



## Full wwPDB EM Validation Report ⓘ

May 5, 2026 – 11:28 pm BST

PDB ID : 9QW9 / pdb\_00009qw9  
EMDB ID : EMD-53415  
Title : Human vault protein - primed conformation  
Authors : Lapenta, F.; Marechal, N.; Durand, A.; Aupic, J.; Cassetta, A.  
Deposited on : 2025-04-14  
Resolution : 3.09 Å(reported)

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

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

A user guide is available at

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

with specific help available everywhere you see the ⓘ symbol.

The types of validation reports are described at

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

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

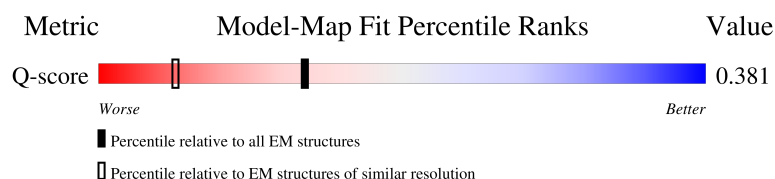
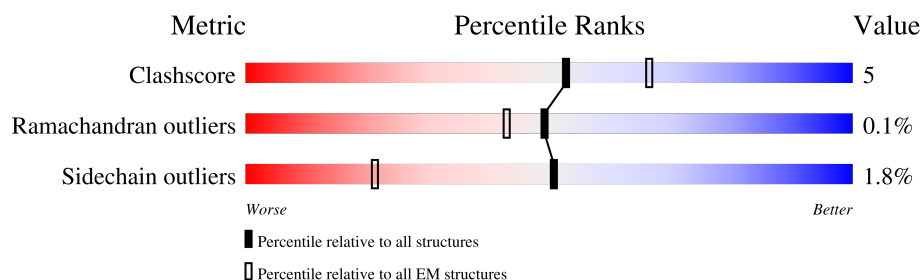
EMDB validation analysis : 0.0.1.dev132  
MolProbity : 4-5-2 with Phenix2.0  
Percentile statistics : 20250101.v01 (using entries in the PDB archive January 1st 2025)  
EM percentile statistics : 202505.v01 (Using data in the EMDB archive up until May 2025)  
MapQ : 1.9.13  
Ideal geometry (proteins)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

# 1 Overall quality at a glance i

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

The reported resolution of this entry is 3.09 Å.

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









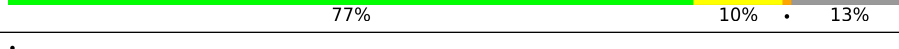
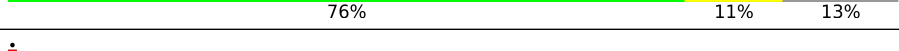
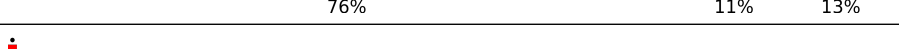
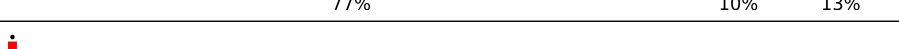
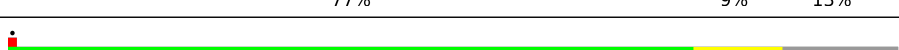

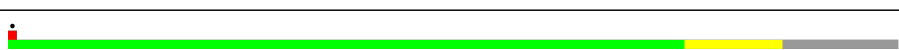

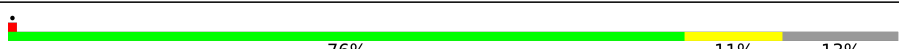





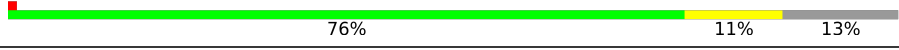
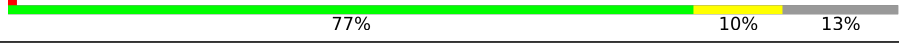



Metric	Whole archive (#Entries)	EM structures (#Entries)	Similar EM resolution (#Entries, resolution range(Å))
Clashscore	229148	23984	-
Ramachandran outliers	224038	23583	-
Sidechain outliers	223484	23102	-
Q-score	-	25397	14003 ( 2.59 - 3.59 )

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

Mol	Chain	Length	Quality of chain
1	A	893	
1	AA	893	
1	AB	893	
1	AC	893	







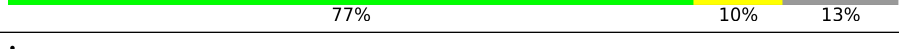
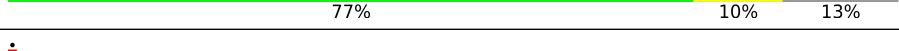
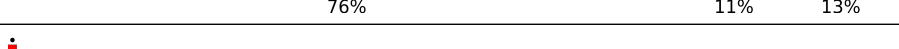
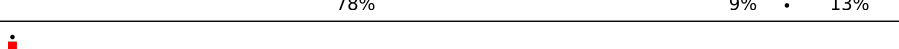
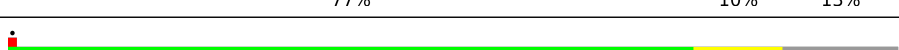

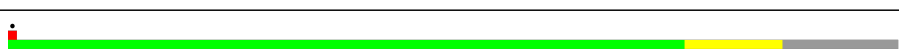

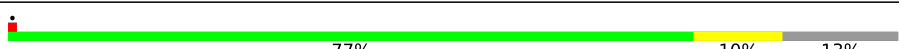





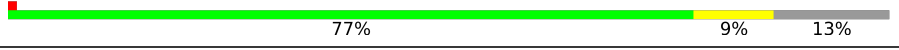
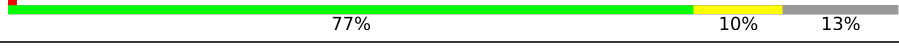



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Mol	Chain	Length	Quality of chain
1	B	893	
1	BA	893	
1	BB	893	
1	C	893	
1	CA	893	
1	CB	893	
1	D	893	
1	DA	893	
1	DB	893	
1	E	893	
1	EA	893	
1	EB	893	
1	F	893	
1	FA	893	
1	FB	893	
1	G	893	
1	GA	893	
1	GB	893	
1	H	893	
1	HA	893	
1	HB	893	
1	I	893	
1	IA	893	
1	IB	893	
1	J	893	



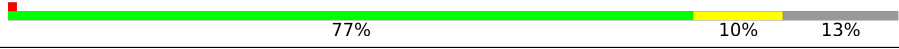




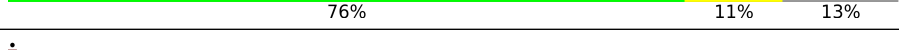
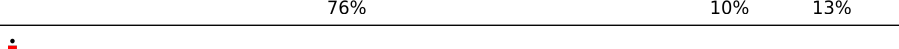
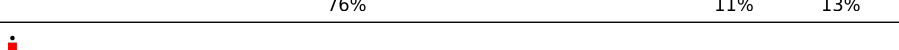

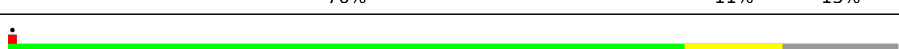


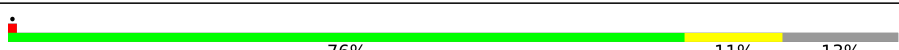





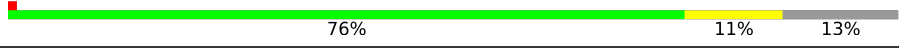



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Mol	Chain	Length	Quality of chain
1	JA	893	
1	JB	893	
1	K	893	
1	KA	893	
1	KB	893	
1	L	893	
1	LA	893	
1	LB	893	
1	M	893	
1	MA	893	
1	MB	893	
1	N	893	
1	NA	893	
1	NB	893	
1	O	893	
1	OA	893	
1	OB	893	
1	P	893	
1	PA	893	
1	PB	893	
1	Q	893	
1	QA	893	
1	QB	893	
1	R	893	
1	RA	893	

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Mol	Chain	Length	Quality of chain
1	RB	893	
1	S	893	
1	SA	893	
1	SB	893	
1	T	893	
1	TA	893	
1	TB	893	
1	UA	893	
1	UB	893	
1	V	893	
1	VA	893	
1	VB	893	
1	W	893	
1	WA	893	
1	WB	893	
1	X	893	
1	XA	893	
1	XB	893	
1	Y	893	
1	YA	893	
1	YB	893	
1	Z	893	
1	ZA	893	
1	ZB	893	

## 2 Entry composition [i](#)

There is only 1 type of molecule in this entry. The entry contains 966108 atoms, of which 483990 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 Major vault protein.

Mol	Chain	Residues	Atoms						AltConf	Trace
1	A	779	Total 12386	C 3890	H 6205	N 1105	O 1176	S 10	0	0
1	AA	779	Total 12386	C 3890	H 6205	N 1105	O 1176	S 10	0	0
1	AB	779	Total 12386	C 3890	H 6205	N 1105	O 1176	S 10	0	0
1	AC	779	Total 12386	C 3890	H 6205	N 1105	O 1176	S 10	0	0
1	B	779	Total 12386	C 3890	H 6205	N 1105	O 1176	S 10	0	0
1	BA	779	Total 12386	C 3890	H 6205	N 1105	O 1176	S 10	0	0
1	BB	779	Total 12386	C 3890	H 6205	N 1105	O 1176	S 10	0	0
1	C	779	Total 12386	C 3890	H 6205	N 1105	O 1176	S 10	0	0
1	CA	779	Total 12386	C 3890	H 6205	N 1105	O 1176	S 10	0	0
1	CB	779	Total 12386	C 3890	H 6205	N 1105	O 1176	S 10	0	0
1	D	779	Total 12386	C 3890	H 6205	N 1105	O 1176	S 10	0	0
1	DA	779	Total 12386	C 3890	H 6205	N 1105	O 1176	S 10	0	0
1	DB	779	Total 12386	C 3890	H 6205	N 1105	O 1176	S 10	0	0
1	E	779	Total 12386	C 3890	H 6205	N 1105	O 1176	S 10	0	0
1	EA	779	Total 12386	C 3890	H 6205	N 1105	O 1176	S 10	0	0
1	EB	779	Total 12386	C 3890	H 6205	N 1105	O 1176	S 10	0	0
1	F	779	Total 12386	C 3890	H 6205	N 1105	O 1176	S 10	0	0

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Mol	Chain	Residues	Atoms						AltConf	Trace
1	FA	779	Total 12386	C 3890	H 6205	N 1105	O 1176	S 10	0	0
1	FB	779	Total 12386	C 3890	H 6205	N 1105	O 1176	S 10	0	0
1	G	779	Total 12386	C 3890	H 6205	N 1105	O 1176	S 10	0	0
1	GA	779	Total 12386	C 3890	H 6205	N 1105	O 1176	S 10	0	0
1	GB	779	Total 12386	C 3890	H 6205	N 1105	O 1176	S 10	0	0
1	H	779	Total 12386	C 3890	H 6205	N 1105	O 1176	S 10	0	0
1	HA	779	Total 12386	C 3890	H 6205	N 1105	O 1176	S 10	0	0
1	HB	779	Total 12386	C 3890	H 6205	N 1105	O 1176	S 10	0	0
1	I	779	Total 12386	C 3890	H 6205	N 1105	O 1176	S 10	0	0
1	IA	779	Total 12386	C 3890	H 6205	N 1105	O 1176	S 10	0	0
1	IB	779	Total 12386	C 3890	H 6205	N 1105	O 1176	S 10	0	0
1	J	779	Total 12386	C 3890	H 6205	N 1105	O 1176	S 10	0	0
1	JA	779	Total 12386	C 3890	H 6205	N 1105	O 1176	S 10	0	0
1	JB	779	Total 12386	C 3890	H 6205	N 1105	O 1176	S 10	0	0
1	K	779	Total 12386	C 3890	H 6205	N 1105	O 1176	S 10	0	0
1	KA	779	Total 12386	C 3890	H 6205	N 1105	O 1176	S 10	0	0
1	KB	779	Total 12386	C 3890	H 6205	N 1105	O 1176	S 10	0	0
1	L	779	Total 12386	C 3890	H 6205	N 1105	O 1176	S 10	0	0
1	LA	779	Total 12386	C 3890	H 6205	N 1105	O 1176	S 10	0	0
1	LB	779	Total 12386	C 3890	H 6205	N 1105	O 1176	S 10	0	0
1	M	779	Total 12386	C 3890	H 6205	N 1105	O 1176	S 10	0	0

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Mol	Chain	Residues	Atoms						AltConf	Trace
1	MA	779	Total 12386	C 3890	H 6205	N 1105	O 1176	S 10	0	0
1	MB	779	Total 12386	C 3890	H 6205	N 1105	O 1176	S 10	0	0
1	N	779	Total 12386	C 3890	H 6205	N 1105	O 1176	S 10	0	0
1	NA	779	Total 12386	C 3890	H 6205	N 1105	O 1176	S 10	0	0
1	NB	779	Total 12386	C 3890	H 6205	N 1105	O 1176	S 10	0	0
1	O	779	Total 12386	C 3890	H 6205	N 1105	O 1176	S 10	0	0
1	OA	779	Total 12386	C 3890	H 6205	N 1105	O 1176	S 10	0	0
1	OB	779	Total 12386	C 3890	H 6205	N 1105	O 1176	S 10	0	0
1	P	779	Total 12386	C 3890	H 6205	N 1105	O 1176	S 10	0	0
1	PA	779	Total 12386	C 3890	H 6205	N 1105	O 1176	S 10	0	0
1	PB	779	Total 12386	C 3890	H 6205	N 1105	O 1176	S 10	0	0
1	Q	779	Total 12386	C 3890	H 6205	N 1105	O 1176	S 10	0	0
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1	QB	779	Total 12386	C 3890	H 6205	N 1105	O 1176	S 10	0	0
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1	RB	779	Total 12386	C 3890	H 6205	N 1105	O 1176	S 10	0	0
1	S	779	Total 12386	C 3890	H 6205	N 1105	O 1176	S 10	0	0
1	SA	779	Total 12386	C 3890	H 6205	N 1105	O 1176	S 10	0	0
1	SB	779	Total 12386	C 3890	H 6205	N 1105	O 1176	S 10	0	0
1	T	779	Total 12386	C 3890	H 6205	N 1105	O 1176	S 10	0	0

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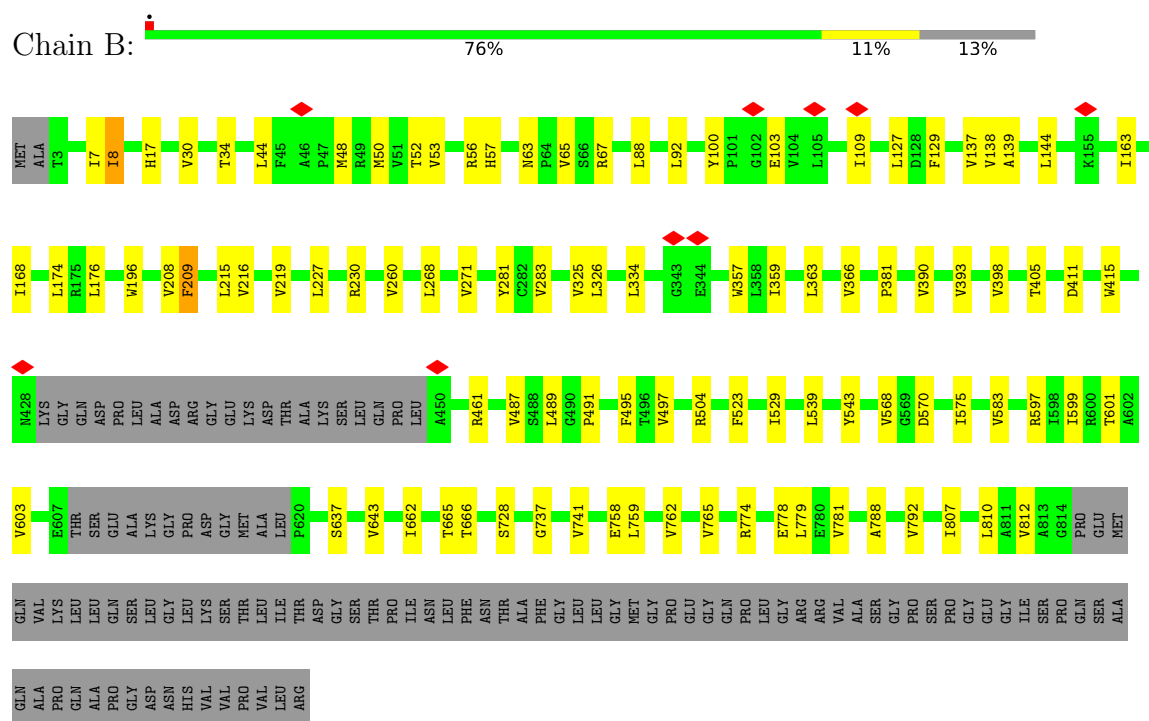
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Mol	Chain	Residues	Atoms						AltConf	Trace
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1	TB	779	Total 12386	C 3890	H 6205	N 1105	O 1176	S 10	0	0
1	UA	779	Total 12386	C 3890	H 6205	N 1105	O 1176	S 10	0	0
1	UB	779	Total 12386	C 3890	H 6205	N 1105	O 1176	S 10	0	0
1	V	779	Total 12386	C 3890	H 6205	N 1105	O 1176	S 10	0	0
1	VA	779	Total 12386	C 3890	H 6205	N 1105	O 1176	S 10	0	0
1	VB	779	Total 12386	C 3890	H 6205	N 1105	O 1176	S 10	0	0
1	W	779	Total 12386	C 3890	H 6205	N 1105	O 1176	S 10	0	0
1	WA	779	Total 12386	C 3890	H 6205	N 1105	O 1176	S 10	0	0
1	WB	779	Total 12386	C 3890	H 6205	N 1105	O 1176	S 10	0	0
1	X	779	Total 12386	C 3890	H 6205	N 1105	O 1176	S 10	0	0
1	XA	779	Total 12386	C 3890	H 6205	N 1105	O 1176	S 10	0	0
1	XB	779	Total 12386	C 3890	H 6205	N 1105	O 1176	S 10	0	0
1	Y	779	Total 12386	C 3890	H 6205	N 1105	O 1176	S 10	0	0
1	YA	779	Total 12386	C 3890	H 6205	N 1105	O 1176	S 10	0	0
1	YB	779	Total 12386	C 3890	H 6205	N 1105	O 1176	S 10	0	0
1	Z	779	Total 12386	C 3890	H 6205	N 1105	O 1176	S 10	0	0
1	ZA	779	Total 12386	C 3890	H 6205	N 1105	O 1176	S 10	0	0
1	ZB	779	Total 12386	C 3890	H 6205	N 1105	O 1176	S 10	0	0

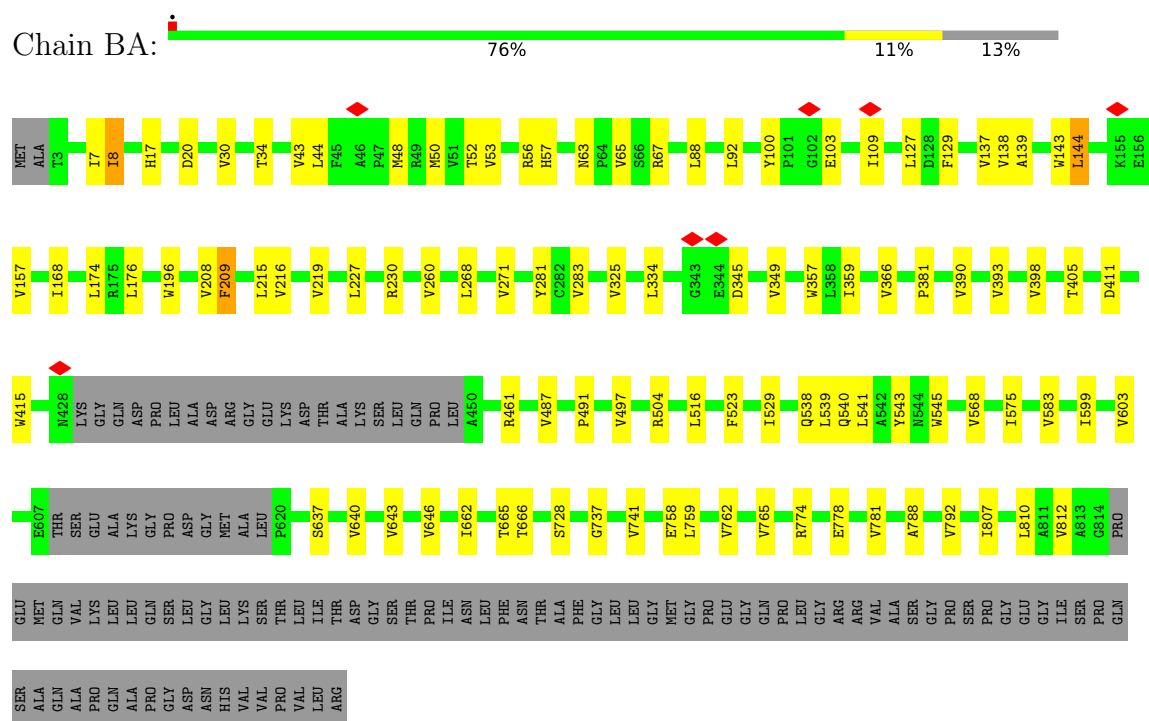




- Molecule 1: Major vault protein

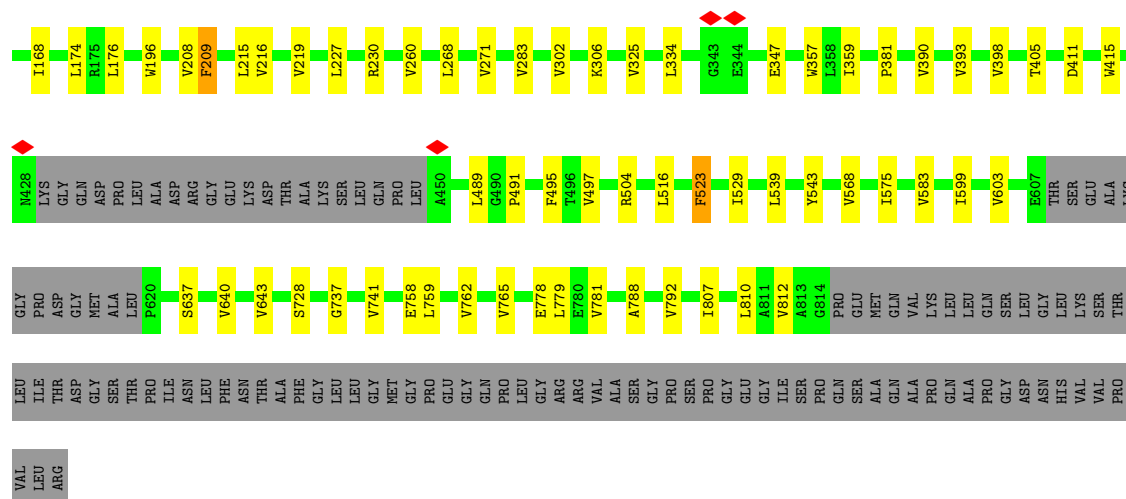


- Molecule 1: Major vault protein

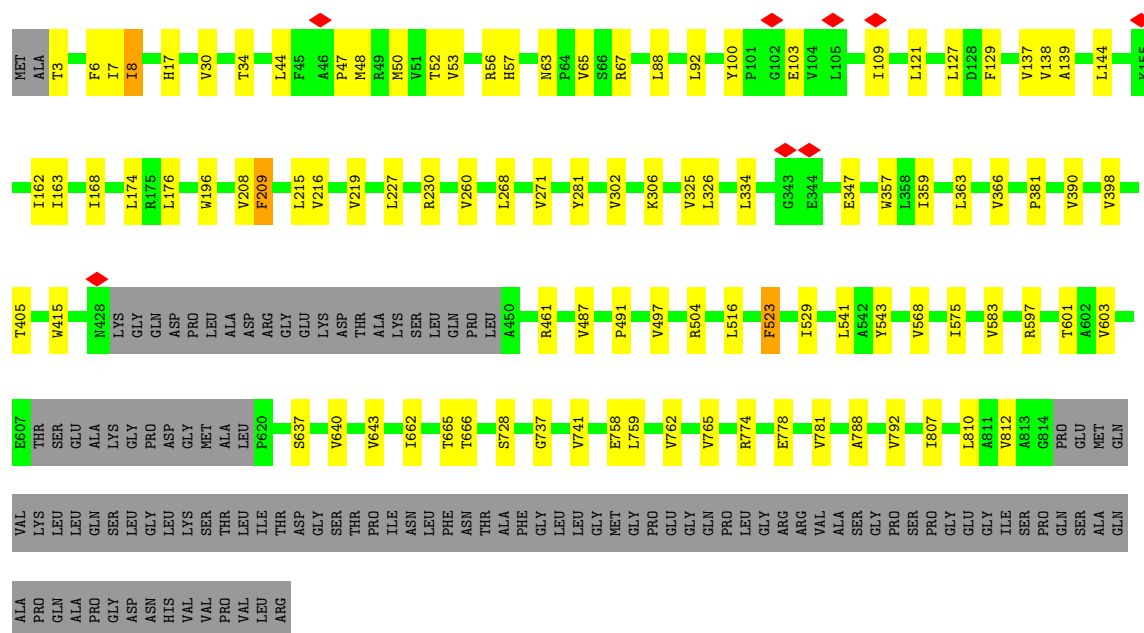
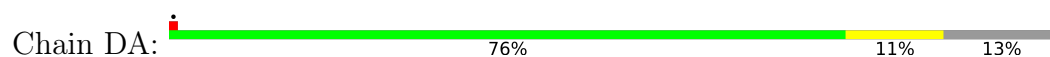




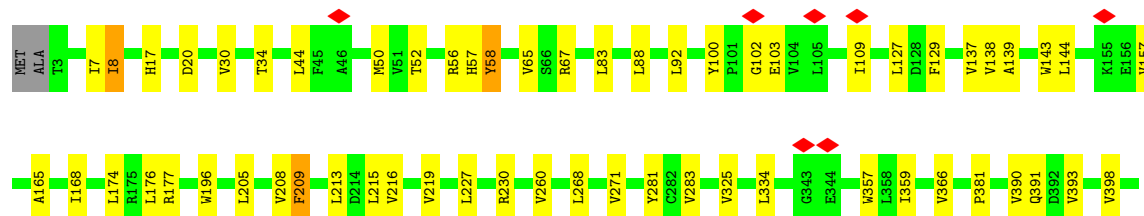
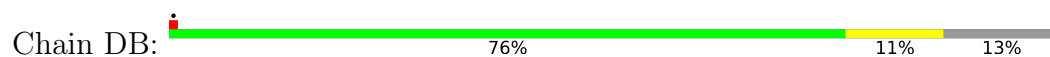


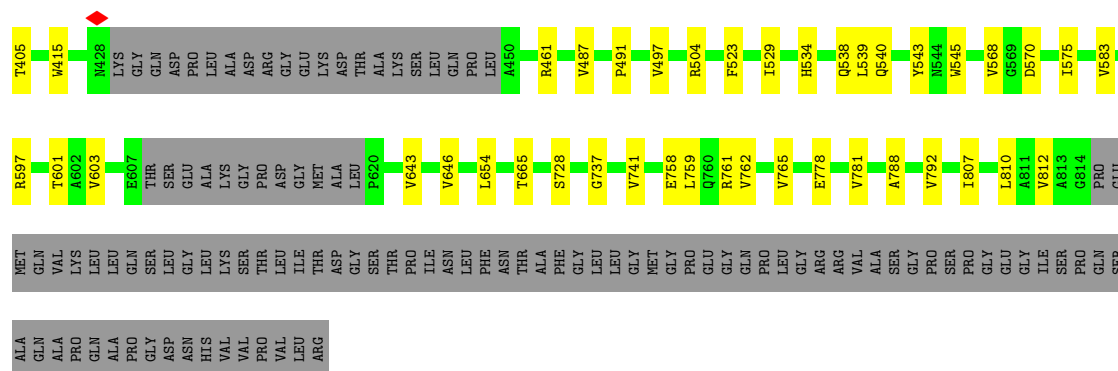


- Molecule 1: Major vault protein

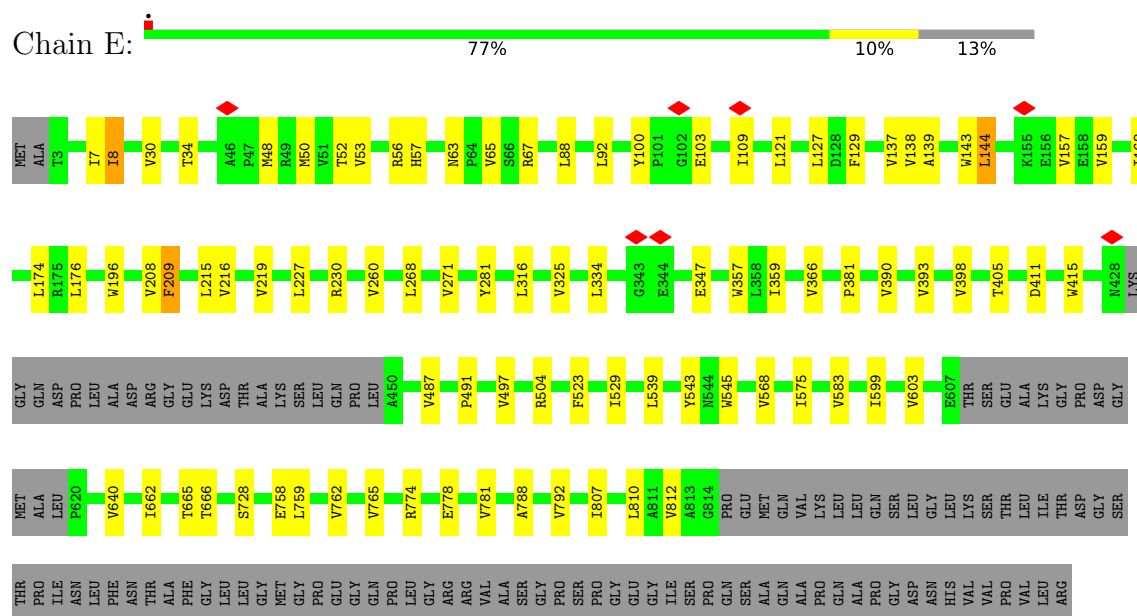


- Molecule 1: Major vault protein

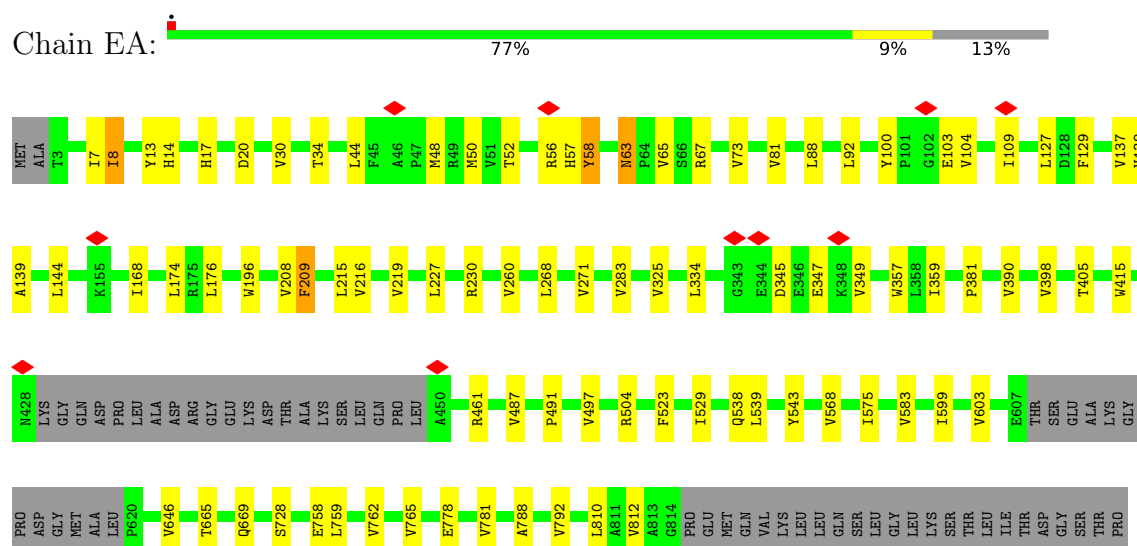




- Molecule 1: Major vault protein



- Molecule 1: Major vault protein

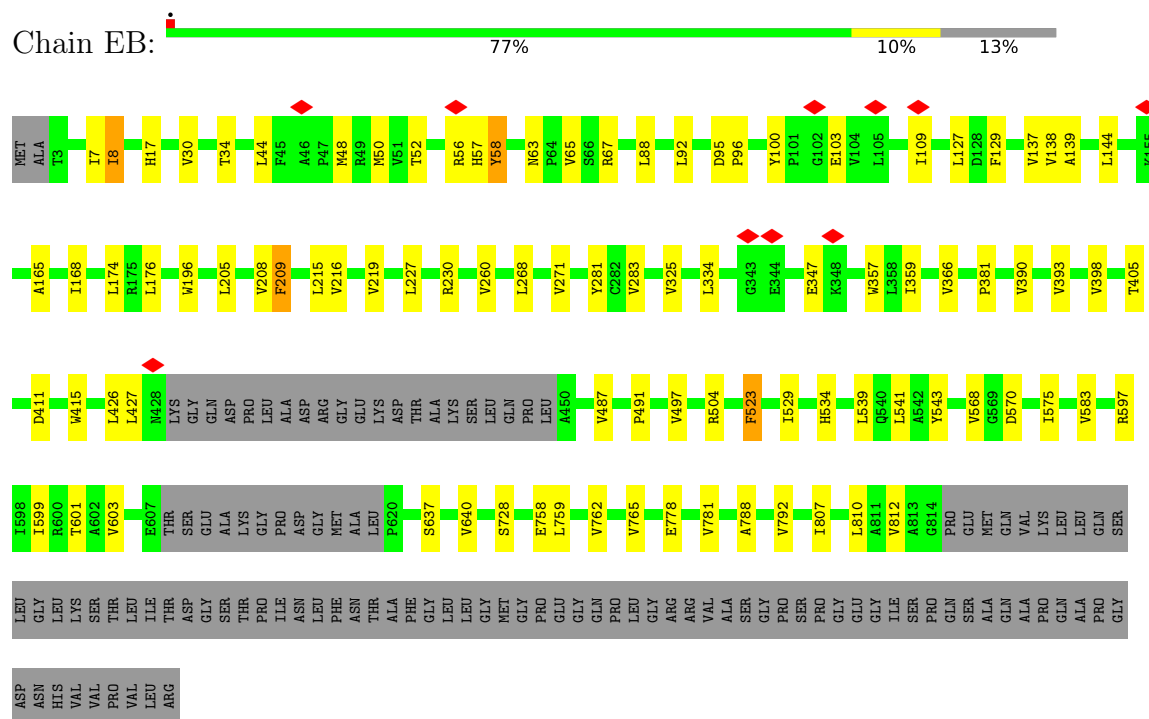




ILE ASN LEU PHE ASN THR ALA PHE GLY LEU LEU GLY MET GLY PRO GLU GLY GLN PRO LEU GLY ARG ARG VAL ALA SER GLY PRO PRO PRO GLY ILE SER PRO GLN SER ALA ALA PRO GLN ALA PRO GLY ASP ASN HIS VAL VAL PRO VAL LEU ARG

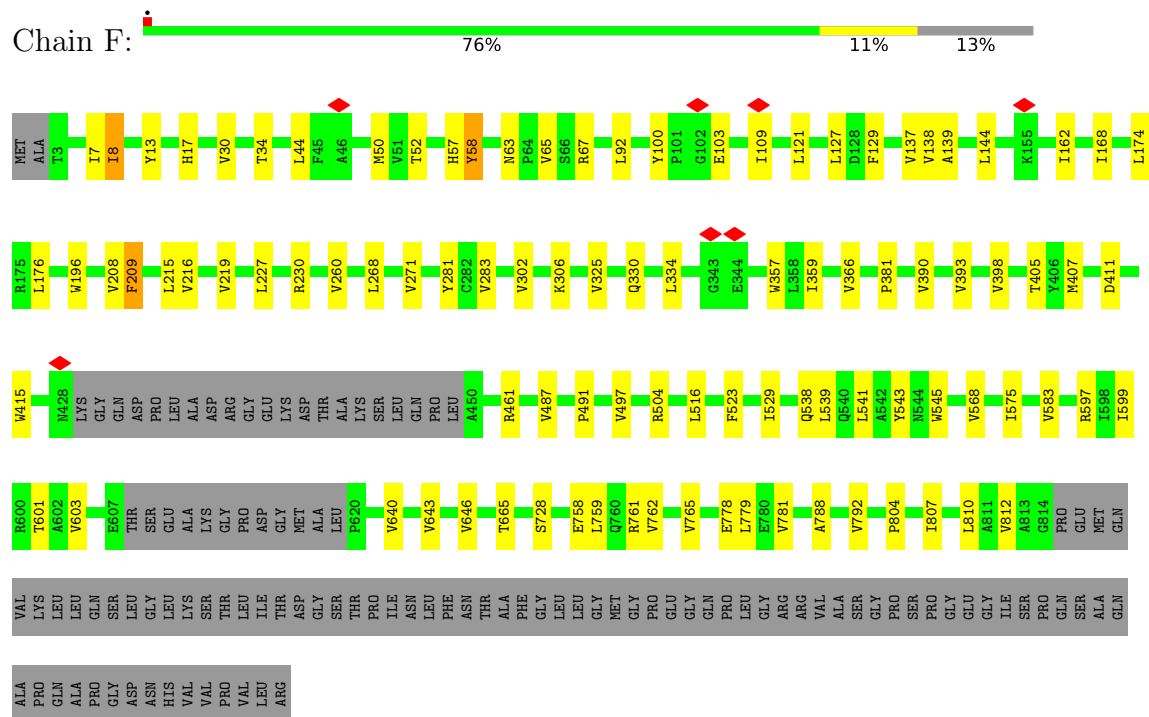
• Molecule 1: Major vault protein

Chain EB:

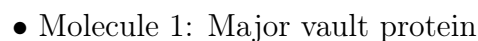


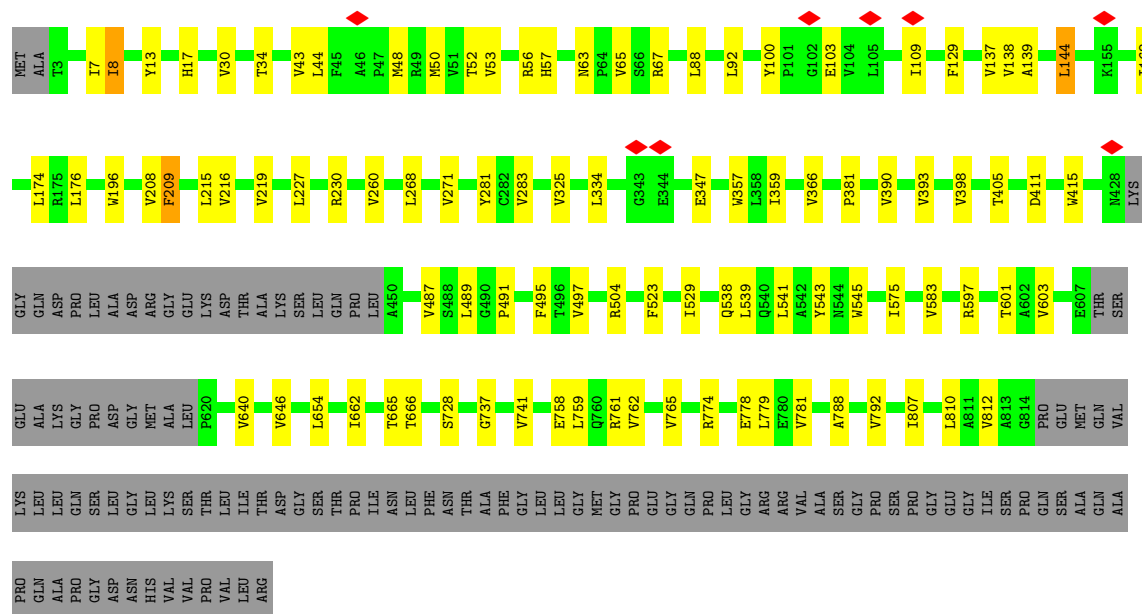
• Molecule 1: Major vault protein

Chain F:

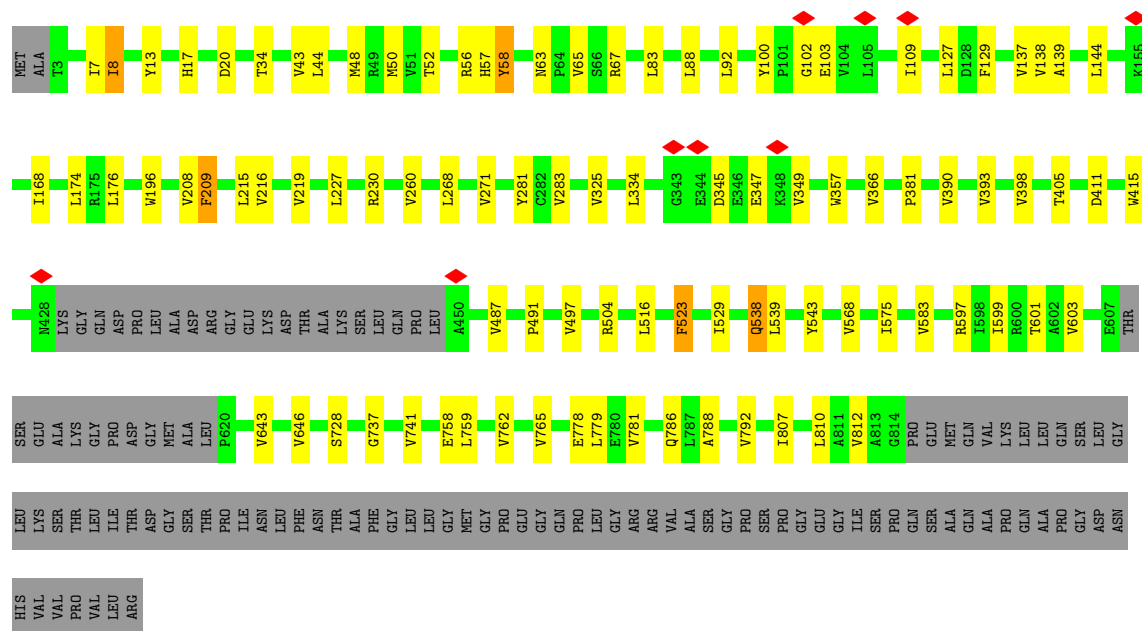
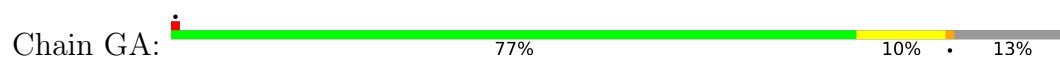


• Molecule 1: Major vault protein

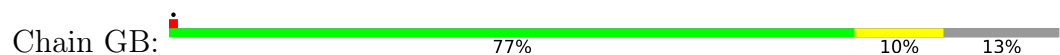


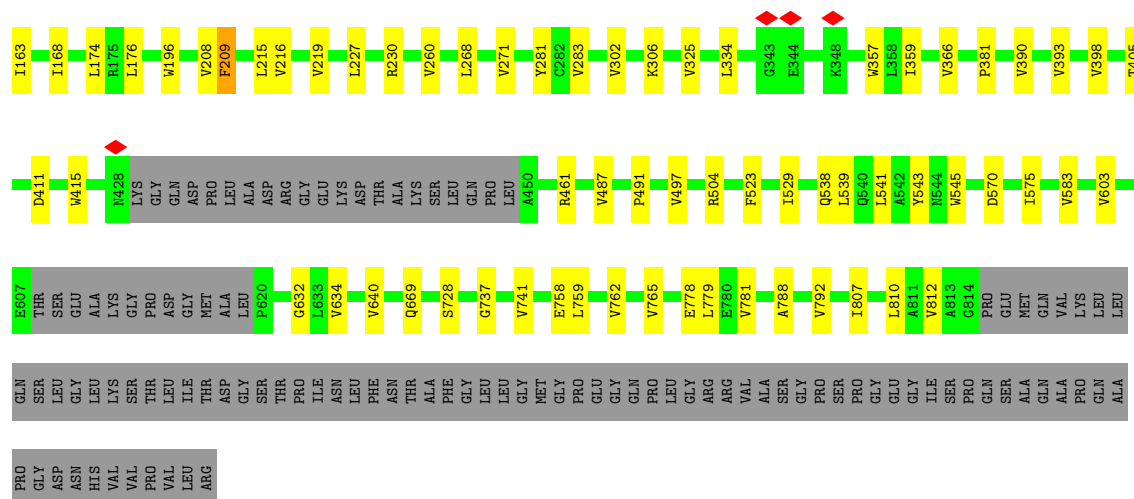


• Molecule 1: Major vault protein



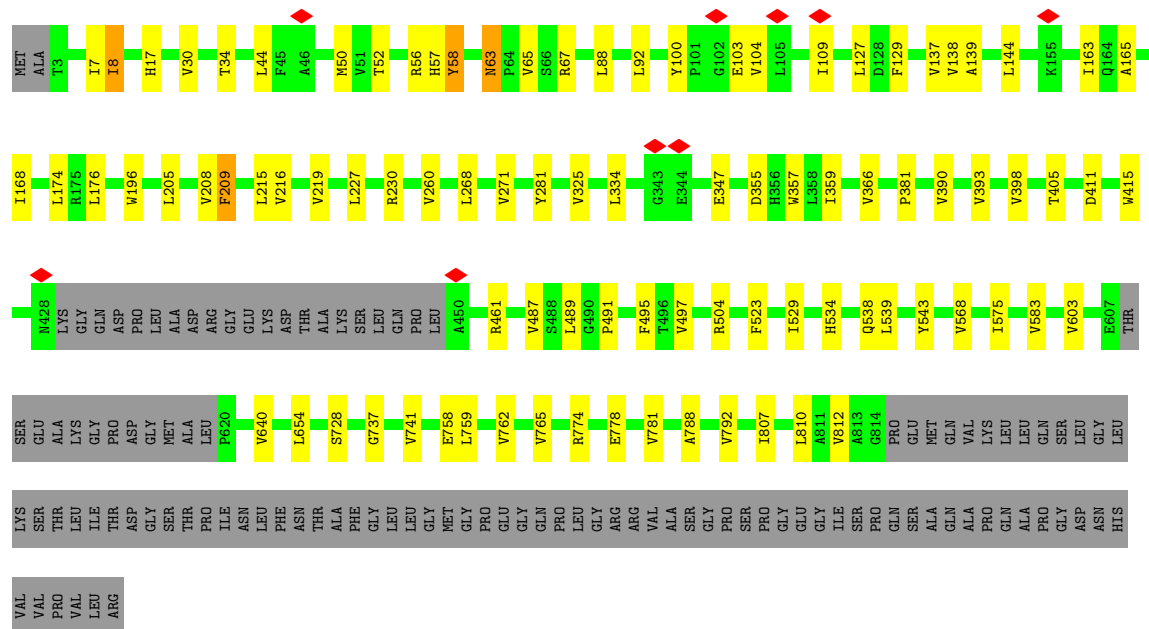
• Molecule 1: Major vault protein

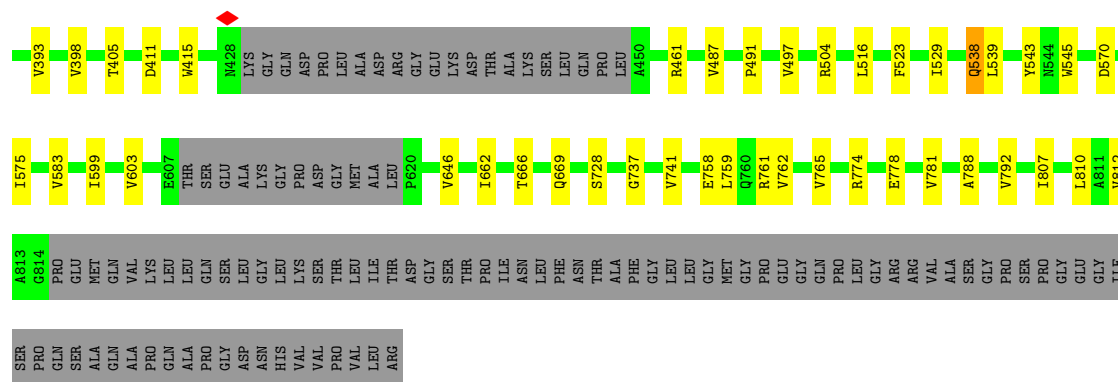


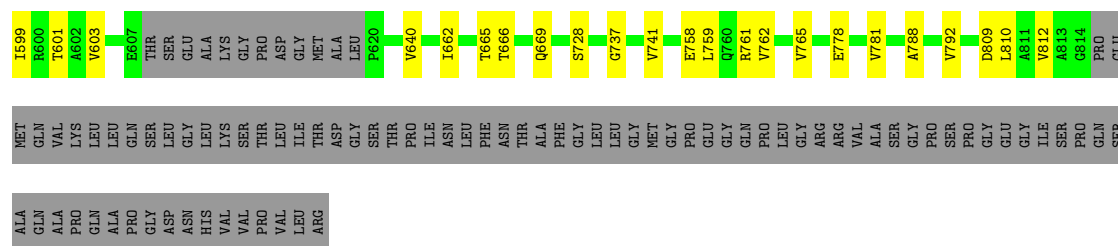


- Molecule 1: Major vault protein

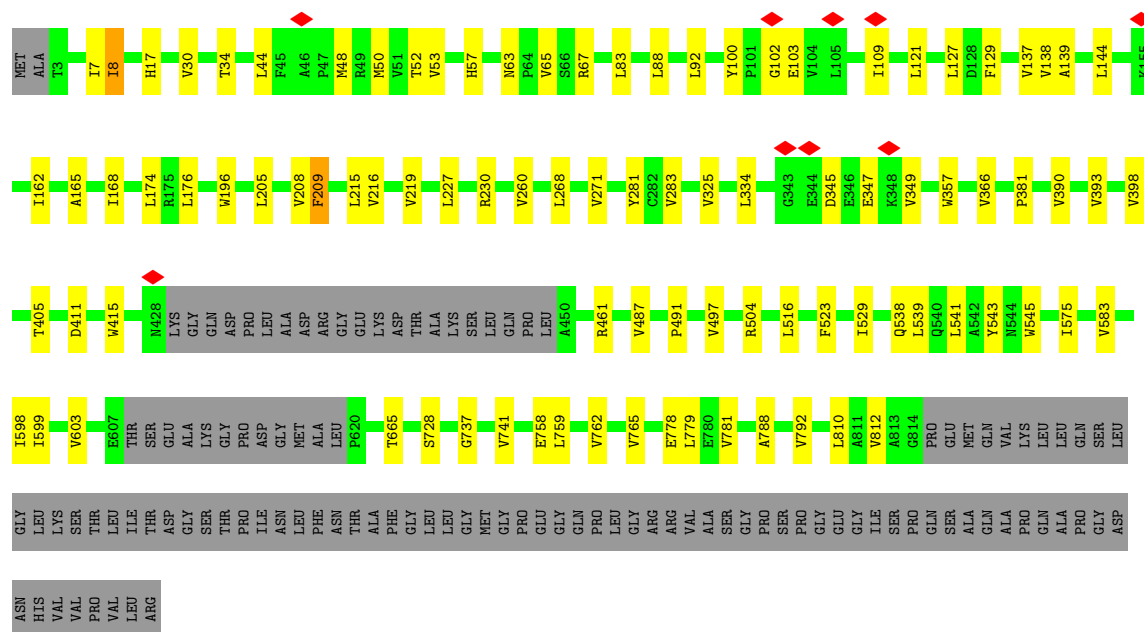
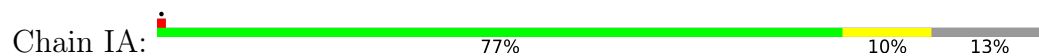
Chain H: 77% 10% 13%



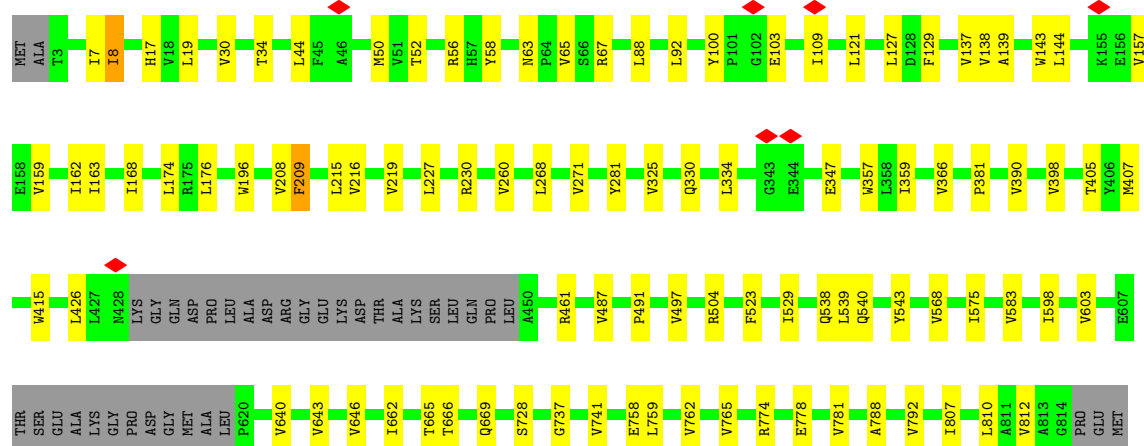
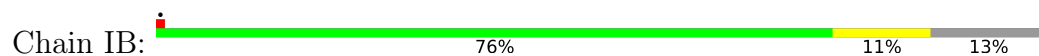


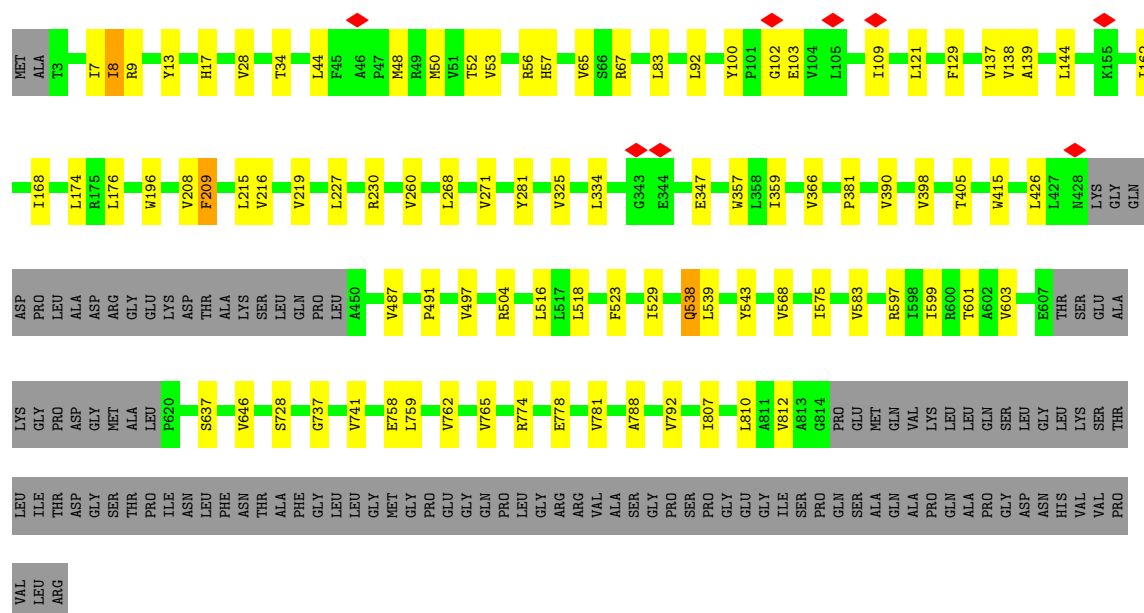


• Molecule 1: Major vault protein

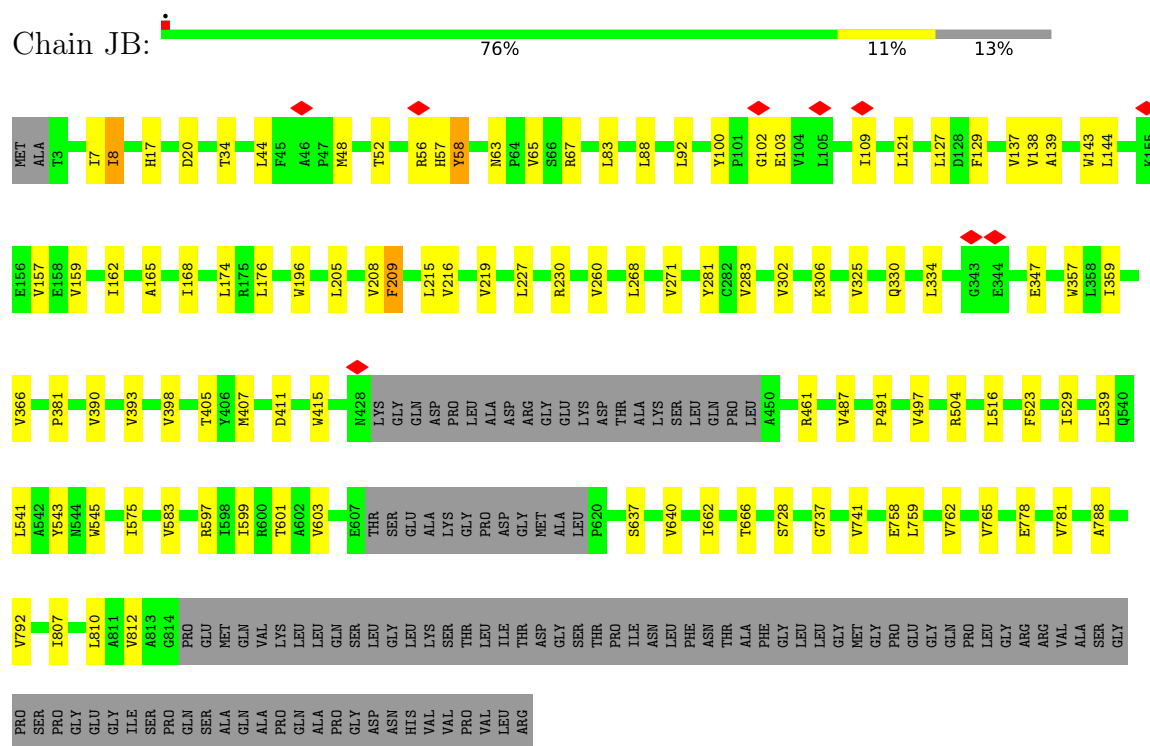


• Molecule 1: Major vault protein

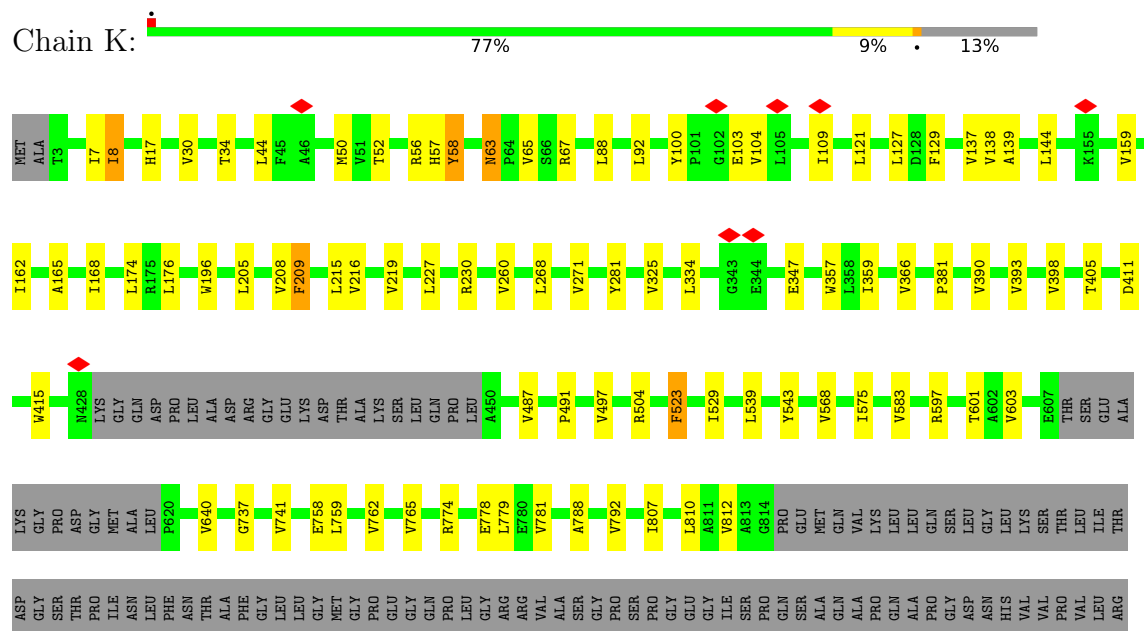




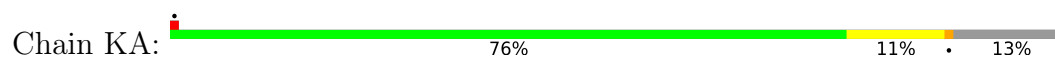
- Molecule 1: Major vault protein



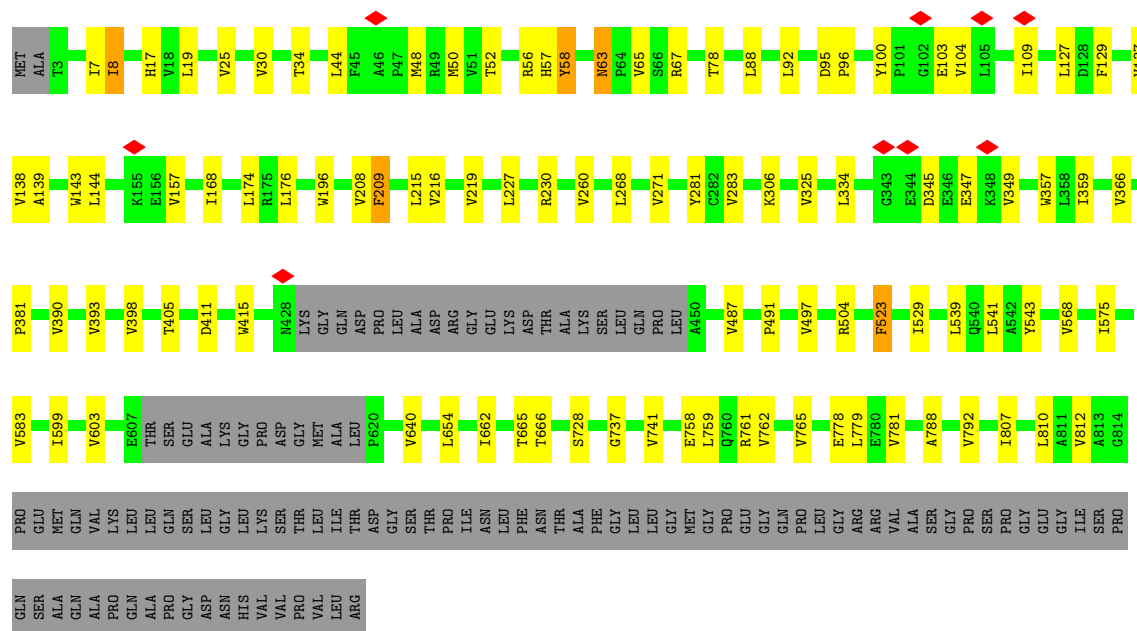
- Molecule 1: Major vault protein



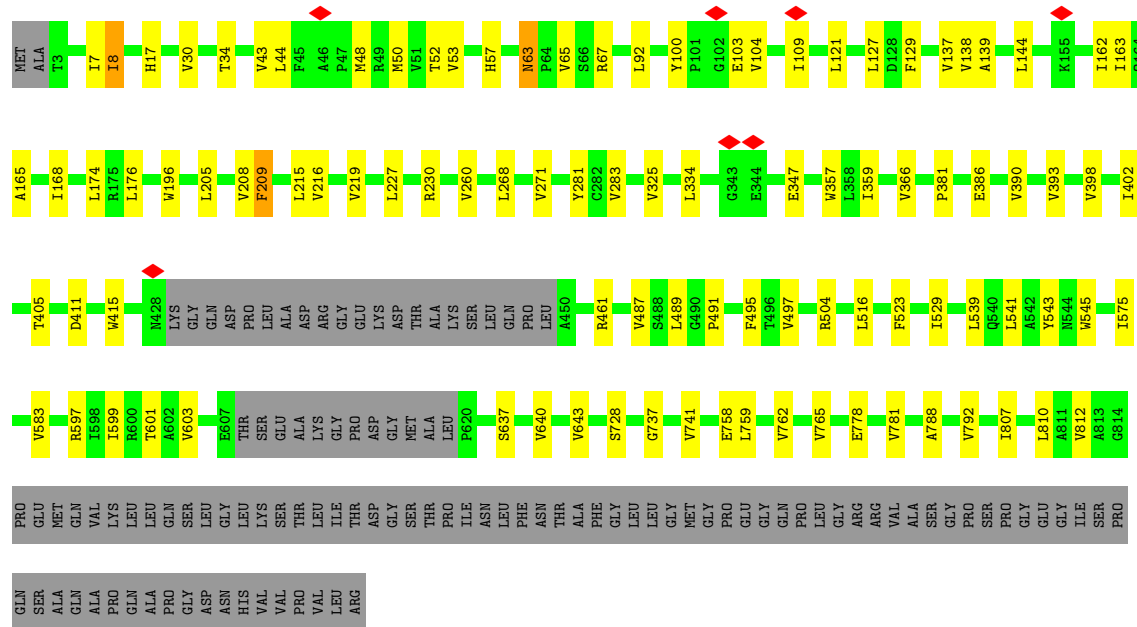
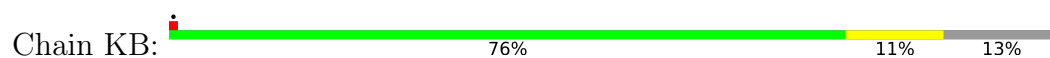
- Molecule 1: Major vault protein



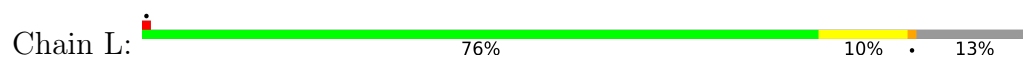


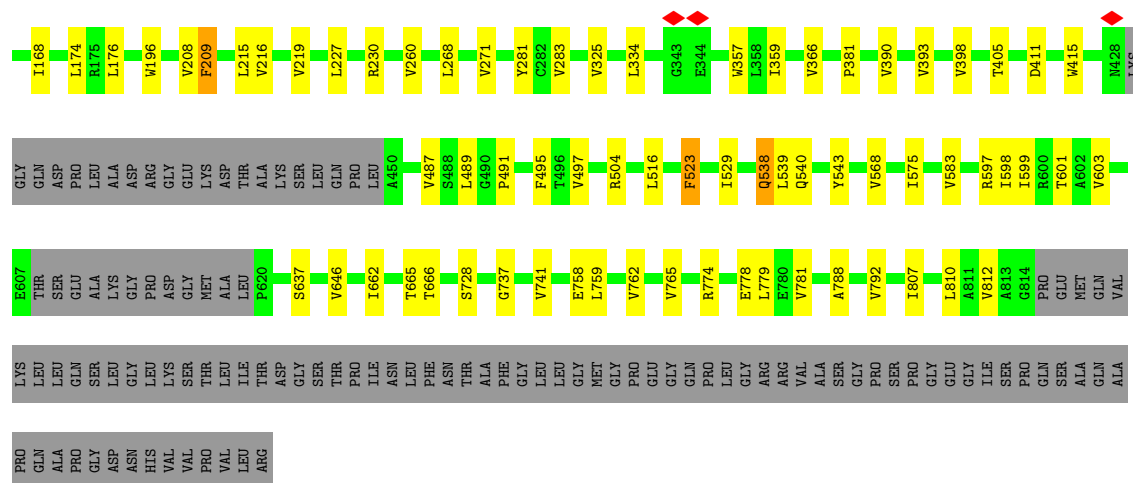


- Molecule 1: Major vault protein



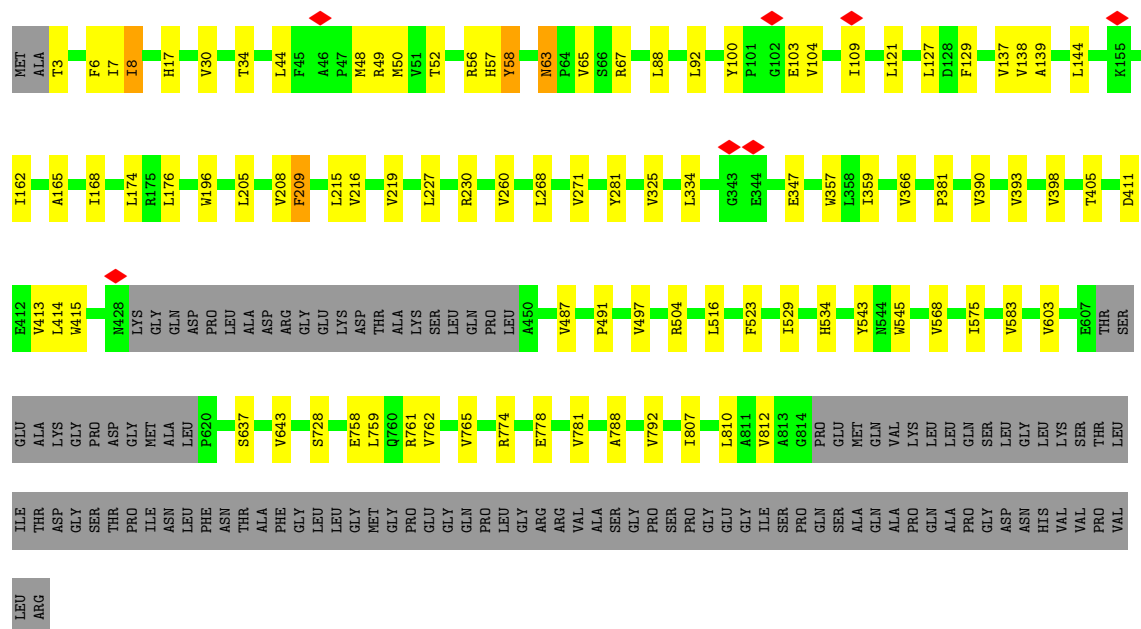
- Molecule 1: Major vault protein





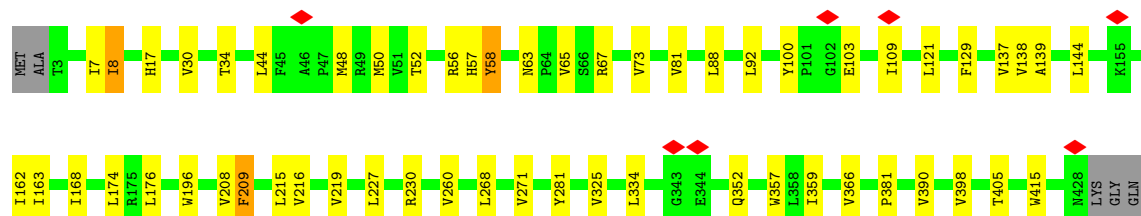
### • Molecule 1: Major vault protein

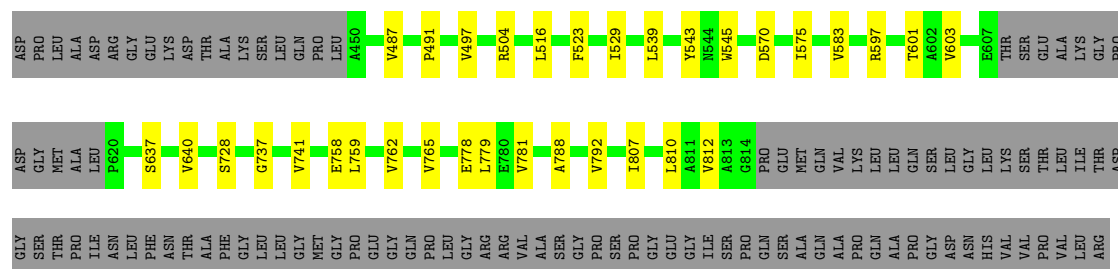
Chain LA: 77% 10% 13%



### • Molecule 1: Major vault protein

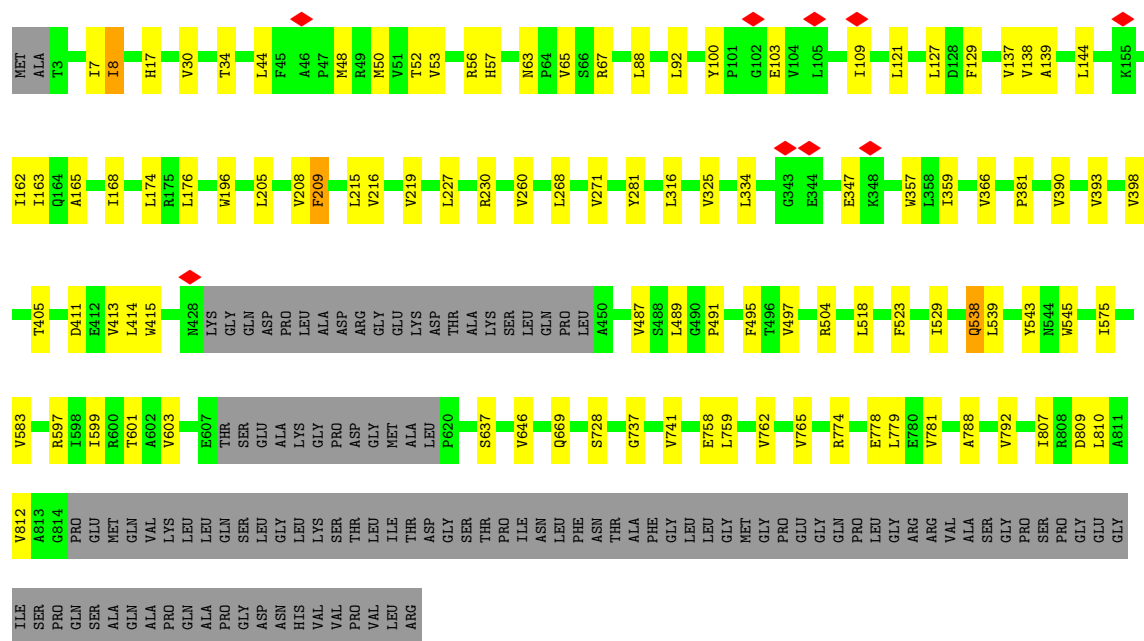
Chain LB: 77% 10% 13%





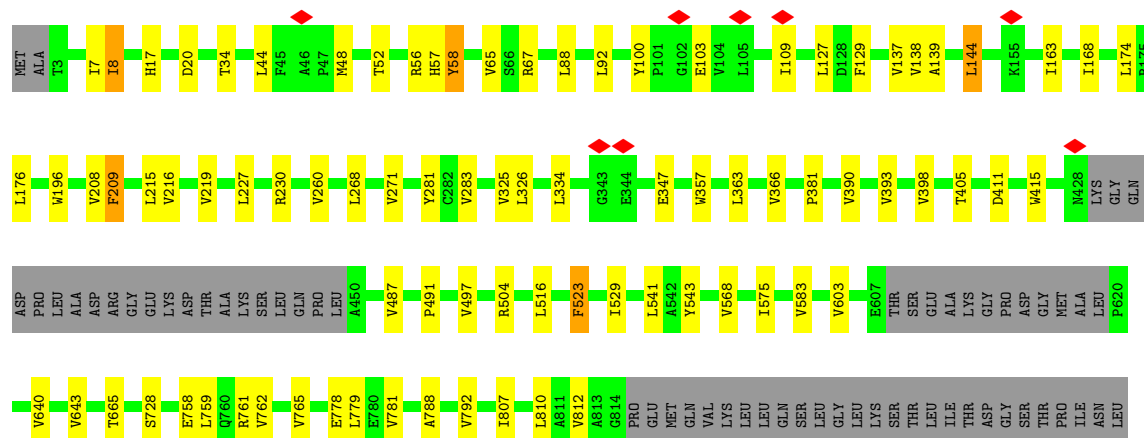
### • Molecule 1: Major vault protein

Chain M:




### • Molecule 1: Major vault protein

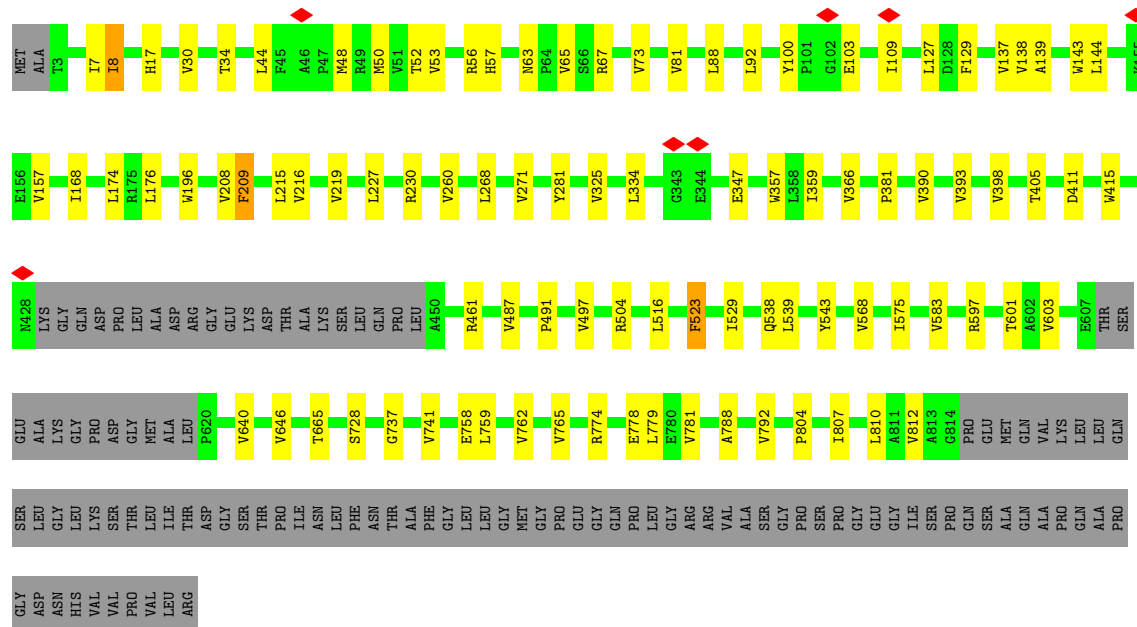
Chain MA:




PHE ASN THR PHE GLY LEU LEU MET GLY PRO GLU GLY GLN PRO PRO LEU LEU ARG ARG VAL ALA SER GLY PRO SER GLY GLY ILE SER PRO GLN SER ALA GLN ALA PRO GLN ALA PRO PRO GLY ASP ASN HIS VAL PRO VAL LEU ARG

