



Full wwPDB EM Validation Report ⓘ

Aug 15, 2024 – 03:58 PM EDT

PDB ID : 8T8O
EMDB ID : EMD-41100
Title : CCW Flagellar Switch Complex - FliF, FliG, FliM, and FliN forming 34-mer C-ring from Salmonella
Authors : Singh, P.K.; Iverson, T.M.
Deposited on : 2023-06-22
Resolution : 4.00 Å(reported)

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

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

A user guide is available at

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

The types of validation reports are described at

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

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

EMDB validation analysis : 0.0.1.dev92
MolProbity : 4.02b-467
Percentile statistics : 20191225.v01 (using entries in the PDB archive December 25th 2019)
MapQ : 1.9.13
Ideal geometry (proteins) : Engh & Huber (2001)
Ideal geometry (DNA, RNA) : Parkinson et al. (1996)
Validation Pipeline (wwPDB-VP) : 2.37.1

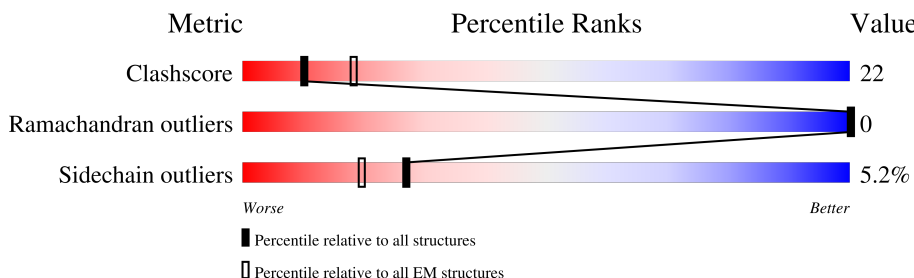
1 Overall quality at a glance

The following experimental techniques were used to determine the structure:

ELECTRON MICROSCOPY

The reported resolution of this entry is 4.00 Å.

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



Metric	Whole archive (#Entries)	EM structures (#Entries)
Clashscore	158937	4297
Ramachandran outliers	154571	4023
Sidechain outliers	154315	3826

The table below summarises the geometric issues observed across the polymeric chains and their fit to the map. The red, orange, yellow and green segments of the bar indicate the fraction of residues that contain outliers for ≥ 3 , 2, 1 and 0 types of geometric quality criteria respectively. A grey segment represents the fraction of residues that are not modelled. The numeric value for each fraction is indicated below the corresponding segment, with a dot representing fractions $\leq 5\%$. The upper red bar (where present) indicates the fraction of residues that have poor fit to the EM map (all-atom inclusion $< 40\%$). The numeric value is given above the bar.

Mol	Chain	Length	Quality of chain
1	A	47	<div> <div>49%</div> <div>100%</div> </div>
1	BC	47	<div> <div>49%</div> <div>100%</div> </div>
1	BF	47	<div> <div>51%</div> <div>100%</div> </div>
1	DB	47	<div> <div>49%</div> <div>100%</div> </div>
1	DE	47	<div> <div>51%</div> <div>100%</div> </div>
1	F	47	<div> <div>49%</div> <div>100%</div> </div>
1	FA	47	<div> <div>49%</div> <div>100%</div> </div>
1	FD	47	<div> <div>51%</div> <div>100%</div> </div>

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Mol	Chain	Length	Quality of chain
1	FG	47	49% 100%
1	HC	47	51% 98%
1	HF	47	51% 100%
1	I	47	49% 100%
1	JB	47	49% 100%
1	JE	47	51% 100%
1	LA	47	49% 100%
1	LD	47	51% 100%
1	LG	47	49% 98%
1	NC	47	51% 100%
1	NF	47	49% 98%
1	PB	47	49% 98%
1	PE	47	51% 98%
1	RA	47	49% 100%
1	RD	47	51% 98%
1	RG	47	49% 100%
1	S	47	49% 100%
1	TC	47	51% 100%
1	TF	47	49% 100%
1	VB	47	49% 100%
1	VE	47	51% 100%
1	XA	47	49% 98%
1	XD	47	51% 96%
1	Z	47	49% 98%
1	ZC	47	51% 100%

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Mol	Chain	Length	Quality of chain
1	ZF	47	<div>49%</div> <div>100%</div>
2	AA	331	<div>9%</div> <div>44%</div> <div>48%</div> <div>5%</div>
2	AD	331	<div>9%</div> <div>47%</div> <div>45%</div> <div>5%</div>
2	AG	331	<div>10%</div> <div>45%</div> <div>47%</div> <div>• •</div>
2	B	331	<div>9%</div> <div>49%</div> <div>44%</div> <div>• •</div>
2	CC	331	<div>9%</div> <div>46%</div> <div>46%</div> <div>5%</div>
2	CF	331	<div>10%</div> <div>45%</div> <div>47%</div> <div>• •</div>
2	EB	331	<div>9%</div> <div>47%</div> <div>47%</div> <div>• •</div>
2	EE	331	<div>10%</div> <div>47%</div> <div>44%</div> <div>5%</div>
2	G	331	<div>10%</div> <div>47%</div> <div>45%</div> <div>• •</div>
2	GA	331	<div>10%</div> <div>47%</div> <div>45%</div> <div>• •</div>
2	GD	331	<div>9%</div> <div>46%</div> <div>46%</div> <div>5%</div>
2	GG	331	<div>10%</div> <div>48%</div> <div>45%</div> <div>• •</div>
2	IC	331	<div>10%</div> <div>45%</div> <div>47%</div> <div>5%</div>
2	IF	331	<div>10%</div> <div>44%</div> <div>48%</div> <div>5%</div>
2	J	331	<div>10%</div> <div>45%</div> <div>47%</div> <div>5%</div>
2	KB	331	<div>9%</div> <div>45%</div> <div>47%</div> <div>5%</div>
2	KE	331	<div>9%</div> <div>48%</div> <div>45%</div> <div>• •</div>
2	MA	331	<div>10%</div> <div>49%</div> <div>44%</div> <div>• •</div>
2	MD	331	<div>10%</div> <div>44%</div> <div>48%</div> <div>• •</div>
2	MG	331	<div>10%</div> <div>47%</div> <div>45%</div> <div>5%</div>
2	OC	331	<div>9%</div> <div>49%</div> <div>43%</div> <div>5%</div>
2	OF	331	<div>10%</div> <div>41%</div> <div>49%</div> <div>6%</div>
2	QB	331	<div>10%</div> <div>44%</div> <div>48%</div> <div>• •</div>
2	QE	331	<div>10%</div> <div>46%</div> <div>48%</div> <div>• •</div>






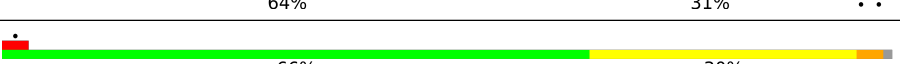
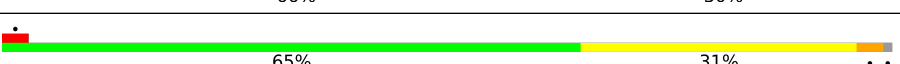
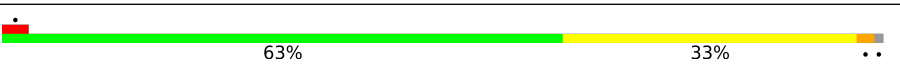


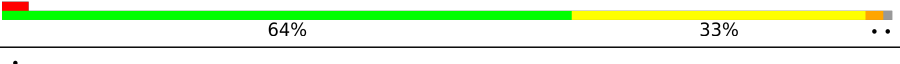
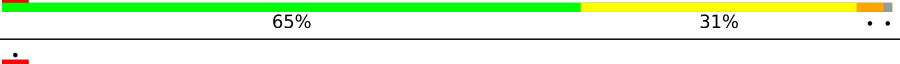

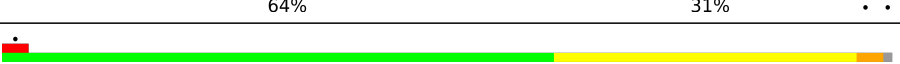
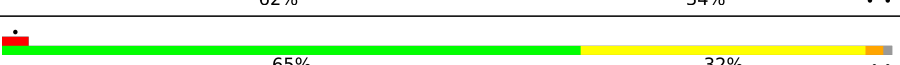



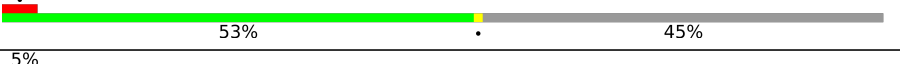






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Mol	Chain	Length	Quality of chain
2	SA	331	
2	SD	331	
2	SG	331	
2	T	331	
2	UC	331	
2	UF	331	
2	WB	331	
2	WE	331	
2	YA	331	
2	YD	331	
3	BA	300	
3	BD	300	
3	BG	300	
3	C	300	
3	DC	300	
3	DF	300	
3	FB	300	
3	FE	300	
3	HA	300	
3	HD	300	
3	HG	300	
3	JC	300	
3	JF	300	
3	K	300	
3	LB	300	







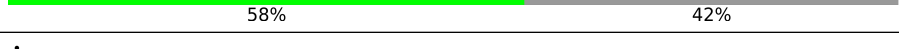
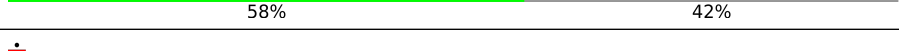
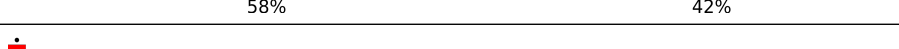
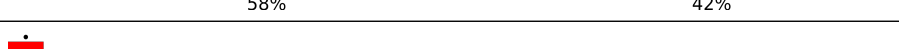
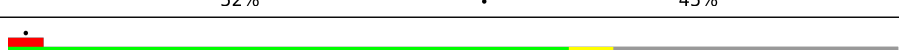

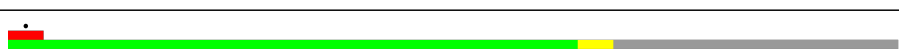

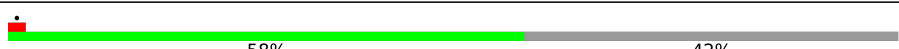





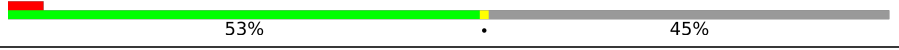
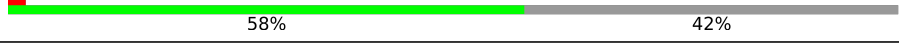



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Mol	Chain	Length	Quality of chain
3	LE	300	
3	M	300	
3	NA	300	
3	ND	300	
3	NG	300	
3	PC	300	
3	PF	300	
3	RB	300	
3	RE	300	
3	TA	300	
3	TD	300	
3	TG	300	
3	V	300	
3	VC	300	
3	VF	300	
3	XB	300	
3	XE	300	
3	ZA	300	
3	ZD	300	
4	AB	137	
4	AC	137	
4	AE	137	
4	AF	137	
4	BB	137	
4	BE	137	







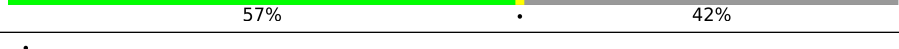
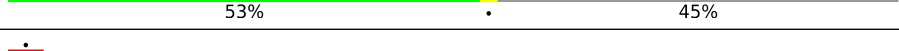
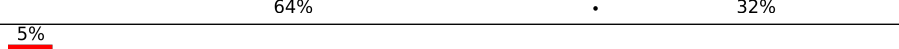
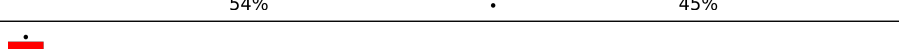
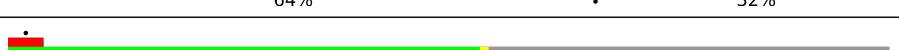

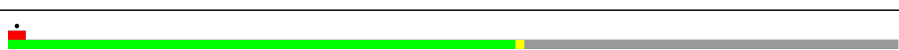

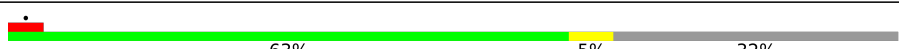





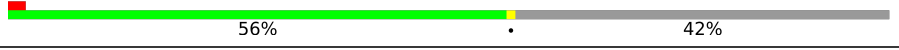
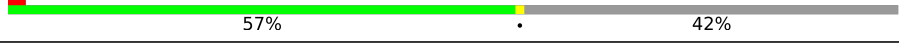



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Mol	Chain	Length	Quality of chain
4	CA	137	
4	CB	137	
4	CD	137	
4	CE	137	
4	CG	137	
4	D	137	
4	DA	137	
4	DD	137	
4	DG	137	
4	E	137	
4	EA	137	
4	EC	137	
4	ED	137	
4	EF	137	
4	EG	137	
4	FC	137	
4	FF	137	
4	GB	137	
4	GC	137	
4	GE	137	
4	GF	137	
4	H	137	
4	HB	137	
4	HE	137	
4	IA	137	


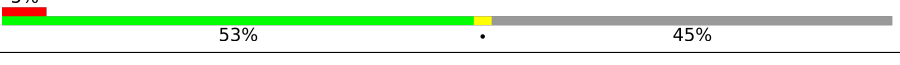



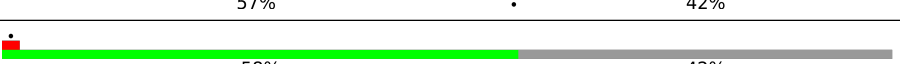




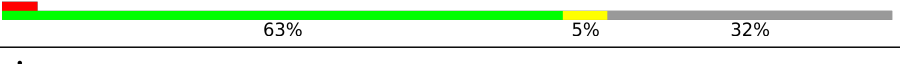
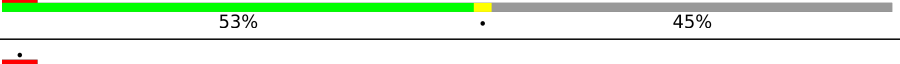

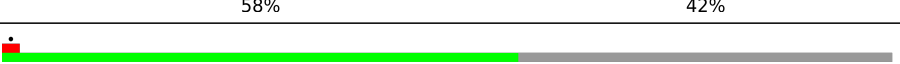
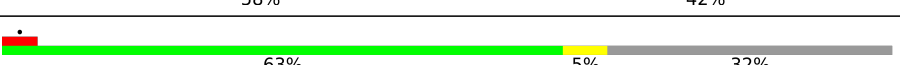
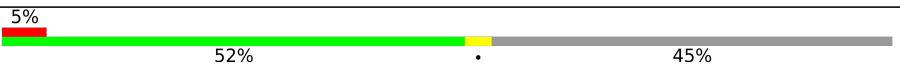









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Mol	Chain	Length	Quality of chain
4	IB	137	
4	ID	137	
4	IE	137	
4	IG	137	
4	JA	137	
4	JD	137	
4	JG	137	
4	KA	137	
4	KC	137	
4	KD	137	
4	KF	137	
4	KG	137	
4	L	137	
4	LC	137	
4	LF	137	
4	MB	137	
4	MC	137	
4	ME	137	
4	MF	137	
4	N	137	
4	NB	137	
4	NE	137	
4	O	137	
4	OA	137	
4	OB	137	



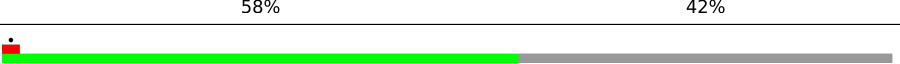
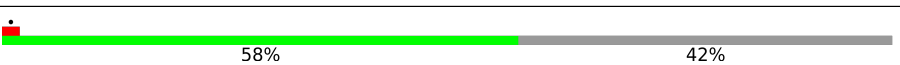
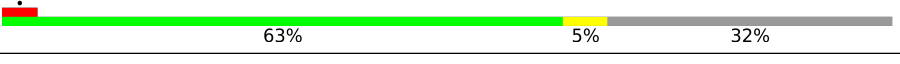

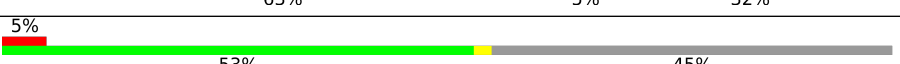

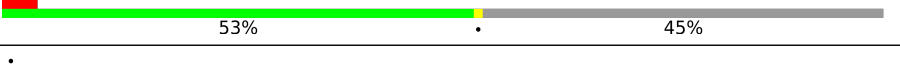
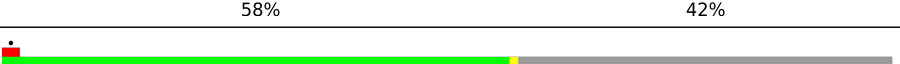
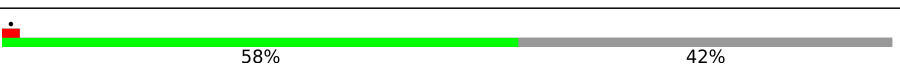
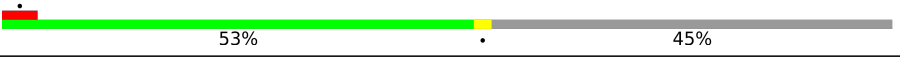

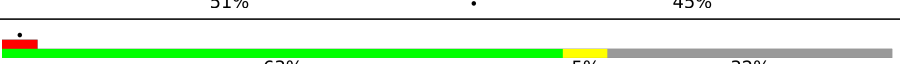

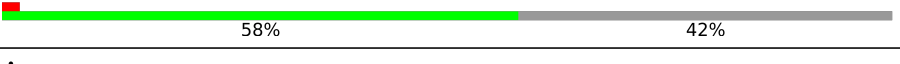
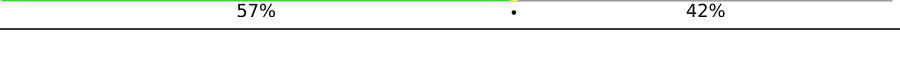




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Mol	Chain	Length	Quality of chain
4	OD	137	
4	OE	137	
4	OG	137	
4	P	137	
4	PA	137	
4	PD	137	
4	PG	137	
4	Q	137	
4	QA	137	
4	QC	137	
4	QD	137	
4	QF	137	
4	QG	137	
4	R	137	
4	RC	137	
4	RF	137	
4	SB	137	
4	SC	137	
4	SE	137	
4	SF	137	
4	TB	137	
4	TE	137	
4	UA	137	
4	UB	137	
4	UD	137	

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Mol	Chain	Length	Quality of chain
4	UE	137	
4	UG	137	
4	VA	137	
4	VD	137	
4	VG	137	
4	W	137	
4	WA	137	
4	WC	137	
4	WD	137	
4	WF	137	
4	WG	137	
4	X	137	
4	XC	137	
4	XF	137	
4	Y	137	
4	YB	137	
4	YC	137	
4	YE	137	
4	YF	137	
4	ZB	137	
4	ZE	137	

2 Entry composition [i](#)

There are 4 unique types of molecules in this entry. The entry contains 205020 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 Flagellar M-ring protein.

Mol	Chain	Residues	Atoms				AltConf	Trace
1	F	47	Total	C	N	O	0	0
			234	140	47	47		
1	A	47	Total	C	N	O	0	0
			234	140	47	47		
1	I	47	Total	C	N	O	0	0
			234	140	47	47		
1	S	47	Total	C	N	O	0	0
			234	140	47	47		
1	Z	47	Total	C	N	O	0	0
			234	140	47	47		
1	FA	47	Total	C	N	O	0	0
			234	140	47	47		
1	LA	47	Total	C	N	O	0	0
			234	140	47	47		
1	RA	47	Total	C	N	O	0	0
			234	140	47	47		
1	XA	47	Total	C	N	O	0	0
			234	140	47	47		
1	DB	47	Total	C	N	O	0	0
			234	140	47	47		
1	JB	47	Total	C	N	O	0	0
			234	140	47	47		
1	PB	47	Total	C	N	O	0	0
			234	140	47	47		
1	VB	47	Total	C	N	O	0	0
			234	140	47	47		
1	BC	47	Total	C	N	O	0	0
			234	140	47	47		
1	HC	47	Total	C	N	O	0	0
			234	140	47	47		
1	NC	47	Total	C	N	O	0	0
			234	140	47	47		
1	TC	47	Total	C	N	O	0	0
			234	140	47	47		

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Mol	Chain	Residues	Atoms				AltConf	Trace
1	ZC	47	Total	C	N	O	0	0
			234	140	47	47		
1	FD	47	Total	C	N	O	0	0
			234	140	47	47		
1	LD	47	Total	C	N	O	0	0
			234	140	47	47		
1	RD	47	Total	C	N	O	0	0
			234	140	47	47		
1	XD	47	Total	C	N	O	0	0
			234	140	47	47		
1	DE	47	Total	C	N	O	0	0
			234	140	47	47		
1	JE	47	Total	C	N	O	0	0
			234	140	47	47		
1	PE	47	Total	C	N	O	0	0
			234	140	47	47		
1	VE	47	Total	C	N	O	0	0
			234	140	47	47		
1	BF	47	Total	C	N	O	0	0
			234	140	47	47		
1	HF	47	Total	C	N	O	0	0
			234	140	47	47		
1	NF	47	Total	C	N	O	0	0
			234	140	47	47		
1	TF	47	Total	C	N	O	0	0
			234	140	47	47		
1	ZF	47	Total	C	N	O	0	0
			234	140	47	47		
1	FG	47	Total	C	N	O	0	0
			234	140	47	47		
1	LG	47	Total	C	N	O	0	0
			234	140	47	47		
1	RG	47	Total	C	N	O	0	0
			234	140	47	47		

- Molecule 2 is a protein called Flagellar motor switch protein FliG.

Mol	Chain	Residues	Atoms					AltConf	Trace
2	G	320	Total	C	N	O	S	0	0
			2496	1552	443	492	9		
2	B	320	Total	C	N	O	S	0	0
			2496	1552	443	492	9		

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Mol	Chain	Residues	Atoms					AltConf	Trace
2	J	320	Total	C	N	O	S	0	0
			2496	1552	443	492	9		
2	T	320	Total	C	N	O	S	0	0
			2496	1552	443	492	9		
2	AA	320	Total	C	N	O	S	0	0
			2496	1552	443	492	9		
2	GA	320	Total	C	N	O	S	0	0
			2496	1552	443	492	9		
2	MA	320	Total	C	N	O	S	0	0
			2496	1552	443	492	9		
2	SA	320	Total	C	N	O	S	0	0
			2496	1552	443	492	9		
2	YA	320	Total	C	N	O	S	0	0
			2496	1552	443	492	9		
2	EB	320	Total	C	N	O	S	0	0
			2496	1552	443	492	9		
2	KB	320	Total	C	N	O	S	0	0
			2496	1552	443	492	9		
2	QB	320	Total	C	N	O	S	0	0
			2496	1552	443	492	9		
2	WB	320	Total	C	N	O	S	0	0
			2496	1552	443	492	9		
2	CC	320	Total	C	N	O	S	0	0
			2496	1552	443	492	9		
2	IC	320	Total	C	N	O	S	0	0
			2496	1552	443	492	9		
2	OC	320	Total	C	N	O	S	0	0
			2496	1552	443	492	9		
2	UC	320	Total	C	N	O	S	0	0
			2496	1552	443	492	9		
2	AD	320	Total	C	N	O	S	0	0
			2496	1552	443	492	9		
2	GD	320	Total	C	N	O	S	0	0
			2496	1552	443	492	9		
2	MD	320	Total	C	N	O	S	0	0
			2496	1552	443	492	9		
2	SD	320	Total	C	N	O	S	0	0
			2496	1552	443	492	9		
2	YD	320	Total	C	N	O	S	0	0
			2496	1552	443	492	9		
2	EE	320	Total	C	N	O	S	0	0
			2496	1552	443	492	9		

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Mol	Chain	Residues	Atoms					AltConf	Trace
2	KE	320	Total	C	N	O	S	0	0
			2496	1552	443	492	9		
2	QE	320	Total	C	N	O	S	0	0
			2496	1552	443	492	9		
2	WE	320	Total	C	N	O	S	0	0
			2496	1552	443	492	9		
2	CF	320	Total	C	N	O	S	0	0
			2496	1552	443	492	9		
2	IF	320	Total	C	N	O	S	0	0
			2496	1552	443	492	9		
2	OF	320	Total	C	N	O	S	0	0
			2496	1552	443	492	9		
2	UF	320	Total	C	N	O	S	0	0
			2496	1552	443	492	9		
2	AG	320	Total	C	N	O	S	0	0
			2496	1552	443	492	9		
2	GG	320	Total	C	N	O	S	0	0
			2496	1552	443	492	9		
2	MG	320	Total	C	N	O	S	0	0
			2496	1552	443	492	9		
2	SG	320	Total	C	N	O	S	0	0
			2496	1552	443	492	9		

- Molecule 3 is a protein called FlIM, Flagellar motor switch protein FlIM.

Mol	Chain	Residues	Atoms					AltConf	Trace
3	M	296	Total	C	N	O	S	0	0
			2086	1314	383	384	5		
3	C	296	Total	C	N	O	S	0	0
			2086	1314	383	384	5		
3	K	296	Total	C	N	O	S	0	0
			2086	1314	383	384	5		
3	V	296	Total	C	N	O	S	0	0
			2086	1314	383	384	5		
3	BA	296	Total	C	N	O	S	0	0
			2086	1314	383	384	5		
3	HA	296	Total	C	N	O	S	0	0
			2086	1314	383	384	5		
3	NA	296	Total	C	N	O	S	0	0
			2086	1314	383	384	5		
3	TA	296	Total	C	N	O	S	0	0
			2086	1314	383	384	5		

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Mol	Chain	Residues	Atoms					AltConf	Trace
3	ZA	296	Total 2086	C 1314	N 383	O 384	S 5	0	0
3	FB	296	Total 2086	C 1314	N 383	O 384	S 5	0	0
3	LB	296	Total 2086	C 1314	N 383	O 384	S 5	0	0
3	RB	296	Total 2086	C 1314	N 383	O 384	S 5	0	0
3	XB	296	Total 2086	C 1314	N 383	O 384	S 5	0	0
3	DC	296	Total 2086	C 1314	N 383	O 384	S 5	0	0
3	JC	296	Total 2086	C 1314	N 383	O 384	S 5	0	0
3	PC	296	Total 2086	C 1314	N 383	O 384	S 5	0	0
3	VC	296	Total 2086	C 1314	N 383	O 384	S 5	0	0
3	BD	296	Total 2086	C 1314	N 383	O 384	S 5	0	0
3	HD	296	Total 2086	C 1314	N 383	O 384	S 5	0	0
3	ND	296	Total 2086	C 1314	N 383	O 384	S 5	0	0
3	TD	296	Total 2086	C 1314	N 383	O 384	S 5	0	0
3	ZD	296	Total 2086	C 1314	N 383	O 384	S 5	0	0
3	FE	296	Total 2086	C 1314	N 383	O 384	S 5	0	0
3	LE	296	Total 2086	C 1314	N 383	O 384	S 5	0	0
3	RE	296	Total 2086	C 1314	N 383	O 384	S 5	0	0
3	XE	296	Total 2086	C 1314	N 383	O 384	S 5	0	0
3	DF	296	Total 2086	C 1314	N 383	O 384	S 5	0	0
3	JF	296	Total 2086	C 1314	N 383	O 384	S 5	0	0
3	PF	296	Total 2086	C 1314	N 383	O 384	S 5	0	0

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Mol	Chain	Residues	Atoms					AltConf	Trace
3	VF	296	Total	C	N	O	S	0	0
			2086	1314	383	384	5		
3	BG	296	Total	C	N	O	S	0	0
			2086	1314	383	384	5		
3	HG	296	Total	C	N	O	S	0	0
			2086	1314	383	384	5		
3	NG	296	Total	C	N	O	S	0	0
			2086	1314	383	384	5		
3	TG	296	Total	C	N	O	S	0	0
			2086	1314	383	384	5		

- Molecule 4 is a protein called Flagellar motor switch protein FliN.

Mol	Chain	Residues	Atoms				AltConf	Trace
4	N	93	Total	C	N	O	0	0
			456	270	93	93		
4	P	79	Total	C	N	O	0	0
			389	230	79	80		
4	Q	75	Total	C	N	O	0	0
			369	218	75	76		
4	D	93	Total	C	N	O	0	0
			456	270	93	93		
4	E	79	Total	C	N	O	0	0
			389	230	79	80		
4	H	75	Total	C	N	O	0	0
			369	218	75	76		
4	L	93	Total	C	N	O	0	0
			456	270	93	93		
4	O	79	Total	C	N	O	0	0
			389	230	79	80		
4	R	75	Total	C	N	O	0	0
			369	218	75	76		
4	W	93	Total	C	N	O	0	0
			456	270	93	93		
4	X	79	Total	C	N	O	0	0
			389	230	79	80		
4	Y	75	Total	C	N	O	0	0
			369	218	75	76		
4	CA	93	Total	C	N	O	0	0
			456	270	93	93		
4	DA	79	Total	C	N	O	0	0
			389	230	79	80		

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Mol	Chain	Residues	Atoms				AltConf	Trace
4	EA	75	Total	C	N	O	0	0
			369	218	75	76		
4	IA	93	Total	C	N	O	0	0
			456	270	93	93		
4	JA	79	Total	C	N	O	0	0
			389	230	79	80		
4	KA	75	Total	C	N	O	0	0
			369	218	75	76		
4	OA	93	Total	C	N	O	0	0
			456	270	93	93		
4	PA	79	Total	C	N	O	0	0
			389	230	79	80		
4	QA	75	Total	C	N	O	0	0
			369	218	75	76		
4	UA	93	Total	C	N	O	0	0
			456	270	93	93		
4	VA	79	Total	C	N	O	0	0
			389	230	79	80		
4	WA	75	Total	C	N	O	0	0
			369	218	75	76		
4	AB	93	Total	C	N	O	0	0
			456	270	93	93		
4	BB	79	Total	C	N	O	0	0
			389	230	79	80		
4	CB	75	Total	C	N	O	0	0
			369	218	75	76		
4	GB	93	Total	C	N	O	0	0
			456	270	93	93		
4	HB	79	Total	C	N	O	0	0
			389	230	79	80		
4	IB	75	Total	C	N	O	0	0
			369	218	75	76		
4	MB	93	Total	C	N	O	0	0
			456	270	93	93		
4	NB	79	Total	C	N	O	0	0
			389	230	79	80		
4	OB	75	Total	C	N	O	0	0
			369	218	75	76		
4	SB	93	Total	C	N	O	0	0
			456	270	93	93		
4	TB	79	Total	C	N	O	0	0
			389	230	79	80		

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Mol	Chain	Residues	Atoms				AltConf	Trace
4	UB	75	Total	C	N	O	0	0
			369	218	75	76		
4	YB	93	Total	C	N	O	0	0
			456	270	93	93		
4	ZB	79	Total	C	N	O	0	0
			389	230	79	80		
4	AC	75	Total	C	N	O	0	0
			369	218	75	76		
4	EC	93	Total	C	N	O	0	0
			456	270	93	93		
4	FC	79	Total	C	N	O	0	0
			389	230	79	80		
4	GC	75	Total	C	N	O	0	0
			369	218	75	76		
4	KC	93	Total	C	N	O	0	0
			456	270	93	93		
4	LC	79	Total	C	N	O	0	0
			389	230	79	80		
4	MC	75	Total	C	N	O	0	0
			369	218	75	76		
4	QC	93	Total	C	N	O	0	0
			456	270	93	93		
4	RC	79	Total	C	N	O	0	0
			389	230	79	80		
4	SC	75	Total	C	N	O	0	0
			369	218	75	76		
4	WC	93	Total	C	N	O	0	0
			456	270	93	93		
4	XC	79	Total	C	N	O	0	0
			389	230	79	80		
4	YC	75	Total	C	N	O	0	0
			369	218	75	76		
4	CD	93	Total	C	N	O	0	0
			456	270	93	93		
4	DD	79	Total	C	N	O	0	0
			389	230	79	80		
4	ED	75	Total	C	N	O	0	0
			369	218	75	76		
4	ID	93	Total	C	N	O	0	0
			456	270	93	93		
4	JD	79	Total	C	N	O	0	0
			389	230	79	80		

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Mol	Chain	Residues	Atoms				AltConf	Trace
4	KD	75	Total	C	N	O	0	0
			369	218	75	76		
4	OD	93	Total	C	N	O	0	0
			456	270	93	93		
4	PD	79	Total	C	N	O	0	0
			389	230	79	80		
4	QD	75	Total	C	N	O	0	0
			369	218	75	76		
4	UD	93	Total	C	N	O	0	0
			456	270	93	93		
4	VD	79	Total	C	N	O	0	0
			389	230	79	80		
4	WD	75	Total	C	N	O	0	0
			369	218	75	76		
4	AE	93	Total	C	N	O	0	0
			456	270	93	93		
4	BE	79	Total	C	N	O	0	0
			389	230	79	80		
4	CE	75	Total	C	N	O	0	0
			369	218	75	76		
4	GE	93	Total	C	N	O	0	0
			456	270	93	93		
4	HE	79	Total	C	N	O	0	0
			389	230	79	80		
4	IE	75	Total	C	N	O	0	0
			369	218	75	76		
4	ME	93	Total	C	N	O	0	0
			456	270	93	93		
4	NE	79	Total	C	N	O	0	0
			389	230	79	80		
4	OE	75	Total	C	N	O	0	0
			369	218	75	76		
4	SE	93	Total	C	N	O	0	0
			456	270	93	93		
4	TE	79	Total	C	N	O	0	0
			389	230	79	80		
4	UE	75	Total	C	N	O	0	0
			369	218	75	76		
4	YE	93	Total	C	N	O	0	0
			456	270	93	93		
4	ZE	79	Total	C	N	O	0	0
			389	230	79	80		

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Mol	Chain	Residues	Atoms				AltConf	Trace
4	AF	75	Total	C	N	O	0	0
			369	218	75	76		
4	EF	93	Total	C	N	O	0	0
			456	270	93	93		
4	FF	79	Total	C	N	O	0	0
			389	230	79	80		
4	GF	75	Total	C	N	O	0	0
			369	218	75	76		
4	KF	93	Total	C	N	O	0	0
			456	270	93	93		
4	LF	79	Total	C	N	O	0	0
			389	230	79	80		
4	MF	75	Total	C	N	O	0	0
			369	218	75	76		
4	QF	93	Total	C	N	O	0	0
			456	270	93	93		
4	RF	79	Total	C	N	O	0	0
			389	230	79	80		
4	SF	75	Total	C	N	O	0	0
			369	218	75	76		
4	WF	93	Total	C	N	O	0	0
			456	270	93	93		
4	XF	79	Total	C	N	O	0	0
			389	230	79	80		
4	YF	75	Total	C	N	O	0	0
			369	218	75	76		
4	CG	93	Total	C	N	O	0	0
			456	270	93	93		
4	DG	79	Total	C	N	O	0	0
			389	230	79	80		
4	EG	75	Total	C	N	O	0	0
			369	218	75	76		
4	IG	93	Total	C	N	O	0	0
			456	270	93	93		
4	JG	79	Total	C	N	O	0	0
			389	230	79	80		
4	KG	75	Total	C	N	O	0	0
			369	218	75	76		
4	OG	93	Total	C	N	O	0	0
			456	270	93	93		
4	PG	79	Total	C	N	O	0	0
			389	230	79	80		

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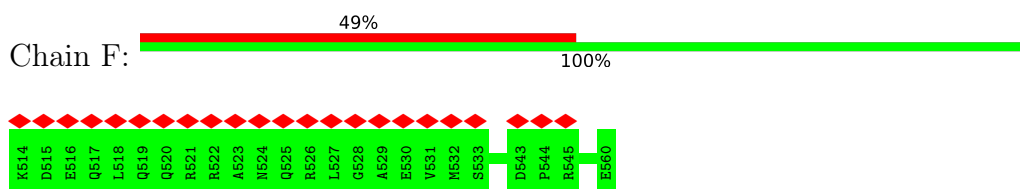
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Mol	Chain	Residues	Atoms				AltConf	Trace
4	QG	75	Total 369	C 218	N 75	O 76	0	0
4	UG	93	Total 456	C 270	N 93	O 93	0	0
4	VG	79	Total 389	C 230	N 79	O 80	0	0
4	WG	75	Total 369	C 218	N 75	O 76	0	0

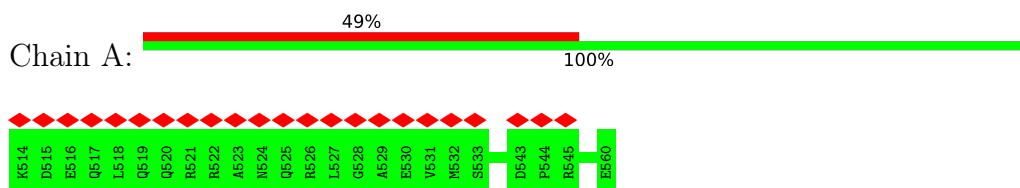
3 Residue-property plots

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

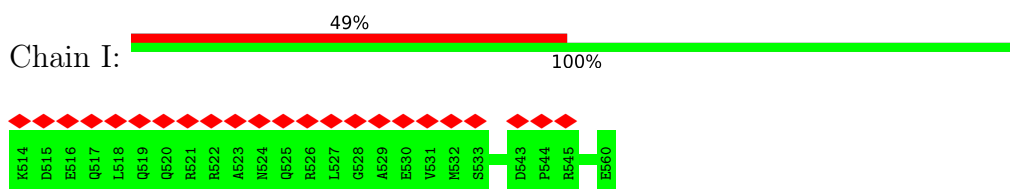
- Molecule 1: Flagellar M-ring protein



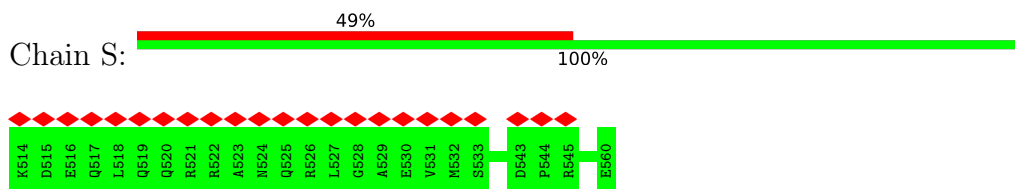
- Molecule 1: Flagellar M-ring protein



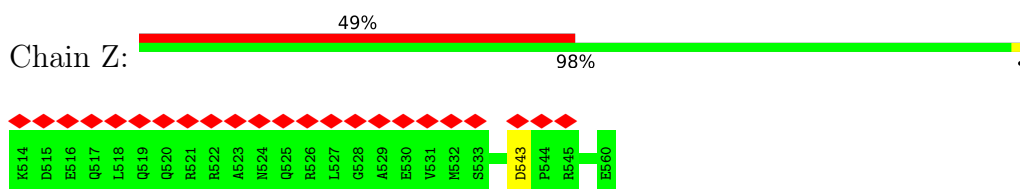
- Molecule 1: Flagellar M-ring protein



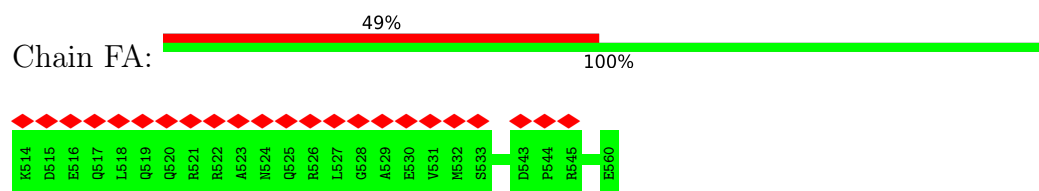
- Molecule 1: Flagellar M-ring protein



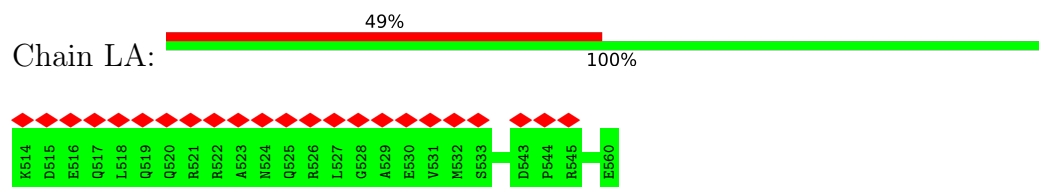
- Molecule 1: Flagellar M-ring protein



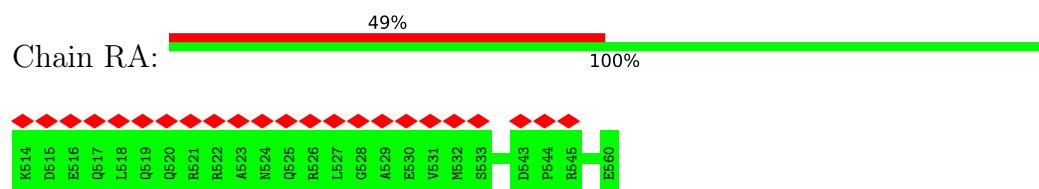
• Molecule 1: Flagellar M-ring protein



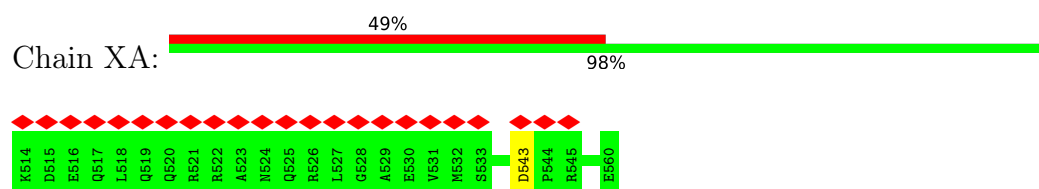
• Molecule 1: Flagellar M-ring protein



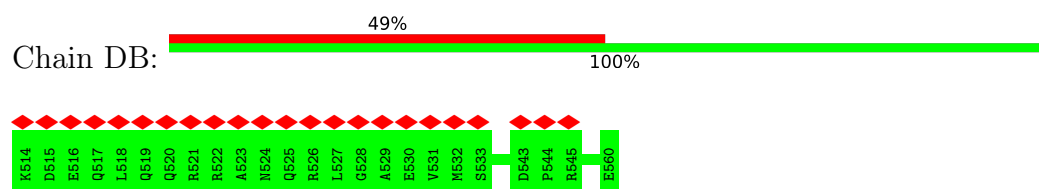
• Molecule 1: Flagellar M-ring protein



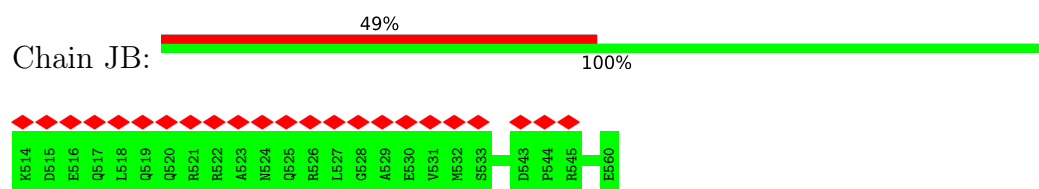
• Molecule 1: Flagellar M-ring protein



• Molecule 1: Flagellar M-ring protein

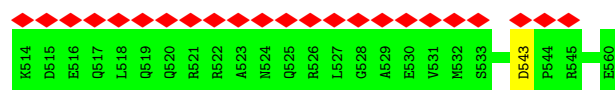


• Molecule 1: Flagellar M-ring protein

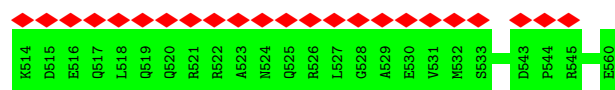


• Molecule 1: Flagellar M-ring protein

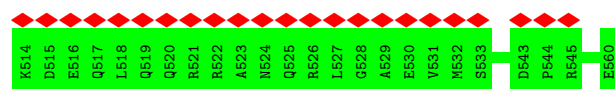




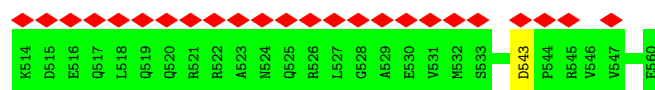
- Molecule 1: Flagellar M-ring protein



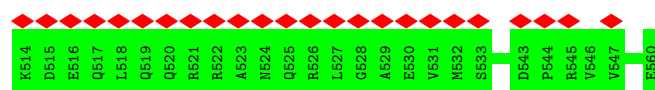
- Molecule 1: Flagellar M-ring protein



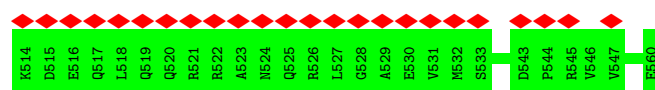
- Molecule 1: Flagellar M-ring protein



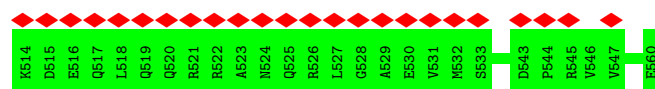
- Molecule 1: Flagellar M-ring protein



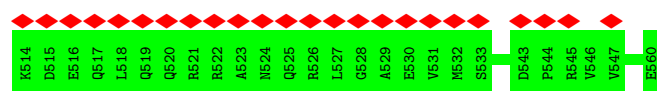
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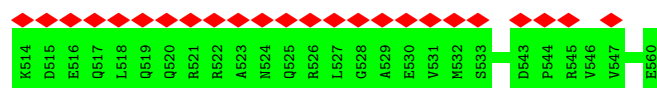
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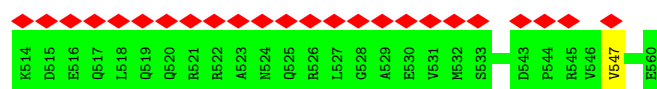
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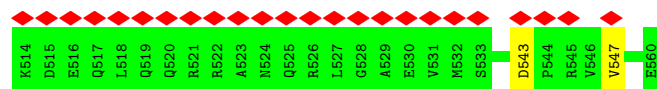
- Molecule 1: Flagellar M-ring protein



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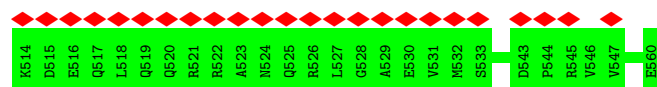
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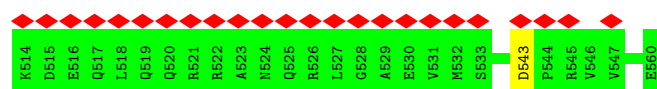
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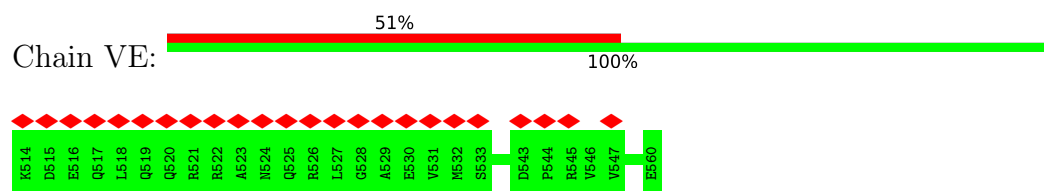
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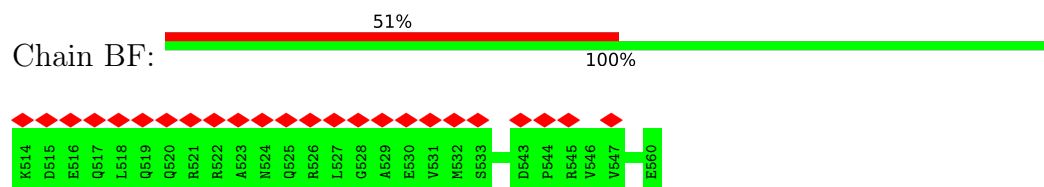
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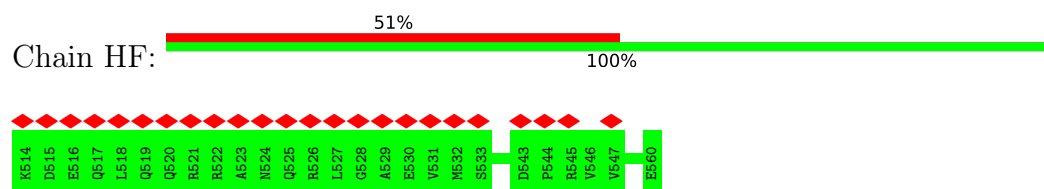
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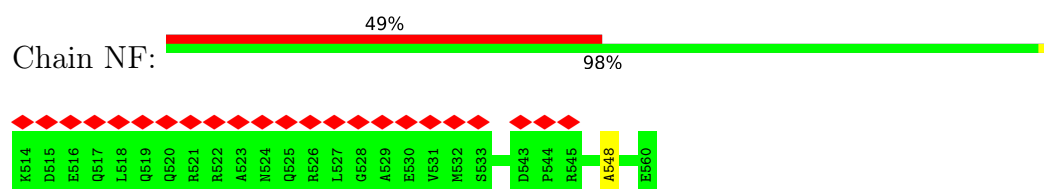
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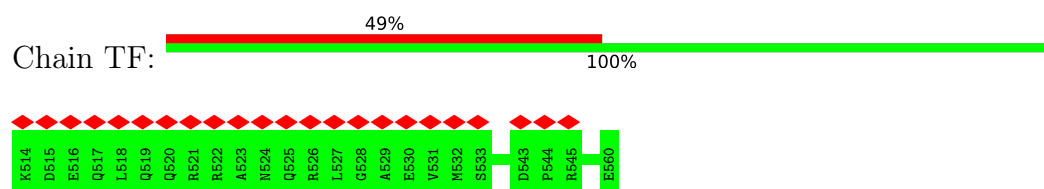
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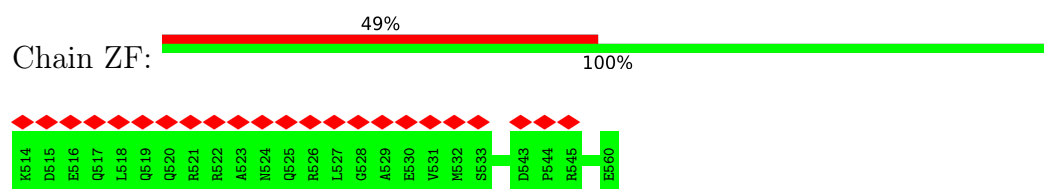
• Molecule 1: Flagellar M-ring protein



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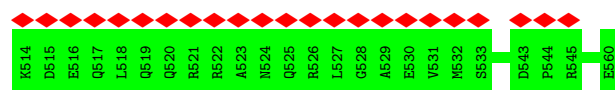


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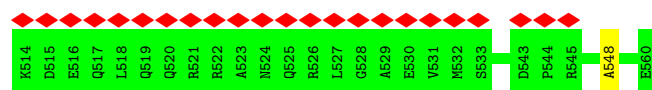


• Molecule 1: Flagellar M-ring protein

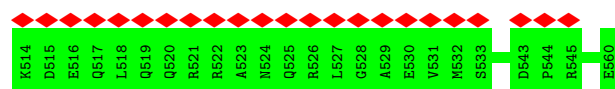




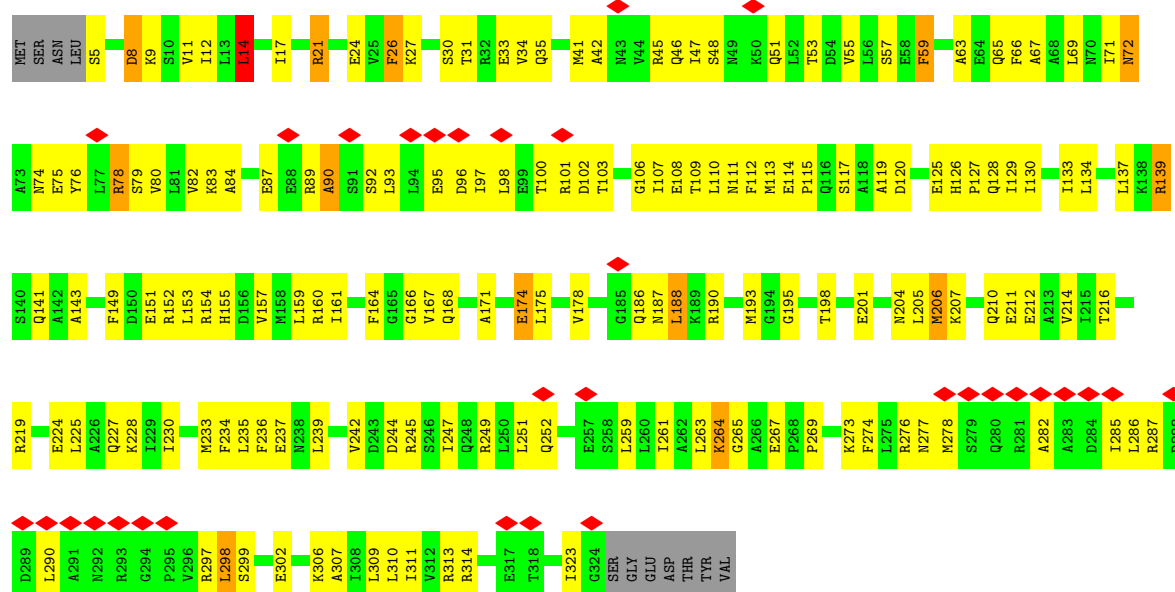
• Molecule 1: Flagellar M-ring protein



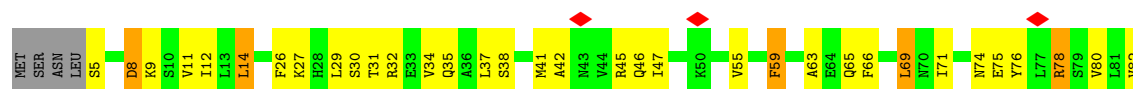
• Molecule 1: Flagellar M-ring protein

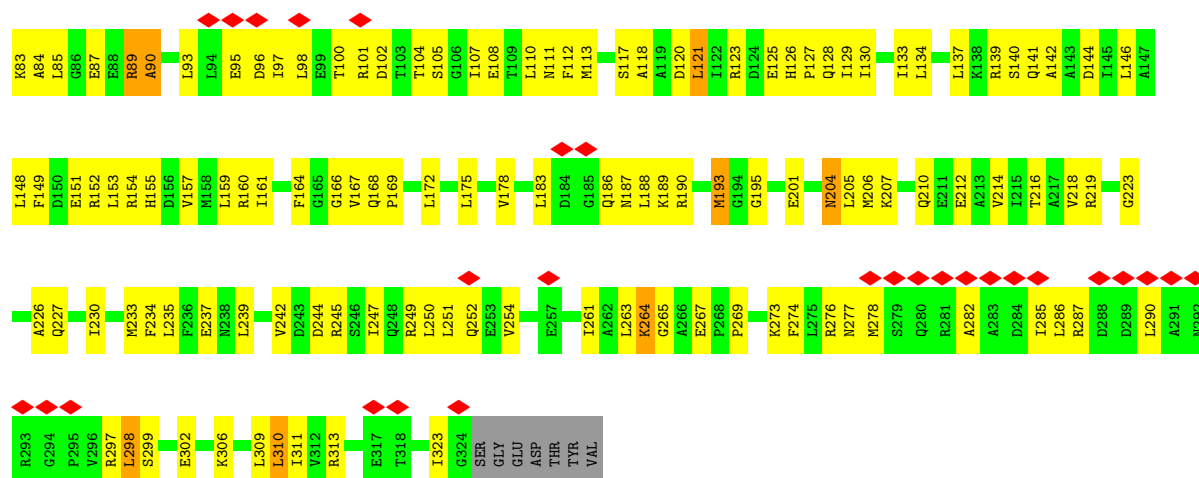


• Molecule 2: Flagellar motor switch protein FliG

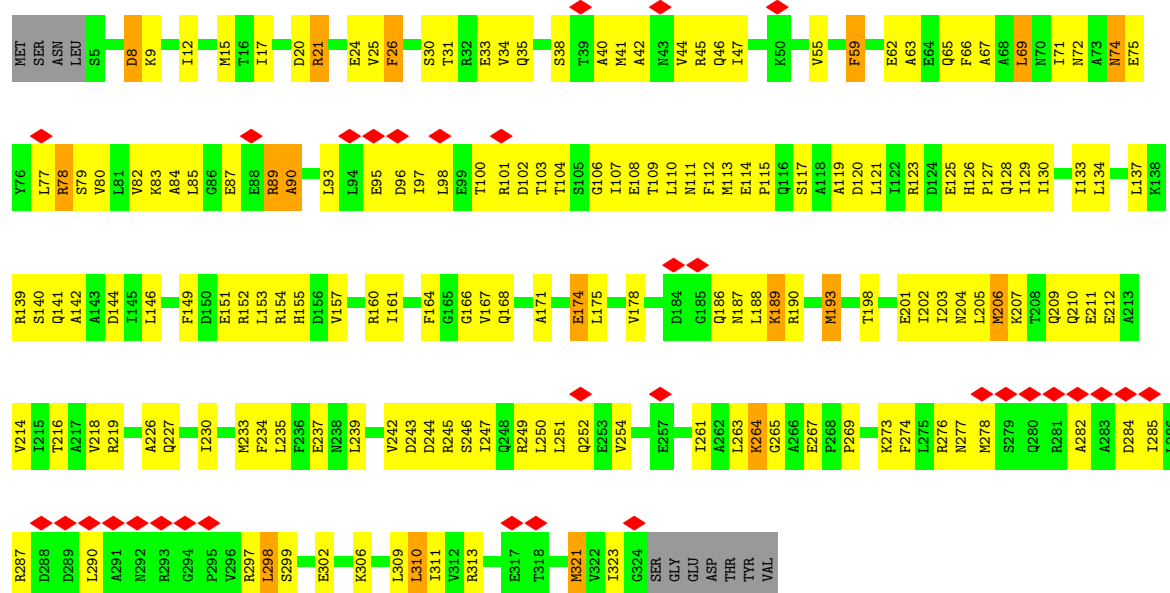
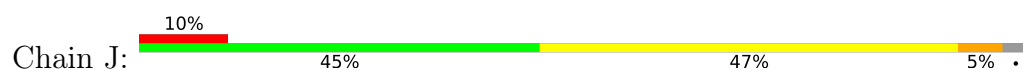


• Molecule 2: Flagellar motor switch protein FliG

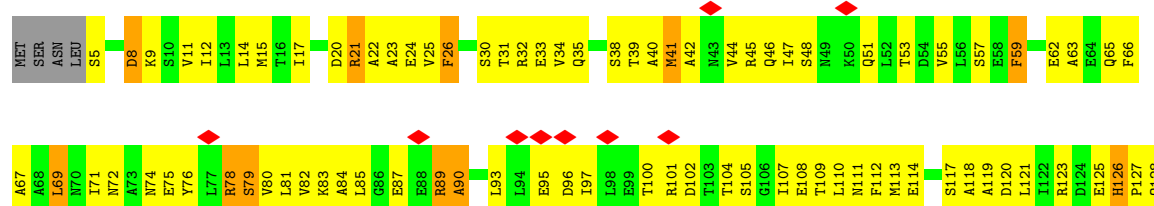
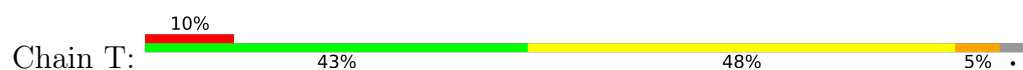


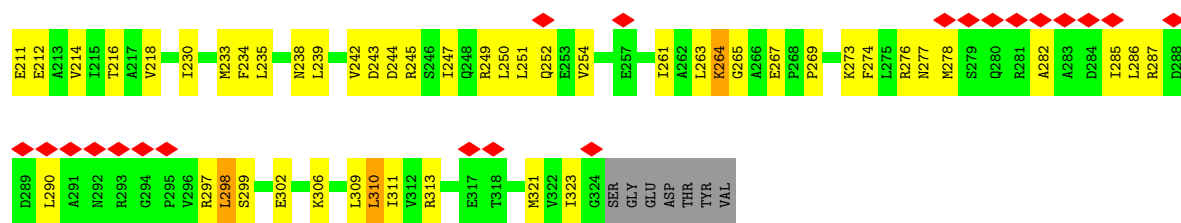


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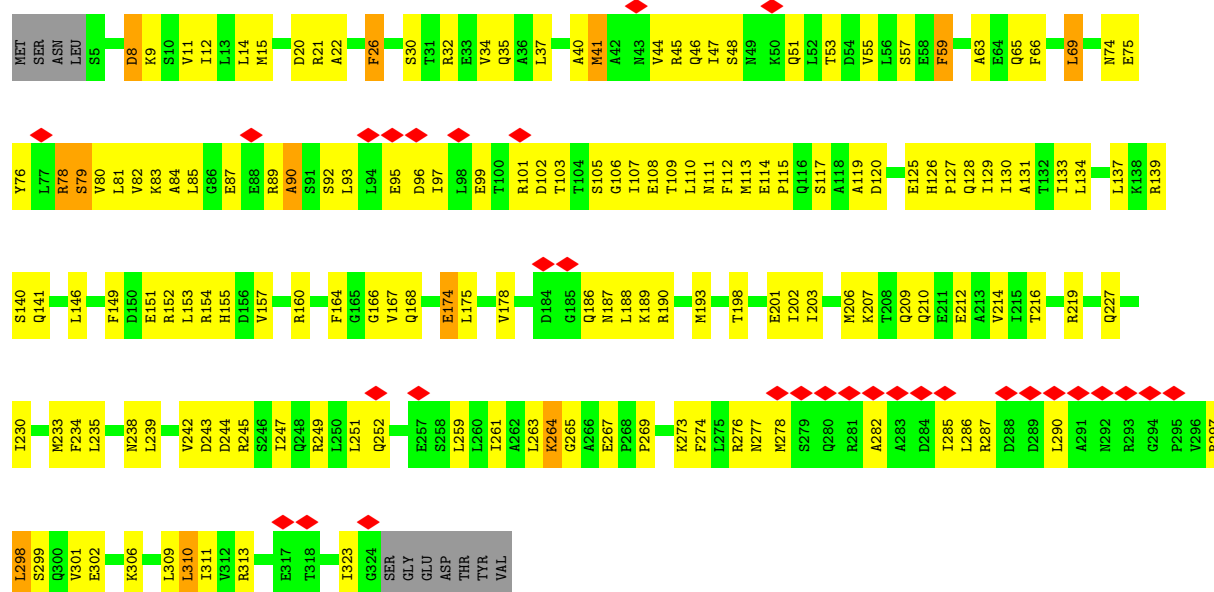


• Molecule 2: Flagellar motor switch protein FlgI

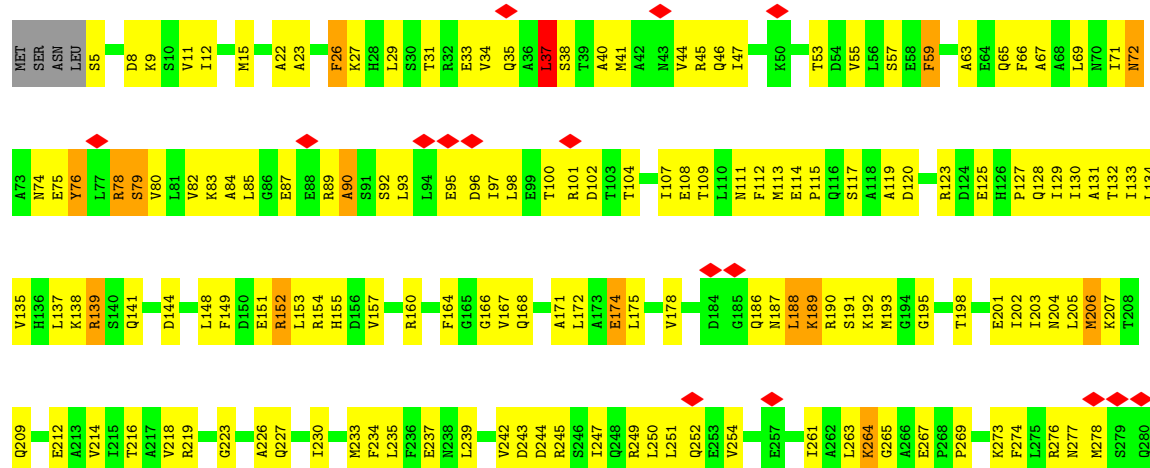


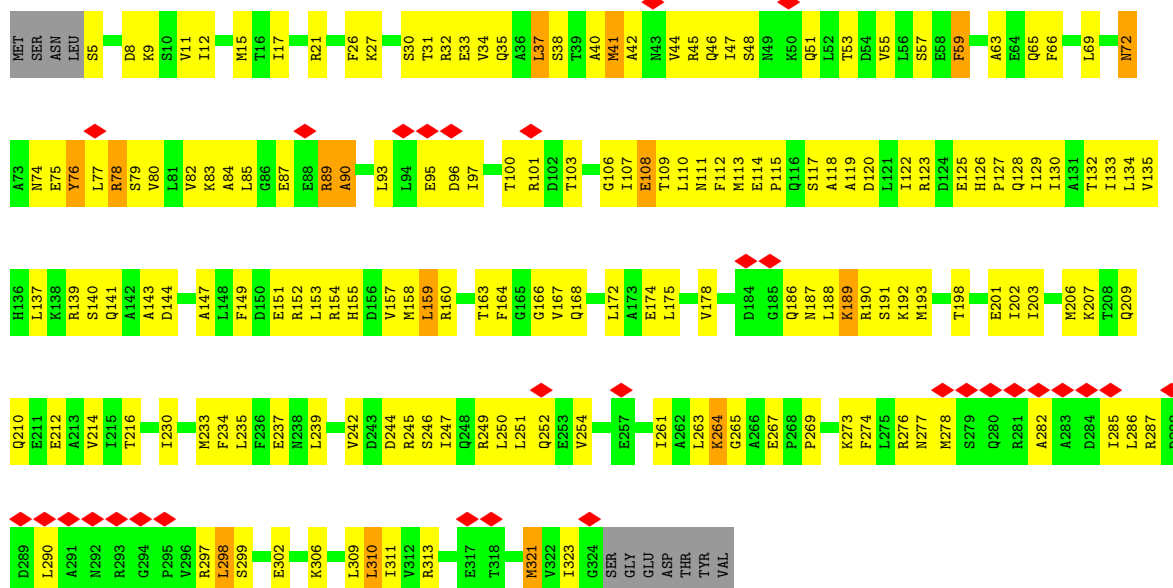


• Molecule 2: Flagellar motor switch protein FlhG

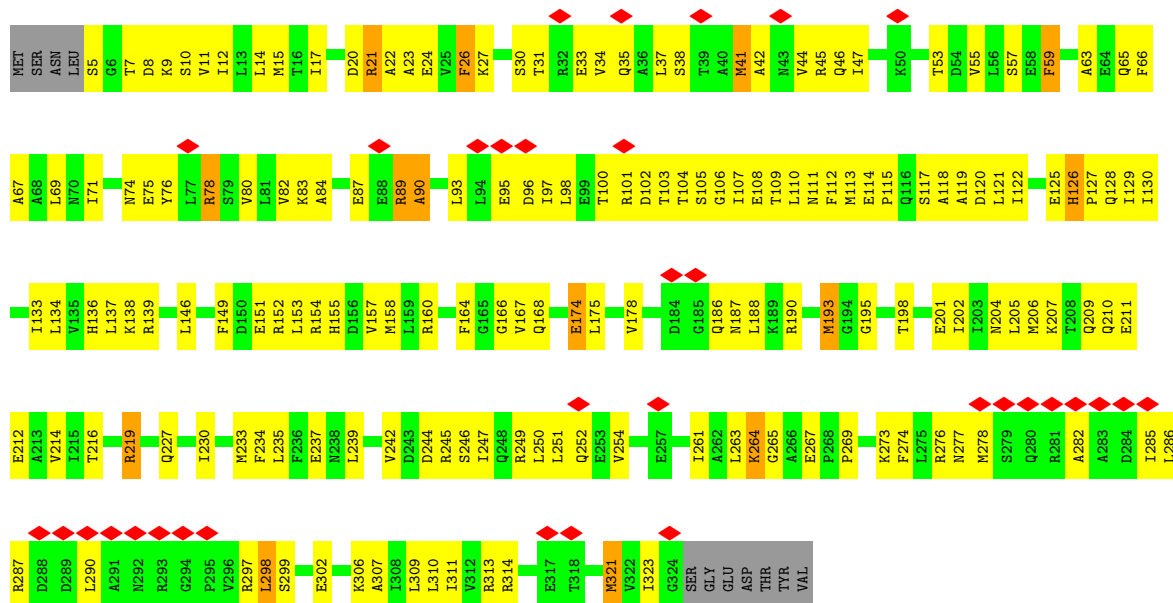


• Molecule 2: Flagellar motor switch protein FlhG



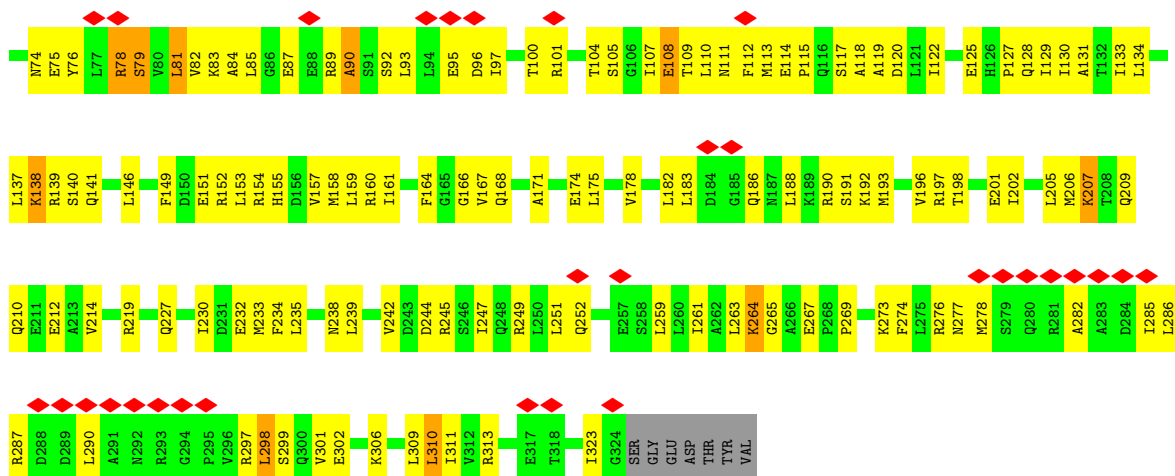


• Molecule 2: Flagellar motor switch protein Flg

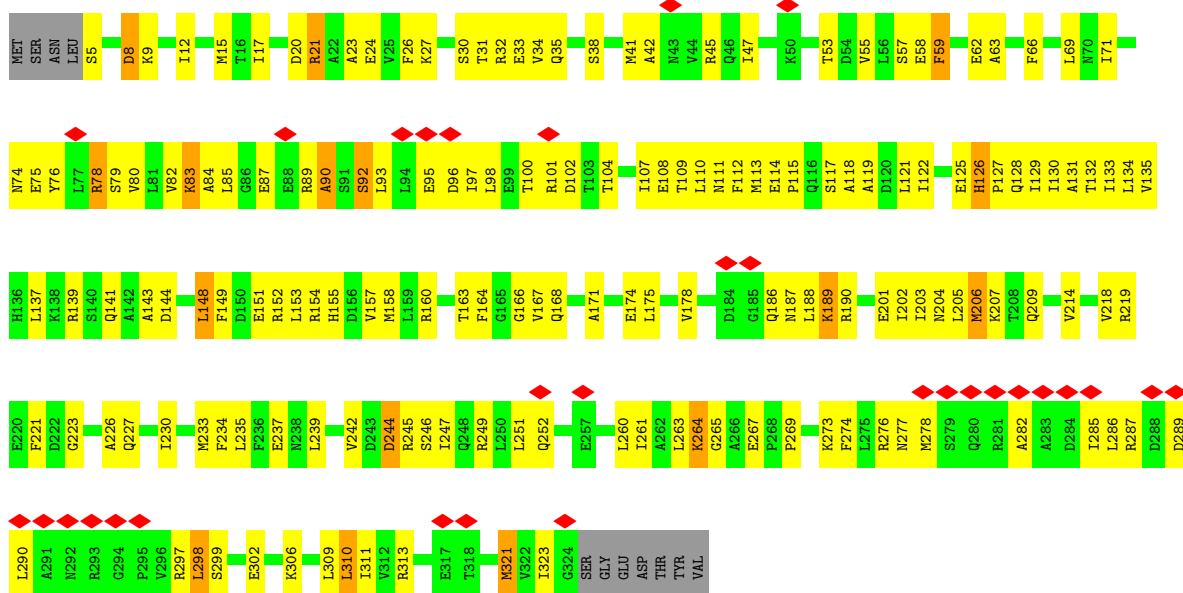


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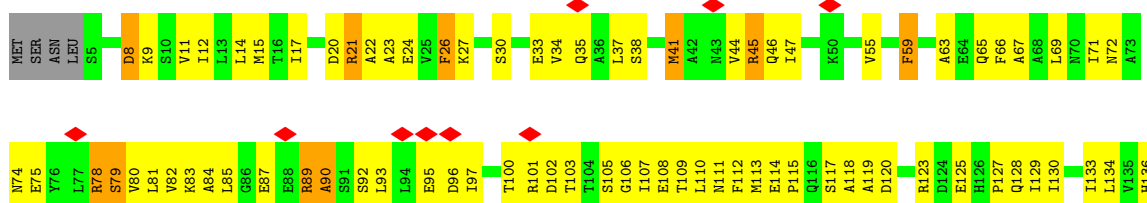
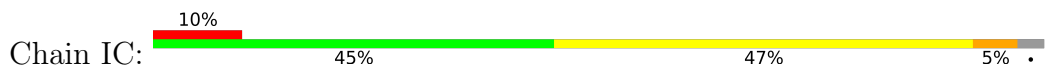


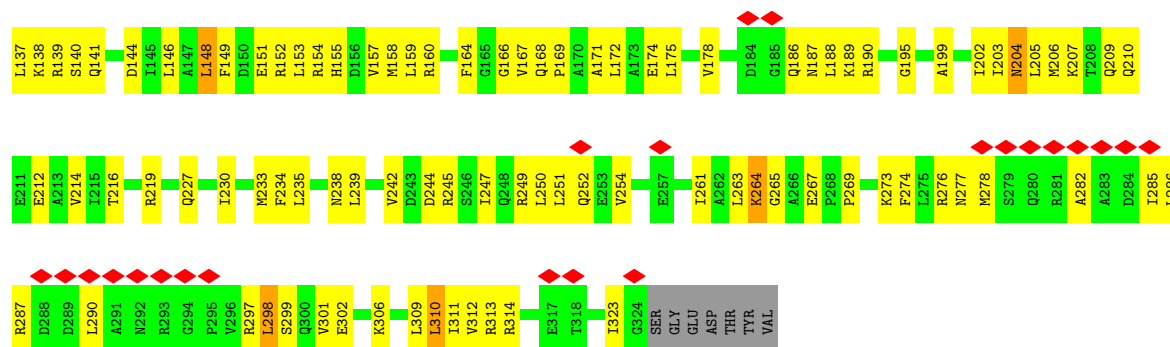


• Molecule 2: Flagellar motor switch protein FlhG

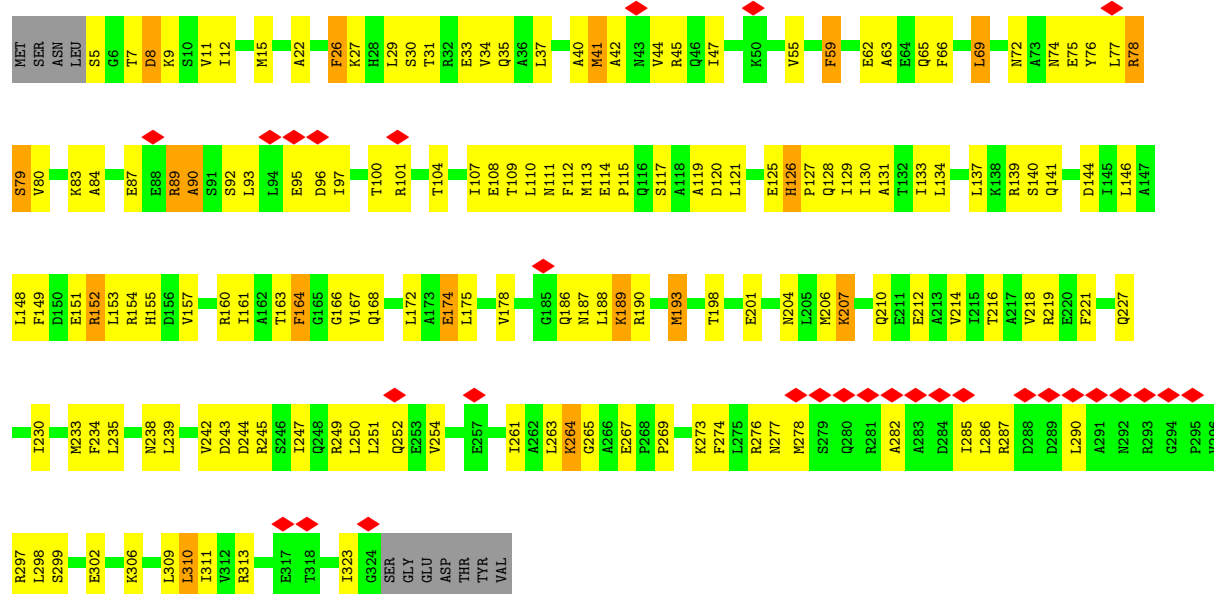


• Molecule 2: Flagellar motor switch protein FlhG

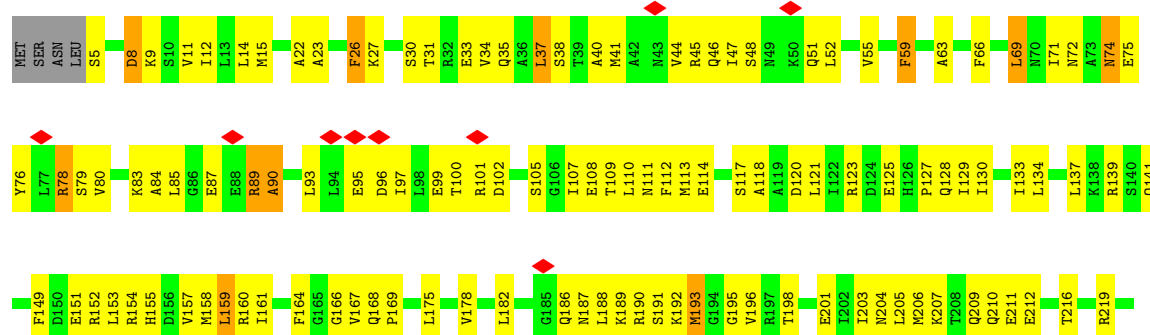


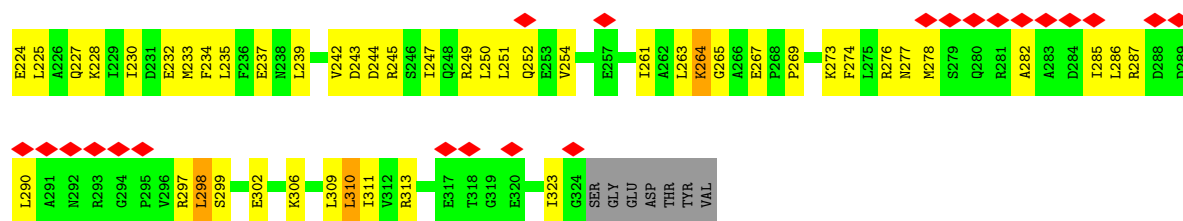


• Molecule 2: Flagellar motor switch protein FltG

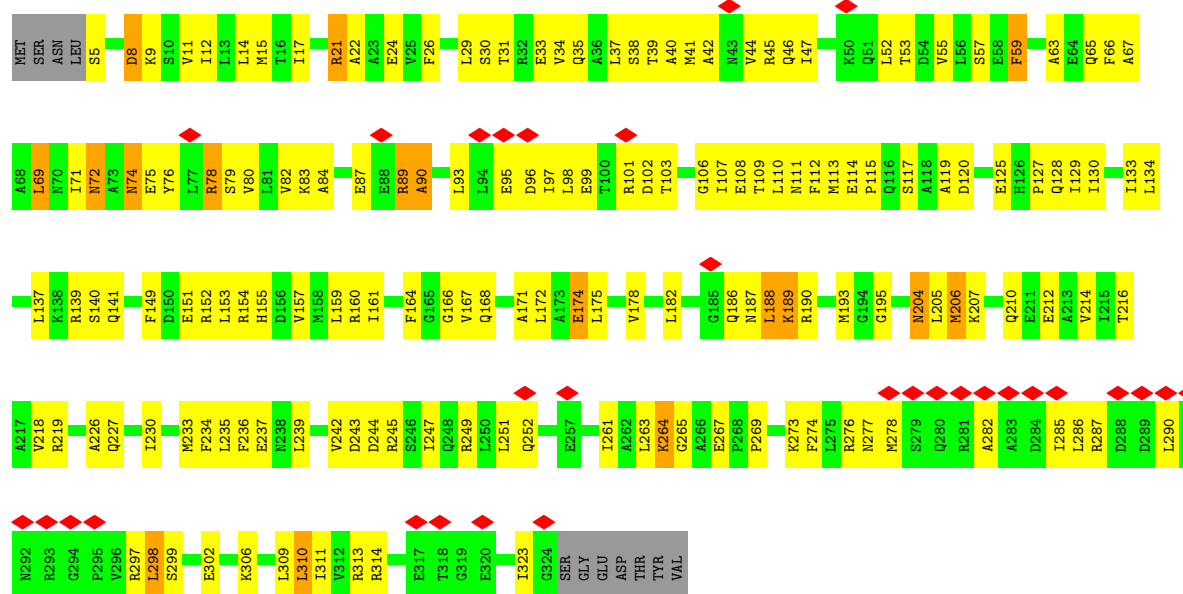


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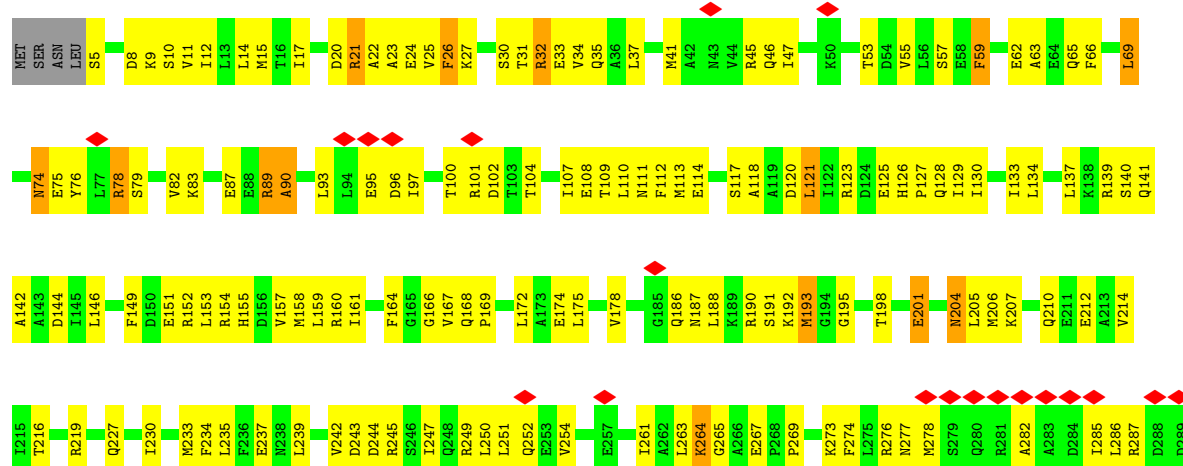




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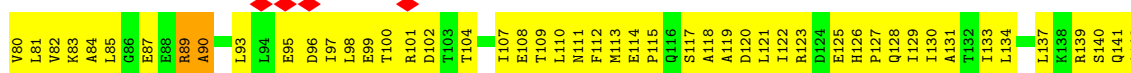
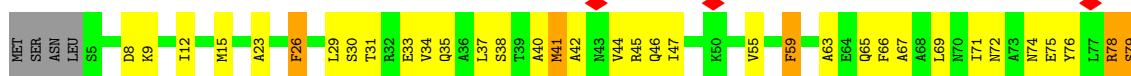


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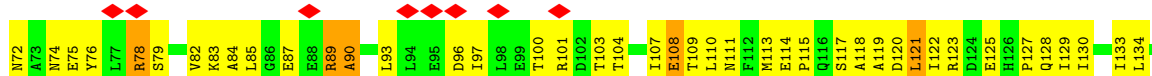
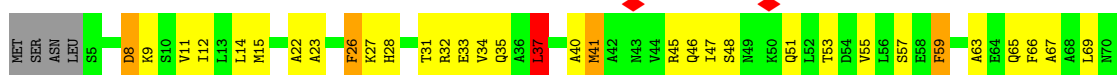




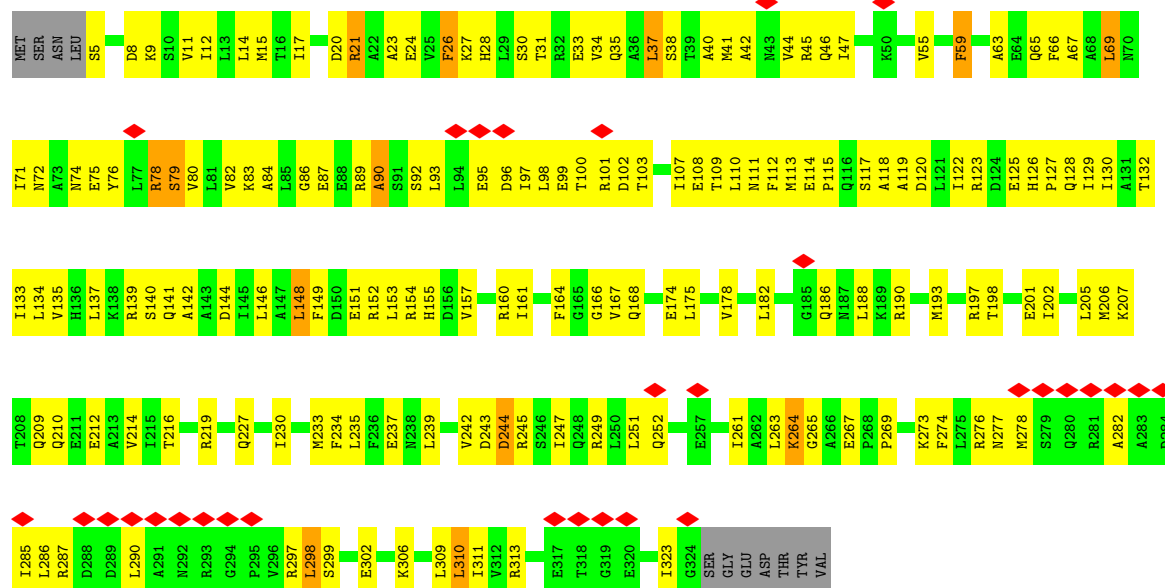
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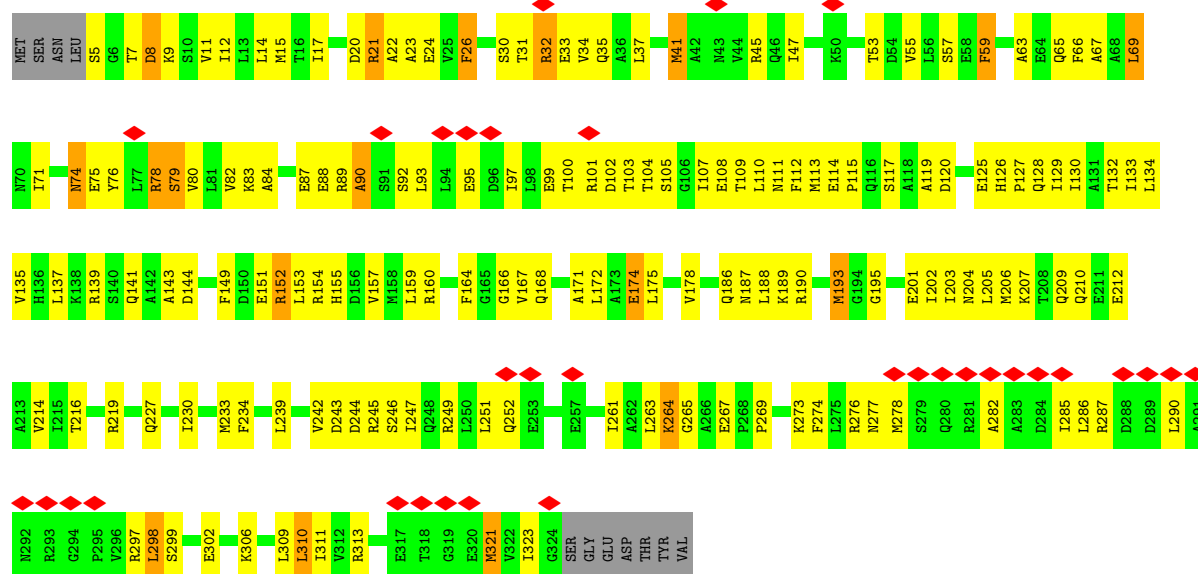
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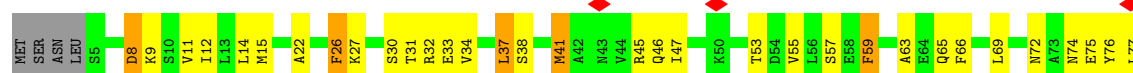
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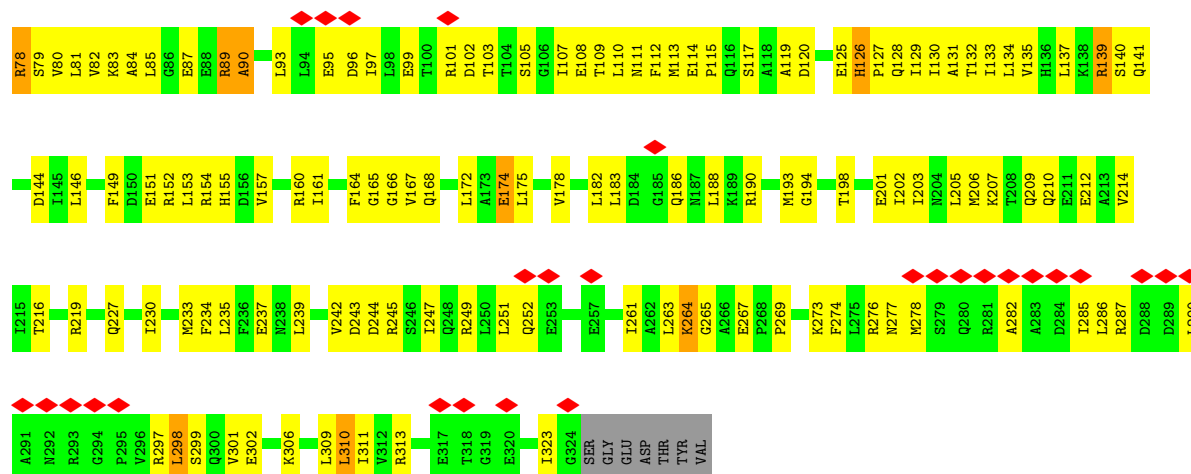


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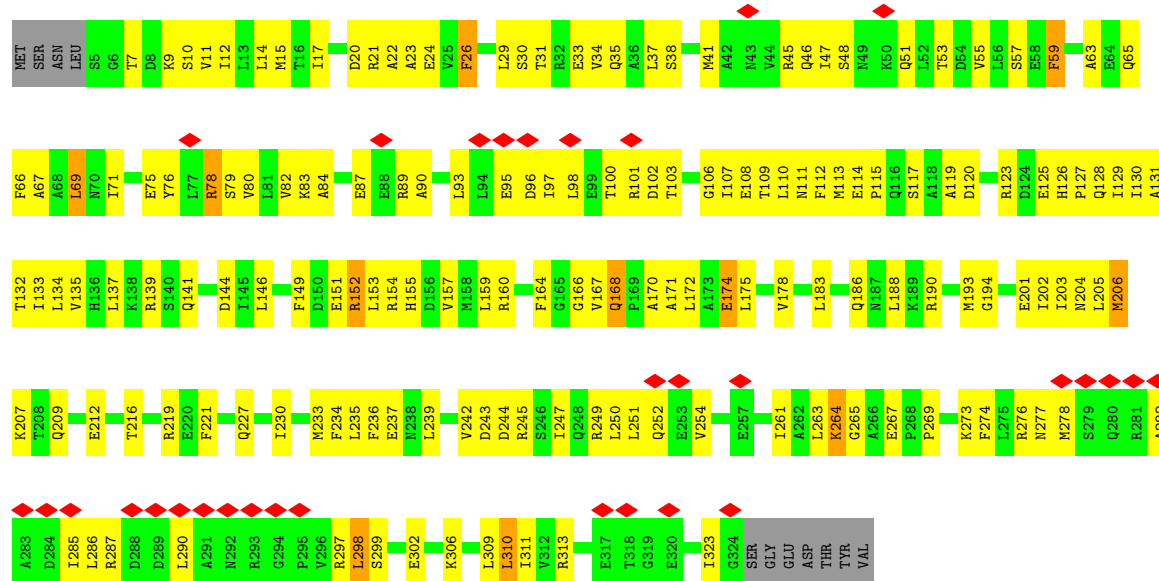


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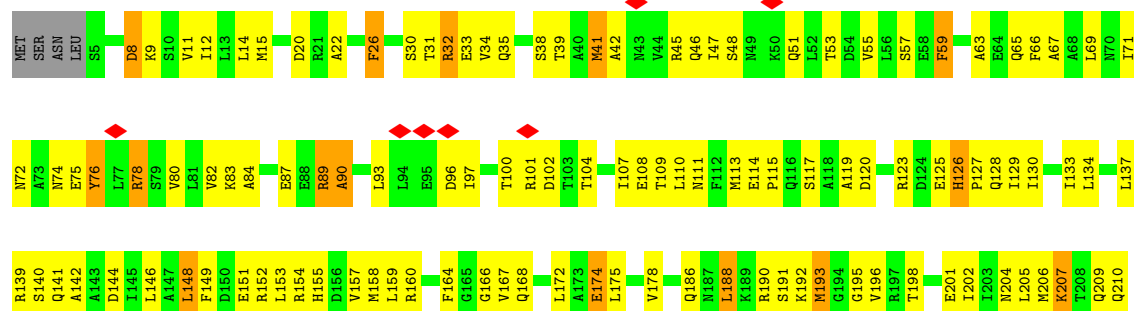


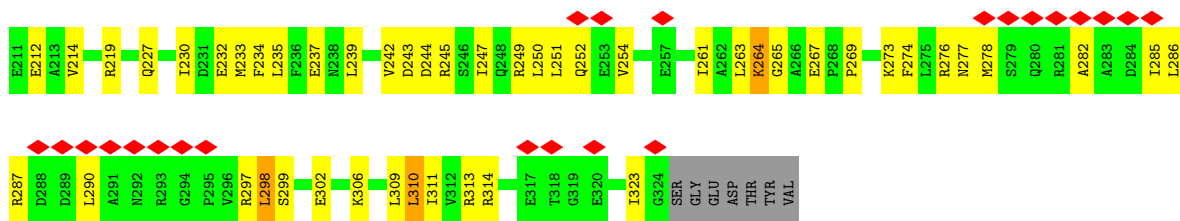


• Molecule 2: Flagellar motor switch protein FliG

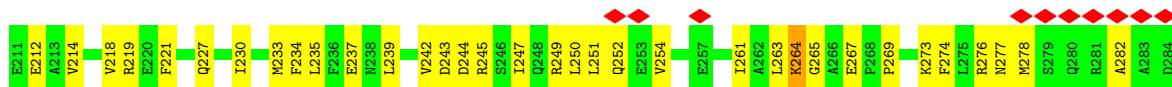
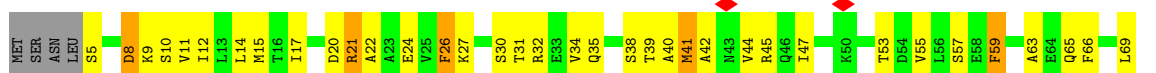


• Molecule 2: Flagellar motor switch protein FliG



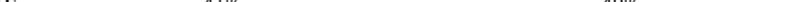


• Molecule 2: Flagellar motor switch protein FlhG

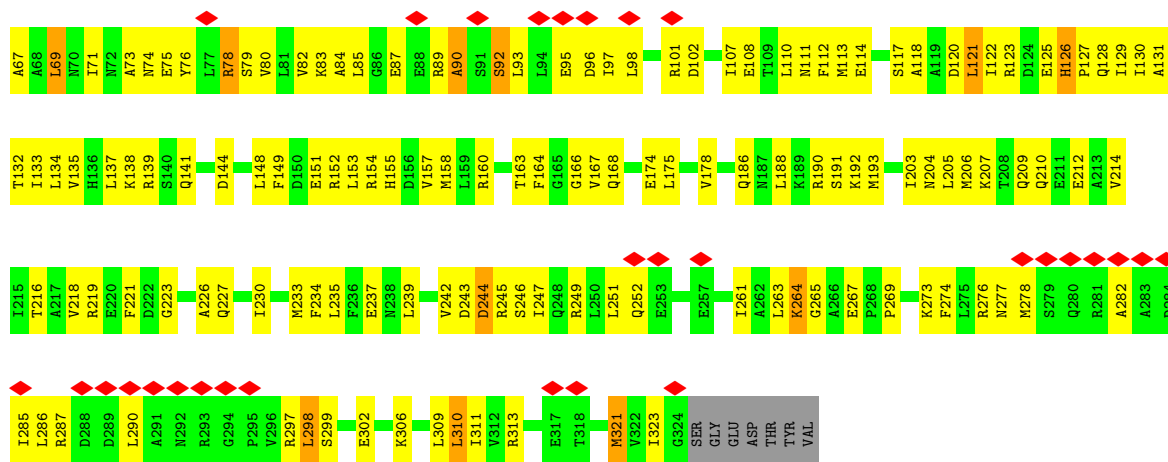


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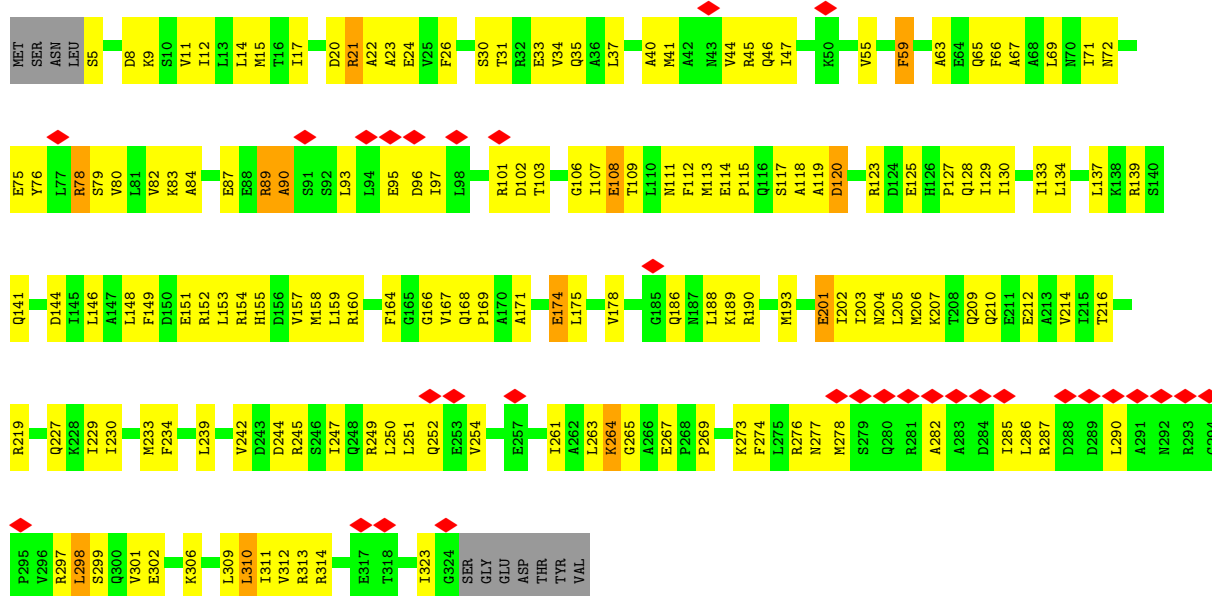


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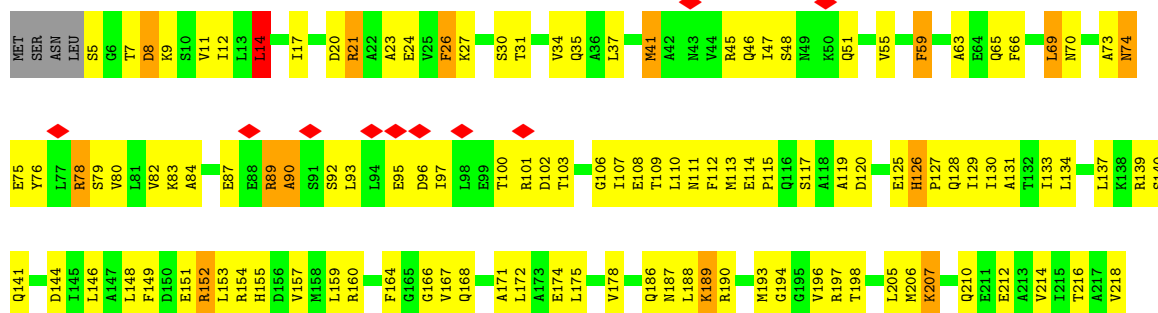


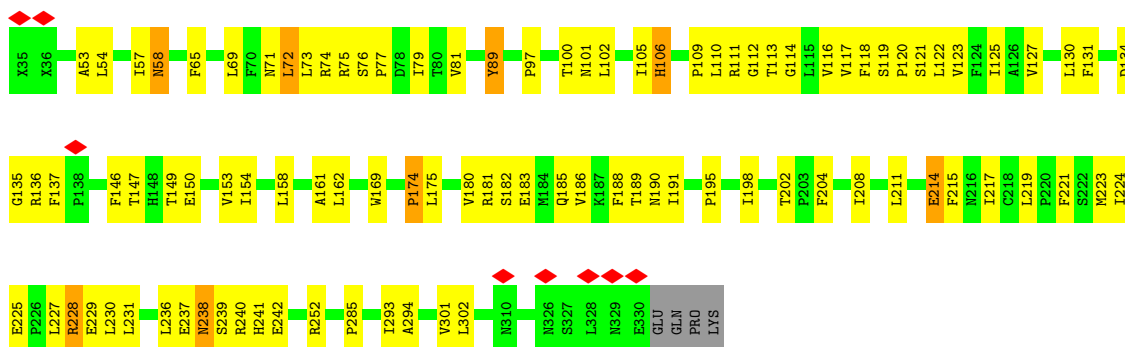


• Molecule 2: Flagellar motor switch protein FliG

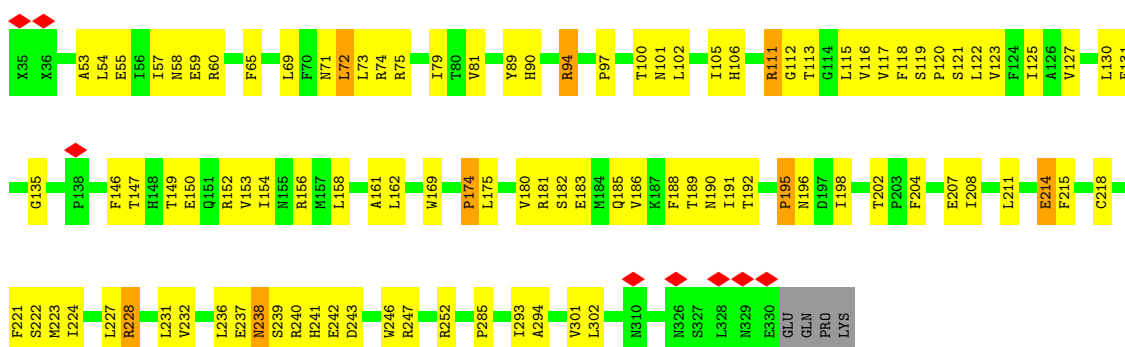


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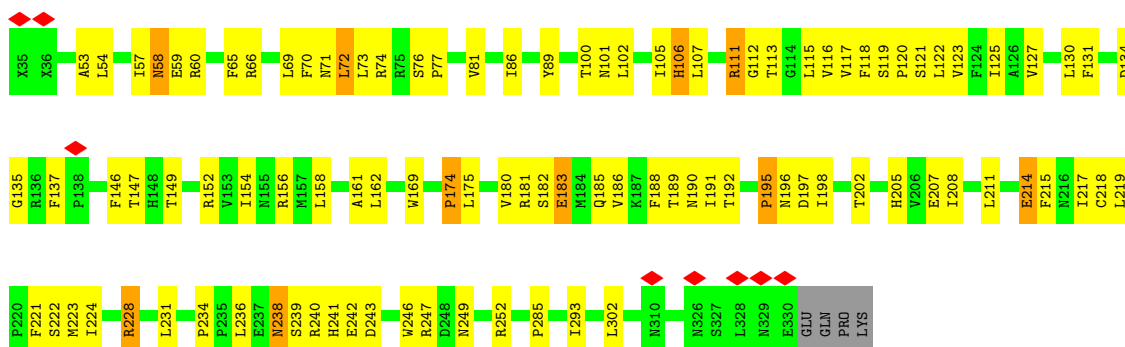




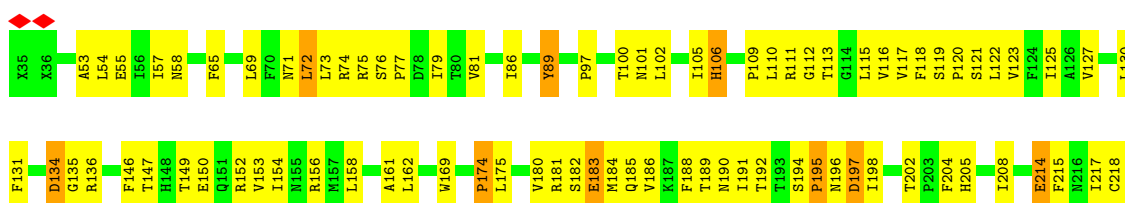
• Molecule 3: FlIM, Flagellar motor switch protein FlIM



• Molecule 3: FlIM, Flagellar motor switch protein FlIM

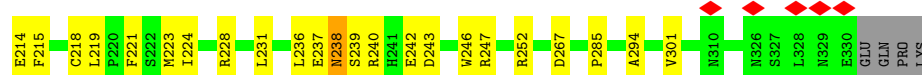


• Molecule 3: FlIM, Flagellar motor switch protein FlIM

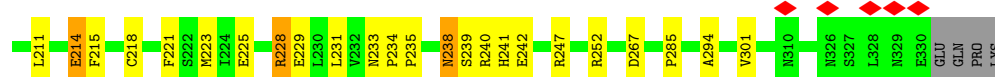
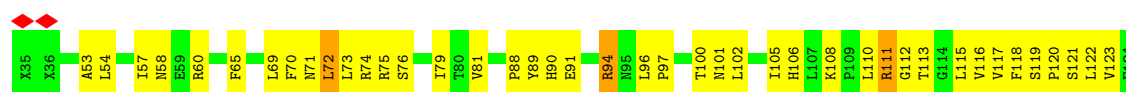




- Molecule 3: FlIM,Flagellar motor switch protein FlIM



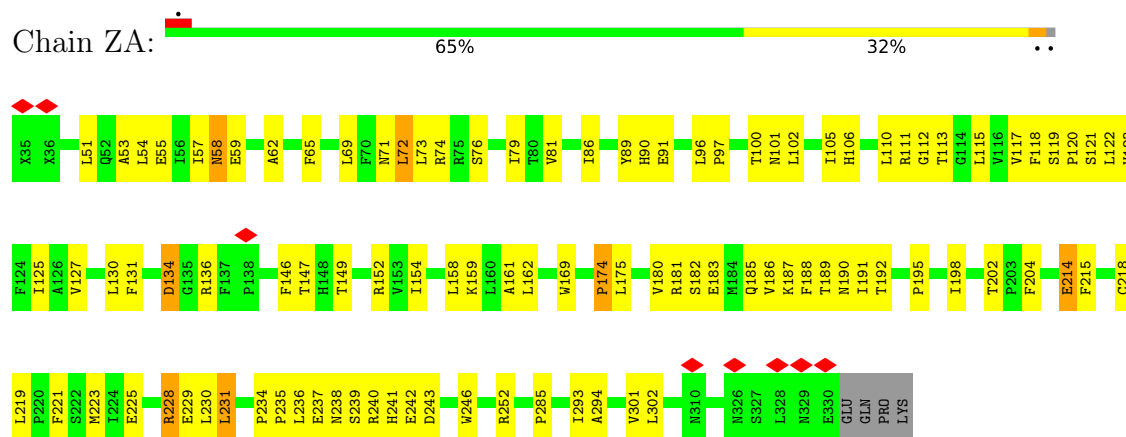
- Molecule 3: FlIM,Flagellar motor switch protein FlIM



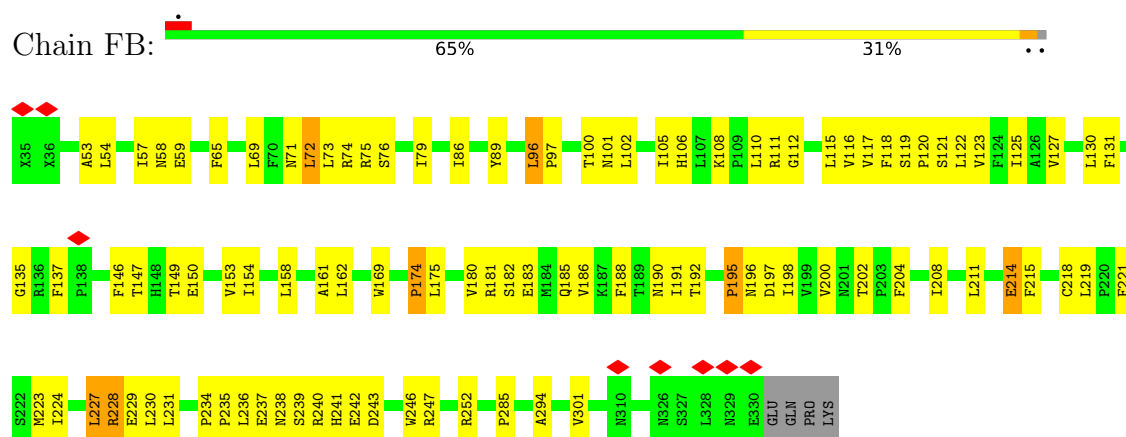
- Molecule 3: FlIM,Flagellar motor switch protein FlIM



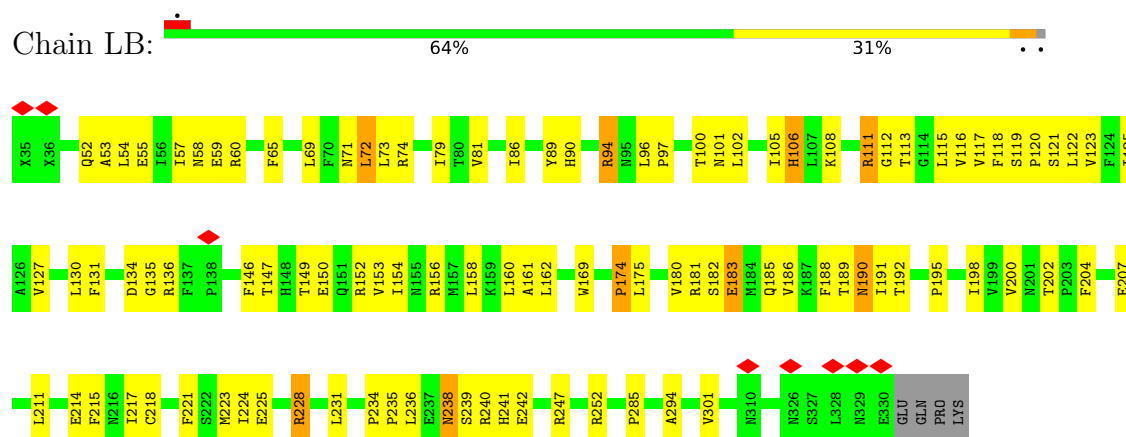
- Molecule 3: FlIM,Flagellar motor switch protein FlIM



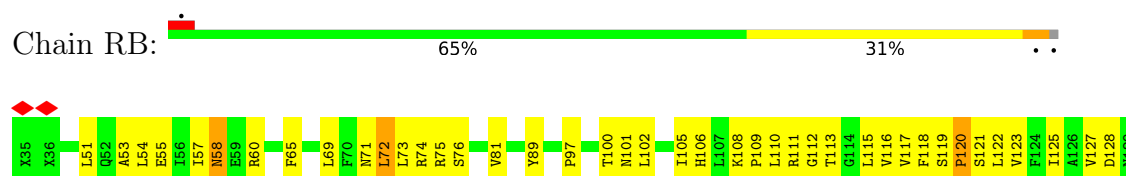
• Molecule 3: FlIM,Flagellar motor switch protein FlIM

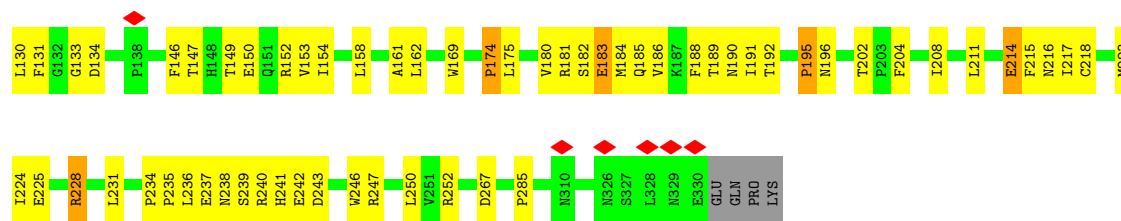


• Molecule 3: FlIM,Flagellar motor switch protein FlIM

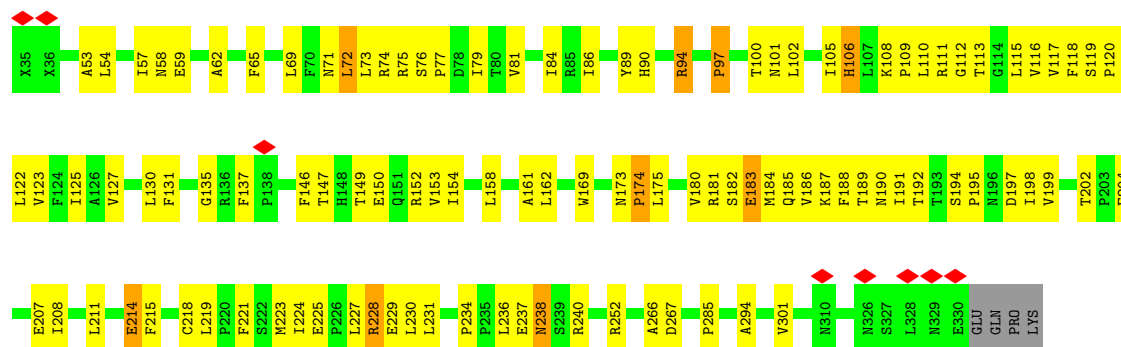


• Molecule 3: FlIM,Flagellar motor switch protein FlIM

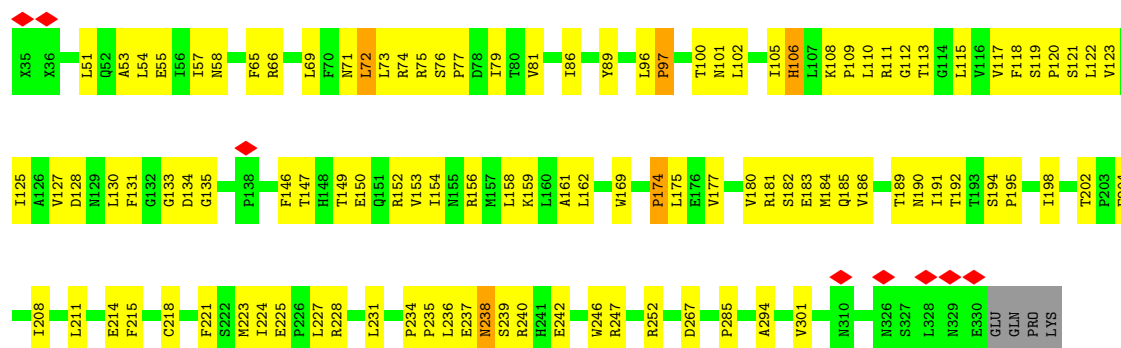




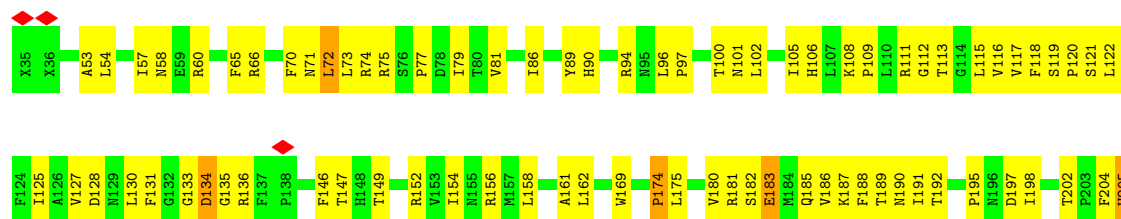
• Molecule 3: FlIM, Flagellar motor switch protein FlIM

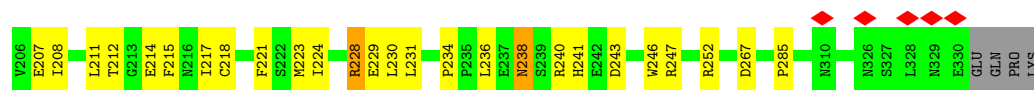


• Molecule 3: FlIM, Flagellar motor switch protein FlIM

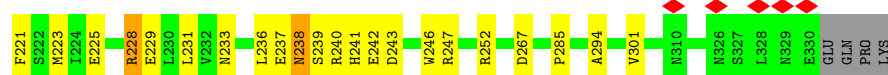
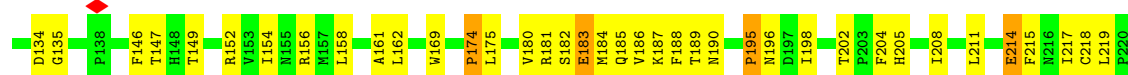


• Molecule 3: FlIM, Flagellar motor switch protein FlIM

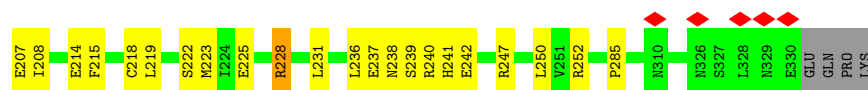




• Molecule 3: FliM, Flagellar motor switch protein FliM



• Molecule 3: FliM, Flagellar motor switch protein FliM

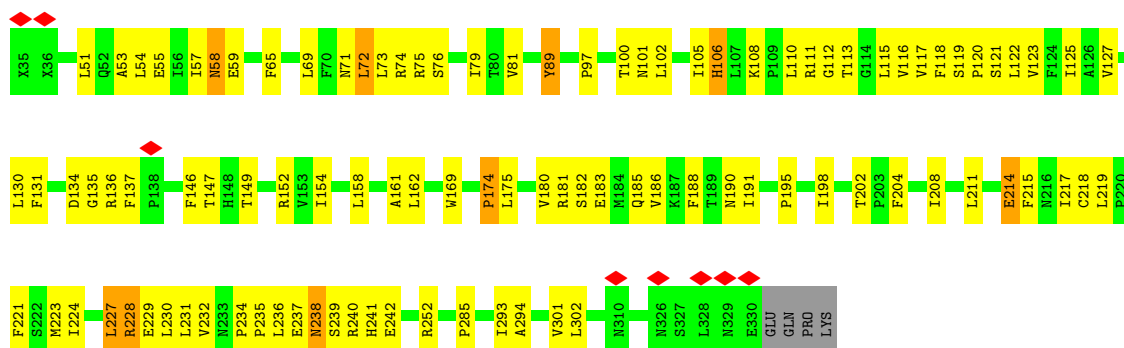


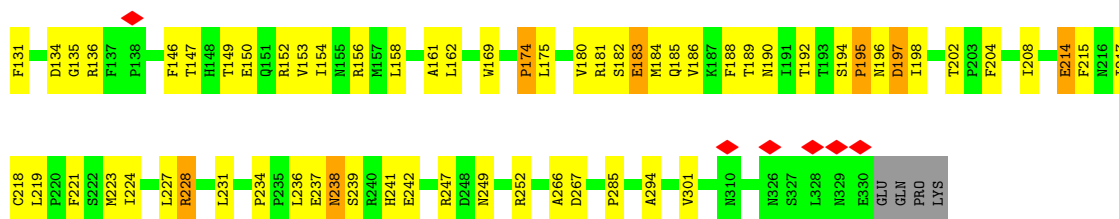
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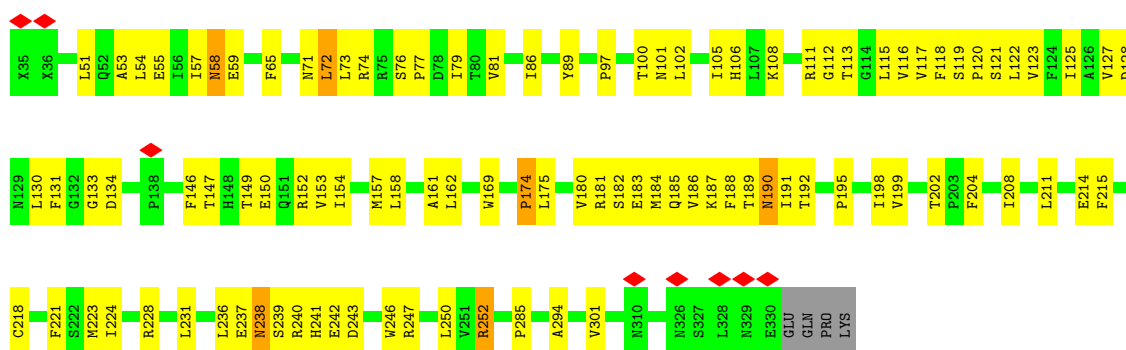
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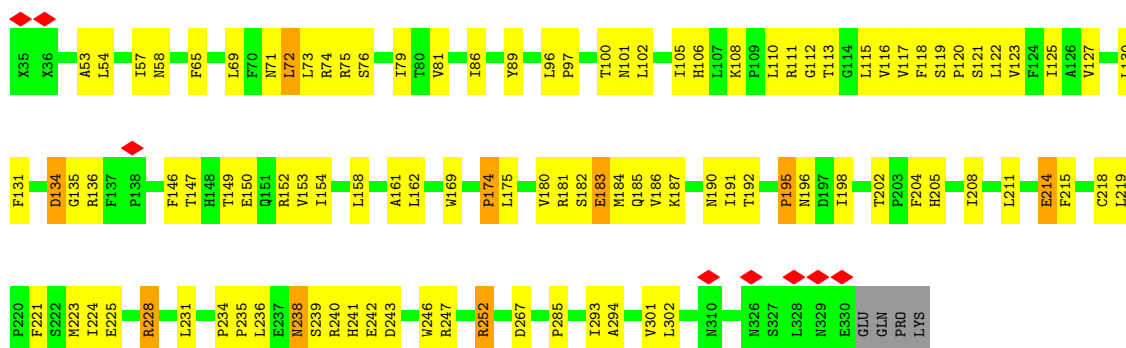




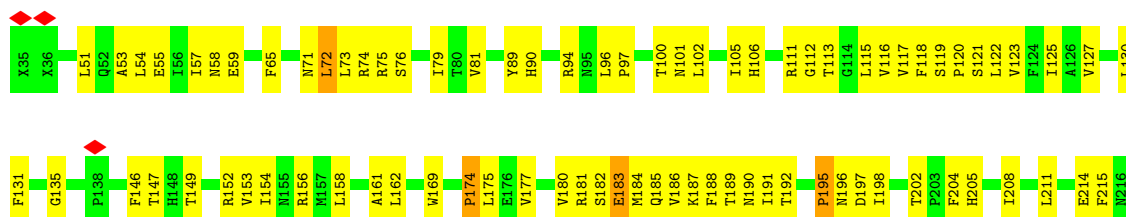
• Molecule 3: FlIM,Flagellar motor switch protein FlIM



• Molecule 3: FlIM,Flagellar motor switch protein FlIM

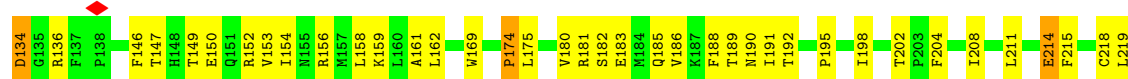


• Molecule 3: FlIM,Flagellar motor switch protein FlIM

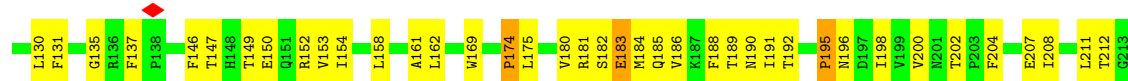
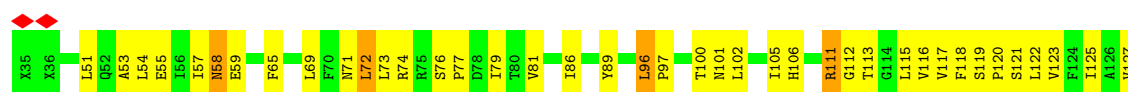




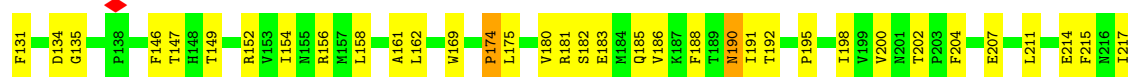
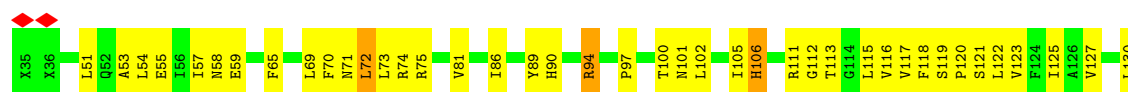
- Molecule 3: FlIM,Flagellar motor switch protein FlIM



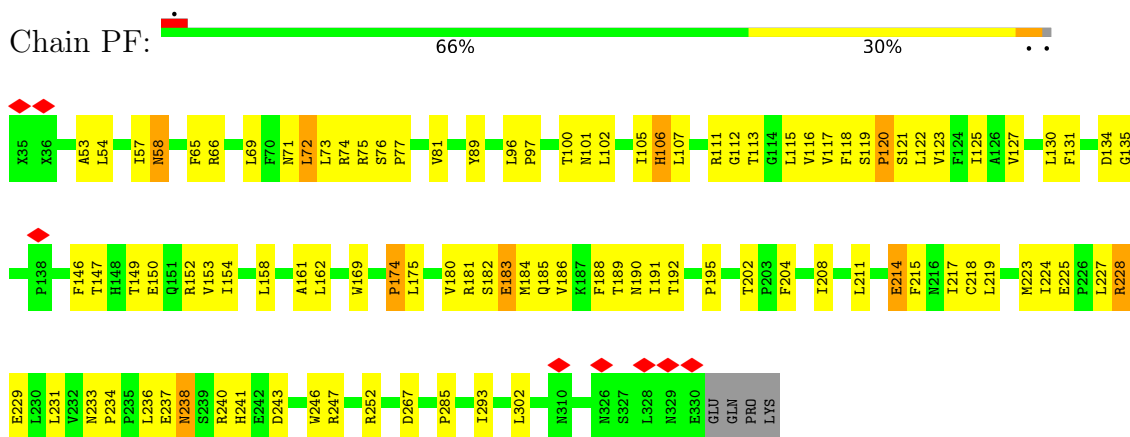
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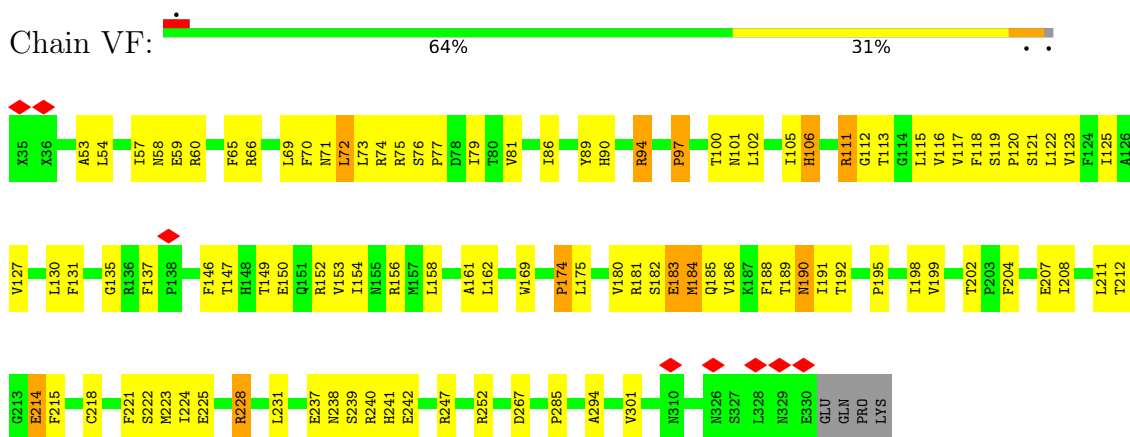
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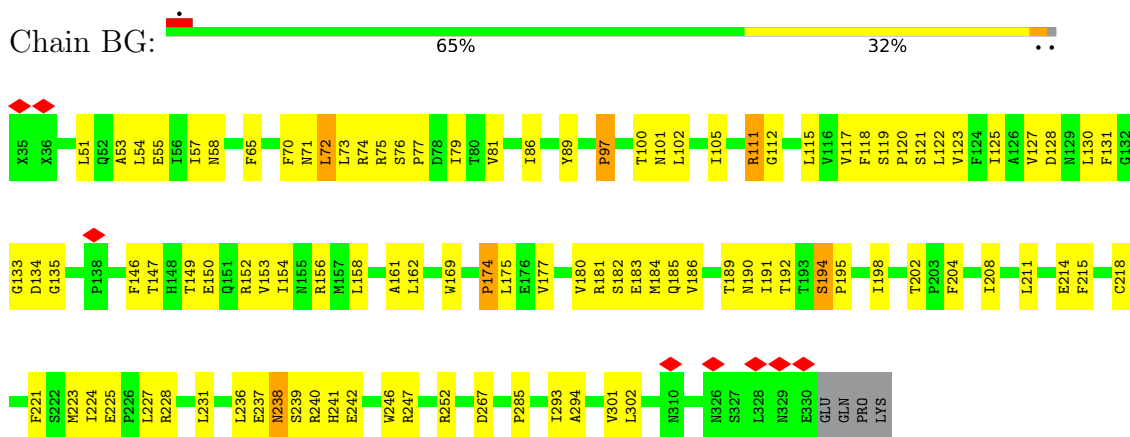
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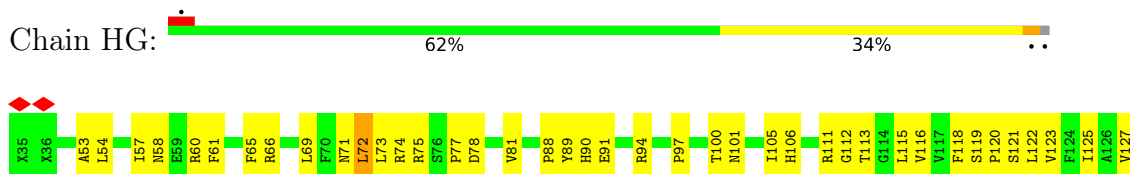
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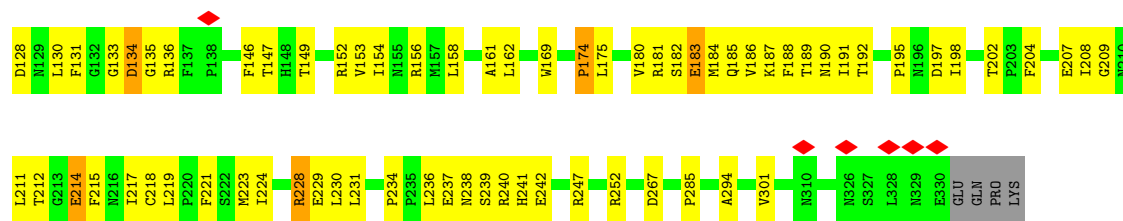


