Full wwPDB NMR Structure Validation Report (i)

Mar 8, 2026 – 05:12 PM UTC

PDB ID	:	5T56 / pdb_00005t56
BMRB ID	:	30165
Title	:	[3]catenane from MccJ25 G12R/I13C/G21C lasso peptide
Authors	:	Link, A.J.; Allen, C.D.
Deposited on	:	2016-08-30

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

We welcome your comments at *validation@mail.wwpdb.org*
A user guide is available at
https://www.wwpdb.org/validation/2017/NMRValidationReportHelp
with specific help available everywhere you see the (i) symbol.

The types of validation reports are described at
http://www.wwpdb.org/validation/2017/FAQs#types.

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

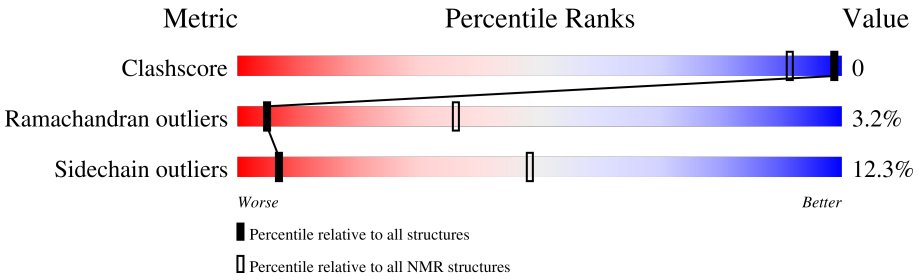
MolProbity	:	4-5-2 with Phenix2.0
Percentile statistics	:	20250101.v01 (using entries in the PDB archive January 1st 2025)
wwPDB-RCI	:	v_1n_11_5_13_A (Berjanski et al., 2005)
PANAV	:	Wang et al. (2010)
wwPDB-ShiftChecker	:	v1.2
BMRB Restraints Analysis	:	v1.2
Ideal geometry (proteins)	:	Engh & Huber (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:
SOLUTION NMR

The overall completeness of chemical shifts assignment is 46%.

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)	NMR archive (#Entries)
Clashscore	229148	14424
Ramachandran outliers	224038	12848
Sidechain outliers	223484	12823

The table below summarises the geometric issues observed across the polymeric chains and their fit to the experimental data. The red, orange, yellow and green segments indicate the fraction of residues that contain outliers for >=3, 2, 1 and 0 types of geometric quality criteria. A cyan segment indicates the fraction of residues that are not part of the well-defined cores, and 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 <=5%

Mol	Chain	Length	Quality of chain
1	A	12	92% 8%
1	C	12	92% 8%
2	B	9	100%
2	D	9	100%

2 Ensemble composition and analysis

This entry contains 20 models. Model 13 is the overall representative, medoid model (most similar to other models). The authors have identified model 1 as representative, based on the following criterion: *target function*.

The following residues are included in the computation of the global validation metrics.

Well-defined (core) protein residues			
Well-defined core	**Residue range (total)**	**Backbone RMSD (Å)**	**Medoid model**
1	A:1-A:12, B:13-B:21, C:22-C:33, D:34-D:42 (42)	1.18	13

Ill-defined regions of proteins are excluded from the global statistics.

Ligands and non-protein polymers are included in the analysis.

The models can be grouped into 4 clusters and 4 single-model clusters were found.

Cluster number	Models
1	2, 4, 11, 13, 15, 16, 18
2	5, 6, 10, 19, 20
3	7, 8
4	3, 9
Single-model clusters	1; 12; 14; 17

3 Entry composition (i)

There are 2 unique types of molecules in this entry. The entry contains 606 atoms, of which 288 are hydrogens and 0 are deuteriums.

- Molecule 1 is a protein called Microcin J25.

Mol	Chain	Residues	Atoms					Trace
1	A	12	Total	C	H	N	O	0
			174	59	83	17	15	
1	C	12	Total	C	H	N	O	0
			174	59	83	17	15	

There are 2 discrepancies between the modelled and reference sequences:

Chain	Residue	Modelled	Actual	Comment	Reference
A	12	ARG	-	expression tag	UNP Q9X2V7
C	33	ARG	-	expression tag	UNP Q9X2V7

- Molecule 2 is a protein called Microcin J25.

Mol	Chain	Residues	Atoms						Trace
2	B	9	Total	C	H	N	O	S	0
			129	44	61	9	13	2	
2	D	9	Total	C	H	N	O	S	0
			129	44	61	9	13	2	

There are 4 discrepancies between the modelled and reference sequences:

Chain	Residue	Modelled	Actual	Comment	Reference
B	13	CYS	-	expression tag	UNP Q9X2V7
B	21	CYS	-	expression tag	UNP Q9X2V7
D	34	CYS	-	expression tag	UNP Q9X2V7
D	42	CYS	-	expression tag	UNP Q9X2V7

4 Residue-property plots

4.1 Average score per residue in the NMR ensemble

These plots are provided for all protein, RNA, DNA and oligosaccharide chains in the entry. The first graphic is the same as shown in the summary in section 1 of this report. The second graphic shows the sequence where residues are colour-coded according to the number of geometric quality criteria for which they contain at least one outlier: green = 0, yellow = 1, orange = 2 and red = 3 or more. Stretches of 2 or more consecutive residues without any outliers are shown as green connectors. Residues which are classified as ill-defined in the NMR ensemble, are shown in cyan with an underline colour-coded according to the previous scheme. Residues which were present in the experimental sample, but not modelled in the final structure are shown in grey.

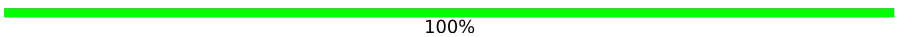
• Molecule 1: Microcin J25

Chain A:

• Molecule 1: Microcin J25

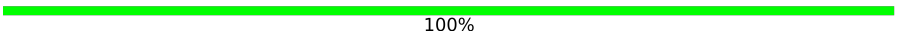
Chain C:

• Molecule 2: Microcin J25

Chain B: 

There are no outlier residues in this chain.

• Molecule 2: Microcin J25

Chain D: 

There are no outlier residues in this chain.

4.2 Scores per residue for each member of the ensemble

Colouring as in section 4.1 above.

4.2.1 Score per residue for model 1

• Molecule 1: Microcin J25

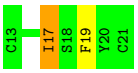
Chain A:

• Molecule 1: Microcin J25

Chain C:

• Molecule 2: Microcin J25

Chain B:



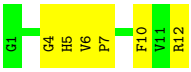
• Molecule 2: Microcin J25

Chain D:

4.2.2 Score per residue for model 2

• Molecule 1: Microcin J25

Chain A:



• Molecule 1: Microcin J25

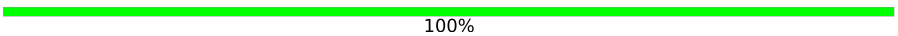
Chain C:

• Molecule 2: Microcin J25

Chain B:

• Molecule 2: Microcin J25

Chain D:



There are no outlier residues in this chain.

4.2.3 Score per residue for model 3

• Molecule 1: Microcin J25

Chain A:

• Molecule 1: Microcin J25

Chain C:

• Molecule 2: Microcin J25

Chain B:

• Molecule 2: Microcin J25

Chain D:

4.2.4 Score per residue for model 4

• Molecule 1: Microcin J25

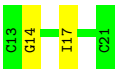
Chain A:

• Molecule 1: Microcin J25

Chain C:

• Molecule 2: Microcin J25

Chain B:



• Molecule 2: Microcin J25

Chain D:

4.2.5 Score per residue for model 5

• Molecule 1: Microcin J25

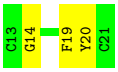
Chain A:

• Molecule 1: Microcin J25

Chain C:

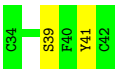
• Molecule 2: Microcin J25

Chain B:



• Molecule 2: Microcin J25

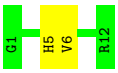
Chain D:



4.2.6 Score per residue for model 6

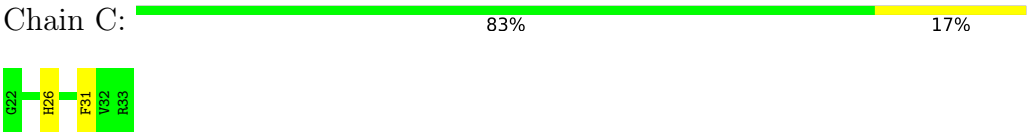
• Molecule 1: Microcin J25

Chain A:



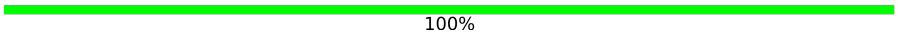
• Molecule 1: Microcin J25

Chain C:



• Molecule 2: Microcin J25

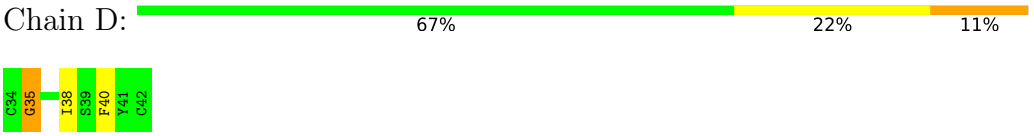
Chain B:



There are no outlier residues in this chain.

• Molecule 2: Microcin J25

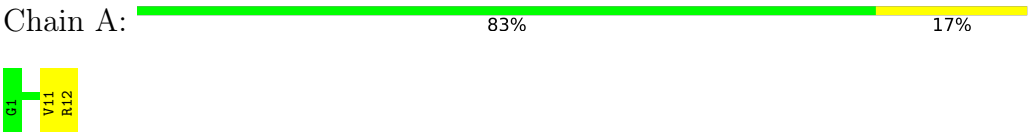
Chain D:



4.2.7 Score per residue for model 7

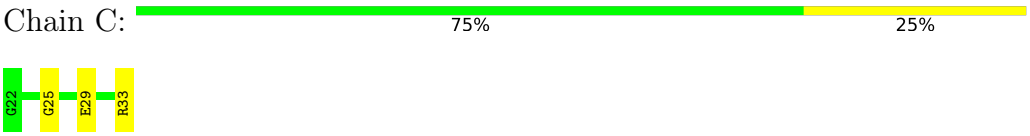
• Molecule 1: Microcin J25

Chain A:



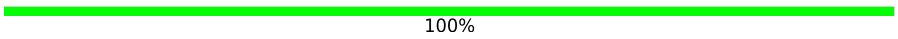
• Molecule 1: Microcin J25

Chain C:



• Molecule 2: Microcin J25

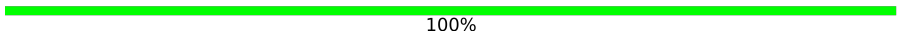
Chain B:



There are no outlier residues in this chain.

• Molecule 2: Microcin J25

Chain D:

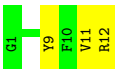


There are no outlier residues in this chain.

4.2.8 Score per residue for model 8

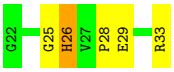
• Molecule 1: Microcin J25

Chain A:



• Molecule 1: Microcin J25

Chain C:

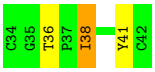


• Molecule 2: Microcin J25

Chain B:

• Molecule 2: Microcin J25

Chain D:



4.2.9 Score per residue for model 9

• Molecule 1: Microcin J25

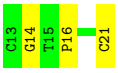
Chain A:

• Molecule 1: Microcin J25

Chain C:

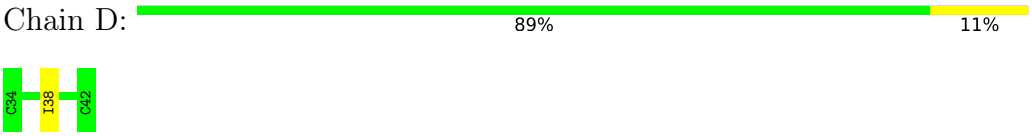
• Molecule 2: Microcin J25

Chain B:



• Molecule 2: Microcin J25

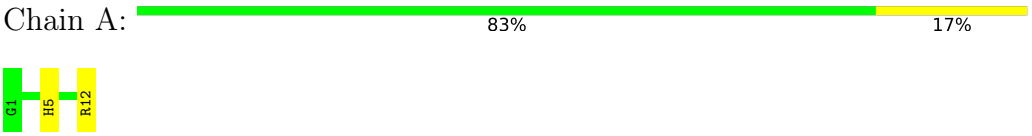
Chain D:



4.2.10 Score per residue for model 10

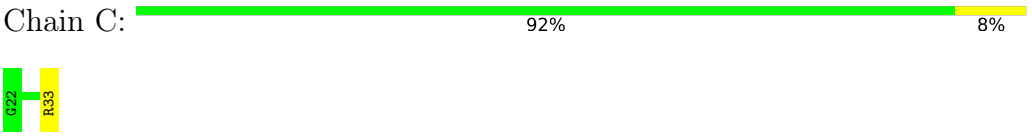
• Molecule 1: Microcin J25

Chain A:



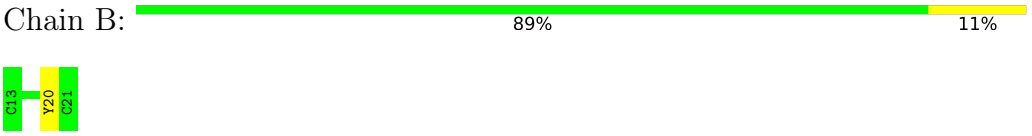
• Molecule 1: Microcin J25

Chain C:



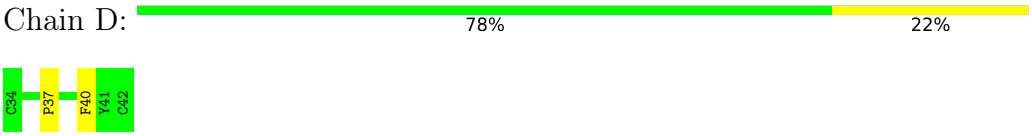
• Molecule 2: Microcin J25

Chain B:



• Molecule 2: Microcin J25

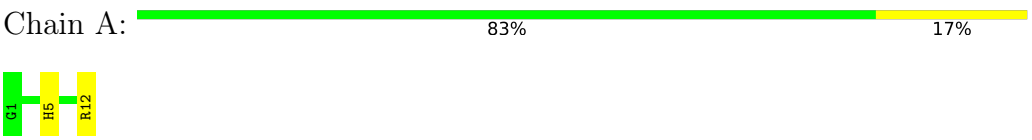
Chain D:



4.2.11 Score per residue for model 11

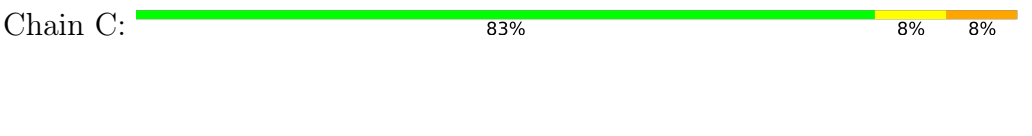
• Molecule 1: Microcin J25

Chain A:



• Molecule 1: Microcin J25

Chain C:

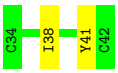


• Molecule 2: Microcin J25

Chain B:

• Molecule 2: Microcin J25

Chain D:



4.2.12 Score per residue for model 12

• Molecule 1: Microcin J25

Chain A:

• Molecule 1: Microcin J25

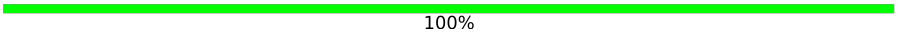
Chain C:

• Molecule 2: Microcin J25

Chain B:

• Molecule 2: Microcin J25

Chain D:

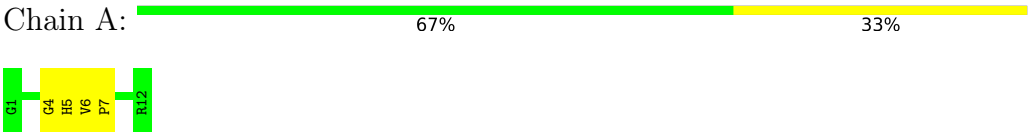


There are no outlier residues in this chain.

4.2.13 Score per residue for model 13 (medoid)

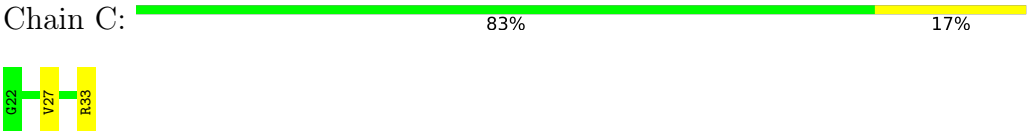
• Molecule 1: Microcin J25

Chain A:



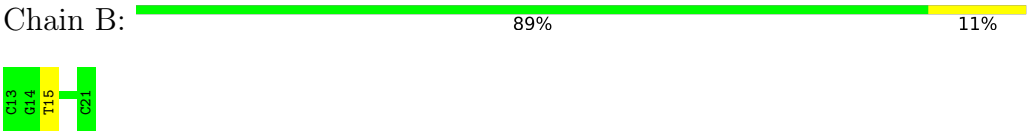
• Molecule 1: Microcin J25

Chain C:



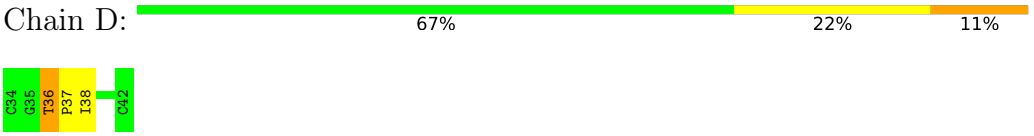
• Molecule 2: Microcin J25

Chain B:



• Molecule 2: Microcin J25

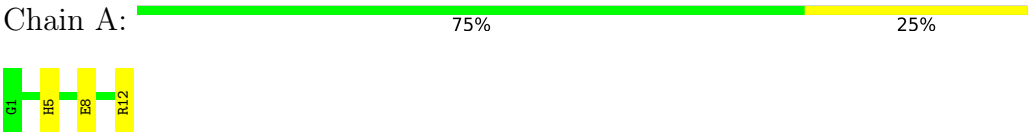
Chain D:



4.2.14 Score per residue for model 14

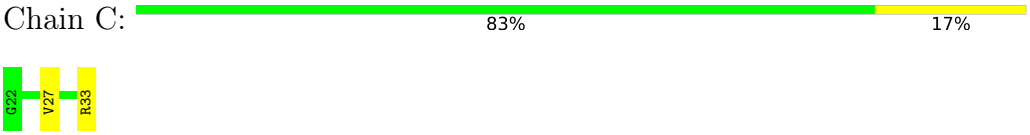
• Molecule 1: Microcin J25

Chain A:



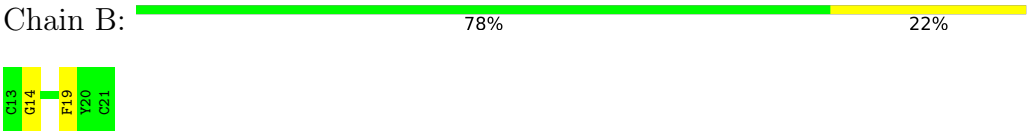
• Molecule 1: Microcin J25

Chain C:



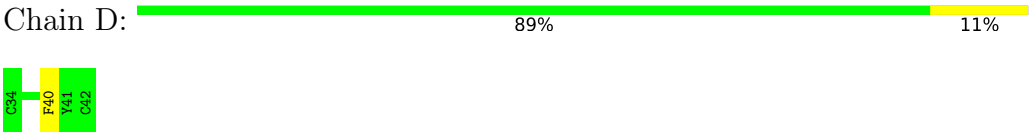
• Molecule 2: Microcin J25

Chain B:



• Molecule 2: Microcin J25

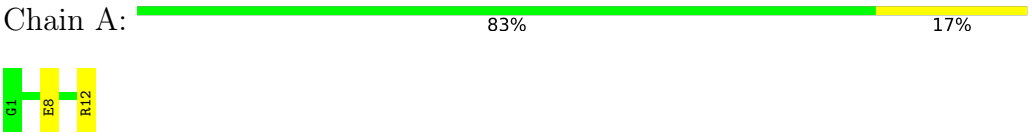
Chain D:



4.2.15 Score per residue for model 15

• Molecule 1: Microcin J25

Chain A:



• Molecule 1: Microcin J25

Chain C:

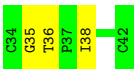
• Molecule 2: Microcin J25

Chain B:

There are no outlier residues in this chain.

• Molecule 2: Microcin J25

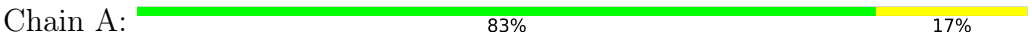
Chain D:



4.2.16 Score per residue for model 16

• Molecule 1: Microcin J25

Chain A:



• Molecule 1: Microcin J25

Chain C:

• Molecule 2: Microcin J25

Chain B:

• Molecule 2: Microcin J25

Chain D:

4.2.17 Score per residue for model 17

• Molecule 1: Microcin J25

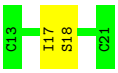
Chain A:

• Molecule 1: Microcin J25

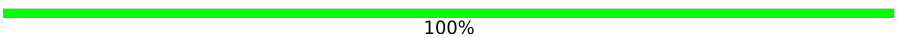
Chain C:

• Molecule 2: Microcin J25

Chain B:



• Molecule 2: Microcin J25

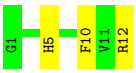
Chain D: 

There are no outlier residues in this chain.

4.2.18 Score per residue for model 18

• Molecule 1: Microcin J25

Chain A:

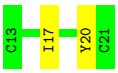


• Molecule 1: Microcin J25

Chain C:

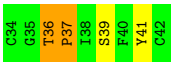
• Molecule 2: Microcin J25

Chain B:



• Molecule 2: Microcin J25

Chain D:



4.2.19 Score per residue for model 19

• Molecule 1: Microcin J25

Chain A:

• Molecule 1: Microcin J25

Chain C:

• Molecule 2: Microcin J25

Chain B:

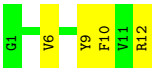
• Molecule 2: Microcin J25

Chain D:

4.2.20 Score per residue for model 20

• Molecule 1: Microcin J25

Chain A:

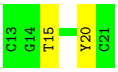


• Molecule 1: Microcin J25

Chain C:

• Molecule 2: Microcin J25

Chain B:



• Molecule 2: Microcin J25

Chain D:

5 Refinement protocol and experimental data overview

The models were refined using the following method: *molecular dynamics*.

Of the 20 calculated structures, 20 were deposited, based on the following criterion: *all calculated structures submitted*.

The following table shows the software used for structure solution, optimisation and refinement.

Software name	Classification	Version
GROMACS	refinement	5.1.1
CYANA	structure calculation	2.1

The following table shows chemical shift validation statistics as aggregates over all chemical shift files. Detailed validation can be found in section 7 of this report.

Chemical shift file(s)	working_cs.cif
Number of chemical shift lists	1
Total number of shifts	238
Number of shifts mapped to atoms	238
Number of unparsed shifts	0
Number of shifts with mapping errors	0
Number of shifts with mapping warnings	0
Assignment completeness (well-defined parts)	46%

6 Model quality (i)

6.1 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 (average) 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	0.70±0.02	0±0/94 (0.0± 0.0%)	1.77±0.12	1±1/124 (0.8± 0.6%)
1	C	0.70±0.02	0±0/94 (0.0± 0.0%)	1.80±0.10	1±1/124 (0.8± 0.9%)
2	B	0.61±0.01	0±0/70 (0.0± 0.0%)	1.59±0.19	1±1/93 (0.8± 1.2%)
2	D	0.61±0.02	0±0/70 (0.0± 0.0%)	1.68±0.18	1±1/93 (0.7± 0.9%)
All	All	0.66	0/6560 (0.0%)	1.73	66/8680 (0.8%)

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

Mol	Chain	Chirality	Planarity
1	C	0.0±0.0	0.3±0.6
1	A	0.0±0.0	0.2±0.4
2	D	0.0±0.0	0.2±0.4
2	B	0.0±0.0	0.1±0.4
All	All	0	19

There are no bond-length outliers.

All unique angle outliers are listed below. They are sorted according to the Z-score of the worst occurrence in the ensemble.

Mol	Chain	Res	Type	Atoms	Z	Observed(^{o})	Ideal(^{o})	Models	
								Worst	Total
2	D	36	THR	CA-CB-CG2	8.49	124.94	110.50	18	3
2	D	41	TYR	CA-C-N	6.64	133.64	121.70	16	1
2	D	41	TYR	C-N-CA	6.64	133.64	121.70	16	1
2	D	41	TYR	N-CA-C	6.48	121.39	112.90	5	1
1	A	5	HIS	CB-CG-CD2	-6.35	122.94	131.20	11	7
2	B	17	ILE	CB-CA-C	6.26	121.56	111.29	1	1
1	C	29	GLU	CA-C-N	5.90	130.78	122.46	7	1
1	C	29	GLU	C-N-CA	5.90	130.78	122.46	7	1
2	B	19	PHE	CA-CB-CG	5.84	119.64	113.80	1	3

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Mol	Chain	Res	Type	Atoms	Z	Observed(^{o})	Ideal(^{o})	Models	
								Worst	Total
1	C	28	PRO	CA-C-N	5.84	133.19	122.79	5	4
1	C	28	PRO	C-N-CA	5.84	133.19	122.79	5	4
1	C	26	HIS	CB-CG-CD2	-5.79	123.67	131.20	6	4
1	A	9	TYR	CA-C-N	5.75	129.04	120.31	8	1
1	A	9	TYR	C-N-CA	5.75	129.04	120.31	8	1
2	B	14	GLY	CA-C-N	5.72	130.01	120.86	14	2
2	B	14	GLY	C-N-CA	5.72	130.01	120.86	14	2
2	B	16	PRO	CA-C-N	5.56	128.69	120.74	9	1
2	B	16	PRO	C-N-CA	5.56	128.69	120.74	9	1
2	D	37	PRO	N-CA-CB	5.56	108.19	103.35	19	1
2	B	15	THR	O-C-N	-5.53	118.17	121.71	12	1
1	A	5	HIS	CA-CB-CG	5.50	119.30	113.80	2	1
2	D	38	ILE	CA-C-O	5.49	123.74	119.46	15	1
1	C	25	GLY	CA-C-N	5.41	130.68	122.08	16	1
1	C	25	GLY	C-N-CA	5.41	130.68	122.08	16	1
1	A	7	PRO	CA-C-N	5.35	128.06	121.64	13	1
1	A	7	PRO	C-N-CA	5.35	128.06	121.64	13	1
2	D	40	PHE	CA-CB-CG	5.34	119.14	113.80	14	2
1	A	5	HIS	ND1-CG-CD2	5.31	111.41	106.10	4	1
2	B	16	PRO	N-CA-CB	5.22	107.47	103.30	2	1
1	A	12	ARG	CD-NE-CZ	5.21	131.70	124.40	1	1
1	A	10	PHE	CA-C-N	5.17	128.89	121.96	17	1
1	A	10	PHE	C-N-CA	5.17	128.89	121.96	17	1
1	C	26	HIS	CA-CB-CG	5.14	118.94	113.80	15	1
2	D	38	ILE	CA-CB-CG1	5.12	119.11	110.40	8	1
2	D	39	SER	CA-C-N	5.10	130.75	122.53	5	1
2	D	39	SER	C-N-CA	5.10	130.75	122.53	5	1
1	A	6	VAL	CA-C-N	5.09	124.85	119.76	9	1
1	A	6	VAL	C-N-CA	5.09	124.85	119.76	9	1
2	B	15	THR	CA-C-N	5.08	124.84	119.76	19	1
2	B	15	THR	C-N-CA	5.08	124.84	119.76	19	1
1	C	25	GLY	N-CA-C	5.06	116.98	110.96	18	1
1	A	7	PRO	N-CA-CB	5.03	106.60	103.23	2	1
1	C	24	ALA	CA-C-N	5.01	128.70	121.54	1	1
1	C	24	ALA	C-N-CA	5.01	128.70	121.54	1	1

There are no chirality outliers.

All unique planar outliers are listed below. They are sorted by the frequency of occurrence in the ensemble.

Mol	Chain	Res	Type	Group	Models (Total)
1	A	4	GLY	Peptide	3
1	C	33	ARG	Sidechain	3
2	D	41	TYR	Sidechain,Peptide	3
1	A	9	TYR	Sidechain	2
1	C	25	GLY	Peptide	2
2	B	14	GLY	Peptide	1
2	B	19	PHE	Peptide	1
2	D	35	GLY	Peptide	1
2	D	40	PHE	Sidechain	1
1	C	30	TYR	Sidechain	1
2	B	20	TYR	Peptide	1

6.2 Too-close contacts

In the following table, the Non-H and H(model) columns list the number of non-hydrogen atoms and hydrogen atoms in each 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 averaged over the ensemble.

Mol	Chain	Non-H	H(model)	H(added)	Clashes
1	A	91	83	82	0±0
All	All	6360	5760	5613	1

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 0.

All unique clashes are listed below, sorted by their clash magnitude.

Atom-1	Atom-2	Clash(Å)	Distance(Å)	Models	
				Worst	Total
1:A:11:VAL:HG22	1:A:12:ARG:N	0.42	2.29	8	1

6.3 Torsion angles

6.3.1 Protein backbone

In the following table, the Percentiles column shows the percent Ramachandran outliers of the chain as a percentile score with respect to all PDB entries followed by that with respect to all NMR 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	10/12 (83%)	8±1 (80±12%)	2±1 (20±11%)	0±0 (1±3%)	15 65
1	C	10/12 (83%)	8±1 (84±12%)	2±1 (16±12%)	0±0 (0±0%)	100 100
2	B	7/9 (78%)	5±1 (74±15%)	1±1 (19±14%)	0±1 (7±8%)	2 16
2	D	7/9 (78%)	5±1 (72±14%)	1±1 (21±15%)	0±1 (7±8%)	2 16
All	All	680/840 (81%)	533 (78%)	125 (18%)	22 (3%)	5 36

All 8 unique Ramachandran outliers are listed below. They are sorted by the frequency of occurrence in the ensemble.

Mol	Chain	Res	Type	Models (Total)
2	D	35	GLY	5
2	B	14	GLY	3
2	B	17	ILE	3
2	B	20	TYR	3
2	D	37	PRO	3
2	D	38	ILE	2
1	A	10	PHE	2
2	B	15	THR	1

6.3.2 Protein sidechains (i)

In the following table, the Percentiles column shows the percent sidechain outliers of the chain as a percentile score with respect to all PDB entries followed by that with respect to all NMR 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	8/8 (100%)	7±1 (82±10%)	1±1 (18±10%)	3 36
1	C	8/8 (100%)	7±1 (83±7%)	1±1 (17±7%)	4 38
2	B	8/8 (100%)	7±0 (93±6%)	1±0 (7±6%)	16 65
2	D	8/8 (100%)	7±1 (92±10%)	1±1 (8±10%)	14 62
All	All	640/640 (100%)	561 (88%)	79 (12%)	7 48

All 21 unique residues with a non-rotameric sidechain are listed below. They are sorted by the frequency of occurrence in the ensemble.

Mol	Chain	Res	Type	Models (Total)
1	A	12	ARG	16
1	C	33	ARG	15

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Mol	Chain	Res	Type	Models (Total)
2	D	38	ILE	6
1	A	8	GLU	5
2	B	15	THR	5
1	C	27	VAL	4
1	A	6	VAL	4
2	B	17	ILE	3
1	C	29	GLU	3
2	D	36	THR	3
1	A	11	VAL	2
1	C	26	HIS	2
1	C	31	PHE	2
2	B	18	SER	2
1	A	10	PHE	1
2	B	21	CYS	1
1	C	32	VAL	1
1	A	5	HIS	1
2	D	34	CYS	1
2	D	37	PRO	1
2	D	39	SER	1

6.3.3 RNA

There are no RNA molecules in this entry.

6.4 Non-standard residues in protein, DNA, RNA chains

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

6.5 Carbohydrates

There are no oligosaccharides in this entry.

6.6 Ligand geometry

There are no ligands in this entry.

6.7 Other polymers

There are no such molecules in this entry.

6.8 Polymer linkage issues (i)

There are no chain breaks in this entry.

7 Chemical shift validation

The completeness of assignment taking into account all chemical shift lists is 46% for the well-defined parts and 46% for the entire structure.

7.1 Chemical shift list 1

File name: working_cs.cif

Chemical shift list name: *nmr_star_coordinates_final3.txt*

7.1.1 Bookkeeping

The following table shows the results of parsing the chemical shift list and reports the number of nuclei with statistically unusual chemical shifts.

Total number of shifts	238
Number of shifts mapped to atoms	238
Number of unparsed shifts	0
Number of shifts with mapping errors	0
Number of shifts with mapping warnings	0
Number of shift outliers (ShiftChecker)	0

7.1.2 Chemical shift referencing

No chemical shift referencing corrections were calculated (not enough data).

7.1.3 Completeness of resonance assignments

The following table shows the completeness of the chemical shift assignments for the well-defined regions of the structure. The overall completeness is 46%, i.e. 238 atoms were assigned a chemical shift out of a possible 520. 0 out of 4 assigned methyl groups (LEU and VAL) were assigned stereospecifically.

	Total	^{1}H	^{13}C	^{15}N
Backbone	72/210 (34%)	72/88 (82%)	0/84 (0%)	0/38 (0%)
Sidechain	132/220 (60%)	132/146 (90%)	0/68 (0%)	0/6 (0%)
Aromatic	34/90 (38%)	34/44 (77%)	0/44 (0%)	0/2 (0%)
Overall	238/520 (46%)	238/278 (86%)	0/196 (0%)	0/46 (0%)

The following table shows the completeness of the chemical shift assignments for the full structure. The overall completeness is 46%, i.e. 238 atoms were assigned a chemical shift out of a possible 520. 0 out of 4 assigned methyl groups (LEU and VAL) were assigned stereospecifically.