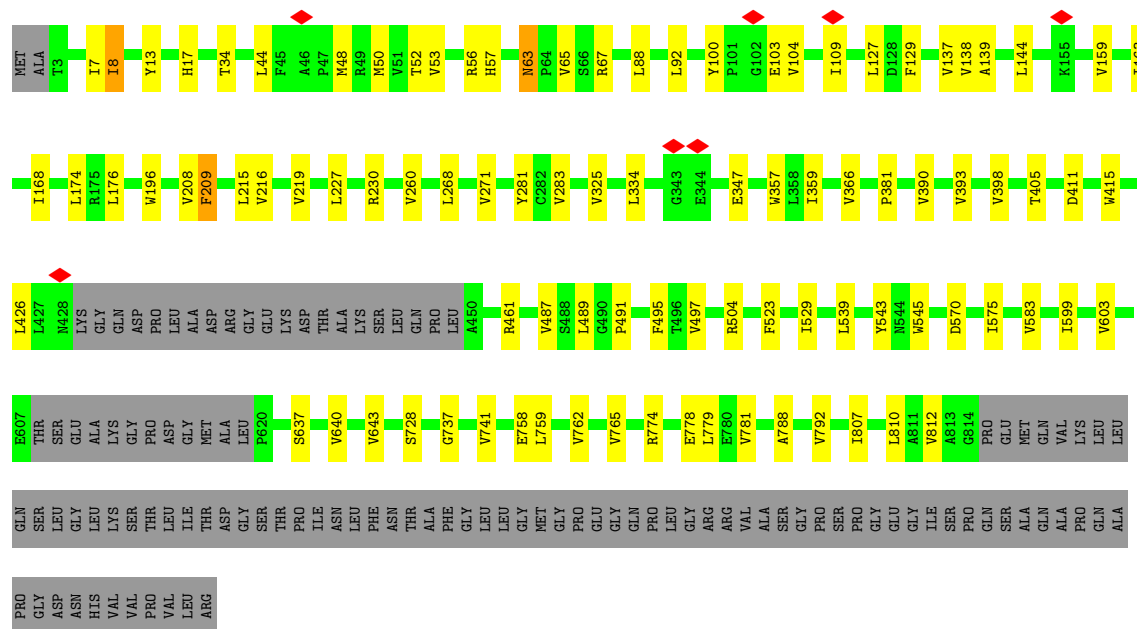
• Molecule 1: Major vault protein

Chain MB:  77% 10% 13%

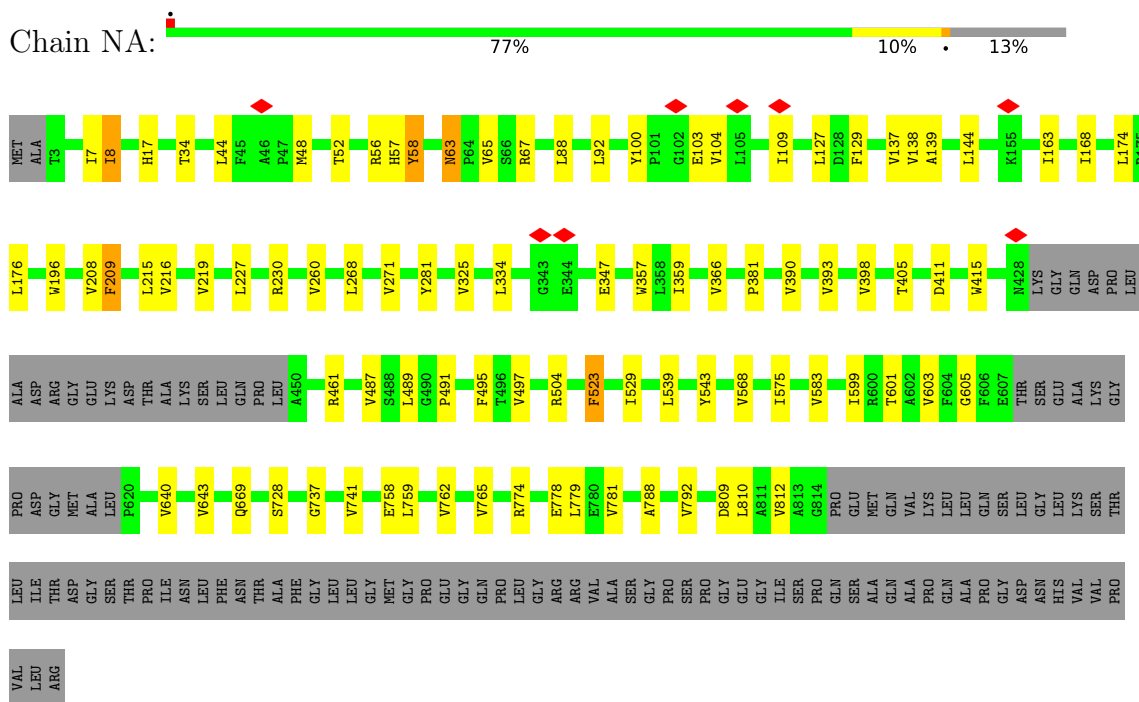


• Molecule 1: Major vault protein

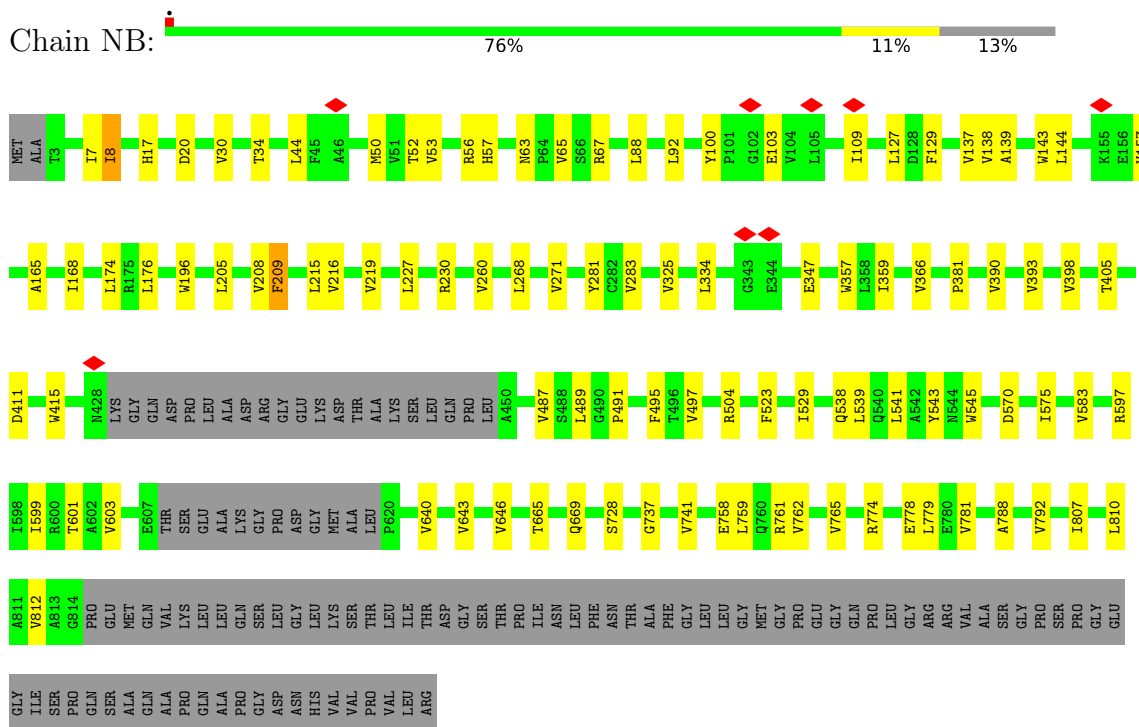
Chain N:  77% 10% 13%



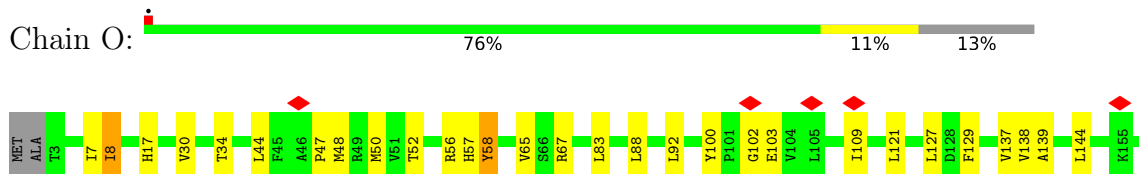
• Molecule 1: Major vault protein

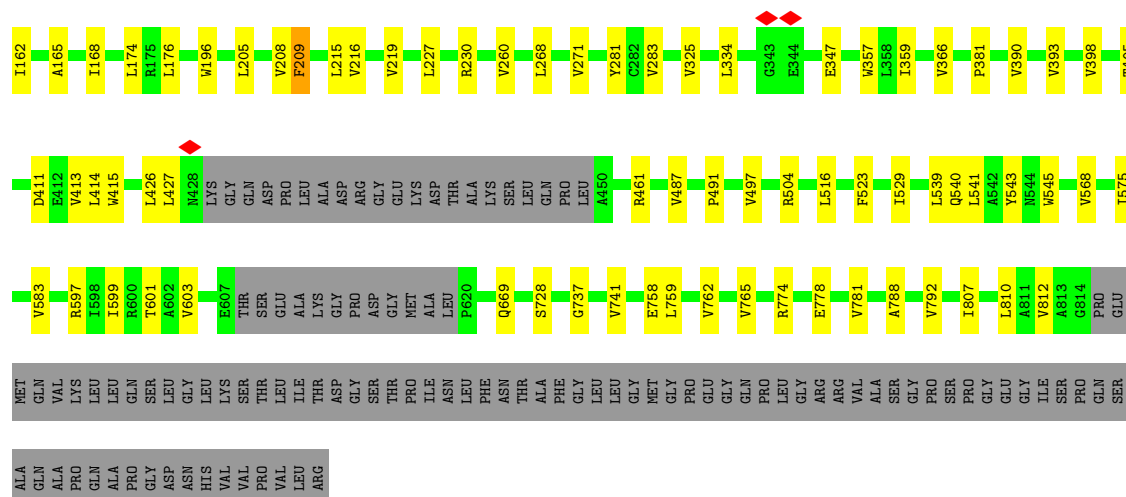


- Molecule 1: Major vault protein



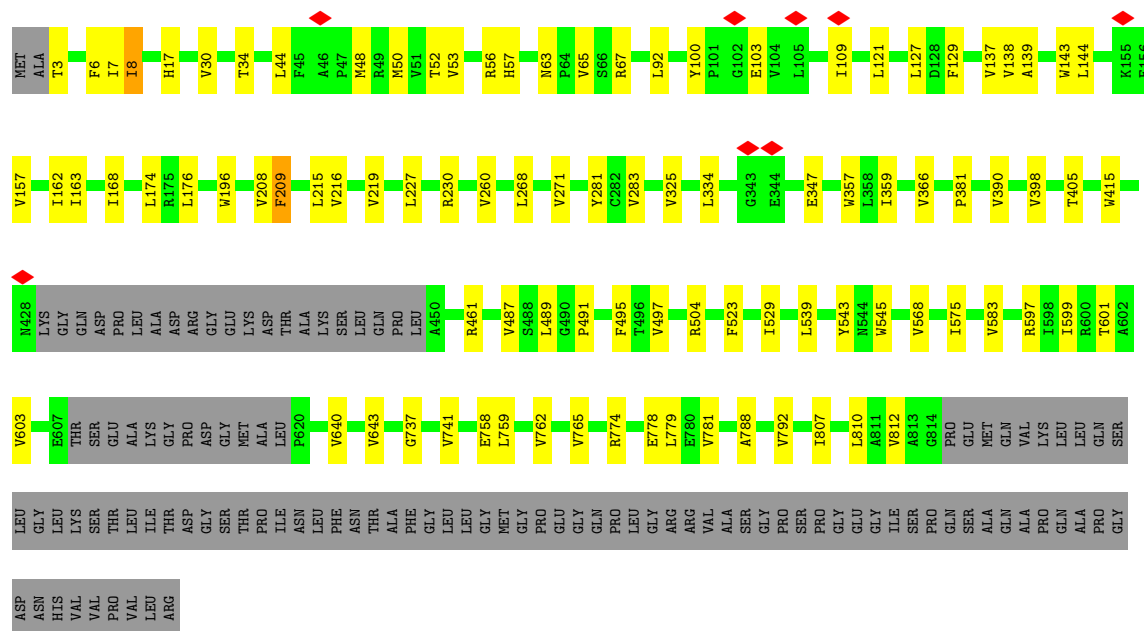
- Molecule 1: Major vault protein





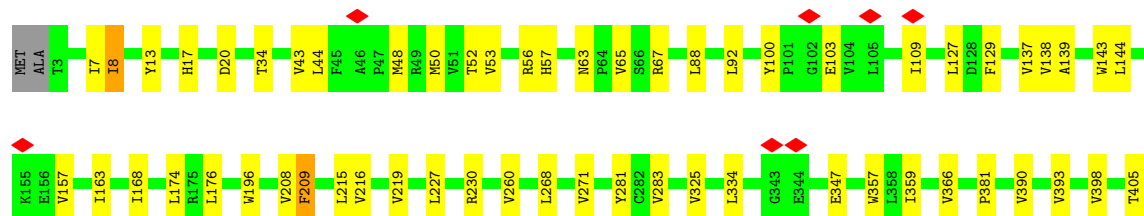
- Molecule 1: Major vault protein

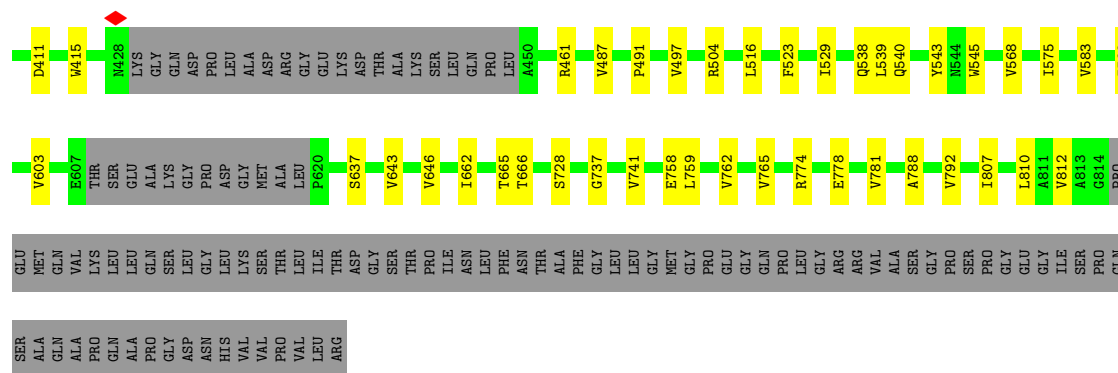
Chain OA: 77% 10% 13%



- Molecule 1: Major vault protein

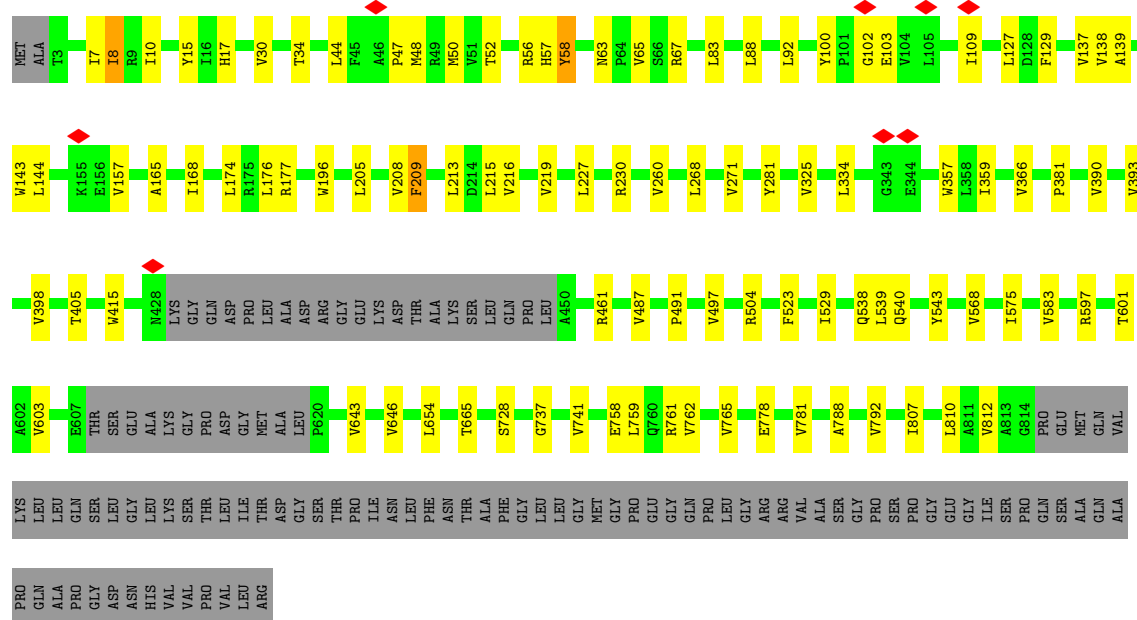
Chain OB: 76% 11% 13%

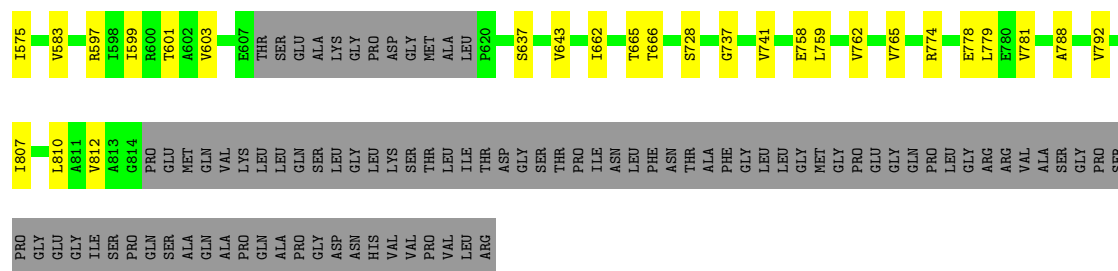




• Molecule 1: Major vault protein

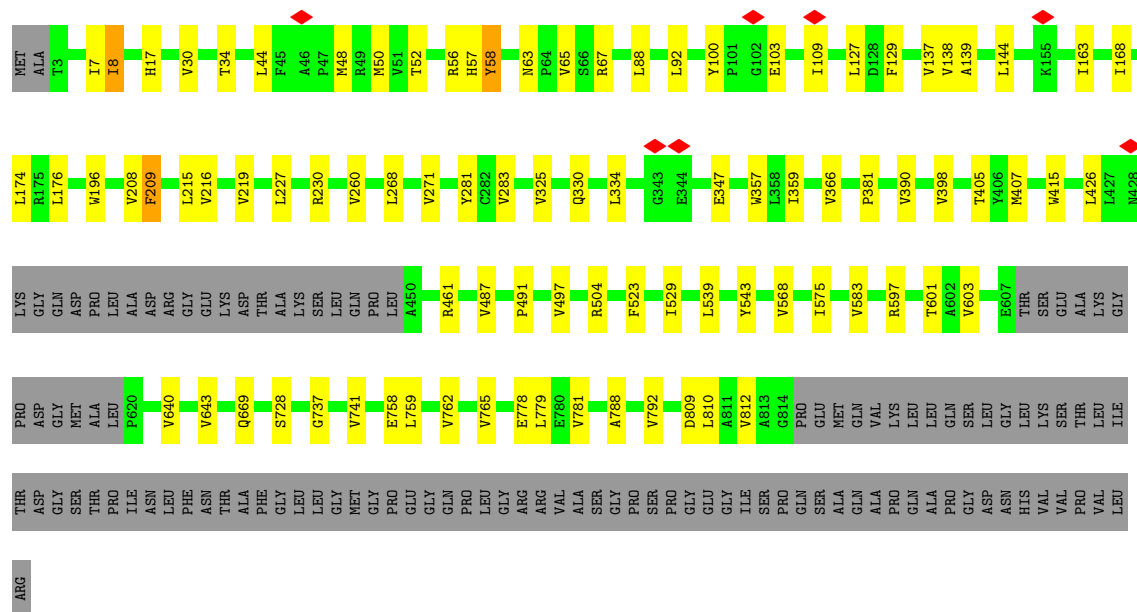
Chain P: 76% 11% 13%





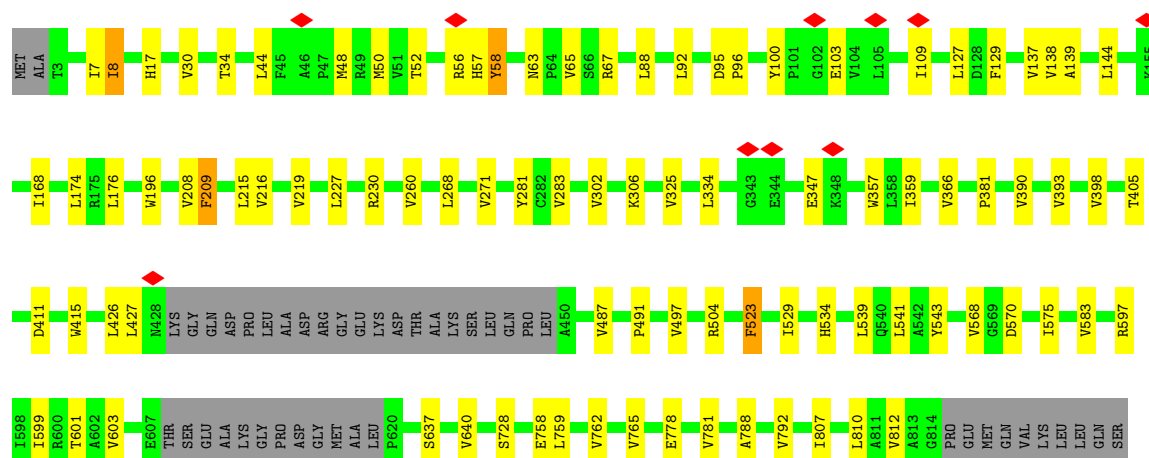
• Molecule 1: Major vault protein

Chain PB: 77% 10% 13%

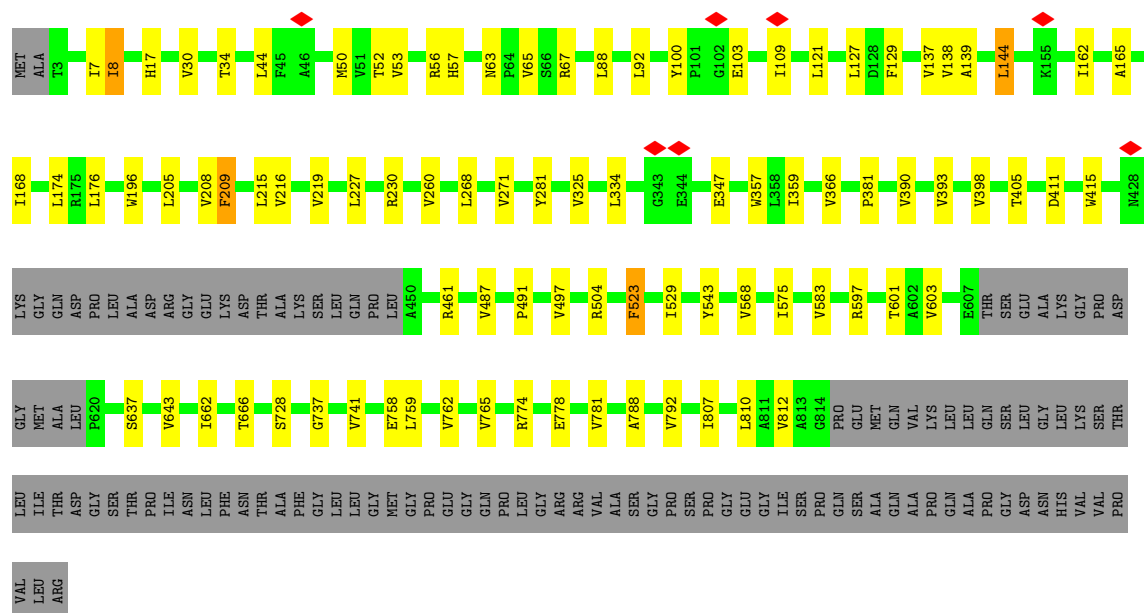


• Molecule 1: Major vault protein

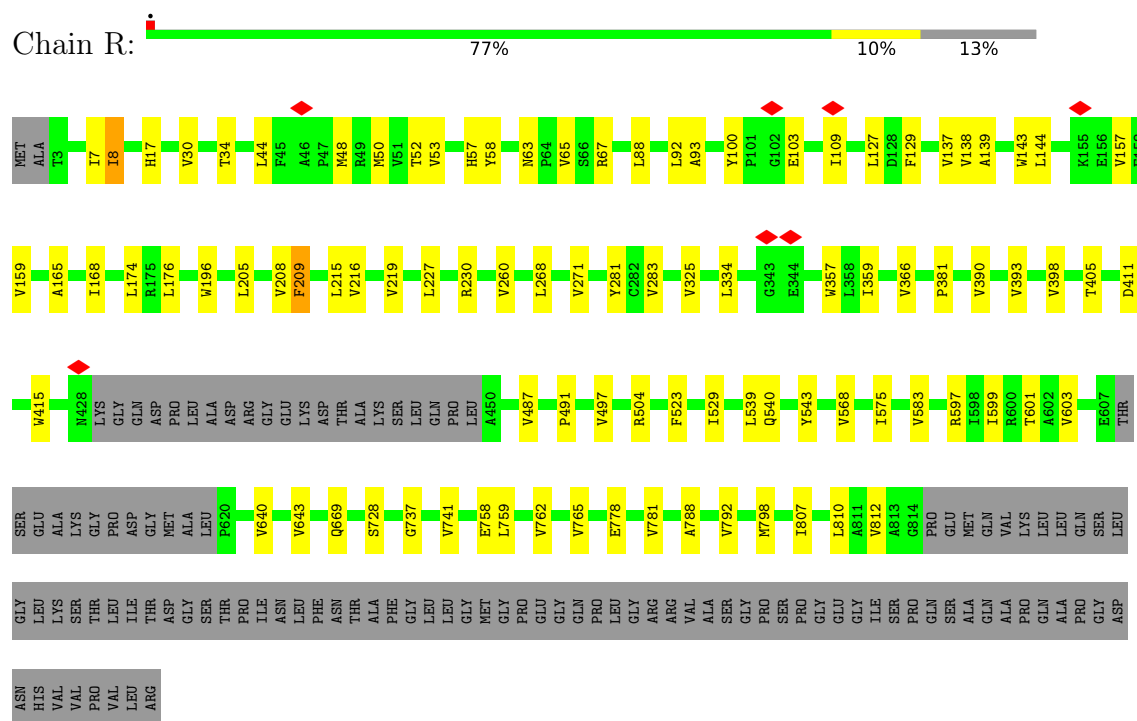
Chain Q: 77% 10% 13%



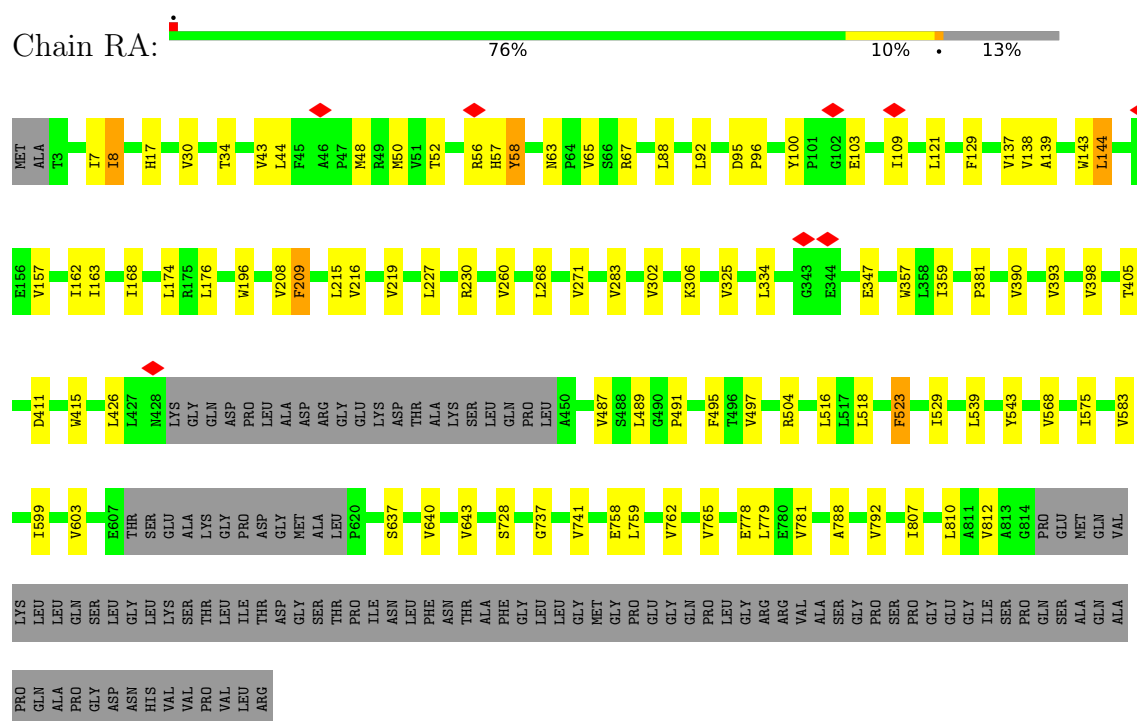




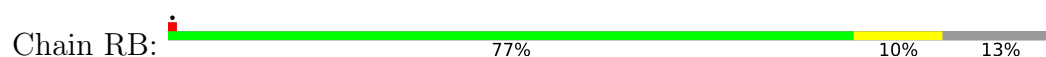
## • Molecule 1: Major vault protein

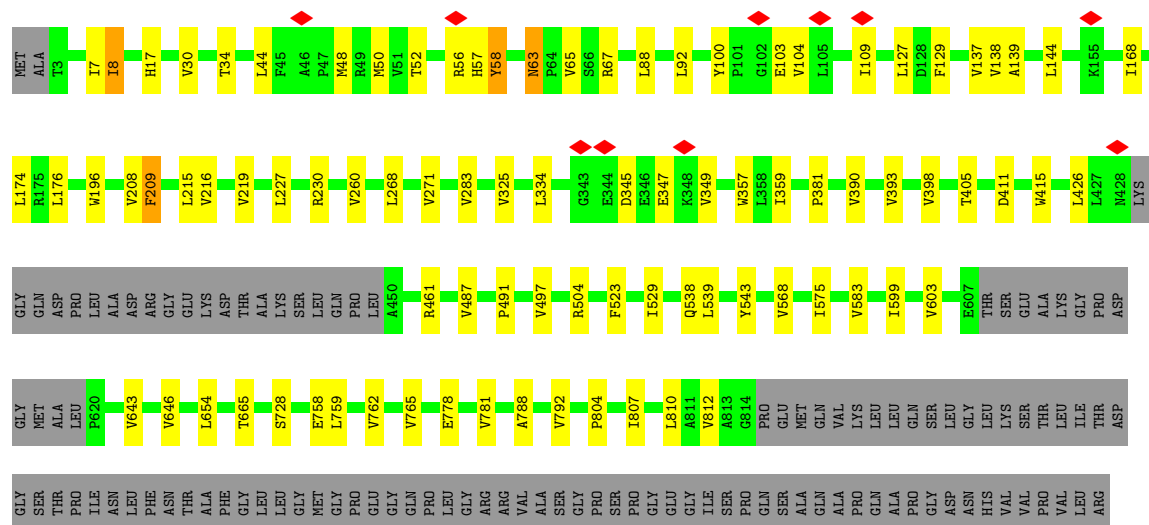


## • Molecule 1: Major vault protein

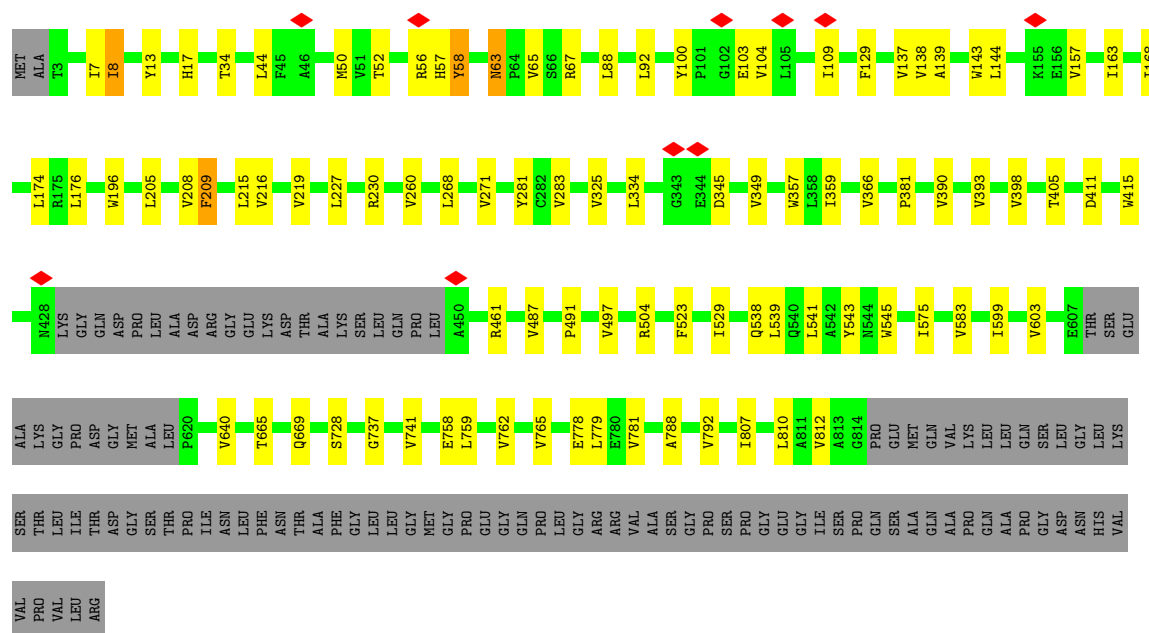
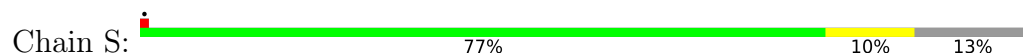


## • Molecule 1: Major vault protein

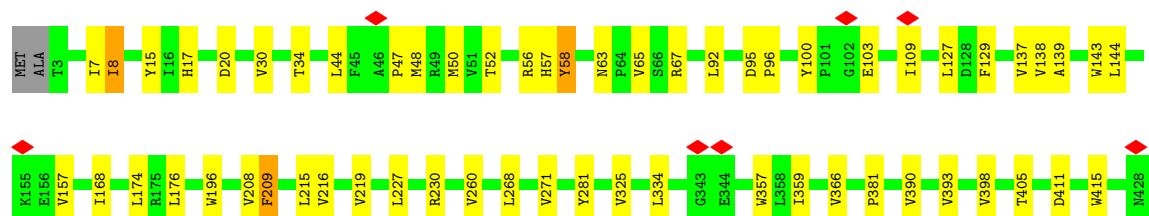
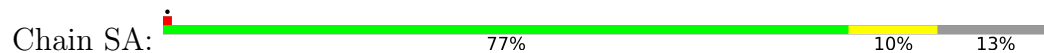


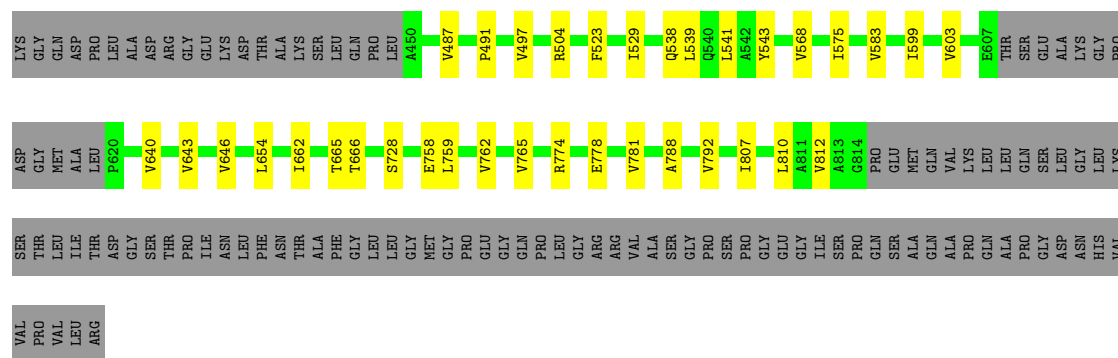


### • Molecule 1: Major vault protein

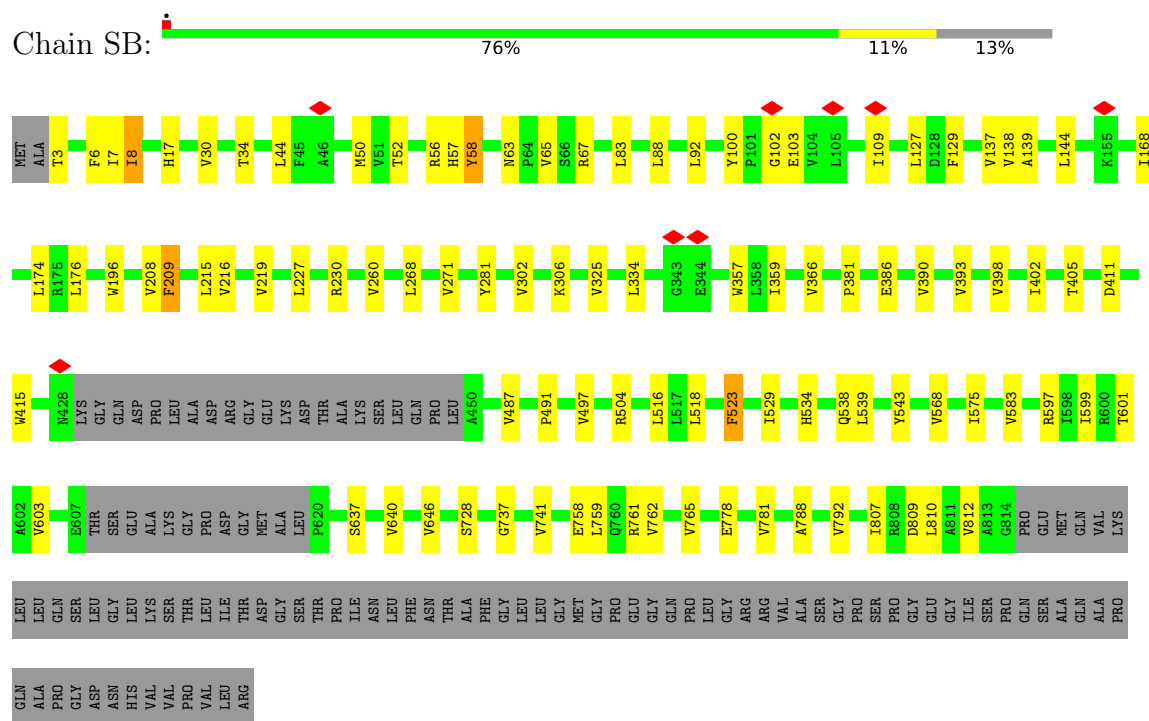


### • Molecule 1: Major vault protein

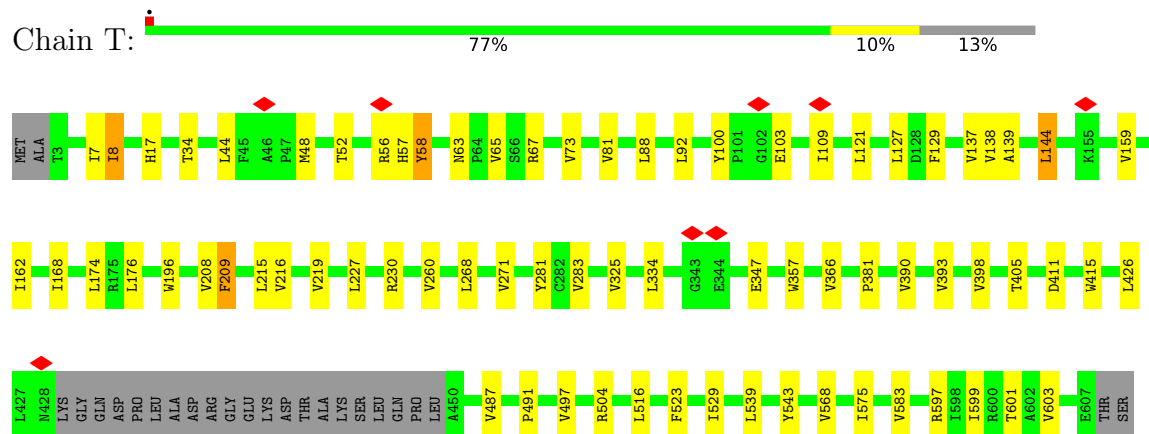


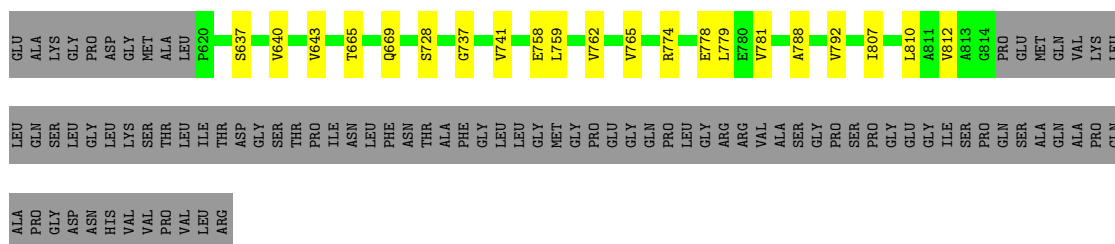


### • Molecule 1: Major vault protein




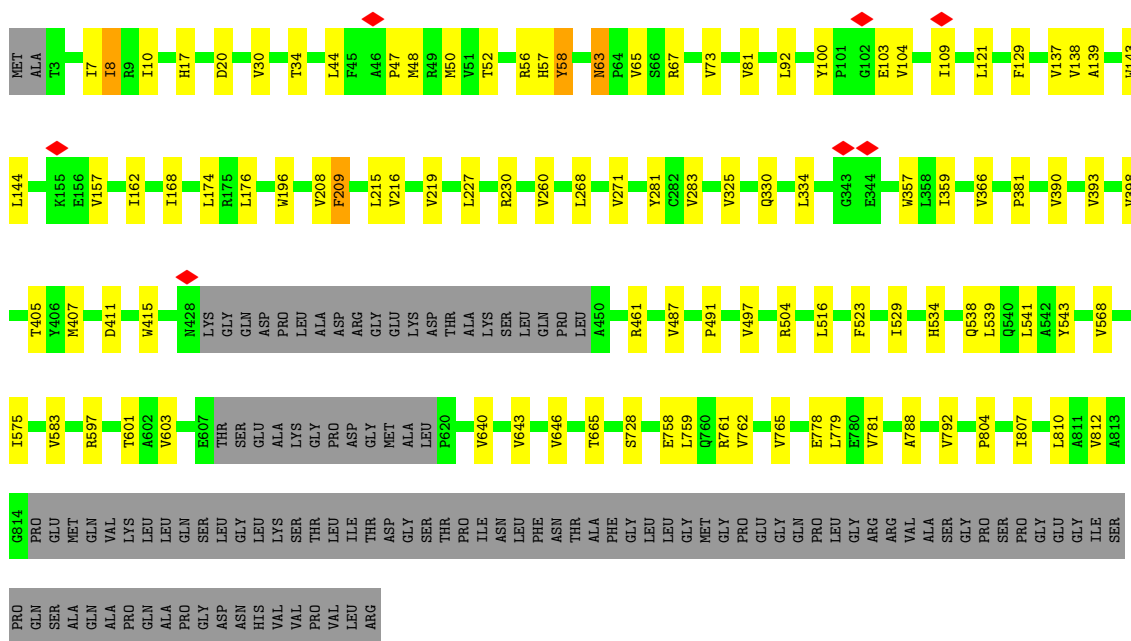
### • Molecule 1: Major vault protein






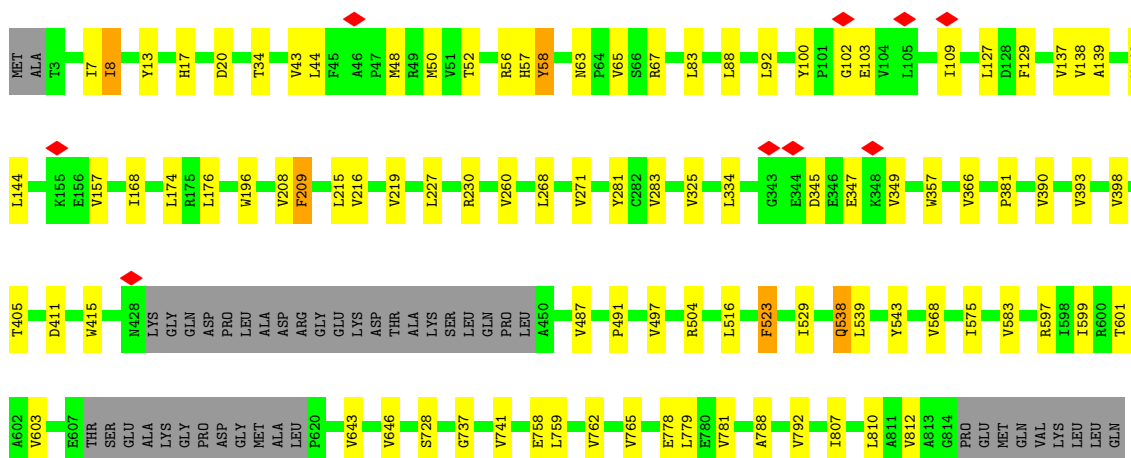
• Molecule 1: Major vault protein

Chain TA:  76% 11% 13%



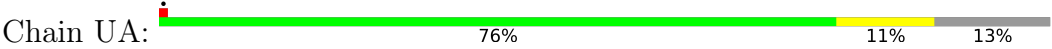
• Molecule 1: Major vault protein

Chain TB:  76% 10% 13%



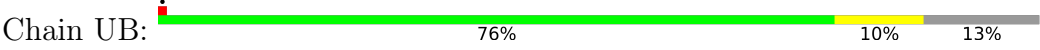
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GLY	ASP	ASN	HIS	VAL	VAL	PRO	VAL	LEU	LEU	ARG																																								

● Molecule 1: Major vault protein



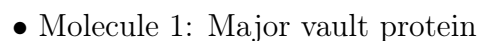
PRO	GLN	ALA	PRO	GLY	ASP	ASN	HIS	VAL	VAL	PRO	VAL	LEU	ARG																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																												
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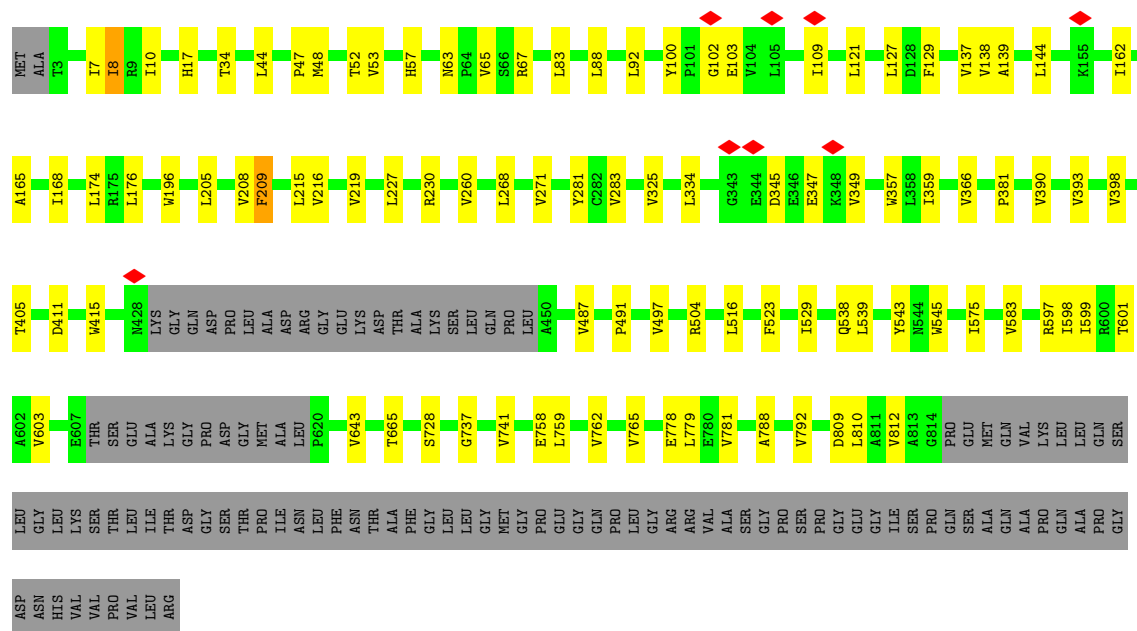
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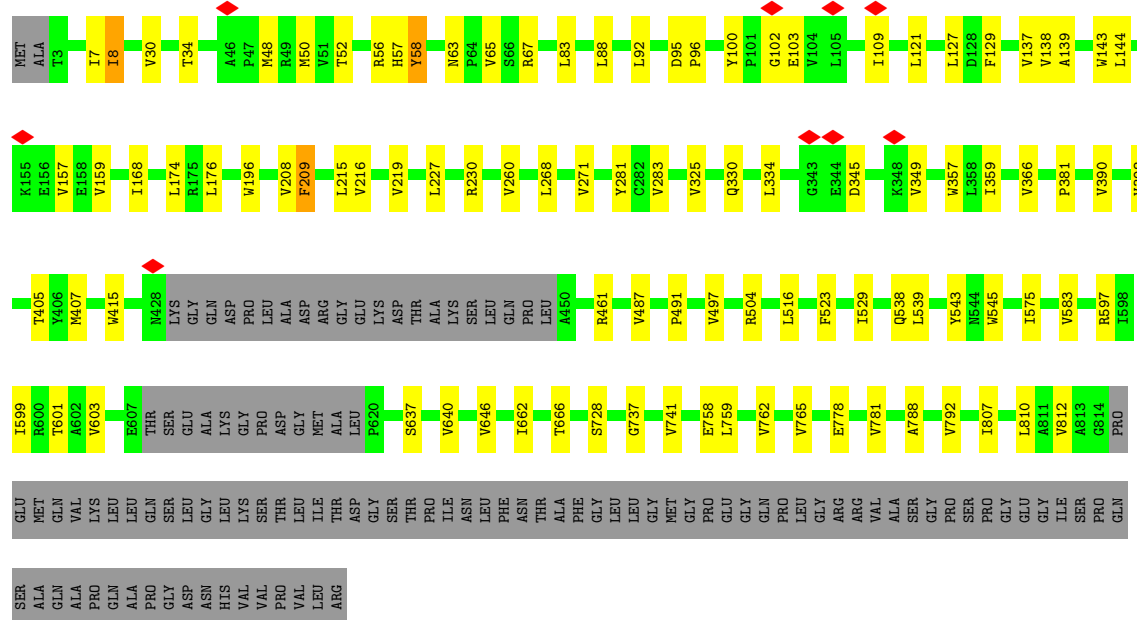
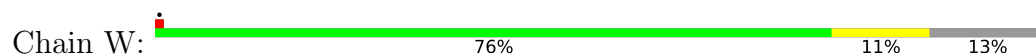
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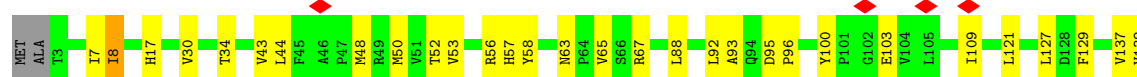
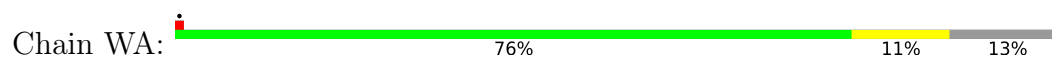




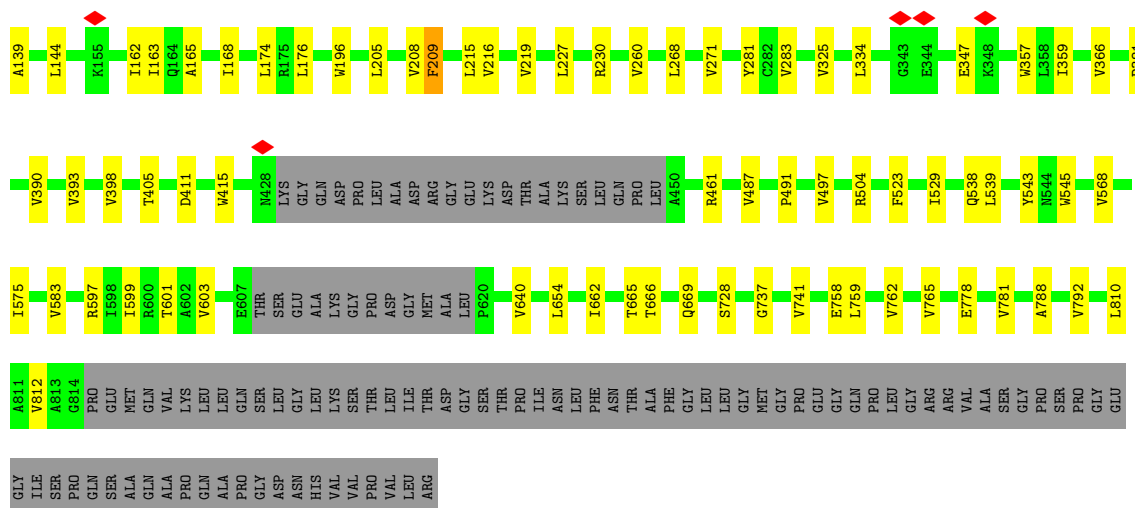
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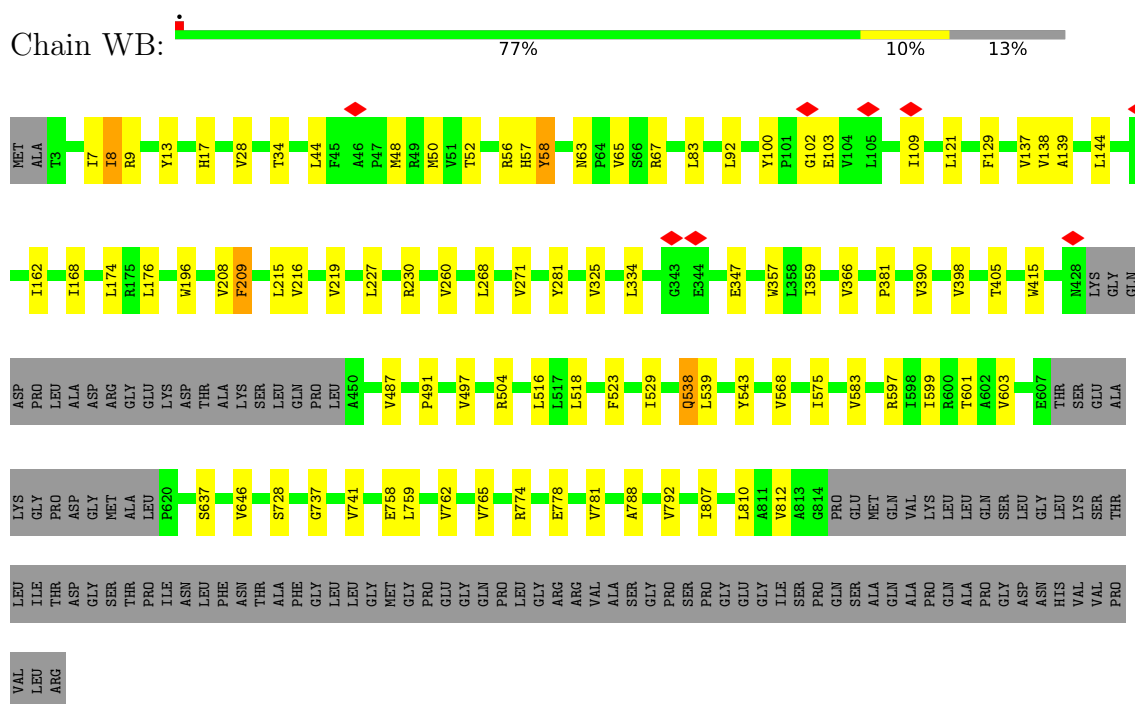
- Molecule 1: Major vault protein



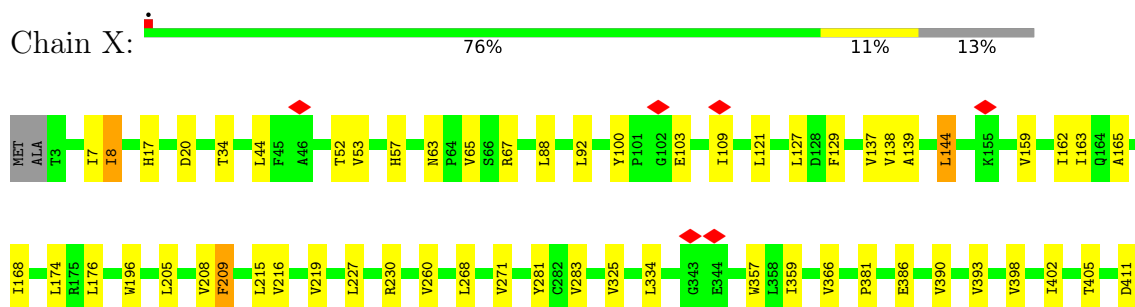


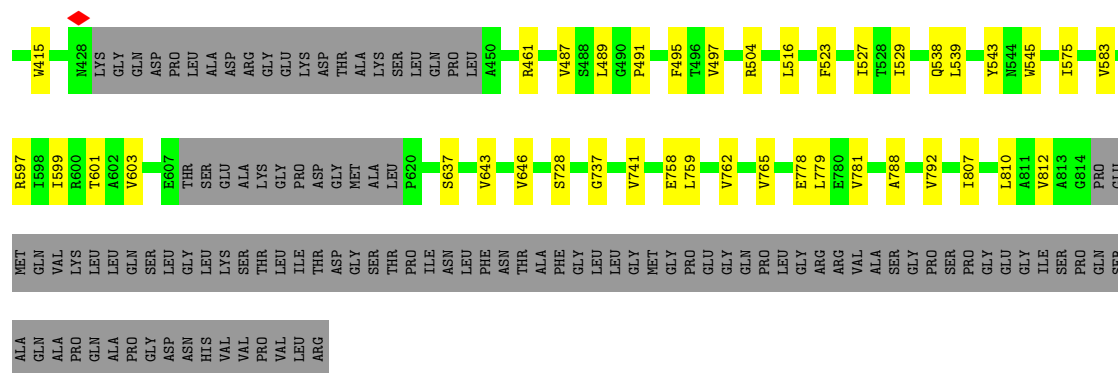


• Molecule 1: Major vault protein



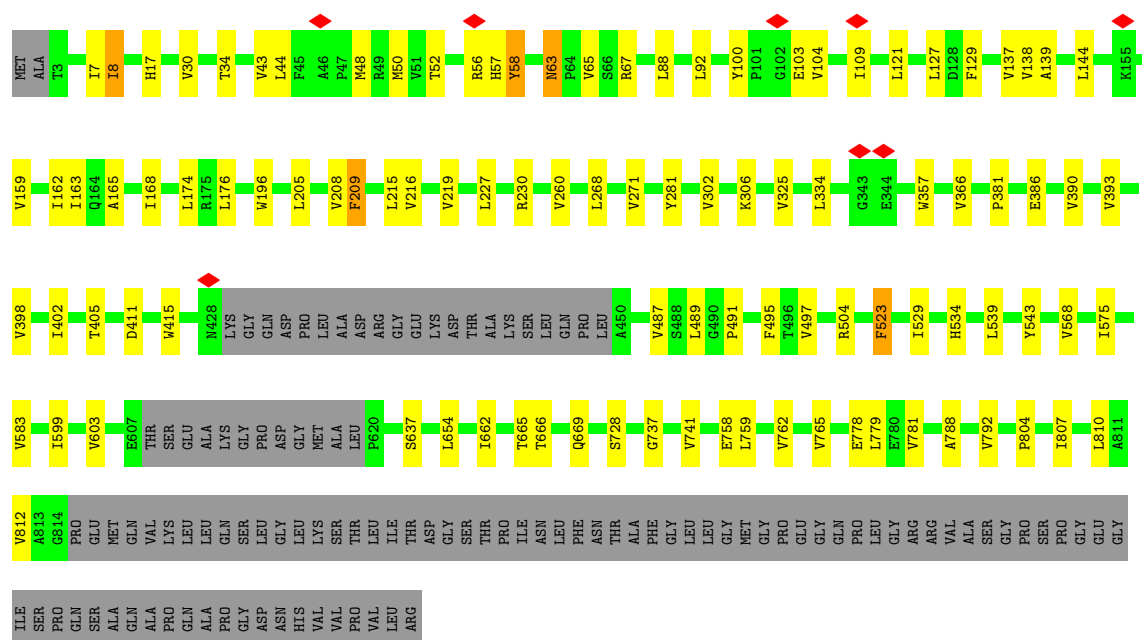
• Molecule 1: Major vault protein





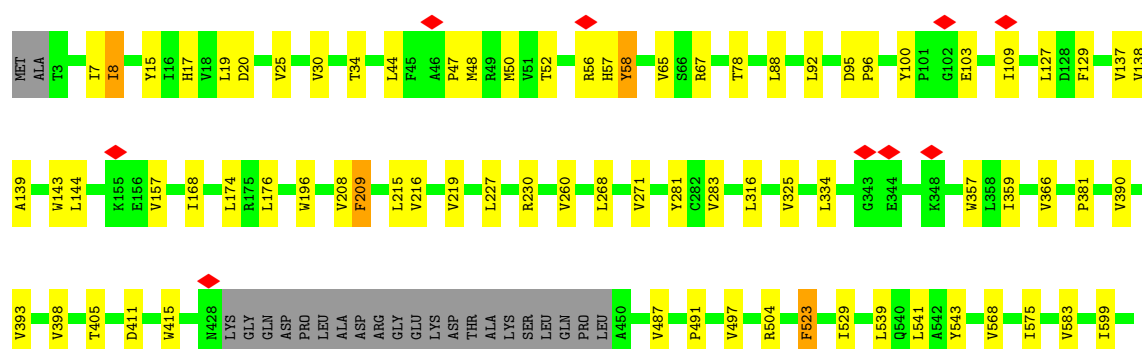
• Molecule 1: Major vault protein

Chain XA: 76% 11% 13%

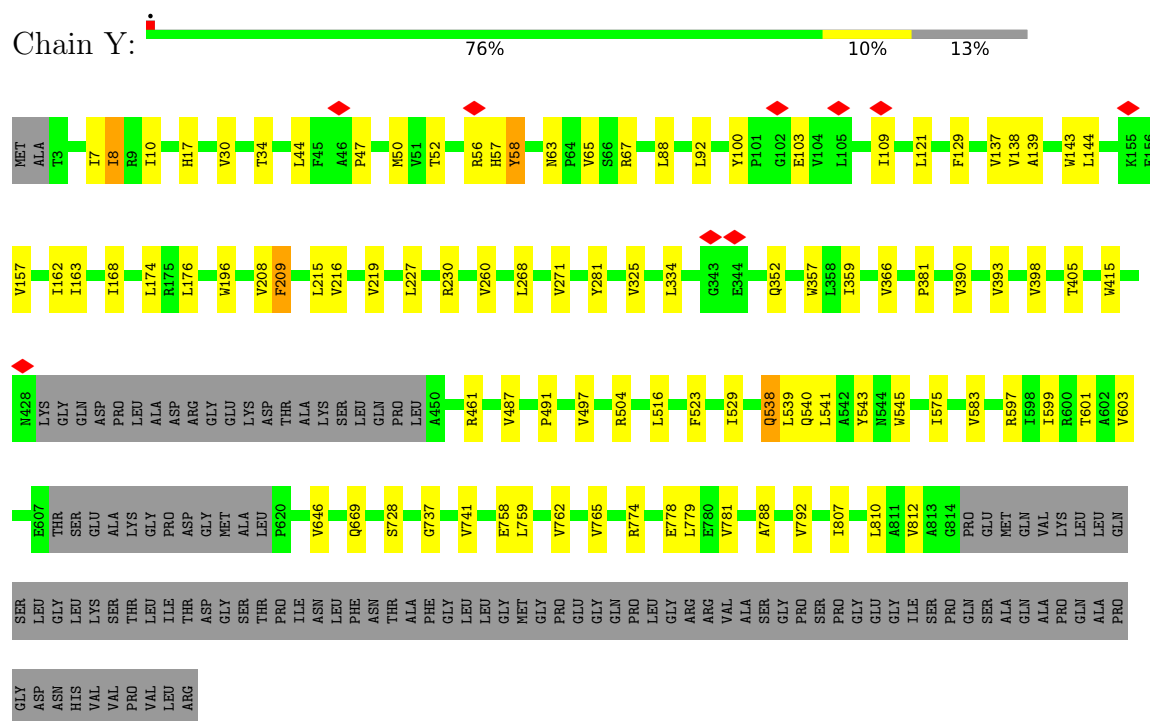


• Molecule 1: Major vault protein

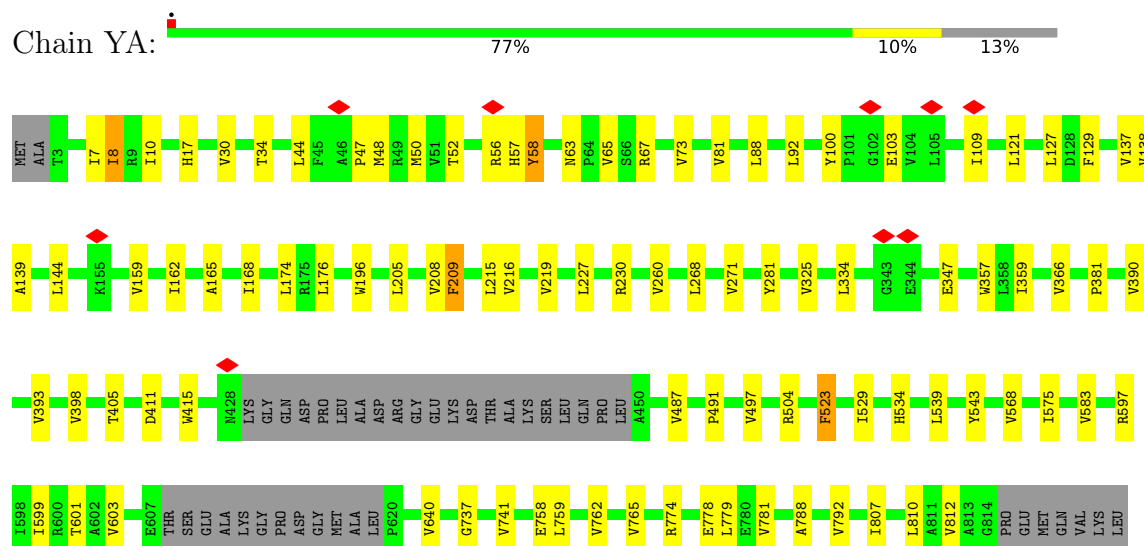
Chain XB: 76% 11% 13%



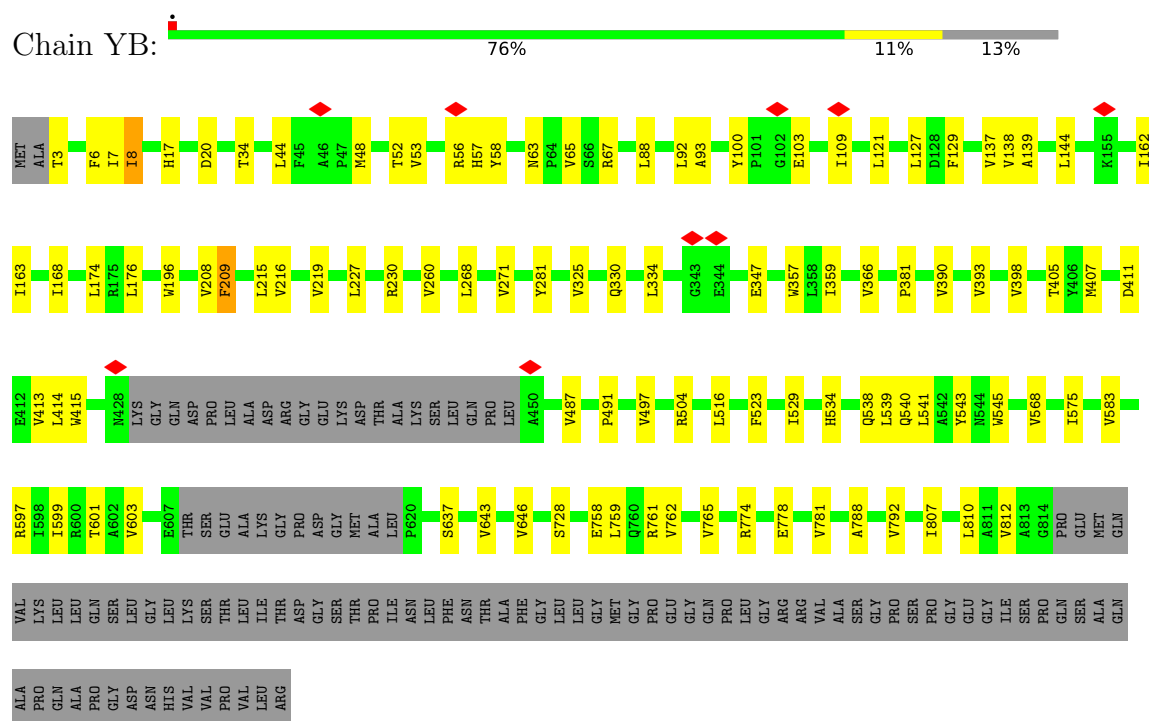
- Molecule 1: Major vault protein



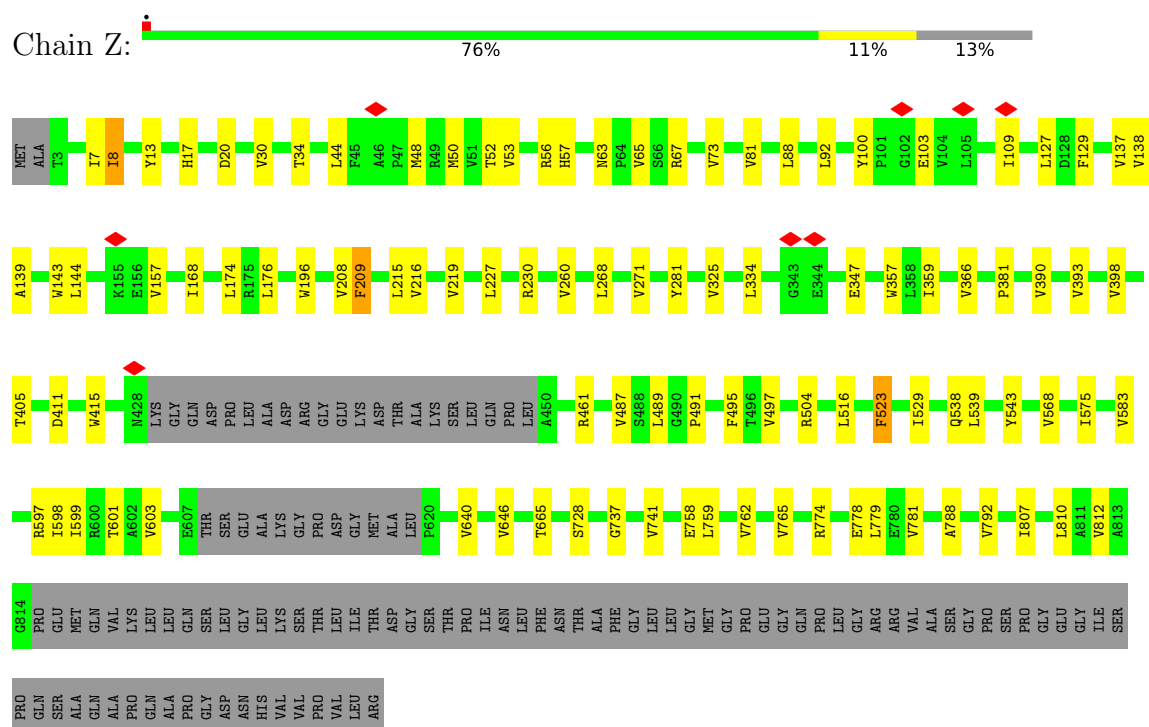
- Molecule 1: Major vault protein



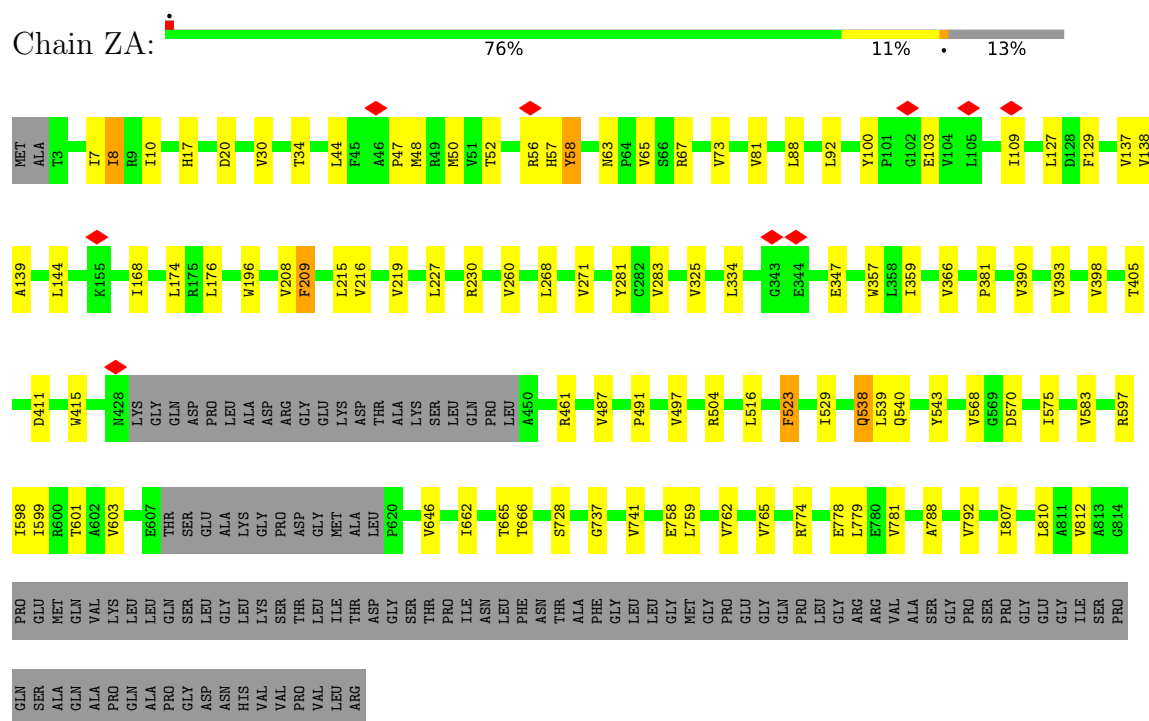
- Molecule 1: Major vault protein



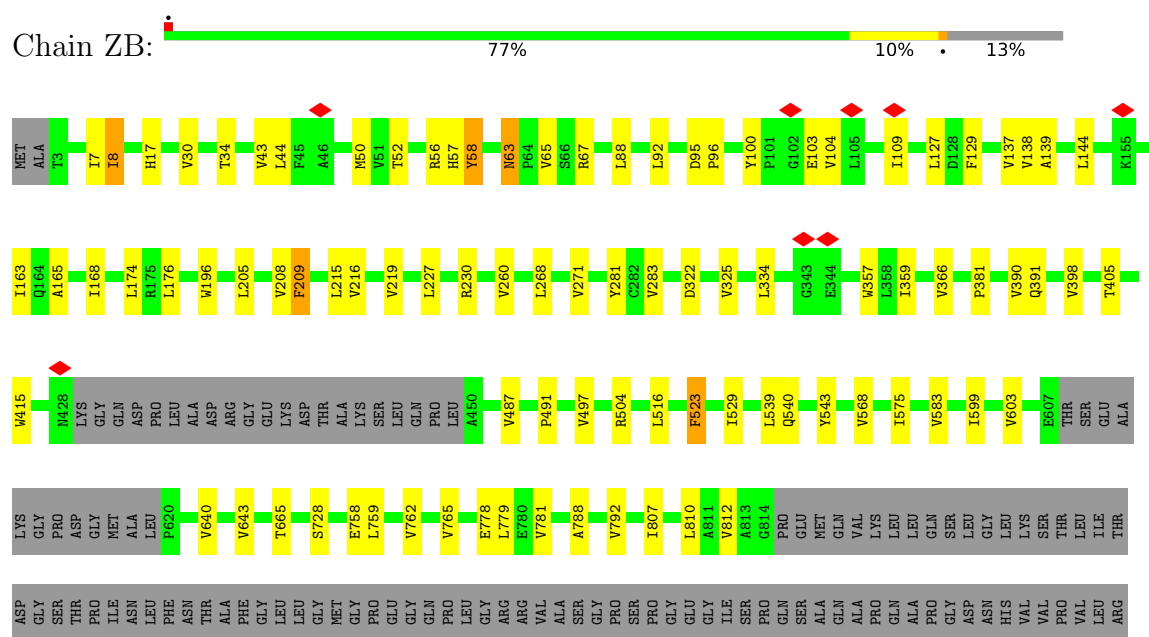
- Molecule 1: Major vault protein



- Molecule 1: Major vault protein



- Molecule 1: Major vault protein



## 4 Experimental information

Property	Value	Source
EM reconstruction method	SINGLE PARTICLE	Depositor
Imposed symmetry	POINT, D39	Depositor
Number of particles used	11172	Depositor
Resolution determination method	FSC 0.143 CUT-OFF	Depositor
CTF correction method	PHASE FLIPPING AND AMPLITUDE CORRECTION	Depositor
Microscope	TFS KRIOS	Depositor
Voltage (kV)	300	Depositor
Electron dose ( $e^-/\text{\AA}^2$ )	50.15	Depositor
Minimum defocus (nm)	551	Depositor
Maximum defocus (nm)	2330	Depositor
Magnification	81000	Depositor
Image detector	GATAN K3 BIOQUANTUM (6k x 4k)	Depositor
Maximum map value	1.881	Depositor
Minimum map value	-0.270	Depositor
Average map value	0.006	Depositor
Map value standard deviation	0.099	Depositor
Recommended contour level	0.6	Depositor
Map size (Å)	861.7984, 861.7984, 861.7984	wwPDB
Map dimensions	1024, 1024, 1024	wwPDB
Map angles (°)	90.0, 90.0, 90.0	wwPDB
Pixel spacing (Å)	0.8416, 0.8416, 0.8416	Depositor

## 5 Model quality [i](#)

### 5.1 Standard geometry [i](#)

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	0.17	0/6291	0.40	0/8533
1	AA	0.17	0/6291	0.40	0/8533
1	AB	0.17	0/6291	0.40	0/8533
1	AC	0.17	0/6291	0.40	0/8533
1	B	0.17	0/6291	0.40	0/8533
1	BA	0.17	0/6291	0.40	0/8533
1	BB	0.17	0/6291	0.40	0/8533
1	C	0.17	0/6291	0.39	0/8533
1	CA	0.17	0/6291	0.40	0/8533
1	CB	0.17	0/6291	0.40	0/8533
1	D	0.17	0/6291	0.40	0/8533
1	DA	0.17	0/6291	0.40	0/8533
1	DB	0.17	0/6291	0.40	0/8533
1	E	0.17	0/6291	0.40	0/8533
1	EA	0.17	0/6291	0.40	0/8533
1	EB	0.17	0/6291	0.39	0/8533
1	F	0.17	0/6291	0.40	0/8533
1	FA	0.17	0/6291	0.40	0/8533
1	FB	0.17	0/6291	0.40	0/8533
1	G	0.17	0/6291	0.40	0/8533
1	GA	0.17	0/6291	0.40	0/8533
1	GB	0.17	0/6291	0.40	0/8533
1	H	0.17	0/6291	0.39	0/8533
1	HA	0.17	0/6291	0.40	0/8533
1	HB	0.17	0/6291	0.40	0/8533
1	I	0.17	0/6291	0.39	0/8533
1	IA	0.17	0/6291	0.40	0/8533
1	IB	0.17	0/6291	0.40	0/8533
1	J	0.17	0/6291	0.40	0/8533
1	JA	0.17	0/6291	0.40	0/8533
1	JB	0.17	0/6291	0.40	0/8533
1	K	0.17	0/6291	0.40	0/8533
1	KA	0.17	0/6291	0.40	0/8533
1	KB	0.17	0/6291	0.40	0/8533

Mol	Chain	Bond lengths		Bond angles	
		RMSZ	# Z  >5	RMSZ	# Z  >5
1	L	0.17	0/6291	0.40	0/8533
1	LA	0.17	0/6291	0.40	0/8533
1	LB	0.17	0/6291	0.40	0/8533
1	M	0.17	0/6291	0.40	0/8533
1	MA	0.17	0/6291	0.40	0/8533
1	MB	0.17	0/6291	0.40	0/8533
1	N	0.17	0/6291	0.40	0/8533
1	NA	0.17	0/6291	0.40	0/8533
1	NB	0.17	0/6291	0.39	0/8533
1	O	0.17	0/6291	0.40	0/8533
1	OA	0.17	0/6291	0.40	0/8533
1	OB	0.17	0/6291	0.40	0/8533
1	P	0.17	0/6291	0.40	0/8533
1	PA	0.17	0/6291	0.40	0/8533
1	PB	0.17	0/6291	0.40	0/8533
1	Q	0.17	0/6291	0.39	0/8533
1	QA	0.17	0/6291	0.40	0/8533
1	QB	0.17	0/6291	0.40	0/8533
1	R	0.17	0/6291	0.40	0/8533
1	RA	0.17	0/6291	0.40	0/8533
1	RB	0.17	0/6291	0.40	0/8533
1	S	0.17	0/6291	0.40	0/8533
1	SA	0.17	0/6291	0.40	0/8533
1	SB	0.17	0/6291	0.40	0/8533
1	T	0.17	0/6291	0.40	0/8533
1	TA	0.17	0/6291	0.40	0/8533
1	TB	0.17	0/6291	0.40	0/8533
1	UA	0.17	0/6291	0.40	0/8533
1	UB	0.17	0/6291	0.40	0/8533
1	V	0.17	0/6291	0.40	0/8533
1	VA	0.17	0/6291	0.40	0/8533
1	VB	0.17	0/6291	0.40	0/8533
1	W	0.17	0/6291	0.39	0/8533
1	WA	0.17	0/6291	0.40	0/8533
1	WB	0.17	0/6291	0.40	0/8533
1	X	0.17	0/6291	0.39	0/8533
1	XA	0.17	0/6291	0.39	0/8533
1	XB	0.17	0/6291	0.40	0/8533
1	Y	0.17	0/6291	0.40	0/8533
1	YA	0.17	0/6291	0.40	0/8533
1	YB	0.17	0/6291	0.40	0/8533
1	Z	0.17	0/6291	0.40	0/8533
1	ZA	0.17	0/6291	0.40	0/8533



Mol	Chain	Bond lengths		Bond angles	
		RMSZ	# Z  >5	RMSZ	# Z  >5
1	ZB	0.17	0/6291	0.40	0/8533
All	All	0.17	0/490698	0.40	0/665574

There are no bond length outliers.

There are no bond angle outliers.

There are no chirality outliers.

There are no planarity outliers.

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

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

Mol	Chain	Non-H	H(model)	H(added)	Clashes	Symm-Clashes
1	A	6181	6205	6203	75	0
1	AA	6181	6205	6203	66	0
1	AB	6181	6205	6203	75	0
1	AC	6181	6205	6203	74	0
1	B	6181	6205	6203	71	0
1	BA	6181	6205	6203	69	0
1	BB	6181	6205	6203	76	0
1	C	6181	6205	6203	65	0
1	CA	6181	6205	6203	66	0
1	CB	6181	6205	6203	76	0
1	D	6181	6205	6203	65	0
1	DA	6181	6205	6203	70	0
1	DB	6181	6205	6203	75	0
1	E	6181	6205	6203	65	0
1	EA	6181	6205	6203	62	0
1	EB	6181	6205	6203	71	0
1	F	6181	6205	6203	70	0
1	FA	6181	6205	6203	72	0
1	FB	6181	6205	6203	69	0
1	G	6181	6205	6203	66	0
1	GA	6181	6205	6203	68	0
1	GB	6181	6205	6203	67	0
1	H	6181	6205	6203	67	0
1	HA	6181	6205	6203	68	0

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Mol	Chain	Non-H	H(model)	H(added)	Clashes	Symm-Clashes
1	HB	6181	6205	6203	70	0
1	I	6181	6205	6203	69	0
1	IA	6181	6205	6203	62	0
1	IB	6181	6205	6203	73	0
1	J	6181	6205	6203	73	0
1	JA	6181	6205	6203	66	0
1	JB	6181	6205	6203	68	0
1	K	6181	6205	6203	70	0
1	KA	6181	6205	6203	75	0
1	KB	6181	6205	6203	66	0
1	L	6181	6205	6203	69	0
1	LA	6181	6205	6203	73	0
1	LB	6181	6205	6203	62	0
1	M	6181	6205	6203	74	0
1	MA	6181	6205	6203	63	0
1	MB	6181	6205	6203	70	0
1	N	6181	6205	6203	70	0
1	NA	6181	6205	6203	72	0
1	NB	6181	6205	6203	68	0
1	O	6181	6205	6203	74	0
1	OA	6181	6205	6203	72	0
1	OB	6181	6205	6203	68	0
1	P	6181	6205	6203	74	0
1	PA	6181	6205	6203	75	0
1	PB	6181	6205	6203	67	0
1	Q	6181	6205	6203	69	0
1	QA	6181	6205	6203	64	0
1	QB	6181	6205	6203	64	0
1	R	6181	6205	6203	70	0
1	RA	6181	6205	6203	66	0
1	RB	6181	6205	6203	61	0
1	S	6181	6205	6203	66	0
1	SA	6181	6205	6203	69	0
1	SB	6181	6205	6203	67	0
1	T	6181	6205	6203	67	0
1	TA	6181	6205	6203	72	0
1	TB	6181	6205	6203	65	0
1	UA	6181	6205	6203	71	0
1	UB	6181	6205	6203	69	0
1	V	6181	6205	6203	69	0
1	VA	6181	6205	6203	69	0
1	VB	6181	6205	6203	65	0

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Mol	Chain	Non-H	H(model)	H(added)	Clashes	Symm-Clashes
1	W	6181	6205	6203	64	0
1	WA	6181	6205	6203	72	0
1	WB	6181	6205	6203	70	0
1	X	6181	6205	6203	70	0
1	XA	6181	6205	6203	73	0
1	XB	6181	6205	6203	76	0
1	Y	6181	6205	6203	67	0
1	YA	6181	6205	6203	75	0
1	YB	6181	6205	6203	77	0
1	Z	6181	6205	6203	70	0
1	ZA	6181	6205	6203	71	0
1	ZB	6181	6205	6203	67	0
All	All	482118	483990	483834	5156	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 5.

All (5156) close contacts within the same asymmetric unit are listed below, sorted by their clash magnitude.

