PHE:H	1.72	0.54
1:A:276:LEU:HD11	1:A:304:LYS:HE3	1.88	0.54
1:A:825:LYS:HZ3	1:A:942:ALA:HA	1.72	0.54
1:A:316:SER:HB3	1:A:595:VAL:HG22	1.89	0.54
1:B:535:LYS:NZ	1:B:554:GLU:OE2	2.40	0.54
1:A:226:LEU:HG	1:A:227:VAL:HG23	1.88	0.54
1:A:890:ALA:HA	1:C:1046:GLY:HA2	1.89	0.54
1:A:403:ARG:NH2	1:B:372:ALA:O	2.31	0.54
1:A:756:TYR:HB3	1:A:759:PHE:HD2	1.73	0.54
1:A:1002:GLN:NE2	1:B:1005:GLN:OE1	2.40	0.54
1:B:365:TYR:O	1:B:369:TYR:HB2	2.08	0.54
1:A:966:LEU:HA	1:A:1000:ARG:HH12	1.72	0.54
1:A:33:THR:OG1	1:A:219:GLY:O	2.26	0.54
1:B:104:TRP:HZ3	1:B:240:THR:HG22	1.72	0.54
1:A:972:ALA:HB2	1:A:996:LEU:HD21	1.90	0.54
1:A:809:PRO:O	1:A:814:LYS:NZ	2.39	0.54
1:A:904:TYR:HE2	1:A:913:GLN:HE21	1.54	0.54
1:B:858:LEU:HD23	1:B:959:LEU:HD12	1.90	0.54
1:B:340:GLU:OE2	1:B:356:LYS:NZ	2.41	0.54
1:B:358:ILE:HB	1:B:395:VAL:HG23	1.89	0.54
1:C:598:ILE:HB	1:C:609:ALA:HB3	1.89	0.54
1:B:384:PRO:HA	1:B:387:LEU:HD23	1.89	0.54
1:A:349:SER:OG	1:A:452:LEU:O	2.25	0.54
1:B:737:ASP:HB3	1:B:740:MET:HG3	1.90	0.54
1:B:358:ILE:HB	1:B:395:VAL:HG23	1.89	0.54
1:B:393:THR:HA	1:B:522:ALA:HA	1.90	0.54
1:A:34:ARG:HH22	1:A:220:PHE:HA	1.71	0.54
1:A:600:PRO:HD3	1:A:692:ILE:HD11	1.90	0.54
1:A:576:VAL:HG12	1:A:587:ILE:HD11	1.89	0.54
1:B:357:ARG:NH2	1:B:396:TYR:OH	2.40	0.54
1:C:727:LEU:HD22	1:C:1025:ALA:HB2	1.89	0.54

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:C:319:ARG:NH2	1:C:590:CYS:SG	2.79	0.54
1:A:774:GLN:OE1	1:A:777:ASN:ND2	2.36	0.54
1:A:1010:GLN:OE1	1:A:1014:ARG:NH2	2.41	0.54
1:B:358:ILE:HB	1:B:395:VAL:HG23	1.90	0.54
1:A:756:TYR:HB3	1:A:759:PHE:HD2	1.72	0.54
1:B:82:PRO:HG2	1:B:84:LEU:HD21	1.89	0.54
1:A:598:ILE:HD11	1:A:678:ILE:HD12	1.89	0.54
1:A:809:PRO:O	1:A:814:LYS:NZ	2.40	0.54
1:B:440:ASN:OD1	1:B:440:ASN:N	2.41	0.54
1:C:21:ARG:HA	1:C:79:PHE:HB3	1.89	0.54
1:A:393:THR:HA	1:A:522:ALA:HA	1.89	0.54
1:C:389:ASP:OD1	1:C:528:LYS:NZ	2.38	0.54
1:B:384:PRO:HA	1:B:387:LEU:HD23	1.90	0.54
1:A:37:TYR:OH	1:A:53:ASP:OD1	2.22	0.54
1:C:596:SER:HB2	1:C:611:LEU:HB3	1.88	0.54
1:C:353:TRP:HE1	1:C:466:ARG:HB2	1.73	0.54
1:C:401:VAL:HG22	1:C:509:ARG:HG2	1.89	0.54
1:C:788:ILE:HD12	1:C:876:ALA:HB2	1.89	0.54
1:B:759:PHE:HA	1:B:762:GLN:HG3	1.88	0.54
1:C:83:VAL:HG22	1:C:239:GLN:HG2	1.90	0.54
1:C:130:VAL:HB	1:C:168:PHE:HB2	1.89	0.54
1:A:1013:ILE:HD13	1:B:1012:LEU:HD12	1.90	0.54
1:B:340:GLU:OE2	1:B:356:LYS:NZ	2.41	0.54
1:B:966:LEU:HA	1:B:1000:ARG:HH12	1.73	0.54
1:C:57:PRO:HG3	1:C:273:ARG:HD2	1.89	0.54
1:A:406:GLU:HG2	1:A:418:ILE:HG13	1.89	0.54
1:B:457:ARG:NH1	1:B:467:ASP:OD2	2.38	0.54
1:C:394:ASN:N	1:C:394:ASN:OD1	2.40	0.54
1:B:293:LEU:HD23	1:B:294:ASP:HB3	1.89	0.54
1:B:314:GLN:NE2	1:B:316:SER:O	2.40	0.54
1:A:858:LEU:HD13	1:A:959:LEU:HB3	1.88	0.54
1:C:994:ASP:HA	1:C:997:ILE:HG12	1.89	0.54
1:C:598:ILE:HB	1:C:609:ALA:HB3	1.89	0.54
1:A:226:LEU:HG	1:A:227:VAL:HG23	1.88	0.54
1:B:37:TYR:OH	1:B:53:ASP:OD1	2.23	0.54
1:B:908:GLY:O	1:B:1038:LYS:NZ	2.35	0.54
1:B:398:ASP:OD2	1:B:423:TYR:OH	2.23	0.54
1:B:1002:GLN:HA	1:B:1005:GLN:HG3	1.88	0.54
1:B:360:ASN:O	1:B:360:ASN:ND2	2.39	0.54
1:C:353:TRP:HE1	1:C:466:ARG:HB2	1.72	0.54
1:C:1051:SER:OG	1:C:1064:HIS:ND1	2.37	0.54

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:140:PHE:HD1	1:A:246:ILE:HG12	1.73	0.54
1:A:742:ILE:O	1:A:1000:ARG:NH1	2.40	0.54
1:B:719:THR:HG22	1:B:1070:ALA:HB2	1.88	0.54
1:A:440:ASN:OD1	1:A:440:ASN:N	2.39	0.54
1:C:193:VAL:HB	1:C:204:TYR:HB2	1.89	0.54
1:C:1031:GLU:HB3	1:C:1037:SER:HB2	1.90	0.54
1:C:276:LEU:HG	1:C:306:PHE:HE1	1.73	0.54
1:B:353:TRP:HH2	1:B:464:PHE:HA	1.73	0.54
1:B:403:ARG:NH2	1:C:372:ALA:O	2.32	0.54
1:B:1051:SER:OG	1:B:1064:HIS:ND1	2.37	0.54
1:B:401:VAL:HG22	1:B:509:ARG:HG2	1.89	0.54
1:B:34:ARG:NH1	1:B:219:GLY:O	2.41	0.54
1:A:1027:THR:HG22	1:A:1042:PHE:HZ	1.73	0.54
1:C:193:VAL:HB	1:C:204:TYR:HB2	1.90	0.54
1:B:295:PRO:HG2	1:B:608:VAL:HG11	1.89	0.54
1:B:596:SER:HB2	1:B:611:LEU:HB3	1.90	0.54
1:B:708:SER:HB3	1:B:711:SER:HB3	1.88	0.54
1:C:52:GLN:OE1	1:C:274:THR:OG1	2.23	0.54
1:A:642:VAL:HG22	1:A:651:ILE:HG22	1.90	0.54
1:A:1051:SER:OG	1:A:1064:HIS:ND1	2.39	0.54
1:C:729:VAL:O	1:C:777:ASN:ND2	2.41	0.54
1:C:915:VAL:HG11	1:C:1108:ASN:HB2	1.90	0.54
1:B:401:VAL:HG12	1:B:509:ARG:HG2	1.90	0.54
1:B:708:SER:HB3	1:B:711:SER:HB3	1.90	0.54
1:B:28:TYR:HB3	1:B:61:ASN:HB3	1.89	0.54
1:C:877:LEU:HD23	1:C:1053:PRO:HG2	1.89	0.54
1:B:52:GLN:OE1	1:B:274:THR:OG1	2.24	0.54
1:B:707:TYR:HD2	1:C:792:PRO:HG3	1.73	0.54
1:B:1010:GLN:HB3	1:B:1014:ARG:HH12	1.72	0.54
1:C:887:THR:HG21	1:C:894:LEU:HG	1.89	0.54
1:B:596:SER:HB2	1:B:611:LEU:HB3	1.90	0.54
1:C:398:ASP:OD1	1:C:398:ASP:N	2.39	0.54
1:B:401:VAL:HG22	1:B:509:ARG:HG2	1.89	0.54
1:B:106:PHE:HB2	1:B:117:LEU:HB3	1.89	0.54
1:B:317:ASN:HD21	1:B:592:PHE:HD1	1.55	0.54
1:A:805:ILE:HD12	1:A:1054:GLN:HE21	1.73	0.54
1:A:1115:ILE:HG22	1:A:1137:VAL:HG23	1.89	0.54
1:A:971:GLY:HA3	1:A:995:ARG:HH12	1.72	0.54
1:B:418:ILE:HD12	1:B:422:ASN:HD22	1.71	0.54
1:C:756:TYR:HB3	1:C:759:PHE:HD1	1.72	0.54
1:B:34:ARG:NH1	1:B:219:GLY:O	2.41	0.54

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:B:738:CYS:O	1:B:742:ILE:HB	2.08	0.54
1:A:338:PHE:HE2	1:A:368:LEU:HD23	1.73	0.54
1:A:802:PHE:HB3	1:A:806:LEU:HD23	1.90	0.54
1:C:398:ASP:OD2	1:C:423:TYR:OH	2.26	0.54
1:A:969:ASN:HD21	1:B:755:GLN:HG2	1.72	0.54
1:B:139:PRO:O	1:B:245:HIS:ND1	2.40	0.54
1:B:442:ASP:OD1	1:B:509:ARG:NH2	2.41	0.54
1:B:726:ILE:HD13	1:B:1061:VAL:HG22	1.90	0.54
1:B:102:ARG:HH21	1:B:246:ILE:HG12	1.73	0.54
1:B:440:ASN:OD1	1:B:440:ASN:N	2.41	0.54
1:A:745:ASP:OD1	1:C:319:ARG:NH1	2.34	0.54
1:C:688:THR:HA	1:C:690:GLN:HG2	1.89	0.54
1:B:177:MET:HG2	1:B:190:ARG:HH22	1.72	0.54
1:A:440:ASN:OD1	1:A:440:ASN:N	2.40	0.54
1:B:28:TYR:HB3	1:B:61:ASN:HB3	1.88	0.54
1:C:278:LYS:NZ	1:C:286:THR:OG1	2.41	0.54
1:C:131:CYS:SG	1:C:132:GLU:N	2.80	0.54
1:A:86:PHE:HB2	1:A:238:PHE:HD1	1.72	0.53
1:C:337:PRO:HG3	1:C:356:LYS:HD3	1.89	0.53
1:A:115:GLN:HB2	1:A:233:ILE:HD13	1.90	0.53
1:C:599:THR:HB	1:C:608:VAL:HG23	1.89	0.53
1:B:226:LEU:HG	1:B:227:VAL:HG23	1.89	0.53
1:C:357:ARG:NH2	1:C:396:TYR:OH	2.41	0.53
1:C:501:TYR:O	1:C:506:GLN:NE2	2.41	0.53
1:C:1051:SER:OG	1:C:1064:HIS:ND1	2.36	0.53
1:B:326:ILE:HD11	1:B:534:VAL:HG22	1.90	0.53
1:B:398:ASP:HB2	1:B:512:VAL:HG12	1.91	0.53
1:C:1033:VAL:HG21	1:C:1053:PRO:HG3	1.89	0.53
1:C:727:LEU:HD21	1:C:1024:LEU:HG	1.90	0.53
1:A:599:THR:HB	1:A:608:VAL:HG23	1.89	0.53
1:A:315:THR:HG23	1:A:595:VAL:HG23	1.90	0.53
1:A:401:VAL:HG22	1:A:509:ARG:HG2	1.89	0.53
1:B:762:GLN:HA	1:B:765:ARG:HE	1.73	0.53
1:A:58:PHE:HD1	1:A:290:ASP:HB2	1.72	0.53
1:A:384:PRO:HA	1:A:387:LEU:HD23	1.90	0.53
1:B:1116:THR:OG1	1:B:1118:ASP:OD1	2.26	0.53
1:C:735:SER:HB3	1:C:861:LEU:HD21	1.90	0.53
1:B:731:MET:H	1:B:774:GLN:HE21	1.55	0.53
1:C:126:VAL:HG22	1:C:172:SER:HB3	1.90	0.53
1:C:393:THR:OG1	1:C:394:ASN:OD1	2.24	0.53
1:A:386:LYS:NZ	1:B:984:LEU:O	2.36	0.53

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:B:726:ILE:HD13	1:B:1061:VAL:HG22	1.90	0.53
1:C:326:ILE:HD12	1:C:539:VAL:HG21	1.90	0.53
1:A:791:THR:HG21	1:A:806:LEU:HD11	1.89	0.53
1:B:44:ARG:NH2	1:B:279:TYR:OH	2.42	0.53
1:C:384:PRO:HA	1:C:387:LEU:HD23	1.89	0.53
1:B:440:ASN:OD1	1:B:440:ASN:N	2.39	0.53
1:C:222:ALA:HB2	1:C:285:ILE:HB	1.90	0.53
1:A:759:PHE:HA	1:A:762:GLN:HE21	1.73	0.53
1:C:1031:GLU:HG3	1:C:1039:ARG:HH21	1.73	0.53
1:A:604:THR:OG1	1:A:605:SER:N	2.41	0.53
1:B:1040:VAL:HG11	1:C:1035:GLY:HA3	1.89	0.53
1:A:34:ARG:NH1	1:A:219:GLY:O	2.41	0.53
1:C:389:ASP:OD1	1:C:528:LYS:NZ	2.41	0.53
1:C:440:ASN:OD1	1:C:440:ASN:N	2.41	0.53
1:A:39:PRO:HG3	1:A:51:THR:HG21	1.90	0.53
1:A:1051:SER:OG	1:A:1064:HIS:ND1	2.35	0.53
1:B:679:GLY:HA2	1:C:864:LEU:HD12	1.89	0.53
1:A:299:THR:HG21	1:A:597:VAL:HG21	1.91	0.53
1:A:358:ILE:HB	1:A:395:VAL:HG23	1.90	0.53
1:B:719:THR:HG22	1:B:1070:ALA:HB2	1.89	0.53
1:A:393:THR:HA	1:A:522:ALA:HA	1.91	0.53
1:C:858:LEU:HD22	1:C:959:LEU:HD12	1.90	0.53
1:A:312:ILE:HD11	1:A:596:SER:HB3	1.90	0.53
1:B:121:ASN:HD21	1:B:177:MET:HE3	1.73	0.53
1:C:484:LYS:HD2	1:C:490:PHE:HB2	1.89	0.53
1:B:357:ARG:NH2	1:B:394:ASN:OD1	2.41	0.53
1:C:966:LEU:HA	1:C:1000:ARG:HH12	1.73	0.53
1:A:52:GLN:OE1	1:A:274:THR:OG1	2.23	0.53
1:A:276:LEU:HD11	1:A:304:LYS:HE2	1.91	0.53
1:A:908:GLY:O	1:A:1038:LYS:NZ	2.30	0.53
1:C:1031:GLU:HB3	1:C:1037:SER:HB2	1.89	0.53
1:C:729:VAL:HG21	1:C:781:VAL:HG11	1.91	0.53
1:B:87:ASN:OD1	1:B:87:ASN:N	2.40	0.53
1:B:384:PRO:HA	1:B:387:LEU:HD23	1.91	0.53
1:A:998:THR:O	1:A:1002:GLN:NE2	2.41	0.53
1:B:452:LEU:HD12	1:B:492:LEU:HD12	1.91	0.53
1:A:130:VAL:HB	1:A:168:PHE:HB3	1.89	0.53
1:C:574:ASP:OD1	1:C:574:ASP:N	2.39	0.53
1:A:358:ILE:HB	1:A:395:VAL:HG23	1.89	0.53
1:A:393:THR:HA	1:A:522:ALA:HA	1.89	0.53
1:B:384:PRO:HA	1:B:387:LEU:HD22	1.90	0.53

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:384:PRO:HA	1:A:387:LEU:HD23	1.91	0.53
1:A:719:THR:HG22	1:A:1070:ALA:HB2	1.89	0.53
1:A:994:ASP:HA	1:A:997:ILE:HG12	1.89	0.53
1:B:408:ARG:NH2	1:C:376:THR:OG1	2.40	0.53
1:B:611:LEU:HD22	1:B:666:ILE:HD11	1.90	0.53
1:A:688:THR:HA	1:A:690:GLN:HG2	1.89	0.53
1:C:326:ILE:HD11	1:C:534:VAL:HG22	1.89	0.53
1:B:289:VAL:HG11	1:B:300:LYS:HB2	1.91	0.53
1:B:994:ASP:HA	1:B:997:ILE:HG12	1.90	0.53
1:B:365:TYR:O	1:B:369:TYR:HB2	2.08	0.53
1:C:729:VAL:HG21	1:C:781:VAL:HG11	1.89	0.53
1:C:788:ILE:HG13	1:C:876:ALA:HB2	1.90	0.53
1:C:905:ARG:HE	1:C:1050:MET:HE3	1.72	0.53
1:C:58:PHE:HD2	1:C:290:ASP:HB2	1.74	0.53
1:C:126:VAL:HB	1:C:172:SER:HB3	1.90	0.53
1:C:37:TYR:OH	1:C:53:ASP:OD1	2.24	0.53
1:C:887:THR:HB	1:C:894:LEU:HD23	1.89	0.53
1:A:809:PRO:HA	1:A:814:LYS:HD2	1.90	0.53
1:A:34:ARG:HH22	1:A:220:PHE:HA	1.74	0.53
1:A:972:ALA:HB2	1:A:996:LEU:HD21	1.90	0.53
1:C:809:PRO:O	1:C:814:LYS:NZ	2.36	0.53
1:A:91:TYR:OH	1:A:191:GLU:OE2	2.18	0.53
1:A:129:LYS:NZ	1:A:168:PHE:O	2.42	0.53
1:A:440:ASN:OD1	1:A:440:ASN:N	2.39	0.53
1:B:1030:SER:HA	1:B:1034:LEU:HD23	1.91	0.53
1:C:994:ASP:HA	1:C:997:ILE:HG12	1.90	0.53
1:A:34:ARG:HH22	1:A:220:PHE:HA	1.72	0.53
1:B:418:ILE:HD12	1:B:422:ASN:HD22	1.73	0.53
1:B:401:VAL:HG22	1:B:509:ARG:HG2	1.89	0.53
1:B:777:ASN:HA	1:B:780:GLU:HG3	1.91	0.53
1:B:384:PRO:HA	1:B:387:LEU:HD23	1.89	0.53
1:A:87:ASN:OD1	1:A:87:ASN:N	2.41	0.53
1:B:411:ALA:HB3	1:B:414:GLN:HG2	1.89	0.53
1:B:571:ASP:OD2	1:C:975:SER:OG	2.25	0.53
1:C:503:VAL:HA	1:C:506:GLN:HG3	1.91	0.53
1:A:735:SER:HA	1:A:767:LEU:HD23	1.89	0.53
1:A:908:GLY:O	1:A:1038:LYS:NZ	2.38	0.53
1:A:314:GLN:HE22	1:A:594:GLY:HA3	1.73	0.53
1:A:759:PHE:HA	1:A:762:GLN:HE21	1.72	0.53
1:C:326:ILE:HD11	1:C:534:VAL:HG22	1.90	0.53
1:C:401:VAL:HG22	1:C:509:ARG:HG2	1.90	0.53

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:C:117:LEU:HD11	1:C:231:ILE:HG12	1.91	0.53
1:B:323:THR:OG1	1:B:324:GLU:OE2	2.21	0.53
1:A:1051:SER:OG	1:A:1064:HIS:ND1	2.33	0.53
1:C:598:ILE:HB	1:C:609:ALA:HB3	1.91	0.53
1:B:457:ARG:NH1	1:B:467:ASP:OD1	2.41	0.53
1:B:960:ASN:O	1:B:964:LYS:HB2	2.09	0.53
1:C:393:THR:OG1	1:C:394:ASN:OD1	2.25	0.53
1:B:358:ILE:HB	1:B:395:VAL:HG23	1.90	0.53
1:C:125:ASN:HB2	1:C:172:SER:H	1.73	0.53
1:C:1116:THR:OG1	1:C:1118:ASP:OD1	2.25	0.53
1:C:598:ILE:HB	1:C:609:ALA:HB3	1.91	0.53
1:C:408:ARG:HD2	1:C:414:GLN:HE21	1.73	0.53
1:C:897:PRO:HB2	1:C:900:MET:HG2	1.91	0.53
1:A:177:MET:SD	1:A:190:ARG:NH2	2.82	0.53
1:A:745:ASP:OD1	1:C:319:ARG:NH1	2.41	0.53
1:B:102:ARG:HH21	1:B:246:ILE:HG12	1.73	0.53
1:A:731:MET:SD	1:A:774:GLN:NE2	2.81	0.53
1:C:897:PRO:HG2	1:C:900:MET:HG2	1.91	0.53
1:A:562:PHE:HE1	1:B:225:PRO:HG2	1.74	0.53
1:B:454:ARG:HH12	1:B:457:ARG:HD2	1.73	0.53
1:A:129:LYS:NZ	1:A:166:CYS:SG	2.67	0.53
1:C:598:ILE:HD11	1:C:678:ILE:HD12	1.90	0.53
1:C:326:ILE:HD11	1:C:534:VAL:HG22	1.90	0.53
1:B:398:ASP:OD2	1:B:423:TYR:OH	2.22	0.53
1:A:994:ASP:HA	1:A:997:ILE:HG12	1.89	0.53
1:C:1010:GLN:NE2	1:C:1011:GLN:OE1	2.42	0.53
1:A:762:GLN:HB3	1:A:765:ARG:HH21	1.74	0.53
1:B:384:PRO:HA	1:B:387:LEU:HD23	1.89	0.53
1:C:357:ARG:NH2	1:C:396:TYR:OH	2.41	0.53
1:C:22:THR:O	1:C:78:ARG:NH1	2.42	0.53
1:A:547:THR:O	1:B:978:ASN:ND2	2.42	0.53
1:A:87:ASN:OD1	1:A:87:ASN:N	2.39	0.53
1:B:440:ASN:OD1	1:B:440:ASN:N	2.39	0.53
1:B:976:VAL:HG13	1:B:979:ASP:HB3	1.91	0.53
1:A:357:ARG:NH2	1:A:396:TYR:OH	2.41	0.53
1:A:729:VAL:HG21	1:A:781:VAL:HG11	1.90	0.53
1:A:994:ASP:HA	1:A:997:ILE:HG12	1.90	0.53
1:B:994:ASP:HA	1:B:997:ILE:HG12	1.91	0.53
1:C:34:ARG:HH22	1:C:220:PHE:HA	1.73	0.53
1:A:960:ASN:O	1:A:964:LYS:HB2	2.09	0.53
1:B:358:ILE:HB	1:B:395:VAL:HG23	1.91	0.53

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:B:326:ILE:HD11	1:B:534:VAL:HG12	1.90	0.53
1:C:984:LEU:HD12	1:C:988:GLU:HB2	1.91	0.53
1:B:598:ILE:HB	1:B:609:ALA:HB3	1.89	0.53
1:C:101:ILE:HD11	1:C:240:THR:HB	1.90	0.53
1:A:599:THR:HB	1:A:608:VAL:HG23	1.91	0.53
1:A:141:LEU:HD13	1:A:154:GLU:HG3	1.91	0.53
1:C:454:ARG:HH21	1:C:491:PRO:HB2	1.74	0.53
1:A:296:LEU:HB2	1:A:608:VAL:HG21	1.91	0.53
1:A:440:ASN:OD1	1:A:440:ASN:N	2.40	0.53
1:A:99:ASN:OD1	1:A:190:ARG:NH2	2.42	0.53
1:B:326:ILE:HD11	1:B:534:VAL:HG22	1.90	0.53
1:A:1115:ILE:HG22	1:A:1137:VAL:HG23	1.90	0.53
1:A:712:ILE:HG12	1:A:1077:THR:HB	1.90	0.53
1:B:383:SER:OG	1:C:983:ARG:O	2.22	0.53
1:C:712:ILE:HG12	1:C:1077:THR:HB	1.90	0.53
1:C:396:TYR:HB2	1:C:514:SER:HB2	1.89	0.53
1:B:729:VAL:HG12	1:B:1059:GLY:HA2	1.89	0.53
1:A:440:ASN:OD1	1:A:440:ASN:N	2.39	0.53
1:C:722:VAL:HG12	1:C:1065:VAL:HB	1.90	0.53
1:A:52:GLN:OE1	1:A:274:THR:OG1	2.23	0.53
1:A:80:ALA:O	1:A:245:HIS:NE2	2.41	0.53
1:A:393:THR:HA	1:A:522:ALA:HA	1.91	0.53
1:C:1031:GLU:HG3	1:C:1039:ARG:HH21	1.74	0.53
1:B:966:LEU:HA	1:B:1000:ARG:HH12	1.73	0.53
1:A:809:PRO:HA	1:A:814:LYS:HD2	1.91	0.53
1:C:759:PHE:HA	1:C:762:GLN:HE21	1.73	0.53
1:B:357:ARG:NH2	1:B:396:TYR:OH	2.42	0.53
1:B:393:THR:HA	1:B:522:ALA:HA	1.89	0.53
1:B:930:ALA:HA	1:B:933:LYS:HG2	1.91	0.53
1:C:52:GLN:OE1	1:C:274:THR:OG1	2.24	0.53
1:C:973:ILE:HG21	1:C:983:ARG:HH12	1.73	0.53
1:A:102:ARG:HE	1:A:246:ILE:HD11	1.73	0.53
1:B:401:VAL:HG22	1:B:509:ARG:HG2	1.90	0.53
1:A:805:ILE:HD12	1:A:878:LEU:HD11	1.91	0.53
1:B:440:ASN:OD1	1:B:440:ASN:N	2.39	0.53
1:C:877:LEU:O	1:C:881:THR:OG1	2.27	0.53
1:A:189:LEU:HD22	1:A:208:THR:HB	1.91	0.53
1:B:503:VAL:HA	1:B:506:GLN:HG3	1.91	0.53
1:C:129:LYS:NZ	1:C:167:THR:OG1	2.42	0.53
1:A:105:ILE:HG23	1:A:239:GLN:HB3	1.90	0.53
1:B:222:ALA:HB2	1:B:285:ILE:HB	1.89	0.53

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:C:729:VAL:HG12	1:C:1059:GLY:HA2	1.91	0.53
1:C:40:ASP:N	1:C:40:ASP:OD1	2.40	0.53
1:C:34:ARG:HH22	1:C:220:PHE:HA	1.74	0.53
1:A:997:ILE:O	1:A:1001:LEU:HB3	2.09	0.53
1:B:440:ASN:OD1	1:B:440:ASN:N	2.39	0.53
1:C:319:ARG:NH2	1:C:590:CYS:SG	2.82	0.53
1:A:756:TYR:HB3	1:A:759:PHE:HD2	1.74	0.53
1:A:976:VAL:HG13	1:A:979:ASP:HB3	1.89	0.53
1:A:1128:VAL:HG21	1:B:918:GLU:HG3	1.90	0.53
1:B:99:ASN:OD1	1:B:190:ARG:NH2	2.41	0.53
1:A:1035:GLY:HA3	1:C:1040:VAL:HG11	1.91	0.52
1:C:322:PRO:HG3	1:C:538:CYS:HB3	1.90	0.52
1:A:64:TRP:HE1	1:A:264:ALA:HB1	1.74	0.52
1:A:106:PHE:HB2	1:A:117:LEU:HB2	1.91	0.52
1:C:777:ASN:HA	1:C:780:GLU:HG3	1.91	0.52
1:A:58:PHE:HD2	1:A:290:ASP:HB2	1.74	0.52
1:A:930:ALA:HA	1:A:933:LYS:HG2	1.90	0.52
1:C:302:THR:O	1:C:304:LYS:NZ	2.43	0.52
1:C:326:ILE:HD11	1:C:534:VAL:HG22	1.91	0.52
1:A:92:PHE:HB3	1:A:192:PHE:HB2	1.91	0.52
1:C:326:ILE:HD13	1:C:533:LEU:HD12	1.91	0.52
1:C:992:GLN:OE1	1:C:995:ARG:NH2	2.42	0.52
1:A:1035:GLY:HA3	1:C:1040:VAL:HG11	1.90	0.52
1:B:719:THR:HG22	1:B:1070:ALA:HB2	1.89	0.52
1:B:965:GLN:OE1	1:C:758:SER:OG	2.26	0.52
1:A:403:ARG:NH1	1:A:405:ASP:OD1	2.42	0.52
1:C:398:ASP:OD2	1:C:423:TYR:OH	2.26	0.52
1:C:117:LEU:HB2	1:C:233:ILE:HD11	1.91	0.52
1:C:994:ASP:HA	1:C:997:ILE:HG12	1.92	0.52
1:C:902:MET:HB3	1:C:916:LEU:HD11	1.91	0.52
1:C:1031:GLU:HG3	1:C:1039:ARG:HH11	1.73	0.52
1:C:357:ARG:NH2	1:C:396:TYR:OH	2.42	0.52
1:A:759:PHE:HA	1:A:762:GLN:HE21	1.74	0.52
1:B:37:TYR:HB3	1:B:223:LEU:HB2	1.91	0.52
1:B:440:ASN:OD1	1:B:440:ASN:N	2.42	0.52
1:C:966:LEU:HA	1:C:1000:ARG:HH12	1.74	0.52
1:C:195:LYS:HG2	1:C:202:LYS:HB2	1.91	0.52
1:C:315:THR:HG23	1:C:595:VAL:HG23	1.91	0.52
1:A:561:PRO:HA	1:A:577:ARG:HH22	1.74	0.52
1:B:92:PHE:HE2	1:B:104:TRP:HE1	1.56	0.52
1:A:35:GLY:HA3	1:A:56:LEU:HB3	1.91	0.52

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:295:PRO:HG2	1:A:608:VAL:HG11	1.90	0.52
1:B:401:VAL:HG22	1:B:509:ARG:HG2	1.90	0.52
1:B:401:VAL:HG22	1:B:509:ARG:HG2	1.91	0.52
1:C:1033:VAL:HG21	1:C:1053:PRO:HG3	1.91	0.52
1:C:599:THR:HB	1:C:608:VAL:HG23	1.92	0.52
1:C:750:SER:HA	1:C:753:LEU:HG	1.90	0.52
1:A:777:ASN:HA	1:A:780:GLU:HG3	1.90	0.52
1:A:102:ARG:HE	1:A:246:ILE:HD11	1.74	0.52
1:A:822:LEU:HD22	1:A:945:LEU:HD11	1.91	0.52
1:C:14:GLN:O	1:C:158:ARG:NE	2.43	0.52
1:A:99:ASN:OD1	1:A:190:ARG:NH2	2.42	0.52
1:A:759:PHE:HA	1:A:762:GLN:HE21	1.75	0.52
1:C:360:ASN:H	1:C:523:THR:HB	1.74	0.52
1:C:858:LEU:HD22	1:C:959:LEU:HD12	1.90	0.52
1:C:40:ASP:N	1:C:40:ASP:OD1	2.42	0.52
1:A:719:THR:HG22	1:A:1070:ALA:HB2	1.90	0.52
1:A:1010:GLN:HE22	1:A:1014:ARG:HD2	1.73	0.52
1:A:1028:LYS:NZ	1:A:1042:PHE:O	2.41	0.52
1:A:1107:ARG:HH21	1:B:904:TYR:HD2	1.55	0.52
1:B:968:SER:OG	1:C:755:GLN:O	2.27	0.52
1:A:822:LEU:HD22	1:A:945:LEU:HD11	1.92	0.52
1:B:112:SER:HB2	1:B:132:GLU:HB3	1.92	0.52
1:B:58:PHE:HD2	1:B:290:ASP:HB2	1.74	0.52
1:A:986:PRO:HD2	1:A:987:PRO:HD3	1.91	0.52
1:A:34:ARG:HH22	1:A:220:PHE:HA	1.73	0.52
1:A:340:GLU:OE2	1:A:356:LYS:NZ	2.43	0.52
1:B:327:VAL:O	1:B:531:THR:OG1	2.25	0.52
1:A:762:GLN:HA	1:A:765:ARG:HE	1.74	0.52
1:B:319:ARG:NH2	1:B:590:CYS:SG	2.82	0.52
1:C:733:LYS:NZ	1:C:862:PRO:O	2.34	0.52
1:A:894:LEU:HD21	1:C:715:PRO:HD3	1.91	0.52
1:A:393:THR:HA	1:A:522:ALA:HA	1.90	0.52
1:A:701:VAL:HG13	1:B:787:GLN:HG2	1.91	0.52
1:C:37:TYR:OH	1:C:53:ASP:OD1	2.27	0.52
1:C:994:ASP:HA	1:C:997:ILE:HG12	1.91	0.52
1:C:458:LYS:HB3	1:C:474:GLN:HB3	1.91	0.52
1:C:887:THR:HG21	1:C:894:LEU:HB2	1.92	0.52
1:B:720:ILE:HD11	1:B:1065:VAL:HB	1.91	0.52
1:C:440:ASN:OD1	1:C:440:ASN:N	2.39	0.52
1:C:877:LEU:O	1:C:881:THR:OG1	2.27	0.52
1:B:760:CYS:O	1:B:764:ASN:HB2	2.09	0.52

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:C:586:ASP:N	1:C:586:ASP:OD1	2.40	0.52
1:C:1030:SER:HA	1:C:1034:LEU:HD23	1.91	0.52
1:B:358:ILE:HB	1:B:395:VAL:HG23	1.91	0.52
1:C:586:ASP:N	1:C:586:ASP:OD1	2.41	0.52
1:C:902:MET:HB3	1:C:916:LEU:HD11	1.91	0.52
1:A:1116:THR:OG1	1:A:1118:ASP:OD1	2.27	0.52
1:B:384:PRO:HA	1:B:387:LEU:HG	1.90	0.52
1:C:599:THR:HB	1:C:608:VAL:HG23	1.91	0.52
1:C:40:ASP:N	1:C:40:ASP:OD1	2.43	0.52
1:B:317:ASN:HD21	1:B:592:PHE:HD2	1.57	0.52
1:B:357:ARG:NH2	1:B:396:TYR:OH	2.42	0.52
1:C:45:SER:HA	1:C:279:TYR:HB3	1.92	0.52
1:A:1083:HIS:HD2	1:A:1137:VAL:H	1.56	0.52
1:B:440:ASN:OD1	1:B:440:ASN:N	2.41	0.52
1:C:791:THR:HG21	1:C:806:LEU:HD11	1.92	0.52
1:B:418:ILE:HD12	1:B:422:ASN:HB2	1.91	0.52
1:B:972:ALA:HB2	1:B:996:LEU:HD21	1.90	0.52
1:C:576:VAL:HG12	1:C:587:ILE:HD11	1.92	0.52
1:C:310:LYS:HG2	1:C:676:ILE:HG21	1.92	0.52
1:A:416:GLY:HA2	1:B:369:TYR:HE2	1.74	0.52
1:B:440:ASN:OD1	1:B:440:ASN:N	2.40	0.52
1:A:784:GLN:NE2	1:C:1041:ASP:OD2	2.42	0.52
1:B:323:THR:OG1	1:B:324:GLU:OE1	2.18	0.52
1:C:326:ILE:HD13	1:C:533:LEU:HA	1.89	0.52
1:B:902:MET:HE1	1:B:1050:MET:HE2	1.92	0.52
1:A:563:GLN:HG2	1:B:43:PHE:HD1	1.75	0.52
1:B:1013:ILE:HD13	1:C:1012:LEU:HG	1.92	0.52
1:C:34:ARG:HH22	1:C:220:PHE:HA	1.74	0.52
1:B:440:ASN:OD1	1:B:440:ASN:N	2.39	0.52
1:A:130:VAL:HB	1:A:168:PHE:HB3	1.92	0.52
1:A:328:ARG:HB2	1:A:543:PHE:HD2	1.74	0.52
1:B:756:TYR:OH	1:B:994:ASP:OD1	2.27	0.52
1:B:737:ASP:HB3	1:B:740:MET:HE3	1.90	0.52
1:A:33:THR:OG1	1:A:219:GLY:O	2.25	0.52
1:C:809:PRO:O	1:C:814:LYS:NZ	2.40	0.52
1:A:858:LEU:HD13	1:A:959:LEU:HB3	1.91	0.52
1:C:276:LEU:HG	1:C:306:PHE:HE1	1.75	0.52
1:C:742:ILE:O	1:C:1000:ARG:NH1	2.34	0.52
1:B:102:ARG:HH21	1:B:246:ILE:HG12	1.74	0.52
1:C:731:MET:HE2	1:C:955:ASN:HD21	1.74	0.52
1:B:393:THR:HA	1:B:522:ALA:HA	1.91	0.52

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:299:THR:HG21	1:A:597:VAL:HG21	1.92	0.52
1:B:440:ASN:OD1	1:B:440:ASN:N	2.39	0.52
1:A:87:ASN:OD1	1:A:87:ASN:N	2.42	0.52
1:C:22:THR:O	1:C:78:ARG:NH1	2.43	0.52
1:A:908:GLY:O	1:A:1038:LYS:NZ	2.34	0.52
1:B:28:TYR:HB3	1:B:61:ASN:HB3	1.90	0.52
1:B:99:ASN:O	1:B:102:ARG:NH2	2.43	0.52
1:C:1067:TYR:HE2	1:C:1108:ASN:HD22	1.58	0.52
1:B:362:VAL:HG21	1:B:526:GLY:H	1.75	0.52
1:C:950:ASP:N	1:C:950:ASP:OD1	2.42	0.52
1:A:37:TYR:OH	1:A:53:ASP:OD1	2.27	0.52
1:B:90:VAL:HG21	1:B:238:PHE:HE2	1.75	0.52
1:A:877:LEU:HD21	1:A:1029:MET:HE1	1.91	0.52
1:B:771:ALA:HA	1:B:774:GLN:HG2	1.92	0.52
1:A:389:ASP:OD1	1:A:528:LYS:NZ	2.40	0.52
1:C:440:ASN:N	1:C:440:ASN:OD1	2.39	0.52
1:A:777:ASN:HA	1:A:780:GLU:HG3	1.92	0.52
1:B:22:THR:OG1	1:B:78:ARG:NH1	2.40	0.52
1:C:901:GLN:HE21	1:C:1050:MET:HE1	1.74	0.52
1:C:99:ASN:OD1	1:C:190:ARG:NH2	2.41	0.52
1:A:147:LYS:HB3	1:A:153:MET:HE2	1.92	0.52
1:C:81:ASN:O	1:C:239:GLN:NE2	2.35	0.52
1:B:350:VAL:HG21	1:B:418:ILE:HD11	1.92	0.52
1:C:1052:PHE:HB2	1:C:1063:LEU:HD12	1.92	0.52
1:A:729:VAL:HG21	1:A:781:VAL:HG11	1.92	0.52
1:A:440:ASN:OD1	1:A:440:ASN:N	2.40	0.52
1:B:314:GLN:HE21	1:B:314:GLN:HA	1.74	0.52
1:A:291:CYS:HB2	1:A:298:GLU:HA	1.91	0.52
1:B:901:GLN:HE21	1:B:905:ARG:HE	1.58	0.52
1:A:106:PHE:HB2	1:A:117:LEU:HB3	1.92	0.52
1:A:599:THR:HB	1:A:608:VAL:HG23	1.91	0.52
1:A:719:THR:HG22	1:A:1070:ALA:HB2	1.92	0.52
1:B:203:ILE:HG13	1:B:226:LEU:HB3	1.91	0.52
1:C:401:VAL:HG22	1:C:509:ARG:HG2	1.92	0.52
1:A:33:THR:OG1	1:A:219:GLY:O	2.28	0.52
1:A:688:THR:HA	1:A:690:GLN:HG2	1.91	0.52
1:B:28:TYR:HB3	1:B:61:ASN:HB3	1.91	0.52
1:B:720:ILE:HD11	1:B:1065:VAL:HG13	1.91	0.52
1:A:34:ARG:NH1	1:A:219:GLY:O	2.42	0.52
1:B:86:PHE:HE1	1:B:89:GLY:HA2	1.74	0.52
1:C:93:ALA:HA	1:C:191:GLU:HA	1.92	0.52

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:C:37:TYR:OH	1:C:54:LEU:O	2.28	0.52
1:C:759:PHE:HA	1:C:762:GLN:HE21	1.74	0.52
1:B:93:ALA:HB3	1:B:266:TYR:HB2	1.91	0.52
1:B:1115:ILE:HG22	1:B:1137:VAL:HG23	1.92	0.52
1:C:599:THR:HB	1:C:608:VAL:HG23	1.92	0.52
1:C:319:ARG:NH2	1:C:590:CYS:SG	2.79	0.52
1:C:877:LEU:O	1:C:881:THR:OG1	2.27	0.52
1:B:317:ASN:HD21	1:B:592:PHE:HD2	1.57	0.52
1:B:902:MET:HE1	1:B:1050:MET:HE2	1.92	0.52
1:B:326:ILE:HD11	1:B:534:VAL:HG22	1.92	0.52
1:B:731:MET:H	1:B:774:GLN:HE22	1.58	0.52
1:A:403:ARG:NH2	1:B:372:ALA:O	2.36	0.52
1:A:994:ASP:HA	1:A:997:ILE:HG12	1.90	0.52
1:A:357:ARG:HH12	1:A:394:ASN:HD22	1.58	0.52
1:B:54:LEU:HB3	1:B:270:LEU:HD13	1.92	0.52
1:C:398:ASP:OD2	1:C:423:TYR:OH	2.20	0.52
1:B:889:GLY:HA3	1:B:1034:LEU:HD11	1.92	0.52
1:A:930:ALA:HA	1:A:933:LYS:HE2	1.91	0.52
1:C:393:THR:OG1	1:C:394:ASN:OD1	2.27	0.52
1:B:950:ASP:OD1	1:B:950:ASP:N	2.42	0.52
1:C:546:LEU:HD21	1:C:573:THR:HG21	1.92	0.52
1:B:87:ASN:C	1:B:87:ASN:HD22	2.15	0.52
1:B:92:PHE:HB3	1:B:192:PHE:HB2	1.92	0.52
1:B:735:SER:HB3	1:B:861:LEU:HD21	1.91	0.52
1:B:48:LEU:HD12	1:B:276:LEU:HD21	1.92	0.52
1:B:326:ILE:HD11	1:B:534:VAL:HG22	1.91	0.52
1:C:403:ARG:NH1	1:C:405:ASP:OD1	2.42	0.52
1:C:1115:ILE:HG22	1:C:1137:VAL:HG23	1.91	0.52
1:A:895:GLN:HE22	1:C:706:ALA:HB3	1.74	0.52
1:A:994:ASP:HA	1:A:997:ILE:HG12	1.90	0.52
1:A:809:PRO:HA	1:A:814:LYS:HD2	1.91	0.52
1:B:440:ASN:OD1	1:B:440:ASN:N	2.40	0.52
1:A:574:ASP:O	1:A:587:ILE:N	2.38	0.52
1:C:53:ASP:OD1	1:C:54:LEU:N	2.43	0.52
1:A:92:PHE:HB3	1:A:192:PHE:HB2	1.90	0.52
1:B:108:THR:OG1	1:B:234:ASN:O	2.28	0.52
1:C:806:LEU:HD12	1:C:807:PRO:HD2	1.92	0.52
1:B:37:TYR:OH	1:B:53:ASP:OD1	2.27	0.52
1:A:951:VAL:HA	1:A:954:GLN:HG3	1.92	0.52
1:B:104:TRP:HZ3	1:B:240:THR:HG22	1.74	0.52
1:B:398:ASP:HB2	1:B:512:VAL:HG12	1.91	0.52

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:C:751:ASN:HA	1:C:754:LEU:HG	1.92	0.52
1:A:1010:GLN:HE22	1:A:1014:ARG:HD2	1.75	0.52
1:B:978:ASN:HA	1:B:981:LEU:HG	1.91	0.52
1:C:276:LEU:HG	1:C:306:PHE:HE1	1.74	0.52
1:A:384:PRO:HG3	1:C:415:THR:HG21	1.91	0.52
1:C:887:THR:HG21	1:C:894:LEU:HB2	1.90	0.52
1:A:133:PHE:HE1	1:A:160:TYR:HB3	1.75	0.52
1:C:326:ILE:HD11	1:C:534:VAL:HG22	1.90	0.52
1:C:326:ILE:HD11	1:C:534:VAL:HG22	1.90	0.52
1:C:1010:GLN:HE22	1:C:1014:ARG:HH21	1.58	0.52
1:A:503:VAL:HA	1:A:506:GLN:HG3	1.91	0.52
1:C:729:VAL:HG12	1:C:1059:GLY:HA2	1.92	0.52
1:A:84:LEU:HD12	1:A:85:PRO:HD2	1.91	0.52
1:C:64:TRP:HE1	1:C:264:ALA:HB1	1.75	0.52
1:B:994:ASP:HA	1:B:997:ILE:HG12	1.92	0.52
1:B:389:ASP:OD1	1:B:389:ASP:N	2.42	0.52
1:B:406:GLU:HB3	1:B:418:ILE:HG21	1.92	0.52
1:C:401:VAL:HG22	1:C:509:ARG:HG2	1.92	0.51
1:B:106:PHE:HB2	1:B:117:LEU:HB3	1.93	0.51
1:B:915:VAL:HG12	1:B:1109:PHE:HD2	1.74	0.51
1:C:53:ASP:OD1	1:C:54:LEU:N	2.43	0.51
1:B:393:THR:HA	1:B:522:ALA:HA	1.92	0.51
1:A:574:ASP:OD1	1:A:574:ASP:N	2.41	0.51
1:C:395:VAL:HG12	1:C:515:PHE:HB3	1.92	0.51
1:B:355:ARG:HH22	1:C:230:PRO:HB2	1.76	0.51
1:C:212:LEU:HD22	1:C:217:PRO:HD3	1.92	0.51
1:A:881:THR:HG22	1:A:882:ILE:HD13	1.90	0.51
1:C:741:TYR:HH	1:C:1003:SER:HG	1.58	0.51
1:A:384:PRO:HA	1:A:387:LEU:HD23	1.92	0.51
1:C:599:THR:HB	1:C:608:VAL:HG23	1.92	0.51
1:B:440:ASN:OD1	1:B:440:ASN:N	2.43	0.51
1:A:905:ARG:NH2	1:A:1049:LEU:O	2.42	0.51
1:A:328:ARG:NH1	1:A:531:THR:O	2.39	0.51
1:A:791:THR:HG21	1:A:806:LEU:HD11	1.91	0.51
1:B:298:GLU:HB3	1:B:315:THR:HG21	1.92	0.51
1:B:440:ASN:OD1	1:B:440:ASN:N	2.40	0.51
1:C:467:ASP:OD2	1:C:467:ASP:N	2.41	0.51
1:A:299:THR:HG21	1:A:597:VAL:HG21	1.92	0.51
1:A:33:THR:OG1	1:A:219:GLY:O	2.25	0.51
1:A:389:ASP:OD1	1:A:528:LYS:NZ	2.42	0.51
1:B:571:ASP:OD2	1:C:975:SER:OG	2.29	0.51

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:C:1116:THR:OG1	1:C:1118:ASP:OD1	2.24	0.51
1:B:733:LYS:NZ	1:B:862:PRO:O	2.43	0.51
1:A:559:PHE:HB2	1:A:584:ILE:HD13	1.92	0.51
1:A:574:ASP:OD1	1:A:574:ASP:N	2.42	0.51
1:A:919:ASN:HD22	1:A:922:LEU:HD21	1.75	0.51
1:B:189:LEU:HD12	1:B:217:PRO:HG2	1.92	0.51
1:B:381:GLY:HA2	1:C:984:LEU:HD13	1.90	0.51
1:B:736:VAL:HG22	1:B:858:LEU:HG	1.92	0.51
1:C:1010:GLN:HE22	1:C:1014:ARG:HH21	1.58	0.51
1:A:986:PRO:HD2	1:A:987:PRO:HD3	1.93	0.51
1:B:777:ASN:HA	1:B:780:GLU:HG3	1.93	0.51
1:C:401:VAL:HG22	1:C:509:ARG:HG2	1.92	0.51
1:A:334:ASN:ND2	1:A:360:ASN:O	2.41	0.51
1:A:887:THR:HB	1:A:894:LEU:HD23	1.91	0.51
1:B:276:LEU:HG	1:B:306:PHE:HE1	1.76	0.51
1:B:566:GLY:HA2	1:C:43:PHE:HB3	1.93	0.51
1:A:52:GLN:OE1	1:A:274:THR:OG1	2.22	0.51
1:B:300:LYS:HE3	1:B:306:PHE:HA	1.93	0.51
1:A:353:TRP:HH2	1:A:464:PHE:HA	1.75	0.51
1:A:1031:GLU:OE1	1:A:1039:ARG:NH1	2.39	0.51
1:A:319:ARG:HH11	1:B:744:GLY:HA3	1.75	0.51
1:C:271:GLN:HG2	1:C:272:PRO:HD2	1.91	0.51
1:C:807:PRO:HA	1:C:816:SER:HA	1.92	0.51
1:A:598:ILE:HB	1:A:609:ALA:HB3	1.92	0.51
1:B:15:CYS:SG	1:B:137:ASN:N	2.80	0.51
1:C:394:ASN:N	1:C:394:ASN:OD1	2.42	0.51
1:C:599:THR:HB	1:C:608:VAL:HG23	1.93	0.51
1:A:68:ILE:H	1:A:78:ARG:HB2	1.74	0.51
1:C:330:PRO:HD3	1:C:579:PRO:HB2	1.93	0.51
1:A:52:GLN:OE1	1:A:274:THR:OG1	2.23	0.51
1:C:809:PRO:HA	1:C:814:LYS:HD2	1.92	0.51
1:A:599:THR:HB	1:A:608:VAL:HG23	1.92	0.51
1:C:599:THR:HB	1:C:608:VAL:HG23	1.92	0.51
1:C:809:PRO:HA	1:C:814:LYS:HD2	1.91	0.51
1:A:45:SER:OG	1:C:567:ARG:O	2.29	0.51
1:C:877:LEU:O	1:C:881:THR:OG1	2.28	0.51
1:A:299:THR:HG21	1:A:597:VAL:HG21	1.92	0.51
1:C:58:PHE:HD2	1:C:290:ASP:HB2	1.75	0.51
1:A:33:THR:OG1	1:A:219:GLY:O	2.27	0.51
1:A:330:PRO:HG3	1:A:579:PRO:HB2	1.91	0.51
1:A:452:LEU:HA	1:A:494:SER:HA	1.93	0.51

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:930:ALA:HA	1:A:933:LYS:HG2	1.91	0.51
1:C:34:ARG:NH1	1:C:219:GLY:O	2.40	0.51
1:C:564:GLN:HG2	1:C:577:ARG:HD2	1.91	0.51
1:A:770:ILE:HD12	1:A:1012:LEU:HD13	1.91	0.51
1:A:41:LYS:HD2	1:C:562:PHE:HE1	1.76	0.51
1:A:384:PRO:HA	1:A:387:LEU:HD23	1.92	0.51
1:B:358:ILE:HB	1:B:395:VAL:HG23	1.92	0.51
1:B:564:GLN:O	1:B:577:ARG:N	2.43	0.51
1:A:755:GLN:NE2	1:C:971:GLY:H	2.08	0.51
1:A:773:GLU:HG2	1:A:1019:ARG:HH21	1.74	0.51
1:C:709:ASN:OD1	1:C:709:ASN:N	2.41	0.51
1:C:112:SER:HB3	1:C:134:GLN:HG3	1.91	0.51
1:C:329:PHE:HZ	1:C:544:ASN:H	1.57	0.51
1:B:390:LEU:HD12	1:B:391:CYS:H	1.76	0.51
1:A:994:ASP:HA	1:A:997:ILE:HG12	1.91	0.51
1:C:34:ARG:HH22	1:C:220:PHE:HA	1.73	0.51
1:C:326:ILE:HD11	1:C:534:VAL:HG22	1.92	0.51
1:A:440:ASN:OD1	1:A:440:ASN:N	2.43	0.51
1:A:574:ASP:OD1	1:A:574:ASP:N	2.43	0.51
1:A:985:ASP:OD1	1:A:985:ASP:N	2.33	0.51
1:C:517:LEU:HG	1:C:518:LEU:HD22	1.92	0.51
1:A:1033:VAL:HG11	1:A:1053:PRO:HG3	1.92	0.51
1:B:1002:GLN:HA	1:B:1005:GLN:HG3	1.92	0.51
1:A:151:SER:HB2	1:A:153:MET:HE3	1.92	0.51
1:A:316:SER:OG	1:A:317:ASN:N	2.44	0.51
1:B:392:PHE:HD1	1:B:517:LEU:HD13	1.75	0.51
1:C:299:THR:HG21	1:C:597:VAL:HG11	1.92	0.51
1:A:533:LEU:HD21	1:A:585:LEU:HD11	1.91	0.51
1:B:390:LEU:HD12	1:B:391:CYS:H	1.75	0.51
1:B:815:ARG:NH1	1:B:867:ASP:OD1	2.43	0.51
1:A:914:ASN:OD1	1:A:914:ASN:N	2.43	0.51
1:A:560:LEU:HD23	1:A:562:PHE:H	1.76	0.51
1:A:334:ASN:ND2	1:A:360:ASN:O	2.41	0.51
1:A:35:GLY:HA3	1:A:56:LEU:HB3	1.93	0.51
1:A:120:VAL:HB	1:A:127:VAL:HB	1.92	0.51
1:C:273:ARG:NH1	1:C:290:ASP:OD2	2.44	0.51
1:A:393:THR:HA	1:A:522:ALA:HA	1.93	0.51
1:A:1115:ILE:HG22	1:A:1137:VAL:HG23	1.91	0.51
1:A:994:ASP:HA	1:A:997:ILE:HG12	1.91	0.51
1:B:326:ILE:HD11	1:B:534:VAL:HG22	1.91	0.51
1:A:384:PRO:HA	1:A:387:LEU:HD23	1.93	0.51

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:C:326:ILE:HD13	1:C:533:LEU:HD12	1.93	0.51
1:C:326:ILE:HD11	1:C:534:VAL:HG22	1.91	0.51
1:C:599:THR:HB	1:C:608:VAL:HG23	1.92	0.51
1:B:33:THR:OG1	1:B:219:GLY:O	2.22	0.51
1:B:105:ILE:HG23	1:B:239:GLN:HB2	1.92	0.51
1:C:599:THR:HB	1:C:608:VAL:HG23	1.93	0.51
1:C:650:LEU:HD13	1:C:653:ALA:HB3	1.92	0.51
1:A:1105:THR:HG21	1:A:1110:TYR:HA	1.93	0.51
1:A:971:GLY:HA3	1:A:995:ARG:HH22	1.76	0.51
1:C:897:PRO:HG2	1:C:900:MET:HB2	1.92	0.51
1:B:785:VAL:HG12	1:B:787:GLN:H	1.74	0.51
1:A:401:VAL:HG22	1:A:509:ARG:HG2	1.92	0.51
1:A:64:TRP:HE1	1:A:264:ALA:HB1	1.76	0.51
1:A:403:ARG:NE	1:A:406:GLU:OE1	2.40	0.51
1:C:966:LEU:HA	1:C:1000:ARG:HH12	1.75	0.51
1:B:726:ILE:HD13	1:B:1061:VAL:HG22	1.92	0.51
1:A:911:VAL:HG11	1:A:1067:TYR:HE1	1.76	0.51
1:A:33:THR:OG1	1:A:219:GLY:O	2.27	0.51
1:A:702:GLU:HA	1:B:788:ILE:HB	1.93	0.51
1:B:722:VAL:HG22	1:B:930:ALA:HB1	1.93	0.51
1:B:1013:ILE:HD13	1:C:1012:LEU:HG	1.92	0.51
1:A:317:ASN:HD21	1:A:592:PHE:HB2	1.75	0.51
1:C:326:ILE:HD13	1:C:533:LEU:HD12	1.93	0.51
1:A:130:VAL:HG21	1:A:231:ILE:HD11	1.93	0.51
1:A:881:THR:HG22	1:A:882:ILE:HD13	1.91	0.51
1:A:58:PHE:HD2	1:A:290:ASP:HB2	1.76	0.51
1:A:111:ASP:OD1	1:A:111:ASP:N	2.43	0.51
1:B:37:TYR:OH	1:B:54:LEU:O	2.24	0.51
1:A:742:ILE:HG21	1:A:997:ILE:HG22	1.92	0.51
1:B:905:ARG:HE	1:B:1050:MET:HB3	1.75	0.51
1:C:737:ASP:OD2	1:C:740:MET:N	2.42	0.51
1:A:370:ASN:N	1:A:370:ASN:HD22	2.08	0.51
1:A:501:TYR:O	1:A:506:GLN:NE2	2.43	0.51
1:C:1073:LYS:HE3	1:C:1075:PHE:HZ	1.75	0.51
1:B:117:LEU:HG	1:B:233:ILE:HD11	1.93	0.51
1:B:33:THR:OG1	1:B:219:GLY:O	2.29	0.51
1:B:577:ARG:HH21	1:B:584:ILE:HD11	1.75	0.51
1:C:931:ILE:HA	1:C:934:ILE:HG22	1.93	0.51
1:A:994:ASP:HA	1:A:997:ILE:HG12	1.91	0.51
1:C:317:ASN:HD21	1:C:592:PHE:HD1	1.59	0.51
1:B:317:ASN:ND2	1:C:737:ASP:OD1	2.44	0.51

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:C:769:GLY:HA2	1:C:772:VAL:HG12	1.93	0.51
1:A:52:GLN:OE1	1:A:274:THR:OG1	2.23	0.51
1:A:547:THR:O	1:B:978:ASN:ND2	2.43	0.51
1:C:102:ARG:HE	1:C:246:ILE:HD11	1.74	0.51
1:C:105:ILE:HB	1:C:239:GLN:HB3	1.92	0.51
1:A:34:ARG:HH22	1:A:220:PHE:HA	1.75	0.51
1:B:726:ILE:HG22	1:B:1061:VAL:HG22	1.92	0.51
1:C:206:LYS:HG2	1:C:224:GLU:H	1.76	0.51
1:C:34:ARG:HH22	1:C:220:PHE:HA	1.75	0.51
1:C:106:PHE:O	1:C:117:LEU:N	2.41	0.51
1:C:614:GLY:N	1:C:647:ALA:O	2.41	0.51
1:C:599:THR:HB	1:C:608:VAL:HG23	1.92	0.51
1:A:644:GLN:NE2	1:A:645:THR:O	2.43	0.51
1:B:614:GLY:N	1:B:647:ALA:O	2.35	0.51
1:C:409:GLN:HE22	1:C:417:ASN:HB3	1.76	0.51
1:B:1051:SER:OG	1:B:1064:HIS:ND1	2.37	0.51
1:C:809:PRO:HA	1:C:814:LYS:HD2	1.92	0.51
1:A:873:TYR:O	1:A:877:LEU:HB2	2.11	0.51
1:B:34:ARG:NH2	1:B:221:SER:H	2.08	0.51
1:B:729:VAL:HG21	1:B:781:VAL:HG11	1.92	0.51
1:A:326:ILE:HD11	1:A:534:VAL:HG22	1.93	0.51
1:A:599:THR:HB	1:A:608:VAL:HG23	1.91	0.51
1:A:864:LEU:HD12	1:A:865:LEU:HD13	1.92	0.51
1:A:997:ILE:O	1:A:1001:LEU:HB3	2.11	0.51
1:B:699:LEU:HG	1:C:873:TYR:HE1	1.76	0.51
1:C:729:VAL:HG12	1:C:1059:GLY:HA2	1.93	0.51
1:A:327:VAL:HA	1:A:542:ASN:HB3	1.92	0.51
1:B:105:ILE:HB	1:B:239:GLN:HB2	1.93	0.51
1:A:740:MET:HE1	1:A:857:GLY:HA3	1.92	0.51
1:B:37:TYR:OH	1:B:53:ASP:OD1	2.29	0.51
1:C:611:LEU:HB2	1:C:650:LEU:HD13	1.92	0.51
1:A:734:THR:HG21	1:A:959:LEU:HD21	1.92	0.51
1:A:806:LEU:HD22	1:A:878:LEU:HD21	1.92	0.51
1:B:729:VAL:HG21	1:B:781:VAL:HG11	1.91	0.51
1:B:506:GLN:HG3	1:B:507:PRO:HD2	1.92	0.51
1:C:383:SER:O	1:C:387:LEU:HB2	2.10	0.51
1:C:37:TYR:OH	1:C:53:ASP:OD1	2.27	0.51
1:A:904:TYR:OH	1:C:1093:GLY:O	2.16	0.51
1:B:405:ASP:N	1:B:405:ASP:OD1	2.43	0.51
1:A:87:ASN:OD1	1:A:87:ASN:N	2.43	0.51
1:A:120:VAL:HB	1:A:127:VAL:HB	1.92	0.51

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:C:37:TYR:OH	1:C:53:ASP:OD1	2.23	0.51
1:B:678:ILE:HD11	1:B:684:ALA:HB2	1.93	0.51
1:A:1116:THR:OG1	1:A:1118:ASP:OD1	2.29	0.51
1:A:1091:ARG:NH1	1:A:1118:ASP:O	2.36	0.51
1:C:393:THR:HA	1:C:522:ALA:HA	1.91	0.51
1:C:402:ILE:HD12	1:C:418:ILE:HG21	1.93	0.51
1:C:896:ILE:HD12	1:C:897:PRO:HD2	1.91	0.51
1:A:57:PRO:HG3	1:A:273:ARG:HG3	1.92	0.51
1:A:905:ARG:NH1	1:A:1049:LEU:O	2.39	0.51
1:A:802:PHE:HB3	1:A:806:LEU:HD23	1.93	0.51
1:B:37:TYR:OH	1:B:195:LYS:NZ	2.43	0.51
1:A:52:GLN:OE1	1:A:274:THR:OG1	2.19	0.51
1:B:712:ILE:HG12	1:B:1094:VAL:HG11	1.92	0.51
1:C:729:VAL:HG21	1:C:781:VAL:HG11	1.92	0.51
1:A:105:ILE:HG22	1:A:239:GLN:HB3	1.93	0.51
1:A:566:GLY:HA2	1:B:43:PHE:HB3	1.93	0.51
1:A:777:ASN:HA	1:A:780:GLU:HG3	1.92	0.51
1:C:877:LEU:O	1:C:881:THR:OG1	2.25	0.51
1:A:56:LEU:HD22	1:A:91:TYR:HD1	1.76	0.51
1:A:1139:ASP:HB3	1:A:1142:GLN:HG2	1.92	0.51
1:B:37:TYR:OH	1:B:195:LYS:NZ	2.41	0.51
1:C:440:ASN:OD1	1:C:440:ASN:N	2.42	0.51
1:A:592:PHE:HZ	1:B:857:GLY:HA2	1.76	0.51
1:B:462:LYS:NZ	1:B:465:GLU:OE1	2.35	0.51
1:C:1118:ASP:OD2	1:C:1118:ASP:N	2.39	0.51
1:B:326:ILE:HD11	1:B:534:VAL:HG22	1.92	0.51
1:B:1013:ILE:HD13	1:C:1012:LEU:HG	1.92	0.51
1:C:1008:VAL:O	1:C:1012:LEU:HB2	2.11	0.51
1:B:102:ARG:HH21	1:B:246:ILE:HG12	1.76	0.51
1:B:129:LYS:HZ3	1:B:169:GLU:HB2	1.75	0.51
1:C:358:ILE:HB	1:C:395:VAL:HG23	1.93	0.51
1:A:102:ARG:NH2	1:A:121:ASN:O	2.43	0.51
1:C:360:ASN:H	1:C:523:THR:HB	1.76	0.51
1:A:111:ASP:OD1	1:A:111:ASP:N	2.41	0.51
1:A:277:LEU:HD12	1:A:285:ILE:HD13	1.92	0.51
1:A:414:GLN:HA	1:A:414:GLN:NE2	2.26	0.51
1:B:360:ASN:H	1:B:523:THR:HB	1.75	0.51
1:C:317:ASN:HD21	1:C:592:PHE:HD2	1.58	0.51
1:C:908:GLY:O	1:C:1038:LYS:NZ	2.37	0.51
1:A:819:GLU:HA	1:A:822:LEU:HG	1.93	0.51
1:C:1052:PHE:HB2	1:C:1063:LEU:HG	1.92	0.51