	Total	^{1}H	^{13}C	^{15}N
Backbone	72/210 (34%)	72/88 (82%)	0/84 (0%)	0/38 (0%)
Sidechain	132/220 (60%)	132/146 (90%)	0/68 (0%)	0/6 (0%)
Aromatic	34/90 (38%)	34/44 (77%)	0/44 (0%)	0/2 (0%)
Overall	238/520 (46%)	238/278 (86%)	0/196 (0%)	0/46 (0%)

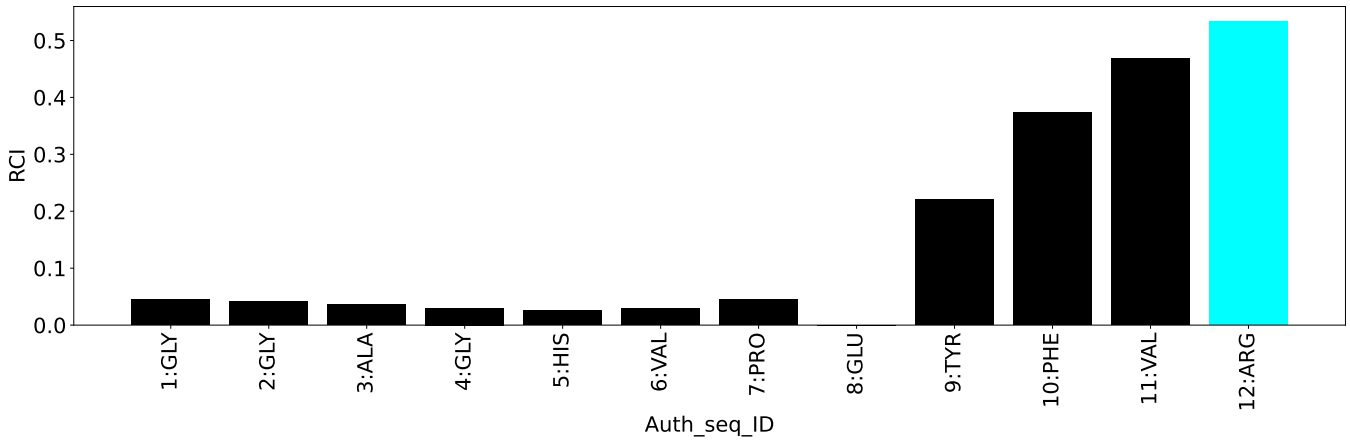
7.1.4 Statistically unusual chemical shifts (i)

There are no statistically unusual chemical shifts.

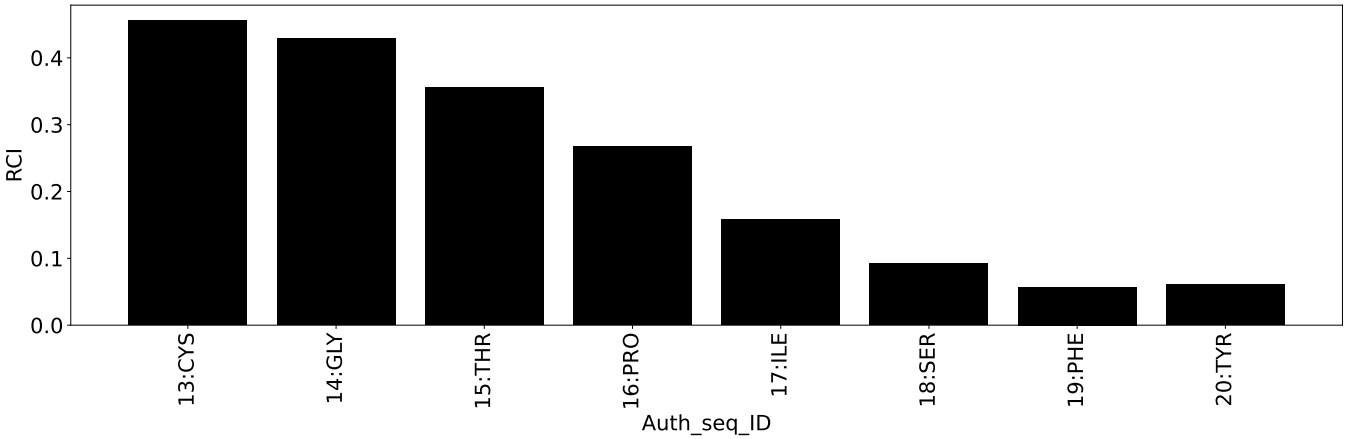
7.1.5 Random Coil Index (RCI) plots (i)

The image below reports *random coil index* values for the protein chains in the structure. The height of each bar gives a probability of a given residue to be disordered, as predicted from the available chemical shifts and the amino acid sequence. A value above 0.2 is an indication of significant predicted disorder. The colour of the bar shows whether the residue is in the well-defined core (black) or in the ill-defined residue ranges (cyan), as described in section 2 on ensemble composition.If well-defined core and ill-defined regions are not identified then it is shown as gray bars.

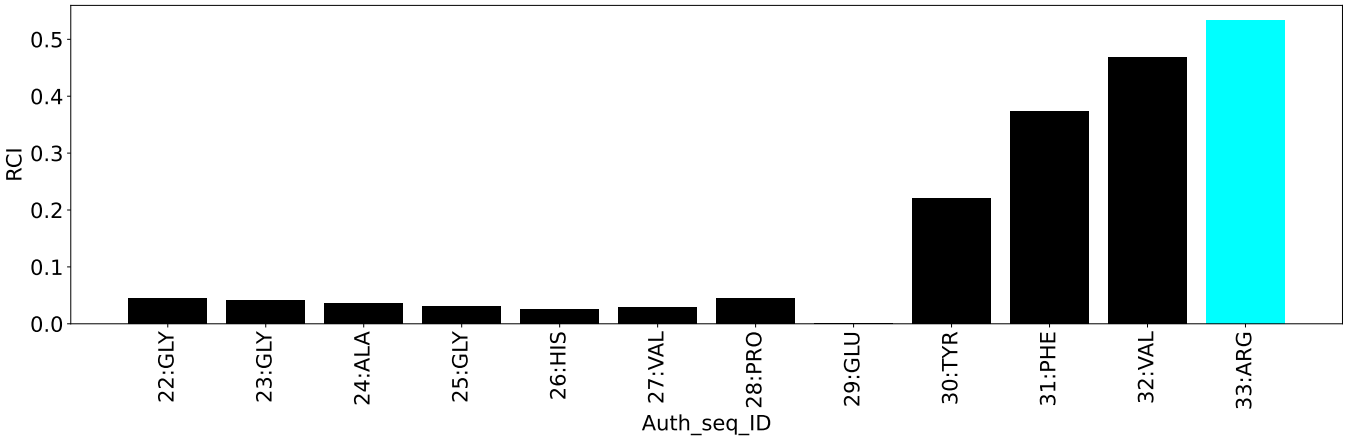
Random coil index (RCI) for chain A:



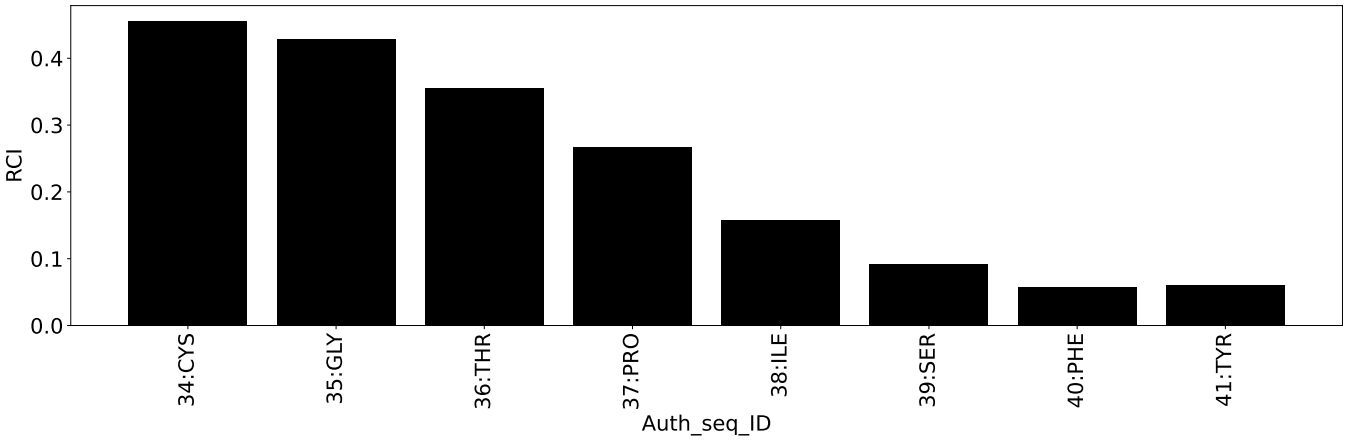
Random coil index (RCI) for chain B:



Random coil index (RCI) for chain C:



Random coil index (RCI) for chain D:



8 NMR restraints analysis

8.1 Conformationally restricting restraints

The following table provides the summary of experimentally observed NMR restraints in different categories. Restraints are classified into different categories based on the sequence separation of the atoms involved.

Description	Value
Total distance restraints	1312
Intra-residue (|i-j|=0)	552
Sequential (|i-j|=1)	332
Medium range (|i-j|>1 and |i-j|<5)	60
Long range (|i-j|≥5)	8
Inter-chain	360
Hydrogen bond restraints	0
Disulfide bond restraints	0
Total dihedral-angle restraints	0
Number of unmapped restraints	0
Number of restraints per residue	31.2
Number of long range restraints per residue[1]	0.2

[1]Long range hydrogen bonds and disulfide bonds are counted as long range restraints while calculating the number of long range restraints per residue

8.2 Residual restraint violations

This section provides the overview of the restraint violations analysis. The violations are binned as small, medium and large violations based on its absolute value. Average number of violations per model is calculated by dividing the total number of violations in each bin by the size of the ensemble.

8.2.1 Average number of distance violations per model

Distance violations less than 0.1 Å
are not included in the calculation.

Bins (Å)	Average number of violations per model	Max (Å)
0.1-0.2 (Small)	26.2	0.2
0.2-0.5 (Medium)	64.1	0.5
>0.5 (Large)	128.2	4.87

8.2.2 Average number of dihedral-angle violations per model (i)

Dihedral-angle violations less than 1° are not included in the calculation.There are no dihedral-angle violations

9 Distance violation analysis (i)

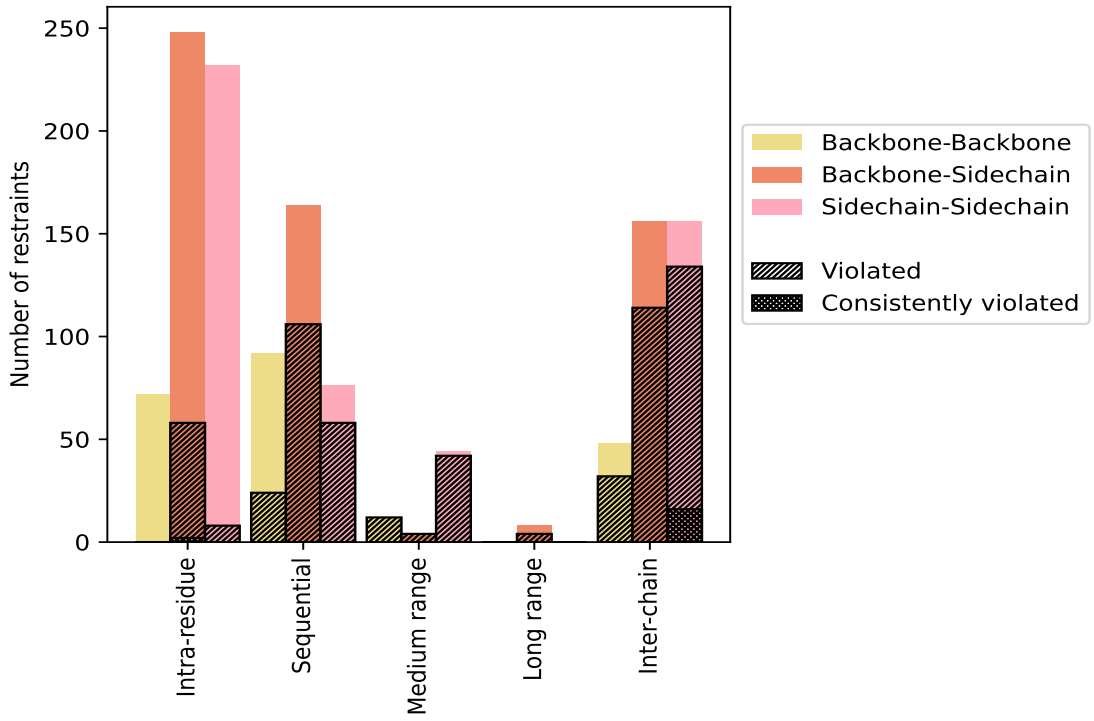
9.1 Summary of distance violations (i)

The following table shows the summary of distance violations in different restraint categories based on the sequence separation of the atoms involved. Each category is further sub-divided into three sub-categories based on the atoms involved. Violations less than 0.1 Å are not included in the statistics.

Restraints type	Count	%[1]	Violated[3]			Consistently Violated[4]		
			Count	%[2]	%[1]	Count	%[2]	%[1]
Intra-residue (|i-j|=0)	552	42.1	66	12.0	5.0	2	0.4	0.2
Backbone-Backbone	72	5.5	0	0.0	0.0	0	0.0	0.0
Backbone-Sidechain	248	18.9	58	23.4	4.4	2	0.8	0.2
Sidechain-Sidechain	232	17.7	8	3.4	0.6	0	0.0	0.0
Sequential (|i-j|=1)	332	25.3	188	56.6	14.3	0	0.0	0.0
Backbone-Backbone	92	7.0	24	26.1	1.8	0	0.0	0.0
Backbone-Sidechain	164	12.5	106	64.6	8.1	0	0.0	0.0
Sidechain-Sidechain	76	5.8	58	76.3	4.4	0	0.0	0.0
Medium range (|i-j|>1 & |i-j|<5)	60	4.6	58	96.7	4.4	0	0.0	0.0
Backbone-Backbone	12	0.9	12	100.0	0.9	0	0.0	0.0
Backbone-Sidechain	4	0.3	4	100.0	0.3	0	0.0	0.0
Sidechain-Sidechain	44	3.4	42	95.5	3.2	0	0.0	0.0
Long range (|i-j|≥5)	8	0.6	4	50.0	0.3	0	0.0	0.0
Backbone-Backbone	0	0.0	0	0.0	0.0	0	0.0	0.0
Backbone-Sidechain	8	0.6	4	50.0	0.3	0	0.0	0.0
Sidechain-Sidechain	0	0.0	0	0.0	0.0	0	0.0	0.0
Inter-chain	360	27.4	280	77.8	21.3	16	4.4	1.2
Backbone-Backbone	48	3.7	32	66.7	2.4	0	0.0	0.0
Backbone-Sidechain	156	11.9	114	73.1	8.7	0	0.0	0.0
Sidechain-Sidechain	156	11.9	134	85.9	10.2	16	10.3	1.2
Hydrogen bond	0	0.0	0	0.0	0.0	0	0.0	0.0
Disulfide bond	0	0.0	0	0.0	0.0	0	0.0	0.0
Total	1312	100.0	596	45.4	45.4	18	1.4	1.4
Backbone-Backbone	224	17.1	68	30.4	5.2	0	0.0	0.0
Backbone-Sidechain	580	44.2	286	49.3	21.8	2	0.3	0.2
Sidechain-Sidechain	508	38.7	242	47.6	18.4	16	3.1	1.2

[1] percentage calculated with respect to the total number of distance restraints, [2] percentage calculated with respect to the number of restraints in a particular restraint category, [3] violated in at least one model, [4] violated in all the models

9.1.1 Bar chart : Distribution of distance restraints and violations



Violated and consistently violated restraints are shown using different hatch patterns in their respective categories. The hydrogen bonds and disulfied bonds are counted in their appropriate category on the x-axis

9.2 Distance violation statistics for each model

The following table provides the distance violation statistics for each model in the ensemble. Violations less than 0.1 Å
are not included in the statistics.

Model ID	Number of violations						Mean (Å)	Max (Å)	SD[6] (Å)	Median (Å)
	IR[1]	SQ[2]	MR[3]	LR[4]	IC[5]	Total				
1	24	76	18	0	142	260	0.84	3.12	0.71	0.58
2	20	56	24	0	98	198	0.78	2.9	0.59	0.63
3	18	56	38	0	122	234	0.89	3.0	0.73	0.64
4	16	62	24	0	128	230	0.75	2.52	0.55	0.55
5	18	58	20	0	148	244	0.99	4.87	0.95	0.68
6	20	46	14	0	114	194	0.78	3.34	0.62	0.66
7	28	60	30	0	100	218	0.82	4.44	0.71	0.59
8	18	56	24	0	110	208	0.87	2.88	0.62	0.74
9	18	72	36	0	112	238	0.94	3.98	0.86	0.59
10	22	52	30	0	104	208	0.99	3.36	0.73	0.79

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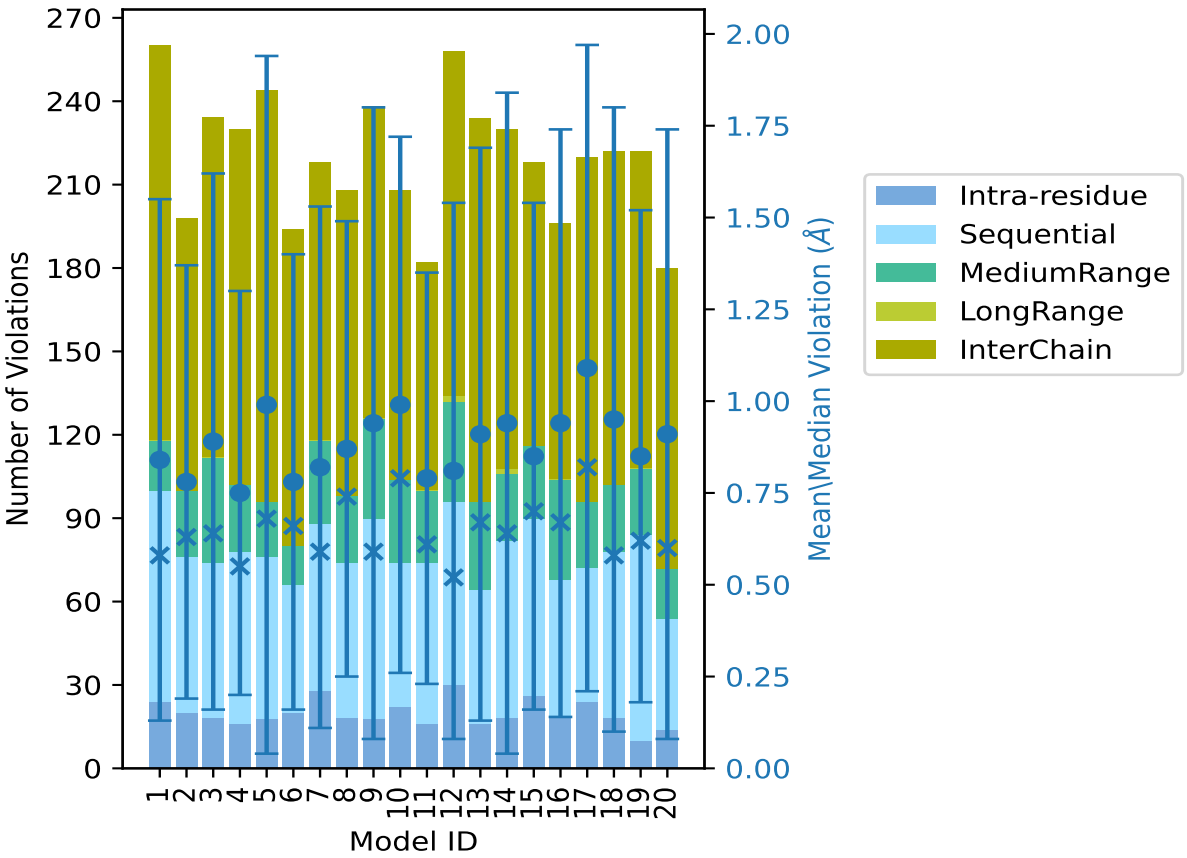
Model ID	Number of violations						Mean (Å)	Max (Å)	SD[6] (Å)	Median (Å)
	IR[1]	SQ[2]	MR[3]	LR[4]	IC[5]	Total				
11	16	58	26	0	82	182	0.79	2.26	0.56	0.61
12	30	66	36	2	124	258	0.81	3.41	0.73	0.52
13	16	48	32	0	138	234	0.91	3.19	0.78	0.67
14	18	64	24	2	122	230	0.94	4.26	0.9	0.64
15	26	64	26	0	102	218	0.85	3.2	0.69	0.7
16	18	50	36	0	92	196	0.94	3.44	0.8	0.67
17	24	48	24	0	124	220	1.09	4.39	0.88	0.82
18	18	60	24	0	120	222	0.95	4.25	0.85	0.58
19	10	74	24	0	114	222	0.85	3.22	0.67	0.62
20	14	40	18	0	108	180	0.91	4.63	0.83	0.6

[1]Intra-residue restraints, [2]Sequential restraints, [3]Medium range restraints, [4]Long range restraints, [5]Inter-chain restraints, [6]Standard deviation

9.2.1 Bar graph : Distance Violation statistics for each model



The mean(dot),median(x) and the standard deviation are shown in blue with respect to the y axis on the right

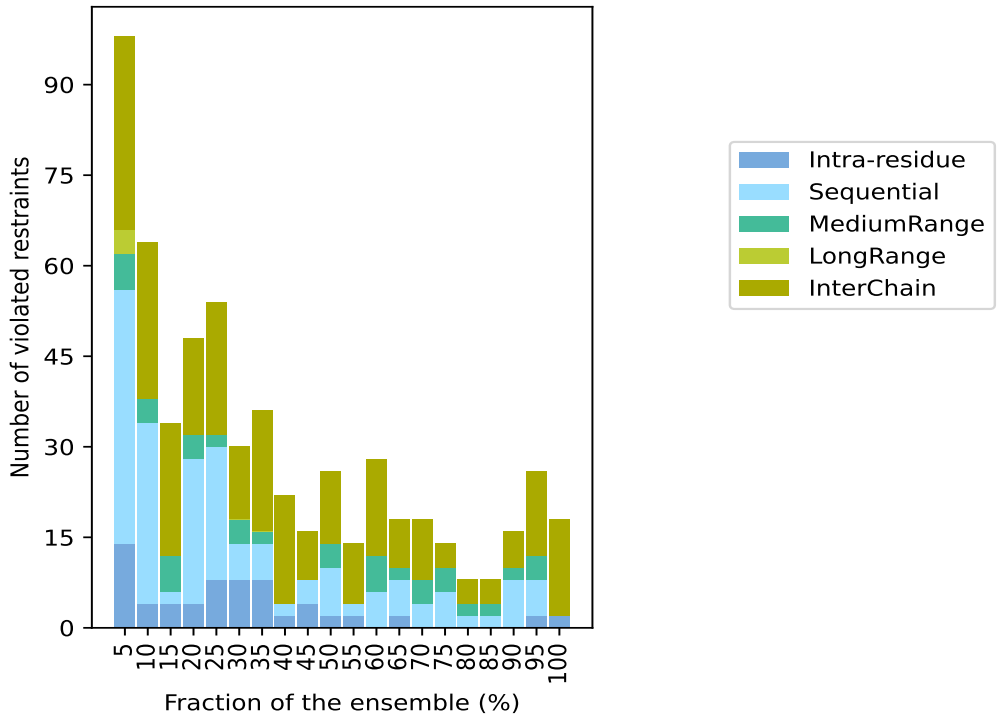
9.3 Distance violation statistics for the ensemble (i)

Violation analysis may find that some restraints are violated in few models and some are violated in most of models. The following table provides this information as number of violated restraints for a given fraction of the ensemble. In total, 716(IR:486, SQ:144, MR:2, LR:4, IC:80) restraints are not violated in the ensemble.

Number of violated restraints						Fraction of the ensemble	
IR[1]	SQ[2]	MR[3]	LR[4]	IC[5]	Total	Count[6]	%
14	42	6	4	32	98	1	5.0
4	30	4	0	26	64	2	10.0
4	2	6	0	22	34	3	15.0
4	24	4	0	16	48	4	20.0
8	22	2	0	22	54	5	25.0
8	6	4	0	12	30	6	30.0
8	6	2	0	20	36	7	35.0
2	2	0	0	18	22	8	40.0
4	4	0	0	8	16	9	45.0
2	8	4	0	12	26	10	50.0
2	2	0	0	10	14	11	55.0
0	6	6	0	16	28	12	60.0
2	6	2	0	8	18	13	65.0
0	4	4	0	10	18	14	70.0
0	6	4	0	4	14	15	75.0
0	2	2	0	4	8	16	80.0
0	2	2	0	4	8	17	85.0
0	8	2	0	6	16	18	90.0
2	6	4	0	14	26	19	95.0
2	0	0	0	16	18	20	100.0

[1]Intra-residue restraints, [2]Sequential restraints, [3]Medium range restraints, [4]Long range restraints, [5]Inter-chain restraints, [6] Number of models with violations

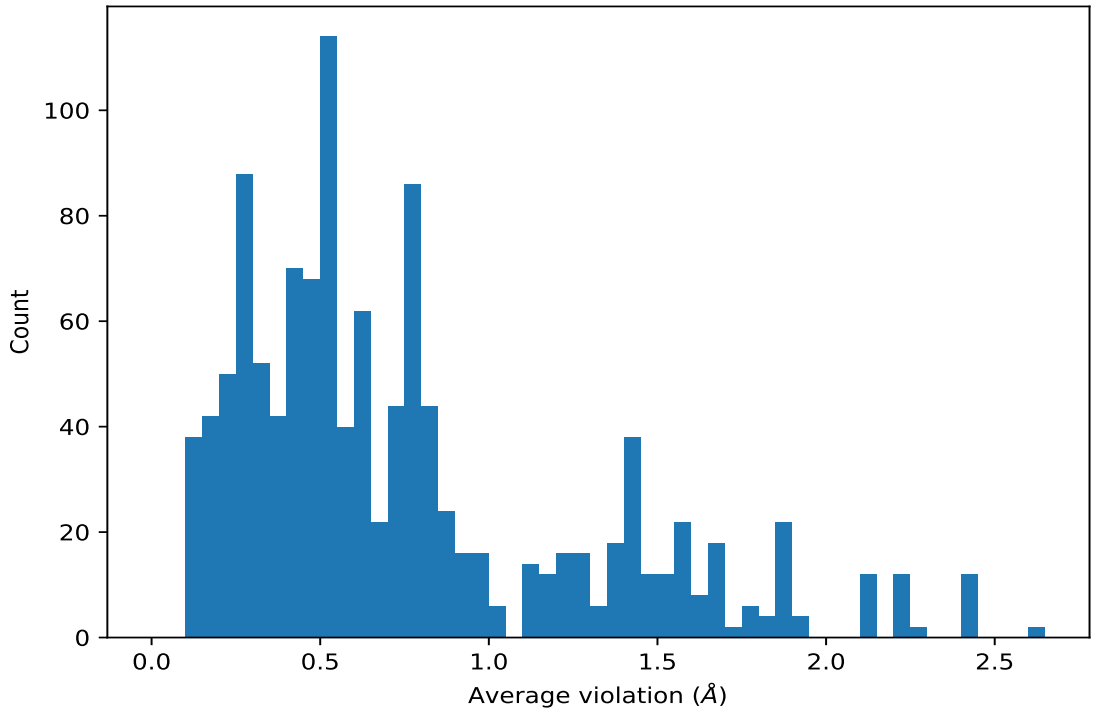
9.3.1 Bar graph : Distance violation statistics for the ensemble



9.4 Most violated distance restraints in the ensemble

9.4.1 Histogram : Distribution of mean distance violations

The following histogram shows the distribution of the average value of the violation. The average is calculated for each restraint that is violated in more than one model over all the violated models in the ensemble



9.4.2 Table: Most violated distance restraints

The following table provides the mean and the standard deviation of theviolation for each restraint sorted by number of violated models and the mean value. The Key (restraint list ID, restraint ID) is the unique identifier for a given restraint. Rows with same key represent combinatorial or ambiguous restraints and are counted as a single restraint.

Key	Atom-1	Atom-2	Models[1]	Mean (Å)	SD[1] (Å)	Median (Å)
(1,857)	2:19:B:PHE:HB3	2:38:D:ILE:HG13	20	2.6	1.08	2.38
(1,1176)	2:19:B:PHE:HB3	2:38:D:ILE:HG13	20	2.6	1.08	2.38
(1,17)	2:19:B:PHE:HB2	2:38:D:ILE:HG13	20	2.29	1.09	1.94
(1,664)	2:19:B:PHE:HB2	2:38:D:ILE:HG13	20	2.29	1.09	1.94
(1,201)	2:17:B:ILE:HG13	2:40:D:PHE:HB3	20	1.71	0.71	1.77
(1,520)	2:17:B:ILE:HG13	2:40:D:PHE:HB3	20	1.71	0.71	1.77
(1,932)	1:33:C:ARG:HA	1:33:C:ARG:HE	20	1.6	0.46	1.53
(1,1251)	1:33:C:ARG:HA	1:33:C:ARG:HE	20	1.6	0.46	1.53
(1,774)	1:29:C:GLU:HG2	2:41:D:TYR:HB2	20	1.56	0.5	1.49
(1,774)	1:29:C:GLU:HG2	2:41:D:TYR:HB3	20	1.56	0.5	1.49
(1,1093)	1:29:C:GLU:HG2	2:41:D:TYR:HB2	20	1.56	0.5	1.49
(1,1093)	1:29:C:GLU:HG2	2:41:D:TYR:HB3	20	1.56	0.5	1.49
(1,118)	1:8:A:GLU:HG2	2:20:B:TYR:HB2	20	1.46	0.57	1.43
(1,118)	1:8:A:GLU:HG2	2:20:B:TYR:HB3	20	1.46	0.57	1.43
(1,437)	1:8:A:GLU:HG2	2:20:B:TYR:HB2	20	1.46	0.57	1.43
(1,437)	1:8:A:GLU:HG2	2:20:B:TYR:HB3	20	1.46	0.57	1.43

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Key	Atom-1	Atom-2	Models[1]	Mean (Å)	SD[1] (Å)	Median (Å)
(1,117)	1:8:A:GLU:HG2	2:20:B:TYR:HD1	20	1.41	0.59	1.4
(1,117)	1:8:A:GLU:HG2	2:20:B:TYR:HD2	20	1.41	0.59	1.4
(1,436)	1:8:A:GLU:HG2	2:20:B:TYR:HD1	20	1.41	0.59	1.4
(1,436)	1:8:A:GLU:HG2	2:20:B:TYR:HD2	20	1.41	0.59	1.4
(1,8)	2:17:B:ILE:HG13	2:40:D:PHE:HB2	20	1.4	0.7	1.36
(1,673)	2:17:B:ILE:HG13	2:40:D:PHE:HB2	20	1.4	0.7	1.36
(1,856)	1:24:C:ALA:HB1	2:40:D:PHE:HB3	20	0.76	0.38	0.68
(1,856)	1:24:C:ALA:HB2	2:40:D:PHE:HB3	20	0.76	0.38	0.68
(1,856)	1:24:C:ALA:HB3	2:40:D:PHE:HB3	20	0.76	0.38	0.68
(1,1175)	1:24:C:ALA:HB1	2:40:D:PHE:HB3	20	0.76	0.38	0.68
(1,1175)	1:24:C:ALA:HB2	2:40:D:PHE:HB3	20	0.76	0.38	0.68
(1,1175)	1:24:C:ALA:HB3	2:40:D:PHE:HB3	20	0.76	0.38	0.68
(1,9)	2:17:B:ILE:HD11	2:40:D:PHE:HE1	19	2.23	1.05	2.51
(1,9)	2:17:B:ILE:HD11	2:40:D:PHE:HE2	19	2.23	1.05	2.51
(1,9)	2:17:B:ILE:HD12	2:40:D:PHE:HE1	19	2.23	1.05	2.51
(1,9)	2:17:B:ILE:HD12	2:40:D:PHE:HE2	19	2.23	1.05	2.51
(1,9)	2:17:B:ILE:HD13	2:40:D:PHE:HE1	19	2.23	1.05	2.51
(1,9)	2:17:B:ILE:HD13	2:40:D:PHE:HE2	19	2.23	1.05	2.51
(1,674)	2:17:B:ILE:HD11	2:40:D:PHE:HE1	19	2.23	1.05	2.51
(1,674)	2:17:B:ILE:HD11	2:40:D:PHE:HE2	19	2.23	1.05	2.51
(1,674)	2:17:B:ILE:HD12	2:40:D:PHE:HE1	19	2.23	1.05	2.51
(1,674)	2:17:B:ILE:HD12	2:40:D:PHE:HE2	19	2.23	1.05	2.51
(1,674)	2:17:B:ILE:HD13	2:40:D:PHE:HE1	19	2.23	1.05	2.51
(1,674)	2:17:B:ILE:HD13	2:40:D:PHE:HE2	19	2.23	1.05	2.51
(1,276)	1:12:A:ARG:HA	1:12:A:ARG:HE	19	1.6	0.58	1.47
(1,595)	1:12:A:ARG:HA	1:12:A:ARG:HE	19	1.6	0.58	1.47
(1,96)	1:6:A:VAL:HG11	2:20:B:TYR:HE1	19	1.58	0.59	1.74
(1,96)	1:6:A:VAL:HG11	2:20:B:TYR:HE2	19	1.58	0.59	1.74
(1,96)	1:6:A:VAL:HG12	2:20:B:TYR:HE1	19	1.58	0.59	1.74
(1,96)	1:6:A:VAL:HG12	2:20:B:TYR:HE2	19	1.58	0.59	1.74
(1,96)	1:6:A:VAL:HG13	2:20:B:TYR:HE1	19	1.58	0.59	1.74
(1,96)	1:6:A:VAL:HG13	2:20:B:TYR:HE2	19	1.58	0.59	1.74
(1,415)	1:6:A:VAL:HG11	2:20:B:TYR:HE1	19	1.58	0.59	1.74
(1,415)	1:6:A:VAL:HG11	2:20:B:TYR:HE2	19	1.58	0.59	1.74
(1,415)	1:6:A:VAL:HG12	2:20:B:TYR:HE1	19	1.58	0.59	1.74
(1,415)	1:6:A:VAL:HG12	2:20:B:TYR:HE2	19	1.58	0.59	1.74
(1,415)	1:6:A:VAL:HG13	2:20:B:TYR:HE1	19	1.58	0.59	1.74
(1,415)	1:6:A:VAL:HG13	2:20:B:TYR:HE2	19	1.58	0.59	1.74
(1,184)	1:7:A:PRO:HB3	1:10:A:PHE:HE1	19	1.54	0.99	1.34
(1,184)	1:7:A:PRO:HB3	1:10:A:PHE:HE2	19	1.54	0.99	1.34
(1,503)	1:7:A:PRO:HB3	1:10:A:PHE:HE1	19	1.54	0.99	1.34
(1,503)	1:7:A:PRO:HB3	1:10:A:PHE:HE2	19	1.54	0.99	1.34