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:O:168:ILE:HG22	1:O:215:LEU:HD21	1.64	0.80
1:MB:168:ILE:HG22	1:MB:215:LEU:HD21	1.64	0.79
1:Z:168:ILE:HG22	1:Z:215:LEU:HD21	1.64	0.79
1:T:168:ILE:HG22	1:T:215:LEU:HD21	1.64	0.79
1:VA:168:ILE:HG22	1:VA:215:LEU:HD21	1.63	0.79
1:I:168:ILE:HG22	1:I:215:LEU:HD21	1.65	0.79
1:H:168:ILE:HG22	1:H:215:LEU:HD21	1.65	0.78
1:JB:168:ILE:HG22	1:JB:215:LEU:HD21	1.64	0.78
1:W:168:ILE:HG22	1:W:215:LEU:HD21	1.66	0.78
1:CB:168:ILE:HG22	1:CB:215:LEU:HD21	1.67	0.77
1:E:48:MET:HA	1:E:48:MET:HE3	1.67	0.77
1:S:13:TYR:HA	1:S:50:MET:HE3	1.69	0.75
1:WB:13:TYR:HA	1:WB:50:MET:HE3	1.68	0.75
1:F:168:ILE:HG22	1:F:215:LEU:HD21	1.67	0.74
1:JA:13:TYR:HA	1:JA:50:MET:HE3	1.68	0.74
1:RB:539:LEU:HD11	1:RB:599:ILE:HD11	1.70	0.74
1:FB:168:ILE:HG22	1:FB:215:LEU:HD21	1.69	0.73
1:C:168:ILE:HG22	1:C:215:LEU:HD21	1.70	0.72
1:RA:168:ILE:HG22	1:RA:215:LEU:HD21	1.71	0.72
1:PA:168:ILE:HG22	1:PA:215:LEU:HD21	1.71	0.72
1:M:539:LEU:HD11	1:M:599:ILE:HD11	1.72	0.72

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:VB:168:ILE:HG22	1:VB:215:LEU:HD21	1.70	0.71
1:YB:168:ILE:HG22	1:YB:215:LEU:HD21	1.71	0.71
1:OB:168:ILE:HG22	1:OB:215:LEU:HD21	1.72	0.71
1:EA:539:LEU:HD11	1:EA:599:ILE:HD11	1.70	0.71
1:GB:168:ILE:HG22	1:GB:215:LEU:HD21	1.73	0.71
1:MA:168:ILE:HG22	1:MA:215:LEU:HD21	1.72	0.71
1:Y:168:ILE:HG22	1:Y:215:LEU:HD21	1.73	0.71
1:GA:168:ILE:HG22	1:GA:215:LEU:HD21	1.74	0.70
1:AB:539:LEU:HD11	1:AB:599:ILE:HD11	1.71	0.70
1:CA:168:ILE:HG22	1:CA:215:LEU:HD21	1.72	0.70
1:QA:168:ILE:HG22	1:QA:215:LEU:HD21	1.71	0.70
1:TA:168:ILE:HG22	1:TA:215:LEU:HD21	1.73	0.70
1:IA:168:ILE:HG22	1:IA:215:LEU:HD21	1.72	0.70
1:FA:168:ILE:HG22	1:FA:215:LEU:HD21	1.73	0.70
1:WA:168:ILE:HG22	1:WA:215:LEU:HD21	1.74	0.70
1:O:539:LEU:HD23	1:O:540:GLN:N	2.07	0.70
1:BA:168:ILE:HG22	1:BA:215:LEU:HD21	1.72	0.69
1:PB:168:ILE:HG22	1:PB:215:LEU:HD21	1.73	0.69
1:Y:539:LEU:HD23	1:Y:540:GLN:N	2.07	0.69
1:A:168:ILE:HG22	1:A:215:LEU:HD21	1.74	0.69
1:HB:168:ILE:HG22	1:HB:215:LEU:HD21	1.74	0.69
1:N:168:ILE:HG22	1:N:215:LEU:HD21	1.73	0.69
1:KB:168:ILE:HG22	1:KB:215:LEU:HD21	1.75	0.69
1:NA:168:ILE:HG22	1:NA:215:LEU:HD21	1.75	0.69
1:KA:168:ILE:HG22	1:KA:215:LEU:HD21	1.75	0.69
1:LA:168:ILE:HG22	1:LA:215:LEU:HD21	1.74	0.69
1:JA:168:ILE:HG22	1:JA:215:LEU:HD21	1.74	0.69
1:S:168:ILE:HG22	1:S:215:LEU:HD21	1.73	0.69
1:TB:168:ILE:HG22	1:TB:215:LEU:HD21	1.73	0.69
1:GA:13:TYR:HA	1:GA:50:MET:HE3	1.74	0.69
1:R:168:ILE:HG22	1:R:215:LEU:HD21	1.74	0.69
1:V:539:LEU:HD23	1:V:540:GLN:N	2.07	0.69
1:ZB:168:ILE:HG22	1:ZB:215:LEU:HD21	1.73	0.69
1:D:168:ILE:HG22	1:D:215:LEU:HD21	1.74	0.69
1:G:168:ILE:HG22	1:G:215:LEU:HD21	1.74	0.69
1:HA:168:ILE:HG22	1:HA:215:LEU:HD21	1.74	0.68
1:WB:168:ILE:HG22	1:WB:215:LEU:HD21	1.75	0.68
1:B:168:ILE:HG22	1:B:215:LEU:HD21	1.74	0.68
1:N:13:TYR:HA	1:N:50:MET:HE3	1.76	0.68
1:A:48:MET:HA	1:A:48:MET:HE3	1.75	0.68
1:SB:168:ILE:HG22	1:SB:215:LEU:HD21	1.74	0.68

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:BB:168:ILE:HG22	1:BB:215:LEU:HD21	1.75	0.68
1:Q:168:ILE:HG22	1:Q:215:LEU:HD21	1.75	0.68
1:TB:13:TYR:HA	1:TB:50:MET:HE3	1.75	0.68
1:X:168:ILE:HG22	1:X:215:LEU:HD21	1.74	0.68
1:QB:168:ILE:HG22	1:QB:215:LEU:HD21	1.74	0.68
1:UB:168:ILE:HG22	1:UB:215:LEU:HD21	1.76	0.68
1:D:48:MET:HA	1:D:48:MET:HE3	1.75	0.67
1:XB:168:ILE:HG22	1:XB:215:LEU:HD21	1.75	0.67
1:DB:168:ILE:HG22	1:DB:215:LEU:HD21	1.76	0.67
1:P:168:ILE:HG22	1:P:215:LEU:HD21	1.74	0.67
1:AC:168:ILE:HG22	1:AC:215:LEU:HD21	1.76	0.67
1:AB:30:VAL:CG2	1:AB:50:MET:HE3	2.25	0.67
1:D:30:VAL:HG21	1:D:50:MET:HE3	1.77	0.67
1:JB:48:MET:HA	1:JB:48:MET:HE3	1.77	0.67
1:ZA:168:ILE:HG22	1:ZA:215:LEU:HD21	1.77	0.67
1:UA:168:ILE:HG22	1:UA:215:LEU:HD21	1.75	0.67
1:V:168:ILE:HG22	1:V:215:LEU:HD21	1.75	0.66
1:AA:168:ILE:HG22	1:AA:215:LEU:HD21	1.76	0.66
1:EB:168:ILE:HG22	1:EB:215:LEU:HD21	1.76	0.66
1:J:168:ILE:HG22	1:J:215:LEU:HD21	1.77	0.66
1:AA:129:PHE:CZ	1:AA:137:VAL:HG21	2.31	0.66
1:E:168:ILE:HG22	1:E:215:LEU:HD21	1.77	0.66
1:LB:168:ILE:HG22	1:LB:215:LEU:HD21	1.77	0.66
1:M:168:ILE:HG22	1:M:215:LEU:HD21	1.76	0.66
1:XA:168:ILE:HG22	1:XA:215:LEU:HD21	1.76	0.66
1:CA:129:PHE:CZ	1:CA:137:VAL:HG21	2.31	0.66
1:EB:129:PHE:CZ	1:EB:137:VAL:HG21	2.31	0.66
1:QB:129:PHE:CZ	1:QB:137:VAL:HG21	2.31	0.66
1:TA:129:PHE:CZ	1:TA:137:VAL:HG21	2.31	0.66
1:YB:539:LEU:HD11	1:YB:599:ILE:HD11	1.78	0.66
1:DA:129:PHE:CZ	1:DA:137:VAL:HG21	2.31	0.66
1:Y:30:VAL:HG21	1:Y:50:MET:HE3	1.77	0.66
1:OB:129:PHE:CZ	1:OB:137:VAL:HG21	2.31	0.66
1:NB:129:PHE:CZ	1:NB:137:VAL:HG21	2.31	0.66
1:PA:129:PHE:CZ	1:PA:137:VAL:HG21	2.31	0.66
1:Q:129:PHE:CZ	1:Q:137:VAL:HG21	2.31	0.66
1:O:129:PHE:CZ	1:O:137:VAL:HG21	2.30	0.66
1:OA:168:ILE:HG22	1:OA:215:LEU:HD21	1.78	0.66
1:RB:168:ILE:HG22	1:RB:215:LEU:HD21	1.78	0.66
1:WA:30:VAL:HG21	1:WA:50:MET:HE3	1.78	0.66
1:AB:168:ILE:HG22	1:AB:215:LEU:HD21	1.78	0.65

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:C:129:PHE:CZ	1:C:137:VAL:HG21	2.31	0.65
1:CB:129:PHE:CZ	1:CB:137:VAL:HG21	2.31	0.65
1:DA:168:ILE:HG22	1:DA:215:LEU:HD21	1.77	0.65
1:W:129:PHE:CZ	1:W:137:VAL:HG21	2.31	0.65
1:XB:129:PHE:CZ	1:XB:137:VAL:HG21	2.31	0.65
1:B:129:PHE:CZ	1:B:137:VAL:HG21	2.31	0.65
1:DB:129:PHE:CZ	1:DB:137:VAL:HG21	2.32	0.65
1:KA:129:PHE:CZ	1:KA:137:VAL:HG21	2.32	0.65
1:QA:129:PHE:CZ	1:QA:137:VAL:HG21	2.31	0.65
1:W:30:VAL:HG21	1:W:50:MET:HE3	1.78	0.65
1:AB:30:VAL:HG21	1:AB:50:MET:HE3	1.78	0.65
1:EA:129:PHE:CZ	1:EA:137:VAL:HG21	2.31	0.65
1:G:129:PHE:CZ	1:G:137:VAL:HG21	2.31	0.65
1:YA:129:PHE:CZ	1:YA:137:VAL:HG21	2.32	0.65
1:BA:30:VAL:HG21	1:BA:50:MET:HE3	1.78	0.65
1:P:129:PHE:CZ	1:P:137:VAL:HG21	2.32	0.65
1:S:129:PHE:CZ	1:S:137:VAL:HG21	2.32	0.65
1:TB:48:MET:HA	1:TB:48:MET:HE3	1.79	0.65
1:UB:129:PHE:CZ	1:UB:137:VAL:HG21	2.32	0.65
1:CB:65:VAL:HG13	1:CB:103:GLU:OE2	1.97	0.65
1:HA:810:LEU:O	1:HA:810:LEU:HD23	1.97	0.65
1:SA:129:PHE:CZ	1:SA:137:VAL:HG21	2.31	0.65
1:UA:129:PHE:CZ	1:UA:137:VAL:HG21	2.32	0.65
1:E:129:PHE:CZ	1:E:137:VAL:HG21	2.32	0.65
1:HB:129:PHE:CZ	1:HB:137:VAL:HG21	2.32	0.65
1:L:810:LEU:O	1:L:810:LEU:HD23	1.97	0.65
1:NB:168:ILE:HG22	1:NB:215:LEU:HD21	1.79	0.65
1:UB:810:LEU:O	1:UB:810:LEU:HD23	1.97	0.65
1:ZB:129:PHE:CZ	1:ZB:137:VAL:HG21	2.32	0.65
1:A:129:PHE:CZ	1:A:137:VAL:HG21	2.32	0.65
1:JB:129:PHE:CZ	1:JB:137:VAL:HG21	2.32	0.65
1:K:168:ILE:HG22	1:K:215:LEU:HD21	1.78	0.65
1:MA:129:PHE:CZ	1:MA:137:VAL:HG21	2.32	0.65
1:ZA:129:PHE:CZ	1:ZA:137:VAL:HG21	2.32	0.65
1:ZB:65:VAL:HG13	1:ZB:103:GLU:OE2	1.97	0.65
1:GA:129:PHE:CZ	1:GA:137:VAL:HG21	2.32	0.65
1:L:129:PHE:CZ	1:L:137:VAL:HG21	2.32	0.65
1:LA:129:PHE:CZ	1:LA:137:VAL:HG21	2.32	0.65
1:PB:129:PHE:CZ	1:PB:137:VAL:HG21	2.31	0.65
1:Y:129:PHE:CZ	1:Y:137:VAL:HG21	2.32	0.65
1:B:810:LEU:O	1:B:810:LEU:HD23	1.96	0.65

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:BA:129:PHE:CZ	1:BA:137:VAL:HG21	2.32	0.65
1:HA:129:PHE:CZ	1:HA:137:VAL:HG21	2.32	0.65
1:SB:129:PHE:CZ	1:SB:137:VAL:HG21	2.32	0.65
1:D:30:VAL:CG2	1:D:50:MET:HE3	2.27	0.65
1:EA:168:ILE:HG22	1:EA:215:LEU:HD21	1.78	0.65
1:IA:129:PHE:CZ	1:IA:137:VAL:HG21	2.32	0.65
1:LB:129:PHE:CZ	1:LB:137:VAL:HG21	2.32	0.65
1:YB:129:PHE:CZ	1:YB:137:VAL:HG21	2.32	0.65
1:Z:129:PHE:CZ	1:Z:137:VAL:HG21	2.31	0.65
1:FA:129:PHE:CZ	1:FA:137:VAL:HG21	2.32	0.64
1:J:65:VAL:HG13	1:J:103:GLU:OE2	1.97	0.64
1:JA:129:PHE:CZ	1:JA:137:VAL:HG21	2.32	0.64
1:AB:129:PHE:CZ	1:AB:137:VAL:HG21	2.32	0.64
1:MB:129:PHE:CZ	1:MB:137:VAL:HG21	2.32	0.64
1:M:810:LEU:O	1:M:810:LEU:HD23	1.97	0.64
1:ZA:810:LEU:O	1:ZA:810:LEU:HD23	1.97	0.64
1:FB:129:PHE:CZ	1:FB:137:VAL:HG21	2.33	0.64
1:T:129:PHE:CZ	1:T:137:VAL:HG21	2.32	0.64
1:VB:129:PHE:CZ	1:VB:137:VAL:HG21	2.33	0.64
1:WB:129:PHE:CZ	1:WB:137:VAL:HG21	2.32	0.64
1:F:129:PHE:CZ	1:F:137:VAL:HG21	2.32	0.64
1:H:129:PHE:CZ	1:H:137:VAL:HG21	2.32	0.64
1:RB:129:PHE:CZ	1:RB:137:VAL:HG21	2.32	0.64
1:AC:129:PHE:CZ	1:AC:137:VAL:HG21	2.33	0.64
1:J:129:PHE:CZ	1:J:137:VAL:HG21	2.33	0.64
1:OA:129:PHE:CZ	1:OA:137:VAL:HG21	2.32	0.64
1:TB:129:PHE:CZ	1:TB:137:VAL:HG21	2.32	0.64
1:IB:129:PHE:CZ	1:IB:137:VAL:HG21	2.32	0.64
1:N:129:PHE:CZ	1:N:137:VAL:HG21	2.33	0.64
1:YA:30:VAL:HG21	1:YA:50:MET:HE3	1.79	0.64
1:D:129:PHE:CZ	1:D:137:VAL:HG21	2.33	0.64
1:EA:65:VAL:HG13	1:EA:103:GLU:OE2	1.98	0.64
1:FA:810:LEU:HD23	1:FA:810:LEU:O	1.98	0.64
1:KB:129:PHE:CZ	1:KB:137:VAL:HG21	2.33	0.64
1:UB:65:VAL:HG13	1:UB:103:GLU:OE2	1.98	0.64
1:V:65:VAL:HG13	1:V:103:GLU:OE2	1.98	0.64
1:WA:129:PHE:CZ	1:WA:137:VAL:HG21	2.33	0.64
1:GB:65:VAL:HG13	1:GB:103:GLU:OE2	1.98	0.64
1:GB:129:PHE:CZ	1:GB:137:VAL:HG21	2.33	0.64
1:HB:810:LEU:HD23	1:HB:810:LEU:O	1.98	0.64
1:MA:65:VAL:HG13	1:MA:103:GLU:OE2	1.98	0.64

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:QA:65:VAL:HG13	1:QA:103:GLU:OE2	1.98	0.64
1:VA:129:PHE:CZ	1:VA:137:VAL:HG21	2.32	0.64
1:XA:129:PHE:CZ	1:XA:137:VAL:HG21	2.32	0.64
1:YA:65:VAL:HG13	1:YA:103:GLU:OE2	1.98	0.64
1:BB:48:MET:HA	1:BB:48:MET:HE3	1.80	0.64
1:J:30:VAL:HG21	1:J:50:MET:HE3	1.79	0.64
1:NA:129:PHE:CZ	1:NA:137:VAL:HG21	2.33	0.64
1:R:129:PHE:CZ	1:R:137:VAL:HG21	2.32	0.64
1:RB:65:VAL:HG13	1:RB:103:GLU:OE2	1.97	0.64
1:T:810:LEU:O	1:T:810:LEU:HD23	1.98	0.64
1:W:810:LEU:O	1:W:810:LEU:HD23	1.98	0.64
1:WA:65:VAL:HG13	1:WA:103:GLU:OE2	1.98	0.64
1:D:810:LEU:O	1:D:810:LEU:HD23	1.98	0.63
1:J:30:VAL:CG2	1:J:50:MET:HE3	2.28	0.63
1:K:65:VAL:HG13	1:K:103:GLU:OE2	1.98	0.63
1:LA:65:VAL:HG13	1:LA:103:GLU:OE2	1.98	0.63
1:M:129:PHE:CZ	1:M:137:VAL:HG21	2.33	0.63
1:S:65:VAL:HG13	1:S:103:GLU:OE2	1.98	0.63
1:ZB:810:LEU:O	1:ZB:810:LEU:HD23	1.98	0.63
1:BA:65:VAL:HG13	1:BA:103:GLU:OE2	1.98	0.63
1:BB:129:PHE:CZ	1:BB:137:VAL:HG21	2.33	0.63
1:H:810:LEU:O	1:H:810:LEU:HD23	1.98	0.63
1:IA:810:LEU:HD23	1:IA:810:LEU:O	1.99	0.63
1:IB:65:VAL:HG13	1:IB:103:GLU:OE2	1.98	0.63
1:RA:810:LEU:O	1:RA:810:LEU:HD23	1.99	0.63
1:SB:810:LEU:HD23	1:SB:810:LEU:O	1.99	0.63
1:X:65:VAL:HG13	1:X:103:GLU:OE2	1.98	0.63
1:AB:810:LEU:O	1:AB:810:LEU:HD23	1.98	0.63
1:L:30:VAL:HG21	1:L:50:MET:HE3	1.81	0.63
1:P:30:VAL:HG21	1:P:50:MET:HE3	1.80	0.63
1:S:810:LEU:HD23	1:S:810:LEU:O	1.99	0.63
1:X:129:PHE:CZ	1:X:137:VAL:HG21	2.33	0.63
1:XA:65:VAL:HG13	1:XA:103:GLU:OE2	1.98	0.63
1:BB:810:LEU:O	1:BB:810:LEU:HD23	1.99	0.63
1:H:65:VAL:HG13	1:H:103:GLU:OE2	1.98	0.63
1:K:129:PHE:CZ	1:K:137:VAL:HG21	2.33	0.63
1:KA:810:LEU:O	1:KA:810:LEU:HD23	1.99	0.63
1:M:30:VAL:HG21	1:M:50:MET:HE3	1.81	0.63
1:OA:65:VAL:HG13	1:OA:103:GLU:OE2	1.99	0.63
1:OB:65:VAL:HG13	1:OB:103:GLU:OE2	1.98	0.63
1:V:129:PHE:CZ	1:V:137:VAL:HG21	2.33	0.63

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:EA:810:LEU:O	1:EA:810:LEU:HD23	1.98	0.63
1:EB:810:LEU:HD23	1:EB:810:LEU:O	1.99	0.63
1:FB:65:VAL:HG13	1:FB:103:GLU:OE2	1.99	0.63
1:KA:65:VAL:HG13	1:KA:103:GLU:OE2	1.99	0.63
1:KB:48:MET:HA	1:KB:48:MET:HE3	1.79	0.63
1:LA:810:LEU:HD23	1:LA:810:LEU:O	1.98	0.63
1:N:810:LEU:HD23	1:N:810:LEU:O	1.99	0.63
1:OA:810:LEU:HD23	1:OA:810:LEU:O	1.98	0.63
1:Q:810:LEU:HD23	1:Q:810:LEU:O	1.99	0.63
1:SA:168:ILE:HG22	1:SA:215:LEU:HD21	1.80	0.63
1:VB:810:LEU:HD23	1:VB:810:LEU:O	1.99	0.63
1:GB:810:LEU:O	1:GB:810:LEU:HD23	1.99	0.63
1:I:129:PHE:CZ	1:I:137:VAL:HG21	2.33	0.63
1:W:48:MET:HA	1:W:48:MET:HE3	1.80	0.63
1:WB:48:MET:HE3	1:WB:48:MET:HA	1.79	0.63
1:C:810:LEU:HD23	1:C:810:LEU:O	1.99	0.63
1:DA:65:VAL:HG13	1:DA:103:GLU:OE2	1.99	0.63
1:EA:48:MET:HE3	1:EA:48:MET:HA	1.81	0.63
1:KB:65:VAL:HG13	1:KB:103:GLU:OE2	1.98	0.63
1:L:168:ILE:HG22	1:L:215:LEU:HD21	1.80	0.63
1:PA:65:VAL:HG13	1:PA:103:GLU:OE2	1.98	0.63
1:QB:65:VAL:HG13	1:QB:103:GLU:OE2	1.99	0.63
1:RA:129:PHE:CZ	1:RA:137:VAL:HG21	2.34	0.63
1:TB:810:LEU:O	1:TB:810:LEU:HD23	1.99	0.63
1:V:810:LEU:HD23	1:V:810:LEU:O	1.99	0.63
1:VA:810:LEU:HD23	1:VA:810:LEU:O	1.98	0.63
1:Z:65:VAL:HG13	1:Z:103:GLU:OE2	1.99	0.63
1:A:810:LEU:HD23	1:A:810:LEU:O	1.99	0.63
1:NA:65:VAL:HG13	1:NA:103:GLU:OE2	1.98	0.63
1:SA:65:VAL:HG13	1:SA:103:GLU:OE2	1.99	0.63
1:EB:65:VAL:HG13	1:EB:103:GLU:OE2	1.98	0.63
1:I:65:VAL:HG13	1:I:103:GLU:OE2	1.98	0.63
1:D:65:VAL:HG13	1:D:103:GLU:OE2	1.99	0.62
1:M:65:VAL:HG13	1:M:103:GLU:OE2	1.99	0.62
1:T:65:VAL:HG13	1:T:103:GLU:OE2	1.99	0.62
1:C:65:VAL:HG13	1:C:103:GLU:OE2	1.98	0.62
1:E:30:VAL:HG21	1:E:50:MET:HE3	1.80	0.62
1:Q:65:VAL:HG13	1:Q:103:GLU:OE2	1.98	0.62
1:TA:810:LEU:HD23	1:TA:810:LEU:O	1.98	0.62
1:HB:65:VAL:HG13	1:HB:103:GLU:OE2	2.00	0.62
1:TA:30:VAL:CG2	1:TA:50:MET:HE3	2.29	0.62

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:XA:30:VAL:CG2	1:XA:50:MET:HE3	2.29	0.62
1:XB:65:VAL:HG13	1:XB:103:GLU:OE2	1.99	0.62
1:Y:30:VAL:CG2	1:Y:50:MET:HE3	2.29	0.62
1:CA:810:LEU:O	1:CA:810:LEU:HD23	2.00	0.62
1:VA:65:VAL:HG13	1:VA:103:GLU:OE2	1.99	0.62
1:XB:810:LEU:HD23	1:XB:810:LEU:O	2.00	0.62
1:AA:810:LEU:HD23	1:AA:810:LEU:O	2.00	0.62
1:R:65:VAL:HG13	1:R:103:GLU:OE2	2.00	0.62
1:RB:345:ASP:OD2	1:RB:349:VAL:HG13	2.00	0.62
1:CA:65:VAL:HG13	1:CA:103:GLU:OE2	1.99	0.62
1:G:65:VAL:HG13	1:G:103:GLU:OE2	1.99	0.62
1:IB:30:VAL:HG21	1:IB:50:MET:HE3	1.80	0.62
1:LB:48:MET:HE3	1:LB:48:MET:HA	1.80	0.62
1:LB:65:VAL:HG13	1:LB:103:GLU:OE2	2.00	0.62
1:NA:48:MET:HA	1:NA:48:MET:HE3	1.82	0.62
1:UA:65:VAL:HG13	1:UA:103:GLU:OE2	2.00	0.62
1:ZA:539:LEU:HD11	1:ZA:599:ILE:HD11	1.81	0.62
1:F:810:LEU:HD23	1:F:810:LEU:O	1.99	0.62
1:MA:810:LEU:O	1:MA:810:LEU:HD23	1.99	0.62
1:MB:65:VAL:HG13	1:MB:103:GLU:OE2	1.99	0.62
1:NB:65:VAL:HG13	1:NB:103:GLU:OE2	2.00	0.62
1:PB:65:VAL:HG13	1:PB:103:GLU:OE2	1.99	0.62
1:RA:65:VAL:HG13	1:RA:103:GLU:OE2	1.99	0.62
1:Y:539:LEU:HD11	1:Y:599:ILE:HD11	1.81	0.62
1:AC:65:VAL:HG13	1:AC:103:GLU:OE2	1.99	0.62
1:B:65:VAL:HG13	1:B:103:GLU:OE2	1.99	0.62
1:BB:65:VAL:HG13	1:BB:103:GLU:OE2	1.99	0.62
1:GA:810:LEU:HD23	1:GA:810:LEU:O	1.99	0.62
1:E:65:VAL:HG13	1:E:103:GLU:OE2	1.99	0.62
1:F:65:VAL:HG13	1:F:103:GLU:OE2	2.00	0.62
1:L:539:LEU:HD11	1:L:599:ILE:HD11	1.81	0.62
1:N:65:VAL:HG13	1:N:103:GLU:OE2	2.00	0.62
1:R:30:VAL:HG21	1:R:50:MET:HE3	1.82	0.62
1:E:810:LEU:O	1:E:810:LEU:HD23	2.00	0.62
1:G:48:MET:HA	1:G:48:MET:HE3	1.81	0.62
1:HA:65:VAL:HG13	1:HA:103:GLU:OE2	1.99	0.62
1:JA:810:LEU:O	1:JA:810:LEU:HD23	2.00	0.62
1:NB:810:LEU:HD23	1:NB:810:LEU:O	2.00	0.62
1:RB:48:MET:HA	1:RB:48:MET:HE3	1.81	0.62
1:Y:810:LEU:HD23	1:Y:810:LEU:O	2.00	0.62
1:PA:810:LEU:O	1:PA:810:LEU:HD23	2.00	0.61

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:Y:65:VAL:HG13	1:Y:103:GLU:OE2	1.99	0.61
1:DB:30:VAL:HG21	1:DB:50:MET:HE3	1.82	0.61
1:MB:30:VAL:HG21	1:MB:50:MET:HE3	1.81	0.61
1:IB:810:LEU:O	1:IB:810:LEU:HD23	1.99	0.61
1:TA:65:VAL:HG13	1:TA:103:GLU:OE2	2.00	0.61
1:WB:539:LEU:HD11	1:WB:599:ILE:HD11	1.82	0.61
1:XB:30:VAL:HG21	1:XB:50:MET:HE3	1.82	0.61
1:MA:48:MET:HE3	1:MA:48:MET:HA	1.82	0.61
1:O:30:VAL:HG21	1:O:50:MET:HE3	1.81	0.61
1:JA:48:MET:HA	1:JA:48:MET:HE3	1.81	0.61
1:WB:810:LEU:O	1:WB:810:LEU:HD23	2.00	0.61
1:RB:810:LEU:O	1:RB:810:LEU:HD23	1.99	0.61
1:YB:65:VAL:HG13	1:YB:103:GLU:OE2	2.00	0.61
1:KB:810:LEU:HD23	1:KB:810:LEU:O	2.01	0.61
1:L:65:VAL:HG13	1:L:103:GLU:OE2	2.00	0.61
1:LB:810:LEU:O	1:LB:810:LEU:HD23	2.00	0.61
1:FB:30:VAL:HG21	1:FB:50:MET:HE3	1.83	0.61
1:V:539:LEU:HD11	1:V:599:ILE:HD11	1.82	0.61
1:ZA:65:VAL:HG13	1:ZA:103:GLU:OE2	2.00	0.61
1:IB:168:ILE:HG22	1:IB:215:LEU:HD21	1.81	0.61
1:JA:9:ARG:HH12	1:KA:19:LEU:HD11	1.64	0.61
1:OB:810:LEU:O	1:OB:810:LEU:HD23	2.00	0.61
1:P:810:LEU:O	1:P:810:LEU:HD23	2.01	0.61
1:PB:810:LEU:HD23	1:PB:810:LEU:O	2.01	0.61
1:UA:810:LEU:O	1:UA:810:LEU:HD23	2.01	0.61
1:AC:810:LEU:O	1:AC:810:LEU:HD23	2.01	0.60
1:BA:810:LEU:O	1:BA:810:LEU:HD23	2.00	0.60
1:JA:539:LEU:HD11	1:JA:599:ILE:HD11	1.82	0.60
1:QB:810:LEU:O	1:QB:810:LEU:HD23	2.02	0.60
1:XA:30:VAL:HG21	1:XA:50:MET:HE3	1.81	0.60
1:MB:48:MET:HA	1:MB:48:MET:HE3	1.83	0.60
1:X:810:LEU:O	1:X:810:LEU:HD23	2.01	0.60
1:YB:810:LEU:O	1:YB:810:LEU:HD23	2.01	0.60
1:CB:810:LEU:HD23	1:CB:810:LEU:O	2.01	0.60
1:NA:810:LEU:HD23	1:NA:810:LEU:O	2.01	0.60
1:TA:30:VAL:HG21	1:TA:50:MET:HE3	1.81	0.60
1:DA:810:LEU:HD23	1:DA:810:LEU:O	2.02	0.60
1:K:30:VAL:HG21	1:K:50:MET:HE3	1.82	0.60
1:QA:810:LEU:HD23	1:QA:810:LEU:O	2.00	0.60
1:SA:810:LEU:O	1:SA:810:LEU:HD23	2.01	0.60
1:J:529:ILE:CD1	1:J:583:VAL:HG11	2.32	0.60

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:T:48:MET:HA	1:T:48:MET:HE3	1.82	0.60
1:XA:529:ILE:CD1	1:XA:583:VAL:HG11	2.32	0.60
1:AA:65:VAL:HG13	1:AA:103:GLU:OE2	2.01	0.59
1:M:529:ILE:CD1	1:M:583:VAL:HG11	2.32	0.59
1:O:65:VAL:HG13	1:O:103:GLU:OE2	2.02	0.59
1:UA:30:VAL:HG21	1:UA:50:MET:HE3	1.85	0.59
1:IA:529:ILE:CD1	1:IA:583:VAL:HG11	2.32	0.59
1:JA:529:ILE:CD1	1:JA:583:VAL:HG11	2.33	0.59
1:SA:529:ILE:CD1	1:SA:583:VAL:HG11	2.33	0.59
1:WA:529:ILE:CD1	1:WA:583:VAL:HG11	2.33	0.59
1:O:539:LEU:HD11	1:O:599:ILE:HD11	1.85	0.59
1:TB:575:ILE:HD12	1:TB:603:VAL:HG13	1.84	0.59
1:Y:575:ILE:HD12	1:Y:603:VAL:HG13	1.85	0.59
1:E:529:ILE:CD1	1:E:583:VAL:HG11	2.33	0.59
1:G:13:TYR:HA	1:G:50:MET:HE3	1.85	0.59
1:K:575:ILE:HD12	1:K:603:VAL:HG13	1.84	0.59
1:AC:529:ILE:HD13	1:AC:583:VAL:HG11	1.84	0.59
1:EB:30:VAL:HG21	1:EB:50:MET:HE3	1.84	0.59
1:VB:529:ILE:CD1	1:VB:583:VAL:HG11	2.33	0.59
1:WB:9:ARG:HH12	1:XB:19:LEU:HD11	1.67	0.59
1:X:539:LEU:HD11	1:X:599:ILE:HD11	1.83	0.59
1:AB:529:ILE:CD1	1:AB:583:VAL:HG11	2.32	0.59
1:D:529:ILE:CD1	1:D:583:VAL:HG11	2.33	0.59
1:KA:30:VAL:HG21	1:KA:50:MET:HE3	1.83	0.59
1:LA:529:ILE:CD1	1:LA:583:VAL:HG11	2.33	0.59
1:Q:30:VAL:HG21	1:Q:50:MET:HE3	1.85	0.59
1:XB:529:ILE:CD1	1:XB:583:VAL:HG11	2.32	0.59
1:A:529:ILE:CD1	1:A:583:VAL:HG11	2.32	0.59
1:BA:30:VAL:CG2	1:BA:50:MET:HE3	2.32	0.59
1:KA:529:ILE:CD1	1:KA:583:VAL:HG11	2.32	0.59
1:NA:529:ILE:HD13	1:NA:583:VAL:HG11	1.85	0.59
1:RA:529:ILE:CD1	1:RA:583:VAL:HG11	2.33	0.59
1:YB:529:ILE:CD1	1:YB:583:VAL:HG11	2.33	0.59
1:E:30:VAL:CG2	1:E:50:MET:HE3	2.33	0.59
1:OA:575:ILE:HD12	1:OA:603:VAL:HG13	1.85	0.59
1:UA:575:ILE:HD12	1:UA:603:VAL:HG13	1.85	0.59
1:W:529:ILE:CD1	1:W:583:VAL:HG11	2.33	0.58
1:Z:48:MET:HE3	1:Z:48:MET:HA	1.84	0.58
1:B:529:ILE:CD1	1:B:583:VAL:HG11	2.33	0.58
1:PB:30:VAL:HG21	1:PB:50:MET:HE3	1.86	0.58
1:PA:529:ILE:CD1	1:PA:583:VAL:HG11	2.33	0.58

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:ZA:575:ILE:HD12	1:ZA:603:VAL:HG13	1.84	0.58
1:O:575:ILE:HD12	1:O:603:VAL:HG13	1.85	0.58
1:OA:529:ILE:CD1	1:OA:583:VAL:HG11	2.32	0.58
1:SB:529:ILE:CD1	1:SB:583:VAL:HG11	2.34	0.58
1:DA:30:VAL:HG21	1:DA:50:MET:HE3	1.86	0.58
1:I:529:ILE:CD1	1:I:583:VAL:HG11	2.33	0.58
1:JB:529:ILE:CD1	1:JB:583:VAL:HG11	2.33	0.58
1:GA:48:MET:HA	1:GA:48:MET:HE3	1.85	0.58
1:T:575:ILE:HD12	1:T:603:VAL:HG13	1.85	0.58
1:UB:575:ILE:HD12	1:UB:603:VAL:HG13	1.85	0.58
1:B:575:ILE:HD12	1:B:603:VAL:HG13	1.84	0.58
1:CB:230:ARG:CB	1:CB:268:LEU:HD11	2.34	0.58
1:D:230:ARG:CB	1:D:268:LEU:HD11	2.34	0.58
1:EA:230:ARG:CB	1:EA:268:LEU:HD11	2.33	0.58
1:EB:765:VAL:HG22	1:FB:759:LEU:HD21	1.86	0.58
1:PA:575:ILE:HD12	1:PA:603:VAL:HG13	1.86	0.58
1:CA:575:ILE:HD12	1:CA:603:VAL:HG13	1.86	0.58
1:LB:575:ILE:HD12	1:LB:603:VAL:HG13	1.84	0.58
1:RB:230:ARG:CB	1:RB:268:LEU:HD11	2.34	0.58
1:XB:230:ARG:CB	1:XB:268:LEU:HD11	2.34	0.58
1:ZB:575:ILE:HD12	1:ZB:603:VAL:HG13	1.86	0.58
1:BA:230:ARG:CB	1:BA:268:LEU:HD11	2.34	0.58
1:FB:810:LEU:HD23	1:FB:810:LEU:O	2.04	0.58
1:M:230:ARG:CB	1:M:268:LEU:HD11	2.34	0.58
1:Q:529:ILE:CD1	1:Q:583:VAL:HG11	2.34	0.58
1:SA:575:ILE:HD12	1:SA:603:VAL:HG13	1.86	0.58
1:WB:529:ILE:CD1	1:WB:583:VAL:HG11	2.33	0.58
1:GB:230:ARG:CB	1:GB:268:LEU:HD11	2.34	0.58
1:L:575:ILE:HD12	1:L:603:VAL:HG13	1.84	0.58
1:QB:30:VAL:HG21	1:QB:50:MET:HE3	1.85	0.58
1:S:230:ARG:CB	1:S:268:LEU:HD11	2.34	0.58
1:TA:575:ILE:HD12	1:TA:603:VAL:HG13	1.86	0.58
1:WA:575:ILE:HD12	1:WA:603:VAL:HG13	1.86	0.58
1:YA:168:ILE:HG22	1:YA:215:LEU:HD21	1.85	0.58
1:EB:529:ILE:CD1	1:EB:583:VAL:HG11	2.34	0.57
1:FA:529:ILE:CD1	1:FA:583:VAL:HG11	2.34	0.57
1:GA:529:ILE:HD13	1:GA:583:VAL:HG11	1.85	0.57
1:NA:230:ARG:CB	1:NA:268:LEU:HD11	2.34	0.57
1:S:575:ILE:HD12	1:S:603:VAL:HG13	1.86	0.57
1:FA:230:ARG:CB	1:FA:268:LEU:HD11	2.34	0.57
1:I:575:ILE:HD12	1:I:603:VAL:HG13	1.86	0.57

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:KA:230:ARG:CB	1:KA:268:LEU:HD11	2.34	0.57
1:P:230:ARG:CB	1:P:268:LEU:HD11	2.34	0.57
1:R:575:ILE:HD12	1:R:603:VAL:HG13	1.87	0.57
1:V:48:MET:HA	1:V:48:MET:HE3	1.86	0.57
1:XA:523:PHE:CD2	1:XA:568:VAL:HG23	2.39	0.57
1:XB:575:ILE:HD12	1:XB:603:VAL:HG13	1.86	0.57
1:A:230:ARG:CB	1:A:268:LEU:HD11	2.34	0.57
1:L:529:ILE:CD1	1:L:583:VAL:HG11	2.35	0.57
1:SB:230:ARG:CB	1:SB:268:LEU:HD11	2.34	0.57
1:T:230:ARG:CB	1:T:268:LEU:HD11	2.34	0.57
1:Z:230:ARG:CB	1:Z:268:LEU:HD11	2.35	0.57
1:CA:230:ARG:CB	1:CA:268:LEU:HD11	2.34	0.57
1:DA:529:ILE:CD1	1:DA:583:VAL:HG11	2.35	0.57
1:O:230:ARG:CB	1:O:268:LEU:HD11	2.34	0.57
1:OA:230:ARG:CB	1:OA:268:LEU:HD11	2.34	0.57
1:OB:230:ARG:CB	1:OB:268:LEU:HD11	2.34	0.57
1:V:230:ARG:CB	1:V:268:LEU:HD11	2.35	0.57
1:HA:575:ILE:HD12	1:HA:603:VAL:HG13	1.85	0.57
1:J:523:PHE:CD2	1:J:568:VAL:HG23	2.39	0.57
1:NB:575:ILE:HD12	1:NB:603:VAL:HG13	1.86	0.57
1:P:65:VAL:HG13	1:P:103:GLU:OE2	2.04	0.57
1:V:529:ILE:CD1	1:V:583:VAL:HG11	2.34	0.57
1:WB:575:ILE:HD12	1:WB:603:VAL:HG13	1.87	0.57
1:AC:230:ARG:CB	1:AC:268:LEU:HD11	2.34	0.57
1:C:230:ARG:CB	1:C:268:LEU:HD11	2.35	0.57
1:DB:230:ARG:CB	1:DB:268:LEU:HD11	2.35	0.57
1:E:529:ILE:HD12	1:E:583:VAL:HG11	1.85	0.57
1:JA:575:ILE:HD12	1:JA:603:VAL:HG13	1.86	0.57
1:PB:230:ARG:CB	1:PB:268:LEU:HD11	2.35	0.57
1:QA:575:ILE:HD12	1:QA:603:VAL:HG13	1.87	0.57
1:SA:529:ILE:HD12	1:SA:583:VAL:HG11	1.86	0.57
1:AB:230:ARG:CB	1:AB:268:LEU:HD11	2.34	0.57
1:DA:230:ARG:CB	1:DA:268:LEU:HD11	2.35	0.57
1:DB:65:VAL:HG13	1:DB:103:GLU:OE2	2.04	0.57
1:EA:575:ILE:HD12	1:EA:603:VAL:HG13	1.87	0.57
1:MB:230:ARG:CB	1:MB:268:LEU:HD11	2.35	0.57
1:R:810:LEU:O	1:R:810:LEU:HD23	2.04	0.57
1:RA:230:ARG:CB	1:RA:268:LEU:HD11	2.34	0.57
1:VB:575:ILE:HD12	1:VB:603:VAL:HG13	1.86	0.57
1:CB:575:ILE:HD12	1:CB:603:VAL:HG13	1.87	0.57
1:GA:575:ILE:HD12	1:GA:603:VAL:HG13	1.86	0.57

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:IA:65:VAL:HG13	1:IA:103:GLU:OE2	2.05	0.57
1:JB:810:LEU:O	1:JB:810:LEU:HD23	2.05	0.57
1:L:230:ARG:CB	1:L:268:LEU:HD11	2.35	0.57
1:OB:529:ILE:CD1	1:OB:583:VAL:HG11	2.35	0.57
1:W:575:ILE:HD12	1:W:603:VAL:HG13	1.87	0.57
1:WA:810:LEU:HD23	1:WA:810:LEU:O	2.05	0.57
1:AA:575:ILE:HD12	1:AA:603:VAL:HG13	1.86	0.57
1:BA:529:ILE:CD1	1:BA:583:VAL:HG11	2.35	0.57
1:FA:575:ILE:HD12	1:FA:603:VAL:HG13	1.87	0.57
1:MB:529:ILE:CD1	1:MB:583:VAL:HG11	2.35	0.57
1:Q:230:ARG:CB	1:Q:268:LEU:HD11	2.35	0.57
1:ZA:230:ARG:CB	1:ZA:268:LEU:HD11	2.35	0.57
1:HB:230:ARG:CB	1:HB:268:LEU:HD11	2.35	0.57
1:VB:65:VAL:HG13	1:VB:103:GLU:OE2	2.05	0.57
1:YA:810:LEU:HD23	1:YA:810:LEU:O	2.05	0.57
1:A:575:ILE:HD12	1:A:603:VAL:HG13	1.86	0.56
1:BA:575:ILE:HD12	1:BA:603:VAL:HG13	1.87	0.56
1:DA:575:ILE:HD12	1:DA:603:VAL:HG13	1.87	0.56
1:F:230:ARG:CB	1:F:268:LEU:HD11	2.35	0.56
1:F:575:ILE:HD12	1:F:603:VAL:HG13	1.86	0.56
1:FA:65:VAL:HG13	1:FA:103:GLU:OE2	2.05	0.56
1:G:230:ARG:CB	1:G:268:LEU:HD11	2.35	0.56
1:HB:575:ILE:HD12	1:HB:603:VAL:HG13	1.86	0.56
1:KA:575:ILE:HD12	1:KA:603:VAL:HG13	1.86	0.56
1:R:529:ILE:CD1	1:R:583:VAL:HG11	2.35	0.56
1:VA:230:ARG:CB	1:VA:268:LEU:HD11	2.35	0.56
1:X:529:ILE:CD1	1:X:583:VAL:HG11	2.35	0.56
1:EA:529:ILE:CD1	1:EA:583:VAL:HG11	2.35	0.56
1:GA:230:ARG:CB	1:GA:268:LEU:HD11	2.35	0.56
1:HA:230:ARG:CB	1:HA:268:LEU:HD11	2.35	0.56
1:IA:575:ILE:HD12	1:IA:603:VAL:HG13	1.86	0.56
1:JA:65:VAL:HG13	1:JA:103:GLU:OE2	2.04	0.56
1:QB:230:ARG:CB	1:QB:268:LEU:HD11	2.35	0.56
1:VB:230:ARG:CB	1:VB:268:LEU:HD11	2.35	0.56
1:X:230:ARG:CB	1:X:268:LEU:HD11	2.36	0.56
1:ZA:529:ILE:CD1	1:ZA:583:VAL:HG11	2.34	0.56
1:AB:65:VAL:HG13	1:AB:103:GLU:OE2	2.06	0.56
1:DB:529:ILE:HD13	1:DB:583:VAL:HG11	1.87	0.56
1:FB:230:ARG:CB	1:FB:268:LEU:HD11	2.36	0.56
1:FB:529:ILE:CD1	1:FB:583:VAL:HG11	2.35	0.56
1:G:575:ILE:HD12	1:G:603:VAL:HG13	1.86	0.56

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:G:810:LEU:O	1:G:810:LEU:HD23	2.06	0.56
1:H:230:ARG:CB	1:H:268:LEU:HD11	2.35	0.56
1:HB:529:ILE:CD1	1:HB:583:VAL:HG11	2.35	0.56
1:I:230:ARG:CB	1:I:268:LEU:HD11	2.35	0.56
1:IB:529:ILE:CD1	1:IB:583:VAL:HG11	2.35	0.56
1:P:575:ILE:HD12	1:P:603:VAL:HG13	1.87	0.56
1:QA:230:ARG:CB	1:QA:268:LEU:HD11	2.35	0.56
1:R:230:ARG:CB	1:R:268:LEU:HD11	2.36	0.56
1:UB:230:ARG:CB	1:UB:268:LEU:HD11	2.35	0.56
1:W:65:VAL:HG13	1:W:103:GLU:OE2	2.05	0.56
1:W:230:ARG:CB	1:W:268:LEU:HD11	2.36	0.56
1:WB:65:VAL:HG13	1:WB:103:GLU:OE2	2.05	0.56
1:YA:575:ILE:HD12	1:YA:603:VAL:HG13	1.87	0.56
1:YB:230:ARG:CB	1:YB:268:LEU:HD11	2.35	0.56
1:YB:575:ILE:HD12	1:YB:603:VAL:HG13	1.87	0.56
1:Z:529:ILE:CD1	1:Z:583:VAL:HG11	2.35	0.56
1:A:83:LEU:HD21	1:A:102:GLY:O	2.06	0.56
1:DB:575:ILE:HD12	1:DB:603:VAL:HG13	1.88	0.56
1:IB:230:ARG:CB	1:IB:268:LEU:HD11	2.35	0.56
1:J:230:ARG:CB	1:J:268:LEU:HD11	2.36	0.56
1:JB:65:VAL:HG13	1:JB:103:GLU:OE2	2.06	0.56
1:MA:230:ARG:CB	1:MA:268:LEU:HD11	2.35	0.56
1:Q:575:ILE:HD12	1:Q:603:VAL:HG13	1.87	0.56
1:AA:230:ARG:CB	1:AA:268:LEU:HD11	2.36	0.56
1:DB:810:LEU:O	1:DB:810:LEU:HD23	2.05	0.56
1:IA:230:ARG:CB	1:IA:268:LEU:HD11	2.35	0.56
1:LA:230:ARG:CB	1:LA:268:LEU:HD11	2.35	0.56
1:MB:810:LEU:HD23	1:MB:810:LEU:O	2.05	0.56
1:N:529:ILE:CD1	1:N:583:VAL:HG11	2.35	0.56
1:QB:529:ILE:CD1	1:QB:583:VAL:HG11	2.35	0.56
1:RB:575:ILE:HD12	1:RB:603:VAL:HG13	1.87	0.56
1:SB:65:VAL:HG13	1:SB:103:GLU:OE2	2.06	0.56
1:Z:575:ILE:HD12	1:Z:603:VAL:HG13	1.88	0.56
1:ZB:529:ILE:CD1	1:ZB:583:VAL:HG11	2.36	0.56
1:C:529:ILE:HD13	1:C:583:VAL:HG11	1.87	0.56
1:EB:575:ILE:HD12	1:EB:603:VAL:HG13	1.87	0.56
1:FB:575:ILE:HD12	1:FB:603:VAL:HG13	1.87	0.56
1:I:810:LEU:HD23	1:I:810:LEU:O	2.05	0.56
1:N:230:ARG:CB	1:N:268:LEU:HD11	2.36	0.56
1:RB:529:ILE:CD1	1:RB:583:VAL:HG11	2.35	0.56
1:Y:230:ARG:CB	1:Y:268:LEU:HD11	2.36	0.56

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:CA:30:VAL:HG21	1:CA:50:MET:HE3	1.88	0.56
1:E:230:ARG:CB	1:E:268:LEU:HD11	2.36	0.56
1:IB:30:VAL:CG2	1:IB:50:MET:HE3	2.36	0.56
1:JA:230:ARG:CB	1:JA:268:LEU:HD11	2.36	0.56
1:LA:48:MET:HE3	1:LA:49:ARG:H	1.71	0.56
1:LB:230:ARG:CB	1:LB:268:LEU:HD11	2.35	0.56
1:NB:230:ARG:CB	1:NB:268:LEU:HD11	2.36	0.56
1:O:83:LEU:HD21	1:O:102:GLY:O	2.06	0.56
1:RA:575:ILE:HD12	1:RA:603:VAL:HG13	1.87	0.56
1:SB:575:ILE:HD12	1:SB:603:VAL:HG13	1.87	0.56
1:UA:230:ARG:CB	1:UA:268:LEU:HD11	2.36	0.56
1:Y:529:ILE:CD1	1:Y:583:VAL:HG11	2.36	0.56
1:YA:30:VAL:CG2	1:YA:50:MET:HE3	2.35	0.56
1:FA:83:LEU:HD21	1:FA:102:GLY:O	2.06	0.56
1:JB:230:ARG:CB	1:JB:268:LEU:HD11	2.36	0.56
1:VB:83:LEU:HD21	1:VB:102:GLY:O	2.06	0.56
1:WA:230:ARG:CB	1:WA:268:LEU:HD11	2.35	0.56
1:AC:575:ILE:HD12	1:AC:603:VAL:HG13	1.88	0.56
1:E:575:ILE:HD12	1:E:603:VAL:HG13	1.86	0.56
1:JA:83:LEU:HD21	1:JA:102:GLY:O	2.06	0.56
1:P:30:VAL:CG2	1:P:50:MET:HE3	2.36	0.56
1:V:523:PHE:CD2	1:V:568:VAL:HG23	2.41	0.56
1:VA:575:ILE:HD12	1:VA:603:VAL:HG13	1.88	0.56
1:XA:575:ILE:HD12	1:XA:603:VAL:HG13	1.88	0.56
1:EB:230:ARG:CB	1:EB:268:LEU:HD11	2.35	0.56
1:IB:575:ILE:HD12	1:IB:603:VAL:HG13	1.87	0.56
1:KB:230:ARG:CB	1:KB:268:LEU:HD11	2.36	0.56
1:KB:529:ILE:CD1	1:KB:583:VAL:HG11	2.36	0.56
1:PB:575:ILE:HD12	1:PB:603:VAL:HG13	1.86	0.56
1:S:529:ILE:HD12	1:S:583:VAL:HG11	1.88	0.56
1:XA:230:ARG:CB	1:XA:268:LEU:HD11	2.36	0.56
1:B:230:ARG:CB	1:B:268:LEU:HD11	2.35	0.55
1:BB:529:ILE:CD1	1:BB:583:VAL:HG11	2.35	0.55
1:GB:575:ILE:HD12	1:GB:603:VAL:HG13	1.86	0.55
1:JB:83:LEU:HD21	1:JB:102:GLY:O	2.06	0.55
1:K:810:LEU:HD23	1:K:810:LEU:O	2.06	0.55
1:O:168:ILE:HG22	1:O:215:LEU:CD2	2.35	0.55
1:SA:230:ARG:CB	1:SA:268:LEU:HD11	2.36	0.55
1:T:529:ILE:CD1	1:T:583:VAL:HG11	2.35	0.55
1:TA:529:ILE:CD1	1:TA:583:VAL:HG11	2.36	0.55
1:A:65:VAL:HG13	1:A:103:GLU:OE2	2.07	0.55

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:BB:575:ILE:HD12	1:BB:603:VAL:HG13	1.89	0.55
1:F:529:ILE:CD1	1:F:583:VAL:HG11	2.36	0.55
1:H:575:ILE:HD12	1:H:603:VAL:HG13	1.88	0.55
1:IA:83:LEU:HD21	1:IA:102:GLY:O	2.06	0.55
1:MA:529:ILE:CD1	1:MA:583:VAL:HG11	2.36	0.55
1:MA:575:ILE:HD12	1:MA:603:VAL:HG13	1.87	0.55
1:NA:575:ILE:HD12	1:NA:603:VAL:HG13	1.88	0.55
1:PA:230:ARG:CB	1:PA:268:LEU:HD11	2.36	0.55
1:QB:575:ILE:HD12	1:QB:603:VAL:HG13	1.87	0.55
1:TA:65:VAL:HG11	1:TA:100:TYR:HB2	1.87	0.55
1:TA:230:ARG:CB	1:TA:268:LEU:HD11	2.35	0.55
1:XA:810:LEU:O	1:XA:810:LEU:HD23	2.06	0.55
1:Z:810:LEU:O	1:Z:810:LEU:HD23	2.06	0.55
1:IB:529:ILE:HD12	1:IB:583:VAL:HG11	1.88	0.55
1:V:575:ILE:HD12	1:V:603:VAL:HG13	1.88	0.55
1:VA:168:ILE:HG22	1:VA:215:LEU:CD2	2.36	0.55
1:C:575:ILE:HD12	1:C:603:VAL:HG13	1.87	0.55
1:G:529:ILE:CD1	1:G:583:VAL:HG11	2.36	0.55
1:IB:523:PHE:CD2	1:IB:568:VAL:HG23	2.41	0.55
1:JB:168:ILE:HG22	1:JB:215:LEU:CD2	2.35	0.55
1:OB:575:ILE:HD12	1:OB:603:VAL:HG13	1.87	0.55
1:UA:529:ILE:CD1	1:UA:583:VAL:HG11	2.36	0.55
1:Z:523:PHE:CD2	1:Z:568:VAL:HG23	2.42	0.55
1:DB:30:VAL:CG2	1:DB:50:MET:HE3	2.37	0.55
1:K:230:ARG:CB	1:K:268:LEU:HD11	2.37	0.55
1:P:529:ILE:HD13	1:P:583:VAL:HG11	1.87	0.55
1:QA:65:VAL:HG11	1:QA:100:TYR:HB2	1.88	0.55
1:S:529:ILE:CD1	1:S:583:VAL:HG11	2.37	0.55
1:X:575:ILE:HD12	1:X:603:VAL:HG13	1.89	0.55
1:BB:230:ARG:CB	1:BB:268:LEU:HD11	2.36	0.55
1:H:523:PHE:CD2	1:H:568:VAL:HG23	2.41	0.55
1:HB:230:ARG:HB2	1:HB:268:LEU:HD11	1.89	0.55
1:MB:523:PHE:CD2	1:MB:568:VAL:HG23	2.42	0.55
1:N:575:ILE:HD12	1:N:603:VAL:HG13	1.89	0.55
1:O:810:LEU:O	1:O:810:LEU:HD23	2.06	0.55
1:W:345:ASP:OD2	1:W:349:VAL:HG13	2.07	0.55
1:GB:529:ILE:CD1	1:GB:583:VAL:HG11	2.37	0.55
1:J:810:LEU:O	1:J:810:LEU:HD23	2.06	0.55
1:LA:575:ILE:HD12	1:LA:603:VAL:HG13	1.87	0.55
1:VA:523:PHE:CD2	1:VA:568:VAL:HG23	2.42	0.55
1:AB:83:LEU:HD21	1:AB:102:GLY:O	2.07	0.55

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:CA:65:VAL:HG11	1:CA:100:TYR:HB2	1.88	0.55
1:J:575:ILE:HD12	1:J:603:VAL:HG13	1.88	0.55
1:OB:48:MET:HE3	1:OB:48:MET:HA	1.88	0.55
1:TB:83:LEU:HD21	1:TB:102:GLY:O	2.07	0.55
1:TB:230:ARG:CB	1:TB:268:LEU:HD11	2.36	0.55
1:X:529:ILE:HD12	1:X:583:VAL:HG11	1.89	0.55
1:GA:65:VAL:HG13	1:GA:103:GLU:OE2	2.07	0.55
1:GA:83:LEU:HD21	1:GA:102:GLY:O	2.07	0.55
1:YA:230:ARG:CB	1:YA:268:LEU:HD11	2.37	0.55
1:A:523:PHE:CD2	1:A:568:VAL:HG23	2.41	0.55
1:D:230:ARG:HB2	1:D:268:LEU:HD11	1.89	0.55
1:EA:271:VAL:HG23	1:EA:271:VAL:O	2.07	0.55
1:NB:529:ILE:CD1	1:NB:583:VAL:HG11	2.37	0.55
1:TB:529:ILE:CD1	1:TB:583:VAL:HG11	2.37	0.55
1:VA:529:ILE:CD1	1:VA:583:VAL:HG11	2.36	0.55
1:XA:48:MET:HE3	1:XA:48:MET:HA	1.89	0.55
1:BB:30:VAL:HG21	1:BB:50:MET:HE3	1.89	0.54
1:C:65:VAL:HG11	1:C:100:TYR:HB2	1.88	0.54
1:D:575:ILE:HD12	1:D:603:VAL:HG13	1.88	0.54
1:JB:271:VAL:O	1:JB:271:VAL:HG23	2.07	0.54
1:T:230:ARG:HB2	1:T:268:LEU:HD11	1.89	0.54
1:WB:83:LEU:HD21	1:WB:102:GLY:O	2.07	0.54
1:YA:529:ILE:HD13	1:YA:583:VAL:HG11	1.88	0.54
1:GB:529:ILE:HD12	1:GB:583:VAL:HG11	1.88	0.54
1:K:529:ILE:HD13	1:K:583:VAL:HG11	1.88	0.54
1:LA:381:PRO:HA	1:LA:405:THR:HG22	1.90	0.54
1:PB:65:VAL:HG11	1:PB:100:TYR:HB2	1.88	0.54
1:RA:230:ARG:HB2	1:RA:268:LEU:HD11	1.89	0.54
1:Z:168:ILE:HG22	1:Z:215:LEU:CD2	2.36	0.54
1:AB:381:PRO:HA	1:AB:405:THR:HG22	1.90	0.54
1:CB:271:VAL:HG23	1:CB:271:VAL:O	2.08	0.54
1:MB:65:VAL:HG11	1:MB:100:TYR:HB2	1.88	0.54
1:MB:575:ILE:HD12	1:MB:603:VAL:HG13	1.88	0.54
1:TA:523:PHE:CD2	1:TA:568:VAL:HG23	2.42	0.54
1:WB:230:ARG:CB	1:WB:268:LEU:HD11	2.36	0.54
1:Z:381:PRO:HA	1:Z:405:THR:HG22	1.90	0.54
1:ZB:230:ARG:CB	1:ZB:268:LEU:HD11	2.36	0.54
1:CB:529:ILE:CD1	1:CB:583:VAL:HG11	2.37	0.54
1:DB:271:VAL:HG23	1:DB:271:VAL:O	2.08	0.54
1:JB:737:GLY:O	1:JB:741:VAL:HG23	2.07	0.54
1:Q:765:VAL:HG22	1:R:759:LEU:HD21	1.89	0.54

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:RB:529:ILE:HD12	1:RB:583:VAL:HG11	1.90	0.54
1:VB:271:VAL:HG23	1:VB:271:VAL:O	2.08	0.54
1:XA:65:VAL:HG11	1:XA:100:TYR:HB2	1.89	0.54
1:CB:230:ARG:HB2	1:CB:268:LEU:HD11	1.90	0.54
1:EB:65:VAL:HG11	1:EB:100:TYR:HB2	1.89	0.54
1:F:271:VAL:HG23	1:F:271:VAL:O	2.08	0.54
1:G:65:VAL:HG11	1:G:100:TYR:HB2	1.88	0.54
1:KA:230:ARG:HB2	1:KA:268:LEU:HD11	1.90	0.54
1:KB:529:ILE:HD12	1:KB:583:VAL:HG11	1.89	0.54
1:LB:271:VAL:HG23	1:LB:271:VAL:O	2.08	0.54
1:LB:529:ILE:CD1	1:LB:583:VAL:HG11	2.37	0.54
1:OA:523:PHE:CD2	1:OA:568:VAL:HG23	2.43	0.54
1:OB:381:PRO:HA	1:OB:405:THR:HG22	1.89	0.54
1:RB:271:VAL:O	1:RB:271:VAL:HG23	2.08	0.54
1:W:271:VAL:HG23	1:W:271:VAL:O	2.07	0.54
1:BA:381:PRO:HA	1:BA:405:THR:HG22	1.90	0.54
1:CA:271:VAL:HG23	1:CA:271:VAL:O	2.07	0.54
1:G:271:VAL:HG23	1:G:271:VAL:O	2.08	0.54
1:JB:575:ILE:HD12	1:JB:603:VAL:HG13	1.88	0.54
1:NA:271:VAL:HG23	1:NA:271:VAL:O	2.08	0.54
1:OB:13:TYR:HA	1:OB:50:MET:HE3	1.89	0.54
1:W:83:LEU:HD21	1:W:102:GLY:O	2.07	0.54
1:ZB:523:PHE:CD2	1:ZB:568:VAL:HG23	2.43	0.54
1:AB:230:ARG:HB2	1:AB:268:LEU:HD11	1.90	0.54
1:BA:65:VAL:HG11	1:BA:100:TYR:HB2	1.89	0.54
1:BB:144:LEU:HD12	1:BB:144:LEU:O	2.07	0.54
1:JA:271:VAL:HG23	1:JA:271:VAL:O	2.08	0.54
1:JB:65:VAL:HG11	1:JB:100:TYR:HB2	1.90	0.54
1:KA:271:VAL:O	1:KA:271:VAL:HG23	2.08	0.54
1:KB:271:VAL:HG23	1:KB:271:VAL:O	2.08	0.54
1:MA:271:VAL:HG23	1:MA:271:VAL:O	2.08	0.54
1:OB:65:VAL:HG11	1:OB:100:TYR:HB2	1.89	0.54
1:SB:302:VAL:HG11	1:SB:306:LYS:HZ2	1.73	0.54
1:UA:65:VAL:HG11	1:UA:100:TYR:HB2	1.88	0.54
1:YA:271:VAL:HG23	1:YA:271:VAL:O	2.08	0.54
1:AA:381:PRO:HA	1:AA:405:THR:HG22	1.90	0.54
1:BB:271:VAL:HG23	1:BB:271:VAL:O	2.08	0.54
1:D:65:VAL:HG11	1:D:100:TYR:HB2	1.90	0.54
1:E:271:VAL:O	1:E:271:VAL:HG23	2.08	0.54
1:EB:271:VAL:HG23	1:EB:271:VAL:O	2.08	0.54
1:HA:230:ARG:HB2	1:HA:268:LEU:HD11	1.90	0.54

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:IA:271:VAL:HG23	1:IA:271:VAL:O	2.08	0.54
1:KB:575:ILE:HD12	1:KB:603:VAL:HG13	1.89	0.54
1:LA:271:VAL:HG23	1:LA:271:VAL:O	2.08	0.54
1:NB:271:VAL:HG23	1:NB:271:VAL:O	2.08	0.54
1:AA:65:VAL:HG11	1:AA:100:TYR:HB2	1.89	0.54
1:BA:271:VAL:HG23	1:BA:271:VAL:O	2.08	0.54
1:C:230:ARG:HB2	1:C:268:LEU:HD11	1.90	0.54
1:DA:271:VAL:HG23	1:DA:271:VAL:O	2.08	0.54
1:DA:381:PRO:HA	1:DA:405:THR:HG22	1.90	0.54
1:DB:83:LEU:HD21	1:DB:102:GLY:O	2.07	0.54
1:FB:230:ARG:HB2	1:FB:268:LEU:HD11	1.90	0.54
1:FB:271:VAL:HG23	1:FB:271:VAL:O	2.08	0.54
1:GA:523:PHE:CD2	1:GA:568:VAL:HG23	2.43	0.54
1:GB:230:ARG:HB2	1:GB:268:LEU:HD11	1.90	0.54
1:IA:381:PRO:HA	1:IA:405:THR:HG22	1.90	0.54
1:M:30:VAL:CG2	1:M:50:MET:HE3	2.38	0.54
1:M:381:PRO:HA	1:M:405:THR:HG22	1.90	0.54
1:N:144:LEU:HD12	1:N:144:LEU:O	2.07	0.54
1:SB:83:LEU:HD21	1:SB:102:GLY:O	2.07	0.54
1:TB:271:VAL:HG23	1:TB:271:VAL:O	2.08	0.54
1:YB:381:PRO:HA	1:YB:405:THR:HG22	1.90	0.54
1:BB:529:ILE:HD12	1:BB:583:VAL:HG11	1.90	0.54
1:F:230:ARG:HB2	1:F:268:LEU:HD11	1.90	0.54
1:FA:271:VAL:HG23	1:FA:271:VAL:O	2.08	0.54
1:H:168:ILE:HG22	1:H:215:LEU:CD2	2.36	0.54
1:IB:65:VAL:HG11	1:IB:100:TYR:HB2	1.90	0.54
1:K:271:VAL:HG23	1:K:271:VAL:O	2.08	0.54
1:MB:271:VAL:HG23	1:MB:271:VAL:O	2.08	0.54
1:O:271:VAL:HG23	1:O:271:VAL:O	2.08	0.54
1:PB:271:VAL:HG23	1:PB:271:VAL:O	2.08	0.54
1:SB:30:VAL:HG21	1:SB:50:MET:HE3	1.89	0.54
1:TA:812:VAL:O	1:TA:812:VAL:HG23	2.08	0.54
1:ZA:381:PRO:HA	1:ZA:405:THR:HG22	1.90	0.54
1:CB:381:PRO:HA	1:CB:405:THR:HG22	1.91	0.53
1:D:271:VAL:HG23	1:D:271:VAL:O	2.08	0.53
1:EA:65:VAL:HG11	1:EA:100:TYR:HB2	1.90	0.53
1:HA:271:VAL:O	1:HA:271:VAL:HG23	2.07	0.53
1:N:381:PRO:HA	1:N:405:THR:HG22	1.90	0.53
1:AB:575:ILE:HD12	1:AB:603:VAL:HG13	1.90	0.53
1:B:812:VAL:HG23	1:B:812:VAL:O	2.09	0.53
1:C:271:VAL:HG23	1:C:271:VAL:O	2.08	0.53

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:EB:230:ARG:HB2	1:EB:268:LEU:HD11	1.90	0.53
1:EB:381:PRO:HA	1:EB:405:THR:HG22	1.90	0.53
1:EB:812:VAL:HG23	1:EB:812:VAL:O	2.09	0.53
1:FB:381:PRO:HA	1:FB:405:THR:HG22	1.90	0.53
1:H:65:VAL:HG11	1:H:100:TYR:HB2	1.89	0.53
1:I:765:VAL:HG22	1:J:759:LEU:HD21	1.91	0.53
1:IA:812:VAL:HG23	1:IA:812:VAL:O	2.09	0.53
1:MB:168:ILE:HG22	1:MB:215:LEU:CD2	2.35	0.53
1:MB:230:ARG:HB2	1:MB:268:LEU:HD11	1.90	0.53
1:NA:812:VAL:HG23	1:NA:812:VAL:O	2.09	0.53
1:O:230:ARG:HB2	1:O:268:LEU:HD11	1.90	0.53
1:P:83:LEU:HD21	1:P:102:GLY:O	2.07	0.53
1:R:812:VAL:HG23	1:R:812:VAL:O	2.08	0.53
1:UB:230:ARG:HB2	1:UB:268:LEU:HD11	1.90	0.53
1:UB:271:VAL:HG23	1:UB:271:VAL:O	2.08	0.53
1:WA:381:PRO:HA	1:WA:405:THR:HG22	1.90	0.53
1:WA:737:GLY:O	1:WA:741:VAL:HG23	2.08	0.53
1:X:538:GLN:HB2	1:X:646:VAL:HG22	1.90	0.53
1:YA:381:PRO:HA	1:YA:405:THR:HG22	1.91	0.53
1:BB:381:PRO:HA	1:BB:405:THR:HG22	1.90	0.53
1:FB:65:VAL:HG11	1:FB:100:TYR:HB2	1.90	0.53
1:I:812:VAL:O	1:I:812:VAL:HG23	2.09	0.53
1:IB:271:VAL:HG23	1:IB:271:VAL:O	2.08	0.53
1:JB:230:ARG:HB2	1:JB:268:LEU:HD11	1.91	0.53
1:L:325:VAL:HG13	1:L:325:VAL:O	2.07	0.53
1:LA:230:ARG:HB2	1:LA:268:LEU:HD11	1.91	0.53
1:M:575:ILE:HD12	1:M:603:VAL:HG13	1.90	0.53
1:MA:65:VAL:HG11	1:MA:100:TYR:HB2	1.91	0.53
1:MA:381:PRO:HA	1:MA:405:THR:HG22	1.90	0.53
1:O:529:ILE:CD1	1:O:583:VAL:HG11	2.38	0.53
1:OB:230:ARG:HB2	1:OB:268:LEU:HD11	1.91	0.53
1:QB:271:VAL:HG23	1:QB:271:VAL:O	2.08	0.53
1:SB:271:VAL:HG23	1:SB:271:VAL:O	2.08	0.53
1:UA:381:PRO:HA	1:UA:405:THR:HG22	1.91	0.53
1:W:65:VAL:HG11	1:W:100:TYR:HB2	1.90	0.53
1:WA:271:VAL:HG23	1:WA:271:VAL:O	2.08	0.53
1:XA:812:VAL:HG23	1:XA:812:VAL:O	2.08	0.53
1:YA:65:VAL:HG11	1:YA:100:TYR:HB2	1.90	0.53
1:AC:230:ARG:HB2	1:AC:268:LEU:HD11	1.90	0.53
1:CB:398:VAL:HG11	1:CB:415:TRP:CE3	2.44	0.53
1:GA:271:VAL:HG23	1:GA:271:VAL:O	2.08	0.53

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:MA:523:PHE:CD2	1:MA:568:VAL:HG23	2.43	0.53
1:OA:812:VAL:O	1:OA:812:VAL:HG23	2.09	0.53
1:PB:230:ARG:HB2	1:PB:268:LEU:HD11	1.91	0.53
1:SA:381:PRO:HA	1:SA:405:THR:HG22	1.91	0.53
1:VA:271:VAL:HG23	1:VA:271:VAL:O	2.08	0.53
1:W:168:ILE:HG22	1:W:215:LEU:CD2	2.36	0.53
1:WB:523:PHE:CD2	1:WB:568:VAL:HG23	2.43	0.53
1:ZB:271:VAL:O	1:ZB:271:VAL:HG23	2.08	0.53
1:AB:529:ILE:HD12	1:AB:583:VAL:HG11	1.90	0.53
1:AC:381:PRO:HA	1:AC:405:THR:HG22	1.91	0.53
1:B:230:ARG:HB2	1:B:268:LEU:HD11	1.91	0.53
1:D:812:VAL:HG23	1:D:812:VAL:O	2.09	0.53
1:DB:230:ARG:HB2	1:DB:268:LEU:HD11	1.91	0.53
1:DB:381:PRO:HA	1:DB:405:THR:HG22	1.91	0.53
1:FA:230:ARG:HB2	1:FA:268:LEU:HD11	1.91	0.53
1:IA:737:GLY:O	1:IA:741:VAL:HG23	2.08	0.53
1:L:812:VAL:HG23	1:L:812:VAL:O	2.09	0.53
1:N:271:VAL:HG23	1:N:271:VAL:O	2.08	0.53
1:NA:230:ARG:HB2	1:NA:268:LEU:HD11	1.91	0.53
1:QB:812:VAL:HG23	1:QB:812:VAL:O	2.09	0.53
1:RB:230:ARG:HB2	1:RB:268:LEU:HD11	1.90	0.53
1:TA:271:VAL:HG23	1:TA:271:VAL:O	2.08	0.53
1:VA:381:PRO:HA	1:VA:405:THR:HG22	1.91	0.53
1:VB:737:GLY:O	1:VB:741:VAL:HG23	2.08	0.53
1:W:381:PRO:HA	1:W:405:THR:HG22	1.91	0.53
1:WA:812:VAL:O	1:WA:812:VAL:HG23	2.08	0.53
1:XA:271:VAL:O	1:XA:271:VAL:HG23	2.08	0.53
1:ZB:812:VAL:HG23	1:ZB:812:VAL:O	2.09	0.53
1:AA:812:VAL:HG23	1:AA:812:VAL:O	2.09	0.53
1:B:381:PRO:HA	1:B:405:THR:HG22	1.90	0.53
1:E:812:VAL:HG23	1:E:812:VAL:O	2.09	0.53
1:F:812:VAL:O	1:F:812:VAL:HG23	2.08	0.53
1:FA:65:VAL:HG11	1:FA:100:TYR:HB2	1.91	0.53
1:GA:812:VAL:O	1:GA:812:VAL:HG23	2.09	0.53
1:GB:812:VAL:HG23	1:GB:812:VAL:O	2.09	0.53
1:H:271:VAL:HG23	1:H:271:VAL:O	2.08	0.53
1:HB:381:PRO:HA	1:HB:405:THR:HG22	1.91	0.53
1:IB:765:VAL:HG22	1:JB:759:LEU:HD21	1.91	0.53
1:JA:230:ARG:HB2	1:JA:268:LEU:HD11	1.91	0.53
1:JA:523:PHE:CD2	1:JA:568:VAL:HG23	2.44	0.53
1:LA:765:VAL:HG22	1:MA:759:LEU:HD21	1.91	0.53

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:MA:230:ARG:HB2	1:MA:268:LEU:HD11	1.91	0.53
1:MB:381:PRO:HA	1:MB:405:THR:HG22	1.90	0.53
1:N:529:ILE:HD12	1:N:583:VAL:HG11	1.90	0.53
1:Q:65:VAL:HG11	1:Q:100:TYR:HB2	1.89	0.53
1:QA:271:VAL:HG23	1:QA:271:VAL:O	2.08	0.53
1:QA:812:VAL:HG23	1:QA:812:VAL:O	2.09	0.53
1:R:381:PRO:HA	1:R:405:THR:HG22	1.91	0.53
1:S:271:VAL:HG23	1:S:271:VAL:O	2.08	0.53
1:SB:812:VAL:O	1:SB:812:VAL:HG23	2.09	0.53
1:T:168:ILE:HG22	1:T:215:LEU:CD2	2.36	0.53
1:T:381:PRO:HA	1:T:405:THR:HG22	1.91	0.53
1:TA:230:ARG:HB2	1:TA:268:LEU:HD11	1.90	0.53
1:TB:65:VAL:HG13	1:TB:103:GLU:OE2	2.07	0.53
1:VB:812:VAL:O	1:VB:812:VAL:HG23	2.09	0.53
1:X:271:VAL:HG23	1:X:271:VAL:O	2.08	0.53
1:A:271:VAL:HG23	1:A:271:VAL:O	2.08	0.53
1:A:381:PRO:HA	1:A:405:THR:HG22	1.91	0.53
1:AC:812:VAL:HG23	1:AC:812:VAL:O	2.09	0.53
1:CA:230:ARG:HB2	1:CA:268:LEU:HD11	1.91	0.53
1:DA:812:VAL:O	1:DA:812:VAL:HG23	2.09	0.53
1:E:381:PRO:HA	1:E:405:THR:HG22	1.90	0.53
1:FA:539:LEU:HD12	1:FA:540:GLN:H	1.74	0.53
1:HA:381:PRO:HA	1:HA:405:THR:HG22	1.91	0.53
1:KA:812:VAL:HG23	1:KA:812:VAL:O	2.09	0.53
1:L:271:VAL:HG23	1:L:271:VAL:O	2.08	0.53
1:LB:812:VAL:HG23	1:LB:812:VAL:O	2.09	0.53
1:NA:65:VAL:HG11	1:NA:100:TYR:HB2	1.91	0.53
1:NB:65:VAL:HG11	1:NB:100:TYR:HB2	1.91	0.53
1:NB:812:VAL:O	1:NB:812:VAL:HG23	2.09	0.53
1:Q:381:PRO:HA	1:Q:405:THR:HG22	1.91	0.53
1:S:381:PRO:HA	1:S:405:THR:HG22	1.91	0.53
1:SA:271:VAL:O	1:SA:271:VAL:HG23	2.07	0.53
1:UA:271:VAL:HG23	1:UA:271:VAL:O	2.08	0.53
1:X:812:VAL:HG23	1:X:812:VAL:O	2.09	0.53
1:Z:230:ARG:HB2	1:Z:268:LEU:HD11	1.90	0.53
1:AA:529:ILE:CD1	1:AA:583:VAL:HG11	2.38	0.53
1:BA:230:ARG:HB2	1:BA:268:LEU:HD11	1.91	0.53
1:EA:230:ARG:HB2	1:EA:268:LEU:HD11	1.91	0.53
1:FA:325:VAL:O	1:FA:325:VAL:HG13	2.09	0.53
1:FA:523:PHE:CD2	1:FA:568:VAL:HG23	2.44	0.53
1:G:381:PRO:HA	1:G:405:THR:HG22	1.90	0.53

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:HB:271:VAL:HG23	1:HB:271:VAL:O	2.08	0.53
1:I:271:VAL:HG23	1:I:271:VAL:O	2.08	0.53
1:IA:230:ARG:HB2	1:IA:268:LEU:HD11	1.91	0.53
1:LA:812:VAL:O	1:LA:812:VAL:HG23	2.09	0.53
1:M:230:ARG:HB2	1:M:268:LEU:HD11	1.90	0.53
1:OB:812:VAL:HG23	1:OB:812:VAL:O	2.09	0.53
1:P:271:VAL:HG23	1:P:271:VAL:O	2.08	0.53
1:P:812:VAL:HG23	1:P:812:VAL:O	2.09	0.53
1:PA:381:PRO:HA	1:PA:405:THR:HG22	1.91	0.53
1:PA:737:GLY:O	1:PA:741:VAL:HG23	2.09	0.53
1:PB:30:VAL:CG2	1:PB:50:MET:HE3	2.39	0.53
1:PB:812:VAL:HG23	1:PB:812:VAL:O	2.09	0.53
1:UB:812:VAL:HG23	1:UB:812:VAL:O	2.09	0.53
1:VA:65:VAL:HG11	1:VA:100:TYR:HB2	1.89	0.53
1:Y:381:PRO:HA	1:Y:405:THR:HG22	1.91	0.53
1:YB:271:VAL:HG23	1:YB:271:VAL:O	2.08	0.53
1:YB:516:LEU:O	1:YB:516:LEU:HD23	2.08	0.53
1:Z:65:VAL:HG11	1:Z:100:TYR:HB2	1.89	0.53
1:ZA:812:VAL:HG23	1:ZA:812:VAL:O	2.09	0.53
1:AA:271:VAL:HG23	1:AA:271:VAL:O	2.08	0.53
1:AB:271:VAL:HG23	1:AB:271:VAL:O	2.08	0.53
1:EA:812:VAL:HG23	1:EA:812:VAL:O	2.09	0.53
1:FA:381:PRO:HA	1:FA:405:THR:HG22	1.91	0.53
1:FB:398:VAL:HG11	1:FB:415:TRP:CE3	2.44	0.53
1:G:812:VAL:HG23	1:G:812:VAL:O	2.09	0.53
1:GA:381:PRO:HA	1:GA:405:THR:HG22	1.91	0.53
1:GB:381:PRO:HA	1:GB:405:THR:HG22	1.91	0.53
1:J:92:LEU:HD12	1:J:92:LEU:N	2.24	0.53
1:J:381:PRO:HA	1:J:405:THR:HG22	1.90	0.53
1:JA:65:VAL:HG11	1:JA:100:TYR:HB2	1.91	0.53
1:M:271:VAL:O	1:M:271:VAL:HG23	2.08	0.53
1:N:812:VAL:HG23	1:N:812:VAL:O	2.09	0.53
1:O:381:PRO:HA	1:O:405:THR:HG22	1.91	0.53
1:OA:381:PRO:HA	1:OA:405:THR:HG22	1.91	0.53
1:OB:271:VAL:O	1:OB:271:VAL:HG23	2.08	0.53
1:Q:812:VAL:O	1:Q:812:VAL:HG23	2.09	0.53
1:QA:230:ARG:HB2	1:QA:268:LEU:HD11	1.90	0.53
1:SB:737:GLY:O	1:SB:741:VAL:HG23	2.09	0.53
1:V:381:PRO:HA	1:V:405:THR:HG22	1.91	0.53
1:V:812:VAL:HG23	1:V:812:VAL:O	2.09	0.53
1:WB:812:VAL:HG23	1:WB:812:VAL:O	2.09	0.53

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:XA:381:PRO:HA	1:XA:405:THR:HG22	1.91	0.53
1:XB:230:ARG:HB2	1:XB:268:LEU:HD11	1.90	0.53
1:Y:230:ARG:HB2	1:Y:268:LEU:HD11	1.91	0.53
1:ZB:381:PRO:HA	1:ZB:405:THR:HG22	1.91	0.53
1:BA:812:VAL:HG23	1:BA:812:VAL:O	2.09	0.53
1:CA:812:VAL:HG23	1:CA:812:VAL:O	2.09	0.53
1:F:65:VAL:HG11	1:F:100:TYR:HB2	1.90	0.53
1:J:812:VAL:HG23	1:J:812:VAL:O	2.09	0.53
1:OA:144:LEU:HD23	1:OA:144:LEU:N	2.24	0.53
1:PA:271:VAL:O	1:PA:271:VAL:HG23	2.08	0.53
1:Q:230:ARG:HB2	1:Q:268:LEU:HD11	1.90	0.53
1:QA:381:PRO:HA	1:QA:405:THR:HG22	1.91	0.53
1:RA:381:PRO:HA	1:RA:405:THR:HG22	1.91	0.53
1:S:812:VAL:HG23	1:S:812:VAL:O	2.09	0.53
1:T:765:VAL:HG22	1:V:759:LEU:HD21	1.91	0.53
1:TA:381:PRO:HA	1:TA:405:THR:HG22	1.91	0.53
1:VA:812:VAL:HG23	1:VA:812:VAL:O	2.09	0.53
1:Y:271:VAL:HG23	1:Y:271:VAL:O	2.08	0.53
1:AC:271:VAL:O	1:AC:271:VAL:HG23	2.08	0.52
1:B:271:VAL:O	1:B:271:VAL:HG23	2.08	0.52
1:C:812:VAL:HG23	1:C:812:VAL:O	2.09	0.52
1:CB:168:ILE:HG22	1:CB:215:LEU:CD2	2.36	0.52
1:EA:381:PRO:HA	1:EA:405:THR:HG22	1.91	0.52
1:FB:737:GLY:O	1:FB:741:VAL:HG23	2.10	0.52
1:K:812:VAL:HG23	1:K:812:VAL:O	2.09	0.52
1:KA:381:PRO:HA	1:KA:405:THR:HG22	1.91	0.52
1:KB:381:PRO:HA	1:KB:405:THR:HG22	1.91	0.52
1:SB:230:ARG:HB2	1:SB:268:LEU:HD11	1.91	0.52
1:TA:366:VAL:HG13	1:TA:366:VAL:O	2.09	0.52
1:ZA:325:VAL:HG13	1:ZA:325:VAL:O	2.09	0.52
1:A:230:ARG:HB2	1:A:268:LEU:HD11	1.92	0.52
1:A:812:VAL:O	1:A:812:VAL:HG23	2.09	0.52
1:CA:381:PRO:HA	1:CA:405:THR:HG22	1.91	0.52
1:CB:812:VAL:O	1:CB:812:VAL:HG23	2.09	0.52
1:H:529:ILE:CD1	1:H:583:VAL:HG11	2.39	0.52
1:IB:812:VAL:HG23	1:IB:812:VAL:O	2.09	0.52
1:JA:381:PRO:HA	1:JA:405:THR:HG22	1.91	0.52
1:JB:381:PRO:HA	1:JB:405:THR:HG22	1.91	0.52
1:K:575:ILE:HG23	1:K:603:VAL:HG22	1.90	0.52
1:KA:65:VAL:HG11	1:KA:100:TYR:HB2	1.91	0.52
1:KB:812:VAL:O	1:KB:812:VAL:HG23	2.09	0.52

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:MA:812:VAL:HG23	1:MA:812:VAL:O	2.09	0.52
1:NA:381:PRO:HA	1:NA:405:THR:HG22	1.91	0.52
1:PA:812:VAL:HG23	1:PA:812:VAL:O	2.09	0.52
1:SA:812:VAL:HG23	1:SA:812:VAL:O	2.09	0.52
1:VB:230:ARG:HB2	1:VB:268:LEU:HD11	1.91	0.52
1:VB:398:VAL:HG11	1:VB:415:TRP:CE3	2.45	0.52
1:XB:812:VAL:HG23	1:XB:812:VAL:O	2.10	0.52
1:EA:529:ILE:HD12	1:EA:583:VAL:HG11	1.90	0.52
1:FA:812:VAL:HG23	1:FA:812:VAL:O	2.09	0.52
1:GB:65:VAL:HG11	1:GB:100:TYR:HB2	1.92	0.52
1:H:381:PRO:HA	1:H:405:THR:HG22	1.91	0.52
1:HB:737:GLY:O	1:HB:741:VAL:HG23	2.10	0.52
1:IA:575:ILE:HG23	1:IA:603:VAL:HG22	1.92	0.52
1:J:271:VAL:O	1:J:271:VAL:HG23	2.08	0.52
1:JB:812:VAL:HG23	1:JB:812:VAL:O	2.09	0.52
1:K:65:VAL:HG11	1:K:100:TYR:HB2	1.90	0.52
1:M:529:ILE:HD12	1:M:583:VAL:HG11	1.90	0.52
1:N:575:ILE:HG23	1:N:603:VAL:HG22	1.91	0.52
1:PB:529:ILE:CD1	1:PB:583:VAL:HG11	2.40	0.52
1:QB:381:PRO:HA	1:QB:405:THR:HG22	1.91	0.52
1:R:271:VAL:O	1:R:271:VAL:HG23	2.08	0.52
1:R:737:GLY:O	1:R:741:VAL:HG23	2.10	0.52
1:T:812:VAL:O	1:T:812:VAL:HG23	2.09	0.52
1:TB:523:PHE:CD2	1:TB:568:VAL:HG23	2.43	0.52
1:TB:812:VAL:HG23	1:TB:812:VAL:O	2.09	0.52
1:W:812:VAL:HG23	1:W:812:VAL:O	2.09	0.52
1:X:381:PRO:HA	1:X:405:THR:HG22	1.91	0.52
1:YA:523:PHE:CD2	1:YA:568:VAL:HG23	2.45	0.52
1:ZA:271:VAL:HG23	1:ZA:271:VAL:O	2.08	0.52
1:B:737:GLY:O	1:B:741:VAL:HG23	2.09	0.52
1:DB:812:VAL:HG23	1:DB:812:VAL:O	2.09	0.52
1:HB:325:VAL:HG13	1:HB:325:VAL:O	2.09	0.52
1:I:381:PRO:HA	1:I:405:THR:HG22	1.90	0.52
1:IA:765:VAL:HG22	1:JA:759:LEU:HD21	1.92	0.52
1:JA:812:VAL:HG23	1:JA:812:VAL:O	2.09	0.52
1:K:381:PRO:HA	1:K:405:THR:HG22	1.91	0.52
1:LA:65:VAL:HG11	1:LA:100:TYR:HB2	1.91	0.52
1:LB:30:VAL:HG21	1:LB:50:MET:HE3	1.90	0.52
1:MB:529:ILE:HD12	1:MB:583:VAL:HG11	1.91	0.52
1:NB:381:PRO:HA	1:NB:405:THR:HG22	1.91	0.52
1:OA:271:VAL:HG23	1:OA:271:VAL:O	2.08	0.52

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:P:381:PRO:HA	1:P:405:THR:HG22	1.91	0.52
1:R:230:ARG:HB2	1:R:268:LEU:HD11	1.91	0.52
1:RA:812:VAL:HG23	1:RA:812:VAL:O	2.09	0.52
1:SB:523:PHE:CD2	1:SB:568:VAL:HG23	2.44	0.52
1:V:271:VAL:HG23	1:V:271:VAL:O	2.08	0.52
1:WA:65:VAL:HG11	1:WA:100:TYR:HB2	1.91	0.52
1:WB:381:PRO:HA	1:WB:405:THR:HG22	1.91	0.52
1:Z:271:VAL:HG23	1:Z:271:VAL:O	2.08	0.52
1:Z:325:VAL:O	1:Z:325:VAL:HG13	2.09	0.52
1:ZA:230:ARG:HB2	1:ZA:268:LEU:HD11	1.92	0.52
1:ZB:65:VAL:HG11	1:ZB:100:TYR:HB2	1.91	0.52
1:AB:48:MET:HA	1:AB:48:MET:HE3	1.91	0.52
1:B:575:ILE:HG23	1:B:603:VAL:HG22	1.91	0.52
1:CA:30:VAL:CG2	1:CA:50:MET:HE3	2.40	0.52
1:CB:144:LEU:HD12	1:CB:144:LEU:O	2.10	0.52
1:D:381:PRO:HA	1:D:405:THR:HG22	1.91	0.52
1:FB:168:ILE:HG22	1:FB:215:LEU:CD2	2.39	0.52
1:FB:539:LEU:HD11	1:FB:541:LEU:HD21	1.92	0.52
1:FB:812:VAL:O	1:FB:812:VAL:HG23	2.09	0.52
1:G:230:ARG:HB2	1:G:268:LEU:HD11	1.92	0.52
1:HB:65:VAL:HG11	1:HB:100:TYR:HB2	1.91	0.52
1:IB:381:PRO:HA	1:IB:405:THR:HG22	1.91	0.52
1:J:65:VAL:HG11	1:J:100:TYR:HB2	1.90	0.52
1:LA:516:LEU:O	1:LA:516:LEU:HD23	2.08	0.52
1:LB:65:VAL:HG11	1:LB:100:TYR:HB2	1.91	0.52
1:LB:230:ARG:HB2	1:LB:268:LEU:HD11	1.92	0.52
1:M:398:VAL:HG11	1:M:415:TRP:CE3	2.44	0.52
1:MA:390:VAL:HG23	1:MA:390:VAL:O	2.10	0.52
1:O:398:VAL:HG11	1:O:415:TRP:CE3	2.44	0.52
1:Q:302:VAL:HG11	1:Q:306:LYS:HZ2	1.74	0.52
1:R:65:VAL:HG11	1:R:100:TYR:HB2	1.90	0.52
1:SA:523:PHE:CD2	1:SA:568:VAL:HG23	2.45	0.52
1:V:529:ILE:HD12	1:V:583:VAL:HG11	1.91	0.52
1:VB:381:PRO:HA	1:VB:405:THR:HG22	1.91	0.52
1:WA:230:ARG:HB2	1:WA:268:LEU:HD11	1.91	0.52
1:WA:575:ILE:HG23	1:WA:603:VAL:HG22	1.91	0.52
1:YB:812:VAL:HG23	1:YB:812:VAL:O	2.09	0.52
1:Z:30:VAL:HG22	1:Z:50:MET:HE2	1.92	0.52
1:Z:529:ILE:HD12	1:Z:583:VAL:HG11	1.92	0.52
1:AB:812:VAL:HG23	1:AB:812:VAL:O	2.09	0.52
1:BB:65:VAL:HG11	1:BB:100:TYR:HB2	1.92	0.52

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:CA:765:VAL:HG22	1:DA:759:LEU:HD21	1.92	0.52
1:DA:65:VAL:HG11	1:DA:100:TYR:HB2	1.92	0.52
1:F:398:VAL:HG11	1:F:415:TRP:CE3	2.45	0.52
1:GB:271:VAL:O	1:GB:271:VAL:HG23	2.08	0.52
1:JA:398:VAL:HG11	1:JA:415:TRP:CE3	2.45	0.52
1:JB:529:ILE:HD12	1:JB:583:VAL:HG11	1.91	0.52
1:K:523:PHE:CD2	1:K:568:VAL:HG23	2.44	0.52
1:KB:390:VAL:HG23	1:KB:390:VAL:O	2.10	0.52
1:KB:737:GLY:O	1:KB:741:VAL:HG23	2.09	0.52
1:LB:381:PRO:HA	1:LB:405:THR:HG22	1.91	0.52
1:PA:523:PHE:CD2	1:PA:568:VAL:HG23	2.45	0.52
1:Q:271:VAL:HG23	1:Q:271:VAL:O	2.08	0.52
1:RB:65:VAL:HG11	1:RB:100:TYR:HB2	1.90	0.52
1:SB:65:VAL:HG11	1:SB:100:TYR:HB2	1.91	0.52
1:SB:381:PRO:HA	1:SB:405:THR:HG22	1.91	0.52
1:T:271:VAL:HG23	1:T:271:VAL:O	2.08	0.52
1:VB:765:VAL:HG22	1:WB:759:LEU:HD21	1.91	0.52
1:W:230:ARG:HB2	1:W:268:LEU:HD11	1.91	0.52
1:XB:381:PRO:HA	1:XB:405:THR:HG22	1.91	0.52
1:Y:398:VAL:HG11	1:Y:415:TRP:CE3	2.45	0.52
1:YA:812:VAL:HG23	1:YA:812:VAL:O	2.09	0.52
1:Z:812:VAL:HG23	1:Z:812:VAL:O	2.10	0.52
1:AB:398:VAL:HG11	1:AB:415:TRP:CE3	2.44	0.52
1:BB:812:VAL:HG23	1:BB:812:VAL:O	2.09	0.52
1:CA:529:ILE:CD1	1:CA:583:VAL:HG11	2.40	0.52
1:CB:65:VAL:HG11	1:CB:100:TYR:HB2	1.92	0.52
1:DB:17:HIS:HB3	1:DB:44:LEU:HD23	1.91	0.52
1:DB:65:VAL:HG11	1:DB:100:TYR:HB2	1.92	0.52
1:GA:65:VAL:HG11	1:GA:100:TYR:HB2	1.92	0.52
1:GA:230:ARG:HB2	1:GA:268:LEU:HD11	1.92	0.52
1:I:65:VAL:HG11	1:I:100:TYR:HB2	1.90	0.52
1:I:168:ILE:HG22	1:I:215:LEU:CD2	2.37	0.52
1:JB:575:ILE:HG23	1:JB:603:VAL:HG22	1.92	0.52
1:L:381:PRO:HA	1:L:405:THR:HG22	1.91	0.52
1:NB:230:ARG:HB2	1:NB:268:LEU:HD11	1.92	0.52
1:OA:765:VAL:HG22	1:PA:759:LEU:HD21	1.92	0.52
1:R:398:VAL:HG11	1:R:415:TRP:CE3	2.44	0.52
1:T:523:PHE:CD2	1:T:568:VAL:HG23	2.45	0.52
1:TB:381:PRO:HA	1:TB:405:THR:HG22	1.91	0.52
1:WB:65:VAL:HG11	1:WB:100:TYR:HB2	1.90	0.52
1:XB:271:VAL:HG23	1:XB:271:VAL:O	2.08	0.52

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:YA:398:VAL:HG11	1:YA:415:TRP:CE3	2.45	0.52
1:YB:48:MET:HA	1:YB:48:MET:HE3	1.91	0.52
1:YB:230:ARG:HB2	1:YB:268:LEU:HD11	1.91	0.52
1:ZA:65:VAL:HG11	1:ZA:100:TYR:HB2	1.92	0.52
1:ZB:390:VAL:HG23	1:ZB:390:VAL:O	2.10	0.52
1:C:523:PHE:CD2	1:C:568:VAL:HG23	2.44	0.52
1:G:529:ILE:HD12	1:G:583:VAL:HG11	1.92	0.52
1:GB:30:VAL:HG21	1:GB:50:MET:HE3	1.91	0.52
1:H:497:VAL:HG13	1:H:497:VAL:O	2.10	0.52
1:HA:812:VAL:HG23	1:HA:812:VAL:O	2.09	0.52
1:HB:812:VAL:HG23	1:HB:812:VAL:O	2.09	0.52
1:I:230:ARG:HB2	1:I:268:LEU:HD11	1.92	0.52
1:I:529:ILE:HD12	1:I:583:VAL:HG11	1.92	0.52
1:IA:529:ILE:HD12	1:IA:583:VAL:HG11	1.91	0.52
1:NA:523:PHE:CD2	1:NA:568:VAL:HG23	2.45	0.52
1:P:523:PHE:CD2	1:P:568:VAL:HG23	2.45	0.52
1:PA:230:ARG:HB2	1:PA:268:LEU:HD11	1.91	0.52
1:QB:65:VAL:HG11	1:QB:100:TYR:HB2	1.91	0.52
1:R:539:LEU:HD12	1:R:540:GLN:N	2.25	0.52
1:RB:381:PRO:HA	1:RB:405:THR:HG22	1.91	0.52
1:RB:812:VAL:HG23	1:RB:812:VAL:O	2.09	0.52
1:TB:325:VAL:HG13	1:TB:325:VAL:O	2.10	0.52
1:VB:529:ILE:HD12	1:VB:583:VAL:HG11	1.91	0.52
1:Y:65:VAL:HG11	1:Y:100:TYR:HB2	1.91	0.52
1:C:381:PRO:HA	1:C:405:THR:HG22	1.91	0.52
1:E:765:VAL:HG22	1:F:759:LEU:HD21	1.92	0.52
1:H:390:VAL:HG23	1:H:390:VAL:O	2.10	0.52
1:H:812:VAL:HG23	1:H:812:VAL:O	2.09	0.52
1:IA:398:VAL:HG11	1:IA:415:TRP:CE3	2.45	0.52
1:J:230:ARG:HB2	1:J:268:LEU:HD11	1.92	0.52
1:M:812:VAL:HG23	1:M:812:VAL:O	2.09	0.52
1:TB:230:ARG:HB2	1:TB:268:LEU:HD11	1.91	0.52
1:UA:812:VAL:HG23	1:UA:812:VAL:O	2.09	0.52
1:VA:390:VAL:HG23	1:VA:390:VAL:O	2.10	0.52
1:WB:230:ARG:HB2	1:WB:268:LEU:HD11	1.91	0.52
1:WB:398:VAL:HG11	1:WB:415:TRP:CE3	2.45	0.52
1:Y:812:VAL:O	1:Y:812:VAL:HG23	2.09	0.52
1:AA:230:ARG:HB2	1:AA:268:LEU:HD11	1.92	0.52
1:BA:144:LEU:O	1:BA:144:LEU:HD12	2.10	0.52
1:DB:497:VAL:HG13	1:DB:497:VAL:O	2.10	0.52
1:F:516:LEU:O	1:F:516:LEU:HD23	2.10	0.52

