



## Full wwPDB EM Validation Report ⓘ

Nov 4, 2025 – 11:41 AM EST

PDB ID : 9NH0 / pdb\_00009nh0  
EMDB ID : EMD-49398  
Title : In situ cryo-EM structure of PR and DotA-IcmX of the Legionella Dot/Icm T4SS machine at C1 symmetry  
Authors : Yue, J.; Liu, J.  
Deposited on : 2025-02-23  
Resolution : 4.63 Å(reported)

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

We welcome your comments at [validation@mail.wwpdb.org](mailto:validation@mail.wwpdb.org)

A user guide is available at

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

The types of validation reports are described at

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

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

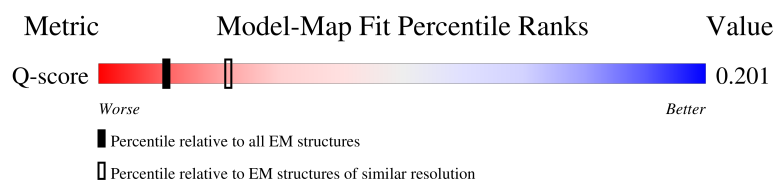
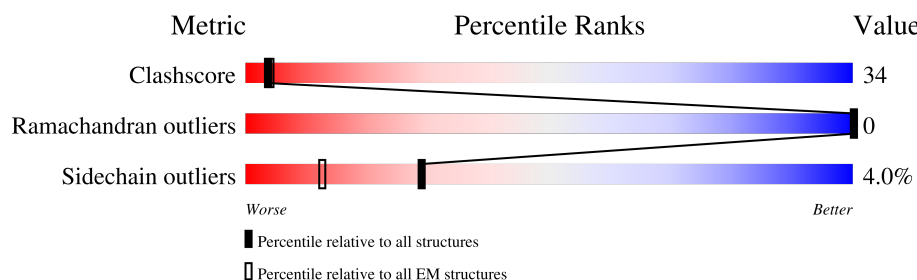
EMDB validation analysis : 0.0.1.dev129  
MolProbity : 4-5-2 with Phenix2.0  
Percentile statistics : 20231227.v01 (using entries in the PDB archive December 27th 2023)  
EM percentile statistics : 202505.v01 (Using data in the EMDb archive up until May 2025)  
MapQ : 1.9.13  
Ideal geometry (proteins) : Engh & Huber (2001)  
Ideal geometry (DNA, RNA) : Parkinson et al. (1996)  
Validation Pipeline (wwPDB-VP) : 2.46

# 1 Overall quality at a glance

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

The reported resolution of this entry is 4.63 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	210492	15764	-
Ramachandran outliers	207382	16835	-
Sidechain outliers	206894	16415	-
Q-score	-	25397	2068 ( 4.13 - 5.13 )

The table below summarises the geometric issues observed across the polymeric chains and their fit to the map. The red, orange, yellow and green segments of the bar indicate the fraction of residues that contain outliers for  $\geq 3$ , 2, 1 and 0 types of geometric quality criteria respectively. A grey segment represents the fraction of residues that are not modelled. The numeric value for each fraction is indicated below the corresponding segment, with a dot representing fractions  $\leq 5\%$ . 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	Aa	16	<div> <div>100%</div> <div>62% 38%</div> </div>
1	Ag	16	<div> <div>100%</div> <div>50% 50%</div> </div>
1	Am	16	<div> <div>100%</div> <div>31% 62% 6%</div> </div>
1	As	16	<div> <div>100%</div> <div>69% 31%</div> </div>

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Mol	Chain	Length	Quality of chain
1	Ay	16	
1	Be	16	
1	Bk	16	
1	Bq	16	
1	Bw	16	
1	Cc	16	
1	Ci	16	
1	Co	16	
1	Cu	16	
1	Da	16	
1	Dg	16	
1	Dm	16	
1	Ds	16	
1	Dy	16	
2	Ab	9	
2	Ah	9	
2	An	9	
2	At	9	
2	Az	9	
2	Bf	9	
2	Bl	9	
2	Br	9	
2	Bx	9	
2	Cd	9	
2	Cj	9	

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Mol	Chain	Length	Quality of chain
2	Cp	9	
2	Cv	9	
2	Db	9	
2	Dh	9	
2	Dn	9	
2	Dt	9	
2	Dz	9	
3	Ac	16	
3	Ai	16	
3	Ao	16	
3	Au	16	
3	Ba	16	
3	Bg	16	
3	Bm	16	
3	Bs	16	
3	By	16	
3	Ce	16	
3	Ck	16	
3	Cq	16	
3	Cw	16	
3	Dc	16	
3	Di	16	
3	Do	16	
3	Du	16	
3	Ea	16	

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Mol	Chain	Length	Quality of chain
4	Ad	5	
4	Aj	5	
4	Ap	5	
4	Av	5	
4	Bb	5	
4	Bh	5	
4	Bn	5	
4	Bt	5	
4	Bz	5	
4	Cf	5	
4	Cl	5	
4	Cr	5	
4	Cx	5	
4	Dd	5	
4	Dj	5	
4	Dp	5	
4	Dv	5	
4	Eb	5	
5	Ae	269	
5	Ak	269	
5	Aq	269	
5	Aw	269	
5	Bc	269	
5	Bi	269	
5	Bo	269	

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Mol	Chain	Length	Quality of chain
5	Bu	269	
5	Ca	269	
5	Cg	269	
5	Cm	269	
5	Cs	269	
5	Cy	269	
5	De	269	
5	Dk	269	
5	Dq	269	
5	Dw	269	
5	Ec	269	
6	Af	361	
6	Al	361	
6	Ar	361	
6	Ax	361	
6	Bd	361	
6	Bj	361	
6	Bp	361	
6	Bv	361	
6	Cb	361	
6	Ch	361	
6	Cn	361	
6	Ct	361	
6	Cz	361	
6	Df	361	

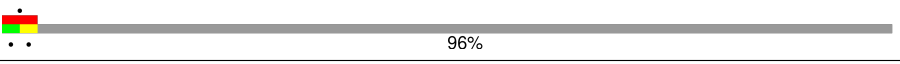
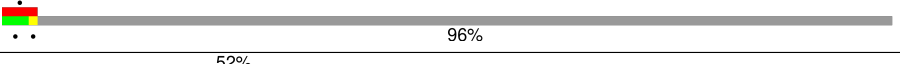










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Mol	Chain	Length	Quality of chain
6	Dl	361	
6	Dr	361	
6	Dx	361	
6	Ed	361	
7	Ee	1048	
7	Ef	1048	
7	Eg	1048	
7	Eh	1048	
7	Ei	1048	
7	Ej	1048	
7	Ek	1048	
7	El	1048	
7	Em	1048	
7	En	1048	
7	Eo	1048	
7	Ep	1048	
7	Eq	1048	
7	Er	1048	
7	Es	1048	
7	Et	1048	
7	Eu	1048	
7	Ev	1048	
7	Fg	1048	
7	Fh	1048	
7	Fi	1048	

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Mol	Chain	Length	Quality of chain
7	Fj	1048	 96%
7	Fk	1048	 96%
8	Ew	466	 52% 43% 44% 11%
8	Ex	466	 50% 43% 45% 11%
8	Ey	466	 48% 40% 48% 11%
8	Ez	466	 50% 44% 43% 11%
8	Fa	466	 49% 42% 45% 11%
9	Fb	1048	 74% 40% 40% 18%
9	Fc	1048	 77% 40% 39% 18%
9	Fd	1048	 75% 39% 41% 18%
9	Fe	1048	 75% 42% 37% 18%
9	Ff	1048	 75% 40% 40% 18%



## 2 Entry composition [i](#)

There are 9 unique types of molecules in this entry. The entry contains 94015 atoms, of which 0 are hydrogens and 0 are deuteriums.

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

- Molecule 1 is a protein called unknown peptide E.

Mol	Chain	Residues	Atoms					AltConf	Trace
1	Aa	16	Total	C	N	O	S	0	0
			117	71	20	25	1		
1	Ag	16	Total	C	N	O	S	0	0
			117	71	20	25	1		
1	Am	16	Total	C	N	O	S	0	0
			117	71	20	25	1		
1	As	16	Total	C	N	O	S	0	0
			117	71	20	25	1		
1	Ay	16	Total	C	N	O	S	0	0
			117	71	20	25	1		
1	Be	16	Total	C	N	O	S	0	0
			117	71	20	25	1		
1	Bk	16	Total	C	N	O	S	0	0
			117	71	20	25	1		
1	Bq	16	Total	C	N	O	S	0	0
			117	71	20	25	1		
1	Bw	16	Total	C	N	O	S	0	0
			117	71	20	25	1		
1	Cc	16	Total	C	N	O	S	0	0
			117	71	20	25	1		
1	Ci	16	Total	C	N	O	S	0	0
			117	71	20	25	1		
1	Co	16	Total	C	N	O	S	0	0
			117	71	20	25	1		
1	Cu	16	Total	C	N	O	S	0	0
			117	71	20	25	1		
1	Da	16	Total	C	N	O	S	0	0
			117	71	20	25	1		
1	Dg	16	Total	C	N	O	S	0	0
			117	71	20	25	1		
1	Dm	16	Total	C	N	O	S	0	0
			117	71	20	25	1		
1	Ds	16	Total	C	N	O	S	0	0
			117	71	20	25	1		

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Mol	Chain	Residues	Atoms					AltConf	Trace
1	Dy	16	Total	C	N	O	S	0	0
			117	71	20	25	1		

- Molecule 2 is a protein called unknown peptide F.

Mol	Chain	Residues	Atoms					AltConf	Trace
2	Ab	9	Total	C	N	O	S	0	0
			61	35	12	12	2		
2	Ah	9	Total	C	N	O	S	0	0
			61	35	12	12	2		
2	An	9	Total	C	N	O	S	0	0
			61	35	12	12	2		
2	At	9	Total	C	N	O	S	0	0
			61	35	12	12	2		
2	Az	9	Total	C	N	O	S	0	0
			61	35	12	12	2		
2	Bf	9	Total	C	N	O	S	0	0
			61	35	12	12	2		
2	Bl	9	Total	C	N	O	S	0	0
			61	35	12	12	2		
2	Br	9	Total	C	N	O	S	0	0
			61	35	12	12	2		
2	Bx	9	Total	C	N	O	S	0	0
			61	35	12	12	2		
2	Cd	9	Total	C	N	O	S	0	0
			61	35	12	12	2		
2	Cj	9	Total	C	N	O	S	0	0
			61	35	12	12	2		
2	Cp	9	Total	C	N	O	S	0	0
			61	35	12	12	2		
2	Cv	9	Total	C	N	O	S	0	0
			61	35	12	12	2		
2	Db	9	Total	C	N	O	S	0	0
			61	35	12	12	2		
2	Dh	9	Total	C	N	O	S	0	0
			61	35	12	12	2		
2	Dn	9	Total	C	N	O	S	0	0
			61	35	12	12	2		
2	Dt	9	Total	C	N	O	S	0	0
			61	35	12	12	2		
2	Dz	9	Total	C	N	O	S	0	0
			61	35	12	12	2		

- Molecule 3 is a protein called unknown peptide G.

Mol	Chain	Residues	Atoms				AltConf	Trace
3	Ac	16	Total	C	N	O	0	0
			125	78	21	26		
3	Ai	16	Total	C	N	O	0	0
			125	78	21	26		
3	Ao	16	Total	C	N	O	0	0
			125	78	21	26		
3	Au	16	Total	C	N	O	0	0
			125	78	21	26		
3	Ba	16	Total	C	N	O	0	0
			125	78	21	26		
3	Bg	16	Total	C	N	O	0	0
			125	78	21	26		
3	Bm	16	Total	C	N	O	0	0
			125	78	21	26		
3	Bs	16	Total	C	N	O	0	0
			125	78	21	26		
3	By	16	Total	C	N	O	0	0
			125	78	21	26		
3	Ce	16	Total	C	N	O	0	0
			125	78	21	26		
3	Ck	16	Total	C	N	O	0	0
			125	78	21	26		
3	Cq	16	Total	C	N	O	0	0
			125	78	21	26		
3	Cw	16	Total	C	N	O	0	0
			125	78	21	26		
3	Dc	16	Total	C	N	O	0	0
			125	78	21	26		
3	Di	16	Total	C	N	O	0	0
			125	78	21	26		
3	Do	16	Total	C	N	O	0	0
			125	78	21	26		
3	Du	16	Total	C	N	O	0	0
			125	78	21	26		
3	Ea	16	Total	C	N	O	0	0
			125	78	21	26		

- Molecule 4 is a protein called unknown peptide H.

Mol	Chain	Residues	Atoms				AltConf	Trace
4	Ad	5	Total	C	N	O	0	0
			36	23	5	8		

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Mol	Chain	Residues	Atoms				AltConf	Trace
4	Aj	5	Total	C	N	O	0	0
			36	23	5	8		
4	Ap	5	Total	C	N	O	0	0
			36	23	5	8		
4	Av	5	Total	C	N	O	0	0
			36	23	5	8		
4	Bb	5	Total	C	N	O	0	0
			36	23	5	8		
4	Bh	5	Total	C	N	O	0	0
			36	23	5	8		
4	Bn	5	Total	C	N	O	0	0
			36	23	5	8		
4	Bt	5	Total	C	N	O	0	0
			36	23	5	8		
4	Bz	5	Total	C	N	O	0	0
			36	23	5	8		
4	Cf	5	Total	C	N	O	0	0
			36	23	5	8		
4	Cl	5	Total	C	N	O	0	0
			36	23	5	8		
4	Cr	5	Total	C	N	O	0	0
			36	23	5	8		
4	Cx	5	Total	C	N	O	0	0
			36	23	5	8		
4	Dd	5	Total	C	N	O	0	0
			36	23	5	8		
4	Dj	5	Total	C	N	O	0	0
			36	23	5	8		
4	Dp	5	Total	C	N	O	0	0
			36	23	5	8		
4	Dv	5	Total	C	N	O	0	0
			36	23	5	8		
4	Eb	5	Total	C	N	O	0	0
			36	23	5	8		

- Molecule 5 is a protein called IcmG (DotF).

Mol	Chain	Residues	Atoms					AltConf	Trace
5	Ae	63	Total	C	N	O	S	0	0
			484	308	84	91	1		
5	Ak	63	Total	C	N	O	S	0	0
			484	308	84	91	1		

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Mol	Chain	Residues	Atoms					AltConf	Trace
5	Aq	63	Total 484	C 308	N 84	O 91	S 1	0	0
5	Aw	63	Total 484	C 308	N 84	O 91	S 1	0	0
5	Bc	63	Total 484	C 308	N 84	O 91	S 1	0	0
5	Bi	63	Total 484	C 308	N 84	O 91	S 1	0	0
5	Bo	63	Total 484	C 308	N 84	O 91	S 1	0	0
5	Bu	63	Total 484	C 308	N 84	O 91	S 1	0	0
5	Ca	63	Total 484	C 308	N 84	O 91	S 1	0	0
5	Cg	63	Total 484	C 308	N 84	O 91	S 1	0	0
5	Cm	63	Total 484	C 308	N 84	O 91	S 1	0	0
5	Cs	63	Total 484	C 308	N 84	O 91	S 1	0	0
5	Cy	63	Total 484	C 308	N 84	O 91	S 1	0	0
5	De	63	Total 484	C 308	N 84	O 91	S 1	0	0
5	Dk	63	Total 484	C 308	N 84	O 91	S 1	0	0
5	Dq	63	Total 484	C 308	N 84	O 91	S 1	0	0
5	Dw	63	Total 484	C 308	N 84	O 91	S 1	0	0
5	Ec	63	Total 484	C 308	N 84	O 91	S 1	0	0

- Molecule 6 is a protein called IcmK (DotH).

Mol	Chain	Residues	Atoms					AltConf	Trace
6	Af	160	Total 1238	C 795	N 207	O 233	S 3	0	0
6	Al	160	Total 1238	C 795	N 207	O 233	S 3	0	0
6	Ar	160	Total 1238	C 795	N 207	O 233	S 3	0	0

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Mol	Chain	Residues	Atoms					AltConf	Trace
6	Ax	160	Total	C	N	O	S	0	0
			1238	795	207	233	3		
6	Bd	160	Total	C	N	O	S	0	0
			1238	795	207	233	3		
6	Bj	160	Total	C	N	O	S	0	0
			1238	795	207	233	3		
6	Bp	160	Total	C	N	O	S	0	0
			1238	795	207	233	3		
6	Bv	160	Total	C	N	O	S	0	0
			1238	795	207	233	3		
6	Cb	160	Total	C	N	O	S	0	0
			1238	795	207	233	3		
6	Ch	160	Total	C	N	O	S	0	0
			1238	795	207	233	3		
6	Cn	160	Total	C	N	O	S	0	0
			1238	795	207	233	3		
6	Ct	160	Total	C	N	O	S	0	0
			1238	795	207	233	3		
6	Cz	160	Total	C	N	O	S	0	0
			1238	795	207	233	3		
6	Df	160	Total	C	N	O	S	0	0
			1238	795	207	233	3		
6	DI	160	Total	C	N	O	S	0	0
			1238	795	207	233	3		
6	Dr	160	Total	C	N	O	S	0	0
			1238	795	207	233	3		
6	Dx	160	Total	C	N	O	S	0	0
			1238	795	207	233	3		
6	Ed	160	Total	C	N	O	S	0	0
			1238	795	207	233	3		

- Molecule 7 is a protein called IcmE (DotG).

Mol	Chain	Residues	Atoms					AltConf	Trace
7	Ee	34	Total	C	N	O	S	0	0
			276	168	47	60	1		
7	Ef	59	Total	C	N	O	S	0	0
			472	287	81	103	1		
7	Eg	34	Total	C	N	O	S	0	0
			276	168	47	60	1		
7	Eh	48	Total	C	N	O	S	0	0
			397	244	68	84	1		

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Mol	Chain	Residues	Atoms					AltConf	Trace
7	Ei	49	Total	C	N	O	S	0	0
			402	247	69	85	1		
7	Ej	58	Total	C	N	O	S	0	0
			464	283	80	100	1		
7	Ek	34	Total	C	N	O	S	0	0
			276	168	47	60	1		
7	El	34	Total	C	N	O	S	0	0
			276	168	47	60	1		
7	Em	46	Total	C	N	O	S	0	0
			381	232	66	82	1		
7	En	34	Total	C	N	O	S	0	0
			276	168	47	60	1		
7	Eo	34	Total	C	N	O	S	0	0
			276	168	47	60	1		
7	Ep	34	Total	C	N	O	S	0	0
			276	168	47	60	1		
7	Eq	57	Total	C	N	O	S	0	0
			459	280	79	99	1		
7	Er	34	Total	C	N	O	S	0	0
			276	168	47	60	1		
7	Es	56	Total	C	N	O	S	0	0
			454	277	78	98	1		
7	Et	48	Total	C	N	O	S	0	0
			397	244	68	84	1		
7	Eu	59	Total	C	N	O	S	0	0
			472	287	81	103	1		
7	Ev	34	Total	C	N	O	S	0	0
			276	168	47	60	1		
7	Fg	43	Total	C	N	O	S	0	0
			334	221	58	54	1		
7	Fh	43	Total	C	N	O	S	0	0
			334	221	58	54	1		
7	Fi	43	Total	C	N	O	S	0	0
			334	221	58	54	1		
7	Fj	43	Total	C	N	O	S	0	0
			334	221	58	54	1		
7	Fk	43	Total	C	N	O	S	0	0
			334	221	58	54	1		

- Molecule 8 is a protein called IcmX (IcmY).

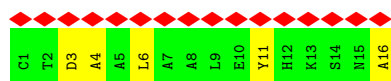
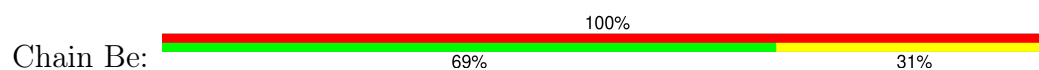
Mol	Chain	Residues	Atoms					AltConf	Trace
8	Ew	417	Total	C	N	O	S	0	0
			3213	2013	542	646	12		
8	Ex	417	Total	C	N	O	S	0	0
			3213	2013	542	646	12		
8	Ey	417	Total	C	N	O	S	0	0
			3213	2013	542	646	12		
8	Ez	417	Total	C	N	O	S	0	0
			3213	2013	542	646	12		
8	Fa	417	Total	C	N	O	S	0	0
			3213	2013	542	646	12		

- Molecule 9 is a protein called DotA.

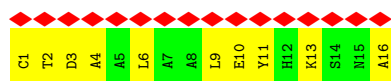
Mol	Chain	Residues	Atoms					AltConf	Trace
9	Fb	857	Total	C	N	O	S	0	0
			6560	4256	1051	1202	51		
9	Fc	857	Total	C	N	O	S	0	0
			6560	4256	1051	1202	51		
9	Fd	857	Total	C	N	O	S	0	0
			6560	4256	1051	1202	51		
9	Fe	857	Total	C	N	O	S	0	0
			6560	4256	1051	1202	51		
9	Ff	857	Total	C	N	O	S	0	0
			6560	4256	1051	1202	51		







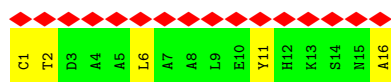
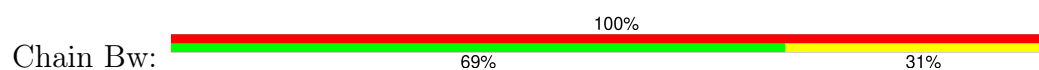
- Molecule 1: unknown peptide E



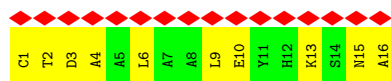
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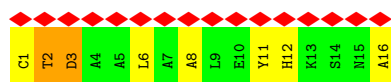
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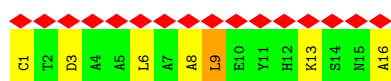
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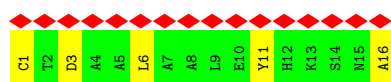
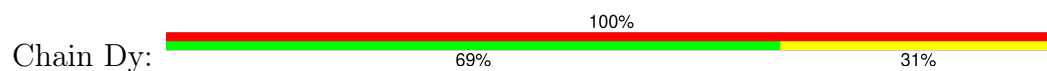
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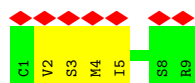
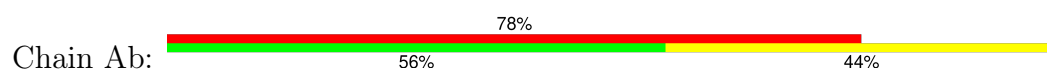
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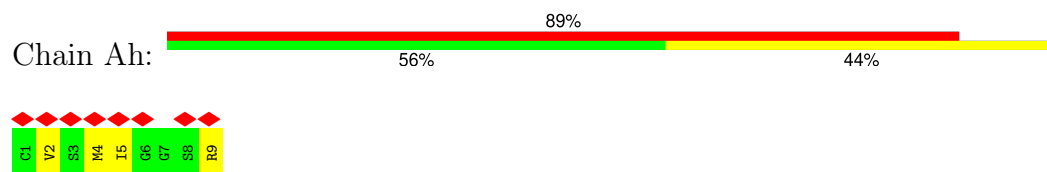
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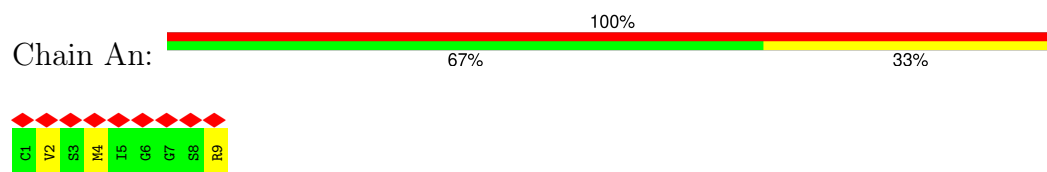
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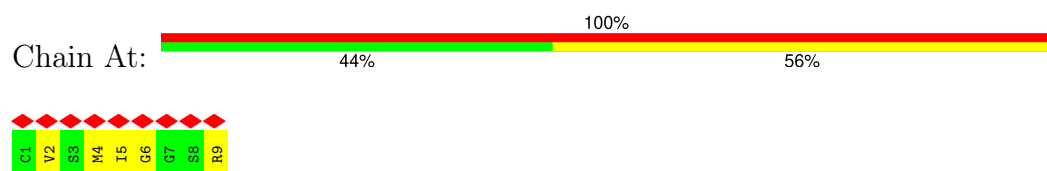
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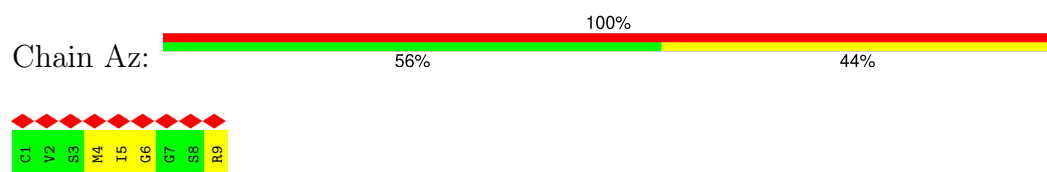
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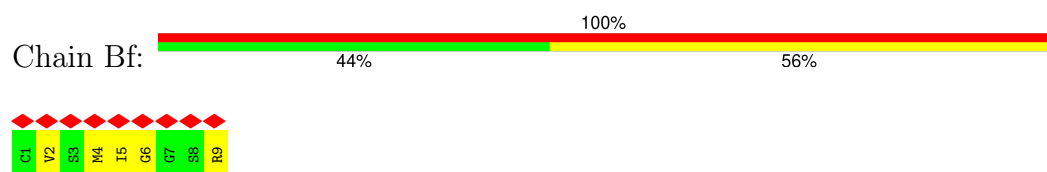
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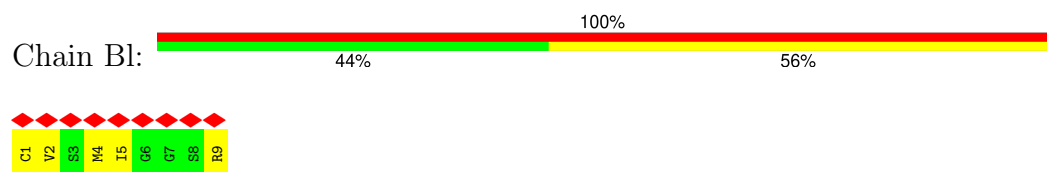
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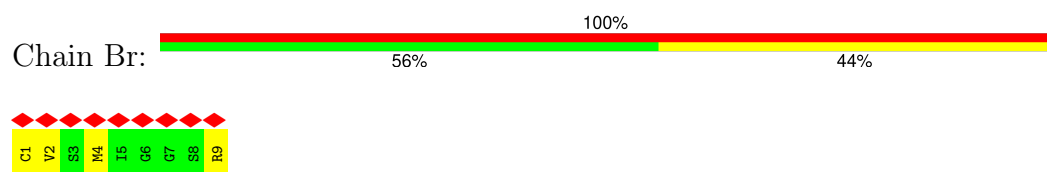
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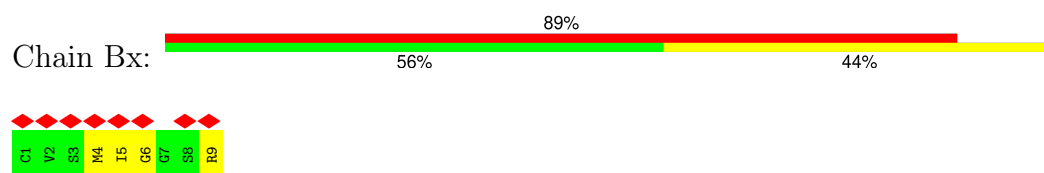
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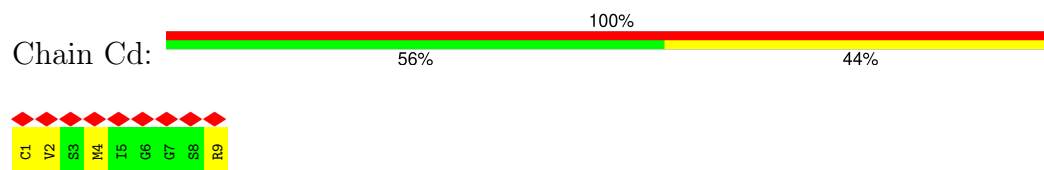
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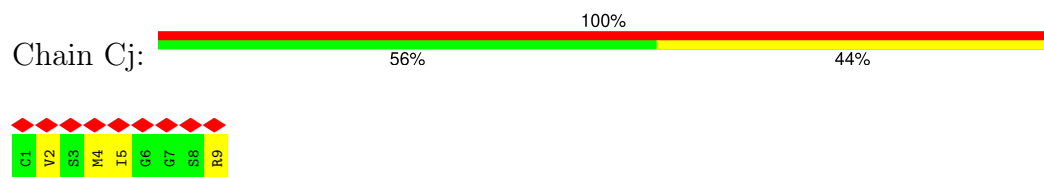
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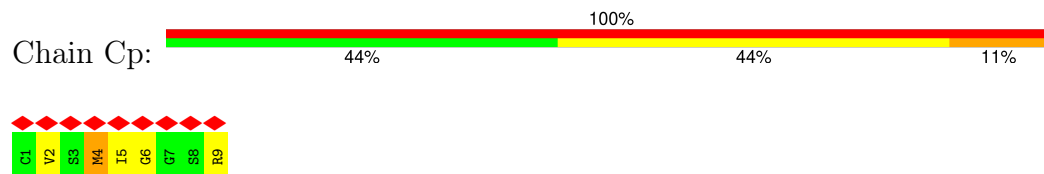
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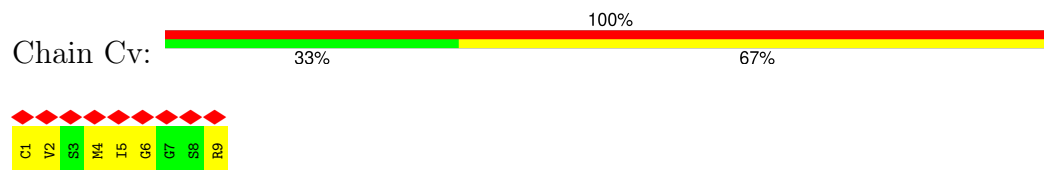
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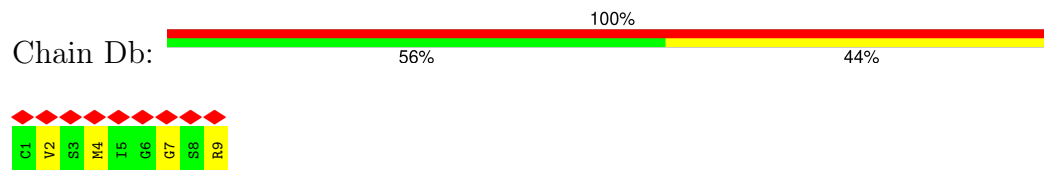
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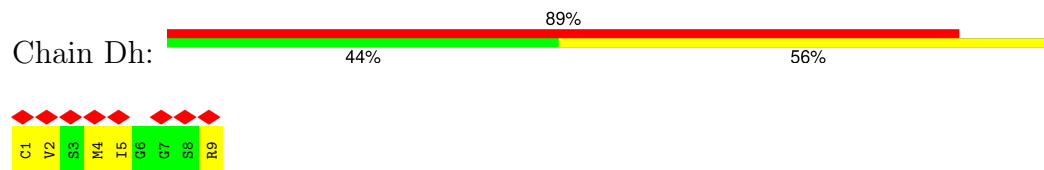
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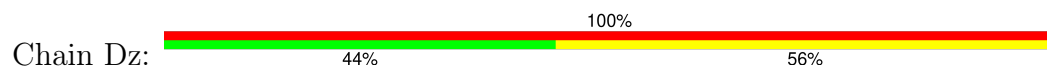
- Molecule 2: unknown peptide F



- Molecule 2: unknown peptide F



- Molecule 2: unknown peptide F



- Molecule 3: unknown peptide G



- Molecule 3: unknown peptide G



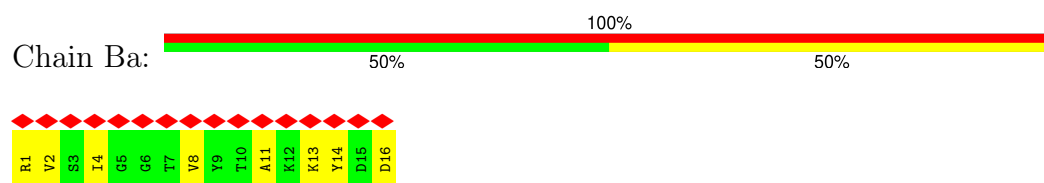
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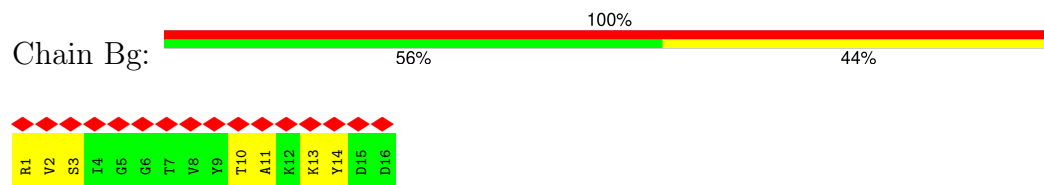
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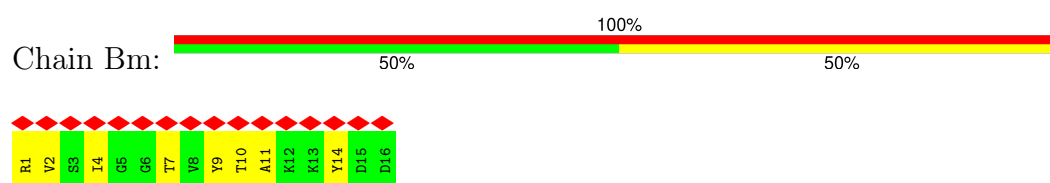
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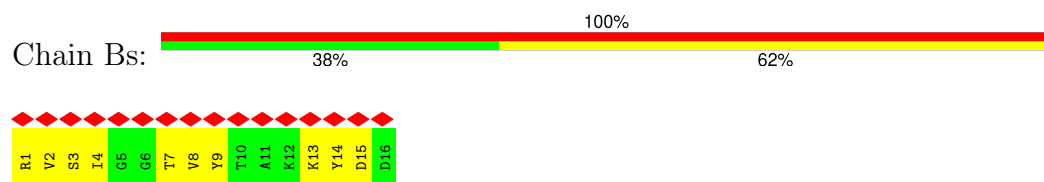
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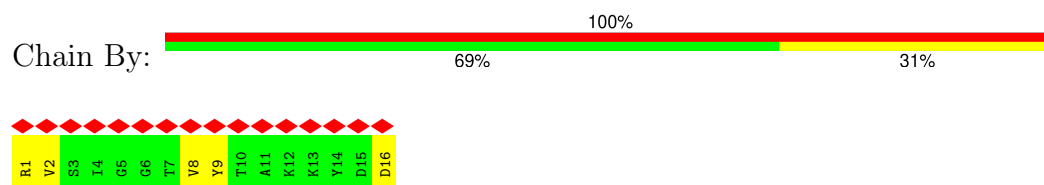
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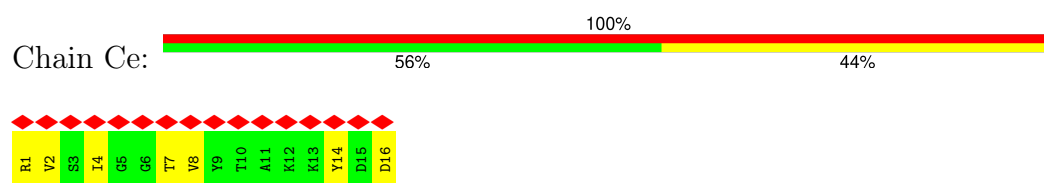
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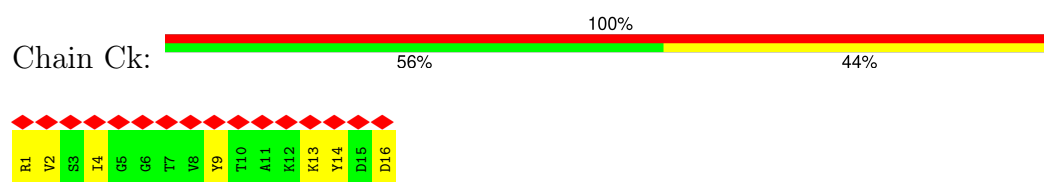
- Molecule 3: unknown peptide G



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- Molecule 3: unknown peptide G



## ● Molecule 3: unknown peptide G



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## ● Molecule 3: unknown peptide G



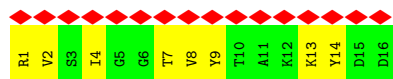
## ● Molecule 3: unknown peptide G



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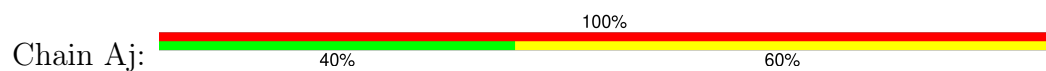




- Molecule 4: unknown peptide H



- Molecule 4: unknown peptide H



- Molecule 4: unknown peptide H



- Molecule 4: unknown peptide H



- Molecule 4: unknown peptide H



- Molecule 4: unknown peptide H



- Molecule 4: unknown peptide H



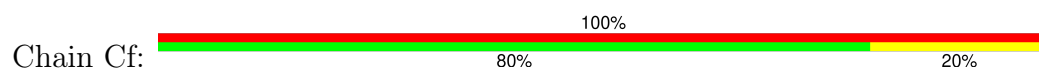
- Molecule 4: unknown peptide H



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- Molecule 4: unknown peptide H



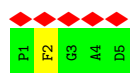
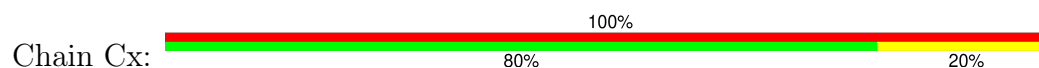
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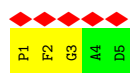
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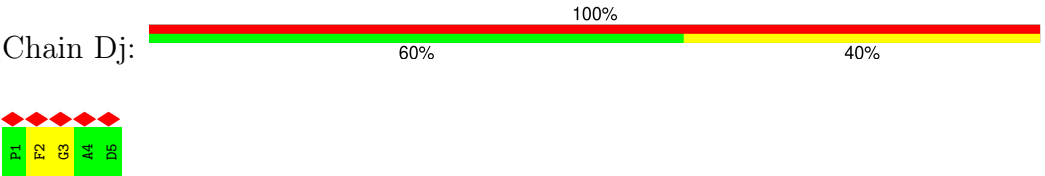
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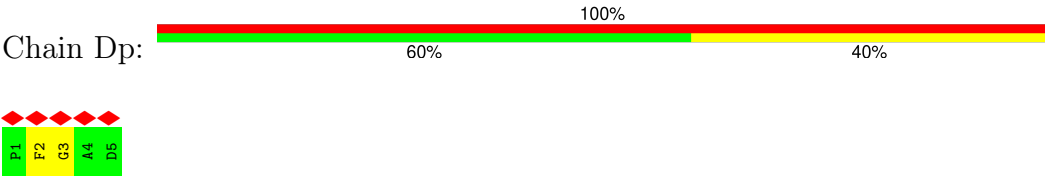
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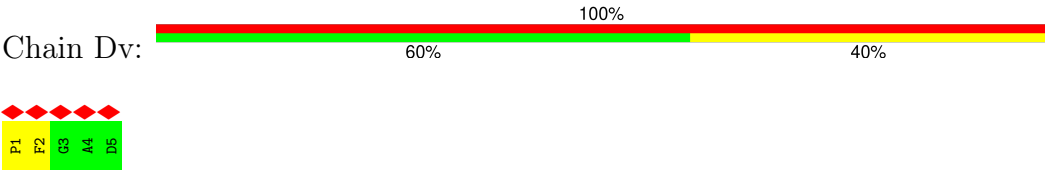
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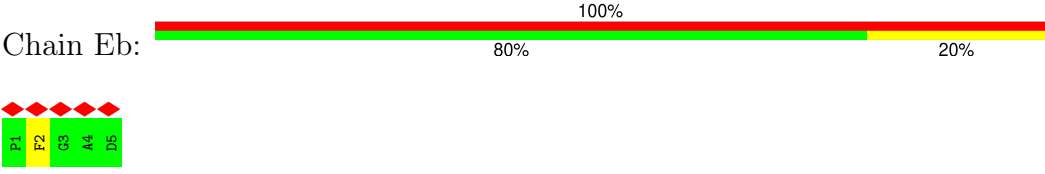
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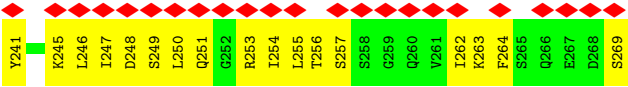
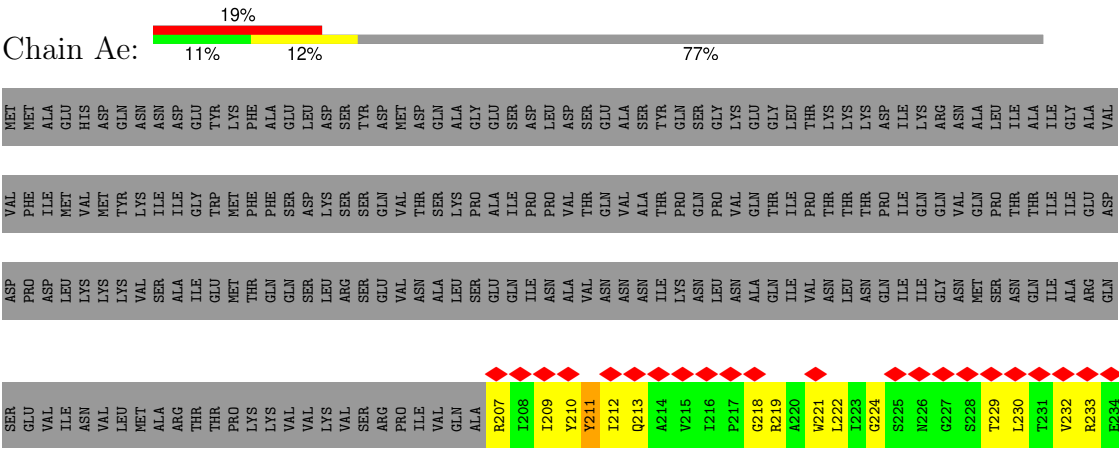
• Molecule 4: unknown peptide H



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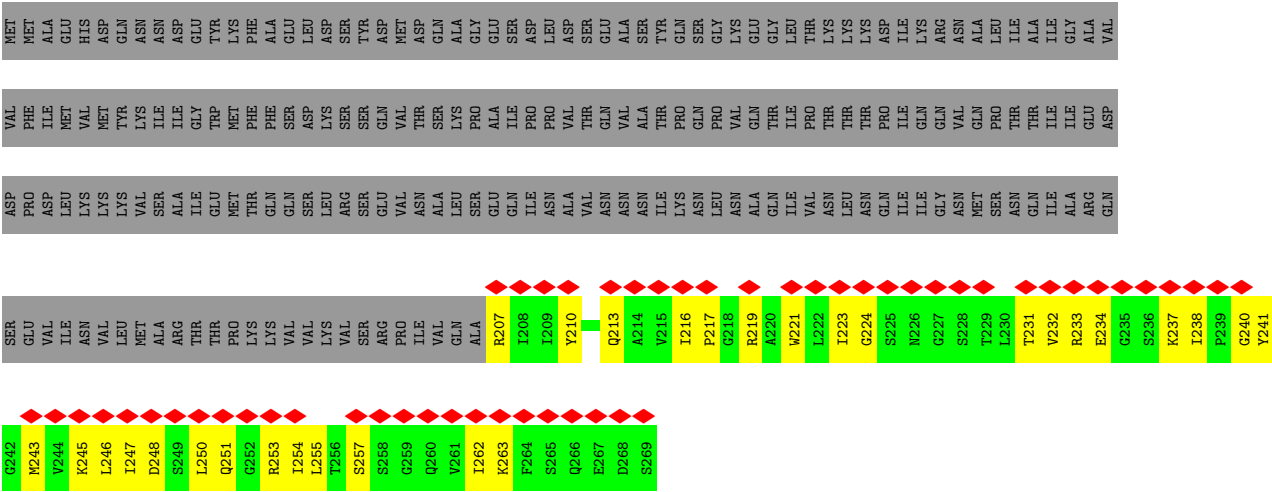


• Molecule 5: IcmG (DotF)

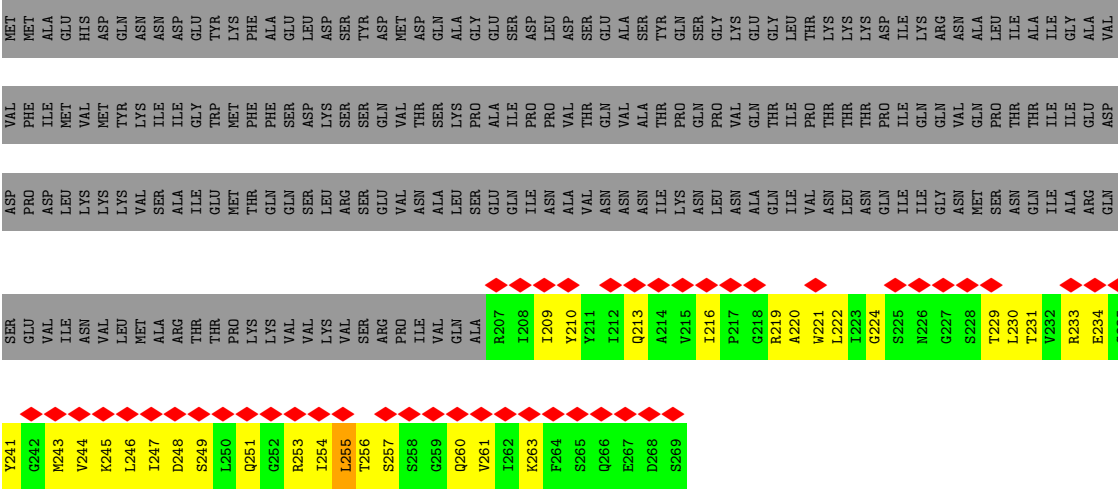


• Molecule 5: IcmG (DotF)

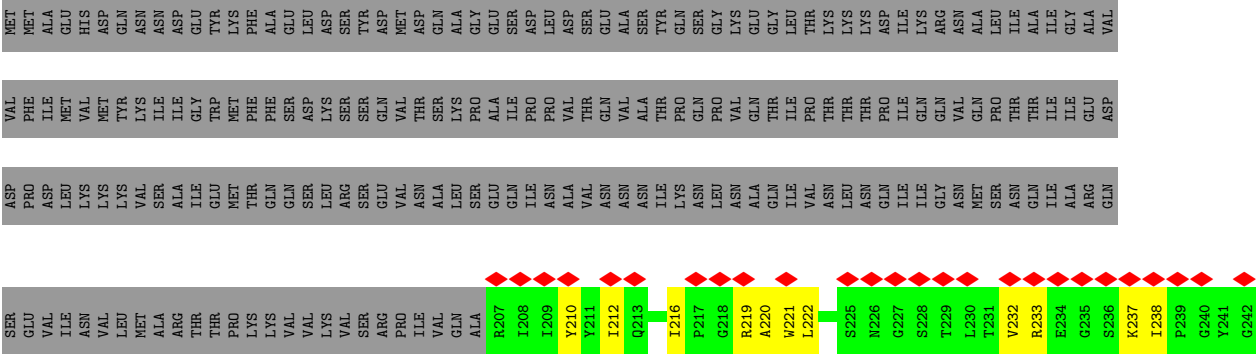




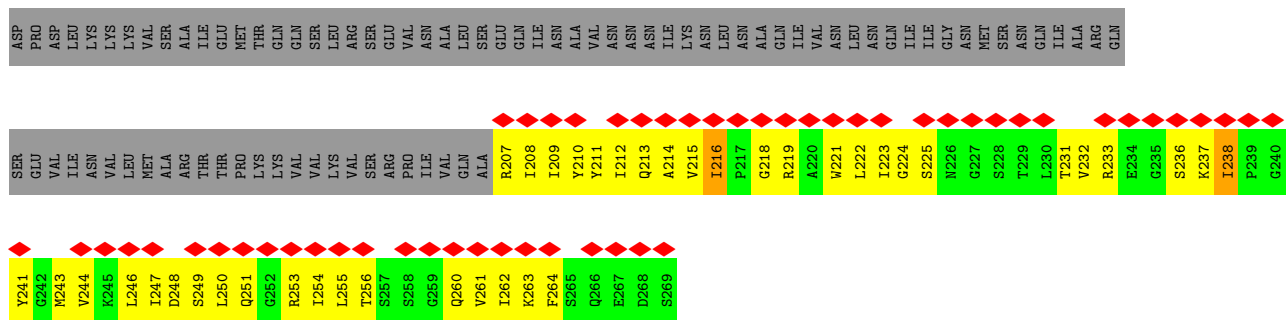
• Molecule 5: IcmG (DotF)



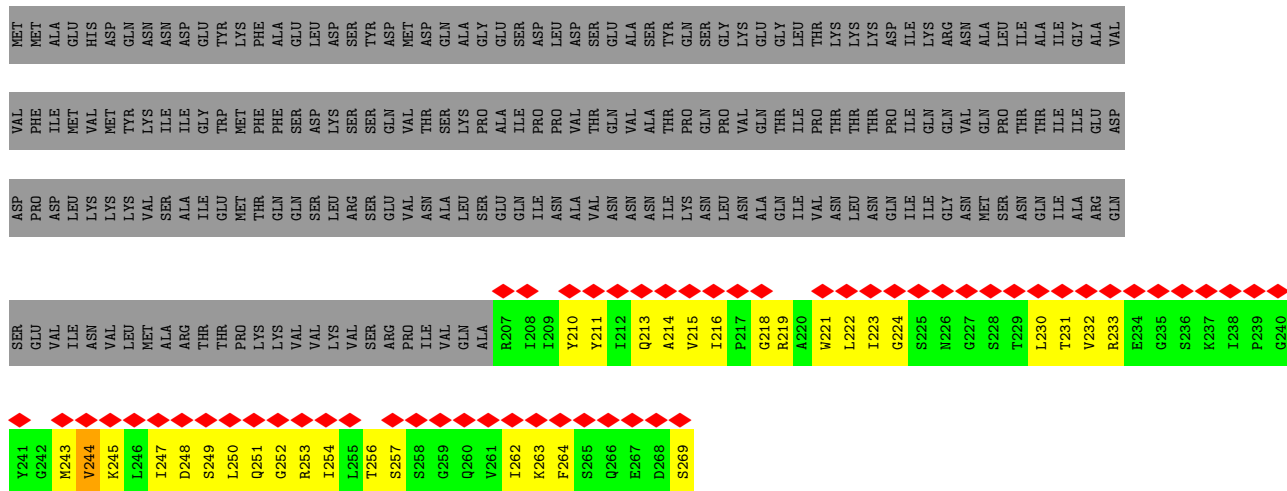
• Molecule 5: IcmG (DotF)



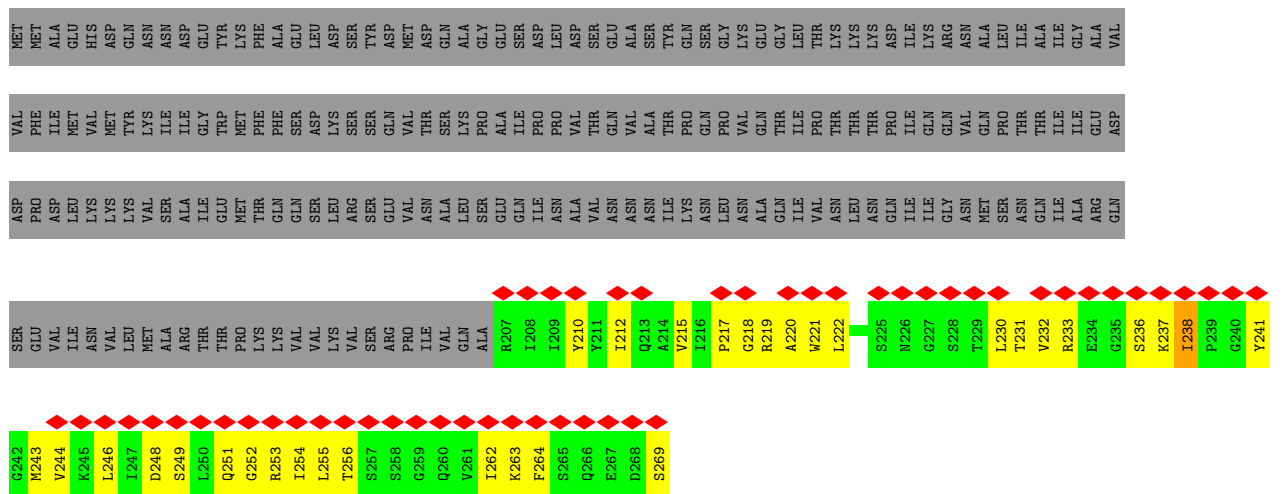




- Molecule 5: IcmG (DotF)



- Molecule 5: IcmG (DotF)



- Molecule 5: IcmG (DotF)



[illegible]

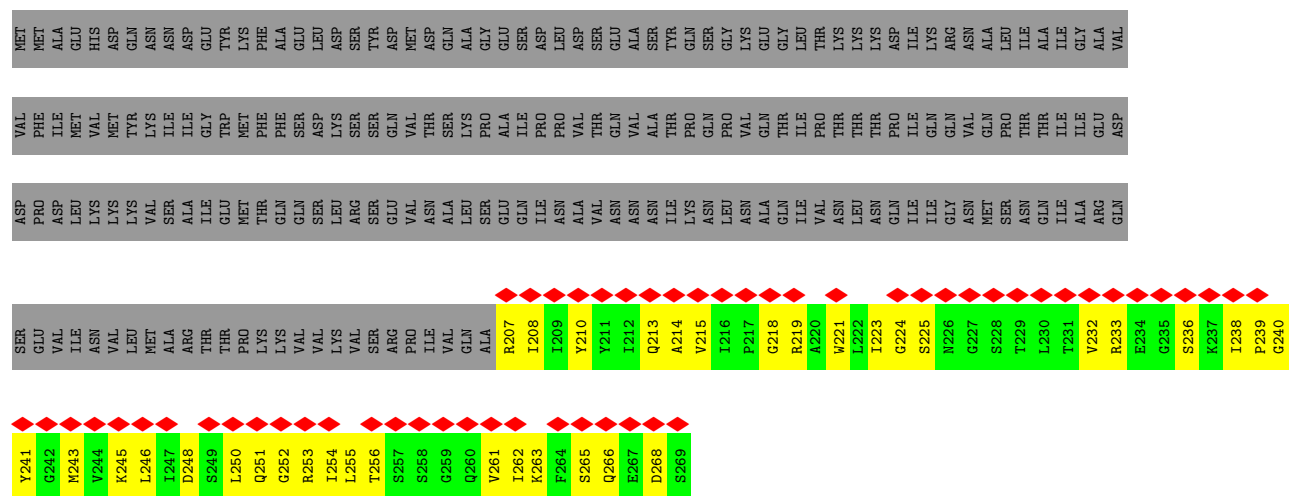
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SER	GLU	VAL	ILE	ASN	LEU	MET	ALA	ARG	THR	PRO	PRO	LYS	LYS	VAL	VAL	LYS	VAL	SER	ARG	ARG	PRO	ILE	ILE	GLN	GLN	ALA																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																							</
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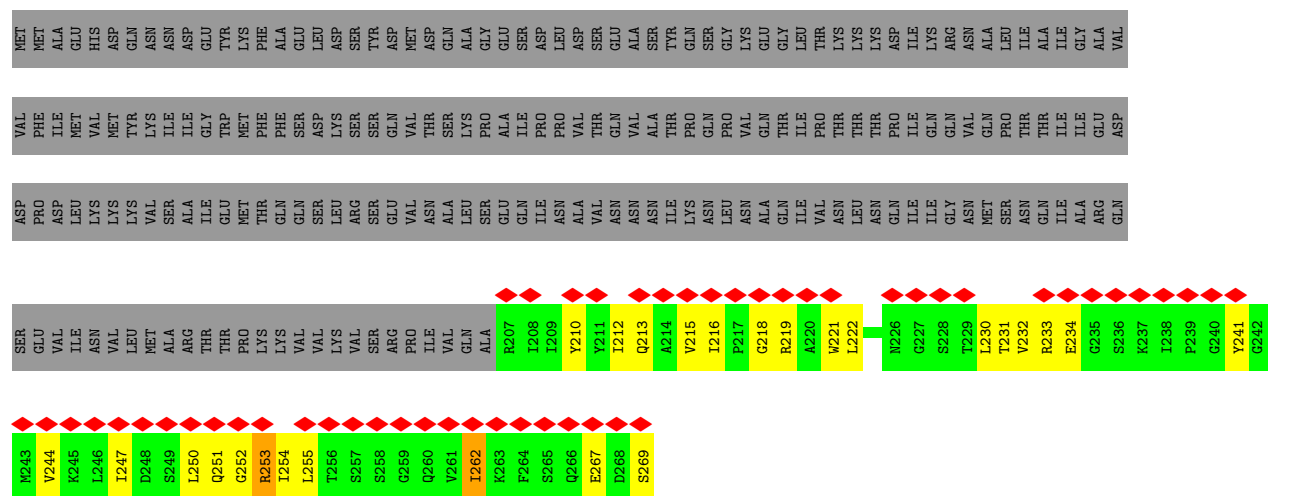




- Molecule 5: IcmG (DotF)



- Molecule 5: IcmG (DotF)

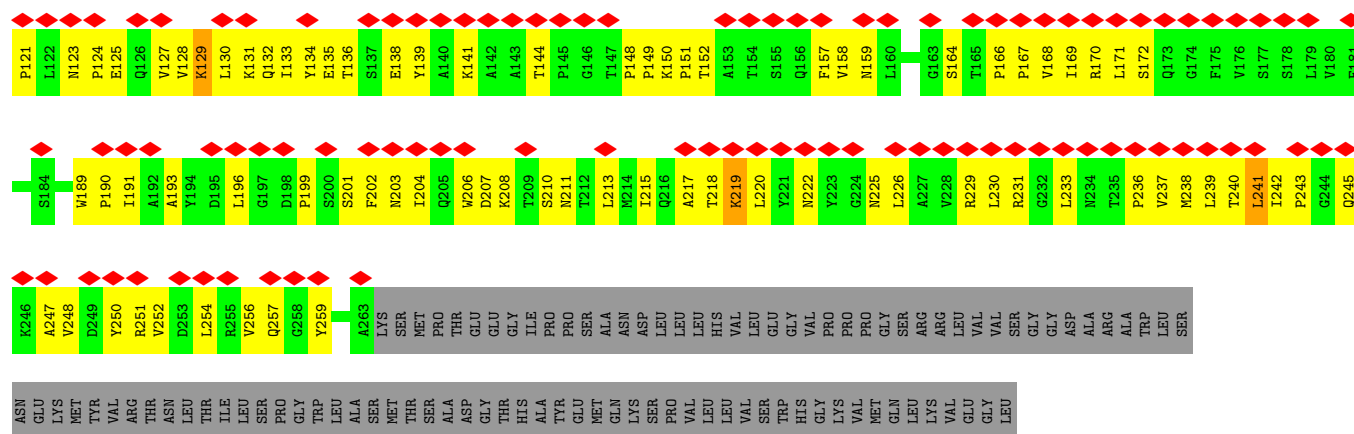


- Molecule 5: IcmG (DotF)

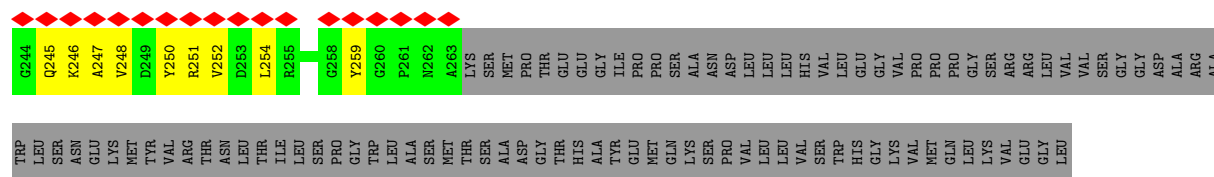
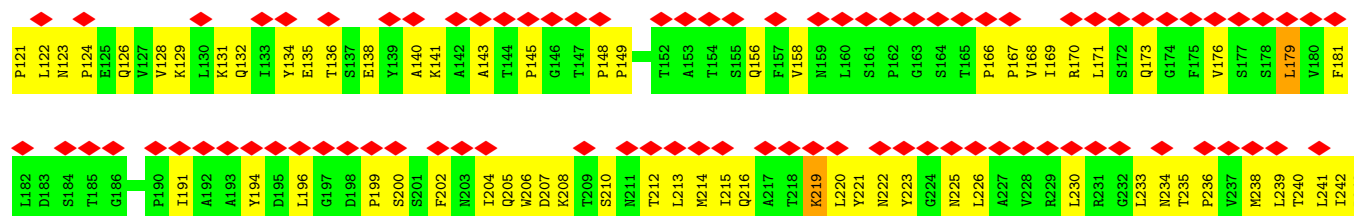
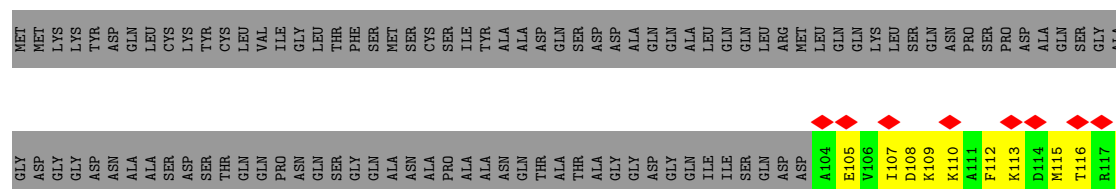


Y241	SER	GLU	VAL	ILE	ASN	VAL	LEU	MET	ALA	ARG	THR	PRO	PRO	LYS	LYS	VAL	VAL	LYS	VAL	SER	ARG	PRO	ILE	VAL	GLN	ALA	R207	R208	T209	Y210	Y211	T212	Q213	A214	V215	T216	P217	Q218	R219	A220	W221	T222	T223	G224	S225	N226	Q227	S228	T229	L230	T231	V232	R233	E234	G235	S236	T237	T238	P239	Q240
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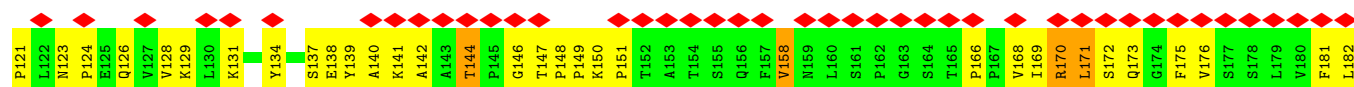
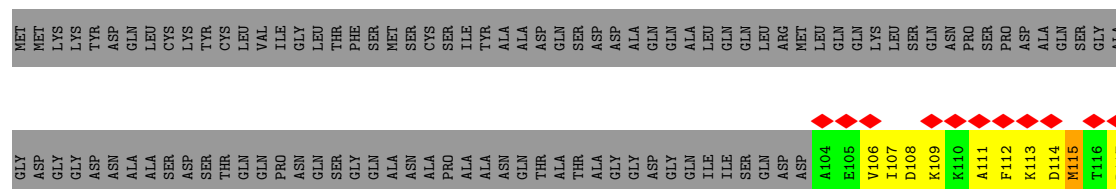
GLY	ASP	GLY	GLY	ASP	ASN	ALA	ALA	SER	ASP	SER	THR	GLN	GLN	PRO	ASN	GLN	SER	GLY	GLN	ALA	ALA	ASN	ASN	GLN	THR	ALA	ALA	THR	ALA	GLY	GLY	ASP	GLN	GLN	ILE	ILE	SER	GLN	ASP	ASP	ASP	A104	E105	V106	I107	D108	K109	K110	A111	F112	K113	D114	M115	T116	R117	N118	L119	V120
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• Molecule 6: IcmK (DotH)

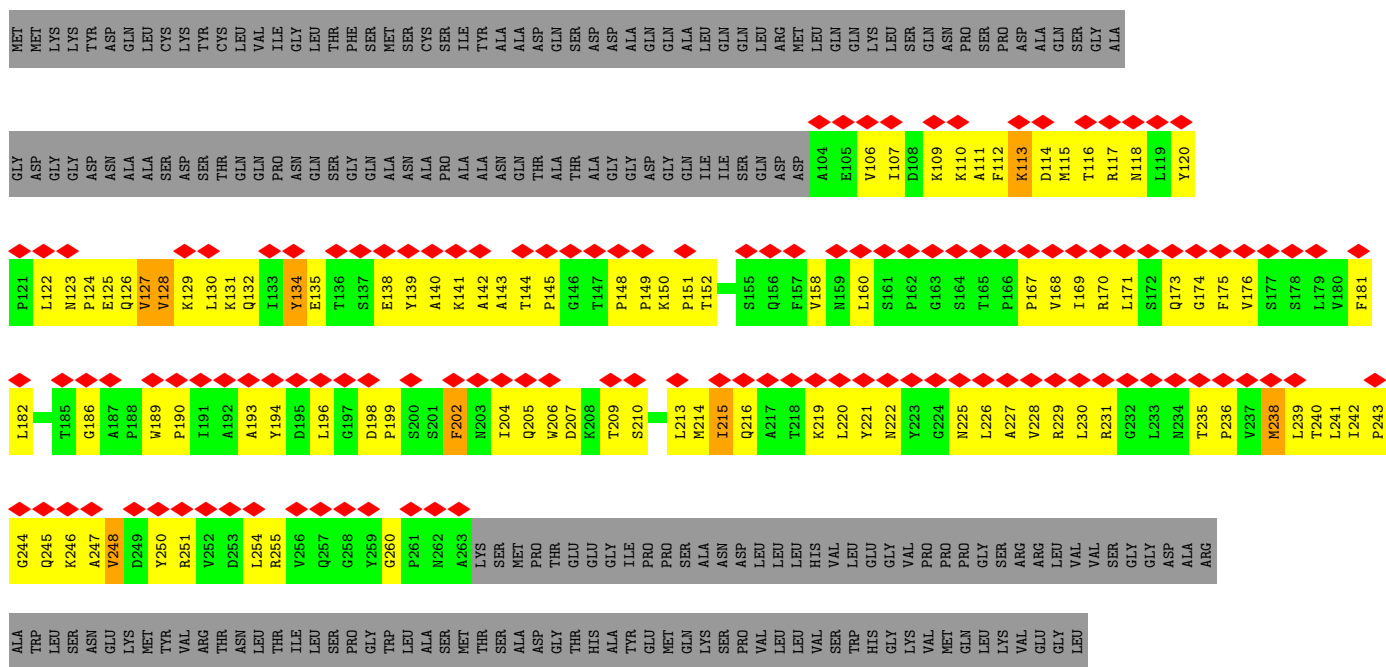


• Molecule 6: IcmK (DotH)

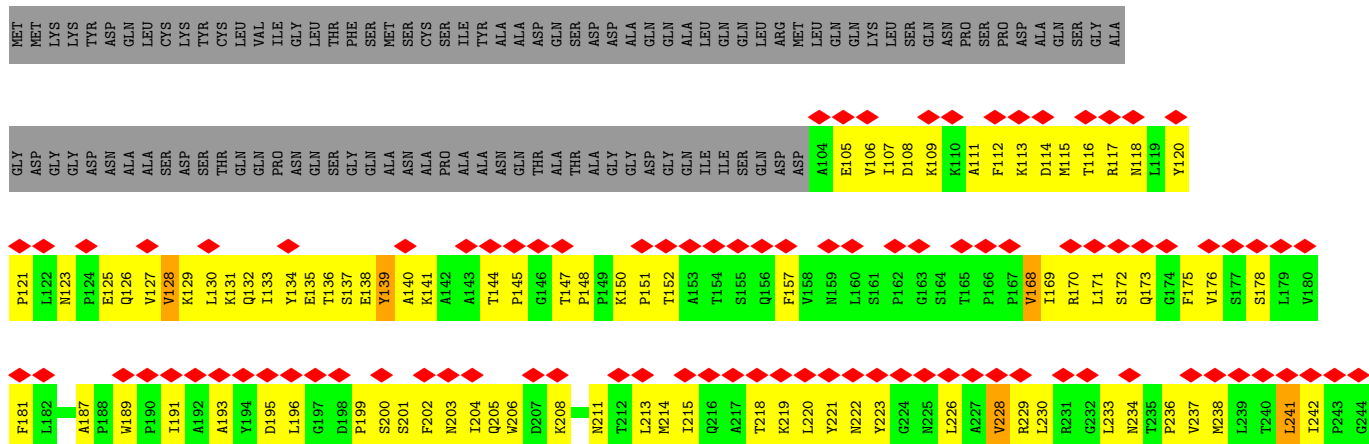




- Molecule 6: IcmK (DotH)



- Molecule 6: IcmK (DotH)





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• Molecule 6: IcmK (DotH)



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• Molecule 6: IcmK (DotH)



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PRO PRO  
SER SER  
ALA ALA  
ASP ASP  
THR THR  
HIS HIS  
ALA ALA  
TYR TYR  
GLU GLU  
MET MET  
SER SER  
GLN GLN  
LYS LYS  
SER SER  
PRO PRO  
VAL VAL  
LEU LEU  
HIS HIS  
VAL VAL  
TRP TRP  
HIS HIS  
GLY GLY  
VAL VAL  
PRO PRO  
PRO PRO  
GLY GLY  
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ARG ARG  
VAL VAL  
LEU LEU  
VAL VAL  
SER SER  
GLY GLY  
GLY GLY  
ASP ASP  
ALA ALA  
ARG ARG

TRP LEU  
SER ASN  
GLU GLU  
LYS LYS  
MET MET  
TYR TYR  
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GLN GLN  
ASP ASP  
THR THR  
HIS HIS  
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TYR TYR  
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HIS HIS  
TRP TRP  
HIS HIS  
GLY GLY  
VAL VAL  
PRO PRO  
PRO PRO  
GLY GLY  
SER SER  
VAL VAL  
VAL VAL  
GLY GLY  
GLY GLY  
LEU LEU

• Molecule 6: IcmK (DotH)



LEU	ASP	L241	F181	P121	MET
ALA	ALA	L242	L182	L122	ASP
ARG	ARG	P243	D183	N123	LYS
ALA	ALA	G244	S184	P124	GLY
LEU	TRP	Q245	T185	E125	TYR
SER	LEU	K246	G186	Q126	ASN
ASN	ASN	A247	A187	V127	ALA
GLU	GLU	V248	P188	V128	ALA
LYS	MET	D249	W189	K129	SER
TYR	TYR	Y250	P190	L130	ASP
VAL	VAL	R251	I191	K131	SER
ARG	ARG	V252	A192	Q132	THR
THR	THR	D253	A193	I133	GLN
LEU	ASN	L254	A194	Y134	LEU
LEU	LEU	R255	Y194	E135	GLY
THR	THR	R256	D195	T136	ASN
ILE	ILE	V256	L196	S137	SER
LEU	LEU	Q257	G197	E138	PHE
SER	SER	G258	D198	Y139	GLN
PRO	PRO	Y259	P199	A140	ALA
GLY	GLY	G260	S200	K141	ALA
TRP	TRP	P261	S201	A142	PRO
LEU	LEU	N262	F202	A143	ALA
ALA	ALA	A263	N203	T144	ILE
SER	SER	LYS	T204	P145	TYR
MET	MET	THR	Q205	G146	ALA
THR	THR	SER	W206	T147	GLN
ALA	ALA	PRO	D207	P148	ASP
ASP	ASP	THR	K208	P149	ASP
GLY	GLY	GLU	T209	K150	GLY
THR	THR	GLU	S210	P151	ASP
HIS	HIS	GLY	N210	GLY	GLN
ALA	ALA	ILE	N211	T152	GLN
TYR	TYR	GLU	T212	A153	ALA
GLU	GLU	PRO	L213	T154	LEU
MET	MET	SER	M214	S155	GLN
GLN	GLN	ALA	I215	Q156	LEU
LYS	LYS	ASN	T215	F157	GLN
SER	SER	ASP	Q216	V158	ARG
PRO	PRO	VAL	A217	N159	MET
VAL	VAL	LEU	T218	L160	LEU
LEU	LEU	LEU	K219	S161	GLN
VAL	VAL	VAL	L220	P162	LYS
TRP	TRP	LEU	Y221	G163	SER
GLU	GLU	GLU	N222	S164	GLN
GLY	GLY	GLY	Y223	T165	ASN
VAL	VAL	VAL	G224	P166	PRO
PRO	PRO	PRO	N225	P167	SER
PRO	PRO	PRO	L226	V168	PRO
GLY	GLY	GLY	A227	I169	ASP
LEU	LEU	SER	V228	R170	ALA
VAL	VAL	ARG	R229	L171	SER
VAL	VAL	LEU	L230	S172	GLY
VAL	VAL	VAL	R231	Q173	GLY
SER	SER	SER	G232	G174	ALA
GLY	GLY	GLY	L233	F175	
GLY	GLY	GLY	M234	V176	
			T235	S177	
			P236	S178	
			V237	L179	
			M238	V180	
			L239		
			T240		

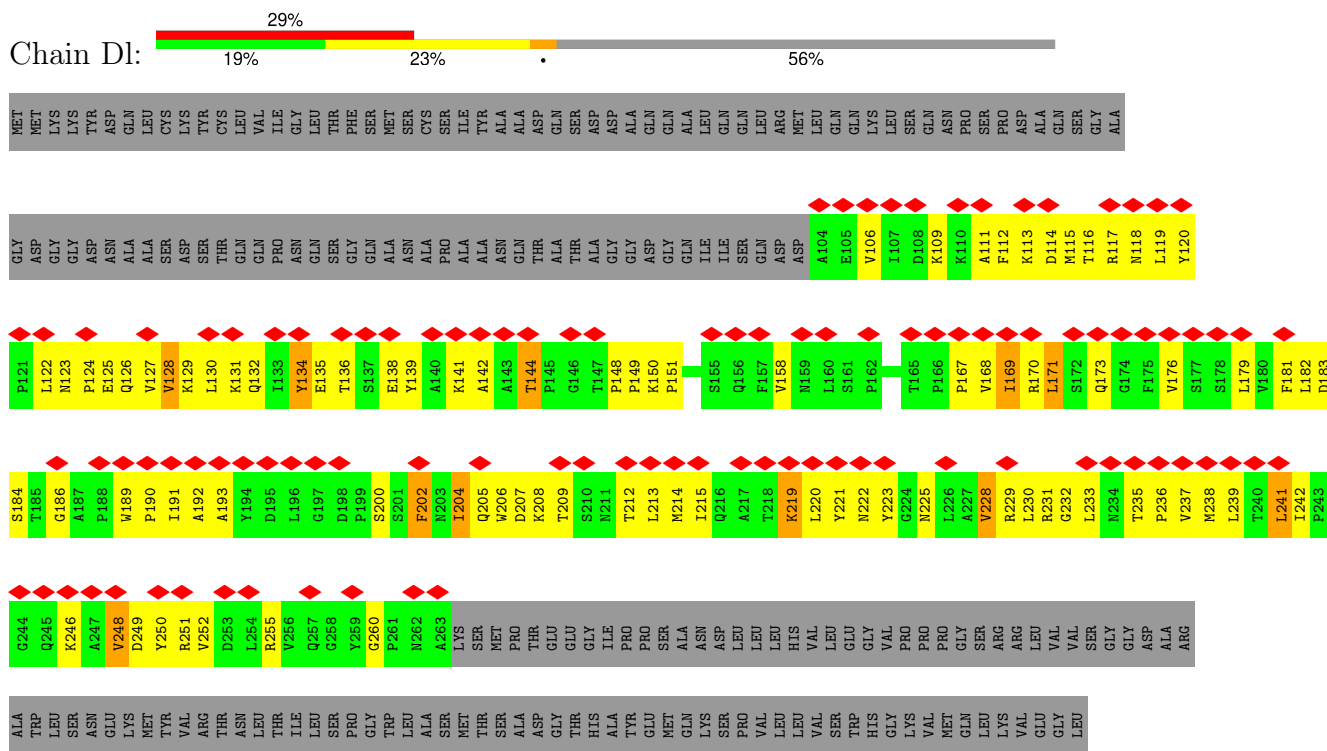
• Molecule 6: IcmK (DotH)



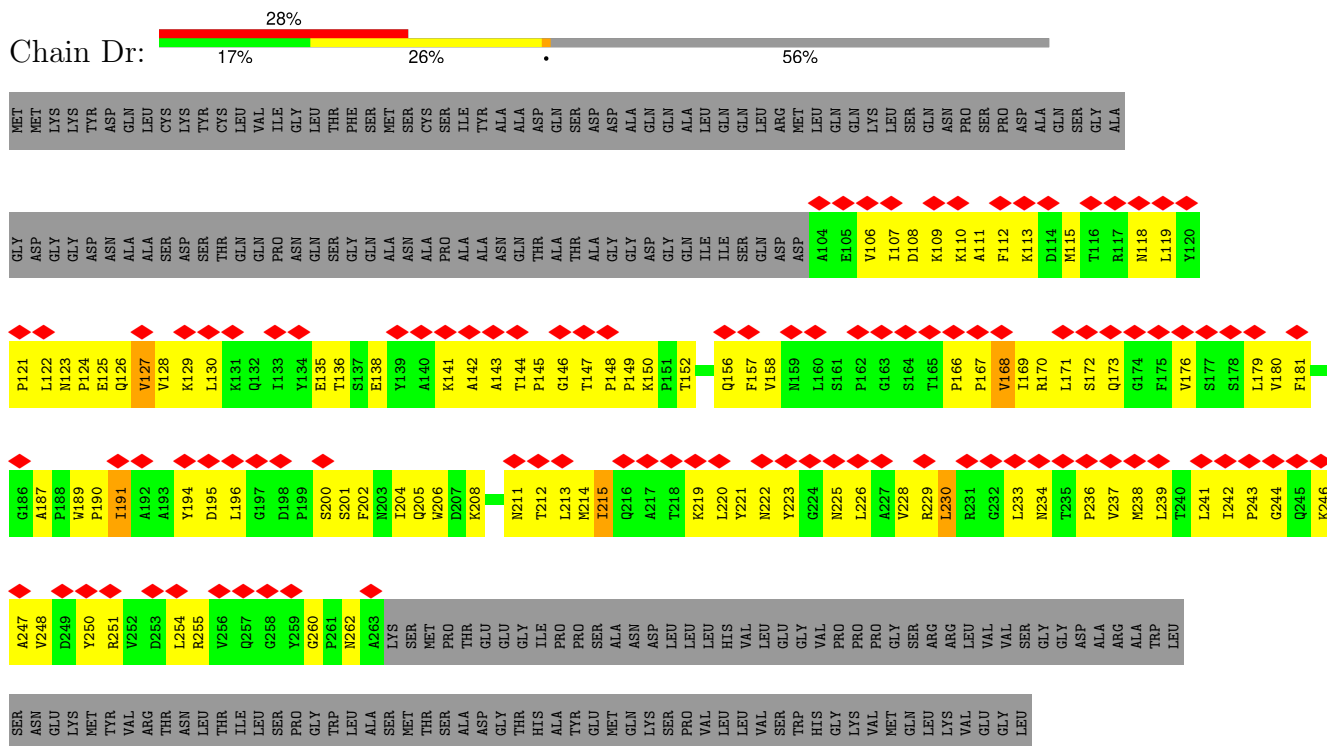
ALA	TRP	LEU	SER	ASN	GLU	LYS	MET	TYR	VAL	ARG	THR	ASN	LEU	THR	ILE	THR	THR	GLY	TRP	LEU	ALA	SER	LYS	GLY	GLU	GLY	VAL	GLY	GLY	VAL	PRO	PRO	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GLY	GL
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• Molecule 6: IcmK (DotH)





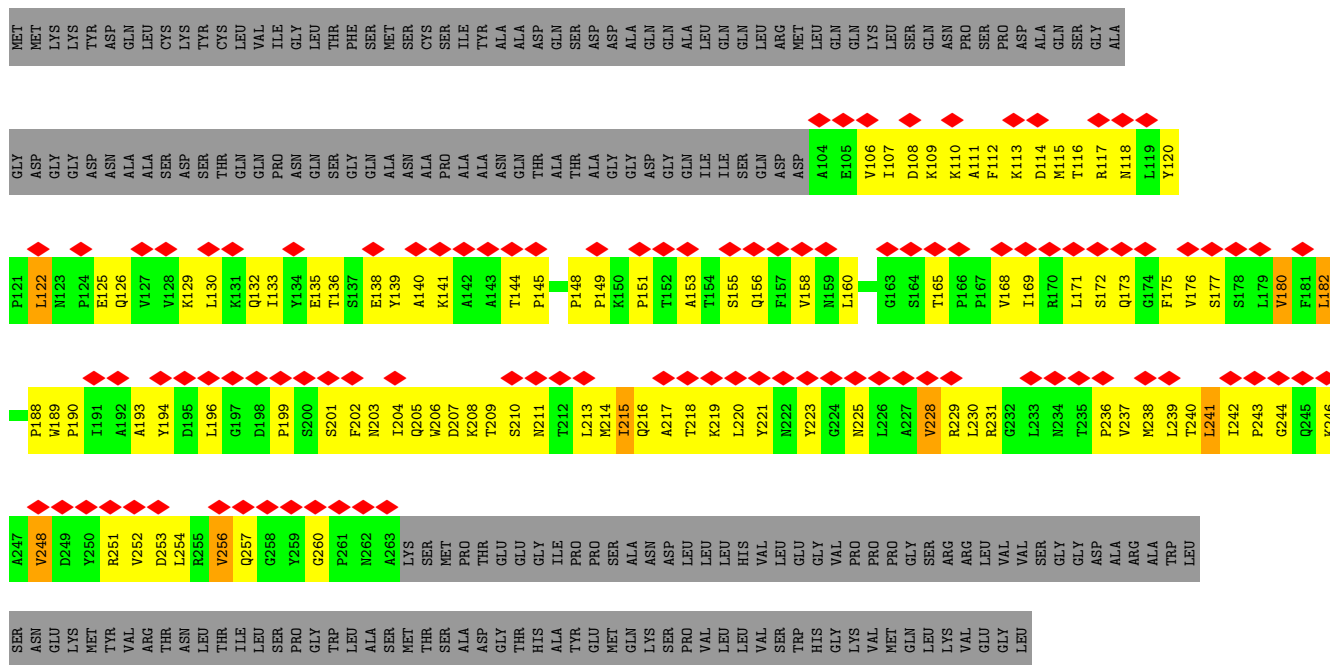
- Molecule 6: IcmK (DotH)



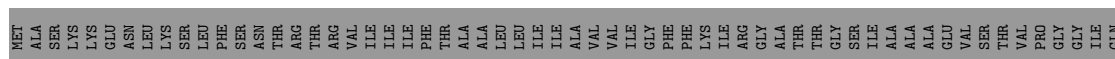
- Molecule 6: IcmK (DotH)



- Molecule 6: IcmK (DotH)



- Molecule 7: IcmE (DotG)



- Molecule 7: IcmE (DotG)



MET	ALA	SER	LYS	GLU	ASN	LEU	SER	PHE	THR	ARG	THR	ARG	VAL	ILE	ILE	PHE	THR	ALA	ALA	LEU	LEU	ILE	ILE	ALA	VAL	VAL	ILE	ILE	GLY	PHE	PHE	LYS	ILE	ALA	ARG	GLY	ALA	THR	THR	GLY	SER	SER	ILE	ALA	ALA	ALA	GLU	GLU	VAL	VAL	THR	THR	PRO	PRO	GLY	GLY	ILE	ILE	TLE	TLE
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- Molecule 7: IcmE (DotG)



- Molecule 7: IcmE (DotG)

Chain Ei:



95%

[illegible]

- Molecule 7: IcmE (DotG)

Chain Ej:  94%

MET	ALA	SER	LYS	GLU	ASN	LEU	LYS	SER	PHE	SER	THR	THR	ARG	ARG	VAL	ILE	ILE	PHE	THR	ALA	ALA	LEU	LEU	ILE	ILE	ALA	VAL	VAL	ILE	ILE	GLY	PHE	PHE	LYS	ILE	ILE	ARG	GLY	ARG	ALA	ALA	THR	THR	GLY	SER	SER	ILE	ALA	ALA	GLU	GLU	VAL	VAL	THR	THR	VAL	PRO	GLY	GLY	ILE	ILE	GLN
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- Molecule 7: IcmE (DotG)

Chain El:  97%

MET	ALA	SER	LYS	GLU	ASN	LEU	SER	THR	THR	ARG	VAL	ILE	ILE	PHE	THR	ALA	ALA	LEU	ILE	ILE	PHE	PHE	LYS	ILE	ARG	GLY	ALA	THR	THR	GLY	SER	ILE	ALA	ALA	ALA	GLU	VAL	SER	THR	VAL	PRO	GLY	GLY	ILE	GLN
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- Molecule 7: IcmE (DotG)







- Molecule 7: IcmE (DotG)



MET	ALA	SER	LYS	GLU	ASN	LEU	SER	PHE	THR	ARG	THR	ARG	VAL	ILE	ILE	PHE	ALA	ALA	VAL	ILE	ILE	GLY	PHE	PHE	LYS	ILE	ILE	ARG	GLY	ALA	ALA	THR	THR	GLY	SER	SER	ILE	ILE	ALA	ALA	GLU	GLU	VAL	VAL	THR	THR	VAL	PRO	GLY	GLY	ILE	ILE	TYR
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MET	ALA	SER	LYS	GLU	ASN	LEU	SER	LEU	THR	THR	ARG	VAL	ILE	ILE	PHE	THR	ALA	ALA	LEU	LEU	ILE	ILE	PHE	PHE	PHE	LYS	ILE	ILE	ARG	GLY	ALA	THR	THR	GLY	SER	ILE	ILE	ALA	ALA	ALA	GLU	VAL	SER	THR	VAL	VAL	PRO	GLY	GLY	ILE	ILE	GLN
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- Molecule 7: IcmE (DotG)

[illegible]



- Molecule 7: IcmE (DotG)

[illegible]

- Molecule 7: IcmE (DotG)

Chain Fh:  96%





- Molecule 7: IcmE (DotG)



- Molecule 7: IcmE (DotG)

Chain Fk:  96%

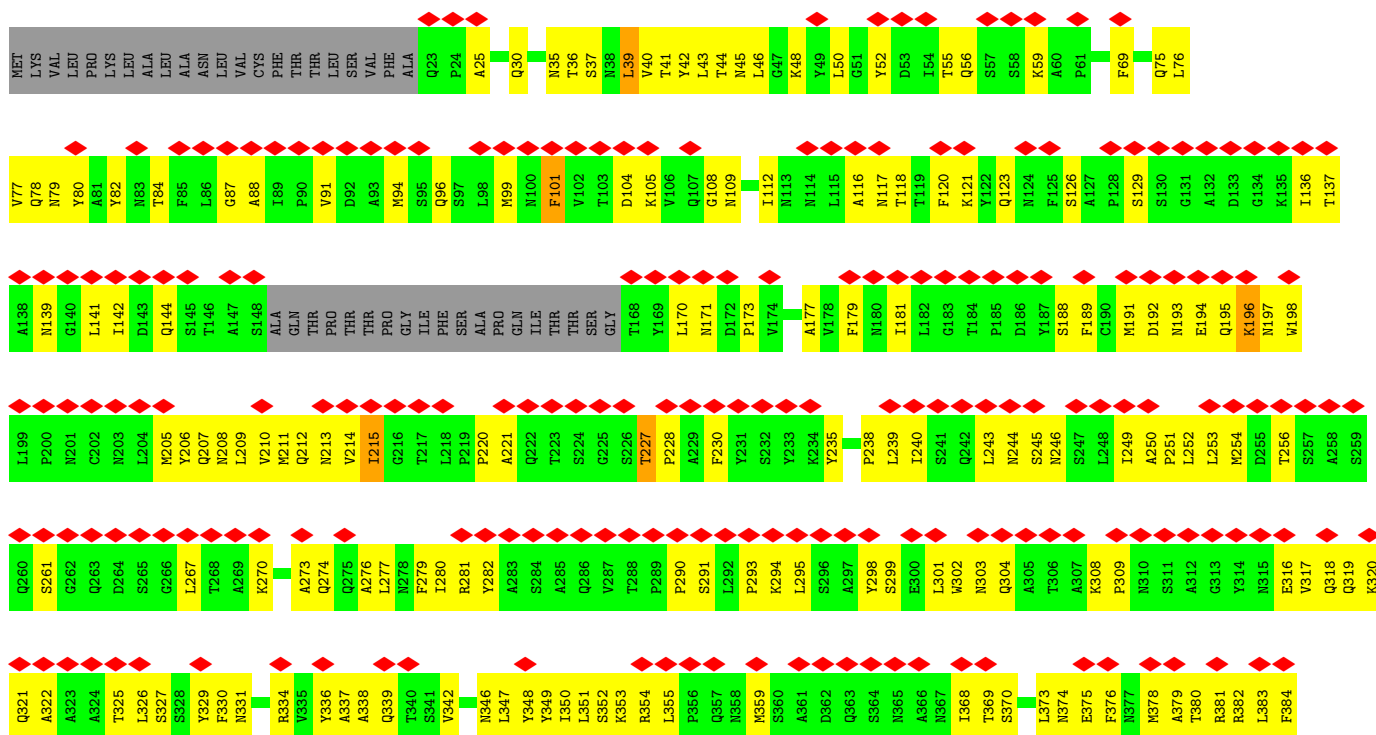
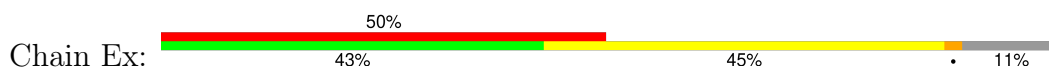
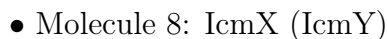


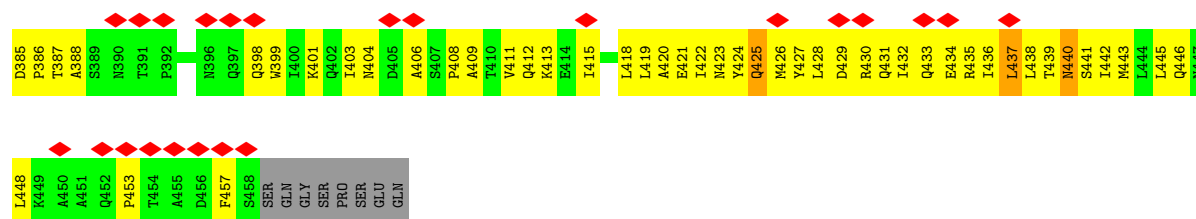


- Molecule 8: IcmX (IcmY)

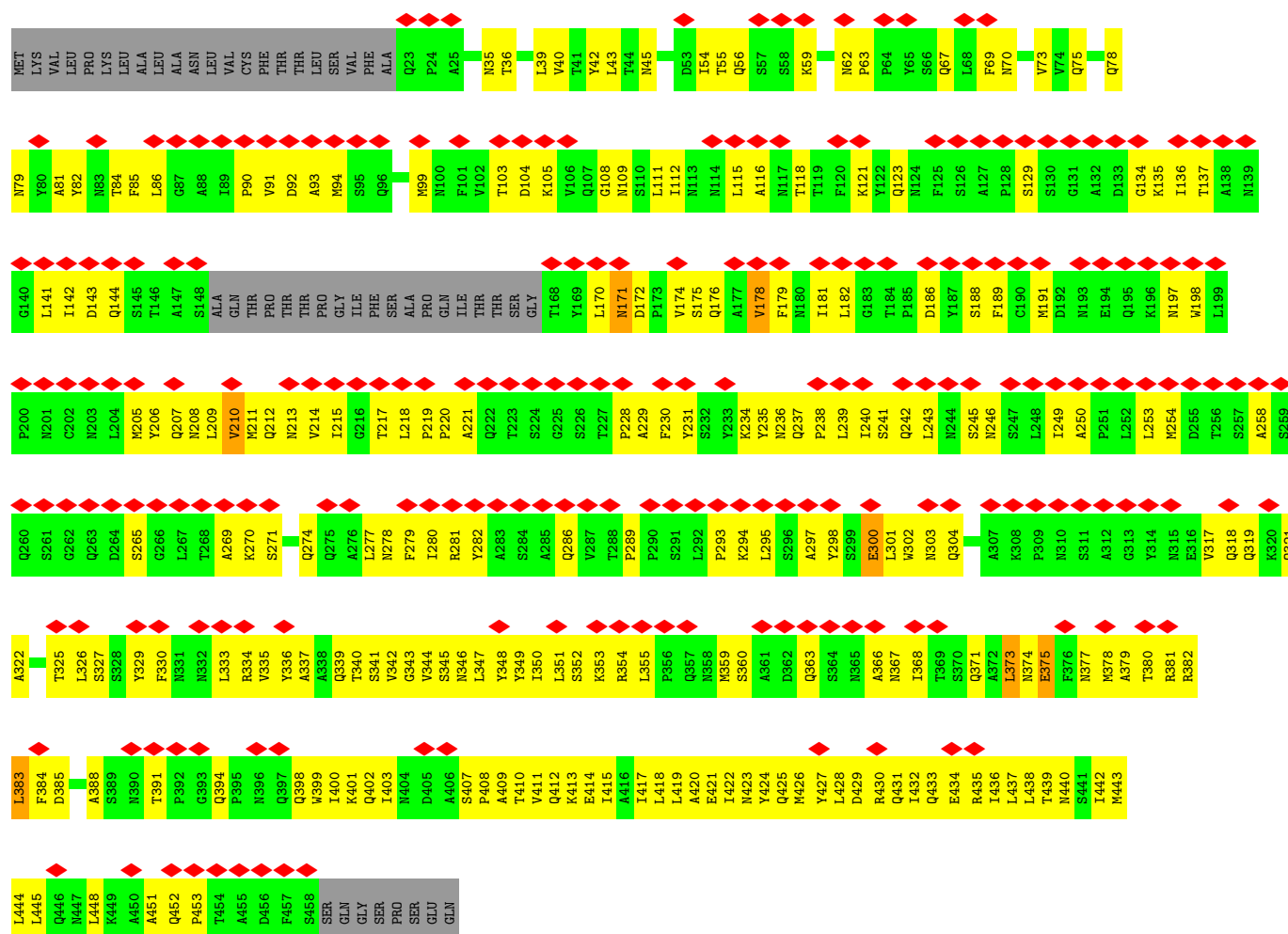
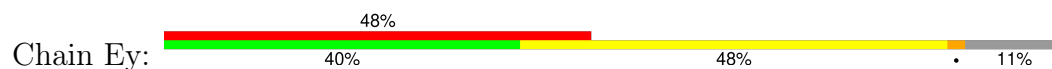
Government	Percentage
Current government	52%
Previous government	43%
Previous government	44%
Previous government	11%



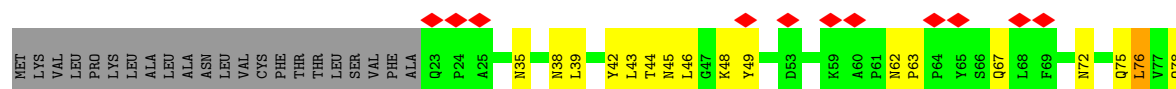


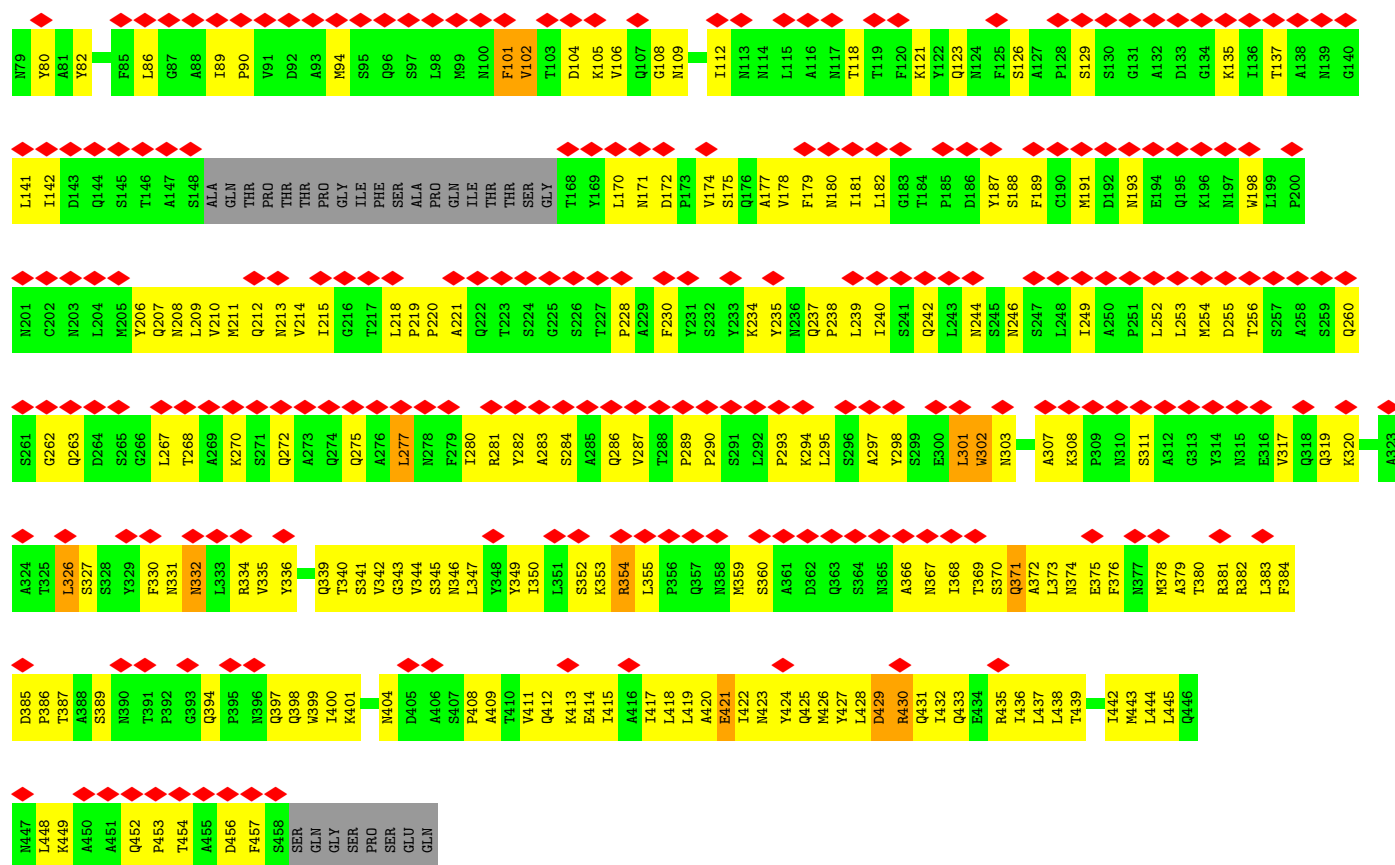


• Molecule 8: IcmX (IcmY)



• Molecule 8: IcmX (IcmY)

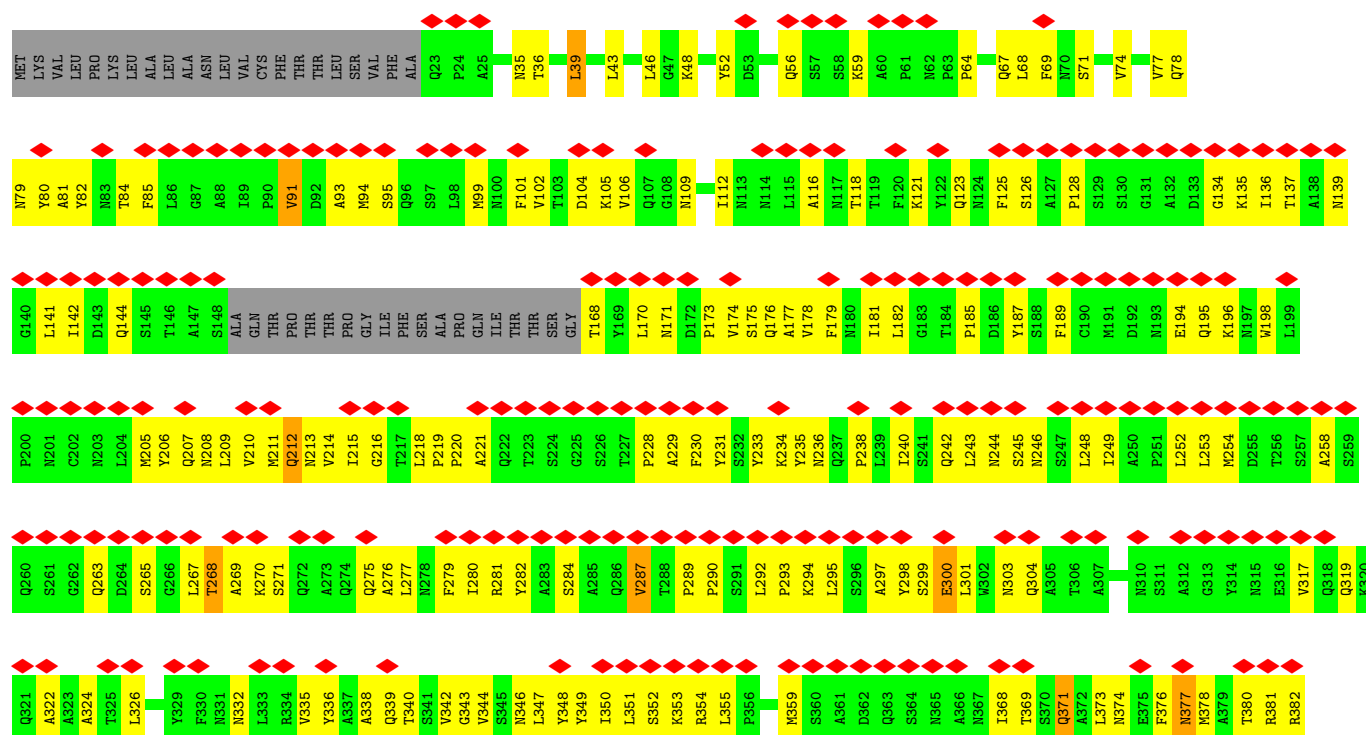


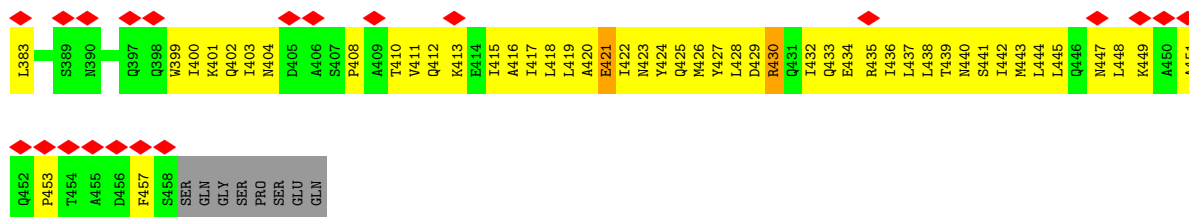


### • Molecule 8: IcmX (IcmY)

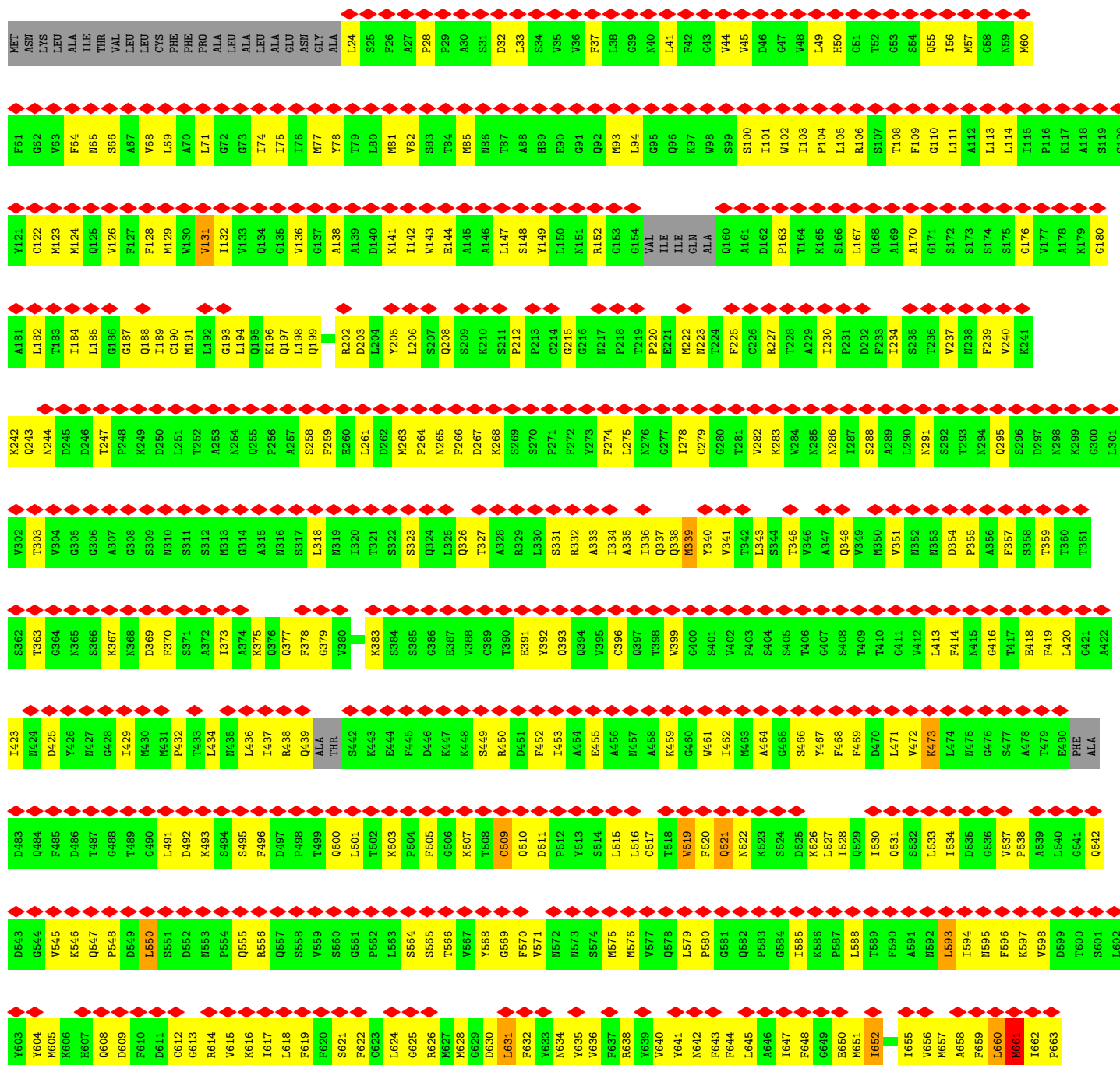
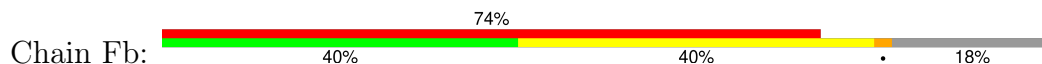