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Key	Atom-1	Atom-2	Models[1]	Mean (Å)	SD[1] (Å)	Median (Å)
(1,752)	1:27:C:VAL:HG11	2:41:D:TYR:HE1	19	1.43	0.58	1.29
(1,752)	1:27:C:VAL:HG11	2:41:D:TYR:HE2	19	1.43	0.58	1.29
(1,752)	1:27:C:VAL:HG12	2:41:D:TYR:HE1	19	1.43	0.58	1.29
(1,752)	1:27:C:VAL:HG12	2:41:D:TYR:HE2	19	1.43	0.58	1.29
(1,752)	1:27:C:VAL:HG13	2:41:D:TYR:HE1	19	1.43	0.58	1.29
(1,752)	1:27:C:VAL:HG13	2:41:D:TYR:HE2	19	1.43	0.58	1.29
(1,1071)	1:27:C:VAL:HG11	2:41:D:TYR:HE1	19	1.43	0.58	1.29
(1,1071)	1:27:C:VAL:HG11	2:41:D:TYR:HE2	19	1.43	0.58	1.29
(1,1071)	1:27:C:VAL:HG12	2:41:D:TYR:HE1	19	1.43	0.58	1.29
(1,1071)	1:27:C:VAL:HG12	2:41:D:TYR:HE2	19	1.43	0.58	1.29
(1,1071)	1:27:C:VAL:HG13	2:41:D:TYR:HE1	19	1.43	0.58	1.29
(1,1071)	1:27:C:VAL:HG13	2:41:D:TYR:HE2	19	1.43	0.58	1.29
(1,773)	1:29:C:GLU:HG2	2:41:D:TYR:HD1	19	1.24	0.42	1.19
(1,773)	1:29:C:GLU:HG2	2:41:D:TYR:HD2	19	1.24	0.42	1.19
(1,1092)	1:29:C:GLU:HG2	2:41:D:TYR:HD1	19	1.24	0.42	1.19
(1,1092)	1:29:C:GLU:HG2	2:41:D:TYR:HD2	19	1.24	0.42	1.19
(1,840)	1:28:C:PRO:HB3	1:31:C:PHE:HE1	19	1.15	0.8	0.94
(1,840)	1:28:C:PRO:HB3	1:31:C:PHE:HE2	19	1.15	0.8	0.94
(1,1159)	1:28:C:PRO:HB3	1:31:C:PHE:HE1	19	1.15	0.8	0.94
(1,1159)	1:28:C:PRO:HB3	1:31:C:PHE:HE2	19	1.15	0.8	0.94
(1,717)	2:34:D:CYS:HB2	2:35:D:GLY:HA3	19	0.81	0.21	0.81
(1,1036)	2:34:D:CYS:HB2	2:35:D:GLY:HA3	19	0.81	0.21	0.81
(1,200)	1:3:A:ALA:HB1	2:19:B:PHE:HB3	19	0.71	0.36	0.54
(1,200)	1:3:A:ALA:HB2	2:19:B:PHE:HB3	19	0.71	0.36	0.54
(1,200)	1:3:A:ALA:HB3	2:19:B:PHE:HB3	19	0.71	0.36	0.54
(1,519)	1:3:A:ALA:HB1	2:19:B:PHE:HB3	19	0.71	0.36	0.54
(1,519)	1:3:A:ALA:HB2	2:19:B:PHE:HB3	19	0.71	0.36	0.54
(1,519)	1:3:A:ALA:HB3	2:19:B:PHE:HB3	19	0.71	0.36	0.54
(1,732)	2:34:D:CYS:HB2	2:35:D:GLY:H	19	0.7	0.19	0.74
(1,1051)	2:34:D:CYS:HB2	2:35:D:GLY:H	19	0.7	0.19	0.74
(1,119)	1:8:A:GLU:HG2	2:19:B:PHE:HA	19	0.58	0.29	0.52
(1,438)	1:8:A:GLU:HG2	2:19:B:PHE:HA	19	0.58	0.29	0.52
(1,775)	1:29:C:GLU:HG2	2:40:D:PHE:HA	19	0.53	0.17	0.59
(1,1094)	1:29:C:GLU:HG2	2:40:D:PHE:HA	19	0.53	0.17	0.59
(1,724)	1:27:C:VAL:HG11	1:28:C:PRO:HG2	19	0.52	0.27	0.44
(1,724)	1:27:C:VAL:HG11	1:28:C:PRO:HG3	19	0.52	0.27	0.44
(1,724)	1:27:C:VAL:HG12	1:28:C:PRO:HG2	19	0.52	0.27	0.44
(1,724)	1:27:C:VAL:HG12	1:28:C:PRO:HG3	19	0.52	0.27	0.44
(1,724)	1:27:C:VAL:HG13	1:28:C:PRO:HG2	19	0.52	0.27	0.44
(1,724)	1:27:C:VAL:HG13	1:28:C:PRO:HG3	19	0.52	0.27	0.44
(1,1043)	1:27:C:VAL:HG11	1:28:C:PRO:HG2	19	0.52	0.27	0.44
(1,1043)	1:27:C:VAL:HG11	1:28:C:PRO:HG3	19	0.52	0.27	0.44

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Key	Atom-1	Atom-2	Models[1]	Mean (Å)	SD[1] (Å)	Median (Å)
(1,1043)	1:27:C:VAL:HG12	1:28:C:PRO:HG2	19	0.52	0.27	0.44
(1,1043)	1:27:C:VAL:HG12	1:28:C:PRO:HG3	19	0.52	0.27	0.44
(1,1043)	1:27:C:VAL:HG13	1:28:C:PRO:HG2	19	0.52	0.27	0.44
(1,1043)	1:27:C:VAL:HG13	1:28:C:PRO:HG3	19	0.52	0.27	0.44
(1,18)	2:19:B:PHE:HE1	2:38:D:ILE:HD11	18	2.44	1.11	2.9
(1,18)	2:19:B:PHE:HE1	2:38:D:ILE:HD12	18	2.44	1.11	2.9
(1,18)	2:19:B:PHE:HE1	2:38:D:ILE:HD13	18	2.44	1.11	2.9
(1,18)	2:19:B:PHE:HE2	2:38:D:ILE:HD11	18	2.44	1.11	2.9
(1,18)	2:19:B:PHE:HE2	2:38:D:ILE:HD12	18	2.44	1.11	2.9
(1,18)	2:19:B:PHE:HE2	2:38:D:ILE:HD13	18	2.44	1.11	2.9
(1,665)	2:19:B:PHE:HE1	2:38:D:ILE:HD11	18	2.44	1.11	2.9
(1,665)	2:19:B:PHE:HE1	2:38:D:ILE:HD12	18	2.44	1.11	2.9
(1,665)	2:19:B:PHE:HE1	2:38:D:ILE:HD13	18	2.44	1.11	2.9
(1,665)	2:19:B:PHE:HE2	2:38:D:ILE:HD11	18	2.44	1.11	2.9
(1,665)	2:19:B:PHE:HE2	2:38:D:ILE:HD12	18	2.44	1.11	2.9
(1,665)	2:19:B:PHE:HE2	2:38:D:ILE:HD13	18	2.44	1.11	2.9
(1,172)	1:7:A:PRO:HB2	1:10:A:PHE:HE1	18	1.94	0.73	1.98
(1,172)	1:7:A:PRO:HB2	1:10:A:PHE:HE2	18	1.94	0.73	1.98
(1,491)	1:7:A:PRO:HB2	1:10:A:PHE:HE1	18	1.94	0.73	1.98
(1,491)	1:7:A:PRO:HB2	1:10:A:PHE:HE2	18	1.94	0.73	1.98
(1,739)	1:31:C:PHE:HD1	2:35:D:GLY:HA3	18	1.47	0.78	1.45
(1,739)	1:31:C:PHE:HD2	2:35:D:GLY:HA3	18	1.47	0.78	1.45
(1,1058)	1:31:C:PHE:HD1	2:35:D:GLY:HA3	18	1.47	0.78	1.45
(1,1058)	1:31:C:PHE:HD2	2:35:D:GLY:HA3	18	1.47	0.78	1.45
(1,61)	2:13:B:CYS:HB2	2:14:B:GLY:HA3	18	0.81	0.23	0.88
(1,380)	2:13:B:CYS:HB2	2:14:B:GLY:HA3	18	0.81	0.23	0.88
(1,127)	1:8:A:GLU:HB3	2:20:B:TYR:HD1	18	0.74	0.33	0.7
(1,127)	1:8:A:GLU:HB3	2:20:B:TYR:HD2	18	0.74	0.33	0.7
(1,446)	1:8:A:GLU:HB3	2:20:B:TYR:HD1	18	0.74	0.33	0.7
(1,446)	1:8:A:GLU:HB3	2:20:B:TYR:HD2	18	0.74	0.33	0.7
(1,68)	1:6:A:VAL:HG11	1:7:A:PRO:HG2	18	0.48	0.18	0.52
(1,68)	1:6:A:VAL:HG11	1:7:A:PRO:HG3	18	0.48	0.18	0.52
(1,68)	1:6:A:VAL:HG12	1:7:A:PRO:HG2	18	0.48	0.18	0.52
(1,68)	1:6:A:VAL:HG12	1:7:A:PRO:HG3	18	0.48	0.18	0.52
(1,68)	1:6:A:VAL:HG13	1:7:A:PRO:HG2	18	0.48	0.18	0.52
(1,68)	1:6:A:VAL:HG13	1:7:A:PRO:HG3	18	0.48	0.18	0.52
(1,387)	1:6:A:VAL:HG11	1:7:A:PRO:HG2	18	0.48	0.18	0.52
(1,387)	1:6:A:VAL:HG11	1:7:A:PRO:HG3	18	0.48	0.18	0.52
(1,387)	1:6:A:VAL:HG12	1:7:A:PRO:HG2	18	0.48	0.18	0.52
(1,387)	1:6:A:VAL:HG12	1:7:A:PRO:HG3	18	0.48	0.18	0.52
(1,387)	1:6:A:VAL:HG13	1:7:A:PRO:HG2	18	0.48	0.18	0.52
(1,387)	1:6:A:VAL:HG13	1:7:A:PRO:HG3	18	0.48	0.18	0.52

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Key	Atom-1	Atom-2	Models[1]	Mean (Å)	SD[1] (Å)	Median (Å)
(1,798)	1:27:C:VAL:HG11	1:28:C:PRO:HB3	18	0.48	0.22	0.48
(1,798)	1:27:C:VAL:HG12	1:28:C:PRO:HB3	18	0.48	0.22	0.48
(1,798)	1:27:C:VAL:HG13	1:28:C:PRO:HB3	18	0.48	0.22	0.48
(1,1117)	1:27:C:VAL:HG11	1:28:C:PRO:HB3	18	0.48	0.22	0.48
(1,1117)	1:27:C:VAL:HG12	1:28:C:PRO:HB3	18	0.48	0.22	0.48
(1,1117)	1:27:C:VAL:HG13	1:28:C:PRO:HB3	18	0.48	0.22	0.48
(1,142)	1:6:A:VAL:HG11	1:7:A:PRO:HB3	18	0.46	0.17	0.47
(1,142)	1:6:A:VAL:HG12	1:7:A:PRO:HB3	18	0.46	0.17	0.47
(1,142)	1:6:A:VAL:HG13	1:7:A:PRO:HB3	18	0.46	0.17	0.47
(1,461)	1:6:A:VAL:HG11	1:7:A:PRO:HB3	18	0.46	0.17	0.47
(1,461)	1:6:A:VAL:HG12	1:7:A:PRO:HB3	18	0.46	0.17	0.47
(1,461)	1:6:A:VAL:HG13	1:7:A:PRO:HB3	18	0.46	0.17	0.47
(1,852)	2:36:D:THR:HG21	2:38:D:ILE:HD11	17	1.87	0.62	1.58
(1,852)	2:36:D:THR:HG21	2:38:D:ILE:HD12	17	1.87	0.62	1.58
(1,852)	2:36:D:THR:HG21	2:38:D:ILE:HD13	17	1.87	0.62	1.58
(1,852)	2:36:D:THR:HG22	2:38:D:ILE:HD11	17	1.87	0.62	1.58
(1,852)	2:36:D:THR:HG22	2:38:D:ILE:HD12	17	1.87	0.62	1.58
(1,852)	2:36:D:THR:HG22	2:38:D:ILE:HD13	17	1.87	0.62	1.58
(1,852)	2:36:D:THR:HG23	2:38:D:ILE:HD11	17	1.87	0.62	1.58
(1,852)	2:36:D:THR:HG23	2:38:D:ILE:HD12	17	1.87	0.62	1.58
(1,852)	2:36:D:THR:HG23	2:38:D:ILE:HD13	17	1.87	0.62	1.58
(1,1171)	2:36:D:THR:HG21	2:38:D:ILE:HD11	17	1.87	0.62	1.58
(1,1171)	2:36:D:THR:HG21	2:38:D:ILE:HD12	17	1.87	0.62	1.58
(1,1171)	2:36:D:THR:HG21	2:38:D:ILE:HD13	17	1.87	0.62	1.58
(1,1171)	2:36:D:THR:HG22	2:38:D:ILE:HD11	17	1.87	0.62	1.58
(1,1171)	2:36:D:THR:HG22	2:38:D:ILE:HD12	17	1.87	0.62	1.58
(1,1171)	2:36:D:THR:HG22	2:38:D:ILE:HD13	17	1.87	0.62	1.58
(1,1171)	2:36:D:THR:HG23	2:38:D:ILE:HD11	17	1.87	0.62	1.58
(1,1171)	2:36:D:THR:HG23	2:38:D:ILE:HD12	17	1.87	0.62	1.58
(1,1171)	2:36:D:THR:HG23	2:38:D:ILE:HD13	17	1.87	0.62	1.58
(1,147)	1:4:A:GLY:HA2	1:5:A:HIS:HD2	17	1.43	0.59	1.52
(1,466)	1:4:A:GLY:HA2	1:5:A:HIS:HD2	17	1.43	0.59	1.52
(1,164)	1:1:A:GLY:H	2:20:B:TYR:HD1	17	0.95	0.36	0.9
(1,164)	1:1:A:GLY:H	2:20:B:TYR:HD2	17	0.95	0.36	0.9
(1,483)	1:1:A:GLY:H	2:20:B:TYR:HD1	17	0.95	0.36	0.9
(1,483)	1:1:A:GLY:H	2:20:B:TYR:HD2	17	0.95	0.36	0.9
(1,783)	1:29:C:GLU:HB3	2:41:D:TYR:HD1	17	0.77	0.38	0.7
(1,783)	1:29:C:GLU:HB3	2:41:D:TYR:HD2	17	0.77	0.38	0.7
(1,1102)	1:29:C:GLU:HB3	2:41:D:TYR:HD1	17	0.77	0.38	0.7
(1,1102)	1:29:C:GLU:HB3	2:41:D:TYR:HD2	17	0.77	0.38	0.7
(1,828)	1:28:C:PRO:HB2	1:31:C:PHE:HE1	16	1.46	0.85	1.52
(1,828)	1:28:C:PRO:HB2	1:31:C:PHE:HE2	16	1.46	0.85	1.52

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Key	Atom-1	Atom-2	Models[1]	Mean (Å)	SD[1] (Å)	Median (Å)
(1,1147)	1:28:C:PRO:HB2	1:31:C:PHE:HE1	16	1.46	0.85	1.52
(1,1147)	1:28:C:PRO:HB2	1:31:C:PHE:HE2	16	1.46	0.85	1.52
(1,727)	1:27:C:VAL:HB	2:41:D:TYR:HE1	16	1.21	0.7	1.16
(1,727)	1:27:C:VAL:HB	2:41:D:TYR:HE2	16	1.21	0.7	1.16
(1,1046)	1:27:C:VAL:HB	2:41:D:TYR:HE1	16	1.21	0.7	1.16
(1,1046)	1:27:C:VAL:HB	2:41:D:TYR:HE2	16	1.21	0.7	1.16
(1,76)	2:13:B:CYS:HB2	2:14:B:GLY:H	16	0.67	0.19	0.7
(1,395)	2:13:B:CYS:HB2	2:14:B:GLY:H	16	0.67	0.19	0.7
(1,50)	1:6:A:VAL:HG21	2:20:B:TYR:HE1	16	0.49	0.19	0.52
(1,50)	1:6:A:VAL:HG21	2:20:B:TYR:HE2	16	0.49	0.19	0.52
(1,50)	1:6:A:VAL:HG22	2:20:B:TYR:HE1	16	0.49	0.19	0.52
(1,50)	1:6:A:VAL:HG22	2:20:B:TYR:HE2	16	0.49	0.19	0.52
(1,50)	1:6:A:VAL:HG23	2:20:B:TYR:HE1	16	0.49	0.19	0.52
(1,50)	1:6:A:VAL:HG23	2:20:B:TYR:HE2	16	0.49	0.19	0.52
(1,369)	1:6:A:VAL:HG21	2:20:B:TYR:HE1	16	0.49	0.19	0.52
(1,369)	1:6:A:VAL:HG21	2:20:B:TYR:HE2	16	0.49	0.19	0.52
(1,369)	1:6:A:VAL:HG22	2:20:B:TYR:HE1	16	0.49	0.19	0.52
(1,369)	1:6:A:VAL:HG22	2:20:B:TYR:HE2	16	0.49	0.19	0.52
(1,369)	1:6:A:VAL:HG23	2:20:B:TYR:HE1	16	0.49	0.19	0.52
(1,369)	1:6:A:VAL:HG23	2:20:B:TYR:HE2	16	0.49	0.19	0.52
(1,196)	2:15:B:THR:HG21	2:17:B:ILE:HD11	15	1.66	0.97	1.84
(1,196)	2:15:B:THR:HG21	2:17:B:ILE:HD12	15	1.66	0.97	1.84
(1,196)	2:15:B:THR:HG21	2:17:B:ILE:HD13	15	1.66	0.97	1.84
(1,196)	2:15:B:THR:HG22	2:17:B:ILE:HD11	15	1.66	0.97	1.84
(1,196)	2:15:B:THR:HG22	2:17:B:ILE:HD12	15	1.66	0.97	1.84
(1,196)	2:15:B:THR:HG22	2:17:B:ILE:HD13	15	1.66	0.97	1.84
(1,196)	2:15:B:THR:HG23	2:17:B:ILE:HD11	15	1.66	0.97	1.84
(1,196)	2:15:B:THR:HG23	2:17:B:ILE:HD12	15	1.66	0.97	1.84
(1,196)	2:15:B:THR:HG23	2:17:B:ILE:HD13	15	1.66	0.97	1.84
(1,515)	2:15:B:THR:HG21	2:17:B:ILE:HD11	15	1.66	0.97	1.84
(1,515)	2:15:B:THR:HG21	2:17:B:ILE:HD12	15	1.66	0.97	1.84
(1,515)	2:15:B:THR:HG21	2:17:B:ILE:HD13	15	1.66	0.97	1.84
(1,515)	2:15:B:THR:HG22	2:17:B:ILE:HD11	15	1.66	0.97	1.84
(1,515)	2:15:B:THR:HG22	2:17:B:ILE:HD12	15	1.66	0.97	1.84
(1,515)	2:15:B:THR:HG22	2:17:B:ILE:HD13	15	1.66	0.97	1.84
(1,515)	2:15:B:THR:HG23	2:17:B:ILE:HD11	15	1.66	0.97	1.84
(1,515)	2:15:B:THR:HG23	2:17:B:ILE:HD12	15	1.66	0.97	1.84
(1,515)	2:15:B:THR:HG23	2:17:B:ILE:HD13	15	1.66	0.97	1.84
(1,757)	1:25:C:GLY:HA3	1:26:C:HIS:HD2	15	1.51	0.37	1.52
(1,1076)	1:25:C:GLY:HA3	1:26:C:HIS:HD2	15	1.51	0.37	1.52
(1,15)	2:19:B:PHE:HE1	2:38:D:ILE:HG13	15	1.5	0.78	1.44
(1,15)	2:19:B:PHE:HE2	2:38:D:ILE:HG13	15	1.5	0.78	1.44

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Key	Atom-1	Atom-2	Models[1]	Mean (Å)	SD[1] (Å)	Median (Å)
(1,662)	2:19:B:PHE:HE1	2:38:D:ILE:HG13	15	1.5	0.78	1.44
(1,662)	2:19:B:PHE:HE2	2:38:D:ILE:HG13	15	1.5	0.78	1.44
(1,101)	1:4:A:GLY:HA3	1:5:A:HIS:HD2	15	1.5	0.53	1.53
(1,420)	1:4:A:GLY:HA3	1:5:A:HIS:HD2	15	1.5	0.53	1.53
(1,803)	1:25:C:GLY:HA2	1:26:C:HIS:HD2	15	1.38	0.43	1.38
(1,1122)	1:25:C:GLY:HA2	1:26:C:HIS:HD2	15	1.38	0.43	1.38
(1,71)	1:6:A:VAL:HB	2:20:B:TYR:HE1	15	1.12	0.42	1.14
(1,71)	1:6:A:VAL:HB	2:20:B:TYR:HE2	15	1.12	0.42	1.14
(1,390)	1:6:A:VAL:HB	2:20:B:TYR:HE1	15	1.12	0.42	1.14
(1,390)	1:6:A:VAL:HB	2:20:B:TYR:HE2	15	1.12	0.42	1.14
(1,126)	1:4:A:GLY:HA2	1:6:A:VAL:H	15	0.52	0.32	0.49
(1,445)	1:4:A:GLY:HA2	1:6:A:VAL:H	15	0.52	0.32	0.49
(1,10)	2:19:B:PHE:HD1	2:38:D:ILE:HD11	14	2.11	0.82	2.19
(1,10)	2:19:B:PHE:HD1	2:38:D:ILE:HD12	14	2.11	0.82	2.19
(1,10)	2:19:B:PHE:HD1	2:38:D:ILE:HD13	14	2.11	0.82	2.19
(1,10)	2:19:B:PHE:HD2	2:38:D:ILE:HD11	14	2.11	0.82	2.19
(1,10)	2:19:B:PHE:HD2	2:38:D:ILE:HD12	14	2.11	0.82	2.19
(1,10)	2:19:B:PHE:HD2	2:38:D:ILE:HD13	14	2.11	0.82	2.19
(1,657)	2:19:B:PHE:HD1	2:38:D:ILE:HD11	14	2.11	0.82	2.19
(1,657)	2:19:B:PHE:HD1	2:38:D:ILE:HD12	14	2.11	0.82	2.19
(1,657)	2:19:B:PHE:HD1	2:38:D:ILE:HD13	14	2.11	0.82	2.19
(1,657)	2:19:B:PHE:HD2	2:38:D:ILE:HD11	14	2.11	0.82	2.19
(1,657)	2:19:B:PHE:HD2	2:38:D:ILE:HD12	14	2.11	0.82	2.19
(1,657)	2:19:B:PHE:HD2	2:38:D:ILE:HD13	14	2.11	0.82	2.19
(1,83)	1:10:A:PHE:HD1	2:14:B:GLY:HA3	14	1.6	0.81	1.48
(1,83)	1:10:A:PHE:HD2	2:14:B:GLY:HA3	14	1.6	0.81	1.48
(1,402)	1:10:A:PHE:HD1	2:14:B:GLY:HA3	14	1.6	0.81	1.48
(1,402)	1:10:A:PHE:HD2	2:14:B:GLY:HA3	14	1.6	0.81	1.48
(1,850)	2:36:D:THR:HB	2:38:D:ILE:HD11	14	1.32	0.97	0.9
(1,850)	2:36:D:THR:HB	2:38:D:ILE:HD12	14	1.32	0.97	0.9
(1,850)	2:36:D:THR:HB	2:38:D:ILE:HD13	14	1.32	0.97	0.9
(1,1169)	2:36:D:THR:HB	2:38:D:ILE:HD11	14	1.32	0.97	0.9
(1,1169)	2:36:D:THR:HB	2:38:D:ILE:HD12	14	1.32	0.97	0.9
(1,1169)	2:36:D:THR:HB	2:38:D:ILE:HD13	14	1.32	0.97	0.9
(1,1)	2:17:B:ILE:HD11	2:40:D:PHE:HD1	14	1.28	0.65	1.16
(1,1)	2:17:B:ILE:HD11	2:40:D:PHE:HD2	14	1.28	0.65	1.16
(1,1)	2:17:B:ILE:HD12	2:40:D:PHE:HD1	14	1.28	0.65	1.16
(1,1)	2:17:B:ILE:HD12	2:40:D:PHE:HD2	14	1.28	0.65	1.16
(1,1)	2:17:B:ILE:HD13	2:40:D:PHE:HD1	14	1.28	0.65	1.16
(1,1)	2:17:B:ILE:HD13	2:40:D:PHE:HD2	14	1.28	0.65	1.16
(1,666)	2:17:B:ILE:HD11	2:40:D:PHE:HD1	14	1.28	0.65	1.16
(1,666)	2:17:B:ILE:HD11	2:40:D:PHE:HD2	14	1.28	0.65	1.16

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Key	Atom-1	Atom-2	Models[1]	Mean (Å)	SD[1] (Å)	Median (Å)
(1,666)	2:17:B:ILE:HD12	2:40:D:PHE:HD1	14	1.28	0.65	1.16
(1,666)	2:17:B:ILE:HD12	2:40:D:PHE:HD2	14	1.28	0.65	1.16
(1,666)	2:17:B:ILE:HD13	2:40:D:PHE:HD1	14	1.28	0.65	1.16
(1,666)	2:17:B:ILE:HD13	2:40:D:PHE:HD2	14	1.28	0.65	1.16
(1,182)	1:4:A:GLY:H	1:6:A:VAL:H	14	0.68	0.44	0.52
(1,501)	1:4:A:GLY:H	1:6:A:VAL:H	14	0.68	0.44	0.52
(1,681)	1:31:C:PHE:HE1	2:37:D:PRO:HD2	14	0.55	0.57	0.36
(1,681)	1:31:C:PHE:HE2	2:37:D:PRO:HD2	14	0.55	0.57	0.36
(1,1000)	1:31:C:PHE:HE1	2:37:D:PRO:HD2	14	0.55	0.57	0.36
(1,1000)	1:31:C:PHE:HE2	2:37:D:PRO:HD2	14	0.55	0.57	0.36
(1,734)	2:39:D:SER:HB2	2:40:D:PHE:H	14	0.53	0.23	0.46
(1,1053)	2:39:D:SER:HB2	2:40:D:PHE:H	14	0.53	0.23	0.46
(1,130)	1:3:A:ALA:HB1	1:4:A:GLY:H	14	0.51	0.18	0.47
(1,130)	1:3:A:ALA:HB2	1:4:A:GLY:H	14	0.51	0.18	0.47
(1,130)	1:3:A:ALA:HB3	1:4:A:GLY:H	14	0.51	0.18	0.47
(1,449)	1:3:A:ALA:HB1	1:4:A:GLY:H	14	0.51	0.18	0.47
(1,449)	1:3:A:ALA:HB2	1:4:A:GLY:H	14	0.51	0.18	0.47
(1,449)	1:3:A:ALA:HB3	1:4:A:GLY:H	14	0.51	0.18	0.47
(1,811)	1:27:C:VAL:H	2:40:D:PHE:HB2	14	0.36	0.19	0.29
(1,1130)	1:27:C:VAL:H	2:40:D:PHE:HB2	14	0.36	0.19	0.29
(1,698)	1:31:C:PHE:HE1	2:35:D:GLY:HA3	13	1.87	1.06	1.92
(1,698)	1:31:C:PHE:HE2	2:35:D:GLY:HA3	13	1.87	1.06	1.92
(1,1017)	1:31:C:PHE:HE1	2:35:D:GLY:HA3	13	1.87	1.06	1.92
(1,1017)	1:31:C:PHE:HE2	2:35:D:GLY:HA3	13	1.87	1.06	1.92
(1,14)	2:19:B:PHE:H	2:38:D:ILE:HD11	13	1.76	0.73	1.94
(1,14)	2:19:B:PHE:H	2:38:D:ILE:HD12	13	1.76	0.73	1.94
(1,14)	2:19:B:PHE:H	2:38:D:ILE:HD13	13	1.76	0.73	1.94
(1,661)	2:19:B:PHE:H	2:38:D:ILE:HD11	13	1.76	0.73	1.94
(1,661)	2:19:B:PHE:H	2:38:D:ILE:HD12	13	1.76	0.73	1.94
(1,661)	2:19:B:PHE:H	2:38:D:ILE:HD13	13	1.76	0.73	1.94
(1,16)	2:19:B:PHE:HD1	2:38:D:ILE:HG13	13	1.37	0.71	1.2
(1,16)	2:19:B:PHE:HD2	2:38:D:ILE:HG13	13	1.37	0.71	1.2
(1,663)	2:19:B:PHE:HD1	2:38:D:ILE:HG13	13	1.37	0.71	1.2
(1,663)	2:19:B:PHE:HD2	2:38:D:ILE:HG13	13	1.37	0.71	1.2
(1,838)	1:25:C:GLY:H	1:27:C:VAL:H	13	0.88	0.41	0.99
(1,1157)	1:25:C:GLY:H	1:27:C:VAL:H	13	0.88	0.41	0.99
(1,312)	2:17:B:ILE:HA	2:17:B:ILE:HD11	13	0.59	0.22	0.56
(1,312)	2:17:B:ILE:HA	2:17:B:ILE:HD12	13	0.59	0.22	0.56
(1,312)	2:17:B:ILE:HA	2:17:B:ILE:HD13	13	0.59	0.22	0.56
(1,631)	2:17:B:ILE:HA	2:17:B:ILE:HD11	13	0.59	0.22	0.56
(1,631)	2:17:B:ILE:HA	2:17:B:ILE:HD12	13	0.59	0.22	0.56
(1,631)	2:17:B:ILE:HA	2:17:B:ILE:HD13	13	0.59	0.22	0.56

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Key	Atom-1	Atom-2	Models[1]	Mean (Å)	SD[1] (Å)	Median (Å)
(1,786)	1:24:C:ALA:HB1	1:25:C:GLY:H	13	0.59	0.11	0.59
(1,786)	1:24:C:ALA:HB2	1:25:C:GLY:H	13	0.59	0.11	0.59
(1,786)	1:24:C:ALA:HB3	1:25:C:GLY:H	13	0.59	0.11	0.59
(1,1105)	1:24:C:ALA:HB1	1:25:C:GLY:H	13	0.59	0.11	0.59
(1,1105)	1:24:C:ALA:HB2	1:25:C:GLY:H	13	0.59	0.11	0.59
(1,1105)	1:24:C:ALA:HB3	1:25:C:GLY:H	13	0.59	0.11	0.59
(1,78)	2:18:B:SER:HB2	2:19:B:PHE:H	13	0.58	0.3	0.42
(1,397)	2:18:B:SER:HB2	2:19:B:PHE:H	13	0.58	0.3	0.42
(1,790)	1:31:C:PHE:HD1	2:36:D:THR:HG21	13	0.42	0.3	0.3
(1,790)	1:31:C:PHE:HD1	2:36:D:THR:HG22	13	0.42	0.3	0.3
(1,790)	1:31:C:PHE:HD1	2:36:D:THR:HG23	13	0.42	0.3	0.3
(1,790)	1:31:C:PHE:HD2	2:36:D:THR:HG21	13	0.42	0.3	0.3
(1,790)	1:31:C:PHE:HD2	2:36:D:THR:HG22	13	0.42	0.3	0.3
(1,790)	1:31:C:PHE:HD2	2:36:D:THR:HG23	13	0.42	0.3	0.3
(1,1109)	1:31:C:PHE:HD1	2:36:D:THR:HG21	13	0.42	0.3	0.3
(1,1109)	1:31:C:PHE:HD1	2:36:D:THR:HG22	13	0.42	0.3	0.3
(1,1109)	1:31:C:PHE:HD1	2:36:D:THR:HG23	13	0.42	0.3	0.3
(1,1109)	1:31:C:PHE:HD2	2:36:D:THR:HG21	13	0.42	0.3	0.3
(1,1109)	1:31:C:PHE:HD2	2:36:D:THR:HG22	13	0.42	0.3	0.3
(1,1109)	1:31:C:PHE:HD2	2:36:D:THR:HG23	13	0.42	0.3	0.3
(1,180)	1:6:A:VAL:HG11	1:7:A:PRO:HD3	13	0.35	0.15	0.37
(1,180)	1:6:A:VAL:HG12	1:7:A:PRO:HD3	13	0.35	0.15	0.37
(1,180)	1:6:A:VAL:HG13	1:7:A:PRO:HD3	13	0.35	0.15	0.37
(1,499)	1:6:A:VAL:HG11	1:7:A:PRO:HD3	13	0.35	0.15	0.37
(1,499)	1:6:A:VAL:HG12	1:7:A:PRO:HD3	13	0.35	0.15	0.37
(1,499)	1:6:A:VAL:HG13	1:7:A:PRO:HD3	13	0.35	0.15	0.37
(1,754)	2:18:B:SER:HB3	2:38:D:ILE:HD11	12	1.56	0.93	1.64
(1,754)	2:18:B:SER:HB3	2:38:D:ILE:HD12	12	1.56	0.93	1.64
(1,754)	2:18:B:SER:HB3	2:38:D:ILE:HD13	12	1.56	0.93	1.64
(1,1073)	2:18:B:SER:HB3	2:38:D:ILE:HD11	12	1.56	0.93	1.64
(1,1073)	2:18:B:SER:HB3	2:38:D:ILE:HD12	12	1.56	0.93	1.64
(1,1073)	2:18:B:SER:HB3	2:38:D:ILE:HD13	12	1.56	0.93	1.64
(1,194)	2:15:B:THR:HB	2:17:B:ILE:HD11	12	1.37	0.49	1.23
(1,194)	2:15:B:THR:HB	2:17:B:ILE:HD12	12	1.37	0.49	1.23
(1,194)	2:15:B:THR:HB	2:17:B:ILE:HD13	12	1.37	0.49	1.23
(1,513)	2:15:B:THR:HB	2:17:B:ILE:HD11	12	1.37	0.49	1.23
(1,513)	2:15:B:THR:HB	2:17:B:ILE:HD12	12	1.37	0.49	1.23
(1,513)	2:15:B:THR:HB	2:17:B:ILE:HD13	12	1.37	0.49	1.23
(1,683)	1:30:C:TYR:HE1	2:39:D:SER:HB2	12	1.35	0.94	1.47
(1,683)	1:30:C:TYR:HE2	2:39:D:SER:HB2	12	1.35	0.94	1.47
(1,1002)	1:30:C:TYR:HE1	2:39:D:SER:HB2	12	1.35	0.94	1.47
(1,1002)	1:30:C:TYR:HE2	2:39:D:SER:HB2	12	1.35	0.94	1.47