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:HB:523:PHE:CD2	1:HB:568:VAL:HG23	2.45	0.52
1:JB:497:VAL:HG13	1:JB:497:VAL:O	2.10	0.52
1:K:398:VAL:HG11	1:K:415:TRP:CE3	2.45	0.52
1:MB:390:VAL:HG23	1:MB:390:VAL:O	2.10	0.52
1:N:65:VAL:HG11	1:N:100:TYR:HB2	1.91	0.52
1:OA:398:VAL:HG11	1:OA:415:TRP:CE3	2.45	0.52
1:QA:52:THR:O	1:QA:109:ILE:HD11	2.09	0.52
1:RB:390:VAL:HG23	1:RB:390:VAL:O	2.10	0.52
1:SA:765:VAL:HG22	1:TA:759:LEU:HD21	1.92	0.52
1:TA:398:VAL:HG11	1:TA:415:TRP:CE3	2.45	0.52
1:TA:516:LEU:O	1:TA:516:LEU:HD23	2.10	0.52
1:V:65:VAL:HG11	1:V:100:TYR:HB2	1.91	0.52
1:W:529:ILE:HD12	1:W:583:VAL:HG11	1.92	0.52
1:WB:497:VAL:HG13	1:WB:497:VAL:O	2.10	0.52
1:XA:230:ARG:HB2	1:XA:268:LEU:HD11	1.92	0.52
1:ZB:230:ARG:HB2	1:ZB:268:LEU:HD11	1.91	0.52
1:AB:765:VAL:HG22	1:BB:759:LEU:HD21	1.92	0.51
1:CA:575:ILE:HG23	1:CA:603:VAL:HG22	1.92	0.51
1:D:398:VAL:HG11	1:D:415:TRP:CE3	2.45	0.51
1:DB:575:ILE:HG23	1:DB:603:VAL:HG22	1.92	0.51
1:FA:30:VAL:HG21	1:FA:50:MET:HE3	1.92	0.51
1:I:398:VAL:HG11	1:I:415:TRP:CE3	2.45	0.51
1:IB:398:VAL:HG11	1:IB:415:TRP:CE3	2.45	0.51
1:L:65:VAL:HG11	1:L:100:TYR:HB2	1.91	0.51
1:MB:812:VAL:HG23	1:MB:812:VAL:O	2.09	0.51
1:P:390:VAL:HG23	1:P:390:VAL:O	2.10	0.51
1:PB:381:PRO:HA	1:PB:405:THR:HG22	1.91	0.51
1:S:230:ARG:HB2	1:S:268:LEU:HD11	1.91	0.51
1:TB:65:VAL:HG11	1:TB:100:TYR:HB2	1.92	0.51
1:UB:381:PRO:HA	1:UB:405:THR:HG22	1.91	0.51
1:VA:529:ILE:HD12	1:VA:583:VAL:HG11	1.92	0.51
1:X:390:VAL:HG23	1:X:390:VAL:O	2.10	0.51
1:Z:30:VAL:CG2	1:Z:50:MET:HE2	2.41	0.51
1:BB:575:ILE:HG23	1:BB:603:VAL:HG22	1.91	0.51
1:CB:575:ILE:HG23	1:CB:603:VAL:HG22	1.92	0.51
1:EA:390:VAL:HG23	1:EA:390:VAL:O	2.10	0.51
1:FB:497:VAL:O	1:FB:497:VAL:HG13	2.10	0.51
1:L:390:VAL:O	1:L:390:VAL:HG23	2.10	0.51
1:MB:325:VAL:O	1:MB:325:VAL:HG13	2.09	0.51
1:PB:575:ILE:HG23	1:PB:603:VAL:HG22	1.92	0.51
1:SB:325:VAL:HG13	1:SB:325:VAL:O	2.09	0.51

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:WB:271:VAL:O	1:WB:271:VAL:HG23	2.08	0.51
1:X:398:VAL:HG11	1:X:415:TRP:CE3	2.45	0.51
1:YB:144:LEU:O	1:YB:144:LEU:HD12	2.10	0.51
1:ZA:398:VAL:HG11	1:ZA:415:TRP:CE3	2.46	0.51
1:ZA:523:PHE:CD2	1:ZA:568:VAL:HG23	2.46	0.51
1:B:398:VAL:HG11	1:B:415:TRP:CE3	2.46	0.51
1:BB:230:ARG:HB2	1:BB:268:LEU:HD11	1.92	0.51
1:CA:144:LEU:HD12	1:CA:144:LEU:O	2.10	0.51
1:F:381:PRO:HA	1:F:405:THR:HG22	1.92	0.51
1:F:529:ILE:HD12	1:F:583:VAL:HG11	1.92	0.51
1:HA:325:VAL:O	1:HA:325:VAL:HG13	2.10	0.51
1:JA:144:LEU:O	1:JA:144:LEU:HD12	2.10	0.51
1:MB:575:ILE:HG23	1:MB:603:VAL:HG22	1.91	0.51
1:OB:778:GLU:HA	1:OB:781:VAL:HG22	1.92	0.51
1:PA:575:ILE:HG23	1:PA:603:VAL:HG22	1.91	0.51
1:PB:17:HIS:CB	1:PB:44:LEU:HD23	2.40	0.51
1:QA:523:PHE:CD2	1:QA:568:VAL:HG23	2.44	0.51
1:RA:398:VAL:HG11	1:RA:415:TRP:CE3	2.45	0.51
1:TA:325:VAL:HG13	1:TA:325:VAL:O	2.09	0.51
1:UA:230:ARG:HB2	1:UA:268:LEU:HD11	1.92	0.51
1:W:575:ILE:HG23	1:W:603:VAL:HG22	1.91	0.51
1:WA:529:ILE:HD12	1:WA:583:VAL:HG11	1.92	0.51
1:YB:539:LEU:HD21	1:YB:541:LEU:HD21	1.92	0.51
1:A:765:VAL:HG22	1:B:759:LEU:HD21	1.91	0.51
1:AC:398:VAL:HG11	1:AC:415:TRP:CE3	2.46	0.51
1:AC:523:PHE:CD2	1:AC:568:VAL:HG23	2.45	0.51
1:JA:497:VAL:HG13	1:JA:497:VAL:O	2.11	0.51
1:L:230:ARG:HB2	1:L:268:LEU:HD11	1.92	0.51
1:L:737:GLY:O	1:L:741:VAL:HG23	2.11	0.51
1:LB:390:VAL:HG23	1:LB:390:VAL:O	2.11	0.51
1:LB:398:VAL:HG11	1:LB:415:TRP:CE3	2.45	0.51
1:NA:398:VAL:HG11	1:NA:415:TRP:CE3	2.46	0.51
1:NB:529:ILE:HD12	1:NB:583:VAL:HG11	1.92	0.51
1:P:575:ILE:HG23	1:P:603:VAL:HG22	1.92	0.51
1:PA:65:VAL:HG12	1:PA:83:LEU:CD2	2.40	0.51
1:QB:398:VAL:HG11	1:QB:415:TRP:CE3	2.45	0.51
1:T:325:VAL:HG13	1:T:325:VAL:O	2.11	0.51
1:VA:398:VAL:HG11	1:VA:415:TRP:CE3	2.46	0.51
1:W:325:VAL:HG13	1:W:325:VAL:O	2.09	0.51
1:WA:398:VAL:HG11	1:WA:415:TRP:CE3	2.45	0.51
1:XA:325:VAL:O	1:XA:325:VAL:HG13	2.11	0.51

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:YA:230:ARG:HB2	1:YA:268:LEU:HD11	1.93	0.51
1:ZA:737:GLY:O	1:ZA:741:VAL:HG23	2.11	0.51
1:ZB:398:VAL:HG11	1:ZB:415:TRP:CE3	2.46	0.51
1:A:575:ILE:HG23	1:A:603:VAL:HG22	1.92	0.51
1:AC:144:LEU:HD12	1:AC:144:LEU:O	2.10	0.51
1:BB:398:VAL:HG11	1:BB:415:TRP:CE3	2.46	0.51
1:CA:737:GLY:O	1:CA:741:VAL:HG23	2.11	0.51
1:CB:325:VAL:HG13	1:CB:325:VAL:O	2.10	0.51
1:DB:523:PHE:CD2	1:DB:568:VAL:HG23	2.46	0.51
1:FB:523:PHE:CD2	1:FB:568:VAL:HG23	2.46	0.51
1:H:398:VAL:HG11	1:H:415:TRP:CE3	2.45	0.51
1:HA:765:VAL:HG22	1:IA:759:LEU:HD21	1.93	0.51
1:JB:325:VAL:O	1:JB:325:VAL:HG13	2.09	0.51
1:M:144:LEU:HD12	1:M:144:LEU:O	2.10	0.51
1:M:575:ILE:HG23	1:M:603:VAL:HG22	1.92	0.51
1:M:765:VAL:HG22	1:N:759:LEU:HD21	1.93	0.51
1:MB:765:VAL:HG22	1:NB:759:LEU:HD21	1.93	0.51
1:PA:325:VAL:O	1:PA:325:VAL:HG13	2.09	0.51
1:PB:737:GLY:O	1:PB:741:VAL:HG23	2.11	0.51
1:QA:325:VAL:HG13	1:QA:325:VAL:O	2.11	0.51
1:QA:737:GLY:O	1:QA:741:VAL:HG23	2.10	0.51
1:R:575:ILE:HG23	1:R:603:VAL:HG22	1.92	0.51
1:R:765:VAL:HG22	1:S:759:LEU:HD21	1.91	0.51
1:RA:65:VAL:HG11	1:RA:100:TYR:HB2	1.91	0.51
1:RA:271:VAL:HG23	1:RA:271:VAL:O	2.08	0.51
1:TB:17:HIS:HB3	1:TB:44:LEU:HD23	1.92	0.51
1:UB:529:ILE:CD1	1:UB:583:VAL:HG11	2.41	0.51
1:V:325:VAL:HG13	1:V:325:VAL:O	2.11	0.51
1:VA:144:LEU:HD12	1:VA:144:LEU:O	2.11	0.51
1:AB:538:GLN:HB2	1:AB:646:VAL:HG22	1.93	0.51
1:AC:30:VAL:HG21	1:AC:50:MET:HE3	1.92	0.51
1:AC:65:VAL:HG11	1:AC:100:TYR:HB2	1.91	0.51
1:AC:325:VAL:O	1:AC:325:VAL:HG13	2.11	0.51
1:BB:17:HIS:HB3	1:BB:44:LEU:HD23	1.93	0.51
1:C:390:VAL:HG23	1:C:390:VAL:O	2.11	0.51
1:CA:523:PHE:CD2	1:CA:568:VAL:HG23	2.46	0.51
1:DA:230:ARG:HB2	1:DA:268:LEU:HD11	1.93	0.51
1:DA:398:VAL:HG11	1:DA:415:TRP:CE3	2.46	0.51
1:EA:523:PHE:CD2	1:EA:568:VAL:HG23	2.46	0.51
1:FB:529:ILE:HD12	1:FB:583:VAL:HG11	1.92	0.51
1:GB:325:VAL:O	1:GB:325:VAL:HG13	2.09	0.51

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:GB:398:VAL:HG11	1:GB:415:TRP:CE3	2.46	0.51
1:HA:529:ILE:CD1	1:HA:583:VAL:HG11	2.41	0.51
1:IB:230:ARG:HB2	1:IB:268:LEU:HD11	1.93	0.51
1:JB:398:VAL:HG11	1:JB:415:TRP:CE3	2.46	0.51
1:KB:65:VAL:HG11	1:KB:100:TYR:HB2	1.93	0.51
1:L:523:PHE:CD2	1:L:568:VAL:HG23	2.46	0.51
1:MA:398:VAL:HG11	1:MA:415:TRP:CE3	2.46	0.51
1:MA:765:VAL:HG22	1:NA:759:LEU:HD21	1.93	0.51
1:P:230:ARG:HB2	1:P:268:LEU:HD11	1.91	0.51
1:P:497:VAL:HG13	1:P:497:VAL:O	2.10	0.51
1:P:765:VAL:HG22	1:Q:759:LEU:HD21	1.93	0.51
1:QB:230:ARG:HB2	1:QB:268:LEU:HD11	1.92	0.51
1:R:48:MET:HE2	1:R:48:MET:HA	1.92	0.51
1:SB:390:VAL:HG23	1:SB:390:VAL:O	2.11	0.51
1:TA:529:ILE:HD12	1:TA:583:VAL:HG11	1.92	0.51
1:TB:390:VAL:HG23	1:TB:390:VAL:O	2.11	0.51
1:UA:529:ILE:HD12	1:UA:583:VAL:HG11	1.92	0.51
1:UB:765:VAL:HG22	1:VB:759:LEU:HD21	1.92	0.51
1:V:398:VAL:HG11	1:V:415:TRP:CE3	2.45	0.51
1:XB:65:VAL:HG11	1:XB:100:TYR:HB2	1.92	0.51
1:Y:390:VAL:HG23	1:Y:390:VAL:O	2.10	0.51
1:CA:398:VAL:HG11	1:CA:415:TRP:CE3	2.46	0.51
1:F:390:VAL:HG23	1:F:390:VAL:O	2.11	0.51
1:GA:398:VAL:HG11	1:GA:415:TRP:CE3	2.46	0.51
1:H:144:LEU:HD12	1:H:144:LEU:O	2.10	0.51
1:H:230:ARG:HB2	1:H:268:LEU:HD11	1.93	0.51
1:I:52:THR:O	1:I:109:ILE:HD11	2.10	0.51
1:KB:398:VAL:HG11	1:KB:415:TRP:CE3	2.45	0.51
1:OA:65:VAL:HG11	1:OA:100:TYR:HB2	1.92	0.51
1:OA:230:ARG:HB2	1:OA:268:LEU:HD11	1.91	0.51
1:PA:398:VAL:HG11	1:PA:415:TRP:CE3	2.46	0.51
1:SB:529:ILE:HD12	1:SB:583:VAL:HG11	1.92	0.51
1:WB:538:GLN:HB2	1:WB:646:VAL:HG22	1.92	0.51
1:YB:398:VAL:HG11	1:YB:415:TRP:CE3	2.46	0.51
1:Z:52:THR:O	1:Z:109:ILE:HD11	2.11	0.51
1:A:398:VAL:HG11	1:A:415:TRP:CE3	2.45	0.51
1:BA:778:GLU:HA	1:BA:781:VAL:HG22	1.92	0.51
1:C:737:GLY:O	1:C:741:VAL:HG23	2.10	0.51
1:EA:325:VAL:O	1:EA:325:VAL:HG13	2.11	0.51
1:G:52:THR:O	1:G:109:ILE:HD11	2.10	0.51
1:GA:529:ILE:CD1	1:GA:583:VAL:HG21	2.41	0.51

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:HA:144:LEU:HD12	1:HA:144:LEU:O	2.11	0.51
1:IA:65:VAL:HG11	1:IA:100:TYR:HB2	1.93	0.51
1:KB:230:ARG:HB2	1:KB:268:LEU:HD11	1.93	0.51
1:LB:529:ILE:HD12	1:LB:583:VAL:HG11	1.93	0.51
1:OA:325:VAL:HG13	1:OA:325:VAL:O	2.11	0.51
1:PB:144:LEU:HD12	1:PB:144:LEU:O	2.10	0.51
1:RB:523:PHE:CD2	1:RB:568:VAL:HG23	2.46	0.51
1:S:65:VAL:HG11	1:S:100:TYR:HB2	1.91	0.51
1:TB:398:VAL:HG11	1:TB:415:TRP:CE3	2.46	0.51
1:UB:516:LEU:HD23	1:UB:516:LEU:O	2.11	0.51
1:V:92:LEU:N	1:V:92:LEU:HD12	2.26	0.51
1:VA:325:VAL:O	1:VA:325:VAL:HG13	2.11	0.51
1:W:497:VAL:HG13	1:W:497:VAL:O	2.10	0.51
1:WB:144:LEU:HD12	1:WB:144:LEU:O	2.10	0.51
1:X:737:GLY:O	1:X:741:VAL:HG23	2.11	0.51
1:AA:529:ILE:HD12	1:AA:583:VAL:HG11	1.93	0.51
1:BA:325:VAL:HG13	1:BA:325:VAL:O	2.11	0.51
1:BA:575:ILE:HG23	1:BA:603:VAL:HG22	1.93	0.51
1:CA:390:VAL:HG23	1:CA:390:VAL:O	2.10	0.51
1:DA:575:ILE:HG23	1:DA:603:VAL:HG22	1.92	0.51
1:FA:529:ILE:HD12	1:FA:583:VAL:HG11	1.92	0.51
1:HB:575:ILE:HG23	1:HB:603:VAL:HG22	1.93	0.51
1:I:575:ILE:HG23	1:I:603:VAL:HG22	1.91	0.51
1:LA:325:VAL:HG13	1:LA:325:VAL:O	2.11	0.51
1:LB:575:ILE:HG23	1:LB:603:VAL:HG22	1.91	0.51
1:NA:144:LEU:HD12	1:NA:144:LEU:O	2.10	0.51
1:NA:575:ILE:HG23	1:NA:603:VAL:HG22	1.93	0.51
1:O:575:ILE:HG23	1:O:603:VAL:HG22	1.92	0.51
1:P:65:VAL:HG11	1:P:100:TYR:HB2	1.92	0.51
1:PB:325:VAL:HG13	1:PB:325:VAL:O	2.10	0.51
1:PB:523:PHE:CD2	1:PB:568:VAL:HG23	2.46	0.51
1:R:523:PHE:CD2	1:R:568:VAL:HG23	2.46	0.51
1:RA:325:VAL:HG13	1:RA:325:VAL:O	2.11	0.51
1:UA:48:MET:HA	1:UA:48:MET:HE3	1.93	0.51
1:UA:325:VAL:HG13	1:UA:325:VAL:O	2.11	0.51
1:UA:765:VAL:HG22	1:VA:759:LEU:HD21	1.93	0.51
1:VA:497:VAL:O	1:VA:497:VAL:HG13	2.10	0.51
1:VB:575:ILE:HG23	1:VB:603:VAL:HG22	1.91	0.51
1:ZA:539:LEU:HD23	1:ZA:540:GLN:N	2.26	0.51
1:B:325:VAL:O	1:B:325:VAL:HG13	2.11	0.51
1:F:325:VAL:HG13	1:F:325:VAL:O	2.10	0.51

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:G:390:VAL:HG23	1:G:390:VAL:O	2.11	0.51
1:G:398:VAL:HG11	1:G:415:TRP:CE3	2.46	0.51
1:H:17:HIS:CB	1:H:44:LEU:HD23	2.41	0.51
1:HA:52:THR:O	1:HA:109:ILE:HD11	2.10	0.51
1:L:398:VAL:HG11	1:L:415:TRP:CE3	2.46	0.51
1:N:390:VAL:HG23	1:N:390:VAL:O	2.12	0.51
1:N:398:VAL:HG11	1:N:415:TRP:CE3	2.46	0.51
1:QB:390:VAL:O	1:QB:390:VAL:HG23	2.11	0.51
1:R:529:ILE:HD12	1:R:583:VAL:HG11	1.92	0.51
1:WB:529:ILE:HD12	1:WB:583:VAL:HG11	1.93	0.51
1:X:144:LEU:HD12	1:X:144:LEU:O	2.11	0.51
1:AA:92:LEU:N	1:AA:92:LEU:HD12	2.25	0.50
1:AC:765:VAL:HG22	1:OA:759:LEU:HD21	1.93	0.50
1:C:398:VAL:HG11	1:C:415:TRP:CE3	2.46	0.50
1:D:529:ILE:HD12	1:D:583:VAL:HG11	1.92	0.50
1:E:575:ILE:HG23	1:E:603:VAL:HG22	1.93	0.50
1:FA:765:VAL:HG22	1:GA:759:LEU:HD21	1.94	0.50
1:J:398:VAL:HG11	1:J:415:TRP:CE3	2.47	0.50
1:L:539:LEU:HD23	1:L:540:GLN:N	2.26	0.50
1:M:325:VAL:O	1:M:325:VAL:HG13	2.11	0.50
1:MA:334:LEU:HD12	1:MA:357:TRP:NE1	2.27	0.50
1:MB:497:VAL:O	1:MB:497:VAL:HG13	2.10	0.50
1:OB:529:ILE:HD12	1:OB:583:VAL:HG11	1.94	0.50
1:PA:65:VAL:HG11	1:PA:100:TYR:HB2	1.92	0.50
1:PB:398:VAL:HG11	1:PB:415:TRP:CE3	2.46	0.50
1:QA:398:VAL:HG11	1:QA:415:TRP:CE3	2.46	0.50
1:QB:497:VAL:HG13	1:QB:497:VAL:O	2.12	0.50
1:RA:497:VAL:HG13	1:RA:497:VAL:O	2.11	0.50
1:RB:398:VAL:HG11	1:RB:415:TRP:CE3	2.46	0.50
1:SB:497:VAL:O	1:SB:497:VAL:HG13	2.10	0.50
1:SB:765:VAL:HG22	1:TB:759:LEU:HD21	1.93	0.50
1:UB:92:LEU:HD12	1:UB:92:LEU:N	2.26	0.50
1:UB:758:GLU:O	1:UB:762:VAL:HG22	2.11	0.50
1:W:398:VAL:HG11	1:W:415:TRP:CE3	2.46	0.50
1:YA:737:GLY:O	1:YA:741:VAL:HG23	2.11	0.50
1:Z:334:LEU:HD12	1:Z:357:TRP:NE1	2.26	0.50
1:A:325:VAL:HG13	1:A:325:VAL:O	2.11	0.50
1:AA:759:LEU:HD21	1:Z:765:VAL:HG22	1.92	0.50
1:B:65:VAL:HG11	1:B:100:TYR:HB2	1.93	0.50
1:BA:398:VAL:HG11	1:BA:415:TRP:CE3	2.46	0.50
1:CA:497:VAL:HG13	1:CA:497:VAL:O	2.11	0.50

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:DA:52:THR:O	1:DA:109:ILE:HD11	2.11	0.50
1:DB:390:VAL:HG23	1:DB:390:VAL:O	2.11	0.50
1:E:65:VAL:HG11	1:E:100:TYR:HB2	1.93	0.50
1:E:390:VAL:HG23	1:E:390:VAL:O	2.11	0.50
1:F:765:VAL:HG22	1:G:759:LEU:HD21	1.93	0.50
1:FA:497:VAL:O	1:FA:497:VAL:HG13	2.11	0.50
1:G:765:VAL:HG22	1:H:759:LEU:HD21	1.93	0.50
1:H:325:VAL:O	1:H:325:VAL:HG13	2.11	0.50
1:IB:325:VAL:HG13	1:IB:325:VAL:O	2.11	0.50
1:JB:390:VAL:HG23	1:JB:390:VAL:O	2.11	0.50
1:K:230:ARG:HB2	1:K:268:LEU:HD11	1.93	0.50
1:KA:390:VAL:HG23	1:KA:390:VAL:O	2.11	0.50
1:KA:778:GLU:HA	1:KA:781:VAL:HG22	1.93	0.50
1:MB:52:THR:O	1:MB:109:ILE:HD11	2.12	0.50
1:N:497:VAL:O	1:N:497:VAL:HG13	2.11	0.50
1:O:325:VAL:HG13	1:O:325:VAL:O	2.10	0.50
1:O:778:GLU:HA	1:O:781:VAL:HG22	1.93	0.50
1:O:812:VAL:HG23	1:O:812:VAL:O	2.09	0.50
1:T:65:VAL:HG11	1:T:100:TYR:HB2	1.91	0.50
1:T:398:VAL:HG11	1:T:415:TRP:CE3	2.46	0.50
1:A:65:VAL:HG11	1:A:100:TYR:HB2	1.93	0.50
1:AB:325:VAL:HG13	1:AB:325:VAL:O	2.11	0.50
1:AB:334:LEU:HD12	1:AB:357:TRP:NE1	2.27	0.50
1:BA:52:THR:O	1:BA:109:ILE:HD11	2.11	0.50
1:D:390:VAL:O	1:D:390:VAL:HG23	2.11	0.50
1:D:497:VAL:HG13	1:D:497:VAL:O	2.12	0.50
1:EB:398:VAL:HG11	1:EB:415:TRP:CE3	2.46	0.50
1:FB:390:VAL:O	1:FB:390:VAL:HG23	2.11	0.50
1:FB:575:ILE:HG23	1:FB:603:VAL:HG22	1.92	0.50
1:HA:65:VAL:HG11	1:HA:100:TYR:HB2	1.93	0.50
1:KB:144:LEU:HD12	1:KB:144:LEU:O	2.11	0.50
1:LA:144:LEU:HD12	1:LA:144:LEU:O	2.11	0.50
1:LA:398:VAL:HG11	1:LA:415:TRP:CE3	2.45	0.50
1:OB:52:THR:O	1:OB:109:ILE:HD11	2.12	0.50
1:OB:144:LEU:HD12	1:OB:144:LEU:O	2.10	0.50
1:PA:144:LEU:HD12	1:PA:144:LEU:O	2.11	0.50
1:SA:398:VAL:HG11	1:SA:415:TRP:CE3	2.46	0.50
1:XB:390:VAL:HG23	1:XB:390:VAL:O	2.11	0.50
1:ZB:144:LEU:HD12	1:ZB:144:LEU:O	2.11	0.50
1:AA:260:VAL:O	1:AA:260:VAL:HG13	2.12	0.50
1:AC:48:MET:HA	1:AC:48:MET:HE3	1.93	0.50

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:AC:759:LEU:HD21	1:ZB:765:VAL:HG22	1.93	0.50
1:CA:325:VAL:O	1:CA:325:VAL:HG13	2.12	0.50
1:E:398:VAL:HG11	1:E:415:TRP:CE3	2.47	0.50
1:IB:17:HIS:HB3	1:IB:44:LEU:HD23	1.94	0.50
1:J:497:VAL:HG13	1:J:497:VAL:O	2.12	0.50
1:JA:529:ILE:HD12	1:JA:583:VAL:HG11	1.93	0.50
1:LB:144:LEU:HD12	1:LB:163:ILE:HD11	1.93	0.50
1:NA:390:VAL:HG23	1:NA:390:VAL:O	2.12	0.50
1:NB:390:VAL:O	1:NB:390:VAL:HG23	2.11	0.50
1:OA:260:VAL:HG13	1:OA:260:VAL:O	2.12	0.50
1:OB:398:VAL:HG11	1:OB:415:TRP:CE3	2.46	0.50
1:OB:575:ILE:HG23	1:OB:603:VAL:HG22	1.93	0.50
1:PB:390:VAL:HG23	1:PB:390:VAL:O	2.10	0.50
1:PB:765:VAL:HG22	1:QB:759:LEU:HD21	1.94	0.50
1:Q:325:VAL:HG13	1:Q:325:VAL:O	2.12	0.50
1:RA:529:ILE:HD12	1:RA:583:VAL:HG11	1.92	0.50
1:S:398:VAL:HG11	1:S:415:TRP:CE3	2.46	0.50
1:SA:65:VAL:HG11	1:SA:100:TYR:HB2	1.91	0.50
1:TB:529:ILE:HD12	1:TB:583:VAL:HG11	1.93	0.50
1:UA:398:VAL:HG11	1:UA:415:TRP:CE3	2.47	0.50
1:W:390:VAL:O	1:W:390:VAL:HG23	2.11	0.50
1:XB:325:VAL:O	1:XB:325:VAL:HG13	2.12	0.50
1:Y:325:VAL:HG13	1:Y:325:VAL:O	2.12	0.50
1:YA:575:ILE:HG23	1:YA:603:VAL:HG22	1.92	0.50
1:AB:52:THR:O	1:AB:109:ILE:HD11	2.12	0.50
1:AC:390:VAL:HG23	1:AC:390:VAL:O	2.12	0.50
1:DA:497:VAL:O	1:DA:497:VAL:HG13	2.12	0.50
1:EB:325:VAL:O	1:EB:325:VAL:HG13	2.12	0.50
1:EB:497:VAL:O	1:EB:497:VAL:HG13	2.12	0.50
1:OA:575:ILE:HG23	1:OA:603:VAL:HG22	1.92	0.50
1:OB:325:VAL:O	1:OB:325:VAL:HG13	2.11	0.50
1:QA:497:VAL:O	1:QA:497:VAL:HG13	2.11	0.50
1:QB:325:VAL:HG13	1:QB:325:VAL:O	2.11	0.50
1:R:260:VAL:HG13	1:R:260:VAL:O	2.12	0.50
1:R:325:VAL:O	1:R:325:VAL:HG13	2.11	0.50
1:RA:144:LEU:O	1:RA:144:LEU:HD12	2.12	0.50
1:RA:523:PHE:CD2	1:RA:568:VAL:HG23	2.47	0.50
1:RB:325:VAL:O	1:RB:325:VAL:HG13	2.11	0.50
1:S:325:VAL:O	1:S:325:VAL:HG13	2.11	0.50
1:VA:230:ARG:HB2	1:VA:268:LEU:HD11	1.93	0.50
1:W:737:GLY:O	1:W:741:VAL:HG23	2.11	0.50

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:Z:390:VAL:HG23	1:Z:390:VAL:O	2.10	0.50
1:B:144:LEU:HD12	1:B:144:LEU:O	2.12	0.50
1:C:497:VAL:O	1:C:497:VAL:HG13	2.12	0.50
1:D:144:LEU:O	1:D:144:LEU:HD12	2.11	0.50
1:EA:765:VAL:HG22	1:FA:759:LEU:HD21	1.94	0.50
1:G:325:VAL:HG13	1:G:325:VAL:O	2.11	0.50
1:GA:325:VAL:HG13	1:GA:325:VAL:O	2.11	0.50
1:GA:390:VAL:O	1:GA:390:VAL:HG23	2.11	0.50
1:GA:539:LEU:HD21	1:GA:599:ILE:HD11	1.94	0.50
1:GB:260:VAL:HG13	1:GB:260:VAL:O	2.12	0.50
1:K:30:VAL:CG2	1:K:50:MET:HE3	2.42	0.50
1:UB:325:VAL:O	1:UB:325:VAL:HG13	2.12	0.50
1:XA:390:VAL:HG23	1:XA:390:VAL:O	2.10	0.50
1:XA:497:VAL:HG13	1:XA:497:VAL:O	2.12	0.50
1:XB:778:GLU:HA	1:XB:781:VAL:HG22	1.93	0.50
1:YB:778:GLU:HA	1:YB:781:VAL:HG22	1.93	0.50
1:Z:497:VAL:HG13	1:Z:497:VAL:O	2.10	0.50
1:Z:575:ILE:HG23	1:Z:603:VAL:HG22	1.92	0.50
1:AA:325:VAL:HG13	1:AA:325:VAL:O	2.11	0.50
1:BB:302:VAL:HG11	1:BB:306:LYS:HZ2	1.76	0.50
1:BB:325:VAL:O	1:BB:325:VAL:HG13	2.12	0.50
1:GA:366:VAL:O	1:GA:366:VAL:HG13	2.10	0.50
1:H:778:GLU:HA	1:H:781:VAL:HG22	1.94	0.50
1:HB:398:VAL:HG11	1:HB:415:TRP:CE3	2.46	0.50
1:J:529:ILE:HD12	1:J:583:VAL:HG11	1.93	0.50
1:JA:390:VAL:O	1:JA:390:VAL:HG23	2.11	0.50
1:LA:497:VAL:O	1:LA:497:VAL:HG13	2.12	0.50
1:LB:65:VAL:HG23	1:LB:67:ARG:HG3	1.94	0.50
1:MB:778:GLU:HA	1:MB:781:VAL:HG22	1.94	0.50
1:NB:325:VAL:HG13	1:NB:325:VAL:O	2.11	0.50
1:O:65:VAL:HG11	1:O:100:TYR:HB2	1.93	0.50
1:PB:17:HIS:HB3	1:PB:44:LEU:HD23	1.94	0.50
1:Q:398:VAL:HG11	1:Q:415:TRP:CE3	2.46	0.50
1:QB:765:VAL:HG22	1:RB:759:LEU:HD21	1.93	0.50
1:S:390:VAL:HG23	1:S:390:VAL:O	2.11	0.50
1:TA:390:VAL:HG23	1:TA:390:VAL:O	2.11	0.50
1:TB:366:VAL:O	1:TB:366:VAL:HG13	2.11	0.50
1:TB:778:GLU:HA	1:TB:781:VAL:HG22	1.94	0.50
1:UB:144:LEU:HD12	1:UB:144:LEU:O	2.10	0.50
1:WB:390:VAL:HG23	1:WB:390:VAL:O	2.11	0.50
1:X:230:ARG:HB2	1:X:268:LEU:HD11	1.93	0.50

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:XA:398:VAL:HG11	1:XA:415:TRP:CE3	2.46	0.50
1:Y:575:ILE:HG23	1:Y:603:VAL:HG22	1.92	0.50
1:YB:765:VAL:HG22	1:ZB:759:LEU:HD21	1.92	0.50
1:B:765:VAL:HG22	1:C:759:LEU:HD21	1.94	0.50
1:BB:334:LEU:HD12	1:BB:357:TRP:NE1	2.26	0.50
1:C:778:GLU:HA	1:C:781:VAL:HG22	1.93	0.50
1:D:325:VAL:HG13	1:D:325:VAL:O	2.11	0.50
1:DA:390:VAL:HG23	1:DA:390:VAL:O	2.11	0.50
1:DB:17:HIS:CB	1:DB:44:LEU:HD23	2.42	0.50
1:DB:92:LEU:HD12	1:DB:92:LEU:N	2.26	0.50
1:FB:765:VAL:HG22	1:GB:759:LEU:HD21	1.94	0.50
1:J:778:GLU:HA	1:J:781:VAL:HG22	1.94	0.50
1:K:737:GLY:O	1:K:741:VAL:HG23	2.12	0.50
1:L:65:VAL:HG23	1:L:67:ARG:HG3	1.94	0.50
1:L:778:GLU:HA	1:L:781:VAL:HG22	1.94	0.50
1:LB:778:GLU:HA	1:LB:781:VAL:HG22	1.93	0.50
1:M:390:VAL:HG23	1:M:390:VAL:O	2.12	0.50
1:M:538:GLN:HB2	1:M:646:VAL:HG22	1.93	0.50
1:MA:325:VAL:O	1:MA:325:VAL:HG13	2.11	0.50
1:N:230:ARG:HB2	1:N:268:LEU:HD11	1.93	0.50
1:PA:529:ILE:HD12	1:PA:583:VAL:HG11	1.94	0.50
1:QB:260:VAL:HG13	1:QB:260:VAL:O	2.12	0.50
1:QB:778:GLU:HA	1:QB:781:VAL:HG22	1.94	0.50
1:T:390:VAL:O	1:T:390:VAL:HG23	2.11	0.50
1:TA:765:VAL:HG22	1:UA:759:LEU:HD21	1.93	0.50
1:UB:398:VAL:HG11	1:UB:415:TRP:CE3	2.47	0.50
1:X:260:VAL:O	1:X:260:VAL:HG13	2.12	0.50
1:YA:260:VAL:O	1:YA:260:VAL:HG13	2.12	0.50
1:YA:765:VAL:HG22	1:ZA:759:LEU:HD21	1.94	0.50
1:ZB:334:LEU:HD12	1:ZB:357:TRP:NE1	2.27	0.50
1:EA:398:VAL:HG11	1:EA:415:TRP:CE3	2.47	0.50
1:EB:390:VAL:HG23	1:EB:390:VAL:O	2.12	0.50
1:FB:325:VAL:HG13	1:FB:325:VAL:O	2.11	0.50
1:JA:538:GLN:HB2	1:JA:646:VAL:HG22	1.93	0.50
1:KA:260:VAL:O	1:KA:260:VAL:HG13	2.12	0.50
1:KA:737:GLY:O	1:KA:741:VAL:HG23	2.12	0.50
1:M:52:THR:O	1:M:109:ILE:HD11	2.12	0.50
1:MA:144:LEU:HD12	1:MA:144:LEU:O	2.11	0.50
1:PA:390:VAL:HG23	1:PA:390:VAL:O	2.11	0.50
1:RB:778:GLU:HA	1:RB:781:VAL:HG22	1.94	0.50
1:TA:575:ILE:HG23	1:TA:603:VAL:HG22	1.93	0.50

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:V:390:VAL:HG23	1:V:390:VAL:O	2.12	0.50
1:VA:17:HIS:CB	1:VA:44:LEU:HD23	2.42	0.50
1:WB:325:VAL:O	1:WB:325:VAL:HG13	2.12	0.50
1:X:92:LEU:HD12	1:X:92:LEU:N	2.27	0.50
1:XA:529:ILE:HD12	1:XA:583:VAL:HG11	1.93	0.50
1:XB:398:VAL:HG11	1:XB:415:TRP:CE3	2.47	0.50
1:YA:325:VAL:O	1:YA:325:VAL:HG13	2.12	0.50
1:YB:325:VAL:HG13	1:YB:325:VAL:O	2.11	0.50
1:ZA:17:HIS:HB3	1:ZA:44:LEU:HD23	1.94	0.50
1:ZA:497:VAL:HG13	1:ZA:497:VAL:O	2.12	0.50
1:AA:737:GLY:O	1:AA:741:VAL:HG23	2.12	0.49
1:DA:260:VAL:O	1:DA:260:VAL:HG13	2.12	0.49
1:DA:325:VAL:HG13	1:DA:325:VAL:O	2.11	0.49
1:E:48:MET:HA	1:E:48:MET:CE	2.41	0.49
1:E:230:ARG:HB2	1:E:268:LEU:HD11	1.94	0.49
1:GB:390:VAL:HG23	1:GB:390:VAL:O	2.11	0.49
1:HA:334:LEU:HD12	1:HA:357:TRP:NE1	2.27	0.49
1:IB:390:VAL:O	1:IB:390:VAL:HG23	2.12	0.49
1:J:325:VAL:O	1:J:325:VAL:HG13	2.12	0.49
1:JA:334:LEU:HD12	1:JA:357:TRP:NE1	2.27	0.49
1:KA:30:VAL:CG2	1:KA:50:MET:HE3	2.42	0.49
1:MB:398:VAL:HG11	1:MB:415:TRP:CE3	2.47	0.49
1:N:325:VAL:HG13	1:N:325:VAL:O	2.12	0.49
1:NB:737:GLY:O	1:NB:741:VAL:HG23	2.12	0.49
1:OA:529:ILE:HD12	1:OA:583:VAL:HG11	1.93	0.49
1:QA:575:ILE:HG23	1:QA:603:VAL:HG22	1.93	0.49
1:QB:52:THR:O	1:QB:109:ILE:HD11	2.12	0.49
1:V:230:ARG:HB2	1:V:268:LEU:HD11	1.93	0.49
1:VB:497:VAL:O	1:VB:497:VAL:HG13	2.12	0.49
1:XB:30:VAL:CG2	1:XB:50:MET:HE3	2.42	0.49
1:Y:778:GLU:HA	1:Y:781:VAL:HG22	1.93	0.49
1:ZA:575:ILE:HG23	1:ZA:603:VAL:HG22	1.93	0.49
1:A:52:THR:O	1:A:109:ILE:HD11	2.12	0.49
1:AC:575:ILE:HG23	1:AC:603:VAL:HG22	1.93	0.49
1:B:778:GLU:HA	1:B:781:VAL:HG22	1.94	0.49
1:BB:260:VAL:O	1:BB:260:VAL:HG13	2.13	0.49
1:C:52:THR:O	1:C:109:ILE:HD11	2.12	0.49
1:FB:52:THR:O	1:FB:109:ILE:HD11	2.12	0.49
1:H:529:ILE:HD12	1:H:583:VAL:HG11	1.95	0.49
1:JB:260:VAL:O	1:JB:260:VAL:HG13	2.12	0.49
1:KB:778:GLU:HA	1:KB:781:VAL:HG22	1.95	0.49

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:NB:260:VAL:HG13	1:NB:260:VAL:O	2.12	0.49
1:OB:390:VAL:HG23	1:OB:390:VAL:O	2.11	0.49
1:RA:260:VAL:HG13	1:RA:260:VAL:O	2.12	0.49
1:S:260:VAL:O	1:S:260:VAL:HG13	2.12	0.49
1:UA:52:THR:O	1:UA:109:ILE:HD11	2.12	0.49
1:UB:737:GLY:O	1:UB:741:VAL:HG23	2.12	0.49
1:Z:398:VAL:HG11	1:Z:415:TRP:CE3	2.47	0.49
1:ZB:325:VAL:O	1:ZB:325:VAL:HG13	2.12	0.49
1:AC:65:VAL:HG23	1:AC:67:ARG:HG3	1.94	0.49
1:AC:737:GLY:O	1:AC:741:VAL:HG23	2.12	0.49
1:B:529:ILE:HD12	1:B:583:VAL:HG11	1.94	0.49
1:BA:765:VAL:HG22	1:CA:759:LEU:HD21	1.95	0.49
1:D:778:GLU:HA	1:D:781:VAL:HG22	1.94	0.49
1:EA:538:GLN:HB2	1:EA:646:VAL:HG22	1.95	0.49
1:EB:260:VAL:O	1:EB:260:VAL:HG13	2.12	0.49
1:F:30:VAL:HG22	1:F:50:MET:HE2	1.94	0.49
1:FB:539:LEU:HD12	1:FB:540:GLN:H	1.76	0.49
1:G:260:VAL:HG13	1:G:260:VAL:O	2.13	0.49
1:H:44:LEU:H	1:H:44:LEU:HD22	1.77	0.49
1:I:325:VAL:HG13	1:I:325:VAL:O	2.12	0.49
1:I:737:GLY:O	1:I:741:VAL:HG23	2.12	0.49
1:IA:260:VAL:O	1:IA:260:VAL:HG13	2.13	0.49
1:J:390:VAL:HG23	1:J:390:VAL:O	2.11	0.49
1:JA:575:ILE:HG23	1:JA:603:VAL:HG22	1.94	0.49
1:K:260:VAL:O	1:K:260:VAL:HG13	2.12	0.49
1:KB:65:VAL:HG23	1:KB:67:ARG:HG3	1.94	0.49
1:KB:325:VAL:HG13	1:KB:325:VAL:O	2.12	0.49
1:M:778:GLU:HA	1:M:781:VAL:HG22	1.94	0.49
1:MB:30:VAL:CG2	1:MB:50:MET:HE3	2.42	0.49
1:NA:737:GLY:O	1:NA:741:VAL:HG23	2.13	0.49
1:NB:92:LEU:N	1:NB:92:LEU:HD12	2.27	0.49
1:QB:523:PHE:CD2	1:QB:568:VAL:HG23	2.48	0.49
1:T:778:GLU:HA	1:T:781:VAL:HG22	1.94	0.49
1:TB:539:LEU:HD21	1:TB:599:ILE:HD11	1.93	0.49
1:UA:778:GLU:HA	1:UA:781:VAL:HG22	1.94	0.49
1:V:497:VAL:O	1:V:497:VAL:HG13	2.12	0.49
1:VB:390:VAL:HG23	1:VB:390:VAL:O	2.11	0.49
1:X:325:VAL:O	1:X:325:VAL:HG13	2.12	0.49
1:ZA:390:VAL:O	1:ZA:390:VAL:HG23	2.10	0.49
1:A:260:VAL:HG13	1:A:260:VAL:O	2.12	0.49
1:BB:778:GLU:HA	1:BB:781:VAL:HG22	1.94	0.49

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:E:497:VAL:HG13	1:E:497:VAL:O	2.12	0.49
1:FA:17:HIS:HB3	1:FA:44:LEU:HD23	1.95	0.49
1:FA:260:VAL:O	1:FA:260:VAL:HG13	2.13	0.49
1:FA:390:VAL:HG23	1:FA:390:VAL:O	2.11	0.49
1:FA:398:VAL:HG11	1:FA:415:TRP:CE3	2.48	0.49
1:H:65:VAL:HG23	1:H:67:ARG:HG3	1.95	0.49
1:IB:144:LEU:N	1:IB:144:LEU:HD23	2.27	0.49
1:IB:778:GLU:HA	1:IB:781:VAL:HG22	1.95	0.49
1:K:390:VAL:HG23	1:K:390:VAL:O	2.12	0.49
1:KA:575:ILE:HG23	1:KA:603:VAL:HG22	1.94	0.49
1:LA:778:GLU:HA	1:LA:781:VAL:HG22	1.93	0.49
1:MA:778:GLU:HA	1:MA:781:VAL:HG22	1.95	0.49
1:O:144:LEU:O	1:O:144:LEU:HD12	2.12	0.49
1:P:398:VAL:HG11	1:P:415:TRP:CE3	2.47	0.49
1:PB:497:VAL:O	1:PB:497:VAL:HG13	2.12	0.49
1:PB:778:GLU:HA	1:PB:781:VAL:HG22	1.95	0.49
1:QA:390:VAL:HG23	1:QA:390:VAL:O	2.11	0.49
1:R:390:VAL:HG23	1:R:390:VAL:O	2.11	0.49
1:TA:334:LEU:HD12	1:TA:357:TRP:NE1	2.27	0.49
1:WA:30:VAL:CG2	1:WA:50:MET:HE3	2.43	0.49
1:WA:334:LEU:HD12	1:WA:357:TRP:NE1	2.28	0.49
1:YB:497:VAL:O	1:YB:497:VAL:HG13	2.12	0.49
1:ZB:65:VAL:HG23	1:ZB:67:ARG:HG3	1.95	0.49
1:AB:65:VAL:HG11	1:AB:100:TYR:HB2	1.95	0.49
1:B:390:VAL:HG23	1:B:390:VAL:O	2.12	0.49
1:BB:390:VAL:HG23	1:BB:390:VAL:O	2.12	0.49
1:DA:778:GLU:HA	1:DA:781:VAL:HG22	1.94	0.49
1:DB:778:GLU:HA	1:DB:781:VAL:HG22	1.94	0.49
1:E:334:LEU:HD12	1:E:357:TRP:NE1	2.27	0.49
1:GB:778:GLU:HA	1:GB:781:VAL:HG22	1.95	0.49
1:HA:737:GLY:O	1:HA:741:VAL:HG23	2.13	0.49
1:J:758:GLU:O	1:J:762:VAL:HG22	2.13	0.49
1:K:65:VAL:HG23	1:K:67:ARG:HG3	1.94	0.49
1:K:778:GLU:HA	1:K:781:VAL:HG22	1.94	0.49
1:LB:325:VAL:O	1:LB:325:VAL:HG13	2.11	0.49
1:NA:325:VAL:O	1:NA:325:VAL:HG13	2.11	0.49
1:PB:44:LEU:H	1:PB:44:LEU:HD22	1.76	0.49
1:Q:390:VAL:HG23	1:Q:390:VAL:O	2.12	0.49
1:Q:778:GLU:HA	1:Q:781:VAL:HG22	1.94	0.49
1:R:539:LEU:HD21	1:R:599:ILE:HD11	1.94	0.49
1:SA:575:ILE:HG23	1:SA:603:VAL:HG22	1.93	0.49

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:SB:260:VAL:O	1:SB:260:VAL:HG13	2.13	0.49
1:T:575:ILE:HG23	1:T:603:VAL:HG22	1.93	0.49
1:TA:260:VAL:O	1:TA:260:VAL:HG13	2.13	0.49
1:UA:390:VAL:HG23	1:UA:390:VAL:O	2.12	0.49
1:WA:260:VAL:HG13	1:WA:260:VAL:O	2.13	0.49
1:YA:778:GLU:HA	1:YA:781:VAL:HG22	1.93	0.49
1:YB:65:VAL:HG11	1:YB:100:TYR:HB2	1.93	0.49
1:YB:529:ILE:HD12	1:YB:583:VAL:HG11	1.94	0.49
1:A:529:ILE:HD12	1:A:583:VAL:HG11	1.93	0.49
1:AB:260:VAL:HG13	1:AB:260:VAL:O	2.12	0.49
1:AB:497:VAL:HG13	1:AB:497:VAL:O	2.12	0.49
1:B:523:PHE:CD2	1:B:568:VAL:HG23	2.46	0.49
1:CA:306:LYS:HZ3	1:CA:308:PHE:HB3	1.77	0.49
1:CB:778:GLU:HA	1:CB:781:VAL:HG22	1.93	0.49
1:FB:260:VAL:HG13	1:FB:260:VAL:O	2.12	0.49
1:FB:778:GLU:HA	1:FB:781:VAL:HG22	1.94	0.49
1:G:778:GLU:HA	1:G:781:VAL:HG22	1.93	0.49
1:HB:9:ARG:NH1	1:IB:19:LEU:HD11	2.28	0.49
1:I:48:MET:HA	1:I:48:MET:HE3	1.95	0.49
1:IA:758:GLU:O	1:IA:762:VAL:HG22	2.13	0.49
1:J:260:VAL:HG13	1:J:260:VAL:O	2.12	0.49
1:KA:398:VAL:HG11	1:KA:415:TRP:CE3	2.47	0.49
1:O:30:VAL:CG2	1:O:50:MET:HE3	2.43	0.49
1:QA:778:GLU:HA	1:QA:781:VAL:HG22	1.94	0.49
1:RB:765:VAL:HG22	1:SB:759:LEU:HD21	1.93	0.49
1:SA:230:ARG:HB2	1:SA:268:LEU:HD11	1.93	0.49
1:UA:575:ILE:HG23	1:UA:603:VAL:HG22	1.93	0.49
1:UB:260:VAL:O	1:UB:260:VAL:HG13	2.12	0.49
1:VA:765:VAL:HG22	1:WA:759:LEU:HD21	1.95	0.49
1:VB:758:GLU:O	1:VB:762:VAL:HG22	2.12	0.49
1:WA:765:VAL:HG22	1:XA:759:LEU:HD21	1.94	0.49
1:A:759:LEU:HD21	1:NA:765:VAL:HG22	1.93	0.49
1:AB:144:LEU:O	1:AB:144:LEU:HD12	2.11	0.49
1:AB:575:ILE:HG23	1:AB:603:VAL:HG22	1.93	0.49
1:AC:778:GLU:HA	1:AC:781:VAL:HG22	1.94	0.49
1:B:260:VAL:HG13	1:B:260:VAL:O	2.13	0.49
1:C:325:VAL:HG13	1:C:325:VAL:O	2.11	0.49
1:DB:398:VAL:HG12	1:DB:491:PRO:HB3	1.93	0.49
1:FA:778:GLU:HA	1:FA:781:VAL:HG22	1.94	0.49
1:G:575:ILE:HG23	1:G:603:VAL:HG22	1.94	0.49
1:GA:17:HIS:HB3	1:GA:44:LEU:HD23	1.95	0.49

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:HB:390:VAL:HG23	1:HB:390:VAL:O	2.11	0.49
1:IA:325:VAL:HG13	1:IA:325:VAL:O	2.12	0.49
1:KA:325:VAL:O	1:KA:325:VAL:HG13	2.12	0.49
1:L:575:ILE:HG23	1:L:603:VAL:HG22	1.94	0.49
1:LB:765:VAL:HG22	1:MB:759:LEU:HD21	1.95	0.49
1:NA:778:GLU:HA	1:NA:781:VAL:HG22	1.94	0.49
1:NB:778:GLU:HA	1:NB:781:VAL:HG22	1.95	0.49
1:O:334:LEU:HD12	1:O:357:TRP:NE1	2.28	0.49
1:P:260:VAL:O	1:P:260:VAL:HG13	2.13	0.49
1:P:325:VAL:O	1:P:325:VAL:HG13	2.13	0.49
1:PA:260:VAL:HG13	1:PA:260:VAL:O	2.12	0.49
1:UB:65:VAL:HG11	1:UB:100:TYR:HB2	1.93	0.49
1:WB:778:GLU:HA	1:WB:781:VAL:HG22	1.94	0.49
1:ZA:778:GLU:HA	1:ZA:781:VAL:HG22	1.94	0.49
1:A:144:LEU:HD23	1:A:144:LEU:N	2.28	0.49
1:BA:390:VAL:O	1:BA:390:VAL:HG23	2.11	0.49
1:C:575:ILE:HG23	1:C:603:VAL:HG22	1.93	0.49
1:CB:260:VAL:O	1:CB:260:VAL:HG13	2.12	0.49
1:D:260:VAL:HG13	1:D:260:VAL:O	2.12	0.49
1:DB:260:VAL:HG13	1:DB:260:VAL:O	2.13	0.49
1:E:260:VAL:O	1:E:260:VAL:HG13	2.13	0.49
1:FA:737:GLY:O	1:FA:741:VAL:HG23	2.12	0.49
1:G:144:LEU:HD12	1:G:144:LEU:O	2.11	0.49
1:HB:778:GLU:HA	1:HB:781:VAL:HG22	1.94	0.49
1:IA:497:VAL:O	1:IA:497:VAL:HG13	2.12	0.49
1:KB:260:VAL:O	1:KB:260:VAL:HG13	2.12	0.49
1:L:765:VAL:HG22	1:M:759:LEU:HD21	1.95	0.49
1:LB:737:GLY:O	1:LB:741:VAL:HG23	2.13	0.49
1:M:65:VAL:HG11	1:M:100:TYR:HB2	1.94	0.49
1:N:778:GLU:HA	1:N:781:VAL:HG22	1.94	0.49
1:O:390:VAL:O	1:O:390:VAL:HG23	2.13	0.49
1:OB:765:VAL:HG22	1:PB:759:LEU:HD21	1.95	0.49
1:PA:334:LEU:HD12	1:PA:357:TRP:NE1	2.28	0.49
1:QB:575:ILE:HG23	1:QB:603:VAL:HG22	1.93	0.49
1:RA:390:VAL:O	1:RA:390:VAL:HG23	2.11	0.49
1:SA:390:VAL:HG23	1:SA:390:VAL:O	2.11	0.49
1:SA:497:VAL:HG13	1:SA:497:VAL:O	2.12	0.49
1:SA:778:GLU:HA	1:SA:781:VAL:HG22	1.94	0.49
1:SB:778:GLU:HA	1:SB:781:VAL:HG22	1.94	0.49
1:TA:758:GLU:O	1:TA:762:VAL:HG22	2.12	0.49
1:VB:65:VAL:HG11	1:VB:100:TYR:HB2	1.95	0.49

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:VB:778:GLU:HA	1:VB:781:VAL:HG22	1.95	0.49
1:WB:260:VAL:HG13	1:WB:260:VAL:O	2.13	0.49
1:XB:737:GLY:O	1:XB:741:VAL:HG23	2.13	0.49
1:ZB:17:HIS:HB3	1:ZB:44:LEU:HD23	1.95	0.49
1:A:65:VAL:HG23	1:A:67:ARG:HG3	1.95	0.49
1:C:260:VAL:O	1:C:260:VAL:HG13	2.13	0.49
1:DA:144:LEU:HD23	1:DA:144:LEU:N	2.28	0.49
1:E:65:VAL:HG23	1:E:67:ARG:HG3	1.95	0.49
1:H:17:HIS:HB3	1:H:44:LEU:HD23	1.95	0.49
1:HA:260:VAL:HG13	1:HA:260:VAL:O	2.13	0.49
1:I:65:VAL:HG23	1:I:67:ARG:HG3	1.95	0.49
1:K:765:VAL:HG22	1:L:759:LEU:HD21	1.93	0.49
1:M:260:VAL:HG13	1:M:260:VAL:O	2.13	0.49
1:N:48:MET:HA	1:N:48:MET:HE3	1.95	0.49
1:N:260:VAL:O	1:N:260:VAL:HG13	2.13	0.49
1:O:260:VAL:HG13	1:O:260:VAL:O	2.12	0.49
1:P:92:LEU:HD12	1:P:92:LEU:N	2.28	0.49
1:Q:575:ILE:HG23	1:Q:603:VAL:HG22	1.95	0.49
1:QA:765:VAL:HG22	1:RA:759:LEU:HD21	1.95	0.49
1:RA:48:MET:HA	1:RA:48:MET:HE3	1.94	0.49
1:RA:765:VAL:HG22	1:SA:759:LEU:HD21	1.95	0.49
1:S:334:LEU:HD12	1:S:357:TRP:NE1	2.28	0.49
1:S:737:GLY:O	1:S:741:VAL:HG23	2.12	0.49
1:SA:325:VAL:O	1:SA:325:VAL:HG13	2.12	0.49
1:UB:390:VAL:HG23	1:UB:390:VAL:O	2.13	0.49
1:XA:159:VAL:O	1:XA:159:VAL:HG12	2.13	0.49
1:XB:260:VAL:O	1:XB:260:VAL:HG13	2.12	0.49
1:ZA:260:VAL:HG13	1:ZA:260:VAL:O	2.13	0.49
1:BB:497:VAL:HG13	1:BB:497:VAL:O	2.11	0.49
1:C:398:VAL:HG12	1:C:491:PRO:HB3	1.94	0.49
1:CB:30:VAL:HG21	1:CB:50:MET:HE3	1.95	0.49
1:D:575:ILE:HG23	1:D:603:VAL:HG22	1.94	0.49
1:DA:523:PHE:CD2	1:DA:568:VAL:HG23	2.48	0.49
1:DA:765:VAL:HG22	1:EA:759:LEU:HD21	1.94	0.49
1:DB:398:VAL:HG11	1:DB:415:TRP:CE3	2.47	0.49
1:DB:529:ILE:CD1	1:DB:583:VAL:HG21	2.43	0.49
1:F:334:LEU:HD12	1:F:357:TRP:NE1	2.28	0.49
1:F:575:ILE:HG23	1:F:603:VAL:HG22	1.94	0.49
1:F:778:GLU:HA	1:F:781:VAL:HG22	1.95	0.49
1:GB:575:ILE:HG23	1:GB:603:VAL:HG22	1.95	0.49
1:HA:398:VAL:HG11	1:HA:415:TRP:CE3	2.47	0.49

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:HB:529:ILE:HD12	1:HB:583:VAL:HG11	1.94	0.49
1:J:65:VAL:HG23	1:J:67:ARG:HG3	1.95	0.49
1:J:737:GLY:O	1:J:741:VAL:HG23	2.13	0.49
1:JA:52:THR:O	1:JA:109:ILE:HD11	2.13	0.49
1:KA:523:PHE:CD2	1:KA:568:VAL:HG23	2.48	0.49
1:LA:529:ILE:HD12	1:LA:583:VAL:HG11	1.94	0.49
1:MA:17:HIS:HB3	1:MA:44:LEU:HD23	1.95	0.49
1:PA:778:GLU:HA	1:PA:781:VAL:HG22	1.94	0.49
1:QA:260:VAL:O	1:QA:260:VAL:HG13	2.13	0.49
1:RA:778:GLU:HA	1:RA:781:VAL:HG22	1.94	0.49
1:S:144:LEU:N	1:S:144:LEU:HD23	2.28	0.49
1:SA:334:LEU:HD12	1:SA:357:TRP:NE1	2.28	0.49
1:UA:260:VAL:HG13	1:UA:260:VAL:O	2.13	0.49
1:UA:523:PHE:CD2	1:UA:568:VAL:HG23	2.48	0.49
1:V:765:VAL:HG22	1:W:759:LEU:HD21	1.95	0.49
1:WA:48:MET:HA	1:WA:48:MET:HE3	1.95	0.49
1:WA:325:VAL:HG13	1:WA:325:VAL:O	2.12	0.49
1:WA:497:VAL:O	1:WA:497:VAL:HG13	2.12	0.49
1:X:65:VAL:HG11	1:X:100:TYR:HB2	1.94	0.49
1:XB:65:VAL:HG23	1:XB:67:ARG:HG3	1.95	0.49
1:YA:497:VAL:O	1:YA:497:VAL:HG13	2.13	0.49
1:AA:390:VAL:HG23	1:AA:390:VAL:O	2.11	0.48
1:AB:778:GLU:HA	1:AB:781:VAL:HG22	1.94	0.48
1:D:523:PHE:CD2	1:D:568:VAL:HG23	2.48	0.48
1:DB:765:VAL:HG22	1:EB:759:LEU:HD21	1.94	0.48
1:F:497:VAL:O	1:F:497:VAL:HG13	2.12	0.48
1:FB:366:VAL:O	1:FB:366:VAL:HG13	2.13	0.48
1:GA:260:VAL:HG13	1:GA:260:VAL:O	2.13	0.48
1:GA:778:GLU:HA	1:GA:781:VAL:HG22	1.94	0.48
1:H:92:LEU:HD12	1:H:92:LEU:N	2.28	0.48
1:HB:260:VAL:O	1:HB:260:VAL:HG13	2.13	0.48
1:IA:390:VAL:HG23	1:IA:390:VAL:O	2.11	0.48
1:JA:778:GLU:HA	1:JA:781:VAL:HG22	1.94	0.48
1:MA:260:VAL:HG13	1:MA:260:VAL:O	2.13	0.48
1:N:737:GLY:O	1:N:741:VAL:HG23	2.13	0.48
1:O:65:VAL:HG23	1:O:67:ARG:HG3	1.95	0.48
1:P:65:VAL:HG23	1:P:67:ARG:HG3	1.96	0.48
1:P:398:VAL:HG12	1:P:491:PRO:HB3	1.94	0.48
1:R:497:VAL:HG13	1:R:497:VAL:O	2.12	0.48
1:R:778:GLU:HA	1:R:781:VAL:HG22	1.94	0.48
1:RA:575:ILE:HG23	1:RA:603:VAL:HG22	1.94	0.48

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:RB:30:VAL:HG22	1:RB:50:MET:HE2	1.95	0.48
1:SB:398:VAL:HG11	1:SB:415:TRP:CE3	2.48	0.48
1:UA:144:LEU:HD12	1:UA:144:LEU:O	2.12	0.48
1:WB:334:LEU:HD12	1:WB:357:TRP:NE1	2.28	0.48
1:X:52:THR:O	1:X:109:ILE:HD11	2.13	0.48
1:XA:758:GLU:O	1:XA:762:VAL:HG22	2.13	0.48
1:Y:260:VAL:O	1:Y:260:VAL:HG13	2.13	0.48
1:Y:765:VAL:HG22	1:Z:759:LEU:HD21	1.95	0.48
1:ZB:260:VAL:HG13	1:ZB:260:VAL:O	2.13	0.48
1:AA:398:VAL:HG11	1:AA:415:TRP:CE3	2.48	0.48
1:AB:65:VAL:HG23	1:AB:67:ARG:HG3	1.95	0.48
1:AB:390:VAL:O	1:AB:390:VAL:HG23	2.12	0.48
1:AB:759:LEU:HD21	1:ZA:765:VAL:HG22	1.94	0.48
1:B:334:LEU:HD12	1:B:357:TRP:NE1	2.28	0.48
1:BB:737:GLY:O	1:BB:741:VAL:HG23	2.13	0.48
1:C:529:ILE:CD1	1:C:583:VAL:HG21	2.43	0.48
1:DB:325:VAL:O	1:DB:325:VAL:HG13	2.12	0.48
1:EB:778:GLU:HA	1:EB:781:VAL:HG22	1.95	0.48
1:IA:778:GLU:HA	1:IA:781:VAL:HG22	1.95	0.48
1:J:56:ARG:HE	1:K:127:LEU:HD21	1.78	0.48
1:LA:260:VAL:O	1:LA:260:VAL:HG13	2.13	0.48
1:LB:497:VAL:HG13	1:LB:497:VAL:O	2.14	0.48
1:MB:334:LEU:HD12	1:MB:357:TRP:NE1	2.27	0.48
1:NB:398:VAL:HG11	1:NB:415:TRP:CE3	2.48	0.48
1:NB:497:VAL:O	1:NB:497:VAL:HG13	2.12	0.48
1:P:529:ILE:CD1	1:P:583:VAL:HG21	2.43	0.48
1:Q:92:LEU:N	1:Q:92:LEU:HD12	2.28	0.48
1:SB:575:ILE:HG23	1:SB:603:VAL:HG22	1.94	0.48
1:UB:17:HIS:HB3	1:UB:44:LEU:HD23	1.95	0.48
1:UB:334:LEU:HD12	1:UB:357:TRP:NE1	2.28	0.48
1:XA:92:LEU:HD12	1:XA:92:LEU:N	2.28	0.48
1:XA:778:GLU:HA	1:XA:781:VAL:HG22	1.94	0.48
1:YB:56:ARG:HE	1:ZB:127:LEU:HD21	1.78	0.48
1:YB:539:LEU:HD23	1:YB:540:GLN:N	2.27	0.48
1:ZB:758:GLU:O	1:ZB:762:VAL:HG22	2.13	0.48
1:ZB:778:GLU:HA	1:ZB:781:VAL:HG22	1.94	0.48
1:AC:529:ILE:CD1	1:AC:583:VAL:HG21	2.44	0.48
1:BA:529:ILE:HD12	1:BA:583:VAL:HG11	1.93	0.48
1:C:765:VAL:HG22	1:D:759:LEU:HD21	1.95	0.48
1:CA:260:VAL:O	1:CA:260:VAL:HG13	2.13	0.48
1:CB:334:LEU:HD12	1:CB:357:TRP:NE1	2.28	0.48

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:DB:65:VAL:HG23	1:DB:67:ARG:HG3	1.95	0.48
1:EA:778:GLU:HA	1:EA:781:VAL:HG22	1.94	0.48
1:H:737:GLY:O	1:H:741:VAL:HG23	2.14	0.48
1:IB:92:LEU:HD12	1:IB:92:LEU:N	2.28	0.48
1:M:65:VAL:HG23	1:M:67:ARG:HG3	1.96	0.48
1:MB:260:VAL:O	1:MB:260:VAL:HG13	2.13	0.48
1:NA:260:VAL:O	1:NA:260:VAL:HG13	2.13	0.48
1:PA:765:VAL:HG22	1:QA:759:LEU:HD21	1.94	0.48
1:R:366:VAL:O	1:R:366:VAL:HG13	2.13	0.48
1:SB:758:GLU:O	1:SB:762:VAL:HG22	2.14	0.48
1:TA:330:GLN:C	1:TA:407:MET:HE1	2.39	0.48
1:V:260:VAL:O	1:V:260:VAL:HG13	2.14	0.48
1:VA:260:VAL:O	1:VA:260:VAL:HG13	2.13	0.48
1:W:260:VAL:HG13	1:W:260:VAL:O	2.12	0.48
1:WA:390:VAL:O	1:WA:390:VAL:HG23	2.12	0.48
1:XA:260:VAL:HG13	1:XA:260:VAL:O	2.12	0.48
1:YB:260:VAL:HG13	1:YB:260:VAL:O	2.13	0.48
1:YB:788:ALA:O	1:YB:792:VAL:HG23	2.14	0.48
1:A:758:GLU:O	1:A:762:VAL:HG22	2.13	0.48
1:BA:92:LEU:HD12	1:BA:92:LEU:N	2.29	0.48
1:BA:334:LEU:HD12	1:BA:357:TRP:NE1	2.29	0.48
1:CB:390:VAL:HG23	1:CB:390:VAL:O	2.13	0.48
1:EB:92:LEU:HD12	1:EB:92:LEU:N	2.28	0.48
1:FB:539:LEU:HD12	1:FB:540:GLN:N	2.28	0.48
1:GA:575:ILE:HG23	1:GA:603:VAL:HG22	1.95	0.48
1:GB:398:VAL:HG12	1:GB:491:PRO:HB3	1.95	0.48
1:I:497:VAL:O	1:I:497:VAL:HG13	2.14	0.48
1:K:325:VAL:O	1:K:325:VAL:HG13	2.12	0.48
1:L:260:VAL:HG13	1:L:260:VAL:O	2.13	0.48
1:LA:788:ALA:O	1:LA:792:VAL:HG23	2.13	0.48
1:LB:260:VAL:HG13	1:LB:260:VAL:O	2.13	0.48
1:O:92:LEU:HD12	1:O:92:LEU:N	2.28	0.48
1:O:737:GLY:O	1:O:741:VAL:HG23	2.14	0.48
1:OA:52:THR:O	1:OA:109:ILE:HD11	2.12	0.48
1:S:765:VAL:HG22	1:T:759:LEU:HD21	1.95	0.48
1:S:778:GLU:HA	1:S:781:VAL:HG22	1.94	0.48
1:VB:65:VAL:HG23	1:VB:67:ARG:HG3	1.96	0.48
1:VB:325:VAL:HG13	1:VB:325:VAL:O	2.12	0.48
1:WA:17:HIS:HB3	1:WA:44:LEU:HD23	1.96	0.48
1:X:65:VAL:HG23	1:X:67:ARG:HG3	1.95	0.48
1:XA:765:VAL:HG22	1:YA:759:LEU:HD21	1.95	0.48

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:Z:260:VAL:HG13	1:Z:260:VAL:O	2.13	0.48
1:A:17:HIS:HB3	1:A:44:LEU:HD23	1.95	0.48
1:A:390:VAL:O	1:A:390:VAL:HG23	2.13	0.48
1:AA:52:THR:O	1:AA:109:ILE:HD11	2.12	0.48
1:BA:48:MET:HA	1:BA:48:MET:HE3	1.94	0.48
1:CB:737:GLY:O	1:CB:741:VAL:HG23	2.13	0.48
1:H:260:VAL:O	1:H:260:VAL:HG13	2.13	0.48
1:H:765:VAL:HG22	1:I:759:LEU:HD21	1.95	0.48
1:JA:765:VAL:HG22	1:KA:759:LEU:HD21	1.95	0.48
1:LA:390:VAL:HG23	1:LA:390:VAL:O	2.13	0.48
1:OA:390:VAL:O	1:OA:390:VAL:HG23	2.13	0.48
1:OB:17:HIS:HB3	1:OB:44:LEU:HD23	1.95	0.48
1:OB:92:LEU:HD12	1:OB:92:LEU:N	2.29	0.48
1:P:334:LEU:HD12	1:P:357:TRP:NE1	2.28	0.48
1:P:778:GLU:HA	1:P:781:VAL:HG22	1.94	0.48
1:R:788:ALA:O	1:R:792:VAL:HG23	2.14	0.48
1:TA:497:VAL:O	1:TA:497:VAL:HG13	2.12	0.48
1:WB:575:ILE:HG23	1:WB:603:VAL:HG22	1.94	0.48
1:Y:788:ALA:O	1:Y:792:VAL:HG23	2.14	0.48
1:BA:398:VAL:HG12	1:BA:491:PRO:HB3	1.95	0.48
1:DA:30:VAL:CG2	1:DA:50:MET:HE3	2.43	0.48
1:E:778:GLU:HA	1:E:781:VAL:HG22	1.94	0.48
1:EB:17:HIS:HB3	1:EB:44:LEU:HD23	1.96	0.48
1:F:168:ILE:HG21	1:F:174:LEU:HB2	1.95	0.48
1:GB:765:VAL:HG22	1:HB:759:LEU:HD21	1.94	0.48
1:HA:575:ILE:HG23	1:HA:603:VAL:HG22	1.96	0.48
1:I:334:LEU:HD12	1:I:357:TRP:NE1	2.29	0.48
1:I:390:VAL:HG23	1:I:390:VAL:O	2.13	0.48
1:J:168:ILE:HG21	1:J:174:LEU:HB2	1.95	0.48
1:J:765:VAL:HG22	1:K:759:LEU:HD21	1.95	0.48
1:OB:260:VAL:HG13	1:OB:260:VAL:O	2.14	0.48
1:PA:56:ARG:HE	1:QA:127:LEU:HD21	1.77	0.48
1:PB:65:VAL:HG23	1:PB:67:ARG:HG3	1.96	0.48
1:QB:737:GLY:O	1:QB:741:VAL:HG23	2.13	0.48
1:T:260:VAL:HG13	1:T:260:VAL:O	2.13	0.48
1:T:737:GLY:O	1:T:741:VAL:HG23	2.14	0.48
1:TB:765:VAL:HG22	1:UB:759:LEU:HD21	1.96	0.48
1:WB:65:VAL:HG23	1:WB:67:ARG:HG3	1.96	0.48
1:WB:758:GLU:O	1:WB:762:VAL:HG22	2.13	0.48
1:XB:575:ILE:HG23	1:XB:603:VAL:HG22	1.93	0.48
1:Y:334:LEU:HD12	1:Y:357:TRP:NE1	2.29	0.48

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:YA:65:VAL:HG23	1:YA:67:ARG:HG3	1.94	0.48
1:YA:788:ALA:O	1:YA:792:VAL:HG23	2.14	0.48
1:D:92:LEU:N	1:D:92:LEU:HD12	2.28	0.48
1:D:765:VAL:HG22	1:E:759:LEU:HD21	1.94	0.48
1:E:325:VAL:O	1:E:325:VAL:HG13	2.12	0.48
1:F:758:GLU:O	1:F:762:VAL:HG22	2.13	0.48
1:GA:168:ILE:HG21	1:GA:174:LEU:HB2	1.95	0.48
1:GA:765:VAL:HG22	1:HA:759:LEU:HD21	1.95	0.48
1:GB:497:VAL:HG13	1:GB:497:VAL:O	2.14	0.48
1:HA:758:GLU:O	1:HA:762:VAL:HG22	2.13	0.48
1:HB:334:LEU:HD12	1:HB:357:TRP:NE1	2.29	0.48
1:HB:765:VAL:HG22	1:IB:759:LEU:HD21	1.95	0.48
1:JA:325:VAL:HG13	1:JA:325:VAL:O	2.13	0.48
1:JA:758:GLU:O	1:JA:762:VAL:HG22	2.14	0.48
1:KA:334:LEU:HD12	1:KA:357:TRP:NE1	2.29	0.48
1:M:497:VAL:HG13	1:M:497:VAL:O	2.13	0.48
1:RA:92:LEU:N	1:RA:92:LEU:HD12	2.29	0.48
1:TA:92:LEU:N	1:TA:92:LEU:HD12	2.29	0.48
1:TB:575:ILE:HG23	1:TB:603:VAL:HG22	1.94	0.48
1:VB:260:VAL:HG13	1:VB:260:VAL:O	2.13	0.48
1:XB:523:PHE:CD2	1:XB:568:VAL:HG23	2.49	0.48
1:XB:765:VAL:HG22	1:YB:759:LEU:HD21	1.95	0.48
1:AA:334:LEU:HD12	1:AA:357:TRP:NE1	2.29	0.48
1:AB:523:PHE:CE1	1:AB:545:TRP:CD1	3.02	0.48
1:BB:765:VAL:HG22	1:CB:759:LEU:HD21	1.96	0.48
1:E:788:ALA:O	1:E:792:VAL:HG23	2.14	0.48
1:FB:788:ALA:O	1:FB:792:VAL:HG23	2.14	0.48
1:HA:92:LEU:HD12	1:HA:92:LEU:N	2.29	0.48
1:HA:398:VAL:HG12	1:HA:491:PRO:HB3	1.96	0.48
1:HA:539:LEU:HD21	1:HA:599:ILE:HD11	1.95	0.48
1:IB:260:VAL:HG13	1:IB:260:VAL:O	2.13	0.48
1:IB:497:VAL:HG13	1:IB:497:VAL:O	2.12	0.48
1:JB:168:ILE:HG21	1:JB:174:LEU:HB2	1.95	0.48
1:KA:65:VAL:HG23	1:KA:67:ARG:HG3	1.95	0.48
1:LA:48:MET:HE3	1:LA:48:MET:HA	1.95	0.48
1:MA:65:VAL:HG23	1:MA:67:ARG:HG3	1.95	0.48
1:NA:497:VAL:O	1:NA:497:VAL:HG13	2.14	0.48
1:NA:529:ILE:CD1	1:NA:583:VAL:HG21	2.44	0.48
1:NB:52:THR:O	1:NB:109:ILE:HD11	2.14	0.48
1:QB:30:VAL:CG2	1:QB:50:MET:HE3	2.44	0.48
1:QB:529:ILE:HD12	1:QB:583:VAL:HG11	1.95	0.48

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:RB:65:VAL:HG23	1:RB:67:ARG:HG3	1.95	0.48
1:S:65:VAL:HG23	1:S:67:ARG:HG3	1.96	0.48
1:SA:56:ARG:NE	1:SA:56:ARG:N	2.62	0.48
1:SA:260:VAL:O	1:SA:260:VAL:HG13	2.13	0.48
1:UA:30:VAL:CG2	1:UA:50:MET:HE3	2.43	0.48
1:UB:65:VAL:HG23	1:UB:67:ARG:HG3	1.94	0.48
1:UB:529:ILE:HD12	1:UB:583:VAL:HG11	1.95	0.48
1:VB:52:THR:O	1:VB:109:ILE:HD11	2.14	0.48
1:WB:765:VAL:HG22	1:XB:759:LEU:HD21	1.96	0.48
1:YA:529:ILE:CD1	1:YA:583:VAL:HG21	2.44	0.48
1:ZB:575:ILE:HG23	1:ZB:603:VAL:HG22	1.95	0.48
1:E:144:LEU:HD12	1:E:144:LEU:O	2.14	0.48
1:EA:398:VAL:HG12	1:EA:491:PRO:HB3	1.96	0.48
1:EB:523:PHE:CD2	1:EB:568:VAL:HG23	2.49	0.48
1:F:260:VAL:O	1:F:260:VAL:HG13	2.13	0.48
1:F:330:GLN:C	1:F:407:MET:HE1	2.38	0.48
1:FA:575:ILE:HG23	1:FA:603:VAL:HG22	1.93	0.48
1:FB:92:LEU:N	1:FB:92:LEU:HD12	2.29	0.48
1:GB:334:LEU:HD12	1:GB:357:TRP:NE1	2.28	0.48
1:H:758:GLU:O	1:H:762:VAL:HG22	2.13	0.48
1:HA:65:VAL:HG23	1:HA:67:ARG:HG3	1.95	0.48
1:I:92:LEU:N	1:I:92:LEU:HD12	2.29	0.48
1:I:260:VAL:HG13	1:I:260:VAL:O	2.13	0.48
1:IB:65:VAL:HG23	1:IB:67:ARG:HG3	1.96	0.48
1:JA:260:VAL:HG13	1:JA:260:VAL:O	2.13	0.48
1:LB:334:LEU:HD12	1:LB:357:TRP:NE1	2.29	0.48
1:M:737:GLY:O	1:M:741:VAL:HG23	2.14	0.48
1:N:168:ILE:HG21	1:N:174:LEU:HB2	1.95	0.48
1:NB:334:LEU:HD12	1:NB:357:TRP:NE1	2.29	0.48
1:OA:737:GLY:O	1:OA:741:VAL:HG23	2.13	0.48
1:Q:17:HIS:HB3	1:Q:44:LEU:HD23	1.96	0.48
1:Q:523:PHE:CD2	1:Q:568:VAL:HG23	2.49	0.48
1:R:92:LEU:HD12	1:R:92:LEU:N	2.29	0.48
1:RB:497:VAL:HG13	1:RB:497:VAL:O	2.14	0.48
1:RB:538:GLN:HB2	1:RB:646:VAL:HG22	1.94	0.48
1:SB:65:VAL:HG23	1:SB:67:ARG:HG3	1.96	0.48
1:TB:260:VAL:HG13	1:TB:260:VAL:O	2.13	0.48
1:WA:144:LEU:HD23	1:WA:144:LEU:N	2.28	0.48
1:WB:92:LEU:HD12	1:WB:92:LEU:N	2.29	0.48
1:XA:65:VAL:HG23	1:XA:67:ARG:HG3	1.95	0.48
1:Z:778:GLU:HA	1:Z:781:VAL:HG22	1.94	0.48

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:AA:497:VAL:O	1:AA:497:VAL:HG13	2.12	0.48
1:BB:65:VAL:HG23	1:BB:67:ARG:HG3	1.95	0.48
1:BB:92:LEU:HD12	1:BB:92:LEU:N	2.29	0.48
1:D:398:VAL:HG12	1:D:491:PRO:HB3	1.96	0.48
1:EA:575:ILE:HG23	1:EA:603:VAL:HG22	1.95	0.48
1:F:92:LEU:N	1:F:92:LEU:HD12	2.28	0.48
1:FA:92:LEU:HD12	1:FA:92:LEU:N	2.29	0.48
1:GB:737:GLY:O	1:GB:741:VAL:HG23	2.13	0.48
1:HA:778:GLU:HA	1:HA:781:VAL:HG22	1.95	0.48
1:I:778:GLU:HA	1:I:781:VAL:HG22	1.95	0.48
1:L:516:LEU:O	1:L:516:LEU:HD23	2.13	0.48
1:N:144:LEU:HD22	1:N:163:ILE:HD11	1.96	0.48
1:N:334:LEU:HD12	1:N:357:TRP:NE1	2.29	0.48
1:OB:65:VAL:HG23	1:OB:67:ARG:HG3	1.96	0.48
1:Q:260:VAL:O	1:Q:260:VAL:HG13	2.13	0.48
1:UB:575:ILE:HG23	1:UB:603:VAL:HG22	1.95	0.48
1:W:30:VAL:CG2	1:W:50:MET:HE3	2.42	0.48
1:WA:523:PHE:CD2	1:WA:568:VAL:HG23	2.49	0.48
1:WB:788:ALA:O	1:WB:792:VAL:HG23	2.14	0.48
1:Y:497:VAL:O	1:Y:497:VAL:HG13	2.13	0.48
1:Y:529:ILE:HD12	1:Y:583:VAL:HG11	1.95	0.48
1:Y:737:GLY:O	1:Y:741:VAL:HG23	2.13	0.48
1:YA:390:VAL:HG23	1:YA:390:VAL:O	2.12	0.48
1:YB:390:VAL:O	1:YB:390:VAL:HG23	2.14	0.48
1:AB:92:LEU:HD12	1:AB:92:LEU:N	2.29	0.47
1:AC:804:PRO:O	1:AC:807:ILE:HG22	2.14	0.47
1:B:56:ARG:HE	1:C:127:LEU:HD21	1.79	0.47
1:B:65:VAL:HG23	1:B:67:ARG:HG3	1.96	0.47
1:BB:168:ILE:HG21	1:BB:174:LEU:HB2	1.96	0.47
1:DB:366:VAL:O	1:DB:366:VAL:HG13	2.14	0.47
1:EA:260:VAL:HG13	1:EA:260:VAL:O	2.13	0.47
1:GA:737:GLY:O	1:GA:741:VAL:HG23	2.14	0.47
1:GB:65:VAL:HG23	1:GB:67:ARG:HG3	1.95	0.47
1:GB:92:LEU:HD12	1:GB:92:LEU:N	2.29	0.47
1:H:168:ILE:HG21	1:H:174:LEU:HB2	1.96	0.47
1:HB:144:LEU:HD12	1:HB:144:LEU:O	2.14	0.47
1:JA:92:LEU:HD12	1:JA:92:LEU:N	2.29	0.47
1:JB:765:VAL:HG22	1:KB:759:LEU:HD21	1.96	0.47
1:M:334:LEU:HD12	1:M:357:TRP:NE1	2.29	0.47
1:MA:758:GLU:O	1:MA:762:VAL:HG22	2.13	0.47
1:NA:65:VAL:HG23	1:NA:67:ARG:HG3	1.95	0.47

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:QA:334:LEU:HD12	1:QA:357:TRP:NE1	2.29	0.47
1:QA:529:ILE:CD1	1:QA:583:VAL:HG11	2.43	0.47
1:QB:144:LEU:O	1:QB:144:LEU:HD12	2.14	0.47
1:TB:65:VAL:HG23	1:TB:67:ARG:HG3	1.96	0.47
1:UB:539:LEU:HD21	1:UB:599:ILE:HD11	1.95	0.47
1:VA:52:THR:O	1:VA:109:ILE:HD11	2.14	0.47
1:W:334:LEU:HD12	1:W:357:TRP:NE1	2.29	0.47
1:Y:56:ARG:HE	1:Z:127:LEU:HD21	1.77	0.47
1:Y:538:GLN:HB2	1:Y:646:VAL:HG22	1.95	0.47
1:YA:52:THR:O	1:YA:109:ILE:HD11	2.14	0.47
1:ZA:516:LEU:O	1:ZA:516:LEU:HD23	2.14	0.47
1:AA:65:VAL:HG23	1:AA:67:ARG:HG3	1.95	0.47
1:AA:778:GLU:HA	1:AA:781:VAL:HG22	1.95	0.47
1:BA:17:HIS:HB3	1:BA:44:LEU:HD23	1.96	0.47
1:BB:144:LEU:HD22	1:BB:163:ILE:HD11	1.96	0.47
1:CA:778:GLU:HA	1:CA:781:VAL:HG22	1.94	0.47
1:DB:737:GLY:O	1:DB:741:VAL:HG23	2.14	0.47
1:E:398:VAL:HG12	1:E:491:PRO:HB3	1.96	0.47
1:EB:398:VAL:HG12	1:EB:491:PRO:HB3	1.96	0.47
1:F:17:HIS:HB3	1:F:44:LEU:HD23	1.96	0.47
1:FB:65:VAL:HG23	1:FB:67:ARG:HG3	1.96	0.47
1:GA:398:VAL:HG12	1:GA:491:PRO:HB3	1.96	0.47
1:HA:529:ILE:HD12	1:HA:583:VAL:HG11	1.95	0.47
1:I:144:LEU:HD23	1:I:144:LEU:N	2.28	0.47
1:IA:92:LEU:N	1:IA:92:LEU:HD12	2.29	0.47
1:J:334:LEU:HD12	1:J:357:TRP:NE1	2.29	0.47
1:JB:778:GLU:HA	1:JB:781:VAL:HG22	1.95	0.47
1:LA:65:VAL:HG23	1:LA:67:ARG:HG3	1.96	0.47
1:LB:144:LEU:N	1:LB:144:LEU:HD23	2.29	0.47
1:LB:788:ALA:O	1:LB:792:VAL:HG23	2.14	0.47
1:N:765:VAL:HG22	1:O:759:LEU:HD21	1.95	0.47
1:RB:575:ILE:HG23	1:RB:603:VAL:HG22	1.94	0.47
1:SA:788:ALA:O	1:SA:792:VAL:HG23	2.14	0.47
1:TA:52:THR:O	1:TA:109:ILE:HD11	2.14	0.47
1:UB:52:THR:O	1:UB:109:ILE:HD11	2.14	0.47
1:VA:17:HIS:HB3	1:VA:44:LEU:HD23	1.97	0.47
1:VB:92:LEU:N	1:VB:92:LEU:HD12	2.29	0.47
1:WA:778:GLU:HA	1:WA:781:VAL:HG22	1.95	0.47
1:XA:17:HIS:HB3	1:XA:44:LEU:HD23	1.96	0.47
1:XA:168:ILE:HG21	1:XA:174:LEU:HB2	1.96	0.47
1:A:778:GLU:HA	1:A:781:VAL:HG22	1.96	0.47

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:AC:260:VAL:O	1:AC:260:VAL:HG13	2.13	0.47
1:AC:788:ALA:O	1:AC:792:VAL:HG23	2.14	0.47
1:D:65:VAL:HG23	1:D:67:ARG:HG3	1.96	0.47
1:DB:334:LEU:HD12	1:DB:357:TRP:NE1	2.29	0.47
1:E:52:THR:O	1:E:109:ILE:HD11	2.14	0.47
1:E:56:ARG:HE	1:F:127:LEU:HD21	1.80	0.47
1:FA:168:ILE:HG21	1:FA:174:LEU:HB2	1.96	0.47
1:HA:390:VAL:HG23	1:HA:390:VAL:O	2.13	0.47
1:IA:543:TYR:CE2	1:IA:575:ILE:HG21	2.50	0.47
1:JB:539:LEU:HD11	1:JB:640:VAL:HG13	1.96	0.47
1:MA:497:VAL:HG13	1:MA:497:VAL:O	2.14	0.47
1:MB:758:GLU:O	1:MB:762:VAL:HG22	2.15	0.47
1:N:52:THR:O	1:N:109:ILE:HD11	2.14	0.47
1:PB:788:ALA:O	1:PB:792:VAL:HG23	2.15	0.47
1:RB:260:VAL:O	1:RB:260:VAL:HG13	2.13	0.47
1:T:144:LEU:O	1:T:144:LEU:HD12	2.14	0.47
1:TA:788:ALA:O	1:TA:792:VAL:HG23	2.14	0.47
1:UA:737:GLY:O	1:UA:741:VAL:HG23	2.15	0.47
1:UB:778:GLU:HA	1:UB:781:VAL:HG22	1.95	0.47
1:VA:168:ILE:HG21	1:VA:174:LEU:HB2	1.96	0.47
1:VA:575:ILE:HG23	1:VA:603:VAL:HG22	1.96	0.47
1:W:65:VAL:HG23	1:W:67:ARG:HG3	1.96	0.47
1:AB:398:VAL:HG12	1:AB:491:PRO:HB3	1.96	0.47
1:AB:737:GLY:O	1:AB:741:VAL:HG23	2.14	0.47
1:CA:330:GLN:C	1:CA:407:MET:HE1	2.40	0.47
1:EA:92:LEU:HD12	1:EA:92:LEU:N	2.29	0.47
1:EA:497:VAL:HG13	1:EA:497:VAL:O	2.14	0.47
1:EB:529:ILE:HD12	1:EB:583:VAL:HG11	1.96	0.47
1:G:334:LEU:HD12	1:G:357:TRP:NE1	2.30	0.47
1:GA:529:ILE:CD1	1:GA:583:VAL:HG11	2.44	0.47
1:H:334:LEU:HD12	1:H:357:TRP:NE1	2.29	0.47
1:K:497:VAL:O	1:K:497:VAL:HG13	2.14	0.47
1:K:529:ILE:CD1	1:K:583:VAL:HG21	2.44	0.47
1:MA:398:VAL:HG12	1:MA:491:PRO:HB3	1.96	0.47
1:MA:575:ILE:HG23	1:MA:603:VAL:HG22	1.95	0.47
1:OB:334:LEU:HD12	1:OB:357:TRP:NE1	2.29	0.47
1:P:737:GLY:O	1:P:741:VAL:HG23	2.14	0.47
1:PB:260:VAL:O	1:PB:260:VAL:HG13	2.13	0.47
1:Q:497:VAL:O	1:Q:497:VAL:HG13	2.14	0.47
1:QA:398:VAL:HG12	1:QA:491:PRO:HB3	1.94	0.47
1:RA:65:VAL:HG23	1:RA:67:ARG:HG3	1.95	0.47