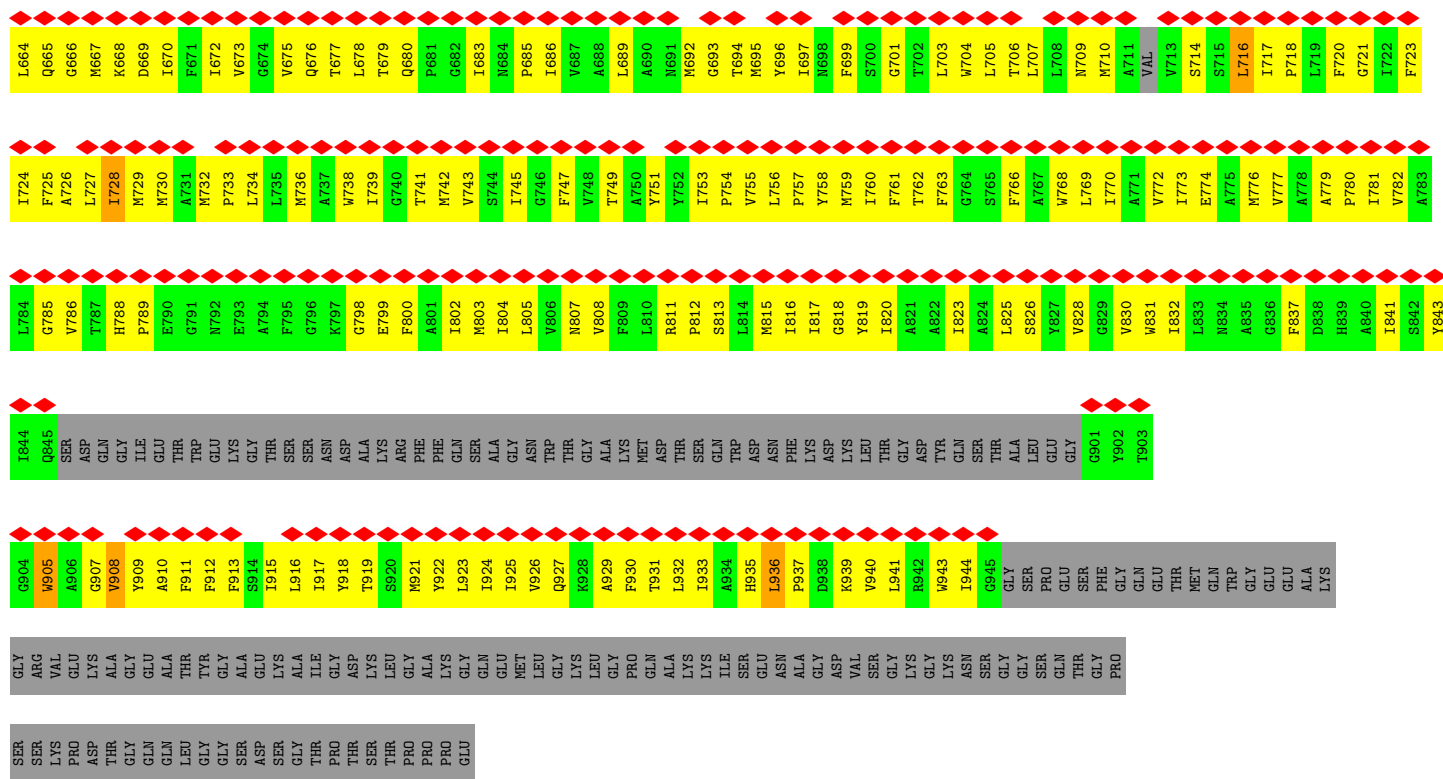
Chain Fa:



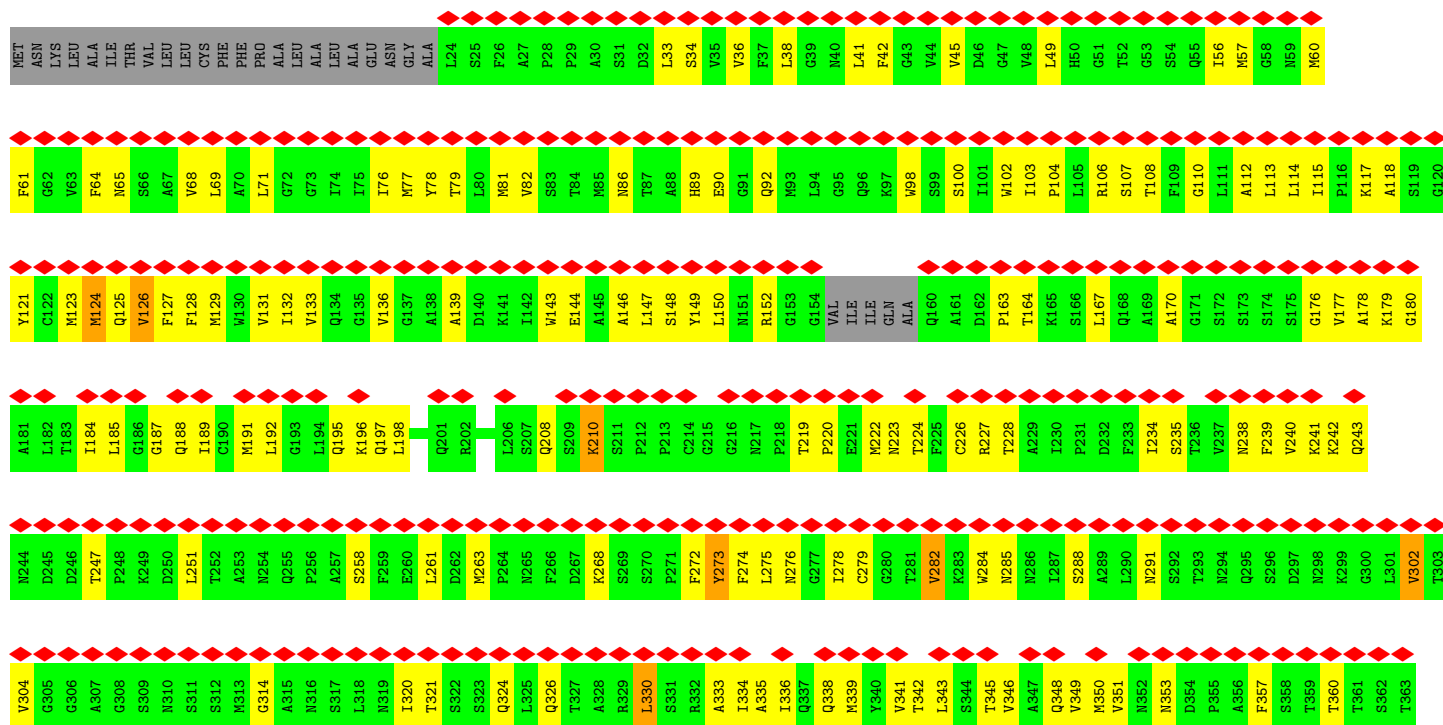
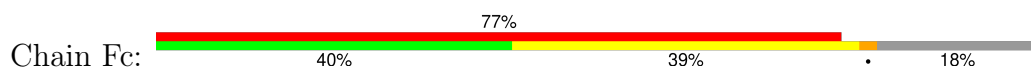


## • Molecule 9: DotA





## • Molecule 9: DotA





ASP	GLN	GLY	ILE	GLU	THR	TRP	GLU	LYS	GLY	THR	SER	SER	ASN	ASP	ALA	LYS	ARG	PHE	PHE	GLN	SER	ALA	GLY	ASN	TRP	THR	THR	GLY	ALA	LYS	MET	ASP	THR	SER	GLN	TRP	ASP	ASN	PHE	LYS	GLN	ASP	LYS	LEU	THR	GLY	ASP	TYR	GLN	SER	THR	ALA	LEU	GLY	GLY	G901	G902	G903	G904	W905	A906
T787	H788	P789	E790	G791	N792	E793	A794	F795	G796	K797	G798	E799	F800	A801	I802	M803	M804	L805	V806	M807	V808	F809	L810	R811	S812	S813	L814	M815	L816	I817	G818	Y819	I820	F821	T822	F823	G824	L825	S826	Y827	L828	W829	L830	W831	V832	W833	L834	A835	G836	F837	D838	H839	A840	I841	S842	Y843	L844	G845	SER		
M667	K668	D669	F670	D671	C672	G673	G674	G675	D676	T677	L678	T679	D680	F681	I682	I683	M684	P685	I686	V687	A688	L689	M690	M691	M692	N693	T694	M695	Y696	I697	L698	F699	S700	G701	T702	L703	W704	L705	T706	L707	L708	W709	W710	A711	VAL	W713	S714	S715	L716	L717	F718	L719	F720	G721	I722	F723	L724	F725	A726		
L727	I728	M729	K730	A731	M732	P733	L734	L735	M736	A737	W738	I739	G740	T741	W742	W743	S744	I745	G746	F747	W748	T749	A750	Y751	Y752	I753	P754	Y755	L756	P757	R758	M759	I760	F761	T762	F763	G764	S765	F766	A767	W768	L769	M770	I771	A772	W773	E774	A775	M776	W777	A778	A779	P780	I781	Y782	A783	L784	G785	Y786		
H607	Q608	D609	F610	D611	C612	G613	R614	V615	K616	L617	L618	F619	F620	S621	F622	C623	L624	G625	R626	M627	M628	G629	D630	L631	F632	Y633	N634	G635	V636	F637	R638	Y639	V640	Y641	N642	F643	F644	L645	A646	L647	F648	G649	E650	M651	I652	N653	S654	L655	V656	M657	A658	L660	R661	I662	P663	L664	G665	G666			
Q547	P548	D549	L550	S551	D552	N553	P554	Q555	R556	Q557	S558	V559	S560	G561	P562	L563	S564	S565	T566	V567	Y568	G569	F570	V571	N572	N573	S574	M575	M576	V577	Q578	L579	Y641	N642	F643	F644	L645	A646	L647	F648	G649	E650	M651	I652	N653	S654	L655	V656	M657	A658	L660	R661	I662	P663	L664	G665	G666				
Q547	P548	D549	L550	S551	D552	N553	P554	Q555	R556	Q557	S558	V559	S560	G561	P562	L563	S564	S565	T566	V567	Y568	G569	F570	V571	N572	N573	S574	M575	M576	V577	Q578	L579	Y641	N642	F643	F644	L645	A646	L647	F648	G649	E650	M651	I652	N653	S654	L655	V656	M657	A658	L660	R661	I662	P663	L664	G665	G666				
Q547	P548	D549	L550	S551	D552	N553	P554	Q555	R556	Q557	S558	V559	S560	G561	P562	L563	S564	S565	T566	V567	Y568	G569	F570	V571	N572	N573	S574	M575	M576	V577	Q578	L579	Y641	N642	F643	F644	L645	A646	L647	F648	G649	E650	M651	I652	N653	S654	L655	V656	M657	A658	L660	R661	I662	P663	L664	G665	G666				
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Q547	P548	D549	L550	S551	D552	N553	P554	Q555	R556	Q557	S558	V559	S560	G561	P562	L563	S564	S565	T566	V567	Y568	G569	F570	V571	N572	N573	S574	M575	M576	V577	Q578	L579	Y641	N642	F643	F644	L645	A646	L647	F648	G649	E650	M651	I652	N653	S654	L655	V656	M657	A658	L660	R661	I662	P663	L664	G665	G666				
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Q547	P548	D549	L550	S551	D552	N553	P554	Q555	R556	Q557	S558	V559	S560	G561	P562	L563	S564	S565	T566	V567	Y568	G569	F570	V571	N572	N573	S574	M575	M576	V577	Q578	L579	Y641	N642	F643	F644	L645	A646	L647	F648	G649	E650	M651	I652	N653	S654	L655	V656	M657	A658	L660	R661	I662	P663	L664	G665	G666				
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Q547	P548	D549	L550	S551	D552	N553	P554	Q555	R556	Q557	S558	V559	S560	G561	P562	L563	S564	S565	T566	V567	Y568	G569	F570	V571	N572	N573	S574	M575	M576	V577	Q578	L579	Y641	N642	F643	F644	L645	A646	L647	F648	G649	E650	M651	I652	N653	S654	L655	V656	M657	A658	L660	R661	I662	P663	L664	G665	G666				
Q547	P548	D549	L550	S551	D552	N553	P554	Q555	R556	Q557	S558	V559	S560	G561	P562	L563	S564	S565	T566	V567	Y568	G569	F570	V571	N572	N573	S574	M575	M576	V577	Q578	L579	Y641	N642	F643	F644	L645	A646	L647	F648	G649	E650	M651	I652	N653	S654	L655	V656	M657	A658	L660	R661	I662	P663	L664	G665	G666				
Q547	P548	D549	L550	S551	D552	N553	P554	Q555	R556	Q557	S558	V559	S560	G561	P562	L563	S564	S565	T566	V567	Y568	G569	F570	V571	N572	N573	S574	M575	M576	V577	Q578	L579	Y641	N642	F643	F644	L645	A646	L647	F648	G649	E650	M651	I652	N653	S654	L655	V656	M657	A658	L660	R661	I662	P663	L664	G665	G666				
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Q547	P548	D549	L550	S551	D552	N553	P554>P																																																						







GLY	ARG	GLY	ARG	L301	T361	A422	ALA	Q542	L664	L784	G904	GLY	ARG	GLY	ARG	L301	T361	A422	ALA	Q542	L664	L784	G904	GLY	ARG	GLY	ARG
SER	SER	VAL	VAL	V302	S362	I423	D483	D543	Q665	G785	W905	SER	SER	VAL	VAL	V302	S362	I423	D483	D543	Q665	G785	W905	SER	SER	VAL	VAL
PRO	PRO	ASP	ASP	T303	T363	Q424	Q484	G544	G666	V786	A906	GLN	GLN	ASP	ASP	T303	T363	Q424	Q484	G544	G666	V786	A906	GLN	GLN	ASP	ASP
THR	THR	THR	THR	V304	G364	Y426	F485	V545	M667	T787	W908	GLY	GLY	GLN	GLN	V304	G364	Y426	F485	V545	M667	T787	W908	GLY	GLY	GLN	GLN
GLY	GLY	ALA	ALA	G305	N365	G428	D486	Q547	D669	H788	Y909	GLU	GLU	ILE	ILE	G305	N365	G428	D486	Q547	D669	H788	Y909	GLU	GLU	ILE	ILE
GLU	GLU	GLU	GLU	G306	S366	T427	G488	P548	L670	W789	A910	ALA	ALA	THR	THR	G306	S366	T427	G488	P548	L670	W789	A910	ALA	ALA	THR	THR
THR	THR	THR	THR	A307	K367	G428	T489	D549	F671	G791	F911	THR	THR	TRP	TRP	A307	K367	G428	T489	D549	F671	G791	F911	THR	THR	TRP	TRP
GLU	GLU	GLY	GLY	G308	N368	I429	G490	L550	F610	T792	W912	GLY	GLY	GLU	GLU	G308	N368	I429	G490	L550	F610	T792	W912	GLY	GLY	GLU	GLU
GLY	GLY	GLY	GLY	S309	D369	M430	G491	S551	D611	G793	F913	GLY	GLY	GLU	GLU	S309	D369	M430	G491	S551	D611	G793	F913	GLY	GLY	GLU	GLU
SER	SER	ALA	ALA	N310	F370	M431	D492	D552	G612	A794	S914	GLY	GLY	THR	THR	N310	F370	M431	D492	D552	G612	A794	S914	GLY	GLY	THR	THR
ASP	ASP	GLU	GLU	S311	S371	P432	K493	N553	G674	F795	W916	THR	THR	THR	THR	S311	S371	P432	K493	N553	G674	F795	W916	THR	THR	THR	THR
GLY	GLY	ALA	ALA	S312	A372	L434	S494	P554	R614	G796	I915	SER	SER	SER	SER	S312	A372	L434	S494	P554	R614	G796	I915	SER	SER	SER	SER
THR	THR	GLY	GLY	M313	I373	L436	S495	Q554	V615	T797	W916	GLY	GLY	GLN	GLN	M313	I373	L436	S495	Q554	V615	T797	W916	GLY	GLY	GLN	GLN
GLY	GLY	PRO	PRO	G314	A374	I437	F496	R556	K616	W738	Y918	ASP	ASP	ALA	ALA	G314	A374	I437	F496	R556	K616	W738	Y918	ASP	ASP	ALA	ALA
THR	THR	THR	THR	A315	Q377	R439	D497	Q557	I617	G740	F919	GLY	GLY	GLY	GLY	A315	Q377	R439	D497	Q557	I617	G740	F919	GLY	GLY	GLY	GLY
PRO	PRO	PRO	PRO	N316	F378	Q439	T499	S558	L618	T741	A921	LEU	LEU	ARG	ARG	N316	F378	Q439	T499	S558	L618	T741	A921	LEU	LEU	ARG	ARG
PRO	PRO	PRO	PRO	S317	G379	Q439	T499	S559	Q680	G742	W922	GLY	GLY	PHE	PHE	S317	G379	Q439	T499	S559	Q680	G742	W922	GLY	GLY	PHE	PHE
GLY	GLY	GLY	GLY	N319	V380	Q500	Q500	S560	F620	H743	S923	GLY	GLY	GLN	GLN	N319	V380	Q500	Q500	S560	F620	H743	S923	GLY	GLY	GLN	GLN
GLY	GLY	GLY	GLY	I320	P381	THR	L501	G561	S821	S744	I924	GLY	GLY	ALA	ALA	I320	P381	THR	L501	G561	S821	S744	I924	GLY	GLY	ALA	ALA
GLY	GLY	GLY	GLY	T321	Y382	S442	T502	P562	C623	L745	W925	GLY	GLY	GLY	GLY	T321	Y382	S442	T502	P562	C623	L745	W925	GLY	GLY	GLY	GLY
GLY	GLY	GLY	GLY	S322	K383	K443	K503	L563	L624	T746	A926	GLY	GLY	TRP	TRP	S322	K383	K443	K503	L563	L624	T746	A926	GLY	GLY	TRP	TRP
GLY	GLY	GLY	GLY	S323	S384	E444	P504	S564	G625	G746	W926	GLY	GLY	ASP	ASP	S323	S384	E444	P504	S564	G625	G746	W926	GLY	GLY	ASP	ASP
GLY	GLY	GLY	GLY	Q324	S385	F445	F505	T566	R626	F747	K928	GLY	GLY	THR	THR	Q324	S385	F445	F505	T566	R626	F747	K928	GLY	GLY	THR	THR
GLY	GLY	GLY	GLY	L325	G386	D446	G506	S567	M627	W748	A929	GLY	GLY	ALA	ALA	L325	G386	D446	G506	S567	M627	W748	A929	GLY	GLY	ALA	ALA
GLY	GLY	GLY	GLY	Q326	E387	K447	K507	Y568	M628	T749	F930	GLY	GLY	ALA	ALA	Q326	E387	K447	K507	Y568	M628	T749	F930	GLY	GLY	ALA	ALA
GLY	GLY	GLY	GLY	T327	V388	K448	T508	G569	L631	A751	W931	GLY	GLY	MET	MET	T327	V388	K448	T508	G569	L631	A751	W931	GLY	GLY	MET	MET
GLY	GLY	GLY	GLY	A328	C389	S449	C509	F570	F632	Y752	I932	GLY	GLY	ASP	ASP	A328	C389	S449	C509	F570	F632	Y752	I932	GLY	GLY	ASP	ASP
GLY	GLY	GLY	GLY	R329	T390	R450	Q510	N571	Y633	T753	W933	GLY	GLY	THR	THR	R329	T390	R450	Q510	N571	Y633	T753	W933	GLY	GLY	THR	THR
GLY	GLY	GLY	GLY	S331	E391	D451	D511	N572	M634	Y754	A934	GLY	GLY	GLN	GLN	S331	E391	D451	D511	N572	M634	Y754	A934	GLY	GLY	GLN	GLN
GLY	GLY	GLY	GLY	R332	Y392	F452	P512	N573	Y635	T755	W935	GLY	GLY	TRP	TRP	R332	Y392	F452	P512	N573	Y635	T755	W935	GLY	GLY	TRP	TRP
GLY	GLY	GLY	GLY	A333	Q393	A454	Y513	S574	Y636	W756	A936	GLY	GLY	ASP	ASP	A333	Q393	A454	Y513	S574	Y636	W756	A936	GLY	GLY	ASP	ASP
GLY	GLY	GLY	GLY	I334	V395	E455	S514	L515	R638	T757	W937	GLY	GLY	PHE	PHE	I334	V395	E455	S514	L515	R638	T757	W937	GLY	GLY	PHE	PHE
GLY	GLY	GLY	GLY	A335	C396	A456	L516	M576	Y639	W758	A938	GLY	GLY	GLY	GLY	A335	C396	A456	L516	M576	Y639	W758	A938	GLY	GLY	GLY	GLY
GLY	GLY	GLY	GLY	I336	Q397	N457	C517	V577	W640	T759	W939	GLY	GLY	ASP	ASP	I336	Q397	N457	C517	V577	W640	T759	W939	GLY	GLY	ASP	ASP
GLY	GLY	GLY	GLY	Q337	T398	A458	T518	Q578	Y641	L760	I941	GLY	GLY	LEU	LEU	Q337	T398	A458	T518	Q578	Y641	L760	I941	GLY	GLY	LEU	LEU
GLY	GLY	GLY	GLY	Q338	W399	K459	W519	L579	M642	F761	W942	GLY	GLY	THR	THR	Q338	W399	K459	W519	L579	M642	F761	W942	GLY	GLY	THR	THR
GLY	GLY	GLY	GLY	M339	G400	G460	F520	P580	N642	W762	A942	GLY	GLY	GLY	GLY	M339	G400	G460	F520	P580	N642	W762	A942	GLY	GLY	GLY	GLY
GLY	GLY	GLY	GLY	Y340	S401	V461	Q521	N582	F643	T763	W943	GLY	GLY	ASP	ASP	Y340	S401	V461	Q521	N582	F643	T763	W943	GLY	GLY	ASP	ASP
GLY	GLY	GLY	GLY	V341	V402	I462	N522	Q581	F644	W764	A944	GLY	GLY	GLN	GLN	V341	V402	I462	N522	Q581	F644	W764	A944	GLY	GLY	GLN	GLN
GLY	GLY	GLY	GLY	T342	P403	M463	K523	P583	L645	G764	W945	GLY	GLY	SER	SER	T342	P403	M463	K523	P583	L645	G764	W945	GLY	GLY	SER	SER
GLY	GLY	GLY	GLY	L343	S404	A464	S524	G584	I647	F766	A946	GLY	GLY	THR	THR	L343	S404	A464	S524	G584	I647	F766	A946	GLY	GLY	THR	THR
GLY	GLY	GLY	GLY	S344	S405	G465	D525	I585	F648	W767	W947	GLY	GLY	ALA	ALA	S344	S405	G465	D525	I585	F648	W767	W947	GLY	GLY	ALA	ALA
GLY	GLY	GLY	GLY	T345	S406	G466	K526	K586	M651	A767	I948	GLY	GLY	LEU	LEU	T345	S406	G466	K526	K586	M651	A767	I948	GLY	GLY	LEU	LEU
GLY	GLY	GLY	GLY	V346	T406	S466	L527	P587	I652	W768	W949	GLY	GLY	GLU	GLU	V346	T406	S466	L527	P587	I652	W768	W949	GLY	GLY	GLU	GLU
GLY	GLY	GLY	GLY	Q348	S408	Y467	I528	T588	N653	T770	A950	GLY	GLY	GLY	GLY	Q348	S408	Y467	I528	T588	N653	T770	A950	GLY	GLY	GLY	GLY
GLY	GLY	GLY	GLY	V349	T409	F469	Q529	T589	S654	A771	W951	GLY	GLY	THR	THR	V349	T409	F469	Q529	T589	S654	A771	W951	GLY	GLY	THR	THR
GLY	GLY	GLY	GLY	M350	T410	D470	I530	F590	I655	W769	W952	GLY	GLY	GLY	GLY	M350	T410	D470	I530	F590	I655	W769	W952	GLY	GLY	GLY	GLY
GLY	GLY	GLY	GLY	V351	G411	L471	Q531	N591	N655	T771	A953	GLY	GLY	GLY	GLY	V351	G411	L471	Q531	N591	N655	T771	A953	GLY	GLY	GLY	GLY
GLY	GLY	GLY	GLY	N352	V412	V472	S532	N592	V656	W772	W954	GLY	GLY	GLY	GLY	N352	V412	V472	S532	N592	V656	W772	W954	GLY	GLY	GLY	GLY
GLY	GLY	GLY	GLY	N353	L413	K473	L533	L593	M657	I773	A955	GLY	GLY	GLY	GLY	N353	L413	K473	L533	L593	M657	I773	A955	GLY	GLY	GLY	GLY
GLY	GLY	GLY	GLY	D354	F414	L474	I534	I594	A658	E774	W956	GLY	GLY	GLY	GLY	D354	F414	L474	I534	I594	A658	E774	W956	GLY	GLY	GLY	GLY
GLY	GLY	GLY	GLY	P355	N415	N475	D535	N595	F659	W775	A957	GLY	GLY	GLY	GLY	P355	N415	N475	D535	N595	F659	W775	A957	GLY	GLY	GLY	GLY
GLY	GLY	GLY	GLY	A356	G416	G476	G536	F596	L660	T776	W958	GLY	GLY	GLY	GLY	A356	G416	G476	G536	F596	L660	T776	W958	GLY	GLY	GLY	GLY
GLY	GLY	GLY	GLY	F357	T417	S477	V537	K597	M661	A778	W959	GLY	GLY	GLY	GLY	F357	T417	S477	V537	K597	M661	A778	W959	GLY	GLY	GLY	GLY
GLY	GLY	GLY	GLY	S358	E418	A478	P538	V598	G661	T779	A960	GLY	GLY	GLY	GLY	S358	E418	A478	P538	V598	G661	T779	A960	GLY	GLY	GLY	GLY
GLY	GLY	GLY	GLY	T359	F419	T479	A539	D599	M662	A779	W960	GLY	GLY	GLY	GLY	T359	F419	T479	A539	D599	M662	A779	W960	GLY	GLY	GLY	GLY
GLY	GLY	GLY	GLY	T360	L420	E480	L540	T600	I662	W780	A961	GLY	GLY	GLY	GLY	T360	L420	E480	L540	T600	I662	W78					

## 4 Experimental information

Property	Value	Source
EM reconstruction method	SINGLE PARTICLE	Depositor
Imposed symmetry	POINT, Not provided	
Number of particles used	37503	Depositor
Resolution determination method	FSC 0.143 CUT-OFF	Depositor
CTF correction method	PHASE FLIPPING ONLY	Depositor
Microscope	FEI TECNAI 12	Depositor
Voltage (kV)	300	Depositor
Electron dose ( $e^-/\text{\AA}^2$ )	73	Depositor
Minimum defocus (nm)	800	Depositor
Maximum defocus (nm)	2000	Depositor
Magnification	Not provided	
Image detector	GATAN K3 (6k x 4k)	Depositor
Maximum map value	0.434	Depositor
Minimum map value	-0.283	Depositor
Average map value	0.002	Depositor
Map value standard deviation	0.022	Depositor
Recommended contour level	0.16	Depositor
Map size ( $\text{\AA}$ )	640.8192, 640.8192, 640.8192	wwPDB
Map dimensions	512, 512, 512	wwPDB
Map angles ( $^\circ$ )	90.0, 90.0, 90.0	wwPDB
Pixel spacing ( $\text{\AA}$ )	1.2516, 1.2516, 1.2516	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	Aa	0.11	0/118	0.44	0/158
1	Ag	0.18	0/118	0.64	0/158
1	Am	0.14	0/118	0.40	0/158
1	As	0.15	0/118	0.38	0/158
1	Ay	0.12	0/118	0.44	0/158
1	Be	0.14	0/118	0.43	0/158
1	Bk	0.13	0/118	0.52	0/158
1	Bq	0.11	0/118	0.42	0/158
1	Bw	0.16	0/118	0.45	0/158
1	Cc	0.13	0/118	0.40	0/158
1	Ci	0.13	0/118	0.47	0/158
1	Co	0.16	0/118	0.69	0/158
1	Cu	0.09	0/118	0.37	0/158
1	Da	0.16	0/118	0.63	0/158
1	Dg	0.16	0/118	0.68	0/158
1	Dm	0.13	0/118	0.49	0/158
1	Ds	0.13	0/118	0.50	0/158
1	Dy	0.13	0/118	0.50	0/158
2	Ab	0.34	0/60	0.53	0/76
2	Ah	0.27	0/60	0.60	0/76
2	An	0.59	0/60	1.04	0/76
2	At	0.20	0/60	0.56	0/76
2	Az	0.23	0/60	0.56	0/76
2	Bf	0.21	0/60	0.44	0/76
2	Bl	0.31	0/60	0.81	0/76
2	Br	0.40	0/60	0.89	0/76
2	Bx	0.21	0/60	0.71	0/76
2	Cd	0.28	0/60	0.61	0/76
2	Cj	0.28	0/60	0.62	0/76
2	Cp	0.24	0/60	0.45	0/76
2	Cv	0.19	0/60	0.43	0/76
2	Db	0.24	0/60	0.63	0/76
2	Dh	0.45	0/60	0.69	0/76
2	Dn	0.30	0/60	0.61	0/76

Mol	Chain	Bond lengths		Bond angles	
		RMSZ	# Z  >5	RMSZ	# Z  >5
2	Dt	0.28	0/60	0.67	0/76
2	Dz	0.21	0/60	0.40	0/76
3	Ac	0.10	0/126	0.29	0/167
3	Ai	0.17	0/126	0.44	0/167
3	Ao	0.18	0/126	0.43	0/167
3	Au	0.13	0/126	0.36	0/167
3	Ba	0.13	0/126	0.37	0/167
3	Bg	0.13	0/126	0.33	0/167
3	Bm	0.17	0/126	0.49	0/167
3	Bs	0.15	0/126	0.34	0/167
3	By	0.12	0/126	0.36	0/167
3	Ce	0.13	0/126	0.39	0/167
3	Ck	0.13	0/126	0.38	0/167
3	Cq	0.15	0/126	0.47	0/167
3	Cw	0.13	0/126	0.39	0/167
3	Dc	0.13	0/126	0.40	0/167
3	Di	0.14	0/126	0.48	0/167
3	Do	0.12	0/126	0.33	0/167
3	Du	0.17	0/126	0.52	0/167
3	Ea	0.11	0/126	0.50	0/167
4	Ad	0.12	0/37	0.22	0/47
4	Aj	0.11	0/37	0.26	0/47
4	Ap	0.14	0/37	0.34	0/47
4	Av	0.09	0/37	0.19	0/47
4	Bb	0.20	0/37	0.39	0/47
4	Bh	0.11	0/37	0.23	0/47
4	Bn	0.11	0/37	0.25	0/47
4	Bt	0.12	0/37	0.31	0/47
4	Bz	0.14	0/37	0.27	0/47
4	Cf	0.17	0/37	0.40	0/47
4	Cl	0.09	0/37	0.20	0/47
4	Cr	0.08	0/37	0.18	0/47
4	Cx	0.12	0/37	0.22	0/47
4	Dd	0.20	0/37	0.45	0/47
4	Dj	0.09	0/37	0.22	0/47
4	Dp	0.08	0/37	0.23	0/47
4	Dv	0.12	0/37	0.18	0/47
4	Eb	0.13	0/37	0.30	0/47
5	Ae	0.20	0/491	0.58	0/660
5	Ak	0.21	0/491	0.59	0/660
5	Aq	0.17	0/491	0.46	0/660
5	Aw	0.20	0/491	0.54	0/660
5	Bc	0.23	0/491	0.61	0/660

Mol	Chain	Bond lengths		Bond angles	
		RMSZ	# Z  >5	RMSZ	# Z  >5
5	Bi	0.18	0/491	0.57	0/660
5	Bo	0.24	0/491	0.63	0/660
5	Bu	0.18	0/491	0.52	0/660
5	Ca	0.18	0/491	0.53	0/660
5	Cg	0.20	0/491	0.45	0/660
5	Cm	0.23	0/491	0.67	0/660
5	Cs	0.25	0/491	0.68	0/660
5	Cy	0.19	0/491	0.48	0/660
5	De	0.22	0/491	0.62	0/660
5	Dk	0.21	0/491	0.60	0/660
5	Dq	0.19	0/491	0.61	0/660
5	Dw	0.18	0/491	0.47	0/660
5	Ec	0.16	0/491	0.46	0/660
6	Af	0.27	0/1269	0.74	7/1734 (0.4%)
6	Al	0.26	0/1269	0.63	0/1734
6	Ar	0.26	0/1269	0.67	0/1734
6	Ax	0.27	0/1269	0.64	3/1734 (0.2%)
6	Bd	0.23	0/1269	0.59	0/1734
6	Bj	0.24	0/1269	0.65	0/1734
6	Bp	0.27	0/1269	0.76	0/1734
6	Bv	0.28	0/1269	0.68	1/1734 (0.1%)
6	Cb	0.23	0/1269	0.64	1/1734 (0.1%)
6	Ch	0.25	0/1269	0.65	2/1734 (0.1%)
6	Cn	0.28	0/1269	0.71	4/1734 (0.2%)
6	Ct	0.25	0/1269	0.64	2/1734 (0.1%)
6	Cz	0.26	0/1269	0.68	1/1734 (0.1%)
6	Df	0.30	0/1269	0.75	4/1734 (0.2%)
6	Dl	0.27	0/1269	0.69	0/1734
6	Dr	0.30	0/1269	0.80	2/1734 (0.1%)
6	Dx	0.26	0/1269	0.82	6/1734 (0.3%)
6	Ed	0.26	0/1269	0.63	0/1734
7	Ee	0.25	0/278	0.71	0/377
7	Ef	0.25	0/475	0.83	0/642
7	Eg	0.32	0/278	0.98	0/377
7	Eh	0.26	0/400	0.75	0/540
7	Ei	0.29	0/405	0.97	1/547 (0.2%)
7	Ej	0.35	0/467	1.14	4/631 (0.6%)
7	Ek	0.26	0/278	0.75	0/377
7	El	0.28	0/278	0.86	0/377
7	Em	0.35	0/384	0.99	1/518 (0.2%)
7	En	0.29	0/278	0.96	2/377 (0.5%)
7	Eo	0.27	0/278	0.83	1/377 (0.3%)
7	Ep	0.31	0/278	0.79	0/377

Mol	Chain	Bond lengths		Bond angles	
		RMSZ	# Z  >5	RMSZ	# Z  >5
7	Eq	0.30	0/462	1.04	3/624 (0.5%)
7	Er	0.36	0/278	1.07	3/377 (0.8%)
7	Es	0.28	0/457	0.88	0/617
7	Et	0.26	0/400	0.85	1/540 (0.2%)
7	Eu	0.28	0/475	0.90	2/642 (0.3%)
7	Ev	0.33	0/278	1.07	2/377 (0.5%)
7	Fg	0.25	0/337	0.74	0/451
7	Fh	0.19	0/337	0.72	1/451 (0.2%)
7	Fi	0.20	0/337	0.68	1/451 (0.2%)
7	Fj	0.19	0/337	0.68	0/451
7	Fk	0.17	0/337	0.49	0/451
8	Ew	0.25	0/3277	0.65	1/4473 (0.0%)
8	Ex	0.24	0/3277	0.64	1/4473 (0.0%)
8	Ey	0.26	0/3277	0.64	0/4473
8	Ez	0.25	0/3277	0.66	6/4473 (0.1%)
8	Fa	0.28	0/3277	0.70	3/4473 (0.1%)
9	Fb	0.23	0/6715	0.65	2/9115 (0.0%)
9	Fc	0.23	0/6715	0.61	4/9115 (0.0%)
9	Fd	0.23	0/6715	0.66	3/9115 (0.0%)
9	Fe	0.23	0/6715	0.65	5/9115 (0.1%)
9	Ff	0.23	0/6715	0.63	1/9115 (0.0%)
All	All	0.24	0/95890	0.66	81/130045 (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
6	Al	0	1
6	Ar	0	1
6	Bp	0	1
6	Bv	0	1
6	Ch	0	1
6	Ct	0	1
6	Df	0	1
6	Dl	0	1
6	Dx	0	1
7	Ej	0	1
7	En	0	1
7	Eq	0	1
7	Er	0	1

*Continued on next page...*



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Mol	Chain	#Chirality outliers	#Planarity outliers
8	Fa	0	1
9	Fb	0	1
9	Fe	0	1
All	All	0	16

There are no bond length outliers.

All (81) bond angle outliers are listed below:

Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
7	Ej	802	LEU	CA-CB-CG	9.24	148.64	116.30
7	Et	801	MET	CB-CG-SD	8.93	139.50	112.70
6	Af	110	LYS	CA-CB-CG	8.64	131.37	114.10
6	Df	146	GLY	CA-C-N	-8.61	104.57	120.94
6	Df	146	GLY	C-N-CA	-8.61	104.57	120.94
7	Eq	783	GLU	CA-CB-CG	7.80	129.70	114.10
6	Af	115	MET	CB-CG-SD	-7.73	89.50	112.70
9	Ff	77	MET	CA-CB-CG	7.23	128.56	114.10
7	Ei	802	LEU	CA-CB-CG	7.21	141.55	116.30
9	Fb	521	GLN	CA-CB-CG	7.07	128.23	114.10
6	Af	110	LYS	CB-CG-CD	6.71	126.74	111.30
8	Ez	354	ARG	CA-C-N	-6.55	108.11	121.48
8	Ez	354	ARG	C-N-CA	-6.55	108.11	121.48
8	Ew	383	LEU	CA-CB-CG	6.47	138.96	116.30
6	Dr	146	GLY	CA-C-N	-6.39	112.11	120.67
6	Dr	146	GLY	C-N-CA	-6.39	112.11	120.67
7	Eq	802	LEU	CA-CB-CG	6.20	138.00	116.30
6	Ch	146	GLY	CA-C-N	-6.16	111.74	120.49
6	Ch	146	GLY	C-N-CA	-6.16	111.74	120.49
7	Em	802	LEU	CA-CB-CG	6.15	137.83	116.30
7	Eu	802	LEU	CA-CB-CG	6.14	137.80	116.30
6	Df	115	MET	CG-SD-CE	-6.08	87.53	100.90
7	Fh	38	LYS	CA-CB-CG	6.02	126.14	114.10
8	Ez	421	GLU	CA-CB-CG	5.90	125.91	114.10
7	Ej	801	MET	CA-CB-CG	5.90	125.90	114.10
6	Cn	146	GLY	CA-C-N	-5.80	104.49	120.97
6	Cn	146	GLY	C-N-CA	-5.80	104.49	120.97
7	Eq	773	GLN	CA-CB-CG	5.79	125.67	114.10
6	Dx	114	ASP	CA-C-N	-5.77	110.90	121.52
6	Dx	114	ASP	C-N-CA	-5.77	110.90	121.52
6	Ax	115	MET	CG-SD-CE	-5.72	88.33	100.90
6	Dx	139	TYR	CA-CB-CG	5.71	124.18	113.90
6	Af	110	LYS	N-CA-CB	5.66	118.22	110.01

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
6	Dx	146	GLY	CA-C-N	-5.66	110.19	120.94
6	Dx	146	GLY	C-N-CA	-5.66	110.19	120.94
7	Ej	780	LYS	CA-CB-CG	5.58	125.25	114.10
6	Af	110	LYS	CG-CD-CE	-5.54	98.55	111.30
9	Fe	916	LEU	CA-CB-CG	5.53	135.67	116.30
6	Bv	139	TYR	CA-CB-CG	5.53	123.86	113.90
6	Cb	119	LEU	CA-CB-CG	5.51	135.58	116.30
9	Fd	928	LYS	CA-CB-CG	5.47	125.04	114.10
8	Ez	430	ARG	N-CA-CB	5.47	118.76	110.28
6	Cn	115	MET	CG-SD-CE	-5.46	88.88	100.90
8	Fa	39	LEU	CA-CB-CG	5.46	135.43	116.30
9	Fe	728	ILE	CA-C-N	-5.46	112.54	120.29
9	Fe	728	ILE	C-N-CA	-5.46	112.54	120.29
8	Ex	215	ILE	CB-CA-C	-5.45	104.88	112.02
6	Cn	115	MET	CA-CB-CG	-5.43	103.24	114.10
6	Ct	146	GLY	CA-C-N	-5.39	113.45	120.67
6	Ct	146	GLY	C-N-CA	-5.39	113.45	120.67
9	Fc	708	LEU	CA-CB-CG	5.34	135.00	116.30
9	Fb	661	MET	CA-CB-CG	5.32	124.73	114.10
8	Fa	430	ARG	CB-CG-CD	-5.29	99.13	111.30
8	Ez	429	ASP	CA-C-N	-5.28	112.28	120.31
8	Ez	429	ASP	C-N-CA	-5.28	112.28	120.31
7	Ev	795	GLN	CA-C-N	-5.28	112.28	120.31
7	Ev	795	GLN	C-N-CA	-5.28	112.28	120.31
7	Eo	802	LEU	CA-CB-CG	5.26	134.70	116.30
6	Df	125	GLU	CA-CB-CG	5.25	124.61	114.10
7	Ej	801	MET	CB-CA-C	5.20	119.51	110.72
9	Fe	77	MET	CA-CB-CG	5.19	124.48	114.10
7	Fi	27	LEU	CA-CB-CG	5.17	134.39	116.30
7	Eu	789	LYS	CA-CB-CG	5.15	124.40	114.10
9	Fe	191	MET	CA-CB-CG	5.15	124.41	114.10
6	Ax	146	GLY	CA-C-N	-5.13	111.19	120.94
6	Ax	146	GLY	C-N-CA	-5.13	111.19	120.94
9	Fc	742	MET	CB-CG-SD	5.12	128.07	112.70
7	En	800	ASP	CA-C-N	-5.11	113.77	122.56
7	En	800	ASP	C-N-CA	-5.11	113.77	122.56
9	Fc	628	MET	CB-CG-SD	5.11	128.03	112.70
6	Af	146	GLY	CA-C-N	5.10	133.34	122.17
6	Af	146	GLY	C-N-CA	5.10	133.34	122.17
9	Fd	742	MET	CB-CG-SD	5.09	127.97	112.70
9	Fc	719	LEU	N-CA-CB	-5.08	108.38	114.17
6	Dx	251	ARG	CA-CB-CG	5.07	124.25	114.10

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
7	Er	809	VAL	CA-C-N	-5.07	113.83	122.56
7	Er	809	VAL	C-N-CA	-5.07	113.83	122.56
8	Fa	212	GLN	CA-CB-CG	5.05	124.20	114.10
7	Er	802	LEU	CA-CB-CG	5.04	133.93	116.30
9	Fd	916	LEU	CA-CB-CG	5.04	133.92	116.30
6	Cz	105	GLU	CA-CB-CG	5.01	124.12	114.10

There are no chirality outliers.

All (16) planarity outliers are listed below:

Mol	Chain	Res	Type	Group
6	Al	219	LYS	Peptide
6	Ar	219	LYS	Peptide
6	Bp	238	MET	Peptide
6	Bv	214	MET	Peptide
6	Ch	219	LYS	Peptide
6	Ct	219	LYS	Peptide
6	Df	117	ARG	Sidechain
6	Dl	219	LYS	Peptide
6	Dx	117	ARG	Sidechain
7	Ej	801	MET	Peptide
7	En	797	ARG	Sidechain
7	Eq	801	MET	Peptide
7	Er	810	GLN	Peptide
8	Fa	377	ASN	Peptide
9	Fb	661	MET	Peptide
9	Fe	730	MET	Peptide

## 5.2 Too-close contacts [i](#)

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

Mol	Chain	Non-H	H(model)	H(added)	Clashes	Symm-Clashes
1	Aa	117	0	111	7	0
1	Ag	117	0	111	10	0
1	Am	117	0	111	16	0
1	As	117	0	111	6	0
1	Ay	117	0	111	9	0

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Mol	Chain	Non-H	H(model)	H(added)	Clashes	Symm-Clashes
1	Be	117	0	111	6	0
1	Bk	117	0	111	11	0
1	Bq	117	0	111	9	0
1	Bw	117	0	111	6	0
1	Cc	117	0	111	10	0
1	Ci	117	0	111	12	0
1	Co	117	0	111	8	0
1	Cu	117	0	111	6	0
1	Da	117	0	111	7	0
1	Dg	117	0	111	9	0
1	Dm	117	0	111	10	0
1	Ds	117	0	111	8	0
1	Dy	117	0	111	5	0
2	Ab	61	0	65	9	0
2	Ah	61	0	65	8	0
2	An	61	0	65	12	0
2	At	61	0	65	7	0
2	Az	61	0	65	10	0
2	Bf	61	0	65	12	0
2	Bl	61	0	65	9	0
2	Br	61	0	65	11	0
2	Bx	61	0	65	6	0
2	Cd	61	0	65	9	0
2	Cj	61	0	65	4	0
2	Cp	61	0	65	7	0
2	Cv	61	0	65	10	0
2	Db	61	0	65	9	0
2	Dh	61	0	65	10	0
2	Dn	61	0	65	10	0
2	Dt	61	0	65	7	0
2	Dz	61	0	65	7	0
3	Ac	125	0	126	9	0
3	Ai	125	0	126	11	0
3	Ao	125	0	126	17	0
3	Au	125	0	126	9	0
3	Ba	125	0	126	9	0
3	Bg	125	0	126	9	0
3	Bm	125	0	126	14	0
3	Bs	125	0	126	15	0
3	By	125	0	126	6	0
3	Ce	125	0	126	9	0
3	Ck	125	0	126	13	0

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Mol	Chain	Non-H	H(model)	H(added)	Clashes	Symm-Clashes
3	Cq	125	0	126	10	0
3	Cw	125	0	126	9	0
3	Dc	125	0	126	5	0
3	Di	125	0	126	8	0
3	Do	125	0	126	15	0
3	Du	125	0	126	13	0
3	Ea	125	0	126	9	0
4	Ad	36	0	30	2	0
4	Aj	36	0	30	4	0
4	Ap	36	0	30	5	0
4	Av	36	0	30	2	0
4	Bb	36	0	30	2	0
4	Bh	36	0	30	5	0
4	Bn	36	0	30	3	0
4	Bt	36	0	30	4	0
4	Bz	36	0	30	2	0
4	Cf	36	0	30	3	0
4	Cl	36	0	30	3	0
4	Cr	36	0	30	2	0
4	Cx	36	0	30	1	0
4	Dd	36	0	30	5	0
4	Dj	36	0	30	2	0
4	Dp	36	0	30	3	0
4	Dv	36	0	30	2	0
4	Eb	36	0	30	1	0
5	Ae	484	0	502	41	0
5	Ak	484	0	502	31	0
5	Aq	484	0	502	33	0
5	Aw	484	0	502	22	0
5	Bc	484	0	502	41	0
5	Bi	484	0	502	37	0
5	Bo	484	0	502	42	0
5	Bu	484	0	502	31	0
5	Ca	484	0	502	37	0
5	Cg	484	0	502	33	0
5	Cm	484	0	502	29	0
5	Cs	484	0	502	25	0
5	Cy	484	0	502	29	0
5	De	484	0	502	35	0
5	Dk	484	0	502	48	0
5	Dq	484	0	502	32	0
5	Dw	484	0	502	25	0

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Mol	Chain	Non-H	H(model)	H(added)	Clashes	Symm-Clashes
5	Ec	484	0	502	23	0
6	Af	1238	0	1252	108	0
6	Al	1238	0	1252	113	0
6	Ar	1238	0	1252	143	0
6	Ax	1238	0	1252	110	0
6	Bd	1238	0	1252	120	0
6	Bj	1238	0	1252	125	0
6	Bp	1238	0	1252	133	0
6	Bv	1238	0	1252	120	0
6	Cb	1238	0	1252	113	0
6	Ch	1238	0	1252	109	0
6	Cn	1238	0	1252	119	0
6	Ct	1238	0	1252	117	0
6	Cz	1238	0	1252	129	0
6	Df	1238	0	1252	126	0
6	Dl	1238	0	1252	129	0
6	Dr	1238	0	1252	134	0
6	Dx	1238	0	1252	121	0
6	Ed	1238	0	1252	121	0
7	Ee	276	0	263	24	0
7	Ef	472	0	453	48	0
7	Eg	276	0	263	30	0
7	Eh	397	0	386	36	0
7	Ei	402	0	391	45	0
7	Ej	464	0	449	53	0
7	Ek	276	0	263	30	0
7	El	276	0	263	36	0
7	Em	381	0	364	56	0
7	En	276	0	263	42	0
7	Eo	276	0	263	40	0
7	Ep	276	0	263	32	0
7	Eq	459	0	444	54	0
7	Er	276	0	263	33	0
7	Es	454	0	439	53	0
7	Et	397	0	386	51	0
7	Eu	472	0	453	37	0
7	Ev	276	0	263	43	0
7	Fg	334	0	378	24	0
7	Fh	334	0	378	26	0
7	Fi	334	0	378	21	0
7	Fj	334	0	378	19	0
7	Fk	334	0	378	17	0

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Mol	Chain	Non-H	H(model)	H(added)	Clashes	Symm-Clashes
8	Ew	3213	0	3131	301	0
8	Ex	3213	0	3131	272	0
8	Ey	3213	0	3131	328	0
8	Ez	3213	0	3131	296	0
8	Fa	3213	0	3131	291	0
9	Fb	6560	0	6545	450	0
9	Fc	6560	0	6545	458	0
9	Fd	6560	0	6545	450	0
9	Fe	6560	0	6545	420	0
9	Ff	6560	0	6545	422	0
All	All	94015	0	93950	6392	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 34.

All (6392) 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)
8:Ew:428:LEU:HB2	8:Fa:430:ARG:HH22	1.06	1.15
9:Fc:929:ALA:O	9:Fc:932:LEU:HB3	1.47	1.14
9:Ff:936:LEU:HA	9:Ff:939:LYS:HE3	1.32	1.12
8:Ey:254:MET:HE3	8:Ey:354:ARG:HB3	1.37	1.03
6:Df:115:MET:HE1	7:Ep:798:THR:HA	1.40	1.01
9:Fe:663:PRO:O	9:Fe:667:MET:HB3	1.59	1.00
8:Ew:418:LEU:HD11	8:Fa:419:LEU:HD13	1.43	0.99
6:Cz:150:LYS:H	6:Cz:247:ALA:HA	1.27	0.99
6:Bd:208:LYS:HD2	7:Ej:823:GLY:HA2	1.43	0.99
8:Fa:209:LEU:HA	8:Fa:212:GLN:HE21	1.27	0.98
6:Cb:111:ALA:HB2	7:Ek:797:ARG:HE	1.29	0.97
8:Fa:419:LEU:HA	8:Fa:422:ILE:HG12	1.46	0.97
6:Bj:126:GLN:HA	6:Bj:129:LYS:HE2	1.47	0.97
8:Ez:433:GLN:HA	8:Ez:436:ILE:HG12	1.45	0.96
8:Ew:425:GLN:HG2	8:Fa:426:MET:HE1	1.45	0.96
2:An:4:MET:HE1	5:Aq:249:SER:H	1.30	0.96
5:Dk:254:ILE:HB	5:Dk:262:ILE:HB	1.46	0.96
6:Df:204:ILE:HD13	6:Df:215:ILE:HG13	1.48	0.95
8:Ew:78:GLN:HG2	8:Ex:383:LEU:HD12	1.48	0.95
9:Fb:918:TYR:O	9:Fb:921:MET:HB2	1.66	0.95
8:Ey:413:LYS:HZ3	8:Ez:404:ASN:HA	1.30	0.94
8:Ez:254:MET:HE3	8:Ez:354:ARG:HB3	1.50	0.93
8:Ew:432:ILE:HG22	8:Fa:437:LEU:HD21	1.50	0.93

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
6:Ct:169:ILE:HG12	6:Ct:239:LEU:HD11	1.49	0.93
9:Fd:756:LEU:HA	9:Fd:759:MET:HG2	1.52	0.92
8:Ew:425:GLN:HG3	8:Fa:430:ARG:HE	1.36	0.91
8:Ew:428:LEU:HB2	8:Fa:430:ARG:NH2	1.85	0.91
5:Dk:213:GLN:HE21	5:Dk:221:TRP:HB2	1.34	0.91
8:Fa:206:TYR:HE2	8:Fa:208:ASN:HB2	1.35	0.91
8:Ew:383:LEU:HD12	8:Fa:78:GLN:HG2	1.53	0.90
8:Ew:334:ARG:HG3	8:Ex:282:TYR:HE1	1.35	0.90
6:Bv:145:PRO:HD3	6:Cb:131:LYS:HE2	1.54	0.90
9:Fe:37:PHE:HZ	9:Ff:831:TRP:HB2	1.37	0.90
8:Ew:330:PHE:HB3	8:Ew:334:ARG:HH21	1.34	0.89
6:Bv:202:PHE:HB3	6:Bv:215:ILE:HD11	1.54	0.89
6:Cb:208:LYS:HD2	7:En:823:GLY:HA2	1.54	0.89
7:Es:788:GLN:HB3	7:Es:792:GLN:HE22	1.35	0.89
6:Ax:111:ALA:HB3	7:Ef:797:ARG:HH11	1.38	0.89
9:Fb:103:ILE:HD12	9:Fb:104:PRO:HD3	1.55	0.89
8:Ey:437:LEU:HD21	8:Ez:432:ILE:HG22	1.54	0.88
8:Fa:412:GLN:HA	8:Fa:415:ILE:HD12	1.54	0.88
9:Fc:766:PHE:HA	9:Fc:769:LEU:HG	1.54	0.88
6:Ax:229:ARG:HH12	6:Ax:236:PRO:HA	1.35	0.88
8:Ey:434:GLU:HA	8:Ey:437:LEU:HD23	1.55	0.88
6:Dr:202:PHE:HB3	6:Dr:215:ILE:HD11	1.53	0.88
8:Ex:301:LEU:HG	8:Ex:326:LEU:HD21	1.55	0.88
3:Bs:4:ILE:HA	6:Ch:132:GLN:HE21	1.37	0.88
9:Fd:213:PRO:HB2	9:Fd:222:MET:HE2	1.54	0.88
6:Dx:108:ASP:HA	7:Es:797:ARG:HH22	1.38	0.88
8:Ex:254:MET:HE3	8:Ex:354:ARG:HB3	1.53	0.88
6:Cn:230:LEU:HD12	6:Cn:231:ARG:H	1.38	0.87
6:Bv:173:GLN:HE21	7:El:815:VAL:HG11	1.37	0.87
9:Fc:336:ILE:HA	9:Fc:339:MET:HG2	1.55	0.86
2:Bl:4:MET:HE1	5:Bo:249:SER:H	1.39	0.86
9:Fd:745:ILE:HG23	9:Fd:919:THR:HG21	1.57	0.86
6:Dr:149:PRO:HB2	6:Dr:248:VAL:HG13	1.57	0.86
8:Ex:188:SER:HA	8:Ex:191:MET:HG2	1.55	0.86
6:Bp:169:ILE:HG12	6:Bp:239:LEU:HD21	1.57	0.86
6:Cb:189:TRP:HE1	6:Cb:232:GLY:H	1.19	0.86
8:Ew:412:GLN:HA	8:Ew:415:ILE:HD12	1.56	0.86
8:Ew:317:VAL:HA	8:Ex:198:TRP:HE1	1.38	0.86
9:Fb:295:GLN:HE22	9:Fb:303:THR:H	1.19	0.86
9:Fb:697:ILE:HD11	9:Fc:906:ALA:HA	1.57	0.86
6:Dx:199:PRO:HD2	7:Ev:816:GLU:HG3	1.56	0.85

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:Bs:2:VAL:HG21	6:Ch:129:LYS:HG2	1.58	0.85
8:Ew:433:GLN:HA	8:Ew:436:ILE:HG12	1.57	0.85
9:Fd:759:MET:HE1	9:Fe:917:ILE:HD11	1.58	0.85
8:Ez:381:ARG:HE	8:Ez:382:ARG:HG3	1.41	0.85
6:Dr:219:LYS:HZ2	6:Dr:221:TYR:H	1.24	0.85
6:Ed:111:ALA:O	6:Ed:115:MET:HB2	1.77	0.85
9:Ff:824:ALA:O	9:Ff:827:TYR:HB3	1.76	0.85
6:Dr:208:LYS:HD2	7:Eu:823:GLY:HA2	1.57	0.85
7:Es:807:GLN:HE21	7:Es:808:LEU:HD23	1.41	0.85
8:Ew:430:ARG:NH2	8:Ex:428:LEU:HB2	1.92	0.85
9:Fb:598:VAL:HG12	9:Ff:714:SER:HB3	1.59	0.85
9:Fb:624:LEU:O	9:Fb:628:MET:HB2	1.76	0.85
6:Bd:201:SER:HA	6:Bd:219:LYS:HG3	1.59	0.85
6:Bv:108:ASP:HA	7:Ej:797:ARG:HH22	1.39	0.85
2:Bf:5:ILE:HG21	6:Bp:140:ALA:HB2	1.59	0.84
5:Bc:264:PHE:HZ	6:Bd:170:ARG:HH12	1.25	0.84
6:Bd:202:PHE:HB2	6:Bd:215:ILE:HD11	1.59	0.84
5:Dq:246:LEU:HB2	5:Dq:255:LEU:HD13	1.59	0.84
1:Am:12:HIS:HD2	3:Ao:9:TYR:HB2	1.41	0.84
8:Ey:277:LEU:HB2	8:Ey:281:ARG:HH21	1.41	0.84
6:Al:202:PHE:HB3	6:Al:215:ILE:HD11	1.60	0.83
9:Fb:929:ALA:O	9:Fb:932:LEU:HB3	1.77	0.83
6:Ct:149:PRO:HB2	6:Ct:248:VAL:HG13	1.61	0.83
8:Ez:426:MET:HE1	8:Fa:425:GLN:HG2	1.58	0.83
8:Fa:382:ARG:HB3	8:Fa:399:TRP:CE3	2.14	0.83
5:Cm:207:ARG:HD2	5:Cm:208:ILE:H	1.43	0.83
6:Ch:202:PHE:HB3	6:Ch:215:ILE:HD11	1.60	0.83
6:Ed:199:PRO:HD2	7:Ee:816:GLU:HG3	1.58	0.83
7:Em:793:GLU:HA	7:Em:796:GLN:HE21	1.43	0.83
9:Fd:663:PRO:O	9:Fd:667:MET:HB3	1.78	0.83
5:Ak:219:ARG:HA	5:Ak:232:VAL:O	1.78	0.83
5:Bu:213:GLN:HG3	6:Bv:168:VAL:HG11	1.61	0.83
8:Fa:400:ILE:HA	8:Fa:403:ILE:HD12	1.59	0.83
8:Fa:421:GLU:O	8:Fa:425:GLN:HB3	1.77	0.83
6:Bd:157:PHE:HD1	6:Bd:255:ARG:HG3	1.43	0.82
7:Ep:809:VAL:HG22	7:Ep:813:LYS:HE3	1.62	0.82
5:Dk:248:ASP:HB3	5:Dk:253:ARG:HG2	1.62	0.82
6:Dr:173:GLN:HE21	7:Et:815:VAL:HG21	1.43	0.82
8:Ex:412:GLN:HA	8:Ex:415:ILE:HD12	1.61	0.82
7:Er:807:GLN:HA	7:Er:810:GLN:NE2	1.93	0.82
8:Ez:359:MET:HE1	8:Ez:369:THR:HG22	1.61	0.82

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
6:Bv:138:GLU:HA	6:Bv:141:LYS:HB3	1.61	0.82
6:Ch:131:LYS:HG2	7:Em:812:TRP:CH2	2.15	0.82
6:Bj:138:GLU:HA	6:Bj:141:LYS:HB3	1.62	0.82
2:Cp:2:VAL:HG22	3:Cq:1:ARG:HE	1.45	0.82
6:Ct:204:ILE:HG12	6:Ct:215:ILE:HG12	1.61	0.82
6:Dx:208:LYS:HD2	7:Ev:823:GLY:HA2	1.60	0.82
8:Ew:327:SER:HB2	8:Ex:211:MET:HE1	1.61	0.82
6:Bj:193:ALA:HB3	6:Bj:229:ARG:HB3	1.61	0.81
3:Cw:2:VAL:HG11	6:Dl:129:LYS:HB3	1.62	0.81
2:Dn:4:MET:HE3	5:Dq:250:LEU:HG	1.61	0.81
6:Dx:216:GLN:HE22	7:Eu:817:THR:HA	1.45	0.81
8:Ey:274:GLN:HA	8:Ey:277:LEU:HG	1.61	0.81
8:Ez:398:GLN:HB2	8:Ez:401:LYS:HD3	1.60	0.81
9:Ff:741:THR:HG23	9:Ff:916:LEU:HD21	1.60	0.81
3:Dc:2:VAL:HG21	6:Dr:129:LYS:HE3	1.63	0.81
9:Fb:745:ILE:HG23	9:Fb:919:THR:HG21	1.62	0.81
9:Fc:86:ASN:HA	9:Fc:89:HIS:HD2	1.44	0.81
9:Fd:528:ILE:HD11	9:Fd:538:PRO:HD3	1.63	0.81
9:Fd:709:ASN:HD21	9:Fe:597:LYS:H	1.23	0.81
5:Bo:219:ARG:HH21	5:Bo:231:THR:HG23	1.46	0.81
8:Ey:419:LEU:HD13	8:Ez:418:LEU:HD13	1.63	0.80
6:Af:131:LYS:HE2	6:Ed:145:PRO:HD3	1.62	0.80
6:Bv:117:ARG:HH12	6:Bv:118:ASN:HB3	1.45	0.80
6:Bd:234:ASN:HD22	6:Bj:188:PRO:HG2	1.44	0.80
9:Fc:760:ILE:HD12	9:Fc:930:PHE:HB3	1.63	0.80
6:Bp:229:ARG:HA	6:Bp:236:PRO:HB3	1.63	0.80
6:Dx:126:GLN:HB3	6:Ed:112:PHE:CZ	2.16	0.80
5:Ae:238:ILE:HD12	5:Ae:239:PRO:HD2	1.64	0.80
9:Fe:771:ALA:HA	9:Fe:774:GLU:HG3	1.63	0.80
8:Ew:398:GLN:HB2	8:Ew:401:LYS:HD3	1.62	0.80
9:Fc:124:MET:HG3	9:Fc:128:PHE:HE1	1.46	0.80
9:Fd:696:TYR:HE1	9:Fd:745:ILE:HB	1.45	0.80
9:Ff:745:ILE:HG23	9:Ff:919:THR:HG21	1.62	0.80
6:Al:219:LYS:HD2	6:Al:222:ASN:HD21	1.46	0.80
7:Em:807:GLN:HA	7:Em:810:GLN:HE21	1.46	0.80
8:Ey:75:GLN:HE21	8:Ey:79:ASN:HD21	1.30	0.80
6:Al:208:LYS:HD2	7:Eg:823:GLY:HA2	1.62	0.80
8:Ex:82:TYR:HB2	8:Ey:383:LEU:HD21	1.64	0.80
8:Ex:437:LEU:HD13	8:Ey:436:ILE:HD13	1.63	0.80
8:Fa:426:MET:HA	8:Fa:429:ASP:OD2	1.82	0.80
6:Af:230:LEU:HD12	6:Af:231:ARG:H	1.46	0.79

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
6:Al:115:MET:HE3	7:Ev:801:MET:HB2	1.61	0.79
6:Dx:189:TRP:HE1	6:Dx:232:GLY:H	1.30	0.79
7:Ev:809:VAL:HA	7:Ev:812:TRP:CD1	2.17	0.79
7:Ep:807:GLN:HA	7:Ep:810:GLN:NE2	1.98	0.79
8:Ey:339:GLN:HE22	8:Ez:283:ALA:HA	1.46	0.79
6:Bd:189:TRP:HE1	6:Bd:232:GLY:H	1.31	0.79
6:Af:121:PRO:HB2	7:Ev:801:MET:HG3	1.63	0.79
9:Ff:188:GLN:HE22	9:Ff:343:LEU:HB2	1.48	0.79
6:Cz:107:ILE:HA	6:Cz:110:LYS:HZ3	1.47	0.79
6:Al:171:LEU:HD12	6:Al:172:SER:H	1.48	0.79
6:Ax:204:ILE:HD11	6:Ax:213:LEU:HD22	1.64	0.79
8:Ew:427:TYR:HD1	8:Ew:428:LEU:HD13	1.48	0.79
8:Ey:412:GLN:HA	8:Ey:415:ILE:HD12	1.64	0.79
8:Ey:419:LEU:HD11	8:Ez:418:LEU:HD22	1.65	0.79
9:Fb:566:THR:HG23	9:Fb:569:GLY:H	1.47	0.79
9:Fb:697:ILE:HD12	9:Fc:909:TYR:HD2	1.47	0.79
9:Ff:537:VAL:HG13	9:Ff:542:GLN:HA	1.64	0.79
6:Ar:219:LYS:HE3	6:Ar:222:ASN:HD21	1.47	0.79
8:Fa:277:LEU:HD23	8:Fa:281:ARG:HH12	1.47	0.79
9:Ff:500:GLN:HA	9:Ff:503:LYS:HG3	1.64	0.79
6:Ax:134:TYR:HE1	7:Eg:809:VAL:HG23	1.48	0.79
6:Ed:169:ILE:HD11	6:Ed:239:LEU:HB3	1.64	0.79
6:Dx:126:GLN:HA	6:Dx:129:LYS:HG2	1.65	0.78
7:Ei:777:ILE:HA	7:Ei:780:LYS:HG2	1.65	0.78
9:Fe:661:MET:HE1	9:Fe:703:LEU:HD21	1.65	0.78
5:Ae:241:TYR:HB3	5:Ae:256:THR:HG21	1.63	0.78
2:An:2:VAL:HA	3:Ao:1:ARG:HE	1.45	0.78
8:Ey:208:ASN:O	8:Ey:212:GLN:HG2	1.84	0.78
5:Bo:210:TYR:HB3	5:Bo:222:LEU:HD13	1.65	0.78
9:Fb:528:ILE:HD11	9:Fb:538:PRO:HD3	1.64	0.78
6:Dl:229:ARG:HH12	6:Dl:233:LEU:HB2	1.48	0.78
8:Ez:430:ARG:HH12	8:Fa:425:GLN:HG3	1.49	0.78
6:Ar:149:PRO:HB2	6:Ar:248:VAL:HG13	1.63	0.78
6:Ed:117:ARG:HH21	7:Eu:797:ARG:HG3	1.48	0.78
7:Em:793:GLU:HA	7:Em:796:GLN:NE2	1.98	0.78
8:Ey:234:LYS:HA	8:Ey:237:GLN:HE21	1.48	0.78
8:Ey:330:PHE:HB2	8:Ey:334:ARG:HH22	1.48	0.78
9:Fe:261:LEU:HD11	9:Fe:265:ASN:HD21	1.49	0.78
8:Ey:207:GLN:HB2	8:Ey:289:PRO:HG2	1.66	0.78
8:Ey:423:ASN:HD22	8:Ez:382:ARG:HH22	1.32	0.78
6:Ed:144:THR:HG21	6:Ed:148:PRO:HG3	1.66	0.78

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
8:Ey:382:ARG:HE	8:Ey:418:LEU:HD23	1.49	0.78
3:Ac:2:VAL:HG21	6:Ar:129:LYS:HD3	1.65	0.77
6:Ar:181:PHE:HD1	6:Ar:254:LEU:HD11	1.49	0.77
9:Fd:697:ILE:HD12	9:Fe:909:TYR:HD1	1.49	0.77
8:Ey:189:PHE:CD1	8:Ey:295:LEU:HD23	2.20	0.77
9:Fb:828:VAL:HG21	9:Ff:41:LEU:HD12	1.64	0.77
9:Fd:573:ASN:HA	9:Fd:576:MET:HE1	1.66	0.77
6:Af:126:GLN:HA	6:Af:129:LYS:HE2	1.65	0.77
9:Fb:721:GLY:HA2	9:Fb:724:ILE:HG12	1.66	0.77
6:Df:121:PRO:HB2	7:Eq:797:ARG:HH12	1.49	0.77
7:Ep:807:GLN:HA	7:Ep:810:GLN:HE21	1.49	0.77
9:Fe:704:TRP:HE1	9:Ff:734:LEU:HB2	1.50	0.77
5:Ec:254:ILE:HB	5:Ec:262:ILE:HB	1.66	0.77
7:Ev:809:VAL:HA	7:Ev:812:TRP:HD1	1.46	0.77
8:Ex:142:ILE:HA	8:Ex:210:VAL:HG23	1.66	0.77
8:Ez:214:VAL:HG22	8:Ez:267:LEU:HD11	1.66	0.77
6:Bp:171:LEU:HD11	6:Bp:241:LEU:HB2	1.67	0.77
5:Aw:264:PHE:HB3	5:Aw:269:SER:HB3	1.67	0.77
6:Bj:129:LYS:HE3	6:Bp:112:PHE:CE2	2.20	0.77
6:Cb:112:PHE:O	6:Cb:116:THR:HG23	1.85	0.77
8:Fa:254:MET:HE3	8:Fa:354:ARG:HB2	1.67	0.77
6:Cn:214:MET:HE1	7:Eo:818:GLN:HE21	1.50	0.77
9:Fe:336:ILE:HA	9:Fe:339:MET:HG2	1.65	0.77
6:Bd:150:LYS:H	6:Bd:247:ALA:HA	1.50	0.76
6:Bv:130:LEU:HA	6:Bv:133:ILE:HD12	1.67	0.76
6:Df:229:ARG:HD2	6:Df:236:PRO:HB3	1.68	0.76
9:Fe:528:ILE:HD11	9:Fe:538:PRO:HD3	1.66	0.76
6:Ct:121:PRO:HB2	7:Eo:801:MET:HE1	1.65	0.76
5:De:245:LYS:HB2	5:De:246:LEU:HD22	1.68	0.76
8:Ex:280:ILE:HB	8:Ex:347:LEU:HD13	1.65	0.76
9:Fb:905:TRP:HE1	9:Ff:702:THR:HG22	1.50	0.76
5:Ca:238:ILE:HD11	5:Ca:244:VAL:HG23	1.67	0.76
7: Ei:799:SER:HA	7: Ei:802:LEU:HD23	1.66	0.76
5:Dq:238:ILE:HD12	5:Dq:239:PRO:HD2	1.67	0.76
8:Ex:55:THR:HA	8:Ey:42:TYR:HE1	1.51	0.76
6:Ar:126:GLN:HA	6:Ar:129:LYS:HZ2	1.49	0.76
5:Bc:246:LEU:HB3	5:Bc:255:LEU:HD23	1.68	0.76
6:Cb:141:LYS:HZ2	6:Ch:127:VAL:HG13	1.50	0.76
6:Cz:169:ILE:HD11	6:Cz:239:LEU:HD23	1.67	0.76
8:Ew:418:LEU:CD1	8:Fa:419:LEU:HD13	2.16	0.76
8:Ey:111:LEU:HD11	8:Ez:386:PRO:HB3	1.67	0.76

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
9:Fd:766:PHE:HA	9:Fd:769:LEU:HG	1.67	0.76
7:Ek:814:GLN:HE22	7:Ek:816:GLU:HB2	1.48	0.76
8:Ey:188:SER:HA	8:Ey:191:MET:HG2	1.67	0.76
7:Em:788:GLN:HG3	9:Fc:235:SER:HB2	1.67	0.76
8:Ez:448:LEU:HD13	8:Fa:340:THR:HG22	1.67	0.76
9:Fc:759:MET:HA	9:Fd:921:MET:HE3	1.67	0.76
1:Ay:2:THR:HG21	5:Bc:253:ARG:HH12	1.50	0.76
6:Dx:117:ARG:NH1	6:Dx:118:ASN:HA	2.01	0.76
7:Eu:786:ALA:HA	7:Eu:789:LYS:HZ3	1.49	0.76
1:As:15:ASN:HB3	2:Az:9:ARG:HH22	1.50	0.76
6:Dx:196:LEU:HD11	6:Dx:202:PHE:HB2	1.67	0.76
9:Fb:123:MET:HA	9:Fb:126:VAL:HG22	1.66	0.76
9:Fe:391:GLU:HG3	9:Fe:393:GLN:H	1.49	0.76
6:Ct:242:ILE:HG13	6:Ct:245:GLN:NE2	2.01	0.75
9:Fd:728:ILE:HD13	9:Fe:729:MET:SD	2.25	0.75
6:Ct:128:VAL:HG23	6:Ct:132:GLN:HE21	1.49	0.75
7:Ev:797:ARG:HG3	7:Ev:801:MET:HE1	1.68	0.75
8:Ex:277:LEU:HD13	8:Ex:281:ARG:HH22	1.51	0.75
8:Ez:301:LEU:HD13	8:Ez:326:LEU:HD21	1.67	0.75
8:Fa:206:TYR:H	8:Fa:209:LEU:HD12	1.51	0.75
9:Fe:37:PHE:CZ	9:Ff:831:TRP:HB2	2.21	0.75
9:Fe:692:MET:HA	9:Fe:695:MET:HG2	1.68	0.75
8:Ew:430:ARG:HH22	8:Ex:428:LEU:HD13	1.51	0.75
9:Fc:57:MET:HB3	9:Fc:61:PHE:HE1	1.51	0.75
9:Fe:37:PHE:HE2	9:Ff:832:ILE:HD13	1.49	0.75
6:Df:126:GLN:HA	6:Df:129:LYS:HG2	1.67	0.75
9:Fc:778:ALA:HA	9:Fc:781:ILE:HD12	1.68	0.75
6:Al:144:THR:HG21	6:Al:148:PRO:HG3	1.69	0.75
6:Ar:115:MET:HE1	7:Ee:801:MET:HB2	1.67	0.75
6:Bv:114:ASP:HB2	6:Bv:117:ARG:HH21	1.50	0.75
8:Ez:437:LEU:HD11	8:Fa:346:ASN:HB3	1.69	0.75
9:Fc:251:LEU:HD11	9:Fc:326:GLN:HE22	1.52	0.75
2:Ab:5:ILE:HD12	6:Al:139:TYR:HD2	1.50	0.75
6:Bv:204:ILE:HB	6:Bv:215:ILE:HD13	1.68	0.75
6:Cz:106:VAL:HA	6:Cz:109:LYS:HG2	1.69	0.75
8:Ex:280:ILE:HG21	8:Ex:351:LEU:HD23	1.67	0.75
9:Fd:251:LEU:HD21	9:Fd:297:ASP:HA	1.68	0.75
6:Cz:171:LEU:HD11	6:Cz:241:LEU:HB3	1.67	0.75
7:Eg:807:GLN:HA	7:Eg:810:GLN:HG3	1.67	0.75
9:Ff:905:TRP:HE3	9:Ff:909:TYR:CE1	2.05	0.75
5:Dq:254:ILE:HB	5:Dq:262:ILE:HB	1.67	0.75

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
8:Ez:427:TYR:HD1	8:Ez:430:ARG:HE	1.33	0.75
9:Fc:825:LEU:HD12	9:Fc:929:ALA:HB2	1.67	0.74
6:Af:149:PRO:HB2	6:Af:248:VAL:HG13	1.69	0.74
6:Bd:106:VAL:HA	6:Bd:109:LYS:HG2	1.69	0.74
5:Ak:238:ILE:HB	5:Ak:241:TYR:HB2	1.69	0.74
6:Ar:131:LYS:HZ2	7:Ef:812:TRP:HZ3	1.33	0.74
6:Ct:106:VAL:HA	6:Ct:109:LYS:HG2	1.69	0.74
6:Ed:201:SER:HB2	6:Ed:219:LYS:HD2	1.67	0.74
8:Ew:245:SER:HA	8:Ew:279:PHE:CE1	2.23	0.74
9:Fc:648:PHE:O	9:Fc:652:ILE:HG13	1.86	0.74
9:Ff:420:LEU:HD21	9:Ff:575:MET:HA	1.70	0.74
6:Ar:208:LYS:HE3	7:Eh:823:GLY:HA2	1.69	0.74
7:Eq:799:SER:HA	7:Eq:802:LEU:HD23	1.69	0.74
8:Ey:423:ASN:HD21	8:Ez:379:ALA:HA	1.52	0.74
8:Ey:424:TYR:HE1	8:Ez:380:THR:HA	1.50	0.74
6:Ch:115:MET:HE1	7:El:801:MET:HG2	1.70	0.74
5:Cs:230:LEU:HD12	5:Cs:231:THR:H	1.51	0.74
8:Ex:428:LEU:HA	8:Ex:431:GLN:HG2	1.67	0.74
9:Fd:239:PHE:HD1	9:Fd:261:LEU:HD13	1.52	0.74
6:Bp:112:PHE:HA	6:Bp:115:MET:HG2	1.70	0.74
6:Df:117:ARG:NH2	7:Eq:797:ARG:HD3	2.03	0.74
9:Ff:540:LEU:HD11	9:Ff:562:PRO:HB2	1.68	0.74
3:Ba:2:VAL:HG21	6:Bp:129:LYS:HG2	1.69	0.74
9:Fb:202:ARG:HG3	9:Fb:230:ILE:HD11	1.68	0.74
6:Cb:214:MET:HE1	7:Em:818:GLN:HG3	1.69	0.74
9:Fb:634:ASN:HA	9:Fb:638:ARG:HG3	1.68	0.74
9:Fc:103:ILE:HG22	9:Fc:106:ARG:HH21	1.53	0.74
9:Fc:110:GLY:HA3	9:Fc:787:THR:HG23	1.68	0.74
9:Fc:540:LEU:HD21	9:Fc:563:LEU:HA	1.68	0.74
9:Fe:694:THR:O	9:Fe:698:ASN:HB2	1.87	0.73
9:Ff:707:LEU:HA	9:Ff:710:MET:SD	2.28	0.73
8:Fa:253:LEU:HD13	8:Fa:373:LEU:HD22	1.70	0.73
5:Cm:216:ILE:HD11	6:Ct:148:PRO:HG2	1.70	0.73
5:Dw:219:ARG:HH21	5:Dw:231:THR:HG23	1.52	0.73
7:Eq:809:VAL:HA	7:Eq:812:TRP:HD1	1.52	0.73
9:Fd:202:ARG:HG3	9:Fd:230:ILE:HD11	1.69	0.73
9:Fc:732:MET:HA	9:Fc:735:LEU:HD12	1.70	0.73
2:An:4:MET:CE	5:Aq:249:SER:H	2.01	0.73
2:Bx:6:GLY:HA2	6:Cn:125:GLU:HG3	1.70	0.73
6:Df:169:ILE:HD11	6:Df:239:LEU:HB3	1.70	0.73
8:Ew:324:ALA:HB2	8:Ex:208:ASN:HD21	1.53	0.73

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
9:Fc:511:ASP:OD2	9:Fc:512:PRO:HA	1.87	0.73
6:Bp:126:GLN:HA	6:Bp:129:LYS:HG3	1.68	0.73
5:Cy:219:ARG:HA	5:Cy:232:VAL:O	1.87	0.73
8:Ey:317:VAL:HA	8:Ez:198:TRP:HZ3	1.52	0.73
8:Fa:427:TYR:O	8:Fa:430:ARG:HB2	1.88	0.73
9:Fb:766:PHE:HB3	9:Fc:928:LYS:NZ	2.04	0.73
6:Af:126:GLN:HA	6:Af:129:LYS:HG3	1.70	0.73
8:Ew:430:ARG:HG2	8:Ex:429:ASP:OD2	1.89	0.73
9:Fb:812:PRO:HA	9:Fb:815:MET:HE3	1.70	0.73
9:Fc:527:LEU:HA	9:Fc:530:ILE:HG12	1.71	0.73
9:Ff:674:GLY:HA2	9:Ff:692:MET:HE1	1.71	0.73
9:Fc:769:LEU:O	9:Fc:773:ILE:HD12	1.89	0.73
6:Bp:238:MET:HE1	6:Bv:178:SER:HB3	1.70	0.73
8:Ey:419:LEU:CD1	8:Ez:418:LEU:HD13	2.19	0.73
6:Al:106:VAL:HA	6:Al:109:LYS:HD3	1.71	0.72
6:Cb:111:ALA:HB2	7:Ek:797:ARG:NE	2.03	0.72
6:Cb:115:MET:HG3	6:Cb:118:ASN:HB2	1.71	0.72
6:Ct:229:ARG:HH11	6:Ct:236:PRO:HA	1.52	0.72
5:Dw:210:TYR:HB3	5:Dw:222:LEU:HD13	1.71	0.72
6:Dx:214:MET:HE1	7:Eu:818:GLN:HE21	1.53	0.72
9:Fc:692:MET:HA	9:Fc:695:MET:HG2	1.71	0.72
9:Fd:212:PRO:HA	9:Fd:215:GLY:H	1.52	0.72
9:Fd:397:GLN:HE21	9:Fd:578:GLN:HB2	1.54	0.72
9:Fb:383:LYS:HA	9:Fb:396:CYS:HA	1.71	0.72
1:Bk:3:ASP:HB2	2:Bl:9:ARG:HH22	1.52	0.72
2:Dt:5:ILE:HD12	6:Ed:139:TYR:HD2	1.55	0.72
8:Ez:302:TRP:HD1	8:Ez:303:ASN:HD22	1.37	0.72
9:Fe:326:GLN:HA	9:Fe:329:ARG:HG2	1.71	0.72
6:Ax:212:THR:HG22	6:Ax:214:MET:HE1	1.72	0.72
5:Dk:238:ILE:HD12	5:Dk:239:PRO:HD2	1.71	0.72
8:Ew:380:THR:HA	8:Fa:424:TYR:CE2	2.24	0.72
9:Fc:42:PHE:HA	9:Fc:57:MET:HG3	1.72	0.72
6:Al:230:LEU:HD12	6:Al:231:ARG:H	1.54	0.72
6:Dl:208:LYS:HD2	7:Et:823:GLY:HA2	1.71	0.72
8:Ez:457:PHE:HZ	8:Fa:293:PRO:HB3	1.54	0.72
9:Fc:500:GLN:HA	9:Fc:503:LYS:HG3	1.68	0.72
6:Cb:144:THR:HG21	6:Cb:148:PRO:HG3	1.72	0.72
8:Ew:330:PHE:HB3	8:Ew:334:ARG:NH2	2.05	0.72
9:Fb:732:MET:HB2	9:Fb:736:MET:HE1	1.71	0.72
6:Ar:221:TYR:CE1	6:Ax:139:TYR:HA	2.23	0.72
6:Bj:225:ASN:HD21	6:Bp:176:VAL:H	1.38	0.72

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
6:Ed:208:LYS:HG3	7:Ee:824:THR:HB	1.70	0.72
8:Ey:421:GLU:O	8:Ey:425:GLN:HG2	1.89	0.72
9:Fd:41:LEU:O	9:Fd:57:MET:HE3	1.89	0.72
9:Fe:662:ILE:HG23	9:Fe:663:PRO:HD3	1.72	0.72
6:Bv:206:TRP:HB2	6:Bv:213:LEU:HD23	1.71	0.72
6:Ed:156:GLN:HE22	6:Ed:158:VAL:HG22	1.55	0.72
8:Ex:189:PHE:HA	8:Ex:295:LEU:HD12	1.72	0.72
6:Bp:168:VAL:HA	6:Bp:240:THR:O	1.90	0.72
2:Cj:2:VAL:HG22	3:Ck:1:ARG:HE	1.54	0.72
5:Cy:230:LEU:HD12	5:Cy:231:THR:H	1.52	0.72
8:Ey:422:ILE:HA	8:Ey:425:GLN:HE21	1.55	0.72
8:Fa:170:LEU:HD13	8:Fa:176:GLN:HA	1.72	0.72
5:Cg:238:ILE:HD11	5:Cg:244:VAL:HG23	1.69	0.72
7:Eg:798:THR:HA	7:Eg:801:MET:HG2	1.72	0.72
8:Ew:350:ILE:HA	8:Ew:353:LYS:HE3	1.72	0.72
8:Ey:277:LEU:HB2	8:Ey:281:ARG:NH2	2.05	0.72
6:Bv:111:ALA:HB3	7:Ej:797:ARG:HH21	1.53	0.71
9:Fc:56:ILE:HA	9:Fd:149:TYR:HE2	1.55	0.71
9:Fc:339:MET:HE1	9:Fc:426:TYR:HB2	1.72	0.71
9:Fd:41:LEU:HB2	9:Fe:828:VAL:HG11	1.70	0.71
9:Ff:527:LEU:HA	9:Ff:530:ILE:HG12	1.72	0.71
6:Bj:202:PHE:HB3	6:Bj:215:ILE:HD11	1.72	0.71
8:Ex:191:MET:HE1	8:Ex:198:TRP:CD2	2.25	0.71
9:Fe:920:SER:O	9:Fe:924:ILE:HD12	1.90	0.71
5:Ae:219:ARG:HA	5:Ae:232:VAL:O	1.89	0.71
2:At:2:VAL:HG22	3:Au:1:ARG:HE	1.54	0.71
6:Bd:115:MET:SD	6:Bd:119:LEU:HD23	2.29	0.71
5:Cm:233:ARG:NH1	5:Cm:236:SER:HB3	2.05	0.71
5:Cs:219:ARG:NH1	5:Cs:233:ARG:HB3	2.04	0.71
8:Ex:330:PHE:O	8:Ex:334:ARG:HG2	1.89	0.71
9:Fd:140:ASP:O	9:Fd:144:GLU:HG3	1.91	0.71
9:Fd:196:LYS:HG2	9:Fd:357:PHE:HB3	1.72	0.71
6:Cb:202:PHE:HB3	6:Cb:215:ILE:HD11	1.72	0.71
8:Ey:424:TYR:CE1	8:Ez:380:THR:HA	2.25	0.71
9:Fd:272:PHE:HB3	9:Fd:275:LEU:HD22	1.71	0.71
9:Ff:335:ALA:O	9:Ff:338:GLN:HG2	1.90	0.71
6:Al:130:LEU:HG	7:Ee:812:TRP:HH2	1.54	0.71
6:Bp:141:LYS:HD3	7:Ek:808:LEU:HD12	1.72	0.71
6:Cz:225:ASN:HB2	6:Cz:238:MET:HE1	1.72	0.71
6:Df:145:PRO:HD2	6:Dl:131:LYS:NZ	2.06	0.71
6:Df:145:PRO:HD2	6:Dl:131:LYS:HZ2	1.56	0.71

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
6:Bp:173:GLN:HE21	7:Ek:815:VAL:HG22	1.55	0.71
6:Cb:138:GLU:HA	6:Cb:141:LYS:HG2	1.70	0.71
6:Cb:204:ILE:HD11	6:Cb:213:LEU:HD22	1.73	0.71
7:Ef:810:GLN:HA	7:Ef:813:LYS:HG2	1.73	0.71
8:Fa:228:PRO:HB3	8:Fa:234:LYS:HE3	1.72	0.71
9:Fb:704:TRP:HH2	9:Fc:731:ALA:HA	1.55	0.71
3:Ao:2:VAL:HG11	6:Bd:129:LYS:HG2	1.73	0.71
9:Fb:469:PHE:CE2	9:Fc:843:TYR:HB2	2.26	0.71
9:Fc:239:PHE:CD2	9:Fc:336:ILE:HG13	2.26	0.71
9:Fd:907:GLY:O	9:Fd:910:ALA:HB3	1.90	0.71
9:Fd:932:LEU:HA	9:Fd:935:HIS:CD2	2.26	0.71
6:Af:132:GLN:HE21	6:Af:136:THR:HG23	1.56	0.71
6:Ct:115:MET:HE1	7:En:801:MET:HB2	1.73	0.71
6:Ed:111:ALA:HB1	7:Et:797:ARG:HH21	1.56	0.71
8:Ew:403:ILE:HG22	8:Fa:413:LYS:HG2	1.72	0.71
6:Ax:134:TYR:CE1	7:Eg:809:VAL:HG23	2.25	0.70
8:Fa:211:MET:HA	8:Fa:214:VAL:HG23	1.73	0.70
9:Fb:103:ILE:HD11	7:Fh:20:ILE:HD12	1.73	0.70
1:Am:15:ASN:HB3	2:At:9:ARG:HH22	1.56	0.70
2:Dh:4:MET:SD	5:Dk:248:ASP:HA	2.32	0.70
7:Ek:793:GLU:HA	7:Ek:796:GLN:HG3	1.72	0.70
8:Ew:383:LEU:HD12	8:Fa:78:GLN:CG	2.20	0.70
8:Ez:182:LEU:HD11	8:Fa:243:LEU:HD12	1.73	0.70
9:Fd:593:LEU:HD11	9:Fd:905:TRP:HB3	1.73	0.70
9:Fe:812:PRO:HA	9:Fe:815:MET:HG2	1.73	0.70
5:Bo:241:TYR:HB3	5:Bo:256:THR:HG21	1.73	0.70
8:Ez:207:GLN:HB2	8:Ez:289:PRO:HG2	1.73	0.70
9:Fb:745:ILE:O	9:Fb:749:THR:HG22	1.90	0.70
9:Fe:111:LEU:HD13	7:Fj:31:ALA:HB2	1.72	0.70
9:Fe:500:GLN:HA	9:Fe:503:LYS:HG2	1.73	0.70
2:Bx:4:MET:HE1	5:Ca:249:SER:H	1.56	0.70
9:Fb:259:PHE:HE2	9:Fb:286:ASN:HB2	1.54	0.70
9:Fc:240:VAL:HG23	9:Fc:241:LYS:HD3	1.73	0.70
9:Ff:177:VAL:HG12	9:Ff:491:LEU:HD23	1.72	0.70
5:Aq:247:ILE:HG23	5:Aq:254:ILE:HD12	1.73	0.70
8:Ew:208:ASN:HD21	8:Fa:324:ALA:HB2	1.55	0.70
8:Ey:189:PHE:HD1	8:Ey:295:LEU:HD23	1.56	0.70
9:Fd:339:MET:SD	9:Fd:426:TYR:HB2	2.31	0.70
9:Ff:655:ILE:HG23	9:Ff:659:PHE:HE1	1.54	0.70
2:At:4:MET:HE2	5:Aw:249:SER:H	1.56	0.70
7:Ef:801:MET:HA	7:Ef:801:MET:HE3	1.73	0.70

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
8:Ew:254:MET:HE3	8:Ew:354:ARG:HB2	1.73	0.70
8:Ey:174:VAL:HG11	8:Ey:345:SER:HA	1.72	0.70
9:Fd:732:MET:O	9:Fd:736:MET:HB2	1.91	0.70
6:Ar:238:MET:HG3	6:Ax:251:ARG:HG3	1.74	0.70
5:Cy:264:PHE:HB3	5:Cy:269:SER:HB3	1.73	0.70
8:Ey:249:ILE:HD12	8:Ey:354:ARG:HH12	1.56	0.70
9:Ff:239:PHE:HD2	9:Ff:261:LEU:HD23	1.55	0.70
6:Al:189:TRP:CG	6:Al:230:LEU:HD11	2.27	0.70
6:Ar:205:GLN:HB2	7:Eg:820:TYR:CZ	2.27	0.70
6:Dl:115:MET:HE1	7:Eq:802:LEU:HD13	1.74	0.70
6:Dr:238:MET:HE1	6:Dx:251:ARG:HB3	1.71	0.70
8:Ex:334:ARG:HD3	8:Ey:282:TYR:CE1	2.27	0.70
8:Ey:186:ASP:H	8:Ey:189:PHE:HE2	1.39	0.70
6:Bj:115:MET:HE1	7:Eh:802:LEU:HD13	1.74	0.70
8:Ew:277:LEU:O	8:Ew:281:ARG:HG3	1.91	0.70
8:Ey:178:VAL:HG23	8:Ey:337:ALA:HB1	1.72	0.70
9:Fd:743:VAL:HG23	9:Fd:747:PHE:HE1	1.57	0.70
7:Fj:17:THR:O	7:Fj:21:ILE:HG12	1.92	0.70
6:Ar:107:ILE:HA	6:Ar:110:LYS:HZ3	1.57	0.70
5:Bo:253:ARG:HA	5:Bo:263:LYS:HZ1	1.57	0.70
6:Dr:219:LYS:HD2	6:Dr:220:LEU:H	1.56	0.70
6:Dx:110:LYS:HZ1	7:Es:794:ILE:HG12	1.56	0.70
6:Bv:246:LYS:HE2	6:Bv:246:LYS:HA	1.73	0.69
7:Eh:799:SER:HA	7:Eh:802:LEU:HD23	1.74	0.69
7:Ei:785:LEU:HD23	9:Fd:241:LYS:HD3	1.73	0.69
8:Ew:178:VAL:HG22	8:Ew:341:SER:HB3	1.74	0.69
9:Fc:590:PHE:HA	9:Fc:593:LEU:HD13	1.74	0.69
9:Fd:921:MET:O	9:Fd:925:ILE:HD12	1.92	0.69
6:Cb:141:LYS:NZ	7:Em:808:LEU:HD13	2.07	0.69
7:Ej:777:ILE:HA	7:Ej:780:LYS:HG2	1.74	0.69
9:Fe:648:PHE:O	9:Fe:652:ILE:HG13	1.91	0.69
2:Ah:2:VAL:HA	3:Ai:1:ARG:HE	1.56	0.69
5:Aq:219:ARG:HH22	6:Ax:150:LYS:HE2	1.56	0.69
1:Da:6:LEU:O	1:Da:10:GLU:HG2	1.92	0.69
6:Dx:225:ASN:HD21	6:Ed:176:VAL:H	1.41	0.69
7:Em:808:LEU:O	7:Em:812:TRP:CD1	2.45	0.69
8:Ew:385:ASP:H	8:Ew:398:GLN:HE22	1.41	0.69
8:Ey:381:ARG:HH22	8:Ey:382:ARG:HD2	1.57	0.69
8:Ew:353:LYS:HG3	8:Ew:354:ARG:HE	1.56	0.69
9:Fc:701:GLY:O	9:Fc:705:LEU:HD22	1.92	0.69
6:Al:149:PRO:HB2	6:Al:248:VAL:HG13	1.73	0.69

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
6:Bd:153:ALA:HB1	6:Bd:251:ARG:HH21	1.58	0.69
6:Ch:166:PRO:HB3	6:Cn:250:TYR:CE2	2.27	0.69
7:Er:807:GLN:HA	7:Er:810:GLN:HE21	1.56	0.69
9:Fe:472:VAL:HG12	9:Fe:684:ASN:HD21	1.58	0.69
6:Ar:158:VAL:HG22	6:Ar:167:PRO:HG2	1.73	0.69
5:Aw:219:ARG:CZ	5:Aw:233:ARG:HB3	2.22	0.69
8:Ey:302:TRP:CZ3	8:Ez:218:LEU:HG	2.28	0.69
8:Ez:429:ASP:HA	8:Ez:432:ILE:HG12	1.72	0.69
9:Fc:653:ASN:HB3	9:Fc:657:MET:HE1	1.75	0.69
9:Fe:596:PHE:CD2	9:Fe:656:VAL:HG11	2.27	0.69
9:Ff:739:ILE:O	9:Ff:743:VAL:HG12	1.92	0.69
3:Bg:2:VAL:HG21	6:Bv:129:LYS:HG2	1.74	0.69
2:Cd:2:VAL:HG22	3:Ce:1:ARG:HE	1.57	0.69
5:Cy:219:ARG:HH21	5:Cy:231:THR:HG23	1.57	0.69
6:Cz:121:PRO:HB2	7:Ep:801:MET:SD	2.33	0.69
6:Dr:202:PHE:HZ	6:Dr:243:PRO:HD3	1.57	0.69
8:Ey:86:LEU:HD12	8:Ey:115:LEU:HD12	1.75	0.69
8:Fa:438:LEU:O	8:Fa:442:ILE:HG12	1.92	0.69
9:Fb:71:LEU:O	9:Fb:75:ILE:HG12	1.92	0.69
9:Fe:332:ARG:HH21	9:Fe:489:THR:HA	1.56	0.69
7:Fh:12:PHE:HA	7:Fh:18:ARG:HE	1.58	0.69
6:Bv:144:THR:HG21	6:Bv:148:PRO:HG3	1.75	0.69
7:Et:796:GLN:HE22	9:Ff:228:THR:HA	1.57	0.69
8:Ez:252:LEU:HD12	8:Ez:373:LEU:HA	1.74	0.69
9:Fc:520:PHE:HE2	9:Fc:526:LYS:HB2	1.58	0.69
6:Cz:219:LYS:HZ2	6:Cz:221:TYR:H	1.41	0.69
6:Dr:111:ALA:HB1	6:Dr:115:MET:CE	2.23	0.69
7:Eq:780:LYS:HA	7:Eq:783:GLU:OE1	1.93	0.69
8:Ey:430:ARG:HH12	8:Ez:428:LEU:HB2	1.58	0.69
9:Ff:793:GLU:HG3	9:Ff:795:PHE:H	1.57	0.69
1:Am:12:HIS:CD2	3:Ao:9:TYR:HB2	2.27	0.68
8:Ex:205:MET:HA	8:Ex:209:LEU:HD23	1.75	0.68
9:Fd:180:GLY:O	9:Fd:184:ILE:HD12	1.93	0.68
9:Fd:932:LEU:O	9:Fd:936:LEU:HD12	1.93	0.68
6:Bd:112:PHE:O	6:Bd:116:THR:HG22	1.93	0.68
2:Br:2:VAL:HG22	3:Bs:1:ARG:HE	1.57	0.68
8:Ey:438:LEU:O	8:Ey:442:ILE:HG12	1.93	0.68
9:Fb:618:LEU:HB2	9:Ff:618:LEU:HD11	1.74	0.68
9:Fe:527:LEU:HA	9:Fe:530:ILE:HG12	1.75	0.68
9:Fe:502:THR:HG23	9:Fe:531:GLN:HE21	1.58	0.68
5:Aw:219:ARG:CZ	6:Bd:150:LYS:HD3	2.23	0.68

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
5:Dw:253:ARG:HH11	5:Dw:255:LEU:HD11	1.58	0.68
9:Fc:104:PRO:O	9:Fc:108:THR:HG23	1.93	0.68
9:Fd:334:ILE:HD12	9:Fd:429:ILE:HD11	1.73	0.68
6:Df:225:ASN:HB2	6:Dl:250:TYR:OH	1.94	0.68
9:Fd:928:LYS:HA	9:Fd:931:THR:HG23	1.75	0.68
5:Aw:222:LEU:HD11	5:Aw:262:ILE:HG21	1.76	0.68
6:Bj:110:LYS:HG3	7:Eh:794:ILE:HG21	1.75	0.68
6:Bp:122:LEU:HD11	6:Bp:130:LEU:HD11	1.76	0.68
6:Ct:126:GLN:HA	6:Ct:129:LYS:HG3	1.74	0.68
2:Dt:2:VAL:HG22	3:Du:1:ARG:HE	1.57	0.68
6:Dx:228:VAL:HG13	6:Dx:237:VAL:HB	1.76	0.68
8:Ew:418:LEU:O	8:Ew:422:ILE:HG13	1.93	0.68
9:Fc:102:TRP:HD1	9:Fc:792:ASN:HB2	1.59	0.68
9:Fd:61:PHE:HA	9:Fd:64:PHE:HB3	1.75	0.68
9:Fe:323:SER:HA	9:Fe:326:GLN:HE21	1.58	0.68
6:Ch:131:LYS:HA	7:Em:812:TRP:HZ3	1.58	0.68
6:Ch:196:LEU:HB2	6:Ch:226:LEU:HD13	1.74	0.68
9:Fc:129:MET:HE1	9:Fc:776:MET:HB3	1.76	0.68
9:Fc:760:ILE:HD12	9:Fc:930:PHE:CB	2.24	0.68
9:Fd:666:GLY:O	9:Fd:670:ILE:HG12	1.93	0.68
6:Cz:170:ARG:HD2	6:Cz:247:ALA:HB3	1.74	0.68
6:Dr:130:LEU:HD23	7:Es:808:LEU:HD12	1.76	0.68
9:Fc:702:THR:O	9:Fc:706:THR:HG23	1.94	0.68
9:Fe:605:MET:HE1	9:Fe:641:TYR:CD2	2.29	0.68
6:Ax:126:GLN:HA	6:Ax:129:LYS:HE2	1.75	0.68
8:Ey:427:TYR:O	8:Ey:430:ARG:HB2	1.94	0.68
9:Fc:698:ASN:HA	9:Fd:905:TRP:CH2	2.29	0.68
9:Fd:227:ARG:NH1	9:Fd:228:THR:HA	2.09	0.68
7:Fi:30:ILE:O	7:Fi:34:ILE:HG12	1.94	0.68
1:Dg:9:LEU:HD12	1:Dg:13:LYS:HE2	1.74	0.68
8:Ez:228:PRO:HG3	8:Ez:234:LYS:HG3	1.76	0.68
9:Fd:192:LEU:HB2	9:Fd:351:VAL:HG21	1.76	0.68
7:Fk:32:VAL:HG13	7:Fk:36:PHE:CE1	2.28	0.68
1:Da:16:ALA:HB3	2:Dh:9:ARG:HG3	1.76	0.67
6:Dr:236:PRO:HG2	6:Dr:238:MET:HE3	1.76	0.67
5:Dw:269:SER:HA	6:Dx:246:LYS:HE3	1.76	0.67
7:Ef:799:SER:O	7:Ef:803:THR:HG23	1.94	0.67
7:Es:808:LEU:HB3	7:Es:812:TRP:CZ2	2.29	0.67
9:Fc:648:PHE:CE2	9:Fc:720:PHE:HB3	2.29	0.67
9:Fd:78:TYR:HA	9:Fd:81:MET:HE2	1.74	0.67
7:Fg:30:ILE:O	7:Fg:34:ILE:HG13	1.93	0.67