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:B:332:ILE:HG12	1:B:524:VAL:HG13	1.91	0.51
1:C:1029:MET:HE2	1:C:1033:VAL:HG21	1.92	0.51
1:A:372:ALA:O	1:C:403:ARG:NH2	2.31	0.51
1:A:81:ASN:O	1:A:239:GLN:NE2	2.36	0.51
1:B:312:ILE:HD11	1:B:676:ILE:HB	1.92	0.51
1:C:326:ILE:HD11	1:C:534:VAL:HG22	1.92	0.51
1:C:777:ASN:HA	1:C:780:GLU:HG3	1.92	0.51
1:A:334:ASN:ND2	1:A:360:ASN:O	2.40	0.51
1:B:738:CYS:O	1:B:742:ILE:HB	2.11	0.51
1:B:1031:GLU:OE2	1:B:1039:ARG:NH2	2.42	0.51
1:A:229:LEU:HD12	1:A:230:PRO:HD2	1.93	0.51
1:B:777:ASN:HA	1:B:780:GLU:HG3	1.92	0.51
1:A:687:GLN:HG2	1:A:693:ILE:HD13	1.92	0.51
1:B:471:GLU:OE2	1:C:113:LYS:NZ	2.40	0.51
1:C:91:TYR:OH	1:C:191:GLU:OE1	2.28	0.51
1:A:762:GLN:HA	1:A:765:ARG:HE	1.76	0.51
1:A:134:GLN:NE2	1:A:162:SER:H	2.10	0.50
1:B:915:VAL:O	1:B:919:ASN:HB2	2.11	0.50
1:C:918:GLU:N	1:C:918:GLU:OE1	2.43	0.50
1:C:177:MET:SD	1:C:190:ARG:NH2	2.85	0.50
1:A:745:ASP:OD2	1:C:319:ARG:NH1	2.44	0.50
1:C:108:THR:O	1:C:237:ARG:NH2	2.44	0.50
1:C:298:GLU:HB3	1:C:315:THR:HG21	1.93	0.50
1:A:983:ARG:HG3	1:A:984:LEU:HD23	1.92	0.50
1:C:45:SER:HA	1:C:279:TYR:HB3	1.93	0.50
1:C:326:ILE:HD13	1:C:533:LEU:HD12	1.94	0.50
1:B:785:VAL:HG12	1:B:787:GLN:H	1.76	0.50
1:B:139:PRO:HG2	1:B:245:HIS:HE1	1.76	0.50
1:B:35:GLY:HA3	1:B:56:LEU:HB3	1.93	0.50
1:A:576:VAL:HG12	1:A:587:ILE:HD11	1.91	0.50
1:B:384:PRO:HA	1:B:387:LEU:HD23	1.92	0.50
1:C:777:ASN:HA	1:C:780:GLU:HG3	1.93	0.50
1:A:382:VAL:HG21	1:A:515:PHE:HZ	1.76	0.50
1:A:115:GLN:HB2	1:A:233:ILE:HG12	1.93	0.50
1:C:973:ILE:HG21	1:C:983:ARG:HH12	1.76	0.50
1:B:15:CYS:SG	1:B:137:ASN:N	2.83	0.50
1:B:465:GLU:OE1	1:C:234:ASN:ND2	2.32	0.50
1:C:916:LEU:HD23	1:C:917:TYR:HD2	1.76	0.50
1:B:401:VAL:HG22	1:B:509:ARG:HG2	1.93	0.50
1:B:738:CYS:O	1:B:742:ILE:HB	2.12	0.50
1:B:1115:ILE:HG22	1:B:1137:VAL:HG23	1.92	0.50

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:1031:GLU:HB3	1:A:1037:SER:HB2	1.93	0.50
1:B:472:ILE:HD13	1:B:484:LYS:HB2	1.91	0.50
1:C:350:VAL:HG22	1:C:400:PHE:HB2	1.94	0.50
1:A:1115:ILE:HG22	1:A:1137:VAL:HG23	1.93	0.50
1:C:33:THR:OG1	1:C:219:GLY:O	2.27	0.50
1:B:1031:GLU:HG3	1:B:1039:ARG:HH22	1.77	0.50
1:C:177:MET:HE3	1:C:178:ASP:H	1.76	0.50
1:C:440:ASN:N	1:C:440:ASN:OD1	2.40	0.50
1:B:276:LEU:HG	1:B:306:PHE:HE1	1.75	0.50
1:B:1139:ASP:HB3	1:B:1142:GLN:HG2	1.94	0.50
1:B:85:PRO:HA	1:B:237:ARG:HE	1.77	0.50
1:C:33:THR:OG1	1:C:219:GLY:O	2.22	0.50
1:C:398:ASP:OD2	1:C:423:TYR:OH	2.26	0.50
1:C:401:VAL:HG22	1:C:509:ARG:HG2	1.94	0.50
1:A:1139:ASP:HB3	1:A:1142:GLN:HG2	1.93	0.50
1:B:984:LEU:HB2	1:B:989:ALA:HB2	1.94	0.50
1:C:319:ARG:NH2	1:C:590:CYS:SG	2.84	0.50
1:A:1155:TYR:O	1:A:1159:HIS:ND1	2.33	0.50
1:B:55:PHE:HB2	1:B:273:ARG:HB2	1.93	0.50
1:A:729:VAL:HG21	1:A:781:VAL:HG11	1.93	0.50
1:C:222:ALA:HB2	1:C:285:ILE:HB	1.93	0.50
1:C:737:ASP:OD2	1:C:740:MET:N	2.42	0.50
1:A:222:ALA:HB2	1:A:285:ILE:HB	1.93	0.50
1:B:569:ILE:HG22	1:C:47:VAL:HG12	1.93	0.50
1:A:745:ASP:OD1	1:C:319:ARG:NH1	2.42	0.50
1:B:577:ARG:HH21	1:B:582:LEU:HD13	1.76	0.50
1:C:330:PRO:HD3	1:C:579:PRO:HB2	1.92	0.50
1:A:1012:LEU:HD12	1:C:1013:ILE:HD13	1.92	0.50
1:B:34:ARG:NH1	1:B:219:GLY:O	2.45	0.50
1:C:106:PHE:HB2	1:C:117:LEU:HB3	1.93	0.50
1:C:491:PRO:HG2	1:C:492:LEU:HD22	1.93	0.50
1:A:389:ASP:OD1	1:A:528:LYS:NZ	2.41	0.50
1:B:117:LEU:HB2	1:B:233:ILE:HD11	1.93	0.50
1:A:401:VAL:HG22	1:A:509:ARG:HG2	1.93	0.50
1:A:751:ASN:HA	1:A:754:LEU:HG	1.93	0.50
1:B:688:THR:HA	1:B:690:GLN:HG2	1.92	0.50
1:A:667:GLY:HA2	1:B:864:LEU:HA	1.93	0.50
1:C:185:ASN:ND2	1:C:211:ASN:OD1	2.45	0.50
1:C:767:LEU:HA	1:C:770:ILE:HG22	1.92	0.50
1:A:985:ASP:OD1	1:A:987:PRO:HD2	2.11	0.50
1:C:129:LYS:NZ	1:C:168:PHE:O	2.41	0.50

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:B:393:THR:OG1	1:B:394:ASN:OD1	2.30	0.50
1:C:58:PHE:HD2	1:C:290:ASP:HB2	1.75	0.50
1:A:1031:GLU:CD	1:C:1039:ARG:HG3	2.37	0.50
1:B:1115:ILE:HG22	1:B:1137:VAL:HG23	1.93	0.50
1:A:1033:VAL:HG11	1:A:1053:PRO:HG3	1.93	0.50
1:A:151:SER:HB2	1:A:153:MET:HE3	1.94	0.50
1:C:966:LEU:HA	1:C:1000:ARG:HH12	1.75	0.50
1:A:1116:THR:OG1	1:A:1118:ASP:OD1	2.29	0.50
1:C:809:PRO:HA	1:C:814:LYS:HD2	1.92	0.50
1:A:53:ASP:OD1	1:A:54:LEU:N	2.44	0.50
1:C:729:VAL:HG21	1:C:781:VAL:HG11	1.92	0.50
1:C:788:ILE:HG13	1:C:876:ALA:HB2	1.93	0.50
1:C:954:GLN:HE22	1:C:1014:ARG:HH11	1.59	0.50
1:A:409:GLN:HA	1:A:414:GLN:HG2	1.93	0.50
1:A:1115:ILE:HG22	1:A:1137:VAL:HG23	1.93	0.50
1:B:351:TYR:HE2	1:B:452:LEU:HB2	1.75	0.50
1:A:334:ASN:ND2	1:A:360:ASN:O	2.41	0.50
1:B:37:TYR:OH	1:B:53:ASP:OD1	2.29	0.50
1:C:809:PRO:HA	1:C:814:LYS:HD2	1.92	0.50
1:B:52:GLN:OE1	1:B:274:THR:OG1	2.27	0.50
1:A:111:ASP:OD1	1:A:111:ASP:N	2.43	0.50
1:A:403:ARG:NH2	1:B:372:ALA:O	2.36	0.50
1:C:467:ASP:OD2	1:C:467:ASP:N	2.42	0.50
1:A:722:VAL:HG22	1:A:930:ALA:HB1	1.91	0.50
1:A:886:TRP:HB3	1:A:1035:GLY:HA2	1.94	0.50
1:C:452:LEU:HD23	1:C:492:LEU:HB3	1.93	0.50
1:C:599:THR:HB	1:C:608:VAL:HG23	1.94	0.50
1:A:14:GLN:O	1:A:158:ARG:NE	2.43	0.50
1:A:1116:THR:OG1	1:A:1118:ASP:OD1	2.29	0.50
1:B:442:ASP:OD2	1:B:509:ARG:NE	2.41	0.50
1:B:102:ARG:HH12	1:B:123:ALA:HB2	1.76	0.50
1:C:39:PRO:HG3	1:C:51:THR:HG21	1.92	0.50
1:A:1116:THR:OG1	1:A:1118:ASP:OD1	2.29	0.50
1:B:34:ARG:HH22	1:B:220:PHE:HA	1.76	0.50
1:B:1030:SER:HA	1:B:1034:LEU:HD23	1.94	0.50
1:A:108:THR:O	1:A:237:ARG:NH1	2.45	0.50
1:B:44:ARG:HH11	1:B:49:HIS:HB3	1.77	0.50
1:B:950:ASP:OD1	1:B:950:ASP:N	2.44	0.50
1:C:393:THR:HB	1:C:520:ALA:HB3	1.93	0.50
1:A:111:ASP:OD1	1:A:111:ASP:N	2.43	0.50
1:C:619:GLU:N	1:C:619:GLU:OE1	2.45	0.50

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:87:ASN:OD1	1:A:87:ASN:N	2.44	0.50
1:B:403:ARG:NH2	1:C:372:ALA:O	2.35	0.50
1:A:108:THR:O	1:A:237:ARG:NH2	2.45	0.50
1:A:574:ASP:OD1	1:A:574:ASP:N	2.45	0.50
1:C:338:PHE:HA	1:C:341:VAL:HG12	1.94	0.50
1:C:1039:ARG:HD2	1:C:1042:PHE:HB2	1.94	0.50
1:A:1116:THR:OG1	1:A:1118:ASP:OD1	2.29	0.50
1:A:726:ILE:HG22	1:A:1061:VAL:HG22	1.93	0.50
1:A:994:ASP:HA	1:A:997:ILE:HG12	1.91	0.50
1:B:440:ASN:OD1	1:B:440:ASN:N	2.42	0.50
1:B:236:THR:O	1:B:237:ARG:NE	2.44	0.50
1:B:33:THR:OG1	1:B:219:GLY:O	2.29	0.50
1:C:330:PRO:HD3	1:C:579:PRO:HB2	1.94	0.50
1:A:503:VAL:HA	1:A:506:GLN:HG3	1.93	0.50
1:C:393:THR:HA	1:C:522:ALA:HA	1.93	0.50
1:A:984:LEU:HB2	1:A:989:ALA:HB2	1.94	0.50
1:B:547:THR:O	1:C:978:ASN:ND2	2.44	0.50
1:C:454:ARG:NH1	1:C:456:PHE:O	2.44	0.50
1:C:330:PRO:HD3	1:C:579:PRO:HB2	1.93	0.50
1:C:726:ILE:HG13	1:C:948:LEU:HD23	1.93	0.50
1:C:18:PHE:HB2	1:C:21:ARG:HB2	1.92	0.50
1:A:951:VAL:HA	1:A:954:GLN:HG3	1.93	0.50
1:B:390:LEU:HD12	1:B:391:CYS:H	1.76	0.50
1:C:392:PHE:HB2	1:C:524:VAL:HG23	1.92	0.50
1:B:52:GLN:OE1	1:B:274:THR:OG1	2.22	0.50
1:B:389:ASP:OD1	1:B:528:LYS:NZ	2.43	0.50
1:C:195:LYS:HG2	1:C:202:LYS:HB2	1.94	0.50
1:B:194:PHE:HB3	1:B:201:PHE:HE1	1.76	0.50
1:C:598:ILE:HD11	1:C:678:ILE:HD13	1.92	0.50
1:A:113:LYS:NZ	1:C:471:GLU:OE2	2.45	0.50
1:B:902:MET:HB3	1:B:916:LEU:HD11	1.93	0.50
1:B:37:TYR:OH	1:B:53:ASP:OD1	2.28	0.50
1:B:398:ASP:OD2	1:B:423:TYR:OH	2.23	0.50
1:B:613:GLN:HA	1:B:648:GLY:HA3	1.94	0.50
1:B:1054:GLN:N	1:B:1061:VAL:O	2.45	0.50
1:C:1091:ARG:HH12	1:C:1121:PHE:HB3	1.75	0.50
1:A:89:GLY:HA3	1:A:270:LEU:HD12	1.94	0.50
1:A:204:TYR:HD1	1:A:225:PRO:HA	1.76	0.50
1:A:566:GLY:HA2	1:B:43:PHE:HB3	1.94	0.50
1:B:303:LEU:HD23	1:B:308:VAL:HG22	1.93	0.50
1:A:777:ASN:HA	1:A:780:GLU:HG3	1.92	0.50

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:C:290:ASP:HB3	1:C:293:LEU:HB2	1.93	0.50
1:C:382:VAL:HG21	1:C:515:PHE:HZ	1.77	0.50
1:A:299:THR:HG21	1:A:597:VAL:HG21	1.93	0.50
1:A:931:ILE:HA	1:A:934:ILE:HG22	1.94	0.50
1:A:1033:VAL:HG11	1:A:1053:PRO:HG3	1.93	0.50
1:C:912:THR:H	1:C:1106:GLN:HE22	1.59	0.50
1:A:1051:SER:OG	1:A:1064:HIS:ND1	2.38	0.50
1:C:326:ILE:HD11	1:C:534:VAL:HG22	1.93	0.50
1:C:58:PHE:HB2	1:C:293:LEU:HD22	1.92	0.50
1:C:741:TYR:OH	1:C:1003:SER:OG	2.28	0.50
1:A:931:ILE:HA	1:A:934:ILE:HG22	1.94	0.50
1:B:105:ILE:HG13	1:B:241:LEU:HD11	1.92	0.50
1:C:871:ALA:HA	1:C:874:THR:HG22	1.93	0.50
1:A:177:MET:SD	1:A:190:ARG:NH2	2.85	0.50
1:A:1005:GLN:HE21	1:C:1006:THR:HG21	1.77	0.50
1:B:650:LEU:HD13	1:B:653:ALA:HB3	1.93	0.50
1:C:611:LEU:HB2	1:C:650:LEU:HD13	1.92	0.50
1:C:360:ASN:H	1:C:523:THR:HB	1.75	0.50
1:C:905:ARG:HH12	1:C:1036:GLN:HB2	1.76	0.50
1:A:52:GLN:OE1	1:A:274:THR:OG1	2.26	0.50
1:A:729:VAL:HG21	1:A:781:VAL:HG11	1.94	0.50
1:C:902:MET:HB3	1:C:916:LEU:HD11	1.93	0.50
1:B:951:VAL:HA	1:B:954:GLN:HG3	1.94	0.50
1:C:212:LEU:HD22	1:C:217:PRO:HD3	1.92	0.50
1:C:214:ARG:HD2	1:C:215:GLY:H	1.75	0.50
1:A:111:ASP:OD2	1:A:111:ASP:N	2.44	0.50
1:A:206:LYS:HG2	1:A:224:GLU:H	1.77	0.50
1:B:699:LEU:HG	1:C:873:TYR:HE1	1.77	0.50
1:B:826:VAL:HB	1:B:1057:PRO:HG2	1.93	0.50
1:C:809:PRO:HA	1:C:814:LYS:HD2	1.92	0.50
1:C:106:PHE:O	1:C:117:LEU:N	2.37	0.50
1:A:533:LEU:HD21	1:A:585:LEU:HD11	1.92	0.50
1:A:353:TRP:HH2	1:A:464:PHE:HA	1.76	0.50
1:A:547:THR:O	1:B:978:ASN:ND2	2.45	0.50
1:A:777:ASN:O	1:A:781:VAL:HG12	2.11	0.50
1:A:360:ASN:HD22	1:A:360:ASN:C	2.16	0.50
1:B:1086:LYS:HA	1:B:1125:ASN:HA	1.93	0.50
1:C:38:TYR:HE2	1:C:224:GLU:HG3	1.77	0.50
1:C:777:ASN:HA	1:C:780:GLU:HG3	1.93	0.50
1:C:389:ASP:OD1	1:C:528:LYS:NZ	2.43	0.50
1:B:37:TYR:OH	1:B:53:ASP:OD1	2.27	0.50

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:C:393:THR:OG1	1:C:394:ASN:N	2.44	0.50
1:C:1051:SER:OG	1:C:1064:HIS:ND1	2.40	0.50
1:B:34:ARG:NH2	1:B:221:SER:H	2.10	0.50
1:B:569:ILE:HA	1:C:47:VAL:HG12	1.92	0.50
1:B:598:ILE:HB	1:B:609:ALA:HB3	1.92	0.50
1:B:664:ILE:HD12	1:B:665:PRO:HD2	1.93	0.50
1:C:374:PHE:HB3	1:C:436:TRP:HD1	1.77	0.50
1:A:398:ASP:OD2	1:A:423:TYR:OH	2.23	0.50
1:B:978:ASN:HA	1:B:981:LEU:HG	1.94	0.50
1:C:401:VAL:HG22	1:C:509:ARG:HG2	1.94	0.50
1:C:1033:VAL:HG21	1:C:1053:PRO:HG3	1.93	0.50
1:C:598:ILE:HB	1:C:609:ALA:HB3	1.94	0.50
1:A:759:PHE:HA	1:A:762:GLN:HE21	1.77	0.50
1:B:22:THR:OG1	1:B:78:ARG:NH1	2.45	0.50
1:B:978:ASN:HA	1:B:981:LEU:HG	1.93	0.50
1:A:722:VAL:HG21	1:A:931:ILE:HD13	1.94	0.50
1:A:47:VAL:HG12	1:C:569:ILE:HA	1.94	0.50
1:A:983:ARG:HH11	1:C:517:LEU:HD11	1.77	0.50
1:B:598:ILE:HB	1:B:609:ALA:HB3	1.94	0.50
1:B:712:ILE:HG12	1:B:1094:VAL:HG11	1.94	0.50
1:A:809:PRO:O	1:A:814:LYS:NZ	2.39	0.50
1:A:1116:THR:OG1	1:A:1118:ASP:OD1	2.30	0.50
1:B:34:ARG:NH2	1:B:221:SER:H	2.10	0.50
1:A:398:ASP:OD2	1:A:423:TYR:OH	2.28	0.50
1:B:431:GLY:HA2	1:B:515:PHE:CZ	2.47	0.50
1:A:278:LYS:HB2	1:A:306:PHE:CE1	2.46	0.50
1:B:679:GLY:HA2	1:C:864:LEU:HD12	1.94	0.50
1:B:962:LEU:HD22	1:B:1007:TYR:HE2	1.75	0.50
1:C:552:LEU:HB2	1:C:585:LEU:HD22	1.94	0.50
1:A:80:ALA:O	1:A:245:HIS:NE2	2.45	0.50
1:C:782:PHE:HE1	1:C:870:ILE:HD11	1.77	0.50
1:A:611:LEU:HD22	1:A:678:ILE:HD11	1.94	0.50
1:A:1106:GLN:NE2	1:A:1111:GLU:OE2	2.45	0.50
1:B:195:LYS:HD3	1:B:204:TYR:HE1	1.77	0.50
1:B:185:ASN:HB2	1:B:213:VAL:HA	1.94	0.50
1:C:326:ILE:HD11	1:C:534:VAL:HG22	1.94	0.50
1:C:426:PRO:HA	1:C:463:PRO:HB3	1.92	0.50
1:C:708:SER:HB3	1:C:711:SER:HB3	1.94	0.50
1:B:326:ILE:HD11	1:B:534:VAL:HG22	1.92	0.50
1:B:119:ILE:HG12	1:B:128:ILE:HG12	1.93	0.50
1:C:599:THR:HB	1:C:608:VAL:HG23	1.94	0.50

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:B:877:LEU:HD21	1:B:1029:MET:HE1	1.94	0.50
1:C:972:ALA:HA	1:C:995:ARG:HH22	1.77	0.50
1:C:330:PRO:HD3	1:C:579:PRO:HB2	1.92	0.50
1:B:431:GLY:HA2	1:B:515:PHE:CZ	2.47	0.50
1:A:204:TYR:HB3	1:A:223:LEU:HG	1.94	0.50
1:A:561:PRO:HA	1:A:577:ARG:HH12	1.77	0.50
1:A:344:ALA:O	1:A:509:ARG:NH1	2.45	0.49
1:A:702:GLU:HA	1:B:788:ILE:HB	1.94	0.49
1:B:14:GLN:N	1:B:255:SER:HG	2.09	0.49
1:B:332:ILE:HA	1:B:524:VAL:HG22	1.94	0.49
1:C:642:VAL:HG13	1:C:651:ILE:HG22	1.93	0.49
1:C:276:LEU:HG	1:C:306:PHE:HE1	1.76	0.49
1:A:111:ASP:OD1	1:A:111:ASP:N	2.45	0.49
1:A:994:ASP:HA	1:A:997:ILE:HG12	1.94	0.49
1:B:950:ASP:OD1	1:B:950:ASP:N	2.42	0.49
1:A:392:PHE:HB2	1:A:524:VAL:HG23	1.94	0.49
1:B:1116:THR:OG1	1:B:1118:ASP:OD1	2.30	0.49
1:A:302:THR:O	1:A:304:LYS:NZ	2.44	0.49
1:A:401:VAL:HG22	1:A:509:ARG:HG2	1.93	0.49
1:A:994:ASP:HA	1:A:997:ILE:HG12	1.93	0.49
1:A:1115:ILE:HG22	1:A:1137:VAL:HG23	1.94	0.49
1:C:37:TYR:OH	1:C:53:ASP:OD1	2.28	0.49
1:C:730:SER:OG	1:C:731:MET:N	2.45	0.49
1:A:566:GLY:HA2	1:B:43:PHE:HB3	1.93	0.49
1:C:483:VAL:HG12	1:C:484:LYS:HG3	1.94	0.49
1:C:119:ILE:HG22	1:C:128:ILE:HD13	1.93	0.49
1:C:966:LEU:HA	1:C:1000:ARG:HH12	1.77	0.49
1:B:394:ASN:OD1	1:B:394:ASN:N	2.45	0.49
1:A:1115:ILE:HG22	1:A:1137:VAL:HG23	1.94	0.49
1:C:561:PRO:O	1:C:577:ARG:NH1	2.42	0.49
1:C:53:ASP:HB3	1:C:55:PHE:HE1	1.77	0.49
1:A:81:ASN:ND2	1:A:138:ASP:OD1	2.45	0.49
1:B:950:ASP:OD1	1:B:950:ASP:N	2.44	0.49
1:C:729:VAL:HG21	1:C:781:VAL:HG11	1.92	0.49
1:C:552:LEU:HD22	1:C:585:LEU:HD13	1.94	0.49
1:B:403:ARG:NH2	1:C:372:ALA:O	2.37	0.49
1:A:386:LYS:NZ	1:B:981:LEU:O	2.35	0.49
1:A:1010:GLN:HE21	1:A:1010:GLN:C	2.21	0.49
1:A:726:ILE:HG22	1:A:1061:VAL:HG22	1.93	0.49
1:A:483:VAL:HG12	1:A:484:LYS:HD3	1.94	0.49
1:B:985:ASP:OD2	1:B:985:ASP:N	2.45	0.49

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:C:752:LEU:HD23	1:C:993:ILE:HG22	1.94	0.49
1:C:440:ASN:N	1:C:440:ASN:OD1	2.41	0.49
1:A:561:PRO:O	1:A:577:ARG:NH1	2.45	0.49
1:A:143:VAL:HG22	1:A:154:GLU:HA	1.94	0.49
1:B:978:ASN:HA	1:B:981:LEU:HG	1.93	0.49
1:B:726:ILE:HD13	1:B:1061:VAL:HG22	1.93	0.49
1:C:809:PRO:HA	1:C:814:LYS:HD2	1.93	0.49
1:C:87:ASN:OD1	1:C:87:ASN:N	2.43	0.49
1:C:650:LEU:HD13	1:C:653:ALA:HB3	1.94	0.49
1:B:332:ILE:HG13	1:B:362:VAL:HG22	1.94	0.49
1:C:1028:LYS:NZ	1:C:1042:PHE:O	2.36	0.49
1:A:574:ASP:OD1	1:A:574:ASP:N	2.45	0.49
1:C:825:LYS:HD2	1:C:942:ALA:HA	1.93	0.49
1:A:189:LEU:HD22	1:A:210:ILE:HD13	1.93	0.49
1:C:484:LYS:HE3	1:C:489:TYR:HA	1.93	0.49
1:A:726:ILE:HD13	1:A:1061:VAL:HG22	1.94	0.49
1:B:276:LEU:HD11	1:B:304:LYS:HE2	1.95	0.49
1:C:52:GLN:OE1	1:C:274:THR:OG1	2.22	0.49
1:A:111:ASP:OD1	1:A:111:ASP:N	2.43	0.49
1:B:353:TRP:HE1	1:B:466:ARG:HB2	1.77	0.49
1:B:403:ARG:HB2	1:B:495:TYR:CE2	2.46	0.49
1:A:765:ARG:HH22	1:C:957:GLN:HE21	1.59	0.49
1:B:517:LEU:HD11	1:C:983:ARG:HH11	1.76	0.49
1:B:596:SER:HB2	1:B:611:LEU:HB3	1.94	0.49
1:B:719:THR:HG23	1:B:1070:ALA:HB2	1.94	0.49
1:B:1116:THR:OG1	1:B:1118:ASP:OD1	2.29	0.49
1:C:147:LYS:HB3	1:C:153:MET:HE2	1.93	0.49
1:C:599:THR:HB	1:C:608:VAL:HG23	1.95	0.49
1:A:735:SER:O	1:A:859:THR:OG1	2.26	0.49
1:B:326:ILE:HD11	1:B:534:VAL:HG22	1.92	0.49
1:B:389:ASP:OD1	1:B:389:ASP:N	2.40	0.49
1:B:433:VAL:HA	1:B:512:VAL:HG12	1.94	0.49
1:A:1115:ILE:HG22	1:A:1137:VAL:HG23	1.93	0.49
1:C:491:PRO:HG2	1:C:492:LEU:HD22	1.93	0.49
1:B:105:ILE:HG13	1:B:241:LEU:HD11	1.95	0.49
1:B:358:ILE:HB	1:B:395:VAL:HG23	1.94	0.49
1:B:699:LEU:HG	1:C:873:TYR:HE1	1.77	0.49
1:A:22:THR:O	1:A:78:ARG:NH1	2.45	0.49
1:B:64:TRP:HE1	1:B:264:ALA:HB1	1.77	0.49
1:B:1116:THR:OG1	1:B:1118:ASP:OD2	2.30	0.49
1:A:791:THR:HG21	1:A:806:LEU:HD11	1.94	0.49

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:B:1046:GLY:HA2	1:C:890:ALA:HA	1.95	0.49
1:A:563:GLN:HG2	1:B:43:PHE:HD1	1.76	0.49
1:C:118:LEU:HD11	1:C:135:PHE:HZ	1.77	0.49
1:C:752:LEU:HD11	1:C:994:ASP:HB3	1.94	0.49
1:B:1051:SER:OG	1:B:1064:HIS:ND1	2.42	0.49
1:B:983:ARG:HG3	1:B:984:LEU:HD22	1.94	0.49
1:C:756:TYR:HB3	1:C:759:PHE:CZ	2.47	0.49
1:B:52:GLN:OE1	1:B:274:THR:OG1	2.25	0.49
1:B:598:ILE:HD11	1:B:611:LEU:HD23	1.95	0.49
1:C:687:GLN:HG2	1:C:693:ILE:HD13	1.94	0.49
1:C:1086:LYS:HE3	1:C:1122:VAL:HG11	1.93	0.49
1:C:105:ILE:HG23	1:C:239:GLN:HB2	1.93	0.49
1:C:546:LEU:HD21	1:C:573:THR:HG21	1.93	0.49
1:C:738:CYS:O	1:C:742:ILE:HB	2.12	0.49
1:A:440:ASN:OD1	1:A:440:ASN:N	2.45	0.49
1:B:1116:THR:OG1	1:B:1118:ASP:OD1	2.29	0.49
1:C:330:PRO:HD3	1:C:579:PRO:HB2	1.94	0.49
1:B:15:CYS:SG	1:B:137:ASN:N	2.83	0.49
1:A:1116:THR:HG22	1:A:1138:TYR:HD1	1.77	0.49
1:B:1030:SER:HA	1:B:1034:LEU:HD23	1.94	0.49
1:B:1107:ARG:NH2	1:C:904:TYR:OH	2.39	0.49
1:C:884:SER:OG	1:C:887:THR:OG1	2.24	0.49
1:B:902:MET:HE1	1:B:1050:MET:HE2	1.94	0.49
1:B:777:ASN:HD21	1:B:1019:ARG:HD2	1.77	0.49
1:C:866:THR:OG1	1:C:867:ASP:OD2	2.30	0.49
1:A:222:ALA:HB2	1:A:285:ILE:HB	1.94	0.49
1:C:191:GLU:HG2	1:C:223:LEU:HD21	1.92	0.49
1:C:394:ASN:OD1	1:C:394:ASN:N	2.45	0.49
1:C:742:ILE:HG21	1:C:753:LEU:HG	1.93	0.49
1:B:521:PRO:HD3	1:C:41:LYS:HE2	1.95	0.49
1:C:1086:LYS:HE3	1:C:1122:VAL:HG11	1.93	0.49
1:C:58:PHE:HD2	1:C:290:ASP:HB2	1.78	0.49
1:C:599:THR:HB	1:C:608:VAL:HG23	1.94	0.49
1:A:1116:THR:OG1	1:A:1118:ASP:OD1	2.30	0.49
1:B:332:ILE:HG13	1:B:362:VAL:HG22	1.94	0.49
1:C:1086:LYS:HE3	1:C:1122:VAL:HG11	1.94	0.49
1:A:37:TYR:OH	1:A:53:ASP:OD1	2.29	0.49
1:A:737:ASP:OD2	1:C:317:ASN:ND2	2.46	0.49
1:A:353:TRP:HH2	1:A:464:PHE:HA	1.78	0.49
1:A:1081:ILE:HG23	1:A:1088:HIS:HB2	1.94	0.49
1:B:972:ALA:HB2	1:B:996:LEU:HD21	1.94	0.49

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:C:58:PHE:CD2	1:C:290:ASP:HB2	2.47	0.49
1:C:777:ASN:HA	1:C:780:GLU:HG3	1.94	0.49
1:B:193:VAL:HB	1:B:204:TYR:HB2	1.93	0.49
1:C:866:THR:H	1:C:869:MET:HE2	1.76	0.49
1:A:642:VAL:HG22	1:A:651:ILE:HG22	1.95	0.49
1:B:106:PHE:HE2	1:B:119:ILE:HD12	1.77	0.49
1:A:791:THR:HG21	1:A:806:LEU:HD11	1.94	0.49
1:C:1033:VAL:HG21	1:C:1053:PRO:HG3	1.93	0.49
1:A:39:PRO:HG3	1:A:51:THR:HG21	1.94	0.49
1:A:1031:GLU:HB3	1:A:1037:SER:HB2	1.95	0.49
1:A:755:GLN:HE22	1:C:971:GLY:H	1.60	0.49
1:B:276:LEU:HG	1:B:306:PHE:HE1	1.78	0.49
1:B:1051:SER:OG	1:B:1064:HIS:ND1	2.38	0.49
1:C:774:GLN:HA	1:C:774:GLN:HE21	1.78	0.49
1:A:222:ALA:HB2	1:A:285:ILE:HB	1.93	0.49
1:B:731:MET:H	1:B:774:GLN:HE22	1.58	0.49
1:B:966:LEU:HA	1:B:1000:ARG:HH12	1.77	0.49
1:C:299:THR:HG22	1:C:597:VAL:HG21	1.94	0.49
1:A:99:ASN:OD1	1:A:190:ARG:NH2	2.41	0.49
1:B:401:VAL:HG22	1:B:509:ARG:HG2	1.95	0.49
1:C:106:PHE:HB3	1:C:235:ILE:HD12	1.95	0.49
1:A:809:PRO:O	1:A:814:LYS:NZ	2.41	0.49
1:B:1051:SER:OG	1:B:1064:HIS:ND1	2.42	0.49
1:C:33:THR:OG1	1:C:219:GLY:O	2.26	0.49
1:C:389:ASP:OD1	1:C:528:LYS:NZ	2.37	0.49
1:A:983:ARG:O	1:C:383:SER:N	2.37	0.49
1:B:326:ILE:HD11	1:B:534:VAL:HG22	1.94	0.49
1:C:619:GLU:N	1:C:619:GLU:OE1	2.46	0.49
1:A:767:LEU:HA	1:A:770:ILE:HG22	1.94	0.49
1:C:564:GLN:OE1	1:C:577:ARG:NH2	2.46	0.49
1:C:733:LYS:HE3	1:C:771:ALA:HB1	1.94	0.49
1:A:906:PHE:HB3	1:A:911:VAL:HB	1.94	0.49
1:B:393:THR:HA	1:B:522:ALA:HA	1.94	0.49
1:B:619:GLU:N	1:B:619:GLU:OE1	2.46	0.49
1:B:731:MET:H	1:B:774:GLN:NE2	2.11	0.49
1:A:517:LEU:HD11	1:B:983:ARG:HD2	1.94	0.49
1:A:687:GLN:HG2	1:A:693:ILE:HD13	1.95	0.49
1:A:517:LEU:HD11	1:B:983:ARG:HH11	1.78	0.49
1:C:1116:THR:OG1	1:C:1118:ASP:OD1	2.29	0.49
1:A:386:LYS:NZ	1:B:981:LEU:O	2.46	0.49
1:A:722:VAL:HG22	1:A:930:ALA:HB1	1.93	0.49

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:B:576:VAL:HG12	1:B:587:ILE:HD11	1.94	0.49
1:B:775:ASP:HB3	1:B:864:LEU:HD13	1.94	0.49
1:C:822:LEU:HD13	1:C:1056:ALA:HB2	1.95	0.49
1:A:91:TYR:OH	1:A:191:GLU:OE2	2.29	0.49
1:A:547:THR:O	1:B:978:ASN:ND2	2.46	0.49
1:B:1079:PRO:HB3	1:C:900:MET:HE1	1.94	0.49
1:C:946:GLY:HA2	1:C:949:GLN:HB2	1.95	0.49
1:B:950:ASP:OD1	1:B:950:ASP:N	2.45	0.49
1:B:58:PHE:HD2	1:B:290:ASP:HB2	1.77	0.49
1:C:728:PRO:HD3	1:C:947:LYS:HG3	1.94	0.49
1:C:756:TYR:HB3	1:C:759:PHE:HD1	1.78	0.49
1:A:315:THR:OG1	1:A:316:SER:N	2.46	0.49
1:A:767:LEU:HD13	1:A:770:ILE:HD11	1.94	0.49
1:B:731:MET:H	1:B:774:GLN:NE2	2.11	0.49
1:C:33:THR:OG1	1:C:219:GLY:O	2.28	0.49
1:B:358:ILE:HB	1:B:395:VAL:HG23	1.94	0.49
1:A:58:PHE:HD2	1:A:290:ASP:HB2	1.77	0.49
1:B:34:ARG:NH2	1:B:221:SER:H	2.11	0.49
1:B:273:ARG:HE	1:B:292:ALA:HB3	1.78	0.49
1:A:729:VAL:HG21	1:A:781:VAL:HG11	1.94	0.49
1:A:36:VAL:HG13	1:A:222:ALA:HA	1.95	0.49
1:A:501:TYR:O	1:A:506:GLN:NE2	2.44	0.49
1:A:382:VAL:HG21	1:A:515:PHE:HZ	1.78	0.49
1:B:21:ARG:HD3	1:B:79:PHE:HB3	1.93	0.49
1:C:731:MET:H	1:C:774:GLN:NE2	2.10	0.49
1:A:785:VAL:HG12	1:A:787:GLN:H	1.77	0.49
1:C:777:ASN:O	1:C:781:VAL:HG12	2.12	0.49
1:A:762:GLN:HA	1:A:765:ARG:HE	1.77	0.49
1:B:52:GLN:OE1	1:B:274:THR:OG1	2.24	0.49
1:A:390:LEU:HD21	1:A:517:LEU:HD12	1.95	0.49
1:A:398:ASP:OD2	1:A:423:TYR:OH	2.25	0.49
1:B:390:LEU:HD12	1:B:391:CYS:H	1.77	0.49
1:A:598:ILE:HD11	1:A:678:ILE:HD13	1.95	0.49
1:B:722:VAL:HG22	1:B:930:ALA:HB1	1.95	0.49
1:B:185:ASN:HB2	1:B:213:VAL:HA	1.95	0.49
1:A:34:ARG:NH2	1:A:221:SER:H	2.11	0.49
1:A:555:SER:HB2	1:A:586:ASP:HB3	1.93	0.49
1:B:37:TYR:OH	1:B:53:ASP:OD1	2.25	0.49
1:B:569:ILE:HG12	1:C:47:VAL:HG22	1.95	0.49
1:C:34:ARG:HH22	1:C:220:PHE:HA	1.77	0.49
1:C:276:LEU:HD11	1:C:304:LYS:HA	1.95	0.49

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:B:389:ASP:OD1	1:B:389:ASP:N	2.42	0.49
1:B:642:VAL:HG22	1:B:651:ILE:HD12	1.94	0.49
1:A:777:ASN:HA	1:A:780:GLU:HG3	1.94	0.49
1:C:102:ARG:HD3	1:C:121:ASN:HB3	1.94	0.49
1:A:490:PHE:HD1	1:A:491:PRO:HD2	1.78	0.49
1:A:735:SER:HA	1:A:767:LEU:HD23	1.94	0.49
1:B:102:ARG:NH1	1:B:154:GLU:OE2	2.38	0.49
1:B:48:LEU:HD12	1:B:276:LEU:HD11	1.95	0.49
1:C:52:GLN:OE1	1:C:274:THR:OG1	2.21	0.49
1:A:599:THR:HB	1:A:608:VAL:HG23	1.94	0.49
1:A:994:ASP:HA	1:A:997:ILE:HG12	1.94	0.49
1:B:40:ASP:OD1	1:B:41:LYS:N	2.43	0.49
1:C:370:ASN:HD22	1:C:384:PRO:HB2	1.77	0.49
1:B:777:ASN:HA	1:B:780:GLU:HG3	1.95	0.49
1:C:970:PHE:O	1:C:995:ARG:NH1	2.46	0.49
1:A:802:PHE:HB3	1:A:806:LEU:HD23	1.94	0.49
1:C:229:LEU:HD12	1:C:230:PRO:HD2	1.94	0.49
1:A:501:TYR:HB3	1:A:505:TYR:HB3	1.93	0.49
1:A:222:ALA:HB2	1:A:285:ILE:HB	1.94	0.49
1:A:1035:GLY:HA3	1:C:1040:VAL:HG11	1.95	0.49
1:C:751:ASN:HA	1:C:754:LEU:HG	1.95	0.49
1:A:330:PRO:HG3	1:A:579:PRO:HB2	1.94	0.49
1:A:886:TRP:HB3	1:A:1035:GLY:HA2	1.95	0.49
1:B:719:THR:HG23	1:B:1070:ALA:HB2	1.95	0.49
1:B:884:SER:OG	1:B:887:THR:OG1	2.25	0.49
1:C:52:GLN:OE1	1:C:274:THR:OG1	2.24	0.49
1:A:390:LEU:HD12	1:A:391:CYS:H	1.77	0.49
1:B:719:THR:HG23	1:B:1070:ALA:HB2	1.95	0.49
1:A:93:ALA:HB3	1:A:266:TYR:HB2	1.95	0.49
1:A:806:LEU:HD22	1:A:878:LEU:HD21	1.94	0.49
1:B:457:ARG:NE	1:B:459:SER:O	2.46	0.49
1:B:777:ASN:HA	1:B:780:GLU:HG3	1.94	0.49
1:C:785:VAL:HG12	1:C:787:GLN:H	1.78	0.49
1:B:719:THR:HG23	1:B:1068:VAL:HB	1.94	0.49
1:B:28:TYR:HB3	1:B:61:ASN:HB3	1.95	0.49
1:B:92:PHE:HE2	1:B:265:TYR:HB2	1.78	0.49
1:A:560:LEU:HD12	1:A:562:PHE:HE1	1.78	0.49
1:B:204:TYR:CZ	1:B:225:PRO:HG3	2.48	0.49
1:B:326:ILE:HD11	1:B:534:VAL:HG22	1.94	0.49
1:B:574:ASP:OD1	1:B:574:ASP:N	2.46	0.49
1:B:472:ILE:HD13	1:B:484:LYS:HB2	1.94	0.49

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:C:825:LYS:HD3	1:C:942:ALA:HA	1.95	0.49
1:A:701:VAL:HG13	1:B:787:GLN:HG2	1.95	0.49
1:A:1050:MET:HE2	1:A:1050:MET:HB3	1.69	0.49
1:B:353:TRP:HH2	1:B:464:PHE:HA	1.78	0.49
1:B:452:LEU:HA	1:B:494:SER:HA	1.95	0.49
1:C:725:GLU:OE2	1:C:1064:HIS:NE2	2.43	0.49
1:A:884:SER:OG	1:A:887:THR:OG1	2.25	0.49
1:C:80:ALA:O	1:C:245:HIS:NE2	2.46	0.49
1:C:770:ILE:HA	1:C:773:GLU:HG3	1.95	0.49
1:B:34:ARG:HH22	1:B:220:PHE:HA	1.78	0.49
1:A:498:GLN:HB2	1:A:501:TYR:HE2	1.78	0.49
1:A:732:THR:OG1	1:A:955:ASN:OD1	2.30	0.49
1:B:596:SER:HB2	1:B:611:LEU:HG	1.94	0.49
1:B:781:VAL:HG13	1:B:782:PHE:HD2	1.77	0.49
1:A:353:TRP:HH2	1:A:464:PHE:HA	1.78	0.49
1:B:718:PHE:HE1	1:B:1108:ASN:HD21	1.58	0.49
1:A:559:PHE:HB2	1:A:584:ILE:HD13	1.94	0.48
1:A:1081:ILE:HG23	1:A:1088:HIS:HB2	1.95	0.48
1:A:1039:ARG:HH12	1:B:1039:ARG:HH11	1.60	0.48
1:A:599:THR:HB	1:A:608:VAL:HG23	1.94	0.48
1:B:756:TYR:HE2	1:B:997:ILE:HD12	1.76	0.48
1:C:440:ASN:OD1	1:C:440:ASN:N	2.40	0.48
1:A:22:THR:O	1:A:78:ARG:NH1	2.46	0.48
1:C:806:LEU:HD12	1:C:807:PRO:HD2	1.94	0.48
1:A:353:TRP:HH2	1:A:464:PHE:HA	1.78	0.48
1:A:503:VAL:HG21	1:C:503:VAL:HB	1.95	0.48
1:A:546:LEU:HD21	1:A:573:THR:HG21	1.95	0.48
1:B:429:PHE:HE2	1:B:514:SER:HA	1.77	0.48
1:B:727:LEU:HD11	1:B:1025:ALA:HB2	1.95	0.48
1:A:971:GLY:HA3	1:A:995:ARG:HH12	1.78	0.48
1:B:98:SER:HB2	1:B:179:LEU:HA	1.95	0.48
1:C:319:ARG:HH11	1:C:592:PHE:HB3	1.77	0.48
1:B:457:ARG:NE	1:B:459:SER:O	2.45	0.48
1:B:317:ASN:ND2	1:C:737:ASP:OD1	2.46	0.48
1:B:984:LEU:HD13	1:B:988:GLU:HG2	1.94	0.48
1:A:765:ARG:NH1	1:C:957:GLN:OE1	2.45	0.48
1:B:775:ASP:HB3	1:B:864:LEU:HD13	1.93	0.48
1:C:1050:MET:HE2	1:C:1050:MET:HB3	1.61	0.48
1:B:332:ILE:HG13	1:B:362:VAL:HG22	1.94	0.48
1:A:395:VAL:HG12	1:A:515:PHE:HB3	1.95	0.48
1:A:774:GLN:OE1	1:A:777:ASN:ND2	2.35	0.48

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:809:PRO:HA	1:A:814:LYS:HD2	1.94	0.48
1:A:31:SER:OG	1:A:60:SER:O	2.25	0.48
1:B:15:CYS:SG	1:B:137:ASN:N	2.81	0.48
1:A:115:GLN:HG3	1:A:233:ILE:HG12	1.95	0.48
1:B:858:LEU:HD13	1:B:959:LEU:HD12	1.94	0.48
1:C:133:PHE:HE1	1:C:160:TYR:HB3	1.77	0.48
1:B:216:LEU:HD13	1:B:266:TYR:HE1	1.77	0.48
1:C:82:PRO:O	1:C:239:GLN:NE2	2.44	0.48
1:A:43:PHE:HE1	1:C:563:GLN:HE21	1.60	0.48
1:A:901:GLN:HE21	1:A:905:ARG:HE	1.59	0.48
1:C:1111:GLU:OE1	1:C:1113:GLN:NE2	2.33	0.48
1:B:406:GLU:HB3	1:B:418:ILE:HG21	1.94	0.48
1:B:1116:THR:OG1	1:B:1118:ASP:OD1	2.31	0.48
1:C:398:ASP:OD2	1:C:423:TYR:OH	2.24	0.48
1:C:1115:ILE:HG22	1:C:1137:VAL:HG23	1.95	0.48
1:A:390:LEU:HD12	1:A:391:CYS:H	1.77	0.48
1:B:390:LEU:HD12	1:B:391:CYS:H	1.79	0.48
1:C:1054:GLN:HB2	1:C:1061:VAL:HG13	1.95	0.48
1:C:985:ASP:OD1	1:C:985:ASP:N	2.34	0.48
1:A:776:LYS:HG3	1:A:1019:ARG:HH22	1.78	0.48
1:A:393:THR:HA	1:A:522:ALA:HA	1.95	0.48
1:A:454:ARG:NH2	1:A:469:SER:O	2.43	0.48
1:B:401:VAL:HG12	1:B:509:ARG:HG2	1.95	0.48
1:A:966:LEU:HA	1:A:1000:ARG:HH12	1.78	0.48
1:B:461:LEU:HD12	1:B:465:GLU:HB3	1.95	0.48
1:A:1005:GLN:HE21	1:C:1006:THR:HG22	1.78	0.48
1:B:293:LEU:HD22	1:B:294:ASP:OD2	2.13	0.48
1:C:28:TYR:HB3	1:C:61:ASN:HB2	1.94	0.48
1:A:295:PRO:HA	1:A:298:GLU:HB2	1.96	0.48
1:A:994:ASP:HA	1:A:997:ILE:HG12	1.94	0.48
1:B:334:ASN:ND2	1:B:360:ASN:O	2.39	0.48
1:B:1013:ILE:HD13	1:C:1012:LEU:HG	1.94	0.48
1:C:393:THR:OG1	1:C:394:ASN:N	2.46	0.48
1:A:777:ASN:HA	1:A:780:GLU:HG3	1.96	0.48
1:A:896:ILE:HD12	1:A:897:PRO:HD2	1.94	0.48
1:A:48:LEU:HD11	1:A:276:LEU:HB3	1.95	0.48
1:A:342:PHE:O	1:A:509:ARG:NH2	2.46	0.48
1:A:345:THR:HG22	1:A:346:ARG:HG2	1.95	0.48
1:A:547:THR:HG22	1:B:982:SER:HB3	1.93	0.48
1:C:64:TRP:HZ2	1:C:214:ARG:HH22	1.62	0.48
1:A:134:GLN:HE22	1:A:162:SER:H	1.59	0.48

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:561:PRO:O	1:A:577:ARG:NH1	2.43	0.48
1:B:483:VAL:HG12	1:B:484:LYS:HG3	1.95	0.48
1:B:751:ASN:HA	1:B:754:LEU:HG	1.94	0.48
1:C:408:ARG:HD2	1:C:414:GLN:HE21	1.78	0.48
1:C:871:ALA:HA	1:C:874:THR:HG22	1.94	0.48
1:A:317:ASN:HD21	1:A:592:PHE:HB2	1.79	0.48
1:B:431:GLY:HA2	1:B:515:PHE:CZ	2.48	0.48
1:B:985:ASP:OD1	1:B:985:ASP:N	2.41	0.48
1:A:822:LEU:HD12	1:A:945:LEU:HD11	1.95	0.48
1:A:759:PHE:HA	1:A:762:GLN:HE21	1.78	0.48
1:A:805:ILE:HD12	1:A:1054:GLN:HE21	1.78	0.48
1:B:52:GLN:OE1	1:B:274:THR:OG1	2.25	0.48
1:C:738:CYS:HB2	1:C:760:CYS:HB2	1.37	0.48
1:C:915:VAL:HG21	1:C:1108:ASN:HD22	1.77	0.48
1:C:985:ASP:OD1	1:C:985:ASP:N	2.39	0.48
1:A:763:LEU:HB2	1:A:1008:VAL:HG11	1.95	0.48
1:A:353:TRP:HH2	1:A:464:PHE:HA	1.78	0.48
1:B:276:LEU:HG	1:B:306:PHE:HE1	1.78	0.48
1:A:722:VAL:HG22	1:A:930:ALA:HB1	1.95	0.48
1:B:701:VAL:HG13	1:C:787:GLN:HG3	1.95	0.48
1:B:330:PRO:HD3	1:B:579:PRO:HB2	1.95	0.48
1:B:517:LEU:HD11	1:C:983:ARG:HH11	1.78	0.48
1:A:873:TYR:HE1	1:C:699:LEU:HG	1.78	0.48
1:B:34:ARG:NH2	1:B:221:SER:H	2.11	0.48
1:B:552:LEU:HD12	1:B:585:LEU:HB2	1.94	0.48
1:C:732:THR:HB	1:C:955:ASN:HD22	1.78	0.48
1:A:390:LEU:HD12	1:A:391:CYS:H	1.78	0.48
1:B:592:PHE:HZ	1:C:740:MET:HE2	1.79	0.48
1:B:1079:PRO:HA	1:C:900:MET:HE1	1.94	0.48
1:C:819:GLU:HA	1:C:822:LEU:HG	1.94	0.48
1:A:315:THR:HG23	1:A:595:VAL:HG23	1.95	0.48
1:B:1106:GLN:NE2	1:B:1111:GLU:OE1	2.45	0.48
1:A:719:THR:HG23	1:A:1070:ALA:HB2	1.94	0.48
1:C:395:VAL:HG12	1:C:515:PHE:HB3	1.96	0.48
1:C:984:LEU:HD23	1:C:988:GLU:HG2	1.95	0.48
1:A:39:PRO:HG3	1:A:51:THR:HG21	1.96	0.48
1:A:972:ALA:HB2	1:A:996:LEU:HD21	1.95	0.48
1:B:483:VAL:HG12	1:B:484:LYS:HG3	1.95	0.48
1:A:722:VAL:HG12	1:A:1065:VAL:HG22	1.96	0.48
1:A:767:LEU:HA	1:A:770:ILE:HG22	1.95	0.48
1:A:53:ASP:OD1	1:A:54:LEU:N	2.45	0.48

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:276:LEU:HD11	1:A:304:LYS:HE2	1.94	0.48
1:A:1115:ILE:HG22	1:A:1137:VAL:HG23	1.96	0.48
1:A:403:ARG:NH2	1:B:372:ALA:O	2.39	0.48
1:B:426:PRO:HD2	1:B:429:PHE:HD2	1.79	0.48
1:B:491:PRO:HG2	1:B:492:LEU:HD22	1.94	0.48
1:C:414:GLN:NE2	1:C:414:GLN:HA	2.28	0.48
1:A:896:ILE:HD12	1:A:897:PRO:HD2	1.94	0.48
1:A:108:THR:O	1:A:237:ARG:NH2	2.46	0.48
1:B:785:VAL:HG12	1:B:787:GLN:H	1.78	0.48
1:B:997:ILE:O	1:B:1001:LEU:HB2	2.14	0.48
1:C:440:ASN:OD1	1:C:440:ASN:N	2.41	0.48
1:A:642:VAL:HG22	1:A:651:ILE:HG22	1.95	0.48
1:A:964:LYS:NZ	1:C:569:ILE:O	2.42	0.48
1:B:1116:THR:OG1	1:B:1118:ASP:OD1	2.30	0.48
1:A:1040:VAL:HG11	1:B:1035:GLY:HA3	1.96	0.48
1:B:90:VAL:HG13	1:B:267:VAL:HG13	1.95	0.48
1:C:729:VAL:HG12	1:C:1059:GLY:HA2	1.95	0.48
1:C:642:VAL:HG22	1:C:651:ILE:HG22	1.95	0.48
1:B:970:PHE:HD2	1:B:996:LEU:HA	1.79	0.48
1:B:64:TRP:HE1	1:B:264:ALA:HB1	1.78	0.48
1:B:726:ILE:HD13	1:B:1061:VAL:HG22	1.95	0.48
1:C:119:ILE:HG12	1:C:128:ILE:HG12	1.96	0.48
1:C:730:SER:HB2	1:C:774:GLN:NE2	2.22	0.48
1:A:989:ALA:O	1:A:993:ILE:HD12	2.13	0.48
1:C:1031:GLU:HB3	1:C:1037:SER:HB2	1.94	0.48
1:A:107:GLY:H	1:A:235:ILE:HD11	1.77	0.48
1:C:785:VAL:HG12	1:C:787:GLN:H	1.78	0.48
1:C:106:PHE:HB2	1:C:117:LEU:HB3	1.96	0.48
1:A:994:ASP:HA	1:A:997:ILE:HG12	1.94	0.48
1:A:403:ARG:NH2	1:B:372:ALA:O	2.38	0.48
1:B:326:ILE:HD13	1:B:533:LEU:HD12	1.96	0.48
1:B:687:GLN:NE2	1:B:688:THR:O	2.47	0.48
1:A:22:THR:O	1:A:78:ARG:NH1	2.46	0.48
1:C:599:THR:HB	1:C:608:VAL:HG23	1.95	0.48
1:A:80:ALA:O	1:A:245:HIS:NE2	2.45	0.48
1:B:642:VAL:HG22	1:B:651:ILE:HG22	1.95	0.48
1:B:83:VAL:HG13	1:B:237:ARG:NH1	2.28	0.48
1:B:237:ARG:NH1	1:B:239:GLN:OE1	2.44	0.48
1:B:977:LEU:HD21	1:B:996:LEU:HG	1.95	0.48
1:C:409:GLN:HE21	1:C:418:ILE:HG12	1.79	0.48
1:A:302:THR:HG21	1:A:315:THR:HG23	1.95	0.48

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:C:37:TYR:OH	1:C:53:ASP:OD1	2.27	0.48
1:C:360:ASN:H	1:C:523:THR:HB	1.79	0.48
1:C:896:ILE:HD12	1:C:897:PRO:HD2	1.95	0.48
1:A:115:GLN:HE22	1:C:468:ILE:HG21	1.77	0.48
1:A:442:ASP:OD1	1:A:448:ASN:ND2	2.47	0.48
1:A:345:THR:HG22	1:A:346:ARG:HG2	1.96	0.48
1:B:389:ASP:OD1	1:B:389:ASP:N	2.40	0.48
1:B:722:VAL:HG22	1:B:930:ALA:HB1	1.94	0.48
1:C:898:PHE:HE2	1:C:1050:MET:HE1	1.78	0.48
1:A:106:PHE:HB3	1:A:235:ILE:HD12	1.94	0.48
1:B:722:VAL:HG12	1:B:1065:VAL:HG22	1.96	0.48
1:B:728:PRO:HD3	1:B:947:LYS:HG3	1.95	0.48
1:B:884:SER:OG	1:B:887:THR:OG1	2.25	0.48
1:A:58:PHE:HD2	1:A:290:ASP:HB2	1.78	0.48
1:A:206:LYS:NZ	1:A:221:SER:OG	2.46	0.48
1:A:332:ILE:HG13	1:A:362:VAL:HG22	1.95	0.48
1:A:858:LEU:HD13	1:A:959:LEU:HD12	1.96	0.48
1:A:1106:GLN:NE2	1:A:1111:GLU:OE2	2.47	0.48
1:B:976:VAL:HG13	1:B:979:ASP:HB3	1.95	0.48
1:B:44:ARG:NH2	1:B:49:HIS:HB2	2.29	0.48
1:B:57:PRO:HB2	1:B:60:SER:HB3	1.96	0.48
1:B:90:VAL:HG13	1:B:267:VAL:HG13	1.96	0.48
1:B:183:GLN:HG3	1:B:187:LYS:HG3	1.95	0.48
1:A:456:PHE:HB3	1:A:473:TYR:CD1	2.48	0.48
1:C:884:SER:OG	1:C:887:THR:OG1	2.27	0.48
1:A:315:THR:HG23	1:A:595:VAL:HG23	1.95	0.48
1:C:394:ASN:OD1	1:C:394:ASN:N	2.46	0.48
1:A:364:ASP:OD2	1:A:364:ASP:N	2.44	0.48
1:B:951:VAL:HA	1:B:954:GLN:HG3	1.94	0.48
1:B:731:MET:H	1:B:774:GLN:NE2	2.12	0.48
1:C:483:VAL:HG12	1:C:484:LYS:HG3	1.96	0.48
1:A:108:THR:OG1	1:A:234:ASN:O	2.30	0.48
1:C:708:SER:HB3	1:C:711:SER:HB3	1.96	0.48
1:A:196:ASN:HD22	1:A:235:ILE:HG22	1.79	0.48
1:A:767:LEU:HA	1:A:770:ILE:HG22	1.95	0.48
1:A:906:PHE:HB3	1:A:911:VAL:HB	1.96	0.48
1:C:212:LEU:HD22	1:C:217:PRO:HD3	1.96	0.48
1:C:784:GLN:HA	1:C:784:GLN:HE21	1.79	0.48
1:C:976:VAL:HG13	1:C:979:ASP:HB3	1.95	0.48
1:A:317:ASN:HD21	1:A:592:PHE:HB2	1.79	0.48
1:A:357:ARG:NH2	1:A:396:TYR:OH	2.47	0.48

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:535:LYS:HD2	1:A:585:LEU:HD11	1.95	0.48
1:A:745:ASP:OD1	1:C:319:ARG:NH1	2.46	0.48
1:B:727:LEU:HD11	1:B:1025:ALA:HB2	1.96	0.48
1:B:785:VAL:HG12	1:B:787:GLN:H	1.78	0.48
1:C:58:PHE:HD2	1:C:290:ASP:HB2	1.77	0.48
1:B:577:ARG:HH21	1:B:582:LEU:HD22	1.78	0.48
1:C:738:CYS:O	1:C:742:ILE:HB	2.13	0.48
1:B:611:LEU:HD13	1:B:650:LEU:HD22	1.96	0.48
1:C:908:GLY:O	1:C:1038:LYS:NZ	2.30	0.48
1:C:1051:SER:OG	1:C:1064:HIS:ND1	2.33	0.48
1:A:950:ASP:N	1:A:950:ASP:OD1	2.45	0.48
1:B:390:LEU:HD21	1:C:983:ARG:HG2	1.96	0.48
1:A:642:VAL:HG22	1:A:651:ILE:HG22	1.96	0.48
1:A:906:PHE:HB3	1:A:911:VAL:HB	1.96	0.48
1:A:296:LEU:O	1:A:299:THR:OG1	2.28	0.48
1:A:599:THR:HB	1:A:608:VAL:HG23	1.94	0.48
1:B:822:LEU:HD21	1:B:1061:VAL:HG21	1.96	0.48
1:B:1116:THR:OG1	1:B:1118:ASP:OD1	2.31	0.48
1:B:990:GLU:HA	1:B:993:ILE:HD12	1.95	0.48
1:A:229:LEU:HD12	1:A:230:PRO:HD2	1.95	0.48
1:A:864:LEU:HD12	1:A:865:LEU:HD13	1.94	0.48
1:B:777:ASN:HA	1:B:780:GLU:HG3	1.94	0.48
1:B:726:ILE:HD13	1:B:1061:VAL:HG22	1.96	0.48
1:B:390:LEU:HD12	1:B:391:CYS:H	1.79	0.48
1:C:130:VAL:HG21	1:C:231:ILE:HD11	1.96	0.48
1:C:389:ASP:OD1	1:C:528:LYS:NZ	2.38	0.48
1:A:106:PHE:HD1	1:A:235:ILE:HD13	1.79	0.48
1:A:599:THR:HB	1:A:608:VAL:HG23	1.94	0.48
1:B:733:LYS:HB3	1:B:861:LEU:HD23	1.96	0.48
1:A:1050:MET:HE2	1:A:1050:MET:HB3	1.67	0.48
1:C:360:ASN:H	1:C:523:THR:HB	1.78	0.48
1:A:200:TYR:OH	1:C:464:PHE:O	2.25	0.48
1:C:871:ALA:HA	1:C:874:THR:HG22	1.96	0.48
1:A:49:HIS:NE2	1:A:51:THR:HB	2.29	0.48
1:A:92:PHE:HE2	1:A:265:TYR:HB2	1.79	0.48
1:B:1091:ARG:NH1	1:B:1120:THR:O	2.45	0.48
1:B:699:LEU:HG	1:C:873:TYR:HE1	1.79	0.48
1:B:902:MET:HE1	1:B:1050:MET:HE2	1.94	0.48
1:C:366:SER:O	1:C:370:ASN:HB2	2.13	0.48
1:B:15:CYS:SG	1:B:137:ASN:N	2.86	0.48
1:B:731:MET:H	1:B:774:GLN:NE2	2.11	0.48

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:C:483:VAL:HG12	1:C:484:LYS:HG3	1.95	0.48
1:C:762:GLN:HA	1:C:765:ARG:HE	1.78	0.48
1:A:106:PHE:HB2	1:A:117:LEU:HB2	1.96	0.48
1:B:1045:LYS:NZ	1:C:889:GLY:O	2.47	0.48
1:A:61:ASN:O	1:A:61:ASN:ND2	2.34	0.48
1:A:931:ILE:HA	1:A:934:ILE:HG22	1.95	0.48
1:B:949:GLN:O	1:B:953:ASN:ND2	2.47	0.48
1:A:38:TYR:HE1	1:A:285:ILE:HG12	1.79	0.48
1:A:105:ILE:HB	1:A:241:LEU:HD21	1.96	0.48
1:C:83:VAL:HG13	1:C:239:GLN:HE21	1.78	0.48
1:A:389:ASP:OD1	1:A:528:LYS:NZ	2.46	0.48
1:A:395:VAL:HG22	1:A:515:PHE:HB3	1.95	0.48
1:A:1033:VAL:HG11	1:A:1053:PRO:HG3	1.96	0.48
1:B:334:ASN:ND2	1:B:360:ASN:O	2.41	0.48
1:A:84:LEU:HD11	1:A:267:VAL:HG11	1.96	0.48
1:A:415:THR:OG1	1:B:369:TYR:OH	2.24	0.48
1:A:765:ARG:NH1	1:C:957:GLN:OE1	2.46	0.48
1:B:1116:THR:OG1	1:B:1118:ASP:OD1	2.31	0.48
1:A:566:GLY:HA2	1:B:43:PHE:HB3	1.96	0.48
1:B:681:GLY:N	1:C:864:LEU:O	2.47	0.48
1:C:34:ARG:NH2	1:C:221:SER:H	2.11	0.48
1:C:52:GLN:OE1	1:C:274:THR:OG1	2.24	0.48
1:C:457:ARG:NE	1:C:467:ASP:OD1	2.46	0.48
1:A:1105:THR:HG21	1:A:1110:TYR:HD2	1.77	0.48
1:B:279:TYR:HE1	1:B:285:ILE:HG12	1.79	0.48
1:B:722:VAL:HG12	1:B:1065:VAL:HG22	1.96	0.48
1:B:738:CYS:O	1:B:742:ILE:HB	2.14	0.48
1:A:858:LEU:HD13	1:A:959:LEU:HD12	1.96	0.48
1:A:1116:THR:OG1	1:A:1118:ASP:OD1	2.31	0.48
1:A:345:THR:HG22	1:A:346:ARG:HG2	1.96	0.48
1:A:567:ARG:HG2	1:A:571:ASP:HA	1.96	0.48
1:A:896:ILE:HD12	1:A:897:PRO:HD2	1.96	0.48
1:C:973:ILE:HG21	1:C:983:ARG:HH12	1.79	0.48
1:A:976:VAL:HG13	1:A:979:ASP:HB3	1.95	0.48
1:C:407:VAL:HA	1:C:410:ILE:HG12	1.96	0.48
1:C:884:SER:OG	1:C:894:LEU:O	2.32	0.48
1:C:980:ILE:HG23	1:C:984:LEU:HD23	1.95	0.48
1:A:1072:GLU:OE1	1:A:1072:GLU:N	2.47	0.48
1:C:650:LEU:HD13	1:C:653:ALA:HB3	1.96	0.48
1:B:401:VAL:HG22	1:B:509:ARG:HG2	1.95	0.48
1:B:577:ARG:HB2	1:B:584:ILE:HG12	1.96	0.48