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:SA:92:LEU:HD12	1:SA:92:LEU:N	2.29	0.47
1:UB:497:VAL:HG13	1:UB:497:VAL:O	2.13	0.47
1:V:52:THR:O	1:V:109:ILE:HD11	2.15	0.47
1:V:144:LEU:N	1:V:144:LEU:HD23	2.28	0.47
1:WB:737:GLY:O	1:WB:741:VAL:HG23	2.14	0.47
1:XA:575:ILE:HG23	1:XA:603:VAL:HG22	1.96	0.47
1:XA:804:PRO:O	1:XA:807:ILE:HG22	2.15	0.47
1:Y:92:LEU:N	1:Y:92:LEU:HD12	2.29	0.47
1:AA:779:LEU:HD21	1:BA:774:ARG:HG2	1.97	0.47
1:AC:758:GLU:O	1:AC:762:VAL:HG22	2.15	0.47
1:DA:737:GLY:O	1:DA:741:VAL:HG23	2.13	0.47
1:F:52:THR:O	1:F:109:ILE:HD11	2.15	0.47
1:F:65:VAL:HG23	1:F:67:ARG:HG3	1.97	0.47
1:GA:65:VAL:HG23	1:GA:67:ARG:HG3	1.96	0.47
1:GA:345:ASP:OD2	1:GA:349:VAL:HG13	2.14	0.47
1:HB:65:VAL:HG23	1:HB:67:ARG:HG3	1.96	0.47
1:JB:65:VAL:HG23	1:JB:67:ARG:HG3	1.96	0.47
1:KA:758:GLU:O	1:KA:762:VAL:HG22	2.15	0.47
1:NA:788:ALA:O	1:NA:792:VAL:HG23	2.15	0.47
1:PB:330:GLN:C	1:PB:407:MET:HE1	2.39	0.47
1:Q:65:VAL:HG23	1:Q:67:ARG:HG3	1.96	0.47
1:QB:65:VAL:HG23	1:QB:67:ARG:HG3	1.96	0.47
1:QB:92:LEU:N	1:QB:92:LEU:HD12	2.29	0.47
1:S:575:ILE:HG23	1:S:603:VAL:HG22	1.95	0.47
1:T:168:ILE:HG21	1:T:174:LEU:HB2	1.96	0.47
1:TA:778:GLU:HA	1:TA:781:VAL:HG22	1.95	0.47
1:TB:17:HIS:CB	1:TB:44:LEU:HD23	2.44	0.47
1:TB:737:GLY:O	1:TB:741:VAL:HG23	2.14	0.47
1:V:778:GLU:HA	1:V:781:VAL:HG22	1.95	0.47
1:VA:44:LEU:H	1:VA:44:LEU:HD22	1.79	0.47
1:W:778:GLU:HA	1:W:781:VAL:HG22	1.96	0.47
1:X:778:GLU:HA	1:X:781:VAL:HG22	1.94	0.47
1:YA:366:VAL:HG13	1:YA:366:VAL:O	2.15	0.47
1:ZB:398:VAL:HG12	1:ZB:491:PRO:HB3	1.95	0.47
1:A:56:ARG:HE	1:B:127:LEU:HD21	1.80	0.47
1:A:127:LEU:HD21	1:NA:56:ARG:HE	1.80	0.47
1:AC:17:HIS:HB3	1:AC:44:LEU:HD23	1.96	0.47
1:B:92:LEU:HD12	1:B:92:LEU:N	2.30	0.47
1:BA:260:VAL:HG13	1:BA:260:VAL:O	2.14	0.47
1:CB:788:ALA:O	1:CB:792:VAL:HG23	2.14	0.47
1:DB:788:ALA:O	1:DB:792:VAL:HG23	2.15	0.47

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:E:92:LEU:HD12	1:E:92:LEU:N	2.29	0.47
1:E:543:TYR:CE2	1:E:575:ILE:HG21	2.50	0.47
1:FA:758:GLU:O	1:FA:762:VAL:HG22	2.13	0.47
1:HB:92:LEU:N	1:HB:92:LEU:HD12	2.30	0.47
1:JA:168:ILE:HG22	1:JA:215:LEU:CD2	2.42	0.47
1:KB:788:ALA:O	1:KB:792:VAL:HG23	2.15	0.47
1:L:497:VAL:O	1:L:497:VAL:HG13	2.13	0.47
1:LA:92:LEU:N	1:LA:92:LEU:HD12	2.29	0.47
1:N:56:ARG:HE	1:O:127:LEU:HD21	1.78	0.47
1:N:65:VAL:HG23	1:N:67:ARG:HG3	1.96	0.47
1:P:788:ALA:O	1:P:792:VAL:HG23	2.15	0.47
1:PA:65:VAL:HG23	1:PA:67:ARG:HG3	1.96	0.47
1:PA:497:VAL:HG13	1:PA:497:VAL:O	2.14	0.47
1:Q:529:ILE:HD12	1:Q:583:VAL:HG11	1.95	0.47
1:RA:334:LEU:HD12	1:RA:357:TRP:NE1	2.30	0.47
1:RA:788:ALA:O	1:RA:792:VAL:HG23	2.14	0.47
1:RB:56:ARG:HE	1:SB:127:LEU:HD21	1.80	0.47
1:S:52:THR:O	1:S:109:ILE:HD11	2.14	0.47
1:SA:219:VAL:HG11	1:SA:227:LEU:HD22	1.97	0.47
1:VA:92:LEU:HD12	1:VA:92:LEU:N	2.29	0.47
1:XA:168:ILE:HG22	1:XA:215:LEU:CD2	2.44	0.47
1:Y:65:VAL:HG23	1:Y:67:ARG:HG3	1.95	0.47
1:Y:144:LEU:HD12	1:Y:163:ILE:HD11	1.96	0.47
1:ZA:398:VAL:HG12	1:ZA:491:PRO:HB3	1.97	0.47
1:AB:281:TYR:CG	1:AB:366:VAL:HG23	2.50	0.47
1:BA:65:VAL:HG23	1:BA:67:ARG:HG3	1.96	0.47
1:CA:52:THR:O	1:CA:109:ILE:HD11	2.14	0.47
1:CA:788:ALA:O	1:CA:792:VAL:HG23	2.15	0.47
1:CB:529:ILE:HD12	1:CB:583:VAL:HG11	1.95	0.47
1:EA:65:VAL:HG23	1:EA:67:ARG:HG3	1.96	0.47
1:EB:30:VAL:CG2	1:EB:50:MET:HE3	2.44	0.47
1:F:30:VAL:CG2	1:F:50:MET:HE2	2.44	0.47
1:FA:539:LEU:HD12	1:FA:540:GLN:N	2.30	0.47
1:FB:398:VAL:HG12	1:FB:491:PRO:HB3	1.97	0.47
1:G:56:ARG:HE	1:H:127:LEU:HD21	1.79	0.47
1:GB:144:LEU:N	1:GB:144:LEU:HD23	2.30	0.47
1:GB:281:TYR:CG	1:GB:366:VAL:HG23	2.50	0.47
1:HA:345:ASP:OD2	1:HA:349:VAL:HG13	2.14	0.47
1:HA:497:VAL:HG13	1:HA:497:VAL:O	2.13	0.47
1:HA:543:TYR:CE2	1:HA:575:ILE:HG21	2.50	0.47
1:HB:497:VAL:O	1:HB:497:VAL:HG13	2.14	0.47

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:IA:52:THR:O	1:IA:109:ILE:HD11	2.14	0.47
1:IA:65:VAL:HG23	1:IA:67:ARG:HG3	1.96	0.47
1:IA:168:ILE:HG21	1:IA:174:LEU:HB2	1.96	0.47
1:IA:334:LEU:HD12	1:IA:357:TRP:NE1	2.30	0.47
1:IB:575:ILE:HG23	1:IB:603:VAL:HG22	1.97	0.47
1:J:788:ALA:O	1:J:792:VAL:HG23	2.15	0.47
1:K:398:VAL:HG12	1:K:491:PRO:HB3	1.96	0.47
1:K:788:ALA:O	1:K:792:VAL:HG23	2.14	0.47
1:KA:92:LEU:HD12	1:KA:92:LEU:N	2.29	0.47
1:KA:788:ALA:O	1:KA:792:VAL:HG23	2.14	0.47
1:LA:758:GLU:O	1:LA:762:VAL:HG22	2.15	0.47
1:MA:487:VAL:HG13	1:MA:487:VAL:O	2.15	0.47
1:MB:92:LEU:HD12	1:MB:92:LEU:N	2.30	0.47
1:N:92:LEU:N	1:N:92:LEU:HD12	2.29	0.47
1:NA:543:TYR:CE2	1:NA:575:ILE:HG21	2.50	0.47
1:NB:788:ALA:O	1:NB:792:VAL:HG23	2.15	0.47
1:O:52:THR:O	1:O:109:ILE:HD11	2.15	0.47
1:O:788:ALA:O	1:O:792:VAL:HG23	2.14	0.47
1:OB:497:VAL:HG13	1:OB:497:VAL:O	2.13	0.47
1:PB:168:ILE:HG21	1:PB:174:LEU:HB2	1.96	0.47
1:RA:168:ILE:HG21	1:RA:174:LEU:HB2	1.96	0.47
1:RB:92:LEU:HD12	1:RB:92:LEU:N	2.29	0.47
1:S:398:VAL:HG12	1:S:491:PRO:HB3	1.95	0.47
1:SB:92:LEU:N	1:SB:92:LEU:HD12	2.29	0.47
1:T:529:ILE:HD12	1:T:583:VAL:HG11	1.96	0.47
1:TA:63:ASN:OD1	1:TA:104:VAL:HG12	2.15	0.47
1:TA:504:ARG:HA	1:TA:504:ARG:NE	2.30	0.47
1:TB:168:ILE:HG21	1:TB:174:LEU:HB2	1.95	0.47
1:UA:758:GLU:O	1:UA:762:VAL:HG22	2.15	0.47
1:UB:788:ALA:O	1:UB:792:VAL:HG23	2.15	0.47
1:V:65:VAL:HG23	1:V:67:ARG:HG3	1.96	0.47
1:VA:65:VAL:HG23	1:VA:67:ARG:HG3	1.95	0.47
1:VA:334:LEU:HD12	1:VA:357:TRP:NE1	2.30	0.47
1:VA:737:GLY:O	1:VA:741:VAL:HG23	2.14	0.47
1:VB:168:ILE:HG21	1:VB:174:LEU:HB2	1.96	0.47
1:WA:65:VAL:HG23	1:WA:67:ARG:HG3	1.95	0.47
1:WB:168:ILE:HG22	1:WB:215:LEU:CD2	2.42	0.47
1:X:17:HIS:HB3	1:X:44:LEU:HD23	1.97	0.47
1:X:788:ALA:O	1:X:792:VAL:HG23	2.15	0.47
1:XA:334:LEU:HD12	1:XA:357:TRP:NE1	2.30	0.47
1:XB:92:LEU:HD12	1:XB:92:LEU:N	2.29	0.47

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:ZA:65:VAL:HG23	1:ZA:67:ARG:HG3	1.95	0.47
1:ZA:543:TYR:CE2	1:ZA:575:ILE:HG21	2.50	0.47
1:ZB:497:VAL:HG13	1:ZB:497:VAL:O	2.14	0.47
1:A:168:ILE:HG21	1:A:174:LEU:HB2	1.96	0.47
1:A:737:GLY:O	1:A:741:VAL:HG23	2.14	0.47
1:AA:765:VAL:HG22	1:BA:759:LEU:HD21	1.96	0.47
1:AB:366:VAL:O	1:AB:366:VAL:HG13	2.14	0.47
1:DA:168:ILE:HG22	1:DA:215:LEU:CD2	2.45	0.47
1:EA:334:LEU:HD12	1:EA:357:TRP:NE1	2.30	0.47
1:EB:575:ILE:HG23	1:EB:603:VAL:HG22	1.95	0.47
1:G:737:GLY:O	1:G:741:VAL:HG23	2.15	0.47
1:G:758:GLU:O	1:G:762:VAL:HG22	2.15	0.47
1:G:788:ALA:O	1:G:792:VAL:HG23	2.15	0.47
1:HB:281:TYR:CG	1:HB:366:VAL:HG23	2.50	0.47
1:J:575:ILE:HG23	1:J:603:VAL:HG22	1.96	0.47
1:LA:56:ARG:HE	1:MA:127:LEU:HD21	1.80	0.47
1:MB:65:VAL:HG23	1:MB:67:ARG:HG3	1.96	0.47
1:N:504:ARG:HA	1:N:504:ARG:NE	2.30	0.47
1:OA:788:ALA:O	1:OA:792:VAL:HG23	2.15	0.47
1:OB:219:VAL:HG11	1:OB:227:LEU:HD22	1.97	0.47
1:P:366:VAL:HG13	1:P:366:VAL:O	2.14	0.47
1:PB:52:THR:O	1:PB:109:ILE:HD11	2.15	0.47
1:RB:17:HIS:HB3	1:RB:44:LEU:HD23	1.97	0.47
1:T:758:GLU:O	1:T:762:VAL:HG22	2.15	0.47
1:T:788:ALA:O	1:T:792:VAL:HG23	2.15	0.47
1:V:168:ILE:HG21	1:V:174:LEU:HB2	1.97	0.47
1:XB:788:ALA:O	1:XB:792:VAL:HG23	2.14	0.47
1:A:92:LEU:N	1:A:92:LEU:HD12	2.30	0.47
1:B:497:VAL:HG13	1:B:497:VAL:O	2.14	0.47
1:BB:17:HIS:CB	1:BB:44:LEU:HD23	2.45	0.47
1:C:65:VAL:HG23	1:C:67:ARG:HG3	1.96	0.47
1:CA:504:ARG:NE	1:CA:504:ARG:HA	2.30	0.47
1:D:63:ASN:OD1	1:D:104:VAL:HG12	2.14	0.47
1:DA:92:LEU:N	1:DA:92:LEU:HD12	2.30	0.47
1:E:219:VAL:HG11	1:E:227:LEU:HD22	1.97	0.47
1:H:575:ILE:HG23	1:H:603:VAL:HG22	1.96	0.47
1:I:168:ILE:HG21	1:I:174:LEU:HB2	1.97	0.47
1:IA:788:ALA:O	1:IA:792:VAL:HG23	2.15	0.47
1:IB:788:ALA:O	1:IB:792:VAL:HG23	2.15	0.47
1:JA:737:GLY:O	1:JA:741:VAL:HG23	2.15	0.47
1:L:788:ALA:O	1:L:792:VAL:HG23	2.15	0.47

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:NB:65:VAL:HG23	1:NB:67:ARG:HG3	1.96	0.47
1:OA:92:LEU:N	1:OA:92:LEU:HD12	2.30	0.47
1:P:56:ARG:HE	1:Q:127:LEU:HD21	1.79	0.47
1:PB:334:LEU:HD12	1:PB:357:TRP:NE1	2.30	0.47
1:T:65:VAL:HG23	1:T:67:ARG:HG3	1.97	0.47
1:T:92:LEU:N	1:T:92:LEU:HD12	2.30	0.47
1:TA:65:VAL:HG23	1:TA:67:ARG:HG3	1.96	0.47
1:UB:543:TYR:CE2	1:UB:575:ILE:HG21	2.50	0.47
1:VA:778:GLU:HA	1:VA:781:VAL:HG22	1.95	0.47
1:VB:788:ALA:O	1:VB:792:VAL:HG23	2.15	0.47
1:WA:56:ARG:HE	1:XA:127:LEU:HD21	1.80	0.47
1:XA:56:ARG:HE	1:YA:127:LEU:HD21	1.78	0.47
1:Y:504:ARG:HA	1:Y:504:ARG:NE	2.30	0.47
1:AA:788:ALA:O	1:AA:792:VAL:HG23	2.15	0.47
1:AB:788:ALA:O	1:AB:792:VAL:HG23	2.15	0.47
1:B:504:ARG:NE	1:B:504:ARG:HA	2.30	0.47
1:BA:516:LEU:HD23	1:BA:516:LEU:O	2.15	0.47
1:C:788:ALA:O	1:C:792:VAL:HG23	2.15	0.47
1:DA:529:ILE:HD12	1:DA:583:VAL:HG11	1.96	0.47
1:EB:56:ARG:HE	1:FB:127:LEU:HD21	1.80	0.47
1:EB:65:VAL:HG23	1:EB:67:ARG:HG3	1.97	0.47
1:F:504:ARG:HA	1:F:504:ARG:NE	2.30	0.47
1:FB:281:TYR:CG	1:FB:366:VAL:HG23	2.50	0.47
1:J:52:THR:O	1:J:109:ILE:HD11	2.15	0.47
1:J:398:VAL:HG12	1:J:491:PRO:HB3	1.96	0.47
1:K:334:LEU:HD12	1:K:357:TRP:NE1	2.29	0.47
1:KB:92:LEU:N	1:KB:92:LEU:HD12	2.29	0.47
1:L:529:ILE:HD12	1:L:583:VAL:HG11	1.97	0.47
1:LB:92:LEU:HD12	1:LB:92:LEU:N	2.30	0.47
1:M:523:PHE:CE1	1:M:545:TRP:CD1	3.02	0.47
1:MB:56:ARG:HE	1:NB:127:LEU:HD21	1.80	0.47
1:MB:168:ILE:HG21	1:MB:174:LEU:HB2	1.97	0.47
1:MB:543:TYR:CE2	1:MB:575:ILE:HG21	2.50	0.47
1:NB:765:VAL:HG22	1:OB:759:LEU:HD21	1.96	0.47
1:OA:30:VAL:HG21	1:OA:50:MET:HE3	1.97	0.47
1:OA:65:VAL:HG23	1:OA:67:ARG:HG3	1.96	0.47
1:PA:92:LEU:HD12	1:PA:92:LEU:N	2.30	0.47
1:S:497:VAL:O	1:S:497:VAL:HG13	2.14	0.47
1:S:504:ARG:HA	1:S:504:ARG:NE	2.30	0.47
1:SA:65:VAL:HG23	1:SA:67:ARG:HG3	1.95	0.47
1:SA:504:ARG:NE	1:SA:504:ARG:HA	2.30	0.47

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:TA:168:ILE:HG21	1:TA:174:LEU:HB2	1.97	0.47
1:UA:65:VAL:HG23	1:UA:67:ARG:HG3	1.96	0.47
1:UA:334:LEU:HD12	1:UA:357:TRP:NE1	2.30	0.47
1:VA:398:VAL:HG12	1:VA:491:PRO:HB3	1.96	0.47
1:Y:144:LEU:HD23	1:Y:144:LEU:N	2.29	0.47
1:YB:504:ARG:NE	1:YB:504:ARG:HA	2.30	0.47
1:Z:17:HIS:HB3	1:Z:44:LEU:HD23	1.97	0.47
1:A:334:LEU:HD12	1:A:357:TRP:NE1	2.30	0.46
1:B:52:THR:O	1:B:109:ILE:HD11	2.15	0.46
1:C:219:VAL:HG11	1:C:227:LEU:HD22	1.97	0.46
1:DB:281:TYR:CG	1:DB:366:VAL:HG23	2.50	0.46
1:FA:53:VAL:HG12	1:FA:93:ALA:HA	1.97	0.46
1:GB:543:TYR:CE2	1:GB:575:ILE:HG21	2.50	0.46
1:IB:543:TYR:CE2	1:IB:575:ILE:HG21	2.50	0.46
1:KA:17:HIS:HB3	1:KA:44:LEU:HD23	1.97	0.46
1:KA:765:VAL:HG22	1:LA:759:LEU:HD21	1.96	0.46
1:LB:504:ARG:NE	1:LB:504:ARG:HA	2.30	0.46
1:M:56:ARG:HE	1:N:127:LEU:HD21	1.80	0.46
1:MA:92:LEU:HD12	1:MA:92:LEU:N	2.30	0.46
1:MB:17:HIS:HB3	1:MB:44:LEU:HD23	1.97	0.46
1:OA:778:GLU:HA	1:OA:781:VAL:HG22	1.96	0.46
1:OB:737:GLY:O	1:OB:741:VAL:HG23	2.15	0.46
1:PB:92:LEU:N	1:PB:92:LEU:HD12	2.30	0.46
1:QA:65:VAL:HG23	1:QA:67:ARG:HG3	1.97	0.46
1:QA:504:ARG:HA	1:QA:504:ARG:NE	2.31	0.46
1:QB:56:ARG:HE	1:RB:127:LEU:HD21	1.80	0.46
1:R:65:VAL:HG23	1:R:67:ARG:HG3	1.96	0.46
1:R:398:VAL:HG12	1:R:491:PRO:HB3	1.97	0.46
1:T:398:VAL:HG12	1:T:491:PRO:HB3	1.96	0.46
1:UA:504:ARG:NE	1:UA:504:ARG:HA	2.30	0.46
1:V:543:TYR:CE2	1:V:575:ILE:HG21	2.50	0.46
1:V:788:ALA:O	1:V:792:VAL:HG23	2.15	0.46
1:VB:543:TYR:CE2	1:VB:575:ILE:HG21	2.49	0.46
1:W:765:VAL:HG22	1:X:759:LEU:HD21	1.96	0.46
1:X:504:ARG:NE	1:X:504:ARG:HA	2.31	0.46
1:Y:168:ILE:HG21	1:Y:174:LEU:HB2	1.97	0.46
1:Y:779:LEU:HD21	1:Z:774:ARG:HG2	1.98	0.46
1:YA:334:LEU:HD12	1:YA:357:TRP:NE1	2.29	0.46
1:Z:398:VAL:HG12	1:Z:491:PRO:HB3	1.97	0.46
1:AB:17:HIS:HB3	1:AB:44:LEU:HD23	1.98	0.46
1:AC:398:VAL:HG12	1:AC:491:PRO:HB3	1.96	0.46

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:BA:737:GLY:O	1:BA:741:VAL:HG23	2.15	0.46
1:D:334:LEU:HD12	1:D:357:TRP:NE1	2.31	0.46
1:DB:7:ILE:O	1:DB:8:ILE:C	2.58	0.46
1:F:302:VAL:HG11	1:F:306:LYS:HZ2	1.80	0.46
1:FA:65:VAL:HG23	1:FA:67:ARG:HG3	1.96	0.46
1:FA:539:LEU:HD11	1:FA:541:LEU:HD21	1.97	0.46
1:JA:65:VAL:HG23	1:JA:67:ARG:HG3	1.96	0.46
1:K:63:ASN:OD1	1:K:104:VAL:HG12	2.15	0.46
1:KB:398:VAL:HG12	1:KB:491:PRO:HB3	1.97	0.46
1:L:52:THR:O	1:L:109:ILE:HD11	2.15	0.46
1:NA:281:TYR:CG	1:NA:366:VAL:HG23	2.51	0.46
1:OB:398:VAL:HG12	1:OB:491:PRO:HB3	1.95	0.46
1:PA:504:ARG:NE	1:PA:504:ARG:HA	2.31	0.46
1:PB:504:ARG:HA	1:PB:504:ARG:NE	2.30	0.46
1:QA:92:LEU:N	1:QA:92:LEU:HD12	2.30	0.46
1:R:504:ARG:NE	1:R:504:ARG:HA	2.30	0.46
1:RA:504:ARG:NE	1:RA:504:ARG:HA	2.30	0.46
1:RB:334:LEU:HD12	1:RB:357:TRP:NE1	2.30	0.46
1:S:281:TYR:CG	1:S:366:VAL:HG23	2.50	0.46
1:SA:398:VAL:HG12	1:SA:491:PRO:HB3	1.96	0.46
1:T:497:VAL:O	1:T:497:VAL:HG13	2.14	0.46
1:T:504:ARG:NE	1:T:504:ARG:HA	2.30	0.46
1:VA:504:ARG:HA	1:VA:504:ARG:NE	2.30	0.46
1:W:52:THR:O	1:W:109:ILE:HD11	2.14	0.46
1:X:168:ILE:HG21	1:X:174:LEU:HB2	1.97	0.46
1:ZB:487:VAL:HG13	1:ZB:487:VAL:O	2.15	0.46
1:ZB:504:ARG:NE	1:ZB:504:ARG:HA	2.30	0.46
1:AC:92:LEU:HD12	1:AC:92:LEU:N	2.30	0.46
1:AC:497:VAL:HG13	1:AC:497:VAL:O	2.14	0.46
1:AC:529:ILE:CD1	1:AC:583:VAL:HG11	2.44	0.46
1:CA:92:LEU:HD12	1:CA:92:LEU:N	2.30	0.46
1:CA:168:ILE:HG22	1:CA:215:LEU:CD2	2.43	0.46
1:CB:56:ARG:HE	1:DB:127:LEU:HD21	1.80	0.46
1:DA:65:VAL:HG23	1:DA:67:ARG:HG3	1.96	0.46
1:EB:281:TYR:CG	1:EB:366:VAL:HG23	2.51	0.46
1:GB:504:ARG:NE	1:GB:504:ARG:HA	2.30	0.46
1:HB:758:GLU:O	1:HB:762:VAL:HG22	2.14	0.46
1:J:144:LEU:N	1:J:144:LEU:HD23	2.30	0.46
1:J:159:VAL:O	1:J:159:VAL:HG12	2.14	0.46
1:J:168:ILE:HG22	1:J:215:LEU:CD2	2.43	0.46
1:JA:398:VAL:HG12	1:JA:491:PRO:HB3	1.98	0.46

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:JB:52:THR:O	1:JB:109:ILE:HD11	2.15	0.46
1:KA:543:TYR:CE2	1:KA:575:ILE:HG21	2.50	0.46
1:N:366:VAL:HG13	1:N:366:VAL:O	2.16	0.46
1:NA:92:LEU:HD12	1:NA:92:LEU:N	2.30	0.46
1:NB:779:LEU:HD21	1:OB:774:ARG:HG2	1.97	0.46
1:OB:168:ILE:HG21	1:OB:174:LEU:HB2	1.97	0.46
1:RA:56:ARG:HE	1:SA:127:LEU:HD21	1.79	0.46
1:UA:788:ALA:O	1:UA:792:VAL:HG23	2.16	0.46
1:WA:52:THR:O	1:WA:109:ILE:HD11	2.16	0.46
1:X:386:GLU:O	1:X:402:ILE:HG23	2.16	0.46
1:YA:92:LEU:HD12	1:YA:92:LEU:N	2.30	0.46
1:YA:281:TYR:CG	1:YA:366:VAL:HG23	2.50	0.46
1:YA:398:VAL:HG12	1:YA:491:PRO:HB3	1.97	0.46
1:Z:758:GLU:O	1:Z:762:VAL:HG22	2.15	0.46
1:ZA:281:TYR:CG	1:ZA:366:VAL:HG23	2.51	0.46
1:A:788:ALA:O	1:A:792:VAL:HG23	2.15	0.46
1:AB:7:ILE:O	1:AB:8:ILE:C	2.59	0.46
1:BA:497:VAL:HG13	1:BA:497:VAL:O	2.14	0.46
1:BB:504:ARG:NE	1:BB:504:ARG:HA	2.30	0.46
1:C:92:LEU:N	1:C:92:LEU:HD12	2.31	0.46
1:CB:17:HIS:HB3	1:CB:44:LEU:HD23	1.98	0.46
1:D:504:ARG:NE	1:D:504:ARG:HA	2.30	0.46
1:EA:56:ARG:HE	1:FA:127:LEU:HD21	1.79	0.46
1:FA:398:VAL:HG12	1:FA:491:PRO:HB3	1.98	0.46
1:G:65:VAL:HG23	1:G:67:ARG:HG3	1.96	0.46
1:HA:168:ILE:HG21	1:HA:174:LEU:HB2	1.98	0.46
1:HB:366:VAL:HG13	1:HB:366:VAL:O	2.16	0.46
1:HB:543:TYR:CE2	1:HB:575:ILE:HG21	2.51	0.46
1:KA:52:THR:O	1:KA:109:ILE:HD11	2.15	0.46
1:KA:398:VAL:HG12	1:KA:491:PRO:HB3	1.98	0.46
1:L:543:TYR:CE2	1:L:575:ILE:HG21	2.50	0.46
1:O:504:ARG:NE	1:O:504:ARG:HA	2.31	0.46
1:PA:788:ALA:O	1:PA:792:VAL:HG23	2.16	0.46
1:QA:219:VAL:HG11	1:QA:227:LEU:HD22	1.97	0.46
1:S:17:HIS:HB3	1:S:44:LEU:HD23	1.98	0.46
1:TA:17:HIS:HB3	1:TA:44:LEU:HD23	1.97	0.46
1:UB:398:VAL:HG12	1:UB:491:PRO:HB3	1.96	0.46
1:UB:504:ARG:NE	1:UB:504:ARG:HA	2.31	0.46
1:VB:539:LEU:HD21	1:VB:599:ILE:HD11	1.97	0.46
1:W:504:ARG:NE	1:W:504:ARG:HA	2.30	0.46
1:WA:543:TYR:CE2	1:WA:575:ILE:HG21	2.50	0.46

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:XA:144:LEU:N	1:XA:144:LEU:HD23	2.30	0.46
1:Y:539:LEU:HD21	1:Y:541:LEU:HD21	1.97	0.46
1:ZA:92:LEU:N	1:ZA:92:LEU:HD12	2.31	0.46
1:AC:543:TYR:CE2	1:AC:575:ILE:HG21	2.50	0.46
1:BA:168:ILE:HG21	1:BA:174:LEU:HB2	1.97	0.46
1:CB:65:VAL:HG23	1:CB:67:ARG:HG3	1.95	0.46
1:CB:765:VAL:HG22	1:DB:759:LEU:HD21	1.97	0.46
1:D:56:ARG:HE	1:E:127:LEU:HD21	1.80	0.46
1:D:219:VAL:HG11	1:D:227:LEU:HD22	1.98	0.46
1:DA:398:VAL:HG12	1:DA:491:PRO:HB3	1.97	0.46
1:EB:334:LEU:HD12	1:EB:357:TRP:NE1	2.30	0.46
1:GA:92:LEU:N	1:GA:92:LEU:HD12	2.30	0.46
1:HA:487:VAL:HG13	1:HA:487:VAL:O	2.16	0.46
1:HB:345:ASP:OD2	1:HB:349:VAL:HG13	2.15	0.46
1:I:56:ARG:HE	1:J:127:LEU:HD21	1.80	0.46
1:IA:168:ILE:HG22	1:IA:215:LEU:CD2	2.43	0.46
1:JB:330:GLN:C	1:JB:407:MET:HE1	2.41	0.46
1:K:281:TYR:CG	1:K:366:VAL:HG23	2.50	0.46
1:MB:804:PRO:O	1:MB:807:ILE:HG22	2.15	0.46
1:NB:504:ARG:NE	1:NB:504:ARG:HA	2.30	0.46
1:NB:758:GLU:O	1:NB:762:VAL:HG22	2.16	0.46
1:O:497:VAL:HG13	1:O:497:VAL:O	2.15	0.46
1:O:765:VAL:HG22	1:P:759:LEU:HD21	1.96	0.46
1:OA:168:ILE:HG21	1:OA:174:LEU:HB2	1.97	0.46
1:OB:543:TYR:CE2	1:OB:575:ILE:HG21	2.51	0.46
1:PB:529:ILE:HD12	1:PB:583:VAL:HG11	1.98	0.46
1:PB:543:TYR:CE2	1:PB:575:ILE:HG21	2.51	0.46
1:Q:398:VAL:HG12	1:Q:491:PRO:HB3	1.97	0.46
1:QB:758:GLU:O	1:QB:762:VAL:HG22	2.15	0.46
1:RB:398:VAL:HG12	1:RB:491:PRO:HB3	1.96	0.46
1:UA:17:HIS:HB3	1:UA:44:LEU:HD23	1.98	0.46
1:V:17:HIS:HB3	1:V:44:LEU:HD23	1.98	0.46
1:V:538:GLN:HB2	1:V:646:VAL:HG22	1.97	0.46
1:W:56:ARG:HE	1:X:127:LEU:HD21	1.80	0.46
1:W:168:ILE:HG21	1:W:174:LEU:HB2	1.96	0.46
1:WA:53:VAL:HG12	1:WA:93:ALA:HA	1.97	0.46
1:XB:758:GLU:O	1:XB:762:VAL:HG22	2.15	0.46
1:YA:17:HIS:HB3	1:YA:44:LEU:HD23	1.98	0.46
1:ZA:504:ARG:NE	1:ZA:504:ARG:HA	2.30	0.46
1:A:30:VAL:HG21	1:A:50:MET:HE3	1.98	0.46
1:AA:219:VAL:HG11	1:AA:227:LEU:HD22	1.98	0.46

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:BA:504:ARG:NE	1:BA:504:ARG:HA	2.31	0.46
1:BB:788:ALA:O	1:BB:792:VAL:HG23	2.16	0.46
1:DA:17:HIS:HB3	1:DA:44:LEU:HD23	1.98	0.46
1:F:788:ALA:O	1:F:792:VAL:HG23	2.15	0.46
1:GA:219:VAL:HG11	1:GA:227:LEU:HD13	1.98	0.46
1:GA:788:ALA:O	1:GA:792:VAL:HG23	2.16	0.46
1:HB:52:THR:O	1:HB:109:ILE:HD11	2.16	0.46
1:JB:523:PHE:CE2	1:JB:545:TRP:CD1	3.04	0.46
1:KB:219:VAL:HG11	1:KB:227:LEU:HD22	1.98	0.46
1:L:56:ARG:HE	1:M:127:LEU:HD21	1.80	0.46
1:M:504:ARG:NE	1:M:504:ARG:HA	2.30	0.46
1:MA:504:ARG:NE	1:MA:504:ARG:HA	2.30	0.46
1:NA:504:ARG:HA	1:NA:504:ARG:NE	2.30	0.46
1:NB:575:ILE:HG23	1:NB:603:VAL:HG22	1.98	0.46
1:OA:17:HIS:HB3	1:OA:44:LEU:HD23	1.98	0.46
1:OA:281:TYR:CG	1:OA:366:VAL:HG23	2.50	0.46
1:OB:758:GLU:O	1:OB:762:VAL:HG22	2.16	0.46
1:PA:168:ILE:HG21	1:PA:174:LEU:HB2	1.97	0.46
1:RA:17:HIS:HB3	1:RA:44:LEU:HD23	1.98	0.46
1:RB:219:VAL:HG11	1:RB:227:LEU:HD22	1.98	0.46
1:S:543:TYR:CE2	1:S:575:ILE:HG21	2.50	0.46
1:SA:543:TYR:CE2	1:SA:575:ILE:HG21	2.50	0.46
1:TA:804:PRO:O	1:TA:807:ILE:HG22	2.16	0.46
1:W:330:GLN:C	1:W:407:MET:HE1	2.41	0.46
1:X:219:VAL:HG11	1:X:227:LEU:HD22	1.98	0.46
1:X:765:VAL:HG22	1:Y:759:LEU:HD21	1.97	0.46
1:XB:504:ARG:HA	1:XB:504:ARG:NE	2.30	0.46
1:YA:504:ARG:HA	1:YA:504:ARG:NE	2.31	0.46
1:YA:543:TYR:CE2	1:YA:575:ILE:HG21	2.51	0.46
1:YB:65:VAL:HG23	1:YB:67:ARG:HG3	1.97	0.46
1:Z:65:VAL:HG23	1:Z:67:ARG:HG3	1.96	0.46
1:Z:168:ILE:HG21	1:Z:174:LEU:HB2	1.97	0.46
1:ZB:529:ILE:HD12	1:ZB:583:VAL:HG11	1.97	0.46
1:AA:17:HIS:HB3	1:AA:44:LEU:HD23	1.98	0.46
1:AA:127:LEU:HD21	1:Z:56:ARG:HE	1.80	0.46
1:AC:504:ARG:NE	1:AC:504:ARG:HA	2.31	0.46
1:B:543:TYR:CE2	1:B:575:ILE:HG21	2.51	0.46
1:BA:281:TYR:CG	1:BA:366:VAL:HG23	2.51	0.46
1:CA:334:LEU:HD12	1:CA:357:TRP:NE1	2.30	0.46
1:DB:168:ILE:HG21	1:DB:174:LEU:HB2	1.98	0.46
1:EA:219:VAL:HG11	1:EA:227:LEU:HD22	1.98	0.46

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:FA:7:ILE:O	1:FA:8:ILE:C	2.59	0.46
1:IA:7:ILE:O	1:IA:8:ILE:C	2.59	0.46
1:JA:788:ALA:O	1:JA:792:VAL:HG23	2.16	0.46
1:K:52:THR:O	1:K:109:ILE:HD11	2.16	0.46
1:K:92:LEU:HD12	1:K:92:LEU:N	2.30	0.46
1:KA:281:TYR:CG	1:KA:366:VAL:HG23	2.51	0.46
1:KA:504:ARG:NE	1:KA:504:ARG:HA	2.31	0.46
1:KB:52:THR:O	1:KB:109:ILE:HD11	2.16	0.46
1:KB:504:ARG:HA	1:KB:504:ARG:NE	2.30	0.46
1:L:92:LEU:N	1:L:92:LEU:HD12	2.31	0.46
1:LB:543:TYR:CE2	1:LB:575:ILE:HG21	2.51	0.46
1:M:281:TYR:CG	1:M:366:VAL:HG23	2.50	0.46
1:MB:398:VAL:HG12	1:MB:491:PRO:HB3	1.97	0.46
1:OA:144:LEU:HD12	1:OA:163:ILE:HD11	1.97	0.46
1:OB:281:TYR:CG	1:OB:366:VAL:HG23	2.51	0.46
1:P:168:ILE:HG21	1:P:174:LEU:HB2	1.97	0.46
1:QA:788:ALA:O	1:QA:792:VAL:HG23	2.15	0.46
1:RB:804:PRO:O	1:RB:807:ILE:HG22	2.15	0.46
1:TB:538:GLN:HB2	1:TB:646:VAL:HG22	1.97	0.46
1:TB:788:ALA:O	1:TB:792:VAL:HG23	2.16	0.46
1:UA:7:ILE:O	1:UA:8:ILE:C	2.59	0.46
1:YA:539:LEU:HD11	1:YA:640:VAL:HG13	1.97	0.46
1:Z:7:ILE:O	1:Z:8:ILE:C	2.59	0.46
1:AA:504:ARG:HA	1:AA:504:ARG:NE	2.30	0.46
1:AA:758:GLU:O	1:AA:762:VAL:HG22	2.16	0.46
1:C:168:ILE:HG21	1:C:174:LEU:HB2	1.97	0.46
1:C:334:LEU:HD12	1:C:357:TRP:NE1	2.30	0.46
1:C:543:TYR:CE2	1:C:575:ILE:HG21	2.51	0.46
1:CA:168:ILE:HG21	1:CA:174:LEU:HB2	1.97	0.46
1:CA:543:TYR:CE2	1:CA:575:ILE:HG21	2.51	0.46
1:D:52:THR:O	1:D:109:ILE:HD11	2.15	0.46
1:E:504:ARG:NE	1:E:504:ARG:HA	2.30	0.46
1:E:523:PHE:CD2	1:E:568:VAL:HG23	2.50	0.46
1:EA:543:TYR:CE2	1:EA:575:ILE:HG21	2.51	0.46
1:G:92:LEU:HD12	1:G:92:LEU:N	2.30	0.46
1:G:504:ARG:NE	1:G:504:ARG:HA	2.30	0.46
1:GA:543:TYR:CE2	1:GA:575:ILE:HG21	2.51	0.46
1:HB:9:ARG:HH12	1:IB:19:LEU:HD11	1.81	0.46
1:IA:504:ARG:HA	1:IA:504:ARG:NE	2.30	0.46
1:IB:281:TYR:CG	1:IB:366:VAL:HG23	2.51	0.46
1:J:281:TYR:CG	1:J:366:VAL:HG23	2.51	0.46

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:J:504:ARG:NE	1:J:504:ARG:HA	2.30	0.46
1:JB:543:TYR:CE2	1:JB:575:ILE:HG21	2.51	0.46
1:KA:345:ASP:OD2	1:KA:349:VAL:HG13	2.15	0.46
1:P:281:TYR:CG	1:P:366:VAL:HG23	2.50	0.46
1:PA:57:HIS:O	1:PA:58:TYR:HD1	1.99	0.46
1:Q:30:VAL:CG2	1:Q:50:MET:HE3	2.44	0.46
1:RA:398:VAL:HG12	1:RA:491:PRO:HB3	1.96	0.46
1:S:63:ASN:OD1	1:S:104:VAL:HG12	2.16	0.46
1:S:168:ILE:HG21	1:S:174:LEU:HB2	1.97	0.46
1:T:366:VAL:O	1:T:366:VAL:HG13	2.16	0.46
1:TB:334:LEU:HD12	1:TB:357:TRP:NE1	2.31	0.46
1:TB:497:VAL:HG13	1:TB:497:VAL:O	2.16	0.46
1:TB:504:ARG:NE	1:TB:504:ARG:HA	2.31	0.46
1:UA:56:ARG:HE	1:VA:127:LEU:HD21	1.79	0.46
1:VA:44:LEU:HD22	1:VA:44:LEU:N	2.31	0.46
1:VA:281:TYR:CG	1:VA:366:VAL:HG23	2.51	0.46
1:VB:504:ARG:NE	1:VB:504:ARG:HA	2.30	0.46
1:WA:168:ILE:HG21	1:WA:174:LEU:HB2	1.98	0.46
1:XA:398:VAL:HG12	1:XA:491:PRO:HB3	1.96	0.46
1:ZA:788:ALA:O	1:ZA:792:VAL:HG23	2.15	0.46
1:A:497:VAL:O	1:A:497:VAL:HG13	2.15	0.46
1:AB:345:ASP:OD2	1:AB:349:VAL:HG13	2.16	0.46
1:B:788:ALA:O	1:B:792:VAL:HG23	2.16	0.46
1:BA:219:VAL:HG11	1:BA:227:LEU:HD22	1.98	0.46
1:BB:398:VAL:HG12	1:BB:491:PRO:HB3	1.98	0.46
1:CA:7:ILE:O	1:CA:8:ILE:C	2.59	0.46
1:CA:65:VAL:HG23	1:CA:67:ARG:HG3	1.96	0.46
1:D:758:GLU:O	1:D:762:VAL:HG22	2.16	0.46
1:DA:281:TYR:CG	1:DA:366:VAL:HG23	2.51	0.46
1:EB:366:VAL:HG13	1:EB:366:VAL:O	2.16	0.46
1:FA:281:TYR:CG	1:FA:366:VAL:HG23	2.51	0.46
1:JA:7:ILE:O	1:JA:8:ILE:C	2.59	0.46
1:JB:92:LEU:HD12	1:JB:92:LEU:N	2.30	0.46
1:L:30:VAL:CG2	1:L:50:MET:HE3	2.44	0.46
1:LA:504:ARG:HA	1:LA:504:ARG:NE	2.30	0.46
1:LA:575:ILE:HG23	1:LA:603:VAL:HG22	1.97	0.46
1:M:788:ALA:O	1:M:792:VAL:HG23	2.15	0.46
1:MA:168:ILE:HG21	1:MA:174:LEU:HB2	1.97	0.46
1:N:788:ALA:O	1:N:792:VAL:HG23	2.16	0.46
1:NA:487:VAL:HG13	1:NA:487:VAL:O	2.16	0.46
1:OB:366:VAL:O	1:OB:366:VAL:HG13	2.16	0.46

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:PA:17:HIS:HB3	1:PA:44:LEU:HD23	1.98	0.46
1:Q:56:ARG:HE	1:R:127:LEU:HD21	1.80	0.46
1:Q:334:LEU:HD12	1:Q:357:TRP:NE1	2.30	0.46
1:Q:366:VAL:O	1:Q:366:VAL:HG13	2.16	0.46
1:R:758:GLU:O	1:R:762:VAL:HG22	2.16	0.46
1:RB:543:TYR:CE2	1:RB:575:ILE:HG21	2.51	0.46
1:RB:788:ALA:O	1:RB:792:VAL:HG23	2.16	0.46
1:SB:168:ILE:HG21	1:SB:174:LEU:HB2	1.97	0.46
1:SB:788:ALA:O	1:SB:792:VAL:HG23	2.15	0.46
1:TB:219:VAL:HG11	1:TB:227:LEU:HD13	1.97	0.46
1:TB:543:TYR:CE2	1:TB:575:ILE:HG21	2.51	0.46
1:UA:398:VAL:HG12	1:UA:491:PRO:HB3	1.97	0.46
1:V:737:GLY:O	1:V:741:VAL:HG23	2.16	0.46
1:WA:281:TYR:CG	1:WA:366:VAL:HG23	2.51	0.46
1:XA:281:TYR:CG	1:XA:366:VAL:HG23	2.51	0.46
1:XB:334:LEU:HD12	1:XB:357:TRP:NE1	2.30	0.46
1:AA:543:TYR:CE2	1:AA:575:ILE:HG21	2.51	0.46
1:AC:56:ARG:HE	1:OA:127:LEU:HD21	1.80	0.46
1:AC:487:VAL:HG13	1:AC:487:VAL:O	2.16	0.46
1:BA:543:TYR:CE2	1:BA:575:ILE:HG21	2.51	0.46
1:BB:281:TYR:CG	1:BB:366:VAL:HG23	2.51	0.46
1:BB:570:ASP:C	1:BB:570:ASP:OD1	2.59	0.46
1:CB:53:VAL:HG12	1:CB:93:ALA:HA	1.98	0.46
1:D:168:ILE:HG21	1:D:174:LEU:HB2	1.98	0.46
1:FA:788:ALA:O	1:FA:792:VAL:HG23	2.16	0.46
1:G:497:VAL:HG13	1:G:497:VAL:O	2.15	0.46
1:GA:504:ARG:NE	1:GA:504:ARG:HA	2.30	0.46
1:H:44:LEU:HD22	1:H:44:LEU:N	2.31	0.46
1:HA:788:ALA:O	1:HA:792:VAL:HG23	2.15	0.46
1:I:523:PHE:CD2	1:I:568:VAL:HG23	2.51	0.46
1:I:543:TYR:CE2	1:I:575:ILE:HG21	2.50	0.46
1:IB:758:GLU:O	1:IB:762:VAL:HG22	2.15	0.46
1:JA:543:TYR:CE2	1:JA:575:ILE:HG21	2.51	0.46
1:JB:7:ILE:O	1:JB:8:ILE:C	2.59	0.46
1:K:17:HIS:HB3	1:K:44:LEU:HD23	1.97	0.46
1:K:529:ILE:CD1	1:K:583:VAL:HG11	2.45	0.46
1:K:539:LEU:HD11	1:K:640:VAL:HG13	1.98	0.46
1:KA:219:VAL:HG11	1:KA:227:LEU:HD22	1.98	0.46
1:LB:487:VAL:HG13	1:LB:487:VAL:O	2.16	0.46
1:M:366:VAL:HG13	1:M:366:VAL:O	2.15	0.46
1:NA:63:ASN:OD1	1:NA:104:VAL:HG12	2.15	0.46

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:NB:543:TYR:CE2	1:NB:575:ILE:HG21	2.51	0.46
1:OA:48:MET:HE2	1:OA:48:MET:HA	1.97	0.46
1:PA:219:VAL:HG11	1:PA:227:LEU:HD22	1.98	0.46
1:PB:398:VAL:HG12	1:PB:491:PRO:HB3	1.97	0.46
1:Q:504:ARG:NE	1:Q:504:ARG:HA	2.31	0.46
1:QA:7:ILE:O	1:QA:8:ILE:C	2.59	0.46
1:S:7:ILE:O	1:S:8:ILE:C	2.59	0.46
1:S:788:ALA:O	1:S:792:VAL:HG23	2.16	0.46
1:TA:7:ILE:O	1:TA:8:ILE:C	2.59	0.46
1:TB:92:LEU:HD12	1:TB:92:LEU:N	2.30	0.46
1:UA:393:VAL:HG22	1:UA:411:ASP:C	2.41	0.46
1:VB:219:VAL:HG11	1:VB:227:LEU:HD22	1.98	0.46
1:WA:144:LEU:HD12	1:WA:163:ILE:HD11	1.97	0.46
1:WA:788:ALA:O	1:WA:792:VAL:HG23	2.16	0.46
1:YB:575:ILE:HG23	1:YB:603:VAL:HG22	1.98	0.46
1:Z:543:TYR:CE2	1:Z:575:ILE:HG21	2.51	0.46
1:A:144:LEU:HD12	1:A:163:ILE:HD11	1.97	0.45
1:B:281:TYR:CG	1:B:366:VAL:HG23	2.52	0.45
1:B:398:VAL:HG12	1:B:491:PRO:HB3	1.98	0.45
1:DA:56:ARG:HE	1:EA:127:LEU:HD21	1.80	0.45
1:DA:168:ILE:HG21	1:DA:174:LEU:HB2	1.98	0.45
1:DA:504:ARG:NE	1:DA:504:ARG:HA	2.31	0.45
1:EB:504:ARG:HA	1:EB:504:ARG:NE	2.31	0.45
1:FB:543:TYR:CE2	1:FB:575:ILE:HG21	2.51	0.45
1:I:281:TYR:CG	1:I:366:VAL:HG23	2.51	0.45
1:I:504:ARG:NE	1:I:504:ARG:HA	2.31	0.45
1:I:788:ALA:O	1:I:792:VAL:HG23	2.16	0.45
1:IA:30:VAL:CG2	1:IA:50:MET:HE3	2.45	0.45
1:IB:504:ARG:NE	1:IB:504:ARG:HA	2.31	0.45
1:JB:504:ARG:NE	1:JB:504:ARG:HA	2.31	0.45
1:K:504:ARG:HA	1:K:504:ARG:NE	2.30	0.45
1:M:92:LEU:HD12	1:M:92:LEU:N	2.30	0.45
1:NB:219:VAL:HG11	1:NB:227:LEU:HD22	1.98	0.45
1:NB:523:PHE:CE2	1:NB:545:TRP:CD1	3.05	0.45
1:OA:56:ARG:HE	1:PA:127:LEU:HD21	1.81	0.45
1:PB:168:ILE:HG22	1:PB:215:LEU:CD2	2.43	0.45
1:QA:168:ILE:HG21	1:QA:174:LEU:HB2	1.98	0.45
1:QB:281:TYR:CG	1:QB:366:VAL:HG23	2.51	0.45
1:QB:398:VAL:HG12	1:QB:491:PRO:HB3	1.97	0.45
1:S:92:LEU:HD12	1:S:92:LEU:N	2.30	0.45
1:T:281:TYR:CG	1:T:366:VAL:HG23	2.51	0.45

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:UA:281:TYR:CG	1:UA:366:VAL:HG23	2.51	0.45
1:V:758:GLU:O	1:V:762:VAL:HG22	2.16	0.45
1:VA:56:ARG:HE	1:WA:127:LEU:HD21	1.80	0.45
1:W:92:LEU:HD12	1:W:92:LEU:N	2.30	0.45
1:XA:63:ASN:OD1	1:XA:104:VAL:HG12	2.15	0.45
1:YB:168:ILE:HG21	1:YB:174:LEU:HB2	1.97	0.45
1:YB:219:VAL:HG11	1:YB:227:LEU:HD22	1.98	0.45
1:Z:13:TYR:HA	1:Z:50:MET:HE3	1.97	0.45
1:Z:504:ARG:NE	1:Z:504:ARG:HA	2.31	0.45
1:AA:398:VAL:HG12	1:AA:491:PRO:HB3	1.98	0.45
1:AB:127:LEU:HD21	1:ZA:56:ARG:HE	1.80	0.45
1:AB:504:ARG:NE	1:AB:504:ARG:HA	2.30	0.45
1:AC:779:LEU:HD21	1:OA:774:ARG:HG2	1.98	0.45
1:C:504:ARG:HA	1:C:504:ARG:NE	2.31	0.45
1:DA:144:LEU:HD12	1:DA:163:ILE:HD11	1.97	0.45
1:EA:788:ALA:O	1:EA:792:VAL:HG23	2.17	0.45
1:FB:7:ILE:O	1:FB:8:ILE:C	2.59	0.45
1:FB:168:ILE:HG21	1:FB:174:LEU:HB2	1.97	0.45
1:GB:7:ILE:O	1:GB:8:ILE:C	2.60	0.45
1:H:281:TYR:CG	1:H:366:VAL:HG23	2.51	0.45
1:IA:398:VAL:HG12	1:IA:491:PRO:HB3	1.97	0.45
1:KB:17:HIS:HB3	1:KB:44:LEU:HD23	1.98	0.45
1:KB:765:VAL:HG22	1:LB:759:LEU:HD21	1.97	0.45
1:L:398:VAL:HG12	1:L:491:PRO:HB3	1.97	0.45
1:LA:7:ILE:O	1:LA:8:ILE:C	2.60	0.45
1:LA:219:VAL:HG11	1:LA:227:LEU:HD22	1.98	0.45
1:MB:281:TYR:CG	1:MB:366:VAL:HG23	2.51	0.45
1:NA:7:ILE:O	1:NA:8:ILE:C	2.59	0.45
1:O:219:VAL:HG11	1:O:227:LEU:HD13	1.98	0.45
1:O:529:ILE:HD12	1:O:583:VAL:HG11	1.98	0.45
1:OA:334:LEU:HD12	1:OA:357:TRP:NE1	2.31	0.45
1:OA:504:ARG:NE	1:OA:504:ARG:HA	2.31	0.45
1:OB:538:GLN:HB2	1:OB:646:VAL:HG22	1.98	0.45
1:P:529:ILE:CD1	1:P:583:VAL:HG11	2.46	0.45
1:R:281:TYR:CG	1:R:366:VAL:HG23	2.50	0.45
1:T:334:LEU:HD12	1:T:357:TRP:NE1	2.31	0.45
1:TB:398:VAL:HG12	1:TB:491:PRO:HB3	1.97	0.45
1:V:30:VAL:HG21	1:V:50:MET:HE3	1.99	0.45
1:V:144:LEU:HD12	1:V:163:ILE:HD11	1.97	0.45
1:VA:758:GLU:O	1:VA:762:VAL:HG22	2.16	0.45
1:W:539:LEU:HD11	1:W:640:VAL:HG13	1.96	0.45

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:W:788:ALA:O	1:W:792:VAL:HG23	2.16	0.45
1:WA:7:ILE:O	1:WA:8:ILE:C	2.59	0.45
1:X:398:VAL:HG12	1:X:491:PRO:HB3	1.97	0.45
1:XB:219:VAL:HG11	1:XB:227:LEU:HD22	1.98	0.45
1:XB:366:VAL:HG13	1:XB:366:VAL:O	2.17	0.45
1:YB:57:HIS:O	1:YB:58:TYR:HD1	2.00	0.45
1:AA:7:ILE:O	1:AA:8:ILE:C	2.59	0.45
1:AA:168:ILE:HG21	1:AA:174:LEU:HB2	1.98	0.45
1:AC:281:TYR:CG	1:AC:366:VAL:HG23	2.51	0.45
1:BB:52:THR:O	1:BB:109:ILE:HD11	2.16	0.45
1:BB:366:VAL:HG13	1:BB:366:VAL:O	2.16	0.45
1:CB:7:ILE:O	1:CB:8:ILE:C	2.60	0.45
1:D:788:ALA:O	1:D:792:VAL:HG23	2.16	0.45
1:DA:7:ILE:O	1:DA:8:ILE:C	2.59	0.45
1:EA:13:TYR:HA	1:EA:50:MET:CE	2.46	0.45
1:EB:219:VAL:HG11	1:EB:227:LEU:HD22	1.98	0.45
1:FA:17:HIS:CB	1:FA:44:LEU:HD23	2.47	0.45
1:FB:504:ARG:NE	1:FB:504:ARG:HA	2.30	0.45
1:GA:583:VAL:O	1:GA:583:VAL:HG22	2.16	0.45
1:HA:504:ARG:NE	1:HA:504:ARG:HA	2.31	0.45
1:HB:398:VAL:HG12	1:HB:491:PRO:HB3	1.97	0.45
1:HB:504:ARG:HA	1:HB:504:ARG:NE	2.30	0.45
1:I:144:LEU:HD12	1:I:163:ILE:HD11	1.97	0.45
1:IB:737:GLY:O	1:IB:741:VAL:HG23	2.16	0.45
1:K:543:TYR:CE2	1:K:575:ILE:HG21	2.51	0.45
1:MA:529:ILE:HD12	1:MA:583:VAL:HG11	1.97	0.45
1:NA:529:ILE:CD1	1:NA:583:VAL:HG11	2.45	0.45
1:P:504:ARG:NE	1:P:504:ARG:HA	2.31	0.45
1:QA:487:VAL:O	1:QA:487:VAL:HG13	2.17	0.45
1:RA:7:ILE:O	1:RA:8:ILE:C	2.59	0.45
1:WA:504:ARG:NE	1:WA:504:ARG:HA	2.30	0.45
1:XA:504:ARG:NE	1:XA:504:ARG:HA	2.31	0.45
1:XB:168:ILE:HG21	1:XB:174:LEU:HB2	1.98	0.45
1:XB:543:TYR:CE2	1:XB:575:ILE:HG21	2.50	0.45
1:A:7:ILE:O	1:A:8:ILE:C	2.60	0.45
1:A:281:TYR:CG	1:A:366:VAL:HG23	2.51	0.45
1:A:543:TYR:CE2	1:A:575:ILE:HG21	2.52	0.45
1:AC:7:ILE:O	1:AC:8:ILE:C	2.59	0.45
1:CB:281:TYR:CG	1:CB:366:VAL:HG23	2.52	0.45
1:CB:543:TYR:CE2	1:CB:575:ILE:HG21	2.52	0.45
1:DB:758:GLU:O	1:DB:762:VAL:HG22	2.16	0.45

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:EA:7:ILE:O	1:EA:8:ILE:C	2.59	0.45
1:EB:788:ALA:O	1:EB:792:VAL:HG23	2.17	0.45
1:G:281:TYR:CG	1:G:366:VAL:HG23	2.51	0.45
1:GA:7:ILE:O	1:GA:8:ILE:C	2.59	0.45
1:GA:334:LEU:HD12	1:GA:357:TRP:NE1	2.32	0.45
1:HA:7:ILE:O	1:HA:8:ILE:C	2.59	0.45
1:HB:788:ALA:O	1:HB:792:VAL:HG23	2.16	0.45
1:IB:7:ILE:O	1:IB:8:ILE:C	2.59	0.45
1:J:17:HIS:HB3	1:J:44:LEU:HD23	1.97	0.45
1:J:63:ASN:OD1	1:J:104:VAL:HG12	2.16	0.45
1:JB:788:ALA:O	1:JB:792:VAL:HG23	2.16	0.45
1:LB:17:HIS:HB3	1:LB:44:LEU:HD23	1.98	0.45
1:MA:7:ILE:O	1:MA:8:ILE:C	2.59	0.45
1:MA:52:THR:O	1:MA:109:ILE:HD11	2.16	0.45
1:NA:539:LEU:HD21	1:NA:599:ILE:HD11	1.98	0.45
1:O:56:ARG:HE	1:P:127:LEU:HD21	1.80	0.45
1:P:758:GLU:O	1:P:762:VAL:HG22	2.16	0.45
1:PA:543:TYR:CE2	1:PA:575:ILE:HG21	2.51	0.45
1:QB:504:ARG:NE	1:QB:504:ARG:HA	2.31	0.45
1:QB:543:TYR:CE2	1:QB:575:ILE:HG21	2.51	0.45
1:SB:334:LEU:HD12	1:SB:357:TRP:NE1	2.32	0.45
1:SB:366:VAL:HG13	1:SB:366:VAL:O	2.17	0.45
1:UA:92:LEU:HD12	1:UA:92:LEU:N	2.31	0.45
1:UB:487:VAL:HG13	1:UB:487:VAL:O	2.16	0.45
1:WB:7:ILE:O	1:WB:8:ILE:C	2.59	0.45
1:XB:17:HIS:HB3	1:XB:44:LEU:HD23	1.97	0.45
1:XB:497:VAL:O	1:XB:497:VAL:HG13	2.16	0.45
1:Y:7:ILE:O	1:Y:8:ILE:C	2.59	0.45
1:Y:487:VAL:O	1:Y:487:VAL:HG13	2.16	0.45
1:ZA:7:ILE:O	1:ZA:8:ILE:C	2.60	0.45
1:ZA:529:ILE:HD12	1:ZA:583:VAL:HG11	1.97	0.45
1:ZB:92:LEU:HD12	1:ZB:92:LEU:N	2.30	0.45
1:A:398:VAL:HG12	1:A:491:PRO:HB3	1.97	0.45
1:A:504:ARG:NE	1:A:504:ARG:HA	2.31	0.45
1:BB:56:ARG:HE	1:CB:127:LEU:HD21	1.81	0.45
1:CA:398:VAL:HG12	1:CA:491:PRO:HB3	1.96	0.45
1:CB:219:VAL:HG11	1:CB:227:LEU:HD13	1.99	0.45
1:CB:497:VAL:O	1:CB:497:VAL:HG13	2.15	0.45
1:FA:504:ARG:HA	1:FA:504:ARG:NE	2.31	0.45
1:G:543:TYR:CE2	1:G:575:ILE:HG21	2.52	0.45
1:GB:788:ALA:O	1:GB:792:VAL:HG23	2.16	0.45

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:HB:7:ILE:O	1:HB:8:ILE:C	2.59	0.45
1:K:56:ARG:HE	1:L:127:LEU:HD21	1.82	0.45
1:KA:7:ILE:O	1:KA:8:ILE:C	2.59	0.45
1:KA:168:ILE:HG21	1:KA:174:LEU:HB2	1.98	0.45
1:KA:487:VAL:HG13	1:KA:487:VAL:O	2.17	0.45
1:L:334:LEU:HD12	1:L:357:TRP:NE1	2.32	0.45
1:L:504:ARG:NE	1:L:504:ARG:HA	2.30	0.45
1:NA:17:HIS:HB3	1:NA:44:LEU:HD23	1.98	0.45
1:NA:758:GLU:O	1:NA:762:VAL:HG22	2.16	0.45
1:O:539:LEU:HD21	1:O:541:LEU:HD21	1.97	0.45
1:Q:788:ALA:O	1:Q:792:VAL:HG23	2.17	0.45
1:QA:539:LEU:HD21	1:QA:599:ILE:HD11	1.99	0.45
1:RA:219:VAL:HG11	1:RA:227:LEU:HD22	1.98	0.45
1:SA:47:PRO:O	1:SA:48:MET:HE2	2.17	0.45
1:V:281:TYR:CG	1:V:366:VAL:HG23	2.51	0.45
1:VA:30:VAL:HG21	1:VA:50:MET:HE3	1.99	0.45
1:VA:489:LEU:HD21	1:VA:495:PHE:CZ	2.52	0.45
1:VB:398:VAL:HG12	1:VB:491:PRO:HB3	1.97	0.45
1:W:7:ILE:O	1:W:8:ILE:C	2.60	0.45
1:XA:7:ILE:O	1:XA:8:ILE:C	2.60	0.45
1:XA:788:ALA:O	1:XA:792:VAL:HG23	2.16	0.45
1:XB:281:TYR:CG	1:XB:366:VAL:HG23	2.51	0.45
1:YA:529:ILE:CD1	1:YA:583:VAL:HG11	2.45	0.45
1:AC:168:ILE:HG21	1:AC:174:LEU:HB2	1.98	0.45
1:C:758:GLU:O	1:C:762:VAL:HG22	2.17	0.45
1:CB:504:ARG:HA	1:CB:504:ARG:NE	2.31	0.45
1:DB:504:ARG:NE	1:DB:504:ARG:HA	2.31	0.45
1:F:543:TYR:CE2	1:F:575:ILE:HG21	2.52	0.45
1:F:804:PRO:O	1:F:807:ILE:HG22	2.15	0.45
1:H:489:LEU:HD21	1:H:495:PHE:CZ	2.52	0.45
1:HA:281:TYR:CG	1:HA:366:VAL:HG23	2.51	0.45
1:HB:56:ARG:HE	1:IB:127:LEU:HD21	1.82	0.45
1:I:758:GLU:O	1:I:762:VAL:HG22	2.16	0.45
1:IB:398:VAL:HG12	1:IB:491:PRO:HB3	1.98	0.45
1:JA:17:HIS:HB3	1:JA:44:LEU:HD23	1.99	0.45
1:JB:398:VAL:HG12	1:JB:491:PRO:HB3	1.98	0.45
1:KA:25:VAL:HG21	1:KA:78:THR:O	2.16	0.45
1:KB:386:GLU:O	1:KB:402:ILE:HG23	2.16	0.45
1:MB:7:ILE:O	1:MB:8:ILE:C	2.59	0.45
1:NB:398:VAL:HG12	1:NB:491:PRO:HB3	1.98	0.45
1:O:398:VAL:HG12	1:O:491:PRO:HB3	1.98	0.45

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:OB:504:ARG:NE	1:OB:504:ARG:HA	2.31	0.45
1:PA:53:VAL:HG21	1:PA:59:CYS:HB2	1.99	0.45
1:PA:281:TYR:CG	1:PA:366:VAL:HG23	2.51	0.45
1:QB:7:ILE:O	1:QB:8:ILE:C	2.59	0.45
1:SB:504:ARG:HA	1:SB:504:ARG:NE	2.31	0.45
1:T:543:TYR:CE2	1:T:575:ILE:HG21	2.51	0.45
1:X:17:HIS:CB	1:X:44:LEU:HD23	2.47	0.45
1:X:575:ILE:HG23	1:X:603:VAL:HG22	1.98	0.45
1:XB:7:ILE:O	1:XB:8:ILE:C	2.59	0.45
1:YA:7:ILE:O	1:YA:8:ILE:C	2.59	0.45
1:Z:92:LEU:HD12	1:Z:92:LEU:N	2.31	0.45
1:ZA:168:ILE:HG21	1:ZA:174:LEU:HB2	1.99	0.45
1:ZB:168:ILE:HG21	1:ZB:174:LEU:HB2	1.98	0.45
1:AA:575:ILE:HG23	1:AA:603:VAL:HG22	1.98	0.45
1:BA:366:VAL:O	1:BA:366:VAL:HG13	2.16	0.45
1:D:7:ILE:O	1:D:8:ILE:C	2.60	0.45
1:DA:643:VAL:O	1:DA:643:VAL:HG12	2.17	0.45
1:DB:583:VAL:O	1:DB:583:VAL:HG22	2.16	0.45
1:EA:504:ARG:NE	1:EA:504:ARG:HA	2.31	0.45
1:G:393:VAL:HG22	1:G:411:ASP:C	2.42	0.45
1:KA:529:ILE:HD12	1:KA:583:VAL:HG11	1.98	0.45
1:KB:543:TYR:CE2	1:KB:575:ILE:HG21	2.52	0.45
1:KB:758:GLU:O	1:KB:762:VAL:HG22	2.17	0.45
1:N:281:TYR:CG	1:N:366:VAL:HG23	2.51	0.45
1:OA:497:VAL:O	1:OA:497:VAL:HG13	2.15	0.45
1:P:583:VAL:HG22	1:P:583:VAL:O	2.16	0.45
1:Q:281:TYR:CG	1:Q:366:VAL:HG23	2.51	0.45
1:R:168:ILE:HG21	1:R:174:LEU:HB2	1.98	0.45
1:S:366:VAL:HG13	1:S:366:VAL:O	2.17	0.45
1:TA:219:VAL:HG11	1:TA:227:LEU:HD22	1.99	0.45
1:TB:281:TYR:CG	1:TB:366:VAL:HG23	2.52	0.45
1:W:543:TYR:CE2	1:W:575:ILE:HG21	2.51	0.45
1:WB:504:ARG:NE	1:WB:504:ARG:HA	2.30	0.45
1:Z:788:ALA:O	1:Z:792:VAL:HG23	2.17	0.45
1:ZA:366:VAL:O	1:ZA:366:VAL:HG13	2.17	0.45
1:AB:543:TYR:CE2	1:AB:575:ILE:HG21	2.52	0.45
1:B:219:VAL:HG11	1:B:227:LEU:HD22	1.98	0.45
1:DA:176:LEU:HD23	1:DA:176:LEU:H	1.82	0.45
1:DA:543:TYR:CE2	1:DA:575:ILE:HG21	2.51	0.45
1:E:7:ILE:O	1:E:8:ILE:C	2.60	0.45
1:G:398:VAL:HG12	1:G:491:PRO:HB3	1.98	0.45

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:GB:168:ILE:HG21	1:GB:174:LEU:HB2	1.98	0.45
1:H:398:VAL:HG12	1:H:491:PRO:HB3	1.97	0.45
1:HA:176:LEU:HD23	1:HA:176:LEU:H	1.82	0.45
1:HB:30:VAL:HG21	1:HB:50:MET:HE3	1.99	0.45
1:IB:121:LEU:HD23	1:IB:162:ILE:HG22	1.99	0.45
1:K:366:VAL:HG13	1:K:366:VAL:O	2.17	0.45
1:KB:575:ILE:HG23	1:KB:603:VAL:HG22	1.96	0.45
1:LA:63:ASN:OD1	1:LA:104:VAL:HG12	2.16	0.45
1:LA:168:ILE:HG21	1:LA:174:LEU:HB2	1.98	0.45
1:LA:334:LEU:HD12	1:LA:357:TRP:NE1	2.32	0.45
1:MB:366:VAL:HG13	1:MB:366:VAL:O	2.17	0.45
1:MB:504:ARG:HA	1:MB:504:ARG:NE	2.31	0.45
1:MB:788:ALA:O	1:MB:792:VAL:HG23	2.17	0.45
1:OA:7:ILE:O	1:OA:8:ILE:C	2.59	0.45
1:OA:758:GLU:O	1:OA:762:VAL:HG22	2.16	0.45
1:R:539:LEU:HD12	1:R:540:GLN:H	1.80	0.45
1:RB:758:GLU:O	1:RB:762:VAL:HG22	2.17	0.45
1:UA:497:VAL:HG13	1:UA:497:VAL:O	2.15	0.45
1:UB:7:ILE:O	1:UB:8:ILE:C	2.59	0.45
1:V:504:ARG:HA	1:V:504:ARG:NE	2.31	0.45
1:WA:758:GLU:O	1:WA:762:VAL:HG22	2.16	0.45
1:WB:543:TYR:CE2	1:WB:575:ILE:HG21	2.51	0.45
1:Z:219:VAL:HG11	1:Z:227:LEU:HD22	1.99	0.45
1:AA:30:VAL:HG21	1:AA:50:MET:HE3	1.98	0.45
1:AA:489:LEU:HD21	1:AA:495:PHE:CZ	2.52	0.45
1:B:7:ILE:O	1:B:8:ILE:C	2.60	0.45
1:B:176:LEU:HD23	1:B:176:LEU:H	1.82	0.45
1:BA:7:ILE:O	1:BA:8:ILE:C	2.60	0.45
1:BA:758:GLU:O	1:BA:762:VAL:HG22	2.16	0.45
1:C:281:TYR:CG	1:C:366:VAL:HG23	2.52	0.45
1:C:366:VAL:HG13	1:C:366:VAL:O	2.17	0.45
1:D:302:VAL:HG11	1:D:306:LYS:HZ2	1.81	0.45
1:DA:216:VAL:O	1:DA:216:VAL:HG13	2.17	0.45
1:DB:487:VAL:O	1:DB:487:VAL:HG13	2.17	0.45
1:E:281:TYR:CG	1:E:366:VAL:HG23	2.52	0.45
1:FB:758:GLU:O	1:FB:762:VAL:HG22	2.17	0.45
1:GB:52:THR:O	1:GB:109:ILE:HD11	2.16	0.45
1:H:504:ARG:NE	1:H:504:ARG:HA	2.30	0.45
1:HB:168:ILE:HG21	1:HB:174:LEU:HB2	1.98	0.45
1:HB:176:LEU:HD23	1:HB:176:LEU:H	1.82	0.45
1:IA:30:VAL:HG21	1:IA:50:MET:HE3	1.99	0.45

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:IA:219:VAL:HG11	1:IA:227:LEU:HD22	1.98	0.45
1:IB:56:ARG:HE	1:JB:127:LEU:CD2	2.30	0.45
1:JA:504:ARG:NE	1:JA:504:ARG:HA	2.31	0.45
1:JB:334:LEU:HD12	1:JB:357:TRP:NE1	2.31	0.45
1:KB:497:VAL:O	1:KB:497:VAL:HG13	2.16	0.45
1:L:281:TYR:CG	1:L:366:VAL:HG23	2.51	0.45
1:O:7:ILE:O	1:O:8:ILE:C	2.60	0.45
1:P:487:VAL:HG13	1:P:487:VAL:O	2.17	0.45
1:RB:504:ARG:NE	1:RB:504:ARG:HA	2.31	0.45
1:SB:17:HIS:HB3	1:SB:44:LEU:HD23	1.98	0.45
1:SB:281:TYR:CG	1:SB:366:VAL:HG23	2.51	0.45
1:T:7:ILE:O	1:T:8:ILE:C	2.59	0.45
1:TB:7:ILE:O	1:TB:8:ILE:C	2.59	0.45
1:Y:543:TYR:CE2	1:Y:575:ILE:HG21	2.51	0.45
1:ZA:662:ILE:O	1:ZA:666:THR:HG23	2.17	0.45
1:ZB:7:ILE:O	1:ZB:8:ILE:C	2.59	0.45
1:AB:176:LEU:HD23	1:AB:176:LEU:H	1.82	0.45
1:BA:788:ALA:O	1:BA:792:VAL:HG23	2.17	0.45
1:BB:7:ILE:O	1:BB:8:ILE:C	2.60	0.45
1:BB:543:TYR:CE2	1:BB:575:ILE:HG21	2.52	0.45
1:CA:487:VAL:O	1:CA:487:VAL:HG13	2.17	0.45
1:DA:788:ALA:O	1:DA:792:VAL:HG23	2.17	0.45
1:DB:219:VAL:HG11	1:DB:227:LEU:HD22	1.99	0.45
1:EB:176:LEU:HD23	1:EB:176:LEU:H	1.82	0.45
1:FB:334:LEU:HD12	1:FB:357:TRP:NE1	2.32	0.45
1:GB:17:HIS:HB3	1:GB:44:LEU:HD23	1.99	0.45
1:H:52:THR:O	1:H:109:ILE:HD11	2.17	0.45
1:I:7:ILE:O	1:I:8:ILE:C	2.59	0.45
1:IB:176:LEU:HD23	1:IB:176:LEU:H	1.82	0.45
1:IB:334:LEU:HD12	1:IB:357:TRP:NE1	2.32	0.45
1:KB:334:LEU:HD12	1:KB:357:TRP:NE1	2.32	0.45
1:L:17:HIS:HB3	1:L:44:LEU:HD23	1.97	0.45
1:LB:7:ILE:O	1:LB:8:ILE:C	2.59	0.45
1:N:219:VAL:HG11	1:N:227:LEU:HD22	1.99	0.45
1:OA:398:VAL:HG12	1:OA:491:PRO:HB3	1.97	0.45
1:P:17:HIS:HB3	1:P:44:LEU:HD23	1.99	0.45
1:P:52:THR:O	1:P:109:ILE:HD11	2.17	0.45
1:QA:366:VAL:HG13	1:QA:366:VAL:O	2.17	0.45
1:R:7:ILE:O	1:R:8:ILE:C	2.59	0.45
1:R:30:VAL:CG2	1:R:50:MET:HE3	2.46	0.45
1:R:543:TYR:CE2	1:R:575:ILE:HG21	2.51	0.45

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:SB:398:VAL:HG12	1:SB:491:PRO:HB3	1.98	0.45
1:V:575:ILE:HG23	1:V:603:VAL:HG22	1.98	0.45
1:W:219:VAL:HG11	1:W:227:LEU:HD22	1.99	0.45
1:WB:281:TYR:CG	1:WB:366:VAL:HG23	2.52	0.45
1:YB:334:LEU:HD12	1:YB:357:TRP:NE1	2.32	0.45
1:Z:281:TYR:CG	1:Z:366:VAL:HG23	2.52	0.45
1:ZA:219:VAL:HG11	1:ZA:227:LEU:HD22	1.99	0.45
1:ZA:538:GLN:HB2	1:ZA:646:VAL:HG22	1.98	0.45
1:ZB:543:TYR:CE2	1:ZB:575:ILE:HG21	2.52	0.45
1:A:366:VAL:O	1:A:366:VAL:HG13	2.18	0.44
1:AA:176:LEU:HD23	1:AA:176:LEU:H	1.82	0.44
1:B:17:HIS:HB3	1:B:44:LEU:HD23	1.99	0.44
1:CA:17:HIS:HB3	1:CA:44:LEU:HD23	1.99	0.44
1:DA:334:LEU:HD12	1:DA:357:TRP:NE1	2.32	0.44
1:DB:52:THR:O	1:DB:109:ILE:HD11	2.17	0.44
1:EB:48:MET:HA	1:EB:48:MET:HE3	1.99	0.44
1:G:366:VAL:O	1:G:366:VAL:HG13	2.17	0.44
1:GA:497:VAL:O	1:GA:497:VAL:HG13	2.17	0.44
1:H:56:ARG:HE	1:I:127:LEU:HD21	1.82	0.44
1:IB:366:VAL:O	1:IB:366:VAL:HG13	2.16	0.44
1:LA:281:TYR:CG	1:LA:366:VAL:HG23	2.52	0.44
1:LB:176:LEU:HD23	1:LB:176:LEU:H	1.82	0.44
1:MA:56:ARG:HE	1:NA:127:LEU:HD21	1.81	0.44
1:NA:583:VAL:HG22	1:NA:583:VAL:O	2.17	0.44
1:NB:7:ILE:O	1:NB:8:ILE:C	2.59	0.44
1:NB:489:LEU:HD21	1:NB:495:PHE:CZ	2.52	0.44
1:O:168:ILE:HG21	1:O:174:LEU:HB2	1.98	0.44
1:PB:7:ILE:O	1:PB:8:ILE:C	2.59	0.44
1:S:168:ILE:HG22	1:S:215:LEU:CD2	2.46	0.44
1:SB:7:ILE:O	1:SB:8:ILE:C	2.59	0.44
1:UA:168:ILE:HG21	1:UA:174:LEU:HB2	1.99	0.44
1:UB:56:ARG:HE	1:VB:127:LEU:CD2	2.31	0.44
1:V:398:VAL:HG12	1:V:491:PRO:HB3	1.98	0.44
1:W:398:VAL:HG12	1:W:491:PRO:HB3	1.98	0.44
1:WB:52:THR:O	1:WB:109:ILE:HD11	2.18	0.44
1:XA:219:VAL:HG11	1:XA:227:LEU:HD22	1.99	0.44
1:YA:56:ARG:HE	1:ZA:127:LEU:HD21	1.82	0.44
1:YB:53:VAL:HG12	1:YB:93:ALA:HA	1.98	0.44
1:ZA:52:THR:O	1:ZA:109:ILE:HD11	2.16	0.44
1:ZA:334:LEU:HD12	1:ZA:357:TRP:NE1	2.32	0.44
1:A:176:LEU:H	1:A:176:LEU:HD23	1.83	0.44

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:A:774:ARG:HG2	1:NA:779:LEU:HD21	1.99	0.44
1:AC:539:LEU:HD21	1:AC:599:ILE:HD11	1.99	0.44
1:BA:53:VAL:HG12	1:BA:57:HIS:O	2.18	0.44
1:CA:176:LEU:HD23	1:CA:176:LEU:H	1.82	0.44
1:DA:208:VAL:HG23	1:DA:209:PHE:CD1	2.52	0.44
1:DB:543:TYR:CE2	1:DB:575:ILE:HG21	2.53	0.44
1:EA:17:HIS:HB3	1:EA:44:LEU:HD23	2.00	0.44
1:EB:7:ILE:O	1:EB:8:ILE:C	2.59	0.44
1:FA:57:HIS:O	1:FA:58:TYR:HD1	2.00	0.44
1:FA:538:GLN:HB2	1:FA:646:VAL:HG22	1.97	0.44
1:G:168:ILE:HG21	1:G:174:LEU:HB2	1.99	0.44
1:G:523:PHE:CE2	1:G:545:TRP:CD1	3.05	0.44
1:H:7:ILE:O	1:H:8:ILE:C	2.60	0.44
1:IB:144:LEU:HD12	1:IB:163:ILE:HD11	1.98	0.44
1:K:359:ILE:HG23	1:K:359:ILE:O	2.17	0.44
1:KB:168:ILE:HG21	1:KB:174:LEU:HB2	1.99	0.44
1:LA:30:VAL:HG21	1:LA:50:MET:HE3	2.00	0.44
1:LA:176:LEU:HD23	1:LA:176:LEU:H	1.83	0.44
1:LB:219:VAL:HG11	1:LB:227:LEU:HD13	1.99	0.44
1:M:7:ILE:O	1:M:8:ILE:C	2.59	0.44
1:NA:168:ILE:HG21	1:NA:174:LEU:HB2	1.98	0.44
1:NA:334:LEU:HD12	1:NA:357:TRP:NE1	2.32	0.44
1:NB:30:VAL:HG21	1:NB:50:MET:HE3	1.98	0.44
1:PA:53:VAL:HG12	1:PA:93:ALA:HA	1.99	0.44
1:Q:219:VAL:HG11	1:Q:227:LEU:HD22	1.98	0.44
1:QA:543:TYR:CE2	1:QA:575:ILE:HG21	2.51	0.44
1:R:219:VAL:HG11	1:R:227:LEU:HD13	2.00	0.44
1:UA:543:TYR:CE2	1:UA:575:ILE:HG21	2.52	0.44
1:V:7:ILE:O	1:V:8:ILE:C	2.60	0.44
1:V:168:ILE:HG22	1:V:215:LEU:CD2	2.46	0.44
1:V:208:VAL:HG23	1:V:209:PHE:CD1	2.52	0.44
1:WA:176:LEU:HD23	1:WA:176:LEU:H	1.83	0.44
1:WB:487:VAL:HG13	1:WB:487:VAL:O	2.17	0.44
1:X:7:ILE:O	1:X:8:ILE:C	2.60	0.44
1:X:121:LEU:HD23	1:X:162:ILE:HG22	1.99	0.44
1:Y:52:THR:O	1:Y:109:ILE:HD11	2.16	0.44
1:YB:281:TYR:CG	1:YB:366:VAL:HG23	2.52	0.44
1:Z:366:VAL:O	1:Z:366:VAL:HG13	2.17	0.44
1:ZB:63:ASN:OD1	1:ZB:104:VAL:HG12	2.17	0.44
1:A:17:HIS:CB	1:A:44:LEU:HD23	2.47	0.44
1:AA:523:PHE:CE2	1:AA:545:TRP:CD1	3.05	0.44

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:AC:774:ARG:HG2	1:ZB:779:LEU:HD21	2.00	0.44
1:BB:487:VAL:HG13	1:BB:487:VAL:O	2.18	0.44
1:CB:52:THR:O	1:CB:109:ILE:HD11	2.17	0.44
1:D:543:TYR:CE2	1:D:575:ILE:HG21	2.52	0.44
1:DA:758:GLU:O	1:DA:762:VAL:HG22	2.17	0.44
1:DB:529:ILE:CD1	1:DB:583:VAL:HG11	2.46	0.44
1:EA:168:ILE:HG21	1:EA:174:LEU:HB2	1.99	0.44
1:EA:176:LEU:HD23	1:EA:176:LEU:H	1.82	0.44
1:F:539:LEU:HD11	1:F:640:VAL:HG13	2.00	0.44
1:GA:281:TYR:CG	1:GA:366:VAL:HG23	2.52	0.44
1:H:30:VAL:HG21	1:H:50:MET:HE3	1.99	0.44
1:I:366:VAL:O	1:I:366:VAL:HG13	2.18	0.44
1:L:219:VAL:HG11	1:L:227:LEU:HD22	1.99	0.44
1:L:662:ILE:O	1:L:666:THR:HG23	2.17	0.44
1:LA:398:VAL:HG12	1:LA:491:PRO:HB3	1.99	0.44
1:M:168:ILE:HG21	1:M:174:LEU:HB2	1.98	0.44
1:OA:168:ILE:HG22	1:OA:215:LEU:CD2	2.45	0.44
1:P:7:ILE:O	1:P:8:ILE:C	2.60	0.44
1:QA:281:TYR:CG	1:QA:366:VAL:HG23	2.52	0.44
1:RA:539:LEU:HD21	1:RA:599:ILE:HD11	2.00	0.44
1:RB:30:VAL:CG2	1:RB:50:MET:HE2	2.47	0.44
1:SA:7:ILE:O	1:SA:8:ILE:C	2.59	0.44
1:SB:219:VAL:HG11	1:SB:227:LEU:HD22	1.99	0.44
1:WA:57:HIS:O	1:WA:58:TYR:HD1	2.00	0.44
1:XA:543:TYR:CE2	1:XA:575:ILE:HG21	2.52	0.44
1:YA:176:LEU:HD23	1:YA:176:LEU:H	1.82	0.44
1:ZA:487:VAL:O	1:ZA:487:VAL:HG13	2.17	0.44
1:AC:583:VAL:HG22	1:AC:583:VAL:O	2.17	0.44
1:BB:758:GLU:O	1:BB:762:VAL:HG22	2.17	0.44
1:C:17:HIS:HB3	1:C:44:LEU:HD23	1.98	0.44
1:C:583:VAL:HG22	1:C:583:VAL:O	2.17	0.44
1:CA:219:VAL:HG11	1:CA:227:LEU:HD22	1.99	0.44
1:CB:92:LEU:N	1:CB:92:LEU:HD12	2.32	0.44
1:DB:168:ILE:HG22	1:DB:215:LEU:CD2	2.47	0.44
1:DB:216:VAL:O	1:DB:216:VAL:HG13	2.18	0.44
1:F:523:PHE:CD2	1:F:568:VAL:HG23	2.52	0.44
1:FB:53:VAL:HG12	1:FB:57:HIS:O	2.18	0.44
1:G:7:ILE:O	1:G:8:ILE:C	2.59	0.44
1:GA:17:HIS:CB	1:GA:44:LEU:HD23	2.47	0.44
1:IB:17:HIS:CB	1:IB:44:LEU:HD23	2.47	0.44
1:K:168:ILE:HG21	1:K:174:LEU:HB2	1.99	0.44

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:K:583:VAL:HG22	1:K:583:VAL:O	2.17	0.44
1:KA:176:LEU:HD23	1:KA:176:LEU:H	1.83	0.44
1:LA:17:HIS:HB3	1:LA:44:LEU:HD23	2.00	0.44
1:LB:758:GLU:O	1:LB:762:VAL:HG22	2.18	0.44
1:N:7:ILE:O	1:N:8:ILE:C	2.60	0.44
1:O:281:TYR:CG	1:O:366:VAL:HG23	2.52	0.44
1:OA:121:LEU:HD23	1:OA:162:ILE:HG22	1.99	0.44
1:P:219:VAL:HG11	1:P:227:LEU:HD22	1.99	0.44
1:PA:398:VAL:HG12	1:PA:491:PRO:HB3	1.97	0.44
1:PA:779:LEU:HD21	1:QA:774:ARG:HG2	2.00	0.44
1:Q:52:THR:O	1:Q:109:ILE:HD11	2.17	0.44
1:RA:52:THR:O	1:RA:109:ILE:HD11	2.17	0.44
1:RA:543:TYR:CE2	1:RA:575:ILE:HG21	2.52	0.44
1:RB:63:ASN:OD1	1:RB:104:VAL:HG12	2.17	0.44
1:S:779:LEU:HD21	1:T:774:ARG:HG2	2.00	0.44
1:SA:281:TYR:CG	1:SA:366:VAL:HG23	2.52	0.44
1:T:487:VAL:O	1:T:487:VAL:HG13	2.17	0.44
1:T:779:LEU:HD21	1:V:774:ARG:HG2	2.00	0.44
1:TA:176:LEU:HD23	1:TA:176:LEU:H	1.82	0.44
1:UB:17:HIS:CB	1:UB:44:LEU:HD23	2.47	0.44
1:UB:168:ILE:HG21	1:UB:174:LEU:HB2	1.98	0.44
1:VA:7:ILE:O	1:VA:8:ILE:C	2.60	0.44
1:X:487:VAL:HG13	1:X:487:VAL:O	2.18	0.44
1:XA:176:LEU:HD23	1:XA:176:LEU:H	1.83	0.44
1:XB:398:VAL:HG12	1:XB:491:PRO:HB3	1.98	0.44
1:XB:529:ILE:HD12	1:XB:583:VAL:HG11	1.98	0.44
1:YB:52:THR:O	1:YB:109:ILE:HD11	2.18	0.44
1:Z:176:LEU:HD23	1:Z:176:LEU:H	1.82	0.44
1:Z:487:VAL:HG13	1:Z:487:VAL:O	2.17	0.44
1:AA:774:ARG:HG2	1:Z:779:LEU:HD21	1.99	0.44
1:AB:56:ARG:HE	1:BB:127:LEU:HD21	1.81	0.44
1:BB:219:VAL:HG11	1:BB:227:LEU:HD22	1.99	0.44
1:D:176:LEU:H	1:D:176:LEU:HD23	1.81	0.44
1:DB:56:ARG:HE	1:EB:127:LEU:HD21	1.80	0.44
1:E:758:GLU:O	1:E:762:VAL:HG22	2.18	0.44
1:EA:63:ASN:OD1	1:EA:104:VAL:HG12	2.17	0.44
1:EB:17:HIS:CB	1:EB:44:LEU:HD23	2.48	0.44
1:F:7:ILE:O	1:F:8:ILE:C	2.59	0.44
1:FA:487:VAL:HG13	1:FA:487:VAL:O	2.18	0.44
1:FB:176:LEU:H	1:FB:176:LEU:HD23	1.82	0.44
1:G:219:VAL:HG11	1:G:227:LEU:HD22	2.00	0.44