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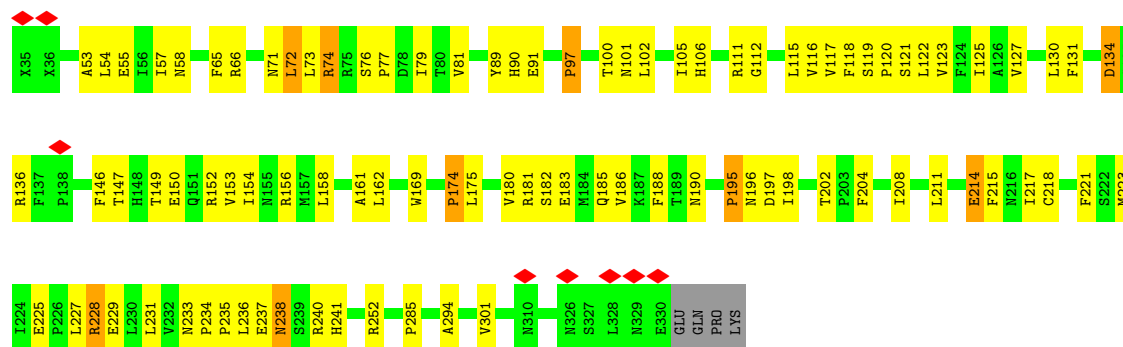


• Molecule 3: FlIM,Flagellar motor switch protein FlIM

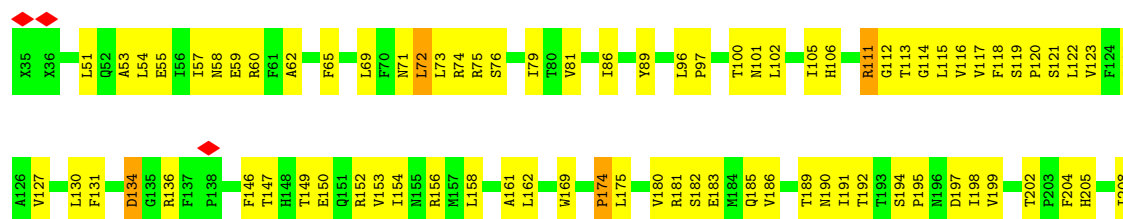




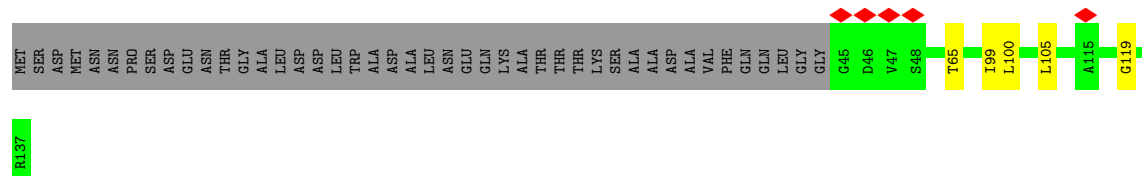
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• Molecule 3: FliM, Flagellar motor switch protein FliM



• Molecule 4: Flagellar motor switch protein FliN

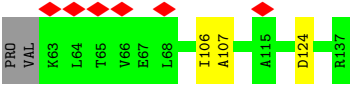
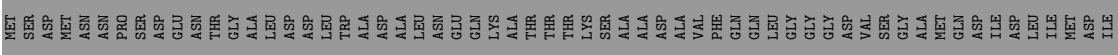


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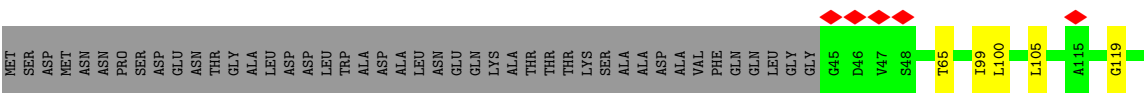




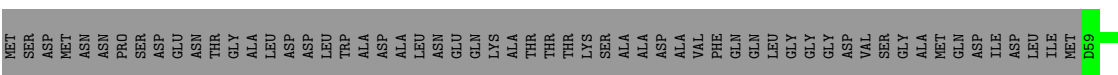
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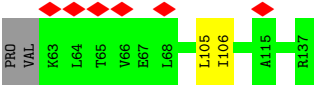
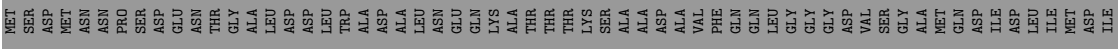
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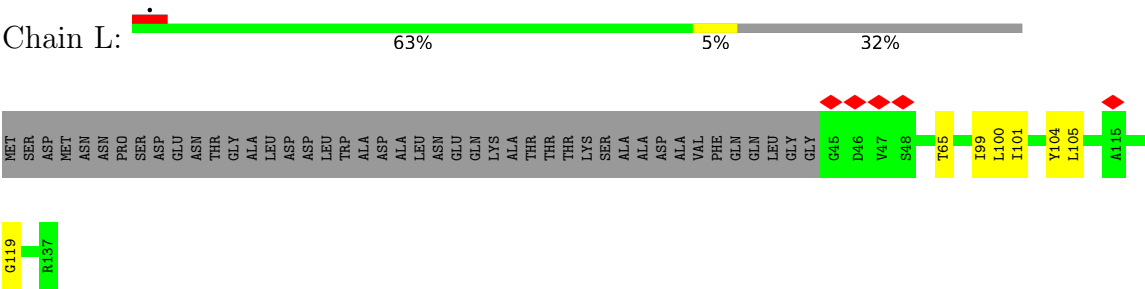
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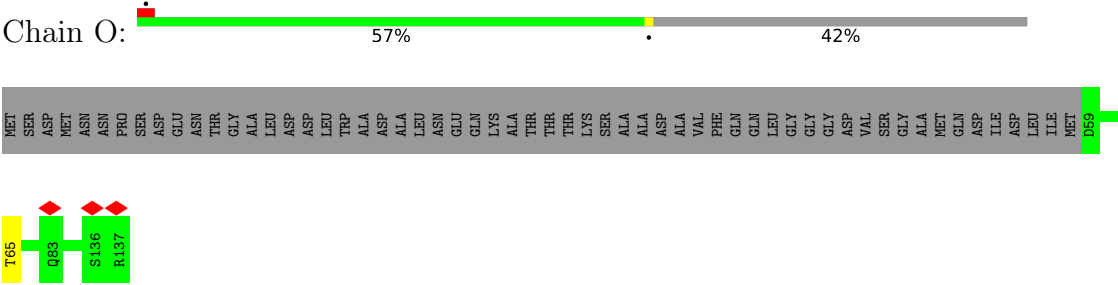
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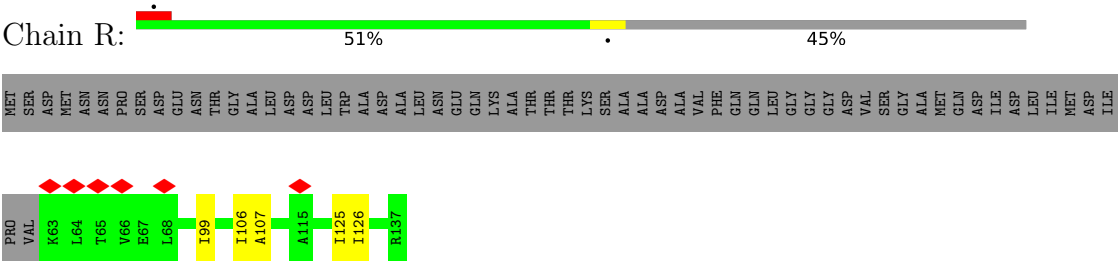
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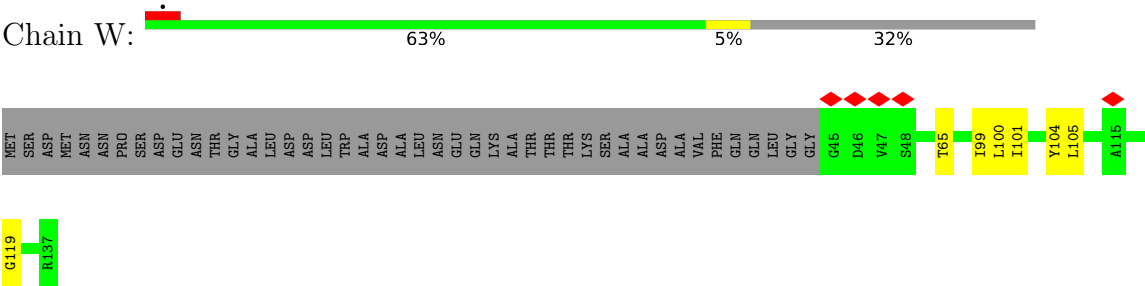
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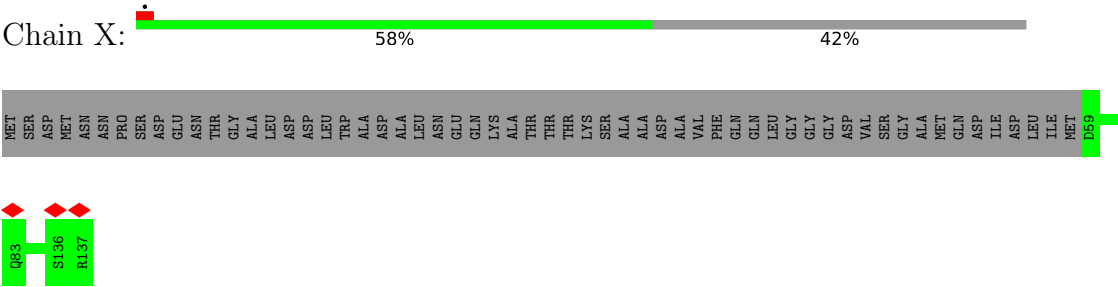
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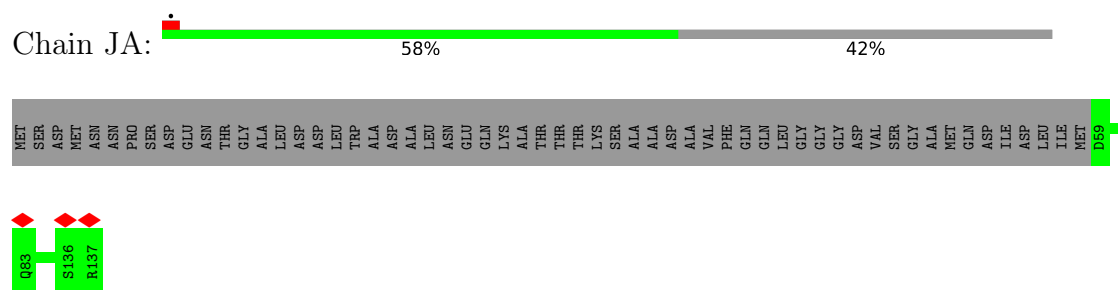
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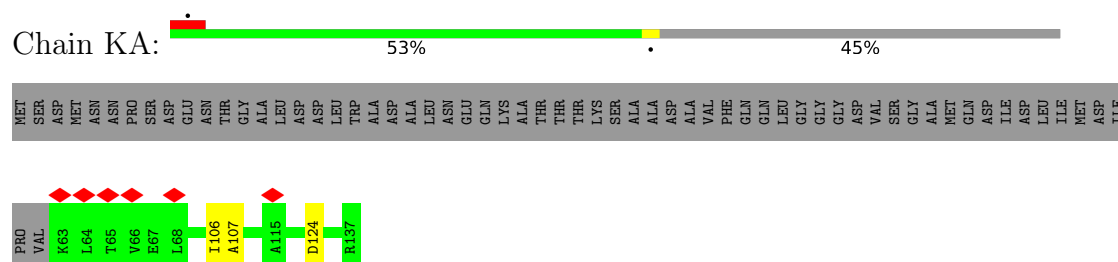
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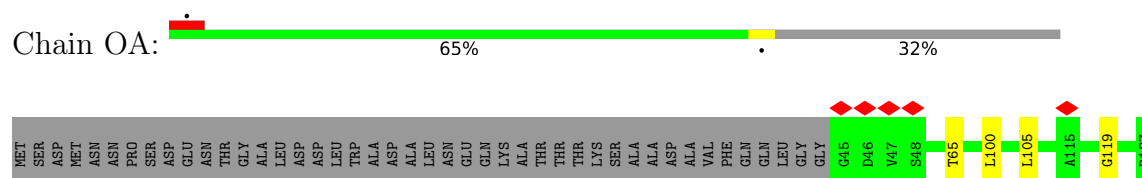
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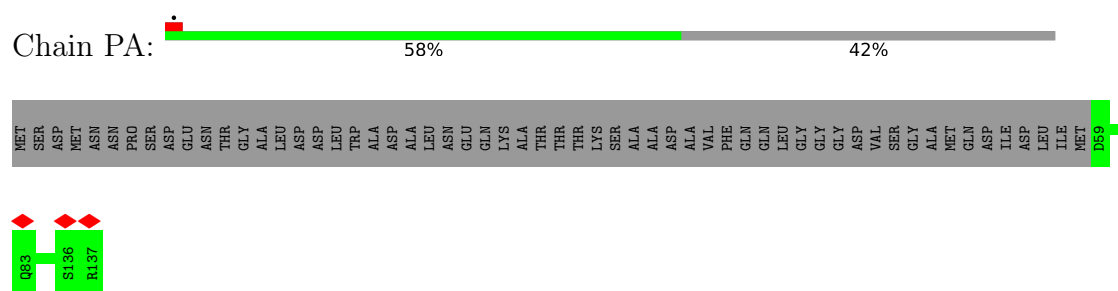
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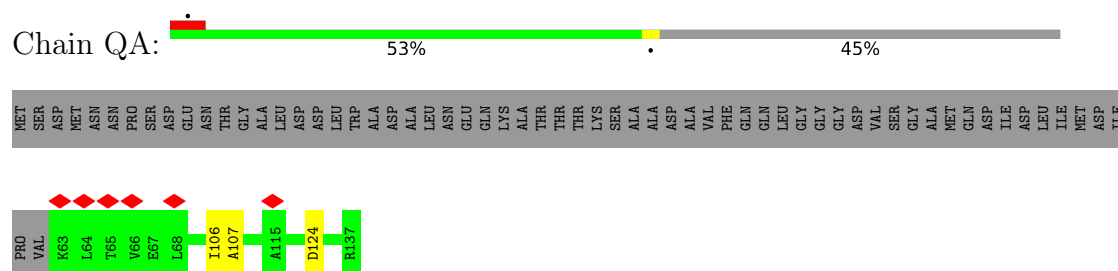
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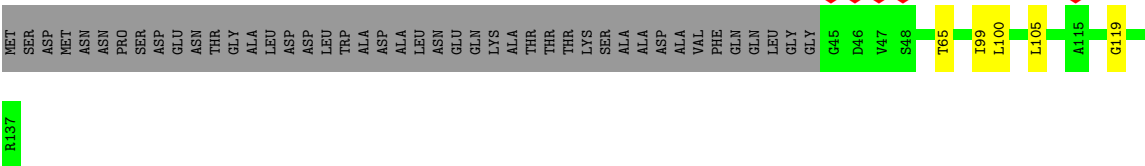
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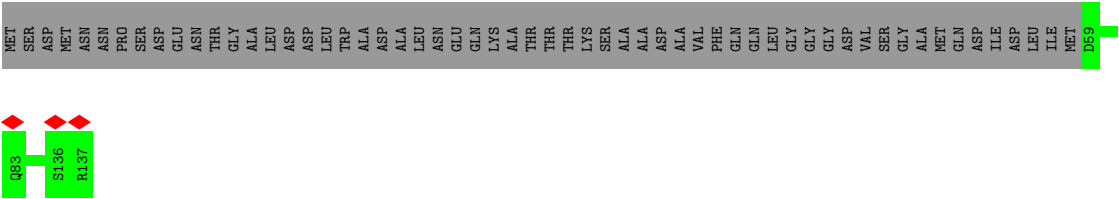
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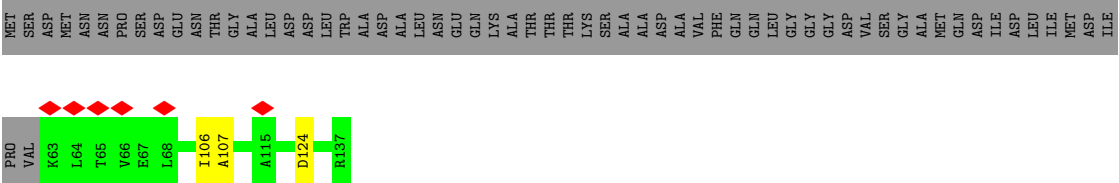
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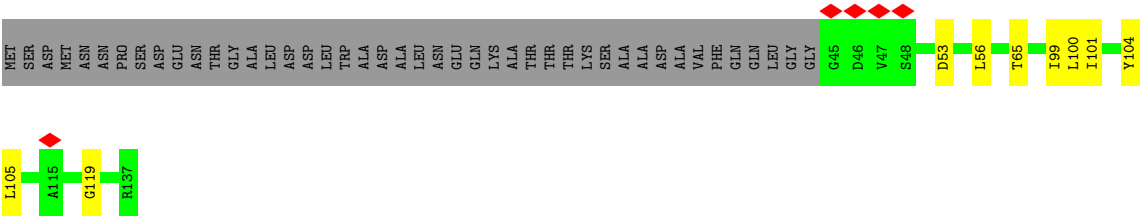
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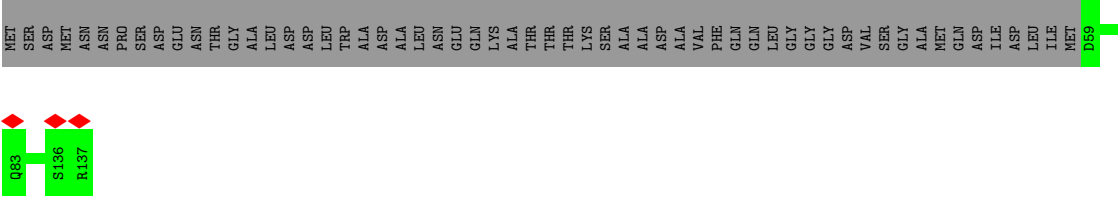
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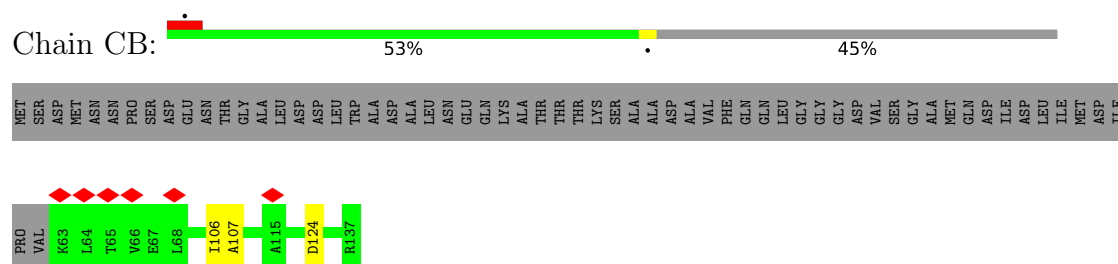
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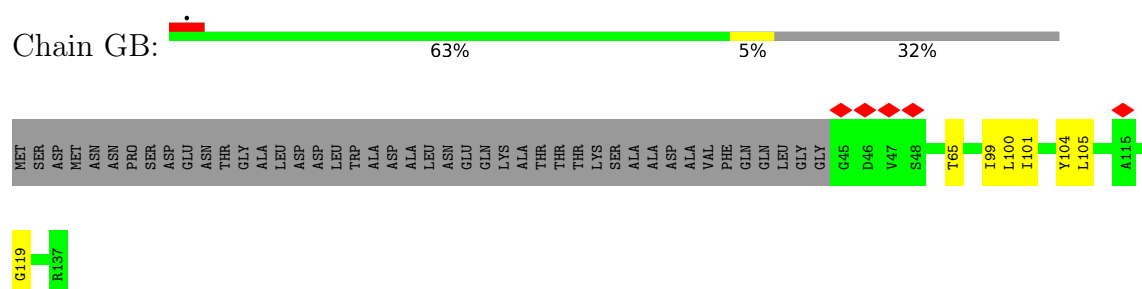
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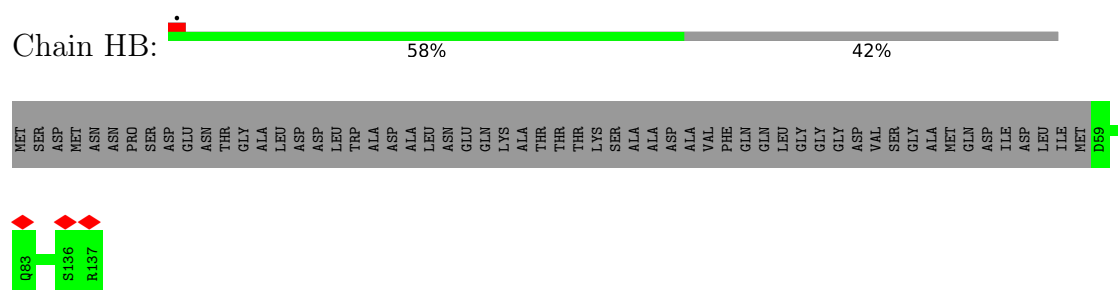
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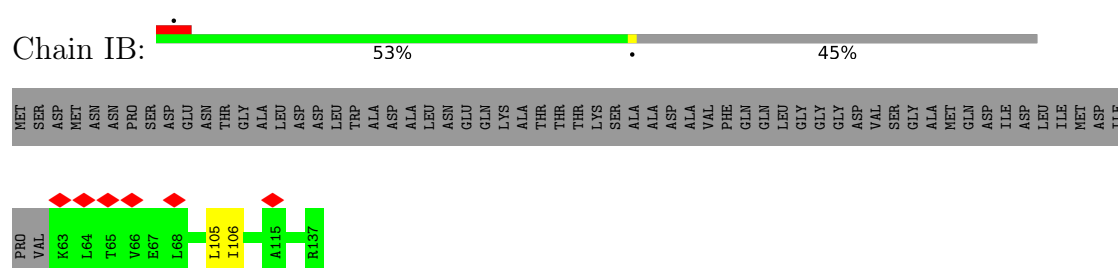
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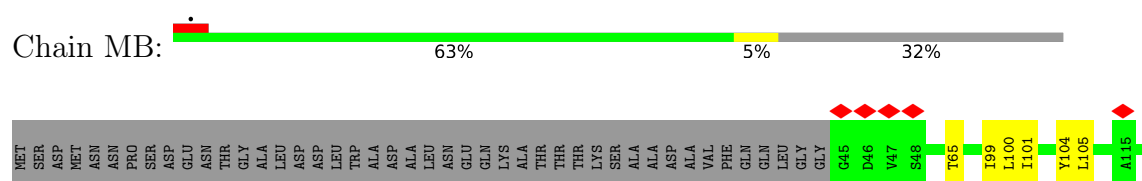
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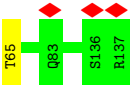
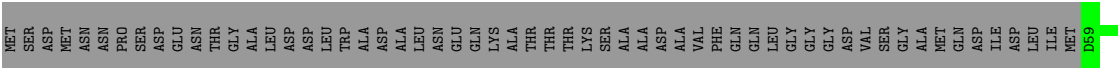


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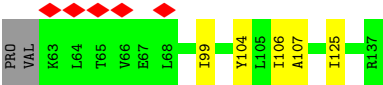
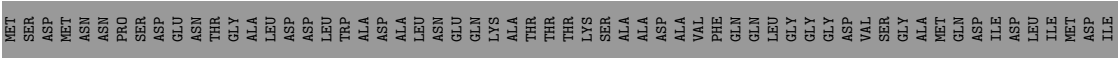




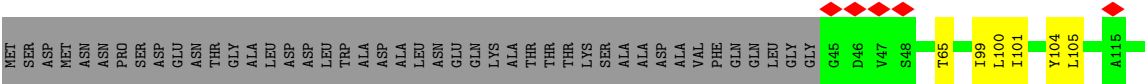
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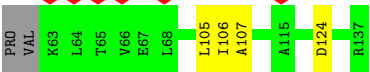
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MET	SER	ASP	MET	ASN	PRO	SER	ASP	GLU	ASN	THR	GLY	ALA	LEU	ASP	ASP	LEU	TRP	ALA	ASP	ALA	LEU	ASN	GLU	GLN	LYS	THR	THR	THR	LYS	SER	ALA	ALA	ASP	ALA	VAL	PHE	GLN	GLN	LEU	GLY	GLY	GLY	ASP	VAL	SER	GLY	ALA	MET	GLN	ILE	ASP	LEU	ILE	MET	ASP	ASP	ILE
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• Molecule 4: Flagellar motor switch protein FliN



MET	SER	ASP	MET	ASN	PRO	SER	ASP	GLU	ASN	THR	GLY	ALA	LEU	ASP	ASP	LEU	TRP	ALA	ASP	ALA	LEU	ASN	GLU	GLN	LYS	THR	THR	THR	LYS	SER	ALA	ALA	ASP	ALA	VAL	PHE	GLN	GLN	LEU	GLY	GLY	G45	D46	V47	S48	T65	V66	I99	L100	I101	Y104	L105	A115
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• Molecule 4: Flagellar motor switch protein FliN



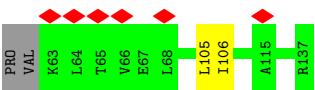
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• Molecule 4: Flagellar motor switch protein FliN



MET	SER	ASP	MET	ASN	PRO	SER	ASP	GLU	ASN	THR	GLY	ALA	LEU	ASP	ASP	LEU	TRP	ALA	ASP	ALA	LEU	ASN	GLU	GLN	LYS	THR	THR	THR	LYS	SER	ALA	ALA	ASP	ALA	VAL	PHE	GLN	GLN	LEU	GLY	GLY	GLY	ASP	VAL	SER	GLY	ALA	MET	GLN	ILE	ASP	LEU	ILE	MET	ASP	ASP	ILE
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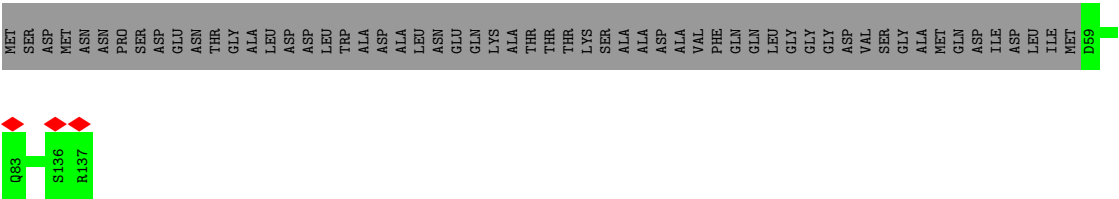
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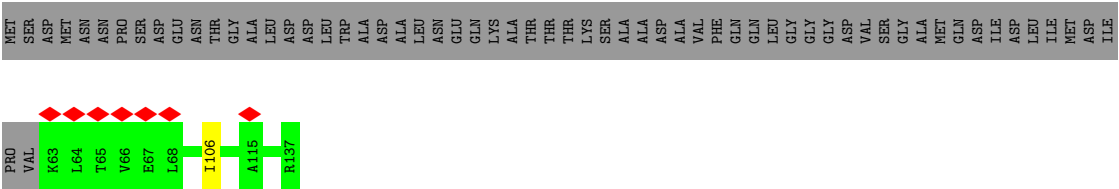
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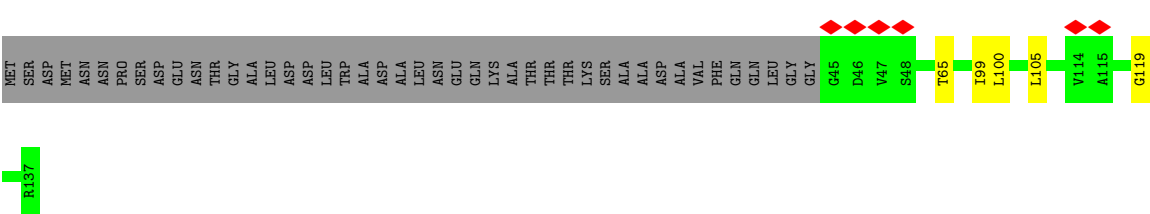
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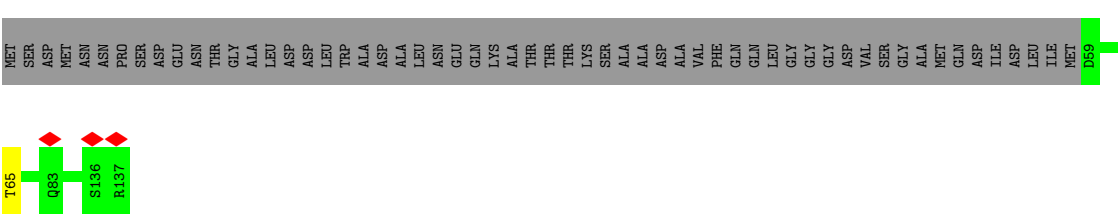
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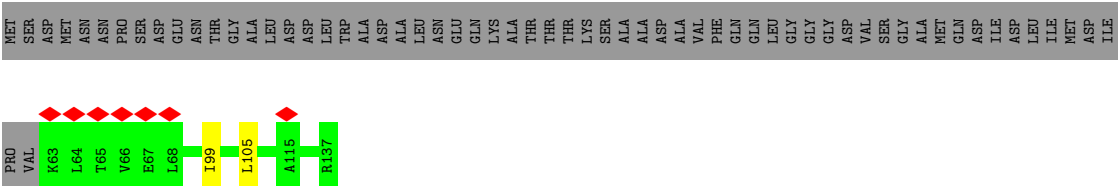
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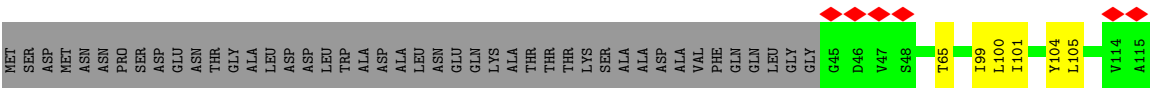
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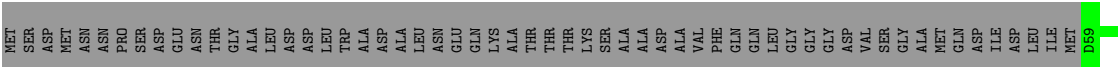
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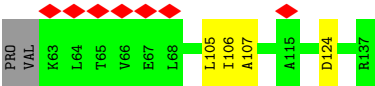
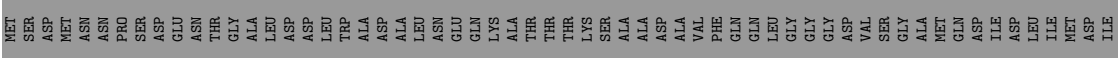
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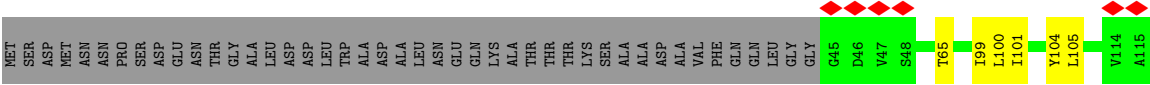
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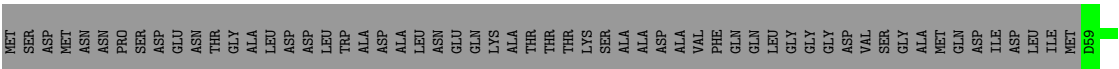
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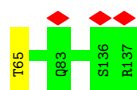


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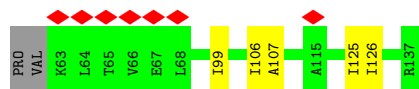
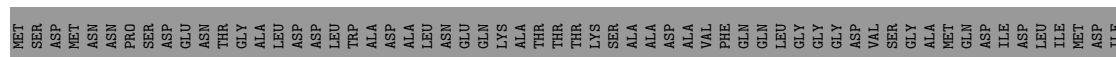


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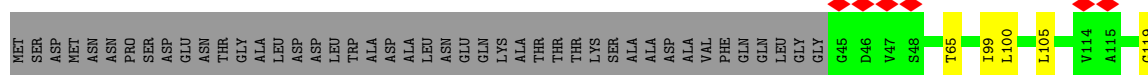




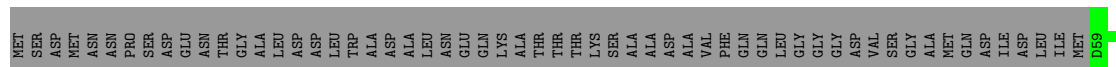
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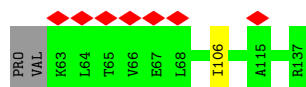
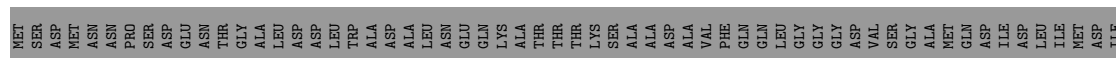
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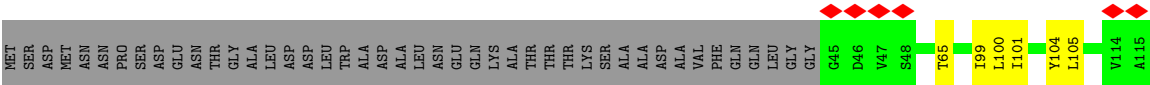


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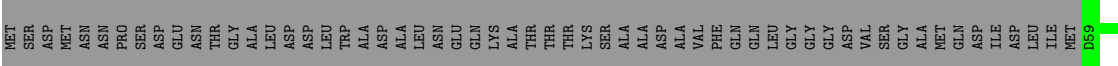


- Molecule 4: Flagellar motor switch protein FliN

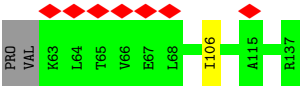
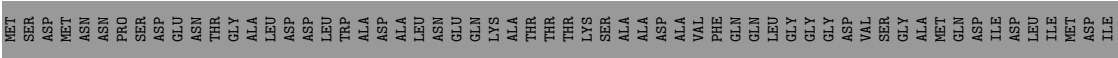




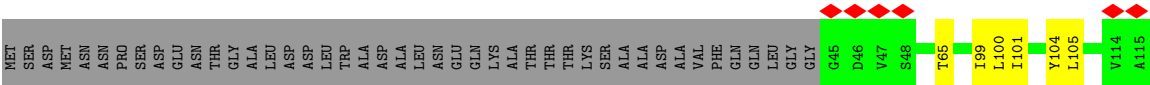
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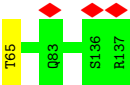
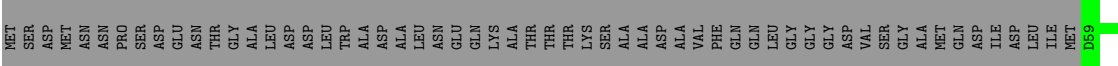
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• Molecule 4: Flagellar motor switch protein FliN



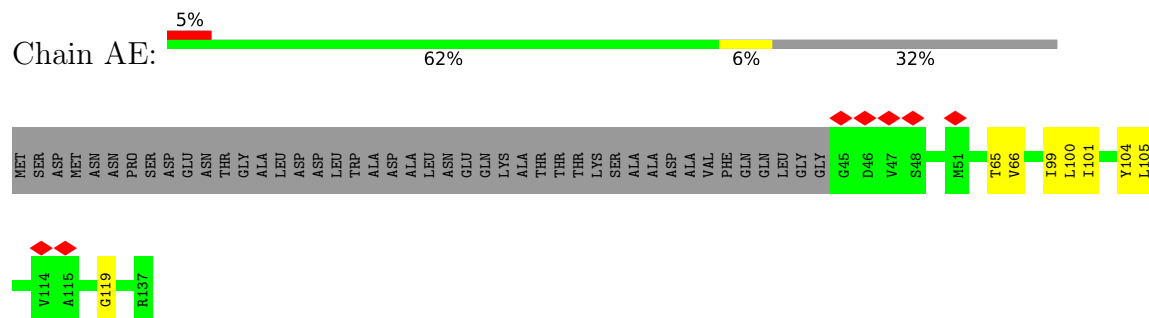
• Molecule 4: Flagellar motor switch protein FliN

- Molecule 4: Flagellar motor switch protein FliN

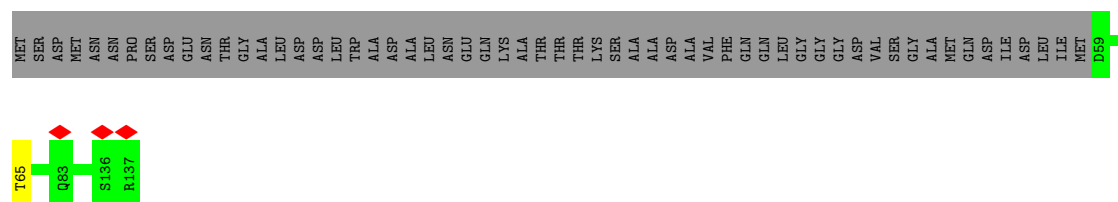
- Molecule 4: Flagellar motor switch protein FliN

- Molecule 4: Flagellar motor switch protein FliN

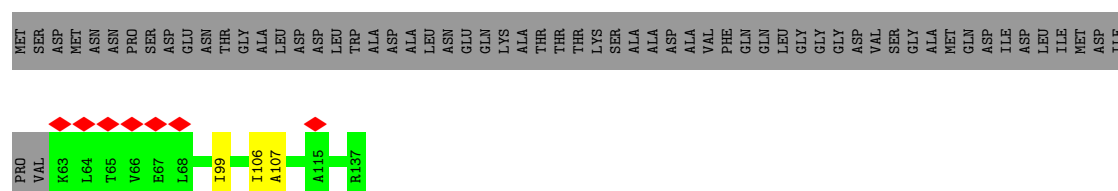
- Molecule 4: Flagellar motor switch protein FliN



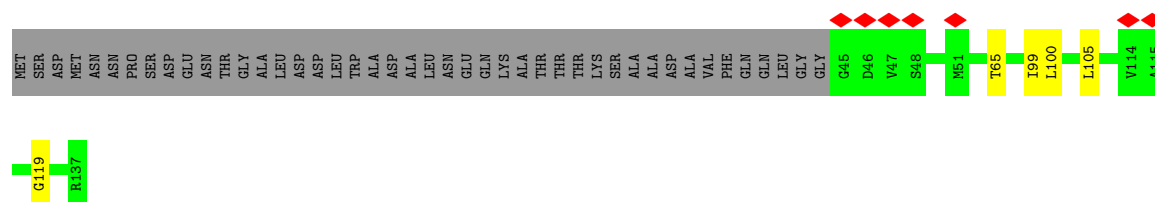
• Molecule 4: Flagellar motor switch protein FliN

Chain BE:  57% 42%

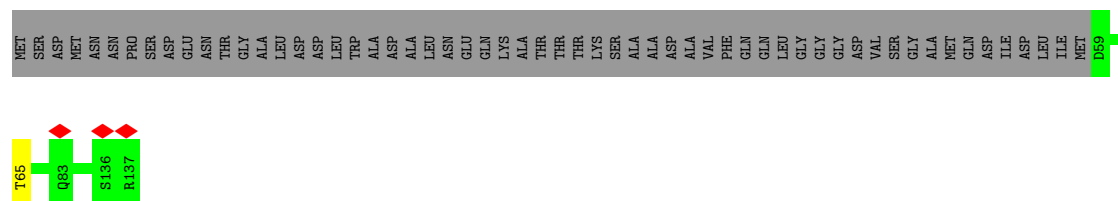
• Molecule 4: Flagellar motor switch protein FliN

Chain CE:  5% 53% 45%

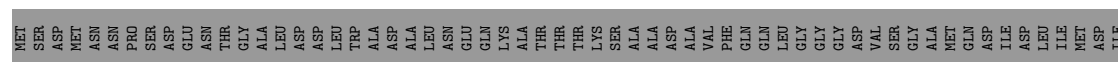
• Molecule 4: Flagellar motor switch protein FliN

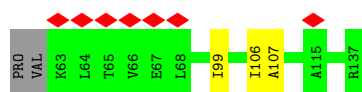
Chain GE:  5% 64% 32%

• Molecule 4: Flagellar motor switch protein FliN

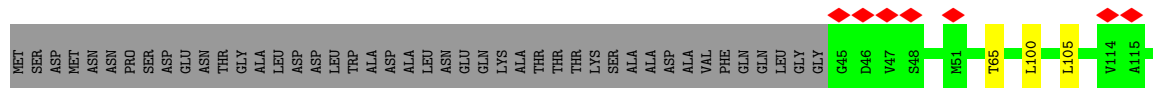
Chain HE:  57% 42%

• Molecule 4: Flagellar motor switch protein FliN

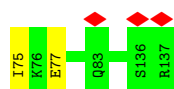
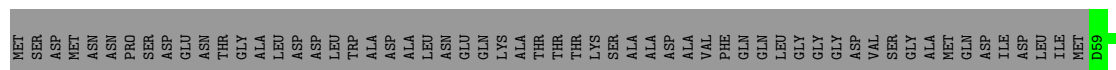
Chain IE:  5% 53% 45%



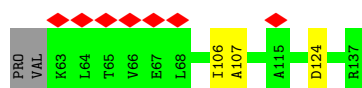
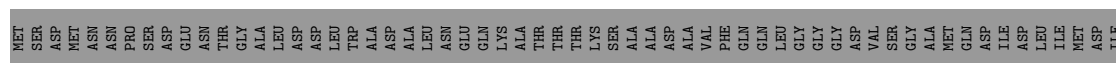
- Molecule 4: Flagellar motor switch protein FliN



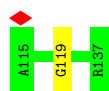
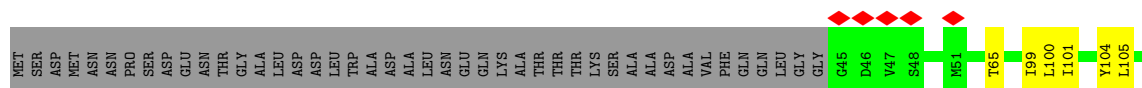
- Molecule 4: Flagellar motor switch protein FliN



- Molecule 4: Flagellar motor switch protein FliN

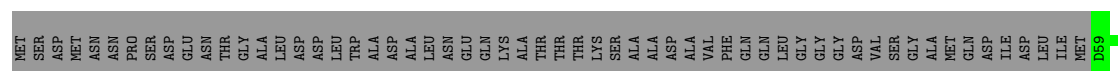


- Molecule 4: Flagellar motor switch protein FliN

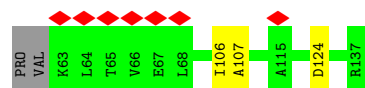
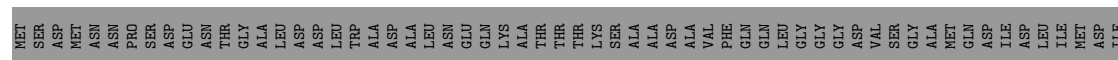


- Molecule 4: Flagellar motor switch protein FliN

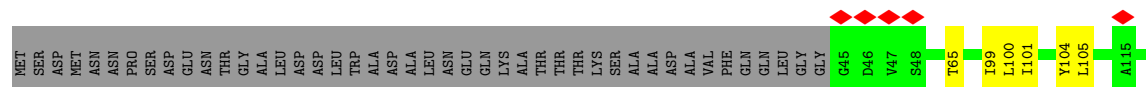




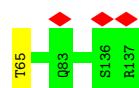
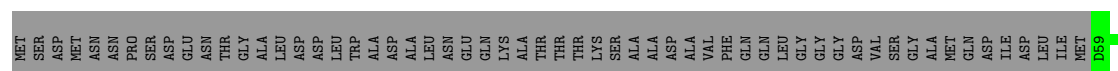
- Molecule 4: Flagellar motor switch protein FliN



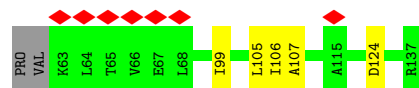
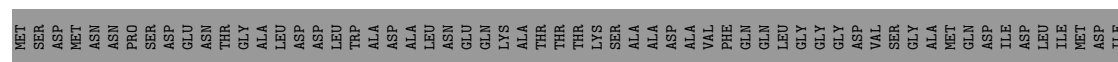
- Molecule 4: Flagellar motor switch protein FliN



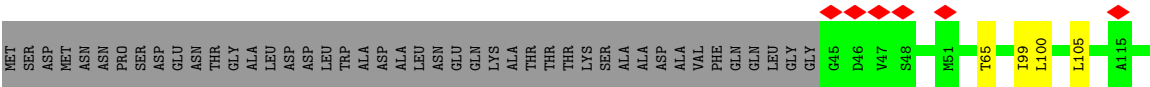
- Molecule 4: Flagellar motor switch protein FliN



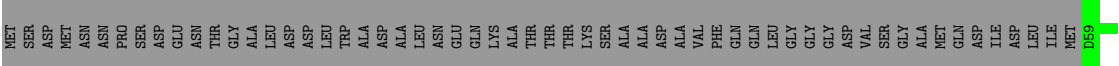
- Molecule 4: Flagellar motor switch protein FliN



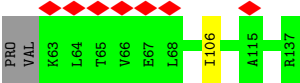
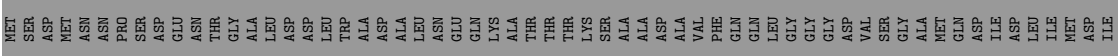
- Molecule 4: Flagellar motor switch protein FliN



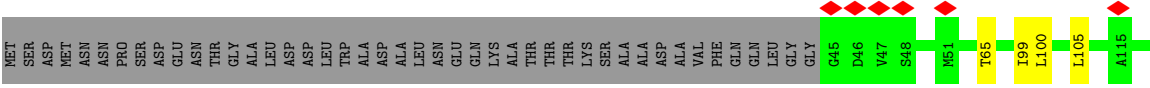
• Molecule 4: Flagellar motor switch protein FliN



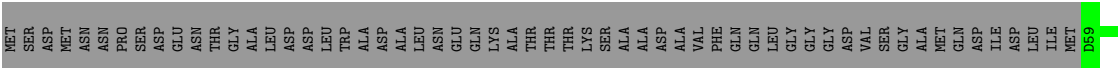
• Molecule 4: Flagellar motor switch protein FliN



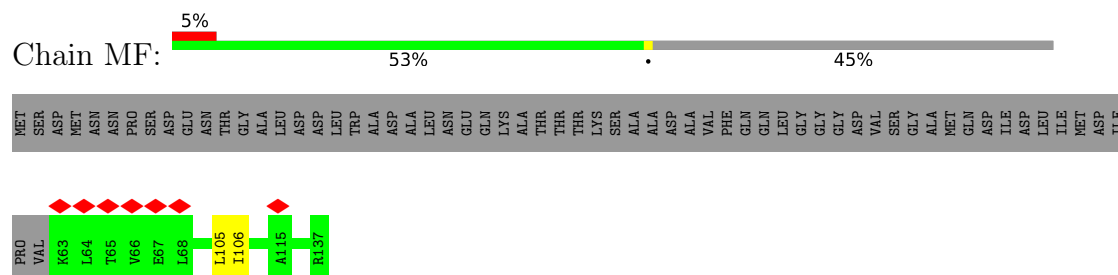
• Molecule 4: Flagellar motor switch protein FliN



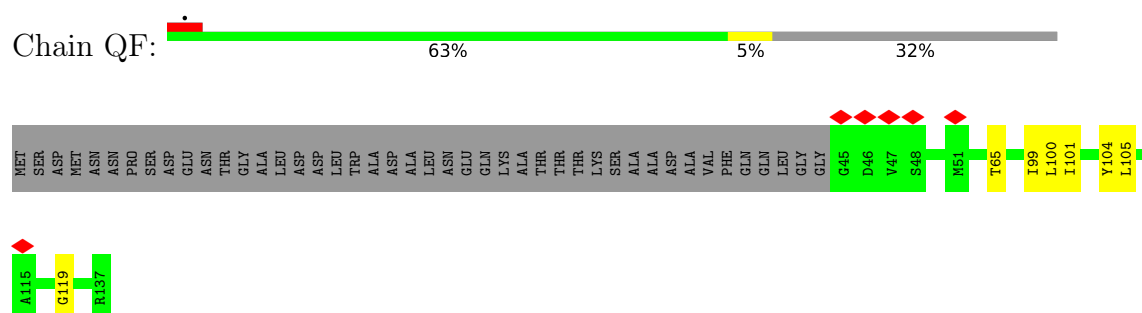
• Molecule 4: Flagellar motor switch protein FliN



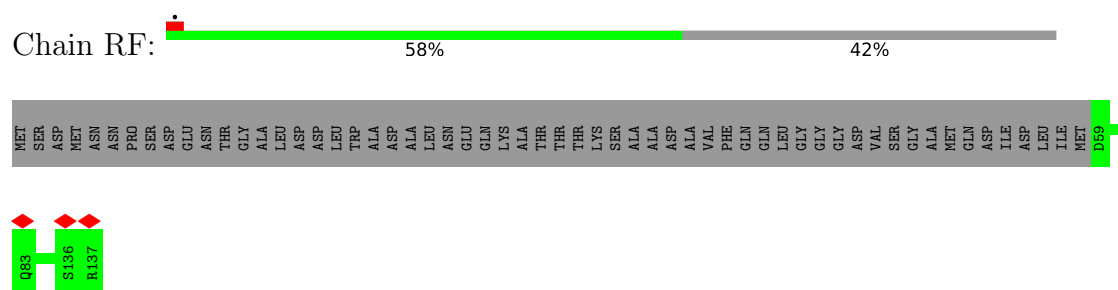
• Molecule 4: Flagellar motor switch protein FliN



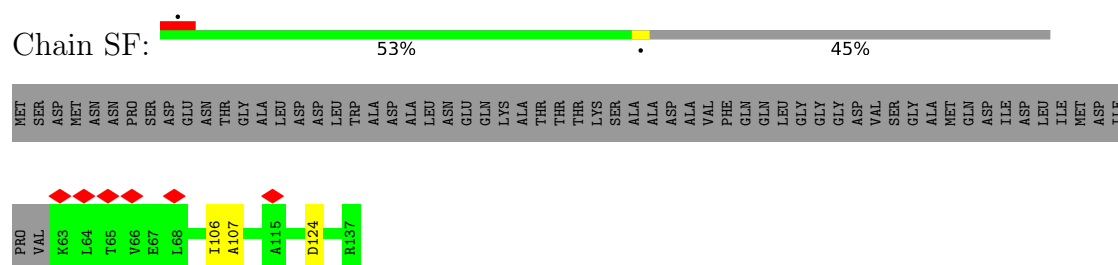
• Molecule 4: Flagellar motor switch protein FliN



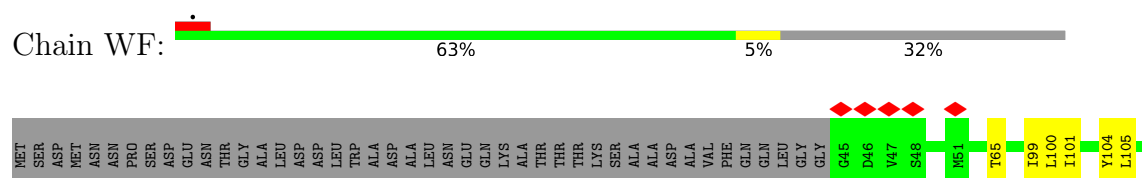
• Molecule 4: Flagellar motor switch protein FliN

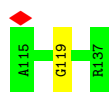


• Molecule 4: Flagellar motor switch protein FliN

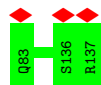
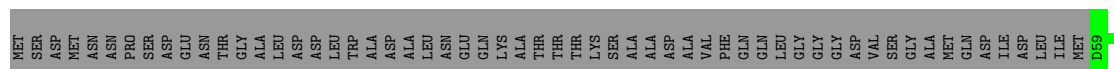


• Molecule 4: Flagellar motor switch protein FliN

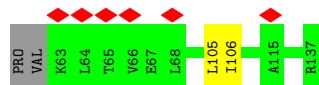
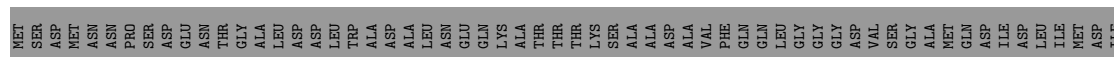




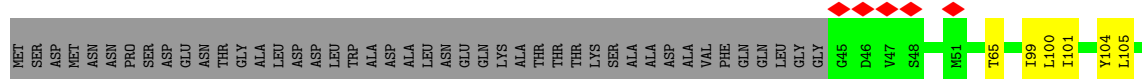
- Molecule 4: Flagellar motor switch protein FliN



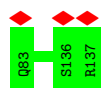
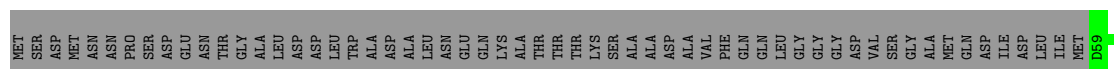
- Molecule 4: Flagellar motor switch protein FliN



- Molecule 4: Flagellar motor switch protein FliN



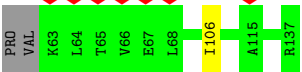
- Molecule 4: Flagellar motor switch protein FliN



- Molecule 4: Flagellar motor switch protein FliN



MET	SER	ASP	MET	ASN	PRO	SER	ASP	GLU	ASN	THR	GLY	ALA	LEU	ASP	ASP	LEU	TRP	ALA	ASP	ALA	LEU	LEU	ASN	GLU	GLN	LYS	ALA	ALA	ALA	ASP	ASP	ALA	VAL	PHE	GLN	GLN	LEU	GLY	GLY	GLY	ASP	VAL	SER	GLY	ALA	MET	GLN	ASP	ILE	LEU	ILE	MET	ASP	ILE
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• Molecule 4: Flagellar motor switch protein FliN



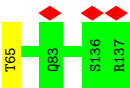
MET	SER	ASP	MET	ASN	PRO	SER	ASP	GLU	ASN	THR	GLY	ALA	LEU	ASP	ASP	LEU	TRP	ALA	ASP	ALA	LEU	LEU	ASN	GLU	GLN	LYS	ALA	ALA	ALA	ASP	ASP	ALA	VAL	PHE	GLN	GLN	LEU	GLY	GLY	G45	D46	V47	S48	M51	T65	I99	L100	L105	A115
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• Molecule 4: Flagellar motor switch protein FliN



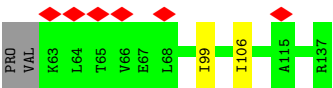
MET	SER	ASP	MET	ASN	PRO	SER	ASP	GLU	ASN	THR	GLY	ALA	LEU	ASP	ASP	LEU	TRP	ALA	ASP	ALA	LEU	LEU	ASN	GLU	GLN	LYS	ALA	ALA	ALA	ASP	ASP	ALA	VAL	PHE	GLN	GLN	LEU	GLY	GLY	G45	D46	V47	S48	M51	T65	I99	L100	L105	D59
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• Molecule 4: Flagellar motor switch protein FliN



MET	SER	ASP	MET	ASN	PRO	SER	ASP	GLU	ASN	THR	GLY	ALA	LEU	ASP	ASP	LEU	TRP	ALA	ASP	ALA	LEU	LEU	ASN	GLU	GLN	LYS	ALA	ALA	ALA	ASP	ASP	ALA	VAL	PHE	GLN	GLN	LEU	GLY	GLY	GLY	ASP	VAL	SER	GLY	ALA	MET	GLN	ASP	ILE	ASP	LEU	ILE	MET	ASP	ILE
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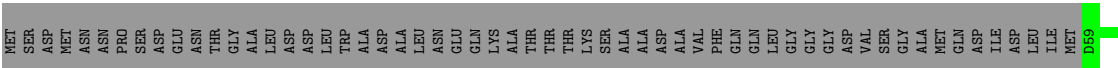
• Molecule 4: Flagellar motor switch protein FliN



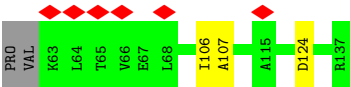
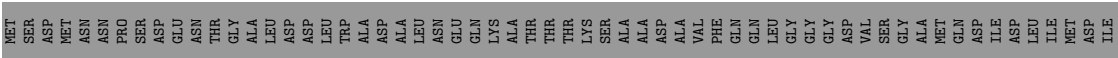
MET	SER	ASP	MET	ASN	PRO	SER	ASP	GLU	ASN	THR	GLY	ALA	LEU	ASP	ASP	LEU	TRP	ALA	ASP	ALA	LEU	LEU	ASN	GLU	GLN	LYS	ALA	ALA	ALA	ASP	ASP	ALA	VAL	PHE	GLN	GLN	LEU	GLY	GLY	G45	D46	V47	S48	M51	T65	I99	L100	L105	A115
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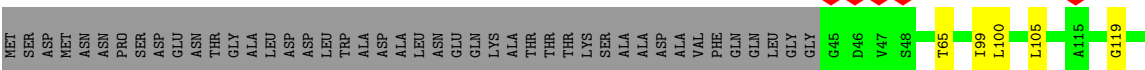
• Molecule 4: Flagellar motor switch protein FliN



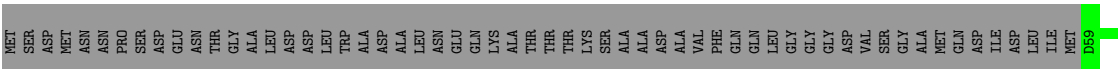
• Molecule 4: Flagellar motor switch protein FliN



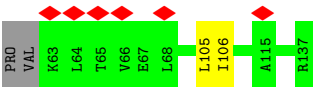
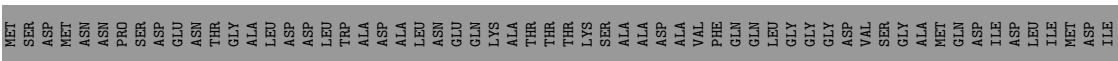
• Molecule 4: Flagellar motor switch protein FliN



• Molecule 4: Flagellar motor switch protein FliN



• Molecule 4: Flagellar motor switch protein FliN



4 Experimental information

Property	Value	Source
EM reconstruction method	SINGLE PARTICLE	Depositor
Imposed symmetry	POINT, C34	Depositor
Number of particles used	51268	Depositor
Resolution determination method	FSC 0.143 CUT-OFF	Depositor
CTF correction method	PHASE FLIPPING AND AMPLITUDE CORRECTION	Depositor
Microscope	FEI TITAN KRIOS	Depositor
Voltage (kV)	300	Depositor
Electron dose ($e^-/\text{\AA}^2$)	51.557	Depositor
Minimum defocus (nm)	1000	Depositor
Maximum defocus (nm)	2000	Depositor
Magnification	Not provided	
Image detector	GATAN K3 (6k x 4k)	Depositor
Maximum map value	0.483	Depositor
Minimum map value	-0.133	Depositor
Average map value	-0.001	Depositor
Map value standard deviation	0.016	Depositor
Recommended contour level	0.147	Depositor
Map size (Å)	1046.784, 1046.784, 1046.784	wwPDB
Map dimensions	768, 768, 768	wwPDB
Map angles (°)	90.0, 90.0, 90.0	wwPDB
Pixel spacing (Å)	1.363, 1.363, 1.363	Depositor

5 Model quality

5.1 Standard geometry

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

Mol	Chain	Bond lengths		Bond angles	
		RMSZ	$\# Z > 5$	RMSZ	$\# Z > 5$
1	A	0.22	0/233	0.35	0/324
1	BC	0.22	0/233	0.34	0/324
1	BF	0.22	0/233	0.34	0/324
1	DB	0.22	0/233	0.34	0/324
1	DE	0.22	0/233	0.34	0/324
1	F	0.22	0/233	0.34	0/324
1	FA	0.22	0/233	0.35	0/324
1	FD	0.22	0/233	0.34	0/324
1	FG	0.22	0/233	0.34	0/324
1	HC	0.22	0/233	0.34	0/324
1	HF	0.22	0/233	0.35	0/324
1	I	0.22	0/233	0.35	0/324
1	JB	0.22	0/233	0.34	0/324
1	JE	0.22	0/233	0.35	0/324
1	LA	0.22	0/233	0.35	0/324
1	LD	0.22	0/233	0.35	0/324
1	LG	0.22	0/233	0.35	0/324
1	NC	0.22	0/233	0.35	0/324
1	NF	0.22	0/233	0.35	0/324
1	PB	0.22	0/233	0.34	0/324
1	PE	0.22	0/233	0.34	0/324
1	RA	0.22	0/233	0.34	0/324
1	RD	0.22	0/233	0.35	0/324
1	RG	0.22	0/233	0.34	0/324
1	S	0.22	0/233	0.35	0/324
1	TC	0.22	0/233	0.34	0/324
1	TF	0.22	0/233	0.34	0/324
1	VB	0.22	0/233	0.34	0/324
1	VE	0.22	0/233	0.34	0/324
1	XA	0.22	0/233	0.35	0/324
1	XD	0.22	0/233	0.36	0/324
1	Z	0.22	0/233	0.34	0/324
1	ZC	0.22	0/233	0.34	0/324
1	ZF	0.22	0/233	0.34	0/324

Mol	Chain	Bond lengths		Bond angles	
		RMSZ	# Z >5	RMSZ	# Z >5
2	AA	0.30	0/2516	0.65	3/3393 (0.1%)
2	AD	0.29	0/2516	0.64	2/3393 (0.1%)
2	AG	0.30	0/2516	0.65	3/3393 (0.1%)
2	B	0.30	0/2516	0.64	3/3393 (0.1%)
2	CC	0.31	0/2516	0.66	3/3393 (0.1%)
2	CF	0.30	0/2516	0.66	6/3393 (0.2%)
2	EB	0.30	0/2516	0.66	2/3393 (0.1%)
2	EE	0.29	0/2516	0.64	2/3393 (0.1%)
2	G	0.30	0/2516	0.65	2/3393 (0.1%)
2	GA	0.29	0/2516	0.62	1/3393 (0.0%)
2	GD	0.29	0/2516	0.62	1/3393 (0.0%)
2	GG	0.29	0/2516	0.64	1/3393 (0.0%)
2	IC	0.29	0/2516	0.65	2/3393 (0.1%)
2	IF	0.30	0/2516	0.65	2/3393 (0.1%)
2	J	0.29	0/2516	0.63	2/3393 (0.1%)
2	KB	0.30	0/2516	0.66	3/3393 (0.1%)
2	KE	0.29	0/2516	0.64	2/3393 (0.1%)
2	MA	0.29	0/2516	0.64	2/3393 (0.1%)
2	MD	0.29	0/2516	0.64	2/3393 (0.1%)
2	MG	0.28	0/2516	0.63	2/3393 (0.1%)
2	OC	0.28	0/2516	0.63	2/3393 (0.1%)
2	OF	0.30	0/2516	0.65	2/3393 (0.1%)
2	QB	0.30	0/2516	0.63	0/3393
2	QE	0.29	0/2516	0.62	1/3393 (0.0%)
2	SA	0.30	0/2516	0.63	2/3393 (0.1%)
2	SD	0.29	0/2516	0.63	4/3393 (0.1%)
2	SG	0.28	0/2516	0.65	1/3393 (0.0%)
2	T	0.30	0/2516	0.65	3/3393 (0.1%)
2	UC	0.30	0/2516	0.66	3/3393 (0.1%)
2	UF	0.28	0/2516	0.60	0/3393
2	WB	0.30	0/2516	0.63	3/3393 (0.1%)
2	WE	0.29	0/2516	0.63	4/3393 (0.1%)
2	YA	0.30	0/2516	0.63	2/3393 (0.1%)
2	YD	0.31	0/2516	0.63	2/3393 (0.1%)
3	BA	0.42	0/2043	0.73	4/2790 (0.1%)
3	BD	0.47	2/2043 (0.1%)	0.73	4/2790 (0.1%)
3	BG	0.52	3/2043 (0.1%)	0.81	7/2790 (0.3%)
3	C	0.44	2/2043 (0.1%)	0.75	4/2790 (0.1%)
3	DC	0.53	3/2043 (0.1%)	0.82	7/2790 (0.3%)
3	DF	0.44	1/2043 (0.0%)	0.74	5/2790 (0.2%)
3	FB	0.44	1/2043 (0.0%)	0.74	5/2790 (0.2%)
3	FE	0.45	1/2043 (0.0%)	0.77	4/2790 (0.1%)
3	HA	0.46	2/2043 (0.1%)	0.76	4/2790 (0.1%)

Mol	Chain	Bond lengths		Bond angles	
		RMSZ	# Z >5	RMSZ	# Z >5
3	HD	0.45	1/2043 (0.0%)	0.75	4/2790 (0.1%)
3	HG	0.53	3/2043 (0.1%)	0.78	6/2790 (0.2%)
3	JC	0.52	4/2043 (0.2%)	0.79	6/2790 (0.2%)
3	JF	0.46	2/2043 (0.1%)	0.75	4/2790 (0.1%)
3	K	0.44	1/2043 (0.0%)	0.73	4/2790 (0.1%)
3	LB	0.46	2/2043 (0.1%)	0.75	4/2790 (0.1%)
3	LE	0.40	0/2043	0.72	3/2790 (0.1%)
3	M	0.47	2/2043 (0.1%)	0.76	5/2790 (0.2%)
3	NA	0.40	0/2043	0.72	3/2790 (0.1%)
3	ND	0.44	1/2043 (0.0%)	0.73	4/2790 (0.1%)
3	NG	0.46	2/2043 (0.1%)	0.77	5/2790 (0.2%)
3	PC	0.46	2/2043 (0.1%)	0.76	5/2790 (0.2%)
3	PF	0.47	2/2043 (0.1%)	0.77	6/2790 (0.2%)
3	RB	0.46	2/2043 (0.1%)	0.78	6/2790 (0.2%)
3	RE	0.43	1/2043 (0.0%)	0.72	3/2790 (0.1%)
3	TA	0.43	1/2043 (0.0%)	0.72	3/2790 (0.1%)
3	TD	0.38	0/2043	0.70	2/2790 (0.1%)
3	TG	0.45	1/2043 (0.0%)	0.74	5/2790 (0.2%)
3	V	0.39	0/2043	0.70	2/2790 (0.1%)
3	VC	0.44	1/2043 (0.0%)	0.74	4/2790 (0.1%)
3	VF	0.47	2/2043 (0.1%)	0.77	6/2790 (0.2%)
3	XB	0.48	2/2043 (0.1%)	0.77	6/2790 (0.2%)
3	XE	0.47	2/2043 (0.1%)	0.75	4/2790 (0.1%)
3	ZA	0.48	2/2043 (0.1%)	0.76	5/2790 (0.2%)
3	ZD	0.42	0/2043	0.73	4/2790 (0.1%)
4	AB	0.25	0/455	0.47	0/630
4	AC	0.27	0/368	0.47	0/508
4	AE	0.25	0/455	0.47	0/630
4	AF	0.26	0/368	0.47	0/508
4	BB	0.27	0/388	0.49	0/536
4	BE	0.26	0/388	0.49	0/536
4	CA	0.25	0/455	0.47	0/630
4	CB	0.26	0/368	0.47	0/508
4	CD	0.25	0/455	0.47	0/630
4	CE	0.26	0/368	0.46	0/508
4	CG	0.25	0/455	0.47	0/630
4	D	0.25	0/455	0.47	0/630
4	DA	0.26	0/388	0.49	0/536
4	DD	0.26	0/388	0.50	0/536
4	DG	0.26	0/388	0.50	0/536
4	E	0.26	0/388	0.50	0/536
4	EA	0.27	0/368	0.47	0/508
4	EC	0.25	0/455	0.47	0/630

Mol	Chain	Bond lengths		Bond angles	
		RMSZ	# Z >5	RMSZ	# Z >5
4	ED	0.27	0/368	0.47	0/508
4	EF	0.25	0/455	0.47	0/630
4	EG	0.27	0/368	0.47	0/508
4	FC	0.27	0/388	0.51	0/536
4	FF	0.27	0/388	0.49	0/536
4	GB	0.25	0/455	0.47	0/630
4	GC	0.27	0/368	0.47	0/508
4	GE	0.25	0/455	0.47	0/630
4	GF	0.27	0/368	0.47	0/508
4	H	0.27	0/368	0.47	0/508
4	HB	0.26	0/388	0.49	0/536
4	HE	0.27	0/388	0.50	0/536
4	IA	0.25	0/455	0.47	0/630
4	IB	0.27	0/368	0.48	0/508
4	ID	0.25	0/455	0.47	0/630
4	IE	0.26	0/368	0.45	0/508
4	IG	0.25	0/455	0.47	0/630
4	JA	0.26	0/388	0.49	0/536
4	JD	0.26	0/388	0.49	0/536
4	JG	0.26	0/388	0.49	0/536
4	KA	0.26	0/368	0.46	0/508
4	KC	0.25	0/455	0.47	0/630
4	KD	0.27	0/368	0.47	0/508
4	KF	0.25	0/455	0.47	0/630
4	KG	0.27	0/368	0.47	0/508
4	L	0.25	0/455	0.47	0/630
4	LC	0.26	0/388	0.49	0/536
4	LF	0.26	0/388	0.50	0/536
4	MB	0.25	0/455	0.47	0/630
4	MC	0.27	0/368	0.46	0/508
4	ME	0.25	0/455	0.47	0/630
4	MF	0.26	0/368	0.47	0/508
4	N	0.25	0/455	0.47	0/630
4	NB	0.26	0/388	0.49	0/536
4	NE	0.26	0/388	0.49	0/536
4	O	0.26	0/388	0.49	0/536
4	OA	0.25	0/455	0.47	0/630
4	OB	0.26	0/368	0.45	0/508
4	OD	0.25	0/455	0.47	0/630
4	OE	0.27	0/368	0.47	0/508
4	OG	0.25	0/455	0.47	0/630
4	P	0.26	0/388	0.51	0/536
4	PA	0.27	0/388	0.50	0/536

Mol	Chain	Bond lengths		Bond angles	
		RMSZ	# Z >5	RMSZ	# Z >5
4	PD	0.26	0/388	0.49	0/536
4	PG	0.27	0/388	0.50	0/536
4	Q	0.27	0/368	0.47	0/508
4	QA	0.27	0/368	0.47	0/508
4	QC	0.25	0/455	0.47	0/630
4	QD	0.26	0/368	0.46	0/508
4	QF	0.25	0/455	0.47	0/630
4	QG	0.27	0/368	0.47	0/508
4	R	0.26	0/368	0.46	0/508
4	RC	0.27	0/388	0.50	0/536
4	RF	0.26	0/388	0.49	0/536
4	SB	0.25	0/455	0.47	0/630
4	SC	0.27	0/368	0.47	0/508
4	SE	0.25	0/455	0.47	0/630
4	SF	0.27	0/368	0.47	0/508
4	TB	0.27	0/388	0.49	0/536
4	TE	0.26	0/388	0.50	0/536
4	UA	0.25	0/455	0.47	0/630
4	UB	0.27	0/368	0.47	0/508
4	UD	0.25	0/455	0.47	0/630
4	UE	0.27	0/368	0.47	0/508
4	UG	0.25	0/455	0.47	0/630
4	VA	0.27	0/388	0.50	0/536
4	VD	0.26	0/388	0.49	0/536
4	VG	0.26	0/388	0.50	0/536
4	W	0.25	0/455	0.47	0/630
4	WA	0.27	0/368	0.47	0/508
4	WC	0.25	0/455	0.47	0/630
4	WD	0.26	0/368	0.46	0/508
4	WF	0.25	0/455	0.47	0/630
4	WG	0.27	0/368	0.47	0/508
4	X	0.27	0/388	0.49	0/536
4	XC	0.27	0/388	0.51	0/536
4	XF	0.26	0/388	0.49	0/536
4	Y	0.27	0/368	0.47	0/508
4	YB	0.25	0/455	0.47	0/630
4	YC	0.26	0/368	0.46	0/508
4	YE	0.25	0/455	0.47	0/630
4	YF	0.27	0/368	0.47	0/508
4	ZB	0.26	0/388	0.49	0/536
4	ZE	0.27	0/388	0.49	0/536
All	All	0.35	51/204102 (0.0%)	0.64	228/278154 (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
2	AA	0	1
2	AD	0	1
2	AG	0	1
2	B	0	1
2	CC	0	1
2	CF	0	1
2	EB	0	1
2	EE	0	1
2	G	0	2
2	GA	0	1
2	GD	0	1
2	GG	0	1
2	IC	0	1
2	IF	0	1
2	J	0	1
2	KB	0	1
2	KE	0	1
2	MA	0	1
2	MD	0	1
2	MG	0	1
2	OC	0	1
2	OF	0	1
2	QB	0	1
2	QE	0	1
2	SA	0	2
2	SD	0	1
2	SG	0	1
2	T	0	2
2	UC	0	1
2	UF	0	1
2	WB	0	1
2	WE	0	1
2	YA	0	1
2	YD	0	1
All	All	0	37

All (51) bond length outliers are listed below:

Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
3	DC	174	PRO	CG-CD	-10.95	1.14	1.50
3	HG	195	PRO	CG-CD	-10.51	1.16	1.50
3	BD	174	PRO	CG-CD	-10.23	1.16	1.50
3	BG	174	PRO	CG-CD	-10.20	1.17	1.50
3	M	174	PRO	CG-CD	-9.84	1.18	1.50
3	XB	174	PRO	CG-CD	-9.73	1.18	1.50
3	JC	195	PRO	CG-CD	-9.02	1.20	1.50
3	JC	174	PRO	CG-CD	-8.78	1.21	1.50
3	RB	174	PRO	CG-CD	-8.45	1.22	1.50
3	VF	174	PRO	CG-CD	-8.42	1.22	1.50
3	PF	174	PRO	CG-CD	-8.38	1.23	1.50
3	HG	174	PRO	CG-CD	-8.06	1.24	1.50
3	TG	174	PRO	CG-CD	-7.70	1.25	1.50
3	ND	174	PRO	CG-CD	-7.42	1.26	1.50
3	VC	174	PRO	CG-CD	-7.40	1.26	1.50
3	FB	174	PRO	CG-CD	-7.27	1.26	1.50
3	K	174	PRO	CG-CD	-7.10	1.27	1.50
3	ZA	195	PRO	CG-CD	-6.96	1.27	1.50
3	DF	174	PRO	CG-CD	-6.95	1.27	1.50
3	HA	174	PRO	CG-CD	-6.77	1.28	1.50
3	FE	174	PRO	CG-CD	-6.74	1.28	1.50
3	ZA	174	PRO	CG-CD	-6.73	1.28	1.50
3	NG	174	PRO	CG-CD	-6.45	1.29	1.50
3	XE	174	PRO	CG-CD	-6.40	1.29	1.50
3	RE	174	PRO	CG-CD	-6.39	1.29	1.50
3	XE	195	PRO	CG-CD	-6.32	1.29	1.50
3	LB	195	PRO	CG-CD	-6.06	1.30	1.50
3	JF	195	PRO	CG-CD	-6.03	1.30	1.50
3	TA	174	PRO	CG-CD	-6.01	1.30	1.50
3	DC	97	PRO	CG-CD	-5.99	1.30	1.50
3	HG	195	PRO	N-CD	5.97	1.56	1.47
3	DC	174	PRO	N-CD	5.90	1.56	1.47
3	BG	97	PRO	CG-CD	-5.89	1.31	1.50
3	BD	174	PRO	N-CD	5.80	1.55	1.47
3	HD	174	PRO	CG-CD	-5.78	1.31	1.50
3	XB	174	PRO	N-CD	5.75	1.55	1.47
3	M	174	PRO	N-CD	5.71	1.55	1.47
3	BG	174	PRO	N-CD	5.61	1.55	1.47
3	LB	174	PRO	CG-CD	-5.55	1.32	1.50
3	PC	174	PRO	CG-CD	-5.54	1.32	1.50
3	NG	97	PRO	CG-CD	-5.53	1.32	1.50
3	JF	174	PRO	CG-CD	-5.50	1.32	1.50
3	PC	97	PRO	CG-CD	-5.45	1.32	1.50

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
3	HA	195	PRO	CG-CD	-5.42	1.32	1.50
3	JC	195	PRO	N-CD	5.39	1.55	1.47
3	VF	174	PRO	N-CD	5.19	1.55	1.47
3	JC	174	PRO	N-CD	5.18	1.55	1.47
3	C	174	PRO	CG-CD	-5.18	1.33	1.50
3	PF	174	PRO	N-CD	5.16	1.55	1.47
3	RB	174	PRO	N-CD	5.15	1.55	1.47
3	C	195	PRO	CG-CD	-5.10	1.33	1.50

All (228) bond angle outliers are listed below:

Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
3	DC	174	PRO	N-CD-CG	-12.15	84.98	103.20
3	FE	195	PRO	CA-N-CD	-11.99	94.71	111.50
3	HA	195	PRO	CA-N-CD	-11.96	94.76	111.50
3	LB	195	PRO	CA-N-CD	-11.92	94.81	111.50
3	JC	195	PRO	CA-N-CD	-11.88	94.87	111.50
3	K	174	PRO	CA-N-CD	-11.87	94.88	111.50
3	FE	174	PRO	CA-N-CD	-11.81	94.96	111.50
3	HG	195	PRO	CA-N-CD	-11.80	94.98	111.50
3	DC	195	PRO	CA-N-CD	-11.78	95.00	111.50
3	ZD	174	PRO	CA-N-CD	-11.78	95.01	111.50
3	HA	174	PRO	CA-N-CD	-11.77	95.02	111.50
3	ZA	195	PRO	CA-N-CD	-11.76	95.04	111.50
3	BG	195	PRO	CA-N-CD	-11.73	95.08	111.50
3	FB	174	PRO	CA-N-CD	-11.72	95.09	111.50
3	DF	174	PRO	CA-N-CD	-11.72	95.09	111.50
3	BA	174	PRO	CA-N-CD	-11.72	95.09	111.50
3	TG	174	PRO	CA-N-CD	-11.71	95.11	111.50
3	JF	174	PRO	CA-N-CD	-11.71	95.11	111.50
3	XE	195	PRO	CA-N-CD	-11.70	95.11	111.50
3	VF	174	PRO	CA-N-CD	-11.70	95.12	111.50
3	HD	174	PRO	CA-N-CD	-11.70	95.12	111.50
3	RE	174	PRO	CA-N-CD	-11.68	95.15	111.50
3	ZA	174	PRO	CA-N-CD	-11.68	95.15	111.50
3	TA	174	PRO	CA-N-CD	-11.67	95.16	111.50
3	VC	174	PRO	CA-N-CD	-11.66	95.18	111.50
3	XE	174	PRO	CA-N-CD	-11.66	95.18	111.50
3	HG	174	PRO	CA-N-CD	-11.65	95.19	111.50
3	PF	174	PRO	CA-N-CD	-11.65	95.19	111.50
3	RB	174	PRO	CA-N-CD	-11.63	95.21	111.50
3	JC	174	PRO	CA-N-CD	-11.63	95.22	111.50

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
3	JF	195	PRO	CA-N-CD	-11.62	95.24	111.50
3	C	174	PRO	CA-N-CD	-11.61	95.24	111.50
3	BG	174	PRO	N-CD-CG	-11.61	85.78	103.20
3	NG	174	PRO	CA-N-CD	-11.57	95.30	111.50
3	XB	174	PRO	N-CD-CG	-11.56	85.86	103.20
3	PC	174	PRO	CA-N-CD	-11.55	95.33	111.50
3	LB	174	PRO	CA-N-CD	-11.54	95.34	111.50
3	C	195	PRO	CA-N-CD	-11.53	95.36	111.50
3	ND	174	PRO	CA-N-CD	-11.52	95.37	111.50
3	BG	174	PRO	CA-N-CD	-11.51	95.38	111.50
3	XB	174	PRO	CA-N-CD	-11.48	95.43	111.50
3	M	174	PRO	CA-N-CD	-11.48	95.43	111.50
3	BD	174	PRO	CA-N-CD	-11.44	95.48	111.50
3	DC	174	PRO	CA-N-CD	-11.43	95.50	111.50
3	HD	195	PRO	CA-N-CD	-11.42	95.51	111.50
3	BD	174	PRO	N-CD-CG	-11.41	86.09	103.20
3	VF	195	PRO	CA-N-CD	-11.36	95.60	111.50
3	HG	195	PRO	N-CD-CG	-11.27	86.30	103.20
3	M	174	PRO	N-CD-CG	-11.19	86.42	103.20
3	TG	195	PRO	CA-N-CD	-11.18	95.85	111.50
3	PF	195	PRO	CA-N-CD	-11.17	95.86	111.50
3	XB	195	PRO	CA-N-CD	-11.16	95.87	111.50
3	LE	195	PRO	CA-N-CD	-11.13	95.92	111.50
3	RB	195	PRO	CA-N-CD	-11.12	95.93	111.50
3	NA	195	PRO	CA-N-CD	-11.11	95.94	111.50
3	M	195	PRO	CA-N-CD	-11.04	96.05	111.50
3	DC	97	PRO	CA-N-CD	-10.87	96.28	111.50
3	BG	97	PRO	CA-N-CD	-10.73	96.48	111.50
3	NG	97	PRO	CA-N-CD	-10.68	96.56	111.50
3	PC	97	PRO	CA-N-CD	-10.67	96.57	111.50
3	JC	174	PRO	N-CD-CG	-10.53	87.41	103.20
3	JC	195	PRO	N-CD-CG	-10.51	87.44	103.20
3	NA	174	PRO	CA-N-CD	-10.33	97.04	111.50
3	ZD	195	PRO	CA-N-CD	-10.25	97.14	111.50
3	RB	174	PRO	N-CD-CG	-10.20	87.90	103.20
3	VC	195	PRO	CA-N-CD	-10.19	97.24	111.50
3	FB	195	PRO	CA-N-CD	-10.18	97.25	111.50
3	VF	174	PRO	N-CD-CG	-10.18	87.93	103.20
3	DF	195	PRO	CA-N-CD	-10.18	97.25	111.50
3	BA	195	PRO	CA-N-CD	-10.16	97.28	111.50
3	PF	174	PRO	N-CD-CG	-10.13	88.00	103.20
3	K	195	PRO	CA-N-CD	-10.08	97.39	111.50

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
3	HG	174	PRO	N-CD-CG	-10.00	88.20	103.20
3	TA	195	PRO	CA-N-CD	-9.82	97.75	111.50
3	ND	195	PRO	CA-N-CD	-9.77	97.82	111.50
3	V	195	PRO	CA-N-CD	-9.76	97.83	111.50
3	TD	195	PRO	CA-N-CD	-9.74	97.86	111.50
3	TG	174	PRO	N-CD-CG	-9.68	88.68	103.20
3	LE	174	PRO	CA-N-CD	-9.64	98.00	111.50
3	TD	174	PRO	CA-N-CD	-9.61	98.05	111.50
3	FB	174	PRO	N-CD-CG	-9.56	88.85	103.20
3	ND	174	PRO	N-CD-CG	-9.46	89.01	103.20
3	VC	174	PRO	N-CD-CG	-9.44	89.04	103.20
3	V	174	PRO	CA-N-CD	-9.39	98.35	111.50
3	DF	174	PRO	N-CD-CG	-9.30	89.25	103.20
3	K	174	PRO	N-CD-CG	-9.28	89.27	103.20
3	ZA	195	PRO	N-CD-CG	-9.24	89.33	103.20
3	FE	174	PRO	N-CD-CG	-9.08	89.57	103.20
3	ZA	174	PRO	N-CD-CG	-9.04	89.64	103.20
3	HA	174	PRO	N-CD-CG	-9.01	89.69	103.20
3	NG	174	PRO	N-CD-CG	-8.86	89.91	103.20
3	XE	174	PRO	N-CD-CG	-8.82	89.97	103.20
3	XE	195	PRO	N-CD-CG	-8.81	89.99	103.20
3	RE	174	PRO	N-CD-CG	-8.80	90.00	103.20
3	LB	195	PRO	N-CD-CG	-8.68	90.19	103.20
3	JF	195	PRO	N-CD-CG	-8.64	90.24	103.20
3	TA	174	PRO	N-CD-CG	-8.61	90.28	103.20
3	PC	195	PRO	CA-N-CD	-8.59	99.47	111.50
3	NG	195	PRO	CA-N-CD	-8.51	99.59	111.50
3	HD	174	PRO	N-CD-CG	-8.34	90.70	103.20
3	PF	120	PRO	CA-N-CD	-8.27	99.92	111.50
3	RE	195	PRO	CA-N-CD	-8.23	99.97	111.50
3	HA	195	PRO	N-CD-CG	-8.19	90.92	103.20
3	PC	174	PRO	N-CD-CG	-8.16	90.95	103.20
3	JF	174	PRO	N-CD-CG	-8.14	90.98	103.20
3	LB	174	PRO	N-CD-CG	-8.10	91.05	103.20
3	RB	120	PRO	CA-N-CD	-8.06	100.21	111.50
3	C	195	PRO	N-CD-CG	-7.99	91.21	103.20
3	DC	195	PRO	N-CD-CG	-7.94	91.28	103.20
3	C	174	PRO	N-CD-CG	-7.93	91.31	103.20
3	BA	174	PRO	N-CD-CG	-7.84	91.44	103.20
3	BG	97	PRO	N-CD-CG	-7.82	91.48	103.20
3	FE	195	PRO	N-CD-CG	-7.79	91.52	103.20
3	PC	97	PRO	N-CD-CG	-7.78	91.53	103.20

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
3	NG	97	PRO	N-CD-CG	-7.76	91.56	103.20
3	BG	195	PRO	N-CD-CG	-7.75	91.57	103.20
3	HD	195	PRO	N-CD-CG	-7.61	91.78	103.20
3	ZD	174	PRO	N-CD-CG	-7.57	91.85	103.20
3	DC	97	PRO	N-CD-CG	-7.56	91.85	103.20
3	VF	97	PRO	CA-N-CD	-7.28	101.31	111.50
3	BD	195	PRO	CA-N-CD	-7.00	101.70	111.50
3	VF	195	PRO	N-CD-CG	-6.98	92.74	103.20
2	IF	41	MET	CA-CB-CG	6.94	125.09	113.30
2	UC	159	LEU	CA-CB-CG	6.90	131.18	115.30
2	OF	159	LEU	CA-CB-CG	6.85	131.05	115.30
3	XB	97	PRO	CA-N-CD	-6.74	102.06	111.50
2	SG	159	LEU	CA-CB-CG	6.72	130.75	115.30
3	DC	174	PRO	CA-CB-CG	-6.72	91.24	104.00
3	PF	195	PRO	N-CD-CG	-6.57	93.34	103.20
2	KB	159	LEU	CA-CB-CG	6.57	130.40	115.30
3	XB	195	PRO	N-CD-CG	-6.54	93.39	103.20
2	WE	188	LEU	CA-CB-CG	6.46	130.16	115.30
3	HG	195	PRO	CA-CB-CG	-6.45	91.74	104.00
3	BG	174	PRO	CA-CB-CG	-6.43	91.79	104.00
2	AG	121	LEU	CA-CB-CG	6.41	130.04	115.30
2	AA	159	LEU	CA-CB-CG	6.40	130.01	115.30
2	B	121	LEU	CA-CB-CG	6.39	130.00	115.30
2	CF	121	LEU	CA-CB-CG	6.39	129.99	115.30
3	BD	174	PRO	CA-CB-CG	-6.33	91.98	104.00
3	TG	195	PRO	N-CD-CG	-6.20	93.91	103.20
3	XB	174	PRO	CA-CB-CG	-6.19	92.23	104.00
3	M	174	PRO	CA-CB-CG	-6.18	92.27	104.00
3	LE	195	PRO	N-CD-CG	-6.17	93.95	103.20
3	NA	195	PRO	N-CD-CG	-6.15	93.98	103.20
3	M	195	PRO	N-CD-CG	-6.12	94.01	103.20
3	RB	195	PRO	N-CD-CG	-6.03	94.16	103.20
2	SA	37	LEU	CA-CB-CG	6.01	129.13	115.30
2	G	14	LEU	CA-CB-CG	5.96	129.01	115.30
3	JC	195	PRO	CA-CB-CG	-5.92	92.75	104.00
2	SD	121	LEU	CA-CB-CG	5.82	128.69	115.30
3	JC	174	PRO	CA-CB-CG	-5.81	92.95	104.00
2	MG	14	LEU	CA-CB-CG	5.80	128.65	115.30
2	EB	121	LEU	CA-CB-CG	5.78	128.59	115.30
2	GD	121	LEU	CA-CB-CG	5.76	128.54	115.30
3	RB	174	PRO	CA-CB-CG	-5.65	93.26	104.00
3	VF	174	PRO	CA-CB-CG	-5.61	93.34	104.00

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
3	PF	174	PRO	CA-CB-CG	-5.59	93.38	104.00
2	EB	310	LEU	CA-CB-CG	5.55	128.07	115.30
2	B	310	LEU	CA-CB-CG	5.55	128.06	115.30
2	UC	310	LEU	CA-CB-CG	5.52	128.00	115.30
2	CF	188	LEU	CA-CB-CG	5.51	127.98	115.30
2	CF	41	MET	CB-CG-SD	5.49	128.87	112.40
3	HG	174	PRO	CA-CB-CG	-5.49	93.57	104.00
2	MD	310	LEU	CA-CB-CG	5.48	127.91	115.30
2	GA	310	LEU	CA-CB-CG	5.46	127.86	115.30
2	J	310	LEU	CA-CB-CG	5.46	127.85	115.30
2	SA	310	LEU	CA-CB-CG	5.46	127.85	115.30
2	KB	310	LEU	CA-CB-CG	5.45	127.84	115.30
2	IF	310	LEU	CA-CB-CG	5.44	127.82	115.30
2	AG	310	LEU	CA-CB-CG	5.44	127.81	115.30
2	SD	310	LEU	CA-CB-CG	5.44	127.81	115.30
2	EE	310	LEU	CA-CB-CG	5.43	127.79	115.30
2	T	310	LEU	CA-CB-CG	5.43	127.78	115.30
2	CC	310	LEU	CA-CB-CG	5.43	127.78	115.30
2	QE	310	LEU	CA-CB-CG	5.42	127.77	115.30
2	AA	310	LEU	CA-CB-CG	5.41	127.75	115.30
2	YD	310	LEU	CA-CB-CG	5.40	127.73	115.30
2	GG	310	LEU	CA-CB-CG	5.40	127.72	115.30
2	AD	310	LEU	CA-CB-CG	5.39	127.70	115.30
2	KE	310	LEU	CA-CB-CG	5.39	127.69	115.30
2	CF	310	LEU	CA-CB-CG	5.39	127.69	115.30
2	MA	310	LEU	CA-CB-CG	5.38	127.67	115.30
2	WE	310	LEU	CA-CB-CG	5.37	127.65	115.30
2	CC	83	LYS	CA-CB-CG	5.37	125.21	113.40
2	IC	310	LEU	CA-CB-CG	5.37	127.65	115.30
2	OC	310	LEU	CA-CB-CG	5.37	127.65	115.30
2	YA	310	LEU	CA-CB-CG	5.36	127.64	115.30
2	WB	310	LEU	CA-CB-CG	5.36	127.63	115.30
2	G	8	ASP	CB-CG-OD1	5.35	123.12	118.30
2	MG	8	ASP	CB-CG-OD1	5.35	123.11	118.30
3	TG	174	PRO	CA-CB-CG	-5.30	93.94	104.00
2	KB	41	MET	CB-CG-SD	5.27	128.20	112.40
2	WB	37	LEU	CA-CB-CG	5.25	127.37	115.30
2	OF	121	LEU	CA-CB-CG	5.25	127.36	115.30
2	WE	8	ASP	CB-CG-OD1	5.22	123.00	118.30
3	FB	174	PRO	CA-CB-CG	-5.21	94.11	104.00
2	AA	8	ASP	CB-CG-OD1	5.20	122.98	118.30
2	T	8	ASP	CB-CG-OD1	5.20	122.98	118.30

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
2	WE	41	MET	CB-CG-SD	5.19	127.97	112.40
3	ND	174	PRO	CA-CB-CG	-5.17	94.18	104.00
3	ZD	195	PRO	N-CD-CG	-5.15	95.47	103.20
3	VC	174	PRO	CA-CB-CG	-5.15	94.22	104.00
2	UC	8	ASP	CB-CG-OD1	5.14	122.93	118.30
2	IC	8	ASP	CB-CG-OD1	5.14	122.92	118.30
3	BA	195	PRO	N-CD-CG	-5.13	95.50	103.20
2	CC	8	ASP	CB-CG-OD1	5.13	122.92	118.30
3	K	174	PRO	CA-CB-CG	-5.11	94.28	104.00
2	CF	188	LEU	CB-CG-CD2	-5.11	102.32	111.00
2	SD	8	ASP	CB-CG-OD1	5.10	122.89	118.30
2	EE	8	ASP	CB-CG-OD1	5.10	122.89	118.30
2	T	41	MET	CB-CG-SD	5.10	127.69	112.40
2	AD	8	ASP	CB-CG-OD1	5.09	122.88	118.30
3	DF	96	LEU	CA-CB-CG	5.09	127.01	115.30
2	YD	8	ASP	CB-CG-OD1	5.09	122.88	118.30
2	SD	37	LEU	CA-CB-CG	5.08	126.99	115.30
2	B	8	ASP	CB-CG-OD1	5.08	122.87	118.30
2	KE	8	ASP	CB-CG-OD1	5.08	122.87	118.30
2	AG	8	ASP	CB-CG-OD1	5.07	122.86	118.30
3	DF	174	PRO	CA-CB-CG	-5.05	94.40	104.00
2	J	8	ASP	CB-CG-OD1	5.05	122.85	118.30
2	MA	8	ASP	CB-CG-OD1	5.05	122.85	118.30
3	FB	96	LEU	CA-CB-CG	5.05	126.91	115.30
2	OC	8	ASP	CB-CG-OD1	5.04	122.84	118.30
2	YA	8	ASP	CB-CG-OD1	5.04	122.84	118.30
2	CF	8	ASP	CB-CG-OD2	5.04	122.83	118.30
3	ZA	195	PRO	CA-CB-CG	-5.03	94.44	104.00
2	MD	41	MET	CB-CG-SD	5.02	127.46	112.40
2	WB	8	ASP	CB-CG-OD1	5.02	122.81	118.30

There are no chirality outliers.

All (37) planarity outliers are listed below:

Mol	Chain	Res	Type	Group
2	AA	90	ALA	Peptide
2	AD	90	ALA	Peptide
2	AG	90	ALA	Peptide
2	B	90	ALA	Peptide
2	CC	90	ALA	Peptide
2	CF	90	ALA	Peptide
2	EB	90	ALA	Peptide

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Mol	Chain	Res	Type	Group
2	EE	90	ALA	Peptide
2	G	139	ARG	Sidechain
2	G	90	ALA	Peptide
2	GA	90	ALA	Peptide
2	GD	90	ALA	Peptide
2	GG	90	ALA	Peptide
2	IC	90	ALA	Peptide
2	IF	90	ALA	Peptide
2	J	90	ALA	Peptide
2	KB	90	ALA	Peptide
2	KE	90	ALA	Peptide
2	MA	90	ALA	Peptide
2	MD	90	ALA	Peptide
2	MG	90	ALA	Peptide
2	OC	90	ALA	Peptide
2	OF	90	ALA	Peptide
2	QB	90	ALA	Peptide
2	QE	90	ALA	Peptide
2	SA	139	ARG	Sidechain
2	SA	90	ALA	Peptide
2	SD	90	ALA	Peptide
2	SG	90	ALA	Peptide
2	T	139	ARG	Sidechain
2	T	90	ALA	Peptide
2	UC	90	ALA	Peptide
2	UF	90	ALA	Peptide
2	WB	90	ALA	Peptide
2	WE	90	ALA	Peptide
2	YA	90	ALA	Peptide
2	YD	90	ALA	Peptide

5.2 Too-close contacts

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

Mol	Chain	Non-H	H(model)	H(added)	Clashes	Symm-Clashes
1	A	234	0	102	0	0
1	BC	234	0	102	0	0
1	BF	234	0	102	0	0

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Mol	Chain	Non-H	H(model)	H(added)	Clashes	Symm-Clashes
1	DB	234	0	102	0	0
1	DE	234	0	102	0	0
1	F	234	0	102	0	0
1	FA	234	0	102	0	0
1	FD	234	0	102	0	0
1	FG	234	0	102	0	0
1	HC	234	0	102	1	0
1	HF	234	0	102	0	0
1	I	234	0	102	0	0
1	JB	234	0	102	0	0
1	JE	234	0	102	0	0
1	LA	234	0	102	0	0
1	LD	234	0	102	0	0
1	LG	234	0	102	1	0
1	NC	234	0	102	0	0
1	NF	234	0	102	1	0
1	PB	234	0	102	1	0
1	PE	234	0	102	1	0
1	RA	234	0	102	0	0
1	RD	234	0	102	2	0
1	RG	234	0	102	0	0
1	S	234	0	102	0	0
1	TC	234	0	102	0	0
1	TF	234	0	102	0	0
1	VB	234	0	102	0	0
1	VE	234	0	102	0	0
1	XA	234	0	102	1	0
1	XD	234	0	102	2	0
1	Z	234	0	102	1	0
1	ZC	234	0	102	0	0
1	ZF	234	0	102	0	0
2	AA	2496	0	2546	176	0
2	AD	2496	0	2546	169	0
2	AG	2496	0	2546	185	0
2	B	2496	0	2546	175	0
2	CC	2496	0	2546	186	0
2	CF	2496	0	2546	185	0
2	EB	2496	0	2546	177	0
2	EE	2496	0	2546	161	0
2	G	2496	0	2546	182	0
2	GA	2496	0	2546	164	0
2	GD	2496	0	2546	181	0

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Mol	Chain	Non-H	H(model)	H(added)	Clashes	Symm-Clashes
2	GG	2496	0	2546	168	0
2	IC	2496	0	2546	181	0
2	IF	2496	0	2546	194	0
2	J	2496	0	2546	175	0
2	KB	2496	0	2546	183	0
2	KE	2496	0	2546	173	0
2	MA	2496	0	2546	178	0
2	MD	2496	0	2546	180	0
2	MG	2496	0	2546	171	0
2	OC	2496	0	2546	166	0
2	OF	2496	0	2546	209	0
2	QB	2496	0	2546	189	0
2	QE	2496	0	2546	170	0
2	SA	2496	0	2546	181	0
2	SD	2496	0	2546	182	0
2	SG	2496	0	2546	190	0
2	T	2496	0	2546	195	0
2	UC	2496	0	2546	179	0
2	UF	2496	0	2546	188	0
2	WB	2496	0	2546	183	0
2	WE	2496	0	2546	179	0
2	YA	2496	0	2546	182	0
2	YD	2496	0	2546	175	0
3	BA	2086	0	1798	101	0
3	BD	2086	0	1797	99	0
3	BG	2086	0	1799	99	0
3	C	2086	0	1798	99	0
3	DC	2086	0	1799	102	0
3	DF	2086	0	1798	95	0
3	FB	2086	0	1798	93	0
3	FE	2086	0	1798	97	0
3	HA	2086	0	1798	86	0
3	HD	2086	0	1798	97	0
3	HG	2086	0	1798	96	0
3	JC	2086	0	1799	100	0
3	JF	2086	0	1798	81	0
3	K	2086	0	1798	96	0
3	LB	2086	0	1798	95	0
3	LE	2086	0	1798	95	0
3	M	2086	0	1798	95	0
3	NA	2086	0	1798	99	0
3	ND	2086	0	1798	87	0

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Mol	Chain	Non-H	H(model)	H(added)	Clashes	Symm-Clashes
3	NG	2086	0	1798	84	0
3	PC	2086	0	1798	95	0
3	PF	2086	0	1798	97	0
3	RB	2086	0	1798	102	0
3	RE	2086	0	1799	92	0
3	TA	2086	0	1799	97	0
3	TD	2086	0	1799	89	0
3	TG	2086	0	1799	104	0
3	V	2086	0	1799	93	0
3	VC	2086	0	1797	97	0
3	VF	2086	0	1798	96	0
3	XB	2086	0	1798	114	0
3	XE	2086	0	1799	93	0
3	ZA	2086	0	1799	96	0
3	ZD	2086	0	1798	96	0
4	AB	456	0	206	5	0
4	AC	369	0	166	2	0
4	AE	456	0	206	6	0
4	AF	369	0	166	4	0
4	BB	389	0	173	0	0
4	BE	389	0	173	1	0
4	CA	456	0	206	4	0
4	CB	369	0	166	2	0
4	CD	456	0	206	4	0
4	CE	369	0	166	6	0
4	CG	456	0	206	5	0
4	D	456	0	206	3	0
4	DA	389	0	173	0	0
4	DD	389	0	173	0	0
4	DG	389	0	173	0	0
4	E	389	0	173	0	0
4	EA	369	0	166	3	0
4	EC	456	0	206	5	0
4	ED	369	0	166	1	0
4	EF	456	0	206	3	0
4	EG	369	0	166	1	0
4	FC	389	0	173	0	0
4	FF	389	0	173	0	0
4	GB	456	0	206	4	0
4	GC	369	0	166	1	0
4	GE	456	0	206	3	0
4	GF	369	0	166	2	0

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Mol	Chain	Non-H	H(model)	H(added)	Clashes	Symm-Clashes
4	H	369	0	166	2	0
4	HB	389	0	173	0	0
4	HE	389	0	173	1	0
4	IA	456	0	206	4	0
4	IB	369	0	166	2	0
4	ID	456	0	206	4	0
4	IE	369	0	166	5	0
4	IG	456	0	206	4	0
4	JA	389	0	173	0	0
4	JD	389	0	173	0	0
4	JG	389	0	173	1	0
4	KA	369	0	166	2	0
4	KC	456	0	206	4	0
4	KD	369	0	166	1	0
4	KF	456	0	206	3	0
4	KG	369	0	166	2	0
4	L	456	0	206	4	0
4	LC	389	0	173	1	0
4	LF	389	0	173	0	0
4	MB	456	0	206	4	0
4	MC	369	0	166	2	0
4	ME	456	0	206	3	0
4	MF	369	0	166	2	0
4	N	456	0	206	4	0
4	NB	389	0	173	1	0
4	NE	389	0	173	1	0
4	O	389	0	173	1	0
4	OA	456	0	206	3	0
4	OB	369	0	166	8	0
4	OD	456	0	206	3	0
4	OE	369	0	166	3	0
4	OG	456	0	206	3	0
4	P	389	0	173	0	0
4	PA	389	0	173	0	0
4	PD	389	0	173	1	0
4	PG	389	0	173	0	0
4	Q	369	0	166	2	0
4	QA	369	0	166	2	0
4	QC	456	0	206	5	0
4	QD	369	0	166	6	0
4	QF	456	0	206	5	0
4	QG	369	0	166	2	0

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Mol	Chain	Non-H	H(model)	H(added)	Clashes	Symm-Clashes
4	R	369	0	166	8	0
4	RC	389	0	173	0	0
4	RF	389	0	173	0	0
4	SB	456	0	206	5	0
4	SC	369	0	166	3	0
4	SE	456	0	206	4	0
4	SF	369	0	166	2	0
4	TB	389	0	173	0	0
4	TE	389	0	173	0	0
4	UA	456	0	206	3	0
4	UB	369	0	166	3	0
4	UD	456	0	206	3	0
4	UE	369	0	166	2	0
4	UG	456	0	206	3	0
4	VA	389	0	173	0	0
4	VD	389	0	173	0	0
4	VG	389	0	173	0	0
4	W	456	0	206	4	0
4	WA	369	0	166	2	0
4	WC	456	0	206	4	0
4	WD	369	0	166	6	0
4	WF	456	0	206	5	0
4	WG	369	0	166	2	0
4	X	389	0	173	0	0
4	XC	389	0	173	1	0
4	XF	389	0	173	0	0
4	Y	369	0	166	2	0
4	YB	456	0	206	6	0
4	YC	369	0	166	8	0
4	YE	456	0	206	4	0
4	YF	369	0	166	2	0
4	ZB	389	0	173	0	0
4	ZE	389	0	173	1	0
All	All	205020	0	169702	8071	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 22.

All (8071) 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)
2:WB:83:LYS:HE3	2:CC:23:ALA:HA	1.33	1.10
2:YD:186:GLN:HE22	3:FE:71:ASN:HA	1.18	1.04
2:WB:89:ARG:HG2	2:WB:90:ALA:H	1.23	1.02
2:QB:80:VAL:HA	2:QB:83:LYS:HD3	1.39	1.01
2:IC:80:VAL:HA	2:IC:83:LYS:HD3	1.43	0.99
2:AA:186:GLN:HE22	3:HA:71:ASN:HA	1.26	0.99
2:UF:202:ILE:HD13	2:AG:139:ARG:HH12	1.22	0.99
2:KB:80:VAL:HA	2:KB:83:LYS:HD3	1.45	0.98
2:EB:89:ARG:HG2	2:EB:90:ALA:H	1.27	0.98
2:EE:89:ARG:HG2	2:EE:90:ALA:H	1.29	0.98
2:UF:202:ILE:HA	2:AG:139:ARG:HH22	1.27	0.97
2:AA:186:GLN:HB3	3:HA:74:ARG:HH12	1.29	0.97
2:IF:186:GLN:HB3	3:PF:74:ARG:HH12	1.27	0.97
2:SD:186:GLN:HB3	3:ZD:74:ARG:HH12	1.27	0.97
2:QE:186:GLN:HE22	3:XE:71:ASN:HA	1.28	0.96
2:G:89:ARG:HG2	2:G:90:ALA:H	1.26	0.95
2:EB:83:LYS:HB3	2:KB:27:LYS:HB2	1.47	0.95
2:YA:89:ARG:HG2	2:YA:90:ALA:H	1.29	0.95
2:CF:128:GLN:HG2	3:DF:131:PHE:HB3	1.43	0.95
2:YD:89:ARG:HD2	2:YD:90:ALA:H	1.28	0.95
2:MA:89:ARG:HD2	2:MA:90:ALA:H	1.28	0.95
2:MA:128:GLN:HG2	3:NA:131:PHE:HB3	1.48	0.95
2:SA:89:ARG:HG2	2:SA:90:ALA:H	1.28	0.94
2:MD:80:VAL:HA	2:MD:83:LYS:HD3	1.49	0.94
2:SD:120:ASP:OD2	3:ZD:76:SER:OG	1.86	0.94
2:MD:128:GLN:HG2	3:ND:131:PHE:HB3	1.50	0.93
2:AA:128:GLN:HG2	3:BA:131:PHE:HB3	1.50	0.92
2:EB:201:GLU:HB3	2:KB:139:ARG:HH21	1.35	0.92
2:AD:17:ILE:HD12	2:AD:21:ARG:HG3	1.52	0.92
2:YA:201:GLU:HG3	2:EB:139:ARG:HD2	1.51	0.92
2:KE:128:GLN:HG2	3:LE:131:PHE:HB3	1.50	0.92
2:GD:128:GLN:HG2	3:HD:131:PHE:HB3	1.52	0.91
2:YD:120:ASP:OD2	3:FE:76:SER:OG	1.87	0.91
2:AD:128:GLN:HG2	3:BD:131:PHE:HB3	1.52	0.91
2:UF:89:ARG:HG2	2:UF:90:ALA:H	1.35	0.91
2:J:128:GLN:HG2	3:K:131:PHE:HB3	1.50	0.90
2:KE:83:LYS:HD3	2:QE:26:PHE:HB3	1.52	0.90
2:UF:80:VAL:HA	2:UF:83:LYS:HE3	1.53	0.90
2:AA:120:ASP:OD2	3:HA:76:SER:OG	1.89	0.90
2:OF:186:GLN:HE22	3:VF:71:ASN:HA	1.35	0.90
2:KB:83:LYS:HG2	2:QB:26:PHE:HB3	1.52	0.90
2:G:128:GLN:HG2	3:M:131:PHE:HB3	1.52	0.90

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:T:45:ARG:HG3	2:T:46:GLN:HG3	1.53	0.90
2:YD:128:GLN:HG2	3:ZD:131:PHE:HB3	1.51	0.90
2:OF:188:LEU:HD21	3:VF:74:ARG:HH11	1.37	0.90
2:UF:128:GLN:HG2	3:VF:131:PHE:HB3	1.54	0.90
2:OC:186:GLN:OE1	3:VC:71:ASN:ND2	2.05	0.89
2:IF:186:GLN:HE22	3:PF:71:ASN:HA	1.36	0.89
2:MD:75:GLU:HA	2:MD:78:ARG:HH21	1.37	0.89
2:YD:201:GLU:HG3	2:EE:139:ARG:HD2	1.52	0.89
2:AD:45:ARG:HG3	2:AD:46:GLN:HG3	1.52	0.89
2:AG:186:GLN:HE22	3:HG:71:ASN:HA	1.38	0.89
2:J:186:GLN:HE22	3:V:71:ASN:HA	1.36	0.89
2:IF:102:ASP:HB2	2:IF:108:GLU:HB2	1.54	0.89
2:AG:113:MET:O	2:AG:141:GLN:NE2	2.05	0.88
2:MG:186:GLN:OE1	3:TG:71:ASN:ND2	2.04	0.88
2:MD:120:ASP:OD2	3:TD:76:SER:OG	1.90	0.88
2:MG:17:ILE:HD12	2:MG:21:ARG:HG3	1.54	0.88
2:KE:186:GLN:HB3	3:RE:74:ARG:HH12	1.37	0.88
2:CF:186:GLN:NE2	3:JF:71:ASN:OD1	2.04	0.88
2:IF:128:GLN:HG2	3:JF:131:PHE:HB3	1.53	0.88
2:G:17:ILE:HD12	2:G:21:ARG:HG3	1.53	0.88
2:GA:108:GLU:HA	2:GA:111:ASN:HD22	1.39	0.88
2:IC:102:ASP:HB2	2:IC:108:GLU:HB2	1.54	0.88
2:MG:128:GLN:HG2	3:NG:131:PHE:HB3	1.52	0.88
2:IC:75:GLU:HA	2:IC:78:ARG:HH21	1.39	0.88
2:GD:186:GLN:HE22	3:ND:71:ASN:HA	1.39	0.88
2:SD:128:GLN:HG2	3:TD:131:PHE:HB3	1.55	0.87
2:OF:128:GLN:HG2	3:PF:131:PHE:HB3	1.54	0.87
2:OF:69:LEU:HD11	2:UF:47:ILE:HG22	1.56	0.87
2:MA:75:GLU:HA	2:MA:78:ARG:HH21	1.37	0.87
2:YA:188:LEU:HD13	3:FB:74:ARG:HD2	1.54	0.87
2:UC:128:GLN:HG2	3:VC:131:PHE:HB3	1.55	0.87
2:SG:128:GLN:HG2	3:TG:131:PHE:HB3	1.55	0.87
2:G:47:ILE:HG22	2:SG:69:LEU:HD11	1.57	0.87
2:SA:188:LEU:HD13	3:ZA:74:ARG:HD2	1.55	0.87
2:WE:128:GLN:HG2	3:XE:131:PHE:HB3	1.55	0.87
2:EB:168:GLN:HE22	3:FB:130:LEU:HD11	1.37	0.87
2:WB:128:GLN:HG2	3:XB:131:PHE:HB3	1.56	0.87
2:GA:201:GLU:O	2:MA:139:ARG:NH2	2.08	0.87
2:EB:188:LEU:HD13	3:LB:74:ARG:HD2	1.58	0.86
2:KB:75:GLU:HA	2:KB:78:ARG:HH21	1.40	0.86
2:QB:128:GLN:HG2	3:RB:131:PHE:HB3	1.55	0.86

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:KE:201:GLU:HG3	2:QE:139:ARG:HD2	1.57	0.86
2:B:75:GLU:HA	2:B:78:ARG:HH21	1.38	0.86
2:T:75:GLU:HA	2:T:78:ARG:HH21	1.38	0.86
2:AG:186:GLN:HB3	3:HG:74:ARG:HH22	1.41	0.86
2:EB:17:ILE:HD12	2:EB:21:ARG:HG3	1.57	0.86
2:AG:188:LEU:HD21	3:HG:74:ARG:HH12	1.40	0.86
2:T:128:GLN:HG2	3:V:131:PHE:HB3	1.55	0.86
2:EB:207:LYS:HG3	2:EB:210:GLN:HG2	1.58	0.86
2:IC:186:GLN:HE22	3:PC:71:ASN:HA	1.40	0.86
2:CC:128:GLN:HG2	3:DC:131:PHE:HB3	1.56	0.86
2:AD:186:GLN:HE22	3:HD:71:ASN:HA	1.41	0.86
2:EB:128:GLN:HG2	3:FB:131:PHE:HB3	1.55	0.85
2:IC:83:LYS:HB3	2:OC:27:LYS:HB2	1.55	0.85
2:EE:128:GLN:HG2	3:FE:131:PHE:HB3	1.58	0.85
2:G:186:GLN:HE22	3:C:71:ASN:HA	1.40	0.85
2:SA:120:ASP:OD1	2:SA:123:ARG:NH1	2.10	0.85
2:YD:186:GLN:HB3	3:FE:74:ARG:HH12	1.39	0.85
2:WE:102:ASP:HB2	2:WE:108:GLU:HB2	1.57	0.85
2:YA:128:GLN:HG2	3:ZA:131:PHE:HB3	1.56	0.85
2:CC:80:VAL:HA	2:CC:83:LYS:HE3	1.57	0.85
2:QB:102:ASP:HB2	2:QB:108:GLU:HB2	1.56	0.85
4:OB:99:ILE:O	3:RB:252:ARG:NH2	2.09	0.85
2:QB:186:GLN:HE22	3:XB:71:ASN:HA	1.40	0.84
2:IC:207:LYS:HG3	2:IC:210:GLN:HG2	1.59	0.84
2:KB:128:GLN:HG2	3:LB:131:PHE:HB3	1.56	0.84
2:MD:207:LYS:HG3	2:MD:210:GLN:HG2	1.59	0.84
2:GG:128:GLN:HG2	3:HG:131:PHE:HB3	1.56	0.84
2:IC:128:GLN:HG2	3:JC:131:PHE:HB3	1.57	0.84
2:UC:188:LEU:HD13	3:BD:74:ARG:HD2	1.58	0.84
2:G:102:ASP:HB2	2:G:108:GLU:HB2	1.60	0.84
2:J:26:PHE:HE1	2:J:34:VAL:HG22	1.42	0.84
2:SA:102:ASP:HB2	2:SA:108:GLU:HB2	1.58	0.84
2:UC:186:GLN:HE22	3:BD:71:ASN:HA	1.42	0.84
2:AG:207:LYS:HG3	2:AG:210:GLN:HG2	1.57	0.84
2:B:102:ASP:HB2	2:B:108:GLU:HB2	1.59	0.84
2:B:207:LYS:HZ2	2:B:210:GLN:HG3	1.43	0.84
2:WE:26:PHE:HE1	2:WE:34:VAL:HG22	1.43	0.84
2:KB:188:LEU:HD13	3:RB:74:ARG:HD2	1.58	0.84
2:AG:128:GLN:HG2	3:BG:131:PHE:HB3	1.57	0.84
2:SA:75:GLU:HA	2:SA:78:ARG:HH21	1.43	0.83
2:SA:108:GLU:HA	2:SA:111:ASN:HD22	1.41	0.83

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:GD:188:LEU:HD13	3:ND:74:ARG:HD2	1.60	0.83
2:MD:102:ASP:HB2	2:MD:108:GLU:HB2	1.60	0.83
2:UF:102:ASP:HB2	2:UF:108:GLU:HB2	1.60	0.83
2:B:186:GLN:HE22	3:K:71:ASN:HA	1.43	0.83
2:YA:75:GLU:HA	2:YA:78:ARG:HH21	1.43	0.83
4:R:99:ILE:O	3:V:252:ARG:NH2	2.12	0.83
2:GA:17:ILE:HD12	2:GA:21:ARG:HG3	1.59	0.83
2:T:83:LYS:HG2	2:AA:26:PHE:HB3	1.61	0.83
2:CC:102:ASP:HB2	2:CC:108:GLU:HB2	1.61	0.83
2:QE:186:GLN:HB3	3:XE:74:ARG:NH2	1.93	0.83
2:WE:186:GLN:HB3	3:DF:74:ARG:HH12	1.43	0.83
2:CF:17:ILE:HD12	2:CF:21:ARG:HG3	1.59	0.83
2:B:128:GLN:HG2	3:C:131:PHE:HB3	1.58	0.83
2:KB:207:LYS:HZ2	2:KB:210:GLN:HG3	1.43	0.83
2:IC:168:GLN:HE22	3:JC:130:LEU:HD11	1.42	0.83
2:GD:207:LYS:HZ2	2:GD:210:GLN:HG3	1.44	0.83
4:CE:99:ILE:O	3:FE:252:ARG:NH2	2.12	0.82
2:CF:26:PHE:CE1	2:CF:34:VAL:HG22	2.14	0.82
2:GA:128:GLN:HG2	3:HA:131:PHE:HB3	1.58	0.82
2:J:102:ASP:HB2	2:J:108:GLU:HB2	1.60	0.82
2:IC:188:LEU:HD13	3:PC:74:ARG:HD2	1.60	0.82
2:WB:206:MET:SD	2:WB:207:LYS:N	2.51	0.82
2:SD:75:GLU:HA	2:SD:78:ARG:HH21	1.44	0.82
2:QE:75:GLU:HA	2:QE:78:ARG:HH21	1.45	0.82
2:SA:128:GLN:HG2	3:TA:131:PHE:HB3	1.59	0.82
2:AD:102:ASP:HB2	2:AD:108:GLU:HB2	1.58	0.82
4:IE:99:ILE:O	3:LE:252:ARG:NH1	2.13	0.82
2:GA:75:GLU:HA	2:GA:78:ARG:HH21	1.44	0.82
2:GG:207:LYS:HG3	2:GG:210:GLN:HG2	1.60	0.81
2:T:102:ASP:HB2	2:T:108:GLU:HB2	1.60	0.81
3:V:122:LEU:HD12	3:V:215:PHE:HB3	1.61	0.81
2:CC:188:LEU:HD13	3:JC:74:ARG:HD2	1.62	0.81
2:EE:75:GLU:HA	2:EE:78:ARG:HH21	1.45	0.81
2:G:188:LEU:HD13	3:C:74:ARG:HD2	1.63	0.81
2:IC:202:ILE:HG13	2:OC:139:ARG:HH22	1.45	0.81
2:AG:17:ILE:HD12	2:AG:21:ARG:HG3	1.61	0.81
2:AG:102:ASP:HB2	2:AG:108:GLU:HB2	1.62	0.81
2:AA:201:GLU:HG3	2:GA:139:ARG:HD2	1.60	0.81
3:JF:122:LEU:HD12	3:JF:215:PHE:HB3	1.63	0.80
3:C:122:LEU:HD12	3:C:215:PHE:HB3	1.63	0.80
3:NA:122:LEU:HD12	3:NA:215:PHE:HB3	1.63	0.80

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:CC:207:LYS:NZ	2:CC:209:GLN:OE1	2.15	0.80
2:KE:69:LEU:O	2:KE:72:ASN:ND2	2.13	0.80
2:B:188:LEU:HD13	3:K:74:ARG:HD2	1.61	0.80
2:T:17:ILE:HD12	2:T:21:ARG:HG3	1.62	0.80
2:YD:21:ARG:HH21	2:YD:24:GLU:HB2	1.45	0.80
2:QE:128:GLN:HG2	3:RE:131:PHE:HB3	1.62	0.80
2:UF:186:GLN:HE22	3:BG:71:ASN:HA	1.46	0.80
2:GG:75:GLU:HA	2:GG:78:ARG:HH21	1.44	0.80
2:OC:69:LEU:HD11	2:UC:47:ILE:HG22	1.64	0.80
2:AD:69:LEU:HD11	2:GD:47:ILE:HG22	1.62	0.80
2:QE:102:ASP:HB2	2:QE:108:GLU:HB2	1.64	0.80
2:WE:75:GLU:HA	2:WE:78:ARG:HH21	1.45	0.80
2:J:45:ARG:HG3	2:J:46:GLN:HG3	1.62	0.80
2:T:186:GLN:HE22	3:BA:71:ASN:HA	1.47	0.80
2:IC:45:ARG:HG3	2:IC:46:GLN:HG3	1.61	0.80
2:B:201:GLU:HB3	2:J:139:ARG:HH21	1.44	0.80
2:YD:45:ARG:HG3	2:YD:46:GLN:HG3	1.62	0.80
2:CF:113:MET:O	2:CF:141:GLN:NE2	2.14	0.80
2:WB:79:SER:O	2:WB:83:LYS:HG2	1.82	0.80
2:QE:21:ARG:HH21	2:QE:24:GLU:HB2	1.45	0.80
4:R:107:ALA:N	3:V:252:ARG:HE	1.79	0.80
2:GD:89:ARG:HH22	2:GD:93:LEU:HG	1.46	0.80
2:SA:69:LEU:HD11	2:YA:47:ILE:HG22	1.64	0.80
2:WB:186:GLN:HE22	3:DC:71:ASN:HA	1.47	0.80
2:KE:102:ASP:HB2	2:KE:108:GLU:HB2	1.62	0.80
2:J:75:GLU:HA	2:J:78:ARG:HH21	1.47	0.79
2:OC:188:LEU:HD13	3:VC:74:ARG:HD2	1.63	0.79
2:MA:188:LEU:HD13	3:TA:74:ARG:HD2	1.64	0.79
2:UC:69:LEU:HD11	2:AD:47:ILE:HG22	1.64	0.79
2:QB:207:LYS:HZ2	2:QB:210:GLN:HG3	1.47	0.79
3:TD:122:LEU:HD12	3:TD:215:PHE:HB3	1.65	0.79
2:MG:188:LEU:HD13	3:TG:74:ARG:HD2	1.65	0.79
2:IF:128:GLN:HE22	2:IF:164:PHE:HD2	1.30	0.79
3:VF:122:LEU:HD12	3:VF:215:PHE:HB3	1.65	0.79
2:QB:188:LEU:HD13	3:XB:74:ARG:HD2	1.63	0.79
2:SD:83:LYS:HE2	2:YD:26:PHE:HB3	1.62	0.79
3:LE:122:LEU:HD12	3:LE:215:PHE:HB3	1.64	0.79
2:QE:69:LEU:HD11	2:WE:47:ILE:HG22	1.65	0.79
2:MD:188:LEU:HD13	3:TD:74:ARG:HD2	1.63	0.79
2:G:35:GLN:NE2	2:SG:96:ASP:OD2	2.15	0.79
2:B:168:GLN:HE22	3:C:130:LEU:HD11	1.48	0.79

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:J:188:LEU:HD13	3:V:74:ARG:HD2	1.64	0.79
3:FE:122:LEU:HD12	3:FE:215:PHE:HB3	1.65	0.79
2:GA:188:LEU:HD13	3:NA:74:ARG:HD2	1.65	0.79
3:RB:120:PRO:HD2	3:RB:121:SER:H	1.48	0.79
2:MG:108:GLU:HA	2:MG:111:ASN:HD22	1.48	0.79
2:EB:65:GLN:NE2	2:KB:47:ILE:O	2.16	0.79
2:AD:188:LEU:HD13	3:HD:74:ARG:HD2	1.64	0.79
4:QD:99:ILE:O	3:TD:252:ARG:NH2	2.16	0.79
2:UF:206:MET:SD	2:UF:207:LYS:N	2.56	0.79
3:PF:120:PRO:HD2	3:PF:121:SER:H	1.47	0.78
2:B:80:VAL:O	2:B:83:LYS:HG2	1.84	0.78
2:T:69:LEU:HD11	2:AA:47:ILE:HG22	1.66	0.78
3:LB:122:LEU:HD12	3:LB:215:PHE:HB3	1.64	0.78
2:QB:107:ILE:HG23	2:QB:137:LEU:HD23	1.65	0.78
2:KE:75:GLU:HA	2:KE:78:ARG:HH21	1.48	0.78
2:KE:207:LYS:HG3	2:KE:210:GLN:HG2	1.64	0.78
2:YA:26:PHE:HE1	2:YA:34:VAL:HG22	1.49	0.78
2:OC:75:GLU:HA	2:OC:78:ARG:HH21	1.48	0.78
3:TD:238:ASN:HB3	3:TD:241:HIS:HD2	1.49	0.78
2:YD:102:ASP:HB2	2:YD:108:GLU:HB2	1.63	0.78
3:BA:122:LEU:HD12	3:BA:215:PHE:HB3	1.66	0.78
3:HG:122:LEU:HD12	3:HG:215:PHE:HB3	1.64	0.78
3:TG:75:ARG:HG3	3:TG:208:ILE:HD11	1.64	0.78
2:SD:120:ASP:OD1	2:SD:123:ARG:NH1	2.17	0.78
2:SD:186:GLN:HE22	3:ZD:71:ASN:HA	1.48	0.78
2:AA:102:ASP:HB2	2:AA:108:GLU:HB2	1.65	0.78
3:JC:122:LEU:HD12	3:JC:215:PHE:HB3	1.65	0.78
2:WE:206:MET:SD	2:WE:207:LYS:N	2.57	0.78
2:J:69:LEU:HD11	2:T:47:ILE:HG22	1.65	0.78
2:GA:69:LEU:HD11	2:MA:47:ILE:HG22	1.66	0.78
3:PC:58:ASN:HD21	3:PC:81:VAL:HG11	1.48	0.78
2:UC:102:ASP:HB2	2:UC:108:GLU:HB2	1.66	0.78
3:ND:97:PRO:O	3:ND:100:THR:OG1	2.02	0.78
2:SG:160:ARG:O	2:SG:164:PHE:HB2	1.83	0.78
2:MA:76:TYR:HE2	2:SA:38:SER:HB3	1.49	0.78
3:XB:122:LEU:HD12	3:XB:215:PHE:HB3	1.66	0.78
2:MD:69:LEU:HD11	2:SD:47:ILE:HG22	1.66	0.78
2:YA:207:LYS:HZ1	2:YA:209:GLN:HB3	1.50	0.77
2:AA:65:GLN:NE2	2:GA:47:ILE:O	2.17	0.77
2:OF:186:GLN:HE21	3:VF:74:ARG:HH22	1.29	0.77
3:RE:122:LEU:HD12	3:RE:215:PHE:HB3	1.67	0.77