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:C:435:ALA:HB2	1:C:510:VAL:HG12	1.96	0.48
1:A:717:ASN:OD1	1:A:718:PHE:N	2.46	0.48
1:B:96:GLU:OE1	1:B:100:ILE:N	2.46	0.48
1:A:574:ASP:OD1	1:A:574:ASP:N	2.46	0.48
1:C:1028:LYS:NZ	1:C:1042:PHE:O	2.47	0.48
1:C:394:ASN:N	1:C:394:ASN:OD1	2.46	0.48
1:A:867:ASP:OD1	1:A:867:ASP:N	2.47	0.48
1:A:931:ILE:HA	1:A:934:ILE:HG22	1.96	0.48
1:C:52:GLN:OE1	1:C:274:THR:OG1	2.23	0.48
1:C:189:LEU:HD22	1:C:217:PRO:HG2	1.96	0.48
1:C:965:GLN:HE21	1:C:965:GLN:CA	2.27	0.48
1:A:574:ASP:OD1	1:A:574:ASP:N	2.46	0.48
1:C:642:VAL:HG22	1:C:651:ILE:HG22	1.95	0.48
1:B:619:GLU:N	1:B:619:GLU:OE1	2.46	0.48
1:A:1081:ILE:HD11	1:A:1135:ASN:HB3	1.96	0.48
1:B:390:LEU:HD12	1:B:391:CYS:H	1.79	0.48
1:A:1115:ILE:HG22	1:A:1137:VAL:HG23	1.96	0.48
1:B:206:LYS:NZ	1:B:221:SER:OG	2.46	0.48
1:B:222:ALA:HB2	1:B:285:ILE:HB	1.96	0.48
1:B:358:ILE:HB	1:B:395:VAL:HG23	1.96	0.48
1:C:206:LYS:HB2	1:C:223:LEU:HA	1.96	0.47
1:A:151:SER:HB2	1:A:153:MET:HE3	1.96	0.47
1:C:390:LEU:HD12	1:C:391:CYS:H	1.79	0.47
1:C:532:ASN:N	1:C:532:ASN:OD1	2.46	0.47
1:C:906:PHE:HD1	1:C:916:LEU:HB2	1.78	0.47
1:A:346:ARG:HA	1:A:509:ARG:HH22	1.79	0.47
1:C:34:ARG:NH2	1:C:221:SER:H	2.10	0.47
1:C:87:ASN:OD1	1:C:87:ASN:N	2.45	0.47
1:B:52:GLN:OE1	1:B:274:THR:OG1	2.26	0.47
1:C:229:LEU:HD22	1:C:231:ILE:HB	1.96	0.47
1:A:393:THR:HA	1:A:522:ALA:HA	1.95	0.47
1:A:869:MET:HE2	1:A:869:MET:HB3	1.69	0.47
1:B:193:VAL:HG23	1:B:270:LEU:HD21	1.96	0.47
1:B:822:LEU:HD23	1:B:1056:ALA:HB2	1.95	0.47
1:C:809:PRO:O	1:C:814:LYS:NZ	2.41	0.47
1:B:131:CYS:HB2	1:B:133:PHE:CE1	2.49	0.47
1:B:610:VAL:N	1:B:651:ILE:O	2.46	0.47
1:C:568:ASP:HB3	1:C:574:ASP:HB2	1.96	0.47
1:C:552:LEU:HD13	1:C:585:LEU:HD22	1.95	0.47
1:A:204:TYR:HD1	1:A:225:PRO:HA	1.78	0.47
1:A:365:TYR:HD1	1:A:368:LEU:HD12	1.79	0.47

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:C:396:TYR:HB2	1:C:514:SER:HB2	1.96	0.47
1:A:15:CYS:SG	1:A:137:ASN:N	2.82	0.47
1:A:566:GLY:HA2	1:B:43:PHE:HB3	1.96	0.47
1:B:719:THR:HG23	1:B:1070:ALA:HB2	1.96	0.47
1:C:177:MET:HE3	1:C:178:ASP:H	1.78	0.47
1:A:14:GLN:O	1:A:158:ARG:NE	2.47	0.47
1:A:738:CYS:HB2	1:A:760:CYS:HB2	1.55	0.47
1:A:765:ARG:HH12	1:C:957:GLN:CD	2.22	0.47
1:A:915:VAL:HG21	1:A:1109:PHE:HE2	1.79	0.47
1:C:36:VAL:O	1:C:223:LEU:HG	2.14	0.47
1:C:901:GLN:HE21	1:C:1050:MET:HE1	1.79	0.47
1:B:960:ASN:O	1:B:964:LYS:HB2	2.14	0.47
1:A:101:ILE:HD13	1:A:240:THR:HG23	1.95	0.47
1:A:390:LEU:HD12	1:A:391:CYS:H	1.79	0.47
1:C:732:THR:HB	1:C:955:ASN:ND2	2.29	0.47
1:C:1116:THR:OG1	1:C:1118:ASP:OD1	2.29	0.47
1:C:1156:PHE:HA	1:C:1159:HIS:CD2	2.48	0.47
1:A:384:PRO:HG3	1:C:415:THR:HG21	1.96	0.47
1:B:777:ASN:HA	1:B:780:GLU:HG3	1.96	0.47
1:C:877:LEU:O	1:C:881:THR:OG1	2.28	0.47
1:B:314:GLN:HE21	1:B:314:GLN:CA	2.27	0.47
1:C:110:LEU:HD11	1:C:237:ARG:HB2	1.96	0.47
1:A:1106:GLN:OE1	1:A:1106:GLN:N	2.47	0.47
1:B:871:ALA:HA	1:B:874:THR:HG22	1.95	0.47
1:C:965:GLN:HE21	1:C:965:GLN:CA	2.21	0.47
1:A:562:PHE:O	1:B:41:LYS:NZ	2.47	0.47
1:A:1115:ILE:HG22	1:A:1137:VAL:HG23	1.96	0.47
1:A:84:LEU:HD13	1:A:238:PHE:HE2	1.79	0.47
1:B:297:SER:HA	1:B:300:LYS:HG2	1.96	0.47
1:A:916:LEU:HD22	1:A:917:TYR:HD1	1.78	0.47
1:B:719:THR:HG23	1:B:1070:ALA:HB2	1.96	0.47
1:C:393:THR:OG1	1:C:394:ASN:N	2.47	0.47
1:A:129:LYS:NZ	1:A:131:CYS:SG	2.71	0.47
1:A:955:ASN:C	1:A:955:ASN:ND2	2.65	0.47
1:C:334:ASN:OD1	1:C:335:LEU:N	2.43	0.47
1:A:390:LEU:HD12	1:A:391:CYS:H	1.78	0.47
1:A:911:VAL:HG11	1:A:1067:TYR:HE1	1.78	0.47
1:C:34:ARG:NH1	1:C:219:GLY:O	2.46	0.47
1:A:106:PHE:HB2	1:A:117:LEU:HB2	1.96	0.47
1:A:957:GLN:HG3	1:B:765:ARG:HH21	1.79	0.47
1:B:483:VAL:HG12	1:B:484:LYS:HG3	1.95	0.47

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:C:404:GLY:HA2	1:C:407:VAL:HG13	1.96	0.47
1:B:52:GLN:OE1	1:B:274:THR:OG1	2.26	0.47
1:C:33:THR:OG1	1:C:219:GLY:O	2.33	0.47
1:C:58:PHE:HD1	1:C:290:ASP:HB2	1.77	0.47
1:C:380:TYR:HE1	1:C:433:VAL:HG12	1.79	0.47
1:A:599:THR:HB	1:A:608:VAL:HG23	1.95	0.47
1:B:398:ASP:OD2	1:B:423:TYR:OH	2.23	0.47
1:C:81:ASN:O	1:C:239:GLN:NE2	2.41	0.47
1:A:423:TYR:HE1	1:A:512:VAL:HG11	1.79	0.47
1:B:398:ASP:OD2	1:B:423:TYR:OH	2.27	0.47
1:C:37:TYR:OH	1:C:53:ASP:OD1	2.25	0.47
1:B:393:THR:OG1	1:B:394:ASN:OD1	2.32	0.47
1:B:431:GLY:HA2	1:B:515:PHE:CZ	2.50	0.47
1:A:972:ALA:HB2	1:A:996:LEU:HD21	1.95	0.47
1:B:37:TYR:OH	1:B:53:ASP:OD1	2.28	0.47
1:B:699:LEU:HG	1:C:873:TYR:HE1	1.79	0.47
1:C:189:LEU:HD12	1:C:217:PRO:HG2	1.96	0.47
1:A:334:ASN:ND2	1:A:360:ASN:O	2.42	0.47
1:A:501:TYR:HB3	1:A:505:TYR:HB3	1.95	0.47
1:A:787:GLN:HG2	1:C:701:VAL:HG13	1.95	0.47
1:C:52:GLN:OE1	1:C:274:THR:OG1	2.29	0.47
1:A:1084:ASP:OD1	1:A:1084:ASP:N	2.39	0.47
1:C:423:TYR:HD1	1:C:466:ARG:HB3	1.79	0.47
1:C:856:ASN:ND2	1:C:858:LEU:HD23	2.29	0.47
1:B:129:LYS:HG2	1:B:133:PHE:HZ	1.78	0.47
1:A:353:TRP:HE1	1:A:466:ARG:HB2	1.80	0.47
1:B:185:ASN:HB2	1:B:213:VAL:HA	1.96	0.47
1:B:332:ILE:HG13	1:B:362:VAL:HG22	1.96	0.47
1:C:203:ILE:HD11	1:C:227:VAL:HB	1.96	0.47
1:A:1100:THR:OG1	1:A:1101:HIS:ND1	2.38	0.47
1:C:1052:PHE:HB2	1:C:1063:LEU:HD12	1.96	0.47
1:B:976:VAL:HG13	1:B:979:ASP:HB3	1.96	0.47
1:C:108:THR:O	1:C:237:ARG:NH1	2.47	0.47
1:A:930:ALA:HA	1:A:933:LYS:HG2	1.96	0.47
1:B:713:ALA:HB2	1:C:895:GLN:OE1	2.14	0.47
1:B:1116:THR:OG1	1:B:1118:ASP:OD1	2.31	0.47
1:C:976:VAL:HG13	1:C:979:ASP:HB3	1.95	0.47
1:B:727:LEU:HD21	1:B:1024:LEU:HD22	1.96	0.47
1:C:37:TYR:OH	1:C:53:ASP:OD2	2.22	0.47
1:A:53:ASP:OD2	1:A:195:LYS:NZ	2.48	0.47
1:B:739:THR:HA	1:B:753:LEU:HD21	1.95	0.47

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:B:777:ASN:HA	1:B:780:GLU:HG3	1.95	0.47
1:C:276:LEU:HD11	1:C:301:CYS:HA	1.96	0.47
1:C:984:LEU:HD23	1:C:988:GLU:HG2	1.96	0.47
1:C:1084:ASP:OD2	1:C:1086:LYS:NZ	2.48	0.47
1:B:295:PRO:HG2	1:B:608:VAL:HG21	1.96	0.47
1:C:222:ALA:HB2	1:C:285:ILE:HB	1.96	0.47
1:C:382:VAL:HG11	1:C:387:LEU:HD13	1.96	0.47
1:C:877:LEU:HG	1:C:1053:PRO:HG2	1.96	0.47
1:B:37:TYR:OH	1:B:54:LEU:O	2.31	0.47
1:B:715:PRO:HD3	1:C:894:LEU:HD11	1.96	0.47
1:C:22:THR:O	1:C:78:ARG:NH1	2.47	0.47
1:C:58:PHE:HD1	1:C:290:ASP:HB2	1.79	0.47
1:A:897:PRO:HG2	1:A:900:MET:HG2	1.94	0.47
1:A:1115:ILE:HG22	1:A:1137:VAL:HG23	1.96	0.47
1:A:474:GLN:HG2	1:A:480:CYS:H	1.79	0.47
1:B:1046:GLY:HA2	1:C:890:ALA:HA	1.96	0.47
1:A:730:SER:OG	1:A:731:MET:N	2.46	0.47
1:B:133:PHE:HD1	1:B:160:TYR:HB3	1.79	0.47
1:B:950:ASP:OD1	1:B:950:ASP:N	2.46	0.47
1:A:117:LEU:HG	1:A:128:ILE:HD11	1.96	0.47
1:B:390:LEU:HD12	1:B:391:CYS:H	1.79	0.47
1:C:101:ILE:HD11	1:C:240:THR:HB	1.96	0.47
1:B:443:SER:HB3	1:B:499:PRO:HG3	1.96	0.47
1:A:53:ASP:OD1	1:A:54:LEU:N	2.45	0.47
1:A:611:LEU:HD22	1:A:678:ILE:HD11	1.95	0.47
1:A:896:ILE:HD12	1:A:897:PRO:HD2	1.95	0.47
1:B:408:ARG:HH21	1:B:409:GLN:HE21	1.62	0.47
1:C:310:LYS:HG2	1:C:676:ILE:HG21	1.96	0.47
1:C:774:GLN:HE22	1:C:1018:ILE:HG21	1.80	0.47
1:C:822:LEU:HD13	1:C:1056:ALA:HB2	1.97	0.47
1:A:950:ASP:N	1:A:950:ASP:OD1	2.46	0.47
1:C:177:MET:HE3	1:C:178:ASP:H	1.80	0.47
1:A:108:THR:O	1:A:237:ARG:NH2	2.47	0.47
1:A:390:LEU:HD12	1:A:391:CYS:H	1.78	0.47
1:C:687:GLN:HG2	1:C:693:ILE:HD13	1.97	0.47
1:A:910:GLY:O	1:A:1106:GLN:OE1	2.31	0.47
1:A:37:TYR:OH	1:A:53:ASP:OD1	2.26	0.47
1:A:574:ASP:OD1	1:A:574:ASP:N	2.46	0.47
1:B:315:THR:HG23	1:B:595:VAL:HG23	1.95	0.47
1:C:389:ASP:OD1	1:C:528:LYS:NZ	2.40	0.47
1:C:884:SER:OG	1:C:894:LEU:O	2.33	0.47

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:896:ILE:HD12	1:A:897:PRO:HD2	1.96	0.47
1:A:785:VAL:HG12	1:A:787:GLN:H	1.78	0.47
1:B:1054:GLN:HB2	1:B:1061:VAL:HG23	1.96	0.47
1:A:384:PRO:HG3	1:C:415:THR:HG21	1.96	0.47
1:A:560:LEU:HD12	1:A:562:PHE:H	1.79	0.47
1:B:457:ARG:HH12	1:B:461:LEU:HD23	1.79	0.47
1:C:978:ASN:HA	1:C:981:LEU:HG	1.97	0.47
1:A:299:THR:HG21	1:A:597:VAL:HG21	1.96	0.47
1:C:1009:THR:O	1:C:1013:ILE:HG12	2.15	0.47
1:A:39:PRO:HG3	1:A:51:THR:HG21	1.96	0.47
1:B:573:THR:HG22	1:B:587:ILE:HD13	1.95	0.47
1:A:330:PRO:HD3	1:A:579:PRO:HB2	1.96	0.47
1:B:517:LEU:HD11	1:C:983:ARG:HD2	1.96	0.47
1:A:858:LEU:HD13	1:A:959:LEU:HD12	1.96	0.47
1:A:553:THR:OG1	1:A:586:ASP:OD1	2.28	0.47
1:B:389:ASP:OD1	1:B:528:LYS:NZ	2.41	0.47
1:C:809:PRO:HA	1:C:814:LYS:HD2	1.96	0.47
1:B:317:ASN:HD21	1:B:592:PHE:HD1	1.61	0.47
1:A:767:LEU:HA	1:A:770:ILE:HG12	1.97	0.47
1:C:966:LEU:HA	1:C:1000:ARG:HH12	1.79	0.47
1:A:751:ASN:HA	1:A:754:LEU:HG	1.97	0.47
1:A:1115:ILE:HG22	1:A:1137:VAL:HG23	1.97	0.47
1:C:58:PHE:HD2	1:C:290:ASP:HB2	1.80	0.47
1:A:971:GLY:HA3	1:A:995:ARG:HH22	1.80	0.47
1:C:1054:GLN:HB2	1:C:1061:VAL:HG12	1.96	0.47
1:A:866:THR:H	1:A:869:MET:HE2	1.78	0.47
1:B:369:TYR:CZ	1:B:384:PRO:HB2	2.50	0.47
1:C:129:LYS:NZ	1:C:168:PHE:O	2.36	0.47
1:A:798:GLY:O	1:A:920:GLN:NE2	2.47	0.47
1:B:912:THR:OG1	1:B:1106:GLN:NE2	2.35	0.47
1:C:431:GLY:HA2	1:C:515:PHE:CE1	2.49	0.47
1:C:730:SER:OG	1:C:731:MET:N	2.48	0.47
1:C:976:VAL:HG23	1:C:979:ASP:HB3	1.97	0.47
1:C:1050:MET:HE2	1:C:1050:MET:HB3	1.60	0.47
1:B:1033:VAL:HG21	1:B:1053:PRO:HG3	1.95	0.47
1:B:931:ILE:HA	1:B:934:ILE:HG22	1.95	0.47
1:C:310:LYS:HG2	1:C:676:ILE:HG21	1.95	0.47
1:C:389:ASP:OD1	1:C:528:LYS:NZ	2.33	0.47
1:A:577:ARG:HD3	1:A:584:ILE:HG12	1.97	0.47
1:C:424:LYS:HD3	1:C:463:PRO:HA	1.96	0.47
1:A:96:GLU:OE1	1:A:99:ASN:ND2	2.39	0.47

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:C:889:GLY:HA3	1:C:1034:LEU:HD22	1.96	0.47
1:B:722:VAL:HG22	1:B:930:ALA:HB1	1.97	0.47
1:B:1019:ARG:O	1:B:1019:ARG:NH1	2.46	0.47
1:A:115:GLN:NE2	1:C:466:ARG:O	2.48	0.47
1:A:931:ILE:HA	1:A:934:ILE:HG22	1.97	0.47
1:A:1031:GLU:HB3	1:A:1037:SER:HB2	1.97	0.47
1:B:366:SER:O	1:B:370:ASN:HB2	2.14	0.47
1:C:34:ARG:HH22	1:C:220:PHE:HA	1.79	0.47
1:C:183:GLN:HG3	1:C:187:LYS:HG3	1.97	0.47
1:A:409:GLN:N	1:A:409:GLN:OE1	2.48	0.47
1:A:678:ILE:HD11	1:A:684:ALA:HB2	1.97	0.47
1:A:711:SER:OG	1:B:895:GLN:OE1	2.32	0.47
1:B:365:TYR:CE2	1:B:387:LEU:HD12	2.50	0.47
1:C:1033:VAL:HG21	1:C:1053:PRO:HG3	1.96	0.47
1:A:577:ARG:HD2	1:A:582:LEU:HA	1.96	0.47
1:B:677:PRO:HB3	1:C:864:LEU:HD21	1.97	0.47
1:B:203:ILE:HD11	1:B:226:LEU:HD23	1.96	0.47
1:A:785:VAL:HG12	1:A:787:GLN:H	1.79	0.47
1:C:613:GLN:HA	1:C:648:GLY:HA3	1.95	0.47
1:C:1156:PHE:O	1:C:1160:THR:OG1	2.32	0.47
1:A:742:ILE:HD11	1:A:753:LEU:HD22	1.96	0.47
1:A:1013:ILE:HG13	1:B:1012:LEU:HG	1.95	0.47
1:C:52:GLN:OE1	1:C:274:THR:OG1	2.25	0.47
1:A:34:ARG:NH2	1:A:221:SER:H	2.12	0.47
1:A:115:GLN:HB2	1:A:233:ILE:HG12	1.96	0.47
1:A:950:ASP:N	1:A:950:ASP:OD1	2.45	0.47
1:C:732:THR:OG1	1:C:955:ASN:ND2	2.47	0.47
1:B:385:THR:OG1	1:C:985:ASP:OD2	2.33	0.47
1:A:642:VAL:HG22	1:A:651:ILE:HG22	1.97	0.47
1:A:805:ILE:HG22	1:A:878:LEU:HD21	1.96	0.47
1:B:785:VAL:HG12	1:B:787:GLN:H	1.79	0.47
1:C:908:GLY:C	1:C:1038:LYS:HZ2	2.23	0.47
1:A:688:THR:HA	1:A:690:GLN:HG2	1.97	0.47
1:A:759:PHE:HA	1:A:762:GLN:HE21	1.79	0.47
1:A:796:ASP:OD1	1:A:796:ASP:N	2.48	0.47
1:B:576:VAL:HG12	1:B:587:ILE:HD11	1.95	0.47
1:C:884:SER:OG	1:C:894:LEU:O	2.33	0.47
1:A:729:VAL:HG21	1:A:781:VAL:HG11	1.96	0.47
1:C:1010:GLN:HE22	1:C:1014:ARG:HH21	1.62	0.47
1:A:916:LEU:O	1:A:920:GLN:HB2	2.15	0.47
1:B:612:TYR:HB2	1:B:649:CYS:HB3	1.95	0.47

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:819:GLU:HA	1:A:822:LEU:HG	1.97	0.47
1:B:417:ASN:O	1:B:421:TYR:HB2	2.15	0.47
1:B:1010:GLN:OE1	1:B:1014:ARG:NH1	2.47	0.47
1:B:1103:PHE:CD1	1:B:1112:PRO:HB3	2.49	0.47
1:C:344:ALA:HB3	1:C:347:PHE:HE2	1.80	0.47
1:A:809:PRO:HA	1:A:814:LYS:HD2	1.97	0.47
1:A:894:LEU:HD21	1:C:715:PRO:HD3	1.95	0.47
1:B:334:ASN:ND2	1:B:360:ASN:O	2.39	0.47
1:A:867:ASP:OD1	1:A:867:ASP:N	2.46	0.47
1:B:83:VAL:HG22	1:B:239:GLN:HE22	1.79	0.47
1:B:517:LEU:HD11	1:C:983:ARG:HH21	1.80	0.47
1:C:738:CYS:O	1:C:742:ILE:HB	2.15	0.47
1:A:403:ARG:NH2	1:B:372:ALA:O	2.33	0.47
1:A:503:VAL:HA	1:A:506:GLN:HG3	1.96	0.47
1:A:895:GLN:NE2	1:C:711:SER:OG	2.46	0.47
1:B:877:LEU:O	1:B:881:THR:HG23	2.15	0.47
1:C:37:TYR:HB3	1:C:223:LEU:HD23	1.96	0.47
1:C:483:VAL:HG12	1:C:484:LYS:HG3	1.97	0.47
1:B:34:ARG:HG3	1:B:216:LEU:HD11	1.97	0.47
1:B:326:ILE:HD13	1:B:533:LEU:HD12	1.97	0.47
1:A:919:ASN:HD22	1:A:922:LEU:HD21	1.79	0.47
1:B:390:LEU:HD12	1:B:391:CYS:H	1.79	0.47
1:C:401:VAL:HG22	1:C:509:ARG:HG2	1.97	0.47
1:C:439:ASN:ND2	1:C:506:GLN:OE1	2.48	0.47
1:C:785:VAL:HG12	1:C:787:GLN:H	1.79	0.47
1:C:871:ALA:HA	1:C:874:THR:HG22	1.95	0.47
1:A:53:ASP:OD2	1:A:195:LYS:NZ	2.47	0.47
1:A:345:THR:HG22	1:A:346:ARG:HG2	1.96	0.47
1:C:379:CYS:HA	1:C:432:CYS:HA	1.97	0.47
1:C:958:ALA:HB1	1:C:1007:TYR:HE2	1.80	0.47
1:A:334:ASN:ND2	1:A:360:ASN:O	2.41	0.47
1:A:751:ASN:HA	1:A:754:LEU:HG	1.97	0.47
1:B:950:ASP:OD1	1:B:950:ASP:N	2.46	0.47
1:B:310:LYS:HE2	1:B:664:ILE:HD11	1.97	0.47
1:B:1091:ARG:NH1	1:B:1118:ASP:O	2.48	0.47
1:A:189:LEU:HD22	1:A:208:THR:HB	1.96	0.47
1:A:722:VAL:HG12	1:A:1065:VAL:HG22	1.97	0.47
1:B:276:LEU:HD11	1:B:304:LYS:HE2	1.96	0.47
1:B:731:MET:H	1:B:774:GLN:NE2	2.13	0.47
1:B:877:LEU:O	1:B:881:THR:OG1	2.30	0.47
1:B:978:ASN:HA	1:B:981:LEU:HG	1.97	0.47

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:B:994:ASP:HA	1:B:997:ILE:HG12	1.97	0.47
1:B:28:TYR:HB3	1:B:61:ASN:HB3	1.96	0.47
1:B:195:LYS:HE3	1:B:202:LYS:HD3	1.97	0.47
1:B:533:LEU:HD21	1:B:585:LEU:HD11	1.95	0.47
1:A:102:ARG:HE	1:A:246:ILE:HD11	1.80	0.47
1:B:483:VAL:HG12	1:B:484:LYS:HG3	1.96	0.47
1:B:950:ASP:OD1	1:B:950:ASP:N	2.46	0.47
1:C:196:ASN:HD22	1:C:235:ILE:HG12	1.78	0.47
1:A:15:CYS:SG	1:A:137:ASN:N	2.87	0.47
1:A:931:ILE:HA	1:A:934:ILE:HG22	1.97	0.47
1:B:390:LEU:HD12	1:B:391:CYS:H	1.80	0.47
1:B:442:ASP:O	1:B:448:ASN:ND2	2.43	0.47
1:C:785:VAL:HG12	1:C:787:GLN:H	1.79	0.47
1:A:194:PHE:HE2	1:A:203:ILE:HG23	1.80	0.47
1:A:752:LEU:HD12	1:A:753:LEU:HD22	1.97	0.47
1:A:984:LEU:HB2	1:A:989:ALA:HB2	1.97	0.47
1:C:190:ARG:CZ	1:C:207:HIS:HE1	2.27	0.47
1:B:574:ASP:OD1	1:B:574:ASP:N	2.47	0.47
1:B:442:ASP:OD2	1:B:509:ARG:NE	2.44	0.47
1:B:560:LEU:HB2	1:B:563:GLN:HE22	1.80	0.47
1:C:914:ASN:HD21	1:C:1111:GLU:CD	2.23	0.47
1:A:574:ASP:OD1	1:A:574:ASP:N	2.47	0.47
1:A:785:VAL:HG12	1:A:787:GLN:H	1.79	0.47
1:B:606:ASN:C	1:B:606:ASN:ND2	2.72	0.47
1:C:395:VAL:HG12	1:C:515:PHE:HB3	1.96	0.47
1:A:599:THR:HB	1:A:608:VAL:HG23	1.96	0.47
1:A:398:ASP:HB2	1:A:512:VAL:HG13	1.96	0.47
1:C:884:SER:OG	1:C:887:THR:OG1	2.26	0.47
1:B:688:THR:HA	1:B:690:GLN:HG2	1.97	0.47
1:B:711:SER:OG	1:C:895:GLN:NE2	2.41	0.47
1:B:295:PRO:HG2	1:B:608:VAL:HG21	1.97	0.47
1:A:777:ASN:HA	1:A:780:GLU:HG3	1.97	0.47
1:C:722:VAL:HG12	1:C:1065:VAL:HG22	1.97	0.47
1:C:966:LEU:HA	1:C:1000:ARG:HH12	1.79	0.47
1:C:334:ASN:OD1	1:C:335:LEU:N	2.43	0.47
1:C:393:THR:OG1	1:C:394:ASN:N	2.48	0.47
1:A:972:ALA:HB2	1:A:996:LEU:HD21	1.96	0.47
1:B:483:VAL:HG12	1:B:484:LYS:HG3	1.96	0.47
1:A:390:LEU:HD12	1:A:391:CYS:H	1.79	0.47
1:A:785:VAL:HG12	1:A:787:GLN:H	1.79	0.47
1:A:1115:ILE:HG22	1:A:1137:VAL:HG23	1.97	0.47

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:C:483:VAL:HG12	1:C:484:LYS:HG3	1.96	0.47
1:A:805:ILE:HD12	1:A:1054:GLN:HE21	1.80	0.47
1:A:999:GLY:O	1:A:1002:GLN:NE2	2.48	0.47
1:C:994:ASP:HA	1:C:997:ILE:HG12	1.97	0.47
1:B:435:ALA:HB2	1:B:510:VAL:HG22	1.97	0.47
1:A:884:SER:OG	1:A:887:THR:OG1	2.27	0.47
1:A:1050:MET:HE2	1:A:1050:MET:HB3	1.78	0.47
1:B:52:GLN:OE1	1:B:274:THR:OG1	2.26	0.47
1:C:34:ARG:HH22	1:C:220:PHE:HA	1.79	0.47
1:A:206:LYS:HB2	1:A:223:LEU:HA	1.96	0.47
1:B:620:VAL:HG11	1:B:642:VAL:HG11	1.97	0.47
1:C:56:LEU:HD22	1:C:91:TYR:CD1	2.50	0.47
1:A:310:LYS:HG3	1:A:600:PRO:HA	1.97	0.47
1:A:316:SER:OG	1:A:317:ASN:N	2.48	0.47
1:A:454:ARG:NH2	1:A:469:SER:O	2.40	0.47
1:A:567:ARG:HD3	1:A:571:ASP:HA	1.96	0.47
1:C:729:VAL:HG12	1:C:1059:GLY:HA2	1.97	0.47
1:C:738:CYS:HB2	1:C:763:LEU:HD11	1.96	0.47
1:A:291:CYS:HB3	1:A:301:CYS:HB2	1.31	0.47
1:B:319:ARG:HH22	1:C:745:ASP:H	1.62	0.47
1:C:483:VAL:HG12	1:C:484:LYS:HG3	1.96	0.47
1:A:390:LEU:HD12	1:A:391:CYS:H	1.79	0.47
1:A:431:GLY:HA2	1:A:515:PHE:CE2	2.50	0.47
1:A:978:ASN:HA	1:A:981:LEU:HG	1.97	0.47
1:B:1045:LYS:HA	1:B:1045:LYS:HD3	1.82	0.47
1:C:503:VAL:HA	1:C:506:GLN:HG3	1.97	0.47
1:B:596:SER:HB2	1:B:611:LEU:HB3	1.96	0.47
1:A:315:THR:OG1	1:A:316:SER:N	2.48	0.47
1:A:701:VAL:HG13	1:B:787:GLN:HG2	1.95	0.47
1:A:978:ASN:HA	1:A:981:LEU:HG	1.97	0.47
1:B:442:ASP:OD2	1:B:509:ARG:NE	2.45	0.47
1:A:785:VAL:HG12	1:A:787:GLN:H	1.79	0.47
1:A:985:ASP:OD2	1:A:987:PRO:HD2	2.15	0.47
1:A:1129:VAL:HG12	1:B:917:TYR:HB3	1.95	0.47
1:C:365:TYR:O	1:C:369:TYR:HB2	2.15	0.47
1:C:976:VAL:HG13	1:C:979:ASP:HB3	1.97	0.47
1:A:189:LEU:HD22	1:A:217:PRO:HG2	1.96	0.47
1:B:574:ASP:OD1	1:B:574:ASP:N	2.46	0.47
1:B:1115:ILE:HG22	1:B:1137:VAL:HG23	1.97	0.47
1:C:1050:MET:HE3	1:C:1050:MET:HB3	1.78	0.47
1:C:787:GLN:OE1	1:C:787:GLN:N	2.48	0.47

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:C:598:ILE:HB	1:C:609:ALA:HB3	1.96	0.47
1:A:905:ARG:NH1	1:A:1049:LEU:O	2.48	0.47
1:B:386:LYS:NZ	1:C:985:ASP:OD2	2.48	0.47
1:C:762:GLN:HA	1:C:765:ARG:HG2	1.96	0.47
1:B:598:ILE:HB	1:B:609:ALA:HB3	1.97	0.47
1:C:58:PHE:HD2	1:C:290:ASP:HB2	1.79	0.47
1:A:599:THR:HB	1:A:608:VAL:HG23	1.95	0.47
1:B:287:ASP:OD1	1:B:288:ALA:N	2.48	0.47
1:B:912:THR:O	1:B:915:VAL:HG12	2.15	0.47
1:C:153:MET:HE1	1:C:155:SER:HB3	1.97	0.47
1:B:1139:ASP:HB3	1:B:1142:GLN:HG3	1.97	0.47
1:B:177:MET:HE1	1:B:190:ARG:HH22	1.79	0.47
1:B:858:LEU:HD13	1:B:959:LEU:HD12	1.96	0.47
1:C:877:LEU:HG	1:C:1053:PRO:HG2	1.97	0.47
1:A:369:TYR:OH	1:C:415:THR:OG1	2.26	0.47
1:B:442:ASP:OD2	1:B:509:ARG:NE	2.43	0.47
1:A:676:ILE:HD12	1:A:677:PRO:HD2	1.96	0.47
1:B:58:PHE:HD2	1:B:290:ASP:HB2	1.79	0.47
1:C:1115:ILE:HG22	1:C:1137:VAL:HG23	1.96	0.47
1:B:421:TYR:HE2	1:B:457:ARG:H	1.63	0.47
1:B:761:THR:O	1:B:765:ARG:HG2	2.15	0.47
1:B:770:ILE:HD11	1:B:1012:LEU:HD22	1.97	0.47
1:C:401:VAL:HA	1:C:509:ARG:HA	1.97	0.47
1:C:664:ILE:HD12	1:C:665:PRO:HD2	1.96	0.47
1:C:773:GLU:OE2	1:C:1019:ARG:NE	2.48	0.47
1:C:34:ARG:NH2	1:C:221:SER:H	2.13	0.47
1:C:58:PHE:HD2	1:C:290:ASP:HB2	1.80	0.47
1:B:433:VAL:HA	1:B:512:VAL:HG23	1.96	0.47
1:B:722:VAL:HG22	1:B:930:ALA:HB1	1.97	0.47
1:B:751:ASN:HA	1:B:754:LEU:HG	1.97	0.47
1:A:360:ASN:H	1:A:523:THR:HB	1.80	0.47
1:A:34:ARG:NH2	1:A:221:SER:H	2.14	0.47
1:B:91:TYR:N	1:B:268:GLY:O	2.37	0.47
1:B:701:VAL:HG13	1:C:787:GLN:HG3	1.96	0.47
1:C:40:ASP:N	1:C:40:ASP:OD1	2.44	0.47
1:C:532:ASN:N	1:C:532:ASN:OD1	2.46	0.47
1:A:738:CYS:HB3	1:A:763:LEU:HD21	1.97	0.47
1:C:291:CYS:HB3	1:C:301:CYS:HB2	1.37	0.47
1:C:722:VAL:HG22	1:C:930:ALA:HB1	1.96	0.47
1:C:931:ILE:HA	1:C:934:ILE:HG22	1.97	0.47
1:A:574:ASP:OD1	1:A:574:ASP:N	2.48	0.47

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:B:611:LEU:HD13	1:B:650:LEU:HD22	1.96	0.47
1:C:642:VAL:HG22	1:C:651:ILE:HG22	1.97	0.47
1:B:332:ILE:HG13	1:B:362:VAL:HG22	1.97	0.47
1:B:1115:ILE:HG22	1:B:1137:VAL:HG23	1.96	0.47
1:C:435:ALA:HB2	1:C:510:VAL:HG12	1.96	0.47
1:A:295:PRO:HA	1:A:298:GLU:HB3	1.97	0.47
1:A:1033:VAL:HG21	1:A:1053:PRO:HG3	1.97	0.47
1:A:719:THR:HG23	1:A:1070:ALA:HB2	1.97	0.47
1:A:919:ASN:HD22	1:A:922:LEU:HD21	1.80	0.47
1:B:189:LEU:HD12	1:B:217:PRO:HG2	1.97	0.47
1:C:300:LYS:NZ	1:C:306:PHE:O	2.47	0.47
1:A:791:THR:HG21	1:A:806:LEU:HD11	1.96	0.47
1:B:398:ASP:HB2	1:B:512:VAL:HG12	1.96	0.47
1:B:573:THR:HG22	1:B:587:ILE:HD13	1.97	0.47
1:B:775:ASP:HB3	1:B:864:LEU:HD13	1.96	0.47
1:C:390:LEU:HD12	1:C:391:CYS:H	1.80	0.47
1:A:732:THR:HG23	1:A:955:ASN:HD22	1.80	0.46
1:B:443:SER:HB3	1:B:499:PRO:HG3	1.97	0.46
1:C:18:PHE:HB2	1:C:21:ARG:HB2	1.98	0.46
1:C:96:GLU:OE1	1:C:96:GLU:N	2.48	0.46
1:A:105:ILE:HG23	1:A:239:GLN:HB3	1.96	0.46
1:A:1002:GLN:HE21	1:B:1005:GLN:NE2	2.12	0.46
1:B:143:VAL:HG11	1:B:152:TRP:CE3	2.51	0.46
1:B:576:VAL:HG12	1:B:587:ILE:HD11	1.96	0.46
1:A:330:PRO:HD3	1:A:544:ASN:HD21	1.80	0.46
1:A:788:ILE:HG23	1:A:876:ALA:HB2	1.95	0.46
1:C:390:LEU:HD12	1:C:391:CYS:H	1.80	0.46
1:A:353:TRP:HH2	1:A:464:PHE:HA	1.79	0.46
1:A:1040:VAL:HG11	1:B:1035:GLY:HA3	1.97	0.46
1:A:1105:THR:HG21	1:A:1110:TYR:HD1	1.79	0.46
1:B:912:THR:O	1:B:915:VAL:HG12	2.15	0.46
1:C:976:VAL:HG13	1:C:979:ASP:HB3	1.97	0.46
1:C:877:LEU:O	1:C:881:THR:OG1	2.28	0.46
1:A:393:THR:HA	1:A:522:ALA:HA	1.97	0.46
1:B:722:VAL:HG12	1:B:1065:VAL:HG22	1.97	0.46
1:C:390:LEU:HD12	1:C:391:CYS:H	1.79	0.46
1:B:135:PHE:HD2	1:B:160:TYR:HE1	1.63	0.46
1:B:866:THR:O	1:B:870:ILE:HG12	2.15	0.46
1:A:483:VAL:HG12	1:A:484:LYS:HG3	1.98	0.46
1:B:1013:ILE:HD13	1:C:1012:LEU:HB3	1.97	0.46
1:C:393:THR:OG1	1:C:394:ASN:OD1	2.32	0.46

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:34:ARG:HG3	1:A:216:LEU:HD11	1.97	0.46
1:A:712:ILE:HG21	1:A:1096:VAL:HG12	1.97	0.46
1:A:1040:VAL:HG11	1:B:1035:GLY:HA3	1.96	0.46
1:B:858:LEU:HD13	1:B:959:LEU:HD12	1.97	0.46
1:B:39:PRO:HG3	1:B:51:THR:HG21	1.96	0.46
1:B:80:ALA:O	1:B:245:HIS:NE2	2.48	0.46
1:A:785:VAL:HG12	1:A:787:GLN:H	1.79	0.46
1:A:919:ASN:HD22	1:A:922:LEU:HD21	1.80	0.46
1:C:326:ILE:HD13	1:C:533:LEU:HD12	1.96	0.46
1:A:759:PHE:HA	1:A:762:GLN:HE21	1.80	0.46
1:A:183:GLN:HG3	1:A:187:LYS:HG3	1.96	0.46
1:A:315:THR:OG1	1:A:316:SER:N	2.48	0.46
1:B:884:SER:OG	1:B:887:THR:OG1	2.26	0.46
1:A:369:TYR:OH	1:C:415:THR:OG1	2.25	0.46
1:A:1030:SER:HA	1:A:1034:LEU:HD23	1.97	0.46
1:B:326:ILE:HD11	1:B:534:VAL:HG22	1.96	0.46
1:C:92:PHE:HB3	1:C:192:PHE:HB2	1.97	0.46
1:A:972:ALA:HB2	1:A:996:LEU:HD21	1.96	0.46
1:B:34:ARG:HH22	1:B:220:PHE:HA	1.80	0.46
1:B:679:GLY:HA2	1:C:864:LEU:HD13	1.97	0.46
1:A:503:VAL:HG21	1:C:503:VAL:HB	1.96	0.46
1:C:1116:THR:OG1	1:C:1118:ASP:OD1	2.32	0.46
1:A:294:ASP:OD1	1:A:297:SER:N	2.45	0.46
1:B:330:PRO:HD3	1:B:579:PRO:HB2	1.97	0.46
1:A:1106:GLN:OE1	1:A:1111:GLU:OE2	2.33	0.46
1:B:34:ARG:NH2	1:B:221:SER:H	2.13	0.46
1:C:785:VAL:HG12	1:C:787:GLN:H	1.80	0.46
1:B:15:CYS:SG	1:B:137:ASN:N	2.84	0.46
1:B:777:ASN:HA	1:B:780:GLU:HG3	1.97	0.46
1:C:951:VAL:HA	1:C:954:GLN:HG3	1.98	0.46
1:A:117:LEU:HD11	1:A:231:ILE:HG12	1.97	0.46
1:B:1115:ILE:HG22	1:B:1137:VAL:HG23	1.96	0.46
1:A:357:ARG:NH2	1:A:396:TYR:OH	2.49	0.46
1:A:598:ILE:HD11	1:A:678:ILE:HD13	1.97	0.46
1:A:1031:GLU:HB3	1:A:1037:SER:HB2	1.95	0.46
1:C:431:GLY:HA2	1:C:515:PHE:CZ	2.51	0.46
1:A:299:THR:HG21	1:A:597:VAL:HG21	1.96	0.46
1:B:330:PRO:HD3	1:B:579:PRO:HB2	1.97	0.46
1:B:715:PRO:HD3	1:C:894:LEU:HD11	1.96	0.46
1:C:106:PHE:HD2	1:C:117:LEU:HD23	1.80	0.46
1:C:426:PRO:HB3	1:C:464:PHE:HB2	1.98	0.46

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:C:738:CYS:HB3	1:C:760:CYS:HB2	1.64	0.46
1:A:15:CYS:SG	1:A:137:ASN:N	2.86	0.46
1:B:37:TYR:OH	1:B:53:ASP:OD1	2.33	0.46
1:C:989:ALA:O	1:C:993:ILE:HG12	2.16	0.46
1:B:334:ASN:ND2	1:B:360:ASN:O	2.40	0.46
1:A:425:LEU:HD12	1:A:429:PHE:CG	2.51	0.46
1:A:688:THR:HA	1:A:690:GLN:HG2	1.97	0.46
1:C:871:ALA:HA	1:C:874:THR:HG22	1.96	0.46
1:A:989:ALA:O	1:A:993:ILE:HG12	2.15	0.46
1:C:606:ASN:O	1:C:606:ASN:ND2	2.37	0.46
1:B:1123:SER:OG	1:C:914:ASN:OD1	2.20	0.46
1:A:730:SER:OG	1:A:731:MET:N	2.46	0.46
1:A:796:ASP:OD1	1:A:796:ASP:N	2.48	0.46
1:B:129:LYS:NZ	1:B:168:PHE:O	2.40	0.46
1:C:596:SER:HB2	1:C:611:LEU:HB3	1.98	0.46
1:A:712:ILE:HD11	1:B:896:ILE:HD12	1.97	0.46
1:C:406:GLU:N	1:C:406:GLU:OE2	2.49	0.46
1:B:728:PRO:HD3	1:B:947:LYS:HG3	1.97	0.46
1:C:1039:ARG:HG2	1:C:1042:PHE:HB2	1.97	0.46
1:A:731:MET:H	1:A:774:GLN:NE2	2.13	0.46
1:B:330:PRO:HD3	1:B:579:PRO:HB2	1.97	0.46
1:C:390:LEU:HD12	1:C:391:CYS:H	1.79	0.46
1:C:276:LEU:HD21	1:C:304:LYS:HA	1.98	0.46
1:C:720:ILE:HD11	1:C:1065:VAL:HG12	1.96	0.46
1:A:108:THR:O	1:A:237:ARG:NH2	2.49	0.46
1:C:326:ILE:HD11	1:C:534:VAL:HG22	1.98	0.46
1:C:785:VAL:HG12	1:C:787:GLN:H	1.80	0.46
1:C:788:ILE:HG13	1:C:876:ALA:HB2	1.98	0.46
1:A:310:LYS:HG3	1:A:600:PRO:HA	1.97	0.46
1:A:34:ARG:NH2	1:A:221:SER:H	2.14	0.46
1:A:796:ASP:OD1	1:A:796:ASP:N	2.49	0.46
1:B:1028:LYS:NZ	1:B:1042:PHE:O	2.44	0.46
1:C:355:ARG:NH2	1:C:398:ASP:OD2	2.49	0.46
1:C:483:VAL:HG12	1:C:484:LYS:HG3	1.96	0.46
1:C:1115:ILE:HG22	1:C:1137:VAL:HG23	1.98	0.46
1:A:756:TYR:HB3	1:A:759:PHE:CD2	2.50	0.46
1:B:332:ILE:HG13	1:B:362:VAL:HG22	1.97	0.46
1:B:619:GLU:N	1:B:619:GLU:OE1	2.49	0.46
1:C:87:ASN:OD1	1:C:87:ASN:N	2.46	0.46
1:C:183:GLN:HG3	1:C:187:LYS:HG3	1.97	0.46
1:A:222:ALA:HB2	1:A:285:ILE:HB	1.98	0.46

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:796:ASP:OD1	1:A:796:ASP:N	2.49	0.46
1:C:31:SER:O	1:C:31:SER:OG	2.31	0.46
1:C:617:CYS:N	1:C:649:CYS:SG	2.87	0.46
1:B:85:PRO:HA	1:B:237:ARG:HD2	1.97	0.46
1:B:564:GLN:HG2	1:B:577:ARG:HD2	1.98	0.46
1:C:303:LEU:HD23	1:C:308:VAL:HG22	1.98	0.46
1:A:954:GLN:HG3	1:A:1014:ARG:HE	1.79	0.46
1:B:58:PHE:HD2	1:B:290:ASP:HB2	1.81	0.46
1:B:296:LEU:O	1:B:299:THR:OG1	2.31	0.46
1:A:87:ASN:OD1	1:A:87:ASN:N	2.49	0.46
1:A:1010:GLN:OE1	1:A:1014:ARG:NH2	2.49	0.46
1:C:966:LEU:HD13	1:C:1000:ARG:HH22	1.79	0.46
1:C:1028:LYS:HB3	1:C:1062:PHE:HE1	1.80	0.46
1:B:562:PHE:CZ	1:C:225:PRO:HG2	2.51	0.46
1:B:565:PHE:O	1:C:43:PHE:HB3	2.15	0.46
1:C:770:ILE:HA	1:C:773:GLU:HG3	1.96	0.46
1:C:105:ILE:HG23	1:C:239:GLN:HB3	1.98	0.46
1:C:130:VAL:HG21	1:C:231:ILE:HD11	1.98	0.46
1:C:393:THR:OG1	1:C:394:ASN:N	2.48	0.46
1:A:688:THR:HA	1:A:690:GLN:HG2	1.98	0.46
1:B:353:TRP:HH2	1:B:464:PHE:HA	1.80	0.46
1:A:403:ARG:NH2	1:B:372:ALA:O	2.33	0.46
1:A:564:GLN:HG2	1:A:565:PHE:CD1	2.50	0.46
1:A:792:PRO:HG3	1:C:707:TYR:HB3	1.98	0.46
1:B:33:THR:OG1	1:B:34:ARG:NH2	2.47	0.46
1:C:93:ALA:HB3	1:C:266:TYR:HB2	1.97	0.46
1:B:719:THR:HG23	1:B:1068:VAL:HB	1.96	0.46
1:A:1031:GLU:OE1	1:C:1039:ARG:HG3	2.14	0.46
1:C:807:PRO:HG2	1:C:875:SER:HB2	1.96	0.46
1:C:951:VAL:HA	1:C:954:GLN:NE2	2.31	0.46
1:B:365:TYR:HE2	1:B:387:LEU:HD12	1.80	0.46
1:A:882:ILE:HD12	1:A:882:ILE:HA	1.79	0.46
1:B:767:LEU:HD21	1:B:1008:VAL:HG12	1.98	0.46
1:B:777:ASN:HA	1:B:780:GLU:HG3	1.97	0.46
1:A:226:LEU:HD23	1:A:227:VAL:HG12	1.97	0.46
1:A:222:ALA:HB2	1:A:285:ILE:HB	1.98	0.46
1:A:289:VAL:HG13	1:A:297:SER:HB3	1.97	0.46
1:B:858:LEU:HD23	1:B:959:LEU:HD12	1.98	0.46
1:B:287:ASP:OD1	1:B:288:ALA:N	2.48	0.46
1:B:751:ASN:HA	1:B:754:LEU:HG	1.97	0.46
1:C:34:ARG:NH2	1:C:221:SER:H	2.12	0.46

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:390:LEU:HD12	1:A:391:CYS:H	1.80	0.46
1:B:722:VAL:HG22	1:B:930:ALA:HB1	1.98	0.46
1:C:330:PRO:HD3	1:C:579:PRO:HB2	1.97	0.46
1:A:58:PHE:HD2	1:A:290:ASP:HB2	1.80	0.46
1:A:796:ASP:OD1	1:A:796:ASP:N	2.49	0.46
1:C:398:ASP:HB2	1:C:512:VAL:HG22	1.97	0.46
1:C:989:ALA:O	1:C:993:ILE:HD12	2.16	0.46
1:A:562:PHE:CE1	1:B:225:PRO:HG2	2.51	0.46
1:B:102:ARG:HD2	1:B:141:LEU:HD22	1.98	0.46
1:A:1105:THR:HG21	1:A:1110:TYR:HD1	1.80	0.46
1:B:48:LEU:HD12	1:B:276:LEU:HD11	1.97	0.46
1:B:680:ALA:H	1:C:864:LEU:HA	1.81	0.46
1:A:462:LYS:HE2	1:A:465:GLU:HG3	1.96	0.46
1:C:58:PHE:HD2	1:C:290:ASP:HB2	1.80	0.46
1:C:439:ASN:ND2	1:C:506:GLN:OE1	2.49	0.46
1:B:58:PHE:HD2	1:B:290:ASP:HB2	1.81	0.46
1:C:48:LEU:HB3	1:C:276:LEU:HD11	1.98	0.46
1:C:315:THR:OG1	1:C:316:SER:N	2.48	0.46
1:C:390:LEU:HD12	1:C:391:CYS:H	1.80	0.46
1:B:390:LEU:HD12	1:B:391:CYS:H	1.80	0.46
1:B:978:ASN:HA	1:B:981:LEU:HG	1.97	0.46
1:A:825:LYS:HZ2	1:A:945:LEU:HG	1.81	0.46
1:A:37:TYR:OH	1:A:54:LEU:O	2.25	0.46
1:A:457:ARG:NH2	1:A:467:ASP:OD2	2.45	0.46
1:C:902:MET:HB3	1:C:916:LEU:HD11	1.98	0.46
1:A:796:ASP:OD1	1:A:796:ASP:N	2.49	0.46
1:B:566:GLY:HA2	1:C:43:PHE:HB3	1.98	0.46
1:B:983:ARG:HG3	1:B:984:LEU:HD23	1.97	0.46
1:C:543:PHE:HE1	1:C:552:LEU:HD11	1.80	0.46
1:B:1030:SER:HA	1:B:1034:LEU:HD23	1.98	0.46
1:C:99:ASN:HD21	1:C:177:MET:HE2	1.81	0.46
1:A:295:PRO:HA	1:A:298:GLU:HB2	1.98	0.46
1:A:1052:PHE:HB2	1:A:1063:LEU:HD12	1.98	0.46
1:C:532:ASN:OD1	1:C:533:LEU:N	2.49	0.46
1:B:41:LYS:HD3	1:B:225:PRO:HG2	1.97	0.46
1:C:49:HIS:CE1	1:C:51:THR:HG22	2.51	0.46
1:C:971:GLY:HA3	1:C:995:ARG:HH21	1.81	0.46
1:A:498:GLN:H	1:A:501:TYR:HE1	1.62	0.46
1:A:1031:GLU:CD	1:C:1039:ARG:HG3	2.41	0.46
1:B:775:ASP:HB3	1:B:864:LEU:HD13	1.98	0.46
1:B:984:LEU:HD12	1:B:989:ALA:HA	1.97	0.46

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:326:ILE:HD11	1:A:534:VAL:HG22	1.96	0.46
1:B:435:ALA:HB2	1:B:510:VAL:HG22	1.96	0.46
1:C:989:ALA:O	1:C:993:ILE:HD12	2.16	0.46
1:A:395:VAL:HG22	1:A:515:PHE:HB3	1.98	0.46
1:A:688:THR:HA	1:A:690:GLN:HG2	1.98	0.46
1:C:560:LEU:HD23	1:C:561:PRO:HD2	1.98	0.46
1:A:989:ALA:O	1:A:993:ILE:HG12	2.15	0.46
1:B:729:VAL:HG21	1:B:781:VAL:HG11	1.98	0.46
1:C:394:ASN:OD1	1:C:394:ASN:N	2.47	0.46
1:C:557:LYS:HE2	1:C:559:PHE:HE2	1.81	0.46
1:A:358:ILE:HB	1:A:395:VAL:HG23	1.98	0.46
1:C:385:THR:OG1	1:C:386:LYS:N	2.48	0.46
1:C:762:GLN:HA	1:C:765:ARG:HE	1.80	0.46
1:C:896:ILE:HD12	1:C:897:PRO:HD2	1.96	0.46
1:C:931:ILE:HA	1:C:934:ILE:HG22	1.97	0.46
1:C:483:VAL:HG12	1:C:484:LYS:HG3	1.98	0.46
1:B:40:ASP:OD1	1:B:42:VAL:HG12	2.16	0.46
1:B:977:LEU:HD22	1:B:993:ILE:HG22	1.98	0.46
1:A:205:SER:HB3	1:A:226:LEU:HD22	1.97	0.46
1:B:882:ILE:HD12	1:B:882:ILE:HA	1.71	0.46
1:A:295:PRO:HA	1:A:298:GLU:HB3	1.97	0.46
1:A:326:ILE:HD11	1:A:534:VAL:HG22	1.97	0.46
1:B:319:ARG:HH22	1:C:745:ASP:H	1.64	0.46
1:C:576:VAL:HG12	1:C:587:ILE:HD11	1.96	0.46
1:B:326:ILE:HD13	1:B:533:LEU:HD12	1.98	0.46
1:C:1050:MET:HE2	1:C:1050:MET:HB3	1.77	0.46
1:A:390:LEU:HD12	1:A:391:CYS:H	1.81	0.46
1:C:240:THR:OG1	1:C:245:HIS:NE2	2.41	0.46
1:B:33:THR:OG1	1:B:219:GLY:O	2.27	0.46
1:B:796:ASP:OD1	1:B:796:ASP:N	2.48	0.46
1:A:1039:ARG:NE	1:B:1031:GLU:OE2	2.44	0.46
1:B:425:LEU:HD23	1:B:429:PHE:CD1	2.51	0.46
1:B:44:ARG:HH12	1:B:49:HIS:CD2	2.34	0.46
1:A:642:VAL:HG22	1:A:651:ILE:HG22	1.98	0.46
1:C:398:ASP:HB2	1:C:512:VAL:HG22	1.98	0.46
1:C:452:LEU:HA	1:C:494:SER:HA	1.98	0.46
1:A:1155:TYR:O	1:A:1159:HIS:ND1	2.34	0.46
1:C:807:PRO:HA	1:C:816:SER:HA	1.96	0.46
1:A:809:PRO:HA	1:A:814:LYS:HD2	1.97	0.46
1:B:442:ASP:OD2	1:B:509:ARG:NE	2.44	0.46
1:A:976:VAL:HG13	1:A:979:ASP:HB3	1.96	0.46

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:C:642:VAL:HG22	1:C:651:ILE:HG22	1.97	0.46
1:B:298:GLU:HB3	1:B:315:THR:HG21	1.98	0.46
1:B:785:VAL:HG12	1:B:787:GLN:H	1.80	0.46
1:C:756:TYR:HB3	1:C:759:PHE:HD2	1.80	0.46
1:C:483:VAL:HG12	1:C:484:LYS:HG3	1.98	0.46
1:B:386:LYS:O	1:B:390:LEU:HB2	2.15	0.46
1:B:429:PHE:HE2	1:B:514:SER:HA	1.80	0.46
1:B:809:PRO:HA	1:B:814:LYS:HD2	1.98	0.46
1:A:133:PHE:CE2	1:A:160:TYR:HB3	2.50	0.46
1:A:728:PRO:HD3	1:A:947:LYS:HG2	1.98	0.46
1:C:483:VAL:HG12	1:C:484:LYS:HG3	1.98	0.46
1:B:403:ARG:NH2	1:C:372:ALA:O	2.37	0.46
1:C:365:TYR:O	1:C:369:TYR:HB2	2.16	0.46
1:C:412:PRO:HD3	1:C:425:LEU:HD23	1.98	0.46
1:A:997:ILE:O	1:A:1001:LEU:HB3	2.16	0.46
1:B:296:LEU:O	1:B:299:THR:OG1	2.29	0.46
1:B:314:GLN:HE22	1:B:594:GLY:HA3	1.81	0.46
1:B:390:LEU:HD12	1:B:391:CYS:H	1.81	0.46
1:C:366:SER:O	1:C:370:ASN:HB2	2.15	0.46
1:B:557:LYS:HB3	1:B:559:PHE:HE2	1.81	0.46
1:C:296:LEU:O	1:C:299:THR:OG1	2.31	0.46
1:A:483:VAL:HG12	1:A:484:LYS:HG3	1.98	0.46
1:A:456:PHE:HB3	1:A:473:TYR:HD1	1.81	0.46
1:A:737:ASP:OD1	1:A:737:ASP:N	2.40	0.46
1:B:37:TYR:HB3	1:B:223:LEU:HB2	1.98	0.46
1:A:882:ILE:HD12	1:A:882:ILE:HA	1.87	0.46
1:B:37:TYR:HB3	1:B:223:LEU:HB2	1.98	0.46
1:B:332:ILE:HG13	1:B:362:VAL:HG22	1.98	0.46
1:A:65:PHE:HE1	1:A:84:LEU:HD11	1.80	0.46
1:B:287:ASP:OD1	1:B:288:ALA:N	2.48	0.46
1:B:966:LEU:HA	1:B:1000:ARG:HH12	1.80	0.46
1:B:877:LEU:O	1:B:881:THR:HG23	2.16	0.46
1:C:393:THR:OG1	1:C:520:ALA:O	2.32	0.46
1:C:948:LEU:HD11	1:C:1059:GLY:HA3	1.96	0.46
1:B:398:ASP:HB2	1:B:512:VAL:HG13	1.98	0.46
1:C:751:ASN:HA	1:C:754:LEU:HG	1.98	0.46
1:A:206:LYS:HB2	1:A:223:LEU:HA	1.97	0.46
1:A:722:VAL:HG22	1:A:930:ALA:HB1	1.98	0.46
1:A:759:PHE:HA	1:A:762:GLN:HG3	1.97	0.46
1:B:105:ILE:HB	1:B:239:GLN:HB2	1.97	0.46
1:B:878:LEU:O	1:B:882:ILE:HG12	2.16	0.46

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:C:777:ASN:HA	1:C:780:GLU:HG3	1.97	0.46
1:B:950:ASP:HA	1:B:953:ASN:ND2	2.30	0.46
1:C:197:ILE:HG13	1:C:198:ASP:OD2	2.16	0.46
1:C:1051:SER:OG	1:C:1064:HIS:ND1	2.35	0.46
1:A:115:GLN:HB2	1:A:233:ILE:HG12	1.97	0.46
1:A:796:ASP:OD1	1:A:796:ASP:N	2.49	0.46
1:C:596:SER:HB2	1:C:611:LEU:HB3	1.97	0.46
1:C:989:ALA:O	1:C:993:ILE:HG12	2.16	0.46
1:C:64:TRP:HE1	1:C:264:ALA:HB1	1.80	0.46
1:C:105:ILE:HG23	1:C:239:GLN:HB3	1.97	0.46
1:A:599:THR:HB	1:A:608:VAL:HG23	1.96	0.46
1:B:408:ARG:HE	1:B:409:GLN:HG3	1.80	0.46
1:C:129:LYS:NZ	1:C:168:PHE:H	2.14	0.46
1:B:406:GLU:HG2	1:B:418:ILE:HG12	1.98	0.46
1:C:106:PHE:HB2	1:C:117:LEU:HB3	1.98	0.46
1:A:386:LYS:NZ	1:B:984:LEU:O	2.47	0.46
1:A:802:PHE:HB3	1:A:806:LEU:HD23	1.98	0.46
1:B:195:LYS:HE3	1:B:202:LYS:HD3	1.98	0.46
1:A:37:TYR:OH	1:A:53:ASP:OD1	2.32	0.46
1:A:326:ILE:HD11	1:A:534:VAL:HG22	1.97	0.46
1:C:358:ILE:HB	1:C:395:VAL:HG23	1.98	0.46
1:C:722:VAL:HG22	1:C:930:ALA:HB1	1.96	0.46
1:B:350:VAL:HA	1:B:400:PHE:HB2	1.98	0.46
1:C:313:TYR:HB2	1:C:597:VAL:HG12	1.98	0.46
1:C:1014:ARG:O	1:C:1018:ILE:HG12	2.16	0.46
1:C:290:ASP:HB3	1:C:293:LEU:HB2	1.97	0.46
1:C:398:ASP:OD2	1:C:423:TYR:OH	2.24	0.46
1:A:214:ARG:O	1:A:214:ARG:NH1	2.49	0.46
1:C:276:LEU:HD13	1:C:301:CYS:HA	1.97	0.46
1:B:751:ASN:HA	1:B:754:LEU:HG	1.97	0.46
1:C:733:LYS:HD3	1:C:771:ALA:HB1	1.97	0.46
1:A:92:PHE:HE1	1:A:94:SER:HB2	1.80	0.46
1:A:328:ARG:NH1	1:A:531:THR:O	2.44	0.46
1:B:287:ASP:OD1	1:B:288:ALA:N	2.49	0.46
1:B:1051:SER:OG	1:B:1064:HIS:ND1	2.39	0.46
1:C:442:ASP:OD2	1:C:509:ARG:NE	2.41	0.46
1:A:985:ASP:O	1:A:989:ALA:CB	2.64	0.46
1:B:216:LEU:HD21	1:B:266:TYR:HE2	1.80	0.46
1:B:513:LEU:HB3	1:B:515:PHE:HE2	1.81	0.46
1:C:338:PHE:HE1	1:C:369:TYR:H	1.63	0.46
1:B:955:ASN:C	1:B:955:ASN:HD22	2.23	0.46

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:C:606:ASN:O	1:C:606:ASN:ND2	2.36	0.46
1:B:1116:THR:OG1	1:B:1118:ASP:OD1	2.32	0.46
1:C:390:LEU:HD12	1:C:391:CYS:H	1.80	0.46
1:C:1073:LYS:HE3	1:C:1075:PHE:HZ	1.81	0.46
1:A:483:VAL:HG12	1:A:484:LYS:HG3	1.98	0.46
1:B:471:GLU:OE2	1:C:113:LYS:NZ	2.44	0.46
1:A:762:GLN:HA	1:A:765:ARG:HH21	1.81	0.46
1:B:950:ASP:OD1	1:B:950:ASP:N	2.47	0.46
1:A:108:THR:O	1:A:237:ARG:NH2	2.49	0.46
1:A:389:ASP:OD1	1:A:528:LYS:NZ	2.47	0.46
1:A:1081:ILE:HD11	1:A:1135:ASN:HB3	1.97	0.46
1:B:503:VAL:HA	1:B:506:GLN:HG3	1.96	0.46
1:B:936:ASP:O	1:B:940:SER:OG	2.24	0.46
1:A:994:ASP:HA	1:A:997:ILE:HG12	1.98	0.46
1:C:302:THR:O	1:C:304:LYS:NZ	2.48	0.46
1:A:100:ILE:HG21	1:A:263:ALA:HB2	1.98	0.46
1:A:332:ILE:HG13	1:A:362:VAL:HG22	1.97	0.46
1:A:486:PHE:O	1:A:487:ASN:ND2	2.49	0.46
1:B:287:ASP:OD1	1:B:288:ALA:N	2.49	0.46
1:C:905:ARG:HE	1:C:1050:MET:HE3	1.80	0.46
1:A:756:TYR:HB3	1:A:759:PHE:HD2	1.81	0.46
1:B:42:VAL:O	1:B:44:ARG:NH1	2.49	0.46
1:B:491:PRO:HG2	1:B:492:LEU:HD22	1.97	0.46
1:C:737:ASP:N	1:C:737:ASP:OD1	2.42	0.46
1:A:100:ILE:HA	1:A:246:ILE:HD12	1.97	0.46
1:C:650:LEU:HD13	1:C:653:ALA:HB3	1.98	0.46
1:B:796:ASP:N	1:B:796:ASP:OD2	2.49	0.46
1:A:869:MET:HE2	1:A:869:MET:HB3	1.92	0.46
1:A:955:ASN:HD22	1:A:955:ASN:HA	1.48	0.46
1:B:366:SER:O	1:B:370:ASN:HB2	2.15	0.46
1:C:785:VAL:HG12	1:C:787:GLN:H	1.80	0.46
1:A:452:LEU:HD23	1:A:492:LEU:HB3	1.97	0.46
1:A:717:ASN:OD1	1:A:718:PHE:N	2.49	0.46
1:A:788:ILE:HB	1:C:702:GLU:HA	1.97	0.46
1:A:976:VAL:HG23	1:A:979:ASP:HB3	1.97	0.46
1:B:778:THR:HA	1:B:781:VAL:HG12	1.98	0.46
1:C:560:LEU:HD12	1:C:561:PRO:HD2	1.98	0.46
1:A:390:LEU:HD12	1:A:391:CYS:H	1.81	0.46
1:C:730:SER:OG	1:C:731:MET:N	2.49	0.46
1:A:168:PHE:HE2	1:C:466:ARG:HH12	1.64	0.46
1:B:984:LEU:HD13	1:B:988:GLU:HG2	1.97	0.46