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
6:Af:138:GLU:CD	6:Ed:220:LEU:HD21	2.20	0.67
5:Ca:246:LEU:HG	5:Ca:255:LEU:HB2	1.76	0.67
8:Ey:444:LEU:HD11	8:Ez:439:THR:HB	1.75	0.67
9:Fc:745:ILE:HG23	9:Fc:919:THR:HG21	1.74	0.67
6:Ar:234:ASN:HD22	6:Ax:188:PRO:HG3	1.59	0.67
6:Cn:128:VAL:HG23	6:Cn:132:GLN:HE21	1.60	0.67
9:Fe:124:MET:HE3	9:Fe:124:MET:O	1.93	0.67
6:Ax:169:ILE:HD13	6:Ax:241:LEU:HD23	1.76	0.67
6:Dl:125:GLU:HG2	6:Dl:126:GLN:OE1	1.94	0.67
6:Ed:117:ARG:HE	7:Eu:797:ARG:HE	1.42	0.67
7:Ek:793:GLU:O	7:Ek:797:ARG:HG2	1.94	0.67
8:Ew:444:LEU:HD22	8:Ex:443:MET:HB3	1.76	0.67
8:Fa:434:GLU:HA	8:Fa:437:LEU:HG	1.76	0.67
7:Fj:25:ALA:O	7:Fj:29:ILE:HG12	1.95	0.67
6:Af:106:VAL:HA	6:Af:109:LYS:HG2	1.76	0.67
6:Bj:189:TRP:HE1	6:Bj:232:GLY:H	1.39	0.67
7:Et:814:GLN:HE22	7:Et:816:GLU:HB3	1.58	0.67
8:Ew:399:TRP:NE1	8:Ew:403:ILE:HD11	2.09	0.67
9:Fd:395:VAL:HB	9:Fd:397:GLN:HE22	1.59	0.67
5:De:264:PHE:HD1	5:De:269:SER:HB3	1.59	0.67
9:Fb:657:MET:HA	9:Fb:660:LEU:HD22	1.75	0.67
9:Fb:808:VAL:HG12	9:Fb:811:ARG:HH22	1.58	0.67
6:Ax:198:ASP:HA	7:EI:816:GLU:HG3	1.76	0.67
2:Bl:4:MET:HE1	5:Bo:249:SER:N	2.09	0.67
1:Dm:12:HIS:CD2	3:Do:9:TYR:HB2	2.30	0.67
8:Ew:78:GLN:HE22	8:Ex:384:PHE:HB2	1.59	0.67
8:Ew:327:SER:CB	8:Ex:211:MET:HE1	2.24	0.67
8:Ey:352:SER:HA	8:Ey:355:LEU:HG	1.75	0.67
8:Ez:105:LYS:HE2	8:Fa:368:ILE:H	1.60	0.67
9:Fc:192:LEU:HB2	9:Fc:351:VAL:HG21	1.75	0.67
9:Ff:662:ILE:HG12	9:Ff:703:LEU:HB3	1.77	0.67
7:Eo:794:ILE:HD12	7:Eo:794:ILE:H	1.59	0.67
8:Fa:189:PHE:HA	8:Fa:295:LEU:HD21	1.76	0.67
8:Fa:441:SER:O	8:Fa:445:LEU:HD22	1.94	0.67
9:Fb:57:MET:O	9:Fb:60:MET:HG2	1.95	0.67
9:Fb:766:PHE:HA	9:Fb:769:LEU:HG	1.75	0.67
9:Fd:468:PHE:CG	9:Fd:757:PRO:HG2	2.30	0.67
9:Fd:819:TYR:CE1	9:Fd:823:ILE:HD11	2.30	0.67
9:Ff:815:MET:HB3	9:Ff:933:ILE:HG13	1.76	0.67
9:Ff:940:VAL:O	9:Ff:944:ILE:HG12	1.95	0.67
6:Ar:132:GLN:O	6:Ar:136:THR:HG23	1.95	0.67

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
6:Ar:219:LYS:HE3	6:Ar:222:ASN:ND2	2.09	0.67
3:Au:2:VAL:HG21	6:Bj:129:LYS:HD3	1.76	0.67
6:Ax:131:LYS:HE3	7:Eg:812:TRP:CH2	2.29	0.67
8:Ew:419:LEU:HD11	8:Ex:415:ILE:HG23	1.76	0.67
8:Ex:77:VAL:HG12	8:Ex:378:MET:HE1	1.76	0.67
8:Ex:245:SER:HA	8:Ex:279:PHE:CE2	2.30	0.67
8:Fa:206:TYR:CE2	8:Fa:208:ASN:HB2	2.26	0.67
9:Fc:604:TYR:HA	9:Fc:641:TYR:OH	1.95	0.67
9:Fd:714:SER:HB3	9:Fe:598:VAL:HG12	1.77	0.67
9:Fe:432:PRO:O	9:Fe:436:LEU:HD22	1.94	0.67
6:Bd:117:ARG:HE	7:Eh:797:ARG:HD2	1.59	0.67
7:En:794:ILE:H	7:En:794:ILE:HD12	1.60	0.67
8:Ex:91:VAL:HA	8:Ey:246:ASN:ND2	2.09	0.67
9:Fc:106:ARG:HG2	9:Fc:106:ARG:HH11	1.60	0.67
9:Fc:136:VAL:HG13	9:Fc:461:TRP:HE1	1.59	0.67
9:Fc:272:PHE:HB3	9:Fc:275:LEU:HD12	1.77	0.67
7:Fi:17:THR:O	7:Fi:21:ILE:HG12	1.95	0.67
6:Ax:115:MET:CE	7:Ef:801:MET:HB2	2.25	0.66
6:Ct:238:MET:HE1	6:Cz:251:ARG:HG3	1.76	0.66
8:Ew:111:LEU:HD23	8:Ex:386:PRO:HB3	1.75	0.66
9:Fb:651:MET:O	9:Fb:655:ILE:HG12	1.95	0.66
7:Fg:18:ARG:O	7:Fg:22:ILE:HG12	1.95	0.66
2:An:4:MET:HE1	5:Aq:249:SER:N	2.06	0.66
6:Ar:176:VAL:HA	6:Ar:216:GLN:OE1	1.95	0.66
8:Ex:56:GLN:HB2	8:Ex:59:LYS:HE3	1.77	0.66
8:Ex:144:GLN:HB2	8:Ex:179:PHE:CZ	2.30	0.66
9:Fb:205:TYR:HA	9:Fb:208:GLN:HE21	1.60	0.66
9:Fc:657:MET:O	9:Fc:661:MET:HB2	1.95	0.66
9:Fd:837:PHE:O	9:Fd:841:ILE:HG12	1.96	0.66
9:Fe:909:TYR:HD2	9:Fe:912:PHE:HE2	1.42	0.66
9:Ff:653:ASN:O	9:Ff:657:MET:HG2	1.96	0.66
9:Ff:770:ILE:HA	9:Ff:773:ILE:HD12	1.77	0.66
7:Fi:19:VAL:HA	7:Fi:22:ILE:HG12	1.77	0.66
5:Bi:216:ILE:HG12	5:Bi:221:TRP:HH2	1.60	0.66
6:Bv:108:ASP:HA	7:Ej:797:ARG:NH2	2.10	0.66
2:Cv:5:ILE:HD12	6:Df:139:TYR:HD2	1.59	0.66
6:Bd:219:LYS:HD2	6:Bd:222:ASN:HD22	1.60	0.66
4:Eb:2:PHE:HB3	5:Ec:251:GLN:HE21	1.60	0.66
8:Ex:418:LEU:O	8:Ex:421:GLU:HG2	1.95	0.66
9:Fb:132:ILE:O	9:Fb:136:VAL:HG23	1.96	0.66
9:Fc:928:LYS:HG3	9:Fc:929:ALA:N	2.08	0.66

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
9:Ff:685:PRO:HD2	9:Ff:827:TYR:HE1	1.60	0.66
6:Cb:110:LYS:HG3	7:Ek:797:ARG:NH2	2.11	0.66
6:Ct:205:GLN:HB3	7:Ep:820:TYR:CZ	2.30	0.66
6:Dr:194:TYR:HE2	6:Dr:196:LEU:HB2	1.59	0.66
7:Ej:806:THR:O	7:Ej:810:GLN:HG2	1.95	0.66
8:Ew:234:LYS:HA	8:Ew:237:GLN:HE21	1.60	0.66
8:Ey:433:GLN:HA	8:Ey:436:ILE:HG12	1.77	0.66
9:Fd:326:GLN:HA	9:Fd:329:ARG:HE	1.60	0.66
9:Fd:652:ILE:HA	9:Fd:655:ILE:HD12	1.77	0.66
9:Fc:693:GLY:HA2	9:Fc:696:TYR:CD2	2.30	0.66
1:Ag:12:HIS:CD2	3:Ai:9:TYR:HB2	2.30	0.66
6:Ar:169:ILE:HD11	6:Ar:239:LEU:HB3	1.77	0.66
6:Cn:208:LYS:HD2	7:Ep:823:GLY:HA2	1.78	0.66
6:Cz:145:PRO:HD3	6:Df:131:LYS:HE2	1.77	0.66
8:Ew:419:LEU:HB2	8:Ex:418:LEU:HD23	1.78	0.66
9:Fb:630:ASP:HA	9:Fb:634:ASN:HB2	1.76	0.66
9:Fc:572:ASN:HA	9:Fc:575:MET:HE3	1.77	0.66
9:Fc:753:ILE:HG23	9:Fc:926:VAL:HG21	1.76	0.66
6:Bp:115:MET:HE1	7:Ei:798:THR:HG23	1.78	0.66
6:Ct:171:LEU:HD12	6:Ct:172:SER:H	1.60	0.66
8:Fa:208:ASN:HA	8:Fa:211:MET:SD	2.35	0.66
9:Fb:188:GLN:HA	9:Fb:191:MET:HG2	1.78	0.66
9:Fb:818:GLY:HA3	9:Fb:936:LEU:HD11	1.78	0.66
9:Fc:180:GLY:HA3	9:Fc:491:LEU:HB3	1.78	0.66
9:Fd:103:ILE:HG13	9:Fd:104:PRO:HD3	1.78	0.66
9:Fe:266:PHE:H	9:Fe:276:ASN:HD21	1.43	0.66
9:Ff:566:THR:HG23	9:Ff:569:GLY:H	1.61	0.66
6:Bp:236:PRO:HG2	6:Bv:251:ARG:HH21	1.61	0.66
6:Bv:233:LEU:HD12	6:Bv:234:ASN:H	1.60	0.66
6:Ct:128:VAL:HG23	6:Ct:132:GLN:NE2	2.11	0.66
8:Ey:419:LEU:C	8:Ey:419:LEU:HD12	2.21	0.66
9:Fb:594:ILE:HG23	9:Ff:705:LEU:HD12	1.78	0.66
9:Fc:268:LYS:HZ3	9:Fd:404:SER:H	1.44	0.66
9:Ff:685:PRO:HG3	9:Ff:831:TRP:HE1	1.59	0.66
6:Bv:136:THR:HA	6:Bv:139:TYR:CD2	2.31	0.66
8:Ey:431:GLN:HE21	8:Ey:435:ARG:HD2	1.61	0.66
8:Ez:418:LEU:HD12	8:Ez:419:LEU:N	2.10	0.66
8:Fa:207:GLN:HB2	8:Fa:289:PRO:HG2	1.78	0.66
5:Dk:269:SER:HA	6:Dl:246:LYS:HZ2	1.60	0.65
7:Ei:805:ALA:O	7:Ei:809:VAL:HG23	1.95	0.65
8:Ez:430:ARG:HH22	8:Fa:425:GLN:HG3	1.61	0.65

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
9:Fb:242:LYS:HD2	9:Fb:261:LEU:HD23	1.79	0.65
9:Fd:917:ILE:HD12	9:Fd:920:SER:HB2	1.78	0.65
1:Co:6:LEU:HA	1:Co:9:LEU:HG	1.77	0.65
3:Cq:2:VAL:HG22	3:Cq:8:VAL:HG13	1.77	0.65
6:Ed:132:GLN:O	6:Ed:136:THR:HG23	1.95	0.65
7:Em:799:SER:O	7:Em:803:THR:HG23	1.97	0.65
7:Ev:792:GLN:HA	7:Ev:795:GLN:NE2	2.11	0.65
8:Ez:430:ARG:HH12	8:Fa:425:GLN:CG	2.08	0.65
9:Fc:814:LEU:HA	9:Fc:817:ILE:HD12	1.76	0.65
9:Fc:825:LEU:HD11	9:Fc:928:LYS:HE3	1.78	0.65
6:Cz:169:ILE:HG12	6:Cz:179:LEU:HD11	1.77	0.65
6:Df:121:PRO:HB2	7:Eq:797:ARG:NH1	2.11	0.65
8:Ex:419:LEU:HA	8:Ex:422:ILE:HG12	1.78	0.65
8:Ey:430:ARG:HH22	8:Ez:428:LEU:HB2	1.60	0.65
8:Ez:102:VAL:HG21	8:Ez:106:VAL:HG12	1.77	0.65
9:Fb:103:ILE:HD12	9:Fb:104:PRO:CD	2.25	0.65
9:Fb:828:VAL:HA	9:Fb:831:TRP:CE3	2.31	0.65
9:Fc:243:GLN:HE22	9:Fc:330:LEU:HD22	1.61	0.65
9:Fe:146:ALA:O	9:Fe:150:LEU:HD22	1.95	0.65
9:Fe:266:PHE:HE2	9:Fe:275:LEU:HB2	1.60	0.65
9:Ff:942:ARG:C	9:Ff:942:ARG:HE	2.03	0.65
6:Bd:181:PHE:HZ	6:Bd:213:LEU:HB2	1.62	0.65
2:Bf:4:MET:HE1	5:Bi:250:LEU:HG	1.78	0.65
6:Bv:219:LYS:HG3	6:Bv:222:ASN:HB2	1.79	0.65
6:Cb:208:LYS:HG3	7:En:824:THR:HG22	1.78	0.65
6:Cz:179:LEU:HD13	6:Cz:254:LEU:HD21	1.79	0.65
7:Ei:782:ASN:HA	9:Fd:241:LYS:HD2	1.78	0.65
8:Ez:334:ARG:HH22	8:Fa:215:ILE:HD13	1.61	0.65
9:Fd:723:PHE:HD1	9:Fd:726:ALA:HB3	1.62	0.65
9:Fe:837:PHE:CG	9:Fe:918:TYR:HE1	2.14	0.65
6:Bd:129:LYS:O	6:Bd:133:ILE:HG12	1.96	0.65
6:Bp:204:ILE:HG13	6:Bp:215:ILE:HD13	1.78	0.65
9:Fb:261:LEU:HD21	9:Fb:265:ASN:HD21	1.62	0.65
9:Fc:107:SER:HA	9:Fc:787:THR:HG22	1.78	0.65
9:Fc:756:LEU:HA	9:Fc:759:MET:HG2	1.79	0.65
5:Bo:250:LEU:HD12	5:Bo:251:GLN:HG3	1.79	0.65
6:Bp:206:TRP:HB2	6:Bp:213:LEU:HD23	1.79	0.65
6:Df:182:LEU:HD13	6:Df:186:GLY:HA2	1.78	0.65
2:Dh:4:MET:HE1	5:Dk:249:SER:H	1.60	0.65
8:Ey:136:ILE:HD13	8:Ey:277:LEU:HD23	1.79	0.65
8:Fa:68:LEU:HA	8:Fa:410:THR:HG23	1.79	0.65

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
9:Fb:537:VAL:HG13	9:Fb:542:GLN:HA	1.79	0.65
7:Fj:36:PHE:HA	7:Fj:39:ILE:HD12	1.79	0.65
5:Ae:213:GLN:HG3	6:Af:168:VAL:HG21	1.79	0.65
5:Ae:246:LEU:HD12	5:Ae:255:LEU:HB2	1.77	0.65
6:Ar:216:GLN:HE21	7:Eg:818:GLN:NE2	1.94	0.65
3:Ba:2:VAL:HG22	3:Ba:8:VAL:HG23	1.78	0.65
6:Bp:112:PHE:O	6:Bp:116:THR:HG23	1.95	0.65
6:Bv:111:ALA:HB3	7:Ej:797:ARG:NH2	2.11	0.65
7:Eg:807:GLN:HB3	7:Eg:810:GLN:HE21	1.61	0.65
8:Ew:420:ALA:O	8:Ew:424:TYR:HD1	1.79	0.65
8:Ey:178:VAL:HG13	8:Ez:240:ILE:HD11	1.79	0.65
9:Fd:260:GLU:HG2	9:Fd:283:LYS:HD2	1.79	0.65
9:Fe:758:TYR:O	9:Fe:762:THR:HG23	1.96	0.65
5:Ak:234:GLU:HA	5:Ak:247:ILE:HD11	1.79	0.65
6:Bd:208:LYS:NZ	7:Ej:821:THR:HG22	2.10	0.65
6:Ch:166:PRO:HB3	6:Cn:250:TYR:HE2	1.60	0.65
6:Cz:112:PHE:HA	6:Cz:115:MET:HE3	1.78	0.65
8:Ew:83:ASN:HA	8:Ew:115:LEU:HD21	1.79	0.65
5:Aq:216:ILE:HD11	6:Ax:148:PRO:HG2	1.77	0.65
6:Bv:126:GLN:HB3	6:Cb:112:PHE:CE2	2.31	0.65
6:Ed:203:ASN:ND2	7:Ev:818:GLN:HE22	1.95	0.65
8:Ew:331:ASN:O	8:Ew:335:VAL:HG22	1.96	0.65
8:Ey:253:LEU:HD22	8:Ey:373:LEU:HD22	1.79	0.65
9:Fb:939:LYS:HG3	9:Fb:940:VAL:N	2.11	0.65
9:Fc:741:THR:O	9:Fc:745:ILE:HG12	1.97	0.65
9:Fc:828:VAL:O	9:Fc:832:ILE:HG12	1.97	0.65
9:Fd:608:GLN:H	9:Fd:634:ASN:HD21	1.43	0.65
6:Bj:182:LEU:HG	6:Bj:255:ARG:HE	1.62	0.65
1:Cc:15:ASN:HB3	2:Cj:9:ARG:HH12	1.61	0.65
6:Dx:111:ALA:HB3	7:Es:797:ARG:NH2	2.12	0.65
8:Ey:36:THR:HA	8:Ey:39:LEU:HD23	1.79	0.65
8:Ey:422:ILE:HD13	8:Ey:425:GLN:HE21	1.61	0.65
8:Ez:206:TYR:CE1	8:Ez:208:ASN:HB2	2.32	0.65
8:Ez:424:TYR:CE2	8:Fa:383:LEU:HB2	2.32	0.65
8:Fa:381:ARG:HG3	8:Fa:382:ARG:HE	1.61	0.65
9:Fc:57:MET:HB3	9:Fc:61:PHE:CE1	2.31	0.65
9:Fc:461:TRP:CD1	9:Fc:461:TRP:H	2.15	0.65
9:Fc:670:ILE:HD12	9:Fc:696:TYR:HD1	1.61	0.65
9:Fe:612:CYS:HB2	9:Fe:626:ARG:HE	1.61	0.65
4:Ad:3:GLY:HA3	1:Dy:16:ALA:HA	1.78	0.64
6:Bv:121:PRO:HB2	7:Ek:801:MET:HB3	1.78	0.64

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
6:Dr:204:ILE:HG13	6:Dr:215:ILE:HD13	1.77	0.64
9:Fc:745:ILE:O	9:Fc:749:THR:HG22	1.97	0.64
9:Fd:905:TRP:CE3	9:Fd:909:TYR:HE1	2.16	0.64
3:Bm:11:ALA:HB1	3:Bm:14:TYR:HE2	1.61	0.64
6:Ct:206:TRP:HB2	6:Ct:213:LEU:HD23	1.80	0.64
7:Ej:771:ALA:O	7:Ej:775:GLN:HG2	1.97	0.64
8:Ex:352:SER:HA	8:Ex:355:LEU:HG	1.79	0.64
8:Ex:424:TYR:O	8:Ex:428:LEU:HD12	1.97	0.64
8:Fa:248:LEU:HD23	8:Fa:279:PHE:HD2	1.61	0.64
9:Fc:673:VAL:O	9:Fc:677:THR:HG23	1.97	0.64
9:Ff:57:MET:HB3	9:Ff:61:PHE:HE1	1.61	0.64
7:Fj:33:VAL:HA	7:Fj:36:PHE:CD1	2.32	0.64
5:Bi:218:GLY:HA3	6:Bp:148:PRO:HD3	1.80	0.64
5:Bo:219:ARG:HG2	6:Bv:148:PRO:HD2	1.77	0.64
6:Cn:189:TRP:CD1	6:Cn:230:LEU:HD11	2.32	0.64
6:Ct:111:ALA:HB3	7:En:797:ARG:CZ	2.27	0.64
2:Dn:2:VAL:HA	3:Do:1:ARG:HE	1.61	0.64
8:Ey:211:MET:HA	8:Ey:214:VAL:HG23	1.80	0.64
5:Ae:219:ARG:HG2	6:Al:148:PRO:HD2	1.78	0.64
6:Af:106:VAL:HG23	6:Af:110:LYS:NZ	2.12	0.64
5:Bi:219:ARG:HH21	5:Bi:231:THR:HG23	1.62	0.64
6:Ch:205:GLN:HB2	6:Ch:214:MET:SD	2.36	0.64
6:Cz:105:GLU:OE2	6:Cz:109:LYS:HD3	1.96	0.64
8:Ew:427:TYR:CD1	8:Ew:428:LEU:HD13	2.32	0.64
8:Ez:428:LEU:HA	8:Ez:431:GLN:HG2	1.79	0.64
9:Fb:33:LEU:HB2	9:Fb:462:ILE:HD11	1.80	0.64
6:Bd:206:TRP:HB2	6:Bd:213:LEU:HD23	1.79	0.64
6:Bj:179:LEU:HB3	6:Bj:254:LEU:HD11	1.80	0.64
5:Bu:216:ILE:HD11	6:Cb:148:PRO:HG2	1.78	0.64
8:Ew:43:LEU:HD11	8:Ex:42:TYR:CD2	2.32	0.64
8:Ew:211:MET:HA	8:Ew:214:VAL:HG23	1.78	0.64
8:Fa:352:SER:HA	8:Fa:355:LEU:HG	1.79	0.64
9:Fd:696:TYR:CZ	9:Fd:742:MET:HA	2.33	0.64
6:Bv:123:ASN:H	6:Bv:126:GLN:NE2	1.96	0.64
1:Cu:16:ALA:HA	4:Dd:3:GLY:HA3	1.80	0.64
6:Dr:107:ILE:O	6:Dr:111:ALA:HB2	1.97	0.64
7:Ef:794:ILE:O	7:Ef:798:THR:HG23	1.97	0.64
8:Ew:380:THR:HA	8:Fa:424:TYR:HE2	1.60	0.64
8:Ex:331:ASN:HD21	8:Ey:211:MET:HB2	1.63	0.64
8:Ey:330:PHE:HB2	8:Ey:334:ARG:NH2	2.13	0.64
8:Ey:336:TYR:O	8:Ey:340:THR:HG23	1.96	0.64

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
8:Ey:423:ASN:HD22	8:Ez:382:ARG:NH2	1.94	0.64
8:Ez:417:ILE:O	8:Ez:420:ALA:HB3	1.98	0.64
9:Fc:136:VAL:HG22	9:Fc:461:TRP:HZ2	1.62	0.64
9:Fc:930:PHE:HA	9:Fc:933:ILE:HD12	1.78	0.64
9:Ff:820:ILE:HA	9:Ff:823:ILE:HD12	1.80	0.64
6:Af:190:PRO:O	6:Af:231:ARG:HG2	1.98	0.64
6:Bp:126:GLN:HA	6:Bp:129:LYS:HE2	1.79	0.64
3:Cq:2:VAL:HG11	6:Df:129:LYS:HE3	1.78	0.64
6:Ct:117:ARG:NH1	6:Ct:121:PRO:HB3	2.13	0.64
8:Ex:136:ILE:HD11	8:Ex:273:ALA:HB1	1.80	0.64
8:Ex:141:LEU:HB3	8:Ex:213:ASN:HB3	1.79	0.64
8:Ey:330:PHE:HA	8:Ey:333:LEU:HG	1.78	0.64
9:Ff:140:ASP:O	9:Ff:144:GLU:HG2	1.97	0.64
9:Ff:330:LEU:O	9:Ff:334:ILE:HG12	1.97	0.64
9:Ff:921:MET:O	9:Ff:925:ILE:HB	1.98	0.64
6:Bv:150:LYS:HG2	6:Bv:152:THR:HG23	1.80	0.64
6:Df:249:ASP:HB3	6:Df:252:VAL:HG23	1.79	0.64
5:Dq:214:ALA:HB3	5:Dq:221:TRP:CD1	2.32	0.64
6:Dx:112:PHE:HA	6:Dx:115:MET:SD	2.38	0.64
8:Ey:437:LEU:HD11	8:Ez:435:ARG:HB3	1.78	0.64
9:Fb:773:ILE:O	9:Fb:777:VAL:HG23	1.97	0.64
9:Fd:124:MET:HE1	9:Fd:780:PRO:HA	1.79	0.64
9:Fd:690:ALA:O	9:Fd:694:THR:HG23	1.97	0.64
9:Ff:259:PHE:HE2	9:Ff:286:ASN:HB2	1.61	0.64
6:Al:114:ASP:HA	6:Al:117:ARG:HG2	1.80	0.64
6:Al:203:ASN:HD21	7:Ef:817:THR:HG23	1.63	0.64
6:Dl:173:GLN:HB2	6:Dl:220:LEU:HA	1.78	0.64
6:Dr:201:SER:C	6:Dr:202:PHE:HD2	2.06	0.64
9:Fb:266:PHE:HE2	9:Fb:275:LEU:HB2	1.62	0.64
9:Fe:49:LEU:HD13	9:Fe:51:GLY:H	1.63	0.64
9:Fe:605:MET:HE3	9:Fe:605:MET:H	1.63	0.64
6:Bv:204:ILE:HD11	6:Bv:213:LEU:HD22	1.80	0.64
6:Ch:131:LYS:HA	7:Em:812:TRP:CZ3	2.32	0.64
6:Cn:129:LYS:HE2	6:Ct:112:PHE:CE2	2.33	0.64
6:Dl:109:LYS:HB2	6:Dl:113:LYS:HZ3	1.63	0.64
6:Dl:205:GLN:HB2	6:Dl:214:MET:HE3	1.79	0.64
8:Ez:208:ASN:HA	8:Ez:211:MET:SD	2.38	0.64
8:Fa:77:VAL:HG12	8:Fa:378:MET:HE1	1.80	0.64
8:Fa:399:TRP:HH2	8:Fa:418:LEU:HD22	1.63	0.64
9:Fb:520:PHE:HE2	9:Fb:526:LYS:HB2	1.63	0.64
9:Fc:841:ILE:HD12	9:Fc:844:ILE:HB	1.79	0.64

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
6:Cz:106:VAL:HG23	6:Cz:110:LYS:NZ	2.14	0.63
2:Dt:5:ILE:HD12	6:Ed:139:TYR:CD2	2.33	0.63
6:Ed:168:VAL:HG13	6:Ed:242:ILE:HD13	1.80	0.63
7:Em:808:LEU:O	7:Em:812:TRP:HD1	1.81	0.63
7:Er:809:VAL:HA	7:Er:812:TRP:CD1	2.32	0.63
8:Ez:174:VAL:O	8:Ez:178:VAL:HG23	1.98	0.63
8:Ez:317:VAL:HG13	8:Fa:198:TRP:HE1	1.63	0.63
8:Ez:418:LEU:HA	8:Ez:421:GLU:OE2	1.97	0.63
9:Fd:664:LEU:O	9:Fd:668:LYS:HB3	1.98	0.63
9:Fe:655:ILE:HG23	9:Fe:659:PHE:CE2	2.33	0.63
9:Fe:663:PRO:O	9:Fe:667:MET:CB	2.41	0.63
9:Ff:665:GLN:HA	9:Ff:668:LYS:HG2	1.80	0.63
7:Fi:16:ARG:O	7:Fi:20:ILE:HG12	1.97	0.63
2:Db:4:MET:HE1	5:De:249:SER:N	2.13	0.63
6:Df:107:ILE:HG23	7:Ep:794:ILE:HG22	1.80	0.63
6:Dx:202:PHE:HB3	6:Dx:215:ILE:HD11	1.80	0.63
8:Ex:441:SER:O	8:Ex:445:LEU:HD22	1.97	0.63
8:Ey:142:ILE:HA	8:Ey:210:VAL:HG23	1.80	0.63
8:Ez:435:ARG:O	8:Ez:439:THR:HG23	1.98	0.63
9:Ff:905:TRP:HB2	9:Ff:909:TYR:HE1	1.62	0.63
6:Ar:169:ILE:HD13	6:Ar:241:LEU:HD13	1.80	0.63
5:Bi:243:MET:HE2	5:Bi:243:MET:HA	1.79	0.63
6:Bp:115:MET:HG3	7:Bi:801:MET:SD	2.38	0.63
6:Cn:234:ASN:HD21	6:Ct:182:LEU:HD21	1.64	0.63
6:Df:225:ASN:HD21	6:Dl:176:VAL:N	1.96	0.63
5:Dq:245:LYS:HB2	5:Dq:246:LEU:HD12	1.79	0.63
6:Dr:122:LEU:HB3	6:Dr:127:VAL:HG12	1.79	0.63
7:El:814:GLN:HE22	7:El:816:GLU:HB3	1.63	0.63
8:Ew:427:TYR:HA	8:Ew:430:ARG:HG3	1.80	0.63
8:Ey:347:LEU:HA	8:Ey:350:ILE:HD12	1.80	0.63
9:Fb:196:LYS:HG2	9:Fb:357:PHE:HB3	1.80	0.63
9:Fb:723:PHE:HZ	9:Ff:724:ILE:HD11	1.63	0.63
9:Fb:768:TRP:HA	9:Fb:811:ARG:HE	1.63	0.63
9:Fc:767:ALA:HA	9:Fc:770:ILE:HD12	1.80	0.63
9:Fd:616:LYS:HD2	9:Fd:621:SER:HB3	1.80	0.63
9:Fe:37:PHE:CE2	9:Ff:832:ILE:HD13	2.33	0.63
9:Ff:663:PRO:O	9:Ff:667:MET:HG3	1.98	0.63
5:Aw:219:ARG:NH1	5:Aw:233:ARG:HB3	2.13	0.63
2:Az:5:ILE:HD11	6:Bj:139:TYR:HE2	1.63	0.63
6:Bd:221:TYR:CE1	6:Bj:139:TYR:HB2	2.33	0.63
6:Bv:173:GLN:HE22	6:Bv:218:THR:C	2.06	0.63

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
6:Cb:115:MET:HE1	7:Ek:798:THR:O	1.99	0.63
8:Ew:340:THR:HG22	8:Fa:448:LEU:HD13	1.79	0.63
8:Ew:444:LEU:O	8:Ew:448:LEU:HG	1.99	0.63
8:Ez:427:TYR:O	8:Ez:430:ARG:HG2	1.98	0.63
9:Fd:752:TYR:HE1	9:Fd:927:GLN:HE22	1.47	0.63
9:Fe:180:GLY:HA3	9:Fe:491:LEU:HB3	1.81	0.63
9:Fe:548:PRO:HD3	9:Fe:568:TYR:HD2	1.63	0.63
9:Ff:756:LEU:O	9:Ff:759:MET:HB3	1.97	0.63
7:Fh:29:ILE:O	7:Fh:33:VAL:HG13	1.97	0.63
7:Fk:23:PHE:O	7:Fk:27:LEU:HD22	1.98	0.63
6:Af:181:PHE:HZ	6:Af:213:LEU:HD12	1.63	0.63
6:Bj:225:ASN:HD21	6:Bp:176:VAL:N	1.97	0.63
6:Bj:228:VAL:HG13	6:Bj:237:VAL:HB	1.79	0.63
6:Df:129:LYS:HE2	6:Df:132:GLN:HE21	1.64	0.63
6:Dl:181:PHE:HE2	6:Dl:191:ILE:HD11	1.64	0.63
6:Dr:121:PRO:HB2	7:Es:801:MET:HB3	1.81	0.63
8:Ez:129:SER:HB2	8:Ez:137:THR:HG21	1.80	0.63
8:Ez:430:ARG:HB3	8:Fa:429:ASP:OD1	1.99	0.63
9:Fb:644:PHE:HB3	9:Fb:720:PHE:CE2	2.34	0.63
9:Fd:624:LEU:HD23	9:Fd:628:MET:HE3	1.80	0.63
9:Fd:636:VAL:O	9:Fd:640:VAL:HG12	1.99	0.63
9:Fe:652:ILE:O	9:Fe:656:VAL:HG22	1.98	0.63
9:Ff:104:PRO:O	9:Ff:108:THR:HG23	1.98	0.63
9:Ff:110:GLY:O	9:Ff:114:LEU:HD12	1.98	0.63
9:Ff:420:LEU:HD11	9:Ff:577:VAL:HG23	1.81	0.63
9:Ff:773:ILE:O	9:Ff:777:VAL:HG23	1.99	0.63
7:Fg:34:ILE:HA	7:Fg:38:LYS:NZ	2.13	0.63
5:Bo:219:ARG:HA	5:Bo:232:VAL:O	1.97	0.63
6:Ch:131:LYS:HG2	7:Em:812:TRP:HH2	1.63	0.63
5:De:213:GLN:HG3	6:Df:168:VAL:HG21	1.81	0.63
8:Ex:75:GLN:HE21	8:Ex:79:ASN:HD21	1.46	0.63
8:Ez:298:TYR:HA	8:Ez:301:LEU:HD11	1.81	0.63
9:Fd:640:VAL:HG22	9:Fd:644:PHE:CZ	2.33	0.63
6:Al:196:LEU:HD13	6:Al:199:PRO:HB3	1.81	0.63
6:Bp:126:GLN:O	6:Bp:130:LEU:HD12	1.98	0.63
5:Cm:213:GLN:HB2	5:Cm:223:ILE:HG22	1.81	0.63
6:Cn:189:TRP:CG	6:Cn:230:LEU:HD11	2.34	0.63
6:Ct:111:ALA:HB2	7:En:794:ILE:HG23	1.80	0.63
7:Eh:794:ILE:H	7:Eh:794:ILE:HD12	1.64	0.63
7:Et:796:GLN:NE2	9:Ff:228:THR:HA	2.14	0.63
8:Ew:78:GLN:HE21	8:Ex:383:LEU:HG	1.64	0.63

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
8:Ey:271:SER:HB2	8:Ey:274:GLN:HG2	1.80	0.63
8:Ey:435:ARG:O	8:Ey:439:THR:HG23	1.99	0.63
5:Bo:216:ILE:HD11	6:Bv:148:PRO:HG2	1.80	0.63
6:Bp:144:THR:OG1	6:Bp:148:PRO:HG3	1.99	0.63
6:Bv:137:SER:HB3	6:Cb:120:TYR:CZ	2.34	0.63
6:Cz:225:ASN:ND2	6:Df:176:VAL:H	1.97	0.63
7: Ei:778:LEU:HD21	9:Fd:240:VAL:HB	1.81	0.63
8:Ex:302:TRP:CZ3	8:Ey:218:LEU:HG	2.34	0.63
8:Ey:343:GLY:HA2	8:Ey:346:ASN:HD21	1.63	0.63
8:Fa:209:LEU:CA	8:Fa:212:GLN:HE21	2.08	0.63
9:Fb:636:VAL:O	9:Fb:640:VAL:HG12	1.97	0.63
9:Fb:699:PHE:CD2	9:Fb:742:MET:HE2	2.33	0.63
9:Fc:554:PRO:HG2	9:Fc:588:LEU:HD21	1.79	0.63
6:Bv:147:THR:HG23	6:Bv:246:LYS:NZ	2.14	0.63
6:Ct:169:ILE:HD12	6:Ct:252:VAL:HG21	1.79	0.63
6:Dl:169:ILE:HG23	6:Dl:241:LEU:HA	1.81	0.63
6:Ed:115:MET:HE2	7:Et:801:MET:HE3	1.81	0.63
7:Et:781:GLN:NE2	9:Ff:241:LYS:HB2	2.14	0.63
8:Ex:78:GLN:OE1	8:Ey:383:LEU:HG	1.99	0.63
8:Fa:381:ARG:HG3	8:Fa:382:ARG:NE	2.14	0.63
8:Fa:435:ARG:O	8:Fa:439:THR:HG23	1.99	0.63
9:Fb:673:VAL:O	9:Fb:677:THR:HG23	1.98	0.63
9:Fb:813:SER:O	9:Fb:817:ILE:HG12	1.99	0.63
9:Fd:648:PHE:O	9:Fd:652:ILE:HG22	1.99	0.63
6:Ax:149:PRO:HB2	6:Ax:248:VAL:HG12	1.81	0.62
6:Bv:114:ASP:HB2	6:Bv:117:ARG:NH2	2.14	0.62
5:Dk:237:LYS:H	5:Dq:266:GLN:HE22	1.46	0.62
6:Dl:170:ARG:HE	6:Dl:249:ASP:HB2	1.63	0.62
8:Ez:277:LEU:HD21	8:Ez:281:ARG:HH21	1.64	0.62
9:Fb:295:GLN:NE2	9:Fb:303:THR:H	1.95	0.62
9:Fb:548:PRO:HD3	9:Fb:568:TYR:CE1	2.34	0.62
9:Fc:125:GLN:O	9:Fc:129:MET:HG2	1.99	0.62
9:Fe:566:THR:HG23	9:Fe:569:GLY:H	1.64	0.62
9:Fe:651:MET:O	9:Fe:655:ILE:HD13	1.99	0.62
9:Ff:773:ILE:HA	9:Ff:776:MET:HG3	1.81	0.62
7:Fj:30:ILE:O	7:Fj:34:ILE:HG12	1.98	0.62
6:Al:150:LYS:HG2	6:Al:152:THR:HG23	1.80	0.62
6:Bd:204:ILE:HD11	6:Bd:213:LEU:HD22	1.81	0.62
2:Bl:2:VAL:HA	3:Bm:1:ARG:HE	1.63	0.62
6:Ch:206:TRP:HB2	6:Ch:213:LEU:HD23	1.81	0.62
8:Ey:448:LEU:HB2	8:Ez:336:TYR:HE1	1.65	0.62

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
8:Fa:214:VAL:HA	8:Fa:267:LEU:HD23	1.80	0.62
9:Fb:190:CYS:O	9:Fb:194:LEU:HB3	1.99	0.62
9:Fb:507:LYS:HE2	9:Fc:630:ASP:HB2	1.80	0.62
9:Fc:612:CYS:HB3	9:Fc:626:ARG:H	1.64	0.62
9:Fd:920:SER:O	9:Fd:924:ILE:HD12	1.99	0.62
5:Ak:248:ASP:HB3	5:Ak:253:ARG:HB3	1.81	0.62
6:Ar:179:LEU:HD22	6:Ar:252:VAL:HG12	1.81	0.62
6:Bv:145:PRO:HG3	6:Cb:131:LYS:HG2	1.81	0.62
6:Bv:228:VAL:HG13	6:Bv:237:VAL:HB	1.80	0.62
3:Cw:2:VAL:HG22	3:Cw:8:VAL:HG12	1.82	0.62
5:Dk:246:LEU:HD22	5:Dk:255:LEU:HD23	1.81	0.62
6:Dr:219:LYS:HD2	6:Dr:220:LEU:N	2.14	0.62
8:Fa:74:VAL:O	8:Fa:78:GLN:HB3	1.99	0.62
9:Fb:647:ILE:HG13	9:Fb:717:ILE:HD12	1.81	0.62
9:Fc:56:ILE:O	9:Fc:60:MET:HG2	2.00	0.62
9:Fc:643:PHE:CE1	9:Fc:647:ILE:HD11	2.35	0.62
9:Fc:716:LEU:HB3	9:Fd:723:PHE:HD2	1.64	0.62
9:Ff:693:GLY:HA3	9:Ff:750:ALA:HB2	1.80	0.62
6:Al:206:TRP:HB2	6:Al:213:LEU:HD23	1.80	0.62
6:Bp:107:ILE:HD11	7:Ei:794:ILE:HG13	1.81	0.62
5:Cs:210:TYR:HA	5:Cs:224:GLY:HA2	1.79	0.62
8:Ew:435:ARG:O	8:Ew:439:THR:HG23	2.00	0.62
9:Fb:527:LEU:HA	9:Fb:530:ILE:HD13	1.81	0.62
9:Fb:663:PRO:HA	9:Fb:699:PHE:HZ	1.65	0.62
9:Fd:501:LEU:H	9:Fd:501:LEU:HD12	1.64	0.62
9:Fd:598:VAL:HG21	9:Fd:652:ILE:HD13	1.82	0.62
9:Fd:806:VAL:O	9:Fd:810:LEU:HG	2.00	0.62
6:Al:169:ILE:HD11	6:Al:239:LEU:HD12	1.81	0.62
8:Ey:178:VAL:HA	8:Ey:181:ILE:HG12	1.82	0.62
9:Fe:323:SER:HA	9:Fe:326:GLN:NE2	2.14	0.62
9:Fe:704:TRP:HA	9:Fe:707:LEU:HD12	1.81	0.62
9:Ff:643:PHE:O	9:Ff:647:ILE:HG12	1.99	0.62
6:Cn:141:LYS:HE3	7:Eo:812:TRP:CE2	2.35	0.62
6:Dx:115:MET:HA	6:Dx:118:ASN:OD1	2.00	0.62
8:Ew:241:SER:HB2	8:Fa:94:MET:HE1	1.81	0.62
8:Ez:89:ILE:HG22	8:Ez:349:TYR:HE1	1.63	0.62
9:Fb:434:LEU:HA	9:Fb:437:ILE:HG22	1.81	0.62
9:Fd:139:ALA:HB2	9:Fd:812:PRO:HB2	1.81	0.62
9:Fd:909:TYR:HA	9:Fd:912:PHE:CD2	2.35	0.62
9:Fe:674:GLY:HA2	9:Fe:692:MET:HE2	1.81	0.62
6:Al:218:THR:HA	7:Ef:815:VAL:HG11	1.81	0.62

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
6:Ar:121:PRO:HB2	7:Ef:801:MET:HE1	1.82	0.62
6:Ar:134:TYR:O	6:Ar:138:GLU:HG3	2.00	0.62
6:Cn:150:LYS:HG2	6:Cn:152:THR:HG23	1.81	0.62
5:De:238:ILE:HD12	5:De:239:PRO:HD2	1.82	0.62
6:Dx:136:THR:HA	6:Dx:139:TYR:CD1	2.34	0.62
8:Ew:283:ALA:HA	8:Fa:339:GLN:HE22	1.63	0.62
8:Ex:428:LEU:O	8:Ex:432:ILE:HG13	1.99	0.62
8:Ez:382:ARG:O	8:Ez:399:TRP:HB3	1.99	0.62
9:Fb:57:MET:HA	9:Fb:60:MET:SD	2.39	0.62
9:Fb:770:ILE:HA	9:Fb:773:ILE:HD12	1.82	0.62
9:Fe:383:LYS:HD2	9:Fe:387:GLU:HB2	1.80	0.62
9:Ff:612:CYS:HB2	9:Ff:626:ARG:HE	1.64	0.62
5:Cm:210:TYR:CG	5:Cm:222:LEU:HD21	2.35	0.62
7:Ee:804:ALA:O	7:Ee:808:LEU:HD22	1.99	0.62
7:En:799:SER:O	7:En:803:THR:HG23	2.00	0.62
7:Es:799:SER:O	7:Es:803:THR:HG23	2.00	0.62
8:Ez:412:GLN:HA	8:Ez:415:ILE:HG12	1.79	0.62
9:Fb:180:GLY:HA3	9:Fb:491:LEU:HB3	1.80	0.62
9:Fd:69:LEU:HD11	7:Fk:31:ALA:HB1	1.82	0.62
7:Fh:17:THR:HG22	7:Fh:21:ILE:HD11	1.82	0.62
7:Fk:34:ILE:O	7:Fk:38:LYS:HG2	1.99	0.62
6:Al:129:LYS:HG3	6:Ar:112:PHE:CZ	2.35	0.62
7:Ei:799:SER:O	7:Ei:803:THR:HG23	2.00	0.62
7:Ej:799:SER:HA	7:Ej:802:LEU:HD23	1.81	0.62
8:Ew:281:ARG:HB3	8:Ew:286:GLN:NE2	2.14	0.62
8:Ex:432:ILE:O	8:Ex:436:ILE:HD12	1.99	0.62
9:Fc:656:VAL:HG12	9:Fc:660:LEU:HD23	1.81	0.62
9:Ff:758:TYR:O	9:Ff:762:THR:HG23	1.99	0.62
9:Ff:402:VAL:HG12	9:Ff:404:SER:H	1.64	0.62
3:Au:2:VAL:HG22	3:Au:8:VAL:HG12	1.81	0.62
1:Ay:9:LEU:O	1:Ay:13:LYS:HG2	2.00	0.62
6:Bp:219:LYS:HE3	6:Bp:222:ASN:H	1.65	0.62
1:Bq:16:ALA:HA	4:Bz:3:GLY:HA3	1.82	0.62
6:Cz:112:PHE:O	6:Cz:116:THR:HG23	2.00	0.62
6:Cz:138:GLU:HA	6:Cz:141:LYS:HG2	1.82	0.62
6:Cz:150:LYS:N	6:Cz:247:ALA:HA	2.08	0.62
6:Dr:229:ARG:HH12	6:Dr:233:LEU:HB2	1.64	0.62
5:Ec:253:ARG:HE	5:Ec:263:LYS:HG3	1.65	0.62
6:Ed:153:ALA:HB1	6:Ed:251:ARG:HH21	1.65	0.62
7:Em:798:THR:O	7:Em:802:LEU:HD22	1.99	0.62
8:Ey:254:MET:HE2	8:Ey:254:MET:N	2.15	0.62

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
8:Fa:84:THR:HB	8:Fa:371:GLN:HA	1.82	0.62
9:Fd:188:GLN:HA	9:Fd:191:MET:HG2	1.82	0.62
9:Fd:493:LYS:HA	9:Fd:493:LYS:HE3	1.81	0.62
9:Ff:124:MET:HA	9:Ff:127:PHE:HD2	1.65	0.62
9:Ff:425:ASP:O	9:Ff:429:ILE:HG13	1.99	0.62
7:Fg:38:LYS:HD3	7:Fg:38:LYS:N	2.15	0.62
6:Bj:128:VAL:HA	6:Bj:131:LYS:HG2	1.82	0.61
6:Ct:195:ASP:OD2	7:Eq:820:TYR:HA	2.00	0.61
6:Cz:205:GLN:HB2	6:Cz:214:MET:SD	2.39	0.61
6:Dx:108:ASP:HA	7:Es:797:ARG:NH2	2.11	0.61
7:Ej:770:SER:HA	7:Ej:773:GLN:HE22	1.65	0.61
8:Ex:196:LYS:HZ2	9:Fc:219:THR:HB	1.65	0.61
8:Ez:430:ARG:NH1	8:Fa:425:GLN:HG3	2.14	0.61
9:Fc:56:ILE:HA	9:Fd:149:TYR:CE2	2.34	0.61
9:Fe:143:TRP:CZ3	9:Fe:147:LEU:HD21	2.35	0.61
9:Fe:673:VAL:O	9:Fe:677:THR:HG23	2.00	0.61
9:Ff:194:LEU:HA	9:Ff:197:GLN:HE21	1.65	0.61
5:Ca:241:TYR:HB3	5:Ca:256:THR:HG21	1.80	0.61
5:Dk:264:PHE:HD2	5:Dk:265:SER:H	1.47	0.61
8:Ew:104:ASP:HA	8:Ew:109:ASN:HD22	1.65	0.61
8:Ez:208:ASN:O	8:Ez:212:GLN:HG2	2.00	0.61
9:Fb:534:ILE:O	9:Fb:568:TYR:HB2	2.00	0.61
9:Fe:745:ILE:O	9:Fe:749:THR:HG22	2.01	0.61
9:Ff:294:ASN:HB2	9:Ff:313:MET:HE2	1.81	0.61
6:Ar:107:ILE:H	6:Ar:107:ILE:HD12	1.64	0.61
5:Bu:219:ARG:HG2	6:Cb:148:PRO:HD2	1.83	0.61
6:Bv:205:GLN:HB2	7:El:820:TYR:CZ	2.35	0.61
6:Cb:132:GLN:HE22	6:Cb:135:GLU:HG3	1.65	0.61
6:Ed:173:GLN:HB2	6:Ed:220:LEU:HB2	1.82	0.61
8:Ew:334:ARG:HG3	8:Ex:282:TYR:CE1	2.26	0.61
8:Ew:457:PHE:HZ	8:Ex:329:TYR:HB2	1.64	0.61
8:Ez:454:THR:HG23	8:Ez:456:ASP:H	1.65	0.61
9:Fb:136:VAL:HG13	9:Fb:461:TRP:HE1	1.65	0.61
9:Fd:122:CYS:HB2	9:Fd:124:MET:HG2	1.81	0.61
9:Fd:702:THR:O	9:Fd:706:THR:HG23	2.00	0.61
9:Ff:242:LYS:HE3	9:Ff:265:ASN:ND2	2.16	0.61
9:Ff:684:ASN:HB3	9:Ff:687:VAL:HG22	1.82	0.61
6:Af:138:GLU:HA	6:Af:141:LYS:HG2	1.83	0.61
6:Bv:151:PRO:HA	6:Bv:248:VAL:HG13	1.82	0.61
6:Cn:206:TRP:HB2	6:Cn:213:LEU:HD23	1.83	0.61
5:De:221:TRP:CD1	5:De:231:THR:HG1	2.19	0.61

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
5:De:247:ILE:HG12	5:De:254:ILE:HD12	1.82	0.61
6:Df:157:PHE:HD1	6:Df:255:ARG:HG3	1.64	0.61
6:Dx:204:ILE:HD13	6:Dx:215:ILE:HD13	1.83	0.61
6:Ed:189:TRP:CD1	6:Ed:230:LEU:HB3	2.35	0.61
8:Ew:96:GLN:HA	8:Ew:99:MET:HE1	1.82	0.61
8:Ex:276:ALA:O	8:Ex:280:ILE:HG23	2.00	0.61
8:Ez:191:MET:HE2	8:Ez:191:MET:HA	1.81	0.61
9:Fb:367:LYS:O	9:Fb:367:LYS:HD2	2.00	0.61
9:Fd:651:MET:O	9:Fd:655:ILE:HG13	2.00	0.61
7:Fi:19:VAL:HB	7:Fi:23:PHE:CZ	2.35	0.61
6:Bj:170:ARG:HB2	6:Bj:249:ASP:H	1.65	0.61
6:Cb:150:LYS:HG2	6:Cb:152:THR:HG23	1.83	0.61
5:Cg:226:ASN:HD21	5:Cg:228:SER:HB3	1.64	0.61
6:Cz:220:LEU:H	6:Cz:220:LEU:HD12	1.66	0.61
5:De:246:LEU:HB2	5:De:255:LEU:HD23	1.81	0.61
7:Es:780:LYS:HZ3	7:Es:783:GLU:HB2	1.65	0.61
8:Ey:75:GLN:O	8:Ey:78:GLN:HG3	2.00	0.61
8:Ez:123:GLN:HE22	8:Ez:256:THR:HG21	1.66	0.61
8:Fa:374:ASN:OD1	8:Fa:378:MET:HE2	2.01	0.61
9:Fb:909:TYR:HA	9:Fb:912:PHE:CD2	2.36	0.61
9:Fd:931:THR:O	9:Fd:934:ALA:HB3	2.00	0.61
9:Ff:325:LEU:O	9:Ff:329:ARG:HG2	2.01	0.61
6:Af:191:ILE:HD13	6:Af:213:LEU:HG	1.82	0.61
6:Cb:201:SER:HB2	6:Cb:219:LYS:HD2	1.82	0.61
6:Cn:106:VAL:HA	6:Cn:109:LYS:HZ2	1.65	0.61
6:Cn:115:MET:CE	7:Em:801:MET:HB2	2.30	0.61
2:Cp:4:MET:H	2:Cp:4:MET:HE3	1.66	0.61
7:Es:777:ILE:O	7:Es:781:GLN:HG3	2.01	0.61
8:Ex:253:LEU:HD13	8:Ex:370:SER:HB2	1.81	0.61
9:Fb:101:ILE:HA	7:Fh:16:ARG:HH22	1.65	0.61
9:Fe:455:GLU:O	9:Fe:459:LYS:HB2	1.99	0.61
6:Bd:126:GLN:HA	6:Bd:129:LYS:HE2	1.83	0.61
5:Cg:215:VAL:HG13	5:Cg:247:ILE:HD11	1.81	0.61
6:Ct:256:VAL:HG12	6:Ct:258:GLY:H	1.64	0.61
6:Dr:111:ALA:HB1	6:Dr:115:MET:HE3	1.82	0.61
7:Ei:794:ILE:H	7:Ei:794:ILE:HD12	1.64	0.61
7:En:797:ARG:HA	7:En:800:ASP:OD1	1.99	0.61
8:Ew:85:PHE:CD2	8:Ew:86:LEU:HD12	2.35	0.61
8:Ey:381:ARG:NH1	8:Ey:382:ARG:HH11	1.98	0.61
8:Fa:336:TYR:O	8:Fa:340:THR:HG23	2.00	0.61
9:Fb:605:MET:HB3	9:Fb:641:TYR:CE2	2.36	0.61

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
9:Fb:729:MET:HA	9:Fb:732:MET:HG3	1.82	0.61
9:Fc:756:LEU:O	9:Fc:760:ILE:HG12	1.99	0.61
5:Aq:245:LYS:HE2	5:Aq:257:SER:HA	1.83	0.61
2:Bl:2:VAL:HG23	3:Bm:1:ARG:HD2	1.82	0.61
6:Cn:117:ARG:HD2	6:Cn:117:ARG:C	2.26	0.61
5:Cy:222:LEU:HD11	5:Cy:262:ILE:HG21	1.82	0.61
6:Df:236:PRO:HG2	6:Dl:251:ARG:HH21	1.66	0.61
6:Dx:115:MET:HG2	7:Es:801:MET:SD	2.41	0.61
6:Ed:117:ARG:NH1	6:Ed:118:ASN:HA	2.16	0.61
7:Ei:802:LEU:O	7:Ei:806:THR:HG23	2.01	0.61
7:En:792:GLN:HA	7:En:795:GLN:NE2	2.16	0.61
8:Fa:444:LEU:HA	8:Fa:447:ASN:HD21	1.66	0.61
7:Fi:33:VAL:HA	7:Fi:36:PHE:CD2	2.35	0.61
5:Ak:254:ILE:HB	5:Ak:262:ILE:HB	1.81	0.61
6:Ax:115:MET:HA	6:Ax:118:ASN:OD1	2.00	0.61
6:Bp:134:TYR:O	6:Bp:138:GLU:HG3	2.01	0.61
3:Ce:2:VAL:HG21	6:Ct:129:LYS:HG2	1.83	0.61
6:Ed:106:VAL:HG13	6:Ed:110:LYS:NZ	2.16	0.61
8:Ew:430:ARG:HH21	8:Ex:428:LEU:HB2	1.64	0.61
9:Fb:726:ALA:HB2	9:Ff:725:PHE:CZ	2.36	0.61
9:Fc:504:PRO:O	9:Fc:509:CYS:HB2	2.01	0.61
9:Fc:924:ILE:HD12	9:Fc:925:ILE:HD13	1.82	0.61
9:Fe:666:GLY:O	9:Fe:670:ILE:HG12	2.01	0.61
5:Ae:219:ARG:H	6:Al:148:PRO:HD2	1.65	0.61
6:Bp:150:LYS:HG2	6:Bp:152:THR:HG23	1.83	0.61
5:Cg:219:ARG:HH21	5:Cg:231:THR:HG23	1.66	0.61
6:Cn:115:MET:HE1	7:Em:798:THR:O	2.01	0.61
2:Db:2:VAL:HA	3:Dc:1:ARG:HE	1.66	0.61
7:El:818:GLN:HG3	7:El:820:TYR:HE1	1.66	0.61
8:Ey:422:ILE:HD13	8:Ey:425:GLN:NE2	2.16	0.61
8:Ez:302:TRP:CZ3	8:Fa:218:LEU:HG	2.36	0.61
8:Fa:374:ASN:HA	8:Fa:377:ASN:OD1	2.00	0.61
9:Fb:741:THR:O	9:Fb:745:ILE:HG13	2.01	0.61
9:Fe:819:TYR:O	9:Fe:823:ILE:HD12	2.01	0.61
9:Ff:579:LEU:HD22	9:Ff:580:PRO:HD2	1.82	0.61
9:Ff:927:GLN:HG3	9:Ff:928:LYS:HD2	1.83	0.61
6:Ar:221:TYR:CZ	6:Ax:142:ALA:HB3	2.36	0.60
6:Dl:219:LYS:HE2	6:Dl:222:ASN:HD21	1.65	0.60
7:Ef:808:LEU:HB3	7:Ef:812:TRP:HE1	1.66	0.60
7:Ep:802:LEU:O	7:Ep:806:THR:HG23	2.01	0.60
8:Ew:425:GLN:HA	8:Fa:430:ARG:HH21	1.66	0.60

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
8:Fa:126:SER:HB2	8:Fa:171:ASN:HA	1.82	0.60
9:Fc:915:ILE:H	9:Fc:915:ILE:HD12	1.65	0.60
9:Fe:197:GLN:HG3	9:Fe:530:ILE:HD11	1.81	0.60
6:Ax:221:TYR:HD2	6:Ax:244:GLY:HA3	1.66	0.60
6:Ed:204:ILE:HD13	6:Ed:215:ILE:HG12	1.82	0.60
7:Eu:794:ILE:HD12	7:Eu:794:ILE:H	1.66	0.60
7:Ev:808:LEU:HA	7:Ev:811:ASP:OD2	2.00	0.60
8:Ew:346:ASN:HD22	8:Ew:435:ARG:HH21	1.48	0.60
9:Fc:811:ARG:O	9:Fc:815:MET:HG3	2.01	0.60
9:Ff:657:MET:O	9:Ff:661:MET:HB2	2.00	0.60
9:Ff:741:THR:O	9:Ff:745:ILE:HG12	2.01	0.60
6:Ar:121:PRO:C	7:Ef:801:MET:HE1	2.26	0.60
6:Dx:230:LEU:HB2	6:Dx:233:LEU:HD13	1.83	0.60
7:Ek:809:VAL:HG22	7:Ek:813:LYS:HE3	1.83	0.60
9:Fb:697:ILE:HD12	9:Fc:909:TYR:CD2	2.35	0.60
9:Fc:757:PRO:HA	9:Fc:760:ILE:HD11	1.82	0.60
6:Ax:170:ARG:NH1	6:Ax:247:ALA:HB1	2.16	0.60
6:Bj:170:ARG:HB2	6:Bj:249:ASP:N	2.16	0.60
6:Dl:113:LYS:HB3	6:Dl:117:ARG:HH22	1.66	0.60
7:Ep:804:ALA:O	7:Ep:808:LEU:HD22	2.01	0.60
7:Ev:805:ALA:O	7:Ev:809:VAL:HG12	2.01	0.60
8:Ey:360:SER:HB2	8:Ey:366:ALA:HA	1.82	0.60
8:Ey:399:TRP:HH2	8:Ey:414:GLU:HB3	1.65	0.60
8:Ey:453:PRO:HD2	8:Ez:290:PRO:HB3	1.83	0.60
9:Fd:739:ILE:O	9:Fd:743:VAL:HG12	2.01	0.60
9:Fd:834:ASN:HA	9:Fd:837:PHE:CE1	2.36	0.60
6:Af:139:TYR:HB3	2:Dz:5:ILE:HD12	1.82	0.60
1:Am:3:ASP:HB2	2:An:9:ARG:HH22	1.67	0.60
6:Ax:151:PRO:HA	6:Ax:248:VAL:HG13	1.82	0.60
5:De:268:ASP:HB2	6:Df:152:THR:HG21	1.83	0.60
5:Dk:210:TYR:HB3	5:Dk:222:LEU:HD11	1.83	0.60
8:Fa:304:GLN:OE1	8:Fa:319:GLN:HA	2.01	0.60
9:Fc:320:ILE:HD11	9:Fc:324:GLN:HB3	1.82	0.60
9:Fd:643:PHE:O	9:Fd:647:ILE:HG13	2.01	0.60
9:Fe:808:VAL:HA	9:Fe:811:ARG:HH12	1.67	0.60
6:Ar:168:VAL:HG13	6:Ar:170:ARG:HH12	1.66	0.60
2:Cv:2:VAL:HG22	3:Cw:1:ARG:HE	1.67	0.60
6:Dr:228:VAL:HG22	6:Dr:237:VAL:HB	1.84	0.60
5:Dw:219:ARG:NH1	5:Dw:233:ARG:HE	1.99	0.60
7:Er:806:THR:HA	7:Er:809:VAL:HG22	1.84	0.60
8:Ew:236:ASN:ND2	8:Fa:182:LEU:HD13	2.17	0.60

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
9:Fb:44:VAL:H	9:Fb:55:GLN:HE22	1.49	0.60
9:Fc:605:MET:HG2	9:Fc:641:TYR:CD2	2.37	0.60
9:Fc:651:MET:HA	9:Fc:651:MET:HE3	1.84	0.60
9:Fd:274:PHE:CZ	9:Fd:515:LEU:HB2	2.36	0.60
7:Fh:34:ILE:O	7:Fh:38:LYS:HD3	2.02	0.60
6:Ar:138:GLU:HA	6:Ar:141:LYS:HB3	1.84	0.60
6:Bd:150:LYS:N	6:Bd:247:ALA:HA	2.17	0.60
1:Be:4:ALA:HB3	2:Bf:9:ARG:HH12	1.67	0.60
5:Bi:210:TYR:HB3	5:Bi:222:LEU:HD12	1.84	0.60
6:Bj:173:GLN:HB2	6:Bj:220:LEU:HD23	1.84	0.60
6:Bj:206:TRP:HB2	6:Bj:213:LEU:HD23	1.84	0.60
6:Dl:238:MET:HE2	6:Dl:238:MET:HA	1.84	0.60
8:Ew:438:LEU:O	8:Ew:442:ILE:HG12	2.01	0.60
8:Ez:207:GLN:O	8:Ez:210:VAL:HG12	2.01	0.60
8:Ez:301:LEU:HD13	8:Ez:326:LEU:CD2	2.31	0.60
8:Fa:189:PHE:HE1	8:Fa:292:LEU:HD23	1.67	0.60
8:Fa:277:LEU:CD2	8:Fa:281:ARG:HH12	2.14	0.60
9:Fb:516:LEU:HA	9:Fb:519:TRP:CE3	2.36	0.60
9:Fe:729:MET:HA	9:Fe:732:MET:HG3	1.83	0.60
9:Fe:765:SER:O	9:Fe:768:TRP:HB3	2.02	0.60
9:Fe:827:TYR:HB3	9:Fe:831:TRP:CZ2	2.37	0.60
9:Ff:589:THR:HG22	9:Ff:665:GLN:HE22	1.67	0.60
1:Aa:16:ALA:HB3	2:Ah:9:ARG:HG3	1.83	0.60
6:Ar:107:ILE:HA	6:Ar:110:LYS:NZ	2.17	0.60
5:Bo:211:TYR:HB2	5:Bo:213:GLN:HE22	1.67	0.60
6:Cb:117:ARG:NH1	6:Cb:118:ASN:HA	2.16	0.60
6:Ct:112:PHE:N	7:En:797:ARG:HH22	2.00	0.60
5:Dk:263:LYS:HA	5:Dk:263:LYS:HE3	1.83	0.60
7:Et:799:SER:O	7:Et:803:THR:HG23	2.02	0.60
8:Ew:206:TYR:CZ	8:Ew:208:ASN:HB3	2.36	0.60
8:Ex:195:GLN:HE22	8:Ex:294:LYS:HD2	1.66	0.60
8:Ez:438:LEU:O	8:Ez:442:ILE:HG12	2.02	0.60
8:Fa:377:ASN:O	8:Fa:381:ARG:HD2	2.00	0.60
9:Fd:644:PHE:HA	9:Fd:647:ILE:HD12	1.84	0.60
9:Fd:709:ASN:ND2	9:Fe:597:LYS:H	1.98	0.60
9:Fe:140:ASP:O	9:Fe:144:GLU:HG2	2.02	0.60
9:Fe:164:THR:HA	9:Fe:579:LEU:HD13	1.83	0.60
9:Fe:461:TRP:CH2	9:Fe:812:PRO:HB3	2.37	0.60
9:Fe:926:VAL:HG13	9:Fe:930:PHE:HE2	1.65	0.60
7:Fk:32:VAL:O	7:Fk:36:PHE:HD1	1.84	0.60
6:Ar:128:VAL:HG23	6:Ar:132:GLN:HE21	1.67	0.60

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
6:Ar:181:PHE:HE2	6:Ar:191:ILE:HD11	1.66	0.60
6:Bj:125:GLU:O	6:Bj:129:LYS:HG3	2.01	0.60
6:Ch:131:LYS:O	6:Ch:135:GLU:HG3	2.02	0.60
5:Dq:232:VAL:HG23	5:Dq:236:SER:HB3	1.84	0.60
5:Dq:253:ARG:HB2	5:Dq:263:LYS:NZ	2.17	0.60
6:Dr:115:MET:HE2	7:Er:798:THR:HA	1.83	0.60
6:Dr:226:LEU:HB3	6:Dr:241:LEU:HD21	1.84	0.60
7:Ev:794:ILE:H	7:Ev:794:ILE:HD12	1.67	0.60
8:Ex:321:GLN:O	8:Ex:325:THR:HG23	2.02	0.60
8:Fa:429:ASP:O	8:Fa:433:GLN:HG2	2.01	0.60
6:Bd:117:ARG:HH21	7:Eh:797:ARG:HD2	1.66	0.60
5:Bu:264:PHE:HB3	5:Bu:269:SER:HB3	1.84	0.60
6:Dl:141:LYS:HE3	7:Es:812:TRP:CE3	2.37	0.60
8:Ew:84:THR:HB	8:Ew:371:GLN:HB2	1.83	0.60
8:Ey:238:PRO:O	8:Ey:242:GLN:HG2	2.01	0.60
8:Ez:272:GLN:HA	8:Ez:275:GLN:HG2	1.84	0.60
8:Fa:81:ALA:HB2	8:Fa:378:MET:CE	2.32	0.60
9:Fb:196:LYS:HE2	9:Fb:359:THR:HB	1.83	0.60
9:Fb:505:PHE:HA	9:Fb:509:CYS:HB3	1.83	0.60
9:Fd:342:THR:HG23	9:Fd:343:LEU:HD12	1.83	0.60
9:Fe:450:ARG:HD3	9:Fe:474:LEU:HD11	1.84	0.60
9:Fe:558:SER:HB3	9:Fe:576:MET:SD	2.42	0.60
9:Fe:741:THR:O	9:Fe:745:ILE:HG12	2.01	0.60
7:Fj:38:LYS:HZ1	7:Fj:42:ALA:HB2	1.66	0.60
6:Al:159:ASN:HD22	6:Al:257:GLN:NE2	2.00	0.59
6:Ar:235:THR:HG23	6:Ax:251:ARG:CZ	2.32	0.59
5:Bc:216:ILE:HD11	6:Bj:148:PRO:HG2	1.84	0.59
6:Cz:115:MET:HA	6:Cz:118:ASN:OD1	2.02	0.59
6:Cz:255:ARG:HH12	6:Cz:257:GLN:HA	1.65	0.59
7:Ef:804:ALA:O	7:Ef:808:LEU:HD22	2.01	0.59
7:Eo:798:THR:O	7:Eo:802:LEU:HD22	2.02	0.59
7:Et:779:GLN:HA	7:Et:782:ASN:HD21	1.67	0.59
8:Ew:170:LEU:HD12	8:Ew:171:ASN:H	1.67	0.59
8:Ew:437:LEU:HD21	8:Ex:432:ILE:HG22	1.83	0.59
8:Ey:419:LEU:HD12	8:Ey:420:ALA:N	2.17	0.59
8:Fa:144:GLN:HB2	8:Fa:179:PHE:CZ	2.36	0.59
9:Fc:103:ILE:HG13	9:Fc:104:PRO:HD3	1.84	0.59
9:Fc:670:ILE:HA	9:Fc:673:VAL:HG22	1.83	0.59
9:Fe:434:LEU:O	9:Fe:438:ARG:HG2	2.01	0.59
9:Ff:117:LYS:HG3	9:Ff:119:SER:H	1.65	0.59
9:Ff:616:LYS:HA	9:Ff:620:PHE:O	2.02	0.59

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
7:Fh:18:ARG:O	7:Fh:22:ILE:HG12	2.02	0.59
6:Af:123:ASN:ND2	6:Af:124:PRO:HD2	2.17	0.59
6:Af:223:TYR:HB2	6:Af:242:ILE:HA	1.84	0.59
5:Bc:207:ARG:HG2	5:Bc:208:ILE:H	1.66	0.59
5:De:213:GLN:HG2	5:De:221:TRP:O	2.02	0.59
8:Ey:82:TYR:HE1	8:Ez:383:LEU:HD23	1.66	0.59
8:Ey:93:ALA:HA	8:Ey:99:MET:SD	2.42	0.59
8:Ey:144:GLN:HB2	8:Ey:179:PHE:CZ	2.37	0.59
8:Ey:419:LEU:O	8:Ey:422:ILE:HB	2.02	0.59
9:Fb:197:GLN:HG3	9:Fb:530:ILE:HD11	1.84	0.59
9:Fb:419:PHE:O	9:Fb:423:ILE:HG12	2.02	0.59
9:Fd:716:LEU:HD21	9:Fe:723:PHE:CD1	2.37	0.59
9:Ff:67:ALA:O	9:Ff:71:LEU:HG	2.01	0.59
9:Ff:631:LEU:O	9:Ff:636:VAL:HG23	2.02	0.59
6:Al:115:MET:CE	7:Ev:801:MET:HB2	2.31	0.59
6:Ax:203:ASN:HD21	6:Ax:205:GLN:HE22	1.49	0.59
6:Bd:202:PHE:HA	6:Bd:218:THR:H	1.67	0.59
6:Bv:229:ARG:CZ	6:Bv:230:LEU:H	2.15	0.59
5:Ca:210:TYR:HB3	5:Ca:222:LEU:HD12	1.82	0.59
6:Cb:126:GLN:HA	6:Cb:129:LYS:HE2	1.84	0.59
5:Cs:219:ARG:CZ	5:Cs:233:ARG:HB3	2.32	0.59
7:Eg:794:ILE:O	7:Eg:798:THR:HG23	2.02	0.59
8:Ew:382:ARG:O	8:Ew:399:TRP:HB3	2.02	0.59
8:Fa:109:ASN:HA	8:Fa:112:ILE:HD12	1.83	0.59
9:Fb:128:PHE:O	9:Fb:132:ILE:HD12	2.02	0.59
9:Fd:811:ARG:HH21	9:Fd:815:MET:HE2	1.66	0.59
6:Bd:119:LEU:HD12	6:Bd:120:TYR:CD2	2.37	0.59
3:Bs:2:VAL:HG22	3:Bs:8:VAL:HG22	1.83	0.59
6:Bv:229:ARG:HH11	6:Bv:236:PRO:HA	1.67	0.59
5:Dk:269:SER:O	6:Dl:150:LYS:HE3	2.02	0.59
6:Dx:129:LYS:O	6:Dx:133:ILE:HG12	2.02	0.59
7:Eq:803:THR:HA	7:Eq:807:GLN:HE22	1.67	0.59
7:Eu:798:THR:O	7:Eu:802:LEU:HD22	2.02	0.59
8:Ew:340:THR:O	8:Ew:344:VAL:HG23	2.03	0.59
9:Fb:131:VAL:HG21	9:Fb:805:LEU:HD11	1.84	0.59
9:Fb:593:LEU:H	9:Fb:593:LEU:HD23	1.67	0.59
9:Fc:816:ILE:O	9:Fc:820:ILE:HG22	2.03	0.59
9:Fe:261:LEU:HB3	9:Fe:282:VAL:HG12	1.84	0.59
6:Ax:219:LYS:HD2	6:Ax:222:ASN:HD21	1.67	0.59
5:Bc:253:ARG:HG2	5:Bc:261:VAL:HG13	1.85	0.59
6:Bp:225:ASN:HD21	6:Bv:176:VAL:N	1.99	0.59

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
5:Ca:222:LEU:HB2	5:Ca:230:LEU:HG	1.85	0.59
6:Cz:193:ALA:HB1	7:Er:820:TYR:HE2	1.67	0.59
6:Dl:214:MET:HE1	7:Es:818:GLN:HE21	1.67	0.59
7:Eo:799:SER:O	7:Eo:803:THR:HG23	2.02	0.59
7:Eq:771:ALA:O	7:Eq:775:GLN:HG3	2.01	0.59
8:Ew:208:ASN:HA	8:Ew:211:MET:SD	2.42	0.59
8:Ex:123:GLN:HE22	8:Ex:256:THR:HG21	1.68	0.59
8:Ex:301:LEU:HD12	8:Ex:322:ALA:HB1	1.83	0.59
8:Ex:382:ARG:HG2	8:Ex:399:TRP:CD1	2.37	0.59
8:Ex:453:PRO:HD3	8:Ey:336:TYR:CD1	2.36	0.59
8:Ey:217:THR:HG22	8:Ey:265:SER:HB3	1.85	0.59
8:Ey:375:GLU:HA	8:Ey:378:MET:HG2	1.83	0.59
9:Fd:339:MET:HE2	9:Fd:339:MET:N	2.17	0.59
9:Fd:735:LEU:HA	9:Fd:738:TRP:CE3	2.38	0.59
9:Fe:334:ILE:HA	9:Fe:337:GLN:HB3	1.83	0.59
9:Fe:760:ILE:HA	9:Fe:763:PHE:CD1	2.36	0.59
9:Ff:812:PRO:HA	9:Ff:815:MET:HE1	1.85	0.59
9:Ff:819:TYR:CE1	9:Ff:823:ILE:HD11	2.36	0.59
1:Aa:15:ASN:HB3	2:Ah:9:ARG:HH22	1.68	0.59
5:Ae:264:PHE:HE2	6:Af:170:ARG:HH12	1.50	0.59
6:Af:129:LYS:HE3	6:Al:112:PHE:CE1	2.37	0.59
5:Ak:219:ARG:H	6:Ar:148:PRO:HD2	1.66	0.59
6:Cb:238:MET:HG3	6:Ch:250:TYR:HB2	1.85	0.59
6:Dl:115:MET:HG3	7:Eq:801:MET:SD	2.42	0.59
8:Ex:252:LEU:HD11	8:Ex:373:LEU:HA	1.84	0.59
9:Fb:696:TYR:HA	9:Fb:742:MET:HE3	1.84	0.59
7:Fg:23:PHE:O	7:Fg:27:LEU:HD12	2.03	0.59
6:Ax:238:MET:HE2	6:Ax:238:MET:HA	1.85	0.59
6:Bp:194:TYR:HB2	6:Bp:228:VAL:HG12	1.85	0.59
6:Cb:141:LYS:HZ3	7:Em:808:LEU:HD13	1.65	0.59
6:Ch:219:LYS:HE2	6:Ch:222:ASN:HD21	1.68	0.59
6:Cn:115:MET:O	6:Cn:119:LEU:HD23	2.02	0.59
6:Dl:220:LEU:HD23	6:Dl:221:TYR:HB3	1.84	0.59
8:Ew:136:ILE:HD11	8:Ew:273:ALA:HB1	1.85	0.59
8:Ew:342:VAL:HG21	8:Ew:442:ILE:HG21	1.83	0.59
8:Ex:214:VAL:HA	8:Ex:267:LEU:HD23	1.85	0.59
8:Ey:430:ARG:HD3	8:Ez:429:ASP:HB3	1.85	0.59
9:Fd:844:ILE:HG22	9:Fd:910:ALA:HB1	1.83	0.59
6:Ar:145:PRO:HD3	6:Ax:131:LYS:NZ	2.17	0.59
5:Aw:212:ILE:HG23	5:Aw:222:LEU:HD22	1.85	0.59
5:Ca:264:PHE:HB3	5:Ca:269:SER:HB3	1.85	0.59

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
6:Ch:141:LYS:HZ3	7:En:808:LEU:HA	1.67	0.59
6:Df:151:PRO:HA	6:Df:248:VAL:HG22	1.83	0.59
6:Df:190:PRO:O	6:Df:231:ARG:HG2	2.03	0.59
7:Ee:794:ILE:H	7:Ee:794:ILE:HD12	1.66	0.59
7:Ei:791:GLN:HG3	7:Ei:794:ILE:HD13	1.85	0.59
8:Ew:431:GLN:O	8:Ew:435:ARG:HG2	2.02	0.59
8:Ey:444:LEU:HD22	8:Ez:443:MET:HB3	1.84	0.59
9:Fd:744:SER:O	9:Fd:748:VAL:HG23	2.02	0.59
9:Fe:640:VAL:HG12	9:Fe:644:PHE:CE2	2.38	0.59
9:Ff:940:VAL:HG12	9:Ff:943:TRP:CZ3	2.38	0.59
6:Ax:176:VAL:HG13	6:Ax:214:MET:HG3	1.84	0.59
6:Df:185:THR:HG21	6:Df:263:ALA:HA	1.85	0.59
5:Dk:214:ALA:HB3	5:Dk:221:TRP:HD1	1.67	0.59
6:Ed:107:ILE:HD11	7:Et:794:ILE:HA	1.84	0.59
8:Ew:334:ARG:HB2	8:Ex:243:LEU:HD11	1.85	0.59
8:Ex:298:TYR:HE2	8:Ey:221:ALA:HB2	1.68	0.59
9:Fc:121:TYR:HB3	9:Fc:125:GLN:NE2	2.18	0.59
9:Fc:124:MET:HA	9:Fc:127:PHE:CD1	2.38	0.59
9:Fd:469:PHE:HD2	9:Fe:843:TYR:CD2	2.21	0.59
9:Fe:129:MET:O	9:Fe:133:VAL:HG12	2.02	0.59
6:Dl:130:LEU:HD23	7:Er:808:LEU:HD12	1.85	0.59
7:Es:784:GLN:O	7:Es:788:GLN:HG2	2.02	0.59
8:Ey:123:GLN:HG2	8:Ey:136:ILE:HB	1.85	0.59
8:Fa:245:SER:HA	8:Fa:279:PHE:CE2	2.38	0.59
9:Fb:391:GLU:HG3	9:Fb:393:GLN:H	1.68	0.59
9:Fb:664:LEU:HD21	9:Fb:738:TRP:CZ2	2.38	0.59
9:Fb:927:GLN:O	9:Fb:931:THR:HG23	2.03	0.59
9:Fc:288:SER:HA	9:Fc:291:ASN:HB3	1.83	0.59
9:Fc:939:LYS:HA	9:Fc:942:ARG:HG2	1.84	0.59
9:Fd:716:LEU:HD21	9:Fe:723:PHE:HD1	1.66	0.59
9:Ff:663:PRO:HG2	9:Ff:738:TRP:CE3	2.37	0.59
6:Ar:204:ILE:HG12	6:Ar:215:ILE:HG23	1.85	0.58
6:Dr:219:LYS:HG2	6:Dr:222:ASN:ND2	2.18	0.58
6:Ed:202:PHE:CE1	6:Ed:241:LEU:HD13	2.37	0.58
7:En:799:SER:HA	7:En:802:LEU:HG	1.84	0.58
8:Ex:173:PRO:HB2	8:Ex:348:TYR:CZ	2.38	0.58
9:Fc:628:MET:HE3	9:Fc:629:GLY:HA2	1.85	0.58
9:Fd:335:ALA:HA	9:Fd:429:ILE:HD13	1.85	0.58
9:Fe:193:GLY:O	9:Fe:197:GLN:HG2	2.02	0.58
9:Fe:808:VAL:HA	9:Fe:811:ARG:NH1	2.18	0.58
7:Fh:35:GLY:O	7:Fh:39:ILE:HG12	2.03	0.58

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
5:Ae:245:LYS:HD3	5:Ae:257:SER:HA	1.85	0.58
1:As:16:ALA:HB3	2:Az:9:ARG:HG3	1.85	0.58
5:Bc:207:ARG:HD2	5:Bc:240:GLY:HA3	1.85	0.58
6:Bd:205:GLN:NE2	7:El:820:TYR:HB2	2.17	0.58
5:Bu:219:ARG:HA	5:Bu:232:VAL:O	2.03	0.58
5:Bu:243:MET:HG2	5:Bu:257:SER:HB3	1.85	0.58
6:Cz:115:MET:CE	7:Eo:801:MET:HG3	2.33	0.58
2:Dn:2:VAL:HA	3:Do:1:ARG:HH21	1.66	0.58
6:Dr:108:ASP:O	6:Dr:111:ALA:HB3	2.03	0.58
6:Ed:204:ILE:HD11	6:Ed:213:LEU:HD22	1.84	0.58
8:Ey:346:ASN:HA	8:Ey:435:ARG:HH12	1.67	0.58
9:Fb:113:LEU:HD13	9:Fb:123:MET:HE3	1.85	0.58
9:Fb:531:GLN:HA	9:Fb:534:ILE:HD12	1.86	0.58
9:Fb:571:VAL:HG12	9:Fb:575:MET:HE1	1.83	0.58
9:Fb:917:ILE:O	9:Fb:921:MET:HG2	2.03	0.58
9:Fb:939:LYS:HZ1	9:Ff:774:GLU:HA	1.69	0.58
9:Fd:668:LYS:O	9:Fd:672:ILE:HG12	2.01	0.58
9:Fe:800:PHE:O	9:Fe:804:ILE:HD12	2.03	0.58
5:Ak:219:ARG:HG2	6:Ar:148:PRO:HD2	1.83	0.58
6:Ax:108:ASP:HA	7:Ef:797:ARG:NH1	2.18	0.58
5:Dk:210:TYR:HB3	5:Dk:222:LEU:HD21	1.84	0.58
6:Dx:196:LEU:HD21	6:Dx:202:PHE:HD1	1.68	0.58
6:Ed:223:TYR:HB2	6:Ed:242:ILE:HG13	1.85	0.58
7:Ef:804:ALA:O	7:Ef:807:GLN:HG3	2.03	0.58
7:Eg:802:LEU:O	7:Eg:806:THR:HG23	2.03	0.58
7:El:780:LYS:O	7:El:784:GLN:HG2	2.02	0.58
7:El:797:ARG:HB2	7:El:801:MET:HE3	1.85	0.58
8:Ew:95:SER:HB3	8:Ew:98:LEU:HD23	1.86	0.58
8:Ew:443:MET:SD	8:Ew:444:LEU:HD23	2.43	0.58
8:Ex:277:LEU:O	8:Ex:280:ILE:HG12	2.03	0.58
8:Ey:123:GLN:NE2	8:Ey:136:ILE:H	2.01	0.58
8:Ey:242:GLN:HE22	8:Ey:269:ALA:H	1.52	0.58
9:Fb:907:GLY:O	9:Fb:910:ALA:HB3	2.03	0.58
9:Fe:667:MET:HE2	9:Fe:912:PHE:HD1	1.69	0.58
6:Df:117:ARG:O	6:Df:121:PRO:HA	2.03	0.58
8:Ew:178:VAL:HA	8:Ew:181:ILE:HD12	1.84	0.58
8:Ez:381:ARG:NE	8:Ez:382:ARG:HG3	2.13	0.58
8:Fa:125:PHE:HA	8:Fa:137:THR:HA	1.85	0.58
9:Fb:434:LEU:HG	9:Fb:438:ARG:HH21	1.67	0.58
9:Fb:613:GLY:HA3	9:Fb:625:GLY:H	1.68	0.58
9:Fd:60:MET:HE1	9:Fe:817:ILE:HD12	1.86	0.58

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
9:Fd:647:ILE:O	9:Fd:651:MET:HG2	2.02	0.58
9:Fe:668:LYS:O	9:Fe:672:ILE:HG12	2.03	0.58
6:Bp:131:LYS:HD3	6:Bp:132:GLN:HE22	1.68	0.58
6:Bp:238:MET:HG3	6:Bv:250:TYR:CE1	2.38	0.58
6:Bv:113:LYS:HA	6:Bv:113:LYS:HE3	1.84	0.58
5:Dk:212:ILE:HG13	5:Dk:262:ILE:HG22	1.85	0.58
3:Do:2:VAL:HG22	3:Do:8:VAL:HB	1.85	0.58
6:Dx:149:PRO:HA	6:Dx:246:LYS:O	2.03	0.58
2:Dz:1:CYS:HA	3:Ea:4:ILE:HD13	1.86	0.58
7:Eq:779:GLN:HA	7:Eq:782:ASN:ND2	2.18	0.58
8:Ew:382:ARG:HB3	8:Ew:399:TRP:CD2	2.38	0.58
8:Ex:412:GLN:HG3	8:Ey:411:VAL:HG21	1.85	0.58
9:Fb:335:ALA:HA	9:Fb:429:ILE:HD11	1.86	0.58
9:Fc:78:TYR:O	9:Fc:81:MET:HG2	2.03	0.58
9:Fe:187:GLY:C	9:Fe:191:MET:HE2	2.29	0.58
9:Fe:769:LEU:HA	9:Fe:772:VAL:HG12	1.85	0.58
9:Ff:283:LYS:HE2	9:Ff:494:SER:HA	1.86	0.58
6:Bd:237:VAL:C	6:Bj:251:ARG:HH22	2.12	0.58
2:Bf:2:VAL:HG22	3:Bg:1:ARG:HE	1.68	0.58
6:Bv:187:ALA:HB1	6:Bv:262:ASN:HB2	1.86	0.58
5:Cm:219:ARG:NH1	5:Cm:233:ARG:HD3	2.19	0.58
6:Cn:144:THR:OG1	6:Cn:148:PRO:HG3	2.03	0.58
6:Cn:203:ASN:HB3	6:Cn:216:GLN:HE21	1.68	0.58
6:Cz:189:TRP:CD1	6:Cz:230:LEU:HD12	2.39	0.58
6:Cz:219:LYS:HD3	6:Cz:220:LEU:N	2.19	0.58
5:De:243:MET:HG3	5:De:245:LYS:HZ2	1.69	0.58
6:Dx:121:PRO:HB2	7:Et:801:MET:HB2	1.85	0.58
7:Eq:809:VAL:HG22	7:Eq:813:LYS:HE3	1.86	0.58
8:Ew:383:LEU:CD1	8:Fa:78:GLN:HG2	2.32	0.58
8:Ey:141:LEU:HG	8:Ey:213:ASN:HB3	1.85	0.58
8:Fa:294:LYS:HA	8:Fa:294:LYS:HE3	1.84	0.58
8:Fa:346:ASN:O	8:Fa:350:ILE:HG13	2.03	0.58
9:Fb:619:PHE:H	9:Fc:618:LEU:HD21	1.67	0.58
9:Fb:657:MET:HB2	9:Fb:661:MET:HE1	1.84	0.58
9:Fb:675:VAL:O	9:Fb:679:THR:HG23	2.04	0.58
9:Fb:781:ILE:H	9:Fb:781:ILE:HD12	1.68	0.58
9:Fb:939:LYS:NZ	9:Ff:774:GLU:HA	2.18	0.58
9:Fc:532:SER:HA	9:Fc:537:VAL:H	1.68	0.58
9:Fc:940:VAL:HG22	9:Fc:943:TRP:CZ3	2.39	0.58
9:Fd:596:PHE:C	9:Fd:597:LYS:HE2	2.28	0.58
9:Ff:304:VAL:HB	9:Ff:318:LEU:H	1.66	0.58

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
7:Fh:25:ALA:O	7:Fh:29:ILE:HG22	2.04	0.58
6:Bd:208:LYS:HZ1	7:Ej:821:THR:HG22	1.68	0.58
6:Bj:208:LYS:HB2	7:Ek:824:THR:HG23	1.85	0.58
6:Ch:149:PRO:HB2	6:Ch:248:VAL:HG13	1.85	0.58
6:Ct:204:ILE:HA	6:Ct:214:MET:O	2.04	0.58
6:Dl:141:LYS:HE3	7:Es:812:TRP:CZ3	2.38	0.58
5:Dq:208:ILE:HG23	5:Dq:225:SER:HB2	1.84	0.58
5:Dq:210:TYR:CE1	5:Dq:238:ILE:HD11	2.39	0.58
6:Dr:125:GLU:C	6:Dr:129:LYS:HZ2	2.11	0.58
8:Ew:384:PHE:HB2	8:Fa:78:GLN:HE22	1.69	0.58
8:Fa:36:THR:HA	8:Fa:39:LEU:HD23	1.85	0.58
8:Fa:295:LEU:O	8:Fa:299:SER:HB2	2.04	0.58
8:Fa:420:ALA:O	8:Fa:424:TYR:HD1	1.87	0.58
9:Fb:699:PHE:HD2	9:Fb:742:MET:HE2	1.66	0.58
9:Fc:38:LEU:HD22	9:Fc:49:LEU:HD13	1.86	0.58
9:Fc:644:PHE:CE1	9:Fc:718:PRO:HD2	2.39	0.58
9:Fc:668:LYS:O	9:Fc:672:ILE:HG12	2.04	0.58
9:Fe:147:LEU:HB3	9:Fe:453:ILE:HG23	1.84	0.58
9:Fe:338:GLN:HA	9:Fe:341:VAL:HG22	1.86	0.58
3:Ao:2:VAL:HG22	3:Ao:8:VAL:HG12	1.85	0.58
6:Ax:144:THR:OG1	6:Ax:148:PRO:HG3	2.03	0.58
6:Dl:214:MET:HE1	7:Es:818:GLN:HG3	1.85	0.58
5:Ec:219:ARG:HH11	5:Ec:233:ARG:NH2	2.02	0.58
6:Ed:156:GLN:HE21	6:Ed:254:LEU:HA	1.68	0.58
6:Ed:199:PRO:HD3	7:Ee:818:GLN:OE1	2.03	0.58
8:Ex:295:LEU:O	8:Ex:299:SER:HB3	2.03	0.58
8:Ey:94:MET:HE1	8:Ez:237:GLN:O	2.04	0.58
8:Ey:346:ASN:O	8:Ey:350:ILE:HG13	2.03	0.58
8:Ey:363:GLN:HA	8:Ey:366:ALA:HB2	1.86	0.58
9:Fb:56:ILE:HG13	9:Fc:149:TYR:HE2	1.68	0.58
9:Fd:287:ILE:HG22	9:Fd:289:ALA:H	1.68	0.58
9:Fe:909:TYR:HA	9:Fe:912:PHE:CD2	2.38	0.58
9:Ff:512:PRO:HG2	9:Ff:513:TYR:CE1	2.38	0.58
5:Ae:245:LYS:HZ3	5:Ae:255:LEU:HB3	1.68	0.58
6:Ar:205:GLN:HG3	7:Eg:818:GLN:NE2	2.18	0.58
6:Bp:125:GLU:O	6:Bp:128:VAL:HG22	2.04	0.58
3:Bs:4:ILE:HG13	6:Ch:132:GLN:NE2	2.19	0.58
6:Cb:114:ASP:O	6:Cb:117:ARG:HB3	2.03	0.58
5:Cg:219:ARG:HG2	6:Cn:148:PRO:HD2	1.85	0.58
6:Ch:115:MET:HE2	7:El:802:LEU:HD22	1.86	0.58
6:Cz:111:ALA:O	6:Cz:115:MET:HG3	2.04	0.58

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:Db:7:GLY:H	4:Dd:1:PRO:HD2	1.69	0.58
5:Dk:213:GLN:HB3	5:Dk:223:ILE:HG22	1.85	0.58
1:Dm:3:ASP:OD1	3:Do:14:TYR:HB3	2.04	0.58
7:Eh:789:LYS:O	7:Eh:789:LYS:HD2	2.04	0.58
7:Ev:809:VAL:HG13	7:Ev:810:GLN:NE2	2.19	0.58
9:Fb:657:MET:HA	9:Fb:660:LEU:CD2	2.33	0.58
9:Fc:703:LEU:O	9:Fc:707:LEU:HG	2.04	0.58
9:Fc:819:TYR:O	9:Fc:823:ILE:HD12	2.04	0.58
9:Fe:287:ILE:H	9:Fe:287:ILE:HD12	1.69	0.58
5:Ae:210:TYR:CE2	5:Ae:238:ILE:HD11	2.38	0.58
5:Ak:216:ILE:HD11	6:Ar:148:PRO:HG2	1.86	0.58
6:Al:129:LYS:HB2	3:Ea:2:VAL:HG11	1.85	0.58
6:Al:199:PRO:HG3	7:Eg:819:VAL:HG22	1.85	0.58
6:Bj:205:GLN:HE22	7:Ej:820:TYR:HB2	1.68	0.58
6:Bv:128:VAL:O	6:Bv:132:GLN:HG2	2.04	0.58
5:Cy:211:TYR:CG	5:Cy:265:SER:HB2	2.38	0.58
6:Dr:145:PRO:HD3	6:Dx:131:LYS:HE2	1.86	0.58
6:Dx:132:GLN:O	6:Dx:136:THR:HG23	2.03	0.58
6:Ed:112:PHE:O	6:Ed:116:THR:HG23	2.04	0.58
6:Ed:189:TRP:NE1	6:Ed:230:LEU:HB3	2.19	0.58
7:Eh:802:LEU:O	7:Eh:806:THR:HG23	2.03	0.58
7:Eo:801:MET:HA	7:Eo:801:MET:HE3	1.85	0.58
7:Eq:774:LEU:HD21	9:Fb:432:PRO:HB2	1.85	0.58
7:Et:805:ALA:HA	7:Et:808:LEU:HG	1.85	0.58
8:Ex:420:ALA:HA	8:Ex:423:ASN:OD1	2.03	0.58
8:Ey:411:VAL:O	8:Ey:415:ILE:HG13	2.03	0.58
9:Fd:377:GLN:HE22	9:Fd:402:VAL:H	1.49	0.58
9:Fd:815:MET:HE1	9:Fd:937:PRO:HG3	1.85	0.58
9:Fd:912:PHE:O	9:Fd:916:LEU:HD22	2.04	0.58
9:Fe:426:TYR:O	9:Fe:430:MET:HG2	2.04	0.58
9:Ff:73:GLY:O	9:Ff:77:MET:HE2	2.04	0.58
3:Bs:3:SER:O	6:Ch:132:GLN:HG3	2.03	0.57
6:Bv:115:MET:CE	7:Ej:802:LEU:HD13	2.33	0.57
6:Ch:133:ILE:HD11	6:Cn:120:TYR:HB2	1.86	0.57
6:Ch:208:LYS:HE2	7:Eo:823:GLY:HA2	1.86	0.57
6:Cz:219:LYS:HD2	6:Cz:222:ASN:CG	2.29	0.57
5:Dw:247:ILE:HG12	5:Dw:254:ILE:HG23	1.85	0.57
6:Ed:141:LYS:HD2	7:Ev:812:TRP:CH2	2.39	0.57
7:Eu:778:LEU:HA	7:Eu:781:GLN:HG3	1.85	0.57
8:Ex:77:VAL:HG13	8:Ex:374:ASN:HD21	1.67	0.57
8:Ex:326:LEU:HD12	8:Ey:218:LEU:HD12	1.86	0.57

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
8:Ey:414:GLU:O	8:Ey:418:LEU:HG	2.04	0.57
8:Ez:342:VAL:HG21	8:Ez:442:ILE:HG13	1.86	0.57
9:Fc:801:ALA:O	9:Fc:805:LEU:HB3	2.04	0.57
9:Fc:814:LEU:CD2	9:Fc:940:VAL:HG21	2.34	0.57
9:Fd:732:MET:SD	9:Fd:733:PRO:HD3	2.43	0.57
9:Fe:326:GLN:O	9:Fe:330:LEU:HD22	2.02	0.57
9:Ff:240:VAL:HA	9:Ff:333:ALA:HB1	1.86	0.57
5:Cs:245:LYS:HD3	5:Cs:245:LYS:N	2.18	0.57
8:Ex:75:GLN:HB2	8:Ey:400:ILE:HD11	1.86	0.57
8:Fa:35:ASN:O	8:Fa:39:LEU:HD22	2.04	0.57
8:Fa:170:LEU:HD12	8:Fa:170:LEU:H	1.69	0.57
9:Fd:185:LEU:HB2	9:Fd:343:LEU:HD23	1.85	0.57
9:Fd:381:PRO:HD3	9:Fd:573:ASN:HB3	1.87	0.57
9:Fe:579:LEU:H	9:Fe:579:LEU:HD12	1.68	0.57
9:Ff:175:SER:HB2	9:Ff:488:GLY:H	1.68	0.57
9:Ff:698:ASN:O	9:Ff:702:THR:HG23	2.04	0.57
9:Ff:829:GLY:HA2	9:Ff:832:ILE:HG12	1.86	0.57
6:Af:201:SER:HB2	6:Af:219:LYS:HD2	1.86	0.57
6:Af:206:TRP:HB2	6:Af:213:LEU:HD23	1.85	0.57
6:Bd:220:LEU:HG	6:Bd:221:TYR:CE2	2.39	0.57
6:Cb:121:PRO:HB2	7:El:801:MET:SD	2.44	0.57
1:Ci:2:THR:O	1:Ci:6:LEU:HD22	2.04	0.57
1:Co:11:TYR:CE1	3:Cq:9:TYR:HB3	2.40	0.57
6:Df:112:PHE:HA	6:Df:115:MET:HG3	1.86	0.57
6:Ed:189:TRP:CZ3	6:Ed:260:GLY:HA2	2.39	0.57
7:Eg:799:SER:O	7:Eg:802:LEU:HD12	2.05	0.57
8:Ew:424:TYR:CE2	8:Ex:380:THR:HA	2.38	0.57
9:Fb:757:PRO:HA	9:Fb:760:ILE:HD12	1.85	0.57
9:Fd:141:LYS:HA	9:Fd:144:GLU:OE1	2.04	0.57
9:Fd:238:ASN:HD22	9:Fd:337:GLN:NE2	2.01	0.57
9:Fd:641:TYR:HA	9:Fd:644:PHE:CD2	2.39	0.57
9:Fe:516:LEU:HA	9:Fe:519:TRP:CE3	2.39	0.57
6:Af:151:PRO:HA	6:Af:248:VAL:HG22	1.87	0.57
6:Af:191:ILE:HG13	6:Af:211:ASN:HA	1.85	0.57
6:Al:115:MET:HE2	7:Ev:798:THR:HA	1.85	0.57
6:Bp:204:ILE:HA	6:Bp:214:MET:O	2.04	0.57
6:Ch:121:PRO:HB2	7:Em:801:MET:CE	2.34	0.57
6:Cz:158:VAL:HB	6:Cz:256:VAL:HA	1.85	0.57
6:Df:122:LEU:HD22	7:Eq:808:LEU:HD11	1.86	0.57
6:Dl:223:TYR:HB2	6:Dl:242:ILE:HA	1.85	0.57
6:Dl:228:VAL:HG13	6:Dl:237:VAL:HB	1.86	0.57

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
6:Ed:189:TRP:HE1	6:Ed:231:ARG:N	2.03	0.57
7:Ej:774:LEU:O	7:Ej:778:LEU:HG	2.04	0.57
8:Ew:377:ASN:O	8:Ew:381:ARG:HG2	2.04	0.57
8:Ex:105:LYS:HZ3	8:Ey:368:ILE:N	2.02	0.57
8:Ey:411:VAL:HA	8:Ey:414:GLU:OE2	2.04	0.57
9:Fb:905:TRP:HD1	9:Ff:698:ASN:HA	1.70	0.57
9:Fd:180:GLY:HA3	9:Fd:491:LEU:HB3	1.86	0.57
9:Fd:511:ASP:HB3	9:Fd:512:PRO:HA	1.85	0.57
9:Fe:781:ILE:H	9:Fe:781:ILE:HD12	1.69	0.57
1:Ag:12:HIS:HD2	3:Ai:9:TYR:HB2	1.69	0.57
6:Ar:221:TYR:HA	6:Ar:243:PRO:HG2	1.84	0.57
5:Bo:246:LEU:HG	5:Bo:253:ARG:NH2	2.19	0.57
6:Ch:113:LYS:HZ3	6:Ch:117:ARG:HB2	1.70	0.57
6:Ct:196:LEU:HD11	6:Ct:202:PHE:CD2	2.40	0.57
7:Ei:809:VAL:HA	7:Ei:812:TRP:CE3	2.39	0.57
7:Ej:774:LEU:HA	7:Ej:777:ILE:HG12	1.86	0.57
8:Ey:238:PRO:HG2	8:Ey:239:LEU:HD12	1.87	0.57
9:Fb:555:GLN:HB2	9:Fb:657:MET:HE1	1.87	0.57
9:Fb:648:PHE:HB2	9:Fb:717:ILE:HD13	1.86	0.57
9:Fe:219:THR:HG23	9:Fe:222:MET:H	1.69	0.57
6:Af:148:PRO:HB2	5:Ec:219:ARG:HG2	1.87	0.57
6:Af:218:THR:HA	7:Ee:815:VAL:HG11	1.87	0.57
6:Bv:128:VAL:HG23	6:Bv:132:GLN:HE21	1.69	0.57
6:Cn:191:ILE:HG12	6:Cn:211:ASN:HA	1.87	0.57
7:Ei:780:LYS:N	7:Ei:780:LYS:HD3	2.20	0.57
7:Ek:797:ARG:O	7:Ek:801:MET:HG3	2.04	0.57
7:Ev:794:ILE:O	7:Ev:798:THR:HG23	2.04	0.57
9:Fc:594:ILE:HG13	9:Fc:905:TRP:HH2	1.69	0.57
9:Fc:658:ALA:HA	9:Fc:662:ILE:HG12	1.87	0.57
9:Fe:73:GLY:O	9:Fe:77:MET:HE2	2.05	0.57
9:Fe:813:SER:O	9:Fe:817:ILE:HG12	2.05	0.57
9:Ff:391:GLU:HG3	9:Ff:393:GLN:H	1.69	0.57
6:Al:238:MET:HG2	6:Ar:251:ARG:CZ	2.33	0.57
6:Ar:199:PRO:HG3	7:Eh:819:VAL:HG22	1.86	0.57
6:Cn:170:ARG:HE	6:Cn:249:ASP:CG	2.13	0.57
6:Cn:219:LYS:HB2	6:Cn:222:ASN:OD1	2.05	0.57
6:Cz:221:TYR:CE1	6:Df:138:GLU:HG3	2.39	0.57
6:Df:180:VAL:HG23	6:Df:211:ASN:HD21	1.69	0.57
6:Dx:134:TYR:O	6:Dx:138:GLU:HG3	2.04	0.57
6:Dx:135:GLU:HA	6:Dx:138:GLU:OE2	2.03	0.57
6:Dx:199:PRO:HG3	7:Ev:817:THR:O	2.05	0.57

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
7:Eo:805:ALA:O	7:Eo:809:VAL:HG12	2.05	0.57
8:Ex:399:TRP:O	8:Ex:403:ILE:HG12	2.05	0.57
9:Fb:45:VAL:HG12	9:Fb:126:VAL:HG12	1.86	0.57
9:Fb:122:CYS:HB2	9:Fb:124:MET:HG2	1.86	0.57
9:Fb:556:ARG:HB3	9:Fb:576:MET:HE1	1.87	0.57
9:Fb:609:ASP:HA	9:Fb:626:ARG:HH12	1.68	0.57
9:Fb:726:ALA:HA	9:Fb:729:MET:SD	2.44	0.57
9:Fc:103:ILE:HG22	9:Fc:106:ARG:NH2	2.20	0.57
9:Fd:769:LEU:O	9:Fd:773:ILE:HG12	2.05	0.57
9:Fe:339:MET:O	9:Fe:343:LEU:HG	2.04	0.57
7:Fk:35:GLY:HA2	7:Fk:38:LYS:HE2	1.85	0.57
6:Af:180:VAL:HG12	6:Af:212:THR:HG22	1.85	0.57
5:Ak:210:TYR:HA	5:Ak:224:GLY:HA2	1.85	0.57
6:Bd:112:PHE:HA	6:Bd:115:MET:HB3	1.87	0.57
1:Bk:16:ALA:HA	4:Bt:3:GLY:HA3	1.86	0.57
6:Bp:170:ARG:HG3	6:Bp:245:GLN:HB2	1.86	0.57
6:Bp:225:ASN:HB2	6:Bv:250:TYR:OH	2.04	0.57
1:Cc:10:GLU:HA	1:Cc:13:LYS:HG2	1.87	0.57
6:Cz:203:ASN:HB3	6:Cz:216:GLN:HG2	1.87	0.57
6:Dx:108:ASP:CA	7:Es:797:ARG:HH22	2.15	0.57
7:Ef:808:LEU:O	7:Ef:812:TRP:HD1	1.88	0.57
8:Ew:109:ASN:HA	8:Ew:112:ILE:HD12	1.86	0.57
8:Ew:205:MET:SD	8:Ew:210:VAL:HB	2.45	0.57
8:Ey:330:PHE:O	8:Ey:334:ARG:HG2	2.05	0.57
8:Ey:350:ILE:HA	8:Ey:353:LYS:HD2	1.87	0.57
9:Fb:148:SER:O	9:Fb:152:ARG:HG2	2.05	0.57
9:Fc:921:MET:O	9:Fc:925:ILE:HG12	2.05	0.57
9:Fd:657:MET:O	9:Fd:657:MET:HE3	2.05	0.57
6:Af:229:ARG:HA	6:Af:236:PRO:HB3	1.87	0.57
2:Ah:4:MET:HE2	4:Aj:2:PHE:HE2	1.68	0.57
1:Am:16:ALA:HB3	2:At:9:ARG:HG3	1.87	0.57
6:Bd:230:LEU:H	6:Bd:233:LEU:HD11	1.69	0.57
6:Bv:200:SER:HB3	7:Em:816:GLU:HG2	1.86	0.57
6:Cz:125:GLU:O	6:Cz:129:LYS:HG3	2.05	0.57
6:Df:125:GLU:O	6:Df:128:VAL:HG22	2.05	0.57
8:Fa:276:ALA:O	8:Fa:280:ILE:HG12	2.05	0.57
9:Fb:45:VAL:HG23	9:Fb:49:LEU:HD23	1.86	0.57
9:Fb:105:LEU:O	9:Fb:108:THR:HG22	2.04	0.57
9:Fd:500:GLN:HA	9:Fd:503:LYS:HG2	1.87	0.57
6:Ch:107:ILE:HG22	7:El:797:ARG:HD3	1.86	0.57
5:Dw:250:LEU:HD12	5:Dw:251:GLN:HG3	1.87	0.57