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:T:120:ASP:OD1	2:T:123:ARG:NH1	2.18	0.77
3:ZA:53:ALA:HB3	3:ZA:231:LEU:HA	1.64	0.77
3:DF:122:LEU:HD12	3:DF:215:PHE:HB3	1.64	0.77
2:IC:65:GLN:NE2	2:OC:47:ILE:O	2.17	0.77
3:TG:247:ARG:HH12	3:TG:251:VAL:HG23	1.49	0.77
3:RB:97:PRO:O	3:RB:100:THR:OG1	2.03	0.77
3:PC:122:LEU:HD12	3:PC:215:PHE:HB3	1.67	0.77
2:YD:75:GLU:HA	2:YD:78:ARG:HH21	1.50	0.77
2:CF:65:GLN:NE2	2:IF:47:ILE:O	2.17	0.77
3:M:122:LEU:HD12	3:M:215:PHE:HB3	1.67	0.77
2:B:69:LEU:HD21	2:J:47:ILE:HB	1.67	0.77
2:GA:69:LEU:HD21	2:MA:47:ILE:HB	1.67	0.77
2:SA:186:GLN:HE22	3:ZA:71:ASN:HA	1.47	0.77
3:TA:122:LEU:HD12	3:TA:215:PHE:HB3	1.66	0.77
2:WB:17:ILE:HD12	2:WB:21:ARG:HG3	1.67	0.77
2:KB:128:GLN:HE22	2:KB:164:PHE:HD2	1.32	0.77
3:PF:97:PRO:O	3:PF:100:THR:OG1	2.02	0.77
2:UC:45:ARG:HG3	2:UC:46:GLN:HG3	1.67	0.77
2:T:188:LEU:HD13	3:BA:74:ARG:HD2	1.67	0.76
2:EB:76:TYR:HE2	2:KB:38:SER:HB2	1.49	0.76
2:KB:201:GLU:OE2	2:QB:139:ARG:NH2	2.19	0.76
3:LB:236:LEU:HB3	3:LB:238:ASN:HD21	1.50	0.76
2:OC:108:GLU:HA	2:OC:111:ASN:HD22	1.50	0.76
3:BG:162:LEU:HD12	3:BG:177:VAL:HG11	1.68	0.76
3:HG:53:ALA:HB3	3:HG:231:LEU:HA	1.67	0.76
3:K:236:LEU:HB3	3:K:238:ASN:HD21	1.50	0.76
3:TD:186:VAL:HG13	3:TD:187:LYS:HG3	1.65	0.76
2:GG:128:GLN:HE22	2:GG:164:PHE:HD2	1.30	0.76
2:G:26:PHE:CE1	2:G:34:VAL:HG22	2.20	0.76
3:M:53:ALA:HB3	3:M:231:LEU:HA	1.65	0.76
2:T:120:ASP:OD2	3:BA:76:SER:OG	2.02	0.76
2:QB:26:PHE:CE1	2:QB:34:VAL:HG22	2.21	0.76
2:WB:205:LEU:HD22	2:CC:139:ARG:HB3	1.68	0.76
2:EE:26:PHE:HE1	2:EE:34:VAL:HG22	1.51	0.76
2:WE:69:LEU:HD11	2:CF:47:ILE:HG22	1.67	0.76
2:WE:128:GLN:HE22	2:WE:164:PHE:HD2	1.33	0.76
2:CF:26:PHE:HE1	2:CF:34:VAL:HG22	1.49	0.76
2:AA:75:GLU:HA	2:AA:78:ARG:HH21	1.50	0.76
3:HA:122:LEU:HD12	3:HA:215:PHE:HB3	1.66	0.76
2:YD:110:LEU:O	2:YD:141:GLN:NE2	2.16	0.76
3:ZD:122:LEU:HD12	3:ZD:215:PHE:HB3	1.66	0.76

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:LE:186:VAL:HG13	3:LE:187:LYS:HG3	1.67	0.76
2:IF:186:GLN:HB3	3:PF:74:ARG:NH1	2.01	0.76
2:IF:207:LYS:HG3	2:IF:210:GLN:HG2	1.67	0.76
2:IC:120:ASP:OD1	3:PC:76:SER:OG	2.02	0.76
3:BD:53:ALA:HB3	3:BD:231:LEU:HA	1.67	0.76
2:AA:128:GLN:HE22	2:AA:164:PHE:HD2	1.33	0.76
2:YA:160:ARG:O	2:YA:164:PHE:HB2	1.84	0.76
2:WB:76:TYR:HE2	2:CC:38:SER:HB3	1.50	0.76
3:XE:53:ALA:HB3	3:XE:231:LEU:HA	1.68	0.76
2:AG:69:LEU:HD11	2:GG:47:ILE:HG22	1.67	0.76
3:K:122:LEU:HD12	3:K:215:PHE:HB3	1.68	0.76
2:CC:17:ILE:HD12	2:CC:21:ARG:HG3	1.68	0.76
2:YD:186:GLN:NE2	3:FE:71:ASN:HA	2.00	0.76
2:MD:83:LYS:HG2	2:SD:26:PHE:HB3	1.66	0.76
2:CF:75:GLU:HA	2:CF:78:ARG:HH21	1.50	0.76
2:UF:65:GLN:NE2	2:AG:47:ILE:O	2.18	0.76
2:GA:207:LYS:HG3	2:GA:210:GLN:HG2	1.68	0.76
2:YA:205:LEU:HD22	2:EB:139:ARG:HB3	1.68	0.76
2:UC:160:ARG:O	2:UC:164:PHE:HB2	1.85	0.76
3:PF:122:LEU:HD12	3:PF:215:PHE:HB3	1.68	0.76
2:AA:21:ARG:HH21	2:AA:24:GLU:HB2	1.49	0.76
2:MA:219:ARG:NH2	2:MA:227:GLN:OE1	2.18	0.76
2:YA:69:LEU:HD11	2:EB:47:ILE:HG22	1.67	0.76
2:YA:186:GLN:HE22	3:FB:71:ASN:HA	1.51	0.76
2:AD:75:GLU:HA	2:AD:78:ARG:HH21	1.51	0.76
3:NG:53:ALA:HB3	3:NG:231:LEU:HA	1.67	0.76
2:AA:219:ARG:NH2	2:AA:227:GLN:OE1	2.19	0.75
3:HD:122:LEU:HD12	3:HD:215:PHE:HB3	1.68	0.75
2:AA:205:LEU:HD22	2:GA:139:ARG:HB3	1.67	0.75
2:MA:207:LYS:HG3	2:MA:210:GLN:HG2	1.69	0.75
3:JF:106:HIS:CE1	3:JF:113:THR:HG22	2.21	0.75
2:GG:186:GLN:HE22	3:NG:71:ASN:HA	1.50	0.75
2:T:26:PHE:HE1	2:T:34:VAL:HG22	1.50	0.75
2:T:80:VAL:HA	2:T:83:LYS:HD3	1.67	0.75
3:JC:53:ALA:HB3	3:JC:231:LEU:HA	1.66	0.75
3:PC:97:PRO:O	3:PC:100:THR:OG1	2.05	0.75
2:SD:22:ALA:O	2:SD:26:PHE:HB2	1.85	0.75
2:QE:188:LEU:HD23	2:QE:188:LEU:H	1.52	0.75
3:JF:102:LEU:HD22	3:JF:117:VAL:HG22	1.66	0.75
3:FB:122:LEU:HD12	3:FB:215:PHE:HB3	1.68	0.75
3:BD:122:LEU:HD12	3:BD:215:PHE:HB3	1.68	0.75

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:MD:89:ARG:HH22	2:MD:93:LEU:HG	1.52	0.75
2:KE:79:SER:O	2:KE:83:LYS:HG2	1.86	0.75
2:QB:75:GLU:HA	2:QB:78:ARG:HH21	1.51	0.75
3:ND:186:VAL:HG13	3:ND:187:LYS:HG3	1.67	0.75
3:PF:112:GLY:HA3	3:PF:223:MET:HG2	1.67	0.75
2:GG:219:ARG:NH2	2:GG:227:GLN:OE1	2.19	0.75
2:J:120:ASP:OD1	2:J:123:ARG:NH1	2.20	0.75
3:K:97:PRO:O	3:K:100:THR:OG1	2.03	0.75
3:PC:186:VAL:HG13	3:PC:187:LYS:HG3	1.68	0.75
2:WB:89:ARG:HG2	2:WB:90:ALA:N	1.99	0.75
2:OC:128:GLN:HG2	3:PC:131:PHE:HB3	1.67	0.75
2:B:96:ASP:OD2	2:J:35:GLN:NE2	2.19	0.74
2:AA:186:GLN:NE2	3:HA:71:ASN:HA	2.02	0.74
2:IC:21:ARG:HH21	2:IC:24:GLU:HB2	1.51	0.74
2:UF:89:ARG:HG2	2:UF:90:ALA:N	2.01	0.74
2:G:89:ARG:HG2	2:G:90:ALA:N	2.02	0.74
2:GD:201:GLU:HB3	2:MD:139:ARG:HH21	1.52	0.74
2:SD:79:SER:O	2:SD:83:LYS:HG2	1.87	0.74
2:SD:108:GLU:HA	2:SD:111:ASN:HD22	1.52	0.74
2:GG:65:GLN:NE2	2:MG:47:ILE:O	2.19	0.74
2:SG:45:ARG:HG3	2:SG:46:GLN:HG3	1.69	0.74
3:LB:102:LEU:HD22	3:LB:117:VAL:HG22	1.68	0.74
2:G:75:GLU:HA	2:G:78:ARG:HH21	1.53	0.74
3:HD:97:PRO:O	3:HD:100:THR:OG1	2.06	0.74
4:SE:100:LEU:HA	4:SE:105:LEU:HA	1.70	0.74
2:J:207:LYS:HG3	2:J:210:GLN:HG2	1.68	0.74
2:AA:160:ARG:O	2:AA:164:PHE:HB2	1.86	0.74
3:XB:106:HIS:CE1	3:XB:113:THR:HG22	2.21	0.74
2:WE:80:VAL:HG12	2:CF:26:PHE:HE2	1.53	0.74
2:AD:207:LYS:HZ2	2:AD:210:GLN:HG3	1.53	0.74
2:EE:160:ARG:O	2:EE:164:PHE:HB2	1.87	0.74
2:AA:111:ASN:OD1	2:AA:141:GLN:NE2	2.20	0.74
2:GA:186:GLN:HE22	3:NA:71:ASN:HA	1.51	0.74
2:EB:201:GLU:HB3	2:KB:139:ARG:NH2	2.02	0.74
3:NG:122:LEU:HD12	3:NG:215:PHE:HB3	1.69	0.74
2:AA:17:ILE:HD12	2:AA:21:ARG:HG3	1.70	0.74
2:YD:17:ILE:HD12	2:YD:21:ARG:HG3	1.70	0.74
2:OF:45:ARG:HG3	2:OF:46:GLN:HG3	1.69	0.74
2:OF:75:GLU:O	2:OF:79:SER:HB3	1.86	0.74
2:GA:96:ASP:OD2	2:MA:35:GLN:NE2	2.20	0.74
2:IC:219:ARG:NH2	2:IC:227:GLN:OE1	2.20	0.74

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:PC:53:ALA:HB3	3:PC:231:LEU:HA	1.69	0.74
2:YA:80:VAL:HA	2:YA:83:LYS:HD3	1.70	0.74
2:EB:96:ASP:OD2	2:KB:35:GLN:NE2	2.21	0.74
3:VC:53:ALA:HB3	3:VC:231:LEU:HA	1.67	0.74
4:OD:100:LEU:HA	4:OD:105:LEU:HA	1.70	0.74
2:YD:65:GLN:NE2	2:EE:47:ILE:O	2.20	0.74
3:V:105:ILE:HG12	3:V:162:LEU:HD11	1.70	0.73
2:MA:128:GLN:HE22	2:MA:164:PHE:HD2	1.35	0.73
2:KE:45:ARG:HG3	2:KE:46:GLN:HG3	1.68	0.73
2:J:17:ILE:HD12	2:J:21:ARG:HG3	1.69	0.73
3:VC:122:LEU:HD12	3:VC:215:PHE:HB3	1.70	0.73
2:WE:186:GLN:HE21	3:DF:71:ASN:HA	1.53	0.73
2:GA:120:ASP:OD1	3:NA:76:SER:OG	2.06	0.73
2:SA:207:LYS:NZ	2:SA:209:GLN:OE1	2.18	0.73
3:ZA:122:LEU:HD12	3:ZA:215:PHE:HB3	1.71	0.73
2:EB:80:VAL:HA	2:EB:83:LYS:HD3	1.70	0.73
3:HD:238:ASN:HB3	3:HD:241:HIS:HD2	1.54	0.73
2:SA:45:ARG:HG3	2:SA:46:GLN:HG3	1.69	0.73
2:EB:310:LEU:HD12	2:EB:311:ILE:HD13	1.71	0.73
2:SD:89:ARG:HH22	2:SD:93:LEU:HD21	1.54	0.73
2:YD:219:ARG:NH2	2:YD:227:GLN:OE1	2.20	0.73
2:OF:108:GLU:HA	2:OF:111:ASN:HD22	1.52	0.73
2:QB:17:ILE:HD12	2:QB:21:ARG:HG3	1.71	0.73
4:YE:100:LEU:HA	4:YE:105:LEU:HA	1.71	0.73
2:G:160:ARG:O	2:G:164:PHE:HB2	1.89	0.73
2:AA:45:ARG:HG3	2:AA:46:GLN:HG3	1.71	0.73
2:CC:186:GLN:HE22	3:JC:71:ASN:HA	1.54	0.73
2:UC:96:ASP:OD2	2:AD:35:GLN:NE2	2.21	0.73
3:RB:122:LEU:HD12	3:RB:215:PHE:HB3	1.70	0.73
2:B:128:GLN:HE22	2:B:164:PHE:HD2	1.35	0.73
2:QB:205:LEU:HD21	2:WB:140:SER:HA	1.71	0.73
2:YD:160:ARG:O	2:YD:164:PHE:HB2	1.89	0.73
2:CF:210:GLN:HE22	2:IF:147:ALA:HB1	1.53	0.73
2:SG:118:ALA:HA	2:SG:121:LEU:HG	1.71	0.73
2:B:152:ARG:HH22	2:B:153:LEU:HB2	1.54	0.72
2:EE:128:GLN:HE22	2:EE:164:PHE:HD2	1.37	0.72
2:CC:128:GLN:HE22	2:CC:164:PHE:HD2	1.36	0.72
2:YD:21:ARG:NH2	2:YD:24:GLU:HB2	2.03	0.72
4:AB:100:LEU:HA	4:AB:105:LEU:HA	1.70	0.72
2:MD:45:ARG:HG3	2:MD:46:GLN:HG3	1.71	0.72
2:SD:186:GLN:HB3	3:ZD:74:ARG:NH1	2.03	0.72

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:EE:201:GLU:HG3	2:KE:139:ARG:HD2	1.71	0.72
2:MA:83:LYS:HG2	2:SA:26:PHE:HB3	1.72	0.72
2:SD:206:MET:SD	2:SD:207:LYS:N	2.62	0.72
2:YD:69:LEU:HD11	2:EE:47:ILE:HG22	1.71	0.72
2:CF:102:ASP:HB2	2:CF:108:GLU:HB2	1.71	0.72
2:IF:158:MET:HE3	2:IF:159:LEU:HD22	1.70	0.72
2:GA:160:ARG:O	2:GA:164:PHE:HB2	1.88	0.72
2:WB:21:ARG:O	2:WB:21:ARG:NE	2.19	0.72
2:UC:75:GLU:HA	2:UC:78:ARG:HH21	1.54	0.72
2:YA:219:ARG:NH2	2:YA:227:GLN:OE1	2.23	0.72
4:WC:100:LEU:HA	4:WC:105:LEU:HA	1.71	0.72
2:GD:128:GLN:HE22	2:GD:164:PHE:HD2	1.35	0.72
2:KE:160:ARG:O	2:KE:164:PHE:HB2	1.90	0.72
2:OF:160:ARG:O	2:OF:164:PHE:HB2	1.88	0.72
2:EB:89:ARG:HG2	2:EB:90:ALA:N	2.02	0.72
2:OC:201:GLU:HG2	2:UC:139:ARG:HD2	1.71	0.72
2:YD:96:ASP:OD2	2:EE:35:GLN:NE2	2.22	0.72
2:EE:89:ARG:HG2	2:EE:90:ALA:N	2.04	0.72
2:QE:160:ARG:O	2:QE:164:PHE:HB2	1.90	0.72
2:CF:89:ARG:HH22	2:CF:93:LEU:HG	1.54	0.72
2:OF:96:ASP:OD2	2:UF:35:GLN:NE2	2.23	0.72
2:MA:310:LEU:HD12	2:MA:311:ILE:HD13	1.72	0.72
4:SB:100:LEU:HA	4:SB:105:LEU:HA	1.72	0.72
4:YC:99:ILE:O	3:BD:252:ARG:NH2	2.23	0.72
2:GD:69:LEU:HD21	2:MD:47:ILE:HB	1.70	0.72
2:J:69:LEU:HD21	2:T:47:ILE:HB	1.72	0.72
2:MA:168:GLN:HE22	3:NA:130:LEU:HD11	1.54	0.72
2:YA:287:ARG:HA	2:YA:290:LEU:HD12	1.72	0.72
2:YA:310:LEU:HD12	2:YA:311:ILE:HD13	1.72	0.72
4:GB:100:LEU:HA	4:GB:105:LEU:HA	1.71	0.72
3:DC:162:LEU:HD12	3:DC:177:VAL:HG11	1.71	0.72
2:GA:310:LEU:HD12	2:GA:311:ILE:HD13	1.72	0.72
2:WB:310:LEU:HD12	2:WB:311:ILE:HD13	1.72	0.72
2:AA:310:LEU:HD12	2:AA:311:ILE:HD13	1.72	0.71
2:SA:310:LEU:HD12	2:SA:311:ILE:HD13	1.72	0.71
2:GD:65:GLN:NE2	2:MD:47:ILE:O	2.22	0.71
3:LE:53:ALA:HB3	3:LE:231:LEU:HA	1.72	0.71
2:IF:201:GLU:OE2	2:OF:139:ARG:NH2	2.23	0.71
2:OF:219:ARG:NH2	2:OF:227:GLN:OE1	2.23	0.71
2:UF:120:ASP:OD1	2:UF:123:ARG:NH1	2.23	0.71
3:HG:186:VAL:HG13	3:HG:187:LYS:HG3	1.71	0.71

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:C:53:ALA:HB3	3:C:231:LEU:HA	1.70	0.71
2:T:310:LEU:HD12	2:T:311:ILE:HD13	1.72	0.71
4:YB:100:LEU:HA	4:YB:105:LEU:HA	1.72	0.71
2:IC:310:LEU:HD12	2:IC:311:ILE:HD13	1.72	0.71
2:OC:310:LEU:HD12	2:OC:311:ILE:HD13	1.72	0.71
2:MD:160:ARG:O	2:MD:164:PHE:HB2	1.89	0.71
2:WE:96:ASP:OD2	2:CF:35:GLN:NE2	2.23	0.71
2:OF:17:ILE:HD12	2:OF:21:ARG:HG3	1.71	0.71
2:OF:97:ILE:O	2:OF:101:ARG:NH2	2.21	0.71
3:PF:53:ALA:HB3	3:PF:231:LEU:HA	1.70	0.71
4:QF:100:LEU:HA	4:QF:105:LEU:HA	1.72	0.71
2:GG:69:LEU:HD21	2:MG:47:ILE:HB	1.73	0.71
2:GG:96:ASP:OD2	2:MG:35:GLN:NE2	2.23	0.71
2:T:107:ILE:HG22	2:T:111:ASN:HD21	1.54	0.71
2:KB:310:LEU:HD12	2:KB:311:ILE:HD13	1.73	0.71
2:QB:287:ARG:HA	2:QB:290:LEU:HD12	1.73	0.71
2:CC:310:LEU:HD12	2:CC:311:ILE:HD13	1.72	0.71
2:IC:287:ARG:HA	2:IC:290:LEU:HD12	1.73	0.71
2:KE:96:ASP:OD2	2:QE:35:GLN:NE2	2.23	0.71
2:AG:96:ASP:OD2	2:GG:35:GLN:NE2	2.22	0.71
2:AG:287:ARG:HA	2:AG:290:LEU:HD12	1.73	0.71
3:TG:122:LEU:HD12	3:TG:215:PHE:HB3	1.70	0.71
2:B:310:LEU:HD12	2:B:311:ILE:HD13	1.72	0.71
2:J:310:LEU:HD12	2:J:311:ILE:HD13	1.72	0.71
2:KB:186:GLN:HE22	3:RB:71:ASN:HA	1.55	0.71
2:QB:74:ASN:O	2:QB:78:ARG:NE	2.23	0.71
2:IC:107:ILE:HD13	2:IC:136:HIS:HB2	1.70	0.71
2:UC:310:LEU:HD12	2:UC:311:ILE:HD13	1.72	0.71
4:YC:107:ALA:N	3:BD:252:ARG:HE	1.86	0.71
2:YA:89:ARG:HG2	2:YA:90:ALA:N	2.03	0.71
2:AD:310:LEU:HD12	2:AD:311:ILE:HD13	1.72	0.71
2:EE:102:ASP:HB2	2:EE:108:GLU:HB3	1.72	0.71
2:KE:168:GLN:HE22	3:LE:130:LEU:HD11	1.54	0.71
3:XE:122:LEU:HD12	3:XE:215:PHE:HB3	1.70	0.71
2:OF:234:PHE:HB3	2:OF:313:ARG:HH21	1.55	0.71
4:CG:100:LEU:HA	4:CG:105:LEU:HA	1.72	0.71
3:NG:97:PRO:O	3:NG:100:THR:OG1	2.07	0.71
2:SG:287:ARG:HA	2:SG:290:LEU:HD12	1.73	0.71
4:W:100:LEU:HA	4:W:105:LEU:HA	1.71	0.71
2:MA:107:ILE:HG22	2:MA:111:ASN:HD21	1.55	0.71
4:EC:100:LEU:HA	4:EC:105:LEU:HA	1.72	0.71

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:IC:234:PHE:HB3	2:IC:313:ARG:HH21	1.56	0.71
2:UC:118:ALA:HA	2:UC:121:LEU:HG	1.70	0.71
2:YD:69:LEU:HD21	2:EE:47:ILE:HB	1.73	0.71
2:AG:128:GLN:HE22	2:AG:164:PHE:HD2	1.35	0.71
2:J:152:ARG:HH22	2:J:153:LEU:HB2	1.56	0.71
2:GA:287:ARG:HA	2:GA:290:LEU:HD12	1.73	0.71
2:SA:89:ARG:HG2	2:SA:90:ALA:N	2.03	0.71
2:EB:186:GLN:HB3	3:LB:74:ARG:HH12	1.55	0.71
2:WB:128:GLN:HE22	2:WB:164:PHE:HD2	1.38	0.71
2:IF:287:ARG:HA	2:IF:290:LEU:HD12	1.73	0.71
3:TG:105:ILE:HG12	3:TG:162:LEU:HD11	1.72	0.71
2:AD:287:ARG:HA	2:AD:290:LEU:HD12	1.73	0.71
2:SD:310:LEU:HD12	2:SD:311:ILE:HD13	1.72	0.71
2:MG:160:ARG:O	2:MG:164:PHE:HB2	1.90	0.71
2:J:21:ARG:O	2:J:21:ARG:NE	2.16	0.71
2:J:287:ARG:HA	2:J:290:LEU:HD12	1.73	0.71
2:MD:310:LEU:HD12	2:MD:311:ILE:HD13	1.73	0.71
2:SD:45:ARG:HG3	2:SD:46:GLN:HG3	1.73	0.71
4:ME:100:LEU:HA	4:ME:105:LEU:HA	1.73	0.71
2:CF:151:GLU:HA	2:CF:154:ARG:HG2	1.71	0.71
2:GG:188:LEU:HD13	3:NG:74:ARG:HG2	1.71	0.71
2:KB:120:ASP:OD1	3:RB:76:SER:OG	2.08	0.71
2:GD:17:ILE:HD12	2:GD:21:ARG:HG3	1.73	0.71
2:OF:207:LYS:HZ2	2:OF:210:GLN:HG3	1.56	0.71
2:GG:310:LEU:HD12	2:GG:311:ILE:HD13	1.72	0.71
2:MG:21:ARG:HH21	2:MG:24:GLU:HB2	1.56	0.71
2:YA:207:LYS:HG3	2:YA:210:GLN:HG2	1.70	0.70
2:EB:21:ARG:HH21	2:EB:24:GLU:HB2	1.56	0.70
2:WB:287:ARG:HA	2:WB:290:LEU:HD12	1.73	0.70
4:CD:100:LEU:HA	4:CD:105:LEU:HA	1.73	0.70
1:RD:547:VAL:HA	2:SD:28:HIS:CE1	2.26	0.70
2:QE:120:ASP:OD1	3:XE:76:SER:OG	2.09	0.70
2:QE:130:ILE:O	2:QE:134:LEU:HG	1.91	0.70
4:WF:100:LEU:HA	4:WF:105:LEU:HA	1.73	0.70
2:B:287:ARG:HA	2:B:290:LEU:HD12	1.73	0.70
2:AA:287:ARG:HA	2:AA:290:LEU:HD12	1.73	0.70
3:BA:105:ILE:HG12	3:BA:162:LEU:HD11	1.71	0.70
4:OA:100:LEU:HA	4:OA:105:LEU:HA	1.73	0.70
2:CC:287:ARG:HA	2:CC:290:LEU:HD12	1.72	0.70
2:SD:234:PHE:HB3	2:SD:313:ARG:HH21	1.56	0.70
2:YD:310:LEU:HD12	2:YD:311:ILE:HD13	1.72	0.70

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:EE:310:LEU:HD12	2:EE:311:ILE:HD13	1.72	0.70
3:FE:102:LEU:HD22	3:FE:117:VAL:HG22	1.73	0.70
2:KE:310:LEU:HD12	2:KE:311:ILE:HD13	1.72	0.70
2:QE:120:ASP:OD2	2:QE:123:ARG:NH1	2.24	0.70
2:QE:310:LEU:HD12	2:QE:311:ILE:HD13	1.72	0.70
2:WE:287:ARG:HA	2:WE:290:LEU:HD12	1.73	0.70
2:UF:127:PRO:HA	2:UF:130:ILE:HG12	1.73	0.70
2:GG:108:GLU:HA	2:GG:111:ASN:HD22	1.56	0.70
2:T:26:PHE:CE1	2:T:34:VAL:HG22	2.26	0.70
2:MA:80:VAL:HA	2:MA:83:LYS:HD3	1.72	0.70
2:YA:128:GLN:HE22	2:YA:164:PHE:HD2	1.37	0.70
2:KB:96:ASP:OD2	2:QB:35:GLN:NE2	2.24	0.70
2:MD:127:PRO:HA	2:MD:130:ILE:HG12	1.74	0.70
2:CF:310:LEU:HD12	2:CF:311:ILE:HD13	1.72	0.70
2:IF:96:ASP:OD2	2:OF:35:GLN:NE2	2.23	0.70
2:IF:310:LEU:HD12	2:IF:311:ILE:HD13	1.72	0.70
2:AG:310:LEU:HD12	2:AG:311:ILE:HD13	1.73	0.70
2:GG:234:PHE:HB3	2:GG:313:ARG:HH21	1.54	0.70
3:C:105:ILE:HG12	3:C:162:LEU:HD11	1.74	0.70
2:SA:287:ARG:HA	2:SA:290:LEU:HD12	1.73	0.70
2:EB:127:PRO:HA	2:EB:130:ILE:HG12	1.71	0.70
2:EB:287:ARG:HA	2:EB:290:LEU:HD12	1.73	0.70
2:KB:287:ARG:HA	2:KB:290:LEU:HD12	1.73	0.70
3:DC:122:LEU:HD12	3:DC:215:PHE:HB3	1.73	0.70
3:VC:105:ILE:HG12	3:VC:162:LEU:HD11	1.72	0.70
2:EE:108:GLU:HA	2:EE:111:ASN:HD22	1.56	0.70
2:EE:207:LYS:HG3	2:EE:210:GLN:HG2	1.73	0.70
2:QE:287:ARG:HA	2:QE:290:LEU:HD12	1.73	0.70
2:WE:310:LEU:HD12	2:WE:311:ILE:HD13	1.72	0.70
2:UF:287:ARG:HA	2:UF:290:LEU:HD12	1.73	0.70
2:GG:287:ARG:HA	2:GG:290:LEU:HD12	1.72	0.70
2:MG:128:GLN:HE22	2:MG:164:PHE:HD2	1.39	0.70
2:G:287:ARG:HA	2:G:290:LEU:HD12	1.73	0.70
3:M:74:ARG:HH11	2:SG:188:LEU:HD21	1.57	0.70
2:MA:287:ARG:HA	2:MA:290:LEU:HD12	1.73	0.70
2:EB:128:GLN:HE22	2:EB:164:PHE:HD2	1.38	0.70
3:RB:190:ASN:ND2	3:XB:234:PRO:O	2.24	0.70
2:UC:287:ARG:HA	2:UC:290:LEU:HD12	1.73	0.70
2:SD:287:ARG:HA	2:SD:290:LEU:HD12	1.73	0.70
3:ZD:53:ALA:HB3	3:ZD:231:LEU:HA	1.73	0.70
2:EE:287:ARG:HA	2:EE:290:LEU:HD12	1.73	0.70

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:B:89:ARG:HH22	2:B:93:LEU:HG	1.55	0.70
3:DC:53:ALA:HB3	3:DC:231:LEU:HA	1.73	0.70
2:YD:205:LEU:HD22	2:EE:139:ARG:HB3	1.71	0.70
2:OF:287:ARG:HA	2:OF:290:LEU:HD12	1.73	0.70
2:UF:69:LEU:HD21	2:AG:47:ILE:HB	1.74	0.70
3:BG:122:LEU:HD12	3:BG:215:PHE:HB3	1.72	0.70
2:GG:207:LYS:HZ1	2:GG:209:GLN:HB3	1.54	0.70
2:MA:97:ILE:O	2:MA:101:ARG:NH2	2.21	0.70
2:IC:128:GLN:HE22	2:IC:164:PHE:HD2	1.35	0.70
3:JC:186:VAL:HG13	3:JC:187:LYS:HG3	1.72	0.70
2:MD:287:ARG:HA	2:MD:290:LEU:HD12	1.73	0.70
3:DF:105:ILE:HG12	3:DF:162:LEU:HD11	1.74	0.70
2:J:234:PHE:HB3	2:J:313:ARG:HH21	1.55	0.70
2:MD:26:PHE:CE1	2:MD:34:VAL:HG22	2.27	0.70
3:ND:105:ILE:HG12	3:ND:162:LEU:HD11	1.73	0.70
3:ND:238:ASN:HB3	3:ND:241:HIS:HD2	1.57	0.70
4:UD:100:LEU:HA	4:UD:105:LEU:HA	1.73	0.70
2:KE:107:ILE:HG23	2:KE:137:LEU:HD23	1.72	0.70
2:QE:207:LYS:NZ	2:QE:209:GLN:OE1	2.18	0.70
2:J:80:VAL:O	2:J:83:LYS:HB3	1.92	0.70
2:T:160:ARG:O	2:T:164:PHE:HB2	1.91	0.70
2:EB:75:GLU:HA	2:EB:78:ARG:HH21	1.56	0.70
2:KB:69:LEU:HD23	2:QB:47:ILE:HG22	1.74	0.70
2:KB:160:ARG:O	2:KB:164:PHE:HB2	1.92	0.70
2:OC:287:ARG:HA	2:OC:290:LEU:HD12	1.73	0.70
2:SD:128:GLN:HE22	2:SD:164:PHE:HD2	1.39	0.70
3:FE:183:GLU:HG3	3:FE:188:PHE:HB2	1.74	0.70
2:WE:160:ARG:O	2:WE:164:PHE:HB2	1.92	0.70
4:EF:100:LEU:HA	4:EF:105:LEU:HA	1.72	0.70
2:GG:89:ARG:HH22	2:GG:93:LEU:HG	1.56	0.70
3:C:102:LEU:HD22	3:C:117:VAL:HG22	1.73	0.70
2:J:65:GLN:NE2	2:T:47:ILE:O	2.23	0.70
2:AA:75:GLU:O	2:AA:79:SER:HB3	1.91	0.70
4:IA:100:LEU:HA	4:IA:105:LEU:HA	1.74	0.70
2:KB:74:ASN:O	2:KB:78:ARG:NE	2.24	0.70
2:KB:127:PRO:HA	2:KB:130:ILE:HD12	1.74	0.70
2:IC:21:ARG:NH2	2:IC:24:GLU:HB2	2.07	0.70
3:ND:240:ARG:NH1	3:ND:240:ARG:HA	2.07	0.70
4:UG:100:LEU:HA	4:UG:105:LEU:HA	1.73	0.70
2:WB:65:GLN:NE2	2:CC:47:ILE:O	2.20	0.69
4:ID:100:LEU:HA	4:ID:105:LEU:HA	1.73	0.69

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:YD:287:ARG:HA	2:YD:290:LEU:HD12	1.73	0.69
2:IF:120:ASP:OD1	3:PF:76:SER:OG	2.10	0.69
2:UF:130:ILE:O	2:UF:134:LEU:HG	1.92	0.69
2:SG:21:ARG:HH21	2:SG:24:GLU:HB2	1.57	0.69
2:B:65:GLN:NE2	2:J:47:ILE:O	2.24	0.69
2:T:89:ARG:HH22	2:T:93:LEU:HG	1.57	0.69
2:YA:21:ARG:HH21	2:YA:24:GLU:HB2	1.58	0.69
2:UF:207:LYS:HZ2	2:UF:210:GLN:HB2	1.57	0.69
4:N:100:LEU:HA	4:N:105:LEU:HA	1.74	0.69
2:T:76:TYR:OH	2:AA:34:VAL:HG13	1.92	0.69
2:MA:45:ARG:HG3	2:MA:46:GLN:HG3	1.74	0.69
2:MA:89:ARG:HD2	2:MA:90:ALA:N	2.06	0.69
2:SA:128:GLN:HE22	2:SA:164:PHE:HD2	1.37	0.69
4:UA:100:LEU:HA	4:UA:105:LEU:HA	1.74	0.69
4:MB:100:LEU:HA	4:MB:105:LEU:HA	1.75	0.69
2:WB:107:ILE:HG23	2:WB:137:LEU:HD23	1.74	0.69
2:WB:160:ARG:O	2:WB:164:PHE:HB2	1.92	0.69
4:QC:100:LEU:HA	4:QC:105:LEU:HA	1.74	0.69
2:SD:160:ARG:O	2:SD:164:PHE:HB2	1.91	0.69
2:WE:310:LEU:HA	2:WE:313:ARG:HG2	1.75	0.69
2:T:287:ARG:HA	2:T:290:LEU:HD12	1.73	0.69
3:PC:105:ILE:HG12	3:PC:162:LEU:HD11	1.74	0.69
2:IC:120:ASP:OD2	2:IC:123:ARG:NH1	2.26	0.69
2:YD:89:ARG:HD2	2:YD:90:ALA:N	2.06	0.69
2:YD:120:ASP:OD1	2:YD:123:ARG:NH1	2.26	0.69
2:MG:287:ARG:HA	2:MG:290:LEU:HD12	1.74	0.69
3:TG:72:LEU:HD23	3:TG:73:LEU:HD12	1.74	0.69
2:IC:310:LEU:HA	2:IC:313:ARG:HG2	1.75	0.69
2:OC:186:GLN:HB3	3:VC:74:ARG:HH12	1.58	0.69
2:YD:26:PHE:HZ	2:YD:34:VAL:HA	1.58	0.69
4:IE:106:ILE:H	3:LE:252:ARG:NH2	1.91	0.69
2:KE:287:ARG:HA	2:KE:290:LEU:HD12	1.73	0.69
2:QE:96:ASP:OD2	2:WE:35:GLN:NE2	2.24	0.69
2:OF:129:ILE:HG23	2:OF:175:LEU:HD21	1.75	0.69
4:OG:100:LEU:HA	4:OG:105:LEU:HA	1.74	0.69
4:CA:100:LEU:HA	4:CA:105:LEU:HA	1.74	0.69
2:EB:186:GLN:HB3	3:LB:74:ARG:NH1	2.07	0.69
2:SD:310:LEU:HA	2:SD:313:ARG:HG2	1.75	0.69
2:GG:120:ASP:OD2	3:NG:76:SER:OG	2.11	0.69
2:GG:160:ARG:O	2:GG:164:PHE:HB2	1.93	0.69
3:C:127:VAL:O	3:C:131:PHE:HB2	1.93	0.69

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:AA:96:ASP:OD2	2:GA:35:GLN:NE2	2.26	0.69
2:SA:120:ASP:OD2	3:ZA:76:SER:OG	2.02	0.69
2:QB:89:ARG:HH22	2:QB:93:LEU:HD21	1.58	0.69
2:IC:69:LEU:HD21	2:OC:47:ILE:HB	1.74	0.69
2:OC:310:LEU:HA	2:OC:313:ARG:HG2	1.75	0.69
3:PC:57:ILE:HD11	3:PC:169:TRP:HA	1.75	0.69
2:UC:205:LEU:HD13	2:AD:139:ARG:HB3	1.75	0.69
2:GD:97:ILE:O	2:GD:101:ARG:NH2	2.21	0.69
2:GD:129:ILE:HG23	2:GD:175:LEU:HD21	1.75	0.69
3:ND:75:ARG:HG3	3:ND:208:ILE:HD11	1.72	0.69
4:AE:100:LEU:HA	4:AE:105:LEU:HA	1.74	0.69
2:CF:310:LEU:HA	2:CF:313:ARG:HG2	1.75	0.69
2:AG:207:LYS:HZ1	2:AG:209:GLN:HB3	1.58	0.69
3:BG:53:ALA:HB3	3:BG:231:LEU:HA	1.73	0.69
2:GG:120:ASP:OD1	2:GG:123:ARG:NH1	2.25	0.69
2:MG:96:ASP:OD2	2:SG:35:GLN:NE2	2.26	0.69
2:SG:137:LEU:HB3	2:SG:141:GLN:HE21	1.56	0.69
2:EB:310:LEU:HA	2:EB:313:ARG:HG2	1.75	0.69
2:QB:234:PHE:HB3	2:QB:313:ARG:HH21	1.56	0.69
2:GD:287:ARG:HA	2:GD:290:LEU:HD12	1.73	0.69
2:KE:65:GLN:NE2	2:QE:47:ILE:O	2.26	0.69
2:OF:79:SER:O	2:OF:83:LYS:HE3	1.93	0.69
2:GA:129:ILE:HG23	2:GA:175:LEU:HD21	1.74	0.69
2:EB:83:LYS:HD2	2:KB:26:PHE:HD1	1.58	0.69
2:MD:310:LEU:HA	2:MD:313:ARG:HG2	1.75	0.69
2:QE:127:PRO:HA	2:QE:130:ILE:HG12	1.75	0.69
2:WE:69:LEU:HD21	2:CF:47:ILE:HB	1.75	0.69
2:CF:152:ARG:HH12	2:CF:153:LEU:HD13	1.56	0.69
2:CF:287:ARG:HA	2:CF:290:LEU:HD12	1.74	0.69
2:OF:107:ILE:HG23	2:OF:137:LEU:HD23	1.74	0.69
2:GG:310:LEU:HA	2:GG:313:ARG:HG2	1.75	0.69
3:NG:105:ILE:HG12	3:NG:162:LEU:HD11	1.74	0.69
2:J:96:ASP:OD2	2:T:35:GLN:NE2	2.25	0.68
2:J:201:GLU:O	2:T:139:ARG:NH2	2.26	0.68
2:AA:69:LEU:HD21	2:GA:47:ILE:HB	1.74	0.68
3:HA:90:HIS:O	3:HA:94:ARG:HG2	1.93	0.68
3:HA:112:GLY:HA3	3:HA:223:MET:HG2	1.75	0.68
2:QB:107:ILE:HD13	2:QB:136:HIS:HB2	1.73	0.68
2:QB:202:ILE:HD13	2:WB:139:ARG:NH1	2.08	0.68
3:VC:112:GLY:HA3	3:VC:223:MET:HG2	1.74	0.68
2:AD:206:MET:SD	2:AD:207:LYS:N	2.66	0.68

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:AD:219:ARG:NH2	2:AD:227:GLN:OE1	2.25	0.68
3:ND:122:LEU:HD12	3:ND:215:PHE:HB3	1.72	0.68
2:YD:310:LEU:HA	2:YD:313:ARG:HG2	1.75	0.68
2:EE:160:ARG:O	2:EE:164:PHE:CB	2.42	0.68
3:PF:57:ILE:HD11	3:PF:169:TRP:HA	1.74	0.68
2:AG:310:LEU:HA	2:AG:313:ARG:HG2	1.75	0.68
2:GG:69:LEU:HD11	2:MG:47:ILE:HG22	1.75	0.68
3:NG:54:LEU:HB2	3:NG:231:LEU:HD23	1.75	0.68
2:SG:129:ILE:HG23	2:SG:175:LEU:HD21	1.75	0.68
4:L:100:LEU:HA	4:L:105:LEU:HA	1.74	0.68
2:AA:21:ARG:NH2	2:AA:24:GLU:HB2	2.07	0.68
2:AA:186:GLN:HB3	3:HA:74:ARG:NH1	2.04	0.68
2:SA:96:ASP:OD2	2:YA:35:GLN:NE2	2.27	0.68
2:YA:310:LEU:HA	2:YA:313:ARG:HG2	1.75	0.68
2:UC:310:LEU:HA	2:UC:313:ARG:HG2	1.75	0.68
2:MD:110:LEU:O	2:MD:141:GLN:NE2	2.25	0.68
3:TD:105:ILE:HG12	3:TD:162:LEU:HD11	1.74	0.68
2:QE:310:LEU:HA	2:QE:313:ARG:HG2	1.75	0.68
3:HG:74:ARG:HB2	3:HG:74:ARG:NH1	2.09	0.68
3:NA:53:ALA:HB3	3:NA:231:LEU:HA	1.73	0.68
2:WB:79:SER:OG	2:WB:83:LYS:NZ	2.27	0.68
2:CC:234:PHE:HB3	2:CC:313:ARG:HH21	1.58	0.68
2:CC:310:LEU:HA	2:CC:313:ARG:HG2	1.76	0.68
2:OC:120:ASP:OD1	3:VC:76:SER:OG	2.11	0.68
2:QE:107:ILE:HG23	2:QE:137:LEU:HD23	1.75	0.68
3:PF:120:PRO:HD2	3:PF:121:SER:N	2.08	0.68
3:VF:53:ALA:HB3	3:VF:231:LEU:HA	1.75	0.68
2:WB:127:PRO:HA	2:WB:130:ILE:HG12	1.75	0.68
2:CC:21:ARG:NH2	2:CC:24:GLU:HB2	2.08	0.68
2:T:310:LEU:HA	2:T:313:ARG:HG2	1.75	0.68
2:AA:129:ILE:HG23	2:AA:175:LEU:HD21	1.75	0.68
3:HA:105:ILE:HG12	3:HA:162:LEU:HD11	1.75	0.68
3:FB:105:ILE:HG12	3:FB:162:LEU:HD11	1.75	0.68
2:SD:107:ILE:HG23	2:SD:137:LEU:HD23	1.75	0.68
3:ZD:57:ILE:HD11	3:ZD:169:TRP:HA	1.75	0.68
2:UF:120:ASP:OD2	3:BG:76:SER:OG	2.11	0.68
2:UF:201:GLU:O	2:AG:139:ARG:NH2	2.26	0.68
2:AA:310:LEU:HA	2:AA:313:ARG:HG2	1.75	0.68
3:NA:105:ILE:HG12	3:NA:162:LEU:HD11	1.75	0.68
2:EB:69:LEU:HD21	2:KB:47:ILE:HB	1.74	0.68
2:GD:134:LEU:HD21	2:GD:146:LEU:HD21	1.76	0.68

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:HD:105:ILE:HG12	3:HD:162:LEU:HD11	1.76	0.68
2:KE:310:LEU:HA	2:KE:313:ARG:HG2	1.75	0.68
2:AG:21:ARG:HH21	2:AG:24:GLU:HB2	1.58	0.68
2:MG:206:MET:SD	2:MG:207:LYS:N	2.67	0.68
2:J:310:LEU:HA	2:J:313:ARG:HG2	1.75	0.68
3:FB:240:ARG:NH1	3:FB:240:ARG:HA	2.09	0.68
2:WB:110:LEU:O	2:WB:141:GLN:NE2	2.27	0.68
2:GD:125:GLU:HA	3:HD:147:THR:HG21	1.76	0.68
3:ZD:112:GLY:HA3	3:ZD:223:MET:HG2	1.76	0.68
3:BG:115:LEU:HD21	3:BG:218:CYS:HB3	1.76	0.68
2:GG:130:ILE:O	2:GG:134:LEU:HG	1.92	0.68
2:AA:120:ASP:OD1	2:AA:123:ARG:NH1	2.25	0.68
2:QB:120:ASP:OD1	3:XB:76:SER:OG	2.12	0.68
4:KC:100:LEU:HA	4:KC:105:LEU:HA	1.75	0.68
3:ZD:236:LEU:HB3	3:ZD:238:ASN:HD21	1.59	0.68
2:EE:21:ARG:HH21	2:EE:24:GLU:HB2	1.59	0.68
2:WE:186:GLN:HB3	3:DF:74:ARG:NH1	2.08	0.68
2:T:69:LEU:HD21	2:AA:47:ILE:HB	1.76	0.68
2:GA:21:ARG:HH21	2:GA:24:GLU:HB2	1.59	0.68
2:SA:310:LEU:HA	2:SA:313:ARG:HG2	1.75	0.68
2:KB:310:LEU:HA	2:KB:313:ARG:HG2	1.76	0.68
2:QB:128:GLN:HE22	2:QB:164:PHE:HD2	1.40	0.68
3:RB:57:ILE:HD11	3:RB:169:TRP:HA	1.76	0.68
2:WB:310:LEU:HA	2:WB:313:ARG:HG2	1.75	0.68
2:UC:168:GLN:HE22	3:VC:130:LEU:HD11	1.59	0.68
3:ND:96:LEU:HD23	3:ND:97:PRO:HD2	1.76	0.68
3:VF:105:ILE:HG12	3:VF:162:LEU:HD11	1.76	0.68
2:GA:37:LEU:O	2:GA:41:MET:HG3	1.94	0.67
2:QB:201:GLU:HG2	2:WB:139:ARG:NH1	2.09	0.67
3:JC:236:LEU:HB3	3:JC:238:ASN:HD21	1.57	0.67
3:VC:225:GLU:HA	3:VC:228:ARG:HB3	1.77	0.67
3:BD:105:ILE:HG12	3:BD:162:LEU:HD11	1.76	0.67
2:GD:234:PHE:HB3	2:GD:313:ARG:HH21	1.58	0.67
2:OF:186:GLN:NE2	3:VF:71:ASN:HA	2.09	0.67
2:UF:96:ASP:OD2	2:AG:35:GLN:NE2	2.27	0.67
2:AG:219:ARG:HD3	2:AG:223:GLY:HA2	1.76	0.67
2:GG:186:GLN:C	3:NG:74:ARG:HH22	1.97	0.67
2:SG:17:ILE:HD12	2:SG:21:ARG:HG3	1.75	0.67
3:V:190:ASN:ND2	3:BA:234:PRO:O	2.27	0.67
3:NA:240:ARG:NH1	3:NA:240:ARG:HA	2.09	0.67
2:SA:22:ALA:O	2:SA:26:PHE:HB2	1.93	0.67

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:WB:75:GLU:HA	2:WB:78:ARG:HH21	1.60	0.67
2:OC:186:GLN:HB3	3:VC:74:ARG:NH1	2.09	0.67
3:PC:112:GLY:HA3	3:PC:223:MET:HG2	1.76	0.67
2:AD:234:PHE:HB3	2:AD:313:ARG:HH21	1.59	0.67
2:MD:219:ARG:NH2	2:MD:227:GLN:OE1	2.25	0.67
2:KE:34:VAL:O	2:KE:38:SER:OG	2.08	0.67
2:WE:22:ALA:O	2:WE:26:PHE:HB2	1.93	0.67
3:VF:240:ARG:NH1	3:VF:240:ARG:HA	2.09	0.67
2:G:26:PHE:HE2	2:SG:80:VAL:HG12	1.58	0.67
2:B:310:LEU:HA	2:B:313:ARG:HG2	1.75	0.67
2:GA:234:PHE:HB3	2:GA:313:ARG:HH21	1.58	0.67
3:RB:112:GLY:HA3	3:RB:223:MET:HG2	1.76	0.67
2:CC:74:ASN:O	2:CC:78:ARG:NE	2.28	0.67
2:CC:127:PRO:HA	2:CC:130:ILE:HG12	1.76	0.67
3:DC:115:LEU:HD21	3:DC:218:CYS:HB3	1.75	0.67
3:HD:127:VAL:O	3:HD:131:PHE:HB2	1.94	0.67
3:TD:238:ASN:HB3	3:TD:241:HIS:CD2	2.29	0.67
4:CE:106:ILE:CB	3:FE:252:ARG:HG2	2.24	0.67
2:CF:186:GLN:HB3	3:JF:74:ARG:HH22	1.59	0.67
2:IF:310:LEU:HA	2:IF:313:ARG:HG2	1.76	0.67
3:K:54:LEU:HA	3:K:57:ILE:HG22	1.74	0.67
2:YA:127:PRO:HA	2:YA:130:ILE:HG12	1.77	0.67
3:LB:105:ILE:HG12	3:LB:162:LEU:HD11	1.76	0.67
2:WB:69:LEU:HD21	2:CC:47:ILE:HB	1.76	0.67
2:IC:110:LEU:O	2:IC:141:GLN:NE2	2.22	0.67
2:OC:125:GLU:HA	3:PC:147:THR:HG21	1.76	0.67
2:UC:203:ILE:HA	2:UC:206:MET:SD	2.33	0.67
2:EE:310:LEU:HA	2:EE:313:ARG:HG2	1.75	0.67
2:QE:45:ARG:HG3	2:QE:46:GLN:HG3	1.76	0.67
3:JF:105:ILE:HG12	3:JF:162:LEU:HD11	1.77	0.67
4:IG:100:LEU:HA	4:IG:105:LEU:HA	1.75	0.67
2:B:125:GLU:HA	3:C:147:THR:HG21	1.77	0.67
2:MA:310:LEU:HA	2:MA:313:ARG:HG2	1.75	0.67
2:IC:129:ILE:HG23	2:IC:175:LEU:HD21	1.76	0.67
2:AD:310:LEU:HA	2:AD:313:ARG:HG2	1.75	0.67
2:YD:41:MET:HG3	2:YD:42:ALA:N	2.10	0.67
2:IF:234:PHE:HB3	2:IF:313:ARG:HH21	1.59	0.67
2:UF:249:ARG:NH1	2:UF:252:GLN:OE1	2.28	0.67
2:AG:188:LEU:HD23	2:AG:188:LEU:H	1.60	0.67
2:J:186:GLN:NE2	3:V:71:ASN:HA	2.10	0.67
2:T:110:LEU:O	2:T:141:GLN:NE2	2.28	0.67

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:AA:249:ARG:NH1	2:AA:252:GLN:OE1	2.28	0.67
3:BA:115:LEU:HD21	3:BA:218:CYS:HB3	1.77	0.67
3:RB:120:PRO:HD2	3:RB:121:SER:N	2.09	0.67
3:XB:229:GLU:OE1	3:XB:230:LEU:HD12	1.94	0.67
2:AD:207:LYS:NZ	2:AD:210:GLN:HG3	2.10	0.67
2:SD:75:GLU:O	2:SD:79:SER:HB3	1.93	0.67
2:EE:21:ARG:O	2:EE:21:ARG:NE	2.21	0.67
2:KE:125:GLU:HA	3:LE:147:THR:HG21	1.76	0.67
3:LE:105:ILE:HG12	3:LE:162:LEU:HD11	1.77	0.67
2:MG:186:GLN:HB3	3:TG:74:ARG:NH1	2.10	0.67
2:B:69:LEU:HD11	2:J:47:ILE:HG22	1.77	0.67
2:B:234:PHE:HB3	2:B:313:ARG:HH21	1.59	0.67
3:NA:54:LEU:HA	3:NA:57:ILE:HG22	1.77	0.67
2:WB:108:GLU:HA	2:WB:111:ASN:HD22	1.59	0.67
3:VC:72:LEU:HD23	3:VC:73:LEU:HD12	1.76	0.67
2:GD:96:ASP:OD2	2:MD:35:GLN:NE2	2.28	0.67
3:ZD:105:ILE:HG12	3:ZD:162:LEU:HD11	1.76	0.67
2:WE:201:GLU:HG3	2:CF:139:ARG:HD2	1.76	0.67
2:CF:69:LEU:HD21	2:IF:47:ILE:HB	1.76	0.67
2:IF:75:GLU:HA	2:IF:78:ARG:HH21	1.60	0.67
2:GG:76:TYR:HB2	2:MG:41:MET:HE3	1.76	0.67
2:MG:7:THR:HG23	2:MG:37:LEU:HD23	1.77	0.67
2:G:249:ARG:NH1	2:G:252:GLN:OE1	2.28	0.67
2:AA:207:LYS:HG3	2:AA:210:GLN:HG2	1.75	0.67
2:GA:125:GLU:HA	3:HA:147:THR:HG21	1.76	0.67
2:SA:160:ARG:O	2:SA:164:PHE:HB2	1.93	0.67
2:QB:129:ILE:HG23	2:QB:175:LEU:HD21	1.77	0.67
2:CC:249:ARG:NH1	2:CC:252:GLN:OE1	2.28	0.67
2:IC:107:ILE:HG23	2:IC:137:LEU:HD23	1.77	0.67
2:OC:37:LEU:O	2:OC:41:MET:HG3	1.95	0.67
2:OC:130:ILE:O	2:OC:134:LEU:HG	1.95	0.67
2:GD:75:GLU:HA	2:GD:78:ARG:HH21	1.58	0.67
2:YD:249:ARG:NH1	2:YD:252:GLN:OE1	2.28	0.67
3:FE:105:ILE:HG12	3:FE:162:LEU:HD11	1.75	0.67
2:UF:69:LEU:HD11	2:AG:47:ILE:HG22	1.77	0.67
3:M:105:ILE:HG12	3:M:162:LEU:HD11	1.76	0.67
3:M:112:GLY:HA3	3:M:223:MET:HG2	1.75	0.67
2:MA:249:ARG:NH1	2:MA:252:GLN:OE1	2.28	0.67
2:SA:219:ARG:HD3	2:SA:223:GLY:HA2	1.75	0.67
3:ZA:240:ARG:HA	3:ZA:240:ARG:NH1	2.10	0.67
2:EB:234:PHE:HB3	2:EB:313:ARG:HH21	1.58	0.67

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:KB:249:ARG:NH1	2:KB:252:GLN:OE1	2.28	0.67
2:GD:69:LEU:HD11	2:MD:47:ILE:HG22	1.75	0.67
2:MD:249:ARG:NH1	2:MD:252:GLN:OE1	2.28	0.67
2:CF:38:SER:HA	2:CF:41:MET:HE1	1.76	0.67
2:MG:249:ARG:NH1	2:MG:252:GLN:OE1	2.28	0.67
2:SG:234:PHE:HB3	2:SG:313:ARG:HH21	1.60	0.67
2:SA:69:LEU:HD21	2:YA:47:ILE:HB	1.75	0.67
2:EB:78:ARG:O	2:EB:82:VAL:HG23	1.95	0.67
3:FE:240:ARG:HA	3:FE:240:ARG:NH1	2.10	0.67
4:GE:100:LEU:HA	4:GE:105:LEU:HA	1.77	0.67
2:MG:21:ARG:O	2:MG:21:ARG:NE	2.23	0.67
2:G:130:ILE:O	2:G:134:LEU:HG	1.95	0.66
2:MA:168:GLN:NE2	3:NA:130:LEU:HD11	2.10	0.66
2:QB:127:PRO:HA	2:QB:130:ILE:HG12	1.76	0.66
2:IC:207:LYS:HZ1	2:IC:209:GLN:HB3	1.60	0.66
2:UC:249:ARG:NH1	2:UC:252:GLN:OE1	2.29	0.66
2:YD:127:PRO:HA	2:YD:130:ILE:HG12	1.76	0.66
2:QE:65:GLN:NE2	2:WE:47:ILE:O	2.24	0.66
2:MG:130:ILE:O	2:MG:134:LEU:HG	1.95	0.66
2:B:249:ARG:NH1	2:B:252:GLN:OE1	2.29	0.66
2:J:127:PRO:HA	2:J:130:ILE:HG12	1.75	0.66
2:QB:249:ARG:NH1	2:QB:252:GLN:OE1	2.28	0.66
2:CC:219:ARG:HD3	2:CC:223:GLY:HA2	1.77	0.66
3:ND:54:LEU:HA	3:ND:57:ILE:HG22	1.77	0.66
2:SD:90:ALA:HA	2:SD:93:LEU:HG	1.78	0.66
3:ZD:115:LEU:HD21	3:ZD:218:CYS:HB3	1.77	0.66
2:QE:212:GLU:O	2:QE:216:THR:HG23	1.95	0.66
2:GG:205:LEU:HD21	2:MG:140:SER:HA	1.76	0.66
2:SA:249:ARG:NH1	2:SA:252:GLN:OE1	2.29	0.66
2:YA:108:GLU:HA	2:YA:111:ASN:HD22	1.61	0.66
3:RB:240:ARG:NH1	3:RB:240:ARG:HA	2.11	0.66
2:IC:249:ARG:NH1	2:IC:252:GLN:OE1	2.29	0.66
2:GD:37:LEU:O	2:GD:41:MET:HG3	1.95	0.66
2:SD:83:LYS:HA	2:YD:23:ALA:HB1	1.76	0.66
2:KE:249:ARG:NH1	2:KE:252:GLN:OE1	2.28	0.66
2:CF:110:LEU:O	2:CF:141:GLN:NE2	2.23	0.66
2:CF:207:LYS:HZ2	2:CF:210:GLN:HG3	1.59	0.66
2:CF:249:ARG:NH1	2:CF:252:GLN:OE1	2.28	0.66
3:DF:240:ARG:NH1	3:DF:240:ARG:HA	2.10	0.66
2:UF:110:LEU:O	2:UF:141:GLN:NE2	2.27	0.66
2:MG:186:GLN:HB3	3:TG:74:ARG:HH12	1.61	0.66

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:B:127:PRO:HA	2:B:130:ILE:HG12	1.76	0.66
2:J:203:ILE:HA	2:J:206:MET:SD	2.34	0.66
2:J:249:ARG:NH1	2:J:252:GLN:OE1	2.28	0.66
3:K:102:LEU:HD22	3:K:117:VAL:HG22	1.77	0.66
2:T:107:ILE:HG23	2:T:137:LEU:HD23	1.76	0.66
2:T:249:ARG:NH1	2:T:252:GLN:OE1	2.28	0.66
2:GA:310:LEU:HA	2:GA:313:ARG:HG2	1.76	0.66
2:YA:78:ARG:O	2:YA:82:VAL:HG23	1.96	0.66
2:YA:249:ARG:NH1	2:YA:252:GLN:OE1	2.28	0.66
2:EB:45:ARG:HG3	2:EB:46:GLN:HG3	1.77	0.66
2:WB:130:ILE:O	2:WB:134:LEU:HG	1.94	0.66
2:WB:188:LEU:H	3:DC:74:ARG:HH21	1.44	0.66
2:WB:249:ARG:NH1	2:WB:252:GLN:OE1	2.28	0.66
3:ND:169:TRP:HD1	3:ND:175:LEU:HD13	1.60	0.66
2:WE:158:MET:HE3	2:WE:159:LEU:HG	1.77	0.66
3:HG:229:GLU:OE1	3:HG:230:LEU:HD12	1.95	0.66
2:SG:125:GLU:HA	3:TG:147:THR:HG21	1.77	0.66
2:J:120:ASP:OD2	3:V:76:SER:OG	2.07	0.66
3:K:112:GLY:HA3	3:K:223:MET:HG2	1.76	0.66
3:TA:57:ILE:HD11	3:TA:169:TRP:HA	1.77	0.66
3:XB:105:ILE:HG12	3:XB:162:LEU:HD11	1.77	0.66
2:UC:234:PHE:HB3	2:UC:313:ARG:HH21	1.59	0.66
2:EE:249:ARG:NH1	2:EE:252:GLN:OE1	2.28	0.66
3:FE:112:GLY:HA3	3:FE:223:MET:HG2	1.77	0.66
3:LE:54:LEU:HA	3:LE:57:ILE:HG22	1.77	0.66
2:QE:249:ARG:NH1	2:QE:252:GLN:OE1	2.29	0.66
3:XE:240:ARG:NH1	3:XE:240:ARG:HA	2.10	0.66
3:DF:53:ALA:HB3	3:DF:231:LEU:HA	1.76	0.66
2:IF:249:ARG:NH1	2:IF:252:GLN:OE1	2.28	0.66
3:M:247:ARG:NH1	3:M:247:ARG:O	2.29	0.66
2:B:160:ARG:O	2:B:164:PHE:HB2	1.95	0.66
4:D:100:LEU:HA	4:D:105:LEU:HA	1.76	0.66
2:T:205:LEU:HD21	2:AA:140:SER:HA	1.77	0.66
2:MA:83:LYS:HB3	2:SA:27:LYS:HB2	1.76	0.66
2:MA:234:PHE:HB3	2:MA:313:ARG:HH21	1.60	0.66
2:YA:96:ASP:OD2	2:EB:35:GLN:NE2	2.29	0.66
2:KB:69:LEU:O	2:KB:72:ASN:ND2	2.28	0.66
3:PC:240:ARG:HA	3:PC:240:ARG:NH1	2.10	0.66
2:UC:125:GLU:HA	3:VC:147:THR:HG21	1.77	0.66
2:UC:129:ILE:HG23	2:UC:175:LEU:HD21	1.76	0.66
2:AD:249:ARG:NH1	2:AD:252:GLN:OE1	2.28	0.66

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:ND:198:ILE:HG23	3:ND:221:PHE:CD1	2.31	0.66
2:EE:125:GLU:HA	3:FE:147:THR:HG21	1.77	0.66
2:EE:203:ILE:HA	2:EE:206:MET:SD	2.36	0.66
3:RE:53:ALA:HB3	3:RE:231:LEU:HA	1.77	0.66
2:OF:107:ILE:HD13	2:OF:136:HIS:HB2	1.76	0.66
2:SG:249:ARG:NH1	2:SG:252:GLN:OE1	2.28	0.66
2:G:201:GLU:HG3	2:B:139:ARG:HD2	1.78	0.66
2:J:219:ARG:NH2	2:J:227:GLN:OE1	2.24	0.66
2:GA:249:ARG:NH1	2:GA:252:GLN:OE1	2.29	0.66
2:MA:107:ILE:HG23	2:MA:137:LEU:HD23	1.78	0.66
3:DC:106:HIS:CE1	3:DC:113:THR:HG22	2.31	0.66
2:IC:199:ALA:HA	2:IC:202:ILE:HD12	1.77	0.66
3:PC:58:ASN:ND2	3:PC:81:VAL:HG11	2.10	0.66
2:AD:21:ARG:O	2:AD:21:ARG:NE	2.21	0.66
2:AD:21:ARG:HH21	2:AD:24:GLU:HB2	1.61	0.66
2:GD:249:ARG:NH1	2:GD:252:GLN:OE1	2.28	0.66
3:LE:198:ILE:HG23	3:LE:221:PHE:CD1	2.30	0.66
3:DF:102:LEU:HD22	3:DF:117:VAL:HG22	1.78	0.66
2:OF:89:ARG:HH22	2:OF:93:LEU:HD21	1.60	0.66
3:PF:240:ARG:HA	3:PF:240:ARG:NH1	2.10	0.66
2:MG:186:GLN:HE22	3:TG:71:ASN:HA	1.60	0.66
3:NG:169:TRP:HD1	3:NG:175:LEU:HD13	1.59	0.66
3:TG:53:ALA:HB3	3:TG:231:LEU:HA	1.77	0.66
2:T:21:ARG:O	2:T:21:ARG:NE	2.19	0.66
2:SA:65:GLN:NE2	2:YA:47:ILE:O	2.25	0.66
2:SA:234:PHE:HB3	2:SA:313:ARG:HH21	1.60	0.66
2:YA:160:ARG:O	2:YA:164:PHE:CB	2.43	0.66
2:AD:130:ILE:O	2:AD:134:LEU:HG	1.95	0.66
3:LE:240:ARG:NH1	3:LE:240:ARG:HA	2.10	0.66
2:WE:127:PRO:HA	2:WE:130:ILE:HG12	1.77	0.66
2:OF:38:SER:HA	2:OF:41:MET:HE2	1.76	0.66
2:GG:249:ARG:NH1	2:GG:252:GLN:OE1	2.29	0.66
2:B:219:ARG:HD3	2:B:223:GLY:HA2	1.78	0.66
2:AA:160:ARG:O	2:AA:164:PHE:CB	2.44	0.66
2:EB:249:ARG:NH1	2:EB:252:GLN:OE1	2.28	0.66
2:QB:107:ILE:HA	2:QB:110:LEU:HD12	1.78	0.66
3:XB:240:ARG:NH1	3:XB:240:ARG:HA	2.11	0.66
2:AD:65:GLN:NE2	2:GD:47:ILE:O	2.26	0.66
3:LE:169:TRP:HD1	3:LE:175:LEU:HD13	1.61	0.66
2:GA:160:ARG:O	2:GA:164:PHE:CB	2.44	0.66
2:AD:96:ASP:OD2	2:GD:35:GLN:NE2	2.29	0.66

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:HD:53:ALA:HB3	3:HD:231:LEU:HA	1.77	0.66
2:SD:249:ARG:NH1	2:SD:252:GLN:OE1	2.28	0.66
2:WE:65:GLN:NE2	2:CF:47:ILE:O	2.24	0.66
2:WE:249:ARG:NH1	2:WE:252:GLN:OE1	2.28	0.66
2:OF:207:LYS:NZ	2:OF:210:GLN:HG3	2.11	0.66
2:J:108:GLU:HA	2:J:111:ASN:HD22	1.60	0.65
3:BA:127:VAL:O	3:BA:131:PHE:HB2	1.96	0.65
2:GA:65:GLN:NE2	2:MA:47:ILE:O	2.29	0.65
2:KB:26:PHE:CE2	2:KB:37:LEU:HD11	2.32	0.65
2:MD:186:GLN:HE22	3:TD:71:ASN:HA	1.61	0.65
2:QE:26:PHE:HE1	2:QE:34:VAL:HG22	1.61	0.65
3:BG:101:ASN:ND2	3:BG:154:ILE:HG21	2.11	0.65
2:T:130:ILE:O	2:T:134:LEU:HG	1.96	0.65
3:HA:115:LEU:HD21	3:HA:218:CYS:HB3	1.77	0.65
3:NA:169:TRP:HD1	3:NA:175:LEU:HD13	1.60	0.65
2:QB:78:ARG:O	2:QB:82:VAL:HG23	1.96	0.65
2:IC:160:ARG:O	2:IC:164:PHE:HB2	1.93	0.65
2:MD:118:ALA:HA	2:MD:121:LEU:HG	1.77	0.65
1:PE:543:ASP:HA	2:QE:21:ARG:HH11	1.61	0.65
3:RE:198:ILE:HG23	3:RE:221:PHE:CD1	2.30	0.65
3:DF:72:LEU:HD23	3:DF:73:LEU:HD12	1.78	0.65
2:OF:249:ARG:NH1	2:OF:252:GLN:OE1	2.28	0.65
2:AG:249:ARG:NH1	2:AG:252:GLN:OE1	2.28	0.65
3:BG:240:ARG:NH1	3:BG:240:ARG:HA	2.11	0.65
3:HG:105:ILE:HG12	3:HG:162:LEU:HD11	1.78	0.65
2:GA:128:GLN:HE22	2:GA:164:PHE:HD2	1.42	0.65
2:GA:203:ILE:HA	2:GA:206:MET:SD	2.37	0.65
2:KB:130:ILE:O	2:KB:134:LEU:HG	1.97	0.65
2:OC:249:ARG:NH1	2:OC:252:GLN:OE1	2.29	0.65
2:OF:125:GLU:HA	3:PF:147:THR:HG21	1.79	0.65
3:PF:127:VAL:O	3:PF:131:PHE:HB2	1.96	0.65
2:G:206:MET:SD	2:G:207:LYS:N	2.70	0.65
2:MA:130:ILE:O	2:MA:134:LEU:HG	1.96	0.65
2:SA:230:ILE:HA	2:SA:233:MET:HE2	1.78	0.65
2:EB:187:ASN:O	2:EB:189:LYS:NZ	2.28	0.65
2:QB:14:LEU:HD12	2:QB:22:ALA:HB1	1.79	0.65
3:DC:240:ARG:NH1	3:DC:240:ARG:HA	2.11	0.65
2:IC:96:ASP:OD2	2:OC:35:GLN:NE2	2.30	0.65
3:ZD:121:SER:O	3:ZD:125:ILE:HG13	1.97	0.65
3:ZD:127:VAL:O	3:ZD:131:PHE:HB2	1.97	0.65
3:JF:53:ALA:HB3	3:JF:231:LEU:HA	1.78	0.65

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:AA:203:ILE:HA	2:AA:206:MET:SD	2.37	0.65
2:IC:69:LEU:HD11	2:OC:47:ILE:HG22	1.76	0.65
3:JC:240:ARG:NH1	3:JC:240:ARG:HA	2.12	0.65
2:AD:129:ILE:HG23	2:AD:175:LEU:HD21	1.78	0.65
2:MD:234:PHE:HB3	2:MD:313:ARG:HH21	1.59	0.65
2:IF:80:VAL:O	2:IF:83:LYS:HG2	1.97	0.65
2:OF:12:ILE:HD11	2:OF:55:VAL:HG11	1.79	0.65
3:HG:240:ARG:NH1	3:HG:240:ARG:HA	2.12	0.65
2:G:47:ILE:O	2:SG:65:GLN:NE2	2.25	0.65
2:G:96:ASP:OD2	2:B:35:GLN:NE2	2.29	0.65
4:R:106:ILE:O	4:R:126:ILE:N	2.29	0.65
2:WE:69:LEU:O	2:WE:72:ASN:ND2	2.29	0.65
2:CF:69:LEU:HD11	2:IF:47:ILE:HG22	1.79	0.65
3:PF:225:GLU:HA	3:PF:228:ARG:HB3	1.77	0.65
3:HA:240:ARG:NH1	3:HA:240:ARG:HA	2.11	0.65
2:MA:219:ARG:HH21	2:MA:230:ILE:HD12	1.62	0.65
3:JC:191:ILE:HG13	3:JC:192:THR:HG23	1.79	0.65
3:M:90:HIS:O	3:M:94:ARG:HG2	1.97	0.65
2:B:230:ILE:HA	2:B:233:MET:HE2	1.79	0.65
2:T:128:GLN:HE22	2:T:164:PHE:HD2	1.42	0.65
3:V:53:ALA:HB3	3:V:231:LEU:HA	1.79	0.65
2:AA:12:ILE:HD11	2:AA:55:VAL:HG11	1.77	0.65
3:DC:101:ASN:ND2	3:DC:154:ILE:HG21	2.12	0.65
3:HD:169:TRP:HD1	3:HD:175:LEU:HD13	1.62	0.65
2:IF:160:ARG:O	2:IF:164:PHE:HB2	1.96	0.65
4:KF:100:LEU:HA	4:KF:105:LEU:HA	1.79	0.65
2:OF:90:ALA:HA	2:OF:93:LEU:HG	1.79	0.65
3:PF:120:PRO:HA	3:PF:123:VAL:HG22	1.79	0.65
2:GA:21:ARG:O	2:GA:21:ARG:NE	2.19	0.65
2:EB:152:ARG:HA	2:EB:155:HIS:CE1	2.32	0.65
3:HD:54:LEU:HA	3:HD:57:ILE:HG22	1.79	0.65
3:HD:229:GLU:OE1	3:HD:230:LEU:HD12	1.96	0.65
3:ND:53:ALA:HB3	3:ND:231:LEU:HA	1.78	0.65
2:QE:30:SER:OG	2:QE:33:GLU:OE1	2.10	0.65
3:XE:127:VAL:O	3:XE:131:PHE:HB2	1.97	0.65
2:UF:188:LEU:H	3:BG:74:ARG:HH21	1.44	0.65
2:AG:127:PRO:HA	2:AG:130:ILE:HG12	1.78	0.65
3:NG:121:SER:O	3:NG:125:ILE:HG13	1.97	0.65
3:C:112:GLY:HA3	3:C:223:MET:HG2	1.79	0.65
3:K:121:SER:O	3:K:125:ILE:HG13	1.96	0.65
2:T:65:GLN:NE2	2:AA:47:ILE:O	2.30	0.65

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:AA:69:LEU:HD11	2:GA:47:ILE:HG22	1.79	0.65
3:TA:121:SER:O	3:TA:125:ILE:HG13	1.97	0.65
2:YA:125:GLU:HA	3:ZA:147:THR:HG21	1.79	0.65
2:EB:125:GLU:HA	3:FB:147:THR:HG21	1.78	0.65
3:FB:54:LEU:HA	3:FB:57:ILE:HG22	1.79	0.65
3:FB:112:GLY:HA3	3:FB:223:MET:HG2	1.79	0.65
2:QB:45:ARG:HG3	2:QB:46:GLN:HG3	1.79	0.65
2:CC:152:ARG:HA	2:CC:155:HIS:CE1	2.32	0.65
3:TD:53:ALA:HB3	3:TD:231:LEU:HA	1.79	0.65
2:YD:151:GLU:HA	2:YD:154:ARG:HG2	1.79	0.65
2:EE:129:ILE:HG23	2:EE:175:LEU:HD21	1.78	0.65
3:LE:57:ILE:HD11	3:LE:169:TRP:HA	1.78	0.65
2:CF:130:ILE:O	2:CF:134:LEU:HG	1.97	0.65
3:PF:96:LEU:HD23	3:PF:97:PRO:HD2	1.79	0.65
2:AG:152:ARG:HA	2:AG:155:HIS:CE1	2.32	0.65
2:AG:188:LEU:HD21	3:HG:74:ARG:NH1	2.09	0.65
2:B:78:ARG:O	2:B:82:VAL:HG23	1.96	0.64
2:J:187:ASN:O	2:J:189:LYS:NZ	2.31	0.64
3:V:127:VAL:O	3:V:131:PHE:HB2	1.98	0.64
3:TA:72:LEU:HD23	3:TA:73:LEU:HD12	1.79	0.64
2:OC:190:ARG:O	2:UC:166:GLY:HA3	1.97	0.64
2:YD:152:ARG:HA	2:YD:155:HIS:CE1	2.32	0.64
3:FE:53:ALA:HB3	3:FE:231:LEU:HA	1.76	0.64
2:QE:234:PHE:HB3	2:QE:313:ARG:HH21	1.61	0.64
2:AG:65:GLN:NE2	2:GG:47:ILE:O	2.24	0.64
2:MG:125:GLU:HA	3:NG:147:THR:HG21	1.78	0.64
3:NG:198:ILE:HG23	3:NG:221:PHE:CD1	2.32	0.64
2:G:21:ARG:O	2:G:21:ARG:NE	2.22	0.64
2:G:129:ILE:HG23	2:G:175:LEU:HD21	1.78	0.64
3:NA:96:LEU:HD23	3:NA:97:PRO:HD2	1.78	0.64
3:TA:53:ALA:HB3	3:TA:231:LEU:HA	1.79	0.64
2:YA:107:ILE:HG22	2:YA:111:ASN:HD21	1.63	0.64
2:EB:130:ILE:O	2:EB:134:LEU:HG	1.97	0.64
3:LB:72:LEU:HD23	3:LB:73:LEU:HD12	1.78	0.64
3:PC:54:LEU:HA	3:PC:57:ILE:HG22	1.79	0.64
2:UC:107:ILE:HG23	2:UC:137:LEU:HD23	1.79	0.64
3:BD:90:HIS:O	3:BD:94:ARG:HG2	1.97	0.64
2:EE:205:LEU:HD13	2:KE:139:ARG:HB3	1.79	0.64
2:KE:244:ASP:OD1	2:KE:276:ARG:NH2	2.31	0.64
3:LE:96:LEU:HD23	3:LE:97:PRO:HD2	1.78	0.64
2:WE:125:GLU:HA	3:XE:147:THR:HG21	1.78	0.64

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:WE:130:ILE:O	2:WE:134:LEU:HG	1.98	0.64
2:IF:34:VAL:O	2:IF:38:SER:OG	2.11	0.64
2:IF:188:LEU:HB3	3:PF:74:ARG:HE	1.59	0.64
2:IF:230:ILE:HA	2:IF:233:MET:HE2	1.78	0.64
3:PF:105:ILE:HG12	3:PF:162:LEU:HD11	1.80	0.64
2:SG:152:ARG:NH2	2:SG:153:LEU:HB2	2.13	0.64
3:K:105:ILE:HG12	3:K:162:LEU:HD11	1.79	0.64
3:V:77:PRO:HB3	3:V:208:ILE:HG12	1.80	0.64
3:ZA:127:VAL:O	3:ZA:131:PHE:HB2	1.97	0.64
2:KB:151:GLU:O	2:KB:155:HIS:ND1	2.30	0.64
2:IC:127:PRO:HA	2:IC:130:ILE:HG12	1.80	0.64
3:DF:238:ASN:HB3	3:DF:241:HIS:HD2	1.63	0.64
2:OF:110:LEU:HB3	2:OF:141:GLN:HE22	1.63	0.64
2:OF:128:GLN:HE22	2:OF:164:PHE:HD2	1.42	0.64
2:MA:129:ILE:HG23	2:MA:175:LEU:HD21	1.79	0.64
2:SA:107:ILE:HG23	2:SA:137:LEU:HD23	1.78	0.64
3:XB:53:ALA:HB3	3:XB:231:LEU:HA	1.79	0.64
3:XB:115:LEU:HD21	3:XB:218:CYS:HB3	1.79	0.64
2:OC:206:MET:SD	2:OC:207:LYS:N	2.71	0.64
2:OC:234:PHE:HB3	2:OC:313:ARG:HH21	1.61	0.64
3:RE:191:ILE:HG13	3:RE:192:THR:HG23	1.79	0.64
2:WE:8:ASP:O	2:WE:12:ILE:HG12	1.97	0.64
3:NG:112:GLY:HA3	3:NG:223:MET:HG2	1.79	0.64
2:J:115:PRO:O	2:J:119:ALA:HB2	1.96	0.64
3:K:169:TRP:HD1	3:K:175:LEU:HD13	1.62	0.64
2:T:244:ASP:OD1	2:T:276:ARG:NH2	2.31	0.64
2:AA:80:VAL:O	2:AA:83:LYS:HG2	1.98	0.64
2:AA:234:PHE:HB3	2:AA:313:ARG:HH21	1.63	0.64
2:SA:129:ILE:HG23	2:SA:175:LEU:HD21	1.78	0.64
2:KB:120:ASP:OD2	2:KB:123:ARG:NH1	2.31	0.64
2:OC:127:PRO:HA	2:OC:130:ILE:HG12	1.80	0.64
2:KE:207:LYS:HZ3	2:KE:209:GLN:HB3	1.61	0.64
3:RE:105:ILE:HG12	3:RE:162:LEU:HD11	1.80	0.64
3:DF:238:ASN:HB3	3:DF:241:HIS:CD2	2.32	0.64
2:T:125:GLU:HA	3:V:147:THR:HG21	1.79	0.64
2:MA:212:GLU:O	2:MA:216:THR:HG22	1.98	0.64
3:FB:127:VAL:O	3:FB:131:PHE:HB2	1.97	0.64
3:VC:54:LEU:HA	3:VC:57:ILE:HG22	1.80	0.64
2:YD:207:LYS:HZ2	2:YD:210:GLN:HG3	1.62	0.64
2:WE:115:PRO:O	2:WE:119:ALA:CB	2.46	0.64
2:CF:127:PRO:HA	2:CF:130:ILE:HG12	1.78	0.64

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:UF:125:GLU:HA	3:VF:147:THR:HG21	1.80	0.64
2:B:83:LYS:HD3	2:J:26:PHE:CD2	2.33	0.64
2:QB:125:GLU:HA	3:RB:147:THR:HG21	1.79	0.64
2:CC:129:ILE:HG23	2:CC:175:LEU:HD21	1.79	0.64
2:SD:187:ASN:O	2:SD:189:LYS:NZ	2.30	0.64
2:KE:168:GLN:NE2	3:LE:130:LEU:HD11	2.13	0.64
2:CF:113:MET:HG3	2:CF:117:SER:HB2	1.78	0.64
2:CF:152:ARG:HA	2:CF:155:HIS:CE1	2.32	0.64
2:CF:186:GLN:CB	3:JF:74:ARG:HH22	2.11	0.64
3:VF:115:LEU:HD21	3:VF:218:CYS:HB3	1.80	0.64
2:AG:71:ILE:O	2:AG:74:ASN:ND2	2.30	0.64
2:MG:234:PHE:HB3	2:MG:313:ARG:HH21	1.63	0.64
3:TG:112:GLY:HA3	3:TG:223:MET:HG2	1.79	0.64
3:M:121:SER:O	3:M:125:ILE:HG13	1.98	0.64
3:C:106:HIS:CE1	3:C:113:THR:HG22	2.33	0.64
2:WB:125:GLU:HA	3:XB:147:THR:HG21	1.80	0.64
2:CC:108:GLU:HA	2:CC:111:ASN:HD22	1.63	0.64
2:IC:83:LYS:HG2	2:OC:26:PHE:HB3	1.79	0.64
2:YD:12:ILE:HD11	2:YD:55:VAL:HG11	1.77	0.64
2:IF:120:ASP:OD2	2:IF:123:ARG:NH1	2.30	0.64
2:AG:129:ILE:HG23	2:AG:175:LEU:HD21	1.79	0.64
2:MG:127:PRO:HA	2:MG:130:ILE:HG12	1.78	0.64
3:K:240:ARG:NH1	3:K:240:ARG:HA	2.12	0.64
3:BA:57:ILE:HD11	3:BA:169:TRP:HA	1.78	0.64
3:NA:225:GLU:HA	3:NA:228:ARG:HB3	1.79	0.64
3:FB:53:ALA:HB3	3:FB:231:LEU:HA	1.80	0.64
2:OC:107:ILE:HG22	2:OC:111:ASN:HD21	1.63	0.64
3:PC:169:TRP:HD1	3:PC:175:LEU:HD13	1.62	0.64
3:BD:121:SER:O	3:BD:125:ILE:HG13	1.98	0.64
2:KE:89:ARG:HH22	2:KE:93:LEU:HG	1.62	0.64
3:RE:57:ILE:HD11	3:RE:169:TRP:HA	1.79	0.64
2:OF:74:ASN:O	2:OF:78:ARG:NE	2.31	0.64
3:C:121:SER:O	3:C:125:ILE:HG13	1.98	0.64
3:NA:127:VAL:O	3:NA:131:PHE:HB2	1.98	0.64
2:YA:69:LEU:O	2:YA:72:ASN:ND2	2.30	0.64
3:JC:96:LEU:HD23	3:JC:97:PRO:HD2	1.79	0.64
2:OC:187:ASN:O	2:OC:189:LYS:NZ	2.31	0.64
2:MD:207:LYS:HD3	2:MD:210:GLN:HE21	1.63	0.64
2:WE:234:PHE:HB3	2:WE:313:ARG:HH21	1.61	0.64
3:XE:115:LEU:HD21	3:XE:218:CYS:HB3	1.80	0.64
2:IF:152:ARG:HA	2:IF:155:HIS:CE1	2.33	0.64

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:OF:127:PRO:HA	2:OF:130:ILE:HG12	1.79	0.64
2:UF:234:PHE:HB3	2:UF:313:ARG:HH21	1.62	0.64
3:M:54:LEU:HA	3:M:57:ILE:HG22	1.78	0.63
2:J:115:PRO:O	2:J:119:ALA:CB	2.45	0.63
2:MA:244:ASP:OD1	2:MA:276:ARG:NH2	2.31	0.63
3:ZA:105:ILE:HG12	3:ZA:162:LEU:HD11	1.80	0.63
2:OC:76:TYR:HE2	2:UC:37:LEU:HD13	1.62	0.63
2:UC:30:SER:OG	2:UC:33:GLU:OE1	2.16	0.63
3:VC:75:ARG:HG3	3:VC:208:ILE:HD11	1.80	0.63
2:MD:212:GLU:O	2:MD:216:THR:HG22	1.98	0.63
3:FE:182:SER:OG	3:FE:183:GLU:N	2.31	0.63
2:QE:69:LEU:HD21	2:WE:47:ILE:HB	1.80	0.63
2:QE:244:ASP:OD1	2:QE:276:ARG:NH2	2.30	0.63
2:CF:244:ASP:OD1	2:CF:276:ARG:NH2	2.30	0.63
2:IF:129:ILE:HG23	2:IF:175:LEU:HD21	1.80	0.63
3:VF:90:HIS:O	3:VF:94:ARG:HG2	1.98	0.63
2:GG:75:GLU:O	2:GG:79:SER:HB3	1.97	0.63
2:T:234:PHE:HB3	2:T:313:ARG:HH21	1.63	0.63
3:V:54:LEU:HA	3:V:57:ILE:HG22	1.80	0.63
3:V:59:GLU:OE1	3:V:59:GLU:N	2.30	0.63
2:GA:201:GLU:HG3	2:MA:139:ARG:CZ	2.28	0.63
2:SA:37:LEU:O	2:SA:41:MET:HG3	1.98	0.63
2:IC:202:ILE:HG13	2:OC:139:ARG:NH2	2.13	0.63
2:YD:26:PHE:CZ	2:YD:37:LEU:HD11	2.33	0.63
2:OF:79:SER:HA	2:OF:82:VAL:HG22	1.80	0.63
2:GG:127:PRO:HA	2:GG:130:ILE:HG12	1.80	0.63
2:GG:201:GLU:HB3	2:MG:139:ARG:HH22	1.63	0.63
2:MG:21:ARG:NH2	2:MG:24:GLU:HB2	2.12	0.63
3:C:54:LEU:HA	3:C:57:ILE:HG22	1.80	0.63
2:SA:76:TYR:HE2	2:YA:26:PHE:CZ	2.17	0.63
3:ZA:72:LEU:HD23	3:ZA:73:LEU:HD12	1.80	0.63
2:KB:212:GLU:O	2:KB:216:THR:HG23	1.98	0.63
2:YD:234:PHE:HB3	2:YD:313:ARG:HH21	1.62	0.63
2:KE:212:GLU:O	2:KE:216:THR:HG23	1.97	0.63
2:WE:110:LEU:O	2:WE:141:GLN:NE2	2.28	0.63
3:XE:54:LEU:HA	3:XE:57:ILE:HG22	1.80	0.63
3:DF:54:LEU:HA	3:DF:57:ILE:HG22	1.79	0.63
2:T:127:PRO:HA	2:T:130:ILE:HG12	1.80	0.63
3:V:169:TRP:HD1	3:V:175:LEU:HD13	1.64	0.63
2:AA:212:GLU:O	2:AA:216:THR:HG22	1.99	0.63
2:GA:11:VAL:HG11	2:GA:41:MET:HB3	1.81	0.63

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:DC:57:ILE:HD11	3:DC:169:TRP:HA	1.81	0.63
2:MD:205:LEU:HD13	2:SD:139:ARG:HB3	1.81	0.63
2:KE:129:ILE:HG23	2:KE:175:LEU:HD21	1.80	0.63
2:MG:37:LEU:O	2:MG:41:MET:HG3	1.98	0.63
3:HA:169:TRP:HD1	3:HA:175:LEU:HD13	1.63	0.63
3:HA:182:SER:OG	3:HA:183:GLU:N	2.31	0.63
2:EB:21:ARG:NH2	2:EB:24:GLU:HB2	2.12	0.63
2:CC:21:ARG:O	2:CC:21:ARG:NE	2.17	0.63
2:CC:187:ASN:O	2:CC:189:LYS:NZ	2.31	0.63
2:UC:22:ALA:O	2:UC:26:PHE:HB2	1.98	0.63
2:AD:67:ALA:O	2:AD:71:ILE:HG12	1.99	0.63
1:RD:547:VAL:HA	2:SD:28:HIS:HE1	1.62	0.63
3:TD:54:LEU:HA	3:TD:57:ILE:HG22	1.79	0.63
2:OF:21:ARG:HH21	2:OF:24:GLU:HB2	1.63	0.63
2:OF:188:LEU:HD23	2:OF:188:LEU:H	1.62	0.63
2:AG:205:LEU:HD22	2:GG:139:ARG:HB3	1.79	0.63
2:SG:102:ASP:HB2	2:SG:108:GLU:HB2	1.79	0.63
2:G:234:PHE:HB3	2:G:313:ARG:HH21	1.64	0.63
2:T:129:ILE:HG23	2:T:175:LEU:HD21	1.80	0.63
3:BA:169:TRP:HD1	3:BA:175:LEU:HD13	1.63	0.63
2:MA:110:LEU:O	2:MA:141:GLN:NE2	2.23	0.63
2:MA:186:GLN:HB3	3:TA:74:ARG:NH1	2.14	0.63
2:SA:244:ASP:OD1	2:SA:276:ARG:NH2	2.31	0.63
2:YA:152:ARG:HH12	2:YA:153:LEU:HB2	1.62	0.63
2:EB:203:ILE:HA	2:EB:206:MET:HE3	1.80	0.63
2:IC:212:GLU:O	2:IC:216:THR:HG22	1.99	0.63
3:JC:105:ILE:HG12	3:JC:162:LEU:HD11	1.80	0.63
2:OC:186:GLN:HE22	3:VC:71:ASN:HA	1.63	0.63
2:OC:212:GLU:O	2:OC:216:THR:HG22	1.99	0.63
4:IE:106:ILE:CB	3:LE:252:ARG:HG2	2.28	0.63
3:NG:127:VAL:O	3:NG:131:PHE:HB2	1.98	0.63
2:B:219:ARG:NH2	2:B:227:GLN:OE1	2.29	0.63
2:J:38:SER:HA	2:J:41:MET:HE1	1.79	0.63
2:J:186:GLN:HB3	3:V:74:ARG:HH12	1.64	0.63
3:V:191:ILE:HG13	3:V:192:THR:HG23	1.81	0.63
3:ZA:54:LEU:HA	3:ZA:57:ILE:HG22	1.81	0.63
2:EB:151:GLU:HA	2:EB:154:ARG:HG2	1.79	0.63
2:KB:65:GLN:NE2	2:QB:47:ILE:O	2.31	0.63
2:KB:129:ILE:HG23	2:KB:175:LEU:HD21	1.79	0.63
2:WB:234:PHE:HB3	2:WB:313:ARG:HH21	1.61	0.63
3:BD:54:LEU:HA	3:BD:57:ILE:HG22	1.79	0.63

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:SD:244:ASP:OD1	2:SD:276:ARG:NH2	2.31	0.63
2:YD:212:GLU:O	2:YD:216:THR:HG22	1.99	0.63
2:CF:188:LEU:HD23	3:JF:74:ARG:HH21	1.64	0.63
3:JF:90:HIS:O	3:JF:94:ARG:HG2	1.98	0.63
2:OF:75:GLU:HA	2:OF:78:ARG:HH21	1.63	0.63
2:OF:120:ASP:OD1	3:VF:76:SER:OG	2.16	0.63
3:PF:75:ARG:HG3	3:PF:208:ILE:HD11	1.81	0.63
3:PF:102:LEU:HD22	3:PF:117:VAL:HG22	1.80	0.63
2:GG:160:ARG:O	2:GG:164:PHE:CB	2.46	0.63
2:B:107:ILE:HG22	2:B:111:ASN:HD21	1.64	0.63
2:B:129:ILE:HG23	2:B:175:LEU:HD21	1.79	0.63
2:B:212:GLU:O	2:B:216:THR:HG22	1.99	0.63
3:NA:112:GLY:HA3	3:NA:223:MET:HG2	1.81	0.63
2:YA:168:GLN:HE22	3:ZA:130:LEU:HD11	1.64	0.63
2:KB:93:LEU:O	2:KB:97:ILE:HG12	1.99	0.63
2:AD:113:MET:O	2:AD:141:GLN:NE2	2.31	0.63
2:QE:205:LEU:HD21	2:WE:140:SER:HA	1.81	0.63
2:OF:120:ASP:OD2	2:OF:123:ARG:NH1	2.31	0.63
2:UF:129:ILE:HG23	2:UF:175:LEU:HD21	1.79	0.63
2:AG:69:LEU:HD21	2:GG:47:ILE:HB	1.81	0.63
2:MG:107:ILE:HG22	2:MG:111:ASN:HD21	1.63	0.63
2:J:129:ILE:HG23	2:J:175:LEU:HD21	1.79	0.63
3:TA:112:GLY:HA3	3:TA:223:MET:HG2	1.81	0.63
3:RB:127:VAL:O	3:RB:131:PHE:HB2	1.98	0.63
3:TD:190:ASN:ND2	3:ZD:234:PRO:O	2.32	0.63
2:EE:110:LEU:O	2:EE:141:GLN:NE2	2.29	0.63
2:EE:234:PHE:HB3	2:EE:313:ARG:HH21	1.63	0.63
2:WE:129:ILE:HG23	2:WE:175:LEU:HD21	1.80	0.63
2:UF:201:GLU:HG3	2:AG:139:ARG:NH1	2.14	0.63
3:ZA:121:SER:O	3:ZA:125:ILE:HG13	1.99	0.62
3:LB:198:ILE:HG23	3:LB:221:PHE:CD1	2.33	0.62
3:JC:115:LEU:HD21	3:JC:218:CYS:HB3	1.81	0.62
3:VC:77:PRO:HB3	3:VC:208:ILE:HB	1.81	0.62
3:TD:121:SER:O	3:TD:125:ILE:HG13	1.99	0.62
2:CF:125:GLU:HA	3:DF:147:THR:HG21	1.80	0.62
2:AG:37:LEU:O	2:AG:41:MET:HG3	1.98	0.62
3:HG:115:LEU:HD21	3:HG:218:CYS:HB3	1.81	0.62
2:SG:21:ARG:NH2	2:SG:24:GLU:HB2	2.13	0.62
2:SG:206:MET:HB3	2:SG:210:GLN:NE2	2.14	0.62
2:G:186:GLN:NE2	3:C:71:ASN:HA	2.13	0.62
2:B:207:LYS:NZ	2:B:210:GLN:HG3	2.14	0.62

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:J:89:ARG:HH22	2:J:93:LEU:HG	1.63	0.62
2:T:71:ILE:O	2:T:74:ASN:ND2	2.32	0.62
2:T:206:MET:SD	2:T:207:LYS:N	2.72	0.62
2:AA:188:LEU:HB2	2:GA:168:GLN:HB3	1.81	0.62
3:RB:105:ILE:HG12	3:RB:162:LEU:HD11	1.81	0.62
2:IC:186:GLN:HB3	3:PC:74:ARG:NH1	2.14	0.62
3:JC:121:SER:O	3:JC:125:ILE:HG13	1.98	0.62
2:UC:26:PHE:CE1	2:UC:34:VAL:HG22	2.34	0.62
2:GD:212:GLU:O	2:GD:216:THR:HG22	1.99	0.62
3:HD:121:SER:O	3:HD:125:ILE:HG13	1.99	0.62
3:ND:102:LEU:HD22	3:ND:117:VAL:HG22	1.80	0.62
3:ND:127:VAL:O	3:ND:131:PHE:HB2	2.00	0.62
2:SD:129:ILE:HG23	2:SD:175:LEU:HD21	1.80	0.62
3:TD:77:PRO:HB3	3:TD:208:ILE:HG12	1.81	0.62
2:YD:151:GLU:O	2:YD:155:HIS:ND1	2.31	0.62
2:QE:186:GLN:HB3	3:XE:74:ARG:HH21	1.63	0.62
2:MG:212:GLU:O	2:MG:216:THR:HG22	1.99	0.62
3:M:191:ILE:HG13	3:M:192:THR:HG23	1.81	0.62
2:B:107:ILE:HG23	2:B:137:LEU:HD23	1.80	0.62
2:J:125:GLU:HA	3:K:147:THR:HG21	1.81	0.62
2:AA:20:ASP:OD1	2:AA:21:ARG:N	2.32	0.62
2:YA:130:ILE:O	2:YA:134:LEU:HG	1.99	0.62
3:FB:238:ASN:HB3	3:FB:241:HIS:HD2	1.64	0.62
2:KB:186:GLN:HB3	3:RB:74:ARG:NH1	2.14	0.62
3:RB:121:SER:O	3:RB:125:ILE:HG13	1.99	0.62
2:WB:37:LEU:O	2:WB:41:MET:HG3	2.00	0.62
3:DC:121:SER:O	3:DC:125:ILE:HG13	1.99	0.62
3:DF:127:VAL:O	3:DF:131:PHE:HB2	1.99	0.62
2:IF:151:GLU:O	2:IF:155:HIS:ND1	2.32	0.62
2:IF:212:GLU:O	2:IF:216:THR:HG22	2.00	0.62
2:MG:187:ASN:O	2:MG:189:LYS:NZ	2.32	0.62
2:B:186:GLN:HB3	3:K:74:ARG:NH1	2.14	0.62
3:BA:121:SER:O	3:BA:125:ILE:HG13	1.99	0.62
2:EB:97:ILE:O	2:EB:101:ARG:NH2	2.27	0.62
2:QB:212:GLU:O	2:QB:216:THR:HG22	2.00	0.62
2:WB:69:LEU:HD11	2:CC:47:ILE:HG22	1.80	0.62
2:AD:186:GLN:HB3	3:HD:74:ARG:HH12	1.65	0.62
2:MD:96:ASP:OD2	2:SD:35:GLN:NE2	2.33	0.62
3:ND:121:SER:O	3:ND:125:ILE:HG13	1.99	0.62
3:ND:207:GLU:OE1	3:ND:207:GLU:N	2.32	0.62
2:SD:160:ARG:O	2:SD:164:PHE:CB	2.48	0.62

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:DF:198:ILE:HG23	3:DF:221:PHE:CD1	2.34	0.62
2:AG:151:GLU:HA	2:AG:154:ARG:HG2	1.82	0.62
2:GG:212:GLU:O	2:GG:216:THR:HG22	1.98	0.62
2:SG:90:ALA:HA	2:SG:93:LEU:HG	1.81	0.62
2:AA:198:THR:O	2:AA:202:ILE:HG12	2.00	0.62
3:NA:57:ILE:HD11	3:NA:169:TRP:HA	1.79	0.62
2:YA:129:ILE:HG23	2:YA:175:LEU:HD21	1.80	0.62
2:EB:160:ARG:O	2:EB:164:PHE:HB2	1.99	0.62
3:LB:53:ALA:HB3	3:LB:231:LEU:HA	1.81	0.62
3:LB:54:LEU:HA	3:LB:57:ILE:HG22	1.82	0.62
3:JC:229:GLU:OE1	3:JC:230:LEU:HD12	2.00	0.62
3:VC:121:SER:O	3:VC:125:ILE:HG13	1.99	0.62
2:MD:120:ASP:OD1	2:MD:123:ARG:NH1	2.33	0.62
3:ND:115:LEU:HD21	3:ND:218:CYS:HB3	1.81	0.62
2:EE:21:ARG:NH2	2:EE:24:GLU:HB2	2.15	0.62
2:EE:212:GLU:O	2:EE:216:THR:HG22	2.00	0.62
2:WE:160:ARG:O	2:WE:164:PHE:CB	2.47	0.62
3:JF:54:LEU:HA	3:JF:57:ILE:HG22	1.80	0.62
3:VF:121:SER:O	3:VF:125:ILE:HG13	2.00	0.62
2:AG:125:GLU:HA	3:BG:147:THR:HG21	1.81	0.62
3:BG:121:SER:O	3:BG:125:ILE:HG13	1.99	0.62
2:GG:67:ALA:O	2:GG:71:ILE:HG12	1.99	0.62
3:TG:54:LEU:HA	3:TG:57:ILE:HG22	1.81	0.62
3:M:57:ILE:HD11	3:M:169:TRP:HA	1.82	0.62
2:AA:26:PHE:CZ	2:AA:37:LEU:HD11	2.35	0.62
2:SA:130:ILE:O	2:SA:134:LEU:HG	2.00	0.62
2:EB:207:LYS:HD3	2:EB:210:GLN:HE21	1.64	0.62
2:WB:129:ILE:HG23	2:WB:175:LEU:HD21	1.82	0.62
3:XB:121:SER:O	3:XB:125:ILE:HG13	2.00	0.62
2:CC:21:ARG:HE	2:CC:21:ARG:C	2.02	0.62
3:JC:54:LEU:HA	3:JC:57:ILE:HG22	1.82	0.62
3:PC:121:SER:O	3:PC:125:ILE:HG13	2.00	0.62
2:UC:108:GLU:HA	2:UC:111:ASN:HD22	1.65	0.62
2:MD:125:GLU:HA	3:ND:147:THR:HG21	1.82	0.62
2:SD:67:ALA:O	2:SD:71:ILE:HG12	2.00	0.62
3:ZD:54:LEU:HA	3:ZD:57:ILE:HG22	1.80	0.62
3:FE:169:TRP:HD1	3:FE:175:LEU:HD13	1.64	0.62
2:CF:93:LEU:O	2:CF:97:ILE:HG12	1.98	0.62
2:OF:118:ALA:HA	2:OF:121:LEU:HG	1.81	0.62
2:GG:111:ASN:HA	2:GG:141:GLN:NE2	2.15	0.62
2:GG:114:GLU:HB2	2:GG:117:SER:OG	1.99	0.62

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:MG:160:ARG:O	2:MG:164:PHE:CB	2.47	0.62
3:BA:54:LEU:HA	3:BA:57:ILE:HG22	1.81	0.62
2:GA:186:GLN:HB3	3:NA:74:ARG:NH1	2.15	0.62
3:NA:72:LEU:HD23	3:NA:73:LEU:HD12	1.82	0.62
2:KB:114:GLU:HB2	2:KB:117:SER:OG	2.00	0.62
3:XB:75:ARG:HG3	3:XB:208:ILE:HD11	1.82	0.62
3:XB:169:TRP:HD1	3:XB:175:LEU:HD13	1.64	0.62
2:CC:186:GLN:HB3	3:JC:74:ARG:NH1	2.15	0.62
3:BD:57:ILE:HD11	3:BD:169:TRP:HA	1.81	0.62
2:GD:207:LYS:NZ	2:GD:210:GLN:HG3	2.15	0.62
3:TD:57:ILE:HD11	3:TD:169:TRP:HA	1.81	0.62
3:LE:183:GLU:OE1	3:LE:185:GLN:N	2.32	0.62
3:RE:96:LEU:HD23	3:RE:97:PRO:HD2	1.81	0.62
2:WE:89:ARG:HH22	2:WE:93:LEU:HG	1.63	0.62
2:CF:83:LYS:HD3	2:IF:26:PHE:CD1	2.34	0.62
2:IF:207:LYS:HZ3	2:IF:209:GLN:HB3	1.64	0.62
2:SG:188:LEU:HD23	2:SG:188:LEU:H	1.64	0.62
2:B:186:GLN:HB3	3:K:74:ARG:HH12	1.64	0.62
2:J:160:ARG:O	2:J:164:PHE:HB2	1.98	0.62
2:MA:186:GLN:HE22	3:TA:71:ASN:HA	1.64	0.62
3:RB:120:PRO:HA	3:RB:123:VAL:HG22	1.80	0.62
3:JC:116:VAL:HG21	3:JC:158:LEU:HD11	1.81	0.62
2:OC:219:ARG:NH2	2:OC:227:GLN:OE1	2.27	0.62
2:KE:186:GLN:HB3	3:RE:74:ARG:NH1	2.12	0.62
2:QE:201:GLU:OE1	2:WE:139:ARG:NH1	2.33	0.62
2:OF:186:GLN:HE21	3:VF:74:ARG:NH2	1.98	0.62
2:OF:230:ILE:HA	2:OF:233:MET:HE2	1.81	0.62
3:PF:169:TRP:HD1	3:PF:175:LEU:HD13	1.64	0.62
3:TG:121:SER:O	3:TG:125:ILE:HG13	2.00	0.62
2:B:205:LEU:HD21	2:J:140:SER:HA	1.81	0.62
3:C:198:ILE:HG23	3:C:221:PHE:CD1	2.35	0.62
3:K:198:ILE:HG23	3:K:221:PHE:CD1	2.34	0.62
3:JF:238:ASN:HB3	3:JF:241:HIS:CD2	2.35	0.62
2:GG:151:GLU:HA	2:GG:154:ARG:HG2	1.82	0.62
2:SG:21:ARG:O	2:SG:21:ARG:NE	2.24	0.62
2:SG:107:ILE:HG23	2:SG:137:LEU:HD23	1.80	0.62
2:J:186:GLN:HB3	3:V:74:ARG:NH1	2.15	0.62
3:V:198:ILE:HG23	3:V:221:PHE:CD1	2.35	0.62
2:AA:31:THR:O	2:AA:34:VAL:HB	2.00	0.62
3:FB:229:GLU:OE1	3:FB:230:LEU:HD12	1.99	0.62
2:AD:186:GLN:NE2	3:HD:71:ASN:HA	2.13	0.62

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:MD:129:ILE:HG23	2:MD:175:LEU:HD21	1.82	0.62
2:EE:37:LEU:O	2:EE:41:MET:HG3	2.00	0.62
3:XE:121:SER:O	3:XE:125:ILE:HG13	2.00	0.62
2:CF:153:LEU:O	2:CF:157:VAL:HG23	2.00	0.62
2:OF:65:GLN:NE2	2:UF:47:ILE:O	2.29	0.62
2:UF:37:LEU:O	2:UF:41:MET:HG3	2.00	0.62
2:AG:234:PHE:HB3	2:AG:313:ARG:HH21	1.65	0.62
3:BG:112:GLY:HA3	3:BG:223:MET:HG2	1.81	0.62
2:GA:204:ASN:OD1	2:GA:233:MET:HB3	1.99	0.61
2:SA:125:GLU:HA	3:TA:147:THR:HG21	1.81	0.61
3:ZA:111:ARG:NH1	3:ZA:111:ARG:HB2	2.15	0.61
2:KB:152:ARG:HA	2:KB:155:HIS:CE1	2.35	0.61
2:KB:187:ASN:O	2:KB:189:LYS:NZ	2.33	0.61
3:XB:90:HIS:O	3:XB:94:ARG:HG2	2.00	0.61
2:OC:72:ASN:ND2	2:UC:41:MET:HB2	2.15	0.61
2:OC:129:ILE:HG23	2:OC:175:LEU:HD21	1.81	0.61
3:TD:72:LEU:HD23	3:TD:73:LEU:HD12	1.81	0.61
2:WE:26:PHE:CE1	2:WE:34:VAL:HG22	2.31	0.61
2:AG:168:GLN:HE22	3:BG:130:LEU:HD11	1.65	0.61
2:GG:129:ILE:HG23	2:GG:175:LEU:HD21	1.80	0.61
2:GG:152:ARG:HA	2:GG:155:HIS:CE1	2.35	0.61
2:B:83:LYS:HD3	2:J:26:PHE:CG	2.35	0.61
2:T:212:GLU:O	2:T:216:THR:HG22	2.00	0.61
2:CC:83:LYS:HD3	2:IC:26:PHE:HB3	1.82	0.61
3:DC:127:VAL:O	3:DC:131:PHE:HB2	2.00	0.61
2:IC:75:GLU:O	2:IC:79:SER:HB3	2.00	0.61
2:IC:186:GLN:NE2	3:PC:71:ASN:HA	2.12	0.61
3:VC:169:TRP:HD1	3:VC:175:LEU:HD13	1.64	0.61
2:AD:168:GLN:HE22	3:BD:130:LEU:HD11	1.65	0.61
2:GD:79:SER:HB2	2:GD:83:LYS:HE2	1.83	0.61
2:QE:12:ILE:HD11	2:QE:55:VAL:HG11	1.82	0.61
3:HG:121:SER:O	3:HG:125:ILE:HG13	2.00	0.61
2:SG:212:GLU:O	2:SG:216:THR:HG22	2.00	0.61
2:KB:234:PHE:HB3	2:KB:313:ARG:HH21	1.64	0.61
2:QB:168:GLN:HE22	3:RB:130:LEU:HD11	1.65	0.61
3:DC:54:LEU:HA	3:DC:57:ILE:HG22	1.83	0.61
3:DC:198:ILE:HG23	3:DC:221:PHE:CD1	2.35	0.61
2:IC:151:GLU:HA	2:IC:154:ARG:HG2	1.81	0.61
2:IC:186:GLN:HB3	3:PC:74:ARG:HH12	1.65	0.61
3:JC:72:LEU:HD23	3:JC:73:LEU:HD12	1.80	0.61
2:IF:125:GLU:HA	3:JF:147:THR:HG21	1.82	0.61

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:OF:69:LEU:HD21	2:UF:47:ILE:HB	1.82	0.61
2:SG:107:ILE:HG22	2:SG:111:ASN:HD21	1.65	0.61
3:M:225:GLU:HA	3:M:228:ARG:HB3	1.82	0.61
2:YA:74:ASN:O	2:YA:78:ARG:NE	2.34	0.61
2:YA:107:ILE:HG23	2:YA:137:LEU:HD23	1.83	0.61
3:RB:53:ALA:HB3	3:RB:231:LEU:HA	1.83	0.61
2:AD:151:GLU:HA	2:AD:154:ARG:HG2	1.82	0.61
2:AD:152:ARG:HA	2:AD:155:HIS:CE1	2.35	0.61
2:AD:186:GLN:HB3	3:HD:74:ARG:NH1	2.16	0.61
2:YD:186:GLN:HB3	3:FE:74:ARG:NH1	2.14	0.61
3:ZD:169:TRP:HD1	3:ZD:175:LEU:HD13	1.65	0.61
2:KE:130:ILE:O	2:KE:134:LEU:HG	1.99	0.61
2:OF:20:ASP:OD2	2:OF:21:ARG:N	2.33	0.61
2:SG:207:LYS:NZ	2:SG:209:GLN:OE1	2.33	0.61
2:B:35:GLN:O	2:B:38:SER:OG	2.18	0.61
2:SA:67:ALA:O	2:SA:71:ILE:HG12	2.00	0.61
2:SA:186:GLN:HB3	3:ZA:74:ARG:NH1	2.16	0.61
2:OC:11:VAL:HG11	2:OC:41:MET:HB3	1.83	0.61
3:PC:225:GLU:HA	3:PC:228:ARG:HB3	1.82	0.61
4:YC:107:ALA:HA	4:YC:125:ILE:HA	1.81	0.61
2:GD:186:GLN:HB3	3:ND:74:ARG:NH1	2.14	0.61
2:MD:273:LYS:HD2	2:MD:276:ARG:HH21	1.66	0.61
2:WE:69:LEU:HD13	2:CF:45:ARG:O	2.01	0.61
2:OF:160:ARG:O	2:OF:164:PHE:CB	2.48	0.61
3:VF:169:TRP:HD1	3:VF:175:LEU:HD13	1.65	0.61
2:GG:20:ASP:OD1	2:GG:21:ARG:N	2.33	0.61
2:J:205:LEU:HD21	2:T:140:SER:HA	1.81	0.61
2:MA:76:TYR:CE2	2:SA:38:SER:HB3	2.35	0.61
2:YA:186:GLN:HB3	3:FB:74:ARG:NH1	2.16	0.61
3:LB:169:TRP:HD1	3:LB:175:LEU:HD13	1.65	0.61
2:QB:20:ASP:OD2	2:QB:21:ARG:N	2.32	0.61
2:CC:130:ILE:O	2:CC:134:LEU:HG	2.01	0.61
3:VC:127:VAL:O	3:VC:131:PHE:HB2	2.01	0.61
2:AD:40:ALA:O	2:AD:44:VAL:HG23	2.00	0.61
2:AD:125:GLU:HA	3:BD:147:THR:HG21	1.83	0.61
3:ZD:198:ILE:HG23	3:ZD:221:PHE:CD1	2.36	0.61
2:QE:125:GLU:HA	3:RE:147:THR:HG21	1.82	0.61
2:WE:207:LYS:NZ	2:WE:209:GLN:OE1	2.33	0.61
3:DF:121:SER:O	3:DF:125:ILE:HG13	2.00	0.61
3:BG:127:VAL:O	3:BG:131:PHE:HB2	2.00	0.61
3:BG:198:ILE:HG23	3:BG:221:PHE:CD1	2.35	0.61

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:MG:310:LEU:HA	2:MG:313:ARG:HG2	1.83	0.61
2:SG:34:VAL:O	2:SG:38:SER:OG	2.14	0.61
2:B:152:ARG:NH2	2:B:153:LEU:HB2	2.14	0.61
2:AA:137:LEU:HB3	2:AA:141:GLN:NE2	2.15	0.61
2:GA:21:ARG:NH2	2:GA:24:GLU:HB2	2.15	0.61
3:HA:118:PHE:CE2	3:HA:158:LEU:HD12	2.36	0.61
2:YA:234:PHE:HB3	2:YA:313:ARG:HH21	1.63	0.61
3:FB:169:TRP:HD1	3:FB:175:LEU:HD13	1.65	0.61
3:LB:100:THR:HG23	3:LB:119:SER:HA	1.83	0.61
2:CC:110:LEU:O	2:CC:141:GLN:NE2	2.34	0.61
2:UC:206:MET:HB3	2:UC:210:GLN:NE2	2.15	0.61
3:VC:238:ASN:HB3	3:VC:241:HIS:ND1	2.16	0.61
2:EE:26:PHE:CE1	2:EE:34:VAL:HG22	2.32	0.61
2:EE:188:LEU:HD23	2:EE:188:LEU:H	1.66	0.61
3:LE:72:LEU:HD23	3:LE:73:LEU:HD12	1.82	0.61
3:RE:112:GLY:HA3	3:RE:223:MET:HG2	1.82	0.61
2:CF:234:PHE:HB3	2:CF:313:ARG:HH21	1.65	0.61
2:IF:130:ILE:O	2:IF:134:LEU:HG	2.01	0.61
3:JF:121:SER:O	3:JF:125:ILE:HG13	2.01	0.61
2:OF:21:ARG:NH2	2:OF:24:GLU:HB2	2.15	0.61
2:AG:80:VAL:O	2:AG:83:LYS:HG2	2.01	0.61
2:AG:130:ILE:O	2:AG:134:LEU:HG	2.00	0.61
2:B:245:ARG:HH12	2:B:249:ARG:HB2	1.65	0.61
2:J:202:ILE:HA	2:T:139:ARG:HH21	1.64	0.61
3:K:119:SER:HB3	3:K:214:GLU:O	2.01	0.61
2:GA:198:THR:O	2:GA:202:ILE:HG12	2.00	0.61
3:HA:121:SER:O	3:HA:125:ILE:HG13	2.01	0.61
2:YA:12:ILE:HD11	2:YA:55:VAL:HG11	1.83	0.61
2:EB:21:ARG:O	2:EB:21:ARG:NE	2.23	0.61
2:EB:244:ASP:OD1	2:EB:276:ARG:NH2	2.33	0.61
2:KB:201:GLU:HB3	2:QB:139:ARG:HH22	1.66	0.61
2:QB:186:GLN:HB3	3:XB:74:ARG:NH1	2.16	0.61
2:WB:168:GLN:HE22	3:XB:130:LEU:HD11	1.66	0.61
2:IC:160:ARG:O	2:IC:164:PHE:CB	2.49	0.61
2:MD:201:GLU:CD	2:SD:139:ARG:HH21	2.03	0.61
2:YD:125:GLU:HA	3:ZD:147:THR:HG21	1.82	0.61
2:QE:245:ARG:HH12	2:QE:249:ARG:HB2	1.65	0.61
2:OF:95:GLU:O	2:OF:99:GLU:HB2	1.99	0.61
2:G:21:ARG:HH21	2:G:24:GLU:HB2	1.65	0.61
2:G:113:MET:O	2:G:141:GLN:NE2	2.33	0.61
3:C:169:TRP:HD1	3:C:175:LEU:HD13	1.65	0.61

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:MA:160:ARG:O	2:MA:164:PHE:HB2	1.99	0.61
2:YA:186:GLN:HB3	3:FB:74:ARG:HH12	1.66	0.61
3:ZA:115:LEU:HD21	3:ZA:218:CYS:HB3	1.83	0.61
2:KB:186:GLN:HB3	3:RB:74:ARG:HH12	1.64	0.61
2:QB:207:LYS:NZ	2:QB:210:GLN:HG3	2.14	0.61
3:JC:127:VAL:O	3:JC:131:PHE:HB2	2.00	0.61
2:OC:160:ARG:O	2:OC:164:PHE:CB	2.49	0.61
2:MD:168:GLN:HE22	3:ND:130:LEU:HD11	1.66	0.61
3:LE:127:VAL:O	3:LE:131:PHE:HB2	2.01	0.61
2:GG:93:LEU:O	2:GG:97:ILE:HG12	2.00	0.61
2:G:125:GLU:HA	3:M:147:THR:HG21	1.83	0.61
2:J:273:LYS:HD2	2:J:276:ARG:HH21	1.65	0.61
3:K:127:VAL:O	3:K:131:PHE:HB2	1.99	0.61
3:V:121:SER:O	3:V:125:ILE:HG13	2.00	0.61
3:HA:183:GLU:HG3	3:HA:188:PHE:HB2	1.82	0.61
3:LB:121:SER:O	3:LB:125:ILE:HG13	2.00	0.61
3:XB:54:LEU:HA	3:XB:57:ILE:HG22	1.83	0.61
2:CC:21:ARG:HH21	2:CC:24:GLU:HB2	1.63	0.61
2:CC:125:GLU:HA	3:DC:147:THR:HG21	1.82	0.61
2:CC:186:GLN:HB3	3:JC:74:ARG:HH12	1.65	0.61
2:CC:201:GLU:CD	2:IC:139:ARG:HH12	2.03	0.61
2:IC:74:ASN:O	2:IC:78:ARG:NE	2.34	0.61
2:IC:152:ARG:HA	2:IC:155:HIS:CE1	2.36	0.61
2:OC:152:ARG:NH2	2:OC:153:LEU:HB2	2.16	0.61
2:AD:212:GLU:O	2:AD:216:THR:HG22	2.01	0.61
2:GD:230:ILE:HA	2:GD:233:MET:HE2	1.82	0.61
3:XE:105:ILE:HG12	3:XE:162:LEU:HD11	1.82	0.61
2:OF:168:GLN:HE22	3:PF:130:LEU:HD11	1.65	0.61
3:TG:115:LEU:HD21	3:TG:218:CYS:HB3	1.82	0.61
3:TG:169:TRP:HD1	3:TG:175:LEU:HD13	1.65	0.61
3:BA:53:ALA:HB3	3:BA:231:LEU:HA	1.83	0.60
3:BA:198:ILE:HG23	3:BA:221:PHE:CD1	2.36	0.60
3:ZA:58:ASN:ND2	3:ZA:59:GLU:OE1	2.34	0.60
3:FB:72:LEU:HD23	3:FB:73:LEU:HD12	1.83	0.60
3:FB:238:ASN:HB3	3:FB:241:HIS:CD2	2.35	0.60
2:KB:158:MET:SD	2:KB:159:LEU:HD13	2.41	0.60
3:RB:169:TRP:HD1	3:RB:175:LEU:HD13	1.66	0.60
3:DC:54:LEU:HB2	3:DC:231:LEU:HD23	1.82	0.60
2:UC:71:ILE:O	2:UC:74:ASN:ND2	2.34	0.60
3:HD:198:ILE:HG23	3:HD:221:PHE:CD1	2.36	0.60
2:MD:26:PHE:HE1	2:MD:34:VAL:HG22	1.66	0.60

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:YD:129:ILE:HG23	2:YD:175:LEU:HD21	1.81	0.60
2:YD:168:GLN:HE22	3:ZD:130:LEU:HD11	1.66	0.60
2:KE:110:LEU:O	2:KE:141:GLN:NE2	2.30	0.60
3:XE:169:TRP:HD1	3:XE:175:LEU:HD13	1.66	0.60
3:BG:57:ILE:HD11	3:BG:169:TRP:HA	1.82	0.60
2:GG:230:ILE:HA	2:GG:233:MET:HE2	1.82	0.60
2:SG:67:ALA:O	2:SG:71:ILE:HG12	2.00	0.60
2:SG:75:GLU:HA	2:SG:78:ARG:HH21	1.67	0.60
2:T:187:ASN:O	2:T:189:LYS:NZ	2.33	0.60
3:TA:96:LEU:HD23	3:TA:97:PRO:HD2	1.82	0.60
2:QB:160:ARG:O	2:QB:164:PHE:HB2	2.01	0.60
2:CC:168:GLN:HE22	3:DC:130:LEU:HD11	1.66	0.60
2:GD:160:ARG:O	2:GD:164:PHE:HB2	2.00	0.60
3:HD:102:LEU:HD22	3:HD:117:VAL:HG22	1.82	0.60
2:KE:26:PHE:CE1	2:KE:34:VAL:HG22	2.35	0.60
2:KE:26:PHE:HE1	2:KE:34:VAL:HG22	1.66	0.60
2:CF:21:ARG:NH2	2:CF:24:GLU:HB2	2.16	0.60
3:JF:127:VAL:O	3:JF:131:PHE:HB2	2.01	0.60
2:OF:30:SER:OG	2:OF:33:GLU:OE1	2.15	0.60
2:GG:245:ARG:HH12	2:GG:249:ARG:HB2	1.65	0.60
3:TG:57:ILE:HD11	3:TG:169:TRP:HA	1.83	0.60
3:V:72:LEU:HD23	3:V:73:LEU:HD12	1.83	0.60
3:V:112:GLY:HA3	3:V:223:MET:HG2	1.81	0.60
2:YA:190:ARG:O	2:EB:166:GLY:HA3	2.01	0.60
2:EB:113:MET:HG3	2:EB:117:SER:HB2	1.83	0.60
3:PC:101:ASN:ND2	3:PC:154:ILE:HG21	2.17	0.60
2:MD:207:LYS:HZ1	2:MD:209:GLN:HB3	1.65	0.60
3:FE:121:SER:O	3:FE:125:ILE:HG13	2.01	0.60
3:FE:191:ILE:HG13	3:FE:192:THR:HG23	1.83	0.60
2:WE:41:MET:HG2	2:WE:42:ALA:N	2.16	0.60
2:OF:212:GLU:O	2:OF:216:THR:HG22	2.01	0.60
2:AG:75:GLU:HA	2:AG:78:ARG:HH21	1.66	0.60
3:BG:54:LEU:HA	3:BG:57:ILE:HG22	1.83	0.60
3:TG:127:VAL:O	3:TG:131:PHE:HB2	2.01	0.60
2:G:47:ILE:HB	2:SG:69:LEU:HD21	1.83	0.60
2:J:21:ARG:NH2	2:J:24:GLU:HB2	2.15	0.60
3:V:57:ILE:HD11	3:V:169:TRP:HA	1.81	0.60
3:BA:72:LEU:HD23	3:BA:73:LEU:HD12	1.83	0.60
2:EB:204:ASN:OD1	2:EB:233:MET:HB3	2.00	0.60
2:CC:160:ARG:O	2:CC:164:PHE:HB2	2.02	0.60
3:NG:57:ILE:HD11	3:NG:169:TRP:HA	1.83	0.60

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:SG:89:ARG:HG2	2:SG:92:SER:HB3	1.84	0.60
2:T:110:LEU:C	2:T:141:GLN:HE22	2.03	0.60
2:AA:127:PRO:HA	2:AA:130:ILE:HG12	1.81	0.60
2:MA:186:GLN:HB3	3:TA:74:ARG:HH12	1.66	0.60
3:RB:54:LEU:HA	3:RB:57:ILE:HG22	1.84	0.60
2:IC:230:ILE:HA	2:IC:233:MET:HE2	1.82	0.60
2:GD:186:GLN:NE2	3:ND:71:ASN:HA	2.14	0.60
2:GD:204:ASN:OD1	2:GD:233:MET:HB3	2.02	0.60
2:EE:187:ASN:O	2:EE:189:LYS:NZ	2.30	0.60
2:KE:188:LEU:HD23	2:KE:188:LEU:H	1.67	0.60
3:DF:169:TRP:HD1	3:DF:175:LEU:HD13	1.67	0.60
2:OF:21:ARG:O	2:OF:21:ARG:NE	2.20	0.60
2:OF:26:PHE:CE1	2:OF:34:VAL:HG22	2.35	0.60
3:PF:121:SER:O	3:PF:125:ILE:HG13	2.00	0.60
2:UF:202:ILE:HD13	2:AG:139:ARG:NH1	2.06	0.60
2:UF:310:LEU:HA	2:UF:313:ARG:HG2	1.83	0.60
3:BG:169:TRP:HD1	3:BG:175:LEU:HD13	1.66	0.60
2:G:89:ARG:NH2	2:B:31:THR:HG23	2.15	0.60
2:B:130:ILE:O	2:B:134:LEU:HG	2.01	0.60
2:J:93:LEU:O	2:J:97:ILE:HG12	2.01	0.60
2:AA:21:ARG:O	2:AA:21:ARG:NE	2.24	0.60
2:GA:187:ASN:O	2:GA:189:LYS:NZ	2.34	0.60
2:QB:230:ILE:HA	2:QB:233:MET:CE	2.32	0.60
3:RB:102:LEU:HD22	3:RB:117:VAL:HG22	1.81	0.60
2:CC:63:ALA:HA	2:CC:66:PHE:CZ	2.37	0.60
2:OC:93:LEU:O	2:OC:97:ILE:HG12	2.02	0.60
3:PC:127:VAL:O	3:PC:131:PHE:HB2	2.01	0.60
2:UC:47:ILE:HD11	2:UC:52:LEU:HB2	1.83	0.60
3:HD:106:HIS:CD2	3:HD:113:THR:HG22	2.36	0.60
3:ND:182:SER:OG	3:ND:183:GLU:N	2.34	0.60
2:SD:110:LEU:HB3	2:SD:141:GLN:HE22	1.67	0.60
2:QE:67:ALA:O	2:QE:71:ILE:HG12	2.01	0.60
2:J:245:ARG:HH12	2:J:249:ARG:HB2	1.65	0.60
2:SA:186:GLN:HB3	3:ZA:74:ARG:HH12	1.66	0.60
2:YA:198:THR:O	2:YA:202:ILE:HG12	2.02	0.60
2:IC:67:ALA:O	2:IC:71:ILE:HG12	2.01	0.60
2:IC:114:GLU:HB2	2:IC:117:SER:OG	2.01	0.60
2:IC:245:ARG:HH12	2:IC:249:ARG:HB2	1.65	0.60
3:PC:198:ILE:HG23	3:PC:221:PHE:CD1	2.36	0.60
2:UC:186:GLN:HB3	3:BD:74:ARG:NH1	2.17	0.60
2:SD:125:GLU:HA	3:TD:147:THR:HG21	1.83	0.60

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:KE:234:PHE:HB3	2:KE:313:ARG:HH21	1.65	0.60
2:WE:30:SER:OG	2:WE:33:GLU:OE1	2.17	0.60
3:VF:183:GLU:OE1	3:VF:185:GLN:N	2.34	0.60
3:VF:238:ASN:HB3	3:VF:241:HIS:ND1	2.17	0.60
2:MG:152:ARG:NH2	2:MG:153:LEU:HB2	2.16	0.60
2:SG:37:LEU:O	2:SG:41:MET:HG3	2.01	0.60
2:GA:130:ILE:O	2:GA:134:LEU:HG	2.02	0.60
2:MA:125:GLU:HA	3:NA:147:THR:HG21	1.82	0.60
2:SA:160:ARG:O	2:SA:164:PHE:CB	2.50	0.60
2:YA:113:MET:O	2:YA:141:GLN:NE2	2.35	0.60
3:FB:57:ILE:HD11	3:FB:169:TRP:HA	1.82	0.60
2:KB:115:PRO:O	2:KB:119:ALA:CB	2.50	0.60
2:QB:90:ALA:HA	2:QB:93:LEU:HG	1.84	0.60
2:UC:245:ARG:HH12	2:UC:249:ARG:HB2	1.66	0.60
2:MD:245:ARG:HH12	2:MD:249:ARG:HB2	1.66	0.60
2:WE:67:ALA:O	2:WE:71:ILE:HG12	2.00	0.60
2:WE:93:LEU:O	2:WE:97:ILE:HG12	2.00	0.60
3:DF:115:LEU:HD11	3:DF:218:CYS:HB3	1.83	0.60
3:JF:198:ILE:HG23	3:JF:221:PHE:CD1	2.36	0.60
2:G:26:PHE:HE1	2:G:34:VAL:HG22	1.65	0.60
3:V:182:SER:OG	3:V:183:GLU:N	2.33	0.60
3:HA:119:SER:HB3	3:HA:214:GLU:O	2.02	0.60
3:TA:127:VAL:O	3:TA:131:PHE:HB2	2.01	0.60
2:EB:205:LEU:HD21	2:KB:140:SER:HA	1.82	0.60
2:QB:37:LEU:O	2:QB:41:MET:HG3	2.01	0.60
2:IC:107:ILE:HA	2:IC:110:LEU:HD12	1.84	0.60
3:HD:115:LEU:HD21	3:HD:218:CYS:HB3	1.83	0.60
2:YD:26:PHE:CZ	2:YD:34:VAL:HA	2.37	0.60
2:QE:107:ILE:HG22	2:QE:111:ASN:HD21	1.67	0.60
2:UF:188:LEU:HD13	3:BG:74:ARG:HE	1.66	0.60
3:VF:59:GLU:N	3:VF:59:GLU:OE1	2.35	0.60
2:AG:113:MET:HE2	2:AG:121:LEU:HD11	1.84	0.60
2:MG:93:LEU:O	2:MG:97:ILE:HG12	2.00	0.60
2:J:89:ARG:HG3	2:J:90:ALA:N	2.16	0.60
2:J:168:GLN:HE22	3:K:130:LEU:HD11	1.66	0.60
2:J:212:GLU:O	2:J:216:THR:HG22	2.01	0.60
2:T:96:ASP:OD2	2:AA:35:GLN:NE2	2.34	0.60
2:GA:245:ARG:HH12	2:GA:249:ARG:HB2	1.67	0.60
2:SA:212:GLU:O	2:SA:216:THR:HG22	2.02	0.60
2:EB:69:LEU:HD11	2:KB:47:ILE:HG22	1.83	0.60
2:EB:111:ASN:HA	2:EB:141:GLN:OE1	2.01	0.60