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Key	Atom-1	Atom-2	Models[1]	Mean (Å)	SD[1] (Å)	Median (Å)
(1,80)	1:7:A:PRO:HG2	1:10:A:PHE:HE1	12	1.21	0.73	1.18
(1,80)	1:7:A:PRO:HG2	1:10:A:PHE:HE2	12	1.21	0.73	1.18
(1,80)	1:7:A:PRO:HG3	1:10:A:PHE:HE1	12	1.21	0.73	1.18
(1,80)	1:7:A:PRO:HG3	1:10:A:PHE:HE2	12	1.21	0.73	1.18
(1,399)	1:7:A:PRO:HG2	1:10:A:PHE:HE1	12	1.21	0.73	1.18
(1,399)	1:7:A:PRO:HG2	1:10:A:PHE:HE2	12	1.21	0.73	1.18
(1,399)	1:7:A:PRO:HG3	1:10:A:PHE:HE1	12	1.21	0.73	1.18
(1,399)	1:7:A:PRO:HG3	1:10:A:PHE:HE2	12	1.21	0.73	1.18
(1,277)	1:7:A:PRO:HG2	1:10:A:PHE:HE1	12	1.16	0.73	1.14
(1,277)	1:7:A:PRO:HG2	1:10:A:PHE:HE2	12	1.16	0.73	1.14
(1,277)	1:7:A:PRO:HG3	1:10:A:PHE:HE1	12	1.16	0.73	1.14
(1,277)	1:7:A:PRO:HG3	1:10:A:PHE:HE2	12	1.16	0.73	1.14
(1,596)	1:7:A:PRO:HG2	1:10:A:PHE:HE1	12	1.16	0.73	1.14
(1,596)	1:7:A:PRO:HG2	1:10:A:PHE:HE2	12	1.16	0.73	1.14
(1,596)	1:7:A:PRO:HG3	1:10:A:PHE:HE1	12	1.16	0.73	1.14
(1,596)	1:7:A:PRO:HG3	1:10:A:PHE:HE2	12	1.16	0.73	1.14
(1,696)	1:30:C:TYR:HD1	2:39:D:SER:HA	12	1.0	0.67	0.72
(1,696)	1:30:C:TYR:HD2	2:39:D:SER:HA	12	1.0	0.67	0.72
(1,1015)	1:30:C:TYR:HD1	2:39:D:SER:HA	12	1.0	0.67	0.72
(1,1015)	1:30:C:TYR:HD2	2:39:D:SER:HA	12	1.0	0.67	0.72
(1,685)	1:32:C:VAL:HA	1:33:C:ARG:HD2	12	0.84	0.4	0.82
(1,685)	1:32:C:VAL:HA	1:33:C:ARG:HD3	12	0.84	0.4	0.82
(1,1004)	1:32:C:VAL:HA	1:33:C:ARG:HD2	12	0.84	0.4	0.82
(1,1004)	1:32:C:VAL:HA	1:33:C:ARG:HD3	12	0.84	0.4	0.82
(1,25)	1:10:A:PHE:HE1	2:16:B:PRO:HD2	12	0.78	0.59	0.69
(1,25)	1:10:A:PHE:HE2	2:16:B:PRO:HD2	12	0.78	0.59	0.69
(1,344)	1:10:A:PHE:HE1	2:16:B:PRO:HD2	12	0.78	0.59	0.69
(1,344)	1:10:A:PHE:HE2	2:16:B:PRO:HD2	12	0.78	0.59	0.69
(1,134)	1:10:A:PHE:HD1	2:15:B:THR:HG21	12	0.7	0.37	0.62
(1,134)	1:10:A:PHE:HD1	2:15:B:THR:HG22	12	0.7	0.37	0.62
(1,134)	1:10:A:PHE:HD1	2:15:B:THR:HG23	12	0.7	0.37	0.62
(1,134)	1:10:A:PHE:HD2	2:15:B:THR:HG21	12	0.7	0.37	0.62
(1,134)	1:10:A:PHE:HD2	2:15:B:THR:HG22	12	0.7	0.37	0.62
(1,134)	1:10:A:PHE:HD2	2:15:B:THR:HG23	12	0.7	0.37	0.62
(1,453)	1:10:A:PHE:HD1	2:15:B:THR:HG21	12	0.7	0.37	0.62
(1,453)	1:10:A:PHE:HD1	2:15:B:THR:HG22	12	0.7	0.37	0.62
(1,453)	1:10:A:PHE:HD1	2:15:B:THR:HG23	12	0.7	0.37	0.62
(1,453)	1:10:A:PHE:HD2	2:15:B:THR:HG21	12	0.7	0.37	0.62
(1,453)	1:10:A:PHE:HD2	2:15:B:THR:HG22	12	0.7	0.37	0.62
(1,453)	1:10:A:PHE:HD2	2:15:B:THR:HG23	12	0.7	0.37	0.62
(1,726)	1:27:C:VAL:HB	2:41:D:TYR:HD1	12	0.68	0.37	0.58
(1,726)	1:27:C:VAL:HB	2:41:D:TYR:HD2	12	0.68	0.37	0.58

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Key	Atom-1	Atom-2	Models[1]	Mean (Å)	SD[1] (Å)	Median (Å)
(1,1045)	1:27:C:VAL:HB	2:41:D:TYR:HD1	12	0.68	0.37	0.58
(1,1045)	1:27:C:VAL:HB	2:41:D:TYR:HD2	12	0.68	0.37	0.58
(1,721)	1:31:C:PHE:HE1	2:37:D:PRO:HD2	12	0.5	0.59	0.31
(1,721)	1:31:C:PHE:HE2	2:37:D:PRO:HD2	12	0.5	0.59	0.31
(1,1040)	1:31:C:PHE:HE1	2:37:D:PRO:HD2	12	0.5	0.59	0.31
(1,1040)	1:31:C:PHE:HE2	2:37:D:PRO:HD2	12	0.5	0.59	0.31
(1,770)	1:29:C:GLU:HB2	2:41:D:TYR:HB2	12	0.48	0.45	0.34
(1,770)	1:29:C:GLU:HB2	2:41:D:TYR:HB3	12	0.48	0.45	0.34
(1,1089)	1:29:C:GLU:HB2	2:41:D:TYR:HB2	12	0.48	0.45	0.34
(1,1089)	1:29:C:GLU:HB2	2:41:D:TYR:HB3	12	0.48	0.45	0.34
(1,836)	1:27:C:VAL:HG11	1:28:C:PRO:HD3	12	0.44	0.21	0.36
(1,836)	1:27:C:VAL:HG12	1:28:C:PRO:HD3	12	0.44	0.21	0.36
(1,836)	1:27:C:VAL:HG13	1:28:C:PRO:HD3	12	0.44	0.21	0.36
(1,1155)	1:27:C:VAL:HG11	1:28:C:PRO:HD3	12	0.44	0.21	0.36
(1,1155)	1:27:C:VAL:HG12	1:28:C:PRO:HD3	12	0.44	0.21	0.36
(1,1155)	1:27:C:VAL:HG13	1:28:C:PRO:HD3	12	0.44	0.21	0.36
(1,686)	2:34:D:CYS:HB3	2:35:D:GLY:HA3	12	0.36	0.14	0.36
(1,1005)	2:34:D:CYS:HB3	2:35:D:GLY:HA3	12	0.36	0.14	0.36
(1,689)	2:18:B:SER:HA	2:38:D:ILE:HG13	11	0.91	0.65	0.76
(1,1008)	2:18:B:SER:HA	2:38:D:ILE:HG13	11	0.91	0.65	0.76
(1,92)	1:5:A:HIS:HB3	1:6:A:VAL:HG21	11	0.53	0.25	0.56
(1,92)	1:5:A:HIS:HB3	1:6:A:VAL:HG22	11	0.53	0.25	0.56
(1,92)	1:5:A:HIS:HB3	1:6:A:VAL:HG23	11	0.53	0.25	0.56
(1,411)	1:5:A:HIS:HB3	1:6:A:VAL:HG21	11	0.53	0.25	0.56
(1,411)	1:5:A:HIS:HB3	1:6:A:VAL:HG22	11	0.53	0.25	0.56
(1,411)	1:5:A:HIS:HB3	1:6:A:VAL:HG23	11	0.53	0.25	0.56
(1,79)	1:7:A:PRO:HA	2:19:B:PHE:HD1	11	0.52	0.28	0.51
(1,79)	1:7:A:PRO:HA	2:19:B:PHE:HD2	11	0.52	0.28	0.51
(1,398)	1:7:A:PRO:HA	2:19:B:PHE:HD1	11	0.52	0.28	0.51
(1,398)	1:7:A:PRO:HA	2:19:B:PHE:HD2	11	0.52	0.28	0.51
(1,86)	1:1:A:GLY:H	2:18:B:SER:HB2	11	0.5	0.32	0.47
(1,405)	1:1:A:GLY:H	2:18:B:SER:HB2	11	0.5	0.32	0.47
(1,968)	2:38:D:ILE:HA	2:38:D:ILE:HD11	11	0.48	0.15	0.49
(1,968)	2:38:D:ILE:HA	2:38:D:ILE:HD12	11	0.48	0.15	0.49
(1,968)	2:38:D:ILE:HA	2:38:D:ILE:HD13	11	0.48	0.15	0.49
(1,1287)	2:38:D:ILE:HA	2:38:D:ILE:HD11	11	0.48	0.15	0.49
(1,1287)	2:38:D:ILE:HA	2:38:D:ILE:HD12	11	0.48	0.15	0.49
(1,1287)	2:38:D:ILE:HA	2:38:D:ILE:HD13	11	0.48	0.15	0.49
(1,84)	1:7:A:PRO:HG2	2:19:B:PHE:HD1	11	0.43	0.19	0.41
(1,84)	1:7:A:PRO:HG2	2:19:B:PHE:HD2	11	0.43	0.19	0.41
(1,84)	1:7:A:PRO:HG3	2:19:B:PHE:HD1	11	0.43	0.19	0.41
(1,84)	1:7:A:PRO:HG3	2:19:B:PHE:HD2	11	0.43	0.19	0.41

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Key	Atom-1	Atom-2	Models[1]	Mean (Å)	SD[1] (Å)	Median (Å)
(1,403)	1:7:A:PRO:HG2	2:19:B:PHE:HD1	11	0.43	0.19	0.41
(1,403)	1:7:A:PRO:HG2	2:19:B:PHE:HD2	11	0.43	0.19	0.41
(1,403)	1:7:A:PRO:HG3	2:19:B:PHE:HD1	11	0.43	0.19	0.41
(1,403)	1:7:A:PRO:HG3	2:19:B:PHE:HD2	11	0.43	0.19	0.41
(1,155)	1:6:A:VAL:H	2:19:B:PHE:HB2	11	0.4	0.25	0.36
(1,474)	1:6:A:VAL:H	2:19:B:PHE:HB2	11	0.4	0.25	0.36
(1,42)	1:10:A:PHE:HE1	2:14:B:GLY:HA3	10	1.81	1.06	1.76
(1,42)	1:10:A:PHE:HE2	2:14:B:GLY:HA3	10	1.81	1.06	1.76
(1,361)	1:10:A:PHE:HE1	2:14:B:GLY:HA3	10	1.81	1.06	1.76
(1,361)	1:10:A:PHE:HE2	2:14:B:GLY:HA3	10	1.81	1.06	1.76
(1,6)	2:17:B:ILE:HG13	2:40:D:PHE:HE1	10	1.28	0.64	1.25
(1,6)	2:17:B:ILE:HG13	2:40:D:PHE:HE2	10	1.28	0.64	1.25
(1,671)	2:17:B:ILE:HG13	2:40:D:PHE:HE1	10	1.28	0.64	1.25
(1,671)	2:17:B:ILE:HG13	2:40:D:PHE:HE2	10	1.28	0.64	1.25
(1,736)	1:28:C:PRO:HG2	1:31:C:PHE:HE1	10	0.83	0.67	0.56
(1,736)	1:28:C:PRO:HG2	1:31:C:PHE:HE2	10	0.83	0.67	0.56
(1,736)	1:28:C:PRO:HG3	1:31:C:PHE:HE1	10	0.83	0.67	0.56
(1,736)	1:28:C:PRO:HG3	1:31:C:PHE:HE2	10	0.83	0.67	0.56
(1,1055)	1:28:C:PRO:HG2	1:31:C:PHE:HE1	10	0.83	0.67	0.56
(1,1055)	1:28:C:PRO:HG2	1:31:C:PHE:HE2	10	0.83	0.67	0.56
(1,1055)	1:28:C:PRO:HG3	1:31:C:PHE:HE1	10	0.83	0.67	0.56
(1,1055)	1:28:C:PRO:HG3	1:31:C:PHE:HE2	10	0.83	0.67	0.56
(1,933)	1:28:C:PRO:HG2	1:31:C:PHE:HE1	10	0.78	0.67	0.52
(1,933)	1:28:C:PRO:HG2	1:31:C:PHE:HE2	10	0.78	0.67	0.52
(1,933)	1:28:C:PRO:HG3	1:31:C:PHE:HE1	10	0.78	0.67	0.52
(1,933)	1:28:C:PRO:HG3	1:31:C:PHE:HE2	10	0.78	0.67	0.52
(1,1252)	1:28:C:PRO:HG2	1:31:C:PHE:HE1	10	0.78	0.67	0.52
(1,1252)	1:28:C:PRO:HG2	1:31:C:PHE:HE2	10	0.78	0.67	0.52
(1,1252)	1:28:C:PRO:HG3	1:31:C:PHE:HE1	10	0.78	0.67	0.52
(1,1252)	1:28:C:PRO:HG3	1:31:C:PHE:HE2	10	0.78	0.67	0.52
(1,124)	1:8:A:GLU:HG3	2:20:B:TYR:HD1	10	0.76	0.41	0.66
(1,124)	1:8:A:GLU:HG3	2:20:B:TYR:HD2	10	0.76	0.41	0.66
(1,443)	1:8:A:GLU:HG3	2:20:B:TYR:HD1	10	0.76	0.41	0.66
(1,443)	1:8:A:GLU:HG3	2:20:B:TYR:HD2	10	0.76	0.41	0.66
(1,133)	2:15:B:THR:HG21	2:16:B:PRO:HD3	10	0.65	0.29	0.6
(1,133)	2:15:B:THR:HG22	2:16:B:PRO:HD3	10	0.65	0.29	0.6
(1,133)	2:15:B:THR:HG23	2:16:B:PRO:HD3	10	0.65	0.29	0.6
(1,452)	2:15:B:THR:HG21	2:16:B:PRO:HD3	10	0.65	0.29	0.6
(1,452)	2:15:B:THR:HG22	2:16:B:PRO:HD3	10	0.65	0.29	0.6
(1,452)	2:15:B:THR:HG23	2:16:B:PRO:HD3	10	0.65	0.29	0.6
(1,748)	1:26:C:HIS:HB3	1:27:C:VAL:HG21	10	0.63	0.4	0.55
(1,748)	1:26:C:HIS:HB3	1:27:C:VAL:HG22	10	0.63	0.4	0.55

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Key	Atom-1	Atom-2	Models[1]	Mean (Å)	SD[1] (Å)	Median (Å)
(1,748)	1:26:C:HIS:HB3	1:27:C:VAL:HG23	10	0.63	0.4	0.55
(1,1067)	1:26:C:HIS:HB3	1:27:C:VAL:HG21	10	0.63	0.4	0.55
(1,1067)	1:26:C:HIS:HB3	1:27:C:VAL:HG22	10	0.63	0.4	0.55
(1,1067)	1:26:C:HIS:HB3	1:27:C:VAL:HG23	10	0.63	0.4	0.55
(1,723)	1:25:C:GLY:H	2:40:D:PHE:HD1	10	0.59	0.33	0.58
(1,723)	1:25:C:GLY:H	2:40:D:PHE:HD2	10	0.59	0.33	0.58
(1,1042)	1:25:C:GLY:H	2:40:D:PHE:HD1	10	0.59	0.33	0.58
(1,1042)	1:25:C:GLY:H	2:40:D:PHE:HD2	10	0.59	0.33	0.58
(1,132)	2:15:B:THR:HG21	2:16:B:PRO:HB3	10	0.53	0.35	0.4
(1,132)	2:15:B:THR:HG22	2:16:B:PRO:HB3	10	0.53	0.35	0.4
(1,132)	2:15:B:THR:HG23	2:16:B:PRO:HB3	10	0.53	0.35	0.4
(1,451)	2:15:B:THR:HG21	2:16:B:PRO:HB3	10	0.53	0.35	0.4
(1,451)	2:15:B:THR:HG22	2:16:B:PRO:HB3	10	0.53	0.35	0.4
(1,451)	2:15:B:THR:HG23	2:16:B:PRO:HB3	10	0.53	0.35	0.4
(1,123)	1:8:A:GLU:HG3	2:19:B:PHE:HA	10	0.42	0.23	0.38
(1,442)	1:8:A:GLU:HG3	2:19:B:PHE:HA	10	0.42	0.23	0.38
(1,684)	1:28:C:PRO:HD2	2:41:D:TYR:HD1	10	0.37	0.18	0.36
(1,684)	1:28:C:PRO:HD2	2:41:D:TYR:HD2	10	0.37	0.18	0.36
(1,1003)	1:28:C:PRO:HD2	2:41:D:TYR:HD1	10	0.37	0.18	0.36
(1,1003)	1:28:C:PRO:HD2	2:41:D:TYR:HD2	10	0.37	0.18	0.36
(1,195)	1:10:A:PHE:HB2	1:11:A:VAL:HG11	10	0.29	0.11	0.3
(1,195)	1:10:A:PHE:HB2	1:11:A:VAL:HG12	10	0.29	0.11	0.3
(1,195)	1:10:A:PHE:HB2	1:11:A:VAL:HG13	10	0.29	0.11	0.3
(1,195)	1:10:A:PHE:HB2	1:11:A:VAL:HG21	10	0.29	0.11	0.3
(1,195)	1:10:A:PHE:HB2	1:11:A:VAL:HG22	10	0.29	0.11	0.3
(1,195)	1:10:A:PHE:HB2	1:11:A:VAL:HG23	10	0.29	0.11	0.3
(1,514)	1:10:A:PHE:HB2	1:11:A:VAL:HG11	10	0.29	0.11	0.3
(1,514)	1:10:A:PHE:HB2	1:11:A:VAL:HG12	10	0.29	0.11	0.3
(1,514)	1:10:A:PHE:HB2	1:11:A:VAL:HG13	10	0.29	0.11	0.3
(1,514)	1:10:A:PHE:HB2	1:11:A:VAL:HG21	10	0.29	0.11	0.3
(1,514)	1:10:A:PHE:HB2	1:11:A:VAL:HG22	10	0.29	0.11	0.3
(1,514)	1:10:A:PHE:HB2	1:11:A:VAL:HG23	10	0.29	0.11	0.3
(1,867)	1:26:C:HIS:HA	1:26:C:HIS:HD2	10	0.2	0.08	0.2
(1,1186)	1:26:C:HIS:HA	1:26:C:HIS:HD2	10	0.2	0.08	0.2
(1,98)	2:17:B:ILE:HD11	2:39:D:SER:HB3	9	1.43	1.05	1.09
(1,98)	2:17:B:ILE:HD12	2:39:D:SER:HB3	9	1.43	1.05	1.09
(1,98)	2:17:B:ILE:HD13	2:39:D:SER:HB3	9	1.43	1.05	1.09
(1,417)	2:17:B:ILE:HD11	2:39:D:SER:HB3	9	1.43	1.05	1.09
(1,417)	2:17:B:ILE:HD12	2:39:D:SER:HB3	9	1.43	1.05	1.09
(1,417)	2:17:B:ILE:HD13	2:39:D:SER:HB3	9	1.43	1.05	1.09
(1,742)	1:22:C:GLY:H	2:39:D:SER:HB2	9	1.03	0.74	0.88
(1,1061)	1:22:C:GLY:H	2:39:D:SER:HB2	9	1.03	0.74	0.88

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Key	Atom-1	Atom-2	Models[1]	Mean (Å)	SD[1] (Å)	Median (Å)
(1,29)	1:11:A:VAL:HA	1:12:A:ARG:HD2	9	0.87	0.29	0.88
(1,29)	1:11:A:VAL:HA	1:12:A:ARG:HD3	9	0.87	0.29	0.88
(1,348)	1:11:A:VAL:HA	1:12:A:ARG:HD2	9	0.87	0.29	0.88
(1,348)	1:11:A:VAL:HA	1:12:A:ARG:HD3	9	0.87	0.29	0.88
(1,65)	1:10:A:PHE:HE1	2:16:B:PRO:HD2	9	0.87	0.53	0.84
(1,65)	1:10:A:PHE:HE2	2:16:B:PRO:HD2	9	0.87	0.53	0.84
(1,384)	1:10:A:PHE:HE1	2:16:B:PRO:HD2	9	0.87	0.53	0.84
(1,384)	1:10:A:PHE:HE2	2:16:B:PRO:HD2	9	0.87	0.53	0.84
(1,826)	1:25:C:GLY:H	2:40:D:PHE:HE1	9	0.71	0.45	0.59
(1,826)	1:25:C:GLY:H	2:40:D:PHE:HE2	9	0.71	0.45	0.59
(1,1145)	1:25:C:GLY:H	2:40:D:PHE:HE1	9	0.71	0.45	0.59
(1,1145)	1:25:C:GLY:H	2:40:D:PHE:HE2	9	0.71	0.45	0.59
(1,144)	1:5:A:HIS:HB2	1:6:A:VAL:HG11	9	0.55	0.64	0.36
(1,144)	1:5:A:HIS:HB2	1:6:A:VAL:HG12	9	0.55	0.64	0.36
(1,144)	1:5:A:HIS:HB2	1:6:A:VAL:HG13	9	0.55	0.64	0.36
(1,463)	1:5:A:HIS:HB2	1:6:A:VAL:HG11	9	0.55	0.64	0.36
(1,463)	1:5:A:HIS:HB2	1:6:A:VAL:HG12	9	0.55	0.64	0.36
(1,463)	1:5:A:HIS:HB2	1:6:A:VAL:HG13	9	0.55	0.64	0.36
(1,923)	1:33:C:ARG:H	1:33:C:ARG:HB2	9	0.19	0.06	0.16
(1,923)	1:33:C:ARG:H	1:33:C:ARG:HB3	9	0.19	0.06	0.16
(1,1242)	1:33:C:ARG:H	1:33:C:ARG:HB2	9	0.19	0.06	0.16
(1,1242)	1:33:C:ARG:H	1:33:C:ARG:HB3	9	0.19	0.06	0.16
(1,243)	1:9:A:TYR:HA	1:9:A:TYR:HE1	9	0.18	0.04	0.17
(1,243)	1:9:A:TYR:HA	1:9:A:TYR:HE2	9	0.18	0.04	0.17
(1,562)	1:9:A:TYR:HA	1:9:A:TYR:HE1	9	0.18	0.04	0.17
(1,562)	1:9:A:TYR:HA	1:9:A:TYR:HE2	9	0.18	0.04	0.17
(1,744)	1:22:C:GLY:H	2:39:D:SER:HA	8	0.92	0.51	0.73
(1,1063)	1:22:C:GLY:H	2:39:D:SER:HA	8	0.92	0.51	0.73
(1,32)	2:16:B:PRO:HA	2:17:B:ILE:HD11	8	0.86	0.55	0.68
(1,32)	2:16:B:PRO:HA	2:17:B:ILE:HD12	8	0.86	0.55	0.68
(1,32)	2:16:B:PRO:HA	2:17:B:ILE:HD13	8	0.86	0.55	0.68
(1,351)	2:16:B:PRO:HA	2:17:B:ILE:HD11	8	0.86	0.55	0.68
(1,351)	2:16:B:PRO:HA	2:17:B:ILE:HD12	8	0.86	0.55	0.68
(1,351)	2:16:B:PRO:HA	2:17:B:ILE:HD13	8	0.86	0.55	0.68
(1,27)	1:9:A:TYR:HE1	2:18:B:SER:HB2	8	0.85	0.63	0.5
(1,27)	1:9:A:TYR:HE2	2:18:B:SER:HB2	8	0.85	0.63	0.5
(1,346)	1:9:A:TYR:HE1	2:18:B:SER:HB2	8	0.85	0.63	0.5
(1,346)	1:9:A:TYR:HE2	2:18:B:SER:HB2	8	0.85	0.63	0.5
(1,706)	1:27:C:VAL:HG21	2:41:D:TYR:HE1	8	0.77	0.22	0.82
(1,706)	1:27:C:VAL:HG21	2:41:D:TYR:HE2	8	0.77	0.22	0.82
(1,706)	1:27:C:VAL:HG22	2:41:D:TYR:HE1	8	0.77	0.22	0.82
(1,706)	1:27:C:VAL:HG22	2:41:D:TYR:HE2	8	0.77	0.22	0.82

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Key	Atom-1	Atom-2	Models[1]	Mean (Å)	SD[1] (Å)	Median (Å)
(1,706)	1:27:C:VAL:HG23	2:41:D:TYR:HE1	8	0.77	0.22	0.82
(1,706)	1:27:C:VAL:HG23	2:41:D:TYR:HE2	8	0.77	0.22	0.82
(1,1025)	1:27:C:VAL:HG21	2:41:D:TYR:HE1	8	0.77	0.22	0.82
(1,1025)	1:27:C:VAL:HG21	2:41:D:TYR:HE2	8	0.77	0.22	0.82
(1,1025)	1:27:C:VAL:HG22	2:41:D:TYR:HE1	8	0.77	0.22	0.82
(1,1025)	1:27:C:VAL:HG22	2:41:D:TYR:HE2	8	0.77	0.22	0.82
(1,1025)	1:27:C:VAL:HG23	2:41:D:TYR:HE1	8	0.77	0.22	0.82
(1,1025)	1:27:C:VAL:HG23	2:41:D:TYR:HE2	8	0.77	0.22	0.82
(1,7)	2:17:B:ILE:HG13	2:40:D:PHE:HD1	8	0.75	0.36	0.8
(1,7)	2:17:B:ILE:HG13	2:40:D:PHE:HD2	8	0.75	0.36	0.8
(1,672)	2:17:B:ILE:HG13	2:40:D:PHE:HD1	8	0.75	0.36	0.8
(1,672)	2:17:B:ILE:HG13	2:40:D:PHE:HD2	8	0.75	0.36	0.8
(1,820)	1:22:C:GLY:H	2:41:D:TYR:HD1	8	0.52	0.34	0.36
(1,820)	1:22:C:GLY:H	2:41:D:TYR:HD2	8	0.52	0.34	0.36
(1,1139)	1:22:C:GLY:H	2:41:D:TYR:HD1	8	0.52	0.34	0.36
(1,1139)	1:22:C:GLY:H	2:41:D:TYR:HD2	8	0.52	0.34	0.36
(1,759)	1:25:C:GLY:H	2:41:D:TYR:H	8	0.4	0.27	0.32
(1,1078)	1:25:C:GLY:H	2:41:D:TYR:H	8	0.4	0.27	0.32
(1,70)	1:6:A:VAL:HB	2:20:B:TYR:HD1	8	0.36	0.21	0.32
(1,70)	1:6:A:VAL:HB	2:20:B:TYR:HD2	8	0.36	0.21	0.32
(1,389)	1:6:A:VAL:HB	2:20:B:TYR:HD1	8	0.36	0.21	0.32
(1,389)	1:6:A:VAL:HB	2:20:B:TYR:HD2	8	0.36	0.21	0.32
(1,43)	1:7:A:PRO:HG2	2:19:B:PHE:HD1	8	0.31	0.16	0.26
(1,43)	1:7:A:PRO:HG2	2:19:B:PHE:HD2	8	0.31	0.16	0.26
(1,43)	1:7:A:PRO:HG3	2:19:B:PHE:HD1	8	0.31	0.16	0.26
(1,43)	1:7:A:PRO:HG3	2:19:B:PHE:HD2	8	0.31	0.16	0.26
(1,362)	1:7:A:PRO:HG2	2:19:B:PHE:HD1	8	0.31	0.16	0.26
(1,362)	1:7:A:PRO:HG2	2:19:B:PHE:HD2	8	0.31	0.16	0.26
(1,362)	1:7:A:PRO:HG3	2:19:B:PHE:HD1	8	0.31	0.16	0.26
(1,362)	1:7:A:PRO:HG3	2:19:B:PHE:HD2	8	0.31	0.16	0.26
(1,60)	1:1:A:GLY:H	2:20:B:TYR:HE1	8	0.26	0.13	0.2
(1,60)	1:1:A:GLY:H	2:20:B:TYR:HE2	8	0.26	0.13	0.2
(1,379)	1:1:A:GLY:H	2:20:B:TYR:HE1	8	0.26	0.13	0.2
(1,379)	1:1:A:GLY:H	2:20:B:TYR:HE2	8	0.26	0.13	0.2
(1,899)	1:30:C:TYR:HA	1:30:C:TYR:HE1	8	0.2	0.04	0.19
(1,899)	1:30:C:TYR:HA	1:30:C:TYR:HE2	8	0.2	0.04	0.19
(1,1218)	1:30:C:TYR:HA	1:30:C:TYR:HE1	8	0.2	0.04	0.19
(1,1218)	1:30:C:TYR:HA	1:30:C:TYR:HE2	8	0.2	0.04	0.19
(1,5)	2:17:B:ILE:HD11	2:40:D:PHE:H	7	1.41	0.67	1.53
(1,5)	2:17:B:ILE:HD12	2:40:D:PHE:H	7	1.41	0.67	1.53
(1,5)	2:17:B:ILE:HD13	2:40:D:PHE:H	7	1.41	0.67	1.53
(1,670)	2:17:B:ILE:HD11	2:40:D:PHE:H	7	1.41	0.67	1.53

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Key	Atom-1	Atom-2	Models[1]	Mean (Å)	SD[1] (Å)	Median (Å)
(1,670)	2:17:B:ILE:HD12	2:40:D:PHE:H	7	1.41	0.67	1.53
(1,670)	2:17:B:ILE:HD13	2:40:D:PHE:H	7	1.41	0.67	1.53
(1,688)	2:37:D:PRO:HA	2:38:D:ILE:HD11	7	0.9	0.52	0.87
(1,688)	2:37:D:PRO:HA	2:38:D:ILE:HD12	7	0.9	0.52	0.87
(1,688)	2:37:D:PRO:HA	2:38:D:ILE:HD13	7	0.9	0.52	0.87
(1,1007)	2:37:D:PRO:HA	2:38:D:ILE:HD11	7	0.9	0.52	0.87
(1,1007)	2:37:D:PRO:HA	2:38:D:ILE:HD12	7	0.9	0.52	0.87
(1,1007)	2:37:D:PRO:HA	2:38:D:ILE:HD13	7	0.9	0.52	0.87
(1,274)	1:12:A:ARG:HB2	1:12:A:ARG:HE	7	0.86	0.39	1.0
(1,593)	1:12:A:ARG:HB2	1:12:A:ARG:HE	7	0.86	0.39	1.0
(1,74)	1:7:A:PRO:HG2	1:10:A:PHE:HD1	7	0.8	0.59	0.72
(1,74)	1:7:A:PRO:HG2	1:10:A:PHE:HD2	7	0.8	0.59	0.72
(1,74)	1:7:A:PRO:HG3	1:10:A:PHE:HD1	7	0.8	0.59	0.72
(1,74)	1:7:A:PRO:HG3	1:10:A:PHE:HD2	7	0.8	0.59	0.72
(1,393)	1:7:A:PRO:HG2	1:10:A:PHE:HD1	7	0.8	0.59	0.72
(1,393)	1:7:A:PRO:HG2	1:10:A:PHE:HD2	7	0.8	0.59	0.72
(1,393)	1:7:A:PRO:HG3	1:10:A:PHE:HD1	7	0.8	0.59	0.72
(1,393)	1:7:A:PRO:HG3	1:10:A:PHE:HD2	7	0.8	0.59	0.72
(1,20)	1:1:A:GLY:HA3	2:18:B:SER:HB2	7	0.75	0.52	0.45
(1,23)	1:1:A:GLY:HA3	2:18:B:SER:HB2	7	0.75	0.52	0.45
(1,339)	1:1:A:GLY:HA3	2:18:B:SER:HB2	7	0.75	0.52	0.45
(1,342)	1:1:A:GLY:HA3	2:18:B:SER:HB2	7	0.75	0.52	0.45
(1,88)	1:1:A:GLY:H	2:18:B:SER:HA	7	0.74	0.4	0.63
(1,407)	1:1:A:GLY:H	2:18:B:SER:HA	7	0.74	0.4	0.63
(1,67)	1:4:A:GLY:H	2:19:B:PHE:HD1	7	0.61	0.19	0.61
(1,67)	1:4:A:GLY:H	2:19:B:PHE:HD2	7	0.61	0.19	0.61
(1,386)	1:4:A:GLY:H	2:19:B:PHE:HD1	7	0.61	0.19	0.61
(1,386)	1:4:A:GLY:H	2:19:B:PHE:HD2	7	0.61	0.19	0.61
(1,162)	1:1:A:GLY:H	2:19:B:PHE:H	7	0.61	0.4	0.5
(1,481)	1:1:A:GLY:H	2:19:B:PHE:H	7	0.61	0.4	0.5
(1,310)	2:17:B:ILE:H	2:17:B:ILE:HD11	7	0.57	0.39	0.39
(1,310)	2:17:B:ILE:H	2:17:B:ILE:HD12	7	0.57	0.39	0.39
(1,310)	2:17:B:ILE:H	2:17:B:ILE:HD13	7	0.57	0.39	0.39
(1,629)	2:17:B:ILE:H	2:17:B:ILE:HD11	7	0.57	0.39	0.39
(1,629)	2:17:B:ILE:H	2:17:B:ILE:HD12	7	0.57	0.39	0.39
(1,629)	2:17:B:ILE:H	2:17:B:ILE:HD13	7	0.57	0.39	0.39
(1,930)	1:33:C:ARG:HB2	1:33:C:ARG:HE	7	0.56	0.43	0.34
(1,1249)	1:33:C:ARG:HB2	1:33:C:ARG:HE	7	0.56	0.43	0.34
(1,103)	1:4:A:GLY:H	2:20:B:TYR:H	7	0.53	0.37	0.44
(1,422)	1:4:A:GLY:H	2:20:B:TYR:H	7	0.53	0.37	0.44
(1,912)	1:31:C:PHE:HE1	2:36:D:THR:HA	7	0.49	0.52	0.3
(1,912)	1:31:C:PHE:HE2	2:36:D:THR:HA	7	0.49	0.52	0.3