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:GA:176:LEU:H	1:GA:176:LEU:HD23	1.82	0.44
1:JA:281:TYR:CG	1:JA:366:VAL:HG23	2.52	0.44
1:JA:487:VAL:O	1:JA:487:VAL:HG13	2.17	0.44
1:LB:398:VAL:HG12	1:LB:491:PRO:HB3	1.99	0.44
1:LB:779:LEU:HD21	1:MB:774:ARG:HG2	1.98	0.44
1:O:543:TYR:CE2	1:O:575:ILE:HG21	2.52	0.44
1:OA:487:VAL:O	1:OA:487:VAL:HG13	2.18	0.44
1:PA:7:ILE:O	1:PA:8:ILE:C	2.60	0.44
1:QB:788:ALA:O	1:QB:792:VAL:HG23	2.17	0.44
1:R:334:LEU:HD12	1:R:357:TRP:NE1	2.33	0.44
1:SB:3:THR:HG22	1:SB:6:PHE:HA	2.00	0.44
1:V:121:LEU:HD23	1:V:162:ILE:HG22	1.99	0.44
1:V:487:VAL:O	1:V:487:VAL:HG13	2.18	0.44
1:VA:63:ASN:OD1	1:VA:104:VAL:HG12	2.17	0.44
1:VA:487:VAL:HG13	1:VA:487:VAL:O	2.18	0.44
1:WA:219:VAL:HG11	1:WA:227:LEU:HD22	2.00	0.44
1:WB:398:VAL:HG12	1:WB:491:PRO:HB3	1.98	0.44
1:X:216:VAL:O	1:X:216:VAL:HG13	2.17	0.44
1:X:543:TYR:CE2	1:X:575:ILE:HG21	2.52	0.44
1:A:121:LEU:HD23	1:A:162:ILE:HG22	1.99	0.44
1:A:208:VAL:HG23	1:A:209:PHE:CD1	2.53	0.44
1:A:219:VAL:HG11	1:A:227:LEU:HD22	2.00	0.44
1:AA:487:VAL:O	1:AA:487:VAL:HG13	2.18	0.44
1:AB:168:ILE:HG21	1:AB:174:LEU:HB2	2.00	0.44
1:AB:487:VAL:O	1:AB:487:VAL:HG13	2.17	0.44
1:AC:127:LEU:HD21	1:ZB:56:ARG:HE	1.82	0.44
1:C:7:ILE:O	1:C:8:ILE:C	2.59	0.44
1:D:393:VAL:HG22	1:D:411:ASP:C	2.43	0.44
1:GB:129:PHE:CE1	1:GB:137:VAL:HG11	2.53	0.44
1:J:219:VAL:HG11	1:J:227:LEU:HD22	1.99	0.44
1:JA:219:VAL:HG11	1:JA:227:LEU:HD22	1.99	0.44
1:K:7:ILE:O	1:K:8:ILE:C	2.60	0.44
1:K:159:VAL:O	1:K:159:VAL:HG12	2.17	0.44
1:L:7:ILE:O	1:L:8:ILE:C	2.60	0.44
1:L:529:ILE:HD13	1:L:583:VAL:HG11	2.00	0.44
1:L:758:GLU:O	1:L:762:VAL:HG22	2.18	0.44
1:LA:366:VAL:O	1:LA:366:VAL:HG13	2.18	0.44
1:M:758:GLU:O	1:M:762:VAL:HG22	2.18	0.44
1:NA:398:VAL:HG12	1:NA:491:PRO:HB3	1.98	0.44
1:QA:17:HIS:HB3	1:QA:44:LEU:HD23	1.99	0.44
1:QA:168:ILE:HG22	1:QA:215:LEU:CD2	2.45	0.44

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:RA:758:GLU:O	1:RA:762:VAL:HG22	2.16	0.44
1:VB:7:ILE:O	1:VB:8:ILE:C	2.59	0.44
1:VB:17:HIS:HB3	1:VB:44:LEU:HD23	1.99	0.44
1:W:121:LEU:HD22	1:W:159:VAL:HG13	2.00	0.44
1:Y:398:VAL:HG12	1:Y:491:PRO:HB3	1.99	0.44
1:YB:7:ILE:O	1:YB:8:ILE:C	2.59	0.44
1:B:48:MET:HE2	1:B:48:MET:HA	1.99	0.44
1:BA:538:GLN:HB2	1:BA:646:VAL:HG22	1.98	0.44
1:CB:366:VAL:O	1:CB:366:VAL:HG13	2.18	0.44
1:DB:176:LEU:HD23	1:DB:176:LEU:H	1.83	0.44
1:E:176:LEU:HD23	1:E:176:LEU:H	1.83	0.44
1:F:393:VAL:HG22	1:F:411:ASP:C	2.43	0.44
1:FB:17:HIS:HB3	1:FB:44:LEU:HD23	2.00	0.44
1:FB:216:VAL:HG13	1:FB:216:VAL:O	2.18	0.44
1:IA:176:LEU:HD23	1:IA:176:LEU:H	1.82	0.44
1:IB:176:LEU:HD21	1:IB:196:TRP:CE2	2.53	0.44
1:IB:330:GLN:C	1:IB:407:MET:HE1	2.42	0.44
1:KA:497:VAL:HG13	1:KA:497:VAL:O	2.16	0.44
1:KB:7:ILE:O	1:KB:8:ILE:C	2.60	0.44
1:KB:30:VAL:CG2	1:KB:50:MET:HE3	2.47	0.44
1:N:398:VAL:HG12	1:N:491:PRO:HB3	1.98	0.44
1:N:543:TYR:CE2	1:N:575:ILE:HG21	2.52	0.44
1:O:366:VAL:O	1:O:366:VAL:HG13	2.18	0.44
1:O:523:PHE:CD2	1:O:568:VAL:HG23	2.52	0.44
1:OB:7:ILE:O	1:OB:8:ILE:C	2.60	0.44
1:PA:44:LEU:N	1:PA:44:LEU:HD22	2.33	0.44
1:PA:176:LEU:HD23	1:PA:176:LEU:H	1.83	0.44
1:Q:17:HIS:CB	1:Q:44:LEU:HD23	2.48	0.44
1:QB:168:ILE:HG21	1:QB:174:LEU:HB2	1.98	0.44
1:RA:121:LEU:HD23	1:RA:162:ILE:HG22	2.00	0.44
1:RA:393:VAL:HG22	1:RA:411:ASP:C	2.42	0.44
1:RB:7:ILE:O	1:RB:8:ILE:C	2.59	0.44
1:SB:219:VAL:HG11	1:SB:227:LEU:HD13	2.00	0.44
1:V:334:LEU:HD12	1:V:357:TRP:NE1	2.33	0.44
1:VB:168:ILE:HG22	1:VB:215:LEU:CD2	2.43	0.44
1:W:281:TYR:CG	1:W:366:VAL:HG23	2.53	0.44
1:ZA:758:GLU:O	1:ZA:762:VAL:HG22	2.17	0.44
1:BB:176:LEU:HD23	1:BB:176:LEU:H	1.82	0.44
1:CB:165:ALA:HB2	1:CB:205:LEU:HD13	2.00	0.44
1:CB:398:VAL:HG12	1:CB:491:PRO:HB3	1.98	0.44
1:E:366:VAL:HG13	1:E:366:VAL:O	2.18	0.44

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:EA:44:LEU:HD22	1:EA:44:LEU:H	1.83	0.44
1:EA:52:THR:O	1:EA:109:ILE:HD11	2.18	0.44
1:FA:366:VAL:HG13	1:FA:366:VAL:O	2.18	0.44
1:FB:176:LEU:HD21	1:FB:196:TRP:CE2	2.53	0.44
1:IA:539:LEU:HD21	1:IA:599:ILE:HD11	1.99	0.44
1:KB:393:VAL:HG22	1:KB:411:ASP:C	2.43	0.44
1:L:176:LEU:H	1:L:176:LEU:HD23	1.82	0.44
1:L:393:VAL:HG22	1:L:411:ASP:C	2.43	0.44
1:M:398:VAL:HG12	1:M:491:PRO:HB3	1.98	0.44
1:MB:219:VAL:HG11	1:MB:227:LEU:HD22	1.99	0.44
1:OB:516:LEU:HD23	1:OB:516:LEU:O	2.18	0.44
1:OB:788:ALA:O	1:OB:792:VAL:HG23	2.17	0.44
1:QA:176:LEU:HD23	1:QA:176:LEU:H	1.82	0.44
1:QB:643:VAL:O	1:QB:643:VAL:HG12	2.17	0.44
1:RA:302:VAL:HG11	1:RA:306:LYS:HZ2	1.83	0.44
1:SB:52:THR:O	1:SB:109:ILE:HD11	2.18	0.44
1:TA:543:TYR:CE2	1:TA:575:ILE:HG21	2.52	0.44
1:TB:345:ASP:OD2	1:TB:349:VAL:HG13	2.18	0.44
1:UB:53:VAL:HG12	1:UB:57:HIS:O	2.17	0.44
1:VA:53:VAL:HG12	1:VA:57:HIS:O	2.18	0.44
1:VA:176:LEU:HD23	1:VA:176:LEU:H	1.83	0.44
1:WA:366:VAL:O	1:WA:366:VAL:HG13	2.17	0.44
1:X:176:LEU:HD23	1:X:176:LEU:H	1.83	0.44
1:X:497:VAL:HG13	1:X:497:VAL:O	2.17	0.44
1:YA:168:ILE:HG21	1:YA:174:LEU:HB2	2.00	0.44
1:YA:487:VAL:O	1:YA:487:VAL:HG13	2.18	0.44
1:AC:334:LEU:HD12	1:AC:357:TRP:NE1	2.33	0.44
1:B:366:VAL:O	1:B:366:VAL:HG13	2.17	0.44
1:B:779:LEU:HD21	1:C:774:ARG:HG2	1.99	0.44
1:DA:219:VAL:HG11	1:DA:227:LEU:HD22	2.00	0.44
1:DA:302:VAL:HG11	1:DA:306:LYS:HZ2	1.83	0.44
1:EA:14:HIS:N	1:EA:50:MET:HE3	2.33	0.44
1:H:176:LEU:HD23	1:H:176:LEU:H	1.82	0.44
1:I:216:VAL:O	1:I:216:VAL:HG13	2.18	0.44
1:IA:281:TYR:CG	1:IA:366:VAL:HG23	2.53	0.44
1:J:10:ILE:HG22	1:J:47:PRO:HB3	2.00	0.44
1:J:44:LEU:N	1:J:44:LEU:HD22	2.33	0.44
1:J:543:TYR:CE2	1:J:575:ILE:HG21	2.53	0.44
1:JA:176:LEU:HD23	1:JA:176:LEU:H	1.83	0.44
1:K:176:LEU:HD21	1:K:196:TRP:CE2	2.53	0.44
1:L:129:PHE:CE1	1:L:137:VAL:HG11	2.53	0.44

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:L:366:VAL:O	1:L:366:VAL:HG13	2.18	0.44
1:LA:523:PHE:CD2	1:LA:568:VAL:HG23	2.53	0.44
1:NA:176:LEU:HD23	1:NA:176:LEU:H	1.82	0.44
1:PA:366:VAL:HG13	1:PA:366:VAL:O	2.17	0.44
1:PB:219:VAL:HG11	1:PB:227:LEU:HD22	1.99	0.44
1:QA:758:GLU:O	1:QA:762:VAL:HG22	2.17	0.44
1:RB:44:LEU:HD22	1:RB:44:LEU:N	2.33	0.44
1:TA:44:LEU:HD22	1:TA:44:LEU:N	2.33	0.44
1:UB:176:LEU:HD21	1:UB:196:TRP:CE2	2.53	0.44
1:UB:393:VAL:HG22	1:UB:411:ASP:C	2.43	0.44
1:UB:538:GLN:HB2	1:UB:646:VAL:HG22	2.00	0.44
1:VA:165:ALA:HB2	1:VA:205:LEU:HD13	2.00	0.44
1:X:334:LEU:HD12	1:X:357:TRP:NE1	2.33	0.44
1:XA:386:GLU:O	1:XA:402:ILE:HG23	2.18	0.44
1:XA:393:VAL:HG22	1:XA:411:ASP:C	2.43	0.44
1:XA:737:GLY:O	1:XA:741:VAL:HG23	2.18	0.44
1:Y:17:HIS:HB3	1:Y:44:LEU:HD23	2.00	0.44
1:YA:583:VAL:HG22	1:YA:583:VAL:O	2.17	0.44
1:ZA:176:LEU:H	1:ZA:176:LEU:HD23	1.83	0.44
1:AC:176:LEU:HD23	1:AC:176:LEU:H	1.82	0.43
1:CA:529:ILE:HD12	1:CA:583:VAL:HG11	1.98	0.43
1:CB:57:HIS:O	1:CB:58:TYR:HD1	2.00	0.43
1:E:168:ILE:HG21	1:E:174:LEU:HB2	2.00	0.43
1:EA:176:LEU:HD21	1:EA:196:TRP:CE2	2.53	0.43
1:EA:487:VAL:O	1:EA:487:VAL:HG13	2.18	0.43
1:FB:219:VAL:HG11	1:FB:227:LEU:HD13	2.00	0.43
1:GB:219:VAL:HG11	1:GB:227:LEU:HD13	2.00	0.43
1:GB:366:VAL:HG13	1:GB:366:VAL:O	2.17	0.43
1:GB:758:GLU:O	1:GB:762:VAL:HG22	2.17	0.43
1:HA:176:LEU:HD21	1:HA:196:TRP:NE1	2.33	0.43
1:HB:17:HIS:HB3	1:HB:44:LEU:HD23	2.00	0.43
1:IA:345:ASP:OD2	1:IA:349:VAL:HG13	2.17	0.43
1:K:44:LEU:HD22	1:K:44:LEU:N	2.33	0.43
1:K:539:LEU:CD1	1:K:640:VAL:HG13	2.48	0.43
1:KA:63:ASN:OD1	1:KA:104:VAL:HG12	2.17	0.43
1:L:538:GLN:HB2	1:L:646:VAL:HG22	1.99	0.43
1:LB:52:THR:O	1:LB:109:ILE:HD11	2.17	0.43
1:MA:543:TYR:CE2	1:MA:575:ILE:HG21	2.53	0.43
1:N:17:HIS:HB3	1:N:44:LEU:HD23	1.99	0.43
1:O:121:LEU:HD23	1:O:162:ILE:HG22	2.00	0.43
1:O:393:VAL:HG22	1:O:411:ASP:C	2.43	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:OA:44:LEU:HD22	1:OA:44:LEU:N	2.33	0.43
1:OA:176:LEU:HD21	1:OA:196:TRP:CE2	2.53	0.43
1:OB:176:LEU:HD23	1:OB:176:LEU:H	1.83	0.43
1:PB:487:VAL:O	1:PB:487:VAL:HG13	2.17	0.43
1:Q:7:ILE:O	1:Q:8:ILE:C	2.59	0.43
1:Q:393:VAL:HG22	1:Q:411:ASP:C	2.43	0.43
1:QB:366:VAL:O	1:QB:366:VAL:HG13	2.18	0.43
1:R:44:LEU:N	1:R:44:LEU:HD22	2.33	0.43
1:S:17:HIS:CB	1:S:44:LEU:HD23	2.48	0.43
1:SA:52:THR:O	1:SA:109:ILE:HD11	2.18	0.43
1:TA:398:VAL:HG12	1:TA:491:PRO:HB3	2.00	0.43
1:TB:758:GLU:O	1:TB:762:VAL:HG22	2.18	0.43
1:V:219:VAL:HG11	1:V:227:LEU:HD22	2.00	0.43
1:W:176:LEU:HD23	1:W:176:LEU:H	1.83	0.43
1:WB:121:LEU:HD23	1:WB:162:ILE:HG22	2.00	0.43
1:XB:487:VAL:O	1:XB:487:VAL:HG13	2.17	0.43
1:Y:219:VAL:HG11	1:Y:227:LEU:HD13	1.99	0.43
1:YB:761:ARG:HH21	1:YB:761:ARG:HG3	1.83	0.43
1:AB:774:ARG:HG2	1:ZA:779:LEU:HD21	2.00	0.43
1:BA:216:VAL:HG13	1:BA:216:VAL:O	2.18	0.43
1:C:529:ILE:CD1	1:C:583:VAL:HG11	2.47	0.43
1:C:539:LEU:HD21	1:C:599:ILE:HD11	1.99	0.43
1:DA:366:VAL:O	1:DA:366:VAL:HG13	2.18	0.43
1:DB:529:ILE:HD13	1:DB:583:VAL:HG21	2.01	0.43
1:F:17:HIS:CB	1:F:44:LEU:HD23	2.48	0.43
1:FA:168:ILE:HG22	1:FA:215:LEU:CD2	2.46	0.43
1:FA:334:LEU:HD12	1:FA:357:TRP:NE1	2.33	0.43
1:GA:538:GLN:HB2	1:GA:646:VAL:HG22	1.99	0.43
1:HB:219:VAL:HG11	1:HB:227:LEU:HD13	2.00	0.43
1:I:17:HIS:HB3	1:I:44:LEU:HD23	1.99	0.43
1:IA:138:VAL:HG12	1:IA:139:ALA:N	2.34	0.43
1:IB:52:THR:O	1:IB:109:ILE:HD11	2.18	0.43
1:JA:168:ILE:HG21	1:JA:174:LEU:HB2	1.99	0.43
1:JA:597:ARG:O	1:JA:601:THR:HG23	2.18	0.43
1:JB:121:LEU:HD22	1:JB:159:VAL:HG13	1.99	0.43
1:JB:219:VAL:HG11	1:JB:227:LEU:HD22	2.00	0.43
1:JB:281:TYR:CG	1:JB:366:VAL:HG23	2.53	0.43
1:KA:366:VAL:HG13	1:KA:366:VAL:O	2.17	0.43
1:KB:176:LEU:HD23	1:KB:176:LEU:H	1.83	0.43
1:KB:176:LEU:HD21	1:KB:196:TRP:NE1	2.33	0.43
1:LB:56:ARG:HE	1:MB:127:LEU:CD2	2.32	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:PA:30:VAL:HG21	1:PA:50:MET:HE3	2.01	0.43
1:QB:53:VAL:HG12	1:QB:57:HIS:O	2.17	0.43
1:QB:176:LEU:HD21	1:QB:196:TRP:CE2	2.53	0.43
1:R:176:LEU:HD23	1:R:176:LEU:H	1.82	0.43
1:RA:176:LEU:H	1:RA:176:LEU:HD23	1.82	0.43
1:S:758:GLU:O	1:S:762:VAL:HG22	2.18	0.43
1:UB:176:LEU:HD23	1:UB:176:LEU:H	1.82	0.43
1:VA:543:TYR:CE2	1:VA:575:ILE:HG21	2.53	0.43
1:VB:281:TYR:CG	1:VB:366:VAL:HG23	2.53	0.43
1:WA:17:HIS:CB	1:WA:44:LEU:HD23	2.48	0.43
1:WB:219:VAL:HG11	1:WB:227:LEU:HD22	1.99	0.43
1:XB:359:ILE:O	1:XB:359:ILE:HG23	2.18	0.43
1:YA:176:LEU:HD21	1:YA:196:TRP:CE2	2.53	0.43
1:YB:176:LEU:HD21	1:YB:196:TRP:CE2	2.54	0.43
1:B:393:VAL:HG22	1:B:411:ASP:C	2.43	0.43
1:DA:176:LEU:HD21	1:DA:196:TRP:CE2	2.53	0.43
1:FA:176:LEU:HD23	1:FA:176:LEU:H	1.83	0.43
1:GA:487:VAL:O	1:GA:487:VAL:HG13	2.18	0.43
1:GB:176:LEU:HD23	1:GB:176:LEU:H	1.83	0.43
1:H:219:VAL:HG11	1:H:227:LEU:HD22	2.00	0.43
1:HA:366:VAL:HG13	1:HA:366:VAL:O	2.17	0.43
1:HA:523:PHE:CE2	1:HA:545:TRP:CD1	3.06	0.43
1:IA:208:VAL:HG23	1:IA:209:PHE:CD1	2.54	0.43
1:IB:168:ILE:HG21	1:IB:174:LEU:HB2	1.99	0.43
1:L:44:LEU:HD22	1:L:44:LEU:N	2.34	0.43
1:MA:176:LEU:HD23	1:MA:176:LEU:H	1.83	0.43
1:MB:53:VAL:HG12	1:MB:57:HIS:O	2.18	0.43
1:MB:216:VAL:O	1:MB:216:VAL:HG13	2.19	0.43
1:N:487:VAL:O	1:N:487:VAL:HG13	2.17	0.43
1:N:758:GLU:O	1:N:762:VAL:HG22	2.18	0.43
1:NA:129:PHE:CE1	1:NA:137:VAL:HG11	2.53	0.43
1:NB:176:LEU:H	1:NB:176:LEU:HD23	1.82	0.43
1:P:44:LEU:HD22	1:P:44:LEU:N	2.34	0.43
1:P:216:VAL:O	1:P:216:VAL:HG13	2.18	0.43
1:QB:216:VAL:HG13	1:QB:216:VAL:O	2.18	0.43
1:UA:487:VAL:HG13	1:UA:487:VAL:O	2.18	0.43
1:UB:219:VAL:HG11	1:UB:227:LEU:HD22	2.00	0.43
1:WA:487:VAL:O	1:WA:487:VAL:HG13	2.19	0.43
1:WB:176:LEU:HD21	1:WB:196:TRP:CE2	2.54	0.43
1:X:393:VAL:HG22	1:X:411:ASP:C	2.43	0.43
1:Y:176:LEU:HD23	1:Y:176:LEU:H	1.82	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:YB:523:PHE:CD2	1:YB:568:VAL:HG23	2.53	0.43
1:Z:216:VAL:O	1:Z:216:VAL:HG13	2.18	0.43
1:ZB:30:VAL:HG21	1:ZB:50:MET:HE3	1.99	0.43
1:CB:17:HIS:CB	1:CB:44:LEU:HD23	2.49	0.43
1:CB:393:VAL:HG22	1:CB:411:ASP:C	2.43	0.43
1:EB:52:THR:O	1:EB:109:ILE:HD11	2.17	0.43
1:EB:359:ILE:HG23	1:EB:359:ILE:O	2.18	0.43
1:F:597:ARG:O	1:F:601:THR:HG23	2.19	0.43
1:FA:52:THR:O	1:FA:109:ILE:HD11	2.18	0.43
1:FA:543:TYR:CE2	1:FA:575:ILE:HG21	2.53	0.43
1:FB:56:ARG:HE	1:GB:127:LEU:CD2	2.31	0.43
1:H:216:VAL:O	1:H:216:VAL:HG13	2.19	0.43
1:HB:779:LEU:HD21	1:IB:774:ARG:HG2	1.99	0.43
1:IB:208:VAL:HG23	1:IB:209:PHE:CD1	2.54	0.43
1:JB:17:HIS:HB3	1:JB:44:LEU:HD23	2.00	0.43
1:JB:302:VAL:HG11	1:JB:306:LYS:HZ2	1.82	0.43
1:K:758:GLU:O	1:K:762:VAL:HG22	2.19	0.43
1:LA:516:LEU:HD23	1:LA:516:LEU:C	2.44	0.43
1:LA:543:TYR:CE2	1:LA:575:ILE:HG21	2.53	0.43
1:LB:168:ILE:HG21	1:LB:174:LEU:HB2	1.99	0.43
1:M:121:LEU:HD23	1:M:162:ILE:HG22	2.00	0.43
1:M:543:TYR:CE2	1:M:575:ILE:HG21	2.53	0.43
1:MB:359:ILE:HG23	1:MB:359:ILE:O	2.18	0.43
1:N:168:ILE:HG22	1:N:215:LEU:CD2	2.46	0.43
1:OA:219:VAL:HG11	1:OA:227:LEU:HD22	2.00	0.43
1:OB:216:VAL:O	1:OB:216:VAL:HG13	2.18	0.43
1:PA:176:LEU:HD21	1:PA:196:TRP:CE2	2.53	0.43
1:PB:176:LEU:HD21	1:PB:196:TRP:CE2	2.53	0.43
1:QA:13:TYR:HA	1:QA:50:MET:HE3	2.00	0.43
1:RB:176:LEU:HD21	1:RB:196:TRP:CE2	2.53	0.43
1:SA:17:HIS:HB3	1:SA:44:LEU:HD23	2.00	0.43
1:SA:176:LEU:HD23	1:SA:176:LEU:H	1.83	0.43
1:SA:366:VAL:HG13	1:SA:366:VAL:O	2.18	0.43
1:T:52:THR:O	1:T:109:ILE:HD11	2.19	0.43
1:UA:53:VAL:HG12	1:UA:57:HIS:O	2.18	0.43
1:V:129:PHE:CE1	1:V:137:VAL:HG11	2.53	0.43
1:V:366:VAL:HG13	1:V:366:VAL:O	2.18	0.43
1:VA:219:VAL:HG11	1:VA:227:LEU:HD22	2.00	0.43
1:YB:208:VAL:HG23	1:YB:209:PHE:CD1	2.54	0.43
1:YB:543:TYR:CE2	1:YB:575:ILE:HG21	2.53	0.43
1:DA:53:VAL:HG12	1:DA:57:HIS:O	2.19	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:DA:487:VAL:O	1:DA:487:VAL:HG13	2.19	0.43
1:E:53:VAL:HG12	1:E:57:HIS:O	2.18	0.43
1:EA:345:ASP:OD2	1:EA:349:VAL:HG13	2.18	0.43
1:EA:758:GLU:O	1:EA:762:VAL:HG22	2.18	0.43
1:EB:543:TYR:CE2	1:EB:575:ILE:HG21	2.53	0.43
1:F:44:LEU:H	1:F:44:LEU:HD22	1.83	0.43
1:F:219:VAL:HG11	1:F:227:LEU:HD22	2.00	0.43
1:FA:219:VAL:HG11	1:FA:227:LEU:HD22	1.99	0.43
1:H:788:ALA:O	1:H:792:VAL:HG23	2.19	0.43
1:I:359:ILE:HG23	1:I:359:ILE:O	2.19	0.43
1:J:7:ILE:O	1:J:8:ILE:C	2.59	0.43
1:K:219:VAL:HG11	1:K:227:LEU:HD13	2.01	0.43
1:L:779:LEU:HD21	1:M:774:ARG:HG2	2.00	0.43
1:MB:17:HIS:CB	1:MB:44:LEU:HD23	2.49	0.43
1:NA:219:VAL:HG11	1:NA:227:LEU:HD22	2.00	0.43
1:NA:366:VAL:HG13	1:NA:366:VAL:O	2.18	0.43
1:OA:543:TYR:CE2	1:OA:575:ILE:HG21	2.52	0.43
1:QB:219:VAL:HG11	1:QB:227:LEU:HD22	2.00	0.43
1:SA:30:VAL:HG21	1:SA:50:MET:HE3	2.00	0.43
1:SB:56:ARG:HE	1:TB:127:LEU:CD2	2.31	0.43
1:TA:597:ARG:O	1:TA:601:THR:HG23	2.19	0.43
1:UB:165:ALA:HB2	1:UB:205:LEU:HD13	2.00	0.43
1:V:330:GLN:C	1:V:407:MET:HE1	2.43	0.43
1:WB:597:ARG:O	1:WB:601:THR:HG23	2.18	0.43
1:X:121:LEU:HD22	1:X:159:VAL:HG13	2.01	0.43
1:YB:92:LEU:N	1:YB:92:LEU:HD12	2.33	0.43
1:ZB:44:LEU:HD22	1:ZB:44:LEU:N	2.34	0.43
1:A:487:VAL:O	1:A:487:VAL:HG13	2.18	0.43
1:AA:539:LEU:HD21	1:AA:599:ILE:HD11	2.01	0.43
1:B:44:LEU:N	1:B:44:LEU:HD22	2.33	0.43
1:BA:176:LEU:HD23	1:BA:176:LEU:H	1.83	0.43
1:CB:168:ILE:HG21	1:CB:174:LEU:HB2	1.99	0.43
1:EB:393:VAL:HG22	1:EB:411:ASP:C	2.43	0.43
1:FA:541:LEU:CD2	1:FA:640:VAL:HG22	2.48	0.43
1:G:216:VAL:HG13	1:G:216:VAL:O	2.19	0.43
1:H:165:ALA:HB2	1:H:205:LEU:HD13	1.99	0.43
1:H:366:VAL:HG13	1:H:366:VAL:O	2.18	0.43
1:HA:176:LEU:HD21	1:HA:196:TRP:CE2	2.54	0.43
1:I:44:LEU:HD22	1:I:44:LEU:N	2.34	0.43
1:I:53:VAL:HG12	1:I:57:HIS:O	2.18	0.43
1:IA:216:VAL:O	1:IA:216:VAL:HG13	2.18	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:J:129:PHE:CE1	1:J:137:VAL:HG11	2.53	0.43
1:J:366:VAL:O	1:J:366:VAL:HG13	2.19	0.43
1:JA:366:VAL:O	1:JA:366:VAL:HG13	2.19	0.43
1:K:219:VAL:HG11	1:K:227:LEU:HD22	2.01	0.43
1:K:393:VAL:HG22	1:K:411:ASP:C	2.44	0.43
1:KB:489:LEU:HD21	1:KB:495:PHE:CZ	2.54	0.43
1:M:44:LEU:HD22	1:M:44:LEU:N	2.34	0.43
1:MA:216:VAL:HG13	1:MA:216:VAL:O	2.19	0.43
1:N:779:LEU:HD21	1:O:774:ARG:HG2	2.00	0.43
1:NA:216:VAL:O	1:NA:216:VAL:HG13	2.19	0.43
1:O:208:VAL:HG23	1:O:209:PHE:CD1	2.54	0.43
1:OA:176:LEU:H	1:OA:176:LEU:HD23	1.82	0.43
1:P:543:TYR:CE2	1:P:575:ILE:HG21	2.53	0.43
1:Q:176:LEU:HD23	1:Q:176:LEU:H	1.82	0.43
1:S:144:LEU:HD12	1:S:163:ILE:HD11	2.00	0.43
1:S:176:LEU:HD23	1:S:176:LEU:H	1.83	0.43
1:T:176:LEU:H	1:T:176:LEU:HD23	1.82	0.43
1:TA:281:TYR:CG	1:TA:366:VAL:HG23	2.53	0.43
1:TB:487:VAL:HG13	1:TB:487:VAL:O	2.17	0.43
1:UB:44:LEU:HD22	1:UB:44:LEU:N	2.34	0.43
1:V:176:LEU:HD21	1:V:196:TRP:CE2	2.54	0.43
1:VB:334:LEU:HD12	1:VB:357:TRP:NE1	2.33	0.43
1:W:523:PHE:CE2	1:W:545:TRP:CD1	3.06	0.43
1:XA:176:LEU:HD21	1:XA:196:TRP:NE1	2.33	0.43
1:XA:208:VAL:HG23	1:XA:209:PHE:CD1	2.54	0.43
1:YA:359:ILE:HG23	1:YA:359:ILE:O	2.18	0.43
1:YA:539:LEU:CD1	1:YA:640:VAL:HG13	2.48	0.43
1:YB:176:LEU:H	1:YB:176:LEU:HD23	1.82	0.43
1:Z:53:VAL:HG12	1:Z:57:HIS:O	2.18	0.43
1:Z:129:PHE:CE1	1:Z:137:VAL:HG11	2.54	0.43
1:ZA:216:VAL:O	1:ZA:216:VAL:HG13	2.18	0.43
1:ZA:393:VAL:HG22	1:ZA:411:ASP:C	2.44	0.43
1:ZB:219:VAL:HG11	1:ZB:227:LEU:HD22	2.01	0.43
1:ZB:788:ALA:O	1:ZB:792:VAL:HG23	2.18	0.43
1:AC:129:PHE:CE1	1:AC:137:VAL:HG11	2.54	0.43
1:BA:345:ASP:OD2	1:BA:349:VAL:HG13	2.19	0.43
1:C:176:LEU:HD23	1:C:176:LEU:H	1.82	0.43
1:CB:523:PHE:CD2	1:CB:568:VAL:HG23	2.53	0.43
1:F:176:LEU:HD21	1:F:196:TRP:NE1	2.33	0.43
1:FA:3:THR:HG22	1:FA:6:PHE:HA	2.00	0.43
1:FB:48:MET:HA	1:FB:48:MET:HE2	2.01	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:G:487:VAL:O	1:G:487:VAL:HG13	2.18	0.43
1:GA:208:VAL:HG23	1:GA:209:PHE:CD1	2.54	0.43
1:H:543:TYR:CE2	1:H:575:ILE:HG21	2.54	0.43
1:HA:17:HIS:HB3	1:HA:44:LEU:HD23	2.01	0.43
1:HA:216:VAL:O	1:HA:216:VAL:HG13	2.19	0.43
1:I:219:VAL:HG11	1:I:227:LEU:HD22	2.00	0.43
1:I:398:VAL:HG12	1:I:491:PRO:HB3	1.99	0.43
1:I:662:ILE:O	1:I:666:THR:HG23	2.18	0.43
1:J:386:GLU:O	1:J:402:ILE:HG23	2.18	0.43
1:J:393:VAL:HG22	1:J:411:ASP:C	2.43	0.43
1:JA:44:LEU:HD22	1:JA:44:LEU:H	1.84	0.43
1:K:597:ARG:O	1:K:601:THR:HG23	2.19	0.43
1:L:216:VAL:O	1:L:216:VAL:HG13	2.18	0.43
1:L:487:VAL:O	1:L:487:VAL:HG13	2.17	0.43
1:LA:176:LEU:HD21	1:LA:196:TRP:CE2	2.54	0.43
1:LA:208:VAL:HG23	1:LA:209:PHE:CD1	2.54	0.43
1:MA:17:HIS:CB	1:MA:44:LEU:HD23	2.49	0.43
1:MB:487:VAL:HG13	1:MB:487:VAL:O	2.17	0.43
1:MB:779:LEU:HD21	1:NB:774:ARG:HG2	2.00	0.43
1:N:176:LEU:HD23	1:N:176:LEU:H	1.83	0.43
1:NA:44:LEU:H	1:NA:44:LEU:HD22	1.84	0.43
1:NA:52:THR:O	1:NA:109:ILE:HD11	2.18	0.43
1:O:176:LEU:HD23	1:O:176:LEU:H	1.83	0.43
1:PB:176:LEU:HD23	1:PB:176:LEU:H	1.83	0.43
1:RB:168:ILE:HG21	1:RB:174:LEU:HB2	2.00	0.43
1:S:219:VAL:HG11	1:S:227:LEU:HD22	2.01	0.43
1:S:345:ASP:OD2	1:S:349:VAL:HG13	2.18	0.43
1:SB:543:TYR:CE2	1:SB:575:ILE:HG21	2.54	0.43
1:T:219:VAL:HG11	1:T:227:LEU:HD13	2.01	0.43
1:TA:176:LEU:HD21	1:TA:196:TRP:NE1	2.33	0.43
1:TA:393:VAL:HG22	1:TA:411:ASP:C	2.44	0.43
1:TA:487:VAL:O	1:TA:487:VAL:HG13	2.19	0.43
1:TB:176:LEU:HD21	1:TB:196:TRP:CE2	2.54	0.43
1:TB:208:VAL:HG23	1:TB:209:PHE:CD1	2.54	0.43
1:WA:393:VAL:HG22	1:WA:411:ASP:C	2.44	0.43
1:YB:366:VAL:O	1:YB:366:VAL:HG13	2.18	0.43
1:YB:758:GLU:O	1:YB:762:VAL:HG22	2.19	0.43
1:ZA:30:VAL:CG2	1:ZA:50:MET:HE3	2.49	0.43
1:ZB:17:HIS:CB	1:ZB:44:LEU:HD23	2.48	0.43
1:A:539:LEU:CD1	1:A:640:VAL:HG13	2.49	0.43
1:AA:44:LEU:HD22	1:AA:44:LEU:N	2.34	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:AA:176:LEU:HD21	1:AA:196:TRP:CE2	2.54	0.43
1:AC:176:LEU:HD21	1:AC:196:TRP:CE2	2.54	0.43
1:AC:366:VAL:HG13	1:AC:366:VAL:O	2.19	0.43
1:AC:643:VAL:O	1:AC:643:VAL:HG12	2.18	0.43
1:B:570:ASP:OD1	1:B:570:ASP:C	2.62	0.43
1:B:662:ILE:O	1:B:666:THR:HG23	2.19	0.43
1:BB:168:ILE:HG22	1:BB:215:LEU:CD2	2.47	0.43
1:C:487:VAL:O	1:C:487:VAL:HG13	2.17	0.43
1:CB:216:VAL:O	1:CB:216:VAL:HG13	2.19	0.43
1:CB:405:THR:HG21	1:DB:393:VAL:O	2.19	0.43
1:D:539:LEU:CD1	1:D:640:VAL:HG13	2.49	0.43
1:E:539:LEU:CD1	1:E:640:VAL:HG13	2.49	0.43
1:FA:56:ARG:HE	1:GA:127:LEU:CD2	2.31	0.43
1:FB:30:VAL:CG2	1:FB:50:MET:HE3	2.46	0.43
1:G:129:PHE:CE1	1:G:137:VAL:HG11	2.53	0.43
1:G:176:LEU:HD23	1:G:176:LEU:H	1.83	0.43
1:GA:216:VAL:O	1:GA:216:VAL:HG13	2.19	0.43
1:GA:643:VAL:O	1:GA:643:VAL:HG12	2.18	0.43
1:HA:138:VAL:HG12	1:HA:139:ALA:N	2.34	0.43
1:I:129:PHE:CE1	1:I:137:VAL:HG11	2.54	0.43
1:I:176:LEU:HD23	1:I:176:LEU:H	1.83	0.43
1:IA:121:LEU:HD23	1:IA:162:ILE:HG22	2.01	0.43
1:IA:523:PHE:CE2	1:IA:545:TRP:CD1	3.07	0.43
1:IB:487:VAL:O	1:IB:487:VAL:HG13	2.18	0.43
1:J:121:LEU:HD23	1:J:162:ILE:HG22	2.01	0.43
1:J:208:VAL:HG23	1:J:209:PHE:CD1	2.54	0.43
1:K:121:LEU:HD23	1:K:162:ILE:HG22	2.00	0.43
1:K:129:PHE:CE1	1:K:137:VAL:HG11	2.54	0.43
1:K:176:LEU:HD23	1:K:176:LEU:H	1.82	0.43
1:KB:487:VAL:O	1:KB:487:VAL:HG13	2.18	0.43
1:LB:216:VAL:O	1:LB:216:VAL:HG13	2.19	0.43
1:MA:138:VAL:HG12	1:MA:139:ALA:N	2.34	0.43
1:MB:176:LEU:HD23	1:MB:176:LEU:H	1.82	0.43
1:N:44:LEU:HD22	1:N:44:LEU:N	2.34	0.43
1:N:523:PHE:CE2	1:N:545:TRP:CD1	3.07	0.43
1:NB:168:ILE:HG21	1:NB:174:LEU:HB2	2.00	0.43
1:O:17:HIS:HB3	1:O:44:LEU:HD23	2.00	0.43
1:O:44:LEU:HD22	1:O:44:LEU:N	2.34	0.43
1:O:165:ALA:HB2	1:O:205:LEU:HD13	2.00	0.43
1:O:219:VAL:HG11	1:O:227:LEU:HD22	2.01	0.43
1:O:426:LEU:HD23	1:O:427:LEU:HD22	2.00	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:Q:168:ILE:HG21	1:Q:174:LEU:HB2	2.00	0.43
1:Q:543:TYR:CE2	1:Q:575:ILE:HG21	2.54	0.43
1:Q:758:GLU:O	1:Q:762:VAL:HG22	2.18	0.43
1:QA:44:LEU:HD22	1:QA:44:LEU:N	2.34	0.43
1:QB:208:VAL:HG23	1:QB:209:PHE:CD1	2.54	0.43
1:R:176:LEU:HD21	1:R:196:TRP:CE2	2.53	0.43
1:RA:44:LEU:HD22	1:RA:44:LEU:N	2.34	0.43
1:RB:176:LEU:HD23	1:RB:176:LEU:H	1.82	0.43
1:RB:487:VAL:O	1:RB:487:VAL:HG13	2.18	0.43
1:S:44:LEU:HD22	1:S:44:LEU:N	2.34	0.43
1:S:487:VAL:HG13	1:S:487:VAL:O	2.18	0.43
1:SB:44:LEU:HD22	1:SB:44:LEU:N	2.33	0.43
1:SB:176:LEU:HD23	1:SB:176:LEU:H	1.83	0.43
1:SB:386:GLU:O	1:SB:402:ILE:HG23	2.19	0.43
1:TB:216:VAL:O	1:TB:216:VAL:HG13	2.19	0.43
1:TB:597:ARG:O	1:TB:601:THR:HG23	2.19	0.43
1:TB:643:VAL:O	1:TB:643:VAL:HG12	2.19	0.43
1:VB:208:VAL:HG23	1:VB:209:PHE:CD1	2.54	0.43
1:WA:398:VAL:HG12	1:WA:491:PRO:HB3	1.99	0.43
1:XA:216:VAL:HG13	1:XA:216:VAL:O	2.19	0.43
1:XB:52:THR:O	1:XB:109:ILE:HD11	2.18	0.43
1:XB:176:LEU:HD21	1:XB:196:TRP:CE2	2.54	0.43
1:YA:44:LEU:HD22	1:YA:44:LEU:N	2.33	0.43
1:YA:159:VAL:O	1:YA:159:VAL:HG12	2.18	0.43
1:YB:219:VAL:HG11	1:YB:227:LEU:HD13	2.01	0.43
1:A:176:LEU:HD21	1:A:196:TRP:CE2	2.54	0.43
1:A:216:VAL:O	1:A:216:VAL:HG13	2.19	0.43
1:AC:17:HIS:CB	1:AC:44:LEU:HD23	2.49	0.43
1:AC:44:LEU:HD22	1:AC:44:LEU:N	2.34	0.43
1:B:758:GLU:O	1:B:762:VAL:HG22	2.19	0.43
1:BA:539:LEU:HD21	1:BA:599:ILE:HD11	2.01	0.43
1:BB:539:LEU:CD1	1:BB:640:VAL:HG13	2.48	0.43
1:C:786:GLN:OE1	1:D:781:VAL:HG11	2.19	0.43
1:D:121:LEU:HD23	1:D:162:ILE:HG22	2.00	0.43
1:D:216:VAL:HG13	1:D:216:VAL:O	2.19	0.43
1:EB:216:VAL:O	1:EB:216:VAL:HG13	2.19	0.43
1:F:129:PHE:CE1	1:F:137:VAL:HG11	2.54	0.43
1:F:176:LEU:HD23	1:F:176:LEU:H	1.83	0.43
1:FB:44:LEU:HD22	1:FB:44:LEU:N	2.33	0.43
1:GA:176:LEU:HD21	1:GA:196:TRP:CE2	2.53	0.43
1:H:63:ASN:OD1	1:H:104:VAL:HG12	2.18	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:HA:208:VAL:HG23	1:HA:209:PHE:CD1	2.54	0.43
1:HA:219:VAL:HG11	1:HA:227:LEU:HD22	2.01	0.43
1:HA:326:LEU:HD12	1:HA:363:LEU:HD21	2.01	0.43
1:HB:176:LEU:HD21	1:HB:196:TRP:CE2	2.54	0.43
1:I:176:LEU:HD21	1:I:196:TRP:CE2	2.54	0.43
1:I:208:VAL:HG23	1:I:209:PHE:CD1	2.54	0.43
1:IB:219:VAL:HG11	1:IB:227:LEU:HD22	2.00	0.43
1:JA:17:HIS:CB	1:JA:44:LEU:HD23	2.49	0.43
1:KA:44:LEU:H	1:KA:44:LEU:HD22	1.84	0.43
1:KB:216:VAL:O	1:KB:216:VAL:HG13	2.19	0.43
1:KB:597:ARG:O	1:KB:601:THR:HG23	2.19	0.43
1:KB:643:VAL:O	1:KB:643:VAL:HG12	2.19	0.43
1:LA:52:THR:O	1:LA:109:ILE:HD11	2.18	0.43
1:LB:208:VAL:HG23	1:LB:209:PHE:CD1	2.54	0.43
1:M:17:HIS:HB3	1:M:44:LEU:HD23	1.99	0.43
1:MA:779:LEU:HD21	1:NA:774:ARG:HG2	2.00	0.43
1:MB:129:PHE:CE1	1:MB:137:VAL:HG11	2.54	0.43
1:NA:643:VAL:O	1:NA:643:VAL:HG12	2.18	0.43
1:OA:366:VAL:O	1:OA:366:VAL:HG13	2.18	0.43
1:OB:53:VAL:HG12	1:OB:57:HIS:O	2.18	0.43
1:P:176:LEU:HD23	1:P:176:LEU:H	1.83	0.43
1:QA:529:ILE:HD12	1:QA:583:VAL:HG11	2.00	0.43
1:QB:121:LEU:HD23	1:QB:162:ILE:HG22	2.00	0.43
1:RA:176:LEU:HD21	1:RA:196:TRP:CE2	2.53	0.43
1:RA:359:ILE:O	1:RA:359:ILE:HG23	2.19	0.43
1:SA:168:ILE:HG21	1:SA:174:LEU:HB2	2.01	0.43
1:T:216:VAL:O	1:T:216:VAL:HG13	2.17	0.43
1:TA:539:LEU:HD11	1:TA:640:VAL:HG13	2.00	0.43
1:UA:176:LEU:HD21	1:UA:196:TRP:NE1	2.34	0.43
1:UA:176:LEU:HD23	1:UA:176:LEU:H	1.83	0.43
1:UA:219:VAL:HG11	1:UA:227:LEU:HD22	2.00	0.43
1:V:143:TRP:HH2	1:V:157:VAL:HG22	1.84	0.43
1:WA:208:VAL:HG23	1:WA:209:PHE:CD1	2.54	0.43
1:WA:216:VAL:O	1:WA:216:VAL:HG13	2.19	0.43
1:WB:539:LEU:HD11	1:WB:599:ILE:CD1	2.49	0.43
1:X:489:LEU:HD21	1:X:495:PHE:CZ	2.54	0.43
1:XA:487:VAL:HG13	1:XA:487:VAL:O	2.18	0.43
1:XB:529:ILE:HD13	1:XB:583:VAL:HG11	2.00	0.43
1:Y:138:VAL:HG12	1:Y:139:ALA:N	2.33	0.43
1:Y:208:VAL:HG23	1:Y:209:PHE:CD1	2.54	0.43
1:YA:216:VAL:O	1:YA:216:VAL:HG13	2.19	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:YB:516:LEU:HD23	1:YB:516:LEU:C	2.44	0.43
1:ZA:129:PHE:CE1	1:ZA:137:VAL:HG11	2.54	0.43
1:ZB:176:LEU:HD23	1:ZB:176:LEU:H	1.83	0.43
1:ZB:219:VAL:HG11	1:ZB:227:LEU:HD13	2.01	0.43
1:AB:758:GLU:O	1:AB:762:VAL:HG22	2.18	0.43
1:B:168:ILE:HG21	1:B:174:LEU:HB2	2.00	0.43
1:BA:523:PHE:CE2	1:BA:545:TRP:CD1	3.07	0.43
1:BB:523:PHE:CE2	1:BB:545:TRP:CD1	3.07	0.43
1:BB:643:VAL:O	1:BB:643:VAL:HG12	2.19	0.43
1:C:138:VAL:HG12	1:C:139:ALA:N	2.34	0.43
1:CB:176:LEU:HD23	1:CB:176:LEU:H	1.83	0.43
1:D:138:VAL:HG12	1:D:139:ALA:N	2.34	0.43
1:DA:138:VAL:HG12	1:DA:139:ALA:N	2.34	0.43
1:G:662:ILE:O	1:G:666:THR:HG23	2.19	0.43
1:GA:44:LEU:HD22	1:GA:44:LEU:N	2.33	0.43
1:GB:208:VAL:HG23	1:GB:209:PHE:CD1	2.54	0.43
1:GB:779:LEU:HD21	1:HB:774:ARG:HG2	2.00	0.43
1:H:208:VAL:HG23	1:H:209:PHE:CD1	2.54	0.43
1:IA:165:ALA:HB2	1:IA:205:LEU:HD13	2.00	0.43
1:IB:643:VAL:O	1:IB:643:VAL:HG12	2.19	0.43
1:J:176:LEU:HD21	1:J:196:TRP:NE1	2.33	0.43
1:JA:53:VAL:HG12	1:JA:57:HIS:O	2.19	0.43
1:JA:176:LEU:HD21	1:JA:196:TRP:CE2	2.54	0.43
1:JA:216:VAL:HG13	1:JA:216:VAL:O	2.19	0.43
1:K:208:VAL:HG23	1:K:209:PHE:CD1	2.54	0.43
1:LA:138:VAL:HG12	1:LA:139:ALA:N	2.34	0.43
1:LA:216:VAL:O	1:LA:216:VAL:HG13	2.19	0.43
1:M:53:VAL:HG12	1:M:57:HIS:O	2.18	0.43
1:M:208:VAL:HG23	1:M:209:PHE:CD1	2.54	0.43
1:MB:208:VAL:HG23	1:MB:209:PHE:CD1	2.54	0.43
1:NA:138:VAL:HG12	1:NA:139:ALA:N	2.34	0.43
1:NB:208:VAL:HG23	1:NB:209:PHE:CD1	2.54	0.43
1:NB:359:ILE:O	1:NB:359:ILE:HG23	2.18	0.43
1:OA:219:VAL:HG11	1:OA:227:LEU:HD13	2.01	0.43
1:OB:44:LEU:N	1:OB:44:LEU:HD22	2.34	0.43
1:P:529:ILE:HD13	1:P:583:VAL:HG21	2.00	0.43
1:PA:345:ASP:OD2	1:PA:349:VAL:HG13	2.18	0.43
1:PA:393:VAL:HG22	1:PA:411:ASP:C	2.44	0.43
1:Q:359:ILE:O	1:Q:359:ILE:HG23	2.18	0.43
1:QB:176:LEU:H	1:QB:176:LEU:HD23	1.82	0.43
1:RB:17:HIS:CB	1:RB:44:LEU:HD23	2.49	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:RB:44:LEU:HD22	1:RB:44:LEU:H	1.83	0.43
1:SA:758:GLU:O	1:SA:762:VAL:HG22	2.18	0.43
1:SB:539:LEU:HD21	1:SB:599:ILE:HD11	1.99	0.43
1:UA:366:VAL:O	1:UA:366:VAL:HG13	2.18	0.43
1:UB:216:VAL:O	1:UB:216:VAL:HG13	2.19	0.43
1:W:539:LEU:HD21	1:W:599:ILE:HD11	2.00	0.43
1:WA:92:LEU:N	1:WA:92:LEU:HD12	2.34	0.43
1:X:44:LEU:HD22	1:X:44:LEU:N	2.34	0.43
1:XA:52:THR:O	1:XA:109:ILE:HD11	2.19	0.43
1:XB:25:VAL:HG21	1:XB:78:THR:O	2.18	0.43
1:XB:176:LEU:HD23	1:XB:176:LEU:H	1.83	0.43
1:YA:208:VAL:HG23	1:YA:209:PHE:CD1	2.54	0.43
1:YB:398:VAL:HG12	1:YB:491:PRO:HB3	2.00	0.43
1:Z:359:ILE:O	1:Z:359:ILE:HG23	2.19	0.43
1:ZA:176:LEU:HD21	1:ZA:196:TRP:NE1	2.34	0.43
1:ZA:529:ILE:HD13	1:ZA:583:VAL:HG11	1.99	0.43
1:A:143:TRP:HH2	1:A:157:VAL:HG22	1.84	0.42
1:A:168:ILE:HG22	1:A:215:LEU:CD2	2.45	0.42
1:AB:176:LEU:HD21	1:AB:196:TRP:CE2	2.54	0.42
1:AB:219:VAL:HG11	1:AB:227:LEU:HD22	2.01	0.42
1:CB:44:LEU:HD22	1:CB:44:LEU:H	1.84	0.42
1:CB:176:LEU:HD21	1:CB:196:TRP:NE1	2.34	0.42
1:CB:426:LEU:HD23	1:CB:427:LEU:HD22	2.00	0.42
1:D:129:PHE:CE1	1:D:137:VAL:HG11	2.54	0.42
1:D:359:ILE:O	1:D:359:ILE:HG23	2.19	0.42
1:D:539:LEU:HD21	1:D:599:ILE:HD11	2.00	0.42
1:EA:44:LEU:HD22	1:EA:44:LEU:N	2.34	0.42
1:EB:570:ASP:OD1	1:EB:570:ASP:C	2.62	0.42
1:F:208:VAL:HG23	1:F:209:PHE:CD1	2.54	0.42
1:F:216:VAL:O	1:F:216:VAL:HG13	2.19	0.42
1:FA:216:VAL:O	1:FA:216:VAL:HG13	2.19	0.42
1:GB:44:LEU:N	1:GB:44:LEU:HD22	2.33	0.42
1:GB:176:LEU:HD21	1:GB:196:TRP:NE1	2.34	0.42
1:GB:523:PHE:CE2	1:GB:545:TRP:CD1	3.07	0.42
1:HA:44:LEU:N	1:HA:44:LEU:HD22	2.33	0.42
1:IA:17:HIS:HB3	1:IA:44:LEU:HD23	2.01	0.42
1:IA:176:LEU:HD21	1:IA:196:TRP:CE2	2.54	0.42
1:IB:539:LEU:CD1	1:IB:640:VAL:HG13	2.49	0.42
1:JA:138:VAL:HG12	1:JA:139:ALA:N	2.34	0.42
1:KA:176:LEU:HD21	1:KA:196:TRP:NE1	2.34	0.42
1:KB:121:LEU:HD23	1:KB:162:ILE:HG22	2.00	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:LA:487:VAL:O	1:LA:487:VAL:HG13	2.18	0.42
1:M:129:PHE:CE1	1:M:137:VAL:HG11	2.54	0.42
1:M:487:VAL:O	1:M:487:VAL:HG13	2.18	0.42
1:MA:219:VAL:HG11	1:MA:227:LEU:HD13	2.00	0.42
1:MB:44:LEU:HD22	1:MB:44:LEU:N	2.34	0.42
1:MB:138:VAL:HG12	1:MB:139:ALA:N	2.34	0.42
1:N:216:VAL:O	1:N:216:VAL:HG13	2.19	0.42
1:OA:779:LEU:HD21	1:PA:774:ARG:HG2	2.01	0.42
1:PA:539:LEU:HD21	1:PA:599:ILE:HD11	2.01	0.42
1:PB:44:LEU:HD22	1:PB:44:LEU:N	2.34	0.42
1:Q:48:MET:HA	1:Q:48:MET:HE3	2.00	0.42
1:R:17:HIS:HB3	1:R:44:LEU:HD23	2.01	0.42
1:S:208:VAL:HG23	1:S:209:PHE:CD1	2.54	0.42
1:S:523:PHE:CE2	1:S:545:TRP:CD1	3.07	0.42
1:TB:52:THR:O	1:TB:109:ILE:HD11	2.19	0.42
1:UA:17:HIS:CB	1:UA:44:LEU:HD23	2.49	0.42
1:VA:208:VAL:HG23	1:VA:209:PHE:CD1	2.54	0.42
1:VA:216:VAL:O	1:VA:216:VAL:HG13	2.19	0.42
1:VA:366:VAL:HG13	1:VA:366:VAL:O	2.19	0.42
1:VB:165:ALA:HB2	1:VB:205:LEU:HD13	2.00	0.42
1:VB:176:LEU:HD23	1:VB:176:LEU:H	1.83	0.42
1:VB:216:VAL:HG13	1:VB:216:VAL:O	2.19	0.42
1:WA:662:ILE:O	1:WA:666:THR:HG23	2.19	0.42
1:X:758:GLU:O	1:X:762:VAL:HG22	2.18	0.42
1:XB:44:LEU:HD22	1:XB:44:LEU:N	2.34	0.42
1:ZA:17:HIS:CB	1:ZA:44:LEU:HD23	2.49	0.42
1:ZA:570:ASP:C	1:ZA:570:ASP:OD1	2.62	0.42
1:A:489:LEU:HD21	1:A:495:PHE:CZ	2.54	0.42
1:AA:359:ILE:HG23	1:AA:359:ILE:O	2.18	0.42
1:AB:53:VAL:HG12	1:AB:57:HIS:O	2.18	0.42
1:AB:138:VAL:HG12	1:AB:139:ALA:N	2.35	0.42
1:AB:208:VAL:HG23	1:AB:209:PHE:CD1	2.54	0.42
1:AC:359:ILE:O	1:AC:359:ILE:HG23	2.19	0.42
1:BA:643:VAL:O	1:BA:643:VAL:HG12	2.19	0.42
1:BB:176:LEU:HD21	1:BB:196:TRP:CE2	2.54	0.42
1:BB:216:VAL:O	1:BB:216:VAL:HG13	2.19	0.42
1:CB:208:VAL:HG23	1:CB:209:PHE:CD1	2.54	0.42
1:CB:359:ILE:O	1:CB:359:ILE:HG23	2.19	0.42
1:CB:523:PHE:CE2	1:CB:545:TRP:CD1	3.07	0.42
1:D:168:ILE:HG22	1:D:215:LEU:CD2	2.44	0.42
1:D:208:VAL:HG23	1:D:209:PHE:CD1	2.54	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:DA:121:LEU:HD23	1:DA:162:ILE:HG22	2.00	0.42
1:DA:129:PHE:CE1	1:DA:137:VAL:HG11	2.54	0.42
1:E:216:VAL:O	1:E:216:VAL:HG13	2.19	0.42
1:EA:208:VAL:HG23	1:EA:209:PHE:CD1	2.54	0.42
1:EB:487:VAL:O	1:EB:487:VAL:HG13	2.19	0.42
1:G:138:VAL:HG12	1:G:139:ALA:N	2.34	0.42
1:G:176:LEU:HD21	1:G:196:TRP:NE1	2.34	0.42
1:G:541:LEU:CD2	1:G:640:VAL:HG22	2.49	0.42
1:GB:121:LEU:HD23	1:GB:162:ILE:HG22	2.00	0.42
1:GB:216:VAL:O	1:GB:216:VAL:HG13	2.19	0.42
1:GB:302:VAL:HG11	1:GB:306:LYS:HZ2	1.84	0.42
1:HB:487:VAL:O	1:HB:487:VAL:HG13	2.18	0.42
1:IA:53:VAL:HG12	1:IA:57:HIS:O	2.19	0.42
1:IA:176:LEU:HD21	1:IA:196:TRP:NE1	2.34	0.42
1:JA:208:VAL:HG23	1:JA:209:PHE:CD1	2.55	0.42
1:JB:138:VAL:HG12	1:JB:139:ALA:N	2.35	0.42
1:JB:176:LEU:HD23	1:JB:176:LEU:H	1.83	0.42
1:JB:208:VAL:HG23	1:JB:209:PHE:CD1	2.54	0.42
1:KB:53:VAL:HG12	1:KB:57:HIS:O	2.19	0.42
1:KB:138:VAL:HG12	1:KB:139:ALA:N	2.34	0.42
1:L:208:VAL:HG23	1:L:209:PHE:CD1	2.55	0.42
1:MA:208:VAL:HG23	1:MA:209:PHE:CD1	2.54	0.42
1:NB:138:VAL:HG12	1:NB:139:ALA:N	2.34	0.42
1:O:216:VAL:HG13	1:O:216:VAL:O	2.19	0.42
1:OA:208:VAL:HG23	1:OA:209:PHE:CD1	2.54	0.42
1:OA:489:LEU:HD21	1:OA:495:PHE:CZ	2.54	0.42
1:OB:138:VAL:HG12	1:OB:139:ALA:N	2.34	0.42
1:PA:63:ASN:OD1	1:PA:104:VAL:HG12	2.19	0.42
1:RB:359:ILE:O	1:RB:359:ILE:HG23	2.19	0.42
1:S:138:VAL:HG12	1:S:139:ALA:N	2.34	0.42
1:SA:56:ARG:N	1:SA:56:ARG:HE	2.17	0.42
1:SB:216:VAL:O	1:SB:216:VAL:HG13	2.19	0.42
1:TA:208:VAL:HG23	1:TA:209:PHE:CD1	2.54	0.42
1:UA:44:LEU:HD22	1:UA:44:LEU:N	2.34	0.42
1:UA:216:VAL:HG13	1:UA:216:VAL:O	2.19	0.42
1:V:176:LEU:HD23	1:V:176:LEU:H	1.82	0.42
1:WA:138:VAL:HG12	1:WA:139:ALA:N	2.34	0.42
1:WA:359:ILE:HG23	1:WA:359:ILE:O	2.19	0.42
1:WB:168:ILE:HG21	1:WB:174:LEU:HB2	2.00	0.42
1:WB:176:LEU:HD23	1:WB:176:LEU:H	1.83	0.42
1:XA:176:LEU:HD21	1:XA:196:TRP:CE2	2.54	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:XA:662:ILE:O	1:XA:666:THR:HG23	2.19	0.42
1:Z:176:LEU:HD21	1:Z:196:TRP:CE2	2.54	0.42
1:ZA:138:VAL:HG12	1:ZA:139:ALA:N	2.34	0.42
1:A:176:LEU:HD21	1:A:196:TRP:NE1	2.35	0.42
1:A:597:ARG:O	1:A:601:THR:HG23	2.19	0.42
1:BB:44:LEU:HD22	1:BB:44:LEU:N	2.34	0.42
1:BB:393:VAL:HG22	1:BB:411:ASP:C	2.44	0.42
1:C:529:ILE:HD13	1:C:583:VAL:HG21	2.00	0.42
1:CA:44:LEU:H	1:CA:44:LEU:HD22	1.84	0.42
1:DA:529:ILE:HD13	1:DA:583:VAL:HG11	2.01	0.42
1:DB:654:LEU:HD22	1:EB:534:HIS:CG	2.54	0.42
1:EA:216:VAL:HG13	1:EA:216:VAL:O	2.19	0.42
1:EB:176:LEU:HD21	1:EB:196:TRP:CE2	2.55	0.42
1:F:44:LEU:HD22	1:F:44:LEU:N	2.34	0.42
1:F:359:ILE:O	1:F:359:ILE:HG23	2.19	0.42
1:F:398:VAL:HG12	1:F:491:PRO:HB3	1.99	0.42
1:F:539:LEU:CD1	1:F:640:VAL:HG13	2.49	0.42
1:G:539:LEU:HD11	1:G:640:VAL:HG13	2.01	0.42
1:GA:393:VAL:HG22	1:GA:411:ASP:C	2.44	0.42
1:GA:597:ARG:O	1:GA:601:THR:HG23	2.18	0.42
1:H:487:VAL:HG13	1:H:487:VAL:O	2.18	0.42
1:HA:393:VAL:HG22	1:HA:411:ASP:C	2.43	0.42
1:HB:208:VAL:HG23	1:HB:209:PHE:CD1	2.54	0.42
1:I:597:ARG:O	1:I:601:THR:HG23	2.20	0.42
1:IB:129:PHE:CE1	1:IB:137:VAL:HG11	2.55	0.42
1:K:165:ALA:HB2	1:K:205:LEU:HD13	2.00	0.42
1:KA:216:VAL:O	1:KA:216:VAL:HG13	2.19	0.42
1:KB:208:VAL:HG23	1:KB:209:PHE:CD1	2.54	0.42
1:KB:523:PHE:CE2	1:KB:545:TRP:CD1	3.07	0.42
1:LA:359:ILE:HG23	1:LA:359:ILE:O	2.19	0.42
1:LB:523:PHE:CE2	1:LB:545:TRP:CD1	3.08	0.42
1:MA:326:LEU:HD12	1:MA:363:LEU:HD21	2.00	0.42
1:MB:176:LEU:HD21	1:MB:196:TRP:CE2	2.54	0.42
1:NA:176:LEU:HD21	1:NA:196:TRP:NE1	2.34	0.42
1:NB:176:LEU:HD21	1:NB:196:TRP:NE1	2.34	0.42
1:NB:216:VAL:HG13	1:NB:216:VAL:O	2.19	0.42
1:OB:359:ILE:O	1:OB:359:ILE:HG23	2.20	0.42
1:Q:426:LEU:HD23	1:Q:427:LEU:HD22	2.02	0.42
1:QA:208:VAL:HG23	1:QA:209:PHE:CD1	2.54	0.42
1:R:57:HIS:O	1:R:58:TYR:HD1	2.02	0.42
1:R:539:LEU:HD11	1:R:640:VAL:HG13	2.01	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:RA:539:LEU:CD1	1:RA:640:VAL:HG13	2.48	0.42
1:RA:779:LEU:HD21	1:SA:774:ARG:HG2	2.01	0.42
1:SB:208:VAL:HG23	1:SB:209:PHE:CD1	2.54	0.42
1:T:176:LEU:HD21	1:T:196:TRP:CE2	2.54	0.42
1:UA:541:LEU:CD2	1:UA:640:VAL:HG22	2.49	0.42
1:UB:345:ASP:OD2	1:UB:349:VAL:HG13	2.19	0.42
1:UB:523:PHE:CE2	1:UB:545:TRP:CD1	3.07	0.42
1:V:17:HIS:CB	1:V:44:LEU:HD23	2.49	0.42
1:VA:144:LEU:HD22	1:VA:163:ILE:HD11	2.02	0.42
1:VB:44:LEU:HD22	1:VB:44:LEU:N	2.34	0.42
1:VB:121:LEU:HD23	1:VB:162:ILE:HG22	2.01	0.42
1:VB:138:VAL:HG12	1:VB:139:ALA:N	2.34	0.42
1:W:359:ILE:O	1:W:359:ILE:HG23	2.19	0.42
1:WA:539:LEU:HD21	1:WA:599:ILE:HD11	2.01	0.42
1:X:168:ILE:HG22	1:X:215:LEU:CD2	2.45	0.42
1:X:176:LEU:HD21	1:X:196:TRP:NE1	2.33	0.42
1:XA:129:PHE:CE1	1:XA:137:VAL:HG11	2.54	0.42
1:Y:758:GLU:O	1:Y:762:VAL:HG22	2.18	0.42
1:YA:597:ARG:O	1:YA:601:THR:HG23	2.19	0.42
1:YA:758:GLU:O	1:YA:762:VAL:HG22	2.19	0.42
1:YB:539:LEU:CD2	1:YB:541:LEU:HG	2.49	0.42
1:ZB:57:HIS:O	1:ZB:58:TYR:HD1	2.02	0.42
1:AA:176:LEU:HD21	1:AA:196:TRP:NE1	2.34	0.42
1:AB:44:LEU:HD22	1:AB:44:LEU:N	2.34	0.42
1:AB:176:LEU:HD21	1:AB:196:TRP:NE1	2.34	0.42
1:B:30:VAL:HG21	1:B:50:MET:HE3	2.01	0.42
1:B:53:VAL:HG12	1:B:57:HIS:O	2.19	0.42
1:B:208:VAL:HG23	1:B:209:PHE:CD1	2.55	0.42
1:B:216:VAL:O	1:B:216:VAL:HG13	2.19	0.42
1:BA:176:LEU:HD21	1:BA:196:TRP:NE1	2.34	0.42
1:BA:208:VAL:HG23	1:BA:209:PHE:CD1	2.55	0.42
1:BA:359:ILE:O	1:BA:359:ILE:HG23	2.20	0.42
1:C:30:VAL:HG21	1:C:50:MET:HE3	2.01	0.42
1:C:53:VAL:HG12	1:C:57:HIS:O	2.19	0.42
1:CB:487:VAL:O	1:CB:487:VAL:HG13	2.19	0.42
1:DA:219:VAL:HG11	1:DA:227:LEU:HD13	2.01	0.42
1:DB:44:LEU:HD22	1:DB:44:LEU:H	1.84	0.42
1:DB:176:LEU:HD21	1:DB:196:TRP:NE1	2.35	0.42
1:DB:208:VAL:HG23	1:DB:209:PHE:CD1	2.55	0.42
1:E:316:LEU:HD12	1:E:316:LEU:HA	1.94	0.42
1:EB:758:GLU:O	1:EB:762:VAL:HG22	2.19	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:FB:208:VAL:HG23	1:FB:209:PHE:CD1	2.55	0.42
1:G:44:LEU:HD22	1:G:44:LEU:N	2.34	0.42
1:GB:138:VAL:HG12	1:GB:139:ALA:N	2.34	0.42
1:GB:219:VAL:HG11	1:GB:227:LEU:HD22	2.01	0.42
1:GB:359:ILE:HG23	1:GB:359:ILE:O	2.19	0.42
1:HA:129:PHE:CE1	1:HA:137:VAL:HG11	2.54	0.42
1:HB:138:VAL:HG12	1:HB:139:ALA:N	2.34	0.42
1:HB:219:VAL:HG11	1:HB:227:LEU:HD22	2.01	0.42
1:I:138:VAL:HG12	1:I:139:ALA:N	2.34	0.42
1:IA:129:PHE:CE1	1:IA:137:VAL:HG11	2.55	0.42
1:J:176:LEU:HD23	1:J:176:LEU:H	1.83	0.42
1:KB:44:LEU:HD22	1:KB:44:LEU:N	2.34	0.42
1:KB:129:PHE:CE1	1:KB:137:VAL:HG11	2.55	0.42
1:KB:165:ALA:HB2	1:KB:205:LEU:HD13	2.01	0.42
1:LB:138:VAL:HG12	1:LB:139:ALA:N	2.34	0.42
1:MA:44:LEU:HD22	1:MA:44:LEU:N	2.34	0.42
1:N:208:VAL:HG23	1:N:209:PHE:CD1	2.54	0.42
1:N:539:LEU:CD1	1:N:640:VAL:HG13	2.48	0.42
1:NA:144:LEU:HD22	1:NA:163:ILE:HD11	2.01	0.42
1:NA:393:VAL:HG22	1:NA:411:ASP:C	2.45	0.42
1:OA:597:ARG:O	1:OA:601:THR:HG23	2.20	0.42
1:P:208:VAL:HG23	1:P:209:PHE:CD1	2.54	0.42
1:PB:597:ARG:O	1:PB:601:THR:HG23	2.20	0.42
1:QA:144:LEU:HD12	1:QA:144:LEU:O	2.19	0.42
1:RA:129:PHE:CE1	1:RA:137:VAL:HG11	2.55	0.42
1:TA:779:LEU:HD21	1:UA:774:ARG:HG2	2.02	0.42
1:TB:44:LEU:HD22	1:TB:44:LEU:N	2.35	0.42
1:V:53:VAL:HG12	1:V:57:HIS:O	2.20	0.42
1:VB:10:ILE:HG22	1:VB:47:PRO:HB3	2.02	0.42
1:VB:129:PHE:CE1	1:VB:137:VAL:HG11	2.55	0.42
1:VB:523:PHE:CE2	1:VB:545:TRP:CD1	3.07	0.42
1:WA:44:LEU:HD22	1:WA:44:LEU:N	2.34	0.42
1:WA:176:LEU:HD21	1:WA:196:TRP:CE2	2.54	0.42
1:XA:121:LEU:HD23	1:XA:162:ILE:HG22	2.00	0.42
1:XB:138:VAL:HG12	1:XB:139:ALA:N	2.34	0.42
1:YA:176:LEU:HD21	1:YA:196:TRP:NE1	2.34	0.42
1:YB:17:HIS:HB3	1:YB:44:LEU:HD23	2.00	0.42
1:Z:17:HIS:CB	1:Z:44:LEU:HD23	2.50	0.42
1:ZA:208:VAL:HG23	1:ZA:209:PHE:CD1	2.54	0.42
1:ZB:176:LEU:HD21	1:ZB:196:TRP:CE2	2.55	0.42
1:A:138:VAL:HG12	1:A:139:ALA:N	2.34	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:AA:53:VAL:HG12	1:AA:57:HIS:O	2.20	0.42
1:AC:144:LEU:HD22	1:AC:163:ILE:HD11	2.02	0.42
1:AC:489:LEU:HD21	1:AC:495:PHE:CZ	2.55	0.42
1:B:176:LEU:HD21	1:B:196:TRP:CE2	2.55	0.42
1:BA:176:LEU:HD21	1:BA:196:TRP:CE2	2.55	0.42
1:C:144:LEU:HD12	1:C:144:LEU:O	2.19	0.42
1:CB:138:VAL:HG12	1:CB:139:ALA:N	2.34	0.42
1:E:176:LEU:HD21	1:E:196:TRP:CE2	2.54	0.42
1:EB:208:VAL:HG23	1:EB:209:PHE:CD1	2.54	0.42
1:F:643:VAL:O	1:F:643:VAL:HG12	2.20	0.42
1:FA:44:LEU:HD22	1:FA:44:LEU:N	2.34	0.42
1:I:393:VAL:HG22	1:I:411:ASP:C	2.44	0.42
1:IB:44:LEU:HD22	1:IB:44:LEU:H	1.85	0.42
1:J:662:ILE:O	1:J:666:THR:HG23	2.19	0.42
1:JB:216:VAL:O	1:JB:216:VAL:HG13	2.19	0.42
1:KA:44:LEU:HD22	1:KA:44:LEU:N	2.33	0.42
1:KA:208:VAL:HG23	1:KA:209:PHE:CD1	2.55	0.42
1:LA:121:LEU:HD23	1:LA:162:ILE:HG22	2.00	0.42
1:M:176:LEU:HD21	1:M:196:TRP:CE2	2.54	0.42
1:M:216:VAL:O	1:M:216:VAL:HG13	2.19	0.42
1:M:359:ILE:O	1:M:359:ILE:HG23	2.19	0.42
1:MA:44:LEU:HD22	1:MA:44:LEU:H	1.85	0.42
1:MA:129:PHE:CE1	1:MA:137:VAL:HG11	2.55	0.42
1:MB:597:ARG:O	1:MB:601:THR:HG23	2.20	0.42
1:N:393:VAL:HG22	1:N:411:ASP:C	2.44	0.42
1:NB:30:VAL:CG2	1:NB:50:MET:HE3	2.50	0.42
1:NB:570:ASP:OD1	1:NB:570:ASP:C	2.61	0.42
1:O:47:PRO:C	1:O:48:MET:HE2	2.45	0.42
1:O:523:PHE:CE2	1:O:545:TRP:CD1	3.08	0.42
1:OA:56:ARG:HE	1:PA:127:LEU:CD2	2.33	0.42
1:OA:539:LEU:CD1	1:OA:640:VAL:HG13	2.49	0.42
1:OB:56:ARG:HE	1:PB:127:LEU:CD2	2.33	0.42
1:OB:523:PHE:CE2	1:OB:545:TRP:CD1	3.07	0.42
1:P:17:HIS:CB	1:P:44:LEU:HD23	2.50	0.42
1:P:359:ILE:O	1:P:359:ILE:HG23	2.20	0.42
1:PA:138:VAL:HG12	1:PA:139:ALA:N	2.34	0.42
1:PB:144:LEU:HD22	1:PB:163:ILE:HD11	2.02	0.42
1:Q:208:VAL:HG23	1:Q:209:PHE:CD1	2.54	0.42
1:R:208:VAL:HG23	1:R:209:PHE:CD1	2.55	0.42
1:S:359:ILE:O	1:S:359:ILE:HG23	2.19	0.42
1:SA:176:LEU:HD21	1:SA:196:TRP:CE2	2.54	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:SB:138:VAL:HG12	1:SB:139:ALA:N	2.34	0.42
1:T:121:LEU:HD22	1:T:159:VAL:HG13	2.01	0.42
1:UA:129:PHE:CE1	1:UA:137:VAL:HG11	2.54	0.42
1:V:138:VAL:HG12	1:V:139:ALA:N	2.34	0.42
1:V:359:ILE:HG23	1:V:359:ILE:O	2.19	0.42
1:VA:138:VAL:HG12	1:VA:139:ALA:N	2.34	0.42
1:WB:44:LEU:N	1:WB:44:LEU:HD22	2.33	0.42
1:X:359:ILE:HG23	1:X:359:ILE:O	2.20	0.42
1:XA:17:HIS:CB	1:XA:44:LEU:HD23	2.49	0.42
1:XB:539:LEU:HD21	1:XB:599:ILE:HD11	2.01	0.42
1:Y:129:PHE:CE1	1:Y:137:VAL:HG11	2.55	0.42
1:Y:176:LEU:HD21	1:Y:196:TRP:CE2	2.55	0.42
1:YA:393:VAL:HG22	1:YA:411:ASP:C	2.45	0.42
1:Z:393:VAL:HG22	1:Z:411:ASP:C	2.45	0.42
1:AA:541:LEU:CD2	1:AA:640:VAL:HG22	2.50	0.42
1:AB:121:LEU:HD23	1:AB:162:ILE:HG22	2.00	0.42
1:BA:44:LEU:HD22	1:BA:44:LEU:N	2.35	0.42
1:CA:17:HIS:CB	1:CA:44:LEU:HD23	2.50	0.42
1:CB:121:LEU:HD23	1:CB:162:ILE:HG22	2.00	0.42
1:DB:56:ARG:HE	1:EB:127:LEU:CD2	2.33	0.42
1:E:208:VAL:HG23	1:E:209:PHE:CD1	2.54	0.42
1:E:359:ILE:HG23	1:E:359:ILE:O	2.20	0.42
1:EA:138:VAL:HG12	1:EA:139:ALA:N	2.35	0.42
1:EA:359:ILE:HG23	1:EA:359:ILE:O	2.19	0.42
1:EB:44:LEU:H	1:EB:44:LEU:HD22	1.84	0.42
1:EB:168:ILE:HG21	1:EB:174:LEU:HB2	2.01	0.42
1:F:487:VAL:O	1:F:487:VAL:HG13	2.18	0.42
1:FA:208:VAL:HG23	1:FA:209:PHE:CD1	2.55	0.42
1:G:359:ILE:O	1:G:359:ILE:HG23	2.20	0.42
1:GA:176:LEU:HD21	1:GA:196:TRP:NE1	2.35	0.42
1:H:144:LEU:HD22	1:H:163:ILE:HD11	2.02	0.42
1:HA:53:VAL:HG12	1:HA:57:HIS:O	2.20	0.42
1:I:165:ALA:HB2	1:I:205:LEU:HD13	2.00	0.42
1:IB:44:LEU:HD22	1:IB:44:LEU:N	2.34	0.42
1:JA:359:ILE:O	1:JA:359:ILE:HG23	2.19	0.42
1:JB:165:ALA:HB2	1:JB:205:LEU:HD13	2.02	0.42
1:JB:359:ILE:HG23	1:JB:359:ILE:O	2.19	0.42
1:JB:487:VAL:O	1:JB:487:VAL:HG13	2.18	0.42
1:JB:539:LEU:HD21	1:JB:599:ILE:HD11	2.02	0.42
1:KA:359:ILE:HG23	1:KA:359:ILE:O	2.19	0.42
1:LA:219:VAL:HG11	1:LA:227:LEU:HD13	2.02	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:LB:176:LEU:HD21	1:LB:196:TRP:CE2	2.55	0.42
1:M:176:LEU:HD23	1:M:176:LEU:H	1.83	0.42
1:MA:788:ALA:O	1:MA:792:VAL:HG23	2.19	0.42
1:MB:539:LEU:HD11	1:MB:640:VAL:HG13	2.02	0.42
1:N:53:VAL:HG12	1:N:57:HIS:O	2.19	0.42
1:NA:359:ILE:HG23	1:NA:359:ILE:O	2.19	0.42
1:NB:53:VAL:HG12	1:NB:57:HIS:O	2.19	0.42
1:O:176:LEU:HD21	1:O:196:TRP:CE2	2.54	0.42
1:O:597:ARG:O	1:O:601:THR:HG23	2.20	0.42
1:PA:208:VAL:HG23	1:PA:209:PHE:CD1	2.54	0.42
1:PA:487:VAL:HG13	1:PA:487:VAL:O	2.19	0.42
1:PA:643:VAL:HG12	1:PA:643:VAL:O	2.20	0.42
1:QA:597:ARG:O	1:QA:601:THR:HG23	2.19	0.42
1:QB:17:HIS:HB3	1:QB:44:LEU:HD23	2.00	0.42
1:R:143:TRP:HH2	1:R:157:VAL:HG22	1.84	0.42
1:R:219:VAL:HG11	1:R:227:LEU:HD22	2.02	0.42
1:RA:57:HIS:O	1:RA:58:TYR:HD1	2.03	0.42
1:RB:216:VAL:O	1:RB:216:VAL:HG13	2.19	0.42
1:T:57:HIS:O	1:T:58:TYR:HD1	2.03	0.42
1:T:539:LEU:HD21	1:T:599:ILE:HD11	2.02	0.42
1:TA:359:ILE:O	1:TA:359:ILE:HG23	2.20	0.42
1:TA:539:LEU:CD1	1:TA:640:VAL:HG13	2.49	0.42
1:TB:176:LEU:HD23	1:TB:176:LEU:H	1.83	0.42
1:UA:138:VAL:HG12	1:UA:139:ALA:N	2.34	0.42
1:UA:208:VAL:HG23	1:UA:209:PHE:CD1	2.55	0.42
1:UA:489:LEU:HD21	1:UA:495:PHE:CZ	2.55	0.42
1:VA:788:ALA:O	1:VA:792:VAL:HG23	2.19	0.42
1:VB:176:LEU:HD21	1:VB:196:TRP:CE2	2.55	0.42
1:VB:345:ASP:OD2	1:VB:349:VAL:HG13	2.19	0.42
1:W:176:LEU:HD21	1:W:196:TRP:CE2	2.54	0.42
1:W:487:VAL:O	1:W:487:VAL:HG13	2.18	0.42
1:W:597:ARG:O	1:W:601:THR:HG23	2.19	0.42
1:WB:208:VAL:HG23	1:WB:209:PHE:CD1	2.55	0.42
1:WB:359:ILE:O	1:WB:359:ILE:HG23	2.19	0.42
1:X:523:PHE:CE2	1:X:545:TRP:CD1	3.08	0.42
1:X:539:LEU:CD1	1:X:599:ILE:HD11	2.50	0.42
1:XA:44:LEU:N	1:XA:44:LEU:HD22	2.35	0.42
1:XA:302:VAL:HG11	1:XA:306:LYS:HZ2	1.85	0.42
1:XA:539:LEU:HD21	1:XA:599:ILE:HD11	2.01	0.42
1:XB:779:LEU:HD21	1:YB:774:ARG:HG2	2.01	0.42
1:YA:56:ARG:HE	1:ZA:127:LEU:CD2	2.33	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:YB:44:LEU:HD22	1:YB:44:LEU:N	2.35	0.42
1:ZB:138:VAL:HG12	1:ZB:139:ALA:N	2.34	0.42
1:A:219:VAL:HG11	1:A:227:LEU:HD13	2.01	0.42
1:AA:208:VAL:HG23	1:AA:209:PHE:CD1	2.54	0.42
1:AB:56:ARG:HE	1:BB:127:LEU:CD2	2.33	0.42
1:AC:216:VAL:O	1:AC:216:VAL:HG13	2.19	0.42
1:BB:44:LEU:HD22	1:BB:44:LEU:H	1.84	0.42
1:C:597:ARG:O	1:C:601:THR:HG23	2.19	0.42
1:CB:56:ARG:HE	1:DB:127:LEU:CD2	2.33	0.42
1:DA:176:LEU:HD21	1:DA:196:TRP:NE1	2.35	0.42
1:E:138:VAL:HG12	1:E:139:ALA:N	2.34	0.42
1:EB:539:LEU:HD11	1:EB:640:VAL:HG13	2.02	0.42
1:FA:176:LEU:HD21	1:FA:196:TRP:NE1	2.35	0.42
1:G:208:VAL:HG23	1:G:209:PHE:CD1	2.55	0.42
1:GA:57:HIS:O	1:GA:58:TYR:HD1	2.03	0.42
1:GA:138:VAL:HG12	1:GA:139:ALA:N	2.34	0.42
1:HA:316:LEU:HD12	1:HA:316:LEU:HA	1.94	0.42
1:HB:643:VAL:O	1:HB:643:VAL:HG12	2.20	0.42
1:I:487:VAL:O	1:I:487:VAL:HG13	2.19	0.42
1:IA:487:VAL:O	1:IA:487:VAL:HG13	2.20	0.42
1:IB:216:VAL:HG13	1:IB:216:VAL:O	2.19	0.42
1:J:176:LEU:HD21	1:J:196:TRP:CE2	2.55	0.42
1:J:216:VAL:HG13	1:J:216:VAL:O	2.19	0.42
1:JA:176:LEU:HD21	1:JA:196:TRP:NE1	2.35	0.42
1:JB:176:LEU:HD21	1:JB:196:TRP:CE2	2.55	0.42
1:JB:597:ARG:O	1:JB:601:THR:HG23	2.20	0.42
1:K:216:VAL:O	1:K:216:VAL:HG13	2.19	0.42
1:KB:63:ASN:OD1	1:KB:104:VAL:HG12	2.19	0.42
1:KB:539:LEU:HD21	1:KB:599:ILE:HD11	2.01	0.42
1:L:597:ARG:O	1:L:601:THR:HG23	2.20	0.42
1:LA:57:HIS:O	1:LA:58:TYR:HD1	2.02	0.42
1:M:176:LEU:HD21	1:M:196:TRP:NE1	2.34	0.42
1:MA:529:ILE:HD13	1:MA:583:VAL:HG11	2.02	0.42
1:NA:44:LEU:HD22	1:NA:44:LEU:N	2.33	0.42
1:NA:489:LEU:HD21	1:NA:495:PHE:CZ	2.55	0.42
1:NA:529:ILE:HD13	1:NA:583:VAL:HG21	2.02	0.42
1:NB:539:LEU:HD21	1:NB:599:ILE:HD11	2.01	0.42
1:OB:176:LEU:HD21	1:OB:196:TRP:NE1	2.34	0.42
1:P:44:LEU:HD22	1:P:44:LEU:H	1.84	0.42
1:PA:662:ILE:O	1:PA:666:THR:HG23	2.19	0.42
1:PA:758:GLU:O	1:PA:762:VAL:HG22	2.19	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:PB:208:VAL:HG23	1:PB:209:PHE:CD1	2.55	0.42
1:PB:281:TYR:CG	1:PB:366:VAL:HG23	2.55	0.42
1:Q:539:LEU:HD21	1:Q:599:ILE:HD11	2.02	0.42
1:Q:597:ARG:O	1:Q:601:THR:HG23	2.20	0.42
1:QB:219:VAL:HG11	1:QB:227:LEU:HD13	2.01	0.42
1:RA:208:VAL:HG23	1:RA:209:PHE:CD1	2.55	0.42
1:RB:52:THR:O	1:RB:109:ILE:HD11	2.19	0.42
1:RB:57:HIS:O	1:RB:58:TYR:HD1	2.02	0.42
1:S:219:VAL:HG11	1:S:227:LEU:HD13	2.00	0.42
1:SA:208:VAL:HG23	1:SA:209:PHE:CD1	2.54	0.42
1:T:208:VAL:HG23	1:T:209:PHE:CD1	2.54	0.42
1:TB:393:VAL:HG22	1:TB:411:ASP:C	2.44	0.42
1:WA:129:PHE:CE1	1:WA:137:VAL:HG11	2.55	0.42
1:WA:597:ARG:O	1:WA:601:THR:HG23	2.20	0.42
1:WB:44:LEU:HD22	1:WB:44:LEU:H	1.84	0.42
1:WB:129:PHE:CE1	1:WB:137:VAL:HG11	2.55	0.42
1:Y:44:LEU:HD22	1:Y:44:LEU:N	2.35	0.42
1:YA:165:ALA:HB2	1:YA:205:LEU:HD13	2.00	0.42
1:Z:44:LEU:HD22	1:Z:44:LEU:N	2.34	0.42
1:Z:138:VAL:HG12	1:Z:139:ALA:N	2.34	0.42
1:Z:208:VAL:HG23	1:Z:209:PHE:CD1	2.54	0.42
1:ZB:208:VAL:HG23	1:ZB:209:PHE:CD1	2.54	0.42
1:AC:138:VAL:HG12	1:AC:139:ALA:N	2.34	0.42
1:AC:393:VAL:HG22	1:AC:411:ASP:C	2.44	0.42
1:B:487:VAL:O	1:B:487:VAL:HG13	2.20	0.42
1:BB:208:VAL:HG23	1:BB:209:PHE:CD1	2.54	0.42
1:C:176:LEU:HD21	1:C:196:TRP:CE2	2.55	0.42
1:CA:44:LEU:HD22	1:CA:44:LEU:N	2.34	0.42
1:CA:48:MET:HA	1:CA:48:MET:HE3	2.02	0.42
1:CA:208:VAL:HG23	1:CA:209:PHE:CD1	2.54	0.42
1:D:17:HIS:HB3	1:D:44:LEU:HD23	2.01	0.42
1:EB:138:VAL:HG12	1:EB:139:ALA:N	2.35	0.42
1:FA:138:VAL:HG12	1:FA:139:ALA:N	2.34	0.42
1:FA:219:VAL:HG11	1:FA:227:LEU:HD13	2.00	0.42
1:GB:487:VAL:O	1:GB:487:VAL:HG13	2.20	0.42
1:J:779:LEU:HD21	1:K:774:ARG:HG2	2.02	0.42
1:JA:129:PHE:CE1	1:JA:137:VAL:HG11	2.54	0.42
1:JB:758:GLU:O	1:JB:762:VAL:HG22	2.20	0.42
1:K:487:VAL:O	1:K:487:VAL:HG13	2.19	0.42
1:LA:523:PHE:CE2	1:LA:545:TRP:CD1	3.08	0.42
1:LB:359:ILE:HG23	1:LB:359:ILE:O	2.19	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:N:489:LEU:HD21	1:N:495:PHE:CZ	2.55	0.42
1:N:643:VAL:O	1:N:643:VAL:HG12	2.20	0.42
1:NA:208:VAL:HG23	1:NA:209:PHE:CD1	2.54	0.42
1:NB:176:LEU:HD21	1:NB:196:TRP:CE2	2.54	0.42
1:O:758:GLU:O	1:O:762:VAL:HG22	2.20	0.42
1:PA:359:ILE:HG23	1:PA:359:ILE:O	2.20	0.42
1:Q:529:ILE:HD13	1:Q:583:VAL:HG11	2.02	0.42
1:QA:53:VAL:HG12	1:QA:57:HIS:O	2.19	0.42
1:QA:176:LEU:HD21	1:QA:196:TRP:CE2	2.55	0.42
1:QA:216:VAL:O	1:QA:216:VAL:HG13	2.19	0.42
1:QB:334:LEU:HD12	1:QB:357:TRP:NE1	2.34	0.42
1:R:216:VAL:HG13	1:R:216:VAL:O	2.18	0.42
1:R:487:VAL:O	1:R:487:VAL:HG13	2.19	0.42
1:RA:138:VAL:HG12	1:RA:139:ALA:N	2.34	0.42
1:S:129:PHE:CE1	1:S:137:VAL:HG11	2.55	0.42
1:SB:176:LEU:HD21	1:SB:196:TRP:CE2	2.55	0.42
1:T:56:ARG:HE	1:V:127:LEU:CD2	2.33	0.42
1:T:393:VAL:HG22	1:T:411:ASP:C	2.45	0.42
1:TB:57:HIS:O	1:TB:58:TYR:HD1	2.03	0.42
1:TB:138:VAL:HG12	1:TB:139:ALA:N	2.34	0.42
1:UA:779:LEU:HD21	1:VA:774:ARG:HG2	2.02	0.42
1:VA:359:ILE:HG23	1:VA:359:ILE:O	2.20	0.42
1:WA:56:ARG:HE	1:XA:127:LEU:CD2	2.33	0.42
1:WB:17:HIS:HB3	1:WB:44:LEU:HD23	2.02	0.42
1:X:208:VAL:HG23	1:X:209:PHE:CD1	2.54	0.42
1:X:597:ARG:O	1:X:601:THR:HG23	2.20	0.42
1:XA:57:HIS:O	1:XA:58:TYR:HD1	2.03	0.42
1:XB:316:LEU:HD12	1:XB:316:LEU:HA	1.94	0.42
1:YA:779:LEU:HD21	1:ZA:774:ARG:HG2	2.01	0.42
1:YB:3:THR:HG22	1:YB:6:PHE:HA	2.02	0.42
1:YB:138:VAL:HG12	1:YB:139:ALA:N	2.34	0.42
1:ZB:52:THR:O	1:ZB:109:ILE:HD11	2.19	0.42
1:AB:216:VAL:O	1:AB:216:VAL:HG13	2.19	0.42
1:AC:127:LEU:CD2	1:ZB:56:ARG:HE	2.33	0.42
1:AC:219:VAL:HG11	1:AC:227:LEU:HD22	2.01	0.42
1:C:208:VAL:HG23	1:C:209:PHE:CD1	2.54	0.42
1:CA:176:LEU:HD21	1:CA:196:TRP:CE2	2.55	0.42
1:CB:643:VAL:O	1:CB:643:VAL:HG12	2.19	0.42
1:EA:176:LEU:HD21	1:EA:196:TRP:NE1	2.35	0.42
1:F:57:HIS:O	1:F:58:TYR:HD1	2.03	0.42
1:F:779:LEU:HD21	1:G:774:ARG:HG2	2.01	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:G:53:VAL:HG12	1:G:57:HIS:O	2.19	0.42
1:G:489:LEU:HD21	1:G:495:PHE:CZ	2.55	0.42
1:HA:359:ILE:O	1:HA:359:ILE:HG23	2.20	0.42
1:HB:73:VAL:HG11	1:HB:81:VAL:HG23	2.01	0.42
1:IB:138:VAL:HG12	1:IB:139:ALA:N	2.35	0.42
1:IB:159:VAL:O	1:IB:159:VAL:HG12	2.20	0.42
1:J:541:LEU:CD2	1:J:640:VAL:HG22	2.50	0.42
1:KA:393:VAL:HG22	1:KA:411:ASP:C	2.45	0.42
1:KA:539:LEU:HD21	1:KA:599:ILE:HD11	2.01	0.42
1:LA:3:THR:HG22	1:LA:6:PHE:HA	2.02	0.42
1:LA:176:LEU:HD21	1:LA:196:TRP:NE1	2.35	0.42
1:MA:176:LEU:HD21	1:MA:196:TRP:CE2	2.55	0.42
1:MA:393:VAL:HG22	1:MA:411:ASP:C	2.45	0.42
1:MB:44:LEU:HD22	1:MB:44:LEU:H	1.84	0.42
1:NA:57:HIS:O	1:NA:58:TYR:HD1	2.03	0.42
1:NA:176:LEU:HD21	1:NA:196:TRP:CE2	2.54	0.42
1:O:56:ARG:HE	1:P:127:LEU:CD2	2.33	0.42
1:OA:143:TRP:HH2	1:OA:157:VAL:HG22	1.84	0.42
1:OB:208:VAL:HG23	1:OB:209:PHE:CD1	2.55	0.42
1:OB:393:VAL:HG22	1:OB:411:ASP:C	2.45	0.42
1:P:129:PHE:CE1	1:P:137:VAL:HG11	2.55	0.42
1:PB:643:VAL:O	1:PB:643:VAL:HG12	2.20	0.42
1:QB:138:VAL:HG12	1:QB:139:ALA:N	2.34	0.42
1:QB:529:ILE:HD13	1:QB:583:VAL:HG11	2.02	0.42
1:R:52:THR:O	1:R:109:ILE:HD11	2.20	0.42
1:TA:57:HIS:O	1:TA:58:TYR:HD1	2.03	0.42
1:TA:516:LEU:HD23	1:TA:516:LEU:C	2.44	0.42
1:TB:44:LEU:HD22	1:TB:44:LEU:H	1.84	0.42
1:UA:662:ILE:O	1:UA:666:THR:HG23	2.19	0.42
1:UB:208:VAL:HG23	1:UB:209:PHE:CD1	2.54	0.42
1:UB:219:VAL:HG11	1:UB:227:LEU:HD13	2.02	0.42
1:VA:129:PHE:CE1	1:VA:137:VAL:HG11	2.55	0.42
1:VB:44:LEU:HD22	1:VB:44:LEU:H	1.84	0.42
1:VB:53:VAL:HG12	1:VB:57:HIS:O	2.19	0.42
1:VB:393:VAL:HG22	1:VB:411:ASP:C	2.45	0.42
1:W:216:VAL:O	1:W:216:VAL:HG13	2.19	0.42
1:WB:57:HIS:O	1:WB:58:TYR:HD1	2.03	0.42
1:X:129:PHE:CE1	1:X:137:VAL:HG11	2.54	0.42
1:XA:489:LEU:HD21	1:XA:495:PHE:CZ	2.55	0.42
1:XB:57:HIS:O	1:XB:58:TYR:HD1	2.03	0.42
1:Y:523:PHE:CE2	1:Y:545:TRP:CD1	3.08	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:YA:121:LEU:HD23	1:YA:162:ILE:HG22	2.00	0.42
1:YA:219:VAL:HG11	1:YA:227:LEU:HD22	2.01	0.42
1:YB:144:LEU:HD22	1:YB:163:ILE:HD11	2.02	0.42
1:ZB:216:VAL:HG13	1:ZB:216:VAL:O	2.19	0.42
1:AA:56:ARG:HE	1:BA:127:LEU:HD21	1.85	0.42
1:AA:127:LEU:CD2	1:Z:56:ARG:HE	2.33	0.42
1:AC:176:LEU:HD21	1:AC:196:TRP:NE1	2.34	0.42
1:B:597:ARG:O	1:B:601:THR:HG23	2.20	0.42
1:BA:17:HIS:CB	1:BA:44:LEU:HD23	2.50	0.42
1:CA:539:LEU:CD1	1:CA:640:VAL:HG13	2.50	0.42
1:G:654:LEU:HD22	1:H:534:HIS:CG	2.55	0.42
1:GA:758:GLU:O	1:GA:762:VAL:HG22	2.18	0.42
1:HA:538:GLN:HB2	1:HA:646:VAL:HG22	2.01	0.42
1:HA:570:ASP:OD1	1:HA:570:ASP:C	2.62	0.42
1:HA:662:ILE:O	1:HA:666:THR:HG23	2.20	0.42
1:HB:216:VAL:O	1:HB:216:VAL:HG13	2.19	0.42
1:HB:393:VAL:HG22	1:HB:411:ASP:C	2.44	0.42
1:J:144:LEU:HD12	1:J:163:ILE:HD11	2.02	0.42
1:J:539:LEU:HD11	1:J:640:VAL:HG13	2.01	0.42
1:K:176:LEU:HD21	1:K:196:TRP:NE1	2.34	0.42
1:K:779:LEU:HD21	1:L:774:ARG:HG2	2.01	0.42
1:KA:583:VAL:O	1:KA:583:VAL:HG12	2.19	0.42
1:L:138:VAL:HG12	1:L:139:ALA:N	2.34	0.42
1:MA:57:HIS:O	1:MA:58:TYR:HD1	2.02	0.42
1:N:63:ASN:OD1	1:N:104:VAL:HG12	2.20	0.42
1:NB:281:TYR:CG	1:NB:366:VAL:HG23	2.55	0.42
1:NB:643:VAL:O	1:NB:643:VAL:HG12	2.20	0.42
1:O:413:VAL:HG22	1:O:414:LEU:N	2.35	0.42
1:OA:53:VAL:HG12	1:OA:57:HIS:O	2.20	0.42
1:OA:216:VAL:O	1:OA:216:VAL:HG13	2.19	0.42
1:OB:129:PHE:CE1	1:OB:137:VAL:HG11	2.55	0.42
1:OB:643:VAL:HG12	1:OB:643:VAL:O	2.19	0.42
1:PA:52:THR:O	1:PA:109:ILE:HD11	2.20	0.42
1:PA:570:ASP:OD1	1:PA:570:ASP:C	2.62	0.42
1:PB:216:VAL:O	1:PB:216:VAL:HG13	2.19	0.42
1:S:143:TRP:HH2	1:S:157:VAL:HG22	1.85	0.42
1:SA:539:LEU:HD21	1:SA:599:ILE:HD11	2.02	0.42
1:TA:643:VAL:O	1:TA:643:VAL:HG12	2.20	0.42
1:UA:523:PHE:CE2	1:UA:545:TRP:CD1	3.07	0.42
1:WA:165:ALA:HB2	1:WA:205:LEU:HD13	2.00	0.42
1:WB:216:VAL:O	1:WB:216:VAL:HG13	2.19	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:XB:393:VAL:HG22	1:XB:411:ASP:C	2.44	0.42
1:Z:516:LEU:HD23	1:Z:516:LEU:O	2.20	0.42
1:A:53:VAL:HG12	1:A:57:HIS:O	2.19	0.41
1:AA:393:VAL:HG22	1:AA:411:ASP:C	2.45	0.41
1:AB:129:PHE:CE1	1:AB:137:VAL:HG11	2.55	0.41
1:AB:359:ILE:O	1:AB:359:ILE:HG23	2.19	0.41
1:B:138:VAL:HG12	1:B:139:ALA:N	2.35	0.41
1:B:326:LEU:HD12	1:B:363:LEU:HD21	2.01	0.41
1:BA:129:PHE:CE1	1:BA:137:VAL:HG11	2.55	0.41
1:BB:159:VAL:O	1:BB:159:VAL:HG12	2.19	0.41
1:C:216:VAL:O	1:C:216:VAL:HG13	2.19	0.41
1:CA:281:TYR:CG	1:CA:366:VAL:HG23	2.55	0.41
1:D:737:GLY:O	1:D:741:VAL:HG23	2.20	0.41
1:D:779:LEU:HD21	1:E:774:ARG:HG2	2.01	0.41
1:DA:662:ILE:O	1:DA:666:THR:HG23	2.19	0.41
1:DB:643:VAL:HG12	1:DB:643:VAL:O	2.20	0.41
1:F:538:GLN:HB2	1:F:646:VAL:HG22	2.02	0.41
1:FB:138:VAL:HG12	1:FB:139:ALA:N	2.35	0.41
1:GA:44:LEU:HD22	1:GA:44:LEU:H	1.85	0.41
1:GA:219:VAL:HG11	1:GA:227:LEU:HD22	2.02	0.41
1:GA:779:LEU:HD21	1:HA:774:ARG:HG2	2.02	0.41
1:H:359:ILE:O	1:H:359:ILE:HG23	2.20	0.41
1:HB:176:LEU:HD21	1:HB:196:TRP:NE1	2.35	0.41
1:J:487:VAL:HG13	1:J:487:VAL:O	2.18	0.41
1:JB:662:ILE:O	1:JB:666:THR:HG23	2.20	0.41
1:K:17:HIS:CB	1:K:44:LEU:HD23	2.50	0.41
1:L:17:HIS:CB	1:L:44:LEU:HD23	2.50	0.41
1:LB:129:PHE:CE1	1:LB:137:VAL:HG11	2.55	0.41
1:M:138:VAL:HG12	1:M:139:ALA:N	2.34	0.41
1:N:570:ASP:OD1	1:N:570:ASP:C	2.62	0.41
1:NB:17:HIS:HB3	1:NB:44:LEU:HD23	2.01	0.41
1:O:138:VAL:HG12	1:O:139:ALA:N	2.34	0.41
1:OA:539:LEU:HD21	1:OA:599:ILE:HD11	2.02	0.41
1:P:57:HIS:O	1:P:58:TYR:HD1	2.03	0.41
1:PB:129:PHE:CE1	1:PB:137:VAL:HG11	2.55	0.41
1:PB:138:VAL:HG12	1:PB:139:ALA:N	2.34	0.41
1:QA:359:ILE:O	1:QA:359:ILE:HG23	2.19	0.41
1:R:129:PHE:CE1	1:R:137:VAL:HG11	2.54	0.41
1:RA:30:VAL:CG2	1:RA:50:MET:HE3	2.50	0.41
1:RA:95:ASP:O	1:RA:96:PRO:C	2.63	0.41
1:RB:208:VAL:HG23	1:RB:209:PHE:CD1	2.55	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:S:57:HIS:O	1:S:58:TYR:HD1	2.03	0.41
1:S:216:VAL:O	1:S:216:VAL:HG13	2.19	0.41
1:S:393:VAL:HG22	1:S:411:ASP:C	2.45	0.41
1:SA:44:LEU:HD22	1:SA:44:LEU:N	2.35	0.41
1:SA:95:ASP:O	1:SA:96:PRO:C	2.62	0.41
1:SA:393:VAL:HG22	1:SA:411:ASP:C	2.45	0.41
1:SA:654:LEU:HD22	1:TA:534:HIS:CG	2.55	0.41
1:SB:57:HIS:O	1:SB:58:TYR:HD1	2.03	0.41
1:SB:597:ARG:O	1:SB:601:THR:HG23	2.20	0.41
1:T:73:VAL:HG11	1:T:81:VAL:HG23	2.02	0.41
1:T:138:VAL:HG12	1:T:139:ALA:N	2.34	0.41
1:TB:779:LEU:HD21	1:UB:774:ARG:HG2	2.01	0.41
1:UA:359:ILE:HG23	1:UA:359:ILE:O	2.20	0.41
1:UB:44:LEU:HD22	1:UB:44:LEU:H	1.84	0.41
1:W:138:VAL:HG12	1:W:139:ALA:N	2.34	0.41
1:X:138:VAL:HG12	1:X:139:ALA:N	2.34	0.41
1:XA:144:LEU:HD12	1:XA:163:ILE:HD11	2.01	0.41
1:XB:143:TRP:HH2	1:XB:157:VAL:HG22	1.85	0.41
1:Y:216:VAL:O	1:Y:216:VAL:HG13	2.19	0.41
1:YB:643:VAL:O	1:YB:643:VAL:HG12	2.19	0.41
1:ZA:44:LEU:HD22	1:ZA:44:LEU:N	2.35	0.41
1:A:44:LEU:HD22	1:A:44:LEU:N	2.34	0.41
1:A:56:ARG:HE	1:B:127:LEU:CD2	2.33	0.41
1:A:129:PHE:CE1	1:A:137:VAL:HG11	2.55	0.41
1:A:779:LEU:HD21	1:B:774:ARG:HG2	2.01	0.41
1:AA:48:MET:HE2	1:AA:48:MET:HA	2.02	0.41
1:AB:144:LEU:HD22	1:AB:163:ILE:HD11	2.02	0.41
1:AC:208:VAL:HG23	1:AC:209:PHE:CD1	2.55	0.41
1:AC:541:LEU:CD2	1:AC:640:VAL:HG22	2.51	0.41
1:BA:662:ILE:O	1:BA:666:THR:HG23	2.20	0.41
1:BB:779:LEU:HD21	1:CB:774:ARG:HG2	2.01	0.41
1:C:359:ILE:O	1:C:359:ILE:HG23	2.20	0.41
1:CA:57:HIS:O	1:CA:58:TYR:HD1	2.03	0.41
1:CA:216:VAL:O	1:CA:216:VAL:HG13	2.19	0.41
1:CB:44:LEU:HD22	1:CB:44:LEU:N	2.33	0.41
1:CB:176:LEU:HD21	1:CB:196:TRP:CE2	2.55	0.41
1:CB:219:VAL:HG11	1:CB:227:LEU:HD22	2.01	0.41
1:CB:597:ARG:O	1:CB:601:THR:HG23	2.20	0.41
1:D:44:LEU:HD22	1:D:44:LEU:N	2.34	0.41
1:DB:129:PHE:CE1	1:DB:137:VAL:HG11	2.55	0.41
1:E:129:PHE:CE1	1:E:137:VAL:HG11	2.54	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:EA:57:HIS:O	1:EA:58:TYR:HD1	2.02	0.41
1:EB:176:LEU:HD21	1:EB:196:TRP:NE1	2.35	0.41
1:EB:426:LEU:HD23	1:EB:427:LEU:HD22	2.01	0.41
1:F:138:VAL:HG12	1:F:139:ALA:N	2.34	0.41
1:F:176:LEU:HD21	1:F:196:TRP:CE2	2.55	0.41
1:FA:129:PHE:CE1	1:FA:137:VAL:HG11	2.54	0.41
1:FA:176:LEU:HD21	1:FA:196:TRP:CE2	2.55	0.41
1:FB:129:PHE:CE1	1:FB:137:VAL:HG11	2.54	0.41
1:H:138:VAL:HG12	1:H:139:ALA:N	2.35	0.41
1:IB:662:ILE:O	1:IB:666:THR:HG23	2.21	0.41
1:J:489:LEU:HD21	1:J:495:PHE:CZ	2.56	0.41
1:KA:17:HIS:CB	1:KA:44:LEU:HD23	2.51	0.41
1:KA:176:LEU:HD21	1:KA:196:TRP:CE2	2.55	0.41
1:LB:219:VAL:HG11	1:LB:227:LEU:HD22	2.02	0.41
1:LB:539:LEU:CD1	1:LB:640:VAL:HG13	2.50	0.41
1:M:489:LEU:HD21	1:M:495:PHE:CZ	2.56	0.41
1:MB:176:LEU:HD21	1:MB:196:TRP:NE1	2.35	0.41
1:MB:737:GLY:O	1:MB:741:VAL:HG23	2.20	0.41
1:N:176:LEU:HD21	1:N:196:TRP:CE2	2.55	0.41
1:N:359:ILE:O	1:N:359:ILE:HG23	2.20	0.41
1:O:176:LEU:HD21	1:O:196:TRP:NE1	2.34	0.41
1:O:359:ILE:O	1:O:359:ILE:HG23	2.19	0.41
1:OB:539:LEU:HD21	1:OB:599:ILE:HD11	2.01	0.41
1:PA:216:VAL:HG13	1:PA:216:VAL:O	2.19	0.41
1:PB:758:GLU:O	1:PB:762:VAL:HG22	2.20	0.41
1:Q:539:LEU:HD11	1:Q:640:VAL:HG13	2.02	0.41
1:RA:487:VAL:O	1:RA:487:VAL:HG13	2.21	0.41
1:RB:138:VAL:HG12	1:RB:139:ALA:N	2.35	0.41
1:SA:359:ILE:HG23	1:SA:359:ILE:O	2.20	0.41
1:SB:129:PHE:CE1	1:SB:137:VAL:HG11	2.55	0.41
1:T:219:VAL:HG11	1:T:227:LEU:HD22	2.01	0.41
1:T:643:VAL:O	1:T:643:VAL:HG12	2.20	0.41
1:TA:10:ILE:HG22	1:TA:47:PRO:HB3	2.03	0.41
1:TA:138:VAL:HG12	1:TA:139:ALA:N	2.34	0.41
1:UB:138:VAL:HG12	1:UB:139:ALA:N	2.34	0.41
1:UB:570:ASP:OD1	1:UB:570:ASP:C	2.62	0.41
1:V:643:VAL:O	1:V:643:VAL:HG12	2.20	0.41
1:VA:176:LEU:HD21	1:VA:196:TRP:CE2	2.54	0.41
1:VB:779:LEU:HD21	1:WB:774:ARG:HG2	2.02	0.41
1:XA:366:VAL:O	1:XA:366:VAL:HG13	2.19	0.41
1:XB:654:LEU:HD22	1:YB:534:HIS:CG	2.55	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:YA:48:MET:HE2	1:YA:48:MET:HA	2.02	0.41
1:YB:216:VAL:HG13	1:YB:216:VAL:O	2.19	0.41
1:Z:597:ARG:O	1:Z:601:THR:HG23	2.20	0.41
1:ZB:359:ILE:O	1:ZB:359:ILE:HG23	2.20	0.41
1:AB:219:VAL:HG11	1:AB:227:LEU:HD13	2.02	0.41
1:BA:393:VAL:HG22	1:BA:411:ASP:C	2.45	0.41
1:BB:138:VAL:HG12	1:BB:139:ALA:N	2.35	0.41
1:BB:489:LEU:HD21	1:BB:495:PHE:CZ	2.55	0.41
1:C:44:LEU:HD22	1:C:44:LEU:N	2.34	0.41
1:CB:413:VAL:HG22	1:CB:414:LEU:N	2.36	0.41
1:D:176:LEU:HD21	1:D:196:TRP:CE2	2.55	0.41
1:EB:56:ARG:HE	1:FB:127:LEU:CD2	2.34	0.41
1:EB:529:ILE:HD13	1:EB:583:VAL:HG11	2.02	0.41
1:F:121:LEU:HD23	1:F:162:ILE:HG22	2.02	0.41
1:FA:597:ARG:O	1:FA:601:THR:HG23	2.20	0.41
1:FB:359:ILE:O	1:FB:359:ILE:HG23	2.20	0.41
1:GB:393:VAL:HG22	1:GB:411:ASP:C	2.45	0.41
1:H:393:VAL:HG22	1:H:411:ASP:C	2.46	0.41
1:HB:44:LEU:HD22	1:HB:44:LEU:N	2.34	0.41
1:HB:56:ARG:HE	1:IB:127:LEU:CD2	2.33	0.41
1:I:539:LEU:CD1	1:I:640:VAL:HG13	2.51	0.41
1:IB:176:LEU:HD21	1:IB:196:TRP:NE1	2.35	0.41
1:IB:359:ILE:O	1:IB:359:ILE:HG23	2.20	0.41
1:J:48:MET:HE2	1:J:48:MET:HA	2.02	0.41
1:JA:44:LEU:HD22	1:JA:44:LEU:N	2.35	0.41
1:JA:121:LEU:HD23	1:JA:162:ILE:HG22	2.00	0.41
1:JB:176:LEU:HD21	1:JB:196:TRP:NE1	2.36	0.41
1:K:57:HIS:O	1:K:58:TYR:HD1	2.03	0.41
1:KB:359:ILE:O	1:KB:359:ILE:HG23	2.20	0.41
1:L:63:ASN:OD1	1:L:104:VAL:HG12	2.19	0.41
1:LA:129:PHE:CE1	1:LA:137:VAL:HG11	2.55	0.41
1:LB:570:ASP:OD1	1:LB:570:ASP:C	2.63	0.41
1:M:219:VAL:HG11	1:M:227:LEU:HD13	2.02	0.41
1:M:219:VAL:HG11	1:M:227:LEU:HD22	2.01	0.41
1:MA:643:VAL:HG12	1:MA:643:VAL:O	2.20	0.41
1:OA:138:VAL:HG12	1:OA:139:ALA:N	2.34	0.41
1:OB:176:LEU:HD21	1:OB:196:TRP:CE2	2.55	0.41
1:P:177:ARG:NH1	1:P:213:LEU:HD11	2.35	0.41
1:Q:44:LEU:HD22	1:Q:44:LEU:N	2.35	0.41
1:QB:129:PHE:CE1	1:QB:137:VAL:HG11	2.55	0.41
1:R:359:ILE:O	1:R:359:ILE:HG23	2.21	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:RA:17:HIS:CB	1:RA:44:LEU:HD23	2.51	0.41
1:SB:359:ILE:O	1:SB:359:ILE:HG23	2.20	0.41
1:T:129:PHE:CE1	1:T:137:VAL:HG11	2.56	0.41
1:UB:144:LEU:HD22	1:UB:163:ILE:HD11	2.03	0.41
1:UB:176:LEU:HD21	1:UB:196:TRP:NE1	2.34	0.41
1:VB:176:LEU:HD21	1:VB:196:TRP:NE1	2.34	0.41
1:WB:28:VAL:O	1:WB:28:VAL:HG13	2.21	0.41
1:X:53:VAL:HG12	1:X:57:HIS:O	2.19	0.41
1:XB:176:LEU:HD21	1:XB:196:TRP:NE1	2.35	0.41
1:XB:208:VAL:HG23	1:XB:209:PHE:CD1	2.55	0.41
1:YA:73:VAL:HG11	1:YA:81:VAL:HG23	2.02	0.41
1:YA:219:VAL:HG11	1:YA:227:LEU:HD13	2.01	0.41
1:YB:359:ILE:O	1:YB:359:ILE:HG23	2.20	0.41
1:YB:523:PHE:CE2	1:YB:545:TRP:CD1	3.08	0.41
1:YB:538:GLN:HB2	1:YB:646:VAL:HG22	2.01	0.41
1:ZB:129:PHE:CE1	1:ZB:137:VAL:HG11	2.55	0.41
1:ZB:643:VAL:O	1:ZB:643:VAL:HG12	2.20	0.41
1:AA:216:VAL:O	1:AA:216:VAL:HG13	2.19	0.41
1:AA:538:GLN:HB2	1:AA:646:VAL:HG22	2.02	0.41
1:B:44:LEU:HD22	1:B:44:LEU:H	1.86	0.41
1:BA:56:ARG:HE	1:CA:127:LEU:CD2	2.33	0.41
1:BB:56:ARG:HE	1:CB:127:LEU:CD2	2.34	0.41
1:BB:73:VAL:HG11	1:BB:81:VAL:HG23	2.02	0.41
1:BB:129:PHE:CE1	1:BB:137:VAL:HG11	2.55	0.41
1:CA:144:LEU:HD22	1:CA:163:ILE:HD11	2.03	0.41
1:CB:53:VAL:HG21	1:CB:59:CYS:HB2	2.03	0.41
1:D:144:LEU:HD22	1:D:163:ILE:HD11	2.02	0.41
1:DA:3:THR:HG22	1:DA:6:PHE:HA	2.02	0.41
1:DB:177:ARG:NH1	1:DB:213:LEU:HD11	2.35	0.41
1:DB:538:GLN:HB2	1:DB:646:VAL:HG22	2.02	0.41
1:FA:44:LEU:HD22	1:FA:44:LEU:H	1.85	0.41
1:GB:539:LEU:CD1	1:GB:640:VAL:HG13	2.50	0.41
1:H:56:ARG:HE	1:I:127:LEU:CD2	2.32	0.41
1:H:57:HIS:O	1:H:58:TYR:HD1	2.03	0.41
1:H:129:PHE:CE1	1:H:137:VAL:HG11	2.56	0.41
1:H:176:LEU:HD21	1:H:196:TRP:CE2	2.55	0.41
1:HA:165:ALA:HB2	1:HA:205:LEU:HD13	2.00	0.41
1:HB:168:ILE:HG22	1:HB:215:LEU:CD2	2.48	0.41
1:IA:779:LEU:HD21	1:JA:774:ARG:HG2	2.02	0.41
1:KA:56:ARG:HE	1:LA:127:LEU:CD2	2.34	0.41
1:KA:529:ILE:HD13	1:KA:583:VAL:HG11	2.00	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:L:57:HIS:O	1:L:58:TYR:HD1	2.03	0.41
1:LB:44:LEU:HD22	1:LB:44:LEU:N	2.35	0.41
1:M:144:LEU:HD22	1:M:163:ILE:HD11	2.02	0.41
1:M:316:LEU:HD12	1:M:316:LEU:HA	1.95	0.41
1:NB:219:VAL:HG11	1:NB:227:LEU:HD13	2.03	0.41
1:O:47:PRO:O	1:O:48:MET:HE2	2.19	0.41
1:O:129:PHE:CE1	1:O:137:VAL:HG11	2.56	0.41
1:OB:523:PHE:CD2	1:OB:568:VAL:HG23	2.55	0.41
1:P:176:LEU:HD21	1:P:196:TRP:NE1	2.35	0.41
1:PA:17:HIS:CB	1:PA:44:LEU:HD23	2.50	0.41
1:PA:489:LEU:HD21	1:PA:495:PHE:CZ	2.56	0.41
1:PB:48:MET:HA	1:PB:48:MET:HE3	2.02	0.41
1:Q:44:LEU:HD22	1:Q:44:LEU:H	1.84	0.41
1:Q:138:VAL:HG12	1:Q:139:ALA:N	2.35	0.41
1:Q:176:LEU:HD21	1:Q:196:TRP:CE2	2.54	0.41
1:QB:44:LEU:HD22	1:QB:44:LEU:N	2.35	0.41
1:R:539:LEU:CD1	1:R:640:VAL:HG13	2.49	0.41
1:SB:538:GLN:HB2	1:SB:646:VAL:HG22	2.03	0.41
1:TA:121:LEU:HD23	1:TA:162:ILE:HG22	2.02	0.41
1:UA:176:LEU:HD21	1:UA:196:TRP:CE2	2.55	0.41
1:UA:539:LEU:HD11	1:UA:640:VAL:HG13	2.01	0.41
1:V:216:VAL:O	1:V:216:VAL:HG13	2.19	0.41
1:VB:48:MET:HE2	1:VB:48:MET:HA	2.02	0.41
1:W:208:VAL:HG23	1:W:209:PHE:CD1	2.55	0.41
1:WA:176:LEU:HD21	1:WA:196:TRP:NE1	2.35	0.41
1:WA:523:PHE:CE2	1:WA:545:TRP:CD1	3.09	0.41
1:WB:529:ILE:HD13	1:WB:583:VAL:HG11	2.03	0.41
1:X:176:LEU:HD21	1:X:196:TRP:CE2	2.55	0.41
1:Y:10:ILE:HG22	1:Y:47:PRO:HB3	2.02	0.41
1:YB:393:VAL:HG22	1:YB:411:ASP:C	2.46	0.41
1:ZB:529:ILE:HD13	1:ZB:583:VAL:HG11	2.02	0.41
1:AB:405:THR:HG21	1:BB:393:VAL:O	2.21	0.41
1:AC:219:VAL:HG11	1:AC:227:LEU:HD13	2.02	0.41
1:B:129:PHE:CE1	1:B:137:VAL:HG11	2.56	0.41
1:B:359:ILE:O	1:B:359:ILE:HG23	2.20	0.41
1:B:643:VAL:HG12	1:B:643:VAL:O	2.20	0.41
1:BA:487:VAL:O	1:BA:487:VAL:HG13	2.20	0.41
1:BB:359:ILE:O	1:BB:359:ILE:HG23	2.20	0.41
1:CA:129:PHE:CE1	1:CA:137:VAL:HG11	2.55	0.41
1:CA:758:GLU:O	1:CA:762:VAL:HG22	2.20	0.41
1:DA:44:LEU:HD22	1:DA:44:LEU:N	2.35	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:DA:326:LEU:HD12	1:DA:363:LEU:HD21	2.02	0.41
1:DB:138:VAL:HG12	1:DB:139:ALA:N	2.35	0.41
1:DB:143:TRP:HH2	1:DB:157:VAL:HG22	1.85	0.41
1:DB:176:LEU:HD21	1:DB:196:TRP:CE2	2.55	0.41
1:F:516:LEU:HD23	1:F:516:LEU:C	2.45	0.41
1:GA:129:PHE:CE1	1:GA:137:VAL:HG11	2.56	0.41
1:I:176:LEU:HD21	1:I:196:TRP:NE1	2.35	0.41
1:IA:393:VAL:HG22	1:IA:411:ASP:C	2.45	0.41
1:KA:779:LEU:HD21	1:LA:774:ARG:HG2	2.01	0.41
1:KB:541:LEU:CD2	1:KB:640:VAL:HG22	2.50	0.41
1:LB:17:HIS:CB	1:LB:44:LEU:HD23	2.50	0.41
1:LB:57:HIS:O	1:LB:58:TYR:HD1	2.04	0.41
1:LB:121:LEU:HD23	1:LB:162:ILE:HG22	2.02	0.41
1:LB:597:ARG:O	1:LB:601:THR:HG23	2.21	0.41
1:MA:176:LEU:HD21	1:MA:196:TRP:NE1	2.35	0.41
1:MA:219:VAL:HG11	1:MA:227:LEU:HD22	2.01	0.41
1:MB:143:TRP:HH2	1:MB:157:VAL:HG22	1.85	0.41
1:MB:516:LEU:O	1:MB:516:LEU:HD23	2.20	0.41
1:MB:538:GLN:HB2	1:MB:646:VAL:HG22	2.02	0.41
1:NA:219:VAL:HG11	1:NA:227:LEU:HD13	2.03	0.41
1:OB:143:TRP:HH2	1:OB:157:VAL:HG22	1.84	0.41
1:P:176:LEU:HD21	1:P:196:TRP:CE2	2.55	0.41
1:P:643:VAL:O	1:P:643:VAL:HG12	2.20	0.41
1:PA:144:LEU:HD22	1:PA:163:ILE:HD11	2.03	0.41
1:PB:539:LEU:HD11	1:PB:640:VAL:HG13	2.03	0.41
1:Q:176:LEU:HD21	1:Q:196:TRP:NE1	2.35	0.41
1:Q:216:VAL:O	1:Q:216:VAL:HG13	2.19	0.41
1:Q:487:VAL:O	1:Q:487:VAL:HG13	2.19	0.41
1:QA:138:VAL:HG12	1:QA:139:ALA:N	2.34	0.41
1:QB:487:VAL:O	1:QB:487:VAL:HG13	2.19	0.41
1:R:165:ALA:HB2	1:R:205:LEU:HD13	2.02	0.41
1:RB:643:VAL:O	1:RB:643:VAL:HG12	2.21	0.41
1:S:539:LEU:CD1	1:S:640:VAL:HG13	2.50	0.41
1:SB:168:ILE:HG22	1:SB:215:LEU:CD2	2.47	0.41
1:SB:393:VAL:HG22	1:SB:411:ASP:C	2.46	0.41
1:T:529:ILE:HD13	1:T:583:VAL:HG11	2.03	0.41
1:UA:654:LEU:HD22	1:VA:534:HIS:CG	2.55	0.41
1:V:44:LEU:HD22	1:V:44:LEU:N	2.36	0.41
1:VA:56:ARG:HE	1:WA:127:LEU:CD2	2.33	0.41
1:W:56:ARG:HE	1:X:127:LEU:CD2	2.34	0.41
1:WB:176:LEU:HD21	1:WB:196:TRP:NE1	2.35	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:X:165:ALA:HB2	1:X:205:LEU:HD13	2.01	0.41
1:XB:47:PRO:O	1:XB:48:MET:HE2	2.21	0.41
1:XB:541:LEU:CD2	1:XB:640:VAL:HG22	2.50	0.41
1:Y:57:HIS:O	1:Y:58:TYR:HD1	2.03	0.41
1:Y:359:ILE:HG23	1:Y:359:ILE:O	2.20	0.41
1:YA:57:HIS:O	1:YA:58:TYR:HD1	2.04	0.41
1:Z:539:LEU:HD21	1:Z:599:ILE:HD11	2.03	0.41
1:ZA:57:HIS:O	1:ZA:58:TYR:HD1	2.04	0.41
1:AB:489:LEU:HD21	1:AB:495:PHE:CZ	2.55	0.41
1:AB:809:ASP:HA	1:AB:812:VAL:HG22	2.03	0.41
1:B:17:HIS:CB	1:B:44:LEU:HD23	2.50	0.41
1:BA:523:PHE:CD2	1:BA:568:VAL:HG23	2.55	0.41
1:CA:138:VAL:HG12	1:CA:139:ALA:N	2.35	0.41
1:CA:176:LEU:HD21	1:CA:196:TRP:NE1	2.36	0.41
1:CA:597:ARG:O	1:CA:601:THR:HG23	2.20	0.41
1:DB:57:HIS:O	1:DB:58:TYR:HD1	2.03	0.41
1:FB:17:HIS:CB	1:FB:44:LEU:HD23	2.50	0.41
1:GA:52:THR:O	1:GA:109:ILE:HD11	2.19	0.41
1:HA:219:VAL:HG11	1:HA:227:LEU:HD13	2.02	0.41
1:I:539:LEU:HD21	1:I:599:ILE:HD11	2.02	0.41
1:J:44:LEU:HD22	1:J:44:LEU:H	1.85	0.41
1:J:57:HIS:O	1:J:58:TYR:HD1	2.03	0.41
1:KB:176:LEU:HD21	1:KB:196:TRP:CE2	2.55	0.41
1:LA:643:VAL:O	1:LA:643:VAL:HG12	2.20	0.41
1:M:393:VAL:HG22	1:M:411:ASP:C	2.45	0.41
1:MB:56:ARG:HE	1:NB:127:LEU:CD2	2.33	0.41
1:NB:56:ARG:HE	1:OB:127:LEU:HD21	1.85	0.41
1:NB:393:VAL:HG22	1:NB:411:ASP:C	2.46	0.41
1:NB:487:VAL:O	1:NB:487:VAL:HG13	2.20	0.41
1:OA:17:HIS:CB	1:OA:44:LEU:HD23	2.50	0.41
1:OB:662:ILE:O	1:OB:666:THR:HG23	2.20	0.41
1:P:538:GLN:HB2	1:P:646:VAL:HG22	2.02	0.41
1:P:597:ARG:O	1:P:601:THR:HG23	2.21	0.41
1:PA:3:THR:HG22	1:PA:6:PHE:HA	2.03	0.41
1:Q:56:ARG:HE	1:R:127:LEU:CD2	2.34	0.41
1:Q:95:ASP:O	1:Q:96:PRO:C	2.63	0.41
1:QA:30:VAL:HG22	1:QA:50:MET:HE2	2.03	0.41
1:SA:129:PHE:CE1	1:SA:137:VAL:HG11	2.56	0.41
1:SA:143:TRP:HH2	1:SA:157:VAL:HG22	1.85	0.41
1:TA:176:LEU:HD21	1:TA:196:TRP:CE2	2.56	0.41
1:VA:393:VAL:HG22	1:VA:411:ASP:C	2.46	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:WB:138:VAL:HG12	1:WB:139:ALA:N	2.34	0.41
1:WB:539:LEU:CD1	1:WB:599:ILE:HD11	2.50	0.41
1:XB:168:ILE:HG22	1:XB:215:LEU:CD2	2.49	0.41
1:XB:583:VAL:O	1:XB:583:VAL:HG12	2.20	0.41
1:Y:219:VAL:HG11	1:Y:227:LEU:HD22	2.02	0.41
1:YA:17:HIS:CB	1:YA:44:LEU:HD23	2.51	0.41
1:YB:129:PHE:CE1	1:YB:137:VAL:HG11	2.55	0.41
1:YB:168:ILE:HG22	1:YB:215:LEU:CD2	2.45	0.41
1:A:643:VAL:O	1:A:643:VAL:HG12	2.21	0.41
1:AA:10:ILE:HG22	1:AA:47:PRO:HB3	2.03	0.41
1:AA:138:VAL:HG12	1:AA:139:ALA:N	2.35	0.41
1:AB:597:ARG:O	1:AB:601:THR:HG23	2.20	0.41
1:C:541:LEU:CD2	1:C:640:VAL:HG22	2.50	0.41
1:CA:643:VAL:O	1:CA:643:VAL:HG12	2.20	0.41
1:DA:47:PRO:C	1:DA:48:MET:HE2	2.46	0.41
1:DB:165:ALA:HB2	1:DB:205:LEU:HD13	2.02	0.41
1:DB:391:GLN:HB2	1:DB:398:VAL:HG22	2.02	0.41
1:E:143:TRP:HH2	1:E:157:VAL:HG22	1.85	0.41
1:E:176:LEU:HD21	1:E:196:TRP:NE1	2.36	0.41
1:EA:73:VAL:HG11	1:EA:81:VAL:HG23	2.03	0.41
1:EB:57:HIS:O	1:EB:58:TYR:HD1	2.03	0.41
1:FA:359:ILE:O	1:FA:359:ILE:HG23	2.20	0.41
1:FA:393:VAL:HG22	1:FA:411:ASP:C	2.45	0.41
1:FB:143:TRP:HH2	1:FB:157:VAL:HG22	1.85	0.41
1:G:538:GLN:HB2	1:G:646:VAL:HG22	2.03	0.41
1:JB:219:VAL:HG11	1:JB:227:LEU:HD13	2.03	0.41
1:K:168:ILE:HG22	1:K:215:LEU:CD2	2.48	0.41
1:KA:56:ARG:HE	1:LA:127:LEU:HD21	1.85	0.41
1:KA:539:LEU:HD11	1:KA:640:VAL:HG13	2.02	0.41
1:LA:168:ILE:HG22	1:LA:215:LEU:CD2	2.48	0.41
1:M:809:ASP:HA	1:M:812:VAL:HG22	2.02	0.41
1:N:129:PHE:CE1	1:N:137:VAL:HG11	2.55	0.41
1:N:138:VAL:HG12	1:N:139:ALA:N	2.35	0.41
1:N:176:LEU:HD21	1:N:196:TRP:NE1	2.35	0.41
1:OA:176:LEU:HD21	1:OA:196:TRP:NE1	2.35	0.41
1:PB:57:HIS:O	1:PB:58:TYR:HD1	2.03	0.41
1:PB:359:ILE:O	1:PB:359:ILE:HG23	2.20	0.41
1:Q:57:HIS:O	1:Q:58:TYR:HD1	2.03	0.41
1:QA:541:LEU:CD2	1:QA:640:VAL:HG22	2.50	0.41
1:RA:737:GLY:O	1:RA:741:VAL:HG23	2.21	0.41
1:RB:129:PHE:CE1	1:RB:137:VAL:HG11	2.56	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:S:176:LEU:HD21	1:S:196:TRP:CE2	2.55	0.41
1:S:176:LEU:HD21	1:S:196:TRP:NE1	2.34	0.41
1:TA:73:VAL:HG11	1:TA:81:VAL:HG23	2.03	0.41
1:TA:143:TRP:HH2	1:TA:157:VAL:HG22	1.85	0.41
1:TB:219:VAL:HG11	1:TB:227:LEU:HD22	2.02	0.41
1:UB:662:ILE:O	1:UB:666:THR:HG23	2.20	0.41
1:WA:95:ASP:O	1:WA:96:PRO:C	2.64	0.41
1:XA:165:ALA:HB2	1:XA:205:LEU:HD13	2.03	0.41
1:YA:129:PHE:CE1	1:YA:137:VAL:HG11	2.55	0.41
1:ZB:144:LEU:HD22	1:ZB:163:ILE:HD11	2.03	0.41
1:ZB:391:GLN:HB2	1:ZB:398:VAL:HG22	2.03	0.41
1:ZB:539:LEU:HD21	1:ZB:599:ILE:HD11	2.03	0.41
1:AA:643:VAL:O	1:AA:643:VAL:HG12	2.21	0.41
1:AB:17:HIS:CB	1:AB:44:LEU:HD23	2.51	0.41
1:AC:56:ARG:HE	1:OA:127:LEU:CD2	2.34	0.41
1:AC:579:VAL:O	1:AC:583:VAL:HG12	2.21	0.41
1:B:405:THR:HG21	1:C:393:VAL:O	2.21	0.41
1:D:57:HIS:O	1:D:58:TYR:HD1	2.03	0.41
1:D:539:LEU:HD11	1:D:640:VAL:HG13	2.03	0.41
1:DA:359:ILE:O	1:DA:359:ILE:HG23	2.21	0.41
1:DA:597:ARG:O	1:DA:601:THR:HG23	2.20	0.41
1:DB:539:LEU:HD12	1:DB:540:GLN:N	2.36	0.41
1:E:393:VAL:HG22	1:E:411:ASP:C	2.45	0.41
1:EA:129:PHE:CE1	1:EA:137:VAL:HG11	2.56	0.41
1:EB:597:ARG:O	1:EB:601:THR:HG23	2.20	0.41
1:F:541:LEU:CD2	1:F:640:VAL:HG22	2.51	0.41
1:FA:47:PRO:C	1:FA:48:MET:HE2	2.46	0.41
1:FB:165:ALA:HB2	1:FB:205:LEU:HD13	2.03	0.41
1:FB:176:LEU:HD21	1:FB:196:TRP:NE1	2.36	0.41
1:G:597:ARG:O	1:G:601:THR:HG23	2.21	0.41
1:G:779:LEU:HD21	1:H:774:ARG:HG2	2.02	0.41
1:HB:121:LEU:HD22	1:HB:159:VAL:HG13	2.02	0.41
1:HB:662:ILE:O	1:HB:666:THR:HG23	2.21	0.41
1:IB:539:LEU:HD11	1:IB:640:VAL:HG13	2.03	0.41
1:J:138:VAL:HG12	1:J:139:ALA:N	2.35	0.41
1:J:529:ILE:HD13	1:J:583:VAL:HG11	2.03	0.41
1:JA:56:ARG:HE	1:KA:127:LEU:HD21	1.85	0.41
1:JB:56:ARG:HE	1:KB:127:LEU:HD21	1.85	0.41
1:KA:541:LEU:CD2	1:KA:640:VAL:HG22	2.51	0.41
1:KB:144:LEU:HD22	1:KB:163:ILE:HD11	2.03	0.41
1:LA:44:LEU:HD22	1:LA:44:LEU:N	2.35	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:MA:541:LEU:CD2	1:MA:640:VAL:HG22	2.51	0.41
1:MB:807:ILE:HD12	1:MB:807:ILE:HA	1.98	0.41
1:NB:143:TRP:HH2	1:NB:157:VAL:HG22	1.85	0.41
1:NB:165:ALA:HB2	1:NB:205:LEU:HD13	2.02	0.41
1:NB:538:GLN:HB2	1:NB:646:VAL:HG22	2.03	0.41
1:OB:17:HIS:CB	1:OB:44:LEU:HD23	2.50	0.41
1:P:47:PRO:C	1:P:48:MET:HE2	2.46	0.41
1:P:138:VAL:HG12	1:P:139:ALA:N	2.35	0.41
1:P:219:VAL:HG11	1:P:227:LEU:HD13	2.03	0.41
1:PB:176:LEU:HD21	1:PB:196:TRP:NE1	2.35	0.41
1:PB:539:LEU:CD1	1:PB:640:VAL:HG13	2.50	0.41
1:QB:393:VAL:HG22	1:QB:411:ASP:C	2.46	0.41
1:QB:597:ARG:O	1:QB:601:THR:HG23	2.21	0.41
1:R:17:HIS:CB	1:R:44:LEU:HD23	2.51	0.41
1:SA:216:VAL:O	1:SA:216:VAL:HG13	2.19	0.41
1:T:176:LEU:HD21	1:T:196:TRP:NE1	2.35	0.41
1:TA:216:VAL:O	1:TA:216:VAL:HG13	2.19	0.41
1:TB:56:ARG:HE	1:UB:127:LEU:CD2	2.33	0.41
1:TB:176:LEU:HD21	1:TB:196:TRP:NE1	2.35	0.41
1:UB:143:TRP:HH2	1:UB:157:VAL:HG22	1.85	0.41
1:VB:597:ARG:O	1:VB:601:THR:HG23	2.21	0.41
1:W:662:ILE:O	1:W:666:THR:HG23	2.20	0.41
1:WB:366:VAL:O	1:WB:366:VAL:HG13	2.20	0.41
1:WB:405:THR:HG21	1:XB:393:VAL:O	2.21	0.41
1:X:144:LEU:HD22	1:X:163:ILE:HD11	2.03	0.41
1:X:405:THR:HG21	1:Y:393:VAL:O	2.21	0.41
1:X:643:VAL:O	1:X:643:VAL:HG12	2.20	0.41
1:XB:95:ASP:O	1:XB:96:PRO:C	2.63	0.41
1:YA:44:LEU:HD22	1:YA:44:LEU:H	1.86	0.41
1:YB:330:GLN:C	1:YB:407:MET:HE1	2.46	0.41
1:Z:176:LEU:HD21	1:Z:196:TRP:NE1	2.35	0.41
1:Z:539:LEU:HD11	1:Z:640:VAL:HG13	2.02	0.41
1:Z:737:GLY:O	1:Z:741:VAL:HG23	2.21	0.41
1:A:529:ILE:HD13	1:A:583:VAL:HG11	2.03	0.41
1:AA:143:TRP:HH2	1:AA:157:VAL:HG22	1.85	0.41
1:AB:527:ILE:HD11	1:AB:539:LEU:HD22	2.03	0.41
1:AC:57:HIS:O	1:AC:58:TYR:HD1	2.04	0.41
1:B:144:LEU:HD22	1:B:163:ILE:HD11	2.03	0.41
1:BA:143:TRP:HH2	1:BA:157:VAL:HG22	1.85	0.41
1:BB:57:HIS:O	1:BB:58:TYR:HD1	2.03	0.41
1:CA:809:ASP:HA	1:CA:812:VAL:HG22	2.03	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:D:44:LEU:HD22	1:D:44:LEU:H	1.86	0.41
1:D:643:VAL:O	1:D:643:VAL:HG12	2.21	0.41
1:DA:17:HIS:CB	1:DA:44:LEU:HD23	2.51	0.41
1:E:121:LEU:HD22	1:E:159:VAL:HG13	2.02	0.41
1:EB:44:LEU:HD22	1:EB:44:LEU:N	2.35	0.41
1:EB:129:PHE:CE1	1:EB:137:VAL:HG11	2.56	0.41
1:EB:539:LEU:HD21	1:EB:599:ILE:HD11	2.02	0.41
1:F:523:PHE:CE2	1:F:545:TRP:CD1	3.08	0.41
1:FB:159:VAL:O	1:FB:159:VAL:HG12	2.20	0.41
1:FB:393:VAL:HG22	1:FB:411:ASP:C	2.46	0.41
1:FB:539:LEU:HD11	1:FB:541:LEU:CD2	2.50	0.41
1:G:176:LEU:HD21	1:G:196:TRP:CE2	2.56	0.41
1:GA:56:ARG:HE	1:HA:127:LEU:CD2	2.33	0.41
1:GB:17:HIS:CB	1:GB:44:LEU:HD23	2.51	0.41
1:GB:56:ARG:HE	1:HB:127:LEU:HD21	1.85	0.41
1:GB:57:HIS:O	1:GB:58:TYR:HD1	2.04	0.41
1:GB:176:LEU:HD21	1:GB:196:TRP:CE2	2.55	0.41
1:GB:570:ASP:OD1	1:GB:570:ASP:C	2.64	0.41
1:HA:30:VAL:CG2	1:HA:50:MET:HE3	2.51	0.41
1:HB:539:LEU:HD21	1:HB:599:ILE:HD11	2.02	0.41
1:HB:597:ARG:O	1:HB:601:THR:HG23	2.21	0.41
1:I:121:LEU:HD23	1:I:162:ILE:HG22	2.03	0.41
1:I:523:PHE:CE2	1:I:545:TRP:CD1	3.08	0.41
1:IA:44:LEU:HD22	1:IA:44:LEU:N	2.35	0.41
1:IB:168:ILE:HG22	1:IB:215:LEU:CD2	2.51	0.41
1:IB:539:LEU:HD12	1:IB:540:GLN:N	2.36	0.41
1:JB:44:LEU:HD22	1:JB:44:LEU:N	2.35	0.41
1:JB:57:HIS:O	1:JB:58:TYR:HD1	2.03	0.41
1:JB:129:PHE:CE1	1:JB:137:VAL:HG11	2.55	0.41
1:JB:143:TRP:HH2	1:JB:157:VAL:HG22	1.85	0.41
1:KA:138:VAL:HG12	1:KA:139:ALA:N	2.35	0.41
1:KA:143:TRP:HH2	1:KA:157:VAL:HG22	1.85	0.41
1:KA:662:ILE:O	1:KA:666:THR:HG23	2.20	0.41
1:KB:281:TYR:CG	1:KB:366:VAL:HG23	2.56	0.41
1:L:168:ILE:HG21	1:L:174:LEU:HB2	2.02	0.41
1:L:176:LEU:HD21	1:L:196:TRP:NE1	2.36	0.41
1:LA:48:MET:HE3	1:LA:48:MET:CA	2.51	0.41
1:LA:393:VAL:HG22	1:LA:411:ASP:C	2.46	0.41
1:LB:73:VAL:HG11	1:LB:81:VAL:HG23	2.03	0.41
1:LB:176:LEU:HD21	1:LB:196:TRP:NE1	2.36	0.41
1:M:539:LEU:HD11	1:M:599:ILE:CD1	2.48	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:M:597:ARG:O	1:M:601:THR:HG23	2.21	0.41
1:MA:168:ILE:HG22	1:MA:215:LEU:CD2	2.46	0.41
1:NB:541:LEU:CD2	1:NB:640:VAL:HG22	2.51	0.41
1:O:405:THR:HG21	1:P:393:VAL:O	2.20	0.41
1:OA:129:PHE:CE1	1:OA:137:VAL:HG11	2.56	0.41
1:OA:643:VAL:O	1:OA:643:VAL:HG12	2.21	0.41
1:OB:44:LEU:HD22	1:OB:44:LEU:H	1.86	0.41
1:OB:487:VAL:O	1:OB:487:VAL:HG13	2.19	0.41
1:P:539:LEU:HD12	1:P:540:GLN:N	2.36	0.41
1:PA:56:ARG:HE	1:QA:127:LEU:CD2	2.34	0.41
1:PA:143:TRP:HH2	1:PA:157:VAL:HG22	1.85	0.41
1:PA:529:ILE:HD13	1:PA:583:VAL:HG11	2.03	0.41
1:PA:597:ARG:O	1:PA:601:THR:HG23	2.21	0.41
1:PB:56:ARG:HE	1:QB:127:LEU:CD2	2.34	0.41
1:PB:809:ASP:HA	1:PB:812:VAL:HG22	2.03	0.41
1:Q:570:ASP:OD1	1:Q:570:ASP:C	2.64	0.41
1:QA:176:LEU:HD21	1:QA:196:TRP:NE1	2.35	0.41
1:QB:165:ALA:HB2	1:QB:205:LEU:HD13	2.03	0.41
1:QB:662:ILE:O	1:QB:666:THR:HG23	2.21	0.41
1:R:53:VAL:HG12	1:R:93:ALA:HA	2.03	0.41
1:R:138:VAL:HG12	1:R:139:ALA:N	2.35	0.41
1:R:159:VAL:O	1:R:159:VAL:HG12	2.20	0.41
1:RA:168:ILE:HG22	1:RA:215:LEU:CD2	2.45	0.41
1:RA:216:VAL:O	1:RA:216:VAL:HG13	2.19	0.41
1:RA:643:VAL:O	1:RA:643:VAL:HG12	2.21	0.41
1:S:56:ARG:HE	1:T:127:LEU:HD21	1.86	0.41
1:SA:47:PRO:C	1:SA:48:MET:HE2	2.45	0.41
1:SA:176:LEU:HD21	1:SA:196:TRP:NE1	2.36	0.41
1:SA:539:LEU:CD1	1:SA:640:VAL:HG13	2.49	0.41
1:SB:176:LEU:HD21	1:SB:196:TRP:NE1	2.35	0.41
1:SB:487:VAL:O	1:SB:487:VAL:HG13	2.20	0.41
1:T:44:LEU:HD22	1:T:44:LEU:N	2.36	0.41
1:T:121:LEU:HD23	1:T:162:ILE:HG22	2.03	0.41
1:T:597:ARG:O	1:T:601:THR:HG23	2.21	0.41
1:TB:129:PHE:CE1	1:TB:137:VAL:HG11	2.56	0.41
1:UA:144:LEU:HD22	1:UA:163:ILE:HD11	2.03	0.41
1:UB:281:TYR:CG	1:UB:366:VAL:HG23	2.56	0.41
1:UB:359:ILE:O	1:UB:359:ILE:HG23	2.20	0.41
1:UB:539:LEU:HD11	1:UB:640:VAL:HG13	2.03	0.41
1:VA:176:LEU:HD21	1:VA:196:TRP:NE1	2.36	0.41
1:W:129:PHE:CE1	1:W:137:VAL:HG11	2.55	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:X:527:ILE:HD11	1:X:539:LEU:HD22	2.03	0.41
1:XA:654:LEU:HD22	1:YA:534:HIS:CG	2.56	0.41
1:XA:779:LEU:HD21	1:YA:774:ARG:HG2	2.03	0.41
1:XB:17:HIS:CB	1:XB:44:LEU:HD23	2.51	0.41
1:XB:56:ARG:HE	1:YB:127:LEU:HD21	1.86	0.41
1:XB:216:VAL:O	1:XB:216:VAL:HG13	2.19	0.41
1:XB:662:ILE:O	1:XB:666:THR:HG23	2.20	0.41
1:Y:17:HIS:CB	1:Y:44:LEU:HD23	2.50	0.41
1:YA:10:ILE:HG22	1:YA:47:PRO:HB3	2.03	0.41
1:YB:597:ARG:O	1:YB:601:THR:HG23	2.21	0.41
1:Z:73:VAL:HG11	1:Z:81:VAL:HG23	2.02	0.41
1:Z:489:LEU:HD21	1:Z:495:PHE:CZ	2.56	0.41
1:ZA:44:LEU:HD22	1:ZA:44:LEU:H	1.86	0.41
1:ZA:597:ARG:O	1:ZA:601:THR:HG23	2.20	0.41
1:ZB:44:LEU:HD22	1:ZB:44:LEU:H	1.85	0.41
1:ZB:176:LEU:HD21	1:ZB:196:TRP:NE1	2.35	0.41
1:A:44:LEU:HD22	1:A:44:LEU:H	1.86	0.41
1:A:393:VAL:HG22	1:A:411:ASP:C	2.45	0.41
1:A:809:ASP:HA	1:A:812:VAL:HG22	2.03	0.41
1:AA:3:THR:HG22	1:AA:6:PHE:HA	2.03	0.41
1:AB:393:VAL:HG22	1:AB:411:ASP:C	2.46	0.41
1:AC:52:THR:O	1:AC:109:ILE:HD11	2.21	0.41
1:B:176:LEU:HD21	1:B:196:TRP:NE1	2.36	0.41
1:B:539:LEU:HD21	1:B:599:ILE:HD11	2.02	0.41
1:C:48:MET:HE2	1:C:48:MET:HA	2.02	0.41
1:CA:359:ILE:HG23	1:CA:359:ILE:O	2.20	0.41
1:CB:758:GLU:O	1:CB:762:VAL:HG22	2.20	0.41
1:E:219:VAL:HG11	1:E:227:LEU:HD13	2.03	0.41
1:E:662:ILE:O	1:E:666:THR:HG23	2.20	0.41
1:EA:30:VAL:CG2	1:EA:50:MET:HE2	2.51	0.41
1:EB:95:ASP:O	1:EB:96:PRO:C	2.63	0.41
1:F:13:TYR:HA	1:F:50:MET:HE3	2.03	0.41
1:FB:487:VAL:O	1:FB:487:VAL:HG13	2.20	0.41
1:GA:529:ILE:HD13	1:GA:583:VAL:HG21	2.03	0.41
1:GB:144:LEU:HD12	1:GB:163:ILE:HD11	2.02	0.41
1:J:17:HIS:CB	1:J:44:LEU:HD23	2.50	0.41
1:J:165:ALA:HB2	1:J:205:LEU:HD13	2.03	0.41
1:JA:28:VAL:O	1:JA:28:VAL:HG13	2.20	0.41
1:KA:48:MET:HE2	1:KA:48:MET:HA	2.02	0.41
1:L:56:ARG:HE	1:M:127:LEU:CD2	2.33	0.41
1:L:359:ILE:O	1:L:359:ILE:HG23	2.20	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:LA:165:ALA:HB2	1:LA:205:LEU:HD13	2.03	0.41
1:LB:281:TYR:CG	1:LB:366:VAL:HG23	2.56	0.41
1:M:779:LEU:HD21	1:N:774:ARG:HG2	2.03	0.41
1:MA:144:LEU:HD22	1:MA:163:ILE:HD11	2.03	0.41
1:MA:281:TYR:CG	1:MA:366:VAL:HG23	2.55	0.41
1:N:539:LEU:HD21	1:N:599:ILE:HD11	2.03	0.41
1:O:487:VAL:O	1:O:487:VAL:HG13	2.19	0.41
1:OA:359:ILE:O	1:OA:359:ILE:HG23	2.21	0.41
1:P:143:TRP:HH2	1:P:157:VAL:HG22	1.86	0.41
1:PA:405:THR:HG21	1:QA:393:VAL:O	2.21	0.41
1:RA:144:LEU:HD22	1:RA:163:ILE:HD11	2.03	0.41
1:RA:489:LEU:HD21	1:RA:495:PHE:CZ	2.56	0.41
1:S:44:LEU:HD22	1:S:44:LEU:H	1.85	0.41
1:SA:487:VAL:O	1:SA:487:VAL:HG13	2.20	0.41
1:SB:539:LEU:CD1	1:SB:640:VAL:HG13	2.51	0.41
1:UA:597:ARG:O	1:UA:601:THR:HG23	2.21	0.41
1:UB:516:LEU:HD23	1:UB:516:LEU:C	2.45	0.41
1:UB:597:ARG:O	1:UB:601:THR:HG23	2.21	0.41
1:W:57:HIS:O	1:W:58:TYR:HD1	2.04	0.41
1:W:538:GLN:HB2	1:W:646:VAL:HG22	2.02	0.41
1:W:758:GLU:O	1:W:762:VAL:HG22	2.20	0.41
1:WA:654:LEU:HD22	1:XA:534:HIS:CG	2.56	0.41
1:XA:138:VAL:HG12	1:XA:139:ALA:N	2.35	0.41
1:XB:56:ARG:HE	1:YB:127:LEU:CD2	2.34	0.41
1:Y:281:TYR:CG	1:Y:366:VAL:HG23	2.55	0.41
1:YA:138:VAL:HG12	1:YA:139:ALA:N	2.35	0.41
1:Z:143:TRP:HH2	1:Z:157:VAL:HG22	1.85	0.41
1:ZA:10:ILE:HG22	1:ZA:47:PRO:HB3	2.02	0.41
1:ZB:165:ALA:HB2	1:ZB:205:LEU:HD13	2.02	0.41
1:ZB:281:TYR:CG	1:ZB:366:VAL:HG23	2.55	0.41
1:B:489:LEU:HD21	1:B:495:PHE:CZ	2.56	0.40
1:BA:138:VAL:HG12	1:BA:139:ALA:N	2.35	0.40
1:BB:176:LEU:HD21	1:BB:196:TRP:NE1	2.35	0.40
1:C:44:LEU:HD22	1:C:44:LEU:H	1.86	0.40
1:CB:129:PHE:CE1	1:CB:137:VAL:HG11	2.56	0.40
1:E:539:LEU:HD21	1:E:599:ILE:HD11	2.02	0.40
1:EA:17:HIS:CB	1:EA:44:LEU:HD23	2.51	0.40
1:EB:165:ALA:HB2	1:EB:205:LEU:HD13	2.03	0.40
1:EB:541:LEU:CD2	1:EB:640:VAL:HG22	2.51	0.40
1:FA:95:ASP:O	1:FA:96:PRO:C	2.64	0.40
1:FB:539:LEU:CD1	1:FB:640:VAL:HG13	2.51	0.40

