



Full wwPDB EM Validation Report ⓘ

Mar 9, 2026 – 09:43 AM UTC

PDB ID : 4A2I / pdb_00004a2i
EMDB ID : EMD-1895
Title : Cryo-electron Microscopy Structure of the 30S Subunit in Complex with the YjeQ Biogenesis Factor
Authors : Jomaa, A.; Stewart, G.; Mears, J.A.; Kireeva, I.; Brown, E.D.; Ortega, J.
Deposited on : 2011-09-27
Resolution : 16.50 Å (reported)
Based on initial models : 2RCN, 2AVY

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

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

A user guide is available at

<https://www.wwpdb.org/validation/2017/EMValidationReportHelp>
with specific help available everywhere you see the ⓘ symbol.

The types of validation reports are described at

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

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

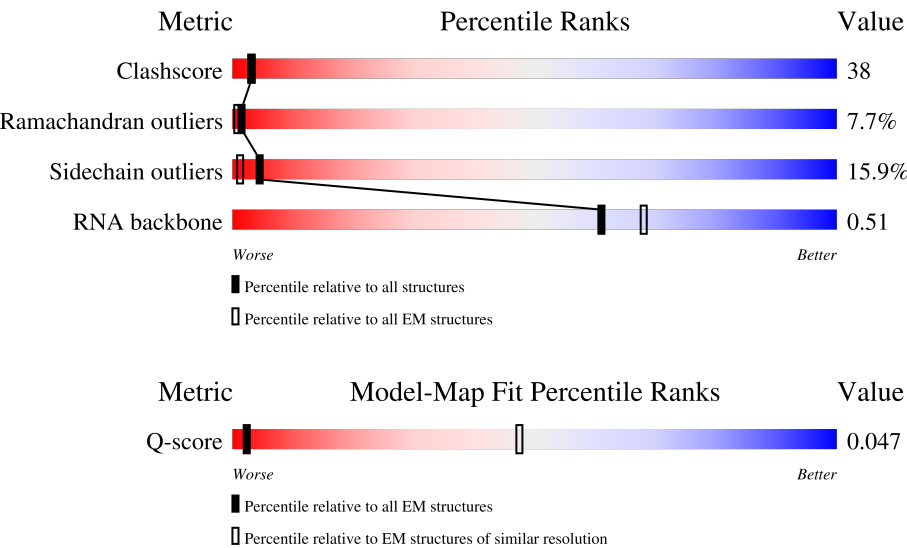
EMDB validation analysis : 0.0.1.dev132
MolProbity : 4-5-2 with Phenix2.0
Percentile statistics : 20250101.v01 (using entries in the PDB archive January 1st 2025)
EM percentile statistics : 202505.v01 (Using data in the EMDB archive up until May 2025)
MapQ : 1.9.13
Ideal geometry (proteins) : Engh &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

1 Overall quality at a glance i

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

The reported resolution of this entry is 16.50 Å.

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



Metric	Whole archive (#Entries)	EM structures (#Entries)	Similar EM resolution (#Entries, resolution range(Å))
Clashscore	229148	23984	-
Ramachandran outliers	224038	23583	-
Sidechain outliers	223484	23102	-
RNA backbone	8273	3508	-
Q-score	-	25397	55 (16.00 - 17.00)

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	A	1530	16% 25% 60% 14% .
2	B	218	38% 21% 59% 18% .
3	C	206	45% 30% 54% 15% .

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Mol	Chain	Length	Quality of chain
4	D	205	
5	E	150	
6	F	100	
7	G	150	
8	H	129	
9	I	127	
10	J	98	
11	K	117	
12	L	123	
13	M	114	
14	N	100	
15	O	88	
16	P	82	
17	Q	80	
18	R	55	
19	S	79	
20	T	85	
21	U	51	
22	V	277	

2 Entry composition

There are 22 unique types of molecules in this entry. The entry contains 53533 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 RNA chain called 16S RIBOSOMAL RNA.

Mol	Chain	Residues	Atoms					AltConf	Trace
1	A	1530	Total	C	N	O	P	0	0
			32831	14642	6024	10635	1530		

- Molecule 2 is a protein called 30S RIBOSOMAL PROTEIN S2.

Mol	Chain	Residues	Atoms					AltConf	Trace
2	B	218	Total	C	N	O	S	0	0
			1704	1081	305	311	7		

- Molecule 3 is a protein called 30S RIBOSOMAL PROTEIN S3.

Mol	Chain	Residues	Atoms					AltConf	Trace
3	C	206	Total	C	N	O	S	0	0
			1624	1028	305	288	3		

- Molecule 4 is a protein called 30S RIBOSOMAL PROTEIN S4.

Mol	Chain	Residues	Atoms					AltConf	Trace
4	D	205	Total	C	N	O	S	0	0
			1643	1026	315	298	4		

- Molecule 5 is a protein called 30S RIBOSOMAL PROTEIN S5.

Mol	Chain	Residues	Atoms					AltConf	Trace
5	E	150	Total	C	N	O	S	0	0
			1105	687	211	201	6		

- Molecule 6 is a protein called 30S RIBOSOMAL PROTEIN S6.

Mol	Chain	Residues	Atoms					AltConf	Trace
6	F	100	Total	C	N	O	S	0	0
			817	515	148	148	6		

- Molecule 7 is a protein called 30S RIBOSOMAL PROTEIN S7.

Mol	Chain	Residues	Atoms					AltConf	Trace
7	G	150	Total	C	N	O	S	0	0
			1174	730	226	214	4		

- Molecule 8 is a protein called 30S RIBOSOMAL PROTEIN S8.

Mol	Chain	Residues	Atoms					AltConf	Trace
8	H	129	Total	C	N	O	S	0	0
			979	616	173	184	6		

- Molecule 9 is a protein called 30S RIBOSOMAL PROTEIN S9.

Mol	Chain	Residues	Atoms					AltConf	Trace
9	I	127	Total	C	N	O	S	0	0
			1022	634	206	179	3		

- Molecule 10 is a protein called 30S RIBOSOMAL PROTEIN S10.

Mol	Chain	Residues	Atoms					AltConf	Trace
10	J	98	Total	C	N	O	S	0	0
			786	493	150	142	1		

- Molecule 11 is a protein called 30S RIBOSOMAL PROTEIN S11.

Mol	Chain	Residues	Atoms					AltConf	Trace
11	K	117	Total	C	N	O	S	0	0
			877	540	174	160	3		

- Molecule 12 is a protein called 30S RIBOSOMAL PROTEIN S12.

Mol	Chain	Residues	Atoms					AltConf	Trace
12	L	123	Total	C	N	O	S	0	0
			955	590	196	165	4		

- Molecule 13 is a protein called 30S RIBOSOMAL PROTEIN S13.

Mol	Chain	Residues	Atoms					AltConf	Trace
13	M	114	Total	C	N	O	S	0	0
			883	546	178	156	3		

- Molecule 14 is a protein called 30S RIBOSOMAL PROTEIN S14.

Mol	Chain	Residues	Atoms					AltConf	Trace
14	N	96	Total	C	N	O	S	0	0
			774	483	160	128	3		

- Molecule 15 is a protein called 30S RIBOSOMAL PROTEIN S15.

Mol	Chain	Residues	Atoms					AltConf	Trace
15	O	88	Total	C	N	O	S	0	0
			716	440	146	129	1		

- Molecule 16 is a protein called 30S RIBOSOMAL PROTEIN S16.

Mol	Chain	Residues	Atoms					AltConf	Trace
16	P	82	Total	C	N	O	S	0	0
			649	406	128	114	1		

- Molecule 17 is a protein called 30S RIBOSOMAL PROTEIN S17.

Mol	Chain	Residues	Atoms					AltConf	Trace
17	Q	80	Total	C	N	O	S	0	0
			648	411	121	113	3		

- Molecule 18 is a protein called 30S RIBOSOMAL PROTEIN S18.

Mol	Chain	Residues	Atoms				AltConf	Trace
18	R	55	Total	C	N	O	0	0
			455	288	86	81		

- Molecule 19 is a protein called 30S RIBOSOMAL PROTEIN S19.

Mol	Chain	Residues	Atoms					AltConf	Trace
19	S	79	Total	C	N	O	S	0	0
			637	408	120	107	2		

- Molecule 20 is a protein called 30S RIBOSOMAL PROTEIN S20.

Mol	Chain	Residues	Atoms					AltConf	Trace
20	T	85	Total	C	N	O	S	0	0
			665	411	137	114	3		

- Molecule 21 is a protein called 30S RIBOSOMAL PROTEIN S21.

Mol	Chain	Residues	Atoms					AltConf	Trace
21	U	51	Total	C	N	O	S	0	0
			425	265	86	73	1		

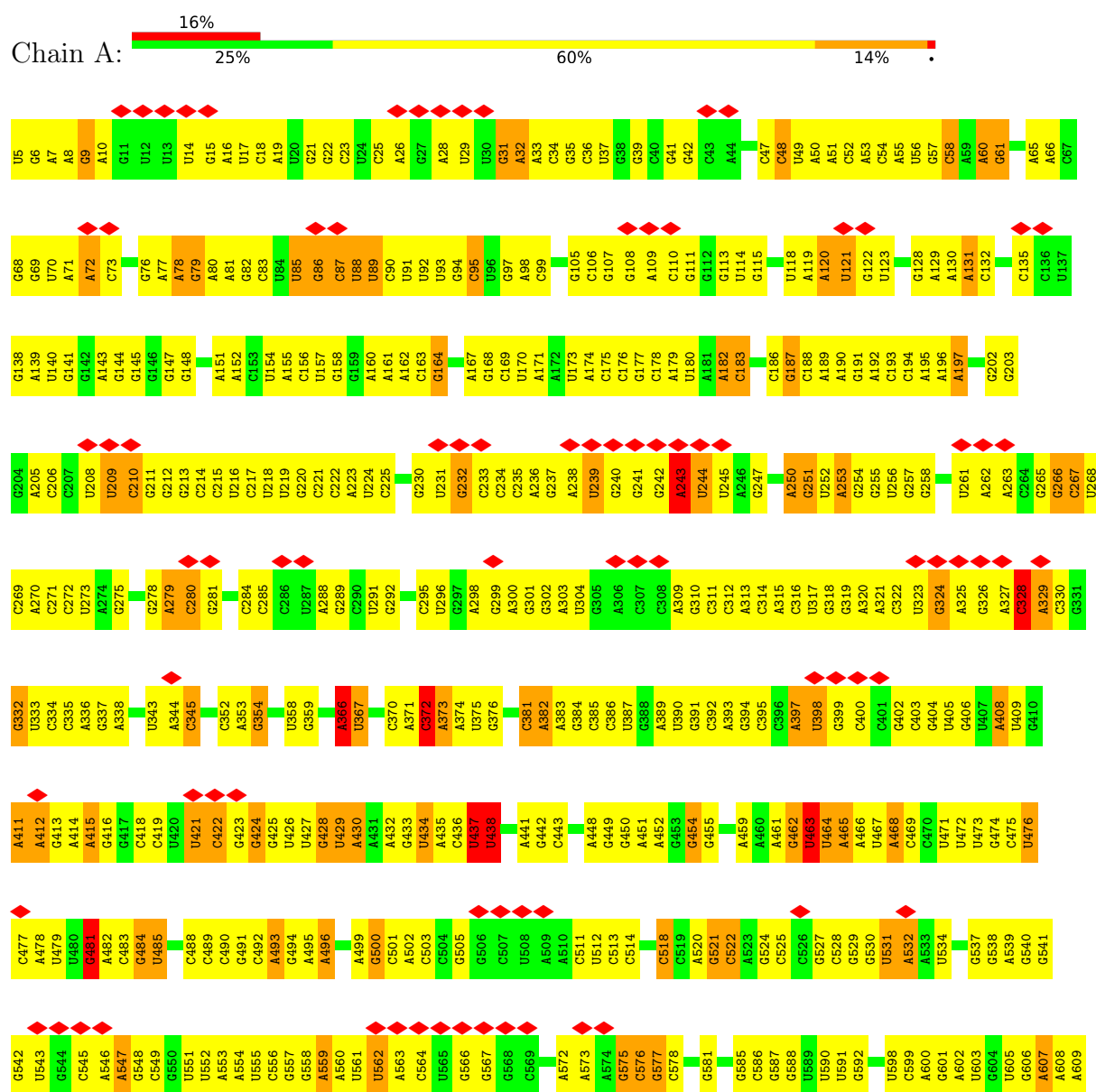
- Molecule 22 is a protein called PUTATIVE RIBOSOME BIOGENESIS GTPASE RSGA.

Mol	Chain	Residues	Atoms					AltConf	Trace
22	V	277	Total	C	N	O	S	0	0
			2164	1364	382	408	10		

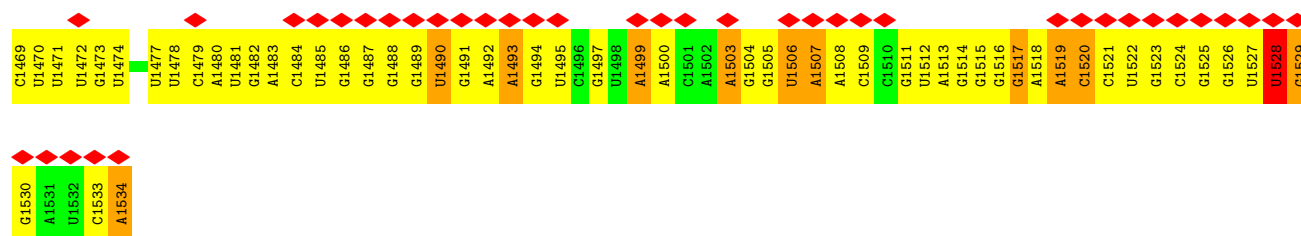
3 Residue-property plots

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.

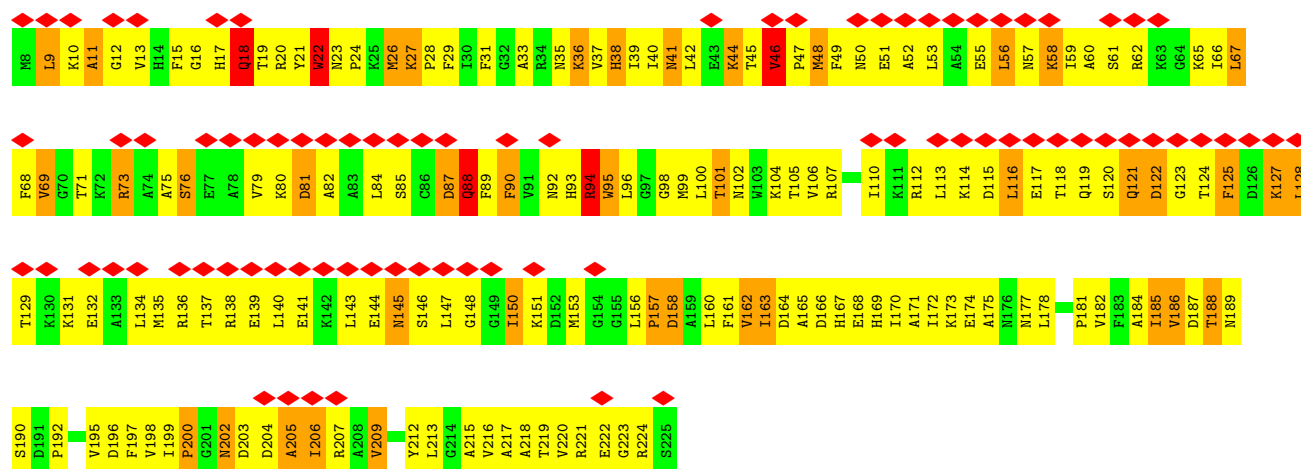
• Molecule 1: 16S RIBOSOMAL RNA



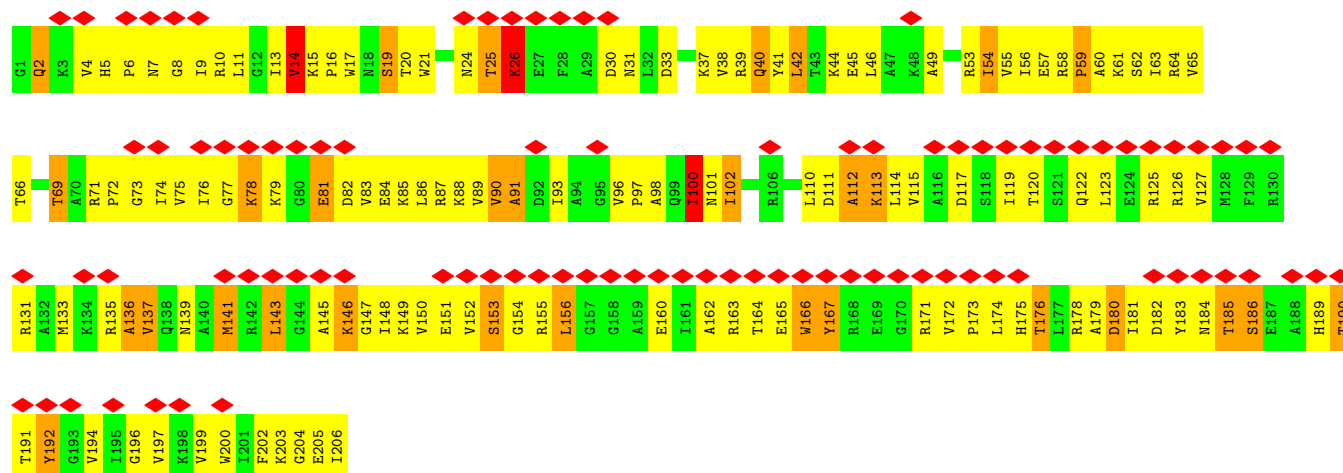
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G1347	U1348	A1349	A1350	C1351	C1352	A1353	G1354	G1355	G1356	G1357	U1358	U1359	G1360	A1361	C1362	A1363	G1364	G1365	A1366	C1367	A1368	A1369	G1370	G1371	U1372	A1373	A1374	A1375	U1376	A1377	G1378	G1379	U1380	U1381	C1382	C1383	U1384	C1385	G1386	G1387	C1388	C1389	U1390	U1391	C1392	U1393	A1394	C1395	G1396	C1397	A1398	A1399	C1400	G1401	C1402	C1403	C1404	G1405	U1406	C1407			
A1287	A1288	G1289	G1290	U1291	G1292	C1293	G1294	U1295	C1296	G1297	U1298	A1299	G1300	U1301	C1302	C1303	G1304	G1305	A1306	U1307	U1308	G1309	G1310	A1311	G1312	U1313	C1314	U1315	G1316	A1317	A1318	A1319	C1320	U1321	C1322	G1323	A1324	C1325	U1326	C1327	C1328	A1329	U1330	G1331	A1332	A1333	G1334	U1335	G1336	G1337	G1338	A1339	A1340	U1341	C1342	G1343	C1344	U1345	A1346				
G1221	G1222	C1223	U1224	A1225	U1226	A1227	C1228	A1229	C1230	G1231	U1232	G1233	C1234	U1235	A1236	C1237	G1238	A1239	U1240	G1241	G1242	C1243	G1244	C1245	A1246	U1247	A1248	C1249	A1250	A1251	U1252	G1253	A1254	G1255	A1256	A1257	G1258	C1259	U1260	A1261	C1262	C1263	A1269	A1270	A1271	G1272	C1273	A1274	G1278	G1279	A1280	C1281	U1282	U1283	C1284	A1285	U1286						
C1149	A1150	A1157	C1158	U1159	G1160	C1161	C1162	A1163	G1164	U1168	A1169	U1170	C1171	A1172	C1173	U1174	G1178	A1179	A1180	G1181	G1182	U1183	G1186	G1187	A1188	U1189	G1190	G1193	U1194	C1195	A1196	A1197	G1198	U1199	C1200	A1201	U1202	C1203	G1206	G1207	C1208	C1209	C1210	U1211	U1212	A1213	G1214	G1215	A1216	C1217	A1218	U1219	G1220										
A1080	A1081	A1082	U1083	G1084	U1085	U1086	U1090	U1091	A1092	A1093	G1094	U1095	C1096	C1097	A1098	G1099	C1100	A1101	A1102	C1103	G1108	C1109	A1110	A1111	C1112	C1113	U1114	U1115	U1116	A1117	U1118	U1121	U1122	U1123	G1124	U1125	U1126	C1127	C1128	C1129	A1130	G1131	C1132	G1133	G1134	U1135	C1136	C1137	G1138	G1139	C1140	C1141	G1142	G1143	U1148								
G1020	A1021	A1022	U1023	G1024	U1025	G1026	C1027	C1028	U1029	A1030	C1031	G1032	G1033	G1034	A1035	C1036	C1037	G1038	U1039	G1040	G1041	A1042	A1043	A1044	C1045	A1046	G1047	U1048	U1049	U1050	C1051	U1052	G1053	C1054	A1055	U1056	G1057	U1058	C1059	U1060	U1062	C1063	G1064	U1065	C1066	A1067	G1068	C1069	U1070	C1071	G1072	U1073	A1074	U1075	U1076	G1077	U1078	G1079					
A958	A959	U960	U961	U965	G966	G967	G968	A969	A970	C970	G971	C972	G973	A974	A975	G976	A977	C979	C980	U981	U982	A983	C984	C985	U986	U987	G988	U989	C990	U991	U992	G993	A994	C995	A996	U997	C998	C999	A1000	C1001	G1002	A1003	A1004	A1005	G1006	U1007	U1008	U1009	U1010	C1011	A1012	G1013	A1014	U1015	A1016	U1017	G1018	A1019					
U891	A892	C893	G894	G895	C896	C897	A900	A901	A902	A906	A907	A908	A909	C910	U911	C912	A913	A914	A915	U916	G917	A918	A919	U920	U921	G922	A923	C924	G925	G926	U927	G928	C929	G930	C931	C934	A935	C936	G939	C940	G941	G945	A946	G947	C948	A949	U950	G951	U952	G953	G954	U955	U956	U957									
A819	U820	G821	U822	C823	G824	A825	U828	G829	G830	A831	G832	G833	U834	U835	G836	U837	C838	C839	C840	C841	U842	U843	G844	A845	G846	G847	C848	U855	C856	G857	G858	G859	A860	G861	U862	U863	A864	A865	C866	G869	U875	A878	C879	C880	G881	C882	C883	U884	G885	G886	G887	G888	A889	G890									
A746	A747	G748	U751	G752	G755	G761	U762	G763	C764	G765	A766	A767	A768	G769	C770	G771	A777	G778	C779	A781	A782	C783	A784	G785	G786	G787	U788	A792	U793	A794	C795	C796	C797	U801	A802	G803	U804	C806	A807	C808	G809	C810	C811	G812	U813	A814	A815	A816	C817	G818													
A676	U677	U678	C679	C680	A681	G682	G683	U684	G685	U686	A687	G691	U692	G693	A694	U697	G698	C699	G700	G703	A704	C708	U709	G710	G711	A712	G713	U714	G715	A716	C719	C720	G721	G722	U723	G724	G725	C726	G730	A731	C732	G733	U734	C735	G736	C737	C738	C739	U740	A741	G742	A743											
U610	C611	C612	C613	C614	U619	C620	A621	A622	C623	C624	U625	G626	G627	G628	A629	A630	C631	U632	G633	C634	A635	U636	C637	U638	G639	A640	U641	A642	C643	U644	G645	A647	G648	A649	G650	C651	U652	U653	U657	C658	U659	C660	G661	U662	A663	G664	A665	G666	G669	G670	G671	A672	A673	G674	A675								



• Molecule 2: 30S RIBOSOMAL PROTEIN S2

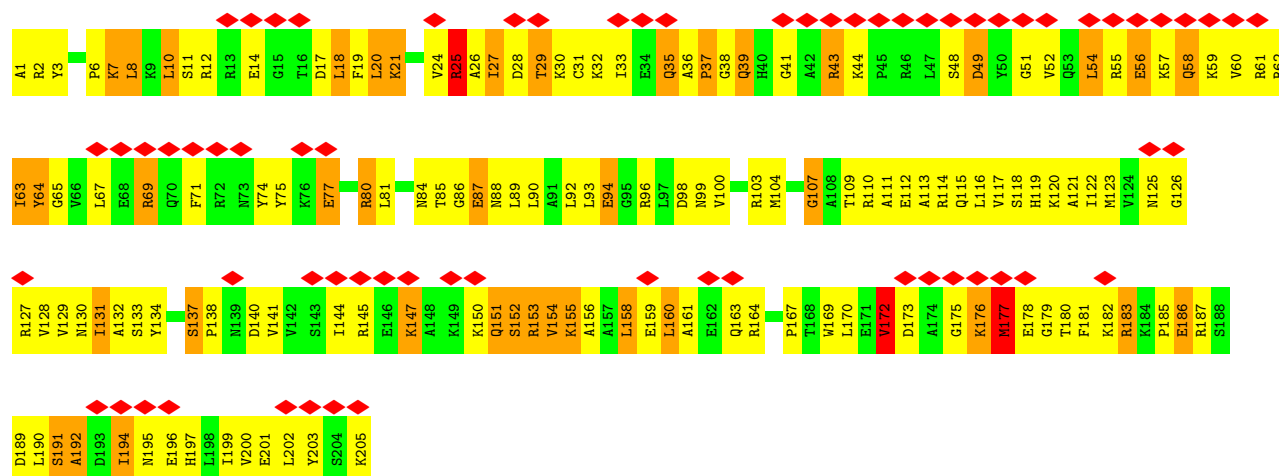


• Molecule 3: 30S RIBOSOMAL PROTEIN S3

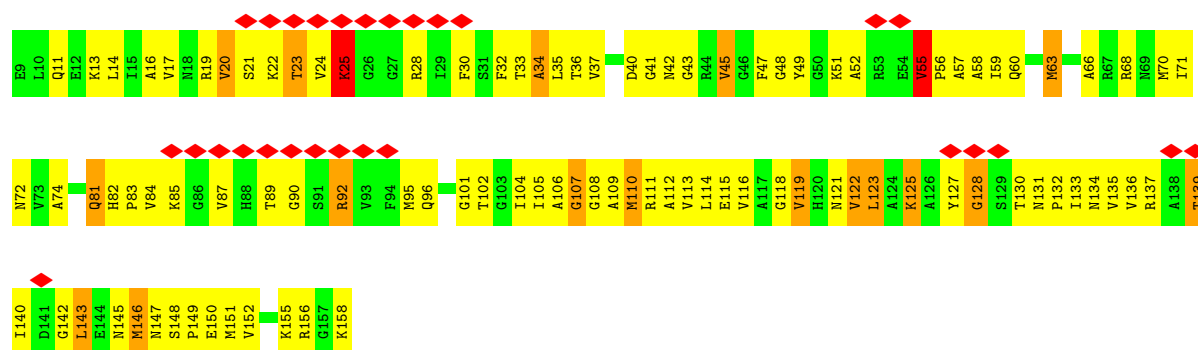


• Molecule 4: 30S RIBOSOMAL PROTEIN S4

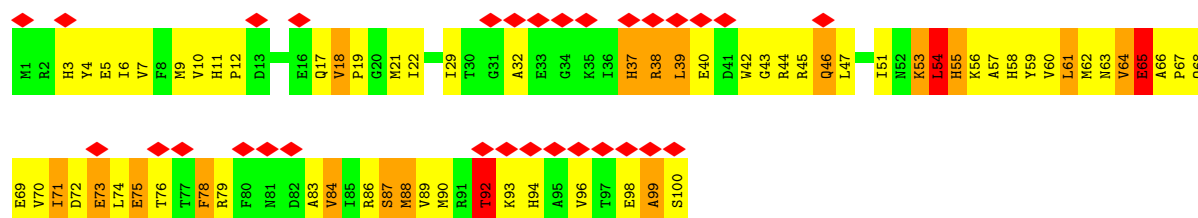




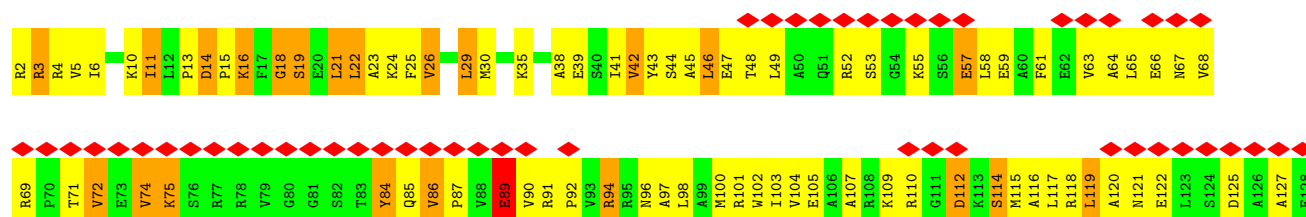
• Molecule 5: 30S RIBOSOMAL PROTEIN S5



• Molecule 6: 30S RIBOSOMAL PROTEIN S6

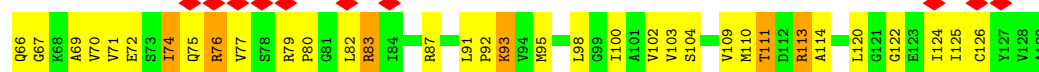
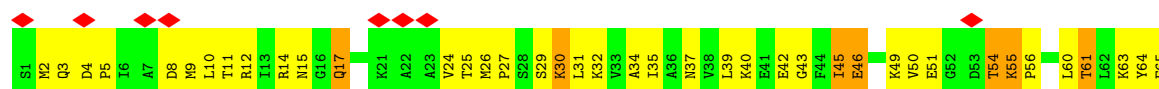
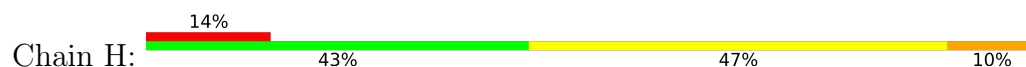


• Molecule 7: 30S RIBOSOMAL PROTEIN S7

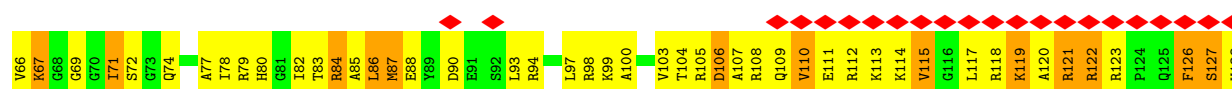
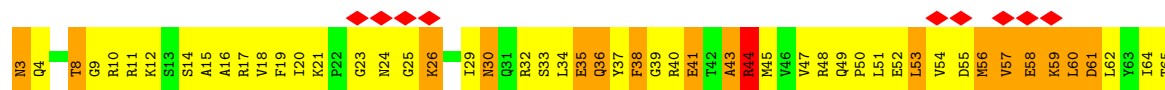




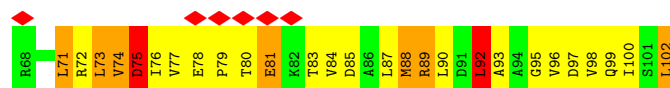
• Molecule 8: 30S RIBOSOMAL PROTEIN S8



• Molecule 9: 30S RIBOSOMAL PROTEIN S9

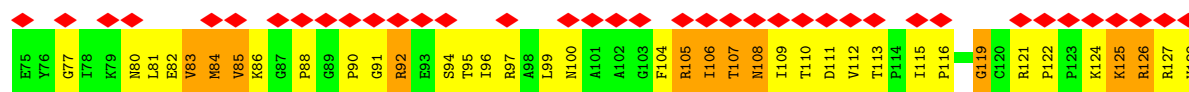


• Molecule 10: 30S RIBOSOMAL PROTEIN S10

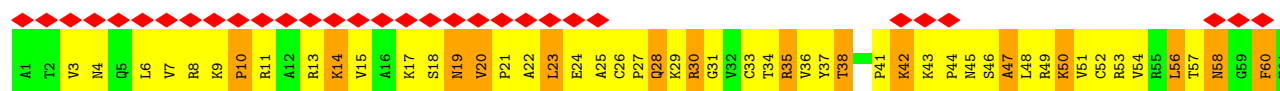


• Molecule 11: 30S RIBOSOMAL PROTEIN S11





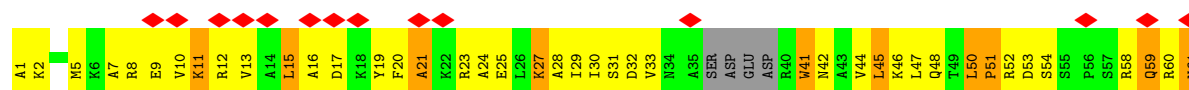
• Molecule 12: 30S RIBOSOMAL PROTEIN S12



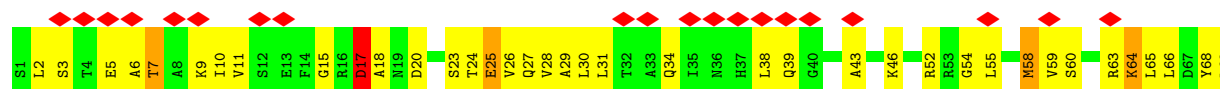
• Molecule 13: 30S RIBOSOMAL PROTEIN S13



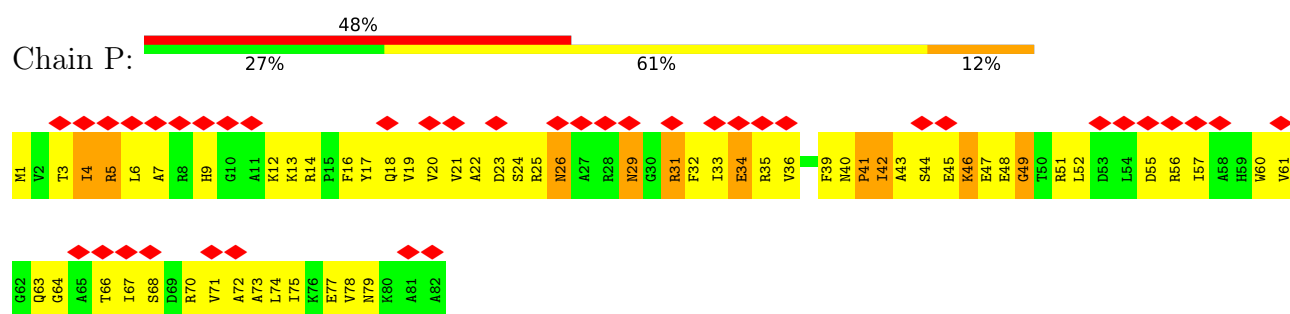
• Molecule 14: 30S RIBOSOMAL PROTEIN S14



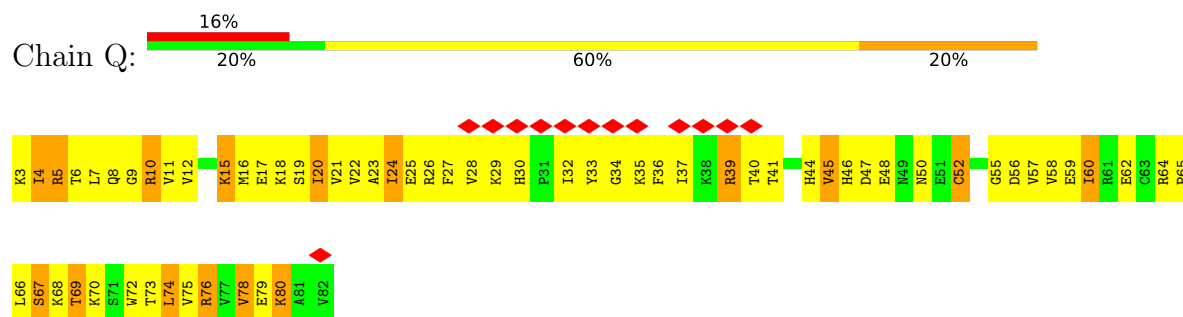
• Molecule 15: 30S RIBOSOMAL PROTEIN S15



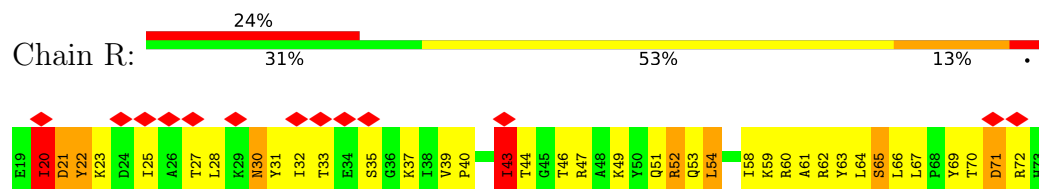
• Molecule 16: 30S RIBOSOMAL PROTEIN S16



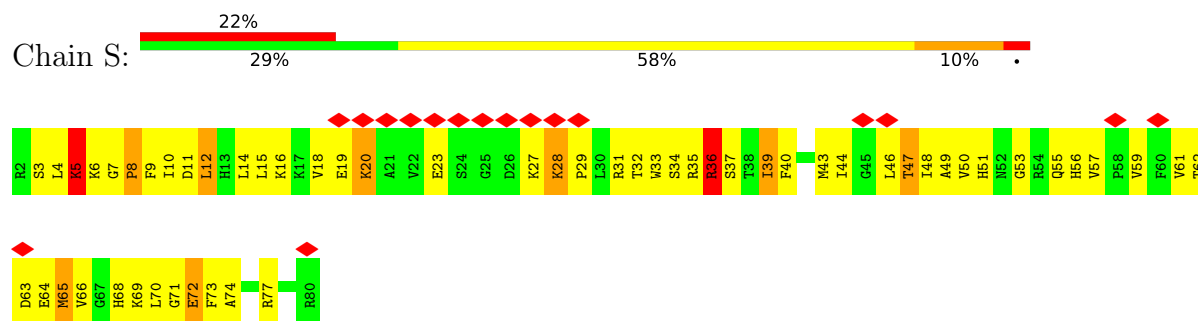
• Molecule 17: 30S RIBOSOMAL PROTEIN S17



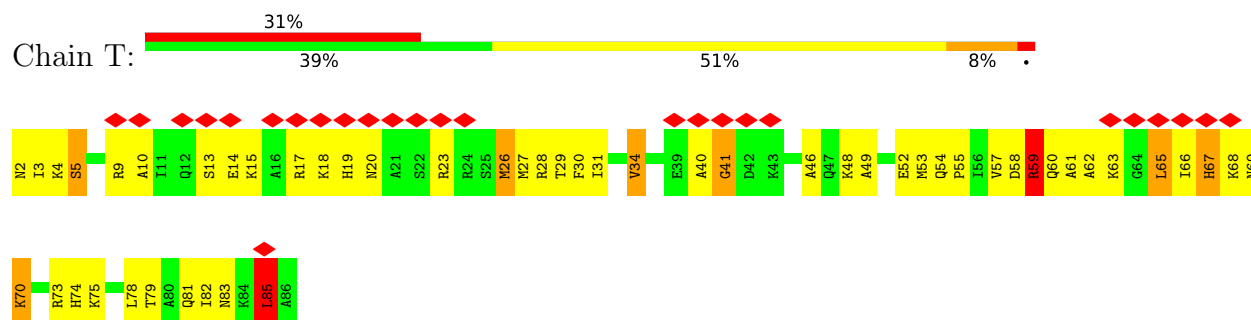
• Molecule 18: 30S RIBOSOMAL PROTEIN S18



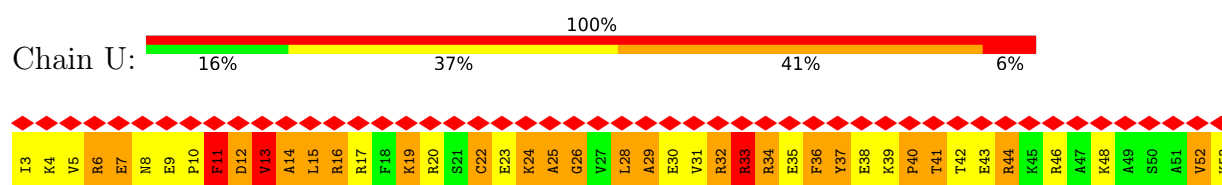
• Molecule 19: 30S RIBOSOMAL PROTEIN S19



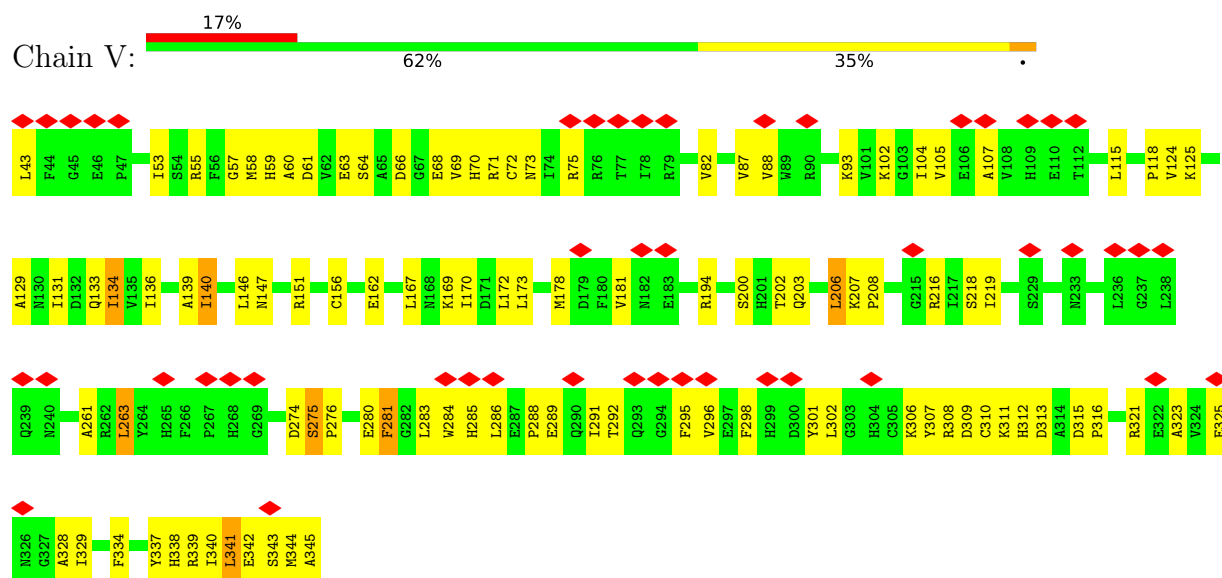
• Molecule 20: 30S RIBOSOMAL PROTEIN S20



• Molecule 21: 30S RIBOSOMAL PROTEIN S21



• Molecule 22: PUTATIVE RIBOSOME BIOGENESIS GTPASE RSGA



4 Experimental information

Property	Value	Source
EM reconstruction method	SINGLE PARTICLE	Depositor
Imposed symmetry	POINT, C1	Depositor
Number of particles used	16228	Depositor
Resolution determination method	Not provided	
CTF correction method	EACH MICROGRAPH	Depositor
Microscope	JEOL 2010F	Depositor
Voltage (kV)	200	Depositor
Electron dose ($e^-/\text{\AA}^2$)	15	Depositor
Minimum defocus (nm)	650	Depositor
Maximum defocus (nm)	3900	Depositor
Magnification	50000	Depositor
Image detector	KODAK SO-163 FILM	Depositor
Maximum map value	0.062	Depositor
Minimum map value	-0.022	Depositor
Average map value	0.000	Depositor
Map value standard deviation	0.004	Depositor
Recommended contour level	0.0088	Depositor
Map size ($\text{\AA}$)	325.12, 325.12, 325.12	wwPDB
Map dimensions	128, 128, 128	wwPDB
Map angles ($^\circ$)	90, 90, 90	wwPDB
Pixel spacing ($\text{\AA}$)	2.54, 2.54, 2.54	Depositor

5 Model quality

5.1 Standard geometry

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

Mol	Chain	Bond lengths		Bond angles	
		RMSZ	# Z >5	RMSZ	# Z >5
1	A	0.27	1/36762 (0.0%)	0.63	27/57350 (0.0%)
2	B	0.31	1/1735 (0.1%)	0.77	1/2338 (0.0%)
3	C	0.28	0/1651	0.74	1/2225 (0.0%)
4	D	0.27	0/1665	0.72	1/2227 (0.0%)
5	E	0.30	0/1118	0.76	1/1504 (0.1%)
6	F	0.30	0/835	0.81	2/1128 (0.2%)
7	G	0.27	0/1187	0.75	0/1591
8	H	0.28	0/989	0.79	1/1326 (0.1%)
9	I	0.28	0/1034	0.75	0/1375
10	J	0.29	0/796	0.71	0/1077
11	K	0.29	0/893	0.80	2/1205 (0.2%)
12	L	0.28	0/969	0.72	0/1300
13	M	0.28	0/892	0.77	0/1193
14	N	0.28	0/785	0.74	0/1043
15	O	0.26	0/724	0.73	1/966 (0.1%)
16	P	0.29	0/659	0.76	0/884
17	Q	0.26	0/657	0.63	0/881
18	R	0.31	0/462	0.71	0/621
19	S	0.29	0/652	0.72	0/877
20	T	0.28	0/671	0.79	1/888 (0.1%)
21	U	1.53	5/430 (1.2%)	1.15	8/570 (1.4%)
22	V	0.36	0/2200	0.86	1/2974 (0.0%)
All	All	0.31	7/57766 (0.0%)	0.68	47/85543 (0.1%)

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

Mol	Chain	#Chirality outliers	#Planarity outliers
1	A	0	16
21	U	0	1
All	All	0	17

All (7) bond length outliers are listed below:

Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
21	U	15	LEU	C-N	-23.62	0.99	1.33
21	U	25	ALA	C-N	-11.79	1.16	1.33
21	U	29	ALA	C-N	11.67	1.49	1.33
21	U	11	PHE	C-N	-9.16	1.21	1.33
21	U	7	GLU	C-N	6.79	1.43	1.33
1	A	463	U	O3'-P	-5.33	1.53	1.61
2	B	46	VAL	CA-CB	5.33	1.56	1.54

All (47) bond angle outliers are listed below:

Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
1	A	328	C	C2'-C3'-O3'	8.63	122.44	109.50
1	A	372	C	C2'-C3'-O3'	7.34	120.51	109.50
21	U	15	LEU	CA-C-N	6.79	136.11	121.64
21	U	15	LEU	C-N-CA	6.79	136.11	121.64
1	A	120	A	C4'-C3'-O3'	-6.61	103.08	113.00
1	A	232	G	C5'-C4'-C3'	-6.61	106.09	116.00
1	A	1528	U	C2'-C3'-O3'	6.57	119.36	109.50
1	A	968	A	C4'-C3'-O3'	-6.50	99.66	109.40
1	A	960	U	C2'-C3'-O3'	6.46	119.20	109.50
2	B	44	LYS	N-CA-C	-6.36	105.16	113.17
1	A	243	A	C4'-C3'-O3'	6.31	118.86	109.40
21	U	29	ALA	CA-C-N	-6.27	111.88	120.28
21	U	29	ALA	C-N-CA	-6.27	111.88	120.28
1	A	811	C	C4'-C3'-O3'	-6.26	103.61	113.00
1	A	765	G	C4'-C3'-O3'	6.08	122.11	113.00
1	A	765	G	N9-C1'-C2'	-6.04	102.94	112.00
1	A	438	U	N1-C1'-C2'	-5.96	105.06	114.00
1	A	1065	U	C2'-C3'-O3'	5.91	118.36	109.50
3	C	137	VAL	N-CA-C	-5.82	108.19	113.71
20	T	65	LEU	CB-CA-C	-5.78	109.89	116.54
21	U	19	LYS	N-CA-C	-5.71	106.28	113.18
1	A	1534	A	C4'-C3'-O3'	5.64	121.46	113.00
1	A	1432	G	C4'-C3'-O3'	-5.63	104.55	113.00
22	V	296	VAL	N-CA-C	5.63	117.04	110.62
1	A	437	U	N1-C1'-C2'	5.60	120.41	112.00
1	A	1004	A	C4'-C3'-O3'	-5.60	104.60	113.00
1	A	1118	U	C5'-C4'-C3'	-5.59	107.61	116.00
4	D	49	ASP	N-CA-C	-5.55	107.52	114.56
6	F	99	ALA	CB-CA-C	-5.44	110.32	116.63
15	O	43	ALA	N-CA-C	-5.34	105.57	111.71
1	A	1049	U	C2'-C3'-O3'	5.33	117.49	109.50

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
1	A	814	A	C5'-C4'-C3'	5.31	123.96	116.00
6	F	18	VAL	CB-CA-C	-5.31	108.66	113.70
1	A	965	U	C4'-C3'-O3'	-5.29	105.06	113.00
1	A	1528	U	C4'-C3'-O3'	5.25	117.27	109.40
5	E	55	VAL	CB-CA-C	-5.24	108.72	113.70
21	U	29	ALA	N-CA-C	-5.24	105.99	112.38
1	A	1319	A	N9-C1'-C2'	5.22	119.83	112.00
1	A	1043	G	N9-C1'-C2'	5.20	119.80	112.00
1	A	1226	C	C2'-C3'-O3'	5.19	117.28	109.50
21	U	46	ARG	N-CA-C	-5.18	107.02	113.55
11	K	105	ARG	CA-C-O	5.16	120.93	117.94
11	K	105	ARG	CB-CA-C	-5.16	109.64	117.07
1	A	855	U	C5'-C4'-C3'	-5.15	108.27	116.00
1	A	366	A	C2'-C3'-O3'	5.14	117.21	109.50
8	H	54	THR	N-CA-C	-5.02	108.90	114.62
21	U	15	LEU	O-C-N	-5.01	116.61	121.87

There are no chirality outliers.