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Key	Atom-1	Atom-2	Models[1]	Mean (Å)	SD[1] (Å)	Median (Å)
(1,1231)	1:31:C:PHE:HE1	2:36:D:THR:HA	7	0.49	0.52	0.3
(1,1231)	1:31:C:PHE:HE2	2:36:D:THR:HA	7	0.49	0.52	0.3
(1,924)	1:33:C:ARG:H	1:33:C:ARG:HE	7	0.38	0.14	0.35
(1,1243)	1:33:C:ARG:H	1:33:C:ARG:HE	7	0.38	0.14	0.35
(1,778)	1:29:C:GLU:HG3	2:41:D:TYR:HB2	7	0.36	0.2	0.22
(1,778)	1:29:C:GLU:HG3	2:41:D:TYR:HB3	7	0.36	0.2	0.22
(1,1097)	1:29:C:GLU:HG3	2:41:D:TYR:HB2	7	0.36	0.2	0.22
(1,1097)	1:29:C:GLU:HG3	2:41:D:TYR:HB3	7	0.36	0.2	0.22
(1,851)	1:31:C:PHE:HB2	1:32:C:VAL:HG11	7	0.29	0.06	0.3
(1,851)	1:31:C:PHE:HB2	1:32:C:VAL:HG12	7	0.29	0.06	0.3
(1,851)	1:31:C:PHE:HB2	1:32:C:VAL:HG13	7	0.29	0.06	0.3
(1,851)	1:31:C:PHE:HB2	1:32:C:VAL:HG21	7	0.29	0.06	0.3
(1,851)	1:31:C:PHE:HB2	1:32:C:VAL:HG22	7	0.29	0.06	0.3
(1,851)	1:31:C:PHE:HB2	1:32:C:VAL:HG23	7	0.29	0.06	0.3
(1,1170)	1:31:C:PHE:HB2	1:32:C:VAL:HG11	7	0.29	0.06	0.3
(1,1170)	1:31:C:PHE:HB2	1:32:C:VAL:HG12	7	0.29	0.06	0.3
(1,1170)	1:31:C:PHE:HB2	1:32:C:VAL:HG13	7	0.29	0.06	0.3
(1,1170)	1:31:C:PHE:HB2	1:32:C:VAL:HG21	7	0.29	0.06	0.3
(1,1170)	1:31:C:PHE:HB2	1:32:C:VAL:HG22	7	0.29	0.06	0.3
(1,1170)	1:31:C:PHE:HB2	1:32:C:VAL:HG23	7	0.29	0.06	0.3
(1,30)	2:13:B:CYS:HB3	2:14:B:GLY:HA3	7	0.28	0.09	0.29
(1,349)	2:13:B:CYS:HB3	2:14:B:GLY:HA3	7	0.28	0.09	0.29
(1,785)	1:25:C:GLY:H	2:40:D:PHE:H	7	0.26	0.12	0.22
(1,1104)	1:25:C:GLY:H	2:40:D:PHE:H	7	0.26	0.12	0.22
(1,802)	1:26:C:HIS:HD2	1:27:C:VAL:HG11	6	1.43	0.75	1.54
(1,802)	1:26:C:HIS:HD2	1:27:C:VAL:HG12	6	1.43	0.75	1.54
(1,802)	1:26:C:HIS:HD2	1:27:C:VAL:HG13	6	1.43	0.75	1.54
(1,1121)	1:26:C:HIS:HD2	1:27:C:VAL:HG11	6	1.43	0.75	1.54
(1,1121)	1:26:C:HIS:HD2	1:27:C:VAL:HG12	6	1.43	0.75	1.54
(1,1121)	1:26:C:HIS:HD2	1:27:C:VAL:HG13	6	1.43	0.75	1.54
(1,170)	1:4:A:GLY:H	2:19:B:PHE:HE1	6	0.83	0.21	0.81
(1,170)	1:4:A:GLY:H	2:19:B:PHE:HE2	6	0.83	0.21	0.81
(1,489)	1:4:A:GLY:H	2:19:B:PHE:HE1	6	0.83	0.21	0.81
(1,489)	1:4:A:GLY:H	2:19:B:PHE:HE2	6	0.83	0.21	0.81
(1,256)	1:10:A:PHE:HE1	2:15:B:THR:HA	6	0.78	0.51	0.57
(1,256)	1:10:A:PHE:HE2	2:15:B:THR:HA	6	0.78	0.51	0.57
(1,575)	1:10:A:PHE:HE1	2:15:B:THR:HA	6	0.78	0.51	0.57
(1,575)	1:10:A:PHE:HE2	2:15:B:THR:HA	6	0.78	0.51	0.57
(1,33)	2:17:B:ILE:HG13	2:39:D:SER:HA	6	0.71	0.38	0.67
(1,352)	2:17:B:ILE:HG13	2:39:D:SER:HA	6	0.71	0.38	0.67
(1,40)	1:9:A:TYR:HD1	2:18:B:SER:HA	6	0.62	0.3	0.61
(1,40)	1:9:A:TYR:HD2	2:18:B:SER:HA	6	0.62	0.3	0.61

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Key	Atom-1	Atom-2	Models[1]	Mean (Å)	SD[1] (Å)	Median (Å)
(1,359)	1:9:A:TYR:HD1	2:18:B:SER:HA	6	0.62	0.3	0.61
(1,359)	1:9:A:TYR:HD2	2:18:B:SER:HA	6	0.62	0.3	0.61
(1,966)	2:38:D:ILE:H	2:38:D:ILE:HD11	6	0.61	0.34	0.66
(1,966)	2:38:D:ILE:H	2:38:D:ILE:HD12	6	0.61	0.34	0.66
(1,966)	2:38:D:ILE:H	2:38:D:ILE:HD13	6	0.61	0.34	0.66
(1,1285)	2:38:D:ILE:H	2:38:D:ILE:HD11	6	0.61	0.34	0.66
(1,1285)	2:38:D:ILE:H	2:38:D:ILE:HD12	6	0.61	0.34	0.66
(1,1285)	2:38:D:ILE:H	2:38:D:ILE:HD13	6	0.61	0.34	0.66
(1,268)	1:12:A:ARG:H	1:12:A:ARG:HE	6	0.61	0.3	0.5
(1,587)	1:12:A:ARG:H	1:12:A:ARG:HE	6	0.61	0.3	0.5
(1,122)	1:8:A:GLU:HG3	2:20:B:TYR:HB2	6	0.6	0.26	0.5
(1,122)	1:8:A:GLU:HG3	2:20:B:TYR:HB3	6	0.6	0.26	0.5
(1,441)	1:8:A:GLU:HG3	2:20:B:TYR:HB2	6	0.6	0.26	0.5
(1,441)	1:8:A:GLU:HG3	2:20:B:TYR:HB3	6	0.6	0.26	0.5
(1,782)	1:25:C:GLY:HA2	1:27:C:VAL:H	6	0.51	0.41	0.3
(1,1101)	1:25:C:GLY:HA2	1:27:C:VAL:H	6	0.51	0.41	0.3
(1,755)	1:22:C:GLY:H	2:39:D:SER:HB3	6	0.45	0.35	0.37
(1,1074)	1:22:C:GLY:H	2:39:D:SER:HB3	6	0.45	0.35	0.37
(1,730)	1:28:C:PRO:HG2	1:31:C:PHE:HD1	6	0.43	0.34	0.25
(1,730)	1:28:C:PRO:HG2	1:31:C:PHE:HD2	6	0.43	0.34	0.25
(1,730)	1:28:C:PRO:HG3	1:31:C:PHE:HD1	6	0.43	0.34	0.25
(1,730)	1:28:C:PRO:HG3	1:31:C:PHE:HD2	6	0.43	0.34	0.25
(1,1049)	1:28:C:PRO:HG2	1:31:C:PHE:HD1	6	0.43	0.34	0.25
(1,1049)	1:28:C:PRO:HG2	1:31:C:PHE:HD2	6	0.43	0.34	0.25
(1,1049)	1:28:C:PRO:HG3	1:31:C:PHE:HD1	6	0.43	0.34	0.25
(1,1049)	1:28:C:PRO:HG3	1:31:C:PHE:HD2	6	0.43	0.34	0.25
(1,715)	2:38:D:ILE:H	2:38:D:ILE:HG13	6	0.42	0.13	0.43
(1,1034)	2:38:D:ILE:H	2:38:D:ILE:HG13	6	0.42	0.13	0.43
(1,87)	1:11:A:VAL:HG11	1:12:A:ARG:HA	6	0.29	0.06	0.31
(1,87)	1:11:A:VAL:HG12	1:12:A:ARG:HA	6	0.29	0.06	0.31
(1,87)	1:11:A:VAL:HG13	1:12:A:ARG:HA	6	0.29	0.06	0.31
(1,87)	1:11:A:VAL:HG21	1:12:A:ARG:HA	6	0.29	0.06	0.31
(1,87)	1:11:A:VAL:HG22	1:12:A:ARG:HA	6	0.29	0.06	0.31
(1,87)	1:11:A:VAL:HG23	1:12:A:ARG:HA	6	0.29	0.06	0.31
(1,406)	1:11:A:VAL:HG11	1:12:A:ARG:HA	6	0.29	0.06	0.31
(1,406)	1:11:A:VAL:HG12	1:12:A:ARG:HA	6	0.29	0.06	0.31
(1,406)	1:11:A:VAL:HG13	1:12:A:ARG:HA	6	0.29	0.06	0.31
(1,406)	1:11:A:VAL:HG21	1:12:A:ARG:HA	6	0.29	0.06	0.31
(1,406)	1:11:A:VAL:HG22	1:12:A:ARG:HA	6	0.29	0.06	0.31
(1,406)	1:11:A:VAL:HG23	1:12:A:ARG:HA	6	0.29	0.06	0.31
(1,174)	1:11:A:VAL:HB	1:12:A:ARG:H	6	0.22	0.09	0.24
(1,493)	1:11:A:VAL:HB	1:12:A:ARG:H	6	0.22	0.09	0.24

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Key	Atom-1	Atom-2	Models[1]	Mean (Å)	SD[1] (Å)	Median (Å)
(1,267)	1:12:A:ARG:H	1:12:A:ARG:HB2	6	0.21	0.05	0.22
(1,267)	1:12:A:ARG:H	1:12:A:ARG:HB3	6	0.21	0.05	0.22
(1,586)	1:12:A:ARG:H	1:12:A:ARG:HB2	6	0.21	0.05	0.22
(1,586)	1:12:A:ARG:H	1:12:A:ARG:HB3	6	0.21	0.05	0.22
(1,676)	1:22:C:GLY:HA3	2:39:D:SER:HB2	5	0.95	0.64	0.72
(1,679)	1:22:C:GLY:HA3	2:39:D:SER:HB2	5	0.95	0.64	0.72
(1,995)	1:22:C:GLY:HA3	2:39:D:SER:HB2	5	0.95	0.64	0.72
(1,998)	1:22:C:GLY:HA3	2:39:D:SER:HB2	5	0.95	0.64	0.72
(1,3)	1:3:A:ALA:HB1	1:32:C:VAL:HA	5	0.9	0.43	0.87
(1,3)	1:3:A:ALA:HB2	1:32:C:VAL:HA	5	0.9	0.43	0.87
(1,3)	1:3:A:ALA:HB3	1:32:C:VAL:HA	5	0.9	0.43	0.87
(1,668)	1:3:A:ALA:HB1	1:32:C:VAL:HA	5	0.9	0.43	0.87
(1,668)	1:3:A:ALA:HB2	1:32:C:VAL:HA	5	0.9	0.43	0.87
(1,668)	1:3:A:ALA:HB3	1:32:C:VAL:HA	5	0.9	0.43	0.87
(1,303)	2:16:B:PRO:HB3	2:16:B:PRO:HD3	5	0.89	0.03	0.89
(1,622)	2:16:B:PRO:HB3	2:16:B:PRO:HD3	5	0.89	0.03	0.89
(1,146)	1:5:A:HIS:HD2	1:6:A:VAL:HG11	5	0.81	0.63	0.52
(1,146)	1:5:A:HIS:HD2	1:6:A:VAL:HG12	5	0.81	0.63	0.52
(1,146)	1:5:A:HIS:HD2	1:6:A:VAL:HG13	5	0.81	0.63	0.52
(1,465)	1:5:A:HIS:HD2	1:6:A:VAL:HG11	5	0.81	0.63	0.52
(1,465)	1:5:A:HIS:HD2	1:6:A:VAL:HG12	5	0.81	0.63	0.52
(1,465)	1:5:A:HIS:HD2	1:6:A:VAL:HG13	5	0.81	0.63	0.52
(1,37)	1:8:A:GLU:HG2	2:20:B:TYR:HE1	5	0.77	0.38	0.69
(1,37)	1:8:A:GLU:HG2	2:20:B:TYR:HE2	5	0.77	0.38	0.69
(1,356)	1:8:A:GLU:HG2	2:20:B:TYR:HE1	5	0.77	0.38	0.69
(1,356)	1:8:A:GLU:HG2	2:20:B:TYR:HE2	5	0.77	0.38	0.69
(1,114)	1:8:A:GLU:HB2	2:20:B:TYR:HB2	5	0.7	0.63	0.5
(1,114)	1:8:A:GLU:HB2	2:20:B:TYR:HB3	5	0.7	0.63	0.5
(1,433)	1:8:A:GLU:HB2	2:20:B:TYR:HB2	5	0.7	0.63	0.5
(1,433)	1:8:A:GLU:HB2	2:20:B:TYR:HB3	5	0.7	0.63	0.5
(1,173)	1:7:A:PRO:HB2	1:10:A:PHE:HD1	5	0.7	0.43	0.81
(1,173)	1:7:A:PRO:HB2	1:10:A:PHE:HD2	5	0.7	0.43	0.81
(1,492)	1:7:A:PRO:HB2	1:10:A:PHE:HD1	5	0.7	0.43	0.81
(1,492)	1:7:A:PRO:HB2	1:10:A:PHE:HD2	5	0.7	0.43	0.81
(1,728)	1:24:C:ALA:HA	2:40:D:PHE:HD1	5	0.62	0.24	0.53
(1,728)	1:24:C:ALA:HA	2:40:D:PHE:HD2	5	0.62	0.24	0.53
(1,990)	1:24:C:ALA:HA	2:40:D:PHE:HD1	5	0.62	0.24	0.53
(1,990)	1:24:C:ALA:HA	2:40:D:PHE:HD2	5	0.62	0.24	0.53
(1,1047)	1:24:C:ALA:HA	2:40:D:PHE:HD1	5	0.62	0.24	0.53
(1,1047)	1:24:C:ALA:HA	2:40:D:PHE:HD2	5	0.62	0.24	0.53
(1,1309)	1:24:C:ALA:HA	2:40:D:PHE:HD1	5	0.62	0.24	0.53
(1,1309)	1:24:C:ALA:HA	2:40:D:PHE:HD2	5	0.62	0.24	0.53

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Key	Atom-1	Atom-2	Models[1]	Mean (Å)	SD[1] (Å)	Median (Å)
(1,780)	1:29:C:GLU:HG3	2:41:D:TYR:HD1	5	0.52	0.15	0.5
(1,780)	1:29:C:GLU:HG3	2:41:D:TYR:HD2	5	0.52	0.15	0.5
(1,1099)	1:29:C:GLU:HG3	2:41:D:TYR:HD1	5	0.52	0.15	0.5
(1,1099)	1:29:C:GLU:HG3	2:41:D:TYR:HD2	5	0.52	0.15	0.5
(1,708)	1:24:C:ALA:H	2:40:D:PHE:HD1	5	0.43	0.29	0.29
(1,708)	1:24:C:ALA:H	2:40:D:PHE:HD2	5	0.43	0.29	0.29
(1,847)	1:24:C:ALA:H	2:40:D:PHE:HD1	5	0.43	0.29	0.29
(1,847)	1:24:C:ALA:H	2:40:D:PHE:HD2	5	0.43	0.29	0.29
(1,1027)	1:24:C:ALA:H	2:40:D:PHE:HD1	5	0.43	0.29	0.29
(1,1027)	1:24:C:ALA:H	2:40:D:PHE:HD2	5	0.43	0.29	0.29
(1,1166)	1:24:C:ALA:H	2:40:D:PHE:HD1	5	0.43	0.29	0.29
(1,1166)	1:24:C:ALA:H	2:40:D:PHE:HD2	5	0.43	0.29	0.29
(1,707)	1:27:C:VAL:HG11	1:28:C:PRO:HD3	5	0.34	0.15	0.43
(1,707)	1:27:C:VAL:HG12	1:28:C:PRO:HD3	5	0.34	0.15	0.43
(1,707)	1:27:C:VAL:HG13	1:28:C:PRO:HD3	5	0.34	0.15	0.43
(1,1026)	1:27:C:VAL:HG11	1:28:C:PRO:HD3	5	0.34	0.15	0.43
(1,1026)	1:27:C:VAL:HG12	1:28:C:PRO:HD3	5	0.34	0.15	0.43
(1,1026)	1:27:C:VAL:HG13	1:28:C:PRO:HD3	5	0.34	0.15	0.43
(1,779)	1:29:C:GLU:HG3	2:40:D:PHE:HA	5	0.29	0.06	0.3
(1,1098)	1:29:C:GLU:HG3	2:40:D:PHE:HA	5	0.29	0.06	0.3
(1,55)	1:3:A:ALA:HB1	1:4:A:GLY:HA3	5	0.28	0.03	0.29
(1,55)	1:3:A:ALA:HB2	1:4:A:GLY:HA3	5	0.28	0.03	0.29
(1,55)	1:3:A:ALA:HB3	1:4:A:GLY:HA3	5	0.28	0.03	0.29
(1,115)	1:3:A:ALA:HB1	1:4:A:GLY:HA3	5	0.28	0.03	0.29
(1,115)	1:3:A:ALA:HB2	1:4:A:GLY:HA3	5	0.28	0.03	0.29
(1,115)	1:3:A:ALA:HB3	1:4:A:GLY:HA3	5	0.28	0.03	0.29
(1,374)	1:3:A:ALA:HB1	1:4:A:GLY:HA3	5	0.28	0.03	0.29
(1,374)	1:3:A:ALA:HB2	1:4:A:GLY:HA3	5	0.28	0.03	0.29
(1,374)	1:3:A:ALA:HB3	1:4:A:GLY:HA3	5	0.28	0.03	0.29
(1,434)	1:3:A:ALA:HB1	1:4:A:GLY:HA3	5	0.28	0.03	0.29
(1,434)	1:3:A:ALA:HB2	1:4:A:GLY:HA3	5	0.28	0.03	0.29
(1,434)	1:3:A:ALA:HB3	1:4:A:GLY:HA3	5	0.28	0.03	0.29
(1,157)	2:18:B:SER:HB3	2:19:B:PHE:H	5	0.27	0.04	0.25
(1,159)	2:18:B:SER:HB3	2:19:B:PHE:H	5	0.27	0.04	0.25
(1,476)	2:18:B:SER:HB3	2:19:B:PHE:H	5	0.27	0.04	0.25
(1,478)	2:18:B:SER:HB3	2:19:B:PHE:H	5	0.27	0.04	0.25
(1,984)	2:41:D:TYR:H	2:41:D:TYR:HD1	5	0.26	0.22	0.16
(1,984)	2:41:D:TYR:H	2:41:D:TYR:HD2	5	0.26	0.22	0.16
(1,1303)	2:41:D:TYR:H	2:41:D:TYR:HD1	5	0.26	0.22	0.16
(1,1303)	2:41:D:TYR:H	2:41:D:TYR:HD2	5	0.26	0.22	0.16
(1,249)	1:10:A:PHE:H	1:10:A:PHE:HD1	5	0.23	0.12	0.16
(1,249)	1:10:A:PHE:H	1:10:A:PHE:HD2	5	0.23	0.12	0.16

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Key	Atom-1	Atom-2	Models[1]	Mean (Å)	SD[1] (Å)	Median (Å)
(1,568)	1:10:A:PHE:H	1:10:A:PHE:HD1	5	0.23	0.12	0.16
(1,568)	1:10:A:PHE:H	1:10:A:PHE:HD2	5	0.23	0.12	0.16
(1,51)	1:6:A:VAL:HG11	1:7:A:PRO:HD3	5	0.2	0.07	0.18
(1,51)	1:6:A:VAL:HG12	1:7:A:PRO:HD3	5	0.2	0.07	0.18
(1,51)	1:6:A:VAL:HG13	1:7:A:PRO:HD3	5	0.2	0.07	0.18
(1,370)	1:6:A:VAL:HG11	1:7:A:PRO:HD3	5	0.2	0.07	0.18
(1,370)	1:6:A:VAL:HG12	1:7:A:PRO:HD3	5	0.2	0.07	0.18
(1,370)	1:6:A:VAL:HG13	1:7:A:PRO:HD3	5	0.2	0.07	0.18
(1,91)	1:2:A:GLY:HA3	1:3:A:ALA:HB1	5	0.18	0.04	0.17
(1,91)	1:2:A:GLY:HA3	1:3:A:ALA:HB2	5	0.18	0.04	0.17
(1,91)	1:2:A:GLY:HA3	1:3:A:ALA:HB3	5	0.18	0.04	0.17
(1,410)	1:2:A:GLY:HA3	1:3:A:ALA:HB1	5	0.18	0.04	0.17
(1,410)	1:2:A:GLY:HA3	1:3:A:ALA:HB2	5	0.18	0.04	0.17
(1,410)	1:2:A:GLY:HA3	1:3:A:ALA:HB3	5	0.18	0.04	0.17
(1,905)	1:31:C:PHE:H	1:31:C:PHE:HD1	5	0.16	0.04	0.16
(1,905)	1:31:C:PHE:H	1:31:C:PHE:HD2	5	0.16	0.04	0.16
(1,1224)	1:31:C:PHE:H	1:31:C:PHE:HD1	5	0.16	0.04	0.16
(1,1224)	1:31:C:PHE:H	1:31:C:PHE:HD2	5	0.16	0.04	0.16
(1,837)	2:34:D:CYS:HB3	2:35:D:GLY:H	5	0.16	0.05	0.14
(1,1156)	2:34:D:CYS:HB3	2:35:D:GLY:H	5	0.16	0.05	0.14
(1,902)	1:31:C:PHE:HA	1:32:C:VAL:H	5	0.16	0.04	0.17
(1,1221)	1:31:C:PHE:HA	1:32:C:VAL:H	5	0.16	0.04	0.17
(1,246)	1:10:A:PHE:HA	1:11:A:VAL:H	5	0.12	0.02	0.1
(1,565)	1:10:A:PHE:HA	1:11:A:VAL:H	5	0.12	0.02	0.1
(1,818)	1:22:C:GLY:H	2:40:D:PHE:H	4	0.99	0.59	0.91
(1,1137)	1:22:C:GLY:H	2:40:D:PHE:H	4	0.99	0.59	0.91
(1,806)	1:31:C:PHE:HD1	1:32:C:VAL:HG11	4	0.77	0.38	0.66
(1,806)	1:31:C:PHE:HD1	1:32:C:VAL:HG12	4	0.77	0.38	0.66
(1,806)	1:31:C:PHE:HD1	1:32:C:VAL:HG13	4	0.77	0.38	0.66
(1,806)	1:31:C:PHE:HD1	1:32:C:VAL:HG21	4	0.77	0.38	0.66
(1,806)	1:31:C:PHE:HD1	1:32:C:VAL:HG22	4	0.77	0.38	0.66
(1,806)	1:31:C:PHE:HD1	1:32:C:VAL:HG23	4	0.77	0.38	0.66
(1,806)	1:31:C:PHE:HD2	1:32:C:VAL:HG11	4	0.77	0.38	0.66
(1,806)	1:31:C:PHE:HD2	1:32:C:VAL:HG12	4	0.77	0.38	0.66
(1,806)	1:31:C:PHE:HD2	1:32:C:VAL:HG13	4	0.77	0.38	0.66
(1,806)	1:31:C:PHE:HD2	1:32:C:VAL:HG21	4	0.77	0.38	0.66
(1,806)	1:31:C:PHE:HD2	1:32:C:VAL:HG22	4	0.77	0.38	0.66
(1,806)	1:31:C:PHE:HD2	1:32:C:VAL:HG23	4	0.77	0.38	0.66
(1,1125)	1:31:C:PHE:HD1	1:32:C:VAL:HG11	4	0.77	0.38	0.66
(1,1125)	1:31:C:PHE:HD1	1:32:C:VAL:HG12	4	0.77	0.38	0.66
(1,1125)	1:31:C:PHE:HD1	1:32:C:VAL:HG13	4	0.77	0.38	0.66
(1,1125)	1:31:C:PHE:HD1	1:32:C:VAL:HG21	4	0.77	0.38	0.66

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Key	Atom-1	Atom-2	Models[1]	Mean (Å)	SD[1] (Å)	Median (Å)
(1,1125)	1:31:C:PHE:HD1	1:32:C:VAL:HG22	4	0.77	0.38	0.66
(1,1125)	1:31:C:PHE:HD1	1:32:C:VAL:HG23	4	0.77	0.38	0.66
(1,1125)	1:31:C:PHE:HD2	1:32:C:VAL:HG11	4	0.77	0.38	0.66
(1,1125)	1:31:C:PHE:HD2	1:32:C:VAL:HG12	4	0.77	0.38	0.66
(1,1125)	1:31:C:PHE:HD2	1:32:C:VAL:HG13	4	0.77	0.38	0.66
(1,1125)	1:31:C:PHE:HD2	1:32:C:VAL:HG21	4	0.77	0.38	0.66
(1,1125)	1:31:C:PHE:HD2	1:32:C:VAL:HG22	4	0.77	0.38	0.66
(1,1125)	1:31:C:PHE:HD2	1:32:C:VAL:HG23	4	0.77	0.38	0.66
(1,690)	1:27:C:VAL:HB	2:41:D:TYR:HB2	4	0.68	0.51	0.51
(1,690)	1:27:C:VAL:HB	2:41:D:TYR:HB3	4	0.68	0.51	0.51
(1,1009)	1:27:C:VAL:HB	2:41:D:TYR:HB2	4	0.68	0.51	0.51
(1,1009)	1:27:C:VAL:HB	2:41:D:TYR:HB3	4	0.68	0.51	0.51
(1,85)	1:10:A:PHE:HD1	2:16:B:PRO:HA	4	0.66	0.37	0.67
(1,85)	1:10:A:PHE:HD2	2:16:B:PRO:HA	4	0.66	0.37	0.67
(1,404)	1:10:A:PHE:HD1	2:16:B:PRO:HA	4	0.66	0.37	0.67
(1,404)	1:10:A:PHE:HD2	2:16:B:PRO:HA	4	0.66	0.37	0.67
(1,697)	1:28:C:PRO:HG2	1:31:C:PHE:HB2	4	0.54	0.38	0.38
(1,697)	1:28:C:PRO:HG3	1:31:C:PHE:HB2	4	0.54	0.38	0.38
(1,731)	1:28:C:PRO:HG2	1:31:C:PHE:HB2	4	0.54	0.38	0.38
(1,731)	1:28:C:PRO:HG3	1:31:C:PHE:HB2	4	0.54	0.38	0.38
(1,1016)	1:28:C:PRO:HG2	1:31:C:PHE:HB2	4	0.54	0.38	0.38
(1,1016)	1:28:C:PRO:HG3	1:31:C:PHE:HB2	4	0.54	0.38	0.38
(1,1050)	1:28:C:PRO:HG2	1:31:C:PHE:HB2	4	0.54	0.38	0.38
(1,1050)	1:28:C:PRO:HG3	1:31:C:PHE:HB2	4	0.54	0.38	0.38
(1,150)	1:10:A:PHE:HD1	1:11:A:VAL:HG11	4	0.5	0.28	0.51
(1,150)	1:10:A:PHE:HD1	1:11:A:VAL:HG12	4	0.5	0.28	0.51
(1,150)	1:10:A:PHE:HD1	1:11:A:VAL:HG13	4	0.5	0.28	0.51
(1,150)	1:10:A:PHE:HD1	1:11:A:VAL:HG21	4	0.5	0.28	0.51
(1,150)	1:10:A:PHE:HD1	1:11:A:VAL:HG22	4	0.5	0.28	0.51
(1,150)	1:10:A:PHE:HD1	1:11:A:VAL:HG23	4	0.5	0.28	0.51
(1,150)	1:10:A:PHE:HD2	1:11:A:VAL:HG11	4	0.5	0.28	0.51
(1,150)	1:10:A:PHE:HD2	1:11:A:VAL:HG12	4	0.5	0.28	0.51
(1,150)	1:10:A:PHE:HD2	1:11:A:VAL:HG13	4	0.5	0.28	0.51
(1,150)	1:10:A:PHE:HD2	1:11:A:VAL:HG21	4	0.5	0.28	0.51
(1,150)	1:10:A:PHE:HD2	1:11:A:VAL:HG22	4	0.5	0.28	0.51
(1,150)	1:10:A:PHE:HD2	1:11:A:VAL:HG23	4	0.5	0.28	0.51
(1,469)	1:10:A:PHE:HD1	1:11:A:VAL:HG11	4	0.5	0.28	0.51
(1,469)	1:10:A:PHE:HD1	1:11:A:VAL:HG12	4	0.5	0.28	0.51
(1,469)	1:10:A:PHE:HD1	1:11:A:VAL:HG13	4	0.5	0.28	0.51
(1,469)	1:10:A:PHE:HD1	1:11:A:VAL:HG21	4	0.5	0.28	0.51
(1,469)	1:10:A:PHE:HD1	1:11:A:VAL:HG22	4	0.5	0.28	0.51
(1,469)	1:10:A:PHE:HD1	1:11:A:VAL:HG23	4	0.5	0.28	0.51

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Key	Atom-1	Atom-2	Models[1]	Mean (Å)	SD[1] (Å)	Median (Å)
(1,469)	1:10:A:PHE:HD2	1:11:A:VAL:HG11	4	0.5	0.28	0.51
(1,469)	1:10:A:PHE:HD2	1:11:A:VAL:HG12	4	0.5	0.28	0.51
(1,469)	1:10:A:PHE:HD2	1:11:A:VAL:HG13	4	0.5	0.28	0.51
(1,469)	1:10:A:PHE:HD2	1:11:A:VAL:HG21	4	0.5	0.28	0.51
(1,469)	1:10:A:PHE:HD2	1:11:A:VAL:HG22	4	0.5	0.28	0.51
(1,469)	1:10:A:PHE:HD2	1:11:A:VAL:HG23	4	0.5	0.28	0.51
(1,781)	2:40:D:PHE:HA	2:41:D:TYR:HD1	4	0.48	0.36	0.32
(1,781)	2:40:D:PHE:HA	2:41:D:TYR:HD2	4	0.48	0.36	0.32
(1,1100)	2:40:D:PHE:HA	2:41:D:TYR:HD1	4	0.48	0.36	0.32
(1,1100)	2:40:D:PHE:HA	2:41:D:TYR:HD2	4	0.48	0.36	0.32
(1,63)	1:10:A:PHE:HD1	2:16:B:PRO:HD2	4	0.46	0.24	0.44
(1,63)	1:10:A:PHE:HD2	2:16:B:PRO:HD2	4	0.46	0.24	0.44
(1,382)	1:10:A:PHE:HD1	2:16:B:PRO:HD2	4	0.46	0.24	0.44
(1,382)	1:10:A:PHE:HD2	2:16:B:PRO:HD2	4	0.46	0.24	0.44
(1,105)	1:4:A:GLY:H	2:20:B:TYR:H	4	0.46	0.32	0.4
(1,424)	1:4:A:GLY:H	2:20:B:TYR:H	4	0.46	0.32	0.4
(1,813)	2:39:D:SER:HB3	2:40:D:PHE:H	4	0.37	0.1	0.36
(1,815)	2:39:D:SER:HB3	2:40:D:PHE:H	4	0.37	0.1	0.36
(1,1132)	2:39:D:SER:HB3	2:40:D:PHE:H	4	0.37	0.1	0.36
(1,1134)	2:39:D:SER:HB3	2:40:D:PHE:H	4	0.37	0.1	0.36
(1,59)	2:17:B:ILE:H	2:17:B:ILE:HG13	4	0.34	0.08	0.36
(1,378)	2:17:B:ILE:H	2:17:B:ILE:HG13	4	0.34	0.08	0.36
(1,743)	1:32:C:VAL:HG11	1:33:C:ARG:HA	4	0.3	0.07	0.32
(1,743)	1:32:C:VAL:HG12	1:33:C:ARG:HA	4	0.3	0.07	0.32
(1,743)	1:32:C:VAL:HG13	1:33:C:ARG:HA	4	0.3	0.07	0.32
(1,743)	1:32:C:VAL:HG21	1:33:C:ARG:HA	4	0.3	0.07	0.32
(1,743)	1:32:C:VAL:HG22	1:33:C:ARG:HA	4	0.3	0.07	0.32
(1,743)	1:32:C:VAL:HG23	1:33:C:ARG:HA	4	0.3	0.07	0.32
(1,1062)	1:32:C:VAL:HG11	1:33:C:ARG:HA	4	0.3	0.07	0.32
(1,1062)	1:32:C:VAL:HG12	1:33:C:ARG:HA	4	0.3	0.07	0.32
(1,1062)	1:32:C:VAL:HG13	1:33:C:ARG:HA	4	0.3	0.07	0.32
(1,1062)	1:32:C:VAL:HG21	1:33:C:ARG:HA	4	0.3	0.07	0.32
(1,1062)	1:32:C:VAL:HG22	1:33:C:ARG:HA	4	0.3	0.07	0.32
(1,1062)	1:32:C:VAL:HG23	1:33:C:ARG:HA	4	0.3	0.07	0.32
(1,735)	1:28:C:PRO:HA	2:40:D:PHE:HD1	4	0.3	0.15	0.28
(1,735)	1:28:C:PRO:HA	2:40:D:PHE:HD2	4	0.3	0.15	0.28
(1,1054)	1:28:C:PRO:HA	2:40:D:PHE:HD1	4	0.3	0.15	0.28
(1,1054)	1:28:C:PRO:HA	2:40:D:PHE:HD2	4	0.3	0.15	0.28
(1,711)	1:24:C:ALA:HB1	1:25:C:GLY:HA3	4	0.26	0.05	0.25
(1,711)	1:24:C:ALA:HB2	1:25:C:GLY:HA3	4	0.26	0.05	0.25
(1,711)	1:24:C:ALA:HB3	1:25:C:GLY:HA3	4	0.26	0.05	0.25
(1,771)	1:24:C:ALA:HB1	1:25:C:GLY:HA3	4	0.26	0.05	0.25