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:G:17:HIS:HB3	1:G:44:LEU:HD23	2.02	0.40
1:G:30:VAL:HG22	1:G:50:MET:HE2	2.03	0.40
1:GA:786:GLN:OE1	1:HA:781:VAL:HG11	2.20	0.40
1:GB:632:GLY:O	1:GB:634:VAL:HG23	2.22	0.40
1:HA:144:LEU:HD22	1:HA:163:ILE:HD11	2.03	0.40
1:I:539:LEU:HD11	1:I:640:VAL:HG13	2.03	0.40
1:I:809:ASP:HA	1:I:812:VAL:HG22	2.03	0.40
1:IB:538:GLN:HB2	1:IB:646:VAL:HG22	2.03	0.40
1:J:539:LEU:HD21	1:J:599:ILE:HD11	2.01	0.40
1:JA:529:ILE:HD13	1:JA:583:VAL:HG11	2.02	0.40
1:JB:541:LEU:CD2	1:JB:640:VAL:HG22	2.51	0.40
1:KA:129:PHE:CE1	1:KA:137:VAL:HG11	2.57	0.40
1:KB:168:ILE:HG22	1:KB:215:LEU:CD2	2.48	0.40
1:LA:413:VAL:HG22	1:LA:414:LEU:N	2.36	0.40
1:M:405:THR:HG21	1:N:393:VAL:O	2.21	0.40
1:M:413:VAL:HG22	1:M:414:LEU:N	2.36	0.40
1:MB:393:VAL:HG22	1:MB:411:ASP:C	2.46	0.40
1:N:539:LEU:HD11	1:N:640:VAL:HG13	2.03	0.40
1:O:17:HIS:CB	1:O:44:LEU:HD23	2.51	0.40
1:OA:523:PHE:CE2	1:OA:545:TRP:CD1	3.09	0.40
1:OB:539:LEU:HD12	1:OB:540:GLN:N	2.37	0.40
1:PA:176:LEU:HD21	1:PA:196:TRP:NE1	2.35	0.40
1:Q:541:LEU:CD2	1:Q:640:VAL:HG22	2.52	0.40
1:QB:176:LEU:HD21	1:QB:196:TRP:NE1	2.35	0.40
1:QB:359:ILE:O	1:QB:359:ILE:HG23	2.22	0.40
1:RA:176:LEU:HD21	1:RA:196:TRP:NE1	2.35	0.40
1:RA:405:THR:HG21	1:SA:393:VAL:O	2.21	0.40
1:RB:393:VAL:HG22	1:RB:411:ASP:C	2.47	0.40
1:SA:57:HIS:O	1:SA:58:TYR:HD1	2.04	0.40
1:SA:219:VAL:HG11	1:SA:227:LEU:HD13	2.03	0.40
1:T:17:HIS:HB3	1:T:44:LEU:HD23	2.02	0.40
1:TA:48:MET:HA	1:TA:48:MET:HE2	2.03	0.40
1:TA:538:GLN:HB2	1:TA:646:VAL:HG22	2.02	0.40
1:TB:143:TRP:HH2	1:TB:157:VAL:HG22	1.86	0.40
1:UB:129:PHE:CE1	1:UB:137:VAL:HG11	2.56	0.40
1:V:597:ARG:O	1:V:601:THR:HG23	2.22	0.40
1:W:143:TRP:HH2	1:W:157:VAL:HG22	1.86	0.40
1:WA:539:LEU:CD1	1:WA:640:VAL:HG13	2.50	0.40
1:X:281:TYR:CG	1:X:366:VAL:HG23	2.57	0.40
1:XB:405:THR:HG21	1:YB:393:VAL:O	2.21	0.40
1:Y:597:ARG:O	1:Y:601:THR:HG23	2.21	0.40

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:YB:487:VAL:O	1:YB:487:VAL:HG13	2.20	0.40
1:ZA:359:ILE:O	1:ZA:359:ILE:HG23	2.21	0.40
1:AA:597:ARG:O	1:AA:601:THR:HG23	2.22	0.40
1:BB:539:LEU:HD11	1:BB:640:VAL:HG13	2.03	0.40
1:CA:95:ASP:O	1:CA:96:PRO:C	2.65	0.40
1:CB:570:ASP:OD1	1:CB:570:ASP:C	2.65	0.40
1:DA:56:ARG:HE	1:EA:127:LEU:CD2	2.34	0.40
1:DB:570:ASP:OD1	1:DB:570:ASP:C	2.64	0.40
1:E:523:PHE:CE2	1:E:545:TRP:CD1	3.09	0.40
1:I:30:VAL:CG2	1:I:50:MET:HE3	2.51	0.40
1:IA:366:VAL:HG13	1:IA:366:VAL:O	2.21	0.40
1:IB:219:VAL:HG11	1:IB:227:LEU:HD13	2.03	0.40
1:JB:121:LEU:HD23	1:JB:162:ILE:HG22	2.03	0.40
1:K:138:VAL:HG12	1:K:139:ALA:N	2.35	0.40
1:KA:57:HIS:O	1:KA:58:TYR:HD1	2.03	0.40
1:KA:95:ASP:O	1:KA:96:PRO:C	2.64	0.40
1:KA:219:VAL:HG11	1:KA:227:LEU:HD13	2.03	0.40
1:KA:654:LEU:HD22	1:LA:534:HIS:CG	2.55	0.40
1:L:73:VAL:HG11	1:L:81:VAL:HG23	2.02	0.40
1:NB:129:PHE:CE1	1:NB:137:VAL:HG11	2.56	0.40
1:OB:144:LEU:HD22	1:OB:163:ILE:HD11	2.04	0.40
1:P:10:ILE:HG22	1:P:47:PRO:HB3	2.03	0.40
1:P:654:LEU:HD22	1:Q:534:HIS:CG	2.56	0.40
1:QA:30:VAL:CG2	1:QA:50:MET:HE2	2.52	0.40
1:R:597:ARG:O	1:R:601:THR:HG23	2.21	0.40
1:R:643:VAL:HG12	1:R:643:VAL:O	2.21	0.40
1:S:541:LEU:CD2	1:S:640:VAL:HG22	2.51	0.40
1:SA:138:VAL:HG12	1:SA:139:ALA:N	2.35	0.40
1:SB:809:ASP:HA	1:SB:812:VAL:HG22	2.04	0.40
1:V:176:LEU:HD21	1:V:196:TRP:NE1	2.35	0.40
1:WB:56:ARG:HE	1:XB:127:LEU:CD2	2.34	0.40
1:Y:176:LEU:HD21	1:Y:196:TRP:NE1	2.36	0.40
1:YA:168:ILE:HG22	1:YA:215:LEU:CD2	2.51	0.40
1:YA:539:LEU:HD21	1:YA:599:ILE:HD11	2.03	0.40
1:YB:121:LEU:HD23	1:YB:162:ILE:HG22	2.04	0.40
1:ZB:539:LEU:HD12	1:ZB:540:GLN:N	2.37	0.40
1:A:3:THR:HG22	1:A:6:PHE:HA	2.03	0.40
1:AA:570:ASP:OD1	1:AA:570:ASP:C	2.64	0.40
1:AC:529:ILE:HD13	1:AC:583:VAL:HG21	2.02	0.40
1:BA:539:LEU:HD12	1:BA:540:GLN:N	2.36	0.40
1:BA:541:LEU:CD2	1:BA:640:VAL:HG22	2.51	0.40

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:BB:95:ASP:O	1:BB:96:PRO:C	2.64	0.40
1:BB:539:LEU:HD12	1:BB:540:GLN:N	2.37	0.40
1:BB:601:THR:O	1:BB:605:GLY:HA2	2.22	0.40
1:C:65:VAL:HG12	1:C:83:LEU:CD2	2.51	0.40
1:C:90:ILE:HD11	1:C:155:LYS:N	2.37	0.40
1:DA:541:LEU:CD2	1:DA:640:VAL:HG22	2.51	0.40
1:E:230:ARG:HB3	1:E:268:LEU:HD11	2.03	0.40
1:EB:539:LEU:CD1	1:EB:640:VAL:HG13	2.51	0.40
1:F:281:TYR:CG	1:F:366:VAL:HG23	2.56	0.40
1:FA:48:MET:HE2	1:FA:48:MET:HA	2.03	0.40
1:FB:597:ARG:O	1:FB:601:THR:HG23	2.22	0.40
1:G:56:ARG:HE	1:H:127:LEU:CD2	2.33	0.40
1:H:654:LEU:HD22	1:I:534:HIS:CG	2.57	0.40
1:HB:129:PHE:CE1	1:HB:137:VAL:HG11	2.56	0.40
1:IB:143:TRP:HH2	1:IB:157:VAL:HG22	1.85	0.40
1:JB:17:HIS:CB	1:JB:44:LEU:HD23	2.51	0.40
1:LA:56:ARG:HE	1:MA:127:LEU:CD2	2.34	0.40
1:NA:809:ASP:HA	1:NA:812:VAL:HG22	2.03	0.40
1:NB:44:LEU:N	1:NB:44:LEU:HD22	2.35	0.40
1:NB:539:LEU:HD11	1:NB:640:VAL:HG13	2.04	0.40
1:O:57:HIS:O	1:O:58:TYR:HD1	2.04	0.40
1:OA:3:THR:HG22	1:OA:6:PHE:HA	2.03	0.40
1:P:165:ALA:HB2	1:P:205:LEU:HD13	2.02	0.40
1:PB:219:VAL:HG11	1:PB:227:LEU:HD13	2.03	0.40
1:Q:129:PHE:CE1	1:Q:137:VAL:HG11	2.56	0.40
1:QA:48:MET:HE2	1:QA:48:MET:HA	2.02	0.40
1:RA:143:TRP:HH2	1:RA:157:VAL:HG22	1.86	0.40
1:RB:654:LEU:HD22	1:SB:534:HIS:CG	2.57	0.40
1:S:539:LEU:HD21	1:S:599:ILE:HD11	2.04	0.40
1:SA:643:VAL:O	1:SA:643:VAL:HG12	2.21	0.40
1:TA:17:HIS:CB	1:TA:44:LEU:HD23	2.52	0.40
1:TA:56:ARG:HE	1:UA:127:LEU:HD21	1.87	0.40
1:TA:168:ILE:HG22	1:TA:215:LEU:CD2	2.48	0.40
1:TA:541:LEU:CD2	1:TA:640:VAL:HG22	2.52	0.40
1:V:56:ARG:HE	1:W:127:LEU:HD21	1.87	0.40
1:VB:643:VAL:HG12	1:VB:643:VAL:O	2.21	0.40
1:VB:809:ASP:HA	1:VB:812:VAL:HG22	2.03	0.40
1:X:779:LEU:HD21	1:Y:774:ARG:HG2	2.03	0.40
1:XB:539:LEU:CD1	1:XB:640:VAL:HG13	2.51	0.40
1:Y:121:LEU:HD23	1:Y:162:ILE:HG22	2.03	0.40
1:Y:143:TRP:HH2	1:Y:157:VAL:HG22	1.86	0.40

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:YB:176:LEU:HD21	1:YB:196:TRP:NE1	2.35	0.40
1:YB:413:VAL:HG22	1:YB:414:LEU:N	2.37	0.40
1:ZA:47:PRO:C	1:ZA:48:MET:HE2	2.46	0.40
1:ZA:176:LEU:HD21	1:ZA:196:TRP:CE2	2.56	0.40
1:ZB:95:ASP:O	1:ZB:96:PRO:C	2.65	0.40
1:AB:127:LEU:CD2	1:ZA:56:ARG:HE	2.34	0.40
1:BB:523:PHE:CD2	1:BB:568:VAL:HG23	2.56	0.40
1:C:176:LEU:HD21	1:C:196:TRP:NE1	2.35	0.40
1:CA:779:LEU:HD21	1:DA:774:ARG:HG2	2.04	0.40
1:CB:539:LEU:CD1	1:CB:640:VAL:HG13	2.51	0.40
1:D:489:LEU:HD21	1:D:495:PHE:CZ	2.57	0.40
1:DB:597:ARG:O	1:DB:601:THR:HG23	2.21	0.40
1:E:487:VAL:O	1:E:487:VAL:HG13	2.20	0.40
1:F:539:LEU:HD21	1:F:599:ILE:HD11	2.03	0.40
1:GB:541:LEU:CD2	1:GB:640:VAL:HG22	2.51	0.40
1:HA:56:ARG:HE	1:IA:127:LEU:CD2	2.35	0.40
1:HA:95:ASP:O	1:HA:96:PRO:C	2.65	0.40
1:JB:393:VAL:HG22	1:JB:411:ASP:C	2.47	0.40
1:L:489:LEU:HD21	1:L:495:PHE:CZ	2.57	0.40
1:M:48:MET:HE2	1:M:48:MET:HA	2.03	0.40
1:M:165:ALA:HB2	1:M:205:LEU:HD13	2.03	0.40
1:N:159:VAL:O	1:N:159:VAL:HG12	2.20	0.40
1:NA:539:LEU:HD11	1:NA:640:VAL:HG13	2.03	0.40
1:NB:597:ARG:O	1:NB:601:THR:HG23	2.21	0.40
1:PB:779:LEU:HD21	1:QB:774:ARG:HG2	2.03	0.40
1:QA:13:TYR:HA	1:QA:50:MET:CE	2.52	0.40
1:QA:570:ASP:OD1	1:QA:570:ASP:C	2.64	0.40
1:R:44:LEU:HD22	1:R:44:LEU:H	1.86	0.40
1:R:393:VAL:HG22	1:R:411:ASP:C	2.46	0.40
1:SA:15:TYR:CD1	1:SA:15:TYR:C	3.00	0.40
1:SA:538:GLN:HB2	1:SA:646:VAL:HG22	2.02	0.40
1:SA:541:LEU:CD2	1:SA:640:VAL:HG22	2.52	0.40
1:SB:17:HIS:CB	1:SB:44:LEU:HD23	2.51	0.40
1:T:516:LEU:O	1:T:516:LEU:HD23	2.21	0.40
1:T:539:LEU:CD1	1:T:640:VAL:HG13	2.51	0.40
1:UA:165:ALA:HB2	1:UA:205:LEU:HD13	2.03	0.40
1:V:393:VAL:HG22	1:V:411:ASP:C	2.46	0.40
1:V:539:LEU:HD23	1:V:540:GLN:H	1.83	0.40
1:VA:230:ARG:HB3	1:VA:268:LEU:HD11	2.02	0.40
1:W:95:ASP:O	1:W:96:PRO:C	2.64	0.40
1:WA:168:ILE:HG22	1:WA:215:LEU:CD2	2.48	0.40

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:WB:56:ARG:HE	1:XB:127:LEU:HD21	1.87	0.40
1:WB:65:VAL:HG12	1:WB:83:LEU:CD1	2.52	0.40
1:X:539:LEU:HD11	1:X:599:ILE:CD1	2.50	0.40
1:XB:539:LEU:HD11	1:XB:640:VAL:HG13	2.03	0.40
1:A:127:LEU:CD2	1:NA:56:ARG:HE	2.34	0.40
1:A:539:LEU:HD12	1:A:540:GLN:N	2.37	0.40
1:AA:17:HIS:CB	1:AA:44:LEU:HD23	2.52	0.40
1:AB:316:LEU:HD12	1:AB:316:LEU:HA	1.95	0.40
1:AB:570:ASP:OD1	1:AB:570:ASP:C	2.64	0.40
1:AC:95:ASP:O	1:AC:96:PRO:C	2.64	0.40
1:AC:539:LEU:CD1	1:AC:640:VAL:HG13	2.52	0.40
1:B:219:VAL:HG11	1:B:227:LEU:HD13	2.04	0.40
1:CB:539:LEU:HD12	1:CB:540:GLN:N	2.37	0.40
1:CB:654:LEU:HD22	1:DB:534:HIS:CG	2.56	0.40
1:DB:359:ILE:O	1:DB:359:ILE:HG23	2.21	0.40
1:DB:523:PHE:CE2	1:DB:545:TRP:CD1	3.10	0.40
1:EA:56:ARG:HE	1:FA:127:LEU:CD2	2.34	0.40
1:EB:219:VAL:HG11	1:EB:227:LEU:HD13	2.04	0.40
1:FA:121:LEU:HD23	1:FA:162:ILE:HG22	2.03	0.40
1:FB:219:VAL:HG11	1:FB:227:LEU:HD22	2.02	0.40
1:GB:44:LEU:HD22	1:GB:44:LEU:H	1.87	0.40
1:GB:143:TRP:HH2	1:GB:157:VAL:HG22	1.85	0.40
1:H:539:LEU:CD1	1:H:640:VAL:HG13	2.52	0.40
1:HB:17:HIS:CB	1:HB:44:LEU:HD23	2.51	0.40
1:HB:95:ASP:O	1:HB:96:PRO:C	2.65	0.40
1:HB:529:ILE:HD13	1:HB:583:VAL:HG11	2.04	0.40
1:IA:48:MET:HA	1:IA:48:MET:HE2	2.03	0.40
1:IA:539:LEU:HD11	1:IA:541:LEU:HD21	2.04	0.40
1:K:56:ARG:HE	1:L:127:LEU:CD2	2.34	0.40
1:KA:405:THR:HG21	1:LA:393:VAL:O	2.21	0.40
1:KB:17:HIS:CB	1:KB:44:LEU:HD23	2.51	0.40
1:M:56:ARG:HE	1:N:127:LEU:CD2	2.33	0.40
1:MB:73:VAL:HG11	1:MB:81:VAL:HG23	2.02	0.40
1:NA:17:HIS:CB	1:NA:44:LEU:HD23	2.52	0.40
1:NA:601:THR:O	1:NA:605:GLY:HA2	2.22	0.40
1:P:15:TYR:C	1:P:15:TYR:CD1	3.00	0.40
1:Q:168:ILE:HG22	1:Q:215:LEU:CD2	2.49	0.40
1:R:176:LEU:HD21	1:R:196:TRP:NE1	2.35	0.40
1:SA:662:ILE:O	1:SA:666:THR:HG23	2.20	0.40
1:UA:44:LEU:HD22	1:UA:44:LEU:H	1.86	0.40
1:UA:56:ARG:HE	1:VA:127:LEU:CD2	2.34	0.40

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:VA:48:MET:HE2	1:VA:48:MET:HA	2.04	0.40
1:VA:539:LEU:CD1	1:VA:640:VAL:HG13	2.51	0.40
1:VB:17:HIS:CB	1:VB:44:LEU:HD23	2.51	0.40
1:VB:359:ILE:O	1:VB:359:ILE:HG23	2.22	0.40
1:VB:487:VAL:O	1:VB:487:VAL:HG13	2.20	0.40
1:WA:121:LEU:HD23	1:WA:162:ILE:HG22	2.03	0.40
1:XB:15:TYR:CD1	1:XB:15:TYR:C	3.00	0.40
1:Z:538:GLN:HB2	1:Z:646:VAL:HG22	2.02	0.40
1:ZA:73:VAL:HG11	1:ZA:81:VAL:HG23	2.03	0.40
1:ZB:539:LEU:CD1	1:ZB:640:VAL:HG13	2.51	0.40

There are no symmetry-related clashes.