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:B:775:ASP:HB3	1:B:864:LEU:HD13	1.98	0.46
1:B:442:ASP:OD2	1:B:509:ARG:NE	2.44	0.46
1:B:699:LEU:HG	1:C:873:TYR:HE1	1.81	0.46
1:C:978:ASN:HA	1:C:981:LEU:HG	1.98	0.46
1:A:390:LEU:HD12	1:A:391:CYS:H	1.81	0.46
1:B:287:ASP:OD1	1:B:288:ALA:N	2.49	0.46
1:B:366:SER:O	1:B:370:ASN:HB2	2.15	0.46
1:C:785:VAL:HG12	1:C:787:GLN:H	1.81	0.46
1:C:1002:GLN:O	1:C:1006:THR:HG23	2.16	0.46
1:A:611:LEU:HD13	1:A:650:LEU:HD13	1.97	0.46
1:B:393:THR:OG1	1:B:394:ASN:N	2.49	0.46
1:C:984:LEU:HD23	1:C:988:GLU:HB2	1.97	0.46
1:A:315:THR:OG1	1:A:316:SER:N	2.49	0.46
1:A:533:LEU:HB3	1:A:535:LYS:HZ3	1.81	0.46
1:A:574:ASP:OD2	1:A:574:ASP:N	2.48	0.46
1:A:1031:GLU:OE2	1:C:1039:ARG:HG3	2.16	0.46
1:B:330:PRO:HD3	1:B:579:PRO:HB2	1.97	0.46
1:B:728:PRO:HD3	1:B:947:LYS:HG3	1.98	0.46
1:B:735:SER:HB2	1:B:861:LEU:HD21	1.98	0.46
1:C:532:ASN:OD1	1:C:533:LEU:N	2.49	0.46
1:A:761:THR:O	1:A:765:ARG:HB3	2.16	0.45
1:A:989:ALA:O	1:A:993:ILE:HG13	2.15	0.45
1:B:497:PHE:CE2	1:B:507:PRO:HB3	2.51	0.45
1:C:858:LEU:HD13	1:C:959:LEU:HB3	1.97	0.45
1:C:1067:TYR:OH	1:C:1108:ASN:O	2.34	0.45
1:B:222:ALA:HB2	1:B:285:ILE:HB	1.98	0.45
1:B:532:ASN:OD1	1:B:533:LEU:N	2.49	0.45
1:A:796:ASP:OD1	1:A:796:ASP:N	2.49	0.45
1:A:722:VAL:HG12	1:A:1065:VAL:HG22	1.99	0.45
1:B:984:LEU:HB2	1:B:989:ALA:HB2	1.98	0.45
1:B:1115:ILE:HG22	1:B:1137:VAL:HG23	1.98	0.45
1:C:642:VAL:HG22	1:C:651:ILE:HG22	1.98	0.45
1:C:84:LEU:HD11	1:C:267:VAL:HG11	1.98	0.45
1:A:658:ASN:C	1:A:658:ASN:ND2	2.73	0.45
1:A:1041:ASP:OD1	1:B:1030:SER:HB2	2.16	0.45
1:B:152:TRP:HB3	1:B:179:LEU:HD12	1.98	0.45
1:B:403:ARG:HG2	1:B:406:GLU:OE1	2.16	0.45
1:B:887:THR:HG21	1:B:894:LEU:HB2	1.99	0.45
1:B:973:ILE:HB	1:B:980:ILE:HD11	1.98	0.45
1:C:100:ILE:HA	1:C:246:ILE:HD12	1.98	0.45
1:A:353:TRP:HH2	1:A:464:PHE:HA	1.81	0.45

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:B:483:VAL:HG12	1:B:484:LYS:HG3	1.98	0.45
1:C:532:ASN:OD1	1:C:533:LEU:N	2.49	0.45
1:C:1051:SER:OG	1:C:1064:HIS:ND1	2.35	0.45
1:A:433:VAL:HG12	1:A:512:VAL:HG22	1.97	0.45
1:B:287:ASP:OD1	1:B:288:ALA:N	2.49	0.45
1:C:1096:VAL:HG11	1:C:1105:THR:HG22	1.97	0.45
1:C:719:THR:HG23	1:C:1070:ALA:HB2	1.98	0.45
1:C:555:SER:HB2	1:C:586:ASP:OD1	2.16	0.45
1:C:596:SER:HB2	1:C:611:LEU:HB3	1.98	0.45
1:C:606:ASN:O	1:C:606:ASN:ND2	2.37	0.45
1:C:735:SER:HB3	1:C:861:LEU:HD21	1.97	0.45
1:C:231:ILE:HG13	1:C:233:ILE:HG22	1.98	0.45
1:C:483:VAL:HG12	1:C:484:LYS:HG3	1.97	0.45
1:A:360:ASN:H	1:A:523:THR:HB	1.81	0.45
1:B:909:ILE:HG23	1:B:911:VAL:HG12	1.98	0.45
1:B:971:GLY:HA3	1:B:995:ARG:HH12	1.81	0.45
1:C:1033:VAL:HG21	1:C:1053:PRO:HG3	1.98	0.45
1:A:737:ASP:OD1	1:A:740:MET:HB3	2.15	0.45
1:B:57:PRO:HG3	1:B:273:ARG:HG3	1.98	0.45
1:B:330:PRO:HD3	1:B:579:PRO:HB2	1.97	0.45
1:A:722:VAL:HG12	1:A:1065:VAL:HG22	1.98	0.45
1:B:809:PRO:HA	1:B:814:LYS:HD2	1.97	0.45
1:B:984:LEU:HD12	1:B:988:GLU:HB2	1.97	0.45
1:B:679:GLY:HA2	1:C:864:LEU:HD13	1.97	0.45
1:C:994:ASP:O	1:C:998:THR:HG23	2.16	0.45
1:A:787:GLN:HG2	1:C:701:VAL:HG13	1.99	0.45
1:A:791:THR:HG21	1:A:806:LEU:HD11	1.97	0.45
1:C:751:ASN:HA	1:C:754:LEU:HG	1.98	0.45
1:A:195:LYS:HE3	1:A:202:LYS:HG3	1.97	0.45
1:A:735:SER:O	1:A:859:THR:OG1	2.30	0.45
1:B:401:VAL:HG22	1:B:509:ARG:HG2	1.96	0.45
1:B:912:THR:O	1:B:915:VAL:HG12	2.16	0.45
1:C:483:VAL:HG12	1:C:484:LYS:HG3	1.98	0.45
1:C:871:ALA:HA	1:C:874:THR:HG22	1.98	0.45
1:C:426:PRO:HD3	1:C:463:PRO:HB3	1.98	0.45
1:B:37:TYR:OH	1:B:53:ASP:OD1	2.33	0.45
1:C:34:ARG:HH21	1:C:221:SER:HB3	1.81	0.45
1:C:287:ASP:OD1	1:C:288:ALA:N	2.49	0.45
1:A:347:PHE:CE1	1:A:509:ARG:HD3	2.51	0.45
1:B:1086:LYS:HE3	1:B:1122:VAL:HG11	1.98	0.45
1:C:735:SER:HB3	1:C:861:LEU:HD21	1.98	0.45

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:C:871:ALA:HA	1:C:874:THR:HG22	1.97	0.45
1:A:989:ALA:O	1:A:993:ILE:HG12	2.16	0.45
1:A:353:TRP:HH2	1:A:464:PHE:HA	1.81	0.45
1:B:978:ASN:HA	1:B:981:LEU:HG	1.99	0.45
1:A:483:VAL:HG12	1:A:484:LYS:HG3	1.99	0.45
1:B:442:ASP:OD2	1:B:509:ARG:NE	2.45	0.45
1:C:366:SER:O	1:C:370:ASN:HB2	2.16	0.45
1:B:722:VAL:HG22	1:B:930:ALA:HB1	1.97	0.45
1:C:767:LEU:HA	1:C:770:ILE:HG12	1.97	0.45
1:C:905:ARG:NH1	1:C:1036:GLN:HB2	2.31	0.45
1:C:871:ALA:HA	1:C:874:THR:HG22	1.97	0.45
1:A:1033:VAL:HA	1:A:1051:SER:HB2	1.99	0.45
1:B:106:PHE:HB2	1:B:117:LEU:HB3	1.99	0.45
1:C:318:PHE:HB3	1:C:591:SER:O	2.16	0.45
1:A:108:THR:OG1	1:A:234:ASN:O	2.33	0.45
1:B:204:TYR:CE1	1:B:225:PRO:HB3	2.51	0.45
1:B:326:ILE:HD13	1:B:533:LEU:HD12	1.98	0.45
1:B:483:VAL:HG12	1:B:484:LYS:HG3	1.98	0.45
1:B:722:VAL:HG12	1:B:1065:VAL:HG22	1.99	0.45
1:B:947:LYS:HE3	1:B:947:LYS:HB2	1.87	0.45
1:C:99:ASN:HB3	1:C:102:ARG:NH2	2.32	0.45
1:A:574:ASP:O	1:A:587:ILE:N	2.37	0.45
1:A:884:SER:OG	1:A:887:THR:OG1	2.30	0.45
1:B:877:LEU:O	1:B:881:THR:HG23	2.17	0.45
1:C:57:PRO:HB2	1:C:60:SER:HB3	1.98	0.45
1:A:483:VAL:HG12	1:A:484:LYS:HG3	1.99	0.45
1:B:674:CYS:HB2	1:B:697:MET:HG3	1.99	0.45
1:B:1115:ILE:HG22	1:B:1137:VAL:HG23	1.98	0.45
1:C:319:ARG:NH2	1:C:590:CYS:SG	2.89	0.45
1:B:1013:ILE:HD13	1:C:1012:LEU:HG	1.97	0.45
1:C:121:ASN:HA	1:C:126:VAL:HA	1.98	0.45
1:C:287:ASP:OD1	1:C:288:ALA:N	2.50	0.45
1:B:113:LYS:HG2	1:B:114:THR:HG23	1.99	0.45
1:C:918:GLU:N	1:C:918:GLU:OE1	2.49	0.45
1:A:569:ILE:HG13	1:B:47:VAL:HG23	1.99	0.45
1:B:966:LEU:HA	1:B:1000:ARG:HH12	1.80	0.45
1:A:476:GLY:H	1:A:487:ASN:HB3	1.81	0.45
1:A:567:ARG:HG2	1:B:42:VAL:HG13	1.98	0.45
1:B:954:GLN:OE1	1:B:1014:ARG:NE	2.49	0.45
1:C:918:GLU:OE2	1:C:918:GLU:N	2.49	0.45
1:B:912:THR:O	1:B:915:VAL:HG12	2.16	0.45

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:642:VAL:HG22	1:A:651:ILE:HG22	1.99	0.45
1:A:712:ILE:HD13	1:A:712:ILE:HA	1.74	0.45
1:B:822:LEU:HG	1:B:945:LEU:HD21	1.98	0.45
1:A:177:MET:HE1	1:A:190:ARG:HH12	1.82	0.45
1:A:950:ASP:N	1:A:950:ASP:OD1	2.46	0.45
1:B:751:ASN:HA	1:B:754:LEU:HG	1.97	0.45
1:C:483:VAL:HG12	1:C:484:LYS:HG3	1.98	0.45
1:C:642:VAL:HG22	1:C:651:ILE:HG22	1.98	0.45
1:B:105:ILE:HD12	1:B:241:LEU:HD21	1.98	0.45
1:A:115:GLN:HB2	1:A:233:ILE:HG12	1.96	0.45
1:A:984:LEU:O	1:C:386:LYS:NZ	2.41	0.45
1:B:712:ILE:HG12	1:B:1094:VAL:HG11	1.98	0.45
1:A:599:THR:HB	1:A:608:VAL:HG23	1.97	0.45
1:B:84:LEU:HB2	1:B:237:ARG:HH12	1.81	0.45
1:A:115:GLN:HE21	1:A:233:ILE:HD13	1.82	0.45
1:B:92:PHE:HE2	1:B:265:TYR:HB2	1.81	0.45
1:B:565:PHE:O	1:C:43:PHE:HB3	2.17	0.45
1:B:777:ASN:HA	1:B:780:GLU:HG3	1.97	0.45
1:A:687:GLN:HG2	1:A:693:ILE:HD13	1.97	0.45
1:B:978:ASN:HA	1:B:981:LEU:HG	1.97	0.45
1:A:787:GLN:HE21	1:C:703:ASN:HB3	1.82	0.45
1:B:327:VAL:O	1:B:531:THR:OG1	2.23	0.45
1:B:722:VAL:HG22	1:B:930:ALA:HB1	1.98	0.45
1:C:320:VAL:HA	1:C:592:PHE:HB2	1.98	0.45
1:C:423:TYR:HE2	1:C:425:LEU:HD23	1.82	0.45
1:A:776:LYS:HZ3	1:A:1019:ARG:HH21	1.63	0.45
1:B:361:CYS:H	1:B:522:ALA:HB1	1.81	0.45
1:B:699:LEU:HD21	1:C:869:MET:HB3	1.98	0.45
1:B:773:GLU:OE2	1:B:774:GLN:NE2	2.46	0.45
1:B:866:THR:H	1:B:869:MET:HE2	1.81	0.45
1:C:976:VAL:HG13	1:C:979:ASP:HB3	1.98	0.45
1:B:202:LYS:HZ1	1:B:225:PRO:HB3	1.81	0.45
1:B:775:ASP:HB3	1:B:864:LEU:HD13	1.99	0.45
1:B:1115:ILE:HG22	1:B:1137:VAL:HG23	1.99	0.45
1:C:1054:GLN:HB2	1:C:1061:VAL:HG13	1.99	0.45
1:B:350:VAL:HG21	1:B:418:ILE:HD11	1.99	0.45
1:C:1031:GLU:HG3	1:C:1039:ARG:HH21	1.82	0.45
1:A:128:ILE:HB	1:A:170:TYR:HD2	1.81	0.45
1:B:175:PHE:HE2	1:B:177:MET:HE2	1.81	0.45
1:A:457:ARG:NH2	1:A:467:ASP:OD2	2.39	0.45
1:A:796:ASP:OD1	1:A:796:ASP:N	2.49	0.45

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:B:598:ILE:HB	1:B:609:ALA:HB3	1.97	0.45
1:C:37:TYR:OH	1:C:54:LEU:O	2.35	0.45
1:C:1096:VAL:HG11	1:C:1105:THR:HG22	1.97	0.45
1:B:15:CYS:SG	1:B:137:ASN:N	2.87	0.45
1:A:574:ASP:OD1	1:A:574:ASP:N	2.48	0.45
1:A:987:PRO:HD2	1:A:988:GLU:OE1	2.16	0.45
1:B:48:LEU:HD12	1:B:276:LEU:HD11	1.98	0.45
1:B:722:VAL:HG12	1:B:1065:VAL:HG22	1.99	0.45
1:C:785:VAL:HG12	1:C:787:GLN:H	1.81	0.45
1:A:1050:MET:HE2	1:A:1050:MET:HB3	1.73	0.45
1:B:353:TRP:HH2	1:B:464:PHE:HD1	1.64	0.45
1:C:299:THR:HG22	1:C:597:VAL:HG21	1.97	0.45
1:C:326:ILE:HD13	1:C:533:LEU:HD12	1.98	0.45
1:A:826:VAL:HB	1:A:1057:PRO:HG2	1.99	0.45
1:C:390:LEU:HD12	1:C:391:CYS:H	1.80	0.45
1:A:984:LEU:HD13	1:A:988:GLU:HG2	1.99	0.45
1:B:931:ILE:HA	1:B:934:ILE:HG22	1.97	0.45
1:A:896:ILE:HD12	1:A:897:PRO:HD2	1.97	0.45
1:B:56:LEU:HD22	1:B:91:TYR:CD2	2.51	0.45
1:B:785:VAL:HG12	1:B:787:GLN:H	1.81	0.45
1:A:220:PHE:HE2	1:A:285:ILE:HG22	1.82	0.45
1:C:825:LYS:HD2	1:C:945:LEU:HD13	1.98	0.45
1:B:206:LYS:HB2	1:B:223:LEU:HA	1.98	0.45
1:B:738:CYS:HB2	1:B:763:LEU:HD21	1.98	0.45
1:B:774:GLN:HA	1:B:777:ASN:ND2	2.32	0.45
1:B:1005:GLN:HA	1:B:1008:VAL:HG12	1.98	0.45
1:C:756:TYR:CE2	1:C:997:ILE:HD12	2.52	0.45
1:C:884:SER:OG	1:C:887:THR:OG1	2.29	0.45
1:C:985:ASP:OD1	1:C:985:ASP:N	2.49	0.45
1:C:931:ILE:HA	1:C:934:ILE:HG22	1.98	0.45
1:A:455:LEU:HD11	1:A:493:GLN:HB3	1.98	0.45
1:B:457:ARG:NH1	1:B:459:SER:O	2.50	0.45
1:B:989:ALA:O	1:B:993:ILE:HG12	2.17	0.45
1:A:989:ALA:O	1:A:993:ILE:HD12	2.16	0.45
1:C:1073:LYS:HE3	1:C:1075:PHE:CZ	2.50	0.45
1:A:825:LYS:HB2	1:A:945:LEU:HD12	1.98	0.45
1:A:353:TRP:HH2	1:A:464:PHE:HA	1.82	0.45
1:B:97:LYS:HG3	1:B:186:PHE:HD1	1.82	0.45
1:C:195:LYS:HD3	1:C:270:LEU:HD12	1.98	0.45
1:C:295:PRO:HG2	1:C:608:VAL:HG11	1.98	0.45
1:A:128:ILE:HB	1:A:170:TYR:HD2	1.81	0.45

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:972:ALA:HB2	1:A:996:LEU:HD21	1.99	0.45
1:A:701:VAL:HG13	1:B:787:GLN:HG2	1.98	0.45
1:B:733:LYS:HB3	1:B:861:LEU:HD23	1.97	0.45
1:C:390:LEU:HD12	1:C:391:CYS:H	1.82	0.45
1:C:52:GLN:OE1	1:C:274:THR:OG1	2.22	0.45
1:C:212:LEU:HD13	1:C:217:PRO:HG3	1.99	0.45
1:C:902:MET:HB3	1:C:916:LEU:HD11	1.98	0.45
1:C:1115:ILE:HG22	1:C:1137:VAL:HG23	1.99	0.45
1:C:105:ILE:HD13	1:C:105:ILE:HA	1.77	0.45
1:C:726:ILE:HG22	1:C:1061:VAL:HG22	1.98	0.45
1:A:84:LEU:HD21	1:A:267:VAL:HG11	1.96	0.45
1:A:299:THR:HG22	1:A:315:THR:HG21	1.97	0.45
1:B:729:VAL:HG21	1:B:781:VAL:HG11	1.98	0.45
1:A:212:LEU:HD13	1:A:217:PRO:HD3	1.99	0.45
1:B:393:THR:OG1	1:B:394:ASN:N	2.49	0.45
1:B:949:GLN:HA	1:B:952:VAL:HG22	1.98	0.45
1:A:756:TYR:HB3	1:A:759:PHE:HD2	1.82	0.45
1:A:1033:VAL:HG11	1:A:1053:PRO:HG3	1.98	0.45
1:C:729:VAL:HG12	1:C:1059:GLY:HA2	1.97	0.45
1:A:796:ASP:OD1	1:A:796:ASP:N	2.50	0.45
1:B:726:ILE:HG22	1:B:1061:VAL:HG22	1.97	0.45
1:C:978:ASN:HA	1:C:981:LEU:HG	1.97	0.45
1:B:712:ILE:HG12	1:B:1094:VAL:HG11	1.99	0.45
1:C:431:GLY:HA2	1:C:515:PHE:CE1	2.52	0.45
1:A:729:VAL:HG21	1:A:781:VAL:HG11	1.99	0.45
1:B:402:ILE:HD12	1:B:403:ARG:H	1.81	0.45
1:A:302:THR:HG21	1:A:315:THR:HB	1.98	0.45
1:C:182:LYS:HD2	1:C:187:LYS:HG3	1.98	0.45
1:C:351:TYR:HB2	1:C:468:ILE:HB	1.98	0.45
1:A:295:PRO:HA	1:A:298:GLU:HB3	1.99	0.45
1:A:722:VAL:HG12	1:A:930:ALA:HB1	1.99	0.45
1:B:785:VAL:HG12	1:B:787:GLN:H	1.81	0.45
1:B:1001:LEU:O	1:B:1005:GLN:HG3	2.16	0.45
1:A:295:PRO:HA	1:A:298:GLU:HB2	1.98	0.45
1:A:456:PHE:HB3	1:A:473:TYR:HD2	1.81	0.45
1:C:788:ILE:HG13	1:C:876:ALA:HB2	1.98	0.45
1:B:56:LEU:HD21	1:B:91:TYR:HB2	1.98	0.45
1:B:56:LEU:HG	1:B:270:LEU:HD13	1.99	0.45
1:C:15:CYS:SG	1:C:137:ASN:N	2.86	0.45
1:C:276:LEU:HG	1:C:306:PHE:HE1	1.81	0.45
1:B:206:LYS:HB3	1:B:223:LEU:HD22	1.97	0.45

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:C:48:LEU:HB3	1:C:276:LEU:HD11	1.98	0.45
1:B:503:VAL:HA	1:B:506:GLN:HG3	1.99	0.45
1:B:276:LEU:HG	1:B:306:PHE:HE1	1.81	0.45
1:B:774:GLN:HE22	1:B:1018:ILE:HG21	1.81	0.45
1:A:189:LEU:HD12	1:A:217:PRO:HG2	1.99	0.45
1:B:366:SER:HB3	1:B:388:ASN:HD21	1.81	0.45
1:B:483:VAL:HG12	1:B:484:LYS:HG3	1.98	0.45
1:A:384:PRO:HG3	1:C:415:THR:HG21	1.98	0.45
1:A:1105:THR:HG21	1:A:1110:TYR:HA	1.99	0.45
1:C:611:LEU:HB2	1:C:650:LEU:HD13	1.99	0.45
1:B:386:LYS:NZ	1:C:981:LEU:O	2.39	0.45
1:C:100:ILE:HG21	1:C:263:ALA:HB2	1.99	0.45
1:A:33:THR:OG1	1:A:219:GLY:O	2.32	0.45
1:A:105:ILE:HD12	1:A:105:ILE:HA	1.81	0.45
1:B:725:GLU:OE1	1:B:725:GLU:N	2.50	0.45
1:B:749:CYS:O	1:B:753:LEU:HB3	2.17	0.45
1:C:773:GLU:HG2	1:C:1019:ARG:HH21	1.80	0.45
1:C:1028:LYS:HE2	1:C:1028:LYS:HB2	1.75	0.45
1:A:774:GLN:HA	1:A:777:ASN:HD22	1.82	0.45
1:B:606:ASN:ND2	1:B:606:ASN:O	2.37	0.45
1:B:751:ASN:HA	1:B:754:LEU:HG	1.98	0.45
1:C:751:ASN:HA	1:C:754:LEU:HG	1.99	0.45
1:B:905:ARG:HE	1:B:1050:MET:HB3	1.81	0.45
1:C:287:ASP:OD1	1:C:288:ALA:N	2.50	0.45
1:C:390:LEU:HD12	1:C:391:CYS:H	1.82	0.45
1:A:290:ASP:HB3	1:A:293:LEU:HB2	1.99	0.45
1:A:738:CYS:SG	1:A:739:THR:N	2.90	0.45
1:A:774:GLN:HE22	1:A:1018:ILE:HG21	1.81	0.45
1:B:403:ARG:NH2	1:C:372:ALA:O	2.32	0.45
1:C:965:GLN:HE21	1:C:1003:SER:HB2	1.81	0.45
1:A:989:ALA:O	1:A:993:ILE:HG12	2.16	0.45
1:B:393:THR:OG1	1:B:394:ASN:N	2.49	0.45
1:C:390:LEU:HD12	1:C:391:CYS:H	1.81	0.45
1:C:393:THR:OG1	1:C:394:ASN:N	2.50	0.45
1:C:912:THR:OG1	1:C:1106:GLN:NE2	2.50	0.45
1:A:287:ASP:OD1	1:A:288:ALA:N	2.50	0.45
1:B:733:LYS:HB3	1:B:861:LEU:HD23	1.99	0.45
1:C:342:PHE:CE1	1:C:511:VAL:HG11	2.52	0.45
1:B:659:SER:HB3	1:B:698:SER:HB2	1.97	0.45
1:A:1081:ILE:HG23	1:A:1088:HIS:HB2	1.98	0.45
1:B:403:ARG:NH2	1:C:372:ALA:O	2.32	0.45

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:B:947:LYS:HE3	1:B:947:LYS:HB2	1.89	0.45
1:A:773:GLU:HG2	1:A:1019:ARG:HH21	1.82	0.45
1:C:310:LYS:NZ	1:C:663:ASP:OD2	2.44	0.45
1:B:989:ALA:O	1:B:993:ILE:HD12	2.16	0.45
1:B:785:VAL:HG12	1:B:787:GLN:H	1.81	0.45
1:C:191:GLU:HB2	1:C:223:LEU:HD11	1.98	0.45
1:C:382:VAL:HG21	1:C:515:PHE:HZ	1.82	0.45
1:C:1116:THR:OG1	1:C:1118:ASP:OD1	2.34	0.45
1:A:483:VAL:HG12	1:A:484:LYS:HG3	1.99	0.45
1:B:382:VAL:HG11	1:B:515:PHE:HZ	1.81	0.45
1:B:393:THR:OG1	1:B:516:GLU:O	2.35	0.45
1:C:1106:GLN:HA	1:C:1106:GLN:NE2	2.28	0.45
1:B:972:ALA:HB2	1:B:996:LEU:HD21	1.99	0.45
1:A:295:PRO:HA	1:A:298:GLU:HB2	1.97	0.45
1:A:390:LEU:HD12	1:A:391:CYS:H	1.80	0.45
1:A:564:GLN:OE1	1:A:564:GLN:N	2.49	0.45
1:C:115:GLN:HB2	1:C:233:ILE:HD12	1.98	0.45
1:C:1038:LYS:HB3	1:C:1038:LYS:HE3	1.76	0.45
1:A:785:VAL:HG12	1:A:787:GLN:H	1.81	0.45
1:B:770:ILE:HA	1:B:773:GLU:HG3	1.98	0.45
1:C:931:ILE:HA	1:C:934:ILE:HG22	1.98	0.45
1:B:676:THR:HA	1:B:690:GLN:HG2	1.99	0.45
1:B:779:GLN:C	1:B:779:GLN:HE21	2.23	0.45
1:C:53:ASP:OD1	1:C:195:LYS:NZ	2.49	0.45
1:C:409:GLN:NE2	1:C:418:ILE:HG12	2.31	0.45
1:C:931:ILE:HA	1:C:934:ILE:HG22	1.99	0.45
1:C:84:LEU:HD11	1:C:267:VAL:HG11	1.98	0.45
1:B:407:VAL:O	1:B:410:ILE:HG22	2.17	0.45
1:A:31:SER:OG	1:A:60:SER:O	2.31	0.45
1:A:791:THR:HG21	1:A:806:LEU:HD11	1.99	0.45
1:A:890:ALA:HA	1:C:1046:GLY:HA2	1.98	0.45
1:B:564:GLN:HG2	1:B:577:ARG:HD2	1.98	0.45
1:B:802:PHE:HE1	1:B:927:PHE:HE2	1.65	0.45
1:C:330:PRO:HD3	1:C:579:PRO:HB2	1.98	0.45
1:C:719:THR:HG23	1:C:1070:ALA:HB2	1.99	0.45
1:C:785:VAL:HG12	1:C:787:GLN:H	1.81	0.45
1:B:412:PRO:HG3	1:B:429:PHE:HD1	1.82	0.45
1:B:415:THR:HG21	1:C:384:PRO:HG3	1.99	0.45
1:B:814:LYS:N	1:B:868:GLU:OE2	2.49	0.45
1:B:193:VAL:HB	1:B:204:TYR:HD2	1.80	0.45
1:C:44:ARG:HD2	1:C:47:VAL:HG21	1.99	0.45

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:C:390:LEU:HD12	1:C:391:CYS:H	1.81	0.45
1:B:404:GLY:HA2	1:B:508:TYR:CD2	2.52	0.45
1:C:355:ARG:NH2	1:C:398:ASP:OD2	2.50	0.45
1:B:35:GLY:HA3	1:B:56:LEU:HD12	1.98	0.45
1:B:106:PHE:O	1:B:117:LEU:N	2.40	0.45
1:B:398:ASP:OD2	1:B:423:TYR:OH	2.25	0.45
1:B:802:PHE:HE1	1:B:927:PHE:HE2	1.65	0.45
1:C:226:LEU:HG	1:C:227:VAL:HG13	1.99	0.45
1:C:390:LEU:HD12	1:C:391:CYS:H	1.82	0.45
1:A:787:GLN:HG2	1:C:701:VAL:HG13	1.98	0.45
1:B:483:VAL:HG12	1:B:484:LYS:HG3	1.98	0.45
1:B:195:LYS:HG3	1:B:197:ILE:HG13	1.98	0.45
1:B:293:LEU:HD22	1:B:294:ASP:OD2	2.16	0.45
1:C:523:THR:HG23	1:C:524:VAL:HG13	1.98	0.45
1:A:1081:ILE:HD11	1:A:1135:ASN:HB3	1.98	0.45
1:B:748:GLU:N	1:B:748:GLU:OE2	2.50	0.45
1:C:106:PHE:HB2	1:C:117:LEU:HB3	1.98	0.45
1:C:989:ALA:O	1:C:993:ILE:HG13	2.16	0.45
1:B:650:LEU:HD21	1:B:678:ILE:HD12	1.98	0.45
1:A:574:ASP:OD1	1:A:574:ASP:N	2.49	0.45
1:C:395:VAL:HG12	1:C:515:PHE:HB3	1.99	0.45
1:A:574:ASP:OD1	1:A:574:ASP:N	2.49	0.45
1:B:983:ARG:HB2	1:B:984:LEU:HD22	1.99	0.45
1:C:966:LEU:HD13	1:C:1000:ARG:HH22	1.81	0.45
1:B:278:LYS:HB2	1:B:306:PHE:CE2	2.52	0.45
1:B:332:ILE:HG13	1:B:362:VAL:HG22	1.98	0.45
1:C:37:TYR:OH	1:C:53:ASP:OD1	2.28	0.45
1:A:325:SER:HA	1:A:540:ASN:HB3	1.99	0.45
1:A:332:ILE:HB	1:A:362:VAL:HG11	1.99	0.45
1:A:732:THR:OG1	1:A:955:ASN:OD1	2.19	0.45
1:B:722:VAL:HG22	1:B:930:ALA:HB1	1.99	0.45
1:C:326:ILE:HD13	1:C:533:LEU:HD12	1.99	0.45
1:C:888:PHE:HE2	1:C:1034:LEU:HB2	1.82	0.45
1:B:733:LYS:HG2	1:B:774:GLN:NE2	2.32	0.45
1:A:919:ASN:HD22	1:A:922:LEU:HD21	1.82	0.45
1:C:923:ILE:HA	1:C:926:GLN:HG3	1.99	0.45
1:A:483:VAL:HG12	1:A:484:LYS:HG3	1.99	0.45
1:C:276:LEU:HD21	1:C:304:LYS:HA	1.99	0.45
1:A:168:PHE:HZ	1:C:466:ARG:HH22	1.65	0.45
1:B:97:LYS:NZ	1:B:183:GLN:O	2.44	0.45
1:B:877:LEU:O	1:B:881:THR:HG23	2.17	0.45

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:B:1013:ILE:HD13	1:C:1012:LEU:HB3	1.99	0.45
1:C:914:ASN:OD1	1:C:914:ASN:N	2.50	0.45
1:A:701:VAL:HG13	1:B:787:GLN:HG2	1.98	0.45
1:C:332:ILE:HG13	1:C:362:VAL:HG22	1.98	0.45
1:A:332:ILE:HG13	1:A:362:VAL:HG22	1.99	0.45
1:A:606:ASN:O	1:A:606:ASN:ND2	2.50	0.45
1:A:47:VAL:HG12	1:C:569:ILE:HA	1.98	0.45
1:C:722:VAL:HG12	1:C:1065:VAL:HG22	1.99	0.45
1:B:887:THR:HG21	1:B:894:LEU:HB2	1.98	0.45
1:C:785:VAL:HG12	1:C:787:GLN:H	1.81	0.45
1:B:287:ASP:OD1	1:B:288:ALA:N	2.50	0.45
1:B:809:PRO:HA	1:B:814:LYS:HD2	1.97	0.45
1:A:683:CYS:O	1:A:695:TYR:N	2.44	0.45
1:C:726:ILE:HG13	1:C:947:LYS:HE2	1.98	0.45
1:C:807:PRO:HA	1:C:816:SER:HA	1.99	0.45
1:A:302:THR:HG21	1:A:315:THR:HB	1.98	0.45
1:A:751:ASN:HA	1:A:754:LEU:HG	1.98	0.45
1:C:781:VAL:HG23	1:C:1026:ALA:HA	1.99	0.45
1:A:983:ARG:HD3	1:C:517:LEU:HD11	1.99	0.45
1:C:1002:GLN:O	1:C:1006:THR:HG23	2.17	0.45
1:B:121:ASN:HA	1:B:126:VAL:HA	1.98	0.45
1:B:731:MET:H	1:B:774:GLN:NE2	2.14	0.45
1:C:102:ARG:HG2	1:C:246:ILE:HD11	1.99	0.45
1:C:247:SER:O	1:C:258:TRP:NE1	2.47	0.45
1:C:796:ASP:OD1	1:C:796:ASP:N	2.48	0.45
1:C:296:LEU:O	1:C:299:THR:OG1	2.35	0.45
1:C:330:PRO:HD3	1:C:579:PRO:HB2	1.99	0.45
1:A:109:THR:OG1	1:A:111:ASP:OD1	2.31	0.45
1:B:612:TYR:HD2	1:B:620:VAL:HG22	1.82	0.45
1:B:733:LYS:HB3	1:B:861:LEU:HD23	1.99	0.45
1:C:393:THR:OG1	1:C:394:ASN:N	2.50	0.45
1:A:1041:ASP:OD1	1:B:1030:SER:HB2	2.16	0.45
1:C:909:ILE:HD11	1:C:1047:TYR:CD1	2.52	0.45
1:A:357:ARG:HD3	1:A:396:TYR:CE1	2.52	0.45
1:A:978:ASN:HA	1:A:981:LEU:HG	1.99	0.45
1:C:99:ASN:HB3	1:C:102:ARG:HH12	1.82	0.45
1:A:642:VAL:HG22	1:A:651:ILE:HG22	1.99	0.45
1:B:366:SER:O	1:B:370:ASN:HB2	2.17	0.45
1:B:722:VAL:HG22	1:B:930:ALA:HB1	1.99	0.45
1:B:1115:ILE:HG22	1:B:1137:VAL:HG23	1.98	0.45
1:C:290:ASP:HB3	1:C:293:LEU:HB2	1.99	0.45

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:C:763:LEU:HB2	1:C:1008:VAL:HG11	1.99	0.45
1:A:787:GLN:HG2	1:C:701:VAL:HG13	1.98	0.45
1:B:442:ASP:OD2	1:B:509:ARG:NE	2.46	0.45
1:B:613:GLN:HA	1:B:648:GLY:HA3	1.99	0.45
1:A:976:VAL:HG13	1:A:979:ASP:HB3	1.99	0.45
1:A:809:PRO:HA	1:A:814:LYS:HD2	1.99	0.45
1:B:326:ILE:HD11	1:B:534:VAL:HG22	1.99	0.45
1:B:408:ARG:HH21	1:B:409:GLN:HE21	1.64	0.45
1:C:323:THR:HB	1:C:537:LYS:HZ3	1.81	0.45
1:A:22:THR:O	1:A:78:ARG:NH1	2.50	0.45
1:B:574:ASP:OD1	1:B:574:ASP:N	2.49	0.45
1:C:216:LEU:HD21	1:C:266:TYR:HE2	1.81	0.44
1:C:773:GLU:HA	1:C:776:LYS:HE3	1.99	0.44
1:B:326:ILE:HD11	1:B:534:VAL:HG22	1.99	0.44
1:C:914:ASN:OD1	1:C:914:ASN:N	2.50	0.44
1:A:1005:GLN:HA	1:A:1008:VAL:HG22	1.98	0.44
1:A:1037:SER:OG	1:A:1039:ARG:HG2	2.16	0.44
1:C:402:ILE:HD12	1:C:403:ARG:H	1.81	0.44
1:C:52:GLN:OE1	1:C:274:THR:OG1	2.30	0.44
1:C:231:ILE:HG13	1:C:233:ILE:HG22	1.99	0.44
1:C:574:ASP:OD1	1:C:574:ASP:N	2.49	0.44
1:C:886:TRP:HH2	1:C:904:TYR:HB3	1.82	0.44
1:A:384:PRO:HG3	1:C:415:THR:HG21	1.98	0.44
1:A:1105:THR:HG21	1:A:1110:TYR:HA	1.98	0.44
1:A:1107:ARG:HE	1:A:1107:ARG:HB3	1.61	0.44
1:B:401:VAL:HG22	1:B:509:ARG:HG2	1.97	0.44
1:C:596:SER:HB2	1:C:611:LEU:HB3	1.98	0.44
1:B:947:LYS:HE3	1:B:947:LYS:HB2	1.86	0.44
1:C:379:CYS:CA	1:C:432:CYS:HB3	2.47	0.44
1:A:1135:ASN:OD1	1:A:1136:THR:N	2.45	0.44
1:A:33:THR:OG1	1:A:219:GLY:O	2.24	0.44
1:C:332:ILE:HG12	1:C:524:VAL:HG21	1.99	0.44
1:C:597:VAL:HG13	1:C:608:VAL:HG13	1.99	0.44
1:A:887:THR:HB	1:A:894:LEU:HD23	1.98	0.44
1:B:1086:LYS:HE3	1:B:1122:VAL:HG11	1.99	0.44
1:B:1116:THR:OG1	1:B:1118:ASP:OD1	2.31	0.44
1:C:906:PHE:CD2	1:C:916:LEU:HB2	2.52	0.44
1:A:39:PRO:HG3	1:A:51:THR:HG21	1.98	0.44
1:A:332:ILE:HG13	1:A:362:VAL:HG22	1.98	0.44
1:C:287:ASP:OD1	1:C:288:ALA:N	2.50	0.44
1:C:560:LEU:O	1:C:577:ARG:NH2	2.50	0.44

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:106:PHE:O	1:A:117:LEU:N	2.42	0.44
1:B:1106:GLN:H	1:B:1106:GLN:HG2	1.53	0.44
1:A:96:GLU:HG3	1:A:100:ILE:HG22	1.99	0.44
1:A:1051:SER:OG	1:A:1064:HIS:ND1	2.41	0.44
1:A:326:ILE:HD11	1:A:534:VAL:HG22	2.00	0.44
1:B:382:VAL:HG11	1:B:515:PHE:HZ	1.82	0.44
1:C:190:ARG:CZ	1:C:207:HIS:CE1	3.00	0.44
1:A:611:LEU:HD13	1:A:650:LEU:HD13	1.99	0.44
1:B:365:TYR:CE2	1:B:387:LEU:HD12	2.52	0.44
1:B:287:ASP:OD1	1:B:288:ALA:N	2.50	0.44
1:B:290:ASP:HB3	1:B:293:LEU:HB2	1.99	0.44
1:A:315:THR:HG23	1:A:595:VAL:HG23	1.99	0.44
1:A:717:ASN:OD1	1:A:718:PHE:N	2.50	0.44
1:A:1002:GLN:O	1:A:1006:THR:HG23	2.18	0.44
1:B:489:TYR:CE2	1:C:372:ALA:HB3	2.51	0.44
1:C:1127:ASP:OD2	1:C:1127:ASP:N	2.50	0.44
1:A:871:ALA:HA	1:A:874:THR:HG22	1.99	0.44
1:C:560:LEU:HD12	1:C:561:PRO:HD2	1.99	0.44
1:B:287:ASP:OD1	1:B:288:ALA:N	2.50	0.44
1:C:129:LYS:HZ3	1:C:169:GLU:HB2	1.81	0.44
1:C:290:ASP:HB3	1:C:293:LEU:HB2	1.99	0.44
1:B:406:GLU:HB3	1:B:418:ILE:HG21	1.98	0.44
1:B:406:GLU:HA	1:B:409:GLN:NE2	2.32	0.44
1:A:785:VAL:HG12	1:A:787:GLN:H	1.81	0.44
1:C:404:GLY:HA2	1:C:407:VAL:HG13	1.98	0.44
1:A:701:VAL:HG13	1:B:787:GLN:HG2	1.99	0.44
1:A:390:LEU:HD12	1:A:391:CYS:H	1.82	0.44
1:C:782:PHE:HE1	1:C:870:ILE:HD11	1.81	0.44
1:A:971:GLY:H	1:B:755:GLN:NE2	2.14	0.44
1:B:656:VAL:HG12	1:B:658:ASN:H	1.82	0.44
1:C:431:GLY:HA2	1:C:515:PHE:HE1	1.82	0.44
1:B:717:ASN:OD1	1:B:718:PHE:N	2.49	0.44
1:C:316:SER:H	1:C:595:VAL:HG22	1.82	0.44
1:C:203:ILE:HG22	1:C:226:LEU:HD23	1.99	0.44
1:A:401:VAL:HG22	1:A:509:ARG:HG2	1.99	0.44
1:A:642:VAL:HG22	1:A:651:ILE:HG22	1.99	0.44
1:B:83:VAL:HG22	1:B:239:GLN:HE21	1.82	0.44
1:C:183:GLN:HG3	1:C:187:LYS:HG3	1.99	0.44
1:B:1115:ILE:HG22	1:B:1137:VAL:HG23	1.99	0.44
1:A:454:ARG:HD3	1:A:457:ARG:HD3	1.99	0.44
1:A:896:ILE:HD12	1:A:897:PRO:HD2	1.98	0.44

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:C:435:ALA:HB2	1:C:510:VAL:HG12	1.99	0.44
1:A:102:ARG:NH1	1:A:121:ASN:O	2.47	0.44
1:B:394:ASN:OD1	1:B:394:ASN:N	2.50	0.44
1:B:972:ALA:HB2	1:B:996:LEU:HD21	1.99	0.44
1:B:986:PRO:HD2	1:B:987:PRO:HD3	1.99	0.44
1:A:574:ASP:O	1:A:587:ILE:N	2.36	0.44
1:A:1028:LYS:HE2	1:A:1028:LYS:HB2	1.85	0.44
1:A:102:ARG:HA	1:A:102:ARG:HD3	1.86	0.44
1:A:753:LEU:HD12	1:A:759:PHE:HZ	1.82	0.44
1:B:1120:THR:OG1	1:B:1121:PHE:N	2.51	0.44
1:A:729:VAL:HG13	1:A:1059:GLY:HA2	1.99	0.44
1:A:733:LYS:HE3	1:A:771:ALA:HB1	1.98	0.44
1:A:773:GLU:HA	1:A:776:LYS:HE3	1.99	0.44
1:A:914:ASN:OD1	1:A:914:ASN:N	2.49	0.44
1:B:697:MET:HE2	1:B:697:MET:HB2	1.85	0.44
1:B:1086:LYS:HA	1:B:1125:ASN:HA	1.99	0.44
1:A:701:VAL:HG13	1:B:787:GLN:HG2	1.99	0.44
1:C:884:SER:OG	1:C:894:LEU:O	2.34	0.44
1:A:393:THR:HA	1:A:522:ALA:HA	1.98	0.44
1:A:58:PHE:HD2	1:A:290:ASP:HB2	1.82	0.44
1:C:989:ALA:O	1:C:993:ILE:HG12	2.17	0.44
1:A:822:LEU:HD11	1:A:1061:VAL:HG21	2.00	0.44
1:A:1094:VAL:HG11	1:A:1107:ARG:HH21	1.82	0.44
1:A:106:PHE:HB2	1:A:117:LEU:HD12	1.99	0.44
1:A:1033:VAL:HG21	1:A:1053:PRO:HG3	1.99	0.44
1:C:42:VAL:HG13	1:C:44:ARG:HH22	1.82	0.44
1:C:403:ARG:HE	1:C:406:GLU:CD	2.25	0.44
1:A:564:GLN:N	1:B:41:LYS:HZ2	2.11	0.44
1:C:454:ARG:NH2	1:C:469:SER:O	2.46	0.44
1:A:774:GLN:HE22	1:A:1018:ILE:HG21	1.82	0.44
1:A:864:LEU:O	1:C:669:GLY:N	2.50	0.44
1:B:310:LYS:HG3	1:B:600:PRO:HA	2.00	0.44
1:A:295:PRO:O	1:A:299:THR:HG23	2.17	0.44
1:A:773:GLU:OE2	1:A:1019:ARG:NH1	2.44	0.44
1:A:887:THR:HG21	1:A:894:LEU:HB2	1.99	0.44
1:A:978:ASN:HA	1:A:981:LEU:HG	2.00	0.44
1:C:719:THR:HG23	1:C:1070:ALA:HB2	1.99	0.44
1:A:129:LYS:HE2	1:A:129:LYS:HB2	1.80	0.44
1:A:886:TRP:HB3	1:A:1035:GLY:HA2	2.00	0.44
1:C:905:ARG:NH1	1:C:1036:GLN:HB2	2.33	0.44
1:B:402:ILE:HD12	1:B:403:ARG:H	1.82	0.44

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:722:VAL:HG12	1:A:930:ALA:HB1	1.99	0.44
1:C:612:TYR:HD1	1:C:620:VAL:HG22	1.82	0.44
1:A:102:ARG:HE	1:A:246:ILE:HD11	1.82	0.44
1:B:212:LEU:HD13	1:B:217:PRO:HG3	2.00	0.44
1:B:722:VAL:HG22	1:B:930:ALA:HB1	1.99	0.44
1:B:1019:ARG:HD2	1:B:1019:ARG:HA	1.69	0.44
1:B:777:ASN:HB3	1:B:1019:ARG:NH2	2.33	0.44
1:C:191:GLU:N	1:C:191:GLU:OE1	2.51	0.44
1:C:642:VAL:HG22	1:C:651:ILE:HG22	1.99	0.44
1:A:43:PHE:HB3	1:C:565:PHE:O	2.17	0.44
1:A:888:PHE:HE2	1:A:1034:LEU:HG	1.83	0.44
1:B:118:LEU:HD21	1:B:135:PHE:HE2	1.81	0.44
1:B:87:ASN:OD1	1:B:87:ASN:N	2.42	0.44
1:B:111:ASP:OD1	1:B:113:LYS:NZ	2.48	0.44
1:A:813:SER:OG	1:A:868:GLU:OE2	2.29	0.44
1:B:611:LEU:HD13	1:B:650:LEU:HD22	1.99	0.44
1:A:717:ASN:OD1	1:A:718:PHE:N	2.50	0.44
1:A:825:LYS:HE2	1:A:825:LYS:HB2	1.87	0.44
1:B:1029:MET:HE2	1:B:1033:VAL:HG21	1.99	0.44
1:C:214:ARG:HD2	1:C:215:GLY:N	2.31	0.44
1:B:1115:ILE:HG22	1:B:1137:VAL:HG23	1.99	0.44
1:A:787:GLN:HG2	1:C:701:VAL:HG13	2.00	0.44
1:B:552:LEU:HD22	1:B:587:ILE:HD13	2.00	0.44
1:B:390:LEU:HD12	1:B:391:CYS:H	1.83	0.44
1:C:106:PHE:O	1:C:117:LEU:N	2.41	0.44
1:C:34:ARG:NH2	1:C:221:SER:H	2.14	0.44
1:A:229:LEU:HG	1:A:231:ILE:HG23	2.00	0.44
1:A:1050:MET:HE2	1:A:1050:MET:HB3	1.73	0.44
1:B:1086:LYS:HE3	1:B:1122:VAL:HG11	1.99	0.44
1:C:523:THR:HG23	1:C:524:VAL:HG13	1.99	0.44
1:B:715:PRO:HD3	1:C:894:LEU:HD21	2.00	0.44
1:C:332:ILE:HG12	1:C:524:VAL:HG21	1.99	0.44
1:B:287:ASP:OD1	1:B:288:ALA:N	2.50	0.44
1:B:442:ASP:OD2	1:B:509:ARG:NE	2.42	0.44
1:C:22:THR:O	1:C:78:ARG:NH1	2.50	0.44
1:A:212:LEU:HD22	1:A:217:PRO:HD3	2.00	0.44
1:B:1086:LYS:HD3	1:B:1122:VAL:HG11	1.98	0.44
1:A:787:GLN:HG2	1:C:701:VAL:HG13	1.99	0.44
1:B:192:PHE:HD1	1:B:203:ILE:HD11	1.82	0.44
1:C:330:PRO:HG3	1:C:579:PRO:HB2	2.00	0.44
1:C:360:ASN:OD1	1:C:523:THR:OG1	2.35	0.44

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:B:877:LEU:O	1:B:881:THR:HG23	2.18	0.44
1:A:195:LYS:HD2	1:A:197:ILE:HD11	1.99	0.44
1:B:298:GLU:HB3	1:B:315:THR:HG21	1.99	0.44
1:C:916:LEU:HD23	1:C:917:TYR:CD2	2.53	0.44
1:B:33:THR:OG1	1:B:34:ARG:NH1	2.51	0.44
1:A:1040:VAL:HG21	1:B:1035:GLY:HA3	1.99	0.44
1:B:220:PHE:HE2	1:B:285:ILE:HG22	1.82	0.44
1:B:296:LEU:O	1:B:299:THR:OG1	2.32	0.44
1:C:58:PHE:HD2	1:C:290:ASP:HB2	1.82	0.44
1:C:276:LEU:HD21	1:C:304:LYS:HA	1.99	0.44
1:B:984:LEU:HB2	1:B:989:ALA:HB2	1.98	0.44
1:C:560:LEU:HD12	1:C:561:PRO:HD2	1.99	0.44
1:A:697:MET:HE2	1:A:697:MET:HB2	1.80	0.44
1:B:394:ASN:HB3	1:B:396:TYR:HE2	1.82	0.44
1:C:759:PHE:HA	1:C:762:GLN:HE21	1.83	0.44
1:C:330:PRO:HD3	1:C:579:PRO:HB2	2.00	0.44
1:C:406:GLU:N	1:C:406:GLU:OE2	2.50	0.44
1:A:287:ASP:OD1	1:A:288:ALA:N	2.51	0.44
1:A:719:THR:HG23	1:A:1068:VAL:HB	1.99	0.44
1:B:465:GLU:OE1	1:B:466:ARG:N	2.49	0.44
1:C:442:ASP:OD2	1:C:509:ARG:NE	2.44	0.44
1:A:751:ASN:HA	1:A:754:LEU:HG	1.99	0.44
1:B:100:ILE:HA	1:B:246:ILE:HD12	2.00	0.44
1:B:351:TYR:HE2	1:B:452:LEU:HB2	1.81	0.44
1:A:1081:ILE:HD11	1:A:1135:ASN:HB3	1.99	0.44
1:B:483:VAL:HG12	1:B:484:LYS:HG3	1.99	0.44
1:C:523:THR:HG23	1:C:524:VAL:HG13	1.98	0.44
1:A:486:PHE:O	1:A:487:ASN:ND2	2.51	0.44
1:A:1028:LYS:NZ	1:A:1042:PHE:O	2.43	0.44
1:C:1050:MET:SD	1:C:1052:PHE:HE1	2.40	0.44
1:B:350:VAL:HG21	1:B:418:ILE:HD11	1.99	0.44
1:B:731:MET:H	1:B:774:GLN:HE22	1.65	0.44
1:A:976:VAL:HG22	1:C:572:THR:HG22	1.98	0.44
1:A:1002:GLN:O	1:A:1006:THR:HG23	2.17	0.44
1:C:277:LEU:HD12	1:C:285:ILE:HD13	1.99	0.44
1:B:751:ASN:HA	1:B:754:LEU:HG	1.99	0.44
1:C:762:GLN:HA	1:C:765:ARG:HG2	2.00	0.44
1:B:1050:MET:HE2	1:B:1050:MET:HB2	1.96	0.44
1:C:1135:ASN:OD1	1:C:1136:THR:N	2.42	0.44
1:A:983:ARG:HH21	1:C:517:LEU:HD11	1.82	0.44
1:A:34:ARG:NH2	1:A:221:SER:H	2.15	0.44

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:C:1054:GLN:HB2	1:C:1061:VAL:HG13	2.00	0.44
1:A:398:ASP:HB2	1:A:512:VAL:HG13	2.00	0.44
1:A:1104:VAL:HG23	1:A:1115:ILE:HD13	1.99	0.44
1:A:756:TYR:HB3	1:A:759:PHE:HD2	1.83	0.44
1:B:989:ALA:O	1:B:993:ILE:HG12	2.17	0.44
1:C:1115:ILE:HG22	1:C:1137:VAL:HG23	1.98	0.44
1:B:92:PHE:HD1	1:B:93:ALA:N	2.16	0.44
1:C:434:ILE:HB	1:C:511:VAL:HG13	1.99	0.44
1:C:896:ILE:HD12	1:C:897:PRO:HD2	1.98	0.44
1:A:850:ILE:HG23	1:A:860:VAL:HG11	2.00	0.44
1:A:949:GLN:HA	1:A:952:VAL:HG12	1.99	0.44
1:C:906:PHE:HA	1:C:909:ILE:HG12	2.00	0.44
1:C:971:GLY:O	1:C:995:ARG:NH1	2.51	0.44
1:B:38:TYR:O	1:B:204:TYR:OH	2.25	0.44
1:C:537:LYS:HD3	1:C:538:CYS:N	2.32	0.44
1:B:781:VAL:HG13	1:B:782:PHE:HD1	1.82	0.44
1:A:890:ALA:HA	1:C:1046:GLY:HA2	1.99	0.44
1:C:785:VAL:HG12	1:C:787:GLN:H	1.82	0.44
1:A:547:THR:O	1:B:978:ASN:ND2	2.51	0.44
1:C:39:PRO:HG3	1:C:51:THR:HG21	1.98	0.44
1:C:402:ILE:HD11	1:C:418:ILE:HG13	1.99	0.44
1:B:468:ILE:HG12	1:C:115:GLN:HE22	1.83	0.44
1:A:976:VAL:HG13	1:A:979:ASP:HB3	2.00	0.44
1:C:914:ASN:OD1	1:C:914:ASN:N	2.49	0.44
1:A:291:CYS:HB2	1:A:301:CYS:HB2	1.79	0.44
1:B:91:TYR:OH	1:B:191:GLU:OE1	2.33	0.44
1:B:431:GLY:HA2	1:B:515:PHE:CZ	2.53	0.44
1:C:106:PHE:HB2	1:C:117:LEU:HD12	1.99	0.44
1:A:658:ASN:C	1:A:658:ASN:HD22	2.26	0.44
1:B:557:LYS:HD2	1:B:557:LYS:HA	1.68	0.44
1:C:777:ASN:O	1:C:781:VAL:HG12	2.17	0.44
1:C:960:ASN:HA	1:C:963:VAL:HG22	1.98	0.44
1:B:650:LEU:HD21	1:B:678:ILE:HD11	1.99	0.44
1:C:971:GLY:HA3	1:C:995:ARG:HH12	1.82	0.44
1:C:1116:THR:OG1	1:C:1118:ASP:OD1	2.34	0.44
1:A:1107:ARG:HE	1:A:1107:ARG:HB3	1.60	0.44
1:A:954:GLN:OE1	1:A:1014:ARG:NE	2.49	0.44
1:B:1019:ARG:HG3	1:B:1019:ARG:HH11	1.83	0.44
1:C:532:ASN:OD1	1:C:533:LEU:N	2.50	0.44
1:A:406:GLU:N	1:A:406:GLU:OE2	2.51	0.44
1:C:33:THR:OG1	1:C:219:GLY:O	2.31	0.44

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:742:ILE:HG13	1:A:997:ILE:HG22	1.99	0.44
1:B:762:GLN:HB3	1:B:765:ARG:HH21	1.82	0.44
1:B:767:LEU:HD13	1:B:770:ILE:HD11	2.00	0.44
1:C:912:THR:OG1	1:C:914:ASN:OD1	2.35	0.44
1:A:796:ASP:OD1	1:A:796:ASP:N	2.45	0.44
1:B:287:ASP:OD1	1:B:288:ALA:N	2.50	0.44
1:B:762:GLN:HA	1:B:765:ARG:HE	1.82	0.44
1:B:915:VAL:HG21	1:B:1108:ASN:HD22	1.82	0.44
1:B:1010:GLN:O	1:B:1014:ARG:HG3	2.18	0.44
1:B:1086:LYS:HE3	1:B:1122:VAL:HG11	2.00	0.44
1:C:882:ILE:HD12	1:C:882:ILE:HA	1.85	0.44
1:C:390:LEU:HD12	1:C:391:CYS:H	1.82	0.44
1:B:517:LEU:HD11	1:C:983:ARG:HD2	1.98	0.44
1:C:287:ASP:OD1	1:C:288:ALA:N	2.50	0.44
1:B:717:ASN:OD1	1:B:718:PHE:N	2.49	0.44
1:C:727:LEU:HD11	1:C:1024:LEU:HG	2.00	0.44
1:A:787:GLN:HG2	1:C:701:VAL:HG13	2.00	0.44
1:A:826:VAL:HG23	1:A:945:LEU:HD12	1.99	0.44
1:A:781:VAL:HG13	1:A:782:PHE:HD1	1.82	0.44
1:B:574:ASP:OD1	1:B:574:ASP:N	2.50	0.44
1:A:1081:ILE:HG23	1:A:1088:HIS:HB2	2.00	0.44
1:B:985:ASP:OD2	1:B:985:ASP:N	2.47	0.44
1:C:738:CYS:O	1:C:742:ILE:HB	2.17	0.44
1:A:756:TYR:HE2	1:A:997:ILE:HD11	1.83	0.44
1:B:722:VAL:HG22	1:B:930:ALA:HB1	2.00	0.44
1:B:877:LEU:O	1:B:881:THR:HG23	2.18	0.44
1:C:290:ASP:HB3	1:C:293:LEU:HB2	1.99	0.44
1:C:513:LEU:HD12	1:C:513:LEU:HA	1.86	0.44
1:C:989:ALA:O	1:C:993:ILE:HG12	2.18	0.44
1:A:611:LEU:HD13	1:A:650:LEU:HD13	2.00	0.44
1:C:566:GLY:N	1:C:575:ALA:O	2.36	0.44
1:A:104:TRP:H	1:A:119:ILE:HB	1.82	0.44
1:B:729:VAL:HG12	1:B:1059:GLY:HA2	2.00	0.44
1:B:762:GLN:HA	1:B:765:ARG:HE	1.83	0.44
1:C:34:ARG:NH2	1:C:221:SER:H	2.16	0.44
1:A:1102:TRP:HD1	1:A:1135:ASN:HD22	1.66	0.44
1:C:767:LEU:HA	1:C:770:ILE:HG22	2.00	0.44
1:A:287:ASP:OD1	1:A:288:ALA:N	2.51	0.44
1:A:642:VAL:HG22	1:A:651:ILE:HG22	2.00	0.44
1:A:1105:THR:HG21	1:A:1110:TYR:HA	2.00	0.44
1:B:506:GLN:HG3	1:B:507:PRO:HD2	1.99	0.44

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:B:972:ALA:HB2	1:B:996:LEU:HD21	2.00	0.44
1:B:369:TYR:CZ	1:B:384:PRO:HB2	2.53	0.44
1:C:407:VAL:O	1:C:410:ILE:HG22	2.17	0.44
1:C:729:VAL:HG21	1:C:781:VAL:HG11	1.99	0.44
1:B:552:LEU:HD12	1:B:585:LEU:HD22	1.99	0.44
1:C:277:LEU:HD12	1:C:285:ILE:HG12	1.99	0.44
1:A:199:GLY:HA2	1:A:232:GLY:HA2	1.99	0.44
1:A:711:SER:OG	1:B:895:GLN:OE1	2.34	0.44
1:B:877:LEU:O	1:B:881:THR:HG23	2.17	0.44
1:A:195:LYS:HG3	1:A:197:ILE:HG23	1.98	0.44
1:C:28:TYR:HB3	1:C:61:ASN:HB3	1.98	0.44
1:A:712:ILE:HG21	1:A:1096:VAL:HG12	1.99	0.44
1:C:712:ILE:HD12	1:C:712:ILE:HA	1.71	0.44
1:C:877:LEU:O	1:C:881:THR:HG23	2.18	0.44
1:B:409:GLN:HA	1:B:414:GLN:HG2	2.00	0.44
1:B:731:MET:H	1:B:774:GLN:NE2	2.16	0.44
1:C:276:LEU:HD22	1:C:306:PHE:HE1	1.82	0.44
1:B:271:GLN:HB2	1:B:273:ARG:HH22	1.82	0.44
1:B:408:ARG:NH1	1:C:408:ARG:HA	2.33	0.44
1:B:587:ILE:HD12	1:B:587:ILE:HA	1.76	0.44
1:C:204:TYR:HA	1:C:225:PRO:HA	1.99	0.44
1:A:289:VAL:HG11	1:A:300:LYS:HD2	2.00	0.44
1:A:737:ASP:OD1	1:A:740:MET:HB3	2.17	0.44
1:C:105:ILE:HG23	1:C:239:GLN:HB2	1.99	0.44
1:A:1046:GLY:HA2	1:B:890:ALA:HA	2.00	0.44
1:B:442:ASP:OD2	1:B:509:ARG:NE	2.47	0.44
1:C:229:LEU:HD22	1:C:231:ILE:HG22	2.00	0.44
1:C:1086:LYS:HE3	1:C:1122:VAL:HG11	1.99	0.44
1:A:39:PRO:HG3	1:A:51:THR:HG21	1.99	0.44
1:A:222:ALA:HB2	1:A:285:ILE:HB	2.00	0.44
1:B:105:ILE:HD11	1:B:241:LEU:HD21	1.98	0.44
1:B:390:LEU:HD12	1:B:391:CYS:H	1.82	0.44
1:A:915:VAL:HG12	1:A:1109:PHE:HD2	1.83	0.44
1:A:1000:ARG:O	1:A:1003:SER:OG	2.27	0.44
1:B:722:VAL:HG22	1:B:930:ALA:HB1	2.00	0.44
1:B:796:ASP:OD1	1:B:796:ASP:N	2.51	0.44
1:A:611:LEU:HD22	1:A:678:ILE:HD11	2.00	0.44
1:A:183:GLN:HG3	1:A:187:LYS:HG3	1.99	0.44
1:C:738:CYS:HB3	1:C:760:CYS:HB2	1.87	0.44
1:A:393:THR:HA	1:A:522:ALA:HA	1.99	0.44
1:A:900:MET:HE1	1:C:1079:PRO:HA	1.99	0.44

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:B:735:SER:HB3	1:B:861:LEU:HD21	2.00	0.44
1:A:287:ASP:OD1	1:A:288:ALA:N	2.50	0.44
1:A:406:GLU:N	1:A:406:GLU:OE1	2.51	0.44
1:A:890:ALA:HA	1:C:1046:GLY:HA2	1.99	0.44
1:B:731:MET:H	1:B:774:GLN:NE2	2.16	0.44
1:B:574:ASP:OD1	1:B:574:ASP:N	2.48	0.44
1:B:785:VAL:HG12	1:B:787:GLN:H	1.82	0.44
1:B:884:SER:OG	1:B:887:THR:OG1	2.27	0.44
1:C:106:PHE:O	1:C:117:LEU:N	2.46	0.44
1:A:712:ILE:O	1:A:1075:PHE:N	2.38	0.44
1:C:290:ASP:HB3	1:C:293:LEU:HB2	1.99	0.44
1:C:299:THR:HG22	1:C:597:VAL:HG21	1.99	0.44
1:A:740:MET:HB2	1:A:740:MET:HE3	1.67	0.44
1:A:960:ASN:O	1:A:964:LYS:HB2	2.18	0.44
1:C:785:VAL:HG12	1:C:787:GLN:H	1.82	0.44
1:C:712:ILE:HD12	1:C:712:ILE:HA	1.79	0.44
1:C:1139:ASP:HB3	1:C:1142:GLN:HG2	2.00	0.44
1:A:598:ILE:HD11	1:A:678:ILE:HD13	1.99	0.44
1:A:1039:ARG:HH21	1:C:1039:ARG:HH11	1.65	0.44
1:A:1005:GLN:HE21	1:C:1006:THR:HG22	1.83	0.44
1:B:1046:GLY:HA2	1:C:890:ALA:HA	2.00	0.44
1:C:431:GLY:HA2	1:C:515:PHE:CE1	2.52	0.44
1:B:513:LEU:HD12	1:B:513:LEU:HA	1.87	0.44
1:C:1045:LYS:HD3	1:C:1045:LYS:HA	1.73	0.44
1:B:781:VAL:HG13	1:B:782:PHE:HD1	1.83	0.44
1:C:598:ILE:HB	1:C:609:ALA:HB3	2.00	0.44
1:B:506:GLN:HG3	1:B:507:PRO:HD2	1.99	0.44
1:C:1107:ARG:H	1:C:1107:ARG:HG3	1.41	0.44
1:A:120:VAL:HB	1:A:127:VAL:HB	1.99	0.44
1:A:774:GLN:HE22	1:A:1018:ILE:HG21	1.82	0.44
1:B:997:ILE:O	1:B:1001:LEU:HB2	2.17	0.44
1:C:96:GLU:HG3	1:C:99:ASN:H	1.82	0.44
1:A:722:VAL:HG22	1:A:930:ALA:HB1	2.00	0.44
1:A:748:GLU:H	1:A:748:GLU:HG3	1.55	0.44
1:A:1135:ASN:OD1	1:A:1136:THR:N	2.43	0.44
1:B:37:TYR:CZ	1:B:195:LYS:HE2	2.53	0.44
1:B:337:PRO:HD2	1:B:358:ILE:HG21	1.99	0.44
1:B:420:ASP:OD2	1:B:424:LYS:NZ	2.51	0.44
1:A:357:ARG:NH2	1:A:396:TYR:OH	2.51	0.44
1:A:409:GLN:HA	1:A:414:GLN:HG2	1.99	0.44
1:A:417:ASN:O	1:A:421:TYR:HB2	2.18	0.44