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Key	Atom-1	Atom-2	Models[1]	Mean (Å)	SD[1] (Å)	Median (Å)
(1,771)	1:24:C:ALA:HB2	1:25:C:GLY:HA3	4	0.26	0.05	0.25
(1,771)	1:24:C:ALA:HB3	1:25:C:GLY:HA3	4	0.26	0.05	0.25
(1,1030)	1:24:C:ALA:HB1	1:25:C:GLY:HA3	4	0.26	0.05	0.25
(1,1030)	1:24:C:ALA:HB2	1:25:C:GLY:HA3	4	0.26	0.05	0.25
(1,1030)	1:24:C:ALA:HB3	1:25:C:GLY:HA3	4	0.26	0.05	0.25
(1,1090)	1:24:C:ALA:HB1	1:25:C:GLY:HA3	4	0.26	0.05	0.25
(1,1090)	1:24:C:ALA:HB2	1:25:C:GLY:HA3	4	0.26	0.05	0.25
(1,1090)	1:24:C:ALA:HB3	1:25:C:GLY:HA3	4	0.26	0.05	0.25
(1,716)	1:22:C:GLY:H	2:41:D:TYR:HE1	4	0.25	0.22	0.13
(1,716)	1:22:C:GLY:H	2:41:D:TYR:HE2	4	0.25	0.22	0.13
(1,1035)	1:22:C:GLY:H	2:41:D:TYR:HE1	4	0.25	0.22	0.13
(1,1035)	1:22:C:GLY:H	2:41:D:TYR:HE2	4	0.25	0.22	0.13
(1,740)	1:28:C:PRO:HG2	2:40:D:PHE:HD1	4	0.23	0.07	0.24
(1,740)	1:28:C:PRO:HG2	2:40:D:PHE:HD2	4	0.23	0.07	0.24
(1,740)	1:28:C:PRO:HG3	2:40:D:PHE:HD1	4	0.23	0.07	0.24
(1,740)	1:28:C:PRO:HG3	2:40:D:PHE:HD2	4	0.23	0.07	0.24
(1,848)	1:24:C:ALA:H	1:25:C:GLY:HA3	4	0.23	0.16	0.15
(1,1059)	1:28:C:PRO:HG2	2:40:D:PHE:HD1	4	0.23	0.07	0.24
(1,1059)	1:28:C:PRO:HG2	2:40:D:PHE:HD2	4	0.23	0.07	0.24
(1,1059)	1:28:C:PRO:HG3	2:40:D:PHE:HD1	4	0.23	0.07	0.24
(1,1059)	1:28:C:PRO:HG3	2:40:D:PHE:HD2	4	0.23	0.07	0.24
(1,1167)	1:24:C:ALA:H	1:25:C:GLY:HA3	4	0.23	0.16	0.15
(1,830)	1:32:C:VAL:HB	1:33:C:ARG:H	4	0.18	0.07	0.15
(1,1149)	1:32:C:VAL:HB	1:33:C:ARG:H	4	0.18	0.07	0.15
(1,747)	1:23:C:GLY:HA3	1:24:C:ALA:HB1	4	0.17	0.04	0.16
(1,747)	1:23:C:GLY:HA3	1:24:C:ALA:HB2	4	0.17	0.04	0.16
(1,747)	1:23:C:GLY:HA3	1:24:C:ALA:HB3	4	0.17	0.04	0.16
(1,1066)	1:23:C:GLY:HA3	1:24:C:ALA:HB1	4	0.17	0.04	0.16
(1,1066)	1:23:C:GLY:HA3	1:24:C:ALA:HB2	4	0.17	0.04	0.16
(1,1066)	1:23:C:GLY:HA3	1:24:C:ALA:HB3	4	0.17	0.04	0.16
(1,211)	1:5:A:HIS:HA	1:5:A:HIS:HD2	4	0.16	0.05	0.15
(1,530)	1:5:A:HIS:HA	1:5:A:HIS:HD2	4	0.16	0.05	0.15
(1,789)	2:36:D:THR:HG21	2:37:D:PRO:HD3	4	0.14	0.02	0.14
(1,789)	2:36:D:THR:HG22	2:37:D:PRO:HD3	4	0.14	0.02	0.14
(1,789)	2:36:D:THR:HG23	2:37:D:PRO:HD3	4	0.14	0.02	0.14
(1,1108)	2:36:D:THR:HG21	2:37:D:PRO:HD3	4	0.14	0.02	0.14
(1,1108)	2:36:D:THR:HG22	2:37:D:PRO:HD3	4	0.14	0.02	0.14
(1,1108)	2:36:D:THR:HG23	2:37:D:PRO:HD3	4	0.14	0.02	0.14
(1,4)	2:13:B:CYS:HB3	1:25:C:GLY:HA2	3	1.38	0.31	1.22
(1,669)	2:13:B:CYS:HB3	1:25:C:GLY:HA2	3	1.38	0.31	1.22
(1,959)	2:37:D:PRO:HB3	2:37:D:PRO:HD3	3	0.83	0.04	0.83
(1,1278)	2:37:D:PRO:HB3	2:37:D:PRO:HD3	3	0.83	0.04	0.83

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Key	Atom-1	Atom-2	Models[1]	Mean (Å)	SD[1] (Å)	Median (Å)
(1,72)	1:3:A:ALA:HA	2:19:B:PHE:HD1	3	0.82	0.34	0.82
(1,72)	1:3:A:ALA:HA	2:19:B:PHE:HD2	3	0.82	0.34	0.82
(1,334)	1:3:A:ALA:HA	2:19:B:PHE:HD1	3	0.82	0.34	0.82
(1,334)	1:3:A:ALA:HA	2:19:B:PHE:HD2	3	0.82	0.34	0.82
(1,391)	1:3:A:ALA:HA	2:19:B:PHE:HD1	3	0.82	0.34	0.82
(1,391)	1:3:A:ALA:HA	2:19:B:PHE:HD2	3	0.82	0.34	0.82
(1,653)	1:3:A:ALA:HA	2:19:B:PHE:HD1	3	0.82	0.34	0.82
(1,653)	1:3:A:ALA:HA	2:19:B:PHE:HD2	3	0.82	0.34	0.82
(1,41)	1:7:A:PRO:HG2	1:10:A:PHE:HB2	3	0.76	0.6	0.57
(1,41)	1:7:A:PRO:HG3	1:10:A:PHE:HB2	3	0.76	0.6	0.57
(1,75)	1:7:A:PRO:HG2	1:10:A:PHE:HB2	3	0.76	0.6	0.57
(1,75)	1:7:A:PRO:HG3	1:10:A:PHE:HB2	3	0.76	0.6	0.57
(1,360)	1:7:A:PRO:HG2	1:10:A:PHE:HB2	3	0.76	0.6	0.57
(1,360)	1:7:A:PRO:HG3	1:10:A:PHE:HB2	3	0.76	0.6	0.57
(1,394)	1:7:A:PRO:HG2	1:10:A:PHE:HB2	3	0.76	0.6	0.57
(1,394)	1:7:A:PRO:HG3	1:10:A:PHE:HB2	3	0.76	0.6	0.57
(1,52)	1:3:A:ALA:H	2:19:B:PHE:HD1	3	0.64	0.12	0.6
(1,52)	1:3:A:ALA:H	2:19:B:PHE:HD2	3	0.64	0.12	0.6
(1,191)	1:3:A:ALA:H	2:19:B:PHE:HD1	3	0.64	0.12	0.6
(1,191)	1:3:A:ALA:H	2:19:B:PHE:HD2	3	0.64	0.12	0.6
(1,371)	1:3:A:ALA:H	2:19:B:PHE:HD1	3	0.64	0.12	0.6
(1,371)	1:3:A:ALA:H	2:19:B:PHE:HD2	3	0.64	0.12	0.6
(1,510)	1:3:A:ALA:H	2:19:B:PHE:HD1	3	0.64	0.12	0.6
(1,510)	1:3:A:ALA:H	2:19:B:PHE:HD2	3	0.64	0.12	0.6
(1,28)	1:7:A:PRO:HD2	2:20:B:TYR:HD1	3	0.61	0.05	0.58
(1,28)	1:7:A:PRO:HD2	2:20:B:TYR:HD2	3	0.61	0.05	0.58
(1,347)	1:7:A:PRO:HD2	2:20:B:TYR:HD1	3	0.61	0.05	0.58
(1,347)	1:7:A:PRO:HD2	2:20:B:TYR:HD2	3	0.61	0.05	0.58
(1,99)	1:1:A:GLY:H	2:18:B:SER:HB3	3	0.54	0.24	0.58
(1,418)	1:1:A:GLY:H	2:18:B:SER:HB3	3	0.54	0.24	0.58
(1,829)	1:28:C:PRO:HB2	1:31:C:PHE:HD1	3	0.38	0.24	0.33
(1,829)	1:28:C:PRO:HB2	1:31:C:PHE:HD2	3	0.38	0.24	0.33
(1,1148)	1:28:C:PRO:HB2	1:31:C:PHE:HD1	3	0.38	0.24	0.33
(1,1148)	1:28:C:PRO:HB2	1:31:C:PHE:HD2	3	0.38	0.24	0.33
(1,765)	1:25:C:GLY:H	2:40:D:PHE:HB3	3	0.38	0.04	0.37
(1,1084)	1:25:C:GLY:H	2:40:D:PHE:HB3	3	0.38	0.04	0.37
(1,761)	1:25:C:GLY:H	2:41:D:TYR:H	3	0.37	0.22	0.32
(1,1080)	1:25:C:GLY:H	2:41:D:TYR:H	3	0.37	0.22	0.32
(1,12)	1:11:A:VAL:HA	1:24:C:ALA:HB1	3	0.31	0.15	0.23
(1,12)	1:11:A:VAL:HA	1:24:C:ALA:HB2	3	0.31	0.15	0.23
(1,12)	1:11:A:VAL:HA	1:24:C:ALA:HB3	3	0.31	0.15	0.23
(1,659)	1:11:A:VAL:HA	1:24:C:ALA:HB1	3	0.31	0.15	0.23

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Key	Atom-1	Atom-2	Models[1]	Mean (Å)	SD[1] (Å)	Median (Å)
(1,659)	1:11:A:VAL:HA	1:24:C:ALA:HB2	3	0.31	0.15	0.23
(1,659)	1:11:A:VAL:HA	1:24:C:ALA:HB3	3	0.31	0.15	0.23
(1,701)	1:27:C:VAL:HB	2:41:D:TYR:H	3	0.29	0.13	0.28
(1,1020)	1:27:C:VAL:HB	2:41:D:TYR:H	3	0.29	0.13	0.28
(1,290)	2:15:B:THR:H	2:15:B:THR:HB	3	0.28	0.06	0.32
(1,609)	2:15:B:THR:H	2:15:B:THR:HB	3	0.28	0.06	0.32
(1,772)	1:24:C:ALA:H	1:25:C:GLY:H	3	0.13	0.02	0.12
(1,1091)	1:24:C:ALA:H	1:25:C:GLY:H	3	0.13	0.02	0.12
(1,693)	1:29:C:GLU:HG2	2:41:D:TYR:HE1	2	1.14	0.9	1.14
(1,693)	1:29:C:GLU:HG2	2:41:D:TYR:HE2	2	1.14	0.9	1.14
(1,1012)	1:29:C:GLU:HG2	2:41:D:TYR:HE1	2	1.14	0.9	1.14
(1,1012)	1:29:C:GLU:HG2	2:41:D:TYR:HE2	2	1.14	0.9	1.14
(1,800)	1:26:C:HIS:HB2	1:27:C:VAL:HG11	2	1.12	0.96	1.12
(1,800)	1:26:C:HIS:HB2	1:27:C:VAL:HG12	2	1.12	0.96	1.12
(1,800)	1:26:C:HIS:HB2	1:27:C:VAL:HG13	2	1.12	0.96	1.12
(1,1119)	1:26:C:HIS:HB2	1:27:C:VAL:HG11	2	1.12	0.96	1.12
(1,1119)	1:26:C:HIS:HB2	1:27:C:VAL:HG12	2	1.12	0.96	1.12
(1,1119)	1:26:C:HIS:HB2	1:27:C:VAL:HG13	2	1.12	0.96	1.12
(1,48)	2:15:B:THR:HG21	2:17:B:ILE:HA	2	0.96	0.02	0.96
(1,48)	2:15:B:THR:HG22	2:17:B:ILE:HA	2	0.96	0.02	0.96
(1,48)	2:15:B:THR:HG23	2:17:B:ILE:HA	2	0.96	0.02	0.96
(1,367)	2:15:B:THR:HG21	2:17:B:ILE:HA	2	0.96	0.02	0.96
(1,367)	2:15:B:THR:HG22	2:17:B:ILE:HA	2	0.96	0.02	0.96
(1,367)	2:15:B:THR:HG23	2:17:B:ILE:HA	2	0.96	0.02	0.96
(1,35)	1:10:A:PHE:HD1	2:16:B:PRO:HB3	2	0.74	0.03	0.74
(1,35)	1:10:A:PHE:HD2	2:16:B:PRO:HB3	2	0.74	0.03	0.74
(1,354)	1:10:A:PHE:HD1	2:16:B:PRO:HB3	2	0.74	0.03	0.74
(1,354)	1:10:A:PHE:HD2	2:16:B:PRO:HB3	2	0.74	0.03	0.74
(1,682)	1:30:C:TYR:HD1	2:39:D:SER:HB2	2	0.64	0.49	0.64
(1,682)	1:30:C:TYR:HD2	2:39:D:SER:HB2	2	0.64	0.49	0.64
(1,1001)	1:30:C:TYR:HD1	2:39:D:SER:HB2	2	0.64	0.49	0.64
(1,1001)	1:30:C:TYR:HD2	2:39:D:SER:HB2	2	0.64	0.49	0.64
(1,719)	1:31:C:PHE:HD1	2:37:D:PRO:HD2	2	0.63	0.43	0.63
(1,719)	1:31:C:PHE:HD2	2:37:D:PRO:HD2	2	0.63	0.43	0.63
(1,797)	1:26:C:HIS:HD2	1:27:C:VAL:HG21	2	0.63	0.09	0.63
(1,797)	1:26:C:HIS:HD2	1:27:C:VAL:HG22	2	0.63	0.09	0.63
(1,797)	1:26:C:HIS:HD2	1:27:C:VAL:HG23	2	0.63	0.09	0.63
(1,1038)	1:31:C:PHE:HD1	2:37:D:PRO:HD2	2	0.63	0.43	0.63
(1,1038)	1:31:C:PHE:HD2	2:37:D:PRO:HD2	2	0.63	0.43	0.63
(1,1116)	1:26:C:HIS:HD2	1:27:C:VAL:HG21	2	0.63	0.09	0.63
(1,1116)	1:26:C:HIS:HD2	1:27:C:VAL:HG22	2	0.63	0.09	0.63
(1,1116)	1:26:C:HIS:HD2	1:27:C:VAL:HG23	2	0.63	0.09	0.63

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Key	Atom-1	Atom-2	Models[1]	Mean (Å)	SD[1] (Å)	Median (Å)
(1,151)	1:2:A:GLY:H	2:19:B:PHE:H	2	0.55	0.3	0.55
(1,470)	1:2:A:GLY:H	2:19:B:PHE:H	2	0.55	0.3	0.55
(1,738)	1:31:C:PHE:HD1	1:32:C:VAL:HA	2	0.54	0.04	0.54
(1,738)	1:31:C:PHE:HD2	1:32:C:VAL:HA	2	0.54	0.04	0.54
(1,1057)	1:31:C:PHE:HD1	1:32:C:VAL:HA	2	0.54	0.04	0.54
(1,1057)	1:31:C:PHE:HD2	1:32:C:VAL:HA	2	0.54	0.04	0.54
(1,777)	1:31:C:PHE:HD1	2:37:D:PRO:HG2	2	0.54	0.22	0.54
(1,777)	1:31:C:PHE:HD1	2:37:D:PRO:HG3	2	0.54	0.22	0.54
(1,777)	1:31:C:PHE:HD2	2:37:D:PRO:HG2	2	0.54	0.22	0.54
(1,777)	1:31:C:PHE:HD2	2:37:D:PRO:HG3	2	0.54	0.22	0.54
(1,1096)	1:31:C:PHE:HD1	2:37:D:PRO:HG2	2	0.54	0.22	0.54
(1,1096)	1:31:C:PHE:HD1	2:37:D:PRO:HG3	2	0.54	0.22	0.54
(1,1096)	1:31:C:PHE:HD2	2:37:D:PRO:HG2	2	0.54	0.22	0.54
(1,1096)	1:31:C:PHE:HD2	2:37:D:PRO:HG3	2	0.54	0.22	0.54
(1,691)	1:31:C:PHE:HD1	2:37:D:PRO:HB3	2	0.52	0.15	0.52
(1,691)	1:31:C:PHE:HD2	2:37:D:PRO:HB3	2	0.52	0.15	0.52
(1,1010)	1:31:C:PHE:HD1	2:37:D:PRO:HB3	2	0.52	0.15	0.52
(1,1010)	1:31:C:PHE:HD2	2:37:D:PRO:HB3	2	0.52	0.15	0.52
(1,844)	1:23:C:GLY:HA2	1:24:C:ALA:HB1	2	0.5	0.07	0.5
(1,844)	1:23:C:GLY:HA2	1:24:C:ALA:HB2	2	0.5	0.07	0.5
(1,844)	1:23:C:GLY:HA2	1:24:C:ALA:HB3	2	0.5	0.07	0.5
(1,1163)	1:23:C:GLY:HA2	1:24:C:ALA:HB1	2	0.5	0.07	0.5
(1,1163)	1:23:C:GLY:HA2	1:24:C:ALA:HB2	2	0.5	0.07	0.5
(1,1163)	1:23:C:GLY:HA2	1:24:C:ALA:HB3	2	0.5	0.07	0.5
(1,129)	1:4:A:GLY:H	2:19:B:PHE:H	2	0.48	0.24	0.48
(1,448)	1:4:A:GLY:H	2:19:B:PHE:H	2	0.48	0.24	0.48
(1,82)	1:10:A:PHE:HD1	1:11:A:VAL:HA	2	0.46	0.2	0.46
(1,82)	1:10:A:PHE:HD2	1:11:A:VAL:HA	2	0.46	0.2	0.46
(1,401)	1:10:A:PHE:HD1	1:11:A:VAL:HA	2	0.46	0.2	0.46
(1,401)	1:10:A:PHE:HD2	1:11:A:VAL:HA	2	0.46	0.2	0.46
(1,34)	1:6:A:VAL:HB	2:20:B:TYR:HB2	2	0.43	0.22	0.43
(1,34)	1:6:A:VAL:HB	2:20:B:TYR:HB3	2	0.43	0.22	0.43
(1,353)	1:6:A:VAL:HB	2:20:B:TYR:HB2	2	0.43	0.22	0.43
(1,353)	1:6:A:VAL:HB	2:20:B:TYR:HB3	2	0.43	0.22	0.43
(1,109)	1:4:A:GLY:H	2:19:B:PHE:HB3	2	0.42	0.01	0.42
(1,428)	1:4:A:GLY:H	2:19:B:PHE:HB3	2	0.42	0.01	0.42
(1,166)	1:4:A:GLY:HA3	1:6:A:VAL:H	2	0.41	0.08	0.41
(1,485)	1:4:A:GLY:HA3	1:6:A:VAL:H	2	0.41	0.08	0.41
(1,148)	1:10:A:PHE:HB3	1:11:A:VAL:HG11	2	0.4	0.09	0.4
(1,148)	1:10:A:PHE:HB3	1:11:A:VAL:HG12	2	0.4	0.09	0.4
(1,148)	1:10:A:PHE:HB3	1:11:A:VAL:HG13	2	0.4	0.09	0.4
(1,148)	1:10:A:PHE:HB3	1:11:A:VAL:HG21	2	0.4	0.09	0.4

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Key	Atom-1	Atom-2	Models[1]	Mean (Å)	SD[1] (Å)	Median (Å)
(1,148)	1:10:A:PHE:HB3	1:11:A:VAL:HG22	2	0.4	0.09	0.4
(1,148)	1:10:A:PHE:HB3	1:11:A:VAL:HG23	2	0.4	0.09	0.4
(1,467)	1:10:A:PHE:HB3	1:11:A:VAL:HG11	2	0.4	0.09	0.4
(1,467)	1:10:A:PHE:HB3	1:11:A:VAL:HG12	2	0.4	0.09	0.4
(1,467)	1:10:A:PHE:HB3	1:11:A:VAL:HG13	2	0.4	0.09	0.4
(1,467)	1:10:A:PHE:HB3	1:11:A:VAL:HG21	2	0.4	0.09	0.4
(1,467)	1:10:A:PHE:HB3	1:11:A:VAL:HG22	2	0.4	0.09	0.4
(1,467)	1:10:A:PHE:HB3	1:11:A:VAL:HG23	2	0.4	0.09	0.4
(1,120)	1:3:A:ALA:HA	1:4:A:GLY:H	2	0.39	0.11	0.39
(1,439)	1:3:A:ALA:HA	1:4:A:GLY:H	2	0.39	0.11	0.39
(1,26)	1:9:A:TYR:HD1	2:18:B:SER:HB2	2	0.38	0.06	0.38
(1,26)	1:9:A:TYR:HD2	2:18:B:SER:HB2	2	0.38	0.06	0.38
(1,345)	1:9:A:TYR:HD1	2:18:B:SER:HB2	2	0.38	0.06	0.38
(1,345)	1:9:A:TYR:HD2	2:18:B:SER:HB2	2	0.38	0.06	0.38
(1,149)	1:11:A:VAL:HG11	1:12:A:ARG:H	2	0.34	0.21	0.34
(1,149)	1:11:A:VAL:HG12	1:12:A:ARG:H	2	0.34	0.21	0.34
(1,149)	1:11:A:VAL:HG13	1:12:A:ARG:H	2	0.34	0.21	0.34
(1,149)	1:11:A:VAL:HG21	1:12:A:ARG:H	2	0.34	0.21	0.34
(1,149)	1:11:A:VAL:HG22	1:12:A:ARG:H	2	0.34	0.21	0.34
(1,149)	1:11:A:VAL:HG23	1:12:A:ARG:H	2	0.34	0.21	0.34
(1,468)	1:11:A:VAL:HG11	1:12:A:ARG:H	2	0.34	0.21	0.34
(1,468)	1:11:A:VAL:HG12	1:12:A:ARG:H	2	0.34	0.21	0.34
(1,468)	1:11:A:VAL:HG13	1:12:A:ARG:H	2	0.34	0.21	0.34
(1,468)	1:11:A:VAL:HG21	1:12:A:ARG:H	2	0.34	0.21	0.34
(1,468)	1:11:A:VAL:HG22	1:12:A:ARG:H	2	0.34	0.21	0.34
(1,468)	1:11:A:VAL:HG23	1:12:A:ARG:H	2	0.34	0.21	0.34
(1,964)	2:38:D:ILE:H	2:38:D:ILE:HB	2	0.32	0.05	0.32
(1,1283)	2:38:D:ILE:H	2:38:D:ILE:HB	2	0.32	0.05	0.32
(1,197)	2:15:B:THR:HB	2:16:B:PRO:HD2	2	0.29	0.07	0.29
(1,516)	2:15:B:THR:HB	2:16:B:PRO:HD2	2	0.29	0.07	0.29
(1,769)	1:23:C:GLY:HA2	1:24:C:ALA:H	2	0.24	0.01	0.24
(1,1088)	1:23:C:GLY:HA2	1:24:C:ALA:H	2	0.24	0.01	0.24
(1,24)	1:7:A:PRO:HD3	2:19:B:PHE:HD1	2	0.23	0.11	0.23
(1,24)	1:7:A:PRO:HD3	2:19:B:PHE:HD2	2	0.23	0.11	0.23
(1,343)	1:7:A:PRO:HD3	2:19:B:PHE:HD1	2	0.23	0.11	0.23
(1,343)	1:7:A:PRO:HD3	2:19:B:PHE:HD2	2	0.23	0.11	0.23
(1,248)	1:10:A:PHE:HB3	1:11:A:VAL:H	2	0.22	0.12	0.22
(1,567)	1:10:A:PHE:HB3	1:11:A:VAL:H	2	0.22	0.12	0.22
(1,89)	2:15:B:THR:HG21	2:16:B:PRO:HA	2	0.2	0.02	0.2
(1,89)	2:15:B:THR:HG22	2:16:B:PRO:HA	2	0.2	0.02	0.2
(1,89)	2:15:B:THR:HG23	2:16:B:PRO:HA	2	0.2	0.02	0.2
(1,125)	2:19:B:PHE:HA	2:20:B:TYR:HD1	2	0.2	0.02	0.2

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Key	Atom-1	Atom-2	Models[1]	Mean (Å)	SD[1] (Å)	Median (Å)
(1,125)	2:19:B:PHE:HA	2:20:B:TYR:HD2	2	0.2	0.02	0.2
(1,408)	2:15:B:THR:HG21	2:16:B:PRO:HA	2	0.2	0.02	0.2
(1,408)	2:15:B:THR:HG22	2:16:B:PRO:HA	2	0.2	0.02	0.2
(1,408)	2:15:B:THR:HG23	2:16:B:PRO:HA	2	0.2	0.02	0.2
(1,444)	2:19:B:PHE:HA	2:20:B:TYR:HD1	2	0.2	0.02	0.2
(1,444)	2:19:B:PHE:HA	2:20:B:TYR:HD2	2	0.2	0.02	0.2
(1,791)	1:26:C:HIS:HA	1:27:C:VAL:HG21	2	0.18	0.01	0.18
(1,791)	1:26:C:HIS:HA	1:27:C:VAL:HG22	2	0.18	0.01	0.18
(1,791)	1:26:C:HIS:HA	1:27:C:VAL:HG23	2	0.18	0.01	0.18
(1,1110)	1:26:C:HIS:HA	1:27:C:VAL:HG21	2	0.18	0.01	0.18
(1,1110)	1:26:C:HIS:HA	1:27:C:VAL:HG22	2	0.18	0.01	0.18
(1,1110)	1:26:C:HIS:HA	1:27:C:VAL:HG23	2	0.18	0.01	0.18
(1,824)	1:27:C:VAL:H	2:41:D:TYR:HD1	2	0.16	0.04	0.16
(1,824)	1:27:C:VAL:H	2:41:D:TYR:HD2	2	0.16	0.04	0.16
(1,1143)	1:27:C:VAL:H	2:41:D:TYR:HD1	2	0.16	0.04	0.16
(1,1143)	1:27:C:VAL:H	2:41:D:TYR:HD2	2	0.16	0.04	0.16
(1,69)	1:11:A:VAL:HG11	1:12:A:ARG:HG2	2	0.13	0.0	0.13
(1,69)	1:11:A:VAL:HG11	1:12:A:ARG:HG3	2	0.13	0.0	0.13
(1,69)	1:11:A:VAL:HG12	1:12:A:ARG:HG2	2	0.13	0.0	0.13
(1,69)	1:11:A:VAL:HG12	1:12:A:ARG:HG3	2	0.13	0.0	0.13
(1,69)	1:11:A:VAL:HG13	1:12:A:ARG:HG2	2	0.13	0.0	0.13
(1,69)	1:11:A:VAL:HG13	1:12:A:ARG:HG3	2	0.13	0.0	0.13
(1,69)	1:11:A:VAL:HG21	1:12:A:ARG:HG2	2	0.13	0.0	0.13
(1,69)	1:11:A:VAL:HG21	1:12:A:ARG:HG3	2	0.13	0.0	0.13
(1,69)	1:11:A:VAL:HG22	1:12:A:ARG:HG2	2	0.13	0.0	0.13
(1,69)	1:11:A:VAL:HG22	1:12:A:ARG:HG3	2	0.13	0.0	0.13
(1,69)	1:11:A:VAL:HG23	1:12:A:ARG:HG2	2	0.13	0.0	0.13
(1,69)	1:11:A:VAL:HG23	1:12:A:ARG:HG3	2	0.13	0.0	0.13
(1,388)	1:11:A:VAL:HG11	1:12:A:ARG:HG2	2	0.13	0.0	0.13
(1,388)	1:11:A:VAL:HG11	1:12:A:ARG:HG3	2	0.13	0.0	0.13
(1,388)	1:11:A:VAL:HG12	1:12:A:ARG:HG2	2	0.13	0.0	0.13
(1,388)	1:11:A:VAL:HG12	1:12:A:ARG:HG3	2	0.13	0.0	0.13
(1,388)	1:11:A:VAL:HG13	1:12:A:ARG:HG2	2	0.13	0.0	0.13
(1,388)	1:11:A:VAL:HG13	1:12:A:ARG:HG3	2	0.13	0.0	0.13
(1,388)	1:11:A:VAL:HG21	1:12:A:ARG:HG2	2	0.13	0.0	0.13
(1,388)	1:11:A:VAL:HG21	1:12:A:ARG:HG3	2	0.13	0.0	0.13
(1,388)	1:11:A:VAL:HG22	1:12:A:ARG:HG2	2	0.13	0.0	0.13
(1,388)	1:11:A:VAL:HG22	1:12:A:ARG:HG3	2	0.13	0.0	0.13
(1,388)	1:11:A:VAL:HG23	1:12:A:ARG:HG2	2	0.13	0.0	0.13
(1,388)	1:11:A:VAL:HG23	1:12:A:ARG:HG3	2	0.13	0.0	0.13
(1,909)	1:31:C:PHE:HA	1:31:C:PHE:HD1	2	0.11	0.01	0.11
(1,909)	1:31:C:PHE:HA	1:31:C:PHE:HD2	2	0.11	0.01	0.11

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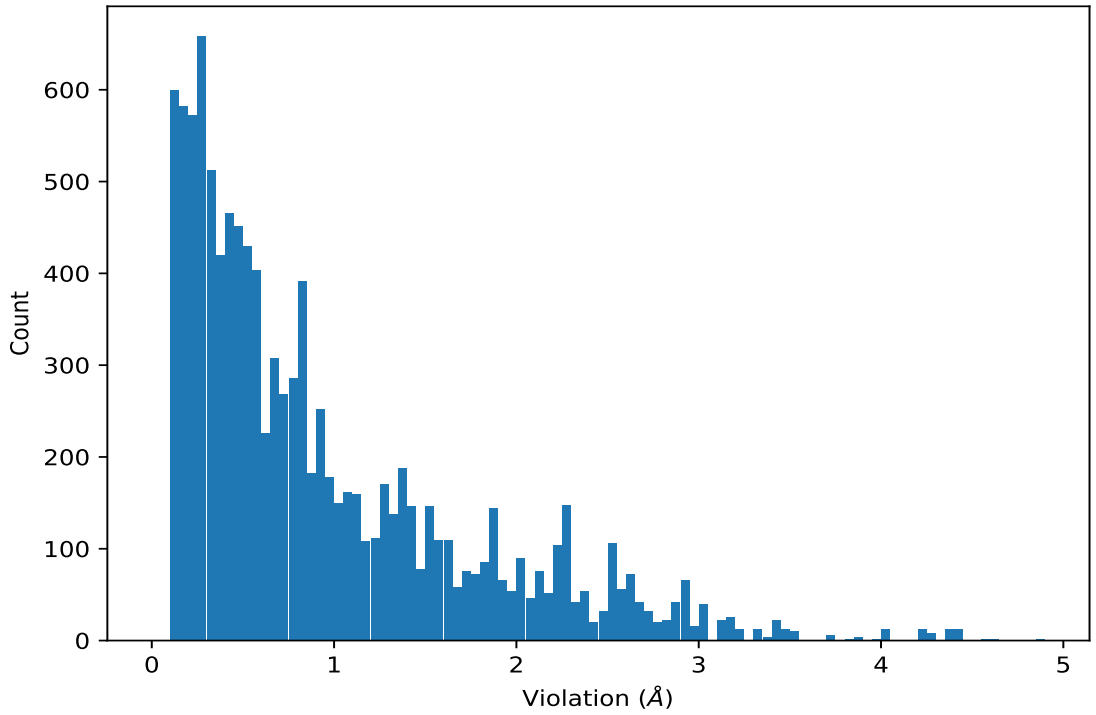
Key	Atom-1	Atom-2	Models[1]	Mean (Å)	SD[1] (Å)	Median (Å)
(1,1228)	1:31:C:PHE:HA	1:31:C:PHE:HD1	2	0.11	0.01	0.11
(1,1228)	1:31:C:PHE:HA	1:31:C:PHE:HD2	2	0.11	0.01	0.11

[1]Number of violated models, [2]Standard deviation

9.5 All violated distance restraints (i)

9.5.1 Histogram : Distribution of distance violations (i)

The following histogram shows the distribution of the absolute value of the violation for all violated restraints in the ensemble.



9.5.2 Table : All distance violations (i)

The following table lists the absolute value of the violation for each restraint in the ensemble sorted by its value. The Key (restraint list ID, restraint ID) is the unique identifier for a given restraint. Rows with same key represent combinatorial or ambiguous restraints and are counted as a single restraint.

Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,1176)	2:19:B:PHE:HB3	2:38:D:ILE:HG13	5	4.87
(1,857)	2:19:B:PHE:HB3	2:38:D:ILE:HG13	5	4.87

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,1176)	2:19:B:PHE:HB3	2:38:D:ILE:HG13	20	4.63
(1,857)	2:19:B:PHE:HB3	2:38:D:ILE:HG13	20	4.63
(1,664)	2:19:B:PHE:HB2	2:38:D:ILE:HG13	5	4.57
(1,17)	2:19:B:PHE:HB2	2:38:D:ILE:HG13	5	4.57
(1,674)	2:17:B:ILE:HD11	2:40:D:PHE:HE1	7	4.44
(1,674)	2:17:B:ILE:HD11	2:40:D:PHE:HE2	7	4.44
(1,674)	2:17:B:ILE:HD12	2:40:D:PHE:HE1	7	4.44
(1,674)	2:17:B:ILE:HD12	2:40:D:PHE:HE2	7	4.44
(1,674)	2:17:B:ILE:HD13	2:40:D:PHE:HE1	7	4.44
(1,674)	2:17:B:ILE:HD13	2:40:D:PHE:HE2	7	4.44
(1,9)	2:17:B:ILE:HD11	2:40:D:PHE:HE1	7	4.44
(1,9)	2:17:B:ILE:HD11	2:40:D:PHE:HE2	7	4.44
(1,9)	2:17:B:ILE:HD12	2:40:D:PHE:HE1	7	4.44
(1,9)	2:17:B:ILE:HD12	2:40:D:PHE:HE2	7	4.44
(1,9)	2:17:B:ILE:HD13	2:40:D:PHE:HE1	7	4.44
(1,9)	2:17:B:ILE:HD13	2:40:D:PHE:HE2	7	4.44
(1,665)	2:19:B:PHE:HE1	2:38:D:ILE:HD11	17	4.39
(1,665)	2:19:B:PHE:HE1	2:38:D:ILE:HD12	17	4.39
(1,665)	2:19:B:PHE:HE1	2:38:D:ILE:HD13	17	4.39
(1,665)	2:19:B:PHE:HE2	2:38:D:ILE:HD11	17	4.39
(1,665)	2:19:B:PHE:HE2	2:38:D:ILE:HD12	17	4.39
(1,665)	2:19:B:PHE:HE2	2:38:D:ILE:HD13	17	4.39
(1,18)	2:19:B:PHE:HE1	2:38:D:ILE:HD11	17	4.39
(1,18)	2:19:B:PHE:HE1	2:38:D:ILE:HD12	17	4.39
(1,18)	2:19:B:PHE:HE1	2:38:D:ILE:HD13	17	4.39
(1,18)	2:19:B:PHE:HE2	2:38:D:ILE:HD11	17	4.39
(1,18)	2:19:B:PHE:HE2	2:38:D:ILE:HD12	17	4.39
(1,18)	2:19:B:PHE:HE2	2:38:D:ILE:HD13	17	4.39
(1,1176)	2:19:B:PHE:HB3	2:38:D:ILE:HG13	14	4.26
(1,857)	2:19:B:PHE:HB3	2:38:D:ILE:HG13	14	4.26
(1,664)	2:19:B:PHE:HB2	2:38:D:ILE:HG13	20	4.25
(1,361)	1:10:A:PHE:HE1	2:14:B:GLY:HA3	18	4.25
(1,361)	1:10:A:PHE:HE2	2:14:B:GLY:HA3	18	4.25
(1,42)	1:10:A:PHE:HE1	2:14:B:GLY:HA3	18	4.25
(1,42)	1:10:A:PHE:HE2	2:14:B:GLY:HA3	18	4.25
(1,17)	2:19:B:PHE:HB2	2:38:D:ILE:HG13	20	4.25
(1,665)	2:19:B:PHE:HE1	2:38:D:ILE:HD11	14	4.21
(1,665)	2:19:B:PHE:HE1	2:38:D:ILE:HD12	14	4.21
(1,665)	2:19:B:PHE:HE1	2:38:D:ILE:HD13	14	4.21
(1,665)	2:19:B:PHE:HE2	2:38:D:ILE:HD11	14	4.21
(1,665)	2:19:B:PHE:HE2	2:38:D:ILE:HD12	14	4.21
(1,665)	2:19:B:PHE:HE2	2:38:D:ILE:HD13	14	4.21

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,18)	2:19:B:PHE:HE1	2:38:D:ILE:HD11	14	4.21
(1,18)	2:19:B:PHE:HE1	2:38:D:ILE:HD12	14	4.21
(1,18)	2:19:B:PHE:HE1	2:38:D:ILE:HD13	14	4.21
(1,18)	2:19:B:PHE:HE2	2:38:D:ILE:HD11	14	4.21
(1,18)	2:19:B:PHE:HE2	2:38:D:ILE:HD12	14	4.21
(1,18)	2:19:B:PHE:HE2	2:38:D:ILE:HD13	14	4.21
(1,674)	2:17:B:ILE:HD11	2:40:D:PHE:HE1	14	4.03
(1,674)	2:17:B:ILE:HD11	2:40:D:PHE:HE2	14	4.03
(1,674)	2:17:B:ILE:HD12	2:40:D:PHE:HE1	14	4.03
(1,674)	2:17:B:ILE:HD12	2:40:D:PHE:HE2	14	4.03
(1,674)	2:17:B:ILE:HD13	2:40:D:PHE:HE1	14	4.03
(1,674)	2:17:B:ILE:HD13	2:40:D:PHE:HE2	14	4.03
(1,9)	2:17:B:ILE:HD11	2:40:D:PHE:HE1	14	4.03
(1,9)	2:17:B:ILE:HD11	2:40:D:PHE:HE2	14	4.03
(1,9)	2:17:B:ILE:HD12	2:40:D:PHE:HE1	14	4.03
(1,9)	2:17:B:ILE:HD12	2:40:D:PHE:HE2	14	4.03
(1,9)	2:17:B:ILE:HD13	2:40:D:PHE:HE1	14	4.03
(1,9)	2:17:B:ILE:HD13	2:40:D:PHE:HE2	14	4.03
(1,1176)	2:19:B:PHE:HB3	2:38:D:ILE:HG13	9	3.98
(1,857)	2:19:B:PHE:HB3	2:38:D:ILE:HG13	9	3.98
(1,1017)	1:31:C:PHE:HE1	2:35:D:GLY:HA3	18	3.85
(1,1017)	1:31:C:PHE:HE2	2:35:D:GLY:HA3	18	3.85
(1,698)	1:31:C:PHE:HE1	2:35:D:GLY:HA3	18	3.85
(1,698)	1:31:C:PHE:HE2	2:35:D:GLY:HA3	18	3.85
(1,664)	2:19:B:PHE:HB2	2:38:D:ILE:HG13	14	3.83
(1,17)	2:19:B:PHE:HB2	2:38:D:ILE:HG13	14	3.83
(1,664)	2:19:B:PHE:HB2	2:38:D:ILE:HG13	9	3.74
(1,17)	2:19:B:PHE:HB2	2:38:D:ILE:HG13	9	3.74
(1,503)	1:7:A:PRO:HB3	1:10:A:PHE:HE1	17	3.72
(1,503)	1:7:A:PRO:HB3	1:10:A:PHE:HE2	17	3.72
(1,184)	1:7:A:PRO:HB3	1:10:A:PHE:HE1	17	3.72
(1,184)	1:7:A:PRO:HB3	1:10:A:PHE:HE2	17	3.72
(1,417)	2:17:B:ILE:HD11	2:39:D:SER:HB3	9	3.54
(1,417)	2:17:B:ILE:HD12	2:39:D:SER:HB3	9	3.54
(1,417)	2:17:B:ILE:HD13	2:39:D:SER:HB3	9	3.54
(1,98)	2:17:B:ILE:HD11	2:39:D:SER:HB3	9	3.54
(1,98)	2:17:B:ILE:HD12	2:39:D:SER:HB3	9	3.54
(1,98)	2:17:B:ILE:HD13	2:39:D:SER:HB3	9	3.54
(1,1017)	1:31:C:PHE:HE1	2:35:D:GLY:HA3	17	3.52
(1,1017)	1:31:C:PHE:HE2	2:35:D:GLY:HA3	17	3.52
(1,698)	1:31:C:PHE:HE1	2:35:D:GLY:HA3	17	3.52
(1,698)	1:31:C:PHE:HE2	2:35:D:GLY:HA3	17	3.52

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,665)	2:19:B:PHE:HE1	2:38:D:ILE:HD11	5	3.48
(1,665)	2:19:B:PHE:HE1	2:38:D:ILE:HD12	5	3.48
(1,665)	2:19:B:PHE:HE1	2:38:D:ILE:HD13	5	3.48
(1,665)	2:19:B:PHE:HE2	2:38:D:ILE:HD11	5	3.48
(1,665)	2:19:B:PHE:HE2	2:38:D:ILE:HD12	5	3.48
(1,665)	2:19:B:PHE:HE2	2:38:D:ILE:HD13	5	3.48
(1,18)	2:19:B:PHE:HE1	2:38:D:ILE:HD11	5	3.48
(1,18)	2:19:B:PHE:HE1	2:38:D:ILE:HD12	5	3.48
(1,18)	2:19:B:PHE:HE1	2:38:D:ILE:HD13	5	3.48
(1,18)	2:19:B:PHE:HE2	2:38:D:ILE:HD11	5	3.48
(1,18)	2:19:B:PHE:HE2	2:38:D:ILE:HD12	5	3.48
(1,18)	2:19:B:PHE:HE2	2:38:D:ILE:HD13	5	3.48
(1,657)	2:19:B:PHE:HD1	2:38:D:ILE:HD11	14	3.44
(1,657)	2:19:B:PHE:HD1	2:38:D:ILE:HD12	14	3.44
(1,657)	2:19:B:PHE:HD1	2:38:D:ILE:HD13	14	3.44
(1,657)	2:19:B:PHE:HD2	2:38:D:ILE:HD11	14	3.44
(1,657)	2:19:B:PHE:HD2	2:38:D:ILE:HD12	14	3.44
(1,657)	2:19:B:PHE:HD2	2:38:D:ILE:HD13	14	3.44
(1,520)	2:17:B:ILE:HG13	2:40:D:PHE:HB3	16	3.44
(1,201)	2:17:B:ILE:HG13	2:40:D:PHE:HB3	16	3.44
(1,10)	2:19:B:PHE:HD1	2:38:D:ILE:HD11	14	3.44
(1,10)	2:19:B:PHE:HD1	2:38:D:ILE:HD12	14	3.44
(1,10)	2:19:B:PHE:HD1	2:38:D:ILE:HD13	14	3.44
(1,10)	2:19:B:PHE:HD2	2:38:D:ILE:HD11	14	3.44
(1,10)	2:19:B:PHE:HD2	2:38:D:ILE:HD12	14	3.44
(1,10)	2:19:B:PHE:HD2	2:38:D:ILE:HD13	14	3.44
(1,491)	1:7:A:PRO:HB2	1:10:A:PHE:HE1	17	3.43
(1,491)	1:7:A:PRO:HB2	1:10:A:PHE:HE2	17	3.43
(1,172)	1:7:A:PRO:HB2	1:10:A:PHE:HE1	17	3.43
(1,172)	1:7:A:PRO:HB2	1:10:A:PHE:HE2	17	3.43
(1,1002)	1:30:C:TYR:HE1	2:39:D:SER:HB2	12	3.41
(1,1002)	1:30:C:TYR:HE2	2:39:D:SER:HB2	12	3.41
(1,683)	1:30:C:TYR:HE1	2:39:D:SER:HB2	12	3.41
(1,683)	1:30:C:TYR:HE2	2:39:D:SER:HB2	12	3.41
(1,673)	2:17:B:ILE:HG13	2:40:D:PHE:HB2	16	3.37
(1,8)	2:17:B:ILE:HG13	2:40:D:PHE:HB2	16	3.37
(1,1176)	2:19:B:PHE:HB3	2:38:D:ILE:HG13	10	3.36
(1,857)	2:19:B:PHE:HB3	2:38:D:ILE:HG13	10	3.36
(1,674)	2:17:B:ILE:HD11	2:40:D:PHE:HE1	6	3.34
(1,674)	2:17:B:ILE:HD11	2:40:D:PHE:HE2	6	3.34
(1,674)	2:17:B:ILE:HD12	2:40:D:PHE:HE1	6	3.34
(1,674)	2:17:B:ILE:HD12	2:40:D:PHE:HE2	6	3.34

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,674)	2:17:B:ILE:HD13	2:40:D:PHE:HE1	6	3.34
(1,674)	2:17:B:ILE:HD13	2:40:D:PHE:HE2	6	3.34
(1,9)	2:17:B:ILE:HD11	2:40:D:PHE:HE1	6	3.34
(1,9)	2:17:B:ILE:HD11	2:40:D:PHE:HE2	6	3.34
(1,9)	2:17:B:ILE:HD12	2:40:D:PHE:HE1	6	3.34
(1,9)	2:17:B:ILE:HD12	2:40:D:PHE:HE2	6	3.34
(1,9)	2:17:B:ILE:HD13	2:40:D:PHE:HE1	6	3.34
(1,9)	2:17:B:ILE:HD13	2:40:D:PHE:HE2	6	3.34
(1,503)	1:7:A:PRO:HB3	1:10:A:PHE:HE1	16	3.24
(1,503)	1:7:A:PRO:HB3	1:10:A:PHE:HE2	16	3.24
(1,184)	1:7:A:PRO:HB3	1:10:A:PHE:HE1	16	3.24
(1,184)	1:7:A:PRO:HB3	1:10:A:PHE:HE2	16	3.24
(1,1073)	2:18:B:SER:HB3	2:38:D:ILE:HD11	19	3.22
(1,1073)	2:18:B:SER:HB3	2:38:D:ILE:HD12	19	3.22
(1,1073)	2:18:B:SER:HB3	2:38:D:ILE:HD13	19	3.22
(1,754)	2:18:B:SER:HB3	2:38:D:ILE:HD11	19	3.22
(1,754)	2:18:B:SER:HB3	2:38:D:ILE:HD12	19	3.22
(1,754)	2:18:B:SER:HB3	2:38:D:ILE:HD13	19	3.22
(1,1176)	2:19:B:PHE:HB3	2:38:D:ILE:HG13	15	3.2
(1,857)	2:19:B:PHE:HB3	2:38:D:ILE:HG13	15	3.2
(1,1169)	2:36:D:THR:HB	2:38:D:ILE:HD11	13	3.19
(1,1169)	2:36:D:THR:HB	2:38:D:ILE:HD12	13	3.19
(1,1169)	2:36:D:THR:HB	2:38:D:ILE:HD13	13	3.19
(1,850)	2:36:D:THR:HB	2:38:D:ILE:HD11	13	3.19
(1,850)	2:36:D:THR:HB	2:38:D:ILE:HD12	13	3.19
(1,850)	2:36:D:THR:HB	2:38:D:ILE:HD13	13	3.19
(1,664)	2:19:B:PHE:HB2	2:38:D:ILE:HG13	10	3.19
(1,17)	2:19:B:PHE:HB2	2:38:D:ILE:HG13	10	3.19
(1,515)	2:15:B:THR:HG21	2:17:B:ILE:HD11	15	3.17
(1,515)	2:15:B:THR:HG21	2:17:B:ILE:HD12	15	3.17
(1,515)	2:15:B:THR:HG21	2:17:B:ILE:HD13	15	3.17
(1,515)	2:15:B:THR:HG22	2:17:B:ILE:HD11	15	3.17
(1,515)	2:15:B:THR:HG22	2:17:B:ILE:HD12	15	3.17
(1,515)	2:15:B:THR:HG22	2:17:B:ILE:HD13	15	3.17
(1,515)	2:15:B:THR:HG23	2:17:B:ILE:HD11	15	3.17
(1,515)	2:15:B:THR:HG23	2:17:B:ILE:HD12	15	3.17
(1,515)	2:15:B:THR:HG23	2:17:B:ILE:HD13	15	3.17
(1,196)	2:15:B:THR:HG21	2:17:B:ILE:HD11	15	3.17
(1,196)	2:15:B:THR:HG21	2:17:B:ILE:HD12	15	3.17
(1,196)	2:15:B:THR:HG21	2:17:B:ILE:HD13	15	3.17
(1,196)	2:15:B:THR:HG22	2:17:B:ILE:HD11	15	3.17
(1,196)	2:15:B:THR:HG22	2:17:B:ILE:HD12	15	3.17

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,196)	2:15:B:THR:HG22	2:17:B:ILE:HD13	15	3.17
(1,196)	2:15:B:THR:HG23	2:17:B:ILE:HD11	15	3.17
(1,196)	2:15:B:THR:HG23	2:17:B:ILE:HD12	15	3.17
(1,196)	2:15:B:THR:HG23	2:17:B:ILE:HD13	15	3.17
(1,1046)	1:27:C:VAL:HB	2:41:D:TYR:HE1	1	3.12
(1,1046)	1:27:C:VAL:HB	2:41:D:TYR:HE2	1	3.12
(1,727)	1:27:C:VAL:HB	2:41:D:TYR:HE1	1	3.12
(1,727)	1:27:C:VAL:HB	2:41:D:TYR:HE2	1	3.12
(1,1171)	2:36:D:THR:HG21	2:38:D:ILE:HD11	5	3.11
(1,1171)	2:36:D:THR:HG21	2:38:D:ILE:HD12	5	3.11
(1,1171)	2:36:D:THR:HG21	2:38:D:ILE:HD13	5	3.11
(1,1171)	2:36:D:THR:HG22	2:38:D:ILE:HD11	5	3.11
(1,1171)	2:36:D:THR:HG22	2:38:D:ILE:HD12	5	3.11
(1,1171)	2:36:D:THR:HG22	2:38:D:ILE:HD13	5	3.11
(1,1171)	2:36:D:THR:HG23	2:38:D:ILE:HD11	5	3.11
(1,1171)	2:36:D:THR:HG23	2:38:D:ILE:HD12	5	3.11
(1,1171)	2:36:D:THR:HG23	2:38:D:ILE:HD13	5	3.11
(1,852)	2:36:D:THR:HG21	2:38:D:ILE:HD11	5	3.11
(1,852)	2:36:D:THR:HG21	2:38:D:ILE:HD12	5	3.11
(1,852)	2:36:D:THR:HG21	2:38:D:ILE:HD13	5	3.11
(1,852)	2:36:D:THR:HG22	2:38:D:ILE:HD11	5	3.11
(1,852)	2:36:D:THR:HG22	2:38:D:ILE:HD12	5	3.11
(1,852)	2:36:D:THR:HG22	2:38:D:ILE:HD13	5	3.11
(1,852)	2:36:D:THR:HG23	2:38:D:ILE:HD11	5	3.11
(1,852)	2:36:D:THR:HG23	2:38:D:ILE:HD12	5	3.11
(1,852)	2:36:D:THR:HG23	2:38:D:ILE:HD13	5	3.11
(1,657)	2:19:B:PHE:HD1	2:38:D:ILE:HD11	5	3.05
(1,657)	2:19:B:PHE:HD1	2:38:D:ILE:HD12	5	3.05
(1,657)	2:19:B:PHE:HD1	2:38:D:ILE:HD13	5	3.05
(1,657)	2:19:B:PHE:HD2	2:38:D:ILE:HD11	5	3.05
(1,657)	2:19:B:PHE:HD2	2:38:D:ILE:HD12	5	3.05
(1,657)	2:19:B:PHE:HD2	2:38:D:ILE:HD13	5	3.05
(1,491)	1:7:A:PRO:HB2	1:10:A:PHE:HE1	16	3.05
(1,491)	1:7:A:PRO:HB2	1:10:A:PHE:HE2	16	3.05
(1,172)	1:7:A:PRO:HB2	1:10:A:PHE:HE1	16	3.05
(1,172)	1:7:A:PRO:HB2	1:10:A:PHE:HE2	16	3.05
(1,10)	2:19:B:PHE:HD1	2:38:D:ILE:HD11	5	3.05
(1,10)	2:19:B:PHE:HD1	2:38:D:ILE:HD12	5	3.05
(1,10)	2:19:B:PHE:HD1	2:38:D:ILE:HD13	5	3.05
(1,10)	2:19:B:PHE:HD2	2:38:D:ILE:HD11	5	3.05
(1,10)	2:19:B:PHE:HD2	2:38:D:ILE:HD12	5	3.05
(1,10)	2:19:B:PHE:HD2	2:38:D:ILE:HD13	5	3.05

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,665)	2:19:B:PHE:HE1	2:38:D:ILE:HD11	18	3.04
(1,665)	2:19:B:PHE:HE1	2:38:D:ILE:HD12	18	3.04
(1,665)	2:19:B:PHE:HE1	2:38:D:ILE:HD13	18	3.04
(1,665)	2:19:B:PHE:HE2	2:38:D:ILE:HD11	18	3.04
(1,665)	2:19:B:PHE:HE2	2:38:D:ILE:HD12	18	3.04
(1,665)	2:19:B:PHE:HE2	2:38:D:ILE:HD13	18	3.04
(1,18)	2:19:B:PHE:HE1	2:38:D:ILE:HD11	18	3.04
(1,18)	2:19:B:PHE:HE1	2:38:D:ILE:HD12	18	3.04
(1,18)	2:19:B:PHE:HE1	2:38:D:ILE:HD13	18	3.04
(1,18)	2:19:B:PHE:HE2	2:38:D:ILE:HD11	18	3.04
(1,18)	2:19:B:PHE:HE2	2:38:D:ILE:HD12	18	3.04
(1,18)	2:19:B:PHE:HE2	2:38:D:ILE:HD13	18	3.04
(1,665)	2:19:B:PHE:HE1	2:38:D:ILE:HD11	3	3.0
(1,665)	2:19:B:PHE:HE1	2:38:D:ILE:HD12	3	3.0
(1,665)	2:19:B:PHE:HE1	2:38:D:ILE:HD13	3	3.0
(1,665)	2:19:B:PHE:HE2	2:38:D:ILE:HD11	3	3.0
(1,665)	2:19:B:PHE:HE2	2:38:D:ILE:HD12	3	3.0
(1,665)	2:19:B:PHE:HE2	2:38:D:ILE:HD13	3	3.0
(1,18)	2:19:B:PHE:HE1	2:38:D:ILE:HD11	3	3.0
(1,18)	2:19:B:PHE:HE1	2:38:D:ILE:HD12	3	3.0
(1,18)	2:19:B:PHE:HE1	2:38:D:ILE:HD13	3	3.0
(1,18)	2:19:B:PHE:HE2	2:38:D:ILE:HD11	3	3.0
(1,18)	2:19:B:PHE:HE2	2:38:D:ILE:HD12	3	3.0
(1,18)	2:19:B:PHE:HE2	2:38:D:ILE:HD13	3	3.0
(1,1147)	1:28:C:PRO:HB2	1:31:C:PHE:HE1	9	2.97
(1,1147)	1:28:C:PRO:HB2	1:31:C:PHE:HE2	9	2.97
(1,828)	1:28:C:PRO:HB2	1:31:C:PHE:HE1	9	2.97
(1,828)	1:28:C:PRO:HB2	1:31:C:PHE:HE2	9	2.97
(1,665)	2:19:B:PHE:HE1	2:38:D:ILE:HD11	12	2.96
(1,665)	2:19:B:PHE:HE1	2:38:D:ILE:HD12	12	2.96
(1,665)	2:19:B:PHE:HE1	2:38:D:ILE:HD13	12	2.96
(1,665)	2:19:B:PHE:HE2	2:38:D:ILE:HD11	12	2.96
(1,665)	2:19:B:PHE:HE2	2:38:D:ILE:HD12	12	2.96
(1,665)	2:19:B:PHE:HE2	2:38:D:ILE:HD13	12	2.96
(1,18)	2:19:B:PHE:HE1	2:38:D:ILE:HD11	12	2.96
(1,18)	2:19:B:PHE:HE1	2:38:D:ILE:HD12	12	2.96
(1,18)	2:19:B:PHE:HE1	2:38:D:ILE:HD13	12	2.96
(1,18)	2:19:B:PHE:HE2	2:38:D:ILE:HD11	12	2.96
(1,18)	2:19:B:PHE:HE2	2:38:D:ILE:HD12	12	2.96
(1,18)	2:19:B:PHE:HE2	2:38:D:ILE:HD13	12	2.96
(1,665)	2:19:B:PHE:HE1	2:38:D:ILE:HD11	7	2.94
(1,665)	2:19:B:PHE:HE1	2:38:D:ILE:HD12	7	2.94

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,665)	2:19:B:PHE:HE1	2:38:D:ILE:HD13	7	2.94
(1,665)	2:19:B:PHE:HE2	2:38:D:ILE:HD11	7	2.94
(1,665)	2:19:B:PHE:HE2	2:38:D:ILE:HD12	7	2.94
(1,665)	2:19:B:PHE:HE2	2:38:D:ILE:HD13	7	2.94
(1,18)	2:19:B:PHE:HE1	2:38:D:ILE:HD11	7	2.94
(1,18)	2:19:B:PHE:HE1	2:38:D:ILE:HD12	7	2.94
(1,18)	2:19:B:PHE:HE1	2:38:D:ILE:HD13	7	2.94
(1,18)	2:19:B:PHE:HE2	2:38:D:ILE:HD11	7	2.94
(1,18)	2:19:B:PHE:HE2	2:38:D:ILE:HD12	7	2.94
(1,18)	2:19:B:PHE:HE2	2:38:D:ILE:HD13	7	2.94
(1,665)	2:19:B:PHE:HE1	2:38:D:ILE:HD11	13	2.93
(1,665)	2:19:B:PHE:HE1	2:38:D:ILE:HD12	13	2.93
(1,665)	2:19:B:PHE:HE1	2:38:D:ILE:HD13	13	2.93
(1,665)	2:19:B:PHE:HE2	2:38:D:ILE:HD11	13	2.93
(1,665)	2:19:B:PHE:HE2	2:38:D:ILE:HD12	13	2.93
(1,665)	2:19:B:PHE:HE2	2:38:D:ILE:HD13	13	2.93
(1,657)	2:19:B:PHE:HD1	2:38:D:ILE:HD11	9	2.93
(1,657)	2:19:B:PHE:HD1	2:38:D:ILE:HD12	9	2.93
(1,657)	2:19:B:PHE:HD1	2:38:D:ILE:HD13	9	2.93
(1,657)	2:19:B:PHE:HD2	2:38:D:ILE:HD11	9	2.93
(1,657)	2:19:B:PHE:HD2	2:38:D:ILE:HD12	9	2.93
(1,657)	2:19:B:PHE:HD2	2:38:D:ILE:HD13	9	2.93
(1,515)	2:15:B:THR:HG21	2:17:B:ILE:HD11	17	2.93
(1,515)	2:15:B:THR:HG21	2:17:B:ILE:HD12	17	2.93
(1,515)	2:15:B:THR:HG21	2:17:B:ILE:HD13	17	2.93
(1,515)	2:15:B:THR:HG22	2:17:B:ILE:HD11	17	2.93
(1,515)	2:15:B:THR:HG22	2:17:B:ILE:HD12	17	2.93
(1,515)	2:15:B:THR:HG22	2:17:B:ILE:HD13	17	2.93
(1,515)	2:15:B:THR:HG23	2:17:B:ILE:HD11	17	2.93
(1,515)	2:15:B:THR:HG23	2:17:B:ILE:HD12	17	2.93
(1,515)	2:15:B:THR:HG23	2:17:B:ILE:HD13	17	2.93
(1,196)	2:15:B:THR:HG21	2:17:B:ILE:HD11	17	2.93
(1,196)	2:15:B:THR:HG21	2:17:B:ILE:HD12	17	2.93
(1,196)	2:15:B:THR:HG21	2:17:B:ILE:HD13	17	2.93
(1,196)	2:15:B:THR:HG22	2:17:B:ILE:HD11	17	2.93
(1,196)	2:15:B:THR:HG22	2:17:B:ILE:HD12	17	2.93
(1,196)	2:15:B:THR:HG22	2:17:B:ILE:HD13	17	2.93
(1,196)	2:15:B:THR:HG23	2:17:B:ILE:HD11	17	2.93
(1,196)	2:15:B:THR:HG23	2:17:B:ILE:HD12	17	2.93
(1,196)	2:15:B:THR:HG23	2:17:B:ILE:HD13	17	2.93
(1,18)	2:19:B:PHE:HE1	2:38:D:ILE:HD11	13	2.93
(1,18)	2:19:B:PHE:HE1	2:38:D:ILE:HD12	13	2.93

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,18)	2:19:B:PHE:HE1	2:38:D:ILE:HD13	13	2.93
(1,18)	2:19:B:PHE:HE2	2:38:D:ILE:HD11	13	2.93
(1,18)	2:19:B:PHE:HE2	2:38:D:ILE:HD12	13	2.93
(1,18)	2:19:B:PHE:HE2	2:38:D:ILE:HD13	13	2.93
(1,10)	2:19:B:PHE:HD1	2:38:D:ILE:HD11	9	2.93
(1,10)	2:19:B:PHE:HD1	2:38:D:ILE:HD12	9	2.93
(1,10)	2:19:B:PHE:HD1	2:38:D:ILE:HD13	9	2.93
(1,10)	2:19:B:PHE:HD2	2:38:D:ILE:HD11	9	2.93
(1,10)	2:19:B:PHE:HD2	2:38:D:ILE:HD12	9	2.93
(1,10)	2:19:B:PHE:HD2	2:38:D:ILE:HD13	9	2.93
(1,665)	2:19:B:PHE:HE1	2:38:D:ILE:HD11	9	2.92
(1,665)	2:19:B:PHE:HE1	2:38:D:ILE:HD12	9	2.92
(1,665)	2:19:B:PHE:HE1	2:38:D:ILE:HD13	9	2.92
(1,665)	2:19:B:PHE:HE2	2:38:D:ILE:HD11	9	2.92
(1,665)	2:19:B:PHE:HE2	2:38:D:ILE:HD12	9	2.92
(1,665)	2:19:B:PHE:HE2	2:38:D:ILE:HD13	9	2.92
(1,18)	2:19:B:PHE:HE1	2:38:D:ILE:HD11	9	2.92
(1,18)	2:19:B:PHE:HE1	2:38:D:ILE:HD12	9	2.92
(1,18)	2:19:B:PHE:HE1	2:38:D:ILE:HD13	9	2.92
(1,18)	2:19:B:PHE:HE2	2:38:D:ILE:HD11	9	2.92
(1,18)	2:19:B:PHE:HE2	2:38:D:ILE:HD12	9	2.92
(1,18)	2:19:B:PHE:HE2	2:38:D:ILE:HD13	9	2.92
(1,1169)	2:36:D:THR:HB	2:38:D:ILE:HD11	5	2.9
(1,1169)	2:36:D:THR:HB	2:38:D:ILE:HD12	5	2.9
(1,1169)	2:36:D:THR:HB	2:38:D:ILE:HD13	5	2.9
(1,1147)	1:28:C:PRO:HB2	1:31:C:PHE:HE1	2	2.9
(1,1147)	1:28:C:PRO:HB2	1:31:C:PHE:HE2	2	2.9
(1,850)	2:36:D:THR:HB	2:38:D:ILE:HD11	5	2.9
(1,850)	2:36:D:THR:HB	2:38:D:ILE:HD12	5	2.9
(1,850)	2:36:D:THR:HB	2:38:D:ILE:HD13	5	2.9
(1,828)	1:28:C:PRO:HB2	1:31:C:PHE:HE1	2	2.9
(1,828)	1:28:C:PRO:HB2	1:31:C:PHE:HE2	2	2.9
(1,665)	2:19:B:PHE:HE1	2:38:D:ILE:HD11	8	2.88
(1,665)	2:19:B:PHE:HE1	2:38:D:ILE:HD12	8	2.88
(1,665)	2:19:B:PHE:HE1	2:38:D:ILE:HD13	8	2.88
(1,665)	2:19:B:PHE:HE2	2:38:D:ILE:HD11	8	2.88
(1,665)	2:19:B:PHE:HE2	2:38:D:ILE:HD12	8	2.88
(1,665)	2:19:B:PHE:HE2	2:38:D:ILE:HD13	8	2.88
(1,18)	2:19:B:PHE:HE1	2:38:D:ILE:HD11	8	2.88
(1,18)	2:19:B:PHE:HE1	2:38:D:ILE:HD12	8	2.88
(1,18)	2:19:B:PHE:HE1	2:38:D:ILE:HD13	8	2.88
(1,18)	2:19:B:PHE:HE2	2:38:D:ILE:HD11	8	2.88

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,18)	2:19:B:PHE:HE2	2:38:D:ILE:HD12	8	2.88
(1,18)	2:19:B:PHE:HE2	2:38:D:ILE:HD13	8	2.88
(1,402)	1:10:A:PHE:HD1	2:14:B:GLY:HA3	1	2.87
(1,402)	1:10:A:PHE:HD2	2:14:B:GLY:HA3	1	2.87
(1,83)	1:10:A:PHE:HD1	2:14:B:GLY:HA3	1	2.87
(1,83)	1:10:A:PHE:HD2	2:14:B:GLY:HA3	1	2.87
(1,674)	2:17:B:ILE:HD11	2:40:D:PHE:HE1	10	2.86
(1,674)	2:17:B:ILE:HD11	2:40:D:PHE:HE2	10	2.86
(1,674)	2:17:B:ILE:HD12	2:40:D:PHE:HE1	10	2.86
(1,674)	2:17:B:ILE:HD12	2:40:D:PHE:HE2	10	2.86
(1,674)	2:17:B:ILE:HD13	2:40:D:PHE:HE1	10	2.86
(1,674)	2:17:B:ILE:HD13	2:40:D:PHE:HE2	10	2.86
(1,9)	2:17:B:ILE:HD11	2:40:D:PHE:HE1	10	2.86
(1,9)	2:17:B:ILE:HD11	2:40:D:PHE:HE2	10	2.86
(1,9)	2:17:B:ILE:HD12	2:40:D:PHE:HE1	10	2.86
(1,9)	2:17:B:ILE:HD12	2:40:D:PHE:HE2	10	2.86
(1,9)	2:17:B:ILE:HD13	2:40:D:PHE:HE1	10	2.86
(1,9)	2:17:B:ILE:HD13	2:40:D:PHE:HE2	10	2.86
(1,663)	2:19:B:PHE:HD1	2:38:D:ILE:HG13	5	2.85
(1,663)	2:19:B:PHE:HD2	2:38:D:ILE:HG13	5	2.85
(1,16)	2:19:B:PHE:HD1	2:38:D:ILE:HG13	5	2.85
(1,16)	2:19:B:PHE:HD2	2:38:D:ILE:HG13	5	2.85
(1,1171)	2:36:D:THR:HG21	2:38:D:ILE:HD11	17	2.84
(1,1171)	2:36:D:THR:HG21	2:38:D:ILE:HD12	17	2.84
(1,1171)	2:36:D:THR:HG21	2:38:D:ILE:HD13	17	2.84
(1,1171)	2:36:D:THR:HG22	2:38:D:ILE:HD11	17	2.84
(1,1171)	2:36:D:THR:HG22	2:38:D:ILE:HD12	17	2.84
(1,1171)	2:36:D:THR:HG22	2:38:D:ILE:HD13	17	2.84
(1,1171)	2:36:D:THR:HG23	2:38:D:ILE:HD11	17	2.84
(1,1171)	2:36:D:THR:HG23	2:38:D:ILE:HD12	17	2.84
(1,1171)	2:36:D:THR:HG23	2:38:D:ILE:HD13	17	2.84
(1,852)	2:36:D:THR:HG21	2:38:D:ILE:HD11	17	2.84
(1,852)	2:36:D:THR:HG21	2:38:D:ILE:HD12	17	2.84
(1,852)	2:36:D:THR:HG21	2:38:D:ILE:HD13	17	2.84
(1,852)	2:36:D:THR:HG22	2:38:D:ILE:HD11	17	2.84
(1,852)	2:36:D:THR:HG22	2:38:D:ILE:HD12	17	2.84
(1,852)	2:36:D:THR:HG22	2:38:D:ILE:HD13	17	2.84
(1,852)	2:36:D:THR:HG23	2:38:D:ILE:HD11	17	2.84
(1,852)	2:36:D:THR:HG23	2:38:D:ILE:HD12	17	2.84
(1,852)	2:36:D:THR:HG23	2:38:D:ILE:HD13	17	2.84
(1,1176)	2:19:B:PHE:HB3	2:38:D:ILE:HG13	3	2.83
(1,857)	2:19:B:PHE:HB3	2:38:D:ILE:HG13	3	2.83