All (17) planarity outliers are listed below:

Mol	Chain	Res	Type	Group
1	A	1006	G	Sidechain
1	A	1028	C	Sidechain
1	A	1319	A	Sidechain
1	A	1331	G	Sidechain
1	A	1441	A	Sidechain
1	A	187	G	Sidechain
1	A	281	G	Sidechain
1	A	437	U	Sidechain
1	A	438	U	Sidechain
1	A	454	G	Sidechain
1	A	481	G	Sidechain
1	A	496	A	Sidechain
1	A	521	G	Sidechain
1	A	575	G	Sidechain
1	A	58	C	Sidechain
1	A	703	G	Sidechain
21	U	11	PHE	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	A	32831	0	16507	1531	0
2	B	1704	0	1732	205	0
3	C	1624	0	1699	170	0
4	D	1643	0	1710	177	0
5	E	1105	0	1148	113	0
6	F	817	0	808	81	0
7	G	1174	0	1230	105	0
8	H	979	0	1034	77	0
9	I	1022	0	1070	150	0
10	J	786	0	828	98	0
11	K	877	0	887	113	0
12	L	955	0	1019	106	0
13	M	883	0	944	88	0
14	N	774	0	827	101	0
15	O	716	0	742	53	0
16	P	649	0	666	81	0
17	Q	648	0	691	83	0
18	R	455	0	478	45	0
19	S	637	0	665	82	0
20	T	665	0	714	56	0
21	U	425	0	448	104	0
22	V	2164	0	2131	451	0
All	All	53533	0	37978	3443	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 38.

All (3443) 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:A:1415:G:H21	22:V:285:HIS:CE1	1.06	1.69
1:A:1418:A:C6	22:V:312:HIS:CD2	1.77	1.68
1:A:1485:U:H4'	22:V:286:LEU:CD1	1.22	1.61
1:A:1485:U:C4'	22:V:286:LEU:HD12	1.22	1.59
1:A:1418:A:C6	22:V:312:HIS:HD2	0.88	1.54

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:1418:A:C2	22:V:292:THR:CG2	1.91	1.53
1:A:1483:A:C1'	22:V:298:PHE:CE2	1.94	1.49
1:A:1415:G:N2	22:V:285:HIS:CE1	1.83	1.45
1:A:1419:G:C5'	22:V:309:ASP:HA	1.17	1.43
1:A:1483:A:H1'	22:V:298:PHE:CE2	1.48	1.42
1:A:1418:A:N3	22:V:292:THR:CG2	1.80	1.39
1:A:1415:G:C2'	22:V:345:ALA:HB1	1.14	1.38
21:U:14:ALA:HA	21:U:16:ARG:NH1	1.37	1.38
1:A:1407:C:H5''	22:V:284:TRP:C	1.46	1.37
1:A:1517:G:C3'	22:V:124:VAL:HG22	1.38	1.36
1:A:1483:A:C1'	22:V:298:PHE:HE2	1.30	1.36
1:A:1418:A:C2	22:V:292:THR:HG23	1.51	1.35
21:U:14:ALA:O	21:U:16:ARG:NE	1.58	1.35
1:A:1415:G:H2'	22:V:345:ALA:CB	1.04	1.34
1:A:1486:G:H4'	22:V:285:HIS:N	1.43	1.32
1:A:1419:G:H5'	22:V:309:ASP:CA	1.57	1.32
1:A:1416:G:C8	22:V:345:ALA:C	2.07	1.32
1:A:1419:G:C5'	22:V:309:ASP:CA	2.08	1.31
1:A:1407:C:C4'	22:V:284:TRP:HA	1.61	1.30
21:U:14:ALA:CA	21:U:16:ARG:NH1	1.92	1.30
1:A:1416:G:H22	22:V:291:ILE:CG1	1.48	1.27
1:A:785:G:N7	22:V:71:ARG:CZ	2.00	1.25
1:A:1418:A:N6	22:V:337:TYR:OH	1.69	1.25
1:A:1420:U:C6	22:V:310:CYS:HA	1.71	1.24
1:A:784:A:H4'	22:V:72:CYS:N	1.53	1.24
1:A:1483:A:O4'	22:V:298:PHE:HE2	1.18	1.24
1:A:1418:A:N1	22:V:312:HIS:CD2	2.03	1.24
1:A:1420:U:OP2	22:V:309:ASP:OD1	1.57	1.23
1:A:1485:U:O4'	22:V:291:ILE:CD1	1.86	1.22
1:A:1418:A:N3	22:V:292:THR:HG21	0.91	1.22
1:A:1418:A:C5	22:V:312:HIS:CD2	2.26	1.21
21:U:14:ALA:N	21:U:16:ARG:NH1	1.88	1.21
1:A:1517:G:O3'	22:V:124:VAL:HG22	1.33	1.20
1:A:1483:A:O4'	22:V:298:PHE:CE2	1.91	1.20
1:A:1416:G:C8	22:V:345:ALA:HB3	1.76	1.20
1:A:1418:A:C2	22:V:292:THR:HG21	1.61	1.20
1:A:1482:G:H5''	22:V:338:HIS:CD2	1.61	1.19
1:A:1420:U:H6	22:V:310:CYS:CA	1.55	1.18
1:A:1420:U:OP2	22:V:309:ASP:CG	1.88	1.17
1:A:793:U:O4	22:V:125:LYS:CB	1.82	1.17
1:A:1485:U:C1'	22:V:291:ILE:HD11	1.75	1.17

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:1416:G:C8	22:V:345:ALA:CA	2.28	1.15
1:A:784:A:O3'	22:V:60:ALA:O	1.62	1.15
1:A:1482:G:C5'	22:V:338:HIS:CD2	2.09	1.15
1:A:1418:A:C4	22:V:292:THR:HG21	1.82	1.15
21:U:14:ALA:CA	21:U:16:ARG:CZ	2.24	1.15
1:A:1407:C:C5'	22:V:284:TRP:C	2.19	1.14
1:A:1486:G:C4'	22:V:285:HIS:N	2.09	1.14
1:A:1407:C:H4'	22:V:284:TRP:CA	1.78	1.13
1:A:1517:G:H5''	22:V:124:VAL:C	1.43	1.13
1:A:1420:U:C6	22:V:311:LYS:N	2.16	1.12
1:A:1484:C:H3'	22:V:340:ILE:HD13	1.32	1.11
1:A:1517:G:C5'	22:V:124:VAL:C	2.06	1.11
1:A:1483:A:H1'	22:V:298:PHE:CD2	1.84	1.11
1:A:1418:A:C6	22:V:337:TYR:OH	1.96	1.10
1:A:783:C:C5'	22:V:58:MET:HB3	1.82	1.10
1:A:785:G:H5'	22:V:69:VAL:HG12	1.33	1.10
22:V:82:VAL:HG11	22:V:129:ALA:HB2	1.31	1.10
1:A:1407:C:C4'	22:V:284:TRP:CA	2.30	1.09
2:B:163:ILE:HG23	2:B:164:ASP:H	1.17	1.09
1:A:1416:G:C8	22:V:345:ALA:CB	2.35	1.08
1:A:1415:G:N2	22:V:285:HIS:ND1	1.95	1.08
1:A:1416:G:N7	22:V:345:ALA:C	2.13	1.07
1:A:785:G:C5'	22:V:69:VAL:HG12	1.83	1.06
1:A:1407:C:H4'	22:V:284:TRP:HA	1.07	1.06
1:A:1416:G:O4'	22:V:345:ALA:HB3	1.55	1.06
1:A:1418:A:C5	22:V:312:HIS:HD2	1.67	1.06
1:A:785:G:H5'	22:V:69:VAL:CG1	1.83	1.06
1:A:1416:G:H22	22:V:291:ILE:HG13	0.91	1.06
1:A:1420:U:C5	22:V:311:LYS:HG2	1.90	1.06
7:G:125:ASP:HB3	7:G:130:LYS:HB3	1.39	1.05
1:A:1486:G:O4'	22:V:285:HIS:HA	1.55	1.05
3:C:78:LYS:HG2	3:C:81:GLU:HB2	1.36	1.05
1:A:1416:G:H8	22:V:345:ALA:O	1.40	1.05
1:A:1407:C:C3'	22:V:284:TRP:HA	1.87	1.04
1:A:1408:A:OP1	22:V:283:LEU:HB3	1.57	1.04
10:J:35:GLN:HB3	10:J:77:VAL:HG23	1.34	1.04
1:A:1416:G:C8	22:V:345:ALA:O	2.10	1.04
1:A:1420:U:H6	22:V:310:CYS:HA	0.90	1.04
8:H:92:PRO:HA	8:H:93:LYS:HZ2	1.24	1.03
1:A:793:U:O4	22:V:125:LYS:HB2	1.25	1.03
1:A:1408:A:H5'	22:V:151:ARG:HH22	1.25	1.02

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:1517:G:O3'	22:V:124:VAL:CG2	2.07	1.01
1:A:1407:C:O3'	22:V:284:TRP:CA	2.07	1.01
1:A:1486:G:C8	22:V:344:MET:SD	2.54	1.01
21:U:14:ALA:N	21:U:16:ARG:CZ	2.23	1.01
1:A:1416:G:N2	22:V:291:ILE:CG1	2.23	1.01
1:A:1483:A:N6	22:V:288:PRO:O	1.93	1.01
1:A:1418:A:N6	22:V:312:HIS:HD2	1.57	1.01
1:A:785:G:C4'	22:V:69:VAL:HG12	1.90	1.00
1:A:1517:G:C3'	22:V:124:VAL:CG2	2.32	1.00
1:A:1420:U:O5'	22:V:307:TYR:HB2	1.61	1.00
1:A:1416:G:H2'	22:V:288:PRO:HD3	1.45	0.99
1:A:1482:G:H2'	22:V:341:LEU:CD2	1.93	0.99
2:B:46:VAL:HG13	2:B:47:PRO:HD3	1.45	0.98
1:A:1416:G:N2	22:V:291:ILE:HG13	1.76	0.98
1:A:1485:U:H4'	22:V:286:LEU:HD11	1.44	0.98
1:A:376:G:H5''	16:P:5:ARG:HB2	1.45	0.98
8:H:103:VAL:HG12	8:H:124:ILE:HA	1.46	0.97
1:A:784:A:H5'	22:V:59:HIS:CD2	2.00	0.97
1:A:1420:U:C4	22:V:311:LYS:HG2	1.99	0.97
11:K:124:LYS:HA	21:U:34:ARG:HB3	1.47	0.96
14:N:63:CYS:HB3	14:N:67:GLY:H	1.30	0.96
1:A:1417:G:O3'	22:V:289:GLU:CD	2.07	0.96
1:A:1420:U:C5'	22:V:307:TYR:HB2	1.95	0.96
1:A:1408:A:C5'	22:V:151:ARG:NH2	2.27	0.96
21:U:29:ALA:HB1	21:U:32:ARG:HH21	1.26	0.96
19:S:49:ALA:HB1	19:S:56:HIS:HB3	1.47	0.96
3:C:58:ARG:HG2	3:C:63:ILE:HG22	1.48	0.96
1:A:1417:G:H4'	22:V:289:GLU:OE1	1.66	0.96
2:B:67:LEU:HD11	2:B:157:PRO:HB3	1.47	0.95
1:A:1313:U:H5''	19:S:5:LYS:HG2	1.47	0.95
1:A:1408:A:OP1	22:V:283:LEU:CB	2.14	0.95
1:A:1408:A:C5'	22:V:151:ARG:HH22	1.64	0.95
6:F:53:LYS:HD3	6:F:54:LEU:H	1.30	0.95
1:A:1417:G:O2'	22:V:289:GLU:N	2.00	0.94
1:A:1419:G:H5'	22:V:309:ASP:HA	0.97	0.94
13:M:11:HIS:H	13:M:44:ILE:HD11	1.32	0.94
13:M:106:ARG:HE	13:M:112:ARG:HG2	1.29	0.94
1:A:1485:U:H1'	22:V:291:ILE:HD11	1.46	0.94
1:A:1086:U:H3	1:A:1099:G:H22	1.14	0.94
1:A:784:A:H5''	22:V:60:ALA:H	1.33	0.93
1:A:1484:C:O5'	22:V:340:ILE:HD12	1.68	0.93

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:785:G:P	22:V:71:ARG:HA	1.91	0.93
1:A:1482:G:O5'	22:V:313:ASP:OD1	1.85	0.93
22:V:173:LEU:HD12	22:V:178:MET:HE2	1.49	0.93
9:I:51:LEU:HD23	9:I:56:MET:HE3	1.51	0.92
1:A:1416:G:H8	22:V:345:ALA:C	1.67	0.92
1:A:1484:C:H2'	22:V:291:ILE:HD12	1.46	0.92
1:A:1416:G:H1	22:V:341:LEU:HA	1.35	0.92
1:A:1420:U:P	22:V:307:TYR:HB3	2.08	0.92
1:A:1415:G:H21	22:V:285:HIS:HE1	0.96	0.92
1:A:1484:C:N3	22:V:341:LEU:CD1	2.33	0.91
21:U:14:ALA:HA	21:U:16:ARG:CZ	1.90	0.91
1:A:1485:U:O4'	22:V:291:ILE:HD12	1.69	0.91
17:Q:18:LYS:HG2	17:Q:48:GLU:HA	1.52	0.91
1:A:1482:G:C5'	22:V:313:ASP:OD1	2.19	0.90
1:A:1484:C:H3'	22:V:340:ILE:CD1	2.00	0.90
10:J:88:MET:HB2	10:J:89:ARG:HH12	1.32	0.90
1:A:1482:G:P	22:V:313:ASP:OD1	2.28	0.90
6:F:88:MET:HE1	6:F:90:MET:HG3	1.52	0.90
1:A:1485:U:OP2	22:V:340:ILE:HD13	1.72	0.90
2:B:150:ILE:HG13	2:B:153:MET:HE3	1.53	0.90
21:U:14:ALA:C	21:U:16:ARG:NE	2.30	0.90
1:A:1483:A:O4'	22:V:337:TYR:CD2	1.99	0.89
14:N:60:ARG:HH21	14:N:62:ARG:HE	1.21	0.89
1:A:1408:A:H5'	22:V:151:ARG:NH2	1.87	0.89
1:A:1485:U:C1'	22:V:291:ILE:CD1	2.46	0.89
1:A:1420:U:P	22:V:307:TYR:CB	2.59	0.89
11:K:22:ILE:HG21	11:K:95:THR:HG21	1.53	0.89
19:S:18:VAL:HG21	19:S:43:MET:HG2	1.53	0.89
1:A:785:G:C3'	22:V:69:VAL:HG12	2.02	0.89
20:T:68:LYS:HG3	20:T:69:ASN:H	1.36	0.89
1:A:1486:G:C5	22:V:344:MET:HA	2.07	0.89
1:A:1483:A:H61	22:V:288:PRO:C	1.80	0.88
9:I:34:LEU:HD11	9:I:47:VAL:HG21	1.56	0.88
1:A:784:A:H4'	22:V:71:ARG:C	1.98	0.88
1:A:784:A:C4'	22:V:72:CYS:N	2.35	0.88
1:A:1415:G:H2'	22:V:345:ALA:HB2	1.53	0.88
1:A:1484:C:N3	22:V:341:LEU:HD13	1.88	0.88
14:N:60:ARG:HD3	14:N:60:ARG:H	1.37	0.88
21:U:13:VAL:C	21:U:16:ARG:HH12	1.81	0.88
1:A:784:A:H5'	22:V:59:HIS:CG	1.76	0.88
19:S:4:LEU:HD13	19:S:9:PHE:H	1.38	0.88

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:B:131:LYS:HA	2:B:134:LEU:HD12	1.56	0.88
4:D:60:VAL:HA	4:D:63:ILE:HD12	1.57	0.87
22:V:82:VAL:CG1	22:V:129:ALA:HB2	2.04	0.87
12:L:8:ARG:HG3	12:L:9:LYS:H	1.40	0.87
13:M:78:ARG:HH21	13:M:79:LEU:HG	1.37	0.87
22:V:87:VAL:HG11	22:V:105:VAL:HG13	1.54	0.87
1:A:1420:U:C6	22:V:310:CYS:CA	2.44	0.86
21:U:13:VAL:HG13	21:U:14:ALA:H	1.38	0.86
1:A:699:C:H2'	1:A:700:G:H5''	1.57	0.86
1:A:1407:C:H5''	22:V:284:TRP:CA	2.06	0.86
6:F:53:LYS:HZ3	6:F:53:LYS:HA	1.41	0.86
1:A:785:G:H4'	22:V:70:HIS:CA	2.04	0.86
1:A:1517:G:N3	22:V:118:PRO:O	2.08	0.86
13:M:48:SER:H	13:M:51:GLN:HB2	1.38	0.86
1:A:1416:G:H8	22:V:345:ALA:CB	1.85	0.86
1:A:1418:A:N6	22:V:312:HIS:CD2	2.37	0.86
1:A:1482:G:C5'	22:V:338:HIS:NE2	1.67	0.86
9:I:29:ILE:HG22	9:I:64:ILE:HB	1.57	0.86
1:A:1517:G:N3	22:V:118:PRO:C	2.34	0.86
16:P:20:VAL:HG23	16:P:35:ARG:HA	1.57	0.86
1:A:1418:A:C2	22:V:312:HIS:CD2	2.64	0.86
1:A:785:G:C8	22:V:71:ARG:NE	2.43	0.85
1:A:1482:G:H2'	22:V:341:LEU:HD23	1.59	0.85
16:P:46:LYS:H	16:P:46:LYS:HD3	1.39	0.85
1:A:243:A:H4'	1:A:244:U:H5'	1.59	0.85
2:B:33:ALA:HA	2:B:38:HIS:HA	1.58	0.85
4:D:160:LEU:H	4:D:160:LEU:HD13	1.41	0.85
8:H:54:THR:HG23	8:H:55:LYS:HG2	1.59	0.85
21:U:14:ALA:O	21:U:16:ARG:CZ	2.25	0.85
1:A:1416:G:C8	22:V:345:ALA:N	2.43	0.84
1:A:1484:C:O2'	22:V:291:ILE:HA	1.77	0.84
3:C:146:LYS:HE3	3:C:202:PHE:HE2	1.42	0.84
7:G:112:ASP:HB2	7:G:118:ARG:HG2	1.55	0.84
17:Q:27:PHE:HB3	17:Q:36:PHE:HB3	1.59	0.84
2:B:163:ILE:HG23	2:B:164:ASP:N	1.92	0.84
1:A:1481:U:O3'	22:V:313:ASP:OD1	1.79	0.84
18:R:52:ARG:HB3	18:R:52:ARG:HH11	1.40	0.84
21:U:13:VAL:C	21:U:16:ARG:NH1	2.35	0.84
21:U:14:ALA:O	21:U:16:ARG:CD	2.25	0.84
1:A:1486:G:C5'	22:V:285:HIS:N	2.40	0.84
10:J:8:ILE:HD12	10:J:100:ILE:HG22	1.58	0.84

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:1483:A:C6	22:V:288:PRO:O	2.31	0.84
15:O:6:ALA:HA	15:O:9:LYS:HE3	1.58	0.83
1:A:1486:G:C4	22:V:344:MET:HA	2.14	0.83
11:K:126:ARG:HB2	21:U:33:ARG:HD2	1.61	0.83
12:L:48:LEU:H	12:L:48:LEU:HD23	1.42	0.83
1:A:783:C:H4'	22:V:58:MET:HB3	1.59	0.83
1:A:1407:C:O3'	22:V:284:TRP:C	2.20	0.83
13:M:33:LEU:HD22	13:M:38:ILE:HB	1.61	0.83
1:A:1101:A:H4'	1:A:1102:A:O5'	1.76	0.83
5:E:105:ILE:HB	5:E:123:LEU:HA	1.61	0.83
1:A:1407:C:O3'	22:V:284:TRP:HA	1.77	0.82
2:B:202:ASN:HD22	2:B:204:ASP:H	1.26	0.82
15:O:25:GLU:HG3	15:O:80:LEU:HD12	1.58	0.82
1:A:664:G:H22	1:A:741:G:H1	1.25	0.82
1:A:784:A:C4'	22:V:71:ARG:C	2.49	0.82
12:L:56:LEU:HD11	12:L:81:ILE:HD12	1.61	0.82
21:U:14:ALA:HA	21:U:16:ARG:HH11	1.03	0.82
1:A:1323:G:H2'	1:A:1324:A:C8	2.15	0.82
3:C:155:ARG:H	3:C:162:ALA:HA	1.45	0.82
6:F:64:VAL:HG12	6:F:65:GLU:H	1.45	0.82
2:B:61:SER:HA	2:B:224:ARG:HA	1.62	0.82
1:A:1415:G:N2	22:V:285:HIS:HE1	1.53	0.82
1:A:33:A:H1'	12:L:27:PRO:HG3	1.61	0.82
1:A:1417:G:O2'	22:V:288:PRO:C	2.17	0.81
7:G:21:LEU:HD23	7:G:21:LEU:H	1.45	0.81
1:A:785:G:O5'	22:V:71:ARG:HA	1.78	0.81
1:A:1414:U:H3	22:V:343:SER:HA	1.46	0.81
3:C:49:ALA:HB1	3:C:75:VAL:HG22	1.63	0.81
1:A:1418:A:P	22:V:289:GLU:OE2	2.38	0.81
1:A:783:C:C4'	22:V:58:MET:HB3	2.09	0.81
1:A:1409:C:N4	1:A:1491:G:H1	1.79	0.81
1:A:1482:G:H5'	22:V:313:ASP:OD1	1.79	0.81
4:D:36:ALA:HA	4:D:41:GLY:HA3	1.61	0.81
1:A:1407:C:C5'	22:V:284:TRP:CA	2.57	0.80
10:J:35:GLN:HB2	10:J:78:GLU:HB2	1.64	0.80
4:D:159:GLU:HG3	4:D:160:LEU:N	1.94	0.80
15:O:7:THR:HG22	15:O:30:LEU:HD11	1.62	0.80
1:A:1211:U:H4'	1:A:1213:A:H1'	1.64	0.80
1:A:1483:A:H5''	22:V:334:PHE:O	1.82	0.80
20:T:30:PHE:HB3	20:T:53:MET:HB3	1.63	0.80
1:A:785:G:H4'	22:V:70:HIS:N	1.96	0.80

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:785:G:C8	22:V:71:ARG:CZ	2.65	0.80
1:A:1419:G:H5'	22:V:309:ASP:CB	2.11	0.80
5:E:87:VAL:HG12	5:E:92:ARG:HA	1.64	0.80
1:A:1005:A:H2'	1:A:1006:G:O4'	1.82	0.79
1:A:1416:G:N9	22:V:345:ALA:HB3	1.97	0.79
1:A:1416:G:O4'	22:V:345:ALA:CB	2.28	0.79
1:A:1419:G:H4'	22:V:308:ARG:O	1.81	0.79
10:J:56:HIS:H	14:N:80:ARG:HH22	1.30	0.79
17:Q:16:MET:HB2	17:Q:19:SER:HB2	1.64	0.79
21:U:36:PHE:HB3	21:U:40:PRO:HD3	1.64	0.79
1:A:785:G:N7	22:V:71:ARG:NH2	2.29	0.79
9:I:12:LYS:H	9:I:109:GLN:HE22	1.30	0.79
17:Q:10:ARG:CZ	17:Q:11:VAL:H	1.95	0.79
1:A:1416:G:C1'	22:V:345:ALA:HB3	2.11	0.79
1:A:1483:A:H62	22:V:341:LEU:HD11	1.46	0.79
22:V:323:ALA:HB1	22:V:329:ILE:HD13	1.62	0.79
1:A:784:A:H5''	22:V:60:ALA:N	1.97	0.79
1:A:785:G:H4'	22:V:69:VAL:C	2.08	0.79
1:A:1485:U:O4'	22:V:286:LEU:HD12	1.81	0.78
5:E:83:PRO:HB3	5:E:96:GLN:HG3	1.65	0.78
9:I:23:GLY:H	9:I:60:LEU:HA	1.48	0.78
13:M:19:THR:HA	13:M:24:VAL:HG23	1.64	0.78
21:U:14:ALA:C	21:U:16:ARG:CZ	2.56	0.78
1:A:1485:U:O4'	22:V:291:ILE:HD11	1.61	0.78
1:A:1483:A:C5'	22:V:334:PHE:O	2.30	0.78
15:O:2:LEU:HD22	15:O:7:THR:HG23	1.65	0.78
17:Q:11:VAL:HA	17:Q:22:VAL:HG22	1.66	0.78
20:T:27:MET:HE1	20:T:66:ILE:HG12	1.66	0.78
1:A:1419:G:O2'	22:V:310:CYS:HB3	1.84	0.78
9:I:110:VAL:HG12	9:I:111:GLU:H	1.47	0.78
17:Q:10:ARG:NH1	17:Q:56:ASP:H	1.79	0.78
1:A:1482:G:H5'	22:V:313:ASP:HA	1.66	0.78
2:B:112:ARG:O	2:B:116:LEU:HB2	1.82	0.78
1:A:65:A:N3	1:A:65:A:H2'	1.99	0.77
1:A:785:G:O5'	22:V:71:ARG:CA	2.31	0.77
1:A:1308:U:H2'	1:A:1309:G:H8	1.50	0.77
3:C:184:ASN:HD22	3:C:185:THR:H	1.32	0.77
8:H:113:ARG:HH21	8:H:114:ALA:HA	1.49	0.77
13:M:11:HIS:N	13:M:44:ILE:HD11	1.99	0.77
14:N:30:ILE:HG21	14:N:41:TRP:HB2	1.65	0.77
1:A:1415:G:C3'	22:V:345:ALA:HB1	2.14	0.77

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
12:L:35:ARG:NH2	12:L:75:GLU:HB3	2.00	0.77
21:U:3:ILE:HA	21:U:19:LYS:HG2	1.64	0.77
1:A:1418:A:C4	22:V:312:HIS:NE2	2.52	0.77
1:A:1421:G:OP1	22:V:306:LYS:HD3	1.84	0.77
4:D:187:ARG:HA	4:D:190:LEU:HD22	1.65	0.77
6:F:38:ARG:HB3	6:F:63:ASN:HB2	1.65	0.77
7:G:87:PRO:HG2	7:G:151:ALA:HB2	1.66	0.77
1:A:78:A:H2'	1:A:79:G:C8	2.20	0.77
21:U:14:ALA:C	21:U:16:ARG:HD2	2.09	0.77
2:B:120:SER:HA	2:B:125:PHE:HB3	1.67	0.77
15:O:81:ILE:HA	15:O:86:LEU:HD12	1.67	0.77
1:A:1420:U:C6	22:V:310:CYS:C	2.63	0.77
2:B:9:LEU:HD22	2:B:11:ALA:H	1.48	0.77
16:P:42:ILE:HB	16:P:46:LYS:HD2	1.67	0.77
21:U:14:ALA:C	21:U:16:ARG:CD	2.58	0.77
1:A:1421:G:O4'	22:V:315:ASP:HA	1.86	0.76
4:D:104:MET:HE2	4:D:170:LEU:HD13	1.67	0.76
1:A:1420:U:O5'	22:V:307:TYR:CB	2.33	0.76
11:K:30:ILE:HG22	11:K:45:THR:HA	1.67	0.76
14:N:12:ARG:HA	14:N:15:LEU:HD11	1.66	0.76
1:A:437:U:H2'	1:A:438:U:O4'	1.85	0.76
1:A:1486:G:C8	22:V:344:MET:CE	2.68	0.76
2:B:162:VAL:HG13	2:B:184:ALA:HB2	1.67	0.76
1:A:1416:G:N2	22:V:291:ILE:HG12	1.99	0.76
1:A:1219:A:H2'	1:A:1220:G:C8	2.21	0.76
13:M:21:ILE:HG22	13:M:64:VAL:HG11	1.67	0.76
21:U:14:ALA:HA	21:U:16:ARG:CD	2.15	0.76
1:A:1416:G:C5	22:V:344:MET:HB2	2.21	0.76
22:V:203:GLN:HA	22:V:206:LEU:HD22	1.67	0.76
1:A:840:C:H2'	1:A:842:U:H5''	1.67	0.75
2:B:41:ASN:HD22	2:B:44:LYS:HE2	1.50	0.75
13:M:23:GLY:HA3	13:M:64:VAL:HG13	1.67	0.75
19:S:35:ARG:HG2	19:S:50:VAL:HG13	1.66	0.75
1:A:1417:G:O3'	22:V:289:GLU:OE2	2.05	0.75
3:C:8:GLY:HA2	3:C:11:LEU:HG	1.67	0.75
1:A:785:G:P	22:V:60:ALA:O	2.43	0.75
11:K:86:LYS:HB2	11:K:112:VAL:HG23	1.68	0.75
13:M:18:LEU:HB3	13:M:29:SER:HB2	1.67	0.75
15:O:70:LYS:HZ1	15:O:74:VAL:HG13	1.51	0.75
1:A:781:A:H2'	1:A:782:A:H5'	1.69	0.75
10:J:19:ASP:O	10:J:22:THR:HG22	1.87	0.75

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
22:V:87:VAL:CG1	22:V:105:VAL:HG13	2.16	0.75
1:A:784:A:O3'	22:V:60:ALA:C	2.30	0.75
16:P:52:LEU:HG	16:P:75:ILE:HG12	1.69	0.75
1:A:785:G:O5'	22:V:70:HIS:C	2.30	0.74
1:A:1482:G:H1'	22:V:311:LYS:O	1.87	0.74
1:A:783:C:H5''	22:V:58:MET:HB3	1.68	0.74
1:A:1103:C:H5''	2:B:96:LEU:HD12	1.68	0.74
1:A:1486:G:C4'	22:V:285:HIS:CA	2.65	0.74
3:C:5:HIS:HD2	3:C:8:GLY:H	1.35	0.74
1:A:1485:U:C4'	22:V:286:LEU:CD1	2.10	0.74
1:A:135:C:O2	16:P:1:MET:HB2	1.87	0.74
1:A:783:C:H5'	22:V:58:MET:HB3	1.69	0.74
1:A:1407:C:C4'	22:V:284:TRP:C	2.58	0.74
1:A:1484:C:C6	22:V:340:ILE:HB	2.22	0.74
13:M:15:VAL:HG23	13:M:33:LEU:HD12	1.70	0.74
2:B:163:ILE:HD11	2:B:209:VAL:HG12	1.70	0.74
11:K:110:THR:HG21	21:U:4:LYS:HD2	1.69	0.74
1:A:785:G:H8	22:V:71:ARG:HB2	1.51	0.74
1:A:1484:C:H1'	22:V:291:ILE:C	1.95	0.74
5:E:108:GLY:H	5:E:110:MET:HE1	1.53	0.74
7:G:67:ASN:HD22	7:G:127:ALA:HA	1.51	0.74
12:L:66:ILE:HD13	12:L:73:LEU:HD12	1.69	0.74
1:A:1483:A:O4'	22:V:298:PHE:CZ	2.40	0.74
1:A:278:G:H21	1:A:279:A:H62	1.36	0.73
1:A:1486:G:H8	22:V:344:MET:CE	2.01	0.73
16:P:71:VAL:HA	16:P:74:LEU:HG	1.69	0.73
1:A:1482:G:C8	22:V:313:ASP:HB2	2.23	0.73
21:U:20:ARG:HA	21:U:24:LYS:HG3	1.71	0.73
1:A:1484:C:C4	22:V:341:LEU:CD1	2.71	0.73
13:M:44:ILE:HD12	13:M:45:SER:H	1.52	0.73
3:C:153:SER:HB2	3:C:196:GLY:H	1.54	0.73
3:C:179:ALA:HB1	3:C:202:PHE:HE1	1.52	0.73
4:D:10:LEU:HD21	4:D:62:ARG:HD3	1.70	0.73
1:A:1517:G:H5''	22:V:125:LYS:N	2.03	0.73
10:J:39:PRO:HA	10:J:74:VAL:HG22	1.70	0.73
16:P:26:ASN:CG	16:P:31:ARG:HB3	2.13	0.73
21:U:40:PRO:HG2	21:U:41:THR:H	1.54	0.73
1:A:1308:U:H2'	1:A:1309:G:C8	2.24	0.73
1:A:764:C:H3'	1:A:765:G:H21	1.51	0.73
1:A:812:G:H2'	1:A:812:G:N3	2.02	0.73
1:A:927:G:H4'	1:A:1503:A:N7	2.03	0.73

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
4:D:84:ASN:ND2	4:D:87:GLU:H	1.87	0.73
1:A:430:A:OP1	4:D:8:LEU:HB2	1.89	0.73
1:A:501:C:H2'	1:A:502:A:H8	1.53	0.73
2:B:204:ASP:CG	2:B:205:ALA:H	1.95	0.73
1:A:1418:A:C5	22:V:312:HIS:NE2	2.57	0.72
1:A:1419:G:C5'	22:V:309:ASP:CB	2.67	0.72
1:A:1461:G:H2'	1:A:1462:C:C6	2.23	0.72
1:A:1484:C:C4	22:V:341:LEU:HD13	2.24	0.72
14:N:63:CYS:HB3	14:N:67:GLY:N	2.04	0.72
11:K:42:GLY:HA3	11:K:73:VAL:HG22	1.71	0.72
4:D:94:GLU:HG3	4:D:103:ARG:HH12	1.55	0.72
4:D:156:ALA:O	4:D:159:GLU:HG2	1.90	0.72
1:A:946:A:H2'	1:A:947:G:C8	2.24	0.72
1:A:1125:U:H2'	1:A:1126:U:H5''	1.70	0.72
5:E:131:ASN:HD21	5:E:133:ILE:HB	1.54	0.72
1:A:429:U:H3'	4:D:8:LEU:HD23	1.70	0.72
1:A:1218:C:H2'	1:A:1219:A:H8	1.55	0.72
1:A:1250:A:H4'	9:I:69:GLY:H	1.54	0.72
11:K:113:THR:HG21	21:U:28:LEU:HD11	1.72	0.72
2:B:67:LEU:H	2:B:67:LEU:HD22	1.52	0.72
1:A:1407:C:C3'	22:V:284:TRP:C	2.62	0.72
1:A:1408:A:OP1	22:V:283:LEU:C	2.28	0.72
9:I:71:ILE:H	9:I:71:ILE:HD12	1.55	0.72
13:M:44:ILE:HA	13:M:47:LEU:HD13	1.71	0.72
16:P:67:ILE:HG13	16:P:71:VAL:HG13	1.71	0.72
1:A:412:A:H1'	1:A:413:G:H8	1.54	0.72
10:J:92:LEU:HD22	10:J:92:LEU:H	1.55	0.72
7:G:100:MET:HA	7:G:103:ILE:HD12	1.71	0.71
8:H:49:LYS:HG3	8:H:50:VAL:H	1.55	0.71
1:A:1218:C:H2'	1:A:1219:A:C8	2.25	0.71
1:A:1471:U:O2'	1:A:1472:U:H5'	1.90	0.71
1:A:1407:C:H4'	22:V:284:TRP:CB	2.20	0.71
11:K:92:ARG:HH11	11:K:92:ARG:HB3	1.55	0.71
20:T:60:GLN:HE21	20:T:61:ALA:H	1.36	0.71
1:A:878:A:H5''	8:H:80:PRO:HG2	1.72	0.71
1:A:1057:G:H5''	3:C:153:SER:HB3	1.72	0.71
6:F:86:ARG:HH11	18:R:63:TYR:HB3	1.55	0.71
1:A:1415:G:C2'	22:V:345:ALA:CB	1.90	0.71
2:B:46:VAL:HA	2:B:49:PHE:HD2	1.55	0.71
2:B:116:LEU:HD11	2:B:139:GLU:HB3	1.71	0.71
13:M:77:LYS:HA	13:M:80:MET:HB3	1.71	0.71

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
13:M:78:ARG:NH2	13:M:79:LEU:HG	2.05	0.71
17:Q:29:LYS:HD3	17:Q:35:LYS:N	2.05	0.71
21:U:29:ALA:CB	21:U:32:ARG:HH21	2.02	0.71
22:V:73:ASN:HD21	22:V:102:LYS:HE3	1.56	0.71
12:L:98:ARG:HA	12:L:98:ARG:HE	1.55	0.71
17:Q:20:ILE:HG13	17:Q:45:VAL:HB	1.73	0.71
14:N:30:ILE:HD12	14:N:30:ILE:H	1.56	0.71
1:A:922:G:H2'	1:A:923:A:C8	2.26	0.70
1:A:1330:U:H2'	1:A:1331:G:H5'	1.72	0.70
11:K:56:LYS:O	11:K:58:THR:HG22	1.90	0.70
3:C:72:PRO:O	3:C:76:ILE:HG12	1.91	0.70
1:A:1179:A:H4'	9:I:104:THR:HA	1.73	0.70
1:A:1409:C:H42	1:A:1491:G:H1	1.39	0.70
4:D:194:ILE:HD13	4:D:195:ASN:N	2.06	0.70
12:L:35:ARG:HG3	12:L:36:VAL:H	1.56	0.70
14:N:50:LEU:HD23	14:N:51:PRO:HD3	1.73	0.70
16:P:42:ILE:HG22	16:P:43:ALA:H	1.55	0.70
20:T:81:GLN:O	20:T:85:LEU:HB3	1.90	0.70
1:A:186:C:H4'	20:T:75:LYS:HB2	1.74	0.70
1:A:239:U:H4'	1:A:239:U:OP1	1.90	0.70
1:A:1182:G:H4'	1:A:1183:U:C5'	2.21	0.70
1:A:1407:C:C3'	22:V:284:TRP:CA	2.56	0.70
1:A:1458:G:H5'	20:T:26:MET:HB2	1.73	0.70
1:A:1483:A:N7	22:V:341:LEU:HD22	2.07	0.70
9:I:56:MET:HA	9:I:59:LYS:HZ2	1.56	0.70
19:S:48:ILE:HB	19:S:59:VAL:HB	1.73	0.70
1:A:1481:U:O2	22:V:313:ASP:HB3	1.90	0.70
3:C:57:GLU:HB2	3:C:64:ARG:HB2	1.73	0.70
5:E:95:MET:HE1	5:E:114:LEU:HD21	1.72	0.70
16:P:7:ALA:HB1	16:P:29:ASN:HB3	1.74	0.70
1:A:390:U:H2'	1:A:391:G:C8	2.27	0.70
2:B:212:TYR:O	2:B:216:VAL:HG23	1.92	0.70
13:M:38:ILE:HG13	13:M:55:LEU:HD21	1.72	0.70
1:A:1417:G:O5'	22:V:288:PRO:HG2	1.90	0.70
9:I:40:ARG:H	9:I:44:ARG:HE	1.39	0.70
22:V:140:ILE:HG21	22:V:173:LEU:HD21	1.74	0.70
1:A:814:A:H5'	1:A:1511:G:H4'	1.73	0.70
1:A:1420:U:H5''	22:V:307:TYR:HB2	1.72	0.70
10:J:52:LEU:HB2	14:N:80:ARG:HD2	1.74	0.70
21:U:14:ALA:CA	21:U:16:ARG:HD2	2.22	0.70
10:J:56:HIS:O	10:J:57:VAL:HG12	1.92	0.70

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
11:K:19:VAL:HG12	11:K:82:GLU:HB2	1.74	0.70
11:K:88:PRO:HD3	21:U:28:LEU:HD13	1.73	0.70
1:A:785:G:H5''	22:V:61:ASP:CG	2.16	0.69
7:G:13:PRO:HB2	7:G:18:GLY:HA2	1.72	0.69
1:A:71:A:H61	1:A:99:C:H1'	1.56	0.69
1:A:785:G:C8	22:V:71:ARG:HB2	2.25	0.69
1:A:1486:G:C5	22:V:344:MET:CA	2.71	0.69
2:B:67:LEU:HD12	2:B:153:MET:HE2	1.74	0.69
7:G:38:ALA:O	7:G:41:ILE:HG22	1.92	0.69
1:A:673:A:H2'	1:A:674:G:C8	2.27	0.69
2:B:141:GLU:O	2:B:145:ASN:HB2	1.93	0.69
12:L:20:VAL:HG13	12:L:94:TYR:HH	1.55	0.69
1:A:279:A:H5''	1:A:280:C:H3'	1.74	0.69
1:A:780:A:O2'	1:A:781:A:H5''	1.91	0.69
1:A:785:G:O5'	22:V:71:ARG:N	2.26	0.69
12:L:80:LEU:HD23	12:L:97:VAL:HG21	1.74	0.69
1:A:1326:U:H2'	1:A:1327:C:C6	2.27	0.69
1:A:1482:G:O5'	22:V:338:HIS:ND1	2.26	0.69
9:I:112:ARG:NH1	9:I:112:ARG:HB2	2.07	0.69
1:A:784:A:H4'	22:V:72:CYS:CA	2.23	0.69
4:D:12:ARG:HD2	4:D:37:PRO:HA	1.75	0.69
5:E:101:GLY:H	5:E:121:ASN:HD21	1.41	0.69
3:C:141:MET:HA	3:C:141:MET:HE3	1.75	0.69
12:L:28:GLN:HG3	12:L:80:LEU:HD21	1.73	0.69
21:U:7:GLU:OE2	21:U:15:LEU:HD22	1.93	0.69
1:A:784:A:H5''	22:V:72:CYS:O	1.93	0.68
1:A:1251:A:H2'	1:A:1252:A:C8	2.28	0.68
13:M:109:LYS:HG3	13:M:110:GLY:H	1.59	0.68
19:S:49:ALA:HA	19:S:57:VAL:O	1.93	0.68
1:A:268:U:H2'	1:A:269:C:C6	2.28	0.68
1:A:1483:A:C4'	22:V:298:PHE:CZ	2.77	0.68
3:C:154:GLY:HA3	3:C:162:ALA:HB1	1.76	0.68
7:G:149:ALA:HB1	11:K:58:THR:HG21	1.75	0.68
12:L:24:GLU:HB3	12:L:26:CYS:SG	2.34	0.68
12:L:79:ILE:HD13	12:L:96:THR:HG22	1.74	0.68
15:O:68:TYR:CZ	15:O:72:LYS:HG3	2.29	0.68
1:A:1412:C:H2'	1:A:1413:A:C8	2.29	0.68
10:J:40:ILE:HG13	10:J:73:LEU:HB3	1.75	0.68
1:A:541:G:O2'	4:D:39:GLN:HB2	1.92	0.68
1:A:1320:C:H41	19:S:36:ARG:HG2	1.57	0.68
1:A:1416:G:N1	22:V:341:LEU:HA	2.00	0.68

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:B:102:ASN:O	2:B:106:VAL:HG23	1.93	0.68
3:C:141:MET:HE1	3:C:147:GLY:H	1.59	0.68
10:J:87:LEU:HB3	10:J:88:MET:HE3	1.75	0.68
10:J:6:ILE:HG12	10:J:102:LEU:HD11	1.74	0.68
1:A:901:A:H5'	1:A:902:G:OP2	1.94	0.68
6:F:37:HIS:CE1	6:F:65:GLU:HB2	2.29	0.68
14:N:66:THR:HG23	14:N:67:GLY:H	1.59	0.68
1:A:1000:A:H2'	1:A:1001:C:C6	2.29	0.68
1:A:1415:G:N7	22:V:342:GLU:O	2.27	0.68
8:H:102:VAL:HG12	8:H:125:ILE:HD12	1.76	0.68
1:A:1021:A:H2'	1:A:1022:A:O4'	1.93	0.68
1:A:1484:C:C2'	22:V:291:ILE:HD12	2.23	0.68
2:B:67:LEU:HD21	2:B:157:PRO:HG3	1.76	0.68
3:C:133:MET:HE2	3:C:152:VAL:HG21	1.76	0.68
4:D:159:GLU:HG3	4:D:160:LEU:H	1.59	0.68
18:R:46:THR:HG23	18:R:51:GLN:HB2	1.75	0.68
1:A:518:C:H2'	1:A:530:G:C8	2.28	0.67
1:A:1071:C:H2'	1:A:1072:G:H8	1.60	0.67
1:A:1208:C:H2'	1:A:1209:C:O4'	1.93	0.67
8:H:111:THR:H	8:H:114:ALA:HB3	1.60	0.67
8:H:111:THR:HG23	8:H:114:ALA:HB2	1.76	0.67
13:M:64:VAL:HA	13:M:68:LEU:HD12	1.76	0.67
1:A:473:U:H2'	1:A:474:G:H8	1.58	0.67
1:A:1419:G:C4'	22:V:308:ARG:O	2.42	0.67
2:B:119:GLN:HE22	2:B:127:LYS:HD3	1.60	0.67
18:R:58:ILE:O	18:R:62:ARG:HG3	1.94	0.67
21:U:14:ALA:HA	21:U:16:ARG:HD2	1.76	0.67
1:A:619:U:N3	4:D:131:ILE:HD12	2.09	0.67
1:A:785:G:O4'	22:V:71:ARG:N	2.27	0.67
3:C:9:ILE:HG23	3:C:10:ARG:HG3	1.76	0.67
1:A:1420:U:C5'	22:V:307:TYR:O	2.43	0.67
1:A:967:C:H3'	1:A:968:A:H5'	1.75	0.67
1:A:1241:G:H2'	1:A:1242:G:H8	1.59	0.67
3:C:46:LEU:HD11	3:C:86:LEU:HD21	1.76	0.67
8:H:64:TYR:HB3	8:H:69:ALA:HA	1.77	0.67
14:N:60:ARG:NH2	14:N:62:ARG:HE	1.92	0.67
1:A:1416:G:N7	22:V:345:ALA:N	2.42	0.67
1:A:1486:G:O4'	22:V:285:HIS:CA	2.37	0.67
6:F:86:ARG:NH1	18:R:63:TYR:HB3	2.09	0.67
20:T:28:ARG:HA	20:T:31:ILE:HD12	1.77	0.67
1:A:501:C:H2'	1:A:502:A:C8	2.30	0.67