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
9:Fb:695:MET:O	9:Fb:695:MET:HE3	2.05	0.57
9:Fb:929:ALA:O	9:Fb:932:LEU:CB	2.53	0.57
9:Fb:932:LEU:HD12	9:Ff:770:ILE:HD12	1.87	0.57
9:Fc:370:PHE:HB2	9:Fc:410:THR:HG21	1.86	0.57
9:Fc:493:LYS:HA	9:Fc:493:LYS:HE3	1.87	0.57
9:Fc:684:ASN:HB3	9:Fc:687:VAL:HG22	1.86	0.57
9:Fd:186:GLY:HA2	9:Fd:189:ILE:HD11	1.87	0.57
9:Fd:590:PHE:HB3	9:Fd:593:LEU:HB2	1.87	0.57
9:Fd:631:LEU:O	9:Fd:636:VAL:HG23	2.05	0.57
9:Fd:745:ILE:HD11	9:Fd:916:LEU:HD13	1.87	0.57
9:Fe:124:MET:HG3	9:Fe:779:ALA:HB1	1.87	0.57
9:Fe:735:LEU:O	9:Fe:739:ILE:HG12	2.05	0.57
5:Ae:248:ASP:HB3	5:Ae:253:ARG:HB3	1.87	0.56
6:Af:109:LYS:O	6:Af:113:LYS:HG3	2.05	0.56
2:Bf:4:MET:HE3	5:Bi:249:SER:H	1.70	0.56
2:Cp:9:ARG:H	2:Cp:9:ARG:HD3	1.70	0.56
6:Cz:112:PHE:CA	6:Cz:115:MET:HE3	2.35	0.56
5:Dq:219:ARG:HG2	6:Dx:148:PRO:HD2	1.87	0.56
5:Dq:253:ARG:HG3	5:Dq:261:VAL:HG13	1.86	0.56
6:Dx:221:TYR:HD2	6:Dx:244:GLY:HA3	1.69	0.56
8:Ex:443:MET:HA	8:Ex:446:GLN:HG3	1.87	0.56
8:Ey:142:ILE:HD11	8:Ey:281:ARG:HH11	1.70	0.56
8:Ey:242:GLN:HB3	8:Ey:278:ASN:ND2	2.20	0.56
8:Ey:297:ALA:O	8:Ey:301:LEU:HD22	2.04	0.56
8:Ey:430:ARG:NH1	8:Ez:428:LEU:HB2	2.20	0.56
8:Ez:126:SER:HB2	8:Ez:171:ASN:HA	1.87	0.56
9:Fc:667:MET:O	9:Fc:670:ILE:HG13	2.05	0.56
9:Fe:180:GLY:O	9:Fe:184:ILE:HG22	2.05	0.56
9:Fe:586:LYS:HD2	9:Fe:587:PRO:HD2	1.86	0.56
9:Fe:769:LEU:O	9:Fe:773:ILE:HG13	2.04	0.56
9:Fe:822:ALA:HB1	9:Fe:930:PHE:CD1	2.40	0.56
9:Ff:71:LEU:O	9:Ff:75:ILE:HG12	2.05	0.56
9:Ff:124:MET:SD	9:Ff:779:ALA:HB1	2.45	0.56
9:Ff:765:SER:O	9:Ff:769:LEU:HG	2.05	0.56
6:Af:131:LYS:HD2	7:Ev:812:TRP:CH2	2.40	0.56
6:Af:250:TYR:CE2	6:Ed:238:MET:HB3	2.40	0.56
5:Bo:253:ARG:CZ	5:Bo:255:LEU:HD22	2.35	0.56
6:Ch:181:PHE:CZ	6:Ch:256:VAL:HG11	2.40	0.56
6:Ct:189:TRP:CD1	6:Ct:230:LEU:HD12	2.39	0.56
6:Df:111:ALA:O	6:Df:115:MET:HG2	2.05	0.56
6:Df:149:PRO:HB2	6:Df:248:VAL:HG13	1.86	0.56

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
6:Ed:110:LYS:HE3	7:Et:790:TYR:OH	2.06	0.56
7:Et:782:ASN:HA	7:Et:785:LEU:HG	1.88	0.56
7:Eu:793:GLU:HA	7:Eu:796:GLN:HE21	1.70	0.56
8:Ew:340:THR:HA	8:Fa:448:LEU:HD22	1.86	0.56
9:Fb:258:SER:HB2	9:Fb:283:LYS:HE3	1.87	0.56
9:Fb:259:PHE:CE2	9:Fb:286:ASN:HB2	2.39	0.56
9:Fb:908:VAL:HA	9:Fb:911:PHE:CD2	2.39	0.56
9:Fb:933:ILE:H	9:Fb:933:ILE:HD12	1.71	0.56
9:Fd:175:SER:HB2	9:Fd:488:GLY:H	1.70	0.56
9:Ff:655:ILE:HG23	9:Ff:659:PHE:CE1	2.39	0.56
9:Ff:710:MET:HA	9:Ff:714:SER:OG	2.05	0.56
6:Al:203:ASN:ND2	7:Ef:817:THR:HG23	2.19	0.56
5:Aw:212:ILE:HB	5:Aw:264:PHE:CD1	2.41	0.56
5:Bc:220:ALA:HB3	5:Bc:232:VAL:HG23	1.87	0.56
5:Bc:222:LEU:HD11	5:Bc:262:ILE:HG21	1.88	0.56
1:Be:4:ALA:HB2	3:Bg:14:TYR:HB2	1.86	0.56
6:Bp:202:PHE:HB3	6:Bp:215:ILE:HD11	1.88	0.56
6:Cn:117:ARG:NH2	7:En:797:ARG:HB2	2.20	0.56
6:Ct:181:PHE:HE2	6:Ct:213:LEU:HD12	1.70	0.56
1:Dm:11:TYR:CD2	3:Do:9:TYR:HB3	2.40	0.56
7:Eg:804:ALA:O	7:Eg:808:LEU:HD22	2.06	0.56
7:Em:806:THR:HA	7:Em:809:VAL:HG12	1.87	0.56
7:Em:812:TRP:CD1	7:Em:812:TRP:N	2.74	0.56
7:Es:805:ALA:HA	7:Es:808:LEU:HG	1.88	0.56
7:Eu:782:ASN:O	7:Eu:785:LEU:HD12	2.05	0.56
8:Ew:298:TYR:O	8:Ew:301:LEU:HD12	2.05	0.56
8:Ey:69:PHE:CE2	8:Ey:402:GLN:HB3	2.41	0.56
8:Ey:381:ARG:HH12	8:Ey:382:ARG:HG3	1.69	0.56
8:Ez:82:TYR:CZ	8:Ez:86:LEU:HD11	2.40	0.56
8:Ez:82:TYR:O	8:Ez:86:LEU:HG	2.05	0.56
8:Ez:189:PHE:HA	8:Ez:295:LEU:HD21	1.87	0.56
8:Ez:336:TYR:O	8:Ez:340:THR:HG23	2.05	0.56
8:Ez:457:PHE:CZ	8:Fa:293:PRO:HB3	2.37	0.56
9:Fb:469:PHE:HE2	9:Fc:843:TYR:HB2	1.71	0.56
9:Fb:605:MET:HG3	9:Fb:638:ARG:HD2	1.88	0.56
9:Fb:734:LEU:HD11	9:Fb:738:TRP:CE2	2.40	0.56
9:Fc:90:GLU:HG3	9:Fc:92:GLN:H	1.69	0.56
9:Fc:674:GLY:HA2	9:Fc:692:MET:HE2	1.86	0.56
9:Fd:177:VAL:HA	9:Fd:491:LEU:HD22	1.87	0.56
9:Fd:593:LEU:HG	9:Fd:905:TRP:HD1	1.71	0.56
9:Fe:288:SER:HA	9:Fe:291:ASN:HB3	1.87	0.56

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
9:Fe:596:PHE:HE2	9:Fe:652:ILE:HG22	1.70	0.56
9:Fe:704:TRP:HE3	9:Fe:707:LEU:HD12	1.69	0.56
9:Fe:753:ILE:HG23	9:Fe:926:VAL:HG11	1.86	0.56
7:Fi:33:VAL:O	7:Fi:37:PHE:HD1	1.89	0.56
6:Al:115:MET:HE1	7:Ev:802:LEU:N	2.21	0.56
6:Ax:115:MET:HE1	7:Ef:801:MET:HB2	1.87	0.56
6:Ax:172:SER:HB3	6:Ax:175:PHE:HB2	1.87	0.56
5:Bo:238:ILE:HD11	5:Bo:244:VAL:HB	1.86	0.56
6:Bp:150:LYS:H	6:Bp:247:ALA:HA	1.70	0.56
6:Cb:181:PHE:CD2	6:Cb:191:ILE:HD11	2.40	0.56
4:Cf:2:PHE:HB3	5:Cg:251:GLN:HE21	1.71	0.56
5:Cg:207:ARG:HH12	5:Cg:240:GLY:HA3	1.71	0.56
6:Ct:170:ARG:HD2	6:Ct:247:ALA:HB3	1.87	0.56
6:Ct:214:MET:HB2	7:Ep:820:TYR:OH	2.06	0.56
7:Ef:781:GLN:NE2	7:Ef:785:LEU:HD11	2.19	0.56
7:Ej:772:GLU:CD	7:Ej:772:GLU:H	2.13	0.56
8:Ew:383:LEU:HG	8:Fa:82:TYR:CE2	2.40	0.56
8:Ew:451:ALA:HB1	9:Fd:619:PHE:HE1	1.71	0.56
8:Ex:104:ASP:HA	8:Ex:109:ASN:HD22	1.71	0.56
8:Ey:431:GLN:O	8:Ey:435:ARG:HG3	2.05	0.56
8:Ez:90:PRO:HB2	8:Fa:246:ASN:HD21	1.71	0.56
8:Ez:317:VAL:HG22	8:Fa:198:TRP:CD1	2.40	0.56
8:Ez:418:LEU:O	8:Ez:422:ILE:HG12	2.05	0.56
9:Fc:339:MET:O	9:Fc:343:LEU:HG	2.05	0.56
9:Ff:702:THR:O	9:Ff:706:THR:HG23	2.05	0.56
5:Ae:245:LYS:NZ	5:Ae:255:LEU:HB3	2.21	0.56
2:Bx:4:MET:SD	5:Ca:248:ASP:HA	2.45	0.56
3:By:2:VAL:HG11	6:Cn:129:LYS:HG2	1.87	0.56
6:Ct:221:TYR:OH	6:Cz:135:GLU:HG2	2.06	0.56
5:De:215:VAL:HG22	5:De:249:SER:HB3	1.87	0.56
6:Df:125:GLU:OE1	6:Df:129:LYS:HD3	2.04	0.56
6:Df:238:MET:HA	6:Df:238:MET:HE3	1.87	0.56
8:Ew:353:LYS:HG3	8:Ew:354:ARG:NE	2.21	0.56
8:Ex:327:SER:CB	8:Ey:211:MET:HE1	2.36	0.56
8:Ey:207:GLN:O	8:Ey:211:MET:HG3	2.06	0.56
9:Fc:188:GLN:O	9:Fc:192:LEU:HG	2.06	0.56
9:Ff:913:PHE:O	9:Ff:917:ILE:HG22	2.05	0.56
7:Fg:23:PHE:CE1	7:Fg:27:LEU:HD11	2.41	0.56
6:Bj:193:ALA:HB1	7:Ek:820:TYR:HE2	1.70	0.56
6:Bp:123:ASN:OD1	6:Bp:124:PRO:HD2	2.05	0.56
2:Br:2:VAL:HG22	3:Bs:1:ARG:NE	2.21	0.56

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
6:Cz:250:TYR:HE2	6:Cz:251:ARG:HH21	1.53	0.56
6:Df:115:MET:SD	7:Ep:802:LEU:HD22	2.44	0.56
6:Df:223:TYR:HB2	6:Df:242:ILE:HA	1.88	0.56
5:Ec:238:ILE:HD11	5:Ec:244:VAL:HG23	1.86	0.56
7:Ef:808:LEU:O	7:Ef:812:TRP:CD1	2.58	0.56
8:Ew:125:PHE:HZ	8:Ew:176:GLN:HG3	1.70	0.56
8:Ew:246:ASN:HD21	8:Ew:250:ALA:HB3	1.70	0.56
8:Ew:399:TRP:CD1	8:Ew:403:ILE:HD11	2.41	0.56
8:Ex:117:ASN:HB3	8:Ex:121:LYS:HD3	1.88	0.56
8:Ex:339:GLN:O	8:Ex:342:VAL:HG22	2.05	0.56
9:Fb:60:MET:O	9:Fb:64:PHE:HB3	2.06	0.56
9:Fb:69:LEU:HD11	7:Fh:38:LYS:HZ3	1.71	0.56
9:Fb:905:TRP:CZ2	9:Ff:705:LEU:HD21	2.39	0.56
9:Fc:176:GLY:HA3	9:Fc:492:ASP:HB3	1.88	0.56
9:Fe:465:GLY:HA2	9:Fe:819:TYR:HE1	1.70	0.56
9:Ff:188:GLN:O	9:Ff:192:LEU:HD13	2.05	0.56
9:Ff:628:MET:HE2	9:Ff:628:MET:N	2.20	0.56
6:Ar:129:LYS:HG3	6:Ax:112:PHE:CZ	2.40	0.56
6:Bj:135:GLU:HA	6:Bj:138:GLU:OE1	2.05	0.56
5:Ca:219:ARG:HG2	6:Ch:148:PRO:HD2	1.87	0.56
5:Cm:264:PHE:HB3	5:Cm:269:SER:HB3	1.85	0.56
6:Df:173:GLN:HB2	6:Df:220:LEU:HD23	1.88	0.56
6:Df:189:TRP:HE1	6:Df:232:GLY:H	1.52	0.56
2:Dh:2:VAL:HA	3:Di:1:ARG:HE	1.71	0.56
5:Dk:213:GLN:CD	6:Dl:168:VAL:HG21	2.31	0.56
5:Dw:212:ILE:HG13	5:Dw:262:ILE:HG22	1.87	0.56
8:Ew:185:PRO:HA	8:Ew:292:LEU:HD12	1.88	0.56
8:Ex:431:GLN:HB3	8:Ex:435:ARG:NH2	2.21	0.56
8:Ey:322:ALA:O	8:Ey:326:LEU:HG	2.06	0.56
9:Fb:199:GLN:HA	9:Fb:230:ILE:HD13	1.88	0.56
9:Fb:766:PHE:HB3	9:Fc:928:LYS:HZ2	1.71	0.56
9:Fe:113:LEU:CD1	9:Fe:124:MET:HB2	2.36	0.56
9:Fe:703:LEU:O	9:Fe:707:LEU:HG	2.06	0.56
6:Bv:126:GLN:HA	6:Bv:129:LYS:HG3	1.87	0.56
6:Cn:176:VAL:HG23	6:Cn:214:MET:HB2	1.88	0.56
5:De:215:VAL:HG11	5:De:252:GLY:HA2	1.87	0.56
6:Dl:221:TYR:CE1	6:Dr:142:ALA:HB3	2.41	0.56
5:Dw:233:ARG:HG2	5:Dw:234:GLU:H	1.70	0.56
7:Ej:789:LYS:HA	7:Ej:792:GLN:OE1	2.06	0.56
8:Ey:346:ASN:HA	8:Ey:435:ARG:NH1	2.20	0.56
8:Ez:426:MET:HA	8:Ez:429:ASP:CG	2.30	0.56

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
9:Fb:493:LYS:HZ2	9:Fc:597:LYS:HE3	1.71	0.56
9:Fb:547:GLN:HA	9:Fb:568:TYR:HE1	1.70	0.56
9:Fb:598:VAL:HG21	9:Fb:652:ILE:HG21	1.88	0.56
9:Fc:108:THR:HG22	7:Fg:27:LEU:HD21	1.87	0.56
9:Fc:128:PHE:O	9:Fc:132:ILE:HG12	2.05	0.56
9:Fc:645:LEU:HA	9:Fc:648:PHE:HB3	1.88	0.56
9:Fd:213:PRO:HB2	9:Fd:222:MET:CE	2.31	0.56
9:Fd:438:ARG:HH12	9:Fd:439:GLN:HG2	1.71	0.56
9:Fe:335:ALA:HB1	9:Fe:426:TYR:HE1	1.71	0.56
9:Fe:641:TYR:CE2	9:Fe:645:LEU:HD21	2.41	0.56
9:Fe:690:ALA:O	9:Fe:694:THR:HG23	2.06	0.56
9:Ff:524:SER:O	9:Ff:528:ILE:HG13	2.06	0.56
7:Fk:35:GLY:O	7:Fk:39:ILE:HG12	2.05	0.56
6:Af:130:LEU:HA	6:Af:133:ILE:HG22	1.86	0.56
6:Ar:123:ASN:HB3	6:Ar:126:GLN:HG3	1.86	0.56
6:Ax:229:ARG:NH1	6:Ax:236:PRO:HA	2.13	0.56
6:Bd:238:MET:N	6:Bj:251:ARG:HH12	2.04	0.56
6:Bp:238:MET:HG3	6:Bv:250:TYR:HE1	1.69	0.56
5:Cm:215:VAL:HG22	5:Cm:220:ALA:HB1	1.88	0.56
6:Cn:238:MET:N	6:Ct:251:ARG:HH12	2.02	0.56
6:Ct:111:ALA:HB1	7:En:798:THR:HG23	1.88	0.56
7:Ej:799:SER:HA	7:Ej:802:LEU:CD2	2.36	0.56
8:Ew:198:TRP:CH2	8:Fa:317:VAL:HA	2.41	0.56
8:Ez:424:TYR:HE2	8:Fa:380:THR:HA	1.70	0.56
9:Fb:220:PRO:HB2	9:Fc:367:LYS:NZ	2.21	0.56
9:Fc:674:GLY:O	9:Fc:678:LEU:HG	2.05	0.56
9:Fe:56:ILE:HB	9:Ff:820:ILE:HD11	1.86	0.56
9:Fe:162:ASP:HB3	9:Fe:165:LYS:HB2	1.88	0.56
9:Fe:636:VAL:O	9:Fe:640:VAL:HG23	2.05	0.56
9:Fe:768:TRP:C	9:Fe:768:TRP:CD1	2.84	0.56
9:Ff:339:MET:O	9:Ff:343:LEU:HG	2.05	0.56
9:Ff:523:LYS:HB2	9:Ff:526:LYS:NZ	2.21	0.56
9:Ff:735:LEU:O	9:Ff:739:ILE:HG12	2.04	0.56
9:Ff:917:ILE:HD12	9:Ff:921:MET:SD	2.45	0.56
5:Aq:244:VAL:HA	5:Aq:256:THR:HA	1.88	0.56
8:Ey:419:LEU:HA	8:Ey:422:ILE:HG12	1.87	0.56
9:Fb:692:MET:HE3	9:Fb:696:TYR:CE2	2.41	0.56
9:Fb:758:TYR:O	9:Fb:762:THR:HG23	2.06	0.56
9:Fc:704:TRP:HH2	9:Fd:733:PRO:HG2	1.70	0.56
9:Fc:760:ILE:HA	9:Fc:763:PHE:CD2	2.41	0.56
9:Fd:78:TYR:HA	9:Fd:81:MET:CE	2.36	0.56

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
9:Fd:567:VAL:O	9:Fd:571:VAL:HG23	2.06	0.56
9:Fe:110:GLY:HA3	9:Fe:787:THR:HG22	1.87	0.56
9:Fe:355:PRO:HG3	9:Fe:375:LYS:HG3	1.88	0.56
9:Fe:605:MET:HE3	9:Fe:605:MET:N	2.20	0.56
9:Ff:34:SER:O	9:Ff:38:LEU:HG	2.06	0.56
9:Ff:73:GLY:C	9:Ff:77:MET:HE2	2.30	0.56
5:Aw:221:TRP:HZ2	6:Bd:151:PRO:HG3	1.70	0.55
6:Bd:189:TRP:CG	6:Bd:230:LEU:HD13	2.40	0.55
4:Bh:2:PHE:HB2	5:Bi:251:GLN:NE2	2.21	0.55
6:Bj:189:TRP:NE1	6:Bj:232:GLY:H	2.04	0.55
5:Bo:221:TRP:CZ2	5:Bo:223:ILE:HD11	2.41	0.55
5:Ca:232:VAL:HG23	5:Ca:236:SER:HB3	1.88	0.55
6:Cb:196:LEU:HD21	6:Cb:199:PRO:HB3	1.88	0.55
6:Ch:121:PRO:HB2	7:Em:801:MET:HE1	1.88	0.55
5:Cm:219:ARG:HH11	5:Cm:233:ARG:HB3	1.70	0.55
6:Dr:115:MET:HA	6:Dr:118:ASN:OD1	2.06	0.55
7:Ej:776:ALA:O	7:Ej:780:LYS:HD2	2.06	0.55
8:Ew:430:ARG:HH22	8:Ex:428:LEU:HB2	1.70	0.55
8:Ex:351:LEU:HD12	8:Ex:352:SER:N	2.21	0.55
8:Ez:382:ARG:HB3	8:Ez:399:TRP:CD1	2.41	0.55
8:Fa:419:LEU:HA	8:Fa:422:ILE:CG1	2.27	0.55
9:Fb:41:LEU:HB2	9:Fc:828:VAL:HG11	1.88	0.55
9:Fe:940:VAL:HA	9:Fe:943:TRP:CE3	2.41	0.55
3:Ai:5:GLY:O	4:Ap:1:PRO:HB3	2.06	0.55
6:Bd:158:VAL:HB	6:Bd:256:VAL:HA	1.87	0.55
6:Bp:160:LEU:HD21	6:Bp:235:THR:HG23	1.88	0.55
1:Bw:2:THR:O	1:Bw:6:LEU:HD22	2.06	0.55
5:Cg:264:PHE:HB3	5:Cg:269:SER:HB3	1.88	0.55
6:Ch:185:THR:HB	6:Ch:259:TYR:CZ	2.42	0.55
6:Cz:115:MET:HE1	7:Eo:797:ARG:HH21	1.71	0.55
6:Cz:134:TYR:HE1	7:Ep:812:TRP:HB3	1.72	0.55
6:Dr:124:PRO:O	6:Dr:128:VAL:HG13	2.07	0.55
6:Dx:106:VAL:HA	6:Dx:109:LYS:NZ	2.22	0.55
6:Dx:117:ARG:HE	7:Et:797:ARG:HG3	1.71	0.55
7:Ej:798:THR:O	7:Ej:802:LEU:HD22	2.06	0.55
7:Et:780:LYS:HA	7:Et:783:GLU:CD	2.31	0.55
8:Ez:211:MET:HA	8:Ez:214:VAL:HB	1.87	0.55
8:Ez:331:ASN:HA	8:Ez:334:ARG:HG2	1.88	0.55
9:Fd:692:MET:O	9:Fd:696:TYR:HB2	2.07	0.55
9:Fe:185:LEU:HB2	9:Fe:343:LEU:HD13	1.88	0.55
9:Fe:540:LEU:HD23	9:Fe:562:PRO:HB2	1.86	0.55

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
7:Fj:16:ARG:HE	7:Fj:17:THR:HG22	1.71	0.55
5:Bc:245:LYS:N	5:Bc:245:LYS:HD2	2.21	0.55
6:Bj:196:LEU:HD21	6:Bj:199:PRO:HB3	1.86	0.55
6:Bp:219:LYS:HE3	6:Bp:221:TYR:H	1.71	0.55
6:Dl:112:PHE:O	6:Dl:116:THR:HG23	2.06	0.55
6:Dl:144:THR:HG21	6:Dl:148:PRO:HG3	1.87	0.55
6:Dl:179:LEU:HB3	6:Dl:181:PHE:CE1	2.42	0.55
6:Dx:111:ALA:HA	6:Dx:114:ASP:OD2	2.06	0.55
7:Es:788:GLN:HB3	7:Es:792:GLN:NE2	2.15	0.55
8:Ew:428:LEU:O	8:Ew:432:ILE:HG12	2.06	0.55
8:Ez:72:ASN:O	8:Ez:75:GLN:HG2	2.06	0.55
8:Ez:327:SER:HB3	8:Fa:215:ILE:HG21	1.88	0.55
8:Ez:360:SER:HB2	8:Ez:366:ALA:HA	1.88	0.55
9:Fc:143:TRP:CZ2	9:Fc:456:ALA:HB1	2.42	0.55
9:Fc:515:LEU:H	9:Fc:515:LEU:HD12	1.71	0.55
9:Fe:512:PRO:HG2	9:Fe:513:TYR:CZ	2.42	0.55
9:Fe:528:ILE:HA	9:Fe:531:GLN:HG2	1.87	0.55
6:Cb:191:ILE:HB	6:Cb:206:TRP:CZ2	2.41	0.55
6:Ch:199:PRO:HD2	7:Eo:816:GLU:OE1	2.06	0.55
6:Df:230:LEU:HD12	6:Df:231:ARG:H	1.71	0.55
5:Dk:244:VAL:HA	5:Dk:256:THR:HA	1.88	0.55
8:Ew:384:PHE:HB2	8:Fa:78:GLN:NE2	2.21	0.55
8:Fa:77:VAL:O	8:Fa:378:MET:HE1	2.06	0.55
9:Fb:675:VAL:HA	9:Fb:678:LEU:HG	1.89	0.55
9:Fb:747:PHE:HA	9:Fb:751:TYR:HB2	1.89	0.55
9:Fd:130:TRP:HD1	9:Fd:134:GLN:NE2	2.04	0.55
9:Fd:378:PHE:HB3	9:Fd:414:PHE:HD2	1.69	0.55
9:Fd:612:CYS:HB2	9:Fd:626:ARG:HE	1.71	0.55
9:Fd:909:TYR:CD2	9:Fd:912:PHE:HE2	2.24	0.55
9:Fe:729:MET:SD	9:Fe:730:MET:N	2.80	0.55
1:Am:11:TYR:CD2	3:Ao:9:TYR:HB3	2.42	0.55
6:Bd:106:VAL:HG23	6:Bd:110:LYS:NZ	2.21	0.55
6:Bj:125:GLU:O	6:Bj:128:VAL:HG22	2.07	0.55
6:Ch:125:GLU:O	6:Ch:129:LYS:HG3	2.06	0.55
6:Cn:242:ILE:HD12	6:Cn:243:PRO:HD2	1.88	0.55
6:Ct:106:VAL:HG23	6:Ct:110:LYS:NZ	2.20	0.55
6:Cz:181:PHE:HE2	6:Cz:191:ILE:HD11	1.71	0.55
6:Dr:112:PHE:HA	6:Dr:115:MET:SD	2.47	0.55
7:Eq:797:ARG:O	7:Eq:797:ARG:HD2	2.07	0.55
8:Ew:383:LEU:HG	8:Fa:82:TYR:HE2	1.71	0.55
8:Ey:111:LEU:HD12	8:Ey:112:ILE:HD13	1.89	0.55

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
8:Ey:430:ARG:NH2	8:Ez:428:LEU:HB2	2.21	0.55
9:Fb:138:ALA:O	9:Fb:142:ILE:HG12	2.06	0.55
9:Fc:663:PRO:HG3	9:Fc:703:LEU:HD11	1.88	0.55
9:Fd:129:MET:O	9:Fd:133:VAL:HG23	2.07	0.55
9:Fd:420:LEU:HD22	9:Fd:579:LEU:HD21	1.88	0.55
9:Fd:469:PHE:O	9:Fd:472:VAL:HB	2.07	0.55
5:Bi:255:LEU:HD13	5:Bi:261:VAL:HG13	1.89	0.55
6:Bj:221:TYR:CE2	6:Bp:142:ALA:HB3	2.42	0.55
6:Bj:223:TYR:HB2	6:Bj:242:ILE:HA	1.88	0.55
6:Bv:115:MET:SD	7:Ej:801:MET:SD	3.04	0.55
6:Cz:229:ARG:NH1	6:Df:212:THR:HG21	2.22	0.55
5:Dq:219:ARG:NH1	5:Dq:233:ARG:HB3	2.22	0.55
6:Dr:111:ALA:HB1	6:Dr:115:MET:HE1	1.89	0.55
7:Eu:798:THR:HA	7:Eu:801:MET:HG2	1.89	0.55
8:Ex:298:TYR:CZ	8:Ey:230:PHE:HE2	2.24	0.55
8:Ey:426:MET:C	8:Ey:426:MET:HE2	2.31	0.55
8:Ez:101:PHE:CE2	8:Fa:252:LEU:HB2	2.42	0.55
9:Fb:585:ILE:H	9:Fb:585:ILE:HD12	1.72	0.55
9:Fb:692:MET:HE3	9:Fb:696:TYR:HE2	1.72	0.55
9:Fc:78:TYR:HA	9:Fc:81:MET:HE3	1.89	0.55
9:Fc:741:THR:HG23	9:Fc:916:LEU:HD11	1.89	0.55
9:Fd:111:LEU:HD13	7:Fk:31:ALA:HB2	1.89	0.55
9:Fd:255:GLN:HG3	9:Fd:290:LEU:HD22	1.89	0.55
6:Af:148:PRO:HD2	5:Ec:218:GLY:HA3	1.89	0.55
5:Bo:264:PHE:HD2	6:Bp:246:LYS:HE2	1.72	0.55
6:Bv:201:SER:HA	6:Bv:219:LYS:HE3	1.88	0.55
5:Cg:219:ARG:H	6:Cn:148:PRO:HD2	1.72	0.55
5:Cm:219:ARG:HD3	6:Ct:246:LYS:NZ	2.22	0.55
5:Cs:213:GLN:HB2	5:Cs:223:ILE:HG22	1.89	0.55
6:Cz:169:ILE:HD13	6:Cz:241:LEU:HG	1.88	0.55
2:Dn:5:ILE:HD12	6:Dx:139:TYR:HB2	1.89	0.55
7:Eq:773:GLN:OE1	7:Eq:777:ILE:HD11	2.06	0.55
8:Ew:39:LEU:HD12	8:Ex:39:LEU:HD11	1.87	0.55
8:Ew:191:MET:HE3	8:Ew:192:ASP:H	1.71	0.55
8:Ew:191:MET:HE1	8:Ew:197:ASN:N	2.22	0.55
8:Fa:74:VAL:O	8:Fa:78:GLN:CB	2.55	0.55
9:Fb:757:PRO:HB3	9:Fb:930:PHE:CE1	2.42	0.55
9:Fc:342:THR:HG23	9:Fc:343:LEU:HD23	1.88	0.55
9:Fd:723:PHE:CD1	9:Fd:726:ALA:HB3	2.42	0.55
9:Fd:735:LEU:O	9:Fd:739:ILE:HG12	2.07	0.55
9:Fe:33:LEU:HD13	9:Fe:462:ILE:HD11	1.88	0.55

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
9:Fe:596:PHE:HD2	9:Fe:656:VAL:HG21	1.72	0.55
9:Fe:766:PHE:O	9:Fe:770:ILE:HG13	2.06	0.55
9:Ff:540:LEU:HD21	9:Ff:563:LEU:HD22	1.88	0.55
9:Ff:642:ASN:HA	9:Ff:645:LEU:HB3	1.89	0.55
7:Fg:34:ILE:HA	7:Fg:38:LYS:HZ1	1.72	0.55
3:Ac:2:VAL:HG21	6:Ar:129:LYS:HB3	1.89	0.55
5:Aq:248:ASP:HB3	5:Aq:253:ARG:HG2	1.87	0.55
5:Bc:245:LYS:HZ2	5:Bc:257:SER:HB3	1.72	0.55
5:Bu:215:VAL:HG11	5:Bu:252:GLY:HA2	1.89	0.55
5:De:211:TYR:HD2	5:De:263:LYS:O	1.90	0.55
6:Df:238:MET:HB3	6:Dl:250:TYR:HE1	1.72	0.55
6:Dx:216:GLN:NE2	7:Eu:817:THR:HA	2.19	0.55
7:Em:780:LYS:HG3	7:Em:781:GLN:NE2	2.22	0.55
7:Em:785:LEU:HA	7:Em:788:GLN:HB3	1.89	0.55
7:Eo:798:THR:HA	7:Eo:801:MET:HG2	1.88	0.55
8:Ex:193:ASN:HB2	8:Ex:194:GLU:OE2	2.07	0.55
8:Ey:304:GLN:HE22	8:Ey:319:GLN:N	2.05	0.55
8:Fa:118:THR:HA	8:Fa:121:LYS:HE3	1.88	0.55
8:Fa:348:TYR:HD1	8:Fa:351:LEU:HD23	1.70	0.55
9:Fc:223:ASN:O	9:Fc:227:ARG:HG2	2.07	0.55
9:Fc:528:ILE:HA	9:Fc:531:GLN:HG2	1.88	0.55
9:Fd:185:LEU:O	9:Fd:189:ILE:HG12	2.06	0.55
9:Fd:586:LYS:HZ3	9:Fd:588:LEU:HA	1.71	0.55
9:Fe:335:ALA:HB1	9:Fe:426:TYR:CE1	2.41	0.55
9:Ff:624:LEU:O	9:Ff:628:MET:HG2	2.07	0.55
9:Ff:915:ILE:O	9:Ff:919:THR:HG23	2.07	0.55
6:Af:199:PRO:HG3	7:Ef:819:VAL:HG13	1.88	0.55
6:Ar:126:GLN:HG2	6:Ar:129:LYS:NZ	2.21	0.55
6:Bj:145:PRO:HG3	6:Bp:131:LYS:HE2	1.88	0.55
6:Ct:108:ASP:HA	7:En:797:ARG:HH21	1.72	0.55
6:Ct:150:LYS:HG2	6:Ct:152:THR:HG23	1.87	0.55
6:Cz:191:ILE:HD12	6:Cz:206:TRP:NE1	2.21	0.55
6:Df:237:VAL:C	6:Dl:251:ARG:HH22	2.14	0.55
6:Dr:138:GLU:HA	6:Dr:141:LYS:HE2	1.87	0.55
6:Dr:226:LEU:HB2	6:Dr:241:LEU:HD11	1.88	0.55
7:Eq:797:ARG:HG3	7:Eq:801:MET:HE2	1.89	0.55
7:Er:794:ILE:HA	7:Er:797:ARG:CZ	2.37	0.55
8:Ex:431:GLN:O	8:Ex:435:ARG:HG2	2.06	0.55
9:Fb:144:GLU:HA	9:Fb:147:LEU:HG	1.88	0.55
9:Fc:467:TYR:O	9:Fc:470:ASP:HB2	2.06	0.55
9:Fc:773:ILE:O	9:Fc:777:VAL:HG13	2.07	0.55

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
9:Fd:237:VAL:HG21	9:Fd:263:MET:HB2	1.89	0.55
9:Fd:634:ASN:HA	9:Fd:638:ARG:HG3	1.89	0.55
9:Ff:77:MET:HE1	7:Fi:32:VAL:HG11	1.89	0.55
9:Ff:182:LEU:HD23	9:Ff:571:VAL:HG11	1.88	0.55
6:Al:118:ASN:O	6:Al:121:PRO:HD3	2.07	0.55
6:Bj:129:LYS:HE3	6:Bp:112:PHE:HE2	1.68	0.55
1:Bk:9:LEU:O	1:Bk:13:LYS:HG2	2.07	0.55
6:Bp:219:LYS:HD2	6:Bp:220:LEU:H	1.72	0.55
6:Cb:208:LYS:HG3	7:En:824:THR:H	1.71	0.55
6:Dr:205:GLN:HG2	7:Et:818:GLN:HE21	1.72	0.55
6:Dr:238:MET:CE	6:Dx:178:SER:HB3	2.37	0.55
6:Dx:136:THR:HA	6:Dx:139:TYR:CE1	2.42	0.55
6:Ed:108:ASP:O	6:Ed:112:PHE:HB2	2.07	0.55
7:Ek:804:ALA:O	7:Ek:807:GLN:HG3	2.06	0.55
8:Ew:189:PHE:HD2	8:Ew:295:LEU:HB2	1.71	0.55
8:Ew:210:VAL:O	8:Ew:214:VAL:HG23	2.07	0.55
8:Ew:318:GLN:H	8:Ew:318:GLN:CD	2.15	0.55
8:Ex:82:TYR:CB	8:Ey:383:LEU:HD21	2.35	0.55
8:Ex:441:SER:HB3	8:Ey:346:ASN:OD1	2.07	0.55
8:Ez:298:TYR:CZ	8:Fa:230:PHE:HE2	2.24	0.55
8:Fa:440:ASN:O	8:Fa:443:MET:HG2	2.07	0.55
9:Fb:617:ILE:HG12	9:Fc:617:ILE:HG22	1.89	0.55
9:Fb:828:VAL:O	9:Fb:832:ILE:HG12	2.07	0.55
9:Fb:925:ILE:HD12	9:Fb:925:ILE:H	1.71	0.55
9:Fc:196:LYS:HD3	9:Fc:197:GLN:HE21	1.72	0.55
9:Fe:392:TYR:HB2	9:Fe:576:MET:HE2	1.89	0.55
9:Fe:814:LEU:HD12	9:Fe:937:PRO:HB3	1.88	0.55
9:Fe:932:LEU:H	9:Fe:932:LEU:HD22	1.71	0.55
9:Ff:262:ASP:HB2	9:Ff:277:GLY:HA2	1.88	0.55
9:Ff:674:GLY:O	9:Ff:678:LEU:HG	2.06	0.55
6:Al:189:TRP:CD1	6:Al:230:LEU:HD11	2.43	0.54
6:Ar:112:PHE:O	6:Ar:116:THR:HG23	2.07	0.54
5:Bi:213:GLN:OE1	5:Bi:223:ILE:HG12	2.07	0.54
6:Bj:205:GLN:NE2	7:Ej:820:TYR:HB2	2.21	0.54
6:Df:199:PRO:HG3	7:Es:819:VAL:HG13	1.88	0.54
6:Df:208:LYS:HB3	7:Es:823:GLY:HA2	1.89	0.54
6:Dr:181:PHE:HE1	6:Dr:237:VAL:HG21	1.72	0.54
7:Eh:798:THR:O	7:Eh:802:LEU:HD22	2.07	0.54
7:Eq:807:GLN:HA	7:Eq:810:GLN:CD	2.32	0.54
8:Ew:174:VAL:O	8:Ew:178:VAL:HG23	2.08	0.54
8:Ew:238:PRO:O	8:Ew:242:GLN:HG2	2.07	0.54

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
8:Ey:82:TYR:O	8:Ey:86:LEU:HG	2.06	0.54
8:Ey:206:TYR:CZ	8:Ey:209:LEU:HB2	2.42	0.54
8:Fa:415:ILE:O	8:Fa:419:LEU:HG	2.07	0.54
9:Fb:642:ASN:O	9:Fb:645:LEU:HD12	2.08	0.54
9:Fc:381:PRO:HB3	9:Fc:577:VAL:HG12	1.89	0.54
9:Fd:147:LEU:HD11	9:Fd:457:ASN:HD21	1.72	0.54
9:Fd:530:ILE:O	9:Fd:534:ILE:HG12	2.06	0.54
9:Ff:290:LEU:HD22	9:Ff:329:ARG:NH2	2.22	0.54
9:Ff:927:GLN:O	9:Ff:931:THR:HG23	2.07	0.54
2:Ab:4:MET:HE1	5:Ae:249:SER:N	2.23	0.54
6:Af:106:VAL:HG23	6:Af:110:LYS:HZ3	1.71	0.54
2:Bf:4:MET:HE2	4:Bh:2:PHE:HE1	1.71	0.54
5:Bi:253:ARG:HB2	5:Bi:263:LYS:NZ	2.22	0.54
6:Bj:171:LEU:HD11	6:Bj:241:LEU:HB3	1.89	0.54
6:Bv:108:ASP:HA	7:Ej:797:ARG:HH12	1.73	0.54
6:Cb:206:TRP:HB2	6:Cb:213:LEU:HD23	1.89	0.54
6:Cn:238:MET:HG3	6:Ct:251:ARG:NH2	2.22	0.54
5:Cy:219:ARG:NH1	5:Cy:233:ARG:HB3	2.22	0.54
6:Dl:132:GLN:O	6:Dl:136:THR:HG23	2.07	0.54
6:Dl:141:LYS:HD3	6:Dl:141:LYS:C	2.32	0.54
6:Dl:237:VAL:C	6:Dr:251:ARG:HH22	2.15	0.54
6:Dx:168:VAL:HG13	6:Dx:242:ILE:HD13	1.88	0.54
6:Ed:189:TRP:CD1	6:Ed:230:LEU:HD23	2.43	0.54
7:Ei:789:LYS:HE2	9:Fd:267:ASP:N	2.23	0.54
8:Ew:144:GLN:HB2	8:Ew:179:PHE:CE1	2.42	0.54
8:Ey:205:MET:HE1	8:Ey:209:LEU:HG	1.90	0.54
8:Ey:281:ARG:C	8:Ey:286:GLN:HE21	2.16	0.54
8:Ey:429:ASP:O	8:Ey:433:GLN:HG2	2.06	0.54
9:Fb:616:LYS:HD2	9:Fb:621:SER:HB3	1.88	0.54
9:Ff:94:LEU:HD13	9:Ff:98:TRP:HD1	1.72	0.54
1:As:9:LEU:HB2	2:At:2:VAL:HG21	1.89	0.54
6:Ax:219:LYS:HD2	6:Ax:222:ASN:ND2	2.23	0.54
5:Bo:215:VAL:HG22	5:Bo:249:SER:HB3	1.90	0.54
6:Dr:206:TRP:HB2	6:Dr:213:LEU:HD23	1.90	0.54
7:Eh:778:LEU:O	7:Eh:781:GLN:HG2	2.06	0.54
7:El:797:ARG:HA	7:El:800:ASP:OD2	2.08	0.54
8:Ey:277:LEU:HD12	8:Ey:278:ASN:N	2.21	0.54
8:Ez:330:PHE:HB3	8:Ez:334:ARG:NH2	2.22	0.54
9:Fb:643:PHE:O	9:Fb:647:ILE:HG12	2.07	0.54
9:Fc:86:ASN:HA	9:Fc:89:HIS:CD2	2.35	0.54
9:Fd:39:GLY:HA2	9:Fd:49:LEU:HD21	1.90	0.54

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
9:Fe:187:GLY:O	9:Fe:191:MET:HE2	2.06	0.54
9:Ff:117:LYS:HZ3	9:Ff:119:SER:N	2.06	0.54
6:Af:216:GLN:HG2	7:Ee:818:GLN:HG2	1.89	0.54
1:Ag:6:LEU:HA	1:Ag:9:LEU:HG	1.90	0.54
5:Ak:253:ARG:C	5:Ak:254:ILE:HD13	2.32	0.54
6:Ar:204:ILE:HG23	6:Ar:215:ILE:HG12	1.89	0.54
5:Bo:207:ARG:HH12	5:Bo:209:ILE:HG13	1.72	0.54
6:Cb:151:PRO:HA	6:Cb:248:VAL:HG13	1.89	0.54
6:Ch:228:VAL:HB	6:Ch:237:VAL:HG13	1.88	0.54
5:Cm:247:ILE:HG23	5:Cm:254:ILE:HD12	1.88	0.54
6:Cn:216:GLN:HG2	7:Eo:818:GLN:NE2	2.23	0.54
5:Cy:221:TRP:CZ3	5:Cy:229:THR:HG22	2.43	0.54
5:De:212:ILE:HG21	5:De:215:VAL:HB	1.89	0.54
3:Do:2:VAL:HB	6:Ed:129:LYS:HB3	1.88	0.54
8:Ex:338:ALA:O	8:Ex:342:VAL:HG13	2.06	0.54
8:Ex:430:ARG:NH1	8:Ey:428:LEU:HB2	2.23	0.54
8:Ez:209:LEU:HD23	8:Ez:213:ASN:HD21	1.72	0.54
8:Ez:430:ARG:NH2	8:Fa:425:GLN:HG3	2.22	0.54
8:Fa:174:VAL:O	8:Fa:178:VAL:HG23	2.08	0.54
8:Fa:249:ILE:C	8:Fa:354:ARG:HH12	2.16	0.54
9:Fb:234:ILE:H	9:Fb:234:ILE:HD12	1.73	0.54
9:Fc:630:ASP:HA	9:Fc:634:ASN:HB2	1.88	0.54
9:Fd:745:ILE:CG2	9:Fd:919:THR:HG21	2.33	0.54
9:Fe:760:ILE:HA	9:Fe:763:PHE:CE1	2.43	0.54
9:Fe:828:VAL:O	9:Fe:832:ILE:HG12	2.08	0.54
6:Ax:111:ALA:HB3	7:Ef:797:ARG:NH1	2.15	0.54
6:Ax:170:ARG:HH11	6:Ax:247:ALA:HB1	1.72	0.54
2:Az:5:ILE:HG21	6:Bj:140:ALA:HB2	1.88	0.54
6:Ch:153:ALA:HB1	6:Ch:251:ARG:NE	2.23	0.54
1:Co:12:HIS:CE1	3:Cq:9:TYR:HB2	2.43	0.54
6:Dr:168:VAL:HG23	6:Dr:242:ILE:HD13	1.90	0.54
5:Dw:215:VAL:HG11	5:Dw:252:GLY:HA2	1.89	0.54
8:Ew:68:LEU:HD21	8:Ew:74:VAL:HG21	1.89	0.54
8:Ew:240:ILE:HG23	8:Fa:94:MET:HE3	1.87	0.54
8:Ey:172:ASP:HB3	8:Ey:175:SER:OG	2.07	0.54
9:Fc:426:TYR:O	9:Fc:430:MET:HG2	2.08	0.54
9:Fe:606:LYS:HD3	9:Fe:606:LYS:N	2.23	0.54
9:Fe:733:PRO:HA	9:Fe:736:MET:HE2	1.90	0.54
9:Fe:936:LEU:O	9:Fe:940:VAL:HG23	2.07	0.54
9:Ff:663:PRO:HG2	9:Ff:738:TRP:HE3	1.73	0.54
9:Ff:717:ILE:HD11	9:Ff:720:PHE:HB2	1.89	0.54

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
6:Al:220:LEU:HD21	6:Ar:138:GLU:OE1	2.07	0.54
6:Ar:126:GLN:HG2	6:Ar:129:LYS:HZ1	1.72	0.54
6:Bd:238:MET:HG2	6:Bj:251:ARG:CZ	2.38	0.54
5:Cs:263:LYS:HD2	5:Cs:263:LYS:C	2.33	0.54
6:Dl:189:TRP:HE1	6:Dl:232:GLY:H	1.54	0.54
6:Dx:189:TRP:NE1	6:Dx:232:GLY:H	2.01	0.54
6:Ed:125:GLU:C	6:Ed:129:LYS:HE3	2.33	0.54
6:Ed:171:LEU:HD23	6:Ed:217:ALA:HB2	1.88	0.54
8:Ew:327:SER:HB2	8:Ex:215:ILE:HD11	1.90	0.54
8:Ew:367:ASN:C	8:Ew:368:ILE:HD13	2.32	0.54
8:Ez:170:LEU:HD23	8:Ez:171:ASN:N	2.22	0.54
8:Ez:206:TYR:CE2	8:Ez:209:LEU:HB2	2.41	0.54
8:Fa:230:PHE:HA	8:Fa:235:TYR:CD2	2.43	0.54
9:Fb:66:SER:O	9:Fb:69:LEU:HD12	2.08	0.54
9:Fd:144:GLU:HA	9:Fd:147:LEU:HG	1.89	0.54
9:Fd:378:PHE:HB3	9:Fd:414:PHE:CD2	2.41	0.54
9:Fd:670:ILE:HD12	9:Fd:696:TYR:CD2	2.43	0.54
9:Ff:129:MET:O	9:Ff:133:VAL:HG13	2.07	0.54
9:Ff:185:LEU:HA	9:Ff:188:GLN:HE21	1.71	0.54
9:Ff:811:ARG:HH21	9:Ff:937:PRO:HB2	1.72	0.54
5:Ak:219:ARG:HH21	5:Ak:231:THR:HG23	1.72	0.54
6:Cn:115:MET:SD	7:Em:802:LEU:HD13	2.47	0.54
6:Ct:107:ILE:HG13	6:Ct:110:LYS:HZ1	1.72	0.54
6:Cz:219:LYS:HD2	6:Cz:222:ASN:OD1	2.07	0.54
6:Df:220:LEU:HD12	6:Dl:138:GLU:OE2	2.08	0.54
7:Ek:799:SER:O	7:Ek:803:THR:HG23	2.08	0.54
7:Eq:802:LEU:O	7:Eq:806:THR:HG23	2.07	0.54
8:Ew:424:TYR:HE2	8:Ex:380:THR:HA	1.73	0.54
8:Ex:327:SER:HB2	8:Ey:211:MET:HE1	1.90	0.54
8:Ex:426:MET:HA	8:Ex:429:ASP:OD1	2.08	0.54
9:Fb:936:LEU:O	9:Fb:940:VAL:HG13	2.08	0.54
9:Fd:447:LYS:HD3	9:Fd:447:LYS:N	2.21	0.54
9:Fe:534:ILE:HG22	9:Fe:568:TYR:CE1	2.42	0.54
3:Ai:11:ALA:HB1	3:Ai:14:TYR:HE2	1.73	0.54
6:Ar:112:PHE:O	6:Ar:115:MET:HG2	2.07	0.54
2:Cv:4:MET:HG2	5:Cy:250:LEU:HD12	1.90	0.54
2:Dh:2:VAL:HG22	3:Di:1:ARG:HE	1.72	0.54
7:Ei:789:LYS:HE2	9:Fd:267:ASP:H	1.73	0.54
8:Ew:67:GLN:HA	8:Ew:413:LYS:NZ	2.23	0.54
8:Ew:427:TYR:CD2	8:Ex:379:ALA:HB2	2.43	0.54
8:Ey:236:ASN:O	8:Ey:240:ILE:HG12	2.08	0.54

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
8:Fa:218:LEU:HD23	8:Fa:218:LEU:H	1.72	0.54
9:Fb:905:TRP:HA	9:Fb:908:VAL:HG22	1.90	0.54
9:Fc:345:THR:O	9:Fc:348:GLN:HG3	2.08	0.54
9:Fc:652:ILE:O	9:Fc:656:VAL:HG23	2.07	0.54
9:Fd:370:PHE:HE2	9:Fd:565:SER:HB3	1.73	0.54
9:Fe:49:LEU:HD22	9:Fe:50:HIS:H	1.72	0.54
9:Fe:342:THR:HG23	9:Fe:343:LEU:HD23	1.90	0.54
9:Fe:605:MET:H	9:Fe:605:MET:CE	2.21	0.54
6:Ax:115:MET:HE1	7:Ef:802:LEU:N	2.23	0.54
5:Dk:211:TYR:CD2	5:Dk:265:SER:HA	2.43	0.54
6:Dx:208:LYS:HB3	7:Ev:824:THR:HG23	1.88	0.54
6:Ed:115:MET:HA	6:Ed:118:ASN:OD1	2.07	0.54
8:Ew:415:ILE:HG22	8:Ew:419:LEU:HD21	1.90	0.54
8:Ez:268:THR:HG22	8:Ez:270:LYS:HZ1	1.72	0.54
9:Fb:56:ILE:H	9:Fb:56:ILE:HD12	1.73	0.54
9:Fb:378:PHE:HB3	9:Fb:414:PHE:CE1	2.43	0.54
9:Fb:800:PHE:HA	9:Fb:803:MET:SD	2.48	0.54
9:Fd:33:LEU:HA	9:Fd:36:VAL:HB	1.90	0.54
9:Fd:290:LEU:HD21	9:Fd:329:ARG:HD3	1.88	0.54
9:Fd:595:ASN:HB3	9:Fd:597:LYS:HE3	1.88	0.54
9:Fe:143:TRP:O	9:Fe:147:LEU:HG	2.07	0.54
9:Fe:179:LYS:HG2	9:Fe:492:ASP:HA	1.90	0.54
9:Fe:939:LYS:HA	9:Fe:942:ARG:HB2	1.90	0.54
9:Ff:608:GLN:HG3	9:Ff:633:TYR:HE2	1.72	0.54
2:Ab:3:SER:CB	6:Ar:128:VAL:HG21	2.38	0.54
6:Bd:226:LEU:HB3	6:Bd:241:LEU:HD21	1.89	0.54
3:Bm:11:ALA:HB1	3:Bm:14:TYR:CE2	2.41	0.54
6:Ch:115:MET:HE1	7:El:801:MET:CG	2.37	0.54
6:Cz:178:SER:HB3	6:Cz:251:ARG:HD3	1.90	0.54
6:Dx:193:ALA:HB1	7:Ev:820:TYR:HE1	1.73	0.54
7:Eh:782:ASN:HA	7:Eh:785:LEU:HD12	1.90	0.54
7:Em:782:ASN:HA	7:Em:785:LEU:HD23	1.90	0.54
7:Eo:799:SER:HA	7:Eo:802:LEU:HD23	1.88	0.54
8:Ex:211:MET:O	8:Ex:215:ILE:HD12	2.08	0.54
8:Ex:301:LEU:CG	8:Ex:326:LEU:HD21	2.32	0.54
8:Ez:180:ASN:HD22	8:Ez:286:GLN:HE21	1.54	0.54
8:Ez:433:GLN:HA	8:Ez:436:ILE:CG1	2.29	0.54
8:Fa:81:ALA:HB2	8:Fa:378:MET:HE2	1.90	0.54
9:Fb:202:ARG:HD2	9:Fb:225:PHE:CE2	2.43	0.54
9:Fb:418:GLU:CD	9:Fb:418:GLU:H	2.15	0.54
9:Fb:500:GLN:HE21	9:Fb:503:LYS:HE2	1.73	0.54

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
9:Fc:768:TRP:O	9:Fc:772:VAL:HG23	2.07	0.54
9:Fd:377:GLN:OE1	9:Fd:402:VAL:HG12	2.07	0.54
9:Fd:465:GLY:HA2	9:Fd:819:TYR:CZ	2.43	0.54
9:Fe:530:ILE:O	9:Fe:534:ILE:HG13	2.07	0.54
9:Ff:219:THR:HG22	9:Ff:222:MET:HE3	1.89	0.54
6:Bd:114:ASP:O	6:Bd:117:ARG:HB3	2.08	0.53
6:Ch:218:THR:O	7:En:815:VAL:HG21	2.08	0.53
6:Df:238:MET:HG3	6:Dl:250:TYR:CD1	2.42	0.53
3:Du:2:VAL:HG22	3:Du:8:VAL:HG22	1.90	0.53
2:Dz:2:VAL:HA	3:Ea:1:ARG:HE	1.73	0.53
7:Eo:799:SER:HA	7:Eo:802:LEU:CD2	2.37	0.53
8:Ew:252:LEU:HD11	8:Ew:373:LEU:HA	1.91	0.53
8:Ex:208:ASN:O	8:Ex:212:GLN:HG2	2.08	0.53
8:Ey:43:LEU:HD21	8:Ez:39:LEU:HD13	1.89	0.53
8:Ez:346:ASN:O	8:Ez:350:ILE:HG13	2.08	0.53
9:Fc:718:PRO:HG3	9:Fd:720:PHE:HE1	1.73	0.53
9:Fc:800:PHE:O	9:Fc:804:ILE:HD12	2.08	0.53
9:Fd:528:ILE:HA	9:Fd:531:GLN:HG3	1.90	0.53
9:Fe:766:PHE:O	9:Fe:769:LEU:HG	2.08	0.53
9:Ff:336:ILE:HA	9:Ff:339:MET:HG2	1.91	0.53
6:Af:129:LYS:CD	3:Du:2:VAL:HG21	2.39	0.53
4:Bb:5:ASP:HA	5:Bc:251:GLN:HG2	1.90	0.53
6:Df:111:ALA:HB3	7:Ep:797:ARG:NH2	2.24	0.53
6:Ed:125:GLU:O	6:Ed:129:LYS:HG2	2.08	0.53
7:Eu:810:GLN:HA	7:Eu:813:LYS:HG3	1.91	0.53
8:Ew:411:VAL:O	8:Ew:415:ILE:HG13	2.08	0.53
8:Ex:35:ASN:O	8:Ex:39:LEU:HD22	2.08	0.53
8:Ex:142:ILE:HD13	8:Ex:281:ARG:HD3	1.90	0.53
8:Ex:194:GLU:HB2	8:Ex:196:LYS:HD3	1.89	0.53
8:Ez:174:VAL:HG13	8:Ez:344:VAL:HG23	1.90	0.53
8:Fa:424:TYR:O	8:Fa:427:TYR:HB3	2.07	0.53
9:Fb:469:PHE:O	9:Fb:473:LYS:HG2	2.08	0.53
9:Fb:664:LEU:HD21	9:Fb:738:TRP:CE2	2.43	0.53
9:Fc:146:ALA:O	9:Fc:150:LEU:HD22	2.08	0.53
9:Fc:673:VAL:HA	9:Fc:676:GLN:HG3	1.90	0.53
9:Fd:334:ILE:O	9:Fd:338:GLN:HG2	2.06	0.53
9:Fd:617:ILE:HB	9:Fd:620:PHE:HB3	1.89	0.53
9:Fd:617:ILE:HD13	9:Fd:620:PHE:CD2	2.44	0.53
9:Fe:132:ILE:HG21	9:Fe:772:VAL:HG23	1.90	0.53
9:Ff:930:PHE:O	9:Ff:933:ILE:HG22	2.07	0.53
5:Ae:207:ARG:HH12	5:Ae:209:ILE:HA	1.73	0.53

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:Ai:11:ALA:HB1	3:Ai:14:TYR:CE2	2.44	0.53
6:Ar:134:TYR:HE1	7:Ef:812:TRP:HB2	1.74	0.53
1:Bq:3:ASP:HA	1:Bq:6:LEU:HG	1.89	0.53
3:By:1:ARG:HG2	3:By:2:VAL:N	2.22	0.53
6:Cb:194:TYR:CD2	6:Cb:213:LEU:HD21	2.43	0.53
5:Cg:218:GLY:HA3	6:Cn:148:PRO:HD3	1.89	0.53
6:Cn:199:PRO:HG3	7:Ep:819:VAL:HG13	1.89	0.53
1:Co:12:HIS:NE2	3:Cq:9:TYR:HB2	2.23	0.53
6:Cz:128:VAL:HA	6:Cz:131:LYS:HG2	1.90	0.53
6:Cz:225:ASN:HB2	6:Cz:238:MET:CE	2.37	0.53
5:De:243:MET:HE2	5:De:243:MET:HA	1.90	0.53
6:Dl:179:LEU:HD13	6:Dl:252:VAL:HB	1.90	0.53
7:Eq:807:GLN:O	7:Eq:810:GLN:HG2	2.08	0.53
7:Eu:779:GLN:O	7:Eu:783:GLU:HG3	2.08	0.53
8:Ew:172:ASP:HB3	8:Ew:175:SER:OG	2.08	0.53
8:Ez:104:ASP:HA	8:Ez:109:ASN:ND2	2.23	0.53
8:Ez:418:LEU:HD12	8:Ez:419:LEU:H	1.73	0.53
9:Fc:708:LEU:O	9:Fc:713:VAL:HG23	2.07	0.53
9:Fc:937:PRO:O	9:Fc:940:VAL:HB	2.08	0.53
9:Ff:644:PHE:HB2	9:Ff:720:PHE:CE1	2.43	0.53
9:Ff:833:LEU:HD12	9:Ff:837:PHE:HB3	1.89	0.53
2:Ab:4:MET:HE1	5:Ae:250:LEU:H	1.74	0.53
6:Bj:208:LYS:HB3	7:Ek:823:GLY:HA2	1.90	0.53
5:Cm:219:ARG:H	6:Ct:148:PRO:HD2	1.74	0.53
5:Cm:253:ARG:C	5:Cm:254:ILE:HD13	2.34	0.53
6:Cn:130:LEU:HG	7:En:812:TRP:CH2	2.44	0.53
5:Dk:213:GLN:NE2	6:Dl:168:VAL:HG21	2.23	0.53
6:Dl:200:SER:HB3	7:Et:816:GLU:HG2	1.91	0.53
7:Eq:809:VAL:HA	7:Eq:812:TRP:CD1	2.38	0.53
8:Ey:410:THR:O	8:Ey:414:GLU:HG3	2.08	0.53
8:Ez:112:ILE:HD12	8:Ez:112:ILE:H	1.72	0.53
9:Fc:718:PRO:HG3	9:Fd:720:PHE:CE1	2.44	0.53
9:Fd:507:LYS:HZ1	9:Fe:630:ASP:HB3	1.73	0.53
9:Fd:644:PHE:HB2	9:Fd:720:PHE:HE2	1.73	0.53
9:Fe:434:LEU:HA	9:Fe:437:ILE:HG12	1.90	0.53
9:Ff:247:THR:HG22	9:Ff:256:PRO:HD2	1.89	0.53
9:Ff:370:PHE:HB2	9:Ff:410:THR:HG21	1.91	0.53
5:Ca:253:ARG:HB3	5:Ca:253:ARG:NH1	2.23	0.53
6:Dl:131:LYS:O	6:Dl:135:GLU:HG2	2.08	0.53
6:Dl:158:VAL:HG13	6:Dl:167:PRO:HG2	1.90	0.53
6:Dl:176:VAL:HG13	6:Dl:214:MET:HG2	1.90	0.53

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
7:Ek:794:ILE:HA	7:Ek:797:ARG:CZ	2.38	0.53
8:Ey:381:ARG:HH12	8:Ey:382:ARG:HH11	1.55	0.53
8:Ez:211:MET:O	8:Ez:215:ILE:HG12	2.09	0.53
9:Fb:464:ALA:HA	9:Fb:467:TYR:HD2	1.74	0.53
9:Fb:595:ASN:HB3	9:Fb:597:LYS:HE2	1.90	0.53
9:Fe:748:VAL:O	9:Fe:753:ILE:HG12	2.09	0.53
9:Fe:755:VAL:O	9:Fe:758:TYR:HB3	2.09	0.53
9:Ff:685:PRO:O	9:Ff:689:LEU:HG	2.08	0.53
2:Ab:5:ILE:HD12	6:Al:139:TYR:CD2	2.38	0.53
6:Af:144:THR:HG21	6:Af:148:PRO:HG3	1.91	0.53
6:Af:145:PRO:HD3	6:Al:131:LYS:HE2	1.91	0.53
6:Ar:171:LEU:HD13	6:Ar:202:PHE:HZ	1.73	0.53
5:Bo:261:VAL:HG12	5:Bo:263:LYS:NZ	2.23	0.53
2:Cd:2:VAL:HG22	3:Ce:1:ARG:NE	2.23	0.53
6:Ch:201:SER:HA	6:Ch:219:LYS:NZ	2.24	0.53
5:Dk:211:TYR:CE2	5:Dk:265:SER:HA	2.44	0.53
6:Dr:223:TYR:HB2	6:Dr:242:ILE:HA	1.90	0.53
7:Ev:802:LEU:O	7:Ev:806:THR:HG23	2.08	0.53
8:Ew:218:LEU:HD12	8:Ew:219:PRO:HD2	1.91	0.53
8:Ez:444:LEU:HD13	8:Ez:448:LEU:HD23	1.91	0.53
9:Fc:615:VAL:HG22	9:Fc:622:PHE:O	2.09	0.53
9:Fe:73:GLY:C	9:Fe:77:MET:HE2	2.33	0.53
6:Ar:206:TRP:HB2	6:Ar:213:LEU:HD23	1.90	0.53
6:Ar:214:MET:HE2	7:Eg:820:TYR:OH	2.09	0.53
6:Ar:230:LEU:HB2	6:Ar:233:LEU:HD23	1.91	0.53
6:Bj:150:LYS:HG2	6:Bj:152:THR:HG23	1.90	0.53
5:Dk:207:ARG:CZ	5:Dk:208:ILE:H	2.21	0.53
6:Dr:225:ASN:HD21	6:Dx:176:VAL:N	2.06	0.53
6:Dx:182:LEU:HD12	6:Dx:183:ASP:N	2.24	0.53
6:Ed:115:MET:HG2	7:Et:801:MET:HE3	1.90	0.53
7:Ej:789:LYS:O	7:Ej:793:GLU:HG2	2.08	0.53
8:Ew:324:ALA:HB2	8:Ex:208:ASN:ND2	2.22	0.53
8:Ex:36:THR:HA	8:Ex:39:LEU:HD23	1.91	0.53
8:Ey:62:ASN:HA	8:Ez:48:LYS:NZ	2.24	0.53
8:Ez:411:VAL:HA	8:Ez:414:GLU:HG2	1.91	0.53
9:Fb:369:ASP:HB3	9:Fb:375:LYS:HD2	1.91	0.53
9:Fc:381:PRO:HD2	9:Fc:573:ASN:ND2	2.23	0.53
9:Fc:698:ASN:HA	9:Fd:905:TRP:CZ3	2.44	0.53
9:Fd:939:LYS:HB3	9:Fd:942:ARG:NH2	2.22	0.53
9:Ff:758:TYR:O	9:Ff:762:THR:HG23	2.09	0.53
9:Ff:924:ILE:HA	9:Ff:927:GLN:HG2	1.91	0.53

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
6:Al:121:PRO:HB3	7:Ee:797:ARG:HH21	1.72	0.53
6:Al:238:MET:N	6:Ar:251:ARG:HH12	2.06	0.53
6:Ar:112:PHE:HA	6:Ar:115:MET:SD	2.49	0.53
6:Ar:225:ASN:HD21	6:Ax:176:VAL:N	2.07	0.53
1:Bk:16:ALA:HB3	2:Br:9:ARG:HG2	1.90	0.53
5:Bo:253:ARG:C	5:Bo:254:ILE:HD13	2.34	0.53
6:Cn:158:VAL:HB	6:Cn:256:VAL:HA	1.91	0.53
6:Df:108:ASP:HA	7:Ep:797:ARG:HH22	1.74	0.53
1:Dm:6:LEU:HB3	5:Dq:246:LEU:HD21	1.91	0.53
6:Ed:129:LYS:HA	6:Ed:132:GLN:HG3	1.90	0.53
7:Et:785:LEU:HD23	9:Ff:241:LYS:HG2	1.91	0.53
8:Ew:301:LEU:HD13	8:Ew:326:LEU:CD2	2.39	0.53
8:Ew:317:VAL:HG12	8:Ex:198:TRP:CD1	2.43	0.53
8:Ey:342:VAL:HG21	8:Ey:442:ILE:HG13	1.91	0.53
8:Ez:420:ALA:O	8:Ez:424:TYR:HD1	1.92	0.53
8:Fa:421:GLU:O	8:Fa:425:GLN:CB	2.52	0.53
9:Fb:339:MET:O	9:Fb:343:LEU:HG	2.08	0.53
9:Fb:501:LEU:H	9:Fb:501:LEU:HD12	1.73	0.53
9:Fb:648:PHE:O	9:Fb:652:ILE:HG13	2.09	0.53
9:Fc:197:GLN:HG3	9:Fc:530:ILE:HD11	1.90	0.53
9:Ff:693:GLY:HA2	9:Ff:696:TYR:CD2	2.43	0.53
5:Ae:246:LEU:CD1	5:Ae:255:LEU:HB2	2.39	0.53
1:Be:4:ALA:HB3	2:Bf:9:ARG:NH1	2.24	0.53
6:Bp:238:MET:HE3	6:Bv:251:ARG:HG3	1.90	0.53
5:Bu:219:ARG:HH21	5:Bu:231:THR:HG23	1.72	0.53
4:Bz:5:ASP:HA	5:Ca:251:GLN:HG2	1.91	0.53
6:Cb:166:PRO:HD2	6:Ch:151:PRO:HB2	1.91	0.53
6:Dl:141:LYS:HD3	6:Dl:141:LYS:O	2.09	0.53
7:Eg:810:GLN:HA	7:Eg:813:LYS:NZ	2.24	0.53
8:Ew:142:ILE:HA	8:Ew:210:VAL:HG23	1.90	0.53
8:Ew:234:LYS:HA	8:Ew:237:GLN:NE2	2.24	0.53
8:Ew:242:GLN:HE22	8:Ew:269:ALA:H	1.55	0.53
8:Ew:428:LEU:CB	8:Fa:430:ARG:HH22	1.99	0.53
8:Ez:254:MET:HE2	8:Ez:254:MET:N	2.24	0.53
8:Fa:210:VAL:O	8:Fa:214:VAL:HG23	2.09	0.53
8:Fa:444:LEU:HA	8:Fa:447:ASN:ND2	2.24	0.53
9:Fb:455:GLU:O	9:Fb:459:LYS:HG2	2.09	0.53
9:Fc:475:ASN:HB3	9:Fc:684:ASN:HD21	1.74	0.53
9:Fd:57:MET:HB3	9:Fd:61:PHE:CE1	2.44	0.53
9:Fe:65:ASN:HD21	9:Fe:116:PRO:HD3	1.73	0.53
9:Fe:78:TYR:HA	9:Fe:81:MET:HE2	1.91	0.53

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
9:Fe:514:SER:O	9:Fe:518:THR:HG23	2.09	0.53
9:Fe:524:SER:O	9:Fe:528:ILE:HG22	2.09	0.53
9:Fe:527:LEU:H	9:Fe:527:LEU:HD12	1.74	0.53
9:Ff:810:LEU:HB3	9:Ff:814:LEU:HD21	1.91	0.53
9:Ff:917:ILE:O	9:Ff:921:MET:HG2	2.09	0.53
6:Ar:205:GLN:HG3	7:Eg:818:GLN:HE21	1.74	0.53
6:Ax:233:LEU:HD11	6:Ax:235:THR:HB	1.91	0.53
6:Cn:133:ILE:HD13	6:Ct:116:THR:HG22	1.89	0.53
6:Df:108:ASP:HA	7:Ep:797:ARG:NH2	2.24	0.53
6:Df:208:LYS:HB2	7:Es:824:THR:HG23	1.91	0.53
6:Df:238:MET:HG3	6:Dl:250:TYR:HD1	1.74	0.53
6:Dl:122:LEU:HD21	7:Er:801:MET:HB2	1.90	0.53
6:Dr:230:LEU:HB2	6:Dr:233:LEU:HD23	1.91	0.53
7:Ef:816:GLU:H	7:Ef:816:GLU:CD	2.17	0.53
7:Ev:809:VAL:HG13	7:Ev:810:GLN:HE21	1.73	0.53
8:Ex:77:VAL:HG12	8:Ex:378:MET:CE	2.38	0.53
8:Ex:235:TYR:O	8:Ex:238:PRO:HD2	2.09	0.53
8:Ey:186:ASP:HA	8:Ey:207:GLN:OE1	2.09	0.53
8:Ez:335:VAL:O	8:Ez:339:GLN:HG3	2.09	0.53
8:Fa:136:ILE:HD12	8:Fa:277:LEU:HD12	1.91	0.53
9:Fd:623:CYS:HB3	9:Fd:626:ARG:HB3	1.90	0.53
9:Ff:34:SER:HA	9:Ff:462:ILE:HD13	1.91	0.53
9:Ff:35:VAL:HA	9:Ff:38:LEU:HD12	1.90	0.53
9:Ff:432:PRO:O	9:Ff:436:LEU:HD22	2.09	0.53
3:Ao:1:ARG:HA	6:Bd:125:GLU:OE1	2.08	0.52
5:Bi:253:ARG:C	5:Bi:254:ILE:HD13	2.34	0.52
5:Ca:219:ARG:H	6:Ch:148:PRO:HD2	1.73	0.52
6:Cn:115:MET:HA	6:Cn:118:ASN:OD1	2.09	0.52
6:Dl:115:MET:HA	6:Dl:118:ASN:OD1	2.10	0.52
7:Em:795:GLN:HA	7:Em:798:THR:HB	1.91	0.52
8:Ey:182:LEU:HD21	8:Ez:240:ILE:HD13	1.90	0.52
8:Ey:448:LEU:HD13	8:Ez:340:THR:HG22	1.91	0.52
8:Fa:449:LYS:HD2	8:Fa:449:LYS:C	2.35	0.52
9:Fc:238:ASN:O	9:Fc:242:LYS:HG2	2.09	0.52
9:Fe:419:PHE:O	9:Fe:423:ILE:HG12	2.08	0.52
9:Ff:66:SER:O	9:Ff:69:LEU:HD12	2.09	0.52
6:Af:205:GLN:HB2	6:Af:214:MET:SD	2.50	0.52
6:Al:117:ARG:HH12	7:Ee:797:ARG:CZ	2.23	0.52
6:Bj:199:PRO:HG3	7:Ek:819:VAL:HG12	1.91	0.52
6:Bp:106:VAL:HG22	6:Bp:110:LYS:HE3	1.91	0.52
5:Bu:213:GLN:NE2	5:Bu:223:ILE:HG23	2.24	0.52

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
6:Ch:158:VAL:HB	6:Ch:256:VAL:HA	1.91	0.52
1:Ci:11:TYR:CD2	3:Ck:9:TYR:HB3	2.44	0.52
6:Cn:115:MET:CE	7:Em:798:THR:HA	2.38	0.52
6:Df:227:ALA:HB2	6:Df:238:MET:HE1	1.92	0.52
6:Df:229:ARG:NH1	6:Df:229:ARG:HA	2.25	0.52
7:Er:794:ILE:HD12	7:Er:794:ILE:H	1.73	0.52
8:Ew:384:PHE:HB3	8:Fa:82:TYR:CD2	2.44	0.52
8:Ex:55:THR:HA	8:Ey:42:TYR:CE1	2.40	0.52
8:Ey:78:GLN:HB2	8:Ez:383:LEU:CD1	2.39	0.52
8:Fa:207:GLN:HA	8:Fa:210:VAL:HG12	1.91	0.52
9:Fb:64:PHE:CZ	9:Fb:780:PRO:HG3	2.44	0.52
9:Fc:38:LEU:O	9:Fc:42:PHE:HD1	1.92	0.52
9:Fc:129:MET:O	9:Fc:133:VAL:HG12	2.09	0.52
9:Fc:179:LYS:HA	9:Fc:550:LEU:HD12	1.91	0.52
9:Fc:813:SER:O	9:Fc:817:ILE:HG13	2.09	0.52
9:Fe:72:GLY:O	9:Fe:76:ILE:HG12	2.10	0.52
9:Fe:176:GLY:HA3	9:Fe:492:ASP:HB2	1.91	0.52
9:Ff:289:ALA:HA	9:Ff:312:SER:HB2	1.90	0.52
6:Ax:123:ASN:HB2	6:Ax:126:GLN:HG3	1.92	0.52
6:Bd:238:MET:HB3	6:Bj:250:TYR:CD1	2.45	0.52
6:Bj:129:LYS:O	6:Bj:133:ILE:HG12	2.09	0.52
6:Bp:151:PRO:HA	6:Bp:248:VAL:HG13	1.91	0.52
6:Ct:107:ILE:H	6:Ct:107:ILE:HD12	1.73	0.52
6:Dx:221:TYR:CD1	6:Ed:139:TYR:HD1	2.27	0.52
7:Ej:790:TYR:O	7:Ej:794:ILE:HG13	2.09	0.52
7:Ek:794:ILE:HA	7:Ek:797:ARG:NH1	2.25	0.52
7:Es:798:THR:O	7:Es:802:LEU:HG	2.09	0.52
8:Ex:302:TRP:HD1	8:Ex:303:ASN:OD1	1.92	0.52
8:Ez:263:GLN:H	8:Ez:268:THR:HG21	1.75	0.52
8:Ez:426:MET:CE	8:Fa:425:GLN:HG2	2.35	0.52
8:Fa:295:LEU:O	8:Fa:299:SER:CB	2.57	0.52
9:Fc:302:VAL:HB	9:Fc:320:ILE:HG23	1.91	0.52
9:Fc:370:PHE:HE2	9:Fc:565:SER:HB3	1.74	0.52
9:Fd:339:MET:HA	9:Fd:342:THR:HG22	1.90	0.52
9:Fe:327:THR:HA	9:Fe:330:LEU:HD23	1.90	0.52
9:Fe:627:MET:HE2	9:Fe:627:MET:C	2.35	0.52
9:Fe:923:LEU:O	9:Fe:926:VAL:HB	2.10	0.52
6:Af:131:LYS:HG2	6:Ed:145:PRO:HG3	1.91	0.52
2:Ah:4:MET:HE1	5:Ak:248:ASP:HA	1.90	0.52
5:Aq:233:ARG:NH1	5:Aq:236:SER:HB2	2.24	0.52
5:Aw:263:LYS:HD2	5:Aw:264:PHE:O	2.09	0.52

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
6:Ax:185:THR:HG21	6:Ax:263:ALA:HA	1.92	0.52
2:Az:6:GLY:HA2	6:Bp:125:GLU:HB3	1.91	0.52
5:Bi:219:ARG:HA	5:Bi:232:VAL:O	2.09	0.52
6:Bj:225:ASN:ND2	6:Bp:176:VAL:HG22	2.25	0.52
6:Bp:106:VAL:HA	6:Bp:109:LYS:HG2	1.91	0.52
6:Cz:182:LEU:HD11	6:Cz:186:GLY:HA2	1.91	0.52
5:Dk:219:ARG:HD3	6:Dr:148:PRO:O	2.09	0.52
5:Dw:222:LEU:HD21	5:Dw:230:LEU:HD23	1.90	0.52
6:Dx:187:ALA:HB1	6:Dx:262:ASN:HB2	1.91	0.52
7:Ef:772:GLU:HA	7:Ef:775:GLN:CD	2.35	0.52
8:Ew:188:SER:HA	8:Ew:191:MET:HB2	1.89	0.52
8:Ew:230:PHE:HE2	8:Fa:298:TYR:CZ	2.28	0.52
8:Ew:428:LEU:HA	8:Ew:431:GLN:HG2	1.90	0.52
9:Fb:109:PHE:HB2	9:Fb:786:VAL:HG11	1.90	0.52
9:Fb:149:TYR:CB	9:Ff:59:ASN:HD22	2.23	0.52
9:Fb:669:ASP:HA	9:Fb:672:ILE:HG12	1.90	0.52
9:Fb:798:GLY:O	9:Fb:802:ILE:HG12	2.08	0.52
9:Fc:804:ILE:O	9:Fc:808:VAL:HG22	2.09	0.52
9:Fd:232:ASP:OD2	9:Fd:234:ILE:HG13	2.10	0.52
9:Fd:680:GLN:OE1	9:Fd:683:ILE:HG12	2.09	0.52
7:Fi:26:ALA:O	7:Fi:30:ILE:HG12	2.08	0.52
6:Al:171:LEU:HG	6:Al:217:ALA:HB2	1.91	0.52
6:Bj:173:GLN:CB	6:Bj:220:LEU:HD23	2.39	0.52
5:Bo:254:ILE:HB	5:Bo:262:ILE:HB	1.92	0.52
6:Df:238:MET:HB3	6:Dl:250:TYR:CE1	2.45	0.52
6:Dl:205:GLN:HB2	6:Dl:214:MET:CE	2.39	0.52
6:Dr:170:ARG:HD3	6:Dr:247:ALA:C	2.35	0.52
1:Ds:9:LEU:O	1:Ds:13:LYS:HG2	2.10	0.52
6:Dx:201:SER:HA	6:Dx:219:LYS:HE3	1.92	0.52
8:Ex:170:LEU:H	8:Ex:170:LEU:HD12	1.74	0.52
9:Fb:182:LEU:HD23	9:Fb:571:VAL:HG21	1.92	0.52
9:Fb:323:SER:O	9:Fb:326:GLN:HG3	2.09	0.52
9:Fb:547:GLN:HA	9:Fb:568:TYR:CE1	2.44	0.52
9:Fe:664:LEU:HD22	9:Fe:665:GLN:OE1	2.10	0.52
9:Ff:146:ALA:O	9:Ff:150:LEU:HD22	2.09	0.52
9:Ff:200:ALA:HA	9:Ff:203:ASP:OD1	2.09	0.52
5:Aq:219:ARG:HH12	6:Ax:150:LYS:HG2	1.75	0.52
6:Ax:173:GLN:HG2	6:Ax:220:LEU:HD22	1.91	0.52
6:Bd:107:ILE:HG13	6:Bd:110:LYS:HZ3	1.74	0.52
6:Bd:204:ILE:HD12	6:Bd:214:MET:O	2.09	0.52
6:Cn:230:LEU:HD12	6:Cn:231:ARG:N	2.15	0.52

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
6:Ed:218:THR:O	7:Ev:815:VAL:HG21	2.10	0.52
8:Ex:429:ASP:O	8:Ex:433:GLN:HG2	2.10	0.52
8:Ey:297:ALA:HA	8:Ey:300:GLU:OE1	2.10	0.52
8:Ey:318:GLN:CD	8:Ey:318:GLN:H	2.15	0.52
8:Ey:440:ASN:O	8:Ey:443:MET:HG3	2.10	0.52
8:Ez:298:TYR:HE1	8:Fa:221:ALA:HB2	1.75	0.52
8:Ez:385:ASP:H	8:Ez:398:GLN:HE22	1.57	0.52
9:Fc:129:MET:N	9:Fc:129:MET:HE2	2.24	0.52
9:Fd:438:ARG:NH1	9:Fd:439:GLN:HG2	2.24	0.52
9:Fd:630:ASP:HA	9:Fd:634:ASN:HB2	1.92	0.52
9:Fd:647:ILE:HG23	9:Fd:651:MET:HE3	1.91	0.52
9:Fe:290:LEU:HD11	9:Fe:437:ILE:HD12	1.91	0.52
9:Fe:635:TYR:O	9:Fe:639:TYR:HB3	2.10	0.52
9:Ff:33:LEU:HB3	9:Ff:462:ILE:HD11	1.91	0.52
6:Al:242:ILE:HD12	6:Al:243:PRO:HD2	1.92	0.52
6:Ar:115:MET:HE1	7:Ee:801:MET:CB	2.36	0.52
6:Ch:242:ILE:HD12	6:Ch:243:PRO:HD2	1.90	0.52
6:Ct:144:THR:OG1	6:Ct:148:PRO:HG3	2.09	0.52
6:Cz:173:GLN:HG2	6:Cz:220:LEU:HA	1.90	0.52
5:De:212:ILE:H	5:De:212:ILE:HD12	1.74	0.52
6:Dx:223:TYR:HB2	6:Dx:242:ILE:HA	1.92	0.52
8:Ew:359:MET:HE1	8:Ew:369:THR:HG22	1.92	0.52
8:Ew:415:ILE:HG23	8:Fa:419:LEU:HD21	1.91	0.52
8:Ex:249:ILE:HD12	8:Ex:250:ALA:N	2.25	0.52
8:Ex:418:LEU:O	8:Ex:422:ILE:HG12	2.10	0.52
8:Ey:249:ILE:HA	8:Ey:354:ARG:HH22	1.75	0.52
8:Fa:212:GLN:HA	8:Fa:215:ILE:HG22	1.91	0.52
9:Fb:650:GLU:OE1	9:Fb:650:GLU:N	2.41	0.52
9:Fc:800:PHE:HA	9:Fc:803:MET:HE2	1.90	0.52
9:Fc:828:VAL:HA	9:Fc:831:TRP:CE3	2.45	0.52
9:Ff:743:VAL:HG22	9:Ff:747:PHE:CE1	2.45	0.52
9:Ff:925:ILE:O	9:Ff:929:ALA:HB2	2.09	0.52
5:Bc:212:ILE:HG13	5:Bc:262:ILE:HG22	1.91	0.52
5:Bo:214:ALA:HB3	5:Bo:221:TRP:HE3	1.75	0.52
6:Cb:204:ILE:HD12	6:Cb:214:MET:O	2.10	0.52
6:Cb:236:PRO:HG2	6:Cb:238:MET:HE1	1.92	0.52
6:Ch:181:PHE:CZ	6:Ch:230:LEU:HG	2.45	0.52
6:Ch:238:MET:HG3	6:Cn:250:TYR:CZ	2.45	0.52
5:Cs:247:ILE:HG12	5:Cs:254:ILE:HG23	1.91	0.52
6:Cz:134:TYR:CE1	7:Ep:812:TRP:HB3	2.44	0.52
6:Dl:149:PRO:HA	6:Dl:246:LYS:O	2.09	0.52

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
8:Ew:105:LYS:HZ2	8:Ex:368:ILE:H	1.57	0.52
8:Ew:457:PHE:CZ	8:Ex:329:TYR:HB2	2.43	0.52
8:Fa:351:LEU:CD1	8:Fa:355:LEU:HD21	2.39	0.52
9:Fb:345:THR:O	9:Fb:348:GLN:HG3	2.09	0.52
9:Fb:655:ILE:O	9:Fb:659:PHE:HD1	1.93	0.52
9:Fc:34:SER:HA	9:Fc:462:ILE:HD11	1.92	0.52
9:Fc:462:ILE:HG13	9:Fc:463:MET:SD	2.49	0.52
9:Fc:633:TYR:HA	9:Fc:637:PHE:HD1	1.75	0.52
9:Fc:695:MET:HE3	9:Fc:698:ASN:HD21	1.74	0.52
9:Fc:725:PHE:HE2	9:Fd:725:PHE:CE2	2.28	0.52
9:Fd:527:LEU:HA	9:Fd:530:ILE:HD13	1.91	0.52
9:Fe:936:LEU:HA	9:Fe:939:LYS:HE3	1.91	0.52
5:Ae:219:ARG:NH1	5:Ae:233:ARG:HD3	2.25	0.52
6:Al:158:VAL:HB	6:Al:256:VAL:HA	1.92	0.52
6:Ar:225:ASN:HB2	6:Ax:250:TYR:CZ	2.44	0.52
6:Ax:171:LEU:HD11	6:Ax:241:LEU:HB3	1.90	0.52
6:Bj:173:GLN:HB2	6:Bj:220:LEU:HA	1.92	0.52
5:Bo:253:ARG:O	5:Bo:254:ILE:HD13	2.10	0.52
5:Bu:213:GLN:OE1	5:Bu:223:ILE:HG12	2.10	0.52
5:Cm:233:ARG:HH12	5:Cm:236:SER:HB3	1.73	0.52
1:Cu:9:LEU:O	1:Cu:13:LYS:HG2	2.09	0.52
5:Cy:216:ILE:HD11	6:Df:148:PRO:HG2	1.92	0.52
5:De:219:ARG:HG2	6:Dl:148:PRO:HD2	1.91	0.52
6:Dl:191:ILE:HD12	6:Dl:206:TRP:NE1	2.24	0.52
6:Ed:130:LEU:HA	6:Ed:133:ILE:HD12	1.91	0.52
6:Ed:168:VAL:HA	6:Ed:240:THR:O	2.10	0.52
7:Et:797:ARG:HH12	7:Et:801:MET:HB3	1.75	0.52
7:Et:809:VAL:O	7:Et:813:LYS:HG3	2.10	0.52
8:Ew:144:GLN:HB2	8:Ew:179:PHE:CZ	2.45	0.52
8:Ew:252:LEU:HD12	8:Ew:252:LEU:O	2.10	0.52
8:Ew:340:THR:HG22	8:Fa:448:LEU:HD22	1.91	0.52
8:Ew:382:ARG:HB3	8:Ew:399:TRP:CE3	2.44	0.52
8:Ex:195:GLN:NE2	8:Ex:294:LYS:HD2	2.24	0.52
8:Ez:331:ASN:ND2	8:Fa:211:MET:HB2	2.24	0.52
9:Fb:180:GLY:O	9:Fb:184:ILE:HD13	2.09	0.52
9:Fb:760:ILE:HD13	9:Fb:930:PHE:HB3	1.90	0.52
9:Fc:304:VAL:HG21	9:Fc:439:GLN:HE22	1.75	0.52
9:Fc:913:PHE:CZ	9:Fc:917:ILE:HD11	2.45	0.52
9:Fd:127:PHE:O	9:Fd:131:VAL:HG23	2.10	0.52
9:Fd:131:VAL:HG21	9:Fd:805:LEU:HD11	1.92	0.52
6:Af:205:GLN:NE2	6:Af:216:GLN:HE22	2.06	0.52

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
6:Bd:170:ARG:NH1	6:Bd:247:ALA:HB3	2.25	0.52
6:Bp:189:TRP:CD1	6:Bp:230:LEU:HD12	2.44	0.52
6:Bp:196:LEU:HD12	6:Bp:204:ILE:HD12	1.91	0.52
6:Ch:196:LEU:HD23	6:Ch:199:PRO:HB3	1.92	0.52
6:Cn:106:VAL:HA	6:Cn:109:LYS:NZ	2.25	0.52
6:Cz:130:LEU:HA	6:Cz:133:ILE:HD12	1.92	0.52
7:Er:802:LEU:O	7:Er:806:THR:HG22	2.10	0.52
8:Ew:346:ASN:O	8:Ew:350:ILE:HG12	2.10	0.52
8:Ew:453:PRO:HD3	8:Ex:336:TYR:CD1	2.45	0.52
8:Ex:206:TYR:CE1	8:Ex:208:ASN:HB3	2.44	0.52
8:Ey:437:LEU:HD13	8:Ez:436:ILE:HD13	1.92	0.52
8:Ez:332:ASN:HA	8:Ez:335:VAL:HG22	1.92	0.52
8:Fa:69:PHE:CZ	8:Fa:402:GLN:HB3	2.45	0.52
9:Fb:909:TYR:HA	9:Fb:912:PHE:HD2	1.75	0.52
9:Fc:748:VAL:O	9:Fc:753:ILE:HG12	2.10	0.52
9:Fc:775:ALA:HB2	9:Fc:808:VAL:HG11	1.92	0.52
9:Fd:377:GLN:HG2	9:Fd:379:GLY:H	1.75	0.52
9:Fd:737:ALA:O	9:Fd:741:THR:HG23	2.10	0.52
9:Fe:251:LEU:HD23	9:Fe:325:LEU:HD11	1.92	0.52
9:Fe:704:TRP:CD1	9:Ff:734:LEU:HD22	2.45	0.52
9:Ff:190:CYS:O	9:Ff:194:LEU:HD13	2.10	0.52
9:Ff:530:ILE:O	9:Ff:534:ILE:HG12	2.10	0.52
9:Ff:824:ALA:O	9:Ff:828:VAL:HG23	2.10	0.52
6:Af:109:LYS:HD3	6:Af:109:LYS:N	2.25	0.51
5:Ak:243:MET:HE3	5:Ak:257:SER:HB3	1.92	0.51
5:Aq:221:TRP:CZ3	5:Aq:231:THR:HB	2.45	0.51
6:Ax:137:SER:O	6:Ax:141:LYS:HG3	2.09	0.51
6:Bv:115:MET:HE3	7:Ej:802:LEU:HD13	1.91	0.51
6:Bv:196:LEU:HD21	6:Bv:202:PHE:HB2	1.92	0.51
5:Cy:210:TYR:HB3	5:Cy:222:LEU:HD12	1.90	0.51
6:Dr:172:SER:HB3	6:Dr:248:VAL:CG1	2.39	0.51
6:Ed:132:GLN:HA	6:Ed:135:GLU:HG2	1.93	0.51
8:Ew:85:PHE:HD2	8:Ew:86:LEU:HD12	1.74	0.51
8:Ey:451:ALA:HA	9:Fb:619:PHE:HD2	1.75	0.51
8:Fa:205:MET:HE2	8:Fa:205:MET:HA	1.92	0.51
9:Fb:594:ILE:H	9:Fb:594:ILE:HD12	1.75	0.51
9:Fb:704:TRP:HA	9:Fb:707:LEU:HD12	1.92	0.51
9:Fb:921:MET:O	9:Fb:925:ILE:HD12	2.10	0.51
9:Fc:425:ASP:O	9:Fc:429:ILE:HG13	2.09	0.51
9:Fc:662:ILE:HB	9:Fc:703:LEU:HD21	1.91	0.51
9:Fc:766:PHE:O	9:Fc:770:ILE:HG13	2.10	0.51

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
9:Fc:814:LEU:HD23	9:Fc:940:VAL:HG21	1.92	0.51
9:Fd:420:LEU:H	9:Fd:420:LEU:HD12	1.74	0.51
9:Fe:193:GLY:HA2	9:Fe:357:PHE:CE1	2.45	0.51
9:Fe:430:MET:HE2	9:Fe:430:MET:N	2.25	0.51
9:Ff:259:PHE:CE2	9:Ff:286:ASN:HB2	2.43	0.51
9:Ff:656:VAL:O	9:Ff:660:LEU:HG	2.10	0.51
6:Ar:196:LEU:HD23	6:Ar:199:PRO:HB3	1.92	0.51
6:Bd:159:ASN:HD21	6:Bd:164:SER:HB2	1.75	0.51
6:Bd:169:ILE:HG23	6:Bd:241:LEU:HA	1.91	0.51
6:Bd:191:ILE:HD12	6:Bd:206:TRP:CE2	2.45	0.51
6:Ch:196:LEU:HD11	6:Ch:202:PHE:CD2	2.45	0.51
6:Cn:192:ALA:HB3	6:Cn:229:ARG:HG3	1.92	0.51
5:Cy:230:LEU:HD12	5:Cy:231:THR:N	2.22	0.51
5:Dk:219:ARG:NH1	5:Dk:233:ARG:HB3	2.24	0.51
6:Dr:189:TRP:CE2	6:Dr:260:GLY:HA2	2.45	0.51
5:Dw:213:GLN:HG2	5:Dw:221:TRP:O	2.11	0.51
6:Dx:189:TRP:CZ2	6:Dx:233:LEU:HD12	2.45	0.51
6:Ed:172:SER:HB3	6:Ed:175:PHE:HB2	1.92	0.51
8:Ew:424:TYR:O	8:Ew:427:TYR:HB3	2.10	0.51
8:Ex:249:ILE:HA	8:Ex:354:ARG:HH12	1.74	0.51
8:Ex:426:MET:C	8:Ex:426:MET:HE2	2.36	0.51
8:Ez:253:LEU:HD22	8:Ez:373:LEU:HD13	1.93	0.51
8:Ez:341:SER:O	8:Ez:345:SER:HB3	2.10	0.51
9:Fc:548:PRO:HD3	9:Fc:568:TYR:HD2	1.75	0.51
9:Fc:693:GLY:O	9:Fc:697:ILE:HG23	2.10	0.51
9:Fc:725:PHE:CE1	9:Fd:729:MET:HG2	2.45	0.51
9:Fd:500:GLN:O	9:Fd:503:LYS:HG2	2.11	0.51
9:Fe:531:GLN:HA	9:Fe:534:ILE:HD12	1.91	0.51
9:Ff:124:MET:HA	9:Ff:127:PHE:CD2	2.45	0.51
7:Fg:35:GLY:O	7:Fg:39:ILE:HG12	2.09	0.51
6:Ax:129:LYS:HD2	6:Bd:112:PHE:CE1	2.46	0.51
6:Ax:229:ARG:NH1	6:Ax:229:ARG:HA	2.26	0.51
6:Bv:134:TYR:O	6:Bv:138:GLU:HG3	2.09	0.51
5:Ca:232:VAL:HG21	5:Ca:244:VAL:HG21	1.92	0.51
6:Dr:204:ILE:HA	6:Dr:214:MET:O	2.10	0.51
3:Du:13:LYS:HD2	3:Du:13:LYS:O	2.11	0.51
7:Eg:807:GLN:HA	7:Eg:810:GLN:CG	2.36	0.51
8:Ey:228:PRO:HB3	8:Ey:234:LYS:HE2	1.91	0.51
8:Ey:329:TYR:CZ	8:Ey:333:LEU:HD21	2.44	0.51
8:Ez:179:PHE:HZ	8:Fa:233:TYR:HA	1.75	0.51
8:Fa:343:GLY:O	8:Fa:347:LEU:HG	2.10	0.51

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
9:Fb:74:ILE:HA	9:Fb:77:MET:HE3	1.92	0.51
9:Fb:378:PHE:HB3	9:Fb:414:PHE:CD1	2.46	0.51
9:Fb:701:GLY:O	9:Fb:705:LEU:HD22	2.11	0.51
9:Fc:738:TRP:HD1	9:Fc:912:PHE:CE2	2.28	0.51
9:Fd:419:PHE:O	9:Fd:423:ILE:HG12	2.10	0.51
9:Fe:213:PRO:O	9:Fe:218:PRO:HB3	2.10	0.51
9:Fe:667:MET:O	9:Fe:670:ILE:HG13	2.10	0.51
9:Fe:708:LEU:HD13	9:Ff:730:MET:SD	2.50	0.51
9:Ff:685:PRO:HG3	9:Ff:831:TRP:NE1	2.22	0.51
9:Ff:921:MET:HE3	9:Ff:924:ILE:HD11	1.92	0.51
5:Aw:212:ILE:HB	5:Aw:264:PHE:CE1	2.44	0.51
6:Ax:115:MET:HB3	6:Ax:119:LEU:HD23	1.91	0.51
5:Bc:213:GLN:HE21	5:Bc:221:TRP:HB3	1.76	0.51
6:Bd:121:PRO:HB2	7:Eh:801:MET:SD	2.50	0.51
1:Bk:16:ALA:HB3	2:Br:9:ARG:HE	1.75	0.51
6:Cb:189:TRP:CD1	6:Cb:230:LEU:HD12	2.45	0.51
6:Ch:132:GLN:HA	6:Ch:135:GLU:OE2	2.11	0.51
6:Cz:149:PRO:HB2	6:Cz:248:VAL:HG12	1.91	0.51
6:Dr:229:ARG:HD3	6:Dr:230:LEU:O	2.11	0.51
6:Dx:150:LYS:HG3	6:Dx:152:THR:HG23	1.91	0.51
7:Es:797:ARG:HA	7:Es:800:ASP:OD2	2.10	0.51
8:Ew:452:GLN:HE22	8:Ex:290:PRO:HG3	1.75	0.51
8:Ex:126:SER:HB2	8:Ex:171:ASN:HA	1.92	0.51
9:Fb:78:TYR:O	9:Fb:82:VAL:HG23	2.11	0.51
9:Fb:243:GLN:O	9:Fb:247:THR:HG22	2.11	0.51
9:Fc:735:LEU:HD23	9:Fc:738:TRP:HZ3	1.74	0.51
9:Fc:819:TYR:CE1	9:Fc:823:ILE:HD11	2.45	0.51
9:Ff:73:GLY:HA2	9:Ff:76:ILE:HG12	1.93	0.51
9:Ff:122:CYS:HB2	9:Ff:125:GLN:HG2	1.91	0.51
9:Ff:345:THR:O	9:Ff:348:GLN:HG3	2.11	0.51
9:Ff:419:PHE:O	9:Ff:423:ILE:HG12	2.11	0.51
9:Ff:689:LEU:HA	9:Ff:692:MET:HG3	1.92	0.51
9:Ff:750:ALA:O	9:Ff:754:PRO:HG2	2.10	0.51
6:Af:238:MET:HB3	6:Al:250:TYR:HD2	1.75	0.51
6:Ar:221:TYR:CZ	6:Ax:139:TYR:HD2	2.28	0.51
6:Bj:134:TYR:O	6:Bj:138:GLU:HG3	2.10	0.51
6:Bp:115:MET:HA	6:Bp:118:ASN:OD1	2.11	0.51
5:Cs:222:LEU:HD12	5:Cs:223:ILE:N	2.26	0.51
5:Cy:221:TRP:HZ3	5:Cy:229:THR:HG22	1.76	0.51
6:Df:191:ILE:HB	6:Df:206:TRP:CZ2	2.45	0.51
2:Dh:5:ILE:HD11	6:Dr:136:THR:O	2.10	0.51

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
6:Dl:225:ASN:HD21	6:Dr:176:VAL:H	1.58	0.51
6:Dr:110:LYS:O	6:Dr:110:LYS:HD3	2.10	0.51
6:Dr:126:GLN:HA	6:Dr:129:LYS:HZ2	1.75	0.51
7:Ej:778:LEU:HA	7:Ej:781:GLN:HG3	1.91	0.51
7:Ep:806:THR:HA	7:Ep:809:VAL:HG12	1.92	0.51
8:Ex:254:MET:N	8:Ex:254:MET:HE2	2.25	0.51
8:Ez:330:PHE:HB3	8:Ez:334:ARG:HH21	1.75	0.51
8:Fa:207:GLN:O	8:Fa:210:VAL:HG12	2.11	0.51
9:Fc:79:THR:HA	9:Fc:82:VAL:HG22	1.91	0.51
9:Fc:180:GLY:O	9:Fc:184:ILE:HG22	2.10	0.51
9:Fc:837:PHE:CZ	9:Fc:914:SER:HB2	2.46	0.51
9:Fd:641:TYR:HA	9:Fd:644:PHE:CE2	2.46	0.51
9:Ff:668:LYS:HG3	9:Ff:669:ASP:N	2.25	0.51
6:Af:176:VAL:N	6:Ed:225:ASN:HD21	2.09	0.51
6:Al:129:LYS:O	6:Al:129:LYS:HD3	2.10	0.51
6:Al:238:MET:HG2	6:Ar:251:ARG:NH1	2.26	0.51
2:An:2:VAL:HA	3:Ao:1:ARG:NE	2.22	0.51
6:Ar:141:LYS:HD2	7:Eg:812:TRP:CZ2	2.45	0.51
6:Ar:194:TYR:HD2	6:Ar:226:LEU:HD11	1.76	0.51
6:Ax:107:ILE:H	6:Ax:107:ILE:HD12	1.76	0.51
6:Bj:138:GLU:CA	6:Bj:141:LYS:HB3	2.37	0.51
6:Ch:189:TRP:CE2	6:Ch:260:GLY:HA2	2.45	0.51
5:Cm:253:ARG:O	5:Cm:254:ILE:HD13	2.11	0.51
6:Cn:106:VAL:O	6:Cn:110:LYS:HG2	2.11	0.51
6:Cn:171:LEU:HD21	6:Cn:241:LEU:HB3	1.92	0.51
3:Cw:5:GLY:HA3	4:Dd:1:PRO:HA	1.93	0.51
6:Dl:151:PRO:HA	6:Dl:248:VAL:HG13	1.92	0.51
6:Dr:172:SER:HB3	6:Dr:248:VAL:HG11	1.91	0.51
8:Ex:43:LEU:HD11	8:Ey:42:TYR:CD2	2.45	0.51
8:Ez:414:GLU:O	8:Ez:418:LEU:HG	2.11	0.51
9:Fb:685:PRO:HB2	9:Fb:830:VAL:HG11	1.91	0.51
9:Fc:41:LEU:HD22	9:Fc:42:PHE:CD1	2.45	0.51
9:Fc:112:ALA:HA	9:Fc:115:ILE:CD1	2.40	0.51
9:Fc:600:THR:HB	9:Fc:645:LEU:HD13	1.92	0.51
9:Fd:434:LEU:HA	9:Fd:437:ILE:HG22	1.91	0.51
9:Fd:703:LEU:O	9:Fd:707:LEU:HG	2.11	0.51
9:Fe:129:MET:O	9:Fe:132:ILE:HG13	2.10	0.51
9:Ff:525:ASP:HA	9:Ff:528:ILE:HD12	1.93	0.51
7:Fh:24:THR:HA	7:Fh:27:LEU:HD12	1.92	0.51
3:Ai:2:VAL:HG21	6:Ax:129:LYS:HD3	1.93	0.51
6:Ar:128:VAL:O	6:Ar:132:GLN:HG2	2.11	0.51

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
6:Bd:190:PRO:O	6:Bd:230:LEU:HD22	2.11	0.51
6:Bp:122:LEU:HB3	6:Bp:127:VAL:HG12	1.92	0.51
5:Bu:251:GLN:HB3	5:Bu:253:ARG:CZ	2.41	0.51
6:Bv:125:GLU:O	6:Bv:129:LYS:HG3	2.11	0.51
6:Ch:125:GLU:C	6:Ch:129:LYS:HE3	2.34	0.51
6:Ct:157:PHE:HB2	6:Ct:255:ARG:HH21	1.76	0.51
6:Cz:150:LYS:HE3	6:Cz:150:LYS:HA	1.92	0.51
6:Df:115:MET:CE	7:Ep:798:THR:HA	2.27	0.51
6:Dr:190:PRO:O	6:Dr:230:LEU:HD22	2.10	0.51
6:Dx:126:GLN:HA	6:Dx:129:LYS:CG	2.39	0.51
5:Ec:233:ARG:HG3	5:Ec:234:GLU:N	2.25	0.51
7:Eu:766:ASP:N	7:Eu:769:SER:HG	2.09	0.51
8:Ey:123:GLN:OE1	8:Ey:135:LYS:HB3	2.11	0.51
8:Ez:432:ILE:HB	8:Ez:433:GLN:NE2	2.26	0.51
9:Fb:263:MET:HE2	9:Fb:279:CYS:O	2.10	0.51
9:Fc:837:PHE:HZ	9:Fc:914:SER:HB2	1.76	0.51
9:Fd:45:VAL:HG12	9:Fd:49:LEU:HD23	1.91	0.51
9:Fd:113:LEU:HA	9:Fd:123:MET:SD	2.50	0.51
9:Ff:111:LEU:HD23	7:Fi:27:LEU:HD12	1.92	0.51
9:Ff:339:MET:SD	9:Ff:426:TYR:HB2	2.50	0.51
6:Ar:121:PRO:HB2	7:Ef:801:MET:CE	2.41	0.51
5:Bi:219:ARG:NH1	5:Bi:233:ARG:HB3	2.26	0.51
6:Bj:238:MET:HE2	6:Bj:238:MET:HA	1.93	0.51
6:Bp:181:PHE:CZ	6:Bp:228:VAL:HG21	2.45	0.51
5:Cg:207:ARG:NH1	5:Cg:240:GLY:HA3	2.25	0.51
5:Cm:219:ARG:HH12	5:Cm:233:ARG:HD3	1.76	0.51
6:Cn:130:LEU:HG	7:En:812:TRP:CZ3	2.45	0.51
6:Cn:205:GLN:NE2	7:Eo:820:TYR:HB2	2.25	0.51
6:Cz:199:PRO:HG3	7:Er:819:VAL:HG13	1.93	0.51
6:Dl:249:ASP:OD2	6:Dl:252:VAL:HG22	2.11	0.51
7:Eu:807:GLN:O	7:Eu:810:GLN:HG2	2.10	0.51
9:Fb:693:GLY:O	9:Fb:697:ILE:HG23	2.11	0.51
9:Fb:766:PHE:HB3	9:Fc:928:LYS:HZ3	1.75	0.51
9:Fc:645:LEU:HD22	9:Fc:648:PHE:HD2	1.76	0.51
9:Fe:378:PHE:HD1	9:Fe:414:PHE:HE2	1.59	0.51
6:Af:133:ILE:HD11	6:Al:120:TYR:HB2	1.93	0.51
6:Af:214:MET:HE1	6:Af:216:GLN:HG3	1.92	0.51
1:Am:2:THR:HG23	5:Aq:253:ARG:NH1	2.26	0.51
6:Ax:228:VAL:HB	6:Ax:237:VAL:HB	1.93	0.51
6:Bj:129:LYS:HE3	6:Bp:112:PHE:CD2	2.46	0.51
2:Br:4:MET:HB2	4:Bt:2:PHE:CZ	2.46	0.51