## 5.3 Torsion angles [i](#)

### 5.3.1 Protein backbone [i](#)

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

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

Mol	Chain	Analysed	Favoured	Allowed	Outliers	Percentiles	
1	A	773/893 (87%)	721 (93%)	51 (7%)	1 (0%)	48	78
1	AA	773/893 (87%)	715 (92%)	57 (7%)	1 (0%)	48	78
1	AB	773/893 (87%)	721 (93%)	52 (7%)	0	100	100
1	AC	773/893 (87%)	718 (93%)	54 (7%)	1 (0%)	48	78
1	B	773/893 (87%)	720 (93%)	53 (7%)	0	100	100
1	BA	773/893 (87%)	718 (93%)	55 (7%)	0	100	100
1	BB	773/893 (87%)	716 (93%)	57 (7%)	0	100	100
1	C	773/893 (87%)	720 (93%)	53 (7%)	0	100	100
1	CA	773/893 (87%)	721 (93%)	51 (7%)	1 (0%)	48	78
1	CB	773/893 (87%)	718 (93%)	54 (7%)	1 (0%)	48	78
1	D	773/893 (87%)	720 (93%)	52 (7%)	1 (0%)	48	78
1	DA	773/893 (87%)	719 (93%)	53 (7%)	1 (0%)	48	78

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Mol	Chain	Analysed	Favoured	Allowed	Outliers	Percentiles	
1	DB	773/893 (87%)	722 (93%)	51 (7%)	0	100	100
1	E	773/893 (87%)	719 (93%)	53 (7%)	1 (0%)	48	78
1	EA	773/893 (87%)	723 (94%)	49 (6%)	1 (0%)	48	78
1	EB	773/893 (87%)	716 (93%)	56 (7%)	1 (0%)	48	78
1	F	773/893 (87%)	720 (93%)	53 (7%)	0	100	100
1	FA	773/893 (87%)	719 (93%)	53 (7%)	1 (0%)	48	78
1	FB	773/893 (87%)	717 (93%)	56 (7%)	0	100	100
1	G	773/893 (87%)	720 (93%)	52 (7%)	1 (0%)	48	78
1	GA	773/893 (87%)	723 (94%)	49 (6%)	1 (0%)	48	78
1	GB	773/893 (87%)	719 (93%)	54 (7%)	0	100	100
1	H	773/893 (87%)	719 (93%)	53 (7%)	1 (0%)	48	78
1	HA	773/893 (87%)	722 (93%)	51 (7%)	0	100	100
1	HB	773/893 (87%)	720 (93%)	53 (7%)	0	100	100
1	I	773/893 (87%)	718 (93%)	54 (7%)	1 (0%)	48	78
1	IA	773/893 (87%)	717 (93%)	55 (7%)	1 (0%)	48	78
1	IB	773/893 (87%)	716 (93%)	56 (7%)	1 (0%)	48	78
1	J	773/893 (87%)	718 (93%)	55 (7%)	0	100	100
1	JA	773/893 (87%)	722 (93%)	50 (6%)	1 (0%)	48	78
1	JB	773/893 (87%)	720 (93%)	52 (7%)	1 (0%)	48	78
1	K	773/893 (87%)	719 (93%)	53 (7%)	1 (0%)	48	78
1	KA	773/893 (87%)	716 (93%)	56 (7%)	1 (0%)	48	78
1	KB	773/893 (87%)	719 (93%)	53 (7%)	1 (0%)	48	78
1	L	773/893 (87%)	720 (93%)	53 (7%)	0	100	100
1	LA	773/893 (87%)	719 (93%)	53 (7%)	1 (0%)	48	78
1	LB	773/893 (87%)	720 (93%)	53 (7%)	0	100	100
1	M	773/893 (87%)	721 (93%)	51 (7%)	1 (0%)	48	78
1	MA	773/893 (87%)	719 (93%)	53 (7%)	1 (0%)	48	78
1	MB	773/893 (87%)	720 (93%)	52 (7%)	1 (0%)	48	78
1	N	773/893 (87%)	719 (93%)	53 (7%)	1 (0%)	48	78
1	NA	773/893 (87%)	719 (93%)	53 (7%)	1 (0%)	48	78
1	NB	773/893 (87%)	715 (92%)	57 (7%)	1 (0%)	48	78

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Mol	Chain	Analysed	Favoured	Allowed	Outliers	Percentiles	
1	O	773/893 (87%)	720 (93%)	52 (7%)	1 (0%)	48	78
1	OA	773/893 (87%)	718 (93%)	54 (7%)	1 (0%)	48	78
1	OB	773/893 (87%)	719 (93%)	53 (7%)	1 (0%)	48	78
1	P	773/893 (87%)	716 (93%)	57 (7%)	0	100	100
1	PA	773/893 (87%)	719 (93%)	53 (7%)	1 (0%)	48	78
1	PB	773/893 (87%)	723 (94%)	49 (6%)	1 (0%)	48	78
1	Q	773/893 (87%)	717 (93%)	55 (7%)	1 (0%)	48	78
1	QA	773/893 (87%)	720 (93%)	53 (7%)	0	100	100
1	QB	773/893 (87%)	723 (94%)	49 (6%)	1 (0%)	48	78
1	R	773/893 (87%)	717 (93%)	56 (7%)	0	100	100
1	RA	773/893 (87%)	719 (93%)	53 (7%)	1 (0%)	48	78
1	RB	773/893 (87%)	722 (93%)	50 (6%)	1 (0%)	48	78
1	S	773/893 (87%)	718 (93%)	55 (7%)	0	100	100
1	SA	773/893 (87%)	717 (93%)	56 (7%)	0	100	100
1	SB	773/893 (87%)	718 (93%)	55 (7%)	0	100	100
1	T	773/893 (87%)	719 (93%)	53 (7%)	1 (0%)	48	78
1	TA	773/893 (87%)	718 (93%)	55 (7%)	0	100	100
1	TB	773/893 (87%)	725 (94%)	47 (6%)	1 (0%)	48	78
1	UA	773/893 (87%)	721 (93%)	52 (7%)	0	100	100
1	UB	773/893 (87%)	719 (93%)	54 (7%)	0	100	100
1	V	773/893 (87%)	719 (93%)	53 (7%)	1 (0%)	48	78
1	VA	773/893 (87%)	722 (93%)	50 (6%)	1 (0%)	48	78
1	VB	773/893 (87%)	722 (93%)	50 (6%)	1 (0%)	48	78
1	W	773/893 (87%)	723 (94%)	50 (6%)	0	100	100
1	WA	773/893 (87%)	720 (93%)	52 (7%)	1 (0%)	48	78
1	WB	773/893 (87%)	720 (93%)	52 (7%)	1 (0%)	48	78
1	X	773/893 (87%)	722 (93%)	51 (7%)	0	100	100
1	XA	773/893 (87%)	718 (93%)	55 (7%)	0	100	100
1	XB	773/893 (87%)	719 (93%)	54 (7%)	0	100	100
1	Y	773/893 (87%)	722 (93%)	51 (7%)	0	100	100
1	YA	773/893 (87%)	719 (93%)	53 (7%)	1 (0%)	48	78

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Mol	Chain	Analysed	Favoured	Allowed	Outliers	Percentiles	
1	YB	773/893 (87%)	719 (93%)	53 (7%)	1 (0%)	48	78
1	Z	773/893 (87%)	719 (93%)	53 (7%)	1 (0%)	48	78
1	ZA	773/893 (87%)	723 (94%)	49 (6%)	1 (0%)	48	78
1	ZB	773/893 (87%)	718 (93%)	55 (7%)	0	100	100
All	All	60294/69654 (87%)	56116 (93%)	4129 (7%)	49 (0%)	49	78

All (49) Ramachandran outliers are listed below:

Mol	Chain	Res	Type
1	M	347	GLU
1	CB	347	GLU
1	E	347	GLU
1	LA	347	GLU
1	MA	347	GLU
1	NA	347	GLU
1	QB	347	GLU
1	T	347	GLU
1	TB	347	GLU
1	YB	347	GLU
1	A	347	GLU
1	AA	347	GLU
1	AC	347	GLU
1	CA	347	GLU
1	D	347	GLU
1	DA	347	GLU
1	EA	347	GLU
1	EB	347	GLU
1	FA	347	GLU
1	G	347	GLU
1	GA	347	GLU
1	H	347	GLU
1	I	347	GLU
1	IA	347	GLU
1	IB	347	GLU
1	JA	347	GLU
1	JB	347	GLU
1	K	347	GLU
1	KA	347	GLU
1	KB	347	GLU
1	MB	347	GLU
1	N	347	GLU

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Mol	Chain	Res	Type
1	NB	347	GLU
1	O	347	GLU
1	OA	347	GLU
1	OB	347	GLU
1	PA	347	GLU
1	PB	347	GLU
1	Q	347	GLU
1	RA	347	GLU
1	RB	347	GLU
1	V	347	GLU
1	VA	347	GLU
1	VB	347	GLU
1	WA	347	GLU
1	WB	347	GLU
1	YA	347	GLU
1	Z	347	GLU
1	ZA	347	GLU

### 5.3.2 Protein sidechains ⓘ

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

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

Mol	Chain	Analysed	Rotameric	Outliers	Percentiles	
1	A	666/755 (88%)	656 (98%)	10 (2%)	57	75
1	AA	666/755 (88%)	656 (98%)	10 (2%)	57	75
1	AB	666/755 (88%)	653 (98%)	13 (2%)	48	72
1	AC	666/755 (88%)	654 (98%)	12 (2%)	51	73
1	B	666/755 (88%)	655 (98%)	11 (2%)	53	74
1	BA	666/755 (88%)	652 (98%)	14 (2%)	47	71
1	BB	666/755 (88%)	657 (99%)	9 (1%)	59	76
1	C	666/755 (88%)	654 (98%)	12 (2%)	51	73
1	CA	666/755 (88%)	655 (98%)	11 (2%)	53	74
1	CB	666/755 (88%)	653 (98%)	13 (2%)	48	72
1	D	666/755 (88%)	653 (98%)	13 (2%)	48	72

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Mol	Chain	Analysed	Rotameric	Outliers	Percentiles	
1	DA	666/755 (88%)	653 (98%)	13 (2%)	48	72
1	DB	666/755 (88%)	653 (98%)	13 (2%)	48	72
1	E	666/755 (88%)	657 (99%)	9 (1%)	59	76
1	EA	666/755 (88%)	653 (98%)	13 (2%)	48	72
1	EB	666/755 (88%)	654 (98%)	12 (2%)	51	73
1	F	666/755 (88%)	655 (98%)	11 (2%)	53	74
1	FA	666/755 (88%)	653 (98%)	13 (2%)	48	72
1	FB	666/755 (88%)	654 (98%)	12 (2%)	51	73
1	G	666/755 (88%)	654 (98%)	12 (2%)	51	73
1	GA	666/755 (88%)	651 (98%)	15 (2%)	44	70
1	GB	666/755 (88%)	654 (98%)	12 (2%)	51	73
1	H	666/755 (88%)	655 (98%)	11 (2%)	53	74
1	HA	666/755 (88%)	652 (98%)	14 (2%)	47	71
1	HB	666/755 (88%)	653 (98%)	13 (2%)	48	72
1	I	666/755 (88%)	654 (98%)	12 (2%)	51	73
1	IA	666/755 (88%)	653 (98%)	13 (2%)	48	72
1	IB	666/755 (88%)	653 (98%)	13 (2%)	48	72
1	J	666/755 (88%)	655 (98%)	11 (2%)	53	74
1	JA	666/755 (88%)	656 (98%)	10 (2%)	57	75
1	JB	666/755 (88%)	652 (98%)	14 (2%)	47	71
1	K	666/755 (88%)	657 (99%)	9 (1%)	59	76
1	KA	666/755 (88%)	652 (98%)	14 (2%)	47	71
1	KB	666/755 (88%)	655 (98%)	11 (2%)	53	74
1	L	666/755 (88%)	650 (98%)	16 (2%)	43	69
1	LA	666/755 (88%)	656 (98%)	10 (2%)	57	75
1	LB	666/755 (88%)	655 (98%)	11 (2%)	53	74
1	M	666/755 (88%)	655 (98%)	11 (2%)	53	74
1	MA	666/755 (88%)	652 (98%)	14 (2%)	47	71
1	MB	666/755 (88%)	656 (98%)	10 (2%)	57	75
1	N	666/755 (88%)	655 (98%)	11 (2%)	53	74
1	NA	666/755 (88%)	656 (98%)	10 (2%)	57	75

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Mol	Chain	Analysed	Rotameric	Outliers	Percentiles	
1	NB	666/755 (88%)	653 (98%)	13 (2%)	48	72
1	O	666/755 (88%)	655 (98%)	11 (2%)	53	74
1	OA	666/755 (88%)	659 (99%)	7 (1%)	65	78
1	OB	666/755 (88%)	653 (98%)	13 (2%)	48	72
1	P	666/755 (88%)	654 (98%)	12 (2%)	51	73
1	PA	666/755 (88%)	654 (98%)	12 (2%)	51	73
1	PB	666/755 (88%)	655 (98%)	11 (2%)	53	74
1	Q	666/755 (88%)	654 (98%)	12 (2%)	51	73
1	QA	666/755 (88%)	654 (98%)	12 (2%)	51	73
1	QB	666/755 (88%)	655 (98%)	11 (2%)	53	74
1	R	666/755 (88%)	655 (98%)	11 (2%)	53	74
1	RA	666/755 (88%)	650 (98%)	16 (2%)	43	69
1	RB	666/755 (88%)	654 (98%)	12 (2%)	51	73
1	S	666/755 (88%)	652 (98%)	14 (2%)	47	71
1	SA	666/755 (88%)	656 (98%)	10 (2%)	57	75
1	SB	666/755 (88%)	652 (98%)	14 (2%)	47	71
1	T	666/755 (88%)	652 (98%)	14 (2%)	47	71
1	TA	666/755 (88%)	654 (98%)	12 (2%)	51	73
1	TB	666/755 (88%)	651 (98%)	15 (2%)	44	70
1	UA	666/755 (88%)	656 (98%)	10 (2%)	57	75
1	UB	666/755 (88%)	654 (98%)	12 (2%)	51	73
1	V	666/755 (88%)	652 (98%)	14 (2%)	47	71
1	VA	666/755 (88%)	659 (99%)	7 (1%)	65	78
1	VB	666/755 (88%)	654 (98%)	12 (2%)	51	73
1	W	666/755 (88%)	653 (98%)	13 (2%)	48	72
1	WA	666/755 (88%)	654 (98%)	12 (2%)	51	73
1	WB	666/755 (88%)	655 (98%)	11 (2%)	53	74
1	X	666/755 (88%)	653 (98%)	13 (2%)	48	72
1	XA	666/755 (88%)	654 (98%)	12 (2%)	51	73
1	XB	666/755 (88%)	653 (98%)	13 (2%)	48	72
1	Y	666/755 (88%)	653 (98%)	13 (2%)	48	72

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Mol	Chain	Analysed	Rotameric	Outliers	Percentiles	
1	YA	666/755 (88%)	657 (99%)	9 (1%)	59	76
1	YB	666/755 (88%)	657 (99%)	9 (1%)	59	76
1	Z	666/755 (88%)	653 (98%)	13 (2%)	48	72
1	ZA	666/755 (88%)	650 (98%)	16 (2%)	43	69
1	ZB	666/755 (88%)	652 (98%)	14 (2%)	47	71
All	All	51948/58890 (88%)	51012 (98%)	936 (2%)	51	73

All (936) residues with a non-rotameric sidechain are listed below:

Mol	Chain	Res	Type
1	A	8	ILE
1	A	34	THR
1	A	58	TYR
1	A	63	ASN
1	A	88	LEU
1	A	209	PHE
1	A	283	VAL
1	A	461	ARG
1	A	761	ARG
1	A	807	ILE
1	AA	8	ILE
1	AA	34	THR
1	AA	63	ASN
1	AA	144	LEU
1	AA	209	PHE
1	AA	665	THR
1	AA	669	GLN
1	AA	728	SER
1	AA	761	ARG
1	AA	807	ILE
1	AB	8	ILE
1	AB	34	THR
1	AB	43	VAL
1	AB	63	ASN
1	AB	88	LEU
1	AB	144	LEU
1	AB	209	PHE
1	AB	518	LEU
1	AB	538	GLN
1	AB	637	SER

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Mol	Chain	Res	Type
1	AB	669	GLN
1	AB	728	SER
1	AB	807	ILE
1	AC	8	ILE
1	AC	34	THR
1	AC	43	VAL
1	AC	58	TYR
1	AC	63	ASN
1	AC	88	LEU
1	AC	209	PHE
1	AC	283	VAL
1	AC	461	ARG
1	AC	523	PHE
1	AC	669	GLN
1	AC	728	SER
1	B	8	ILE
1	B	34	THR
1	B	63	ASN
1	B	88	LEU
1	B	209	PHE
1	B	283	VAL
1	B	461	ARG
1	B	637	SER
1	B	665	THR
1	B	728	SER
1	B	807	ILE
1	BA	8	ILE
1	BA	20	ASP
1	BA	34	THR
1	BA	43	VAL
1	BA	63	ASN
1	BA	88	LEU
1	BA	144	LEU
1	BA	209	PHE
1	BA	283	VAL
1	BA	461	ARG
1	BA	637	SER
1	BA	665	THR
1	BA	728	SER
1	BA	807	ILE
1	BB	8	ILE
1	BB	34	THR

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Mol	Chain	Res	Type
1	BB	58	TYR
1	BB	63	ASN
1	BB	209	PHE
1	BB	283	VAL
1	BB	637	SER
1	BB	728	SER
1	BB	807	ILE
1	C	8	ILE
1	C	20	ASP
1	C	34	THR
1	C	63	ASN
1	C	88	LEU
1	C	144	LEU
1	C	209	PHE
1	C	516	LEU
1	C	523	PHE
1	C	637	SER
1	C	728	SER
1	C	807	ILE
1	CA	8	ILE
1	CA	34	THR
1	CA	58	TYR
1	CA	63	ASN
1	CA	88	LEU
1	CA	209	PHE
1	CA	283	VAL
1	CA	426	LEU
1	CA	461	ARG
1	CA	669	GLN
1	CA	728	SER
1	CB	8	ILE
1	CB	34	THR
1	CB	63	ASN
1	CB	88	LEU
1	CB	144	LEU
1	CB	209	PHE
1	CB	283	VAL
1	CB	461	ARG
1	CB	516	LEU
1	CB	538	GLN
1	CB	669	GLN
1	CB	728	SER

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Mol	Chain	Res	Type
1	CB	807	ILE
1	D	8	ILE
1	D	34	THR
1	D	58	TYR
1	D	63	ASN
1	D	88	LEU
1	D	144	LEU
1	D	209	PHE
1	D	283	VAL
1	D	516	LEU
1	D	523	PHE
1	D	637	SER
1	D	728	SER
1	D	807	ILE
1	DA	8	ILE
1	DA	34	THR
1	DA	63	ASN
1	DA	88	LEU
1	DA	127	LEU
1	DA	209	PHE
1	DA	461	ARG
1	DA	516	LEU
1	DA	523	PHE
1	DA	637	SER
1	DA	665	THR
1	DA	728	SER
1	DA	807	ILE
1	DB	8	ILE
1	DB	20	ASP
1	DB	34	THR
1	DB	58	TYR
1	DB	88	LEU
1	DB	144	LEU
1	DB	209	PHE
1	DB	283	VAL
1	DB	461	ARG
1	DB	665	THR
1	DB	728	SER
1	DB	761	ARG
1	DB	807	ILE
1	E	8	ILE
1	E	34	THR

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Mol	Chain	Res	Type
1	E	63	ASN
1	E	88	LEU
1	E	144	LEU
1	E	209	PHE
1	E	665	THR
1	E	728	SER
1	E	807	ILE
1	EA	8	ILE
1	EA	20	ASP
1	EA	34	THR
1	EA	58	TYR
1	EA	63	ASN
1	EA	88	LEU
1	EA	144	LEU
1	EA	209	PHE
1	EA	283	VAL
1	EA	461	ARG
1	EA	665	THR
1	EA	669	GLN
1	EA	728	SER
1	EB	8	ILE
1	EB	34	THR
1	EB	58	TYR
1	EB	63	ASN
1	EB	88	LEU
1	EB	144	LEU
1	EB	209	PHE
1	EB	283	VAL
1	EB	523	PHE
1	EB	637	SER
1	EB	728	SER
1	EB	807	ILE
1	F	8	ILE
1	F	34	THR
1	F	58	TYR
1	F	63	ASN
1	F	144	LEU
1	F	209	PHE
1	F	283	VAL
1	F	461	ARG
1	F	665	THR
1	F	728	SER

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Mol	Chain	Res	Type
1	F	761	ARG
1	FA	8	ILE
1	FA	34	THR
1	FA	63	ASN
1	FA	88	LEU
1	FA	144	LEU
1	FA	209	PHE
1	FA	516	LEU
1	FA	518	LEU
1	FA	523	PHE
1	FA	637	SER
1	FA	728	SER
1	FA	761	ARG
1	FA	807	ILE
1	FB	8	ILE
1	FB	20	ASP
1	FB	34	THR
1	FB	63	ASN
1	FB	88	LEU
1	FB	144	LEU
1	FB	209	PHE
1	FB	283	VAL
1	FB	669	GLN
1	FB	728	SER
1	FB	798	MET
1	FB	807	ILE
1	G	8	ILE
1	G	34	THR
1	G	43	VAL
1	G	63	ASN
1	G	88	LEU
1	G	144	LEU
1	G	209	PHE
1	G	283	VAL
1	G	665	THR
1	G	728	SER
1	G	761	ARG
1	G	807	ILE
1	GA	8	ILE
1	GA	20	ASP
1	GA	34	THR
1	GA	43	VAL

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Mol	Chain	Res	Type
1	GA	58	TYR
1	GA	63	ASN
1	GA	88	LEU
1	GA	144	LEU
1	GA	209	PHE
1	GA	283	VAL
1	GA	516	LEU
1	GA	523	PHE
1	GA	538	GLN
1	GA	728	SER
1	GA	807	ILE
1	GB	8	ILE
1	GB	34	THR
1	GB	58	TYR
1	GB	63	ASN
1	GB	88	LEU
1	GB	209	PHE
1	GB	283	VAL
1	GB	461	ARG
1	GB	538	GLN
1	GB	669	GLN
1	GB	728	SER
1	GB	807	ILE
1	H	8	ILE
1	H	34	THR
1	H	58	TYR
1	H	63	ASN
1	H	88	LEU
1	H	209	PHE
1	H	355	ASP
1	H	461	ARG
1	H	538	GLN
1	H	728	SER
1	H	807	ILE
1	HA	8	ILE
1	HA	20	ASP
1	HA	34	THR
1	HA	63	ASN
1	HA	88	LEU
1	HA	209	PHE
1	HA	316	LEU
1	HA	461	ARG

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Mol	Chain	Res	Type
1	HA	516	LEU
1	HA	538	GLN
1	HA	669	GLN
1	HA	728	SER
1	HA	761	ARG
1	HA	807	ILE
1	HB	8	ILE
1	HB	58	TYR
1	HB	63	ASN
1	HB	88	LEU
1	HB	144	LEU
1	HB	209	PHE
1	HB	283	VAL
1	HB	426	LEU
1	HB	637	SER
1	HB	665	THR
1	HB	669	GLN
1	HB	728	SER
1	HB	807	ILE
1	I	8	ILE
1	I	34	THR
1	I	63	ASN
1	I	88	LEU
1	I	209	PHE
1	I	283	VAL
1	I	461	ARG
1	I	538	GLN
1	I	665	THR
1	I	669	GLN
1	I	728	SER
1	I	761	ARG
1	IA	8	ILE
1	IA	34	THR
1	IA	63	ASN
1	IA	88	LEU
1	IA	144	LEU
1	IA	209	PHE
1	IA	283	VAL
1	IA	461	ARG
1	IA	516	LEU
1	IA	538	GLN
1	IA	598	ILE

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Mol	Chain	Res	Type
1	IA	665	THR
1	IA	728	SER
1	IB	8	ILE
1	IB	34	THR
1	IB	58	TYR
1	IB	63	ASN
1	IB	88	LEU
1	IB	209	PHE
1	IB	426	LEU
1	IB	461	ARG
1	IB	598	ILE
1	IB	665	THR
1	IB	669	GLN
1	IB	728	SER
1	IB	807	ILE
1	J	8	ILE
1	J	34	THR
1	J	58	TYR
1	J	63	ASN
1	J	88	LEU
1	J	209	PHE
1	J	523	PHE
1	J	637	SER
1	J	665	THR
1	J	728	SER
1	J	807	ILE
1	JA	8	ILE
1	JA	34	THR
1	JA	209	PHE
1	JA	426	LEU
1	JA	516	LEU
1	JA	518	LEU
1	JA	538	GLN
1	JA	637	SER
1	JA	728	SER
1	JA	807	ILE
1	JB	8	ILE
1	JB	20	ASP
1	JB	34	THR
1	JB	58	TYR
1	JB	63	ASN
1	JB	88	LEU

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Mol	Chain	Res	Type
1	JB	144	LEU
1	JB	209	PHE
1	JB	283	VAL
1	JB	461	ARG
1	JB	516	LEU
1	JB	637	SER
1	JB	728	SER
1	JB	807	ILE
1	K	8	ILE
1	K	34	THR
1	K	58	TYR
1	K	63	ASN
1	K	88	LEU
1	K	144	LEU
1	K	209	PHE
1	K	523	PHE
1	K	807	ILE
1	KA	8	ILE
1	KA	34	THR
1	KA	58	TYR
1	KA	63	ASN
1	KA	88	LEU
1	KA	144	LEU
1	KA	209	PHE
1	KA	283	VAL
1	KA	306	LYS
1	KA	523	PHE
1	KA	665	THR
1	KA	728	SER
1	KA	761	ARG
1	KA	807	ILE
1	KB	8	ILE
1	KB	34	THR
1	KB	43	VAL
1	KB	63	ASN
1	KB	209	PHE
1	KB	283	VAL
1	KB	461	ARG
1	KB	516	LEU
1	KB	637	SER
1	KB	728	SER
1	KB	807	ILE

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Mol	Chain	Res	Type
1	L	8	ILE
1	L	20	ASP
1	L	34	THR
1	L	58	TYR
1	L	63	ASN
1	L	88	LEU
1	L	144	LEU
1	L	209	PHE
1	L	283	VAL
1	L	523	PHE
1	L	538	GLN
1	L	598	ILE
1	L	637	SER
1	L	665	THR
1	L	728	SER
1	L	807	ILE
1	LA	8	ILE
1	LA	34	THR
1	LA	58	TYR
1	LA	63	ASN
1	LA	88	LEU
1	LA	209	PHE
1	LA	637	SER
1	LA	728	SER
1	LA	761	ARG
1	LA	807	ILE
1	LB	8	ILE
1	LB	34	THR
1	LB	58	TYR
1	LB	63	ASN
1	LB	88	LEU
1	LB	209	PHE
1	LB	352	GLN
1	LB	516	LEU
1	LB	637	SER
1	LB	728	SER
1	LB	807	ILE
1	M	8	ILE
1	M	34	THR
1	M	63	ASN
1	M	88	LEU
1	M	209	PHE

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Mol	Chain	Res	Type
1	M	518	LEU
1	M	538	GLN
1	M	637	SER
1	M	669	GLN
1	M	728	SER
1	M	807	ILE
1	MA	8	ILE
1	MA	20	ASP
1	MA	34	THR
1	MA	58	TYR
1	MA	88	LEU
1	MA	144	LEU
1	MA	209	PHE
1	MA	283	VAL
1	MA	516	LEU
1	MA	523	PHE
1	MA	665	THR
1	MA	728	SER
1	MA	761	ARG
1	MA	807	ILE
1	MB	8	ILE
1	MB	34	THR
1	MB	63	ASN
1	MB	88	LEU
1	MB	144	LEU
1	MB	209	PHE
1	MB	461	ARG
1	MB	523	PHE
1	MB	665	THR
1	MB	728	SER
1	N	8	ILE
1	N	34	THR
1	N	63	ASN
1	N	88	LEU
1	N	209	PHE
1	N	283	VAL
1	N	426	LEU
1	N	461	ARG
1	N	637	SER
1	N	728	SER
1	N	807	ILE
1	NA	8	ILE

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Mol	Chain	Res	Type
1	NA	34	THR
1	NA	58	TYR
1	NA	63	ASN
1	NA	88	LEU
1	NA	209	PHE
1	NA	461	ARG
1	NA	523	PHE
1	NA	669	GLN
1	NA	728	SER
1	NB	8	ILE
1	NB	20	ASP
1	NB	34	THR
1	NB	63	ASN
1	NB	88	LEU
1	NB	144	LEU
1	NB	209	PHE
1	NB	283	VAL
1	NB	665	THR
1	NB	669	GLN
1	NB	728	SER
1	NB	761	ARG
1	NB	807	ILE
1	O	8	ILE
1	O	34	THR
1	O	58	TYR
1	O	88	LEU
1	O	209	PHE
1	O	283	VAL
1	O	461	ARG
1	O	516	LEU
1	O	669	GLN
1	O	728	SER
1	O	807	ILE
1	OA	8	ILE
1	OA	34	THR
1	OA	63	ASN
1	OA	209	PHE
1	OA	283	VAL
1	OA	461	ARG
1	OA	807	ILE
1	OB	8	ILE
1	OB	20	ASP

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Mol	Chain	Res	Type
1	OB	34	THR
1	OB	43	VAL
1	OB	63	ASN
1	OB	88	LEU
1	OB	209	PHE
1	OB	283	VAL
1	OB	461	ARG
1	OB	637	SER
1	OB	665	THR
1	OB	728	SER
1	OB	807	ILE
1	P	8	ILE
1	P	34	THR
1	P	58	TYR
1	P	63	ASN
1	P	88	LEU
1	P	144	LEU
1	P	209	PHE
1	P	461	ARG
1	P	665	THR
1	P	728	SER
1	P	761	ARG
1	P	807	ILE
1	PA	8	ILE
1	PA	34	THR
1	PA	63	ASN
1	PA	88	LEU
1	PA	209	PHE
1	PA	283	VAL
1	PA	426	LEU
1	PA	461	ARG
1	PA	637	SER
1	PA	665	THR
1	PA	728	SER
1	PA	807	ILE
1	PB	8	ILE
1	PB	34	THR
1	PB	58	TYR
1	PB	63	ASN
1	PB	88	LEU
1	PB	209	PHE
1	PB	283	VAL

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Mol	Chain	Res	Type
1	PB	426	LEU
1	PB	461	ARG
1	PB	669	GLN
1	PB	728	SER
1	Q	8	ILE
1	Q	34	THR
1	Q	58	TYR
1	Q	63	ASN
1	Q	88	LEU
1	Q	144	LEU
1	Q	209	PHE
1	Q	283	VAL
1	Q	523	PHE
1	Q	637	SER
1	Q	728	SER
1	Q	807	ILE
1	QA	8	ILE
1	QA	34	THR
1	QA	63	ASN
1	QA	88	LEU
1	QA	144	LEU
1	QA	209	PHE
1	QA	426	LEU
1	QA	516	LEU
1	QA	523	PHE
1	QA	637	SER
1	QA	728	SER
1	QA	807	ILE
1	QB	8	ILE
1	QB	34	THR
1	QB	63	ASN
1	QB	88	LEU
1	QB	144	LEU
1	QB	209	PHE
1	QB	461	ARG
1	QB	523	PHE
1	QB	637	SER
1	QB	728	SER
1	QB	807	ILE
1	R	8	ILE
1	R	34	THR
1	R	63	ASN

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Mol	Chain	Res	Type
1	R	88	LEU
1	R	144	LEU
1	R	209	PHE
1	R	283	VAL
1	R	669	GLN
1	R	728	SER
1	R	798	MET
1	R	807	ILE
1	RA	8	ILE
1	RA	34	THR
1	RA	43	VAL
1	RA	58	TYR
1	RA	63	ASN
1	RA	88	LEU
1	RA	144	LEU
1	RA	209	PHE
1	RA	283	VAL
1	RA	426	LEU
1	RA	516	LEU
1	RA	518	LEU
1	RA	523	PHE
1	RA	637	SER
1	RA	728	SER
1	RA	807	ILE
1	RB	8	ILE
1	RB	34	THR
1	RB	58	TYR
1	RB	63	ASN
1	RB	88	LEU
1	RB	144	LEU
1	RB	209	PHE
1	RB	283	VAL
1	RB	426	LEU
1	RB	461	ARG
1	RB	665	THR
1	RB	728	SER
1	S	8	ILE
1	S	34	THR
1	S	58	TYR
1	S	63	ASN
1	S	88	LEU
1	S	205	LEU

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Mol	Chain	Res	Type
1	S	209	PHE
1	S	283	VAL
1	S	461	ARG
1	S	538	GLN
1	S	665	THR
1	S	669	GLN
1	S	728	SER
1	S	807	ILE
1	SA	8	ILE
1	SA	20	ASP
1	SA	34	THR
1	SA	58	TYR
1	SA	63	ASN
1	SA	144	LEU
1	SA	209	PHE
1	SA	665	THR
1	SA	728	SER
1	SA	807	ILE
1	SB	8	ILE
1	SB	34	THR
1	SB	58	TYR
1	SB	63	ASN
1	SB	88	LEU
1	SB	144	LEU
1	SB	209	PHE
1	SB	516	LEU
1	SB	518	LEU
1	SB	523	PHE
1	SB	637	SER
1	SB	728	SER
1	SB	761	ARG
1	SB	807	ILE
1	T	8	ILE
1	T	34	THR
1	T	58	TYR
1	T	63	ASN
1	T	88	LEU
1	T	144	LEU
1	T	209	PHE
1	T	283	VAL
1	T	426	LEU
1	T	637	SER

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Mol	Chain	Res	Type
1	T	665	THR
1	T	669	GLN
1	T	728	SER
1	T	807	ILE
1	TA	8	ILE
1	TA	20	ASP
1	TA	34	THR
1	TA	58	TYR
1	TA	63	ASN
1	TA	144	LEU
1	TA	209	PHE
1	TA	283	VAL
1	TA	461	ARG
1	TA	665	THR
1	TA	728	SER
1	TA	761	ARG
1	TB	8	ILE
1	TB	20	ASP
1	TB	34	THR
1	TB	43	VAL
1	TB	58	TYR
1	TB	63	ASN
1	TB	88	LEU
1	TB	144	LEU
1	TB	209	PHE
1	TB	283	VAL
1	TB	516	LEU
1	TB	523	PHE
1	TB	538	GLN
1	TB	728	SER
1	TB	807	ILE
1	UA	8	ILE
1	UA	34	THR
1	UA	63	ASN
1	UA	88	LEU
1	UA	209	PHE
1	UA	283	VAL
1	UA	665	THR
1	UA	728	SER
1	UA	761	ARG
1	UA	807	ILE
1	UB	8	ILE

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Mol	Chain	Res	Type
1	UB	20	ASP
1	UB	34	THR
1	UB	63	ASN
1	UB	88	LEU
1	UB	144	LEU
1	UB	209	PHE
1	UB	461	ARG
1	UB	538	GLN
1	UB	669	GLN
1	UB	728	SER
1	UB	807	ILE
1	V	8	ILE
1	V	20	ASP
1	V	34	THR
1	V	63	ASN
1	V	205	LEU
1	V	209	PHE
1	V	461	ARG
1	V	523	PHE
1	V	538	GLN
1	V	549	VAL
1	V	598	ILE
1	V	665	THR
1	V	728	SER
1	V	807	ILE
1	VA	8	ILE
1	VA	34	THR
1	VA	63	ASN
1	VA	88	LEU
1	VA	209	PHE
1	VA	461	ARG
1	VA	807	ILE
1	VB	8	ILE
1	VB	34	THR
1	VB	63	ASN
1	VB	88	LEU
1	VB	144	LEU
1	VB	209	PHE
1	VB	283	VAL
1	VB	516	LEU
1	VB	538	GLN
1	VB	598	ILE

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Mol	Chain	Res	Type
1	VB	665	THR
1	VB	728	SER
1	W	8	ILE
1	W	34	THR
1	W	58	TYR
1	W	63	ASN
1	W	88	LEU
1	W	144	LEU
1	W	209	PHE
1	W	283	VAL
1	W	461	ARG
1	W	516	LEU
1	W	637	SER
1	W	728	SER
1	W	807	ILE
1	WA	8	ILE
1	WA	34	THR
1	WA	43	VAL
1	WA	63	ASN
1	WA	88	LEU
1	WA	209	PHE
1	WA	283	VAL
1	WA	461	ARG
1	WA	538	GLN
1	WA	665	THR
1	WA	669	GLN
1	WA	728	SER
1	WB	8	ILE
1	WB	34	THR
1	WB	58	TYR
1	WB	63	ASN
1	WB	209	PHE
1	WB	516	LEU
1	WB	518	LEU
1	WB	538	GLN
1	WB	637	SER
1	WB	728	SER
1	WB	807	ILE
1	X	8	ILE
1	X	20	ASP
1	X	34	THR
1	X	63	ASN

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Mol	Chain	Res	Type
1	X	88	LEU
1	X	144	LEU
1	X	209	PHE
1	X	283	VAL
1	X	461	ARG
1	X	516	LEU
1	X	637	SER
1	X	728	SER
1	X	807	ILE
1	XA	8	ILE
1	XA	34	THR
1	XA	43	VAL
1	XA	58	TYR
1	XA	63	ASN
1	XA	88	LEU
1	XA	209	PHE
1	XA	523	PHE
1	XA	637	SER
1	XA	665	THR
1	XA	669	GLN
1	XA	728	SER
1	XB	8	ILE
1	XB	20	ASP
1	XB	34	THR
1	XB	58	TYR
1	XB	88	LEU
1	XB	144	LEU
1	XB	209	PHE
1	XB	283	VAL
1	XB	523	PHE
1	XB	665	THR
1	XB	728	SER
1	XB	761	ARG
1	XB	807	ILE
1	Y	8	ILE
1	Y	34	THR
1	Y	58	TYR
1	Y	63	ASN
1	Y	88	LEU
1	Y	209	PHE
1	Y	352	GLN
1	Y	461	ARG

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Mol	Chain	Res	Type
1	Y	516	LEU
1	Y	538	GLN
1	Y	669	GLN
1	Y	728	SER
1	Y	807	ILE
1	YA	8	ILE
1	YA	34	THR
1	YA	58	TYR
1	YA	63	ASN
1	YA	88	LEU
1	YA	144	LEU
1	YA	209	PHE
1	YA	523	PHE
1	YA	807	ILE
1	YB	8	ILE
1	YB	20	ASP
1	YB	34	THR
1	YB	63	ASN
1	YB	88	LEU
1	YB	209	PHE
1	YB	637	SER
1	YB	728	SER
1	YB	807	ILE
1	Z	8	ILE
1	Z	20	ASP
1	Z	34	THR
1	Z	63	ASN
1	Z	88	LEU
1	Z	144	LEU
1	Z	209	PHE
1	Z	461	ARG
1	Z	523	PHE
1	Z	598	ILE
1	Z	665	THR
1	Z	728	SER
1	Z	807	ILE
1	ZA	8	ILE
1	ZA	20	ASP
1	ZA	34	THR
1	ZA	58	TYR
1	ZA	63	ASN
1	ZA	88	LEU

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Mol	Chain	Res	Type
1	ZA	144	LEU
1	ZA	209	PHE
1	ZA	283	VAL
1	ZA	461	ARG
1	ZA	523	PHE
1	ZA	538	GLN
1	ZA	598	ILE
1	ZA	665	THR
1	ZA	728	SER
1	ZA	807	ILE
1	ZB	8	ILE
1	ZB	34	THR
1	ZB	43	VAL
1	ZB	58	TYR
1	ZB	63	ASN
1	ZB	88	LEU
1	ZB	209	PHE
1	ZB	283	VAL
1	ZB	322	ASP
1	ZB	516	LEU
1	ZB	523	PHE
1	ZB	665	THR
1	ZB	728	SER
1	ZB	807	ILE

Sometimes sidechains can be flipped to improve hydrogen bonding and reduce clashes. All (386) such sidechains are listed below:

Mol	Chain	Res	Type
1	A	85	HIS
1	A	118	ASN
1	A	329	GLN
1	A	378	GLN
1	A	592	HIS
1	A	630	GLN
1	A	797	GLN
1	AA	329	GLN
1	AA	592	HIS
1	AA	630	GLN
1	AA	797	GLN
1	AB	85	HIS
1	AB	118	ASN
1	AB	329	GLN

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Mol	Chain	Res	Type
1	AB	351	HIS
1	AB	534	HIS
1	AB	630	GLN
1	AC	118	ASN
1	AC	329	GLN
1	AC	592	HIS
1	AC	630	GLN
1	B	85	HIS
1	B	118	ASN
1	B	329	GLN
1	BA	85	HIS
1	BA	118	ASN
1	BA	329	GLN
1	BA	464	HIS
1	BA	534	HIS
1	BA	592	HIS
1	BA	630	GLN
1	BA	797	GLN
1	BB	85	HIS
1	BB	118	ASN
1	BB	329	GLN
1	BB	464	HIS
1	BB	592	HIS
1	BB	630	GLN
1	C	329	GLN
1	C	546	HIS
1	C	592	HIS
1	C	797	GLN
1	CA	85	HIS
1	CA	118	ASN
1	CA	329	GLN
1	CA	592	HIS
1	CA	630	GLN
1	CA	797	GLN
1	CB	85	HIS
1	CB	329	GLN
1	CB	630	GLN
1	CB	797	GLN
1	D	85	HIS
1	D	118	ASN
1	D	329	GLN
1	D	464	HIS

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Mol	Chain	Res	Type
1	D	534	HIS
1	D	592	HIS
1	D	630	GLN
1	D	797	GLN
1	DA	85	HIS
1	DA	329	GLN
1	DA	592	HIS
1	DA	797	GLN
1	DB	85	HIS
1	DB	630	GLN
1	DB	749	GLN
1	E	118	ASN
1	E	329	GLN
1	E	534	HIS
1	E	592	HIS
1	E	630	GLN
1	EA	85	HIS
1	EA	329	GLN
1	EA	464	HIS
1	EA	534	HIS
1	EA	630	GLN
1	EA	797	GLN
1	EB	85	HIS
1	EB	329	GLN
1	EB	592	HIS
1	EB	630	GLN
1	EB	797	GLN
1	F	118	ASN
1	F	329	GLN
1	F	534	HIS
1	F	630	GLN
1	FA	85	HIS
1	FA	329	GLN
1	FA	592	HIS
1	FA	630	GLN
1	FA	797	GLN
1	FB	85	HIS
1	FB	118	ASN
1	FB	329	GLN
1	FB	592	HIS
1	FB	630	GLN
1	FB	797	GLN

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Mol	Chain	Res	Type
1	G	118	ASN
1	G	329	GLN
1	G	630	GLN
1	GA	85	HIS
1	GA	118	ASN
1	GA	329	GLN
1	GA	534	HIS
1	GA	592	HIS
1	GA	630	GLN
1	GA	797	GLN
1	GB	85	HIS
1	GB	329	GLN
1	GB	378	GLN
1	GB	534	HIS
1	GB	630	GLN
1	H	118	ASN
1	H	329	GLN
1	H	592	HIS
1	H	630	GLN
1	HA	118	ASN
1	HA	329	GLN
1	HA	592	HIS
1	HA	630	GLN
1	HA	797	GLN
1	HB	118	ASN
1	HB	329	GLN
1	HB	630	GLN
1	HB	797	GLN
1	I	118	ASN
1	I	329	GLN
1	I	592	HIS
1	IA	85	HIS
1	IA	118	ASN
1	IA	329	GLN
1	IA	464	HIS
1	IA	534	HIS
1	IA	592	HIS
1	IA	630	GLN
1	IA	797	GLN
1	IB	118	ASN
1	IB	329	GLN
1	IB	464	HIS

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Mol	Chain	Res	Type
1	IB	592	HIS
1	IB	630	GLN
1	J	85	HIS
1	J	118	ASN
1	J	329	GLN
1	J	592	HIS
1	J	630	GLN
1	JA	85	HIS
1	JA	118	ASN
1	JA	329	GLN
1	JA	534	HIS
1	JA	630	GLN
1	JA	797	GLN
1	JB	85	HIS
1	JB	118	ASN
1	JB	329	GLN
1	JB	534	HIS
1	JB	592	HIS
1	JB	630	GLN
1	K	85	HIS
1	K	118	ASN
1	K	329	GLN
1	K	464	HIS
1	K	592	HIS
1	KA	329	GLN
1	KA	546	HIS
1	KA	592	HIS
1	KA	749	GLN
1	KA	797	GLN
1	KB	85	HIS
1	KB	329	GLN
1	KB	592	HIS
1	KB	630	GLN
1	KB	797	GLN
1	L	85	HIS
1	L	329	GLN
1	L	592	HIS
1	LA	85	HIS
1	LA	118	ASN
1	LA	329	GLN
1	LA	378	GLN
1	LA	464	HIS

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Mol	Chain	Res	Type
1	LA	592	HIS
1	LA	749	GLN
1	LB	85	HIS
1	LB	118	ASN
1	LB	329	GLN
1	LB	534	HIS
1	LB	546	HIS
1	LB	592	HIS
1	LB	630	GLN
1	LB	797	GLN
1	M	85	HIS
1	M	329	GLN
1	M	534	HIS
1	M	592	HIS
1	M	630	GLN
1	MA	329	GLN
1	MA	592	HIS
1	MB	118	ASN
1	MB	329	GLN
1	MB	378	GLN
1	MB	534	HIS
1	MB	592	HIS
1	MB	630	GLN
1	MB	797	GLN
1	N	118	ASN
1	N	329	GLN
1	N	464	HIS
1	N	592	HIS
1	N	630	GLN
1	NA	85	HIS
1	NA	118	ASN
1	NA	329	GLN
1	NA	592	HIS
1	NA	630	GLN
1	NB	118	ASN
1	NB	329	GLN
1	NB	546	HIS
1	NB	592	HIS
1	NB	630	GLN
1	O	85	HIS
1	O	118	ASN
1	O	329	GLN

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Mol	Chain	Res	Type
1	O	534	HIS
1	O	592	HIS
1	O	630	GLN
1	O	797	GLN
1	OA	118	ASN
1	OA	329	GLN
1	OA	534	HIS
1	OA	592	HIS
1	OA	630	GLN
1	OA	797	GLN
1	OB	118	ASN
1	OB	329	GLN
1	OB	534	HIS
1	OB	592	HIS
1	OB	630	GLN
1	P	85	HIS
1	P	329	GLN
1	P	534	HIS
1	P	592	HIS
1	P	630	GLN
1	PA	85	HIS
1	PA	329	GLN
1	PA	630	GLN
1	PA	797	GLN
1	PB	118	ASN
1	PB	329	GLN
1	PB	592	HIS
1	PB	797	GLN
1	Q	329	GLN
1	Q	592	HIS
1	Q	630	GLN
1	Q	797	GLN
1	QA	329	GLN
1	QA	592	HIS
1	QA	630	GLN
1	QA	797	GLN
1	QB	118	ASN
1	QB	329	GLN
1	QB	592	HIS
1	QB	797	GLN
1	R	118	ASN
1	R	329	GLN

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Mol	Chain	Res	Type
1	R	797	GLN
1	RA	63	ASN
1	RA	118	ASN
1	RA	329	GLN
1	RA	464	HIS
1	RA	534	HIS
1	RA	592	HIS
1	RA	630	GLN
1	RA	797	GLN
1	RB	85	HIS
1	RB	118	ASN
1	RB	329	GLN
1	RB	534	HIS
1	RB	592	HIS
1	RB	630	GLN
1	S	85	HIS
1	S	329	GLN
1	S	378	GLN
1	S	534	HIS
1	S	592	HIS
1	S	630	GLN
1	SA	118	ASN
1	SA	329	GLN
1	SA	534	HIS
1	SA	630	GLN
1	SA	797	GLN
1	SB	118	ASN
1	SB	329	GLN
1	SB	592	HIS
1	SB	797	GLN
1	T	118	ASN
1	T	329	GLN
1	T	534	HIS
1	T	797	GLN
1	TA	85	HIS
1	TA	118	ASN
1	TA	329	GLN
1	TA	464	HIS
1	TB	85	HIS
1	TB	118	ASN
1	TB	329	GLN
1	TB	630	GLN

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Mol	Chain	Res	Type
1	UA	118	ASN
1	UA	329	GLN
1	UB	118	ASN
1	UB	329	GLN
1	UB	464	HIS
1	UB	592	HIS
1	UB	630	GLN
1	V	118	ASN
1	V	329	GLN
1	V	630	GLN
1	VA	85	HIS
1	VA	118	ASN
1	VA	329	GLN
1	VA	592	HIS
1	VA	797	GLN
1	VB	118	ASN
1	VB	329	GLN
1	VB	534	HIS
1	VB	592	HIS
1	VB	630	GLN
1	W	85	HIS
1	W	118	ASN
1	W	329	GLN
1	W	534	HIS
1	W	592	HIS
1	W	797	GLN
1	WA	85	HIS
1	WA	329	GLN
1	WA	534	HIS
1	WB	85	HIS
1	WB	118	ASN
1	WB	329	GLN
1	WB	534	HIS
1	WB	592	HIS
1	X	329	GLN
1	X	592	HIS
1	X	797	GLN
1	XA	85	HIS
1	XA	118	ASN
1	XA	329	GLN
1	XA	630	GLN
1	XB	329	GLN

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Mol	Chain	Res	Type
1	XB	534	HIS
1	XB	546	HIS
1	XB	592	HIS
1	XB	749	GLN
1	XB	797	GLN
1	Y	329	GLN
1	Y	351	HIS
1	Y	534	HIS
1	Y	592	HIS
1	Y	797	GLN
1	YA	85	HIS
1	YA	329	GLN
1	YA	378	GLN
1	YA	464	HIS
1	YA	592	HIS
1	YA	630	GLN
1	YA	797	GLN
1	YB	118	ASN
1	YB	329	GLN
1	YB	378	GLN
1	YB	464	HIS
1	YB	630	GLN
1	YB	749	GLN
1	Z	118	ASN
1	Z	329	GLN
1	Z	630	GLN
1	Z	797	GLN
1	ZA	85	HIS
1	ZA	118	ASN
1	ZA	329	GLN
1	ZA	592	HIS
1	ZA	797	GLN
1	ZB	85	HIS
1	ZB	329	GLN
1	ZB	592	HIS
1	ZB	630	GLN

### 5.3.3 RNA ⓘ

There are no RNA molecules in this entry.

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

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

## 5.5 Carbohydrates [i](#)

There are no oligosaccharides in this entry.

## 5.6 Ligand geometry [i](#)

There are no ligands in this entry.

## 5.7 Other polymers [i](#)

There are no such residues in this entry.

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

There are no chain breaks in this entry.

## 6 Map visualisation [i](#)

This section contains visualisations of the EMDB entry EMD-53415. These allow visual inspection of the internal detail of the map and identification of artifacts.

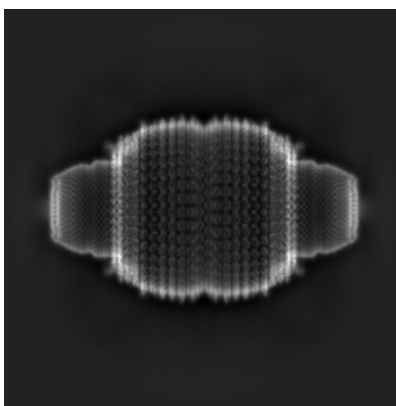
Images derived from a raw map, generated by summing the deposited half-maps, are presented below the corresponding image components of the primary map to allow further visual inspection and comparison with those of the primary map.

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

#### 6.1.1 Primary map



X

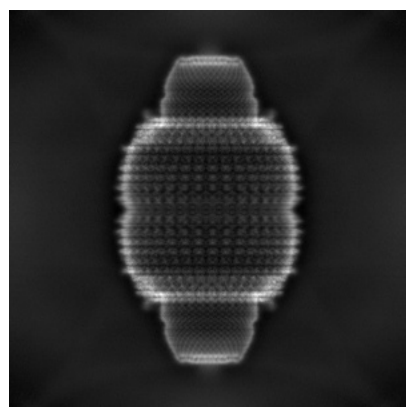


Y

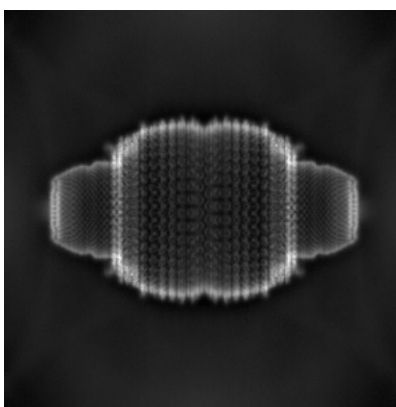


Z

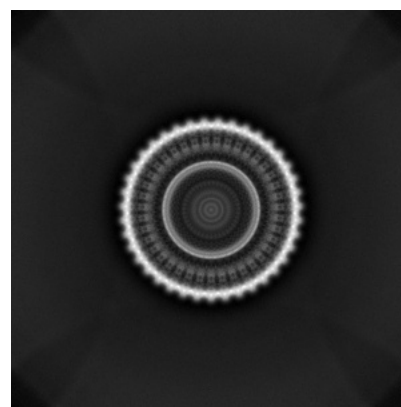
#### 6.1.2 Raw map



X



Y

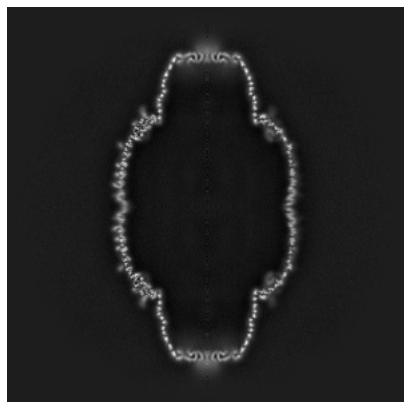


Z

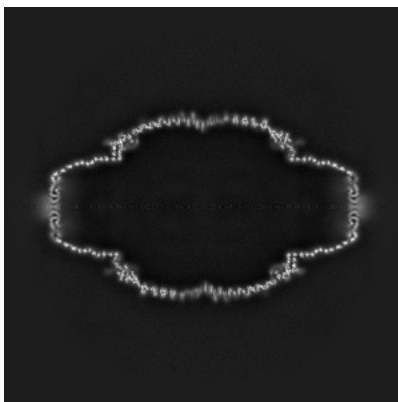
The images above show the map projected in three orthogonal directions.

## 6.2 Central slices [i](#)

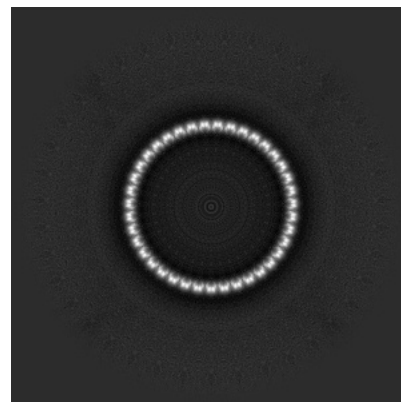
### 6.2.1 Primary map



X Index: 512

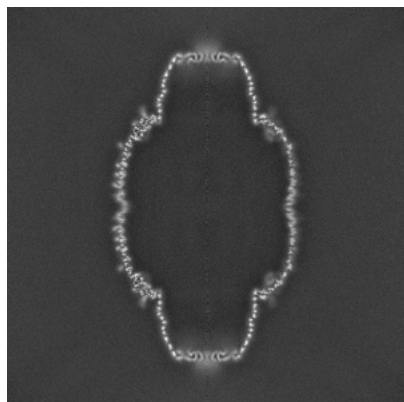


Y Index: 512

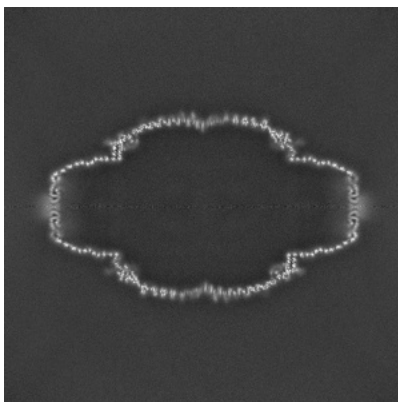


Z Index: 512

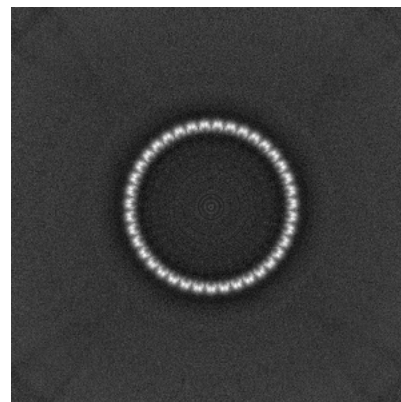
### 6.2.2 Raw map



X Index: 512



Y Index: 512

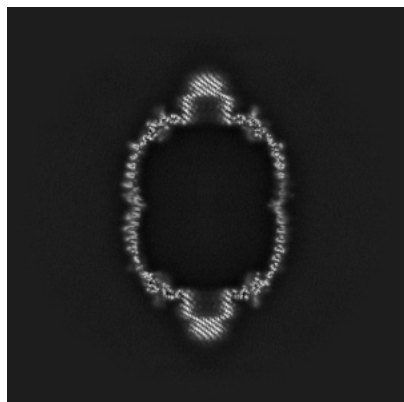


Z Index: 512

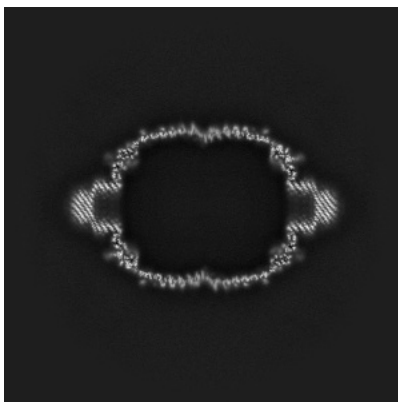
The images above show central slices of the map in three orthogonal directions.

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

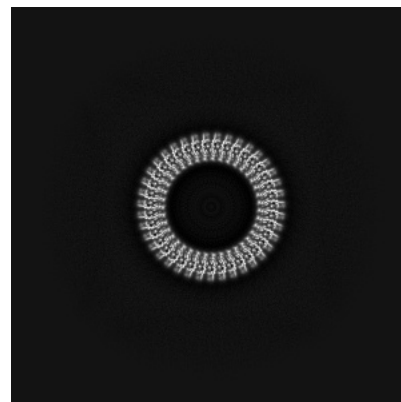
### 6.3.1 Primary map



X Index: 402

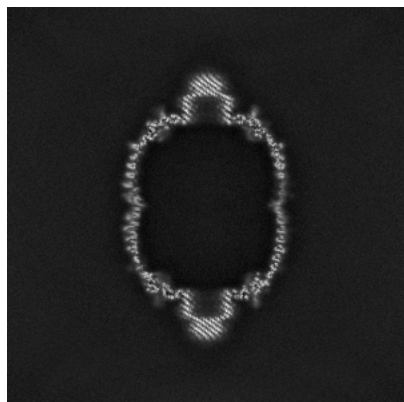


Y Index: 622

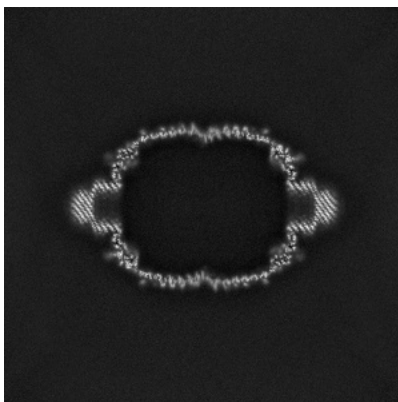


Z Index: 728

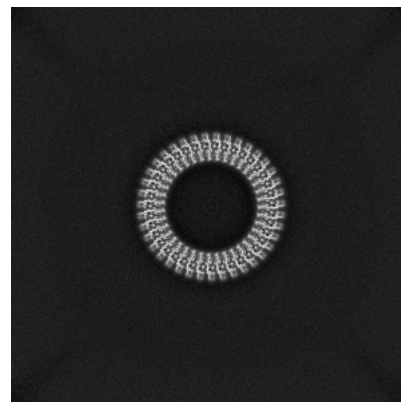
### 6.3.2 Raw map



X Index: 402



Y Index: 622

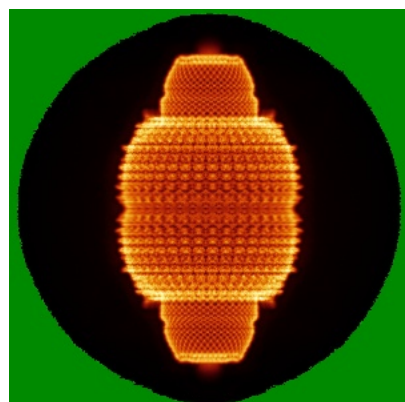


Z Index: 296

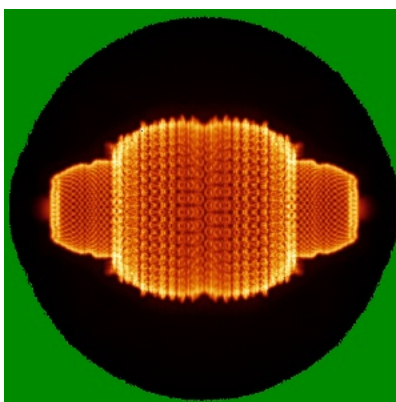
The images above show the largest variance slices of the map in three orthogonal directions.

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

### 6.4.1 Primary map



X

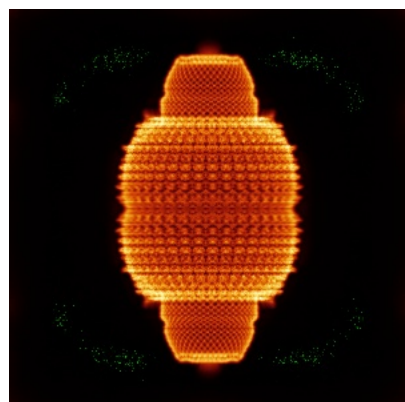


Y

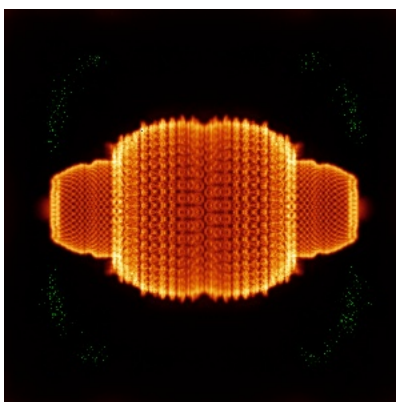


Z

### 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](#)

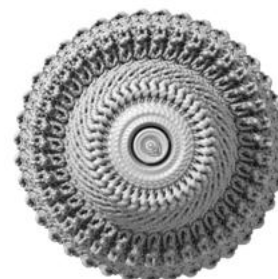
### 6.5.1 Primary map



X



Y



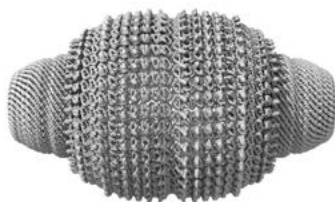
Z

The images above show the 3D surface view of the map at the recommended contour level 0.6. These images, in conjunction with the slice images, may facilitate assessment of whether an appropriate contour level has been provided.

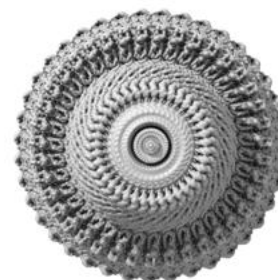
### 6.5.2 Raw map



X



Y



Z

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.



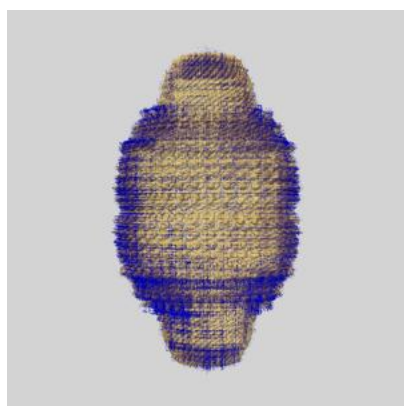
## 6.6 Mask visualisation [i](#)

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

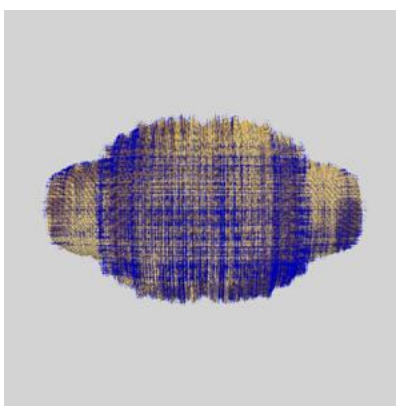
A mask typically either:

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

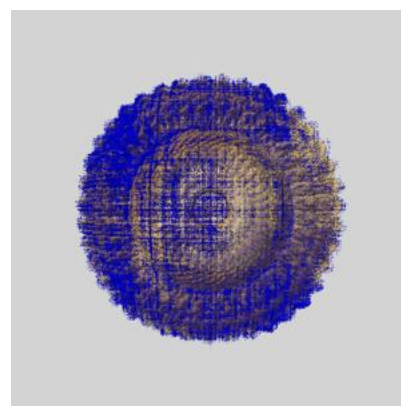
### 6.6.1 emd\_53415\_msk\_1.map [i](#)



X



Y

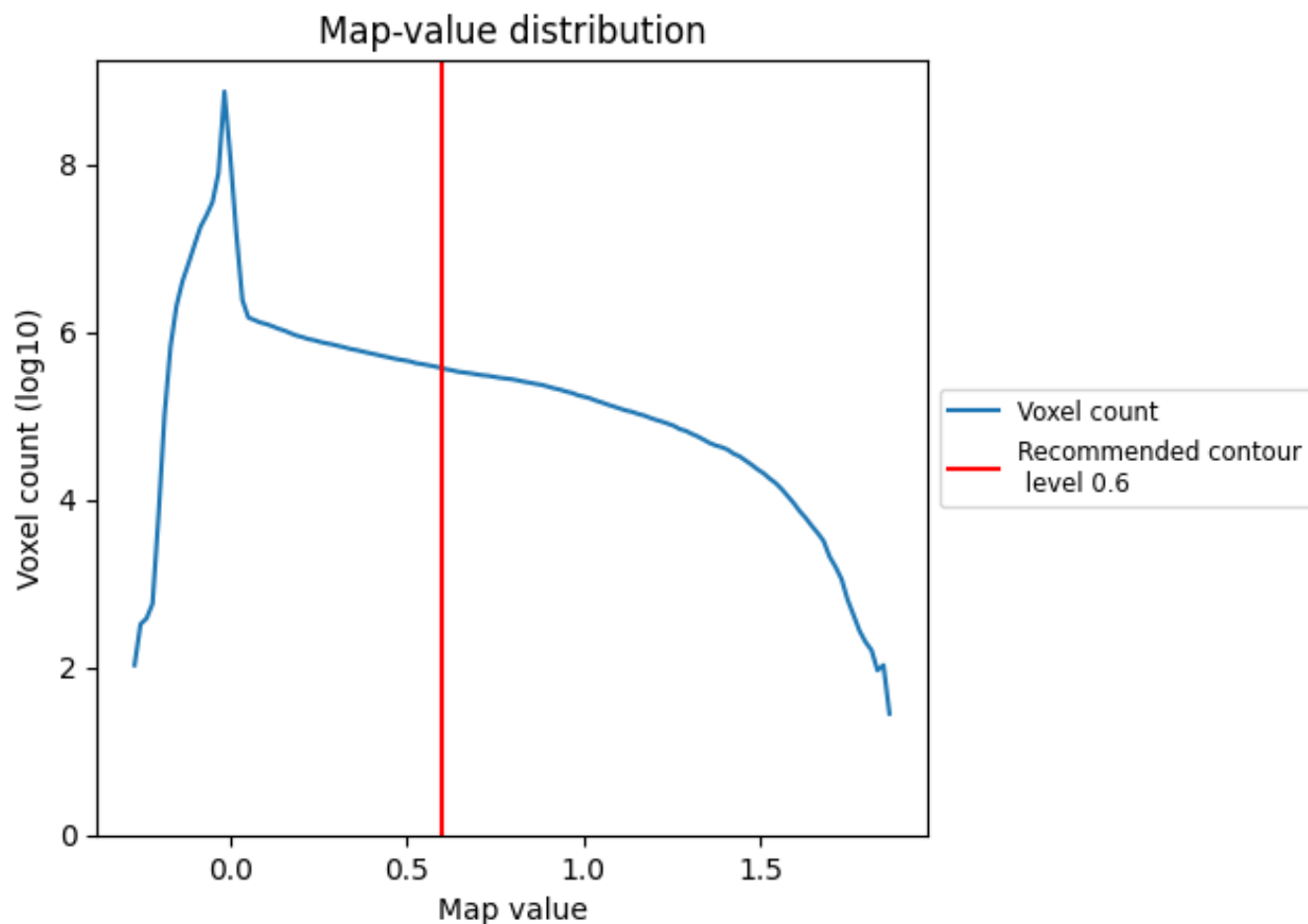


Z

## 7 Map analysis [i](#)

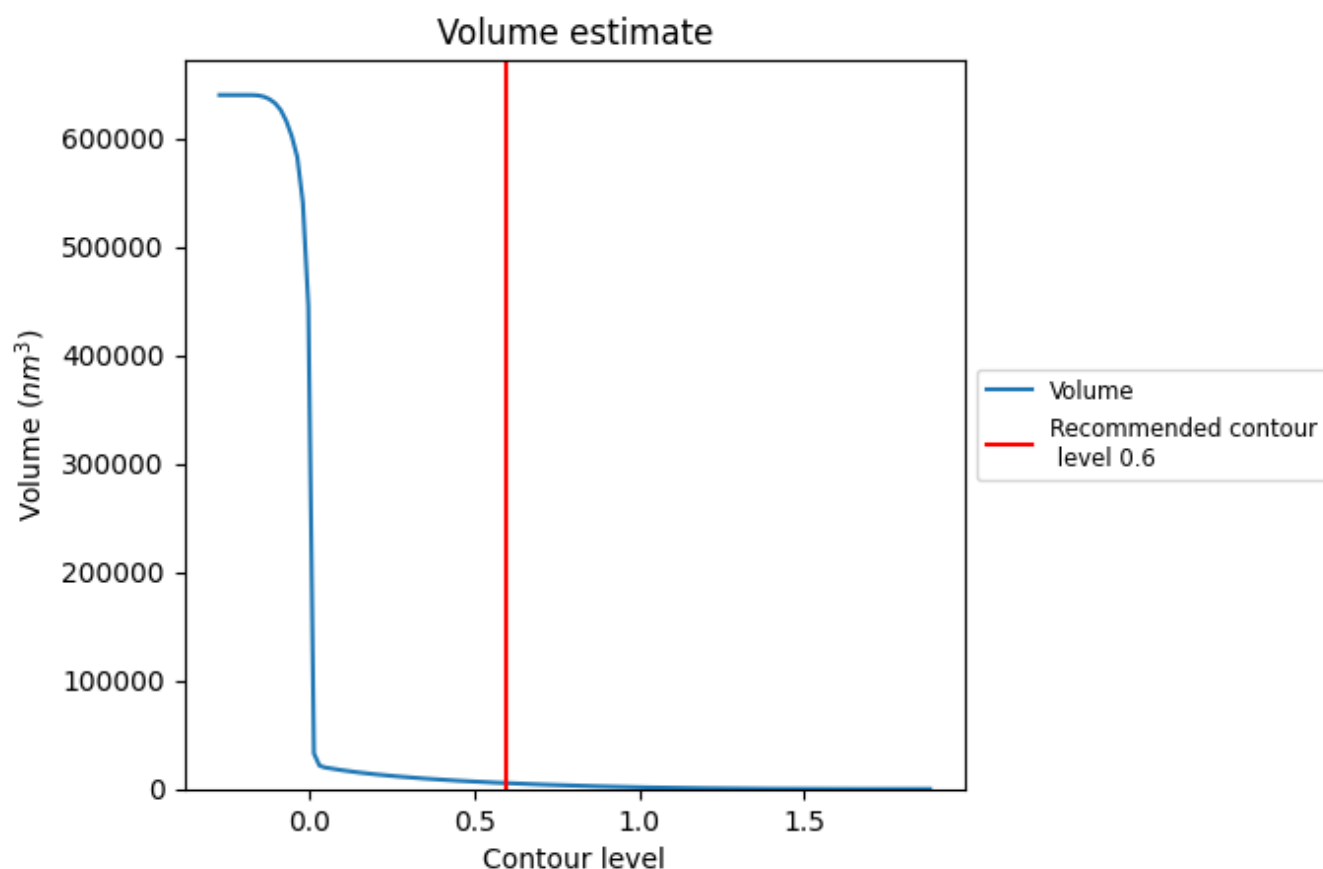
This section contains the results of statistical analysis of the map.

### 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.

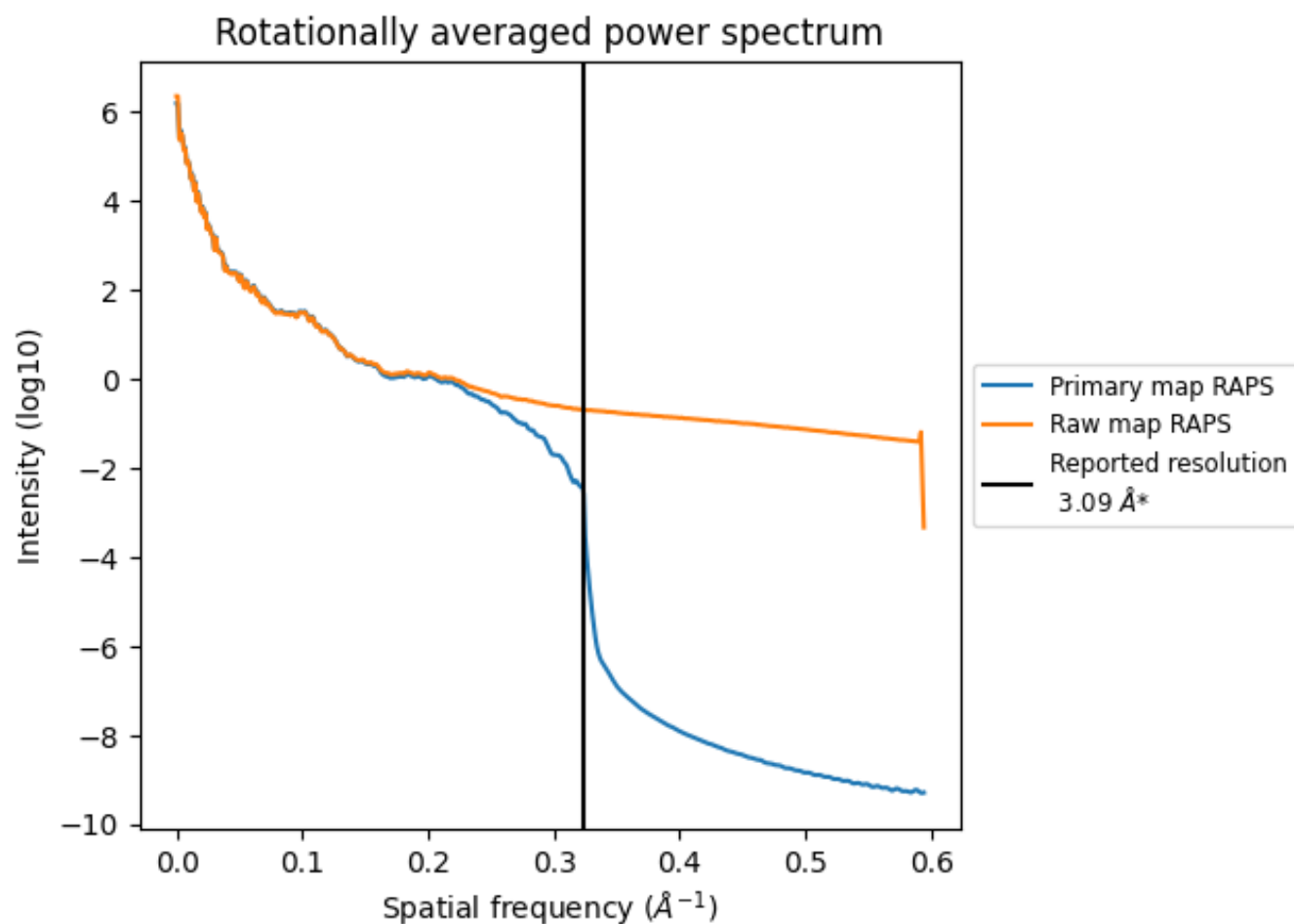
## 7.2 Volume estimate [i](#)



The volume at the recommended contour level is 5349  $\text{nm}^3$ ; this corresponds to an approximate mass of 4832 kDa.

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

### 7.3 Rotationally averaged power spectrum ⓘ

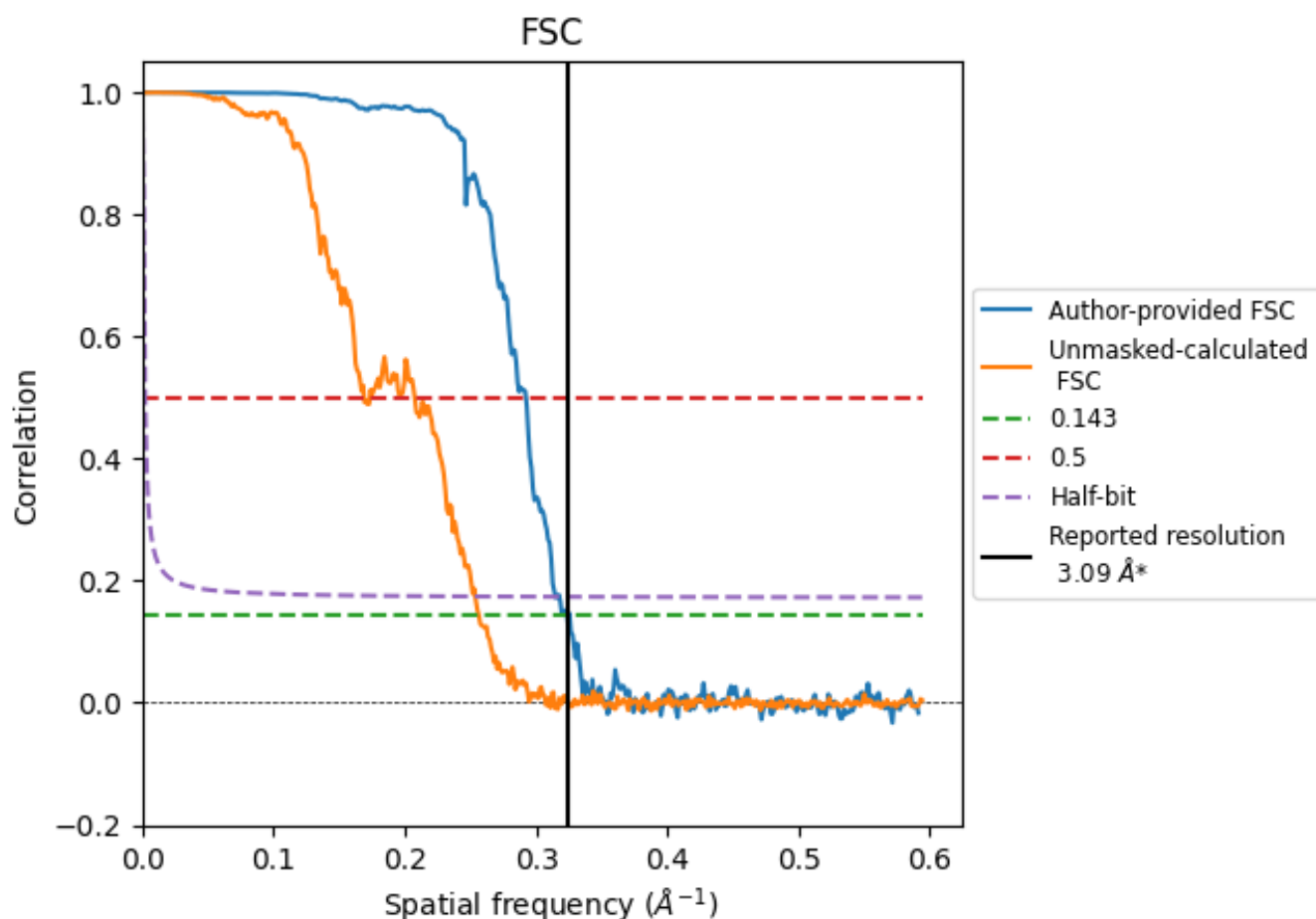


\*Reported resolution corresponds to spatial frequency of 0.324 Å<sup>-1</sup>

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

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

### 8.1 FSC [i](#)



\*Reported resolution corresponds to spatial frequency of 0.324  $\text{\AA}^{-1}$

## 8.2 Resolution estimates [i](#)

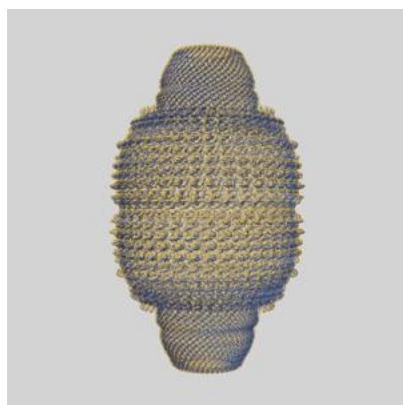
Resolution estimate (Å)	Estimation criterion (FSC cut-off)		
	0.143	0.5	Half-bit
Reported by author	3.09	-	-
Author-provided FSC curve	3.09	3.42	3.18
Unmasked-calculated*	3.90	5.91	3.94

\*Resolution estimate based on FSC curve calculated by comparison of deposited half-maps. The value from deposited half-maps intersecting FSC 0.143 CUT-OFF 3.90 differs from the reported value 3.09 by more than 10 %

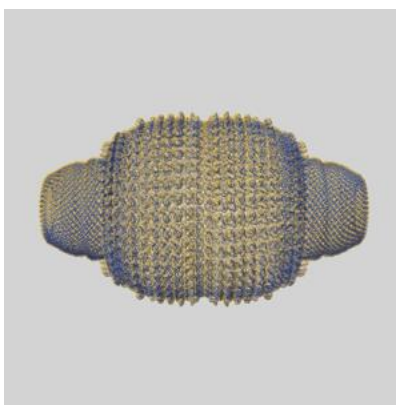
## 9 Map-model fit [i](#)

This section contains information regarding the fit between EMDB map EMD-53415 and PDB model 9QW9. Per-residue inclusion information can be found in section [3](#) on page [10](#).

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



X



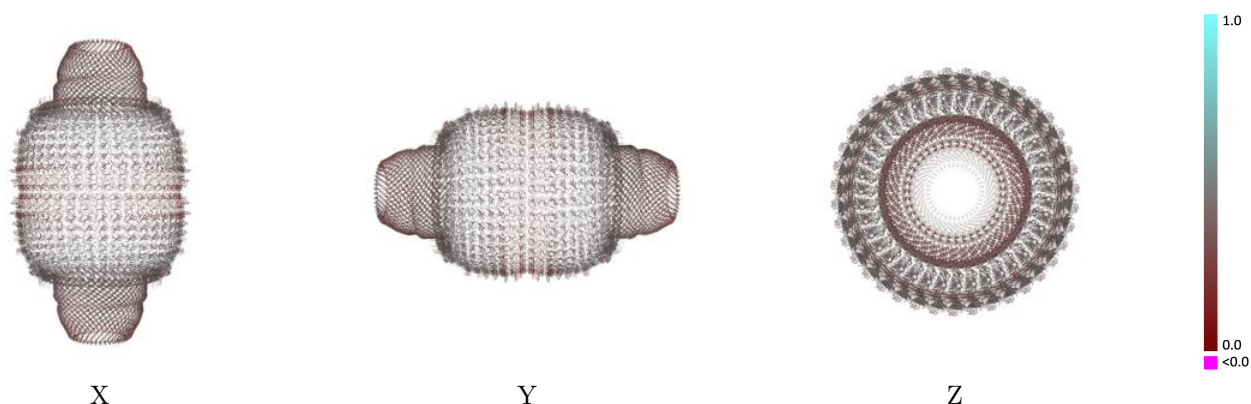
Y



Z

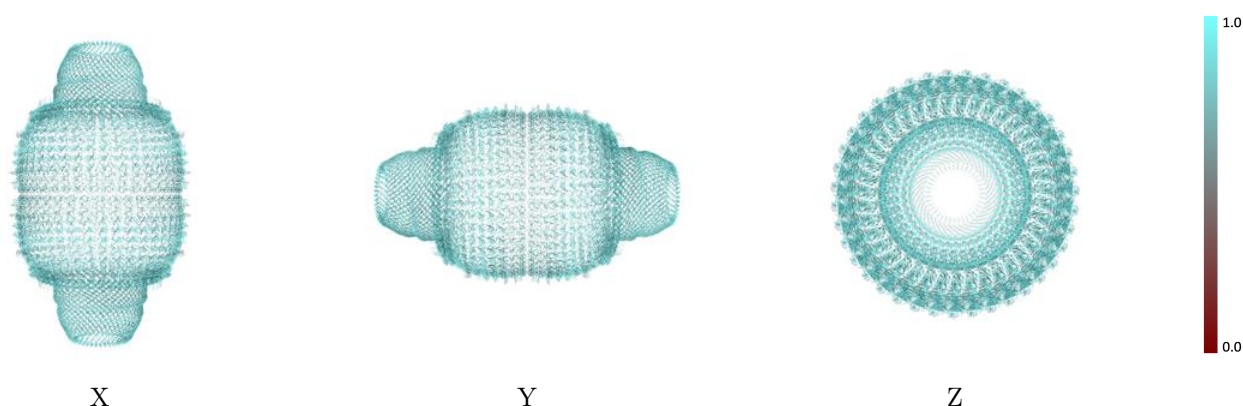
The images above show the 3D surface view of the map at the recommended contour level 0.6 at 50% transparency in yellow overlaid with a ribbon representation of the model coloured in blue. These images allow for the visual assessment of the quality of fit between the atomic model and the map.

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



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

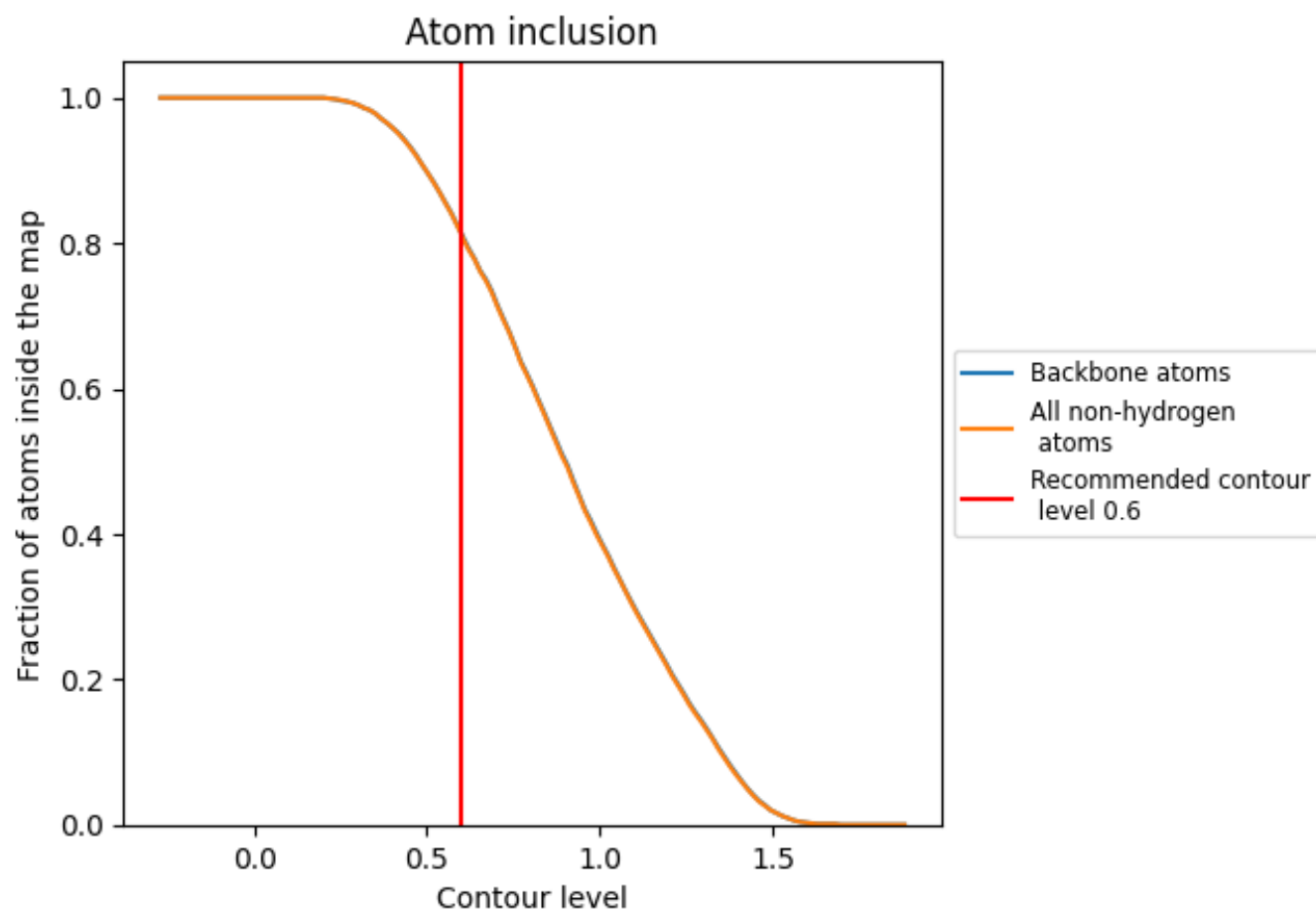
## 9.3 Atom inclusion mapped to coordinate model [i](#)



The images above show the model with each residue coloured according to its atom inclusion. This shows to what extent they are inside the map at the recommended contour level (0.6).









































































## 9.4 Atom inclusion [i](#)



At the recommended contour level, 81% of all backbone atoms, 81% 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 (0.6) and Q-score for the entire model and for each chain.

Chain	Atom inclusion	Q-score
All	 0.8120	 0.3810
A	 0.8240	 0.3810
AA	 0.8250	 0.3800
AB	 0.8240	 0.3820
AC	 0.8220	 0.3810
B	 0.8250	 0.3810
BA	 0.8240	 0.3800
BB	 0.8260	 0.3820
C	 0.8240	 0.3810
CA	 0.8250	 0.3800
CB	 0.8250	 0.3810
D	 0.8230	 0.3810
DA	 0.8250	 0.3810
DB	 0.8260	 0.3800
E	 0.8250	 0.3810
EA	 0.8250	 0.3810
EB	 0.8240	 0.3800
F	 0.8250	 0.3820
FA	 0.8230	 0.3810
FB	 0.8260	 0.3800
G	 0.8240	 0.3820
GA	 0.8260	 0.3810
GB	 0.8240	 0.3810
H	 0.8240	 0.3820
HA	 0.8250	 0.3810
HB	 0.8250	 0.3810
I	 0.8250	 0.3810
IA	 0.8240	 0.3810
IB	 0.8260	 0.3810
J	 0.8250	 0.3800
JA	 0.8240	 0.3800
JB	 0.8240	 0.3800
K	 0.8250	 0.3800
KA	 0.8240	 0.3800
KB	 0.8250	 0.3800





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Chain	Atom inclusion	Q-score
L	 0.8240	 0.3800
LA	 0.8240	 0.3810
LB	 0.8250	 0.3800
M	 0.8250	 0.3800
MA	 0.8230	 0.3810
MB	 0.8260	 0.3800
N	 0.8250	 0.3800
NA	 0.8230	 0.3810
NB	 0.8250	 0.3800
O	 0.8250	 0.3800
OA	 0.8230	 0.3820
OB	 0.8250	 0.3810
P	 0.8250	 0.3800
PA	 0.8250	 0.3820
PB	 0.8250	 0.3810
Q	 0.8230	 0.3800
QA	 0.8260	 0.3820
QB	 0.8260	 0.3810
R	 0.8250	 0.3800
RA	 0.8230	 0.3820
RB	 0.8260	 0.3810
S	 0.8250	 0.3810
SA	 0.8240	 0.3820
SB	 0.8230	 0.3810
T	 0.8250	 0.3810
TA	 0.8260	 0.3820
TB	 0.8260	 0.3810
UA	 0.8250	 0.3820
UB	 0.8260	 0.3810
V	 0.8250	 0.3810
VA	 0.8240	 0.3810
VB	 0.8240	 0.3810
W	 0.8250	 0.3800
WA	 0.8250	 0.3810
WB	 0.8240	 0.3810
X	 0.8250	 0.3800
XA	 0.8250	 0.3810
XB	 0.8250	 0.3810
Y	 0.8250	 0.3790
YA	 0.8250	 0.3810
YB	 0.8250	 0.3820
Z	 0.8250	 0.3800

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Chain	Atom inclusion	Q-score
ZA	 0.8240	 0.3820
ZB	 0.8250	 0.3820