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:1060:U:H2'	1:A:1061:G:H8	1.58	0.67
1:A:1418:A:C4	22:V:312:HIS:CD2	2.83	0.67
2:B:202:ASN:ND2	2:B:204:ASP:H	1.92	0.67
10:J:71:LEU:H	10:J:71:LEU:HD12	1.58	0.67
1:A:1416:G:C6	22:V:344:MET:HB2	2.30	0.67
6:F:10:VAL:HA	6:F:84:VAL:HA	1.77	0.67
13:M:9:PRO:O	13:M:44:ILE:HG12	1.94	0.67
1:A:17:U:H2'	1:A:18:C:C6	2.30	0.67
3:C:33:ASP:HB2	14:N:64:ARG:HD3	1.75	0.67
1:A:1236:A:H4'	1:A:1304:G:H4'	1.76	0.66
3:C:77:GLY:HA3	3:C:82:ASP:H	1.59	0.66
16:P:51:ARG:HB3	16:P:51:ARG:NH1	2.10	0.66
1:A:1060:U:H5''	10:J:53:ILE:HG22	1.75	0.66
2:B:198:VAL:HG22	2:B:200:PRO:HD3	1.76	0.66
5:E:109:ALA:HB3	5:E:135:VAL:HG23	1.76	0.66
21:U:3:ILE:HG12	21:U:19:LYS:HG2	1.77	0.66
13:M:106:ARG:NE	13:M:112:ARG:HG2	2.08	0.66
20:T:48:LYS:O	20:T:52:GLU:HB2	1.96	0.66
1:A:1482:G:O4'	22:V:312:HIS:O	2.13	0.66
1:A:922:G:H4'	5:E:24:VAL:HA	1.77	0.66
1:A:1302:C:O4'	13:M:16:ILE:HD11	1.96	0.66
1:A:1420:U:C5	22:V:309:ASP:O	2.49	0.66
10:J:37:ARG:NE	10:J:77:VAL:HG11	2.11	0.66
13:M:106:ARG:HH11	13:M:106:ARG:HA	1.61	0.66
1:A:1026:G:H2'	1:A:1027:C:C6	2.31	0.66
11:K:69:CYS:C	11:K:71:ASP:H	2.01	0.66
14:N:60:ARG:HE	14:N:62:ARG:NE	1.93	0.66
1:A:1484:C:C3'	22:V:340:ILE:HD13	2.20	0.66
2:B:52:ALA:O	2:B:56:LEU:HD22	1.96	0.66
9:I:33:SER:HB3	9:I:36:GLN:HB2	1.76	0.66
17:Q:10:ARG:HH11	17:Q:55:GLY:H	1.43	0.66
1:A:1483:A:H4'	22:V:298:PHE:HZ	1.61	0.66
1:A:1408:A:OP1	22:V:283:LEU:CG	2.44	0.66
5:E:23:THR:HG23	5:E:28:ARG:HD3	1.78	0.66
1:A:1123:U:O2'	1:A:1124:G:H5'	1.96	0.66
1:A:1415:G:C2'	22:V:345:ALA:HB2	2.16	0.66
3:C:176:THR:HB	3:C:179:ALA:HB2	1.77	0.66
10:J:53:ILE:HG12	10:J:63:ASP:HB2	1.78	0.66
10:J:65:TYR:HB3	14:N:95:LEU:HD11	1.77	0.66
1:A:1484:C:N3	22:V:341:LEU:HD12	2.10	0.65
2:B:53:LEU:HD21	2:B:212:TYR:HE1	1.60	0.65

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:131:A:H2'	1:A:132:C:C6	2.32	0.65
1:A:238:A:H2'	1:A:239:U:H5''	1.77	0.65
1:A:559:A:H4'	1:A:560:A:H3'	1.77	0.65
1:A:1029:U:H5''	1:A:1030:U:H5	1.60	0.65
1:A:1062:U:H2'	1:A:1063:C:C6	2.31	0.65
4:D:19:PHE:HB2	4:D:110:ARG:HH12	1.61	0.65
6:F:3:HIS:HA	6:F:65:GLU:HA	1.77	0.65
10:J:11:LYS:HG2	10:J:97:ASP:HB3	1.78	0.65
1:A:205:A:H2'	1:A:206:C:H6	1.60	0.65
1:A:1342:C:H2'	1:A:1343:G:H8	1.61	0.65
1:A:1407:C:C5'	22:V:284:TRP:HB2	2.26	0.65
1:A:1484:C:C2	22:V:341:LEU:HD13	2.30	0.65
15:O:31:LEU:HD12	15:O:58:MET:HB2	1.78	0.65
21:U:36:PHE:HB3	21:U:40:PRO:CD	2.26	0.65
1:A:875:U:O2'	8:H:14:ARG:HD2	1.97	0.65
13:M:52:ILE:HD12	13:M:55:LEU:HD12	1.78	0.65
14:N:30:ILE:HB	14:N:44:VAL:HG11	1.78	0.65
1:A:1513:A:H2'	1:A:1514:G:C8	2.32	0.65
1:A:278:G:N2	1:A:279:A:H62	1.93	0.65
3:C:146:LYS:HE3	3:C:202:PHE:CE2	2.30	0.65
5:E:104:ILE:HD13	5:E:115:GLU:HG3	1.78	0.65
7:G:144:ALA:C	7:G:146:ALA:H	2.04	0.65
11:K:105:ARG:NH2	21:U:10:PRO:HG3	2.12	0.65
22:V:207:LYS:HB3	22:V:208:PRO:HD3	1.77	0.65
1:A:1017:U:H2'	1:A:1018:G:C8	2.31	0.65
4:D:123:MET:HA	4:D:128:VAL:HA	1.78	0.65
5:E:104:ILE:HD11	5:E:114:LEU:HB2	1.79	0.65
1:A:539:A:H2'	1:A:540:G:C8	2.32	0.65
1:A:1300:G:H1'	1:A:1301:U:H5	1.61	0.65
1:A:1342:C:H2'	1:A:1343:G:C8	2.32	0.65
1:A:1482:G:H2'	22:V:341:LEU:HD22	1.76	0.65
3:C:146:LYS:HD3	3:C:204:GLY:HA2	1.79	0.65
1:A:269:C:H2'	1:A:270:A:C8	2.32	0.65
3:C:13:ILE:O	3:C:14:VAL:HG22	1.96	0.65
3:C:185:THR:HG22	3:C:186:SER:H	1.62	0.65
14:N:12:ARG:HH11	14:N:60:ARG:NH1	1.95	0.65
1:A:620:C:H1'	4:D:131:ILE:HG21	1.79	0.64
1:A:1420:U:O3'	22:V:316:PRO:O	2.14	0.64
1:A:1225:A:H3'	1:A:1226:C:H6	1.61	0.64
1:A:1416:G:N1	22:V:341:LEU:HD12	2.12	0.64
1:A:1416:G:OP2	22:V:345:ALA:O	2.15	0.64

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:1486:G:C4'	22:V:285:HIS:HA	2.27	0.64
3:C:113:LYS:HE2	3:C:117:ASP:OD2	1.97	0.64
8:H:5:PRO:HB2	8:H:32:LYS:NZ	2.11	0.64
8:H:63:LYS:HG2	8:H:70:VAL:HG21	1.80	0.64
1:A:36:C:H5''	12:L:119:LYS:HB3	1.79	0.64
1:A:188:C:H2'	1:A:189:A:O4'	1.97	0.64
1:A:270:A:H2'	1:A:271:C:C6	2.32	0.64
1:A:562:U:H1'	12:L:11:ARG:HB3	1.78	0.64
1:A:1101:A:H61	2:B:173:LYS:HD2	1.63	0.64
8:H:30:LYS:HA	8:H:30:LYS:NZ	2.11	0.64
14:N:11:LYS:HZ2	14:N:11:LYS:HA	1.61	0.64
14:N:60:ARG:HD3	14:N:60:ARG:N	2.11	0.64
20:T:59:ARG:HB2	20:T:59:ARG:HH11	1.62	0.64
2:B:41:ASN:ND2	2:B:44:LYS:HB2	2.13	0.64
4:D:84:ASN:HD22	4:D:87:GLU:H	1.43	0.64
7:G:148:LYS:HG3	7:G:151:ALA:HB3	1.79	0.64
13:M:3:ILE:HG23	13:M:56:ARG:HG2	1.78	0.64
19:S:39:ILE:HG21	19:S:65:MET:HB3	1.80	0.64
1:A:105:G:H2'	1:A:106:C:C6	2.33	0.64
1:A:1221:G:H5''	19:S:35:ARG:HH11	1.62	0.64
1:A:1419:G:N2	22:V:313:ASP:N	2.46	0.64
1:A:1485:U:OP2	22:V:340:ILE:CD1	2.44	0.64
4:D:2:ARG:HG3	4:D:114:ARG:CZ	2.27	0.64
1:A:522:C:H41	12:L:49:ARG:HH22	1.46	0.64
1:A:783:C:C5'	22:V:58:MET:CB	2.69	0.64
1:A:1074:G:H5'	2:B:104:LYS:HZ1	1.63	0.64
6:F:40:GLU:HG3	6:F:42:TRP:HE1	1.60	0.64
6:F:53:LYS:HA	6:F:53:LYS:NZ	2.12	0.64
8:H:5:PRO:HB2	8:H:32:LYS:HZ1	1.62	0.64
8:H:9:MET:HE1	8:H:32:LYS:O	1.97	0.64
10:J:88:MET:HB2	10:J:89:ARG:NH1	2.11	0.64
14:N:51:PRO:HB2	14:N:54:SER:HB3	1.79	0.64
1:A:817:C:H1'	1:A:819:A:H5'	1.80	0.64
2:B:65:LYS:HD2	2:B:157:PRO:HA	1.80	0.64
3:C:61:LYS:O	3:C:96:VAL:HB	1.98	0.64
17:Q:59:GLU:H	17:Q:74:LEU:HD23	1.62	0.64
1:A:87:C:H2'	1:A:88:U:H4'	1.79	0.64
10:J:87:LEU:HD22	10:J:90:LEU:HD12	1.79	0.64
11:K:88:PRO:HA	11:K:92:ARG:HE	1.63	0.64
13:M:103:THR:HG22	13:M:104:ASN:H	1.62	0.64
20:T:15:LYS:HD3	20:T:18:LYS:HE3	1.79	0.64

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:269:C:H2'	1:A:270:A:H8	1.63	0.64
1:A:1326:U:H2'	1:A:1327:C:H6	1.61	0.64
1:A:1416:G:C4	22:V:344:MET:HB2	2.33	0.64
4:D:88:ASN:O	4:D:92:LEU:HD23	1.97	0.64
16:P:68:SER:HB3	16:P:71:VAL:HG12	1.80	0.64
19:S:31:ARG:HG3	19:S:56:HIS:NE2	2.13	0.64
1:A:1517:G:C4	22:V:118:PRO:O	2.51	0.64
2:B:118:THR:O	2:B:121:GLN:HB3	1.98	0.64
9:I:20:ILE:HG23	9:I:60:LEU:HD13	1.79	0.64
13:M:1:ALA:O	13:M:8:ILE:HG22	1.98	0.64
21:U:39:LYS:N	21:U:40:PRO:HD2	2.13	0.64
21:U:40:PRO:O	21:U:44:ARG:HB2	1.98	0.64
1:A:1391:U:H2'	1:A:1392:G:C8	2.34	0.63
3:C:178:ARG:O	3:C:178:ARG:HG2	1.96	0.63
1:A:1418:A:P	22:V:289:GLU:CD	2.81	0.63
1:A:203:G:H1'	1:A:465:A:N6	2.13	0.63
2:B:163:ILE:O	2:B:185:ILE:HG13	1.99	0.63
3:C:5:HIS:CD2	3:C:8:GLY:H	2.16	0.63
5:E:45:VAL:O	5:E:71:ILE:HG22	1.98	0.63
1:A:312:C:H2'	1:A:313:A:C8	2.33	0.63
1:A:785:G:C5'	22:V:69:VAL:CG1	2.58	0.63
1:A:1182:G:H4'	1:A:1183:U:H5'	1.79	0.63
1:A:1483:A:H4'	22:V:298:PHE:CZ	2.34	0.63
4:D:56:GLU:HG2	4:D:195:ASN:HD22	1.63	0.63
5:E:140:ILE:HA	5:E:143:LEU:HD12	1.81	0.63
15:O:29:ALA:HA	15:O:84:LEU:HD21	1.80	0.63
21:U:14:ALA:CA	21:U:16:ARG:NE	2.59	0.63
22:V:170:ILE:HB	22:V:178:MET:HE1	1.79	0.63
1:A:9:G:H5'	5:E:107:GLY:HA3	1.81	0.63
1:A:1416:G:H1'	22:V:285:HIS:CE1	2.34	0.63
1:A:1468:A:H2'	1:A:1469:C:O4'	1.98	0.63
6:F:92:THR:HG23	6:F:93:LYS:H	1.64	0.63
9:I:40:ARG:N	9:I:44:ARG:HE	1.96	0.63
9:I:113:LYS:HA	9:I:120:ALA:HB2	1.81	0.63
11:K:105:ARG:HH21	21:U:10:PRO:HG3	1.63	0.63
12:L:20:VAL:HG13	12:L:94:TYR:OH	1.99	0.63
16:P:42:ILE:HG22	16:P:43:ALA:N	2.14	0.63
1:A:785:G:N7	22:V:71:ARG:NH1	2.45	0.63
1:A:1484:C:C5	22:V:340:ILE:HB	2.32	0.63
2:B:148:GLY:O	2:B:151:LYS:HE2	1.99	0.63
6:F:9:MET:HA	6:F:58:HIS:O	1.99	0.63

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
9:I:51:LEU:HB3	9:I:56:MET:HG2	1.81	0.63
20:T:60:GLN:H	20:T:60:GLN:CD	2.07	0.63
1:A:859:G:H2'	1:A:860:A:C8	2.32	0.63
1:A:1038:C:H2'	1:A:1039:G:H8	1.64	0.63
1:A:1486:G:C2	22:V:344:MET:O	2.51	0.63
3:C:151:GLU:O	3:C:197:VAL:HA	1.99	0.63
16:P:9:HIS:HE1	16:P:29:ASN:HB2	1.63	0.63
17:Q:10:ARG:HA	17:Q:10:ARG:HE	1.62	0.63
17:Q:26:ARG:HE	17:Q:39:ARG:NH1	1.97	0.63
1:A:79:G:H2'	1:A:80:A:C8	2.34	0.63
1:A:1011:C:H2'	1:A:1012:A:H8	1.64	0.63
9:I:56:MET:HE2	9:I:57:VAL:H	1.64	0.63
10:J:67:ILE:HD11	14:N:95:LEU:HD22	1.81	0.63
21:U:36:PHE:HB3	21:U:40:PRO:CG	2.29	0.63
22:V:151:ARG:HB3	22:V:151:ARG:HH11	1.63	0.63
1:A:441:A:H61	1:A:493:A:H62	1.44	0.62
1:A:528:C:H41	12:L:45:ASN:CG	2.07	0.62
1:A:983:A:H5'	1:A:984:C:OP2	1.99	0.62
1:A:1416:G:C4	22:V:344:MET:CB	2.82	0.62
1:A:1482:G:N3	22:V:311:LYS:O	2.31	0.62
2:B:202:ASN:HD22	2:B:204:ASP:N	1.95	0.62
17:Q:10:ARG:HH11	17:Q:55:GLY:N	1.97	0.62
1:A:10:A:OP2	5:E:130:THR:HB	1.99	0.62
1:A:1118:U:H1'	1:A:1179:A:C4	2.34	0.62
1:A:1409:C:C5'	22:V:151:ARG:HE	2.10	0.62
2:B:75:ALA:O	2:B:79:VAL:HB	1.98	0.62
5:E:114:LEU:HD22	5:E:119:VAL:HG21	1.80	0.62
10:J:15:HIS:HA	10:J:18:ILE:HG22	1.79	0.62
14:N:15:LEU:HD12	14:N:16:ALA:N	2.14	0.62
1:A:476:U:H2'	1:A:477:C:C6	2.35	0.62
1:A:981:U:H4'	14:N:60:ARG:HG3	1.81	0.62
1:A:1447:A:H5'	1:A:1448:C:H5	1.63	0.62
1:A:1485:U:O4	22:V:339:ARG:O	2.17	0.62
4:D:99:ASN:ND2	4:D:110:ARG:HE	1.97	0.62
8:H:11:THR:HA	8:H:14:ARG:NH1	2.14	0.62
14:N:71:GLY:O	14:N:79:SER:HA	1.98	0.62
16:P:67:ILE:HD11	16:P:72:ALA:HA	1.81	0.62
21:U:14:ALA:CA	21:U:16:ARG:CD	2.78	0.62
1:A:982:U:H5''	14:N:5:MET:HE2	1.80	0.62
1:A:121:U:OP1	1:A:121:U:H3'	1.99	0.62
1:A:531:U:H5'	1:A:531:U:H6	1.63	0.62

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:882:C:O2'	1:A:883:C:H5'	1.99	0.62
1:A:918:A:H2'	1:A:919:A:C8	2.35	0.62
1:A:1513:A:H2'	1:A:1514:G:H8	1.65	0.62
14:N:11:LYS:O	14:N:15:LEU:HG	2.00	0.62
21:U:26:GLY:C	21:U:28:LEU:H	2.05	0.62
4:D:99:ASN:CG	4:D:103:ARG:HH21	2.06	0.62
4:D:163:GLN:HB2	4:D:164:ARG:NH1	2.14	0.62
8:H:30:LYS:HA	8:H:30:LYS:HZ2	1.63	0.62
12:L:43:LYS:N	12:L:44:PRO:HD2	2.14	0.62
1:A:505:G:H5'	1:A:534:U:H2'	1.81	0.62
1:A:1485:U:C5'	22:V:286:LEU:HD12	2.20	0.62
1:A:1505:G:H4'	1:A:1506:U:H5''	1.81	0.62
9:I:30:ASN:ND2	9:I:65:THR:HA	2.14	0.62
11:K:86:LYS:HG3	11:K:113:THR:HA	1.82	0.62
1:A:60:A:H4'	1:A:61:G:O5'	1.99	0.62
1:A:78:A:H2'	1:A:79:G:H8	1.63	0.62
1:A:524:G:H2'	1:A:525:C:C6	2.33	0.62
1:A:1074:G:H5'	2:B:104:LYS:NZ	2.15	0.62
1:A:1225:A:H3'	1:A:1226:C:C6	2.35	0.62
1:A:1483:A:N1	22:V:288:PRO:O	2.33	0.62
14:N:78:LEU:HB2	14:N:83:VAL:HG22	1.82	0.62
20:T:4:LYS:HD2	20:T:5:SER:N	2.13	0.62
22:V:87:VAL:HG13	22:V:107:ALA:O	2.00	0.62
1:A:8:A:H2'	4:D:205:LYS:O	2.00	0.62
1:A:132:C:H5''	20:T:68:LYS:HE3	1.82	0.62
1:A:662:U:H2'	1:A:663:A:C8	2.34	0.62
1:A:1109:C:H42	1:A:1110:A:N6	1.98	0.62
2:B:17:HIS:HB2	2:B:202:ASN:OD1	2.00	0.62
3:C:184:ASN:HD22	3:C:185:THR:N	1.98	0.62
19:S:5:LYS:N	19:S:5:LYS:HE3	2.15	0.62
7:G:44:SER:O	7:G:48:THR:HG23	1.99	0.62
15:O:87:ARG:NH1	15:O:87:ARG:HA	2.15	0.62
17:Q:26:ARG:HH21	17:Q:39:ARG:CZ	2.13	0.62
1:A:37:U:P	12:L:119:LYS:HB2	2.40	0.61
1:A:1171:A:H2'	1:A:1172:C:H6	1.65	0.61
1:A:1518:A:H2'	1:A:1519:A:C8	2.35	0.61
7:G:49:LEU:HB2	7:G:57:GLU:HG3	1.82	0.61
22:V:170:ILE:HB	22:V:178:MET:CE	2.29	0.61
1:A:678:U:H2'	1:A:679:C:C6	2.34	0.61
1:A:785:G:H8	22:V:71:ARG:NE	1.96	0.61
1:A:1219:A:H2'	1:A:1220:G:H8	1.65	0.61

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:1420:U:H5	22:V:309:ASP:O	1.83	0.61
2:B:27:LYS:HB3	2:B:28:PRO:HD3	1.81	0.61
3:C:69:THR:HG21	3:C:75:VAL:HG21	1.81	0.61
9:I:10:ARG:HB2	9:I:14:SER:O	2.00	0.61
12:L:85:ARG:HA	12:L:93:ARG:HA	1.80	0.61
1:A:785:G:H4'	22:V:70:HIS:HA	1.83	0.61
1:A:1285:A:H4'	1:A:1286:U:C5	2.36	0.61
1:A:1407:C:H5''	22:V:284:TRP:CB	2.30	0.61
2:B:69:VAL:O	2:B:163:ILE:HG22	2.00	0.61
4:D:94:GLU:HG3	4:D:103:ARG:NH1	2.16	0.61
5:E:33:THR:O	5:E:34:ALA:HB3	2.00	0.61
5:E:35:LEU:HD11	5:E:136:VAL:HG11	1.81	0.61
11:K:69:CYS:O	11:K:73:VAL:HG23	2.01	0.61
22:V:66:ASP:OD1	22:V:68:GLU:HG2	1.99	0.61
1:A:80:A:H2'	1:A:81:A:O4'	2.01	0.61
1:A:784:A:N7	22:V:71:ARG:NH1	2.40	0.61
3:C:45:GLU:C	3:C:46:LEU:HD12	2.25	0.61
4:D:20:LEU:H	4:D:20:LEU:HD12	1.65	0.61
5:E:85:LYS:HE2	5:E:92:ARG:NH1	2.16	0.61
19:S:4:LEU:HD13	19:S:9:PHE:N	2.13	0.61
1:A:69:G:H2'	1:A:70:U:C6	2.34	0.61
1:A:1060:U:H2'	1:A:1061:G:C8	2.35	0.61
1:A:1419:G:H22	22:V:313:ASP:N	1.98	0.61
10:J:25:ILE:HG23	10:J:29:ALA:HB3	1.83	0.61
15:O:11:VAL:HG23	15:O:26:VAL:HG11	1.82	0.61
1:A:212:G:H2'	1:A:213:G:H8	1.65	0.61
1:A:1295:U:H2'	1:A:1296:C:C6	2.35	0.61
2:B:76:SER:O	2:B:79:VAL:HG12	2.01	0.61
9:I:87:MET:HE3	9:I:94:ARG:HE	1.66	0.61
10:J:80:THR:H	10:J:84:VAL:HG11	1.66	0.61
13:M:68:LEU:O	13:M:72:ILE:HD13	1.99	0.61
13:M:98:GLY:H	13:M:99:GLN:HE21	1.49	0.61
18:R:52:ARG:HB3	18:R:52:ARG:NH1	2.14	0.61
1:A:1254:A:OP1	10:J:47:GLU:HG3	2.01	0.61
1:A:1307:U:H2'	1:A:1308:U:C6	2.36	0.61
1:A:1415:G:O6	22:V:342:GLU:C	2.43	0.61
1:A:1418:A:N6	22:V:337:TYR:CZ	2.67	0.61
5:E:152:VAL:HG21	8:H:98:LEU:HD23	1.83	0.61
13:M:5:GLY:O	13:M:6:ILE:HG12	2.01	0.61
22:V:87:VAL:HG12	22:V:88:VAL:N	2.16	0.61
1:A:1482:G:C5'	22:V:313:ASP:HA	2.30	0.61

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
13:M:88:LEU:O	13:M:92:ARG:HG3	2.00	0.61
17:Q:59:GLU:H	17:Q:74:LEU:CD2	2.14	0.61
19:S:11:ASP:CG	19:S:34:SER:HB2	2.26	0.61
1:A:501:C:H1'	1:A:549:C:H1'	1.81	0.60
1:A:590:U:H2'	1:A:591:U:H6	1.66	0.60
1:A:1281:C:H5'	1:A:1282:C:H5	1.65	0.60
1:A:1417:G:O3'	22:V:289:GLU:OE1	2.18	0.60
3:C:59:PRO:HG2	3:C:60:ALA:H	1.65	0.60
1:A:832:G:O2'	1:A:833:G:H5'	2.01	0.60
1:A:858:G:O6	1:A:869:G:H3'	2.01	0.60
1:A:1300:G:H1'	1:A:1301:U:C5	2.36	0.60
2:B:128:LEU:HG	2:B:132:GLU:HG2	1.83	0.60
4:D:123:MET:HE3	4:D:126:GLY:HA2	1.81	0.60
4:D:154:VAL:HG13	4:D:155:LYS:H	1.66	0.60
7:G:3:ARG:HB3	7:G:3:ARG:NH1	2.16	0.60
7:G:59:GLU:O	7:G:63:VAL:HG23	2.01	0.60
9:I:15:ALA:O	9:I:66:VAL:HG23	2.01	0.60
12:L:64:SER:OG	12:L:96:THR:HG23	2.00	0.60
14:N:20:PHE:HA	14:N:24:ALA:HB2	1.83	0.60
19:S:50:VAL:O	19:S:56:HIS:HA	2.00	0.60
22:V:321:ARG:O	22:V:325:GLU:HG3	2.01	0.60
1:A:763:G:H2'	1:A:764:C:C6	2.36	0.60
5:E:96:GLN:HB3	5:E:123:LEU:CD1	2.31	0.60
5:E:110:MET:HB3	5:E:139:THR:HG21	1.82	0.60
14:N:63:CYS:SG	14:N:66:THR:HG22	2.41	0.60
1:A:105:G:H2'	1:A:106:C:H6	1.66	0.60
1:A:1074:G:H2'	1:A:1075:U:C6	2.35	0.60
1:A:1482:G:O4'	22:V:313:ASP:HA	2.01	0.60
4:D:58:GLN:HA	4:D:61:ARG:HB2	1.83	0.60
12:L:78:VAL:HG12	12:L:101:LEU:HD23	1.82	0.60
1:A:398:U:H2'	1:A:399:G:H8	1.66	0.60
1:A:763:G:H2'	1:A:764:C:H6	1.64	0.60
1:A:1257:A:H2'	1:A:1257:A:N3	2.16	0.60
2:B:96:LEU:HD21	2:B:146:SER:HB2	1.82	0.60
7:G:97:ALA:HA	7:G:100:MET:HE3	1.82	0.60
7:G:117:LEU:HD22	7:G:120:ALA:HB3	1.83	0.60
9:I:35:GLU:HA	9:I:39:GLY:HA3	1.82	0.60
11:K:124:LYS:CA	21:U:34:ARG:HB3	2.29	0.60
19:S:7:GLY:H	19:S:8:PRO:HD3	1.67	0.60
1:A:1193:G:O2'	1:A:1194:U:H5'	2.01	0.60
1:A:1260:G:H4'	1:A:1284:C:H5'	1.84	0.60

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:1419:G:H22	22:V:313:ASP:H	1.48	0.60
4:D:7:LYS:HE2	4:D:20:LEU:HD22	1.83	0.60
4:D:20:LEU:O	4:D:21:LYS:HE3	2.02	0.60
4:D:120:LYS:HB2	4:D:145:ARG:HH21	1.66	0.60
5:E:155:LYS:HA	8:H:65:PHE:CD1	2.36	0.60
9:I:18:VAL:HG11	9:I:82:ILE:HG12	1.83	0.60
9:I:117:LEU:HD22	9:I:123:ARG:HG2	1.83	0.60
13:M:98:GLY:H	13:M:99:GLN:NE2	2.00	0.60
21:U:14:ALA:N	21:U:16:ARG:HH12	1.81	0.60
1:A:590:U:H2'	1:A:591:U:C6	2.35	0.60
1:A:1011:C:H2'	1:A:1012:A:C8	2.37	0.60
15:O:55:LEU:O	15:O:59:VAL:HG23	2.01	0.60
20:T:54:GLN:N	20:T:55:PRO:HD2	2.16	0.60
1:A:57:G:H2'	1:A:58:C:C6	2.37	0.60
1:A:801:U:H2'	1:A:802:A:H8	1.66	0.60
1:A:1482:G:C2	22:V:311:LYS:HB3	2.37	0.60
1:A:1485:U:H1'	22:V:286:LEU:HB2	1.82	0.60
4:D:191:SER:O	4:D:192:ALA:HB2	2.01	0.60
6:F:40:GLU:HG3	6:F:42:TRP:NE1	2.16	0.60
7:G:22:LEU:O	7:G:25:PHE:HB3	2.01	0.60
7:G:64:ALA:HA	7:G:127:ALA:HB2	1.82	0.60
13:M:21:ILE:CG2	13:M:64:VAL:HG11	2.32	0.60
1:A:1225:A:H4'	19:S:77:ARG:NH2	2.16	0.60
1:A:1323:G:H2'	1:A:1324:A:H8	1.65	0.60
1:A:1484:C:H2'	22:V:340:ILE:HG21	1.82	0.60
11:K:95:THR:HG23	11:K:96:ILE:H	1.65	0.60
15:O:60:SER:HA	15:O:63:ARG:NH1	2.17	0.60
1:A:1418:A:OP1	22:V:289:GLU:OE2	2.19	0.60
1:A:1438:G:O2'	1:A:1439:G:H5'	2.00	0.60
2:B:31:PHE:HB3	2:B:39:ILE:HB	1.84	0.60
2:B:166:ASP:OD1	2:B:190:SER:HA	2.02	0.60
3:C:5:HIS:CD2	14:N:88:MET:HB3	2.37	0.60
3:C:30:ASP:O	3:C:33:ASP:HB3	2.01	0.60
19:S:28:LYS:HB2	19:S:29:PRO:HD2	1.84	0.60
19:S:50:VAL:HB	19:S:57:VAL:HG22	1.84	0.60
8:H:42:GLU:HG3	8:H:100:ILE:HD13	1.83	0.59
16:P:46:LYS:H	16:P:46:LYS:CD	2.10	0.59
1:A:429:U:H1'	1:A:430:A:H5''	1.84	0.59
1:A:699:C:C2'	1:A:700:G:H5''	2.30	0.59
1:A:1074:G:H2'	1:A:1075:U:H6	1.67	0.59
1:A:1416:G:H1	22:V:341:LEU:HD12	1.66	0.59

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:1421:G:P	22:V:316:PRO:O	2.60	0.59
2:B:37:VAL:HG22	2:B:38:HIS:N	2.17	0.59
4:D:90:LEU:HA	4:D:93:LEU:HD12	1.83	0.59
5:E:156:ARG:HD2	8:H:42:GLU:O	2.02	0.59
12:L:35:ARG:HA	12:L:35:ARG:HE	1.67	0.59
17:Q:7:LEU:HD13	17:Q:24:ILE:HG12	1.84	0.59
20:T:60:GLN:NE2	20:T:61:ALA:H	1.99	0.59
1:A:93:U:H3'	1:A:94:G:H5''	1.83	0.59
1:A:1090:U:H2'	1:A:1091:U:C6	2.37	0.59
1:A:1111:A:HO2'	1:A:1112:C:H6	1.50	0.59
1:A:1361:G:H2'	1:A:1362:A:H5''	1.84	0.59
12:L:19:ASN:HB2	12:L:93:ARG:NH1	2.18	0.59
17:Q:28:VAL:C	17:Q:36:PHE:HA	2.27	0.59
1:A:194:C:O2'	1:A:195:A:H5'	2.01	0.59
4:D:11:SER:OG	4:D:17:ASP:HA	2.02	0.59
4:D:118:SER:HA	4:D:130:ASN:HB2	1.85	0.59
7:G:11:ILE:HG12	7:G:24:LYS:HE2	1.85	0.59
9:I:17:ARG:O	9:I:64:ILE:HA	2.02	0.59
10:J:15:HIS:HD2	10:J:18:ILE:HG22	1.65	0.59
15:O:78:THR:HA	15:O:81:ILE:HG12	1.85	0.59
1:A:720:C:H5''	18:R:40:PRO:HA	1.84	0.59
1:A:926:G:N2	1:A:1505:G:H2'	2.18	0.59
1:A:1048:G:H4'	14:N:2:LYS:NZ	2.18	0.59
2:B:53:LEU:HD21	2:B:212:TYR:CE1	2.37	0.59
4:D:59:LYS:O	4:D:63:ILE:HG13	2.02	0.59
10:J:38:GLY:O	10:J:74:VAL:HA	2.02	0.59
1:A:95:C:O2	1:A:95:C:H2'	2.02	0.59
1:A:436:C:O2'	1:A:437:U:H5'	2.03	0.59
1:A:1026:G:H2'	1:A:1027:C:H6	1.67	0.59
1:A:1278:G:H4'	1:A:1279:G:O5'	2.01	0.59
4:D:94:GLU:HA	4:D:103:ARG:HH22	1.67	0.59
4:D:138:PRO:HA	4:D:181:PHE:HD2	1.67	0.59
12:L:42:LYS:HE3	12:L:90:PRO:HD3	1.84	0.59
1:A:626:G:H2'	1:A:627:G:C8	2.38	0.59
1:A:1419:G:N2	22:V:313:ASP:H	1.99	0.59
1:A:1483:A:N7	22:V:341:LEU:CD2	2.66	0.59
4:D:69:ARG:HA	4:D:69:ARG:HE	1.68	0.59
7:G:74:VAL:HG12	7:G:75:LYS:H	1.68	0.59
10:J:26:VAL:O	10:J:30:LYS:HB3	2.02	0.59
11:K:83:VAL:HG23	11:K:109:ILE:HA	1.84	0.59
1:A:1243:C:H2'	1:A:1244:G:H8	1.67	0.59

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:1485:U:H4'	22:V:286:LEU:CG	2.22	0.59
8:H:12:ARG:HD3	8:H:26:MET:HE3	1.84	0.59
1:A:677:U:H3	1:A:713:G:H22	1.51	0.59
3:C:112:ALA:HB1	3:C:199:VAL:HG23	1.84	0.59
6:F:99:ALA:O	6:F:100:SER:HB2	2.01	0.59
8:H:8:ASP:O	8:H:12:ARG:HG3	2.03	0.59
9:I:18:VAL:HG13	9:I:64:ILE:HG12	1.85	0.59
10:J:56:HIS:N	14:N:80:ARG:HH22	1.98	0.59
14:N:61:ASN:O	14:N:62:ARG:HB2	2.03	0.59
16:P:51:ARG:O	16:P:52:LEU:HD13	2.03	0.59
1:A:241:G:O2'	1:A:242:G:H5'	2.03	0.58
1:A:1408:A:OP1	22:V:283:LEU:HG	2.03	0.58
1:A:1482:G:O4'	22:V:313:ASP:CA	2.50	0.58
9:I:18:VAL:HG22	9:I:64:ILE:HG12	1.85	0.58
13:M:89:ARG:HB2	13:M:96:VAL:HG12	1.84	0.58
16:P:1:MET:N	16:P:24:SER:HB3	2.18	0.58
17:Q:10:ARG:HA	17:Q:10:ARG:NE	2.18	0.58
1:A:118:U:O4	1:A:288:A:H2'	2.02	0.58
2:B:26:MET:HE3	2:B:192:PRO:HG3	1.84	0.58
4:D:25:ARG:NH1	4:D:30:LYS:HG2	2.18	0.58
7:G:53:SER:HB2	7:G:55:LYS:NZ	2.18	0.58
11:K:18:GLY:O	11:K:81:LEU:HB2	2.04	0.58
16:P:1:MET:HE1	16:P:64:GLY:O	2.03	0.58
1:A:370:C:H2'	1:A:371:A:H8	1.67	0.58
1:A:967:C:H3'	1:A:968:A:C5'	2.33	0.58
1:A:1100:C:OP2	2:B:94:ARG:HD3	2.04	0.58
2:B:41:ASN:ND2	2:B:44:LYS:HE2	2.17	0.58
2:B:67:LEU:HD13	2:B:67:LEU:N	2.18	0.58
3:C:183:TYR:HA	3:C:199:VAL:O	2.03	0.58
4:D:173:ASP:HB3	4:D:178:GLU:HB3	1.85	0.58
21:U:14:ALA:HA	21:U:16:ARG:NE	2.18	0.58
21:U:13:VAL:HG13	21:U:14:ALA:N	2.16	0.58
1:A:586:C:O2'	1:A:587:G:H5'	2.04	0.58
1:A:987:G:H2'	1:A:988:G:H8	1.68	0.58
1:A:1444:U:H2'	1:A:1445:U:C6	2.38	0.58
1:A:1483:A:C6	22:V:292:THR:N	2.53	0.58
5:E:156:ARG:HB2	8:H:43:GLY:HA3	1.85	0.58
8:H:92:PRO:HA	8:H:93:LYS:NZ	2.09	0.58
13:M:48:SER:O	13:M:52:ILE:HG22	2.03	0.58
1:A:451:A:H4'	1:A:452:A:O4'	2.03	0.58
1:A:555:U:H2'	1:A:556:C:C6	2.39	0.58

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:1241:G:H2'	1:A:1242:G:C8	2.39	0.58
1:A:1415:G:C2	22:V:285:HIS:HE1	2.22	0.58
1:A:1446:A:H2'	1:A:1447:A:H5''	1.85	0.58
8:H:64:TYR:HA	8:H:70:VAL:HG23	1.85	0.58
8:H:79:ARG:HB2	8:H:80:PRO:HD2	1.84	0.58
20:T:67:HIS:O	20:T:70:LYS:HG2	2.02	0.58
1:A:376:G:H5''	16:P:5:ARG:CB	2.28	0.58
1:A:1035:A:H2'	1:A:1036:A:H8	1.69	0.58
1:A:1116:U:O2'	1:A:1117:A:H5'	2.03	0.58
1:A:1163:A:H2'	1:A:1164:G:H8	1.68	0.58
1:A:1217:C:OP1	14:N:8:ARG:HD3	2.03	0.58
1:A:107:G:O6	20:T:9:ARG:HD3	2.03	0.58
1:A:865:A:H5'	1:A:1078:U:O4	2.04	0.58
1:A:1142:G:H2'	1:A:1143:G:O4'	2.03	0.58
1:A:1163:A:H2'	1:A:1164:G:C8	2.39	0.58
1:A:1338:G:H2'	1:A:1339:A:C8	2.38	0.58
17:Q:17:GLU:C	17:Q:19:SER:H	2.11	0.58
1:A:312:C:H2'	1:A:313:A:H8	1.68	0.58
1:A:1463:U:H2'	1:A:1464:U:C6	2.39	0.58
4:D:24:VAL:HG23	4:D:25:ARG:HD2	1.86	0.58
4:D:196:GLU:HA	4:D:199:ILE:HD12	1.85	0.58
6:F:42:TRP:HB2	6:F:59:TYR:HB2	1.85	0.58
9:I:82:ILE:O	9:I:86:LEU:HD22	2.04	0.58
1:A:940:C:H2'	1:A:941:G:H8	1.69	0.58
2:B:18:GLN:O	2:B:37:VAL:HG23	2.03	0.58
4:D:170:LEU:HB2	4:D:180:THR:O	2.04	0.58
6:F:4:TYR:O	6:F:63:ASN:HA	2.04	0.58
1:A:922:G:N3	1:A:1398:A:H2	2.02	0.57
1:A:1508:A:H2'	1:A:1509:C:C6	2.39	0.57
4:D:169:TRP:NE1	4:D:170:LEU:HD23	2.18	0.57
11:K:52:ARG:HD2	11:K:53:GLY:N	2.19	0.57
12:L:22:ALA:HB2	12:L:56:LEU:HD21	1.85	0.57
14:N:30:ILE:CG2	14:N:41:TRP:HB2	2.34	0.57
22:V:87:VAL:CG1	22:V:88:VAL:N	2.67	0.57
1:A:384:G:H2'	1:A:385:C:C6	2.39	0.57
1:A:560:A:H4'	1:A:561:U:H5''	1.86	0.57
1:A:1077:G:N2	1:A:1079:G:H3'	2.19	0.57
1:A:1172:C:H2'	1:A:1173:U:C6	2.39	0.57
1:A:1284:C:H3'	1:A:1285:A:H5''	1.84	0.57
1:A:1329:A:OP1	13:M:28:ARG:HB2	2.04	0.57
2:B:147:LEU:O	2:B:150:ILE:HG22	2.03	0.57

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
8:H:45:ILE:C	8:H:63:LYS:HZ1	2.12	0.57
9:I:80:HIS:O	9:I:84:ARG:HB2	2.04	0.57
12:L:19:ASN:O	12:L:20:VAL:HG23	2.03	0.57
1:A:57:G:H2'	1:A:58:C:H6	1.67	0.57
1:A:190:A:H8	1:A:190:A:O5'	1.87	0.57
1:A:272:C:H2'	1:A:273:U:H6	1.68	0.57
1:A:1238:A:C2	1:A:1241:G:H1'	2.38	0.57
2:B:119:GLN:O	2:B:124:THR:HG23	2.04	0.57
4:D:137:SER:HB2	4:D:138:PRO:HD2	1.85	0.57
5:E:104:ILE:CD1	5:E:114:LEU:HB2	2.34	0.57
17:Q:46:HIS:HB2	17:Q:70:LYS:HE3	1.85	0.57
1:A:239:U:C5'	1:A:239:U:H6	2.18	0.57
1:A:625:U:H4'	16:P:16:PHE:CE2	2.39	0.57
1:A:672:U:H2'	1:A:673:A:C8	2.39	0.57
1:A:674:G:H2'	1:A:675:A:H8	1.70	0.57
1:A:920:U:H2'	1:A:921:U:C6	2.38	0.57
4:D:25:ARG:HH12	4:D:30:LYS:HG2	1.69	0.57
4:D:111:ALA:O	4:D:114:ARG:HB3	2.04	0.57
6:F:9:MET:HE1	18:R:64:LEU:HA	1.84	0.57
7:G:96:ASN:O	7:G:100:MET:HG3	2.04	0.57
1:A:168:G:O2'	1:A:169:C:H5'	2.04	0.57
1:A:1091:U:H2'	1:A:1093:A:OP2	2.04	0.57
9:I:56:MET:CE	9:I:57:VAL:H	2.17	0.57
9:I:118:ARG:HH22	9:I:122:ARG:HH21	1.52	0.57
14:N:12:ARG:HH21	14:N:58:ARG:HH12	1.53	0.57
1:A:85:U:H4'	1:A:86:G:H4'	1.86	0.57
1:A:160:A:H2'	1:A:161:A:O4'	2.04	0.57
1:A:202:G:H2'	1:A:203:G:C8	2.39	0.57
6:F:17:GLN:HG2	6:F:21:MET:HG3	1.86	0.57
1:A:203:G:H1'	1:A:465:A:H62	1.69	0.57
1:A:270:A:H2'	1:A:271:C:H6	1.68	0.57
1:A:1055:A:H4'	3:C:160:GLU:CG	2.35	0.57
1:A:1201:A:H1'	1:A:1202:U:OP2	2.04	0.57
1:A:1422:G:O2'	1:A:1423:G:H5'	2.05	0.57
5:E:13:LYS:HD2	5:E:112:ALA:HB1	1.85	0.57
9:I:18:VAL:HG22	9:I:64:ILE:HG23	1.85	0.57
16:P:23:ASP:O	16:P:26:ASN:HB2	2.05	0.57
21:U:13:VAL:O	21:U:14:ALA:HB2	2.05	0.57
1:A:796:C:H5'	11:K:128:VAL:HG13	1.86	0.57
1:A:967:C:OP1	1:A:969:A:H5'	2.04	0.57
1:A:1053:G:H4'	1:A:1054:C:H5'	1.87	0.57

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:1416:G:C2	22:V:344:MET:HG3	2.39	0.57
4:D:60:VAL:HB	4:D:194:ILE:HG13	1.86	0.57
7:G:132:THR:HG22	7:G:136:LYS:HG3	1.87	0.57
9:I:51:LEU:HD11	9:I:82:ILE:HG21	1.87	0.57
16:P:18:GLN:NE2	16:P:35:ARG:HD2	2.20	0.57
1:A:91:U:H2'	1:A:92:U:C6	2.40	0.57
1:A:1170:A:H8	1:A:1170:A:O5'	1.88	0.57
1:A:1380:U:O4	7:G:2:ARG:HA	2.04	0.57
5:E:90:GLY:O	5:E:128:GLY:HA3	2.04	0.57
6:F:18:VAL:N	6:F:19:PRO:HD2	2.19	0.57
9:I:118:ARG:NH1	9:I:122:ARG:HE	2.03	0.57
11:K:58:THR:HG23	11:K:61:ALA:HB2	1.85	0.57
1:A:450:G:H4'	16:P:41:PRO:HB2	1.86	0.57
11:K:25:SER:HB3	11:K:28:ASN:O	2.04	0.57
14:N:24:ALA:HB1	14:N:27:LYS:HE3	1.87	0.57
17:Q:75:VAL:HG23	17:Q:76:ARG:H	1.70	0.57
19:S:44:ILE:HA	19:S:61:VAL:HB	1.86	0.57
20:T:34:VAL:HG22	20:T:49:ALA:HB1	1.87	0.57
1:A:72:A:H2'	1:A:73:C:H6	1.70	0.56
1:A:978:A:H5'	1:A:1362:A:N6	2.19	0.56
1:A:1102:A:H2'	1:A:1103:C:C6	2.40	0.56
3:C:90:VAL:CG2	3:C:98:ALA:HB3	2.34	0.56
15:O:60:SER:HA	15:O:63:ARG:HH12	1.69	0.56
1:A:16:A:O2'	1:A:17:U:H5'	2.05	0.56
1:A:390:U:H2'	1:A:391:G:H8	1.66	0.56
1:A:676:A:H1'	11:K:116:PRO:HB3	1.86	0.56
1:A:784:A:C5	22:V:71:ARG:NH1	2.73	0.56
1:A:878:A:H1'	8:H:3:GLN:HE22	1.70	0.56
1:A:987:G:H2'	1:A:988:G:C8	2.40	0.56
1:A:1330:U:C2'	1:A:1331:G:H5'	2.35	0.56
1:A:1418:A:C5	22:V:292:THR:OG1	2.59	0.56
1:A:1522:U:O2'	1:A:1523:G:H5'	2.06	0.56
3:C:39:ARG:HG3	3:C:54:ILE:HG21	1.87	0.56
13:M:22:TYR:HB3	13:M:69:ARG:NH2	2.20	0.56
14:N:9:GLU:HB2	14:N:60:ARG:NH2	2.20	0.56
14:N:30:ILE:C	14:N:32:ASP:H	2.13	0.56
16:P:12:LYS:HD2	16:P:13:LYS:HE3	1.87	0.56
1:A:129:A:H1'	1:A:130:A:C8	2.40	0.56
1:A:177:G:H5''	20:T:59:ARG:HH21	1.70	0.56
1:A:924:C:O2'	1:A:925:G:H5'	2.05	0.56
1:A:1225:A:O5'	13:M:102:LYS:HB3	2.05	0.56

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:1294:G:H2'	1:A:1295:U:C6	2.40	0.56
3:C:71:ARG:HB3	3:C:74:ILE:HG22	1.87	0.56
7:G:114:SER:C	7:G:118:ARG:HG3	2.30	0.56
11:K:61:ALA:O	11:K:64:VAL:HG12	2.06	0.56
11:K:88:PRO:HD3	21:U:28:LEU:CD1	2.34	0.56
11:K:113:THR:CG2	21:U:28:LEU:HD21	2.36	0.56
11:K:126:ARG:HB2	21:U:33:ARG:CD	2.32	0.56
1:A:810:C:O2'	1:A:811:C:H5'	2.06	0.56
1:A:883:C:O2'	1:A:884:U:H5'	2.05	0.56
1:A:1483:A:H5'	22:V:334:PHE:O	2.05	0.56
2:B:125:PHE:HD1	2:B:125:PHE:H	1.53	0.56
4:D:98:ASP:CG	4:D:132:ALA:HB1	2.30	0.56
10:J:30:LYS:HB2	10:J:36:VAL:HG21	1.85	0.56
10:J:67:ILE:HG13	14:N:95:LEU:HD13	1.87	0.56
11:K:32:THR:HA	11:K:43:TRP:HA	1.88	0.56
11:K:95:THR:HG23	11:K:96:ILE:N	2.20	0.56
14:N:30:ILE:HG21	14:N:44:VAL:HG21	1.88	0.56
1:A:176:C:H2'	1:A:177:G:N3	2.20	0.56
1:A:678:U:H2'	1:A:679:C:H6	1.70	0.56
1:A:1246:A:H2'	1:A:1247:U:C6	2.41	0.56
1:A:1479:C:H2'	1:A:1480:A:H8	1.71	0.56
1:A:1490:U:H5'	1:A:1491:G:OP2	2.05	0.56
7:G:4:ARG:NE	7:G:5:VAL:H	2.03	0.56
7:G:13:PRO:HB2	7:G:18:GLY:CA	2.35	0.56
7:G:72:VAL:HA	7:G:89:GLU:HA	1.86	0.56
13:M:95:PRO:HD3	13:M:108:ARG:HG2	1.87	0.56
1:A:335:C:H2'	1:A:336:A:C8	2.41	0.56
1:A:629:A:H2'	1:A:630:A:O4'	2.06	0.56
1:A:785:G:C5'	22:V:61:ASP:CG	2.78	0.56
1:A:803:G:H2'	1:A:804:U:C6	2.40	0.56
1:A:844:G:N2	1:A:845:A:H62	2.03	0.56
1:A:973:G:H1'	10:J:56:HIS:ND1	2.20	0.56
1:A:1115:U:H2'	1:A:1116:U:C6	2.41	0.56
2:B:216:VAL:O	2:B:220:VAL:HG23	2.06	0.56
7:G:4:ARG:CZ	7:G:5:VAL:H	2.18	0.56
7:G:85:GLN:O	7:G:86:VAL:C	2.49	0.56
1:A:86:G:H4'	1:A:86:G:OP1	2.05	0.56
1:A:769:G:H4'	1:A:1513:A:H4'	1.87	0.56
1:A:1008:U:H2'	1:A:1009:U:H4'	1.88	0.56
2:B:38:HIS:O	2:B:39:ILE:HD13	2.06	0.56
3:C:46:LEU:HD23	3:C:75:VAL:HG13	1.86	0.56