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:781:VAL:HG13	1:A:782:PHE:HD1	1.82	0.44
1:A:931:ILE:HA	1:A:934:ILE:HG22	2.00	0.44
1:C:287:ASP:OD2	1:C:288:ALA:N	2.50	0.44
1:C:358:ILE:HB	1:C:395:VAL:HG23	2.00	0.44
1:C:730:SER:OG	1:C:731:MET:N	2.51	0.44
1:A:58:PHE:CD2	1:A:290:ASP:HB2	2.53	0.44
1:B:393:THR:OG1	1:B:516:GLU:O	2.36	0.44
1:A:642:VAL:HG22	1:A:651:ILE:HG22	2.00	0.44
1:B:785:VAL:HG12	1:B:787:GLN:H	1.82	0.44
1:B:902:MET:HE1	1:B:1050:MET:HE2	1.98	0.44
1:A:566:GLY:HA2	1:B:43:PHE:HB3	1.99	0.44
1:A:973:ILE:HG21	1:A:983:ARG:NH1	2.33	0.44
1:A:984:LEU:HB2	1:A:989:ALA:HB2	2.00	0.44
1:B:314:GLN:HE22	1:B:594:GLY:HA3	1.82	0.44
1:A:58:PHE:HD2	1:A:290:ASP:HB2	1.83	0.44
1:C:594:GLY:H	1:C:613:GLN:NE2	2.16	0.44
1:C:763:LEU:HB2	1:C:1008:VAL:HG11	2.00	0.44
1:A:985:ASP:OD1	1:A:987:PRO:HD2	2.17	0.44
1:A:1031:GLU:HB3	1:A:1037:SER:HB2	2.00	0.44
1:C:131:CYS:HB2	1:C:133:PHE:CE1	2.53	0.44
1:A:357:ARG:HD2	1:A:359:SER:OG	2.17	0.44
1:B:677:PRO:HB3	1:C:864:LEU:HD11	2.00	0.44
1:C:409:GLN:HA	1:C:414:GLN:HE21	1.82	0.44
1:C:914:ASN:ND2	1:C:1106:GLN:OE1	2.45	0.44
1:A:33:THR:OG1	1:A:219:GLY:O	2.35	0.44
1:A:40:ASP:OD1	1:A:40:ASP:N	2.51	0.44
1:B:951:VAL:HA	1:B:954:GLN:HG3	1.98	0.44
1:C:483:VAL:HG12	1:C:484:LYS:HG3	1.99	0.44
1:C:688:THR:HA	1:C:690:GLN:HG2	2.00	0.44
1:C:398:ASP:OD2	1:C:423:TYR:OH	2.24	0.44
1:B:906:PHE:CD2	1:B:916:LEU:HB2	2.53	0.44
1:C:1033:VAL:HG21	1:C:1053:PRO:HG3	2.00	0.44
1:A:395:VAL:HG12	1:A:515:PHE:HB3	1.99	0.44
1:C:1086:LYS:HE3	1:C:1122:VAL:HG11	1.98	0.44
1:B:195:LYS:HD2	1:B:195:LYS:HA	1.70	0.44
1:B:884:SER:HG	1:B:887:THR:HG1	1.61	0.44
1:A:402:ILE:HD12	1:A:403:ARG:H	1.83	0.44
1:A:701:VAL:HG13	1:B:787:GLN:HG2	1.99	0.44
1:C:1051:SER:OG	1:C:1064:HIS:ND1	2.39	0.44
1:A:189:LEU:HD13	1:A:210:ILE:HD13	2.00	0.44
1:A:730:SER:OG	1:A:731:MET:N	2.49	0.44