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:EB:245:ARG:HH12	2:EB:249:ARG:HB2	1.65	0.60
2:IC:125:GLU:HA	3:JC:147:THR:HG21	1.84	0.60
2:GD:245:ARG:HH12	2:GD:249:ARG:HB2	1.66	0.60
2:MD:65:GLN:NE2	2:SD:47:ILE:O	2.30	0.60
2:MD:186:GLN:HB3	3:TD:74:ARG:NH1	2.16	0.60
2:MD:187:ASN:O	2:MD:189:LYS:NZ	2.34	0.60
2:SD:168:GLN:HE22	3:TD:130:LEU:HD11	1.66	0.60
2:SD:245:ARG:HH12	2:SD:249:ARG:HB2	1.67	0.60
3:TD:127:VAL:O	3:TD:131:PHE:HB2	2.02	0.60
3:TD:183:GLU:OE1	3:TD:185:GLN:N	2.35	0.60
2:YD:21:ARG:HE	2:YD:21:ARG:C	2.05	0.60
2:CF:21:ARG:O	2:CF:21:ARG:NE	2.21	0.60
3:DF:182:SER:OG	3:DF:183:GLU:N	2.35	0.60
3:VF:57:ILE:HD11	3:VF:169:TRP:HA	1.84	0.60
2:GG:80:VAL:O	2:GG:83:LYS:HG2	2.02	0.60
3:HG:54:LEU:HA	3:HG:57:ILE:HG22	1.82	0.60
2:G:21:ARG:NH2	2:G:24:GLU:HB2	2.17	0.59
2:G:212:GLU:O	2:G:216:THR:HG22	2.02	0.59
3:M:127:VAL:O	3:M:131:PHE:HB2	2.02	0.59
2:T:108:GLU:HA	2:T:111:ASN:HD22	1.67	0.59
3:BA:174:PRO:O	3:BA:174:PRO:HD2	2.02	0.59
3:TA:240:ARG:NH1	3:TA:240:ARG:HA	2.17	0.59
2:YA:83:LYS:O	2:EB:27:LYS:HE2	2.02	0.59
2:YA:153:LEU:O	2:YA:157:VAL:HG23	2.02	0.59
3:LB:57:ILE:HD11	3:LB:169:TRP:HA	1.84	0.59
2:IC:93:LEU:O	2:IC:97:ILE:HG12	2.02	0.59
2:AD:190:ARG:O	2:GD:166:GLY:HA3	2.01	0.59
2:AD:273:LYS:HD2	2:AD:276:ARG:HH21	1.67	0.59
2:GD:21:ARG:NH2	2:GD:24:GLU:HB2	2.16	0.59
2:GD:230:ILE:HA	2:GD:233:MET:CE	2.32	0.59
2:SD:188:LEU:HB3	3:ZD:74:ARG:HE	1.66	0.59
2:KE:93:LEU:O	2:KE:97:ILE:HG12	2.02	0.59
2:IF:160:ARG:O	2:IF:164:PHE:CB	2.50	0.59
2:OF:273:LYS:HD2	2:OF:276:ARG:HH21	1.67	0.59
2:GG:97:ILE:O	2:GG:101:ARG:NH2	2.25	0.59
2:G:76:TYR:HE1	2:B:38:SER:HB3	1.67	0.59
2:G:245:ARG:HH12	2:G:249:ARG:HB2	1.67	0.59
2:T:93:LEU:O	2:T:97:ILE:HG12	2.02	0.59
2:T:198:THR:O	2:T:202:ILE:HG12	2.02	0.59
2:SA:245:ARG:HH12	2:SA:249:ARG:HB2	1.66	0.59
2:EB:168:GLN:NE2	3:FB:130:LEU:HD11	2.13	0.59

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:FB:75:ARG:HG3	3:FB:208:ILE:HD11	1.84	0.59
2:QB:83:LYS:HD2	2:WB:26:PHE:CD1	2.35	0.59
3:JC:169:TRP:HD1	3:JC:175:LEU:HD13	1.67	0.59
2:UC:186:GLN:NE2	3:BD:71:ASN:HA	2.16	0.59
2:GD:93:LEU:O	2:GD:97:ILE:HG12	2.01	0.59
2:GD:186:GLN:HB3	3:ND:74:ARG:HH12	1.67	0.59
2:GD:310:LEU:HA	2:GD:313:ARG:HG2	1.83	0.59
2:SD:11:VAL:HG11	2:SD:41:MET:HB3	1.84	0.59
3:RE:127:VAL:O	3:RE:131:PHE:HB2	2.02	0.59
2:OF:93:LEU:O	2:OF:97:ILE:HG12	2.01	0.59
2:AG:89:ARG:HD2	2:AG:90:ALA:H	1.67	0.59
2:GG:188:LEU:H	3:NG:74:ARG:NH2	2.00	0.59
3:HG:169:TRP:HD1	3:HG:175:LEU:HD13	1.68	0.59
2:MG:95:GLU:HG3	2:MG:112:PHE:CE2	2.37	0.59
2:G:273:LYS:HD2	2:G:276:ARG:HH21	1.67	0.59
2:T:186:GLN:HB3	3:BA:74:ARG:NH1	2.17	0.59
3:BA:236:LEU:HB3	3:BA:238:ASN:OD1	2.02	0.59
2:GA:186:GLN:HB3	3:NA:74:ARG:HH12	1.68	0.59
2:QB:310:LEU:HA	2:QB:313:ARG:HG2	1.84	0.59
2:WB:96:ASP:OD2	2:CC:35:GLN:NE2	2.35	0.59
2:CC:202:ILE:HD13	2:IC:139:ARG:NH2	2.17	0.59
2:OC:168:GLN:HE22	3:PC:130:LEU:HD11	1.67	0.59
2:GD:26:PHE:CE1	2:GD:34:VAL:HG22	2.38	0.59
2:MD:93:LEU:O	2:MD:97:ILE:HG12	2.01	0.59
3:ND:57:ILE:HD11	3:ND:169:TRP:HA	1.84	0.59
2:SD:127:PRO:HA	2:SD:130:ILE:HG12	1.83	0.59
3:ZD:75:ARG:HG3	3:ZD:208:ILE:HD11	1.84	0.59
3:FE:190:ASN:ND2	3:LE:234:PRO:O	2.35	0.59
2:KE:22:ALA:O	2:KE:26:PHE:HB2	2.01	0.59
3:DF:96:LEU:HD23	3:DF:97:PRO:HD2	1.83	0.59
2:IF:115:PRO:O	2:IF:119:ALA:CB	2.50	0.59
2:OF:107:ILE:HA	2:OF:110:LEU:HD12	1.83	0.59
2:UF:245:ARG:HH12	2:UF:249:ARG:HB2	1.67	0.59
2:SG:245:ARG:HH12	2:SG:249:ARG:HB2	1.67	0.59
2:G:230:ILE:HA	2:G:233:MET:HE2	1.84	0.59
2:J:152:ARG:NH2	2:J:153:LEU:HB2	2.18	0.59
2:GA:89:ARG:HD2	2:GA:90:ALA:H	1.67	0.59
2:MA:273:LYS:HD2	2:MA:276:ARG:HH21	1.67	0.59
2:SA:297:ARG:HG3	2:SA:299:SER:H	1.67	0.59
2:YA:76:TYR:OH	2:EB:34:VAL:HG13	2.02	0.59
2:KB:297:ARG:HG3	2:KB:299:SER:H	1.67	0.59

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:IC:11:VAL:HG11	2:IC:41:MET:HB3	1.83	0.59
2:AD:153:LEU:O	2:AD:157:VAL:HG23	2.03	0.59
2:AD:187:ASN:O	2:AD:189:LYS:NZ	2.35	0.59
3:BD:115:LEU:HD21	3:BD:218:CYS:HB3	1.85	0.59
2:GD:83:LYS:HD3	2:GD:83:LYS:N	2.17	0.59
2:SD:93:LEU:O	2:SD:97:ILE:HG12	2.01	0.59
2:WE:245:ARG:HH12	2:WE:249:ARG:HB2	1.66	0.59
2:IF:203:ILE:HA	2:IF:206:MET:HE2	1.83	0.59
2:OF:186:GLN:HB3	3:VF:74:ARG:NH1	2.17	0.59
2:AG:160:ARG:O	2:AG:164:PHE:HB2	2.01	0.59
3:HG:116:VAL:HG21	3:HG:158:LEU:HD11	1.84	0.59
2:SG:310:LEU:HA	2:SG:313:ARG:HG2	1.84	0.59
3:TA:118:PHE:CE2	3:TA:158:LEU:HD12	2.38	0.59
3:ZA:174:PRO:O	3:ZA:174:PRO:HD2	2.03	0.59
2:KB:85:LEU:HD22	2:KB:85:LEU:H	1.67	0.59
2:QB:273:LYS:HD2	2:QB:276:ARG:HH21	1.68	0.59
2:WB:160:ARG:O	2:WB:164:PHE:CB	2.50	0.59
3:XB:127:VAL:O	3:XB:131:PHE:HB2	2.03	0.59
2:CC:219:ARG:NH2	2:CC:227:GLN:OE1	2.34	0.59
2:IC:37:LEU:O	2:IC:41:MET:HG3	2.03	0.59
2:IC:204:ASN:OD1	2:IC:233:MET:HB3	2.02	0.59
3:PC:116:VAL:HG21	3:PC:158:LEU:HD11	1.85	0.59
2:AD:21:ARG:NH2	2:AD:24:GLU:HB2	2.17	0.59
2:SD:74:ASN:O	2:SD:78:ARG:NE	2.35	0.59
2:KE:89:ARG:HG3	2:KE:90:ALA:N	2.17	0.59
3:JF:112:GLY:HA3	3:JF:223:MET:HG2	1.84	0.59
3:PF:54:LEU:HA	3:PF:57:ILE:HG22	1.83	0.59
3:HG:101:ASN:ND2	3:HG:154:ILE:HG21	2.18	0.59
3:NG:174:PRO:HD2	3:NG:174:PRO:O	2.02	0.59
3:M:101:ASN:ND2	3:M:154:ILE:HG21	2.18	0.59
3:M:115:LEU:HD21	3:M:218:CYS:HB3	1.83	0.59
2:B:89:ARG:HG3	2:B:90:ALA:N	2.16	0.59
2:GA:78:ARG:O	2:GA:82:VAL:HG23	2.02	0.59
2:MA:245:ARG:HH12	2:MA:249:ARG:HB2	1.68	0.59
3:NA:121:SER:O	3:NA:125:ILE:HG13	2.03	0.59
2:SA:230:ILE:HA	2:SA:233:MET:CE	2.33	0.59
2:YA:245:ARG:HH12	2:YA:249:ARG:HB2	1.66	0.59
2:KB:83:LYS:HD2	2:QB:26:PHE:CD2	2.36	0.59
2:QB:93:LEU:O	2:QB:97:ILE:HG12	2.03	0.59
3:RB:111:ARG:HA	3:RB:111:ARG:CZ	2.33	0.59
2:WB:75:GLU:O	2:WB:79:SER:HB3	2.02	0.59

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:CC:151:GLU:HA	2:CC:154:ARG:HG2	1.83	0.59
2:IC:108:GLU:HA	2:IC:111:ASN:HD22	1.67	0.59
3:JC:97:PRO:O	3:JC:100:THR:OG1	2.17	0.59
3:PC:72:LEU:HD23	3:PC:73:LEU:HD12	1.82	0.59
2:MD:110:LEU:C	2:MD:141:GLN:HE22	2.05	0.59
2:MD:153:LEU:O	2:MD:157:VAL:HG23	2.02	0.59
2:IF:297:ARG:HG3	2:IF:299:SER:H	1.68	0.59
3:JF:169:TRP:HD1	3:JF:175:LEU:HD13	1.67	0.59
3:JF:182:SER:OG	3:JF:183:GLU:N	2.36	0.59
2:OF:245:ARG:HH12	2:OF:249:ARG:HB2	1.67	0.59
3:VF:127:VAL:O	3:VF:131:PHE:HB2	2.03	0.59
3:TG:174:PRO:HD2	3:TG:174:PRO:O	2.03	0.59
2:G:224:GLU:HA	2:G:227:GLN:OE1	2.03	0.59
2:T:160:ARG:O	2:T:164:PHE:CB	2.50	0.59
2:T:186:GLN:HB3	3:BA:74:ARG:HH12	1.68	0.59
2:T:273:LYS:HD2	2:T:276:ARG:HH21	1.67	0.59
2:AA:168:GLN:HE22	3:BA:130:LEU:HD11	1.67	0.59
2:AA:245:ARG:HH12	2:AA:249:ARG:HB2	1.67	0.59
2:GA:21:ARG:HE	2:GA:21:ARG:C	2.06	0.59
2:KB:151:GLU:HA	2:KB:154:ARG:HG2	1.84	0.59
2:WB:110:LEU:C	2:WB:141:GLN:HE22	2.06	0.59
3:XB:183:GLU:OE1	3:XB:185:GLN:N	2.35	0.59
2:YD:206:MET:SD	2:YD:207:LYS:N	2.76	0.59
2:EE:273:LYS:HD2	2:EE:276:ARG:HH21	1.68	0.59
2:KE:273:LYS:HD2	2:KE:276:ARG:HH21	1.68	0.59
3:DF:101:ASN:ND2	3:DF:154:ILE:HG21	2.18	0.59
2:OF:230:ILE:HA	2:OF:233:MET:CE	2.33	0.59
2:GG:83:LYS:O	2:MG:27:LYS:HE2	2.03	0.59
2:GG:186:GLN:NE2	3:NG:71:ASN:HA	2.17	0.59
2:G:310:LEU:HA	2:G:313:ARG:HG2	1.83	0.59
2:J:130:ILE:O	2:J:134:LEU:HG	2.03	0.59
2:T:245:ARG:HH12	2:T:249:ARG:HB2	1.66	0.59
3:HA:54:LEU:HA	3:HA:57:ILE:HG22	1.84	0.59
2:MA:37:LEU:O	2:MA:41:MET:HG3	2.03	0.59
2:SA:186:GLN:NE2	3:ZA:71:ASN:HA	2.18	0.59
2:YA:21:ARG:NH2	2:YA:24:GLU:HB2	2.17	0.59
2:KB:89:ARG:HG3	2:KB:90:ALA:N	2.17	0.59
3:LB:127:VAL:O	3:LB:131:PHE:HB2	2.02	0.59
3:LB:174:PRO:O	3:LB:174:PRO:HD2	2.03	0.59
3:RB:174:PRO:HD2	3:RB:174:PRO:O	2.03	0.59
2:CC:273:LYS:HD2	2:CC:276:ARG:HH21	1.68	0.59

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:IC:153:LEU:O	2:IC:157:VAL:HG23	2.03	0.59
2:OC:245:ARG:HH12	2:OC:249:ARG:HB2	1.66	0.59
2:UC:89:ARG:HH22	2:UC:93:LEU:HD21	1.67	0.59
2:AD:93:LEU:O	2:AD:97:ILE:HG12	2.03	0.59
3:TD:224:ILE:O	3:TD:228:ARG:N	2.36	0.59
2:QE:297:ARG:HG3	2:QE:299:SER:H	1.68	0.59
2:AG:89:ARG:HD2	2:AG:90:ALA:N	2.18	0.59
2:AG:273:LYS:HD2	2:AG:276:ARG:HH21	1.68	0.59
2:GG:153:LEU:O	2:GG:157:VAL:HG23	2.02	0.59
3:HG:127:VAL:O	3:HG:131:PHE:HB2	2.03	0.59
3:NG:225:GLU:HA	3:NG:228:ARG:HB3	1.84	0.59
2:SG:168:GLN:HE22	3:TG:130:LEU:HD11	1.65	0.59
2:G:26:PHE:CE2	2:SG:80:VAL:HG12	2.38	0.59
2:G:78:ARG:O	2:G:82:VAL:HG23	2.03	0.59
3:K:90:HIS:O	3:K:94:ARG:HG2	2.02	0.59
2:T:251:LEU:HD11	2:T:277:ASN:HD22	1.68	0.59
2:AA:74:ASN:O	2:AA:78:ARG:NE	2.36	0.59
2:AA:297:ARG:HG3	2:AA:299:SER:H	1.68	0.59
3:FB:174:PRO:HD2	3:FB:174:PRO:O	2.03	0.59
2:KB:89:ARG:HH22	2:KB:93:LEU:HG	1.66	0.59
2:KB:245:ARG:HH12	2:KB:249:ARG:HB2	1.66	0.59
2:CC:55:VAL:O	2:CC:59:PHE:HB3	2.03	0.59
3:DC:169:TRP:HD1	3:DC:175:LEU:HD13	1.66	0.59
2:UC:38:SER:O	2:UC:41:MET:HG2	2.03	0.59
3:VC:174:PRO:O	3:VC:174:PRO:HD2	2.03	0.59
3:HD:111:ARG:HH21	3:ND:241:HIS:CE1	2.21	0.59
2:MD:187:ASN:N	3:TD:74:ARG:HH22	2.00	0.59
3:ND:119:SER:HB3	3:ND:214:GLU:O	2.02	0.59
3:TD:115:LEU:HD21	3:TD:218:CYS:HB3	1.85	0.59
2:EE:297:ARG:HG3	2:EE:299:SER:H	1.68	0.59
3:XE:182:SER:OG	3:XE:183:GLU:N	2.36	0.59
2:CF:245:ARG:HH12	2:CF:249:ARG:HB2	1.65	0.59
3:PF:101:ASN:ND2	3:PF:154:ILE:HG21	2.18	0.59
3:PF:183:GLU:OE1	3:PF:185:GLN:N	2.36	0.59
2:UF:198:THR:O	2:UF:202:ILE:HG12	2.02	0.59
2:AG:89:ARG:HG2	2:AG:92:SER:HB3	1.85	0.59
3:HG:57:ILE:HD11	3:HG:169:TRP:HA	1.84	0.59
3:NG:116:VAL:HG21	3:NG:158:LEU:HD11	1.84	0.59
2:G:21:ARG:HE	2:G:21:ARG:C	2.04	0.59
3:C:72:LEU:HD23	3:C:73:LEU:HD12	1.85	0.59
3:C:174:PRO:HD2	3:C:174:PRO:O	2.03	0.59

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:J:201:GLU:HG3	2:T:139:ARG:CZ	2.33	0.59
2:GA:89:ARG:HD2	2:GA:90:ALA:N	2.18	0.59
3:ZA:169:TRP:HD1	3:ZA:175:LEU:HD13	1.68	0.59
3:LB:182:SER:OG	3:LB:183:GLU:N	2.33	0.59
2:QB:186:GLN:HB3	3:XB:74:ARG:HH12	1.66	0.59
3:JC:90:HIS:O	3:JC:94:ARG:HG2	2.02	0.59
2:UC:75:GLU:O	2:UC:79:SER:HB3	2.02	0.59
2:UC:186:GLN:HB3	3:BD:74:ARG:HH12	1.67	0.59
2:AD:80:VAL:O	2:AD:83:LYS:HG2	2.02	0.59
2:SD:79:SER:HA	2:SD:82:VAL:HG22	1.85	0.59
2:WE:113:MET:O	2:WE:141:GLN:NE2	2.36	0.59
2:CF:206:MET:HG2	2:IF:143:ALA:HB1	1.84	0.59
2:IF:151:GLU:HA	2:IF:154:ARG:HG2	1.85	0.59
2:AG:251:LEU:HD11	2:AG:277:ASN:HD22	1.68	0.59
3:HG:198:ILE:HG23	3:HG:221:PHE:CD1	2.37	0.59
2:B:273:LYS:HD2	2:B:276:ARG:HH21	1.67	0.58
4:R:106:ILE:CB	3:V:252:ARG:HD3	2.32	0.58
2:CC:230:ILE:HA	2:CC:233:MET:CE	2.33	0.58
3:JC:101:ASN:ND2	3:JC:154:ILE:HG21	2.18	0.58
2:UC:93:LEU:O	2:UC:97:ILE:HG12	2.03	0.58
2:AD:245:ARG:HH12	2:AD:249:ARG:HB2	1.67	0.58
3:BD:101:ASN:ND2	3:BD:154:ILE:HG21	2.18	0.58
3:TD:169:TRP:HD1	3:TD:175:LEU:HD13	1.68	0.58
3:FE:119:SER:HB3	3:FE:214:GLU:O	2.02	0.58
3:RE:174:PRO:HD2	3:RE:174:PRO:O	2.03	0.58
2:IF:80:VAL:HG12	2:OF:26:PHE:CE2	2.38	0.58
2:OF:83:LYS:HA	2:UF:23:ALA:HB1	1.84	0.58
2:OF:186:GLN:HB3	3:VF:74:ARG:HH12	1.68	0.58
2:UF:114:GLU:HB2	2:UF:117:SER:OG	2.03	0.58
2:UF:219:ARG:NH2	2:UF:227:GLN:OE1	2.30	0.58
3:VF:174:PRO:O	3:VF:174:PRO:HD2	2.03	0.58
2:GG:158:MET:HE3	2:GG:159:LEU:HG	1.83	0.58
3:TG:102:LEU:HD22	3:TG:117:VAL:HG22	1.85	0.58
2:G:83:LYS:O	2:B:27:LYS:HE3	2.03	0.58
3:M:120:PRO:HA	3:M:123:VAL:HG22	1.84	0.58
2:B:113:MET:HG3	2:B:117:SER:HB2	1.85	0.58
2:J:83:LYS:HD3	2:T:26:PHE:CG	2.38	0.58
2:GA:212:GLU:O	2:GA:216:THR:HG22	2.03	0.58
2:MA:96:ASP:OD2	2:SA:35:GLN:NE2	2.36	0.58
2:SA:219:ARG:NH2	2:SA:227:GLN:OE1	2.32	0.58
2:YA:158:MET:HE3	2:YA:159:LEU:HG	1.85	0.58

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:EB:74:ASN:O	2:EB:78:ARG:NE	2.36	0.58
2:WB:245:ARG:HH12	2:WB:249:ARG:HB2	1.67	0.58
2:OC:80:VAL:O	2:OC:83:LYS:HG2	2.03	0.58
2:OC:97:ILE:O	2:OC:101:ARG:NH2	2.30	0.58
2:OC:273:LYS:HD2	2:OC:276:ARG:HH21	1.69	0.58
2:UC:11:VAL:O	2:UC:15:MET:HE2	2.02	0.58
2:GD:21:ARG:HH21	2:GD:24:GLU:HB2	1.68	0.58
2:YD:207:LYS:NZ	2:YD:210:GLN:HG3	2.19	0.58
2:QE:129:ILE:HG23	2:QE:175:LEU:HD21	1.84	0.58
2:QE:230:ILE:HA	2:QE:233:MET:CE	2.32	0.58
3:XE:174:PRO:HD2	3:XE:174:PRO:O	2.03	0.58
2:UF:152:ARG:HH22	2:UF:153:LEU:HB2	1.68	0.58
2:AG:245:ARG:HH12	2:AG:249:ARG:HB2	1.68	0.58
2:MG:11:VAL:HG11	2:MG:41:MET:HB3	1.85	0.58
3:NG:101:ASN:ND2	3:NG:154:ILE:HG21	2.17	0.58
2:SG:109:THR:O	2:SG:113:MET:HG3	2.02	0.58
2:SG:153:LEU:O	2:SG:157:VAL:HG23	2.03	0.58
4:R:107:ALA:H	3:V:252:ARG:HE	1.52	0.58
3:HA:174:PRO:O	3:HA:174:PRO:HD2	2.03	0.58
2:YA:203:ILE:HA	2:YA:206:MET:SD	2.44	0.58
3:FB:96:LEU:HD23	3:FB:97:PRO:HD2	1.85	0.58
3:XB:57:ILE:HD11	3:XB:169:TRP:HA	1.84	0.58
2:UC:205:LEU:HD21	2:AD:140:SER:HA	1.85	0.58
3:VC:57:ILE:HD11	3:VC:169:TRP:HA	1.85	0.58
3:BD:174:PRO:O	3:BD:174:PRO:HD2	2.04	0.58
2:MD:33:GLU:O	2:MD:37:LEU:HG	2.03	0.58
4:QD:104:TYR:O	3:TD:252:ARG:NH1	2.35	0.58
2:YD:188:LEU:HB2	2:EE:168:GLN:HB3	1.85	0.58
3:ZD:101:ASN:ND2	3:ZD:154:ILE:HG21	2.19	0.58
3:FE:54:LEU:HA	3:FE:57:ILE:HG22	1.85	0.58
3:LE:121:SER:O	3:LE:125:ILE:HG13	2.03	0.58
2:WE:89:ARG:HG3	2:WE:90:ALA:N	2.18	0.58
2:WE:230:ILE:HA	2:WE:233:MET:CE	2.34	0.58
2:OF:33:GLU:O	2:OF:37:LEU:HG	2.02	0.58
2:OF:297:ARG:HG3	2:OF:299:SER:H	1.68	0.58
2:OF:310:LEU:HA	2:OF:313:ARG:HG2	1.84	0.58
3:PF:174:PRO:O	3:PF:174:PRO:HD2	2.04	0.58
3:HG:90:HIS:O	3:HG:94:ARG:HG2	2.04	0.58
2:MG:111:ASN:HA	2:MG:141:GLN:OE1	2.02	0.58
2:MG:245:ARG:HH12	2:MG:249:ARG:HB2	1.68	0.58
2:G:67:ALA:O	2:G:71:ILE:HG12	2.03	0.58

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:G:108:GLU:HA	2:G:111:ASN:HD22	1.68	0.58
2:GA:273:LYS:HD2	2:GA:276:ARG:HH21	1.68	0.58
2:MA:76:TYR:HB2	2:SA:41:MET:HE3	1.84	0.58
2:MA:251:LEU:HD11	2:MA:277:ASN:HD22	1.69	0.58
3:NA:120:PRO:HA	3:NA:123:VAL:HG22	1.85	0.58
2:SA:168:GLN:HE22	3:TA:130:LEU:HD11	1.68	0.58
3:TA:169:TRP:HD1	3:TA:175:LEU:HD13	1.68	0.58
3:ZA:101:ASN:ND2	3:ZA:154:ILE:HG21	2.19	0.58
2:KB:273:LYS:HD2	2:KB:276:ARG:HH21	1.68	0.58
3:LB:71:ASN:HD22	3:LB:160:LEU:HD21	1.68	0.58
3:LB:238:ASN:HB3	3:LB:241:HIS:HD2	1.69	0.58
2:QB:245:ARG:HH12	2:QB:249:ARG:HB2	1.67	0.58
2:QB:297:ARG:HG3	2:QB:299:SER:H	1.67	0.58
2:IC:130:ILE:O	2:IC:134:LEU:HG	2.02	0.58
3:PC:183:GLU:OE1	3:PC:184:MET:N	2.37	0.58
2:UC:158:MET:SD	2:UC:159:LEU:HD13	2.43	0.58
2:UC:168:GLN:NE2	3:VC:130:LEU:HD11	2.18	0.58
3:VC:59:GLU:OE1	3:VC:59:GLU:N	2.36	0.58
3:BD:72:LEU:HD23	3:BD:73:LEU:HD12	1.84	0.58
3:BD:169:TRP:HD1	3:BD:175:LEU:HD13	1.67	0.58
3:ND:101:ASN:ND2	3:ND:154:ILE:HG21	2.18	0.58
2:SD:83:LYS:HE3	2:YD:23:ALA:HA	1.85	0.58
2:SD:273:LYS:HD2	2:SD:276:ARG:HH21	1.68	0.58
2:YD:74:ASN:O	2:YD:78:ARG:NE	2.36	0.58
2:KE:113:MET:O	2:KE:141:GLN:NE2	2.37	0.58
2:WE:188:LEU:HB2	2:CF:168:GLN:HB3	1.85	0.58
2:CF:207:LYS:NZ	2:CF:210:GLN:HG3	2.19	0.58
3:JF:57:ILE:HD11	3:JF:169:TRP:HA	1.84	0.58
2:MG:219:ARG:NH2	2:MG:227:GLN:OE1	2.33	0.58
2:B:93:LEU:O	2:B:97:ILE:HG12	2.02	0.58
3:V:115:LEU:HD21	3:V:218:CYS:HB3	1.85	0.58
2:AA:125:GLU:HA	3:BA:147:THR:HG21	1.84	0.58
3:BA:112:GLY:HA3	3:BA:223:MET:HG2	1.84	0.58
2:SA:69:LEU:HD13	2:YA:45:ARG:O	2.02	0.58
3:TA:105:ILE:HG12	3:TA:162:LEU:HD11	1.85	0.58
3:RB:101:ASN:ND2	3:RB:154:ILE:HG21	2.18	0.58
2:CC:31:THR:O	2:CC:34:VAL:HB	2.04	0.58
2:CC:89:ARG:HG2	2:CC:92:SER:HB3	1.86	0.58
2:IC:84:ALA:O	2:IC:87:GLU:HB3	2.04	0.58
3:JC:57:ILE:HD11	3:JC:169:TRP:HA	1.85	0.58
3:JC:198:ILE:HG23	3:JC:221:PHE:CD1	2.39	0.58

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:BD:120:PRO:HA	3:BD:123:VAL:HG22	1.85	0.58
2:GD:244:ASP:OD1	2:GD:276:ARG:NH2	2.36	0.58
2:MD:74:ASN:O	2:MD:78:ARG:NE	2.36	0.58
3:TD:198:ILE:HG23	3:TD:221:PHE:HD1	1.67	0.58
2:YD:245:ARG:HH12	2:YD:249:ARG:HB2	1.68	0.58
3:ZD:183:GLU:OE1	3:ZD:185:GLN:N	2.37	0.58
2:EE:251:LEU:HD11	2:EE:277:ASN:HD22	1.69	0.58
3:LE:97:PRO:O	3:LE:100:THR:OG1	2.16	0.58
2:QE:93:LEU:O	2:QE:97:ILE:HG12	2.03	0.58
2:WE:193:MET:O	2:CF:164:PHE:HD1	1.86	0.58
2:WE:273:LYS:HD2	2:WE:276:ARG:HH21	1.68	0.58
2:CF:251:LEU:HD11	2:CF:277:ASN:HD22	1.69	0.58
2:OF:69:LEU:HD13	2:UF:45:ARG:O	2.03	0.58
2:UF:107:ILE:HG23	2:UF:137:LEU:HD23	1.84	0.58
3:VF:54:LEU:HA	3:VF:57:ILE:HG22	1.84	0.58
3:VF:207:GLU:N	3:VF:207:GLU:OE1	2.37	0.58
3:HG:72:LEU:HD23	3:HG:73:LEU:HD12	1.85	0.58
2:G:26:PHE:HB3	2:SG:83:LYS:HB2	1.86	0.58
2:AA:273:LYS:HD2	2:AA:276:ARG:HH21	1.68	0.58
3:BA:106:HIS:CE1	3:BA:113:THR:HG22	2.39	0.58
2:MA:115:PRO:O	2:MA:119:ALA:CB	2.51	0.58
3:NA:198:ILE:HG23	3:NA:221:PHE:CD1	2.38	0.58
2:SA:26:PHE:CE1	2:SA:34:VAL:HG22	2.38	0.58
2:EB:251:LEU:HD11	2:EB:277:ASN:HD22	1.69	0.58
2:QB:251:LEU:HD11	2:QB:277:ASN:HD22	1.69	0.58
2:WB:251:LEU:HD11	2:WB:277:ASN:HD22	1.68	0.58
3:XB:173:ASN:HD22	3:XB:174:PRO:HD3	1.68	0.58
2:IC:244:ASP:OD1	2:IC:276:ARG:NH2	2.36	0.58
2:UC:153:LEU:O	2:UC:157:VAL:HG23	2.03	0.58
2:AD:89:ARG:HH11	2:AD:89:ARG:HG3	1.69	0.58
2:AD:251:LEU:HD11	2:AD:277:ASN:HD22	1.69	0.58
3:ND:174:PRO:HD2	3:ND:174:PRO:O	2.03	0.58
2:YD:153:LEU:O	2:YD:157:VAL:HG23	2.04	0.58
3:ZD:183:GLU:OE1	3:ZD:184:MET:N	2.36	0.58
2:KE:245:ARG:HH12	2:KE:249:ARG:HB2	1.67	0.58
2:IF:95:GLU:O	2:IF:99:GLU:HB2	2.03	0.58
2:IF:273:LYS:HD2	2:IF:276:ARG:HH21	1.68	0.58
2:OF:78:ARG:O	2:OF:82:VAL:HG13	2.03	0.58
3:K:174:PRO:HD2	3:K:174:PRO:O	2.03	0.58
2:AA:21:ARG:HE	2:AA:21:ARG:C	2.05	0.58
3:BA:183:GLU:OE1	3:BA:185:GLN:N	2.36	0.58

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:QB:152:ARG:HH22	2:QB:153:LEU:HB2	1.69	0.58
2:CC:245:ARG:HH12	2:CC:249:ARG:HB2	1.68	0.58
2:IC:251:LEU:HD11	2:IC:277:ASN:HD22	1.69	0.58
2:OC:15:MET:HE2	2:OC:15:MET:N	2.18	0.58
3:VC:240:ARG:NH1	3:VC:240:ARG:HA	2.19	0.58
2:AD:230:ILE:HA	2:AD:233:MET:CE	2.34	0.58
3:BD:191:ILE:HG13	3:BD:192:THR:HG23	1.86	0.58
2:GD:83:LYS:HD2	2:MD:23:ALA:HA	1.85	0.58
3:HD:174:PRO:HD2	3:HD:174:PRO:O	2.03	0.58
3:TD:111:ARG:CZ	3:TD:111:ARG:HA	2.33	0.58
2:YD:297:ARG:HG3	2:YD:299:SER:H	1.69	0.58
3:FE:127:VAL:O	3:FE:131:PHE:HB2	2.04	0.58
2:KE:97:ILE:O	2:KE:101:ARG:NH2	2.27	0.58
3:LE:120:PRO:HA	3:LE:123:VAL:HG22	1.84	0.58
2:QE:109:THR:O	2:QE:113:MET:HG3	2.04	0.58
2:CF:83:LYS:HD3	2:IF:26:PHE:HD1	1.68	0.58
2:CF:273:LYS:HD2	2:CF:276:ARG:HH21	1.68	0.58
2:OF:74:ASN:OD1	2:OF:78:ARG:NH2	2.36	0.58
2:UF:168:GLN:HE22	3:VF:130:LEU:HD11	1.68	0.58
2:AG:151:GLU:O	2:AG:155:HIS:ND1	2.36	0.58
2:GG:115:PRO:O	2:GG:119:ALA:CB	2.51	0.58
2:GG:244:ASP:OD1	2:GG:276:ARG:NH2	2.36	0.58
2:MG:26:PHE:CE1	2:MG:34:VAL:HG22	2.38	0.58
2:MG:129:ILE:HG23	2:MG:175:LEU:HD21	1.84	0.58
2:G:186:GLN:HB3	3:C:74:ARG:NH1	2.18	0.58
2:B:160:ARG:O	2:B:164:PHE:CB	2.52	0.58
3:BA:183:GLU:OE1	3:BA:184:MET:N	2.36	0.58
3:HA:53:ALA:HB3	3:HA:231:LEU:HA	1.85	0.58
3:HA:57:ILE:HD11	3:HA:169:TRP:HA	1.85	0.58
3:HA:72:LEU:HD23	3:HA:73:LEU:HD12	1.84	0.58
2:MA:9:LYS:HG2	2:MA:55:VAL:HG22	1.86	0.58
2:YA:113:MET:HG3	2:YA:117:SER:HB2	1.84	0.58
3:ZA:182:SER:OG	3:ZA:183:GLU:N	2.36	0.58
3:FB:101:ASN:ND2	3:FB:154:ILE:HG21	2.19	0.58
2:QE:251:LEU:HD11	2:QE:277:ASN:HD22	1.69	0.58
2:WE:251:LEU:HD11	2:WE:277:ASN:HD22	1.68	0.58
3:XE:58:ASN:ND2	3:XE:59:GLU:OE1	2.36	0.58
2:CF:129:ILE:HG23	2:CF:175:LEU:HD21	1.85	0.58
2:OF:251:LEU:HD11	2:OF:277:ASN:HD22	1.69	0.58
2:AG:297:ARG:HG3	2:AG:299:SER:H	1.69	0.58
2:MG:80:VAL:O	2:MG:83:LYS:HG2	2.04	0.58

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:RB:75:ARG:HG3	3:RB:208:ILE:HD11	1.86	0.58
2:WB:93:LEU:O	2:WB:97:ILE:HG12	2.04	0.58
2:WB:206:MET:HB2	2:CC:143:ALA:HB1	1.86	0.58
3:DC:225:GLU:HA	3:DC:228:ARG:HB3	1.85	0.58
2:UC:230:ILE:HA	2:UC:233:MET:HE2	1.85	0.58
2:AD:297:ARG:HG3	2:AD:299:SER:H	1.69	0.58
2:EE:11:VAL:HG11	2:EE:41:MET:HB3	1.85	0.58
2:QE:7:THR:HG23	2:QE:37:LEU:HD23	1.86	0.58
2:QE:37:LEU:O	2:QE:41:MET:HG3	2.03	0.58
2:WE:83:LYS:HB2	2:CF:26:PHE:HB3	1.86	0.58
2:WE:120:ASP:OD1	2:WE:123:ARG:NH1	2.37	0.58
2:WE:198:THR:O	2:WE:202:ILE:HG12	2.04	0.58
2:WE:230:ILE:HA	2:WE:233:MET:HE2	1.85	0.58
3:DF:57:ILE:HD11	3:DF:169:TRP:HA	1.85	0.58
2:OF:153:LEU:O	2:OF:157:VAL:HG23	2.04	0.58
2:AG:69:LEU:HD13	2:GG:45:ARG:O	2.03	0.58
2:MG:75:GLU:O	2:MG:79:SER:HB3	2.02	0.58
2:MG:146:LEU:HD11	2:MG:157:VAL:HG11	1.86	0.58
2:MG:273:LYS:HD2	2:MG:276:ARG:HH21	1.69	0.58
2:B:206:MET:SD	2:B:207:LYS:N	2.77	0.58
2:B:297:ARG:HG3	2:B:299:SER:H	1.69	0.58
2:J:40:ALA:O	2:J:44:VAL:HG23	2.04	0.58
2:AA:251:LEU:HD11	2:AA:277:ASN:HD22	1.68	0.58
2:GA:93:LEU:O	2:GA:97:ILE:HG12	2.04	0.58
2:GA:230:ILE:HA	2:GA:233:MET:HE2	1.86	0.58
2:SA:80:VAL:HA	2:YA:26:PHE:CE2	2.39	0.58
2:YA:151:GLU:HA	2:YA:154:ARG:HG2	1.85	0.58
2:YA:273:LYS:HD2	2:YA:276:ARG:HH21	1.68	0.58
2:EB:207:LYS:HZ1	2:EB:209:GLN:HB3	1.66	0.58
3:FB:120:PRO:HA	3:FB:123:VAL:HG22	1.86	0.58
2:WB:273:LYS:HD2	2:WB:276:ARG:HH21	1.68	0.58
3:XB:174:PRO:HD2	3:XB:174:PRO:O	2.04	0.58
2:AD:107:ILE:HG23	2:AD:137:LEU:HD23	1.86	0.58
3:BD:112:GLY:HA3	3:BD:223:MET:HG2	1.85	0.58
2:MD:97:ILE:O	2:MD:101:ARG:NH2	2.32	0.58
2:KE:251:LEU:HD11	2:KE:277:ASN:HD22	1.68	0.58
3:RE:169:TRP:HD1	3:RE:175:LEU:HD13	1.68	0.58
3:RE:182:SER:OG	3:RE:183:GLU:N	2.37	0.58
2:WE:186:GLN:CB	3:DF:74:ARG:HH12	2.16	0.58
3:XE:90:HIS:CE1	3:XE:91:GLU:HG2	2.39	0.58
3:DF:174:PRO:HD2	3:DF:174:PRO:O	2.03	0.58

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:IF:93:LEU:O	2:IF:97:ILE:HG12	2.03	0.58
2:UF:273:LYS:HD2	2:UF:276:ARG:HH21	1.68	0.58
3:VF:101:ASN:ND2	3:VF:154:ILE:HG21	2.18	0.58
2:MG:251:LEU:HD11	2:MG:277:ASN:HD22	1.69	0.58
2:SG:26:PHE:CE1	2:SG:34:VAL:HG22	2.39	0.58
2:SG:74:ASN:O	2:SG:78:ARG:NE	2.36	0.58
2:SG:273:LYS:HD2	2:SG:276:ARG:HH21	1.69	0.58
2:G:45:ARG:O	2:SG:69:LEU:HD13	2.04	0.57
2:SA:251:LEU:HD11	2:SA:277:ASN:HD22	1.68	0.57
2:YA:187:ASN:O	2:YA:189:LYS:NZ	2.37	0.57
2:YA:251:LEU:HD11	2:YA:277:ASN:HD22	1.69	0.57
2:CC:251:LEU:HD11	2:CC:277:ASN:HD22	1.68	0.57
3:DC:190:ASN:ND2	3:JC:234:PRO:O	2.37	0.57
3:PC:229:GLU:OE2	3:PC:233:ASN:ND2	2.37	0.57
2:SD:251:LEU:HD11	2:SD:277:ASN:HD22	1.69	0.57
2:YD:230:ILE:HA	2:YD:233:MET:HE2	1.85	0.57
2:YD:273:LYS:HD2	2:YD:276:ARG:HH21	1.68	0.57
2:QE:168:GLN:HE21	2:QE:170:ALA:HB3	1.69	0.57
2:CF:75:GLU:O	2:CF:79:SER:HB3	2.04	0.57
2:IF:245:ARG:HH12	2:IF:249:ARG:HB2	1.68	0.57
3:JF:174:PRO:HD2	3:JF:174:PRO:O	2.03	0.57
2:UF:230:ILE:HA	2:UF:233:MET:HE2	1.86	0.57
2:G:251:LEU:HD11	2:G:277:ASN:HD22	1.69	0.57
3:M:169:TRP:HD1	3:M:175:LEU:HD13	1.69	0.57
2:J:297:ARG:HG3	2:J:299:SER:H	1.69	0.57
2:AA:26:PHE:CE1	2:AA:34:VAL:HG22	2.40	0.57
2:AA:93:LEU:O	2:AA:97:ILE:HG12	2.04	0.57
2:GA:297:ARG:HG3	2:GA:299:SER:H	1.68	0.57
2:SA:115:PRO:O	2:SA:119:ALA:CB	2.52	0.57
2:YA:93:LEU:O	2:YA:97:ILE:HG12	2.04	0.57
2:KB:108:GLU:HA	2:KB:111:ASN:HD22	1.68	0.57
2:QB:114:GLU:HB2	2:QB:117:SER:OG	2.03	0.57
2:WB:188:LEU:HD13	3:DC:74:ARG:HE	1.67	0.57
2:CC:76:TYR:HE1	2:CC:83:LYS:NZ	2.01	0.57
2:CC:201:GLU:HG2	2:IC:139:ARG:NH2	2.18	0.57
2:IC:205:LEU:HD21	2:OC:140:SER:HA	1.86	0.57
2:OC:251:LEU:HD11	2:OC:277:ASN:HD22	1.69	0.57
3:PC:174:PRO:O	3:PC:174:PRO:HD2	2.03	0.57
2:UC:80:VAL:O	2:UC:83:LYS:HG2	2.04	0.57
2:UC:251:LEU:HD11	2:UC:277:ASN:HD22	1.69	0.57
3:BD:229:GLU:OE1	3:BD:230:LEU:HD12	2.04	0.57

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:YD:78:ARG:O	2:YD:82:VAL:HG23	2.05	0.57
3:ZD:134:ASP:OD2	3:ZD:135:GLY:N	2.37	0.57
3:ZD:182:SER:OG	3:ZD:183:GLU:N	2.38	0.57
2:KE:151:GLU:HA	2:KE:154:ARG:HG2	1.86	0.57
3:LE:106:HIS:CD2	3:LE:113:THR:HG22	2.39	0.57
2:MG:76:TYR:HB2	2:SG:41:MET:HE3	1.85	0.57
2:SG:251:LEU:HD11	2:SG:277:ASN:HD22	1.69	0.57
2:G:219:ARG:HH12	2:G:227:GLN:HG3	1.69	0.57
2:YA:85:LEU:HD22	2:YA:85:LEU:H	1.69	0.57
3:ZA:119:SER:HB3	3:ZA:214:GLU:O	2.04	0.57
2:EB:273:LYS:HD2	2:EB:276:ARG:HH21	1.69	0.57
3:LB:90:HIS:O	3:LB:94:ARG:HG2	2.03	0.57
3:JC:174:PRO:HD2	3:JC:174:PRO:O	2.04	0.57
2:OC:107:ILE:HG23	2:OC:137:LEU:HD23	1.87	0.57
2:UC:244:ASP:OD1	2:UC:276:ARG:NH2	2.38	0.57
2:GD:206:MET:SD	2:GD:207:LYS:N	2.77	0.57
2:MD:121:LEU:HD22	3:TD:70:PHE:CE1	2.39	0.57
2:MD:130:ILE:O	2:MD:134:LEU:HG	2.04	0.57
3:ND:72:LEU:HD23	3:ND:73:LEU:HD12	1.85	0.57
3:ND:221:PHE:HD2	3:ND:225:GLU:OE1	1.87	0.57
2:SD:297:ARG:HG3	2:SD:299:SER:H	1.69	0.57
3:FE:174:PRO:O	3:FE:174:PRO:HD2	2.03	0.57
2:QE:76:TYR:HE1	2:WE:26:PHE:CZ	2.23	0.57
3:RE:240:ARG:NH1	3:RE:240:ARG:HA	2.19	0.57
3:BG:54:LEU:HB2	3:BG:231:LEU:HD23	1.86	0.57
3:HG:183:GLU:OE1	3:HG:184:MET:N	2.38	0.57
2:SG:204:ASN:OD1	2:SG:233:MET:HB3	2.04	0.57
2:SG:244:ASP:OD1	2:SG:276:ARG:NH2	2.37	0.57
3:ZA:118:PHE:CE2	3:ZA:158:LEU:HD12	2.40	0.57
2:QB:190:ARG:O	2:WB:166:GLY:HA3	2.04	0.57
3:XB:59:GLU:OE1	3:XB:59:GLU:N	2.37	0.57
2:CC:62:GLU:OE2	2:IC:45:ARG:NH2	2.36	0.57
2:UC:224:GLU:HA	2:UC:227:GLN:OE1	2.04	0.57
2:UC:273:LYS:HD2	2:UC:276:ARG:HH21	1.69	0.57
3:BD:127:VAL:O	3:BD:131:PHE:HB2	2.04	0.57
3:ND:183:GLU:HG3	3:ND:188:PHE:HB2	1.86	0.57
3:TD:182:SER:OG	3:TD:183:GLU:N	2.37	0.57
3:ZD:174:PRO:O	3:ZD:174:PRO:HD2	2.03	0.57
2:KE:203:ILE:HG12	2:KE:206:MET:HE3	1.86	0.57
2:CF:96:ASP:OD2	2:IF:35:GLN:NE2	2.37	0.57
3:BG:105:ILE:HG12	3:BG:162:LEU:HD11	1.86	0.57

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:GG:202:ILE:CG1	2:MG:139:ARG:HH21	2.16	0.57
2:G:207:LYS:HZ2	2:G:210:GLN:HG3	1.70	0.57
3:BA:79:ILE:HG22	3:BA:204:PHE:HB3	1.86	0.57
2:GA:251:LEU:HD11	2:GA:277:ASN:HD22	1.69	0.57
2:MA:201:GLU:HG3	2:SA:139:ARG:NH2	2.19	0.57
3:TA:190:ASN:ND2	3:ZA:234:PRO:O	2.37	0.57
2:KB:125:GLU:HA	3:LB:147:THR:HG21	1.84	0.57
2:KB:160:ARG:O	2:KB:164:PHE:CB	2.51	0.57
3:XB:109:PRO:O	3:XB:111:ARG:NH1	2.37	0.57
2:CC:201:GLU:HG2	2:IC:139:ARG:HH22	1.68	0.57
2:UC:212:GLU:O	2:UC:216:THR:HG23	2.04	0.57
4:YC:106:ILE:O	4:YC:126:ILE:N	2.37	0.57
2:AD:95:GLU:HG3	2:AD:112:PHE:CE2	2.40	0.57
2:GD:78:ARG:O	2:GD:82:VAL:HG23	2.05	0.57
2:GD:273:LYS:HD2	2:GD:276:ARG:HH21	1.69	0.57
3:NG:72:LEU:HD23	3:NG:73:LEU:HD12	1.87	0.57
3:M:240:ARG:NH1	3:M:240:ARG:HA	2.19	0.57
4:EA:106:ILE:O	3:HA:252:ARG:HG2	2.04	0.57
3:HA:127:VAL:O	3:HA:131:PHE:HB2	2.04	0.57
2:EB:160:ARG:O	2:EB:164:PHE:CB	2.53	0.57
3:LB:101:ASN:ND2	3:LB:154:ILE:HG21	2.19	0.57
2:WB:207:LYS:HZ2	2:WB:210:GLN:HB2	1.69	0.57
2:UC:33:GLU:O	2:UC:37:LEU:HG	2.04	0.57
3:BD:225:GLU:HA	3:BD:228:ARG:HB3	1.87	0.57
3:ND:120:PRO:HA	3:ND:123:VAL:HG22	1.85	0.57
2:QE:69:LEU:HD13	2:WE:45:ARG:O	2.05	0.57
2:QE:152:ARG:HH22	2:QE:153:LEU:HB2	1.69	0.57
2:WE:78:ARG:O	2:WE:82:VAL:HG23	2.04	0.57
3:XE:101:ASN:ND2	3:XE:154:ILE:HG21	2.20	0.57
3:XE:119:SER:HB3	3:XE:214:GLU:O	2.04	0.57
3:VF:112:GLY:HA3	3:VF:223:MET:HG2	1.86	0.57
2:MG:168:GLN:HE22	3:NG:130:LEU:HD11	1.68	0.57
3:M:119:SER:HB3	3:M:214:GLU:O	2.05	0.57
3:K:72:LEU:HD23	3:K:73:LEU:HD12	1.87	0.57
3:K:120:PRO:HA	3:K:123:VAL:HG22	1.85	0.57
3:TA:174:PRO:HD2	3:TA:174:PRO:O	2.04	0.57
2:YA:115:PRO:O	2:YA:119:ALA:CB	2.52	0.57
2:EB:219:ARG:NH2	2:EB:227:GLN:OE1	2.30	0.57
3:RB:216:ASN:O	3:RB:217:ILE:HD13	2.04	0.57
2:CC:297:ARG:HG3	2:CC:299:SER:H	1.69	0.57
2:OC:230:ILE:HA	2:OC:233:MET:HE2	1.86	0.57

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:EE:21:ARG:HE	2:EE:21:ARG:C	2.04	0.57
2:EE:205:LEU:HD21	2:KE:140:SER:HA	1.85	0.57
2:QE:219:ARG:NH2	2:QE:227:GLN:OE1	2.33	0.57
2:IF:230:ILE:HA	2:IF:233:MET:CE	2.34	0.57
2:UF:251:LEU:HD11	2:UF:277:ASN:HD22	1.69	0.57
2:GG:251:LEU:HD11	2:GG:277:ASN:HD22	1.69	0.57
2:MG:151:GLU:HB3	2:MG:155:HIS:CE1	2.40	0.57
2:B:251:LEU:HD11	2:B:277:ASN:HD22	1.69	0.57
3:BA:134:ASP:OD2	3:BA:135:GLY:N	2.37	0.57
3:NA:116:VAL:HG21	3:NA:158:LEU:HD11	1.87	0.57
3:ZA:57:ILE:HD11	3:ZA:169:TRP:HA	1.86	0.57
2:EB:93:LEU:O	2:EB:97:ILE:HG12	2.04	0.57
2:KB:26:PHE:CZ	2:KB:37:LEU:HD11	2.40	0.57
3:LB:183:GLU:HG3	3:LB:188:PHE:HB2	1.86	0.57
2:WB:152:ARG:HH22	2:WB:153:LEU:HB2	1.68	0.57
3:XB:227:LEU:HD13	3:XB:230:LEU:HD13	1.85	0.57
2:CC:89:ARG:HD2	2:CC:90:ALA:H	1.70	0.57
2:UC:230:ILE:HA	2:UC:233:MET:CE	2.35	0.57
2:GD:75:GLU:O	2:GD:79:SER:HB3	2.05	0.57
2:GD:219:ARG:NH2	2:GD:227:GLN:OE1	2.35	0.57
2:MD:297:ARG:HG3	2:MD:299:SER:H	1.69	0.57
2:KE:107:ILE:HG22	2:KE:111:ASN:HD21	1.69	0.57
2:QE:273:LYS:HD2	2:QE:276:ARG:HH21	1.70	0.57
3:DF:183:GLU:HG3	3:DF:188:PHE:HB2	1.85	0.57
2:IF:251:LEU:HD11	2:IF:277:ASN:HD22	1.68	0.57
2:OF:152:ARG:HH22	2:OF:153:LEU:HB2	1.70	0.57
2:GG:273:LYS:HD2	2:GG:276:ARG:HH21	1.69	0.57
3:M:174:PRO:HD2	3:M:174:PRO:O	2.03	0.57
3:C:240:ARG:NH1	3:C:240:ARG:HA	2.20	0.57
2:T:152:ARG:HH22	2:T:153:LEU:HB2	1.70	0.57
2:SA:152:ARG:HH22	2:SA:153:LEU:HB2	1.69	0.57
3:FB:86:ILE:HG13	3:FB:200:VAL:HG12	1.85	0.57
2:KB:115:PRO:O	2:KB:119:ALA:HB2	2.05	0.57
3:XB:72:LEU:HD23	3:XB:73:LEU:HD12	1.86	0.57
3:DC:118:PHE:CE2	3:DC:158:LEU:HD12	2.39	0.57
3:VC:115:LEU:HD21	3:VC:218:CYS:HB3	1.87	0.57
2:AD:160:ARG:O	2:AD:164:PHE:HB2	2.03	0.57
2:GD:153:LEU:O	2:GD:157:VAL:HG23	2.05	0.57
2:YD:251:LEU:HD11	2:YD:277:ASN:HD22	1.68	0.57
2:WE:168:GLN:HE22	3:XE:130:LEU:HD11	1.69	0.57
3:DF:191:ILE:HG13	3:DF:192:THR:HG23	1.86	0.57

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:JF:240:ARG:NH1	3:JF:240:ARG:HA	2.20	0.57
3:NG:229:GLU:OE2	3:NG:233:ASN:ND2	2.38	0.57
2:G:186:GLN:HB3	3:C:74:ARG:HH12	1.68	0.57
3:C:57:ILE:HD11	3:C:169:TRP:HA	1.86	0.57
2:SA:95:GLU:HG3	2:SA:112:PHE:CE2	2.39	0.57
2:SA:205:LEU:HD21	2:YA:140:SER:HA	1.87	0.57
3:LB:112:GLY:HA3	3:LB:223:MET:HG2	1.85	0.57
2:IC:230:ILE:HA	2:IC:233:MET:CE	2.35	0.57
2:OC:63:ALA:HA	2:OC:66:PHE:CZ	2.39	0.57
3:VC:102:LEU:HD22	3:VC:117:VAL:HG22	1.86	0.57
2:GD:190:ARG:O	2:MD:166:GLY:HA3	2.05	0.57
2:SD:152:ARG:HH22	2:SD:153:LEU:HB2	1.70	0.57
2:EE:130:ILE:O	2:EE:134:LEU:HG	2.05	0.57
3:RE:101:ASN:ND2	3:RE:154:ILE:HG21	2.19	0.57
2:WE:201:GLU:OE1	2:CF:139:ARG:NH1	2.38	0.57
2:CF:80:VAL:O	2:CF:83:LYS:HG2	2.05	0.57
2:IF:115:PRO:O	2:IF:119:ALA:HB2	2.05	0.57
2:OF:188:LEU:HB2	2:UF:168:GLN:HB3	1.87	0.57
2:OF:205:LEU:HD21	2:UF:140:SER:HA	1.87	0.57
2:AG:21:ARG:O	2:AG:21:ARG:NE	2.25	0.57
2:GG:78:ARG:O	2:GG:82:VAL:HG23	2.04	0.57
2:GG:107:ILE:HG23	2:GG:137:LEU:HD23	1.87	0.57
3:NG:182:SER:OG	3:NG:183:GLU:N	2.38	0.57
2:SG:80:VAL:O	2:SG:83:LYS:HG2	2.05	0.57
2:SG:230:ILE:HA	2:SG:233:MET:HE2	1.87	0.57
3:TG:119:SER:HB3	3:TG:214:GLU:O	2.05	0.57
2:AA:107:ILE:HG23	2:AA:137:LEU:HD23	1.87	0.56
2:SA:204:ASN:OD1	2:SA:233:MET:HB3	2.04	0.56
2:SA:273:LYS:HD2	2:SA:276:ARG:HH21	1.70	0.56
2:IC:115:PRO:O	2:IC:119:ALA:CB	2.53	0.56
2:IC:242:VAL:HG13	2:IC:323:ILE:HG12	1.87	0.56
2:IC:273:LYS:HD2	2:IC:276:ARG:HH21	1.69	0.56
2:OC:76:TYR:CE2	2:UC:37:LEU:HD13	2.40	0.56
2:UC:206:MET:HB2	2:UC:211:GLU:HG2	1.87	0.56
2:QE:48:SER:OG	2:QE:51:GLN:OE1	2.23	0.56
3:RE:121:SER:O	3:RE:125:ILE:HG13	2.05	0.56
2:CF:21:ARG:HH21	2:CF:24:GLU:HB2	1.69	0.56
2:IF:95:GLU:HG3	2:IF:112:PHE:CE2	2.39	0.56
2:GG:125:GLU:HA	3:HG:147:THR:HG21	1.87	0.56
3:HG:174:PRO:HD2	3:HG:174:PRO:O	2.04	0.56
3:HG:183:GLU:OE1	3:HG:185:GLN:N	2.38	0.56

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:TG:101:ASN:ND2	3:TG:154:ILE:HG21	2.19	0.56
3:TG:240:ARG:NH1	3:TG:240:ARG:HA	2.20	0.56
2:G:297:ARG:HG3	2:G:299:SER:H	1.69	0.56
2:B:186:GLN:NE2	3:K:71:ASN:HA	2.17	0.56
2:T:297:ARG:HG3	2:T:299:SER:H	1.69	0.56
3:V:236:LEU:HB3	3:V:238:ASN:OD1	2.05	0.56
2:GA:205:LEU:HD21	2:MA:140:SER:HA	1.87	0.56
2:MA:187:ASN:N	3:TA:74:ARG:HH22	2.03	0.56
3:FB:115:LEU:HD21	3:FB:218:CYS:HB3	1.86	0.56
2:KB:251:LEU:HD11	2:KB:277:ASN:HD22	1.69	0.56
2:QB:206:MET:SD	2:QB:207:LYS:N	2.78	0.56
3:RB:182:SER:OG	3:RB:183:GLU:N	2.38	0.56
2:WB:107:ILE:HG22	2:WB:111:ASN:HD21	1.70	0.56
2:WB:198:THR:O	2:WB:202:ILE:HG12	2.05	0.56
3:XB:102:LEU:HD22	3:XB:117:VAL:HG22	1.87	0.56
2:CC:89:ARG:HD2	2:CC:90:ALA:N	2.19	0.56
2:UC:242:VAL:HG13	2:UC:323:ILE:HG12	1.87	0.56
2:MD:251:LEU:HD11	2:MD:277:ASN:HD22	1.70	0.56
2:EE:93:LEU:O	2:EE:97:ILE:HG12	2.05	0.56
2:QE:242:VAL:HG13	2:QE:323:ILE:HG12	1.87	0.56
2:CF:78:ARG:O	2:CF:82:VAL:HG23	2.05	0.56
2:UF:80:VAL:O	2:UF:83:LYS:HG2	2.05	0.56
2:MG:21:ARG:HE	2:MG:21:ARG:C	2.06	0.56
2:SG:89:ARG:HD2	2:SG:90:ALA:H	1.69	0.56
2:G:115:PRO:O	2:G:119:ALA:CB	2.53	0.56
2:B:113:MET:O	2:B:141:GLN:NE2	2.38	0.56
3:C:106:HIS:CE1	3:C:113:THR:CG2	2.88	0.56
2:AA:89:ARG:HH11	2:AA:89:ARG:HG3	1.71	0.56
2:GA:190:ARG:O	2:MA:166:GLY:HA3	2.04	0.56
2:MA:22:ALA:O	2:MA:26:PHE:HB2	2.04	0.56
2:MA:203:ILE:HG12	2:MA:206:MET:HE3	1.86	0.56
3:TA:198:ILE:HG23	3:TA:221:PHE:CD1	2.40	0.56
2:YA:201:GLU:OE1	2:EB:139:ARG:NH1	2.38	0.56
3:ZA:106:HIS:CD2	3:ZA:113:THR:HG22	2.39	0.56
2:EB:129:ILE:HG23	2:EB:175:LEU:HD21	1.88	0.56
3:RB:115:LEU:HD21	3:RB:218:CYS:HB3	1.87	0.56
3:RB:183:GLU:OE1	3:RB:184:MET:N	2.38	0.56
3:XB:101:ASN:ND2	3:XB:154:ILE:HG21	2.20	0.56
2:CC:230:ILE:HA	2:CC:233:MET:HE2	1.88	0.56
2:IC:55:VAL:O	2:IC:59:PHE:HB3	2.06	0.56
2:OC:244:ASP:OD1	2:OC:276:ARG:NH2	2.39	0.56

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:UC:130:ILE:HD11	2:UC:157:VAL:HG22	1.86	0.56
2:UC:193:MET:O	2:AD:164:PHE:HD1	1.88	0.56
2:AD:74:ASN:O	2:AD:78:ARG:NE	2.38	0.56
2:AD:242:VAL:HG13	2:AD:323:ILE:HG12	1.88	0.56
3:BD:116:VAL:HG21	3:BD:158:LEU:HD11	1.88	0.56
2:GD:251:LEU:HD11	2:GD:277:ASN:HD22	1.70	0.56
3:HD:100:THR:HG23	3:HD:119:SER:HA	1.87	0.56
2:MD:186:GLN:HB3	3:TD:74:ARG:HH12	1.70	0.56
3:ND:90:HIS:O	3:ND:94:ARG:HG2	2.05	0.56
3:TD:236:LEU:HB3	3:TD:238:ASN:OD1	2.05	0.56
2:YD:80:VAL:O	2:YD:83:LYS:HG2	2.04	0.56
2:EE:80:VAL:O	2:EE:83:LYS:HG2	2.06	0.56
2:EE:245:ARG:HH12	2:EE:249:ARG:HB2	1.68	0.56
3:LE:183:GLU:OE1	3:LE:184:MET:N	2.38	0.56
2:WE:115:PRO:O	2:WE:119:ALA:HB2	2.05	0.56
2:WE:242:VAL:HG13	2:WE:323:ILE:HG12	1.87	0.56
3:JF:191:ILE:HG13	3:JF:192:THR:HG23	1.87	0.56
2:OF:206:MET:SD	2:OF:207:LYS:N	2.78	0.56
3:PF:106:HIS:CE1	3:PF:113:THR:HG22	2.40	0.56
3:PF:182:SER:OG	3:PF:183:GLU:N	2.38	0.56
3:VF:183:GLU:OE1	3:VF:184:MET:N	2.39	0.56
2:AG:186:GLN:NE2	3:HG:71:ASN:HA	2.15	0.56
3:HG:112:GLY:HA3	3:HG:223:MET:HG2	1.87	0.56
2:MG:133:ILE:O	2:MG:137:LEU:HG	2.05	0.56
3:NG:119:SER:HB3	3:NG:214:GLU:O	2.05	0.56
2:G:80:VAL:O	2:G:83:LYS:HG2	2.06	0.56
2:B:71:ILE:O	2:B:74:ASN:ND2	2.39	0.56
2:T:69:LEU:HD13	2:AA:45:ARG:O	2.06	0.56
3:V:100:THR:HG23	3:V:119:SER:HA	1.87	0.56
2:GA:30:SER:OG	2:GA:33:GLU:OE1	2.20	0.56
2:GA:186:GLN:NE2	3:NA:71:ASN:HA	2.20	0.56
2:MA:95:GLU:HG3	2:MA:112:PHE:CE2	2.41	0.56
2:KB:26:PHE:CZ	2:KB:34:VAL:HG22	2.41	0.56
3:RB:183:GLU:OE1	3:RB:185:GLN:N	2.38	0.56
3:DC:105:ILE:HG12	3:DC:162:LEU:HD11	1.88	0.56
3:PC:183:GLU:OE1	3:PC:185:GLN:N	2.38	0.56
2:AD:47:ILE:HD11	2:AD:52:LEU:HB2	1.87	0.56
2:AD:128:GLN:HG2	3:BD:131:PHE:CB	2.32	0.56
3:HD:57:ILE:HD11	3:HD:169:TRP:HA	1.87	0.56
3:HD:72:LEU:HD23	3:HD:73:LEU:HD12	1.87	0.56
2:SD:83:LYS:CE	2:YD:26:PHE:HB3	2.35	0.56

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:YD:115:PRO:O	2:YD:119:ALA:CB	2.53	0.56
2:YD:128:GLN:HE22	2:YD:164:PHE:HD2	1.51	0.56
2:KE:207:LYS:NZ	2:KE:209:GLN:HB3	2.21	0.56
3:XE:57:ILE:HD11	3:XE:169:TRP:HA	1.86	0.56
2:CF:20:ASP:OD1	2:CF:21:ARG:N	2.38	0.56
2:OF:109:THR:O	2:OF:113:MET:HG3	2.05	0.56
3:BG:174:PRO:O	3:BG:174:PRO:HD2	2.05	0.56
2:GG:17:ILE:HD12	2:GG:21:ARG:HG3	1.86	0.56
2:J:251:LEU:HD11	2:J:277:ASN:HD22	1.70	0.56
2:AA:168:GLN:NE2	3:BA:130:LEU:HD11	2.21	0.56
2:MA:153:LEU:O	2:MA:157:VAL:HG23	2.06	0.56
3:NA:69:LEU:O	3:NA:73:LEU:HB2	2.06	0.56
3:NA:182:SER:OG	3:NA:183:GLU:N	2.38	0.56
2:YA:187:ASN:N	3:FB:74:ARG:HH22	2.03	0.56
2:YA:242:VAL:HG13	2:YA:323:ILE:HG12	1.87	0.56
3:XB:120:PRO:HA	3:XB:123:VAL:HG22	1.88	0.56
2:UC:128:GLN:HE22	2:UC:164:PHE:HD2	1.54	0.56
2:AD:108:GLU:HA	2:AD:111:ASN:HD22	1.71	0.56
3:BD:182:SER:OG	3:BD:183:GLU:N	2.39	0.56
2:GD:152:ARG:HH22	2:GD:153:LEU:HB2	1.71	0.56
2:MD:151:GLU:HA	2:MD:154:ARG:HG2	1.87	0.56
2:YD:168:GLN:NE2	3:ZD:130:LEU:HD11	2.21	0.56
3:FE:118:PHE:CE2	3:FE:158:LEU:HD12	2.40	0.56
2:QE:22:ALA:O	2:QE:26:PHE:HB2	2.05	0.56
2:WE:114:GLU:HB2	2:WE:117:SER:OG	2.06	0.56
2:UF:110:LEU:C	2:UF:141:GLN:HE22	2.08	0.56
3:NG:238:ASN:HB3	3:NG:241:HIS:ND1	2.21	0.56
3:M:182:SER:OG	3:M:183:GLU:N	2.39	0.56
2:T:168:GLN:NE2	3:V:130:LEU:HD11	2.20	0.56
3:BA:116:VAL:HG21	3:BA:158:LEU:HD11	1.86	0.56
2:MA:12:ILE:HD11	2:MA:55:VAL:HG11	1.88	0.56
3:TA:101:ASN:ND2	3:TA:154:ILE:HG21	2.20	0.56
2:YA:76:TYR:HE1	2:EB:38:SER:HB2	1.71	0.56
2:EB:21:ARG:HE	2:EB:21:ARG:C	2.07	0.56
2:QB:95:GLU:HG3	2:QB:112:PHE:CE2	2.41	0.56
3:RB:109:PRO:O	3:RB:111:ARG:NH1	2.38	0.56
2:OC:95:GLU:HG3	2:OC:112:PHE:CE2	2.40	0.56
2:OC:133:ILE:O	2:OC:137:LEU:HG	2.05	0.56
3:PC:182:SER:OG	3:PC:183:GLU:N	2.37	0.56
3:HD:119:SER:HB3	3:HD:214:GLU:O	2.06	0.56
3:TD:183:GLU:OE1	3:TD:184:MET:N	2.39	0.56

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:KE:153:LEU:O	2:KE:157:VAL:HG23	2.06	0.56
2:QE:114:GLU:HB2	2:QE:117:SER:OG	2.06	0.56
3:XE:118:PHE:CE2	3:XE:158:LEU:HD12	2.41	0.56
2:MG:168:GLN:NE2	3:NG:130:LEU:HD11	2.21	0.56
2:SG:108:GLU:HA	2:SG:111:ASN:HD22	1.70	0.56
2:MA:120:ASP:OD1	3:TA:76:SER:OG	2.23	0.56
3:FB:121:SER:O	3:FB:125:ILE:HG13	2.05	0.56
2:KB:155:HIS:O	2:KB:159:LEU:HD22	2.05	0.56
3:RB:119:SER:HB3	3:RB:214:GLU:O	2.05	0.56
2:OC:114:GLU:HB2	2:OC:117:SER:OG	2.06	0.56
2:AD:188:LEU:HD13	3:HD:74:ARG:CD	2.36	0.56
2:GD:128:GLN:HG2	3:HD:131:PHE:CB	2.32	0.56
2:QE:26:PHE:CE1	2:QE:34:VAL:HG22	2.41	0.56
2:QE:80:VAL:O	2:QE:83:LYS:HG2	2.05	0.56
3:RE:75:ARG:HG3	3:RE:208:ILE:HD11	1.86	0.56
2:IF:89:ARG:HH22	2:IF:93:LEU:HD21	1.71	0.56
2:OF:193:MET:O	2:UF:164:PHE:HD1	1.89	0.56
2:UF:32:ARG:NH2	2:UF:33:GLU:OE1	2.38	0.56
2:AG:203:ILE:HG12	2:AG:206:MET:HE3	1.88	0.56
2:GG:21:ARG:HH21	2:GG:24:GLU:HB2	1.71	0.56
2:GG:55:VAL:O	2:GG:59:PHE:HB3	2.06	0.56
2:SG:9:LYS:HG2	2:SG:55:VAL:HG22	1.87	0.56
2:G:65:GLN:NE2	2:B:47:ILE:O	2.36	0.56
2:G:95:GLU:HG3	2:G:112:PHE:CE2	2.40	0.56
2:B:230:ILE:HA	2:B:233:MET:CE	2.36	0.56
2:T:83:LYS:O	2:AA:27:LYS:HE3	2.06	0.56
3:V:238:ASN:HB3	3:V:241:HIS:ND1	2.21	0.56
2:GA:9:LYS:HG2	2:GA:55:VAL:HG22	1.88	0.56
2:GA:151:GLU:HB3	2:GA:155:HIS:CE1	2.41	0.56
2:EB:190:ARG:O	2:KB:166:GLY:HA3	2.04	0.56
2:KB:168:GLN:HE22	3:LB:130:LEU:HD11	1.71	0.56
2:CC:153:LEU:O	2:CC:157:VAL:HG23	2.06	0.56
3:DC:112:GLY:HA3	3:DC:223:MET:HG2	1.88	0.56
3:DC:182:SER:OG	3:DC:183:GLU:N	2.39	0.56
2:OC:55:VAL:O	2:OC:59:PHE:HB3	2.05	0.56
2:MD:203:ILE:HA	2:MD:206:MET:HE2	1.87	0.56
3:ND:238:ASN:HB3	3:ND:241:HIS:CD2	2.39	0.56
3:ZD:72:LEU:HD23	3:ZD:73:LEU:HD12	1.86	0.56
3:FE:115:LEU:HD11	3:FE:218:CYS:HB3	1.86	0.56
2:QE:230:ILE:HA	2:QE:233:MET:HE2	1.87	0.56
3:XE:65:PHE:CE1	3:XE:161:ALA:HB2	2.41	0.56

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:XE:111:ARG:NE	3:XE:111:ARG:HA	2.20	0.56
3:BG:239:SER:HA	3:BG:242:GLU:OE2	2.06	0.56
2:GG:204:ASN:OD1	2:GG:233:MET:HB3	2.06	0.56
3:M:86:ILE:HD11	3:M:221:PHE:HE1	1.71	0.56
3:K:57:ILE:HD11	3:K:169:TRP:HA	1.86	0.56
2:GA:230:ILE:HA	2:GA:233:MET:CE	2.36	0.56
2:MA:160:ARG:O	2:MA:164:PHE:CB	2.54	0.56
3:NA:191:ILE:HG13	3:NA:192:THR:HG23	1.88	0.56
2:YA:26:PHE:CE1	2:YA:34:VAL:HG22	2.37	0.56
3:ZA:236:LEU:HB3	3:ZA:238:ASN:OD1	2.06	0.56
2:CC:151:GLU:O	2:CC:155:HIS:ND1	2.39	0.56
2:IC:69:LEU:HD13	2:OC:45:ARG:O	2.06	0.56
3:VC:106:HIS:CD2	3:VC:113:THR:HG22	2.40	0.56
2:AD:230:ILE:HA	2:AD:233:MET:HE2	1.88	0.56
2:KE:128:GLN:HE22	2:KE:164:PHE:HD2	1.52	0.56
2:QE:33:GLU:O	2:QE:37:LEU:HG	2.06	0.56
2:IF:76:TYR:HD2	2:OF:41:MET:HE2	1.71	0.56
2:IF:83:LYS:HD3	2:OF:26:PHE:CD2	2.41	0.56
2:IF:114:GLU:HB2	2:IF:117:SER:OG	2.05	0.56
3:JF:100:THR:HG23	3:JF:119:SER:HA	1.87	0.56
2:UF:186:GLN:HB3	3:BG:74:ARG:NH1	2.21	0.56
3:BG:118:PHE:CE2	3:BG:158:LEU:HD12	2.41	0.56
2:G:128:GLN:CG	3:M:131:PHE:HB3	2.32	0.56
3:LB:119:SER:HB3	3:LB:214:GLU:O	2.05	0.56
3:LB:120:PRO:HA	3:LB:123:VAL:HG22	1.87	0.56
2:WB:219:ARG:NH2	2:WB:227:GLN:OE1	2.31	0.56
2:WB:242:VAL:HG13	2:WB:323:ILE:HG12	1.88	0.56
2:OC:9:LYS:HG2	2:OC:55:VAL:HG22	1.88	0.56
2:AD:168:GLN:NE2	3:BD:130:LEU:HD11	2.21	0.56
2:MD:190:ARG:O	2:SD:166:GLY:HA3	2.05	0.56
2:SD:37:LEU:O	2:SD:41:MET:HG3	2.05	0.56
2:WE:80:VAL:HG12	2:CF:26:PHE:CE2	2.36	0.56
3:XE:102:LEU:HD22	3:XE:117:VAL:HG22	1.88	0.56
2:CF:152:ARG:HH22	2:CF:153:LEU:HB2	1.71	0.56
2:AG:21:ARG:NH2	2:AG:24:GLU:HB2	2.20	0.56
2:AG:126:HIS:CE1	3:BG:127:VAL:HG13	2.41	0.56
2:G:114:GLU:HB2	2:G:117:SER:OG	2.06	0.55
2:T:128:GLN:HG2	3:V:131:PHE:CB	2.34	0.55
2:SA:11:VAL:HG11	2:SA:41:MET:HB3	1.86	0.55
2:SA:187:ASN:O	2:SA:189:LYS:NZ	2.39	0.55
2:KB:55:VAL:O	2:KB:59:PHE:HB3	2.07	0.55

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:LB:97:PRO:O	3:LB:100:THR:OG1	2.18	0.55
2:QB:33:GLU:O	2:QB:37:LEU:HG	2.06	0.55
2:QB:55:VAL:O	2:QB:59:PHE:HB3	2.06	0.55
2:CC:83:LYS:HB3	2:IC:27:LYS:HB2	1.88	0.55
2:AD:21:ARG:HE	2:AD:21:ARG:C	2.06	0.55
2:AD:115:PRO:O	2:AD:119:ALA:CB	2.54	0.55
3:BD:119:SER:HB3	3:BD:214:GLU:O	2.05	0.55
2:GD:9:LYS:HG2	2:GD:55:VAL:HG22	1.89	0.55
2:EE:230:ILE:HA	2:EE:233:MET:HE2	1.86	0.55
3:FE:57:ILE:HD11	3:FE:169:TRP:HA	1.87	0.55
2:KE:115:PRO:O	2:KE:119:ALA:CB	2.54	0.55
2:KE:242:VAL:HG13	2:KE:323:ILE:HG12	1.87	0.55
2:QE:188:LEU:HB2	2:WE:168:GLN:HB3	1.89	0.55
3:DF:112:GLY:HA3	3:DF:223:MET:HG2	1.86	0.55
2:UF:190:ARG:O	2:AG:166:GLY:HA3	2.05	0.55
2:SG:55:VAL:O	2:SG:59:PHE:HB3	2.07	0.55
2:G:188:LEU:HD13	3:C:74:ARG:CD	2.35	0.55
2:G:190:ARG:O	2:B:166:GLY:HA3	2.05	0.55
2:T:21:ARG:HH21	2:T:24:GLU:HB2	1.71	0.55
3:HA:224:ILE:O	3:HA:228:ARG:N	2.39	0.55
3:TA:120:PRO:HA	3:TA:123:VAL:HG22	1.87	0.55
2:EB:95:GLU:HG3	2:EB:112:PHE:CE2	2.41	0.55
2:EB:152:ARG:HH22	2:EB:153:LEU:HB2	1.71	0.55
2:KB:110:LEU:O	2:KB:113:MET:N	2.36	0.55
2:KB:168:GLN:NE2	3:LB:130:LEU:HD11	2.21	0.55
2:QB:128:GLN:HG2	3:RB:131:PHE:CB	2.34	0.55
2:CC:110:LEU:C	2:CC:141:GLN:HE22	2.10	0.55
3:JC:236:LEU:HB3	3:JC:238:ASN:ND2	2.20	0.55
2:AD:55:VAL:O	2:AD:59:PHE:HB3	2.06	0.55
2:MD:201:GLU:HG3	2:SD:139:ARG:HE	1.71	0.55
3:ND:69:LEU:O	3:ND:73:LEU:HB2	2.06	0.55
3:TD:119:SER:HB3	3:TD:214:GLU:O	2.06	0.55
2:QE:55:VAL:O	2:QE:59:PHE:HB3	2.07	0.55
2:WE:76:TYR:HE2	2:CF:26:PHE:CE2	2.24	0.55
2:WE:153:LEU:O	2:WE:157:VAL:HG23	2.05	0.55
3:JF:120:PRO:HA	3:JF:123:VAL:HG22	1.87	0.55
2:UF:188:LEU:H	3:BG:74:ARG:NH2	2.04	0.55
3:VF:120:PRO:HA	3:VF:123:VAL:HG22	1.87	0.55
2:MG:89:ARG:HG3	2:MG:89:ARG:HH11	1.72	0.55
2:SG:130:ILE:HD11	2:SG:157:VAL:HG22	1.87	0.55
2:G:139:ARG:NH2	2:SG:201:GLU:HG3	2.21	0.55

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:K:101:ASN:ND2	3:K:154:ILE:HG21	2.21	0.55
2:GA:205:LEU:HD13	2:MA:139:ARG:HB3	1.88	0.55
2:YA:83:LYS:HG2	2:EB:26:PHE:HB3	1.88	0.55
3:RB:72:LEU:HD23	3:RB:73:LEU:HD12	1.87	0.55
3:RB:106:HIS:CD2	3:RB:113:THR:HG22	2.40	0.55
2:WB:138:LYS:HE2	2:WB:138:LYS:HA	1.88	0.55
3:DC:174:PRO:HD2	3:DC:174:PRO:O	2.05	0.55
2:AD:69:LEU:HD21	2:GD:47:ILE:HB	1.88	0.55
3:HD:240:ARG:NH1	3:HD:240:ARG:HA	2.22	0.55
3:LE:69:LEU:O	3:LE:73:LEU:HB2	2.06	0.55
2:CF:41:MET:HG2	2:CF:42:ALA:N	2.21	0.55
2:CF:242:VAL:HG13	2:CF:323:ILE:HG12	1.87	0.55
3:DF:120:PRO:HA	3:DF:123:VAL:HG22	1.87	0.55
2:IF:97:ILE:O	2:IF:101:ARG:NH2	2.34	0.55
3:JF:119:SER:HB3	3:JF:214:GLU:O	2.06	0.55
3:JF:207:GLU:N	3:JF:207:GLU:OE2	2.39	0.55
2:AG:168:GLN:NE2	3:BG:130:LEU:HD11	2.21	0.55
3:NG:115:LEU:HD21	3:NG:218:CYS:HB3	1.88	0.55
2:SG:128:GLN:CG	3:TG:131:PHE:HB3	2.34	0.55
2:B:26:PHE:CZ	2:B:34:VAL:HG22	2.42	0.55
2:B:245:ARG:O	2:B:245:ARG:NH1	2.37	0.55
3:K:53:ALA:HB3	3:K:231:LEU:HA	1.86	0.55
2:MA:242:VAL:HG13	2:MA:323:ILE:HG12	1.88	0.55
3:NA:97:PRO:O	3:NA:100:THR:OG1	2.17	0.55
2:EB:242:VAL:HG13	2:EB:323:ILE:HG12	1.88	0.55
2:QB:153:LEU:O	2:QB:157:VAL:HG23	2.06	0.55
3:PC:102:LEU:HD22	3:PC:117:VAL:HG22	1.88	0.55
2:SD:114:GLU:HB2	2:SD:117:SER:OG	2.07	0.55
2:YD:55:VAL:O	2:YD:59:PHE:HB3	2.07	0.55
2:YD:107:ILE:HG23	2:YD:137:LEU:HD23	1.88	0.55
3:ZD:79:ILE:HG22	3:ZD:204:PHE:HB3	1.87	0.55
2:EE:151:GLU:HB3	2:EE:155:HIS:CE1	2.41	0.55
3:LE:182:SER:OG	3:LE:183:GLU:N	2.39	0.55
3:RE:120:PRO:HA	3:RE:123:VAL:HG22	1.88	0.55
3:DF:221:PHE:HD2	3:DF:225:GLU:OE1	1.89	0.55
3:JF:101:ASN:ND2	3:JF:154:ILE:HG21	2.20	0.55
3:PF:183:GLU:OE1	3:PF:184:MET:N	2.38	0.55
2:MG:107:ILE:HG23	2:MG:137:LEU:HD23	1.87	0.55
2:SG:95:GLU:HG3	2:SG:112:PHE:CE2	2.41	0.55
3:TG:86:ILE:HG12	3:TG:198:ILE:HD11	1.89	0.55
3:TG:182:SER:OG	3:TG:183:GLU:N	2.40	0.55

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:G:55:VAL:O	2:G:59:PHE:HB3	2.06	0.55
2:G:93:LEU:O	2:G:97:ILE:HG12	2.07	0.55
2:AA:186:GLN:HE22	3:HA:71:ASN:CA	2.10	0.55
3:BA:75:ARG:HG3	3:BA:208:ILE:HD11	1.89	0.55
3:BA:101:ASN:ND2	3:BA:154:ILE:HG21	2.22	0.55
3:HA:102:LEU:HD22	3:HA:117:VAL:HG22	1.89	0.55
3:NA:54:LEU:HB2	3:NA:231:LEU:HD23	1.88	0.55
2:YA:97:ILE:O	2:YA:101:ARG:NH2	2.24	0.55
2:IC:158:MET:HE3	2:IC:159:LEU:HG	1.87	0.55
3:PC:119:SER:HB3	3:PC:214:GLU:O	2.06	0.55
3:PC:236:LEU:HB3	3:PC:238:ASN:HD21	1.71	0.55
3:VC:182:SER:OG	3:VC:183:GLU:N	2.40	0.55
2:GD:107:ILE:HG23	2:GD:137:LEU:HD23	1.87	0.55
2:SD:78:ARG:O	2:SD:82:VAL:HG13	2.05	0.55
2:EE:9:LYS:HG2	2:EE:55:VAL:HG22	1.87	0.55
2:KE:37:LEU:O	2:KE:41:MET:HG3	2.06	0.55
3:RE:97:PRO:O	3:RE:100:THR:OG1	2.18	0.55
2:WE:31:THR:O	2:WE:34:VAL:HB	2.06	0.55
2:WE:126:HIS:CE1	3:XE:127:VAL:HG13	2.42	0.55
2:OF:188:LEU:HD21	3:VF:74:ARG:HD2	1.87	0.55
2:UF:128:GLN:CG	3:VF:131:PHE:HB3	2.33	0.55
2:MG:128:GLN:CG	3:NG:131:PHE:HB3	2.33	0.55
3:NG:183:GLU:HG3	3:NG:188:PHE:HB2	1.87	0.55
3:TG:106:HIS:CD2	3:TG:113:THR:HG22	2.41	0.55
2:G:151:GLU:HA	2:G:154:ARG:HG2	1.89	0.55
2:J:31:THR:O	2:J:34:VAL:HB	2.07	0.55
2:T:97:ILE:O	2:T:101:ARG:NH2	2.34	0.55
3:V:118:PHE:CE2	3:V:158:LEU:HD12	2.42	0.55
3:BA:182:SER:OG	3:BA:183:GLU:N	2.37	0.55
3:HA:183:GLU:OE1	3:HA:184:MET:N	2.40	0.55
2:WB:55:VAL:O	2:WB:59:PHE:HB3	2.06	0.55
2:WB:128:GLN:CG	3:XB:131:PHE:HB3	2.35	0.55
2:CC:107:ILE:HG23	2:CC:137:LEU:HD23	1.87	0.55
2:OC:242:VAL:HG13	2:OC:323:ILE:HG12	1.88	0.55
2:OC:297:ARG:HG3	2:OC:299:SER:H	1.72	0.55
2:GD:242:VAL:HG13	2:GD:323:ILE:HG12	1.88	0.55
3:HD:120:PRO:HA	3:HD:123:VAL:HG22	1.88	0.55
3:RE:115:LEU:HD21	3:RE:218:CYS:HB3	1.89	0.55
2:WE:107:ILE:HG23	2:WE:137:LEU:HD23	1.89	0.55
2:CF:188:LEU:HD23	2:CF:188:LEU:H	1.72	0.55
2:CF:245:ARG:O	2:CF:245:ARG:NH1	2.37	0.55

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:IF:187:ASN:O	2:IF:189:LYS:NZ	2.38	0.55
3:VF:102:LEU:HD22	3:VF:117:VAL:HG22	1.88	0.55
2:AG:55:VAL:O	2:AG:59:PHE:HB3	2.07	0.55
3:HG:191:ILE:HG13	3:HG:192:THR:HG23	1.89	0.55
3:C:182:SER:OG	3:C:183:GLU:N	2.40	0.55
2:J:55:VAL:O	2:J:59:PHE:HB3	2.07	0.55
2:GA:80:VAL:O	2:GA:83:LYS:HG2	2.06	0.55
2:MA:151:GLU:HA	2:MA:154:ARG:HG2	1.88	0.55
2:SA:55:VAL:O	2:SA:59:PHE:HB3	2.07	0.55
2:SA:130:ILE:HD11	2:SA:157:VAL:HG22	1.88	0.55
2:CC:96:ASP:OD2	2:IC:35:GLN:NE2	2.39	0.55
3:JC:182:SER:OG	3:JC:183:GLU:N	2.37	0.55
3:VC:119:SER:HB3	3:VC:214:GLU:O	2.05	0.55
2:GD:130:ILE:HD11	2:GD:157:VAL:HG22	1.88	0.55
2:MD:151:GLU:HB3	2:MD:155:HIS:CE1	2.41	0.55
3:ND:100:THR:HG23	3:ND:119:SER:HA	1.89	0.55
2:SD:212:GLU:O	2:SD:216:THR:HG23	2.06	0.55
2:YD:69:LEU:HD13	2:EE:45:ARG:O	2.07	0.55
2:EE:120:ASP:OD2	3:LE:76:SER:OG	2.24	0.55
3:RE:59:GLU:N	3:RE:59:GLU:OE1	2.40	0.55
3:RE:118:PHE:CE2	3:RE:158:LEU:HD12	2.42	0.55
3:RE:221:PHE:HD2	3:RE:225:GLU:OE2	1.89	0.55
2:WE:55:VAL:O	2:WE:59:PHE:HB3	2.07	0.55
2:WE:151:GLU:HA	2:WE:154:ARG:HG2	1.89	0.55
3:PF:100:THR:HG23	3:PF:119:SER:HA	1.89	0.55
2:AG:107:ILE:HG23	2:AG:137:LEU:HD23	1.88	0.55
2:AG:111:ASN:OD1	2:AG:138:LYS:N	2.31	0.55
3:BG:182:SER:OG	3:BG:183:GLU:N	2.40	0.55
2:GG:37:LEU:O	2:GG:41:MET:HG2	2.06	0.55
2:GG:242:VAL:HG13	2:GG:323:ILE:HG12	1.88	0.55
2:MG:55:VAL:O	2:MG:59:PHE:HB3	2.07	0.55
3:NG:236:LEU:HB3	3:NG:238:ASN:OD1	2.07	0.55
2:SG:11:VAL:HG11	2:SG:41:MET:HB3	1.89	0.55
2:SG:89:ARG:HD2	2:SG:90:ALA:N	2.22	0.55
2:G:153:LEU:O	2:G:157:VAL:HG23	2.07	0.55
2:G:245:ARG:O	2:G:245:ARG:NH1	2.37	0.55
3:M:74:ARG:HD2	2:SG:188:LEU:HD21	1.89	0.55
2:B:97:ILE:O	2:B:101:ARG:NH2	2.34	0.55
2:T:242:VAL:HG13	2:T:323:ILE:HG12	1.88	0.55
2:AA:55:VAL:O	2:AA:59:PHE:HB3	2.06	0.55
2:GA:152:ARG:HH22	2:GA:153:LEU:HB2	1.72	0.55

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:MA:102:ASP:HB2	2:MA:108:GLU:HG2	1.88	0.55
2:EB:55:VAL:O	2:EB:59:PHE:HB3	2.07	0.55
2:EB:113:MET:O	2:EB:141:GLN:NE2	2.39	0.55
3:RB:236:LEU:HB3	3:RB:238:ASN:OD1	2.05	0.55
2:WB:134:LEU:HD13	2:WB:146:LEU:HD21	1.88	0.55
2:IC:105:SER:OG	2:IC:107:ILE:HD12	2.06	0.55
2:OC:168:GLN:NE2	3:PC:130:LEU:HD11	2.22	0.55
2:UC:74:ASN:O	2:UC:78:ARG:NE	2.40	0.55
2:UC:110:LEU:O	2:UC:141:GLN:NE2	2.37	0.55
2:SD:14:LEU:HD12	2:SD:22:ALA:HB1	1.89	0.55
2:SD:97:ILE:O	2:SD:101:ARG:NH2	2.30	0.55
2:QE:95:GLU:HG3	2:QE:112:PHE:CE2	2.42	0.55
2:QE:167:VAL:HG13	3:RE:130:LEU:O	2.06	0.55
2:IF:74:ASN:O	2:IF:78:ARG:NE	2.40	0.55
2:IF:90:ALA:HA	2:IF:93:LEU:HG	1.88	0.55
3:PF:115:LEU:HD21	3:PF:218:CYS:HB3	1.89	0.55
2:G:242:VAL:HG13	2:G:323:ILE:HG12	1.88	0.55
2:B:153:LEU:O	2:B:157:VAL:HG23	2.07	0.55
2:J:151:GLU:HA	2:J:154:ARG:HG2	1.89	0.55
2:SA:80:VAL:O	2:SA:83:LYS:HG2	2.06	0.55
2:QB:186:GLN:NE2	3:XB:71:ASN:HA	2.15	0.55
2:OC:151:GLU:HB3	2:OC:155:HIS:CE1	2.41	0.55
2:UC:89:ARG:HH11	2:UC:89:ARG:HG3	1.72	0.55
3:VC:58:ASN:OD1	3:VC:81:VAL:HG11	2.07	0.55
2:MD:152:ARG:HH22	2:MD:153:LEU:HB2	1.72	0.55
2:KE:12:ILE:HD11	2:KE:55:VAL:HG11	1.88	0.55
2:UF:55:VAL:O	2:UF:59:PHE:HB3	2.07	0.55
2:UF:186:GLN:NE2	3:BG:71:ASN:HA	2.21	0.55
2:AG:11:VAL:O	2:AG:15:MET:HE3	2.07	0.55
2:AG:212:GLU:O	2:AG:216:THR:HG23	2.06	0.55
3:BG:225:GLU:HA	3:BG:228:ARG:HB3	1.88	0.55
2:B:204:ASN:OD1	2:B:233:MET:HB3	2.07	0.55
2:AA:242:VAL:HG13	2:AA:323:ILE:HG12	1.87	0.55
2:EB:76:TYR:OH	2:KB:34:VAL:HG13	2.07	0.55
3:XB:100:THR:HG23	3:XB:119:SER:HA	1.88	0.55
2:CC:76:TYR:HE1	2:CC:83:LYS:HZ1	1.55	0.55
3:JC:106:HIS:CD2	3:JC:113:THR:HG22	2.42	0.55
2:OC:160:ARG:O	2:OC:164:PHE:HB3	2.06	0.55
2:UC:63:ALA:HA	2:UC:66:PHE:CZ	2.41	0.55
2:AD:114:GLU:HB2	2:AD:117:SER:OG	2.06	0.55
3:HD:182:SER:OG	3:HD:183:GLU:N	2.40	0.55

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:EE:110:LEU:C	2:EE:141:GLN:HE22	2.09	0.55
2:EE:168:GLN:HE22	3:FE:130:LEU:HD11	1.72	0.55
2:QE:63:ALA:HA	2:QE:66:PHE:CZ	2.42	0.55
2:CF:188:LEU:HB2	2:IF:168:GLN:HB3	1.89	0.55
2:IF:34:VAL:O	2:IF:37:LEU:HD12	2.07	0.55
2:IF:55:VAL:O	2:IF:59:PHE:HB3	2.06	0.55
2:IF:69:LEU:HD13	2:OF:45:ARG:O	2.07	0.55
2:UF:12:ILE:HD11	2:UF:55:VAL:HG11	1.89	0.55
2:SG:7:THR:HG23	2:SG:37:LEU:HD23	1.89	0.55
2:SG:93:LEU:O	2:SG:97:ILE:HG12	2.07	0.55
2:SG:158:MET:SD	2:SG:159:LEU:HD13	2.46	0.55
2:J:190:ARG:O	2:T:166:GLY:HA3	2.06	0.54
2:J:246:SER:HB3	2:J:321:MET:HE3	1.89	0.54
2:GA:26:PHE:CE1	2:GA:34:VAL:HG22	2.42	0.54
2:GA:206:MET:HB3	2:GA:210:GLN:NE2	2.22	0.54
4:QA:106:ILE:O	3:TA:252:ARG:HG2	2.07	0.54
2:SA:63:ALA:HA	2:SA:66:PHE:CZ	2.42	0.54
2:YA:151:GLU:HA	2:YA:154:ARG:HE	1.72	0.54
2:EB:41:MET:HG2	2:EB:42:ALA:N	2.21	0.54
2:EB:205:LEU:CD2	2:KB:140:SER:HA	2.37	0.54
2:CC:74:ASN:OD1	2:CC:78:ARG:NH2	2.40	0.54
2:CC:80:VAL:CA	2:CC:83:LYS:HE3	2.34	0.54
2:CC:187:ASN:N	3:JC:74:ARG:HH22	2.05	0.54
3:DC:96:LEU:HD23	3:DC:97:PRO:CD	2.37	0.54
2:IC:134:LEU:HD13	2:IC:146:LEU:HD21	1.89	0.54
2:IC:187:ASN:N	3:PC:74:ARG:HH22	2.05	0.54
2:UC:128:GLN:CG	3:VC:131:PHE:HB3	2.34	0.54
2:AD:128:GLN:CG	3:BD:131:PHE:HB3	2.32	0.54
3:ZD:116:VAL:HG21	3:ZD:158:LEU:HD11	1.89	0.54
2:KE:34:VAL:O	2:KE:37:LEU:HD12	2.06	0.54
3:LE:54:LEU:HB2	3:LE:231:LEU:HD23	1.89	0.54
3:XE:72:LEU:HD23	3:XE:73:LEU:HD12	1.88	0.54
3:XE:106:HIS:CD2	3:XE:113:THR:HG22	2.42	0.54
2:OF:168:GLN:NE2	3:PF:130:LEU:HD11	2.23	0.54
2:UF:242:VAL:HG13	2:UF:323:ILE:HG12	1.88	0.54
3:HG:182:SER:OG	3:HG:183:GLU:N	2.40	0.54
2:SG:63:ALA:HA	2:SG:66:PHE:CZ	2.42	0.54
2:T:55:VAL:O	2:T:59:PHE:HB3	2.07	0.54
2:T:151:GLU:HB3	2:T:155:HIS:CE1	2.42	0.54
2:AA:151:GLU:HB3	2:AA:155:HIS:CE1	2.43	0.54
3:BA:240:ARG:HA	3:BA:240:ARG:NH1	2.21	0.54

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:MA:203:ILE:HA	2:MA:206:MET:CE	2.37	0.54
2:SA:93:LEU:O	2:SA:97:ILE:HG12	2.07	0.54
3:TA:97:PRO:O	3:TA:100:THR:OG1	2.19	0.54
2:QB:89:ARG:HG3	2:QB:90:ALA:N	2.23	0.54
3:RB:100:THR:HG23	3:RB:119:SER:HA	1.89	0.54
3:XB:183:GLU:OE1	3:XB:184:MET:N	2.40	0.54
2:CC:20:ASP:O	2:CC:23:ALA:HB3	2.06	0.54
2:CC:69:LEU:HD13	2:IC:44:VAL:HG12	1.89	0.54
2:CC:79:SER:HA	2:CC:82:VAL:CG1	2.37	0.54
2:IC:89:ARG:HG3	2:IC:89:ARG:HH11	1.72	0.54
2:UC:55:VAL:O	2:UC:59:PHE:HB3	2.07	0.54
2:GD:168:GLN:NE2	3:HD:130:LEU:HD11	2.21	0.54
3:HD:236:LEU:HB3	3:HD:238:ASN:HD21	1.70	0.54
2:MD:230:ILE:HA	2:MD:233:MET:CE	2.37	0.54
2:YD:95:GLU:HG3	2:YD:112:PHE:CE2	2.42	0.54
2:CF:55:VAL:O	2:CF:59:PHE:HB3	2.07	0.54
2:IF:26:PHE:CZ	2:IF:34:VAL:HG22	2.43	0.54
2:IF:30:SER:OG	2:IF:33:GLU:OE1	2.17	0.54
2:IF:83:LYS:HD3	2:OF:26:PHE:CE2	2.41	0.54
3:JF:238:ASN:HB3	3:JF:241:HIS:HD2	1.72	0.54
2:UF:89:ARG:HH22	2:AG:31:THR:HG23	1.72	0.54
2:AG:93:LEU:O	2:AG:97:ILE:HG12	2.07	0.54
3:HG:97:PRO:O	3:HG:100:THR:OG1	2.16	0.54
2:SG:152:ARG:HH22	2:SG:153:LEU:HB2	1.71	0.54
3:TG:100:THR:HG23	3:TG:119:SER:HA	1.90	0.54
2:G:72:ASN:HD22	2:B:41:MET:HG3	1.73	0.54
2:B:110:LEU:O	2:B:141:GLN:NE2	2.29	0.54
2:J:66:PHE:HB3	2:T:45:ARG:HD2	1.88	0.54
2:J:95:GLU:HG3	2:J:112:PHE:CE2	2.42	0.54
3:TA:180:VAL:HG23	3:TA:181:ARG:H	1.72	0.54
3:RB:225:GLU:HA	3:RB:228:ARG:HB3	1.89	0.54
3:PC:120:PRO:HA	3:PC:123:VAL:HG22	1.90	0.54
2:UC:207:LYS:HG3	2:UC:210:GLN:HG2	1.90	0.54
2:UC:219:ARG:HH12	2:UC:227:GLN:HG3	1.73	0.54
3:VC:120:PRO:HA	3:VC:123:VAL:HG22	1.88	0.54
2:AD:204:ASN:OD1	2:AD:233:MET:HB3	2.06	0.54
2:SD:153:LEU:O	2:SD:157:VAL:HG23	2.08	0.54
2:SD:242:VAL:HG13	2:SD:323:ILE:HG12	1.89	0.54
2:YD:114:GLU:HB2	2:YD:117:SER:OG	2.06	0.54
2:EE:55:VAL:O	2:EE:59:PHE:HB3	2.08	0.54
2:KE:205:LEU:HD22	2:QE:139:ARG:HB3	1.89	0.54

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:WE:245:ARG:O	2:WE:245:ARG:NH1	2.37	0.54
2:CF:107:ILE:HG23	2:CF:137:LEU:HD23	1.90	0.54
2:CF:186:GLN:NE2	3:JF:71:ASN:HA	2.22	0.54
2:OF:115:PRO:O	2:OF:119:ALA:CB	2.55	0.54
2:UF:134:LEU:HD13	2:UF:146:LEU:HD21	1.90	0.54
2:AG:63:ALA:HA	2:AG:66:PHE:CZ	2.42	0.54
2:MG:190:ARG:O	2:SG:166:GLY:HA3	2.07	0.54
2:SG:141:GLN:HA	2:SG:144:ASP:OD2	2.07	0.54
2:SG:219:ARG:NH2	2:SG:227:GLN:OE1	2.31	0.54
3:M:72:LEU:HD23	3:M:73:LEU:HD12	1.90	0.54
3:M:116:VAL:HG21	3:M:158:LEU:HD11	1.88	0.54
3:V:183:GLU:HG3	3:V:188:PHE:HB2	1.89	0.54
2:AA:114:GLU:HB2	2:AA:117:SER:OG	2.07	0.54
3:BA:224:ILE:O	3:BA:228:ARG:N	2.40	0.54
2:GA:187:ASN:N	3:NA:74:ARG:HH22	2.05	0.54
3:HA:97:PRO:O	3:HA:100:THR:OG1	2.17	0.54
3:HA:191:ILE:HG13	3:HA:192:THR:HG23	1.88	0.54
2:MA:202:ILE:HD13	2:SA:139:ARG:NH2	2.21	0.54
2:SA:151:GLU:HB3	2:SA:155:HIS:CE1	2.42	0.54
3:TA:236:LEU:HB3	3:TA:238:ASN:OD1	2.07	0.54
3:TA:238:ASN:HB3	3:TA:241:HIS:ND1	2.22	0.54
2:YA:297:ARG:HG3	2:YA:299:SER:H	1.72	0.54
4:CB:106:ILE:O	3:FB:252:ARG:HG2	2.07	0.54
2:EB:297:ARG:HG3	2:EB:299:SER:H	1.72	0.54
2:KB:245:ARG:O	2:KB:245:ARG:NH1	2.37	0.54
2:IC:95:GLU:HG3	2:IC:112:PHE:CE2	2.43	0.54
3:PC:77:PRO:HB3	3:PC:208:ILE:HB	1.89	0.54
2:UC:109:THR:O	2:UC:113:MET:HG3	2.08	0.54
3:VC:97:PRO:O	3:VC:100:THR:OG1	2.19	0.54
2:GD:55:VAL:O	2:GD:59:PHE:HB3	2.08	0.54
2:GD:160:ARG:O	2:GD:164:PHE:CB	2.55	0.54
2:GD:168:GLN:OE1	2:GD:168:GLN:N	2.38	0.54
3:ND:96:LEU:HD23	3:ND:97:PRO:CD	2.37	0.54
2:SD:109:THR:O	2:SD:113:MET:HG3	2.07	0.54
2:SD:168:GLN:NE2	3:TD:130:LEU:HD11	2.21	0.54
4:WD:107:ALA:HB3	3:ZD:252:ARG:HH11	1.71	0.54
2:EE:206:MET:HB3	2:EE:210:GLN:NE2	2.23	0.54
2:QE:128:GLN:HE22	2:QE:164:PHE:HD2	1.54	0.54
3:DF:224:ILE:O	3:DF:228:ARG:N	2.41	0.54
2:IF:168:GLN:HE21	2:IF:171:ALA:H	1.56	0.54
2:OF:31:THR:O	2:OF:34:VAL:HB	2.08	0.54

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:OF:89:ARG:HG3	2:OF:90:ALA:N	2.23	0.54
3:NG:102:LEU:HD22	3:NG:117:VAL:HG22	1.90	0.54
3:C:100:THR:HG23	3:C:119:SER:HA	1.89	0.54
3:C:134:ASP:OD1	3:C:136:ARG:N	2.26	0.54
2:J:201:GLU:HG3	2:T:139:ARG:NH2	2.23	0.54
2:T:30:SER:OG	2:T:33:GLU:OE1	2.21	0.54
3:V:101:ASN:ND2	3:V:154:ILE:HG21	2.22	0.54
2:AA:95:GLU:HG3	2:AA:112:PHE:CE2	2.43	0.54
2:AA:187:ASN:O	2:AA:189:LYS:NZ	2.37	0.54
2:GA:55:VAL:O	2:GA:59:PHE:HB3	2.08	0.54
2:MA:114:GLU:HB2	2:MA:117:SER:OG	2.07	0.54
2:MA:127:PRO:HA	2:MA:130:ILE:HG12	1.89	0.54
2:SA:187:ASN:N	3:ZA:74:ARG:HH22	2.06	0.54
2:YA:142:ALA:O	2:YA:146:LEU:HG	2.08	0.54
3:ZA:65:PHE:CE1	3:ZA:161:ALA:HB2	2.42	0.54
3:ZA:191:ILE:HG13	3:ZA:192:THR:HG23	1.89	0.54
2:EB:110:LEU:HD13	2:EB:137:LEU:HD21	1.90	0.54
3:FB:100:THR:HG23	3:FB:119:SER:HA	1.90	0.54
3:LB:116:VAL:HG21	3:LB:158:LEU:HD11	1.89	0.54
2:WB:83:LYS:HE2	2:CC:26:PHE:HB3	1.90	0.54
2:CC:126:HIS:CE1	3:DC:127:VAL:HG13	2.42	0.54
3:PC:97:PRO:O	3:PC:97:PRO:HD2	2.07	0.54
2:AD:69:LEU:HD13	2:GD:45:ARG:O	2.07	0.54
2:MD:83:LYS:O	2:SD:27:LYS:HD2	2.07	0.54
2:SD:72:ASN:ND2	2:YD:41:MET:HB2	2.23	0.54
3:TD:120:PRO:HA	3:TD:123:VAL:HG22	1.89	0.54
2:YD:93:LEU:O	2:YD:97:ILE:HG12	2.07	0.54
2:EE:230:ILE:HA	2:EE:233:MET:CE	2.38	0.54
2:KE:9:LYS:HG2	2:KE:55:VAL:HG22	1.89	0.54
2:KE:186:GLN:NE2	3:RE:71:ASN:HA	2.22	0.54
2:WE:297:ARG:HG3	2:WE:299:SER:H	1.73	0.54
3:DF:106:HIS:CD2	3:DF:113:THR:HG22	2.43	0.54
2:OF:128:GLN:HG2	3:PF:131:PHE:CB	2.33	0.54
3:VF:182:SER:OG	3:VF:183:GLU:N	2.41	0.54
2:SG:151:GLU:HB3	2:SG:155:HIS:CE1	2.43	0.54
2:SG:168:GLN:NE2	3:TG:130:LEU:HD11	2.23	0.54
3:M:198:ILE:HG23	3:M:221:PHE:CD1	2.43	0.54
2:J:230:ILE:HA	2:J:233:MET:CE	2.37	0.54
2:MA:11:VAL:HG11	2:MA:41:MET:HB3	1.89	0.54
2:MA:93:LEU:O	2:MA:97:ILE:HG12	2.08	0.54
2:QB:96:ASP:OD2	2:WB:35:GLN:NE2	2.40	0.54

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:WB:168:GLN:NE2	3:XB:130:LEU:HD11	2.22	0.54
2:CC:93:LEU:O	2:CC:97:ILE:HG12	2.07	0.54
2:CC:168:GLN:NE2	3:DC:130:LEU:HD11	2.21	0.54
2:OC:22:ALA:O	2:OC:26:PHE:HB2	2.07	0.54
2:OC:167:VAL:HG21	3:PC:131:PHE:CD1	2.43	0.54
2:GD:26:PHE:HE1	2:GD:34:VAL:HG22	1.72	0.54
2:MD:55:VAL:O	2:MD:59:PHE:HB3	2.07	0.54
2:MD:69:LEU:O	2:MD:72:ASN:ND2	2.34	0.54
2:MD:201:GLU:OE1	2:SD:139:ARG:NH2	2.41	0.54
4:QD:106:ILE:H	3:TD:252:ARG:CZ	2.21	0.54
2:SD:89:ARG:HG3	2:SD:90:ALA:N	2.22	0.54
2:SD:151:GLU:HB3	2:SD:155:HIS:CE1	2.42	0.54
2:KE:130:ILE:HD11	2:KE:157:VAL:HG22	1.89	0.54
2:QE:168:GLN:NE2	2:QE:171:ALA:H	2.06	0.54
2:WE:20:ASP:OD1	2:WE:20:ASP:N	2.41	0.54
2:WE:109:THR:O	2:WE:113:MET:HG3	2.07	0.54
3:XE:191:ILE:HG13	3:XE:192:THR:HG23	1.89	0.54
3:PF:106:HIS:CE1	3:PF:113:THR:CG2	2.91	0.54
2:UF:63:ALA:HA	2:UF:66:PHE:CZ	2.43	0.54
2:UF:93:LEU:O	2:UF:97:ILE:HG12	2.08	0.54
2:UF:153:LEU:O	2:UF:157:VAL:HG23	2.08	0.54
2:UF:207:LYS:NZ	2:UF:209:GLN:OE1	2.34	0.54
2:AG:9:LYS:HG2	2:AG:55:VAL:HG22	1.88	0.54
2:MG:74:ASN:O	2:MG:78:ARG:NE	2.40	0.54
2:MG:242:VAL:HG13	2:MG:323:ILE:HG12	1.88	0.54
2:AA:122:ILE:HB	2:AA:130:ILE:HG22	1.90	0.54
3:NA:79:ILE:HG22	3:NA:204:PHE:HB3	1.89	0.54
2:SA:12:ILE:HD11	2:SA:55:VAL:HG11	1.90	0.54
2:SA:153:LEU:O	2:SA:157:VAL:HG23	2.08	0.54
2:EB:107:ILE:HG23	2:EB:137:LEU:HD23	1.89	0.54
2:EB:230:ILE:HA	2:EB:233:MET:CE	2.38	0.54
2:KB:178:VAL:HG11	3:LB:149:THR:HG22	1.90	0.54
2:QB:83:LYS:HD2	2:WB:26:PHE:HD1	1.73	0.54
2:WB:190:ARG:O	2:CC:166:GLY:HA3	2.07	0.54
2:WB:297:ARG:HG3	2:WB:299:SER:H	1.73	0.54
2:CC:190:ARG:O	2:IC:166:GLY:HA3	2.07	0.54
2:IC:188:LEU:HB3	3:PC:74:ARG:NE	2.23	0.54
2:OC:26:PHE:CE1	2:OC:34:VAL:HG22	2.42	0.54
2:UC:152:ARG:HH22	2:UC:153:LEU:HB2	1.72	0.54
3:BD:198:ILE:HG23	3:BD:221:PHE:CD1	2.42	0.54
2:GD:187:ASN:N	3:ND:74:ARG:HH22	2.06	0.54

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:ND:116:VAL:HG21	3:ND:158:LEU:HD11	1.90	0.54
2:KE:55:VAL:O	2:KE:59:PHE:HB3	2.08	0.54
2:KE:75:GLU:O	2:KE:79:SER:HB3	2.08	0.54
2:KE:245:ARG:O	2:KE:245:ARG:NH1	2.37	0.54
2:WE:205:LEU:HD21	2:CF:140:SER:HA	1.90	0.54
2:CF:205:LEU:HD22	2:IF:139:ARG:HB3	1.90	0.54
3:DF:100:THR:HG23	3:DF:119:SER:HA	1.88	0.54
2:OF:80:VAL:HG11	2:OF:93:LEU:HD12	1.89	0.54
3:PF:119:SER:HB3	3:PF:214:GLU:O	2.06	0.54
2:AG:153:LEU:O	2:AG:157:VAL:HG23	2.06	0.54
2:GG:190:ARG:O	2:MG:166:GLY:HA3	2.08	0.54
2:GG:202:ILE:HG12	2:MG:139:ARG:HH21	1.71	0.54
2:G:31:THR:O	2:G:34:VAL:HB	2.08	0.54
2:G:207:LYS:NZ	2:G:210:GLN:HG3	2.23	0.54
2:G:230:ILE:HA	2:G:233:MET:CE	2.38	0.54
3:K:69:LEU:O	3:K:73:LEU:HB2	2.07	0.54
3:K:111:ARG:NH2	3:K:222:SER:O	2.40	0.54
2:T:11:VAL:O	2:T:15:MET:HE2	2.07	0.54
2:AA:98:LEU:HB3	2:AA:108:GLU:OE1	2.08	0.54
2:IC:63:ALA:HA	2:IC:66:PHE:CZ	2.42	0.54
2:SD:55:VAL:O	2:SD:59:PHE:HB3	2.08	0.54
2:YD:152:ARG:HA	2:YD:155:HIS:HE1	1.73	0.54
2:YD:188:LEU:HD23	2:YD:188:LEU:H	1.73	0.54
2:CF:95:GLU:HG3	2:CF:112:PHE:CE2	2.42	0.54
2:IF:204:ASN:OD1	2:IF:233:MET:HB3	2.07	0.54
2:IF:218:VAL:HG12	2:IF:226:ALA:HB2	1.90	0.54
2:UF:41:MET:SD	2:UF:42:ALA:N	2.81	0.54
2:GG:230:ILE:HA	2:GG:233:MET:CE	2.37	0.54
3:HG:74:ARG:HB2	3:HG:74:ARG:HH11	1.73	0.54
2:B:55:VAL:O	2:B:59:PHE:HB3	2.08	0.54
2:B:98:LEU:HB3	2:B:108:GLU:OE1	2.07	0.54
3:C:119:SER:HB3	3:C:214:GLU:O	2.07	0.54
2:AA:133:ILE:O	2:AA:137:LEU:HG	2.06	0.54
2:GA:15:MET:N	2:GA:15:MET:HE2	2.23	0.54
3:HA:77:PRO:HB3	3:HA:208:ILE:HB	1.90	0.54
2:SA:83:LYS:O	2:YA:27:LYS:HE3	2.08	0.54
3:TA:182:SER:OG	3:TA:183:GLU:N	2.39	0.54
2:YA:168:GLN:NE2	3:ZA:130:LEU:HD11	2.22	0.54
2:EB:230:ILE:HA	2:EB:233:MET:HE2	1.89	0.54
2:KB:207:LYS:NZ	2:KB:209:GLN:HB3	2.22	0.54
3:LB:69:LEU:HD22	3:LB:73:LEU:HD13	1.90	0.54

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:LB:238:ASN:HB3	3:LB:241:HIS:CD2	2.43	0.54
2:QB:168:GLN:NE2	3:RB:130:LEU:HD11	2.23	0.54
2:UC:128:GLN:HG2	3:VC:131:PHE:CB	2.34	0.54
2:SD:69:LEU:HD13	2:YD:45:ARG:O	2.08	0.54
2:SD:190:ARG:O	2:YD:166:GLY:HA3	2.08	0.54
2:EE:204:ASN:OD1	2:EE:233:MET:HB3	2.07	0.54
2:KE:95:GLU:HG3	2:KE:112:PHE:CE2	2.43	0.54
2:QE:11:VAL:O	2:QE:15:MET:HE3	2.08	0.54
2:WE:63:ALA:HA	2:WE:66:PHE:CZ	2.43	0.54
2:WE:204:ASN:OD1	2:WE:233:MET:HB3	2.08	0.54
3:XE:111:ARG:HA	3:XE:111:ARG:CZ	2.37	0.54
4:AF:106:ILE:O	3:DF:252:ARG:HG2	2.08	0.54
2:IF:198:THR:O	2:IF:202:ILE:HG12	2.08	0.54
2:OF:11:VAL:O	2:OF:15:MET:HE3	2.08	0.54
2:UF:83:LYS:O	2:AG:27:LYS:HD2	2.08	0.54
2:UF:97:ILE:O	2:UF:101:ARG:NH2	2.32	0.54
2:SG:21:ARG:HE	2:SG:21:ARG:C	2.08	0.54
3:M:74:ARG:HG3	2:SG:188:LEU:HD11	1.90	0.54
2:B:151:GLU:HB3	2:B:155:HIS:CE1	2.43	0.54
3:C:120:PRO:HA	3:C:123:VAL:HG22	1.89	0.54
3:TA:119:SER:HB3	3:TA:214:GLU:O	2.08	0.54
3:ZA:120:PRO:HA	3:ZA:123:VAL:HG22	1.90	0.54
2:WB:63:ALA:HA	2:WB:66:PHE:CZ	2.43	0.54
2:IC:203:ILE:HA	2:IC:206:MET:CE	2.38	0.54
2:OC:89:ARG:HH11	2:OC:89:ARG:HG3	1.71	0.54
2:UC:107:ILE:HG22	2:UC:111:ASN:HD21	1.73	0.54
2:AD:63:ALA:HA	2:AD:66:PHE:CZ	2.42	0.54
3:HD:190:ASN:HD22	3:ND:236:LEU:HD13	1.72	0.54
2:MD:205:LEU:HD21	2:SD:140:SER:HA	1.90	0.54
2:YD:37:LEU:HD12	2:YD:38:SER:N	2.22	0.54
2:EE:186:GLN:HB3	3:LE:74:ARG:NH2	2.23	0.54
2:QE:78:ARG:O	2:QE:82:VAL:HG12	2.08	0.54
2:CF:31:THR:O	2:CF:34:VAL:HB	2.08	0.54
2:CF:188:LEU:CD2	3:JF:74:ARG:HH21	2.21	0.54
2:IF:205:LEU:HD13	2:OF:139:ARG:HB3	1.89	0.54
2:SG:128:GLN:HG2	3:TG:131:PHE:CB	2.34	0.54
2:SG:128:GLN:HE22	2:SG:164:PHE:HD2	1.56	0.54
2:G:63:ALA:HA	2:G:66:PHE:CZ	2.43	0.53
2:G:168:GLN:HE22	3:M:130:LEU:HD11	1.72	0.53
3:M:86:ILE:HD11	3:M:221:PHE:CE1	2.43	0.53
2:J:187:ASN:N	3:V:74:ARG:HH22	2.06	0.53

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:K:106:HIS:CD2	3:K:113:THR:HG22	2.42	0.53
3:K:239:SER:HA	3:K:242:GLU:OE1	2.08	0.53
2:T:21:ARG:NH2	2:T:24:GLU:HB2	2.23	0.53
2:AA:78:ARG:O	2:AA:82:VAL:HG23	2.08	0.53
3:BA:238:ASN:HB3	3:BA:241:HIS:ND1	2.23	0.53
2:MA:55:VAL:O	2:MA:59:PHE:HB3	2.09	0.53
2:SA:11:VAL:O	2:SA:15:MET:HE3	2.08	0.53
3:TA:54:LEU:HA	3:TA:57:ILE:HG22	1.90	0.53
2:YA:8:ASP:O	2:YA:12:ILE:HG12	2.08	0.53
2:EB:118:ALA:HA	2:EB:121:LEU:HG	1.89	0.53
2:WB:153:LEU:O	2:WB:157:VAL:HG23	2.09	0.53
3:XB:119:SER:HB3	3:XB:214:GLU:O	2.08	0.53
2:CC:114:GLU:HB2	2:CC:117:SER:OG	2.07	0.53
2:CC:204:ASN:OD1	2:CC:233:MET:HB3	2.08	0.53
2:OC:111:ASN:HA	2:OC:141:GLN:OE1	2.08	0.53
2:SD:12:ILE:HD11	2:SD:55:VAL:HG11	1.89	0.53
4:CE:107:ALA:H	3:FE:252:ARG:NH2	2.05	0.53
3:LE:119:SER:HB3	3:LE:214:GLU:O	2.08	0.53
2:QE:245:ARG:O	2:QE:245:ARG:NH1	2.38	0.53
2:WE:142:ALA:O	2:WE:146:LEU:HG	2.08	0.53
2:CF:63:ALA:HA	2:CF:66:PHE:CZ	2.43	0.53
2:CF:69:LEU:HD13	2:IF:45:ARG:O	2.08	0.53
3:PF:72:LEU:HD23	3:PF:73:LEU:HD12	1.90	0.53
2:AG:78:ARG:O	2:AG:82:VAL:HG12	2.07	0.53
2:GG:174:GLU:HG2	3:HG:152:ARG:HG2	1.89	0.53
2:MG:282:ALA:HA	2:MG:285:ILE:HG12	1.90	0.53
2:B:11:VAL:O	2:B:14:LEU:HD12	2.08	0.53
2:B:142:ALA:O	2:B:146:LEU:HG	2.08	0.53
2:J:230:ILE:HA	2:J:233:MET:HE2	1.88	0.53
3:K:182:SER:OG	3:K:183:GLU:N	2.40	0.53
2:T:76:TYR:HD2	2:T:79:SER:OG	1.91	0.53
2:AA:63:ALA:HA	2:AA:66:PHE:CZ	2.43	0.53
2:YA:84:ALA:O	2:YA:87:GLU:HB3	2.09	0.53
2:KB:202:ILE:HD13	2:QB:139:ARG:NH1	2.23	0.53
2:QB:21:ARG:O	2:QB:21:ARG:NE	2.21	0.53
2:QB:21:ARG:NH2	2:QB:24:GLU:HB2	2.24	0.53
3:PC:100:THR:HG23	3:PC:119:SER:HA	1.90	0.53
2:UC:95:GLU:HG3	2:UC:112:PHE:CE2	2.42	0.53
3:BD:240:ARG:NH1	3:BD:240:ARG:HA	2.23	0.53
3:HD:116:VAL:HG21	3:HD:158:LEU:HD11	1.90	0.53
2:MD:230:ILE:HA	2:MD:233:MET:HE2	1.90	0.53

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:YD:302:GLU:O	2:YD:306:LYS:HG2	2.09	0.53
2:KE:108:GLU:HA	2:KE:111:ASN:HD22	1.73	0.53
2:KE:114:GLU:HB2	2:KE:117:SER:OG	2.07	0.53
2:CF:21:ARG:HE	2:CF:21:ARG:C	2.09	0.53
4:SF:106:ILE:O	3:VF:252:ARG:HG2	2.08	0.53
3:VF:100:THR:HG23	3:VF:119:SER:HA	1.89	0.53
3:VF:221:PHE:HD2	3:VF:225:GLU:OE1	1.91	0.53
3:BG:119:SER:HB3	3:BG:214:GLU:O	2.07	0.53
2:GG:63:ALA:HA	2:GG:66:PHE:CZ	2.44	0.53
2:AA:152:ARG:HH22	2:AA:153:LEU:HB2	1.74	0.53
3:NA:96:LEU:HD23	3:NA:97:PRO:CD	2.39	0.53
2:SA:69:LEU:O	2:SA:72:ASN:ND2	2.41	0.53
2:SA:133:ILE:O	2:SA:137:LEU:HG	2.08	0.53
2:SA:190:ARG:O	2:YA:166:GLY:HA3	2.07	0.53
2:KB:11:VAL:O	2:KB:15:MET:HE2	2.08	0.53
2:QB:204:ASN:OD1	2:QB:233:MET:HB3	2.08	0.53
2:QB:230:ILE:HA	2:QB:233:MET:HE2	1.90	0.53
3:DC:119:SER:HB3	3:DC:214:GLU:O	2.07	0.53
3:DC:246:TRP:HB2	3:DC:247:ARG:HH12	1.74	0.53
2:GD:110:LEU:O	2:GD:141:GLN:NE2	2.40	0.53
2:MD:114:GLU:HB2	2:MD:117:SER:OG	2.09	0.53
2:YD:242:VAL:HG13	2:YD:323:ILE:HG12	1.90	0.53
2:QE:204:ASN:OD1	2:QE:233:MET:HB3	2.08	0.53
3:DF:59:GLU:OE2	3:DF:59:GLU:N	2.40	0.53
2:IF:85:LEU:HD22	2:IF:85:LEU:H	1.73	0.53
2:OF:79:SER:O	2:OF:83:LYS:HG2	2.08	0.53
3:PF:202:THR:OG1	3:PF:217:ILE:HG22	2.08	0.53
2:UF:151:GLU:HA	2:UF:154:ARG:HG2	1.91	0.53
2:SG:107:ILE:H	2:SG:107:ILE:HD12	1.74	0.53
2:SG:158:MET:HE3	2:SG:159:LEU:HD22	1.90	0.53
2:SG:245:ARG:O	2:SG:245:ARG:NH1	2.38	0.53
3:TG:120:PRO:HA	3:TG:123:VAL:HG22	1.89	0.53
2:G:69:LEU:HD13	2:B:45:ARG:O	2.08	0.53
2:G:107:ILE:HG23	2:G:137:LEU:HD23	1.89	0.53
2:J:168:GLN:NE2	3:K:130:LEU:HD11	2.23	0.53
2:T:107:ILE:HG22	2:T:111:ASN:ND2	2.21	0.53
2:MA:26:PHE:CE1	2:MA:34:VAL:HG22	2.42	0.53
2:MA:26:PHE:HE1	2:MA:34:VAL:HG22	1.74	0.53
3:TA:115:LEU:HD21	3:TA:218:CYS:HB3	1.90	0.53
2:QB:219:ARG:NH2	2:QB:227:GLN:OE1	2.35	0.53
2:QB:302:GLU:O	2:QB:306:LYS:HG2	2.08	0.53

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:WB:152:ARG:NH2	2:WB:153:LEU:HB2	2.23	0.53
2:IC:190:ARG:O	2:OC:166:GLY:HA3	2.08	0.53
2:UC:151:GLU:HA	2:UC:154:ARG:HG2	1.89	0.53
2:MD:188:LEU:HB3	3:TD:74:ARG:NE	2.23	0.53
2:EE:245:ARG:O	2:EE:245:ARG:NH1	2.37	0.53
3:FE:120:PRO:HA	3:FE:123:VAL:HG22	1.90	0.53
2:QE:151:GLU:HA	2:QE:154:ARG:HG2	1.90	0.53
2:AG:239:LEU:HD11	2:AG:309:LEU:HD11	1.90	0.53
3:NG:120:PRO:HA	3:NG:123:VAL:HG22	1.89	0.53
2:G:128:GLN:HG2	3:M:131:PHE:CB	2.33	0.53
3:M:207:GLU:OE1	3:M:207:GLU:N	2.42	0.53
3:M:252:ARG:HG2	4:WG:106:ILE:O	2.09	0.53
2:B:151:GLU:HA	2:B:154:ARG:HG2	1.90	0.53
2:J:9:LYS:HG2	2:J:55:VAL:HG22	1.90	0.53
3:V:120:PRO:HA	3:V:123:VAL:HG22	1.91	0.53
2:GA:130:ILE:HD11	2:GA:157:VAL:HG22	1.90	0.53
2:MA:107:ILE:HG22	2:MA:111:ASN:ND2	2.23	0.53
2:SA:111:ASN:OD1	2:SA:138:LYS:N	2.35	0.53
2:SA:168:GLN:NE2	3:TA:130:LEU:HD11	2.23	0.53
2:YA:55:VAL:O	2:YA:59:PHE:HB3	2.08	0.53
2:YA:67:ALA:O	2:YA:71:ILE:HG12	2.09	0.53
3:FB:119:SER:HB3	3:FB:214:GLU:O	2.08	0.53
2:KB:188:LEU:HB3	3:RB:74:ARG:NE	2.23	0.53
4:MB:101:ILE:N	4:MB:104:TYR:O	2.33	0.53
2:QB:151:GLU:HB3	2:QB:155:HIS:CE1	2.44	0.53
2:IC:21:ARG:O	2:IC:21:ARG:NE	2.26	0.53
2:OC:146:LEU:HD11	2:OC:157:VAL:HG11	1.90	0.53
3:VC:239:SER:HA	3:VC:242:GLU:OE1	2.09	0.53
2:GD:297:ARG:HG3	2:GD:299:SER:H	1.74	0.53
2:MD:168:GLN:NE2	3:ND:130:LEU:HD11	2.22	0.53
2:YD:30:SER:O	2:YD:34:VAL:HG23	2.09	0.53
2:YD:31:THR:O	2:YD:34:VAL:HB	2.09	0.53
2:EE:152:ARG:HH22	2:EE:153:LEU:HB2	1.73	0.53
2:KE:186:GLN:HE21	3:RE:74:ARG:HH22	1.57	0.53
2:WE:239:LEU:HD11	2:WE:309:LEU:HD11	1.90	0.53
3:XE:190:ASN:OD1	3:DF:236:LEU:HB2	2.09	0.53
2:CF:297:ARG:HG3	2:CF:299:SER:H	1.72	0.53
2:IF:207:LYS:NZ	2:IF:209:GLN:HB3	2.22	0.53
2:GG:9:LYS:HG2	2:GG:55:VAL:HG22	1.90	0.53
3:HG:120:PRO:HA	3:HG:123:VAL:HG22	1.89	0.53
2:G:302:GLU:O	2:G:306:LYS:HG2	2.09	0.53