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
4:D:117:VAL:O	4:D:130:ASN:HA	2.05	0.56
11:K:97:ARG:HB3	11:K:97:ARG:NH1	2.21	0.56
12:L:8:ARG:HG3	12:L:9:LYS:N	2.17	0.56
13:M:2:ARG:HB3	13:M:6:ILE:O	2.06	0.56
14:N:66:THR:HG23	14:N:67:GLY:N	2.20	0.56
16:P:20:VAL:HG22	16:P:21:VAL:N	2.21	0.56
21:U:14:ALA:H	21:U:16:ARG:CZ	2.15	0.56
1:A:372:C:H4'	1:A:373:A:H5'	1.86	0.56
1:A:555:U:H2'	1:A:556:C:H6	1.71	0.56
1:A:764:C:H2'	1:A:765:G:H5'	1.87	0.56
1:A:1027:C:H2'	1:A:1028:C:H6	1.70	0.56
1:A:1280:A:O4'	10:J:43:PRO:HG3	2.06	0.56
1:A:1420:U:O2'	22:V:315:ASP:O	2.07	0.56
2:B:144:GLU:O	2:B:148:GLY:HA3	2.06	0.56
4:D:32:LYS:O	4:D:35:GLN:HB2	2.05	0.56
6:F:100:SER:HA	18:R:23:LYS:NZ	2.21	0.56
14:N:42:ASN:O	14:N:46:LYS:HG3	2.05	0.56
17:Q:45:VAL:HG12	17:Q:46:HIS:H	1.70	0.56
1:A:332:G:OP2	20:T:2:ASN:HB3	2.06	0.56
1:A:449:G:H2'	1:A:450:G:C8	2.40	0.56
1:A:1392:G:H2'	1:A:1393:U:C6	2.41	0.56
1:A:1418:A:C4	22:V:292:THR:CG2	2.62	0.56
3:C:154:GLY:HA2	3:C:163:ARG:O	2.06	0.56
6:F:3:HIS:HB2	6:F:92:THR:CB	2.35	0.56
6:F:54:LEU:CD1	6:F:55:HIS:H	2.18	0.56
9:I:23:GLY:O	9:I:24:ASN:HB2	2.06	0.56
11:K:58:THR:HG23	11:K:61:ALA:CB	2.36	0.56
1:A:313:A:H2'	1:A:314:C:C6	2.40	0.56
1:A:985:C:H2'	1:A:986:U:C6	2.41	0.56
1:A:1135:U:O2	1:A:1135:U:H2'	2.06	0.56
1:A:1492:A:H2'	1:A:1493:A:O4'	2.06	0.56
9:I:50:PRO:HG2	9:I:51:LEU:HD12	1.88	0.56
12:L:3:VAL:O	12:L:7:VAL:HG23	2.06	0.56
12:L:83:GLY:HA2	12:L:94:TYR:HD1	1.70	0.56
12:L:105:GLY:HA3	12:L:117:GLY:HA3	1.88	0.56
14:N:65:GLN:HB2	14:N:78:LEU:HD22	1.87	0.56
17:Q:10:ARG:CZ	17:Q:56:ASP:H	2.19	0.56
1:A:218:U:H2'	1:A:219:U:C6	2.41	0.55
1:A:389:A:H3'	1:A:390:U:H6	1.71	0.55
1:A:408:A:H5'	4:D:21:LYS:HE2	1.88	0.55
1:A:1320:C:N4	19:S:36:ARG:HG2	2.21	0.55

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:1421:G:OP1	22:V:316:PRO:O	2.25	0.55
2:B:10:LYS:C	2:B:12:GLY:H	2.14	0.55
4:D:99:ASN:HD21	4:D:110:ARG:HE	1.52	0.55
9:I:30:ASN:HD22	9:I:30:ASN:N	2.04	0.55
19:S:32:THR:HB	19:S:49:ALA:O	2.06	0.55
21:U:38:GLU:C	21:U:40:PRO:HD2	2.31	0.55
1:A:32:A:H2'	1:A:33:A:C8	2.42	0.55
1:A:521:G:OP2	12:L:50:LYS:HE3	2.06	0.55
1:A:1129:C:H5''	9:I:17:ARG:HH22	1.70	0.55
1:A:1244:G:H2'	1:A:1245:C:C6	2.41	0.55
4:D:117:VAL:HG12	4:D:130:ASN:HA	1.88	0.55
5:E:82:HIS:HB2	5:E:83:PRO:HD2	1.88	0.55
7:G:19:SER:OG	7:G:22:LEU:HB2	2.06	0.55
11:K:92:ARG:HB3	11:K:92:ARG:NH1	2.19	0.55
12:L:37:TYR:HB2	12:L:51:VAL:HG23	1.88	0.55
14:N:15:LEU:HD12	14:N:16:ALA:H	1.71	0.55
14:N:50:LEU:CD2	14:N:51:PRO:HD3	2.36	0.55
16:P:41:PRO:O	16:P:42:ILE:HD13	2.05	0.55
19:S:59:VAL:HG11	19:S:70:LEU:HD11	1.88	0.55
1:A:764:C:C2'	1:A:765:G:H5'	2.37	0.55
1:A:1171:A:H2'	1:A:1172:C:C6	2.40	0.55
1:A:1407:C:C5'	22:V:284:TRP:CB	2.84	0.55
8:H:26:MET:HB2	8:H:27:PRO:HD2	1.86	0.55
9:I:51:LEU:HD11	9:I:82:ILE:HD13	1.87	0.55
12:L:30:ARG:CB	12:L:30:ARG:HH11	2.19	0.55
14:N:27:LYS:HD2	14:N:27:LYS:C	2.30	0.55
17:Q:74:LEU:HD22	17:Q:75:VAL:N	2.21	0.55
18:R:23:LYS:C	18:R:25:ILE:H	2.14	0.55
1:A:940:C:H2'	1:A:941:G:C8	2.40	0.55
1:A:1324:A:H2'	1:A:1325:C:H6	1.72	0.55
1:A:1416:G:C4	22:V:344:MET:HB3	2.42	0.55
21:U:36:PHE:CB	21:U:40:PRO:HD3	2.35	0.55
1:A:163:C:H2'	1:A:164:G:O4'	2.06	0.55
1:A:278:G:H21	1:A:279:A:N6	2.04	0.55
1:A:1111:A:O2'	1:A:1112:C:H6	1.90	0.55
1:A:1243:C:H2'	1:A:1244:G:C8	2.41	0.55
3:C:166:TRP:HA	3:C:166:TRP:CE3	2.42	0.55
7:G:45:ALA:HB1	7:G:120:ALA:HB2	1.88	0.55
7:G:67:ASN:ND2	7:G:127:ALA:HA	2.21	0.55
10:J:55:PRO:HA	14:N:80:ARG:NH2	2.21	0.55
1:A:372:C:H1'	1:A:373:A:OP2	2.06	0.55

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:921:U:H2'	1:A:922:G:O4'	2.06	0.55
1:A:1512:U:O2'	1:A:1513:A:H5'	2.07	0.55
5:E:45:VAL:CG1	5:E:116:VAL:HG23	2.36	0.55
9:I:61:ASP:O	9:I:62:LEU:HD13	2.05	0.55
21:U:14:ALA:N	21:U:16:ARG:NH2	2.55	0.55
1:A:6:G:H4'	1:A:298:A:H4'	1.88	0.55
1:A:1409:C:OP2	22:V:283:LEU:HD23	2.07	0.55
2:B:138:ARG:HA	2:B:141:GLU:OE2	2.06	0.55
10:J:11:LYS:NZ	10:J:99:GLN:H	2.05	0.55
13:M:106:ARG:NH1	13:M:109:LYS:HD2	2.21	0.55
15:O:54:GLY:O	15:O:58:MET:HG2	2.07	0.55
1:A:398:U:H2'	1:A:399:G:C8	2.42	0.55
1:A:1112:C:O2	3:C:178:ARG:N	2.40	0.55
2:B:67:LEU:HD21	2:B:157:PRO:CG	2.37	0.55
3:C:126:ARG:HH22	3:C:190:THR:HG22	1.72	0.55
6:F:64:VAL:HG12	6:F:65:GLU:N	2.17	0.55
7:G:53:SER:HB2	7:G:55:LYS:HZ2	1.72	0.55
13:M:13:HIS:HB2	13:M:16:ILE:CG2	2.37	0.55
1:A:370:C:H2'	1:A:371:A:C8	2.42	0.55
1:A:801:U:H2'	1:A:802:A:C8	2.42	0.55
1:A:845:A:H3'	1:A:846:G:O4'	2.06	0.55
1:A:859:G:H2'	1:A:860:A:H8	1.71	0.55
3:C:53:ARG:HG2	3:C:54:ILE:H	1.72	0.55
3:C:151:GLU:HB2	3:C:200:TRP:HZ3	1.71	0.55
5:E:132:PRO:HA	5:E:135:VAL:HG22	1.88	0.55
16:P:12:LYS:HD2	16:P:13:LYS:HG3	1.89	0.55
19:S:49:ALA:HB1	19:S:56:HIS:CB	2.30	0.55
1:A:393:A:H5'	1:A:483:C:O2'	2.06	0.55
1:A:1160:G:H2'	1:A:1161:C:H6	1.72	0.55
1:A:1332:A:H2'	1:A:1333:A:C8	2.42	0.55
1:A:1448:C:H2'	1:A:1449:C:C6	2.42	0.55
3:C:42:LEU:O	3:C:46:LEU:HB2	2.07	0.55
3:C:133:MET:HE2	3:C:152:VAL:CG2	2.37	0.55
6:F:61:LEU:HD12	6:F:63:ASN:OD1	2.07	0.55
17:Q:16:MET:CB	17:Q:19:SER:HB2	2.35	0.55
1:A:97:G:H2'	1:A:98:A:O4'	2.07	0.54
1:A:716:A:N3	11:K:119:GLY:HA2	2.22	0.54
1:A:736:C:H4'	6:F:88:MET:HE3	1.89	0.54
1:A:973:G:H3'	1:A:974:A:H5''	1.89	0.54
1:A:1027:C:H2'	1:A:1028:C:C6	2.41	0.54
1:A:1420:U:C4	22:V:311:LYS:CG	2.83	0.54

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:B:112:ARG:HA	2:B:115:ASP:OD2	2.07	0.54
3:C:21:TRP:CB	3:C:58:ARG:HB2	2.37	0.54
4:D:25:ARG:HH11	4:D:30:LYS:HE3	1.70	0.54
9:I:38:PHE:O	9:I:44:ARG:HG2	2.07	0.54
11:K:110:THR:CG2	21:U:4:LYS:HA	2.37	0.54
14:N:17:ASP:HA	14:N:21:ALA:CB	2.37	0.54
16:P:51:ARG:C	16:P:52:LEU:HD22	2.31	0.54
17:Q:44:HIS:O	17:Q:72:TRP:HB2	2.07	0.54
1:A:37:U:OP1	12:L:119:LYS:HE3	2.07	0.54
1:A:930:C:H2'	1:A:931:C:O4'	2.08	0.54
1:A:1250:A:H4'	9:I:69:GLY:N	2.22	0.54
1:A:1427:C:O2'	1:A:1428:A:H5'	2.07	0.54
2:B:113:LEU:HG	2:B:143:LEU:HB3	1.89	0.54
7:G:115:MET:HE3	7:G:119:LEU:HB2	1.90	0.54
8:H:9:MET:HA	8:H:26:MET:HE2	1.89	0.54
10:J:24:GLU:OE2	10:J:90:LEU:HD22	2.06	0.54
10:J:53:ILE:HG23	14:N:84:ARG:CZ	2.37	0.54
14:N:81:ILE:HD12	14:N:82:LYS:H	1.72	0.54
22:V:301:TYR:CE2	22:V:329:ILE:HD11	2.43	0.54
1:A:173:U:H5'	1:A:197:A:O4'	2.07	0.54
1:A:189:A:H2'	1:A:190:A:C8	2.43	0.54
1:A:1363:A:H2'	1:A:1363:A:N3	2.22	0.54
3:C:78:LYS:CG	3:C:81:GLU:HB2	2.24	0.54
8:H:50:VAL:HG22	8:H:51:GLU:N	2.22	0.54
9:I:113:LYS:HG2	9:I:114:LYS:N	2.22	0.54
13:M:64:VAL:HG12	13:M:65:GLU:H	1.71	0.54
15:O:84:LEU:HD22	15:O:86:LEU:HD11	1.89	0.54
22:V:133:GLN:O	22:V:218:SER:HA	2.08	0.54
22:V:140:ILE:HD11	22:V:167:LEU:HD11	1.89	0.54
1:A:532:A:H62	3:C:191:THR:HB	1.72	0.54
1:A:1346:A:N1	1:A:1374:A:H5''	2.22	0.54
1:A:1491:G:H5''	1:A:1492:A:OP2	2.07	0.54
2:B:67:LEU:HA	2:B:89:PHE:O	2.07	0.54
4:D:144:ILE:CD1	4:D:154:VAL:HG21	2.38	0.54
6:F:18:VAL:O	6:F:22:ILE:HG13	2.07	0.54
8:H:92:PRO:CA	8:H:93:LYS:HZ2	2.10	0.54
9:I:49:GLN:NE2	9:I:79:ARG:HD2	2.22	0.54
15:O:70:LYS:NZ	15:O:74:VAL:HG13	2.22	0.54
1:A:477:C:H2'	1:A:478:A:C8	2.42	0.54
1:A:636:U:H2'	1:A:637:C:C6	2.43	0.54
1:A:960:U:O2'	1:A:1223:C:H4'	2.08	0.54

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:1035:A:H2'	1:A:1036:A:C8	2.43	0.54
2:B:46:VAL:O	2:B:49:PHE:HB2	2.07	0.54
2:B:67:LEU:HD11	2:B:157:PRO:CB	2.30	0.54
5:E:134:ASN:O	5:E:137:ARG:HG2	2.07	0.54
17:Q:45:VAL:HG11	17:Q:74:LEU:HB2	1.89	0.54
17:Q:64:ARG:HG3	17:Q:65:PRO:HD2	1.89	0.54
19:S:20:LYS:O	19:S:23:GLU:HB3	2.07	0.54
19:S:29:PRO:HA	19:S:47:THR:HB	1.89	0.54
21:U:39:LYS:N	21:U:40:PRO:CD	2.69	0.54
22:V:73:ASN:ND2	22:V:102:LYS:HE3	2.22	0.54
1:A:602:A:O2'	1:A:603:U:H5'	2.07	0.54
1:A:1418:A:C2	22:V:312:HIS:NE2	2.75	0.54
1:A:1483:A:N6	22:V:288:PRO:C	2.51	0.54
7:G:100:MET:O	7:G:104:VAL:HG23	2.08	0.54
11:K:15:VAL:HG21	11:K:41:LEU:HD11	1.88	0.54
11:K:54:SER:HA	11:K:56:LYS:HE3	1.89	0.54
11:K:113:THR:HG21	21:U:28:LEU:HD21	1.88	0.54
16:P:40:ASN:HB3	16:P:49:GLY:O	2.08	0.54
20:T:82:ILE:HG13	20:T:83:ASN:N	2.22	0.54
1:A:766:A:H2'	1:A:767:A:O4'	2.08	0.54
1:A:1417:G:C4'	22:V:289:GLU:OE1	2.47	0.54
1:A:1435:G:H2'	1:A:1436:U:C6	2.43	0.54
3:C:156:LEU:HD12	3:C:156:LEU:H	1.72	0.54
6:F:46:GLN:HA	6:F:56:LYS:HG3	1.89	0.54
12:L:20:VAL:O	12:L:23:LEU:HG	2.08	0.54
19:S:69:LYS:O	19:S:72:GLU:HB2	2.07	0.54
1:A:1100:C:O2'	1:A:1101:A:H5'	2.07	0.54
1:A:1310:G:H2'	1:A:1311:A:O4'	2.07	0.54
1:A:1473:G:H2'	1:A:1474:U:O4'	2.07	0.54
1:A:1508:A:H2'	1:A:1509:C:H6	1.73	0.54
4:D:17:ASP:OD2	4:D:27:ILE:HG12	2.07	0.54
5:E:56:PRO:O	5:E:60:GLN:HG3	2.07	0.54
5:E:95:MET:HE2	5:E:114:LEU:HD11	1.90	0.54
6:F:53:LYS:CD	6:F:54:LEU:H	2.11	0.54
17:Q:32:ILE:HG13	17:Q:33:TYR:N	2.21	0.54
18:R:33:THR:HG23	18:R:35:SER:H	1.73	0.54
1:A:190:A:H2'	1:A:191:G:O4'	2.07	0.54
1:A:415:A:H3'	1:A:416:G:H8	1.72	0.54
1:A:1101:A:N6	2:B:173:LYS:HD2	2.22	0.54
1:A:1381:U:O2'	1:A:1382:C:H5'	2.08	0.54
1:A:1485:U:C5'	22:V:286:LEU:CD1	2.84	0.54

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
5:E:95:MET:HE3	5:E:139:THR:HG22	1.89	0.54
12:L:58:ASN:H	12:L:58:ASN:ND2	2.05	0.54
16:P:42:ILE:HB	16:P:46:LYS:CD	2.36	0.54
1:A:621:A:H2'	1:A:622:A:C8	2.43	0.54
1:A:638:U:H2'	1:A:639:G:O4'	2.08	0.54
1:A:1040:U:H2'	1:A:1041:G:C8	2.43	0.54
1:A:1103:C:C5'	2:B:96:LEU:HD12	2.37	0.54
1:A:1407:C:H5''	22:V:284:TRP:HB2	1.89	0.54
1:A:1419:G:H5'	22:V:309:ASP:N	2.19	0.54
1:A:1420:U:H5''	22:V:307:TYR:O	2.07	0.54
1:A:1488:G:O2'	1:A:1489:G:H5'	2.09	0.54
3:C:6:PRO:HG3	3:C:200:TRP:HE1	1.73	0.54
3:C:156:LEU:HD11	3:C:163:ARG:O	2.08	0.54
8:H:29:SER:CB	8:H:32:LYS:HZ2	2.20	0.54
9:I:43:ALA:C	9:I:45:MET:H	2.16	0.54
9:I:44:ARG:N	9:I:44:ARG:HH11	2.06	0.54
10:J:7:ARG:HA	10:J:75:ASP:HB2	1.90	0.54
14:N:42:ASN:HB3	14:N:46:LYS:HE2	1.89	0.54
17:Q:18:LYS:HA	17:Q:47:ASP:O	2.08	0.54
20:T:74:HIS:O	20:T:78:LEU:HD12	2.08	0.54
21:U:37:TYR:C	21:U:40:PRO:HD2	2.33	0.54
22:V:68:GLU:HG3	22:V:70:HIS:HE1	1.73	0.54
22:V:323:ALA:CB	22:V:329:ILE:HD13	2.34	0.54
22:V:328:ALA:C	22:V:329:ILE:HD12	2.33	0.54
1:A:785:G:C3'	22:V:69:VAL:CG1	2.74	0.53
1:A:1039:G:H2'	1:A:1040:U:C6	2.43	0.53
1:A:1048:G:H4'	14:N:2:LYS:HZ2	1.73	0.53
4:D:138:PRO:HA	4:D:181:PHE:CD2	2.43	0.53
11:K:33:ILE:O	11:K:41:LEU:HB2	2.07	0.53
11:K:88:PRO:HA	11:K:92:ARG:NE	2.23	0.53
12:L:86:VAL:HG11	12:L:89:LEU:HD23	1.90	0.53
21:U:29:ALA:HB1	21:U:32:ARG:NH2	2.10	0.53
1:A:77:A:H2'	1:A:78:A:C8	2.43	0.53
1:A:187:G:H4'	20:T:79:THR:HG21	1.89	0.53
1:A:532:A:H62	3:C:191:THR:CG2	2.22	0.53
1:A:865:A:H2'	1:A:866:C:C6	2.43	0.53
1:A:1117:A:H5''	9:I:105:ARG:HH12	1.73	0.53
1:A:1307:U:H2'	1:A:1308:U:H6	1.72	0.53
1:A:1484:C:H3'	22:V:340:ILE:HG21	1.89	0.53
2:B:148:GLY:HA2	2:B:151:LYS:HD3	1.89	0.53
2:B:185:ILE:HA	2:B:199:ILE:HB	1.90	0.53

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:B:185:ILE:HG22	2:B:199:ILE:HB	1.91	0.53
3:C:38:VAL:HG23	3:C:39:ARG:N	2.23	0.53
3:C:186:SER:HB3	3:C:197:VAL:CG1	2.38	0.53
5:E:81:GLN:HE21	5:E:147:ASN:C	2.17	0.53
5:E:150:GLU:C	5:E:152:VAL:H	2.16	0.53
12:L:56:LEU:HB3	12:L:58:ASN:ND2	2.24	0.53
1:A:513:C:H2'	1:A:514:C:H6	1.73	0.53
1:A:626:G:H2'	1:A:627:G:H8	1.73	0.53
1:A:784:A:C5'	22:V:72:CYS:O	2.56	0.53
1:A:784:A:H2'	1:A:785:G:H8	1.69	0.53
1:A:1283:U:H2'	1:A:1284:C:C6	2.44	0.53
7:G:110:ARG:HH22	7:G:121:ASN:HB3	1.72	0.53
9:I:11:ARG:HA	9:I:105:ARG:NE	2.23	0.53
9:I:24:ASN:O	9:I:61:ASP:HA	2.09	0.53
19:S:4:LEU:CD1	19:S:9:PHE:H	2.17	0.53
21:U:36:PHE:HB3	21:U:40:PRO:HG3	1.90	0.53
1:A:275:G:C5'	17:Q:15:LYS:HG2	2.39	0.53
2:B:71:THR:HG23	2:B:93:HIS:C	2.33	0.53
5:E:55:VAL:N	5:E:56:PRO:HD2	2.23	0.53
8:H:5:PRO:O	8:H:32:LYS:HE3	2.09	0.53
1:A:252:U:H2'	1:A:253:A:H8	1.74	0.53
1:A:499:A:H4'	1:A:500:G:OP1	2.09	0.53
1:A:846:G:H2'	1:A:846:G:N3	2.23	0.53
1:A:1415:G:H3'	22:V:345:ALA:O	2.08	0.53
1:A:1507:A:H2'	1:A:1508:A:C8	2.44	0.53
3:C:39:ARG:HG3	3:C:54:ILE:HD13	1.91	0.53
3:C:111:ASP:O	3:C:115:VAL:HG23	2.08	0.53
6:F:47:LEU:HD13	6:F:51:ILE:HG22	1.91	0.53
8:H:43:GLY:O	8:H:63:LYS:HE3	2.09	0.53
9:I:38:PHE:O	9:I:41:GLU:HB2	2.09	0.53
9:I:113:LYS:HG2	9:I:114:LYS:H	1.74	0.53
12:L:20:VAL:O	12:L:20:VAL:HG12	2.08	0.53
16:P:51:ARG:HB3	16:P:51:ARG:HH11	1.71	0.53
22:V:131:ILE:HD12	22:V:219:ILE:HG23	1.90	0.53
1:A:214:C:H2'	1:A:215:C:C6	2.44	0.53
1:A:664:G:N2	1:A:741:G:H1	2.02	0.53
1:A:860:A:H2'	1:A:861:G:O4'	2.09	0.53
1:A:1409:C:N4	1:A:1491:G:N1	2.54	0.53
1:A:1420:U:H5''	22:V:307:TYR:H	1.72	0.53
11:K:110:THR:HG22	21:U:4:LYS:HA	1.90	0.53
14:N:81:ILE:O	14:N:84:ARG:HB3	2.09	0.53

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
18:R:44:THR:HG22	18:R:46:THR:HB	1.90	0.53
19:S:18:VAL:CG2	19:S:43:MET:HG2	2.32	0.53
1:A:399:G:H2'	1:A:400:C:C6	2.43	0.53
2:B:66:ILE:C	2:B:67:LEU:HD13	2.34	0.53
2:B:96:LEU:HB2	2:B:99:MET:HE3	1.91	0.53
3:C:77:GLY:HA3	3:C:81:GLU:HB3	1.90	0.53
7:G:16:LYS:C	7:G:16:LYS:HD3	2.34	0.53
18:R:25:ILE:HG13	18:R:67:LEU:HD11	1.91	0.53
20:T:53:MET:O	20:T:57:VAL:HG23	2.08	0.53
1:A:783:C:H5''	22:V:58:MET:CB	2.35	0.53
1:A:1042:A:H2'	1:A:1043:G:O4'	2.08	0.53
1:A:1132:C:H2'	1:A:1133:G:O4'	2.09	0.53
2:B:87:ASP:O	2:B:88:GLN:HG3	2.07	0.53
5:E:71:ILE:HG12	5:E:72:ASN:N	2.24	0.53
5:E:74:ALA:HB1	5:E:148:SER:HB3	1.91	0.53
7:G:45:ALA:HA	7:G:48:THR:OG1	2.08	0.53
7:G:112:ASP:CB	7:G:118:ARG:HG2	2.33	0.53
10:J:81:GLU:HA	10:J:84:VAL:HG22	1.89	0.53
17:Q:4:ILE:HD12	17:Q:5:ARG:H	1.73	0.53
1:A:195:A:H1'	1:A:222:C:O2'	2.09	0.53
1:A:824:G:O2'	1:A:825:A:H5'	2.08	0.53
1:A:1005:A:H3'	1:A:1006:G:H8	1.74	0.53
1:A:1071:C:H2'	1:A:1072:G:C8	2.42	0.53
1:A:1420:U:H5''	22:V:307:TYR:CA	2.39	0.53
1:A:1439:G:H2'	1:A:1440:U:O4'	2.08	0.53
11:K:51:PHE:CE1	11:K:61:ALA:HB1	2.43	0.53
11:K:69:CYS:C	11:K:71:ASP:N	2.66	0.53
17:Q:80:LYS:O	17:Q:80:LYS:HE3	2.09	0.53
22:V:134:ILE:HD11	22:V:136:ILE:HG12	1.90	0.53
1:A:403:C:O2'	1:A:404:G:H5'	2.08	0.53
1:A:411:A:O2'	1:A:412:A:H4'	2.09	0.53
1:A:423:G:H2'	1:A:424:G:O4'	2.09	0.53
1:A:1170:A:H2'	1:A:1171:A:O4'	2.09	0.53
1:A:1221:G:H5''	19:S:35:ARG:NH1	2.23	0.53
1:A:1483:A:C4'	22:V:298:PHE:HZ	2.19	0.53
3:C:55:VAL:HG12	3:C:56:ILE:N	2.23	0.53
8:H:76:ARG:HH11	8:H:76:ARG:HG3	1.74	0.53
16:P:57:ILE:O	16:P:61:VAL:HG22	2.09	0.53
1:A:128:G:H2'	1:A:129:A:C8	2.44	0.52
1:A:196:A:O5'	20:T:63:LYS:HE2	2.09	0.52
1:A:520:A:OP2	12:L:47:ALA:HB1	2.09	0.52

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:742:G:O2'	1:A:743:A:H5'	2.08	0.52
1:A:1485:U:C6	22:V:291:ILE:HD12	2.43	0.52
1:A:1521:C:O2'	1:A:1522:U:H5'	2.10	0.52
2:B:113:LEU:HD23	2:B:113:LEU:C	2.33	0.52
5:E:149:PRO:HA	8:H:98:LEU:HD21	1.91	0.52
6:F:40:GLU:CG	6:F:42:TRP:HE1	2.22	0.52
6:F:43:GLY:O	6:F:58:HIS:HA	2.09	0.52
7:G:91:ARG:HB3	7:G:92:PRO:HD2	1.91	0.52
9:I:30:ASN:ND2	9:I:30:ASN:N	2.56	0.52
9:I:90:ASP:O	9:I:93:LEU:HG	2.09	0.52
10:J:88:MET:C	10:J:90:LEU:H	2.17	0.52
15:O:2:LEU:HD23	15:O:3:SER:N	2.24	0.52
15:O:15:GLY:HA3	15:O:20:ASP:OD1	2.09	0.52
1:A:202:G:H2'	1:A:203:G:H8	1.74	0.52
1:A:997:U:O2'	1:A:998:C:H5'	2.09	0.52
2:B:23:ASN:C	2:B:23:ASN:HD22	2.16	0.52
2:B:67:LEU:HD12	2:B:153:MET:CE	2.39	0.52
3:C:135:ARG:C	3:C:137:VAL:H	2.17	0.52
3:C:155:ARG:N	3:C:162:ALA:HA	2.21	0.52
4:D:54:LEU:O	4:D:54:LEU:HD22	2.09	0.52
4:D:58:GLN:OE1	4:D:62:ARG:HG3	2.09	0.52
4:D:107:GLY:HA2	4:D:112:GLU:OE1	2.10	0.52
7:G:16:LYS:HG3	7:G:43:TYR:OH	2.08	0.52
8:H:49:LYS:HG3	8:H:50:VAL:N	2.21	0.52
12:L:35:ARG:CZ	12:L:75:GLU:HB3	2.39	0.52
1:A:53:A:C2	1:A:54:C:H1'	2.44	0.52
1:A:746:A:H2'	1:A:747:A:C8	2.44	0.52
1:A:956:U:O2'	1:A:957:U:H5'	2.09	0.52
3:C:146:LYS:HG3	3:C:202:PHE:HD2	1.74	0.52
11:K:111:ASP:O	18:R:72:ARG:HD3	2.09	0.52
15:O:66:LEU:HB3	15:O:77:TYR:HE1	1.74	0.52
17:Q:4:ILE:HD12	17:Q:4:ILE:H	1.74	0.52
1:A:405:U:O4	4:D:1:ALA:HA	2.09	0.52
1:A:600:A:H2'	1:A:601:G:H8	1.74	0.52
1:A:610:U:O2	1:A:610:U:O4'	2.28	0.52
1:A:783:C:H5''	22:V:58:MET:H	1.75	0.52
1:A:889:A:H61	1:A:907:A:H3'	1.74	0.52
1:A:1060:U:H5''	10:J:53:ILE:CG2	2.40	0.52
1:A:1308:U:OP1	13:M:95:PRO:HA	2.09	0.52
1:A:1409:C:H4'	22:V:151:ARG:HE	1.73	0.52
1:A:1484:C:H2'	22:V:291:ILE:CD1	2.29	0.52

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
4:D:186:GLU:O	4:D:190:LEU:HD13	2.09	0.52
5:E:109:ALA:CB	5:E:135:VAL:HG23	2.39	0.52
12:L:90:PRO:C	12:L:92:VAL:H	2.18	0.52
13:M:106:ARG:HD3	13:M:110:GLY:O	2.10	0.52
17:Q:60:ILE:HD13	17:Q:60:ILE:N	2.25	0.52
20:T:60:GLN:CD	20:T:60:GLN:N	2.67	0.52
1:A:317:U:H2'	1:A:318:G:H8	1.74	0.52
1:A:1130:A:H5'	9:I:19:PHE:CE2	2.44	0.52
1:A:1452:C:H4'	1:A:1453:G:C5'	2.39	0.52
2:B:68:PHE:HA	2:B:161:PHE:O	2.10	0.52
4:D:25:ARG:HD3	4:D:26:ALA:H	1.73	0.52
4:D:122:ILE:O	4:D:128:VAL:HG23	2.10	0.52
7:G:30:MET:SD	7:G:35:LYS:HB2	2.50	0.52
11:K:28:ASN:ND2	11:K:29:THR:N	2.58	0.52
11:K:70:ALA:HB1	11:K:74:LYS:HE3	1.89	0.52
12:L:30:ARG:HH11	12:L:30:ARG:HB3	1.75	0.52
15:O:70:LYS:HZ1	15:O:74:VAL:HA	1.75	0.52
22:V:263:LEU:HD12	22:V:263:LEU:C	2.33	0.52
1:A:36:C:H2'	1:A:37:U:O4'	2.09	0.52
1:A:279:A:H4'	1:A:280:C:OP2	2.10	0.52
1:A:474:G:H2'	1:A:475:C:C6	2.45	0.52
1:A:712:A:O2'	1:A:713:G:H5'	2.10	0.52
1:A:1376:U:H2'	1:A:1377:A:H8	1.75	0.52
1:A:1420:U:C5'	22:V:307:TYR:CB	2.80	0.52
2:B:53:LEU:HA	2:B:56:LEU:HD23	1.92	0.52
5:E:14:LEU:HA	5:E:36:THR:HG22	1.91	0.52
6:F:100:SER:HA	18:R:23:LYS:HD2	1.91	0.52
10:J:22:THR:HG23	10:J:23:ALA:N	2.24	0.52
13:M:44:ILE:HD12	13:M:45:SER:N	2.23	0.52
1:A:1040:U:H2'	1:A:1041:G:H8	1.74	0.52
1:A:1309:G:N7	13:M:97:ARG:NH2	2.57	0.52
1:A:1420:U:H5''	22:V:307:TYR:CB	2.39	0.52
1:A:1499:A:O2'	1:A:1500:A:H5'	2.09	0.52
2:B:76:SER:OG	2:B:92:ASN:HB2	2.10	0.52
6:F:6:ILE:HD13	6:F:89:VAL:HB	1.89	0.52
7:G:142:ARG:C	7:G:146:ALA:HB3	2.34	0.52
10:J:39:PRO:CA	10:J:74:VAL:HG22	2.39	0.52
11:K:36:ARG:HG2	11:K:37:GLN:H	1.74	0.52
13:M:18:LEU:O	13:M:21:ILE:HG13	2.10	0.52
18:R:62:ARG:HD3	18:R:69:TYR:HA	1.90	0.52
19:S:29:PRO:CA	19:S:47:THR:HB	2.39	0.52

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:263:A:P	20:T:73:ARG:HH22	2.32	0.52
1:A:320:A:H2'	1:A:321:A:C8	2.44	0.52
1:A:408:A:H3'	1:A:409:U:H6	1.74	0.52
1:A:472:U:H2'	1:A:473:U:C6	2.44	0.52
2:B:49:PHE:O	2:B:53:LEU:HD23	2.10	0.52
2:B:96:LEU:H	2:B:99:MET:HE3	1.73	0.52
2:B:101:THR:HG23	2:B:102:ASN:H	1.75	0.52
4:D:117:VAL:HG12	4:D:130:ASN:C	2.35	0.52
5:E:40:ASP:C	5:E:42:ASN:H	2.17	0.52
7:G:14:ASP:OD1	7:G:22:LEU:HD12	2.10	0.52
7:G:143:MET:O	7:G:147:ASN:HB2	2.09	0.52
9:I:56:MET:SD	9:I:57:VAL:N	2.82	0.52
11:K:81:LEU:HD22	11:K:104:PHE:HB3	1.92	0.52
11:K:92:ARG:HH11	11:K:92:ARG:CB	2.23	0.52
12:L:22:ALA:HB2	12:L:56:LEU:CD2	2.40	0.52
12:L:106:VAL:CG2	12:L:116:TYR:HB3	2.40	0.52
15:O:6:ALA:O	15:O:9:LYS:HB3	2.10	0.52
21:U:5:VAL:HG12	21:U:6:ARG:N	2.25	0.52
1:A:16:A:O2'	5:E:20:VAL:HG13	2.10	0.52
1:A:56:U:H2'	1:A:57:G:C8	2.45	0.52
1:A:936:C:H1'	1:A:1382:C:H42	1.74	0.52
1:A:1033:G:H2'	1:A:1034:G:O4'	2.10	0.52
1:A:1366:C:H2'	1:A:1367:C:C6	2.45	0.52
4:D:96:ARG:O	4:D:100:VAL:HG23	2.09	0.52
9:I:11:ARG:HA	9:I:105:ARG:HE	1.75	0.52
11:K:88:PRO:HD3	21:U:28:LEU:HD22	1.91	0.52
14:N:9:GLU:O	14:N:13:VAL:HG23	2.10	0.52
17:Q:24:ILE:HB	17:Q:41:THR:HB	1.92	0.52
17:Q:56:ASP:HB3	17:Q:79:GLU:O	2.09	0.52
19:S:62:THR:HB	19:S:64:GLU:OE2	2.10	0.52
1:A:337:G:H2'	1:A:338:A:C8	2.45	0.52
1:A:590:U:OP1	8:H:30:LYS:HE2	2.10	0.52
1:A:785:G:N7	22:V:71:ARG:NE	2.44	0.52
1:A:815:A:H4'	1:A:817:C:C4	2.45	0.52
1:A:920:U:H2'	1:A:921:U:H6	1.74	0.52
3:C:77:GLY:CA	3:C:81:GLU:HB3	2.40	0.52
4:D:18:LEU:HB2	4:D:20:LEU:HD11	1.92	0.52
11:K:111:ASP:HB3	21:U:3:ILE:HD13	1.92	0.52
13:M:106:ARG:HE	13:M:112:ARG:CG	2.14	0.52
16:P:71:VAL:HA	16:P:74:LEU:CG	2.39	0.52
21:U:14:ALA:H	21:U:16:ARG:NH2	2.07	0.52

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:120:A:H2'	1:A:121:U:H5''	1.91	0.51
1:A:266:G:O2'	1:A:267:C:H3'	2.10	0.51
1:A:731:G:OP1	1:A:766:A:H1'	2.10	0.51
2:B:204:ASP:CG	2:B:205:ALA:N	2.63	0.51
3:C:76:ILE:C	3:C:82:ASP:HB2	2.36	0.51
5:E:81:GLN:HG2	5:E:148:SER:HA	1.92	0.51
5:E:84:VAL:HG22	5:E:85:LYS:H	1.73	0.51
7:G:45:ALA:HB2	7:G:116:ALA:O	2.10	0.51
9:I:49:GLN:HE21	9:I:79:ARG:HD2	1.75	0.51
9:I:99:LYS:HD3	9:I:100:ALA:N	2.26	0.51
11:K:63:GLN:HB3	11:K:94:SER:OG	2.10	0.51
16:P:12:LYS:C	16:P:14:ARG:H	2.17	0.51
1:A:205:A:H2'	1:A:206:C:C6	2.44	0.51
1:A:275:G:H5'	17:Q:15:LYS:HG2	1.92	0.51
1:A:785:G:C4'	22:V:70:HIS:C	2.83	0.51
1:A:1390:U:H2'	1:A:1391:U:C6	2.45	0.51
2:B:95:TRP:CZ3	2:B:171:ALA:HA	2.44	0.51
6:F:55:HIS:O	6:F:56:LYS:HG3	2.09	0.51
6:F:67:PRO:HG2	6:F:70:VAL:HG22	1.92	0.51
1:A:255:G:H5'	17:Q:17:GLU:O	2.10	0.51
1:A:785:G:H4'	22:V:70:HIS:C	2.35	0.51
1:A:979:C:H2'	1:A:980:C:O4'	2.10	0.51
1:A:1122:U:H2'	1:A:1123:U:C6	2.45	0.51
1:A:1423:G:H2'	1:A:1424:U:O4'	2.10	0.51
1:A:1524:C:H2'	1:A:1525:G:C8	2.46	0.51
2:B:9:LEU:HD11	2:B:11:ALA:HB3	1.93	0.51
3:C:141:MET:CE	3:C:147:GLY:H	2.23	0.51
4:D:11:SER:HB2	4:D:20:LEU:HD11	1.92	0.51
4:D:197:HIS:HA	4:D:200:VAL:HG22	1.92	0.51
6:F:64:VAL:CG1	6:F:65:GLU:H	2.15	0.51
7:G:19:SER:OG	7:G:21:LEU:HG	2.10	0.51
11:K:109:ILE:H	21:U:5:VAL:HB	1.75	0.51
12:L:95:HIS:HD1	12:L:96:THR:N	2.09	0.51
13:M:27:THR:O	13:M:30:LYS:HB3	2.11	0.51
17:Q:35:LYS:O	17:Q:37:ILE:HG13	2.10	0.51
21:U:26:GLY:C	21:U:28:LEU:N	2.66	0.51
1:A:642:A:H2'	1:A:643:C:C6	2.46	0.51
1:A:785:G:H5'	22:V:69:VAL:HG13	1.83	0.51
1:A:958:A:N1	19:S:53:GLY:HA3	2.25	0.51
1:A:975:A:H5''	1:A:976:G:O5'	2.10	0.51
1:A:1247:U:O2'	1:A:1248:A:H5'	2.10	0.51

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:1485:U:O2'	22:V:286:LEU:N	2.43	0.51
2:B:162:VAL:HG13	2:B:184:ALA:CB	2.39	0.51
4:D:115:GLN:HE21	4:D:119:HIS:CE1	2.28	0.51
4:D:131:ILE:O	4:D:134:TYR:HB2	2.10	0.51
4:D:183:ARG:HH22	4:D:186:GLU:H	1.58	0.51
8:H:104:SER:O	8:H:122:GLY:HA3	2.10	0.51
12:L:121:PRO:C	12:L:123:ALA:H	2.18	0.51
14:N:27:LYS:HD2	14:N:28:ALA:N	2.26	0.51
18:R:52:ARG:HH11	18:R:52:ARG:CB	2.15	0.51
21:U:43:GLU:OE1	21:U:43:GLU:HA	2.10	0.51
1:A:309:A:H2'	1:A:310:G:H8	1.76	0.51
1:A:751:U:H2'	1:A:752:G:O4'	2.11	0.51
1:A:1190:G:O2'	3:C:2:GLN:HB2	2.10	0.51
1:A:1477:U:H2'	1:A:1478:U:C6	2.46	0.51
1:A:1484:C:C2'	22:V:340:ILE:HG21	2.41	0.51
2:B:53:LEU:HD12	2:B:219:THR:HG21	1.92	0.51
2:B:55:GLU:HG3	2:B:197:PHE:CZ	2.45	0.51
6:F:32:ALA:HB1	6:F:70:VAL:HG11	1.93	0.51
7:G:110:ARG:HH22	7:G:121:ASN:CB	2.23	0.51
11:K:22:ILE:CG2	11:K:95:THR:HG21	2.35	0.51
11:K:56:LYS:O	11:K:58:THR:N	2.43	0.51
11:K:80:ASN:HD21	11:K:107:THR:CG2	2.23	0.51
11:K:83:VAL:HG21	11:K:109:ILE:HG12	1.92	0.51
13:M:106:ARG:CZ	13:M:109:LYS:HD2	2.40	0.51
22:V:75:ARG:HD3	22:V:104:ILE:HD11	1.92	0.51
1:A:693:G:H2'	1:A:694:A:O4'	2.10	0.51
1:A:736:C:H2'	1:A:737:C:C6	2.45	0.51
1:A:909:A:H2'	1:A:910:C:O4'	2.11	0.51
1:A:1299:A:H62	1:A:1302:C:H5	1.57	0.51
1:A:1482:G:C4'	22:V:313:ASP:HA	2.40	0.51
13:M:78:ARG:HH12	19:S:64:GLU:HG2	1.75	0.51
19:S:50:VAL:H	19:S:57:VAL:H	1.57	0.51
1:A:237:G:O2'	1:A:238:A:H5'	2.11	0.51
1:A:413:G:O6	4:D:32:LYS:HE3	2.11	0.51
1:A:644:U:H4'	8:H:83:ARG:HH22	1.75	0.51
1:A:764:C:H3'	1:A:765:G:N2	2.25	0.51
3:C:186:SER:O	3:C:197:VAL:HG12	2.11	0.51
7:G:14:ASP:CG	7:G:15:PRO:HD2	2.35	0.51
7:G:66:GLU:HA	7:G:69:ARG:HE	1.75	0.51
9:I:93:LEU:HD13	9:I:97:LEU:HD23	1.92	0.51
9:I:110:VAL:HG12	9:I:111:GLU:N	2.20	0.51

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
11:K:56:LYS:H	11:K:56:LYS:HD3	1.75	0.51
14:N:17:ASP:HA	14:N:21:ALA:HB2	1.92	0.51
22:V:134:ILE:HG13	22:V:134:ILE:O	2.10	0.51
1:A:71:A:N6	1:A:99:C:H1'	2.24	0.51
1:A:939:G:H4'	7:G:101:ARG:NH2	2.25	0.51
1:A:1051:C:H2'	1:A:1052:U:C6	2.46	0.51
1:A:1098:C:C2'	1:A:1099:G:H5'	2.40	0.51
1:A:1289:A:H2'	1:A:1290:G:H5'	1.91	0.51
2:B:37:VAL:HG22	2:B:38:HIS:H	1.76	0.51
2:B:95:TRP:HH2	2:B:100:LEU:HB2	1.75	0.51
4:D:116:LEU:HB3	4:D:122:ILE:HD11	1.93	0.51
4:D:123:MET:HG3	4:D:127:ARG:C	2.36	0.51
5:E:37:VAL:HG11	5:E:113:VAL:HG12	1.92	0.51
5:E:110:MET:O	5:E:113:VAL:HG22	2.10	0.51
7:G:3:ARG:HB3	7:G:3:ARG:CZ	2.40	0.51
11:K:110:THR:CG2	21:U:4:LYS:HD2	2.40	0.51
15:O:80:LEU:HD23	15:O:80:LEU:C	2.36	0.51
18:R:62:ARG:HD2	18:R:69:TYR:CD1	2.46	0.51
22:V:202:THR:O	22:V:203:GLN:HB2	2.11	0.51
1:A:177:G:P	20:T:59:ARG:HE	2.33	0.51
1:A:208:U:H2'	1:A:210:C:C5	2.45	0.51
1:A:1460:C:H2'	1:A:1461:G:O4'	2.11	0.51
4:D:64:TYR:H	4:D:64:TYR:HD1	1.59	0.51
4:D:71:PHE:O	4:D:74:TYR:HB2	2.11	0.51
4:D:172:VAL:HA	4:D:178:GLU:O	2.11	0.51
9:I:32:ARG:NH2	9:I:36:GLN:HG2	2.25	0.51
11:K:80:ASN:HD21	11:K:107:THR:HG23	1.76	0.51
18:R:60:ARG:HG3	18:R:61:ALA:N	2.25	0.51
1:A:838:G:H2'	1:A:839:C:O4'	2.11	0.51
1:A:1320:C:H41	19:S:36:ARG:HE	1.58	0.51
1:A:1412:C:H2'	1:A:1413:A:H8	1.73	0.51
2:B:163:ILE:CD1	2:B:209:VAL:HG12	2.38	0.51
3:C:21:TRP:HB3	3:C:58:ARG:HB2	1.93	0.51
3:C:78:LYS:HE3	3:C:81:GLU:OE2	2.11	0.51
3:C:149:LYS:HB3	3:C:200:TRP:HB2	1.94	0.51
4:D:112:GLU:OE2	4:D:153:ARG:HD3	2.11	0.51
7:G:11:ILE:HD12	7:G:11:ILE:H	1.75	0.51
8:H:60:LEU:HD12	8:H:60:LEU:N	2.26	0.51
10:J:55:PRO:O	10:J:56:HIS:HB3	2.10	0.51
11:K:125:LYS:O	11:K:126:ARG:O	2.29	0.51
12:L:3:VAL:HG23	12:L:4:ASN:OD1	2.11	0.51