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
6:Bv:193:ALA:HB1	7:Em:820:TYR:HE2	1.76	0.51
6:Cn:157:PHE:HA	6:Cn:255:ARG:HB2	1.93	0.51
6:Cz:214:MET:HE1	7:Eq:818:GLN:CD	2.36	0.51
6:Dx:225:ASN:HD21	6:Ed:176:VAL:N	2.08	0.51
7:Ej:797:ARG:HG3	7:Ej:798:THR:N	2.26	0.51
7:Em:804:ALA:O	7:Em:808:LEU:HG	2.11	0.51
7:Er:801:MET:HE2	7:Er:801:MET:O	2.10	0.51
7:Et:781:GLN:HE22	7:Et:785:LEU:HD23	1.76	0.51
8:Ew:129:SER:HB3	8:Ew:137:THR:HG21	1.93	0.51
8:Ew:424:TYR:O	8:Ew:428:LEU:HD22	2.10	0.51
8:Ex:207:GLN:O	8:Ex:211:MET:HG3	2.11	0.51
8:Fa:433:GLN:HA	8:Fa:436:ILE:HG12	1.92	0.51
9:Fb:556:ARG:HB3	9:Fb:576:MET:CE	2.41	0.51
9:Fc:612:CYS:HB3	9:Fc:626:ARG:N	2.25	0.51
9:Fd:170:ALA:HB2	9:Fd:177:VAL:HG22	1.93	0.51
9:Fd:347:ALA:HA	9:Fd:350:MET:HG2	1.92	0.51
9:Fe:77:MET:O	9:Fe:81:MET:HG3	2.11	0.51
9:Fe:435:ASN:O	9:Fe:439:GLN:HG2	2.11	0.51
9:Fe:908:VAL:O	9:Fe:911:PHE:HB3	2.11	0.51
7:Fh:5:LYS:HA	7:Fh:8:LEU:HG	1.93	0.51
5:Ae:219:ARG:HH22	6:Al:150:LYS:HD2	1.76	0.51
2:Ah:4:MET:SD	5:Ak:248:ASP:HA	2.51	0.51
5:Ak:247:ILE:HG23	5:Ak:254:ILE:HD12	1.92	0.51
6:Ax:138:GLU:HA	6:Ax:141:LYS:HD2	1.92	0.51
6:Ax:203:ASN:ND2	6:Ax:205:GLN:HE22	2.09	0.51
6:Cn:218:THR:OG1	6:Cn:219:LYS:HD2	2.11	0.51
6:Ct:199:PRO:HG2	7:Eq:816:GLU:OE1	2.10	0.51
6:Df:173:GLN:CD	7:Er:815:VAL:HG21	2.35	0.51
6:Di:190:PRO:O	6:Di:231:ARG:HG2	2.11	0.51
6:Ed:238:MET:HE3	6:Ed:238:MET:HA	1.92	0.51
7:Eh:789:LYS:HD2	7:Eh:789:LYS:C	2.36	0.51
7:Eu:797:ARG:HG2	7:Eu:801:MET:SD	2.51	0.51
8:Ew:89:ILE:HD12	8:Ew:90:PRO:HD2	1.92	0.51
8:Ey:341:SER:HA	8:Ey:344:VAL:HG22	1.92	0.51
9:Fb:163:PRO:HB2	9:Fb:579:LEU:HD21	1.92	0.51
9:Fb:377:GLN:HE21	9:Fb:379:GLY:HA2	1.76	0.51
9:Fb:921:MET:O	9:Fb:924:ILE:HG13	2.11	0.51
9:Fc:706:THR:O	9:Fc:710:MET:HG2	2.11	0.51
9:Fe:287:ILE:HG13	9:Fe:332:ARG:NH1	2.26	0.51
9:Ff:147:LEU:HB3	9:Ff:453:ILE:HG23	1.91	0.51
9:Ff:664:LEU:HD21	9:Ff:738:TRP:CE2	2.46	0.51

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
9:Ff:837:PHE:O	9:Ff:841:ILE:HG12	2.11	0.51
2:Ab:2:VAL:HG23	3:Ac:1:ARG:HG3	1.93	0.50
6:Al:190:PRO:O	6:Al:231:ARG:HG2	2.11	0.50
5:Bc:253:ARG:HA	5:Bc:262:ILE:O	2.11	0.50
6:Bd:238:MET:HB3	6:Bj:250:TYR:HD1	1.76	0.50
5:Bo:246:LEU:HD12	5:Bo:247:ILE:N	2.26	0.50
6:Bp:134:TYR:HA	6:Bv:120:TYR:HE2	1.77	0.50
6:Bv:107:ILE:HD12	6:Bv:107:ILE:H	1.76	0.50
6:Bv:181:PHE:CD2	6:Bv:230:LEU:HD21	2.46	0.50
6:Cb:183:ASP:HB3	6:Cb:189:TRP:CE3	2.46	0.50
2:Cv:6:GLY:CA	6:Dl:125:GLU:HB3	2.41	0.50
6:Cz:106:VAL:HG23	6:Cz:110:LYS:HZ2	1.76	0.50
2:Db:2:VAL:HG13	3:Dc:1:ARG:HD2	1.93	0.50
6:Df:191:ILE:HD12	6:Df:206:TRP:NE1	2.26	0.50
7:Ee:805:ALA:O	7:Ee:809:VAL:HG12	2.11	0.50
8:Ew:42:TYR:HD2	8:Fa:43:LEU:HD11	1.76	0.50
8:Ew:423:ASN:HD22	8:Ex:425:GLN:HE22	1.57	0.50
8:Ex:25:ALA:HB1	8:Ex:30:GLN:HE22	1.75	0.50
8:Ex:433:GLN:HA	8:Ex:436:ILE:CD1	2.41	0.50
8:Ey:399:TRP:CE2	8:Ey:403:ILE:HD11	2.46	0.50
8:Ez:141:LEU:HB2	8:Ez:213:ASN:HB3	1.92	0.50
8:Ez:219:PRO:HG2	8:Ez:230:PHE:HB3	1.93	0.50
8:Ez:424:TYR:CZ	8:Fa:383:LEU:HB2	2.45	0.50
9:Fb:222:MET:HE1	9:Fb:519:TRP:HD1	1.76	0.50
9:Fd:617:ILE:HG22	9:Fd:618:LEU:H	1.76	0.50
9:Fd:659:PHE:HA	9:Fd:662:ILE:HD11	1.92	0.50
9:Fd:759:MET:CE	9:Fe:917:ILE:HD11	2.38	0.50
9:Fe:148:SER:O	9:Fe:152:ARG:HD3	2.11	0.50
9:Fe:332:ARG:O	9:Fe:336:ILE:HG13	2.11	0.50
9:Fe:589:THR:OG1	9:Fe:664:LEU:HD21	2.10	0.50
9:Fe:632:PHE:HZ	9:Ff:628:MET:HB3	1.76	0.50
9:Fe:729:MET:SD	9:Fe:730:MET:HB3	2.51	0.50
7:Fh:19:VAL:HB	7:Fh:23:PHE:CZ	2.46	0.50
6:Ar:168:VAL:HA	6:Ar:240:THR:HG23	1.94	0.50
6:Ax:158:VAL:HB	6:Ax:256:VAL:HA	1.93	0.50
5:Bi:246:LEU:HD12	5:Bi:247:ILE:N	2.26	0.50
6:Bj:201:SER:HB2	6:Bj:219:LYS:HD2	1.93	0.50
6:Bj:229:ARG:O	6:Bj:229:ARG:HD3	2.11	0.50
5:Bo:208:ILE:HG23	5:Bo:225:SER:HB2	1.93	0.50
5:Bu:219:ARG:NH1	5:Bu:233:ARG:HB3	2.26	0.50
5:Cg:219:ARG:HA	5:Cg:232:VAL:O	2.11	0.50

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
6:Df:106:VAL:O	6:Df:110:LYS:HG3	2.11	0.50
6:Dl:169:ILE:HD12	6:Dl:249:ASP:HB3	1.93	0.50
6:Ed:112:PHE:CE1	7:Et:801:MET:HE2	2.45	0.50
6:Ed:208:LYS:HE3	7:Ee:823:GLY:C	2.36	0.50
7:Eu:797:ARG:O	7:Eu:801:MET:HG2	2.12	0.50
8:Ew:77:VAL:HG12	8:Ew:378:MET:HE1	1.93	0.50
8:Ex:205:MET:SD	8:Ex:210:VAL:HB	2.52	0.50
8:Ex:409:ALA:O	8:Ex:413:LYS:HG3	2.12	0.50
8:Ey:136:ILE:CD1	8:Ey:277:LEU:HD23	2.42	0.50
8:Ez:67:GLN:HB3	8:Fa:404:ASN:OD1	2.12	0.50
8:Ez:179:PHE:CZ	8:Fa:233:TYR:HA	2.46	0.50
8:Ez:228:PRO:HG2	8:Ez:235:TYR:HB2	1.94	0.50
8:Fa:78:GLN:HE22	8:Fa:79:ASN:ND2	2.09	0.50
9:Fc:751:TYR:CE1	9:Fd:910:ALA:HA	2.46	0.50
9:Fd:57:MET:HA	9:Fd:60:MET:HB3	1.93	0.50
9:Fd:268:LYS:HB3	9:Fe:403:PRO:HG2	1.93	0.50
9:Fd:418:GLU:CD	9:Fd:418:GLU:H	2.19	0.50
9:Fd:464:ALA:HA	9:Fd:467:TYR:CE1	2.46	0.50
9:Fd:546:LYS:HE3	9:Fd:650:GLU:HG3	1.93	0.50
9:Fd:936:LEU:HA	9:Fd:939:LYS:HE3	1.94	0.50
9:Fe:258:SER:HA	9:Fe:285:ASN:HA	1.94	0.50
9:Fe:759:MET:HG2	9:Ff:917:ILE:HD11	1.92	0.50
9:Ff:71:LEU:HA	9:Ff:74:ILE:HD12	1.93	0.50
6:Af:105:GLU:O	6:Af:109:LYS:HE2	2.11	0.50
6:Af:107:ILE:HA	6:Af:110:LYS:HE2	1.92	0.50
6:Bv:208:LYS:HE2	7:Em:823:GLY:HA2	1.94	0.50
5:Cg:220:ALA:HB2	5:Cg:247:ILE:HD12	1.92	0.50
6:Ch:182:LEU:HB3	6:Ch:255:ARG:HE	1.77	0.50
3:Ck:2:VAL:HG21	6:Cz:129:LYS:HG2	1.94	0.50
4:Cr:2:PHE:HB2	5:Cs:251:GLN:HE21	1.76	0.50
6:Cz:177:SER:HA	6:Cz:250:TYR:O	2.12	0.50
6:Dx:114:ASP:O	6:Dx:117:ARG:HB3	2.12	0.50
7:Eq:798:THR:O	7:Eq:802:LEU:HD22	2.11	0.50
8:Ew:412:GLN:NE2	8:Ex:412:GLN:HB3	2.27	0.50
8:Ex:424:TYR:CE2	8:Ey:380:THR:HA	2.46	0.50
8:Ey:321:GLN:O	8:Ey:325:THR:HG23	2.11	0.50
8:Fa:185:PRO:HA	8:Fa:292:LEU:HD23	1.94	0.50
8:Fa:350:ILE:HA	8:Fa:353:LYS:HE2	1.94	0.50
9:Fb:191:MET:HE1	9:Fb:264:PRO:HG3	1.94	0.50
9:Fb:843:TYR:CZ	9:Ff:469:PHE:HB3	2.47	0.50
9:Fc:123:MET:O	9:Fc:126:VAL:HB	2.11	0.50

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
9:Fc:179:LYS:HD3	9:Fc:550:LEU:HD12	1.93	0.50
9:Fc:243:GLN:O	9:Fc:247:THR:HG22	2.11	0.50
9:Fc:514:SER:O	9:Fc:517:CYS:HB2	2.12	0.50
9:Fc:631:LEU:O	9:Fc:635:TYR:HB3	2.11	0.50
9:Fd:64:PHE:O	9:Fd:68:VAL:HG12	2.11	0.50
9:Fd:64:PHE:CZ	9:Fd:780:PRO:HG3	2.47	0.50
9:Fd:462:ILE:O	9:Fd:761:PHE:HZ	1.95	0.50
9:Ff:77:MET:SD	7:Fi:28:LEU:HD11	2.52	0.50
9:Ff:274:PHE:CZ	9:Ff:515:LEU:HD12	2.46	0.50
9:Ff:338:GLN:HA	9:Ff:341:VAL:HG22	1.92	0.50
9:Ff:498:PRO:O	9:Ff:501:LEU:HG	2.12	0.50
7:Fg:15:THR:HG22	7:Fg:18:ARG:HH22	1.76	0.50
1:Aa:11:TYR:CD2	3:Ac:9:TYR:HB3	2.46	0.50
6:Af:132:GLN:NE2	6:Af:136:THR:HG23	2.25	0.50
6:Af:191:ILE:HD12	6:Af:206:TRP:NE1	2.26	0.50
6:Ax:114:ASP:O	6:Ax:117:ARG:HG3	2.11	0.50
2:Bl:9:ARG:NH2	3:Bm:14:TYR:HB2	2.26	0.50
5:Bu:254:ILE:HB	5:Bu:262:ILE:CG2	2.42	0.50
6:Bv:199:PRO:HD2	7:Em:816:GLU:HB3	1.94	0.50
5:Cg:219:ARG:HH12	5:Cg:233:ARG:HD3	1.75	0.50
5:Cg:226:ASN:ND2	5:Cg:228:SER:HB3	2.26	0.50
6:Cn:115:MET:HE3	7:Em:801:MET:HB2	1.92	0.50
6:Ct:227:ALA:HA	6:Cz:251:ARG:HH22	1.75	0.50
6:Ed:160:LEU:HD12	6:Ed:160:LEU:O	2.11	0.50
7:Eq:781:GLN:HG3	9:Fb:341:VAL:HG11	1.94	0.50
9:Fb:223:ASN:O	9:Fb:227:ARG:HG3	2.11	0.50
9:Fc:770:ILE:HD11	9:Fd:928:LYS:NZ	2.26	0.50
9:Fc:908:VAL:O	9:Fc:911:PHE:HB3	2.11	0.50
9:Fe:77:MET:SD	7:Fj:28:LEU:HD21	2.52	0.50
9:Fe:734:LEU:O	9:Fe:737:ALA:HB3	2.11	0.50
9:Ff:455:GLU:O	9:Ff:459:LYS:HG2	2.12	0.50
9:Ff:516:LEU:O	9:Ff:520:PHE:HB2	2.11	0.50
9:Ff:940:VAL:HA	9:Ff:943:TRP:CE3	2.46	0.50
6:Al:229:ARG:HA	6:Al:236:PRO:HB3	1.94	0.50
5:Bc:247:ILE:HA	5:Bc:254:ILE:HD13	1.94	0.50
5:Bi:247:ILE:HG23	5:Bi:254:ILE:HD12	1.93	0.50
6:Df:173:GLN:HA	6:Df:217:ALA:O	2.11	0.50
6:Dl:221:TYR:HE1	6:Dr:142:ALA:HB3	1.76	0.50
6:Dr:180:VAL:HG12	6:Dr:212:THR:HG22	1.93	0.50
7:Ek:805:ALA:O	7:Ek:809:VAL:HG12	2.11	0.50
7:Et:789:LYS:HA	7:Et:789:LYS:HZ3	1.75	0.50

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
8:Ey:419:LEU:HD11	8:Ez:418:LEU:CD2	2.39	0.50
8:Ez:317:VAL:HG22	8:Fa:198:TRP:HD1	1.76	0.50
8:Ez:347:LEU:HA	8:Ez:350:ILE:HD12	1.93	0.50
8:Fa:281:ARG:NH1	8:Fa:281:ARG:HB2	2.25	0.50
8:Fa:425:GLN:HA	8:Fa:428:LEU:HG	1.94	0.50
9:Fb:64:PHE:O	9:Fb:68:VAL:HG13	2.11	0.50
9:Fb:103:ILE:HG22	9:Fb:106:ARG:HE	1.76	0.50
9:Fb:605:MET:HB3	9:Fb:641:TYR:HE2	1.76	0.50
9:Fb:905:TRP:HE1	9:Ff:702:THR:CG2	2.20	0.50
9:Fc:812:PRO:O	9:Fc:816:ILE:HG12	2.12	0.50
9:Fd:35:VAL:HA	9:Fd:38:LEU:HB2	1.93	0.50
9:Fd:596:PHE:HZ	9:Fd:653:ASN:HD22	1.59	0.50
9:Fe:139:ALA:HB3	9:Fe:461:TRP:NE1	2.26	0.50
7:Fh:30:ILE:O	7:Fh:34:ILE:HG22	2.11	0.50
2:Ab:4:MET:HE3	5:Ae:250:LEU:HG	1.92	0.50
4:Aj:4:ALA:HB3	5:Ak:251:GLN:NE2	2.27	0.50
3:Ao:2:VAL:HG21	6:Bd:129:LYS:CG	2.42	0.50
5:Bi:219:ARG:HG2	6:Bp:148:PRO:HD2	1.93	0.50
6:Bp:122:LEU:HD12	6:Bp:127:VAL:HG12	1.94	0.50
6:Cb:153:ALA:HB1	6:Cb:251:ARG:HH21	1.77	0.50
5:Cg:231:THR:HG21	6:Cn:151:PRO:HD2	1.94	0.50
6:Ch:223:TYR:HB2	6:Ch:242:ILE:HA	1.93	0.50
5:Cm:233:ARG:CZ	5:Cm:236:SER:HB3	2.41	0.50
6:Df:216:GLN:HB3	7:Er:818:GLN:NE2	2.26	0.50
6:Df:251:ARG:HG2	6:Df:251:ARG:HH11	1.75	0.50
5:Dk:219:ARG:HA	5:Dk:232:VAL:O	2.12	0.50
6:Dr:135:GLU:O	6:Dr:138:GLU:HG2	2.10	0.50
6:Ed:182:LEU:O	6:Ed:256:VAL:HG23	2.12	0.50
7:En:797:ARG:HD2	7:En:801:MET:HE1	1.94	0.50
7:Es:785:LEU:HD12	7:Es:789:LYS:HD3	1.93	0.50
7:Et:780:LYS:NZ	7:Eu:775:GLN:HE21	2.10	0.50
8:Ey:210:VAL:O	8:Ey:214:VAL:HG23	2.12	0.50
8:Ez:277:LEU:HA	8:Ez:280:ILE:HG22	1.93	0.50
9:Fb:651:MET:O	9:Fb:652:ILE:C	2.55	0.50
9:Fd:243:GLN:O	9:Fd:247:THR:HG22	2.12	0.50
9:Fd:735:LEU:HA	9:Fd:738:TRP:CZ3	2.46	0.50
9:Fe:420:LEU:HD22	9:Fe:420:LEU:H	1.77	0.50
3:Au:2:VAL:HG21	6:Bj:129:LYS:CD	2.40	0.50
5:Bc:213:GLN:HG2	5:Bc:221:TRP:O	2.11	0.50
1:Bq:3:ASP:OD1	3:Bs:14:TYR:HB3	2.12	0.50
6:Ch:200:SER:OG	7:Eo:816:GLU:HG2	2.11	0.50

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
6:Dr:166:PRO:HG3	6:Dx:250:TYR:CD1	2.46	0.50
7: Ei:780:LYS:HD3	7: Ei:780:LYS:H	1.77	0.50
8:Ew:298:TYR:HA	8:Ew:301:LEU:HD11	1.93	0.50
8:Ex:191:MET:HE1	8:Ex:198:TRP:CG	2.46	0.50
8:Ey:143:ASP:HB3	8:Ey:176:GLN:NE2	2.26	0.50
8:Ez:420:ALA:N	8:Fa:418:LEU:HD21	2.27	0.50
8:Fa:332:ASN:HA	8:Fa:335:VAL:HG22	1.93	0.50
9:Fb:185:LEU:O	9:Fb:189:ILE:HG12	2.12	0.50
9:Fc:222:MET:HA	9:Fc:222:MET:HE3	1.94	0.50
9:Fc:321:THR:HG23	9:Fc:324:GLN:HB2	1.93	0.50
9:Fc:595:ASN:HB3	9:Fc:597:LYS:HE2	1.94	0.50
9:Fc:926:VAL:O	9:Fc:929:ALA:HB3	2.12	0.50
9:Fe:260:GLU:HG3	9:Fe:283:LYS:HG2	1.93	0.50
6:Af:238:MET:HG2	6:Al:251:ARG:NH2	2.27	0.50
6:Al:115:MET:O	6:Al:119:LEU:HD12	2.12	0.50
6:Al:229:ARG:HD2	6:Al:229:ARG:C	2.37	0.50
1: Ci:1:CYS:SG	4: Cl:2:PHE:HB3	2.52	0.50
5:Cm:220:ALA:HB3	5:Cm:232:VAL:HG13	1.94	0.50
6:Cn:114:ASP:O	6:Cn:117:ARG:HB3	2.12	0.50
6:Cz:129:LYS:O	6:Cz:133:ILE:HG13	2.11	0.50
6:Df:225:ASN:HD21	6:Dl:176:VAL:H	1.57	0.50
5:Dk:264:PHE:CD1	6:Dl:246:LYS:HE3	2.46	0.50
7:Es:797:ARG:O	7:Es:801:MET:HG3	2.11	0.50
8:Ex:424:TYR:O	8:Ex:427:TYR:HB3	2.12	0.50
8:Ey:220:PRO:HD2	8:Ey:228:PRO:O	2.12	0.50
8:Ez:418:LEU:HA	8:Ez:421:GLU:CD	2.36	0.50
9:Fb:449:SER:O	9:Fb:452:PHE:HE1	1.95	0.50
9:Fb:493:LYS:HA	9:Fb:493:LYS:HE3	1.93	0.50
9:Fc:641:TYR:CE1	9:Fc:645:LEU:HG	2.47	0.50
9:Fd:670:ILE:HA	9:Fd:673:VAL:HG22	1.94	0.50
9:Fe:67:ALA:O	9:Fe:71:LEU:HD12	2.11	0.50
9:Fe:915:ILE:HA	9:Fe:918:TYR:HB2	1.94	0.50
9:Ff:339:MET:HE1	9:Ff:422:ALA:O	2.12	0.50
9:Ff:514:SER:O	9:Ff:517:CYS:HB2	2.11	0.50
2:Ah:2:VAL:HA	3: Ai:1:ARG:HH21	1.76	0.50
6:Al:204:ILE:HD11	6:Al:213:LEU:HB3	1.92	0.50
5:Aw:220:ALA:HB2	5:Aw:247:ILE:HD12	1.94	0.50
6:Ax:169:ILE:HD11	6:Ax:239:LEU:HB3	1.94	0.50
6:Ax:199:PRO:HD2	7: Ei:816:GLU:HB2	1.94	0.50
6:Bp:196:LEU:HD22	6:Bp:199:PRO:HA	1.94	0.50
1:Bq:16:ALA:HB3	2:Bx:9:ARG:HG3	1.94	0.50

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
6:Bv:147:THR:HG23	6:Bv:246:LYS:HZ1	1.76	0.50
6:Ch:181:PHE:CE2	6:Ch:256:VAL:HG11	2.47	0.50
6:Cn:167:PRO:HB2	6:Cn:239:LEU:HD23	1.94	0.50
6:Cz:115:MET:CE	7:Eo:797:ARG:HH21	2.24	0.50
6:Ed:109:LYS:O	6:Ed:113:LYS:HG2	2.12	0.50
7:Ei:787:GLU:O	7:Ei:791:GLN:HB2	2.12	0.50
7:Es:780:LYS:HA	7:Es:783:GLU:HG2	1.94	0.50
8:Ew:48:LYS:HG2	8:Fa:64:PRO:HG3	1.93	0.50
8:Ew:425:GLN:CG	8:Fa:426:MET:HE1	2.30	0.50
8:Ex:317:VAL:HG23	8:Ex:318:GLN:OE1	2.11	0.50
8:Ez:320:LYS:HZ3	8:Fa:198:TRP:CG	2.29	0.50
9:Fb:709:ASN:HD21	9:Fc:597:LYS:H	1.59	0.50
9:Fb:753:ILE:HG23	9:Fb:926:VAL:HG11	1.94	0.50
9:Fc:339:MET:CE	9:Fc:426:TYR:HB2	2.41	0.50
9:Fc:738:TRP:HD1	9:Fc:912:PHE:HE2	1.60	0.50
9:Fd:507:LYS:HD2	9:Fe:635:TYR:HB2	1.93	0.50
9:Fe:182:LEU:HD23	9:Fe:571:VAL:HG21	1.93	0.50
9:Fe:243:GLN:O	9:Fe:247:THR:HG22	2.12	0.50
9:Fe:667:MET:CE	9:Fe:915:ILE:HD11	2.42	0.50
9:Ff:354:ASP:HB3	9:Ff:357:PHE:HD1	1.77	0.50
6:Af:110:LYS:HA	6:Af:113:LYS:HE2	1.94	0.49
6:Af:130:LEU:HD21	7:Ev:805:ALA:HB1	1.94	0.49
6:Bp:150:LYS:N	6:Bp:247:ALA:HA	2.27	0.49
6:Bp:189:TRP:CG	6:Bp:230:LEU:HD12	2.47	0.49
6:Bp:194:TYR:HE1	6:Bp:196:LEU:HB2	1.76	0.49
6:Bp:209:THR:HG22	7:El:824:THR:HG21	1.93	0.49
6:Cb:130:LEU:HD23	7:El:808:LEU:HD12	1.94	0.49
6:Cn:237:VAL:C	6:Ct:251:ARG:HH22	2.20	0.49
2:Cp:5:ILE:HD11	6:Cz:136:THR:O	2.11	0.49
6:Df:221:TYR:CE2	6:Dl:139:TYR:HB2	2.47	0.49
6:Df:236:PRO:C	6:Dl:251:ARG:NH2	2.70	0.49
6:Dl:111:ALA:HA	6:Dl:114:ASP:OD1	2.12	0.49
6:Dl:189:TRP:CG	6:Dl:230:LEU:HD13	2.47	0.49
5:Dq:213:GLN:NE2	5:Dq:223:ILE:HB	2.27	0.49
7:Eq:807:GLN:H	7:Eq:807:GLN:CD	2.19	0.49
8:Ex:215:ILE:HD12	8:Ex:215:ILE:H	1.76	0.49
8:Ey:174:VAL:O	8:Ey:178:VAL:HG12	2.11	0.49
8:Fa:300:GLU:HA	8:Fa:303:ASN:HD21	1.76	0.49
9:Fc:461:TRP:CZ3	9:Fc:812:PRO:HB3	2.47	0.49
9:Fc:825:LEU:HD22	9:Fc:828:VAL:HG23	1.94	0.49
9:Fd:596:PHE:CD1	9:Fd:656:VAL:HG11	2.47	0.49

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
9:Ff:605:MET:HB2	9:Ff:638:ARG:HG2	1.94	0.49
9:Ff:762:THR:HB	9:Ff:766:PHE:CZ	2.47	0.49
7:Fj:24:THR:O	7:Fj:27:LEU:HD12	2.11	0.49
6:Af:129:LYS:HG2	3:Du:2:VAL:HG21	1.93	0.49
6:Ar:207:ASP:OD2	6:Ar:210:SER:HB3	2.12	0.49
6:Ar:220:LEU:HD12	6:Ax:138:GLU:OE2	2.12	0.49
6:Df:108:ASP:HA	7:Ep:797:ARG:NH1	2.27	0.49
6:Df:190:PRO:O	6:Df:230:LEU:HD12	2.12	0.49
3:Du:7:THR:HB	3:Du:9:TYR:CE2	2.48	0.49
6:Dx:204:ILE:CD1	6:Dx:215:ILE:HD13	2.42	0.49
7:El:804:ALA:O	7:El:807:GLN:HB3	2.12	0.49
8:Ex:121:LYS:HA	8:Ex:121:LYS:HE3	1.94	0.49
8:Ey:134:GLY:N	8:Ey:258:ALA:HB2	2.28	0.49
9:Fc:697:ILE:HG13	9:Fc:698:ASN:N	2.27	0.49
9:Fd:760:ILE:HG22	9:Fd:933:ILE:HD11	1.95	0.49
9:Fe:73:GLY:HA2	9:Fe:76:ILE:HG12	1.94	0.49
9:Fe:702:THR:HG23	9:Ff:905:TRP:NE1	2.26	0.49
9:Ff:56:ILE:HG13	9:Ff:57:MET:SD	2.51	0.49
9:Ff:335:ALA:HA	9:Ff:338:GLN:CD	2.37	0.49
6:Af:132:GLN:HG2	3:Du:4:ILE:HA	1.94	0.49
6:Al:230:LEU:HD12	6:Al:231:ARG:N	2.25	0.49
6:Ar:246:LYS:HA	6:Ar:246:LYS:HE2	1.94	0.49
6:Bd:179:LEU:HD12	6:Bd:254:LEU:HD11	1.94	0.49
5:Bu:244:VAL:HA	5:Bu:256:THR:HA	1.93	0.49
6:Bv:136:THR:HA	6:Bv:139:TYR:CE2	2.47	0.49
6:Cn:214:MET:CE	7:Eo:818:GLN:HE21	2.21	0.49
6:Cz:216:GLN:HB3	7:Eq:818:GLN:CD	2.38	0.49
6:Cz:216:GLN:HB3	7:Eq:818:GLN:OE1	2.12	0.49
6:Df:238:MET:HG2	6:Dl:251:ARG:CZ	2.42	0.49
7:Ef:773:GLN:O	7:Ef:777:ILE:HG12	2.12	0.49
7:Ep:797:ARG:HG2	7:Ep:801:MET:HG2	1.94	0.49
7:Eq:808:LEU:O	7:Eq:812:TRP:CD1	2.65	0.49
8:Ey:55:THR:HA	8:Ez:42:TYR:CE1	2.47	0.49
8:Ey:228:PRO:HB2	8:Ey:235:TYR:HB2	1.94	0.49
8:Ey:317:VAL:HA	8:Ez:198:TRP:CZ3	2.40	0.49
8:Ez:298:TYR:HA	8:Ez:301:LEU:CD1	2.42	0.49
8:Fa:297:ALA:HA	8:Fa:300:GLU:OE1	2.13	0.49
9:Fc:112:ALA:HA	9:Fc:115:ILE:HD11	1.94	0.49
9:Fe:273:TYR:O	9:Fe:276:ASN:HB2	2.12	0.49
9:Fe:433:THR:O	9:Fe:437:ILE:HG12	2.12	0.49
9:Ff:94:LEU:HD13	9:Ff:98:TRP:CD1	2.47	0.49

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
9:Ff:766:PHE:O	9:Ff:770:ILE:HG12	2.12	0.49
6:Bj:168:VAL:HG13	6:Bj:242:ILE:HD13	1.95	0.49
6:Bj:189:TRP:CE2	6:Bj:260:GLY:HA2	2.47	0.49
5:Bu:211:TYR:HA	5:Bu:262:ILE:HD11	1.93	0.49
6:Bv:246:LYS:O	6:Bv:246:LYS:HD3	2.12	0.49
6:Cn:128:VAL:HG23	6:Cn:132:GLN:NE2	2.24	0.49
6:Cz:145:PRO:HG3	6:Df:131:LYS:HG2	1.94	0.49
6:Cz:178:SER:HB3	6:Cz:251:ARG:CD	2.43	0.49
5:Dk:217:PRO:HG2	6:Dr:143:ALA:HB1	1.94	0.49
6:Dx:204:ILE:HA	6:Dx:214:MET:O	2.12	0.49
5:Ec:253:ARG:HH21	5:Ec:263:LYS:HD2	1.78	0.49
7:Eh:792:GLN:HB2	7:Eh:795:GLN:HE21	1.77	0.49
7:El:805:ALA:O	7:El:809:VAL:HG12	2.12	0.49
7:Es:771:ALA:HA	7:Es:774:LEU:HD23	1.95	0.49
8:Ew:246:ASN:ND2	8:Ew:250:ALA:HB3	2.27	0.49
8:Ew:334:ARG:HD2	8:Ex:282:TYR:OH	2.12	0.49
8:Ew:448:LEU:HD13	8:Ex:339:GLN:HB3	1.95	0.49
8:Ex:409:ALA:O	8:Ex:412:GLN:HG2	2.11	0.49
8:Ey:246:ASN:HA	8:Ey:249:ILE:HG22	1.94	0.49
8:Ey:253:LEU:C	8:Ey:254:MET:HE2	2.37	0.49
8:Ey:408:PRO:O	8:Ey:411:VAL:HG22	2.12	0.49
9:Fb:643:PHE:CE2	9:Fc:605:MET:HE1	2.48	0.49
9:Fb:935:HIS:O	9:Fb:939:LYS:HG2	2.12	0.49
9:Fc:588:LEU:HD12	9:Fc:661:MET:HE3	1.95	0.49
9:Fd:79:THR:HA	9:Fd:82:VAL:HG22	1.95	0.49
9:Fd:263:MET:HA	9:Fd:264:PRO:C	2.38	0.49
9:Fd:819:TYR:O	9:Fd:823:ILE:HG12	2.12	0.49
9:Ff:109:PHE:O	9:Ff:113:LEU:HD13	2.12	0.49
9:Ff:596:PHE:CE2	9:Ff:656:VAL:HG21	2.47	0.49
7:Fg:34:ILE:HG23	7:Fg:38:LYS:HZ1	1.77	0.49
6:Bj:188:PRO:HB3	6:Bj:211:ASN:ND2	2.27	0.49
6:Ch:181:PHE:HE2	6:Ch:237:VAL:HG11	1.78	0.49
6:Cn:121:PRO:HG2	7:En:801:MET:HG3	1.95	0.49
5:Cs:219:ARG:HD2	6:Cz:147:THR:HG23	1.93	0.49
5:De:219:ARG:NH1	5:De:233:ARG:HB3	2.27	0.49
5:Dq:250:LEU:HD12	5:Dq:251:GLN:HG3	1.95	0.49
6:Dr:196:LEU:HD12	6:Dr:204:ILE:HD12	1.94	0.49
6:Dr:223:TYR:HB2	6:Dr:242:ILE:HG13	1.94	0.49
7:El:792:GLN:HA	7:El:795:GLN:NE2	2.28	0.49
7:Eq:797:ARG:HD2	7:Eq:797:ARG:C	2.37	0.49
8:Ew:307:ALA:HB1	8:Ew:314:TYR:CD1	2.48	0.49

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
8:Ew:339:GLN:HE22	8:Fa:448:LEU:HB2	1.77	0.49
8:Ex:261:SER:HA	8:Ex:270:LYS:HE3	1.94	0.49
8:Ez:331:ASN:HD21	8:Fa:211:MET:HB2	1.77	0.49
8:Ez:375:GLU:O	8:Ez:378:MET:HE3	2.12	0.49
8:Fa:400:ILE:HG13	8:Fa:401:LYS:HD2	1.95	0.49
9:Fb:113:LEU:O	9:Fb:122:CYS:HB3	2.13	0.49
9:Fb:664:LEU:HA	9:Fb:667:MET:HG3	1.95	0.49
9:Fc:704:TRP:CH2	9:Fd:733:PRO:HG2	2.47	0.49
9:Fc:735:LEU:HD23	9:Fc:738:TRP:CZ3	2.47	0.49
9:Fc:909:TYR:O	9:Fc:912:PHE:HB2	2.12	0.49
9:Fd:667:MET:HA	9:Fd:670:ILE:HD11	1.94	0.49
9:Fe:546:LYS:N	9:Fe:546:LYS:HD3	2.27	0.49
9:Fe:590:PHE:HB3	9:Fe:593:LEU:HB3	1.93	0.49
9:Fe:653:ASN:O	9:Fe:657:MET:HG2	2.12	0.49
9:Ff:234:ILE:H	9:Ff:234:ILE:HD12	1.78	0.49
9:Ff:720:PHE:O	9:Ff:723:PHE:HD2	1.94	0.49
6:Af:181:PHE:CZ	6:Af:213:LEU:HD12	2.45	0.49
2:Bl:1:CYS:HA	3:Bm:4:ILE:HD13	1.94	0.49
6:Cb:221:TYR:CD1	6:Ch:139:TYR:HD1	2.30	0.49
5:Cs:245:LYS:NZ	5:Cs:257:SER:HB3	2.27	0.49
6:Ct:233:LEU:HD23	6:Ct:234:ASN:H	1.77	0.49
6:Df:108:ASP:HA	7:Ep:797:ARG:HH12	1.78	0.49
5:Dk:208:ILE:HG23	5:Dk:225:SER:HB2	1.94	0.49
7:Eg:810:GLN:HA	7:Eg:813:LYS:HE2	1.95	0.49
7:En:807:GLN:NE2	7:En:808:LEU:HD22	2.28	0.49
8:Ew:206:TYR:OH	8:Ew:208:ASN:HB3	2.12	0.49
8:Ex:94:MET:HE3	8:Ey:241:SER:HB2	1.94	0.49
8:Fa:445:LEU:HA	8:Fa:448:LEU:HG	1.93	0.49
9:Fb:45:VAL:HG11	9:Fb:129:MET:HG2	1.94	0.49
9:Fc:185:LEU:O	9:Fc:189:ILE:HG12	2.13	0.49
9:Fc:465:GLY:HA3	9:Fc:761:PHE:HD2	1.78	0.49
9:Fe:338:GLN:NE2	9:Fe:425:ASP:HB2	2.28	0.49
9:Fe:807:ASN:HB2	9:Fe:941:LEU:HD11	1.95	0.49
7:Fg:31:ALA:HA	7:Fg:34:ILE:HD12	1.95	0.49
4:Ap:5:ASP:HA	5:Aq:251:GLN:HG2	1.95	0.49
6:Ar:166:PRO:HG3	6:Ax:250:TYR:HD1	1.77	0.49
3:Bs:1:ARG:HG2	3:Bs:2:VAL:N	2.28	0.49
5:Ca:219:ARG:NH1	5:Ca:233:ARG:HB3	2.28	0.49
6:Cb:129:LYS:HD2	6:Ch:112:PHE:CZ	2.48	0.49
1:Cc:6:LEU:O	1:Cc:10:GLU:HG2	2.12	0.49
6:Df:238:MET:N	6:Dl:251:ARG:HH22	2.11	0.49

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:Dg:13:LYS:N	1:Dg:13:LYS:HD3	2.28	0.49
6:Dl:236:PRO:HG2	6:Dr:251:ARG:NE	2.27	0.49
2:Dn:4:MET:HE1	5:Dq:248:ASP:OD1	2.13	0.49
6:Dr:122:LEU:HD13	6:Dr:126:GLN:HB3	1.95	0.49
8:Ex:42:TYR:HA	8:Ex:45:ASN:ND2	2.28	0.49
8:Ex:120:PHE:HA	8:Ex:123:GLN:HB2	1.95	0.49
8:Ex:346:ASN:O	8:Ex:350:ILE:HG22	2.13	0.49
8:Ey:35:ASN:O	8:Ey:39:LEU:HD22	2.13	0.49
8:Ez:350:ILE:HG22	8:Ez:354:ARG:HH21	1.78	0.49
9:Fb:420:LEU:HD22	9:Fb:420:LEU:H	1.76	0.49
9:Fb:661:MET:HB3	9:Fb:665:GLN:CG	2.43	0.49
9:Fb:776:MET:C	9:Fb:776:MET:HE2	2.37	0.49
9:Fc:124:MET:HG3	9:Fc:128:PHE:CE1	2.35	0.49
9:Fc:124:MET:HE3	9:Fc:783:ALA:HB2	1.95	0.49
9:Fc:136:VAL:HG22	9:Fc:461:TRP:CZ2	2.46	0.49
9:Fd:708:LEU:HD13	9:Fe:730:MET:SD	2.52	0.49
9:Fd:768:TRP:HA	9:Fd:811:ARG:HH12	1.77	0.49
9:Fe:725:PHE:O	9:Fe:728:ILE:HG22	2.12	0.49
9:Fe:744:SER:O	9:Fe:748:VAL:HG23	2.11	0.49
9:Ff:117:LYS:HZ3	9:Ff:119:SER:H	1.60	0.49
9:Ff:693:GLY:O	9:Ff:697:ILE:HG12	2.13	0.49
6:Al:106:VAL:HG23	6:Al:110:LYS:NZ	2.27	0.49
3:Ao:2:VAL:HG21	6:Bd:129:LYS:HD3	1.95	0.49
6:Ax:115:MET:HE3	7:Ef:798:THR:HA	1.95	0.49
6:Bp:204:ILE:CG1	6:Bp:215:ILE:HD13	2.42	0.49
5:Ca:231:THR:HG21	6:Ch:151:PRO:HG2	1.94	0.49
6:Cb:110:LYS:HA	6:Cb:113:LYS:HG2	1.94	0.49
6:Ch:207:ASP:HB3	7:En:820:TYR:OH	2.13	0.49
6:Ct:225:ASN:HB2	6:Cz:250:TYR:OH	2.12	0.49
1:Dg:9:LEU:CD1	1:Dg:13:LYS:HE2	2.42	0.49
1:Dg:16:ALA:HA	4:Dp:3:GLY:HA3	1.93	0.49
6:Dl:169:ILE:HD13	6:Dl:252:VAL:HG21	1.95	0.49
6:Dl:169:ILE:HG13	6:Dl:171:LEU:HG	1.94	0.49
6:Dr:126:GLN:HA	6:Dr:129:LYS:NZ	2.28	0.49
8:Ew:415:ILE:O	8:Ew:419:LEU:HG	2.12	0.49
8:Ex:430:ARG:HG2	8:Ey:429:ASP:HB2	1.95	0.49
8:Ez:189:PHE:HE2	8:Fa:231:TYR:CE2	2.30	0.49
8:Ez:320:LYS:HZ3	8:Fa:198:TRP:CD1	2.31	0.49
9:Fb:106:ARG:NH2	9:Fb:789:PRO:HB3	2.28	0.49
9:Fb:147:LEU:HB2	9:Fb:453:ILE:HG23	1.93	0.49
9:Fb:597:LYS:HD3	9:Fb:597:LYS:N	2.27	0.49

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
9:Fb:915:ILE:O	9:Fb:919:THR:HG23	2.13	0.49
9:Fc:672:ILE:O	9:Fc:675:VAL:HG12	2.13	0.49
9:Fd:730:MET:C	9:Fd:730:MET:HE2	2.38	0.49
9:Fe:383:LYS:HG2	9:Fe:387:GLU:H	1.78	0.49
9:Fe:697:ILE:HD11	9:Fe:746:GLY:HA3	1.95	0.49
3:Ac:4:ILE:O	6:Ar:132:GLN:HB3	2.12	0.49
6:Bd:225:ASN:HD21	6:Bj:176:VAL:HG23	1.78	0.49
6:Bj:187:ALA:HB1	6:Bj:262:ASN:HB2	1.93	0.49
6:Bp:135:GLU:HA	6:Bp:138:GLU:OE1	2.13	0.49
2:Br:9:ARG:HD2	3:Bs:15:ASP:HA	1.95	0.49
6:Bv:111:ALA:O	6:Bv:115:MET:HG3	2.12	0.49
5:Cg:233:ARG:HG2	5:Cg:236:SER:OG	2.13	0.49
6:Ch:126:GLN:HB2	6:Cn:112:PHE:CE2	2.48	0.49
5:Cs:264:PHE:HB3	5:Cs:269:SER:HB3	1.94	0.49
6:Ct:128:VAL:CG2	6:Ct:132:GLN:HE21	2.23	0.49
6:Ct:220:LEU:HG	6:Ct:221:TYR:CE2	2.48	0.49
5:Cy:219:ARG:HD3	6:Df:148:PRO:O	2.13	0.49
6:Cz:126:GLN:HA	6:Cz:129:LYS:HD3	1.95	0.49
1:Ds:1:CYS:HA	4:Dv:2:PHE:CD1	2.47	0.49
6:Dx:225:ASN:ND2	6:Ed:176:VAL:H	2.08	0.49
7:Er:809:VAL:HB	7:Er:813:LYS:NZ	2.28	0.49
8:Ew:433:GLN:HE22	8:Fa:433:GLN:HE21	1.61	0.49
8:Ex:129:SER:HB2	8:Ex:137:THR:HG21	1.94	0.49
8:Ex:246:ASN:O	8:Ex:251:PRO:HD3	2.13	0.49
8:Ex:385:ASP:HB3	8:Ex:388:ALA:HB2	1.95	0.49
8:Fa:104:ASP:HA	8:Fa:109:ASN:ND2	2.27	0.49
8:Fa:173:PRO:HA	8:Fa:176:GLN:HB3	1.95	0.49
8:Fa:254:MET:N	8:Fa:254:MET:HE2	2.27	0.49
9:Fb:693:GLY:HA2	9:Fb:696:TYR:CD2	2.48	0.49
9:Fb:825:LEU:HA	9:Fb:828:VAL:HG23	1.95	0.49
9:Fd:49:LEU:HD12	9:Fd:50:HIS:H	1.78	0.49
9:Fd:641:TYR:CE1	9:Fd:645:LEU:HD21	2.48	0.49
9:Fe:261:LEU:HD11	9:Fe:265:ASN:ND2	2.23	0.49
9:Fe:695:MET:HB3	9:Fe:699:PHE:CE1	2.47	0.49
9:Fe:756:LEU:HA	9:Fe:759:MET:HE1	1.93	0.49
6:Al:135:GLU:HA	6:Al:138:GLU:HG3	1.94	0.49
6:Al:226:LEU:HD12	6:Al:239:LEU:HB2	1.94	0.49
6:Ar:181:PHE:CD1	6:Ar:254:LEU:HD11	2.38	0.49
5:Bc:268:ASP:OD1	6:Bd:152:THR:HG21	2.13	0.49
5:Ca:219:ARG:HD3	6:Ch:148:PRO:O	2.13	0.49
6:Ct:134:TYR:HE1	7:Eo:812:TRP:HB2	1.78	0.49

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
6:Ct:141:LYS:HZ3	7:Ep:811:ASP:HB2	1.78	0.49
6:Cz:129:LYS:HE2	6:Df:112:PHE:CE2	2.48	0.49
6:Dx:182:LEU:HD11	6:Dx:186:GLY:HA2	1.95	0.49
6:Ed:203:ASN:HB3	6:Ed:216:GLN:HG2	1.95	0.49
7:Es:807:GLN:HE21	7:Es:808:LEU:CD2	2.20	0.49
7:Ev:799:SER:HA	7:Ev:802:LEU:HD23	1.95	0.49
8:Ew:294:LYS:O	8:Ew:298:TYR:HB3	2.12	0.49
8:Ex:277:LEU:HD13	8:Ex:281:ARG:NH2	2.25	0.49
8:Ey:170:LEU:HD12	8:Ey:171:ASN:H	1.78	0.49
9:Fd:757:PRO:HG3	9:Fd:930:PHE:CE2	2.48	0.49
9:Fe:191:MET:HB3	9:Fe:233:PHE:HB2	1.95	0.49
9:Fe:339:MET:HE1	9:Fe:422:ALA:O	2.13	0.49
9:Ff:525:ASP:OD1	9:Ff:526:LYS:HD3	2.12	0.49
9:Ff:804:ILE:O	9:Ff:807:ASN:HB3	2.12	0.49
9:Ff:809:PHE:CD1	9:Ff:810:LEU:HD23	2.47	0.49
5:Bc:210:TYR:HB3	5:Bc:222:LEU:HD12	1.94	0.48
6:Bd:225:ASN:ND2	6:Bj:176:VAL:H	2.11	0.48
3:Bm:1:ARG:N	3:Bm:10:THR:HG23	2.28	0.48
6:Cb:194:TYR:HA	6:Cb:229:ARG:HH12	1.77	0.48
5:Cg:246:LEU:HD12	5:Cg:247:ILE:N	2.27	0.48
6:Cn:141:LYS:C	6:Cn:141:LYS:HD3	2.38	0.48
6:Df:181:PHE:HD1	6:Df:254:LEU:HD12	1.78	0.48
6:Ed:106:VAL:HG13	6:Ed:110:LYS:HZ2	1.77	0.48
6:Ed:141:LYS:NZ	7:Ev:811:ASP:HB2	2.28	0.48
7:Eh:822:GLU:OE1	7:Eh:822:GLU:HA	2.12	0.48
7:Em:787:GLU:O	7:Em:791:GLN:HB2	2.13	0.48
7:Eo:811:ASP:HB2	7:Eo:812:TRP:CE3	2.49	0.48
8:Ew:242:GLN:HE22	8:Ew:269:ALA:N	2.11	0.48
8:Ez:453:PRO:HA	9:Ff:619:PHE:HZ	1.78	0.48
8:Fa:381:ARG:HG3	8:Fa:382:ARG:CD	2.43	0.48
9:Fb:661:MET:HB3	9:Fb:665:GLN:NE2	2.29	0.48
9:Fb:686:ILE:O	9:Fb:689:LEU:HG	2.12	0.48
9:Fb:723:PHE:CE1	9:Ff:716:LEU:HA	2.48	0.48
9:Fb:811:ARG:HG2	9:Fb:815:MET:CE	2.43	0.48
9:Fc:827:TYR:HB3	9:Fc:831:TRP:CZ2	2.48	0.48
9:Fd:624:LEU:O	9:Fd:628:MET:HG2	2.12	0.48
9:Fd:709:ASN:HD21	9:Fe:597:LYS:N	2.02	0.48
9:Fe:644:PHE:HE1	9:Fe:718:PRO:HB2	1.78	0.48
9:Ff:921:MET:SD	9:Ff:921:MET:N	2.85	0.48
6:Al:110:LYS:HA	6:Al:113:LYS:HG2	1.94	0.48
6:Ax:189:TRP:CD1	6:Ax:230:LEU:HD12	2.48	0.48

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
6:Bj:190:PRO:O	6:Bj:231:ARG:HG2	2.13	0.48
5:Bo:219:ARG:NH1	5:Bo:233:ARG:HB3	2.28	0.48
6:Bp:221:TYR:HD2	6:Bp:244:GLY:HA3	1.77	0.48
5:Cg:212:ILE:HG21	5:Cg:215:VAL:HG23	1.95	0.48
5:Cs:219:ARG:HG2	6:Cz:148:PRO:HD2	1.95	0.48
5:Cy:212:ILE:HB	5:Cy:264:PHE:CD2	2.48	0.48
6:Df:138:GLU:HA	6:Df:141:LYS:HB3	1.95	0.48
6:Dl:141:LYS:HE3	7:Es:812:TRP:CD2	2.49	0.48
6:Dx:111:ALA:HB3	7:Es:797:ARG:HH21	1.78	0.48
8:Ew:286:GLN:HE22	8:Ew:287:VAL:HG13	1.77	0.48
8:Ez:420:ALA:HB2	8:Fa:399:TRP:CZ3	2.48	0.48
9:Fb:37:PHE:HE1	9:Fc:831:TRP:HB2	1.78	0.48
9:Fc:263:MET:HE2	9:Fc:279:CYS:O	2.14	0.48
9:Fc:531:GLN:HA	9:Fc:534:ILE:HD12	1.95	0.48
9:Fc:710:MET:HA	9:Fc:714:SER:OG	2.14	0.48
9:Fd:556:ARG:NH1	9:Fd:575:MET:HG3	2.27	0.48
9:Fd:816:ILE:O	9:Fd:820:ILE:HG13	2.14	0.48
9:Fe:322:SER:O	9:Fe:326:GLN:HG3	2.12	0.48
9:Fe:593:LEU:HD23	9:Fe:594:ILE:CD1	2.43	0.48
9:Ff:113:LEU:O	9:Ff:122:CYS:HB3	2.13	0.48
9:Ff:167:LEU:HD21	9:Ff:178:ALA:H	1.78	0.48
9:Ff:339:MET:CE	9:Ff:426:TYR:HB2	2.43	0.48
9:Ff:600:THR:HG22	9:Ff:648:PHE:HE2	1.77	0.48
5:Ae:250:LEU:HD12	5:Ae:251:GLN:HG3	1.94	0.48
6:Al:201:SER:C	6:Al:202:PHE:HD1	2.20	0.48
6:Bd:117:ARG:HH12	6:Bd:121:PRO:HB3	1.78	0.48
5:Bi:236:SER:H	5:Bi:244:VAL:HG12	1.78	0.48
6:Cb:238:MET:HG3	6:Ch:250:TYR:CB	2.44	0.48
6:Ch:196:LEU:HD11	6:Ch:202:PHE:HD2	1.78	0.48
1:Co:13:LYS:HD2	1:Co:13:LYS:C	2.38	0.48
6:Ct:134:TYR:CE1	7:Eo:812:TRP:HB2	2.47	0.48
6:Ct:242:ILE:HD12	6:Ct:243:PRO:HD2	1.94	0.48
5:De:211:TYR:O	5:De:223:ILE:HG22	2.12	0.48
5:Dw:219:ARG:HA	5:Dw:232:VAL:O	2.12	0.48
6:Dx:125:GLU:C	6:Dx:129:LYS:HZ2	2.21	0.48
7:Ee:809:VAL:HA	7:Ee:812:TRP:CE3	2.48	0.48
8:Ew:381:ARG:HG3	8:Ew:382:ARG:NH1	2.28	0.48
8:Ew:423:ASN:ND2	8:Ex:425:GLN:HE22	2.12	0.48
8:Ew:424:TYR:CZ	8:Ex:379:ALA:O	2.66	0.48
8:Ex:239:LEU:H	8:Ex:239:LEU:HD12	1.78	0.48
8:Ey:174:VAL:HG21	8:Ez:246:ASN:ND2	2.28	0.48

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
8:Fa:187:TYR:HA	8:Fa:206:TYR:CG	2.48	0.48
9:Fb:149:TYR:HB2	9:Ff:59:ASN:HD22	1.78	0.48
9:Fb:187:GLY:HA2	9:Fb:496:PHE:HE2	1.76	0.48
9:Fb:761:PHE:HE1	9:Fb:815:MET:HG3	1.78	0.48
9:Fc:588:LEU:HB3	9:Fc:665:GLN:CD	2.38	0.48
9:Fd:369:ASP:HB3	9:Fd:375:LYS:HA	1.95	0.48
9:Fd:469:PHE:O	9:Fd:469:PHE:CD1	2.67	0.48
9:Fd:732:MET:O	9:Fd:736:MET:CB	2.60	0.48
9:Fd:743:VAL:CG2	9:Fd:747:PHE:HE1	2.23	0.48
9:Fe:907:GLY:O	9:Fe:910:ALA:HB3	2.13	0.48
9:Ff:778:ALA:O	9:Ff:782:VAL:HG23	2.13	0.48
6:Af:125:GLU:O	6:Af:129:LYS:HG3	2.14	0.48
6:Ar:132:GLN:HA	6:Ar:135:GLU:OE2	2.13	0.48
6:Bj:219:LYS:HD3	6:Bj:222:ASN:ND2	2.28	0.48
5:Bo:212:ILE:HB	5:Bo:264:PHE:CD1	2.48	0.48
5:Bo:246:LEU:HD12	5:Bo:247:ILE:H	1.77	0.48
6:Cn:135:GLU:HA	6:Cn:138:GLU:OE2	2.13	0.48
6:Cn:166:PRO:HG3	6:Ct:250:TYR:CD2	2.49	0.48
6:Dx:190:PRO:HA	6:Dx:211:ASN:HB3	1.95	0.48
6:Ed:203:ASN:HD21	7:Ev:818:GLN:HE22	1.61	0.48
7:Eg:807:GLN:CD	7:Eg:807:GLN:H	2.20	0.48
8:Ew:338:ALA:HB2	8:Ex:243:LEU:HD22	1.94	0.48
8:Ex:327:SER:OG	8:Ey:211:MET:HE1	2.13	0.48
8:Ey:103:THR:H	8:Ez:253:LEU:HD21	1.77	0.48
8:Fa:359:MET:SD	8:Fa:369:THR:HG22	2.53	0.48
8:Fa:401:LYS:HD2	8:Fa:401:LYS:N	2.28	0.48
9:Fb:414:PHE:HB3	9:Fb:418:GLU:OE1	2.13	0.48
9:Fb:596:PHE:CE2	9:Fb:656:VAL:HG21	2.48	0.48
9:Fb:655:ILE:HG22	9:Fb:659:PHE:CE1	2.49	0.48
9:Fb:944:ILE:HD11	9:Ff:781:ILE:HD12	1.94	0.48
9:Fc:905:TRP:HB3	9:Fc:909:TYR:CE2	2.48	0.48
9:Fd:693:GLY:O	9:Fd:697:ILE:HG23	2.14	0.48
9:Fe:713:VAL:HG11	9:Ff:596:PHE:HZ	1.78	0.48
9:Ff:743:VAL:HG22	9:Ff:747:PHE:CZ	2.49	0.48
1:Aa:12:HIS:CE1	3:Ac:9:TYR:HB2	2.48	0.48
3:Ao:4:ILE:O	6:Bd:132:GLN:HB2	2.14	0.48
5:Aq:209:ILE:HA	5:Aq:241:TYR:OH	2.14	0.48
6:Ax:121:PRO:HB2	7:Eg:801:MET:HE1	1.96	0.48
1:Ay:16:ALA:HA	4:Bh:3:GLY:HA3	1.94	0.48
5:Bi:212:ILE:HG21	5:Bi:215:VAL:HG23	1.96	0.48
6:Bp:113:LYS:HZ2	6:Bp:114:ASP:HA	1.79	0.48

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
6:Bp:190:PRO:O	6:Bp:231:ARG:HG2	2.12	0.48
6:Bv:115:MET:HA	6:Bv:118:ASN:ND2	2.28	0.48
6:Cz:172:SER:HB2	6:Cz:248:VAL:HB	1.95	0.48
6:Dx:106:VAL:HA	6:Dx:109:LYS:HZ1	1.78	0.48
6:Dx:126:GLN:O	6:Dx:130:LEU:HG	2.14	0.48
2:Dz:4:MET:HG2	5:Ec:250:LEU:HD12	1.95	0.48
7:Er:801:MET:C	7:Er:801:MET:SD	2.96	0.48
7:Es:770:SER:O	7:Es:774:LEU:HD22	2.13	0.48
8:Ew:271:SER:H	8:Ew:274:GLN:HB2	1.78	0.48
8:Ex:75:GLN:O	8:Ex:78:GLN:HG3	2.14	0.48
8:Ex:252:LEU:HD12	8:Ex:252:LEU:O	2.13	0.48
8:Ey:399:TRP:O	8:Ey:403:ILE:HG12	2.12	0.48
9:Fc:643:PHE:O	9:Fc:647:ILE:HG13	2.13	0.48
9:Fd:65:ASN:HA	9:Fd:68:VAL:HG12	1.95	0.48
9:Fd:74:ILE:O	9:Fd:77:MET:HE3	2.12	0.48
9:Fd:256:PRO:HG2	9:Fd:259:PHE:CE2	2.49	0.48
9:Fd:685:PRO:HD2	9:Fd:827:TYR:CE2	2.49	0.48
9:Fd:727:LEU:HA	9:Fd:730:MET:SD	2.53	0.48
9:Ff:182:LEU:HD23	9:Ff:571:VAL:HG21	1.96	0.48
9:Ff:238:ASN:O	9:Ff:242:LYS:HD3	2.13	0.48
9:Ff:241:LYS:O	9:Ff:241:LYS:HD3	2.13	0.48
9:Ff:668:LYS:O	9:Ff:672:ILE:HG12	2.13	0.48
9:Ff:811:ARG:NH2	9:Ff:937:PRO:HB2	2.28	0.48
6:Af:153:ALA:HB1	6:Af:251:ARG:HG2	1.96	0.48
5:Ak:253:ARG:O	5:Ak:254:ILE:HD13	2.12	0.48
6:Al:115:MET:SD	7:Ev:802:LEU:HD13	2.52	0.48
6:Ax:111:ALA:HB3	7:Ef:797:ARG:HD3	1.95	0.48
6:Ax:189:TRP:CG	6:Ax:230:LEU:HD12	2.49	0.48
1:Cc:1:CYS:HA	2:Cd:9:ARG:CZ	2.43	0.48
1:Cc:1:CYS:N	3:Ce:16:ASP:HB2	2.28	0.48
6:Dl:126:GLN:O	6:Dl:129:LYS:HG3	2.13	0.48
6:Dl:179:LEU:O	6:Dl:212:THR:HA	2.14	0.48
6:Dl:183:ASP:HB3	6:Dl:189:TRP:CE3	2.49	0.48
1:Dm:1:CYS:SG	4:Dp:2:PHE:HB3	2.53	0.48
5:Dq:253:ARG:HD2	5:Dq:255:LEU:HG	1.95	0.48
6:Dx:196:LEU:HD22	6:Dx:199:PRO:HA	1.96	0.48
5:Ec:213:GLN:HE21	5:Ec:221:TRP:HB3	1.77	0.48
7:Eg:809:VAL:HG22	7:Eg:813:LYS:NZ	2.28	0.48
8:Ew:277:LEU:HA	8:Ew:280:ILE:HG22	1.94	0.48
8:Ew:297:ALA:O	8:Ew:301:LEU:HG	2.13	0.48
8:Ey:42:TYR:HA	8:Ey:45:ASN:ND2	2.29	0.48

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
8:Ey:93:ALA:HB3	8:Ey:172:ASP:HB2	1.96	0.48
8:Ez:419:LEU:HA	8:Ez:422:ILE:HG12	1.96	0.48
9:Fb:468:PHE:CE2	9:Fb:823:ILE:HD13	2.49	0.48
9:Fe:90:GLU:HG3	9:Fe:92:GLN:H	1.78	0.48
9:Fe:113:LEU:HD12	9:Fe:124:MET:HB2	1.94	0.48
9:Fe:670:ILE:HA	9:Fe:673:VAL:HG22	1.96	0.48
7:Fg:16:ARG:O	7:Fg:20:ILE:HG12	2.13	0.48
7:Fj:35:GLY:O	7:Fj:39:ILE:HG13	2.13	0.48
6:Af:112:PHE:O	6:Af:115:MET:HB2	2.14	0.48
6:Ax:134:TYR:HA	6:Bd:120:TYR:HE1	1.78	0.48
6:Bd:145:PRO:HG3	6:Bj:131:LYS:HZ3	1.78	0.48
6:Ch:115:MET:HE2	7:El:802:LEU:CD2	2.43	0.48
6:Ct:115:MET:HE1	7:En:801:MET:CB	2.43	0.48
5:Cy:212:ILE:HD11	5:Cy:262:ILE:HG22	1.96	0.48
6:Cz:151:PRO:HA	6:Cz:248:VAL:HG13	1.95	0.48
6:Df:130:LEU:HA	6:Df:133:ILE:HD12	1.96	0.48
8:Ew:35:ASN:CG	8:Fa:36:THR:HB	2.39	0.48
8:Ey:174:VAL:HG21	8:Ez:246:ASN:HD22	1.78	0.48
8:Ez:334:ARG:HG3	8:Fa:282:TYR:HE2	1.78	0.48
9:Fb:467:TYR:O	9:Fb:471:LEU:HD13	2.14	0.48
9:Fb:665:GLN:HA	9:Fb:668:LYS:HE3	1.95	0.48
9:Fd:332:ARG:O	9:Fd:336:ILE:HG13	2.13	0.48
9:Fd:469:PHE:O	9:Fd:469:PHE:HD1	1.97	0.48
9:Fd:777:VAL:O	9:Fd:780:PRO:HD2	2.14	0.48
9:Fe:185:LEU:O	9:Fe:189:ILE:HG12	2.13	0.48
9:Fe:825:LEU:HA	9:Fe:828:VAL:HG22	1.96	0.48
9:Fe:909:TYR:HA	9:Fe:912:PHE:CE2	2.49	0.48
9:Ff:804:ILE:HA	9:Ff:807:ASN:HB3	1.94	0.48
6:Ar:109:LYS:HD2	6:Ar:109:LYS:C	2.39	0.48
3:Bg:13:LYS:HD3	3:Bg:13:LYS:HA	1.65	0.48
6:Bj:230:LEU:HB3	6:Bj:233:LEU:HD13	1.94	0.48
1:Bk:6:LEU:O	1:Bk:10:GLU:HG3	2.13	0.48
5:Bo:248:ASP:HB3	5:Bo:253:ARG:HB3	1.96	0.48
6:Bp:173:GLN:NE2	7:Ek:815:VAL:HG22	2.26	0.48
6:Cb:115:MET:HG2	6:Cb:119:LEU:HD12	1.95	0.48
6:Ch:176:VAL:HG23	6:Ch:214:MET:HB2	1.95	0.48
6:Ct:171:LEU:HD11	6:Ct:217:ALA:N	2.28	0.48
7:Ef:768:ALA:O	7:Ef:772:GLU:HG2	2.13	0.48
7:El:819:VAL:C	7:El:820:TYR:HD1	2.21	0.48
7:Eq:792:GLN:HA	7:Eq:795:GLN:OE1	2.14	0.48
8:Ew:75:GLN:OE1	8:Ex:401:LYS:HE2	2.14	0.48

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
8:Ew:104:ASP:HA	8:Ew:109:ASN:ND2	2.28	0.48
8:Ew:352:SER:HA	8:Ew:355:LEU:HG	1.96	0.48
8:Ew:414:GLU:O	8:Ew:417:ILE:HG13	2.13	0.48
8:Ey:105:LYS:NZ	8:Ez:367:ASN:HA	2.29	0.48
8:Ez:452:GLN:HG2	8:Fa:336:TYR:CE2	2.49	0.48
8:Fa:351:LEU:HD12	8:Fa:355:LEU:HD21	1.94	0.48
9:Fb:71:LEU:HD23	9:Fb:74:ILE:HD11	1.95	0.48
9:Fd:82:VAL:HA	9:Fd:85:MET:HG3	1.95	0.48
9:Fd:106:ARG:HH11	9:Fd:789:PRO:HA	1.79	0.48
9:Fd:128:PHE:O	9:Fd:132:ILE:HG12	2.14	0.48
9:Fd:151:ASN:HB3	9:Fd:152:ARG:NH1	2.29	0.48
9:Fd:321:THR:HG23	9:Fd:324:GLN:H	1.77	0.48
9:Fd:354:ASP:OD2	9:Fd:356:ALA:HB3	2.14	0.48
9:Fd:812:PRO:HA	9:Fd:815:MET:HG2	1.94	0.48
9:Fe:596:PHE:CE2	9:Fe:652:ILE:HG22	2.49	0.48
9:Fe:728:ILE:HD12	9:Ff:730:MET:HB3	1.95	0.48
9:Fe:755:VAL:HG22	9:Fe:759:MET:SD	2.54	0.48
9:Fe:911:PHE:CD1	9:Fe:911:PHE:C	2.92	0.48
9:Ff:56:ILE:C	9:Ff:60:MET:HE3	2.39	0.48
9:Ff:732:MET:O	9:Ff:736:MET:HB3	2.13	0.48
5:Ae:264:PHE:HB3	5:Ae:269:SER:HB3	1.95	0.48
5:Bc:230:LEU:HA	6:Bd:165:THR:HG21	1.96	0.48
5:Bc:234:GLU:OE2	6:Bj:147:THR:HG21	2.14	0.48
6:Bd:115:MET:HA	6:Bd:118:ASN:OD1	2.14	0.48
6:Bv:168:VAL:HG23	6:Bv:242:ILE:HD13	1.95	0.48
6:Cb:167:PRO:HD2	6:Cb:238:MET:O	2.14	0.48
6:Cb:205:GLN:HB2	6:Cb:214:MET:SD	2.54	0.48
5:Cg:232:VAL:HG23	5:Cg:236:SER:HB2	1.96	0.48
6:Cz:250:TYR:C	6:Cz:251:ARG:HG2	2.38	0.48
6:Df:144:THR:HG23	6:Df:148:PRO:HG3	1.94	0.48
6:Dr:150:LYS:N	6:Dr:247:ALA:HA	2.29	0.48
6:Dr:205:GLN:HB2	6:Dr:214:MET:SD	2.54	0.48
6:Dr:219:LYS:NZ	6:Dr:221:TYR:H	2.05	0.48
7:Eh:806:THR:HA	7:Eh:809:VAL:HG12	1.96	0.48
7:El:807:GLN:HG3	7:El:808:LEU:HD23	1.95	0.48
8:Ex:342:VAL:HG11	8:Ex:442:ILE:HD13	1.95	0.48
8:Fa:141:LEU:HB3	8:Fa:213:ASN:HB3	1.94	0.48
9:Fb:24:LEU:HD23	7:Fi:40:ARG:NH1	2.29	0.48
9:Fb:617:ILE:HG22	9:Fb:618:LEU:H	1.79	0.48
9:Fb:648:PHE:CZ	9:Ff:716:LEU:HD11	2.49	0.48
9:Fb:804:ILE:O	9:Fb:808:VAL:HG22	2.14	0.48

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
9:Fc:338:GLN:NE2	9:Fc:425:ASP:HB2	2.29	0.48
9:Fc:346:VAL:HG13	9:Fc:350:MET:HE3	1.96	0.48
9:Fc:462:ILE:HA	9:Fc:761:PHE:HZ	1.78	0.48
9:Fc:548:PRO:HD3	9:Fc:568:TYR:CD2	2.49	0.48
9:Fc:721:GLY:HA2	9:Fc:724:ILE:HG12	1.95	0.48
9:Fc:842:SER:HA	9:Fc:845:GLN:HE21	1.79	0.48
9:Fd:628:MET:HE2	9:Fd:628:MET:N	2.29	0.48
9:Fd:732:MET:HA	9:Fd:735:LEU:HD23	1.96	0.48
9:Fd:762:THR:CG2	9:Fd:766:PHE:HE2	2.25	0.48
9:Ff:528:ILE:HG22	9:Ff:538:PRO:HD3	1.95	0.48
9:Ff:936:LEU:O	9:Ff:940:VAL:HG13	2.14	0.48
6:Al:158:VAL:O	6:Al:257:GLN:HG3	2.13	0.48
6:Al:189:TRP:CH2	6:Al:233:LEU:HD21	2.48	0.48
6:Ax:221:TYR:HA	6:Ax:243:PRO:HG2	1.96	0.48
6:Cb:132:GLN:NE2	6:Cb:135:GLU:HG3	2.28	0.48
3:Ck:2:VAL:HG21	6:Cz:129:LYS:HD2	1.95	0.48
5:Cm:253:ARG:NH1	5:Cm:253:ARG:HB2	2.29	0.48
6:Cn:151:PRO:HA	6:Cn:248:VAL:HG13	1.96	0.48
6:Ct:111:ALA:HB3	7:En:797:ARG:NH2	2.28	0.48
6:Ct:149:PRO:HA	6:Ct:246:LYS:HD2	1.95	0.48
6:Cz:109:LYS:N	6:Cz:109:LYS:HD2	2.29	0.48
6:Cz:125:GLU:O	6:Cz:128:VAL:HG22	2.13	0.48
8:Ew:78:GLN:HE21	8:Ex:383:LEU:CG	2.27	0.48
8:Ew:278:ASN:HA	8:Ew:281:ARG:HD2	1.96	0.48
8:Ew:421:GLU:HG3	8:Fa:423:ASN:ND2	2.28	0.48
8:Ex:295:LEU:O	8:Ex:299:SER:CB	2.62	0.48
8:Ez:408:PRO:HA	8:Ez:411:VAL:HG22	1.93	0.48
9:Fb:468:PHE:O	9:Fb:472:VAL:HG23	2.13	0.48
9:Fb:643:PHE:CD2	9:Fc:605:MET:HE1	2.49	0.48
9:Fb:936:LEU:HA	9:Fb:939:LYS:HD3	1.94	0.48
9:Fc:353:ASN:O	9:Fc:375:LYS:HE2	2.14	0.48
9:Fc:378:PHE:HB3	9:Fc:414:PHE:CD2	2.49	0.48
9:Fc:776:MET:HE2	9:Fc:776:MET:C	2.39	0.48
9:Fd:563:LEU:HD12	9:Fd:563:LEU:O	2.13	0.48
9:Fd:589:THR:HG23	9:Fd:590:PHE:HD2	1.78	0.48
9:Fd:762:THR:HG23	9:Fd:766:PHE:CE2	2.49	0.48
9:Fe:71:LEU:O	9:Fe:75:ILE:HG12	2.14	0.48
9:Fe:266:PHE:H	9:Fe:276:ASN:ND2	2.11	0.48
7:Fg:36:PHE:CD1	7:Fg:36:PHE:C	2.92	0.48
6:Ar:171:LEU:HD13	6:Ar:202:PHE:CZ	2.49	0.47
5:Aw:210:TYR:CD2	5:Aw:238:ILE:HD11	2.49	0.47

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
6:Bd:117:ARG:NH1	6:Bd:118:ASN:HA	2.28	0.47
5:Bi:217:PRO:HG2	6:Bp:143:ALA:HB1	1.96	0.47
3:Bm:2:VAL:HG21	6:Cb:129:LYS:HG2	1.96	0.47
6:Bp:126:GLN:HB3	6:Bv:112:PHE:CE1	2.49	0.47
6:Bp:194:TYR:CB	6:Bp:228:VAL:HG12	2.42	0.47
3:By:2:VAL:HG21	6:Cn:129:LYS:HG2	1.96	0.47
2:Cd:4:MET:HB2	4:Cf:2:PHE:CZ	2.49	0.47
6:Dl:126:GLN:HA	6:Dl:129:LYS:HE3	1.95	0.47
6:Dr:195:ASP:OD1	7:Eu:820:TYR:HA	2.14	0.47
1:Ds:3:ASP:HB3	3:Du:14:TYR:HD2	1.79	0.47
6:Ed:122:LEU:HB3	6:Ed:126:GLN:HB3	1.94	0.47
7:En:798:THR:O	7:En:802:LEU:HG	2.13	0.47
7:Er:793:GLU:HA	7:Er:796:GLN:NE2	2.29	0.47
8:Ew:389:SER:HB3	8:Ew:394:GLN:OE1	2.14	0.47
8:Ey:91:VAL:HG11	8:Ey:116:ALA:HB1	1.96	0.47
8:Ey:430:ARG:HH21	8:Ez:425:GLN:HA	1.79	0.47
8:Ez:179:PHE:HD2	8:Fa:240:ILE:HD13	1.77	0.47
8:Ez:389:SER:HB3	8:Ez:394:GLN:OE1	2.14	0.47
8:Ez:419:LEU:HD13	8:Fa:419:LEU:HB3	1.95	0.47
8:Fa:340:THR:O	8:Fa:344:VAL:HG23	2.14	0.47
8:Fa:349:TYR:O	8:Fa:353:LYS:HG3	2.14	0.47
9:Fb:239:PHE:HE2	9:Fb:340:TYR:HB2	1.79	0.47
9:Fb:348:GLN:HA	9:Fb:351:VAL:HG12	1.96	0.47
9:Fb:911:PHE:O	9:Fb:915:ILE:HG22	2.13	0.47
9:Fc:694:THR:HA	9:Fc:697:ILE:HG12	1.96	0.47
9:Fd:140:ASP:HA	9:Fd:143:TRP:HE3	1.78	0.47
9:Fd:278:ILE:CD1	9:Fd:516:LEU:HD11	2.44	0.47
9:Fd:663:PRO:O	9:Fd:667:MET:HE3	2.14	0.47
9:Fd:924:ILE:HA	9:Fd:927:GLN:HB2	1.96	0.47
9:Fe:641:TYR:O	9:Fe:645:LEU:HG	2.13	0.47
9:Fe:766:PHE:HZ	9:Ff:929:ALA:HA	1.79	0.47
9:Ff:110:GLY:HA3	9:Ff:787:THR:HG23	1.96	0.47
9:Ff:490:GLY:O	9:Ff:493:LYS:HG2	2.13	0.47
9:Ff:534:ILE:HA	9:Ff:567:VAL:HG22	1.95	0.47
7:Fi:30:ILE:HA	7:Fi:33:VAL:HG22	1.96	0.47
6:Al:170:ARG:HG2	6:Al:245:GLN:OE1	2.14	0.47
6:Ar:191:ILE:HB	6:Ar:206:TRP:CZ2	2.48	0.47
6:Ax:182:LEU:HD12	6:Ax:183:ASP:N	2.29	0.47
6:Ax:190:PRO:HA	6:Ax:211:ASN:HB3	1.96	0.47
6:Cb:115:MET:CG	6:Cb:118:ASN:HB2	2.43	0.47
6:Cn:129:LYS:O	6:Cn:133:ILE:HG12	2.14	0.47

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
6:Cn:134:TYR:O	6:Cn:138:GLU:HG3	2.13	0.47
2:Dt:4:MET:HE2	2:Dt:4:MET:HA	1.96	0.47
5:Dw:219:ARG:H	6:Ed:148:PRO:HD2	1.79	0.47
6:Dx:167:PRO:HG2	6:Dx:239:LEU:HD12	1.96	0.47
6:Ed:151:PRO:HA	6:Ed:248:VAL:HG13	1.97	0.47
6:Ed:202:PHE:HZ	6:Ed:243:PRO:HD3	1.79	0.47
7:Ee:795:GLN:HA	7:Ee:798:THR:OG1	2.14	0.47
7:Ek:809:VAL:HA	7:Ek:812:TRP:HE3	1.78	0.47
8:Ew:280:ILE:HG21	8:Ew:351:LEU:HD22	1.96	0.47
8:Ew:382:ARG:HH11	8:Ew:382:ARG:HG3	1.79	0.47
8:Ey:56:GLN:HB2	8:Ey:59:LYS:HG2	1.97	0.47
8:Ez:82:TYR:HB2	8:Fa:383:LEU:HD21	1.95	0.47
8:Ez:448:LEU:HB2	8:Fa:336:TYR:HE1	1.79	0.47
9:Fb:425:ASP:O	9:Fb:429:ILE:HG23	2.14	0.47
9:Fb:516:LEU:HA	9:Fb:519:TRP:CZ3	2.50	0.47
9:Fb:609:ASP:HB3	9:Fb:626:ARG:HH22	1.79	0.47
9:Fb:670:ILE:HG21	9:Fb:742:MET:HE1	1.96	0.47
9:Fb:692:MET:O	9:Fb:696:TYR:HD2	1.97	0.47
9:Fb:732:MET:SD	9:Fb:733:PRO:HD3	2.54	0.47
9:Fc:419:PHE:O	9:Fc:423:ILE:HG12	2.15	0.47
9:Fd:104:PRO:HG3	7:Fk:20:ILE:HD11	1.96	0.47
9:Fd:736:MET:O	9:Fd:736:MET:HE3	2.13	0.47
9:Fd:804:ILE:O	9:Fd:808:VAL:HG22	2.13	0.47
9:Fe:516:LEU:HA	9:Fe:519:TRP:CZ3	2.48	0.47
9:Fe:640:VAL:HG12	9:Fe:644:PHE:HE2	1.79	0.47
9:Fe:658:ALA:O	9:Fe:662:ILE:HG22	2.15	0.47
9:Fe:729:MET:O	9:Fe:732:MET:HG3	2.14	0.47
9:Ff:321:THR:HG23	9:Ff:324:GLN:H	1.79	0.47
9:Ff:761:PHE:HA	9:Ff:933:ILE:CD1	2.44	0.47
6:Al:238:MET:HG3	6:Ar:250:TYR:HB3	1.96	0.47
6:Ar:171:LEU:HD22	6:Ar:202:PHE:CE2	2.50	0.47
6:Ar:221:TYR:CZ	6:Ax:139:TYR:HA	2.49	0.47
6:Ar:236:PRO:HD2	6:Ax:251:ARG:HE	1.78	0.47
1:Be:16:ALA:HB1	4:Bn:3:GLY:H	1.78	0.47
6:Bp:145:PRO:HD3	6:Bv:131:LYS:HE2	1.96	0.47
6:Bv:115:MET:HE1	7:Ej:802:LEU:HA	1.96	0.47
6:Cb:196:LEU:HD11	6:Cb:199:PRO:HB3	1.96	0.47
6:Cn:130:LEU:HA	6:Cn:133:ILE:HD11	1.95	0.47
6:Cn:238:MET:H	6:Ct:251:ARG:HH12	1.60	0.47
6:Ct:227:ALA:HA	6:Cz:251:ARG:NH2	2.29	0.47
6:Dl:190:PRO:O	6:Dl:230:LEU:HD22	2.14	0.47

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
5:Dq:207:ARG:HH21	5:Dq:240:GLY:HA3	1.79	0.47
5:Dq:215:VAL:HG11	5:Dq:252:GLY:HA2	1.96	0.47
6:Dx:132:GLN:O	6:Dx:135:GLU:HG2	2.14	0.47
6:Ed:122:LEU:HD23	7:Eu:801:MET:HE3	1.96	0.47
7:Ei:801:MET:SD	7:Ei:801:MET:C	2.98	0.47
8:Ew:187:TYR:HB3	8:Ew:206:TYR:CE2	2.50	0.47
8:Ew:189:PHE:CD2	8:Ew:295:LEU:HD23	2.49	0.47
8:Ex:457:PHE:HZ	8:Ey:293:PRO:HB3	1.80	0.47
8:Ey:359:MET:HB2	8:Ey:367:ASN:HB2	1.96	0.47
9:Fb:176:GLY:HA3	9:Fb:492:ASP:HB2	1.96	0.47
9:Fc:98:TRP:CD1	9:Fc:103:ILE:HD13	2.49	0.47
9:Fc:594:ILE:HG13	9:Fc:905:TRP:CH2	2.49	0.47
9:Fd:692:MET:O	9:Fd:696:TYR:CB	2.62	0.47
9:Fd:909:TYR:HA	9:Fd:912:PHE:CE2	2.49	0.47
9:Fe:242:LYS:HG3	9:Fe:261:LEU:HD11	1.95	0.47
6:Af:125:GLU:OE2	6:Af:129:LYS:HD3	2.13	0.47
5:Ak:263:LYS:HA	5:Ak:263:LYS:HE2	1.96	0.47
1:Am:2:THR:HG21	5:Aq:255:LEU:HD11	1.96	0.47
2:Bf:2:VAL:HG13	3:Bg:1:ARG:HD2	1.95	0.47
6:Ct:189:TRP:CG	6:Ct:230:LEU:HD12	2.49	0.47
6:Dr:125:GLU:HA	6:Dr:128:VAL:HG22	1.97	0.47
6:Ed:122:LEU:HB3	6:Ed:126:GLN:CB	2.43	0.47
7:Eh:781:GLN:O	7:Eh:785:LEU:HG	2.15	0.47
8:Ew:86:LEU:HD22	8:Ew:115:LEU:HD23	1.95	0.47
8:Ew:233:TYR:HB3	8:Ew:234:LYS:HZ2	1.78	0.47
8:Ew:298:TYR:HE2	8:Ex:221:ALA:HA	1.78	0.47
8:Ex:80:TYR:HB2	8:Ex:374:ASN:OD1	2.14	0.47
8:Ey:219:PRO:HG2	8:Ey:230:PHE:HB3	1.96	0.47
8:Ey:444:LEU:O	8:Ey:448:LEU:HG	2.13	0.47
8:Ez:189:PHE:HE1	8:Ez:293:PRO:O	1.98	0.47
9:Fb:661:MET:HB3	9:Fb:665:GLN:HG3	1.95	0.47
9:Fc:420:LEU:HD12	9:Fc:421:GLY:N	2.30	0.47
9:Fd:696:TYR:CE1	9:Fd:745:ILE:HB	2.37	0.47
9:Fd:704:TRP:HE3	9:Fd:707:LEU:HD12	1.79	0.47
9:Fe:354:ASP:OD1	9:Fe:357:PHE:HD2	1.98	0.47
9:Fe:604:TYR:HD1	9:Fe:605:MET:N	2.12	0.47
9:Fe:664:LEU:O	9:Fe:668:LYS:HB3	2.14	0.47
9:Fe:672:ILE:HA	9:Fe:675:VAL:HG12	1.96	0.47
9:Ff:624:LEU:HA	9:Ff:627:MET:CE	2.44	0.47
6:Af:205:GLN:HE22	6:Af:216:GLN:HE22	1.62	0.47
6:Al:149:PRO:CB	6:Al:248:VAL:HG13	2.44	0.47

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
5:Aq:210:TYR:HA	5:Aq:224:GLY:HA2	1.95	0.47
5:Bi:222:LEU:HD11	5:Bi:262:ILE:HG21	1.97	0.47
6:Bp:160:LEU:HD12	6:Bp:160:LEU:O	2.14	0.47
5:Bu:262:ILE:HD12	5:Bu:263:LYS:H	1.79	0.47
6:Bv:106:VAL:HA	6:Bv:109:LYS:HZ3	1.80	0.47
6:Ct:115:MET:HE2	7:En:798:THR:HA	1.95	0.47
6:Cz:209:THR:HG22	7:Er:824:THR:HG21	1.96	0.47
6:Dx:106:VAL:O	6:Dx:110:LYS:HG2	2.14	0.47
7:Ef:772:GLU:O	7:Ef:775:GLN:HG2	2.14	0.47
7:Eh:780:LYS:HD2	7:Eh:784:GLN:HG2	1.95	0.47
7:Eu:776:ALA:O	7:Eu:779:GLN:HG2	2.15	0.47
7:Eu:784:GLN:O	7:Eu:788:GLN:HG2	2.14	0.47
8:Ex:40:VAL:O	8:Ex:43:LEU:HG	2.14	0.47
8:Ey:334:ARG:HD2	8:Ez:282:TYR:CE1	2.49	0.47
8:Ez:268:THR:CG2	8:Ez:270:LYS:HZ1	2.28	0.47
9:Fb:331:SER:O	9:Fb:334:ILE:HG22	2.14	0.47
9:Fb:332:ARG:O	9:Fb:336:ILE:HG12	2.15	0.47
9:Fb:655:ILE:HG22	9:Fb:659:PHE:HE1	1.79	0.47
9:Fc:188:GLN:HA	9:Fc:191:MET:HG2	1.95	0.47
9:Fc:346:VAL:O	9:Fc:350:MET:HG2	2.14	0.47
9:Fc:399:TRP:CE3	9:Fc:416:GLY:HA3	2.50	0.47
9:Fc:465:GLY:H	9:Fc:761:PHE:HE2	1.61	0.47
9:Ff:391:GLU:O	9:Ff:394:GLN:HG2	2.14	0.47
7:Fi:35:GLY:O	7:Fi:39:ILE:HG13	2.14	0.47
6:Ax:252:VAL:HG12	6:Ax:254:LEU:HD23	1.95	0.47
3:Bm:4:ILE:HA	6:Cb:132:GLN:HG2	1.97	0.47
6:Bp:111:ALA:HA	6:Bp:114:ASP:OD1	2.15	0.47
6:Bp:226:LEU:HD12	6:Bp:227:ALA:N	2.30	0.47
5:Bu:219:ARG:H	6:Cb:148:PRO:HD2	1.79	0.47
6:Bv:113:LYS:O	6:Bv:116:THR:HG22	2.15	0.47
6:Cb:107:ILE:HG22	7:Ek:797:ARG:HD3	1.96	0.47
5:Cm:210:TYR:HD2	5:Cm:224:GLY:HA3	1.79	0.47
1:Co:3:ASP:O	1:Co:6:LEU:HD12	2.15	0.47
6:Cz:107:ILE:HG22	7:Eo:797:ARG:HD3	1.96	0.47
5:Dk:219:ARG:HG2	6:Dr:148:PRO:HD2	1.96	0.47
6:Dl:192:ALA:HB3	6:Dl:229:ARG:HG3	1.97	0.47
6:Dr:225:ASN:HD21	6:Dx:176:VAL:H	1.61	0.47
6:Ed:241:LEU:H	6:Ed:241:LEU:HD12	1.80	0.47
7:Ef:809:VAL:HA	7:Ef:812:TRP:CD1	2.49	0.47
8:Ex:99:MET:HB2	8:Ex:116:ALA:HB1	1.96	0.47
8:Ey:188:SER:HB2	8:Ey:294:LYS:HD2	1.96	0.47

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
8:Ey:343:GLY:O	8:Ey:347:LEU:HG	2.15	0.47
8:Ey:430:ARG:HH22	8:Ez:428:LEU:HD12	1.78	0.47
8:Ez:35:ASN:O	8:Ez:39:LEU:HD23	2.14	0.47
8:Ez:284:SER:HB2	8:Ez:344:VAL:HG12	1.95	0.47
8:Ez:426:MET:C	8:Ez:426:MET:HE2	2.38	0.47
9:Fc:33:LEU:HA	9:Fc:36:VAL:HG12	1.97	0.47
9:Fc:222:MET:HE1	9:Fc:519:TRP:HD1	1.80	0.47
9:Fc:693:GLY:HA2	9:Fc:696:TYR:CE2	2.49	0.47
9:Fc:825:LEU:HD12	9:Fc:929:ALA:CB	2.39	0.47
9:Fe:378:PHE:CE2	9:Fe:565:SER:HA	2.49	0.47
9:Fe:628:MET:HA	9:Fe:628:MET:HE2	1.96	0.47
9:Fe:837:PHE:CE1	9:Fe:841:ILE:HG21	2.50	0.47
9:Ff:809:PHE:HD1	9:Ff:810:LEU:HD23	1.80	0.47
9:Ff:924:ILE:O	9:Ff:927:GLN:HG2	2.15	0.47
6:Af:238:MET:N	6:Al:251:ARG:HH12	2.13	0.47
1:Ag:9:LEU:HD12	1:Ag:10:GLU:N	2.30	0.47
6:Ar:115:MET:O	6:Ar:119:LEU:HD12	2.13	0.47
6:Ax:109:LYS:N	6:Ax:109:LYS:HD3	2.30	0.47
6:Ax:191:ILE:HD12	6:Ax:206:TRP:NE1	2.30	0.47
6:Ax:208:LYS:HE2	7:Ei:823:GLY:HA2	1.97	0.47
6:Bp:186:GLY:N	6:Bp:255:ARG:HH12	2.12	0.47
1:Cc:9:LEU:HD13	2:Cd:2:VAL:HG23	1.96	0.47
6:Ch:235:THR:HG23	6:Cn:251:ARG:HD2	1.97	0.47
1:Ci:3:ASP:OD2	3:Ck:14:TYR:HB3	2.14	0.47
5:Cm:238:ILE:O	5:Cm:238:ILE:HG13	2.13	0.47
5:Cs:230:LEU:CD1	5:Cs:231:THR:H	2.24	0.47
5:Cy:219:ARG:NH2	6:Df:150:LYS:HD3	2.29	0.47
6:Cz:189:TRP:CG	6:Cz:230:LEU:HD12	2.50	0.47
1:Da:16:ALA:HA	4:Dj:3:GLY:HA3	1.96	0.47
5:De:255:LEU:HD13	5:De:261:VAL:HG13	1.95	0.47
6:Dl:144:THR:CG2	6:Dl:148:PRO:HG3	2.44	0.47
6:Dr:106:VAL:HA	6:Dr:109:LYS:NZ	2.30	0.47
6:Dr:150:LYS:HG2	6:Dr:152:THR:HG23	1.95	0.47
6:Dr:208:LYS:HB3	7:Eu:824:THR:HG23	1.96	0.47
7:Ei:787:GLU:HA	7:Ei:790:TYR:HD2	1.80	0.47
7:Em:782:ASN:HB3	9:Fc:241:LYS:HE2	1.97	0.47
7:Eq:781:GLN:OE1	9:Fb:337:GLN:HB3	2.15	0.47
7:Er:808:LEU:HB3	7:Er:812:TRP:CZ2	2.49	0.47
7:Et:779:GLN:HA	7:Et:782:ASN:ND2	2.29	0.47
8:Ew:376:PHE:HE2	8:Fa:85:PHE:CE2	2.33	0.47
8:Ew:418:LEU:HD12	8:Ew:418:LEU:C	2.40	0.47

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
8:Ex:42:TYR:HA	8:Ex:45:ASN:HD21	1.79	0.47
8:Ey:54:ILE:HG23	8:Ez:45:ASN:HB2	1.96	0.47
8:Ey:129:SER:H	8:Ey:137:THR:HG21	1.79	0.47
8:Ey:191:MET:HE1	8:Ey:198:TRP:CG	2.49	0.47
8:Ey:269:ALA:HB1	8:Ey:274:GLN:HB2	1.96	0.47
8:Ey:298:TYR:CZ	8:Ez:230:PHE:HE1	2.33	0.47
8:Ez:78:GLN:HB2	8:Fa:383:LEU:CD1	2.44	0.47
8:Fa:408:PRO:HA	8:Fa:411:VAL:HG22	1.96	0.47
9:Fb:264:PRO:HB2	9:Fb:266:PHE:CZ	2.50	0.47
9:Fb:644:PHE:HB3	9:Fb:720:PHE:HE2	1.77	0.47
9:Fb:730:MET:HE2	9:Ff:708:LEU:HD22	1.97	0.47
9:Fc:68:VAL:HA	9:Fc:71:LEU:HG	1.97	0.47
9:Fc:820:ILE:HA	9:Fc:823:ILE:HD12	1.96	0.47
9:Fd:297:ASP:HB2	9:Fd:299:LYS:HG2	1.97	0.47
9:Fd:459:LYS:HE2	9:Fd:463:MET:HG2	1.97	0.47
9:Fd:546:LYS:CE	9:Fd:650:GLU:HG3	2.45	0.47
9:Fd:721:GLY:HA2	9:Fd:724:ILE:HG12	1.95	0.47
9:Fd:797:LYS:HB3	9:Fd:800:PHE:CE2	2.49	0.47
9:Fe:612:CYS:HB2	9:Fe:626:ARG:NE	2.27	0.47
9:Fe:912:PHE:O	9:Fe:916:LEU:HD22	2.14	0.47
9:Ff:802:ILE:O	9:Ff:806:VAL:HG22	2.15	0.47
9:Ff:936:LEU:N	9:Ff:937:PRO:HD2	2.29	0.47
6:Al:115:MET:HE3	7:Ev:801:MET:HE2	1.96	0.47
6:Al:125:GLU:O	6:Al:128:VAL:HG22	2.15	0.47
5:Bi:216:ILE:HG12	5:Bi:221:TRP:CH2	2.44	0.47
5:Bi:251:GLN:HB3	5:Bi:253:ARG:HH21	1.79	0.47
5:Bo:219:ARG:HD3	6:Bv:148:PRO:O	2.14	0.47
5:Dq:207:ARG:NH1	5:Dq:208:ILE:H	2.13	0.47
7:Et:805:ALA:O	7:Et:809:VAL:HG12	2.14	0.47
8:Ew:67:GLN:HA	8:Ew:413:LYS:HZ2	1.80	0.47
8:Ew:254:MET:CE	8:Ew:354:ARG:HB2	2.43	0.47
8:Ex:80:TYR:HB3	8:Ex:359:MET:HE2	1.97	0.47
8:Ey:343:GLY:HA2	8:Ey:346:ASN:ND2	2.29	0.47
8:Ey:374:ASN:O	8:Ey:378:MET:HE2	2.14	0.47
8:Ey:400:ILE:HG13	8:Ey:401:LYS:N	2.29	0.47
8:Fa:123:GLN:HA	8:Fa:135:LYS:HE3	1.96	0.47
8:Fa:177:ALA:O	8:Fa:181:ILE:HG12	2.14	0.47
9:Fb:759:MET:HB3	9:Fb:763:PHE:CZ	2.49	0.47
9:Fc:185:LEU:HD12	9:Fc:189:ILE:HD11	1.97	0.47
9:Fd:644:PHE:HB3	9:Fd:717:ILE:HD11	1.97	0.47
5:Ae:253:ARG:CZ	5:Ae:253:ARG:HB2	2.44	0.47

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
6:Ax:191:ILE:HG13	6:Ax:211:ASN:HA	1.96	0.47
6:Bd:182:LEU:HD11	6:Bd:186:GLY:HA2	1.97	0.47
6:Bd:196:LEU:HD21	6:Bd:202:PHE:HD2	1.80	0.47
6:Bj:254:LEU:H	6:Bj:254:LEU:HD12	1.80	0.47
5:Cm:219:ARG:HD3	6:Ct:246:LYS:HZ3	1.78	0.47
6:Ct:128:VAL:O	6:Ct:132:GLN:HG2	2.15	0.47
1:Cu:1:CYS:HA	4:Cx:2:PHE:CD2	2.50	0.47
5:De:245:LYS:HD2	5:De:245:LYS:N	2.29	0.47
5:Dk:215:VAL:HG22	5:Dk:249:SER:HB3	1.97	0.47
6:Dx:130:LEU:HD12	7:Et:808:LEU:HD12	1.97	0.47
6:Ed:252:VAL:HG13	6:Ed:254:LEU:HD21	1.97	0.47
8:Ey:327:SER:OG	8:Ez:211:MET:HE1	2.15	0.47
8:Fa:381:ARG:HG3	8:Fa:382:ARG:HD3	1.97	0.47
9:Fb:203:ASP:HA	9:Fb:206:LEU:HG	1.96	0.47
9:Fb:693:GLY:O	9:Fb:696:TYR:HB2	2.15	0.47
9:Fc:57:MET:SD	9:Fc:60:MET:HE2	2.55	0.47
9:Fc:825:LEU:HD11	9:Fc:928:LYS:CE	2.45	0.47
9:Fd:829:GLY:HA2	9:Fd:832:ILE:HG12	1.97	0.47
9:Fe:753:ILE:HG13	9:Fe:922:TYR:CZ	2.50	0.47
9:Ff:624:LEU:HA	9:Ff:627:MET:HE1	1.96	0.47
1:Ag:11:TYR:CE2	3:Ai:9:TYR:HB3	2.50	0.47
6:Bj:117:ARG:HH21	7:Ei:797:ARG:HB3	1.79	0.47
1:Cc:16:ALA:HB3	2:Cj:9:ARG:HG3	1.97	0.47
6:Ct:196:LEU:HD21	6:Ct:202:PHE:HB2	1.96	0.47
6:Dx:191:ILE:HD13	6:Dx:213:LEU:HG	1.97	0.47
6:Dx:238:MET:HE3	6:Ed:251:ARG:HB2	1.97	0.47
6:Ed:111:ALA:CB	7:Et:797:ARG:HH21	2.27	0.47
7:Ef:797:ARG:O	7:Ef:801:MET:HG2	2.14	0.47
7:Es:784:GLN:OE1	7:Es:784:GLN:HA	2.14	0.47
7:Eu:804:ALA:O	7:Eu:807:GLN:HG3	2.15	0.47
8:Ew:68:LEU:HD12	8:Ew:69:PHE:H	1.80	0.47
8:Ew:118:THR:HA	8:Ew:121:LYS:HG2	1.96	0.47
8:Ex:101:PHE:HZ	8:Ex:116:ALA:HA	1.79	0.47
8:Ex:316:GLU:O	8:Ex:320:LYS:HG3	2.15	0.47
8:Ey:105:LYS:HZ3	8:Ez:367:ASN:HA	1.79	0.47
8:Ey:407:SER:O	8:Ey:411:VAL:HG13	2.15	0.47
8:Ez:420:ALA:O	8:Ez:423:ASN:HB3	2.14	0.47
8:Fa:208:ASN:O	8:Fa:212:GLN:HG3	2.14	0.47
9:Fb:831:TRP:HE3	9:Ff:37:PHE:HE2	1.63	0.47
9:Fc:716:LEU:HB3	9:Fd:723:PHE:CD2	2.48	0.47
9:Fd:146:ALA:O	9:Fd:150:LEU:HD22	2.15	0.47

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
9:Fd:339:MET:O	9:Fd:343:LEU:HD13	2.15	0.47
9:Fd:427:ASN:HA	9:Fd:430:MET:HE3	1.97	0.47
9:Fd:566:THR:HG23	9:Fd:569:GLY:H	1.81	0.47
9:Fe:132:ILE:HG22	9:Fe:808:VAL:HB	1.96	0.47
9:Fe:725:PHE:CZ	9:Ff:722:ILE:HG23	2.50	0.47
9:Ff:338:GLN:HE22	9:Ff:429:ILE:HB	1.80	0.47
9:Ff:434:LEU:O	9:Ff:437:ILE:HG22	2.15	0.47
7:Fj:14:ASN:HB3	7:Fj:17:THR:HG23	1.96	0.47
7:Fj:20:ILE:HG13	7:Fj:21:ILE:HD13	1.97	0.47
6:Af:117:ARG:HE	7:Ev:797:ARG:CZ	2.29	0.46
6:Af:238:MET:HG2	6:Al:251:ARG:CZ	2.45	0.46
2:An:4:MET:HE1	5:Aq:248:ASP:HA	1.96	0.46
6:Ar:171:LEU:HD21	6:Ar:241:LEU:HD12	1.97	0.46
6:Ar:173:GLN:NE2	6:Ar:220:LEU:HD23	2.30	0.46
6:Ar:181:PHE:HA	6:Ar:254:LEU:HD12	1.97	0.46
5:Aw:254:ILE:HD13	5:Aw:262:ILE:HB	1.97	0.46
6:Bd:219:LYS:HD2	6:Bd:222:ASN:ND2	2.27	0.46
6:Bj:115:MET:HE1	7:Eh:802:LEU:HA	1.96	0.46
1:Cc:16:ALA:HA	4:Cl:3:GLY:HA3	1.97	0.46
6:Cn:233:LEU:HD13	6:Cn:235:THR:O	2.15	0.46
6:Ct:122:LEU:HG	7:Eo:808:LEU:HD11	1.97	0.46
6:Cz:108:ASP:HA	7:Eo:797:ARG:CZ	2.45	0.46
5:De:218:GLY:HA3	6:Dl:148:PRO:HD3	1.96	0.46
5:De:264:PHE:CD1	5:De:269:SER:HB3	2.46	0.46
5:Dq:219:ARG:HD3	6:Dx:148:PRO:O	2.14	0.46
6:Dr:171:LEU:HB2	6:Dr:243:PRO:HA	1.97	0.46
1:Ds:16:ALA:HB3	2:Dz:9:ARG:HG2	1.97	0.46
6:Dx:238:MET:HE3	6:Ed:251:ARG:CB	2.45	0.46
7:Eh:789:LYS:HA	7:Eh:792:GLN:HE22	1.80	0.46
7:El:798:THR:O	7:El:802:LEU:HD23	2.15	0.46
7:Eo:797:ARG:O	7:Eo:801:MET:HG2	2.15	0.46
7:Es:774:LEU:O	7:Es:778:LEU:HG	2.15	0.46
8:Ez:177:ALA:O	8:Ez:181:ILE:HG12	2.15	0.46
8:Fa:426:MET:C	8:Fa:426:MET:HE2	2.40	0.46
9:Fb:500:GLN:O	9:Fb:503:LYS:HB3	2.13	0.46
9:Fb:615:VAL:HG12	9:Fb:622:PHE:O	2.15	0.46
9:Fb:635:TYR:CE2	9:Ff:507:LYS:HG3	2.50	0.46
9:Fc:728:ILE:CD1	9:Fd:730:MET:HB3	2.45	0.46
9:Fd:326:GLN:HB3	9:Fd:329:ARG:HH21	1.79	0.46
9:Fd:905:TRP:CE3	9:Fd:909:TYR:CE1	3.00	0.46
9:Fe:922:TYR:O	9:Fe:926:VAL:HG23	2.16	0.46