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:C:116:VAL:HG21	3:C:158:LEU:HD11	1.89	0.53
2:T:168:GLN:HE22	3:V:130:LEU:HD11	1.73	0.53
2:MA:78:ARG:O	2:MA:82:VAL:HG23	2.08	0.53
2:MA:84:ALA:O	2:MA:87:GLU:HB3	2.08	0.53
2:EB:63:ALA:HA	2:EB:66:PHE:CZ	2.43	0.53
4:OB:106:ILE:CB	3:RB:252:ARG:HG2	2.39	0.53
2:WB:128:GLN:HG2	3:XB:131:PHE:CB	2.35	0.53
2:WB:151:GLU:HA	2:WB:154:ARG:HG2	1.89	0.53
2:WB:207:LYS:NZ	2:WB:209:GLN:OE1	2.36	0.53
3:JC:100:THR:HG23	3:JC:119:SER:HA	1.91	0.53
3:PC:115:LEU:HD21	3:PC:218:CYS:HB3	1.91	0.53
3:VC:101:ASN:ND2	3:VC:154:ILE:HG21	2.24	0.53
2:GD:110:LEU:C	2:GD:141:GLN:HE22	2.12	0.53
4:KD:106:ILE:O	3:ND:252:ARG:HG2	2.08	0.53
2:MD:95:GLU:O	2:MD:99:GLU:HB2	2.07	0.53
2:SD:128:GLN:CG	3:TD:131:PHE:HB3	2.35	0.53
3:ZD:249:ASN:O	3:ZD:252:ARG:HB2	2.08	0.53
2:KE:107:ILE:H	2:KE:107:ILE:HD12	1.72	0.53
2:IF:107:ILE:HG23	2:IF:137:LEU:HD23	1.90	0.53
3:JF:58:ASN:OD1	3:JF:81:VAL:HG11	2.08	0.53
2:OF:21:ARG:HE	2:OF:21:ARG:C	2.11	0.53
2:OF:114:GLU:HB2	2:OF:117:SER:OG	2.08	0.53
3:VF:111:ARG:NH2	3:VF:222:SER:O	2.41	0.53
2:MG:206:MET:SD	2:MG:210:GLN:HB2	2.48	0.53
2:J:121:LEU:HD21	3:V:70:PHE:CZ	2.43	0.53
2:J:133:ILE:O	2:J:137:LEU:HG	2.09	0.53
2:GA:302:GLU:O	2:GA:306:LYS:HG2	2.09	0.53
2:EB:152:ARG:NH2	2:EB:153:LEU:HB2	2.23	0.53
2:EB:245:ARG:O	2:EB:245:ARG:NH1	2.38	0.53
3:FB:116:VAL:HG21	3:FB:158:LEU:HD11	1.88	0.53
2:KB:190:ARG:O	2:QB:166:GLY:HA3	2.08	0.53
2:KB:198:THR:O	2:KB:202:ILE:HG12	2.08	0.53
2:KB:302:GLU:O	2:KB:306:LYS:HG2	2.09	0.53
3:DC:202:THR:OG1	3:DC:202:THR:O	2.27	0.53
2:IC:26:PHE:CE1	2:IC:34:VAL:HG22	2.44	0.53
2:GD:21:ARG:O	2:GD:21:ARG:NE	2.24	0.53
3:HD:238:ASN:HB3	3:HD:241:HIS:CD2	2.39	0.53
3:TD:101:ASN:ND2	3:TD:154:ILE:HG21	2.24	0.53
2:EE:186:GLN:HB3	3:LE:74:ARG:NH1	2.24	0.53
2:KE:186:GLN:HE22	3:RE:71:ASN:HA	1.74	0.53
2:IF:80:VAL:HG11	2:IF:93:LEU:HD12	1.90	0.53

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:PF:229:GLU:OE2	3:PF:233:ASN:ND2	2.42	0.53
2:MG:239:LEU:HD11	2:MG:309:LEU:HD11	1.91	0.53
2:SG:111:ASN:HA	2:SG:141:GLN:OE1	2.08	0.53
2:J:239:LEU:HD11	2:J:309:LEU:HD11	1.91	0.53
2:GA:128:GLN:HG2	3:HA:131:PHE:CB	2.36	0.53
2:GA:168:GLN:HE22	3:HA:130:LEU:HD11	1.72	0.53
2:SA:302:GLU:O	2:SA:306:LYS:HG2	2.09	0.53
2:QB:108:GLU:HG3	2:QB:112:PHE:HE2	1.74	0.53
2:WB:188:LEU:H	3:DC:74:ARG:NH2	2.04	0.53
3:JC:120:PRO:HA	3:JC:123:VAL:HG22	1.91	0.53
2:YD:264:LYS:HD2	2:YD:265:GLY:N	2.24	0.53
2:QE:98:LEU:HB3	2:QE:108:GLU:OE1	2.09	0.53
3:XE:120:PRO:HA	3:XE:123:VAL:HG22	1.89	0.53
2:IF:302:GLU:O	2:IF:306:LYS:HG2	2.09	0.53
3:JF:116:VAL:HG21	3:JF:158:LEU:HD11	1.90	0.53
2:OF:55:VAL:O	2:OF:59:PHE:HB3	2.08	0.53
2:OF:204:ASN:OD1	2:OF:233:MET:HB3	2.09	0.53
2:UF:95:GLU:HG3	2:UF:112:PHE:CE2	2.44	0.53
2:UF:152:ARG:NH2	2:UF:153:LEU:HB2	2.23	0.53
2:UF:198:THR:O	2:UF:201:GLU:HG2	2.09	0.53
4:YF:106:ILE:O	3:BG:252:ARG:HG2	2.09	0.53
2:AG:21:ARG:HE	2:AG:21:ARG:C	2.10	0.53
3:BG:224:ILE:O	3:BG:228:ARG:N	2.41	0.53
2:SG:97:ILE:O	2:SG:101:ARG:NH2	2.29	0.53
2:SG:230:ILE:HA	2:SG:233:MET:CE	2.38	0.53
3:K:207:GLU:OE1	3:K:207:GLU:N	2.42	0.53
3:NA:174:PRO:HD2	3:NA:174:PRO:O	2.09	0.53
3:TA:59:GLU:OE1	3:TA:59:GLU:N	2.42	0.53
2:YA:110:LEU:O	2:YA:141:GLN:NE2	2.36	0.53
2:YA:113:MET:HG2	2:YA:118:ALA:HB2	1.89	0.53
3:ZA:225:GLU:HA	3:ZA:228:ARG:HB3	1.89	0.53
3:LB:225:GLU:HA	3:LB:228:ARG:HB3	1.91	0.53
4:OB:106:ILE:H	3:RB:252:ARG:CZ	2.22	0.53
2:QB:63:ALA:HA	2:QB:66:PHE:CZ	2.44	0.53
2:IC:89:ARG:HG3	2:IC:90:ALA:N	2.24	0.53
2:IC:107:ILE:O	2:IC:111:ASN:ND2	2.42	0.53
2:OC:193:MET:SD	2:OC:193:MET:N	2.82	0.53
2:UC:188:LEU:HB3	3:BD:74:ARG:NE	2.24	0.53
3:VC:190:ASN:ND2	3:BD:234:PRO:O	2.41	0.53
3:HD:116:VAL:HG12	3:HD:217:ILE:HD13	1.91	0.53
2:MD:95:GLU:HG3	2:MD:112:PHE:CE2	2.43	0.53

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:ZD:247:ARG:HH11	3:ZD:247:ARG:HA	1.74	0.53
2:EE:130:ILE:HD11	2:EE:157:VAL:HG22	1.90	0.53
2:EE:264:LYS:HD2	2:EE:265:GLY:N	2.24	0.53
2:KE:109:THR:O	2:KE:113:MET:HG3	2.08	0.53
2:WE:120:ASP:OD2	3:DF:76:SER:OG	2.19	0.53
2:IF:89:ARG:HG3	2:IF:89:ARG:HH11	1.74	0.53
3:JF:229:GLU:OE1	3:JF:230:LEU:HD12	2.09	0.53
2:UF:239:LEU:HD11	2:UF:309:LEU:HD11	1.91	0.53
2:AG:302:GLU:O	2:AG:306:LYS:HG2	2.09	0.53
2:SG:297:ARG:HG3	2:SG:299:SER:H	1.74	0.53
2:G:48:SER:OG	2:G:51:GLN:OE1	2.27	0.53
2:G:74:ASN:O	2:G:78:ARG:NE	2.42	0.53
3:C:239:SER:HA	3:C:242:GLU:OE1	2.09	0.53
4:KA:106:ILE:O	3:NA:252:ARG:HG2	2.09	0.53
2:SA:151:GLU:HA	2:SA:154:ARG:HG2	1.90	0.53
2:YA:65:GLN:NE2	2:EB:47:ILE:O	2.33	0.53
3:ZA:112:GLY:HA3	3:ZA:223:MET:HG2	1.89	0.53
2:KB:264:LYS:HD2	2:KB:265:GLY:N	2.24	0.53
3:LB:59:GLU:OE1	3:LB:59:GLU:N	2.42	0.53
2:QB:9:LYS:HG2	2:QB:55:VAL:HG22	1.91	0.53
3:XB:112:GLY:HA3	3:XB:223:MET:HG2	1.92	0.53
2:OC:84:ALA:O	2:OC:87:GLU:HB3	2.09	0.53
2:OC:89:ARG:HG3	2:OC:90:ALA:N	2.25	0.53
2:GD:188:LEU:HB3	3:ND:74:ARG:NE	2.24	0.53
3:ND:240:ARG:HA	3:ND:240:ARG:HH11	1.73	0.53
3:TD:100:THR:HG23	3:TD:119:SER:HA	1.90	0.53
2:YD:21:ARG:O	2:YD:21:ARG:NE	2.25	0.53
4:CE:106:ILE:H	3:FE:252:ARG:CZ	2.22	0.53
2:KE:297:ARG:HG3	2:KE:299:SER:H	1.74	0.53
3:LE:86:ILE:HD11	3:LE:221:PHE:CE1	2.44	0.53
3:RE:54:LEU:HA	3:RE:57:ILE:HG22	1.91	0.53
3:RE:119:SER:HB3	3:RE:214:GLU:O	2.09	0.53
2:WE:219:ARG:NH2	2:WE:227:GLN:OE1	2.30	0.53
2:UF:69:LEU:HD13	2:AG:45:ARG:O	2.09	0.53
2:UF:167:VAL:HG21	3:VF:131:PHE:CD1	2.44	0.53
3:BG:120:PRO:HA	3:BG:123:VAL:HG22	1.91	0.53
2:MG:153:LEU:O	2:MG:157:VAL:HG23	2.08	0.53
2:MG:297:ARG:HG3	2:MG:299:SER:H	1.73	0.53
2:B:239:LEU:HD11	2:B:309:LEU:HD11	1.92	0.52
2:B:302:GLU:O	2:B:306:LYS:HG2	2.09	0.52
3:C:238:ASN:HB3	3:C:241:HIS:ND1	2.24	0.52

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:T:40:ALA:O	2:T:44:VAL:HG23	2.09	0.52
2:T:95:GLU:HG3	2:T:112:PHE:CE2	2.44	0.52
2:AA:130:ILE:O	2:AA:134:LEU:HG	2.10	0.52
3:NA:106:HIS:CD2	3:NA:113:THR:HG22	2.45	0.52
2:YA:245:ARG:O	2:YA:245:ARG:NH1	2.37	0.52
2:QB:65:GLN:NE2	2:WB:47:ILE:O	2.36	0.52
2:WB:69:LEU:HD13	2:CC:45:ARG:O	2.09	0.52
2:WB:87:GLU:HG2	2:WB:89:ARG:N	2.24	0.52
2:IC:9:LYS:HG2	2:IC:55:VAL:HG22	1.91	0.52
2:AD:130:ILE:HD11	2:AD:157:VAL:HG22	1.90	0.52
2:MD:75:GLU:O	2:MD:79:SER:HB3	2.09	0.52
2:MD:302:GLU:O	2:MD:306:LYS:HG2	2.09	0.52
4:WD:106:ILE:C	3:ZD:252:ARG:HD3	2.30	0.52
2:YD:245:ARG:O	2:YD:245:ARG:NH1	2.37	0.52
3:ZD:120:PRO:HA	3:ZD:123:VAL:HG22	1.90	0.52
2:EE:302:GLU:O	2:EE:306:LYS:HG2	2.09	0.52
2:CF:219:ARG:NH2	2:CF:227:GLN:OE1	2.32	0.52
2:CF:239:LEU:HD11	2:CF:309:LEU:HD11	1.91	0.52
3:C:101:ASN:ND2	3:C:154:ILE:HG21	2.24	0.52
2:T:201:GLU:CG	2:AA:139:ARG:HH21	2.23	0.52
2:T:302:GLU:O	2:T:306:LYS:HG2	2.10	0.52
2:WB:79:SER:HA	2:WB:82:VAL:HG22	1.90	0.52
2:WB:201:GLU:HG2	2:CC:139:ARG:HD2	1.92	0.52
2:CC:122:ILE:HB	2:CC:130:ILE:HG22	1.91	0.52
3:BD:102:LEU:HD22	3:BD:117:VAL:HG22	1.91	0.52
3:HD:101:ASN:ND2	3:HD:154:ILE:HG21	2.24	0.52
2:SD:198:THR:O	2:SD:202:ILE:HG12	2.09	0.52
3:ZD:134:ASP:OD2	3:ZD:136:ARG:N	2.29	0.52
2:QE:115:PRO:O	2:QE:119:ALA:CB	2.58	0.52
2:CF:207:LYS:NZ	2:CF:209:GLN:HB3	2.25	0.52
2:OF:302:GLU:O	2:OF:306:LYS:HG2	2.09	0.52
2:UF:186:GLN:C	3:BG:74:ARG:HH22	2.13	0.52
3:VF:198:ILE:HG13	3:VF:221:PHE:CE1	2.45	0.52
2:GG:31:THR:O	2:GG:34:VAL:HB	2.09	0.52
2:GG:168:GLN:NE2	3:HG:130:LEU:HD11	2.25	0.52
3:TG:86:ILE:HD11	3:TG:221:PHE:CE1	2.45	0.52
2:B:76:TYR:HE2	2:J:26:PHE:CE2	2.27	0.52
2:GA:76:TYR:HB2	2:MA:41:MET:HE3	1.91	0.52
2:GA:89:ARG:HG2	2:GA:92:SER:HB3	1.92	0.52
3:HA:120:PRO:HA	3:HA:123:VAL:HG22	1.89	0.52
2:SA:128:GLN:HG2	3:TA:131:PHE:CB	2.37	0.52

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:YA:75:GLU:O	2:YA:79:SER:HB3	2.08	0.52
2:YA:202:ILE:HD13	2:EB:139:ARG:HD3	1.91	0.52
2:QB:31:THR:O	2:QB:34:VAL:HB	2.09	0.52
2:QB:100:THR:OG1	2:QB:101:ARG:NH2	2.43	0.52
2:WB:186:GLN:C	3:DC:74:ARG:HH22	2.12	0.52
3:XB:182:SER:OG	3:XB:183:GLU:N	2.41	0.52
2:IC:11:VAL:O	2:IC:15:MET:HE3	2.09	0.52
2:IC:111:ASN:OD1	2:IC:138:LYS:N	2.23	0.52
2:AD:31:THR:O	2:AD:34:VAL:HB	2.10	0.52
2:AD:302:GLU:O	2:AD:306:LYS:HG2	2.10	0.52
3:BD:180:VAL:HG23	3:BD:181:ARG:H	1.75	0.52
2:GD:14:LEU:HD12	2:GD:22:ALA:HB1	1.92	0.52
2:MD:69:LEU:HD21	2:SD:47:ILE:HB	1.91	0.52
3:ND:112:GLY:HA3	3:ND:223:MET:HG2	1.91	0.52
2:SD:186:GLN:C	3:ZD:74:ARG:HH22	2.12	0.52
2:YD:126:HIS:CD2	3:ZD:127:VAL:HG13	2.43	0.52
3:FE:86:ILE:HG13	3:FE:198:ILE:HD11	1.91	0.52
3:LE:191:ILE:HG13	3:LE:192:THR:HG23	1.90	0.52
2:QE:107:ILE:HD12	2:QE:107:ILE:H	1.73	0.52
3:DF:119:SER:HB3	3:DF:214:GLU:O	2.09	0.52
2:UF:9:LYS:HG2	2:UF:55:VAL:HG22	1.90	0.52
2:UF:115:PRO:O	2:UF:119:ALA:CB	2.57	0.52
2:MG:114:GLU:HB2	2:MG:117:SER:OG	2.08	0.52
3:NG:118:PHE:CE2	3:NG:158:LEU:HD12	2.45	0.52
2:B:264:LYS:HD2	2:B:265:GLY:N	2.24	0.52
2:J:21:ARG:HH21	2:J:25:VAL:N	2.08	0.52
2:T:153:LEU:O	2:T:157:VAL:HG23	2.08	0.52
3:HA:101:ASN:ND2	3:HA:154:ILE:HG21	2.25	0.52
3:TA:100:THR:HG23	3:TA:119:SER:HA	1.91	0.52
3:FB:182:SER:OG	3:FB:183:GLU:N	2.39	0.52
3:LB:58:ASN:OD1	3:LB:81:VAL:HG11	2.09	0.52
2:QB:264:LYS:HD2	2:QB:265:GLY:N	2.25	0.52
2:WB:30:SER:O	2:WB:34:VAL:HG23	2.09	0.52
2:CC:302:GLU:O	2:CC:306:LYS:HG2	2.09	0.52
3:DC:191:ILE:HG13	3:DC:192:THR:HG23	1.91	0.52
3:DC:224:ILE:O	3:DC:228:ARG:N	2.43	0.52
2:IC:297:ARG:HG3	2:IC:299:SER:H	1.74	0.52
2:MD:178:VAL:HG11	3:ND:149:THR:HG22	1.91	0.52
2:SD:107:ILE:H	2:SD:107:ILE:HD12	1.74	0.52
2:SD:264:LYS:HD2	2:SD:265:GLY:N	2.25	0.52
3:TD:112:GLY:HA3	3:TD:223:MET:HG2	1.92	0.52

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:QE:76:TYR:HE1	2:WE:26:PHE:CE2	2.26	0.52
3:RE:183:GLU:HG3	3:RE:188:PHE:HB2	1.90	0.52
2:CF:98:LEU:HB3	2:CF:108:GLU:OE1	2.09	0.52
3:DF:214:GLU:OE1	3:DF:215:PHE:N	2.43	0.52
2:IF:188:LEU:H	3:PF:74:ARG:HH21	1.55	0.52
2:OF:83:LYS:HZ3	2:UF:26:PHE:HD2	1.54	0.52
2:UF:128:GLN:HG2	3:VF:131:PHE:CB	2.34	0.52
2:UF:264:LYS:HD2	2:UF:265:GLY:N	2.25	0.52
2:AG:264:LYS:HD2	2:AG:265:GLY:N	2.25	0.52
2:GG:69:LEU:HD13	2:MG:45:ARG:O	2.09	0.52
2:GG:95:GLU:HG3	2:GG:112:PHE:CE2	2.44	0.52
2:GG:134:LEU:HD13	2:GG:146:LEU:HD21	1.91	0.52
2:MG:245:ARG:O	2:MG:245:ARG:NH1	2.37	0.52
2:G:41:MET:HG2	2:G:42:ALA:N	2.25	0.52
2:G:151:GLU:HB3	2:G:155:HIS:CE1	2.45	0.52
2:B:76:TYR:HE2	2:J:26:PHE:CZ	2.28	0.52
2:B:190:ARG:O	2:J:166:GLY:HA3	2.09	0.52
2:J:67:ALA:O	2:J:71:ILE:HG12	2.09	0.52
2:T:114:GLU:HB2	2:T:117:SER:OG	2.08	0.52
2:MA:69:LEU:HD13	2:SA:45:ARG:O	2.08	0.52
3:NA:75:ARG:HG3	3:NA:208:ILE:HD11	1.90	0.52
2:YA:128:GLN:HG2	3:ZA:131:PHE:CB	2.37	0.52
2:KB:153:LEU:O	2:KB:157:VAL:HG23	2.09	0.52
2:KB:203:ILE:HG12	2:KB:206:MET:HE3	1.92	0.52
2:CC:141:GLN:HA	2:CC:144:ASP:OD2	2.10	0.52
2:IC:203:ILE:HA	2:IC:206:MET:HE2	1.91	0.52
3:HD:75:ARG:HG3	3:HD:208:ILE:HD11	1.92	0.52
2:MD:167:VAL:HG13	3:ND:130:LEU:O	2.10	0.52
2:SD:128:GLN:HG2	3:TD:131:PHE:CB	2.34	0.52
2:SD:151:GLU:HA	2:SD:154:ARG:HG2	1.91	0.52
2:SD:239:LEU:HD11	2:SD:309:LEU:HD11	1.91	0.52
2:SD:302:GLU:O	2:SD:306:LYS:HG2	2.09	0.52
3:RE:96:LEU:HD23	3:RE:97:PRO:CD	2.40	0.52
3:RE:190:ASN:ND2	3:XE:234:PRO:O	2.43	0.52
3:XE:198:ILE:HG23	3:XE:221:PHE:CD1	2.45	0.52
3:XE:225:GLU:HA	3:XE:228:ARG:HB3	1.91	0.52
2:CF:168:GLN:HE21	2:CF:171:ALA:H	1.57	0.52
3:DF:116:VAL:HG21	3:DF:158:LEU:HD11	1.91	0.52
2:OF:151:GLU:HA	2:OF:154:ARG:HG2	1.92	0.52
2:AG:120:ASP:OD1	2:AG:123:ARG:NH1	2.43	0.52
2:MG:126:HIS:CE1	3:NG:127:VAL:HG13	2.45	0.52

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:MG:244:ASP:OD1	2:MG:276:ARG:NH2	2.43	0.52
3:NG:190:ASN:ND2	3:TG:234:PRO:O	2.42	0.52
2:G:130:ILE:HD11	2:G:157:VAL:HG22	1.92	0.52
2:B:126:HIS:CD2	3:C:127:VAL:HG13	2.44	0.52
3:C:116:VAL:HG12	3:C:217:ILE:HD13	1.92	0.52
2:J:153:LEU:O	2:J:157:VAL:HG23	2.09	0.52
2:J:264:LYS:HD2	2:J:265:GLY:N	2.25	0.52
2:J:302:GLU:O	2:J:306:LYS:HG2	2.10	0.52
2:T:186:GLN:NE2	3:BA:71:ASN:HA	2.19	0.52
2:T:264:LYS:HD2	2:T:265:GLY:N	2.25	0.52
2:YA:207:LYS:NZ	2:YA:209:GLN:HB3	2.21	0.52
2:YA:239:LEU:HD11	2:YA:309:LEU:HD11	1.91	0.52
2:EB:11:VAL:O	2:EB:15:MET:HE3	2.09	0.52
3:FB:191:ILE:HG13	3:FB:192:THR:HG23	1.91	0.52
2:KB:78:ARG:O	2:KB:82:VAL:HG23	2.09	0.52
3:RB:238:ASN:HB3	3:RB:241:HIS:ND1	2.24	0.52
2:WB:87:GLU:HG2	2:WB:89:ARG:H	1.74	0.52
2:WB:151:GLU:HB3	2:WB:155:HIS:CE1	2.45	0.52
2:CC:9:LYS:HG2	2:CC:55:VAL:HG22	1.91	0.52
2:UC:264:LYS:HD2	2:UC:265:GLY:N	2.25	0.52
2:AD:264:LYS:HD2	2:AD:265:GLY:N	2.25	0.52
2:MD:31:THR:O	2:MD:34:VAL:HB	2.09	0.52
2:MD:264:LYS:HD2	2:MD:265:GLY:N	2.25	0.52
2:SD:230:ILE:HA	2:SD:233:MET:HE2	1.91	0.52
3:TD:102:LEU:HD22	3:TD:117:VAL:HG22	1.90	0.52
3:LE:111:ARG:HB2	3:LE:111:ARG:NH1	2.24	0.52
3:LE:112:GLY:HA3	3:LE:223:MET:HG2	1.91	0.52
2:QE:152:ARG:NH2	2:QE:153:LEU:HB2	2.25	0.52
2:QE:302:GLU:O	2:QE:306:LYS:HG2	2.09	0.52
2:CF:9:LYS:HG2	2:CF:55:VAL:HG22	1.92	0.52
3:JF:59:GLU:N	3:JF:59:GLU:OE2	2.43	0.52
2:UF:89:ARG:HH21	2:UF:93:LEU:HD21	1.74	0.52
2:AG:128:GLN:HG2	3:BG:131:PHE:CB	2.36	0.52
3:HG:106:HIS:CD2	3:HG:113:THR:HG22	2.44	0.52
2:MG:11:VAL:O	2:MG:14:LEU:HD12	2.10	0.52
2:SG:239:LEU:HD11	2:SG:309:LEU:HD11	1.92	0.52
3:M:100:THR:HG23	3:M:119:SER:HA	1.91	0.52
2:AA:302:GLU:O	2:AA:306:LYS:HG2	2.09	0.52
2:MA:188:LEU:HB3	3:TA:74:ARG:NE	2.25	0.52
2:SA:188:LEU:HD13	3:ZA:74:ARG:CD	2.32	0.52
2:YA:264:LYS:HD2	2:YA:265:GLY:N	2.25	0.52

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:ZA:198:ILE:HG23	3:ZA:221:PHE:CD1	2.45	0.52
2:EB:84:ALA:O	2:EB:87:GLU:HB3	2.10	0.52
2:EB:151:GLU:O	2:EB:155:HIS:ND1	2.43	0.52
2:KB:239:LEU:HD11	2:KB:309:LEU:HD11	1.91	0.52
2:WB:31:THR:O	2:WB:34:VAL:HB	2.10	0.52
3:XB:65:PHE:CE1	3:XB:161:ALA:HB2	2.45	0.52
2:CC:186:GLN:NE2	3:JC:71:ASN:HA	2.23	0.52
3:JC:96:LEU:HD23	3:JC:97:PRO:CD	2.39	0.52
2:AD:89:ARG:HG3	2:AD:90:ALA:N	2.25	0.52
2:AD:205:LEU:O	2:AD:205:LEU:HD23	2.10	0.52
2:GD:264:LYS:HD2	2:GD:265:GLY:N	2.25	0.52
2:MD:30:SER:OG	2:MD:33:GLU:OE1	2.28	0.52
2:MD:78:ARG:O	2:MD:82:VAL:HG23	2.10	0.52
2:MD:203:ILE:HA	2:MD:206:MET:CE	2.38	0.52
2:EE:14:LEU:HD12	2:EE:22:ALA:HB1	1.92	0.52
3:FE:117:VAL:O	3:FE:215:PHE:HB2	2.10	0.52
2:KE:83:LYS:HB3	2:QE:23:ALA:O	2.09	0.52
2:QE:264:LYS:HD2	2:QE:265:GLY:N	2.25	0.52
3:RE:102:LEU:HD22	3:RE:117:VAL:HG22	1.91	0.52
4:UE:106:ILE:O	3:XE:252:ARG:HG2	2.09	0.52
2:CF:221:PHE:CE2	2:IF:159:LEU:HD11	2.45	0.52
2:IF:157:VAL:O	2:IF:161:ILE:HG12	2.09	0.52
2:OF:239:LEU:HD11	2:OF:309:LEU:HD11	1.91	0.52
2:UF:208:THR:HA	2:UF:211:GLU:HB2	1.90	0.52
2:UF:297:ARG:HG3	2:UF:299:SER:H	1.73	0.52
3:VF:225:GLU:HA	3:VF:228:ARG:HB3	1.91	0.52
3:BG:202:THR:O	3:BG:202:THR:OG1	2.27	0.52
3:NG:100:THR:HG23	3:NG:119:SER:HA	1.91	0.52
3:C:77:PRO:HB3	3:C:208:ILE:HB	1.91	0.52
3:C:97:PRO:O	3:C:100:THR:OG1	2.19	0.52
2:T:239:LEU:HD11	2:T:309:LEU:HD11	1.91	0.52
2:AA:89:ARG:HG3	2:AA:90:ALA:N	2.24	0.52
2:AA:264:LYS:HD2	2:AA:265:GLY:N	2.25	0.52
2:GA:239:LEU:HD11	2:GA:309:LEU:HD11	1.92	0.52
3:HA:202:THR:O	3:HA:202:THR:OG1	2.28	0.52
2:MA:297:ARG:HG3	2:MA:299:SER:H	1.74	0.52
3:NA:102:LEU:HD22	3:NA:117:VAL:HG22	1.92	0.52
2:SA:127:PRO:O	2:SA:130:ILE:HG12	2.09	0.52
2:YA:120:ASP:OD2	3:FB:76:SER:OG	2.19	0.52
2:KB:31:THR:O	2:KB:34:VAL:HB	2.09	0.52
2:KB:187:ASN:N	3:RB:74:ARG:HH22	2.08	0.52

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:KB:203:ILE:HA	2:KB:206:MET:CE	2.40	0.52
3:LB:236:LEU:HB3	3:LB:238:ASN:ND2	2.23	0.52
2:WB:9:LYS:HG2	2:WB:55:VAL:HG22	1.92	0.52
2:WB:186:GLN:NE2	3:DC:71:ASN:HA	2.22	0.52
2:WB:230:ILE:HA	2:WB:233:MET:HE2	1.92	0.52
2:CC:75:GLU:HA	2:CC:78:ARG:HH21	1.75	0.52
2:UC:297:ARG:HG3	2:UC:299:SER:H	1.74	0.52
3:ND:191:ILE:HG13	3:ND:192:THR:HG23	1.91	0.52
2:EE:95:GLU:HG3	2:EE:112:PHE:CE2	2.45	0.52
2:EE:153:LEU:O	2:EE:157:VAL:HG23	2.10	0.52
3:FE:97:PRO:O	3:FE:100:THR:OG1	2.17	0.52
2:KE:264:LYS:HD2	2:KE:265:GLY:N	2.25	0.52
2:WE:83:LYS:HD3	2:CF:26:PHE:CD2	2.45	0.52
3:XE:110:LEU:HG	3:XE:223:MET:HB3	1.92	0.52
2:IF:8:ASP:O	2:IF:12:ILE:HG12	2.10	0.52
2:IF:264:LYS:HD2	2:IF:265:GLY:N	2.24	0.52
3:JF:118:PHE:CE2	3:JF:158:LEU:HD12	2.45	0.52
2:OF:151:GLU:HB3	2:OF:155:HIS:CE1	2.44	0.52
2:OF:264:LYS:HD2	2:OF:265:GLY:N	2.25	0.52
2:GG:297:ARG:HG3	2:GG:299:SER:H	1.74	0.52
2:MG:128:GLN:HG2	3:NG:131:PHE:CB	2.34	0.52
3:NG:97:PRO:O	3:NG:97:PRO:HD2	2.08	0.52
3:TG:96:LEU:HD23	3:TG:97:PRO:HD2	1.91	0.52
2:G:239:LEU:HD11	2:G:309:LEU:HD11	1.91	0.52
2:B:11:VAL:HG11	2:B:41:MET:HB3	1.91	0.52
2:AA:137:LEU:HB3	2:AA:141:GLN:HE21	1.75	0.52
2:GA:264:LYS:HD2	2:GA:265:GLY:N	2.24	0.52
2:MA:130:ILE:HD11	2:MA:157:VAL:HG22	1.92	0.52
2:SA:264:LYS:HD2	2:SA:265:GLY:N	2.25	0.52
2:YA:151:GLU:HB3	2:YA:155:HIS:CE1	2.45	0.52
2:EB:264:LYS:HD2	2:EB:265:GLY:N	2.25	0.52
3:FB:198:ILE:HG23	3:FB:221:PHE:CD1	2.45	0.52
2:KB:152:ARG:HA	2:KB:155:HIS:HE1	1.75	0.52
3:LB:118:PHE:CE2	3:LB:158:LEU:HD12	2.45	0.52
2:WB:115:PRO:O	2:WB:119:ALA:HB2	2.10	0.52
3:XB:202:THR:OG1	3:XB:202:THR:O	2.28	0.52
2:CC:202:ILE:HD13	2:IC:139:ARG:HH21	1.74	0.52
2:CC:264:LYS:HD2	2:CC:265:GLY:N	2.25	0.52
3:DC:120:PRO:HA	3:DC:123:VAL:HG22	1.91	0.52
2:AD:9:LYS:HG2	2:AD:55:VAL:HG22	1.92	0.52
2:AD:278:MET:N	2:AD:278:MET:SD	2.83	0.52

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:BD:100:THR:HG23	3:BD:119:SER:HA	1.90	0.52
3:HD:118:PHE:CE2	3:HD:158:LEU:HD12	2.44	0.52
2:MD:12:ILE:HD11	2:MD:55:VAL:HG11	1.91	0.52
2:MD:205:LEU:CD2	2:SD:140:SER:HA	2.40	0.52
2:YD:230:ILE:HA	2:YD:233:MET:CE	2.40	0.52
3:LE:96:LEU:HD23	3:LE:97:PRO:CD	2.39	0.52
2:QE:9:LYS:HG2	2:QE:55:VAL:HG22	1.91	0.52
2:IF:33:GLU:O	2:IF:37:LEU:HG	2.09	0.52
2:AG:278:MET:N	2:AG:278:MET:SD	2.83	0.52
2:GG:239:LEU:HD11	2:GG:309:LEU:HD11	1.92	0.52
2:GG:264:LYS:HD2	2:GG:265:GLY:N	2.25	0.52
2:MG:264:LYS:HD2	2:MG:265:GLY:N	2.25	0.52
3:NG:180:VAL:HG23	3:NG:181:ARG:H	1.75	0.52
3:M:180:VAL:HG23	3:M:181:ARG:H	1.75	0.52
2:B:242:VAL:HG13	2:B:323:ILE:HG12	1.90	0.52
4:R:106:ILE:H	3:V:252:ARG:NE	2.07	0.52
2:GA:153:LEU:O	2:GA:157:VAL:HG23	2.10	0.52
2:MA:207:LYS:HZ3	2:MA:209:GLN:HB3	1.76	0.52
3:NA:119:SER:HB3	3:NA:214:GLU:O	2.09	0.52
2:SA:34:VAL:O	2:SA:37:LEU:HD12	2.10	0.52
2:SA:101:ARG:O	2:SA:104:THR:HG23	2.10	0.52
2:SA:239:LEU:HD11	2:SA:309:LEU:HD11	1.92	0.52
3:FB:59:GLU:N	3:FB:59:GLU:OE2	2.43	0.52
3:FB:97:PRO:O	3:FB:100:THR:OG1	2.22	0.52
2:OC:110:LEU:HB3	2:OC:141:GLN:HE22	1.75	0.52
2:GD:142:ALA:O	2:GD:146:LEU:HG	2.09	0.52
2:GD:167:VAL:HG13	3:HD:130:LEU:O	2.10	0.52
3:TD:118:PHE:CE2	3:TD:158:LEU:HD12	2.44	0.52
2:EE:83:LYS:O	2:KE:27:LYS:HE2	2.10	0.52
2:EE:114:GLU:HB2	2:EE:117:SER:OG	2.08	0.52
3:FE:116:VAL:HG21	3:FE:158:LEU:HD11	1.92	0.52
3:LE:225:GLU:HA	3:LE:228:ARG:HB3	1.92	0.52
3:JF:227:LEU:HD13	3:JF:230:LEU:HD13	1.91	0.52
2:UF:108:GLU:HA	2:UF:111:ASN:HD22	1.75	0.52
2:UF:278:MET:N	2:UF:278:MET:SD	2.83	0.52
2:AG:83:LYS:HD3	2:GG:26:PHE:HD1	1.74	0.52
2:GG:11:VAL:O	2:GG:15:MET:HE3	2.10	0.52
2:MG:83:LYS:O	2:SG:27:LYS:HE2	2.10	0.52
2:MG:230:ILE:HA	2:MG:233:MET:HE2	1.92	0.52
3:NG:90:HIS:CE1	3:NG:91:GLU:HG2	2.45	0.52
2:G:264:LYS:HD2	2:G:265:GLY:N	2.25	0.51

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:M:238:ASN:HB3	3:M:241:HIS:ND1	2.25	0.51
2:GA:244:ASP:HA	2:GA:247:ILE:HG12	1.93	0.51
2:MA:74:ASN:O	2:MA:78:ARG:NE	2.44	0.51
2:MA:133:ILE:O	2:MA:137:LEU:HG	2.10	0.51
2:MA:245:ARG:O	2:MA:245:ARG:NH1	2.37	0.51
4:WA:106:ILE:O	3:ZA:252:ARG:HG2	2.10	0.51
2:YA:126:HIS:CE1	3:ZA:127:VAL:HG13	2.45	0.51
3:JC:134:ASP:OD2	3:JC:136:ARG:N	2.37	0.51
2:UC:190:ARG:O	2:AD:166:GLY:HA3	2.10	0.51
2:GD:114:GLU:HB2	2:GD:117:SER:OG	2.09	0.51
2:GD:151:GLU:HA	2:GD:154:ARG:HG2	1.91	0.51
3:ND:106:HIS:CD2	3:ND:113:THR:HG22	2.45	0.51
2:SD:79:SER:OG	2:SD:83:LYS:NZ	2.42	0.51
2:YD:9:LYS:HG2	2:YD:55:VAL:HG22	1.92	0.51
2:QE:239:LEU:HD11	2:QE:309:LEU:HD11	1.91	0.51
2:WE:80:VAL:O	2:WE:83:LYS:HG2	2.10	0.51
2:WE:151:GLU:HA	2:WE:154:ARG:HE	1.74	0.51
2:IF:278:MET:N	2:IF:278:MET:SD	2.83	0.51
2:UF:67:ALA:O	2:UF:71:ILE:HG12	2.11	0.51
3:VF:119:SER:HB3	3:VF:214:GLU:O	2.08	0.51
2:SG:264:LYS:HD2	2:SG:265:GLY:N	2.25	0.51
3:TG:117:VAL:O	3:TG:215:PHE:HB2	2.10	0.51
2:G:310:LEU:HD12	2:G:311:ILE:HD13	1.92	0.51
2:J:151:GLU:HB3	2:J:155:HIS:CE1	2.45	0.51
2:J:188:LEU:HB3	3:V:74:ARG:NE	2.25	0.51
2:AA:239:LEU:HD11	2:AA:309:LEU:HD11	1.92	0.51
3:NA:101:ASN:ND2	3:NA:154:ILE:HG21	2.26	0.51
2:EB:128:GLN:CG	3:FB:131:PHE:HB3	2.35	0.51
3:LB:202:THR:O	3:LB:202:THR:OG1	2.28	0.51
2:WB:107:ILE:H	2:WB:107:ILE:HD12	1.75	0.51
3:XB:185:GLN:HG3	3:XB:186:VAL:H	1.76	0.51
4:YB:101:ILE:N	4:YB:104:TYR:O	2.34	0.51
2:CC:79:SER:HA	2:CC:82:VAL:HG12	1.92	0.51
2:CC:95:GLU:HG3	2:CC:112:PHE:CE2	2.45	0.51
3:DC:72:LEU:HD23	3:DC:73:LEU:HD12	1.91	0.51
2:OC:264:LYS:HD2	2:OC:265:GLY:N	2.25	0.51
2:UC:187:ASN:N	3:BD:74:ARG:HH22	2.08	0.51
2:SD:205:LEU:CD2	2:YD:140:SER:HA	2.40	0.51
2:SD:245:ARG:O	2:SD:245:ARG:NH1	2.37	0.51
2:YD:239:LEU:HD11	2:YD:309:LEU:HD11	1.92	0.51
2:EE:11:VAL:O	2:EE:15:MET:HE3	2.09	0.51

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:KE:239:LEU:HD11	2:KE:309:LEU:HD11	1.92	0.51
3:LE:101:ASN:ND2	3:LE:154:ILE:HG21	2.24	0.51
3:LE:115:LEU:HD21	3:LE:218:CYS:HB3	1.92	0.51
3:LE:116:VAL:HG21	3:LE:158:LEU:HD11	1.92	0.51
2:QE:151:GLU:HB3	2:QE:155:HIS:CE1	2.45	0.51
2:WE:264:LYS:HD2	2:WE:265:GLY:N	2.25	0.51
2:IF:239:LEU:HD11	2:IF:309:LEU:HD11	1.92	0.51
3:JF:72:LEU:HD23	3:JF:73:LEU:HD12	1.91	0.51
3:BG:97:PRO:HD2	3:BG:97:PRO:O	2.09	0.51
2:GG:69:LEU:O	2:GG:72:ASN:ND2	2.40	0.51
2:GG:128:GLN:HG2	3:HG:131:PHE:CB	2.36	0.51
3:HG:100:THR:HG23	3:HG:119:SER:HA	1.92	0.51
3:HG:119:SER:HB3	3:HG:214:GLU:O	2.10	0.51
2:MG:110:LEU:HB3	2:MG:141:GLN:HE22	1.74	0.51
2:G:8:ASP:O	2:G:12:ILE:HG12	2.10	0.51
2:B:8:ASP:O	2:B:12:ILE:HG12	2.11	0.51
2:B:107:ILE:H	2:B:107:ILE:HD12	1.75	0.51
2:AA:115:PRO:O	2:AA:119:ALA:CB	2.59	0.51
2:GA:114:GLU:HB2	2:GA:117:SER:OG	2.10	0.51
2:MA:152:ARG:HH12	2:MA:153:LEU:HB2	1.76	0.51
2:MA:230:ILE:HA	2:MA:233:MET:HE2	1.92	0.51
2:SA:152:ARG:NH2	2:SA:153:LEU:HB2	2.24	0.51
2:SA:245:ARG:O	2:SA:245:ARG:NH1	2.38	0.51
2:EB:282:ALA:HA	2:EB:285:ILE:HG12	1.92	0.51
2:CC:278:MET:SD	2:CC:278:MET:N	2.83	0.51
2:IC:264:LYS:HD2	2:IC:265:GLY:N	2.25	0.51
3:JC:119:SER:HB3	3:JC:214:GLU:O	2.10	0.51
2:OC:239:LEU:HD11	2:OC:309:LEU:HD11	1.92	0.51
2:UC:130:ILE:O	2:UC:134:LEU:HD13	2.11	0.51
3:BD:96:LEU:HD23	3:BD:97:PRO:HD2	1.92	0.51
2:GD:8:ASP:O	2:GD:12:ILE:HG12	2.11	0.51
2:GD:151:GLU:HB3	2:GD:155:HIS:CE1	2.44	0.51
3:HD:227:LEU:HD13	3:HD:230:LEU:HD13	1.92	0.51
2:MD:239:LEU:HD11	2:MD:309:LEU:HD11	1.91	0.51
3:ND:74:ARG:O	3:ND:75:ARG:HD2	2.10	0.51
2:SD:230:ILE:HA	2:SD:233:MET:CE	2.40	0.51
3:TD:190:ASN:OD1	3:ZD:236:LEU:HB2	2.10	0.51
4:OE:106:ILE:O	3:RE:252:ARG:HG2	2.10	0.51
2:QE:108:GLU:HA	2:QE:111:ASN:HD22	1.76	0.51
3:RE:55:GLU:HA	3:RE:58:ASN:ND2	2.26	0.51
3:RE:180:VAL:HG23	3:RE:181:ARG:H	1.75	0.51

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:XE:54:LEU:HD13	3:XE:231:LEU:HD21	1.93	0.51
3:XE:58:ASN:OD1	3:XE:81:VAL:HG11	2.10	0.51
2:IF:153:LEU:O	2:IF:157:VAL:HG23	2.11	0.51
2:IF:190:ARG:O	2:OF:166:GLY:HA3	2.11	0.51
2:IF:205:LEU:CD2	2:OF:140:SER:HA	2.41	0.51
2:OF:128:GLN:CG	3:PF:131:PHE:HB3	2.35	0.51
2:UF:168:GLN:NE2	3:VF:130:LEU:HD11	2.25	0.51
3:VF:185:GLN:HG3	3:VF:186:VAL:H	1.76	0.51
2:SG:130:ILE:O	2:SG:134:LEU:HD13	2.10	0.51
2:SG:151:GLU:HA	2:SG:154:ARG:HG2	1.93	0.51
2:SG:160:ARG:O	2:SG:164:PHE:CB	2.56	0.51
2:G:168:GLN:NE2	3:M:130:LEU:HD11	2.25	0.51
2:J:66:PHE:CB	2:T:45:ARG:HD2	2.40	0.51
2:AA:38:SER:HA	2:AA:41:MET:HE1	1.92	0.51
3:BA:120:PRO:HA	3:BA:123:VAL:HG22	1.91	0.51
2:YA:69:LEU:HD13	2:EB:45:ARG:O	2.10	0.51
3:ZA:102:LEU:HD22	3:ZA:117:VAL:HG22	1.93	0.51
2:KB:97:ILE:O	2:KB:101:ARG:NH2	2.33	0.51
2:WB:264:LYS:HD2	2:WB:265:GLY:N	2.25	0.51
2:IC:12:ILE:HD11	2:IC:55:VAL:HG11	1.92	0.51
3:PC:214:GLU:OE1	3:PC:215:PHE:N	2.43	0.51
2:UC:167:VAL:HG13	3:VC:130:LEU:O	2.10	0.51
2:UC:278:MET:N	2:UC:278:MET:SD	2.84	0.51
3:VC:100:THR:HG23	3:VC:119:SER:HA	1.92	0.51
2:GD:21:ARG:HE	2:GD:21:ARG:C	2.10	0.51
2:SD:205:LEU:HD21	2:YD:140:SER:HA	1.93	0.51
3:FE:101:ASN:ND2	3:FE:154:ILE:HG21	2.25	0.51
2:KE:278:MET:N	2:KE:278:MET:SD	2.84	0.51
3:RE:100:THR:HG23	3:RE:119:SER:HA	1.91	0.51
2:CF:278:MET:SD	2:CF:278:MET:N	2.84	0.51
3:DF:77:PRO:HB3	3:DF:208:ILE:HB	1.92	0.51
3:JF:183:GLU:HG3	3:JF:188:PHE:HB2	1.92	0.51
2:UF:244:ASP:OD1	2:UF:276:ARG:NH2	2.43	0.51
3:VF:101:ASN:HB2	3:VF:146:PHE:CE2	2.45	0.51
2:GG:115:PRO:O	2:GG:119:ALA:HB2	2.10	0.51
3:HG:180:VAL:HG23	3:HG:181:ARG:H	1.75	0.51
2:G:152:ARG:HH12	2:G:153:LEU:HB2	1.75	0.51
2:B:128:GLN:HG2	3:C:131:PHE:CB	2.37	0.51
3:K:183:GLU:HG3	3:K:188:PHE:HB2	1.91	0.51
2:T:85:LEU:H	2:T:85:LEU:HD22	1.74	0.51
2:AA:188:LEU:HD23	2:AA:188:LEU:H	1.75	0.51

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:HA:86:ILE:HG13	3:HA:198:ILE:HD11	1.92	0.51
3:TA:183:GLU:HG3	3:TA:188:PHE:HB2	1.92	0.51
2:EB:188:LEU:HB3	3:LB:74:ARG:NE	2.26	0.51
3:LB:117:VAL:O	3:LB:215:PHE:HB2	2.10	0.51
2:QB:128:GLN:CG	3:RB:131:PHE:HB3	2.35	0.51
2:WB:79:SER:O	2:WB:82:VAL:HG22	2.11	0.51
2:WB:114:GLU:HB2	2:WB:117:SER:OG	2.10	0.51
3:XB:198:ILE:HG13	3:XB:221:PHE:CE1	2.45	0.51
4:AC:106:ILE:O	3:DC:252:ARG:HG2	2.11	0.51
2:CC:118:ALA:HA	2:CC:121:LEU:HG	1.92	0.51
3:DC:106:HIS:CE1	3:DC:113:THR:CG2	2.94	0.51
2:OC:121:LEU:HD13	3:VC:70:PHE:CZ	2.46	0.51
2:GD:239:LEU:HD11	2:GD:309:LEU:HD11	1.92	0.51
2:SD:76:TYR:HD1	2:YD:41:MET:CE	2.23	0.51
3:LE:180:VAL:HG23	3:LE:181:ARG:H	1.75	0.51
2:QE:278:MET:N	2:QE:278:MET:SD	2.84	0.51
2:WE:278:MET:SD	2:WE:278:MET:N	2.84	0.51
3:VF:97:PRO:O	3:VF:100:THR:OG1	2.18	0.51
2:GG:12:ILE:HD11	2:GG:55:VAL:HG11	1.93	0.51
2:GG:203:ILE:HA	2:GG:206:MET:CE	2.40	0.51
2:MG:75:GLU:HA	2:MG:78:ARG:HH21	1.75	0.51
3:TG:59:GLU:OE1	3:TG:59:GLU:N	2.43	0.51
2:T:76:TYR:CE2	2:AA:26:PHE:CE2	2.99	0.51
2:T:149:PHE:CD2	2:T:153:LEU:HD22	2.46	0.51
2:AA:153:LEU:O	2:AA:157:VAL:HG23	2.11	0.51
3:BA:202:THR:O	3:BA:202:THR:OG1	2.28	0.51
3:NA:169:TRP:CD1	3:NA:175:LEU:HD13	2.45	0.51
2:WB:278:MET:SD	2:WB:278:MET:N	2.83	0.51
2:IC:168:GLN:NE2	3:JC:130:LEU:HD11	2.21	0.51
3:VC:69:LEU:O	3:VC:73:LEU:HB2	2.11	0.51
3:BD:207:GLU:N	3:BD:207:GLU:OE1	2.44	0.51
2:MD:128:GLN:HE22	2:MD:164:PHE:HD2	1.59	0.51
3:TD:116:VAL:HG21	3:TD:158:LEU:HD11	1.93	0.51
2:EE:168:GLN:NE2	3:FE:130:LEU:HD11	2.25	0.51
2:EE:186:GLN:OE1	3:LE:71:ASN:HA	2.10	0.51
2:EE:278:MET:N	2:EE:278:MET:SD	2.83	0.51
2:KE:74:ASN:O	2:KE:78:ARG:NE	2.43	0.51
2:KE:128:GLN:HG2	3:LE:131:PHE:CB	2.34	0.51
3:LE:224:ILE:O	3:LE:228:ARG:N	2.44	0.51
2:WE:178:VAL:HG11	3:XE:149:THR:HG22	1.92	0.51
3:BG:97:PRO:O	3:BG:100:THR:OG1	2.21	0.51

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:SG:244:ASP:HA	2:SG:247:ILE:HG12	1.93	0.51
2:G:11:VAL:O	2:G:14:LEU:HD12	2.10	0.51
2:B:188:LEU:HB3	3:K:74:ARG:NE	2.26	0.51
2:T:126:HIS:CE1	3:V:127:VAL:HG13	2.45	0.51
2:T:187:ASN:N	3:BA:74:ARG:HH22	2.07	0.51
2:AA:30:SER:O	2:AA:34:VAL:HG23	2.11	0.51
3:HA:58:ASN:ND2	3:HA:59:GLU:OE2	2.43	0.51
4:KA:107:ALA:HB1	4:KA:124:ASP:O	2.10	0.51
2:EB:69:LEU:HD13	2:KB:45:ARG:O	2.11	0.51
2:CC:83:LYS:HA	2:IC:23:ALA:HB1	1.91	0.51
2:OC:278:MET:N	2:OC:278:MET:SD	2.84	0.51
3:VC:96:LEU:HD23	3:VC:97:PRO:HD2	1.92	0.51
2:GD:102:ASP:HB3	2:GD:107:ILE:HB	1.93	0.51
2:YD:63:ALA:HA	2:YD:66:PHE:CZ	2.46	0.51
2:WE:110:LEU:O	2:WE:113:MET:N	2.42	0.51
3:XE:66:ARG:HB2	3:XE:79:ILE:HD12	1.92	0.51
2:UF:302:GLU:O	2:UF:306:LYS:HG2	2.11	0.51
2:AG:108:GLU:HA	2:AG:111:ASN:HD22	1.74	0.51
2:MG:109:THR:O	2:MG:113:MET:HG3	2.11	0.51
2:SG:84:ALA:O	2:SG:87:GLU:HB3	2.10	0.51
2:G:204:ASN:OD1	2:G:233:MET:HB3	2.11	0.51
3:M:59:GLU:N	3:M:59:GLU:OE2	2.44	0.51
3:C:118:PHE:CE2	3:C:158:LEU:HD12	2.45	0.51
2:J:83:LYS:HD3	2:T:26:PHE:CB	2.41	0.51
2:T:38:SER:OG	2:T:39:THR:N	2.43	0.51
3:HA:101:ASN:HD22	3:HA:146:PHE:HE2	1.59	0.51
2:QB:310:LEU:HD12	2:QB:311:ILE:HD13	1.92	0.51
2:CC:239:LEU:HD11	2:CC:309:LEU:HD11	1.92	0.51
2:IC:78:ARG:O	2:IC:82:VAL:HG23	2.11	0.51
2:IC:278:MET:N	2:IC:278:MET:SD	2.84	0.51
3:JC:109:PRO:O	3:JC:111:ARG:NH1	2.43	0.51
3:JC:118:PHE:CE2	3:JC:158:LEU:HD12	2.46	0.51
2:UC:9:LYS:HG2	2:UC:55:VAL:HG22	1.92	0.51
2:GD:130:ILE:O	2:GD:134:LEU:HD13	2.10	0.51
3:HD:89:TYR:HE2	3:HD:191:ILE:HG13	1.75	0.51
2:YD:109:THR:O	2:YD:113:MET:HG3	2.10	0.51
2:YD:167:VAL:HG13	3:ZD:130:LEU:O	2.11	0.51
2:YD:278:MET:N	2:YD:278:MET:SD	2.83	0.51
4:AE:101:ILE:N	4:AE:104:TYR:O	2.33	0.51
2:QE:75:GLU:HA	2:QE:78:ARG:NH2	2.21	0.51
2:WE:30:SER:O	2:WE:34:VAL:HG23	2.11	0.51

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:WE:151:GLU:HB3	2:WE:155:HIS:CE1	2.45	0.51
2:CF:101:ARG:O	2:CF:104:THR:HG23	2.10	0.51
1:NF:548:ALA:HB1	2:OF:70:ASN:HD22	1.75	0.51
2:OF:278:MET:N	2:OF:278:MET:SD	2.84	0.51
3:PF:65:PHE:CE1	3:PF:161:ALA:HB2	2.46	0.51
2:GG:141:GLN:HA	2:GG:144:ASP:OD2	2.10	0.51
2:GG:245:ARG:O	2:GG:245:ARG:NH1	2.38	0.51
2:MG:278:MET:SD	2:MG:278:MET:N	2.83	0.51
2:SG:12:ILE:HD11	2:SG:55:VAL:HG11	1.92	0.51
3:TG:58:ASN:OD1	3:TG:81:VAL:HG11	2.09	0.51
2:G:143:ALA:HB1	2:SG:206:MET:HG2	1.93	0.51
3:M:96:LEU:HD23	3:M:97:PRO:HD2	1.92	0.51
3:C:79:ILE:HG22	3:C:204:PHE:HB3	1.93	0.51
2:AA:207:LYS:NZ	2:AA:209:GLN:HB3	2.26	0.51
3:BA:119:SER:HB3	3:BA:214:GLU:O	2.11	0.51
3:BA:134:ASP:OD2	3:BA:136:ARG:N	2.29	0.51
2:GA:95:GLU:HG3	2:GA:112:PHE:CE2	2.46	0.51
2:GA:168:GLN:NE2	3:HA:130:LEU:HD11	2.26	0.51
3:TA:75:ARG:HG3	3:TA:208:ILE:HD11	1.92	0.51
2:YA:186:GLN:NE2	3:FB:71:ASN:HA	2.22	0.51
2:YA:188:LEU:HD13	3:FB:74:ARG:CD	2.33	0.51
3:LB:191:ILE:HG13	3:LB:192:THR:HG23	1.92	0.51
3:LB:240:ARG:NH1	3:LB:240:ARG:HA	2.25	0.51
2:QB:126:HIS:CE1	3:RB:127:VAL:HG13	2.46	0.51
2:WB:26:PHE:CE2	2:WB:37:LEU:HD11	2.46	0.51
3:JC:190:ASN:OD1	3:PC:236:LEU:N	2.44	0.51
3:PC:90:HIS:CE1	3:PC:91:GLU:HG2	2.46	0.51
3:PC:202:THR:O	3:PC:202:THR:OG1	2.29	0.51
2:UC:239:LEU:HD11	2:UC:309:LEU:HD11	1.91	0.51
2:AD:239:LEU:HD11	2:AD:309:LEU:HD11	1.92	0.51
2:SD:278:MET:N	2:SD:278:MET:SD	2.84	0.51
2:YD:86:GLY:O	2:YD:197:ARG:NH1	2.44	0.51
2:EE:151:GLU:HA	2:EE:154:ARG:HG2	1.93	0.51
2:KE:79:SER:HA	2:KE:82:VAL:CG1	2.41	0.51
3:LE:79:ILE:HG22	3:LE:204:PHE:HB3	1.93	0.51
3:LE:134:ASP:OD2	3:LE:136:ARG:N	2.39	0.51
3:RE:101:ASN:HB2	3:RE:146:PHE:HE1	1.76	0.51
2:WE:152:ARG:HH12	2:WE:153:LEU:HB2	1.75	0.51
2:IF:80:VAL:HG12	2:OF:26:PHE:HE2	1.75	0.51
2:IF:205:LEU:HD21	2:OF:140:SER:HA	1.92	0.51
3:PF:96:LEU:HD23	3:PF:97:PRO:CD	2.40	0.51

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:UF:188:LEU:HD22	3:BG:74:ARG:HH21	1.76	0.51
2:G:168:GLN:HB3	2:SG:188:LEU:HB2	1.93	0.51
2:G:278:MET:N	2:G:278:MET:SD	2.83	0.51
3:K:117:VAL:O	3:K:215:PHE:HB2	2.11	0.51
2:AA:230:ILE:HA	2:AA:233:MET:CE	2.41	0.51
2:AA:278:MET:N	2:AA:278:MET:SD	2.83	0.51
3:BA:106:HIS:CE1	3:BA:113:THR:CG2	2.93	0.51
2:GA:69:LEU:HD13	2:MA:45:ARG:O	2.10	0.51
2:GA:108:GLU:HA	2:GA:111:ASN:ND2	2.17	0.51
2:MA:190:ARG:O	2:SA:166:GLY:HA3	2.10	0.51
2:MA:278:MET:SD	2:MA:278:MET:N	2.83	0.51
3:NA:229:GLU:OE2	3:NA:233:ASN:ND2	2.44	0.51
3:FB:183:GLU:HG3	3:FB:188:PHE:HB2	1.93	0.51
3:LB:180:VAL:HG23	3:LB:181:ARG:H	1.74	0.51
3:DC:239:SER:HA	3:DC:242:GLU:OE1	2.11	0.51
2:GD:32:ARG:NH2	2:GD:33:GLU:OE1	2.41	0.51
2:GD:310:LEU:HD12	2:GD:311:ILE:HD13	1.92	0.51
3:HD:239:SER:HA	3:HD:242:GLU:OE1	2.11	0.51
2:MD:67:ALA:O	2:MD:71:ILE:HG12	2.11	0.51
2:YD:128:GLN:HG2	3:ZD:131:PHE:CB	2.34	0.51
2:YD:133:ILE:O	2:YD:137:LEU:HG	2.11	0.51
3:ZD:224:ILE:O	3:ZD:228:ARG:N	2.44	0.51
2:CF:210:GLN:O	2:CF:214:VAL:HG12	2.09	0.51
2:IF:203:ILE:HA	2:IF:206:MET:CE	2.41	0.51
2:OF:205:LEU:CD2	2:UF:140:SER:HA	2.41	0.51
2:UF:151:GLU:HB3	2:UF:155:HIS:CE1	2.45	0.51
2:AG:8:ASP:O	2:AG:12:ILE:HG12	2.11	0.51
2:GG:278:MET:N	2:GG:278:MET:SD	2.84	0.51
3:HG:65:PHE:CE1	3:HG:161:ALA:HB2	2.46	0.51
3:HG:134:ASP:OD2	3:HG:136:ARG:N	2.38	0.51
2:SG:107:ILE:HD12	2:SG:107:ILE:N	2.26	0.51
2:SG:310:LEU:HD12	2:SG:311:ILE:HD13	1.92	0.51
2:T:167:VAL:HG13	3:V:130:LEU:O	2.10	0.50
2:T:186:GLN:C	3:BA:74:ARG:HH22	2.14	0.50
2:T:278:MET:N	2:T:278:MET:SD	2.83	0.50
2:AA:69:LEU:HD13	2:GA:45:ARG:O	2.11	0.50
2:MA:131:ALA:HA	2:MA:134:LEU:HD12	1.94	0.50
2:MA:264:LYS:HD2	2:MA:265:GLY:N	2.25	0.50
2:YA:244:ASP:HA	2:YA:247:ILE:HG12	1.94	0.50
2:EB:239:LEU:HD11	2:EB:309:LEU:HD11	1.92	0.50
2:QB:134:LEU:HD13	2:QB:146:LEU:HD21	1.93	0.50

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:QB:187:ASN:N	3:XB:74:ARG:HH22	2.08	0.50
2:WB:186:GLN:HB3	3:DC:74:ARG:NH1	2.26	0.50
2:IC:128:GLN:HG2	3:JC:131:PHE:CB	2.37	0.50
2:UC:12:ILE:HD11	2:UC:55:VAL:HG11	1.92	0.50
2:GD:133:ILE:O	2:GD:137:LEU:HG	2.11	0.50
2:GD:278:MET:N	2:GD:278:MET:SD	2.85	0.50
2:MD:152:ARG:HH12	2:MD:153:LEU:HD13	1.74	0.50
2:SD:186:GLN:NE2	3:ZD:71:ASN:HA	2.23	0.50
3:TD:51:LEU:O	3:TD:55:GLU:HG3	2.11	0.50
2:YD:67:ALA:O	2:YD:71:ILE:HG12	2.12	0.50
2:YD:110:LEU:C	2:YD:141:GLN:HE22	2.10	0.50
2:QE:128:GLN:HG2	3:RE:131:PHE:CB	2.37	0.50
2:QE:188:LEU:HD21	3:XE:74:ARG:CZ	2.42	0.50
4:YE:101:ILE:N	4:YE:104:TYR:O	2.34	0.50
2:CF:264:LYS:HD2	2:CF:265:GLY:N	2.25	0.50
3:DF:118:PHE:CE2	3:DF:158:LEU:HD12	2.46	0.50
3:DF:180:VAL:HG23	3:DF:181:ARG:H	1.76	0.50
2:IF:111:ASN:HA	2:IF:141:GLN:NE2	2.26	0.50
3:JF:180:VAL:HG23	3:JF:181:ARG:H	1.75	0.50
3:PF:118:PHE:CE2	3:PF:158:LEU:HD12	2.46	0.50
2:UF:244:ASP:HA	2:UF:247:ILE:HG12	1.94	0.50
2:MG:8:ASP:O	2:MG:12:ILE:HG12	2.10	0.50
2:MG:186:GLN:NE2	3:TG:71:ASN:HA	2.25	0.50
3:NG:221:PHE:HD2	3:NG:225:GLU:OE2	1.94	0.50
2:SG:302:GLU:O	2:SG:306:LYS:HG2	2.11	0.50
2:G:26:PHE:CD2	2:SG:83:LYS:HD3	2.46	0.50
3:C:183:GLU:HG3	3:C:188:PHE:HB2	1.93	0.50
2:J:12:ILE:HD11	2:J:55:VAL:HG11	1.93	0.50
3:V:224:ILE:O	3:V:228:ARG:N	2.44	0.50
2:AA:207:LYS:CG	2:AA:210:GLN:HG2	2.42	0.50
3:NA:115:LEU:HD21	3:NA:218:CYS:HB3	1.93	0.50
2:SA:9:LYS:HG2	2:SA:55:VAL:HG22	1.93	0.50
2:SA:111:ASN:HA	2:SA:141:GLN:OE1	2.10	0.50
2:YA:110:LEU:HD13	2:YA:137:LEU:HD21	1.92	0.50
2:EB:244:ASP:HA	2:EB:247:ILE:HG12	1.93	0.50
3:FB:214:GLU:OE1	3:FB:215:PHE:N	2.44	0.50
2:KB:84:ALA:O	2:KB:87:GLU:HB3	2.11	0.50
2:QB:239:LEU:HD11	2:QB:309:LEU:HD11	1.92	0.50
3:RB:183:GLU:HG3	3:RB:188:PHE:HB2	1.94	0.50
2:WB:302:GLU:O	2:WB:306:LYS:HG2	2.12	0.50
3:XB:116:VAL:HG21	3:XB:158:LEU:HD11	1.93	0.50

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:XB:118:PHE:CE2	3:XB:158:LEU:HD12	2.46	0.50
2:CC:245:ARG:O	2:CC:245:ARG:NH1	2.37	0.50
3:VC:94:ARG:HH11	3:VC:94:ARG:HG3	1.75	0.50
2:GD:63:ALA:HA	2:GD:66:PHE:CZ	2.47	0.50
2:GD:159:LEU:HD21	3:HD:137:PHE:CD1	2.45	0.50
3:ND:118:PHE:CE2	3:ND:158:LEU:HD12	2.47	0.50
2:EE:141:GLN:HA	2:EE:144:ASP:OD2	2.11	0.50
3:FE:247:ARG:NH1	3:FE:250:LEU:HD22	2.27	0.50
2:KE:151:GLU:HB3	2:KE:155:HIS:CE1	2.45	0.50
3:LE:190:ASN:OD1	3:RE:236:LEU:HB2	2.11	0.50
2:QE:107:ILE:HD12	2:QE:107:ILE:N	2.26	0.50
2:QE:153:LEU:O	2:QE:157:VAL:HG23	2.11	0.50
2:WE:101:ARG:O	2:WE:104:THR:HG23	2.11	0.50
2:OF:72:ASN:HD22	2:UF:41:MET:CE	2.25	0.50
2:OF:193:MET:O	2:OF:195:GLY:N	2.43	0.50
2:AG:84:ALA:O	2:AG:87:GLU:HB3	2.10	0.50
2:AG:245:ARG:O	2:AG:245:ARG:NH1	2.37	0.50
2:MG:187:ASN:N	3:TG:74:ARG:HH22	2.09	0.50
3:NG:54:LEU:HA	3:NG:57:ILE:HG22	1.94	0.50
3:K:65:PHE:CE1	3:K:161:ALA:HB2	2.46	0.50
3:K:75:ARG:HG3	3:K:208:ILE:HD11	1.93	0.50
2:T:9:LYS:HG2	2:T:55:VAL:HG22	1.93	0.50
2:GA:207:LYS:H	2:GA:210:GLN:HE21	1.60	0.50
2:YA:17:ILE:HD12	2:YA:21:ARG:HG3	1.93	0.50
2:YA:30:SER:O	2:YA:34:VAL:HG23	2.11	0.50
2:EB:9:LYS:HG2	2:EB:55:VAL:HG22	1.94	0.50
2:EB:101:ARG:O	2:EB:104:THR:HG23	2.12	0.50
3:FB:180:VAL:HG23	3:FB:181:ARG:H	1.76	0.50
3:RB:190:ASN:OD1	3:XB:236:LEU:HB2	2.10	0.50
3:XB:97:PRO:O	3:XB:100:THR:OG1	2.18	0.50
2:CC:12:ILE:HD11	2:CC:55:VAL:HG11	1.91	0.50
2:UC:151:GLU:HB3	2:UC:155:HIS:CE1	2.45	0.50
2:AD:11:VAL:O	2:AD:15:MET:HE3	2.10	0.50
2:GD:127:PRO:O	2:GD:130:ILE:HG12	2.10	0.50
2:GD:168:GLN:HE22	3:HD:130:LEU:HD11	1.76	0.50
3:ZD:119:SER:HB3	3:ZD:214:GLU:O	2.10	0.50
2:EE:239:LEU:HD11	2:EE:309:LEU:HD11	1.92	0.50
2:CF:282:ALA:HA	2:CF:285:ILE:HG12	1.92	0.50
2:OF:101:ARG:O	2:OF:104:THR:HG23	2.12	0.50
2:OF:310:LEU:HD12	2:OF:311:ILE:HD13	1.92	0.50
2:UF:87:GLU:HG2	2:UF:89:ARG:N	2.26	0.50

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:AG:67:ALA:O	2:AG:71:ILE:HG12	2.11	0.50
3:BG:72:LEU:HD23	3:BG:73:LEU:HD12	1.93	0.50
2:GG:244:ASP:HA	2:GG:247:ILE:HG12	1.93	0.50
2:G:87:GLU:HG2	2:G:89:ARG:N	2.26	0.50
3:M:102:LEU:HD22	3:M:117:VAL:HG22	1.94	0.50
3:C:237:GLU:HG2	3:C:238:ASN:H	1.76	0.50
2:J:84:ALA:O	2:J:87:GLU:HB3	2.11	0.50
2:J:101:ARG:O	2:J:104:THR:HG23	2.11	0.50
2:J:178:VAL:HG11	3:K:149:THR:HG22	1.94	0.50
2:AA:151:GLU:HA	2:AA:154:ARG:HE	1.76	0.50
2:YA:11:VAL:O	2:YA:15:MET:HE3	2.11	0.50
3:ZA:58:ASN:OD1	3:ZA:81:VAL:HG11	2.11	0.50
3:ZA:238:ASN:HB3	3:ZA:241:HIS:ND1	2.26	0.50
3:FB:202:THR:OG1	3:FB:202:THR:O	2.30	0.50
3:RB:180:VAL:HG23	3:RB:181:ARG:H	1.76	0.50
2:IC:302:GLU:O	2:IC:306:LYS:HG2	2.12	0.50
2:OC:230:ILE:HA	2:OC:233:MET:CE	2.42	0.50
2:OC:245:ARG:O	2:OC:245:ARG:NH1	2.38	0.50
2:UC:69:LEU:H	2:UC:69:LEU:HD12	1.76	0.50
3:FE:101:ASN:HD22	3:FE:146:PHE:HE2	1.59	0.50
3:FE:202:THR:O	3:FE:202:THR:OG1	2.28	0.50
2:KE:302:GLU:O	2:KE:306:LYS:HG2	2.12	0.50
3:RE:72:LEU:HD23	3:RE:73:LEU:HD12	1.93	0.50
3:JF:224:ILE:O	3:JF:228:ARG:N	2.45	0.50
2:UF:26:PHE:CE1	2:UF:34:VAL:HG22	2.46	0.50
2:AG:244:ASP:HA	2:AG:247:ILE:HG12	1.94	0.50
2:G:83:LYS:HD3	2:B:26:PHE:CD1	2.47	0.50
2:B:84:ALA:O	2:B:87:GLU:HB3	2.11	0.50
2:B:159:LEU:HD21	3:C:137:PHE:CD1	2.46	0.50
2:J:30:SER:O	2:J:34:VAL:HG23	2.10	0.50
2:J:41:MET:HG2	2:J:42:ALA:N	2.27	0.50
3:K:247:ARG:HG2	3:K:247:ARG:HH11	1.76	0.50
2:T:30:SER:O	2:T:34:VAL:HG23	2.11	0.50
2:T:151:GLU:HA	2:T:154:ARG:HG2	1.93	0.50
2:GA:63:ALA:HA	2:GA:66:PHE:CZ	2.47	0.50
2:GA:151:GLU:HA	2:GA:154:ARG:HG2	1.93	0.50
2:GA:245:ARG:O	2:GA:245:ARG:NH1	2.37	0.50
2:MA:151:GLU:HB3	2:MA:155:HIS:CE1	2.47	0.50
2:SA:278:MET:N	2:SA:278:MET:SD	2.84	0.50
2:YA:130:ILE:HD11	2:YA:157:VAL:HG22	1.93	0.50
3:ZA:79:ILE:HG22	3:ZA:204:PHE:HB3	1.92	0.50

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:EB:114:GLU:HB2	2:EB:117:SER:OG	2.11	0.50
2:KB:230:ILE:HA	2:KB:233:MET:HE2	1.93	0.50
2:QB:30:SER:OG	2:QB:33:GLU:OE1	2.20	0.50
2:WB:67:ALA:O	2:WB:71:ILE:HG12	2.11	0.50
3:DC:97:PRO:HD2	3:DC:97:PRO:O	2.11	0.50
2:IC:21:ARG:HE	2:IC:21:ARG:C	2.10	0.50
2:UC:302:GLU:O	2:UC:306:LYS:HG2	2.12	0.50
3:VC:116:VAL:HG21	3:VC:158:LEU:HD11	1.93	0.50
3:BD:202:THR:O	3:BD:202:THR:OG1	2.29	0.50
3:HD:65:PHE:CE1	3:HD:161:ALA:HB2	2.47	0.50
2:MD:9:LYS:HG2	2:MD:55:VAL:HG22	1.92	0.50
2:SD:133:ILE:O	2:SD:137:LEU:HG	2.12	0.50
3:ZD:202:THR:O	3:ZD:202:THR:OG1	2.28	0.50
2:KE:115:PRO:O	2:KE:119:ALA:HB2	2.12	0.50
2:WE:244:ASP:HA	2:WE:247:ILE:HG12	1.93	0.50
2:CF:126:HIS:ND1	2:CF:127:PRO:HD2	2.27	0.50
2:IF:76:TYR:HD2	2:OF:41:MET:CE	2.25	0.50
2:AG:128:GLN:CG	3:BG:131:PHE:HB3	2.36	0.50
2:MG:244:ASP:HA	2:MG:247:ILE:HG12	1.94	0.50
3:NG:202:THR:OG1	3:NG:202:THR:O	2.29	0.50
2:J:26:PHE:CE1	2:J:34:VAL:HG22	2.34	0.50
2:J:107:ILE:HG23	2:J:137:LEU:HD23	1.93	0.50
2:T:79:SER:HA	2:T:82:VAL:CG1	2.42	0.50
2:AA:83:LYS:O	2:GA:27:LYS:HD2	2.12	0.50
2:AA:109:THR:O	2:AA:113:MET:HG3	2.11	0.50
3:NA:180:VAL:HG23	3:NA:181:ARG:H	1.76	0.50
2:EB:11:VAL:HG11	2:EB:41:MET:HB3	1.94	0.50
2:QB:84:ALA:O	2:QB:87:GLU:HB3	2.12	0.50
2:WB:244:ASP:HA	2:WB:247:ILE:HG12	1.94	0.50
2:CC:63:ALA:HA	2:CC:66:PHE:CE2	2.46	0.50
2:IC:8:ASP:O	2:IC:12:ILE:HG12	2.12	0.50
3:JC:112:GLY:HA3	3:JC:223:MET:HG2	1.93	0.50
2:OC:186:GLN:NE2	3:VC:71:ASN:HA	2.26	0.50
3:PC:118:PHE:CE2	3:PC:158:LEU:HD12	2.46	0.50
2:UC:83:LYS:HD3	2:AD:26:PHE:HD1	1.77	0.50
2:UC:105:SER:OG	2:UC:107:ILE:HD11	2.12	0.50
2:UC:205:LEU:CD2	2:AD:140:SER:HA	2.41	0.50
2:UC:245:ARG:O	2:UC:245:ARG:NH1	2.38	0.50
2:AD:69:LEU:HD11	2:GD:47:ILE:CG2	2.36	0.50
2:GD:76:TYR:HD2	2:MD:41:MET:CE	2.24	0.50
2:GD:109:THR:O	2:GD:113:MET:HG3	2.11	0.50

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:GD:302:GLU:O	2:GD:306:LYS:HG2	2.12	0.50
3:HD:101:ASN:HB2	3:HD:146:PHE:CE2	2.47	0.50
2:MD:278:MET:SD	2:MD:278:MET:N	2.85	0.50
2:YD:130:ILE:O	2:YD:134:LEU:HG	2.11	0.50
2:EE:127:PRO:O	2:EE:130:ILE:HG12	2.12	0.50
2:EE:128:GLN:CG	3:FE:131:PHE:HB3	2.37	0.50
2:EE:244:ASP:HA	2:EE:247:ILE:HG12	1.93	0.50
2:KE:146:LEU:HD11	2:KE:157:VAL:HG11	1.93	0.50
3:DF:96:LEU:HD23	3:DF:97:PRO:CD	2.42	0.50
2:IF:9:LYS:HG2	2:IF:55:VAL:HG22	1.92	0.50
2:OF:130:ILE:O	2:OF:134:LEU:HD13	2.12	0.50
3:VF:202:THR:O	3:VF:202:THR:OG1	2.28	0.50
2:AG:26:PHE:CE1	2:AG:34:VAL:HG22	2.47	0.50
2:AG:95:GLU:HG3	2:AG:112:PHE:CE2	2.47	0.50
2:MG:230:ILE:HA	2:MG:233:MET:CE	2.41	0.50
2:B:244:ASP:HA	2:B:247:ILE:HG12	1.94	0.50
3:C:190:ASN:O	3:K:236:LEU:HD12	2.12	0.50
2:J:98:LEU:HB3	2:J:108:GLU:OE1	2.11	0.50
2:J:205:LEU:CD2	2:T:140:SER:HA	2.41	0.50
2:T:190:ARG:O	2:AA:166:GLY:HA3	2.11	0.50
2:AA:244:ASP:HA	2:AA:247:ILE:HG12	1.94	0.50
2:GA:278:MET:N	2:GA:278:MET:SD	2.84	0.50
2:MA:239:LEU:HD11	2:MA:309:LEU:HD11	1.92	0.50
3:TA:101:ASN:HB2	3:TA:146:PHE:HE1	1.77	0.50
2:YA:115:PRO:O	2:YA:119:ALA:HB2	2.10	0.50
2:QB:151:GLU:HA	2:QB:154:ARG:HG2	1.91	0.50
2:QB:198:THR:O	2:QB:202:ILE:HG12	2.11	0.50
3:RB:239:SER:HA	3:RB:242:GLU:OE1	2.11	0.50
2:WB:239:LEU:HD11	2:WB:309:LEU:HD11	1.93	0.50
3:XB:224:ILE:O	3:XB:228:ARG:N	2.45	0.50
2:CC:188:LEU:HB3	3:JC:74:ARG:NE	2.26	0.50
4:EC:101:ILE:N	4:EC:104:TYR:O	2.34	0.50
2:UC:80:VAL:HG11	2:UC:93:LEU:HD12	1.92	0.50
2:UC:90:ALA:HA	2:UC:93:LEU:HG	1.94	0.50
2:UC:107:ILE:H	2:UC:107:ILE:HD12	1.76	0.50
2:AD:8:ASP:O	2:AD:12:ILE:HG12	2.11	0.50
3:HD:134:ASP:OD1	3:HD:135:GLY:N	2.45	0.50
3:ND:202:THR:OG1	3:ND:202:THR:O	2.29	0.50
2:SD:48:SER:OG	2:SD:51:GLN:OE1	2.28	0.50
2:EE:186:GLN:HB3	3:LE:74:ARG:CZ	2.42	0.50
2:QE:76:TYR:HE2	2:WE:38:SER:HB3	1.77	0.50

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:JF:97:PRO:O	3:JF:100:THR:OG1	2.18	0.50
2:OF:244:ASP:HA	2:OF:247:ILE:HG12	1.94	0.50
3:PF:54:LEU:HB2	3:PF:231:LEU:HD23	1.93	0.50
2:AG:160:ARG:O	2:AG:164:PHE:CB	2.60	0.50
3:HG:247:ARG:HG2	3:HG:247:ARG:HH11	1.76	0.50
2:MG:310:LEU:HD12	2:MG:311:ILE:HD13	1.92	0.50
2:SG:187:ASN:O	2:SG:189:LYS:NZ	2.42	0.50
2:SG:278:MET:N	2:SG:278:MET:SD	2.84	0.50
2:G:65:GLN:HG3	2:B:46:GLN:HG2	1.94	0.50
2:SA:78:ARG:O	2:SA:82:VAL:HG12	2.11	0.50
2:SA:244:ASP:HA	2:SA:247:ILE:HG12	1.94	0.50
2:YA:230:ILE:HA	2:YA:233:MET:CE	2.41	0.50
2:YA:278:MET:N	2:YA:278:MET:SD	2.84	0.50
2:EB:141:GLN:HA	2:EB:144:ASP:OD2	2.12	0.50
2:EB:153:LEU:O	2:EB:157:VAL:HG23	2.11	0.50
2:IC:245:ARG:O	2:IC:245:ARG:NH1	2.38	0.50
3:JC:180:VAL:HG23	3:JC:181:ARG:H	1.76	0.50
3:VC:117:VAL:O	3:VC:215:PHE:HB2	2.12	0.50
2:AD:26:PHE:CZ	2:AD:34:VAL:HG22	2.47	0.50
2:AD:245:ARG:O	2:AD:245:ARG:NH1	2.37	0.50
2:GD:69:LEU:HD13	2:MD:45:ARG:O	2.12	0.50
2:MD:107:ILE:HG23	2:MD:137:LEU:HD23	1.93	0.50
3:ND:117:VAL:O	3:ND:215:PHE:HB2	2.12	0.50
4:QD:106:ILE:CB	3:TD:252:ARG:HG2	2.42	0.50
2:SD:87:GLU:HG2	2:SD:89:ARG:N	2.27	0.50
2:EE:63:ALA:HA	2:EE:66:PHE:CZ	2.46	0.50
2:CF:114:GLU:HB2	2:CF:117:SER:OG	2.12	0.50
3:JF:117:VAL:O	3:JF:215:PHE:HB2	2.12	0.50
2:UF:310:LEU:HD12	2:UF:311:ILE:HD13	1.93	0.50
2:AG:149:PHE:CD2	2:AG:153:LEU:HD22	2.47	0.50
2:MG:302:GLU:O	2:MG:306:LYS:HG2	2.12	0.50
2:SG:174:GLU:HG2	3:TG:152:ARG:HD2	1.92	0.50
3:M:65:PHE:CE1	3:M:161:ALA:HB2	2.47	0.50
3:C:229:GLU:OE1	3:C:230:LEU:HD12	2.12	0.50
2:J:15:MET:SD	2:J:15:MET:N	2.85	0.50
2:T:63:ALA:HA	2:T:66:PHE:CZ	2.47	0.50
2:T:84:ALA:O	2:T:87:GLU:HB3	2.11	0.50
3:HA:180:VAL:HG23	3:HA:181:ARG:H	1.77	0.50
2:SA:33:GLU:O	2:SA:37:LEU:HG	2.11	0.50
3:TA:96:LEU:HD23	3:TA:97:PRO:CD	2.41	0.50
2:YA:9:LYS:HG2	2:YA:55:VAL:HG22	1.94	0.50

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:EB:12:ILE:HD11	2:EB:55:VAL:HG11	1.94	0.50
2:EB:31:THR:O	2:EB:34:VAL:HB	2.12	0.50
3:FB:118:PHE:CE2	3:FB:158:LEU:HD12	2.46	0.50
2:QB:244:ASP:HA	2:QB:247:ILE:HG12	1.94	0.50
3:RB:191:ILE:HG13	3:RB:192:THR:HG23	1.94	0.50
2:WB:12:ILE:HD11	2:WB:55:VAL:HG11	1.93	0.50
3:XB:173:ASN:ND2	3:XB:174:PRO:HD3	2.27	0.50
3:XB:183:GLU:HG3	3:XB:188:PHE:HB2	1.94	0.50
2:CC:84:ALA:O	2:CC:87:GLU:HB3	2.11	0.50
2:CC:128:GLN:CG	3:DC:131:PHE:HB3	2.36	0.50
3:JC:183:GLU:HG3	3:JC:188:PHE:HB2	1.93	0.50
2:UC:244:ASP:HA	2:UC:247:ILE:HG12	1.93	0.50
3:BD:183:GLU:HG3	3:BD:188:PHE:HB2	1.92	0.50
2:MD:63:ALA:HA	2:MD:66:PHE:CZ	2.47	0.50
3:FE:238:ASN:HB3	3:FE:241:HIS:ND1	2.27	0.50
2:KE:203:ILE:HA	2:KE:206:MET:CE	2.40	0.50
3:RE:247:ARG:HG2	3:RE:247:ARG:HH11	1.76	0.50
2:IF:244:ASP:HA	2:IF:247:ILE:HG12	1.94	0.50
2:UF:45:ARG:HG3	2:UF:46:GLN:HG3	1.93	0.50
2:AG:203:ILE:HA	2:AG:206:MET:CE	2.42	0.50
3:HG:238:ASN:HB3	3:HG:241:HIS:ND1	2.27	0.50
2:G:244:ASP:HA	2:G:247:ILE:HG12	1.94	0.49
2:MA:63:ALA:HA	2:MA:66:PHE:CZ	2.46	0.49
2:MA:108:GLU:HA	2:MA:111:ASN:HD22	1.76	0.49
2:YA:119:ALA:HB1	2:YA:149:PHE:CE1	2.47	0.49
2:YA:230:ILE:HA	2:YA:233:MET:HE2	1.93	0.49
3:ZA:190:ASN:ND2	3:FB:234:PRO:O	2.45	0.49
2:EB:278:MET:N	2:EB:278:MET:SD	2.84	0.49
2:QB:122:ILE:HB	2:QB:130:ILE:HG22	1.93	0.49
2:QB:186:GLN:C	3:XB:74:ARG:HH22	2.16	0.49
2:QB:278:MET:N	2:QB:278:MET:SD	2.84	0.49
2:CC:174:GLU:HG2	3:DC:152:ARG:HG2	1.94	0.49
2:IC:174:GLU:HG2	3:JC:152:ARG:HG2	1.94	0.49
2:OC:244:ASP:HA	2:OC:247:ILE:HG12	1.94	0.49
2:GD:95:GLU:HG3	2:GD:112:PHE:CE2	2.47	0.49
3:HD:183:GLU:HG3	3:HD:188:PHE:HB2	1.93	0.49
2:MD:245:ARG:O	2:MD:245:ARG:NH1	2.38	0.49
2:SD:63:ALA:HA	2:SD:66:PHE:CZ	2.47	0.49
2:SD:130:ILE:O	2:SD:134:LEU:HD13	2.12	0.49
3:ZD:190:ASN:O	3:FE:236:LEU:HD12	2.12	0.49
2:EE:101:ARG:O	2:EE:104:THR:HG23	2.12	0.49

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:QE:221:PHE:CE2	2:WE:159:LEU:HD21	2.46	0.49
3:JF:65:PHE:CE1	3:JF:161:ALA:HB2	2.47	0.49
2:OF:30:SER:O	2:OF:34:VAL:HG23	2.12	0.49
3:BG:190:ASN:ND2	3:HG:234:PRO:O	2.45	0.49
2:GG:302:GLU:O	2:GG:306:LYS:HG2	2.12	0.49
2:SG:8:ASP:O	2:SG:12:ILE:HG12	2.11	0.49
2:G:75:GLU:O	2:G:79:SER:HB3	2.11	0.49
3:M:229:GLU:OE1	3:M:230:LEU:HD12	2.12	0.49
2:J:83:LYS:HD3	2:T:26:PHE:HB3	1.94	0.49
2:J:167:VAL:HG13	3:K:130:LEU:O	2.11	0.49
4:O:65:THR:O	4:R:99:ILE:HA	2.12	0.49
2:T:110:LEU:HD13	2:T:137:LEU:HD21	1.93	0.49
2:AA:174:GLU:HG2	3:BA:152:ARG:HG2	1.94	0.49
2:AA:230:ILE:HA	2:AA:233:MET:HE2	1.94	0.49
2:GA:121:LEU:HD13	3:NA:70:PHE:CZ	2.47	0.49
2:GA:152:ARG:NH2	2:GA:153:LEU:HB2	2.26	0.49
2:SA:87:GLU:HG2	2:SA:89:ARG:N	2.27	0.49
3:TA:100:THR:O	3:TA:102:LEU:HD13	2.12	0.49
2:YA:302:GLU:O	2:YA:306:LYS:HG2	2.12	0.49
2:EB:149:PHE:CD2	2:EB:153:LEU:HD22	2.47	0.49
2:WB:230:ILE:HA	2:WB:233:MET:CE	2.42	0.49
3:XB:221:PHE:HD2	3:XB:225:GLU:OE1	1.96	0.49
3:DC:102:LEU:HD22	3:DC:117:VAL:HG22	1.94	0.49
2:OC:12:ILE:HD11	2:OC:55:VAL:HG11	1.93	0.49
3:PC:239:SER:O	3:PC:242:GLU:HG2	2.12	0.49
2:AD:110:LEU:O	2:AD:113:MET:N	2.43	0.49
2:AD:133:ILE:O	2:AD:137:LEU:HG	2.12	0.49
3:HD:69:LEU:O	3:HD:73:LEU:HB2	2.12	0.49
2:MD:204:ASN:OD1	2:MD:233:MET:HB3	2.12	0.49
2:EE:31:THR:O	2:EE:34:VAL:HB	2.12	0.49
2:KE:230:ILE:HA	2:KE:233:MET:CE	2.43	0.49
2:QE:133:ILE:O	2:QE:137:LEU:HG	2.12	0.49
2:IF:63:ALA:HA	2:IF:66:PHE:CZ	2.47	0.49
2:OF:63:ALA:HA	2:OF:66:PHE:CZ	2.46	0.49
2:OF:133:ILE:O	2:OF:137:LEU:HG	2.12	0.49
3:VF:116:VAL:HG21	3:VF:158:LEU:HD11	1.92	0.49
2:AG:230:ILE:HA	2:AG:233:MET:HE2	1.94	0.49
2:G:263:LEU:HD11	2:G:274:PHE:HD2	1.78	0.49
3:M:183:GLU:HG3	3:M:188:PHE:HB2	1.93	0.49
2:B:278:MET:SD	2:B:278:MET:N	2.84	0.49
4:H:106:ILE:O	3:K:252:ARG:HG2	2.12	0.49

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:K:202:THR:OG1	3:K:202:THR:O	2.29	0.49
2:T:188:LEU:HB3	3:BA:74:ARG:NE	2.27	0.49
2:T:201:GLU:HG3	2:AA:139:ARG:HH21	1.77	0.49
2:AA:9:LYS:HG2	2:AA:55:VAL:HG22	1.93	0.49
2:AA:151:GLU:HA	2:AA:154:ARG:HG2	1.94	0.49
2:GA:75:GLU:HA	2:GA:78:ARG:NH2	2.22	0.49
2:GA:101:ARG:O	2:GA:104:THR:HG23	2.12	0.49
2:MA:244:ASP:HA	2:MA:247:ILE:HG12	1.94	0.49
2:MA:302:GLU:O	2:MA:306:LYS:HG2	2.12	0.49
3:NA:58:ASN:OD1	3:NA:81:VAL:HG11	2.12	0.49
3:TA:58:ASN:OD1	3:TA:81:VAL:HG11	2.13	0.49
3:TA:79:ILE:HG22	3:TA:204:PHE:HB3	1.94	0.49
2:YA:107:ILE:H	2:YA:107:ILE:HD12	1.77	0.49
2:QB:21:ARG:HH21	2:QB:24:GLU:HB2	1.76	0.49
2:WB:206:MET:SD	2:WB:210:GLN:HB3	2.52	0.49
2:IC:239:LEU:HD11	2:IC:309:LEU:HD11	1.92	0.49
2:UC:127:PRO:O	2:UC:130:ILE:HG12	2.12	0.49
3:VC:190:ASN:OD1	3:BD:236:LEU:HB2	2.12	0.49
2:AD:186:GLN:C	3:HD:74:ARG:HH22	2.15	0.49
3:BD:238:ASN:HB3	3:BD:241:HIS:ND1	2.27	0.49
3:TD:183:GLU:HG3	3:TD:188:PHE:HB2	1.94	0.49
2:KE:33:GLU:O	2:KE:37:LEU:HG	2.12	0.49
2:KE:69:LEU:HD11	2:QE:45:ARG:O	2.13	0.49
2:KE:152:ARG:HH12	2:KE:153:LEU:HB2	1.76	0.49
3:LE:221:PHE:HD2	3:LE:225:GLU:OE2	1.95	0.49
3:RE:79:ILE:HG22	3:RE:204:PHE:HB3	1.93	0.49
2:WE:302:GLU:O	2:WE:306:LYS:HG2	2.13	0.49
2:CF:8:ASP:O	2:CF:12:ILE:HG12	2.12	0.49
2:CF:38:SER:OG	2:CF:39:THR:N	2.46	0.49
2:CF:244:ASP:HA	2:CF:247:ILE:HG12	1.94	0.49
2:IF:210:GLN:O	2:IF:214:VAL:HG12	2.11	0.49
2:OF:87:GLU:HG2	2:OF:89:ARG:N	2.26	0.49
3:PF:116:VAL:HG21	3:PF:158:LEU:HD11	1.95	0.49
2:UF:202:ILE:CD1	2:AG:139:ARG:HH12	2.10	0.49
3:BG:102:LEU:HD22	3:BG:117:VAL:HG22	1.94	0.49
2:MG:63:ALA:HA	2:MG:66:PHE:CZ	2.46	0.49
2:MG:89:ARG:HG3	2:MG:90:ALA:N	2.27	0.49
2:MG:167:VAL:HG13	3:NG:130:LEU:O	2.12	0.49
2:G:107:ILE:O	2:G:111:ASN:ND2	2.45	0.49
2:B:187:ASN:N	3:K:74:ARG:HH22	2.10	0.49
2:AA:119:ALA:HB1	2:AA:149:PHE:CE1	2.47	0.49

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:GA:146:LEU:HD21	2:GA:157:VAL:HG11	1.94	0.49
2:MA:107:ILE:HD12	2:MA:107:ILE:H	1.78	0.49
2:YA:95:GLU:HG3	2:YA:112:PHE:CE2	2.47	0.49
2:KB:278:MET:N	2:KB:278:MET:SD	2.84	0.49
2:WB:75:GLU:HA	2:WB:78:ARG:NH2	2.24	0.49
2:CC:149:PHE:CD2	2:CC:153:LEU:HD22	2.47	0.49
2:CC:244:ASP:HA	2:CC:247:ILE:HG12	1.94	0.49
3:DC:75:ARG:HG3	3:DC:208:ILE:HD11	1.95	0.49
2:IC:244:ASP:HA	2:IC:247:ILE:HG12	1.94	0.49
2:GD:244:ASP:HA	2:GD:247:ILE:HG12	1.94	0.49
3:HD:112:GLY:HA3	3:HD:223:MET:HG2	1.92	0.49
2:MD:69:LEU:HD13	2:SD:45:ARG:O	2.12	0.49
2:SD:152:ARG:NH2	2:SD:153:LEU:HB2	2.27	0.49
3:LE:169:TRP:CD1	3:LE:175:LEU:HD13	2.46	0.49
3:XE:180:VAL:HG23	3:XE:181:ARG:H	1.76	0.49
3:XE:238:ASN:HB3	3:XE:241:HIS:ND1	2.27	0.49
2:CF:128:GLN:CG	3:DF:131:PHE:HB3	2.31	0.49
2:CF:230:ILE:HA	2:CF:233:MET:CE	2.42	0.49
2:CF:230:ILE:HA	2:CF:233:MET:HE2	1.93	0.49
2:IF:127:PRO:HA	2:IF:130:ILE:HD12	1.94	0.49
2:AG:131:ALA:HA	2:AG:134:LEU:HD12	1.95	0.49
2:AG:174:GLU:HG2	3:BG:152:ARG:HG2	1.95	0.49
2:SG:11:VAL:O	2:SG:15:MET:HE3	2.11	0.49
2:SG:138:LYS:HB2	2:SG:141:GLN:HG2	1.93	0.49
3:TG:69:LEU:O	3:TG:73:LEU:HB2	2.12	0.49
3:TG:97:PRO:O	3:TG:100:THR:OG1	2.19	0.49
3:M:54:LEU:HD13	3:M:231:LEU:HD21	1.94	0.49
2:B:193:MET:O	2:J:164:PHE:HD1	1.95	0.49
2:MA:263:LEU:HD11	2:MA:274:PHE:HD2	1.78	0.49
2:EB:128:GLN:HG2	3:FB:131:PHE:CB	2.35	0.49
2:KB:69:LEU:HD13	2:QB:44:VAL:HG12	1.93	0.49
2:KB:76:TYR:CE1	2:QB:26:PHE:CE2	3.01	0.49
2:KB:167:VAL:HG13	3:LB:130:LEU:O	2.12	0.49
2:QB:8:ASP:O	2:QB:12:ILE:HG12	2.11	0.49
3:RB:118:PHE:CE2	3:RB:158:LEU:HD12	2.48	0.49
2:WB:263:LEU:HD11	2:WB:274:PHE:HD2	1.78	0.49
2:CC:160:ARG:O	2:CC:164:PHE:CB	2.60	0.49
2:AD:83:LYS:O	2:GD:27:LYS:HE3	2.12	0.49
2:AD:95:GLU:O	2:AD:99:GLU:HB2	2.13	0.49
2:SD:71:ILE:O	2:SD:74:ASN:ND2	2.45	0.49
3:ZD:190:ASN:O	3:ZD:190:ASN:ND2	2.46	0.49

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:KE:183:LEU:O	2:KE:186:GLN:N	2.45	0.49
2:IF:15:MET:N	2:IF:15:MET:SD	2.84	0.49
3:PF:180:VAL:HG23	3:PF:181:ARG:H	1.76	0.49
3:VF:72:LEU:HD23	3:VF:73:LEU:HD12	1.93	0.49
2:GG:8:ASP:O	2:GG:12:ILE:HG12	2.12	0.49
2:GG:84:ALA:O	2:GG:87:GLU:HB3	2.12	0.49
3:C:106:HIS:HE1	3:C:113:THR:CG2	2.26	0.49
3:C:180:VAL:HG23	3:C:181:ARG:H	1.77	0.49
4:H:105:LEU:O	3:K:252:ARG:HD3	2.13	0.49
2:J:21:ARG:HE	2:J:21:ARG:C	2.10	0.49
2:J:63:ALA:HA	2:J:66:PHE:CZ	2.47	0.49
2:J:109:THR:O	2:J:113:MET:HG3	2.12	0.49
2:GA:84:ALA:O	2:GA:87:GLU:HB3	2.12	0.49
3:TA:190:ASN:O	3:ZA:236:LEU:HD12	2.13	0.49
3:TA:202:THR:OG1	3:TA:202:THR:O	2.31	0.49
1:XA:543:ASP:HA	2:YA:21:ARG:HH11	1.76	0.49
2:EB:187:ASN:N	3:LB:74:ARG:HH22	2.11	0.49
3:FB:79:ILE:HG22	3:FB:204:PHE:HB3	1.94	0.49
3:FB:190:ASN:O	3:LB:236:LEU:HD12	2.13	0.49
2:KB:80:VAL:CA	2:KB:83:LYS:HD3	2.32	0.49
2:KB:244:ASP:HA	2:KB:247:ILE:HG12	1.94	0.49
4:OB:107:ALA:HB3	3:RB:252:ARG:HE	1.76	0.49
2:QB:167:VAL:HG13	3:RB:130:LEU:O	2.12	0.49
4:UB:107:ALA:HB1	4:UB:124:ASP:O	2.12	0.49
2:WB:26:PHE:CZ	2:WB:34:VAL:HG22	2.47	0.49
2:WB:78:ARG:O	2:WB:82:VAL:HG13	2.12	0.49
2:WB:81:LEU:O	2:WB:85:LEU:HD13	2.12	0.49
2:WB:109:THR:O	2:WB:113:MET:HG3	2.12	0.49
2:CC:263:LEU:HD11	2:CC:274:PHE:HD2	1.78	0.49
2:UC:107:ILE:HD12	2:UC:107:ILE:N	2.27	0.49
2:AD:263:LEU:HD11	2:AD:274:PHE:HD2	1.78	0.49
2:MD:244:ASP:OD1	2:MD:276:ARG:NH2	2.46	0.49
2:SD:195:GLY:HA2	2:YD:161:ILE:O	2.12	0.49
2:SD:219:ARG:NH2	2:SD:227:GLN:OE1	2.36	0.49
3:TD:106:HIS:CD2	3:TD:113:THR:HG22	2.47	0.49
2:EE:69:LEU:HD13	2:KE:45:ARG:O	2.12	0.49
3:LE:102:LEU:HD22	3:LE:117:VAL:HG22	1.93	0.49
3:LE:185:GLN:HG3	3:LE:186:VAL:H	1.78	0.49
2:QE:183:LEU:O	2:QE:186:GLN:N	2.46	0.49
2:IF:10:SER:O	2:IF:14:LEU:HD22	2.13	0.49
2:UF:79:SER:HA	2:UF:82:VAL:HG12	1.94	0.49

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:UF:87:GLU:HG2	2:UF:89:ARG:H	1.76	0.49
2:GG:167:VAL:HG13	3:HG:130:LEU:O	2.11	0.49
2:SG:111:ASN:HA	2:SG:141:GLN:CD	2.33	0.49
2:SG:206:MET:HB3	2:SG:210:GLN:HE21	1.76	0.49
2:G:30:SER:O	2:G:34:VAL:HG23	2.12	0.49
3:C:134:ASP:OD1	3:C:135:GLY:N	2.45	0.49
2:J:69:LEU:HD13	2:T:45:ARG:O	2.12	0.49
2:J:244:ASP:HA	2:J:247:ILE:HG12	1.95	0.49
3:K:224:ILE:O	3:K:228:ARG:N	2.46	0.49
3:V:106:HIS:CE1	3:V:113:THR:HG22	2.47	0.49
2:AA:41:MET:HG2	2:AA:42:ALA:N	2.27	0.49
2:AA:167:VAL:HG13	3:BA:130:LEU:O	2.13	0.49
2:AA:245:ARG:O	2:AA:245:ARG:NH1	2.37	0.49
2:AA:263:LEU:HD11	2:AA:274:PHE:HD2	1.77	0.49
3:ZA:180:VAL:HG23	3:ZA:181:ARG:H	1.76	0.49
2:EB:8:ASP:O	2:EB:12:ILE:HG12	2.12	0.49
2:EB:302:GLU:O	2:EB:306:LYS:HG2	2.12	0.49
2:KB:12:ILE:HD11	2:KB:55:VAL:HG11	1.94	0.49
2:KB:201:GLU:HB3	2:QB:139:ARG:HH12	1.78	0.49
2:KB:207:LYS:HZ1	2:KB:209:GLN:HB3	1.77	0.49
3:LB:65:PHE:CE1	3:LB:161:ALA:HB2	2.48	0.49
2:WB:245:ARG:O	2:WB:245:ARG:NH1	2.37	0.49
3:XB:101:ASN:HB2	3:XB:146:PHE:CE2	2.47	0.49
2:OC:8:ASP:O	2:OC:12:ILE:HG12	2.13	0.49
2:UC:209:GLN:O	2:UC:212:GLU:HG3	2.13	0.49
3:BD:59:GLU:N	3:BD:59:GLU:OE2	2.45	0.49
2:GD:10:SER:O	2:GD:14:LEU:HD22	2.13	0.49
2:MD:84:ALA:O	2:MD:87:GLU:HB3	2.12	0.49
2:EE:219:ARG:NH2	2:EE:227:GLN:OE1	2.32	0.49
3:FE:224:ILE:O	3:FE:228:ARG:N	2.46	0.49
2:QE:10:SER:O	2:QE:14:LEU:HD22	2.11	0.49
2:QE:141:GLN:HA	2:QE:144:ASP:OD2	2.11	0.49
2:WE:168:GLN:NE2	3:XE:130:LEU:HD11	2.26	0.49
2:CF:214:VAL:O	2:CF:218:VAL:HG23	2.13	0.49
3:DF:51:LEU:O	3:DF:55:GLU:HG3	2.13	0.49
3:JF:202:THR:OG1	3:JF:217:ILE:HB	2.12	0.49
2:OF:26:PHE:CZ	2:OF:34:VAL:HG22	2.48	0.49
2:OF:158:MET:SD	2:OF:159:LEU:HD23	2.53	0.49
2:MG:107:ILE:H	2:MG:107:ILE:HD12	1.78	0.49
3:TG:118:PHE:CE2	3:TG:158:LEU:HD12	2.48	0.49
2:G:127:PRO:O	2:G:130:ILE:HG12	2.12	0.49

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:V:111:ARG:NH2	3:V:222:SER:O	2.45	0.49
3:V:119:SER:HB3	3:V:214:GLU:O	2.11	0.49
2:GA:8:ASP:O	2:GA:12:ILE:HG12	2.11	0.49
2:GA:178:VAL:HG11	3:HA:149:THR:HG22	1.95	0.49
2:MA:128:GLN:HG2	3:NA:131:PHE:CB	2.33	0.49
2:QB:74:ASN:OD1	2:QB:78:ARG:NH2	2.46	0.49
3:XB:214:GLU:OE1	3:XB:215:PHE:N	2.46	0.49
3:JC:238:ASN:HB3	3:JC:241:HIS:ND1	2.28	0.49
2:OC:302:GLU:O	2:OC:306:LYS:HG2	2.12	0.49
2:UC:34:VAL:O	2:UC:37:LEU:HD12	2.12	0.49
2:AD:159:LEU:HD21	3:BD:137:PHE:CD1	2.48	0.49
2:MD:8:ASP:O	2:MD:12:ILE:HG12	2.13	0.49
2:YD:84:ALA:O	2:YD:87:GLU:HB3	2.13	0.49
2:YD:244:ASP:HA	2:YD:247:ILE:HG12	1.94	0.49
2:EE:107:ILE:HG23	2:EE:137:LEU:HD23	1.94	0.49
3:FE:65:PHE:CE1	3:FE:161:ALA:HB2	2.48	0.49
2:QE:205:LEU:HD13	2:WE:139:ARG:HB3	1.93	0.49
3:XE:134:ASP:OD1	3:XE:136:ARG:N	2.29	0.49
3:XE:202:THR:OG1	3:XE:202:THR:O	2.30	0.49
2:CF:110:LEU:HD13	2:CF:137:LEU:HD21	1.94	0.49
2:CF:182:LEU:O	2:CF:186:GLN:HB2	2.12	0.49
2:CF:302:GLU:O	2:CF:306:LYS:HG2	2.12	0.49
2:IF:168:GLN:HG2	2:IF:170:ALA:H	1.77	0.49
3:PF:183:GLU:HG3	3:PF:188:PHE:HB2	1.93	0.49
2:UF:151:GLU:HA	2:UF:154:ARG:HE	1.77	0.49
2:AG:97:ILE:O	2:AG:101:ARG:NH2	2.30	0.49
2:GG:205:LEU:CD2	2:MG:140:SER:HA	2.42	0.49
2:MG:48:SER:OG	2:MG:51:GLN:OE1	2.29	0.49
3:NG:285:PRO:HA	4:OG:119:GLY:HA2	1.95	0.49
2:SG:155:HIS:O	2:SG:159:LEU:HD22	2.13	0.49
2:G:133:ILE:O	2:G:137:LEU:HG	2.13	0.49
2:G:186:GLN:C	3:C:74:ARG:HH22	2.15	0.49
2:J:278:MET:SD	2:J:278:MET:N	2.85	0.49
2:T:107:ILE:H	2:T:107:ILE:HD12	1.78	0.49
2:T:109:THR:O	2:T:113:MET:HG3	2.13	0.49
2:AA:84:ALA:O	2:AA:87:GLU:HB3	2.12	0.49
2:AA:261:ILE:O	2:AA:264:LYS:HG3	2.13	0.49
3:NA:202:THR:O	3:NA:202:THR:OG1	2.31	0.49
2:SA:261:ILE:O	2:SA:264:LYS:HG3	2.13	0.49
2:YA:63:ALA:HA	2:YA:66:PHE:CZ	2.48	0.49
2:EB:117:SER:O	2:EB:121:LEU:HG	2.13	0.49

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:KB:186:GLN:NE2	3:RB:71:ASN:HA	2.25	0.49
2:QB:67:ALA:O	2:QB:71:ILE:HG12	2.13	0.49
3:RB:65:PHE:CE1	3:RB:161:ALA:HB2	2.48	0.49
3:XB:79:ILE:HG22	3:XB:204:PHE:HB3	1.95	0.49
2:CC:109:THR:O	2:CC:113:MET:HG3	2.13	0.49
2:IC:85:LEU:HD22	2:IC:85:LEU:H	1.77	0.49
2:OC:107:ILE:H	2:OC:107:ILE:HD12	1.77	0.49
2:GD:245:ARG:O	2:GD:245:ARG:NH1	2.38	0.49
4:ID:101:ILE:N	4:ID:104:TYR:O	2.34	0.49
2:MD:101:ARG:O	2:MD:104:THR:HG23	2.13	0.49
2:SD:9:LYS:HG2	2:SD:55:VAL:HG22	1.94	0.49
2:SD:65:GLN:NE2	2:YD:47:ILE:O	2.45	0.49
2:SD:107:ILE:HD12	2:SD:107:ILE:N	2.28	0.49
2:SD:118:ALA:HA	2:SD:121:LEU:CD1	2.43	0.49
2:YD:206:MET:HB2	2:EE:143:ALA:HB1	1.93	0.49
2:EE:65:GLN:HG3	2:KE:46:GLN:HG2	1.95	0.49
2:CF:10:SER:O	2:CF:14:LEU:HD22	2.13	0.49
2:IF:186:GLN:NE2	3:PF:71:ASN:HA	2.16	0.49
3:JF:190:ASN:ND2	3:PF:234:PRO:O	2.45	0.49
2:OF:130:ILE:HD11	2:OF:157:VAL:HG22	1.95	0.49
2:UF:245:ARG:O	2:UF:245:ARG:NH1	2.37	0.49
2:SG:152:ARG:HH22	2:SG:153:LEU:HD12	1.77	0.49
3:M:79:ILE:HG22	3:M:204:PHE:HB3	1.94	0.49
3:M:118:PHE:CE2	3:M:158:LEU:HD12	2.48	0.49
2:T:128:GLN:CG	3:V:131:PHE:HB3	2.35	0.49
2:T:244:ASP:HA	2:T:247:ILE:HG12	1.95	0.49
2:AA:207:LYS:HZ3	2:AA:209:GLN:HB3	1.77	0.49
2:MA:15:MET:SD	2:MA:15:MET:N	2.86	0.49
2:SA:84:ALA:O	2:SA:87:GLU:HB3	2.13	0.49
3:FB:190:ASN:ND2	3:LB:234:PRO:O	2.45	0.49
3:LB:221:PHE:HD2	3:LB:225:GLU:OE1	1.95	0.49
2:QB:21:ARG:HE	2:QB:21:ARG:C	2.11	0.49
3:XB:69:LEU:O	3:XB:73:LEU:HB2	2.13	0.49
2:CC:167:VAL:HG13	3:DC:130:LEU:O	2.13	0.49
1:HC:543:ASP:HA	2:IC:21:ARG:HH11	1.77	0.49
3:JC:75:ARG:HG3	3:JC:208:ILE:HD11	1.94	0.49
2:UC:160:ARG:O	2:UC:164:PHE:CB	2.58	0.49
3:VC:118:PHE:CE2	3:VC:158:LEU:HD12	2.48	0.49
2:AD:127:PRO:O	2:AD:130:ILE:HG12	2.11	0.49
3:ND:58:ASN:OD1	3:ND:81:VAL:HG11	2.12	0.49
3:ZD:117:VAL:O	3:ZD:215:PHE:HB2	2.13	0.49

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:EE:100:THR:OG1	2:EE:101:ARG:NH2	2.45	0.49
2:KE:107:ILE:HD12	2:KE:107:ILE:N	2.28	0.49
3:LE:238:ASN:HB3	3:LE:241:HIS:ND1	2.28	0.49
3:RE:202:THR:O	3:RE:202:THR:OG1	2.29	0.49
3:RE:238:ASN:HB3	3:RE:241:HIS:ND1	2.27	0.49
2:WE:74:ASN:O	2:WE:78:ARG:NE	2.46	0.49
2:WE:193:MET:O	2:WE:195:GLY:N	2.46	0.49
3:DF:65:PHE:CE1	3:DF:161:ALA:HB2	2.48	0.49
3:DF:185:GLN:HG3	3:DF:186:VAL:H	1.78	0.49
2:IF:263:LEU:HD11	2:IF:274:PHE:HD2	1.77	0.49
2:OF:142:ALA:O	2:OF:146:LEU:HD22	2.13	0.49
2:UF:83:LYS:HA	2:AG:23:ALA:HB1	1.94	0.49
2:UF:95:GLU:O	2:UF:99:GLU:HB2	2.12	0.49
3:VF:97:PRO:O	3:VF:97:PRO:HD2	2.13	0.49
3:VF:214:GLU:OE1	3:VF:215:PHE:N	2.46	0.49
3:VF:239:SER:O	3:VF:242:GLU:HG2	2.13	0.49
3:BG:58:ASN:OD1	3:BG:81:VAL:HG11	2.12	0.49
2:G:110:LEU:O	2:G:113:MET:N	2.43	0.48
2:B:100:THR:OG1	2:B:101:ARG:NH2	2.46	0.48
3:C:214:GLU:OE1	3:C:215:PHE:N	2.46	0.48
2:T:8:ASP:O	2:T:12:ILE:HG12	2.12	0.48
2:T:152:ARG:NH2	2:T:153:LEU:HB2	2.27	0.48
3:V:123:VAL:HG21	3:V:146:PHE:CZ	2.48	0.48
3:BA:102:LEU:HD22	3:BA:117:VAL:HG22	1.95	0.48
2:SA:80:VAL:HA	2:YA:26:PHE:HE2	1.76	0.48
2:SA:202:ILE:HD13	2:YA:139:ARG:HH11	1.78	0.48
2:YA:100:THR:OG1	2:YA:101:ARG:NH2	2.46	0.48
4:IB:106:ILE:O	3:LB:252:ARG:HG2	2.12	0.48
3:PC:294:ALA:O	3:PC:301:VAL:N	2.46	0.48
2:UC:133:ILE:O	2:UC:137:LEU:HG	2.13	0.48
3:BD:239:SER:O	3:BD:242:GLU:HG2	2.13	0.48
2:SD:76:TYR:OH	2:YD:34:VAL:HG13	2.12	0.48
2:SD:96:ASP:OD2	2:YD:35:GLN:NE2	2.45	0.48
2:SD:117:SER:O	2:SD:121:LEU:HG	2.13	0.48
3:TD:123:VAL:HG21	3:TD:146:PHE:CZ	2.48	0.48
2:EE:87:GLU:HG2	2:EE:89:ARG:N	2.28	0.48
2:KE:244:ASP:HA	2:KE:247:ILE:HG12	1.94	0.48
2:QE:244:ASP:HA	2:QE:247:ILE:HG12	1.94	0.48
3:DF:202:THR:OG1	3:DF:202:THR:O	2.30	0.48
2:OF:152:ARG:NH2	2:OF:153:LEU:HB2	2.28	0.48
2:UF:210:GLN:O	2:UF:214:VAL:HG12	2.13	0.48

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:AG:167:VAL:HG13	3:BG:130:LEU:O	2.13	0.48
2:SG:167:VAL:HG13	3:TG:130:LEU:O	2.12	0.48
2:G:159:LEU:HD21	3:M:137:PHE:CD1	2.48	0.48
2:B:107:ILE:HD12	2:B:107:ILE:N	2.28	0.48
3:C:229:GLU:OE1	3:C:230:LEU:N	2.46	0.48
3:K:100:THR:HG23	3:K:119:SER:HA	1.95	0.48
2:T:230:ILE:HA	2:T:233:MET:CE	2.43	0.48
3:NA:180:VAL:HG23	3:NA:181:ARG:N	2.28	0.48
2:SA:74:ASN:O	2:SA:78:ARG:NE	2.46	0.48
3:TA:191:ILE:HG13	3:TA:192:THR:HG23	1.94	0.48
2:KB:76:TYR:HE2	2:QB:38:SER:HA	1.79	0.48
2:QB:65:GLN:HG3	2:WB:46:GLN:HG2	1.95	0.48
2:WB:84:ALA:O	2:WB:87:GLU:HB3	2.14	0.48
2:CC:128:GLN:HG2	3:DC:131:PHE:CB	2.35	0.48
2:OC:74:ASN:O	2:OC:78:ARG:NE	2.46	0.48
2:OC:96:ASP:OD2	2:UC:35:GLN:NE2	2.45	0.48
2:UC:168:GLN:OE1	2:UC:168:GLN:N	2.41	0.48
2:AD:76:TYR:OH	2:GD:34:VAL:O	2.29	0.48
2:AD:244:ASP:HA	2:AD:247:ILE:HG12	1.94	0.48
3:HD:79:ILE:HG22	3:HD:204:PHE:HB3	1.94	0.48
3:ND:180:VAL:HG23	3:ND:181:ARG:N	2.28	0.48
2:SD:207:LYS:NZ	2:SD:210:GLN:HB2	2.27	0.48
3:FE:185:GLN:HG2	3:FE:187:LYS:HZ2	1.78	0.48
2:KE:298:LEU:HB2	2:KE:302:GLU:OE2	2.13	0.48
2:WE:48:SER:OG	2:WE:51:GLN:OE1	2.27	0.48
2:MG:151:GLU:HA	2:MG:154:ARG:HG2	1.95	0.48
2:SG:178:VAL:HG11	3:TG:149:THR:HG22	1.96	0.48
2:G:139:ARG:NH2	2:SG:202:ILE:HD13	2.27	0.48
2:B:168:GLN:HE22	3:C:130:LEU:CD1	2.23	0.48
3:C:117:VAL:O	3:C:215:PHE:HB2	2.13	0.48
2:AA:37:LEU:HA	2:AA:40:ALA:HB3	1.95	0.48
2:SA:8:ASP:O	2:SA:12:ILE:HG12	2.13	0.48
2:SA:172:LEU:HD12	2:SA:172:LEU:HA	1.70	0.48
2:KB:83:LYS:HA	2:QB:23:ALA:HB1	1.95	0.48
2:QB:152:ARG:NH2	2:QB:153:LEU:HB2	2.28	0.48
2:QB:188:LEU:HB3	3:XB:74:ARG:NE	2.28	0.48
2:CC:203:ILE:HA	2:CC:206:MET:HE2	1.95	0.48
3:VC:86:ILE:HG13	3:VC:198:ILE:HD11	1.95	0.48
2:YD:33:GLU:O	2:YD:37:LEU:HG	2.13	0.48
2:YD:174:GLU:OE2	3:ZD:152:ARG:HG2	2.13	0.48
2:EE:8:ASP:O	2:EE:12:ILE:HG12	2.13	0.48

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:KE:127:PRO:HA	2:KE:130:ILE:HG12	1.96	0.48
3:JF:106:HIS:CE1	3:JF:113:THR:CG2	2.94	0.48
3:PF:247:ARG:HG2	3:PF:247:ARG:HH11	1.77	0.48
2:UF:263:LEU:HD11	2:UF:274:PHE:HD2	1.78	0.48
3:VF:118:PHE:CE2	3:VF:158:LEU:HD12	2.48	0.48
2:MG:263:LEU:HD11	2:MG:274:PHE:HD2	1.78	0.48
2:G:27:LYS:HE3	2:SG:83:LYS:O	2.13	0.48
2:G:174:GLU:HG2	3:M:152:ARG:HG2	1.95	0.48
2:B:95:GLU:HG3	2:B:112:PHE:CE2	2.48	0.48
3:K:118:PHE:CE2	3:K:158:LEU:HD12	2.49	0.48
3:V:69:LEU:O	3:V:73:LEU:HB2	2.14	0.48
2:AA:110:LEU:O	2:AA:113:MET:N	2.45	0.48
2:AA:126:HIS:ND1	2:AA:127:PRO:HD2	2.29	0.48
2:SA:31:THR:O	2:SA:34:VAL:HB	2.13	0.48
2:YA:167:VAL:HG13	3:ZA:130:LEU:O	2.13	0.48
3:FB:237:GLU:HG2	3:FB:238:ASN:H	1.79	0.48
2:KB:8:ASP:O	2:KB:12:ILE:HG12	2.12	0.48
4:NB:65:THR:O	4:OB:99:ILE:HA	2.13	0.48
2:CC:66:PHE:HB3	2:IC:45:ARG:HD2	1.94	0.48
3:DC:58:ASN:OD1	3:DC:81:VAL:HG11	2.14	0.48
2:UC:155:HIS:O	2:UC:159:LEU:HD22	2.13	0.48
3:VC:202:THR:O	3:VC:202:THR:OG1	2.32	0.48
2:AD:172:LEU:HD12	2:AD:172:LEU:HA	1.71	0.48
2:GD:126:HIS:CD2	3:HD:127:VAL:HG13	2.48	0.48
2:MD:89:ARG:HG3	2:MD:90:ALA:N	2.28	0.48
2:MD:98:LEU:HB3	2:MD:108:GLU:OE1	2.14	0.48
2:YD:263:LEU:HD11	2:YD:274:PHE:HD2	1.77	0.48
3:ZD:65:PHE:CE1	3:ZD:161:ALA:HB2	2.48	0.48
3:ZD:77:PRO:HB3	3:ZD:208:ILE:HB	1.95	0.48
2:EE:109:THR:O	2:EE:113:MET:HG3	2.13	0.48
2:EE:128:GLN:HG2	3:FE:131:PHE:CB	2.37	0.48
2:EE:133:ILE:O	2:EE:137:LEU:HG	2.14	0.48
3:FE:72:LEU:HD23	3:FE:73:LEU:HD12	1.95	0.48
2:QE:178:VAL:HG11	3:RE:149:THR:HG22	1.96	0.48
3:RE:54:LEU:HB2	3:RE:231:LEU:HD23	1.95	0.48
2:WE:196:VAL:HG12	2:WE:232:GLU:OE2	2.14	0.48
3:DF:86:ILE:HG13	3:DF:200:VAL:HG12	1.95	0.48
2:IF:89:ARG:HG3	2:IF:90:ALA:N	2.29	0.48
2:UF:230:ILE:HA	2:UF:233:MET:CE	2.43	0.48
3:VF:86:ILE:HG13	3:VF:198:ILE:HD11	1.95	0.48
2:AG:263:LEU:HD11	2:AG:274:PHE:HD2	1.78	0.48

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:BG:117:VAL:O	3:BG:215:PHE:HB2	2.13	0.48
2:GG:65:GLN:HE21	2:MG:47:ILE:H	1.60	0.48
3:HG:118:PHE:CE2	3:HG:158:LEU:HD12	2.49	0.48
2:MG:65:GLN:HG3	2:SG:46:GLN:HG2	1.96	0.48
3:TG:225:GLU:HA	3:TG:228:ARG:HB3	1.95	0.48
2:T:230:ILE:HA	2:T:233:MET:HE2	1.95	0.48
3:V:106:HIS:CE1	3:V:113:THR:CG2	2.96	0.48
2:GA:53:THR:O	2:GA:57:SER:OG	2.29	0.48
2:GA:89:ARG:CD	2:GA:90:ALA:H	2.27	0.48
2:GA:100:THR:OG1	2:GA:101:ARG:NH2	2.46	0.48
2:GA:127:PRO:O	2:GA:130:ILE:HG12	2.12	0.48
2:GA:201:GLU:HG3	2:MA:139:ARG:NH2	2.28	0.48
2:KB:174:GLU:HG2	3:LB:152:ARG:HD2	1.95	0.48
2:QB:11:VAL:HG11	2:QB:41:MET:HB3	1.96	0.48
2:QB:133:ILE:O	2:QB:137:LEU:HG	2.13	0.48
2:WB:244:ASP:OD1	2:WB:276:ARG:NH2	2.47	0.48
2:CC:78:ARG:O	2:CC:82:VAL:HG12	2.13	0.48
2:CC:101:ARG:O	2:CC:104:THR:HG23	2.13	0.48
2:OC:101:ARG:O	2:OC:104:THR:HG23	2.13	0.48
2:OC:128:GLN:CG	3:PC:131:PHE:HB3	2.41	0.48
2:OC:204:ASN:OD1	2:OC:233:MET:HB3	2.12	0.48
3:BD:79:ILE:HG22	3:BD:204:PHE:HB3	1.94	0.48
3:ND:247:ARG:HG2	3:ND:247:ARG:HH11	1.78	0.48
2:SD:100:THR:OG1	2:SD:101:ARG:NH2	2.45	0.48
2:YD:115:PRO:O	2:YD:119:ALA:HB2	2.13	0.48
2:EE:67:ALA:O	2:EE:71:ILE:HG12	2.14	0.48
2:EE:210:GLN:O	2:EE:214:VAL:HG12	2.12	0.48
2:KE:120:ASP:OD2	3:RE:76:SER:N	2.46	0.48
3:RE:106:HIS:CD2	3:RE:113:THR:HG22	2.49	0.48
2:WE:15:MET:SD	2:WE:15:MET:N	2.86	0.48
2:CF:11:VAL:O	2:CF:15:MET:HE3	2.12	0.48
2:UF:8:ASP:O	2:UF:12:ILE:HG12	2.13	0.48
2:GG:40:ALA:O	2:GG:44:VAL:HG23	2.14	0.48
2:SG:209:GLN:O	2:SG:212:GLU:HG3	2.14	0.48
3:M:202:THR:O	3:M:202:THR:OG1	2.30	0.48
2:J:114:GLU:HB2	2:J:117:SER:OG	2.13	0.48
2:T:159:LEU:HD21	3:V:137:PHE:CD1	2.49	0.48
3:V:207:GLU:OE1	3:V:207:GLU:N	2.46	0.48
3:BA:192:THR:OG1	3:BA:194:SER:O	2.31	0.48
2:GA:188:LEU:HB3	3:NA:74:ARG:NE	2.28	0.48
3:HA:180:VAL:HG23	3:HA:181:ARG:N	2.29	0.48

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:MA:65:GLN:NE2	2:SA:47:ILE:O	2.43	0.48
3:NA:100:THR:HG23	3:NA:119:SER:HA	1.96	0.48
2:SA:186:GLN:C	3:ZA:74:ARG:HH22	2.17	0.48
3:FB:96:LEU:HD23	3:FB:97:PRO:CD	2.43	0.48
2:KB:9:LYS:HG2	2:KB:55:VAL:HG22	1.95	0.48
3:LB:190:ASN:ND2	3:RB:234:PRO:O	2.46	0.48
2:WB:107:ILE:HD12	2:WB:107:ILE:N	2.28	0.48
2:CC:98:LEU:HB3	2:CC:108:GLU:OE1	2.13	0.48
2:CC:121:LEU:HD22	3:JC:70:PHE:CE1	2.48	0.48
2:CC:131:ALA:HA	2:CC:134:LEU:HD12	1.95	0.48
2:OC:83:LYS:HB2	2:UC:23:ALA:O	2.13	0.48
2:UC:87:GLU:HG2	2:UC:89:ARG:N	2.29	0.48
4:SE:101:ILE:N	4:SE:104:TYR:O	2.36	0.48
2:WE:107:ILE:O	2:WE:111:ASN:ND2	2.47	0.48
2:IF:73:ALA:HA	2:OF:41:MET:SD	2.54	0.48
2:OF:115:PRO:O	2:OF:119:ALA:HB2	2.14	0.48
3:VF:65:PHE:CE1	3:VF:161:ALA:HB2	2.47	0.48
2:GG:210:GLN:O	2:GG:214:VAL:HG12	2.14	0.48
3:TG:111:ARG:NH2	3:TG:222:SER:O	2.46	0.48
2:G:120:ASP:OD1	3:C:76:SER:OG	2.30	0.48
2:B:133:ILE:O	2:B:137:LEU:HG	2.14	0.48
3:C:110:LEU:HG	3:C:223:MET:HB3	1.95	0.48
2:J:128:GLN:CG	3:K:131:PHE:HB3	2.34	0.48
3:K:169:TRP:CD1	3:K:175:LEU:HD13	2.47	0.48
2:T:210:GLN:O	2:T:214:VAL:HG12	2.14	0.48
3:HA:169:TRP:CD1	3:HA:175:LEU:HD13	2.46	0.48
3:FB:239:SER:O	3:FB:242:GLU:HG2	2.13	0.48
2:KB:230:ILE:HA	2:KB:233:MET:CE	2.43	0.48
2:QB:210:GLN:O	2:QB:214:VAL:HG12	2.14	0.48
2:WB:188:LEU:HD22	3:DC:74:ARG:HH21	1.78	0.48
2:IC:210:GLN:O	2:IC:214:VAL:HG12	2.14	0.48
2:OC:107:ILE:HD12	2:OC:107:ILE:N	2.29	0.48
2:UC:113:MET:HE1	2:UC:121:LEU:HD21	1.96	0.48
2:UC:178:VAL:HG11	3:VC:149:THR:HG22	1.95	0.48
2:AD:120:ASP:OD1	3:HD:76:SER:OG	2.30	0.48
2:AD:167:VAL:HG21	3:BD:131:PHE:CD1	2.49	0.48
3:HD:180:VAL:HG23	3:HD:181:ARG:H	1.78	0.48
2:MD:133:ILE:O	2:MD:137:LEU:HG	2.13	0.48
2:EE:65:GLN:NE2	2:KE:47:ILE:O	2.45	0.48
2:EE:99:GLU:HA	2:EE:108:GLU:HG2	1.95	0.48
2:EE:152:ARG:NH2	2:EE:153:LEU:HB2	2.28	0.48

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:FE:180:VAL:HG23	3:FE:181:ARG:H	1.77	0.48
3:LE:174:PRO:O	3:LE:174:PRO:HD2	2.12	0.48
2:QE:65:GLN:HE21	2:WE:47:ILE:H	1.60	0.48
2:CF:149:PHE:CG	2:CF:153:LEU:HD23	2.49	0.48
3:DF:79:ILE:HG22	3:DF:204:PHE:HB3	1.96	0.48
2:IF:31:THR:O	2:IF:34:VAL:HB	2.14	0.48
2:OF:67:ALA:O	2:OF:71:ILE:HG12	2.14	0.48
2:OF:84:ALA:O	2:OF:87:GLU:HB3	2.14	0.48
3:PF:101:ASN:HB2	3:PF:146:PHE:CE2	2.49	0.48
4:QG:106:ILE:HA	3:TG:252:ARG:HG3	1.95	0.48
2:B:31:THR:O	2:B:34:VAL:HB	2.14	0.48
2:B:63:ALA:HA	2:B:66:PHE:CZ	2.48	0.48
2:B:168:GLN:OE1	2:B:168:GLN:N	2.39	0.48
2:J:30:SER:OG	2:J:33:GLU:OE1	2.30	0.48
2:J:108:GLU:HG3	2:J:112:PHE:HE2	1.79	0.48
2:T:53:THR:O	2:T:57:SER:OG	2.29	0.48
2:MA:128:GLN:CG	3:NA:131:PHE:HB3	2.33	0.48
2:SA:65:GLN:HE21	2:YA:47:ILE:H	1.62	0.48
2:YA:76:TYR:HH	2:EB:34:VAL:HG13	1.78	0.48
2:YA:80:VAL:HA	2:YA:83:LYS:CD	2.42	0.48
3:ZA:96:LEU:HD23	3:ZA:97:PRO:HD2	1.95	0.48
2:EB:133:ILE:O	2:EB:137:LEU:HG	2.14	0.48
3:FB:227:LEU:HD13	3:FB:230:LEU:HD13	1.95	0.48
2:KB:41:MET:HG2	2:KB:42:ALA:N	2.29	0.48
2:QB:146:LEU:HD11	2:QB:157:VAL:HG11	1.94	0.48
2:WB:81:LEU:H	2:WB:81:LEU:HD12	1.78	0.48
2:WB:83:LYS:HB3	2:CC:23:ALA:HB1	1.95	0.48
2:CC:97:ILE:O	2:CC:101:ARG:NH2	2.34	0.48
3:DC:117:VAL:O	3:DC:215:PHE:HB2	2.14	0.48
2:UC:26:PHE:HE1	2:UC:34:VAL:HG22	1.77	0.48
2:UC:128:GLN:NE2	2:UC:164:PHE:HD2	2.12	0.48
2:AD:26:PHE:HE2	2:AD:37:LEU:HD12	1.79	0.48
3:BD:65:PHE:CE1	3:BD:161:ALA:HB2	2.48	0.48
2:GD:205:LEU:HD22	2:MD:140:SER:HA	1.94	0.48
2:MD:40:ALA:O	2:MD:44:VAL:HG23	2.14	0.48
2:SD:149:PHE:CD2	2:SD:153:LEU:HD22	2.49	0.48
2:SD:261:ILE:O	2:SD:264:LYS:HG3	2.14	0.48
3:ZD:238:ASN:HB3	3:ZD:241:HIS:ND1	2.28	0.48
2:KE:69:LEU:HD23	2:QE:47:ILE:HB	1.96	0.48
3:LE:202:THR:O	3:LE:202:THR:OG1	2.30	0.48
3:RE:180:VAL:HG23	3:RE:181:ARG:N	2.29	0.48