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:33:A:H2'	1:A:34:C:C6	2.46	0.50
1:A:492:C:H2'	1:A:493:A:H5''	1.92	0.50
1:A:545:C:O2'	1:A:546:A:H5'	2.10	0.50
1:A:1197:A:O2'	1:A:1198:G:H5'	2.10	0.50
1:A:1206:G:H4'	3:C:192:TYR:HA	1.93	0.50
1:A:1332:A:H2'	1:A:1333:A:H8	1.74	0.50
3:C:100:ILE:HG23	3:C:100:ILE:O	2.10	0.50
6:F:12:PRO:HG3	6:F:54:LEU:HG	1.94	0.50
16:P:40:ASN:HD21	16:P:42:ILE:HG13	1.75	0.50
22:V:151:ARG:HB3	22:V:151:ARG:NH1	2.25	0.50
1:A:90:C:H2'	1:A:91:U:C6	2.47	0.50
1:A:807:A:H2'	1:A:808:C:C6	2.47	0.50
1:A:824:G:H2'	1:A:825:A:H8	1.75	0.50
1:A:840:C:C2'	1:A:842:U:H5''	2.38	0.50
1:A:1420:U:H5''	22:V:307:TYR:N	2.26	0.50
2:B:95:TRP:CE3	2:B:171:ALA:HA	2.46	0.50
3:C:83:VAL:HG12	3:C:87:ARG:HE	1.75	0.50
3:C:89:VAL:C	3:C:91:ALA:H	2.19	0.50
7:G:72:VAL:HB	7:G:144:ALA:HB3	1.92	0.50
8:H:8:ASP:OD2	8:H:12:ARG:HD2	2.11	0.50
9:I:87:MET:HE3	9:I:94:ARG:NE	2.24	0.50
10:J:11:LYS:HZ3	10:J:99:GLN:H	1.58	0.50
11:K:122:PRO:HG2	21:U:35:GLU:HG2	1.93	0.50
12:L:54:VAL:HG12	12:L:56:LEU:HD12	1.93	0.50
13:M:10:ASP:HA	13:M:44:ILE:CD1	2.41	0.50
13:M:106:ARG:HH11	13:M:106:ARG:CA	2.24	0.50
1:A:216:U:H2'	1:A:217:C:C6	2.46	0.50
1:A:219:U:H2'	1:A:220:G:C8	2.46	0.50
1:A:275:G:O5'	17:Q:15:LYS:HG2	2.11	0.50
1:A:663:A:O2'	1:A:664:G:H5'	2.11	0.50
1:A:1285:A:N6	1:A:1355:G:H4'	2.26	0.50
1:A:1410:A:OP1	22:V:147:ASN:OD1	2.29	0.50
1:A:1434:A:H2'	1:A:1435:G:C8	2.46	0.50
1:A:1490:U:H3'	1:A:1491:G:H8	1.77	0.50
21:U:3:ILE:CA	21:U:19:LYS:HG2	2.39	0.50
1:A:22:G:H4'	1:A:885:G:C8	2.47	0.50
1:A:49:U:O2'	1:A:50:A:H2'	2.11	0.50
1:A:476:U:H2'	1:A:477:C:H6	1.73	0.50
1:A:613:C:H2'	1:A:614:C:C6	2.46	0.50
1:A:857:C:H2'	1:A:858:G:O4'	2.10	0.50
1:A:1483:A:N6	22:V:341:LEU:HD11	2.22	0.50

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
9:I:11:ARG:HG3	9:I:11:ARG:O	2.10	0.50
10:J:67:ILE:HG12	14:N:94:GLY:O	2.11	0.50
12:L:103:CYS:SG	12:L:104:SER:N	2.83	0.50
1:A:90:C:H2'	1:A:91:U:H6	1.76	0.50
1:A:546:A:H4'	1:A:548:G:O3'	2.12	0.50
1:A:633:G:H2'	1:A:634:C:C6	2.47	0.50
2:B:100:LEU:O	2:B:178:LEU:HG	2.11	0.50
2:B:113:LEU:O	2:B:117:GLU:HG3	2.12	0.50
3:C:71:ARG:NH2	3:C:74:ILE:HB	2.26	0.50
4:D:51:GLY:O	4:D:55:ARG:HB2	2.12	0.50
5:E:110:MET:SD	5:E:110:MET:N	2.85	0.50
7:G:23:ALA:O	7:G:26:VAL:HG22	2.11	0.50
12:L:20:VAL:HG22	12:L:94:TYR:CE1	2.46	0.50
22:V:134:ILE:HG12	22:V:156:CYS:SG	2.51	0.50
1:A:33:A:H2'	1:A:34:C:H6	1.77	0.50
1:A:529:G:H22	12:L:47:ALA:CB	2.24	0.50
1:A:556:C:O2'	1:A:557:G:H5'	2.11	0.50
1:A:642:A:H2'	1:A:643:C:H6	1.76	0.50
2:B:132:GLU:O	2:B:136:ARG:HG3	2.11	0.50
4:D:2:ARG:O	4:D:3:TYR:HB2	2.11	0.50
11:K:49:SER:HA	11:K:68:ARG:HH21	1.77	0.50
12:L:31:GLY:HA3	12:L:54:VAL:CG1	2.41	0.50
16:P:52:LEU:HG	16:P:75:ILE:CG1	2.40	0.50
22:V:43:LEU:HD12	22:V:93:LYS:HD2	1.94	0.50
1:A:235:C:H2'	1:A:236:A:C8	2.47	0.50
1:A:251:G:H4'	1:A:252:U:H5'	1.93	0.50
1:A:1101:A:N7	2:B:170:ILE:HD12	2.26	0.50
1:A:1172:C:H2'	1:A:1173:U:H6	1.76	0.50
1:A:1343:G:H2'	1:A:1344:C:C6	2.46	0.50
2:B:71:THR:CG2	2:B:94:ARG:HH21	2.25	0.50
3:C:110:LEU:O	3:C:203:LYS:HE2	2.10	0.50
5:E:89:THR:HG21	5:E:134:ASN:ND2	2.27	0.50
5:E:92:ARG:HB3	5:E:127:TYR:HB2	1.94	0.50
6:F:54:LEU:HD13	6:F:55:HIS:H	1.77	0.50
7:G:72:VAL:HB	7:G:144:ALA:CB	2.41	0.50
13:M:13:HIS:O	13:M:16:ILE:HG22	2.11	0.50
13:M:89:ARG:HH11	13:M:94:LEU:HD13	1.76	0.50
1:A:1482:G:C4	22:V:311:LYS:O	2.65	0.50
3:C:150:VAL:HG12	3:C:151:GLU:H	1.77	0.50
4:D:84:ASN:HD22	4:D:87:GLU:N	2.10	0.50
4:D:169:TRP:HB3	4:D:183:ARG:HH21	1.77	0.50

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
5:E:17:VAL:HA	5:E:34:ALA:HA	1.93	0.50
5:E:155:LYS:HE3	8:H:70:VAL:HG13	1.94	0.50
10:J:41:PRO:O	10:J:42:LEU:HB2	2.12	0.50
16:P:5:ARG:H	16:P:5:ARG:HD2	1.77	0.50
17:Q:23:ALA:HB1	17:Q:40:THR:HG23	1.94	0.50
21:U:40:PRO:C	21:U:42:THR:N	2.69	0.50
1:A:16:A:H4'	5:E:21:SER:H	1.76	0.50
1:A:300:A:H2'	1:A:301:G:O4'	2.11	0.50
1:A:971:G:OP1	1:A:971:G:H3'	2.12	0.50
1:A:1137:C:O2'	1:A:1138:G:H5''	2.12	0.50
1:A:1484:C:C2'	22:V:291:ILE:CD1	2.87	0.50
2:B:67:LEU:HD22	2:B:67:LEU:N	2.21	0.50
2:B:114:LYS:HA	2:B:117:GLU:OE1	2.12	0.50
3:C:119:ILE:O	3:C:123:LEU:HG	2.12	0.50
4:D:17:ASP:HB2	4:D:27:ILE:HG23	1.94	0.50
9:I:51:LEU:CD1	9:I:82:ILE:HG21	2.42	0.50
10:J:57:VAL:HG13	10:J:58:ASN:N	2.27	0.50
11:K:88:PRO:HA	11:K:92:ARG:HD2	1.93	0.50
1:A:214:C:H2'	1:A:215:C:H6	1.76	0.49
1:A:685:G:O4'	11:K:40:ALA:HB3	2.11	0.49
1:A:841:C:H3'	1:A:843:U:OP2	2.11	0.49
1:A:1366:C:H2'	1:A:1367:C:H6	1.76	0.49
1:A:1372:U:OP1	9:I:72:SER:HB2	2.12	0.49
7:G:103:ILE:O	7:G:107:ALA:HB2	2.11	0.49
16:P:67:ILE:HG13	16:P:71:VAL:CG1	2.41	0.49
22:V:173:LEU:HD11	22:V:181:VAL:HG21	1.94	0.49
1:A:537:G:H2'	1:A:538:G:C8	2.47	0.49
1:A:685:G:O2'	1:A:686:U:H5'	2.12	0.49
1:A:708:C:H2'	1:A:709:U:C6	2.47	0.49
1:A:1007:U:H2'	1:A:1008:U:H6	1.77	0.49
1:A:1169:A:H2'	1:A:1170:A:C8	2.47	0.49
2:B:76:SER:HA	2:B:92:ASN:HB2	1.94	0.49
2:B:218:ALA:HA	2:B:221:ARG:HE	1.76	0.49
3:C:71:ARG:O	3:C:74:ILE:HG22	2.11	0.49
6:F:6:ILE:HB	6:F:89:VAL:HB	1.93	0.49
7:G:61:PHE:O	7:G:65:LEU:HD13	2.11	0.49
17:Q:10:ARG:NH2	17:Q:11:VAL:HG23	2.27	0.49
17:Q:18:LYS:HE3	17:Q:48:GLU:HG2	1.93	0.49
1:A:17:U:H2'	1:A:18:C:H6	1.73	0.49
1:A:110:C:H2'	1:A:111:G:O4'	2.12	0.49
1:A:182:A:H1'	1:A:183:C:C5	2.47	0.49

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:711:G:O2'	1:A:712:A:H5'	2.11	0.49
1:A:969:A:N3	1:A:970:C:O2	2.45	0.49
1:A:1108:G:H5'	3:C:175:HIS:CD2	2.47	0.49
1:A:1142:G:H3'	1:A:1143:G:H8	1.76	0.49
1:A:1324:A:H2'	1:A:1325:C:C6	2.46	0.49
1:A:1420:U:P	22:V:309:ASP:OD1	2.67	0.49
1:A:1482:G:O4'	22:V:312:HIS:C	2.55	0.49
2:B:107:ARG:HA	2:B:110:ILE:CD1	2.42	0.49
3:C:150:VAL:HG12	3:C:151:GLU:N	2.27	0.49
3:C:154:GLY:HA2	3:C:163:ARG:H	1.77	0.49
4:D:24:VAL:O	4:D:25:ARG:C	2.55	0.49
17:Q:10:ARG:NE	17:Q:10:ARG:CA	2.76	0.49
22:V:82:VAL:HG21	22:V:129:ALA:HB3	1.94	0.49
1:A:16:A:N1	1:A:919:A:H2	2.10	0.49
1:A:72:A:H2'	1:A:73:C:C6	2.47	0.49
1:A:272:C:H2'	1:A:273:U:C6	2.45	0.49
1:A:560:A:H5'	1:A:566:G:N2	2.28	0.49
1:A:837:U:H2'	1:A:838:G:H8	1.77	0.49
1:A:1284:C:H3'	1:A:1285:A:C5'	2.42	0.49
2:B:131:LYS:HE3	2:B:135:MET:HE2	1.94	0.49
2:B:195:VAL:HG12	2:B:197:PHE:H	1.78	0.49
4:D:69:ARG:HE	4:D:69:ARG:CA	2.25	0.49
5:E:114:LEU:C	5:E:116:VAL:H	2.21	0.49
6:F:98:GLU:HG2	6:F:99:ALA:H	1.76	0.49
9:I:118:ARG:HH22	9:I:122:ARG:NH2	2.10	0.49
11:K:52:ARG:O	11:K:54:SER:N	2.46	0.49
18:R:63:TYR:C	18:R:65:SER:H	2.19	0.49
22:V:286:LEU:HB2	22:V:291:ILE:HD11	1.94	0.49
1:A:131:A:H2'	1:A:132:C:H6	1.75	0.49
1:A:785:G:C4'	22:V:71:ARG:N	2.75	0.49
1:A:840:C:C2	1:A:842:U:H4'	2.47	0.49
3:C:85:LYS:O	3:C:89:VAL:HG23	2.13	0.49
7:G:144:ALA:C	7:G:146:ALA:N	2.70	0.49
9:I:34:LEU:HD11	9:I:47:VAL:CG2	2.36	0.49
9:I:53:LEU:O	9:I:53:LEU:HD13	2.12	0.49
12:L:48:LEU:H	12:L:48:LEU:CD2	2.18	0.49
13:M:89:ARG:NH1	13:M:94:LEU:HD13	2.27	0.49
19:S:20:LYS:HB3	19:S:20:LYS:HZ2	1.78	0.49
21:U:8:ASN:O	21:U:9:GLU:HB2	2.12	0.49
1:A:98:A:H2'	1:A:99:C:C6	2.47	0.49
1:A:575:G:H4'	1:A:576:C:O5'	2.13	0.49

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:677:U:H2'	1:A:678:U:C6	2.48	0.49
1:A:1225:A:H4'	19:S:77:ARG:HH22	1.75	0.49
1:A:1481:U:H2'	1:A:1482:G:C8	2.47	0.49
1:A:1485:U:C1'	22:V:286:LEU:HB2	2.42	0.49
9:I:18:VAL:HG21	9:I:82:ILE:CG1	2.43	0.49
12:L:35:ARG:O	12:L:52:CYS:HB2	2.13	0.49
19:S:12:LEU:O	19:S:15:LEU:HB3	2.13	0.49
19:S:51:HIS:HA	19:S:55:GLN:O	2.12	0.49
20:T:68:LYS:HG3	20:T:69:ASN:ND2	2.27	0.49
1:A:915:A:H2'	1:A:916:U:H5'	1.94	0.49
1:A:1096:C:O2'	1:A:1097:C:H5'	2.12	0.49
8:H:10:LEU:HA	8:H:74:ILE:HD11	1.95	0.49
8:H:17:GLN:NE2	8:H:69:ALA:HB1	2.28	0.49
11:K:15:VAL:HG11	11:K:35:ASP:HB2	1.94	0.49
1:A:79:G:H2'	1:A:80:A:O4'	2.13	0.49
1:A:218:U:H2'	1:A:219:U:H6	1.77	0.49
1:A:782:A:H2'	1:A:783:C:O4'	2.13	0.49
1:A:900:A:H2'	1:A:901:A:C8	2.47	0.49
1:A:1060:U:H4'	10:J:54:SER:HB2	1.95	0.49
1:A:1234:C:H2'	1:A:1235:U:C6	2.48	0.49
1:A:1482:G:C1'	22:V:311:LYS:O	2.59	0.49
2:B:59:ILE:HD12	2:B:60:ALA:N	2.27	0.49
7:G:94:ARG:HD3	7:G:98:LEU:HD11	1.94	0.49
12:L:113:ARG:HA	12:L:118:VAL:HG23	1.93	0.49
15:O:78:THR:HA	15:O:81:ILE:CD1	2.42	0.49
21:U:29:ALA:HA	21:U:32:ARG:HB2	1.94	0.49
1:A:1110:A:C2'	1:A:1111:A:H5'	2.42	0.49
1:A:1415:G:H3'	22:V:345:ALA:C	2.38	0.49
1:A:1452:C:H5'	1:A:1453:G:C4	2.47	0.49
1:A:1484:C:C6	22:V:337:TYR:CD1	3.01	0.49
2:B:175:ALA:C	2:B:177:ASN:H	2.20	0.49
2:B:187:ASP:CG	2:B:188:THR:N	2.70	0.49
3:C:113:LYS:HG3	3:C:117:ASP:OD2	2.13	0.49
4:D:29:THR:O	4:D:30:LYS:HB2	2.13	0.49
4:D:77:GLU:O	4:D:81:LEU:HG	2.11	0.49
6:F:37:HIS:HE1	6:F:65:GLU:HB2	1.73	0.49
9:I:18:VAL:HG21	9:I:82:ILE:HG13	1.95	0.49
9:I:112:ARG:HB2	9:I:112:ARG:HH11	1.77	0.49
13:M:58:GLU:HA	13:M:61:LYS:HE2	1.94	0.49
14:N:74:ARG:O	14:N:74:ARG:HD3	2.13	0.49
18:R:40:PRO:HD2	18:R:43:ILE:HD11	1.94	0.49

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:31:G:H2'	1:A:48:C:H5	1.77	0.49
1:A:327:A:H1'	1:A:329:A:O4'	2.13	0.49
1:A:664:G:H5''	18:R:52:ARG:NH2	2.27	0.49
1:A:1178:G:H2'	1:A:1180:A:OP2	2.13	0.49
1:A:1271:A:H2'	1:A:1272:G:H8	1.78	0.49
2:B:31:PHE:HA	2:B:41:ASN:HB2	1.94	0.49
2:B:162:VAL:HG21	2:B:168:GLU:HB2	1.95	0.49
6:F:39:LEU:HD13	6:F:40:GLU:H	1.77	0.49
13:M:38:ILE:HG13	13:M:55:LEU:CD2	2.41	0.49
15:O:87:ARG:HA	15:O:87:ARG:HH11	1.78	0.49
16:P:9:HIS:CE1	16:P:29:ASN:HB2	2.44	0.49
17:Q:60:ILE:HB	17:Q:72:TRP:HE3	1.78	0.49
1:A:87:C:N3	1:A:88:U:H1'	2.28	0.48
1:A:254:G:O2'	1:A:255:G:H5'	2.13	0.48
1:A:265:G:H2'	1:A:267:C:H5	1.78	0.48
1:A:755:G:OP2	15:O:64:LYS:HG2	2.13	0.48
1:A:806:C:H2'	1:A:807:A:H8	1.77	0.48
1:A:1343:G:H2'	1:A:1344:C:H6	1.78	0.48
2:B:36:LYS:N	2:B:36:LYS:HE3	2.27	0.48
2:B:46:VAL:CG1	2:B:47:PRO:HD3	2.31	0.48
2:B:215:ALA:O	2:B:218:ALA:HB3	2.13	0.48
3:C:78:LYS:H	3:C:81:GLU:CB	2.25	0.48
5:E:41:GLY:HA2	5:E:118:GLY:HA2	1.94	0.48
5:E:111:ARG:O	5:E:115:GLU:HG3	2.13	0.48
9:I:32:ARG:HD3	9:I:37:TYR:HD1	1.77	0.48
13:M:15:VAL:HG13	13:M:29:SER:OG	2.13	0.48
16:P:20:VAL:HG21	16:P:32:PHE:CD2	2.48	0.48
1:A:235:C:H2'	1:A:236:A:H8	1.78	0.48
1:A:537:G:H2'	1:A:538:G:H8	1.78	0.48
1:A:720:C:C5'	18:R:40:PRO:HA	2.43	0.48
1:A:723:U:O4'	21:U:48:LYS:HD3	2.13	0.48
1:A:1281:C:H5'	1:A:1282:C:C5	2.48	0.48
1:A:1478:U:H2'	1:A:1479:C:C6	2.48	0.48
1:A:1486:G:C8	22:V:344:MET:HE2	2.45	0.48
2:B:206:ILE:O	2:B:209:VAL:HG23	2.13	0.48
11:K:125:LYS:O	21:U:33:ARG:NH2	2.45	0.48
18:R:70:THR:OG1	18:R:71:ASP:N	2.46	0.48
20:T:61:ALA:HA	20:T:66:ILE:O	2.13	0.48
22:V:82:VAL:HG21	22:V:129:ALA:CB	2.43	0.48
1:A:179:A:H2'	1:A:180:U:C6	2.48	0.48
1:A:182:A:H1'	1:A:183:C:H5	1.79	0.48

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:220:G:O2'	1:A:221:C:H5'	2.12	0.48
1:A:370:C:O2'	1:A:371:A:H5'	2.13	0.48
1:A:812:G:N3	1:A:812:G:C2'	2.76	0.48
1:A:1137:C:H1'	1:A:1138:G:N2	2.28	0.48
1:A:1285:A:H62	1:A:1355:G:H4'	1.76	0.48
6:F:67:PRO:O	6:F:70:VAL:HG22	2.13	0.48
13:M:22:TYR:HB3	13:M:69:ARG:HH22	1.78	0.48
14:N:12:ARG:HH11	14:N:60:ARG:HH12	1.62	0.48
17:Q:8:GLN:HB3	17:Q:57:VAL:HG13	1.95	0.48
19:S:64:GLU:OE2	19:S:65:MET:HG3	2.12	0.48
1:A:34:C:H2'	1:A:35:G:C8	2.47	0.48
1:A:157:U:O2'	1:A:158:G:H5'	2.13	0.48
1:A:513:C:H2'	1:A:514:C:C6	2.49	0.48
1:A:953:G:H2'	1:A:954:G:O4'	2.14	0.48
1:A:1086:U:H4'	1:A:1389:C:H5''	1.94	0.48
1:A:1130:A:O2'	1:A:1131:G:H5'	2.13	0.48
1:A:1314:C:H2'	1:A:1315:U:C6	2.49	0.48
1:A:1456:A:H2'	1:A:1457:G:O4'	2.14	0.48
1:A:1464:U:H2'	1:A:1465:A:C8	2.48	0.48
2:B:9:LEU:HD22	2:B:11:ALA:N	2.24	0.48
2:B:22:TRP:CG	2:B:23:ASN:N	2.81	0.48
4:D:10:LEU:HD21	4:D:62:ARG:CD	2.41	0.48
4:D:194:ILE:HD11	4:D:199:ILE:HD11	1.96	0.48
5:E:45:VAL:HG23	5:E:71:ILE:CG2	2.43	0.48
5:E:63:MET:O	5:E:66:ALA:HB3	2.14	0.48
8:H:87:ARG:O	8:H:91:LEU:HG	2.14	0.48
9:I:4:GLN:NE2	9:I:21:LYS:HE3	2.28	0.48
9:I:56:MET:HA	9:I:59:LYS:HB2	1.94	0.48
10:J:55:PRO:HA	14:N:80:ARG:HH22	1.78	0.48
11:K:83:VAL:CG2	11:K:109:ILE:HG12	2.44	0.48
17:Q:68:LYS:O	17:Q:69:THR:CB	2.61	0.48
21:U:3:ILE:HA	21:U:19:LYS:CG	2.37	0.48
21:U:40:PRO:HG2	21:U:41:THR:N	2.27	0.48
22:V:276:PRO:O	22:V:280:GLU:HG3	2.13	0.48
1:A:56:U:H2'	1:A:57:G:H8	1.77	0.48
1:A:659:U:H2'	1:A:660:C:C6	2.49	0.48
1:A:864:A:H2'	1:A:865:A:C8	2.49	0.48
1:A:957:U:H3	1:A:960:U:H5''	1.77	0.48
1:A:1018:G:H2'	1:A:1019:A:C8	2.48	0.48
1:A:1109:C:N4	1:A:1110:A:N6	2.62	0.48
1:A:1526:G:O2'	1:A:1527:U:H5'	2.13	0.48

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:B:15:PHE:HD1	2:B:16:GLY:H	1.61	0.48
6:F:100:SER:HA	18:R:23:LYS:CE	2.42	0.48
7:G:14:ASP:OD2	7:G:15:PRO:HD2	2.14	0.48
8:H:24:VAL:CG1	8:H:60:LEU:HB2	2.43	0.48
8:H:29:SER:HB3	8:H:32:LYS:HZ2	1.78	0.48
9:I:87:MET:SD	9:I:94:ARG:HG3	2.53	0.48
12:L:35:ARG:HG3	12:L:36:VAL:N	2.26	0.48
13:M:109:LYS:HG3	13:M:110:GLY:N	2.28	0.48
17:Q:20:ILE:HG21	17:Q:52:CYS:SG	2.53	0.48
17:Q:29:LYS:HA	17:Q:35:LYS:C	2.39	0.48
1:A:208:U:C2'	1:A:209:U:H5''	2.43	0.48
1:A:333:U:H2'	1:A:334:C:H6	1.79	0.48
1:A:462:G:H3'	1:A:463:U:H5''	1.95	0.48
1:A:672:U:H2'	1:A:673:A:H8	1.78	0.48
1:A:878:A:C5'	8:H:80:PRO:HG2	2.43	0.48
2:B:163:ILE:CG2	2:B:164:ASP:H	1.98	0.48
3:C:115:VAL:CG1	3:C:137:VAL:HG13	2.43	0.48
3:C:122:GLN:HE22	3:C:136:ALA:HB1	1.78	0.48
10:J:65:TYR:C	14:N:98:ALA:HB2	2.39	0.48
12:L:58:ASN:HD22	12:L:58:ASN:N	2.11	0.48
1:A:5:U:H1'	1:A:6:G:C2	2.48	0.48
1:A:174:A:O2'	1:A:175:C:H5'	2.13	0.48
1:A:193:C:H2'	1:A:194:C:C6	2.49	0.48
1:A:588:G:H5'	8:H:2:MET:O	2.14	0.48
1:A:600:A:H2'	1:A:601:G:C8	2.48	0.48
1:A:684:U:O2'	11:K:39:ASN:HB3	2.14	0.48
1:A:1072:G:H2'	1:A:1073:U:C6	2.48	0.48
1:A:1329:A:O2'	1:A:1330:U:H5'	2.13	0.48
1:A:1419:G:H5''	22:V:309:ASP:CB	2.44	0.48
1:A:1484:C:H1'	22:V:291:ILE:O	2.12	0.48
1:A:1506:U:H4'	11:K:128:VAL:OXT	2.13	0.48
2:B:35:ASN:O	2:B:36:LYS:HB2	2.13	0.48
5:E:104:ILE:HG23	5:E:104:ILE:O	2.14	0.48
7:G:137:ARG:HG2	7:G:141:HIS:CE1	2.48	0.48
8:H:29:SER:HB3	8:H:32:LYS:HG3	1.95	0.48
17:Q:10:ARG:NH1	17:Q:11:VAL:HB	2.28	0.48
18:R:64:LEU:CB	18:R:66:LEU:HG	2.44	0.48
22:V:55:ARG:NH1	22:V:57:GLY:O	2.47	0.48
1:A:335:C:H2'	1:A:336:A:H8	1.78	0.48
1:A:434:U:O2	1:A:434:U:H2'	2.14	0.48
1:A:541:G:H2'	1:A:542:G:H8	1.79	0.48

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:677:U:H2'	1:A:678:U:H6	1.77	0.48
1:A:806:C:H2'	1:A:807:A:C8	2.48	0.48
1:A:911:U:H2'	1:A:912:C:C6	2.49	0.48
1:A:1002:G:H2'	1:A:1002:G:N3	2.27	0.48
1:A:1397:C:H4'	1:A:1398:A:OP2	2.13	0.48
3:C:61:LYS:O	3:C:97:PRO:HD2	2.14	0.48
3:C:78:LYS:H	3:C:81:GLU:HB3	1.79	0.48
4:D:7:LYS:HG3	4:D:20:LEU:HB2	1.94	0.48
6:F:66:ALA:HB1	6:F:67:PRO:HD2	1.96	0.48
18:R:46:THR:CG2	18:R:51:GLN:HB2	2.43	0.48
19:S:35:ARG:HG2	19:S:50:VAL:CG1	2.39	0.48
1:A:232:G:H1'	1:A:262:A:N1	2.29	0.48
1:A:252:U:H2'	1:A:253:A:C8	2.48	0.48
1:A:321:A:O2'	1:A:322:C:H5'	2.14	0.48
1:A:522:C:H41	12:L:49:ARG:NH2	2.10	0.48
1:A:734:G:H2'	1:A:735:C:C6	2.49	0.48
1:A:1367:C:O2'	1:A:1368:A:H5'	2.14	0.48
1:A:1414:U:N3	22:V:343:SER:HA	2.24	0.48
1:A:1484:C:C3'	22:V:340:ILE:HG21	2.44	0.48
1:A:1486:G:H4'	22:V:285:HIS:CA	2.30	0.48
2:B:98:GLY:C	2:B:100:LEU:H	2.22	0.48
4:D:93:LEU:O	4:D:96:ARG:HB3	2.14	0.48
12:L:80:LEU:O	12:L:97:VAL:HG23	2.14	0.48
14:N:15:LEU:C	14:N:17:ASP:H	2.21	0.48
16:P:29:ASN:N	16:P:29:ASN:HD22	2.11	0.48
1:A:303:A:H2'	1:A:304:U:O4'	2.14	0.48
1:A:784:A:C3'	22:V:72:CYS:N	2.76	0.48
1:A:986:U:H1'	19:S:53:GLY:O	2.14	0.48
1:A:1081:A:O2'	1:A:1082:A:H5'	2.14	0.48
1:A:1220:G:H2'	1:A:1221:G:H8	1.77	0.48
1:A:1369:C:H2'	1:A:1370:G:C8	2.49	0.48
2:B:9:LEU:H	2:B:9:LEU:HD12	1.78	0.48
2:B:45:THR:HA	2:B:48:MET:HG3	1.95	0.48
2:B:79:VAL:HG13	2:B:90:PHE:HD2	1.79	0.48
2:B:162:VAL:HG11	2:B:172:ILE:HG12	1.96	0.48
2:B:162:VAL:O	2:B:184:ALA:HA	2.14	0.48
2:B:163:ILE:HG13	2:B:185:ILE:HD11	1.96	0.48
3:C:21:TRP:CD1	3:C:58:ARG:HD2	2.49	0.48
3:C:100:ILE:C	3:C:100:ILE:HD13	2.38	0.48
6:F:43:GLY:HA2	6:F:58:HIS:CE1	2.49	0.48
7:G:71:THR:HA	7:G:90:VAL:HG22	1.96	0.48

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
8:H:64:TYR:CB	8:H:69:ALA:HA	2.44	0.48
9:I:23:GLY:N	9:I:60:LEU:HA	2.24	0.48
9:I:64:ILE:HG22	9:I:65:THR:N	2.28	0.48
11:K:63:GLN:O	11:K:67:GLU:HB3	2.14	0.48
11:K:121:ARG:HG3	11:K:121:ARG:NH1	2.29	0.48
13:M:33:LEU:HB3	13:M:38:ILE:O	2.14	0.48
13:M:84:CYS:HA	19:S:72:GLU:O	2.13	0.48
19:S:4:LEU:N	19:S:5:LYS:HE3	2.29	0.48
22:V:274:ASP:O	22:V:275:SER:HB3	2.13	0.48
1:A:255:G:H2'	1:A:256:U:C6	2.48	0.47
1:A:1258:G:C4	1:A:1278:G:N2	2.82	0.47
1:A:1486:G:H5'	22:V:285:HIS:N	2.27	0.47
1:A:1504:G:H4'	1:A:1505:G:C4	2.49	0.47
6:F:75:GLU:O	6:F:79:ARG:HG2	2.14	0.47
7:G:131:GLY:O	7:G:134:VAL:HG12	2.13	0.47
8:H:11:THR:HG22	8:H:15:ASN:ND2	2.29	0.47
13:M:78:ARG:HH22	13:M:82:LEU:HD11	1.78	0.47
20:T:40:ALA:O	20:T:41:GLY:C	2.56	0.47
20:T:49:ALA:HA	20:T:52:GLU:HB3	1.96	0.47
1:A:78:A:O2'	1:A:79:G:H5'	2.15	0.47
1:A:421:U:H1'	3:C:125:ARG:HH12	1.79	0.47
1:A:607:A:H2'	1:A:608:A:C8	2.50	0.47
1:A:1014:A:C2	1:A:1219:A:H1'	2.49	0.47
1:A:1118:U:H1'	1:A:1179:A:C5	2.50	0.47
1:A:1414:U:H2'	1:A:1415:G:H8	1.79	0.47
2:B:27:LYS:C	2:B:29:PHE:H	2.21	0.47
2:B:80:LYS:O	2:B:84:LEU:N	2.47	0.47
4:D:94:GLU:HA	4:D:94:GLU:OE1	2.14	0.47
10:J:88:MET:CE	10:J:88:MET:H	2.27	0.47
15:O:7:THR:CG2	15:O:30:LEU:HD11	2.38	0.47
16:P:71:VAL:CA	16:P:74:LEU:HG	2.41	0.47
17:Q:10:ARG:NE	17:Q:11:VAL:H	2.12	0.47
1:A:238:A:C2'	1:A:239:U:H5''	2.43	0.47
1:A:317:U:H2'	1:A:318:G:C8	2.49	0.47
1:A:636:U:H2'	1:A:637:C:H6	1.78	0.47
1:A:1215:G:H5'	1:A:1215:G:H8	1.78	0.47
1:A:1348:U:H2'	1:A:1349:A:H8	1.80	0.47
1:A:1432:G:H8	1:A:1432:G:OP2	1.97	0.47
1:A:1484:C:C2'	22:V:291:ILE:HA	2.41	0.47
2:B:172:ILE:O	2:B:175:ALA:HB3	2.13	0.47
16:P:36:VAL:HG13	16:P:36:VAL:O	2.14	0.47

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
20:T:68:LYS:CG	20:T:69:ASN:H	2.12	0.47
21:U:31:VAL:O	21:U:32:ARG:C	2.57	0.47
1:A:239:U:H5''	1:A:239:U:H6	1.78	0.47
1:A:343:U:O2'	1:A:344:A:H2'	2.14	0.47
1:A:719:C:H1'	18:R:37:LYS:HE3	1.96	0.47
1:A:946:A:H2'	1:A:947:G:H8	1.75	0.47
1:A:1198:G:H2'	1:A:1199:U:C6	2.50	0.47
1:A:1481:U:O2	22:V:313:ASP:CB	2.61	0.47
3:C:5:HIS:CD2	3:C:7:ASN:HB2	2.49	0.47
3:C:66:THR:HA	3:C:101:ASN:O	2.14	0.47
4:D:160:LEU:H	4:D:160:LEU:CD1	2.20	0.47
9:I:38:PHE:C	9:I:44:ARG:HG2	2.39	0.47
10:J:5:ARG:HD3	10:J:5:ARG:N	2.30	0.47
10:J:48:ARG:HA	10:J:66:GLU:HA	1.95	0.47
10:J:57:VAL:HG13	10:J:58:ASN:CG	2.40	0.47
11:K:15:VAL:HG21	11:K:41:LEU:CD1	2.44	0.47
11:K:121:ARG:HG3	11:K:121:ARG:HH11	1.79	0.47
14:N:59:GLN:N	14:N:59:GLN:NE2	2.61	0.47
14:N:79:SER:O	14:N:81:ILE:HD12	2.14	0.47
16:P:33:ILE:N	16:P:33:ILE:HD12	2.29	0.47
17:Q:56:ASP:C	17:Q:79:GLU:HB3	2.40	0.47
22:V:280:GLU:O	22:V:281:PHE:C	2.57	0.47
1:A:109:A:H4'	1:A:110:C:OP2	2.14	0.47
1:A:1416:G:P	22:V:345:ALA:O	2.73	0.47
1:A:1452:C:H4'	1:A:1453:G:H5''	1.96	0.47
1:A:1483:A:H2'	22:V:295:PHE:CE1	1.92	0.47
2:B:163:ILE:CG2	2:B:164:ASP:N	2.64	0.47
3:C:112:ALA:O	3:C:113:LYS:C	2.57	0.47
4:D:163:GLN:HB2	4:D:164:ARG:HH12	1.80	0.47
4:D:176:LYS:O	4:D:177:MET:C	2.57	0.47
6:F:18:VAL:HG21	6:F:58:HIS:CG	2.49	0.47
6:F:70:VAL:HA	6:F:73:GLU:HG3	1.96	0.47
11:K:21:HIS:CE1	11:K:34:THR:HG21	2.50	0.47
11:K:92:ARG:NH2	11:K:111:ASP:OD1	2.47	0.47
14:N:72:PHE:CE1	14:N:77:GLY:HA2	2.49	0.47
15:O:28:VAL:HG22	15:O:65:LEU:HB3	1.96	0.47
16:P:39:PHE:HE2	16:P:70:ARG:HH21	1.63	0.47
20:T:4:LYS:HD2	20:T:5:SER:H	1.79	0.47
1:A:87:C:H2'	1:A:88:U:C4'	2.44	0.47
1:A:333:U:H2'	1:A:334:C:C6	2.49	0.47
1:A:577:G:O2'	1:A:578:C:H5'	2.14	0.47

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:1131:G:O2'	1:A:1132:C:H5'	2.15	0.47
1:A:1348:U:H4'	9:I:121:ARG:HH11	1.79	0.47
1:A:1352:C:H2'	1:A:1353:G:C8	2.50	0.47
1:A:1418:A:C4	22:V:292:THR:OG1	2.68	0.47
1:A:1448:C:H2'	1:A:1449:C:H6	1.79	0.47
3:C:11:LEU:HD22	3:C:17:TRP:CD1	2.50	0.47
3:C:139:ASN:O	3:C:143:LEU:HD22	2.14	0.47
6:F:5:GLU:HG3	6:F:63:ASN:OD1	2.15	0.47
6:F:38:ARG:HH21	6:F:96:VAL:CG1	2.28	0.47
8:H:64:TYR:CA	8:H:70:VAL:HG23	2.45	0.47
10:J:41:PRO:O	10:J:71:LEU:HD13	2.13	0.47
16:P:26:ASN:OD1	16:P:31:ARG:HB3	2.14	0.47
16:P:33:ILE:HD12	16:P:33:ILE:H	1.79	0.47
16:P:39:PHE:CE1	16:P:74:LEU:HD22	2.49	0.47
1:A:21:G:H2'	1:A:22:G:C8	2.49	0.47
1:A:88:U:O2'	1:A:89:U:C6	2.68	0.47
1:A:147:G:H2'	1:A:148:G:C8	2.49	0.47
1:A:426:U:H2'	1:A:427:U:C6	2.49	0.47
1:A:437:U:H5''	4:D:151:GLN:NE2	2.29	0.47
1:A:649:A:H2'	1:A:650:G:O4'	2.14	0.47
1:A:662:U:O2'	1:A:836:G:H5''	2.15	0.47
1:A:674:G:H2'	1:A:675:A:C8	2.49	0.47
1:A:708:C:H2'	1:A:709:U:H6	1.80	0.47
1:A:751:U:H4'	15:O:23:SER:HA	1.97	0.47
1:A:948:C:O2'	1:A:949:A:H5'	2.15	0.47
1:A:1071:C:O2'	1:A:1072:G:H5'	2.15	0.47
1:A:1294:G:H2'	1:A:1295:U:H6	1.79	0.47
1:A:1376:U:H2'	1:A:1377:A:C8	2.49	0.47
1:A:1484:C:H6	22:V:337:TYR:HA	1.80	0.47
2:B:60:ALA:HB3	2:B:223:GLY:HA3	1.95	0.47
2:B:82:ALA:O	2:B:217:ALA:HB2	2.15	0.47
2:B:158:ASP:O	2:B:181:PRO:HD2	2.15	0.47
3:C:90:VAL:HA	3:C:93:ILE:HG21	1.97	0.47
3:C:178:ARG:O	3:C:206:ILE:HA	2.14	0.47
4:D:58:GLN:CD	4:D:62:ARG:HG3	2.39	0.47
4:D:81:LEU:CD1	4:D:92:LEU:HD21	2.45	0.47
4:D:100:VAL:CG1	4:D:104:MET:HE3	2.45	0.47
7:G:58:LEU:HB3	7:G:59:GLU:OE2	2.14	0.47
7:G:94:ARG:CD	7:G:98:LEU:HD11	2.45	0.47
9:I:30:ASN:HD21	9:I:65:THR:HA	1.80	0.47
10:J:8:ILE:H	10:J:75:ASP:HB2	1.80	0.47

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
11:K:60:PHE:HA	11:K:63:GLN:OE1	2.14	0.47
11:K:82:GLU:HG3	11:K:107:THR:OG1	2.13	0.47
12:L:35:ARG:HH21	12:L:75:GLU:HB3	1.78	0.47
14:N:60:ARG:HE	14:N:62:ARG:CZ	2.28	0.47
19:S:3:SER:O	19:S:4:LEU:HG	2.14	0.47
19:S:4:LEU:O	19:S:6:LYS:N	2.48	0.47
19:S:43:MET:O	19:S:46:LEU:HB2	2.14	0.47
21:U:35:GLU:HB2	21:U:37:TYR:CZ	2.50	0.47
22:V:147:ASN:O	22:V:151:ARG:HG3	2.15	0.47
1:A:591:U:H2'	1:A:592:G:H8	1.79	0.47
1:A:894:G:O2'	1:A:895:G:H5'	2.15	0.47
1:A:1060:U:C5'	10:J:53:ILE:HG22	2.42	0.47
1:A:1180:A:P	9:I:98:ARG:HH22	2.38	0.47
2:B:156:LEU:H	2:B:156:LEU:HD12	1.80	0.47
3:C:8:GLY:HA2	3:C:11:LEU:CG	2.41	0.47
3:C:71:ARG:O	3:C:75:VAL:HG23	2.15	0.47
3:C:75:VAL:O	3:C:75:VAL:HG12	2.14	0.47
3:C:120:THR:HA	3:C:123:LEU:HD12	1.96	0.47
6:F:71:ILE:HG13	6:F:72:ASP:N	2.29	0.47
7:G:87:PRO:CG	7:G:151:ALA:HB2	2.41	0.47
10:J:6:ILE:HG12	10:J:102:LEU:CD1	2.45	0.47
22:V:139:ALA:HB1	22:V:169:LYS:HD2	1.97	0.47
1:A:301:G:H2'	1:A:302:G:H8	1.79	0.47
1:A:608:A:H2'	1:A:609:A:O4'	2.14	0.47
1:A:901:A:H3'	1:A:902:G:O4'	2.15	0.47
1:A:975:A:H4'	1:A:976:G:OP2	2.14	0.47
1:A:1418:A:N1	22:V:292:THR:HG23	2.16	0.47
2:B:116:LEU:CD1	2:B:139:GLU:HB3	2.44	0.47
3:C:46:LEU:CD2	3:C:75:VAL:HG13	2.44	0.47
4:D:167:PRO:HB2	4:D:170:LEU:HG	1.97	0.47
8:H:34:ALA:HB1	8:H:109:VAL:HG21	1.97	0.47
8:H:111:THR:HG23	8:H:114:ALA:CB	2.44	0.47
10:J:22:THR:CG2	10:J:23:ALA:N	2.77	0.47
13:M:11:HIS:O	13:M:12:LYS:HG2	2.14	0.47
17:Q:22:VAL:HG21	17:Q:58:VAL:HG21	1.96	0.47
17:Q:24:ILE:O	17:Q:40:THR:HA	2.14	0.47
19:S:37:SER:O	19:S:69:LYS:HG2	2.15	0.47
1:A:34:C:H2'	1:A:35:G:H8	1.79	0.47
1:A:551:U:O2'	1:A:552:U:H5'	2.15	0.47
1:A:1096:C:H2'	1:A:1097:C:C6	2.50	0.47
1:A:1121:U:H2'	1:A:1122:U:O4'	2.15	0.47

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:B:13:VAL:HG11	2:B:207:ARG:HB2	1.95	0.47
2:B:160:LEU:O	2:B:182:VAL:HA	2.14	0.47
4:D:145:ARG:C	4:D:147:LYS:H	2.22	0.47
4:D:170:LEU:HD12	4:D:170:LEU:O	2.15	0.47
7:G:132:THR:HA	7:G:135:LYS:HB3	1.97	0.47
9:I:40:ARG:HA	9:I:44:ARG:HH21	1.80	0.47
12:L:50:LYS:HE2	12:L:50:LYS:N	2.30	0.47
13:M:21:ILE:HG22	13:M:64:VAL:CG1	2.41	0.47
16:P:29:ASN:N	16:P:29:ASN:ND2	2.60	0.47
20:T:17:ARG:C	20:T:17:ARG:HD2	2.40	0.47
21:U:40:PRO:O	21:U:42:THR:N	2.48	0.47
1:A:230:G:O2'	1:A:231:U:H5'	2.15	0.46
1:A:563:A:H2'	1:A:567:G:C8	2.50	0.46
1:A:1018:G:H2'	1:A:1019:A:H8	1.80	0.46
2:B:50:ASN:O	2:B:51:GLU:C	2.58	0.46
2:B:95:TRP:CH2	2:B:100:LEU:HB2	2.50	0.46
3:C:84:GLU:HA	3:C:87:ARG:NH2	2.30	0.46
5:E:19:ARG:O	5:E:20:VAL:HB	2.14	0.46
9:I:49:GLN:N	9:I:50:PRO:HD2	2.30	0.46
10:J:67:ILE:CG1	14:N:95:LEU:HD13	2.45	0.46
12:L:106:VAL:HG22	12:L:117:GLY:H	1.79	0.46
13:M:13:HIS:HB2	13:M:16:ILE:HG22	1.97	0.46
13:M:53:ASP:OD1	13:M:56:ARG:HD2	2.15	0.46
19:S:10:ILE:HD11	19:S:15:LEU:HD13	1.96	0.46
1:A:785:G:C5'	22:V:61:ASP:CB	2.92	0.46
1:A:785:G:O5'	22:V:70:HIS:O	2.33	0.46
1:A:1252:A:H2'	1:A:1253:G:H5''	1.98	0.46
1:A:1410:A:H2'	1:A:1411:C:C6	2.50	0.46
1:A:1482:G:O5'	22:V:313:ASP:CG	2.57	0.46
4:D:147:LYS:H	4:D:147:LYS:HE3	1.80	0.46
4:D:177:MET:H	4:D:177:MET:HE2	1.79	0.46
10:J:18:ILE:HD12	10:J:72:ARG:HG3	1.97	0.46
1:A:114:U:H2'	1:A:115:G:C8	2.50	0.46
1:A:844:G:N7	1:A:846:G:N3	2.63	0.46
1:A:1220:G:H4'	19:S:33:TRP:O	2.14	0.46
1:A:1446:A:C2'	1:A:1447:A:H5''	2.46	0.46
1:A:1517:G:H1'	22:V:118:PRO:C	2.41	0.46
1:A:1527:U:O2'	1:A:1528:U:H5'	2.16	0.46
3:C:171:ARG:NH1	3:C:173:PRO:HG3	2.31	0.46
5:E:33:THR:O	5:E:34:ALA:CB	2.61	0.46
7:G:139:ASP:HA	7:G:142:ARG:HH12	1.80	0.46