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
9:Ff:655:ILE:O	9:Ff:659:PHE:HD1	1.98	0.46
9:Ff:700:SER:HB2	9:Ff:742:MET:HE2	1.96	0.46
5:Ae:253:ARG:HH12	5:Ae:263:LYS:HG2	1.80	0.46
6:Ax:166:PRO:HG3	6:Bd:250:TYR:CD2	2.51	0.46
6:Bj:170:ARG:HD2	6:Bj:247:ALA:HB3	1.97	0.46
6:Bv:170:ARG:O	6:Bv:249:ASP:HB2	2.15	0.46
5:Cg:222:LEU:HD21	5:Cg:254:ILE:HD11	1.97	0.46
6:Cn:229:ARG:HH11	6:Cn:230:LEU:H	1.63	0.46
3:Di:1:ARG:HG2	3:Di:2:VAL:N	2.29	0.46
8:Ew:323:ALA:C	8:Ex:212:GLN:HE22	2.23	0.46
8:Ex:75:GLN:HE21	8:Ex:79:ASN:ND2	2.12	0.46
8:Ex:84:THR:O	8:Ex:88:ALA:HB2	2.15	0.46
9:Fb:588:LEU:HD13	9:Fb:661:MET:HG2	1.97	0.46
9:Fb:630:ASP:HB3	9:Ff:507:LYS:HE2	1.96	0.46
9:Fb:924:ILE:CD1	9:Ff:763:PHE:HE1	2.27	0.46
9:Fd:65:ASN:HA	9:Fd:68:VAL:CG1	2.44	0.46
9:Fe:213:PRO:HB2	9:Fe:222:MET:HG2	1.95	0.46
9:Fe:769:LEU:O	9:Fe:772:VAL:HG12	2.16	0.46
9:Ff:191:MET:SD	9:Ff:263:MET:HG2	2.55	0.46
9:Ff:339:MET:SD	9:Ff:339:MET:N	2.88	0.46
9:Ff:726:ALA:O	9:Ff:730:MET:HG3	2.16	0.46
9:Ff:928:LYS:HA	9:Ff:931:THR:HG23	1.97	0.46
6:Af:109:LYS:C	6:Af:113:LYS:HZ3	2.24	0.46
1:Ag:16:ALA:HB1	4:Ap:3:GLY:H	1.81	0.46
6:Ar:204:ILE:HG12	6:Ar:215:ILE:HG12	1.97	0.46
6:Bd:109:LYS:HD2	6:Bd:109:LYS:N	2.31	0.46
6:Bp:227:ALA:HA	6:Bp:238:MET:HB2	1.96	0.46
6:Cb:176:VAL:HG23	6:Cb:214:MET:HE2	1.98	0.46
5:Cg:219:ARG:NH2	6:Cn:150:LYS:HD2	2.31	0.46
6:Ch:115:MET:HE3	7:El:798:THR:HA	1.98	0.46
6:Cn:141:LYS:HE3	7:Eo:812:TRP:CZ2	2.49	0.46
6:Cn:255:ARG:HD2	6:Cn:255:ARG:HA	1.67	0.46
5:Dk:246:LEU:HD21	5:Dk:248:ASP:HB2	1.96	0.46
5:Dq:253:ARG:HB2	5:Dq:263:LYS:HZ1	1.80	0.46
5:Dw:253:ARG:C	5:Dw:254:ILE:HD13	2.40	0.46
6:Dx:122:LEU:HD11	6:Dx:130:LEU:HD11	1.98	0.46
6:Dx:203:ASN:HD21	7:Eu:817:THR:HG23	1.80	0.46
7:Ek:794:ILE:O	7:Ek:798:THR:HG23	2.15	0.46
7:Eu:778:LEU:O	7:Eu:781:GLN:HB2	2.14	0.46
8:Ew:64:PRO:HG3	8:Ex:48:LYS:HG2	1.96	0.46
8:Ew:301:LEU:HB2	8:Ew:326:LEU:HD21	1.96	0.46

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
8:Ex:40:VAL:O	8:Ex:44:THR:HG23	2.15	0.46
8:Ex:342:VAL:HG21	8:Ex:442:ILE:HD13	1.98	0.46
8:Ex:382:ARG:HG2	8:Ex:399:TRP:CG	2.50	0.46
8:Ex:435:ARG:O	8:Ex:439:THR:HG23	2.15	0.46
8:Ey:346:ASN:HA	8:Ey:435:ARG:NH2	2.30	0.46
8:Ez:426:MET:HA	8:Ez:429:ASP:OD2	2.15	0.46
8:Fa:56:GLN:HB3	8:Fa:59:LYS:HG2	1.96	0.46
9:Fb:452:PHE:HD1	9:Fb:452:PHE:H	1.62	0.46
9:Fb:464:ALA:HB3	9:Fb:761:PHE:CE2	2.50	0.46
9:Fb:741:THR:HG23	9:Fb:916:LEU:HD11	1.96	0.46
9:Fc:273:TYR:O	9:Fc:276:ASN:HB2	2.16	0.46
9:Fc:520:PHE:CE2	9:Fc:526:LYS:HB2	2.44	0.46
9:Fc:827:TYR:HA	9:Fc:830:VAL:HG12	1.97	0.46
9:Fd:45:VAL:HG11	9:Fd:129:MET:HG2	1.97	0.46
9:Fd:748:VAL:O	9:Fd:753:ILE:HG12	2.16	0.46
5:Aq:233:ARG:HB2	6:Ax:147:THR:HG21	1.95	0.46
6:Bd:144:THR:HG21	6:Bd:148:PRO:HG3	1.96	0.46
5:Bi:264:PHE:HB3	5:Bi:269:SER:HB3	1.98	0.46
6:Bj:117:ARG:NH2	7: Ei:797:ARG:HB3	2.31	0.46
6:Bj:147:THR:OG1	6:Bj:246:LYS:HE2	2.15	0.46
6:Bp:225:ASN:HB2	6:Bv:250:TYR:HH	1.80	0.46
6:Cn:189:TRP:CE2	6:Cn:260:GLY:HA2	2.50	0.46
2:Cp:4:MET:SD	5:Cs:248:ASP:HA	2.56	0.46
5:Cs:253:ARG:C	5:Cs:254:ILE:HD13	2.41	0.46
6:Ct:121:PRO:CB	7:Eo:801:MET:HE1	2.39	0.46
6:Df:122:LEU:CD2	7:Eq:808:LEU:HD11	2.45	0.46
6:Dr:115:MET:SD	7:Er:801:MET:SD	3.13	0.46
6:Dx:137:SER:HG	6:Ed:120:TYR:HE1	1.61	0.46
7:Em:781:GLN:O	7:Em:785:LEU:HD23	2.16	0.46
7:Em:808:LEU:O	7:Em:811:ASP:N	2.49	0.46
7:Eq:774:LEU:HD11	9:Fb:432:PRO:HG2	1.96	0.46
7:Eq:822:GLU:OE1	7:Eq:822:GLU:HA	2.15	0.46
8:Ex:293:PRO:HD3	8:Ex:329:TYR:OH	2.15	0.46
8:Ey:67:GLN:HA	8:Ey:413:LYS:NZ	2.31	0.46
8:Ez:175:SER:HB3	8:Fa:244:ASN:HD22	1.80	0.46
8:Ez:341:SER:HA	8:Ez:344:VAL:HG22	1.97	0.46
9:Fb:609:ASP:HA	9:Fb:626:ARG:NH1	2.30	0.46
9:Fb:908:VAL:HA	9:Fb:911:PHE:CE2	2.50	0.46
9:Fc:261:LEU:HB2	9:Fc:284:TRP:HH2	1.80	0.46
9:Fc:420:LEU:CD1	9:Fc:579:LEU:HD21	2.45	0.46
9:Fc:430:MET:N	9:Fc:430:MET:HE2	2.31	0.46

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
9:Fc:837:PHE:O	9:Fc:841:ILE:HG22	2.16	0.46
9:Fe:756:LEU:O	9:Fe:760:ILE:HG12	2.15	0.46
9:Ff:463:MET:N	9:Ff:463:MET:HE3	2.30	0.46
5:Ae:247:ILE:HD12	5:Ae:254:ILE:HD12	1.97	0.46
6:Al:121:PRO:HB3	7:Ee:797:ARG:NH2	2.30	0.46
6:Al:240:THR:C	6:Al:241:LEU:HD23	2.40	0.46
5:Bc:219:ARG:HD3	6:Bj:148:PRO:O	2.15	0.46
6:Bd:182:LEU:HD12	6:Bd:183:ASP:N	2.30	0.46
5:Bi:246:LEU:HG	5:Bi:255:LEU:HD23	1.98	0.46
6:Bp:107:ILE:N	6:Bp:110:LYS:HZ1	2.13	0.46
3:Ce:2:VAL:HG22	3:Ce:8:VAL:HG23	1.98	0.46
6:Cz:114:ASP:O	6:Cz:117:ARG:HG3	2.16	0.46
6:Cz:150:LYS:HA	6:Cz:150:LYS:CE	2.46	0.46
1:Da:15:ASN:HB3	2:Dh:9:ARG:NH2	2.30	0.46
6:Df:170:ARG:HG3	6:Df:248:VAL:HA	1.98	0.46
1:Dm:3:ASP:HA	1:Dm:6:LEU:HG	1.97	0.46
6:Dr:109:LYS:O	6:Dr:113:LYS:HG2	2.14	0.46
1:Ds:6:LEU:O	1:Ds:10:GLU:HG2	2.16	0.46
6:Dx:122:LEU:HG	7:Et:808:LEU:HD11	1.96	0.46
6:Dx:182:LEU:HG	6:Dx:255:ARG:HE	1.81	0.46
6:Ed:180:VAL:HG13	6:Ed:253:ASP:HA	1.97	0.46
6:Ed:228:VAL:HG13	6:Ed:237:VAL:HB	1.96	0.46
7:Ef:782:ASN:HA	7:Ef:785:LEU:HG	1.97	0.46
8:Ew:107:GLN:HG3	8:Ex:387:THR:HG22	1.97	0.46
8:Ew:115:LEU:HD12	8:Ew:118:THR:OG1	2.16	0.46
8:Ew:172:ASP:O	8:Ew:176:GLN:HG2	2.16	0.46
8:Ey:85:PHE:HE2	8:Ez:376:PHE:HE1	1.61	0.46
8:Ey:327:SER:HA	8:Ey:330:PHE:CD2	2.50	0.46
8:Ez:89:ILE:HG22	8:Ez:349:TYR:CE1	2.48	0.46
8:Ez:399:TRP:HH2	8:Ez:414:GLU:HB3	1.81	0.46
9:Fb:274:PHE:CE2	9:Fb:515:LEU:HB2	2.50	0.46
9:Fb:666:GLY:HA3	9:Fb:699:PHE:CZ	2.51	0.46
9:Fb:725:PHE:HA	9:Fb:728:ILE:HG22	1.97	0.46
9:Fc:500:GLN:HA	9:Fc:503:LYS:CG	2.39	0.46
9:Fd:377:GLN:O	9:Fd:412:VAL:HA	2.16	0.46
9:Fd:813:SER:O	9:Fd:817:ILE:HG13	2.15	0.46
9:Fe:234:ILE:HD12	9:Fe:234:ILE:H	1.81	0.46
9:Fe:756:LEU:HA	9:Fe:759:MET:CE	2.46	0.46
6:Af:112:PHE:CD1	6:Af:112:PHE:C	2.94	0.46
6:Ar:126:GLN:HA	6:Ar:129:LYS:NZ	2.26	0.46
6:Ar:156:GLN:O	6:Ar:254:LEU:HA	2.16	0.46

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
6:Ar:181:PHE:HZ	6:Ar:213:LEU:HD12	1.80	0.46
6:Ar:225:ASN:HB2	6:Ax:250:TYR:OH	2.16	0.46
6:Bd:196:LEU:HD23	6:Bd:199:PRO:HB3	1.97	0.46
6:Bd:208:LYS:HG2	7:Ej:824:THR:HG23	1.98	0.46
6:Bj:239:LEU:HD21	6:Bj:254:LEU:HD21	1.96	0.46
6:Bv:229:ARG:HH22	6:Bv:233:LEU:HB3	1.81	0.46
6:Cb:141:LYS:HZ1	7:Em:808:LEU:HD13	1.80	0.46
6:Ch:107:ILE:HG13	6:Ch:110:LYS:HZ3	1.81	0.46
6:Cn:190:PRO:O	6:Cn:231:ARG:HG2	2.15	0.46
6:Cz:144:THR:HA	6:Df:131:LYS:HZ3	1.81	0.46
2:Db:7:GLY:O	4:Dd:1:PRO:HB2	2.15	0.46
1:Dy:1:CYS:HA	2:Dz:9:ARG:CZ	2.45	0.46
6:Ed:149:PRO:HB2	6:Ed:248:VAL:HG12	1.97	0.46
7:Eo:808:LEU:HB3	7:Eo:812:TRP:CZ2	2.51	0.46
7:Ep:806:THR:HA	7:Ep:809:VAL:CG1	2.45	0.46
8:Ew:380:THR:HB	8:Fa:82:TYR:OH	2.16	0.46
8:Ey:293:PRO:HG3	8:Ey:329:TYR:CE2	2.51	0.46
8:Ez:246:ASN:HA	8:Ez:249:ILE:HG13	1.96	0.46
9:Fb:546:LYS:HG3	9:Fb:650:GLU:OE2	2.16	0.46
9:Fb:729:MET:HE2	9:Fb:729:MET:O	2.16	0.46
9:Fb:819:TYR:O	9:Fb:823:ILE:HG22	2.16	0.46
9:Fd:143:TRP:HB2	9:Fd:816:ILE:HG12	1.97	0.46
9:Fe:526:LYS:HA	9:Fe:526:LYS:HE3	1.96	0.46
9:Fe:702:THR:O	9:Fe:705:LEU:HD12	2.16	0.46
9:Fe:771:ALA:O	9:Fe:774:GLU:HB2	2.14	0.46
9:Ff:290:LEU:HA	9:Ff:313:MET:SD	2.55	0.46
9:Ff:428:GLY:HA2	9:Ff:431:MET:HG2	1.97	0.46
9:Ff:464:ALA:HB1	9:Ff:819:TYR:CE2	2.51	0.46
5:Ae:218:GLY:HA3	6:Al:148:PRO:HD3	1.96	0.46
3:Ao:1:ARG:HH22	3:Ao:3:SER:HA	1.80	0.46
1:Ay:3:ASP:OD1	3:Ba:16:ASP:HB2	2.16	0.46
5:Bi:216:ILE:HD11	6:Bp:148:PRO:HG2	1.97	0.46
3:Bm:2:VAL:HG21	6:Cb:129:LYS:HD3	1.96	0.46
2:Br:4:MET:CE	5:Bu:250:LEU:H	2.29	0.46
6:Cb:230:LEU:HB2	6:Cb:233:LEU:HD23	1.98	0.46
1:Cc:4:ALA:HB2	3:Ce:14:TYR:HB2	1.98	0.46
5:Cg:210:TYR:HA	5:Cg:224:GLY:HA2	1.98	0.46
6:Ch:123:ASN:ND2	6:Ch:124:PRO:HD2	2.31	0.46
3:Cw:1:ARG:H3	3:Cw:10:THR:HG23	1.81	0.46
5:Dk:253:ARG:HE	5:Dk:255:LEU:HD21	1.80	0.46
6:Dr:173:GLN:CD	6:Dr:220:LEU:HD23	2.40	0.46

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
5:Dw:269:SER:C	6:Dx:150:LYS:HD2	2.40	0.46
6:Dx:107:ILE:HG23	6:Dx:110:LYS:NZ	2.31	0.46
6:Ed:209:THR:HG22	7:Ee:824:THR:HG21	1.96	0.46
7:Eg:810:GLN:HA	7:Eg:813:LYS:CE	2.46	0.46
8:Ey:349:TYR:HB3	8:Ey:435:ARG:NH1	2.31	0.46
8:Ez:189:PHE:CG	8:Ez:295:LEU:HD21	2.51	0.46
8:Ez:294:LYS:HE2	8:Ez:294:LYS:HB2	1.55	0.46
8:Ez:419:LEU:HD11	8:Fa:415:ILE:HG23	1.97	0.46
9:Fb:102:TRP:CE3	9:Fb:105:LEU:HD13	2.50	0.46
9:Fb:631:LEU:HD21	9:Fc:628:MET:HE2	1.98	0.46
9:Fb:937:PRO:O	9:Fb:941:LEU:HG	2.15	0.46
9:Fc:144:GLU:HA	9:Fc:147:LEU:HG	1.97	0.46
9:Fc:745:ILE:CG2	9:Fc:919:THR:HG21	2.44	0.46
9:Fd:448:LYS:HB2	9:Fd:448:LYS:HE2	1.67	0.46
9:Fe:655:ILE:O	9:Fe:659:PHE:HD2	1.99	0.46
9:Fe:757:PRO:HA	9:Fe:760:ILE:CG1	2.45	0.46
9:Ff:187:GLY:O	9:Ff:191:MET:HG3	2.16	0.46
9:Ff:460:GLY:C	9:Ff:463:MET:HE1	2.41	0.46
9:Ff:799:GLU:HA	9:Ff:802:ILE:HD12	1.98	0.46
7:Fh:8:LEU:HD12	7:Fh:9:LYS:N	2.30	0.46
6:Af:216:GLN:NE2	7:Ee:818:GLN:HG2	2.31	0.46
6:Af:228:VAL:HB	6:Af:237:VAL:HG13	1.98	0.46
5:Ak:250:LEU:HD12	5:Ak:251:GLN:HG3	1.96	0.46
1:Ay:16:ALA:C	2:Bf:9:ARG:HG3	2.41	0.46
6:Bj:141:LYS:HZ2	7:Ej:812:TRP:CD1	2.34	0.46
6:Cb:166:PRO:HG3	6:Ch:250:TYR:CE1	2.50	0.46
6:Ch:108:ASP:HA	7:El:797:ARG:NH1	2.30	0.46
6:Cz:252:VAL:HG12	6:Cz:254:LEU:HD13	1.97	0.46
5:De:223:ILE:HD12	5:De:228:SER:O	2.16	0.46
5:Dq:218:GLY:HA3	6:Dx:148:PRO:HD3	1.97	0.46
7:Ej:775:GLN:O	7:Ej:778:LEU:HD12	2.16	0.46
8:Ew:298:TYR:HA	8:Ew:301:LEU:CD1	2.45	0.46
8:Ey:81:ALA:HB2	8:Ey:378:MET:HE3	1.98	0.46
8:Ez:302:TRP:CD1	8:Ez:302:TRP:C	2.94	0.46
8:Ez:400:ILE:HG13	8:Ez:401:LYS:N	2.30	0.46
8:Fa:252:LEU:HG	8:Fa:373:LEU:HA	1.98	0.46
9:Fc:163:PRO:HB2	9:Fc:579:LEU:HD12	1.98	0.46
9:Fc:639:TYR:CD1	9:Fc:639:TYR:C	2.93	0.46
9:Fc:751:TYR:C	9:Fc:754:PRO:HD2	2.41	0.46
9:Fe:370:PHE:CD1	9:Fe:377:GLN:HB3	2.51	0.46
9:Fe:505:PHE:HA	9:Fe:509:CYS:SG	2.55	0.46

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
9:Fe:705:LEU:HA	9:Fe:708:LEU:HB2	1.98	0.46
9:Fe:940:VAL:HA	9:Fe:943:TRP:CZ3	2.51	0.46
9:Ff:636:VAL:O	9:Ff:640:VAL:HG13	2.16	0.46
5:Ae:222:LEU:HD11	5:Ae:262:ILE:HG21	1.98	0.46
6:Af:179:LEU:HD22	6:Af:252:VAL:HG22	1.98	0.46
6:Al:129:LYS:O	6:Al:133:ILE:HG22	2.16	0.46
6:Al:238:MET:HB3	6:Ar:250:TYR:HD2	1.81	0.46
6:Ar:169:ILE:HD11	6:Ar:239:LEU:HD22	1.98	0.46
6:Bp:174:GLY:HA2	6:Bp:216:GLN:HE21	1.80	0.46
6:Bp:198:ASP:HB3	6:Bp:202:PHE:HE2	1.79	0.46
5:Bu:230:LEU:HD12	5:Bu:231:THR:N	2.31	0.46
6:Bv:117:ARG:NH1	6:Bv:118:ASN:HB3	2.21	0.46
5:Cg:253:ARG:HD3	5:Cg:255:LEU:HD11	1.98	0.46
1:Ci:8:ALA:HB1	3:Ck:1:ARG:HD2	1.97	0.46
6:Cn:226:LEU:HB2	6:Cn:241:LEU:HD11	1.98	0.46
4:Dd:2:PHE:HB3	5:De:251:GLN:NE2	2.31	0.46
1:Dm:12:HIS:HD2	3:Do:9:TYR:HD2	1.64	0.46
8:Ew:319:GLN:OE1	8:Ew:320:LYS:HG2	2.16	0.46
8:Ew:413:LYS:CE	8:Ex:404:ASN:HA	2.44	0.46
8:Ew:437:LEU:CD2	8:Ex:432:ILE:HG22	2.45	0.46
8:Ex:448:LEU:HD21	8:Ey:340:THR:HG22	1.96	0.46
8:Ey:118:THR:HA	8:Ey:121:LYS:HG2	1.97	0.46
8:Ey:281:ARG:HG3	8:Ey:286:GLN:NE2	2.31	0.46
8:Ez:302:TRP:HD1	8:Ez:303:ASN:ND2	2.08	0.46
8:Ez:326:LEU:N	8:Ez:326:LEU:HD23	2.30	0.46
8:Ez:418:LEU:HD12	8:Ez:419:LEU:HD23	1.98	0.46
9:Fc:124:MET:O	9:Fc:128:PHE:HD1	1.98	0.46
9:Fc:239:PHE:HD1	9:Fc:261:LEU:HG	1.81	0.46
9:Fc:240:VAL:O	9:Fc:243:GLN:HG3	2.15	0.46
9:Fc:452:PHE:CZ	9:Fc:470:ASP:HB3	2.51	0.46
9:Fc:935:HIS:O	9:Fc:939:LYS:HD2	2.16	0.46
9:Fd:526:LYS:HA	9:Fd:529:GLN:HE21	1.81	0.46
9:Fd:674:GLY:O	9:Fd:678:LEU:HG	2.15	0.46
9:Fd:740:GLY:HA2	9:Fd:743:VAL:HG12	1.98	0.46
9:Fe:105:LEU:O	9:Fe:108:THR:HG22	2.15	0.46
9:Fe:137:GLY:HA2	9:Fe:140:ASP:OD1	2.16	0.46
9:Fe:918:TYR:O	9:Fe:921:MET:HB2	2.16	0.46
9:Ff:237:VAL:HG21	9:Ff:263:MET:HB2	1.97	0.46
9:Ff:323:SER:HA	9:Ff:326:GLN:HG2	1.96	0.46
9:Ff:937:PRO:O	9:Ff:940:VAL:HG22	2.16	0.46
5:Ak:217:PRO:HG2	6:Ar:143:ALA:HB1	1.98	0.46

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:Bk:11:TYR:CD2	3:Bm:9:TYR:HB3	2.51	0.46
6:Bp:139:TYR:O	6:Bp:143:ALA:HB2	2.15	0.46
3:Bs:13:LYS:HD3	3:Bs:13:LYS:HA	1.67	0.46
6:Cb:156:GLN:HE22	6:Cb:167:PRO:HG3	1.81	0.46
6:Cb:189:TRP:HB3	6:Cb:230:LEU:HD12	1.97	0.46
5:Cs:210:TYR:CB	5:Cs:222:LEU:HD11	2.46	0.46
6:Df:134:TYR:HE2	7:Eq:812:TRP:HB2	1.81	0.46
6:Df:200:SER:HB3	7:Es:816:GLU:OE2	2.16	0.46
6:Dr:202:PHE:CE1	6:Dr:241:LEU:HB2	2.51	0.46
5:Ec:264:PHE:HB3	5:Ec:269:SER:HB3	1.98	0.46
7:Eh:810:GLN:O	7:Eh:814:GLN:HG2	2.15	0.46
7:Er:801:MET:SD	7:Er:802:LEU:N	2.89	0.46
8:Ew:228:PRO:HB2	8:Ew:232:SER:OG	2.14	0.46
8:Ex:94:MET:CE	8:Ey:241:SER:HB2	2.46	0.46
8:Ey:335:VAL:O	8:Ey:339:GLN:HG2	2.16	0.46
8:Ez:189:PHE:HA	8:Ez:295:LEU:HD11	1.98	0.46
8:Ez:219:PRO:HA	8:Ez:220:PRO:HD3	1.82	0.46
8:Fa:101:PHE:HZ	8:Fa:116:ALA:HA	1.81	0.46
9:Fb:288:SER:HA	9:Fb:291:ASN:HB3	1.98	0.46
9:Fb:550:LEU:H	9:Fb:550:LEU:HD12	1.80	0.46
9:Fb:939:LYS:HG3	9:Fb:940:VAL:H	1.79	0.46
9:Fc:148:SER:O	9:Fc:152:ARG:HG2	2.16	0.46
9:Fc:251:LEU:HD11	9:Fc:326:GLN:NE2	2.25	0.46
9:Fc:724:ILE:O	9:Fc:728:ILE:HG22	2.15	0.46
9:Fc:820:ILE:HA	9:Fc:823:ILE:CD1	2.46	0.46
9:Fc:933:ILE:HD12	9:Fc:933:ILE:H	1.80	0.46
9:Fd:331:SER:O	9:Fd:334:ILE:HG13	2.16	0.46
9:Fe:167:LEU:HD21	9:Fe:579:LEU:HD11	1.98	0.46
9:Ff:505:PHE:CG	9:Ff:524:SER:HB3	2.50	0.46
5:Ak:246:LEU:HD22	5:Ak:255:LEU:HG	1.98	0.45
6:Ax:140:ALA:HB1	6:Bd:127:VAL:HB	1.97	0.45
6:Ax:184:SER:HA	6:Ax:255:ARG:CZ	2.46	0.45
6:Ax:190:PRO:O	6:Ax:231:ARG:HG2	2.15	0.45
2:Az:5:ILE:HD11	6:Bj:139:TYR:CE2	2.49	0.45
6:Bd:195:ASP:HB2	6:Bd:229:ARG:HH22	1.81	0.45
1:Be:3:ASP:O	1:Be:6:LEU:HD12	2.16	0.45
6:Cb:132:GLN:NE2	6:Cb:136:THR:HG23	2.31	0.45
2:Cd:4:MET:HB2	4:Cf:2:PHE:HZ	1.80	0.45
6:Cn:208:LYS:HD3	6:Cn:208:LYS:HA	1.72	0.45
5:De:207:ARG:NH2	5:De:240:GLY:HA3	2.31	0.45
6:Dl:106:VAL:HA	6:Dl:109:LYS:NZ	2.30	0.45

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
6:Dl:119:LEU:C	6:Dl:120:TYR:HD2	2.24	0.45
7:Eo:805:ALA:HA	7:Eo:808:LEU:HG	1.97	0.45
7:Eq:772:GLU:HA	7:Eq:775:GLN:CD	2.40	0.45
8:Ew:430:ARG:HH22	8:Ex:428:LEU:CD1	2.25	0.45
8:Ex:121:LYS:NZ	8:Ex:171:ASN:HB3	2.31	0.45
8:Ex:294:LYS:O	8:Ex:298:TYR:HB3	2.16	0.45
8:Ex:308:LYS:HG2	8:Ex:309:PRO:HD2	1.98	0.45
8:Ex:424:TYR:CZ	8:Ey:379:ALA:O	2.70	0.45
8:Ey:191:MET:HE1	8:Ey:198:TRP:CD2	2.50	0.45
8:Ez:420:ALA:O	8:Ez:424:TYR:CD1	2.69	0.45
9:Fb:594:ILE:HB	9:Fb:660:LEU:HD12	1.97	0.45
9:Fb:658:ALA:O	9:Fb:662:ILE:HG12	2.16	0.45
9:Fb:709:ASN:ND2	9:Fc:597:LYS:H	2.14	0.45
9:Fc:178:ALA:HB3	9:Fc:179:LYS:HZ3	1.81	0.45
9:Fc:187:GLY:HA2	9:Fc:496:PHE:HE1	1.81	0.45
9:Fc:274:PHE:CE2	9:Fc:515:LEU:HB2	2.51	0.45
9:Fc:761:PHE:CD1	9:Fc:761:PHE:C	2.95	0.45
9:Fc:788:HIS:ND1	9:Fc:789:PRO:HD2	2.31	0.45
9:Fd:49:LEU:HD22	9:Fd:129:MET:HE2	1.98	0.45
9:Fd:689:LEU:HA	9:Fd:692:MET:HE2	1.98	0.45
9:Fe:287:ILE:HG13	9:Fe:332:ARG:HH12	1.81	0.45
9:Fe:503:LYS:HA	9:Fe:503:LYS:HD2	1.78	0.45
9:Fe:921:MET:O	9:Fe:925:ILE:HG13	2.17	0.45
9:Ff:322:SER:O	9:Ff:326:GLN:HG2	2.16	0.45
9:Ff:462:ILE:C	9:Ff:761:PHE:HZ	2.24	0.45
9:Ff:705:LEU:O	9:Ff:708:LEU:HB2	2.15	0.45
9:Ff:911:PHE:HA	9:Ff:914:SER:OG	2.17	0.45
9:Ff:926:VAL:HA	9:Ff:929:ALA:HB3	1.97	0.45
1:Aa:16:ALA:HA	4:Aj:3:GLY:HA3	1.98	0.45
3:Ac:13:LYS:HD3	3:Ac:13:LYS:HA	1.66	0.45
1:Ag:16:ALA:HA	4:Ap:3:GLY:HA3	1.97	0.45
1:Am:11:TYR:CE2	3:Ao:9:TYR:HB3	2.51	0.45
6:Ar:131:LYS:HG3	6:Ar:135:GLU:OE1	2.17	0.45
5:Bu:264:PHE:CE2	6:Bv:246:LYS:HB3	2.51	0.45
6:Bv:105:GLU:O	6:Bv:108:ASP:HB3	2.16	0.45
6:Cn:121:PRO:HB2	7:En:801:MET:HG3	1.96	0.45
6:Ct:200:SER:O	6:Ct:218:THR:HB	2.17	0.45
6:Df:172:SER:OG	6:Df:248:VAL:HG12	2.16	0.45
5:Dw:213:GLN:HB2	6:Dx:170:ARG:NH2	2.31	0.45
6:Dx:196:LEU:HB2	6:Dx:226:LEU:HD13	1.97	0.45
6:Ed:171:LEU:HB2	6:Ed:243:PRO:HA	1.98	0.45

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
6:Ed:221:TYR:CD2	6:Ed:244:GLY:HA3	2.50	0.45
7:El:812:TRP:HA	7:El:812:TRP:CE3	2.51	0.45
7:En:793:GLU:HA	7:En:796:GLN:HE21	1.80	0.45
8:Ew:54:ILE:HG23	8:Ex:45:ASN:OD1	2.15	0.45
8:Ew:174:VAL:HG21	8:Ew:345:SER:HB3	1.97	0.45
8:Ew:415:ILE:CG2	8:Fa:419:LEU:HD21	2.47	0.45
8:Ey:108:GLY:O	8:Ey:112:ILE:HG12	2.16	0.45
8:Ez:101:PHE:CD2	8:Fa:252:LEU:HB2	2.51	0.45
8:Ez:424:TYR:O	8:Ez:428:LEU:HG	2.16	0.45
8:Fa:209:LEU:HA	8:Fa:212:GLN:NE2	2.12	0.45
9:Fb:704:TRP:CH2	9:Fc:731:ALA:HA	2.43	0.45
9:Fc:56:ILE:H	9:Fc:56:ILE:HG13	1.61	0.45
9:Fc:429:ILE:O	9:Fc:432:PRO:HD2	2.16	0.45
9:Fc:633:TYR:HA	9:Fc:637:PHE:CD1	2.50	0.45
9:Fd:72:GLY:O	9:Fd:76:ILE:HG13	2.16	0.45
9:Fe:38:LEU:HD21	9:Fe:129:MET:HE1	1.97	0.45
9:Fe:429:ILE:HB	9:Fe:430:MET:HE2	1.98	0.45
9:Fe:759:MET:HB3	9:Ff:921:MET:HE2	1.98	0.45
9:Ff:225:PHE:CD2	9:Ff:519:TRP:HB3	2.51	0.45
2:Ab:4:MET:CE	5:Ae:250:LEU:H	2.30	0.45
5:Aq:234:GLU:HA	5:Aq:247:ILE:HD11	1.98	0.45
6:Ar:138:GLU:O	6:Ar:141:LYS:HB3	2.17	0.45
6:Bd:125:GLU:O	6:Bd:129:LYS:HG3	2.17	0.45
6:Bd:129:LYS:HB3	6:Bj:112:PHE:HZ	1.81	0.45
5:Bo:253:ARG:HG2	5:Bo:263:LYS:NZ	2.31	0.45
5:Bo:263:LYS:HA	5:Bo:263:LYS:HE3	1.98	0.45
6:Bv:205:GLN:HB2	7:El:820:TYR:OH	2.15	0.45
6:Ct:108:ASP:HA	7:En:797:ARG:NH2	2.31	0.45
1:Dg:3:ASP:CG	3:Di:14:TYR:HB3	2.41	0.45
1:Dm:1:CYS:HA	4:Dp:2:PHE:CG	2.52	0.45
3:Ea:13:LYS:HD3	3:Ea:13:LYS:HA	1.67	0.45
5:Ec:212:ILE:HG22	5:Ec:263:LYS:O	2.16	0.45
7:Ee:792:GLN:HA	7:Ee:795:GLN:NE2	2.31	0.45
7:Et:797:ARG:HA	7:Et:800:ASP:OD1	2.16	0.45
7:Eu:809:VAL:HG22	7:Eu:813:LYS:HE3	1.98	0.45
8:Ew:244:ASN:HD22	8:Fa:175:SER:HB3	1.81	0.45
8:Ew:301:LEU:HD13	8:Ew:326:LEU:HD22	1.98	0.45
8:Ey:179:PHE:HA	8:Ey:182:LEU:HG	1.98	0.45
8:Ez:211:MET:HB3	8:Ez:287:VAL:O	2.17	0.45
9:Fb:94:LEU:HD23	9:Fb:94:LEU:HA	1.75	0.45
9:Fb:237:VAL:HA	9:Fb:265:ASN:HD22	1.81	0.45

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
9:Fb:453:ILE:H	9:Fb:453:ILE:HD12	1.80	0.45
9:Fb:596:PHE:C	9:Fb:597:LYS:HD3	2.41	0.45
9:Fb:668:LYS:O	9:Fb:672:ILE:HG12	2.16	0.45
9:Fb:680:GLN:HB2	9:Fb:683:ILE:HG22	1.98	0.45
9:Fb:760:ILE:HA	9:Fb:763:PHE:CD2	2.51	0.45
9:Fb:785:GLY:HA2	9:Fb:788:HIS:ND1	2.32	0.45
9:Fc:192:LEU:HD23	9:Fc:234:ILE:HD11	1.97	0.45
9:Fc:725:PHE:O	9:Fc:725:PHE:HD1	1.98	0.45
9:Fc:911:PHE:O	9:Fc:915:ILE:HD12	2.16	0.45
9:Fd:696:TYR:OH	9:Fd:745:ILE:HD12	2.17	0.45
7:Fj:28:LEU:HD23	7:Fj:29:ILE:HD13	1.98	0.45
6:Af:230:LEU:HD12	6:Af:231:ARG:N	2.23	0.45
1:Ay:3:ASP:HA	1:Ay:6:LEU:HG	1.97	0.45
5:Bc:233:ARG:HD3	5:Bi:269:SER:OXT	2.17	0.45
6:Bd:109:LYS:O	6:Bd:113:LYS:HG2	2.17	0.45
6:Bd:117:ARG:NE	7:Eh:797:ARG:HD2	2.30	0.45
5:Bo:207:ARG:HH12	5:Bo:209:ILE:HA	1.82	0.45
1:Bq:6:LEU:HD12	1:Bq:7:ALA:N	2.31	0.45
2:Br:9:ARG:HA	2:Br:9:ARG:HD3	1.88	0.45
3:By:2:VAL:HG21	6:Cn:129:LYS:CG	2.45	0.45
6:Ch:238:MET:HE2	6:Cn:250:TYR:CE1	2.51	0.45
1:Ci:12:HIS:CE1	3:Ck:9:TYR:HD2	2.34	0.45
6:Cn:130:LEU:HA	6:Cn:133:ILE:CG1	2.45	0.45
6:Cn:204:ILE:HD12	6:Cn:215:ILE:HG12	1.98	0.45
6:Ct:115:MET:CE	7:En:798:THR:HA	2.47	0.45
6:Ct:179:LEU:O	6:Ct:212:THR:HA	2.16	0.45
1:Dg:3:ASP:O	1:Dg:6:LEU:HD12	2.16	0.45
6:Dl:106:VAL:HA	6:Dl:109:LYS:HZ2	1.81	0.45
6:Dl:191:ILE:HB	6:Dl:206:TRP:CZ2	2.51	0.45
6:Dx:122:LEU:CD2	7:Et:801:MET:HG2	2.47	0.45
5:Ec:221:TRP:CH2	6:Ed:165:THR:HB	2.51	0.45
7:Ep:797:ARG:HA	7:Ep:800:ASP:OD1	2.17	0.45
8:Ey:381:ARG:HG2	8:Ey:381:ARG:HH11	1.81	0.45
8:Ey:382:ARG:NE	8:Ey:418:LEU:HB3	2.31	0.45
8:Ez:76:LEU:HG	8:Ez:80:TYR:CE2	2.52	0.45
8:Ez:331:ASN:O	8:Ez:335:VAL:HG22	2.16	0.45
9:Fb:631:LEU:CD2	9:Fc:628:MET:HE2	2.46	0.45
9:Fc:278:ILE:HD13	9:Fc:513:TYR:CE1	2.51	0.45
9:Fc:930:PHE:HA	9:Fc:933:ILE:CD1	2.43	0.45
9:Fe:692:MET:SD	9:Fe:692:MET:N	2.89	0.45
9:Fe:745:ILE:HD11	9:Fe:916:LEU:HD13	1.99	0.45

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
9:Fe:757:PRO:HA	9:Fe:760:ILE:HD11	1.99	0.45
6:Al:115:MET:CE	7:Ev:801:MET:HE2	2.47	0.45
6:Ar:179:LEU:O	6:Ar:212:THR:HA	2.17	0.45
6:Bj:158:VAL:HB	6:Bj:256:VAL:HA	1.98	0.45
5:Ca:246:LEU:HD11	5:Ca:255:LEU:HD23	1.99	0.45
6:Ch:204:ILE:HD11	6:Ch:213:LEU:HB3	1.97	0.45
6:Cn:190:PRO:HA	6:Cn:211:ASN:HB3	1.98	0.45
6:Cn:256:VAL:HG12	6:Cn:258:GLY:H	1.82	0.45
6:Ct:166:PRO:HB3	6:Ct:238:MET:HB3	1.97	0.45
5:De:213:GLN:OE1	5:De:223:ILE:HB	2.16	0.45
6:Dx:113:LYS:HA	6:Dx:113:LYS:HD3	1.74	0.45
6:Dx:151:PRO:HA	6:Dx:248:VAL:HG13	1.98	0.45
7:Ei:809:VAL:O	7:Ei:813:LYS:HG2	2.16	0.45
7:Er:792:GLN:CD	7:Er:792:GLN:H	2.24	0.45
7:Er:794:ILE:HG23	7:Er:797:ARG:NH2	2.31	0.45
7:Es:771:ALA:O	7:Es:775:GLN:HG2	2.16	0.45
7:Ev:805:ALA:O	7:Ev:808:LEU:HD12	2.16	0.45
8:Ew:250:ALA:N	8:Ew:251:PRO:HD2	2.32	0.45
8:Ew:384:PHE:HB3	8:Fa:82:TYR:CE2	2.51	0.45
8:Ew:424:TYR:CE2	8:Ex:383:LEU:HD22	2.51	0.45
8:Ex:384:PHE:HA	8:Ex:398:GLN:HE22	1.81	0.45
8:Ey:374:ASN:HA	8:Ey:377:ASN:OD1	2.16	0.45
8:Ey:437:LEU:CD1	8:Ez:436:ILE:HD13	2.47	0.45
8:Ez:78:GLN:O	8:Fa:383:LEU:HD11	2.16	0.45
8:Ez:343:GLY:O	8:Ez:347:LEU:HG	2.16	0.45
8:Ez:359:MET:HE3	8:Ez:367:ASN:O	2.17	0.45
8:Fa:238:PRO:HG2	8:Fa:263:GLN:HB3	1.99	0.45
9:Fb:77:MET:O	9:Fb:81:MET:HG2	2.17	0.45
9:Fb:632:PHE:O	9:Fb:636:VAL:HB	2.16	0.45
9:Fc:117:LYS:HD3	9:Fc:118:ALA:H	1.82	0.45
9:Fc:179:LYS:HB2	9:Fc:492:ASP:HA	1.98	0.45
9:Fd:66:SER:OG	7:Fk:38:LYS:HG3	2.15	0.45
9:Fd:515:LEU:H	9:Fd:515:LEU:HD22	1.81	0.45
9:Ff:251:LEU:HD21	9:Ff:326:GLN:HE22	1.81	0.45
9:Ff:674:GLY:HA2	9:Ff:692:MET:CE	2.43	0.45
7:Fh:4:LYS:HB2	7:Fh:4:LYS:HE2	1.77	0.45
7:Fk:30:ILE:O	7:Fk:34:ILE:HG22	2.16	0.45
6:Af:170:ARG:HD3	6:Af:247:ALA:C	2.42	0.45
5:Ak:219:ARG:NH1	5:Ak:233:ARG:HD3	2.30	0.45
2:An:2:VAL:HG13	3:Ao:1:ARG:HD2	1.98	0.45
5:Bc:211:TYR:CE1	5:Bc:265:SER:HA	2.52	0.45

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
6:Bj:173:GLN:HA	6:Bj:217:ALA:O	2.16	0.45
6:Bj:229:ARG:HH12	6:Bj:231:ARG:HD3	1.80	0.45
2:Br:4:MET:HE1	5:Bu:249:SER:N	2.31	0.45
5:Ca:215:VAL:HG11	5:Ca:252:GLY:HA2	1.98	0.45
6:Cb:106:VAL:HA	6:Cb:109:LYS:HD3	1.99	0.45
6:Cb:188:PRO:HB3	6:Cb:211:ASN:ND2	2.32	0.45
3:Ce:1:ARG:HG2	3:Ce:2:VAL:N	2.31	0.45
5:Cg:219:ARG:NH1	5:Cg:233:ARG:HD3	2.32	0.45
6:Cn:145:PRO:HD3	6:Ct:131:LYS:HE2	1.98	0.45
6:Dl:225:ASN:HD21	6:Dr:176:VAL:N	2.14	0.45
6:Dx:215:ILE:HD12	6:Dx:216:GLN:N	2.31	0.45
5:Ec:213:GLN:HE21	5:Ec:221:TRP:HE3	1.63	0.45
8:Ey:280:ILE:HG21	8:Ey:351:LEU:HD22	1.97	0.45
8:Ey:349:TYR:HB3	8:Ey:435:ARG:HH12	1.80	0.45
8:Ey:374:ASN:C	8:Ey:378:MET:HE2	2.41	0.45
9:Fb:65:ASN:HA	9:Fb:68:VAL:HG22	1.97	0.45
9:Fb:143:TRP:CZ2	9:Fb:147:LEU:HD23	2.52	0.45
9:Fb:718:PRO:HB3	9:Fc:719:LEU:HD12	1.98	0.45
9:Fb:730:MET:CE	9:Ff:708:LEU:HD22	2.46	0.45
9:Fc:41:LEU:HD22	9:Fc:42:PHE:CE1	2.51	0.45
9:Fc:127:PHE:O	9:Fc:131:VAL:HG23	2.16	0.45
9:Fc:514:SER:O	9:Fc:518:THR:HG23	2.16	0.45
9:Fd:833:LEU:O	9:Fd:837:PHE:CD1	2.70	0.45
9:Fe:45:VAL:HG11	9:Fe:129:MET:HG2	1.97	0.45
9:Ff:771:ALA:HA	9:Ff:774:GLU:HB3	1.97	0.45
9:Ff:806:VAL:O	9:Ff:810:LEU:HG	2.17	0.45
7:Fg:24:THR:HA	7:Fg:27:LEU:CD1	2.46	0.45
6:Al:193:ALA:HB3	6:Al:229:ARG:HB3	1.99	0.45
5:Bc:218:GLY:HA3	6:Bj:148:PRO:HD3	1.97	0.45
5:Bc:236:SER:N	5:Bc:243:MET:HE1	2.31	0.45
6:Bp:241:LEU:HD12	6:Bp:241:LEU:O	2.17	0.45
1:Bq:11:TYR:CD2	3:Bs:9:TYR:HB3	2.51	0.45
5:Bu:254:ILE:HB	5:Bu:262:ILE:HG22	1.97	0.45
6:Bv:181:PHE:HD2	6:Bv:230:LEU:HD21	1.82	0.45
6:Cb:105:GLU:HG2	6:Cb:109:LYS:NZ	2.31	0.45
6:Cn:185:THR:HG21	6:Cn:263:ALA:HA	1.99	0.45
5:Cy:255:LEU:HD13	5:Cy:261:VAL:HG13	1.97	0.45
6:Cz:225:ASN:HD21	6:Df:176:VAL:H	1.64	0.45
1:Da:3:ASP:HB2	3:Dc:14:TYR:HB3	1.98	0.45
6:Dl:113:LYS:HA	6:Dl:113:LYS:HD3	1.78	0.45
7:Ek:792:GLN:HA	7:Ek:795:GLN:OE1	2.16	0.45

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
7:Es:779:GLN:HA	7:Es:782:ASN:OD1	2.17	0.45
8:Ew:230:PHE:HA	8:Ew:235:TYR:CD2	2.52	0.45
8:Ex:104:ASP:HA	8:Ex:109:ASN:ND2	2.30	0.45
8:Ey:63:PRO:HA	8:Ez:48:LYS:HZ1	1.82	0.45
8:Ey:346:ASN:HA	8:Ey:435:ARG:HH22	1.80	0.45
8:Ez:62:ASN:HA	8:Fa:48:LYS:NZ	2.31	0.45
8:Ez:428:LEU:O	8:Ez:432:ILE:HG12	2.16	0.45
9:Fb:604:TYR:HA	9:Fb:641:TYR:OH	2.17	0.45
9:Fb:908:VAL:HG23	9:Fb:912:PHE:HE2	1.81	0.45
9:Fc:666:GLY:O	9:Fc:670:ILE:HG12	2.16	0.45
9:Fd:147:LEU:HB2	9:Fd:453:ILE:HG23	1.98	0.45
9:Fd:618:LEU:O	9:Fd:619:PHE:HD2	2.00	0.45
9:Fe:56:ILE:HG13	9:Fe:57:MET:SD	2.56	0.45
9:Ff:203:ASP:HA	9:Ff:206:LEU:HG	1.98	0.45
6:Af:145:PRO:HG2	6:Al:135:GLU:HG2	1.98	0.45
1:Am:6:LEU:HD12	1:Am:7:ALA:N	2.32	0.45
6:Ar:124:PRO:O	6:Ar:128:VAL:HG12	2.17	0.45
6:Ar:128:VAL:O	6:Ar:131:LYS:HB3	2.16	0.45
6:Ar:145:PRO:HD3	6:Ax:131:LYS:HZ3	1.82	0.45
6:Ar:191:ILE:HD12	6:Ar:206:TRP:NE1	2.32	0.45
5:Aw:269:SER:O	6:Ax:150:LYS:HE3	2.17	0.45
5:Bc:207:ARG:HH21	5:Bc:209:ILE:HA	1.82	0.45
5:Bc:247:ILE:HG12	5:Bc:254:ILE:HD12	1.98	0.45
6:Bd:107:ILE:HA	6:Bd:110:LYS:HZ3	1.82	0.45
6:Bd:190:PRO:HA	6:Bd:211:ASN:HB3	1.98	0.45
5:Bu:219:ARG:HH22	6:Cb:150:LYS:HD2	1.82	0.45
6:Ch:190:PRO:O	6:Ch:231:ARG:HG2	2.17	0.45
5:Cm:219:ARG:HG2	6:Ct:148:PRO:HD2	1.97	0.45
6:Cn:128:VAL:CG2	6:Cn:132:GLN:HE21	2.27	0.45
6:Cn:198:ASP:HA	7:Ep:816:GLU:OE1	2.16	0.45
6:Cn:225:ASN:HB3	6:Ct:250:TYR:OH	2.17	0.45
6:Ct:236:PRO:HB2	6:Cz:251:ARG:NH1	2.32	0.45
2:Dh:1:CYS:HA	3:Di:4:ILE:HD13	1.99	0.45
5:Ec:210:TYR:HD1	5:Ec:224:GLY:HA2	1.82	0.45
7:Ej:785:LEU:O	7:Ej:789:LYS:HG2	2.16	0.45
7:Ek:794:ILE:HG23	7:Ek:797:ARG:NH2	2.32	0.45
8:Ey:90:PRO:HB2	8:Ez:246:ASN:ND2	2.31	0.45
8:Fa:219:PRO:HG3	8:Fa:235:TYR:CE2	2.51	0.45
9:Fb:261:LEU:HB3	9:Fb:282:VAL:HG13	1.99	0.45
9:Fb:745:ILE:HG13	9:Fb:745:ILE:H	1.62	0.45
9:Fb:751:TYR:CE2	9:Fc:910:ALA:HA	2.52	0.45

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
9:Fc:170:ALA:HA	9:Fc:177:VAL:HG22	1.98	0.45
9:Fd:263:MET:H	9:Fd:263:MET:HE3	1.82	0.45
9:Fd:747:PHE:HB3	9:Fd:751:TYR:HB2	1.97	0.45
9:Fe:28:PRO:HB2	9:Fe:32:ASP:OD2	2.17	0.45
9:Fe:419:PHE:HE2	9:Fe:575:MET:HB3	1.81	0.45
9:Fe:756:LEU:HA	9:Fe:759:MET:SD	2.55	0.45
9:Ff:812:PRO:HA	9:Ff:815:MET:CE	2.46	0.45
7:Fh:28:LEU:O	7:Fh:32:VAL:HG13	2.16	0.45
4:Ap:2:PHE:HB2	5:Aq:251:GLN:NE2	2.32	0.45
1:As:16:ALA:HA	4:Bb:3:GLY:HA3	1.99	0.45
5:Aw:251:GLN:HB3	5:Aw:253:ARG:CZ	2.47	0.45
6:Ax:237:VAL:HG12	6:Ax:239:LEU:HD11	1.98	0.45
3:Bg:1:ARG:HA	6:Bv:125:GLU:OE2	2.16	0.45
6:Cb:117:ARG:CZ	7:El:801:MET:HE1	2.47	0.45
6:Ch:181:PHE:CD1	6:Ch:189:TRP:HB3	2.52	0.45
6:Cn:131:LYS:HA	7:En:812:TRP:CZ3	2.51	0.45
5:Cs:246:LEU:HD12	5:Cs:247:ILE:N	2.32	0.45
6:Ct:199:PRO:HG3	7:Eq:819:VAL:HG13	1.99	0.45
6:Cz:181:PHE:HD1	6:Cz:254:LEU:HD23	1.82	0.45
5:De:210:TYR:CE2	5:De:238:ILE:HD11	2.51	0.45
5:Dk:216:ILE:C	5:Dk:216:ILE:HD12	2.42	0.45
6:Dr:107:ILE:O	6:Dr:111:ALA:CB	2.64	0.45
6:Dr:238:MET:HE2	6:Dx:178:SER:HB3	1.97	0.45
5:Ec:212:ILE:HD12	5:Ec:222:LEU:HD22	1.99	0.45
7:Ef:781:GLN:HG3	9:Fe:337:GLN:CD	2.42	0.45
7:Eg:806:THR:HA	7:Eg:809:VAL:HG12	1.98	0.45
7:Eh:793:GLU:HA	7:Eh:796:GLN:HE21	1.82	0.45
7: Ei:782:ASN:HA	9:Fd:241:LYS:CD	2.45	0.45
7:Em:792:GLN:HA	7:Em:795:GLN:OE1	2.17	0.45
7:Em:793:GLU:HG2	7:Em:794:ILE:N	2.32	0.45
7:Eo:795:GLN:HA	7:Eo:798:THR:OG1	2.16	0.45
7:Eq:777:ILE:HD13	7:Eq:780:LYS:NZ	2.31	0.45
8:Ew:50:LEU:HB3	8:Ew:52:TYR:CE1	2.52	0.45
8:Ey:67:GLN:HA	8:Ey:413:LYS:HZ2	1.82	0.45
8:Ey:229:ALA:HB1	8:Ey:231:TYR:CE2	2.52	0.45
8:Ey:349:TYR:O	8:Ey:353:LYS:HD2	2.16	0.45
8:Ez:220:PRO:HD2	8:Ez:228:PRO:O	2.17	0.45
8:Ez:352:SER:HA	8:Ez:355:LEU:CD2	2.47	0.45
8:Ez:409:ALA:O	8:Ez:413:LYS:HG3	2.17	0.45
9:Fb:762:THR:O	9:Fb:766:PHE:CD1	2.70	0.45
9:Fb:766:PHE:HE2	9:Fc:925:ILE:HG23	1.82	0.45

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
9:Fc:268:LYS:NZ	9:Fd:404:SER:H	2.12	0.45
9:Fd:376:GLN:HB2	9:Fd:413:LEU:HD11	1.99	0.45
9:Fe:261:LEU:HB2	9:Fe:284:TRP:HZ3	1.81	0.45
9:Ff:599:ASP:OD2	9:Ff:602:LEU:HD13	2.16	0.45
7:Fg:18:ARG:O	7:Fg:21:ILE:HG12	2.17	0.45
6:Af:138:GLU:CD	6:Af:138:GLU:C	2.85	0.45
1:Am:12:HIS:CE1	3:Ao:1:ARG:CZ	3.00	0.45
5:Aq:219:ARG:NH1	6:Ax:150:LYS:HG2	2.31	0.45
5:Aq:221:TRP:CE3	5:Aq:231:THR:HB	2.52	0.45
6:Bd:195:ASP:HB2	6:Bj:214:MET:HE1	1.98	0.45
2:Bf:4:MET:SD	2:Bf:5:ILE:HD13	2.57	0.45
6:Bv:123:ASN:HB2	6:Bv:126:GLN:CD	2.42	0.45
6:Bv:200:SER:CB	7:Em:816:GLU:HG2	2.46	0.45
5:Ca:254:ILE:HD12	5:Ca:262:ILE:HG21	1.98	0.45
6:Cb:219:LYS:HB3	6:Cb:222:ASN:OD1	2.17	0.45
6:Ch:107:ILE:H	6:Ch:107:ILE:HD12	1.81	0.45
6:Ch:226:LEU:N	6:Ch:238:MET:HE1	2.32	0.45
6:Cn:124:PRO:O	6:Cn:128:VAL:HG12	2.16	0.45
6:Cn:141:LYS:HE3	7:Eo:812:TRP:CD2	2.52	0.45
6:Df:107:ILE:HD12	6:Df:107:ILE:H	1.82	0.45
6:Dl:118:ASN:ND2	7:Eq:802:LEU:HD11	2.32	0.45
6:Dl:204:ILE:HD11	6:Dl:213:LEU:HD22	2.00	0.45
6:Dl:238:MET:SD	6:Dl:239:LEU:N	2.90	0.45
6:Dr:111:ALA:HB2	7:Er:797:ARG:CZ	2.47	0.45
6:Dx:154:THR:O	6:Dx:252:VAL:HA	2.17	0.45
6:Ed:115:MET:SD	7:Et:801:MET:HE3	2.57	0.45
6:Ed:188:PRO:HB3	6:Ed:211:ASN:ND2	2.32	0.45
7:Ej:779:GLN:O	7:Ej:783:GLU:HG3	2.17	0.45
7:Eu:804:ALA:O	7:Eu:808:LEU:HG	2.17	0.45
8:Ey:90:PRO:HB2	8:Ez:246:ASN:HD21	1.82	0.45
8:Ez:297:ALA:O	8:Ez:301:LEU:HG	2.16	0.45
8:Ez:336:TYR:HA	8:Ez:339:GLN:OE1	2.17	0.45
8:Fa:347:LEU:HA	8:Fa:350:ILE:HD12	1.99	0.45
9:Fb:355:PRO:HD3	9:Fb:375:LYS:HG2	1.99	0.45
9:Fb:546:LYS:HG3	9:Fb:650:GLU:HG3	1.98	0.45
9:Fd:290:LEU:HD21	9:Fd:329:ARG:HG2	1.99	0.45
9:Fd:296:SER:HA	9:Fd:302:VAL:HG23	1.98	0.45
9:Fd:432:PRO:O	9:Fd:436:LEU:HD23	2.17	0.45
9:Fd:516:LEU:HA	9:Fd:519:TRP:CE3	2.52	0.45
9:Fe:534:ILE:HG22	9:Fe:568:TYR:HE1	1.80	0.45
9:Ff:179:LYS:NZ	9:Ff:494:SER:HB2	2.32	0.45

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
9:Ff:724:ILE:O	9:Ff:728:ILE:HG12	2.17	0.45
1:Ag:1:CYS:SG	4:Aj:2:PHE:HB3	2.57	0.44
6:Al:225:ASN:HD21	6:Ar:176:VAL:N	2.15	0.44
5:Bc:264:PHE:HB3	5:Bc:269:SER:HB3	1.99	0.44
6:Bj:132:GLN:NE2	6:Bj:133:ILE:HG23	2.33	0.44
6:Bj:155:SER:HB2	6:Bj:253:ASP:HB2	1.99	0.44
6:Bp:123:ASN:O	6:Bp:126:GLN:HB2	2.17	0.44
6:Cb:194:TYR:CA	6:Cb:229:ARG:HH12	2.30	0.44
1:Ci:16:ALA:HA	4:Cr:3:GLY:HA3	1.99	0.44
6:Ct:236:PRO:HG2	6:Cz:251:ARG:HH11	1.82	0.44
6:Cz:207:ASP:HB3	7:Eq:822:GLU:OE2	2.17	0.44
6:Df:208:LYS:HA	6:Df:208:LYS:HD3	1.90	0.44
6:Dr:248:VAL:HG23	6:Dr:250:TYR:CD1	2.52	0.44
6:Ed:202:PHE:CD1	6:Ed:241:LEU:HD13	2.52	0.44
7:Eh:799:SER:O	7:Eh:803:THR:HG23	2.17	0.44
7:Ev:798:THR:O	7:Ev:799:SER:C	2.59	0.44
8:Ew:211:MET:HB3	8:Ew:287:VAL:O	2.17	0.44
8:Ew:286:GLN:NE2	8:Ew:287:VAL:HG13	2.33	0.44
8:Ew:338:ALA:O	8:Ew:342:VAL:HG23	2.17	0.44
8:Ew:381:ARG:HH12	8:Ew:382:ARG:HD3	1.82	0.44
8:Ex:108:GLY:O	8:Ex:112:ILE:HG12	2.17	0.44
8:Ex:348:TYR:HA	8:Ex:351:LEU:HG	2.00	0.44
8:Ex:440:ASN:O	8:Ex:443:MET:SD	2.75	0.44
8:Ey:205:MET:SD	8:Ey:210:VAL:HB	2.57	0.44
8:Ey:329:TYR:CE1	8:Ey:333:LEU:HD21	2.52	0.44
8:Fa:220:PRO:HD2	8:Fa:228:PRO:O	2.18	0.44
8:Fa:229:ALA:HB1	8:Fa:231:TYR:CE2	2.53	0.44
9:Fb:464:ALA:HB3	9:Fb:761:PHE:HE2	1.82	0.44
9:Fd:54:SER:O	9:Fd:57:MET:HE1	2.17	0.44
9:Fd:179:LYS:HB2	9:Fd:550:LEU:HD13	1.98	0.44
9:Fd:731:ALA:C	9:Fd:733:PRO:HD2	2.42	0.44
9:Fd:912:PHE:C	9:Fd:916:LEU:HD22	2.42	0.44
9:Fe:501:LEU:HB3	9:Fe:527:LEU:HD23	1.99	0.44
9:Fe:714:SER:C	9:Fe:716:LEU:HD23	2.42	0.44
9:Ff:282:VAL:HA	9:Ff:494:SER:HB3	1.98	0.44
7:Fh:34:ILE:HG12	7:Fh:38:LYS:HZ2	1.82	0.44
6:Af:172:SER:HB3	6:Af:248:VAL:HG12	1.97	0.44
6:Bj:205:GLN:OE1	7:Ej:818:GLN:HG2	2.18	0.44
5:Bu:245:LYS:HE3	5:Bu:257:SER:HA	2.00	0.44
6:Bv:203:ASN:O	6:Bv:215:ILE:HD12	2.16	0.44
6:Bv:238:MET:HB3	6:Cb:250:TYR:HD2	1.82	0.44

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
6:Cn:128:VAL:O	6:Cn:132:GLN:HG2	2.17	0.44
6:Cn:229:ARG:HD3	6:Cn:230:LEU:N	2.32	0.44
5:Dk:210:TYR:CD2	5:Dk:238:ILE:HD11	2.52	0.44
6:Ed:190:PRO:O	6:Ed:231:ARG:HG2	2.18	0.44
6:Ed:251:ARG:HE	6:Ed:251:ARG:HB3	1.64	0.44
7:Eh:793:GLU:O	7:Eh:797:ARG:HG3	2.18	0.44
7:Eq:779:GLN:HA	7:Eq:782:ASN:HD22	1.82	0.44
8:Ew:243:LEU:HD11	8:Fa:338:ALA:HB2	2.00	0.44
8:Ew:399:TRP:HE1	8:Ew:414:GLU:HG2	1.82	0.44
8:Ex:177:ALA:O	8:Ex:181:ILE:HG12	2.17	0.44
8:Ex:436:ILE:O	8:Ex:437:LEU:C	2.59	0.44
8:Ey:84:THR:HB	8:Ey:371:GLN:HA	1.98	0.44
8:Ey:277:LEU:HA	8:Ey:280:ILE:HG22	1.99	0.44
8:Ey:414:GLU:O	8:Ey:417:ILE:HG13	2.17	0.44
8:Ey:424:TYR:CZ	8:Ez:383:LEU:HB3	2.53	0.44
8:Ez:331:ASN:HA	8:Ez:334:ARG:HE	1.82	0.44
9:Fb:755:VAL:O	9:Fb:758:TYR:HB3	2.16	0.44
9:Fc:121:TYR:HB3	9:Fc:125:GLN:HE21	1.82	0.44
9:Fc:378:PHE:CD1	9:Fc:414:PHE:HE2	2.36	0.44
9:Fc:381:PRO:HD3	9:Fc:399:TRP:CZ3	2.52	0.44
9:Fc:801:ALA:O	9:Fc:805:LEU:CB	2.65	0.44
9:Fc:818:GLY:C	9:Fc:933:ILE:HG13	2.42	0.44
9:Fd:593:LEU:HD21	9:Fd:905:TRP:HB3	1.99	0.44
9:Fd:670:ILE:HD12	9:Fd:696:TYR:HD2	1.80	0.44
9:Fd:674:GLY:HA2	9:Fd:677:THR:HG22	1.99	0.44
9:Fe:164:THR:HG21	9:Fe:583:PRO:HA	1.99	0.44
9:Fe:350:MET:SD	9:Fe:567:VAL:HG22	2.57	0.44
9:Fe:378:PHE:CD1	9:Fe:414:PHE:HE2	2.35	0.44
9:Ff:824:ALA:O	9:Ff:827:TYR:CB	2.56	0.44
9:Ff:829:GLY:O	9:Ff:832:ILE:HG12	2.16	0.44
9:Ff:831:TRP:CD1	9:Ff:831:TRP:N	2.84	0.44
9:Ff:922:TYR:C	9:Ff:922:TYR:CD1	2.95	0.44
5:Ae:210:TYR:HA	5:Ae:224:GLY:HA2	2.00	0.44
5:Bc:253:ARG:HD3	5:Bc:255:LEU:HD13	1.99	0.44
6:Bj:141:LYS:NZ	7:Ej:812:TRP:CD1	2.85	0.44
6:Bj:220:LEU:HD12	6:Bp:138:GLU:OE1	2.17	0.44
6:Bp:193:ALA:HB3	6:Bp:229:ARG:HD2	2.00	0.44
2:Br:4:MET:HE1	5:Bu:248:ASP:OD1	2.16	0.44
6:Cb:122:LEU:HB3	6:Cb:126:GLN:HB2	2.00	0.44
6:Cb:126:GLN:O	6:Cb:129:LYS:HB2	2.17	0.44
6:Cb:178:SER:O	6:Cb:251:ARG:HA	2.18	0.44

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
5:Cg:246:LEU:HD12	5:Cg:247:ILE:H	1.82	0.44
6:Ch:112:PHE:HA	6:Ch:115:MET:HB2	2.00	0.44
6:Cz:221:TYR:HE1	6:Df:138:GLU:HG3	1.80	0.44
6:Dr:115:MET:O	6:Dr:119:LEU:HG	2.17	0.44
1:Ds:6:LEU:HA	1:Ds:9:LEU:HB3	2.00	0.44
6:Dx:206:TRP:HB2	6:Dx:213:LEU:HD23	2.00	0.44
6:Ed:193:ALA:HB3	6:Ed:229:ARG:HD2	1.99	0.44
6:Ed:196:LEU:HD21	6:Ed:202:PHE:O	2.17	0.44
7:Ef:779:GLN:HA	7:Ef:782:ASN:OD1	2.18	0.44
7:Eq:780:LYS:HA	7:Eq:783:GLU:CD	2.42	0.44
8:Ew:281:ARG:HB3	8:Ew:286:GLN:HE21	1.82	0.44
8:Ex:173:PRO:HB2	8:Ex:348:TYR:CE1	2.51	0.44
8:Ex:334:ARG:HD3	8:Ey:282:TYR:HE1	1.82	0.44
8:Ex:420:ALA:O	8:Ex:424:TYR:HD1	2.01	0.44
8:Ez:63:PRO:HA	8:Fa:48:LYS:HZ1	1.82	0.44
8:Ez:188:SER:O	8:Ez:191:MET:HB3	2.17	0.44
8:Ez:238:PRO:HB3	8:Ez:262:GLY:HA3	1.99	0.44
8:Fa:195:GLN:C	8:Fa:196:LYS:HD3	2.41	0.44
8:Fa:211:MET:HB3	8:Fa:287:VAL:O	2.17	0.44
8:Fa:297:ALA:O	8:Fa:301:LEU:HG	2.17	0.44
9:Fb:434:LEU:O	9:Fb:438:ARG:HG3	2.17	0.44
9:Fc:33:LEU:HA	9:Fc:36:VAL:CG1	2.48	0.44
9:Fc:220:PRO:O	9:Fc:224:THR:HG23	2.17	0.44
9:Fc:837:PHE:CE2	9:Fc:918:TYR:HB2	2.52	0.44
9:Fd:84:THR:HA	9:Fd:87:THR:HG22	1.99	0.44
9:Fd:212:PRO:HB3	9:Fd:215:GLY:O	2.17	0.44
9:Fd:426:TYR:HA	9:Fd:429:ILE:HG22	2.00	0.44
9:Fd:579:LEU:HB3	9:Fd:580:PRO:HD2	1.99	0.44
9:Fd:708:LEU:HA	9:Fe:730:MET:SD	2.58	0.44
9:Fe:167:LEU:HD12	9:Fe:585:ILE:HD11	1.97	0.44
9:Fe:931:THR:O	9:Fe:935:HIS:HB3	2.18	0.44
9:Ff:242:LYS:HE3	9:Ff:265:ASN:HD22	1.82	0.44
9:Ff:266:PHE:HB3	9:Ff:270:SER:OG	2.17	0.44
9:Ff:776:MET:HE2	9:Ff:776:MET:C	2.42	0.44
7:Fh:35:GLY:HA2	7:Fh:38:LYS:NZ	2.32	0.44
6:Al:131:LYS:O	6:Al:135:GLU:HG3	2.17	0.44
6:Ax:184:SER:HA	6:Ax:255:ARG:NH1	2.32	0.44
5:Ca:218:GLY:HA3	6:Ch:148:PRO:HD3	2.00	0.44
5:Ca:219:ARG:HH12	5:Ca:233:ARG:HD3	1.82	0.44
6:Cn:230:LEU:HD23	6:Cn:233:LEU:HD12	1.99	0.44
6:Ct:238:MET:CE	6:Cz:251:ARG:HG3	2.45	0.44

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
5:De:212:ILE:HG23	5:De:222:LEU:HD22	1.98	0.44
6:Df:128:VAL:O	6:Df:132:GLN:HG3	2.18	0.44
2:Dh:2:VAL:HG22	3:Di:1:ARG:NE	2.32	0.44
3:Di:7:THR:HB	3:Di:9:TYR:CE2	2.53	0.44
6:Dr:238:MET:CE	6:Dx:251:ARG:HB3	2.43	0.44
7:Es:773:GLN:HG3	9:Fb:580:PRO:HB2	2.00	0.44
8:Ew:348:TYR:HD1	8:Ew:351:LEU:HD23	1.83	0.44
8:Ew:359:MET:HE3	8:Ew:367:ASN:C	2.42	0.44
8:Ex:50:LEU:HB3	8:Ex:52:TYR:HE2	1.82	0.44
8:Ex:84:THR:O	8:Ex:88:ALA:CB	2.65	0.44
8:Ey:104:ASP:HA	8:Ey:109:ASN:HD22	1.81	0.44
8:Ey:408:PRO:HA	8:Ey:411:VAL:HG22	1.98	0.44
8:Ey:431:GLN:HA	8:Ey:434:GLU:OE1	2.17	0.44
8:Ez:105:LYS:HE2	8:Fa:368:ILE:N	2.29	0.44
8:Fa:382:ARG:HB3	8:Fa:399:TRP:HE3	1.76	0.44
9:Fb:267:ASP:OD1	9:Fb:267:ASP:C	2.61	0.44
9:Fb:354:ASP:OD2	9:Fb:373:ILE:HD12	2.17	0.44
9:Fb:615:VAL:O	9:Fb:621:SER:HA	2.17	0.44
9:Fc:261:LEU:HB2	9:Fc:284:TRP:CH2	2.52	0.44
9:Fc:429:ILE:C	9:Fc:430:MET:HE2	2.42	0.44
9:Fc:788:HIS:ND1	9:Fc:790:GLU:HG3	2.32	0.44
9:Fc:936:LEU:HA	9:Fc:939:LYS:HD2	1.99	0.44
9:Fd:100:SER:HB3	7:Fk:16:ARG:HH12	1.82	0.44
9:Fe:57:MET:HE1	9:Ff:824:ALA:CB	2.47	0.44
9:Fe:200:ALA:O	9:Fe:204:LEU:HD23	2.17	0.44
9:Fe:448:LYS:HA	9:Fe:451:ASP:OD2	2.17	0.44
9:Fe:659:PHE:CE1	9:Fe:707:LEU:HD22	2.53	0.44
9:Fe:667:MET:HG3	9:Fe:911:PHE:CE2	2.53	0.44
9:Ff:912:PHE:O	9:Ff:916:LEU:HG	2.18	0.44
6:Af:115:MET:HE1	7:Eu:802:LEU:HA	2.00	0.44
6:Af:176:VAL:H	6:Ed:225:ASN:HD21	1.66	0.44
6:Af:200:SER:H	7:Ef:816:GLU:HG3	1.82	0.44
1:Ag:12:HIS:CE1	3:Ai:1:ARG:CZ	3.00	0.44
6:Al:171:LEU:HD23	6:Al:202:PHE:CE2	2.53	0.44
6:Bd:156:GLN:HE22	6:Bd:158:VAL:HG22	1.82	0.44
3:Bm:7:THR:HB	3:Bm:9:TYR:CE2	2.53	0.44
2:Br:1:CYS:HA	3:Bs:4:ILE:HD13	2.00	0.44
5:Cm:250:LEU:HD12	5:Cm:251:GLN:HG3	2.00	0.44
5:Cs:216:ILE:C	5:Cs:216:ILE:HD12	2.43	0.44
2:Cv:2:VAL:HG22	3:Cw:1:ARG:NE	2.31	0.44
5:Cy:234:GLU:HA	5:Cy:247:ILE:HD11	1.99	0.44

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:Da:9:LEU:HB2	2:Db:2:VAL:HG21	1.99	0.44
5:Dk:216:ILE:HD11	6:Dr:148:PRO:HG2	1.99	0.44
5:Dk:256:THR:HG22	5:Dk:260:GLN:O	2.18	0.44
5:Ec:251:GLN:HB2	5:Ec:253:ARG:HG2	2.00	0.44
6:Ed:155:SER:HB2	6:Ed:253:ASP:OD2	2.18	0.44
7:Es:810:GLN:HA	7:Es:813:LYS:HE3	1.99	0.44
8:Ew:429:ASP:HA	8:Ew:432:ILE:HG12	2.00	0.44
8:Ey:294:LYS:O	8:Ey:298:TYR:HB3	2.16	0.44
8:Ez:118:THR:HA	8:Ez:121:LYS:HG2	1.99	0.44
8:Fa:216:GLY:HA3	8:Fa:265:SER:O	2.17	0.44
9:Fb:334:ILE:HD12	9:Fb:334:ILE:HA	1.84	0.44
9:Fb:528:ILE:HA	9:Fb:531:GLN:HG2	1.99	0.44
9:Fb:800:PHE:O	9:Fb:804:ILE:HG22	2.17	0.44
9:Fd:115:ILE:HG12	9:Fd:123:MET:HE1	1.98	0.44
9:Fd:751:TYR:CE2	9:Fe:910:ALA:HA	2.53	0.44
9:Fd:915:ILE:O	9:Fd:919:THR:HG23	2.17	0.44
9:Ff:128:PHE:CD2	9:Ff:779:ALA:HB2	2.52	0.44
9:Ff:651:MET:O	9:Ff:655:ILE:HG12	2.17	0.44
6:Af:129:LYS:HE3	6:Al:112:PHE:CD1	2.53	0.44
2:An:2:VAL:HA	3:Ao:1:ARG:HH21	1.82	0.44
6:Ar:171:LEU:HD11	6:Ar:241:LEU:HD12	2.00	0.44
6:Ax:121:PRO:HB2	7:Eg:801:MET:CE	2.47	0.44
1:Ay:4:ALA:HB2	3:Ba:14:TYR:HB2	2.00	0.44
5:Bc:219:ARG:HG2	6:Bj:148:PRO:HD2	1.98	0.44
5:Bi:251:GLN:HB3	5:Bi:253:ARG:NH2	2.32	0.44
5:Bu:218:GLY:HA3	6:Cb:148:PRO:HD3	1.99	0.44
6:Cb:156:GLN:NE2	6:Cb:167:PRO:HG3	2.33	0.44
6:Ch:123:ASN:CG	6:Ch:124:PRO:HD2	2.42	0.44
1:Cu:15:ASN:HB3	2:Db:9:ARG:NH1	2.32	0.44
2:Cv:6:GLY:HA3	6:Dl:125:GLU:HB3	1.99	0.44
6:Cz:170:ARG:HD2	6:Cz:247:ALA:CB	2.47	0.44
6:Df:130:LEU:HD23	7:Eq:808:LEU:HD12	1.99	0.44
7:Eg:808:LEU:O	7:Eg:812:TRP:CD1	2.70	0.44
7:Eo:804:ALA:O	7:Eo:808:LEU:HG	2.17	0.44
7:Er:800:ASP:OD1	7:Er:800:ASP:C	2.61	0.44
8:Ew:199:LEU:HD12	8:Ew:199:LEU:H	1.81	0.44
8:Ew:246:ASN:ND2	8:Fa:91:VAL:HA	2.33	0.44
8:Ew:418:LEU:HD12	8:Ew:419:LEU:N	2.32	0.44
8:Ex:330:PHE:HB2	8:Ex:334:ARG:NH1	2.33	0.44
8:Ex:430:ARG:CZ	8:Ey:428:LEU:HB2	2.48	0.44
8:Ey:92:ASP:OD2	8:Ez:244:ASN:HB3	2.18	0.44

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
8:Ez:102:VAL:HA	8:Fa:253:LEU:HD21	2.00	0.44
8:Ez:298:TYR:O	8:Ez:301:LEU:HD12	2.17	0.44
8:Fa:411:VAL:O	8:Fa:415:ILE:HG13	2.17	0.44
9:Fb:205:TYR:HA	9:Fb:208:GLN:NE2	2.31	0.44
9:Fb:264:PRO:HD3	9:Fb:278:ILE:HG22	2.00	0.44
9:Fb:434:LEU:HG	9:Fb:438:ARG:HE	1.82	0.44
9:Fb:707:LEU:HA	9:Fb:710:MET:SD	2.58	0.44
9:Fd:205:TYR:HB3	9:Fd:214:CYS:SG	2.58	0.44
9:Fd:299:LYS:HG2	9:Fd:299:LYS:H	1.61	0.44
9:Fe:196:LYS:HE3	9:Fe:357:PHE:CD1	2.53	0.44
9:Fe:527:LEU:O	9:Fe:531:GLN:HG2	2.18	0.44
9:Fe:932:LEU:HB3	9:Fe:936:LEU:HD23	1.99	0.44
9:Ff:147:LEU:HA	9:Ff:150:LEU:HD23	2.00	0.44
9:Ff:936:LEU:O	9:Ff:939:LYS:HG2	2.17	0.44
6:Ar:205:GLN:HB2	7:Eg:820:TYR:CE2	2.51	0.44
2:Az:4:MET:HE2	5:Bc:250:LEU:HB2	1.99	0.44
5:Bi:247:ILE:N	5:Bi:247:ILE:HD12	2.33	0.44
5:Ca:253:ARG:HG3	5:Ca:263:LYS:CD	2.48	0.44
6:Cb:107:ILE:HA	6:Cb:110:LYS:NZ	2.33	0.44
6:Cn:160:LEU:HD11	6:Cn:237:VAL:HG22	1.99	0.44
6:Cn:196:LEU:HD12	6:Cn:204:ILE:HG12	1.99	0.44
5:Cs:245:LYS:HZ1	5:Cs:257:SER:HB3	1.83	0.44
6:Df:193:ALA:HB1	7:Es:820:TYR:HE2	1.81	0.44
6:Dl:124:PRO:O	6:Dl:128:VAL:HG12	2.18	0.44
6:Dr:113:LYS:HD3	6:Dr:113:LYS:HA	1.64	0.44
2:Dt:4:MET:SD	2:Dt:4:MET:O	2.76	0.44
6:Ed:126:GLN:HA	6:Ed:129:LYS:HG2	2.00	0.44
7:Ei:779:GLN:HA	7:Ei:782:ASN:HB3	1.98	0.44
8:Ew:139:ASN:OD1	8:Ew:141:LEU:HG	2.17	0.44
8:Ew:298:TYR:CZ	8:Ex:230:PHE:HE2	2.36	0.44
8:Ew:404:ASN:OD1	8:Fa:67:GLN:HB3	2.18	0.44
8:Ey:144:GLN:NE2	8:Ey:205:MET:HG3	2.33	0.44
8:Ey:318:GLN:HA	8:Ey:321:GLN:HB3	1.98	0.44
8:Ez:353:LYS:HD2	8:Ez:371:GLN:HG2	2.00	0.44
8:Fa:77:VAL:O	8:Fa:80:TYR:HB2	2.17	0.44
8:Fa:268:THR:HB	8:Fa:270:LYS:NZ	2.33	0.44
9:Fb:546:LYS:HG3	9:Fb:650:GLU:CD	2.42	0.44
9:Fc:69:LEU:HD21	9:Fc:114:LEU:HB3	1.98	0.44
9:Fc:124:MET:HA	9:Fc:127:PHE:CE1	2.52	0.44
9:Fc:178:ALA:HB3	9:Fc:179:LYS:NZ	2.33	0.44
9:Fc:240:VAL:HG12	9:Fc:333:ALA:HB1	1.99	0.44

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
9:Fd:148:SER:O	9:Fd:152:ARG:HG2	2.17	0.44
9:Fd:325:LEU:HG	9:Fd:329:ARG:CZ	2.48	0.44
9:Fd:414:PHE:HB3	9:Fd:418:GLU:OE1	2.18	0.44
9:Fd:725:PHE:HA	9:Fd:728:ILE:HG13	1.99	0.44
9:Fe:143:TRP:CE3	9:Fe:147:LEU:HD21	2.52	0.44
9:Fe:370:PHE:CE2	9:Fe:562:PRO:HA	2.52	0.44
9:Fe:559:VAL:HG21	9:Fe:563:LEU:HB3	2.00	0.44
9:Ff:57:MET:HA	9:Ff:60:MET:HE3	2.00	0.44
9:Ff:72:GLY:HA3	9:Ff:784:LEU:HD21	2.00	0.44
9:Ff:612:CYS:HB2	9:Ff:626:ARG:NE	2.30	0.44
2:Ab:4:MET:CE	5:Ae:250:LEU:HG	2.48	0.44
6:Al:112:PHE:CD2	6:Al:113:LYS:HD3	2.52	0.44
1:Am:16:ALA:HA	4:Av:3:GLY:HA3	2.00	0.44
2:An:4:MET:SD	5:Aq:248:ASP:HA	2.57	0.44
5:Aq:213:GLN:O	6:Ar:170:ARG:HD3	2.18	0.44
5:Aq:219:ARG:HG2	6:Ax:148:PRO:HG2	1.99	0.44
6:Ax:106:VAL:HA	6:Ax:109:LYS:HG2	2.00	0.44
6:Bd:208:LYS:HZ3	7:Ej:821:THR:HG22	1.83	0.44
6:Bp:196:LEU:HD22	6:Bp:199:PRO:CA	2.48	0.44
6:Ch:145:PRO:HB3	6:Cn:132:GLN:OE1	2.17	0.44
6:Cz:178:SER:OG	6:Cz:212:THR:HG23	2.18	0.44
2:Dh:5:ILE:HG12	6:Dx:124:PRO:HB3	1.99	0.44
6:Di:134:TYR:O	6:Di:138:GLU:HG3	2.17	0.44
6:Di:179:LEU:HB3	6:Di:181:PHE:HE1	1.80	0.44
6:Di:189:TRP:CE2	6:Di:260:GLY:HA2	2.53	0.44
6:Dr:115:MET:SD	7:Er:801:MET:HG2	2.58	0.44
6:Dx:189:TRP:CG	6:Dx:230:LEU:HD13	2.52	0.44
7:Er:814:GLN:N	7:Er:814:GLN:CD	2.76	0.44
7:Et:789:LYS:HE3	9:Ff:267:ASP:OD2	2.18	0.44
7:Et:789:LYS:HZ2	7:Et:792:GLN:HE22	1.65	0.44
8:Ew:253:LEU:HD22	8:Ew:373:LEU:HD12	1.98	0.44
8:Ew:419:LEU:HD13	8:Ex:419:LEU:HD21	1.99	0.44
8:Ey:246:ASN:HD21	8:Ey:250:ALA:HB3	1.83	0.44
8:Ez:277:LEU:HD21	8:Ez:281:ARG:NH2	2.32	0.44
9:Fb:65:ASN:O	9:Fb:69:LEU:HG	2.18	0.44
9:Fc:546:LYS:HD3	9:Fc:546:LYS:N	2.33	0.44
9:Fc:782:VAL:O	9:Fc:786:VAL:HG23	2.17	0.44
9:Fd:697:ILE:HD12	9:Fe:909:TYR:CD1	2.40	0.44
9:Fe:77:MET:HE1	7:Fj:32:VAL:HG11	1.99	0.44
9:Fe:632:PHE:CZ	9:Ff:628:MET:HB3	2.51	0.44
9:Fe:807:ASN:HB2	9:Fe:941:LEU:HD21	1.99	0.44

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
9:Fe:926:VAL:HG13	9:Fe:930:PHE:CE2	2.50	0.44
7:Fh:36:PHE:O	7:Fh:40:ARG:HG2	2.18	0.44
6:Af:196:LEU:HA	6:Af:226:LEU:HD13	2.00	0.44
6:Al:225:ASN:HD21	6:Ar:176:VAL:H	1.64	0.44
6:Ar:135:GLU:HA	6:Ar:138:GLU:OE2	2.17	0.44
6:Ax:182:LEU:HD11	6:Ax:186:GLY:HA2	1.99	0.44
3:Ba:13:LYS:HA	3:Ba:13:LYS:HD2	1.70	0.44
6:Bd:225:ASN:HB3	6:Bj:175:PHE:CD2	2.53	0.44
5:Bi:248:ASP:HB3	5:Bi:253:ARG:HG2	1.99	0.44
6:Bj:230:LEU:HB3	6:Bj:233:LEU:CD1	2.47	0.44
6:Bp:106:VAL:C	6:Bp:110:LYS:HZ1	2.26	0.44
6:Ch:128:VAL:O	6:Ch:132:GLN:HG2	2.16	0.44
6:Ch:186:GLY:HA2	6:Ch:255:ARG:NH2	2.33	0.44
1:Ci:12:HIS:NE2	3:Ck:9:TYR:HB2	2.33	0.44
6:Cz:202:PHE:HA	6:Cz:216:GLN:O	2.18	0.44
5:Dk:233:ARG:HG3	5:Dk:234:GLU:OE1	2.18	0.44
5:Dk:264:PHE:HD2	5:Dk:265:SER:N	2.13	0.44
6:Dl:202:PHE:HB3	6:Dl:215:ILE:HG23	2.00	0.44
6:Dl:238:MET:N	6:Dr:251:ARG:HH12	2.15	0.44
6:Dr:173:GLN:HE21	7:Et:815:VAL:CG2	2.24	0.44
6:Dx:105:GLU:OE1	6:Dx:105:GLU:N	2.51	0.44
6:Ed:205:GLN:OE1	6:Ed:214:MET:HG3	2.17	0.44
7:Eh:792:GLN:O	7:Eh:795:GLN:HG3	2.17	0.44
7:Ej:812:TRP:CD1	7:Ej:812:TRP:N	2.85	0.44
7:Em:808:LEU:HA	7:Em:811:ASP:OD2	2.18	0.44
7:Eq:809:VAL:O	7:Eq:813:LYS:HG3	2.17	0.44
7:Er:822:GLU:N	7:Er:822:GLU:OE2	2.51	0.44
7:Eu:776:ALA:HA	7:Eu:779:GLN:NE2	2.33	0.44
8:Ex:243:LEU:HD21	8:Ex:282:TYR:CD1	2.52	0.44
8:Ey:55:THR:HA	8:Ez:42:TYR:CZ	2.53	0.44
8:Ey:349:TYR:HE1	8:Ey:431:GLN:NE2	2.15	0.44
8:Ez:253:LEU:C	8:Ez:254:MET:HE2	2.42	0.44
8:Ez:263:GLN:H	8:Ez:268:THR:CG2	2.30	0.44
8:Ez:397:GLN:HB3	8:Ez:401:LYS:HE2	1.99	0.44
8:Fa:36:THR:HA	8:Fa:39:LEU:CD2	2.46	0.44
9:Fb:243:GLN:HG2	9:Fb:333:ALA:HB2	2.00	0.44
9:Fb:647:ILE:O	9:Fb:651:MET:HG2	2.18	0.44
9:Fb:725:PHE:O	9:Fb:729:MET:HG3	2.18	0.44
9:Fc:124:MET:HE2	9:Fc:124:MET:HB2	1.89	0.44
9:Fd:615:VAL:HG11	9:Fd:624:LEU:HD12	2.00	0.44
9:Fe:60:MET:HA	9:Fe:63:VAL:HG22	1.99	0.44