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:WE:119:ALA:HB1	2:WE:149:PHE:CE1	2.48	0.48
2:CF:108:GLU:HG3	2:CF:112:PHE:HE2	1.77	0.48
2:CF:151:GLU:O	2:CF:155:HIS:ND1	2.47	0.48
3:JF:70:PHE:HD1	3:JF:75:ARG:O	1.96	0.48
2:AG:144:ASP:O	2:AG:148:LEU:HD12	2.14	0.48
2:G:87:GLU:HG2	2:G:89:ARG:H	1.79	0.48
2:G:151:GLU:HA	2:G:154:ARG:HE	1.79	0.48
2:G:201:GLU:OE1	2:B:139:ARG:NH1	2.46	0.48
2:B:83:LYS:HB3	2:J:26:PHE:HB3	1.96	0.48
3:BA:169:TRP:CD1	3:BA:175:LEU:HD13	2.48	0.48
3:BA:190:ASN:ND2	3:BA:190:ASN:O	2.47	0.48
3:BA:240:ARG:HA	3:BA:240:ARG:CZ	2.44	0.48
2:WB:101:ARG:O	2:WB:104:THR:HG23	2.14	0.48
2:WB:188:LEU:HD22	3:DC:74:ARG:HE	1.79	0.48
3:XB:240:ARG:HA	3:XB:240:ARG:HH11	1.77	0.48
2:CC:261:ILE:O	2:CC:264:LYS:HG3	2.14	0.48
2:IC:103:THR:C	2:IC:106:GLY:H	2.17	0.48
3:JC:86:ILE:HD11	3:JC:221:PHE:CE1	2.49	0.48
3:VC:207:GLU:N	3:VC:207:GLU:OE2	2.47	0.48
3:VC:247:ARG:HG2	3:VC:247:ARG:HH11	1.78	0.48
2:AD:167:VAL:HG13	3:BD:130:LEU:O	2.14	0.48
2:AD:174:GLU:HG2	3:BD:152:ARG:HG2	1.95	0.48
2:AD:261:ILE:O	2:AD:264:LYS:HG3	2.14	0.48
2:SD:79:SER:O	2:SD:82:VAL:HG22	2.14	0.48
2:YD:69:LEU:HD11	2:EE:47:ILE:CG2	2.41	0.48
2:QE:26:PHE:CE1	2:QE:29:LEU:HD11	2.49	0.48
3:RE:58:ASN:OD1	3:RE:81:VAL:HG11	2.13	0.48
2:WE:9:LYS:HG2	2:WE:55:VAL:HG22	1.94	0.48
2:CF:207:LYS:HZ1	2:CF:209:GLN:HB3	1.79	0.48
2:IF:152:ARG:HA	2:IF:155:HIS:HE1	1.79	0.48
2:IF:201:GLU:HB3	2:OF:139:ARG:HH21	1.79	0.48
2:OF:245:ARG:O	2:OF:245:ARG:NH1	2.37	0.48
2:UF:48:SER:OG	2:UF:51:GLN:OE1	2.28	0.48
2:UF:100:THR:OG1	2:UF:101:ARG:NH2	2.47	0.48
3:VF:224:ILE:O	3:VF:228:ARG:N	2.47	0.48
2:AG:48:SER:OG	2:AG:51:GLN:OE1	2.31	0.48
2:AG:178:VAL:HG11	3:BG:149:THR:HG22	1.96	0.48
2:GG:128:GLN:CG	3:HG:131:PHE:HB3	2.35	0.48
2:GG:151:GLU:O	2:GG:155:HIS:ND1	2.47	0.48
2:MG:69:LEU:HD13	2:SG:45:ARG:O	2.14	0.48
2:SG:30:SER:OG	2:SG:33:GLU:OE1	2.27	0.48

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:G:76:TYR:HE2	2:B:26:PHE:CE1	2.31	0.48
2:B:9:LYS:HG2	2:B:55:VAL:HG22	1.95	0.48
3:C:69:LEU:O	3:C:73:LEU:HB2	2.13	0.48
2:J:82:VAL:HG13	2:T:23:ALA:HB2	1.96	0.48
2:T:151:GLU:HA	2:T:154:ARG:HE	1.79	0.48
3:BA:122:LEU:CD1	3:BA:215:PHE:HB3	2.41	0.48
2:GA:172:LEU:HD12	2:GA:172:LEU:HA	1.73	0.48
3:HA:185:GLN:HG3	3:HA:186:VAL:H	1.79	0.48
2:MA:151:GLU:HA	2:MA:154:ARG:HE	1.79	0.48
2:EB:108:GLU:HA	2:EB:111:ASN:HD22	1.78	0.48
3:LB:86:ILE:HD12	3:LB:200:VAL:HG23	1.96	0.48
2:QB:10:SER:O	2:QB:14:LEU:HD22	2.13	0.48
2:QB:31:THR:HA	2:QB:34:VAL:HG23	1.95	0.48
2:WB:174:GLU:OE2	3:XB:152:ARG:HG2	2.14	0.48
2:CC:242:VAL:HG13	2:CC:323:ILE:HG12	1.96	0.48
2:IC:17:ILE:HD12	2:IC:21:ARG:HG3	1.95	0.48
2:IC:195:GLY:HA2	2:OC:161:ILE:O	2.14	0.48
2:OC:206:MET:SD	2:OC:210:GLN:HB2	2.54	0.48
3:PC:65:PHE:CE1	3:PC:161:ALA:HB2	2.48	0.48
2:AD:41:MET:HG2	2:AD:42:ALA:N	2.29	0.48
3:BD:118:PHE:CE2	3:BD:158:LEU:HD12	2.48	0.48
2:SD:159:LEU:HD21	3:TD:137:PHE:CD1	2.49	0.48
2:SD:244:ASP:HA	2:SD:247:ILE:HG12	1.95	0.48
3:LE:180:VAL:HG23	3:LE:181:ARG:N	2.29	0.48
2:WE:84:ALA:O	2:WE:87:GLU:HB3	2.14	0.48
2:CF:30:SER:O	2:CF:34:VAL:HG23	2.14	0.48
3:BG:185:GLN:HG3	3:BG:186:VAL:H	1.79	0.48
3:BG:190:ASN:O	3:HG:236:LEU:HD13	2.14	0.48
2:MG:188:LEU:HD22	3:TG:74:ARG:NH1	2.29	0.48
2:G:261:ILE:O	2:G:264:LYS:HG3	2.14	0.47
3:C:101:ASN:HB2	3:C:146:PHE:CE2	2.48	0.47
3:K:180:VAL:HG23	3:K:181:ARG:N	2.29	0.47
2:AA:40:ALA:O	2:AA:44:VAL:HG23	2.13	0.47
2:AA:128:GLN:HG2	3:BA:131:PHE:CB	2.34	0.47
3:BA:111:ARG:CZ	3:BA:111:ARG:HA	2.44	0.47
3:BA:180:VAL:HG23	3:BA:181:ARG:N	2.29	0.47
2:GA:7:THR:HG23	2:GA:37:LEU:HD23	1.96	0.47
2:GA:133:ILE:O	2:GA:137:LEU:HG	2.13	0.47
2:MA:115:PRO:O	2:MA:119:ALA:HB2	2.13	0.47
3:NA:190:ASN:O	3:TA:236:LEU:HD12	2.13	0.47
3:NA:214:GLU:OE1	3:NA:215:PHE:N	2.47	0.47

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:YA:89:ARG:NH2	2:EB:31:THR:HG23	2.29	0.47
2:EB:203:ILE:HA	2:EB:206:MET:CE	2.44	0.47
2:KB:76:TYR:CD2	2:QB:41:MET:HE3	2.49	0.47
2:KB:210:GLN:O	2:KB:214:VAL:HG12	2.14	0.47
2:QB:117:SER:O	2:QB:121:LEU:HG	2.14	0.47
2:QB:230:ILE:HA	2:QB:233:MET:HE1	1.96	0.47
2:WB:8:ASP:O	2:WB:12:ILE:HG12	2.14	0.47
2:WB:65:GLN:HE21	2:CC:47:ILE:H	1.62	0.47
2:CC:15:MET:SD	2:CC:15:MET:N	2.87	0.47
3:JC:202:THR:OG1	3:JC:202:THR:O	2.32	0.47
2:OC:267:GLU:HB3	2:OC:269:PRO:HD2	1.96	0.47
2:UC:151:GLU:HA	2:UC:154:ARG:HE	1.79	0.47
2:SD:167:VAL:HG13	3:TD:130:LEU:O	2.14	0.47
2:EE:20:ASP:OD2	2:EE:21:ARG:N	2.47	0.47
2:EE:190:ARG:O	2:KE:166:GLY:HA3	2.14	0.47
2:KE:151:GLU:HA	2:KE:154:ARG:HE	1.79	0.47
3:RE:101:ASN:HB2	3:RE:146:PHE:CE1	2.49	0.47
2:WE:152:ARG:HH22	2:WE:153:LEU:HB2	1.79	0.47
2:CF:89:ARG:HG3	2:CF:90:ALA:N	2.29	0.47
3:DF:207:GLU:N	3:DF:207:GLU:OE2	2.47	0.47
2:IF:87:GLU:HG2	2:IF:89:ARG:N	2.29	0.47
2:OF:26:PHE:HZ	2:OF:34:VAL:HG13	1.78	0.47
2:OF:121:LEU:HD22	3:VF:70:PHE:CE1	2.49	0.47
2:OF:167:VAL:HG13	3:PF:130:LEU:O	2.14	0.47
2:OF:178:VAL:HG11	3:PF:149:THR:HG22	1.96	0.47
2:UF:101:ARG:O	2:UF:104:THR:HG23	2.14	0.47
2:UF:204:ASN:OD1	2:UF:233:MET:HB3	2.14	0.47
2:AG:188:LEU:HB2	2:GG:168:GLN:HB3	1.95	0.47
2:AG:261:ILE:O	2:AG:264:LYS:HG3	2.14	0.47
2:GG:267:GLU:HB3	2:GG:269:PRO:HD2	1.96	0.47
3:NG:294:ALA:O	3:NG:301:VAL:N	2.45	0.47
2:G:267:GLU:HB3	2:G:269:PRO:HD2	1.96	0.47
3:K:58:ASN:OD1	3:K:81:VAL:HG11	2.14	0.47
2:T:219:ARG:NH2	2:T:227:GLN:OE1	2.37	0.47
2:T:245:ARG:O	2:T:245:ARG:NH1	2.37	0.47
3:V:102:LEU:HD22	3:V:117:VAL:HG22	1.96	0.47
2:AA:155:HIS:O	2:AA:159:LEU:HG	2.14	0.47
2:GA:128:GLN:CG	3:HA:131:PHE:HB3	2.37	0.47
3:HA:59:GLU:OE2	3:HA:59:GLU:N	2.47	0.47
2:MA:210:GLN:O	2:MA:214:VAL:HG12	2.15	0.47
2:QB:101:ARG:O	2:QB:104:THR:HG23	2.14	0.47

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:QB:244:ASP:OD1	2:QB:276:ARG:NH2	2.47	0.47
2:WB:159:LEU:HD21	3:XB:137:PHE:CD1	2.49	0.47
2:WB:167:VAL:HG13	3:XB:130:LEU:O	2.14	0.47
3:XB:207:GLU:OE1	3:XB:207:GLU:N	2.47	0.47
2:CC:30:SER:O	2:CC:34:VAL:HG23	2.14	0.47
2:CC:69:LEU:HD11	2:IC:47:ILE:HB	1.95	0.47
3:DC:180:VAL:HG23	3:DC:181:ARG:N	2.29	0.47
2:GD:128:GLN:CG	3:HD:131:PHE:HB3	2.33	0.47
2:MD:193:MET:N	2:MD:193:MET:SD	2.88	0.47
2:MD:244:ASP:HA	2:MD:247:ILE:HG12	1.95	0.47
3:ND:65:PHE:CE1	3:ND:161:ALA:HB2	2.49	0.47
1:XD:547:VAL:HA	2:YD:28:HIS:NE2	2.29	0.47
2:EE:12:ILE:HD11	2:EE:55:VAL:HG11	1.94	0.47
2:EE:167:VAL:HG13	3:FE:130:LEU:O	2.14	0.47
2:KE:167:VAL:HG13	3:LE:130:LEU:O	2.13	0.47
2:KE:263:LEU:HD11	2:KE:274:PHE:HD2	1.79	0.47
2:WE:65:GLN:HE21	2:CF:47:ILE:H	1.61	0.47
2:WE:128:GLN:CG	3:XE:131:PHE:HB3	2.37	0.47
2:WE:206:MET:SD	2:WE:210:GLN:HB3	2.53	0.47
3:XE:239:SER:O	3:XE:242:GLU:HG2	2.14	0.47
2:OF:242:VAL:HG13	2:OF:323:ILE:HG12	1.97	0.47
3:VF:58:ASN:OD1	3:VF:81:VAL:HG11	2.14	0.47
2:AG:83:LYS:HB2	2:GG:23:ALA:O	2.14	0.47
2:GG:111:ASN:HA	2:GG:141:GLN:HE22	1.78	0.47
2:MG:151:GLU:HA	2:MG:154:ARG:HE	1.78	0.47
3:TG:116:VAL:HG21	3:TG:158:LEU:HD11	1.96	0.47
2:J:100:THR:OG1	2:J:101:ARG:NH2	2.47	0.47
2:J:126:HIS:CE1	3:K:127:VAL:HG13	2.48	0.47
2:J:151:GLU:HA	2:J:154:ARG:HE	1.80	0.47
2:J:261:ILE:O	2:J:264:LYS:HG3	2.14	0.47
2:T:261:ILE:O	2:T:264:LYS:HG3	2.14	0.47
3:BA:109:PRO:O	3:BA:111:ARG:NH1	2.46	0.47
3:NA:118:PHE:CE2	3:NA:158:LEU:HD12	2.49	0.47
3:NA:150:GLU:O	3:NA:153:VAL:HG12	2.14	0.47
2:SA:79:SER:HA	2:SA:82:VAL:CG1	2.44	0.47
2:SA:167:VAL:HG13	3:TA:130:LEU:O	2.14	0.47
3:RB:60:ARG:HE	3:RB:60:ARG:HB2	1.55	0.47
3:JC:111:ARG:CZ	3:JC:111:ARG:HA	2.44	0.47
3:PC:183:GLU:HG3	3:PC:188:PHE:HB2	1.95	0.47
2:AD:187:ASN:N	3:HD:74:ARG:HH22	2.13	0.47
3:BD:54:LEU:HD13	3:BD:231:LEU:HD21	1.94	0.47

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:BD:227:LEU:HD13	3:BD:230:LEU:HD13	1.97	0.47
2:GD:152:ARG:NH2	2:GD:153:LEU:HB2	2.29	0.47
3:HD:113:THR:OG1	3:HD:191:ILE:HG22	2.14	0.47
2:MD:160:ARG:O	2:MD:164:PHE:CB	2.62	0.47
3:TD:207:GLU:N	3:TD:207:GLU:OE2	2.47	0.47
2:YD:142:ALA:O	2:YD:146:LEU:HG	2.14	0.47
3:ZD:97:PRO:O	3:ZD:100:THR:OG1	2.21	0.47
3:ZD:102:LEU:HD22	3:ZD:117:VAL:HG22	1.96	0.47
3:FE:239:SER:O	3:FE:242:GLU:HG2	2.13	0.47
2:KE:186:GLN:HE22	3:RE:71:ASN:CG	2.18	0.47
2:KE:219:ARG:NH2	2:KE:227:GLN:OE1	2.32	0.47
2:QE:261:ILE:O	2:QE:264:LYS:HG3	2.14	0.47
3:RE:224:ILE:O	3:RE:228:ARG:N	2.48	0.47
3:XE:70:PHE:O	3:XE:74:ARG:NH1	2.47	0.47
2:CF:12:ILE:HD11	2:CF:55:VAL:HG11	1.96	0.47
4:MF:106:ILE:O	3:PF:252:ARG:HG2	2.14	0.47
2:OF:151:GLU:HA	2:OF:154:ARG:HE	1.79	0.47
3:PF:180:VAL:HG23	3:PF:181:ARG:N	2.29	0.47
2:GG:21:ARG:NH2	2:GG:24:GLU:HB2	2.29	0.47
2:SG:87:GLU:HG2	2:SG:89:ARG:N	2.30	0.47
2:SG:183:LEU:O	2:SG:186:GLN:N	2.47	0.47
3:TG:74:ARG:O	3:TG:75:ARG:HD2	2.13	0.47
3:TG:202:THR:OG1	3:TG:202:THR:O	2.32	0.47
2:G:139:ARG:HH21	2:SG:202:ILE:HD13	1.80	0.47
2:G:187:ASN:N	3:C:74:ARG:HH22	2.12	0.47
2:J:85:LEU:H	2:J:85:LEU:HD22	1.79	0.47
2:J:267:GLU:HB3	2:J:269:PRO:HD2	1.96	0.47
4:R:107:ALA:HA	4:R:125:ILE:HA	1.96	0.47
2:T:12:ILE:HD11	2:T:55:VAL:HG11	1.97	0.47
2:T:101:ARG:O	2:T:104:THR:HG23	2.15	0.47
3:V:174:PRO:HD2	3:V:174:PRO:O	2.14	0.47
3:BA:246:TRP:HB2	3:BA:247:ARG:HH12	1.79	0.47
2:GA:115:PRO:O	2:GA:119:ALA:CB	2.62	0.47
3:HA:65:PHE:CE1	3:HA:161:ALA:HB2	2.49	0.47
2:MA:167:VAL:HG13	3:NA:130:LEU:O	2.14	0.47
2:MA:174:GLU:HG2	3:NA:152:ARG:HG2	1.95	0.47
2:MA:298:LEU:HB2	2:MA:302:GLU:OE1	2.14	0.47
3:ZA:86:ILE:HD11	3:ZA:221:PHE:HE1	1.79	0.47
2:KB:34:VAL:O	2:KB:37:LEU:HD12	2.15	0.47
2:KB:53:THR:O	2:KB:57:SER:OG	2.28	0.47
3:LB:52:GLN:HA	3:LB:55:GLU:HG3	1.96	0.47

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:QB:109:THR:O	2:QB:113:MET:HG3	2.14	0.47
3:RB:116:VAL:HG21	3:RB:158:LEU:HD11	1.95	0.47
2:WB:100:THR:OG1	2:WB:101:ARG:NH2	2.46	0.47
2:OC:167:VAL:HG13	3:PC:130:LEU:O	2.14	0.47
3:PC:285:PRO:HA	4:QC:119:GLY:HA2	1.97	0.47
2:UC:89:ARG:HG3	2:UC:90:ALA:N	2.29	0.47
2:GD:167:VAL:HG21	3:HD:131:PHE:CD1	2.49	0.47
2:MD:172:LEU:HD12	2:MD:172:LEU:HA	1.78	0.47
2:SD:108:GLU:HA	2:SD:111:ASN:ND2	2.23	0.47
2:EE:263:LEU:HD11	2:EE:274:PHE:HD2	1.79	0.47
2:KE:186:GLN:CB	3:RE:74:ARG:HH12	2.18	0.47
2:CF:152:ARG:NH2	2:CF:153:LEU:HB2	2.28	0.47
2:IF:107:ILE:O	2:IF:111:ASN:ND2	2.48	0.47
2:IF:178:VAL:HG11	3:JF:149:THR:HG22	1.96	0.47
3:VF:97:PRO:HD2	3:VF:100:THR:OG1	2.14	0.47
2:AG:65:GLN:HE21	2:GG:47:ILE:H	1.61	0.47
2:AG:89:ARG:CD	2:AG:90:ALA:H	2.26	0.47
2:AG:141:GLN:HA	2:AG:144:ASP:OD2	2.15	0.47
2:MG:267:GLU:HB3	2:MG:269:PRO:HD2	1.96	0.47
2:SG:193:MET:O	2:SG:195:GLY:N	2.48	0.47
3:TG:60:ARG:HE	3:TG:60:ARG:HB2	1.55	0.47
2:G:47:ILE:H	2:SG:65:GLN:HE21	1.62	0.47
3:C:237:GLU:HG2	3:C:238:ASN:N	2.28	0.47
2:J:209:GLN:O	2:J:212:GLU:HG3	2.15	0.47
3:HA:106:HIS:CD2	3:HA:113:THR:HG22	2.49	0.47
2:MA:75:GLU:O	2:MA:79:SER:HB3	2.14	0.47
2:MA:119:ALA:HB1	2:MA:149:PHE:CE1	2.49	0.47
2:SA:219:ARG:NH1	2:SA:223:GLY:O	2.47	0.47
2:KB:107:ILE:HG23	2:KB:137:LEU:HD23	1.96	0.47
2:KB:128:GLN:NE2	2:KB:164:PHE:HD2	2.06	0.47
3:LB:224:ILE:O	3:LB:228:ARG:N	2.47	0.47
2:QB:195:GLY:HA2	2:WB:161:ILE:O	2.14	0.47
2:QB:206:MET:SD	2:QB:210:GLN:HB2	2.53	0.47
2:QB:245:ARG:O	2:QB:245:ARG:NH1	2.38	0.47
3:RB:101:ASN:HD22	3:RB:146:PHE:HE2	1.61	0.47
2:CC:83:LYS:O	2:IC:27:LYS:HD2	2.14	0.47
3:DC:111:ARG:HA	3:DC:111:ARG:CZ	2.44	0.47
2:IC:267:GLU:HB3	2:IC:269:PRO:HD2	1.96	0.47
2:OC:63:ALA:HA	2:OC:66:PHE:CE2	2.49	0.47
2:OC:109:THR:O	2:OC:113:MET:HG3	2.14	0.47
3:VC:191:ILE:HG13	3:VC:192:THR:HG23	1.97	0.47

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:VC:237:GLU:HG2	3:VC:238:ASN:H	1.79	0.47
2:GD:139:ARG:HD3	2:GD:139:ARG:HA	1.72	0.47
2:YD:160:ARG:O	2:YD:164:PHE:CB	2.61	0.47
3:ZD:100:THR:HG23	3:ZD:119:SER:HA	1.95	0.47
2:KE:133:ILE:O	2:KE:137:LEU:HG	2.14	0.47
2:QE:149:PHE:CD2	2:QE:153:LEU:HD22	2.49	0.47
2:CF:159:LEU:HD21	3:DF:137:PHE:CD1	2.49	0.47
2:IF:141:GLN:HA	2:IF:144:ASP:OD2	2.14	0.47
2:OF:9:LYS:HG2	2:OF:55:VAL:HG22	1.95	0.47
2:UF:79:SER:HA	2:UF:82:VAL:CG1	2.43	0.47
2:UF:188:LEU:HD22	3:BG:74:ARG:HE	1.79	0.47
3:BG:65:PHE:CE1	3:BG:161:ALA:HB2	2.50	0.47
2:GG:113:MET:HE1	2:GG:118:ALA:HA	1.95	0.47
2:MG:174:GLU:OE2	3:NG:152:ARG:HG2	2.14	0.47
2:B:149:PHE:CD2	2:B:153:LEU:HD22	2.49	0.47
2:B:167:VAL:HG13	3:C:130:LEU:O	2.15	0.47
3:C:202:THR:OG1	3:C:202:THR:O	2.32	0.47
3:K:180:VAL:HG23	3:K:181:ARG:H	1.79	0.47
3:K:198:ILE:HG23	3:K:221:PHE:HD1	1.78	0.47
2:GA:12:ILE:HD11	2:GA:55:VAL:HG11	1.96	0.47
2:GA:141:GLN:HA	2:GA:144:ASP:OD2	2.15	0.47
2:MA:87:GLU:HG2	2:MA:89:ARG:N	2.29	0.47
2:SA:128:GLN:CG	3:TA:131:PHE:HB3	2.37	0.47
3:FB:65:PHE:CE1	3:FB:161:ALA:HB2	2.49	0.47
2:CC:115:PRO:O	2:CC:119:ALA:CB	2.63	0.47
3:DC:185:GLN:HG3	3:DC:186:VAL:H	1.80	0.47
2:OC:153:LEU:O	2:OC:157:VAL:HG23	2.14	0.47
2:UC:8:ASP:O	2:UC:12:ILE:HG12	2.13	0.47
2:UC:267:GLU:HB3	2:UC:269:PRO:HD2	1.96	0.47
2:AD:267:GLU:HB3	2:AD:269:PRO:HD2	1.96	0.47
2:GD:101:ARG:O	2:GD:104:THR:HG23	2.15	0.47
2:GD:113:MET:HE1	2:GD:121:LEU:HD21	1.97	0.47
3:HD:202:THR:O	3:HD:202:THR:OG1	2.33	0.47
2:MD:109:THR:O	2:MD:113:MET:HG3	2.14	0.47
2:YD:26:PHE:CE1	2:YD:34:VAL:HG22	2.49	0.47
3:RE:185:GLN:HG3	3:RE:186:VAL:H	1.80	0.47
3:RE:285:PRO:HA	4:SE:119:GLY:HA2	1.97	0.47
2:WE:32:ARG:NE	2:WE:33:GLU:OE2	2.46	0.47
2:WE:115:PRO:O	2:WE:119:ALA:HB3	2.12	0.47
2:UF:15:MET:N	2:UF:15:MET:SD	2.87	0.47
2:UF:74:ASN:O	2:UF:78:ARG:NE	2.46	0.47

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:UF:267:GLU:HB3	2:UF:269:PRO:HD2	1.96	0.47
3:BG:180:VAL:HG23	3:BG:181:ARG:N	2.30	0.47
2:GG:133:ILE:O	2:GG:137:LEU:HG	2.15	0.47
3:HG:185:GLN:HG3	3:HG:186:VAL:H	1.78	0.47
2:MG:210:GLN:O	2:MG:214:VAL:HG12	2.15	0.47
2:SG:105:SER:OG	2:SG:107:ILE:HD11	2.15	0.47
2:G:84:ALA:O	2:G:87:GLU:HB3	2.14	0.47
2:G:115:PRO:O	2:G:119:ALA:HB2	2.13	0.47
2:G:225:LEU:HA	2:G:228:LYS:HD2	1.96	0.47
3:M:74:ARG:NH1	2:SG:188:LEU:HD21	2.27	0.47
4:Q:106:ILE:O	3:C:252:ARG:HG2	2.15	0.47
2:B:206:MET:SD	2:B:210:GLN:HB2	2.54	0.47
3:C:198:ILE:HG23	3:C:221:PHE:HD1	1.79	0.47
2:J:186:GLN:C	3:V:74:ARG:HH22	2.18	0.47
2:J:242:VAL:HG13	2:J:323:ILE:HG12	1.95	0.47
2:J:282:ALA:HA	2:J:285:ILE:HG12	1.97	0.47
2:T:89:ARG:HG3	2:T:90:ALA:N	2.30	0.47
2:T:267:GLU:HB3	2:T:269:PRO:HD2	1.96	0.47
3:V:135:GLY:O	3:V:211:LEU:HD21	2.14	0.47
3:V:169:TRP:CD1	3:V:175:LEU:HD13	2.48	0.47
2:AA:65:GLN:HE21	2:GA:47:ILE:H	1.63	0.47
3:BA:58:ASN:OD1	3:BA:81:VAL:HG11	2.14	0.47
3:BA:100:THR:HG23	3:BA:119:SER:HA	1.96	0.47
3:BA:180:VAL:HG23	3:BA:181:ARG:H	1.79	0.47
4:CA:101:ILE:N	4:CA:104:TYR:O	2.33	0.47
2:GA:14:LEU:HD12	2:GA:22:ALA:HB1	1.97	0.47
2:GA:167:VAL:HG13	3:HA:130:LEU:O	2.15	0.47
2:GA:267:GLU:HB3	2:GA:269:PRO:HD2	1.96	0.47
2:MA:65:GLN:HG3	2:SA:46:GLN:HG2	1.97	0.47
2:MA:85:LEU:H	2:MA:85:LEU:HD22	1.79	0.47
2:MA:230:ILE:HA	2:MA:233:MET:CE	2.44	0.47
3:NA:239:SER:O	3:NA:242:GLU:HG2	2.15	0.47
2:YA:87:GLU:HG2	2:YA:89:ARG:N	2.30	0.47
2:YA:178:VAL:HG11	3:ZA:149:THR:HG22	1.97	0.47
2:YA:186:GLN:C	3:FB:74:ARG:HH22	2.18	0.47
2:YA:244:ASP:OD1	2:YA:276:ARG:NH2	2.48	0.47
2:QB:160:ARG:O	2:QB:164:PHE:CB	2.62	0.47
4:UB:106:ILE:O	3:XB:252:ARG:HG2	2.15	0.47
2:WB:76:TYR:CE2	2:CC:38:SER:HB3	2.41	0.47
2:CC:20:ASP:OD1	2:CC:20:ASP:N	2.47	0.47
2:CC:76:TYR:CE2	2:IC:38:SER:HB3	2.50	0.47

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:CC:267:GLU:HB3	2:CC:269:PRO:HD2	1.96	0.47
2:IC:107:ILE:HD13	2:IC:136:HIS:CB	2.44	0.47
2:OC:157:VAL:O	2:OC:161:ILE:HG12	2.15	0.47
2:UC:76:TYR:HE2	2:AD:26:PHE:CE1	2.33	0.47
2:UC:114:GLU:HB2	2:UC:117:SER:OG	2.14	0.47
2:UC:152:ARG:NH2	2:UC:153:LEU:HB2	2.29	0.47
2:AD:83:LYS:HB2	2:GD:26:PHE:HB3	1.95	0.47
3:BD:294:ALA:O	3:BD:301:VAL:N	2.46	0.47
2:MD:141:GLN:HA	2:MD:144:ASP:OD2	2.15	0.47
2:MD:267:GLU:HB3	2:MD:269:PRO:HD2	1.96	0.47
3:ND:224:ILE:O	3:ND:228:ARG:N	2.48	0.47
4:PD:65:THR:O	4:QD:99:ILE:HA	2.15	0.47
2:SD:122:ILE:HB	2:SD:130:ILE:HG22	1.95	0.47
2:YD:119:ALA:HB1	2:YD:149:PHE:CE1	2.50	0.47
3:ZD:118:PHE:CE2	3:ZD:158:LEU:HD12	2.50	0.47
3:ZD:237:GLU:HG2	3:ZD:238:ASN:H	1.80	0.47
3:FE:180:VAL:HG23	3:FE:181:ARG:N	2.30	0.47
2:KE:84:ALA:O	2:KE:87:GLU:HB3	2.14	0.47
2:KE:95:GLU:O	2:KE:99:GLU:HB2	2.14	0.47
2:QE:128:GLN:CG	3:RE:131:PHE:HB3	2.39	0.47
3:RE:185:GLN:HG2	3:RE:187:LYS:HZ2	1.79	0.47
2:WE:244:ASP:OD1	2:WE:276:ARG:NH2	2.46	0.47
2:CF:107:ILE:O	2:CF:111:ASN:ND2	2.48	0.47
3:DF:117:VAL:O	3:DF:215:PHE:HB2	2.15	0.47
2:IF:98:LEU:HB3	2:IF:108:GLU:OE1	2.14	0.47
3:JF:115:LEU:HD11	3:JF:218:CYS:HB3	1.96	0.47
3:JF:202:THR:OG1	3:JF:202:THR:O	2.32	0.47
2:OF:76:TYR:CE2	2:UF:38:SER:HA	2.49	0.47
2:OF:87:GLU:HG2	2:OF:89:ARG:H	1.80	0.47
2:OF:261:ILE:O	2:OF:264:LYS:HG3	2.14	0.47
2:OF:267:GLU:HB3	2:OF:269:PRO:HD2	1.96	0.47
2:AG:75:GLU:O	2:AG:79:SER:HB3	2.14	0.47
3:BG:183:GLU:OE2	3:BG:184:MET:N	2.48	0.47
2:GG:69:LEU:HD11	2:MG:47:ILE:CG2	2.44	0.47
2:GG:188:LEU:HD22	3:NG:74:ARG:HH21	1.79	0.47
4:KG:106:ILE:O	3:NG:252:ARG:HG2	2.15	0.47
2:MG:9:LYS:HG2	2:MG:55:VAL:HG22	1.95	0.47
2:MG:188:LEU:HB3	3:TG:74:ARG:NE	2.30	0.47
2:SG:120:ASP:OD1	2:SG:123:ARG:NH1	2.47	0.47
2:SG:151:GLU:HA	2:SG:154:ARG:HE	1.80	0.47
3:TG:180:VAL:HG23	3:TG:181:ARG:H	1.79	0.47

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:TG:247:ARG:NH1	3:TG:251:VAL:HG23	2.23	0.47
2:T:111:ASN:CG	2:T:138:LYS:HG2	2.35	0.47
3:BA:118:PHE:CE2	3:BA:158:LEU:HD12	2.50	0.47
2:GA:174:GLU:HG2	3:HA:152:ARG:HG2	1.97	0.47
3:NA:207:GLU:OE1	3:NA:207:GLU:N	2.48	0.47
2:SA:114:GLU:HB2	2:SA:117:SER:OG	2.14	0.47
3:TA:111:ARG:NH2	3:TA:222:SER:O	2.48	0.47
2:YA:31:THR:O	2:YA:34:VAL:HB	2.15	0.47
2:EB:167:VAL:HG13	3:FB:130:LEU:O	2.14	0.47
3:LB:106:HIS:CD2	3:LB:113:THR:HG22	2.50	0.47
2:QB:7:THR:HG23	2:QB:37:LEU:HD23	1.96	0.47
2:QB:76:TYR:HE2	2:WB:38:SER:HB3	1.79	0.47
2:QB:83:LYS:HE2	2:QB:83:LYS:HB2	1.45	0.47
2:QB:267:GLU:HB3	2:QB:269:PRO:HD2	1.96	0.47
2:WB:74:ASN:O	2:WB:78:ARG:NE	2.48	0.47
2:WB:267:GLU:HB3	2:WB:269:PRO:HD2	1.96	0.47
3:DC:118:PHE:HE2	3:DC:158:LEU:HD12	1.79	0.47
2:IC:115:PRO:O	2:IC:119:ALA:HB2	2.14	0.47
2:IC:298:LEU:HB2	2:IC:302:GLU:OE2	2.15	0.47
2:OC:100:THR:OG1	2:OC:101:ARG:NH2	2.47	0.47
3:PC:79:ILE:HG22	3:PC:204:PHE:HB3	1.95	0.47
2:UC:298:LEU:HB2	2:UC:302:GLU:OE2	2.15	0.47
3:BD:229:GLU:OE1	3:BD:230:LEU:N	2.48	0.47
2:GD:195:GLY:HA2	2:MD:161:ILE:O	2.14	0.47
2:GD:267:GLU:HB3	2:GD:269:PRO:HD2	1.96	0.47
2:GD:298:LEU:HB2	2:GD:302:GLU:OE2	2.15	0.47
2:MD:207:LYS:CG	2:MD:210:GLN:HG2	2.39	0.47
2:SD:33:GLU:O	2:SD:37:LEU:HG	2.14	0.47
2:SD:119:ALA:HB1	2:SD:149:PHE:CE1	2.50	0.47
2:YD:65:GLN:HE21	2:EE:47:ILE:H	1.62	0.47
2:KE:8:ASP:O	2:KE:12:ILE:HG12	2.15	0.47
2:QE:84:ALA:O	2:QE:87:GLU:HB3	2.15	0.47
2:WE:38:SER:OG	2:WE:39:THR:N	2.47	0.47
2:WE:133:ILE:O	2:WE:137:LEU:HG	2.14	0.47
2:WE:167:VAL:HG13	3:XE:130:LEU:O	2.14	0.47
2:IF:128:GLN:HG2	3:JF:131:PHE:CB	2.36	0.47
2:UF:174:GLU:HG2	3:VF:152:ARG:HG2	1.97	0.47
2:GG:298:LEU:HB2	2:GG:302:GLU:OE2	2.15	0.47
3:NG:117:VAL:O	3:NG:215:PHE:HB2	2.15	0.47
2:G:76:TYR:HB2	2:B:41:MET:HE1	1.97	0.47
3:M:234:PRO:O	3:TG:190:ASN:ND2	2.47	0.47

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:B:85:LEU:H	2:B:85:LEU:HD22	1.80	0.47
2:B:178:VAL:HG11	3:C:149:THR:HG22	1.97	0.47
3:BA:117:VAL:O	3:BA:215:PHE:HB2	2.14	0.47
3:BA:185:GLN:HG3	3:BA:186:VAL:H	1.79	0.47
2:SA:108:GLU:HG3	2:SA:112:PHE:HE2	1.79	0.47
2:EB:14:LEU:HD12	2:EB:22:ALA:HB1	1.97	0.47
3:FB:69:LEU:O	3:FB:73:LEU:HB2	2.14	0.47
2:KB:48:SER:OG	2:KB:51:GLN:OE1	2.32	0.47
4:MB:65:THR:O	4:MB:99:ILE:HA	2.15	0.47
3:RB:202:THR:O	3:RB:202:THR:OG1	2.33	0.47
2:WB:15:MET:SD	2:WB:15:MET:N	2.88	0.47
2:WB:34:VAL:O	2:WB:37:LEU:HD12	2.15	0.47
2:CC:62:GLU:OE2	2:IC:45:ARG:NH1	2.48	0.47
3:DC:106:HIS:HE1	3:DC:113:THR:CG2	2.27	0.47
2:OC:126:HIS:CE1	3:PC:127:VAL:HG13	2.50	0.47
2:UC:89:ARG:HH22	2:UC:93:LEU:CD2	2.27	0.47
2:AD:98:LEU:HB3	2:AD:108:GLU:OE1	2.15	0.47
2:GD:141:GLN:HA	2:GD:144:ASP:OD2	2.15	0.47
2:MD:30:SER:O	2:MD:34:VAL:HG23	2.14	0.47
2:MD:261:ILE:O	2:MD:264:LYS:HG3	2.14	0.47
3:ZD:122:LEU:CD1	3:ZD:215:PHE:HB3	2.41	0.47
3:ZD:169:TRP:CD1	3:ZD:175:LEU:HD13	2.49	0.47
3:ZD:185:GLN:HG3	3:ZD:186:VAL:H	1.79	0.47
2:EE:30:SER:O	2:EE:34:VAL:HG23	2.15	0.47
2:EE:75:GLU:O	2:EE:79:SER:HB3	2.14	0.47
3:FE:169:TRP:CD1	3:FE:175:LEU:HD13	2.47	0.47
3:RE:135:GLY:O	3:RE:211:LEU:HD21	2.15	0.47
2:CF:84:ALA:O	2:CF:87:GLU:HB3	2.14	0.47
2:IF:53:THR:O	2:IF:57:SER:OG	2.28	0.47
2:UF:159:LEU:HD21	3:VF:137:PHE:CD1	2.49	0.47
2:AG:73:ALA:O	2:AG:76:TYR:HB3	2.14	0.47
2:GG:119:ALA:HB1	2:GG:149:PHE:CE1	2.50	0.47
3:HG:190:ASN:O	3:NG:236:LEU:HD12	2.15	0.47
2:MG:107:ILE:HD12	2:MG:107:ILE:N	2.29	0.47
2:J:72:ASN:ND2	2:T:41:MET:HB2	2.30	0.47
2:J:198:THR:O	2:J:202:ILE:HG12	2.14	0.47
2:T:105:SER:OG	2:T:107:ILE:HD11	2.15	0.47
2:T:198:THR:CG2	2:AA:139:ARG:HH22	2.28	0.47
3:V:66:ARG:HG3	3:V:77:PRO:O	2.15	0.47
2:AA:128:GLN:CG	3:BA:131:PHE:HB3	2.35	0.47
2:MA:48:SER:OG	2:MA:51:GLN:OE1	2.30	0.47

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:MA:152:ARG:HH22	2:MA:153:LEU:HB2	1.80	0.47
3:NA:185:GLN:HG3	3:NA:186:VAL:H	1.80	0.47
2:SA:178:VAL:HG11	3:TA:149:THR:HG22	1.97	0.47
3:TA:240:ARG:HA	3:TA:240:ARG:CZ	2.45	0.47
2:YA:156:ASP:HA	2:YA:159:LEU:HD12	1.96	0.47
3:ZA:110:LEU:HG	3:ZA:223:MET:HB3	1.97	0.47
2:EB:221:PHE:CE2	2:KB:159:LEU:HD11	2.50	0.47
2:KB:37:LEU:HA	2:KB:40:ALA:HB3	1.97	0.47
3:DC:65:PHE:CE1	3:DC:161:ALA:HB2	2.50	0.47
3:DC:183:GLU:OE2	3:DC:184:MET:N	2.48	0.47
2:IC:167:VAL:HG21	3:JC:131:PHE:CD1	2.49	0.47
3:JC:65:PHE:CE1	3:JC:161:ALA:HB2	2.50	0.47
3:VC:239:SER:O	3:VC:242:GLU:HG2	2.15	0.47
2:SD:83:LYS:HB3	2:YD:23:ALA:HA	1.97	0.47
2:SD:138:LYS:HD3	2:SD:138:LYS:HA	1.76	0.47
3:TD:101:ASN:HB2	3:TD:146:PHE:CE2	2.50	0.47
4:HE:65:THR:O	4:IE:99:ILE:HA	2.15	0.47
2:KE:79:SER:HA	2:KE:82:VAL:HG12	1.97	0.47
2:KE:99:GLU:OE2	2:KE:103:THR:HG21	2.15	0.47
3:LE:54:LEU:HD13	3:LE:231:LEU:HD21	1.97	0.47
3:LE:100:THR:HG23	3:LE:119:SER:HA	1.96	0.47
2:QE:172:LEU:HD12	2:QE:172:LEU:HA	1.66	0.47
2:CF:128:GLN:HG2	3:DF:131:PHE:CB	2.32	0.47
2:IF:245:ARG:O	2:IF:245:ARG:NH1	2.37	0.47
2:AG:12:ILE:HD11	2:AG:55:VAL:HG11	1.97	0.47
2:GG:83:LYS:HD2	2:MG:26:PHE:C	2.36	0.47
2:MG:188:LEU:HD13	3:TG:74:ARG:CD	2.42	0.47
3:NG:58:ASN:OD1	3:NG:81:VAL:HG11	2.13	0.47
3:NG:190:ASN:OD1	3:TG:236:LEU:HB2	2.15	0.47
2:SG:267:GLU:HB3	2:SG:269:PRO:HD2	1.96	0.47
3:M:247:ARG:NH1	3:M:251:VAL:HG23	2.30	0.46
2:T:133:ILE:O	2:T:137:LEU:HG	2.15	0.46
2:GA:149:PHE:CG	2:GA:153:LEU:HD13	2.50	0.46
2:GA:209:GLN:O	2:GA:212:GLU:HG3	2.15	0.46
2:SA:100:THR:OG1	2:SA:101:ARG:NH2	2.47	0.46
2:SA:109:THR:O	2:SA:113:MET:HG3	2.16	0.46
2:YA:53:THR:O	2:YA:57:SER:OG	2.30	0.46
2:YA:69:LEU:HD21	2:EB:47:ILE:HB	1.97	0.46
3:RB:169:TRP:CD1	3:RB:175:LEU:HD13	2.50	0.46
3:XB:111:ARG:HA	3:XB:111:ARG:CZ	2.46	0.46
2:CC:8:ASP:O	2:CC:12:ILE:HG12	2.15	0.46

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:IC:81:LEU:O	2:IC:85:LEU:HD13	2.15	0.46
2:UC:30:SER:O	2:UC:34:VAL:HG23	2.16	0.46
2:UC:95:GLU:O	2:UC:99:GLU:HG2	2.15	0.46
2:AD:218:VAL:HG12	2:AD:226:ALA:HB2	1.97	0.46
2:GD:172:LEU:HD12	2:GD:172:LEU:HA	1.76	0.46
2:SD:34:VAL:O	2:SD:37:LEU:HD12	2.15	0.46
2:YD:83:LYS:HB2	2:EE:26:PHE:HB3	1.96	0.46
2:YD:186:GLN:HB3	3:FE:74:ARG:HH22	1.80	0.46
2:EE:174:GLU:HG2	3:FE:152:ARG:HG2	1.97	0.46
2:KE:127:PRO:O	2:KE:130:ILE:HG12	2.14	0.46
2:KE:178:VAL:HG11	3:LE:149:THR:HG22	1.97	0.46
2:WE:174:GLU:HG2	3:XE:152:ARG:HG2	1.96	0.46
2:CF:267:GLU:HB3	2:CF:269:PRO:HD2	1.96	0.46
2:IF:174:GLU:HG2	3:JF:152:ARG:HG2	1.96	0.46
3:JF:285:PRO:HA	4:KF:119:GLY:HA2	1.97	0.46
2:OF:76:TYR:HD2	2:UF:41:MET:CE	2.27	0.46
3:BG:162:LEU:HD13	3:BG:162:LEU:HA	1.78	0.46
3:HG:224:ILE:O	3:HG:228:ARG:N	2.48	0.46
3:NG:169:TRP:CD1	3:NG:175:LEU:HD13	2.46	0.46
2:B:168:GLN:HB2	2:B:169:PRO:HD2	1.97	0.46
2:MA:267:GLU:HB3	2:MA:269:PRO:HD2	1.96	0.46
2:KB:242:VAL:HG13	2:KB:323:ILE:HG12	1.97	0.46
2:KB:267:GLU:HB3	2:KB:269:PRO:HD2	1.96	0.46
3:RB:185:GLN:HG3	3:RB:186:VAL:H	1.80	0.46
3:XB:117:VAL:O	3:XB:215:PHE:HB2	2.15	0.46
2:IC:14:LEU:HD12	2:IC:22:ALA:HB1	1.96	0.46
2:IC:65:GLN:HE21	2:OC:47:ILE:H	1.63	0.46
3:JC:113:THR:OG1	3:JC:191:ILE:HG22	2.15	0.46
3:PC:54:LEU:HD13	3:PC:231:LEU:HD21	1.96	0.46
3:PC:190:ASN:OD1	3:VC:236:LEU:N	2.47	0.46
2:GD:20:ASP:OD1	2:GD:20:ASP:N	2.47	0.46
2:GD:149:PHE:CD2	2:GD:153:LEU:HD22	2.50	0.46
2:GD:198:THR:HG23	2:MD:139:ARG:HH22	1.81	0.46
2:SD:267:GLU:HB3	2:SD:269:PRO:HD2	1.96	0.46
3:TD:202:THR:OG1	3:TD:202:THR:O	2.33	0.46
2:QE:160:ARG:O	2:QE:164:PHE:CB	2.61	0.46
2:WE:100:THR:OG1	2:WE:101:ARG:NH2	2.48	0.46
2:CF:122:ILE:HB	2:CF:130:ILE:HG22	1.96	0.46
2:IF:205:LEU:HD23	2:IF:205:LEU:O	2.15	0.46
2:OF:53:THR:O	2:OF:57:SER:OG	2.29	0.46
2:AG:219:ARG:HH21	2:AG:230:ILE:HD12	1.80	0.46

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:HG:202:THR:O	3:HG:202:THR:OG1	2.32	0.46
2:MG:178:VAL:HG11	3:NG:149:THR:HG22	1.98	0.46
2:SG:114:GLU:HB2	2:SG:117:SER:OG	2.15	0.46
2:B:195:GLY:HA2	2:J:161:ILE:O	2.16	0.46
2:B:210:GLN:O	2:B:214:VAL:HG12	2.16	0.46
2:T:31:THR:O	2:T:34:VAL:HB	2.14	0.46
2:T:38:SER:HA	2:T:41:MET:HE1	1.97	0.46
2:T:128:GLN:NE2	2:T:164:PHE:HD2	2.11	0.46
2:T:188:LEU:HD13	3:BA:74:ARG:CD	2.43	0.46
3:V:58:ASN:OD1	3:V:81:VAL:HG11	2.14	0.46
3:BA:65:PHE:CE1	3:BA:161:ALA:HB2	2.51	0.46
2:GA:20:ASP:OD2	2:GA:21:ARG:N	2.46	0.46
2:MA:8:ASP:O	2:MA:12:ILE:HG12	2.14	0.46
2:MA:53:THR:O	2:MA:57:SER:OG	2.31	0.46
2:MA:203:ILE:HA	2:MA:206:MET:HE2	1.96	0.46
3:XB:252:ARG:HA	3:XB:252:ARG:HD2	1.67	0.46
2:OC:151:GLU:HA	2:OC:154:ARG:HG2	1.96	0.46
3:PC:117:VAL:O	3:PC:215:PHE:HB2	2.16	0.46
2:AD:14:LEU:HD12	2:AD:22:ALA:HB1	1.96	0.46
2:AD:115:PRO:O	2:AD:119:ALA:HB2	2.16	0.46
2:MD:15:MET:SD	2:MD:15:MET:N	2.88	0.46
2:MD:76:TYR:OH	2:SD:34:VAL:HG13	2.15	0.46
2:SD:8:ASP:O	2:SD:12:ILE:HG12	2.15	0.46
3:TD:65:PHE:CE1	3:TD:161:ALA:HB2	2.50	0.46
3:ZD:69:LEU:O	3:ZD:73:LEU:HB2	2.16	0.46
2:EE:267:GLU:HB3	2:EE:269:PRO:HD2	1.96	0.46
2:KE:131:ALA:HA	2:KE:134:LEU:HD12	1.97	0.46
3:LE:58:ASN:OD1	3:LE:81:VAL:HG11	2.14	0.46
3:XE:54:LEU:HB2	3:XE:231:LEU:HD23	1.95	0.46
2:CF:113:MET:CE	2:CF:121:LEU:HD11	2.45	0.46
2:CF:167:VAL:HG13	3:DF:130:LEU:O	2.15	0.46
2:CF:183:LEU:O	2:CF:186:GLN:N	2.49	0.46
3:DF:150:GLU:O	3:DF:153:VAL:HG12	2.15	0.46
2:IF:65:GLN:NE2	2:OF:47:ILE:O	2.45	0.46
2:UF:65:GLN:HE21	2:AG:47:ILE:H	1.63	0.46
2:UF:69:LEU:HD11	2:AG:47:ILE:CG2	2.45	0.46
2:UF:149:PHE:CD2	2:UF:153:LEU:HD22	2.50	0.46
3:BG:239:SER:O	3:BG:242:GLU:HG2	2.15	0.46
3:HG:183:GLU:HG3	3:HG:188:PHE:HB2	1.97	0.46
2:SG:89:ARG:CD	2:SG:90:ALA:H	2.28	0.46
3:TG:285:PRO:HA	4:UG:119:GLY:HA2	1.97	0.46

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:M:62:ALA:O	3:M:79:ILE:HD12	2.16	0.46
3:M:229:GLU:OE1	3:M:230:LEU:N	2.49	0.46
3:C:65:PHE:CE1	3:C:161:ALA:HB2	2.50	0.46
2:J:128:GLN:HG2	3:K:131:PHE:CB	2.33	0.46
2:AA:67:ALA:O	2:AA:71:ILE:HG12	2.14	0.46
2:AA:282:ALA:HA	2:AA:285:ILE:HG12	1.98	0.46
2:MA:76:TYR:HE2	2:SA:38:SER:CB	2.25	0.46
2:MA:127:PRO:O	2:MA:130:ILE:HG12	2.15	0.46
3:NA:190:ASN:O	3:NA:190:ASN:ND2	2.48	0.46
2:SA:35:GLN:O	2:SA:38:SER:OG	2.31	0.46
2:YA:174:GLU:HG2	3:ZA:152:ARG:HG2	1.97	0.46
2:EB:26:PHE:CE1	2:EB:34:VAL:HG22	2.50	0.46
2:EB:267:GLU:HB3	2:EB:269:PRO:HD2	1.96	0.46
2:KB:33:GLU:O	2:KB:37:LEU:HG	2.14	0.46
2:KB:63:ALA:HA	2:KB:66:PHE:CZ	2.51	0.46
2:KB:76:TYR:HE1	2:QB:26:PHE:CE2	2.34	0.46
2:QB:115:PRO:O	2:QB:119:ALA:CB	2.64	0.46
2:QB:149:PHE:CD2	2:QB:153:LEU:HD22	2.50	0.46
2:CC:201:GLU:CG	2:IC:139:ARG:HH12	2.29	0.46
3:DC:236:LEU:HB3	3:DC:238:ASN:HD21	1.79	0.46
2:OC:210:GLN:O	2:OC:214:VAL:HG12	2.15	0.46
2:UC:193:MET:O	2:UC:195:GLY:N	2.49	0.46
2:GD:89:ARG:HG3	2:GD:90:ALA:N	2.30	0.46
2:GD:151:GLU:HA	2:GD:154:ARG:HE	1.79	0.46
2:GD:206:MET:SD	2:GD:210:GLN:HB2	2.55	0.46
4:OD:101:ILE:N	4:OD:104:TYR:O	2.36	0.46
2:SD:101:ARG:O	2:SD:104:THR:HG23	2.15	0.46
2:YD:122:ILE:HB	2:YD:130:ILE:HG22	1.97	0.46
3:FE:185:GLN:HG3	3:FE:186:VAL:H	1.81	0.46
2:QE:111:ASN:HA	2:QE:141:GLN:OE1	2.16	0.46
3:RE:190:ASN:OD1	3:XE:236:LEU:HB2	2.16	0.46
3:XE:285:PRO:HA	4:YE:119:GLY:HA2	1.98	0.46
2:UF:71:ILE:O	2:UF:74:ASN:ND2	2.49	0.46
2:AG:221:PHE:CE2	2:GG:159:LEU:HD21	2.49	0.46
2:GG:66:PHE:HA	2:MG:45:ARG:O	2.15	0.46
2:MG:84:ALA:O	2:MG:87:GLU:HB3	2.14	0.46
2:SG:298:LEU:HB2	2:SG:302:GLU:OE2	2.15	0.46
3:TG:238:ASN:HB3	3:TG:241:HIS:ND1	2.30	0.46
2:G:167:VAL:HG13	3:M:130:LEU:O	2.14	0.46
2:B:69:LEU:HD13	2:J:45:ARG:O	2.16	0.46
2:B:101:ARG:O	2:B:104:THR:HG23	2.15	0.46

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:B:117:SER:O	2:B:121:LEU:HG	2.15	0.46
4:CA:65:THR:O	4:CA:99:ILE:HA	2.15	0.46
2:SA:115:PRO:O	2:SA:119:ALA:HB3	2.15	0.46
3:TA:106:HIS:CD2	3:TA:113:THR:HG22	2.51	0.46
2:YA:267:GLU:HB3	2:YA:269:PRO:HD2	1.96	0.46
2:KB:167:VAL:HG21	3:LB:131:PHE:CD1	2.50	0.46
3:LB:86:ILE:HD11	3:LB:221:PHE:CE1	2.50	0.46
2:QB:119:ALA:HB1	2:QB:149:PHE:CE1	2.51	0.46
2:WB:33:GLU:O	2:WB:37:LEU:HG	2.16	0.46
2:WB:210:GLN:O	2:WB:214:VAL:HG12	2.16	0.46
3:XB:97:PRO:HD2	3:XB:100:THR:OG1	2.14	0.46
2:CC:178:VAL:HG11	3:DC:149:THR:HG22	1.97	0.46
2:IC:167:VAL:HG13	3:JC:130:LEU:O	2.15	0.46
2:UC:84:ALA:O	2:UC:87:GLU:HB3	2.15	0.46
2:AD:30:SER:O	2:AD:34:VAL:HG23	2.16	0.46
2:GD:178:VAL:HG11	3:HD:149:THR:HG22	1.96	0.46
2:MD:282:ALA:HA	2:MD:285:ILE:HG12	1.97	0.46
2:SD:15:MET:N	2:SD:15:MET:SD	2.87	0.46
2:YD:206:MET:SD	2:YD:210:GLN:HB2	2.56	0.46
2:EE:178:VAL:HG11	3:FE:149:THR:HG22	1.97	0.46
3:FE:106:HIS:CD2	3:FE:113:THR:HG22	2.50	0.46
2:KE:30:SER:O	2:KE:34:VAL:HG23	2.15	0.46
3:LE:285:PRO:HA	4:ME:119:GLY:HA2	1.98	0.46
3:XE:117:VAL:O	3:XE:215:PHE:HB2	2.16	0.46
2:OF:187:ASN:O	2:OF:189:LYS:NZ	2.45	0.46
3:PF:185:GLN:HG3	3:PF:186:VAL:H	1.80	0.46
3:PF:202:THR:HG1	3:PF:204:PHE:HE1	1.61	0.46
2:AG:267:GLU:HB3	2:AG:269:PRO:HD2	1.96	0.46
3:BG:101:ASN:HB2	3:BG:146:PHE:CE2	2.51	0.46
3:NG:65:PHE:CE1	3:NG:161:ALA:HB2	2.51	0.46
2:G:103:THR:C	2:G:106:GLY:H	2.18	0.46
3:M:236:LEU:HD13	3:TG:190:ASN:O	2.15	0.46
3:M:240:ARG:HA	3:M:240:ARG:CZ	2.46	0.46
2:B:105:SER:OG	2:B:107:ILE:HD11	2.15	0.46
2:B:267:GLU:HB3	2:B:269:PRO:HD2	1.96	0.46
3:C:225:GLU:HA	3:C:228:ARG:HB3	1.98	0.46
4:D:65:THR:O	4:D:99:ILE:HA	2.16	0.46
2:T:100:THR:OG1	2:T:101:ARG:NH2	2.49	0.46
3:HA:198:ILE:HD12	3:HA:199:VAL:N	2.31	0.46
2:SA:131:ALA:HA	2:SA:134:LEU:HD12	1.97	0.46
3:TA:101:ASN:HB2	3:TA:146:PHE:CE1	2.50	0.46

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:YA:76:TYR:HB2	2:EB:41:MET:HE3	1.97	0.46
4:OB:107:ALA:H	3:RB:252:ARG:NH2	2.14	0.46
2:QB:76:TYR:HB2	2:WB:41:MET:HE3	1.98	0.46
2:QB:80:VAL:HG11	2:QB:93:LEU:HD12	1.97	0.46
2:WB:183:LEU:O	2:WB:186:GLN:N	2.48	0.46
3:DC:154:ILE:HD13	3:DC:154:ILE:HA	1.79	0.46
2:UC:48:SER:OG	2:UC:51:GLN:OE1	2.28	0.46
3:VC:122:LEU:CD1	3:VC:215:PHE:HB3	2.44	0.46
4:YC:107:ALA:H	3:BD:252:ARG:HH21	1.63	0.46
2:AD:65:GLN:HE21	2:GD:47:ILE:H	1.61	0.46
2:YD:20:ASP:O	2:YD:23:ALA:HB3	2.16	0.46
2:YD:267:GLU:HB3	2:YD:269:PRO:HD2	1.96	0.46
4:BE:65:THR:O	4:CE:99:ILE:HA	2.16	0.46
2:EE:195:GLY:HA2	2:KE:161:ILE:O	2.16	0.46
2:WE:12:ILE:HD11	2:WE:55:VAL:HG11	1.97	0.46
4:KF:65:THR:O	4:KF:99:ILE:HA	2.16	0.46
2:OF:190:ARG:O	2:UF:166:GLY:HA3	2.15	0.46
3:VF:183:GLU:HG3	3:VF:188:PHE:HB2	1.96	0.46
3:VF:267:ASP:HA	4:WF:65:THR:HA	1.98	0.46
2:AG:107:ILE:O	2:AG:111:ASN:ND2	2.49	0.46
2:AG:190:ARG:O	2:GG:166:GLY:HA3	2.15	0.46
3:BG:267:ASP:HA	4:CG:65:THR:HA	1.98	0.46
3:HG:75:ARG:HH22	3:HG:209:GLY:HA2	1.79	0.46
2:MG:30:SER:O	2:MG:34:VAL:HG23	2.15	0.46
2:SG:8:ASP:OD1	2:SG:47:ILE:HD11	2.15	0.46
2:B:12:ILE:HD11	2:B:55:VAL:HG11	1.98	0.46
3:C:58:ASN:OD1	3:C:81:VAL:HG11	2.16	0.46
2:J:8:ASP:O	2:J:12:ILE:HG12	2.15	0.46
2:J:174:GLU:HG2	3:K:152:ARG:HG2	1.97	0.46
2:J:263:LEU:HD11	2:J:274:PHE:HD2	1.81	0.46
3:V:101:ASN:HB2	3:V:146:PHE:CE2	2.51	0.46
3:V:116:VAL:HG12	3:V:217:ILE:HG13	1.97	0.46
2:GA:282:ALA:HA	2:GA:285:ILE:HG12	1.98	0.46
2:SA:242:VAL:HG13	2:SA:323:ILE:HG12	1.97	0.46
3:TA:60:ARG:HE	3:TA:60:ARG:HB2	1.57	0.46
2:KB:83:LYS:CG	2:QB:26:PHE:HB3	2.37	0.46
3:RB:247:ARG:O	3:RB:250:LEU:HB3	2.16	0.46
2:WB:32:ARG:NE	2:WB:33:GLU:OE2	2.48	0.46
3:XB:267:ASP:HA	4:YB:65:THR:HA	1.98	0.46
2:CC:71:ILE:O	2:CC:74:ASN:ND2	2.49	0.46
3:PC:185:GLN:HG3	3:PC:186:VAL:H	1.81	0.46

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:UC:198:THR:HG23	2:UC:201:GLU:OE1	2.16	0.46
3:VC:285:PRO:HA	4:WC:119:GLY:HA2	1.98	0.46
2:AD:84:ALA:O	2:AD:87:GLU:HB3	2.15	0.46
2:GD:15:MET:N	2:GD:15:MET:SD	2.89	0.46
2:GD:100:THR:OG1	2:GD:101:ARG:NH2	2.48	0.46
2:MD:126:HIS:CE1	3:ND:127:VAL:HG13	2.50	0.46
3:TD:174:PRO:HD2	3:TD:174:PRO:O	2.15	0.46
2:YD:20:ASP:OD1	2:YD:20:ASP:N	2.49	0.46
4:AE:65:THR:O	4:AE:99:ILE:HA	2.16	0.46
2:EE:84:ALA:O	2:EE:87:GLU:HB3	2.16	0.46
2:KE:267:GLU:HB3	2:KE:269:PRO:HD2	1.96	0.46
3:LE:214:GLU:OE1	3:LE:215:PHE:N	2.49	0.46
2:QE:110:LEU:O	2:QE:141:GLN:NE2	2.46	0.46
2:WE:206:MET:SD	2:WE:207:LYS:HG2	2.56	0.46
3:XE:183:GLU:HG3	3:XE:188:PHE:HB2	1.97	0.46
2:IF:267:GLU:HB3	2:IF:269:PRO:HD2	1.96	0.46
2:OF:79:SER:C	2:OF:83:LYS:HE3	2.36	0.46
2:MG:149:PHE:CD2	2:MG:153:LEU:HD22	2.51	0.46
2:MG:196:VAL:HG12	2:MG:232:GLU:OE2	2.15	0.46
2:SG:53:THR:O	2:SG:57:SER:OG	2.31	0.46
2:G:119:ALA:HB1	2:G:149:PHE:CE1	2.50	0.46
2:G:152:ARG:HH22	2:G:153:LEU:HB2	1.80	0.46
2:B:168:GLN:NE2	3:C:130:LEU:HD11	2.24	0.46
2:B:282:ALA:HA	2:B:285:ILE:HG12	1.98	0.46
3:K:115:LEU:HD11	3:K:218:CYS:HB3	1.97	0.46
2:AA:158:MET:SD	2:AA:159:LEU:HD23	2.56	0.46
2:GA:186:GLN:C	3:NA:74:ARG:HH22	2.20	0.46
3:HA:100:THR:HG23	3:HA:119:SER:HA	1.98	0.46
2:MA:115:PRO:O	2:MA:119:ALA:HB3	2.16	0.46
2:YA:133:ILE:O	2:YA:137:LEU:HG	2.16	0.46
4:AB:101:ILE:N	4:AB:104:TYR:O	2.33	0.46
2:QB:282:ALA:HA	2:QB:285:ILE:HG12	1.98	0.46
2:CC:201:GLU:HG2	2:IC:139:ARG:NH1	2.31	0.46
2:OC:74:ASN:O	2:OC:77:LEU:HB3	2.15	0.46
3:BD:75:ARG:HG3	3:BD:208:ILE:HD11	1.97	0.46
2:EE:78:ARG:O	2:EE:82:VAL:HG23	2.16	0.46
2:KE:11:VAL:O	2:KE:14:LEU:HB3	2.16	0.46
2:KE:160:ARG:O	2:KE:164:PHE:CB	2.61	0.46
2:QE:203:ILE:HA	2:QE:206:MET:HE2	1.98	0.46
2:QE:267:GLU:HB3	2:QE:269:PRO:HD2	1.96	0.46
4:AF:105:LEU:O	3:DF:252:ARG:HD3	2.15	0.46

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:IF:188:LEU:N	3:PF:74:ARG:HH21	2.13	0.46
2:UF:133:ILE:O	2:UF:137:LEU:HG	2.16	0.46
2:AG:204:ASN:HB3	2:AG:205:LEU:HD12	1.98	0.46
3:BG:97:PRO:HD2	3:BG:100:THR:OG1	2.16	0.46
2:G:307:ALA:O	2:G:310:LEU:HG	2.16	0.46
2:B:30:SER:O	2:B:34:VAL:HG23	2.16	0.46
2:J:74:ASN:O	2:J:78:ARG:NE	2.48	0.46
3:K:190:ASN:ND2	3:V:234:PRO:O	2.48	0.46
2:T:41:MET:HG2	2:T:42:ALA:N	2.31	0.46
3:V:202:THR:OG1	3:V:217:ILE:HG22	2.16	0.46
2:AA:152:ARG:NH2	2:AA:153:LEU:HB2	2.31	0.46
2:AA:267:GLU:HB3	2:AA:269:PRO:HD2	1.96	0.46
2:SA:115:PRO:O	2:SA:119:ALA:HB2	2.15	0.46
3:TA:185:GLN:HG2	3:TA:187:LYS:HZ2	1.81	0.46
2:EB:172:LEU:HA	2:EB:172:LEU:HD12	1.72	0.46
2:KB:128:GLN:CG	3:LB:131:PHE:HB3	2.36	0.46
2:QB:178:VAL:HG11	3:RB:149:THR:HG22	1.98	0.46
2:QB:242:VAL:HG13	2:QB:323:ILE:HG12	1.97	0.46
3:RB:110:LEU:HG	3:RB:223:MET:HB3	1.98	0.46
2:CC:58:GLU:O	2:CC:62:GLU:HB2	2.16	0.46
3:DC:110:LEU:HG	3:DC:223:MET:HB3	1.96	0.46
2:IC:89:ARG:HG3	2:IC:89:ARG:NH1	2.31	0.46
2:AD:66:PHE:HA	2:GD:45:ARG:O	2.16	0.46
2:AD:89:ARG:HG3	2:AD:89:ARG:NH1	2.30	0.46
2:GD:79:SER:CB	2:GD:83:LYS:HE2	2.46	0.46
3:HD:123:VAL:HG21	3:HD:146:PHE:CZ	2.51	0.46
4:ID:65:THR:O	4:ID:99:ILE:HA	2.15	0.46
3:TD:59:GLU:N	3:TD:59:GLU:OE2	2.48	0.46
3:TD:117:VAL:O	3:TD:215:PHE:HB2	2.16	0.46
3:ZD:58:ASN:OD1	3:ZD:81:VAL:HG11	2.16	0.46
3:ZD:294:ALA:O	3:ZD:301:VAL:N	2.49	0.46
2:KE:174:GLU:HG2	3:LE:152:ARG:HG2	1.97	0.46
2:QE:30:SER:O	2:QE:34:VAL:HG23	2.16	0.46
3:XE:185:GLN:HG3	3:XE:186:VAL:H	1.81	0.46
2:CF:167:VAL:HG21	3:DF:131:PHE:CD1	2.51	0.46
3:JF:51:LEU:O	3:JF:55:GLU:HG3	2.16	0.46
3:JF:106:HIS:HE1	3:JF:113:THR:CG2	2.28	0.46
2:OF:282:ALA:HA	2:OF:285:ILE:HG12	1.98	0.46
3:PF:169:TRP:CD1	3:PF:175:LEU:HD13	2.49	0.46
2:UF:81:LEU:H	2:UF:81:LEU:HD12	1.80	0.46
2:AG:107:ILE:HG22	2:AG:111:ASN:HD21	1.81	0.46

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:AG:117:SER:O	2:AG:121:LEU:HG	2.15	0.46
2:GG:188:LEU:HD22	3:NG:74:ARG:NH2	2.31	0.46
3:NG:134:ASP:OD2	3:NG:136:ARG:N	2.40	0.46
2:G:12:ILE:HD11	2:G:55:VAL:HG11	1.97	0.46
2:G:160:ARG:O	2:G:164:PHE:CB	2.61	0.46
2:B:201:GLU:HB3	2:J:139:ARG:NH2	2.21	0.46
2:J:110:LEU:HB3	2:J:141:GLN:HE22	1.81	0.46
4:W:65:THR:O	4:W:99:ILE:HA	2.16	0.46
2:GA:210:GLN:O	2:GA:214:VAL:HG12	2.16	0.46
3:HA:117:VAL:O	3:HA:215:PHE:HB2	2.16	0.46
3:HA:185:GLN:HG2	3:HA:187:LYS:HZ2	1.80	0.46
2:MA:126:HIS:ND1	2:MA:127:PRO:HD2	2.31	0.46
3:NA:90:HIS:O	3:NA:94:ARG:HD2	2.15	0.46
2:SA:267:GLU:HB3	2:SA:269:PRO:HD2	1.96	0.46
3:TA:195:PRO:HD2	3:TA:196:ASN:N	2.31	0.46
2:YA:101:ARG:O	2:YA:104:THR:HG23	2.16	0.46
2:YA:120:ASP:OD1	2:YA:123:ARG:NH1	2.49	0.46
2:EB:65:GLN:HE21	2:KB:47:ILE:H	1.64	0.46
3:RB:180:VAL:HG23	3:RB:181:ARG:N	2.31	0.46
3:DC:267:ASP:HA	4:EC:65:THR:HA	1.98	0.46
2:OC:282:ALA:HA	2:OC:285:ILE:HG12	1.98	0.46
3:PC:135:GLY:O	3:PC:211:LEU:HD21	2.15	0.46
2:UC:204:ASN:OD1	2:UC:233:MET:HB3	2.15	0.46
3:BD:110:LEU:HG	3:BD:223:MET:HB3	1.98	0.46
2:GD:83:LYS:CB	2:MD:23:ALA:HA	2.46	0.46
2:GD:108:GLU:HA	2:GD:111:ASN:HD22	1.81	0.46
2:MD:167:VAL:HG21	3:ND:131:PHE:CD1	2.51	0.46
2:MD:174:GLU:HG2	3:ND:152:ARG:HG2	1.97	0.46
2:MD:207:LYS:NZ	2:MD:209:GLN:HB3	2.31	0.46
2:SD:87:GLU:HG2	2:SD:89:ARG:H	1.81	0.46
3:TD:58:ASN:OD1	3:TD:81:VAL:HG11	2.16	0.46
1:XD:543:ASP:HA	2:YD:21:ARG:HH11	1.81	0.46
2:WE:128:GLN:HG2	3:XE:131:PHE:CB	2.37	0.46
2:WE:128:GLN:NE2	2:WE:164:PHE:HD2	2.07	0.46
2:WE:130:ILE:HD11	2:WE:157:VAL:HG22	1.98	0.46
2:WE:267:GLU:HB3	2:WE:269:PRO:HD2	1.96	0.46
2:CF:83:LYS:HG3	2:CF:84:ALA:N	2.29	0.46
3:DF:285:PRO:HA	4:EF:119:GLY:HA2	1.98	0.46
2:IF:22:ALA:O	2:IF:26:PHE:HB2	2.16	0.46
2:IF:109:THR:O	2:IF:113:MET:HG3	2.16	0.46
2:IF:133:ILE:O	2:IF:137:LEU:HG	2.16	0.46

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:VF:69:LEU:O	3:VF:73:LEU:HB2	2.16	0.46
3:VF:101:ASN:HB2	3:VF:146:PHE:HE2	1.81	0.46
3:BG:154:ILE:HD13	3:BG:154:ILE:HA	1.78	0.46
2:GG:30:SER:OG	2:GG:33:GLU:OE1	2.28	0.46
2:MG:31:THR:O	2:MG:34:VAL:HB	2.15	0.46
2:G:198:THR:O	2:G:201:GLU:HG2	2.16	0.45
2:T:107:ILE:HD12	2:T:107:ILE:N	2.30	0.45
3:V:202:THR:OG1	3:V:202:THR:O	2.33	0.45
2:AA:33:GLU:O	2:AA:37:LEU:HG	2.15	0.45
2:AA:210:GLN:O	2:AA:214:VAL:HG12	2.16	0.45
3:BA:101:ASN:HB2	3:BA:146:PHE:CE1	2.51	0.45
2:GA:159:LEU:HD12	2:GA:159:LEU:HA	1.82	0.45
2:MA:107:ILE:O	2:MA:111:ASN:ND2	2.49	0.45
2:MA:178:VAL:HG11	3:NA:149:THR:HG22	1.98	0.45
2:SA:149:PHE:CD2	2:SA:153:LEU:HD22	2.51	0.45
3:TA:55:GLU:HA	3:TA:58:ASN:ND2	2.30	0.45
3:ZA:123:VAL:HG21	3:ZA:146:PHE:CZ	2.51	0.45
3:ZA:134:ASP:OD1	3:ZA:136:ARG:N	2.37	0.45
3:ZA:183:GLU:HG3	3:ZA:188:PHE:HB2	1.98	0.45
2:EB:87:GLU:HG2	2:EB:89:ARG:N	2.31	0.45
3:FB:195:PRO:HD2	3:FB:196:ASN:N	2.31	0.45
2:QB:76:TYR:HE2	2:WB:38:SER:CB	2.29	0.45
3:RB:101:ASN:HB2	3:RB:146:PHE:CE2	2.51	0.45
2:WB:69:LEU:HD11	2:CC:47:ILE:CG2	2.46	0.45
3:XB:97:PRO:O	3:XB:97:PRO:HD2	2.16	0.45
2:CC:26:PHE:CZ	2:CC:34:VAL:HG22	2.51	0.45
2:CC:89:ARG:CD	2:CC:90:ALA:H	2.29	0.45
3:JC:66:ARG:HG3	3:JC:77:PRO:O	2.16	0.45
2:OC:107:ILE:HG22	2:OC:111:ASN:ND2	2.29	0.45
2:OC:144:ASP:O	2:OC:148:LEU:HD12	2.16	0.45
2:UC:120:ASP:OD1	2:UC:123:ARG:NH1	2.49	0.45
3:VC:185:GLN:HG3	3:VC:186:VAL:H	1.81	0.45
2:GD:174:GLU:HG2	3:HD:152:ARG:HG2	1.98	0.45
3:ND:79:ILE:HG22	3:ND:204:PHE:HB3	1.97	0.45
3:ND:236:LEU:HB3	3:ND:238:ASN:OD1	2.16	0.45
2:YD:37:LEU:HD12	2:YD:38:SER:H	1.81	0.45
3:ZD:73:LEU:HD23	3:ZD:208:ILE:HD12	1.98	0.45
3:FE:59:GLU:N	3:FE:59:GLU:OE2	2.48	0.45
4:IE:107:ALA:O	3:LE:252:ARG:NH1	2.49	0.45
2:KE:15:MET:N	2:KE:15:MET:SD	2.88	0.45
3:RE:65:PHE:CE1	3:RE:161:ALA:HB2	2.51	0.45

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:CF:174:GLU:HG2	3:DF:152:ARG:HG2	1.98	0.45
2:IF:26:PHE:CE2	2:IF:37:LEU:HD11	2.51	0.45
2:IF:128:GLN:CG	3:JF:131:PHE:HB3	2.35	0.45
2:IF:242:VAL:HG13	2:IF:323:ILE:HG12	1.97	0.45
3:JF:86:ILE:HD12	3:JF:200:VAL:HG23	1.97	0.45
2:OF:307:ALA:O	2:OF:310:LEU:HG	2.16	0.45
2:UF:84:ALA:O	2:UF:87:GLU:HB3	2.16	0.45
2:UF:167:VAL:HG13	3:VF:130:LEU:O	2.16	0.45
2:AG:7:THR:HG23	2:AG:37:LEU:HD23	1.97	0.45
2:GG:26:PHE:CZ	2:GG:34:VAL:HG13	2.51	0.45
2:GG:30:SER:O	2:GG:34:VAL:HG23	2.16	0.45
3:M:190:ASN:OD1	3:C:236:LEU:HB2	2.17	0.45
2:J:210:GLN:O	2:J:214:VAL:HG12	2.17	0.45
2:T:119:ALA:HB1	2:T:149:PHE:CE1	2.51	0.45
3:V:60:ARG:HE	3:V:60:ARG:HB2	1.60	0.45
3:BA:69:LEU:O	3:BA:73:LEU:HB2	2.17	0.45
3:NA:189:THR:OG1	3:NA:190:ASN:N	2.50	0.45
2:SA:53:THR:O	2:SA:57:SER:OG	2.31	0.45
3:TA:185:GLN:HG3	3:TA:186:VAL:H	1.81	0.45
2:EB:188:LEU:HD22	3:LB:74:ARG:NH1	2.31	0.45
3:XB:135:GLY:O	3:XB:211:LEU:HD21	2.15	0.45
3:XB:237:GLU:HG2	3:XB:238:ASN:N	2.31	0.45
4:AC:105:LEU:O	3:DC:252:ARG:HD3	2.16	0.45
2:UC:14:LEU:HD12	2:UC:22:ALA:HA	1.97	0.45
2:UC:195:GLY:HA2	2:AD:161:ILE:O	2.16	0.45
4:XC:65:THR:O	4:YC:99:ILE:HA	2.17	0.45
3:BD:189:THR:OG1	3:BD:190:ASN:N	2.49	0.45
3:ND:59:GLU:OE2	3:ND:59:GLU:N	2.49	0.45
2:SD:110:LEU:HD13	2:SD:137:LEU:HD21	1.99	0.45
3:TD:185:GLN:HG3	3:TD:186:VAL:H	1.81	0.45
2:YD:282:ALA:HA	2:YD:285:ILE:HG12	1.98	0.45
2:KE:261:ILE:O	2:KE:264:LYS:HG3	2.17	0.45
2:QE:69:LEU:HD11	2:WE:47:ILE:CG2	2.39	0.45
2:QE:230:ILE:HA	2:QE:233:MET:HE1	1.99	0.45
2:WE:298:LEU:HB2	2:WE:302:GLU:OE1	2.15	0.45
2:CF:168:GLN:NE2	3:DF:130:LEU:HD11	2.31	0.45
2:IF:40:ALA:O	2:IF:44:VAL:HG23	2.16	0.45
2:IF:168:GLN:NE2	2:IF:171:ALA:H	2.14	0.45
3:JF:190:ASN:OD1	3:PF:236:LEU:HB2	2.14	0.45
2:UF:85:LEU:H	2:UF:85:LEU:HD22	1.81	0.45
2:GG:14:LEU:HD12	2:GG:22:ALA:HB1	1.98	0.45

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:MG:107:ILE:HG22	2:MG:111:ASN:ND2	2.29	0.45
2:SG:127:PRO:O	2:SG:130:ILE:HG12	2.15	0.45
3:TG:101:ASN:HB2	3:TG:146:PHE:CE1	2.51	0.45
3:M:60:ARG:HE	3:M:60:ARG:HB2	1.57	0.45
2:J:245:ARG:O	2:J:245:ARG:NH1	2.38	0.45
2:T:282:ALA:O	2:T:286:LEU:HD23	2.16	0.45
3:V:86:ILE:HD11	3:V:221:PHE:CE1	2.51	0.45
2:AA:66:PHE:HA	2:GA:45:ARG:O	2.17	0.45
4:EA:105:LEU:O	3:HA:252:ARG:HD3	2.17	0.45
2:SA:119:ALA:HB1	2:SA:149:PHE:CE1	2.51	0.45
2:YA:172:LEU:HD12	2:YA:172:LEU:HA	1.76	0.45
3:ZA:117:VAL:O	3:ZA:215:PHE:HB2	2.16	0.45
2:EB:263:LEU:HD11	2:EB:274:PHE:HD2	1.82	0.45
2:KB:141:GLN:HA	2:KB:144:ASP:OD2	2.15	0.45
4:OB:107:ALA:HA	4:OB:125:ILE:HA	1.97	0.45
2:QB:27:LYS:HG3	2:QB:27:LYS:O	2.15	0.45
2:QB:103:THR:C	2:QB:106:GLY:H	2.20	0.45
2:WB:115:PRO:O	2:WB:119:ALA:CB	2.63	0.45
2:CC:117:SER:O	2:CC:121:LEU:HG	2.15	0.45
2:CC:119:ALA:HB1	2:CC:149:PHE:CE1	2.52	0.45
2:CC:186:GLN:C	3:JC:74:ARG:HH22	2.20	0.45
2:IC:109:THR:O	2:IC:113:MET:HG3	2.17	0.45
2:OC:41:MET:SD	2:OC:42:ALA:N	2.89	0.45
2:OC:89:ARG:HG3	2:OC:89:ARG:NH1	2.31	0.45
2:OC:261:ILE:O	2:OC:264:LYS:HG3	2.16	0.45
3:PC:180:VAL:HG23	3:PC:181:ARG:H	1.82	0.45
2:UC:149:PHE:CD2	2:UC:153:LEU:HD22	2.52	0.45
3:BD:62:ALA:O	3:BD:79:ILE:HD12	2.17	0.45
3:BD:97:PRO:O	3:BD:100:THR:OG1	2.20	0.45
2:GD:261:ILE:H	2:GD:261:ILE:HG13	1.61	0.45
3:HD:117:VAL:O	3:HD:215:PHE:HB2	2.17	0.45
2:MD:85:LEU:H	2:MD:85:LEU:HD22	1.81	0.45
2:MD:128:GLN:CG	3:ND:131:PHE:HB3	2.33	0.45
3:ND:180:VAL:HG23	3:ND:181:ARG:H	1.81	0.45
2:YD:100:THR:OG1	2:YD:101:ARG:NH2	2.49	0.45
2:YD:167:VAL:HG21	3:ZD:131:PHE:CD1	2.52	0.45
2:EE:149:PHE:CG	2:EE:153:LEU:HD13	2.51	0.45
3:FE:285:PRO:HA	4:GE:119:GLY:HA2	1.99	0.45
2:KE:63:ALA:HA	2:KE:66:PHE:CZ	2.50	0.45
2:QE:31:THR:O	2:QE:34:VAL:HB	2.16	0.45
2:QE:190:ARG:O	2:WE:166:GLY:HA3	2.15	0.45

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:RE:127:VAL:HG23	3:RE:153:VAL:HG11	1.98	0.45
3:XE:86:ILE:HD11	3:XE:221:PHE:HE1	1.81	0.45
3:XE:101:ASN:HD22	3:XE:146:PHE:HE2	1.64	0.45
3:XE:236:LEU:HD23	3:XE:238:ASN:HD21	1.80	0.45
2:CF:168:GLN:HE22	3:DF:130:LEU:HD11	1.81	0.45
2:IF:69:LEU:O	2:IF:72:ASN:ND2	2.46	0.45
2:OF:105:SER:OG	2:OF:107:ILE:HD12	2.15	0.45
2:OF:149:PHE:CD2	2:OF:153:LEU:HD22	2.51	0.45
3:PF:285:PRO:HA	4:QF:119:GLY:HA2	1.99	0.45
2:AG:14:LEU:HD12	2:AG:22:ALA:HB1	1.97	0.45
2:AG:76:TYR:HE1	2:GG:26:PHE:CE1	2.34	0.45
3:HG:219:LEU:HD23	3:HG:219:LEU:HA	1.80	0.45
3:HG:239:SER:O	3:HG:242:GLU:HG2	2.16	0.45
3:NG:214:GLU:OE1	3:NG:215:PHE:N	2.50	0.45
2:SG:30:SER:O	2:SG:34:VAL:HG23	2.17	0.45
3:M:185:GLN:HG3	3:M:186:VAL:H	1.82	0.45
2:T:195:GLY:HA2	2:AA:161:ILE:O	2.16	0.45
3:V:185:GLN:HG3	3:V:186:VAL:H	1.81	0.45
2:MA:99:GLU:OE2	2:MA:103:THR:HG21	2.17	0.45
3:TA:180:VAL:HG23	3:TA:181:ARG:N	2.30	0.45
3:ZA:219:LEU:HD23	3:ZA:219:LEU:HA	1.81	0.45
3:FB:185:GLN:HG3	3:FB:186:VAL:H	1.82	0.45
2:KB:147:ALA:O	2:KB:154:ARG:NH1	2.49	0.45
3:LB:285:PRO:HA	4:MB:119:GLY:HA2	1.98	0.45
2:WB:151:GLU:HA	2:WB:154:ARG:HE	1.81	0.45
2:OC:7:THR:HG23	2:OC:37:LEU:HD23	1.99	0.45
2:OC:30:SER:OG	2:OC:33:GLU:OE1	2.21	0.45
2:OC:188:LEU:HD22	3:VC:74:ARG:NH1	2.32	0.45
2:UC:263:LEU:HD11	2:UC:274:PHE:HD2	1.82	0.45
3:VC:105:ILE:HD12	3:VC:105:ILE:HA	1.80	0.45
2:AD:75:GLU:O	2:AD:79:SER:HB3	2.17	0.45
2:AD:76:TYR:HD1	2:GD:41:MET:HE3	1.81	0.45
3:HD:239:SER:O	3:HD:242:GLU:HG2	2.16	0.45
3:HD:285:PRO:HA	4:ID:119:GLY:HA2	1.99	0.45
2:EE:88:GLU:OE1	2:EE:88:GLU:N	2.50	0.45
2:QE:202:ILE:CD1	2:WE:139:ARG:HH11	2.29	0.45
3:RE:195:PRO:HD2	3:RE:196:ASN:N	2.31	0.45
2:CF:133:ILE:O	2:CF:137:LEU:HG	2.17	0.45
2:CF:261:ILE:O	2:CF:264:LYS:HG3	2.17	0.45
4:SF:107:ALA:HB1	4:SF:124:ASP:O	2.16	0.45
2:UF:206:MET:SD	2:UF:210:GLN:HB3	2.56	0.45

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:AG:83:LYS:HD3	2:GG:26:PHE:CD1	2.51	0.45
2:AG:230:ILE:HA	2:AG:233:MET:CE	2.47	0.45
2:SG:242:VAL:HG13	2:SG:323:ILE:HG12	1.97	0.45
3:M:247:ARG:HH12	3:M:251:VAL:HG23	1.80	0.45
3:M:285:PRO:HA	4:N:119:GLY:HA2	1.99	0.45
3:C:123:VAL:HG21	3:C:146:PHE:CZ	2.52	0.45
2:J:87:GLU:HG2	2:J:89:ARG:N	2.32	0.45
4:L:65:THR:O	4:L:99:ILE:HA	2.17	0.45
2:T:178:VAL:HG11	3:V:149:THR:HG22	1.99	0.45
2:AA:97:ILE:O	2:AA:101:ARG:NH2	2.36	0.45
3:BA:183:GLU:HG3	3:BA:188:PHE:HB2	1.98	0.45
3:HA:285:PRO:HA	4:IA:119:GLY:HA2	1.98	0.45
3:NA:183:GLU:HG3	3:NA:188:PHE:HB2	1.99	0.45
2:SA:203:ILE:HA	2:SA:206:MET:HE2	1.99	0.45
3:ZA:239:SER:O	3:ZA:242:GLU:HG2	2.15	0.45
2:KB:95:GLU:HG3	2:KB:112:PHE:CE2	2.51	0.45
2:QB:87:GLU:HG2	2:QB:89:ARG:N	2.32	0.45
2:WB:118:ALA:O	2:WB:122:ILE:HG12	2.15	0.45
2:OC:29:LEU:HD13	2:OC:33:GLU:HB3	1.99	0.45
2:OC:69:LEU:HD21	2:UC:47:ILE:HB	1.98	0.45
2:AD:30:SER:OG	2:AD:33:GLU:OE1	2.32	0.45
2:GD:205:LEU:O	2:GD:205:LEU:HD23	2.17	0.45
2:GD:282:ALA:HA	2:GD:285:ILE:HG12	1.98	0.45
2:MD:115:PRO:O	2:MD:119:ALA:CB	2.64	0.45
2:MD:122:ILE:HB	2:MD:130:ILE:HG22	1.98	0.45
2:SD:65:GLN:HG3	2:YD:46:GLN:HG2	1.99	0.45
2:SD:85:LEU:HD22	2:SD:85:LEU:H	1.80	0.45
2:SD:188:LEU:H	3:ZD:74:ARG:NH2	2.15	0.45
4:WD:106:ILE:CB	3:ZD:252:ARG:HD2	2.46	0.45
2:YD:89:ARG:CD	2:YD:90:ALA:H	2.13	0.45
2:YD:207:LYS:NZ	2:YD:209:GLN:OE1	2.47	0.45
3:LE:118:PHE:CE2	3:LE:158:LEU:HD12	2.52	0.45
3:LE:190:ASN:ND2	3:RE:234:PRO:O	2.49	0.45
3:LE:267:ASP:HA	4:ME:65:THR:HA	1.99	0.45
2:QE:130:ILE:HG13	2:QE:131:ALA:N	2.32	0.45
3:DF:135:GLY:O	3:DF:211:LEU:HD21	2.17	0.45
2:OF:38:SER:O	2:OF:41:MET:HG2	2.16	0.45
2:OF:210:GLN:O	2:OF:214:VAL:HG12	2.16	0.45
2:UF:40:ALA:O	2:UF:44:VAL:HG23	2.17	0.45
2:UF:98:LEU:HB3	2:UF:108:GLU:OE1	2.16	0.45
2:AG:69:LEU:HD11	2:GG:47:ILE:CG2	2.42	0.45

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:AG:186:GLN:HB3	3:HG:74:ARG:NH2	2.21	0.45
3:BG:135:GLY:O	3:BG:211:LEU:HD21	2.17	0.45
2:SG:263:LEU:HD11	2:SG:274:PHE:HD2	1.82	0.45
3:TG:101:ASN:HB2	3:TG:146:PHE:HE1	1.81	0.45
3:TG:224:ILE:O	3:TG:228:ARG:N	2.50	0.45
2:G:164:PHE:HD1	2:SG:193:MET:O	1.99	0.45
3:M:70:PHE:CE1	2:SG:121:LEU:HD22	2.51	0.45
3:M:71:ASN:HA	2:SG:186:GLN:OE1	2.17	0.45
3:C:240:ARG:HA	3:C:240:ARG:CZ	2.47	0.45
2:T:117:SER:O	2:T:121:LEU:HG	2.16	0.45
2:AA:69:LEU:HD11	2:GA:47:ILE:CG2	2.45	0.45
2:AA:107:ILE:O	2:AA:111:ASN:ND2	2.49	0.45
2:GA:207:LYS:NZ	2:GA:209:GLN:HB3	2.32	0.45
2:GA:242:VAL:HG13	2:GA:323:ILE:HG12	1.97	0.45
2:SA:98:LEU:HB3	2:SA:108:GLU:OE1	2.16	0.45
2:EB:99:GLU:HA	2:EB:99:GLU:OE2	2.17	0.45
3:FB:237:GLU:HG2	3:FB:238:ASN:N	2.30	0.45
2:QB:263:LEU:HD11	2:QB:274:PHE:HD2	1.82	0.45
2:WB:66:PHE:HA	2:CC:45:ARG:O	2.17	0.45
2:WB:83:LYS:O	2:CC:27:LYS:NZ	2.36	0.45
2:CC:133:ILE:O	2:CC:137:LEU:HG	2.17	0.45
2:CC:246:SER:HB3	2:CC:321:MET:HE3	1.99	0.45
3:DC:101:ASN:HB2	3:DC:146:PHE:CE2	2.52	0.45
2:IC:133:ILE:O	2:IC:137:LEU:HG	2.17	0.45
2:AD:107:ILE:O	2:AD:111:ASN:ND2	2.49	0.45
3:BD:86:ILE:HD11	3:BD:221:PHE:CE1	2.51	0.45
2:GD:263:LEU:HD11	2:GD:274:PHE:HD2	1.82	0.45
2:YD:87:GLU:HG2	2:YD:89:ARG:N	2.31	0.45
2:EE:115:PRO:O	2:EE:119:ALA:CB	2.64	0.45
3:LE:65:PHE:CE1	3:LE:161:ALA:HB2	2.51	0.45
3:LE:239:SER:O	3:LE:242:GLU:HG2	2.17	0.45
3:LE:243:ASP:HA	3:LE:246:TRP:CE3	2.51	0.45
3:RE:240:ARG:HA	3:RE:240:ARG:CZ	2.47	0.45
2:WE:14:LEU:HD12	2:WE:22:ALA:HB1	1.98	0.45
2:IF:67:ALA:O	2:IF:71:ILE:HG12	2.17	0.45
2:IF:167:VAL:HG13	3:JF:130:LEU:O	2.17	0.45
2:OF:41:MET:HG3	2:OF:42:ALA:N	2.31	0.45
3:VF:180:VAL:HG23	3:VF:181:ARG:H	1.82	0.45
2:AG:85:LEU:HD22	2:AG:85:LEU:H	1.81	0.45
2:AG:242:VAL:HG13	2:AG:323:ILE:HG12	1.98	0.45
3:BG:75:ARG:HG3	3:BG:208:ILE:HD11	1.98	0.45

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:GG:168:GLN:HE22	3:HG:130:LEU:HD11	1.81	0.45
2:MG:167:VAL:HG21	3:NG:131:PHE:CD1	2.52	0.45
2:G:83:LYS:HD3	2:B:26:PHE:HD1	1.82	0.45
3:M:294:ALA:O	3:M:301:VAL:N	2.47	0.45
2:B:128:GLN:CG	3:C:131:PHE:HB3	2.37	0.45
2:B:151:GLU:HA	2:B:154:ARG:HE	1.81	0.45
2:B:263:LEU:HD11	2:B:274:PHE:HD2	1.81	0.45
3:K:237:GLU:HG2	3:K:238:ASN:H	1.82	0.45
4:Y:107:ALA:HB1	4:Y:124:ASP:O	2.17	0.45
2:AA:15:MET:N	2:AA:15:MET:SD	2.90	0.45
2:GA:87:GLU:HG2	2:GA:89:ARG:N	2.32	0.45
2:GA:151:GLU:HA	2:GA:154:ARG:HE	1.81	0.45
3:NA:101:ASN:HB2	3:NA:146:PHE:HE2	1.82	0.45
2:SA:26:PHE:HA	2:SA:29:LEU:CD2	2.46	0.45
3:TA:285:PRO:HA	4:UA:119:GLY:HA2	1.99	0.45
3:ZA:111:ARG:HE	3:FB:241:HIS:CE1	2.35	0.45
3:ZA:185:GLN:HG3	3:ZA:186:VAL:H	1.82	0.45
2:EB:261:ILE:O	2:EB:264:LYS:HG3	2.17	0.45
3:FB:111:ARG:NE	3:FB:111:ARG:HA	2.32	0.45
3:FB:169:TRP:CD1	3:FB:175:LEU:HD13	2.50	0.45
3:LB:115:LEU:HD11	3:LB:218:CYS:HB3	1.98	0.45
2:IC:219:ARG:HH22	2:IC:230:ILE:HD12	1.82	0.45
2:OC:188:LEU:HB3	3:VC:74:ARG:NE	2.32	0.45
3:BD:214:GLU:OE1	3:BD:215:PHE:N	2.50	0.45
2:MD:26:PHE:HA	2:MD:29:LEU:CD2	2.47	0.45
2:MD:38:SER:HA	2:MD:41:MET:HE2	1.99	0.45
3:ND:190:ASN:ND2	3:TD:234:PRO:O	2.49	0.45
2:SD:128:GLN:NE2	2:SD:164:PHE:HD2	2.11	0.45
2:YD:15:MET:N	2:YD:15:MET:SD	2.88	0.45
2:IF:32:ARG:NE	2:IF:33:GLU:OE2	2.50	0.45
2:OF:155:HIS:O	2:OF:159:LEU:HG	2.16	0.45
2:UF:178:VAL:HG11	3:VF:149:THR:HG22	1.99	0.45
2:AG:66:PHE:HA	2:GG:45:ARG:O	2.17	0.45
2:AG:98:LEU:HB3	2:AG:108:GLU:OE1	2.16	0.45
2:AG:207:LYS:NZ	2:AG:209:GLN:HB3	2.30	0.45
2:GG:261:ILE:O	2:GG:264:LYS:HG3	2.17	0.45
3:TG:96:LEU:HD23	3:TG:97:PRO:CD	2.47	0.45
3:TG:113:THR:OG1	3:TG:191:ILE:HG22	2.17	0.45
3:TG:191:ILE:HG13	3:TG:192:THR:HG23	1.98	0.45
3:M:74:ARG:NH1	2:SG:186:GLN:HB3	2.32	0.45
3:M:237:GLU:HG2	3:M:238:ASN:N	2.32	0.45

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:B:205:LEU:CD2	2:J:140:SER:HA	2.44	0.45
3:C:54:LEU:HD13	3:C:231:LEU:HD21	1.98	0.45
3:C:75:ARG:HG3	3:C:208:ILE:HD11	1.98	0.45
2:T:79:SER:HA	2:T:82:VAL:HG12	1.98	0.45
2:T:134:LEU:HD13	2:T:146:LEU:HD21	1.99	0.45
2:T:167:VAL:HG21	3:V:131:PHE:CD1	2.51	0.45
3:V:180:VAL:HG23	3:V:181:ARG:H	1.82	0.45
2:GA:263:LEU:HD11	2:GA:274:PHE:HD2	1.82	0.45
3:NA:54:LEU:HD13	3:NA:231:LEU:HD21	1.98	0.45
2:YA:107:ILE:HA	2:YA:110:LEU:HD12	1.98	0.45
3:XB:180:VAL:HG23	3:XB:181:ARG:H	1.82	0.45
2:IC:178:VAL:HG11	3:JC:149:THR:HG22	1.99	0.45
3:JC:207:GLU:HG3	3:JC:212:THR:HG22	1.99	0.45
3:JC:285:PRO:HA	4:KC:119:GLY:HA2	1.99	0.45
2:OC:30:SER:O	2:OC:34:VAL:HG23	2.17	0.45
2:OC:263:LEU:HD11	2:OC:274:PHE:HD2	1.81	0.45
3:VC:134:ASP:OD2	3:VC:136:ARG:N	2.39	0.45
3:VC:237:GLU:HG2	3:VC:238:ASN:N	2.31	0.45
2:AD:79:SER:O	2:AD:83:LYS:HB3	2.17	0.45
3:BD:285:PRO:HA	4:CD:119:GLY:HA2	1.99	0.45
2:GD:76:TYR:HD1	2:GD:79:SER:OG	1.99	0.45
3:HD:58:ASN:OD1	3:HD:81:VAL:HG11	2.16	0.45
2:MD:100:THR:OG1	2:MD:101:ARG:NH2	2.50	0.45
2:SD:151:GLU:HA	2:SD:154:ARG:HE	1.81	0.45
3:TD:66:ARG:HG3	3:TD:77:PRO:O	2.17	0.45
2:YD:11:VAL:O	2:YD:14:LEU:HB3	2.17	0.45
2:EE:246:SER:HB3	2:EE:321:MET:HE3	1.99	0.45
3:FE:198:ILE:HG13	3:FE:221:PHE:CD1	2.52	0.45
2:KE:210:GLN:O	2:KE:214:VAL:HG12	2.16	0.45
3:DF:97:PRO:O	3:DF:100:THR:OG1	2.20	0.45
2:IF:65:GLN:HG3	2:OF:46:GLN:HG2	1.98	0.45
2:IF:188:LEU:HD22	3:PF:74:ARG:NE	2.32	0.45
2:OF:83:LYS:O	2:UF:27:LYS:HE3	2.17	0.45
2:OF:146:LEU:HD11	2:OF:157:VAL:HG11	1.98	0.45
2:OF:263:LEU:HD11	2:OF:274:PHE:HD2	1.82	0.45
2:AG:132:THR:O	2:AG:135:VAL:HB	2.17	0.45
2:AG:167:VAL:HG21	3:BG:131:PHE:CD1	2.51	0.45
2:AG:234:PHE:CD2	2:AG:313:ARG:HB3	2.52	0.45
3:BG:100:THR:HG23	3:BG:119:SER:HA	1.99	0.45
3:NG:234:PRO:HA	3:NG:235:PRO:HD3	1.85	0.45
3:TG:192:THR:OG1	3:TG:194:SER:O	2.34	0.45

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:G:53:THR:O	2:G:57:SER:OG	2.33	0.45
3:M:239:SER:O	3:M:242:GLU:HG2	2.16	0.45
3:C:122:LEU:CD1	3:C:215:PHE:HB3	2.42	0.45
3:C:285:PRO:HA	4:D:119:GLY:HA2	1.99	0.45
3:K:238:ASN:HB3	3:K:241:HIS:ND1	2.31	0.45
2:T:102:ASP:O	2:T:107:ILE:HD13	2.17	0.45
2:T:168:GLN:OE1	2:T:168:GLN:N	2.44	0.45
3:TA:118:PHE:HE2	3:TA:158:LEU:HD12	1.80	0.45
2:YA:107:ILE:HD12	2:YA:107:ILE:N	2.31	0.45
4:AB:65:THR:O	4:AB:99:ILE:HA	2.17	0.45
2:WB:85:LEU:H	2:WB:85:LEU:HD22	1.81	0.45
2:WB:133:ILE:O	2:WB:137:LEU:HG	2.16	0.45
3:XB:58:ASN:OD1	3:XB:81:VAL:HG11	2.16	0.45
2:IC:119:ALA:HB1	2:IC:149:PHE:CE1	2.52	0.45
2:OC:87:GLU:HG2	2:OC:89:ARG:N	2.32	0.45
3:PC:106:HIS:CE1	3:PC:113:THR:CG2	3.00	0.45
2:AD:12:ILE:HD11	2:AD:55:VAL:HG11	1.98	0.45
2:AD:109:THR:O	2:AD:113:MET:HG3	2.16	0.45
3:HD:110:LEU:HG	3:HD:223:MET:HB3	1.98	0.45
3:HD:224:ILE:O	3:HD:228:ARG:N	2.50	0.45
3:LE:135:GLY:O	3:LE:211:LEU:HD21	2.17	0.45
2:WE:76:TYR:HD1	2:CF:41:MET:CE	2.30	0.45
2:CF:65:GLN:HE21	2:IF:47:ILE:H	1.64	0.45
2:CF:113:MET:CG	2:CF:117:SER:HB2	2.47	0.45
2:CF:168:GLN:HG2	2:CF:170:ALA:H	1.81	0.45
2:IF:76:TYR:CD2	2:OF:41:MET:HE2	2.51	0.45
2:IF:163:THR:HG21	3:JF:134:ASP:OD2	2.17	0.45
2:UF:109:THR:O	2:UF:113:MET:HG3	2.16	0.45
2:GG:171:ALA:HB1	3:HG:131:PHE:CE2	2.52	0.45
2:SG:40:ALA:O	2:SG:44:VAL:HG23	2.16	0.45
3:TG:51:LEU:O	3:TG:55:GLU:HG3	2.17	0.45
3:TG:169:TRP:CD1	3:TG:175:LEU:HD13	2.50	0.45
2:G:45:ARG:O	2:SG:66:PHE:HA	2.17	0.45
2:G:111:ASN:HA	2:G:141:GLN:OE1	2.16	0.45
3:M:252:ARG:HD3	4:WG:105:LEU:O	2.17	0.45
3:C:169:TRP:CD1	3:C:175:LEU:HD13	2.50	0.45
2:T:174:GLU:HG2	3:V:152:ARG:HG2	1.99	0.45
2:AA:178:VAL:HG11	3:BA:149:THR:HG22	1.99	0.45
3:HA:240:ARG:HA	3:HA:240:ARG:HH11	1.78	0.45
2:EB:206:MET:HG2	2:KB:143:ALA:HB1	1.99	0.45
3:FB:135:GLY:O	3:FB:211:LEU:HD21	2.16	0.45

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:KB:263:LEU:HD11	2:KB:274:PHE:HD2	1.82	0.45
2:QB:15:MET:N	2:QB:15:MET:SD	2.89	0.45
3:XB:105:ILE:HD12	3:XB:105:ILE:HA	1.76	0.45
3:XB:106:HIS:HE1	3:XB:113:THR:CG2	2.29	0.45
3:XB:110:LEU:HG	3:XB:223:MET:HB3	1.99	0.45
3:DC:97:PRO:O	3:DC:100:THR:OG1	2.20	0.45
3:DC:239:SER:O	3:DC:242:GLU:HG2	2.17	0.45
3:DC:294:ALA:O	3:DC:301:VAL:N	2.50	0.45
2:IC:207:LYS:HD3	2:IC:210:GLN:HE21	1.82	0.45
2:IC:261:ILE:O	2:IC:264:LYS:HG3	2.17	0.45
3:JC:128:ASP:OD1	3:JC:133:GLY:HA3	2.17	0.45
3:JC:224:ILE:O	3:JC:228:ARG:N	2.50	0.45
3:JC:229:GLU:OE1	3:JC:230:LEU:N	2.50	0.45
3:PC:237:GLU:HG2	3:PC:238:ASN:H	1.81	0.45
3:BD:96:LEU:HD23	3:BD:97:PRO:CD	2.47	0.45
2:GD:12:ILE:HD11	2:GD:55:VAL:HG11	1.98	0.45
2:MD:263:LEU:HD11	2:MD:274:PHE:HD2	1.82	0.45
2:EE:242:VAL:HG13	2:EE:323:ILE:HG12	1.99	0.45
3:FE:58:ASN:ND2	3:FE:59:GLU:OE2	2.50	0.45
3:FE:105:ILE:HD12	3:FE:105:ILE:HA	1.83	0.45
2:KE:78:ARG:O	2:KE:82:VAL:HG12	2.16	0.45
2:WE:191:SER:OG	2:WE:192:LYS:N	2.50	0.45
2:IF:196:VAL:HG12	2:IF:232:GLU:OE2	2.17	0.45
3:VF:135:GLY:O	3:VF:211:LEU:HD21	2.16	0.45
2:AG:114:GLU:HB2	2:AG:117:SER:OG	2.17	0.45
2:MG:197:ARG:NH1	2:MG:232:GLU:HB3	2.32	0.45
2:SG:108:GLU:HG3	2:SG:112:PHE:HE2	1.82	0.45
2:G:14:LEU:HA	2:G:17:ILE:HG12	1.99	0.44
3:M:69:LEU:O	3:M:73:LEU:HB2	2.17	0.44
3:M:214:GLU:OE1	3:M:215:PHE:N	2.51	0.44
2:J:247:ILE:HD13	2:J:273:LYS:HE3	1.99	0.44
3:K:185:GLN:HG3	3:K:186:VAL:H	1.82	0.44
2:T:87:GLU:HG2	2:T:89:ARG:N	2.32	0.44
4:Y:106:ILE:O	3:BA:252:ARG:HG2	2.18	0.44
2:AA:89:ARG:HG3	2:AA:89:ARG:NH1	2.31	0.44
3:BA:77:PRO:HB3	3:BA:208:ILE:HB	1.99	0.44
3:BA:219:LEU:HD23	3:BA:219:LEU:HA	1.82	0.44
2:GA:107:ILE:HG23	2:GA:137:LEU:HD23	1.99	0.44
3:HA:118:PHE:HE2	3:HA:158:LEU:HD12	1.80	0.44
3:HA:294:ALA:O	3:HA:301:VAL:N	2.49	0.44
3:NA:60:ARG:HE	3:NA:60:ARG:HB2	1.56	0.44

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:SA:198:THR:HG23	2:SA:201:GLU:OE1	2.17	0.44
2:SA:263:LEU:HD11	2:SA:274:PHE:HD2	1.82	0.44
3:TA:113:THR:OG1	3:TA:191:ILE:HG22	2.17	0.44
2:EB:81:LEU:O	2:EB:85:LEU:HD13	2.17	0.44
2:QB:41:MET:SD	2:QB:42:ALA:N	2.90	0.44
2:QB:66:PHE:HA	2:WB:45:ARG:O	2.16	0.44
2:QB:127:PRO:O	2:QB:130:ILE:HG13	2.18	0.44
3:RB:204:PHE:N	3:RB:204:PHE:CD1	2.85	0.44
3:RB:234:PRO:HA	3:RB:235:PRO:HD3	1.86	0.44
2:WB:178:VAL:HG11	3:XB:149:THR:HG22	1.99	0.44
2:CC:167:VAL:HG21	3:DC:131:PHE:CD1	2.52	0.44
2:IC:45:ARG:CG	2:IC:46:GLN:HG3	2.42	0.44
2:IC:66:PHE:HA	2:OC:45:ARG:O	2.17	0.44
2:IC:87:GLU:HG2	2:IC:89:ARG:N	2.33	0.44
2:UC:111:ASN:HA	2:UC:141:GLN:OE1	2.17	0.44
3:VC:180:VAL:HG23	3:VC:181:ARG:H	1.82	0.44
2:AD:205:LEU:HD22	2:GD:140:SER:HA	1.99	0.44
3:TD:135:GLY:O	3:TD:211:LEU:HD21	2.17	0.44
4:WD:106:ILE:O	4:WD:126:ILE:N	2.38	0.44
3:ZD:189:THR:OG1	3:ZD:190:ASN:N	2.50	0.44
2:KE:119:ALA:HB1	2:KE:149:PHE:CE1	2.52	0.44
2:QE:83:LYS:HD3	2:WE:26:PHE:CD2	2.52	0.44
2:QE:87:GLU:HG2	2:QE:89:ARG:N	2.33	0.44
2:CF:152:ARG:HH12	2:CF:153:LEU:CD1	2.28	0.44
2:CF:263:LEU:HD11	2:CF:274:PHE:HD2	1.82	0.44
3:PF:106:HIS:HE1	3:PF:113:THR:CG2	2.30	0.44
3:VF:117:VAL:O	3:VF:215:PHE:HB2	2.17	0.44
3:VF:240:ARG:HA	3:VF:240:ARG:HH11	1.81	0.44
3:VF:285:PRO:HA	4:WF:119:GLY:HA2	1.99	0.44
2:GG:219:ARG:HH22	2:GG:230:ILE:HD12	1.81	0.44
2:MG:115:PRO:O	2:MG:119:ALA:CB	2.65	0.44
2:G:109:THR:O	2:G:113:MET:HG3	2.17	0.44
2:J:119:ALA:HB1	2:J:149:PHE:CE1	2.52	0.44
3:K:101:ASN:HB2	3:K:146:PHE:CE1	2.52	0.44
2:T:48:SER:OG	2:T:51:GLN:OE1	2.29	0.44
3:HA:237:GLU:HG2	3:HA:238:ASN:N	2.32	0.44
2:MA:30:SER:O	2:MA:34:VAL:HG23	2.17	0.44
3:NA:127:VAL:HG23	3:NA:153:VAL:HG21	2.00	0.44
3:NA:252:ARG:HA	3:NA:252:ARG:HD2	1.79	0.44
2:SA:76:TYR:HE2	2:YA:26:PHE:CE2	2.35	0.44
4:UA:65:THR:O	4:UA:99:ILE:HA	2.17	0.44

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:YA:188:LEU:HB3	3:FB:74:ARG:NE	2.33	0.44
2:EB:108:GLU:OE2	2:EB:112:PHE:HE2	1.98	0.44
2:KB:17:ILE:HD12	2:KB:21:ARG:HB2	1.99	0.44
3:RB:110:LEU:HD11	3:RB:223:MET:O	2.17	0.44
3:XB:106:HIS:HE1	3:XB:113:THR:HG22	1.77	0.44
3:XB:237:GLU:HG2	3:XB:238:ASN:H	1.82	0.44
2:CC:100:THR:OG1	2:CC:101:ARG:NH2	2.49	0.44
2:IC:207:LYS:NZ	2:IC:209:GLN:HB3	2.30	0.44
2:OC:26:PHE:HE1	2:OC:34:VAL:HA	1.82	0.44
2:OC:115:PRO:O	2:OC:119:ALA:CB	2.65	0.44
2:UC:83:LYS:HD3	2:AD:26:PHE:CD1	2.52	0.44
4:ED:106:ILE:O	3:HD:252:ARG:HG2	2.17	0.44
2:GD:205:LEU:CD2	2:MD:140:SER:HA	2.47	0.44
2:MD:65:GLN:HG3	2:SD:46:GLN:HG2	2.00	0.44
2:MD:79:SER:O	2:MD:83:LYS:HG3	2.17	0.44
2:MD:186:GLN:C	3:TD:74:ARG:HH22	2.20	0.44
2:YD:30:SER:OG	2:YD:33:GLU:OE1	2.32	0.44
2:EE:151:GLU:HA	2:EE:154:ARG:HE	1.81	0.44
2:KE:83:LYS:HA	2:QE:23:ALA:HB1	1.99	0.44
2:KE:152:ARG:HH22	2:KE:153:LEU:HB2	1.82	0.44
2:QE:80:VAL:HG12	2:WE:26:PHE:CE2	2.52	0.44
3:DF:180:VAL:HG23	3:DF:181:ARG:N	2.33	0.44
3:PF:202:THR:OG1	3:PF:202:THR:O	2.35	0.44
2:UF:30:SER:O	2:UF:34:VAL:HG23	2.16	0.44
2:UF:66:PHE:HA	2:AG:45:ARG:O	2.17	0.44
2:UF:298:LEU:HB2	2:UF:302:GLU:OE1	2.16	0.44
3:VF:190:ASN:O	3:BG:236:LEU:HD13	2.16	0.44
2:AG:53:THR:O	2:AG:57:SER:OG	2.32	0.44
2:AG:122:ILE:HB	2:AG:130:ILE:HG22	1.98	0.44
2:AG:133:ILE:O	2:AG:137:LEU:HG	2.17	0.44
2:AG:261:ILE:H	2:AG:261:ILE:HG13	1.60	0.44
2:GG:89:ARG:HG3	2:GG:90:ALA:N	2.32	0.44
3:HG:116:VAL:HG12	3:HG:217:ILE:HD13	1.98	0.44
2:MG:159:LEU:HD12	2:MG:159:LEU:HA	1.82	0.44
2:SG:31:THR:O	2:SG:34:VAL:HB	2.17	0.44
3:TG:247:ARG:NH1	3:TG:247:ARG:O	2.51	0.44
2:B:107:ILE:HG22	2:B:111:ASN:ND2	2.30	0.44
2:GA:131:ALA:HA	2:GA:134:LEU:HD12	1.98	0.44
2:GA:188:LEU:HD13	3:NA:74:ARG:CD	2.43	0.44
3:ZA:294:ALA:O	3:ZA:301:VAL:N	2.50	0.44
3:LB:55:GLU:HA	3:LB:58:ASN:ND2	2.33	0.44

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:DC:54:LEU:HD13	3:DC:231:LEU:HD21	2.00	0.44
3:DC:135:GLY:O	3:DC:211:LEU:HD21	2.17	0.44
2:IC:83:LYS:HD2	2:OC:26:PHE:CD2	2.52	0.44
3:PC:252:ARG:HA	3:PC:252:ARG:HD2	1.74	0.44
2:UC:76:TYR:HD1	2:AD:41:MET:HE1	1.82	0.44
3:VC:96:LEU:HD23	3:VC:97:PRO:CD	2.47	0.44
3:VC:111:ARG:NH2	3:VC:222:SER:O	2.51	0.44
2:AD:97:ILE:O	2:AD:101:ARG:NH2	2.37	0.44
2:AD:111:ASN:HA	2:AD:141:GLN:OE1	2.18	0.44
3:BD:69:LEU:O	3:BD:73:LEU:HB2	2.17	0.44
3:BD:185:GLN:HG3	3:BD:186:VAL:H	1.82	0.44
2:GD:307:ALA:O	2:GD:310:LEU:HG	2.18	0.44
3:ND:135:GLY:O	3:ND:211:LEU:HD21	2.17	0.44
4:QD:106:ILE:O	4:QD:126:ILE:N	2.40	0.44
3:ZD:237:GLU:HG2	3:ZD:238:ASN:N	2.33	0.44
3:FE:183:GLU:OE1	3:FE:184:MET:N	2.51	0.44
3:FE:190:ASN:O	3:FE:190:ASN:OD1	2.36	0.44
2:QE:263:LEU:HD11	2:QE:274:PHE:HD2	1.82	0.44
2:CF:81:LEU:O	2:CF:85:LEU:HD13	2.17	0.44
3:DF:127:VAL:HG23	3:DF:153:VAL:HG21	1.97	0.44
3:PF:191:ILE:HG13	3:PF:192:THR:HG23	1.98	0.44
3:VF:106:HIS:CD2	3:VF:113:THR:HG22	2.52	0.44
3:VF:169:TRP:CD1	3:VF:175:LEU:HD13	2.50	0.44
2:AG:87:GLU:HG2	2:AG:89:ARG:N	2.33	0.44
3:BG:238:ASN:HB3	3:BG:241:HIS:ND1	2.33	0.44
3:TG:185:GLN:HG3	3:TG:186:VAL:H	1.82	0.44
2:G:72:ASN:ND2	2:B:41:MET:HG3	2.31	0.44
2:T:72:ASN:HD21	2:AA:41:MET:HB2	1.83	0.44
2:AA:167:VAL:HG21	3:BA:131:PHE:CD1	2.53	0.44
2:GA:97:ILE:O	2:GA:101:ARG:NH2	2.31	0.44
2:GA:109:THR:O	2:GA:113:MET:HG3	2.18	0.44
2:MA:107:ILE:HD12	2:MA:107:ILE:N	2.32	0.44
3:NA:238:ASN:HB3	3:NA:241:HIS:ND1	2.32	0.44
2:SA:144:ASP:O	2:SA:148:LEU:HD12	2.17	0.44
3:TA:185:GLN:HG2	3:TA:187:LYS:NZ	2.33	0.44
3:FB:294:ALA:O	3:FB:301:VAL:N	2.49	0.44
2:KB:188:LEU:HB3	3:RB:74:ARG:CZ	2.47	0.44
2:QB:105:SER:OG	2:QB:107:ILE:HD12	2.16	0.44
2:QB:246:SER:HB3	2:QB:321:MET:HE3	1.99	0.44
2:QB:307:ALA:O	2:QB:310:LEU:HG	2.16	0.44
2:IC:30:SER:O	2:IC:34:VAL:HG23	2.17	0.44

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:IC:30:SER:OG	2:IC:33:GLU:OE1	2.29	0.44
2:IC:188:LEU:HB3	3:PC:74:ARG:CZ	2.47	0.44
2:IC:263:LEU:HD11	2:IC:274:PHE:HD2	1.81	0.44
2:OC:65:GLN:NE2	2:UC:47:ILE:H	2.15	0.44
2:OC:107:ILE:O	2:OC:111:ASN:ND2	2.51	0.44
2:OC:151:GLU:HA	2:OC:154:ARG:HE	1.81	0.44
3:BD:240:ARG:HA	3:BD:240:ARG:CZ	2.48	0.44
3:HD:135:GLY:O	3:HD:211:LEU:HD21	2.17	0.44
2:MD:186:GLN:NE2	3:TD:71:ASN:HA	2.29	0.44
2:SD:174:GLU:HG2	3:TD:152:ARG:HG2	1.98	0.44
2:YD:128:GLN:CG	3:ZD:131:PHE:HB3	2.34	0.44
2:EE:234:PHE:CD2	2:EE:313:ARG:HB3	2.52	0.44
2:KE:31:THR:O	2:KE:34:VAL:HB	2.18	0.44
2:KE:126:HIS:O	2:KE:130:ILE:HG23	2.18	0.44
2:WE:144:ASP:O	2:WE:148:LEU:HD12	2.18	0.44
3:DF:101:ASN:HB2	3:DF:146:PHE:CE2	2.53	0.44
2:IF:37:LEU:HD12	2:IF:38:SER:N	2.33	0.44
2:IF:37:LEU:O	2:IF:41:MET:HG2	2.17	0.44
2:IF:111:ASN:HA	2:IF:141:GLN:CD	2.38	0.44
2:OF:83:LYS:NZ	2:UF:26:PHE:CD2	2.78	0.44
2:OF:100:THR:OG1	2:OF:101:ARG:NH2	2.50	0.44
2:UF:81:LEU:O	2:UF:85:LEU:HD13	2.18	0.44
3:BG:101:ASN:HD21	3:BG:154:ILE:HG21	1.81	0.44
2:MG:172:LEU:HD12	2:MG:172:LEU:HA	1.70	0.44
3:NG:66:ARG:HG3	3:NG:77:PRO:O	2.17	0.44
3:M:267:ASP:HA	4:N:65:THR:HA	2.00	0.44
2:B:110:LEU:HD13	2:B:137:LEU:HD21	2.00	0.44
3:C:111:ARG:HA	3:C:111:ARG:CZ	2.48	0.44
3:C:239:SER:O	3:C:242:GLU:HG2	2.18	0.44
2:AA:53:THR:O	2:AA:57:SER:OG	2.33	0.44
3:HA:239:SER:O	3:HA:242:GLU:HG2	2.18	0.44
2:MA:261:ILE:O	2:MA:264:LYS:HG3	2.17	0.44
3:NA:101:ASN:HB2	3:NA:146:PHE:CE2	2.52	0.44
3:NA:198:ILE:HD12	3:NA:198:ILE:HA	1.91	0.44
2:YA:128:GLN:CG	3:ZA:131:PHE:HB3	2.37	0.44
2:YA:263:LEU:HD11	2:YA:274:PHE:HD2	1.82	0.44
2:KB:87:GLU:HG2	2:KB:89:ARG:N	2.33	0.44
2:KB:282:ALA:HA	2:KB:285:ILE:HG12	2.00	0.44
3:LB:79:ILE:HG22	3:LB:204:PHE:HB3	2.00	0.44
3:LB:127:VAL:HG23	3:LB:153:VAL:HG11	1.99	0.44
2:QB:107:ILE:HD13	2:QB:136:HIS:CB	2.44	0.44

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:RB:267:ASP:HA	4:SB:65:THR:HA	2.00	0.44
2:WB:282:ALA:HA	2:WB:285:ILE:HG12	2.00	0.44
3:BD:267:ASP:HA	4:CD:65:THR:HA	1.99	0.44
3:HD:101:ASN:HB2	3:HD:146:PHE:HE2	1.82	0.44
3:HD:185:GLN:HG3	3:HD:186:VAL:H	1.83	0.44
2:MD:198:THR:O	2:MD:202:ILE:HG12	2.17	0.44
2:MD:246:SER:HB3	2:MD:321:MET:HE3	2.00	0.44
2:SD:26:PHE:CE1	2:SD:34:VAL:HG22	2.52	0.44
2:SD:178:VAL:HG11	3:TD:149:THR:HG22	2.00	0.44
2:SD:188:LEU:HD22	3:ZD:74:ARG:CZ	2.48	0.44
3:FE:247:ARG:HH11	3:FE:250:LEU:HD22	1.82	0.44
2:KE:81:LEU:O	2:KE:85:LEU:HD13	2.18	0.44
2:KE:107:ILE:HA	2:KE:110:LEU:HD12	2.00	0.44
2:KE:110:LEU:O	2:KE:113:MET:N	2.48	0.44
2:QE:188:LEU:HD23	2:QE:188:LEU:N	2.28	0.44
3:XE:237:GLU:HG2	3:XE:238:ASN:H	1.83	0.44
3:DF:183:GLU:OE1	3:DF:184:MET:N	2.50	0.44
3:HG:237:GLU:HG2	3:HG:238:ASN:N	2.33	0.44
2:SG:149:PHE:CD2	2:SG:153:LEU:HD22	2.53	0.44
2:G:282:ALA:HA	2:G:285:ILE:HG12	2.00	0.44
3:M:58:ASN:OD1	3:M:81:VAL:HG11	2.18	0.44
2:B:141:GLN:HA	2:B:144:ASP:OD2	2.18	0.44
2:J:20:ASP:OD1	2:J:20:ASP:N	2.50	0.44
2:T:31:THR:HA	2:T:34:VAL:CG2	2.48	0.44
2:T:118:ALA:HA	2:T:121:LEU:CD1	2.46	0.44
2:T:263:LEU:HD11	2:T:274:PHE:HD2	1.83	0.44
3:V:195:PRO:HD2	3:V:196:ASN:N	2.32	0.44
2:AA:107:ILE:HG22	2:AA:111:ASN:HD21	1.82	0.44
2:AA:138:LYS:HD3	2:AA:138:LYS:HA	1.85	0.44
2:AA:298:LEU:O	2:AA:302:GLU:HG2	2.18	0.44
2:MA:87:GLU:HG2	2:MA:89:ARG:H	1.82	0.44
2:MA:188:LEU:HD13	3:TA:74:ARG:CD	2.43	0.44
3:NA:135:GLY:O	3:NA:211:LEU:HD21	2.18	0.44
2:EB:79:SER:O	2:EB:83:LYS:HG3	2.17	0.44
2:EB:87:GLU:HG2	2:EB:89:ARG:H	1.82	0.44
2:KB:79:SER:O	2:KB:83:LYS:HG3	2.17	0.44
2:QB:193:MET:N	2:QB:193:MET:SD	2.91	0.44
3:RB:247:ARG:O	3:RB:247:ARG:HD2	2.17	0.44
2:WB:20:ASP:OD1	2:WB:20:ASP:N	2.51	0.44
2:IC:282:ALA:O	2:IC:286:LEU:HD23	2.18	0.44
3:JC:207:GLU:OE1	3:JC:207:GLU:N	2.51	0.44

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:OC:128:GLN:HE22	2:OC:164:PHE:HD1	1.61	0.44
2:UC:167:VAL:HG21	3:VC:131:PHE:CD1	2.53	0.44
3:VC:247:ARG:HH21	3:VC:250:LEU:HD22	1.82	0.44
3:HD:108:LYS:HE2	3:HD:108:LYS:HB2	1.81	0.44
3:ND:183:GLU:OE1	3:ND:184:MET:N	2.51	0.44
2:YD:178:VAL:HG11	3:ZD:149:THR:HG22	2.00	0.44
2:EE:207:LYS:H	2:EE:210:GLN:HE21	1.66	0.44
3:FE:79:ILE:HG22	3:FE:204:PHE:HB3	2.00	0.44
3:FE:198:ILE:HD12	3:FE:199:VAL:N	2.33	0.44
2:KE:126:HIS:CE1	3:LE:127:VAL:HG13	2.53	0.44
2:KE:234:PHE:CD2	2:KE:313:ARG:HB3	2.53	0.44
3:LE:150:GLU:O	3:LE:153:VAL:HG12	2.18	0.44
2:WE:83:LYS:O	2:CF:27:LYS:HD2	2.18	0.44
2:WE:87:GLU:HG2	2:WE:89:ARG:N	2.33	0.44
2:IF:48:SER:OG	2:IF:51:GLN:OE1	2.32	0.44
2:IF:76:TYR:HE1	2:OF:26:PHE:CZ	2.36	0.44
2:OF:76:TYR:HD2	2:UF:41:MET:HE3	1.83	0.44
3:PF:150:GLU:O	3:PF:153:VAL:HG12	2.18	0.44
2:UF:234:PHE:CD2	2:UF:313:ARG:HB3	2.53	0.44
2:AG:282:ALA:O	2:AG:286:LEU:HD23	2.18	0.44
2:MG:120:ASP:OD1	3:TG:76:SER:OG	2.36	0.44
2:SG:307:ALA:O	2:SG:310:LEU:HG	2.18	0.44
3:M:96:LEU:HD23	3:M:97:PRO:CD	2.47	0.44
2:B:80:VAL:HG12	2:J:26:PHE:CE2	2.52	0.44
2:J:207:LYS:NZ	2:J:209:GLN:HB3	2.33	0.44
2:GA:247:ILE:HD13	2:GA:273:LYS:HE3	1.99	0.44
2:MA:282:ALA:HA	2:MA:285:ILE:HG12	2.00	0.44
2:YA:198:THR:O	2:YA:201:GLU:HG2	2.17	0.44
2:YA:282:ALA:O	2:YA:286:LEU:HD23	2.18	0.44
2:EB:89:ARG:NH2	2:KB:31:THR:HG23	2.32	0.44
2:EB:103:THR:C	2:EB:106:GLY:H	2.21	0.44
2:EB:168:GLN:HB2	2:EB:169:PRO:HD2	1.99	0.44
4:GB:65:THR:O	4:GB:99:ILE:HA	2.17	0.44
2:KB:38:SER:HA	2:KB:41:MET:HE1	2.00	0.44
2:KB:158:MET:HE3	2:KB:159:LEU:HD22	1.98	0.44
2:QB:69:LEU:HD12	2:WB:45:ARG:O	2.18	0.44
2:IC:108:GLU:HG3	2:IC:112:PHE:HE2	1.83	0.44
3:JC:54:LEU:HD13	3:JC:231:LEU:HD21	2.00	0.44
3:JC:116:VAL:HG12	3:JC:217:ILE:HD13	1.99	0.44
3:JC:267:ASP:HA	4:KC:65:THR:HA	1.99	0.44
3:VC:101:ASN:HB2	3:VC:146:PHE:CE1	2.52	0.44