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
10:J:35:GLN:H	10:J:78:GLU:HB3	1.80	0.46
13:M:72:ILE:O	13:M:76:ILE:HG13	2.15	0.46
15:O:87:ARG:C	15:O:88:ARG:HG2	2.41	0.46
16:P:6:LEU:HD11	16:P:71:VAL:HB	1.96	0.46
16:P:40:ASN:HA	16:P:41:PRO:HD3	1.73	0.46
1:A:471:U:H2'	1:A:472:U:H6	1.80	0.46
1:A:1171:A:O2'	1:A:1172:C:H5'	2.15	0.46
2:B:128:LEU:HD13	2:B:129:THR:H	1.80	0.46
4:D:81:LEU:HD11	4:D:92:LEU:HD21	1.97	0.46
5:E:89:THR:HG21	5:E:134:ASN:HD21	1.81	0.46
6:F:9:MET:HE1	18:R:64:LEU:CA	2.45	0.46
7:G:110:ARG:HH12	7:G:122:GLU:N	2.13	0.46
7:G:130:LYS:N	7:G:134:VAL:HG11	2.31	0.46
9:I:3:ASN:HB3	9:I:4:GLN:H	1.54	0.46
9:I:44:ARG:O	9:I:47:VAL:HG22	2.15	0.46
20:T:57:VAL:C	20:T:60:GLN:HE22	2.22	0.46
21:U:3:ILE:HA	21:U:19:LYS:HE3	1.97	0.46
1:A:212:G:H2'	1:A:213:G:C8	2.47	0.46
1:A:244:U:O4	1:A:906:A:H1'	2.16	0.46
1:A:301:G:H2'	1:A:302:G:C8	2.50	0.46
1:A:812:G:H4'	1:A:812:G:OP1	2.16	0.46
1:A:1085:U:H3'	1:A:1086:U:C5	2.51	0.46
1:A:1315:U:H3'	1:A:1316:G:C8	2.51	0.46
1:A:1425:U:H2'	1:A:1426:G:H8	1.81	0.46
3:C:126:ARG:HH22	3:C:190:THR:CG2	2.28	0.46
8:H:31:LEU:O	8:H:35:ILE:HG13	2.15	0.46
11:K:109:ILE:O	11:K:110:THR:HG23	2.15	0.46
14:N:10:VAL:HG12	14:N:11:LYS:NZ	2.31	0.46
16:P:4:ILE:HA	16:P:20:VAL:O	2.15	0.46
20:T:70:LYS:HB2	20:T:70:LYS:NZ	2.31	0.46
22:V:64:SER:HB3	22:V:66:ASP:OD1	2.16	0.46
1:A:123:U:OP1	1:A:312:C:H5'	2.16	0.46
1:A:792:A:H1'	1:A:794:A:N7	2.31	0.46
1:A:950:U:H2'	1:A:951:G:C8	2.50	0.46
1:A:968:A:H5'	1:A:968:A:N3	2.31	0.46
1:A:1016:A:H5'	1:A:1218:C:H4'	1.97	0.46
1:A:1113:C:H2'	1:A:1114:C:H6	1.81	0.46
1:A:1410:A:H2'	1:A:1411:C:H6	1.81	0.46
1:A:1416:G:C2'	22:V:288:PRO:HD3	2.31	0.46
1:A:1494:G:H2'	1:A:1495:U:C6	2.50	0.46
3:C:14:VAL:HG23	3:C:15:LYS:HG2	1.98	0.46

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:C:17:TRP:C	3:C:19:SER:H	2.24	0.46
3:C:112:ALA:HB2	3:C:182:ASP:O	2.16	0.46
3:C:131:ARG:CZ	3:C:131:ARG:HB3	2.45	0.46
4:D:94:GLU:CA	4:D:103:ARG:HH22	2.27	0.46
6:F:42:TRP:CZ2	6:F:61:LEU:HD23	2.51	0.46
7:G:35:LYS:O	7:G:39:GLU:HG2	2.16	0.46
9:I:10:ARG:HA	9:I:77:ALA:HB1	1.98	0.46
9:I:71:ILE:O	9:I:72:SER:C	2.59	0.46
14:N:74:ARG:HD3	14:N:74:ARG:C	2.41	0.46
1:A:552:U:H2'	1:A:553:A:C8	2.51	0.46
1:A:647:C:O2'	1:A:648:A:H5'	2.16	0.46
1:A:1430:A:H2'	1:A:1431:A:O4'	2.16	0.46
2:B:119:GLN:HA	2:B:124:THR:HG23	1.97	0.46
4:D:64:TYR:N	4:D:64:TYR:CD1	2.84	0.46
10:J:57:VAL:HG22	10:J:58:ASN:H	1.80	0.46
20:T:74:HIS:O	20:T:78:LEU:HB2	2.16	0.46
1:A:26:A:H61	1:A:558:G:H1'	1.81	0.46
1:A:418:C:H2'	1:A:419:C:H6	1.79	0.46
1:A:489:C:H2'	1:A:490:C:C6	2.51	0.46
1:A:601:G:H2'	1:A:602:A:C8	2.51	0.46
1:A:692:U:H2'	1:A:694:A:OP2	2.16	0.46
1:A:1124:G:H5''	10:J:38:GLY:HA3	1.97	0.46
1:A:1231:G:H2'	1:A:1232:U:H6	1.80	0.46
1:A:1409:C:C4'	22:V:151:ARG:HE	2.27	0.46
3:C:77:GLY:O	3:C:79:LYS:N	2.46	0.46
5:E:35:LEU:HD23	5:E:36:THR:N	2.30	0.46
5:E:84:VAL:HG21	5:E:142:GLY:O	2.15	0.46
8:H:29:SER:HB3	8:H:32:LYS:CG	2.46	0.46
1:A:627:G:H2'	1:A:628:G:C8	2.51	0.46
1:A:628:G:H2'	1:A:629:A:C8	2.51	0.46
1:A:778:G:O2'	1:A:779:C:H5'	2.16	0.46
1:A:841:C:H6	1:A:843:U:OP1	1.99	0.46
1:A:916:U:H2'	1:A:917:G:H8	1.80	0.46
1:A:1085:U:H3'	1:A:1086:U:H5	1.80	0.46
1:A:1256:A:O4'	1:A:1278:G:N2	2.49	0.46
1:A:1272:G:H2'	1:A:1273:C:C6	2.50	0.46
1:A:1420:U:C1'	22:V:311:LYS:H	2.27	0.46
3:C:131:ARG:HB3	3:C:131:ARG:NH1	2.31	0.46
4:D:117:VAL:HG12	4:D:130:ASN:CA	2.45	0.46
7:G:22:LEU:O	7:G:26:VAL:HG13	2.14	0.46
10:J:33:GLY:HA3	10:J:83:THR:OG1	2.16	0.46

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
14:N:20:PHE:CG	14:N:24:ALA:HB2	2.51	0.46
17:Q:20:ILE:CG1	17:Q:45:VAL:HB	2.43	0.46
19:S:61:VAL:HA	19:S:65:MET:SD	2.56	0.46
21:U:5:VAL:CG2	21:U:19:LYS:HZ1	2.28	0.46
1:A:366:A:O2'	1:A:394:G:N2	2.49	0.46
1:A:1348:U:H4'	9:I:121:ARG:NH1	2.31	0.46
2:B:138:ARG:HA	2:B:141:GLU:CD	2.40	0.46
4:D:160:LEU:HD22	4:D:161:ALA:N	2.31	0.46
5:E:108:GLY:N	5:E:110:MET:SD	2.89	0.46
7:G:42:VAL:O	7:G:46:LEU:HB2	2.15	0.46
9:I:15:ALA:O	9:I:66:VAL:HA	2.16	0.46
10:J:80:THR:HG22	10:J:81:GLU:H	1.81	0.46
11:K:12:ARG:N	11:K:12:ARG:HD2	2.30	0.46
11:K:52:ARG:HD2	11:K:53:GLY:H	1.80	0.46
14:N:45:LEU:O	14:N:48:GLN:HB3	2.16	0.46
15:O:78:THR:HA	15:O:81:ILE:CG1	2.46	0.46
19:S:18:VAL:HB	19:S:43:MET:HE3	1.97	0.46
19:S:31:ARG:HG3	19:S:56:HIS:HE2	1.81	0.46
20:T:23:ARG:HB2	20:T:65:LEU:HD11	1.98	0.46
1:A:123:U:H5''	1:A:311:C:O2'	2.16	0.45
1:A:155:A:H2'	1:A:156:C:O4'	2.16	0.45
1:A:299:G:H2'	1:A:300:A:C8	2.51	0.45
1:A:325:A:H2'	1:A:326:G:O4'	2.16	0.45
1:A:389:A:H2'	1:A:389:A:N3	2.30	0.45
1:A:736:C:H2'	1:A:737:C:H6	1.82	0.45
1:A:767:A:H2'	1:A:768:A:C8	2.50	0.45
1:A:1348:U:H2'	1:A:1349:A:C8	2.51	0.45
1:A:1494:G:O2'	1:A:1495:U:H5'	2.15	0.45
3:C:5:HIS:O	3:C:9:ILE:HG22	2.16	0.45
3:C:40:GLN:CD	3:C:44:LYS:HD2	2.42	0.45
3:C:131:ARG:HH22	3:C:135:ARG:NH1	2.14	0.45
3:C:146:LYS:HG3	3:C:202:PHE:CD2	2.51	0.45
4:D:25:ARG:HD3	4:D:26:ALA:N	2.31	0.45
6:F:5:GLU:HA	6:F:63:ASN:HA	1.97	0.45
6:F:38:ARG:HH21	6:F:96:VAL:HG11	1.81	0.45
8:H:40:LYS:HA	8:H:45:ILE:HG13	1.98	0.45
12:L:49:ARG:HH12	12:L:88:ASP:CB	2.29	0.45
16:P:26:ASN:ND2	16:P:31:ARG:HB3	2.31	0.45
21:U:34:ARG:HD2	21:U:34:ARG:C	2.40	0.45
22:V:162:GLU:OE1	22:V:216:ARG:NH1	2.44	0.45
1:A:491:G:H2'	1:A:492:C:C6	2.51	0.45

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:950:U:H2'	1:A:951:G:H8	1.81	0.45
1:A:982:U:H4'	1:A:983:A:O4'	2.16	0.45
1:A:1010:U:H2'	1:A:1011:C:C6	2.52	0.45
2:B:56:LEU:O	2:B:59:ILE:HG13	2.16	0.45
3:C:24:ASN:O	3:C:26:LYS:N	2.49	0.45
4:D:54:LEU:C	4:D:54:LEU:HD13	2.41	0.45
4:D:117:VAL:HG22	4:D:122:ILE:HG13	1.97	0.45
7:G:102:TRP:N	7:G:102:TRP:CD1	2.83	0.45
10:J:80:THR:H	10:J:84:VAL:CG1	2.29	0.45
11:K:60:PHE:HA	11:K:63:GLN:CD	2.42	0.45
13:M:10:ASP:HA	13:M:44:ILE:HD11	1.98	0.45
1:A:234:C:H2'	1:A:235:C:C6	2.52	0.45
1:A:890:G:O2'	1:A:906:A:N6	2.49	0.45
1:A:1101:A:O2'	1:A:1102:A:OP2	2.33	0.45
1:A:1418:A:C2	22:V:302:LEU:HD21	2.52	0.45
3:C:131:ARG:HH22	3:C:135:ARG:CZ	2.29	0.45
4:D:36:ALA:C	4:D:38:GLY:H	2.24	0.45
8:H:51:GLU:O	8:H:56:PRO:HA	2.16	0.45
8:H:65:PHE:CD2	8:H:66:GLN:HG3	2.51	0.45
11:K:88:PRO:HD3	21:U:28:LEU:CD2	2.47	0.45
12:L:98:ARG:CB	12:L:116:TYR:HA	2.46	0.45
15:O:24:THR:HB	15:O:69:LEU:HD21	1.97	0.45
15:O:58:MET:HG2	15:O:58:MET:H	1.59	0.45
15:O:70:LYS:HZ1	15:O:74:VAL:CG1	2.22	0.45
1:A:22:G:H2'	1:A:23:C:C6	2.52	0.45
1:A:138:G:O2'	1:A:139:A:H5'	2.17	0.45
1:A:382:A:H2'	1:A:383:A:C8	2.51	0.45
1:A:620:C:C6	4:D:131:ILE:HD13	2.51	0.45
1:A:746:A:H2'	1:A:747:A:H8	1.80	0.45
1:A:844:G:H21	1:A:845:A:H62	1.61	0.45
1:A:947:G:H2'	1:A:948:C:C6	2.51	0.45
1:A:1057:G:H4'	3:C:194:VAL:O	2.15	0.45
1:A:1126:U:O2'	1:A:1280:A:H2'	2.17	0.45
1:A:1342:C:H5'	9:I:127:SER:HA	1.98	0.45
3:C:57:GLU:HB2	3:C:64:ARG:CB	2.45	0.45
3:C:90:VAL:HG21	3:C:98:ALA:HB3	1.97	0.45
4:D:201:GLU:OE2	5:E:104:ILE:HG22	2.17	0.45
5:E:25:LYS:C	5:E:25:LYS:HE2	2.41	0.45
5:E:82:HIS:CD2	8:H:95:MET:HG3	2.51	0.45
6:F:45:ARG:HG2	6:F:46:GLN:N	2.31	0.45
10:J:89:ARG:NE	10:J:89:ARG:HA	2.31	0.45

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
11:K:57:SER:O	11:K:58:THR:C	2.60	0.45
17:Q:66:LEU:HD11	17:Q:73:THR:HG22	1.98	0.45
1:A:1110:A:N7	1:A:1111:A:N6	2.58	0.45
1:A:1415:G:O6	22:V:343:SER:N	2.49	0.45
1:A:1438:G:C2'	1:A:1439:G:H5'	2.47	0.45
2:B:61:SER:HA	2:B:224:ARG:CA	2.42	0.45
2:B:102:ASN:OD1	2:B:105:THR:HG22	2.16	0.45
3:C:72:PRO:HG2	3:C:73:GLY:H	1.81	0.45
4:D:1:ALA:O	4:D:67:LEU:HD21	2.16	0.45
4:D:100:VAL:HA	4:D:103:ARG:HD2	1.99	0.45
11:K:22:ILE:HD13	11:K:95:THR:CG2	2.46	0.45
11:K:88:PRO:HA	11:K:92:ARG:CD	2.47	0.45
18:R:31:TYR:CG	18:R:54:LEU:HD21	2.52	0.45
18:R:64:LEU:C	18:R:66:LEU:H	2.23	0.45
1:A:167:A:O2'	1:A:168:G:H5'	2.16	0.45
1:A:373:A:H1'	1:A:481:G:H1'	1.99	0.45
1:A:429:U:H4'	1:A:430:A:O5'	2.16	0.45
1:A:512:U:H2'	1:A:513:C:C6	2.51	0.45
1:A:586:C:C2'	1:A:587:G:H5'	2.46	0.45
1:A:734:G:O2'	18:R:59:LYS:HD3	2.17	0.45
1:A:1056:U:O2'	1:A:1057:G:H5'	2.16	0.45
1:A:1148:U:H5''	9:I:8:THR:HG23	1.99	0.45
1:A:1404:C:H2'	1:A:1405:G:C8	2.52	0.45
4:D:29:THR:OG1	4:D:30:LYS:HD3	2.17	0.45
9:I:56:MET:HE2	9:I:56:MET:HB2	1.76	0.45
9:I:57:VAL:C	9:I:59:LYS:H	2.25	0.45
10:J:71:LEU:HD12	10:J:71:LEU:N	2.27	0.45
10:J:89:ARG:HA	10:J:89:ARG:CZ	2.47	0.45
14:N:5:MET:C	14:N:7:ALA:H	2.24	0.45
15:O:27:GLN:O	15:O:31:LEU:HD23	2.16	0.45
17:Q:19:SER:O	17:Q:20:ILE:HG23	2.16	0.45
1:A:22:G:O2'	1:A:23:C:H5'	2.16	0.45
1:A:448:A:H2'	1:A:449:G:C8	2.51	0.45
1:A:471:U:H2'	1:A:472:U:C6	2.52	0.45
1:A:927:G:O2'	1:A:928:G:H5'	2.17	0.45
1:A:1314:C:H2'	1:A:1315:U:H6	1.81	0.45
2:B:73:ARG:C	2:B:75:ALA:H	2.25	0.45
2:B:113:LEU:CG	2:B:143:LEU:HB3	2.47	0.45
3:C:59:PRO:HD2	3:C:62:SER:O	2.17	0.45
4:D:32:LYS:HB3	4:D:35:GLN:HE21	1.81	0.45
5:E:45:VAL:HG12	5:E:116:VAL:HG23	1.98	0.45

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
8:H:49:LYS:CG	8:H:50:VAL:H	2.27	0.45
10:J:73:LEU:CD1	10:J:75:ASP:HB3	2.47	0.45
12:L:106:VAL:CG1	12:L:116:TYR:HB3	2.47	0.45
15:O:6:ALA:O	15:O:10:ILE:HG22	2.16	0.45
15:O:11:VAL:HA	15:O:26:VAL:HG13	1.98	0.45
1:A:132:C:H5''	20:T:68:LYS:CE	2.47	0.45
1:A:143:A:H2	1:A:220:G:H22	1.65	0.45
1:A:664:G:H5''	18:R:52:ARG:CZ	2.46	0.45
1:A:734:G:N2	18:R:63:TYR:CE2	2.84	0.45
1:A:834:U:H2'	1:A:835:U:C6	2.52	0.45
1:A:1110:A:H2'	1:A:1111:A:H5'	1.99	0.45
1:A:1320:C:H2'	1:A:1321:U:O4'	2.16	0.45
2:B:132:GLU:HG3	2:B:136:ARG:HD2	1.99	0.45
3:C:176:THR:O	3:C:179:ALA:HB3	2.17	0.45
7:G:125:ASP:HB3	7:G:130:LYS:CB	2.27	0.45
9:I:51:LEU:HD12	9:I:51:LEU:N	2.32	0.45
9:I:66:VAL:CG2	9:I:74:GLN:HG3	2.47	0.45
10:J:87:LEU:H	10:J:88:MET:CE	2.30	0.45
12:L:119:LYS:O	12:L:119:LYS:HG3	2.16	0.45
14:N:25:GLU:O	14:N:29:ILE:HG13	2.16	0.45
14:N:50:LEU:CG	14:N:51:PRO:HD3	2.46	0.45
20:T:54:GLN:N	20:T:55:PRO:CD	2.80	0.45
22:V:200:SER:O	22:V:203:GLN:NE2	2.50	0.45
1:A:295:C:H2'	1:A:296:U:C6	2.52	0.45
1:A:473:U:C2	1:A:474:G:N7	2.85	0.45
1:A:501:C:O2'	1:A:502:A:H5'	2.17	0.45
1:A:657:U:H4'	15:O:27:GLN:HG3	1.99	0.45
1:A:687:A:C2	1:A:704:A:C5	3.05	0.45
1:A:1038:C:H2'	1:A:1039:G:C8	2.47	0.45
1:A:1239:A:H1'	1:A:1241:G:C5	2.52	0.45
3:C:19:SER:HB2	3:C:39:ARG:HH22	1.82	0.45
5:E:57:ALA:O	5:E:60:GLN:HB2	2.16	0.45
8:H:98:LEU:H	8:H:98:LEU:HD12	1.82	0.45
10:J:88:MET:HE3	10:J:88:MET:H	1.82	0.45
12:L:58:ASN:ND2	12:L:58:ASN:N	2.64	0.45
13:M:3:ILE:HA	13:M:56:ARG:HG2	1.98	0.45
13:M:49:GLU:O	13:M:53:ASP:HB3	2.17	0.45
13:M:75:SER:O	13:M:78:ARG:HB3	2.16	0.45
16:P:43:ALA:H	16:P:46:LYS:HD2	1.81	0.45
20:T:70:LYS:HA	20:T:73:ARG:CZ	2.47	0.45
1:A:87:C:C2	1:A:88:U:H1'	2.51	0.45

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:820:U:H4'	1:A:821:G:OP2	2.16	0.45
1:A:913:A:H1'	1:A:914:A:O4'	2.17	0.45
1:A:1300:G:O2'	1:A:1301:U:P	2.75	0.45
1:A:1415:G:O6	22:V:341:LEU:O	2.34	0.45
1:A:1515:G:O2'	1:A:1516:G:H5'	2.17	0.45
2:B:33:ALA:HA	2:B:38:HIS:CA	2.39	0.45
3:C:33:ASP:O	3:C:37:LYS:HE2	2.16	0.45
4:D:2:ARG:NH1	4:D:114:ARG:HD2	2.32	0.45
4:D:77:GLU:OE1	4:D:80:ARG:NH2	2.48	0.45
4:D:191:SER:O	4:D:192:ALA:CB	2.65	0.45
9:I:38:PHE:O	9:I:39:GLY:C	2.59	0.45
11:K:70:ALA:HA	11:K:74:LYS:HD3	1.99	0.45
16:P:3:THR:HG22	16:P:66:THR:HB	1.98	0.45
16:P:12:LYS:O	16:P:14:ARG:HG3	2.17	0.45
17:Q:26:ARG:HH21	17:Q:39:ARG:NH2	2.15	0.45
18:R:64:LEU:C	18:R:66:LEU:N	2.74	0.45
1:A:652:U:H1'	1:A:653:U:C5	2.52	0.44
1:A:766:A:H2	1:A:1525:G:N3	2.15	0.44
1:A:1211:U:H1'	1:A:1213:A:C2	2.52	0.44
1:A:1238:A:C8	1:A:1303:C:H1'	2.52	0.44
1:A:1306:A:N6	1:A:1331:G:H1'	2.32	0.44
13:M:77:LYS:O	13:M:81:ASP:N	2.44	0.44
13:M:78:ARG:HE	13:M:79:LEU:HG	1.82	0.44
14:N:20:PHE:HA	14:N:24:ALA:CB	2.47	0.44
14:N:63:CYS:HB2	14:N:79:SER:OG	2.17	0.44
19:S:18:VAL:HG13	19:S:19:GLU:N	2.32	0.44
22:V:261:ALA:HA	22:V:275:SER:HB3	1.98	0.44
1:A:113:G:O4'	1:A:354:G:H4'	2.18	0.44
1:A:191:G:H2'	1:A:192:A:C8	2.52	0.44
1:A:692:U:H5	11:K:27:ASN:HD22	1.64	0.44
1:A:1096:C:H2'	1:A:1097:C:H6	1.82	0.44
1:A:1237:C:H4'	1:A:1334:G:N2	2.31	0.44
1:A:1386:G:O2'	1:A:1387:G:H5'	2.17	0.44
1:A:1486:G:O4'	22:V:344:MET:SD	2.76	0.44
2:B:21:TYR:O	2:B:22:TRP:O	2.34	0.44
2:B:128:LEU:CG	2:B:132:GLU:HG2	2.46	0.44
2:B:174:GLU:O	2:B:178:LEU:HD23	2.16	0.44
3:C:41:TYR:HA	3:C:44:LYS:HD3	1.98	0.44
4:D:104:MET:SD	4:D:179:GLY:HA3	2.57	0.44
4:D:152:SER:O	4:D:155:LYS:HG3	2.17	0.44
5:E:16:ALA:O	5:E:34:ALA:HA	2.17	0.44

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
5:E:85:LYS:HE2	5:E:92:ARG:HH11	1.82	0.44
6:F:7:VAL:HG23	6:F:60:VAL:O	2.17	0.44
9:I:64:ILE:HD13	9:I:78:ILE:HG21	1.98	0.44
10:J:80:THR:N	10:J:84:VAL:HG11	2.29	0.44
16:P:34:GLU:CD	16:P:56:ARG:HD3	2.42	0.44
16:P:42:ILE:CB	16:P:46:LYS:HD2	2.42	0.44
21:U:5:VAL:HG22	21:U:19:LYS:NZ	2.31	0.44
1:A:175:C:H2'	1:A:176:C:H6	1.83	0.44
1:A:284:C:H2'	1:A:285:C:H6	1.81	0.44
1:A:336:A:O2'	1:A:337:G:H5'	2.17	0.44
1:A:669:G:O2'	1:A:670:G:H5'	2.17	0.44
1:A:1051:C:H2'	1:A:1052:U:H6	1.82	0.44
1:A:1462:C:H2'	1:A:1463:U:C6	2.52	0.44
1:A:1482:G:C4'	22:V:312:HIS:O	2.65	0.44
2:B:67:LEU:HD23	2:B:160:LEU:HG	1.99	0.44
4:D:75:TYR:CE2	4:D:203:TYR:HB3	2.51	0.44
4:D:99:ASN:ND2	4:D:110:ARG:NE	2.64	0.44
9:I:79:ARG:O	9:I:83:THR:HG22	2.17	0.44
13:M:14:ALA:HB1	13:M:33:LEU:HD11	1.99	0.44
13:M:77:LYS:C	13:M:77:LYS:HD3	2.42	0.44
15:O:17:ASP:HB2	15:O:18:ALA:H	1.59	0.44
16:P:42:ILE:CG2	16:P:43:ALA:H	2.20	0.44
16:P:70:ARG:O	16:P:74:LEU:HG	2.17	0.44
18:R:49:LYS:HA	18:R:52:ARG:HD2	2.00	0.44
22:V:43:LEU:CD1	22:V:93:LYS:HD2	2.47	0.44
1:A:238:A:H3'	1:A:239:U:H5''	2.00	0.44
1:A:552:U:H5'	12:L:82:ARG:HH11	1.82	0.44
1:A:682:G:O2'	1:A:683:G:H5'	2.17	0.44
1:A:803:G:H2'	1:A:804:U:H6	1.82	0.44
1:A:981:U:H4'	14:N:60:ARG:CG	2.45	0.44
3:C:49:ALA:HB1	3:C:75:VAL:CG2	2.40	0.44
6:F:51:ILE:HG23	6:F:51:ILE:O	2.17	0.44
7:G:47:GLU:C	7:G:49:LEU:N	2.76	0.44
7:G:117:LEU:C	7:G:119:LEU:N	2.75	0.44
9:I:24:ASN:C	9:I:26:LYS:H	2.26	0.44
9:I:25:GLY:HA3	9:I:57:VAL:O	2.16	0.44
10:J:42:LEU:HB2	10:J:71:LEU:HD13	1.99	0.44
11:K:88:PRO:CD	21:U:28:LEU:HD22	2.48	0.44
11:K:100:ASN:ND2	21:U:12:ASP:CG	2.76	0.44
12:L:34:THR:HG21	12:L:53:ARG:CZ	2.47	0.44
12:L:80:LEU:HD13	12:L:101:LEU:HD11	1.99	0.44

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
17:Q:20:ILE:CD1	17:Q:45:VAL:HB	2.47	0.44
18:R:20:ILE:HG23	18:R:53:GLN:NE2	2.32	0.44
1:A:119:A:H4'	1:A:120:A:O4'	2.17	0.44
1:A:394:G:H2'	1:A:395:C:H6	1.82	0.44
1:A:463:U:H5'	1:A:464:U:OP2	2.17	0.44
1:A:922:G:H2'	1:A:923:A:H8	1.75	0.44
1:A:985:C:H2'	1:A:986:U:H6	1.82	0.44
1:A:1023:U:H2'	1:A:1024:G:H8	1.82	0.44
1:A:1483:A:C4'	22:V:298:PHE:CE2	2.91	0.44
1:A:1485:U:P	22:V:340:ILE:HD13	2.56	0.44
2:B:119:GLN:HE21	2:B:119:GLN:HB3	1.59	0.44
2:B:134:LEU:HA	2:B:137:THR:OG1	2.17	0.44
5:E:114:LEU:CD2	5:E:119:VAL:HG21	2.47	0.44
6:F:7:VAL:HB	6:F:61:LEU:HD22	2.00	0.44
6:F:10:VAL:CG1	6:F:83:ALA:HB1	2.48	0.44
7:G:71:THR:O	7:G:72:VAL:HG13	2.17	0.44
12:L:43:LYS:N	12:L:44:PRO:CD	2.80	0.44
15:O:15:GLY:C	15:O:17:ASP:H	2.26	0.44
17:Q:7:LEU:C	17:Q:9:GLY:H	2.24	0.44
22:V:173:LEU:HD12	22:V:178:MET:CE	2.35	0.44
1:A:71:A:N1	1:A:99:C:H1'	2.33	0.44
1:A:224:U:H2'	1:A:225:C:H6	1.82	0.44
1:A:232:G:H2'	1:A:233:C:O4'	2.17	0.44
1:A:323:U:H2'	1:A:324:G:O4'	2.18	0.44
1:A:383:A:H2'	1:A:384:G:O4'	2.18	0.44
1:A:397:A:H5'	1:A:398:U:OP1	2.18	0.44
1:A:432:A:H2'	1:A:433:G:H5'	1.99	0.44
1:A:454:G:O2'	1:A:455:G:H5'	2.17	0.44
1:A:661:G:O2'	1:A:662:U:H5'	2.18	0.44
1:A:1230:C:H2'	1:A:1231:G:H8	1.81	0.44
1:A:1238:A:H2	1:A:1241:G:H1'	1.82	0.44
3:C:71:ARG:HB3	3:C:74:ILE:CG2	2.47	0.44
4:D:18:LEU:HD22	4:D:63:ILE:HG12	2.00	0.44
4:D:137:SER:CB	4:D:138:PRO:HD2	2.47	0.44
5:E:81:GLN:OE1	5:E:81:GLN:N	2.50	0.44
6:F:7:VAL:HG11	18:R:64:LEU:HD21	1.99	0.44
6:F:79:ARG:HH21	6:F:87:SER:HB2	1.81	0.44
8:H:46:GLU:HA	8:H:63:LYS:HZ3	1.81	0.44
10:J:81:GLU:O	10:J:85:ASP:HB2	2.17	0.44
11:K:62:ALA:O	11:K:65:ALA:HB3	2.18	0.44
11:K:69:CYS:O	11:K:71:ASP:N	2.51	0.44

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
20:T:49:ALA:O	20:T:52:GLU:HB3	2.16	0.44
1:A:203:G:N2	1:A:205:A:H61	2.16	0.44
1:A:828:U:H2'	1:A:829:G:O5'	2.16	0.44
1:A:924:C:H2'	1:A:925:G:H8	1.83	0.44
1:A:1325:C:O2'	1:A:1326:U:H5'	2.18	0.44
1:A:1338:G:H2'	1:A:1339:A:H8	1.83	0.44
1:A:1350:A:H2'	1:A:1351:U:C6	2.53	0.44
1:A:1376:U:P	7:G:24:LYS:HD3	2.57	0.44
1:A:1458:G:H2'	1:A:1459:G:H8	1.83	0.44
2:B:59:ILE:CD1	2:B:66:ILE:HD11	2.48	0.44
2:B:147:LEU:O	2:B:151:LYS:HG3	2.17	0.44
3:C:90:VAL:HA	3:C:93:ILE:CG2	2.48	0.44
3:C:113:LYS:HE2	3:C:117:ASP:CG	2.42	0.44
4:D:113:ALA:O	4:D:117:VAL:HG23	2.16	0.44
5:E:33:THR:O	5:E:58:ALA:HB1	2.17	0.44
5:E:36:THR:O	5:E:48:GLY:N	2.51	0.44
5:E:49:TYR:HE2	5:E:133:ILE:HG12	1.83	0.44
17:Q:45:VAL:HG13	17:Q:60:ILE:CG2	2.48	0.44
20:T:10:ALA:O	20:T:13:SER:HB3	2.17	0.44
21:U:34:ARG:NH1	21:U:39:LYS:HE3	2.33	0.44
1:A:82:G:C6	1:A:88:U:O2	2.71	0.44
1:A:191:G:H2'	1:A:192:A:H8	1.83	0.44
1:A:238:A:C3'	1:A:239:U:H5''	2.48	0.44
1:A:251:G:H4'	1:A:252:U:C5'	2.48	0.44
1:A:601:G:H2'	1:A:602:A:H8	1.83	0.44
1:A:740:U:O3'	15:O:38:LEU:HD21	2.18	0.44
1:A:784:A:H4'	22:V:72:CYS:C	2.43	0.44
1:A:947:G:H2'	1:A:948:C:H6	1.81	0.44
1:A:1014:A:H2	1:A:1219:A:H1'	1.83	0.44
1:A:1197:A:P	1:A:1197:A:H3'	2.57	0.44
1:A:1246:A:H2'	1:A:1247:U:O4'	2.17	0.44
2:B:101:THR:HG23	2:B:102:ASN:N	2.32	0.44
2:B:119:GLN:O	2:B:125:PHE:HB3	2.17	0.44
3:C:40:GLN:O	3:C:44:LYS:HG3	2.18	0.44
5:E:105:ILE:HD12	5:E:123:LEU:HB3	1.99	0.44
7:G:148:LYS:HA	7:G:151:ALA:HB3	2.00	0.44
14:N:84:ARG:HH11	14:N:84:ARG:HG3	1.83	0.44
19:S:4:LEU:C	19:S:5:LYS:HE3	2.42	0.44
19:S:39:ILE:CG2	19:S:65:MET:HB3	2.46	0.44
1:A:26:A:N6	1:A:558:G:H1'	2.33	0.44
1:A:31:G:H2'	1:A:48:C:C5	2.53	0.44