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:C:912:THR:O	1:C:915:VAL:HG22	2.18	0.44
1:A:319:ARG:NH1	1:B:744:GLY:HA3	2.33	0.43
1:A:436:TRP:HE1	1:A:509:ARG:HH21	1.65	0.43
1:A:822:LEU:HD11	1:A:1061:VAL:HG11	2.00	0.43
1:A:902:MET:HE2	1:A:902:MET:HB2	1.85	0.43
1:C:18:PHE:HB2	1:C:21:ARG:HB2	2.00	0.43
1:B:37:TYR:HB2	1:B:204:TYR:CE2	2.53	0.43
1:B:569:ILE:HG22	1:C:47:VAL:HB	2.00	0.43
1:C:334:ASN:OD1	1:C:335:LEU:N	2.43	0.43
1:A:774:GLN:OE1	1:A:777:ASN:ND2	2.51	0.43
1:C:220:PHE:HE2	1:C:285:ILE:HG22	1.83	0.43
1:C:193:VAL:HG13	1:C:223:LEU:HD21	1.99	0.43
1:C:737:ASP:OD1	1:C:740:MET:HB3	2.17	0.43
1:C:964:LYS:O	1:C:967:SER:OG	2.32	0.43
1:B:1010:GLN:HE22	1:B:1014:ARG:HD2	1.83	0.43
1:C:785:VAL:HG12	1:C:787:GLN:H	1.82	0.43
1:A:1050:MET:HE2	1:A:1050:MET:HB3	1.67	0.43
1:C:909:ILE:HA	1:C:1038:LYS:NZ	2.32	0.43
1:A:395:VAL:HG22	1:A:515:PHE:HB3	2.00	0.43
1:B:563:GLN:N	1:B:563:GLN:OE1	2.51	0.43
1:C:358:ILE:HB	1:C:395:VAL:HG23	2.00	0.43
1:A:1084:ASP:OD1	1:A:1084:ASP:N	2.41	0.43
1:B:37:TYR:OH	1:B:53:ASP:OD2	2.35	0.43
1:A:517:LEU:HD12	1:A:518:LEU:HB2	1.99	0.43
1:C:1049:LEU:HD11	1:C:1067:TYR:HB2	1.99	0.43
1:A:722:VAL:HG22	1:A:930:ALA:HB1	1.99	0.43
1:A:788:ILE:HG23	1:A:876:ALA:HB2	1.99	0.43
1:B:129:LYS:NZ	1:B:166:CYS:SG	2.87	0.43
1:C:22:THR:O	1:C:78:ARG:NH1	2.51	0.43
1:C:33:THR:HG22	1:C:58:PHE:HE1	1.83	0.43
1:C:302:THR:O	1:C:304:LYS:NZ	2.51	0.43
1:A:931:ILE:HA	1:A:934:ILE:HG22	2.00	0.43
1:B:332:ILE:HG13	1:B:362:VAL:HG22	1.99	0.43
1:B:398:ASP:HB2	1:B:512:VAL:HG13	2.00	0.43
1:B:576:VAL:HG12	1:B:587:ILE:HD11	1.99	0.43
1:B:906:PHE:HD2	1:B:916:LEU:HB2	1.83	0.43
1:C:365:TYR:HD1	1:C:387:LEU:HB3	1.83	0.43
1:C:966:LEU:HA	1:C:1000:ARG:HH12	1.82	0.43
1:A:61:ASN:C	1:A:61:ASN:ND2	2.74	0.43
1:C:358:ILE:HG22	1:C:524:VAL:HG11	1.99	0.43
1:C:394:ASN:N	1:C:394:ASN:OD1	2.51	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:C:483:VAL:HG12	1:C:484:LYS:HG3	2.00	0.43
1:C:523:THR:HG23	1:C:524:VAL:HG13	1.99	0.43
1:A:80:ALA:O	1:A:245:HIS:NE2	2.45	0.43
1:C:226:LEU:HG	1:C:227:VAL:HG13	2.00	0.43
1:C:871:ALA:HA	1:C:874:THR:HG22	1.98	0.43
1:B:191:GLU:HB2	1:B:223:LEU:HD21	1.99	0.43
1:B:222:ALA:HB2	1:B:285:ILE:HB	2.00	0.43
1:B:1033:VAL:HG21	1:B:1053:PRO:HG3	1.98	0.43
1:C:785:VAL:HG12	1:C:787:GLN:H	1.82	0.43
1:B:767:LEU:HA	1:B:770:ILE:HG22	1.99	0.43
1:B:864:LEU:HG	1:B:865:LEU:HD23	2.00	0.43
1:C:1135:ASN:OD1	1:C:1136:THR:N	2.44	0.43
1:A:1010:GLN:HG3	1:A:1014:ARG:NH1	2.32	0.43
1:A:896:ILE:HD12	1:A:897:PRO:HD2	2.00	0.43
1:B:884:SER:OG	1:B:887:THR:OG1	2.29	0.43
1:B:997:ILE:O	1:B:1001:LEU:HB2	2.17	0.43
1:C:394:ASN:N	1:C:394:ASN:OD1	2.51	0.43
1:C:722:VAL:HG22	1:C:930:ALA:HB1	1.99	0.43
1:B:574:ASP:OD1	1:B:574:ASP:N	2.50	0.43
1:C:105:ILE:HD11	1:C:110:LEU:HD22	2.00	0.43
1:C:330:PRO:HD3	1:C:579:PRO:HB2	2.00	0.43
1:A:406:GLU:N	1:A:406:GLU:OE1	2.51	0.43
1:A:562:PHE:O	1:B:41:LYS:NZ	2.40	0.43
1:B:431:GLY:HA2	1:B:515:PHE:CE2	2.54	0.43
1:C:126:VAL:HB	1:C:172:SER:HB3	2.00	0.43
1:B:743:CYS:HB3	1:B:977:LEU:HD21	2.01	0.43
1:B:769:GLY:HA2	1:B:772:VAL:HG12	2.01	0.43
1:C:596:SER:HB2	1:C:611:LEU:HD23	1.99	0.43
1:A:393:THR:OG1	1:A:516:GLU:O	2.33	0.43
1:B:382:VAL:HG11	1:B:515:PHE:HZ	1.83	0.43
1:C:989:ALA:O	1:C:993:ILE:HG12	2.18	0.43
1:B:392:PHE:CD1	1:B:517:LEU:HD13	2.53	0.43
1:C:99:ASN:O	1:C:102:ARG:NH1	2.50	0.43
1:A:425:LEU:HD22	1:A:426:PRO:HD2	1.99	0.43
1:A:882:ILE:HD12	1:A:882:ILE:HA	1.78	0.43
1:B:960:ASN:O	1:B:964:LYS:HB2	2.18	0.43
1:C:97:LYS:HD2	1:C:97:LYS:HA	1.84	0.43
1:B:989:ALA:O	1:B:993:ILE:HG12	2.18	0.43
1:A:332:ILE:HG13	1:A:362:VAL:HG22	1.99	0.43
1:B:728:PRO:HD3	1:B:947:LYS:HG3	2.00	0.43
1:B:731:MET:H	1:B:774:GLN:NE2	2.16	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:92:PHE:HE2	1:A:265:TYR:HB2	1.83	0.43
1:A:287:ASP:OD2	1:A:288:ALA:N	2.51	0.43
1:A:934:ILE:HG13	1:A:938:LEU:HD23	1.99	0.43
1:B:562:PHE:HE1	1:C:41:LYS:HE3	1.83	0.43
1:B:735:SER:HB2	1:B:861:LEU:HD21	1.99	0.43
1:C:777:ASN:HA	1:C:780:GLU:HG3	1.99	0.43
1:B:513:LEU:HG	1:B:515:PHE:CE1	2.54	0.43
1:B:866:THR:O	1:B:870:ILE:HG12	2.19	0.43
1:C:295:PRO:HB3	1:C:595:VAL:HG22	2.01	0.43
1:B:183:GLN:HG3	1:B:187:LYS:HG3	2.00	0.43
1:B:1002:GLN:HA	1:B:1005:GLN:NE2	2.33	0.43
1:C:871:ALA:HA	1:C:874:THR:HG22	2.00	0.43
1:C:905:ARG:NH1	1:C:1036:GLN:HB2	2.34	0.43
1:A:503:VAL:HA	1:A:506:GLN:HG3	1.99	0.43
1:A:650:LEU:HD12	1:A:650:LEU:HA	1.90	0.43
1:A:764:ASN:O	1:A:768:THR:HG23	2.18	0.43
1:B:130:VAL:HB	1:B:168:PHE:HB2	2.00	0.43
1:B:517:LEU:HD12	1:B:518:LEU:HB2	1.98	0.43
1:C:796:ASP:OD1	1:C:796:ASP:N	2.49	0.43
1:A:61:ASN:C	1:A:61:ASN:ND2	2.74	0.43
1:B:287:ASP:OD1	1:B:288:ALA:N	2.50	0.43
1:C:483:VAL:HG12	1:C:484:LYS:HG3	2.00	0.43
1:C:962:LEU:HD12	1:C:962:LEU:HA	1.87	0.43
1:A:222:ALA:HB2	1:A:285:ILE:HB	1.99	0.43
1:A:326:ILE:HD11	1:A:534:VAL:HG22	2.00	0.43
1:A:759:PHE:HA	1:A:762:GLN:HE21	1.83	0.43
1:B:563:GLN:N	1:B:563:GLN:OE1	2.51	0.43
1:B:735:SER:HB2	1:B:861:LEU:HD21	2.01	0.43
1:A:310:LYS:HG3	1:A:600:PRO:HA	1.99	0.43
1:C:472:ILE:HG23	1:C:491:PRO:HG2	1.99	0.43
1:B:677:PRO:HB3	1:C:864:LEU:HD21	2.00	0.43
1:B:707:TYR:CD2	1:C:792:PRO:HG3	2.51	0.43
1:C:617:CYS:N	1:C:649:CYS:SG	2.91	0.43
1:C:390:LEU:HD12	1:C:391:CYS:H	1.83	0.43
1:A:426:PRO:HD3	1:A:463:PRO:HB3	2.00	0.43
1:A:745:ASP:OD2	1:C:549:THR:OG1	2.36	0.43
1:B:738:CYS:HB2	1:B:760:CYS:HB2	1.51	0.43
1:C:730:SER:OG	1:C:731:MET:N	2.51	0.43
1:A:332:ILE:HG13	1:A:362:VAL:HG22	2.00	0.43
1:A:858:LEU:HD13	1:A:959:LEU:HD12	2.00	0.43
1:B:407:VAL:O	1:B:410:ILE:HG22	2.18	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:B:986:PRO:HD2	1:B:987:PRO:HD3	2.00	0.43
1:A:115:GLN:HE22	1:C:468:ILE:HG12	1.82	0.43
1:B:731:MET:HG3	1:B:1018:ILE:HG13	2.00	0.43
1:B:93:ALA:HB3	1:B:266:TYR:HD2	1.82	0.43
1:C:537:LYS:HE3	1:C:537:LYS:HB3	1.89	0.43
1:A:298:GLU:HB3	1:A:315:THR:HG21	2.00	0.43
1:A:611:LEU:HD22	1:A:678:ILE:HD11	2.00	0.43
1:B:483:VAL:HG12	1:B:484:LYS:HG3	2.00	0.43
1:B:569:ILE:HA	1:C:47:VAL:HG12	2.00	0.43
1:C:383:SER:O	1:C:387:LEU:CB	2.67	0.43
1:C:424:LYS:HA	1:C:424:LYS:HD3	1.87	0.43
1:C:596:SER:HB2	1:C:611:LEU:HB3	1.99	0.43
1:C:733:LYS:HD3	1:C:861:LEU:HB2	2.00	0.43
1:C:916:LEU:HD22	1:C:917:TYR:CD1	2.53	0.43
1:A:737:ASP:OD2	1:C:317:ASN:ND2	2.50	0.43
1:C:84:LEU:HD11	1:C:267:VAL:HG11	2.00	0.43
1:A:129:LYS:HG2	1:A:133:PHE:HZ	1.82	0.43
1:A:578:ASP:OD2	1:A:581:THR:OG1	2.35	0.43
1:C:1004:LEU:HD12	1:C:1004:LEU:HA	1.83	0.43
1:B:825:LYS:HE3	1:B:825:LYS:HB2	1.81	0.43
1:B:908:GLY:O	1:B:1038:LYS:NZ	2.50	0.43
1:B:407:VAL:O	1:B:410:ILE:HG22	2.19	0.43
1:B:513:LEU:HD12	1:B:513:LEU:HA	1.90	0.43
1:B:912:THR:O	1:B:915:VAL:HG12	2.18	0.43
1:C:442:ASP:OD2	1:C:509:ARG:NE	2.46	0.43
1:A:483:VAL:HG12	1:A:484:LYS:HG3	2.00	0.43
1:A:498:GLN:HB3	1:A:501:TYR:HE2	1.83	0.43
1:A:701:VAL:HG13	1:B:787:GLN:HG2	2.00	0.43
1:B:785:VAL:HG12	1:B:787:GLN:H	1.83	0.43
1:C:287:ASP:OD2	1:C:288:ALA:N	2.51	0.43
1:C:994:ASP:O	1:C:998:THR:HG23	2.18	0.43
1:C:1086:LYS:HE3	1:C:1122:VAL:HG11	1.99	0.43
1:A:950:ASP:N	1:A:950:ASP:OD1	2.49	0.43
1:A:887:THR:HB	1:A:894:LEU:HD23	2.00	0.43
1:A:1081:ILE:HG23	1:A:1088:HIS:HB2	2.00	0.43
1:B:796:ASP:OD1	1:B:796:ASP:N	2.50	0.43
1:B:396:TYR:CZ	1:C:230:PRO:HG3	2.54	0.43
1:B:878:LEU:O	1:B:882:ILE:HG12	2.18	0.43
1:A:290:ASP:OD2	1:A:290:ASP:N	2.50	0.43
1:A:985:ASP:O	1:A:989:ALA:HB2	2.18	0.43
1:C:951:VAL:HA	1:C:954:GLN:HG3	2.00	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:1019:ARG:HA	1:A:1019:ARG:HH11	1.82	0.43
1:B:513:LEU:HD12	1:B:513:LEU:HA	1.87	0.43
1:B:713:ALA:HB2	1:C:895:GLN:CD	2.44	0.43
1:B:1038:LYS:HD3	1:B:1038:LYS:HA	1.71	0.43
1:C:733:LYS:HD3	1:C:861:LEU:HB2	2.00	0.43
1:C:738:CYS:HB2	1:C:760:CYS:HB2	1.46	0.43
1:A:1094:VAL:HG23	1:A:1096:VAL:HG13	2.01	0.43
1:B:384:PRO:HA	1:B:387:LEU:HD23	2.01	0.43
1:B:777:ASN:HD21	1:B:1019:ARG:HD2	1.83	0.43
1:C:503:VAL:HA	1:C:506:GLN:HG3	2.00	0.43
1:A:442:ASP:OD2	1:A:509:ARG:NE	2.47	0.43
1:B:574:ASP:OD1	1:B:574:ASP:N	2.52	0.43
1:C:287:ASP:OD2	1:C:288:ALA:N	2.51	0.43
1:A:731:MET:HG3	1:A:1018:ILE:HG13	2.00	0.43
1:B:58:PHE:HD2	1:B:290:ASP:HB2	1.83	0.43
1:A:731:MET:H	1:A:774:GLN:NE2	2.15	0.43
1:B:983:ARG:HG3	1:B:984:LEU:HD23	2.00	0.43
1:C:642:VAL:HG22	1:C:651:ILE:HG22	2.00	0.43
1:A:726:ILE:HD12	1:A:1061:VAL:HG12	2.00	0.43
1:B:598:ILE:HB	1:B:609:ALA:HB3	1.99	0.43
1:B:1096:VAL:HG13	1:B:1103:PHE:HB2	2.01	0.43
1:C:1135:ASN:OD1	1:C:1136:THR:N	2.41	0.43
1:A:401:VAL:HG22	1:A:509:ARG:HG2	2.00	0.43
1:B:989:ALA:O	1:B:993:ILE:HG12	2.18	0.43
1:A:1083:HIS:CD2	1:A:1137:VAL:H	2.36	0.43
1:B:533:LEU:HD21	1:B:585:LEU:HD11	2.01	0.43
1:B:557:LYS:HE2	1:B:559:PHE:HE1	1.84	0.43
1:C:738:CYS:HB3	1:C:760:CYS:HB2	1.90	0.43
1:C:906:PHE:CD1	1:C:916:LEU:HB2	2.53	0.43
1:A:52:GLN:OE1	1:A:274:THR:OG1	2.24	0.43
1:A:332:ILE:HG13	1:A:362:VAL:HG22	2.01	0.43
1:A:395:VAL:HG12	1:A:515:PHE:HB3	1.99	0.43
1:B:398:ASP:HB2	1:B:512:VAL:HG13	2.01	0.43
1:B:738:CYS:O	1:B:742:ILE:HB	2.19	0.43
1:A:369:TYR:OH	1:C:415:THR:OG1	2.17	0.43
1:A:730:SER:OG	1:A:731:MET:N	2.51	0.43
1:B:395:VAL:HG12	1:B:515:PHE:HB3	2.00	0.43
1:B:1017:GLU:OE1	1:C:1019:ARG:NH1	2.52	0.43
1:C:287:ASP:OD1	1:C:288:ALA:N	2.51	0.43
1:C:574:ASP:N	1:C:574:ASP:OD1	2.51	0.43
1:B:388:ASN:OD1	1:B:388:ASN:N	2.52	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:B:393:THR:HG21	1:B:518:LEU:HB2	2.00	0.43
1:B:781:VAL:HG13	1:B:782:PHE:HD2	1.83	0.43
1:A:1002:GLN:HA	1:A:1005:GLN:HG3	2.00	0.43
1:B:89:GLY:HA3	1:B:270:LEU:HD12	2.00	0.43
1:C:819:GLU:HA	1:C:822:LEU:HG	1.99	0.43
1:B:735:SER:HB2	1:B:861:LEU:HD21	2.00	0.43
1:A:565:PHE:O	1:B:43:PHE:HB3	2.19	0.43
1:B:231:ILE:HG13	1:B:233:ILE:HG12	1.98	0.43
1:C:914:ASN:OD1	1:C:914:ASN:N	2.52	0.43
1:C:950:ASP:O	1:C:954:GLN:NE2	2.50	0.43
1:A:334:ASN:ND2	1:A:360:ASN:O	2.42	0.43
1:C:337:PRO:HD2	1:C:358:ILE:HD12	2.00	0.43
1:A:334:ASN:ND2	1:A:360:ASN:O	2.44	0.43
1:A:408:ARG:HH12	1:A:409:GLN:HE21	1.65	0.43
1:A:785:VAL:HG12	1:A:787:GLN:H	1.83	0.43
1:C:58:PHE:HD2	1:C:290:ASP:HB2	1.84	0.43
1:C:193:VAL:HG22	1:C:204:TYR:HB2	2.00	0.43
1:C:1116:THR:OG1	1:C:1118:ASP:OD2	2.36	0.43
1:A:222:ALA:HB2	1:A:285:ILE:HB	1.99	0.43
1:B:52:GLN:HE22	1:B:273:ARG:N	2.17	0.43
1:B:751:ASN:HA	1:B:754:LEU:HG	2.01	0.43
1:B:973:ILE:HG21	1:B:983:ARG:HH12	1.83	0.43
1:B:1096:VAL:HG11	1:B:1105:THR:HG22	2.01	0.43
1:C:1086:LYS:HE3	1:C:1122:VAL:HG11	2.00	0.43
1:B:418:ILE:HD12	1:B:422:ASN:HD22	1.83	0.43
1:C:574:ASP:OD1	1:C:574:ASP:N	2.50	0.43
1:C:613:GLN:HA	1:C:648:GLY:HA3	2.01	0.43
1:A:606:ASN:O	1:A:606:ASN:ND2	2.46	0.43
1:B:731:MET:HG3	1:B:1018:ILE:HG13	2.00	0.43
1:C:105:ILE:HD11	1:C:110:LEU:HD22	2.00	0.43
1:A:727:LEU:HD23	1:A:728:PRO:HD2	1.99	0.43
1:B:1091:ARG:NH1	1:B:1118:ASP:O	2.52	0.43
1:B:1091:ARG:NH1	1:B:1120:THR:O	2.50	0.43
1:C:774:GLN:HA	1:C:777:ASN:ND2	2.34	0.43
1:C:736:VAL:HG21	1:C:1007:TYR:HE2	1.83	0.43
1:C:976:VAL:HG13	1:C:979:ASP:HB3	2.00	0.43
1:A:100:ILE:HA	1:A:246:ILE:HD12	2.01	0.43
1:C:87:ASN:OD1	1:C:87:ASN:N	2.49	0.43
1:A:41:LYS:HD2	1:C:562:PHE:HE1	1.83	0.43
1:A:1090:PRO:HD3	1:A:1095:PHE:HE2	1.82	0.43
1:C:276:LEU:HD23	1:C:306:PHE:HE1	1.82	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:C:884:SER:OG	1:C:887:THR:OG1	2.28	0.43
1:A:866:THR:OG1	1:A:867:ASP:OD1	2.35	0.43
1:B:385:THR:OG1	1:C:985:ASP:OD2	2.36	0.43
1:C:738:CYS:SG	1:C:739:THR:N	2.91	0.43
1:C:785:VAL:HG12	1:C:787:GLN:H	1.83	0.43
1:C:976:VAL:HG13	1:C:979:ASP:HB3	1.99	0.43
1:A:395:VAL:HG12	1:A:515:PHE:HB3	2.00	0.43
1:A:357:ARG:HG3	1:A:396:TYR:CE1	2.53	0.43
1:A:642:VAL:HG22	1:A:651:ILE:HG22	2.00	0.43
1:A:1040:VAL:HG11	1:B:1035:GLY:HA3	2.01	0.43
1:A:1072:GLU:HG3	1:B:894:LEU:HD13	2.00	0.43
1:C:537:LYS:HE3	1:C:537:LYS:HB3	1.89	0.43
1:A:106:PHE:HB3	1:A:235:ILE:HD12	2.01	0.43
1:A:108:THR:HB	1:A:114:THR:HG21	2.00	0.43
1:A:666:ILE:HD11	1:A:672:ALA:HB2	2.00	0.43
1:B:361:CYS:H	1:B:522:ALA:HB1	1.83	0.43
1:B:912:THR:HG22	1:B:914:ASN:H	1.84	0.43
1:C:1002:GLN:O	1:C:1006:THR:HG23	2.18	0.43
1:B:435:ALA:HB2	1:B:510:VAL:HG22	2.01	0.43
1:B:777:ASN:HD21	1:B:1019:ARG:HD2	1.84	0.43
1:B:877:LEU:O	1:B:881:THR:HG23	2.18	0.43
1:C:402:ILE:HD12	1:C:403:ARG:H	1.83	0.43
1:C:781:VAL:HG13	1:C:782:PHE:CD2	2.52	0.43
1:A:676:THR:HA	1:A:690:GLN:HG2	2.00	0.43
1:C:356:LYS:HE2	1:C:358:ILE:HD11	2.01	0.43
1:C:403:ARG:HB3	1:C:495:TYR:HE1	1.84	0.43
1:A:89:GLY:HA3	1:A:270:LEU:HD12	2.01	0.43
1:A:980:ILE:HG23	1:A:984:LEU:HD13	2.01	0.43
1:B:106:PHE:HB2	1:B:117:LEU:HB3	2.00	0.43
1:B:733:LYS:HB3	1:B:861:LEU:HD23	1.99	0.43
1:B:915:VAL:HG12	1:B:1109:PHE:HD2	1.84	0.43
1:A:287:ASP:OD2	1:A:288:ALA:N	2.52	0.43
1:A:867:ASP:OD1	1:A:867:ASP:N	2.51	0.43
1:A:1115:ILE:HG22	1:A:1137:VAL:HG23	2.00	0.43
1:B:117:LEU:HD12	1:B:233:ILE:HD11	2.00	0.43
1:C:48:LEU:HB3	1:C:276:LEU:HD11	2.00	0.43
1:B:877:LEU:O	1:B:881:THR:HG23	2.18	0.43
1:C:273:ARG:NH1	1:C:290:ASP:OD2	2.51	0.43
1:C:497:PHE:CE2	1:C:507:PRO:HB3	2.53	0.43
1:A:369:TYR:CZ	1:A:384:PRO:HB2	2.53	0.43
1:A:873:TYR:O	1:A:877:LEU:HB2	2.18	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:B:720:ILE:HD11	1:B:1065:VAL:HG12	2.00	0.43
1:C:408:ARG:HH11	1:C:414:GLN:NE2	2.16	0.43
1:C:871:ALA:HA	1:C:874:THR:HG22	2.00	0.43
1:A:287:ASP:OD1	1:A:288:ALA:N	2.51	0.43
1:A:401:VAL:HG22	1:A:509:ARG:HG2	2.00	0.43
1:B:48:LEU:HD23	1:B:276:LEU:HD11	2.00	0.43
1:C:574:ASP:N	1:C:574:ASP:OD1	2.52	0.43
1:B:57:PRO:HB2	1:B:60:SER:HB3	2.01	0.43
1:B:731:MET:H	1:B:774:GLN:NE2	2.16	0.43
1:B:989:ALA:O	1:B:993:ILE:HG12	2.18	0.43
1:C:126:VAL:HB	1:C:172:SER:HB3	2.00	0.43
1:C:916:LEU:HD22	1:C:917:TYR:HD1	1.83	0.43
1:A:911:VAL:HG11	1:A:1067:TYR:HE1	1.84	0.43
1:B:738:CYS:O	1:B:742:ILE:HB	2.18	0.43
1:B:796:ASP:OD1	1:B:796:ASP:N	2.51	0.43
1:A:753:LEU:HD12	1:A:759:PHE:CZ	2.53	0.43
1:B:182:LYS:HD3	1:B:187:LYS:HD2	2.00	0.43
1:B:969:ASN:ND2	1:B:972:ALA:O	2.52	0.43
1:C:965:GLN:HG3	1:C:970:PHE:HZ	1.84	0.43
1:A:781:VAL:HG13	1:A:782:PHE:CD1	2.54	0.43
1:B:960:ASN:O	1:B:964:LYS:HB2	2.19	0.43
1:C:462:LYS:O	1:C:465:GLU:HG3	2.18	0.43
1:C:736:VAL:HG11	1:C:1004:LEU:HD11	1.99	0.43
1:A:404:GLY:HA3	1:A:504:GLY:HA2	2.01	0.43
1:A:759:PHE:HA	1:A:762:GLN:HE21	1.84	0.43
1:B:189:LEU:HD22	1:B:208:THR:HB	2.00	0.43
1:C:131:CYS:SG	1:C:163:ALA:HB1	2.58	0.43
1:C:470:THR:O	1:C:470:THR:OG1	2.32	0.43
1:A:193:VAL:HG21	1:A:270:LEU:HD21	2.00	0.43
1:A:896:ILE:HD12	1:A:897:PRO:HD2	1.99	0.43
1:B:394:ASN:OD1	1:B:394:ASN:N	2.52	0.43
1:C:722:VAL:HG22	1:C:930:ALA:HB1	2.01	0.43
1:A:106:PHE:O	1:A:117:LEU:N	2.43	0.43
1:A:578:ASP:OD2	1:A:581:THR:OG1	2.34	0.43
1:C:878:LEU:HD21	1:C:1054:GLN:HE22	1.83	0.43
1:C:358:ILE:HG22	1:C:524:VAL:HG11	2.00	0.43
1:C:687:GLN:NE2	1:C:688:THR:O	2.52	0.43
1:A:347:PHE:HB2	1:A:401:VAL:HG23	2.00	0.43
1:A:1023:ASN:O	1:A:1027:THR:OG1	2.35	0.43
1:B:567:ARG:HD3	1:B:571:ASP:HA	2.00	0.43
1:B:569:ILE:H	1:B:569:ILE:HG13	1.65	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:C:902:MET:HE1	1:C:1050:MET:HE2	2.01	0.43
1:A:802:PHE:HB3	1:A:806:LEU:HD23	2.01	0.43
1:B:382:VAL:HG11	1:B:515:PHE:HZ	1.83	0.43
1:C:222:ALA:HB2	1:C:285:ILE:HB	2.00	0.43
1:C:1033:VAL:HG21	1:C:1053:PRO:HG3	2.00	0.43
1:B:287:ASP:OD1	1:B:288:ALA:N	2.51	0.43
1:B:751:ASN:HA	1:B:754:LEU:HG	2.00	0.43
1:C:201:PHE:HE2	1:C:203:ILE:HD11	1.83	0.43
1:A:978:ASN:HA	1:A:981:LEU:HG	2.00	0.43
1:B:802:PHE:HE1	1:B:927:PHE:HE2	1.67	0.43
1:B:989:ALA:O	1:B:993:ILE:HG12	2.18	0.43
1:C:273:ARG:HH21	1:C:292:ALA:HB3	1.84	0.43
1:C:1115:ILE:HG22	1:C:1137:VAL:HG23	2.00	0.43
1:A:989:ALA:O	1:A:993:ILE:HG12	2.18	0.43
1:B:287:ASP:OD1	1:B:288:ALA:N	2.51	0.43
1:B:326:ILE:HD13	1:B:533:LEU:HD12	2.01	0.43
1:B:407:VAL:O	1:B:410:ILE:HG22	2.19	0.43
1:B:418:ILE:HD12	1:B:422:ASN:HD22	1.83	0.43
1:B:931:ILE:HA	1:B:934:ILE:HG22	2.00	0.43
1:C:537:LYS:HE3	1:C:537:LYS:HB3	1.91	0.43
1:A:332:ILE:HG13	1:A:362:VAL:HG22	2.00	0.43
1:C:331:ASN:OD1	1:C:331:ASN:N	2.44	0.43
1:A:611:LEU:HD22	1:A:678:ILE:HD11	2.01	0.43
1:C:717:ASN:OD1	1:C:718:PHE:N	2.49	0.43
1:A:825:LYS:HD3	1:A:942:ALA:HA	2.01	0.43
1:A:395:VAL:HG22	1:A:515:PHE:HB3	2.01	0.43
1:B:1086:LYS:HE3	1:B:1122:VAL:HG11	2.00	0.43
1:C:44:ARG:HB3	1:C:279:TYR:CE2	2.54	0.43
1:C:912:THR:OG1	1:C:1106:GLN:OE1	2.32	0.43
1:C:1045:LYS:HD3	1:C:1045:LYS:HA	1.81	0.43
1:B:129:LYS:NZ	1:B:168:PHE:O	2.51	0.43
1:A:390:LEU:HD12	1:A:391:CYS:H	1.83	0.43
1:A:781:VAL:HG13	1:A:782:PHE:CD1	2.54	0.43
1:B:409:GLN:HA	1:B:414:GLN:HG2	2.00	0.43
1:C:83:VAL:HG11	1:C:237:ARG:HH11	1.83	0.43
1:A:414:GLN:HA	1:A:414:GLN:HE21	1.83	0.43
1:A:1081:ILE:HD11	1:A:1135:ASN:HB3	2.00	0.43
1:A:353:TRP:HZ3	1:A:355:ARG:HB2	1.84	0.43
1:B:35:GLY:HA3	1:B:56:LEU:HB3	1.99	0.43
1:C:58:PHE:HB2	1:C:293:LEU:HD22	2.00	0.43
1:C:807:PRO:HA	1:C:816:SER:HA	1.99	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:483:VAL:HG12	1:A:484:LYS:HG3	2.01	0.43
1:B:393:THR:OG1	1:B:394:ASN:N	2.51	0.43
1:A:1106:GLN:OE1	1:A:1106:GLN:N	2.50	0.43
1:A:1028:LYS:HE2	1:A:1028:LYS:HB2	1.87	0.43
1:B:80:ALA:O	1:B:245:HIS:NE2	2.52	0.43
1:C:882:ILE:HD12	1:C:882:ILE:HA	1.87	0.43
1:C:950:ASP:N	1:C:950:ASP:OD1	2.51	0.43
1:A:97:LYS:HA	1:A:97:LYS:HD2	1.80	0.43
1:A:1081:ILE:HD11	1:A:1135:ASN:HB3	2.00	0.43
1:B:513:LEU:HD12	1:B:513:LEU:HA	1.91	0.43
1:B:776:LYS:NZ	1:B:1019:ARG:HE	2.16	0.43
1:B:989:ALA:O	1:B:993:ILE:HG12	2.19	0.43
1:A:170:TYR:CE2	1:A:229:LEU:HD11	2.54	0.43
1:B:731:MET:H	1:B:774:GLN:NE2	2.16	0.43
1:C:404:GLY:HA2	1:C:407:VAL:HG13	1.99	0.43
1:A:1107:ARG:HH21	1:B:896:ILE:HD11	1.84	0.43
1:C:351:TYR:HE2	1:C:452:LEU:HB2	1.83	0.43
1:C:402:ILE:HG23	1:C:407:VAL:HG12	2.00	0.43
1:C:404:GLY:HA2	1:C:407:VAL:HG13	1.99	0.43
1:C:642:VAL:HG22	1:C:651:ILE:HG22	2.01	0.43
1:A:395:VAL:HG22	1:A:515:PHE:HB3	2.01	0.43
1:A:775:ASP:CG	1:A:864:LEU:HG	2.44	0.43
1:A:332:ILE:HG13	1:A:362:VAL:HG22	2.01	0.43
1:A:918:GLU:HG3	1:C:1128:VAL:HG21	2.01	0.43
1:B:1015:ALA:HA	1:B:1018:ILE:HG22	2.01	0.43
1:C:729:VAL:HG21	1:C:781:VAL:HG11	2.01	0.43
1:B:962:LEU:HD12	1:B:962:LEU:HA	1.89	0.43
1:A:483:VAL:HG12	1:A:484:LYS:HG2	2.01	0.43
1:A:809:PRO:HA	1:A:814:LYS:HD2	2.00	0.43
1:B:785:VAL:HG12	1:B:787:GLN:H	1.83	0.43
1:A:727:LEU:HD11	1:A:1024:LEU:HG	2.01	0.43
1:B:357:ARG:O	1:B:357:ARG:NH1	2.52	0.43
1:C:229:LEU:HD23	1:C:229:LEU:HA	1.90	0.43
1:B:985:ASP:OD2	1:B:987:PRO:HD2	2.19	0.43
1:B:34:ARG:NH2	1:B:221:SER:H	2.17	0.43
1:B:276:LEU:HD11	1:B:304:LYS:HE2	2.01	0.43
1:A:562:PHE:CE2	1:B:41:LYS:HD2	2.54	0.43
1:A:726:ILE:HG13	1:A:948:LEU:HD23	2.00	0.43
1:A:886:TRP:HB3	1:A:1035:GLY:HA2	2.00	0.43
1:B:92:PHE:HE1	1:B:265:TYR:HB2	1.84	0.43
1:B:390:LEU:HG	1:B:392:PHE:CE1	2.54	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:B:452:LEU:HA	1:B:494:SER:HA	2.01	0.43
1:A:334:ASN:ND2	1:A:360:ASN:O	2.46	0.43
1:A:886:TRP:HH2	1:A:904:TYR:HB3	1.83	0.43
1:B:592:PHE:HZ	1:C:740:MET:HE2	1.84	0.43
1:B:722:VAL:HG12	1:B:1065:VAL:HG22	2.01	0.43
1:C:730:SER:OG	1:C:731:MET:N	2.51	0.43
1:A:64:TRP:HE1	1:A:264:ALA:HB1	1.83	0.43
1:A:678:ILE:HD11	1:A:684:ALA:HB2	2.01	0.43
1:B:777:ASN:O	1:B:781:VAL:HG12	2.19	0.43
1:C:83:VAL:HG22	1:C:239:GLN:HG2	2.01	0.43
1:A:543:PHE:HZ	1:A:552:LEU:HD21	1.83	0.43
1:A:751:ASN:HA	1:A:754:LEU:HG	2.01	0.43
1:C:574:ASP:OD1	1:C:574:ASP:N	2.51	0.43
1:C:912:THR:O	1:C:915:VAL:HG12	2.19	0.43
1:A:1031:GLU:OE1	1:C:1039:ARG:HD3	2.18	0.43
1:B:31:SER:OG	1:B:60:SER:O	2.32	0.43
1:B:574:ASP:OD1	1:B:574:ASP:N	2.51	0.43
1:B:598:ILE:HD11	1:B:611:LEU:HD23	2.00	0.43
1:C:302:THR:O	1:C:304:LYS:NZ	2.52	0.43
1:A:334:ASN:ND2	1:A:360:ASN:O	2.44	0.43
1:A:1090:PRO:HD3	1:A:1095:PHE:HE2	1.84	0.43
1:C:409:GLN:HA	1:C:414:GLN:HG3	2.01	0.43
1:B:877:LEU:O	1:B:881:THR:HG23	2.19	0.43
1:A:197:ILE:HB	1:A:202:LYS:HE3	2.00	0.43
1:B:280:ASN:OD1	1:B:284:THR:N	2.52	0.43
1:A:1081:ILE:HD12	1:A:1081:ILE:HA	1.84	0.43
1:C:439:ASN:ND2	1:C:506:GLN:OE1	2.49	0.43
1:C:574:ASP:N	1:C:574:ASP:OD1	2.52	0.43
1:A:731:MET:H	1:A:774:GLN:NE2	2.17	0.43
1:A:918:GLU:HG3	1:C:1128:VAL:HG21	2.00	0.43
1:B:874:THR:HG21	1:B:1055:SER:HB2	2.01	0.43
1:B:323:THR:HB	1:B:537:LYS:HZ3	1.83	0.43
1:A:767:LEU:HA	1:A:770:ILE:HG12	2.01	0.43
1:B:576:VAL:HG12	1:B:587:ILE:HD11	2.00	0.43
1:C:775:ASP:HB3	1:C:864:LEU:HD12	2.00	0.43
1:B:720:ILE:HD11	1:B:1065:VAL:HG12	2.00	0.43
1:B:722:VAL:HG22	1:B:930:ALA:HB1	2.01	0.43
1:C:39:PRO:HG3	1:C:51:THR:HG21	2.01	0.43
1:C:730:SER:OG	1:C:731:MET:N	2.52	0.43
1:C:966:LEU:HD13	1:C:1000:ARG:NH1	2.33	0.43
1:C:985:ASP:HB2	1:C:986:PRO:HD2	2.01	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:204:TYR:HD1	1:A:225:PRO:HA	1.83	0.43
1:A:985:ASP:O	1:A:989:ALA:HB2	2.19	0.43
1:A:989:ALA:O	1:A:993:ILE:HG13	2.19	0.43
1:C:960:ASN:O	1:C:964:LYS:HG2	2.18	0.43
1:B:34:ARG:NH2	1:B:221:SER:H	2.17	0.43
1:B:112:SER:HB2	1:B:132:GLU:HB3	2.01	0.43
1:C:725:GLU:OE1	1:C:1064:HIS:NE2	2.52	0.43
1:B:1028:LYS:O	1:B:1032:CYS:HB2	2.18	0.43
1:C:310:LYS:HG3	1:C:600:PRO:HA	2.00	0.43
1:C:717:ASN:OD1	1:C:718:PHE:N	2.49	0.43
1:A:1081:ILE:HD11	1:A:1135:ASN:HB3	2.01	0.43
1:B:37:TYR:OH	1:B:53:ASP:OD1	2.31	0.43
1:B:382:VAL:HG11	1:B:515:PHE:HZ	1.84	0.43
1:C:407:VAL:O	1:C:410:ILE:HG22	2.19	0.43
1:B:989:ALA:O	1:B:993:ILE:HG12	2.19	0.43
1:A:308:VAL:H	1:A:602:THR:HG1	1.67	0.43
1:A:560:LEU:HD12	1:A:562:PHE:HE1	1.84	0.43
1:A:751:ASN:HA	1:A:754:LEU:HG	2.00	0.43
1:A:972:ALA:HB2	1:A:996:LEU:HD21	2.01	0.43
1:A:407:VAL:O	1:A:410:ILE:HG22	2.19	0.43
1:A:521:PRO:HD3	1:B:41:LYS:HZ2	1.83	0.43
1:B:781:VAL:HG13	1:B:782:PHE:CD2	2.54	0.43
1:B:796:ASP:OD1	1:B:796:ASP:N	2.50	0.43
1:C:994:ASP:O	1:C:998:THR:HG23	2.19	0.43
1:A:873:TYR:HE1	1:C:699:LEU:HG	1.84	0.43
1:A:201:PHE:HB3	1:A:229:LEU:HB2	2.00	0.42
1:A:328:ARG:HD3	1:A:543:PHE:HE2	1.84	0.42
1:C:334:ASN:OD1	1:C:335:LEU:N	2.45	0.42
1:A:295:PRO:HA	1:A:298:GLU:HB3	2.02	0.42
1:A:345:THR:O	1:A:509:ARG:NH2	2.52	0.42
1:A:403:ARG:HG2	1:A:497:PHE:HE1	1.83	0.42
1:A:578:ASP:OD2	1:A:581:THR:OG1	2.32	0.42
1:A:950:ASP:N	1:A:950:ASP:OD1	2.49	0.42
1:C:48:LEU:HB3	1:C:276:LEU:HD11	2.00	0.42
1:C:58:PHE:HD1	1:C:290:ASP:HB2	1.84	0.42
1:A:204:TYR:HB3	1:A:223:LEU:HG	2.01	0.42
1:B:106:PHE:HB3	1:B:235:ILE:HG21	2.00	0.42
1:C:722:VAL:HG22	1:C:930:ALA:HB1	2.01	0.42
1:C:1045:LYS:HA	1:C:1045:LYS:HD3	1.77	0.42
1:A:22:THR:O	1:A:78:ARG:NH1	2.52	0.42
1:A:457:ARG:HH12	1:A:461:LEU:HB3	1.83	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:C:334:ASN:O	1:C:362:VAL:N	2.42	0.42
1:A:674:CYS:HB2	1:A:697:MET:HE1	2.01	0.42
1:A:1050:MET:HE2	1:A:1050:MET:HB3	1.69	0.42
1:C:117:LEU:HD12	1:C:117:LEU:HA	1.77	0.42
1:A:751:ASN:HA	1:A:754:LEU:HG	2.00	0.42
1:B:407:VAL:O	1:B:410:ILE:HG22	2.19	0.42
1:C:404:GLY:HA2	1:C:407:VAL:HG13	2.00	0.42
1:C:574:ASP:OD1	1:C:574:ASP:N	2.51	0.42
1:B:751:ASN:HA	1:B:754:LEU:HG	2.00	0.42
1:C:825:LYS:HE3	1:C:825:LYS:HB2	1.87	0.42
1:C:923:ILE:HA	1:C:926:GLN:HG3	2.01	0.42
1:B:455:LEU:HG	1:B:456:PHE:CD2	2.54	0.42
1:B:717:ASN:OD1	1:B:718:PHE:N	2.50	0.42
1:C:36:VAL:HG11	1:C:220:PHE:CZ	2.54	0.42
1:C:759:PHE:HA	1:C:762:GLN:OE1	2.19	0.42
1:A:332:ILE:HG13	1:A:362:VAL:HG22	2.01	0.42
1:B:351:TYR:HE1	1:B:452:LEU:HB2	1.84	0.42
1:C:537:LYS:HE3	1:C:537:LYS:HB3	1.90	0.42
1:C:970:PHE:HD2	1:C:996:LEU:HA	1.84	0.42
1:B:361:CYS:HB3	1:B:524:VAL:O	2.19	0.42
1:C:287:ASP:OD1	1:C:288:ALA:N	2.52	0.42
1:C:1073:LYS:HE3	1:C:1075:PHE:CZ	2.54	0.42
1:A:740:MET:HE1	1:C:319:ARG:HD3	2.01	0.42
1:C:290:ASP:HB3	1:C:293:LEU:HB2	2.00	0.42
1:A:462:LYS:HE2	1:A:465:GLU:HG3	2.00	0.42
1:A:884:SER:OG	1:A:894:LEU:O	2.37	0.42
1:B:407:VAL:O	1:B:410:ILE:HG22	2.19	0.42
1:B:517:LEU:HD11	1:C:983:ARG:HD2	2.01	0.42
1:C:1019:ARG:HA	1:C:1019:ARG:HH11	1.84	0.42
1:A:483:VAL:HG12	1:A:484:LYS:HG3	2.00	0.42
1:B:406:GLU:HB3	1:B:418:ILE:HG21	2.01	0.42
1:C:737:ASP:OD2	1:C:737:ASP:N	2.45	0.42
1:A:102:ARG:HE	1:A:246:ILE:HD11	1.84	0.42
1:B:989:ALA:O	1:B:993:ILE:HG12	2.19	0.42
1:C:985:ASP:OD1	1:C:985:ASP:N	2.44	0.42
1:B:327:VAL:HB	1:B:531:THR:HG23	2.01	0.42
1:B:402:ILE:HD12	1:B:402:ILE:HA	1.85	0.42
1:C:287:ASP:OD1	1:C:288:ALA:N	2.51	0.42
1:C:884:SER:OG	1:C:887:THR:OG1	2.28	0.42
1:A:47:VAL:HG12	1:C:569:ILE:HA	2.01	0.42
1:A:882:ILE:HD12	1:A:882:ILE:HA	1.91	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:B:303:LEU:HD23	1:B:308:VAL:HG22	2.00	0.42
1:B:592:PHE:HZ	1:C:740:MET:HE2	1.83	0.42
1:B:712:ILE:HG12	1:B:1094:VAL:HG11	2.00	0.42
1:A:401:VAL:HG22	1:A:509:ARG:HG2	2.01	0.42
1:C:332:ILE:HG23	1:C:362:VAL:HG11	2.01	0.42
1:A:396:TYR:HB2	1:A:514:SER:HB3	2.00	0.42
1:C:97:LYS:HD3	1:C:186:PHE:HE1	1.84	0.42
1:C:326:ILE:HD13	1:C:533:LEU:HD12	2.01	0.42
1:C:404:GLY:HA2	1:C:407:VAL:HG13	2.00	0.42
1:B:557:LYS:HE2	1:B:559:PHE:HE1	1.84	0.42
1:C:984:LEU:HD12	1:C:988:GLU:HB2	2.01	0.42
1:A:183:GLN:HG3	1:A:187:LYS:HG3	2.00	0.42
1:A:287:ASP:OD2	1:A:288:ALA:N	2.52	0.42
1:A:287:ASP:OD2	1:A:288:ALA:N	2.52	0.42
1:B:1010:GLN:O	1:B:1014:ARG:HG3	2.19	0.42
1:B:303:LEU:HD23	1:B:308:VAL:HG22	2.02	0.42
1:C:796:ASP:OD1	1:C:796:ASP:N	2.49	0.42
1:C:1077:THR:OG1	1:C:1078:ALA:N	2.52	0.42
1:C:560:LEU:HD12	1:C:561:PRO:HD2	2.01	0.42
1:A:328:ARG:NH1	1:A:531:THR:O	2.41	0.42
1:A:435:ALA:HB2	1:A:510:VAL:HG22	2.01	0.42
1:A:902:MET:HE1	1:A:1050:MET:SD	2.58	0.42
1:C:784:GLN:HA	1:C:784:GLN:NE2	2.35	0.42
1:A:355:ARG:HD2	1:A:396:TYR:HB3	2.01	0.42
1:B:738:CYS:O	1:B:742:ILE:HB	2.19	0.42
1:A:870:ILE:O	1:A:874:THR:HG23	2.20	0.42
1:B:189:LEU:HD11	1:B:208:THR:HB	2.01	0.42
1:B:366:SER:O	1:B:370:ASN:HB2	2.19	0.42
1:C:28:TYR:HB3	1:C:61:ASN:HB3	1.99	0.42
1:A:287:ASP:OD1	1:A:288:ALA:N	2.52	0.42
1:A:380:TYR:HE1	1:A:433:VAL:HG12	1.84	0.42
1:B:822:LEU:HD23	1:B:1056:ALA:HB2	2.01	0.42
1:C:762:GLN:HA	1:C:765:ARG:HG2	2.00	0.42
1:B:309:GLU:O	1:B:313:TYR:OH	2.34	0.42
1:B:462:LYS:HE2	1:B:462:LYS:H	1.84	0.42
1:B:756:TYR:HB3	1:B:759:PHE:HD2	1.84	0.42
1:B:817:PHE:HE2	1:B:935:GLN:HE21	1.67	0.42
1:B:902:MET:HE1	1:B:1050:MET:HE2	2.01	0.42
1:C:954:GLN:OE1	1:C:1014:ARG:NH1	2.52	0.42
1:A:719:THR:HG23	1:A:1068:VAL:HB	2.00	0.42
1:A:883:THR:HB	1:C:705:VAL:HG11	2.01	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:C:404:GLY:HA2	1:C:407:VAL:HG13	2.00	0.42
1:C:989:ALA:O	1:C:993:ILE:HG12	2.19	0.42
1:A:806:LEU:HD13	1:A:807:PRO:HD2	2.00	0.42
1:A:904:TYR:HE1	1:C:1094:VAL:HG22	1.84	0.42
1:C:36:VAL:HG11	1:C:220:PHE:CZ	2.54	0.42
1:C:141:LEU:HD13	1:C:246:ILE:HD13	2.01	0.42
1:A:365:TYR:H	1:A:527:PRO:HD3	1.85	0.42
1:B:716:THR:N	1:B:1071:GLN:O	2.52	0.42
1:B:778:THR:HA	1:B:781:VAL:HG12	2.00	0.42
1:A:287:ASP:OD2	1:A:288:ALA:N	2.52	0.42
1:C:564:GLN:O	1:C:577:ARG:N	2.51	0.42
1:A:712:ILE:HG21	1:A:1096:VAL:HG12	2.01	0.42
1:B:733:LYS:HB3	1:B:861:LEU:HD23	2.01	0.42
1:B:785:VAL:HG12	1:B:787:GLN:H	1.84	0.42
1:C:87:ASN:OD1	1:C:87:ASN:N	2.46	0.42
1:C:331:ASN:OD1	1:C:331:ASN:N	2.44	0.42
1:A:1050:MET:HE2	1:A:1050:MET:HB3	1.69	0.42
1:A:1105:THR:HG21	1:A:1110:TYR:HD2	1.84	0.42
1:C:986:PRO:O	1:C:990:GLU:HG2	2.19	0.42
1:B:989:ALA:O	1:B:993:ILE:HG12	2.19	0.42
1:B:802:PHE:HB3	1:B:806:LEU:HD23	2.00	0.42
1:C:106:PHE:O	1:C:117:LEU:N	2.46	0.42
1:C:882:ILE:HD12	1:C:882:ILE:HA	1.87	0.42
1:A:106:PHE:HB3	1:A:235:ILE:HD12	2.00	0.42
1:A:287:ASP:OD1	1:A:288:ALA:N	2.53	0.42
1:B:393:THR:OG1	1:B:516:GLU:O	2.34	0.42
1:B:407:VAL:O	1:B:410:ILE:HG22	2.20	0.42
1:A:897:PRO:HD2	1:C:712:ILE:HD13	2.01	0.42
1:B:188:ASN:OD1	1:B:207:HIS:NE2	2.45	0.42
1:B:992:GLN:N	1:B:992:GLN:OE1	2.51	0.42
1:C:276:LEU:HD11	1:C:304:LYS:HA	2.02	0.42
1:C:749:CYS:O	1:C:753:LEU:HB2	2.19	0.42
1:C:758:SER:O	1:C:762:GLN:HG3	2.20	0.42
1:C:884:SER:OG	1:C:894:LEU:O	2.37	0.42
1:C:497:PHE:CE2	1:C:507:PRO:HB3	2.53	0.42
1:C:1001:LEU:HD23	1:C:1001:LEU:HA	1.92	0.42
1:A:395:VAL:HG12	1:A:515:PHE:HB3	2.01	0.42
1:A:1012:LEU:HB3	1:C:1013:ILE:HD13	2.02	0.42
1:B:453:TYR:HD1	1:B:495:TYR:CZ	2.37	0.42
1:B:720:ILE:HD11	1:B:1065:VAL:HG12	2.01	0.42
1:A:883:THR:HB	1:C:705:VAL:HG11	2.01	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:B:468:ILE:HB	1:C:115:GLN:HE22	1.84	0.42
1:C:331:ASN:OD1	1:C:331:ASN:N	2.44	0.42
1:A:785:VAL:HG12	1:A:787:GLN:H	1.83	0.42
1:C:353:TRP:HE1	1:C:466:ARG:HB2	1.83	0.42
1:C:574:ASP:OD1	1:C:574:ASP:N	2.52	0.42
1:A:406:GLU:N	1:A:406:GLU:OE1	2.52	0.42
1:A:788:ILE:HG23	1:A:876:ALA:HB2	2.01	0.42
1:B:557:LYS:HE2	1:B:559:PHE:HE1	1.85	0.42
1:C:194:PHE:HB3	1:C:201:PHE:CE2	2.54	0.42
1:C:382:VAL:HG21	1:C:515:PHE:HZ	1.82	0.42
1:C:1033:VAL:HG21	1:C:1053:PRO:HG3	2.00	0.42
1:B:44:ARG:HB2	1:B:279:TYR:CD2	2.54	0.42
1:B:726:ILE:HG22	1:B:1061:VAL:HG22	2.01	0.42
1:C:642:VAL:HG22	1:C:651:ILE:HG22	2.01	0.42
1:C:931:ILE:HA	1:C:934:ILE:HG22	2.01	0.42
1:A:52:GLN:OE1	1:A:274:THR:OG1	2.27	0.42
1:C:106:PHE:HD1	1:C:235:ILE:HG21	1.84	0.42
1:C:877:LEU:HD21	1:C:1029:MET:HE1	2.01	0.42
1:A:295:PRO:HA	1:A:298:GLU:HB2	2.01	0.42
1:A:731:MET:H	1:A:774:GLN:NE2	2.17	0.42
1:A:642:VAL:HG22	1:A:651:ILE:HG22	2.01	0.42
1:B:701:VAL:HG13	1:C:787:GLN:HG2	2.01	0.42
1:B:971:GLY:O	1:B:995:ARG:NH2	2.44	0.42
1:C:33:THR:HB	1:C:220:PHE:HD1	1.84	0.42
1:B:206:LYS:HE3	1:B:206:LYS:HB3	1.81	0.42
1:B:717:ASN:OD1	1:B:718:PHE:N	2.48	0.42
1:C:717:ASN:OD1	1:C:718:PHE:N	2.50	0.42
1:A:989:ALA:O	1:A:993:ILE:HG12	2.19	0.42
1:B:382:VAL:HG11	1:B:515:PHE:HZ	1.83	0.42
1:B:406:GLU:HB3	1:B:418:ILE:HG21	2.00	0.42
1:B:858:LEU:HD22	1:B:959:LEU:HD12	2.01	0.42
1:A:293:LEU:HD23	1:A:294:ASP:HB3	2.02	0.42
1:A:614:GLY:N	1:A:647:ALA:O	2.42	0.42
1:A:787:GLN:HG2	1:C:701:VAL:HG13	2.02	0.42
1:C:48:LEU:HD11	1:C:278:LYS:HZ3	1.84	0.42
1:A:715:PRO:HD3	1:B:894:LEU:HD11	2.01	0.42
1:A:740:MET:HE1	1:A:857:GLY:HA3	2.02	0.42
1:A:867:ASP:OD1	1:A:867:ASP:N	2.52	0.42
1:B:395:VAL:HG12	1:B:515:PHE:HB3	2.00	0.42
1:B:884:SER:HA	1:B:896:ILE:HG22	2.00	0.42
1:C:712:ILE:HD12	1:C:712:ILE:HA	1.70	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:C:912:THR:O	1:C:915:VAL:HG22	2.19	0.42
1:A:537:LYS:HE3	1:A:537:LYS:HB3	1.92	0.42
1:B:565:PHE:O	1:C:43:PHE:HB3	2.19	0.42
1:A:212:LEU:HD13	1:A:217:PRO:HD3	2.01	0.42
1:A:1039:ARG:NH1	1:C:1039:ARG:HH12	2.18	0.42
1:C:795:LYS:HB3	1:C:797:PHE:CE2	2.55	0.42
1:A:331:ASN:OD1	1:A:331:ASN:N	2.44	0.42
1:A:578:ASP:OD2	1:A:581:THR:OG1	2.31	0.42
1:A:592:PHE:HZ	1:B:740:MET:HE3	1.84	0.42
1:A:962:LEU:HD12	1:A:962:LEU:HA	1.90	0.42
1:B:722:VAL:HG22	1:B:930:ALA:HB1	2.00	0.42
1:B:934:ILE:HD12	1:B:934:ILE:HA	1.76	0.42
1:A:457:ARG:HH12	1:A:461:LEU:HD23	1.84	0.42
1:B:740:MET:HE2	1:B:740:MET:HB2	1.89	0.42
1:B:1128:VAL:HG11	1:C:918:GLU:HG3	2.02	0.42
1:C:906:PHE:CD2	1:C:916:LEU:HB2	2.54	0.42
1:A:870:ILE:O	1:A:874:THR:HG23	2.19	0.42
1:C:906:PHE:CD1	1:C:916:LEU:HB2	2.54	0.42
1:A:984:LEU:HD12	1:A:988:GLU:HB2	2.02	0.42
1:B:978:ASN:HA	1:B:981:LEU:HG	2.02	0.42
1:C:762:GLN:HB3	1:C:765:ARG:HH21	1.84	0.42
1:A:894:LEU:HD23	1:C:713:ALA:HB3	2.01	0.42
1:C:403:ARG:NH1	1:C:405:ASP:OD2	2.52	0.42
1:C:877:LEU:O	1:C:881:THR:OG1	2.32	0.42
1:C:332:ILE:HG13	1:C:362:VAL:HG22	2.01	0.42
1:C:342:PHE:CE1	1:C:511:VAL:HG11	2.55	0.42
1:B:552:LEU:HD13	1:B:587:ILE:HD13	2.01	0.42
1:A:806:LEU:HD22	1:A:878:LEU:HD21	2.00	0.42
1:B:149:ASN:HB2	1:B:153:MET:HE1	2.01	0.42
1:C:200:TYR:HA	1:C:230:PRO:HA	2.02	0.42
1:B:117:LEU:HB2	1:B:233:ILE:HD11	2.02	0.42
1:A:950:ASP:O	1:A:954:GLN:HB3	2.19	0.42
1:B:715:PRO:HD3	1:C:894:LEU:HD21	2.01	0.42
1:A:106:PHE:HB3	1:A:235:ILE:HG12	2.02	0.42
1:A:574:ASP:O	1:A:587:ILE:N	2.39	0.42
1:C:722:VAL:HG22	1:C:930:ALA:HB1	2.01	0.42
1:C:882:ILE:HD12	1:C:882:ILE:HA	1.85	0.42
1:A:1089:PHE:HE2	1:B:917:TYR:HD2	1.67	0.42
1:B:317:ASN:HD21	1:B:592:PHE:HD2	1.66	0.42
1:B:552:LEU:HD12	1:B:585:LEU:HB2	2.02	0.42
1:B:574:ASP:OD1	1:B:574:ASP:N	2.46	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:B:989:ALA:O	1:B:993:ILE:HG12	2.20	0.42
1:C:717:ASN:HB3	1:C:1071:GLN:HG2	2.01	0.42
1:A:908:GLY:O	1:A:1038:LYS:NZ	2.30	0.42
1:B:431:GLY:HA2	1:B:515:PHE:CE2	2.54	0.42
1:A:1129:VAL:HG23	1:A:1132:ILE:HB	2.01	0.42
1:B:594:GLY:H	1:B:613:GLN:NE2	2.18	0.42
1:C:220:PHE:HE2	1:C:285:ILE:HG22	1.84	0.42
1:C:1116:THR:OG1	1:C:1118:ASP:OD1	2.37	0.42
1:A:406:GLU:N	1:A:406:GLU:OE1	2.52	0.42
1:A:809:PRO:HA	1:A:814:LYS:HD2	2.01	0.42
1:B:44:ARG:HD3	1:B:279:TYR:HE2	1.83	0.42
1:A:1086:LYS:HB3	1:A:1086:LYS:HE3	1.87	0.42
1:B:400:PHE:HE2	1:B:402:ILE:HD13	1.84	0.42
1:B:418:ILE:O	1:B:422:ASN:HB2	2.20	0.42
1:B:769:GLY:HA2	1:B:772:VAL:HG12	2.01	0.42
1:A:543:PHE:HD2	1:A:576:VAL:HG11	1.84	0.42
1:B:332:ILE:HG13	1:B:362:VAL:HG22	2.00	0.42
1:C:796:ASP:OD1	1:C:796:ASP:N	2.49	0.42
1:A:328:ARG:NH1	1:A:531:THR:O	2.44	0.42
1:A:598:ILE:HD11	1:A:678:ILE:HD13	2.01	0.42
1:A:989:ALA:O	1:A:993:ILE:HG12	2.19	0.42
1:C:394:ASN:OD1	1:C:394:ASN:N	2.52	0.42
1:C:730:SER:OG	1:C:731:MET:N	2.52	0.42
1:A:1025:ALA:O	1:A:1029:MET:HB2	2.20	0.42
1:C:574:ASP:N	1:C:574:ASP:OD1	2.52	0.42
1:A:175:PHE:HE2	1:A:177:MET:HG3	1.85	0.42
1:B:462:LYS:HE2	1:B:462:LYS:H	1.85	0.42
1:C:48:LEU:HB3	1:C:276:LEU:HD11	2.01	0.42
1:C:796:ASP:OD1	1:C:796:ASP:N	2.50	0.42
1:A:57:PRO:HB3	1:A:273:ARG:HH12	1.84	0.42
1:A:97:LYS:HD2	1:A:97:LYS:HA	1.87	0.42
1:A:547:THR:O	1:B:978:ASN:ND2	2.52	0.42
1:B:332:ILE:HG13	1:B:362:VAL:HG22	2.01	0.42
1:C:52:GLN:HE22	1:C:273:ARG:N	2.18	0.42
1:C:222:ALA:HB2	1:C:285:ILE:HB	2.02	0.42
1:C:398:ASP:OD1	1:C:423:TYR:OH	2.31	0.42
1:A:206:LYS:HG2	1:A:224:GLU:H	1.83	0.42
1:A:642:VAL:HG22	1:A:651:ILE:HG22	2.02	0.42
1:C:287:ASP:OD1	1:C:288:ALA:N	2.52	0.42
1:A:905:ARG:HE	1:A:1050:MET:HE2	1.83	0.42
1:B:106:PHE:HB3	1:B:235:ILE:HD12	2.02	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:C:871:ALA:HA	1:C:874:THR:HG22	2.01	0.42
1:B:350:VAL:HG21	1:B:418:ILE:HD11	2.01	0.42
1:C:86:PHE:HE2	1:C:90:VAL:HG22	1.84	0.42
1:A:426:PRO:HG2	1:A:429:PHE:HB2	2.01	0.42
1:A:870:ILE:O	1:A:874:THR:HG23	2.20	0.42
1:B:796:ASP:OD2	1:B:796:ASP:N	2.53	0.42
1:B:1115:ILE:HG22	1:B:1137:VAL:HG13	2.01	0.42
1:C:445:VAL:HG22	1:C:499:PRO:HG2	2.02	0.42
1:C:825:LYS:HE2	1:C:825:LYS:HB2	1.81	0.42
1:A:776:LYS:NZ	1:A:1019:ARG:HH21	2.18	0.42
1:B:1039:ARG:NE	1:C:1031:GLU:OE1	2.40	0.42
1:A:997:ILE:O	1:A:1001:LEU:HB3	2.19	0.42
1:C:366:SER:O	1:C:370:ASN:HB2	2.20	0.42
1:A:788:ILE:HG13	1:A:876:ALA:HB2	2.02	0.42
1:B:1046:GLY:HA2	1:C:890:ALA:HA	2.02	0.42
1:C:63:THR:HB	1:C:267:VAL:HG13	2.02	0.42
1:C:402:ILE:HD11	1:C:418:ILE:HG13	2.00	0.42
1:C:462:LYS:H	1:C:462:LYS:HD2	1.85	0.42
1:A:115:GLN:NE2	1:C:468:ILE:HG12	2.34	0.42
1:B:102:ARG:HD2	1:B:141:LEU:HD22	2.02	0.42
1:B:958:ALA:O	1:B:961:THR:OG1	2.33	0.42
1:C:441:LEU:HB3	1:C:509:ARG:HH22	1.85	0.42
1:B:279:TYR:HE1	1:B:285:ILE:HG12	1.85	0.42
1:B:435:ALA:HB2	1:B:510:VAL:HG22	2.02	0.42
1:B:983:ARG:HE	1:B:983:ARG:HB2	1.53	0.42
1:A:278:LYS:HB2	1:A:306:PHE:CE2	2.54	0.42
1:B:319:ARG:HH22	1:C:745:ASP:H	1.66	0.42
1:A:43:PHE:HB3	1:C:565:PHE:O	2.18	0.42
1:A:426:PRO:HD3	1:A:463:PRO:HB3	2.02	0.42
1:A:764:ASN:O	1:A:768:THR:HG23	2.18	0.42
1:A:43:PHE:HE2	1:A:283:GLY:HA3	1.85	0.42
1:A:206:LYS:NZ	1:A:208:THR:HG22	2.34	0.42
1:B:91:TYR:HD1	1:B:193:VAL:HG22	1.84	0.42
1:C:1086:LYS:HA	1:C:1125:ASN:HA	2.01	0.42
1:A:369:TYR:CE2	1:C:416:GLY:HA2	2.54	0.42
1:C:439:ASN:HD21	1:C:499:PRO:HA	1.84	0.42
1:C:611:LEU:HD13	1:C:678:ILE:HD11	2.01	0.42
1:A:1033:VAL:HA	1:A:1051:SER:HB2	2.02	0.42
1:B:774:GLN:HA	1:B:777:ASN:OD1	2.20	0.42
1:B:1002:GLN:O	1:B:1005:GLN:NE2	2.52	0.42
1:C:33:THR:HB	1:C:220:PHE:HD1	1.85	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:312:ILE:HD13	1:A:598:ILE:HD13	2.01	0.42
1:A:735:SER:HB2	1:A:861:LEU:HD21	2.01	0.42
1:B:462:LYS:N	1:B:465:GLU:OE2	2.38	0.42
1:B:1120:THR:OG1	1:B:1121:PHE:N	2.51	0.42
1:A:314:GLN:HE22	1:A:594:GLY:HA3	1.84	0.42
1:A:406:GLU:N	1:A:406:GLU:OE1	2.53	0.42
1:C:1114:ILE:HD13	1:C:1114:ILE:HA	1.92	0.42
1:A:787:GLN:HG2	1:C:701:VAL:HG13	2.02	0.42
1:B:902:MET:HE1	1:B:1050:MET:HE2	2.01	0.42
1:A:905:ARG:HB3	1:A:1049:LEU:HD12	2.00	0.42
1:B:733:LYS:HB3	1:B:861:LEU:HD23	2.00	0.42
1:C:409:GLN:HA	1:C:414:GLN:HG3	2.02	0.42
1:B:315:THR:OG1	1:B:316:SER:N	2.52	0.42
1:B:513:LEU:HD12	1:B:513:LEU:HA	1.87	0.42
1:B:1116:THR:OG1	1:B:1118:ASP:OD2	2.36	0.42
1:C:722:VAL:HG22	1:C:930:ALA:HB1	2.01	0.42
1:C:816:SER:N	1:C:819:GLU:OE1	2.36	0.42
1:C:1051:SER:OG	1:C:1064:HIS:ND1	2.35	0.42
1:C:212:LEU:HD13	1:C:217:PRO:HG3	2.02	0.42
1:C:332:ILE:HG13	1:C:362:VAL:HG22	2.01	0.42
1:C:796:ASP:OD1	1:C:796:ASP:N	2.50	0.42
1:A:970:PHE:HD2	1:A:996:LEU:HA	1.83	0.42
1:B:720:ILE:HD11	1:B:1065:VAL:HG12	2.00	0.42
1:B:802:PHE:HE1	1:B:927:PHE:HE2	1.67	0.42
1:B:370:ASN:HD22	1:B:384:PRO:HB2	1.84	0.42
1:C:1047:TYR:HB2	1:C:1067:TYR:HB3	2.02	0.42
1:B:182:LYS:HD3	1:B:187:LYS:HD2	2.02	0.42
1:A:289:VAL:HG11	1:A:300:LYS:HB2	2.01	0.42
1:A:129:LYS:NZ	1:A:166:CYS:SG	2.70	0.42
1:A:636:TYR:HB3	1:A:651:ILE:HD12	2.00	0.42
1:A:314:GLN:HE22	1:A:594:GLY:HA3	1.84	0.42
1:A:375:SER:OG	1:A:435:ALA:O	2.35	0.42
1:A:537:LYS:HE3	1:A:537:LYS:HB3	1.91	0.42
1:A:1041:ASP:OD1	1:B:1030:SER:HB2	2.20	0.42
1:B:802:PHE:HB3	1:B:806:LEU:HD23	2.02	0.42
1:C:108:THR:O	1:C:237:ARG:NH2	2.45	0.42
1:C:203:ILE:HD13	1:C:227:VAL:HG13	2.01	0.42
1:A:408:ARG:HH11	1:A:414:GLN:HE21	1.66	0.42
1:B:15:CYS:SG	1:B:137:ASN:N	2.91	0.42
1:C:86:PHE:CZ	1:C:89:GLY:HA2	2.55	0.42
1:B:295:PRO:O	1:B:299:THR:HG23	2.19	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:B:418:ILE:HD12	1:B:422:ASN:HD22	1.85	0.42
1:C:291:CYS:HB3	1:C:301:CYS:HB2	1.34	0.42
1:C:862:PRO:HA	1:C:863:PRO:HD3	1.93	0.42
1:C:994:ASP:O	1:C:998:THR:HG23	2.20	0.42
1:A:18:PHE:HB2	1:A:21:ARG:HB2	2.02	0.42
1:A:864:LEU:HA	1:C:667:GLY:HA2	2.00	0.42
1:A:882:ILE:HD12	1:A:882:ILE:HA	1.89	0.42
1:B:362:VAL:HG21	1:B:526:GLY:H	1.85	0.42
1:A:569:ILE:HG13	1:B:47:VAL:HG12	2.00	0.42
1:A:884:SER:OG	1:A:887:THR:OG1	2.35	0.42
1:C:402:ILE:HD11	1:C:418:ILE:HG13	2.01	0.42
1:A:785:VAL:HG12	1:A:787:GLN:H	1.83	0.42
1:A:984:LEU:HB2	1:A:989:ALA:HB2	2.02	0.42
1:B:731:MET:H	1:B:774:GLN:NE2	2.18	0.42
1:B:735:SER:HB2	1:B:861:LEU:HD21	2.02	0.42
1:C:877:LEU:O	1:C:881:THR:OG1	2.31	0.42
1:C:994:ASP:O	1:C:998:THR:HG23	2.20	0.42
1:B:715:PRO:HD3	1:C:894:LEU:HD11	2.01	0.42
1:A:43:PHE:HB3	1:C:565:PHE:O	2.19	0.42
1:A:983:ARG:HH11	1:C:517:LEU:HD11	1.84	0.42
1:B:1039:ARG:HD2	1:B:1042:PHE:HB2	2.02	0.42
1:A:414:GLN:HG3	1:A:415:THR:H	1.83	0.42
1:A:1010:GLN:CD	1:A:1014:ARG:HH22	2.28	0.42
1:C:884:SER:OG	1:C:887:THR:OG1	2.30	0.42
1:A:273:ARG:HD2	1:A:292:ALA:HB3	2.01	0.42
1:A:442:ASP:OD2	1:A:509:ARG:NE	2.45	0.42
1:B:407:VAL:O	1:B:410:ILE:HG22	2.20	0.42
1:A:882:ILE:HD12	1:A:882:ILE:HA	1.78	0.42
1:C:426:PRO:HD3	1:C:463:PRO:HB3	2.01	0.42
1:C:1105:THR:HG21	1:C:1110:TYR:HD2	1.85	0.42
1:A:745:ASP:OD1	1:A:745:ASP:N	2.53	0.42
1:A:986:PRO:O	1:A:990:GLU:HB2	2.19	0.42
1:B:733:LYS:HB3	1:B:861:LEU:HD23	2.02	0.42
1:A:193:VAL:HG23	1:A:270:LEU:HD21	2.02	0.42
1:B:455:LEU:HG	1:B:456:PHE:CD2	2.54	0.42
1:B:934:ILE:HD12	1:B:934:ILE:HA	1.80	0.42
1:C:105:ILE:HB	1:C:239:GLN:HB3	2.02	0.42
1:A:475:ALA:HB3	1:A:487:ASN:HB3	2.01	0.42
1:A:674:CYS:HB2	1:A:697:MET:HE3	2.00	0.42
1:A:1050:MET:HE2	1:A:1050:MET:HB3	1.73	0.42
1:A:586:ASP:OD1	1:A:586:ASP:N	2.51	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:606:ASN:C	1:A:606:ASN:ND2	2.78	0.42
1:B:195:LYS:HE3	1:B:197:ILE:HD12	2.02	0.42
1:B:985:ASP:HB2	1:B:986:PRO:HD2	2.02	0.42
1:C:642:VAL:HG22	1:C:651:ILE:HG22	2.02	0.42
1:A:740:MET:HE2	1:C:592:PHE:CE2	2.55	0.42
1:B:738:CYS:HB2	1:B:763:LEU:HD11	2.00	0.42
1:B:989:ALA:O	1:B:993:ILE:HG12	2.20	0.42
1:C:434:ILE:HB	1:C:511:VAL:HG13	2.02	0.42
1:C:722:VAL:HG22	1:C:930:ALA:HB1	2.01	0.42
1:A:61:ASN:C	1:A:61:ASN:ND2	2.76	0.42
1:A:205:SER:HB3	1:A:226:LEU:HD22	2.02	0.42
1:A:1005:GLN:HE22	1:B:1005:GLN:HG2	1.85	0.42
1:C:130:VAL:HB	1:C:168:PHE:HB3	2.02	0.42
1:A:406:GLU:N	1:A:406:GLU:OE1	2.53	0.42
1:A:781:VAL:HG13	1:A:782:PHE:HD1	1.84	0.42
1:A:957:GLN:HE21	1:B:765:ARG:NH1	2.18	0.42
1:B:310:LYS:HG3	1:B:600:PRO:HA	2.01	0.42
1:B:1039:ARG:HD2	1:B:1042:PHE:HB2	2.01	0.42
1:C:129:LYS:NZ	1:C:167:THR:H	2.16	0.42
1:A:1039:ARG:HG3	1:B:1031:GLU:CD	2.45	0.42
1:B:33:THR:OG1	1:B:34:ARG:NH1	2.53	0.42
1:B:326:ILE:HD13	1:B:533:LEU:HD12	2.02	0.42
1:C:915:VAL:HG21	1:C:1108:ASN:HB3	2.00	0.42
1:A:730:SER:OG	1:A:731:MET:N	2.51	0.42
1:A:759:PHE:HA	1:A:762:GLN:HE21	1.84	0.42
1:C:1012:LEU:HD13	1:C:1012:LEU:HA	1.89	0.42
1:A:415:THR:OG1	1:B:369:TYR:OH	2.26	0.42
1:A:68:ILE:H	1:A:78:ARG:HB2	1.84	0.42
1:A:742:ILE:HD12	1:A:742:ILE:HA	1.80	0.42
1:A:58:PHE:CD2	1:A:290:ASP:HB2	2.55	0.42
1:A:550:GLY:HA2	1:A:589:PRO:HA	2.02	0.42
1:B:276:LEU:HG	1:B:306:PHE:HE1	1.85	0.42
1:B:788:ILE:HG23	1:B:876:ALA:HB2	2.02	0.42
1:C:796:ASP:OD1	1:C:796:ASP:N	2.50	0.42
1:B:339:GLY:O	1:B:343:ASN:HB2	2.19	0.42
1:B:731:MET:H	1:B:774:GLN:HE22	1.67	0.42
1:C:532:ASN:OD1	1:C:533:LEU:N	2.53	0.42
1:C:774:GLN:HA	1:C:777:ASN:ND2	2.35	0.42
1:B:38:TYR:HE2	1:B:224:GLU:HG2	1.85	0.42
1:B:903:ALA:HB1	1:B:913:GLN:HG2	2.01	0.42
1:C:131:CYS:HB2	1:C:133:PHE:CE1	2.55	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:C:898:PHE:HA	1:C:901:GLN:HB2	2.02	0.42
1:C:985:ASP:OD1	1:C:988:GLU:HG3	2.20	0.42
1:A:308:VAL:H	1:A:602:THR:HG1	1.67	0.42
1:B:1002:GLN:O	1:B:1006:THR:HG23	2.19	0.42
1:A:398:ASP:OD2	1:A:398:ASP:N	2.53	0.42
1:A:701:VAL:HG13	1:B:787:GLN:HG2	2.01	0.42
1:B:202:LYS:HD2	1:B:202:LYS:HA	1.89	0.42
1:B:574:ASP:OD2	1:B:574:ASP:N	2.53	0.42
1:A:756:TYR:HB3	1:A:759:PHE:CD2	2.55	0.42
1:A:809:PRO:HA	1:A:814:LYS:HD2	2.00	0.42
1:B:360:ASN:O	1:B:360:ASN:ND2	2.34	0.42
1:A:222:ALA:HB2	1:A:285:ILE:HB	2.02	0.42
1:B:733:LYS:HB3	1:B:861:LEU:HD23	2.02	0.42
1:B:884:SER:OG	1:B:894:LEU:O	2.37	0.42
1:C:796:ASP:OD1	1:C:796:ASP:N	2.50	0.42
1:A:326:ILE:HD11	1:A:534:VAL:HG22	2.01	0.42
1:A:401:VAL:HG22	1:A:509:ARG:HG2	2.00	0.42
1:B:1106:GLN:OE1	1:B:1106:GLN:N	2.49	0.42
1:C:1009:THR:O	1:C:1013:ILE:HG12	2.20	0.42
1:A:1107:ARG:HH21	1:B:896:ILE:HD11	1.85	0.42
1:B:44:ARG:HH22	1:B:49:HIS:CG	2.37	0.42
1:B:538:CYS:HB3	1:B:590:CYS:HB3	1.52	0.42
1:C:195:LYS:HE3	1:C:195:LYS:HB3	1.88	0.42
1:A:1105:THR:HG21	1:A:1110:TYR:HD1	1.85	0.42
1:C:1028:LYS:NZ	1:C:1042:PHE:O	2.53	0.42
1:A:738:CYS:HB3	1:A:763:LEU:HD11	2.02	0.42
1:B:223:LEU:HD12	1:B:223:LEU:HA	1.92	0.42
1:A:791:THR:HG21	1:A:806:LEU:HD11	2.02	0.42
1:A:409:GLN:HA	1:A:414:GLN:HG2	2.01	0.42
1:A:809:PRO:HA	1:A:814:LYS:HD2	2.00	0.42
1:B:44:ARG:HD2	1:B:279:TYR:HE2	1.85	0.42
1:C:796:ASP:OD1	1:C:796:ASP:N	2.50	0.42
1:A:47:VAL:HG12	1:C:569:ILE:HA	2.02	0.42
1:A:815:ARG:NH1	1:A:867:ASP:OD1	2.50	0.42
1:B:44:ARG:HB3	1:B:279:TYR:HD2	1.85	0.42
1:C:882:ILE:HD12	1:C:882:ILE:HA	1.88	0.42
1:A:909:ILE:HA	1:A:1038:LYS:NZ	2.35	0.42
1:C:319:ARG:NH2	1:C:590:CYS:SG	2.93	0.42
1:C:980:ILE:HD13	1:C:980:ILE:HA	1.90	0.42
1:A:592:PHE:CE2	1:B:740:MET:HG3	2.55	0.42
1:A:890:ALA:HA	1:C:1046:GLY:HA2	2.01	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:191:GLU:HG2	1:A:223:LEU:HD22	2.02	0.42
1:B:390:LEU:HG	1:B:392:PHE:HE1	1.85	0.42
1:B:106:PHE:HB3	1:B:235:ILE:HD12	2.02	0.42
1:B:999:GLY:HA2	1:B:1002:GLN:HG3	2.02	0.42
1:A:205:SER:O	1:A:206:LYS:HD2	2.20	0.42
1:B:106:PHE:HB3	1:B:235:ILE:HD13	2.02	0.42
1:C:457:ARG:NH2	1:C:461:LEU:HG	2.35	0.42
1:C:1002:GLN:O	1:C:1006:THR:HG23	2.19	0.42
1:C:822:LEU:HD13	1:C:1056:ALA:HB2	2.02	0.42
1:C:994:ASP:O	1:C:998:THR:HG23	2.20	0.42
1:C:1010:GLN:HE22	1:C:1014:ARG:HH12	1.68	0.42
1:A:972:ALA:HB2	1:A:996:LEU:HD21	2.02	0.42
1:B:462:LYS:HE2	1:B:462:LYS:H	1.85	0.42
1:C:720:ILE:HD11	1:C:1065:VAL:HG22	2.02	0.42
1:C:788:ILE:HG13	1:C:876:ALA:HB2	2.02	0.42
1:A:720:ILE:HD11	1:A:1065:VAL:HG12	2.02	0.42
1:A:58:PHE:CD2	1:A:290:ASP:HB2	2.54	0.42
1:A:331:ASN:OD1	1:A:331:ASN:N	2.44	0.42
1:A:606:ASN:C	1:A:606:ASN:ND2	2.78	0.42
1:A:1143:PRO:HA	1:A:1146:ASP:HB2	2.02	0.42
1:C:296:LEU:O	1:C:299:THR:OG1	2.35	0.42
1:B:442:ASP:OD2	1:B:509:ARG:NE	2.44	0.42
1:B:730:SER:OG	1:B:731:MET:N	2.52	0.42
1:B:884:SER:OG	1:B:887:THR:OG1	2.29	0.42
1:B:809:PRO:HA	1:B:814:LYS:HD2	2.01	0.42
1:B:983:ARG:HG3	1:B:984:LEU:HD23	2.02	0.42
1:B:994:ASP:HA	1:B:997:ILE:HG12	2.02	0.42
1:C:105:ILE:N	1:C:239:GLN:O	2.35	0.42
1:A:541:PHE:N	1:A:548:GLY:O	2.45	0.42
1:B:139:PRO:HG2	1:B:245:HIS:CE1	2.55	0.42
1:C:879:ALA:O	1:C:883:THR:OG1	2.32	0.42
1:C:1008:VAL:O	1:C:1012:LEU:HD22	2.20	0.42
1:A:774:GLN:HE22	1:A:1018:ILE:HG21	1.85	0.42
1:A:787:GLN:HG2	1:C:701:VAL:HG13	2.02	0.42
1:C:819:GLU:HA	1:C:822:LEU:HG	2.02	0.42
1:A:214:ARG:HD2	1:A:214:ARG:HA	1.84	0.42
1:A:1004:LEU:O	1:A:1008:VAL:HG12	2.20	0.42
1:B:454:ARG:HH12	1:B:457:ARG:HB2	1.84	0.42
1:C:37:TYR:HA	1:C:223:LEU:H	1.84	0.42
1:B:406:GLU:HB3	1:B:418:ILE:HG21	2.02	0.42
1:C:751:ASN:HA	1:C:754:LEU:HG	2.02	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:406:GLU:N	1:A:406:GLU:OE1	2.53	0.42
1:A:730:SER:OG	1:A:731:MET:N	2.51	0.42
1:A:915:VAL:HG11	1:A:1109:PHE:CE2	2.55	0.42
1:A:972:ALA:HB2	1:A:996:LEU:HD21	2.02	0.42
1:B:720:ILE:HD11	1:B:1065:VAL:HG12	2.01	0.42
1:A:113:LYS:HA	1:A:113:LYS:HD2	1.86	0.42
1:A:357:ARG:NH2	1:A:396:TYR:OH	2.53	0.42
1:A:393:THR:OG1	1:A:516:GLU:O	2.37	0.42
1:A:435:ALA:HB2	1:A:510:VAL:HG22	2.02	0.42
1:A:758:SER:O	1:A:762:GLN:HG3	2.20	0.42
1:B:200:TYR:HA	1:B:230:PRO:HA	2.01	0.42
1:B:901:GLN:O	1:B:905:ARG:HG2	2.20	0.42
1:C:393:THR:OG1	1:C:394:ASN:N	2.52	0.42
1:C:759:PHE:HD2	1:C:762:GLN:HE21	1.67	0.42
1:C:1002:GLN:O	1:C:1006:THR:HG23	2.19	0.42
1:A:962:LEU:HD12	1:A:962:LEU:HA	1.89	0.42
1:C:407:VAL:O	1:C:410:ILE:HG22	2.20	0.42
1:B:328:ARG:NH1	1:B:580:GLN:OE1	2.50	0.41
1:A:975:SER:O	1:A:1000:ARG:NH2	2.53	0.41
1:A:984:LEU:HB3	1:A:988:GLU:HG2	2.01	0.41
1:B:709:ASN:ND2	1:C:796:ASP:OD2	2.50	0.41
1:B:934:ILE:HD12	1:B:934:ILE:HA	1.85	0.41
1:A:737:ASP:OD2	1:A:740:MET:HB3	2.20	0.41
1:B:366:SER:O	1:B:370:ASN:HB2	2.19	0.41
1:A:64:TRP:HE1	1:A:264:ALA:HB1	1.84	0.41
1:A:1107:ARG:HE	1:A:1107:ARG:HB3	1.65	0.41
1:B:102:ARG:HD2	1:B:141:LEU:HD22	2.02	0.41
1:B:398:ASP:HB2	1:B:512:VAL:HG13	2.02	0.41
1:A:403:ARG:NH2	1:B:372:ALA:O	2.53	0.41
1:A:883:THR:HB	1:C:705:VAL:HG11	2.02	0.41
1:A:901:GLN:HE21	1:A:1050:MET:HE1	1.85	0.41
1:B:85:PRO:HA	1:B:237:ARG:HH11	1.85	0.41
1:A:406:GLU:N	1:A:406:GLU:OE1	2.53	0.41
1:C:962:LEU:HD21	1:C:1007:TYR:CG	2.55	0.41
1:A:947:LYS:HE2	1:A:947:LYS:HB2	1.95	0.41
1:B:759:PHE:HD1	1:B:762:GLN:HE21	1.67	0.41
1:B:1128:VAL:HG21	1:C:918:GLU:HG3	2.01	0.41
1:C:560:LEU:HD23	1:C:562:PHE:H	1.85	0.41
1:C:738:CYS:O	1:C:742:ILE:HB	2.20	0.41
1:A:332:ILE:HG13	1:A:362:VAL:HG22	2.02	0.41
1:A:701:VAL:HG13	1:B:787:GLN:HG2	2.02	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:B:513:LEU:HD12	1:B:513:LEU:HA	1.90	0.41
1:A:999:GLY:HA2	1:A:1002:GLN:HG3	2.01	0.41
1:B:407:VAL:O	1:B:410:ILE:HG22	2.19	0.41
1:B:972:ALA:HB2	1:B:996:LEU:HD21	2.02	0.41
1:B:993:ILE:O	1:B:997:ILE:HG12	2.20	0.41
1:B:1115:ILE:HG22	1:B:1137:VAL:HG23	2.02	0.41
1:A:994:ASP:HA	1:A:997:ILE:HG12	2.02	0.41
1:A:1135:ASN:OD1	1:A:1136:THR:N	2.43	0.41
1:A:58:PHE:HD2	1:A:290:ASP:HB2	1.84	0.41
1:A:1002:GLN:O	1:A:1006:THR:HG23	2.20	0.41
1:C:966:LEU:HA	1:C:1000:ARG:HH12	1.85	0.41
1:A:383:SER:OG	1:B:988:GLU:OE1	2.38	0.41
1:B:185:ASN:HB2	1:B:213:VAL:HA	2.02	0.41
1:C:886:TRP:HH2	1:C:904:TYR:HB3	1.85	0.41
1:B:886:TRP:HZ2	1:B:904:TYR:HD2	1.68	0.41
1:B:904:TYR:HD1	1:B:904:TYR:HA	1.61	0.41
1:B:1090:PRO:HD3	1:B:1095:PHE:HE2	1.85	0.41
1:A:105:ILE:HD12	1:A:105:ILE:HA	1.80	0.41
1:B:117:LEU:HB2	1:B:233:ILE:HD11	2.02	0.41
1:B:701:VAL:HG13	1:C:787:GLN:HG2	2.01	0.41
1:C:229:LEU:HG	1:C:231:ILE:HG23	2.02	0.41
1:A:978:ASN:HA	1:A:981:LEU:HG	2.01	0.41
1:B:701:VAL:HG13	1:C:787:GLN:HG2	2.01	0.41
1:C:586:ASP:OD1	1:C:586:ASP:N	2.49	0.41
1:C:989:ALA:O	1:C:993:ILE:HG12	2.20	0.41
1:A:1002:GLN:NE2	1:B:1005:GLN:OE1	2.50	0.41
1:B:1004:LEU:O	1:B:1008:VAL:HG12	2.20	0.41
1:C:822:LEU:HD13	1:C:1056:ALA:HB2	2.02	0.41
1:C:950:ASP:OD1	1:C:950:ASP:N	2.50	0.41
1:A:129:LYS:HE3	1:A:129:LYS:HB3	1.86	0.41
1:A:332:ILE:HG13	1:A:362:VAL:HG22	2.02	0.41
1:A:569:ILE:HD11	1:B:964:LYS:NZ	2.35	0.41
1:B:48:LEU:HD12	1:B:48:LEU:HA	1.93	0.41
1:A:1079:PRO:HB3	1:B:900:MET:HE1	2.02	0.41
1:C:185:ASN:ND2	1:C:211:ASN:OD1	2.52	0.41
1:A:332:ILE:HG13	1:A:362:VAL:HG22	2.02	0.41
1:A:170:TYR:HE2	1:A:231:ILE:HD11	1.85	0.41
1:A:295:PRO:O	1:A:299:THR:HG23	2.20	0.41
1:A:1116:THR:OG1	1:A:1118:ASP:OD1	2.37	0.41
1:B:297:SER:HA	1:B:300:LYS:HG2	2.02	0.41
1:B:426:PRO:HD3	1:B:463:PRO:HB3	2.01	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:B:58:PHE:HD2	1:B:290:ASP:HB2	1.84	0.41
1:C:1038:LYS:HA	1:C:1038:LYS:HD3	1.79	0.41
1:A:431:GLY:HA2	1:A:515:PHE:CE2	2.55	0.41
1:B:118:LEU:HD21	1:B:135:PHE:HE1	1.85	0.41
1:A:105:ILE:HD13	1:A:105:ILE:HG21	1.83	0.41
1:C:1031:GLU:HG3	1:C:1039:ARG:NH2	2.35	0.41
1:A:195:LYS:HE2	1:A:195:LYS:HB3	1.76	0.41
1:B:934:ILE:HD12	1:B:934:ILE:HA	1.74	0.41
1:C:884:SER:OG	1:C:887:THR:OG1	2.28	0.41
1:A:549:THR:HB	1:B:745:ASP:HB3	2.01	0.41
1:A:436:TRP:HZ3	1:A:511:VAL:HG12	1.85	0.41
1:C:796:ASP:OD1	1:C:796:ASP:N	2.50	0.41
1:A:457:ARG:NH2	1:A:467:ASP:OD2	2.50	0.41
1:A:332:ILE:HG13	1:A:362:VAL:HG22	2.02	0.41
1:B:196:ASN:ND2	1:B:233:ILE:O	2.42	0.41
1:B:679:GLY:HA2	1:C:864:LEU:HD23	2.02	0.41
1:C:867:ASP:N	1:C:867:ASP:OD1	2.51	0.41
1:C:871:ALA:HA	1:C:874:THR:HG22	2.02	0.41
1:A:393:THR:HB	1:A:520:ALA:HB3	2.02	0.41
1:A:396:TYR:CZ	1:B:230:PRO:HG3	2.55	0.41
1:B:187:LYS:NZ	1:B:211:ASN:OD1	2.52	0.41
1:B:751:ASN:HA	1:B:754:LEU:HG	2.01	0.41
1:B:44:ARG:HB2	1:B:279:TYR:CD2	2.55	0.41
1:A:951:VAL:HA	1:A:954:GLN:HG3	2.02	0.41
1:A:1002:GLN:O	1:A:1006:THR:HG23	2.20	0.41
1:B:212:LEU:HD13	1:B:217:PRO:HG3	2.02	0.41
1:B:365:TYR:CE1	1:B:387:LEU:HD12	2.55	0.41
1:C:366:SER:O	1:C:370:ASN:HB2	2.20	0.41
1:C:407:VAL:O	1:C:410:ILE:HG22	2.20	0.41
1:C:775:ASP:HB3	1:C:864:LEU:HD12	2.02	0.41
1:C:777:ASN:OD1	1:C:778:THR:N	2.53	0.41
1:B:1014:ARG:O	1:B:1018:ILE:HG12	2.20	0.41
1:C:48:LEU:HB3	1:C:276:LEU:HD11	2.02	0.41
1:C:393:THR:OG1	1:C:394:ASN:OD1	2.37	0.41
1:C:934:ILE:HD12	1:C:934:ILE:HA	1.89	0.41
1:A:131:CYS:HB2	1:A:133:PHE:CD1	2.55	0.41
1:A:547:THR:OG1	1:B:978:ASN:OD1	2.29	0.41
1:A:39:PRO:HG3	1:A:51:THR:HG21	2.01	0.41
1:B:1051:SER:OG	1:B:1064:HIS:ND1	2.50	0.41
1:C:537:LYS:HE3	1:C:537:LYS:HB3	1.87	0.41
1:B:404:GLY:HA2	1:B:407:VAL:HG13	2.03	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:B:901:GLN:O	1:B:905:ARG:HG2	2.21	0.41
1:B:1083:HIS:CG	1:B:1137:VAL:HG12	2.55	0.41
1:C:291:CYS:HB2	1:C:301:CYS:HB2	1.27	0.41
1:C:796:ASP:OD1	1:C:796:ASP:N	2.51	0.41
1:A:568:ASP:OD1	1:A:569:ILE:N	2.51	0.41
1:B:453:TYR:HE2	1:B:455:LEU:HB2	1.85	0.41
1:B:777:ASN:O	1:B:781:VAL:HG12	2.20	0.41
1:C:304:LYS:HA	1:C:304:LYS:HE3	2.01	0.41
1:C:728:PRO:HB3	1:C:948:LEU:HD13	2.02	0.41
1:A:222:ALA:HB2	1:A:285:ILE:HB	2.02	0.41
1:A:909:ILE:HG12	1:A:1047:TYR:HB3	2.02	0.41
1:C:912:THR:O	1:C:915:VAL:HG12	2.20	0.41
1:A:406:GLU:OE1	1:A:406:GLU:N	2.53	0.41
1:A:785:VAL:HG12	1:A:787:GLN:H	1.84	0.41
1:A:220:PHE:HE2	1:A:285:ILE:HG22	1.83	0.41
1:A:730:SER:OG	1:A:731:MET:N	2.52	0.41
1:A:809:PRO:HA	1:A:814:LYS:HD2	2.02	0.41
1:A:317:ASN:HD21	1:A:592:PHE:HD2	1.67	0.41
1:B:533:LEU:HD21	1:B:585:LEU:HD11	2.02	0.41
1:C:36:VAL:HG21	1:C:220:PHE:CZ	2.55	0.41
1:A:498:GLN:HB2	1:A:501:TYR:CE1	2.55	0.41
1:B:598:ILE:HB	1:B:609:ALA:HB3	2.01	0.41
1:B:1090:PRO:HD3	1:B:1095:PHE:HE2	1.85	0.41
1:C:272:PRO:O	1:C:273:ARG:HD3	2.20	0.41
1:C:369:TYR:CZ	1:C:384:PRO:HB2	2.56	0.41
1:A:100:ILE:HG23	1:A:247:SER:HB3	2.01	0.41
1:B:600:PRO:HD3	1:B:692:ILE:HD11	2.02	0.41
1:A:884:SER:HG	1:A:887:THR:HG1	1.63	0.41
1:A:1025:ALA:O	1:A:1029:MET:HB2	2.20	0.41
1:B:966:LEU:HD22	1:B:1000:ARG:HH12	1.84	0.41
1:B:415:THR:HG21	1:C:384:PRO:HG3	2.02	0.41
1:B:465:GLU:OE1	1:C:234:ASN:ND2	2.47	0.41
1:A:15:CYS:SG	1:A:137:ASN:N	2.88	0.41
1:A:315:THR:HB	1:A:595:VAL:HG23	2.01	0.41
1:A:317:ASN:HD21	1:A:592:PHE:HB2	1.85	0.41
1:C:737:ASP:HA	1:C:764:ASN:HD21	1.86	0.41
1:A:537:LYS:HE3	1:A:537:LYS:HB3	1.95	0.41
1:A:788:ILE:HG13	1:A:876:ALA:HB2	2.02	0.41
1:A:1050:MET:HB3	1:A:1050:MET:HE2	1.69	0.41
1:B:366:SER:O	1:B:370:ASN:HB2	2.19	0.41
1:C:223:LEU:HD12	1:C:223:LEU:HA	1.84	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:C:431:GLY:HA2	1:C:515:PHE:CZ	2.56	0.41
1:A:731:MET:H	1:A:774:GLN:NE2	2.18	0.41
1:B:600:PRO:HD3	1:B:692:ILE:HD11	2.02	0.41
1:B:390:LEU:HG	1:B:392:PHE:HE1	1.85	0.41
1:B:1013:ILE:HD13	1:C:1012:LEU:HG	2.02	0.41
1:C:223:LEU:HD12	1:C:223:LEU:HA	1.85	0.41
1:A:1090:PRO:HD3	1:A:1095:PHE:HE2	1.86	0.41
1:B:735:SER:HB2	1:B:861:LEU:HD21	2.02	0.41
1:C:758:SER:O	1:C:762:GLN:HG3	2.21	0.41
1:C:914:ASN:OD1	1:C:914:ASN:N	2.53	0.41
1:B:796:ASP:OD1	1:B:796:ASP:N	2.50	0.41
1:A:227:VAL:HG12	1:A:229:LEU:HD22	2.03	0.41
1:B:701:VAL:HG13	1:C:787:GLN:HG3	2.02	0.41
1:A:557:LYS:HE2	1:A:559:PHE:HE1	1.85	0.41
1:B:407:VAL:O	1:B:410:ILE:HG22	2.21	0.41
1:B:439:ASN:OD1	1:B:440:ASN:N	2.51	0.41
1:A:541:PHE:HD2	1:A:543:PHE:CE1	2.39	0.41
1:A:561:PRO:O	1:A:577:ARG:NH1	2.54	0.41
1:B:102:ARG:HH12	1:B:177:MET:HE3	1.84	0.41
1:C:335:LEU:HD22	1:C:364:ASP:HA	2.01	0.41
1:C:1031:GLU:HB3	1:C:1037:SER:HB2	2.01	0.41
1:A:1040:VAL:HG11	1:B:1035:GLY:HA3	2.01	0.41
1:B:189:LEU:HD12	1:B:217:PRO:HG2	2.01	0.41
1:B:884:SER:OG	1:B:894:LEU:O	2.39	0.41
1:A:1002:GLN:O	1:A:1006:THR:HG23	2.20	0.41
1:B:775:ASP:OD1	1:B:775:ASP:N	2.51	0.41
1:B:802:PHE:HE1	1:B:927:PHE:HE2	1.69	0.41
1:B:934:ILE:HD12	1:B:934:ILE:HA	1.78	0.41
1:B:1039:ARG:HD2	1:B:1042:PHE:HB2	2.01	0.41
1:C:34:ARG:NH2	1:C:221:SER:H	2.19	0.41
1:A:1081:ILE:HD11	1:A:1135:ASN:HB3	2.02	0.41
1:C:796:ASP:OD1	1:C:796:ASP:N	2.50	0.41
1:C:204:TYR:HB3	1:C:223:LEU:HG	2.02	0.41
1:A:674:CYS:HB2	1:A:697:MET:HG2	2.01	0.41
1:C:52:GLN:HE22	1:C:273:ARG:N	2.18	0.41
1:C:351:TYR:HE2	1:C:452:LEU:HB2	1.86	0.41
1:A:576:VAL:HG12	1:A:587:ILE:HD11	2.01	0.41
1:A:947:LYS:HE3	1:A:947:LYS:HB2	1.96	0.41
1:B:517:LEU:HD23	1:B:518:LEU:HB2	2.01	0.41
1:C:1086:LYS:HE3	1:C:1122:VAL:HG11	2.01	0.41
1:A:91:TYR:HB3	1:A:268:GLY:HA3	2.03	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:B:722:VAL:HG21	1:B:931:ILE:HD13	2.01	0.41
1:C:86:PHE:HB2	1:C:238:PHE:HD2	1.85	0.41
1:A:758:SER:O	1:A:762:GLN:HG3	2.20	0.41
1:A:946:GLY:HA2	1:A:949:GLN:HB3	2.02	0.41
1:C:201:PHE:HD2	1:C:229:LEU:HD22	1.85	0.41
1:C:312:ILE:HD12	1:C:598:ILE:HD12	2.01	0.41
1:A:537:LYS:HE3	1:A:537:LYS:HB3	1.95	0.41
1:C:537:LYS:HE3	1:C:537:LYS:HB3	1.89	0.41
1:A:805:ILE:HG13	1:A:1054:GLN:NE2	2.36	0.41
1:A:1039:ARG:HG3	1:B:1031:GLU:OE1	2.20	0.41
1:B:642:VAL:HG22	1:B:651:ILE:HG22	2.02	0.41
1:C:203:ILE:HD13	1:C:227:VAL:HG13	2.03	0.41
1:C:796:ASP:OD1	1:C:796:ASP:N	2.50	0.41
1:C:884:SER:OG	1:C:894:LEU:O	2.38	0.41
1:A:54:LEU:HD12	1:A:88:ASP:HB3	2.02	0.41
1:A:317:ASN:HD21	1:A:592:PHE:HD2	1.68	0.41
1:A:498:GLN:HB2	1:A:501:TYR:CE2	2.56	0.41
1:A:950:ASP:N	1:A:950:ASP:OD1	2.52	0.41
1:B:914:ASN:OD1	1:B:914:ASN:N	2.53	0.41
1:C:989:ALA:O	1:C:993:ILE:HG13	2.21	0.41
1:C:382:VAL:HG21	1:C:515:PHE:HZ	1.85	0.41
1:B:650:LEU:HD13	1:B:653:ALA:HB3	2.02	0.41
1:B:719:THR:HG23	1:B:1068:VAL:HB	2.02	0.41
1:A:365:TYR:CE2	1:A:387:LEU:HD12	2.56	0.41
1:C:105:ILE:HD11	1:C:110:LEU:HD22	2.03	0.41
1:C:1031:GLU:HG3	1:C:1039:ARG:HH21	1.84	0.41
1:B:276:LEU:HD21	1:B:301:CYS:HA	2.02	0.41
1:C:436:TRP:HZ3	1:C:511:VAL:HG12	1.85	0.41
1:C:730:SER:OG	1:C:731:MET:N	2.54	0.41
1:B:442:ASP:OD1	1:B:509:ARG:NH2	2.44	0.41
1:B:877:LEU:O	1:B:881:THR:HG23	2.20	0.41
1:B:985:ASP:O	1:B:989:ALA:HB2	2.20	0.41
1:C:106:PHE:HB3	1:C:235:ILE:HD13	2.02	0.41
1:C:856:ASN:HD22	1:C:858:LEU:HD23	1.85	0.41
1:A:298:GLU:O	1:A:302:THR:HG23	2.20	0.41
1:A:1029:MET:HE1	1:A:1053:PRO:HB3	2.02	0.41
1:B:720:ILE:HD12	1:B:720:ILE:HA	1.86	0.41
1:B:994:ASP:O	1:B:998:THR:HG23	2.21	0.41
1:B:353:TRP:HH2	1:B:464:PHE:HD1	1.67	0.41
1:B:701:VAL:HG13	1:C:787:GLN:HG3	2.02	0.41
1:B:759:PHE:HA	1:B:762:GLN:HE21	1.85	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:C:731:MET:HE2	1:C:955:ASN:HD21	1.85	0.41
1:C:796:ASP:OD1	1:C:796:ASP:N	2.50	0.41
1:C:105:ILE:HG22	1:C:239:GLN:O	2.21	0.41
1:C:1032:CYS:HA	1:C:1048:HIS:CE1	2.55	0.41
1:A:439:ASN:HD21	1:A:506:GLN:HB3	1.84	0.41
1:B:327:VAL:HB	1:B:531:THR:HG23	2.03	0.41
1:B:730:SER:OG	1:B:731:MET:N	2.53	0.41
1:B:970:PHE:HA	1:C:756:TYR:CE2	2.51	0.41
1:C:276:LEU:HD11	1:C:304:LYS:HA	2.02	0.41
1:B:332:ILE:HG13	1:B:362:VAL:HG22	2.02	0.41
1:B:369:TYR:CZ	1:B:384:PRO:HB2	2.55	0.41
1:C:392:PHE:HB2	1:C:524:VAL:HG13	2.03	0.41
1:C:906:PHE:CD2	1:C:916:LEU:HB2	2.55	0.41
1:C:906:PHE:HA	1:C:909:ILE:HG12	2.02	0.41
1:A:529:LYS:H	1:A:529:LYS:HG2	1.68	0.41
1:A:1081:ILE:HD12	1:A:1081:ILE:HA	1.83	0.41
1:B:762:GLN:HB3	1:B:765:ARG:HH21	1.84	0.41
1:A:870:ILE:O	1:A:874:THR:HG23	2.20	0.41
1:B:296:LEU:O	1:B:299:THR:OG1	2.32	0.41
1:A:960:ASN:O	1:A:964:LYS:HB2	2.20	0.41
1:B:517:LEU:HD11	1:C:983:ARG:NH2	2.35	0.41
1:A:33:THR:OG1	1:A:219:GLY:O	2.30	0.41
1:B:406:GLU:N	1:B:406:GLU:OE1	2.53	0.41
1:B:1115:ILE:HG22	1:B:1137:VAL:HG23	2.02	0.41
1:B:353:TRP:CH2	1:B:464:PHE:HA	2.54	0.41
1:B:809:PRO:HA	1:B:814:LYS:HD2	2.01	0.41
1:C:36:VAL:HG11	1:C:220:PHE:CZ	2.56	0.41
1:C:276:LEU:HD21	1:C:304:LYS:HA	2.02	0.41
1:C:796:ASP:OD1	1:C:796:ASP:N	2.50	0.41
1:C:912:THR:O	1:C:915:VAL:HG22	2.21	0.41
1:C:957:GLN:O	1:C:961:THR:HG23	2.20	0.41
1:A:895:GLN:NE2	1:C:713:ALA:HB2	2.35	0.41
1:B:884:SER:OG	1:B:887:THR:OG1	2.26	0.41
1:B:951:VAL:HA	1:B:954:GLN:HG3	2.02	0.41
1:C:222:ALA:HB2	1:C:285:ILE:HB	2.02	0.41
1:C:480:CYS:C	1:C:481:ASN:HD22	2.27	0.41
1:C:574:ASP:OD1	1:C:574:ASP:N	2.52	0.41
1:C:407:VAL:O	1:C:410:ILE:HG22	2.20	0.41
1:C:722:VAL:HG22	1:C:930:ALA:HB1	2.03	0.41
1:C:788:ILE:HG13	1:C:876:ALA:HB2	2.02	0.41
1:A:642:VAL:HG22	1:A:651:ILE:HG22	2.02	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:957:GLN:O	1:A:961:THR:HG23	2.21	0.41
1:B:970:PHE:HD2	1:B:996:LEU:HA	1.85	0.41
1:C:52:GLN:HE22	1:C:273:ARG:N	2.18	0.41
1:C:195:LYS:HE3	1:C:195:LYS:HB2	1.75	0.41
1:A:115:GLN:HB2	1:A:233:ILE:HG12	2.03	0.41
1:B:775:ASP:HB3	1:B:864:LEU:HD13	2.01	0.41
1:B:1010:GLN:HE22	1:B:1014:ARG:NH2	2.19	0.41
1:C:220:PHE:HE2	1:C:285:ILE:HG22	1.85	0.41
1:C:406:GLU:OE1	1:C:406:GLU:N	2.52	0.41
1:C:557:LYS:HE2	1:C:559:PHE:HE1	1.85	0.41
1:C:985:ASP:HB2	1:C:986:PRO:HD2	2.03	0.41
1:A:1106:GLN:OE1	1:A:1106:GLN:N	2.51	0.41
1:B:172:SER:OG	1:B:173:GLN:N	2.53	0.41
1:B:577:ARG:HG2	1:B:584:ILE:HG13	2.03	0.41
1:A:326:ILE:HD11	1:A:534:VAL:HB	2.02	0.41
1:A:1050:MET:HB2	1:A:1050:MET:HE2	1.75	0.41
1:B:728:PRO:HB2	1:B:1018:ILE:HD11	2.02	0.41
1:C:1004:LEU:O	1:C:1008:VAL:HG23	2.20	0.41
1:A:406:GLU:N	1:A:406:GLU:OE1	2.53	0.41
1:A:758:SER:O	1:A:762:GLN:HG3	2.21	0.41
1:B:105:ILE:HD12	1:B:241:LEU:HD21	2.02	0.41
1:C:402:ILE:HG23	1:C:407:VAL:HG12	2.02	0.41
1:A:826:VAL:HB	1:A:1057:PRO:HG2	2.02	0.41
1:B:431:GLY:HA2	1:B:515:PHE:CE2	2.55	0.41
1:C:642:VAL:HG22	1:C:651:ILE:HG22	2.02	0.41
1:A:168:PHE:HZ	1:C:466:ARG:HH22	1.68	0.41
1:A:821:LEU:HD23	1:A:821:LEU:HA	1.93	0.41
1:A:989:ALA:O	1:A:993:ILE:HG12	2.21	0.41
1:B:827:THR:O	1:B:827:THR:OG1	2.32	0.41
1:C:56:LEU:HD22	1:C:91:TYR:CD1	2.56	0.41
1:C:614:GLY:N	1:C:647:ALA:O	2.44	0.41
1:C:871:ALA:HA	1:C:874:THR:HG22	2.03	0.41
1:C:878:LEU:HD23	1:C:878:LEU:HA	1.90	0.41
1:B:406:GLU:N	1:B:406:GLU:OE1	2.54	0.41
1:B:574:ASP:OD1	1:B:574:ASP:N	2.45	0.41
1:B:1112:PRO:C	1:B:1113:GLN:HE21	2.29	0.41
1:C:604:THR:OG1	1:C:605:SER:N	2.53	0.41
1:C:1084:ASP:OD1	1:C:1086:LYS:NZ	2.53	0.41
1:B:303:LEU:HD23	1:B:308:VAL:HG22	2.02	0.41
1:A:99:ASN:ND2	1:A:177:MET:SD	2.94	0.41
1:B:442:ASP:OD2	1:B:509:ARG:NE	2.49	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:B:866:THR:O	1:B:870:ILE:HG12	2.21	0.41
1:C:587:ILE:HD12	1:C:587:ILE:HA	1.82	0.41
1:C:729:VAL:HG21	1:C:781:VAL:HG11	2.02	0.41
1:B:756:TYR:HE2	1:B:997:ILE:HD12	1.85	0.41
1:C:1116:THR:OG1	1:C:1118:ASP:OD2	2.36	0.41
1:A:358:ILE:HB	1:A:395:VAL:HG23	2.03	0.41
1:A:781:VAL:HG13	1:A:782:PHE:CD1	2.55	0.41
1:B:905:ARG:HH21	1:B:1050:MET:HA	1.86	0.41
1:C:731:MET:H	1:C:774:GLN:NE2	2.16	0.41
1:B:701:VAL:HG23	1:C:787:GLN:HG2	2.02	0.41
1:A:978:ASN:HA	1:A:981:LEU:HG	2.02	0.41
1:A:1051:SER:OG	1:A:1064:HIS:ND1	2.39	0.41
1:B:395:VAL:HG12	1:B:515:PHE:HB3	2.02	0.41
1:B:406:GLU:N	1:B:406:GLU:OE1	2.54	0.41
1:C:555:SER:HB2	1:C:586:ASP:OD1	2.20	0.41
1:C:720:ILE:HD11	1:C:1065:VAL:HG12	2.02	0.41
1:A:546:LEU:HD21	1:A:573:THR:HG21	2.02	0.41
1:A:976:VAL:HG13	1:A:979:ASP:HB3	2.01	0.41
1:B:774:GLN:HA	1:B:777:ASN:HD22	1.86	0.41
1:C:420:ASP:HB2	1:C:460:ASN:OD1	2.20	0.41
1:A:427:ASP:OD2	1:C:986:PRO:HB2	2.20	0.41
1:A:563:GLN:HG2	1:B:43:PHE:HB2	2.02	0.41
1:A:1010:GLN:HE22	1:A:1014:ARG:NH2	2.17	0.41
1:C:537:LYS:HE3	1:C:537:LYS:HB3	1.94	0.41
1:A:330:PRO:HD3	1:A:579:PRO:HB2	2.03	0.41
1:B:462:LYS:HE2	1:B:462:LYS:H	1.86	0.41
1:C:914:ASN:N	1:C:914:ASN:OD1	2.52	0.41
1:A:425:LEU:HD22	1:A:426:PRO:HD2	2.03	0.41
1:A:758:SER:O	1:A:762:GLN:HG3	2.21	0.41
1:C:223:LEU:HD12	1:C:223:LEU:HA	1.94	0.41
1:C:383:SER:HB2	1:C:387:LEU:HB2	2.03	0.41
1:C:398:ASP:HB2	1:C:512:VAL:HG13	2.03	0.41
1:C:436:TRP:HZ3	1:C:511:VAL:HG12	1.85	0.41
1:B:406:GLU:N	1:B:406:GLU:OE1	2.54	0.41
1:C:337:PRO:HB3	1:C:340:GLU:OE2	2.21	0.41
1:B:296:LEU:O	1:B:299:THR:OG1	2.33	0.41
1:A:773:GLU:HB2	1:A:776:LYS:HZ1	1.85	0.41
1:C:866:THR:O	1:C:870:ILE:HG12	2.21	0.41
1:A:295:PRO:O	1:A:299:THR:HG23	2.21	0.41
1:C:457:ARG:NE	1:C:467:ASP:OD1	2.52	0.41
1:A:334:ASN:ND2	1:A:360:ASN:O	2.43	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:C:326:ILE:HD13	1:C:533:LEU:HD12	2.03	0.41
1:C:393:THR:OG1	1:C:516:GLU:O	2.38	0.41
1:A:951:VAL:HG13	1:A:954:GLN:HE21	1.86	0.41
1:B:102:ARG:HH21	1:B:246:ILE:HG12	1.86	0.41
1:B:406:GLU:N	1:B:406:GLU:OE1	2.54	0.41
1:B:562:PHE:HD1	1:C:41:LYS:HG2	1.85	0.41
1:B:738:CYS:HB2	1:B:760:CYS:HB2	1.39	0.41
1:B:767:LEU:HD21	1:B:1008:VAL:HG12	2.02	0.41
1:B:962:LEU:HD22	1:B:1007:TYR:CZ	2.56	0.41
1:C:331:ASN:OD1	1:C:331:ASN:N	2.44	0.41
1:C:574:ASP:OD1	1:C:574:ASP:N	2.51	0.41
1:C:773:GLU:HB2	1:C:1019:ARG:HH12	1.85	0.41
1:C:818:ILE:O	1:C:822:LEU:HG	2.20	0.41
1:C:91:TYR:HE2	1:C:93:ALA:HB2	1.86	0.41
1:A:202:LYS:HG3	1:A:228:ASP:OD1	2.21	0.41
1:A:429:PHE:HE2	1:A:514:SER:HA	1.86	0.41
1:A:557:LYS:HE2	1:A:559:PHE:HE2	1.86	0.41
1:B:503:VAL:HA	1:B:506:GLN:HG3	2.03	0.41
1:B:767:LEU:HD21	1:B:1008:VAL:HG12	2.03	0.41
1:C:985:ASP:OD1	1:C:985:ASP:N	2.53	0.41
1:B:386:LYS:HA	1:B:389:ASP:HB3	2.02	0.41
1:B:884:SER:OG	1:B:894:LEU:O	2.38	0.41
1:B:978:ASN:HA	1:B:981:LEU:HG	2.02	0.41
1:A:722:VAL:HG22	1:A:930:ALA:HB1	2.02	0.41
1:B:598:ILE:HD11	1:B:666:ILE:HD13	2.02	0.41
1:A:22:THR:OG1	1:A:78:ARG:NH1	2.53	0.41
1:A:377:PHE:H	1:C:408:ARG:NH2	2.18	0.41
1:A:398:ASP:HB2	1:A:512:VAL:HG13	2.03	0.41
1:B:731:MET:HG3	1:B:1018:ILE:HG13	2.03	0.41
1:B:989:ALA:O	1:B:993:ILE:HG12	2.21	0.41
1:C:884:SER:OG	1:C:887:THR:OG1	2.32	0.41
1:A:89:GLY:HA3	1:A:270:LEU:HD12	2.03	0.41
1:A:328:ARG:NH1	1:A:531:THR:O	2.54	0.41
1:C:310:LYS:HG2	1:C:664:ILE:HG21	2.03	0.41
1:C:316:SER:OG	1:C:317:ASN:N	2.54	0.41
1:C:806:LEU:HD12	1:C:807:PRO:HD2	2.02	0.41
1:A:987:PRO:HD2	1:A:988:GLU:OE1	2.21	0.41
1:B:290:ASP:HB3	1:B:293:LEU:HB2	2.01	0.41
1:B:407:VAL:O	1:B:410:ILE:HG22	2.21	0.41
1:C:536:ASN:HA	1:C:553:THR:HG22	2.01	0.41
1:C:912:THR:O	1:C:915:VAL:HG12	2.21	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:31:SER:OG	1:A:60:SER:O	2.31	0.41
1:C:462:LYS:H	1:C:462:LYS:HE2	1.86	0.41
1:A:403:ARG:HD2	1:A:505:TYR:CD1	2.55	0.41
1:A:1107:ARG:HE	1:A:1107:ARG:HB3	1.66	0.41
1:A:287:ASP:OD1	1:A:288:ALA:N	2.54	0.41
1:A:326:ILE:HD11	1:A:534:VAL:HG22	2.03	0.41
1:A:862:PRO:HA	1:A:863:PRO:HD3	1.95	0.41
1:B:27:ALA:HB3	1:B:64:TRP:HE3	1.86	0.41
1:B:1014:ARG:O	1:B:1018:ILE:HG12	2.21	0.41
1:A:599:THR:HB	1:A:608:VAL:HG23	2.02	0.41
1:A:737:ASP:OD1	1:A:740:MET:HB3	2.21	0.41
1:B:517:LEU:HD12	1:B:518:LEU:HB3	2.02	0.41
1:B:822:LEU:HD11	1:B:1061:VAL:HG21	2.03	0.41
1:A:22:THR:O	1:A:78:ARG:NH1	2.53	0.41
1:A:402:ILE:HD12	1:A:403:ARG:H	1.85	0.41
1:B:84:LEU:H	1:B:237:ARG:CZ	2.34	0.41
1:B:537:LYS:HE3	1:B:537:LYS:HB3	1.90	0.41
1:C:586:ASP:OD1	1:C:586:ASP:N	2.53	0.41
1:C:670:ASN:HD22	1:C:671:SER:N	2.18	0.41
1:B:326:ILE:HD13	1:B:533:LEU:HD12	2.02	0.41
1:B:379:CYS:HA	1:B:432:CYS:HA	2.02	0.41
1:A:406:GLU:OE1	1:A:406:GLU:N	2.53	0.41
1:A:407:VAL:O	1:A:410:ILE:HG22	2.20	0.41
1:A:1006:THR:OG1	1:A:1007:TYR:N	2.52	0.41
1:B:326:ILE:HD13	1:B:533:LEU:HD12	2.03	0.41
1:A:598:ILE:HB	1:A:609:ALA:HB3	2.02	0.41
1:B:708:SER:HB3	1:B:711:SER:HB2	2.03	0.41
1:C:452:LEU:HD23	1:C:452:LEU:HA	1.94	0.41
1:C:791:THR:HG21	1:C:806:LEU:HD21	2.02	0.41
1:B:153:MET:HE3	1:B:153:MET:HB3	1.83	0.41
1:C:435:ALA:HB2	1:C:510:VAL:HG12	2.03	0.41
1:C:818:ILE:O	1:C:822:LEU:HG	2.20	0.41
1:C:878:LEU:HD21	1:C:1052:PHE:HB3	2.02	0.41
1:A:560:LEU:HD21	1:B:224:GLU:OE2	2.20	0.41
1:A:993:ILE:O	1:A:997:ILE:HG12	2.21	0.41
1:A:1050:MET:HE2	1:A:1050:MET:HB3	1.83	0.41
1:B:993:ILE:O	1:B:997:ILE:HG12	2.21	0.41
1:C:976:VAL:HG13	1:C:979:ASP:HB3	2.03	0.41
1:B:1039:ARG:HD3	1:C:1031:GLU:OE2	2.20	0.41
1:C:452:LEU:HG	1:C:494:SER:HA	2.03	0.41
1:A:365:TYR:HD1	1:A:387:LEU:HB3	1.84	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:B:720:ILE:HD11	1:B:1065:VAL:HG12	2.02	0.41
1:B:818:ILE:O	1:B:822:LEU:HD23	2.20	0.41
1:C:84:LEU:HD22	1:C:238:PHE:CZ	2.56	0.41
1:C:1028:LYS:NZ	1:C:1042:PHE:O	2.48	0.41
1:A:762:GLN:HA	1:A:765:ARG:HG2	2.02	0.41
1:B:296:LEU:O	1:B:299:THR:OG1	2.36	0.41
1:B:390:LEU:HD21	1:B:517:LEU:HD11	2.03	0.41
1:B:827:THR:O	1:B:827:THR:OG1	2.27	0.41
1:C:1115:ILE:HG22	1:C:1137:VAL:HG23	2.03	0.41
1:A:866:THR:O	1:A:870:ILE:HG22	2.21	0.41
1:B:676:THR:HA	1:B:690:GLN:HG2	2.03	0.41
1:A:1107:ARG:HE	1:A:1107:ARG:HB3	1.68	0.41
1:B:201:PHE:CE2	1:B:235:ILE:HD12	2.55	0.41
1:C:775:ASP:OD1	1:C:775:ASP:N	2.54	0.41
1:A:1081:ILE:HG12	1:A:1095:PHE:CE2	2.56	0.41
1:A:299:THR:HG21	1:A:597:VAL:HG21	2.02	0.41
1:A:765:ARG:CZ	1:C:957:GLN:HE21	2.34	0.41
1:B:498:GLN:HB2	1:B:501:TYR:CE2	2.56	0.41
1:A:984:LEU:HD23	1:A:988:GLU:HG2	2.02	0.41
1:B:189:LEU:HD23	1:B:217:PRO:HG2	2.01	0.41
1:B:276:LEU:HD11	1:B:304:LYS:HA	2.03	0.41
1:C:877:LEU:HD21	1:C:1029:MET:SD	2.60	0.41
1:B:722:VAL:HG12	1:B:1065:VAL:HG22	2.02	0.41
1:C:722:VAL:HG22	1:C:930:ALA:HB1	2.03	0.41
1:C:882:ILE:HD12	1:C:882:ILE:HA	1.87	0.41
1:A:295:PRO:O	1:A:299:THR:HG23	2.21	0.41
1:B:293:LEU:HD22	1:B:294:ASP:OD2	2.20	0.41
1:B:314:GLN:HE22	1:B:594:GLY:HA3	1.86	0.41
1:C:402:ILE:HD11	1:C:418:ILE:HG13	2.02	0.41
1:C:774:GLN:HE22	1:C:1018:ILE:HG21	1.85	0.41
1:A:233:ILE:HD12	1:A:233:ILE:HA	1.94	0.41
1:A:276:LEU:HD13	1:A:276:LEU:HA	1.86	0.41
1:B:533:LEU:HD21	1:B:585:LEU:HD11	2.03	0.41
1:B:763:LEU:HB2	1:B:1008:VAL:HG11	2.03	0.41
1:A:720:ILE:HD11	1:A:1065:VAL:HG12	2.02	0.41
1:B:177:MET:HG2	1:B:190:ARG:HH22	1.85	0.41
1:B:598:ILE:HB	1:B:609:ALA:HB3	2.02	0.41
1:B:722:VAL:HG12	1:B:1065:VAL:HG22	2.02	0.41
1:A:759:PHE:O	1:A:762:GLN:NE2	2.54	0.41
1:A:989:ALA:O	1:A:993:ILE:HG12	2.21	0.41
1:B:406:GLU:N	1:B:406:GLU:OE1	2.54	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:300:LYS:NZ	1:A:306:PHE:O	2.54	0.41
1:B:279:TYR:HE1	1:B:285:ILE:HG12	1.86	0.41
1:C:332:ILE:HG13	1:C:362:VAL:HG22	2.02	0.41
1:C:1031:GLU:HG3	1:C:1039:ARG:NH2	2.35	0.41
1:A:34:ARG:HG3	1:A:216:LEU:HD11	2.02	0.41
1:B:513:LEU:HD12	1:B:513:LEU:HA	1.93	0.41
1:C:722:VAL:HG12	1:C:1065:VAL:HG22	2.02	0.41
1:A:302:THR:HG21	1:A:315:THR:HA	2.03	0.41
1:A:541:PHE:N	1:A:548:GLY:O	2.52	0.41
1:B:277:LEU:HD12	1:B:285:ILE:HD13	2.02	0.41
1:B:471:GLU:OE1	1:C:113:LYS:NZ	2.41	0.41
1:C:192:PHE:HB3	1:C:194:PHE:HE2	1.86	0.41
1:C:592:PHE:HD1	1:C:593:GLY:HA2	1.86	0.41
1:C:906:PHE:HB3	1:C:911:VAL:HB	2.03	0.41
1:C:1108:ASN:OD1	1:C:1108:ASN:N	2.54	0.41
1:B:89:GLY:HA3	1:B:270:LEU:HD12	2.03	0.41
1:B:374:PHE:HB2	1:B:377:PHE:CE1	2.55	0.41
1:B:552:LEU:HD12	1:B:552:LEU:HA	1.94	0.41
1:C:973:ILE:HG21	1:C:983:ARG:HH22	1.86	0.41
1:A:334:ASN:ND2	1:A:360:ASN:O	2.44	0.41
1:A:867:ASP:N	1:A:867:ASP:OD1	2.54	0.41
1:A:1105:THR:HG21	1:A:1110:TYR:HA	2.03	0.41
1:B:332:ILE:HG13	1:B:362:VAL:HG22	2.02	0.41
1:B:406:GLU:N	1:B:406:GLU:OE1	2.54	0.41
1:B:1045:LYS:O	1:B:1066:THR:HG21	2.21	0.41
1:B:1107:ARG:HG3	1:B:1108:ASN:OD1	2.21	0.41
1:C:1002:GLN:O	1:C:1006:THR:HG23	2.21	0.41
1:C:1010:GLN:NE2	1:C:1014:ARG:HH12	2.19	0.41
1:A:997:ILE:O	1:A:1001:LEU:HB3	2.21	0.41
1:B:431:GLY:HA2	1:B:515:PHE:CZ	2.56	0.41
1:B:442:ASP:OD2	1:B:509:ARG:NE	2.48	0.41
1:B:758:SER:O	1:B:762:GLN:HG3	2.21	0.41
1:B:759:PHE:HA	1:B:762:GLN:HE21	1.86	0.41
1:C:295:PRO:O	1:C:299:THR:HG23	2.21	0.41
1:A:406:GLU:OE1	1:A:406:GLU:N	2.54	0.41
1:A:962:LEU:HD22	1:A:1007:TYR:CE2	2.56	0.41
1:C:1073:LYS:HE3	1:C:1075:PHE:CZ	2.55	0.41
1:B:406:GLU:N	1:B:406:GLU:OE1	2.54	0.41
1:B:462:LYS:H	1:B:462:LYS:HE2	1.86	0.41
1:B:740:MET:HE3	1:B:740:MET:HB3	1.97	0.41
1:B:1106:GLN:H	1:B:1106:GLN:HG2	1.56	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:C:759:PHE:HA	1:C:762:GLN:HE21	1.86	0.41
1:A:43:PHE:HB3	1:C:565:PHE:O	2.20	0.41
1:A:47:VAL:HG12	1:C:569:ILE:HA	2.03	0.41
1:B:48:LEU:HD12	1:B:48:LEU:HA	1.94	0.41
1:B:537:LYS:HE3	1:B:537:LYS:HB3	1.87	0.41
1:B:733:LYS:HB3	1:B:861:LEU:HD23	2.01	0.41
1:B:775:ASP:HB3	1:B:864:LEU:HD13	2.03	0.41
1:B:429:PHE:CE2	1:B:514:SER:HA	2.55	0.41
1:B:785:VAL:HG12	1:B:787:GLN:H	1.86	0.41
1:B:905:ARG:NE	1:B:1049:LEU:O	2.52	0.41
1:C:406:GLU:HG2	1:C:418:ILE:HD11	2.03	0.41
1:C:429:PHE:HE1	1:C:514:SER:HA	1.86	0.41
1:C:557:LYS:HE2	1:C:559:PHE:HE1	1.86	0.41
1:C:867:ASP:N	1:C:867:ASP:OD1	2.52	0.41
1:A:769:GLY:HA2	1:A:772:VAL:HG22	2.01	0.41
1:A:994:ASP:O	1:A:998:THR:HG23	2.21	0.41
1:B:484:LYS:HD2	1:B:490:PHE:HB2	2.02	0.41
1:B:905:ARG:NH1	1:B:1049:LEU:O	2.54	0.41
1:B:1106:GLN:OE1	1:B:1106:GLN:N	2.54	0.41
1:C:151:SER:HB2	1:C:153:MET:HE3	2.02	0.41
1:C:1135:ASN:OD1	1:C:1136:THR:N	2.44	0.41
1:A:781:VAL:HG13	1:A:782:PHE:CD1	2.56	0.41
1:A:1079:PRO:HB3	1:B:900:MET:HE1	2.03	0.41
1:B:222:ALA:HB2	1:B:285:ILE:HB	2.02	0.41
1:B:299:THR:HG22	1:B:597:VAL:HG21	2.01	0.41
1:B:425:LEU:HD12	1:B:429:PHE:CD2	2.56	0.41
1:B:1046:GLY:HA2	1:C:890:ALA:HA	2.03	0.41
1:A:712:ILE:HG23	1:A:1075:PHE:HB2	2.02	0.41
1:B:324:GLU:OE1	1:B:324:GLU:N	2.54	0.41
1:B:716:THR:HG21	1:B:1073:LYS:HG3	2.03	0.41
1:C:407:VAL:HG21	1:C:508:TYR:CD2	2.52	0.41
1:C:976:VAL:HG13	1:C:979:ASP:HB3	2.02	0.41
1:B:517:LEU:HD12	1:B:518:LEU:HB2	2.02	0.41
1:B:699:LEU:HG	1:C:873:TYR:HE1	1.86	0.41
1:B:909:ILE:HD11	1:B:1047:TYR:CD1	2.56	0.41
1:C:424:LYS:HD3	1:C:424:LYS:HA	1.86	0.41
1:C:462:LYS:O	1:C:465:GLU:HG3	2.21	0.41
1:C:712:ILE:HD12	1:C:712:ILE:HA	1.82	0.41
1:C:722:VAL:HG22	1:C:930:ALA:HB1	2.01	0.41
1:C:914:ASN:HD21	1:C:1106:GLN:CD	2.28	0.41
1:A:392:PHE:HB2	1:A:524:VAL:HG22	2.03	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:431:GLY:HA2	1:A:515:PHE:CE1	2.55	0.41
1:A:559:PHE:HB2	1:A:584:ILE:HD13	2.02	0.41
1:A:714:ILE:HD12	1:A:1107:ARG:HA	2.01	0.41
1:B:406:GLU:N	1:B:406:GLU:OE1	2.54	0.41
1:B:604:THR:OG1	1:B:605:SER:N	2.53	0.41
1:B:701:VAL:HG13	1:C:787:GLN:HG2	2.03	0.41
1:C:722:VAL:HG22	1:C:930:ALA:HB1	2.03	0.41
1:C:914:ASN:O	1:C:918:GLU:HB2	2.21	0.41
1:A:89:GLY:HA3	1:A:270:LEU:HD12	2.03	0.41
1:A:521:PRO:HA	1:A:564:GLN:HE22	1.86	0.41
1:A:734:THR:O	1:A:767:LEU:HD12	2.21	0.41
1:A:1081:ILE:HG23	1:A:1088:HIS:HB2	2.03	0.41
1:B:576:VAL:HG12	1:B:587:ILE:HD11	2.03	0.41
1:B:759:PHE:HA	1:B:762:GLN:HE21	1.86	0.41
1:C:14:GLN:O	1:C:158:ARG:NE	2.54	0.41
1:A:106:PHE:HD1	1:A:238:PHE:HB3	1.86	0.41
1:A:557:LYS:HE2	1:A:559:PHE:HE2	1.86	0.41
1:C:976:VAL:HG13	1:C:979:ASP:HB3	2.03	0.41
1:A:128:ILE:HB	1:A:170:TYR:HD2	1.84	0.41
1:A:730:SER:OG	1:A:731:MET:N	2.52	0.41
1:A:1081:ILE:HD12	1:A:1081:ILE:HA	1.80	0.41
1:B:379:CYS:HA	1:B:432:CYS:HA	2.02	0.41
1:B:670:ASN:HD22	1:B:671:SER:N	2.17	0.41
1:C:912:THR:O	1:C:915:VAL:HG12	2.21	0.41
1:A:222:ALA:HB2	1:A:285:ILE:HB	2.03	0.41
1:A:406:GLU:OE1	1:A:406:GLU:N	2.54	0.41
1:B:293:LEU:HD22	1:B:294:ASP:OD2	2.20	0.41
1:B:365:TYR:O	1:B:369:TYR:HB2	2.21	0.41
1:C:387:LEU:HD13	1:C:392:PHE:HZ	1.86	0.41
1:C:501:TYR:HB3	1:C:505:TYR:HB3	2.03	0.41
1:A:1002:GLN:O	1:A:1006:THR:HG23	2.21	0.41
1:C:247:SER:O	1:C:258:TRP:NE1	2.52	0.41
1:C:330:PRO:HD3	1:C:579:PRO:HB2	2.03	0.41
1:C:912:THR:O	1:C:915:VAL:HG12	2.21	0.41
1:A:86:PHE:CE1	1:A:90:VAL:HG12	2.56	0.41
1:A:594:GLY:H	1:A:613:GLN:HE21	1.68	0.41
1:A:759:PHE:HA	1:A:762:GLN:HE21	1.86	0.41
1:B:54:LEU:HD12	1:B:270:LEU:HD12	2.03	0.41
1:B:293:LEU:HD23	1:B:294:ASP:HB3	2.03	0.41
1:B:406:GLU:N	1:B:406:GLU:OE1	2.54	0.41
1:C:410:ILE:HG21	1:C:433:VAL:HG21	2.03	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:14:GLN:O	1:A:158:ARG:NE	2.52	0.41
1:B:431:GLY:HA2	1:B:515:PHE:CE2	2.55	0.41
1:B:596:SER:HB2	1:B:611:LEU:HG	2.02	0.41
1:C:330:PRO:HD3	1:C:579:PRO:HB2	2.03	0.41
1:C:380:TYR:HE1	1:C:433:VAL:HG12	1.85	0.41
1:A:206:LYS:HE3	1:A:208:THR:HG23	2.02	0.41
1:B:280:ASN:OD1	1:B:284:THR:N	2.53	0.41
1:B:955:ASN:O	1:B:959:LEU:HD22	2.22	0.41
1:C:457:ARG:HH22	1:C:461:LEU:HG	1.86	0.41
1:A:1081:ILE:HD11	1:A:1135:ASN:HB3	2.01	0.41
1:B:431:GLY:HA2	1:B:515:PHE:CE2	2.56	0.41
1:B:774:GLN:HE22	1:B:1018:ILE:HG21	1.86	0.41
1:B:866:THR:O	1:B:870:ILE:HG12	2.22	0.41
1:A:734:THR:O	1:A:767:LEU:HD12	2.21	0.41
1:B:327:VAL:HB	1:B:531:THR:HG23	2.03	0.41
1:A:57:PRO:HB3	1:A:273:ARG:HH12	1.86	0.41
1:A:759:PHE:HA	1:A:762:GLN:HE21	1.86	0.41
1:A:890:ALA:HA	1:C:1046:GLY:HA2	2.03	0.41
1:B:330:PRO:HD3	1:B:579:PRO:HB2	2.02	0.41
1:C:788:ILE:HG23	1:C:876:ALA:HB2	2.03	0.41
1:A:866:THR:OG1	1:A:867:ASP:N	2.54	0.41
1:A:883:THR:HB	1:C:705:VAL:HG11	2.02	0.41
1:B:351:TYR:HE1	1:B:452:LEU:HB2	1.85	0.41
1:B:1090:PRO:HD3	1:B:1095:PHE:HE2	1.86	0.41
1:C:431:GLY:HA2	1:C:515:PHE:CE1	2.56	0.41
1:C:878:LEU:HD21	1:C:1052:PHE:HB3	2.03	0.41
1:A:299:THR:HG21	1:A:597:VAL:HG21	2.03	0.41
1:C:369:TYR:CZ	1:C:384:PRO:HB2	2.56	0.41
1:A:740:MET:HE1	1:A:857:GLY:HA3	2.02	0.41
1:B:183:GLN:HG3	1:B:187:LYS:HG3	2.03	0.41
1:A:861:LEU:HD12	1:A:862:PRO:HD2	2.03	0.41
1:B:669:GLY:O	1:B:670:ILE:HD13	2.21	0.41
1:C:370:ASN:ND2	1:C:384:PRO:HB2	2.36	0.41
1:A:733:LYS:HB3	1:A:861:LEU:HD23	2.02	0.41
1:A:809:PRO:HA	1:A:814:LYS:HD2	2.03	0.41
1:B:884:SER:OG	1:B:894:LEU:O	2.39	0.41
1:C:726:ILE:HG23	1:C:947:LYS:HB3	2.03	0.41
1:C:869:MET:HE2	1:C:869:MET:HB3	1.93	0.41
1:A:726:ILE:HD12	1:A:1061:VAL:HG12	2.02	0.41
1:C:809:PRO:O	1:C:814:LYS:NZ	2.39	0.41
1:A:1090:PRO:HD3	1:A:1095:PHE:HE2	1.85	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:B:537:LYS:HE3	1:B:537:LYS:HB3	1.89	0.41
1:B:592:PHE:CE2	1:C:740:MET:HG3	2.56	0.41
1:B:730:SER:OG	1:B:731:MET:N	2.52	0.41
1:A:701:VAL:HG13	1:B:787:GLN:HG2	2.03	0.41
1:A:758:SER:O	1:A:762:GLN:HG3	2.21	0.41
1:A:999:GLY:HA2	1:A:1002:GLN:HG3	2.02	0.41
1:A:183:GLN:HG3	1:A:187:LYS:HG3	2.02	0.41
1:B:39:PRO:HG3	1:B:51:THR:HG21	2.03	0.41
1:A:365:TYR:CE2	1:A:387:LEU:HD12	2.55	0.41
1:C:206:LYS:HD2	1:C:207:HIS:N	2.36	0.41
1:C:365:TYR:CE1	1:C:387:LEU:HD12	2.56	0.41
1:C:902:MET:HB3	1:C:916:LEU:HD11	2.02	0.41
1:A:984:LEU:HG	1:A:989:ALA:HB2	2.02	0.41
1:B:973:ILE:HG21	1:B:983:ARG:HH12	1.86	0.41
1:C:902:MET:HB3	1:C:916:LEU:HD11	2.03	0.41
1:C:980:ILE:HD13	1:C:980:ILE:HA	1.94	0.41
1:B:183:GLN:HG3	1:B:187:LYS:HG3	2.02	0.41
1:C:85:PRO:HA	1:C:237:ARG:HG2	2.03	0.41
1:A:1028:LYS:NZ	1:A:1042:PHE:O	2.48	0.41
1:B:742:ILE:HD13	1:B:742:ILE:HA	1.95	0.41
1:B:802:PHE:HE1	1:B:927:PHE:HE2	1.68	0.41
1:A:287:ASP:OD1	1:A:288:ALA:N	2.54	0.41
1:A:402:ILE:HD11	1:A:418:ILE:HG13	2.03	0.41
1:A:1090:PRO:HD3	1:A:1095:PHE:HE2	1.86	0.41
1:C:205:SER:N	1:C:224:GLU:O	2.39	0.41
1:C:406:GLU:OE1	1:C:406:GLU:N	2.54	0.41
1:C:730:SER:OG	1:C:731:MET:N	2.54	0.41
1:C:758:SER:O	1:C:762:GLN:HG3	2.21	0.41
1:A:735:SER:HB2	1:A:861:LEU:HD21	2.02	0.41
1:A:862:PRO:HA	1:A:863:PRO:HD3	1.97	0.41
1:B:722:VAL:HG12	1:B:1065:VAL:HG22	2.02	0.41
1:C:31:SER:O	1:C:31:SER:OG	2.36	0.41
1:C:121:ASN:HA	1:C:126:VAL:HA	2.02	0.41
1:B:738:CYS:O	1:B:742:ILE:HG12	2.21	0.40
1:A:212:LEU:HD22	1:A:217:PRO:HD3	2.03	0.40
1:C:714:ILE:HG12	1:C:1075:PHE:HD2	1.86	0.40
1:A:296:LEU:O	1:A:299:THR:OG1	2.35	0.40
1:A:559:PHE:CD1	1:A:584:ILE:HG21	2.56	0.40
1:B:1005:GLN:O	1:B:1009:THR:HG23	2.21	0.40
1:A:442:ASP:OD2	1:A:509:ARG:NE	2.47	0.40
1:C:867:ASP:N	1:C:867:ASP:OD1	2.52	0.40