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:VC:123:VAL:HG21	3:VC:146:PHE:CZ	2.53	0.44
2:AD:26:PHE:HD2	2:AD:29:LEU:HD21	1.82	0.44
2:GD:53:THR:O	2:GD:57:SER:OG	2.29	0.44
2:GD:74:ASN:O	2:GD:78:ARG:NE	2.51	0.44
3:HD:237:GLU:HG2	3:HD:238:ASN:H	1.83	0.44
2:MD:247:ILE:HD13	2:MD:273:LYS:HE3	1.99	0.44
2:YD:107:ILE:O	2:YD:111:ASN:ND2	2.51	0.44
2:EE:209:GLN:O	2:EE:212:GLU:HG3	2.17	0.44
3:FE:100:THR:HG23	3:FE:119:SER:HA	1.99	0.44
2:KE:128:GLN:CG	3:LE:131:PHE:HB3	2.33	0.44
3:LE:117:VAL:O	3:LE:215:PHE:HB2	2.17	0.44
2:QE:282:ALA:O	2:QE:286:LEU:HD23	2.18	0.44
4:SE:65:THR:O	4:SE:99:ILE:HA	2.18	0.44
3:XE:116:VAL:HG21	3:XE:158:LEU:HD11	2.00	0.44
4:YE:65:THR:O	4:YE:99:ILE:HA	2.18	0.44
2:CF:107:ILE:HA	2:CF:110:LEU:HD12	2.00	0.44
3:DF:207:GLU:HG3	3:DF:212:THR:HG22	2.00	0.44
2:IF:78:ARG:O	2:IF:82:VAL:HG12	2.17	0.44
2:IF:83:LYS:HG3	2:IF:84:ALA:N	2.32	0.44
3:JF:169:TRP:CD1	3:JF:175:LEU:HD13	2.50	0.44
2:OF:174:GLU:HG2	3:PF:152:ARG:HG2	2.00	0.44
3:PF:237:GLU:HG2	3:PF:238:ASN:N	2.32	0.44
2:UF:53:THR:O	2:UF:57:SER:OG	2.32	0.44
2:UF:89:ARG:NH2	2:AG:31:THR:HG23	2.33	0.44
4:WF:101:ILE:N	4:WF:104:TYR:O	2.34	0.44
2:AG:219:ARG:NH2	2:AG:227:GLN:OE1	2.47	0.44
3:BG:51:LEU:O	3:BG:55:GLU:HG3	2.18	0.44
3:HG:60:ARG:HE	3:HG:60:ARG:HB2	1.62	0.44
2:SG:282:ALA:HA	2:SG:285:ILE:HG12	2.00	0.44
2:G:234:PHE:CD2	2:G:313:ARG:HB3	2.53	0.44
3:M:117:VAL:O	3:M:215:PHE:HB2	2.18	0.44
2:T:141:GLN:HA	2:T:144:ASP:OD2	2.18	0.44
4:W:101:ILE:N	4:W:104:TYR:O	2.36	0.44
3:HA:243:ASP:HA	3:HA:246:TRP:CE3	2.53	0.44
3:HA:246:TRP:HB2	3:HA:247:ARG:HH12	1.83	0.44
2:MA:126:HIS:O	2:MA:130:ILE:HG23	2.17	0.44
3:NA:195:PRO:HD2	3:NA:196:ASN:N	2.32	0.44
3:TA:117:VAL:O	3:TA:215:PHE:HB2	2.17	0.44
2:YA:41:MET:HG2	2:YA:42:ALA:N	2.33	0.44
2:YA:107:ILE:HG22	2:YA:111:ASN:ND2	2.32	0.44
2:YA:282:ALA:HA	2:YA:285:ILE:HG12	2.00	0.44

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:EB:53:THR:O	2:EB:57:SER:OG	2.32	0.44
3:FB:102:LEU:HD22	3:FB:117:VAL:HG22	2.00	0.44
3:LB:101:ASN:HD22	3:LB:146:PHE:HE1	1.65	0.44
2:WB:97:ILE:O	2:WB:101:ARG:NH2	2.31	0.44
2:WB:298:LEU:HB2	2:WB:302:GLU:OE1	2.18	0.44
2:CC:219:ARG:NH1	2:CC:223:GLY:O	2.51	0.44
2:IC:282:ALA:HA	2:IC:285:ILE:HG12	2.00	0.44
2:OC:119:ALA:HB1	2:OC:149:PHE:CE1	2.53	0.44
3:PC:238:ASN:HB3	3:PC:241:HIS:ND1	2.33	0.44
3:PC:247:ARG:O	3:PC:247:ARG:HD2	2.18	0.44
3:VC:65:PHE:CE1	3:VC:161:ALA:HB2	2.53	0.44
2:AD:195:GLY:HA2	2:GD:161:ILE:O	2.18	0.44
2:GD:97:ILE:O	2:GD:100:THR:OG1	2.35	0.44
3:HD:219:LEU:HD23	3:HD:219:LEU:HA	1.80	0.44
4:UD:65:THR:O	4:UD:99:ILE:HA	2.17	0.44
2:YD:98:LEU:HB3	2:YD:108:GLU:OE1	2.18	0.44
2:YD:210:GLN:O	2:YD:214:VAL:HG12	2.17	0.44
2:EE:53:THR:O	2:EE:57:SER:OG	2.29	0.44
2:KE:207:LYS:CG	2:KE:210:GLN:HG2	2.43	0.44
2:QE:65:GLN:NE2	2:WE:47:ILE:H	2.16	0.44
2:QE:115:PRO:O	2:QE:119:ALA:HB2	2.18	0.44
2:WE:205:LEU:CD2	2:CF:140:SER:HA	2.48	0.44
2:WE:263:LEU:HD11	2:WE:274:PHE:HD2	1.82	0.44
3:XE:123:VAL:HG21	3:XE:146:PHE:CZ	2.53	0.44
2:CF:100:THR:OG1	2:CF:101:ARG:NH2	2.51	0.44
3:DF:195:PRO:HD2	3:DF:196:ASN:N	2.31	0.44
3:DF:229:GLU:OE1	3:DF:230:LEU:HD12	2.18	0.44
2:IF:282:ALA:O	2:IF:286:LEU:HD23	2.18	0.44
2:OF:246:SER:HB3	2:OF:321:MET:HE3	2.00	0.44
3:PF:236:LEU:HD23	3:PF:238:ASN:HD21	1.82	0.44
2:UF:107:ILE:O	2:UF:111:ASN:ND2	2.50	0.44
2:UF:282:ALA:HA	2:UF:285:ILE:HG12	1.99	0.44
3:VF:79:ILE:HG22	3:VF:204:PHE:HB3	1.99	0.44
3:BG:191:ILE:HG13	3:BG:192:THR:HG23	1.99	0.44
2:GG:263:LEU:HD11	2:GG:274:PHE:HD2	1.82	0.44
2:MG:298:LEU:HB2	2:MG:302:GLU:OE2	2.18	0.44
2:SG:210:GLN:O	2:SG:214:VAL:HG12	2.18	0.44
2:B:261:ILE:O	2:B:264:LYS:HG3	2.18	0.44
2:T:214:VAL:O	2:T:218:VAL:HG23	2.18	0.44
3:BA:73:LEU:HD23	3:BA:208:ILE:HD12	1.99	0.44
2:MA:186:GLN:NE2	3:TA:71:ASN:HA	2.31	0.44

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:SA:282:ALA:O	2:SA:286:LEU:HD23	2.18	0.44
3:ZA:185:GLN:HG2	3:ZA:187:LYS:HZ2	1.83	0.44
3:ZA:229:GLU:OE1	3:ZA:230:LEU:N	2.51	0.44
2:KB:133:ILE:O	2:KB:137:LEU:HG	2.17	0.44
2:KB:234:PHE:CD2	2:KB:313:ARG:HB3	2.53	0.44
2:QB:234:PHE:CD2	2:QB:313:ARG:HB3	2.53	0.44
3:RB:239:SER:O	3:RB:242:GLU:HG2	2.18	0.44
2:WB:261:ILE:O	2:WB:264:LYS:HG3	2.17	0.44
3:XB:236:LEU:HB3	3:XB:238:ASN:HD21	1.83	0.44
2:UC:121:LEU:HD22	3:BD:70:PHE:CE2	2.53	0.44
3:VC:240:ARG:HA	3:VC:240:ARG:CZ	2.48	0.44
2:GD:210:GLN:O	2:GD:214:VAL:HG12	2.17	0.44
3:HD:198:ILE:HD12	3:HD:198:ILE:HA	1.90	0.44
3:HD:240:ARG:HA	3:HD:240:ARG:CZ	2.48	0.44
2:MD:83:LYS:HA	2:SD:23:ALA:HB1	2.00	0.44
2:MD:107:ILE:O	2:MD:111:ASN:ND2	2.51	0.44
2:SD:209:GLN:O	2:SD:212:GLU:HG3	2.18	0.44
2:SD:298:LEU:O	2:SD:302:GLU:HG2	2.18	0.44
3:TD:195:PRO:HD2	3:TD:196:ASN:N	2.32	0.44
3:ZD:192:THR:OG1	3:ZD:194:SER:O	2.34	0.44
2:QE:134:LEU:HD13	2:QE:146:LEU:HD21	1.99	0.44
3:RE:202:THR:OG1	3:RE:217:ILE:HB	2.18	0.44
2:WE:282:ALA:O	2:WE:286:LEU:HD23	2.18	0.44
2:CF:84:ALA:HA	2:IF:27:LYS:HE2	1.98	0.44
2:CF:144:ASP:O	2:CF:148:LEU:HD12	2.17	0.44
2:CF:207:LYS:NZ	2:CF:209:GLN:OE1	2.50	0.44
2:IF:282:ALA:HA	2:IF:285:ILE:HG12	1.99	0.44
3:JF:240:ARG:HA	3:JF:240:ARG:CZ	2.48	0.44
2:UF:121:LEU:HG	3:BG:70:PHE:CZ	2.53	0.44
2:UF:186:GLN:HB3	3:BG:74:ARG:HH12	1.82	0.44
2:AG:218:VAL:HG12	2:AG:226:ALA:HB2	1.99	0.44
2:GG:207:LYS:HD3	2:GG:210:GLN:HE21	1.83	0.44
2:GG:282:ALA:HA	2:GG:285:ILE:HG12	2.00	0.44
3:HG:207:GLU:OE1	3:HG:207:GLU:N	2.51	0.44
2:SG:76:TYR:HD2	2:SG:79:SER:OG	2.00	0.44
3:TG:86:ILE:HD11	3:TG:221:PHE:HE1	1.81	0.44
2:B:41:MET:HG2	2:B:42:ALA:N	2.33	0.43
2:AA:196:VAL:HG12	2:AA:232:GLU:OE2	2.18	0.43
2:YA:14:LEU:HD12	2:YA:22:ALA:HB1	2.00	0.43
2:YA:76:TYR:CE2	2:EB:26:PHE:CE2	3.06	0.43
2:EB:159:LEU:HD21	3:FB:137:PHE:CD1	2.53	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:KB:103:THR:C	2:KB:106:GLY:H	2.21	0.43
3:LB:202:THR:OG1	3:LB:217:ILE:HB	2.18	0.43
2:QB:31:THR:HA	2:QB:34:VAL:CG2	2.48	0.43
2:QB:261:ILE:O	2:QB:264:LYS:HG3	2.18	0.43
3:XB:101:ASN:HB2	3:XB:146:PHE:HE2	1.83	0.43
2:CC:247:ILE:HD13	2:CC:273:LYS:HE3	2.00	0.43
3:JC:243:ASP:HA	3:JC:246:TRP:CE3	2.52	0.43
2:UC:85:LEU:HD22	2:UC:85:LEU:H	1.81	0.43
3:VC:101:ASN:HB2	3:VC:146:PHE:HE1	1.83	0.43
2:AD:11:VAL:O	2:AD:15:MET:HG2	2.17	0.43
2:AD:282:ALA:O	2:AD:286:LEU:HD23	2.18	0.43
3:ND:189:THR:OG1	3:ND:190:ASN:N	2.51	0.43
4:OD:65:THR:O	4:OD:99:ILE:HA	2.18	0.43
2:SD:11:VAL:O	2:SD:14:LEU:HB3	2.18	0.43
2:SD:167:VAL:HG21	3:TD:131:PHE:CD1	2.53	0.43
2:SD:210:GLN:O	2:SD:214:VAL:HG12	2.18	0.43
3:TD:109:PRO:O	3:TD:111:ARG:NH1	2.52	0.43
3:TD:169:TRP:CD1	3:TD:175:LEU:HD13	2.51	0.43
3:ZD:183:GLU:HG3	3:ZD:188:PHE:HB2	1.98	0.43
3:FE:86:ILE:HD12	3:FE:86:ILE:HA	1.93	0.43
3:FE:243:ASP:HA	3:FE:246:TRP:CE3	2.53	0.43
3:LE:101:ASN:HB2	3:LE:146:PHE:CE2	2.53	0.43
2:CF:14:LEU:HD12	2:CF:22:ALA:HB1	1.98	0.43
2:CF:132:THR:O	2:CF:135:VAL:HB	2.18	0.43
2:CF:190:ARG:O	2:IF:166:GLY:HA3	2.17	0.43
2:IF:167:VAL:HG21	3:JF:131:PHE:CD1	2.53	0.43
4:QF:65:THR:O	4:QF:99:ILE:HA	2.18	0.43
2:AG:163:THR:HG21	3:BG:134:ASP:OD2	2.18	0.43
2:AG:219:ARG:NH2	2:AG:230:ILE:HD12	2.33	0.43
2:SG:133:ILE:O	2:SG:137:LEU:HG	2.18	0.43
2:SG:171:ALA:HB1	3:TG:131:PHE:CZ	2.53	0.43
2:SG:247:ILE:HD13	2:SG:273:LYS:HE3	2.00	0.43
3:TG:65:PHE:CE1	3:TG:161:ALA:HB2	2.53	0.43
2:G:178:VAL:HG11	3:M:149:THR:HG22	2.00	0.43
2:G:311:ILE:HA	2:G:314:ARG:HG2	2.00	0.43
3:M:243:ASP:HA	3:M:246:TRP:CE3	2.52	0.43
2:J:298:LEU:O	2:J:302:GLU:HG2	2.18	0.43
2:T:75:GLU:O	2:T:79:SER:HB3	2.18	0.43
3:BA:190:ASN:O	3:HA:236:LEU:HD13	2.18	0.43
2:YA:144:ASP:O	2:YA:148:LEU:HD12	2.17	0.43
3:ZA:229:GLU:OE1	3:ZA:230:LEU:HD12	2.17	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:KB:107:ILE:O	2:KB:111:ASN:ND2	2.51	0.43
1:PB:543:ASP:HA	2:QB:21:ARG:HH11	1.83	0.43
2:QB:130:ILE:O	2:QB:134:LEU:HG	2.18	0.43
3:RB:285:PRO:HA	4:SB:119:GLY:HA2	2.00	0.43
2:WB:14:LEU:HD12	2:WB:22:ALA:HB1	1.99	0.43
2:WB:171:ALA:HB1	3:XB:131:PHE:CE2	2.53	0.43
3:XB:185:GLN:HG2	3:XB:187:LYS:HZ2	1.83	0.43
3:XB:198:ILE:HG13	3:XB:221:PHE:CD1	2.53	0.43
2:CC:234:PHE:CD2	2:CC:313:ARG:HB3	2.53	0.43
2:CC:282:ALA:O	2:CC:286:LEU:HD23	2.18	0.43
3:DC:237:GLU:HG2	3:DC:238:ASN:N	2.33	0.43
2:OC:187:ASN:N	3:VC:74:ARG:HH22	2.16	0.43
3:VC:108:LYS:HE2	3:VC:108:LYS:HB2	1.81	0.43
2:AD:298:LEU:O	2:AD:302:GLU:HG2	2.18	0.43
3:BD:117:VAL:O	3:BD:215:PHE:HB2	2.17	0.43
2:GD:83:LYS:HD2	2:MD:23:ALA:CA	2.46	0.43
2:GD:118:ALA:HA	2:GD:121:LEU:CD1	2.48	0.43
3:ND:111:ARG:NH1	3:ND:111:ARG:HB2	2.32	0.43
2:SD:250:LEU:O	2:SD:254:VAL:HG12	2.18	0.43
2:YD:113:MET:HE1	2:YD:118:ALA:HA	1.99	0.43
2:EE:205:LEU:CD2	2:KE:140:SER:HA	2.48	0.43
2:EE:282:ALA:O	2:EE:286:LEU:HD23	2.18	0.43
2:QE:66:PHE:HA	2:WE:45:ARG:O	2.18	0.43
2:QE:250:LEU:O	2:QE:254:VAL:HG12	2.19	0.43
3:RE:237:GLU:HG2	3:RE:238:ASN:N	2.32	0.43
2:WE:167:VAL:HG21	3:XE:131:PHE:CD1	2.53	0.43
3:XE:74:ARG:HD3	3:XE:74:ARG:HA	1.50	0.43
3:DF:198:ILE:HD12	3:DF:198:ILE:HA	1.90	0.43
2:IF:182:LEU:HD23	2:IF:182:LEU:HA	1.90	0.43
2:IF:195:GLY:HA2	2:OF:161:ILE:O	2.18	0.43
2:IF:207:LYS:NZ	2:IF:210:GLN:NE2	2.66	0.43
2:IF:234:PHE:CD2	2:IF:313:ARG:HB3	2.53	0.43
2:IF:261:ILE:H	2:IF:261:ILE:HG13	1.60	0.43
3:JF:185:GLN:HG3	3:JF:186:VAL:H	1.83	0.43
2:OF:5:SER:O	2:OF:9:LYS:HG3	2.18	0.43
2:OF:20:ASP:O	2:OF:23:ALA:HB3	2.17	0.43
2:OF:191:SER:OG	2:OF:192:LYS:N	2.51	0.43
3:PF:189:THR:OG1	3:PF:190:ASN:N	2.51	0.43
2:UF:168:GLN:HE21	2:UF:171:ALA:H	1.66	0.43
2:UF:307:ALA:O	2:UF:310:LEU:HG	2.18	0.43
2:GG:26:PHE:CZ	2:GG:34:VAL:HG22	2.52	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:MG:152:ARG:HH22	2:MG:153:LEU:HB2	1.83	0.43
4:QG:107:ALA:HB1	4:QG:124:ASP:O	2.18	0.43
2:SG:14:LEU:HD12	2:SG:22:ALA:HB1	2.00	0.43
2:SG:78:ARG:O	2:SG:82:VAL:HG23	2.18	0.43
2:SG:282:ALA:O	2:SG:286:LEU:HD23	2.18	0.43
3:M:127:VAL:HG23	3:M:153:VAL:HG11	2.00	0.43
3:K:79:ILE:HG22	3:K:204:PHE:HB3	1.99	0.43
2:T:21:ARG:HE	2:T:21:ARG:C	2.11	0.43
3:V:65:PHE:CE1	3:V:161:ALA:HB2	2.53	0.43
3:V:189:THR:OG1	3:V:190:ASN:N	2.51	0.43
2:AA:234:PHE:CD2	2:AA:313:ARG:HB3	2.53	0.43
3:BA:106:HIS:HE1	3:BA:113:THR:CG2	2.31	0.43
3:BA:150:GLU:O	3:BA:153:VAL:HG12	2.19	0.43
2:GA:67:ALA:O	2:GA:71:ILE:HG12	2.18	0.43
2:MA:167:VAL:HG21	3:NA:131:PHE:CD1	2.53	0.43
2:SA:85:LEU:H	2:SA:85:LEU:HD22	1.82	0.43
3:TA:65:PHE:CE1	3:TA:161:ALA:HB2	2.52	0.43
3:TA:225:GLU:HA	3:TA:228:ARG:HB3	2.00	0.43
2:YA:152:ARG:NH1	2:YA:153:LEU:HB2	2.30	0.43
2:EB:76:TYR:CE2	2:KB:38:SER:HB2	2.40	0.43
3:FB:180:VAL:HG23	3:FB:181:ARG:N	2.32	0.43
2:KB:282:ALA:O	2:KB:286:LEU:HD23	2.18	0.43
2:QB:282:ALA:O	2:QB:286:LEU:HD23	2.19	0.43
2:WB:37:LEU:HD12	2:WB:38:SER:N	2.33	0.43
2:CC:87:GLU:HG2	2:CC:89:ARG:N	2.33	0.43
3:PC:239:SER:HA	3:PC:242:GLU:OE1	2.19	0.43
2:UC:110:LEU:O	2:UC:113:MET:N	2.48	0.43
2:UC:282:ALA:HA	2:UC:285:ILE:HG12	2.00	0.43
3:BD:108:LYS:HE2	3:BD:108:LYS:HB2	1.83	0.43
2:GD:234:PHE:CD2	2:GD:313:ARG:HB3	2.53	0.43
2:MD:87:GLU:HG2	2:MD:89:ARG:N	2.33	0.43
2:MD:242:VAL:HG13	2:MD:323:ILE:HG12	1.99	0.43
2:SD:115:PRO:O	2:SD:119:ALA:CB	2.66	0.43
3:TD:69:LEU:O	3:TD:73:LEU:HB2	2.17	0.43
2:YD:282:ALA:O	2:YD:286:LEU:HD23	2.19	0.43
3:ZD:204:PHE:N	3:ZD:204:PHE:CD1	2.86	0.43
2:KE:172:LEU:HD12	2:KE:172:LEU:HA	1.86	0.43
3:LE:234:PRO:HA	3:LE:235:PRO:HD3	1.86	0.43
3:RE:219:LEU:HD23	3:RE:219:LEU:HA	1.82	0.43
2:CF:69:LEU:HD11	2:IF:47:ILE:CG2	2.46	0.43
2:IF:12:ILE:HD11	2:IF:55:VAL:HG11	1.99	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:IF:186:GLN:C	3:PF:74:ARG:HH22	2.22	0.43
3:VF:123:VAL:HG21	3:VF:146:PHE:CZ	2.53	0.43
2:AG:247:ILE:HD13	2:AG:273:LYS:HE3	2.00	0.43
3:HG:127:VAL:HG23	3:HG:153:VAL:HG11	2.00	0.43
2:MG:198:THR:CG2	2:SG:139:ARG:HH12	2.31	0.43
2:MG:221:PHE:CE2	2:SG:159:LEU:HD11	2.52	0.43
2:MG:307:ALA:O	2:MG:310:LEU:HG	2.18	0.43
3:TG:240:ARG:HA	3:TG:240:ARG:CZ	2.48	0.43
2:G:76:TYR:CE1	2:B:38:SER:HB3	2.51	0.43
3:M:189:THR:OG1	3:M:190:ASN:N	2.51	0.43
2:B:80:VAL:HA	2:B:83:LYS:HE3	2.01	0.43
1:Z:543:ASP:HA	2:AA:21:ARG:HH11	1.83	0.43
2:GA:31:THR:O	2:GA:34:VAL:HB	2.18	0.43
2:MA:110:LEU:O	2:MA:113:MET:N	2.48	0.43
2:YA:210:GLN:O	2:YA:214:VAL:HG12	2.19	0.43
2:YA:250:LEU:O	2:YA:254:VAL:HG12	2.19	0.43
2:EB:85:LEU:HD22	2:EB:85:LEU:H	1.82	0.43
2:EB:167:VAL:HG21	3:FB:131:PHE:CD1	2.53	0.43
2:EB:250:LEU:O	2:EB:254:VAL:HG12	2.19	0.43
2:KB:26:PHE:CZ	2:KB:34:VAL:HA	2.53	0.43
2:KB:246:SER:HB3	2:KB:321:MET:HE3	2.00	0.43
2:KB:261:ILE:O	2:KB:264:LYS:HG3	2.18	0.43
3:LB:294:ALA:O	3:LB:301:VAL:N	2.51	0.43
2:QB:108:GLU:HG3	2:QB:112:PHE:CE2	2.53	0.43
3:RB:123:VAL:HG21	3:RB:146:PHE:CZ	2.54	0.43
2:WB:21:ARG:HE	2:WB:21:ARG:C	2.13	0.43
2:CC:85:LEU:H	2:CC:85:LEU:HD22	1.83	0.43
3:JC:116:VAL:HG12	3:JC:217:ILE:CD1	2.49	0.43
3:JC:185:GLN:HG3	3:JC:186:VAL:H	1.82	0.43
4:QC:65:THR:O	4:QC:99:ILE:HA	2.19	0.43
2:AD:89:ARG:HH22	2:AD:93:LEU:HG	1.82	0.43
2:AD:282:ALA:HA	2:AD:285:ILE:HG12	2.00	0.43
2:MD:152:ARG:NH2	2:MD:153:LEU:HB2	2.33	0.43
2:SD:263:LEU:HD11	2:SD:274:PHE:HD2	1.83	0.43
2:SD:282:ALA:O	2:SD:286:LEU:HD23	2.18	0.43
3:TD:180:VAL:HG23	3:TD:181:ARG:H	1.83	0.43
2:YD:76:TYR:HE2	2:EE:26:PHE:CZ	2.37	0.43
2:YD:188:LEU:CD2	3:FE:74:ARG:HH11	2.30	0.43
2:YD:234:PHE:CD2	2:YD:313:ARG:HB3	2.53	0.43
2:YD:261:ILE:O	2:YD:264:LYS:HG3	2.18	0.43
3:ZD:123:VAL:HG21	3:ZD:146:PHE:CZ	2.53	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:EE:87:GLU:HG2	2:EE:89:ARG:H	1.82	0.43
2:EE:193:MET:O	2:KE:164:PHE:CD1	2.72	0.43
2:EE:282:ALA:HA	2:EE:285:ILE:HG12	1.99	0.43
3:XE:229:GLU:OE1	3:XE:230:LEU:N	2.52	0.43
4:EF:65:THR:O	4:EF:99:ILE:HA	2.17	0.43
2:IF:83:LYS:CG	2:IF:84:ALA:N	2.81	0.43
2:IF:87:GLU:HG2	2:IF:89:ARG:H	1.83	0.43
2:IF:158:MET:CE	2:IF:159:LEU:HD22	2.42	0.43
2:IF:246:SER:HB3	2:IF:321:MET:HE3	2.00	0.43
2:OF:234:PHE:CD2	2:OF:313:ARG:HB3	2.53	0.43
2:UF:11:VAL:O	2:UF:14:LEU:HB3	2.18	0.43
2:AG:282:ALA:HA	2:AG:285:ILE:HG12	2.00	0.43
2:AG:298:LEU:O	2:AG:302:GLU:HG2	2.18	0.43
3:BG:118:PHE:HE2	3:BG:158:LEU:HD12	1.83	0.43
2:GG:109:THR:O	2:GG:113:MET:HG3	2.18	0.43
2:MG:107:ILE:O	2:MG:111:ASN:ND2	2.50	0.43
2:MG:144:ASP:O	2:MG:148:LEU:HD12	2.18	0.43
3:NG:185:GLN:HG3	3:NG:186:VAL:H	1.83	0.43
4:UG:65:THR:O	4:UG:99:ILE:HA	2.19	0.43
2:G:108:GLU:HG3	2:G:112:PHE:HE2	1.83	0.43
3:M:123:VAL:HG21	3:M:146:PHE:CZ	2.53	0.43
2:B:234:PHE:CD2	2:B:313:ARG:HB3	2.54	0.43
2:T:72:ASN:ND2	2:AA:41:MET:HB2	2.33	0.43
2:T:78:ARG:O	2:T:82:VAL:HG12	2.18	0.43
3:BA:97:PRO:O	3:BA:100:THR:OG1	2.22	0.43
3:HA:69:LEU:O	3:HA:73:LEU:HB2	2.18	0.43
2:MA:282:ALA:O	2:MA:286:LEU:HD23	2.18	0.43
2:SA:87:GLU:HG2	2:SA:89:ARG:H	1.82	0.43
2:SA:206:MET:HE1	2:SA:214:VAL:HG11	2.01	0.43
2:SA:234:PHE:CD2	2:SA:313:ARG:HB3	2.54	0.43
3:TA:73:LEU:HD23	3:TA:208:ILE:HD13	2.00	0.43
2:YA:128:GLN:NE2	2:YA:164:PHE:HD2	2.09	0.43
3:ZA:189:THR:OG1	3:ZA:190:ASN:N	2.51	0.43
2:EB:30:SER:O	2:EB:34:VAL:HG23	2.18	0.43
2:EB:102:ASP:HB3	2:EB:107:ILE:HB	2.01	0.43
2:EB:130:ILE:HG13	2:EB:131:ALA:N	2.34	0.43
2:QB:205:LEU:CD2	2:WB:140:SER:HA	2.44	0.43
4:SB:65:THR:O	4:SB:99:ILE:HA	2.18	0.43
2:WB:282:ALA:O	2:WB:286:LEU:HD23	2.18	0.43
2:CC:214:VAL:O	2:CC:218:VAL:HG23	2.18	0.43
2:CC:230:ILE:HA	2:CC:233:MET:HE1	2.00	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:CC:282:ALA:HA	2:CC:285:ILE:HG12	2.01	0.43
3:DC:100:THR:HG23	3:DC:119:SER:HA	1.98	0.43
3:JC:101:ASN:HB2	3:JC:146:PHE:CE2	2.54	0.43
4:KC:65:THR:O	4:KC:99:ILE:HA	2.19	0.43
3:PC:108:LYS:HE2	3:PC:108:LYS:HB2	1.82	0.43
2:UC:158:MET:HE3	2:UC:159:LEU:HD22	2.00	0.43
2:UC:207:LYS:NZ	2:UC:209:GLN:OE1	2.51	0.43
2:AD:119:ALA:HB1	2:AD:149:PHE:CE1	2.53	0.43
3:ND:101:ASN:HB2	3:ND:146:PHE:CE2	2.53	0.43
3:ND:185:GLN:HG3	3:ND:186:VAL:H	1.84	0.43
2:SD:282:ALA:HA	2:SD:285:ILE:HG12	1.99	0.43
2:YD:115:PRO:O	2:YD:119:ALA:HB3	2.18	0.43
2:YD:144:ASP:O	2:YD:148:LEU:HD12	2.18	0.43
3:ZD:180:VAL:HG23	3:ZD:181:ARG:N	2.34	0.43
2:KE:105:SER:OG	2:KE:107:ILE:HD11	2.19	0.43
2:KE:108:GLU:HG3	2:KE:112:PHE:HE2	1.83	0.43
3:LE:75:ARG:HG3	3:LE:208:ILE:HD11	2.01	0.43
2:QE:83:LYS:HB2	2:WE:26:PHE:HB3	2.01	0.43
2:QE:174:GLU:HG2	3:RE:152:ARG:HG2	1.99	0.43
2:WE:210:GLN:O	2:WE:214:VAL:HG12	2.19	0.43
3:XE:204:PHE:N	3:XE:204:PHE:CD1	2.85	0.43
2:IF:108:GLU:HG3	2:IF:112:PHE:HE2	1.83	0.43
3:VF:127:VAL:HG23	3:VF:153:VAL:HG11	2.01	0.43
2:GG:65:GLN:NE2	2:MG:47:ILE:H	2.17	0.43
2:GG:203:ILE:HA	2:GG:206:MET:HE2	2.00	0.43
2:GG:250:LEU:O	2:GG:254:VAL:HG12	2.19	0.43
3:HG:75:ARG:HG3	3:HG:208:ILE:HD11	2.01	0.43
4:IG:65:THR:O	4:IG:99:ILE:HA	2.19	0.43
2:MG:234:PHE:CD2	2:MG:313:ARG:HB3	2.53	0.43
3:NG:79:ILE:HG22	3:NG:204:PHE:HB3	1.99	0.43
2:G:298:LEU:O	2:G:302:GLU:HG2	2.19	0.43
2:B:282:ALA:O	2:B:286:LEU:HD23	2.19	0.43
3:K:237:GLU:HG2	3:K:238:ASN:N	2.34	0.43
2:T:204:ASN:OD1	2:T:233:MET:HB3	2.17	0.43
2:T:234:PHE:CD2	2:T:313:ARG:HB3	2.53	0.43
2:T:250:LEU:O	2:T:254:VAL:HG12	2.19	0.43
2:AA:235:LEU:HD21	2:AA:237:GLU:HB3	2.00	0.43
3:BA:105:ILE:HD12	3:BA:105:ILE:HA	1.76	0.43
2:MA:89:ARG:CD	2:MA:90:ALA:H	2.14	0.43
2:MA:198:THR:O	2:MA:202:ILE:HG12	2.19	0.43
2:MA:234:PHE:CD2	2:MA:313:ARG:HB3	2.54	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:SA:89:ARG:CG	2:SA:90:ALA:N	2.80	0.43
2:SA:195:GLY:HA2	2:YA:161:ILE:O	2.18	0.43
2:SA:247:ILE:HD13	2:SA:273:LYS:HE3	2.00	0.43
2:SA:298:LEU:O	2:SA:302:GLU:HG2	2.19	0.43
2:YA:234:PHE:CD2	2:YA:313:ARG:HB3	2.54	0.43
3:ZA:96:LEU:HD23	3:ZA:97:PRO:CD	2.48	0.43
3:FB:122:LEU:CD1	3:FB:215:PHE:HB3	2.44	0.43
3:LB:207:GLU:N	3:LB:207:GLU:OE1	2.51	0.43
4:OB:104:TYR:O	3:RB:252:ARG:NH1	2.46	0.43
3:RB:150:GLU:O	3:RB:153:VAL:HG12	2.18	0.43
2:WB:76:TYR:HD2	2:CC:41:MET:CE	2.32	0.43
3:DC:79:ILE:HG22	3:DC:204:PHE:HB3	1.99	0.43
3:DC:211:LEU:H	3:DC:211:LEU:HG	1.67	0.43
2:IC:128:GLN:CG	3:JC:131:PHE:HB3	2.38	0.43
2:UC:250:LEU:O	2:UC:254:VAL:HG12	2.19	0.43
3:VC:198:ILE:HD12	3:VC:199:VAL:N	2.34	0.43
3:BD:58:ASN:OD1	3:BD:81:VAL:HG11	2.18	0.43
3:BD:127:VAL:HG23	3:BD:153:VAL:HG11	2.00	0.43
3:BD:154:ILE:HD13	3:BD:154:ILE:HA	1.87	0.43
3:BD:243:ASP:HA	3:BD:246:TRP:CE3	2.54	0.43
2:GD:83:LYS:HB3	2:MD:23:ALA:O	2.18	0.43
2:MD:298:LEU:O	2:MD:302:GLU:HG2	2.18	0.43
2:EE:172:LEU:HD12	2:EE:172:LEU:HA	1.73	0.43
2:EE:261:ILE:O	2:EE:264:LYS:HG3	2.18	0.43
2:KE:139:ARG:HE	2:KE:139:ARG:HB2	1.67	0.43
2:QE:34:VAL:O	2:QE:38:SER:OG	2.22	0.43
2:WE:234:PHE:CD2	2:WE:313:ARG:HB3	2.54	0.43
3:XE:294:ALA:O	3:XE:301:VAL:N	2.49	0.43
2:CF:203:ILE:HG23	2:CF:206:MET:HE1	2.01	0.43
3:JF:190:ASN:O	3:PF:236:LEU:HD13	2.18	0.43
3:PF:204:PHE:N	3:PF:204:PHE:CD1	2.86	0.43
3:PF:214:GLU:OE1	3:PF:215:PHE:N	2.51	0.43
2:AG:76:TYR:CE1	2:AG:80:VAL:HG13	2.53	0.43
2:GG:21:ARG:HE	2:GG:21:ARG:C	2.20	0.43
2:GG:167:VAL:HG21	3:HG:131:PHE:CD1	2.53	0.43
3:HG:58:ASN:OD1	3:HG:81:VAL:HG11	2.19	0.43
3:HG:180:VAL:HG23	3:HG:181:ARG:N	2.33	0.43
2:MG:103:THR:C	2:MG:106:GLY:H	2.21	0.43
2:B:219:ARG:NH2	2:B:230:ILE:HD12	2.34	0.43
2:J:21:ARG:HH21	2:J:24:GLU:HB2	1.82	0.43
2:J:234:PHE:CD2	2:J:313:ARG:HB3	2.54	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:K:204:PHE:CD1	3:K:204:PHE:N	2.86	0.43
3:K:240:ARG:HA	3:K:240:ARG:HH11	1.80	0.43
2:AA:282:ALA:O	2:AA:286:LEU:HD23	2.19	0.43
2:GA:20:ASP:O	2:GA:23:ALA:HB3	2.17	0.43
2:GA:282:ALA:O	2:GA:286:LEU:HD23	2.19	0.43
2:SA:235:LEU:HD21	2:SA:237:GLU:HB3	2.01	0.43
2:SA:282:ALA:HA	2:SA:285:ILE:HG12	2.01	0.43
3:TA:189:THR:OG1	3:TA:190:ASN:N	2.51	0.43
2:YA:207:LYS:CG	2:YA:210:GLN:HG2	2.43	0.43
2:EB:120:ASP:OD1	2:EB:123:ARG:NH1	2.52	0.43
2:EB:214:VAL:O	2:EB:218:VAL:HG23	2.19	0.43
3:LB:240:ARG:HA	3:LB:240:ARG:CZ	2.49	0.43
2:QB:111:ASN:OD1	2:QB:138:LYS:N	2.40	0.43
2:QB:311:ILE:HA	2:QB:314:ARG:HG2	2.01	0.43
3:RB:189:THR:OG1	3:RB:190:ASN:N	2.52	0.43
2:WB:167:VAL:HG21	3:XB:131:PHE:CD1	2.54	0.43
2:CC:201:GLU:HG2	2:IC:139:ARG:HH12	1.83	0.43
3:DC:189:THR:OG1	3:DC:190:ASN:N	2.51	0.43
4:EC:65:THR:O	4:EC:99:ILE:HA	2.18	0.43
2:IC:250:LEU:O	2:IC:254:VAL:HG12	2.19	0.43
2:OC:282:ALA:O	2:OC:286:LEU:HD23	2.19	0.43
2:UC:89:ARG:HG3	2:UC:89:ARG:NH1	2.33	0.43
3:ND:108:LYS:HE2	3:ND:108:LYS:HB2	1.81	0.43
3:ND:202:THR:OG1	3:ND:217:ILE:HB	2.19	0.43
2:SD:53:THR:O	2:SD:57:SER:OG	2.30	0.43
3:ZD:111:ARG:O	3:ZD:223:MET:HG2	2.19	0.43
2:KE:282:ALA:O	2:KE:286:LEU:HD23	2.18	0.43
2:QE:146:LEU:HD13	2:QE:146:LEU:HA	1.87	0.43
3:RE:185:GLN:HG2	3:RE:187:LYS:NZ	2.34	0.43
2:WE:186:GLN:OE1	2:WE:186:GLN:N	2.52	0.43
3:XE:77:PRO:HB3	3:XE:208:ILE:HB	2.01	0.43
3:XE:237:GLU:HG2	3:XE:238:ASN:N	2.32	0.43
2:CF:118:ALA:HA	2:CF:121:LEU:HG	2.00	0.43
2:CF:250:LEU:O	2:CF:254:VAL:HG12	2.19	0.43
2:IF:247:ILE:HD13	2:IF:273:LYS:HE3	2.00	0.43
2:OF:298:LEU:O	2:OF:302:GLU:HG2	2.19	0.43
3:PF:101:ASN:HD22	3:PF:146:PHE:HE2	1.66	0.43
2:UF:141:GLN:HA	2:UF:144:ASP:OD2	2.19	0.43
2:AG:118:ALA:HA	2:AG:121:LEU:CD1	2.48	0.43
2:MG:20:ASP:O	2:MG:23:ALA:HB3	2.19	0.43
2:MG:102:ASP:HB2	2:MG:108:GLU:HG2	2.00	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:G:206:MET:HG3	2:G:211:GLU:HG2	1.99	0.43
2:G:235:LEU:HD21	2:G:237:GLU:HB3	2.01	0.43
2:B:26:PHE:HD2	2:B:29:LEU:HD21	1.83	0.43
2:B:235:LEU:HD21	2:B:237:GLU:HB3	2.00	0.43
3:C:89:TYR:HE2	3:C:191:ILE:HG13	1.83	0.43
2:J:107:ILE:O	2:J:111:ASN:ND2	2.52	0.43
2:T:65:GLN:HG3	2:AA:46:GLN:HG2	1.99	0.43
2:GA:202:ILE:HA	2:MA:139:ARG:HH12	1.83	0.43
3:NA:65:PHE:CE1	3:NA:161:ALA:HB2	2.54	0.43
2:SA:80:VAL:HG12	2:YA:26:PHE:CZ	2.53	0.43
2:SA:219:ARG:NH2	2:SA:230:ILE:HD12	2.34	0.43
3:TA:204:PHE:CD1	3:TA:204:PHE:N	2.87	0.43
2:YA:9:LYS:O	2:YA:12:ILE:HG13	2.19	0.43
2:YA:114:GLU:HB2	2:YA:117:SER:OG	2.18	0.43
2:YA:235:LEU:HD21	2:YA:237:GLU:HB3	2.01	0.43
3:FB:243:ASP:HA	3:FB:246:TRP:CE3	2.54	0.43
2:KB:30:SER:O	2:KB:34:VAL:HG23	2.19	0.43
2:QB:12:ILE:HD11	2:QB:55:VAL:HG11	2.00	0.43
3:RB:51:LEU:O	3:RB:55:GLU:HG3	2.17	0.43
2:WB:130:ILE:HG13	2:WB:131:ALA:N	2.34	0.43
2:IC:144:ASP:O	2:IC:148:LEU:HD12	2.19	0.43
3:JC:58:ASN:OD1	3:JC:81:VAL:HG11	2.18	0.43
2:OC:31:THR:O	2:OC:34:VAL:HB	2.19	0.43
2:OC:198:THR:CG2	2:UC:139:ARG:HH12	2.32	0.43
4:SC:105:LEU:O	3:VC:252:ARG:NE	2.50	0.43
2:UC:27:LYS:O	2:UC:27:LYS:HG3	2.19	0.43
2:AD:178:VAL:HG11	3:BD:149:THR:HG22	2.00	0.43
2:AD:234:PHE:CD2	2:AD:313:ARG:HB3	2.54	0.43
3:BD:123:VAL:HG21	3:BD:146:PHE:CZ	2.54	0.43
3:HD:69:LEU:HD23	3:HD:69:LEU:HA	1.84	0.43
2:MD:130:ILE:HG13	2:MD:131:ALA:N	2.33	0.43
2:MD:234:PHE:CD2	2:MD:313:ARG:HB3	2.54	0.43
2:MD:250:LEU:O	2:MD:254:VAL:HG12	2.19	0.43
3:TD:239:SER:O	3:TD:242:GLU:HG2	2.18	0.43
3:ZD:194:SER:OG	3:ZD:197:ASP:N	2.51	0.43
3:FE:123:VAL:HG21	3:FE:146:PHE:CZ	2.53	0.43
2:KE:282:ALA:HA	2:KE:285:ILE:HG12	2.00	0.43
2:QE:282:ALA:HA	2:QE:285:ILE:HG12	2.00	0.43
3:RE:189:THR:OG1	3:RE:190:ASN:N	2.52	0.43
2:WE:76:TYR:CE2	2:CF:26:PHE:CE2	3.06	0.43
2:WE:188:LEU:HD23	3:DF:74:ARG:CZ	2.49	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:DF:86:ILE:HD11	3:DF:221:PHE:CZ	2.54	0.43
2:IF:84:ALA:O	2:IF:87:GLU:HB3	2.19	0.43
2:IF:261:ILE:O	2:IF:264:LYS:HG3	2.18	0.43
2:OF:282:ALA:O	2:OF:286:LEU:HD23	2.19	0.43
2:OF:311:ILE:HA	2:OF:314:ARG:HG2	2.01	0.43
3:PF:123:VAL:HG21	3:PF:146:PHE:CZ	2.54	0.43
2:UF:160:ARG:O	2:UF:164:PHE:HB2	2.19	0.43
2:UF:282:ALA:O	2:UF:286:LEU:HD23	2.18	0.43
2:AG:7:THR:CG2	2:AG:37:LEU:HD23	2.49	0.43
3:BG:86:ILE:HD11	3:BG:221:PHE:CE1	2.54	0.43
3:BG:214:GLU:OE1	3:BG:215:PHE:N	2.52	0.43
4:CG:65:THR:O	4:CG:99:ILE:HA	2.19	0.43
2:GG:65:GLN:HG3	2:MG:46:GLN:HG2	2.01	0.43
2:GG:282:ALA:O	2:GG:286:LEU:HD23	2.18	0.43
2:MG:89:ARG:HG3	2:MG:89:ARG:NH1	2.31	0.43
2:SG:234:PHE:CD2	2:SG:313:ARG:HB3	2.53	0.43
2:SG:311:ILE:HA	2:SG:314:ARG:HG2	2.01	0.43
3:TG:105:ILE:HD12	3:TG:105:ILE:HA	1.80	0.43
2:G:207:LYS:HZ2	2:G:207:LYS:HG3	1.55	0.43
2:T:14:LEU:HD12	2:T:22:ALA:HB1	2.01	0.43
3:V:249:ASN:O	3:V:252:ARG:HB3	2.19	0.43
2:AA:87:GLU:HG2	2:AA:89:ARG:N	2.34	0.43
2:AA:247:ILE:HD13	2:AA:273:LYS:HE3	2.00	0.43
3:BA:294:ALA:O	3:BA:301:VAL:N	2.50	0.43
2:GA:261:ILE:O	2:GA:264:LYS:HG3	2.18	0.43
2:MA:81:LEU:O	2:MA:85:LEU:HD13	2.19	0.43
2:SA:171:ALA:HB1	3:TA:131:PHE:CE2	2.54	0.43
2:SA:174:GLU:HG2	3:TA:152:ARG:HG2	2.00	0.43
3:ZA:101:ASN:HB2	3:ZA:146:PHE:CE2	2.54	0.43
3:ZA:204:PHE:N	3:ZA:204:PHE:CD1	2.86	0.43
3:FB:101:ASN:HB2	3:FB:146:PHE:CE2	2.54	0.43
2:KB:100:THR:OG1	2:KB:101:ARG:NH2	2.52	0.43
2:KB:122:ILE:HD12	2:KB:130:ILE:HG23	2.01	0.43
2:KB:126:HIS:O	2:KB:129:ILE:N	2.52	0.43
2:KB:163:THR:HG21	3:LB:134:ASP:OD2	2.19	0.43
2:KB:250:LEU:O	2:KB:254:VAL:HG12	2.19	0.43
2:WB:235:LEU:HB3	2:WB:238:ASN:ND2	2.34	0.43
3:XB:150:GLU:O	3:XB:153:VAL:HG12	2.19	0.43
4:YB:65:THR:O	4:YB:99:ILE:HA	2.19	0.43
3:DC:285:PRO:HA	4:EC:119:GLY:HA2	2.01	0.43
3:JC:96:LEU:HA	3:JC:97:PRO:HD3	1.91	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
4:SC:107:ALA:HB1	4:SC:124:ASP:O	2.18	0.43
2:UC:182:LEU:HD23	2:UC:182:LEU:HA	1.90	0.43
2:UC:234:PHE:CD2	2:UC:313:ARG:HB3	2.54	0.43
3:VC:204:PHE:N	3:VC:204:PHE:CD1	2.86	0.43
2:AD:87:GLU:HG2	2:AD:89:ARG:N	2.34	0.43
3:BD:204:PHE:N	3:BD:204:PHE:CD1	2.86	0.43
2:GD:11:VAL:HG11	2:GD:41:MET:HB3	2.01	0.43
2:GD:282:ALA:O	2:GD:286:LEU:HD23	2.19	0.43
2:GD:311:ILE:HA	2:GD:314:ARG:HG2	2.00	0.43
3:HD:51:LEU:O	3:HD:55:GLU:HG3	2.19	0.43
3:ZD:101:ASN:HB2	3:ZD:146:PHE:CE2	2.53	0.43
3:ZD:267:ASP:HA	4:AE:65:THR:HA	2.00	0.43
2:EE:17:ILE:HD12	2:EE:21:ARG:HG3	2.01	0.43
3:FE:237:GLU:HG2	3:FE:238:ASN:N	2.33	0.43
3:FE:294:ALA:O	3:FE:301:VAL:N	2.50	0.43
2:KE:14:LEU:HD12	2:KE:22:ALA:HB1	2.00	0.43
2:KE:87:GLU:HG2	2:KE:89:ARG:N	2.34	0.43
3:LE:101:ASN:HB2	3:LE:146:PHE:HE2	1.82	0.43
2:QE:203:ILE:O	2:QE:206:MET:HG3	2.18	0.43
3:RE:118:PHE:HE2	3:RE:158:LEU:HD12	1.84	0.43
3:XE:211:LEU:H	3:XE:211:LEU:HG	1.68	0.43
2:CF:209:GLN:O	2:CF:212:GLU:HG3	2.19	0.43
3:JF:101:ASN:HD22	3:JF:146:PHE:HE1	1.65	0.43
3:JF:239:SER:O	3:JF:242:GLU:HG2	2.19	0.43
2:OF:250:LEU:O	2:OF:254:VAL:HG12	2.19	0.43
2:UF:183:LEU:O	2:UF:186:GLN:N	2.51	0.43
2:SG:111:ASN:OD1	2:SG:141:GLN:NE2	2.52	0.43
3:TG:243:ASP:HA	3:TG:246:TRP:CE3	2.53	0.43
2:G:115:PRO:O	2:G:119:ALA:HB3	2.18	0.43
2:G:139:ARG:CZ	2:SG:201:GLU:HG3	2.49	0.43
2:G:167:VAL:HG21	3:M:131:PHE:CD1	2.54	0.43
3:C:211:LEU:H	3:C:211:LEU:HG	1.64	0.43
3:K:123:VAL:HG21	3:K:146:PHE:CZ	2.54	0.43
3:K:189:THR:OG1	3:K:190:ASN:N	2.52	0.43
3:BA:123:VAL:HG21	3:BA:146:PHE:CZ	2.54	0.43
2:YA:119:ALA:HB1	2:YA:149:PHE:HE1	1.83	0.43
2:EB:66:PHE:HA	2:KB:45:ARG:O	2.18	0.43
2:KB:172:LEU:HD12	2:KB:172:LEU:HA	1.71	0.43
3:LB:60:ARG:HE	3:LB:60:ARG:HB2	1.57	0.43
3:RB:58:ASN:OD1	3:RB:81:VAL:HG11	2.19	0.43
3:RB:69:LEU:O	3:RB:73:LEU:HB2	2.19	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:RB:237:GLU:HG2	3:RB:238:ASN:N	2.34	0.43
2:WB:209:GLN:O	2:WB:212:GLU:HG3	2.19	0.43
3:XB:190:ASN:ND2	3:DC:234:PRO:O	2.51	0.43
3:DC:162:LEU:HA	3:DC:162:LEU:HD13	1.79	0.43
2:IC:20:ASP:OD2	2:IC:21:ARG:N	2.52	0.43
3:JC:198:ILE:HG23	3:JC:221:PHE:HD1	1.84	0.43
3:JC:205:HIS:N	3:JC:205:HIS:ND1	2.66	0.43
2:OC:62:GLU:HG2	2:UC:45:ARG:HH22	1.83	0.43
3:VC:150:GLU:O	3:VC:153:VAL:HG12	2.19	0.43
2:AD:53:THR:O	2:AD:57:SER:OG	2.30	0.43
2:AD:103:THR:C	2:AD:106:GLY:H	2.22	0.43
2:GD:118:ALA:HA	2:GD:121:LEU:HD12	2.01	0.43
2:MD:41:MET:HG2	2:MD:42:ALA:N	2.34	0.43
2:SD:76:TYR:HD1	2:YD:41:MET:HE2	1.83	0.43
3:ZD:285:PRO:HA	4:AE:119:GLY:HA2	2.01	0.43
2:KE:77:LEU:O	2:KE:80:VAL:HG22	2.19	0.43
3:LE:108:LYS:HE2	3:LE:108:LYS:HB2	1.82	0.43
2:QE:298:LEU:O	2:QE:302:GLU:HG2	2.19	0.43
2:CF:172:LEU:HD12	2:CF:172:LEU:HA	1.69	0.43
3:DF:204:PHE:N	3:DF:204:PHE:CD1	2.86	0.43
2:IF:108:GLU:HA	2:IF:111:ASN:HD22	1.83	0.43
3:PF:69:LEU:O	3:PF:73:LEU:HB2	2.19	0.43
3:BG:285:PRO:HA	4:CG:119:GLY:HA2	2.01	0.43
2:MG:5:SER:O	2:MG:9:LYS:HG3	2.19	0.43
2:MG:119:ALA:HB1	2:MG:149:PHE:HE1	1.84	0.43
2:SG:32:ARG:HG3	2:SG:33:GLU:N	2.34	0.43
3:M:190:ASN:O	3:C:236:LEU:HD22	2.18	0.42
2:B:219:ARG:HH21	2:B:230:ILE:HD12	1.84	0.42
3:C:109:PRO:O	3:C:111:ARG:NH1	2.52	0.42
2:J:142:ALA:O	2:J:146:LEU:HD23	2.19	0.42
2:J:250:LEU:O	2:J:254:VAL:HG12	2.19	0.42
2:T:235:LEU:HD21	2:T:237:GLU:HB3	2.00	0.42
2:T:282:ALA:HA	2:T:285:ILE:HG12	2.01	0.42
2:T:298:LEU:O	2:T:302:GLU:HG2	2.18	0.42
2:AA:113:MET:HE1	2:AA:118:ALA:HA	2.00	0.42
3:BA:204:PHE:N	3:BA:204:PHE:CD1	2.85	0.42
3:BA:239:SER:O	3:BA:242:GLU:HG2	2.19	0.42
4:IA:65:THR:O	4:IA:99:ILE:HA	2.19	0.42
2:MA:11:VAL:O	2:MA:14:LEU:HB3	2.19	0.42
2:MA:298:LEU:HA	2:MA:301:VAL:HG23	2.00	0.42
3:NA:96:LEU:HA	3:NA:97:PRO:HD3	1.91	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:NA:117:VAL:O	3:NA:215:PHE:HB2	2.19	0.42
2:SA:250:LEU:O	2:SA:254:VAL:HG12	2.19	0.42
2:YA:182:LEU:HD23	2:YA:182:LEU:HA	1.91	0.42
3:ZA:111:ARG:HB2	3:ZA:111:ARG:HH11	1.83	0.42
3:ZA:237:GLU:HG2	3:ZA:238:ASN:N	2.34	0.42
2:KB:128:GLN:HG2	3:LB:131:PHE:CB	2.37	0.42
2:QB:128:GLN:NE2	2:QB:164:PHE:HD2	2.12	0.42
2:QB:134:LEU:HD23	2:QB:137:LEU:HD12	2.00	0.42
2:QB:235:LEU:HD21	2:QB:237:GLU:HB3	2.01	0.42
2:WB:182:LEU:HD23	2:WB:182:LEU:HA	1.89	0.42
3:XB:77:PRO:HB3	3:XB:208:ILE:HB	2.01	0.42
3:DC:214:GLU:OE1	3:DC:215:PHE:N	2.52	0.42
2:IC:100:THR:OG1	2:IC:101:ARG:NH2	2.52	0.42
2:IC:115:PRO:O	2:IC:119:ALA:HB3	2.19	0.42
3:JC:202:THR:OG1	3:JC:217:ILE:HB	2.19	0.42
3:JC:211:LEU:H	3:JC:211:LEU:HG	1.68	0.42
3:PC:62:ALA:O	3:PC:79:ILE:HD12	2.19	0.42
3:PC:198:ILE:HD12	3:PC:198:ILE:HA	1.89	0.42
2:AD:65:GLN:HG3	2:GD:46:GLN:HG2	2.00	0.42
2:GD:188:LEU:HD22	3:ND:74:ARG:NH1	2.34	0.42
3:HD:236:LEU:HB3	3:HD:238:ASN:ND2	2.34	0.42
2:MD:142:ALA:O	2:MD:146:LEU:HD23	2.19	0.42
3:ND:169:TRP:CD1	3:ND:175:LEU:HD13	2.46	0.42
2:SD:97:ILE:O	2:SD:100:THR:OG1	2.37	0.42
4:WD:107:ALA:CB	3:ZD:252:ARG:HH11	2.32	0.42
2:EE:20:ASP:O	2:EE:23:ALA:HB3	2.19	0.42
2:QE:76:TYR:OH	2:WE:34:VAL:HG13	2.19	0.42
3:RE:154:ILE:HD13	3:RE:154:ILE:HA	1.84	0.42
3:RE:204:PHE:N	3:RE:204:PHE:CD1	2.87	0.42
3:XE:113:THR:OG1	3:XE:191:ILE:HG22	2.19	0.42
2:CF:178:VAL:HG11	3:DF:149:THR:HG22	2.01	0.42
3:DF:69:LEU:O	3:DF:73:LEU:HB2	2.18	0.42
4:GF:106:ILE:O	3:JF:252:ARG:HG2	2.19	0.42
2:IF:17:ILE:HD12	2:IF:21:ARG:HB2	1.99	0.42
2:OF:119:ALA:HB1	2:OF:149:PHE:CE1	2.54	0.42
2:AG:246:SER:HB3	2:AG:321:MET:HE3	2.00	0.42
4:CG:101:ILE:N	4:CG:104:TYR:O	2.34	0.42
2:GG:115:PRO:O	2:GG:119:ALA:HB3	2.19	0.42
2:GG:298:LEU:HA	2:GG:301:VAL:HG23	2.01	0.42
3:HG:267:ASP:HA	4:IG:65:THR:HA	2.00	0.42
2:MG:87:GLU:HG2	2:MG:89:ARG:N	2.33	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:MG:193:MET:O	2:SG:164:PHE:CD1	2.72	0.42
2:MG:205:LEU:HG	2:SG:139:ARG:HB3	2.00	0.42
2:MG:235:LEU:HD21	2:MG:237:GLU:HB3	2.00	0.42
3:TG:96:LEU:HA	3:TG:97:PRO:HD3	1.92	0.42
2:G:282:ALA:O	2:G:286:LEU:HD23	2.18	0.42
4:Q:107:ALA:HB1	4:Q:124:ASP:O	2.19	0.42
2:B:144:ASP:O	2:B:148:LEU:HD12	2.19	0.42
3:K:59:GLU:OE1	3:K:59:GLU:N	2.52	0.42
2:T:20:ASP:O	2:T:23:ALA:HB3	2.19	0.42
2:AA:38:SER:OG	2:AA:39:THR:N	2.52	0.42
2:GA:234:PHE:CD2	2:GA:313:ARG:HB3	2.55	0.42
3:HA:123:VAL:HG21	3:HA:146:PHE:CZ	2.54	0.42
2:SA:83:LYS:HZ1	2:SA:84:ALA:HB2	1.85	0.42
3:TA:96:LEU:HA	3:TA:97:PRO:HD3	1.91	0.42
3:TA:128:ASP:OD1	3:TA:133:GLY:HA3	2.19	0.42
2:EB:210:GLN:O	2:EB:214:VAL:HG12	2.20	0.42
3:LB:189:THR:OG1	3:LB:190:ASN:N	2.52	0.42
2:WB:234:PHE:CD2	2:WB:313:ARG:HB3	2.54	0.42
2:CC:221:PHE:CE2	2:IC:159:LEU:HD21	2.53	0.42
2:CC:235:LEU:HD21	2:CC:237:GLU:HB3	2.01	0.42
2:IC:186:GLN:C	3:PC:74:ARG:HH22	2.22	0.42
2:IC:188:LEU:HD22	3:PC:74:ARG:NH1	2.35	0.42
3:JC:123:VAL:HG21	3:JC:146:PHE:CZ	2.55	0.42
2:OC:172:LEU:HD12	2:OC:172:LEU:HA	1.70	0.42
2:OC:250:LEU:O	2:OC:254:VAL:HG12	2.19	0.42
2:UC:186:GLN:C	3:BD:74:ARG:HH22	2.23	0.42
2:UC:282:ALA:O	2:UC:286:LEU:HD23	2.18	0.42
3:VC:189:THR:OG1	3:VC:190:ASN:N	2.52	0.42
2:MD:83:LYS:HB3	2:SD:27:LYS:HB2	2.01	0.42
3:ND:228:ARG:O	3:ND:232:VAL:HG23	2.19	0.42
2:SD:128:GLN:NE2	2:SD:164:PHE:CD2	2.87	0.42
2:KE:182:LEU:HA	2:KE:182:LEU:HD23	1.78	0.42
4:OE:106:ILE:HA	3:RE:252:ARG:CG	2.50	0.42
2:QE:188:LEU:HD21	3:XE:74:ARG:NE	2.34	0.42
3:XE:159:LYS:O	3:XE:159:LYS:HD2	2.19	0.42
3:XE:229:GLU:OE1	3:XE:230:LEU:HD12	2.19	0.42
2:CF:141:GLN:HA	2:CF:144:ASP:OD2	2.20	0.42
2:CF:234:PHE:CD2	2:CF:313:ARG:HB3	2.54	0.42
3:DF:58:ASN:OD1	3:DF:81:VAL:HG11	2.19	0.42
2:OF:65:GLN:HG3	2:UF:46:GLN:HG2	2.01	0.42
2:OF:113:MET:SD	2:OF:121:LEU:HD11	2.59	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:PF:54:LEU:HD13	3:PF:231:LEU:HD21	2.00	0.42
3:PF:267:ASP:HA	4:QF:65:THR:HA	2.01	0.42
4:WF:65:THR:O	4:WF:99:ILE:HA	2.19	0.42
2:AG:26:PHE:HA	2:AG:29:LEU:HG	2.00	0.42
2:AG:110:LEU:O	2:AG:113:MET:N	2.42	0.42
4:JG:65:THR:O	4:KG:99:ILE:HA	2.19	0.42
2:MG:311:ILE:HA	2:MG:314:ARG:HG2	2.00	0.42
2:G:46:GLN:HG2	2:SG:65:GLN:HG3	2.00	0.42
3:M:54:LEU:HB2	3:M:231:LEU:HD23	2.02	0.42
2:B:247:ILE:HD13	2:B:273:LYS:HE3	2.01	0.42
2:B:250:LEU:O	2:B:254:VAL:HG12	2.19	0.42
3:V:111:ARG:O	3:V:223:MET:HG2	2.19	0.42
4:EA:107:ALA:HB1	4:EA:124:ASP:O	2.20	0.42
2:GA:250:LEU:O	2:GA:254:VAL:HG12	2.19	0.42
2:MA:186:GLN:C	3:TA:74:ARG:HH22	2.23	0.42
2:SA:198:THR:O	2:SA:202:ILE:HG12	2.19	0.42
2:YA:21:ARG:HE	2:YA:21:ARG:C	2.21	0.42
3:FB:239:SER:HA	3:FB:242:GLU:OE1	2.19	0.42
2:KB:235:LEU:HD21	2:KB:237:GLU:HB3	2.00	0.42
2:QB:53:THR:O	2:QB:57:SER:OG	2.33	0.42
2:QB:151:GLU:HA	2:QB:154:ARG:HE	1.83	0.42
2:QB:174:GLU:HG2	3:RB:152:ARG:HG2	2.00	0.42
3:RB:54:LEU:HD13	3:RB:231:LEU:HD21	2.01	0.42
2:OC:178:VAL:HG11	3:PC:149:THR:HG22	2.00	0.42
2:OC:188:LEU:HD13	3:VC:74:ARG:CD	2.40	0.42
2:UC:31:THR:O	2:UC:34:VAL:HB	2.19	0.42
2:UC:100:THR:OG1	2:UC:101:ARG:NH2	2.52	0.42
2:UC:161:ILE:O	2:UC:164:PHE:HB3	2.19	0.42
4:YC:106:ILE:C	3:BD:252:ARG:HE	2.22	0.42
2:AD:182:LEU:HD23	2:AD:182:LEU:HA	1.92	0.42
2:GD:65:GLN:HG3	2:MD:46:GLN:HG2	2.02	0.42
2:GD:83:LYS:CD	2:MD:23:ALA:HA	2.47	0.42
2:GD:235:LEU:HD21	2:GD:237:GLU:HB3	2.01	0.42
2:MD:128:GLN:HG2	3:ND:131:PHE:CB	2.33	0.42
3:ND:123:VAL:HG21	3:ND:146:PHE:CZ	2.54	0.42
2:YD:190:ARG:O	2:EE:166:GLY:HA3	2.19	0.42
3:ZD:62:ALA:O	3:ZD:79:ILE:HD12	2.19	0.42
2:EE:171:ALA:HB1	3:FE:131:PHE:CE2	2.54	0.42
4:GE:65:THR:O	4:GE:99:ILE:HA	2.19	0.42
2:KE:53:THR:O	2:KE:57:SER:OG	2.32	0.42
2:KE:107:ILE:HG22	2:KE:111:ASN:ND2	2.33	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:KE:230:ILE:HA	2:KE:233:MET:HE2	2.01	0.42
2:KE:298:LEU:HA	2:KE:301:VAL:HG23	2.01	0.42
3:LE:294:ALA:O	3:LE:301:VAL:N	2.52	0.42
2:QE:107:ILE:HG22	2:QE:111:ASN:ND2	2.32	0.42
2:QE:234:PHE:CD2	2:QE:313:ARG:HB3	2.55	0.42
3:RE:117:VAL:O	3:RE:215:PHE:HB2	2.18	0.42
3:RE:198:ILE:HD12	3:RE:198:ILE:HA	1.89	0.42
2:CF:95:GLU:O	2:CF:99:GLU:HB2	2.18	0.42
3:DF:219:LEU:HD22	3:DF:223:MET:HE1	2.00	0.42
3:JF:55:GLU:HA	3:JF:58:ASN:ND2	2.34	0.42
2:OF:72:ASN:HD22	2:UF:41:MET:HE1	1.84	0.42
2:UF:83:LYS:HB2	2:AG:23:ALA:O	2.20	0.42
2:AG:113:MET:CE	2:AG:121:LEU:HD11	2.48	0.42
2:AG:221:PHE:HZ	3:HG:136:ARG:HH22	1.67	0.42
3:BG:294:ALA:O	3:BG:301:VAL:N	2.50	0.42
2:GG:178:VAL:HG11	3:HG:149:THR:HG22	2.01	0.42
2:GG:261:ILE:H	2:GG:261:ILE:HG13	1.60	0.42
3:HG:207:GLU:HG3	3:HG:212:THR:HG22	2.01	0.42
3:NG:202:THR:OG1	3:NG:217:ILE:HB	2.18	0.42
2:SG:107:ILE:HG22	2:SG:111:ASN:ND2	2.31	0.42
3:TG:198:ILE:HD12	3:TG:199:VAL:N	2.33	0.42
3:TG:237:GLU:HG2	3:TG:238:ASN:N	2.33	0.42
2:G:66:PHE:HA	2:B:45:ARG:O	2.19	0.42
2:G:164:PHE:CD1	2:SG:193:MET:O	2.72	0.42
2:G:247:ILE:HD13	2:G:273:LYS:HE3	2.01	0.42
3:M:180:VAL:HG23	3:M:181:ARG:N	2.35	0.42
2:B:193:MET:O	2:J:164:PHE:CD1	2.71	0.42
3:C:185:GLN:HG3	3:C:186:VAL:H	1.85	0.42
3:K:60:ARG:HE	3:K:60:ARG:HB2	1.55	0.42
2:T:67:ALA:O	2:T:71:ILE:HG12	2.18	0.42
2:AA:20:ASP:O	2:AA:23:ALA:HB3	2.18	0.42
3:BA:189:THR:OG1	3:BA:190:ASN:N	2.52	0.42
3:BA:194:SER:OG	3:BA:197:ASP:N	2.53	0.42
2:GA:41:MET:SD	2:GA:42:ALA:N	2.92	0.42
3:HA:219:LEU:HD23	3:HA:219:LEU:HA	1.88	0.42
3:ZA:69:LEU:O	3:ZA:73:LEU:HB2	2.18	0.42
2:KB:186:GLN:C	3:RB:74:ARG:HH22	2.22	0.42
2:KB:247:ILE:HD13	2:KB:273:LYS:HE3	2.00	0.42
3:LB:190:ASN:OD1	3:LB:190:ASN:O	2.37	0.42
3:LB:239:SER:O	3:LB:242:GLU:HG2	2.19	0.42
3:XB:285:PRO:HA	4:YB:119:GLY:HA2	2.01	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:CC:163:THR:HG21	3:DC:134:ASP:OD2	2.19	0.42
3:DC:51:LEU:O	3:DC:55:GLU:HG3	2.19	0.42
4:GC:106:ILE:O	3:JC:252:ARG:HG2	2.19	0.42
2:IC:69:LEU:HD11	2:OC:47:ILE:CG2	2.45	0.42
3:JC:58:ASN:OD1	3:JC:81:VAL:HG21	2.20	0.42
2:OC:186:GLN:C	3:VC:74:ARG:HH22	2.23	0.42
2:UC:69:LEU:HD12	2:UC:69:LEU:N	2.35	0.42
3:VC:86:ILE:O	3:VC:87:GLN:HG3	2.19	0.42
4:YC:106:ILE:CB	3:BD:252:ARG:HG2	2.50	0.42
2:AD:115:PRO:O	2:AD:119:ALA:HB3	2.18	0.42
2:AD:206:MET:SD	2:AD:207:LYS:HG2	2.60	0.42
2:GD:107:ILE:O	2:GD:111:ASN:ND2	2.52	0.42
3:HD:154:ILE:HD13	3:HD:154:ILE:HA	1.84	0.42
2:SD:193:MET:O	2:YD:164:PHE:CD1	2.72	0.42
3:TD:69:LEU:HD23	3:TD:69:LEU:HA	1.83	0.42
3:ZD:108:LYS:HE2	3:ZD:108:LYS:HB2	1.80	0.42
2:EE:7:THR:HG23	2:EE:37:LEU:HD23	2.02	0.42
2:EE:76:TYR:HE2	2:KE:26:PHE:CZ	2.38	0.42
2:EE:119:ALA:HB1	2:EE:149:PHE:CE1	2.55	0.42
2:EE:126:HIS:CD2	2:EE:127:PRO:HD2	2.55	0.42
2:EE:247:ILE:HD13	2:EE:273:LYS:HE3	1.99	0.42
2:QE:247:ILE:HD13	2:QE:273:LYS:HE3	2.00	0.42
2:WE:235:LEU:HD21	2:WE:237:GLU:HB3	2.00	0.42
2:WE:250:LEU:O	2:WE:254:VAL:HG12	2.19	0.42
2:WE:282:ALA:HA	2:WE:285:ILE:HG12	2.01	0.42
3:XE:198:ILE:HD12	3:XE:198:ILE:HA	1.90	0.42
2:CF:53:THR:O	2:CF:57:SER:OG	2.32	0.42
3:DF:294:ALA:O	3:DF:301:VAL:N	2.51	0.42
2:OF:8:ASP:OD1	2:OF:47:ILE:HD11	2.20	0.42
2:OF:14:LEU:HD12	2:OF:22:ALA:HB1	2.02	0.42
2:OF:107:ILE:O	2:OF:111:ASN:ND2	2.51	0.42
2:UF:311:ILE:HA	2:UF:314:ARG:HG2	2.00	0.42
2:AG:65:GLN:NE2	2:GG:47:ILE:H	2.17	0.42
2:GG:229:ILE:HD13	2:GG:229:ILE:HA	1.92	0.42
2:MG:221:PHE:HZ	3:TG:136:ARG:HH22	1.67	0.42
2:MG:285:ILE:HG13	2:MG:286:LEU:N	2.34	0.42
2:SG:95:GLU:O	2:SG:99:GLU:HB2	2.19	0.42
2:SG:250:LEU:O	2:SG:254:VAL:HG12	2.19	0.42
3:TG:294:ALA:O	3:TG:301:VAL:N	2.51	0.42
2:G:5:SER:O	2:G:9:LYS:HG3	2.19	0.42
3:C:154:ILE:HD13	3:C:154:ILE:HA	1.85	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:C:293:ILE:HA	3:C:302:LEU:O	2.19	0.42
2:T:163:THR:HG21	3:V:134:ASP:OD2	2.20	0.42
3:HA:204:PHE:N	3:HA:204:PHE:CD1	2.87	0.42
2:MA:235:LEU:HB3	2:MA:238:ASN:ND2	2.35	0.42
3:NA:267:ASP:HA	4:OA:65:THR:HA	2.01	0.42
3:TA:224:ILE:O	3:TA:228:ARG:N	2.52	0.42
2:EB:40:ALA:O	2:EB:44:VAL:HG23	2.19	0.42
3:RB:108:LYS:HE2	3:RB:108:LYS:HB2	1.83	0.42
2:CC:132:THR:O	2:CC:135:VAL:HB	2.19	0.42
2:CC:144:ASP:O	2:CC:148:LEU:HD12	2.19	0.42
3:JC:79:ILE:HG22	3:JC:204:PHE:HB3	2.01	0.42
3:PC:101:ASN:HB2	3:PC:146:PHE:CE2	2.54	0.42
3:BD:237:GLU:HG2	3:BD:238:ASN:N	2.34	0.42
2:GD:21:ARG:HH21	2:GD:25:VAL:N	2.16	0.42
2:YD:132:THR:O	2:YD:135:VAL:HB	2.20	0.42
2:EE:188:LEU:HD11	3:LE:74:ARG:HG3	2.01	0.42
2:KE:119:ALA:HB1	2:KE:149:PHE:HE1	1.84	0.42
2:CF:87:GLU:HG2	2:CF:89:ARG:N	2.35	0.42
2:UF:247:ILE:HD13	2:UF:273:LYS:HE3	2.01	0.42
3:VF:77:PRO:HB3	3:VF:208:ILE:HB	2.00	0.42
3:BG:123:VAL:HG21	3:BG:146:PHE:CZ	2.55	0.42
3:HG:116:VAL:HG12	3:HG:217:ILE:CD1	2.49	0.42
3:HG:123:VAL:HG21	3:HG:146:PHE:CZ	2.55	0.42
3:NG:154:ILE:HD13	3:NG:154:ILE:HA	1.86	0.42
2:SG:103:THR:C	2:SG:106:GLY:H	2.23	0.42
3:TG:194:SER:OG	3:TG:197:ASP:N	2.52	0.42
2:G:26:PHE:CE2	2:SG:76:TYR:CE2	3.08	0.42
2:G:98:LEU:HB3	2:G:108:GLU:OE1	2.19	0.42
3:M:96:LEU:HA	3:M:97:PRO:HD3	1.91	0.42
3:C:189:THR:OG1	3:C:190:ASN:N	2.53	0.42
2:T:20:ASP:OD2	2:T:21:ARG:N	2.51	0.42
2:T:182:LEU:HD23	2:T:182:LEU:HA	1.92	0.42
3:BA:89:TYR:HE2	3:BA:191:ILE:HG12	1.84	0.42
2:GA:214:VAL:O	2:GA:218:VAL:HG23	2.19	0.42
3:HA:198:ILE:HD12	3:HA:199:VAL:H	1.84	0.42
2:MA:201:GLU:O	2:SA:139:ARG:NH2	2.51	0.42
3:ZA:86:ILE:HD11	3:ZA:221:PHE:CE1	2.55	0.42
3:ZA:97:PRO:O	3:ZA:100:THR:OG1	2.34	0.42
3:FB:110:LEU:HG	3:FB:223:MET:HB3	2.01	0.42
3:FB:234:PRO:HA	3:FB:235:PRO:HD3	1.91	0.42
3:LB:111:ARG:O	3:LB:223:MET:HG2	2.20	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:QB:98:LEU:HB3	2:QB:108:GLU:OE1	2.20	0.42
2:QB:127:PRO:HA	2:QB:130:ILE:CG1	2.47	0.42
2:QB:128:GLN:NE2	2:QB:164:PHE:CD2	2.87	0.42
2:OC:235:LEU:HB3	2:OC:238:ASN:ND2	2.35	0.42
3:PC:75:ARG:HG3	3:PC:208:ILE:HD11	2.02	0.42
3:PC:195:PRO:HD2	3:PC:196:ASN:N	2.35	0.42
2:UC:188:LEU:HB3	3:BD:74:ARG:CZ	2.50	0.42
3:VC:194:SER:OG	3:VC:197:ASP:N	2.53	0.42
2:AD:65:GLN:NE2	2:GD:47:ILE:H	2.17	0.42
2:GD:31:THR:O	2:GD:34:VAL:HB	2.19	0.42
3:ND:195:PRO:HD2	3:ND:196:ASN:N	2.34	0.42
2:SD:234:PHE:CD2	2:SD:313:ARG:HB3	2.54	0.42
3:TD:293:ILE:HA	3:TD:302:LEU:O	2.20	0.42
2:YD:247:ILE:HD13	2:YD:273:LYS:HE3	2.00	0.42
3:ZD:195:PRO:HD2	3:ZD:196:ASN:N	2.34	0.42
2:EE:32:ARG:HG3	2:EE:33:GLU:N	2.35	0.42
3:FE:58:ASN:OD1	3:FE:81:VAL:HG11	2.20	0.42
3:FE:189:THR:OG1	3:FE:190:ASN:N	2.52	0.42
3:FE:204:PHE:N	3:FE:204:PHE:CD1	2.87	0.42
2:QE:151:GLU:HA	2:QE:154:ARG:HE	1.84	0.42
3:RE:90:HIS:O	3:RE:94:ARG:HG2	2.20	0.42
2:CF:5:SER:O	2:CF:9:LYS:HG3	2.20	0.42
3:DF:243:ASP:HA	3:DF:246:TRP:CE3	2.55	0.42
2:IF:214:VAL:O	2:IF:218:VAL:HG23	2.20	0.42
2:OF:89:ARG:HH22	2:OF:93:LEU:CD2	2.31	0.42
2:OF:247:ILE:HD13	2:OF:273:LYS:HE3	2.01	0.42
3:VF:75:ARG:HG3	3:VF:208:ILE:HD11	2.01	0.42
3:VF:150:GLU:O	3:VF:153:VAL:HG12	2.18	0.42
3:BG:190:ASN:OD1	3:HG:236:LEU:HB2	2.20	0.42
3:HG:214:GLU:OE1	3:HG:215:PHE:N	2.52	0.42
2:MG:247:ILE:HD13	2:MG:273:LYS:HE3	2.01	0.42
2:SG:193:MET:HG2	2:SG:194:GLY:N	2.35	0.42
2:J:103:THR:C	2:J:106:GLY:H	2.23	0.42
2:J:202:ILE:HD12	2:T:139:ARG:HE	1.84	0.42
2:J:218:VAL:HG12	2:J:226:ALA:HB2	2.02	0.42
2:AA:128:GLN:NE2	2:AA:164:PHE:HD2	2.10	0.42
2:GA:171:ALA:HB1	3:HA:131:PHE:CE2	2.54	0.42
3:HA:207:GLU:HG3	3:HA:212:THR:HG22	2.00	0.42
3:NA:134:ASP:OD2	3:NA:136:ARG:N	2.41	0.42
2:SA:132:THR:O	2:SA:135:VAL:HB	2.19	0.42
3:ZA:190:ASN:OD1	3:FB:236:LEU:HB2	2.20	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:EB:191:SER:OG	2:EB:192:LYS:N	2.53	0.42
2:KB:5:SER:O	2:KB:9:LYS:HG3	2.20	0.42
2:KB:298:LEU:O	2:KB:302:GLU:HG2	2.20	0.42
2:WB:138:LYS:HD3	2:WB:139:ARG:N	2.35	0.42
2:CC:41:MET:HG2	2:CC:42:ALA:N	2.34	0.42
2:CC:298:LEU:O	2:CC:302:GLU:HG2	2.18	0.42
3:DC:86:ILE:HD11	3:DC:221:PHE:CE1	2.53	0.42
2:IC:35:GLN:HA	2:IC:38:SER:OG	2.20	0.42
2:OC:5:SER:O	2:OC:9:LYS:HG3	2.20	0.42
2:OC:174:GLU:HG2	3:PC:152:ARG:HG2	2.01	0.42
2:OC:221:PHE:CE2	2:UC:159:LEU:HD11	2.54	0.42
2:UC:76:TYR:HD1	2:AD:41:MET:CE	2.33	0.42
2:UC:201:GLU:HG3	2:AD:139:ARG:CZ	2.50	0.42
3:VC:62:ALA:O	3:VC:79:ILE:HD12	2.19	0.42
3:VC:79:ILE:HG22	3:VC:204:PHE:HB3	2.02	0.42
3:BD:198:ILE:HD12	3:BD:198:ILE:HA	1.82	0.42
2:GD:250:LEU:O	2:GD:254:VAL:HG12	2.18	0.42
2:MD:151:GLU:HA	2:MD:154:ARG:HE	1.84	0.42
2:SD:163:THR:HG21	3:TD:134:ASP:OD2	2.19	0.42
2:SD:204:ASN:OD1	2:SD:233:MET:HB3	2.19	0.42
2:YD:235:LEU:HD21	2:YD:237:GLU:HB3	2.01	0.42
3:RE:123:VAL:HG21	3:RE:146:PHE:CZ	2.54	0.42
2:CF:66:PHE:HA	2:IF:45:ARG:O	2.19	0.42
3:DF:169:TRP:CD1	3:DF:175:LEU:HD13	2.52	0.42
3:DF:237:GLU:HG2	3:DF:238:ASN:N	2.34	0.42
3:JF:214:GLU:OE1	3:JF:215:PHE:N	2.52	0.42
2:OF:235:LEU:HD21	2:OF:237:GLU:HB3	2.01	0.42
3:PF:238:ASN:HB3	3:PF:241:HIS:ND1	2.35	0.42
3:PF:243:ASP:HA	3:PF:246:TRP:CE3	2.53	0.42
2:UF:115:PRO:O	2:UF:119:ALA:HB2	2.19	0.42
2:AG:83:LYS:HG3	2:AG:84:ALA:N	2.35	0.42
3:BG:237:GLU:HG2	3:BG:238:ASN:H	1.84	0.42
2:GG:247:ILE:HD13	2:GG:273:LYS:HE3	2.02	0.42
2:MG:137:LEU:HD13	2:MG:141:GLN:NE2	2.35	0.42
3:NG:150:GLU:O	3:NG:153:VAL:HG12	2.20	0.42
3:NG:180:VAL:HG23	3:NG:181:ARG:N	2.35	0.42
2:SG:85:LEU:HD22	2:SG:85:LEU:H	1.84	0.42
2:G:47:ILE:H	2:SG:65:GLN:NE2	2.18	0.42
3:M:237:GLU:HG2	3:M:238:ASN:H	1.84	0.42
2:J:62:GLU:O	2:J:65:GLN:HB3	2.20	0.42
2:J:244:ASP:OD1	2:J:276:ARG:NH2	2.53	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:T:81:LEU:O	2:T:85:LEU:HD13	2.20	0.42
2:AA:89:ARG:HH22	2:AA:93:LEU:HG	1.84	0.42
2:AA:155:HIS:O	2:AA:158:MET:HE3	2.20	0.42
3:BA:127:VAL:CG2	3:BA:153:VAL:HG11	2.49	0.42
2:MA:76:TYR:CE1	2:SA:26:PHE:HE2	2.38	0.42
2:MA:298:LEU:HA	2:MA:301:VAL:CG2	2.50	0.42
2:SA:40:ALA:O	2:SA:44:VAL:HG23	2.19	0.42
2:SA:89:ARG:NH2	2:YA:31:THR:HG23	2.35	0.42
2:SA:151:GLU:HA	2:SA:154:ARG:HE	1.84	0.42
2:YA:87:GLU:HG2	2:YA:89:ARG:H	1.85	0.42
2:EB:178:VAL:HG11	3:FB:149:THR:HG22	2.02	0.42
2:KB:40:ALA:O	2:KB:44:VAL:HG23	2.20	0.42
2:CC:188:LEU:HD22	3:JC:74:ARG:NH1	2.35	0.42
3:DC:122:LEU:CD1	3:DC:215:PHE:HB3	2.47	0.42
3:DC:234:PRO:HA	3:DC:235:PRO:HD3	1.86	0.42
3:PC:169:TRP:CD1	3:PC:175:LEU:HD13	2.49	0.42
2:UC:198:THR:O	2:UC:201:GLU:HG2	2.20	0.42
2:AD:188:LEU:HB3	3:HD:74:ARG:NE	2.35	0.42
2:GD:120:ASP:OD1	2:GD:123:ARG:NH1	2.53	0.42
2:SD:110:LEU:HB3	2:SD:141:GLN:NE2	2.32	0.42
3:ZD:239:SER:O	3:ZD:242:GLU:HG2	2.20	0.42
3:FE:55:GLU:HA	3:FE:58:ASN:HD22	1.85	0.42
3:FE:101:ASN:HB2	3:FE:146:PHE:CE2	2.55	0.42
3:FE:237:GLU:HG2	3:FE:238:ASN:H	1.85	0.42
2:KE:193:MET:O	2:QE:164:PHE:CD1	2.73	0.42
3:LE:195:PRO:HD2	3:LE:196:ASN:N	2.32	0.42
2:QE:20:ASP:OD1	2:QE:21:ARG:N	2.52	0.42
3:RE:229:GLU:OE1	3:RE:230:LEU:HD12	2.20	0.42
3:XE:219:LEU:HD23	3:XE:219:LEU:HA	1.79	0.42
3:JF:135:GLY:O	3:JF:211:LEU:HD21	2.20	0.42
3:JF:204:PHE:N	3:JF:204:PHE:CD1	2.87	0.42
3:VF:190:ASN:O	3:VF:190:ASN:OD1	2.38	0.42
2:AG:11:VAL:HG11	2:AG:41:MET:HB3	2.02	0.42
2:AG:102:ASP:HB3	2:AG:107:ILE:HB	2.01	0.42
2:AG:203:ILE:HA	2:AG:206:MET:HE3	2.01	0.42
2:AG:235:LEU:HD21	2:AG:237:GLU:HB3	2.01	0.42
3:HG:294:ALA:O	3:HG:301:VAL:N	2.50	0.42
2:MG:65:GLN:NE2	2:SG:47:ILE:O	2.48	0.42
2:G:47:ILE:CG2	2:SG:69:LEU:HD11	2.37	0.42
2:G:205:LEU:CD2	2:B:140:SER:HA	2.50	0.42
2:B:76:TYR:HE1	2:J:38:SER:HB2	1.85	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:B:107:ILE:HA	2:B:110:LEU:HD12	2.01	0.42
2:B:167:VAL:HG21	3:C:131:PHE:CD1	2.55	0.42
2:B:188:LEU:HD13	3:K:74:ARG:CD	2.42	0.42
2:B:218:VAL:HG12	2:B:226:ALA:HB2	2.02	0.42
2:J:89:ARG:NH2	2:J:93:LEU:HG	2.33	0.42
3:BA:195:PRO:HD2	3:BA:196:ASN:N	2.34	0.42
3:BA:267:ASP:HA	4:CA:65:THR:HA	2.01	0.42
2:MA:95:GLU:O	2:MA:99:GLU:HB2	2.20	0.42
2:SA:66:PHE:HA	2:YA:45:ARG:O	2.19	0.42
2:SA:79:SER:HA	2:SA:82:VAL:HG12	2.01	0.42
2:SA:107:ILE:O	2:SA:111:ASN:ND2	2.53	0.42
3:TA:154:ILE:HD13	3:TA:154:ILE:HA	1.85	0.42
3:ZA:154:ILE:HA	3:ZA:154:ILE:HD13	1.80	0.42
3:ZA:234:PRO:HA	3:ZA:235:PRO:HD3	1.89	0.42
2:EB:69:LEU:HD11	2:KB:47:ILE:CG2	2.48	0.42
3:LB:204:PHE:CD1	3:LB:204:PHE:N	2.87	0.42
3:LB:234:PRO:HA	3:LB:235:PRO:HD3	1.89	0.42
2:QB:206:MET:HG3	2:QB:211:GLU:HG2	2.01	0.42
2:QB:209:GLN:O	2:QB:212:GLU:HG3	2.20	0.42
3:RB:120:PRO:CD	3:RB:121:SER:N	2.81	0.42
2:OC:152:ARG:HH22	2:OC:153:LEU:HB2	1.83	0.42
3:PC:243:ASP:HA	3:PC:246:TRP:CE3	2.55	0.42
4:WC:65:THR:O	4:WC:99:ILE:HA	2.19	0.42
3:BD:219:LEU:HD23	3:BD:219:LEU:HA	1.86	0.42
3:HD:134:ASP:OD1	3:HD:136:ARG:N	2.27	0.42
2:MD:81:LEU:O	2:MD:85:LEU:HD13	2.20	0.42
2:MD:235:LEU:HD21	2:MD:237:GLU:HB3	2.01	0.42
2:SD:31:THR:O	2:SD:34:VAL:HB	2.20	0.42
2:SD:84:ALA:O	2:SD:87:GLU:HB3	2.19	0.42
2:SD:144:ASP:O	2:SD:148:LEU:HD12	2.19	0.42
3:TD:78:ASP:O	3:TD:206:VAL:HA	2.20	0.42
3:TD:189:THR:OG1	3:TD:190:ASN:N	2.53	0.42
2:YD:66:PHE:HA	2:EE:45:ARG:O	2.19	0.42
2:YD:261:ILE:H	2:YD:261:ILE:HG13	1.67	0.42
4:CE:106:ILE:H	3:FE:252:ARG:NH1	2.18	0.42
2:EE:74:ASN:O	2:EE:78:ARG:NE	2.53	0.42
2:WE:65:GLN:NE2	2:CF:47:ILE:H	2.18	0.42
3:XE:189:THR:OG1	3:XE:190:ASN:N	2.52	0.42
2:CF:127:PRO:HA	2:CF:130:ILE:CG1	2.46	0.42
2:CF:285:ILE:HG13	2:CF:286:LEU:N	2.35	0.42
2:OF:244:ASP:OD1	2:OF:276:ARG:NH2	2.53	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:BG:162:LEU:HD12	3:BG:177:VAL:CG1	2.44	0.42
3:BG:192:THR:OG1	3:BG:194:SER:O	2.34	0.42
2:MG:130:ILE:HD11	2:MG:157:VAL:HG22	2.01	0.42
3:M:101:ASN:HB2	3:M:146:PHE:CE2	2.55	0.42
2:B:298:LEU:O	2:B:302:GLU:HG2	2.19	0.42
2:J:206:MET:O	2:J:211:GLU:HG3	2.20	0.42
3:K:111:ARG:O	3:K:223:MET:HG2	2.20	0.42
3:V:107:LEU:HD11	3:V:219:LEU:HD21	2.01	0.42
2:AA:229:ILE:HD13	2:AA:229:ILE:HA	1.88	0.42
2:AA:259:LEU:HD11	2:AA:274:PHE:HE2	1.85	0.42
2:GA:65:GLN:HG3	2:MA:46:GLN:HG2	2.01	0.42
2:GA:128:GLN:NE2	2:GA:164:PHE:HD2	2.12	0.42
2:MA:105:SER:OG	2:MA:107:ILE:HD11	2.19	0.42
2:MA:146:LEU:HD21	2:MA:157:VAL:HG11	2.02	0.42
2:MA:188:LEU:HD22	3:TA:74:ARG:NH1	2.34	0.42
2:SA:202:ILE:CD1	2:YA:139:ARG:HH11	2.32	0.42
3:ZA:62:ALA:O	3:ZA:79:ILE:HD12	2.20	0.42
2:KB:191:SER:OG	2:KB:192:LYS:N	2.53	0.42
2:QB:69:LEU:HD12	2:WB:45:ARG:C	2.40	0.42
3:XB:111:ARG:O	3:XB:223:MET:HG2	2.20	0.42
3:XB:191:ILE:HG13	3:XB:192:THR:HG23	2.02	0.42
3:XB:198:ILE:HD12	3:XB:199:VAL:N	2.35	0.42
3:DC:123:VAL:HG21	3:DC:146:PHE:CZ	2.55	0.42
2:IC:235:LEU:HB3	2:IC:238:ASN:ND2	2.35	0.42
2:UC:5:SER:O	2:UC:9:LYS:HG3	2.20	0.42
2:AD:38:SER:OG	2:AD:39:THR:N	2.52	0.42
2:GD:5:SER:O	2:GD:9:LYS:HG3	2.20	0.42
3:HD:211:LEU:H	3:HD:211:LEU:HG	1.68	0.42
3:HD:293:ILE:HA	3:HD:302:LEU:O	2.19	0.42
2:MD:218:VAL:HG12	2:MD:226:ALA:HB2	2.02	0.42
2:SD:202:ILE:HD11	2:YD:139:ARG:HH11	1.85	0.42
3:TD:154:ILE:HD13	3:TD:154:ILE:HA	1.84	0.42
2:KE:235:LEU:HD21	2:KE:237:GLU:HB3	2.01	0.42
4:AF:107:ALA:HB1	4:AF:124:ASP:O	2.19	0.42
3:JF:86:ILE:HD11	3:JF:221:PHE:HE1	1.85	0.42
2:OF:196:VAL:HG12	2:OF:232:GLU:OE2	2.19	0.42
3:BG:128:ASP:OD1	3:BG:133:GLY:HA3	2.20	0.42
2:GG:5:SER:O	2:GG:9:LYS:HG3	2.20	0.42
2:GG:130:ILE:HD11	2:GG:157:VAL:HG22	2.02	0.42
3:NG:55:GLU:HA	3:NG:58:ASN:ND2	2.35	0.42
3:NG:123:VAL:HG21	3:NG:146:PHE:CZ	2.55	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:NG:204:PHE:N	3:NG:204:PHE:CD1	2.86	0.42
4:OG:65:THR:O	4:OG:99:ILE:HA	2.19	0.42
3:TG:79:ILE:HG22	3:TG:204:PHE:HB3	2.02	0.42
2:G:100:THR:OG1	2:G:101:ARG:NH2	2.53	0.41
2:B:87:GLU:HG2	2:B:89:ARG:N	2.35	0.41
3:K:54:LEU:HD13	3:K:231:LEU:HD21	2.02	0.41
3:K:106:HIS:NE2	3:K:113:THR:HG22	2.35	0.41
2:T:261:ILE:H	2:T:261:ILE:HG13	1.67	0.41
2:AA:202:ILE:HD11	2:GA:139:ARG:HH11	1.85	0.41
2:GA:5:SER:O	2:GA:9:LYS:HG3	2.19	0.41
3:HA:105:ILE:HG22	3:HA:114:GLY:O	2.20	0.41
2:MA:76:TYR:CE1	2:SA:26:PHE:CE2	3.08	0.41
3:NA:110:LEU:HG	3:NA:223:MET:HB3	2.02	0.41
2:EB:285:ILE:HG13	2:EB:286:LEU:N	2.35	0.41
3:FB:117:VAL:O	3:FB:215:PHE:HB2	2.19	0.41
3:FB:204:PHE:CD1	3:FB:204:PHE:N	2.86	0.41
4:GB:101:ILE:N	4:GB:104:TYR:O	2.36	0.41
2:KB:65:GLN:HG3	2:QB:46:GLN:HG2	2.02	0.41
3:LB:214:GLU:OE1	3:LB:215:PHE:N	2.53	0.41
2:QB:250:LEU:O	2:QB:254:VAL:HG12	2.19	0.41
2:WB:196:VAL:HG12	2:WB:232:GLU:OE2	2.20	0.41
2:CC:83:LYS:HG3	2:IC:23:ALA:HA	2.01	0.41
2:CC:130:ILE:HG13	2:CC:131:ALA:N	2.35	0.41
3:JC:102:LEU:HD22	3:JC:117:VAL:HG22	2.02	0.41
3:JC:117:VAL:O	3:JC:215:PHE:HB2	2.20	0.41
3:JC:135:GLY:O	3:JC:211:LEU:HD21	2.20	0.41
2:OC:193:MET:O	2:UC:164:PHE:CD1	2.72	0.41
3:PC:237:GLU:HG2	3:PC:238:ASN:N	2.34	0.41
2:UC:235:LEU:HD21	2:UC:237:GLU:HB3	2.01	0.41
3:VC:105:ILE:HG22	3:VC:114:GLY:O	2.20	0.41
2:AD:78:ARG:O	2:AD:82:VAL:HG23	2.20	0.41
2:AD:235:LEU:HD21	2:AD:237:GLU:HB3	2.01	0.41
3:BD:101:ASN:HB2	3:BD:146:PHE:CE2	2.54	0.41
2:GD:193:MET:O	2:MD:164:PHE:CD1	2.73	0.41
3:HD:294:ALA:O	3:HD:301:VAL:N	2.53	0.41
2:SD:183:LEU:O	2:SD:186:GLN:N	2.53	0.41
3:TD:101:ASN:HB2	3:TD:146:PHE:HE2	1.84	0.41
4:WD:107:ALA:N	3:ZD:252:ARG:HH11	2.18	0.41
2:YD:75:GLU:O	2:YD:79:SER:HB3	2.19	0.41
3:FE:128:ASP:OD1	3:FE:133:GLY:HA3	2.20	0.41
2:KE:141:GLN:HA	2:KE:144:ASP:OD2	2.20	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:QE:108:GLU:HG3	2:QE:112:PHE:HE2	1.84	0.41
2:QE:119:ALA:HB1	2:QE:149:PHE:CE1	2.55	0.41
3:RE:183:GLU:OE1	3:RE:184:MET:N	2.53	0.41
3:XE:86:ILE:HD11	3:XE:221:PHE:CE1	2.55	0.41
2:IF:235:LEU:HD21	2:IF:237:GLU:HB3	2.01	0.41
2:OF:79:SER:O	2:OF:82:VAL:HG22	2.19	0.41
2:OF:182:LEU:HD23	2:OF:182:LEU:HA	1.86	0.41
3:PF:58:ASN:OD1	3:PF:81:VAL:HG11	2.20	0.41
2:UF:199:ALA:HB3	2:UF:229:ILE:HD11	2.01	0.41
3:VF:60:ARG:HE	3:VF:60:ARG:HB2	1.61	0.41
4:YF:105:LEU:O	3:BG:252:ARG:HD3	2.19	0.41
2:AG:65:GLN:HG3	2:GG:46:GLN:HG2	2.02	0.41
3:BG:79:ILE:HG22	3:BG:204:PHE:HB3	2.02	0.41
3:HG:135:GLY:O	3:HG:211:LEU:HD21	2.19	0.41
1:LG:548:ALA:HB1	2:MG:70:ASN:HD22	1.85	0.41
3:NG:105:ILE:HA	3:NG:105:ILE:HD12	1.79	0.41
3:NG:195:PRO:HD2	3:NG:196:ASN:N	2.35	0.41
3:NG:240:ARG:HA	3:NG:240:ARG:NE	2.36	0.41
2:SG:235:LEU:HD21	2:SG:237:GLU:HB3	2.02	0.41
2:G:126:HIS:O	2:G:130:ILE:HG23	2.20	0.41
2:G:195:GLY:HA2	2:B:161:ILE:O	2.19	0.41
3:M:97:PRO:O	3:M:100:THR:OG1	2.20	0.41
3:M:204:PHE:N	3:M:204:PHE:CD1	2.86	0.41
2:B:118:ALA:HA	2:B:121:LEU:CD1	2.50	0.41
2:B:167:VAL:HG21	3:C:131:PHE:HA	2.01	0.41
2:J:167:VAL:HG21	3:K:131:PHE:CD1	2.55	0.41
2:J:171:ALA:HB1	3:K:131:PHE:CZ	2.55	0.41
3:K:55:GLU:HA	3:K:58:ASN:ND2	2.35	0.41
3:K:285:PRO:HA	4:L:119:GLY:HA2	2.02	0.41
3:BA:65:PHE:CZ	3:BA:161:ALA:HB2	2.55	0.41
3:HA:150:GLU:O	3:HA:154:ILE:HG12	2.20	0.41
2:MA:80:VAL:HA	2:MA:83:LYS:CD	2.47	0.41
2:YA:195:GLY:HA2	2:EB:161:ILE:O	2.20	0.41
3:ZA:237:GLU:HG2	3:ZA:238:ASN:H	1.85	0.41
2:EB:83:LYS:HG2	2:KB:26:PHE:HB3	2.03	0.41
2:KB:207:LYS:NZ	2:KB:210:GLN:HG3	2.25	0.41
3:LB:69:LEU:O	3:LB:73:LEU:HB2	2.20	0.41
2:QB:167:VAL:HG21	3:RB:131:PHE:CD1	2.54	0.41
2:WB:298:LEU:HA	2:WB:301:VAL:HG23	2.01	0.41
3:XB:106:HIS:CE1	3:XB:113:THR:CG2	2.96	0.41
3:XB:294:ALA:O	3:XB:301:VAL:N	2.52	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:IC:110:LEU:O	2:IC:113:MET:N	2.50	0.41
2:OC:247:ILE:HD13	2:OC:273:LYS:HE3	2.01	0.41
2:OC:261:ILE:H	2:OC:261:ILE:HG13	1.60	0.41
4:SC:106:ILE:HA	3:VC:252:ARG:HG3	2.02	0.41
2:UC:225:LEU:HA	2:UC:228:LYS:HD2	2.01	0.41
2:AD:127:PRO:HA	2:AD:130:ILE:HG12	2.02	0.41
2:MD:108:GLU:HA	2:MD:111:ASN:HD22	1.84	0.41
2:MD:191:SER:OG	2:MD:192:LYS:N	2.53	0.41
2:MD:210:GLN:O	2:MD:214:VAL:HG12	2.20	0.41
2:SD:193:MET:HG2	2:SD:194:GLY:N	2.36	0.41
2:SD:206:MET:SD	2:SD:210:GLN:HB3	2.60	0.41
2:YD:40:ALA:O	2:YD:44:VAL:HG23	2.19	0.41
2:YD:182:LEU:HD23	2:YD:182:LEU:HA	1.80	0.41
2:KE:110:LEU:O	2:KE:113:MET:HB2	2.19	0.41
2:KE:186:GLN:NE2	3:RE:74:ARG:HH22	2.17	0.41
2:KE:198:THR:O	2:KE:202:ILE:HG12	2.20	0.41
3:LE:86:ILE:HD11	3:LE:221:PHE:HE1	1.84	0.41
2:QE:132:THR:O	2:QE:135:VAL:HB	2.21	0.41
3:XE:214:GLU:OE1	3:XE:215:PHE:N	2.54	0.41
2:CF:206:MET:CE	2:CF:214:VAL:HG11	2.50	0.41
3:DF:101:ASN:HD22	3:DF:146:PHE:HE2	1.68	0.41
2:OF:199:ALA:HB3	2:OF:229:ILE:HD11	2.02	0.41
2:UF:76:TYR:HB2	2:AG:41:MET:HE3	2.01	0.41
2:UF:261:ILE:O	2:UF:264:LYS:HG3	2.21	0.41
3:BG:54:LEU:HD13	3:BG:231:LEU:HD21	2.02	0.41
2:MG:100:THR:OG1	2:MG:101:ARG:NH2	2.53	0.41
2:MG:261:ILE:O	2:MG:264:LYS:HG3	2.20	0.41
3:NG:237:GLU:HG2	3:NG:238:ASN:N	2.36	0.41
2:G:9:LYS:HG2	2:G:55:VAL:HG22	2.01	0.41
2:B:5:SER:O	2:B:9:LYS:HG3	2.20	0.41
3:C:101:ASN:HB2	3:C:146:PHE:HE2	1.86	0.41
3:C:105:ILE:HG22	3:C:114:GLY:O	2.21	0.41
2:J:141:GLN:HA	2:J:144:ASP:OD2	2.20	0.41
2:J:235:LEU:HD21	2:J:237:GLU:HB3	2.01	0.41
3:K:116:VAL:HG21	3:K:158:LEU:HD11	2.01	0.41
3:K:239:SER:O	3:K:242:GLU:HG2	2.20	0.41
3:V:198:ILE:HD12	3:V:198:ILE:HA	1.88	0.41
3:V:214:GLU:OE1	3:V:215:PHE:N	2.53	0.41
3:BA:110:LEU:HG	3:BA:223:MET:HB3	2.01	0.41
3:BA:111:ARG:O	3:BA:223:MET:HG2	2.21	0.41
2:GA:95:GLU:O	2:GA:99:GLU:HB2	2.21	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:HA:58:ASN:OD1	3:HA:81:VAL:HG11	2.19	0.41
3:HA:237:GLU:HG2	3:HA:238:ASN:H	1.86	0.41
2:MA:83:LYS:HA	2:SA:23:ALA:HB1	2.02	0.41
2:MA:207:LYS:NZ	2:MA:209:GLN:HB3	2.35	0.41
2:MA:247:ILE:HD13	2:MA:273:LYS:HE3	2.02	0.41
2:SA:218:VAL:HG12	2:SA:226:ALA:HB2	2.03	0.41
2:EB:76:TYR:HD2	2:KB:41:MET:CE	2.33	0.41
2:KB:83:LYS:HE2	2:KB:83:LYS:HB2	1.65	0.41
3:RB:224:ILE:O	3:RB:228:ARG:N	2.53	0.41
2:WB:40:ALA:O	2:WB:44:VAL:HG23	2.19	0.41
3:XB:198:ILE:HD12	3:XB:199:VAL:H	1.86	0.41
2:CC:188:LEU:HB3	3:JC:74:ARG:CZ	2.51	0.41
2:IC:172:LEU:HD12	2:IC:172:LEU:HA	1.74	0.41
2:IC:247:ILE:HD13	2:IC:273:LYS:HE3	2.02	0.41
2:OC:40:ALA:O	2:OC:44:VAL:HG23	2.19	0.41
2:OC:62:GLU:HG2	2:UC:45:ARG:NH2	2.36	0.41
2:OC:119:ALA:HB1	2:OC:149:PHE:HE1	1.84	0.41
2:UC:191:SER:OG	2:UC:192:LYS:N	2.52	0.41
2:AD:247:ILE:HD13	2:AD:273:LYS:HE3	2.01	0.41
2:MD:207:LYS:HE2	2:MD:207:LYS:HB2	1.80	0.41
3:ND:204:PHE:N	3:ND:204:PHE:CD1	2.87	0.41
2:SD:137:LEU:HB3	2:SD:141:GLN:OE1	2.19	0.41
3:TD:108:LYS:HE2	3:TD:108:LYS:HB2	1.83	0.41
2:YD:298:LEU:O	2:YD:302:GLU:HG2	2.20	0.41
2:EE:298:LEU:O	2:EE:302:GLU:HG2	2.20	0.41
3:LE:219:LEU:HD23	3:LE:219:LEU:HA	1.86	0.41
2:WE:38:SER:HA	2:WE:41:MET:HE2	2.03	0.41
2:CF:40:ALA:O	2:CF:44:VAL:HG23	2.21	0.41
2:CF:108:GLU:HG3	2:CF:112:PHE:CE2	2.54	0.41
3:DF:211:LEU:H	3:DF:211:LEU:HG	1.68	0.41
2:IF:35:GLN:O	2:IF:38:SER:HB2	2.19	0.41
2:IF:113:MET:HE2	2:IF:113:MET:HB3	1.93	0.41
2:IF:188:LEU:HD13	3:PF:74:ARG:CD	2.50	0.41
3:JF:237:GLU:HG2	3:JF:238:ASN:N	2.36	0.41
2:OF:188:LEU:CD2	3:VF:74:ARG:HH11	2.20	0.41
2:UF:89:ARG:HH12	2:AG:31:THR:HG23	1.85	0.41
2:AG:130:ILE:HG13	2:AG:131:ALA:N	2.35	0.41
3:BG:237:GLU:HG2	3:BG:238:ASN:N	2.35	0.41
3:HG:88:PRO:HG2	3:HG:91:GLU:HB2	2.02	0.41
2:MG:12:ILE:HD11	2:MG:55:VAL:HG11	2.00	0.41
2:SG:128:GLN:NE2	2:SG:164:PHE:HD2	2.18	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:G:76:TYR:OH	2:B:34:VAL:HG13	2.20	0.41
3:C:204:PHE:N	3:C:204:PHE:CD1	2.87	0.41
3:NA:88:PRO:HG2	3:NA:91:GLU:HB2	2.01	0.41
3:NA:294:ALA:O	3:NA:301:VAL:N	2.53	0.41
4:QA:107:ALA:HB1	4:QA:124:ASP:O	2.21	0.41
3:TA:51:LEU:O	3:TA:55:GLU:HG3	2.20	0.41
3:ZA:101:ASN:HD22	3:ZA:146:PHE:HE2	1.68	0.41
3:ZA:159:LYS:HD2	3:ZA:159:LYS:O	2.20	0.41
2:EB:107:ILE:O	2:EB:111:ASN:ND2	2.52	0.41
3:FB:224:ILE:O	3:FB:228:ARG:N	2.54	0.41
2:KB:69:LEU:HD23	2:QB:47:ILE:CG2	2.46	0.41
2:KB:119:ALA:HB1	2:KB:149:PHE:HE1	1.84	0.41
2:KB:132:THR:O	2:KB:135:VAL:HB	2.21	0.41
2:KB:188:LEU:HD22	3:RB:74:ARG:NH1	2.36	0.41
3:LB:108:LYS:HE2	3:LB:108:LYS:HB2	1.82	0.41
3:RB:195:PRO:HD2	3:RB:196:ASN:N	2.35	0.41
3:RB:211:LEU:H	3:RB:211:LEU:HG	1.68	0.41
2:WB:149:PHE:CD2	2:WB:153:LEU:HD22	2.55	0.41
2:WB:197:ARG:NH1	2:WB:232:GLU:HB3	2.35	0.41
3:XB:189:THR:OG1	3:XB:190:ASN:N	2.53	0.41
2:CC:32:ARG:NH2	2:CC:33:GLU:OE1	2.43	0.41
3:DC:66:ARG:HG3	3:DC:77:PRO:O	2.21	0.41
3:DC:86:ILE:HD11	3:DC:221:PHE:CZ	2.56	0.41
2:OC:128:GLN:HG2	3:PC:131:PHE:CB	2.45	0.41
2:OC:214:VAL:O	2:OC:218:VAL:HG23	2.20	0.41
2:OC:234:PHE:CD2	2:OC:313:ARG:HB3	2.55	0.41
3:HD:169:TRP:CD1	3:HD:175:LEU:HD13	2.47	0.41
2:MD:243:ASP:O	2:MD:247:ILE:HG23	2.20	0.41
2:SD:235:LEU:HD21	2:SD:237:GLU:HB3	2.02	0.41
2:YD:141:GLN:HA	2:YD:144:ASP:OD2	2.20	0.41
3:ZD:219:LEU:HD23	3:ZD:219:LEU:HA	1.80	0.41
2:EE:202:ILE:CD1	2:KE:139:ARG:HD3	2.50	0.41
3:FE:77:PRO:HB3	3:FE:208:ILE:HB	2.02	0.41
3:FE:150:GLU:O	3:FE:154:ILE:HG12	2.20	0.41
2:KE:167:VAL:HG21	3:LE:131:PHE:CD1	2.55	0.41
2:QE:65:GLN:HG3	2:WE:46:GLN:HG2	2.02	0.41
3:RE:243:ASP:HA	3:RE:246:TRP:CE3	2.55	0.41
2:WE:53:THR:O	2:WE:57:SER:OG	2.33	0.41
4:ZE:65:THR:O	4:AF:99:ILE:HA	2.20	0.41
2:CF:261:ILE:H	2:CF:261:ILE:HG13	1.67	0.41
3:JF:86:ILE:HD11	3:JF:221:PHE:CE1	2.55	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:JF:294:ALA:O	3:JF:301:VAL:N	2.52	0.41
2:OF:119:ALA:HB1	2:OF:149:PHE:HE1	1.85	0.41
2:UF:65:GLN:HG3	2:AG:46:GLN:HG2	2.03	0.41
3:VF:237:GLU:HG2	3:VF:238:ASN:N	2.36	0.41
4:EG:106:ILE:O	3:HG:252:ARG:HG2	2.21	0.41
2:MG:14:LEU:HA	2:MG:17:ILE:HG12	2.02	0.41
2:MG:119:ALA:HB1	2:MG:149:PHE:CE1	2.55	0.41
2:SG:171:ALA:HB1	3:TG:131:PHE:CE2	2.55	0.41
3:TG:122:LEU:CD1	3:TG:215:PHE:HB3	2.46	0.41
3:TG:204:PHE:N	3:TG:204:PHE:CD1	2.86	0.41
3:TG:237:GLU:HG2	3:TG:238:ASN:H	1.85	0.41
2:G:30:SER:OG	2:G:33:GLU:OE1	2.34	0.41
2:G:166:GLY:HA3	2:SG:190:ARG:O	2.20	0.41
3:C:236:LEU:HD12	3:C:236:LEU:HA	1.92	0.41
3:K:294:ALA:O	3:K:301:VAL:N	2.52	0.41
2:GA:188:LEU:HD22	3:NA:74:ARG:NH1	2.35	0.41
3:HA:128:ASP:OD1	3:HA:133:GLY:HA3	2.20	0.41
2:SA:37:LEU:HD12	2:SA:38:SER:N	2.35	0.41
3:TA:150:GLU:O	3:TA:153:VAL:HG12	2.20	0.41
2:EB:5:SER:O	2:EB:9:LYS:HG3	2.20	0.41
2:EB:188:LEU:HB3	3:LB:74:ARG:CZ	2.50	0.41
3:LB:183:GLU:CD	3:LB:185:GLN:H	2.24	0.41
2:WB:89:ARG:NH2	2:CC:31:THR:HG23	2.36	0.41
3:XB:127:VAL:HG23	3:XB:153:VAL:HG11	2.01	0.41
2:CC:53:THR:O	2:CC:57:SER:OG	2.31	0.41
2:IC:130:ILE:HD11	2:IC:157:VAL:HG22	2.03	0.41
2:IC:298:LEU:HA	2:IC:301:VAL:CG2	2.50	0.41
4:LC:65:THR:O	4:MC:99:ILE:HA	2.20	0.41
4:MC:105:LEU:O	3:PC:252:ARG:NE	2.52	0.41
2:UC:247:ILE:HD13	2:UC:273:LYS:HE3	2.01	0.41
3:VC:195:PRO:HD2	3:VC:196:ASN:N	2.35	0.41
2:AD:5:SER:O	2:AD:9:LYS:HG3	2.20	0.41
2:AD:72:ASN:O	2:GD:41:MET:HE1	2.20	0.41
2:GD:87:GLU:HG2	2:GD:89:ARG:N	2.36	0.41
2:EE:5:SER:O	2:EE:9:LYS:HG3	2.20	0.41
2:EE:99:GLU:O	2:EE:103:THR:HG23	2.20	0.41
2:EE:128:GLN:NE2	2:EE:164:PHE:HD2	2.13	0.41
2:EE:207:LYS:NZ	2:EE:209:GLN:HB3	2.36	0.41
2:WE:247:ILE:HD13	2:WE:273:LYS:HE3	2.02	0.41
3:XE:202:THR:HG1	3:XE:204:PHE:HE1	1.65	0.41
3:XE:243:ASP:HA	3:XE:246:TRP:CE3	2.55	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:DF:96:LEU:HA	3:DF:97:PRO:HD3	1.92	0.41
3:DF:111:ARG:NH2	3:DF:222:SER:O	2.53	0.41
2:IF:5:SER:O	2:IF:9:LYS:HG3	2.20	0.41
2:IF:26:PHE:HE2	2:IF:37:LEU:HD11	1.86	0.41
2:IF:76:TYR:HD1	2:IF:79:SER:OG	2.03	0.41
2:IF:188:LEU:H	3:PF:74:ARG:NH2	2.18	0.41
2:IF:298:LEU:O	2:IF:302:GLU:HG2	2.20	0.41
3:JF:69:LEU:O	3:JF:73:LEU:HB2	2.20	0.41
2:OF:202:ILE:CD1	2:UF:139:ARG:HH11	2.34	0.41
2:AG:5:SER:O	2:AG:9:LYS:HG3	2.20	0.41
3:BG:77:PRO:HB3	3:BG:208:ILE:HB	2.03	0.41
3:BG:101:ASN:HB2	3:BG:146:PHE:HE2	1.85	0.41
3:HG:58:ASN:OD1	3:HG:81:VAL:HG21	2.19	0.41
3:HG:229:GLU:OE1	3:HG:230:LEU:N	2.54	0.41
2:MG:171:ALA:HB1	3:NG:131:PHE:CE2	2.55	0.41
3:TG:123:VAL:HG21	3:TG:146:PHE:CZ	2.56	0.41
3:TG:134:ASP:OD2	3:TG:136:ARG:N	2.40	0.41
3:TG:219:LEU:HD23	3:TG:219:LEU:HA	1.88	0.41
2:G:152:ARG:NH1	2:G:153:LEU:HB2	2.34	0.41
2:J:65:GLN:HG3	2:T:46:GLN:HG2	2.02	0.41
2:J:193:MET:N	2:J:193:MET:SD	2.93	0.41
2:T:14:LEU:HD11	2:T:25:VAL:HG11	2.03	0.41
2:T:83:LYS:HG3	2:AA:23:ALA:HA	2.03	0.41
3:V:293:ILE:HA	3:V:302:LEU:O	2.21	0.41
2:AA:83:LYS:HB2	2:GA:23:ALA:O	2.20	0.41
2:MA:207:LYS:CG	2:MA:210:GLN:HG2	2.45	0.41
3:NA:234:PRO:HA	3:NA:235:PRO:HD3	1.86	0.41
2:SA:26:PHE:O	2:SA:29:LEU:HG	2.20	0.41
2:SA:191:SER:OG	2:SA:192:LYS:N	2.54	0.41
3:TA:69:LEU:O	3:TA:73:LEU:HB2	2.19	0.41
3:ZA:51:LEU:O	3:ZA:55:GLU:HG3	2.20	0.41
3:ZA:90:HIS:CE1	3:ZA:91:GLU:HG2	2.55	0.41
3:ZA:243:ASP:HA	3:ZA:246:TRP:CE3	2.55	0.41
2:EB:128:GLN:NE2	2:EB:164:PHE:CD2	2.87	0.41
3:FB:219:LEU:HD23	3:FB:219:LEU:HA	1.87	0.41
2:KB:128:GLN:NE2	2:KB:164:PHE:CD2	2.86	0.41
2:KB:261:ILE:H	2:KB:261:ILE:HG13	1.67	0.41
2:QB:5:SER:O	2:QB:9:LYS:HG3	2.19	0.41
2:QB:30:SER:O	2:QB:34:VAL:HG23	2.21	0.41
2:QB:188:LEU:HD13	3:XB:74:ARG:CD	2.42	0.41
2:QB:188:LEU:HD22	3:XB:74:ARG:NH1	2.35	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:QB:298:LEU:O	2:QB:302:GLU:HG2	2.20	0.41
2:WB:105:SER:OG	2:WB:107:ILE:HD11	2.21	0.41
3:XB:194:SER:OG	3:XB:197:ASP:N	2.54	0.41
2:CC:260:LEU:HD22	2:CC:289:ASP:HB3	2.03	0.41
2:IC:298:LEU:HA	2:IC:301:VAL:HG23	2.01	0.41
3:JC:60:ARG:HE	3:JC:60:ARG:HB2	1.62	0.41
3:JC:180:VAL:HG23	3:JC:181:ARG:N	2.34	0.41
3:PC:204:PHE:CD1	3:PC:204:PHE:N	2.87	0.41
2:MD:128:GLN:NE2	2:MD:164:PHE:HD2	2.18	0.41
2:MD:202:ILE:HD13	2:MD:202:ILE:N	2.35	0.41
3:ND:246:TRP:HB2	3:ND:247:ARG:HH12	1.86	0.41
3:TD:237:GLU:HG2	3:TD:238:ASN:N	2.35	0.41
2:EE:261:ILE:H	2:EE:261:ILE:HG13	1.67	0.41
2:QE:83:LYS:HG3	2:QE:84:ALA:N	2.36	0.41
2:QE:171:ALA:HB1	3:RE:131:PHE:CE2	2.55	0.41
3:RE:51:LEU:O	3:RE:55:GLU:HG3	2.21	0.41
4:UE:107:ALA:HB1	4:UE:124:ASP:O	2.21	0.41
2:WE:66:PHE:HA	2:CF:45:ARG:O	2.21	0.41
2:WE:89:ARG:NH2	2:WE:93:LEU:HG	2.33	0.41
2:CF:11:VAL:O	2:CF:15:MET:HG2	2.20	0.41
2:CF:298:LEU:HA	2:CF:301:VAL:HG23	2.03	0.41
2:IF:103:THR:C	2:IF:106:GLY:H	2.24	0.41
2:IF:127:PRO:HA	2:IF:130:ILE:CD1	2.51	0.41
2:IF:207:LYS:HD3	2:IF:210:GLN:HE21	1.86	0.41
2:OF:167:VAL:HG21	3:PF:131:PHE:CD1	2.55	0.41
3:BG:150:GLU:O	3:BG:153:VAL:HG12	2.21	0.41
3:HG:69:LEU:O	3:HG:73:LEU:HB2	2.20	0.41
3:TG:62:ALA:O	3:TG:79:ILE:HD12	2.21	0.41
2:G:171:ALA:HB1	3:M:131:PHE:CE2	2.55	0.41
2:G:210:GLN:O	2:G:214:VAL:HG12	2.21	0.41
2:G:259:LEU:HD11	2:G:274:PHE:HE2	1.86	0.41
2:J:201:GLU:C	2:T:139:ARG:NH2	2.73	0.41
2:T:5:SER:O	2:T:9:LYS:HG3	2.20	0.41
3:V:285:PRO:HA	4:W:119:GLY:HA2	2.02	0.41
2:AA:103:THR:C	2:AA:106:GLY:H	2.24	0.41
2:AA:191:SER:OG	2:AA:192:LYS:N	2.53	0.41
2:AA:198:THR:O	2:AA:201:GLU:HG2	2.21	0.41
2:GA:30:SER:O	2:GA:34:VAL:HG23	2.21	0.41
2:GA:235:LEU:HB3	2:GA:238:ASN:ND2	2.36	0.41
4:CB:107:ALA:HB1	4:CB:124:ASP:O	2.21	0.41
2:EB:247:ILE:HD13	2:EB:273:LYS:HE3	2.02	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:LB:150:GLU:O	3:LB:153:VAL:HG12	2.21	0.41
4:UB:105:LEU:O	3:XB:252:ARG:NE	2.53	0.41
2:WB:113:MET:HE1	2:WB:118:ALA:HA	2.03	0.41
3:XB:108:LYS:HE2	3:XB:108:LYS:HB2	1.83	0.41
3:DC:150:GLU:O	3:DC:153:VAL:HG12	2.20	0.41
2:IC:168:GLN:HB2	2:IC:169:PRO:HD2	2.01	0.41
2:OC:76:TYR:HA	2:OC:79:SER:OG	2.21	0.41
3:PC:189:THR:OG1	3:PC:190:ASN:N	2.54	0.41
2:GD:30:SER:O	2:GD:34:VAL:HG23	2.21	0.41
2:GD:110:LEU:HD13	2:GD:137:LEU:HD21	2.03	0.41
2:GD:247:ILE:HD13	2:GD:273:LYS:HE3	2.02	0.41
2:SD:37:LEU:HA	2:SD:40:ALA:HB3	2.03	0.41
2:EE:243:ASP:O	2:EE:247:ILE:HG23	2.21	0.41
3:FE:115:LEU:HD13	3:FE:117:VAL:HG23	2.03	0.41
3:FE:211:LEU:H	3:FE:211:LEU:HG	1.71	0.41
4:OE:107:ALA:HB1	4:OE:124:ASP:O	2.20	0.41
3:RE:294:ALA:O	3:RE:301:VAL:N	2.53	0.41
2:WE:172:LEU:HD12	2:WE:172:LEU:HA	1.77	0.41
3:DF:54:LEU:HD13	3:DF:231:LEU:HD21	2.01	0.41
2:IF:89:ARG:HG3	2:IF:89:ARG:NH1	2.35	0.41
2:UF:5:SER:O	2:UF:9:LYS:HG3	2.21	0.41
2:UF:113:MET:HE2	2:UF:113:MET:HB3	1.90	0.41
2:UF:191:SER:OG	2:UF:192:LYS:N	2.54	0.41
2:AG:74:ASN:O	2:AG:78:ARG:NE	2.53	0.41
2:GG:234:PHE:CE2	2:GG:312:VAL:HB	2.55	0.41
3:HG:54:LEU:HD13	3:HG:231:LEU:HD21	2.03	0.41
2:MG:73:ALA:O	2:MG:76:TYR:HB3	2.20	0.41
2:SG:113:MET:SD	2:SG:121:LEU:HD11	2.60	0.41
2:G:161:ILE:O	2:SG:195:GLY:HA2	2.21	0.41
2:B:118:ALA:HA	2:B:121:LEU:HG	2.01	0.41
3:C:58:ASN:OD1	3:C:81:VAL:HG21	2.20	0.41
3:C:224:ILE:O	3:C:228:ARG:N	2.54	0.41
3:BA:54:LEU:HD13	3:BA:231:LEU:HD21	2.03	0.41
2:MA:109:THR:O	2:MA:113:MET:HG3	2.21	0.41
2:MA:119:ALA:HB1	2:MA:149:PHE:HE1	1.86	0.41
2:MA:201:GLU:HG3	2:SA:139:ARG:CZ	2.51	0.41
3:NA:91:GLU:HA	3:NA:94:ARG:HD2	2.01	0.41
3:TA:105:ILE:HD12	3:TA:105:ILE:HA	1.77	0.41
2:YA:167:VAL:HG21	3:ZA:131:PHE:CD1	2.55	0.41
2:YA:247:ILE:HD13	2:YA:273:LYS:HE3	2.02	0.41
2:EB:100:THR:OG1	2:EB:101:ARG:NH2	2.53	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:EB:207:LYS:HE2	2:EB:207:LYS:HB2	1.78	0.41
2:EB:221:PHE:HZ	3:LB:136:ARG:HH22	1.67	0.41
3:FB:190:ASN:O	3:FB:190:ASN:OD1	2.38	0.41
2:KB:109:THR:O	2:KB:113:MET:HG3	2.21	0.41
2:KB:203:ILE:HA	2:KB:206:MET:HE3	2.01	0.41
3:LB:135:GLY:O	3:LB:211:LEU:HD21	2.20	0.41
2:QB:118:ALA:HA	2:QB:121:LEU:CD1	2.50	0.41
2:QB:247:ILE:HD13	2:QB:273:LYS:HE3	2.01	0.41
2:WB:95:GLU:HG3	2:WB:112:PHE:CE2	2.55	0.41
2:WB:247:ILE:HD13	2:WB:273:LYS:HE3	2.01	0.41
3:DC:108:LYS:HE2	3:DC:108:LYS:HB2	1.83	0.41
2:OC:69:LEU:H	2:OC:69:LEU:HD12	1.85	0.41
2:UC:76:TYR:OH	2:AD:34:VAL:HG13	2.21	0.41
2:UC:196:VAL:HG12	2:UC:232:GLU:OE2	2.20	0.41
3:VC:219:LEU:HD23	3:VC:219:LEU:HA	1.90	0.41
2:MD:83:LYS:HE2	2:MD:83:LYS:HB2	1.74	0.41
3:ZD:86:ILE:HD11	3:ZD:221:PHE:CE1	2.56	0.41
3:ZD:266:ALA:O	4:AE:66:VAL:N	2.30	0.41
2:EE:105:SER:OG	2:EE:107:ILE:HD11	2.21	0.41
3:FE:127:VAL:HG23	3:FE:153:VAL:HG11	2.02	0.41
3:FE:214:GLU:OE1	3:FE:215:PHE:N	2.54	0.41
2:QE:243:ASP:O	2:QE:247:ILE:HG23	2.21	0.41
3:RE:116:VAL:HG12	3:RE:217:ILE:CD1	2.51	0.41
2:WE:152:ARG:NH1	2:WE:153:LEU:HB2	2.35	0.41
2:WE:209:GLN:O	2:WE:212:GLU:HG3	2.21	0.41
3:XE:70:PHE:HD1	3:XE:75:ARG:O	2.04	0.41
2:CF:235:LEU:HD21	2:CF:237:GLU:HB3	2.01	0.41
4:GF:106:ILE:HA	3:JF:252:ARG:CG	2.50	0.41
2:OF:21:ARG:HH21	2:OF:25:VAL:N	2.18	0.41
3:PF:293:ILE:HA	3:PF:302:LEU:O	2.20	0.41
2:UF:78:ARG:O	2:UF:82:VAL:HG12	2.21	0.41
3:VF:294:ALA:O	3:VF:301:VAL:N	2.52	0.41
3:BG:211:LEU:H	3:BG:211:LEU:HG	1.67	0.41
3:NG:211:LEU:H	3:NG:211:LEU:HG	1.65	0.41
2:SG:5:SER:O	2:SG:9:LYS:HG3	2.20	0.41
2:SG:87:GLU:HG2	2:SG:89:ARG:H	1.84	0.41
3:TG:214:GLU:OE1	3:TG:215:PHE:N	2.54	0.41
3:M:169:TRP:CD1	3:M:175:LEU:HD13	2.54	0.41
2:B:172:LEU:HD12	2:B:172:LEU:HA	1.77	0.41
2:B:188:LEU:HB3	3:K:74:ARG:CZ	2.51	0.41
2:B:188:LEU:HD22	3:K:74:ARG:NH1	2.36	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:J:77:LEU:O	2:J:80:VAL:HG22	2.20	0.41
2:J:115:PRO:O	2:J:119:ALA:HB3	2.19	0.41
3:K:195:PRO:HD2	3:K:196:ASN:N	2.35	0.41
3:K:243:ASP:HA	3:K:246:TRP:CE3	2.56	0.41
4:L:101:ILE:N	4:L:104:TYR:O	2.35	0.41
2:T:247:ILE:HD13	2:T:273:LYS:HE3	2.03	0.41
2:AA:79:SER:O	2:AA:83:LYS:HB3	2.20	0.41
2:AA:202:ILE:CD1	2:GA:139:ARG:HH11	2.34	0.41
3:BA:86:ILE:HD11	3:BA:221:PHE:CE1	2.56	0.41
2:GA:144:ASP:O	2:GA:148:LEU:HD12	2.20	0.41
2:GA:298:LEU:O	2:GA:302:GLU:HG2	2.20	0.41
3:NA:127:VAL:CG2	3:NA:153:VAL:HG11	2.50	0.41
3:NA:190:ASN:O	3:NA:190:ASN:CG	2.58	0.41
3:NA:204:PHE:CD1	3:NA:204:PHE:N	2.89	0.41
3:NA:285:PRO:HA	4:OA:119:GLY:HA2	2.02	0.41
2:SA:65:GLN:NE2	2:YA:47:ILE:H	2.18	0.41
3:TA:54:LEU:HB2	3:TA:231:LEU:HD23	2.02	0.41
3:TA:65:PHE:CZ	3:TA:161:ALA:HB2	2.56	0.41
3:TA:90:HIS:O	3:TA:94:ARG:HG2	2.21	0.41
3:TA:162:LEU:HD12	3:TA:177:VAL:HG11	2.02	0.41
3:ZA:113:THR:OG1	3:ZA:191:ILE:HG22	2.20	0.41
3:ZA:202:THR:O	3:ZA:202:THR:OG1	2.30	0.41
3:FB:123:VAL:HG21	3:FB:146:PHE:CZ	2.56	0.41
2:KB:113:MET:HE2	2:KB:118:ALA:HA	2.03	0.41
4:SB:101:ILE:N	4:SB:104:TYR:O	2.37	0.41
2:WB:298:LEU:HA	2:WB:301:VAL:CG2	2.50	0.41
3:XB:123:VAL:HG21	3:XB:146:PHE:CZ	2.56	0.41
2:CC:205:LEU:CD2	2:IC:140:SER:HA	2.51	0.41
2:CC:221:PHE:HZ	3:JC:136:ARG:HH22	1.69	0.41
3:DC:97:PRO:HD2	3:DC:100:THR:OG1	2.21	0.41
3:DC:159:LYS:HD2	3:DC:159:LYS:O	2.21	0.41
2:IC:171:ALA:HB1	3:JC:131:PHE:CE2	2.56	0.41
3:JC:154:ILE:HD13	3:JC:154:ILE:HA	1.77	0.41
3:PC:105:ILE:HG22	3:PC:114:GLY:O	2.21	0.41
3:PC:202:THR:OG1	3:PC:217:ILE:HB	2.21	0.41
2:UC:261:ILE:O	2:UC:264:LYS:HG3	2.21	0.41
2:AD:210:GLN:O	2:AD:214:VAL:HG12	2.21	0.41
3:BD:180:VAL:HG23	3:BD:181:ARG:N	2.34	0.41
4:CD:65:THR:O	4:CD:99:ILE:HA	2.20	0.41
2:GD:20:ASP:O	2:GD:23:ALA:HB3	2.21	0.41
2:MD:119:ALA:HB1	2:MD:149:PHE:CE1	2.56	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:ND:61:PHE:HE2	3:ND:204:PHE:HE2	1.69	0.41
2:SD:234:PHE:CE2	2:SD:312:VAL:HB	2.55	0.41
2:SD:243:ASP:O	2:SD:247:ILE:HG23	2.21	0.41
3:TD:202:THR:OG1	3:TD:217:ILE:HB	2.21	0.41
2:YD:5:SER:O	2:YD:9:LYS:HG3	2.21	0.41
2:YD:99:GLU:O	2:YD:103:THR:HG23	2.21	0.41
2:YD:186:GLN:HE21	3:FE:74:ARG:NH2	2.19	0.41
2:EE:193:MET:HA	2:KE:165:GLY:HA2	2.02	0.41
3:FE:154:ILE:HA	3:FE:157:MET:HG3	2.02	0.41
2:KE:76:TYR:HB2	2:QE:41:MET:HE3	2.03	0.41
2:KE:190:ARG:O	2:QE:166:GLY:HA3	2.21	0.41
3:LE:110:LEU:HG	3:LE:223:MET:HB3	2.03	0.41
2:QE:17:ILE:HD12	2:QE:21:ARG:HG3	2.03	0.41
2:QE:89:ARG:HH11	2:QE:89:ARG:HG3	1.84	0.41
2:QE:203:ILE:HA	2:QE:206:MET:CE	2.51	0.41
2:QE:235:LEU:HD21	2:QE:237:GLU:HB3	2.02	0.41
2:WE:110:LEU:O	2:WE:113:MET:HB2	2.21	0.41
2:WE:186:GLN:NE2	3:DF:71:ASN:CG	2.74	0.41
2:WE:190:ARG:O	2:CF:166:GLY:HA3	2.20	0.41
2:WE:202:ILE:HD13	2:CF:139:ARG:HD3	2.02	0.41
2:WE:261:ILE:O	2:WE:264:LYS:HG3	2.21	0.41
2:WE:311:ILE:HA	2:WE:314:ARG:HG2	2.03	0.41
3:XE:154:ILE:HD13	3:XE:154:ILE:HA	1.79	0.41
3:XE:234:PRO:HA	3:XE:235:PRO:HD3	1.95	0.41
2:CF:182:LEU:HD23	2:CF:182:LEU:HA	1.91	0.41
2:IF:72:ASN:OD1	2:OF:41:MET:HB2	2.20	0.41
2:IF:201:GLU:HB3	2:OF:139:ARG:HE	1.85	0.41
4:MF:105:LEU:O	3:PF:252:ARG:HD3	2.21	0.41
2:OF:77:LEU:O	2:OF:80:VAL:HG22	2.21	0.41
2:OF:195:GLY:HA2	2:UF:161:ILE:O	2.21	0.41
3:PF:135:GLY:O	3:PF:211:LEU:HD21	2.21	0.41
3:PF:224:ILE:O	3:PF:228:ARG:N	2.53	0.41
3:PF:237:GLU:HG2	3:PF:238:ASN:H	1.85	0.41
3:VF:189:THR:OG1	3:VF:190:ASN:N	2.54	0.41
3:VF:191:ILE:HG13	3:VF:192:THR:HG23	2.02	0.41
2:AG:207:LYS:CG	2:AG:210:GLN:HG2	2.41	0.41
3:BG:111:ARG:O	3:BG:223:MET:HG2	2.21	0.41
3:BG:189:THR:OG1	3:BG:190:ASN:N	2.54	0.41
3:BG:293:ILE:HA	3:BG:302:LEU:O	2.21	0.41
2:GG:102:ASP:HB2	2:GG:108:GLU:HG3	2.02	0.41
2:GG:298:LEU:HA	2:GG:301:VAL:CG2	2.50	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:HG:61:PHE:CE2	3:HG:204:PHE:HE2	2.39	0.41
3:HG:128:ASP:OD1	3:HG:133:GLY:HA3	2.21	0.41
3:HG:285:PRO:HA	4:IG:119:GLY:HA2	2.03	0.41
3:NG:198:ILE:HD12	3:NG:198:ILE:HA	1.90	0.41
2:SG:161:ILE:O	2:SG:164:PHE:HB3	2.21	0.41
3:TG:105:ILE:HG22	3:TG:114:GLY:O	2.21	0.41
3:TG:189:THR:OG1	3:TG:190:ASN:N	2.54	0.41
2:G:139:ARG:HH21	2:SG:202:ILE:HA	1.84	0.41
4:N:65:THR:O	4:N:99:ILE:HA	2.20	0.41
2:B:26:PHE:HE2	2:B:37:LEU:HD12	1.85	0.41
3:C:150:GLU:O	3:C:153:VAL:HG12	2.20	0.41
2:J:75:GLU:O	2:J:79:SER:HB3	2.20	0.41
3:K:135:GLY:O	3:K:211:LEU:HD21	2.21	0.41
3:K:191:ILE:HG13	3:K:192:THR:HG23	2.02	0.41
3:K:228:ARG:O	3:K:232:VAL:HG23	2.21	0.41
2:T:171:ALA:HB1	3:V:131:PHE:CZ	2.56	0.41
2:MA:40:ALA:O	2:MA:44:VAL:HG23	2.21	0.41
2:MA:103:THR:C	2:MA:106:GLY:H	2.24	0.41
2:MA:201:GLU:HG3	2:SA:139:ARG:HH22	1.84	0.41
3:NA:111:ARG:O	3:NA:223:MET:HG2	2.21	0.41
2:SA:83:LYS:HB2	2:YA:26:PHE:HB3	2.03	0.41
2:SA:188:LEU:HB3	3:ZA:74:ARG:NE	2.36	0.41
2:YA:115:PRO:O	2:YA:119:ALA:HB3	2.20	0.41
2:YA:196:VAL:HG12	2:YA:232:GLU:OE2	2.20	0.41
2:EB:65:GLN:HG3	2:KB:46:GLN:HG2	2.03	0.41
2:EB:207:LYS:NZ	2:EB:209:GLN:HB3	2.32	0.41
3:FB:154:ILE:HA	3:FB:154:ILE:HD13	1.85	0.41
3:FB:285:PRO:HA	4:GB:119:GLY:HA2	2.02	0.41
3:XB:211:LEU:H	3:XB:211:LEU:HG	1.69	0.41
2:OC:163:THR:HG21	3:PC:134:ASP:OD2	2.21	0.41
2:UC:76:TYR:HE1	2:AD:38:SER:CB	2.33	0.41
2:UC:243:ASP:O	2:UC:247:ILE:HG23	2.21	0.41
3:VC:113:THR:OG1	3:VC:191:ILE:HG22	2.21	0.41
2:AD:171:ALA:HB1	3:BD:131:PHE:CE2	2.56	0.41
2:GD:117:SER:O	2:GD:121:LEU:HG	2.20	0.41
2:GD:159:LEU:HD21	3:HD:137:PHE:CE1	2.56	0.41
3:HD:59:GLU:OE1	3:HD:59:GLU:N	2.54	0.41
3:HD:228:ARG:O	3:HD:232:VAL:HG23	2.21	0.41
3:ZD:150:GLU:O	3:ZD:153:VAL:HG12	2.20	0.41
2:EE:132:THR:O	2:EE:135:VAL:HB	2.21	0.41
2:KE:132:THR:O	2:KE:135:VAL:HB	2.21	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:KE:193:MET:HG2	2:KE:194:GLY:N	2.36	0.41
3:LE:240:ARG:HA	3:LE:240:ARG:HH11	1.85	0.41
3:RE:162:LEU:HD12	3:RE:177:VAL:HG11	2.02	0.41
2:WE:243:ASP:O	2:WE:247:ILE:HG23	2.21	0.41
3:XE:150:GLU:O	3:XE:153:VAL:HG12	2.21	0.41
2:CF:120:ASP:OD1	2:CF:123:ARG:NH1	2.54	0.41
3:PF:211:LEU:H	3:PF:211:LEU:HG	1.66	0.41
2:UF:119:ALA:HB1	2:UF:149:PHE:CE1	2.56	0.41
2:UF:130:ILE:HG13	2:UF:131:ALA:N	2.35	0.41
2:UF:171:ALA:HB1	3:VF:131:PHE:CE2	2.56	0.41
2:UF:207:LYS:NZ	2:UF:210:GLN:HB2	2.30	0.41
3:VF:66:ARG:HG3	3:VF:77:PRO:O	2.21	0.41
3:VF:198:ILE:HD12	3:VF:199:VAL:N	2.35	0.41
3:HG:66:ARG:NH1	3:HG:78:ASP:HA	2.35	0.41
3:HG:101:ASN:HB2	3:HG:146:PHE:CE2	2.55	0.41
3:HG:106:HIS:NE2	3:HG:113:THR:HG22	2.35	0.41
2:MG:83:LYS:HD2	2:SG:26:PHE:C	2.42	0.41
2:MG:193:MET:HG2	2:MG:194:GLY:N	2.36	0.41
2:MG:214:VAL:O	2:MG:218:VAL:HG23	2.21	0.41
2:MG:259:LEU:HD11	2:MG:274:PHE:HE2	1.86	0.41
2:SG:99:GLU:OE1	2:SG:99:GLU:HA	2.21	0.41
3:TG:150:GLU:O	3:TG:153:VAL:HG12	2.21	0.41
2:G:69:LEU:HD21	2:B:47:ILE:HB	2.03	0.40
2:G:110:LEU:HD13	2:G:137:LEU:HD21	2.02	0.40
3:M:195:PRO:HD2	3:M:196:ASN:N	2.36	0.40
3:C:219:LEU:HD23	3:C:219:LEU:HA	1.83	0.40
2:T:74:ASN:O	2:T:78:ARG:NE	2.54	0.40
3:V:106:HIS:HE1	3:V:113:THR:CG2	2.34	0.40
3:V:117:VAL:O	3:V:215:PHE:HB2	2.20	0.40
2:AA:5:SER:O	2:AA:9:LYS:HG3	2.21	0.40
2:GA:74:ASN:O	2:GA:78:ARG:NE	2.54	0.40
2:GA:243:ASP:O	2:GA:247:ILE:HG23	2.21	0.40
3:HA:198:ILE:HG13	3:HA:221:PHE:CD1	2.56	0.40
3:HA:198:ILE:HG13	3:HA:221:PHE:CE1	2.55	0.40
2:SA:5:SER:O	2:SA:9:LYS:HG3	2.21	0.40
2:SA:243:ASP:O	2:SA:247:ILE:HG23	2.20	0.40
4:WA:107:ALA:HB1	4:WA:124:ASP:O	2.20	0.40
2:YA:83:LYS:HB3	2:EB:27:LYS:HB2	2.02	0.40
3:ZA:96:LEU:HA	3:ZA:97:PRO:HD3	1.91	0.40
3:ZA:100:THR:HG23	3:ZA:119:SER:HA	2.03	0.40
3:ZA:293:ILE:HA	3:ZA:302:LEU:O	2.21	0.40