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:65:A:N3	1:A:65:A:C2'	2.74	0.44
1:A:68:G:H5'	1:A:171:A:O2'	2.17	0.44
1:A:93:U:H2'	1:A:95:C:H5	1.83	0.44
1:A:532:A:N6	3:C:191:THR:HB	2.33	0.44
1:A:539:A:H2'	1:A:540:G:H8	1.78	0.44
1:A:813:U:O2'	1:A:814:A:H5'	2.18	0.44
1:A:844:G:H2'	1:A:845:A:H5''	2.00	0.44
1:A:878:A:H2'	1:A:879:C:C6	2.53	0.44
1:A:1029:U:H5''	1:A:1030:U:C5	2.47	0.44
2:B:101:THR:HG22	2:B:174:GLU:OE1	2.18	0.44
2:B:206:ILE:HD13	2:B:206:ILE:C	2.43	0.44
5:E:106:ALA:HB1	5:E:110:MET:HB2	2.00	0.44
6:F:45:ARG:C	6:F:56:LYS:HG2	2.42	0.44
7:G:55:LYS:HE2	7:G:57:GLU:OE1	2.18	0.44
9:I:48:ARG:HA	9:I:51:LEU:HD13	1.98	0.44
12:L:22:ALA:HB1	12:L:29:LYS:HG3	2.00	0.44
12:L:52:CYS:SG	12:L:66:ILE:HD11	2.57	0.44
15:O:34:GLN:HG2	15:O:58:MET:HE1	2.00	0.44
16:P:18:GLN:NE2	16:P:35:ARG:HH11	2.16	0.44
1:A:41:G:H2'	1:A:42:G:H8	1.83	0.43
1:A:267:C:OP2	17:Q:68:LYS:HD2	2.17	0.43
1:A:538:G:OP2	12:L:111:GLN:HB2	2.18	0.43
1:A:771:G:H2'	1:A:772:U:C6	2.53	0.43
1:A:1007:U:H2'	1:A:1008:U:C6	2.52	0.43
1:A:1435:G:H2'	1:A:1436:U:H6	1.81	0.43
4:D:125:ASN:HA	4:D:141:VAL:HG23	2.00	0.43
4:D:190:LEU:O	4:D:192:ALA:N	2.51	0.43
6:F:38:ARG:CB	6:F:63:ASN:HB2	2.43	0.43
8:H:11:THR:HG23	8:H:14:ARG:NH1	2.33	0.43
8:H:46:GLU:HB2	8:H:61:THR:OG1	2.18	0.43
9:I:115:VAL:HG21	10:J:62:ARG:HB2	1.98	0.43
11:K:30:ILE:HG22	11:K:45:THR:CA	2.41	0.43
16:P:7:ALA:HB1	16:P:29:ASN:CB	2.45	0.43
17:Q:30:HIS:H	17:Q:35:LYS:H	1.66	0.43
19:S:4:LEU:HD13	19:S:8:PRO:HA	1.99	0.43
19:S:31:ARG:NH2	19:S:55:GLN:HE22	2.16	0.43
19:S:48:ILE:HG22	19:S:49:ALA:N	2.33	0.43
19:S:62:THR:OG1	19:S:65:MET:HE3	2.18	0.43
1:A:8:A:H1'	5:E:107:GLY:HA2	2.00	0.43
1:A:210:C:H4'	1:A:211:G:H5''	2.00	0.43
1:A:442:G:H2'	1:A:443:C:C6	2.53	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:785:G:O3'	22:V:69:VAL:C	2.61	0.43
1:A:1082:A:O2'	1:A:1083:U:H5'	2.18	0.43
2:B:67:LEU:HD23	2:B:160:LEU:CG	2.48	0.43
3:C:112:ALA:HB1	3:C:199:VAL:CG2	2.47	0.43
3:C:135:ARG:C	3:C:137:VAL:N	2.77	0.43
5:E:89:THR:HG22	5:E:90:GLY:N	2.33	0.43
7:G:29:LEU:HD21	7:G:41:ILE:CG2	2.48	0.43
9:I:32:ARG:HG2	9:I:32:ARG:HH11	1.83	0.43
10:J:15:HIS:CD2	10:J:19:ASP:HB2	2.52	0.43
11:K:124:LYS:O	21:U:33:ARG:NH2	2.51	0.43
17:Q:62:GLU:HG3	17:Q:72:TRP:CH2	2.53	0.43
1:A:93:U:H2'	1:A:95:C:C5	2.54	0.43
1:A:284:C:H2'	1:A:285:C:C6	2.53	0.43
1:A:532:A:H62	3:C:191:THR:CB	2.31	0.43
1:A:538:G:H2'	1:A:539:A:C8	2.54	0.43
1:A:777:A:H2'	1:A:778:G:C8	2.53	0.43
1:A:893:C:H2'	1:A:894:G:H8	1.83	0.43
1:A:1008:U:H2'	1:A:1009:U:C4'	2.47	0.43
1:A:1254:A:H61	1:A:1283:U:H3	1.66	0.43
1:A:1416:G:N3	22:V:286:LEU:O	2.51	0.43
1:A:1505:G:H5''	1:A:1506:U:O5'	2.17	0.43
7:G:29:LEU:HD12	7:G:104:VAL:CG1	2.48	0.43
9:I:52:GLU:C	9:I:54:VAL:N	2.75	0.43
11:K:41:LEU:C	11:K:73:VAL:HG13	2.43	0.43
12:L:41:PRO:HG3	12:L:46:SER:O	2.18	0.43
13:M:79:LEU:HD22	13:M:86:ARG:NH2	2.33	0.43
14:N:92:ILE:HG21	14:N:95:LEU:HD23	1.99	0.43
19:S:44:ILE:HA	19:S:61:VAL:CG1	2.48	0.43
20:T:66:ILE:HG23	20:T:70:LYS:CD	2.48	0.43
21:U:5:VAL:HG13	21:U:15:LEU:HD23	2.00	0.43
21:U:28:LEU:C	21:U:30:GLU:N	2.76	0.43
22:V:43:LEU:HD13	22:V:43:LEU:C	2.43	0.43
1:A:15:G:O2'	5:E:21:SER:HB2	2.18	0.43
1:A:140:U:H2'	1:A:141:G:H8	1.82	0.43
1:A:408:A:OP1	4:D:109:THR:HG21	2.18	0.43
1:A:627:G:H2'	1:A:628:G:H8	1.83	0.43
1:A:709:U:H2'	1:A:710:G:H8	1.83	0.43
1:A:1124:G:C5'	10:J:38:GLY:HA3	2.49	0.43
4:D:154:VAL:HG13	4:D:155:LYS:N	2.33	0.43
4:D:169:TRP:CE2	4:D:185:PRO:HB3	2.54	0.43
7:G:47:GLU:C	7:G:49:LEU:H	2.25	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
8:H:35:ILE:O	8:H:39:LEU:HD12	2.18	0.43
9:I:51:LEU:C	9:I:56:MET:HG2	2.43	0.43
12:L:38:THR:HA	12:L:49:ARG:O	2.19	0.43
13:M:21:ILE:C	13:M:23:GLY:H	2.25	0.43
17:Q:80:LYS:HE3	17:Q:80:LYS:H	1.82	0.43
19:S:57:VAL:HG21	19:S:74:ALA:HB2	2.01	0.43
20:T:15:LYS:HD3	20:T:15:LYS:HA	1.86	0.43
20:T:19:HIS:O	20:T:23:ARG:HG2	2.18	0.43
1:A:611:C:H2'	1:A:612:C:H6	1.84	0.43
1:A:973:G:O2'	10:J:56:HIS:HA	2.18	0.43
1:A:1033:G:H3'	1:A:1034:G:H5''	2.01	0.43
1:A:1099:G:H2'	1:A:1100:C:O4'	2.19	0.43
2:B:123:GLY:C	2:B:125:PHE:H	2.27	0.43
2:B:167:HIS:C	2:B:169:HIS:H	2.25	0.43
4:D:109:THR:HG23	4:D:112:GLU:H	1.82	0.43
4:D:151:GLN:HB3	4:D:154:VAL:CG1	2.49	0.43
4:D:163:GLN:N	4:D:163:GLN:OE1	2.52	0.43
5:E:40:ASP:O	5:E:42:ASN:N	2.45	0.43
13:M:103:THR:O	13:M:104:ASN:C	2.61	0.43
15:O:52:ARG:HG3	15:O:55:LEU:HD23	1.99	0.43
16:P:68:SER:HB3	16:P:71:VAL:CG1	2.49	0.43
17:Q:17:GLU:C	17:Q:19:SER:N	2.76	0.43
19:S:29:PRO:HB3	19:S:47:THR:HB	2.01	0.43
19:S:62:THR:HG22	19:S:63:ASP:N	2.33	0.43
1:A:177:G:N3	1:A:177:G:O4'	2.51	0.43
1:A:392:C:H2'	1:A:393:A:H8	1.82	0.43
1:A:554:A:H2'	1:A:555:U:C6	2.53	0.43
1:A:989:U:H2'	1:A:990:C:H6	1.83	0.43
1:A:1256:A:O2'	1:A:1257:A:H5''	2.18	0.43
1:A:1402:C:H2'	1:A:1403:C:O4'	2.19	0.43
5:E:17:VAL:HB	5:E:34:ALA:HB2	2.00	0.43
6:F:88:MET:HE2	6:F:89:VAL:N	2.33	0.43
11:K:15:VAL:O	11:K:16:SER:HB2	2.17	0.43
12:L:56:LEU:HB2	12:L:60:PHE:O	2.19	0.43
13:M:56:ARG:C	13:M:59:VAL:HG12	2.44	0.43
16:P:44:SER:O	16:P:45:GLU:C	2.60	0.43
21:U:11:PHE:O	21:U:11:PHE:HD1	2.02	0.43
1:A:25:C:H2'	1:A:26:A:C8	2.53	0.43
1:A:397:A:H3'	1:A:397:A:N3	2.33	0.43
1:A:489:C:H2'	1:A:490:C:H6	1.84	0.43
1:A:1291:U:O2'	1:A:1292:G:H5'	2.18	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:1348:U:OP1	9:I:111:GLU:HB2	2.18	0.43
2:B:27:LYS:C	2:B:29:PHE:N	2.77	0.43
2:B:145:ASN:HD22	2:B:145:ASN:HA	1.55	0.43
4:D:169:TRP:CB	4:D:183:ARG:HH21	2.32	0.43
4:D:169:TRP:O	4:D:182:LYS:HB3	2.19	0.43
5:E:155:LYS:HG3	8:H:65:PHE:HB2	2.01	0.43
6:F:68:GLN:O	6:F:71:ILE:HG13	2.19	0.43
9:I:21:LYS:O	9:I:60:LEU:HB2	2.19	0.43
9:I:66:VAL:HG22	9:I:67:LYS:N	2.34	0.43
10:J:40:ILE:CG1	10:J:73:LEU:HB3	2.46	0.43
15:O:39:GLN:HE21	15:O:39:GLN:HB2	1.65	0.43
19:S:68:HIS:NE2	19:S:72:GLU:HG3	2.33	0.43
1:A:224:U:H2'	1:A:225:C:C6	2.54	0.43
1:A:384:G:H2'	1:A:385:C:H6	1.80	0.43
1:A:473:U:C2	1:A:474:G:C8	3.07	0.43
1:A:1226:C:H4'	1:A:1227:A:OP1	2.18	0.43
1:A:1286:U:OP1	1:A:1286:U:C2	2.72	0.43
1:A:1356:G:H2'	1:A:1357:A:C8	2.54	0.43
1:A:1420:U:C6	22:V:309:ASP:O	2.72	0.43
1:A:1482:G:O5'	22:V:338:HIS:CG	2.64	0.43
1:A:1485:U:O2'	22:V:286:LEU:HG	2.18	0.43
1:A:1486:G:H2'	1:A:1487:G:C8	2.53	0.43
2:B:31:PHE:CD2	2:B:41:ASN:HA	2.53	0.43
3:C:24:ASN:O	3:C:25:THR:C	2.62	0.43
4:D:14:GLU:OE2	4:D:58:GLN:HG3	2.18	0.43
4:D:84:ASN:ND2	4:D:86:GLY:N	2.67	0.43
4:D:100:VAL:HA	4:D:103:ARG:CD	2.48	0.43
4:D:145:ARG:C	4:D:147:LYS:N	2.76	0.43
5:E:145:ASN:O	5:E:146:MET:C	2.62	0.43
9:I:24:ASN:CA	9:I:26:LYS:HZ3	2.32	0.43
12:L:23:LEU:C	12:L:25:ALA:N	2.77	0.43
12:L:117:GLY:O	12:L:118:VAL:HG13	2.19	0.43
1:A:261:U:H2'	1:A:263:A:OP2	2.18	0.43
1:A:493:A:N3	1:A:493:A:O4'	2.52	0.43
1:A:564:C:H1'	17:Q:32:ILE:O	2.19	0.43
1:A:644:U:O2'	1:A:645:G:H5'	2.19	0.43
1:A:1073:U:O3'	2:B:104:LYS:NZ	2.51	0.43
1:A:1452:C:H5'	1:A:1453:G:N9	2.34	0.43
3:C:88:LYS:C	3:C:88:LYS:HD3	2.44	0.43
5:E:56:PRO:O	5:E:59:ILE:HG22	2.19	0.43
7:G:94:ARG:NH1	7:G:98:LEU:HD21	2.34	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
9:I:20:ILE:HA	9:I:61:ASP:O	2.19	0.43
9:I:33:SER:C	9:I:35:GLU:N	2.77	0.43
10:J:7:ARG:O	10:J:8:ILE:HD13	2.18	0.43
11:K:57:SER:O	11:K:90:PRO:HG2	2.19	0.43
12:L:56:LEU:HB3	12:L:58:ASN:HD21	1.83	0.43
16:P:7:ALA:O	16:P:17:TYR:HA	2.17	0.43
17:Q:28:VAL:O	17:Q:28:VAL:HG13	2.19	0.43
1:A:19:A:OP1	5:E:134:ASN:ND2	2.51	0.43
1:A:95:C:O2	1:A:95:C:C2'	2.67	0.43
1:A:382:A:O2'	1:A:383:A:H5'	2.19	0.43
1:A:697:U:H2'	1:A:698:G:H5'	2.00	0.43
1:A:811:C:O2'	1:A:901:A:N1	2.48	0.43
1:A:914:A:O2'	1:A:915:A:H5'	2.19	0.43
1:A:1050:G:C2'	1:A:1051:C:H5'	2.49	0.43
1:A:1081:A:OP1	5:E:22:LYS:HB2	2.19	0.43
1:A:1114:C:H2'	1:A:1115:U:O4'	2.19	0.43
1:A:1262:C:H2'	1:A:1263:C:O4'	2.19	0.43
2:B:53:LEU:HD12	2:B:219:THR:CG2	2.49	0.43
2:B:94:ARG:H	2:B:94:ARG:HG2	1.50	0.43
2:B:116:LEU:HD22	2:B:140:LEU:HG	1.99	0.43
2:B:164:ASP:OD2	2:B:203:ASP:HB2	2.18	0.43
3:C:21:TRP:CH2	14:N:93:PRO:HG3	2.54	0.43
3:C:39:ARG:HH21	3:C:56:ILE:HD12	1.84	0.43
5:E:30:PHE:CD1	5:E:30:PHE:N	2.87	0.43
5:E:35:LEU:HD12	5:E:133:ILE:HA	2.01	0.43
5:E:114:LEU:HD13	5:E:122:VAL:HG21	2.01	0.43
8:H:69:ALA:HB3	8:H:72:GLU:OE2	2.19	0.43
9:I:9:GLY:HA2	9:I:80:HIS:CD2	2.53	0.43
11:K:81:LEU:O	11:K:106:ILE:HA	2.19	0.43
16:P:75:ILE:O	16:P:78:VAL:HG12	2.19	0.43
17:Q:23:ALA:C	17:Q:24:ILE:HD12	2.44	0.43
1:A:473:U:N3	1:A:474:G:N7	2.67	0.42
1:A:847:G:H2'	1:A:848:C:C6	2.54	0.42
1:A:969:A:H2'	1:A:970:C:O2	2.18	0.42
1:A:970:C:C5	1:A:1231:G:H1'	2.54	0.42
1:A:1386:G:H2'	1:A:1387:G:H8	1.83	0.42
2:B:165:ALA:CB	2:B:186:VAL:HG12	2.48	0.42
3:C:40:GLN:OE1	3:C:44:LYS:HD2	2.19	0.42
4:D:12:ARG:HG3	4:D:33:ILE:HA	2.00	0.42
10:J:88:MET:SD	10:J:88:MET:N	2.85	0.42
12:L:14:LYS:NZ	12:L:17:LYS:HE2	2.34	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
13:M:63:VAL:HB	13:M:68:LEU:HG	2.00	0.42
13:M:106:ARG:NH1	13:M:106:ARG:HG2	2.32	0.42
16:P:22:ALA:HB2	16:P:32:PHE:HB3	2.01	0.42
17:Q:50:ASN:C	17:Q:52:CYS:H	2.27	0.42
19:S:16:LYS:HA	19:S:16:LYS:HD3	1.83	0.42
20:T:20:ASN:O	20:T:65:LEU:HD21	2.18	0.42
22:V:66:ASP:OD1	22:V:68:GLU:CG	2.66	0.42
1:A:22:G:H2'	1:A:23:C:H6	1.83	0.42
1:A:321:A:H2'	1:A:322:C:H6	1.85	0.42
1:A:394:G:H2'	1:A:395:C:C6	2.53	0.42
1:A:402:G:H2'	1:A:403:C:H6	1.84	0.42
1:A:619:U:C2	4:D:131:ILE:HD12	2.53	0.42
1:A:1389:C:H2'	1:A:1390:U:C6	2.54	0.42
2:B:18:GLN:HG2	2:B:189:ASN:CG	2.44	0.42
3:C:111:ASP:OD2	3:C:114:LEU:HG	2.19	0.42
3:C:156:LEU:H	3:C:156:LEU:CD1	2.31	0.42
4:D:147:LYS:HE3	4:D:147:LYS:N	2.34	0.42
9:I:26:LYS:H	9:I:26:LYS:HG2	1.56	0.42
9:I:64:ILE:HD13	9:I:78:ILE:CG2	2.49	0.42
9:I:82:ILE:HG22	9:I:86:LEU:HD21	2.01	0.42
10:J:59:LYS:C	10:J:61:ALA:H	2.28	0.42
11:K:28:ASN:HD21	11:K:30:ILE:HG23	1.83	0.42
17:Q:12:VAL:HG13	17:Q:21:VAL:HB	2.01	0.42
19:S:16:LYS:HA	19:S:19:GLU:OE1	2.19	0.42
21:U:20:ARG:O	21:U:20:ARG:HG2	2.20	0.42
1:A:242:G:H2'	1:A:243:A:H5''	2.01	0.42
1:A:474:G:H2'	1:A:475:C:H6	1.83	0.42
1:A:785:G:O2'	22:V:69:VAL:O	2.31	0.42
1:A:796:C:O2'	1:A:797:C:H5'	2.19	0.42
1:A:818:G:O2'	1:A:819:A:H5''	2.19	0.42
1:A:861:G:H2'	1:A:862:C:H6	1.84	0.42
1:A:1518:A:P	22:V:124:VAL:HG22	2.54	0.42
4:D:89:LEU:O	4:D:92:LEU:HB2	2.18	0.42
6:F:53:LYS:HZ2	6:F:54:LEU:N	2.17	0.42
14:N:73:LEU:O	14:N:75:LYS:N	2.52	0.42
16:P:25:ARG:HD3	16:P:25:ARG:N	2.34	0.42
16:P:46:LYS:HD3	16:P:46:LYS:N	2.19	0.42
18:R:64:LEU:HB3	18:R:66:LEU:HG	2.01	0.42
1:A:144:G:H2'	1:A:145:G:O4'	2.19	0.42
1:A:236:A:H2'	1:A:237:G:H8	1.85	0.42
1:A:488:C:H2'	1:A:489:C:H6	1.84	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:599:C:O2'	1:A:600:A:H5'	2.19	0.42
1:A:740:U:O2'	1:A:741:G:H5'	2.19	0.42
1:A:920:U:O2'	1:A:921:U:H5'	2.19	0.42
1:A:972:C:H4'	10:J:59:LYS:HB3	2.02	0.42
1:A:1052:U:H5'	1:A:1053:G:OP2	2.19	0.42
1:A:1065:U:H1'	1:A:1066:C:OP2	2.19	0.42
1:A:1157:A:H4'	1:A:1158:C:O5'	2.19	0.42
1:A:1361:G:H2'	1:A:1362:A:C5'	2.49	0.42
1:A:1418:A:C4	22:V:292:THR:CB	3.02	0.42
1:A:1470:U:O2'	1:A:1471:U:H5'	2.20	0.42
2:B:58:LYS:HD3	2:B:62:ARG:NH2	2.34	0.42
2:B:61:SER:HB3	2:B:223:GLY:O	2.20	0.42
2:B:110:ILE:O	2:B:113:LEU:HB3	2.18	0.42
3:C:15:LYS:HA	3:C:16:PRO:HD2	1.97	0.42
3:C:102:ILE:C	3:C:102:ILE:HD12	2.45	0.42
4:D:54:LEU:HA	4:D:57:LYS:HB3	2.01	0.42
4:D:132:ALA:C	4:D:134:TYR:H	2.27	0.42
5:E:105:ILE:HB	5:E:123:LEU:CA	2.42	0.42
6:F:11:HIS:CG	6:F:12:PRO:HD2	2.54	0.42
9:I:86:LEU:HD13	9:I:86:LEU:N	2.34	0.42
10:J:9:ARG:C	10:J:10:LEU:HD12	2.44	0.42
11:K:26:PHE:HE1	11:K:88:PRO:HG2	1.85	0.42
12:L:21:PRO:HD2	12:L:94:TYR:OH	2.19	0.42
12:L:34:THR:HB	12:L:53:ARG:HB2	2.02	0.42
12:L:101:LEU:HB3	12:L:102:ASP:H	1.61	0.42
15:O:46:LYS:HB2	15:O:46:LYS:HE3	1.89	0.42
16:P:21:VAL:HG21	16:P:60:TRP:CD1	2.53	0.42
20:T:57:VAL:O	20:T:61:ALA:HB2	2.19	0.42
1:A:255:G:H2'	1:A:256:U:H6	1.84	0.42
1:A:421:U:H5'	1:A:422:C:C5	2.55	0.42
1:A:598:U:H2'	1:A:599:C:C6	2.53	0.42
1:A:680:C:H2'	1:A:681:A:H8	1.85	0.42
1:A:839:C:O2'	1:A:840:C:H5'	2.19	0.42
1:A:1113:C:H2'	1:A:1114:C:C6	2.55	0.42
1:A:1261:A:H2'	1:A:1262:C:O4'	2.20	0.42
2:B:79:VAL:HG11	2:B:92:ASN:HB3	2.01	0.42
2:B:80:LYS:HB3	2:B:90:PHE:CE2	2.54	0.42
5:E:70:MET:HB3	5:E:71:ILE:H	1.65	0.42
5:E:109:ALA:O	5:E:113:VAL:HG13	2.18	0.42
5:E:137:ARG:HG2	5:E:137:ARG:H	1.72	0.42
9:I:10:ARG:HA	9:I:77:ALA:CB	2.50	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
17:Q:11:VAL:CA	17:Q:22:VAL:HG22	2.44	0.42
17:Q:67:SER:HB2	17:Q:70:LYS:HB3	2.02	0.42
21:U:36:PHE:C	21:U:40:PRO:HD3	2.44	0.42
1:A:113:G:H21	1:A:353:A:H8	1.68	0.42
1:A:412:A:H1'	1:A:413:G:C8	2.43	0.42
1:A:478:A:H2'	1:A:479:U:O4'	2.19	0.42
1:A:512:U:O2'	1:A:513:C:H5'	2.19	0.42
1:A:552:U:H4'	12:L:82:ARG:HG2	2.01	0.42
1:A:585:G:O2'	1:A:586:C:H5'	2.20	0.42
1:A:818:G:C3'	1:A:819:A:H5''	2.50	0.42
1:A:923:A:H2'	1:A:924:C:C6	2.55	0.42
1:A:1273:C:H2'	1:A:1274:A:O4'	2.20	0.42
1:A:1519:A:H3'	1:A:1520:C:C5'	2.49	0.42
3:C:185:THR:O	3:C:186:SER:HB2	2.20	0.42
4:D:54:LEU:HA	4:D:202:LEU:HD22	2.00	0.42
4:D:84:ASN:OD1	5:E:101:GLY:HA2	2.19	0.42
9:I:11:ARG:NH1	9:I:106:ASP:OD2	2.53	0.42
9:I:61:ASP:C	9:I:62:LEU:HD22	2.45	0.42
12:L:41:PRO:HD3	12:L:47:ALA:O	2.19	0.42
13:M:47:LEU:HD23	13:M:51:GLN:C	2.44	0.42
21:U:5:VAL:O	21:U:6:ARG:HB2	2.19	0.42
1:A:408:A:OP1	4:D:111:ALA:HB3	2.19	0.42
1:A:462:G:H2'	1:A:463:U:C6	2.55	0.42
1:A:785:G:H4'	22:V:69:VAL:HG12	1.94	0.42
1:A:1057:G:H2'	1:A:1058:G:O4'	2.19	0.42
1:A:1299:A:N6	1:A:1302:C:H5	2.18	0.42
1:A:1374:A:H4'	7:G:30:MET:HE1	2.02	0.42
3:C:54:ILE:O	3:C:54:ILE:HG12	2.19	0.42
3:C:160:GLU:O	3:C:160:GLU:HG2	2.19	0.42
7:G:13:PRO:HA	7:G:23:ALA:HB2	2.00	0.42
9:I:83:THR:HG23	9:I:84:ARG:N	2.35	0.42
10:J:30:LYS:HG3	10:J:36:VAL:HB	2.01	0.42
10:J:48:ARG:HB3	10:J:66:GLU:HB3	2.01	0.42
12:L:6:LEU:O	12:L:10:PRO:HG3	2.19	0.42
14:N:1:ALA:O	14:N:2:LYS:HB2	2.20	0.42
14:N:79:SER:O	14:N:80:ARG:C	2.62	0.42
1:A:41:G:H2'	1:A:42:G:C8	2.54	0.42
1:A:343:U:H2'	1:A:345:C:C5	2.55	0.42
1:A:591:U:H2'	1:A:592:G:C8	2.55	0.42
1:A:783:C:H5'	22:V:58:MET:CB	2.43	0.42
1:A:787:A:O2'	1:A:788:U:H5'	2.20	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:893:C:H2'	1:A:894:G:C8	2.55	0.42
1:A:1454:G:H2'	1:A:1455:G:O4'	2.19	0.42
2:B:9:LEU:CD1	2:B:11:ALA:HB3	2.50	0.42
2:B:71:THR:HG22	2:B:94:ARG:HH21	1.84	0.42
2:B:187:ASP:H	2:B:190:SER:HB2	1.85	0.42
3:C:41:TYR:HA	3:C:44:LYS:CD	2.49	0.42
5:E:37:VAL:HA	5:E:47:PHE:HA	2.01	0.42
5:E:101:GLY:H	5:E:121:ASN:ND2	2.14	0.42
6:F:7:VAL:HA	6:F:60:VAL:O	2.20	0.42
7:G:115:MET:CE	7:G:119:LEU:HB2	2.50	0.42
9:I:66:VAL:HG21	9:I:74:GLN:HG3	2.02	0.42
9:I:118:ARG:HH12	9:I:122:ARG:HE	1.67	0.42
14:N:12:ARG:NH2	14:N:58:ARG:HH12	2.18	0.42
14:N:68:ARG:HH11	14:N:70:HIS:HB2	1.85	0.42
22:V:75:ARG:CD	22:V:104:ILE:HD11	2.49	0.42
1:A:237:G:H2'	1:A:238:A:H8	1.84	0.42
1:A:415:A:N3	1:A:415:A:O4'	2.52	0.42
1:A:783:C:H4'	22:V:58:MET:CB	2.41	0.42
1:A:861:G:H2'	1:A:862:C:C6	2.55	0.42
1:A:1418:A:H2	22:V:292:THR:HG23	1.55	0.42
1:A:1506:U:H4'	11:K:128:VAL:C	2.44	0.42
3:C:4:VAL:HG21	3:C:9:ILE:HD13	2.02	0.42
6:F:40:GLU:CD	6:F:42:TRP:HE1	2.28	0.42
7:G:87:PRO:HB2	7:G:144:ALA:CB	2.50	0.42
11:K:22:ILE:HD13	11:K:95:THR:HG21	2.01	0.42
12:L:4:ASN:O	12:L:8:ARG:HG2	2.19	0.42
13:M:84:CYS:SG	13:M:86:ARG:HB3	2.59	0.42
14:N:20:PHE:HA	14:N:24:ALA:H	1.85	0.42
14:N:45:LEU:HD23	14:N:46:LYS:N	2.35	0.42
14:N:89:ARG:HB2	14:N:91:GLU:HG3	2.02	0.42
14:N:93:PRO:O	14:N:94:GLY:C	2.63	0.42
16:P:12:LYS:C	16:P:14:ARG:N	2.75	0.42
19:S:51:HIS:C	19:S:53:GLY:H	2.28	0.42
21:U:52:VAL:HG13	21:U:53:LYS:N	2.34	0.42
1:A:233:C:O2'	1:A:234:C:H5'	2.20	0.42
1:A:250:A:H1'	1:A:252:U:C5	2.55	0.42
1:A:405:U:OP2	4:D:114:ARG:NH2	2.53	0.42
1:A:605:U:H2'	1:A:606:G:C8	2.55	0.42
1:A:666:G:H5'	1:A:726:C:H1'	2.02	0.42
1:A:957:U:O2	1:A:959:A:H8	2.03	0.42
1:A:999:C:O2'	1:A:1000:A:H5'	2.20	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:1129:C:H5''	9:I:17:ARG:NH2	2.34	0.42
1:A:1416:G:C2	22:V:344:MET:CB	3.03	0.42
1:A:1484:C:O4'	22:V:295:PHE:CD1	2.49	0.42
1:A:1517:G:H2'	1:A:1518:A:C8	2.55	0.42
3:C:55:VAL:HG12	3:C:56:ILE:H	1.84	0.42
4:D:49:ASP:O	4:D:52:VAL:HG22	2.19	0.42
4:D:99:ASN:HB3	4:D:103:ARG:HH21	1.85	0.42
7:G:138:GLU:HA	7:G:141:HIS:HB2	2.02	0.42
9:I:56:MET:O	9:I:58:GLU:N	2.52	0.42
18:R:21:ASP:CG	18:R:22:TYR:N	2.77	0.42
19:S:20:LYS:O	19:S:20:LYS:HD2	2.18	0.42
21:U:3:ILE:HG23	21:U:19:LYS:HE3	2.01	0.42
21:U:3:ILE:CG1	21:U:19:LYS:HG2	2.46	0.42
1:A:29:U:H5'	1:A:296:U:OP1	2.19	0.41
1:A:162:A:H2'	1:A:163:C:O4'	2.20	0.41
1:A:237:G:H2'	1:A:238:A:C8	2.54	0.41
1:A:441:A:N6	1:A:493:A:H62	2.15	0.41
1:A:581:G:OP1	15:O:64:LYS:HE3	2.20	0.41
1:A:761:G:H2'	1:A:762:U:C6	2.55	0.41
1:A:924:C:HO2'	1:A:925:G:H5'	1.85	0.41
1:A:1244:G:H2'	1:A:1245:C:H6	1.81	0.41
1:A:1252:A:H2'	1:A:1253:G:C5'	2.49	0.41
1:A:1419:G:C5'	22:V:308:ARG:O	2.67	0.41
3:C:13:ILE:HG22	3:C:14:VAL:HG13	2.01	0.41
3:C:139:ASN:C	3:C:141:MET:N	2.76	0.41
4:D:36:ALA:C	4:D:38:GLY:N	2.77	0.41
4:D:159:GLU:HG3	4:D:160:LEU:HD13	2.01	0.41
8:H:24:VAL:HG13	8:H:60:LEU:HB2	2.00	0.41
9:I:25:GLY:HA2	9:I:60:LEU:O	2.19	0.41
9:I:51:LEU:HD21	9:I:62:LEU:HG	2.01	0.41
9:I:119:LYS:C	9:I:121:ARG:H	2.28	0.41
10:J:16:ARG:C	10:J:17:LEU:HD22	2.44	0.41
11:K:33:ILE:HG13	11:K:73:VAL:HG21	2.01	0.41
12:L:8:ARG:CZ	12:L:9:LYS:HE3	2.50	0.41
15:O:5:GLU:O	15:O:9:LYS:HB2	2.20	0.41
19:S:12:LEU:CD2	19:S:15:LEU:HD23	2.49	0.41
21:U:5:VAL:HG22	21:U:19:LYS:HZ1	1.85	0.41
1:A:373:A:C1'	1:A:481:G:H1'	2.50	0.41
1:A:432:A:C2'	1:A:433:G:H5'	2.49	0.41
1:A:691:G:H2'	1:A:692:U:C6	2.55	0.41
1:A:978:A:O2'	1:A:1322:C:H5	2.02	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:1324:A:H2'	1:A:1325:C:O4'	2.19	0.41
1:A:1419:G:H5'	22:V:309:ASP:HB3	1.93	0.41
2:B:42:LEU:O	2:B:46:VAL:HG12	2.20	0.41
2:B:121:GLN:NE2	2:B:122:ASP:HB2	2.35	0.41
7:G:44:SER:C	7:G:48:THR:HG23	2.45	0.41
9:I:33:SER:C	9:I:35:GLU:H	2.27	0.41
10:J:36:VAL:CG2	10:J:76:ILE:HG22	2.50	0.41
12:L:9:LYS:HG3	12:L:9:LYS:O	2.20	0.41
21:U:22:CYS:O	21:U:23:GLU:HG2	2.20	0.41
1:A:36:C:O2'	1:A:37:U:H5'	2.20	0.41
1:A:60:A:H1'	1:A:61:G:O4'	2.20	0.41
1:A:65:A:N7	1:A:381:C:C4	2.89	0.41
1:A:113:G:H2'	1:A:114:U:C6	2.55	0.41
1:A:399:G:H2'	1:A:400:C:H6	1.85	0.41
1:A:411:A:C4	1:A:413:G:H1'	2.55	0.41
1:A:714:G:H2'	1:A:715:A:C8	2.56	0.41
1:A:1067:A:H1'	1:A:1068:G:C8	2.55	0.41
1:A:1202:U:H2'	1:A:1203:C:H5'	2.02	0.41
1:A:1390:U:H2'	1:A:1391:U:H6	1.84	0.41
1:A:1517:G:O3'	22:V:124:VAL:HG13	2.19	0.41
1:A:1529:G:H3'	1:A:1529:G:OP2	2.20	0.41
2:B:81:ASP:O	2:B:85:SER:HB2	2.19	0.41
3:C:39:ARG:NH2	3:C:56:ILE:HD12	2.35	0.41
3:C:172:VAL:HG11	3:C:200:TRP:HB3	2.02	0.41
5:E:143:LEU:O	5:E:146:MET:HB2	2.20	0.41
7:G:19:SER:HG	7:G:22:LEU:HB2	1.86	0.41
7:G:46:LEU:O	7:G:46:LEU:HD13	2.20	0.41
9:I:20:ILE:HD12	9:I:20:ILE:N	2.35	0.41
9:I:64:ILE:HG22	9:I:65:THR:H	1.86	0.41
9:I:74:GLN:HE21	9:I:74:GLN:HB2	1.67	0.41
9:I:83:THR:HA	9:I:86:LEU:CD2	2.51	0.41
9:I:85:ALA:O	9:I:88:GLU:HB2	2.20	0.41
12:L:98:ARG:HA	12:L:98:ARG:NE	2.29	0.41
14:N:86:ALA:HB3	14:N:92:ILE:HD11	2.02	0.41
18:R:27:THR:O	18:R:28:LEU:HD12	2.20	0.41
19:S:35:ARG:HB3	19:S:71:GLY:HA2	2.02	0.41
1:A:502:A:O2'	1:A:503:C:H5'	2.21	0.41
1:A:821:G:H2'	1:A:822:U:C6	2.56	0.41
1:A:1248:A:H2	9:I:71:ILE:HD11	1.85	0.41
1:A:1320:C:P	19:S:69:LYS:HZ1	2.44	0.41
1:A:1437:A:H2'	1:A:1438:G:H8	1.86	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:B:175:ALA:C	2:B:177:ASN:N	2.79	0.41
3:C:122:GLN:O	3:C:127:VAL:HG22	2.20	0.41
9:I:98:ARG:HA	9:I:103:VAL:HG22	2.02	0.41
11:K:95:THR:HG23	11:K:96:ILE:HG13	2.02	0.41
14:N:5:MET:HE3	14:N:62:ARG:NH2	2.35	0.41
14:N:81:ILE:HD12	14:N:82:LYS:N	2.36	0.41
15:O:2:LEU:HD13	15:O:7:THR:HG23	2.02	0.41
16:P:45:GLU:C	16:P:47:GLU:H	2.29	0.41
22:V:206:LEU:HD12	22:V:206:LEU:HA	1.85	0.41
1:A:202:G:H1'	1:A:468:A:H8	1.85	0.41
1:A:337:G:O2'	1:A:338:A:H5'	2.21	0.41
1:A:557:G:H2'	1:A:558:G:O4'	2.20	0.41
1:A:709:U:H2'	1:A:710:G:C8	2.54	0.41
1:A:818:G:C2'	1:A:819:A:H5''	2.51	0.41
1:A:1037:C:H2'	1:A:1038:C:C6	2.55	0.41
1:A:1358:U:OP1	14:N:73:LEU:HA	2.20	0.41
1:A:1528:U:H6	1:A:1528:U:O5'	2.03	0.41
2:B:9:LEU:H	2:B:9:LEU:CD1	2.33	0.41
3:C:110:LEU:HB3	3:C:203:LYS:HZ3	1.86	0.41
3:C:155:ARG:NH1	3:C:192:TYR:HB2	2.36	0.41
4:D:63:ILE:C	4:D:65:GLY:H	2.29	0.41
6:F:44:ARG:HA	6:F:57:ALA:O	2.21	0.41
7:G:112:ASP:H	7:G:118:ARG:HD3	1.86	0.41
9:I:17:ARG:HE	9:I:65:THR:HB	1.85	0.41
10:J:30:LYS:HD2	10:J:30:LYS:O	2.21	0.41
13:M:16:ILE:CG2	13:M:17:ALA:N	2.84	0.41
13:M:58:GLU:HA	13:M:61:LYS:CE	2.49	0.41
16:P:52:LEU:HD22	16:P:52:LEU:N	2.36	0.41
17:Q:26:ARG:NH2	17:Q:28:VAL:HG11	2.36	0.41
1:A:28:A:H2'	1:A:29:U:O4'	2.21	0.41
1:A:114:U:O2'	1:A:115:G:H5'	2.19	0.41
1:A:152:A:N6	1:A:170:U:C2	2.88	0.41
1:A:154:U:H2'	1:A:155:A:C8	2.56	0.41
1:A:335:C:H4'	1:A:1434:A:H5'	2.03	0.41
1:A:499:A:H4'	1:A:500:G:H5'	2.03	0.41
1:A:619:U:O2	4:D:129:VAL:HG13	2.20	0.41
1:A:847:G:H2'	1:A:848:C:H6	1.86	0.41
1:A:1072:G:H2'	1:A:1073:U:H6	1.84	0.41
1:A:1126:U:O2'	1:A:1127:G:H5'	2.21	0.41
1:A:1238:A:N7	1:A:1303:C:H1'	2.35	0.41
1:A:1313:U:H5''	19:S:5:LYS:CG	2.33	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:1371:G:OP1	9:I:12:LYS:HG2	2.20	0.41
1:A:1391:U:H2'	1:A:1392:G:H8	1.80	0.41
4:D:158:LEU:HD23	4:D:158:LEU:N	2.36	0.41
5:E:95:MET:CE	5:E:114:LEU:HD11	2.51	0.41
6:F:74:LEU:HG	6:F:78:PHE:CD2	2.55	0.41
7:G:97:ALA:O	7:G:98:LEU:C	2.63	0.41
10:J:87:LEU:C	10:J:87:LEU:HD13	2.46	0.41
12:L:49:ARG:C	12:L:50:LYS:HE2	2.45	0.41
12:L:71:HIS:CG	12:L:72:ASN:N	2.86	0.41
14:N:17:ASP:HA	14:N:21:ALA:HB3	2.03	0.41
18:R:60:ARG:HA	18:R:63:TYR:CD1	2.54	0.41
19:S:64:GLU:O	19:S:66:VAL:HG23	2.20	0.41
20:T:53:MET:SD	20:T:57:VAL:HG21	2.60	0.41
22:V:68:GLU:CG	22:V:70:HIS:HE1	2.31	0.41
1:A:186:C:H2'	1:A:187:G:O4'	2.21	0.41
1:A:319:G:O2'	1:A:320:A:H5'	2.21	0.41
1:A:374:A:H5''	1:A:452:A:N1	2.36	0.41
1:A:415:A:N1	1:A:428:G:O6	2.53	0.41
1:A:484:G:H4'	1:A:485:U:H5'	2.02	0.41
1:A:634:C:O2'	1:A:635:A:H5'	2.20	0.41
1:A:980:C:H2'	1:A:981:U:H5'	2.03	0.41
1:A:993:G:C2'	1:A:995:C:H41	2.34	0.41
1:A:1181:G:P	1:A:1181:G:O4'	2.78	0.41
1:A:1217:C:H2'	1:A:1218:C:C6	2.56	0.41
1:A:1420:U:O4'	22:V:310:CYS:C	2.45	0.41
1:A:1457:G:H2'	1:A:1458:G:H8	1.86	0.41
2:B:116:LEU:HD13	2:B:140:LEU:HD23	2.02	0.41
4:D:36:ALA:O	4:D:38:GLY:N	2.54	0.41
7:G:141:HIS:O	7:G:144:ALA:O	2.39	0.41
9:I:9:GLY:O	9:I:16:ALA:HB3	2.20	0.41
9:I:59:LYS:HD2	9:I:60:LEU:HD23	2.03	0.41
13:M:103:THR:HG22	13:M:104:ASN:N	2.33	0.41
17:Q:3:LYS:HD2	17:Q:3:LYS:O	2.21	0.41
22:V:53:ILE:HD11	22:V:63:GLU:HB2	2.03	0.41
22:V:167:LEU:C	22:V:167:LEU:HD13	2.46	0.41
1:A:425:G:O2'	1:A:426:U:H5'	2.21	0.41
1:A:612:C:H2'	1:A:613:C:C6	2.55	0.41
1:A:622:A:H2'	1:A:623:C:H5'	2.02	0.41
1:A:959:A:H2'	1:A:960:U:O4'	2.21	0.41
1:A:1098:C:H2'	1:A:1099:G:O4'	2.20	0.41
1:A:1311:A:H2'	1:A:1312:G:O4'	2.21	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:B:45:THR:HA	2:B:48:MET:CG	2.50	0.41
2:B:131:LYS:O	2:B:134:LEU:HB2	2.21	0.41
8:H:75:GLN:O	8:H:126:CYS:HB2	2.21	0.41
9:I:52:GLU:N	9:I:56:MET:HG2	2.35	0.41
9:I:62:LEU:HD22	9:I:62:LEU:N	2.35	0.41
10:J:6:ILE:HA	10:J:102:LEU:CD2	2.51	0.41
11:K:22:ILE:HA	11:K:31:VAL:HG22	2.03	0.41
12:L:113:ARG:HD3	12:L:121:PRO:HD3	2.02	0.41
14:N:11:LYS:C	14:N:13:VAL:N	2.77	0.41
14:N:65:GLN:HA	14:N:65:GLN:HE21	1.86	0.41
21:U:33:ARG:CZ	21:U:34:ARG:HG3	2.51	0.41
22:V:115:LEU:HD11	22:V:281:PHE:CD1	2.56	0.41
1:A:56:U:O5'	1:A:56:U:H6	2.04	0.41
1:A:130:A:O4'	17:Q:64:ARG:HD2	2.21	0.41
1:A:179:A:H2'	1:A:180:U:H6	1.83	0.41
1:A:358:U:H2'	1:A:359:G:C8	2.55	0.41
1:A:367:U:OP1	1:A:395:C:H1'	2.21	0.41
1:A:537:G:H5''	12:L:109:ARG:NH1	2.36	0.41
1:A:542:G:O2'	1:A:543:U:H5'	2.20	0.41
1:A:635:A:H2'	1:A:636:U:C6	2.56	0.41
1:A:738:C:H2'	1:A:739:C:C6	2.56	0.41
1:A:769:G:O2'	1:A:770:C:H5'	2.21	0.41
1:A:812:G:O2'	1:A:813:U:H6	2.04	0.41
1:A:861:G:O2'	1:A:862:C:H5'	2.20	0.41
1:A:896:C:O2'	1:A:897:C:H5'	2.20	0.41
1:A:1047:G:O2'	1:A:1048:G:H5'	2.21	0.41
1:A:1082:A:H2'	1:A:1083:U:C6	2.56	0.41
1:A:1195:C:H2'	1:A:1197:A:O4'	2.21	0.41
1:A:1220:G:H21	19:S:53:GLY:HA2	1.84	0.41
1:A:1432:G:H1'	1:A:1468:A:N6	2.36	0.41
1:A:1468:A:O2'	1:A:1469:C:H5'	2.21	0.41
2:B:42:LEU:HA	2:B:45:THR:OG1	2.21	0.41
2:B:204:ASP:O	2:B:205:ALA:HB3	2.21	0.41
2:B:205:ALA:O	2:B:209:VAL:HG22	2.20	0.41
2:B:209:VAL:O	2:B:213:LEU:HB2	2.21	0.41
3:C:148:ILE:HA	3:C:200:TRP:O	2.20	0.41
4:D:26:ALA:O	4:D:27:ILE:O	2.39	0.41
4:D:43:ARG:HB3	4:D:44:LYS:H	1.49	0.41
4:D:87:GLU:O	4:D:88:ASN:C	2.62	0.41
5:E:108:GLY:H	5:E:110:MET:CE	2.28	0.41
9:I:107:ALA:O	9:I:109:GLN:HG2	2.21	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
9:I:118:ARG:HG2	9:I:118:ARG:O	2.21	0.41
10:J:5:ARG:HG3	10:J:79:PRO:HD3	2.01	0.41
11:K:81:LEU:CD2	11:K:104:PHE:HB3	2.50	0.41
11:K:84:MET:HE2	11:K:110:THR:OG1	2.21	0.41
11:K:85:VAL:O	11:K:111:ASP:HA	2.20	0.41
12:L:79:ILE:C	12:L:79:ILE:HD12	2.46	0.41
12:L:83:GLY:HA2	12:L:94:TYR:CD1	2.52	0.41
14:N:51:PRO:CB	14:N:54:SER:HB3	2.48	0.41
14:N:73:LEU:O	14:N:74:ARG:C	2.64	0.41
15:O:52:ARG:HA	15:O:55:LEU:HB3	2.03	0.41
16:P:20:VAL:HG22	16:P:21:VAL:H	1.83	0.41
16:P:67:ILE:CD1	16:P:72:ALA:HA	2.49	0.41
16:P:73:ALA:HB1	16:P:77:GLU:OE2	2.21	0.41
19:S:28:LYS:H	19:S:28:LYS:HD3	1.84	0.41
19:S:57:VAL:HG21	19:S:74:ALA:CB	2.51	0.41
20:T:57:VAL:O	20:T:57:VAL:HG12	2.21	0.41
20:T:58:ASP:HA	20:T:61:ALA:HB3	2.03	0.41
22:V:308:ARG:HA	22:V:308:ARG:HD3	1.83	0.41
1:A:9:G:OP2	5:E:125:LYS:HD3	2.20	0.41
1:A:35:G:H2'	1:A:36:C:H6	1.86	0.41
1:A:70:U:H1'	1:A:71:A:N7	2.36	0.41
1:A:291:U:O2'	1:A:292:G:H5'	2.21	0.41
1:A:314:C:O2'	1:A:315:A:H5'	2.21	0.41
1:A:495:A:H4'	1:A:496:A:O5'	2.21	0.41
1:A:628:G:O2'	1:A:629:A:H5'	2.21	0.41
1:A:737:C:H2'	1:A:738:C:H6	1.86	0.41
1:A:829:G:O2'	1:A:830:G:H5'	2.21	0.41
1:A:841:C:H3'	1:A:843:U:P	2.60	0.41
1:A:1073:U:O2	2:B:102:ASN:ND2	2.53	0.41
1:A:1150:A:H1'	1:A:1280:A:N6	2.36	0.41
1:A:1231:G:H2'	1:A:1232:U:C6	2.55	0.41
1:A:1269:A:H1'	1:A:1326:U:O4'	2.20	0.41
1:A:1521:C:H2'	1:A:1522:U:C6	2.56	0.41
2:B:186:VAL:HG22	2:B:198:VAL:HG23	2.03	0.41
3:C:173:PRO:C	3:C:175:HIS:H	2.30	0.41
5:E:131:ASN:ND2	5:E:134:ASN:H	2.19	0.41
6:F:29:ILE:HD13	6:F:64:VAL:CG1	2.51	0.41
7:G:57:GLU:OE1	7:G:57:GLU:HA	2.21	0.41
7:G:101:ARG:HB3	7:G:105:GLU:OE2	2.20	0.41
8:H:66:GLN:HB3	8:H:67:GLY:H	1.67	0.41
9:I:4:GLN:HE21	9:I:21:LYS:HE3	1.84	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
9:I:126:PHE:O	9:I:128:LYS:N	2.54	0.41
10:J:73:LEU:O	10:J:74:VAL:HB	2.21	0.41
17:Q:57:VAL:N	17:Q:79:GLU:HB3	2.35	0.41
22:V:140:ILE:HD13	22:V:140:ILE:N	2.35	0.41
1:A:5:U:H1'	1:A:6:G:N1	2.35	0.40
1:A:8:A:H1'	5:E:106:ALA:C	2.46	0.40
1:A:17:U:H1'	1:A:1080:A:N3	2.36	0.40
1:A:140:U:H2'	1:A:141:G:C8	2.56	0.40
1:A:178:C:O2'	1:A:179:A:H5'	2.20	0.40
1:A:328:C:H4'	1:A:329:A:H5''	2.02	0.40
1:A:374:A:H2'	1:A:375:U:C6	2.56	0.40
1:A:386:C:C2'	1:A:387:U:H5'	2.51	0.40
1:A:547:A:H4'	1:A:548:G:O5'	2.21	0.40
1:A:633:G:H2'	1:A:634:C:H6	1.86	0.40
1:A:635:A:O2'	1:A:636:U:H5'	2.21	0.40
1:A:637:C:H2'	1:A:638:U:C6	2.56	0.40
1:A:842:U:OP1	1:A:842:U:H3'	2.21	0.40
1:A:1340:A:H2'	1:A:1341:U:H6	1.86	0.40
1:A:1347:G:N2	1:A:1373:G:H2'	2.36	0.40
1:A:1401:G:H2'	1:A:1402:C:O4'	2.21	0.40
2:B:107:ARG:HA	2:B:110:ILE:HD12	2.04	0.40
4:D:120:LYS:CB	4:D:145:ARG:HH21	2.32	0.40
5:E:55:VAL:N	5:E:56:PRO:CD	2.84	0.40
6:F:3:HIS:HB2	6:F:92:THR:OG1	2.21	0.40
14:N:19:TYR:HD1	14:N:23:ARG:HD3	1.86	0.40
15:O:70:LYS:NZ	15:O:74:VAL:HA	2.36	0.40
17:Q:10:ARG:CZ	17:Q:56:ASP:N	2.84	0.40
17:Q:76:ARG:HE	17:Q:78:VAL:HG22	1.85	0.40
20:T:27:MET:O	20:T:31:ILE:HG13	2.21	0.40
20:T:57:VAL:C	20:T:60:GLN:NE2	2.79	0.40
22:V:82:VAL:CG2	22:V:129:ALA:HB3	2.51	0.40
1:A:640:A:O2'	1:A:641:U:H5'	2.22	0.40
1:A:812:G:OP1	1:A:812:G:C4'	2.68	0.40
1:A:1121:U:O2'	1:A:1122:U:H5'	2.20	0.40
1:A:1408:A:OP1	22:V:283:LEU:CA	2.69	0.40
2:B:56:LEU:HA	2:B:59:ILE:CG1	2.52	0.40
6:F:29:ILE:HG23	6:F:66:ALA:HB2	2.03	0.40
6:F:46:GLN:HE21	6:F:46:GLN:HB2	1.50	0.40
9:I:126:PHE:O	9:I:127:SER:C	2.64	0.40
15:O:52:ARG:O	15:O:55:LEU:HB3	2.21	0.40
16:P:39:PHE:HE2	16:P:70:ARG:NH2	2.18	0.40

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
18:R:23:LYS:C	18:R:25:ILE:N	2.79	0.40
19:S:62:THR:N	19:S:65:MET:HE3	2.35	0.40
1:A:16:A:H4'	5:E:21:SER:N	2.35	0.40
1:A:192:A:O2'	1:A:193:C:H5'	2.20	0.40
1:A:645:G:O2'	1:A:646:G:H5'	2.22	0.40
1:A:730:G:O2'	1:A:766:A:H5'	2.20	0.40
1:A:768:A:H4'	1:A:1523:G:N2	2.35	0.40
1:A:792:A:O2'	1:A:794:A:N7	2.50	0.40
1:A:880:C:H2'	1:A:881:G:H8	1.86	0.40
1:A:945:G:N1	1:A:1337:G:C2	2.89	0.40
1:A:1036:A:H2'	1:A:1036:A:N3	2.36	0.40
1:A:1304:G:H2'	1:A:1305:G:C1'	2.51	0.40
1:A:1391:U:O2'	1:A:1392:G:H5'	2.21	0.40
1:A:1453:G:N2	1:A:1454:G:C8	2.90	0.40
1:A:1487:G:O2'	1:A:1488:G:H5'	2.21	0.40
1:A:1499:A:C1'	1:A:1520:C:H5'	2.51	0.40
3:C:6:PRO:HG2	3:C:183:TYR:CD2	2.57	0.40
3:C:181:ILE:HG22	3:C:182:ASP:N	2.35	0.40
5:E:35:LEU:CD1	5:E:133:ILE:HA	2.50	0.40
5:E:51:LYS:O	5:E:52:ALA:HB2	2.21	0.40
5:E:131:ASN:ND2	5:E:134:ASN:N	2.69	0.40
10:J:37:ARG:N	10:J:77:VAL:HG22	2.37	0.40
12:L:80:LEU:HD13	12:L:101:LEU:CD1	2.51	0.40
14:N:30:ILE:C	14:N:32:ASP:N	2.79	0.40
17:Q:10:ARG:NE	17:Q:56:ASP:O	2.55	0.40
17:Q:30:HIS:N	17:Q:35:LYS:H	2.19	0.40
18:R:28:LEU:C	18:R:30:ASN:N	2.79	0.40
1:A:208:U:H2'	1:A:209:U:H5''	2.04	0.40
1:A:492:C:C2'	1:A:493:A:H5''	2.51	0.40
1:A:648:A:H2'	1:A:649:A:C8	2.56	0.40
1:A:659:U:O2'	1:A:660:C:H5'	2.21	0.40
1:A:662:U:H2'	1:A:663:A:H8	1.81	0.40
1:A:711:G:H2'	1:A:712:A:H8	1.85	0.40
1:A:1057:G:O2'	1:A:1058:G:H5'	2.21	0.40
1:A:1271:A:H2'	1:A:1272:G:C8	2.55	0.40
1:A:1288:A:N1	1:A:1371:G:H1'	2.36	0.40
1:A:1368:A:O2'	1:A:1369:C:H5'	2.22	0.40
2:B:120:SER:HA	2:B:125:PHE:CB	2.44	0.40
2:B:150:ILE:HG13	2:B:153:MET:CE	2.39	0.40
3:C:166:TRP:HB3	3:C:167:TYR:H	1.49	0.40
4:D:58:GLN:O	4:D:62:ARG:HG2	2.22	0.40

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
4:D:119:HIS:C	4:D:121:ALA:H	2.29	0.40
4:D:138:PRO:C	4:D:140:ASP:H	2.29	0.40
7:G:13:PRO:HB2	7:G:18:GLY:C	2.46	0.40
7:G:114:SER:O	7:G:118:ARG:HG3	2.21	0.40
9:I:24:ASN:O	9:I:61:ASP:HB2	2.21	0.40
12:L:30:ARG:HB3	12:L:57:THR:CG2	2.50	0.40
12:L:48:LEU:HD23	12:L:48:LEU:N	2.24	0.40
15:O:70:LYS:HG2	15:O:77:TYR:CG	2.56	0.40
16:P:6:LEU:CD1	16:P:71:VAL:HB	2.52	0.40
16:P:6:LEU:HD21	16:P:74:LEU:HD11	2.02	0.40
17:Q:68:LYS:O	17:Q:69:THR:HB	2.21	0.40
19:S:35:ARG:C	19:S:37:SER:H	2.29	0.40
19:S:48:ILE:HG21	19:S:70:LEU:HD21	2.04	0.40
20:T:60:GLN:C	20:T:62:ALA:H	2.28	0.40
21:U:24:LYS:O	21:U:26:GLY:N	2.54	0.40
1:A:167:A:H2'	1:A:168:G:H8	1.86	0.40
1:A:223:A:H2'	1:A:224:U:C6	2.57	0.40
1:A:438:U:H2'	1:A:494:G:O6	2.21	0.40
1:A:529:G:H22	12:L:47:ALA:HB2	1.85	0.40
1:A:1074:G:O2'	1:A:1075:U:H5'	2.21	0.40
1:A:1283:U:H2'	1:A:1284:C:H6	1.86	0.40
1:A:1517:G:O3'	22:V:124:VAL:CG1	2.70	0.40
2:B:162:VAL:CG1	2:B:184:ALA:HB2	2.46	0.40
4:D:48:SER:OG	4:D:49:ASP:N	2.52	0.40
6:F:42:TRP:N	6:F:42:TRP:CD1	2.89	0.40
7:G:94:ARG:CZ	7:G:98:LEU:HD21	2.51	0.40
7:G:109:LYS:N	7:G:109:LYS:HD2	2.36	0.40
8:H:110:MET:HB3	8:H:114:ALA:HB3	2.03	0.40
9:I:38:PHE:HD1	9:I:38:PHE:HA	1.77	0.40
9:I:93:LEU:C	9:I:93:LEU:HD12	2.47	0.40
12:L:87:LYS:NZ	12:L:87:LYS:HA	2.36	0.40
12:L:107:LYS:HD2	12:L:107:LYS:N	2.37	0.40
18:R:31:TYR:C	18:R:39:VAL:HG13	2.47	0.40
19:S:31:ARG:HG2	19:S:31:ARG:HH11	1.86	0.40
21:U:40:PRO:C	21:U:42:THR:H	2.29	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	
2	B	216/218 (99%)	145 (67%)	53 (24%)	18 (8%)	0	9
3	C	204/206 (99%)	135 (66%)	49 (24%)	20 (10%)	0	7
4	D	203/205 (99%)	131 (64%)	54 (27%)	18 (9%)	0	8
5	E	148/150 (99%)	107 (72%)	34 (23%)	7 (5%)	2	16
6	F	98/100 (98%)	69 (70%)	23 (24%)	6 (6%)	1	13
7	G	148/150 (99%)	103 (70%)	37 (25%)	8 (5%)	1	15
8	H	127/129 (98%)	105 (83%)	19 (15%)	3 (2%)	4	27
9	I	125/127 (98%)	86 (69%)	28 (22%)	11 (9%)	0	8
10	J	96/98 (98%)	63 (66%)	21 (22%)	12 (12%)	0	4
11	K	115/117 (98%)	75 (65%)	27 (24%)	13 (11%)	0	5
12	L	121/123 (98%)	74 (61%)	30 (25%)	17 (14%)	0	3
13	M	112/114 (98%)	87 (78%)	14 (12%)	11 (10%)	0	7
14	N	92/100 (92%)	59 (64%)	24 (26%)	9 (10%)	0	7
15	O	86/88 (98%)	66 (77%)	18 (21%)	2 (2%)	5	28
16	P	80/82 (98%)	59 (74%)	17 (21%)	4 (5%)	1	16
17	Q	78/80 (98%)	48 (62%)	26 (33%)	4 (5%)	1	15
18	R	53/55 (96%)	27 (51%)	19 (36%)	7 (13%)	0	4
19	S	77/79 (98%)	57 (74%)	14 (18%)	6 (8%)	1	10
20	T	83/85 (98%)	69 (83%)	8 (10%)	6 (7%)	1	11
21	U	49/51 (96%)	22 (45%)	12 (24%)	15 (31%)	0	0
22	V	265/277 (96%)	256 (97%)	7 (3%)	2 (1%)	16	54
All	All	2576/2634 (98%)	1843 (72%)	534 (21%)	199 (8%)	1	10

All (199) Ramachandran outliers are listed below:

Mol	Chain	Res	Type
2	B	19	THR
2	B	22	TRP
2	B	163	ILE
2	B	186	VAL
3	C	2	GLN
3	C	91	ALA
3	C	153	SER
4	D	18	LEU
4	D	31	CYS
4	D	191	SER
6	F	92	THR
9	I	8	THR
9	I	43	ALA
10	J	57	VAL
11	K	56	LYS
11	K	126	ARG
12	L	10	PRO
12	L	23	LEU
13	M	6	ILE
13	M	14	ALA
14	N	52	ARG
21	U	14	ALA
21	U	22	CYS
2	B	18	GLN
3	C	19	SER
3	C	25	THR
3	C	54	ILE
3	C	81	GLU
3	C	100	ILE
3	C	180	ASP
4	D	25	ARG
4	D	107	GLY
4	D	152	SER
4	D	172	VAL
4	D	192	ALA
5	E	20	VAL
5	E	128	GLY
6	F	62	MET
6	F	64	VAL
6	F	65	GLU
7	G	84	TYR
7	G	129	ASN
9	I	57	VAL

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Mol	Chain	Res	Type
9	I	71	ILE
9	I	108	ARG
10	J	75	ASP
10	J	92	LEU
11	K	50	GLY
11	K	53	GLY
11	K	125	LYS
12	L	19	ASN
12	L	72	ASN
12	L	84	GLY
13	M	3	ILE
13	M	66	GLY
13	M	104	ASN
14	N	51	PRO
14	N	61	ASN
15	O	73	ASP
16	P	79	ASN
17	Q	6	THR
17	Q	34	GLY
18	R	32	ILE
19	S	5	LYS
20	T	85	LEU
21	U	32	ARG
21	U	34	ARG
2	B	20	ARG
2	B	76	SER
3	C	59	PRO
3	C	78	LYS
3	C	136	ALA
4	D	27	ILE
4	D	28	ASP
4	D	43	ARG
4	D	151	GLN
4	D	177	MET
5	E	43	GLY
6	F	94	HIS
7	G	89	GLU
9	I	44	ARG
9	I	55	ASP
9	I	127	SER
10	J	62	ARG
10	J	93	ALA

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Mol	Chain	Res	Type
11	K	70	ALA
11	K	77	GLY
11	K	91	GLY
12	L	13	ARG
12	L	60	PHE
12	L	122	LYS
13	M	7	ASN
13	M	22	TYR
13	M	65	GLU
13	M	98	GLY
14	N	21	ALA
14	N	74	ARG
14	N	80	ARG
17	Q	5	ARG
18	R	21	ASP
18	R	71	ASP
20	T	59	ARG
21	U	6	ARG
21	U	24	LYS
21	U	25	ALA
21	U	40	PRO
2	B	11	ALA
2	B	41	ASN
2	B	88	GLN
2	B	200	PRO
3	C	65	VAL
3	C	186	SER
4	D	29	THR
5	E	25	LYS
5	E	34	ALA
6	F	54	LEU
7	G	18	GLY
7	G	57	GLU
7	G	75	LYS
7	G	86	VAL
8	H	46	GLU
9	I	67	LYS
10	J	56	HIS
10	J	61	ALA
10	J	74	VAL
11	K	57	SER
11	K	108	ASN