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:C:934:ILE:HD12	1:C:934:ILE:HA	1.89	0.40
1:A:115:GLN:HE21	1:C:468:ILE:HG12	1.86	0.40
1:B:733:LYS:HD2	1:B:771:ALA:HB1	2.03	0.40
1:C:966:LEU:HD12	1:C:1000:ARG:NH1	2.36	0.40
1:A:642:VAL:HG22	1:A:651:ILE:HG22	2.03	0.40
1:A:775:ASP:HB3	1:A:864:LEU:HD13	2.03	0.40
1:B:513:LEU:HD12	1:B:513:LEU:HA	1.91	0.40
1:C:802:PHE:HB3	1:C:806:LEU:HD23	2.03	0.40
1:C:983:ARG:HB2	1:C:984:LEU:HD12	2.03	0.40
1:A:295:PRO:O	1:A:299:THR:HG23	2.21	0.40
1:A:896:ILE:HD12	1:C:712:ILE:HD11	2.03	0.40
1:A:347:PHE:HB2	1:A:401:VAL:HG23	2.02	0.40
1:A:740:MET:HE2	1:C:592:PHE:CE2	2.56	0.40
1:B:993:ILE:O	1:B:997:ILE:HG12	2.21	0.40
1:B:1090:PRO:HD3	1:B:1095:PHE:HE2	1.87	0.40
1:C:195:LYS:HD2	1:C:196:ASN:H	1.85	0.40
1:C:697:MET:HB3	1:C:697:MET:HE3	1.84	0.40
1:C:886:TRP:HH2	1:C:904:TYR:HB3	1.86	0.40
1:C:947:LYS:HE2	1:C:947:LYS:HB2	1.94	0.40
1:A:452:LEU:HD23	1:A:492:LEU:HB3	2.03	0.40
1:B:117:LEU:HD11	1:B:128:ILE:HG23	2.03	0.40
1:B:738:CYS:O	1:B:742:ILE:HG12	2.21	0.40
1:B:742:ILE:O	1:B:1000:ARG:NH1	2.55	0.40
1:C:818:ILE:O	1:C:822:LEU:HG	2.21	0.40
1:C:889:GLY:HA3	1:C:1034:LEU:HD11	2.03	0.40
1:C:1004:LEU:O	1:C:1008:VAL:HG13	2.21	0.40
1:A:369:TYR:CE1	1:A:384:PRO:HB2	2.56	0.40
1:B:347:PHE:CE2	1:B:509:ARG:HD3	2.56	0.40
1:B:712:ILE:HG12	1:B:1094:VAL:HG11	2.03	0.40
1:C:409:GLN:HA	1:C:414:GLN:HG2	2.02	0.40
1:C:599:THR:HB	1:C:608:VAL:HG23	2.04	0.40
1:A:917:TYR:HB3	1:C:1129:VAL:HG22	2.02	0.40
1:B:336:CYS:HB3	1:B:358:ILE:HD12	2.02	0.40
1:C:84:LEU:HD12	1:C:267:VAL:HG21	2.03	0.40
1:A:962:LEU:HD22	1:A:1007:TYR:CE2	2.56	0.40
1:B:303:LEU:HD23	1:B:308:VAL:HG22	2.03	0.40
1:B:733:LYS:HB3	1:B:861:LEU:HD23	2.03	0.40
1:C:1045:LYS:HD3	1:C:1045:LYS:HA	1.86	0.40
1:B:954:GLN:OE1	1:B:1014:ARG:NE	2.54	0.40
1:B:1106:GLN:HE22	1:B:1111:GLU:CD	2.27	0.40
1:C:223:LEU:HA	1:C:223:LEU:HD12	1.90	0.40

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:993:ILE:O	1:A:997:ILE:HG12	2.21	0.40
1:A:1080:ALA:HB3	1:A:1129:VAL:HG11	2.02	0.40
1:C:993:ILE:O	1:C:997:ILE:HG12	2.21	0.40
1:A:117:LEU:HD22	1:A:235:ILE:HD11	2.02	0.40
1:A:1128:VAL:HG21	1:B:918:GLU:HG3	2.03	0.40
1:B:406:GLU:OE1	1:B:406:GLU:N	2.54	0.40
1:C:278:LYS:NZ	1:C:287:ASP:OD2	2.52	0.40
1:A:126:VAL:HB	1:A:172:SER:HB3	2.03	0.40
1:A:594:GLY:H	1:A:613:GLN:HE21	1.69	0.40
1:A:1033:VAL:HG11	1:A:1053:PRO:HG3	2.03	0.40
1:B:1052:PHE:HA	1:B:1053:PRO:HD3	1.95	0.40
1:B:330:PRO:HD3	1:B:579:PRO:HB2	2.03	0.40
1:B:713:ALA:HB2	1:C:895:GLN:NE2	2.36	0.40
1:C:87:ASN:OD1	1:C:87:ASN:N	2.49	0.40
1:C:398:ASP:OD2	1:C:423:TYR:OH	2.29	0.40
1:C:412:PRO:HG3	1:C:429:PHE:HD2	1.86	0.40
1:B:426:PRO:HD3	1:B:463:PRO:HB3	2.02	0.40
1:C:34:ARG:NH2	1:C:221:SER:H	2.19	0.40
1:C:199:GLY:HA2	1:C:232:GLY:HA2	2.03	0.40
1:C:722:VAL:HG22	1:C:930:ALA:HB1	2.03	0.40
1:C:106:PHE:HB2	1:C:117:LEU:HB3	2.01	0.40
1:A:645:THR:OG1	1:A:648:GLY:O	2.33	0.40
1:A:734:THR:O	1:A:767:LEU:HD12	2.21	0.40
1:A:1115:ILE:HG22	1:A:1137:VAL:HG13	2.03	0.40
1:B:314:GLN:HA	1:B:314:GLN:NE2	2.36	0.40
1:B:866:THR:HG22	1:B:869:MET:HE3	2.03	0.40
1:C:1083:HIS:HE1	1:C:1136:THR:HG23	1.87	0.40
1:A:328:ARG:HD2	1:A:533:LEU:HB2	2.01	0.40
1:A:813:SER:OG	1:A:868:GLU:OE2	2.26	0.40
1:B:54:LEU:HB3	1:B:270:LEU:HB3	2.02	0.40
1:B:436:TRP:CE3	1:B:509:ARG:HD2	2.57	0.40
1:C:994:ASP:O	1:C:998:THR:HG23	2.21	0.40
1:A:43:PHE:CG	1:C:563:GLN:HG2	2.56	0.40
1:A:233:ILE:HG13	1:A:234:ASN:N	2.36	0.40
1:A:730:SER:OG	1:A:731:MET:N	2.53	0.40
1:A:993:ILE:O	1:A:997:ILE:HG12	2.22	0.40
1:B:720:ILE:HD11	1:B:1065:VAL:HG12	2.02	0.40
1:B:867:ASP:OD1	1:B:867:ASP:N	2.55	0.40
1:C:884:SER:OG	1:C:894:LEU:O	2.39	0.40
1:B:870:ILE:O	1:B:874:THR:HG23	2.21	0.40
1:A:549:THR:HG22	1:B:745:ASP:HB3	2.03	0.40

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:C:730:SER:OG	1:C:731:MET:N	2.55	0.40
1:A:353:TRP:NE1	1:A:466:ARG:HB2	2.31	0.40
1:A:916:LEU:HD22	1:A:917:TYR:CD1	2.56	0.40
1:B:52:GLN:HE22	1:B:273:ARG:N	2.19	0.40
1:C:358:ILE:HB	1:C:395:VAL:HG23	2.01	0.40
1:C:1002:GLN:O	1:C:1006:THR:HG23	2.22	0.40
1:A:1090:PRO:HD3	1:A:1095:PHE:HE2	1.86	0.40
1:B:800:PHE:HD1	1:B:927:PHE:HD2	1.69	0.40
1:B:884:SER:OG	1:B:894:LEU:O	2.39	0.40
1:C:909:ILE:HA	1:C:1038:LYS:HZ1	1.86	0.40
1:A:572:THR:HG22	1:B:976:VAL:HG23	2.03	0.40
1:A:613:GLN:C	1:A:613:GLN:NE2	2.79	0.40
1:B:273:ARG:HD3	1:B:273:ARG:HA	1.88	0.40
1:B:273:ARG:HD2	1:B:292:ALA:HB3	2.04	0.40
1:B:909:ILE:HD11	1:B:1047:TYR:CD1	2.57	0.40
1:C:814:LYS:HA	1:C:814:LYS:HD3	1.86	0.40
1:B:49:HIS:HE1	1:B:51:THR:HB	1.86	0.40
1:B:280:ASN:OD1	1:B:284:THR:N	2.53	0.40
1:C:537:LYS:HE3	1:C:537:LYS:HB2	1.94	0.40
1:A:497:PHE:CE1	1:A:507:PRO:HB3	2.56	0.40
1:B:326:ILE:HD13	1:B:533:LEU:HD12	2.04	0.40
1:B:713:ALA:HB2	1:C:895:GLN:CD	2.46	0.40
1:C:730:SER:OG	1:C:731:MET:N	2.55	0.40
1:C:498:GLN:HE21	1:C:501:TYR:HD2	1.68	0.40
1:C:742:ILE:HD12	1:C:742:ILE:HA	1.81	0.40
1:B:497:PHE:CE2	1:B:507:PRO:HB3	2.56	0.40
1:C:970:PHE:HD2	1:C:996:LEU:HD13	1.86	0.40
1:A:369:TYR:OH	1:C:415:THR:OG1	2.23	0.40
1:B:431:GLY:HA2	1:B:515:PHE:CE2	2.56	0.40
1:B:452:LEU:HA	1:B:494:SER:HA	2.03	0.40
1:A:1105:THR:HG21	1:A:1110:TYR:HA	2.03	0.40
1:B:212:LEU:HD13	1:B:217:PRO:HG3	2.04	0.40
1:C:1083:HIS:HE1	1:C:1136:THR:HG23	1.86	0.40
1:A:905:ARG:NE	1:A:1049:LEU:O	2.53	0.40
1:B:954:GLN:OE1	1:B:1014:ARG:NH1	2.54	0.40
1:C:738:CYS:O	1:C:742:ILE:HB	2.21	0.40
1:A:983:ARG:NH2	1:C:517:LEU:HD11	2.36	0.40
1:B:559:PHE:HB3	1:B:563:GLN:HB2	2.02	0.40
1:B:598:ILE:HD11	1:B:611:LEU:HD23	2.03	0.40
1:B:751:ASN:HA	1:B:754:LEU:HG	2.02	0.40
1:A:1046:GLY:HA2	1:B:890:ALA:HA	2.02	0.40

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:C:290:ASP:HB3	1:C:293:LEU:HB2	2.03	0.40
1:A:819:GLU:HA	1:A:822:LEU:HG	2.03	0.40
1:B:350:VAL:HG11	1:B:418:ILE:HD11	2.03	0.40
1:B:569:ILE:H	1:B:569:ILE:HG13	1.59	0.40
1:B:323:THR:HB	1:B:537:LYS:HZ3	1.86	0.40
1:A:642:VAL:HG13	1:A:651:ILE:HG22	2.03	0.40
1:A:712:ILE:HG23	1:A:1075:PHE:HB2	2.02	0.40
1:A:895:GLN:OE1	1:C:711:SER:OG	2.28	0.40
1:B:642:VAL:HG22	1:B:651:ILE:HG22	2.04	0.40
1:B:21:ARG:HG3	1:B:79:PHE:HD2	1.86	0.40
1:C:14:GLN:HA	1:C:138:ASP:HB2	2.04	0.40
1:C:207:HIS:O	1:C:207:HIS:ND1	2.54	0.40
1:C:370:ASN:HD22	1:C:384:PRO:HD2	1.86	0.40
1:C:452:LEU:HB3	1:C:492:LEU:HD11	2.03	0.40
1:A:390:LEU:HD12	1:A:391:CYS:H	1.85	0.40
1:A:884:SER:OG	1:A:887:THR:OG1	2.30	0.40
1:C:887:THR:HG21	1:C:894:LEU:HB2	2.02	0.40
1:A:189:LEU:HD12	1:A:217:PRO:HG2	2.03	0.40
1:B:295:PRO:O	1:B:299:THR:HG23	2.21	0.40
1:B:406:GLU:N	1:B:406:GLU:OE1	2.55	0.40
1:B:763:LEU:O	1:B:767:LEU:HD22	2.21	0.40
1:C:431:GLY:HA2	1:C:515:PHE:CE2	2.55	0.40
1:C:858:LEU:HD13	1:C:959:LEU:HD12	2.02	0.40
1:C:906:PHE:CD1	1:C:916:LEU:HB2	2.56	0.40
1:B:884:SER:OG	1:B:887:THR:OG1	2.25	0.40
1:C:43:PHE:CE1	1:C:283:GLY:HA3	2.57	0.40
1:C:770:ILE:HA	1:C:773:GLU:HG3	2.03	0.40
1:A:1028:LYS:HE2	1:A:1028:LYS:HB2	1.81	0.40
1:C:52:GLN:HE22	1:C:273:ARG:N	2.19	0.40
1:C:303:LEU:HD21	1:C:313:TYR:CE2	2.57	0.40
1:A:425:LEU:HD13	1:A:426:PRO:HD2	2.03	0.40
1:B:431:GLY:HA2	1:B:515:PHE:CZ	2.56	0.40
1:B:513:LEU:HD12	1:B:513:LEU:HA	1.89	0.40
1:C:407:VAL:O	1:C:410:ILE:HG22	2.21	0.40
1:C:720:ILE:HD11	1:C:1065:VAL:HG12	2.03	0.40
1:B:99:ASN:HB3	1:B:102:ARG:CZ	2.51	0.40
1:B:802:PHE:HB3	1:B:806:LEU:HD23	2.04	0.40
1:B:962:LEU:HD12	1:B:962:LEU:HA	1.86	0.40
1:C:332:ILE:HG13	1:C:362:VAL:HG22	2.03	0.40
1:A:597:VAL:HG13	1:A:608:VAL:HG13	2.03	0.40
1:B:37:TYR:OH	1:B:54:LEU:O	2.25	0.40

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:C:741:TYR:CE1	1:C:962:LEU:HD12	2.56	0.40
1:C:960:ASN:OD1	1:C:964:LYS:NZ	2.51	0.40
1:A:730:SER:OG	1:A:731:MET:N	2.52	0.40
1:B:406:GLU:N	1:B:406:GLU:OE1	2.54	0.40
1:A:42:VAL:HG12	1:C:567:ARG:NH2	2.37	0.40
1:A:1089:PHE:HB3	1:B:913:GLN:HE21	1.86	0.40
1:B:280:ASN:OD1	1:B:284:THR:N	2.53	0.40
1:B:676:THR:HA	1:B:690:GLN:HG2	2.02	0.40
1:B:1110:TYR:CZ	1:B:1112:PRO:HG3	2.56	0.40
1:C:878:LEU:HD12	1:C:878:LEU:HA	1.88	0.40
1:A:331:ASN:OD1	1:A:331:ASN:N	2.44	0.40
1:A:1045:LYS:HD3	1:A:1045:LYS:HA	1.83	0.40
1:B:44:ARG:HB2	1:B:279:TYR:CD2	2.57	0.40
1:B:133:PHE:HD1	1:B:160:TYR:HB3	1.86	0.40
1:C:869:MET:HE2	1:C:869:MET:HB3	1.99	0.40
1:B:195:LYS:HD2	1:B:202:LYS:HD3	2.03	0.40
1:B:962:LEU:HD12	1:B:962:LEU:HA	1.95	0.40
1:C:110:LEU:HD22	1:C:135:PHE:HE1	1.86	0.40
1:A:805:ILE:HG13	1:A:1054:GLN:NE2	2.36	0.40
1:B:462:LYS:H	1:B:462:LYS:HE2	1.86	0.40
1:C:1083:HIS:HE1	1:C:1136:THR:HG23	1.87	0.40
1:C:406:GLU:OE1	1:C:406:GLU:N	2.54	0.40
1:A:291:CYS:HB3	1:A:301:CYS:HB2	1.35	0.40
1:A:650:LEU:HD12	1:A:650:LEU:HA	1.95	0.40
1:A:1081:ILE:HD11	1:A:1135:ASN:HB3	2.02	0.40
1:C:295:PRO:O	1:C:299:THR:HG23	2.22	0.40
1:A:278:LYS:HB2	1:A:306:PHE:CE2	2.56	0.40
1:A:454:ARG:HD3	1:A:457:ARG:HD3	2.03	0.40
1:B:278:LYS:HB2	1:B:306:PHE:CZ	2.57	0.40
1:C:295:PRO:O	1:C:299:THR:HG23	2.21	0.40
1:A:683:CYS:SG	1:A:697:MET:HE3	2.61	0.40
1:B:598:ILE:HB	1:B:609:ALA:HB3	2.02	0.40
1:B:742:ILE:HD13	1:B:742:ILE:HA	1.95	0.40
1:A:1155:TYR:O	1:A:1159:HIS:ND1	2.34	0.40
1:B:596:SER:O	1:B:611:LEU:N	2.41	0.40
1:C:1004:LEU:O	1:C:1008:VAL:HG13	2.21	0.40
1:C:1031:GLU:HB3	1:C:1037:SER:HB2	2.03	0.40
1:A:54:LEU:HD23	1:A:197:ILE:HD11	2.02	0.40
1:C:1031:GLU:OE2	1:C:1039:ARG:NH2	2.55	0.40
1:C:106:PHE:O	1:C:117:LEU:N	2.40	0.40
1:C:131:CYS:HB2	1:C:133:PHE:CZ	2.56	0.40

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:B:570:ALA:O	1:C:966:LEU:HD23	2.21	0.40
1:B:1115:ILE:HG22	1:B:1137:VAL:HG23	2.03	0.40
1:A:462:LYS:HE2	1:A:462:LYS:H	1.85	0.40
1:A:993:ILE:O	1:A:997:ILE:HG12	2.21	0.40
1:A:712:ILE:HA	1:A:712:ILE:HD13	1.78	0.40
1:A:802:PHE:HE2	1:A:927:PHE:HE2	1.70	0.40
1:B:200:TYR:HA	1:B:230:PRO:HA	2.03	0.40
1:B:462:LYS:H	1:B:462:LYS:HE2	1.87	0.40
1:A:978:ASN:HA	1:A:981:LEU:HG	2.04	0.40
1:B:439:ASN:OD1	1:B:440:ASN:N	2.53	0.40
1:B:735:SER:HB2	1:B:861:LEU:HD21	2.03	0.40
1:B:108:THR:OG1	1:B:234:ASN:O	2.39	0.40
1:B:759:PHE:CD1	1:B:1001:LEU:HD11	2.56	0.40
1:C:717:ASN:OD1	1:C:718:PHE:N	2.51	0.40
1:C:950:ASP:O	1:C:954:GLN:HG2	2.21	0.40
1:A:229:LEU:HG	1:A:231:ILE:HG12	2.04	0.40
1:B:332:ILE:HG13	1:B:362:VAL:HG22	2.02	0.40
1:C:882:ILE:HD12	1:C:882:ILE:HA	1.86	0.40
1:A:403:ARG:HG2	1:A:497:PHE:HE1	1.85	0.40
1:B:131:CYS:HB2	1:B:133:PHE:CE1	2.56	0.40
1:B:321:GLN:OE1	1:B:321:GLN:N	2.53	0.40
1:B:874:THR:O	1:B:878:LEU:HD23	2.21	0.40
1:C:337:PRO:HD3	1:C:358:ILE:HD12	2.02	0.40
1:A:389:ASP:HA	1:A:528:LYS:HE3	2.03	0.40
1:B:388:ASN:OD1	1:B:388:ASN:O	2.39	0.40
1:B:429:PHE:HE2	1:B:514:SER:HA	1.86	0.40
1:B:513:LEU:HA	1:B:513:LEU:HD12	1.84	0.40
1:B:973:ILE:HD12	1:B:973:ILE:HG23	1.86	0.40
1:A:189:LEU:HB3	1:A:208:THR:HB	2.04	0.40
1:A:547:THR:O	1:B:978:ASN:ND2	2.54	0.40
1:A:759:PHE:HA	1:A:762:GLN:HE21	1.85	0.40
1:B:326:ILE:HD13	1:B:533:LEU:HD12	2.04	0.40
1:B:439:ASN:OD1	1:B:440:ASN:N	2.53	0.40
1:C:106:PHE:O	1:C:117:LEU:N	2.44	0.40
1:A:58:PHE:CD2	1:A:290:ASP:HB2	2.56	0.40
1:B:408:ARG:HH12	1:C:376:THR:HA	1.87	0.40
1:B:415:THR:HG21	1:C:384:PRO:HG2	2.04	0.40
1:C:195:LYS:HG3	1:C:197:ILE:HG22	2.04	0.40
1:A:1094:VAL:N	1:A:1105:THR:O	2.43	0.40
1:C:303:LEU:HD12	1:C:308:VAL:HG12	2.04	0.40
1:A:114:THR:HG22	1:C:469:SER:HB2	2.02	0.40

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:B:873:TYR:O	1:B:877:LEU:HB2	2.22	0.40
1:B:1090:PRO:HD3	1:B:1095:PHE:HE2	1.87	0.40
1:C:22:THR:OG1	1:C:78:ARG:NH1	2.52	0.40
1:A:862:PRO:HA	1:A:863:PRO:HD3	1.97	0.40
1:B:194:PHE:HB3	1:B:201:PHE:HE1	1.86	0.40
1:B:503:VAL:HA	1:B:506:GLN:HG3	2.02	0.40
1:B:1002:GLN:O	1:B:1006:THR:HG23	2.21	0.40
1:C:955:ASN:HD22	1:C:955:ASN:HA	1.64	0.40
1:A:80:ALA:O	1:A:245:HIS:NE2	2.55	0.40
1:A:867:ASP:OD1	1:A:867:ASP:N	2.54	0.40
1:B:276:LEU:HD23	1:B:289:VAL:HB	2.04	0.40
1:C:34:ARG:NH2	1:C:221:SER:H	2.19	0.40
1:C:738:CYS:HB3	1:C:760:CYS:HB2	1.67	0.40
1:B:915:VAL:O	1:B:919:ASN:HB2	2.22	0.40
1:C:989:ALA:O	1:C:993:ILE:HG12	2.21	0.40
1:A:58:PHE:HB2	1:A:293:LEU:HD21	2.04	0.40
1:C:33:THR:HB	1:C:220:PHE:HD1	1.86	0.40
1:C:92:PHE:HE1	1:C:265:TYR:HB2	1.86	0.40
1:C:96:GLU:HG2	1:C:99:ASN:HA	2.03	0.40
1:C:106:PHE:O	1:C:117:LEU:N	2.54	0.40
1:C:131:CYS:HB2	1:C:133:PHE:CE1	2.56	0.40
1:B:742:ILE:HD13	1:B:742:ILE:HA	1.93	0.40
1:B:756:TYR:HE2	1:B:997:ILE:HD12	1.86	0.40
1:C:58:PHE:CD2	1:C:290:ASP:HB2	2.56	0.40
1:C:537:LYS:HE3	1:C:537:LYS:HB3	1.90	0.40
1:C:936:ASP:O	1:C:940:SER:OG	2.32	0.40
1:B:204:TYR:CE2	1:B:225:PRO:HG3	2.57	0.40
1:C:537:LYS:HE3	1:C:537:LYS:HB3	1.89	0.40
1:C:97:LYS:HD2	1:C:97:LYS:HA	1.86	0.40
1:A:431:GLY:HA2	1:A:515:PHE:CE2	2.56	0.40
1:B:730:SER:OG	1:B:731:MET:N	2.53	0.40
1:B:738:CYS:O	1:B:742:ILE:HB	2.21	0.40
1:C:606:ASN:C	1:C:606:ASN:ND2	2.77	0.40
1:C:1002:GLN:O	1:C:1006:THR:HG23	2.22	0.40
1:A:676:THR:HA	1:A:690:GLN:HG2	2.02	0.40
1:A:969:ASN:OD1	1:B:755:GLN:NE2	2.54	0.40
1:A:1102:TRP:HD1	1:A:1135:ASN:HD22	1.70	0.40
1:B:954:GLN:HG3	1:B:1014:ARG:NE	2.37	0.40
1:A:106:PHE:HD2	1:A:117:LEU:HD22	1.85	0.40
1:A:546:LEU:HD21	1:A:573:THR:HG21	2.03	0.40
1:A:278:LYS:HD2	1:A:306:PHE:CE2	2.57	0.40

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:B:551:VAL:HG12	1:B:553:THR:HG23	2.03	0.40
1:C:379:CYS:HA	1:C:432:CYS:HA	2.03	0.40
1:B:339:GLY:O	1:B:343:ASN:HB2	2.21	0.40
1:B:342:PHE:HE1	1:B:511:VAL:HG11	1.85	0.40
1:B:725:GLU:HG3	1:B:1064:HIS:CD2	2.56	0.40
1:B:726:ILE:HG22	1:B:1061:VAL:HG22	2.04	0.40
1:C:106:PHE:CD1	1:C:117:LEU:HD23	2.56	0.40
1:A:206:LYS:HD3	1:A:224:GLU:H	1.86	0.40
1:A:733:LYS:HB3	1:A:861:LEU:HD23	2.04	0.40
1:A:1002:GLN:O	1:A:1006:THR:HG23	2.21	0.40
1:B:97:LYS:HD2	1:B:97:LYS:HA	1.81	0.40
1:C:332:ILE:HG13	1:C:362:VAL:HG22	2.03	0.40
1:C:738:CYS:SG	1:C:739:THR:N	2.94	0.40
1:C:866:THR:O	1:C:870:ILE:HG12	2.22	0.40
1:C:1033:VAL:HG21	1:C:1053:PRO:HG3	2.04	0.40
1:A:563:GLN:HG2	1:B:43:PHE:CD1	2.55	0.40
1:C:719:THR:HG23	1:C:1070:ALA:HB2	2.03	0.40
1:A:206:LYS:HD2	1:A:223:LEU:HA	2.03	0.40
1:A:1105:THR:HG21	1:A:1110:TYR:HA	2.04	0.40
1:B:574:ASP:OD1	1:B:574:ASP:N	2.46	0.40
1:B:592:PHE:HZ	1:C:740:MET:HE2	1.86	0.40
1:B:862:PRO:HA	1:B:863:PRO:HD3	2.00	0.40
1:C:295:PRO:O	1:C:299:THR:HG23	2.22	0.40
1:C:342:PHE:HE2	1:C:511:VAL:HG11	1.86	0.40
1:B:389:ASP:HA	1:B:528:LYS:HE3	2.04	0.40
1:C:983:ARG:HE	1:C:983:ARG:HB2	1.54	0.40
1:A:309:GLU:OE2	1:A:309:GLU:N	2.55	0.40
1:B:196:ASN:OD1	1:B:196:ASN:N	2.54	0.40
1:C:369:TYR:CZ	1:C:384:PRO:HB2	2.56	0.40
1:A:1002:GLN:O	1:A:1006:THR:HG23	2.21	0.40
1:B:365:TYR:HE2	1:B:387:LEU:HD12	1.87	0.40
1:B:993:ILE:O	1:B:997:ILE:HG12	2.22	0.40
1:C:143:VAL:HG13	1:C:154:GLU:HA	2.03	0.40
1:C:332:ILE:HG13	1:C:362:VAL:HG22	2.03	0.40
1:C:402:ILE:HD11	1:C:418:ILE:HG13	2.02	0.40
1:C:405:ASP:N	1:C:405:ASP:OD1	2.52	0.40

There are no symmetry-related clashes.

5.3 Torsion angles ⓘ

5.3.1 Protein backbone ⓘ

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

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

Mol	Chain	Analysed	Favoured	Allowed	Outliers	Percentiles	
1	1-A	1067/1230 (87%)	1024 (96%)	42 (4%)	1 (0%)	48	77
1	1-B	1067/1230 (87%)	1020 (96%)	47 (4%)	0	100	100
1	1-C	1067/1230 (87%)	1011 (95%)	54 (5%)	2 (0%)	44	73
1	2-A	1067/1230 (87%)	1025 (96%)	41 (4%)	1 (0%)	48	77
1	2-B	1067/1230 (87%)	1019 (96%)	47 (4%)	1 (0%)	48	77
1	2-C	1067/1230 (87%)	1013 (95%)	53 (5%)	1 (0%)	48	77
1	3-A	1073/1230 (87%)	1038 (97%)	34 (3%)	1 (0%)	48	77
1	3-B	1073/1230 (87%)	1033 (96%)	39 (4%)	1 (0%)	48	77
1	3-C	1073/1230 (87%)	1035 (96%)	37 (3%)	1 (0%)	48	77
1	4-A	1073/1230 (87%)	1038 (97%)	34 (3%)	1 (0%)	48	77
1	4-B	1073/1230 (87%)	1037 (97%)	35 (3%)	1 (0%)	48	77
1	4-C	1073/1230 (87%)	1038 (97%)	34 (3%)	1 (0%)	48	77
1	5-A	1073/1230 (87%)	1038 (97%)	34 (3%)	1 (0%)	48	77
1	5-B	1073/1230 (87%)	1033 (96%)	39 (4%)	1 (0%)	48	77
1	5-C	1073/1230 (87%)	1036 (97%)	36 (3%)	1 (0%)	48	77
1	6-A	1073/1230 (87%)	1036 (97%)	36 (3%)	1 (0%)	48	77
1	6-B	1073/1230 (87%)	1035 (96%)	37 (3%)	1 (0%)	48	77
1	6-C	1073/1230 (87%)	1039 (97%)	33 (3%)	1 (0%)	48	77
1	7-A	1073/1230 (87%)	1037 (97%)	35 (3%)	1 (0%)	48	77
1	7-B	1073/1230 (87%)	1036 (97%)	36 (3%)	1 (0%)	48	77
1	7-C	1073/1230 (87%)	1034 (96%)	38 (4%)	1 (0%)	48	77
1	8-A	1073/1230 (87%)	1037 (97%)	35 (3%)	1 (0%)	48	77
1	8-B	1073/1230 (87%)	1035 (96%)	37 (3%)	1 (0%)	48	77
1	8-C	1073/1230 (87%)	1037 (97%)	35 (3%)	1 (0%)	48	77
1	9-A	1067/1230 (87%)	1023 (96%)	43 (4%)	1 (0%)	48	77

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Mol	Chain	Analysed	Favoured	Allowed	Outliers	Percentiles	
1	9-B	1067/1230 (87%)	1022 (96%)	45 (4%)	0	100	100
1	9-C	1067/1230 (87%)	1016 (95%)	49 (5%)	2 (0%)	44	73
1	10-A	1073/1230 (87%)	1037 (97%)	35 (3%)	1 (0%)	48	77
1	10-B	1073/1230 (87%)	1037 (97%)	35 (3%)	1 (0%)	48	77
1	10-C	1073/1230 (87%)	1038 (97%)	34 (3%)	1 (0%)	48	77
1	11-A	1067/1230 (87%)	1026 (96%)	40 (4%)	1 (0%)	48	77
1	11-B	1067/1230 (87%)	1019 (96%)	48 (4%)	0	100	100
1	11-C	1067/1230 (87%)	1013 (95%)	51 (5%)	3 (0%)	37	67
1	12-A	1073/1230 (87%)	1039 (97%)	33 (3%)	1 (0%)	48	77
1	12-B	1073/1230 (87%)	1037 (97%)	35 (3%)	1 (0%)	48	77
1	12-C	1073/1230 (87%)	1037 (97%)	35 (3%)	1 (0%)	48	77
1	13-A	1073/1230 (87%)	1040 (97%)	32 (3%)	1 (0%)	48	77
1	13-B	1073/1230 (87%)	1036 (97%)	36 (3%)	1 (0%)	48	77
1	13-C	1073/1230 (87%)	1036 (97%)	36 (3%)	1 (0%)	48	77
1	14-A	1073/1230 (87%)	1038 (97%)	34 (3%)	1 (0%)	48	77
1	14-B	1073/1230 (87%)	1036 (97%)	36 (3%)	1 (0%)	48	77
1	14-C	1073/1230 (87%)	1038 (97%)	34 (3%)	1 (0%)	48	77
1	15-A	1073/1230 (87%)	1037 (97%)	35 (3%)	1 (0%)	48	77
1	15-B	1073/1230 (87%)	1039 (97%)	33 (3%)	1 (0%)	48	77
1	15-C	1073/1230 (87%)	1036 (97%)	36 (3%)	1 (0%)	48	77
1	16-A	1073/1230 (87%)	1036 (97%)	36 (3%)	1 (0%)	48	77
1	16-B	1073/1230 (87%)	1037 (97%)	35 (3%)	1 (0%)	48	77
1	16-C	1073/1230 (87%)	1037 (97%)	35 (3%)	1 (0%)	48	77
1	17-A	1073/1230 (87%)	1038 (97%)	34 (3%)	1 (0%)	48	77
1	17-B	1073/1230 (87%)	1037 (97%)	35 (3%)	1 (0%)	48	77
1	17-C	1073/1230 (87%)	1032 (96%)	38 (4%)	3 (0%)	37	67
1	18-A	1073/1230 (87%)	1035 (96%)	37 (3%)	1 (0%)	48	77
1	18-B	1073/1230 (87%)	1037 (97%)	35 (3%)	1 (0%)	48	77
1	18-C	1073/1230 (87%)	1040 (97%)	32 (3%)	1 (0%)	48	77
1	19-A	1073/1230 (87%)	1040 (97%)	32 (3%)	1 (0%)	48	77
1	19-B	1073/1230 (87%)	1037 (97%)	35 (3%)	1 (0%)	48	77

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Mol	Chain	Analysed	Favoured	Allowed	Outliers	Percentiles	
1	19-C	1073/1230 (87%)	1038 (97%)	33 (3%)	2 (0%)	44	73
1	20-A	1073/1230 (87%)	1038 (97%)	34 (3%)	1 (0%)	48	77
1	20-B	1073/1230 (87%)	1035 (96%)	37 (3%)	1 (0%)	48	77
1	20-C	1073/1230 (87%)	1040 (97%)	32 (3%)	1 (0%)	48	77
All	All	64308/73800 (87%)	62001 (96%)	2243 (4%)	64 (0%)	50	77

All (64) Ramachandran outliers are listed below:

Mol	Chain	Res	Type
1	11-C	319	ARG
1	1-C	590	CYS
1	17-C	445	VAL
1	19-C	385	THR
1	2-A	544	ASN
1	1-A	544	ASN
1	1-C	532	ASN
1	2-C	532	ASN
1	9-A	544	ASN
1	9-C	532	ASN
1	11-A	544	ASN
1	11-C	532	ASN
1	13-A	987	PRO
1	20-A	987	PRO
1	3-A	987	PRO
1	3-B	987	PRO
1	3-C	987	PRO
1	4-A	987	PRO
1	4-B	987	PRO
1	4-C	987	PRO
1	5-A	987	PRO
1	5-B	987	PRO
1	5-C	987	PRO
1	6-A	987	PRO
1	6-B	987	PRO
1	6-C	987	PRO
1	7-A	987	PRO
1	7-B	987	PRO
1	7-C	987	PRO
1	8-B	987	PRO
1	8-C	987	PRO
1	10-A	987	PRO

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Mol	Chain	Res	Type
1	10-B	987	PRO
1	10-C	987	PRO
1	12-A	987	PRO
1	12-B	987	PRO
1	12-C	987	PRO
1	13-B	987	PRO
1	13-C	987	PRO
1	14-A	987	PRO
1	14-B	987	PRO
1	14-C	987	PRO
1	15-A	987	PRO
1	15-B	987	PRO
1	15-C	987	PRO
1	16-B	987	PRO
1	16-C	987	PRO
1	17-A	987	PRO
1	17-B	987	PRO
1	17-C	440	ASN
1	17-C	987	PRO
1	18-A	987	PRO
1	18-B	987	PRO
1	18-C	987	PRO
1	19-A	987	PRO
1	19-B	987	PRO
1	19-C	987	PRO
1	20-B	987	PRO
1	20-C	987	PRO
1	2-B	150	LYS
1	8-A	987	PRO
1	9-C	150	LYS
1	11-C	150	LYS
1	16-A	987	PRO

5.3.2 Protein sidechains [i](#)

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

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

Mol	Chain	Analysed	Rotameric	Outliers	Percentiles	
1	1-A	944/1067 (88%)	934 (99%)	10 (1%)	70	90
1	1-B	944/1067 (88%)	934 (99%)	10 (1%)	70	90
1	1-C	944/1067 (88%)	935 (99%)	9 (1%)	73	91
1	2-A	944/1067 (88%)	934 (99%)	10 (1%)	70	90
1	2-B	944/1067 (88%)	934 (99%)	10 (1%)	70	90
1	2-C	944/1067 (88%)	935 (99%)	9 (1%)	73	91
1	3-A	949/1067 (89%)	939 (99%)	10 (1%)	70	90
1	3-B	949/1067 (89%)	939 (99%)	10 (1%)	70	90
1	3-C	949/1067 (89%)	940 (99%)	9 (1%)	75	92
1	4-A	949/1067 (89%)	939 (99%)	10 (1%)	70	90
1	4-B	949/1067 (89%)	939 (99%)	10 (1%)	70	90
1	4-C	949/1067 (89%)	940 (99%)	9 (1%)	75	92
1	5-A	949/1067 (89%)	939 (99%)	10 (1%)	70	90
1	5-B	949/1067 (89%)	939 (99%)	10 (1%)	70	90
1	5-C	949/1067 (89%)	940 (99%)	9 (1%)	75	92
1	6-A	949/1067 (89%)	939 (99%)	10 (1%)	70	90
1	6-B	949/1067 (89%)	939 (99%)	10 (1%)	70	90
1	6-C	949/1067 (89%)	940 (99%)	9 (1%)	75	92
1	7-A	949/1067 (89%)	939 (99%)	10 (1%)	70	90
1	7-B	949/1067 (89%)	939 (99%)	10 (1%)	70	90
1	7-C	949/1067 (89%)	940 (99%)	9 (1%)	75	92
1	8-A	949/1067 (89%)	939 (99%)	10 (1%)	70	90
1	8-B	949/1067 (89%)	939 (99%)	10 (1%)	70	90
1	8-C	949/1067 (89%)	940 (99%)	9 (1%)	75	92
1	9-A	944/1067 (88%)	934 (99%)	10 (1%)	70	90
1	9-B	944/1067 (88%)	934 (99%)	10 (1%)	70	90
1	9-C	944/1067 (88%)	935 (99%)	9 (1%)	73	91
1	10-A	949/1067 (89%)	939 (99%)	10 (1%)	70	90
1	10-B	949/1067 (89%)	939 (99%)	10 (1%)	70	90
1	10-C	949/1067 (89%)	940 (99%)	9 (1%)	75	92
1	11-A	944/1067 (88%)	934 (99%)	10 (1%)	70	90
1	11-B	944/1067 (88%)	934 (99%)	10 (1%)	70	90

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Mol	Chain	Analysed	Rotameric	Outliers	Percentiles	
1	11-C	944/1067 (88%)	935 (99%)	9 (1%)	73	91
1	12-A	949/1067 (89%)	939 (99%)	10 (1%)	70	90
1	12-B	949/1067 (89%)	939 (99%)	10 (1%)	70	90
1	12-C	949/1067 (89%)	940 (99%)	9 (1%)	75	92
1	13-A	949/1067 (89%)	939 (99%)	10 (1%)	70	90
1	13-B	949/1067 (89%)	939 (99%)	10 (1%)	70	90
1	13-C	949/1067 (89%)	940 (99%)	9 (1%)	75	92
1	14-A	949/1067 (89%)	939 (99%)	10 (1%)	70	90
1	14-B	949/1067 (89%)	939 (99%)	10 (1%)	70	90
1	14-C	949/1067 (89%)	940 (99%)	9 (1%)	75	92
1	15-A	949/1067 (89%)	939 (99%)	10 (1%)	70	90
1	15-B	949/1067 (89%)	939 (99%)	10 (1%)	70	90
1	15-C	949/1067 (89%)	940 (99%)	9 (1%)	75	92
1	16-A	949/1067 (89%)	939 (99%)	10 (1%)	70	90
1	16-B	949/1067 (89%)	939 (99%)	10 (1%)	70	90
1	16-C	949/1067 (89%)	940 (99%)	9 (1%)	75	92
1	17-A	949/1067 (89%)	939 (99%)	10 (1%)	70	90
1	17-B	949/1067 (89%)	939 (99%)	10 (1%)	70	90
1	17-C	949/1067 (89%)	940 (99%)	9 (1%)	75	92
1	18-A	949/1067 (89%)	939 (99%)	10 (1%)	70	90
1	18-B	949/1067 (89%)	939 (99%)	10 (1%)	70	90
1	18-C	949/1067 (89%)	940 (99%)	9 (1%)	75	92
1	19-A	949/1067 (89%)	939 (99%)	10 (1%)	70	90
1	19-B	949/1067 (89%)	939 (99%)	10 (1%)	70	90
1	19-C	949/1067 (89%)	940 (99%)	9 (1%)	75	92
1	20-A	949/1067 (89%)	939 (99%)	10 (1%)	70	90
1	20-B	949/1067 (89%)	939 (99%)	10 (1%)	70	90
1	20-C	949/1067 (89%)	940 (99%)	9 (1%)	75	92
All	All	56880/64020 (89%)	56300 (99%)	580 (1%)	71	91

All (580) residues with a non-rotameric sidechain are listed below:

Mol	Chain	Res	Type
1	1-A	95	THR
1	1-A	315	THR
1	1-A	703	ASN
1	1-A	727	LEU
1	1-A	765	ARG
1	1-A	878	LEU
1	1-A	916	LEU
1	1-A	965	GLN
1	1-A	1010	GLN
1	1-A	1018	ILE
1	1-B	104	TRP
1	1-B	314	GLN
1	1-B	569	ILE
1	1-B	775	ASP
1	1-B	796	ASP
1	1-B	861	LEU
1	1-B	901	GLN
1	1-B	914	ASN
1	1-B	945	LEU
1	1-B	1138	TYR
1	1-C	223	LEU
1	1-C	650	LEU
1	1-C	753	LEU
1	1-C	762	GLN
1	1-C	774	GLN
1	1-C	775	ASP
1	1-C	861	LEU
1	1-C	1005	GLN
1	1-C	1129	VAL
1	2-A	95	THR
1	2-A	315	THR
1	2-A	703	ASN
1	2-A	727	LEU
1	2-A	765	ARG
1	2-A	878	LEU
1	2-A	916	LEU
1	2-A	965	GLN
1	2-A	1010	GLN
1	2-A	1018	ILE
1	2-B	104	TRP
1	2-B	314	GLN
1	2-B	569	ILE
1	2-B	775	ASP

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Mol	Chain	Res	Type
1	2-B	796	ASP
1	2-B	861	LEU
1	2-B	901	GLN
1	2-B	914	ASN
1	2-B	945	LEU
1	2-B	1138	TYR
1	2-C	223	LEU
1	2-C	650	LEU
1	2-C	753	LEU
1	2-C	762	GLN
1	2-C	774	GLN
1	2-C	775	ASP
1	2-C	861	LEU
1	2-C	1005	GLN
1	2-C	1129	VAL
1	3-A	95	THR
1	3-A	315	THR
1	3-A	703	ASN
1	3-A	727	LEU
1	3-A	765	ARG
1	3-A	878	LEU
1	3-A	916	LEU
1	3-A	965	GLN
1	3-A	1010	GLN
1	3-A	1018	ILE
1	3-B	104	TRP
1	3-B	314	GLN
1	3-B	569	ILE
1	3-B	775	ASP
1	3-B	796	ASP
1	3-B	861	LEU
1	3-B	901	GLN
1	3-B	914	ASN
1	3-B	945	LEU
1	3-B	1138	TYR
1	3-C	223	LEU
1	3-C	650	LEU
1	3-C	753	LEU
1	3-C	762	GLN
1	3-C	774	GLN
1	3-C	775	ASP
1	3-C	861	LEU

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Mol	Chain	Res	Type
1	3-C	1005	GLN
1	3-C	1129	VAL
1	4-A	95	THR
1	4-A	315	THR
1	4-A	703	ASN
1	4-A	727	LEU
1	4-A	765	ARG
1	4-A	878	LEU
1	4-A	916	LEU
1	4-A	965	GLN
1	4-A	1010	GLN
1	4-A	1018	ILE
1	4-B	104	TRP
1	4-B	314	GLN
1	4-B	569	ILE
1	4-B	775	ASP
1	4-B	796	ASP
1	4-B	861	LEU
1	4-B	901	GLN
1	4-B	914	ASN
1	4-B	945	LEU
1	4-B	1138	TYR
1	4-C	223	LEU
1	4-C	650	LEU
1	4-C	753	LEU
1	4-C	762	GLN
1	4-C	774	GLN
1	4-C	775	ASP
1	4-C	861	LEU
1	4-C	1005	GLN
1	4-C	1129	VAL
1	5-A	95	THR
1	5-A	315	THR
1	5-A	703	ASN
1	5-A	727	LEU
1	5-A	765	ARG
1	5-A	878	LEU
1	5-A	916	LEU
1	5-A	965	GLN
1	5-A	1010	GLN
1	5-A	1018	ILE
1	5-B	104	TRP

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Mol	Chain	Res	Type
1	5-B	314	GLN
1	5-B	569	ILE
1	5-B	775	ASP
1	5-B	796	ASP
1	5-B	861	LEU
1	5-B	901	GLN
1	5-B	914	ASN
1	5-B	945	LEU
1	5-B	1138	TYR
1	5-C	223	LEU
1	5-C	650	LEU
1	5-C	753	LEU
1	5-C	762	GLN
1	5-C	774	GLN
1	5-C	775	ASP
1	5-C	861	LEU
1	5-C	1005	GLN
1	5-C	1129	VAL
1	6-A	95	THR
1	6-A	315	THR
1	6-A	703	ASN
1	6-A	727	LEU
1	6-A	765	ARG
1	6-A	878	LEU
1	6-A	916	LEU
1	6-A	965	GLN
1	6-A	1010	GLN
1	6-A	1018	ILE
1	6-B	104	TRP
1	6-B	314	GLN
1	6-B	569	ILE
1	6-B	775	ASP
1	6-B	796	ASP
1	6-B	861	LEU
1	6-B	901	GLN
1	6-B	914	ASN
1	6-B	945	LEU
1	6-B	1138	TYR
1	6-C	223	LEU
1	6-C	650	LEU
1	6-C	753	LEU
1	6-C	762	GLN

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Mol	Chain	Res	Type
1	6-C	774	GLN
1	6-C	775	ASP
1	6-C	861	LEU
1	6-C	1005	GLN
1	6-C	1129	VAL
1	7-A	95	THR
1	7-A	315	THR
1	7-A	703	ASN
1	7-A	727	LEU
1	7-A	765	ARG
1	7-A	878	LEU
1	7-A	916	LEU
1	7-A	965	GLN
1	7-A	1010	GLN
1	7-A	1018	ILE
1	7-B	104	TRP
1	7-B	314	GLN
1	7-B	569	ILE
1	7-B	775	ASP
1	7-B	796	ASP
1	7-B	861	LEU
1	7-B	901	GLN
1	7-B	914	ASN
1	7-B	945	LEU
1	7-B	1138	TYR
1	7-C	223	LEU
1	7-C	650	LEU
1	7-C	753	LEU
1	7-C	762	GLN
1	7-C	774	GLN
1	7-C	775	ASP
1	7-C	861	LEU
1	7-C	1005	GLN
1	7-C	1129	VAL
1	8-A	95	THR
1	8-A	315	THR
1	8-A	703	ASN
1	8-A	727	LEU
1	8-A	765	ARG
1	8-A	878	LEU
1	8-A	916	LEU
1	8-A	965	GLN

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Mol	Chain	Res	Type
1	8-A	1010	GLN
1	8-A	1018	ILE
1	8-B	104	TRP
1	8-B	314	GLN
1	8-B	569	ILE
1	8-B	775	ASP
1	8-B	796	ASP
1	8-B	861	LEU
1	8-B	901	GLN
1	8-B	914	ASN
1	8-B	945	LEU
1	8-B	1138	TYR
1	8-C	223	LEU
1	8-C	650	LEU
1	8-C	753	LEU
1	8-C	762	GLN
1	8-C	774	GLN
1	8-C	775	ASP
1	8-C	861	LEU
1	8-C	1005	GLN
1	8-C	1129	VAL
1	9-A	95	THR
1	9-A	315	THR
1	9-A	703	ASN
1	9-A	727	LEU
1	9-A	765	ARG
1	9-A	878	LEU
1	9-A	916	LEU
1	9-A	965	GLN
1	9-A	1010	GLN
1	9-A	1018	ILE
1	9-B	104	TRP
1	9-B	314	GLN
1	9-B	569	ILE
1	9-B	775	ASP
1	9-B	796	ASP
1	9-B	861	LEU
1	9-B	901	GLN
1	9-B	914	ASN
1	9-B	945	LEU
1	9-B	1138	TYR
1	9-C	223	LEU

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Mol	Chain	Res	Type
1	9-C	650	LEU
1	9-C	753	LEU
1	9-C	762	GLN
1	9-C	774	GLN
1	9-C	775	ASP
1	9-C	861	LEU
1	9-C	1005	GLN
1	9-C	1129	VAL
1	10-A	95	THR
1	10-A	315	THR
1	10-A	703	ASN
1	10-A	727	LEU
1	10-A	765	ARG
1	10-A	878	LEU
1	10-A	916	LEU
1	10-A	965	GLN
1	10-A	1010	GLN
1	10-A	1018	ILE
1	10-B	104	TRP
1	10-B	314	GLN
1	10-B	569	ILE
1	10-B	775	ASP
1	10-B	796	ASP
1	10-B	861	LEU
1	10-B	901	GLN
1	10-B	914	ASN
1	10-B	945	LEU
1	10-B	1138	TYR
1	10-C	223	LEU
1	10-C	650	LEU
1	10-C	753	LEU
1	10-C	762	GLN
1	10-C	774	GLN
1	10-C	775	ASP
1	10-C	861	LEU
1	10-C	1005	GLN
1	10-C	1129	VAL
1	11-A	95	THR
1	11-A	315	THR
1	11-A	703	ASN
1	11-A	727	LEU
1	11-A	765	ARG

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Mol	Chain	Res	Type
1	11-A	878	LEU
1	11-A	916	LEU
1	11-A	965	GLN
1	11-A	1010	GLN
1	11-A	1018	ILE
1	11-B	104	TRP
1	11-B	314	GLN
1	11-B	569	ILE
1	11-B	775	ASP
1	11-B	796	ASP
1	11-B	861	LEU
1	11-B	901	GLN
1	11-B	914	ASN
1	11-B	945	LEU
1	11-B	1138	TYR
1	11-C	223	LEU
1	11-C	650	LEU
1	11-C	753	LEU
1	11-C	762	GLN
1	11-C	774	GLN
1	11-C	775	ASP
1	11-C	861	LEU
1	11-C	1005	GLN
1	11-C	1129	VAL
1	12-A	95	THR
1	12-A	315	THR
1	12-A	703	ASN
1	12-A	727	LEU
1	12-A	765	ARG
1	12-A	878	LEU
1	12-A	916	LEU
1	12-A	965	GLN
1	12-A	1010	GLN
1	12-A	1018	ILE
1	12-B	104	TRP
1	12-B	314	GLN
1	12-B	569	ILE
1	12-B	775	ASP
1	12-B	796	ASP
1	12-B	861	LEU
1	12-B	901	GLN
1	12-B	914	ASN

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Mol	Chain	Res	Type
1	12-B	945	LEU
1	12-B	1138	TYR
1	12-C	223	LEU
1	12-C	650	LEU
1	12-C	753	LEU
1	12-C	762	GLN
1	12-C	774	GLN
1	12-C	775	ASP
1	12-C	861	LEU
1	12-C	1005	GLN
1	12-C	1129	VAL
1	13-A	95	THR
1	13-A	315	THR
1	13-A	703	ASN
1	13-A	727	LEU
1	13-A	765	ARG
1	13-A	878	LEU
1	13-A	916	LEU
1	13-A	965	GLN
1	13-A	1010	GLN
1	13-A	1018	ILE
1	13-B	104	TRP
1	13-B	314	GLN
1	13-B	569	ILE
1	13-B	775	ASP
1	13-B	796	ASP
1	13-B	861	LEU
1	13-B	901	GLN
1	13-B	914	ASN
1	13-B	945	LEU
1	13-B	1138	TYR
1	13-C	223	LEU
1	13-C	650	LEU
1	13-C	753	LEU
1	13-C	762	GLN
1	13-C	774	GLN
1	13-C	775	ASP
1	13-C	861	LEU
1	13-C	1005	GLN
1	13-C	1129	VAL
1	14-A	95	THR
1	14-A	315	THR

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Mol	Chain	Res	Type
1	14-A	703	ASN
1	14-A	727	LEU
1	14-A	765	ARG
1	14-A	878	LEU
1	14-A	916	LEU
1	14-A	965	GLN
1	14-A	1010	GLN
1	14-A	1018	ILE
1	14-B	104	TRP
1	14-B	314	GLN
1	14-B	569	ILE
1	14-B	775	ASP
1	14-B	796	ASP
1	14-B	861	LEU
1	14-B	901	GLN
1	14-B	914	ASN
1	14-B	945	LEU
1	14-B	1138	TYR
1	14-C	223	LEU
1	14-C	650	LEU
1	14-C	753	LEU
1	14-C	762	GLN
1	14-C	774	GLN
1	14-C	775	ASP
1	14-C	861	LEU
1	14-C	1005	GLN
1	14-C	1129	VAL
1	15-A	95	THR
1	15-A	315	THR
1	15-A	703	ASN
1	15-A	727	LEU
1	15-A	765	ARG
1	15-A	878	LEU
1	15-A	916	LEU
1	15-A	965	GLN
1	15-A	1010	GLN
1	15-A	1018	ILE
1	15-B	104	TRP
1	15-B	314	GLN
1	15-B	569	ILE
1	15-B	775	ASP
1	15-B	796	ASP