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
9:Fe:64:PHE:O	9:Fe:68:VAL:HG13	2.18	0.44
9:Fe:461:TRP:CZ3	9:Fe:462:ILE:HG22	2.53	0.44
9:Ff:101:ILE:O	9:Ff:104:PRO:HD2	2.18	0.44
9:Ff:186:GLY:O	9:Ff:189:ILE:HG22	2.18	0.44
9:Ff:534:ILE:HG22	9:Ff:568:TYR:CD2	2.53	0.44
9:Ff:575:MET:HE2	9:Ff:575:MET:C	2.43	0.44
9:Ff:814:LEU:HD12	9:Ff:937:PRO:HB3	2.00	0.44
6:Af:126:GLN:CA	6:Af:129:LYS:HE2	2.40	0.43
6:Af:215:ILE:HD13	6:Af:215:ILE:HA	1.88	0.43
6:Al:166:PRO:HG3	6:Ar:250:TYR:CD2	2.53	0.43
6:Ar:138:GLU:C	6:Ar:141:LYS:HB3	2.42	0.43
5:Aw:219:ARG:HA	5:Aw:232:VAL:O	2.19	0.43
6:Bp:149:PRO:HA	6:Bp:246:LYS:O	2.18	0.43
5:Cg:213:GLN:OE1	5:Cg:213:GLN:HA	2.18	0.43
6:Ch:129:LYS:O	6:Ch:133:ILE:HG22	2.17	0.43
6:Ch:171:LEU:HD21	6:Ch:241:LEU:HB3	2.00	0.43
6:Cn:234:ASN:ND2	6:Ct:182:LEU:HD21	2.31	0.43
5:Cy:246:LEU:HB2	5:Cy:255:LEU:HD23	1.99	0.43
6:Cz:140:ALA:HB1	6:Df:127:VAL:HB	2.00	0.43
6:Cz:229:ARG:HH12	6:Df:212:THR:HG21	1.82	0.43
6:Df:112:PHE:O	6:Df:115:MET:HB2	2.16	0.43
6:Dr:181:PHE:HD1	6:Dr:230:LEU:HD11	1.83	0.43
6:Dr:195:ASP:O	6:Dr:226:LEU:HD12	2.16	0.43
6:Ed:171:LEU:HD21	6:Ed:215:ILE:HG21	1.99	0.43
7:En:797:ARG:NH1	7:En:801:MET:HE1	2.33	0.43
8:Ew:343:GLY:O	8:Ew:347:LEU:HG	2.18	0.43
8:Ex:220:PRO:HD2	8:Ex:228:PRO:O	2.18	0.43
8:Ey:230:PHE:CD1	8:Ey:231:TYR:HD2	2.36	0.43
8:Ey:413:LYS:NZ	8:Ez:404:ASN:HA	2.15	0.43
9:Fc:437:ILE:HD12	9:Fc:437:ILE:HA	1.88	0.43
9:Fc:725:PHE:HE1	9:Fd:729:MET:HG2	1.83	0.43
9:Fc:729:MET:O	9:Fc:732:MET:HB3	2.18	0.43
9:Fc:753:ILE:HG13	9:Fc:922:TYR:CZ	2.53	0.43
9:Fd:176:GLY:HA3	9:Fd:492:ASP:HB2	2.00	0.43
9:Fd:337:GLN:O	9:Fd:341:VAL:HG12	2.18	0.43
9:Fd:468:PHE:CD1	9:Fd:757:PRO:HG2	2.53	0.43
9:Fd:813:SER:HA	9:Fd:816:ILE:HG22	2.00	0.43
9:Fe:85:MET:C	9:Fe:85:MET:HE2	2.43	0.43
9:Fe:196:LYS:HE3	9:Fe:357:PHE:HD1	1.82	0.43
9:Fe:196:LYS:HD2	9:Fe:357:PHE:O	2.18	0.43
9:Fe:620:PHE:CZ	9:Ff:615:VAL:HA	2.53	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
9:Fe:644:PHE:CE1	9:Fe:718:PRO:HB2	2.53	0.43
9:Ff:35:VAL:HB	9:Ff:49:LEU:HD21	2.00	0.43
6:Af:130:LEU:HA	6:Af:133:ILE:CG2	2.48	0.43
2:Bf:4:MET:HE2	4:Bh:2:PHE:CE1	2.53	0.43
6:Cb:229:ARG:HB3	6:Cb:236:PRO:HB3	1.99	0.43
6:Ch:117:ARG:O	6:Ch:121:PRO:HA	2.18	0.43
6:Ct:229:ARG:HD2	6:Ct:236:PRO:HB3	2.00	0.43
6:Ct:242:ILE:HG13	6:Ct:245:GLN:HE22	1.79	0.43
6:Cz:242:ILE:HD12	6:Cz:243:PRO:HD2	2.00	0.43
6:Dr:108:ASP:O	6:Dr:112:PHE:CD1	2.70	0.43
2:Dt:5:ILE:HD13	6:Ed:140:ALA:HB2	2.01	0.43
6:Dx:117:ARG:NE	7:Et:797:ARG:HG3	2.32	0.43
1:Dy:11:TYR:CD2	3:Ea:9:TYR:HB3	2.53	0.43
7:Ei:793:GLU:HA	7:Ei:796:GLN:HG3	2.00	0.43
7:Es:788:GLN:HB2	7:Es:789:LYS:HD2	2.01	0.43
8:Ey:78:GLN:HE22	8:Ez:384:PHE:HB2	1.83	0.43
8:Ey:424:TYR:O	8:Ey:427:TYR:HB3	2.19	0.43
8:Ez:211:MET:C	8:Ez:211:MET:HE2	2.43	0.43
8:Fa:271:SER:O	8:Fa:275:GLN:HG3	2.18	0.43
8:Fa:451:ALA:HA	9:Fe:619:PHE:CE2	2.53	0.43
9:Fc:164:THR:HA	9:Fc:167:LEU:HG	2.01	0.43
9:Fc:612:CYS:HB2	9:Fc:623:CYS:HB2	1.42	0.43
9:Fd:68:VAL:HG21	9:Fd:780:PRO:HB2	2.00	0.43
9:Fd:913:PHE:O	9:Fd:917:ILE:HG22	2.18	0.43
9:Fe:469:PHE:CE2	9:Ff:843:TYR:HB3	2.53	0.43
9:Ff:65:ASN:O	9:Ff:69:LEU:HG	2.18	0.43
5:Ae:213:GLN:HG2	5:Ae:221:TRP:O	2.18	0.43
6:Af:189:TRP:CE2	6:Af:260:GLY:HA2	2.53	0.43
6:Al:129:LYS:HG3	6:Ar:112:PHE:HZ	1.82	0.43
6:Bj:191:ILE:HB	6:Bj:206:TRP:CZ2	2.53	0.43
6:Bp:115:MET:N	6:Bp:115:MET:SD	2.90	0.43
6:Bp:134:TYR:HA	6:Bv:120:TYR:CE2	2.52	0.43
6:Bp:145:PRO:HB3	6:Bv:132:GLN:HE22	1.84	0.43
6:Bp:238:MET:CE	6:Bv:178:SER:HB3	2.42	0.43
3:Bs:7:THR:HB	3:Bs:9:TYR:CE2	2.53	0.43
5:Bu:216:ILE:HD13	6:Bv:223:TYR:CE2	2.53	0.43
2:Bx:6:GLY:CA	6:Cn:125:GLU:HG3	2.46	0.43
6:Ch:115:MET:CE	7:El:798:THR:HA	2.48	0.43
6:Ch:186:GLY:HA2	6:Ch:255:ARG:HH22	1.82	0.43
5:Cm:232:VAL:HG23	5:Cm:236:SER:OG	2.18	0.43
2:Cp:6:GLY:HA2	6:Df:125:GLU:HB2	2.00	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
6:Dl:205:GLN:OE1	7:Es:820:TYR:HB2	2.18	0.43
2:Dn:2:VAL:HA	3:Do:1:ARG:NH2	2.33	0.43
3:Do:13:LYS:HD3	3:Do:13:LYS:HA	1.69	0.43
5:Dw:219:ARG:HH11	5:Dw:233:ARG:HE	1.65	0.43
6:Dx:122:LEU:HB3	6:Dx:127:VAL:CG2	2.48	0.43
7:Ef:781:GLN:HG3	9:Fe:337:GLN:CG	2.48	0.43
7:Eh:777:ILE:HG21	9:Fe:580:PRO:HB3	2.01	0.43
7:Ei:814:GLN:HE22	7:Ei:816:GLU:HB3	1.83	0.43
7:Ej:797:ARG:HA	7:Ej:800:ASP:OD2	2.18	0.43
7:Eu:776:ALA:HA	7:Eu:779:GLN:CD	2.43	0.43
8:Ew:59:LYS:H	8:Ew:59:LYS:HD2	1.82	0.43
8:Ew:208:ASN:ND2	8:Fa:324:ALA:HB2	2.30	0.43
8:Ew:332:ASN:OD1	8:Ew:332:ASN:C	2.61	0.43
8:Ew:379:ALA:O	8:Fa:424:TYR:CE2	2.70	0.43
8:Ex:194:GLU:HG2	8:Ex:196:LYS:NZ	2.33	0.43
8:Ey:348:TYR:CD1	8:Ey:351:LEU:HD23	2.53	0.43
8:Ez:260:GLN:O	8:Ez:270:LYS:HD3	2.18	0.43
8:Ez:308:LYS:HB2	8:Ez:311:SER:HB2	2.00	0.43
8:Fa:429:ASP:O	8:Fa:432:ILE:HG13	2.19	0.43
9:Fb:434:LEU:HG	9:Fb:438:ARG:NH2	2.33	0.43
9:Fc:57:MET:HE1	9:Fd:824:ALA:CB	2.47	0.43
9:Fe:143:TRP:CZ2	9:Fe:456:ALA:HB1	2.53	0.43
9:Fe:278:ILE:HD12	9:Fe:278:ILE:HA	1.88	0.43
9:Fe:502:THR:HA	9:Fe:505:PHE:HD2	1.83	0.43
9:Fe:675:VAL:HA	9:Fe:678:LEU:HG	2.00	0.43
5:Aq:245:LYS:C	5:Aq:246:LEU:HD23	2.43	0.43
2:Az:5:ILE:HD13	2:Az:5:ILE:N	2.32	0.43
6:Cb:207:ASP:OD2	6:Cb:210:SER:HB3	2.18	0.43
6:Ch:122:LEU:HD23	7:Em:801:MET:HE3	1.99	0.43
3:Cq:2:VAL:HG21	6:Df:129:LYS:HE3	1.99	0.43
6:Cz:233:LEU:HD11	6:Cz:235:THR:HB	1.99	0.43
6:Dl:193:ALA:HB1	7:Et:820:TYR:HE2	1.82	0.43
6:Dl:202:PHE:HB3	6:Dl:215:ILE:CG2	2.48	0.43
6:Dr:173:GLN:NE2	7:Et:815:VAL:HG21	2.22	0.43
6:Dr:181:PHE:CG	6:Dr:254:LEU:HD11	2.53	0.43
6:Dr:200:SER:HB3	7:Eu:816:GLU:OE1	2.17	0.43
7:Ep:795:GLN:O	7:Ep:798:THR:HB	2.17	0.43
8:Ew:298:TYR:CE2	8:Ex:221:ALA:HA	2.54	0.43
8:Ew:384:PHE:HD2	8:Fa:82:TYR:CD1	2.37	0.43
8:Ew:426:MET:HE1	8:Ex:425:GLN:HB2	2.00	0.43
8:Ex:76:LEU:HA	8:Ex:79:ASN:HD22	1.83	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
8:Ex:427:TYR:CD2	8:Ey:379:ALA:HB2	2.53	0.43
8:Ey:40:VAL:HG22	8:Ez:38:ASN:OD1	2.18	0.43
9:Fb:716:LEU:HA	9:Fc:723:PHE:CE2	2.53	0.43
9:Fc:261:LEU:HB3	9:Fc:282:VAL:HG13	1.99	0.43
9:Fd:428:GLY:O	9:Fd:431:MET:HG3	2.19	0.43
9:Fd:641:TYR:CZ	9:Fd:645:LEU:HD21	2.53	0.43
9:Fe:33:LEU:O	9:Fe:37:PHE:HB2	2.17	0.43
9:Ff:57:MET:O	9:Ff:60:MET:HG2	2.17	0.43
9:Ff:274:PHE:HZ	9:Ff:519:TRP:CZ2	2.36	0.43
9:Ff:378:PHE:CE2	9:Ff:565:SER:HA	2.54	0.43
5:Aq:222:LEU:HD23	5:Aq:222:LEU:H	1.83	0.43
5:Aq:261:VAL:HB	5:Aq:263:LYS:HE3	2.00	0.43
3:Au:4:ILE:HG12	6:Bd:146:GLY:N	2.34	0.43
5:Bo:210:TYR:HA	5:Bo:224:GLY:HA2	2.01	0.43
6:Bv:108:ASP:CA	7:Ej:797:ARG:HH22	2.18	0.43
5:Ca:246:LEU:HD11	5:Ca:253:ARG:HH21	1.83	0.43
5:Ca:253:ARG:HE	5:Ca:255:LEU:CD2	2.31	0.43
1:Ci:12:HIS:CD2	3:Ck:9:TYR:HB2	2.54	0.43
5:Dq:241:TYR:HB3	5:Dq:256:THR:HG21	2.00	0.43
7:En:792:GLN:H	7:En:792:GLN:CD	2.27	0.43
7:Ev:792:GLN:HA	7:Ev:795:GLN:CD	2.43	0.43
8:Ew:43:LEU:O	8:Ew:46:LEU:HG	2.18	0.43
8:Ew:198:TRP:CZ3	8:Fa:317:VAL:HA	2.54	0.43
8:Ew:305:ALA:HA	8:Ew:319:GLN:HG2	1.99	0.43
8:Ew:399:TRP:O	8:Ew:403:ILE:HG12	2.18	0.43
8:Ex:43:LEU:O	8:Ex:46:LEU:HG	2.19	0.43
8:Ey:69:PHE:CZ	8:Ey:402:GLN:HB3	2.53	0.43
8:Ey:245:SER:HA	8:Ey:279:PHE:CZ	2.53	0.43
8:Ey:424:TYR:O	8:Ey:428:LEU:HD22	2.18	0.43
8:Ez:80:TYR:C	8:Ez:374:ASN:HD21	2.26	0.43
8:Fa:134:GLY:N	8:Fa:258:ALA:HB2	2.34	0.43
9:Fb:194:LEU:O	9:Fb:198:LEU:HD22	2.18	0.43
9:Fb:367:LYS:HD2	9:Fb:367:LYS:C	2.43	0.43
9:Fb:378:PHE:HB2	9:Fb:570:PHE:CE2	2.53	0.43
9:Fc:429:ILE:HB	9:Fc:430:MET:HE2	2.01	0.43
9:Fd:617:ILE:O	9:Fd:618:LEU:C	2.60	0.43
9:Fd:762:THR:HG23	9:Fd:766:PHE:HE2	1.83	0.43
9:Fd:781:ILE:H	9:Fd:781:ILE:HG13	1.45	0.43
9:Fe:555:GLN:CD	9:Fe:588:LEU:HD12	2.42	0.43
9:Ff:197:GLN:OE1	9:Ff:520:PHE:HZ	2.02	0.43
9:Ff:348:GLN:HA	9:Ff:351:VAL:HG12	2.01	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
9:Ff:470:ASP:O	9:Ff:473:LYS:HG2	2.17	0.43
9:Ff:486:ASP:OD1	9:Ff:486:ASP:C	2.62	0.43
6:Bd:129:LYS:O	6:Bd:132:GLN:HG3	2.19	0.43
1:Bk:3:ASP:O	1:Bk:6:LEU:HD12	2.18	0.43
6:Bv:171:LEU:HD12	6:Bv:172:SER:H	1.83	0.43
2:Cd:1:CYS:HA	3:Ce:4:ILE:HD13	2.01	0.43
6:Ch:115:MET:HG2	7:El:802:LEU:HD22	2.01	0.43
5:Cm:210:TYR:CD2	5:Cm:224:GLY:HA3	2.54	0.43
6:Ct:169:ILE:HB	6:Ct:241:LEU:HD12	2.01	0.43
5:Ec:256:THR:HG22	5:Ec:260:GLN:O	2.19	0.43
7:Es:793:GLU:HA	7:Es:796:GLN:HE21	1.83	0.43
8:Ew:436:ILE:CD1	8:Fa:437:LEU:HD23	2.47	0.43
8:Ex:142:ILE:HG23	8:Ex:267:LEU:HD21	2.00	0.43
8:Ex:337:ALA:HB3	8:Ey:243:LEU:HD11	2.00	0.43
8:Ey:142:ILE:HA	8:Ey:210:VAL:CG2	2.48	0.43
8:Ey:245:SER:HA	8:Ey:279:PHE:CE2	2.53	0.43
8:Ey:419:LEU:HA	8:Ey:422:ILE:CG1	2.49	0.43
8:Fa:449:LYS:HD2	8:Fa:449:LYS:O	2.17	0.43
9:Fb:110:GLY:O	9:Fb:114:LEU:HD23	2.18	0.43
9:Fb:335:ALA:O	9:Fb:338:GLN:HG3	2.18	0.43
9:Fb:729:MET:HE2	9:Fb:729:MET:C	2.43	0.43
9:Fb:807:ASN:ND2	9:Fb:941:LEU:HD13	2.34	0.43
9:Fc:375:LYS:HB2	9:Fc:376:GLN:NE2	2.34	0.43
9:Fc:525:ASP:HA	9:Fc:528:ILE:HG22	2.00	0.43
9:Fc:665:GLN:HE21	9:Fc:665:GLN:HB3	1.66	0.43
9:Fc:755:VAL:HG12	9:Fc:759:MET:SD	2.59	0.43
9:Fd:75:ILE:HD13	9:Fd:75:ILE:HA	1.84	0.43
9:Fd:655:ILE:CG1	9:Fd:710:MET:HE3	2.49	0.43
9:Fe:136:VAL:HG22	9:Fe:461:TRP:HZ2	1.83	0.43
9:Ff:59:ASN:OD1	9:Ff:59:ASN:C	2.61	0.43
9:Ff:452:PHE:HD1	9:Ff:452:PHE:H	1.67	0.43
9:Ff:677:THR:HA	9:Ff:680:GLN:OE1	2.18	0.43
9:Ff:804:ILE:O	9:Ff:808:VAL:HG22	2.19	0.43
7:Fi:36:PHE:O	7:Fi:40:ARG:HG2	2.18	0.43
1:Aa:11:TYR:CE2	3:Ac:9:TYR:HB3	2.53	0.43
6:Bd:169:ILE:HD13	6:Bd:179:LEU:HD21	2.01	0.43
6:Bj:115:MET:HE2	7:Eh:801:MET:HB3	2.00	0.43
6:Cb:221:TYR:HD2	6:Cb:244:GLY:HA3	1.84	0.43
6:Cb:237:VAL:C	6:Cb:238:MET:HE2	2.44	0.43
6:Cn:107:ILE:CD1	7:Em:794:ILE:HA	2.48	0.43
6:Cn:121:PRO:HB2	7:En:801:MET:CG	2.48	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
6:Ct:129:LYS:HE3	6:Cz:112:PHE:CE2	2.54	0.43
5:Cy:218:GLY:HA3	6:Df:148:PRO:HD3	2.01	0.43
6:Dl:126:GLN:NE2	6:Dr:112:PHE:CE1	2.87	0.43
6:Dl:225:ASN:ND2	6:Dr:176:VAL:HG22	2.34	0.43
5:Dq:236:SER:N	5:Dq:243:MET:HE1	2.34	0.43
6:Dr:208:LYS:HD3	6:Dr:208:LYS:HA	1.83	0.43
5:Dw:233:ARG:HH12	6:Ed:246:LYS:HD3	1.83	0.43
6:Dx:191:ILE:HB	6:Dx:206:TRP:CZ2	2.54	0.43
7:En:805:ALA:O	7:En:809:VAL:HG23	2.19	0.43
7:Ep:795:GLN:HA	7:Ep:798:THR:HB	2.00	0.43
7:Et:779:GLN:HG2	7:Et:780:LYS:H	1.84	0.43
8:Ey:103:THR:HG22	8:Ez:255:ASP:HB2	1.99	0.43
8:Ey:381:ARG:HH12	8:Ey:382:ARG:CG	2.32	0.43
9:Fb:530:ILE:H	9:Fb:530:ILE:HD12	1.83	0.43
9:Fb:675:VAL:HG23	9:Fb:678:LEU:HD11	2.00	0.43
9:Fc:147:LEU:HD21	9:Fc:457:ASN:HB3	2.00	0.43
9:Fc:770:ILE:HD11	9:Fd:928:LYS:HZ2	1.84	0.43
9:Fd:34:SER:HA	9:Fd:462:ILE:HD13	2.01	0.43
9:Fd:41:LEU:HD12	9:Fe:828:VAL:HG21	2.00	0.43
9:Fd:222:MET:O	9:Fd:222:MET:SD	2.76	0.43
9:Fd:278:ILE:HD11	9:Fd:516:LEU:HD11	1.99	0.43
9:Fd:739:ILE:HA	9:Fd:742:MET:CE	2.49	0.43
9:Fd:936:LEU:N	9:Fd:937:PRO:HD2	2.33	0.43
9:Fe:113:LEU:HD12	9:Fe:113:LEU:O	2.19	0.43
9:Fe:502:THR:HG23	9:Fe:531:GLN:NE2	2.30	0.43
9:Fe:665:GLN:HA	9:Fe:668:LYS:HE3	2.01	0.43
9:Fe:685:PRO:HB2	9:Fe:830:VAL:HG11	2.00	0.43
9:Ff:239:PHE:HE1	9:Ff:340:TYR:HB3	1.84	0.43
9:Ff:532:SER:HA	9:Ff:537:VAL:H	1.82	0.43
6:Af:144:THR:CG2	6:Af:148:PRO:HG3	2.49	0.43
5:Aw:255:LEU:HD13	5:Aw:261:VAL:HG13	2.01	0.43
6:Ax:113:LYS:HA	6:Ax:113:LYS:HD3	1.81	0.43
1:Ay:6:LEU:HA	1:Ay:9:LEU:HG	2.00	0.43
6:Bd:105:GLU:OE2	6:Bd:109:LYS:HD3	2.18	0.43
6:Bj:107:ILE:HB	7:Eh:797:ARG:NH2	2.34	0.43
6:Bj:245:GLN:HG3	6:Bj:246:LYS:H	1.84	0.43
5:Cg:253:ARG:HB2	5:Cg:263:LYS:HE3	2.01	0.43
6:Ct:112:PHE:HD1	6:Ct:115:MET:HB2	1.83	0.43
5:Cy:251:GLN:HB3	5:Cy:253:ARG:CZ	2.48	0.43
6:Cz:149:PRO:O	6:Cz:151:PRO:HD3	2.19	0.43
6:Cz:189:TRP:O	6:Cz:211:ASN:HB2	2.18	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
5:Dq:213:GLN:HE22	5:Dq:223:ILE:HB	1.81	0.43
6:Dr:157:PHE:HD1	6:Dr:255:ARG:HB2	1.84	0.43
6:Dr:190:PRO:HA	6:Dr:211:ASN:HB3	2.00	0.43
6:Dr:238:MET:HE1	6:Dx:178:SER:HB3	2.00	0.43
8:Ew:78:GLN:C	8:Ew:78:GLN:CD	2.87	0.43
8:Ew:198:TRP:HH2	8:Fa:317:VAL:HA	1.83	0.43
8:Ex:211:MET:O	8:Ex:214:VAL:HB	2.18	0.43
8:Ey:40:VAL:O	8:Ey:43:LEU:HD12	2.18	0.43
8:Ey:75:GLN:HB2	8:Ez:400:ILE:HD11	2.00	0.43
8:Ey:382:ARG:O	8:Ey:399:TRP:HB3	2.19	0.43
8:Ez:170:LEU:HD23	8:Ez:171:ASN:H	1.83	0.43
8:Ez:294:LYS:O	8:Ez:298:TYR:HB3	2.18	0.43
8:Ez:445:LEU:HA	8:Ez:448:LEU:HG	2.01	0.43
8:Fa:416:ALA:O	8:Fa:419:LEU:HD12	2.18	0.43
9:Fb:132:ILE:HD12	9:Fb:132:ILE:H	1.83	0.43
9:Fb:266:PHE:CE2	9:Fb:275:LEU:HB2	2.47	0.43
9:Fb:826:SER:O	9:Fb:830:VAL:HG23	2.18	0.43
9:Fc:378:PHE:HB3	9:Fc:414:PHE:HD2	1.82	0.43
9:Fd:302:VAL:HG21	9:Fd:325:LEU:HD22	2.01	0.43
9:Fd:323:SER:HA	9:Fd:326:GLN:HG3	2.01	0.43
9:Fd:810:LEU:O	9:Fd:814:LEU:HG	2.19	0.43
9:Ff:183:THR:HG23	9:Ff:496:PHE:HB2	2.01	0.43
7:Fi:25:ALA:O	7:Fi:29:ILE:HG12	2.18	0.43
6:Af:208:LYS:HE2	7:Ef:823:GLY:HA2	2.00	0.43
6:Al:123:ASN:CG	6:Al:124:PRO:HD2	2.43	0.43
6:Al:168:VAL:C	6:Al:169:ILE:HD13	2.43	0.43
6:Al:208:LYS:HD3	6:Al:208:LYS:HA	1.74	0.43
6:Ar:173:GLN:HE21	6:Ar:220:LEU:HD23	1.84	0.43
3:Au:1:ARG:HG2	3:Au:2:VAL:N	2.34	0.43
6:Ax:220:LEU:HD12	6:Bd:138:GLU:OE2	2.18	0.43
5:Bc:247:ILE:HG12	5:Bc:254:ILE:CD1	2.49	0.43
5:Bc:248:ASP:CB	5:Bc:253:ARG:HH21	2.31	0.43
6:Bv:191:ILE:HG12	6:Bv:211:ASN:HA	2.00	0.43
1:Bw:6:LEU:CD1	5:Ca:246:LEU:HD23	2.49	0.43
6:Cb:149:PRO:HB2	6:Cb:248:VAL:HG12	2.01	0.43
6:Cb:190:PRO:O	6:Cb:231:ARG:HG2	2.19	0.43
5:Cs:218:GLY:HA3	6:Cz:148:PRO:HD3	2.00	0.43
6:Ct:200:SER:HB3	7:Eq:816:GLU:OE2	2.19	0.43
6:Cz:106:VAL:HG23	6:Cz:110:LYS:HZ1	1.80	0.43
6:Dx:122:LEU:HB3	6:Dx:127:VAL:HG22	2.01	0.43
6:Ed:114:ASP:O	6:Ed:117:ARG:HB3	2.18	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
7: Ei:809: VAL: HA	7: Ei:812: TRP: CZ3	2.53	0.43
7: Em:808: LEU: C	7: Em:812: TRP: CD1	2.96	0.43
8: Ex:80: TYR: CD2	8: Ex:359: MET: HG3	2.53	0.43
8: Ex:84: THR: HG23	8: Ex:359: MET: HE1	2.01	0.43
8: Ex:139: ASN: HD21	8: Ex:141: LEU: HD13	1.83	0.43
8: Ex:440: ASN: HA	8: Ex:443: MET: HG3	2.00	0.43
8: Ey:191: MET: HE3	8: Ey:197: ASN: O	2.18	0.43
8: Ey:211: MET: HA	8: Ey:214: VAL: CG2	2.48	0.43
8: Fa:128: PRO: HB3	8: Fa:168: THR: HA	2.01	0.43
8: Fa:211: MET: O	8: Fa:215: ILE: HG22	2.18	0.43
9: Fb:908: VAL: HG23	9: Fb:912: PHE: CE2	2.53	0.43
9: Fc:226: CYS: HB3	9: Fc:227: ARG: HH21	1.84	0.43
9: Fc:278: ILE: HD12	9: Fc:278: ILE: HA	1.88	0.43
9: Fc:675: VAL: O	9: Fc:679: THR: HG23	2.19	0.43
9: Fc:940: VAL: O	9: Fc:944: ILE: HD12	2.19	0.43
9: Fd:320: ILE: HD11	9: Fd:325: LEU: HB2	2.01	0.43
9: Fd:391: GLU: HG3	9: Fd:393: GLN: H	1.84	0.43
9: Fd:650: GLU: HG2	9: Fd:651: MET: HE2	2.01	0.43
9: Fd:826: SER: O	9: Fd:830: VAL: HG12	2.18	0.43
9: Fd:912: PHE: O	9: Fd:913: PHE: C	2.62	0.43
9: Fe:738: TRP: O	9: Fe:741: THR: HB	2.19	0.43
9: Ff:924: ILE: O	9: Ff:928: LYS: HG2	2.18	0.43
9: Ff:928: LYS: HD2	9: Ff:928: LYS: N	2.34	0.43
6: Af:106: VAL: HG23	6: Af:110: LYS: HZ2	1.82	0.43
6: Af:221: TYR: CD1	6: Al:139: TYR: HD1	2.37	0.43
5: Aq:220: ALA: HB2	5: Aq:247: ILE: HG21	2.00	0.43
6: Ax:168: VAL: HG13	6: Ax:242: ILE: HD13	2.01	0.43
6: Bd:181: PHE: CZ	6: Bd:213: LEU: HB2	2.47	0.43
4: Bh:2: PHE: HB2	5: Bi:251: GLN: HE22	1.84	0.43
1: Bk:1: CYS: HA	4: Bn:2: PHE: CG	2.54	0.43
1: Bw:11: TYR: HD2	3: By:9: TYR: HB3	1.84	0.43
2: Bx:5: ILE: HG21	6: Ch:140: ALA: HB2	2.00	0.43
5: Ca:217: PRO: HD3	6: Ch:139: TYR: OH	2.18	0.43
6: Cb:181: PHE: CD2	6: Cb:181: PHE: N	2.86	0.43
6: Ct:225: ASN: ND2	6: Cz:175: PHE: HB3	2.34	0.43
6: Cz:238: MET: HG3	6: Df:250: TYR: CD2	2.54	0.43
1: Dg:16: ALA: HB3	2: Dn:9: ARG: HG3	2.01	0.43
6: Dl:141: LYS: HE3	7: Es:812: TRP: CH2	2.53	0.43
6: Dl:221: TYR: HE2	6: Dr:138: GLU: HB2	1.83	0.43
6: Dr:169: ILE: HG13	6: Dr:171: LEU: HD22	1.99	0.43
6: Dr:169: ILE: HG23	6: Dr:241: LEU: HA	2.01	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
5:Dw:218:GLY:HA3	6:Ed:148:PRO:HD3	1.99	0.43
6:Dx:182:LEU:HD11	6:Dx:186:GLY:CA	2.49	0.43
7:Ee:802:LEU:HD12	7:Ee:803:THR:N	2.34	0.43
7:Eu:806:THR:HA	7:Eu:809:VAL:HG12	2.01	0.43
8:Ew:125:PHE:HA	8:Ew:137:THR:HA	2.00	0.43
9:Fb:124:MET:HE1	9:Fb:780:PRO:HA	2.01	0.43
9:Fb:283:LYS:HD2	9:Fb:493:LYS:HG3	2.01	0.43
9:Fb:648:PHE:CE2	9:Fb:652:ILE:HD11	2.54	0.43
9:Fb:710:MET:HA	9:Fb:714:SER:HB3	2.01	0.43
9:Fb:799:GLU:O	9:Fb:803:MET:HG3	2.18	0.43
9:Fc:278:ILE:HD13	9:Fc:513:TYR:CZ	2.54	0.43
9:Fc:532:SER:HA	9:Fc:537:VAL:N	2.33	0.43
9:Fc:663:PRO:HA	9:Fc:699:PHE:HZ	1.83	0.43
9:Fc:921:MET:HA	9:Fc:921:MET:HE3	1.99	0.43
9:Fc:940:VAL:HG22	9:Fc:943:TRP:CH2	2.54	0.43
9:Fd:785:GLY:HA2	9:Fd:788:HIS:CD2	2.54	0.43
9:Fd:843:TYR:HD2	9:Fd:843:TYR:H	1.66	0.43
9:Fe:274:PHE:CZ	9:Fe:515:LEU:HG	2.53	0.43
9:Fe:726:ALA:O	9:Fe:727:LEU:C	2.62	0.43
9:Ff:459:LYS:HA	9:Ff:459:LYS:HD3	1.80	0.43
9:Ff:704:TRP:CE3	9:Ff:707:LEU:HD23	2.54	0.43
7:Fg:37:PHE:HB2	7:Fg:38:LYS:HZ2	1.84	0.43
3:Ac:3:SER:O	6:Ar:132:GLN:HG3	2.19	0.42
6:Af:117:ARG:HE	7:Ev:797:ARG:NE	2.17	0.42
6:Al:141:LYS:HD2	7:Ef:812:TRP:CZ2	2.53	0.42
3:Ba:2:VAL:HG21	6:Bp:129:LYS:CG	2.45	0.42
3:Bg:3:SER:O	6:Bv:132:GLN:HG3	2.19	0.42
6:Bj:204:ILE:HD11	6:Bj:213:LEU:HD22	2.01	0.42
3:Bm:2:VAL:HG21	6:Cb:129:LYS:CG	2.49	0.42
6:Cb:213:LEU:HD23	6:Cb:213:LEU:HA	1.79	0.42
6:Ch:181:PHE:CE2	6:Ch:237:VAL:HG11	2.54	0.42
6:Ch:191:ILE:HD12	6:Ch:206:TRP:NE1	2.34	0.42
3:Ck:13:LYS:HD3	3:Ck:13:LYS:HA	1.66	0.42
6:Cn:117:ARG:O	6:Cn:121:PRO:HA	2.19	0.42
2:Cv:6:GLY:HA2	6:Dl:125:GLU:HB3	2.00	0.42
6:Cz:219:LYS:NZ	6:Df:142:ALA:HB2	2.34	0.42
6:Df:221:TYR:HB2	6:Dl:142:ALA:HB1	2.01	0.42
6:Df:242:ILE:HD12	6:Df:243:PRO:HD2	2.00	0.42
6:Dl:176:VAL:HG13	6:Dl:214:MET:CG	2.49	0.42
3:Do:2:VAL:HG21	6:Ed:129:LYS:HD3	2.01	0.42
5:Dw:219:ARG:NH2	5:Dw:231:THR:HG23	2.28	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:Dy:3:ASP:O	1:Dy:6:LEU:HD12	2.18	0.42
6:Ed:117:ARG:HH21	7:Eu:797:ARG:CG	2.24	0.42
6:Ed:207:ASP:OD2	6:Ed:210:SER:HB3	2.19	0.42
6:Ed:218:THR:HA	7:Ev:815:VAL:HG11	2.01	0.42
7:El:793:GLU:HA	7:El:796:GLN:HE21	1.84	0.42
7:Er:797:ARG:NH1	7:Er:798:THR:HG23	2.34	0.42
8:Ew:278:ASN:HA	8:Ew:281:ARG:CD	2.49	0.42
8:Ey:186:ASP:N	8:Ey:189:PHE:HE2	2.10	0.42
8:Fa:74:VAL:HA	8:Fa:381:ARG:HH22	1.84	0.42
8:Fa:235:TYR:O	8:Fa:238:PRO:HD2	2.19	0.42
9:Fc:125:GLN:NE2	9:Fc:126:VAL:HG23	2.34	0.42
9:Fc:666:GLY:HA2	9:Fc:669:ASP:OD2	2.19	0.42
9:Fc:697:ILE:HD13	9:Fc:751:TYR:OH	2.19	0.42
9:Fd:33:LEU:HG	9:Fd:459:LYS:O	2.18	0.42
9:Fd:427:ASN:HD21	9:Fd:485:PHE:HB3	1.83	0.42
9:Fd:671:PHE:O	9:Fd:675:VAL:HG23	2.19	0.42
9:Fd:725:PHE:HA	9:Fd:728:ILE:CG1	2.49	0.42
9:Fd:908:VAL:O	9:Fd:911:PHE:HB3	2.18	0.42
9:Fd:911:PHE:O	9:Fd:915:ILE:HG12	2.18	0.42
9:Fe:373:ILE:HG22	9:Fe:533:LEU:HD13	2.01	0.42
9:Fe:627:MET:HA	9:Fe:630:ASP:HB2	2.01	0.42
9:Fe:731:ALA:O	9:Fe:735:LEU:HB2	2.19	0.42
9:Ff:768:TRP:CD1	9:Ff:811:ARG:HB2	2.54	0.42
6:Af:107:ILE:H	6:Af:107:ILE:HD12	1.84	0.42
6:Af:129:LYS:HD3	3:Du:2:VAL:HG21	2.01	0.42
6:Af:145:PRO:HG3	6:Al:131:LYS:HG2	2.00	0.42
1:Am:2:THR:O	1:Am:6:LEU:HG	2.19	0.42
1:Am:9:LEU:O	1:Am:13:LYS:HG2	2.19	0.42
6:Bd:157:PHE:CD1	6:Bd:255:ARG:HG3	2.35	0.42
6:Bd:208:LYS:HZ2	7:Ej:822:GLU:C	2.27	0.42
2:Bl:5:ILE:HD13	6:Bv:140:ALA:HB2	2.00	0.42
6:Bp:220:LEU:HD12	6:Bv:138:GLU:OE1	2.19	0.42
5:Ca:237:LYS:HD3	5:Ca:243:MET:HB3	2.00	0.42
6:Cb:200:SER:HB3	7:En:816:GLU:HG3	2.01	0.42
6:Cb:214:MET:HE1	7:Em:818:GLN:HE21	1.84	0.42
6:Ch:122:LEU:HD12	7:Em:808:LEU:HD11	2.00	0.42
6:Ct:207:ASP:OD2	6:Ct:210:SER:HB3	2.20	0.42
5:Cy:212:ILE:HG23	5:Cy:222:LEU:HD22	2.00	0.42
6:Cz:169:ILE:HD12	6:Cz:240:THR:O	2.19	0.42
1:Dm:7:ALA:HA	1:Dm:10:GLU:OE1	2.19	0.42
5:Dq:265:SER:HB3	5:Dq:268:ASP:HB3	2.01	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
6:Dr:229:ARG:HD3	6:Dr:229:ARG:C	2.45	0.42
6:Dx:107:ILE:H	6:Dx:107:ILE:HD12	1.84	0.42
7:Eh:799:SER:HA	7:Eh:802:LEU:CD2	2.47	0.42
7:Ek:795:GLN:H	7:Ek:795:GLN:CD	2.28	0.42
7:En:795:GLN:HA	7:En:798:THR:OG1	2.19	0.42
7:Et:807:GLN:OE1	7:Et:808:LEU:HD23	2.19	0.42
8:Ew:49:TYR:HB3	8:Fa:52:TYR:CD1	2.54	0.42
8:Ey:219:PRO:HA	8:Ey:220:PRO:HD3	1.81	0.42
8:Ey:383:LEU:HD23	8:Ey:384:PHE:N	2.34	0.42
8:Ez:448:LEU:HD12	8:Ez:449:LYS:N	2.34	0.42
8:Fa:280:ILE:CD1	8:Fa:351:LEU:HD22	2.49	0.42
8:Fa:300:GLU:HA	8:Fa:303:ASN:ND2	2.33	0.42
9:Fc:651:MET:O	9:Fc:655:ILE:HG12	2.19	0.42
9:Fe:78:TYR:HA	9:Fe:81:MET:CE	2.49	0.42
9:Fe:606:LYS:HB2	9:Fe:608:GLN:HE22	1.83	0.42
9:Ff:165:LYS:HD2	9:Ff:485:PHE:HB2	2.01	0.42
9:Ff:323:SER:O	9:Ff:327:THR:HG23	2.19	0.42
9:Ff:807:ASN:OD1	9:Ff:807:ASN:C	2.62	0.42
3:Ba:11:ALA:HB1	3:Ba:14:TYR:CE2	2.54	0.42
6:Bd:138:GLU:HA	6:Bd:141:LYS:HB3	2.00	0.42
6:Bd:195:ASP:O	6:Bd:226:LEU:HD12	2.19	0.42
6:Bd:196:LEU:HD21	6:Bd:202:PHE:CD2	2.54	0.42
6:Bj:182:LEU:HD11	6:Bj:186:GLY:HA2	2.02	0.42
5:Bo:218:GLY:HA3	6:Bv:148:PRO:HD3	2.02	0.42
6:Bv:108:ASP:OD1	6:Bv:108:ASP:C	2.61	0.42
6:Cb:184:SER:HA	6:Cb:255:ARG:HH21	1.84	0.42
5:Cm:207:ARG:CZ	5:Cm:240:GLY:HA3	2.50	0.42
6:Cn:157:PHE:CD1	6:Cn:157:PHE:C	2.96	0.42
6:Cz:187:ALA:HB1	6:Cz:262:ASN:HB2	2.01	0.42
1:Dg:8:ALA:HB1	3:Di:1:ARG:HD3	2.01	0.42
6:Dr:144:THR:HG23	6:Dr:148:PRO:HG3	2.01	0.42
6:Dx:122:LEU:HD11	6:Dx:130:LEU:CD1	2.49	0.42
6:Dx:122:LEU:HD22	7:Et:801:MET:HG2	2.02	0.42
2:Dz:2:VAL:HG22	3:Ea:1:ARG:HE	1.84	0.42
7:Ep:805:ALA:O	7:Ep:809:VAL:HG12	2.19	0.42
8:Ew:78:GLN:NE2	8:Ex:384:PHE:HB2	2.31	0.42
8:Ew:179:PHE:HB2	8:Ex:240:ILE:HD13	2.02	0.42
8:Ew:211:MET:HA	8:Ew:214:VAL:CG2	2.46	0.42
8:Ew:218:LEU:HD21	8:Fa:326:LEU:HD12	2.00	0.42
8:Ew:292:LEU:HA	8:Ew:329:TYR:OH	2.19	0.42
8:Ew:293:PRO:HB3	8:Fa:457:PHE:CZ	2.54	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
8:Ew:385:ASP:H	8:Ew:398:GLN:NE2	2.14	0.42
8:Ex:302:TRP:HZ3	8:Ey:218:LEU:HG	1.78	0.42
8:Ey:82:TYR:CE2	8:Ez:384:PHE:HB3	2.55	0.42
8:Ey:382:ARG:HG2	8:Ey:399:TRP:CE2	2.55	0.42
8:Ey:415:ILE:O	8:Ey:418:LEU:HD12	2.19	0.42
9:Fb:693:GLY:HA3	9:Fb:749:THR:HG23	2.01	0.42
9:Fc:136:VAL:HG13	9:Fc:461:TRP:NE1	2.32	0.42
9:Fc:258:SER:HA	9:Fc:285:ASN:HA	2.00	0.42
9:Fc:705:LEU:HA	9:Fc:708:LEU:CD1	2.50	0.42
9:Fc:739:ILE:O	9:Fc:743:VAL:HG13	2.19	0.42
9:Fc:912:PHE:O	9:Fc:916:LEU:HD22	2.20	0.42
9:Fd:242:LYS:HZ1	9:Fd:259:PHE:C	2.27	0.42
9:Fd:452:PHE:CZ	9:Fd:470:ASP:HB3	2.53	0.42
9:Fd:461:TRP:CZ2	9:Fd:812:PRO:HB3	2.55	0.42
9:Fd:639:TYR:OH	9:Fe:605:MET:HB2	2.19	0.42
9:Fd:648:PHE:HB2	9:Fd:717:ILE:HD13	2.00	0.42
9:Fd:776:MET:SD	9:Fd:777:VAL:HG23	2.59	0.42
9:Ff:202:ARG:HA	9:Ff:202:ARG:HD2	1.85	0.42
9:Ff:378:PHE:HB3	9:Ff:414:PHE:HD2	1.84	0.42
9:Ff:434:LEU:HA	9:Ff:437:ILE:HG22	2.00	0.42
9:Ff:652:ILE:O	9:Ff:656:VAL:HG23	2.19	0.42
9:Ff:734:LEU:O	9:Ff:734:LEU:HD12	2.19	0.42
9:Ff:750:ALA:C	9:Ff:751:TYR:HD1	2.28	0.42
9:Ff:914:SER:HA	9:Ff:917:ILE:HG22	2.01	0.42
7:Fj:34:ILE:HD13	7:Fj:37:PHE:HE1	1.83	0.42
7:Fk:30:ILE:HD13	7:Fk:30:ILE:HA	1.88	0.42
6:Af:220:LEU:HG	6:Al:138:GLU:OE1	2.20	0.42
1:Ag:3:ASP:HA	1:Ag:6:LEU:HG	2.01	0.42
6:Ar:158:VAL:HG23	6:Ar:254:LEU:HB2	2.02	0.42
6:Ar:194:TYR:HE2	6:Ar:196:LEU:HB2	1.84	0.42
5:Bc:237:LYS:NZ	5:Bc:243:MET:HB2	2.34	0.42
6:Bd:119:LEU:HD12	6:Bd:120:TYR:CG	2.54	0.42
5:Bi:260:GLN:H	5:Bi:260:GLN:CD	2.27	0.42
6:Bp:251:ARG:HG2	6:Bp:251:ARG:HH11	1.85	0.42
5:Bu:210:TYR:HA	5:Bu:224:GLY:HA2	2.01	0.42
6:Bv:189:TRP:CG	6:Bv:230:LEU:HD12	2.54	0.42
5:Cm:219:ARG:HH22	6:Ct:150:LYS:HD2	1.84	0.42
6:Ct:179:LEU:HB3	6:Ct:254:LEU:HD11	2.01	0.42
6:Cz:219:LYS:HD3	6:Cz:219:LYS:C	2.44	0.42
6:Df:191:ILE:HD11	6:Df:211:ASN:C	2.44	0.42
1:Dy:1:CYS:HB2	3:Ea:16:ASP:HB3	2.01	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
6:Ed:106:VAL:HA	6:Ed:109:LYS:HG2	2.00	0.42
6:Ed:206:TRP:HB2	6:Ed:213:LEU:HD23	2.01	0.42
7:Ee:799:SER:HA	7:Ee:802:LEU:HG	2.01	0.42
7:Ef:809:VAL:HA	7:Ef:812:TRP:HD1	1.83	0.42
7:Eh:793:GLU:O	7:Eh:796:GLN:HG3	2.19	0.42
7: Ei:800:ASP:OD1	7: Ei:800:ASP:C	2.62	0.42
8:Ew:302:TRP:CE2	8:Ex:220:PRO:HA	2.55	0.42
8:Ew:346:ASN:HA	8:Ew:435:ARG:NH2	2.34	0.42
8:Ey:428:LEU:O	8:Ey:432:ILE:HG12	2.20	0.42
8:Ey:452:GLN:HA	8:Ey:453:PRO:HD3	1.94	0.42
8:Fa:254:MET:HE3	8:Fa:354:ARG:CB	2.45	0.42
8:Fa:399:TRP:CD1	8:Fa:399:TRP:C	2.96	0.42
9:Fb:103:ILE:N	9:Fb:104:PRO:HD2	2.34	0.42
9:Fb:185:LEU:HD12	9:Fb:189:ILE:HD11	2.00	0.42
9:Fb:193:GLY:O	9:Fb:197:GLN:HG2	2.19	0.42
9:Fb:240:VAL:HG13	9:Fb:337:GLN:HG3	2.01	0.42
9:Fc:77:MET:O	9:Fc:81:MET:HE3	2.20	0.42
9:Fc:705:LEU:HA	9:Fc:708:LEU:HD12	2.01	0.42
9:Fc:716:LEU:HG	9:Fd:603:TYR:CZ	2.54	0.42
9:Fd:301:LEU:HD21	9:Fd:319:ASN:HD21	1.84	0.42
9:Fd:513:TYR:HB3	9:Fd:516:LEU:HG	2.02	0.42
9:Fe:169:ALA:HB1	9:Fe:485:PHE:CD1	2.54	0.42
9:Fe:429:ILE:C	9:Fe:430:MET:HE2	2.44	0.42
9:Fe:452:PHE:O	9:Fe:455:GLU:HG2	2.19	0.42
9:Ff:320:ILE:HD12	9:Ff:325:LEU:HB2	2.01	0.42
9:Ff:768:TRP:HD1	9:Ff:811:ARG:HB2	1.84	0.42
5:Ae:212:ILE:HB	5:Ae:264:PHE:CD1	2.55	0.42
6:Al:252:VAL:HG12	6:Al:254:LEU:HG	2.01	0.42
6:Ar:131:LYS:HD2	7:Ef:812:TRP:CH2	2.55	0.42
6:Bd:123:ASN:HB3	6:Bd:126:GLN:HG3	2.01	0.42
6:Bj:141:LYS:NZ	7:Ej:812:TRP:HD1	2.18	0.42
6:Bp:182:LEU:HD12	6:Bp:186:GLY:C	2.44	0.42
6:Bv:115:MET:HA	6:Bv:118:ASN:HD21	1.85	0.42
5:Ca:212:ILE:HD13	5:Ca:220:ALA:HB1	2.02	0.42
6:Cb:191:ILE:HA	6:Cb:230:LEU:HD13	2.01	0.42
6:Cn:133:ILE:CD1	6:Ct:116:THR:HG22	2.50	0.42
6:Ct:194:TYR:HE2	6:Ct:196:LEU:HB2	1.84	0.42
5:Ec:262:ILE:HD13	5:Ec:262:ILE:HA	1.90	0.42
7:Ef:771:ALA:HA	7:Ef:774:LEU:HG	2.01	0.42
7:Es:774:LEU:HD11	9:Fb:579:LEU:CD1	2.50	0.42
8:Ew:233:TYR:HB3	8:Ew:234:LYS:NZ	2.34	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
8:Ew:317:VAL:CA	8:Ex:198:TRP:HE1	2.20	0.42
8:Ew:408:PRO:HA	8:Ew:411:VAL:HG22	2.01	0.42
8:Ew:413:LYS:HE3	8:Ex:403:ILE:O	2.20	0.42
8:Ew:444:LEU:CD2	8:Ex:443:MET:HB3	2.46	0.42
8:Ex:69:PHE:HE2	8:Ex:403:ILE:HD13	1.85	0.42
8:Ey:62:ASN:HA	8:Ez:48:LYS:HZ1	1.84	0.42
8:Ey:398:GLN:HB2	8:Ey:401:LYS:HG3	2.01	0.42
8:Ez:187:TYR:HB3	8:Ez:206:TYR:CE1	2.55	0.42
8:Fa:350:ILE:HG13	8:Fa:350:ILE:H	1.64	0.42
9:Fb:77:MET:O	9:Fb:81:MET:HE2	2.18	0.42
9:Fb:318:LEU:HD21	9:Fb:439:GLN:C	2.44	0.42
9:Fb:450:ARG:HD2	9:Fb:450:ARG:HA	1.81	0.42
9:Fb:542:GLN:N	9:Fb:542:GLN:CD	2.78	0.42
9:Fb:672:ILE:O	9:Fb:676:GLN:HG2	2.19	0.42
9:Fc:61:PHE:HA	9:Fc:64:PHE:HB3	2.02	0.42
9:Fc:243:GLN:NE2	9:Fc:330:LEU:HD22	2.30	0.42
9:Fc:357:PHE:HZ	9:Fc:530:ILE:HD13	1.85	0.42
9:Fd:71:LEU:O	9:Fd:74:ILE:HG23	2.19	0.42
9:Fd:671:PHE:CD2	9:Fd:672:ILE:HD13	2.55	0.42
9:Fe:130:TRP:O	9:Fe:134:GLN:HG2	2.19	0.42
9:Fe:693:GLY:O	9:Fe:697:ILE:HG12	2.20	0.42
9:Ff:301:LEU:HG	9:Ff:319:ASN:OD1	2.19	0.42
9:Ff:579:LEU:HD23	9:Ff:579:LEU:HA	1.90	0.42
9:Ff:753:ILE:H	9:Ff:753:ILE:HG13	1.62	0.42
9:Ff:753:ILE:CG2	9:Ff:926:VAL:HG11	2.50	0.42
7:Fg:9:LYS:HA	7:Fg:12:PHE:CD2	2.54	0.42
7:Fh:19:VAL:HA	7:Fh:22:ILE:CG1	2.49	0.42
5:Ae:212:ILE:HA	5:Ae:222:LEU:HD22	2.02	0.42
5:Aq:237:LYS:HD3	5:Aq:243:MET:HB3	2.02	0.42
6:Ar:221:TYR:CA	6:Ar:243:PRO:HG2	2.50	0.42
6:Bj:106:VAL:HA	6:Bj:109:LYS:HD3	2.02	0.42
6:Bp:207:ASP:OD2	6:Bp:210:SER:HB3	2.20	0.42
6:Cb:203:ASN:O	6:Cb:215:ILE:HD12	2.20	0.42
2:Cj:5:ILE:CD1	6:Ct:139:TYR:HB3	2.50	0.42
6:Ct:132:GLN:N	6:Ct:132:GLN:CD	2.77	0.42
6:Ct:154:THR:HG23	6:Ct:252:VAL:HG22	2.00	0.42
6:Ct:220:LEU:HG	6:Ct:221:TYR:CD2	2.54	0.42
2:Cv:5:ILE:HD12	6:Df:139:TYR:CD2	2.47	0.42
6:Df:179:LEU:HB3	6:Df:254:LEU:HD11	2.00	0.42
5:Dk:214:ALA:HB3	5:Dk:221:TRP:CD1	2.51	0.42
6:Dl:236:PRO:HB2	6:Dr:251:ARG:NH2	2.35	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:Dm:6:LEU:HD12	1:Dm:7:ALA:N	2.34	0.42
6:Dr:187:ALA:HB1	6:Dr:262:ASN:HB2	2.01	0.42
6:Dx:220:LEU:HB3	6:Dx:221:TYR:CE2	2.55	0.42
6:Ed:115:MET:CE	7:Et:801:MET:HE3	2.49	0.42
7:Ej:788:GLN:O	7:Ej:792:GLN:OE1	2.38	0.42
8:Ew:175:SER:HB3	8:Ex:244:ASN:ND2	2.35	0.42
8:Ex:424:TYR:OH	8:Ey:383:LEU:N	2.52	0.42
8:Ey:409:ALA:O	8:Ey:412:GLN:HG2	2.20	0.42
8:Ez:123:GLN:O	8:Ez:135:LYS:HD2	2.19	0.42
9:Fb:370:PHE:CE1	9:Fb:565:SER:HB3	2.54	0.42
9:Fb:714:SER:HB2	9:Fc:598:VAL:HG12	2.02	0.42
9:Fc:100:SER:HB3	7:Fg:16:ARG:NH1	2.35	0.42
9:Fc:648:PHE:HE1	9:Fc:724:ILE:HD11	1.83	0.42
9:Fd:49:LEU:HG	9:Fd:51:GLY:H	1.85	0.42
9:Fd:195:GLN:NE2	9:Fd:232:ASP:HA	2.34	0.42
9:Fd:635:TYR:O	9:Fd:639:TYR:HB3	2.19	0.42
9:Fe:624:LEU:O	9:Fe:628:MET:HB2	2.19	0.42
9:Fe:632:PHE:N	9:Fe:632:PHE:CD1	2.88	0.42
9:Ff:464:ALA:HB1	9:Ff:819:TYR:HE2	1.83	0.42
9:Ff:547:GLN:HA	9:Ff:568:TYR:HE1	1.85	0.42
7:Fi:3:SER:HB3	7:Fi:6:GLU:OE1	2.20	0.42
6:Af:203:ASN:HB3	6:Af:216:GLN:OE1	2.19	0.42
5:Ak:213:GLN:HB2	5:Ak:223:ILE:HG22	2.02	0.42
6:Ar:105:GLU:O	6:Ar:108:ASP:HB3	2.19	0.42
6:Ar:238:MET:HB3	6:Ax:250:TYR:CZ	2.55	0.42
6:Bj:229:ARG:HD3	6:Bj:229:ARG:C	2.45	0.42
3:Bm:2:VAL:HG11	6:Cb:129:LYS:HG2	2.02	0.42
4:Bn:4:ALA:HB3	5:Bo:251:GLN:NE2	2.35	0.42
6:Bp:138:GLU:O	6:Bp:141:LYS:HB3	2.20	0.42
6:Bp:158:VAL:HG13	6:Bp:167:PRO:HG2	2.01	0.42
6:Ch:119:LEU:HD22	7:El:805:ALA:HB1	2.02	0.42
1:Co:15:ASN:HB3	2:Cv:9:ARG:NH1	2.35	0.42
6:Ct:182:LEU:HD12	6:Ct:253:ASP:OD2	2.20	0.42
1:Da:11:TYR:HD2	3:Dc:9:TYR:HB3	1.84	0.42
2:Db:4:MET:HE1	5:De:249:SER:H	1.80	0.42
5:Dk:216:ILE:CD1	6:Dr:148:PRO:HG2	2.50	0.42
6:Dr:123:ASN:ND2	6:Dr:124:PRO:HD2	2.35	0.42
1:Ds:11:TYR:CD2	3:Du:9:TYR:HB3	2.55	0.42
6:Ed:207:ASP:HB3	7:Ev:820:TYR:CE2	2.55	0.42
7:Ei:798:THR:O	7:Ei:802:LEU:HD22	2.20	0.42
7:Eq:776:ALA:C	7:Eq:780:LYS:HZ3	2.26	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
7:Er:813:LYS:HD3	7:Er:813:LYS:N	2.35	0.42
7:Es:809:VAL:O	7:Es:813:LYS:HE3	2.19	0.42
8:Ew:342:VAL:HG13	8:Ew:438:LEU:HD11	2.00	0.42
8:Ex:220:PRO:HG3	8:Ex:227:THR:HG23	2.00	0.42
8:Ex:252:LEU:HD11	8:Ex:373:LEU:CA	2.49	0.42
8:Ex:359:MET:N	8:Ex:359:MET:SD	2.93	0.42
8:Ez:239:LEU:HA	8:Ez:242:GLN:NE2	2.35	0.42
8:Fa:342:VAL:HG11	8:Fa:442:ILE:HG13	2.00	0.42
9:Fb:734:LEU:HD12	9:Fb:734:LEU:O	2.18	0.42
9:Fb:922:TYR:O	9:Fb:923:LEU:C	2.63	0.42
9:Fc:240:VAL:HA	9:Fc:333:ALA:HB1	2.00	0.42
9:Fc:599:ASP:OD1	9:Fc:599:ASP:C	2.62	0.42
9:Fc:614:ARG:HG2	9:Fc:623:CYS:HB3	2.02	0.42
9:Fc:924:ILE:HA	9:Fc:927:GLN:HB3	2.01	0.42
9:Fe:266:PHE:CE2	9:Fe:275:LEU:HB2	2.48	0.42
9:Fe:275:LEU:HD21	9:Fe:519:TRP:HH2	1.85	0.42
9:Fe:701:GLY:O	9:Fe:705:LEU:HG	2.20	0.42
9:Fe:728:ILE:CD1	9:Ff:730:MET:HB3	2.50	0.42
9:Ff:193:GLY:O	9:Ff:197:GLN:HG3	2.20	0.42
9:Ff:426:TYR:O	9:Ff:430:MET:HG2	2.19	0.42
7:Fi:8:LEU:HB3	7:Fi:12:PHE:CZ	2.54	0.42
1:Aa:9:LEU:O	1:Aa:13:LYS:HG2	2.19	0.42
6:Af:199:PRO:HD2	7:Ef:816:GLU:HG2	2.00	0.42
6:Al:151:PRO:HA	6:Al:248:VAL:HG23	2.02	0.42
6:Ax:231:ARG:HD3	6:Ax:231:ARG:HA	1.92	0.42
6:Bd:230:LEU:HB2	6:Bd:233:LEU:CD2	2.50	0.42
6:Bj:115:MET:O	6:Bj:119:LEU:HD22	2.20	0.42
5:Bo:219:ARG:H	6:Bv:148:PRO:HD2	1.84	0.42
6:Bp:189:TRP:CE2	6:Bp:260:GLY:HA2	2.55	0.42
1:Bw:1:CYS:HB2	3:By:16:ASP:HB3	2.02	0.42
5:Ca:238:ILE:HB	5:Ca:241:TYR:HB2	2.00	0.42
6:Cb:168:VAL:C	6:Cb:169:ILE:HD13	2.45	0.42
1:Ci:1:CYS:HA	4:Cl:2:PHE:CG	2.55	0.42
3:Cq:2:VAL:HG21	6:Df:129:LYS:CD	2.50	0.42
3:Cw:12:LYS:O	3:Cw:13:LYS:HG3	2.19	0.42
5:Cy:212:ILE:HB	5:Cy:264:PHE:CE2	2.54	0.42
6:Cz:216:GLN:HB3	7:Eq:818:GLN:NE2	2.34	0.42
5:De:219:ARG:HH21	5:De:231:THR:HB	1.84	0.42
6:Dl:204:ILE:HD13	6:Dl:215:ILE:CD1	2.49	0.42
6:Dr:126:GLN:HB3	6:Dx:112:PHE:CE1	2.54	0.42
6:Dr:167:PRO:O	6:Dr:239:LEU:HG	2.19	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
6:Dr:221:TYR:CE1	6:Dx:139:TYR:CD2	3.07	0.42
1:Ds:3:ASP:HB3	3:Du:14:TYR:CD2	2.54	0.42
7:Ej:771:ALA:O	7:Ej:774:LEU:HD12	2.19	0.42
7:Ej:777:ILE:HG13	9:Fd:338:GLN:HE21	1.85	0.42
8:Ew:170:LEU:N	8:Ew:176:GLN:HE22	2.17	0.42
8:Ew:242:GLN:NE2	8:Ew:269:ALA:H	2.18	0.42
8:Ew:399:TRP:HH2	8:Fa:417:ILE:HA	1.84	0.42
8:Ex:69:PHE:CE2	8:Ex:403:ILE:HD13	2.54	0.42
8:Ex:194:GLU:HG2	8:Ex:196:LYS:HZ2	1.85	0.42
8:Ex:298:TYR:CE2	8:Ey:221:ALA:HB2	2.53	0.42
8:Ey:174:VAL:HG23	8:Ez:244:ASN:HD21	1.84	0.42
8:Fa:69:PHE:CE2	8:Fa:402:GLN:HB3	2.54	0.42
8:Fa:301:LEU:HB2	8:Fa:326:LEU:HD21	2.01	0.42
8:Fa:432:ILE:O	8:Fa:436:ILE:HG12	2.20	0.42
9:Fb:373:ILE:HD11	9:Fb:533:LEU:HB2	2.01	0.42
9:Fb:912:PHE:O	9:Fb:913:PHE:C	2.63	0.42
9:Fb:913:PHE:CE1	9:Ff:751:TYR:HB3	2.55	0.42
9:Fc:208:GLN:HA	9:Fc:210:LYS:NZ	2.35	0.42
9:Fd:128:PHE:CD2	9:Fd:128:PHE:N	2.88	0.42
9:Fd:662:ILE:H	9:Fd:662:ILE:HD12	1.85	0.42
9:Fe:596:PHE:CD1	9:Fe:596:PHE:C	2.97	0.42
9:Fe:667:MET:HG3	9:Fe:911:PHE:CZ	2.55	0.42
9:Fe:741:THR:HG23	9:Fe:916:LEU:HD11	2.01	0.42
9:Fe:804:ILE:O	9:Fe:808:VAL:HG22	2.20	0.42
9:Ff:221:GLU:H	9:Ff:221:GLU:CD	2.27	0.42
9:Ff:290:LEU:HD12	9:Ff:290:LEU:H	1.85	0.42
9:Ff:447:LYS:H	9:Ff:447:LYS:CD	2.33	0.42
9:Ff:468:PHE:CG	9:Ff:469:PHE:N	2.86	0.42
9:Ff:600:THR:HG22	9:Ff:648:PHE:CE2	2.55	0.42
9:Ff:624:LEU:HD12	9:Ff:625:GLY:N	2.35	0.42
7:Fi:27:LEU:HA	7:Fi:30:ILE:HG12	2.01	0.42
6:Af:150:LYS:N	6:Af:247:ALA:HA	2.35	0.42
5:Ak:219:ARG:HD3	6:Ar:148:PRO:O	2.19	0.42
6:Al:196:LEU:O	6:Al:199:PRO:HD3	2.20	0.42
5:Aq:260:GLN:H	5:Aq:260:GLN:HG3	1.61	0.42
6:Ar:118:ASN:O	6:Ar:121:PRO:HD3	2.19	0.42
6:Ar:181:PHE:CZ	6:Ar:213:LEU:HD12	2.55	0.42
6:Ax:199:PRO:HG3	7:Ei:817:THR:O	2.19	0.42
5:Bc:265:SER:HB3	5:Bc:268:ASP:HB3	2.02	0.42
6:Bd:170:ARG:HG3	6:Bd:249:ASP:HB2	2.02	0.42
5:Bi:218:GLY:HA3	6:Bp:148:PRO:CD	2.48	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
6:Bj:171:LEU:HB2	6:Bj:243:PRO:HB3	2.02	0.42
6:Bp:107:ILE:CD1	7:El:794:ILE:HG13	2.50	0.42
6:Bp:126:GLN:CA	6:Bp:129:LYS:HG3	2.44	0.42
6:Bp:199:PRO:HG3	7:El:819:VAL:HG13	2.02	0.42
6:Bv:135:GLU:HA	6:Bv:138:GLU:OE1	2.20	0.42
6:Bv:137:SER:HB3	6:Cb:120:TYR:CE2	2.55	0.42
6:Bv:173:GLN:O	6:Bv:175:PHE:HD1	2.03	0.42
6:Bv:202:PHE:CE2	6:Bv:241:LEU:HB2	2.54	0.42
6:Cb:191:ILE:HB	6:Cb:206:TRP:CH2	2.55	0.42
6:Ch:204:ILE:HD12	6:Ch:214:MET:O	2.20	0.42
6:Ch:209:THR:HG22	7:Eo:824:THR:HG21	2.01	0.42
3:Cq:6:GLY:HA2	6:Df:132:GLN:OE1	2.20	0.42
6:Ct:223:TYR:HB3	6:Ct:242:ILE:HD13	2.02	0.42
5:Dk:253:ARG:NE	5:Dk:255:LEU:HD21	2.35	0.42
5:Dq:253:ARG:HB2	5:Dq:263:LYS:HZ2	1.84	0.42
7:El:797:ARG:HB2	7:El:801:MET:CE	2.50	0.42
8:Ew:220:PRO:HG3	8:Ew:227:THR:HG23	2.02	0.42
8:Ex:37:SER:O	8:Ex:41:THR:HG22	2.20	0.42
8:Ey:302:TRP:HH2	8:Ez:218:LEU:O	2.03	0.42
8:Ey:350:ILE:HG22	8:Ey:354:ARG:NH2	2.35	0.42
8:Ey:437:LEU:H	8:Ey:437:LEU:HD22	1.85	0.42
8:Ez:108:GLY:O	8:Ez:112:ILE:HD12	2.20	0.42
8:Ez:381:ARG:HH21	8:Ez:382:ARG:HH11	1.67	0.42
9:Fb:618:LEU:HB2	9:Ff:618:LEU:HD21	2.00	0.42
9:Fb:726:ALA:O	9:Fb:727:LEU:C	2.62	0.42
9:Fb:739:ILE:O	9:Fb:743:VAL:HG13	2.20	0.42
9:Fb:768:TRP:HB2	9:Fb:815:MET:HE1	2.02	0.42
9:Fc:139:ALA:HB2	9:Fc:812:PRO:HB2	2.02	0.42
9:Fc:147:LEU:HB2	9:Fc:453:ILE:HG23	2.01	0.42
9:Fc:219:THR:CG2	9:Fc:222:MET:HG2	2.49	0.42
9:Fc:420:LEU:HA	9:Fc:423:ILE:HG12	2.02	0.42
9:Fc:767:ALA:HA	9:Fc:770:ILE:CD1	2.49	0.42
9:Fd:218:PRO:HB2	9:Fd:223:ASN:OD1	2.20	0.42
9:Fd:500:GLN:HA	9:Fd:503:LYS:CG	2.48	0.42
9:Fd:828:VAL:O	9:Fd:832:ILE:HG12	2.19	0.42
9:Fe:223:ASN:O	9:Fe:227:ARG:HG2	2.20	0.42
9:Fe:268:LYS:HD3	9:Ff:403:PRO:HB3	2.01	0.42
9:Fe:718:PRO:O	9:Fe:719:LEU:HB2	2.18	0.42
9:Fe:725:PHE:HA	9:Fe:728:ILE:HG22	2.01	0.42
9:Ff:234:ILE:HG22	9:Ff:340:TYR:OH	2.20	0.42
9:Ff:525:ASP:HA	9:Ff:528:ILE:CD1	2.50	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
9:Ff:753:ILE:HA	9:Ff:756:LEU:HB3	2.01	0.42
9:Ff:837:PHE:CG	9:Ff:838:ASP:N	2.87	0.42
5:Ak:207:ARG:HH22	5:Ak:240:GLY:HA3	1.85	0.42
1:Am:4:ALA:H	2:An:9:ARG:HH22	1.67	0.42
1:Am:4:ALA:H	2:An:9:ARG:NH2	2.17	0.42
6:Ar:113:LYS:HD3	6:Ar:113:LYS:HA	1.75	0.42
6:Ar:200:SER:HB3	7:Eh:816:GLU:HG3	2.01	0.42
6:Ar:230:LEU:HD23	6:Ar:230:LEU:HA	1.90	0.42
5:Bc:248:ASP:HB3	5:Bc:253:ARG:HE	1.84	0.42
5:Bc:248:ASP:CG	5:Bc:253:ARG:HH21	2.28	0.42
6:Bj:122:LEU:HD12	7:Ei:808:LEU:HD11	2.02	0.42
1:Bq:1:CYS:HA	4:Bt:2:PHE:CG	2.55	0.42
6:Bv:108:ASP:HA	7:Ej:797:ARG:NH1	2.34	0.42
5:Ca:222:LEU:HD11	5:Ca:262:ILE:HG21	2.01	0.42
6:Cn:183:ASP:HB3	6:Cn:189:TRP:CE3	2.55	0.42
6:Ct:115:MET:HE2	7:En:798:THR:O	2.19	0.42
1:Cu:3:ASP:O	1:Cu:6:LEU:HD12	2.19	0.42
6:Cz:214:MET:C	6:Cz:215:ILE:HD12	2.45	0.42
2:Db:4:MET:HE1	5:De:250:LEU:H	1.84	0.42
5:De:219:ARG:HA	5:De:232:VAL:O	2.20	0.42
5:De:219:ARG:CZ	6:Dl:150:LYS:HB3	2.50	0.42
6:Dl:238:MET:SD	6:Dr:250:TYR:CD2	3.13	0.42
2:Dn:9:ARG:HD3	3:Do:15:ASP:OD1	2.20	0.42
5:Dw:216:ILE:HG23	5:Dw:221:TRP:CZ3	2.55	0.42
6:Ed:221:TYR:CE2	6:Ed:244:GLY:HA3	2.55	0.42
7:Ee:793:GLU:HA	7:Ee:796:GLN:HE21	1.85	0.42
7:Eh:797:ARG:O	7:Eh:801:MET:HG2	2.20	0.42
7:Em:809:VAL:O	7:Em:813:LYS:HG2	2.20	0.42
8:Ew:62:ASN:HB2	8:Ex:48:LYS:NZ	2.35	0.42
8:Ew:240:ILE:HD11	8:Fa:175:SER:HA	2.01	0.42
8:Ex:274:GLN:O	8:Ex:277:LEU:HD12	2.20	0.42
8:Ey:82:TYR:CD2	8:Ez:384:PHE:HB3	2.55	0.42
8:Ez:234:LYS:HA	8:Ez:237:GLN:CD	2.45	0.42
8:Fa:185:PRO:HG2	8:Fa:206:TYR:HA	2.02	0.42
8:Fa:248:LEU:HD21	8:Fa:279:PHE:HB3	2.02	0.42
9:Fb:399:TRP:CE3	9:Fb:416:GLY:HA3	2.55	0.42
9:Fb:530:ILE:HG22	9:Fb:534:ILE:HD11	2.01	0.42
9:Fc:76:ILE:HD12	9:Fc:76:ILE:HA	1.88	0.42
9:Fc:274:PHE:CD2	9:Fc:515:LEU:HD13	2.55	0.42
9:Fc:596:PHE:CD1	9:Fc:656:VAL:HG11	2.55	0.42
9:Fc:698:ASN:HA	9:Fd:905:TRP:HH2	1.82	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
9:Fc:810:LEU:HB3	9:Fc:814:LEU:CD1	2.50	0.42
9:Fd:37:PHE:HE2	9:Fe:831:TRP:HE3	1.68	0.42
9:Fd:94:LEU:HD23	9:Fd:94:LEU:HA	1.90	0.42
9:Fd:143:TRP:CD1	9:Fd:143:TRP:C	2.98	0.42
9:Fd:147:LEU:HD12	9:Fd:148:SER:N	2.35	0.42
9:Fd:220:PRO:O	9:Fd:224:THR:HG23	2.19	0.42
9:Fd:294:ASN:HD21	9:Fd:304:VAL:HG23	1.85	0.42
9:Fd:673:VAL:HA	9:Fd:676:GLN:HG2	2.01	0.42
9:Fe:87:THR:HG23	9:Fe:92:GLN:O	2.20	0.42
9:Fe:720:PHE:O	9:Fe:722:ILE:N	2.53	0.42
9:Fe:912:PHE:O	9:Fe:913:PHE:C	2.63	0.42
9:Ff:194:LEU:HD21	9:Ff:278:ILE:HG21	2.01	0.42
7:Fk:36:PHE:HA	7:Fk:39:ILE:CG1	2.50	0.42
3:Ao:1:ARG:HG2	3:Ao:2:VAL:N	2.35	0.41
2:Az:6:GLY:CA	6:Bp:125:GLU:HB3	2.49	0.41
6:Bj:107:ILE:HG23	7:Eh:794:ILE:HG13	2.01	0.41
6:Bj:123:ASN:HB3	6:Bj:126:GLN:HG3	2.01	0.41
1:Bk:2:THR:O	1:Bk:6:LEU:HG	2.20	0.41
5:Bo:253:ARG:C	5:Bo:253:ARG:NE	2.77	0.41
6:Bp:123:ASN:HB3	6:Bp:126:GLN:CD	2.45	0.41
6:Bp:144:THR:HA	6:Bp:145:PRO:HD3	1.91	0.41
1:Ci:11:TYR:CE2	3:Ck:9:TYR:HB3	2.55	0.41
6:Cn:251:ARG:HH22	6:Cn:253:ASP:N	2.18	0.41
5:Cs:254:ILE:O	5:Cs:261:VAL:HA	2.18	0.41
6:Cz:107:ILE:HA	6:Cz:110:LYS:NZ	2.28	0.41
6:Cz:238:MET:HE3	6:Df:250:TYR:CE2	2.55	0.41
6:Df:204:ILE:HD11	6:Df:213:LEU:CD1	2.50	0.41
5:Dk:247:ILE:HG13	5:Dk:254:ILE:HD12	2.01	0.41
6:Dl:208:LYS:HB3	7:Et:824:THR:HG22	2.01	0.41
6:Dr:149:PRO:HA	6:Dr:246:LYS:O	2.20	0.41
6:Dr:221:TYR:HD2	6:Dr:244:GLY:HA3	1.84	0.41
5:Dw:219:ARG:HG2	6:Ed:148:PRO:HD2	2.03	0.41
6:Dx:110:LYS:HG3	6:Dx:111:ALA:N	2.35	0.41
7:Ei:777:ILE:HB	7:Ej:778:LEU:HD13	2.02	0.41
7:Ei:789:LYS:HA	7:Ei:792:GLN:HE22	1.84	0.41
7:Eo:822:GLU:OE1	7:Eo:822:GLU:N	2.52	0.41
7:Eq:777:ILE:HA	7:Eq:780:LYS:HZ3	1.85	0.41
8:Ew:39:LEU:HD11	8:Ex:39:LEU:HD21	2.02	0.41
8:Ex:207:GLN:OE1	8:Ex:291:SER:HA	2.20	0.41
8:Ex:436:ILE:HG22	8:Ex:440:ASN:OD1	2.20	0.41
8:Ey:298:TYR:HE2	8:Ez:221:ALA:HA	1.85	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
8:Ey:302:TRP:HZ3	8:Ez:218:LEU:HG	1.82	0.41
8:Ez:188:SER:OG	8:Ez:295:LEU:HG	2.19	0.41
8:Ez:193:ASN:HD21	9:Ff:217:ASN:H	1.67	0.41
8:Ez:368:ILE:HA	8:Ez:368:ILE:HD13	1.83	0.41
8:Fa:373:LEU:O	8:Fa:376:PHE:HB3	2.20	0.41
9:Fb:93:MET:HE1	9:Fb:100:SER:OG	2.19	0.41
9:Fb:612:CYS:SG	9:Fb:614:ARG:HG2	2.60	0.41
9:Fb:693:GLY:HA2	9:Fb:696:TYR:HD2	1.85	0.41
9:Fc:334:ILE:HD12	9:Fc:335:ALA:N	2.34	0.41
9:Fc:431:MET:HE3	9:Fc:432:PRO:HD3	2.02	0.41
9:Fc:455:GLU:O	9:Fc:458:ALA:HB3	2.20	0.41
9:Fc:728:ILE:HD11	9:Fd:730:MET:HB3	2.02	0.41
9:Fc:756:LEU:HA	9:Fc:756:LEU:HD12	1.84	0.41
9:Fd:426:TYR:CD2	9:Fd:430:MET:HE2	2.55	0.41
9:Fd:617:ILE:H	9:Fd:617:ILE:HD12	1.85	0.41
9:Fe:124:MET:HG2	9:Fe:783:ALA:HB2	2.01	0.41
9:Ff:547:GLN:HA	9:Ff:568:TYR:CE1	2.54	0.41
9:Ff:553:ASN:HB3	9:Ff:556:ARG:CZ	2.50	0.41
9:Ff:554:PRO:HB3	9:Ff:587:PRO:HA	2.02	0.41
9:Ff:703:LEU:HA	9:Ff:706:THR:OG1	2.20	0.41
6:Ar:221:TYR:CE1	6:Ax:142:ALA:HB3	2.56	0.41
2:At:6:GLY:CA	6:Bj:125:GLU:HB2	2.50	0.41
3:Au:6:GLY:HA2	6:Bj:132:GLN:HE22	1.85	0.41
5:Aw:219:ARG:HH12	5:Aw:233:ARG:HD3	1.85	0.41
6:Ax:129:LYS:HE2	6:Ax:129:LYS:HB3	1.79	0.41
3:Ba:1:ARG:HG2	3:Ba:2:VAL:N	2.35	0.41
5:Bu:210:TYR:O	5:Bu:262:ILE:HD11	2.20	0.41
6:Bv:168:VAL:C	6:Bv:169:ILE:HD13	2.44	0.41
1:Bw:6:LEU:HD11	5:Ca:246:LEU:HD23	2.02	0.41
6:Ct:112:PHE:CD1	6:Ct:112:PHE:O	2.74	0.41
6:Ct:126:GLN:HB3	6:Cz:112:PHE:CE1	2.55	0.41
6:Cz:129:LYS:O	6:Cz:132:GLN:HG3	2.20	0.41
5:Dk:218:GLY:HA3	6:Dr:148:PRO:HD3	2.02	0.41
5:Dq:207:ARG:NH2	5:Dq:240:GLY:HA3	2.35	0.41
6:Dr:220:LEU:HD12	6:Dx:138:GLU:OE2	2.20	0.41
7:Ef:770:SER:O	7:Ef:774:LEU:HG	2.19	0.41
7:Em:812:TRP:HD1	7:Em:812:TRP:N	2.16	0.41
8:Ew:205:MET:HE1	8:Ew:209:LEU:HG	2.02	0.41
8:Ew:383:LEU:N	8:Fa:424:TYR:OH	2.53	0.41
8:Ex:210:VAL:O	8:Ex:214:VAL:HG23	2.20	0.41
8:Ex:294:LYS:HD3	8:Ex:294:LYS:HA	1.77	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
8:Ex:441:SER:CB	8:Ey:346:ASN:OD1	2.68	0.41
8:Ey:228:PRO:HB3	8:Ey:234:LYS:HG3	2.02	0.41
8:Ey:302:TRP:HD1	8:Ey:303:ASN:HD22	1.67	0.41
8:Ey:426:MET:SD	8:Ez:425:GLN:NE2	2.94	0.41
8:Ey:431:GLN:HA	8:Ey:434:GLU:HG2	2.02	0.41
8:Ez:75:GLN:HB3	8:Fa:400:ILE:HD11	2.02	0.41
8:Fa:290:PRO:HG3	8:Fa:336:TYR:HE2	1.86	0.41
9:Fb:756:LEU:O	9:Fb:760:ILE:HG13	2.20	0.41
9:Fc:268:LYS:HE2	9:Fd:403:PRO:HA	2.02	0.41
9:Fc:274:PHE:CG	9:Fc:515:LEU:HD13	2.56	0.41
9:Fc:500:GLN:HB2	9:Fc:503:LYS:HZ2	1.85	0.41
9:Fc:747:PHE:O	9:Fc:751:TYR:HB2	2.20	0.41
9:Fc:766:PHE:O	9:Fc:766:PHE:HD1	2.03	0.41
9:Fd:179:LYS:HD2	9:Fd:179:LYS:O	2.21	0.41
9:Fe:90:GLU:HG3	9:Fe:91:GLY:N	2.35	0.41
9:Fe:640:VAL:O	9:Fe:644:PHE:CD2	2.73	0.41
9:Fe:678:LEU:HD12	9:Fe:679:THR:HG23	2.01	0.41
6:Af:121:PRO:HB3	7:Ev:800:ASP:OD1	2.20	0.41
6:Ar:170:ARG:HD2	6:Ar:247:ALA:HB3	2.00	0.41
6:Ar:181:PHE:CE2	6:Ar:191:ILE:HD11	2.51	0.41
5:Aw:237:LYS:HD3	5:Aw:243:MET:HB3	2.03	0.41
1:Ay:6:LEU:O	1:Ay:9:LEU:HG	2.21	0.41
6:Bd:115:MET:HE1	7:Eg:802:LEU:HB3	2.02	0.41
1:Be:11:TYR:HE2	3:Bg:11:ALA:HA	1.84	0.41
6:Bj:156:GLN:OE1	6:Bj:156:GLN:N	2.53	0.41
6:Bj:182:LEU:HD11	6:Bj:186:GLY:C	2.45	0.41
6:Bp:219:LYS:HD2	6:Bp:220:LEU:N	2.35	0.41
6:Cb:125:GLU:O	6:Cb:129:LYS:HG3	2.20	0.41
6:Ch:130:LEU:CD2	7:Em:808:LEU:HD12	2.50	0.41
6:Ch:181:PHE:CD1	6:Ch:181:PHE:C	2.97	0.41
3:Ck:4:ILE:HG12	6:Ct:146:GLY:N	2.35	0.41
6:Ct:126:GLN:HG2	6:Ct:129:LYS:HE2	2.02	0.41
1:Cu:2:THR:O	1:Cu:6:LEU:HG	2.20	0.41
2:Cv:1:CYS:HA	3:Cw:4:ILE:HD13	2.01	0.41
6:Df:191:ILE:HD13	6:Df:213:LEU:CD2	2.50	0.41
6:Df:220:LEU:HD23	6:Df:220:LEU:HA	1.83	0.41
6:Df:233:LEU:H	6:Df:233:LEU:HD12	1.85	0.41
6:Di:229:ARG:C	6:Di:229:ARG:HD3	2.45	0.41
6:Dr:170:ARG:HG3	6:Dr:248:VAL:HA	2.03	0.41
6:Dr:173:GLN:OE1	6:Dr:220:LEU:HD23	2.21	0.41
6:Dr:236:PRO:HG2	6:Dr:238:MET:CE	2.47	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:Du:1:ARG:HG2	3:Du:2:VAL:N	2.35	0.41
7:Ej:790:TYR:CE2	7:Ej:794:ILE:HD11	2.55	0.41
7:Es:799:SER:O	7:Es:802:LEU:HD12	2.20	0.41
8:Ey:170:LEU:HD21	8:Ey:175:SER:HB2	2.02	0.41
8:Ey:427:TYR:C	8:Ey:427:TYR:CD2	2.97	0.41
8:Fa:102:VAL:HB	8:Fa:106:VAL:HG21	2.03	0.41
8:Fa:206:TYR:CE2	8:Fa:209:LEU:HG	2.55	0.41
9:Fb:517:CYS:SG	9:Fb:522:ASN:HA	2.60	0.41
9:Fb:768:TRP:NE1	9:Fb:812:PRO:HG3	2.36	0.41
9:Fc:78:TYR:O	9:Fc:82:VAL:HG13	2.20	0.41
9:Fc:339:MET:HA	9:Fc:342:THR:HG22	2.02	0.41
9:Fc:346:VAL:O	9:Fc:349:VAL:HG12	2.20	0.41
9:Fc:530:ILE:O	9:Fc:534:ILE:HG13	2.20	0.41
9:Fc:766:PHE:CE2	9:Fd:925:ILE:HG23	2.55	0.41
9:Fd:510:GLN:N	9:Fd:510:GLN:OE1	2.53	0.41
9:Fe:642:ASN:HA	9:Fe:645:LEU:HD12	2.03	0.41
9:Fe:800:PHE:O	9:Fe:803:MET:HE3	2.21	0.41
9:Ff:140:ASP:OD1	9:Ff:461:TRP:HD1	2.03	0.41
9:Ff:686:ILE:O	9:Ff:689:LEU:HD12	2.21	0.41
7:Fh:35:GLY:HA2	7:Fh:38:LYS:HZ3	1.85	0.41
6:Ax:181:PHE:CE2	6:Ax:228:VAL:HG11	2.55	0.41
6:Ax:191:ILE:HB	6:Ax:206:TRP:CZ2	2.55	0.41
5:Bi:219:ARG:NH2	6:Bp:150:LYS:HD2	2.35	0.41
6:Bj:166:PRO:HG3	6:Bp:250:TYR:CE1	2.55	0.41
6:Bp:149:PRO:HB3	6:Bp:247:ALA:C	2.46	0.41
1:Bw:16:ALA:HB3	2:Cd:9:ARG:HG2	2.02	0.41
5:Ca:233:ARG:CZ	5:Ca:236:SER:HB2	2.50	0.41
6:Cb:123:ASN:HB3	6:Cb:126:GLN:HE22	1.86	0.41
6:Cn:115:MET:HE1	7:Em:798:THR:HA	2.01	0.41
6:Cn:214:MET:HE1	7:Eo:818:GLN:NE2	2.26	0.41
6:Cn:238:MET:HE2	6:Ct:250:TYR:CD2	2.55	0.41
6:Ct:125:GLU:O	6:Ct:129:LYS:HG3	2.20	0.41
6:Ct:170:ARG:HB3	6:Ct:247:ALA:O	2.20	0.41
5:Cy:214:ALA:HB1	6:Cz:242:ILE:HG21	2.02	0.41
6:Cz:220:LEU:HD13	6:Df:138:GLU:CD	2.45	0.41
6:Dl:135:GLU:HA	6:Dl:138:GLU:OE2	2.20	0.41
6:Dl:207:ASP:OD2	6:Dl:207:ASP:C	2.63	0.41
6:Dr:108:ASP:HB3	6:Dr:112:PHE:CZ	2.54	0.41
6:Dr:148:PRO:HA	6:Dr:149:PRO:HD3	1.95	0.41
5:Ec:233:ARG:HG3	5:Ec:234:GLU:H	1.86	0.41
7:Em:799:SER:HA	7:Em:802:LEU:HD23	2.02	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
7:Em:805:ALA:HA	7:Em:808:LEU:HG	2.02	0.41
7:Ep:806:THR:O	7:Ep:810:GLN:HG3	2.21	0.41
7:Er:799:SER:HB2	9:Fb:363:THR:CB	2.51	0.41
7:Et:810:GLN:HA	7:Et:813:LYS:NZ	2.36	0.41
7:Eu:780:LYS:O	7:Eu:784:GLN:HG2	2.19	0.41
8:Ew:39:LEU:CD1	8:Ex:39:LEU:HD11	2.48	0.41
8:Ew:228:PRO:HG2	8:Ew:235:TYR:HB2	2.02	0.41
8:Ex:330:PHE:HB2	8:Ex:334:ARG:HH12	1.84	0.41
8:Ex:349:TYR:CZ	8:Ex:353:LYS:HD3	2.56	0.41
8:Ex:432:ILE:HA	8:Ex:435:ARG:HG2	2.02	0.41
8:Ey:443:MET:SD	8:Ey:444:LEU:HD23	2.60	0.41
8:Fa:139:ASN:O	8:Fa:142:ILE:HG13	2.21	0.41
8:Fa:349:TYR:O	8:Fa:349:TYR:HD1	2.03	0.41
9:Fb:268:LYS:C	9:Fb:268:LYS:HD2	2.46	0.41
9:Fb:608:GLN:OE1	9:Fb:608:GLN:N	2.53	0.41
9:Fb:769:LEU:O	9:Fb:773:ILE:HG13	2.20	0.41
9:Fb:818:GLY:C	9:Fb:933:ILE:HG13	2.45	0.41
9:Fc:65:ASN:O	9:Fc:68:VAL:HG12	2.20	0.41
9:Fc:195:GLN:HA	9:Fc:198:LEU:HD12	2.02	0.41
9:Fc:527:LEU:O	9:Fc:531:GLN:HG2	2.20	0.41
9:Fd:768:TRP:HB2	9:Fd:811:ARG:NH2	2.35	0.41
9:Ff:194:LEU:HA	9:Ff:197:GLN:HG3	2.02	0.41
7:Fh:25:ALA:HA	7:Fh:28:LEU:HG	2.01	0.41
7:Fj:15:THR:HA	7:Fj:18:ARG:CD	2.51	0.41
6:Af:156:GLN:HE22	6:Af:252:VAL:HB	1.86	0.41
6:Al:123:ASN:ND2	6:Al:124:PRO:HD2	2.36	0.41
6:Al:191:ILE:HG13	6:Al:211:ASN:HA	2.02	0.41
6:Al:245:GLN:CD	6:Al:247:ALA:H	2.28	0.41
6:Ar:250:TYR:O	6:Ar:251:ARG:HG2	2.20	0.41
3:Au:2:VAL:HG21	6:Bj:129:LYS:HG2	2.03	0.41
6:Ax:124:PRO:O	6:Ax:128:VAL:HG12	2.21	0.41
6:Ax:203:ASN:HB3	6:Ax:216:GLN:OE1	2.21	0.41
5:Bi:238:ILE:HD11	5:Bi:244:VAL:HB	2.02	0.41
5:Bi:246:LEU:HD12	5:Bi:247:ILE:H	1.85	0.41
6:Bj:228:VAL:CG1	6:Bj:237:VAL:HB	2.48	0.41
6:Bv:195:ASP:O	6:Bv:226:LEU:HD12	2.20	0.41
6:Cb:251:ARG:HE	6:Cb:251:ARG:HB3	1.55	0.41
6:Ch:113:LYS:NZ	6:Ch:117:ARG:HB2	2.36	0.41
6:Ch:187:ALA:HB1	6:Ch:262:ASN:HB2	2.02	0.41
6:Cn:123:ASN:HB2	6:Cn:126:GLN:OE1	2.21	0.41
6:Dl:184:SER:HA	6:Dl:255:ARG:NH2	2.35	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:Do:5:GLY:HA3	4:Dv:1:PRO:HB3	2.02	0.41
6:Dr:191:ILE:HD13	6:Dr:213:LEU:HG	2.01	0.41
2:Dt:1:CYS:HA	3:Du:4:ILE:HD13	2.02	0.41
6:Ed:115:MET:HE1	7:Et:801:MET:HB3	2.03	0.41
6:Ed:129:LYS:O	6:Ed:132:GLN:HG3	2.20	0.41
7:Em:800:ASP:OD2	7:Em:800:ASP:C	2.64	0.41
7:Ev:793:GLU:HA	7:Ev:796:GLN:OE1	2.21	0.41
8:Ew:77:VAL:HG12	8:Ew:378:MET:SD	2.61	0.41
8:Ey:270:LYS:HD2	8:Ey:270:LYS:N	2.36	0.41
8:Ez:307:ALA:HB3	8:Ez:319:GLN:HG3	2.01	0.41
8:Ez:354:ARG:HG3	8:Ez:372:ALA:HB3	2.01	0.41
8:Ez:430:ARG:CZ	8:Fa:425:GLN:HG3	2.50	0.41
8:Fa:242:GLN:HE21	8:Fa:269:ALA:HB3	1.86	0.41
8:Fa:322:ALA:O	8:Fa:326:LEU:HG	2.20	0.41
9:Fb:237:VAL:HA	9:Fb:265:ASN:ND2	2.34	0.41
9:Fb:521:GLN:O	9:Fb:521:GLN:CD	2.63	0.41
9:Fc:112:ALA:O	9:Fc:123:MET:HE3	2.20	0.41
9:Fc:224:THR:O	9:Fc:228:THR:HG22	2.20	0.41
9:Fc:594:ILE:HD12	9:Fc:594:ILE:H	1.85	0.41
9:Fc:628:MET:O	9:Fc:632:PHE:HD1	2.02	0.41
9:Fc:735:LEU:O	9:Fc:739:ILE:HB	2.20	0.41
9:Fd:469:PHE:CD1	9:Fd:469:PHE:C	2.99	0.41
9:Fd:751:TYR:C	9:Fd:754:PRO:HD2	2.45	0.41
9:Fe:263:MET:HE3	9:Fe:279:CYS:O	2.20	0.41
9:Ff:130:TRP:CD1	9:Ff:130:TRP:C	2.98	0.41
7:Fi:19:VAL:HG23	7:Fi:20:ILE:HD13	2.02	0.41
7:Fj:30:ILE:HD13	7:Fj:30:ILE:HA	1.89	0.41
5:Ae:254:ILE:HD13	5:Ae:254:ILE:HA	1.85	0.41
6:Af:198:ASP:OD2	6:Af:201:SER:HB3	2.20	0.41
6:Af:218:THR:O	7:Ee:815:VAL:HG21	2.20	0.41
6:Al:226:LEU:CD1	6:Al:239:LEU:HB2	2.51	0.41
6:Al:237:VAL:C	6:Ar:251:ARG:HH22	2.27	0.41
6:Ar:194:TYR:CD2	6:Ar:226:LEU:HD11	2.56	0.41
6:Bd:126:GLN:O	6:Bd:129:LYS:HB2	2.20	0.41
6:Bj:225:ASN:ND2	6:Bp:175:PHE:HD1	2.18	0.41
1:Bk:4:ALA:H	2:Bl:9:ARG:NH2	2.19	0.41
6:Bv:126:GLN:HA	6:Bv:129:LYS:HE2	2.02	0.41
6:Bv:220:LEU:HD12	6:Cb:138:GLU:OE1	2.20	0.41
5:Cg:229:THR:HG23	6:Ch:165:THR:OG1	2.21	0.41
6:Cn:131:LYS:HG3	7:En:812:TRP:CH2	2.56	0.41
2:Cp:5:ILE:HD12	6:Cz:139:TYR:HB3	2.02	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
6:Cz:181:PHE:HZ	6:Cz:213:LEU:HG	1.86	0.41
5:Dk:269:SER:HA	6:Dl:246:LYS:NZ	2.34	0.41
6:Dl:123:ASN:HB2	6:Dl:126:GLN:HG2	2.02	0.41
6:Dx:177:SER:HB3	6:Dx:215:ILE:HG23	2.03	0.41
7:Ei:781:GLN:HA	7:Ei:784:GLN:HG2	2.03	0.41
7:Eq:775:GLN:HG2	9:Fb:327:THR:OG1	2.20	0.41
7:Es:778:LEU:HA	7:Es:781:GLN:CD	2.46	0.41
8:Ey:342:VAL:HG11	8:Ey:442:ILE:HG13	2.02	0.41
8:Ez:177:ALA:HA	8:Ez:286:GLN:NE2	2.35	0.41
9:Fb:111:LEU:HD13	7:Fh:31:ALA:HB2	2.02	0.41
9:Fb:751:TYR:C	9:Fb:754:PRO:HD2	2.45	0.41
9:Fb:760:ILE:HG13	9:Fb:760:ILE:H	1.73	0.41
9:Fb:939:LYS:HE2	9:Fb:943:TRP:CZ2	2.54	0.41
9:Fc:187:GLY:O	9:Fc:191:MET:SD	2.79	0.41
9:Fc:391:GLU:O	9:Fc:394:GLN:HG3	2.20	0.41
9:Fc:648:PHE:CE1	9:Fc:652:ILE:HG12	2.55	0.41
9:Fd:596:PHE:CE2	9:Fd:598:VAL:HG22	2.55	0.41
9:Fd:736:MET:HA	9:Fd:739:ILE:CG1	2.51	0.41
9:Fe:34:SER:HA	9:Fe:462:ILE:HD13	2.02	0.41
9:Fe:652:ILE:HG13	9:Fe:652:ILE:H	1.63	0.41
9:Fe:729:MET:HA	9:Fe:732:MET:CG	2.47	0.41
9:Fe:763:PHE:CE2	9:Ff:924:ILE:HD11	2.56	0.41
9:Ff:239:PHE:CD1	9:Ff:336:ILE:HG13	2.56	0.41
9:Ff:768:TRP:O	9:Ff:772:VAL:HG23	2.21	0.41
4:Ad:4:ALA:HB3	5:Ae:251:GLN:NE2	2.36	0.41
6:Af:190:PRO:HA	6:Af:211:ASN:HB3	2.02	0.41
6:Af:238:MET:HB3	6:Al:250:TYR:CD2	2.55	0.41
2:Ah:5:ILE:HD13	6:Ar:140:ALA:N	2.36	0.41
3:Ai:1:ARG:HH22	3:Ai:3:SER:HA	1.86	0.41
5:Ak:219:ARG:NH1	5:Ak:233:ARG:HB3	2.35	0.41
6:Al:236:PRO:HG2	6:Ar:251:ARG:NE	2.36	0.41
5:Aw:216:ILE:HD11	6:Bd:148:PRO:HG2	2.01	0.41
6:Ax:176:VAL:CG1	6:Ax:214:MET:HG3	2.48	0.41
6:Bd:132:GLN:O	6:Bd:135:GLU:HG2	2.21	0.41
3:Bg:10:THR:HG22	3:Bg:11:ALA:O	2.20	0.41
6:Bj:181:PHE:HZ	6:Bj:213:LEU:HB2	1.85	0.41
6:Bp:126:GLN:HB3	6:Bv:112:PHE:CZ	2.55	0.41
6:Bp:141:LYS:CD	7:Ek:808:LEU:HD12	2.46	0.41
5:Cg:219:ARG:HD3	6:Cn:148:PRO:O	2.21	0.41
6:Cz:112:PHE:N	6:Cz:115:MET:HE3	2.35	0.41
6:Df:134:TYR:O	6:Df:138:GLU:HB3	2.21	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
6:Df:179:LEU:HD13	6:Df:239:LEU:HD22	2.02	0.41
6:Dl:149:PRO:HB2	6:Dl:248:VAL:HG12	2.03	0.41
6:Dl:182:LEU:HD12	6:Dl:186:GLY:C	2.45	0.41
6:Dl:238:MET:HG2	6:Dr:250:TYR:HD2	1.86	0.41
2:Dn:2:VAL:HG13	3:Do:1:ARG:HE	1.85	0.41
6:Dx:173:GLN:HB2	6:Dx:220:LEU:HD13	2.03	0.41
7:Ei:793:GLU:O	7:Ei:796:GLN:HG3	2.21	0.41
7:Em:780:LYS:HA	7:Em:783:GLU:OE1	2.21	0.41
7:Eq:794:ILE:HB	7:Eq:795:GLN:NE2	2.36	0.41
8:Ew:298:TYR:CZ	8:Ex:230:PHE:CE2	3.08	0.41
8:Ex:349:TYR:CE1	8:Ex:353:LYS:HD3	2.55	0.41
8:Ex:442:ILE:HA	8:Ex:445:LEU:HD23	2.02	0.41
8:Ey:423:ASN:O	8:Ey:426:MET:SD	2.79	0.41
8:Ez:94:MET:N	8:Ez:94:MET:SD	2.93	0.41
8:Fa:142:ILE:HG13	8:Fa:142:ILE:H	1.75	0.41
8:Fa:353:LYS:O	8:Fa:371:GLN:HB3	2.20	0.41
9:Fb:392:TYR:HB2	9:Fb:576:MET:HE3	2.01	0.41
9:Fb:510:GLN:CG	9:Fb:511:ASP:H	2.33	0.41
9:Fb:520:PHE:CE2	9:Fb:526:LYS:HB2	2.50	0.41
9:Fc:132:ILE:O	9:Fc:136:VAL:HG23	2.21	0.41
9:Fc:461:TRP:CD1	9:Fc:461:TRP:N	2.87	0.41
9:Fc:462:ILE:HA	9:Fc:761:PHE:CZ	2.56	0.41
9:Fc:734:LEU:HB3	9:Fc:738:TRP:CZ2	2.56	0.41
9:Fc:928:LYS:HG3	9:Fc:929:ALA:H	1.84	0.41
9:Fd:427:ASN:HA	9:Fd:430:MET:CE	2.51	0.41
9:Fd:588:LEU:HD11	9:Fd:657:MET:HE1	2.02	0.41
9:Fd:697:ILE:HG22	9:Fd:746:GLY:HA3	2.03	0.41
9:Fd:822:ALA:HB1	9:Fd:930:PHE:CD1	2.55	0.41
9:Fe:143:TRP:CE2	9:Fe:147:LEU:HD11	2.56	0.41
9:Fe:648:PHE:CE1	9:Fe:652:ILE:HG12	2.56	0.41
9:Ff:117:LYS:NZ	9:Ff:119:SER:H	2.19	0.41
9:Ff:612:CYS:HA	9:Ff:626:ARG:HB2	2.02	0.41
9:Ff:648:PHE:O	9:Ff:652:ILE:HG12	2.20	0.41
9:Ff:694:THR:O	9:Ff:698:ASN:HB2	2.19	0.41
6:Af:181:PHE:HD2	6:Af:211:ASN:O	2.04	0.41
6:Af:204:ILE:HD12	6:Af:214:MET:O	2.21	0.41
6:Ax:225:ASN:HD21	6:Bd:176:VAL:H	1.67	0.41
6:Bd:158:VAL:HB	6:Bd:256:VAL:HG13	2.03	0.41
6:Bj:179:LEU:HD12	6:Bj:252:VAL:HB	2.03	0.41
6:Bp:113:LYS:NZ	6:Bp:117:ARG:HG3	2.36	0.41
6:Bp:181:PHE:CE2	6:Bp:213:LEU:HD12	2.56	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
5:Ca:253:ARG:HB2	5:Ca:263:LYS:HE3	2.03	0.41
6:Cb:121:PRO:HG2	7:El:801:MET:HB3	2.03	0.41
6:Cb:132:GLN:HE21	6:Cb:136:THR:HG23	1.85	0.41
5:Cs:237:LYS:HA	5:Cs:237:LYS:HD3	1.94	0.41
6:Ct:111:ALA:HB2	7:En:794:ILE:CG2	2.50	0.41
6:Ct:171:LEU:HD12	6:Ct:172:SER:N	2.32	0.41
5:Cy:245:LYS:HD3	5:Cy:257:SER:HB3	2.03	0.41
6:Df:141:LYS:HD3	7:Er:812:TRP:HZ3	1.86	0.41
6:Dr:221:TYR:HA	6:Dr:243:PRO:HG2	2.01	0.41
7:Ej:809:VAL:O	7:Ej:813:LYS:HE3	2.19	0.41
8:Ew:106:VAL:HG22	8:Ew:109:ASN:HB3	2.03	0.41
8:Ew:177:ALA:O	8:Ew:181:ILE:HG13	2.20	0.41
8:Ew:191:MET:HE2	8:Ew:195:GLN:HA	2.02	0.41
8:Ey:70:ASN:HB3	8:Ey:73:VAL:HG22	2.01	0.41
8:Ey:385:ASP:HB3	8:Ey:388:ALA:HB2	2.02	0.41
8:Ez:422:ILE:HA	8:Ez:425:GLN:CD	2.45	0.41
9:Fb:132:ILE:HG12	9:Fb:772:VAL:HG23	2.03	0.41
9:Fb:734:LEU:HD11	9:Fb:738:TRP:CZ2	2.56	0.41
9:Fb:768:TRP:HD1	9:Fb:811:ARG:NE	2.19	0.41
9:Fc:392:TYR:OH	9:Fc:585:ILE:HB	2.20	0.41
9:Fc:673:VAL:HA	9:Fc:676:GLN:HE21	1.85	0.41
9:Fc:692:MET:N	9:Fc:692:MET:SD	2.94	0.41
9:Fc:732:MET:O	9:Fc:736:MET:HG2	2.21	0.41
9:Fc:905:TRP:HB3	9:Fc:909:TYR:CZ	2.56	0.41
9:Fd:56:ILE:HG13	9:Fd:57:MET:SD	2.61	0.41
9:Fd:241:LYS:HB3	9:Fd:241:LYS:HZ2	1.86	0.41
9:Fd:769:LEU:HA	9:Fd:772:VAL:HG12	2.02	0.41
9:Fd:797:LYS:HB3	9:Fd:800:PHE:CZ	2.56	0.41
9:Fe:751:TYR:C	9:Fe:754:PRO:HD2	2.45	0.41
9:Fe:917:ILE:HA	9:Fe:920:SER:OG	2.20	0.41
9:Ff:116:PRO:HB2	9:Ff:120:GLY:HA2	2.02	0.41
7:Fg:30:ILE:HD13	7:Fg:30:ILE:HA	1.93	0.41
5:Ae:211:TYR:C	5:Ae:222:LEU:HD13	2.45	0.41
6:Al:157:PHE:CD1	6:Al:157:PHE:N	2.86	0.41
6:Al:166:PRO:HA	6:Al:167:PRO:HD3	1.94	0.41
6:Ar:225:ASN:HB2	6:Ax:250:TYR:HH	1.86	0.41
2:Az:4:MET:HE1	5:Bc:250:LEU:H	1.86	0.41
6:Bd:156:GLN:HE22	6:Bd:158:VAL:CG2	2.33	0.41
6:Bd:169:ILE:HD13	6:Bd:252:VAL:HG21	2.03	0.41
6:Bd:196:LEU:HD22	6:Bd:204:ILE:HG22	2.02	0.41
2:Bf:6:GLY:HA3	6:Bv:125:GLU:CD	2.46	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
6:Bj:178:SER:HB3	6:Bj:251:ARG:HG2	2.03	0.41
6:Bj:223:TYR:CZ	6:Bp:175:PHE:HZ	2.39	0.41
6:Bp:120:TYR:C	6:Bp:122:LEU:H	2.29	0.41
6:Bp:221:TYR:HA	6:Bp:243:PRO:HG2	2.03	0.41
1:Bq:1:CYS:HA	4:Bt:2:PHE:CD2	2.56	0.41
5:Bu:214:ALA:HB3	5:Bu:221:TRP:CE3	2.56	0.41
6:Bv:106:VAL:HA	6:Bv:109:LYS:NZ	2.36	0.41
6:Bv:157:PHE:C	6:Bv:157:PHE:CD2	2.99	0.41
5:Ca:221:TRP:CZ3	6:Cb:242:ILE:HG21	2.56	0.41
6:Cb:162:PRO:HA	6:Ch:251:ARG:HH22	1.85	0.41
6:Cb:173:GLN:HA	6:Cb:217:ALA:O	2.21	0.41
2:Cd:4:MET:HE1	5:Cg:250:LEU:H	1.86	0.41
6:Ch:181:PHE:HA	6:Ch:254:LEU:HD11	2.02	0.41
1:Ci:3:ASP:OD1	3:Ck:16:ASP:HB2	2.21	0.41
6:Cn:130:LEU:HA	6:Cn:133:ILE:HG12	2.03	0.41
1:Co:3:ASP:OD2	3:Cq:14:TYR:HB3	2.21	0.41
6:Ct:108:ASP:CA	7:En:797:ARG:HH21	2.34	0.41
6:Ct:170:ARG:HA	6:Ct:245:GLN:OE1	2.21	0.41
6:Ct:229:ARG:NH1	6:Ct:230:LEU:H	2.18	0.41
3:Cw:4:ILE:HD11	6:Df:146:GLY:H	1.85	0.41
5:Cy:208:ILE:HD11	5:Cy:225:SER:H	1.85	0.41
5:Cy:212:ILE:HG12	5:Cy:254:ILE:HD11	2.03	0.41
6:Cz:158:VAL:HG13	6:Cz:167:PRO:HG2	2.03	0.41
6:Df:220:LEU:HD12	6:Dl:138:GLU:CD	2.46	0.41
5:Dk:268:ASP:HA	6:Dl:150:LYS:HD2	2.03	0.41
6:Dr:110:LYS:HD3	6:Dr:110:LYS:C	2.46	0.41
6:Dr:125:GLU:O	6:Dr:128:VAL:HG22	2.20	0.41
6:Dr:156:GLN:O	6:Dr:254:LEU:HA	2.20	0.41
6:Dr:206:TRP:CB	6:Dr:213:LEU:HD23	2.50	0.41
6:Dx:236:PRO:HG2	6:Dx:238:MET:SD	2.61	0.41
3:Ea:3:SER:OG	3:Ea:9:TYR:HD2	2.04	0.41
5:Ec:238:ILE:HD11	5:Ec:244:VAL:CG2	2.51	0.41
6:Ed:194:TYR:CD2	6:Ed:194:TYR:C	2.99	0.41
7:Ej:781:GLN:O	7:Ej:785:LEU:HG	2.21	0.41
7:El:793:GLU:O	7:El:797:ARG:HG2	2.20	0.41
7:Em:782:ASN:HA	7:Em:785:LEU:CD2	2.50	0.41
7:Eq:808:LEU:HA	7:Eq:811:ASP:OD1	2.21	0.41
7:Es:806:THR:O	7:Es:810:GLN:HG3	2.21	0.41
8:Ew:45:ASN:O	8:Ew:49:TYR:HD1	2.04	0.41
8:Ew:336:TYR:CD1	8:Fa:453:PRO:HD3	2.55	0.41
8:Ew:382:ARG:HB2	8:Fa:424:TYR:OH	2.21	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
8:Ey:142:ILE:HD11	8:Ey:281:ARG:NH1	2.35	0.41
8:Ey:211:MET:O	8:Ey:215:ILE:HG12	2.21	0.41
8:Ey:230:PHE:HA	8:Ey:235:TYR:CD2	2.56	0.41
8:Ey:277:LEU:HD13	8:Ey:281:ARG:CZ	2.50	0.41
8:Ey:298:TYR:CE2	8:Ez:230:PHE:HE1	2.39	0.41
8:Ey:391:THR:HB	8:Ey:394:GLN:CD	2.46	0.41
8:Ey:415:ILE:HG13	8:Ey:415:ILE:H	1.76	0.41
8:Ey:418:LEU:HD12	8:Ey:419:LEU:N	2.36	0.41
8:Ez:415:ILE:HA	8:Ez:418:LEU:CG	2.51	0.41
8:Fa:77:VAL:HG12	8:Fa:378:MET:CE	2.50	0.41
8:Fa:206:TYR:N	8:Fa:209:LEU:HD12	2.27	0.41
8:Fa:448:LEU:HA	8:Fa:451:ALA:HB3	2.02	0.41
9:Fb:24:LEU:HD23	9:Fb:24:LEU:HA	1.93	0.41
9:Fb:212:PRO:HA	9:Fb:215:GLY:H	1.85	0.41
9:Fb:689:LEU:HD13	9:Fb:922:TYR:CE2	2.56	0.41
9:Fb:837:PHE:O	9:Fb:841:ILE:HG12	2.21	0.41
9:Fc:314:GLY:HA3	9:Fc:439:GLN:O	2.21	0.41
9:Fc:338:GLN:HA	9:Fc:341:VAL:HG12	2.03	0.41
9:Fc:766:PHE:CG	9:Fd:928:LYS:NZ	2.83	0.41
9:Fd:129:MET:HA	9:Fd:132:ILE:HG12	2.03	0.41
9:Fd:668:LYS:HB3	9:Fd:668:LYS:HE2	1.84	0.41
9:Fd:813:SER:O	9:Fd:816:ILE:HG22	2.21	0.41
9:Fe:24:LEU:N	7:Fk:36:PHE:CE2	2.89	0.41
9:Fe:766:PHE:CD1	9:Fe:766:PHE:C	2.98	0.41
9:Fe:790:GLU:H	9:Fe:790:GLU:HG3	1.67	0.41
9:Ff:65:ASN:HA	9:Ff:68:VAL:HG22	2.03	0.41
9:Ff:203:ASP:O	9:Ff:206:LEU:HD12	2.21	0.41
9:Ff:236:THR:HG21	9:Ff:266:PHE:CE1	2.56	0.41
9:Ff:249:LYS:HD2	9:Ff:249:LYS:HA	1.94	0.41
9:Ff:296:SER:HA	9:Ff:302:VAL:HG13	2.02	0.41
9:Ff:573:ASN:HA	9:Ff:576:MET:SD	2.60	0.41
9:Ff:701:GLY:O	9:Ff:705:LEU:HD23	2.21	0.41
7:Fk:32:VAL:HG13	7:Fk:36:PHE:HE1	1.79	0.41
5:Ae:219:ARG:NH2	6:Al:150:LYS:HD2	2.36	0.41
3:Au:6:GLY:HA2	6:Bj:132:GLN:NE2	2.36	0.41
5:Aw:222:LEU:HD11	5:Aw:262:ILE:CG2	2.49	0.41
6:Bd:203:ASN:O	6:Bd:215:ILE:HD12	2.21	0.41
5:Bi:268:ASP:O	6:Bj:247:ALA:HB2	2.21	0.41
5:Bo:237:LYS:HD3	5:Bo:237:LYS:HA	1.86	0.41
6:Bp:199:PRO:HD2	7:El:816:GLU:HG2	2.02	0.41
6:Cb:115:MET:SD	7:Ek:802:LEU:HD12	2.61	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
6:Ch:183:ASP:HB3	6:Ch:189:TRP:CE3	2.56	0.41
6:Ch:229:ARG:HG2	6:Ch:236:PRO:HB3	2.02	0.41
5:Cm:213:GLN:HB3	5:Cm:221:TRP:O	2.21	0.41
5:Cs:252:GLY:O	5:Cs:253:ARG:HD2	2.21	0.41
5:Dk:210:TYR:CE2	5:Dk:238:ILE:HD11	2.56	0.41
5:Dk:254:ILE:HD13	5:Dk:254:ILE:HA	1.95	0.41
5:Dq:210:TYR:HD2	5:Dq:224:GLY:HA2	1.86	0.41
3:Ea:1:ARG:HG2	3:Ea:2:VAL:N	2.35	0.41
6:Ed:135:GLU:HA	6:Ed:138:GLU:OE1	2.20	0.41
7:Eo:814:GLN:HE22	7:Eo:816:GLU:HB3	1.86	0.41
7:Et:781:GLN:C	7:Et:781:GLN:CD	2.89	0.41
8:Ew:119:THR:OG1	8:Ew:355:LEU:HD12	2.20	0.41
8:Ex:436:ILE:HD12	8:Ex:436:ILE:H	1.86	0.41
8:Ex:438:LEU:HG	8:Ex:442:ILE:HD11	2.03	0.41
8:Ey:175:SER:HB3	8:Ez:244:ASN:ND2	2.36	0.41
8:Fa:43:LEU:O	8:Fa:46:LEU:HG	2.21	0.41
8:Fa:248:LEU:HD23	8:Fa:279:PHE:CD2	2.47	0.41
9:Fb:779:ALA:O	9:Fb:782:VAL:HG22	2.21	0.41
9:Fb:831:TRP:CE3	9:Ff:37:PHE:HE2	2.39	0.41
9:Fc:596:PHE:H	9:Fc:597:LYS:HZ3	1.69	0.41
9:Fc:722:ILE:O	9:Fc:725:PHE:HB3	2.21	0.41
9:Fd:189:ILE:CD1	9:Fd:567:VAL:HG11	2.51	0.41
9:Fd:586:LYS:HD3	9:Fd:586:LYS:O	2.20	0.41
9:Fd:747:PHE:HA	9:Fd:751:TYR:HD1	1.86	0.41
9:Fd:920:SER:O	9:Fd:923:LEU:HB3	2.20	0.41
9:Fd:923:LEU:HD23	9:Fd:924:ILE:N	2.36	0.41
9:Fd:929:ALA:O	9:Fd:932:LEU:HD13	2.22	0.41
9:Fe:751:TYR:HB3	9:Ff:913:PHE:CD1	2.56	0.41
9:Fe:913:PHE:HA	9:Fe:916:LEU:HD23	2.03	0.41
9:Ff:338:GLN:NE2	9:Ff:429:ILE:HD12	2.36	0.41
9:Ff:635:TYR:O	9:Ff:639:TYR:HB3	2.22	0.41
9:Ff:801:ALA:O	9:Ff:804:ILE:HG12	2.21	0.41
7:Fj:38:LYS:O	7:Fj:38:LYS:HD2	2.21	0.41
6:Al:106:VAL:HG23	6:Al:110:LYS:HZ2	1.86	0.40
6:Al:132:GLN:O	6:Al:136:THR:HG23	2.21	0.40
5:Aq:222:LEU:HD23	5:Aq:230:LEU:O	2.21	0.40
6:Ar:242:ILE:HG13	6:Ar:245:GLN:HG2	2.03	0.40
6:Bd:113:LYS:HD3	6:Bd:113:LYS:HA	1.66	0.40
5:Bo:262:ILE:C	5:Bo:263:LYS:HD2	2.46	0.40
6:Bp:205:GLN:HB2	6:Bp:214:MET:SD	2.61	0.40
5:Bu:247:ILE:HA	5:Bu:254:ILE:HG12	2.03	0.40

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
5:Ca:262:ILE:HD13	5:Ca:262:ILE:HA	1.89	0.40
6:Cb:114:ASP:O	6:Cb:118:ASN:OD1	2.40	0.40
5:Cg:212:ILE:HG12	5:Cg:254:ILE:HD11	2.02	0.40
5:Cy:221:TRP:HH2	6:Cz:165:THR:HB	1.85	0.40
6:Cz:190:PRO:O	6:Cz:231:ARG:HG2	2.21	0.40
5:Dw:267:GLU:OE2	5:Dw:267:GLU:N	2.50	0.40
6:Dx:189:TRP:CZ2	6:Dx:260:GLY:HA2	2.57	0.40
5:Ec:263:LYS:HE2	5:Ec:263:LYS:HB2	1.89	0.40
6:Ed:173:GLN:HG3	6:Ed:220:LEU:HB3	2.02	0.40
6:Ed:229:ARG:HB3	6:Ed:236:PRO:HB3	2.03	0.40
7:Et:782:ASN:O	7:Et:785:LEU:HD12	2.20	0.40
8:Ew:426:MET:HE2	8:Ew:426:MET:C	2.46	0.40
8:Ew:427:TYR:CA	8:Ew:430:ARG:HG3	2.50	0.40
8:Ex:69:PHE:HZ	8:Ex:406:ALA:HB2	1.85	0.40
8:Ex:96:GLN:HA	8:Ex:99:MET:SD	2.61	0.40
8:Ex:304:GLN:OE1	8:Ex:319:GLN:HA	2.21	0.40
8:Ex:359:MET:SD	8:Ex:369:THR:HG22	2.61	0.40
8:Ex:426:MET:O	8:Ex:430:ARG:HG3	2.21	0.40
8:Ey:217:THR:H	8:Ey:265:SER:HB3	1.85	0.40
8:Ey:414:GLU:HA	8:Ey:417:ILE:HG12	2.02	0.40
8:Ez:179:PHE:CZ	8:Fa:236:ASN:HB2	2.56	0.40
8:Fa:93:ALA:HA	8:Fa:99:MET:HG2	2.02	0.40
8:Fa:378:MET:N	8:Fa:378:MET:SD	2.94	0.40
9:Fb:28:PRO:HB2	9:Fb:32:ASP:OD2	2.20	0.40
9:Fb:66:SER:OG	7:Fh:38:LYS:HE2	2.21	0.40
9:Fb:82:VAL:O	9:Fb:85:MET:HG3	2.21	0.40
9:Fb:239:PHE:CE2	9:Fb:340:TYR:HB2	2.55	0.40
9:Fb:617:ILE:O	9:Fb:618:LEU:C	2.65	0.40
9:Fb:706:THR:O	9:Fb:710:MET:HG3	2.22	0.40
9:Fb:723:PHE:O	9:Fb:726:ALA:HB3	2.21	0.40
9:Fb:741:THR:HG22	9:Fb:745:ILE:HD11	2.03	0.40
9:Fc:920:SER:O	9:Fc:923:LEU:HB3	2.20	0.40
9:Fd:192:LEU:CB	9:Fd:351:VAL:HG21	2.48	0.40
9:Fd:218:PRO:HB2	9:Fd:223:ASN:CG	2.46	0.40
9:Fd:643:PHE:CE2	9:Fd:647:ILE:HD11	2.55	0.40
9:Fd:760:ILE:HD13	9:Fd:763:PHE:HE2	1.85	0.40
9:Fe:113:LEU:HD11	9:Fe:124:MET:HB2	2.02	0.40
9:Fe:685:PRO:O	9:Fe:689:LEU:HG	2.21	0.40
9:Fe:732:MET:N	9:Fe:733:PRO:HD2	2.36	0.40
9:Ff:94:LEU:HD21	7:Fi:21:ILE:HD11	2.03	0.40
9:Ff:127:PHE:O	9:Ff:131:VAL:HG23	2.21	0.40

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
9:Ff:290:LEU:HD11	9:Ff:437:ILE:HD12	2.03	0.40
9:Ff:810:LEU:O	9:Ff:814:LEU:HG	2.20	0.40
6:Af:230:LEU:HB3	6:Af:233:LEU:HD13	2.03	0.40
5:Ak:237:LYS:HA	5:Ak:237:LYS:HD3	1.89	0.40
5:Ak:251:GLN:HB3	5:Ak:253:ARG:NH1	2.36	0.40
6:Ar:108:ASP:OD1	6:Ar:108:ASP:C	2.65	0.40
6:Ar:122:LEU:HD22	7:Ef:801:MET:CE	2.51	0.40
3:Ba:4:ILE:HD12	6:Bp:132:GLN:HG3	2.04	0.40
6:Bd:130:LEU:HA	6:Bd:133:ILE:HD11	2.03	0.40
6:Bv:106:VAL:HG23	6:Bv:109:LYS:HD2	2.03	0.40
6:Ch:237:VAL:HG22	6:Ch:239:LEU:HD11	2.04	0.40
6:Cz:115:MET:HE1	7:Eo:801:MET:HG3	2.03	0.40
6:Df:108:ASP:HA	7:Ep:797:ARG:CZ	2.51	0.40
6:Df:169:ILE:HD13	6:Df:241:LEU:CD2	2.50	0.40
5:Dw:241:TYR:CE2	5:Dw:262:ILE:HD11	2.56	0.40
6:Dx:202:PHE:CG	6:Dx:241:LEU:HD12	2.56	0.40
6:Dx:207:ASP:OD2	6:Dx:210:SER:HB3	2.22	0.40
7:Es:789:LYS:HD2	7:Es:789:LYS:N	2.36	0.40
8:Ew:307:ALA:HB1	8:Ew:314:TYR:CE1	2.56	0.40
8:Ew:348:TYR:CD1	8:Ew:351:LEU:HD23	2.55	0.40
8:Ex:192:ASP:OD2	8:Ex:196:LYS:HG2	2.22	0.40
8:Ey:59:LYS:HG3	8:Ez:49:TYR:OH	2.22	0.40
8:Ey:349:TYR:CZ	8:Ey:353:LYS:HE3	2.56	0.40
8:Ez:277:LEU:O	8:Ez:280:ILE:HG22	2.21	0.40
8:Fa:253:LEU:C	8:Fa:254:MET:HE2	2.47	0.40
9:Fb:239:PHE:CD1	9:Fb:261:LEU:HG	2.55	0.40
9:Fb:709:ASN:HD21	9:Fc:597:LYS:N	2.20	0.40
9:Fb:816:ILE:O	9:Fb:820:ILE:HG22	2.20	0.40
9:Fe:196:LYS:HE2	9:Fe:196:LYS:HB2	1.79	0.40
9:Fe:373:ILE:HD12	9:Fe:373:ILE:N	2.36	0.40
9:Fe:470:ASP:O	9:Fe:473:LYS:HB2	2.20	0.40
9:Fe:599:ASP:C	9:Fe:599:ASP:OD1	2.64	0.40
9:Ff:27:ALA:HA	9:Ff:28:PRO:HD2	1.97	0.40
9:Ff:79:THR:HG21	9:Ff:788:HIS:CE1	2.57	0.40
9:Ff:734:LEU:HD11	9:Ff:738:TRP:NE1	2.36	0.40
7:Fg:29:ILE:HD12	7:Fg:29:ILE:HA	1.91	0.40
7:Fg:34:ILE:HG13	7:Fg:34:ILE:H	1.70	0.40
7:Fh:29:ILE:O	7:Fh:32:VAL:HG22	2.21	0.40
7:Fk:30:ILE:HA	7:Fk:33:VAL:HG22	2.03	0.40
5:Ae:229:THR:O	5:Ae:230:LEU:HD23	2.20	0.40
6:Af:181:PHE:CD2	6:Af:181:PHE:N	2.89	0.40

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
6:Af:236:PRO:HG2	6:Al:251:ARG:CZ	2.51	0.40
6:Ar:121:PRO:CB	7:Ef:801:MET:HE1	2.50	0.40
6:Ar:222:ASN:HB3	6:Ax:142:ALA:HB1	2.04	0.40
2:At:5:ILE:HD11	6:Bd:136:THR:O	2.21	0.40
5:Bc:241:TYR:HB3	5:Bc:256:THR:HG21	2.03	0.40
6:Bd:191:ILE:HD12	6:Bd:206:TRP:NE1	2.36	0.40
6:Bj:181:PHE:CD2	6:Bj:181:PHE:N	2.89	0.40
6:Bj:181:PHE:CE2	6:Bj:191:ILE:HD11	2.57	0.40
6:Bj:221:TYR:HE2	6:Bp:142:ALA:HB3	1.87	0.40
1:Bq:11:TYR:HD2	3:Bs:9:TYR:HB3	1.86	0.40
6:Bv:111:ALA:HB2	7:Ej:794:ILE:HG23	2.04	0.40
6:Bv:191:ILE:HB	6:Bv:206:TRP:CZ2	2.57	0.40
6:Cb:221:TYR:HA	6:Cb:243:PRO:HG2	2.03	0.40
6:Cz:108:ASP:HA	7:Eo:797:ARG:NH1	2.37	0.40
2:Dn:2:VAL:HA	3:Do:1:ARG:NE	2.31	0.40
6:Dx:108:ASP:HA	7:Es:797:ARG:HH12	1.85	0.40
6:Dx:167:PRO:HD2	6:Dx:238:MET:O	2.21	0.40
6:Ed:242:ILE:N	6:Ed:242:ILE:HD12	2.36	0.40
7:El:789:LYS:HD2	9:Fd:267:ASP:HB3	2.03	0.40
7:Ej:808:LEU:O	7:Ej:812:TRP:CD1	2.74	0.40
7:El:792:GLN:O	7:El:795:GLN:HG2	2.21	0.40
7:El:806:THR:O	7:El:810:GLN:HG3	2.21	0.40
7:Ev:792:GLN:H	7:Ev:792:GLN:CD	2.29	0.40
8:Ew:359:MET:HE3	8:Ew:367:ASN:O	2.22	0.40
8:Ew:368:ILE:H	8:Fa:105:LYS:NZ	2.19	0.40
8:Ew:384:PHE:N	8:Fa:82:TYR:CE2	2.90	0.40
8:Ex:408:PRO:HA	8:Ex:411:VAL:HG22	2.03	0.40
8:Ey:349:TYR:HD1	8:Ey:435:ARG:CZ	2.34	0.40
8:Ey:410:THR:O	8:Ey:413:LYS:HB2	2.21	0.40
8:Ey:412:GLN:HG3	8:Ez:411:VAL:HG21	2.03	0.40
8:Ez:43:LEU:HA	8:Ez:46:LEU:HG	2.03	0.40
8:Ez:43:LEU:HD12	8:Ez:44:THR:N	2.35	0.40
8:Ez:175:SER:O	8:Fa:240:ILE:HD11	2.21	0.40
8:Ez:353:LYS:HD2	8:Ez:353:LYS:HA	1.67	0.40
8:Ez:456:ASP:OD1	8:Ez:456:ASP:C	2.64	0.40
8:Fa:68:LEU:HD21	8:Fa:71:SER:HB3	2.03	0.40
9:Fb:44:VAL:N	9:Fb:55:GLN:HE22	2.17	0.40
9:Fb:167:LEU:O	9:Fb:170:ALA:HB3	2.21	0.40
9:Fb:222:MET:HE1	9:Fb:519:TRP:CD1	2.56	0.40
9:Fb:337:GLN:O	9:Fb:341:VAL:HG12	2.22	0.40
9:Fb:694:THR:O	9:Fb:697:ILE:HG12	2.22	0.40

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
9:Fd:77:MET:SD	9:Fd:81:MET:HE2	2.62	0.40
9:Fd:635:TYR:CE1	9:Fe:610:PHE:HD2	2.39	0.40
9:Fd:715:SER:HB2	9:Fd:724:ILE:HG13	2.03	0.40
9:Fe:619:PHE:CD1	9:Fe:619:PHE:N	2.88	0.40
9:Fe:669:ASP:HA	9:Fe:672:ILE:CG1	2.52	0.40
9:Fe:905:TRP:HA	9:Fe:908:VAL:HG23	2.03	0.40
9:Ff:537:VAL:HA	9:Ff:538:PRO:HA	1.90	0.40
9:Ff:594:ILE:HD12	9:Ff:594:ILE:H	1.86	0.40
9:Ff:684:ASN:HB3	9:Ff:687:VAL:CG2	2.50	0.40
7:Fg:29:ILE:O	7:Fg:33:VAL:HG23	2.22	0.40
5:Ae:219:ARG:HG2	6:Al:148:PRO:CD	2.49	0.40
6:Af:170:ARG:HG2	6:Af:249:ASP:CG	2.46	0.40
6:Al:207:ASP:OD2	6:Al:210:SER:HB3	2.21	0.40
6:Ar:126:GLN:CA	6:Ar:129:LYS:HZ2	2.27	0.40
6:Ar:223:TYR:CE1	6:Ax:175:PHE:HE2	2.39	0.40
1:As:1:CYS:SG	4:Av:2:PHE:HB3	2.62	0.40
5:Bc:264:PHE:CD2	6:Bd:246:LYS:HB2	2.57	0.40
6:Bd:149:PRO:HA	6:Bd:246:LYS:O	2.20	0.40
6:Bd:233:LEU:HD22	6:Bd:235:THR:O	2.21	0.40
6:Bj:118:ASN:O	6:Bj:121:PRO:HD3	2.20	0.40
6:Bp:115:MET:HE3	7:Et:802:LEU:HD13	2.04	0.40
5:Ca:215:VAL:HG22	5:Ca:249:SER:HB3	2.04	0.40
6:Cb:141:LYS:NZ	6:Ch:127:VAL:HG13	2.29	0.40
1:Cc:3:ASP:CG	3:Ce:14:TYR:HB3	2.47	0.40
6:Ch:222:ASN:OD1	6:Ch:222:ASN:N	2.54	0.40
6:Ct:208:LYS:HE2	7:Eq:823:GLY:HA2	2.03	0.40
6:Cz:166:PRO:HA	6:Cz:167:PRO:HD3	1.97	0.40
6:Cz:170:ARG:HG3	6:Cz:247:ALA:O	2.22	0.40
6:Df:183:ASP:HB3	6:Df:189:TRP:CE3	2.56	0.40
6:Df:190:PRO:HG2	6:Df:231:ARG:HG3	2.03	0.40
1:Dg:1:CYS:SG	4:Dj:2:PHE:HB3	2.61	0.40
6:Dl:189:TRP:CD2	6:Dl:230:LEU:HD13	2.57	0.40
6:Dl:209:THR:HG22	7:Et:824:THR:HG21	2.04	0.40
8:Ew:426:MET:O	8:Ew:430:ARG:HG3	2.20	0.40
8:Ew:434:GLU:HG2	8:Ex:350:ILE:HD12	2.02	0.40
8:Ex:87:GLY:HA3	8:Ex:118:THR:HG22	2.03	0.40
8:Ex:144:GLN:HB2	8:Ex:179:PHE:CE2	2.57	0.40
8:Ex:326:LEU:N	8:Ex:326:LEU:HD23	2.37	0.40
8:Ex:376:PHE:CZ	8:Ex:380:THR:HG21	2.55	0.40
8:Ey:430:ARG:NH2	8:Ez:425:GLN:HA	2.36	0.40
8:Ez:327:SER:OG	8:Fa:211:MET:HE1	2.20	0.40

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
8:Ez:354:ARG:HA	8:Ez:370:SER:HB2	2.02	0.40
8:Ez:411:VAL:O	8:Ez:415:ILE:HG12	2.21	0.40
8:Ez:445:LEU:HG	8:Fa:284:SER:HA	2.03	0.40
8:Fa:194:GLU:HG2	8:Fa:196:LYS:NZ	2.36	0.40
9:Fb:49:LEU:HD12	9:Fb:50:HIS:N	2.36	0.40
9:Fb:128:PHE:N	9:Fb:128:PHE:CD1	2.88	0.40
9:Fb:466:SER:HB2	9:Fc:839:HIS:ND1	2.36	0.40
9:Fb:618:LEU:HG	9:Fc:618:LEU:HD11	2.03	0.40
9:Fb:667:MET:HA	9:Fb:670:ILE:HG22	2.02	0.40
9:Fb:905:TRP:CD1	9:Ff:698:ASN:HA	2.54	0.40
9:Fb:905:TRP:NE1	9:Ff:702:THR:HG22	2.29	0.40
9:Fc:148:SER:HB2	9:Fc:152:ARG:HH12	1.87	0.40
9:Fc:610:PHE:CD1	9:Fc:629:GLY:HA3	2.56	0.40
9:Fd:28:PRO:HB2	9:Fd:32:ASP:OD2	2.22	0.40
9:Fd:81:MET:O	9:Fd:84:THR:HG22	2.22	0.40
9:Fd:332:ARG:NH2	9:Fd:489:THR:HA	2.36	0.40
9:Fd:597:LYS:HE2	9:Fd:597:LYS:N	2.35	0.40
9:Fe:64:PHE:HB2	9:Ff:817:ILE:HD11	2.04	0.40
9:Fe:383:LYS:HE2	9:Fe:383:LYS:HB3	1.91	0.40
9:Fe:727:LEU:HD23	9:Fe:728:ILE:N	2.36	0.40
9:Ff:224:THR:HA	9:Ff:227:ARG:NH1	2.36	0.40
9:Ff:275:LEU:O	9:Ff:278:ILE:HG12	2.20	0.40
9:Ff:378:PHE:CD1	9:Ff:414:PHE:HE2	2.40	0.40
9:Ff:614:ARG:HA	9:Ff:623:CYS:HA	2.03	0.40
9:Ff:810:LEU:O	9:Ff:811:ARG:C	2.65	0.40
5:Ae:219:ARG:HD3	6:Al:148:PRO:O	2.22	0.40
5:Ae:245:LYS:HZ1	5:Ae:255:LEU:HD12	1.87	0.40
5:Ak:216:ILE:HG12	5:Ak:221:TRP:CH2	2.56	0.40
5:Ak:246:LEU:HD21	5:Ak:248:ASP:HB2	2.02	0.40
5:Ak:253:ARG:HB2	5:Ak:253:ARG:HH11	1.87	0.40
6:Al:119:LEU:HD21	7:Ev:805:ALA:HB3	2.03	0.40
6:Al:159:ASN:OD1	6:Al:164:SER:HB3	2.20	0.40
1:As:13:LYS:HA	1:As:13:LYS:HD2	1.84	0.40
6:Bd:168:VAL:HA	6:Bd:240:THR:HG23	2.03	0.40
6:Bj:145:PRO:HB3	6:Bp:132:GLN:OE1	2.21	0.40
5:Bo:256:THR:HG22	5:Bo:260:GLN:O	2.22	0.40
5:Bo:261:VAL:HG12	5:Bo:263:LYS:HZ3	1.86	0.40
5:Cg:219:ARG:HH22	6:Cn:150:LYS:HD2	1.87	0.40
6:Cn:117:ARG:HD2	6:Cn:117:ARG:O	2.21	0.40
6:Ct:129:LYS:O	6:Ct:133:ILE:HG22	2.22	0.40
6:Df:195:ASP:O	6:Df:226:LEU:HD12	2.21	0.40

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
6:Ed:126:GLN:HA	6:Ed:129:LYS:CG	2.52	0.40
8:Ez:415:ILE:HA	8:Ez:418:LEU:HD21	2.02	0.40
9:Fb:672:ILE:HA	9:Fb:675:VAL:HG12	2.02	0.40
9:Fb:909:TYR:O	9:Fb:910:ALA:C	2.65	0.40
9:Fc:69:LEU:HD11	9:Fc:114:LEU:O	2.21	0.40
9:Fc:579:LEU:HD23	9:Fc:579:LEU:HA	1.67	0.40
9:Fd:105:LEU:HA	9:Fd:108:THR:HG22	2.04	0.40
9:Fd:936:LEU:HA	9:Fd:939:LYS:CE	2.51	0.40
9:Fe:602:LEU:HD22	9:Fe:603:TYR:CE2	2.56	0.40
9:Fe:618:LEU:C	9:Fe:619:PHE:HD1	2.30	0.40
9:Ff:131:VAL:HG12	9:Ff:809:PHE:HB2	2.04	0.40
9:Ff:399:TRP:CE3	9:Ff:416:GLY:HA3	2.57	0.40
9:Ff:905:TRP:O	9:Ff:909:TYR:CD1	2.75	0.40
7:Fg:17:THR:O	7:Fg:20:ILE:HG13	2.21	0.40

There are no symmetry-related clashes.