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Mol	Chain	Res	Type
12	L	14	LYS
12	L	15	VAL
12	L	47	ALA
12	L	56	LEU
12	L	67	GLY
12	L	120	ARG
14	N	31	SER
14	N	94	GLY
16	P	42	ILE
17	Q	69	THR
18	R	22	TYR
18	R	47	ARG
19	S	27	LYS
19	S	36	ARG
19	S	65	MET
21	U	26	GLY
21	U	33	ARG
21	U	36	PHE
21	U	37	TYR
21	U	41	THR
22	V	281	PHE
2	B	24	PRO
2	B	27	LYS
2	B	87	ASP
3	C	14	VAL
3	C	145	ALA
3	C	167	TYR
5	E	146	MET
9	I	122	ARG
10	J	36	VAL
11	K	127	ARG
12	L	42	LYS
12	L	121	PRO
13	M	105	ALA
15	O	17	ASP
18	R	20	ILE
19	S	8	PRO
20	T	5	SER
20	T	46	ALA
2	B	94	ARG
2	B	205	ALA
3	C	26	LYS

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Mol	Chain	Res	Type
3	C	112	ALA
4	D	6	PRO
5	E	107	GLY
7	G	6	ILE
10	J	95	GLY
11	K	106	ILE
14	N	33	VAL
19	S	39	ILE
21	U	13	VAL
22	V	275	SER
2	B	157	PRO
3	C	90	VAL
4	D	63	ILE
4	D	175	GLY
11	K	119	GLY
16	P	41	PRO
10	J	41	PRO
16	P	49	GLY
20	T	3	ILE
4	D	37	PRO
8	H	71	VAL
10	J	96	VAL
12	L	62	VAL
13	M	23	GLY
20	T	41	GLY
21	U	52	VAL
2	B	150	ILE
8	H	77	VAL
9	I	110	VAL
18	R	43	ILE

5.3.2 Protein sidechains ⓘ

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

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

Mol	Chain	Analysed	Rotameric	Outliers	Percentiles
2	B	180/180 (100%)	143 (79%)	37 (21%)	1 6

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Mol	Chain	Analysed	Rotameric	Outliers	Percentiles	
3	C	170/170 (100%)	145 (85%)	25 (15%)	3	13
4	D	172/172 (100%)	137 (80%)	35 (20%)	1	7
5	E	113/113 (100%)	93 (82%)	20 (18%)	2	9
6	F	87/87 (100%)	68 (78%)	19 (22%)	1	6
7	G	123/123 (100%)	101 (82%)	22 (18%)	2	9
8	H	104/104 (100%)	88 (85%)	16 (15%)	2	12
9	I	105/105 (100%)	83 (79%)	22 (21%)	1	6
10	J	86/86 (100%)	70 (81%)	16 (19%)	1	8
11	K	90/90 (100%)	74 (82%)	16 (18%)	2	9
12	L	103/103 (100%)	89 (86%)	14 (14%)	3	14
13	M	92/92 (100%)	72 (78%)	20 (22%)	1	6
14	N	79/83 (95%)	66 (84%)	13 (16%)	2	10
15	O	76/76 (100%)	70 (92%)	6 (8%)	11	32
16	P	65/65 (100%)	54 (83%)	11 (17%)	2	10
17	Q	74/74 (100%)	59 (80%)	15 (20%)	1	7
18	R	48/48 (100%)	42 (88%)	6 (12%)	4	16
19	S	70/70 (100%)	60 (86%)	10 (14%)	3	13
20	T	65/65 (100%)	57 (88%)	8 (12%)	4	17
21	U	44/44 (100%)	36 (82%)	8 (18%)	2	9
22	V	235/235 (100%)	227 (97%)	8 (3%)	32	54
All	All	2181/2185 (100%)	1834 (84%)	347 (16%)	4	11

All (347) residues with a non-rotameric sidechain are listed below:

Mol	Chain	Res	Type
2	B	9	LEU
2	B	18	GLN
2	B	22	TRP
2	B	26	MET
2	B	36	LYS
2	B	38	HIS
2	B	40	ILE
2	B	46	VAL
2	B	48	MET
2	B	56	LEU

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Mol	Chain	Res	Type
2	B	57	ASN
2	B	58	LYS
2	B	67	LEU
2	B	69	VAL
2	B	73	ARG
2	B	81	ASP
2	B	88	GLN
2	B	90	PHE
2	B	94	ARG
2	B	95	TRP
2	B	101	THR
2	B	116	LEU
2	B	121	GLN
2	B	122	ASP
2	B	125	PHE
2	B	127	LYS
2	B	128	LEU
2	B	145	ASN
2	B	158	ASP
2	B	162	VAL
2	B	185	ILE
2	B	188	THR
2	B	196	ASP
2	B	202	ASN
2	B	206	ILE
2	B	209	VAL
2	B	222	GLU
3	C	14	VAL
3	C	20	THR
3	C	26	LYS
3	C	31	ASN
3	C	40	GLN
3	C	42	LEU
3	C	69	THR
3	C	100	ILE
3	C	102	ILE
3	C	113	LYS
3	C	141	MET
3	C	143	LEU
3	C	146	LYS
3	C	156	LEU
3	C	164	THR

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Mol	Chain	Res	Type
3	C	165	GLU
3	C	166	TRP
3	C	174	LEU
3	C	176	THR
3	C	180	ASP
3	C	185	THR
3	C	189	HIS
3	C	190	THR
3	C	192	TYR
3	C	205	GLU
4	D	7	LYS
4	D	8	LEU
4	D	10	LEU
4	D	20	LEU
4	D	21	LYS
4	D	25	ARG
4	D	35	GLN
4	D	39	GLN
4	D	54	LEU
4	D	56	GLU
4	D	58	GLN
4	D	64	TYR
4	D	69	ARG
4	D	77	GLU
4	D	80	ARG
4	D	85	THR
4	D	87	GLU
4	D	94	GLU
4	D	131	ILE
4	D	133	SER
4	D	137	SER
4	D	147	LYS
4	D	150	LYS
4	D	153	ARG
4	D	154	VAL
4	D	155	LYS
4	D	158	LEU
4	D	160	LEU
4	D	172	VAL
4	D	176	LYS
4	D	177	MET
4	D	183	ARG

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Mol	Chain	Res	Type
4	D	186	GLU
4	D	189	ASP
4	D	194	ILE
5	E	11	GLN
5	E	23	THR
5	E	25	LYS
5	E	32	PHE
5	E	45	VAL
5	E	55	VAL
5	E	63	MET
5	E	68	ARG
5	E	81	GLN
5	E	92	ARG
5	E	102	THR
5	E	110	MET
5	E	119	VAL
5	E	122	VAL
5	E	123	LEU
5	E	125	LYS
5	E	139	THR
5	E	143	LEU
5	E	151	MET
5	E	158	LYS
6	F	37	HIS
6	F	38	ARG
6	F	39	LEU
6	F	46	GLN
6	F	53	LYS
6	F	54	LEU
6	F	55	HIS
6	F	61	LEU
6	F	65	GLU
6	F	69	GLU
6	F	71	ILE
6	F	73	GLU
6	F	75	GLU
6	F	76	THR
6	F	78	PHE
6	F	84	VAL
6	F	87	SER
6	F	88	MET
6	F	92	THR

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Mol	Chain	Res	Type
7	G	3	ARG
7	G	10	LYS
7	G	11	ILE
7	G	14	ASP
7	G	16	LYS
7	G	19	SER
7	G	21	LEU
7	G	22	LEU
7	G	26	VAL
7	G	29	LEU
7	G	42	VAL
7	G	46	LEU
7	G	52	ARG
7	G	68	VAL
7	G	72	VAL
7	G	74	VAL
7	G	84	TYR
7	G	89	GLU
7	G	94	ARG
7	G	112	ASP
7	G	114	SER
7	G	119	LEU
8	H	4	ASP
8	H	17	GLN
8	H	25	THR
8	H	30	LYS
8	H	37	ASN
8	H	45	ILE
8	H	55	LYS
8	H	61	THR
8	H	74	ILE
8	H	76	ARG
8	H	82	LEU
8	H	83	ARG
8	H	93	LYS
8	H	111	THR
8	H	113	ARG
8	H	120	LEU
9	I	3	ASN
9	I	26	LYS
9	I	30	ASN
9	I	35	GLU

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Mol	Chain	Res	Type
9	I	36	GLN
9	I	38	PHE
9	I	41	GLU
9	I	44	ARG
9	I	53	LEU
9	I	56	MET
9	I	58	GLU
9	I	59	LYS
9	I	60	LEU
9	I	61	ASP
9	I	84	ARG
9	I	86	LEU
9	I	87	MET
9	I	106	ASP
9	I	115	VAL
9	I	119	LYS
9	I	121	ARG
9	I	126	PHE
10	J	11	LYS
10	J	14	ASP
10	J	18	ILE
10	J	40	ILE
10	J	48	ARG
10	J	50	THR
10	J	57	VAL
10	J	71	LEU
10	J	73	LEU
10	J	75	ASP
10	J	81	GLU
10	J	88	MET
10	J	89	ARG
10	J	92	LEU
10	J	98	VAL
10	J	102	LEU
11	K	33	ILE
11	K	36	ARG
11	K	52	ARG
11	K	56	LYS
11	K	58	THR
11	K	71	ASP
11	K	73	VAL
11	K	74	LYS

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Mol	Chain	Res	Type
11	K	83	VAL
11	K	84	MET
11	K	85	VAL
11	K	92	ARG
11	K	99	LEU
11	K	107	THR
11	K	108	ASN
11	K	115	ILE
12	L	18	SER
12	L	20	VAL
12	L	28	GLN
12	L	30	ARG
12	L	33	CYS
12	L	35	ARG
12	L	38	THR
12	L	50	LYS
12	L	58	ASN
12	L	71	HIS
12	L	87	LYS
12	L	103	CYS
12	L	107	LYS
12	L	118	VAL
13	M	2	ARG
13	M	11	HIS
13	M	18	LEU
13	M	27	THR
13	M	41	ASP
13	M	43	LYS
13	M	44	ILE
13	M	57	ASP
13	M	62	PHE
13	M	64	VAL
13	M	67	ASP
13	M	68	LEU
13	M	81	ASP
13	M	90	HIS
13	M	91	ARG
13	M	97	ARG
13	M	99	GLN
13	M	101	THR
13	M	102	LYS
13	M	106	ARG

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Mol	Chain	Res	Type
14	N	11	LYS
14	N	15	LEU
14	N	27	LYS
14	N	41	TRP
14	N	45	LEU
14	N	47	LEU
14	N	50	LEU
14	N	53	ASP
14	N	59	GLN
14	N	65	GLN
14	N	73	LEU
14	N	74	ARG
14	N	81	ILE
15	O	7	THR
15	O	17	ASP
15	O	25	GLU
15	O	58	MET
15	O	64	LYS
15	O	70	LYS
16	P	4	ILE
16	P	5	ARG
16	P	19	VAL
16	P	26	ASN
16	P	29	ASN
16	P	31	ARG
16	P	34	GLU
16	P	46	LYS
16	P	48	GLU
16	P	55	ASP
16	P	63	GLN
17	Q	4	ILE
17	Q	10	ARG
17	Q	15	LYS
17	Q	20	ILE
17	Q	24	ILE
17	Q	25	GLU
17	Q	39	ARG
17	Q	45	VAL
17	Q	52	CYS
17	Q	60	ILE
17	Q	67	SER
17	Q	74	LEU

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Mol	Chain	Res	Type
17	Q	76	ARG
17	Q	78	VAL
17	Q	80	LYS
18	R	20	ILE
18	R	30	ASN
18	R	43	ILE
18	R	52	ARG
18	R	54	LEU
18	R	65	SER
19	S	5	LYS
19	S	12	LEU
19	S	14	LEU
19	S	20	LYS
19	S	28	LYS
19	S	36	ARG
19	S	40	PHE
19	S	47	THR
19	S	72	GLU
19	S	73	PHE
20	T	14	GLU
20	T	26	MET
20	T	29	THR
20	T	34	VAL
20	T	59	ARG
20	T	67	HIS
20	T	70	LYS
20	T	85	LEU
21	U	11	PHE
21	U	12	ASP
21	U	13	VAL
21	U	16	ARG
21	U	17	ARG
21	U	28	LEU
21	U	33	ARG
21	U	44	ARG
22	V	134	ILE
22	V	140	ILE
22	V	146	LEU
22	V	172	LEU
22	V	194	ARG
22	V	206	LEU
22	V	263	LEU

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Mol	Chain	Res	Type
22	V	341	LEU

Sometimes sidechains can be flipped to improve hydrogen bonding and reduce clashes. All (80) such sidechains are listed below:

Mol	Chain	Res	Type
2	B	17	HIS
2	B	23	ASN
2	B	35	ASN
2	B	108	GLN
2	B	119	GLN
2	B	121	GLN
2	B	145	ASN
2	B	167	HIS
2	B	202	ASN
3	C	2	GLN
3	C	5	HIS
3	C	40	GLN
3	C	68	HIS
3	C	122	GLN
3	C	139	ASN
3	C	184	ASN
4	D	35	GLN
4	D	84	ASN
4	D	99	ASN
4	D	115	GLN
4	D	135	GLN
4	D	151	GLN
4	D	195	ASN
5	E	69	ASN
5	E	72	ASN
5	E	120	HIS
5	E	121	ASN
5	E	131	ASN
5	E	134	ASN
6	F	14	GLN
6	F	17	GLN
6	F	46	GLN
6	F	58	HIS
7	G	67	ASN
7	G	85	GLN
7	G	121	ASN
8	H	3	GLN

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Mol	Chain	Res	Type
8	H	17	GLN
9	I	4	GLN
9	I	24	ASN
9	I	30	ASN
9	I	31	GLN
9	I	36	GLN
9	I	49	GLN
9	I	109	GLN
10	J	15	HIS
10	J	20	GLN
10	J	64	GLN
10	J	99	GLN
11	K	28	ASN
11	K	37	GLN
11	K	100	ASN
11	K	118	ASN
12	L	58	ASN
13	M	51	GLN
13	M	99	GLN
14	N	59	GLN
14	N	61	ASN
14	N	65	GLN
15	O	27	GLN
15	O	34	GLN
15	O	39	GLN
16	P	9	HIS
16	P	18	GLN
16	P	29	ASN
18	R	53	GLN
19	S	42	ASN
19	S	51	HIS
19	S	55	GLN
19	S	56	HIS
20	T	20	ASN
20	T	54	GLN
20	T	60	GLN
22	V	70	HIS
22	V	73	ASN
22	V	190	ASN
22	V	203	GLN
22	V	285	HIS
22	V	293	GLN

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Mol	Chain	Res	Type
22	V	312	HIS

5.3.3 RNA ⓘ

Mol	Chain	Analysed	Backbone Outliers	Pucker Outliers
1	A	1529/1530 (99%)	279 (18%)	25 (1%)

All (279) RNA backbone outliers are listed below:

Mol	Chain	Res	Type
1	A	7	A
1	A	9	G
1	A	14	U
1	A	31	G
1	A	32	A
1	A	39	G
1	A	47	C
1	A	48	C
1	A	51	A
1	A	52	C
1	A	55	A
1	A	61	G
1	A	66	A
1	A	72	A
1	A	76	G
1	A	78	A
1	A	79	G
1	A	83	C
1	A	85	U
1	A	86	G
1	A	87	C
1	A	88	U
1	A	89	U
1	A	95	C
1	A	108	G
1	A	121	U
1	A	122	G
1	A	131	A
1	A	151	A
1	A	164	G
1	A	182	A

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Mol	Chain	Res	Type
1	A	183	C
1	A	197	A
1	A	209	U
1	A	210	C
1	A	239	U
1	A	240	G
1	A	243	A
1	A	244	U
1	A	245	U
1	A	247	G
1	A	250	A
1	A	251	G
1	A	253	A
1	A	257	G
1	A	258	G
1	A	266	G
1	A	267	C
1	A	280	C
1	A	289	G
1	A	316	C
1	A	324	G
1	A	328	C
1	A	329	A
1	A	330	C
1	A	332	G
1	A	345	C
1	A	352	C
1	A	354	G
1	A	367	U
1	A	372	C
1	A	373	A
1	A	381	C
1	A	382	A
1	A	397	A
1	A	398	U
1	A	406	G
1	A	408	A
1	A	411	A
1	A	412	A
1	A	414	A
1	A	415	A
1	A	421	U

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Mol	Chain	Res	Type
1	A	422	C
1	A	424	G
1	A	429	U
1	A	430	A
1	A	434	U
1	A	435	A
1	A	438	U
1	A	459	A
1	A	461	A
1	A	462	G
1	A	463	U
1	A	464	U
1	A	465	A
1	A	466	A
1	A	467	U
1	A	468	A
1	A	469	C
1	A	476	U
1	A	481	G
1	A	482	A
1	A	484	G
1	A	485	U
1	A	493	A
1	A	500	G
1	A	511	C
1	A	518	C
1	A	522	C
1	A	527	G
1	A	531	U
1	A	532	A
1	A	547	A
1	A	559	A
1	A	562	U
1	A	572	A
1	A	573	A
1	A	576	C
1	A	577	G
1	A	607	A
1	A	639	G
1	A	650	G
1	A	653	U
1	A	665	A

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Mol	Chain	Res	Type
1	A	693	G
1	A	700	G
1	A	721	G
1	A	724	G
1	A	731	G
1	A	733	G
1	A	747	A
1	A	748	G
1	A	755	G
1	A	777	A
1	A	781	A
1	A	782	A
1	A	793	U
1	A	794	A
1	A	812	G
1	A	815	A
1	A	817	C
1	A	818	G
1	A	819	A
1	A	828	U
1	A	829	G
1	A	841	C
1	A	842	U
1	A	843	U
1	A	844	G
1	A	845	A
1	A	902	G
1	A	914	A
1	A	926	G
1	A	927	G
1	A	931	C
1	A	934	C
1	A	935	A
1	A	960	U
1	A	961	U
1	A	966	G
1	A	968	A
1	A	969	A
1	A	971	G
1	A	974	A
1	A	976	G
1	A	977	A

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Mol	Chain	Res	Type
1	A	983	A
1	A	991	U
1	A	992	U
1	A	993	G
1	A	1002	G
1	A	1004	A
1	A	1009	U
1	A	1022	A
1	A	1025	U
1	A	1026	G
1	A	1030	U
1	A	1031	C
1	A	1034	G
1	A	1035	A
1	A	1043	G
1	A	1044	A
1	A	1045	C
1	A	1049	U
1	A	1050	G
1	A	1051	C
1	A	1053	G
1	A	1054	C
1	A	1055	A
1	A	1065	U
1	A	1066	C
1	A	1070	U
1	A	1094	G
1	A	1095	U
1	A	1101	A
1	A	1102	A
1	A	1108	G
1	A	1111	A
1	A	1112	C
1	A	1126	U
1	A	1130	A
1	A	1136	C
1	A	1137	C
1	A	1138	G
1	A	1139	G
1	A	1140	C
1	A	1158	C
1	A	1159	U

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Mol	Chain	Res	Type
1	A	1168	U
1	A	1169	A
1	A	1174	G
1	A	1179	A
1	A	1181	G
1	A	1183	U
1	A	1196	A
1	A	1197	A
1	A	1201	A
1	A	1202	U
1	A	1209	C
1	A	1212	U
1	A	1213	A
1	A	1214	C
1	A	1215	G
1	A	1225	A
1	A	1226	C
1	A	1227	A
1	A	1228	C
1	A	1238	A
1	A	1240	U
1	A	1241	G
1	A	1253	G
1	A	1256	A
1	A	1258	G
1	A	1261	A
1	A	1270	G
1	A	1278	G
1	A	1279	G
1	A	1280	A
1	A	1281	C
1	A	1285	A
1	A	1286	U
1	A	1287	A
1	A	1298	U
1	A	1300	G
1	A	1301	U
1	A	1305	G
1	A	1316	G
1	A	1317	C
1	A	1320	C
1	A	1322	C

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Mol	Chain	Res	Type
1	A	1323	G
1	A	1336	C
1	A	1338	G
1	A	1346	A
1	A	1347	G
1	A	1353	G
1	A	1362	A
1	A	1363	A
1	A	1364	U
1	A	1378	C
1	A	1379	G
1	A	1380	U
1	A	1381	U
1	A	1398	A
1	A	1403	C
1	A	1404	C
1	A	1409	C
1	A	1419	G
1	A	1432	G
1	A	1446	A
1	A	1447	A
1	A	1451	U
1	A	1452	C
1	A	1490	U
1	A	1493	A
1	A	1497	G
1	A	1499	A
1	A	1503	A
1	A	1506	U
1	A	1507	A
1	A	1517	G
1	A	1519	A
1	A	1520	C
1	A	1528	U
1	A	1529	G
1	A	1530	G
1	A	1533	C
1	A	1534	A

All (25) RNA pucker outliers are listed below:

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Mol	Chain	Res	Type
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Mol	Chain	Res	Type
1	A	51	A
1	A	60	A
1	A	243	A
1	A	279	A
1	A	328	C
1	A	366	A
1	A	372	C
1	A	428	G
1	A	429	U
1	A	484	G
1	A	576	C
1	A	913	A
1	A	960	U
1	A	975	A
1	A	1025	U
1	A	1049	U
1	A	1065	U
1	A	1101	A
1	A	1181	G
1	A	1201	A
1	A	1226	C
1	A	1300	G
1	A	1302	C
1	A	1397	C
1	A	1528	U

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

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

5.5 Carbohydrates [i](#)

There are no oligosaccharides in this entry.

5.6 Ligand geometry [i](#)

There are no ligands in this entry.

5.7 Other polymers [i](#)

There are no such residues in this entry.

5.8 Polymer linkage issues [i](#)

The following chains have linkage breaks:

Mol	Chain	Number of breaks
22	V	5
21	U	2

All chain breaks are listed below:

Model	Chain	Residue-1	Atom-1	Residue-2	Atom-2	Distance (Å)
1	V	245:ASN	C	259:THR	N	14.12
1	V	284:TRP	C	285:HIS	N	12.64
1	V	110:GLU	C	112:THR	N	11.93
1	V	93:LYS	C	101:VAL	N	6.18
1	V	118:PRO	C	124:VAL	N	4.46
1	U	25:ALA	C	26:GLY	N	1.16
1	U	15:LEU	C	16:ARG	N	0.99

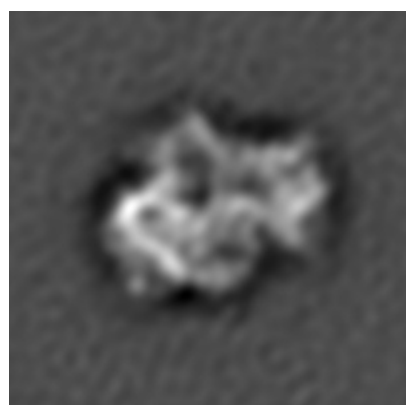
6 Map visualisation [i](#)

This section contains visualisations of the EMDB entry EMD-1895. These allow visual inspection of the internal detail of the map and identification of artifacts.

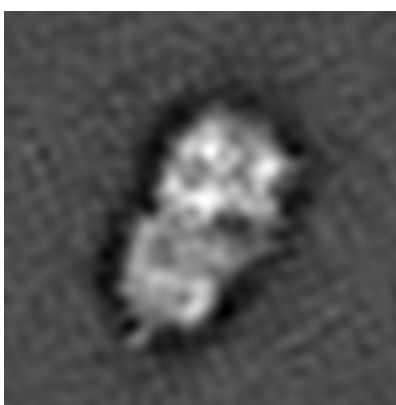
No raw map or half-maps were deposited for this entry and therefore no images, graphs, etc. pertaining to the raw map can be shown.

6.1 Orthogonal projections [i](#)

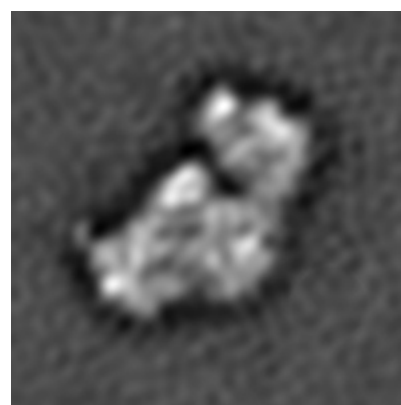
6.1.1 Primary map



X



Y



Z

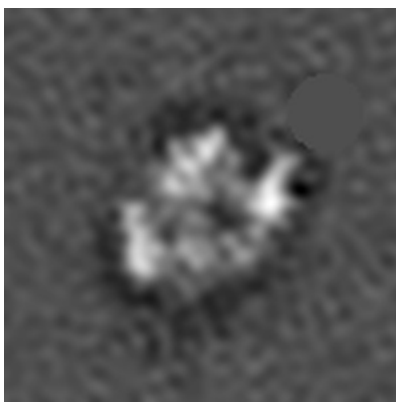
The images above show the map projected in three orthogonal directions.

6.2 Central slices [i](#)

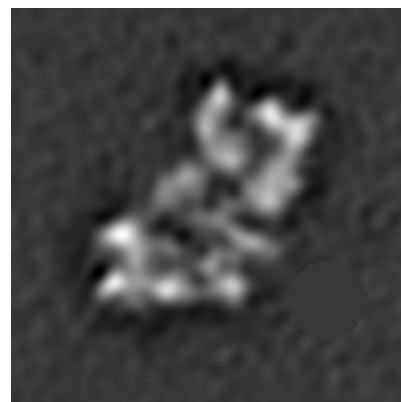
6.2.1 Primary map



X Index: 64



Y Index: 64



Z Index: 64

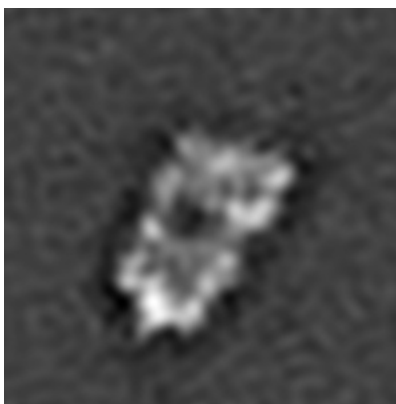
The images above show central slices of the map in three orthogonal directions.

6.3 Largest variance slices [i](#)

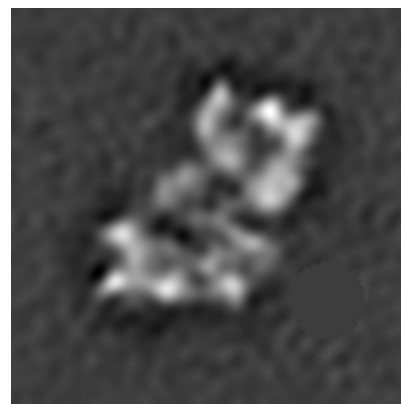
6.3.1 Primary map



X Index: 78



Y Index: 50

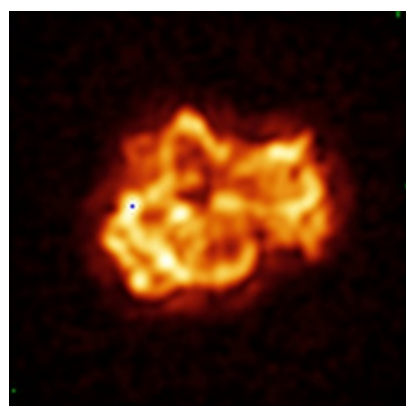


Z Index: 65

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

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

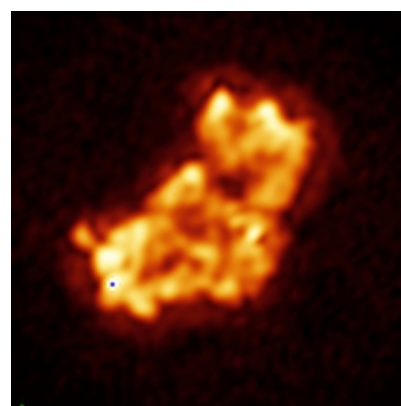
6.4.1 Primary map



X



Y

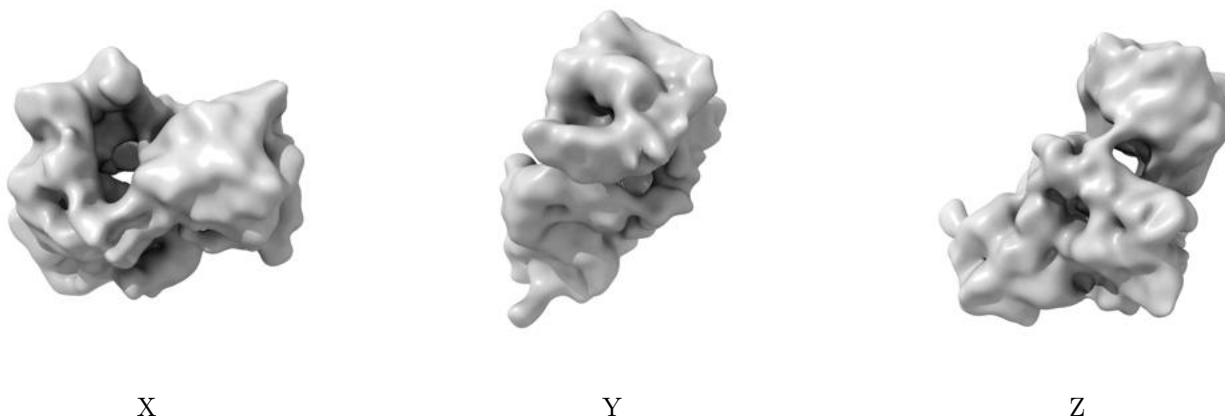


Z

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

6.5 Orthogonal surface views [i](#)

6.5.1 Primary map



The images above show the 3D surface view of the map at the recommended contour level 0.0088. These images, in conjunction with the slice images, may facilitate assessment of whether an appropriate contour level has been provided.

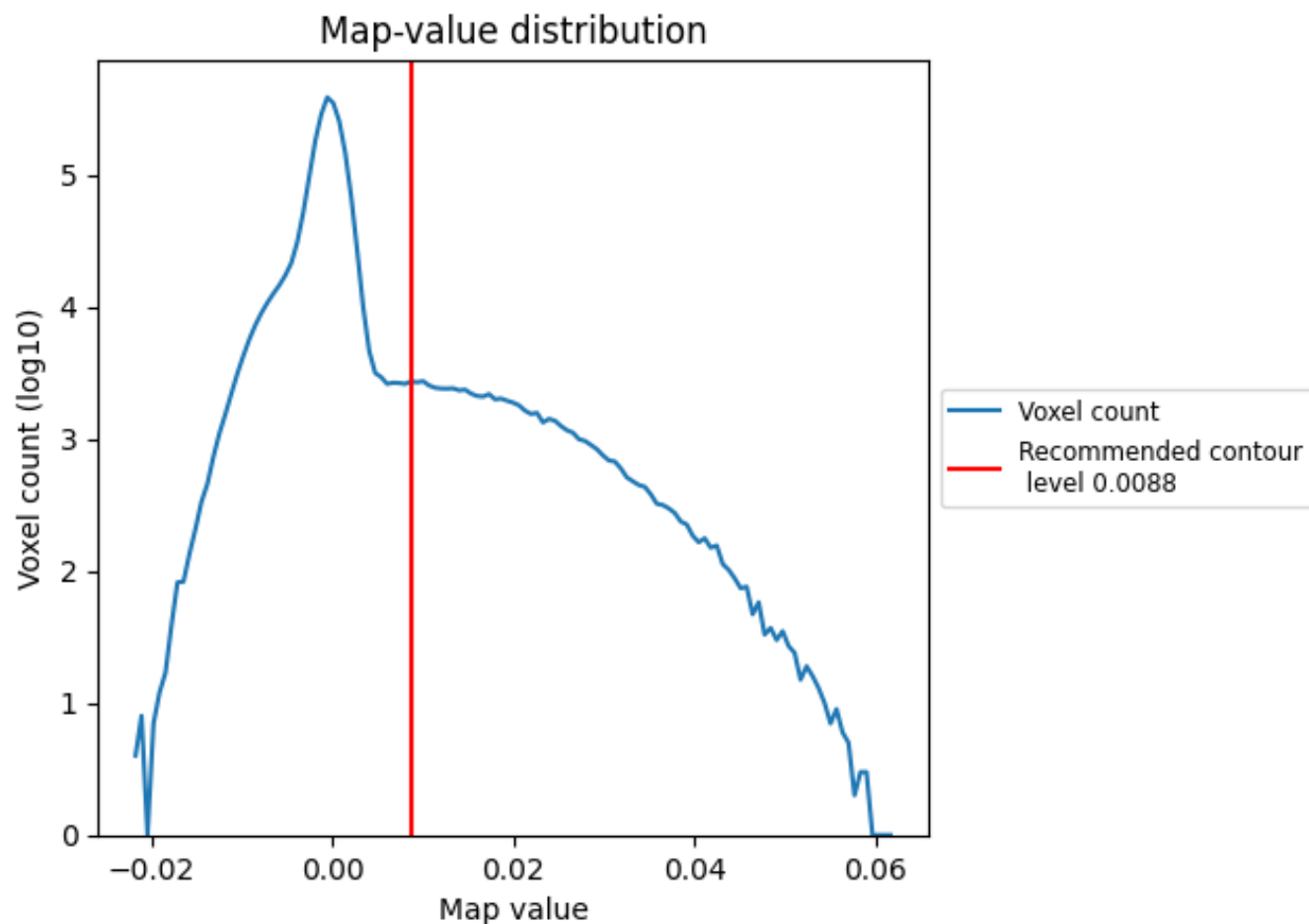
6.6 Mask visualisation [i](#)

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

7 Map analysis [i](#)

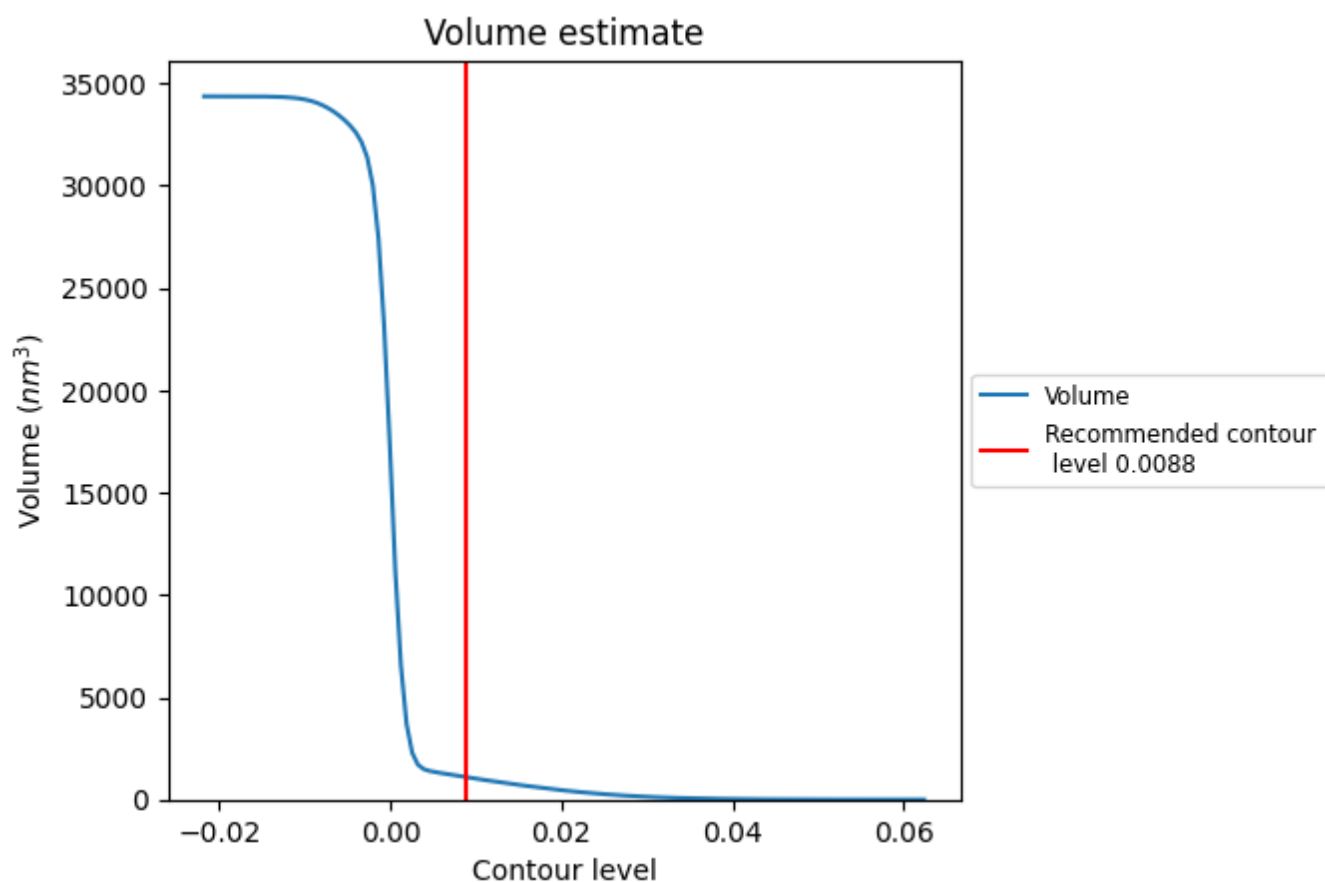
This section contains the results of statistical analysis of the map.

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.

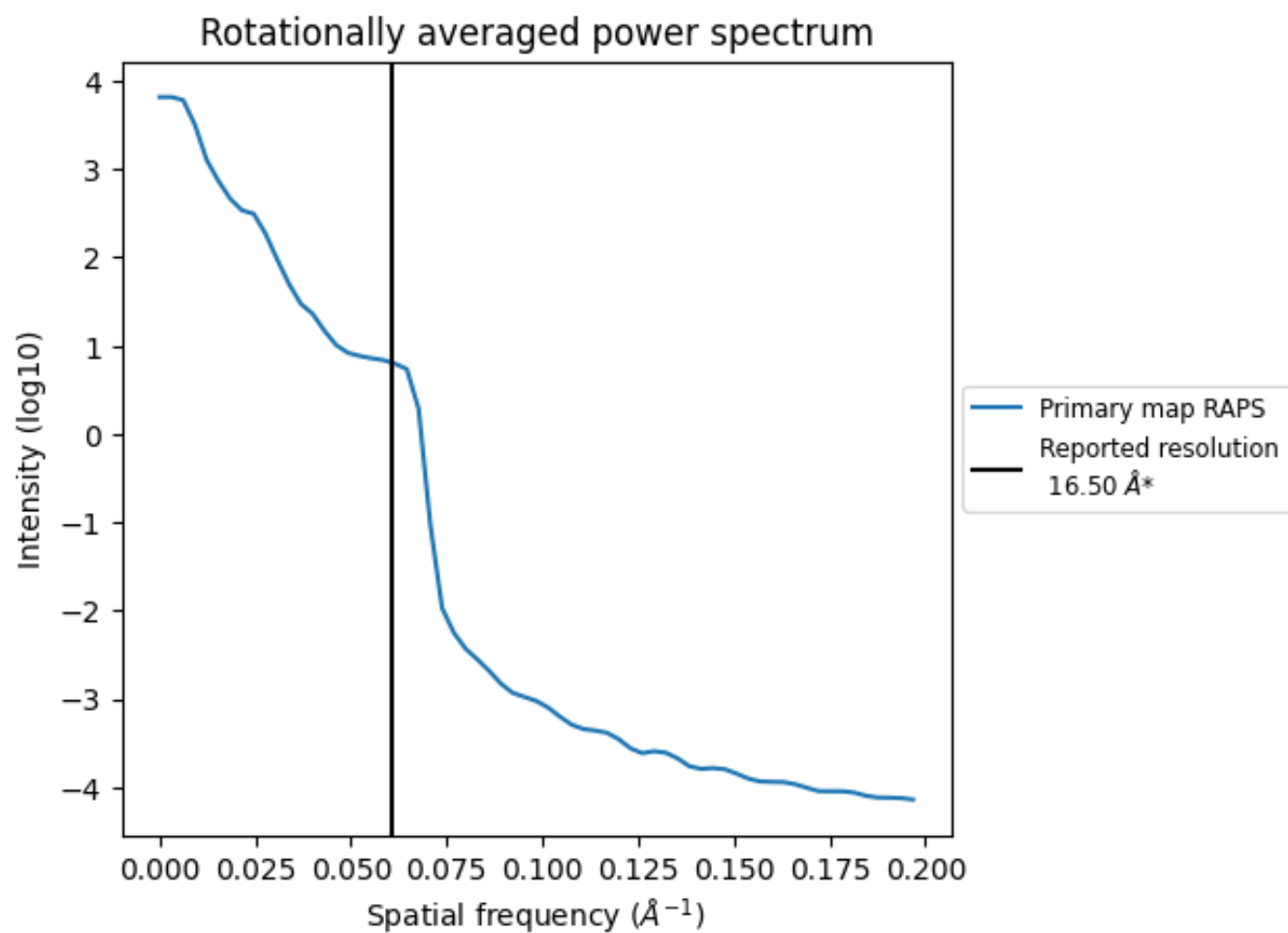
7.2 Volume estimate [i](#)



The volume at the recommended contour level is 1105 nm³; this corresponds to an approximate mass of 998 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 ⓘ



*Reported resolution corresponds to spatial frequency of 0.061 Å⁻¹

8 Fourier-Shell correlation

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

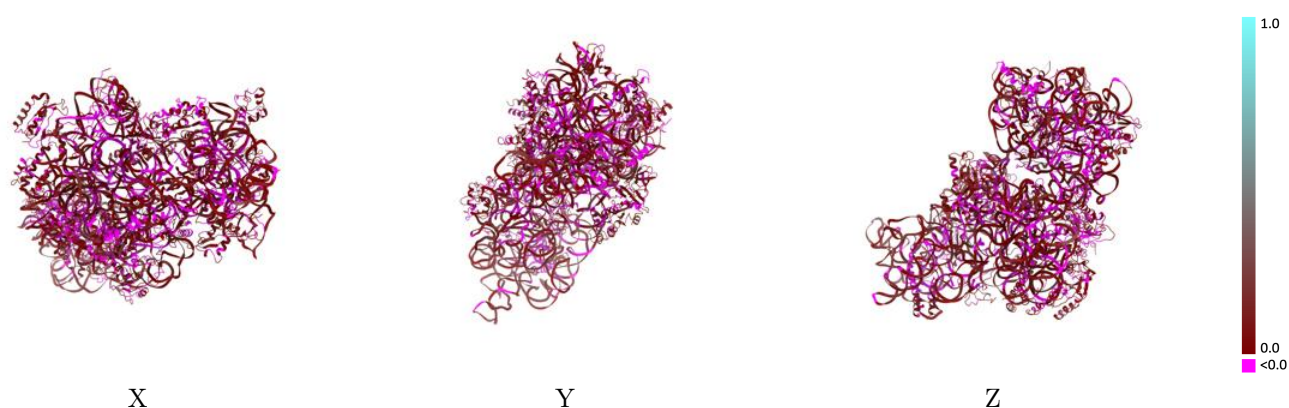
9 Map-model fit [i](#)

This section contains information regarding the fit between EMDB map EMD-1895 and PDB model 4A2I. Per-residue inclusion information can be found in section 3 on page 8.

9.1 Map-model overlay [i](#)

This section was not generated.

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



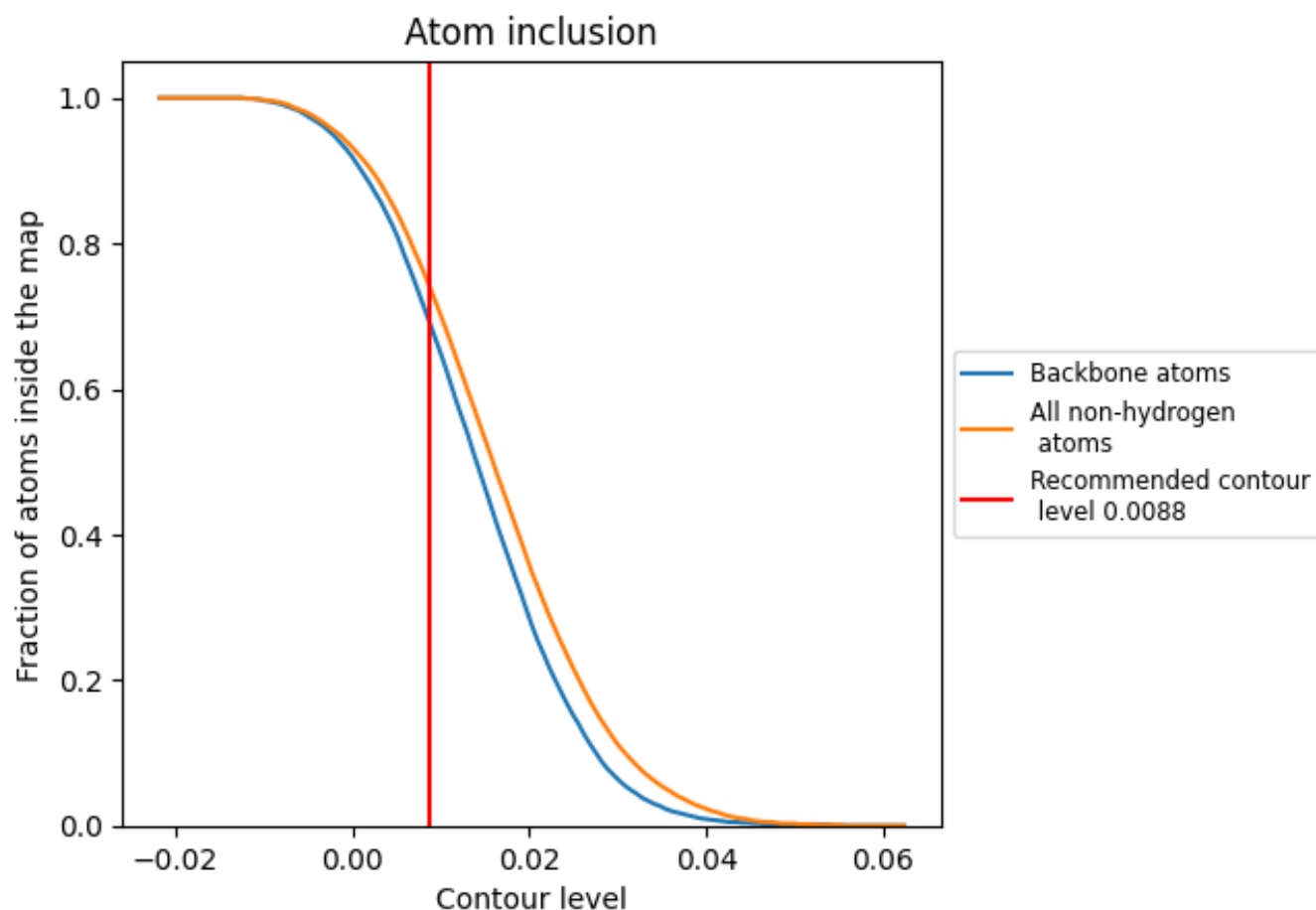
The images above show the model with each residue coloured according to its Q-score. This shows their resolvability in the map with higher Q-score values reflecting better resolvability. Please note: Q-score is calculating the resolvability of atoms, and thus high values are only expected at resolutions 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 (0.0088).

9.4 Atom inclusion [i](#)



At the recommended contour level, 69% of all backbone atoms, 74% 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 (0.0088) and Q-score for the entire model and for each chain.

Chain	Atom inclusion	Q-score
All	 0.7390	 0.0470
A	 0.7970	 0.0590
B	 0.5950	 0.0290
C	 0.5100	 0.0350
D	 0.6060	 0.0280
E	 0.7790	 0.0510
F	 0.6760	 0.0590
G	 0.4970	 -0.0070
H	 0.8110	 0.0410
I	 0.7370	 0.0300
J	 0.7100	 0.0180
K	 0.4480	 -0.0000
L	 0.6470	 0.0260
M	 0.5840	 0.0510
N	 0.7210	 -0.0110
O	 0.6650	 0.0460
P	 0.5250	 -0.0100
Q	 0.8100	 0.0390
R	 0.7320	 0.0380
S	 0.7250	 0.0450
T	 0.6950	 -0.0200
U	 0.0000	 -0.0690
V	 0.7800	 0.0610