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:FB:150:GLU:O	3:FB:153:VAL:HG12	2.20	0.40
4:IB:105:LEU:O	3:LB:252:ARG:HD3	2.20	0.40
2:WB:65:GLN:NE2	2:CC:47:ILE:H	2.18	0.40
3:DC:69:LEU:O	3:DC:73:LEU:HB2	2.21	0.40
2:IC:234:PHE:CE2	2:IC:312:VAL:HB	2.56	0.40
3:JC:189:THR:OG1	3:JC:190:ASN:N	2.54	0.40
3:VC:96:LEU:HA	3:VC:97:PRO:HD3	1.91	0.40
2:GD:243:ASP:O	2:GD:247:ILE:HG23	2.21	0.40
2:MD:149:PHE:CG	2:MD:153:LEU:HD23	2.56	0.40
2:YD:198:THR:O	2:YD:202:ILE:HG12	2.21	0.40
3:FE:108:LYS:HE2	3:FE:108:LYS:HB2	1.83	0.40
2:KE:11:VAL:HG11	2:KE:41:MET:HB3	2.02	0.40
2:KE:152:ARG:NH1	2:KE:153:LEU:HB2	2.36	0.40
2:KE:243:ASP:O	2:KE:247:ILE:HG23	2.21	0.40
2:KE:298:LEU:HA	2:KE:301:VAL:CG2	2.50	0.40
3:XE:101:ASN:HB2	3:XE:146:PHE:CE2	2.56	0.40
2:CF:76:TYR:HE1	2:IF:38:SER:CB	2.33	0.40
2:CF:171:ALA:HB1	3:DF:131:PHE:CE2	2.56	0.40
3:DF:115:LEU:HD13	3:DF:117:VAL:HG23	2.03	0.40
2:IF:186:GLN:CA	3:PF:74:ARG:HH22	2.34	0.40
2:IF:243:ASP:O	2:IF:247:ILE:HG23	2.21	0.40
2:OF:128:GLN:NE2	2:OF:164:PHE:HD2	2.15	0.40
2:OF:223:GLY:O	2:OF:226:ALA:HB3	2.21	0.40
2:UF:209:GLN:O	2:UF:212:GLU:HG3	2.21	0.40
3:VF:207:GLU:HG3	3:VF:212:THR:HG22	2.02	0.40
2:AG:110:LEU:HD13	2:AG:137:LEU:HD21	2.02	0.40
2:AG:214:VAL:O	2:AG:218:VAL:HG23	2.21	0.40
2:AG:219:ARG:NH1	2:AG:223:GLY:O	2.54	0.40
3:BG:246:TRP:HB2	3:BG:247:ARG:HH12	1.86	0.40
2:GG:207:LYS:NZ	2:GG:209:GLN:HB3	2.29	0.40
3:HG:66:ARG:HG3	3:HG:77:PRO:O	2.21	0.40
3:HG:101:ASN:HD22	3:HG:146:PHE:HE2	1.69	0.40
2:MG:285:ILE:HG13	2:MG:286:LEU:HD22	2.03	0.40
3:NG:77:PRO:HB3	3:NG:208:ILE:HB	2.03	0.40
2:G:139:ARG:HH21	2:SG:202:ILE:CD1	2.34	0.40
2:J:243:ASP:O	2:J:247:ILE:HG23	2.21	0.40
3:K:150:GLU:O	3:K:153:VAL:HG12	2.22	0.40
3:K:293:ILE:HA	3:K:302:LEU:O	2.22	0.40
2:T:128:GLN:NE2	2:T:164:PHE:CD2	2.88	0.40
3:V:101:ASN:HB2	3:V:146:PHE:HE2	1.85	0.40
3:V:219:LEU:HD23	3:V:219:LEU:HA	1.91	0.40

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:V:239:SER:O	3:V:242:GLU:HG2	2.21	0.40
2:AA:75:GLU:OE1	2:AA:76:TYR:HD2	2.03	0.40
2:AA:132:THR:O	2:AA:135:VAL:HB	2.21	0.40
3:BA:54:LEU:HB2	3:BA:231:LEU:HD23	2.02	0.40
2:GA:137:LEU:HB3	2:GA:141:GLN:OE1	2.20	0.40
3:HA:189:THR:OG1	3:HA:190:ASN:N	2.54	0.40
3:HA:267:ASP:HA	4:IA:65:THR:HA	2.03	0.40
2:MA:20:ASP:OD1	2:MA:21:ARG:N	2.54	0.40
2:MA:152:ARG:NH1	2:MA:153:LEU:HB2	2.35	0.40
3:NA:65:PHE:CZ	3:NA:161:ALA:HB2	2.56	0.40
3:TA:127:VAL:HG23	3:TA:153:VAL:HG11	2.03	0.40
3:TA:183:GLU:OE1	3:TA:184:MET:N	2.54	0.40
3:LB:96:LEU:HA	3:LB:97:PRO:HD3	1.96	0.40
2:WB:259:LEU:HD11	2:WB:274:PHE:HE2	1.87	0.40
3:XB:54:LEU:HD23	3:XB:84:ILE:HD12	2.04	0.40
2:CC:218:VAL:HG12	2:CC:226:ALA:HB2	2.03	0.40
3:DC:109:PRO:O	3:DC:111:ARG:NH1	2.54	0.40
2:IC:113:MET:HE2	2:IC:118:ALA:HA	2.02	0.40
2:OC:108:GLU:OE2	2:OC:112:PHE:HE2	2.04	0.40
3:BD:86:ILE:HD11	3:BD:221:PHE:HE1	1.85	0.40
3:HD:204:PHE:N	3:HD:204:PHE:CD1	2.87	0.40
2:MD:260:LEU:HD22	2:MD:289:ASP:HB3	2.03	0.40
2:YD:128:GLN:NE2	2:YD:164:PHE:HD2	2.15	0.40
3:FE:51:LEU:O	3:FE:55:GLU:HG3	2.21	0.40
2:KE:247:ILE:HD13	2:KE:273:LYS:HE3	2.02	0.40
2:QE:103:THR:C	2:QE:106:GLY:H	2.25	0.40
3:XE:105:ILE:HD12	3:XE:105:ILE:HA	1.86	0.40
2:CF:103:THR:C	2:CF:106:GLY:H	2.24	0.40
3:DF:189:THR:OG1	3:DF:190:ASN:N	2.54	0.40
3:JF:198:ILE:HD12	3:JF:198:ILE:HA	1.82	0.40
2:OF:141:GLN:HA	2:OF:144:ASP:OD2	2.20	0.40
2:OF:206:MET:SD	2:OF:210:GLN:HB2	2.61	0.40
3:PF:66:ARG:HG3	3:PF:77:PRO:O	2.21	0.40
2:UF:132:THR:O	2:UF:135:VAL:HB	2.20	0.40
2:AG:83:LYS:HA	2:GG:23:ALA:HB1	2.02	0.40
2:GG:207:LYS:HE2	2:GG:207:LYS:HB2	1.86	0.40
2:SG:243:ASP:O	2:SG:247:ILE:HG23	2.20	0.40
3:TG:55:GLU:HA	3:TG:58:ASN:ND2	2.37	0.40
2:G:76:TYR:CE2	2:G:80:VAL:HG13	2.56	0.40
3:M:219:LEU:HD23	3:M:219:LEU:HA	1.89	0.40
2:B:120:ASP:OD1	2:B:123:ARG:NH1	2.54	0.40

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:J:204:ASN:OD1	2:J:233:MET:HB3	2.20	0.40
3:BA:234:PRO:HA	3:BA:235:PRO:HD3	1.89	0.40
3:BA:246:TRP:HB2	3:BA:247:ARG:NH1	2.36	0.40
2:GA:206:MET:HB2	2:GA:211:GLU:HG2	2.04	0.40
3:HA:51:LEU:O	3:HA:55:GLU:HG3	2.21	0.40
3:HA:183:GLU:OE1	3:HA:183:GLU:C	2.60	0.40
2:MA:110:LEU:O	2:MA:113:MET:HB2	2.21	0.40
2:MA:259:LEU:HD11	2:MA:274:PHE:HE2	1.87	0.40
3:NA:108:LYS:HE2	3:NA:108:LYS:HB2	1.82	0.40
2:YA:72:ASN:HB2	2:EB:15:MET:SD	2.62	0.40
2:EB:113:MET:SD	2:EB:121:LEU:HD11	2.62	0.40
3:FB:111:ARG:O	3:FB:223:MET:HG2	2.21	0.40
2:KB:77:LEU:O	2:KB:80:VAL:HG22	2.21	0.40
2:KB:119:ALA:HB1	2:KB:149:PHE:CE1	2.57	0.40
3:LB:190:ASN:O	3:RB:236:LEU:HD12	2.21	0.40
3:RB:128:ASP:OD1	3:RB:133:GLY:HA3	2.21	0.40
2:WB:21:ARG:NH2	2:WB:24:GLU:HB2	2.36	0.40
2:WB:21:ARG:HH21	2:WB:25:VAL:N	2.19	0.40
2:WB:120:ASP:HB3	3:DC:76:SER:OG	2.21	0.40
3:XB:185:GLN:HG2	3:XB:187:LYS:NZ	2.36	0.40
2:CC:171:ALA:HB1	3:DC:131:PHE:CE2	2.57	0.40
2:CC:201:GLU:HG2	2:IC:139:ARG:CZ	2.51	0.40
2:OC:130:ILE:HD11	2:OC:157:VAL:HG22	2.04	0.40
3:PC:107:LEU:HD11	3:PC:219:LEU:HD21	2.03	0.40
4:QC:101:ILE:N	4:QC:104:TYR:O	2.36	0.40
2:UC:40:ALA:O	2:UC:44:VAL:HG23	2.22	0.40
3:BD:105:ILE:HD12	3:BD:105:ILE:HA	1.90	0.40
2:GD:62:GLU:O	2:GD:65:GLN:HB3	2.22	0.40
2:GD:191:SER:OG	2:GD:192:LYS:N	2.53	0.40
3:HD:58:ASN:OD1	3:HD:81:VAL:HG21	2.20	0.40
2:SD:33:GLU:H	2:SD:33:GLU:HG3	1.58	0.40
2:SD:311:ILE:HA	2:SD:314:ARG:HG2	2.04	0.40
3:TD:285:PRO:HA	4:UD:119:GLY:HA2	2.04	0.40
2:YD:71:ILE:O	2:YD:74:ASN:ND2	2.53	0.40
4:NE:75:ILE:O	4:NE:77:GLU:N	2.54	0.40
2:QE:53:THR:O	2:QE:57:SER:OG	2.33	0.40
2:QE:193:MET:HG2	2:QE:194:GLY:N	2.36	0.40
3:XE:118:PHE:HE2	3:XE:158:LEU:HD12	1.84	0.40
2:CF:206:MET:HG2	2:IF:143:ALA:CB	2.49	0.40
3:DF:227:LEU:HD12	3:DF:230:LEU:HD22	2.03	0.40
2:OF:83:LYS:HB3	2:UF:23:ALA:HA	2.03	0.40

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:OF:186:GLN:HE22	3:VF:71:ASN:CA	2.18	0.40
2:OF:243:ASP:O	2:OF:247:ILE:HG23	2.21	0.40
2:UF:88:GLU:N	2:UF:88:GLU:OE1	2.55	0.40
2:UF:261:ILE:H	2:UF:261:ILE:HG13	1.67	0.40
3:HG:162:LEU:HA	3:HG:162:LEU:HD13	1.87	0.40
3:HG:189:THR:OG1	3:HG:190:ASN:N	2.55	0.40
2:MG:78:ARG:O	2:MG:82:VAL:HG23	2.21	0.40
2:MG:130:ILE:HG13	2:MG:131:ALA:N	2.36	0.40
2:MG:207:LYS:NZ	2:MG:210:GLN:HG3	2.36	0.40
3:TG:183:GLU:CD	3:TG:185:GLN:H	2.23	0.40
2:B:183:LEU:O	2:B:186:GLN:N	2.54	0.40
2:B:219:ARG:NH1	2:B:223:GLY:O	2.55	0.40
3:C:294:ALA:O	3:C:301:VAL:N	2.52	0.40
2:T:62:GLU:O	2:T:65:GLN:HB3	2.22	0.40
3:V:243:ASP:HA	3:V:246:TRP:CE3	2.57	0.40
2:AA:45:ARG:CG	2:AA:46:GLN:HG3	2.48	0.40
3:BA:55:GLU:HA	3:BA:58:ASN:ND2	2.36	0.40
3:BA:202:THR:OG1	3:BA:217:ILE:HB	2.20	0.40
2:MA:188:LEU:HB3	3:TA:74:ARG:CZ	2.51	0.40
2:MA:243:ASP:O	2:MA:247:ILE:HG23	2.21	0.40
3:TA:123:VAL:HG21	3:TA:146:PHE:CZ	2.57	0.40
3:TA:162:LEU:HA	3:TA:162:LEU:HD13	1.79	0.40
2:YA:5:SER:O	2:YA:9:LYS:HG3	2.20	0.40
2:YA:261:ILE:O	2:YA:264:LYS:HG3	2.20	0.40
3:ZA:285:PRO:HA	4:AB:119:GLY:HA2	2.03	0.40
2:EB:234:PHE:CD2	2:EB:313:ARG:HB3	2.55	0.40
2:EB:235:LEU:HB3	2:EB:238:ASN:ND2	2.35	0.40
3:LB:180:VAL:HG23	3:LB:181:ARG:N	2.37	0.40
3:LB:185:GLN:HG3	3:LB:186:VAL:H	1.85	0.40
2:WB:191:SER:OG	2:WB:192:LYS:N	2.54	0.40
3:XB:86:ILE:HD12	3:XB:86:ILE:HA	1.94	0.40
3:XB:219:LEU:HD22	3:XB:223:MET:HE1	2.03	0.40
3:XB:266:ALA:O	4:YB:66:VAL:N	2.29	0.40
2:CC:5:SER:O	2:CC:9:LYS:HG3	2.22	0.40
2:CC:171:ALA:HB1	3:DC:131:PHE:CZ	2.56	0.40
3:DC:128:ASP:OD1	3:DC:133:GLY:HA3	2.21	0.40
2:OC:243:ASP:O	2:OC:247:ILE:HG23	2.21	0.40
3:PC:240:ARG:HA	3:PC:240:ARG:HH11	1.85	0.40
3:PC:267:ASP:HA	4:QC:65:THR:HA	2.03	0.40
4:WC:101:ILE:N	4:WC:104:TYR:O	2.36	0.40
2:AD:108:GLU:HG3	2:AD:112:PHE:HE2	1.85	0.40

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:AD:206:MET:SD	2:AD:210:GLN:HB2	2.61	0.40
2:AD:243:ASP:O	2:AD:247:ILE:HG23	2.21	0.40
2:GD:168:GLN:HB2	2:GD:169:PRO:HD2	2.03	0.40
3:HD:234:PRO:HA	3:HD:235:PRO:HD3	1.90	0.40
2:SD:202:ILE:HD11	2:YD:139:ARG:NH1	2.36	0.40
2:YD:34:VAL:O	2:YD:37:LEU:HD12	2.22	0.40
2:YD:65:GLN:NE2	2:EE:47:ILE:H	2.20	0.40
2:YD:108:GLU:HG3	2:YD:112:PHE:HE2	1.86	0.40
2:YD:243:ASP:O	2:YD:247:ILE:HG23	2.22	0.40
3:ZD:105:ILE:HD12	3:ZD:105:ILE:HA	1.89	0.40
3:ZD:202:THR:OG1	3:ZD:217:ILE:HB	2.22	0.40
3:FE:122:LEU:CD1	3:FE:215:PHE:HB3	2.44	0.40
3:FE:150:GLU:O	3:FE:153:VAL:HG12	2.21	0.40
3:LE:204:PHE:N	3:LE:204:PHE:CD1	2.88	0.40
3:LE:236:LEU:HB3	3:LE:238:ASN:OD1	2.21	0.40
2:QE:126:HIS:CD2	2:QE:127:PRO:HD2	2.57	0.40
2:QE:128:GLN:NE2	2:QE:164:PHE:HD2	2.19	0.40
2:CF:243:ASP:O	2:CF:247:ILE:HG23	2.21	0.40
2:IF:149:PHE:CD2	2:IF:153:LEU:HD22	2.55	0.40
2:OF:207:LYS:NZ	2:OF:209:GLN:HB3	2.36	0.40
2:OF:219:ARG:HH22	2:OF:230:ILE:HD12	1.85	0.40
4:QF:101:ILE:N	4:QF:104:TYR:O	2.37	0.40
2:AG:243:ASP:O	2:AG:247:ILE:HG23	2.21	0.40
2:GG:128:GLN:NE2	2:GG:164:PHE:CD2	2.87	0.40
2:GG:188:LEU:HD22	3:NG:74:ARG:HE	1.85	0.40
2:GG:234:PHE:CD2	2:GG:313:ARG:HB3	2.57	0.40
2:GG:311:ILE:HA	2:GG:314:ARG:HG2	2.04	0.40
2:G:8:ASP:HA	2:G:11:VAL:HG22	2.04	0.40
2:B:186:GLN:C	3:K:74:ARG:HH22	2.25	0.40
2:J:126:HIS:ND1	3:K:127:VAL:HG13	2.37	0.40
2:T:129:ILE:HG23	2:T:175:LEU:CD2	2.51	0.40
2:T:130:ILE:HD11	2:T:157:VAL:HG22	2.04	0.40
3:V:240:ARG:HA	3:V:240:ARG:NE	2.36	0.40
2:AA:85:LEU:HD22	2:AA:85:LEU:H	1.85	0.40
2:AA:115:PRO:O	2:AA:119:ALA:HB3	2.21	0.40
3:HA:55:GLU:HA	3:HA:58:ASN:HD22	1.86	0.40
2:MA:76:TYR:OH	2:SA:34:VAL:HG13	2.21	0.40
2:MA:244:ASP:OD1	2:MA:273:LYS:NZ	2.50	0.40
3:NA:194:SER:OG	3:NA:197:ASP:N	2.55	0.40
2:SA:76:TYR:CE2	2:YA:26:PHE:CZ	3.04	0.40
4:AB:53:ASP:O	4:AB:56:LEU:N	2.54	0.40

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:FB:108:LYS:HE2	3:FB:108:LYS:HB2	1.81	0.40
3:RB:243:ASP:HA	3:RB:246:TRP:CE3	2.57	0.40
2:WB:11:VAL:O	2:WB:14:LEU:HB3	2.22	0.40
3:XB:62:ALA:O	3:XB:79:ILE:HD12	2.21	0.40
2:CC:219:ARG:NH2	2:CC:230:ILE:HD12	2.37	0.40
2:IC:311:ILE:HA	2:IC:314:ARG:HG2	2.04	0.40
3:JC:108:LYS:HE2	3:JC:108:LYS:HB2	1.83	0.40
3:JC:204:PHE:CD1	3:JC:204:PHE:N	2.88	0.40
2:OC:72:ASN:HD22	2:UC:41:MET:HB2	1.85	0.40
2:OC:131:ALA:HA	2:OC:134:LEU:HD12	2.04	0.40
2:UC:128:GLN:NE2	2:UC:164:PHE:CD2	2.89	0.40
2:UC:168:GLN:HB2	2:UC:169:PRO:HD2	2.03	0.40
2:AD:205:LEU:CD2	2:GD:140:SER:HA	2.52	0.40
2:AD:311:ILE:HA	2:AD:314:ARG:HG2	2.04	0.40
3:BD:107:LEU:HD11	3:BD:219:LEU:HD21	2.02	0.40
3:BD:128:ASP:OD1	3:BD:133:GLY:HA3	2.21	0.40
2:GD:261:ILE:O	2:GD:264:LYS:HG3	2.20	0.40
2:MD:31:THR:HA	2:MD:34:VAL:CG2	2.51	0.40
2:SD:103:THR:O	2:SD:183:LEU:HD22	2.21	0.40
2:EE:113:MET:HE2	2:EE:113:MET:HB3	1.96	0.40
2:EE:159:LEU:HD12	2:EE:159:LEU:HA	1.81	0.40
3:LE:123:VAL:HG21	3:LE:146:PHE:CZ	2.56	0.40
3:LE:293:ILE:HA	3:LE:302:LEU:O	2.22	0.40
2:QE:100:THR:OG1	2:QE:101:ARG:NH2	2.55	0.40
2:WE:11:VAL:O	2:WE:14:LEU:HB3	2.20	0.40
3:DF:219:LEU:HD23	3:DF:219:LEU:HA	1.80	0.40
3:PF:107:LEU:HD11	3:PF:219:LEU:HD21	2.04	0.40
2:UF:115:PRO:O	2:UF:119:ALA:HB3	2.21	0.40
3:VF:94:ARG:HG2	3:VF:94:ARG:H	1.73	0.40
2:AG:191:SER:OG	2:AG:192:LYS:N	2.55	0.40
3:BG:190:ASN:O	3:BG:190:ASN:OD1	2.40	0.40
2:GG:102:ASP:O	2:GG:107:ILE:HG12	2.20	0.40
2:GG:103:THR:C	2:GG:106:GLY:H	2.25	0.40
2:GG:168:GLN:HB2	2:GG:169:PRO:HD2	2.03	0.40

There are no symmetry-related clashes.

5.3 Torsion angles

5.3.1 Protein backbone

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

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

Mol	Chain	Analysed	Favoured	Allowed	Outliers	Percentiles	
1	A	45/47 (96%)	44 (98%)	1 (2%)	0	100	100
1	BC	45/47 (96%)	44 (98%)	1 (2%)	0	100	100
1	BF	45/47 (96%)	44 (98%)	1 (2%)	0	100	100
1	DB	45/47 (96%)	44 (98%)	1 (2%)	0	100	100
1	DE	45/47 (96%)	44 (98%)	1 (2%)	0	100	100
1	F	45/47 (96%)	44 (98%)	1 (2%)	0	100	100
1	FA	45/47 (96%)	44 (98%)	1 (2%)	0	100	100
1	FD	45/47 (96%)	44 (98%)	1 (2%)	0	100	100
1	FG	45/47 (96%)	44 (98%)	1 (2%)	0	100	100
1	HC	45/47 (96%)	44 (98%)	1 (2%)	0	100	100
1	HF	45/47 (96%)	44 (98%)	1 (2%)	0	100	100
1	I	45/47 (96%)	44 (98%)	1 (2%)	0	100	100
1	JB	45/47 (96%)	44 (98%)	1 (2%)	0	100	100
1	JE	45/47 (96%)	44 (98%)	1 (2%)	0	100	100
1	LA	45/47 (96%)	44 (98%)	1 (2%)	0	100	100
1	LD	45/47 (96%)	44 (98%)	1 (2%)	0	100	100
1	LG	45/47 (96%)	44 (98%)	1 (2%)	0	100	100
1	NC	45/47 (96%)	44 (98%)	1 (2%)	0	100	100
1	NF	45/47 (96%)	44 (98%)	1 (2%)	0	100	100
1	PB	45/47 (96%)	44 (98%)	1 (2%)	0	100	100
1	PE	45/47 (96%)	44 (98%)	1 (2%)	0	100	100
1	RA	45/47 (96%)	44 (98%)	1 (2%)	0	100	100
1	RD	45/47 (96%)	44 (98%)	1 (2%)	0	100	100
1	RG	45/47 (96%)	44 (98%)	1 (2%)	0	100	100
1	S	45/47 (96%)	44 (98%)	1 (2%)	0	100	100

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Mol	Chain	Analysed	Favoured	Allowed	Outliers	Percentiles	
1	TC	45/47 (96%)	44 (98%)	1 (2%)	0	100	100
1	TF	45/47 (96%)	44 (98%)	1 (2%)	0	100	100
1	VB	45/47 (96%)	44 (98%)	1 (2%)	0	100	100
1	VE	45/47 (96%)	44 (98%)	1 (2%)	0	100	100
1	XA	45/47 (96%)	44 (98%)	1 (2%)	0	100	100
1	XD	45/47 (96%)	44 (98%)	1 (2%)	0	100	100
1	Z	45/47 (96%)	44 (98%)	1 (2%)	0	100	100
1	ZC	45/47 (96%)	44 (98%)	1 (2%)	0	100	100
1	ZF	45/47 (96%)	44 (98%)	1 (2%)	0	100	100
2	AA	318/331 (96%)	309 (97%)	9 (3%)	0	100	100
2	AD	318/331 (96%)	307 (96%)	11 (4%)	0	100	100
2	AG	318/331 (96%)	309 (97%)	9 (3%)	0	100	100
2	B	318/331 (96%)	311 (98%)	7 (2%)	0	100	100
2	CC	318/331 (96%)	310 (98%)	8 (2%)	0	100	100
2	CF	318/331 (96%)	308 (97%)	10 (3%)	0	100	100
2	EB	318/331 (96%)	310 (98%)	8 (2%)	0	100	100
2	EE	318/331 (96%)	309 (97%)	9 (3%)	0	100	100
2	G	318/331 (96%)	310 (98%)	8 (2%)	0	100	100
2	GA	318/331 (96%)	309 (97%)	9 (3%)	0	100	100
2	GD	318/331 (96%)	310 (98%)	8 (2%)	0	100	100
2	GG	318/331 (96%)	309 (97%)	9 (3%)	0	100	100
2	IC	318/331 (96%)	310 (98%)	8 (2%)	0	100	100
2	IF	318/331 (96%)	307 (96%)	11 (4%)	0	100	100
2	J	318/331 (96%)	308 (97%)	10 (3%)	0	100	100
2	KB	318/331 (96%)	308 (97%)	10 (3%)	0	100	100
2	KE	318/331 (96%)	306 (96%)	12 (4%)	0	100	100
2	MA	318/331 (96%)	310 (98%)	8 (2%)	0	100	100
2	MD	318/331 (96%)	305 (96%)	13 (4%)	0	100	100
2	MG	318/331 (96%)	312 (98%)	6 (2%)	0	100	100
2	OC	318/331 (96%)	311 (98%)	7 (2%)	0	100	100
2	OF	318/331 (96%)	309 (97%)	9 (3%)	0	100	100

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Mol	Chain	Analysed	Favoured	Allowed	Outliers	Percentiles	
2	QB	318/331 (96%)	308 (97%)	10 (3%)	0	100	100
2	QE	318/331 (96%)	310 (98%)	8 (2%)	0	100	100
2	SA	318/331 (96%)	308 (97%)	10 (3%)	0	100	100
2	SD	318/331 (96%)	309 (97%)	9 (3%)	0	100	100
2	SG	318/331 (96%)	309 (97%)	9 (3%)	0	100	100
2	T	318/331 (96%)	309 (97%)	9 (3%)	0	100	100
2	UC	318/331 (96%)	308 (97%)	10 (3%)	0	100	100
2	UF	318/331 (96%)	309 (97%)	9 (3%)	0	100	100
2	WB	318/331 (96%)	309 (97%)	9 (3%)	0	100	100
2	WE	318/331 (96%)	308 (97%)	10 (3%)	0	100	100
2	YA	318/331 (96%)	309 (97%)	9 (3%)	0	100	100
2	YD	318/331 (96%)	308 (97%)	10 (3%)	0	100	100
3	BA	279/300 (93%)	260 (93%)	19 (7%)	0	100	100
3	BD	279/300 (93%)	261 (94%)	18 (6%)	0	100	100
3	BG	279/300 (93%)	262 (94%)	17 (6%)	0	100	100
3	C	279/300 (93%)	260 (93%)	19 (7%)	0	100	100
3	DC	279/300 (93%)	263 (94%)	16 (6%)	0	100	100
3	DF	279/300 (93%)	262 (94%)	17 (6%)	0	100	100
3	FB	279/300 (93%)	261 (94%)	18 (6%)	0	100	100
3	FE	279/300 (93%)	258 (92%)	21 (8%)	0	100	100
3	HA	279/300 (93%)	260 (93%)	19 (7%)	0	100	100
3	HD	279/300 (93%)	260 (93%)	19 (7%)	0	100	100
3	HG	279/300 (93%)	257 (92%)	22 (8%)	0	100	100
3	JC	279/300 (93%)	262 (94%)	17 (6%)	0	100	100
3	JF	279/300 (93%)	259 (93%)	20 (7%)	0	100	100
3	K	279/300 (93%)	262 (94%)	17 (6%)	0	100	100
3	LB	279/300 (93%)	261 (94%)	18 (6%)	0	100	100
3	LE	279/300 (93%)	261 (94%)	18 (6%)	0	100	100
3	M	279/300 (93%)	262 (94%)	17 (6%)	0	100	100
3	NA	279/300 (93%)	261 (94%)	18 (6%)	0	100	100
3	ND	279/300 (93%)	260 (93%)	19 (7%)	0	100	100

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Mol	Chain	Analysed	Favoured	Allowed	Outliers	Percentiles	
3	NG	279/300 (93%)	261 (94%)	18 (6%)	0	100	100
3	PC	279/300 (93%)	259 (93%)	20 (7%)	0	100	100
3	PF	279/300 (93%)	264 (95%)	15 (5%)	0	100	100
3	RB	279/300 (93%)	261 (94%)	18 (6%)	0	100	100
3	RE	279/300 (93%)	262 (94%)	17 (6%)	0	100	100
3	TA	279/300 (93%)	263 (94%)	16 (6%)	0	100	100
3	TD	279/300 (93%)	258 (92%)	21 (8%)	0	100	100
3	TG	279/300 (93%)	261 (94%)	18 (6%)	0	100	100
3	V	279/300 (93%)	260 (93%)	19 (7%)	0	100	100
3	VC	279/300 (93%)	260 (93%)	19 (7%)	0	100	100
3	VF	279/300 (93%)	259 (93%)	20 (7%)	0	100	100
3	XB	279/300 (93%)	257 (92%)	22 (8%)	0	100	100
3	XE	279/300 (93%)	260 (93%)	19 (7%)	0	100	100
3	ZA	279/300 (93%)	259 (93%)	20 (7%)	0	100	100
3	ZD	279/300 (93%)	259 (93%)	20 (7%)	0	100	100
4	AB	91/137 (66%)	88 (97%)	3 (3%)	0	100	100
4	AC	73/137 (53%)	69 (94%)	4 (6%)	0	100	100
4	AE	91/137 (66%)	89 (98%)	2 (2%)	0	100	100
4	AF	73/137 (53%)	69 (94%)	4 (6%)	0	100	100
4	BB	77/137 (56%)	67 (87%)	10 (13%)	0	100	100
4	BE	77/137 (56%)	69 (90%)	8 (10%)	0	100	100
4	CA	91/137 (66%)	89 (98%)	2 (2%)	0	100	100
4	CB	73/137 (53%)	69 (94%)	4 (6%)	0	100	100
4	CD	91/137 (66%)	90 (99%)	1 (1%)	0	100	100
4	CE	73/137 (53%)	68 (93%)	5 (7%)	0	100	100
4	CG	91/137 (66%)	89 (98%)	2 (2%)	0	100	100
4	D	91/137 (66%)	89 (98%)	2 (2%)	0	100	100
4	DA	77/137 (56%)	68 (88%)	9 (12%)	0	100	100
4	DD	77/137 (56%)	69 (90%)	8 (10%)	0	100	100
4	DG	77/137 (56%)	70 (91%)	7 (9%)	0	100	100
4	E	77/137 (56%)	68 (88%)	9 (12%)	0	100	100

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Mol	Chain	Analysed	Favoured	Allowed	Outliers	Percentiles	
4	EA	73/137 (53%)	69 (94%)	4 (6%)	0	100	100
4	EC	91/137 (66%)	88 (97%)	3 (3%)	0	100	100
4	ED	73/137 (53%)	69 (94%)	4 (6%)	0	100	100
4	EF	91/137 (66%)	91 (100%)	0	0	100	100
4	EG	73/137 (53%)	69 (94%)	4 (6%)	0	100	100
4	FC	77/137 (56%)	70 (91%)	7 (9%)	0	100	100
4	FF	77/137 (56%)	67 (87%)	10 (13%)	0	100	100
4	GB	91/137 (66%)	89 (98%)	2 (2%)	0	100	100
4	GC	73/137 (53%)	69 (94%)	4 (6%)	0	100	100
4	GE	91/137 (66%)	90 (99%)	1 (1%)	0	100	100
4	GF	73/137 (53%)	70 (96%)	3 (4%)	0	100	100
4	H	73/137 (53%)	70 (96%)	3 (4%)	0	100	100
4	HB	77/137 (56%)	67 (87%)	10 (13%)	0	100	100
4	HE	77/137 (56%)	69 (90%)	8 (10%)	0	100	100
4	IA	91/137 (66%)	90 (99%)	1 (1%)	0	100	100
4	IB	73/137 (53%)	69 (94%)	4 (6%)	0	100	100
4	ID	91/137 (66%)	89 (98%)	2 (2%)	0	100	100
4	IE	73/137 (53%)	67 (92%)	6 (8%)	0	100	100
4	IG	91/137 (66%)	89 (98%)	2 (2%)	0	100	100
4	JA	77/137 (56%)	69 (90%)	8 (10%)	0	100	100
4	JD	77/137 (56%)	69 (90%)	8 (10%)	0	100	100
4	JG	77/137 (56%)	66 (86%)	11 (14%)	0	100	100
4	KA	73/137 (53%)	68 (93%)	5 (7%)	0	100	100
4	KC	91/137 (66%)	89 (98%)	2 (2%)	0	100	100
4	KD	73/137 (53%)	70 (96%)	3 (4%)	0	100	100
4	KF	91/137 (66%)	89 (98%)	2 (2%)	0	100	100
4	KG	73/137 (53%)	69 (94%)	4 (6%)	0	100	100
4	L	91/137 (66%)	89 (98%)	2 (2%)	0	100	100
4	LC	77/137 (56%)	67 (87%)	10 (13%)	0	100	100
4	LF	77/137 (56%)	67 (87%)	10 (13%)	0	100	100
4	MB	91/137 (66%)	89 (98%)	2 (2%)	0	100	100

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Mol	Chain	Analysed	Favoured	Allowed	Outliers	Percentiles	
4	MC	73/137 (53%)	69 (94%)	4 (6%)	0	100	100
4	ME	91/137 (66%)	90 (99%)	1 (1%)	0	100	100
4	MF	73/137 (53%)	68 (93%)	5 (7%)	0	100	100
4	N	91/137 (66%)	89 (98%)	2 (2%)	0	100	100
4	NB	77/137 (56%)	68 (88%)	9 (12%)	0	100	100
4	NE	77/137 (56%)	68 (88%)	9 (12%)	0	100	100
4	O	77/137 (56%)	69 (90%)	8 (10%)	0	100	100
4	OA	91/137 (66%)	89 (98%)	2 (2%)	0	100	100
4	OB	73/137 (53%)	68 (93%)	5 (7%)	0	100	100
4	OD	91/137 (66%)	89 (98%)	2 (2%)	0	100	100
4	OE	73/137 (53%)	69 (94%)	4 (6%)	0	100	100
4	OG	91/137 (66%)	90 (99%)	1 (1%)	0	100	100
4	P	77/137 (56%)	69 (90%)	8 (10%)	0	100	100
4	PA	77/137 (56%)	67 (87%)	10 (13%)	0	100	100
4	PD	77/137 (56%)	66 (86%)	11 (14%)	0	100	100
4	PG	77/137 (56%)	68 (88%)	9 (12%)	0	100	100
4	Q	73/137 (53%)	69 (94%)	4 (6%)	0	100	100
4	QA	73/137 (53%)	69 (94%)	4 (6%)	0	100	100
4	QC	91/137 (66%)	90 (99%)	1 (1%)	0	100	100
4	QD	73/137 (53%)	67 (92%)	6 (8%)	0	100	100
4	QF	91/137 (66%)	90 (99%)	1 (1%)	0	100	100
4	QG	73/137 (53%)	69 (94%)	4 (6%)	0	100	100
4	R	73/137 (53%)	68 (93%)	5 (7%)	0	100	100
4	RC	77/137 (56%)	69 (90%)	8 (10%)	0	100	100
4	RF	77/137 (56%)	68 (88%)	9 (12%)	0	100	100
4	SB	91/137 (66%)	90 (99%)	1 (1%)	0	100	100
4	SC	73/137 (53%)	70 (96%)	3 (4%)	0	100	100
4	SE	91/137 (66%)	89 (98%)	2 (2%)	0	100	100
4	SF	73/137 (53%)	69 (94%)	4 (6%)	0	100	100
4	TB	77/137 (56%)	69 (90%)	8 (10%)	0	100	100
4	TE	77/137 (56%)	70 (91%)	7 (9%)	0	100	100

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Mol	Chain	Analysed	Favoured	Allowed	Outliers	Percentiles	
4	UA	91/137 (66%)	89 (98%)	2 (2%)	0	100	100
4	UB	73/137 (53%)	68 (93%)	5 (7%)	0	100	100
4	UD	91/137 (66%)	89 (98%)	2 (2%)	0	100	100
4	UE	73/137 (53%)	69 (94%)	4 (6%)	0	100	100
4	UG	91/137 (66%)	90 (99%)	1 (1%)	0	100	100
4	VA	77/137 (56%)	70 (91%)	7 (9%)	0	100	100
4	VD	77/137 (56%)	68 (88%)	9 (12%)	0	100	100
4	VG	77/137 (56%)	66 (86%)	11 (14%)	0	100	100
4	W	91/137 (66%)	89 (98%)	2 (2%)	0	100	100
4	WA	73/137 (53%)	69 (94%)	4 (6%)	0	100	100
4	WC	91/137 (66%)	90 (99%)	1 (1%)	0	100	100
4	WD	73/137 (53%)	67 (92%)	6 (8%)	0	100	100
4	WF	91/137 (66%)	89 (98%)	2 (2%)	0	100	100
4	WG	73/137 (53%)	69 (94%)	4 (6%)	0	100	100
4	X	77/137 (56%)	67 (87%)	10 (13%)	0	100	100
4	XC	77/137 (56%)	67 (87%)	10 (13%)	0	100	100
4	XF	77/137 (56%)	68 (88%)	9 (12%)	0	100	100
4	Y	73/137 (53%)	69 (94%)	4 (6%)	0	100	100
4	YB	91/137 (66%)	89 (98%)	2 (2%)	0	100	100
4	YC	73/137 (53%)	68 (93%)	5 (7%)	0	100	100
4	YE	91/137 (66%)	89 (98%)	2 (2%)	0	100	100
4	YF	73/137 (53%)	69 (94%)	4 (6%)	0	100	100
4	ZB	77/137 (56%)	68 (88%)	9 (12%)	0	100	100
4	ZE	77/137 (56%)	67 (87%)	10 (13%)	0	100	100
All	All	30022/37026 (81%)	28539 (95%)	1483 (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		
2	AA	271/282 (96%)	256 (94%)	15 (6%)	21	50	
2	AD	271/282 (96%)	255 (94%)	16 (6%)	19	48	
2	AG	271/282 (96%)	258 (95%)	13 (5%)	25	53	
2	B	271/282 (96%)	260 (96%)	11 (4%)	30	57	
2	CC	271/282 (96%)	258 (95%)	13 (5%)	25	53	
2	CF	271/282 (96%)	259 (96%)	12 (4%)	28	55	
2	EB	271/282 (96%)	259 (96%)	12 (4%)	28	55	
2	EE	271/282 (96%)	255 (94%)	16 (6%)	19	48	
2	G	271/282 (96%)	257 (95%)	14 (5%)	23	51	
2	GA	271/282 (96%)	256 (94%)	15 (6%)	21	50	
2	GD	271/282 (96%)	257 (95%)	14 (5%)	23	51	
2	GG	271/282 (96%)	258 (95%)	13 (5%)	25	53	
2	IC	271/282 (96%)	256 (94%)	15 (6%)	21	50	
2	IF	271/282 (96%)	257 (95%)	14 (5%)	23	51	
2	J	271/282 (96%)	256 (94%)	15 (6%)	21	50	
2	KB	271/282 (96%)	258 (95%)	13 (5%)	25	53	
2	KE	271/282 (96%)	259 (96%)	12 (4%)	28	55	
2	MA	271/282 (96%)	258 (95%)	13 (5%)	25	53	
2	MD	271/282 (96%)	259 (96%)	12 (4%)	28	55	
2	MG	271/282 (96%)	255 (94%)	16 (6%)	19	48	
2	OC	271/282 (96%)	254 (94%)	17 (6%)	18	46	
2	OF	271/282 (96%)	253 (93%)	18 (7%)	16	45	
2	QB	271/282 (96%)	257 (95%)	14 (5%)	23	51	
2	QE	271/282 (96%)	258 (95%)	13 (5%)	25	53	
2	SA	271/282 (96%)	255 (94%)	16 (6%)	19	48	
2	SD	271/282 (96%)	255 (94%)	16 (6%)	19	48	
2	SG	271/282 (96%)	258 (95%)	13 (5%)	25	53	
2	T	271/282 (96%)	255 (94%)	16 (6%)	19	48	
2	UC	271/282 (96%)	259 (96%)	12 (4%)	28	55	
2	UF	271/282 (96%)	258 (95%)	13 (5%)	25	53	

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Mol	Chain	Analysed	Rotameric	Outliers	Percentiles	
2	WB	271/282 (96%)	255 (94%)	16 (6%)	19	48
2	WE	271/282 (96%)	258 (95%)	13 (5%)	25	53
2	YA	271/282 (96%)	255 (94%)	16 (6%)	19	48
2	YD	271/282 (96%)	256 (94%)	15 (6%)	21	50
3	BA	177/257 (69%)	166 (94%)	11 (6%)	18	46
3	BD	177/257 (69%)	168 (95%)	9 (5%)	24	52
3	BG	177/257 (69%)	170 (96%)	7 (4%)	31	57
3	C	177/257 (69%)	169 (96%)	8 (4%)	27	55
3	DC	177/257 (69%)	170 (96%)	7 (4%)	31	57
3	DF	177/257 (69%)	168 (95%)	9 (5%)	24	52
3	FB	177/257 (69%)	168 (95%)	9 (5%)	24	52
3	FE	177/257 (69%)	169 (96%)	8 (4%)	27	55
3	HA	177/257 (69%)	169 (96%)	8 (4%)	27	55
3	HD	177/257 (69%)	169 (96%)	8 (4%)	27	55
3	HG	177/257 (69%)	168 (95%)	9 (5%)	24	52
3	JC	177/257 (69%)	167 (94%)	10 (6%)	21	49
3	JF	177/257 (69%)	168 (95%)	9 (5%)	24	52
3	K	177/257 (69%)	168 (95%)	9 (5%)	24	52
3	LB	177/257 (69%)	166 (94%)	11 (6%)	18	46
3	LE	177/257 (69%)	167 (94%)	10 (6%)	21	49
3	M	177/257 (69%)	170 (96%)	7 (4%)	31	57
3	NA	177/257 (69%)	167 (94%)	10 (6%)	21	49
3	ND	177/257 (69%)	169 (96%)	8 (4%)	27	55
3	NG	177/257 (69%)	165 (93%)	12 (7%)	16	44
3	PC	177/257 (69%)	167 (94%)	10 (6%)	21	49
3	PF	177/257 (69%)	166 (94%)	11 (6%)	18	46
3	RB	177/257 (69%)	170 (96%)	7 (4%)	31	57
3	RE	177/257 (69%)	167 (94%)	10 (6%)	21	49
3	TA	177/257 (69%)	168 (95%)	9 (5%)	24	52
3	TD	177/257 (69%)	165 (93%)	12 (7%)	16	44
3	TG	177/257 (69%)	170 (96%)	7 (4%)	31	57

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Mol	Chain	Analysed	Rotameric	Outliers	Percentiles	
3	V	177/257 (69%)	164 (93%)	13 (7%)	14	42
3	VC	177/257 (69%)	168 (95%)	9 (5%)	24	52
3	VF	177/257 (69%)	165 (93%)	12 (7%)	16	44
3	XB	177/257 (69%)	169 (96%)	8 (4%)	27	55
3	XE	177/257 (69%)	170 (96%)	7 (4%)	31	57
3	ZA	177/257 (69%)	170 (96%)	7 (4%)	31	57
3	ZD	177/257 (69%)	165 (93%)	12 (7%)	16	44
All	All	15232/18326 (83%)	14437 (95%)	795 (5%)	27	51

All (795) residues with a non-rotameric sidechain are listed below:

Mol	Chain	Res	Type
2	G	14	LEU
2	G	21	ARG
2	G	26	PHE
2	G	59	PHE
2	G	72	ASN
2	G	78	ARG
2	G	92	SER
2	G	174	GLU
2	G	188	LEU
2	G	193	MET
2	G	206	MET
2	G	236	PHE
2	G	264	LYS
2	G	298	LEU
3	M	58	ASN
3	M	72	LEU
3	M	89	TYR
3	M	94	ARG
3	M	111	ARG
3	M	134	ASP
3	M	228	ARG
2	B	14	LEU
2	B	32	ARG
2	B	59	PHE
2	B	69	LEU
2	B	78	ARG
2	B	89	ARG
2	B	189	LYS

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Mol	Chain	Res	Type
2	B	193	MET
2	B	204	ASN
2	B	264	LYS
2	B	298	LEU
3	C	58	ASN
3	C	72	LEU
3	C	89	TYR
3	C	106	HIS
3	C	214	GLU
3	C	227	LEU
3	C	228	ARG
3	C	238	ASN
2	J	21	ARG
2	J	26	PHE
2	J	59	PHE
2	J	69	LEU
2	J	74	ASN
2	J	78	ARG
2	J	89	ARG
2	J	174	GLU
2	J	189	LYS
2	J	193	MET
2	J	206	MET
2	J	264	LYS
2	J	284	ASP
2	J	298	LEU
2	J	321	MET
3	K	72	LEU
3	K	89	TYR
3	K	94	ARG
3	K	111	ARG
3	K	156	ARG
3	K	214	GLU
3	K	227	LEU
3	K	228	ARG
3	K	238	ASN
2	T	21	ARG
2	T	26	PHE
2	T	32	ARG
2	T	59	PHE
2	T	69	LEU
2	T	78	ARG

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Mol	Chain	Res	Type
2	T	79	SER
2	T	89	ARG
2	T	126	HIS
2	T	158	MET
2	T	174	GLU
2	T	189	LYS
2	T	193	MET
2	T	207	LYS
2	T	264	LYS
2	T	298	LEU
3	V	58	ASN
3	V	72	LEU
3	V	89	TYR
3	V	106	HIS
3	V	111	ARG
3	V	156	ARG
3	V	183	GLU
3	V	197	ASP
3	V	205	HIS
3	V	214	GLU
3	V	228	ARG
3	V	238	ASN
3	V	247	ARG
2	AA	21	ARG
2	AA	26	PHE
2	AA	37	LEU
2	AA	59	PHE
2	AA	69	LEU
2	AA	72	ASN
2	AA	78	ARG
2	AA	79	SER
2	AA	89	ARG
2	AA	92	SER
2	AA	138	LYS
2	AA	193	MET
2	AA	244	ASP
2	AA	264	LYS
2	AA	298	LEU
3	BA	72	LEU
3	BA	89	TYR
3	BA	106	HIS
3	BA	134	ASP

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Mol	Chain	Res	Type
3	BA	156	ARG
3	BA	183	GLU
3	BA	197	ASP
3	BA	205	HIS
3	BA	214	GLU
3	BA	228	ARG
3	BA	238	ASN
2	GA	21	ARG
2	GA	26	PHE
2	GA	41	MET
2	GA	59	PHE
2	GA	69	LEU
2	GA	74	ASN
2	GA	78	ARG
2	GA	79	SER
2	GA	108	GLU
2	GA	152	ARG
2	GA	189	LYS
2	GA	193	MET
2	GA	264	LYS
2	GA	298	LEU
2	GA	321	MET
3	HA	58	ASN
3	HA	72	LEU
3	HA	89	TYR
3	HA	94	ARG
3	HA	111	ARG
3	HA	134	ASP
3	HA	156	ARG
3	HA	238	ASN
2	MA	26	PHE
2	MA	32	ARG
2	MA	41	MET
2	MA	59	PHE
2	MA	69	LEU
2	MA	78	ARG
2	MA	79	SER
2	MA	92	SER
2	MA	174	GLU
2	MA	189	LYS
2	MA	193	MET
2	MA	264	LYS

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Mol	Chain	Res	Type
2	MA	298	LEU
3	NA	72	LEU
3	NA	89	TYR
3	NA	94	ARG
3	NA	111	ARG
3	NA	134	ASP
3	NA	205	HIS
3	NA	214	GLU
3	NA	228	ARG
3	NA	238	ASN
3	NA	247	ARG
2	SA	26	PHE
2	SA	37	LEU
2	SA	59	PHE
2	SA	72	ASN
2	SA	76	TYR
2	SA	78	ARG
2	SA	79	SER
2	SA	92	SER
2	SA	152	ARG
2	SA	174	GLU
2	SA	188	LEU
2	SA	189	LYS
2	SA	193	MET
2	SA	206	MET
2	SA	264	LYS
2	SA	321	MET
3	TA	72	LEU
3	TA	89	TYR
3	TA	111	ARG
3	TA	134	ASP
3	TA	183	GLU
3	TA	214	GLU
3	TA	228	ARG
3	TA	238	ASN
3	TA	250	LEU
2	YA	21	ARG
2	YA	26	PHE
2	YA	59	PHE
2	YA	74	ASN
2	YA	78	ARG
2	YA	79	SER

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Mol	Chain	Res	Type
2	YA	92	SER
2	YA	108	GLU
2	YA	126	HIS
2	YA	148	LEU
2	YA	174	GLU
2	YA	188	LEU
2	YA	189	LYS
2	YA	193	MET
2	YA	264	LYS
2	YA	298	LEU
3	ZA	58	ASN
3	ZA	72	LEU
3	ZA	89	TYR
3	ZA	134	ASP
3	ZA	214	GLU
3	ZA	228	ARG
3	ZA	231	LEU
2	EB	21	ARG
2	EB	26	PHE
2	EB	32	ARG
2	EB	59	PHE
2	EB	72	ASN
2	EB	78	ARG
2	EB	92	SER
2	EB	126	HIS
2	EB	189	LYS
2	EB	193	MET
2	EB	264	LYS
2	EB	298	LEU
3	FB	58	ASN
3	FB	72	LEU
3	FB	89	TYR
3	FB	106	HIS
3	FB	197	ASP
3	FB	214	GLU
3	FB	227	LEU
3	FB	228	ARG
3	FB	247	ARG
2	KB	32	ARG
2	KB	37	LEU
2	KB	59	PHE
2	KB	72	ASN

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Mol	Chain	Res	Type
2	KB	76	TYR
2	KB	78	ARG
2	KB	89	ARG
2	KB	108	GLU
2	KB	189	LYS
2	KB	193	MET
2	KB	264	LYS
2	KB	298	LEU
2	KB	321	MET
3	LB	72	LEU
3	LB	89	TYR
3	LB	94	ARG
3	LB	106	HIS
3	LB	111	ARG
3	LB	156	ARG
3	LB	183	GLU
3	LB	190	ASN
3	LB	228	ARG
3	LB	238	ASN
3	LB	247	ARG
2	QB	21	ARG
2	QB	26	PHE
2	QB	41	MET
2	QB	59	PHE
2	QB	78	ARG
2	QB	89	ARG
2	QB	126	HIS
2	QB	158	MET
2	QB	174	GLU
2	QB	193	MET
2	QB	219	ARG
2	QB	264	LYS
2	QB	298	LEU
2	QB	321	MET
3	RB	58	ASN
3	RB	72	LEU
3	RB	89	TYR
3	RB	134	ASP
3	RB	183	GLU
3	RB	214	GLU
3	RB	228	ARG
2	WB	21	ARG

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Mol	Chain	Res	Type
2	WB	27	LYS
2	WB	32	ARG
2	WB	37	LEU
2	WB	59	PHE
2	WB	78	ARG
2	WB	79	SER
2	WB	81	LEU
2	WB	92	SER
2	WB	108	GLU
2	WB	138	LYS
2	WB	158	MET
2	WB	193	MET
2	WB	207	LYS
2	WB	264	LYS
2	WB	298	LEU
3	XB	72	LEU
3	XB	89	TYR
3	XB	94	ARG
3	XB	106	HIS
3	XB	183	GLU
3	XB	214	GLU
3	XB	228	ARG
3	XB	238	ASN
2	CC	21	ARG
2	CC	59	PHE
2	CC	78	ARG
2	CC	92	SER
2	CC	126	HIS
2	CC	148	LEU
2	CC	158	MET
2	CC	189	LYS
2	CC	206	MET
2	CC	244	ASP
2	CC	264	LYS
2	CC	298	LEU
2	CC	321	MET
3	DC	72	LEU
3	DC	89	TYR
3	DC	106	HIS
3	DC	156	ARG
3	DC	194	SER
3	DC	227	LEU

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Mol	Chain	Res	Type
3	DC	238	ASN
2	IC	21	ARG
2	IC	26	PHE
2	IC	41	MET
2	IC	45	ARG
2	IC	59	PHE
2	IC	72	ASN
2	IC	78	ARG
2	IC	79	SER
2	IC	89	ARG
2	IC	92	SER
2	IC	148	LEU
2	IC	189	LYS
2	IC	204	ASN
2	IC	264	LYS
2	IC	298	LEU
3	JC	72	LEU
3	JC	89	TYR
3	JC	134	ASP
3	JC	156	ARG
3	JC	183	GLU
3	JC	197	ASP
3	JC	205	HIS
3	JC	228	ARG
3	JC	238	ASN
3	JC	247	ARG
2	OC	26	PHE
2	OC	41	MET
2	OC	59	PHE
2	OC	69	LEU
2	OC	78	ARG
2	OC	79	SER
2	OC	89	ARG
2	OC	92	SER
2	OC	126	HIS
2	OC	152	ARG
2	OC	164	PHE
2	OC	174	GLU
2	OC	189	LYS
2	OC	193	MET
2	OC	207	LYS
2	OC	264	LYS

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Mol	Chain	Res	Type
2	OC	298	LEU
3	PC	72	LEU
3	PC	89	TYR
3	PC	106	HIS
3	PC	111	ARG
3	PC	156	ARG
3	PC	183	GLU
3	PC	205	HIS
3	PC	214	GLU
3	PC	228	ARG
3	PC	238	ASN
2	UC	26	PHE
2	UC	37	LEU
2	UC	59	PHE
2	UC	69	LEU
2	UC	72	ASN
2	UC	74	ASN
2	UC	78	ARG
2	UC	89	ARG
2	UC	189	LYS
2	UC	193	MET
2	UC	264	LYS
2	UC	298	LEU
3	VC	72	LEU
3	VC	89	TYR
3	VC	94	ARG
3	VC	106	HIS
3	VC	111	ARG
3	VC	134	ASP
3	VC	156	ARG
3	VC	205	HIS
3	VC	228	ARG
2	AD	21	ARG
2	AD	59	PHE
2	AD	69	LEU
2	AD	72	ASN
2	AD	74	ASN
2	AD	78	ARG
2	AD	89	ARG
2	AD	174	GLU
2	AD	188	LEU
2	AD	189	LYS

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Mol	Chain	Res	Type
2	AD	193	MET
2	AD	204	ASN
2	AD	206	MET
2	AD	236	PHE
2	AD	264	LYS
2	AD	298	LEU
3	BD	58	ASN
3	BD	72	LEU
3	BD	89	TYR
3	BD	94	ARG
3	BD	134	ASP
3	BD	228	ARG
3	BD	238	ASN
3	BD	247	ARG
3	BD	252	ARG
2	GD	21	ARG
2	GD	26	PHE
2	GD	32	ARG
2	GD	59	PHE
2	GD	69	LEU
2	GD	74	ASN
2	GD	78	ARG
2	GD	89	ARG
2	GD	158	MET
2	GD	193	MET
2	GD	201	GLU
2	GD	204	ASN
2	GD	264	LYS
2	GD	298	LEU
3	HD	58	ASN
3	HD	72	LEU
3	HD	89	TYR
3	HD	106	HIS
3	HD	214	GLU
3	HD	227	LEU
3	HD	228	ARG
3	HD	238	ASN
2	MD	26	PHE
2	MD	59	PHE
2	MD	78	ARG
2	MD	79	SER
2	MD	89	ARG

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Mol	Chain	Res	Type
2	MD	174	GLU
2	MD	189	LYS
2	MD	193	MET
2	MD	264	LYS
2	MD	284	ASP
2	MD	298	LEU
2	MD	321	MET
3	ND	72	LEU
3	ND	89	TYR
3	ND	94	ARG
3	ND	183	GLU
3	ND	214	GLU
3	ND	227	LEU
3	ND	228	ARG
3	ND	238	ASN
2	SD	26	PHE
2	SD	32	ARG
2	SD	37	LEU
2	SD	41	MET
2	SD	59	PHE
2	SD	78	ARG
2	SD	89	ARG
2	SD	108	GLU
2	SD	158	MET
2	SD	174	GLU
2	SD	189	LYS
2	SD	197	ARG
2	SD	207	LYS
2	SD	209	GLN
2	SD	264	LYS
2	SD	298	LEU
3	TD	58	ASN
3	TD	72	LEU
3	TD	89	TYR
3	TD	106	HIS
3	TD	156	ARG
3	TD	183	GLU
3	TD	205	HIS
3	TD	212	THR
3	TD	221	PHE
3	TD	228	ARG
3	TD	238	ASN

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Mol	Chain	Res	Type
3	TD	247	ARG
2	YD	21	ARG
2	YD	26	PHE
2	YD	27	LYS
2	YD	37	LEU
2	YD	59	PHE
2	YD	69	LEU
2	YD	72	ASN
2	YD	78	ARG
2	YD	79	SER
2	YD	92	SER
2	YD	148	LEU
2	YD	193	MET
2	YD	244	ASP
2	YD	264	LYS
2	YD	298	LEU
3	ZD	58	ASN
3	ZD	72	LEU
3	ZD	89	TYR
3	ZD	106	HIS
3	ZD	111	ARG
3	ZD	156	ARG
3	ZD	183	GLU
3	ZD	197	ASP
3	ZD	214	GLU
3	ZD	227	LEU
3	ZD	228	ARG
3	ZD	238	ASN
2	EE	21	ARG
2	EE	26	PHE
2	EE	32	ARG
2	EE	41	MET
2	EE	59	PHE
2	EE	69	LEU
2	EE	74	ASN
2	EE	78	ARG
2	EE	79	SER
2	EE	92	SER
2	EE	152	ARG
2	EE	174	GLU
2	EE	193	MET
2	EE	264	LYS

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Mol	Chain	Res	Type
2	EE	298	LEU
2	EE	321	MET
3	FE	58	ASN
3	FE	72	LEU
3	FE	89	TYR
3	FE	111	ARG
3	FE	134	ASP
3	FE	190	ASN
3	FE	238	ASN
3	FE	252	ARG
2	KE	26	PHE
2	KE	32	ARG
2	KE	37	LEU
2	KE	41	MET
2	KE	59	PHE
2	KE	78	ARG
2	KE	89	ARG
2	KE	126	HIS
2	KE	139	ARG
2	KE	174	GLU
2	KE	264	LYS
2	KE	298	LEU
3	LE	72	LEU
3	LE	89	TYR
3	LE	134	ASP
3	LE	183	GLU
3	LE	205	HIS
3	LE	214	GLU
3	LE	228	ARG
3	LE	238	ASN
3	LE	247	ARG
3	LE	252	ARG
2	QE	26	PHE
2	QE	59	PHE
2	QE	69	LEU
2	QE	78	ARG
2	QE	79	SER
2	QE	152	ARG
2	QE	159	LEU
2	QE	168	GLN
2	QE	174	GLU
2	QE	206	MET

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Mol	Chain	Res	Type
2	QE	236	PHE
2	QE	264	LYS
2	QE	298	LEU
3	RE	72	LEU
3	RE	89	TYR
3	RE	111	ARG
3	RE	156	ARG
3	RE	183	GLU
3	RE	197	ASP
3	RE	205	HIS
3	RE	228	ARG
3	RE	238	ASN
3	RE	250	LEU
2	WE	26	PHE
2	WE	32	ARG
2	WE	59	PHE
2	WE	76	TYR
2	WE	78	ARG
2	WE	89	ARG
2	WE	126	HIS
2	WE	148	LEU
2	WE	174	GLU
2	WE	193	MET
2	WE	207	LYS
2	WE	264	LYS
2	WE	298	LEU
3	XE	58	ASN
3	XE	72	LEU
3	XE	89	TYR
3	XE	134	ASP
3	XE	156	ARG
3	XE	214	GLU
3	XE	228	ARG
2	CF	21	ARG
2	CF	26	PHE
2	CF	32	ARG
2	CF	59	PHE
2	CF	72	ASN
2	CF	78	ARG
2	CF	89	ARG
2	CF	174	GLU
2	CF	193	MET

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Mol	Chain	Res	Type
2	CF	201	GLU
2	CF	264	LYS
2	CF	298	LEU
3	DF	58	ASN
3	DF	72	LEU
3	DF	89	TYR
3	DF	111	ARG
3	DF	183	GLU
3	DF	214	GLU
3	DF	228	ARG
3	DF	238	ASN
3	DF	247	ARG
2	IF	27	LYS
2	IF	32	ARG
2	IF	37	LEU
2	IF	59	PHE
2	IF	69	LEU
2	IF	72	ASN
2	IF	78	ARG
2	IF	89	ARG
2	IF	174	GLU
2	IF	189	LYS
2	IF	193	MET
2	IF	264	LYS
2	IF	298	LEU
2	IF	321	MET
3	JF	72	LEU
3	JF	89	TYR
3	JF	94	ARG
3	JF	106	HIS
3	JF	111	ARG
3	JF	156	ARG
3	JF	190	ASN
3	JF	228	ARG
3	JF	247	ARG
2	OF	21	ARG
2	OF	59	PHE
2	OF	69	LEU
2	OF	72	ASN
2	OF	74	ASN
2	OF	76	TYR
2	OF	78	ARG

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Mol	Chain	Res	Type
2	OF	79	SER
2	OF	83	LYS
2	OF	89	ARG
2	OF	108	GLU
2	OF	120	ASP
2	OF	126	HIS
2	OF	174	GLU
2	OF	193	MET
2	OF	264	LYS
2	OF	298	LEU
2	OF	321	MET
3	PF	58	ASN
3	PF	72	LEU
3	PF	89	TYR
3	PF	106	HIS
3	PF	111	ARG
3	PF	134	ASP
3	PF	183	GLU
3	PF	214	GLU
3	PF	227	LEU
3	PF	228	ARG
3	PF	238	ASN
2	UF	26	PHE
2	UF	27	LYS
2	UF	32	ARG
2	UF	59	PHE
2	UF	79	SER
2	UF	81	LEU
2	UF	92	SER
2	UF	158	MET
2	UF	174	GLU
2	UF	193	MET
2	UF	207	LYS
2	UF	264	LYS
2	UF	298	LEU
3	VF	72	LEU
3	VF	89	TYR
3	VF	94	ARG
3	VF	106	HIS
3	VF	111	ARG
3	VF	156	ARG
3	VF	183	GLU

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Mol	Chain	Res	Type
3	VF	184	MET
3	VF	190	ASN
3	VF	214	GLU
3	VF	228	ARG
3	VF	247	ARG
2	AG	21	ARG
2	AG	32	ARG
2	AG	59	PHE
2	AG	69	LEU
2	AG	78	ARG
2	AG	92	SER
2	AG	126	HIS
2	AG	158	MET
2	AG	193	MET
2	AG	244	ASP
2	AG	264	LYS
2	AG	298	LEU
2	AG	321	MET
3	BG	72	LEU
3	BG	89	TYR
3	BG	111	ARG
3	BG	156	ARG
3	BG	194	SER
3	BG	227	LEU
3	BG	238	ASN
2	GG	21	ARG
2	GG	59	PHE
2	GG	78	ARG
2	GG	89	ARG
2	GG	108	GLU
2	GG	120	ASP
2	GG	148	LEU
2	GG	174	GLU
2	GG	189	LYS
2	GG	193	MET
2	GG	201	GLU
2	GG	264	LYS
2	GG	298	LEU
3	HG	72	LEU
3	HG	89	TYR
3	HG	111	ARG
3	HG	134	ASP

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Mol	Chain	Res	Type
3	HG	156	ARG
3	HG	183	GLU
3	HG	197	ASP
3	HG	214	GLU
3	HG	228	ARG
2	MG	14	LEU
2	MG	21	ARG
2	MG	26	PHE
2	MG	41	MET
2	MG	59	PHE
2	MG	69	LEU
2	MG	74	ASN
2	MG	78	ARG
2	MG	89	ARG
2	MG	92	SER
2	MG	126	HIS
2	MG	152	ARG
2	MG	189	LYS
2	MG	207	LYS
2	MG	264	LYS
2	MG	298	LEU
3	NG	72	LEU
3	NG	74	ARG
3	NG	89	TYR
3	NG	106	HIS
3	NG	111	ARG
3	NG	134	ASP
3	NG	156	ARG
3	NG	197	ASP
3	NG	214	GLU
3	NG	227	LEU
3	NG	228	ARG
3	NG	238	ASN
2	SG	21	ARG
2	SG	26	PHE
2	SG	41	MET
2	SG	59	PHE
2	SG	69	LEU
2	SG	74	ASN
2	SG	76	TYR
2	SG	78	ARG
2	SG	152	ARG

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Mol	Chain	Res	Type
2	SG	204	ASN
2	SG	264	LYS
2	SG	298	LEU
2	SG	321	MET
3	TG	72	LEU
3	TG	89	TYR
3	TG	111	ARG
3	TG	134	ASP
3	TG	156	ARG
3	TG	205	HIS
3	TG	228	ARG

Sometimes sidechains can be flipped to improve hydrogen bonding and reduce clashes. All (67) such sidechains are listed below:

Mol	Chain	Res	Type
2	B	74	ASN
3	C	106	HIS
3	C	190	ASN
2	J	126	HIS
3	K	238	ASN
2	T	126	HIS
3	V	106	HIS
2	AA	186	GLN
3	BA	106	HIS
3	BA	190	ASN
2	GA	111	ASN
3	NA	190	ASN
2	SA	111	ASN
3	TA	190	ASN
2	YA	210	GLN
2	EB	210	GLN
3	FB	241	HIS
3	LB	71	ASN
3	LB	106	HIS
3	LB	238	ASN
3	LB	241	HIS
2	QB	72	ASN
2	QB	126	HIS
3	RB	106	HIS
3	RB	190	ASN
3	XB	106	HIS
3	XB	173	ASN

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Mol	Chain	Res	Type
3	XB	190	ASN
3	DC	106	HIS
3	DC	190	ASN
2	OC	126	HIS
3	PC	58	ASN
3	PC	106	HIS
3	VC	71	ASN
2	AD	72	ASN
3	HD	190	ASN
3	HD	238	ASN
2	MD	126	HIS
2	MD	210	GLN
3	ND	241	HIS
2	SD	28	HIS
3	TD	241	HIS
2	YD	186	GLN
3	ZD	190	ASN
2	KE	186	GLN
2	KE	210	GLN
2	QE	210	GLN
2	WE	126	HIS
2	WE	186	GLN
3	XE	90	HIS
2	CF	210	GLN
3	DF	241	HIS
2	IF	128	GLN
2	IF	141	GLN
2	IF	210	GLN
3	JF	106	HIS
3	JF	241	HIS
2	OF	72	ASN
2	OF	186	GLN
3	PF	106	HIS
2	UF	72	ASN
2	AG	186	GLN
2	GG	141	GLN
2	MG	126	HIS
3	NG	90	HIS
2	SG	141	GLN
3	TG	71	ASN

5.3.3 RNA [i](#)

There are no RNA molecules in this entry.

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

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

5.5 Carbohydrates [i](#)

There are no monosaccharides 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.

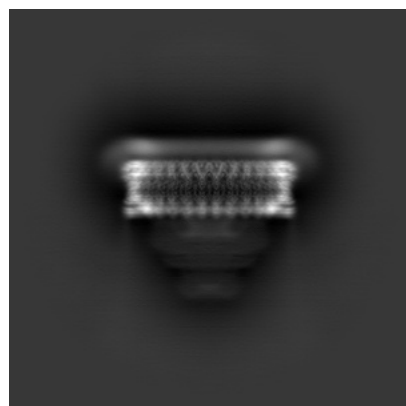
6 Map visualisation [i](#)

This section contains visualisations of the EMDB entry EMD-41100. These allow visual inspection of the internal detail of the map and identification of artifacts.

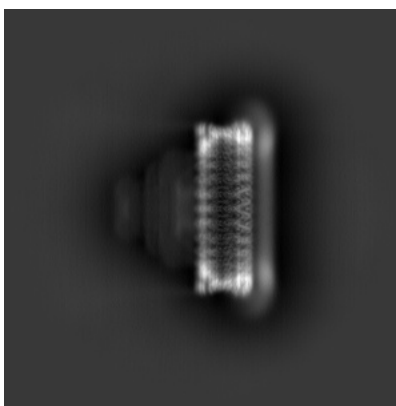
Images derived from a raw map, generated by summing the deposited half-maps, are presented below the corresponding image components of the primary map to allow further visual inspection and comparison with those of the primary map.

6.1 Orthogonal projections [i](#)

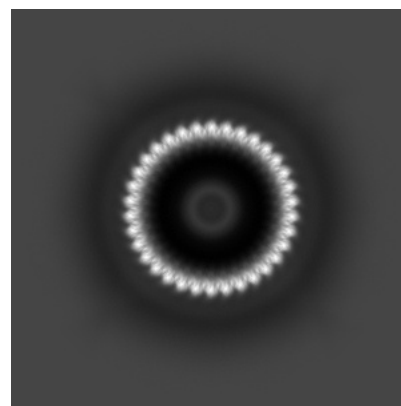
6.1.1 Primary map



X

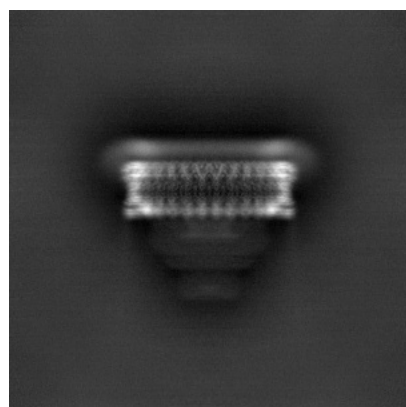


Y

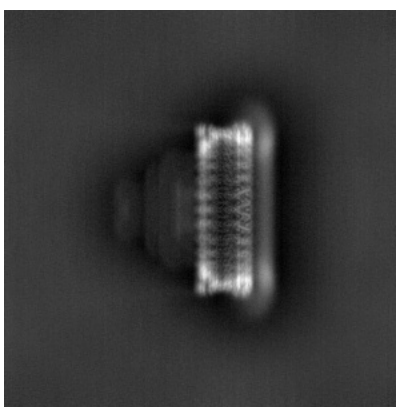


Z

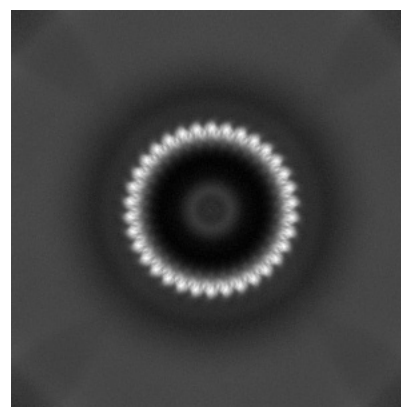
6.1.2 Raw map



X



Y

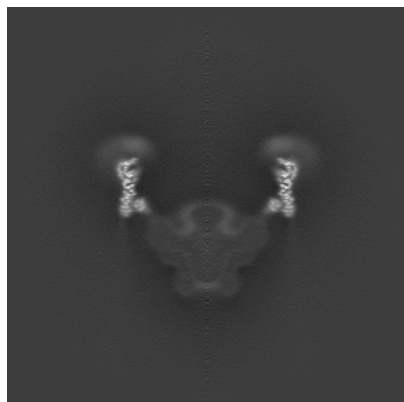


Z

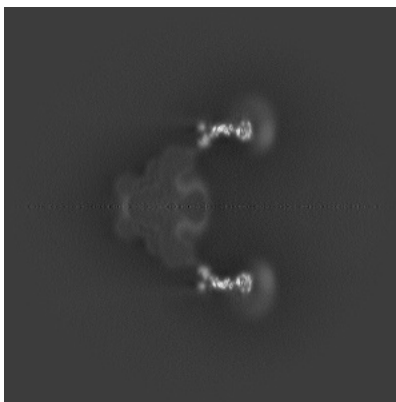
The images above show the map projected in three orthogonal directions.

6.2 Central slices [i](#)

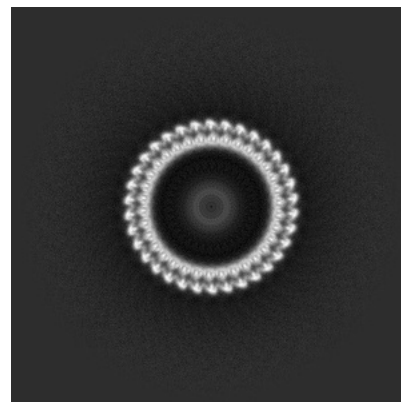
6.2.1 Primary map



X Index: 384

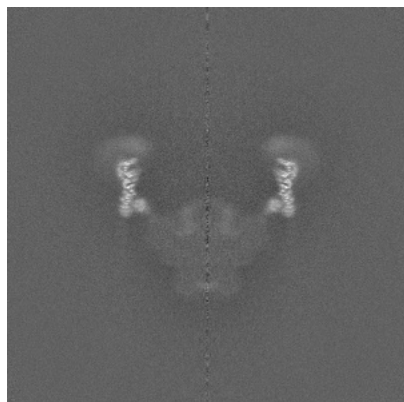


Y Index: 384

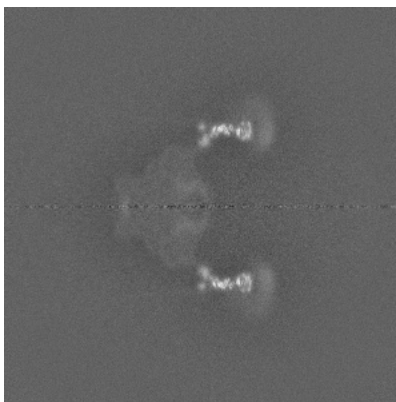


Z Index: 384

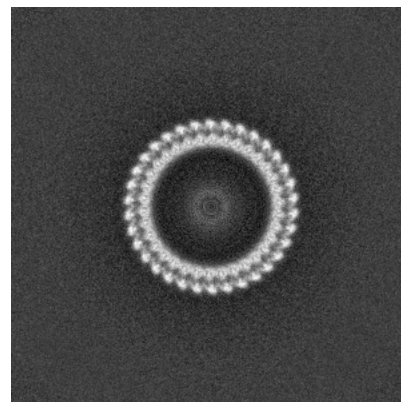
6.2.2 Raw map



X Index: 384



Y Index: 384

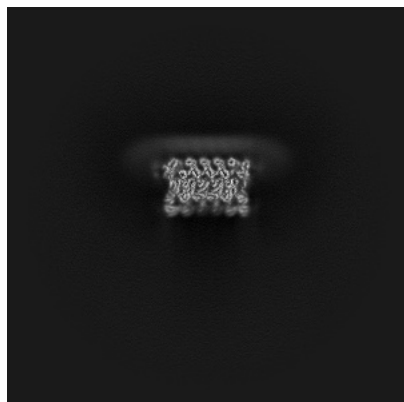


Z Index: 384

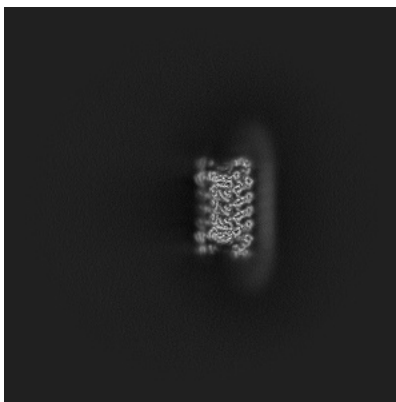
The images above show central slices of the map in three orthogonal directions.

6.3 Largest variance slices [i](#)

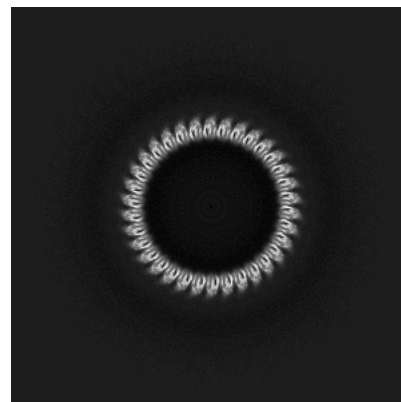
6.3.1 Primary map



X Index: 241

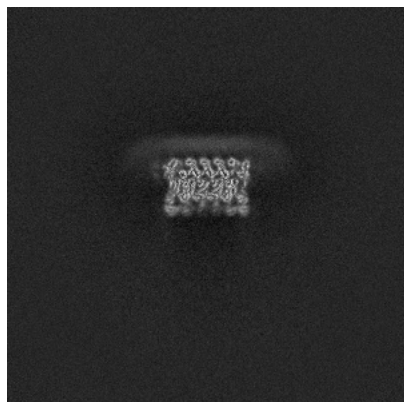


Y Index: 242

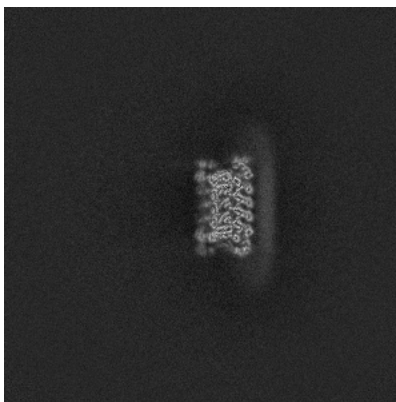


Z Index: 449

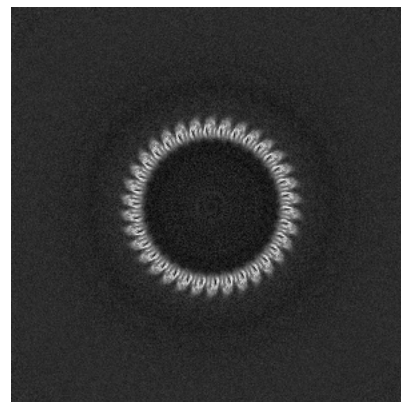
6.3.2 Raw map



X Index: 241



Y Index: 526

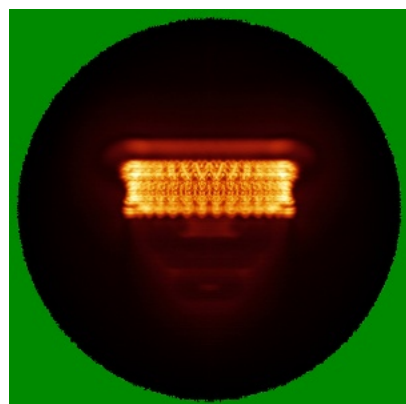


Z Index: 449

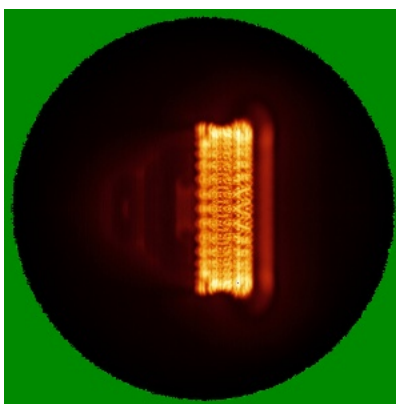
The images above show the largest variance slices of the map in three orthogonal directions.

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

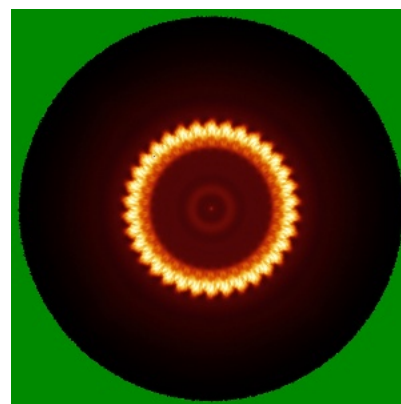
6.4.1 Primary map



X

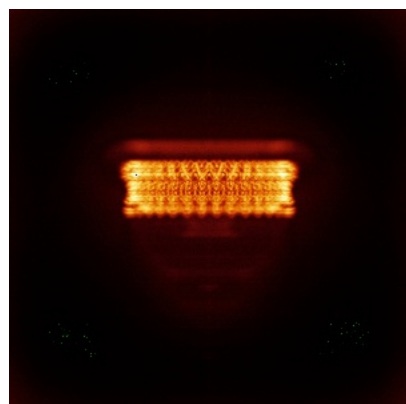


Y

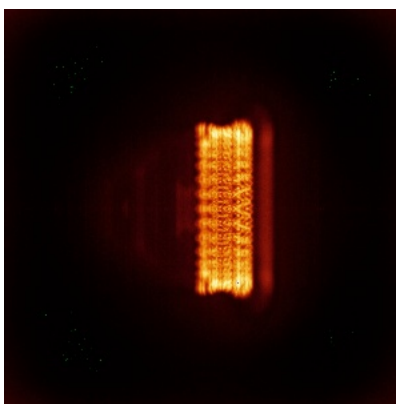


Z

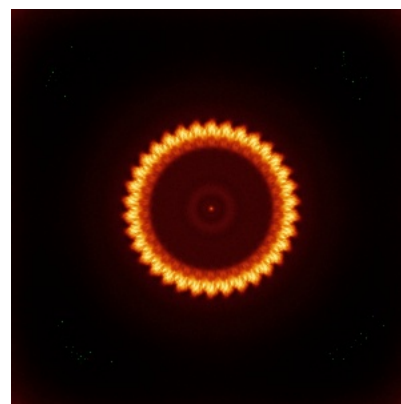
6.4.2 Raw map



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.

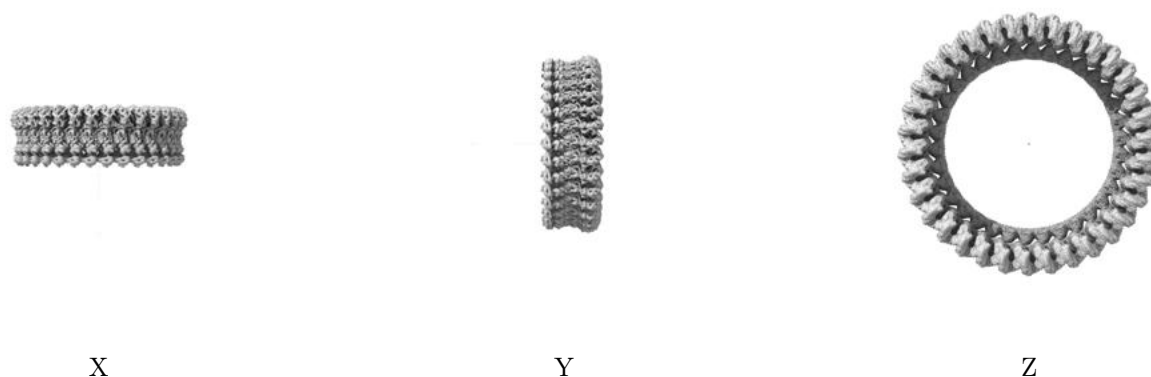
6.5 Orthogonal surface views [i](#)

6.5.1 Primary map



The images above show the 3D surface view of the map at the recommended contour level 0.147. 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.

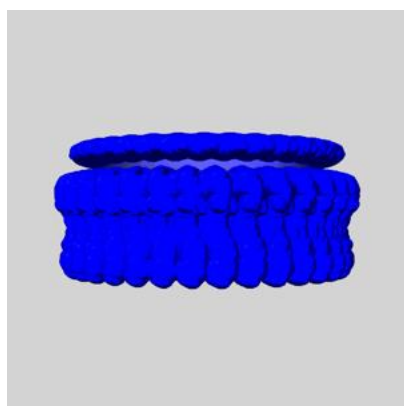
6.6 Mask visualisation [i](#)

This section shows the 3D surface view of the primary map at 50% transparency overlaid with the specified mask at 0% transparency

A mask typically either:

- Encompasses the whole structure
- Separates out a domain, a functional unit, a monomer or an area of interest from a larger structure

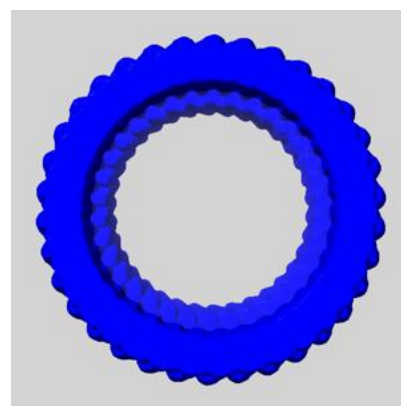
6.6.1 emd_41100_msk_1.map [i](#)



X



Y

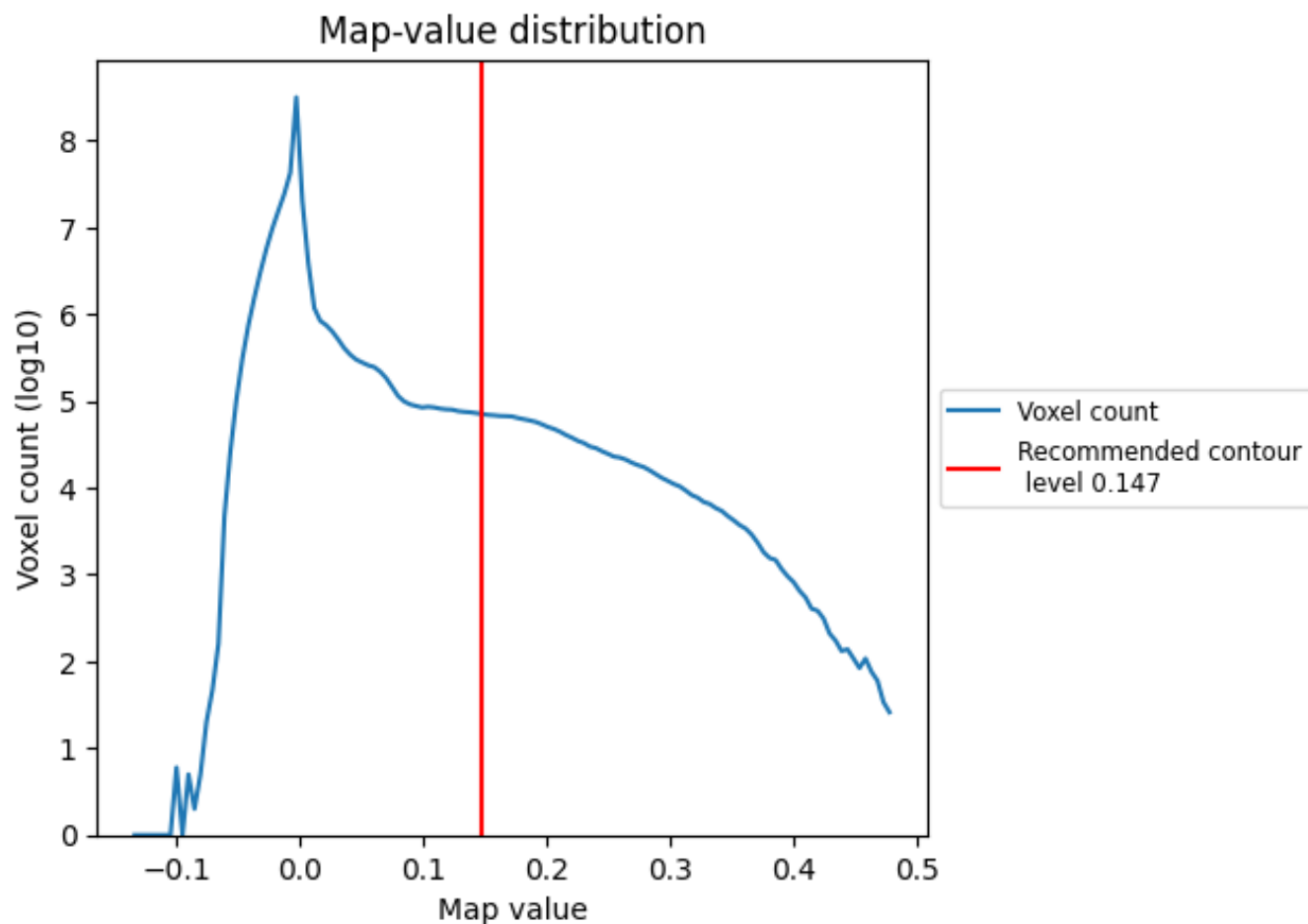


Z

7 Map analysis [i](#)

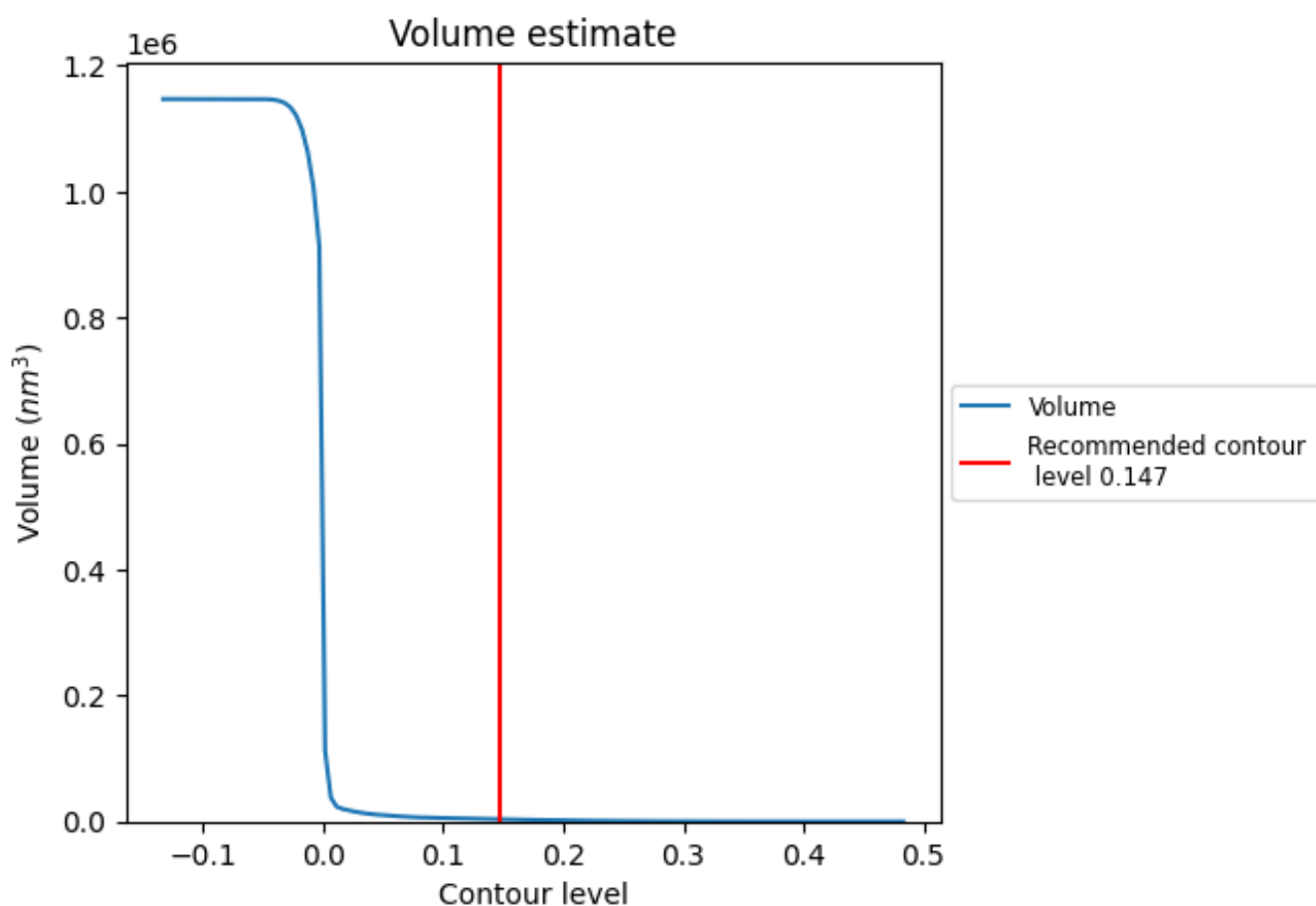
This section contains the results of statistical analysis of the map.

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.

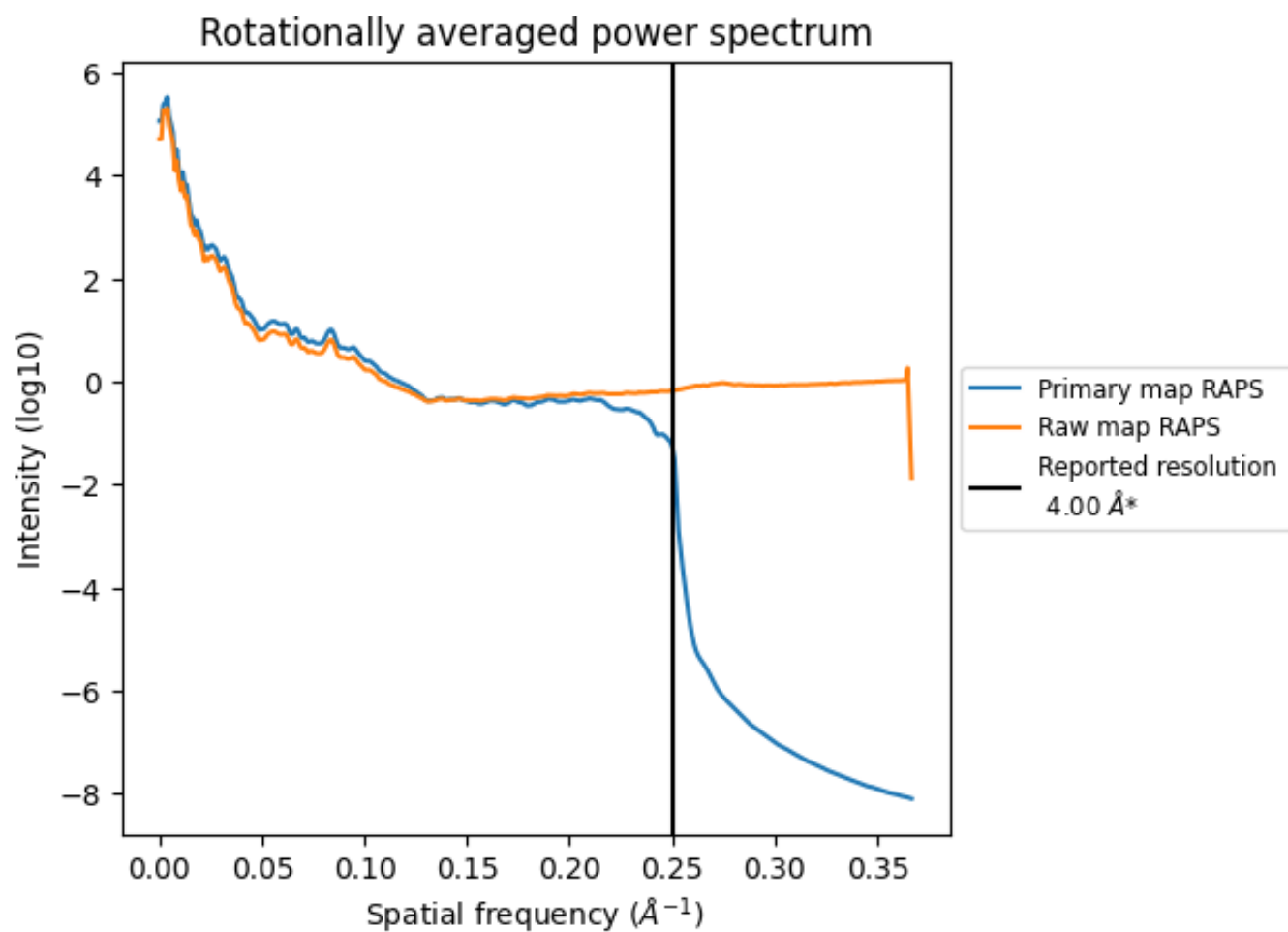
7.2 Volume estimate [i](#)



The volume at the recommended contour level is 3549 nm^3 ; this corresponds to an approximate mass of 3206 kDa.

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

7.3 Rotationally averaged power spectrum [i](#)

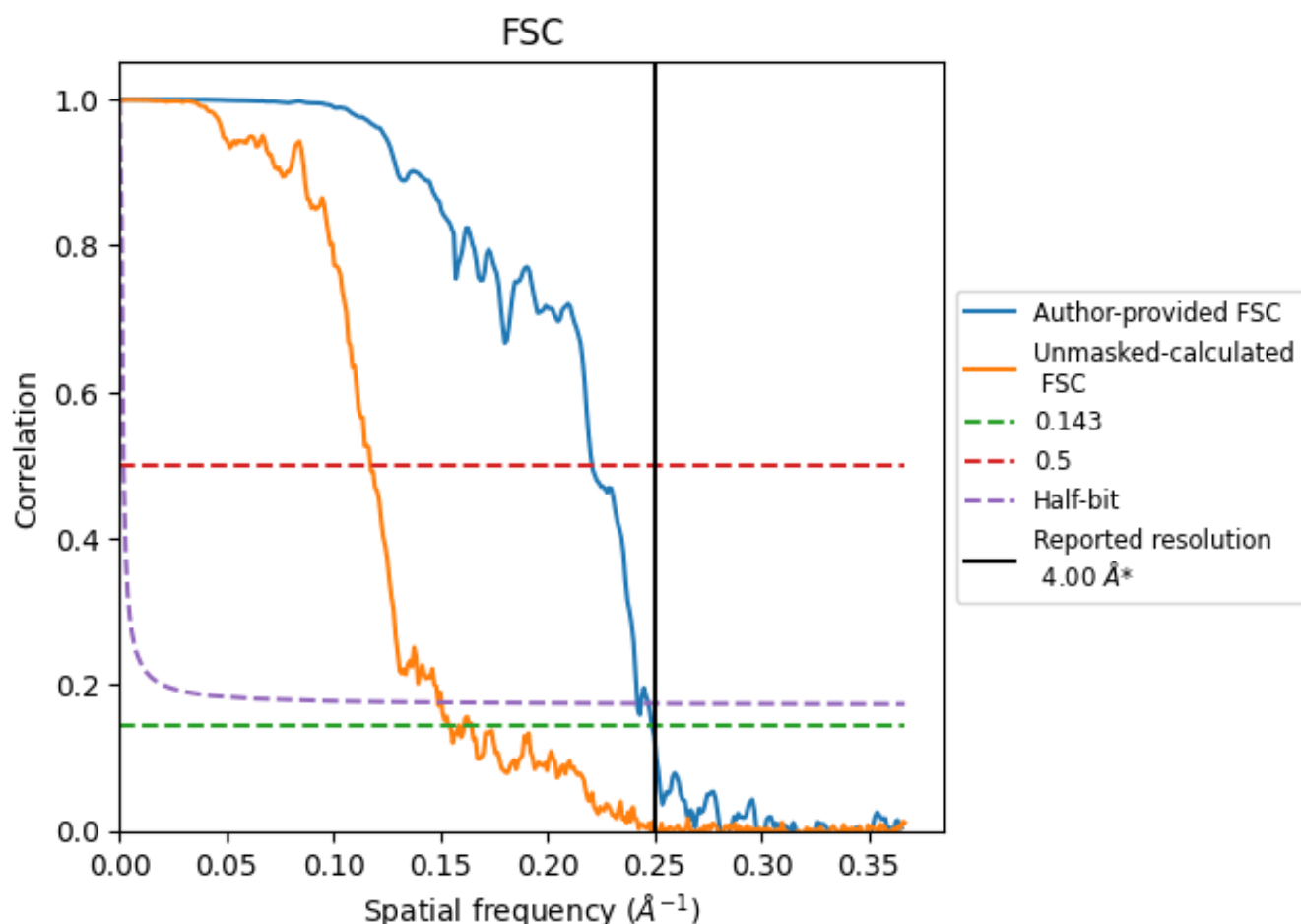


*Reported resolution corresponds to spatial frequency of 0.250 \AA^{-1}

8 Fourier-Shell correlation [i](#)

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

8.1 FSC [i](#)



*Reported resolution corresponds to spatial frequency of 0.250 Å⁻¹

8.2 Resolution estimates [i](#)

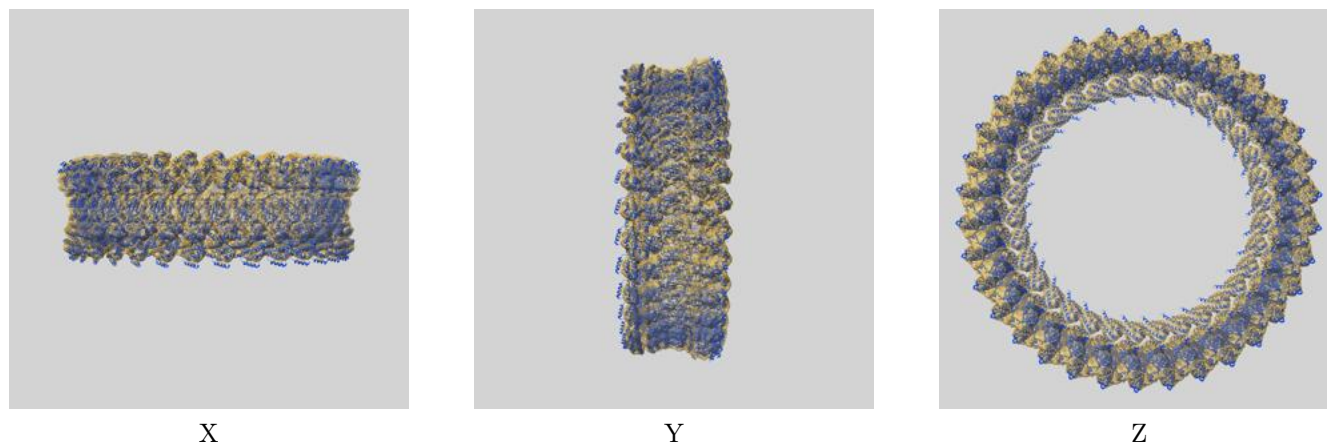
Resolution estimate (Å)	Estimation criterion (FSC cut-off)		
	0.143	0.5	Half-bit
Reported by author	4.00	-	-
Author-provided FSC curve	4.02	4.53	4.13
Unmasked-calculated*	6.50	8.53	6.72

*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 6.50 differs from the reported value 4.0 by more than 10 %

9 Map-model fit [i](#)

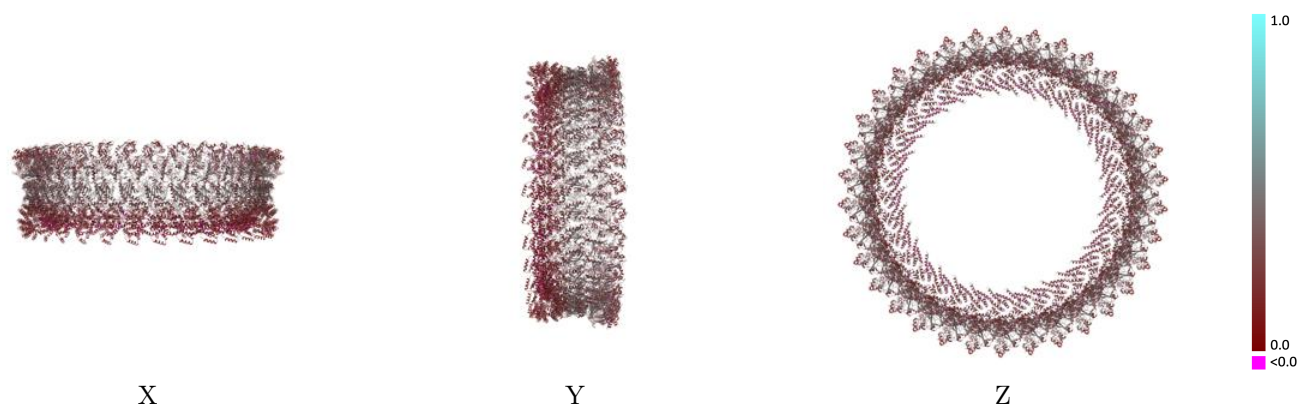
This section contains information regarding the fit between EMDB map EMD-41100 and PDB model 8T8O. Per-residue inclusion information can be found in section [3](#) on page [22](#).

9.1 Map-model overlay [i](#)



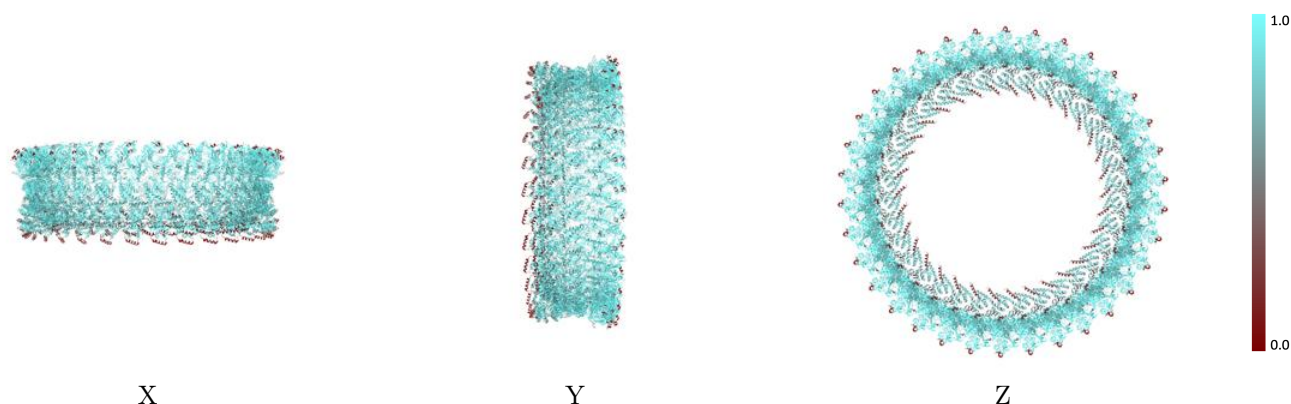
The images above show the 3D surface view of the map at the recommended contour level 0.147 at 50% transparency in yellow overlaid with a ribbon representation of the model coloured in blue. These images allow for the visual assessment of the quality of fit between the atomic model and the map.

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



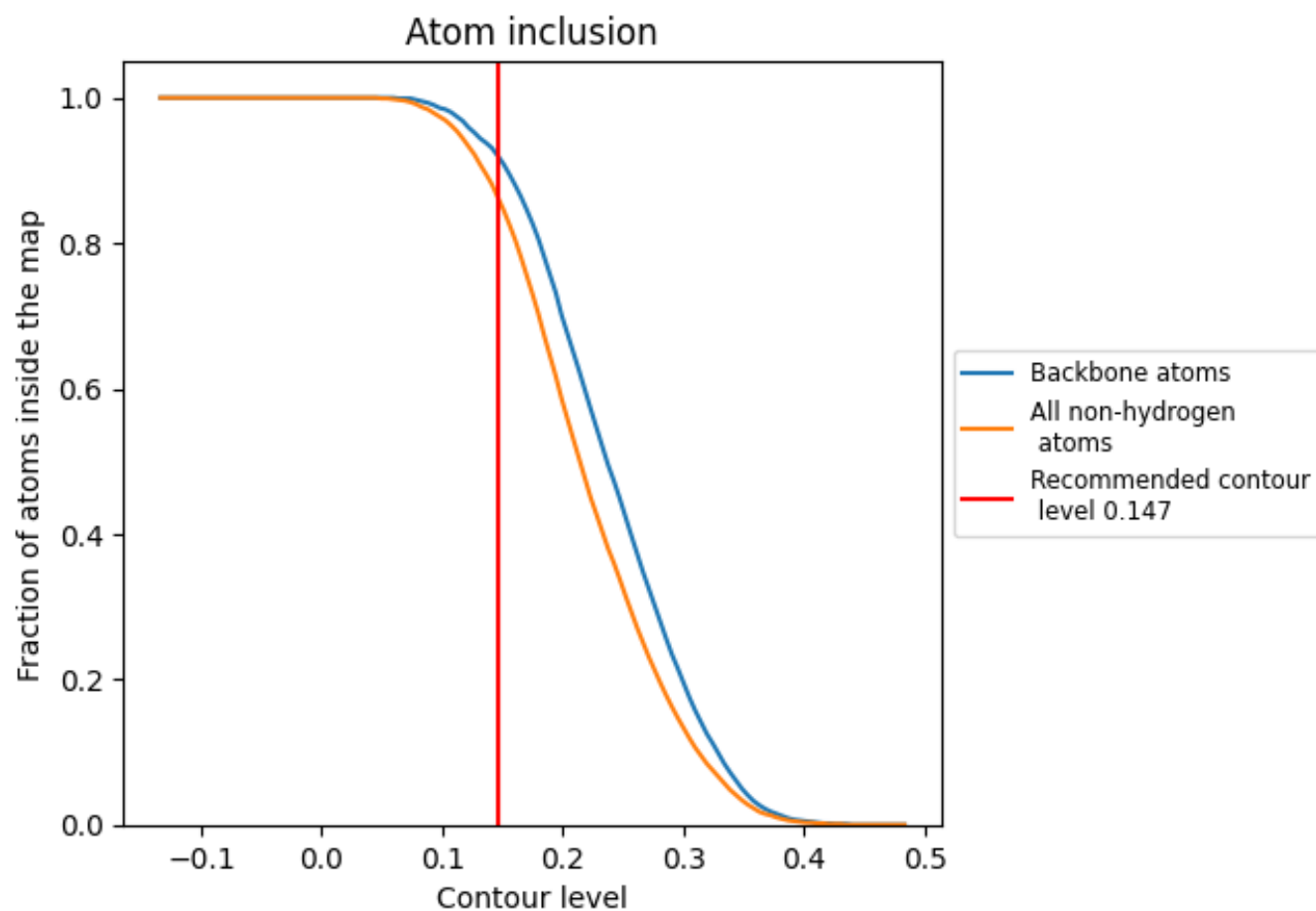
The images above show the model with each residue coloured according its Q-score. This shows their resolvability in the map with higher Q-score values reflecting better resolvability. Please note: Q-score is calculating the resolvability of atoms, and thus high values are only expected at resolutions 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 (0.147).







































































9.4 Atom inclusion [i](#)



At the recommended contour level, 92% of all backbone atoms, 86% of all non-hydrogen atoms, are inside the map.

9.5 Map-model fit summary ⓘ





















































































The table lists the average atom inclusion at the recommended contour level (0.147) and Q-score for the entire model and for each chain.

Chain	Atom inclusion	Q-score
All	 0.8600	 0.2720
A	 0.4960	 0.1610
AA	 0.8160	 0.1970
AB	 0.9230	 0.3140
AC	 0.9190	 0.3650
AD	 0.8140	 0.1950
AE	 0.9120	 0.3140
AF	 0.9050	 0.3600
AG	 0.8120	 0.1960
B	 0.8120	 0.1970
BA	 0.9220	 0.3430
BB	 0.9510	 0.3260
BC	 0.4790	 0.1520
BD	 0.9160	 0.3400
BE	 0.9490	 0.3260
BF	 0.4830	 0.1540
BG	 0.9190	 0.3420
C	 0.9250	 0.3410
CA	 0.9230	 0.3130
CB	 0.9130	 0.3630
CC	 0.8150	 0.1950
CD	 0.9190	 0.3140
CE	 0.9000	 0.3580
CF	 0.8090	 0.1950
CG	 0.9170	 0.3100
D	 0.9250	 0.3140
DA	 0.9560	 0.3220
DB	 0.4910	 0.1550
DC	 0.9180	 0.3380
DD	 0.9490	 0.3270
DE	 0.4830	 0.1530
DF	 0.9200	 0.3450
DG	 0.9540	 0.3200
E	 0.9590	 0.3230
EA	 0.9050	 0.3640























































































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Chain	Atom inclusion	Q-score
EB	 0.8150	 0.1970
EC	 0.9190	 0.3170
ED	 0.9080	 0.3630
EE	 0.8150	 0.1950
EF	 0.9170	 0.3130
EG	 0.9020	 0.3570
F	 0.5000	 0.1610
FA	 0.4960	 0.1560
FB	 0.9240	 0.3430
FC	 0.9460	 0.3260
FD	 0.4790	 0.1490
FE	 0.9150	 0.3410
FF	 0.9510	 0.3230
FG	 0.4910	 0.1630
G	 0.8150	 0.1970
GA	 0.8190	 0.1970
GB	 0.9210	 0.3140
GC	 0.9080	 0.3640
GD	 0.8120	 0.1930
GE	 0.9140	 0.3130
GF	 0.9020	 0.3620
GG	 0.8100	 0.1960
H	 0.9050	 0.3600
HA	 0.9230	 0.3430
HB	 0.9540	 0.3260
HC	 0.4700	 0.1550
HD	 0.9150	 0.3400
HE	 0.9460	 0.3250
HF	 0.4830	 0.1540
HG	 0.9190	 0.3410
I	 0.5000	 0.1600
IA	 0.9230	 0.3130
IB	 0.9130	 0.3640
IC	 0.8160	 0.1970
ID	 0.9140	 0.3150
IE	 0.9000	 0.3570
IF	 0.8120	 0.1960
IG	 0.9170	 0.3110
J	 0.8130	 0.1980
JA	 0.9590	 0.3260
JB	 0.4910	 0.1520
JC	 0.9180	 0.3390





















































































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Chain	Atom inclusion	Q-score
JD	 0.9490	 0.3270
JE	 0.4790	 0.1530
JF	 0.9190	 0.3430
JG	 0.9560	 0.3180
K	 0.9230	 0.3410
KA	 0.9050	 0.3590
KB	 0.8140	 0.1980
KC	 0.9190	 0.3140
KD	 0.9080	 0.3620
KE	 0.8130	 0.1940
KF	 0.9170	 0.3170
KG	 0.9050	 0.3550
L	 0.9250	 0.3140
LA	 0.4960	 0.1550
LB	 0.9220	 0.3420
LC	 0.9430	 0.3200
LD	 0.4790	 0.1540
LE	 0.9170	 0.3400
LF	 0.9540	 0.3230
LG	 0.4960	 0.1610
M	 0.9220	 0.3390
MA	 0.8160	 0.1960
MB	 0.9250	 0.3140
MC	 0.9110	 0.3610
MD	 0.8120	 0.1940
ME	 0.9140	 0.3130
MF	 0.9050	 0.3630
MG	 0.8130	 0.1970
N	 0.9190	 0.3110
NA	 0.9220	 0.3420
NB	 0.9510	 0.3260
NC	 0.4700	 0.1550
ND	 0.9150	 0.3400
NE	 0.9510	 0.3270
NF	 0.4870	 0.1580
NG	 0.9210	 0.3440
O	 0.9540	 0.3220
OA	 0.9230	 0.3130
OB	 0.9160	 0.3560
OC	 0.8100	 0.1950
OD	 0.9140	 0.3160
OE	 0.9020	 0.3610





















































































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Chain	Atom inclusion	Q-score
OF	 0.8120	 0.1980
OG	 0.9170	 0.3120
P	 0.9560	 0.3240
PA	 0.9560	 0.3250
PB	 0.4830	 0.1520
PC	 0.9180	 0.3410
PD	 0.9460	 0.3260
PE	 0.4830	 0.1530
PF	 0.9220	 0.3440
PG	 0.9540	 0.3230
Q	 0.9050	 0.3580
QA	 0.9050	 0.3620
QB	 0.8100	 0.1970
QC	 0.9140	 0.3150
QD	 0.9050	 0.3540
QE	 0.8090	 0.1950
QF	 0.9170	 0.3150
QG	 0.9050	 0.3570
R	 0.9080	 0.3500
RA	 0.4960	 0.1560
RB	 0.9220	 0.3410
RC	 0.9460	 0.3250
RD	 0.4660	 0.1530
RE	 0.9200	 0.3420
RF	 0.9510	 0.3220
RG	 0.5000	 0.1610
S	 0.5000	 0.1560
SA	 0.8160	 0.1960
SB	 0.9230	 0.3160
SC	 0.9130	 0.3610
SD	 0.8090	 0.1950
SE	 0.9170	 0.3140
SF	 0.9080	 0.3560
SG	 0.8150	 0.1950
T	 0.8120	 0.1970
TA	 0.9200	 0.3430
TB	 0.9460	 0.3250
TC	 0.4740	 0.1550
TD	 0.9150	 0.3400
TE	 0.9540	 0.3250
TF	 0.4870	 0.1620
TG	 0.9190	 0.3410





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Chain	Atom inclusion	Q-score
UA	 0.9230	 0.3140
UB	 0.9110	 0.3620
UC	 0.8100	 0.1940
UD	 0.9140	 0.3170
UE	 0.9050	 0.3610
UF	 0.8100	 0.1950
UG	 0.9230	 0.3110
V	 0.9220	 0.3420
VA	 0.9560	 0.3260
VB	 0.4790	 0.1530
VC	 0.9140	 0.3400
VD	 0.9490	 0.3230
VE	 0.4830	 0.1530
VF	 0.9210	 0.3410
VG	 0.9560	 0.3270
W	 0.9230	 0.3110
WA	 0.9130	 0.3630
WB	 0.8110	 0.1960
WC	 0.9190	 0.3140
WD	 0.9000	 0.3510
WE	 0.8060	 0.1950
WF	 0.9140	 0.3100
WG	 0.9050	 0.3550
X	 0.9590	 0.3210
XA	 0.4960	 0.1560
XB	 0.9230	 0.3390
XC	 0.9460	 0.3280
XD	 0.4700	 0.1550
XE	 0.9210	 0.3420
XF	 0.9560	 0.3240
Y	 0.9050	 0.3620
YA	 0.8120	 0.1980
YB	 0.9210	 0.3120
YC	 0.9130	 0.3510
YD	 0.8140	 0.1970
YE	 0.9190	 0.3130
YF	 0.9050	 0.3560
Z	 0.5000	 0.1560
ZA	 0.9220	 0.3430
ZB	 0.9490	 0.3250
ZC	 0.4660	 0.1520
ZD	 0.9170	 0.3410

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Chain	Atom inclusion	Q-score
ZE	 0.9510	 0.3260
ZF	 0.4910	 0.1590