## 5.3 Torsion angles

### 5.3.1 Protein backbone

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

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

Mol	Chain	Analysed	Favoured	Allowed	Outliers	Percentiles	
1	Aa	14/16 (88%)	14 (100%)	0	0	100	100
1	Ag	14/16 (88%)	14 (100%)	0	0	100	100
1	Am	14/16 (88%)	14 (100%)	0	0	100	100
1	As	14/16 (88%)	14 (100%)	0	0	100	100
1	Ay	14/16 (88%)	14 (100%)	0	0	100	100
1	Be	14/16 (88%)	14 (100%)	0	0	100	100
1	Bk	14/16 (88%)	14 (100%)	0	0	100	100
1	Bq	14/16 (88%)	14 (100%)	0	0	100	100
1	Bw	14/16 (88%)	14 (100%)	0	0	100	100

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Mol	Chain	Analysed	Favoured	Allowed	Outliers	Percentiles	
1	Cc	14/16 (88%)	14 (100%)	0	0	100	100
1	Ci	14/16 (88%)	14 (100%)	0	0	100	100
1	Co	14/16 (88%)	14 (100%)	0	0	100	100
1	Cu	14/16 (88%)	14 (100%)	0	0	100	100
1	Da	14/16 (88%)	14 (100%)	0	0	100	100
1	Dg	14/16 (88%)	14 (100%)	0	0	100	100
1	Dm	14/16 (88%)	14 (100%)	0	0	100	100
1	Ds	14/16 (88%)	14 (100%)	0	0	100	100
1	Dy	14/16 (88%)	14 (100%)	0	0	100	100
2	Ab	7/9 (78%)	7 (100%)	0	0	100	100
2	Ah	7/9 (78%)	7 (100%)	0	0	100	100
2	An	7/9 (78%)	7 (100%)	0	0	100	100
2	At	7/9 (78%)	7 (100%)	0	0	100	100
2	Az	7/9 (78%)	7 (100%)	0	0	100	100
2	Bf	7/9 (78%)	7 (100%)	0	0	100	100
2	Bl	7/9 (78%)	7 (100%)	0	0	100	100
2	Br	7/9 (78%)	7 (100%)	0	0	100	100
2	Bx	7/9 (78%)	7 (100%)	0	0	100	100
2	Cd	7/9 (78%)	7 (100%)	0	0	100	100
2	Cj	7/9 (78%)	7 (100%)	0	0	100	100
2	Cp	7/9 (78%)	7 (100%)	0	0	100	100
2	Cv	7/9 (78%)	7 (100%)	0	0	100	100
2	Db	7/9 (78%)	6 (86%)	1 (14%)	0	100	100
2	Dh	7/9 (78%)	7 (100%)	0	0	100	100
2	Dn	7/9 (78%)	7 (100%)	0	0	100	100
2	Dt	7/9 (78%)	7 (100%)	0	0	100	100
2	Dz	7/9 (78%)	7 (100%)	0	0	100	100
3	Ac	14/16 (88%)	12 (86%)	2 (14%)	0	100	100
3	Ai	14/16 (88%)	12 (86%)	2 (14%)	0	100	100
3	Ao	14/16 (88%)	12 (86%)	2 (14%)	0	100	100
3	Au	14/16 (88%)	12 (86%)	2 (14%)	0	100	100

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Mol	Chain	Analysed	Favoured	Allowed	Outliers	Percentiles	
3	Ba	14/16 (88%)	12 (86%)	2 (14%)	0	100	100
3	Bg	14/16 (88%)	13 (93%)	1 (7%)	0	100	100
3	Bm	14/16 (88%)	12 (86%)	2 (14%)	0	100	100
3	Bs	14/16 (88%)	14 (100%)	0	0	100	100
3	By	14/16 (88%)	13 (93%)	1 (7%)	0	100	100
3	Ce	14/16 (88%)	12 (86%)	2 (14%)	0	100	100
3	Ck	14/16 (88%)	12 (86%)	2 (14%)	0	100	100
3	Cq	14/16 (88%)	12 (86%)	2 (14%)	0	100	100
3	Cw	14/16 (88%)	13 (93%)	1 (7%)	0	100	100
3	Dc	14/16 (88%)	13 (93%)	1 (7%)	0	100	100
3	Di	14/16 (88%)	12 (86%)	2 (14%)	0	100	100
3	Do	14/16 (88%)	13 (93%)	1 (7%)	0	100	100
3	Du	14/16 (88%)	13 (93%)	1 (7%)	0	100	100
3	Ea	14/16 (88%)	13 (93%)	1 (7%)	0	100	100
4	Ad	3/5 (60%)	3 (100%)	0	0	100	100
4	Aj	3/5 (60%)	3 (100%)	0	0	100	100
4	Ap	3/5 (60%)	3 (100%)	0	0	100	100
4	Av	3/5 (60%)	3 (100%)	0	0	100	100
4	Bb	3/5 (60%)	3 (100%)	0	0	100	100
4	Bh	3/5 (60%)	3 (100%)	0	0	100	100
4	Bn	3/5 (60%)	3 (100%)	0	0	100	100
4	Bt	3/5 (60%)	3 (100%)	0	0	100	100
4	Bz	3/5 (60%)	3 (100%)	0	0	100	100
4	Cf	3/5 (60%)	3 (100%)	0	0	100	100
4	Cl	3/5 (60%)	3 (100%)	0	0	100	100
4	Cr	3/5 (60%)	3 (100%)	0	0	100	100
4	Cx	3/5 (60%)	3 (100%)	0	0	100	100
4	Dd	3/5 (60%)	3 (100%)	0	0	100	100
4	Dj	3/5 (60%)	3 (100%)	0	0	100	100
4	Dp	3/5 (60%)	3 (100%)	0	0	100	100
4	Dv	3/5 (60%)	3 (100%)	0	0	100	100

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Mol	Chain	Analysed	Favoured	Allowed	Outliers	Percentiles	
4	Eb	3/5 (60%)	3 (100%)	0	0	100	100
5	Ae	61/269 (23%)	59 (97%)	2 (3%)	0	100	100
5	Ak	61/269 (23%)	60 (98%)	1 (2%)	0	100	100
5	Aq	61/269 (23%)	55 (90%)	6 (10%)	0	100	100
5	Aw	61/269 (23%)	56 (92%)	5 (8%)	0	100	100
5	Bc	61/269 (23%)	55 (90%)	6 (10%)	0	100	100
5	Bi	61/269 (23%)	58 (95%)	3 (5%)	0	100	100
5	Bo	61/269 (23%)	58 (95%)	3 (5%)	0	100	100
5	Bu	61/269 (23%)	56 (92%)	5 (8%)	0	100	100
5	Ca	61/269 (23%)	55 (90%)	6 (10%)	0	100	100
5	Cg	61/269 (23%)	58 (95%)	3 (5%)	0	100	100
5	Cm	61/269 (23%)	58 (95%)	3 (5%)	0	100	100
5	Cs	61/269 (23%)	53 (87%)	8 (13%)	0	100	100
5	Cy	61/269 (23%)	56 (92%)	5 (8%)	0	100	100
5	De	61/269 (23%)	55 (90%)	6 (10%)	0	100	100
5	Dk	61/269 (23%)	56 (92%)	5 (8%)	0	100	100
5	Dq	61/269 (23%)	57 (93%)	4 (7%)	0	100	100
5	Dw	61/269 (23%)	60 (98%)	1 (2%)	0	100	100
5	Ec	61/269 (23%)	60 (98%)	1 (2%)	0	100	100
6	Af	158/361 (44%)	153 (97%)	5 (3%)	0	100	100
6	Al	158/361 (44%)	149 (94%)	9 (6%)	0	100	100
6	Ar	158/361 (44%)	151 (96%)	7 (4%)	0	100	100
6	Ax	158/361 (44%)	149 (94%)	9 (6%)	0	100	100
6	Bd	158/361 (44%)	152 (96%)	6 (4%)	0	100	100
6	Bj	158/361 (44%)	148 (94%)	10 (6%)	0	100	100
6	Bp	158/361 (44%)	147 (93%)	11 (7%)	0	100	100
6	Bv	158/361 (44%)	150 (95%)	8 (5%)	0	100	100
6	Cb	158/361 (44%)	151 (96%)	7 (4%)	0	100	100
6	Ch	158/361 (44%)	151 (96%)	7 (4%)	0	100	100
6	Cn	158/361 (44%)	146 (92%)	12 (8%)	0	100	100
6	Ct	158/361 (44%)	147 (93%)	11 (7%)	0	100	100

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Mol	Chain	Analysed	Favoured	Allowed	Outliers	Percentiles	
6	Cz	158/361 (44%)	150 (95%)	8 (5%)	0	100	100
6	Df	158/361 (44%)	149 (94%)	9 (6%)	0	100	100
6	Dl	158/361 (44%)	152 (96%)	6 (4%)	0	100	100
6	Dr	158/361 (44%)	152 (96%)	6 (4%)	0	100	100
6	Dx	158/361 (44%)	155 (98%)	3 (2%)	0	100	100
6	Ed	158/361 (44%)	149 (94%)	9 (6%)	0	100	100
7	Ee	32/1048 (3%)	32 (100%)	0	0	100	100
7	Ef	57/1048 (5%)	57 (100%)	0	0	100	100
7	Eg	32/1048 (3%)	32 (100%)	0	0	100	100
7	Eh	46/1048 (4%)	44 (96%)	2 (4%)	0	100	100
7	Ei	47/1048 (4%)	45 (96%)	2 (4%)	0	100	100
7	Ej	56/1048 (5%)	56 (100%)	0	0	100	100
7	Ek	32/1048 (3%)	31 (97%)	1 (3%)	0	100	100
7	El	32/1048 (3%)	32 (100%)	0	0	100	100
7	Em	44/1048 (4%)	43 (98%)	1 (2%)	0	100	100
7	En	32/1048 (3%)	32 (100%)	0	0	100	100
7	Eo	32/1048 (3%)	32 (100%)	0	0	100	100
7	Ep	32/1048 (3%)	32 (100%)	0	0	100	100
7	Eq	55/1048 (5%)	52 (94%)	3 (6%)	0	100	100
7	Er	32/1048 (3%)	31 (97%)	1 (3%)	0	100	100
7	Es	54/1048 (5%)	53 (98%)	1 (2%)	0	100	100
7	Et	46/1048 (4%)	44 (96%)	2 (4%)	0	100	100
7	Eu	57/1048 (5%)	56 (98%)	1 (2%)	0	100	100
7	Ev	32/1048 (3%)	30 (94%)	2 (6%)	0	100	100
7	Fg	41/1048 (4%)	41 (100%)	0	0	100	100
7	Fh	41/1048 (4%)	41 (100%)	0	0	100	100
7	Fi	41/1048 (4%)	40 (98%)	1 (2%)	0	100	100
7	Fj	41/1048 (4%)	41 (100%)	0	0	100	100
7	Fk	41/1048 (4%)	40 (98%)	1 (2%)	0	100	100
8	Ew	413/466 (89%)	396 (96%)	17 (4%)	0	100	100
8	Ex	413/466 (89%)	398 (96%)	15 (4%)	0	100	100

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Mol	Chain	Analysed	Favoured	Allowed	Outliers	Percentiles	
8	Ey	413/466 (89%)	394 (95%)	19 (5%)	0	100	100
8	Ez	413/466 (89%)	388 (94%)	25 (6%)	0	100	100
8	Fa	413/466 (89%)	394 (95%)	19 (5%)	0	100	100
9	Fb	847/1048 (81%)	810 (96%)	37 (4%)	0	100	100
9	Fc	847/1048 (81%)	805 (95%)	42 (5%)	0	100	100
9	Fd	847/1048 (81%)	799 (94%)	48 (6%)	0	100	100
9	Fe	847/1048 (81%)	793 (94%)	54 (6%)	0	100	100
9	Ff	847/1048 (81%)	794 (94%)	53 (6%)	0	100	100
All	All	11881/43842 (27%)	11290 (95%)	591 (5%)	0	100	100

There are no Ramachandran outliers to report.

### 5.3.2 Protein sidechains ⓘ

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

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

Mol	Chain	Analysed	Rotameric	Outliers	Percentiles	
1	Aa	11/11 (100%)	11 (100%)	0	100	100
1	Ag	11/11 (100%)	11 (100%)	0	100	100
1	Am	11/11 (100%)	10 (91%)	1 (9%)	7	24
1	As	11/11 (100%)	11 (100%)	0	100	100
1	Ay	11/11 (100%)	11 (100%)	0	100	100
1	Be	11/11 (100%)	11 (100%)	0	100	100
1	Bk	11/11 (100%)	11 (100%)	0	100	100
1	Bq	11/11 (100%)	11 (100%)	0	100	100
1	Bw	11/11 (100%)	11 (100%)	0	100	100
1	Cc	11/11 (100%)	10 (91%)	1 (9%)	7	24
1	Ci	11/11 (100%)	9 (82%)	2 (18%)	1	8
1	Co	11/11 (100%)	11 (100%)	0	100	100
1	Cu	11/11 (100%)	11 (100%)	0	100	100

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Mol	Chain	Analysed	Rotameric	Outliers	Percentiles	
1	Da	11/11 (100%)	11 (100%)	0	100	100
1	Dg	11/11 (100%)	10 (91%)	1 (9%)	7	24
1	Dm	11/11 (100%)	10 (91%)	1 (9%)	7	24
1	Ds	11/11 (100%)	10 (91%)	1 (9%)	7	24
1	Dy	11/11 (100%)	11 (100%)	0	100	100
2	Ab	7/7 (100%)	7 (100%)	0	100	100
2	Ah	7/7 (100%)	7 (100%)	0	100	100
2	An	7/7 (100%)	7 (100%)	0	100	100
2	At	7/7 (100%)	7 (100%)	0	100	100
2	Az	7/7 (100%)	7 (100%)	0	100	100
2	Bf	7/7 (100%)	7 (100%)	0	100	100
2	Bl	7/7 (100%)	7 (100%)	0	100	100
2	Br	7/7 (100%)	7 (100%)	0	100	100
2	Bx	7/7 (100%)	7 (100%)	0	100	100
2	Cd	7/7 (100%)	7 (100%)	0	100	100
2	Cj	7/7 (100%)	6 (86%)	1 (14%)	2	13
2	Cp	7/7 (100%)	6 (86%)	1 (14%)	2	13
2	Cv	7/7 (100%)	7 (100%)	0	100	100
2	Db	7/7 (100%)	7 (100%)	0	100	100
2	Dh	7/7 (100%)	7 (100%)	0	100	100
2	Dn	7/7 (100%)	7 (100%)	0	100	100
2	Dt	7/7 (100%)	7 (100%)	0	100	100
2	Dz	7/7 (100%)	7 (100%)	0	100	100
3	Ac	13/13 (100%)	12 (92%)	1 (8%)	10	30
3	Ai	13/13 (100%)	12 (92%)	1 (8%)	10	30
3	Ao	13/13 (100%)	12 (92%)	1 (8%)	10	30
3	Au	13/13 (100%)	13 (100%)	0	100	100
3	Ba	13/13 (100%)	13 (100%)	0	100	100
3	Bg	13/13 (100%)	13 (100%)	0	100	100
3	Bm	13/13 (100%)	13 (100%)	0	100	100
3	Bs	13/13 (100%)	13 (100%)	0	100	100

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Mol	Chain	Analysed	Rotameric	Outliers	Percentiles	
3	By	13/13 (100%)	12 (92%)	1 (8%)	10	30
3	Ce	13/13 (100%)	12 (92%)	1 (8%)	10	30
3	Ck	13/13 (100%)	13 (100%)	0	100	100
3	Cq	13/13 (100%)	13 (100%)	0	100	100
3	Cw	13/13 (100%)	13 (100%)	0	100	100
3	Dc	13/13 (100%)	12 (92%)	1 (8%)	10	30
3	Di	13/13 (100%)	12 (92%)	1 (8%)	10	30
3	Do	13/13 (100%)	11 (85%)	2 (15%)	2	12
3	Du	13/13 (100%)	13 (100%)	0	100	100
3	Ea	13/13 (100%)	12 (92%)	1 (8%)	10	30
4	Ad	3/3 (100%)	3 (100%)	0	100	100
4	Aj	3/3 (100%)	3 (100%)	0	100	100
4	Ap	3/3 (100%)	3 (100%)	0	100	100
4	Av	3/3 (100%)	3 (100%)	0	100	100
4	Bb	3/3 (100%)	3 (100%)	0	100	100
4	Bh	3/3 (100%)	3 (100%)	0	100	100
4	Bn	3/3 (100%)	3 (100%)	0	100	100
4	Bt	3/3 (100%)	3 (100%)	0	100	100
4	Bz	3/3 (100%)	3 (100%)	0	100	100
4	Cf	3/3 (100%)	3 (100%)	0	100	100
4	Cl	3/3 (100%)	3 (100%)	0	100	100
4	Cr	3/3 (100%)	3 (100%)	0	100	100
4	Cx	3/3 (100%)	3 (100%)	0	100	100
4	Dd	3/3 (100%)	3 (100%)	0	100	100
4	Dj	3/3 (100%)	3 (100%)	0	100	100
4	Dp	3/3 (100%)	3 (100%)	0	100	100
4	Dv	3/3 (100%)	3 (100%)	0	100	100
4	Eb	3/3 (100%)	3 (100%)	0	100	100
5	Ae	53/237 (22%)	52 (98%)	1 (2%)	52	70
5	Ak	53/237 (22%)	52 (98%)	1 (2%)	52	70
5	Aq	53/237 (22%)	50 (94%)	3 (6%)	17	39

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Mol	Chain	Analysed	Rotameric	Outliers	Percentiles	
5	Aw	53/237 (22%)	51 (96%)	2 (4%)	28	49
5	Bc	53/237 (22%)	50 (94%)	3 (6%)	17	39
5	Bi	53/237 (22%)	52 (98%)	1 (2%)	52	70
5	Bo	53/237 (22%)	49 (92%)	4 (8%)	11	30
5	Bu	53/237 (22%)	51 (96%)	2 (4%)	28	49
5	Ca	53/237 (22%)	52 (98%)	1 (2%)	52	70
5	Cg	53/237 (22%)	49 (92%)	4 (8%)	11	30
5	Cm	53/237 (22%)	47 (89%)	6 (11%)	4	18
5	Cs	53/237 (22%)	53 (100%)	0	100	100
5	Cy	53/237 (22%)	51 (96%)	2 (4%)	28	49
5	De	53/237 (22%)	51 (96%)	2 (4%)	28	49
5	Dk	53/237 (22%)	53 (100%)	0	100	100
5	Dq	53/237 (22%)	53 (100%)	0	100	100
5	Dw	53/237 (22%)	50 (94%)	3 (6%)	17	39
5	Ec	53/237 (22%)	53 (100%)	0	100	100
6	Af	137/300 (46%)	127 (93%)	10 (7%)	11	31
6	Al	137/300 (46%)	131 (96%)	6 (4%)	24	46
6	Ar	137/300 (46%)	135 (98%)	2 (2%)	60	75
6	Ax	137/300 (46%)	128 (93%)	9 (7%)	14	35
6	Bd	137/300 (46%)	132 (96%)	5 (4%)	30	51
6	Bj	137/300 (46%)	132 (96%)	5 (4%)	30	51
6	Bp	137/300 (46%)	128 (93%)	9 (7%)	14	35
6	Bv	137/300 (46%)	130 (95%)	7 (5%)	20	42
6	Cb	137/300 (46%)	128 (93%)	9 (7%)	14	35
6	Ch	137/300 (46%)	130 (95%)	7 (5%)	20	42
6	Cn	137/300 (46%)	127 (93%)	10 (7%)	11	31
6	Ct	137/300 (46%)	135 (98%)	2 (2%)	60	75
6	Cz	137/300 (46%)	130 (95%)	7 (5%)	20	42
6	Df	137/300 (46%)	128 (93%)	9 (7%)	14	35
6	Dl	137/300 (46%)	125 (91%)	12 (9%)	8	25
6	Dr	137/300 (46%)	128 (93%)	9 (7%)	14	35

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Mol	Chain	Analysed	Rotameric	Outliers	Percentiles	
6	Dx	137/300 (46%)	131 (96%)	6 (4%)	24	46
6	Ed	137/300 (46%)	127 (93%)	10 (7%)	11	31
7	Ee	31/765 (4%)	29 (94%)	2 (6%)	14	35
7	Ef	51/765 (7%)	48 (94%)	3 (6%)	16	38
7	Eg	31/765 (4%)	31 (100%)	0	100	100
7	Eh	44/765 (6%)	44 (100%)	0	100	100
7	Ei	44/765 (6%)	39 (89%)	5 (11%)	4	17
7	Ej	50/765 (6%)	48 (96%)	2 (4%)	27	48
7	Ek	31/765 (4%)	31 (100%)	0	100	100
7	El	31/765 (4%)	31 (100%)	0	100	100
7	Em	42/765 (6%)	38 (90%)	4 (10%)	7	22
7	En	31/765 (4%)	29 (94%)	2 (6%)	14	35
7	Eo	31/765 (4%)	30 (97%)	1 (3%)	34	54
7	Ep	31/765 (4%)	31 (100%)	0	100	100
7	Eq	50/765 (6%)	48 (96%)	2 (4%)	27	48
7	Er	31/765 (4%)	31 (100%)	0	100	100
7	Es	50/765 (6%)	44 (88%)	6 (12%)	4	17
7	Et	44/765 (6%)	42 (96%)	2 (4%)	23	46
7	Eu	51/765 (7%)	50 (98%)	1 (2%)	50	68
7	Ev	31/765 (4%)	31 (100%)	0	100	100
7	Fg	36/765 (5%)	36 (100%)	0	100	100
7	Fh	36/765 (5%)	36 (100%)	0	100	100
7	Fi	36/765 (5%)	36 (100%)	0	100	100
7	Fj	36/765 (5%)	34 (94%)	2 (6%)	17	39
7	Fk	36/765 (5%)	34 (94%)	2 (6%)	17	39
8	Ew	360/401 (90%)	350 (97%)	10 (3%)	38	58
8	Ex	360/401 (90%)	349 (97%)	11 (3%)	35	55
8	Ey	360/401 (90%)	352 (98%)	8 (2%)	47	65
8	Ez	360/401 (90%)	348 (97%)	12 (3%)	33	54
8	Fa	360/401 (90%)	353 (98%)	7 (2%)	52	70
9	Fb	715/858 (83%)	691 (97%)	24 (3%)	32	53

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Mol	Chain	Analysed	Rotameric	Outliers	Percentiles	
9	Fc	715/858 (83%)	682 (95%)	33 (5%)	23	45
9	Fd	715/858 (83%)	693 (97%)	22 (3%)	35	55
9	Fe	715/858 (83%)	676 (94%)	39 (6%)	18	40
9	Ff	715/858 (83%)	688 (96%)	27 (4%)	28	49
All	All	10292/34168 (30%)	9877 (96%)	415 (4%)	29	48

All (415) residues with a non-rotameric sidechain are listed below:

Mol	Chain	Res	Type
3	Ac	8	VAL
5	Ae	211	TYR
6	Af	139	TYR
6	Af	144	THR
6	Af	147	THR
6	Af	168	VAL
6	Af	179	LEU
6	Af	202	PHE
6	Af	223	TYR
6	Af	226	LEU
6	Af	237	VAL
6	Af	256	VAL
3	Ai	8	VAL
5	Ak	245	LYS
6	Al	112	PHE
6	Al	127	VAL
6	Al	129	LYS
6	Al	134	TYR
6	Al	241	LEU
6	Al	259	TYR
1	Am	9	LEU
3	Ao	1	ARG
5	Aq	229	THR
5	Aq	238	ILE
5	Aq	255	LEU
6	Ar	179	LEU
6	Ar	259	TYR
5	Aw	246	LEU
5	Aw	247	ILE
6	Ax	144	THR
6	Ax	158	VAL
6	Ax	170	ARG

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Mol	Chain	Res	Type
6	Ax	171	LEU
6	Ax	202	PHE
6	Ax	213	LEU
6	Ax	220	LEU
6	Ax	248	VAL
6	Ax	259	TYR
5	Bc	211	TYR
5	Bc	238	ILE
5	Bc	264	PHE
6	Bd	122	LEU
6	Bd	144	THR
6	Bd	147	THR
6	Bd	202	PHE
6	Bd	241	LEU
5	Bi	238	ILE
6	Bj	119	LEU
6	Bj	125	GLU
6	Bj	128	VAL
6	Bj	179	LEU
6	Bj	250	TYR
5	Bo	216	ILE
5	Bo	236	SER
5	Bo	238	ILE
5	Bo	243	MET
6	Bp	113	LYS
6	Bp	127	VAL
6	Bp	128	VAL
6	Bp	134	TYR
6	Bp	202	PHE
6	Bp	215	ILE
6	Bp	242	ILE
6	Bp	248	VAL
6	Bp	254	LEU
5	Bu	222	LEU
5	Bu	244	VAL
6	Bv	127	VAL
6	Bv	128	VAL
6	Bv	168	VAL
6	Bv	221	TYR
6	Bv	228	VAL
6	Bv	241	LEU
6	Bv	248	VAL

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Mol	Chain	Res	Type
3	By	8	VAL
5	Ca	238	ILE
6	Cb	127	VAL
6	Cb	134	TYR
6	Cb	144	THR
6	Cb	168	VAL
6	Cb	176	VAL
6	Cb	177	SER
6	Cb	228	VAL
6	Cb	241	LEU
6	Cb	248	VAL
1	Cc	2	THR
3	Ce	7	THR
5	Cg	208	ILE
5	Cg	236	SER
5	Cg	238	ILE
5	Cg	247	ILE
6	Ch	172	SER
6	Ch	176	VAL
6	Ch	226	LEU
6	Ch	239	LEU
6	Ch	241	LEU
6	Ch	242	ILE
6	Ch	259	TYR
1	Ci	2	THR
1	Ci	3	ASP
2	Cj	4	MET
5	Cm	207	ARG
5	Cm	211	TYR
5	Cm	232	VAL
5	Cm	236	SER
5	Cm	246	LEU
5	Cm	250	LEU
6	Cn	107	ILE
6	Cn	128	VAL
6	Cn	129	LYS
6	Cn	141	LYS
6	Cn	144	THR
6	Cn	171	LEU
6	Cn	176	VAL
6	Cn	222	ASN
6	Cn	223	TYR

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Mol	Chain	Res	Type
6	Cn	259	TYR
2	Cp	4	MET
6	Ct	128	VAL
6	Ct	202	PHE
5	Cy	244	VAL
5	Cy	255	LEU
6	Cz	105	GLU
6	Cz	168	VAL
6	Cz	173	GLN
6	Cz	202	PHE
6	Cz	213	LEU
6	Cz	242	ILE
6	Cz	248	VAL
3	Dc	8	VAL
5	De	233	ARG
5	De	244	VAL
6	Df	120	TYR
6	Df	122	LEU
6	Df	160	LEU
6	Df	180	VAL
6	Df	202	PHE
6	Df	241	LEU
6	Df	242	ILE
6	Df	248	VAL
6	Df	256	VAL
1	Dg	9	LEU
3	Di	16	ASP
6	Dl	127	VAL
6	Dl	128	VAL
6	Dl	134	TYR
6	Dl	144	THR
6	Dl	169	ILE
6	Dl	171	LEU
6	Dl	202	PHE
6	Dl	204	ILE
6	Dl	228	VAL
6	Dl	235	THR
6	Dl	241	LEU
6	Dl	248	VAL
1	Dm	10	GLU
3	Do	7	THR
3	Do	8	VAL

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Mol	Chain	Res	Type
6	Dr	127	VAL
6	Dr	147	THR
6	Dr	158	VAL
6	Dr	168	VAL
6	Dr	179	LEU
6	Dr	191	ILE
6	Dr	215	ILE
6	Dr	230	LEU
6	Dr	234	ASN
1	Ds	9	LEU
5	Dw	244	VAL
5	Dw	253	ARG
5	Dw	262	ILE
6	Dx	158	VAL
6	Dx	165	THR
6	Dx	170	ARG
6	Dx	209	THR
6	Dx	228	VAL
6	Dx	248	VAL
3	Ea	8	VAL
6	Ed	122	LEU
6	Ed	177	SER
6	Ed	180	VAL
6	Ed	182	LEU
6	Ed	215	ILE
6	Ed	228	VAL
6	Ed	241	LEU
6	Ed	248	VAL
6	Ed	256	VAL
6	Ed	257	GLN
7	Ee	797	ARG
7	Ee	817	THR
7	Ef	780	LYS
7	Ef	794	ILE
7	Ef	797	ARG
7	Ei	782	ASN
7	Ei	790	TYR
7	Ei	799	SER
7	Ei	812	TRP
7	Ei	817	THR
7	Ej	799	SER
7	Ej	817	THR

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Mol	Chain	Res	Type
7	Em	788	GLN
7	Em	799	SER
7	Em	811	ASP
7	Em	819	VAL
7	En	795	GLN
7	En	817	THR
7	Eo	815	VAL
7	Eq	769	SER
7	Eq	778	LEU
7	Es	785	LEU
7	Es	799	SER
7	Es	807	GLN
7	Es	812	TRP
7	Es	817	THR
7	Es	821	THR
7	Et	788	GLN
7	Et	789	LYS
7	Eu	818	GLN
8	Ew	89	ILE
8	Ew	106	VAL
8	Ew	119	THR
8	Ew	170	LEU
8	Ew	210	VAL
8	Ew	294	LYS
8	Ew	326	LEU
8	Ew	334	ARG
8	Ew	423	ASN
8	Ew	428	LEU
8	Ex	39	LEU
8	Ex	101	PHE
8	Ex	196	LYS
8	Ex	197	ASN
8	Ex	227	THR
8	Ex	375	GLU
8	Ex	381	ARG
8	Ex	425	GLN
8	Ex	434	GLU
8	Ex	437	LEU
8	Ex	440	ASN
8	Ey	171	ASN
8	Ey	178	VAL
8	Ey	210	VAL

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Mol	Chain	Res	Type
8	Ey	300	GLU
8	Ey	373	LEU
8	Ey	375	GLU
8	Ey	383	LEU
8	Ey	445	LEU
8	Ez	76	LEU
8	Ez	101	PHE
8	Ez	102	VAL
8	Ez	142	ILE
8	Ez	172	ASP
8	Ez	277	LEU
8	Ez	301	LEU
8	Ez	302	TRP
8	Ez	326	LEU
8	Ez	332	ASN
8	Ez	371	GLN
8	Ez	387	THR
8	Fa	91	VAL
8	Fa	95	SER
8	Fa	268	THR
8	Fa	287	VAL
8	Fa	300	GLU
8	Fa	371	GLN
8	Fa	421	GLU
9	Fb	131	VAL
9	Fb	141	LYS
9	Fb	244	ASN
9	Fb	339	MET
9	Fb	413	LEU
9	Fb	436	LEU
9	Fb	473	LYS
9	Fb	495	SER
9	Fb	509	CYS
9	Fb	519	TRP
9	Fb	545	VAL
9	Fb	550	LEU
9	Fb	564	SER
9	Fb	593	LEU
9	Fb	631	LEU
9	Fb	652	ILE
9	Fb	660	LEU
9	Fb	703	LEU

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Mol	Chain	Res	Type
9	Fb	716	LEU
9	Fb	728	ILE
9	Fb	774	GLU
9	Fb	905	TRP
9	Fb	908	VAL
9	Fb	936	LEU
9	Fc	45	VAL
9	Fc	113	LEU
9	Fc	124	MET
9	Fc	126	VAL
9	Fc	210	LYS
9	Fc	273	TYR
9	Fc	282	VAL
9	Fc	302	VAL
9	Fc	330	LEU
9	Fc	360	THR
9	Fc	384	SER
9	Fc	434	LEU
9	Fc	491	LEU
9	Fc	508	THR
9	Fc	509	CYS
9	Fc	559	VAL
9	Fc	590	PHE
9	Fc	643	PHE
9	Fc	652	ILE
9	Fc	705	LEU
9	Fc	716	LEU
9	Fc	717	ILE
9	Fc	727	LEU
9	Fc	728	ILE
9	Fc	729	MET
9	Fc	738	TRP
9	Fc	755	VAL
9	Fc	759	MET
9	Fc	760	ILE
9	Fc	834	ASN
9	Fc	935	HIS
9	Fc	936	LEU
9	Fc	943	TRP
9	Fd	41	LEU
9	Fd	50	HIS
9	Fd	74	ILE

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Mol	Chain	Res	Type
9	Fd	150	LEU
9	Fd	227	ARG
9	Fd	260	GLU
9	Fd	270	SER
9	Fd	281	THR
9	Fd	341	VAL
9	Fd	369	ASP
9	Fd	509	CYS
9	Fd	514	SER
9	Fd	559	VAL
9	Fd	586	LYS
9	Fd	593	LEU
9	Fd	676	GLN
9	Fd	703	LEU
9	Fd	708	LEU
9	Fd	725	PHE
9	Fd	728	ILE
9	Fd	738	TRP
9	Fd	781	ILE
9	Fe	35	VAL
9	Fe	36	VAL
9	Fe	38	LEU
9	Fe	56	ILE
9	Fe	127	PHE
9	Fe	150	LEU
9	Fe	151	ASN
9	Fe	196	LYS
9	Fe	261	LEU
9	Fe	320	ILE
9	Fe	325	LEU
9	Fe	388	VAL
9	Fe	390	THR
9	Fe	436	LEU
9	Fe	491	LEU
9	Fe	500	GLN
9	Fe	513	TYR
9	Fe	515	LEU
9	Fe	519	TRP
9	Fe	533	LEU
9	Fe	560	SER
9	Fe	563	LEU
9	Fe	572	ASN

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Mol	Chain	Res	Type
9	Fe	639	TYR
9	Fe	660	LEU
9	Fe	662	ILE
9	Fe	703	LEU
9	Fe	709	ASN
9	Fe	710	MET
9	Fe	717	ILE
9	Fe	728	ILE
9	Fe	759	MET
9	Fe	763	PHE
9	Fe	811	ARG
9	Fe	839	HIS
9	Fe	905	TRP
9	Fe	915	ILE
9	Fe	935	HIS
9	Fe	943	TRP
9	Ff	49	LEU
9	Ff	136	VAL
9	Ff	182	LEU
9	Ff	261	LEU
9	Ff	290	LEU
9	Ff	320	ILE
9	Ff	330	LEU
9	Ff	378	PHE
9	Ff	387	GLU
9	Ff	509	CYS
9	Ff	545	VAL
9	Ff	577	VAL
9	Ff	579	LEU
9	Ff	598	VAL
9	Ff	602	LEU
9	Ff	603	TYR
9	Ff	617	ILE
9	Ff	644	PHE
9	Ff	689	LEU
9	Ff	724	ILE
9	Ff	729	MET
9	Ff	736	MET
9	Ff	774	GLU
9	Ff	787	THR
9	Ff	820	ILE
9	Ff	925	ILE

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Mol	Chain	Res	Type
9	Ff	943	TRP
7	Fj	11	LEU
7	Fj	17	THR
7	Fk	1	MET
7	Fk	27	LEU

Sometimes sidechains can be flipped to improve hydrogen bonding and reduce clashes. All (194) such sidechains are listed below:

Mol	Chain	Res	Type
6	Af	123	ASN
6	Af	132	GLN
6	Af	216	GLN
1	Ag	12	HIS
5	Ak	260	GLN
6	Al	132	GLN
6	Al	159	ASN
6	Al	203	ASN
1	Am	12	HIS
6	Ar	126	GLN
6	Ar	173	GLN
6	Ar	203	ASN
6	Ar	216	GLN
6	Ar	257	GLN
6	Ar	262	ASN
5	Aw	251	GLN
6	Ax	123	ASN
6	Ax	205	GLN
6	Ax	262	ASN
5	Bc	213	GLN
6	Bd	156	GLN
6	Bj	205	GLN
6	Bj	222	ASN
6	Bj	225	ASN
6	Bj	257	GLN
6	Bj	262	ASN
5	Bo	213	GLN
5	Bo	251	GLN
6	Bp	126	GLN
6	Bp	132	GLN
6	Bp	156	GLN
6	Bp	173	GLN
6	Bp	222	ASN

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Mol	Chain	Res	Type
6	Bp	245	GLN
6	Bv	123	ASN
6	Bv	126	GLN
6	Bv	132	GLN
6	Bv	173	GLN
6	Bv	257	GLN
6	Cb	132	GLN
6	Cb	203	ASN
6	Cb	262	ASN
6	Ch	132	GLN
6	Ch	262	ASN
5	Cm	213	GLN
5	Cm	266	GLN
6	Cn	123	ASN
6	Cn	203	ASN
6	Cn	205	GLN
5	Cs	251	GLN
5	Cs	260	GLN
5	Cy	251	GLN
6	Cz	225	ASN
6	Cz	234	ASN
6	Df	203	ASN
6	Df	211	ASN
6	Df	262	ASN
5	Dk	213	GLN
6	Dl	132	GLN
6	Dl	222	ASN
6	Dl	225	ASN
6	Dr	262	ASN
5	Dw	260	GLN
6	Dx	132	GLN
6	Dx	203	ASN
6	Dx	262	ASN
5	Ec	251	GLN
6	Ed	123	ASN
6	Ed	156	GLN
6	Ed	203	ASN
6	Ed	262	ASN
7	Eg	795	GLN
7	Eg	810	GLN
7	Eg	814	GLN
7	Eh	784	GLN

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Mol	Chain	Res	Type
7	Eh	792	GLN
7	Ei	779	GLN
7	Ei	792	GLN
7	Ek	814	GLN
7	El	814	GLN
7	Em	810	GLN
7	Eo	818	GLN
7	Ep	810	GLN
7	Eq	791	GLN
7	Eq	795	GLN
7	Er	810	GLN
7	Er	814	GLN
7	Es	775	GLN
7	Es	807	GLN
7	Et	781	GLN
7	Et	782	ASN
7	Et	792	GLN
7	Et	796	GLN
7	Et	814	GLN
7	Eu	775	GLN
7	Eu	818	GLN
7	Ev	791	GLN
7	Ev	810	GLN
8	Ew	78	GLN
8	Ew	79	ASN
8	Ew	208	ASN
8	Ew	212	GLN
8	Ew	237	GLN
8	Ew	242	GLN
8	Ew	246	ASN
8	Ew	286	GLN
8	Ew	412	GLN
8	Ew	431	GLN
8	Ex	62	ASN
8	Ex	79	ASN
8	Ex	123	GLN
8	Ex	171	ASN
8	Ex	263	GLN
8	Ex	374	ASN
8	Ex	425	GLN
8	Ex	447	ASN
8	Ey	72	ASN

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Mol	Chain	Res	Type
8	Ey	75	GLN
8	Ey	237	GLN
8	Ey	246	ASN
8	Ey	274	GLN
8	Ey	286	GLN
8	Ey	339	GLN
8	Ey	412	GLN
8	Ey	423	ASN
8	Ey	425	GLN
8	Ey	431	GLN
8	Ez	45	ASN
8	Ez	72	ASN
8	Ez	79	ASN
8	Ez	139	ASN
8	Ez	144	GLN
8	Ez	180	ASN
8	Ez	193	ASN
8	Ez	212	GLN
8	Ez	213	ASN
8	Ez	236	ASN
8	Ez	303	ASN
8	Ez	363	GLN
8	Ez	374	ASN
8	Ez	425	GLN
8	Fa	79	ASN
8	Fa	83	ASN
8	Fa	212	GLN
8	Fa	213	ASN
8	Fa	222	GLN
8	Fa	260	GLN
8	Fa	286	GLN
8	Fa	303	ASN
8	Fa	339	GLN
8	Fa	433	GLN
8	Fa	447	ASN
9	Fb	295	GLN
9	Fb	415	ASN
9	Fb	542	GLN
9	Fb	709	ASN
9	Fb	807	ASN
9	Fc	92	GLN
9	Fc	197	GLN

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Mol	Chain	Res	Type
9	Fc	286	ASN
9	Fc	326	GLN
9	Fc	338	GLN
9	Fc	484	GLN
9	Fc	521	GLN
9	Fc	529	GLN
9	Fd	134	GLN
9	Fd	197	GLN
9	Fd	337	GLN
9	Fd	397	GLN
9	Fd	457	ASN
9	Fd	709	ASN
9	Fd	927	GLN
9	Fe	65	ASN
9	Fe	92	GLN
9	Fe	96	GLN
9	Fe	151	ASN
9	Fe	265	ASN
9	Fe	291	ASN
9	Fe	368	ASN
9	Fe	435	ASN
9	Fe	510	GLN
9	Fe	531	GLN
9	Fe	698	ASN
9	Ff	59	ASN
9	Ff	96	GLN
9	Ff	188	GLN
9	Ff	197	GLN
9	Ff	199	GLN
9	Ff	294	ASN
9	Ff	316	ASN
9	Ff	337	GLN
9	Ff	338	GLN
9	Ff	927	GLN
7	Fg	7	ASN

### 5.3.3 RNA ⓘ

There are no RNA molecules in this entry.

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

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

## 5.5 Carbohydrates [i](#)

There are no oligosaccharides in this entry.

## 5.6 Ligand geometry [i](#)

There are no ligands in this entry.

## 5.7 Other polymers [i](#)

There are no such residues in this entry.

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

There are no chain breaks in this entry.

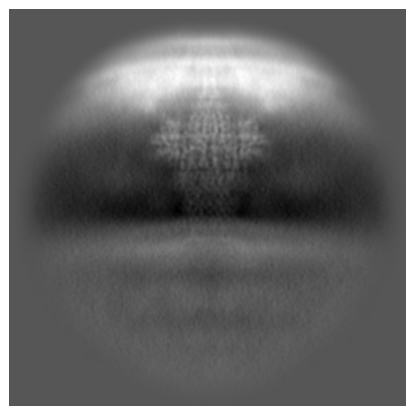
## 6 Map visualisation [i](#)

This section contains visualisations of the EMDB entry EMD-49398. These allow visual inspection of the internal detail of the map and identification of artifacts.

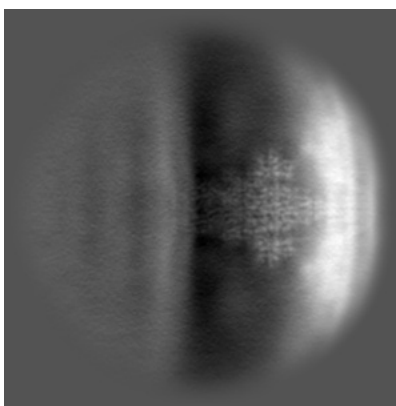
Images derived from a raw map, generated by summing the deposited half-maps, are presented below the corresponding image components of the primary map to allow further visual inspection and comparison with those of the primary map.

### 6.1 Orthogonal projections [i](#)

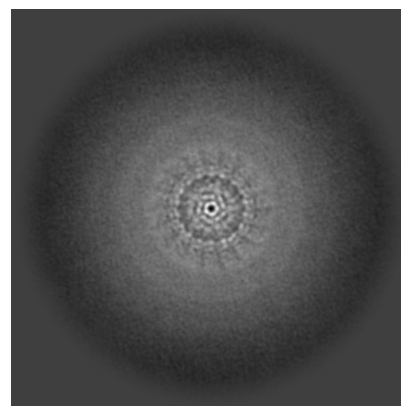
#### 6.1.1 Primary map



X

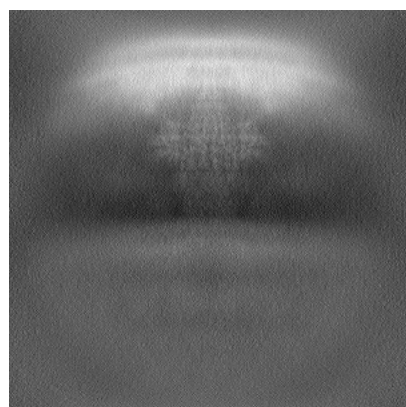


Y

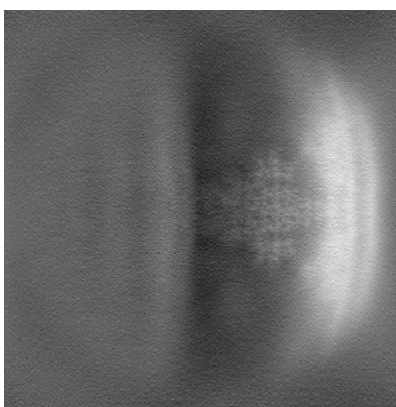


Z

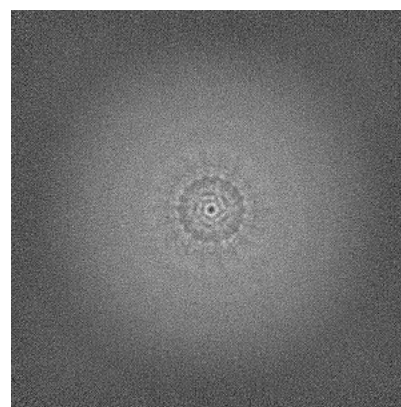
#### 6.1.2 Raw map



X



Y

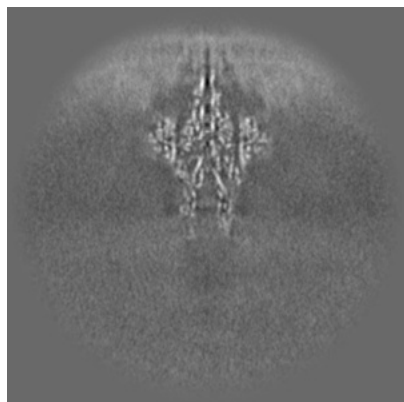


Z

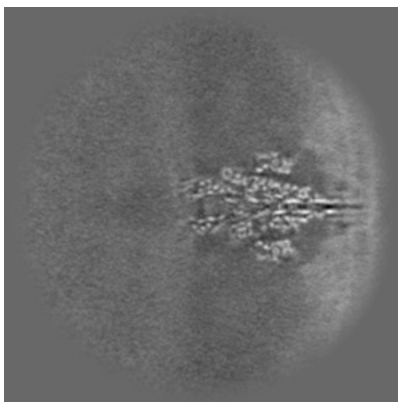
The images above show the map projected in three orthogonal directions.

## 6.2 Central slices [i](#)

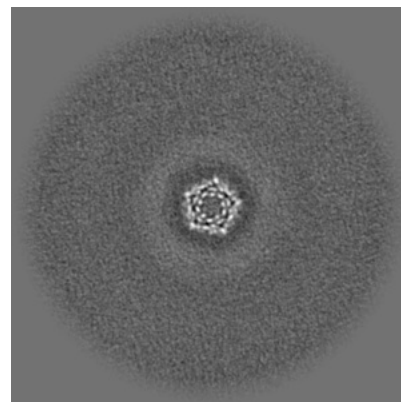
### 6.2.1 Primary map



X Index: 256

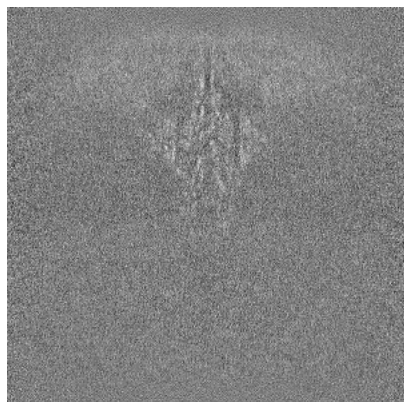


Y Index: 256

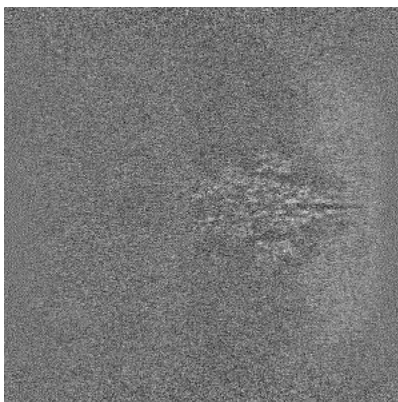


Z Index: 256

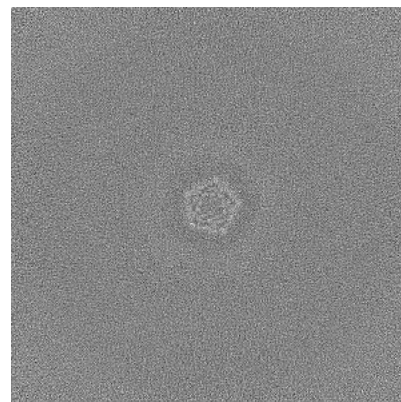
### 6.2.2 Raw map



X Index: 256



Y Index: 256



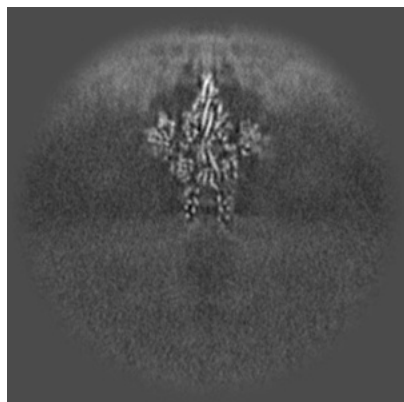
Z Index: 256

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

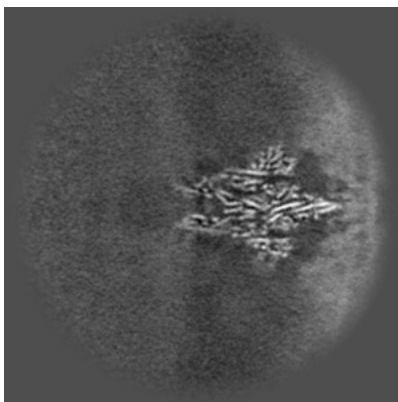


## 6.3 Largest variance slices [i](#)

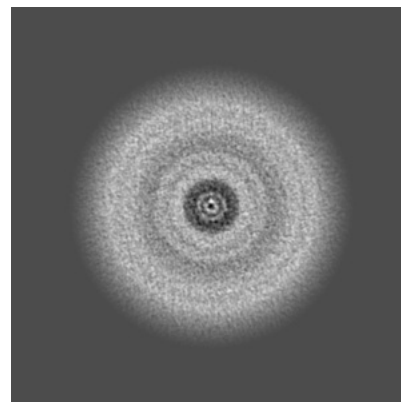
### 6.3.1 Primary map



X Index: 250

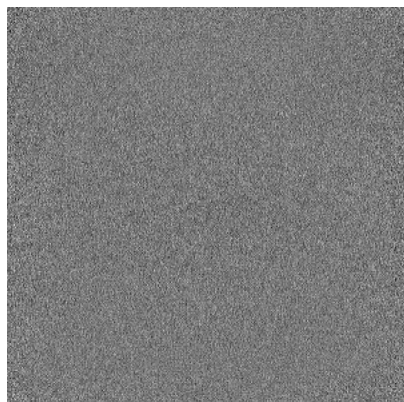


Y Index: 263

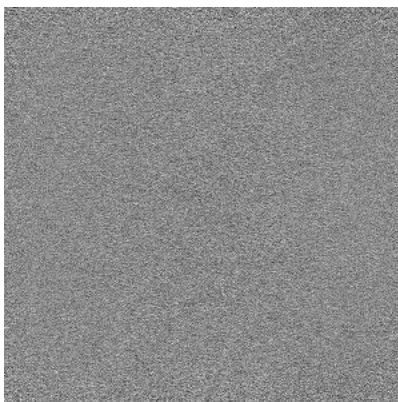


Z Index: 430

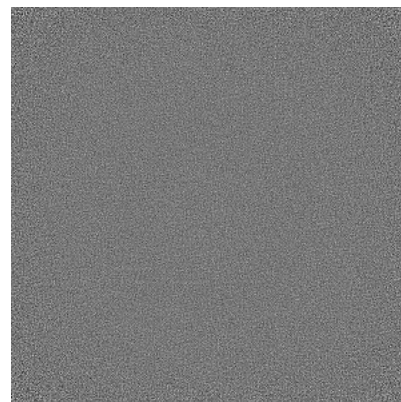
### 6.3.2 Raw map



X Index: 0



Y Index: 0

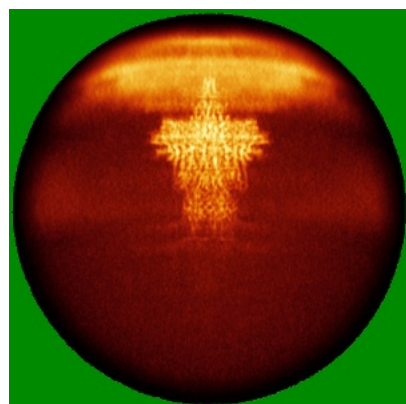


Z Index: 511

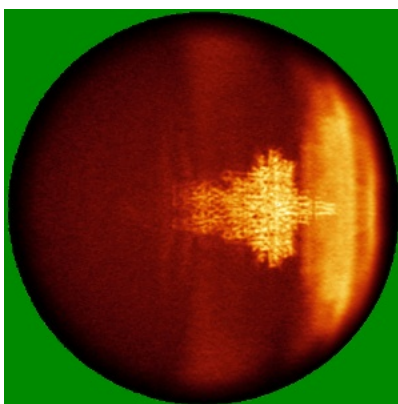
The images above show the largest variance slices of the map in three orthogonal directions.

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

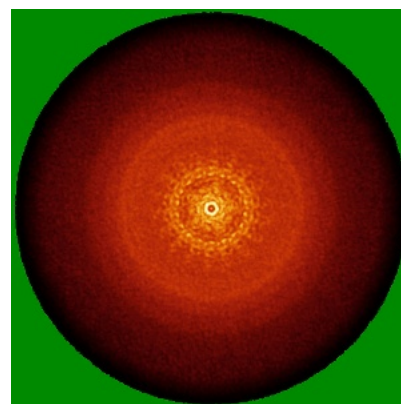
### 6.4.1 Primary map



X

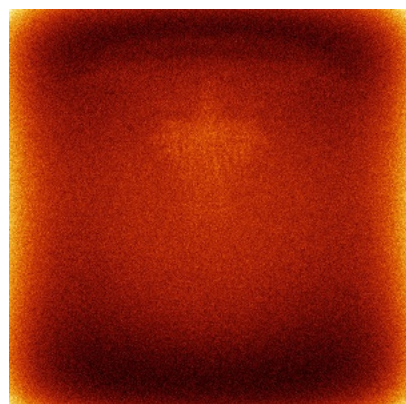


Y

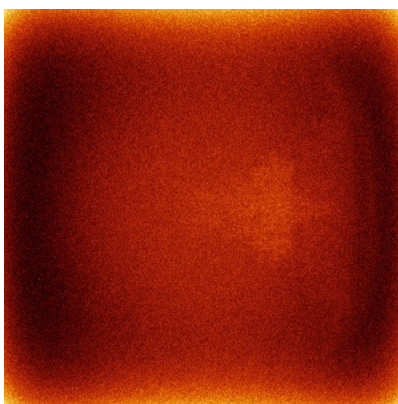


Z

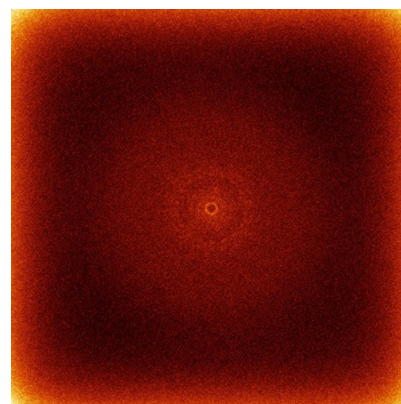
### 6.4.2 Raw map



X



Y



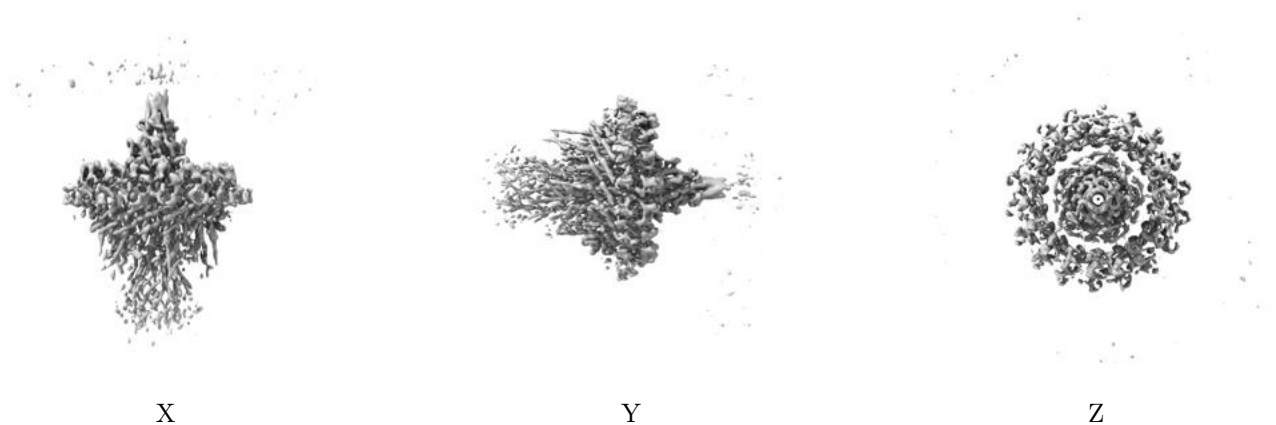
Z

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



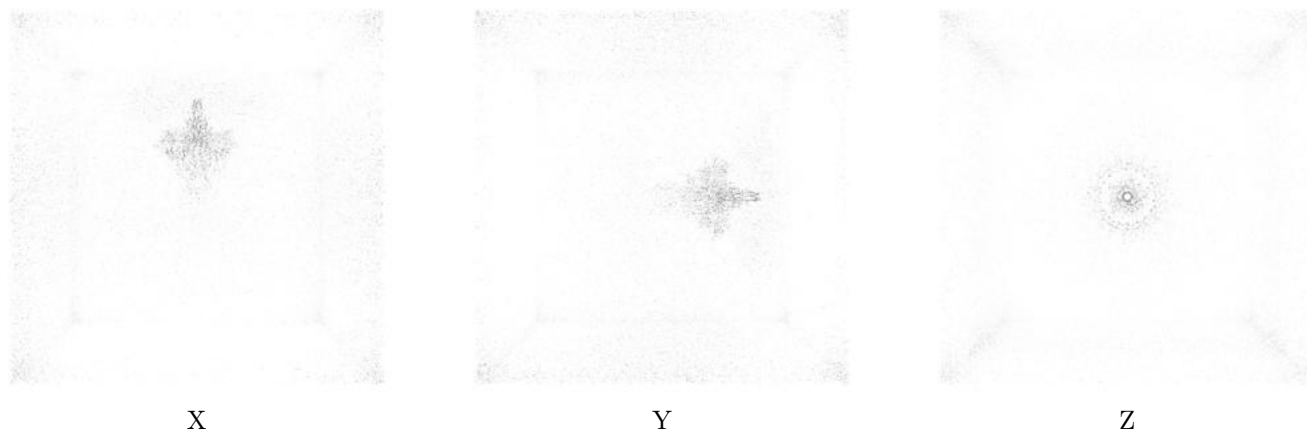
## 6.5 Orthogonal surface views [i](#)

### 6.5.1 Primary map



The images above show the 3D surface view of the map at the recommended contour level 0.16. These images, in conjunction with the slice images, may facilitate assessment of whether an appropriate contour level has been provided.

### 6.5.2 Raw map



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

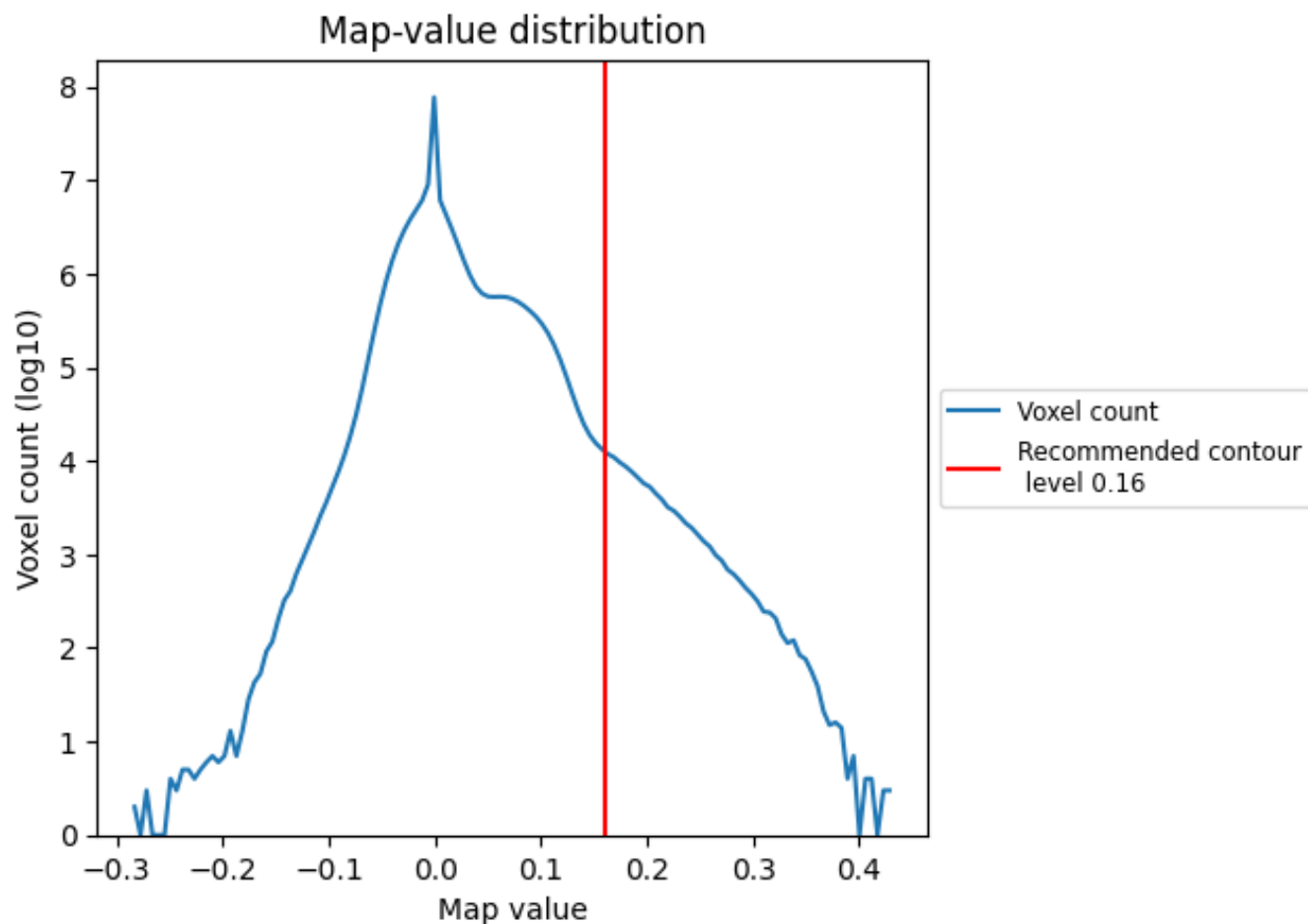
## 6.6 Mask visualisation [i](#)

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

## 7 Map analysis [i](#)

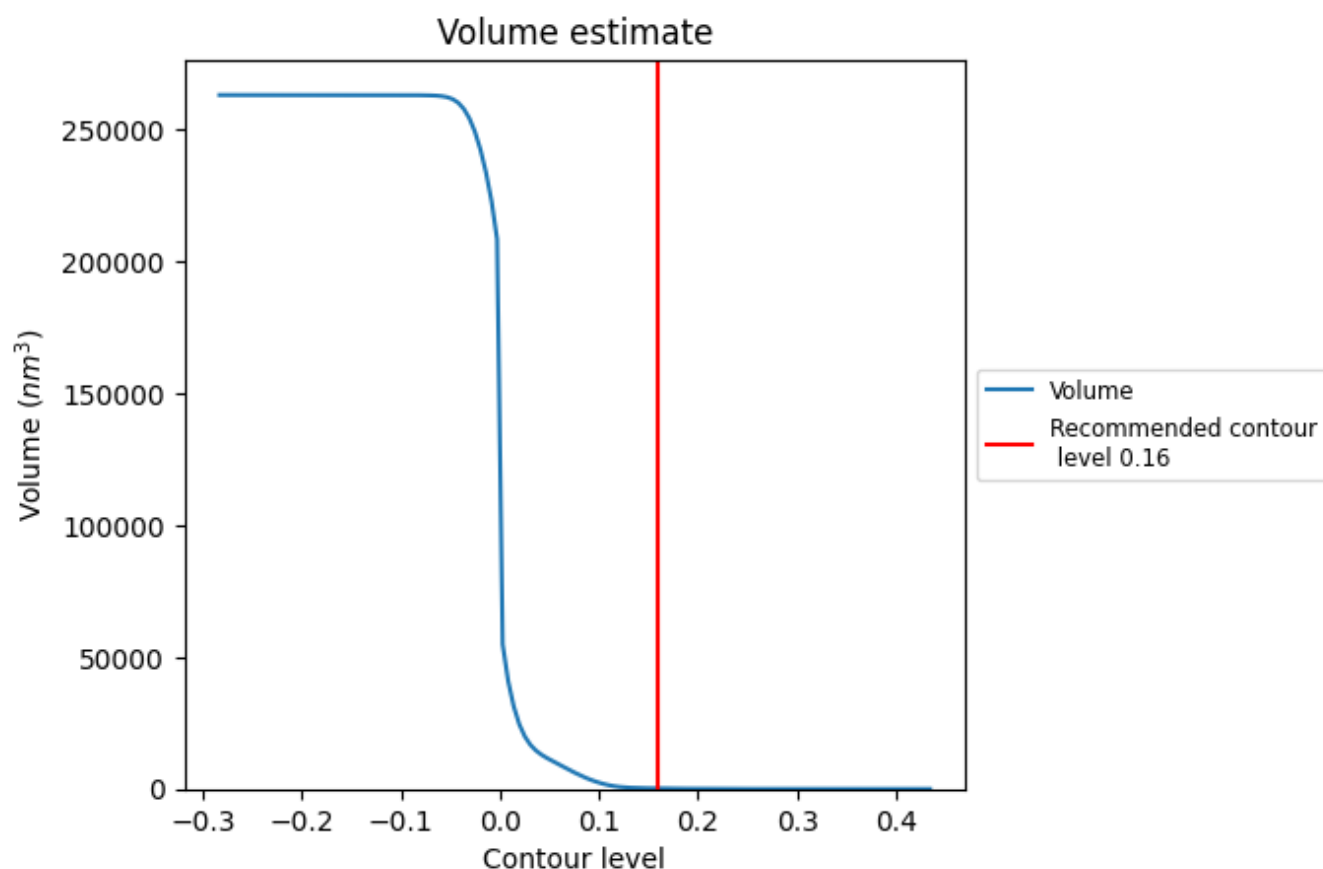
This section contains the results of statistical analysis of the map.

### 7.1 Map-value distribution [i](#)



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

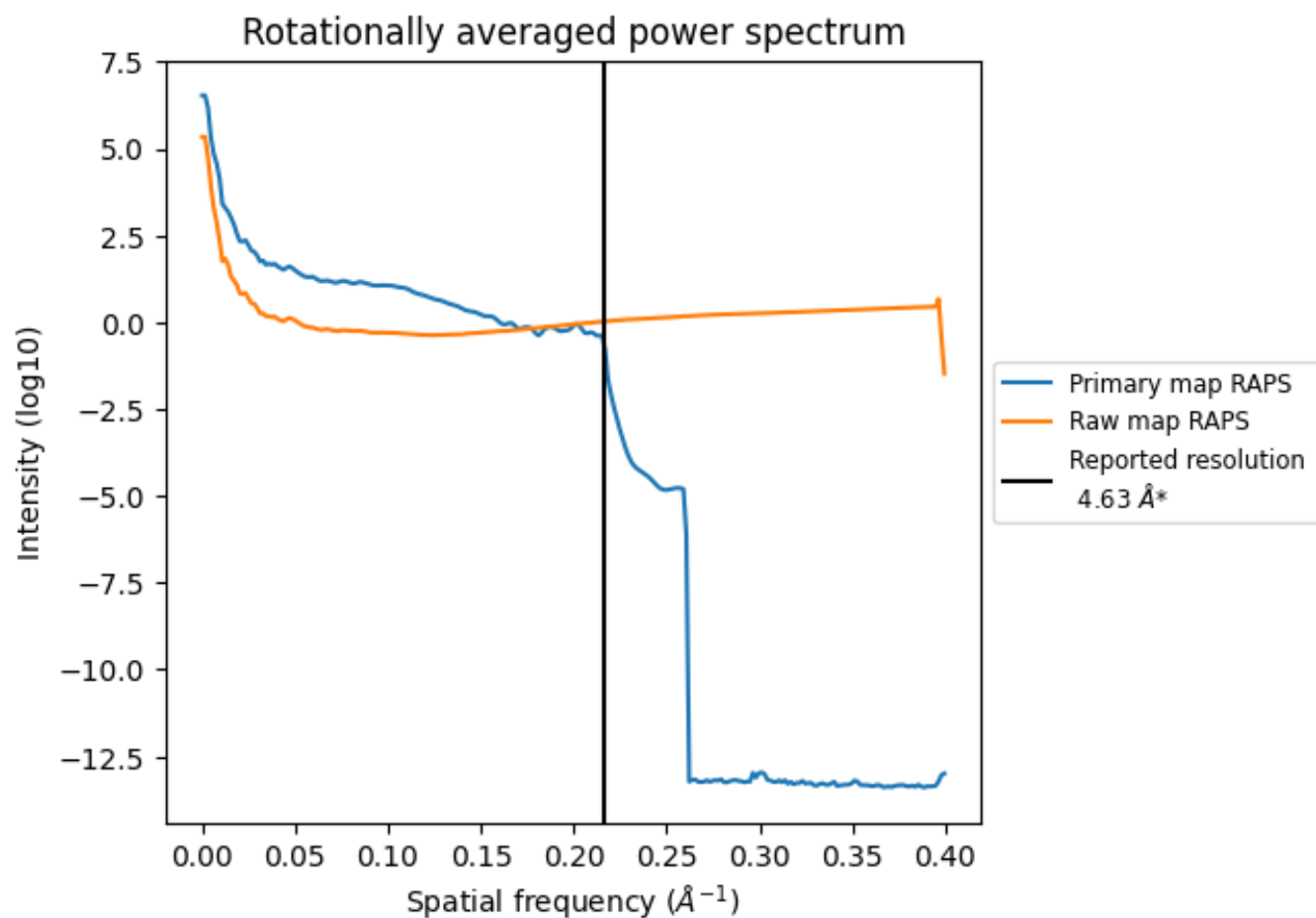
## 7.2 Volume estimate [i](#)



The volume at the recommended contour level is 207  $\text{nm}^3$ ; this corresponds to an approximate mass of 187 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 ⓘ

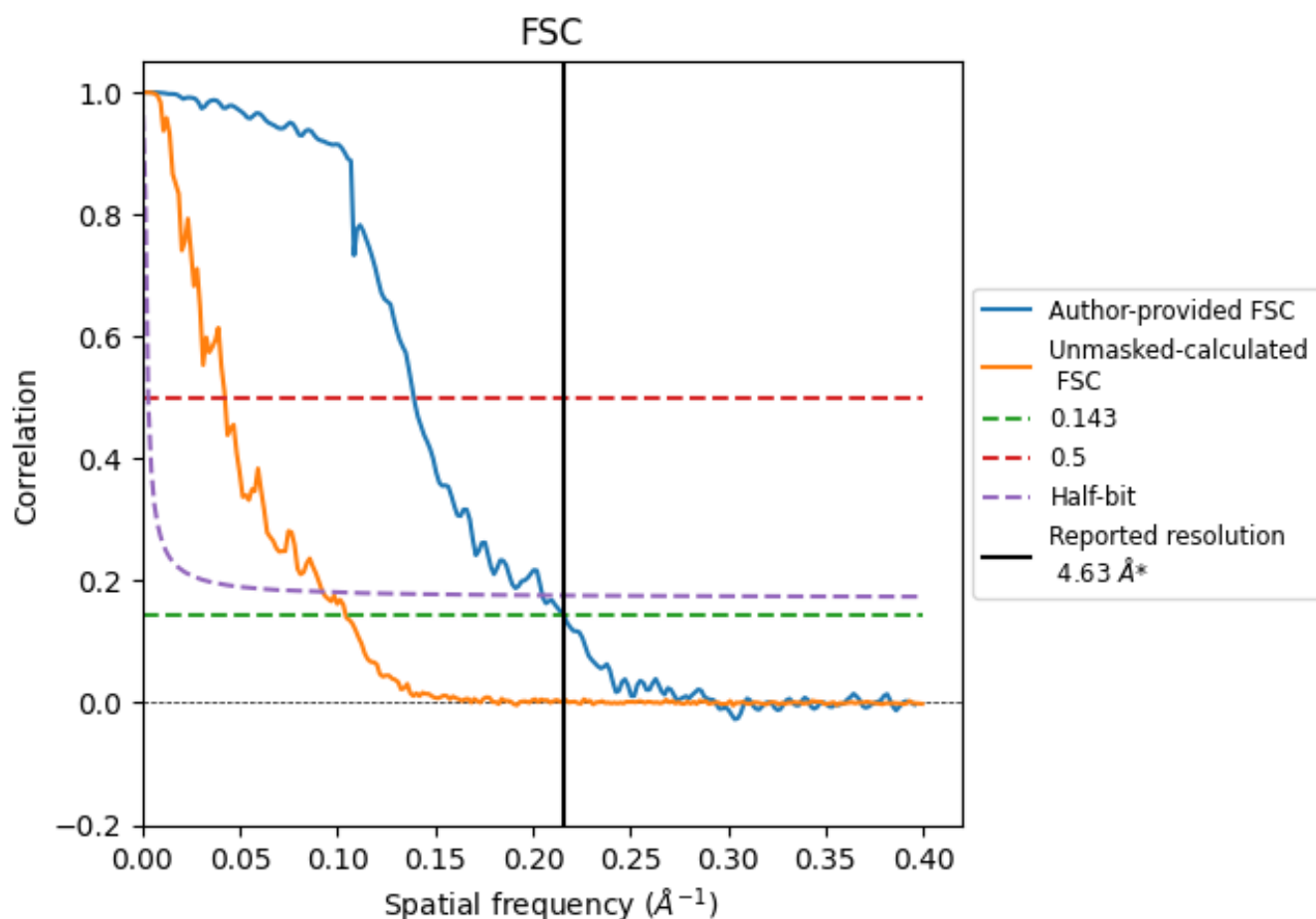


\*Reported resolution corresponds to spatial frequency of 0.216 Å<sup>-1</sup>

## 8 Fourier-Shell correlation [i](#)

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

### 8.1 FSC [i](#)



\*Reported resolution corresponds to spatial frequency of 0.216  $\text{\AA}^{-1}$

## 8.2 Resolution estimates [i](#)

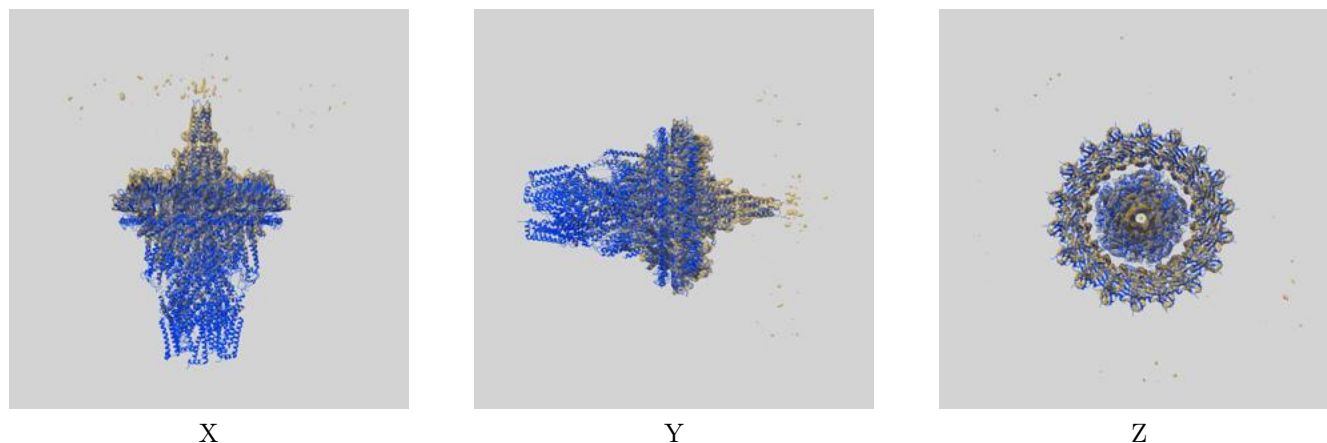
Resolution estimate (Å)	Estimation criterion (FSC cut-off)		
	0.143	0.5	Half-bit
Reported by author	4.63	-	-
Author-provided FSC curve	4.63	7.19	4.88
Unmasked-calculated*	9.59	23.64	10.82

\*Resolution estimate based on FSC curve calculated by comparison of deposited half-maps. The value from deposited half-maps intersecting FSC 0.143 CUT-OFF 9.59 differs from the reported value 4.63 by more than 10 %

## 9 Map-model fit [i](#)

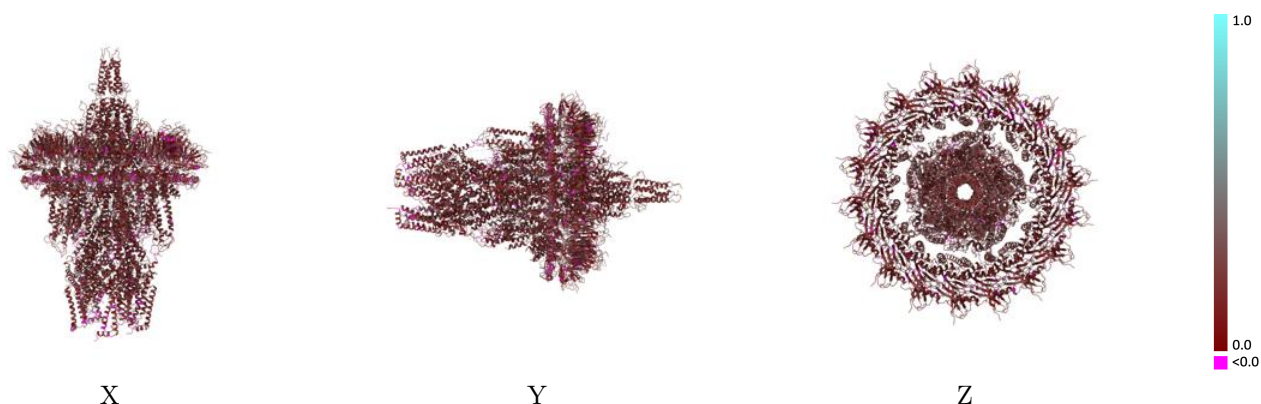
This section contains information regarding the fit between EMDB map EMD-49398 and PDB model 9NH0. Per-residue inclusion information can be found in section [3](#) on page [17](#).

### 9.1 Map-model overlay [i](#)



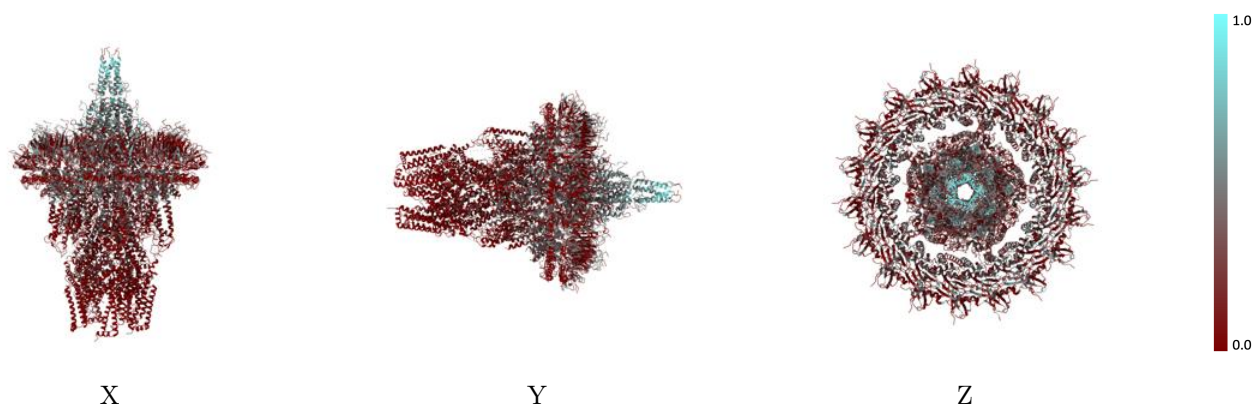
The images above show the 3D surface view of the map at the recommended contour level 0.16 at 50% transparency in yellow overlaid with a ribbon representation of the model coloured in blue. These images allow for the visual assessment of the quality of fit between the atomic model and the map.

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



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

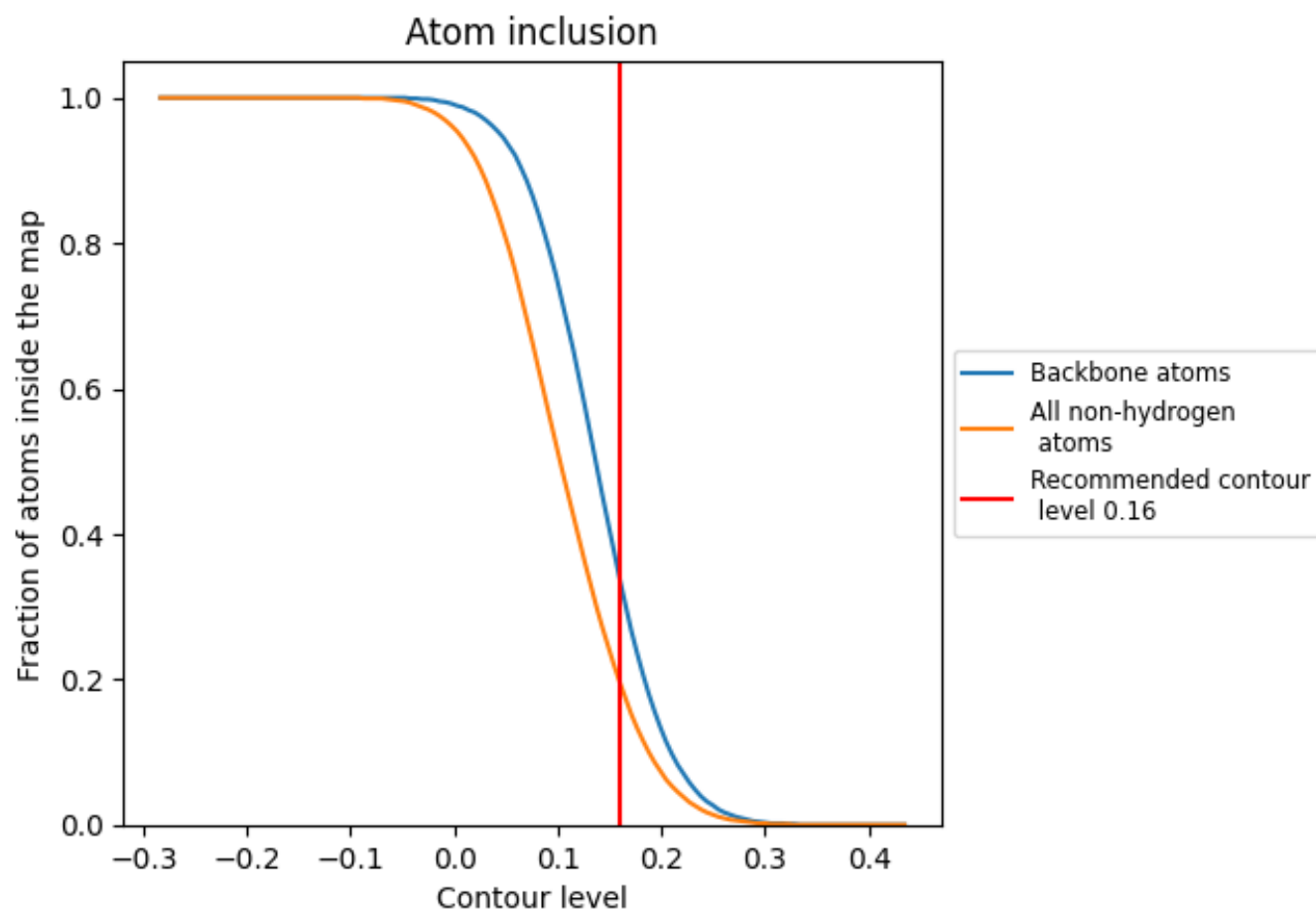
## 9.3 Atom inclusion mapped to coordinate model [i](#)



The images above show the model with each residue coloured according to its atom inclusion. This shows to what extent they are inside the map at the recommended contour level (0.16).









































































## 9.4 Atom inclusion [i](#)



At the recommended contour level, 34% of all backbone atoms, 20% 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 (0.16) and Q-score for the entire model and for each chain.

Chain	Atom inclusion	Q-score
All	 0.1950	 0.2010
Aa	 0.0000	 0.1650
Ab	 0.1020	 0.2220
Ac	 0.0000	 0.0490
Ad	 0.0000	 0.0010
Ae	 0.1820	 0.2010
Af	 0.2550	 0.2080
Ag	 0.0000	 0.1660
Ah	 0.0850	 0.2400
Ai	 0.0000	 0.1260
Aj	 0.0000	 0.0980
Ak	 0.1630	 0.1880
Al	 0.2810	 0.1980
Am	 0.0170	 0.1750
An	 0.0510	 0.1430
Ao	 0.0000	 0.0930
Ap	 0.0000	 0.1540
Aq	 0.1720	 0.1620
Ar	 0.2430	 0.1920
As	 0.0000	 0.1420
At	 0.0340	 0.1620
Au	 0.0000	 0.1150
Av	 0.0000	 0.0400
Aw	 0.1610	 0.1890
Ax	 0.2470	 0.1940
Ay	 0.0000	 0.2180
Az	 0.0000	 0.1730
Ba	 0.0000	 0.1710
Bb	 0.0000	 0.0750
Bc	 0.1000	 0.1690
Bd	 0.2420	 0.2050
Be	 0.0000	 0.1880
Bf	 0.0510	 0.2300
Bg	 0.0000	 0.0930
Bh	 0.0000	 0.0110























































































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Chain	Atom inclusion	Q-score
Bi	 0.1360	 0.1830
Bj	 0.2520	 0.2000
Bk	 0.0000	 0.1890
Bl	 0.0000	 0.1900
Bm	 0.0000	 0.0510
Bn	 0.0000	 -0.0730
Bo	 0.1480	 0.1810
Bp	 0.2170	 0.2080
Bq	 0.0000	 0.1350
Br	 0.0170	 0.2190
Bs	 0.0000	 0.1200
Bt	 0.0000	 -0.0550
Bu	 0.1060	 0.1760
Bv	 0.2710	 0.2130
Bw	 0.0000	 0.1840
Bx	 0.1190	 0.1440
By	 0.0000	 0.1100
Bz	 0.0000	 0.0110
Ca	 0.1500	 0.1780
Cb	 0.2840	 0.2030
Cc	 0.0000	 0.1760
Cd	 0.0340	 0.2270
Ce	 0.0000	 0.1640
Cf	 0.0000	 0.0570
Cg	 0.1290	 0.1830
Ch	 0.2440	 0.2080
Ci	 0.0000	 0.1620
Cj	 0.0000	 0.2090
Ck	 0.0000	 0.1040
Cl	 0.0000	 0.0000
Cm	 0.1440	 0.1950
Cn	 0.2430	 0.2060
Co	 0.0000	 0.2150
Cp	 0.0000	 0.1820
Cq	 0.0000	 0.1010
Cr	 0.0000	 0.0150
Cs	 0.1840	 0.1950
Ct	 0.2650	 0.2070
Cu	 0.0000	 0.1530
Cv	 0.0000	 0.0810
Cw	 0.0000	 0.1330
Cx	 0.0000	 0.2220















































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Chain	Atom inclusion	Q-score
Cy	 0.1590	 0.2020
Cz	 0.2540	 0.1970
Da	 0.0170	 0.1990
Db	 0.0340	 0.1430
Dc	 0.0000	 0.1050
Dd	 0.0000	 0.0980
De	 0.1440	 0.1760
Df	 0.2800	 0.2120
Dg	 0.0000	 0.1630
Dh	 0.0510	 0.1860
Di	 0.0000	 0.1560
Dj	 0.0000	 -0.0640
Dk	 0.1400	 0.2010
Dl	 0.3170	 0.2130
Dm	 0.0000	 0.1880
Dn	 0.0510	 0.2040
Do	 0.0000	 0.1220
Dp	 0.0000	 0.0190
Dq	 0.1780	 0.2090
Dr	 0.3140	 0.2180
Ds	 0.0000	 0.1690
Dt	 0.0340	 0.2350
Du	 0.0000	 0.0600
Dv	 0.0000	 0.0040
Dw	 0.1570	 0.1980
Dx	 0.2860	 0.2140
Dy	 0.0090	 0.1720
Dz	 0.0510	 0.2220
Ea	 0.0000	 0.0810
Eb	 0.0000	 0.0330
Ec	 0.1630	 0.1890
Ed	 0.3060	 0.2140
Ee	 0.3820	 0.2380
Ef	 0.2420	 0.2250
Eg	 0.3160	 0.2360
Eh	 0.2420	 0.2220
Ei	 0.2620	 0.1950
Ej	 0.2610	 0.2150
Ek	 0.3530	 0.2600
El	 0.4410	 0.2450
Em	 0.2930	 0.2330
En	 0.4410	 0.2370

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Chain	Atom inclusion	Q-score
Eo	 0.3460	 0.2380
Ep	 0.3570	 0.2350
Eq	 0.3080	 0.2110
Er	 0.3310	 0.2280
Es	 0.2450	 0.2300
Et	 0.3110	 0.2280
Eu	 0.2330	 0.2250
Ev	 0.4520	 0.2440
Ew	 0.3370	 0.2190
Ex	 0.3450	 0.2180
Ey	 0.3480	 0.2260
Ez	 0.3470	 0.2230
Fa	 0.3440	 0.2270
Fb	 0.1150	 0.1980
Fc	 0.0960	 0.1970
Fd	 0.1050	 0.1970
Fe	 0.1070	 0.1970
Ff	 0.1080	 0.1980
Fg	 0.0000	 0.1920
Fh	 0.0030	 0.1620
Fi	 0.0000	 0.1850
Fj	 0.0000	 0.1730
Fk	 0.0000	 0.1810