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Mol	Chain	Res	Type
1	15-B	861	LEU
1	15-B	901	GLN
1	15-B	914	ASN
1	15-B	945	LEU
1	15-B	1138	TYR
1	15-C	223	LEU
1	15-C	650	LEU
1	15-C	753	LEU
1	15-C	762	GLN
1	15-C	774	GLN
1	15-C	775	ASP
1	15-C	861	LEU
1	15-C	1005	GLN
1	15-C	1129	VAL
1	16-A	95	THR
1	16-A	315	THR
1	16-A	703	ASN
1	16-A	727	LEU
1	16-A	765	ARG
1	16-A	878	LEU
1	16-A	916	LEU
1	16-A	965	GLN
1	16-A	1010	GLN
1	16-A	1018	ILE
1	16-B	104	TRP
1	16-B	314	GLN
1	16-B	569	ILE
1	16-B	775	ASP
1	16-B	796	ASP
1	16-B	861	LEU
1	16-B	901	GLN
1	16-B	914	ASN
1	16-B	945	LEU
1	16-B	1138	TYR
1	16-C	223	LEU
1	16-C	650	LEU
1	16-C	753	LEU
1	16-C	762	GLN
1	16-C	774	GLN
1	16-C	775	ASP
1	16-C	861	LEU
1	16-C	1005	GLN

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Mol	Chain	Res	Type
1	16-C	1129	VAL
1	17-A	95	THR
1	17-A	315	THR
1	17-A	703	ASN
1	17-A	727	LEU
1	17-A	765	ARG
1	17-A	878	LEU
1	17-A	916	LEU
1	17-A	965	GLN
1	17-A	1010	GLN
1	17-A	1018	ILE
1	17-B	104	TRP
1	17-B	314	GLN
1	17-B	569	ILE
1	17-B	775	ASP
1	17-B	796	ASP
1	17-B	861	LEU
1	17-B	901	GLN
1	17-B	914	ASN
1	17-B	945	LEU
1	17-B	1138	TYR
1	17-C	223	LEU
1	17-C	650	LEU
1	17-C	753	LEU
1	17-C	762	GLN
1	17-C	774	GLN
1	17-C	775	ASP
1	17-C	861	LEU
1	17-C	1005	GLN
1	17-C	1129	VAL
1	18-A	95	THR
1	18-A	315	THR
1	18-A	703	ASN
1	18-A	727	LEU
1	18-A	765	ARG
1	18-A	878	LEU
1	18-A	916	LEU
1	18-A	965	GLN
1	18-A	1010	GLN
1	18-A	1018	ILE
1	18-B	104	TRP
1	18-B	314	GLN

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Mol	Chain	Res	Type
1	18-B	569	ILE
1	18-B	775	ASP
1	18-B	796	ASP
1	18-B	861	LEU
1	18-B	901	GLN
1	18-B	914	ASN
1	18-B	945	LEU
1	18-B	1138	TYR
1	18-C	223	LEU
1	18-C	650	LEU
1	18-C	753	LEU
1	18-C	762	GLN
1	18-C	774	GLN
1	18-C	775	ASP
1	18-C	861	LEU
1	18-C	1005	GLN
1	18-C	1129	VAL
1	19-A	95	THR
1	19-A	315	THR
1	19-A	703	ASN
1	19-A	727	LEU
1	19-A	765	ARG
1	19-A	878	LEU
1	19-A	916	LEU
1	19-A	965	GLN
1	19-A	1010	GLN
1	19-A	1018	ILE
1	19-B	104	TRP
1	19-B	314	GLN
1	19-B	569	ILE
1	19-B	775	ASP
1	19-B	796	ASP
1	19-B	861	LEU
1	19-B	901	GLN
1	19-B	914	ASN
1	19-B	945	LEU
1	19-B	1138	TYR
1	19-C	223	LEU
1	19-C	650	LEU
1	19-C	753	LEU
1	19-C	762	GLN
1	19-C	774	GLN

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Mol	Chain	Res	Type
1	19-C	775	ASP
1	19-C	861	LEU
1	19-C	1005	GLN
1	19-C	1129	VAL
1	20-A	95	THR
1	20-A	315	THR
1	20-A	703	ASN
1	20-A	727	LEU
1	20-A	765	ARG
1	20-A	878	LEU
1	20-A	916	LEU
1	20-A	965	GLN
1	20-A	1010	GLN
1	20-A	1018	ILE
1	20-B	104	TRP
1	20-B	314	GLN
1	20-B	569	ILE
1	20-B	775	ASP
1	20-B	796	ASP
1	20-B	861	LEU
1	20-B	901	GLN
1	20-B	914	ASN
1	20-B	945	LEU
1	20-B	1138	TYR
1	20-C	223	LEU
1	20-C	650	LEU
1	20-C	753	LEU
1	20-C	762	GLN
1	20-C	774	GLN
1	20-C	775	ASP
1	20-C	861	LEU
1	20-C	1005	GLN
1	20-C	1129	VAL

Sometimes sidechains can be flipped to improve hydrogen bonding and reduce clashes. All (534) such sidechains are listed below:

Mol	Chain	Res	Type
1	1-A	115	GLN
1	1-A	146	HIS
1	1-A	173	GLN
1	1-A	321	GLN
1	1-A	949	GLN

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Mol	Chain	Res	Type
1	1-A	955	ASN
1	1-B	52	GLN
1	1-B	314	GLN
1	1-B	544	ASN
1	1-B	613	GLN
1	1-B	755	GLN
1	1-B	1088	HIS
1	1-B	1101	HIS
1	1-B	1113	GLN
1	1-B	1142	GLN
1	1-C	61	ASN
1	1-C	185	ASN
1	1-C	606	ASN
1	1-C	641	ASN
1	1-C	644	GLN
1	1-C	856	ASN
1	1-C	914	ASN
1	1-C	1113	GLN
1	2-A	115	GLN
1	2-A	137	ASN
1	2-A	146	HIS
1	2-A	271	GLN
1	2-A	321	GLN
1	2-A	394	ASN
1	2-A	474	GLN
1	2-A	536	ASN
1	2-A	644	GLN
1	2-A	755	GLN
1	2-A	992	GLN
1	2-A	1048	HIS
1	2-B	30	ASN
1	2-B	422	ASN
1	2-B	644	GLN
1	2-B	853	GLN
1	2-B	872	GLN
1	2-B	901	GLN
1	2-B	907	ASN
1	2-B	957	GLN
1	2-B	1058	HIS
1	2-C	61	ASN
1	2-C	115	GLN
1	2-C	134	GLN

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Mol	Chain	Res	Type
1	2-C	245	HIS
1	2-C	370	ASN
1	2-C	563	GLN
1	2-C	658	ASN
1	2-C	777	ASN
1	2-C	949	GLN
1	3-A	536	ASN
1	3-A	563	GLN
1	3-A	580	GLN
1	3-A	606	ASN
1	3-A	777	ASN
1	3-A	779	GLN
1	3-A	955	ASN
1	3-A	1002	GLN
1	3-A	1011	GLN
1	3-A	1134	ASN
1	3-B	99	ASN
1	3-B	125	ASN
1	3-B	317	ASN
1	3-B	370	ASN
1	3-B	409	GLN
1	3-B	414	GLN
1	3-B	439	ASN
1	3-B	670	ASN
1	3-B	762	GLN
1	3-B	784	GLN
1	3-B	895	GLN
1	3-B	901	GLN
1	3-B	949	GLN
1	3-B	1005	GLN
1	3-B	1011	GLN
1	3-B	1113	GLN
1	3-C	69	HIS
1	3-C	99	ASN
1	3-C	121	ASN
1	3-C	125	ASN
1	3-C	245	HIS
1	3-C	314	GLN
1	3-C	409	GLN
1	3-C	670	ASN
1	3-C	919	ASN
1	3-C	992	GLN

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Mol	Chain	Res	Type
1	3-C	1011	GLN
1	4-A	66	HIS
1	4-A	69	HIS
1	4-A	125	ASN
1	4-A	613	GLN
1	4-A	1011	GLN
1	4-B	271	GLN
1	4-B	317	ASN
1	4-B	409	GLN
1	4-B	422	ASN
1	4-B	439	ASN
1	4-B	607	GLN
1	4-B	762	GLN
1	4-B	777	ASN
1	4-B	901	GLN
1	4-C	125	ASN
1	4-C	196	ASN
1	4-C	207	HIS
1	4-C	239	GLN
1	4-C	439	ASN
1	4-C	762	GLN
1	4-C	965	GLN
1	5-A	115	GLN
1	5-A	185	ASN
1	5-A	239	GLN
1	5-A	414	GLN
1	5-A	437	ASN
1	5-A	439	ASN
1	5-A	493	GLN
1	5-A	506	GLN
1	5-A	613	GLN
1	5-A	949	GLN
1	5-A	992	GLN
1	5-A	1083	HIS
1	5-A	1088	HIS
1	5-A	1134	ASN
1	5-B	99	ASN
1	5-B	173	GLN
1	5-B	211	ASN
1	5-B	439	ASN
1	5-B	487	ASN
1	5-B	506	GLN

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Mol	Chain	Res	Type
1	5-B	613	GLN
1	5-B	690	GLN
1	5-B	784	GLN
1	5-B	856	ASN
1	5-B	872	GLN
1	5-B	914	ASN
1	5-B	1011	GLN
1	5-C	146	HIS
1	5-C	211	ASN
1	5-C	370	ASN
1	5-C	687	GLN
1	5-C	762	GLN
1	5-C	774	GLN
1	5-C	872	GLN
1	5-C	1071	GLN
1	5-C	1113	GLN
1	6-A	99	ASN
1	6-A	185	ASN
1	6-A	211	ASN
1	6-A	271	GLN
1	6-A	437	ASN
1	6-A	919	ASN
1	6-A	1011	GLN
1	6-A	1134	ASN
1	6-B	125	ASN
1	6-B	439	ASN
1	6-B	506	GLN
1	6-B	762	GLN
1	6-B	935	GLN
1	6-B	1010	GLN
1	6-B	1083	HIS
1	6-B	1113	GLN
1	6-C	239	GLN
1	6-C	370	ASN
1	6-C	409	GLN
1	6-C	762	GLN
1	6-C	957	GLN
1	6-C	1002	GLN
1	7-A	314	GLN
1	7-A	414	GLN
1	7-A	498	GLN
1	7-A	955	ASN

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Mol	Chain	Res	Type
1	7-A	1134	ASN
1	7-B	125	ASN
1	7-B	314	GLN
1	7-B	409	GLN
1	7-B	414	GLN
1	7-B	439	ASN
1	7-B	762	GLN
1	7-B	872	GLN
1	7-B	895	GLN
1	7-B	914	ASN
1	7-B	992	GLN
1	7-B	1113	GLN
1	7-C	125	ASN
1	7-C	670	ASN
1	7-C	690	GLN
1	7-C	1106	GLN
1	8-A	125	ASN
1	8-A	239	GLN
1	8-A	439	ASN
1	8-A	777	ASN
1	8-A	1134	ASN
1	8-B	69	HIS
1	8-B	125	ASN
1	8-B	211	ASN
1	8-B	314	GLN
1	8-B	764	ASN
1	8-B	856	ASN
1	8-B	957	GLN
1	8-B	1113	GLN
1	8-C	125	ASN
1	8-C	370	ASN
1	8-C	439	ASN
1	8-C	784	GLN
1	9-A	30	ASN
1	9-A	81	ASN
1	9-A	146	HIS
1	9-A	173	GLN
1	9-A	207	HIS
1	9-A	321	GLN
1	9-A	644	GLN
1	9-A	658	ASN
1	9-A	755	GLN

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Mol	Chain	Res	Type
1	9-A	777	ASN
1	9-A	853	GLN
1	9-B	49	HIS
1	9-B	69	HIS
1	9-B	115	GLN
1	9-B	173	GLN
1	9-B	439	ASN
1	9-B	644	GLN
1	9-B	658	ASN
1	9-B	856	ASN
1	9-B	895	GLN
1	9-B	926	GLN
1	9-B	957	GLN
1	9-B	1048	HIS
1	9-B	1088	HIS
1	9-B	1125	ASN
1	9-C	61	ASN
1	9-C	188	ASN
1	9-C	321	GLN
1	9-C	644	GLN
1	9-C	949	GLN
1	9-C	969	ASN
1	9-C	1088	HIS
1	9-C	1113	GLN
1	10-A	211	ASN
1	10-A	271	GLN
1	10-A	370	ASN
1	10-A	762	GLN
1	10-A	955	ASN
1	10-A	1011	GLN
1	10-A	1083	HIS
1	10-B	125	ASN
1	10-B	185	ASN
1	10-B	211	ASN
1	10-B	314	GLN
1	10-B	409	GLN
1	10-B	762	GLN
1	10-B	949	GLN
1	10-B	953	ASN
1	10-B	1036	GLN
1	10-B	1083	HIS
1	10-B	1113	GLN

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Mol	Chain	Res	Type
1	10-C	125	ASN
1	10-C	207	HIS
1	10-C	762	GLN
1	10-C	954	GLN
1	10-C	955	ASN
1	10-C	1010	GLN
1	10-C	1011	GLN
1	11-A	146	HIS
1	11-A	173	GLN
1	11-A	321	GLN
1	11-A	394	ASN
1	11-A	474	GLN
1	11-A	613	GLN
1	11-A	658	ASN
1	11-A	969	ASN
1	11-A	1005	GLN
1	11-B	115	GLN
1	11-B	137	ASN
1	11-B	173	GLN
1	11-B	314	GLN
1	11-B	450	ASN
1	11-B	498	GLN
1	11-B	544	ASN
1	11-B	644	GLN
1	11-B	856	ASN
1	11-B	957	GLN
1	11-B	978	ASN
1	11-B	1005	GLN
1	11-B	1071	GLN
1	11-B	1101	HIS
1	11-B	1113	GLN
1	11-C	115	GLN
1	11-C	245	HIS
1	11-C	271	GLN
1	11-C	658	ASN
1	11-C	784	GLN
1	11-C	853	GLN
1	11-C	856	ASN
1	11-C	954	GLN
1	11-C	1005	GLN
1	11-C	1058	HIS
1	11-C	1088	HIS

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Mol	Chain	Res	Type
1	11-C	1113	GLN
1	12-A	394	ASN
1	12-A	414	GLN
1	12-A	762	GLN
1	12-A	919	ASN
1	12-A	957	GLN
1	12-B	185	ASN
1	12-B	211	ASN
1	12-B	370	ASN
1	12-B	506	GLN
1	12-B	764	ASN
1	12-B	784	GLN
1	12-B	853	GLN
1	12-B	872	GLN
1	12-B	935	GLN
1	12-B	1113	GLN
1	12-C	69	HIS
1	12-C	115	GLN
1	12-C	125	ASN
1	12-C	414	GLN
1	12-C	955	ASN
1	12-C	957	GLN
1	13-A	125	ASN
1	13-A	414	GLN
1	13-A	498	GLN
1	13-A	506	GLN
1	13-A	920	GLN
1	13-A	957	GLN
1	13-A	1005	GLN
1	13-A	1134	ASN
1	13-B	69	HIS
1	13-B	409	GLN
1	13-B	856	ASN
1	13-B	872	GLN
1	13-B	895	GLN
1	13-B	1083	HIS
1	13-C	414	GLN
1	13-C	506	GLN
1	13-C	670	ASN
1	13-C	710	ASN
1	13-C	762	GLN
1	13-C	853	GLN

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Mol	Chain	Res	Type
1	13-C	907	ASN
1	13-C	955	ASN
1	13-C	1113	GLN
1	14-A	115	GLN
1	14-A	134	GLN
1	14-A	207	HIS
1	14-A	394	ASN
1	14-A	437	ASN
1	14-A	439	ASN
1	14-A	544	ASN
1	14-A	564	GLN
1	14-A	606	ASN
1	14-A	777	ASN
1	14-A	954	GLN
1	14-A	1010	GLN
1	14-A	1011	GLN
1	14-A	1134	ASN
1	14-B	125	ASN
1	14-B	409	GLN
1	14-B	414	GLN
1	14-B	439	ASN
1	14-B	762	GLN
1	14-B	777	ASN
1	14-B	1106	GLN
1	14-B	1142	GLN
1	14-C	125	ASN
1	14-C	134	GLN
1	14-C	321	GLN
1	14-C	437	ASN
1	14-C	670	ASN
1	14-C	954	GLN
1	14-C	965	GLN
1	14-C	1002	GLN
1	14-C	1010	GLN
1	14-C	1113	GLN
1	15-A	69	HIS
1	15-A	99	ASN
1	15-A	125	ASN
1	15-A	146	HIS
1	15-A	414	GLN
1	15-A	439	ASN
1	15-A	498	GLN

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Mol	Chain	Res	Type
1	15-A	536	ASN
1	15-A	606	ASN
1	15-A	670	ASN
1	15-A	1106	GLN
1	15-A	1159	HIS
1	15-B	314	GLN
1	15-B	317	ASN
1	15-B	409	GLN
1	15-B	439	ASN
1	15-B	762	GLN
1	15-B	1058	HIS
1	15-B	1113	GLN
1	15-C	69	HIS
1	15-C	125	ASN
1	15-C	196	ASN
1	15-C	239	GLN
1	15-C	314	GLN
1	15-C	370	ASN
1	15-C	409	GLN
1	15-C	437	ASN
1	15-C	544	ASN
1	15-C	762	GLN
1	15-C	1106	GLN
1	16-A	125	ASN
1	16-A	185	ASN
1	16-A	536	ASN
1	16-A	613	GLN
1	16-A	762	GLN
1	16-A	1002	GLN
1	16-A	1054	GLN
1	16-A	1134	ASN
1	16-B	125	ASN
1	16-B	563	GLN
1	16-B	670	ASN
1	16-B	687	GLN
1	16-B	774	GLN
1	16-B	872	GLN
1	16-B	992	GLN
1	16-B	1005	GLN
1	16-B	1083	HIS
1	16-B	1113	GLN
1	16-C	125	ASN

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Mol	Chain	Res	Type
1	16-C	481	ASN
1	16-C	690	GLN
1	16-C	784	GLN
1	16-C	901	GLN
1	16-C	992	GLN
1	16-C	1113	GLN
1	17-A	239	GLN
1	17-A	409	GLN
1	17-A	670	ASN
1	17-A	919	ASN
1	17-A	955	ASN
1	17-A	1005	GLN
1	17-A	1010	GLN
1	17-A	1106	GLN
1	17-A	1134	ASN
1	17-B	314	GLN
1	17-B	439	ASN
1	17-B	762	GLN
1	17-B	784	GLN
1	17-B	992	GLN
1	17-B	1011	GLN
1	17-B	1083	HIS
1	17-B	1113	GLN
1	17-C	69	HIS
1	17-C	125	ASN
1	17-C	314	GLN
1	17-C	360	ASN
1	17-C	370	ASN
1	17-C	414	GLN
1	17-C	762	GLN
1	17-C	954	GLN
1	17-C	992	GLN
1	17-C	1005	GLN
1	18-A	314	GLN
1	18-A	409	GLN
1	18-A	439	ASN
1	18-A	536	ASN
1	18-A	564	GLN
1	18-A	762	GLN
1	18-A	1054	GLN
1	18-A	1134	ASN
1	18-B	125	ASN

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Mol	Chain	Res	Type
1	18-B	409	GLN
1	18-B	439	ASN
1	18-B	540	ASN
1	18-B	762	GLN
1	18-B	784	GLN
1	18-B	856	ASN
1	18-B	872	GLN
1	18-B	935	GLN
1	18-C	99	ASN
1	18-C	125	ASN
1	18-C	134	GLN
1	18-C	207	HIS
1	18-C	317	ASN
1	18-C	388	ASN
1	18-C	540	ASN
1	18-C	670	ASN
1	18-C	762	GLN
1	18-C	992	GLN
1	19-A	49	HIS
1	19-A	115	GLN
1	19-A	414	GLN
1	19-A	439	ASN
1	19-A	544	ASN
1	19-A	613	GLN
1	19-A	777	ASN
1	19-A	895	GLN
1	19-A	919	ASN
1	19-A	955	ASN
1	19-A	1005	GLN
1	19-A	1010	GLN
1	19-A	1083	HIS
1	19-B	125	ASN
1	19-B	271	GLN
1	19-B	409	GLN
1	19-B	655	HIS
1	19-B	935	GLN
1	19-B	1106	GLN
1	19-B	1113	GLN
1	19-C	125	ASN
1	19-C	437	ASN
1	19-C	487	ASN
1	19-C	540	ASN

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Mol	Chain	Res	Type
1	19-C	1159	HIS
1	20-A	125	ASN
1	20-A	409	GLN
1	20-A	613	GLN
1	20-A	1134	ASN
1	20-B	125	ASN
1	20-B	185	ASN
1	20-B	211	ASN
1	20-B	314	GLN
1	20-B	370	ASN
1	20-B	422	ASN
1	20-B	613	GLN
1	20-B	762	GLN
1	20-B	920	GLN
1	20-C	125	ASN
1	20-C	211	ASN
1	20-C	370	ASN
1	20-C	670	ASN
1	20-C	710	ASN
1	20-C	774	GLN
1	20-C	777	ASN
1	20-C	872	GLN
1	20-C	895	GLN
1	20-C	954	GLN
1	20-C	1108	ASN

5.3.3 RNA ⓘ

There are no RNA molecules in this entry.

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

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

5.5 Carbohydrates ⓘ

There are no oligosaccharides in this entry.

5.6 Ligand geometry [i](#)

There are no ligands in this entry.

5.7 Other polymers [i](#)

There are no such residues in this entry.

5.8 Polymer linkage issues [i](#)

The following chains have linkage breaks:

Mol	Chain	Number of breaks
1	1-C	1

All chain breaks are listed below:

Model	Chain	Residue-1	Atom-1	Residue-2	Atom-2	Distance (Å)
1	C	320:VAL	C	321:GLN	N	1.70

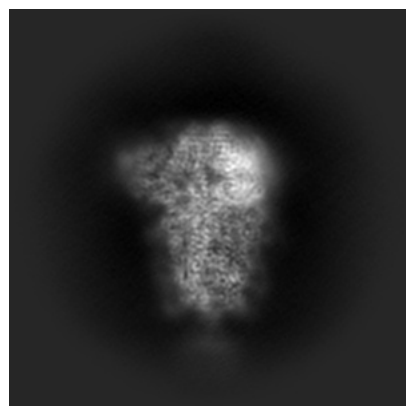
6 Map visualisation [i](#)

This section contains visualisations of the EMDB entry EMD-51280. These allow visual inspection of the internal detail of the map and identification of artifacts.

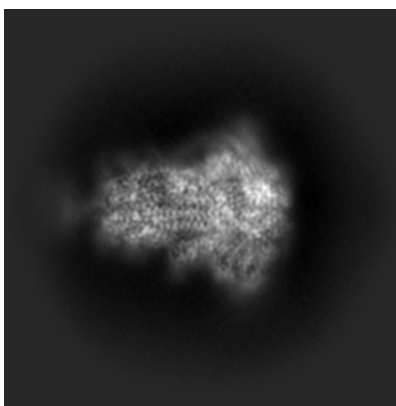
Images derived from a raw map, generated by summing the deposited half-maps, are presented below the corresponding image components of the primary map to allow further visual inspection and comparison with those of the primary map.

6.1 Orthogonal projections [i](#)

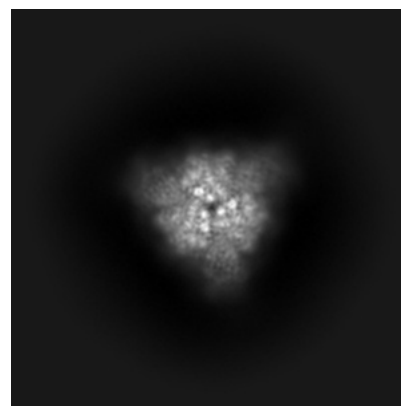
6.1.1 Primary map



X

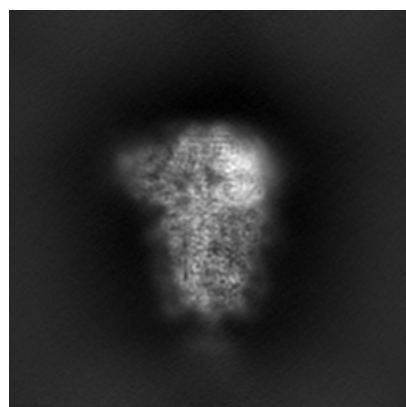


Y

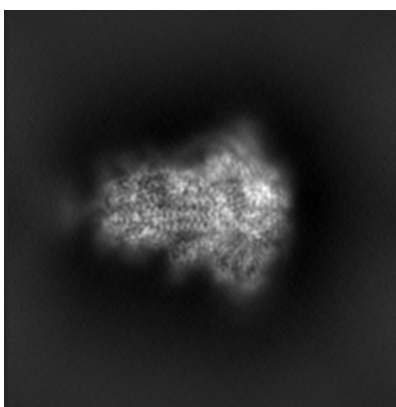


Z

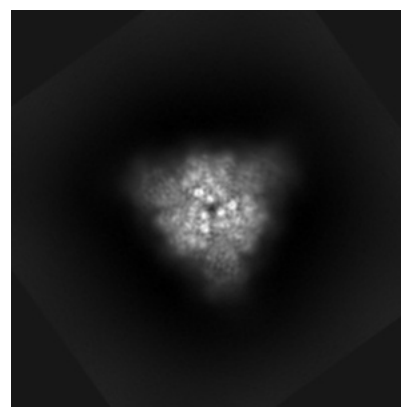
6.1.2 Raw map



X



Y

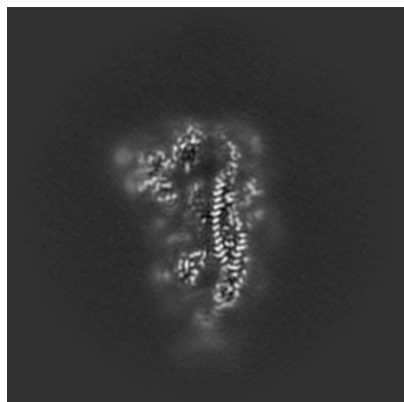


Z

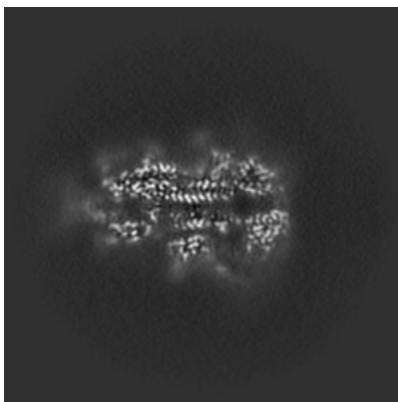
The images above show the map projected in three orthogonal directions.

6.2 Central slices [i](#)

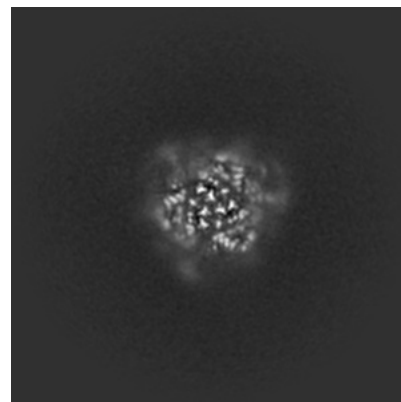
6.2.1 Primary map



X Index: 112

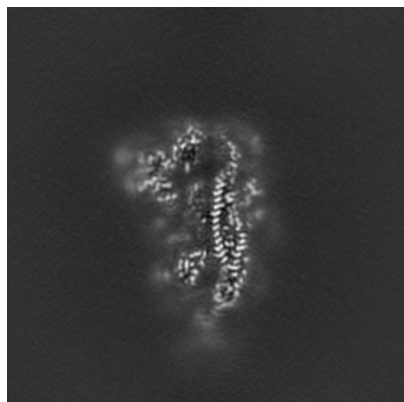


Y Index: 112

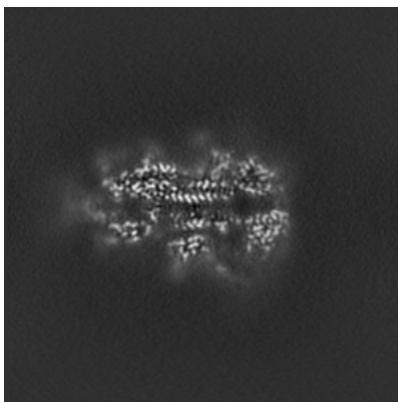


Z Index: 112

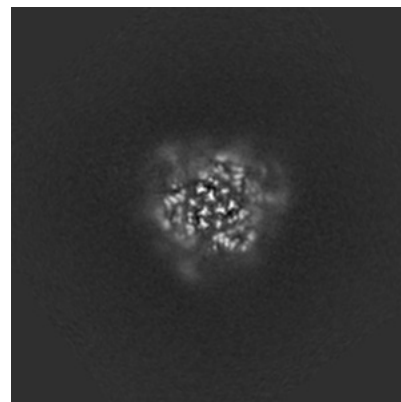
6.2.2 Raw map



X Index: 112



Y Index: 112

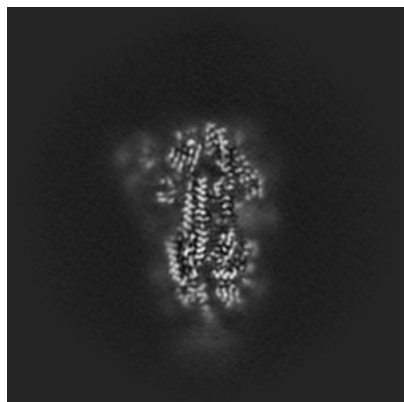


Z Index: 112

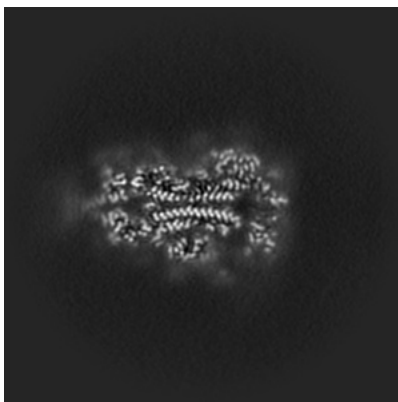
The images above show central slices of the map in three orthogonal directions.

6.3 Largest variance slices [i](#)

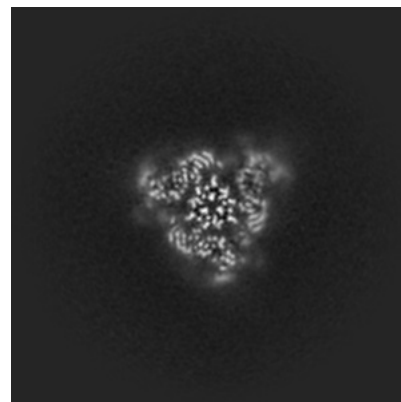
6.3.1 Primary map



X Index: 107

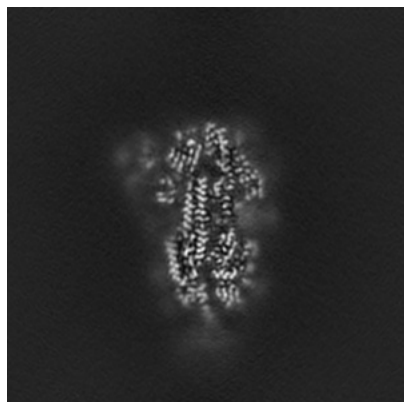


Y Index: 109

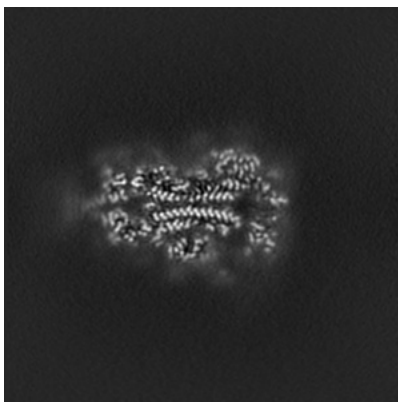


Z Index: 119

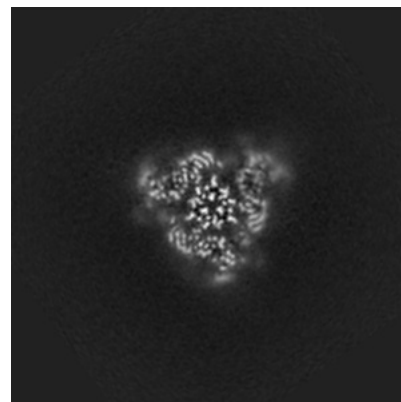
6.3.2 Raw map



X Index: 107



Y Index: 109

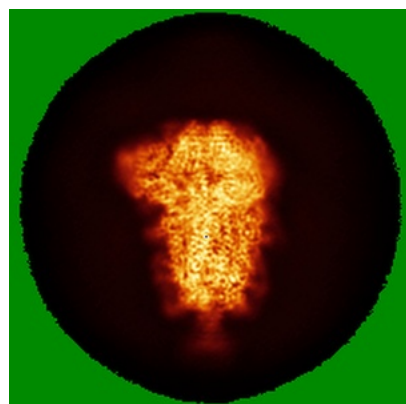


Z Index: 119

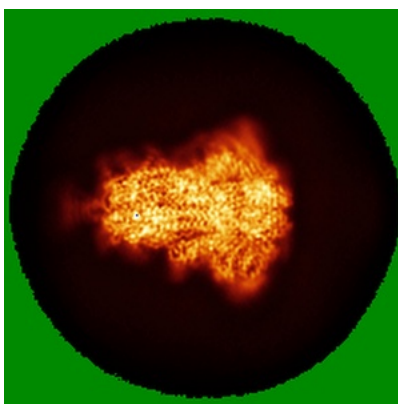
The images above show the largest variance slices of the map in three orthogonal directions.

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

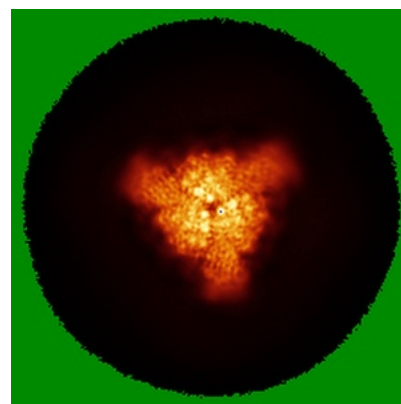
6.4.1 Primary map



X

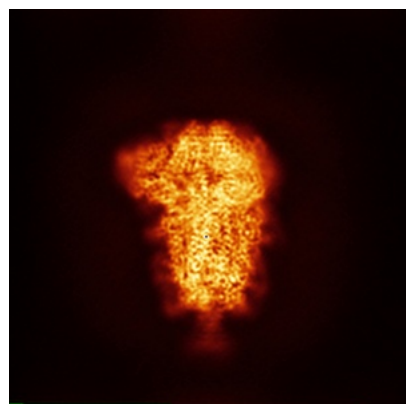


Y

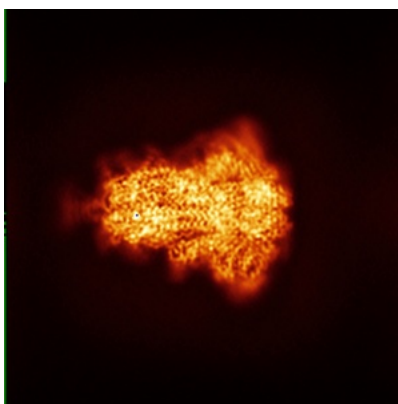


Z

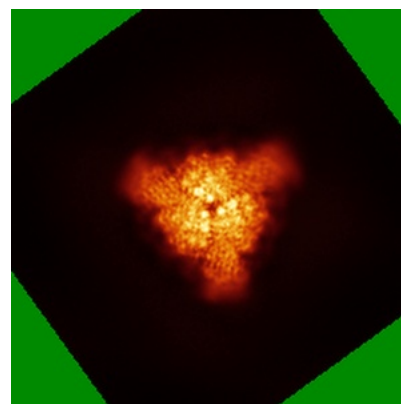
6.4.2 Raw map



X



Y

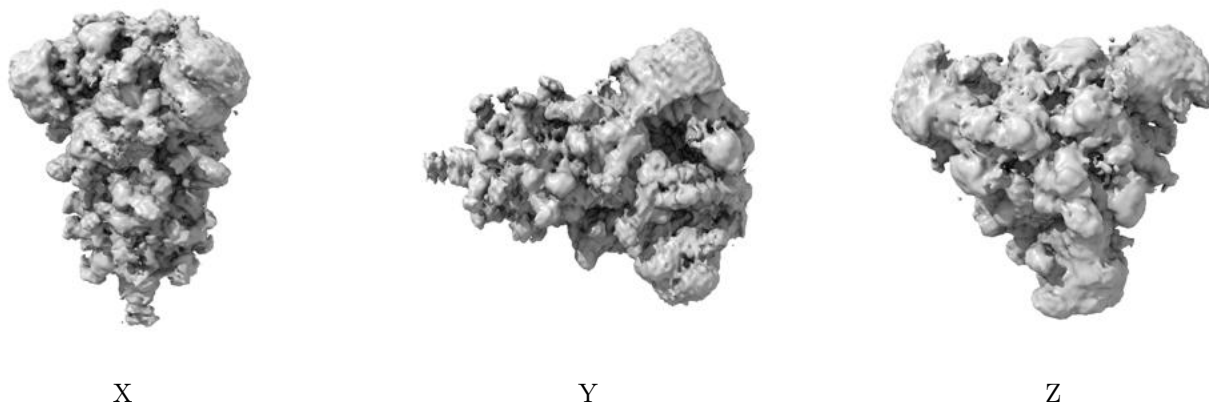


Z

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

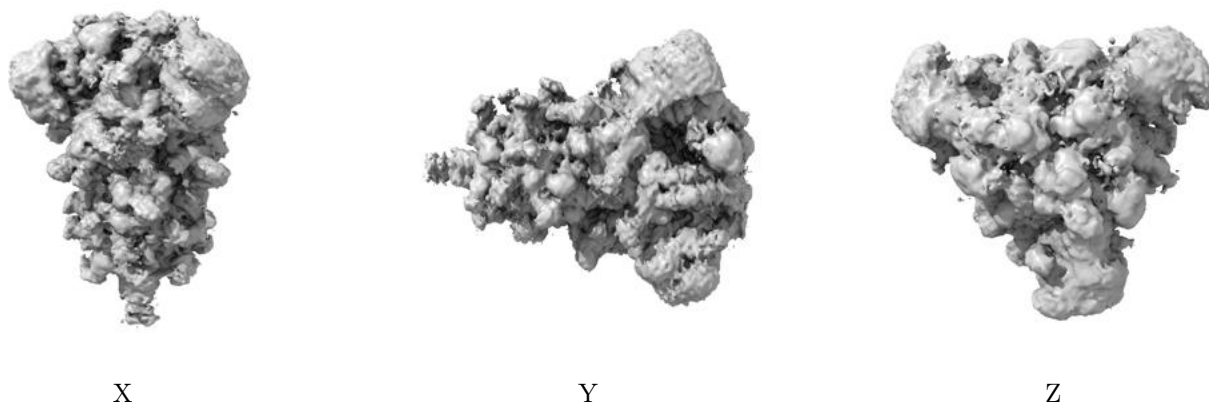
6.5 Orthogonal surface views [i](#)

6.5.1 Primary map



The images above show the 3D surface view of the map at the recommended contour level 0.15. These images, in conjunction with the slice images, may facilitate assessment of whether an appropriate contour level has been provided.

6.5.2 Raw map



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

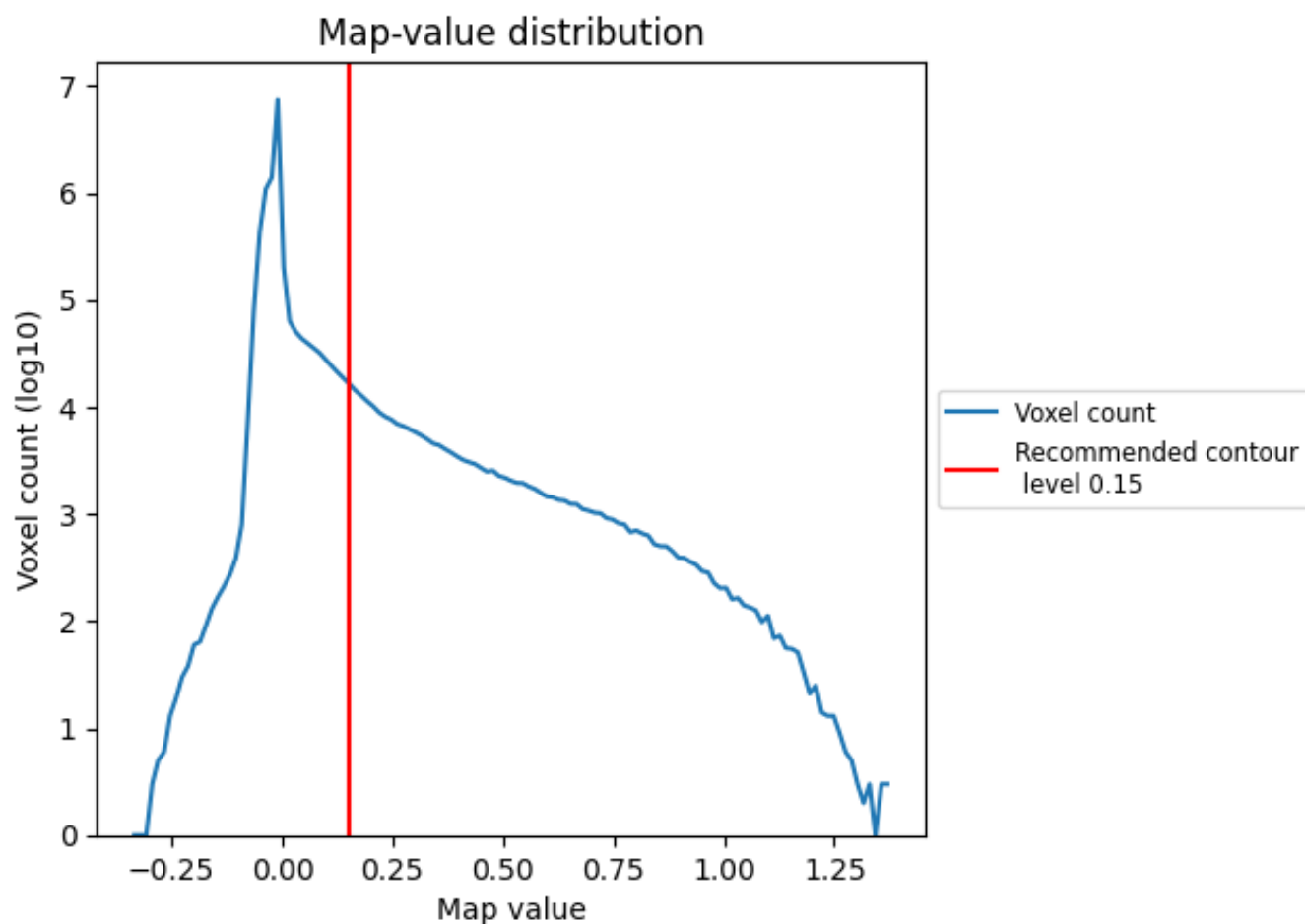
6.6 Mask visualisation [i](#)

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

7 Map analysis [i](#)

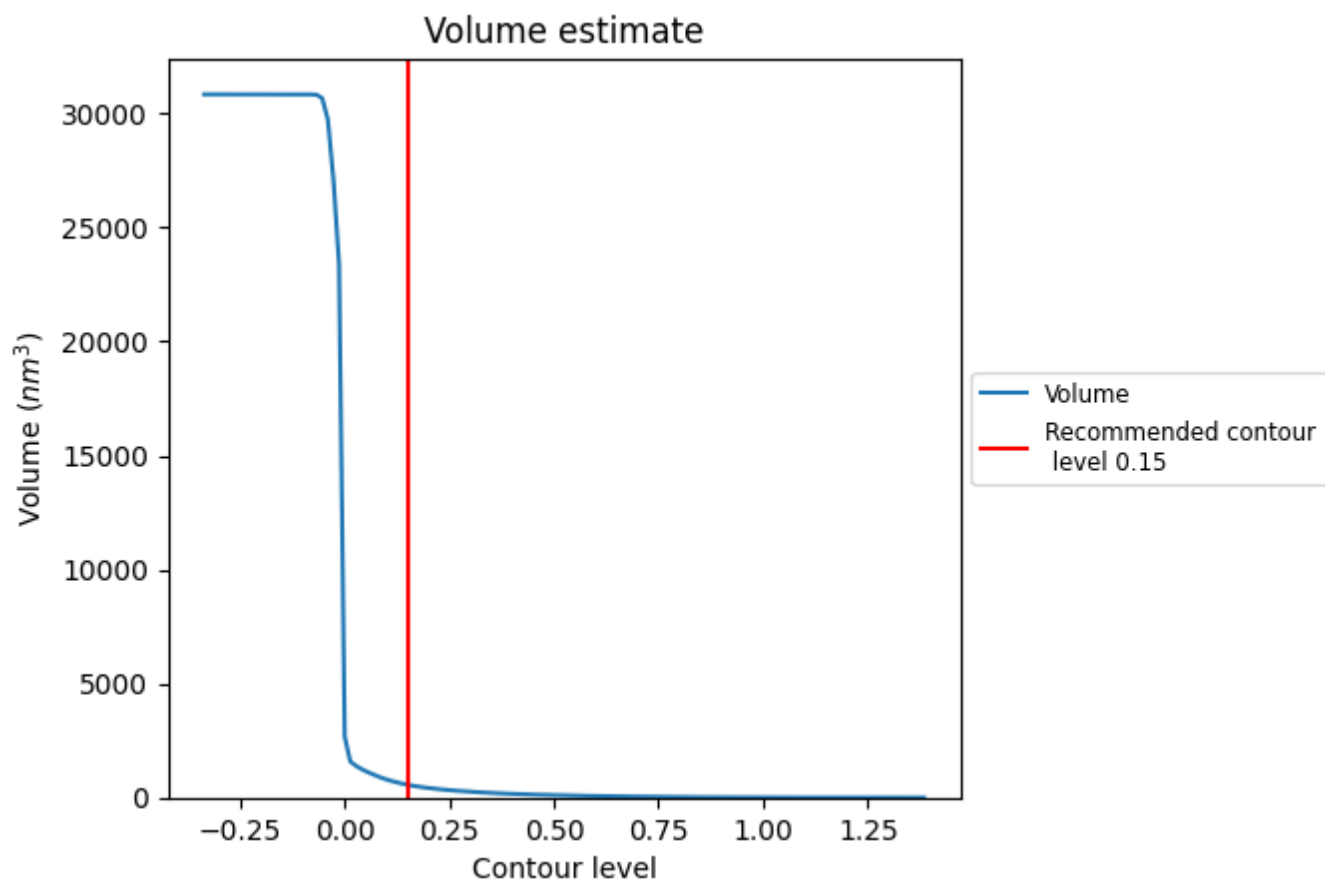
This section contains the results of statistical analysis of the map.

7.1 Map-value distribution [i](#)



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

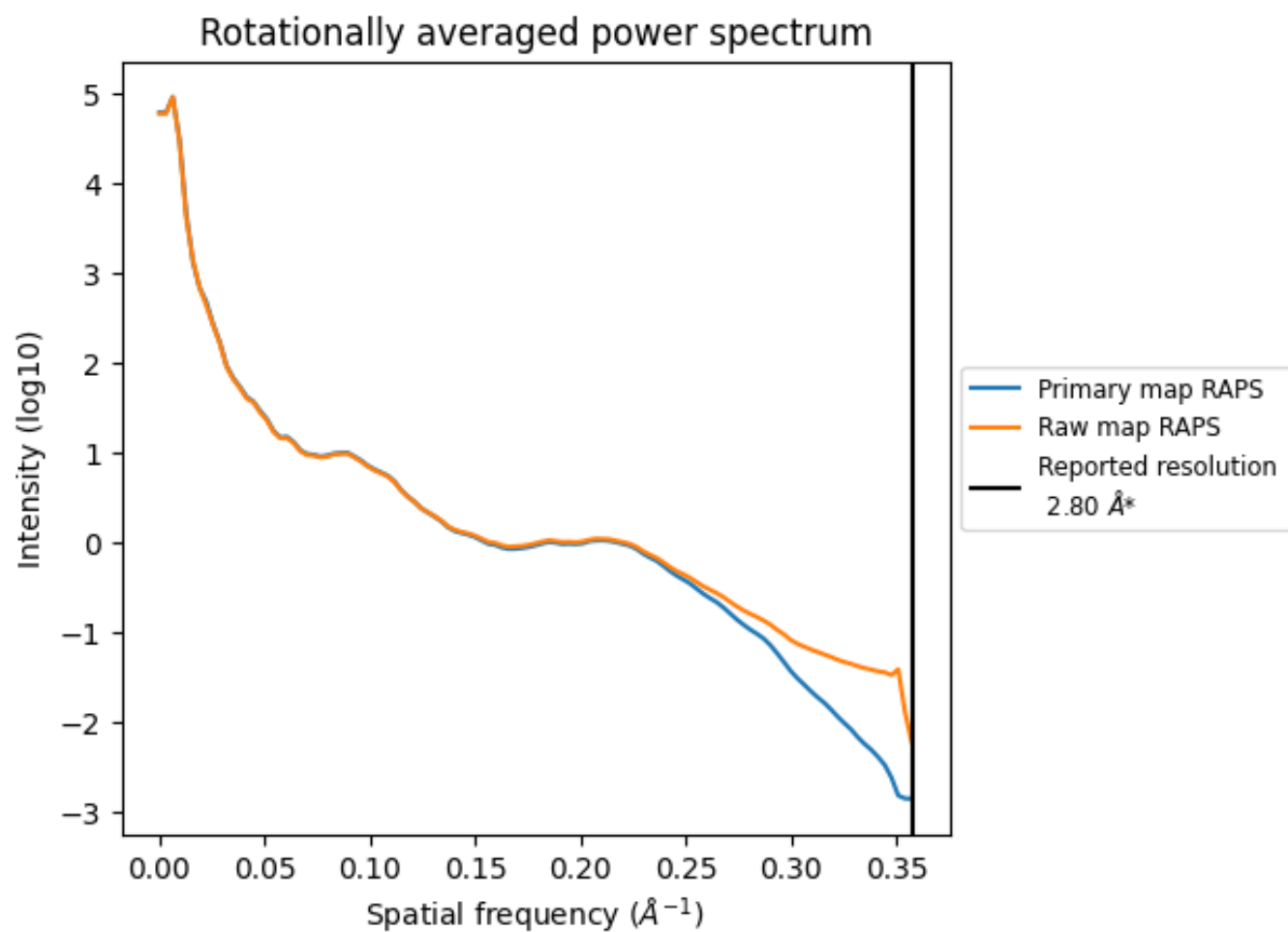
7.2 Volume estimate [i](#)



The volume at the recommended contour level is 564 nm³; this corresponds to an approximate mass of 510 kDa.

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

7.3 Rotationally averaged power spectrum [i](#)

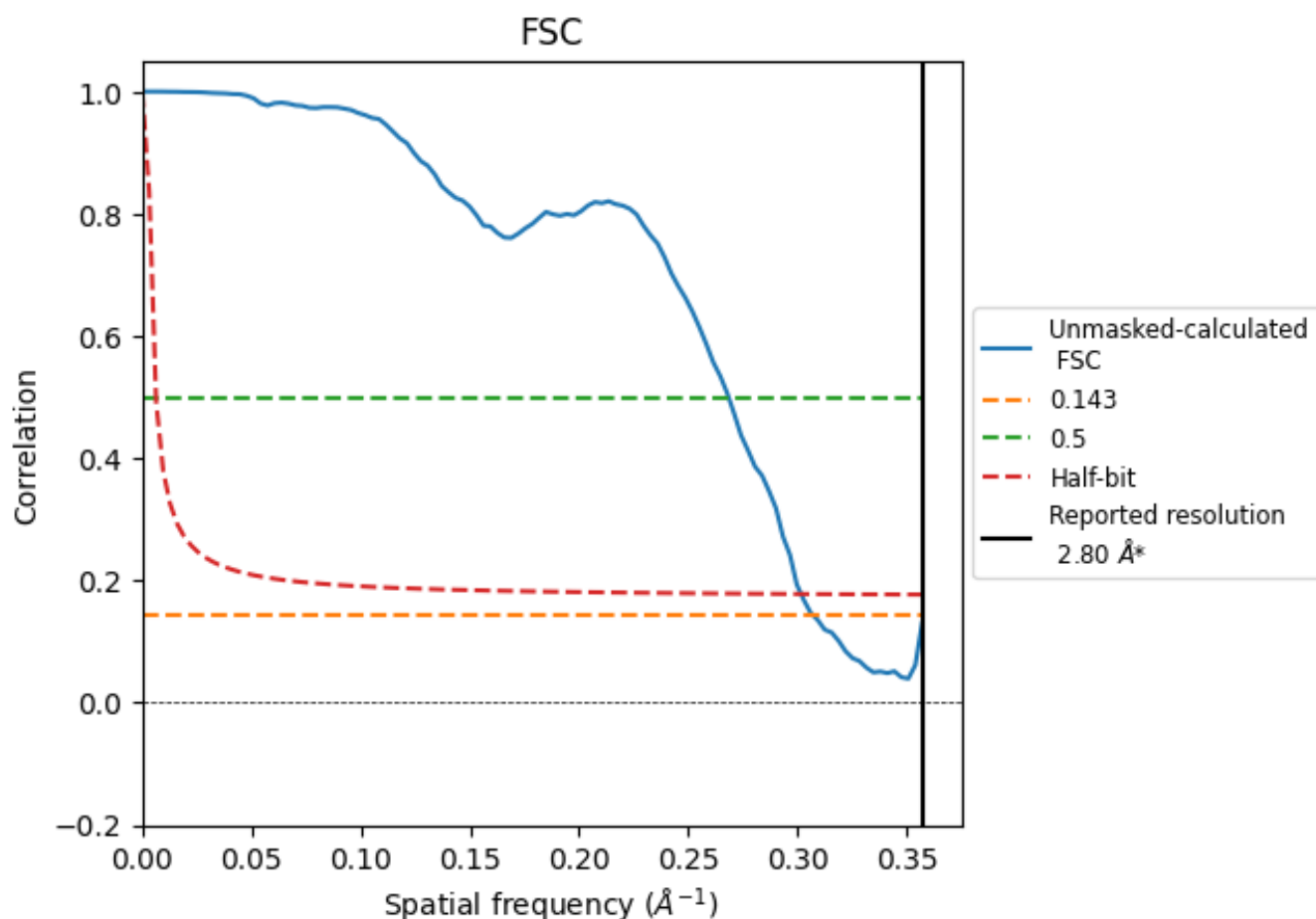


*Reported resolution corresponds to spatial frequency of 0.357 Å⁻¹

8 Fourier-Shell correlation [i](#)

Fourier-Shell Correlation (FSC) is the most commonly used method to estimate the resolution of single-particle and subtomogram-averaged maps. The shape of the curve depends on the imposed symmetry, mask and whether or not the two 3D reconstructions used were processed from a common reference. The reported resolution is shown as a black line. A curve is displayed for the half-bit criterion in addition to lines showing the 0.143 gold standard cut-off and 0.5 cut-off.

8.1 FSC [i](#)



*Reported resolution corresponds to spatial frequency of 0.357 \AA^{-1}

8.2 Resolution estimates [i](#)

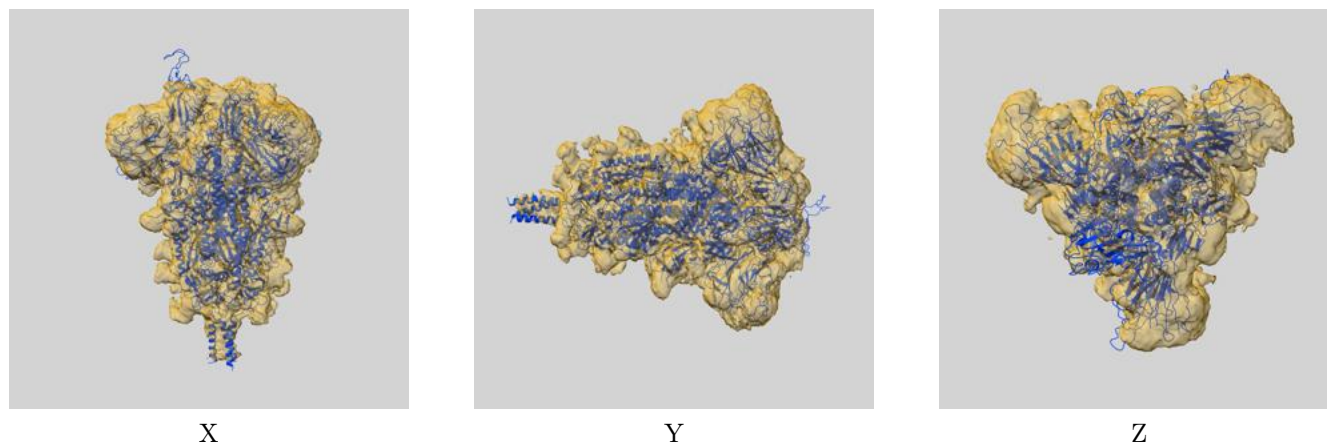
Resolution estimate (Å)	Estimation criterion (FSC cut-off)		
	0.143	0.5	Half-bit
Reported by author	2.80	-	-
Author-provided FSC curve	-	-	-
Unmasked-calculated*	3.26	3.73	3.31

*Resolution estimate based on FSC curve calculated by comparison of deposited half-maps. The value from deposited half-maps intersecting FSC 0.143 CUT-OFF 3.26 differs from the reported value 2.8 by more than 10 %

9 Map-model fit [i](#)

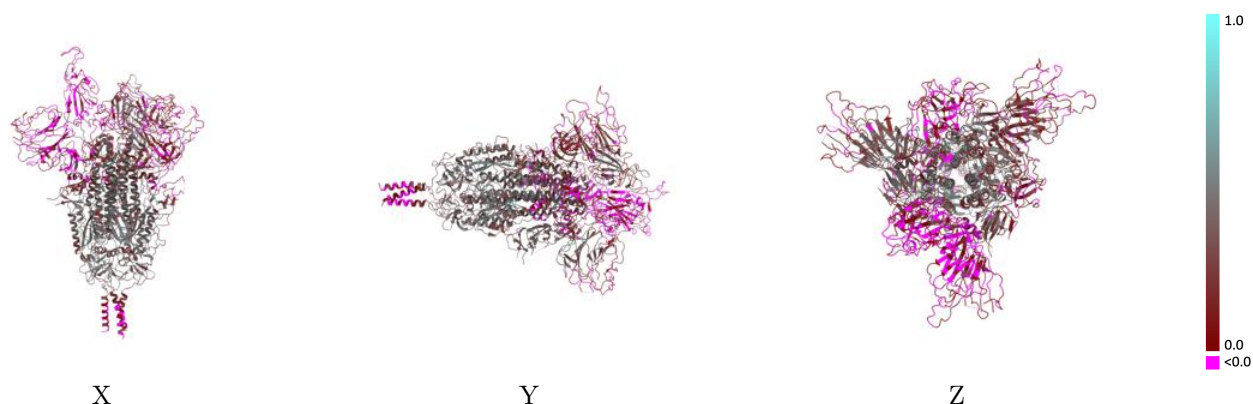
This section contains information regarding the fit between EMDB map EMD-51280 and PDB model 9GDY. Per-residue inclusion information can be found in section [3](#) on page [12](#).

9.1 Map-model overlay [i](#)



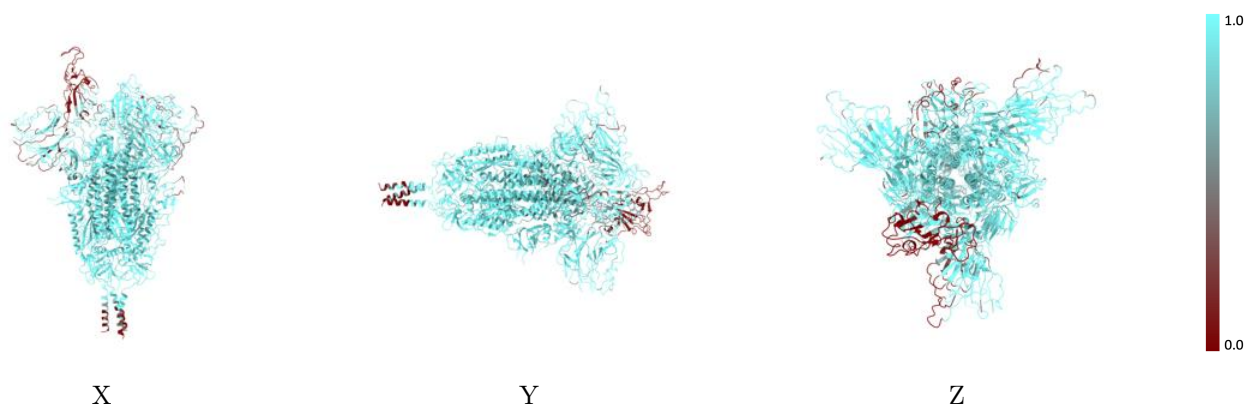
The images above show the 3D surface view of the map at the recommended contour level 0.15 at 50% transparency in yellow overlaid with a ribbon representation of the model coloured in blue. These images allow for the visual assessment of the quality of fit between the atomic model and the map.

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



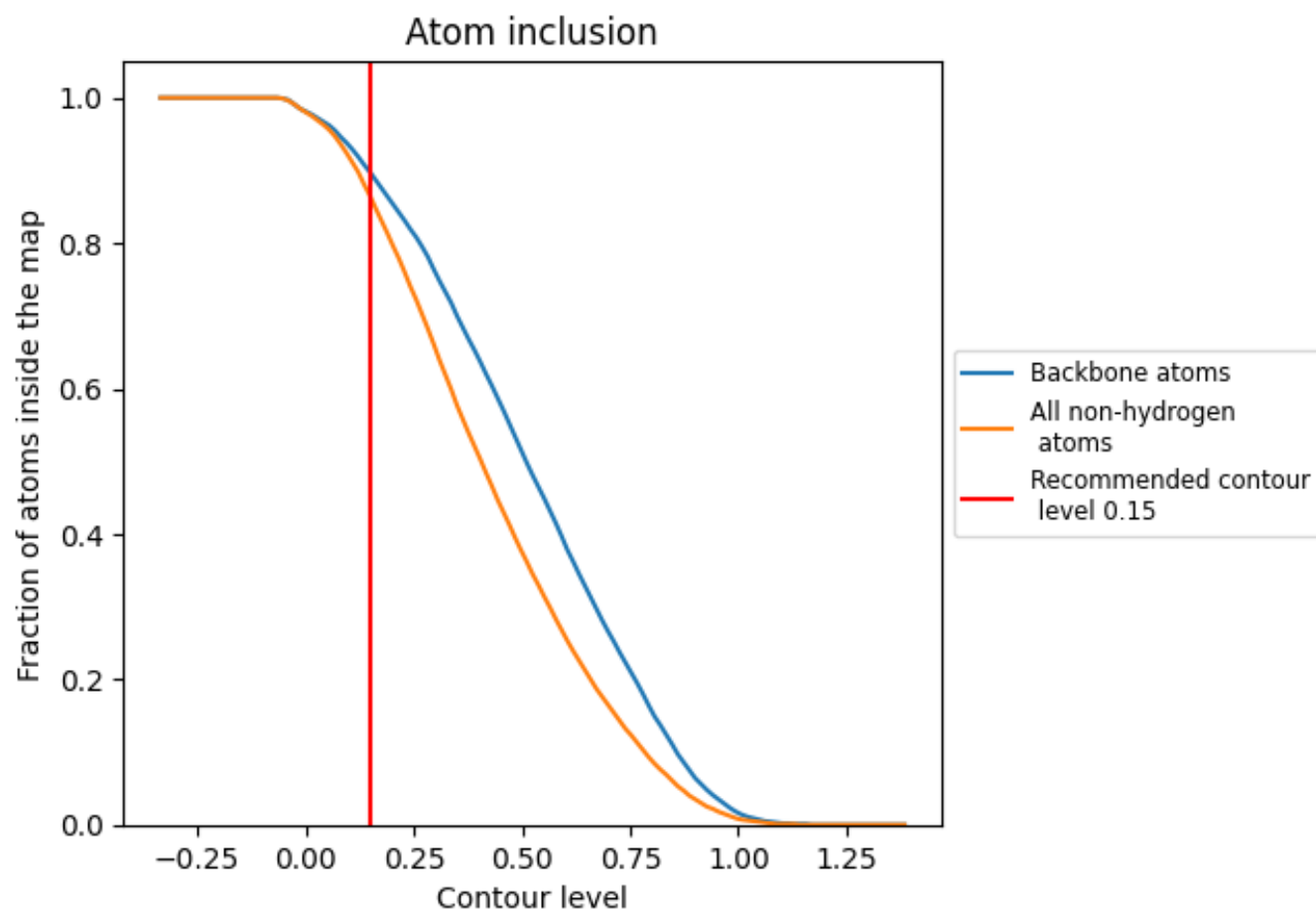
The images above show the model with each residue coloured according to its Q-score. This shows their resolvability in the map with higher Q-score values reflecting better resolvability. Please note: Q-score is calculating the resolvability of atoms, and thus high values are only expected at resolutions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

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



The images above show the model with each residue coloured according to its atom inclusion. This shows to what extent they are inside the map at the recommended contour level (0.15).

9.4 Atom inclusion [i](#)



At the recommended contour level, 90% of all backbone atoms, 86% of all non-hydrogen atoms, are inside the map.

9.5 Map-model fit summary ⓘ

The table lists the average atom inclusion at the recommended contour level (0.15) and Q-score for the entire model and for each chain.

Chain	Atom inclusion	Q-score
All	<div></div> 0.8640	<div></div> 0.2390
A	<div></div> 0.8260	<div></div> 0.2590
B	<div></div> 0.8900	<div></div> 0.2410
C	<div></div> 0.8750	<div></div> 0.2170