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,664)	2:19:B:PHE:HB2	2:38:D:ILE:HG13	15	2.81
(1,17)	2:19:B:PHE:HB2	2:38:D:ILE:HG13	15	2.81
(1,1058)	1:31:C:PHE:HD1	2:35:D:GLY:HA3	5	2.8
(1,1058)	1:31:C:PHE:HD2	2:35:D:GLY:HA3	5	2.8
(1,739)	1:31:C:PHE:HD1	2:35:D:GLY:HA3	5	2.8
(1,739)	1:31:C:PHE:HD2	2:35:D:GLY:HA3	5	2.8
(1,503)	1:7:A:PRO:HB3	1:10:A:PHE:HE1	12	2.77
(1,503)	1:7:A:PRO:HB3	1:10:A:PHE:HE2	12	2.77
(1,184)	1:7:A:PRO:HB3	1:10:A:PHE:HE1	12	2.77
(1,184)	1:7:A:PRO:HB3	1:10:A:PHE:HE2	12	2.77
(1,674)	2:17:B:ILE:HD11	2:40:D:PHE:HE1	17	2.76
(1,674)	2:17:B:ILE:HD11	2:40:D:PHE:HE2	17	2.76
(1,674)	2:17:B:ILE:HD12	2:40:D:PHE:HE1	17	2.76
(1,674)	2:17:B:ILE:HD12	2:40:D:PHE:HE2	17	2.76
(1,674)	2:17:B:ILE:HD13	2:40:D:PHE:HE1	17	2.76
(1,674)	2:17:B:ILE:HD13	2:40:D:PHE:HE2	17	2.76
(1,9)	2:17:B:ILE:HD11	2:40:D:PHE:HE1	17	2.76
(1,9)	2:17:B:ILE:HD11	2:40:D:PHE:HE2	17	2.76
(1,9)	2:17:B:ILE:HD12	2:40:D:PHE:HE1	17	2.76
(1,9)	2:17:B:ILE:HD12	2:40:D:PHE:HE2	17	2.76
(1,9)	2:17:B:ILE:HD13	2:40:D:PHE:HE1	17	2.76
(1,9)	2:17:B:ILE:HD13	2:40:D:PHE:HE2	17	2.76
(1,1008)	2:18:B:SER:HA	2:38:D:ILE:HG13	5	2.74
(1,689)	2:18:B:SER:HA	2:38:D:ILE:HG13	5	2.74
(1,402)	1:10:A:PHE:HD1	2:14:B:GLY:HA3	17	2.74
(1,402)	1:10:A:PHE:HD2	2:14:B:GLY:HA3	17	2.74
(1,83)	1:10:A:PHE:HD1	2:14:B:GLY:HA3	17	2.74
(1,83)	1:10:A:PHE:HD2	2:14:B:GLY:HA3	17	2.74
(1,515)	2:15:B:THR:HG21	2:17:B:ILE:HD11	3	2.73
(1,515)	2:15:B:THR:HG21	2:17:B:ILE:HD12	3	2.73
(1,515)	2:15:B:THR:HG21	2:17:B:ILE:HD13	3	2.73
(1,515)	2:15:B:THR:HG22	2:17:B:ILE:HD11	3	2.73
(1,515)	2:15:B:THR:HG22	2:17:B:ILE:HD12	3	2.73
(1,515)	2:15:B:THR:HG22	2:17:B:ILE:HD13	3	2.73
(1,515)	2:15:B:THR:HG23	2:17:B:ILE:HD11	3	2.73
(1,515)	2:15:B:THR:HG23	2:17:B:ILE:HD12	3	2.73
(1,515)	2:15:B:THR:HG23	2:17:B:ILE:HD13	3	2.73
(1,503)	1:7:A:PRO:HB3	1:10:A:PHE:HE1	13	2.73
(1,503)	1:7:A:PRO:HB3	1:10:A:PHE:HE2	13	2.73
(1,196)	2:15:B:THR:HG21	2:17:B:ILE:HD11	3	2.73
(1,196)	2:15:B:THR:HG21	2:17:B:ILE:HD12	3	2.73
(1,196)	2:15:B:THR:HG21	2:17:B:ILE:HD13	3	2.73

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,196)	2:15:B:THR:HG22	2:17:B:ILE:HD11	3	2.73
(1,196)	2:15:B:THR:HG22	2:17:B:ILE:HD12	3	2.73
(1,196)	2:15:B:THR:HG22	2:17:B:ILE:HD13	3	2.73
(1,196)	2:15:B:THR:HG23	2:17:B:ILE:HD11	3	2.73
(1,196)	2:15:B:THR:HG23	2:17:B:ILE:HD12	3	2.73
(1,196)	2:15:B:THR:HG23	2:17:B:ILE:HD13	3	2.73
(1,184)	1:7:A:PRO:HB3	1:10:A:PHE:HE1	13	2.73
(1,184)	1:7:A:PRO:HB3	1:10:A:PHE:HE2	13	2.73
(1,662)	2:19:B:PHE:HE1	2:38:D:ILE:HG13	14	2.7
(1,662)	2:19:B:PHE:HE2	2:38:D:ILE:HG13	14	2.7
(1,15)	2:19:B:PHE:HE1	2:38:D:ILE:HG13	14	2.7
(1,15)	2:19:B:PHE:HE2	2:38:D:ILE:HG13	14	2.7
(1,661)	2:19:B:PHE:H	2:38:D:ILE:HD11	14	2.69
(1,661)	2:19:B:PHE:H	2:38:D:ILE:HD12	14	2.69
(1,661)	2:19:B:PHE:H	2:38:D:ILE:HD13	14	2.69
(1,14)	2:19:B:PHE:H	2:38:D:ILE:HD11	14	2.69
(1,14)	2:19:B:PHE:H	2:38:D:ILE:HD12	14	2.69
(1,14)	2:19:B:PHE:H	2:38:D:ILE:HD13	14	2.69
(1,666)	2:17:B:ILE:HD11	2:40:D:PHE:HD1	7	2.68
(1,666)	2:17:B:ILE:HD11	2:40:D:PHE:HD2	7	2.68
(1,666)	2:17:B:ILE:HD12	2:40:D:PHE:HD1	7	2.68
(1,666)	2:17:B:ILE:HD12	2:40:D:PHE:HD2	7	2.68
(1,666)	2:17:B:ILE:HD13	2:40:D:PHE:HD1	7	2.68
(1,666)	2:17:B:ILE:HD13	2:40:D:PHE:HD2	7	2.68
(1,1)	2:17:B:ILE:HD11	2:40:D:PHE:HD1	7	2.68
(1,1)	2:17:B:ILE:HD11	2:40:D:PHE:HD2	7	2.68
(1,1)	2:17:B:ILE:HD12	2:40:D:PHE:HD1	7	2.68
(1,1)	2:17:B:ILE:HD12	2:40:D:PHE:HD2	7	2.68
(1,1)	2:17:B:ILE:HD13	2:40:D:PHE:HD1	7	2.68
(1,1)	2:17:B:ILE:HD13	2:40:D:PHE:HD2	7	2.68
(1,1071)	1:27:C:VAL:HG11	2:41:D:TYR:HE1	18	2.67
(1,1071)	1:27:C:VAL:HG11	2:41:D:TYR:HE2	18	2.67
(1,1071)	1:27:C:VAL:HG12	2:41:D:TYR:HE1	18	2.67
(1,1071)	1:27:C:VAL:HG12	2:41:D:TYR:HE2	18	2.67
(1,1071)	1:27:C:VAL:HG13	2:41:D:TYR:HE1	18	2.67
(1,1071)	1:27:C:VAL:HG13	2:41:D:TYR:HE2	18	2.67
(1,1058)	1:31:C:PHE:HD1	2:35:D:GLY:HA3	1	2.67
(1,1058)	1:31:C:PHE:HD2	2:35:D:GLY:HA3	1	2.67
(1,752)	1:27:C:VAL:HG11	2:41:D:TYR:HE1	18	2.67
(1,752)	1:27:C:VAL:HG11	2:41:D:TYR:HE2	18	2.67
(1,752)	1:27:C:VAL:HG12	2:41:D:TYR:HE1	18	2.67
(1,752)	1:27:C:VAL:HG12	2:41:D:TYR:HE2	18	2.67

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,752)	1:27:C:VAL:HG13	2:41:D:TYR:HE1	18	2.67
(1,752)	1:27:C:VAL:HG13	2:41:D:TYR:HE2	18	2.67
(1,739)	1:31:C:PHE:HD1	2:35:D:GLY:HA3	1	2.67
(1,739)	1:31:C:PHE:HD2	2:35:D:GLY:HA3	1	2.67
(1,661)	2:19:B:PHE:H	2:38:D:ILE:HD11	19	2.67
(1,661)	2:19:B:PHE:H	2:38:D:ILE:HD12	19	2.67
(1,661)	2:19:B:PHE:H	2:38:D:ILE:HD13	19	2.67
(1,14)	2:19:B:PHE:H	2:38:D:ILE:HD11	19	2.67
(1,14)	2:19:B:PHE:H	2:38:D:ILE:HD12	19	2.67
(1,14)	2:19:B:PHE:H	2:38:D:ILE:HD13	19	2.67
(1,1176)	2:19:B:PHE:HB3	2:38:D:ILE:HG13	13	2.66
(1,857)	2:19:B:PHE:HB3	2:38:D:ILE:HG13	13	2.66
(1,674)	2:17:B:ILE:HD11	2:40:D:PHE:HE1	13	2.65
(1,674)	2:17:B:ILE:HD11	2:40:D:PHE:HE2	13	2.65
(1,674)	2:17:B:ILE:HD12	2:40:D:PHE:HE1	13	2.65
(1,674)	2:17:B:ILE:HD12	2:40:D:PHE:HE2	13	2.65
(1,674)	2:17:B:ILE:HD13	2:40:D:PHE:HE1	13	2.65
(1,674)	2:17:B:ILE:HD13	2:40:D:PHE:HE2	13	2.65
(1,662)	2:19:B:PHE:HE1	2:38:D:ILE:HG13	12	2.65
(1,662)	2:19:B:PHE:HE2	2:38:D:ILE:HG13	12	2.65
(1,657)	2:19:B:PHE:HD1	2:38:D:ILE:HD11	3	2.65
(1,657)	2:19:B:PHE:HD1	2:38:D:ILE:HD12	3	2.65
(1,657)	2:19:B:PHE:HD1	2:38:D:ILE:HD13	3	2.65
(1,657)	2:19:B:PHE:HD2	2:38:D:ILE:HD11	3	2.65
(1,657)	2:19:B:PHE:HD2	2:38:D:ILE:HD12	3	2.65
(1,657)	2:19:B:PHE:HD2	2:38:D:ILE:HD13	3	2.65
(1,491)	1:7:A:PRO:HB2	1:10:A:PHE:HE1	12	2.65
(1,491)	1:7:A:PRO:HB2	1:10:A:PHE:HE2	12	2.65
(1,172)	1:7:A:PRO:HB2	1:10:A:PHE:HE1	12	2.65
(1,172)	1:7:A:PRO:HB2	1:10:A:PHE:HE2	12	2.65
(1,15)	2:19:B:PHE:HE1	2:38:D:ILE:HG13	12	2.65
(1,15)	2:19:B:PHE:HE2	2:38:D:ILE:HG13	12	2.65
(1,10)	2:19:B:PHE:HD1	2:38:D:ILE:HD11	3	2.65
(1,10)	2:19:B:PHE:HD1	2:38:D:ILE:HD12	3	2.65
(1,10)	2:19:B:PHE:HD1	2:38:D:ILE:HD13	3	2.65
(1,10)	2:19:B:PHE:HD2	2:38:D:ILE:HD11	3	2.65
(1,10)	2:19:B:PHE:HD2	2:38:D:ILE:HD12	3	2.65
(1,10)	2:19:B:PHE:HD2	2:38:D:ILE:HD13	3	2.65
(1,9)	2:17:B:ILE:HD11	2:40:D:PHE:HE1	13	2.65
(1,9)	2:17:B:ILE:HD11	2:40:D:PHE:HE2	13	2.65
(1,9)	2:17:B:ILE:HD12	2:40:D:PHE:HE1	13	2.65
(1,9)	2:17:B:ILE:HD12	2:40:D:PHE:HE2	13	2.65

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,9)	2:17:B:ILE:HD13	2:40:D:PHE:HE1	13	2.65
(1,9)	2:17:B:ILE:HD13	2:40:D:PHE:HE2	13	2.65
(1,399)	1:7:A:PRO:HG2	1:10:A:PHE:HE1	16	2.64
(1,399)	1:7:A:PRO:HG2	1:10:A:PHE:HE2	16	2.64
(1,399)	1:7:A:PRO:HG3	1:10:A:PHE:HE1	16	2.64
(1,399)	1:7:A:PRO:HG3	1:10:A:PHE:HE2	16	2.64
(1,80)	1:7:A:PRO:HG2	1:10:A:PHE:HE1	16	2.64
(1,80)	1:7:A:PRO:HG2	1:10:A:PHE:HE2	16	2.64
(1,80)	1:7:A:PRO:HG3	1:10:A:PHE:HE1	16	2.64
(1,80)	1:7:A:PRO:HG3	1:10:A:PHE:HE2	16	2.64
(1,1159)	1:28:C:PRO:HB3	1:31:C:PHE:HE1	3	2.63
(1,1159)	1:28:C:PRO:HB3	1:31:C:PHE:HE2	3	2.63
(1,840)	1:28:C:PRO:HB3	1:31:C:PHE:HE1	3	2.63
(1,840)	1:28:C:PRO:HB3	1:31:C:PHE:HE2	3	2.63
(1,1017)	1:31:C:PHE:HE1	2:35:D:GLY:HA3	5	2.62
(1,1017)	1:31:C:PHE:HE2	2:35:D:GLY:HA3	5	2.62
(1,698)	1:31:C:PHE:HE1	2:35:D:GLY:HA3	5	2.62
(1,698)	1:31:C:PHE:HE2	2:35:D:GLY:HA3	5	2.62
(1,657)	2:19:B:PHE:HD1	2:38:D:ILE:HD11	17	2.62
(1,657)	2:19:B:PHE:HD1	2:38:D:ILE:HD12	17	2.62
(1,657)	2:19:B:PHE:HD1	2:38:D:ILE:HD13	17	2.62
(1,657)	2:19:B:PHE:HD2	2:38:D:ILE:HD11	17	2.62
(1,657)	2:19:B:PHE:HD2	2:38:D:ILE:HD12	17	2.62
(1,657)	2:19:B:PHE:HD2	2:38:D:ILE:HD13	17	2.62
(1,595)	1:12:A:ARG:HA	1:12:A:ARG:HE	12	2.62
(1,276)	1:12:A:ARG:HA	1:12:A:ARG:HE	12	2.62
(1,10)	2:19:B:PHE:HD1	2:38:D:ILE:HD11	17	2.62
(1,10)	2:19:B:PHE:HD1	2:38:D:ILE:HD12	17	2.62
(1,10)	2:19:B:PHE:HD1	2:38:D:ILE:HD13	17	2.62
(1,10)	2:19:B:PHE:HD2	2:38:D:ILE:HD11	17	2.62
(1,10)	2:19:B:PHE:HD2	2:38:D:ILE:HD12	17	2.62
(1,10)	2:19:B:PHE:HD2	2:38:D:ILE:HD13	17	2.62
(1,417)	2:17:B:ILE:HD11	2:39:D:SER:HB3	2	2.61
(1,417)	2:17:B:ILE:HD12	2:39:D:SER:HB3	2	2.61
(1,417)	2:17:B:ILE:HD13	2:39:D:SER:HB3	2	2.61
(1,98)	2:17:B:ILE:HD11	2:39:D:SER:HB3	2	2.61
(1,98)	2:17:B:ILE:HD12	2:39:D:SER:HB3	2	2.61
(1,98)	2:17:B:ILE:HD13	2:39:D:SER:HB3	2	2.61
(1,1147)	1:28:C:PRO:HB2	1:31:C:PHE:HE1	3	2.6
(1,1147)	1:28:C:PRO:HB2	1:31:C:PHE:HE2	3	2.6
(1,828)	1:28:C:PRO:HB2	1:31:C:PHE:HE1	3	2.6
(1,828)	1:28:C:PRO:HB2	1:31:C:PHE:HE2	3	2.6

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,1061)	1:22:C:GLY:H	2:39:D:SER:HB2	1	2.59
(1,742)	1:22:C:GLY:H	2:39:D:SER:HB2	1	2.59
(1,596)	1:7:A:PRO:HG2	1:10:A:PHE:HE1	16	2.59
(1,596)	1:7:A:PRO:HG2	1:10:A:PHE:HE2	16	2.59
(1,596)	1:7:A:PRO:HG3	1:10:A:PHE:HE1	16	2.59
(1,596)	1:7:A:PRO:HG3	1:10:A:PHE:HE2	16	2.59
(1,520)	2:17:B:ILE:HG13	2:40:D:PHE:HB3	10	2.59
(1,277)	1:7:A:PRO:HG2	1:10:A:PHE:HE1	16	2.59
(1,277)	1:7:A:PRO:HG2	1:10:A:PHE:HE2	16	2.59
(1,277)	1:7:A:PRO:HG3	1:10:A:PHE:HE1	16	2.59
(1,277)	1:7:A:PRO:HG3	1:10:A:PHE:HE2	16	2.59
(1,201)	2:17:B:ILE:HG13	2:40:D:PHE:HB3	10	2.59
(1,436)	1:8:A:GLU:HG2	2:20:B:TYR:HD1	13	2.58
(1,436)	1:8:A:GLU:HG2	2:20:B:TYR:HD2	13	2.58
(1,117)	1:8:A:GLU:HG2	2:20:B:TYR:HD1	13	2.58
(1,117)	1:8:A:GLU:HG2	2:20:B:TYR:HD2	13	2.58
(1,1171)	2:36:D:THR:HG21	2:38:D:ILE:HD11	9	2.57
(1,1171)	2:36:D:THR:HG21	2:38:D:ILE:HD12	9	2.57
(1,1171)	2:36:D:THR:HG21	2:38:D:ILE:HD13	9	2.57
(1,1171)	2:36:D:THR:HG22	2:38:D:ILE:HD11	9	2.57
(1,1171)	2:36:D:THR:HG22	2:38:D:ILE:HD12	9	2.57
(1,1171)	2:36:D:THR:HG22	2:38:D:ILE:HD13	9	2.57
(1,1171)	2:36:D:THR:HG23	2:38:D:ILE:HD11	9	2.57
(1,1171)	2:36:D:THR:HG23	2:38:D:ILE:HD12	9	2.57
(1,1171)	2:36:D:THR:HG23	2:38:D:ILE:HD13	9	2.57
(1,1171)	2:36:D:THR:HG21	2:38:D:ILE:HD11	15	2.57
(1,1171)	2:36:D:THR:HG21	2:38:D:ILE:HD12	15	2.57
(1,1171)	2:36:D:THR:HG21	2:38:D:ILE:HD13	15	2.57
(1,1171)	2:36:D:THR:HG22	2:38:D:ILE:HD11	15	2.57
(1,1171)	2:36:D:THR:HG22	2:38:D:ILE:HD12	15	2.57
(1,1171)	2:36:D:THR:HG22	2:38:D:ILE:HD13	15	2.57
(1,1171)	2:36:D:THR:HG23	2:38:D:ILE:HD11	15	2.57
(1,1171)	2:36:D:THR:HG23	2:38:D:ILE:HD12	15	2.57
(1,1171)	2:36:D:THR:HG23	2:38:D:ILE:HD13	15	2.57
(1,852)	2:36:D:THR:HG21	2:38:D:ILE:HD11	9	2.57
(1,852)	2:36:D:THR:HG21	2:38:D:ILE:HD12	9	2.57
(1,852)	2:36:D:THR:HG21	2:38:D:ILE:HD13	9	2.57
(1,852)	2:36:D:THR:HG22	2:38:D:ILE:HD11	9	2.57
(1,852)	2:36:D:THR:HG22	2:38:D:ILE:HD12	9	2.57
(1,852)	2:36:D:THR:HG22	2:38:D:ILE:HD13	9	2.57
(1,852)	2:36:D:THR:HG23	2:38:D:ILE:HD11	9	2.57
(1,852)	2:36:D:THR:HG23	2:38:D:ILE:HD12	9	2.57

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,852)	2:36:D:THR:HG23	2:38:D:ILE:HD13	9	2.57
(1,852)	2:36:D:THR:HG21	2:38:D:ILE:HD11	15	2.57
(1,852)	2:36:D:THR:HG21	2:38:D:ILE:HD12	15	2.57
(1,852)	2:36:D:THR:HG21	2:38:D:ILE:HD13	15	2.57
(1,852)	2:36:D:THR:HG22	2:38:D:ILE:HD11	15	2.57
(1,852)	2:36:D:THR:HG22	2:38:D:ILE:HD12	15	2.57
(1,852)	2:36:D:THR:HG22	2:38:D:ILE:HD13	15	2.57
(1,852)	2:36:D:THR:HG23	2:38:D:ILE:HD11	15	2.57
(1,852)	2:36:D:THR:HG23	2:38:D:ILE:HD12	15	2.57
(1,852)	2:36:D:THR:HG23	2:38:D:ILE:HD13	15	2.57
(1,595)	1:12:A:ARG:HA	1:12:A:ARG:HE	5	2.57
(1,276)	1:12:A:ARG:HA	1:12:A:ARG:HE	5	2.57
(1,1251)	1:33:C:ARG:HA	1:33:C:ARG:HE	12	2.56
(1,932)	1:33:C:ARG:HA	1:33:C:ARG:HE	12	2.56
(1,671)	2:17:B:ILE:HG13	2:40:D:PHE:HE1	7	2.55
(1,671)	2:17:B:ILE:HG13	2:40:D:PHE:HE2	7	2.55
(1,515)	2:15:B:THR:HG21	2:17:B:ILE:HD11	16	2.55
(1,515)	2:15:B:THR:HG21	2:17:B:ILE:HD12	16	2.55
(1,515)	2:15:B:THR:HG21	2:17:B:ILE:HD13	16	2.55
(1,515)	2:15:B:THR:HG22	2:17:B:ILE:HD11	16	2.55
(1,515)	2:15:B:THR:HG22	2:17:B:ILE:HD12	16	2.55
(1,515)	2:15:B:THR:HG22	2:17:B:ILE:HD13	16	2.55
(1,515)	2:15:B:THR:HG23	2:17:B:ILE:HD11	16	2.55
(1,515)	2:15:B:THR:HG23	2:17:B:ILE:HD12	16	2.55
(1,515)	2:15:B:THR:HG23	2:17:B:ILE:HD13	16	2.55
(1,491)	1:7:A:PRO:HB2	1:10:A:PHE:HE1	13	2.55
(1,491)	1:7:A:PRO:HB2	1:10:A:PHE:HE2	13	2.55
(1,196)	2:15:B:THR:HG21	2:17:B:ILE:HD11	16	2.55
(1,196)	2:15:B:THR:HG21	2:17:B:ILE:HD12	16	2.55
(1,196)	2:15:B:THR:HG21	2:17:B:ILE:HD13	16	2.55
(1,196)	2:15:B:THR:HG22	2:17:B:ILE:HD11	16	2.55
(1,196)	2:15:B:THR:HG22	2:17:B:ILE:HD12	16	2.55
(1,196)	2:15:B:THR:HG22	2:17:B:ILE:HD13	16	2.55
(1,196)	2:15:B:THR:HG23	2:17:B:ILE:HD11	16	2.55
(1,196)	2:15:B:THR:HG23	2:17:B:ILE:HD12	16	2.55
(1,196)	2:15:B:THR:HG23	2:17:B:ILE:HD13	16	2.55
(1,172)	1:7:A:PRO:HB2	1:10:A:PHE:HE1	13	2.55
(1,172)	1:7:A:PRO:HB2	1:10:A:PHE:HE2	13	2.55
(1,6)	2:17:B:ILE:HG13	2:40:D:PHE:HE1	7	2.55
(1,6)	2:17:B:ILE:HG13	2:40:D:PHE:HE2	7	2.55
(1,661)	2:19:B:PHE:H	2:38:D:ILE:HD11	3	2.54
(1,661)	2:19:B:PHE:H	2:38:D:ILE:HD12	3	2.54

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,661)	2:19:B:PHE:H	2:38:D:ILE:HD13	3	2.54
(1,657)	2:19:B:PHE:HD1	2:38:D:ILE:HD11	13	2.54
(1,657)	2:19:B:PHE:HD1	2:38:D:ILE:HD12	13	2.54
(1,657)	2:19:B:PHE:HD1	2:38:D:ILE:HD13	13	2.54
(1,657)	2:19:B:PHE:HD2	2:38:D:ILE:HD11	13	2.54
(1,657)	2:19:B:PHE:HD2	2:38:D:ILE:HD12	13	2.54
(1,657)	2:19:B:PHE:HD2	2:38:D:ILE:HD13	13	2.54
(1,14)	2:19:B:PHE:H	2:38:D:ILE:HD11	3	2.54
(1,14)	2:19:B:PHE:H	2:38:D:ILE:HD12	3	2.54
(1,14)	2:19:B:PHE:H	2:38:D:ILE:HD13	3	2.54
(1,10)	2:19:B:PHE:HD1	2:38:D:ILE:HD11	13	2.54
(1,10)	2:19:B:PHE:HD1	2:38:D:ILE:HD12	13	2.54
(1,10)	2:19:B:PHE:HD1	2:38:D:ILE:HD13	13	2.54
(1,10)	2:19:B:PHE:HD2	2:38:D:ILE:HD11	13	2.54
(1,10)	2:19:B:PHE:HD2	2:38:D:ILE:HD12	13	2.54
(1,10)	2:19:B:PHE:HD2	2:38:D:ILE:HD13	13	2.54
(1,674)	2:17:B:ILE:HD11	2:40:D:PHE:HE1	18	2.53
(1,674)	2:17:B:ILE:HD11	2:40:D:PHE:HE2	18	2.53
(1,674)	2:17:B:ILE:HD12	2:40:D:PHE:HE1	18	2.53
(1,674)	2:17:B:ILE:HD12	2:40:D:PHE:HE2	18	2.53
(1,674)	2:17:B:ILE:HD13	2:40:D:PHE:HE1	18	2.53
(1,674)	2:17:B:ILE:HD13	2:40:D:PHE:HE2	18	2.53
(1,9)	2:17:B:ILE:HD11	2:40:D:PHE:HE1	18	2.53
(1,9)	2:17:B:ILE:HD11	2:40:D:PHE:HE2	18	2.53
(1,9)	2:17:B:ILE:HD12	2:40:D:PHE:HE1	18	2.53
(1,9)	2:17:B:ILE:HD12	2:40:D:PHE:HE2	18	2.53
(1,9)	2:17:B:ILE:HD13	2:40:D:PHE:HE1	18	2.53
(1,9)	2:17:B:ILE:HD13	2:40:D:PHE:HE2	18	2.53
(1,1073)	2:18:B:SER:HB3	2:38:D:ILE:HD11	3	2.52
(1,1073)	2:18:B:SER:HB3	2:38:D:ILE:HD12	3	2.52
(1,1073)	2:18:B:SER:HB3	2:38:D:ILE:HD13	3	2.52
(1,754)	2:18:B:SER:HB3	2:38:D:ILE:HD11	3	2.52
(1,754)	2:18:B:SER:HB3	2:38:D:ILE:HD12	3	2.52
(1,754)	2:18:B:SER:HB3	2:38:D:ILE:HD13	3	2.52
(1,674)	2:17:B:ILE:HD11	2:40:D:PHE:HE1	4	2.52
(1,674)	2:17:B:ILE:HD11	2:40:D:PHE:HE2	4	2.52
(1,674)	2:17:B:ILE:HD12	2:40:D:PHE:HE1	4	2.52
(1,674)	2:17:B:ILE:HD12	2:40:D:PHE:HE2	4	2.52
(1,674)	2:17:B:ILE:HD13	2:40:D:PHE:HE1	4	2.52
(1,674)	2:17:B:ILE:HD13	2:40:D:PHE:HE2	4	2.52
(1,674)	2:17:B:ILE:HD11	2:40:D:PHE:HE1	12	2.52
(1,674)	2:17:B:ILE:HD11	2:40:D:PHE:HE2	12	2.52

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,674)	2:17:B:ILE:HD12	2:40:D:PHE:HE1	12	2.52
(1,674)	2:17:B:ILE:HD12	2:40:D:PHE:HE2	12	2.52
(1,674)	2:17:B:ILE:HD13	2:40:D:PHE:HE1	12	2.52
(1,674)	2:17:B:ILE:HD13	2:40:D:PHE:HE2	12	2.52
(1,664)	2:19:B:PHE:HB2	2:38:D:ILE:HG13	3	2.52
(1,520)	2:17:B:ILE:HG13	2:40:D:PHE:HB3	17	2.52
(1,201)	2:17:B:ILE:HG13	2:40:D:PHE:HB3	17	2.52
(1,17)	2:19:B:PHE:HB2	2:38:D:ILE:HG13	3	2.52
(1,9)	2:17:B:ILE:HD11	2:40:D:PHE:HE1	4	2.52
(1,9)	2:17:B:ILE:HD11	2:40:D:PHE:HE2	4	2.52
(1,9)	2:17:B:ILE:HD12	2:40:D:PHE:HE1	4	2.52
(1,9)	2:17:B:ILE:HD12	2:40:D:PHE:HE2	4	2.52
(1,9)	2:17:B:ILE:HD13	2:40:D:PHE:HE1	4	2.52
(1,9)	2:17:B:ILE:HD13	2:40:D:PHE:HE2	4	2.52
(1,9)	2:17:B:ILE:HD11	2:40:D:PHE:HE1	12	2.52
(1,9)	2:17:B:ILE:HD11	2:40:D:PHE:HE2	12	2.52
(1,9)	2:17:B:ILE:HD12	2:40:D:PHE:HE1	12	2.52
(1,9)	2:17:B:ILE:HD12	2:40:D:PHE:HE2	12	2.52
(1,9)	2:17:B:ILE:HD13	2:40:D:PHE:HE1	12	2.52
(1,9)	2:17:B:ILE:HD13	2:40:D:PHE:HE2	12	2.52
(1,674)	2:17:B:ILE:HD11	2:40:D:PHE:HE1	9	2.51
(1,674)	2:17:B:ILE:HD11	2:40:D:PHE:HE2	9	2.51
(1,674)	2:17:B:ILE:HD12	2:40:D:PHE:HE1	9	2.51
(1,674)	2:17:B:ILE:HD12	2:40:D:PHE:HE2	9	2.51
(1,674)	2:17:B:ILE:HD13	2:40:D:PHE:HE1	9	2.51
(1,674)	2:17:B:ILE:HD13	2:40:D:PHE:HE2	9	2.51
(1,9)	2:17:B:ILE:HD11	2:40:D:PHE:HE1	9	2.51
(1,9)	2:17:B:ILE:HD11	2:40:D:PHE:HE2	9	2.51
(1,9)	2:17:B:ILE:HD12	2:40:D:PHE:HE1	9	2.51
(1,9)	2:17:B:ILE:HD12	2:40:D:PHE:HE2	9	2.51
(1,9)	2:17:B:ILE:HD13	2:40:D:PHE:HE1	9	2.51
(1,9)	2:17:B:ILE:HD13	2:40:D:PHE:HE2	9	2.51
(1,1000)	1:31:C:PHE:HE1	2:37:D:PRO:HD2	20	2.5
(1,1000)	1:31:C:PHE:HE2	2:37:D:PRO:HD2	20	2.5
(1,681)	1:31:C:PHE:HE1	2:37:D:PRO:HD2	20	2.5
(1,681)	1:31:C:PHE:HE2	2:37:D:PRO:HD2	20	2.5
(1,520)	2:17:B:ILE:HG13	2:40:D:PHE:HB3	18	2.49
(1,201)	2:17:B:ILE:HG13	2:40:D:PHE:HB3	18	2.49
(1,665)	2:19:B:PHE:HE1	2:38:D:ILE:HD11	6	2.48
(1,665)	2:19:B:PHE:HE1	2:38:D:ILE:HD12	6	2.48
(1,665)	2:19:B:PHE:HE1	2:38:D:ILE:HD13	6	2.48
(1,665)	2:19:B:PHE:HE2	2:38:D:ILE:HD11	6	2.48

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,665)	2:19:B:PHE:HE2	2:38:D:ILE:HD12	6	2.48
(1,665)	2:19:B:PHE:HE2	2:38:D:ILE:HD13	6	2.48
(1,18)	2:19:B:PHE:HE1	2:38:D:ILE:HD11	6	2.48
(1,18)	2:19:B:PHE:HE1	2:38:D:ILE:HD12	6	2.48
(1,18)	2:19:B:PHE:HE1	2:38:D:ILE:HD13	6	2.48
(1,18)	2:19:B:PHE:HE2	2:38:D:ILE:HD11	6	2.48
(1,18)	2:19:B:PHE:HE2	2:38:D:ILE:HD12	6	2.48
(1,18)	2:19:B:PHE:HE2	2:38:D:ILE:HD13	6	2.48
(1,1017)	1:31:C:PHE:HE1	2:35:D:GLY:HA3	1	2.46
(1,1017)	1:31:C:PHE:HE2	2:35:D:GLY:HA3	1	2.46
(1,698)	1:31:C:PHE:HE1	2:35:D:GLY:HA3	1	2.46
(1,698)	1:31:C:PHE:HE2	2:35:D:GLY:HA3	1	2.46
(1,664)	2:19:B:PHE:HB2	2:38:D:ILE:HG13	19	2.46
(1,17)	2:19:B:PHE:HB2	2:38:D:ILE:HG13	19	2.46
(1,415)	1:6:A:VAL:HG11	2:20:B:TYR:HE1	9	2.45
(1,415)	1:6:A:VAL:HG11	2:20:B:TYR:HE2	9	2.45
(1,415)	1:6:A:VAL:HG12	2:20:B:TYR:HE1	9	2.45
(1,415)	1:6:A:VAL:HG12	2:20:B:TYR:HE2	9	2.45
(1,415)	1:6:A:VAL:HG13	2:20:B:TYR:HE1	9	2.45
(1,415)	1:6:A:VAL:HG13	2:20:B:TYR:HE2	9	2.45
(1,96)	1:6:A:VAL:HG11	2:20:B:TYR:HE1	9	2.45
(1,96)	1:6:A:VAL:HG11	2:20:B:TYR:HE2	9	2.45
(1,96)	1:6:A:VAL:HG12	2:20:B:TYR:HE1	9	2.45
(1,96)	1:6:A:VAL:HG12	2:20:B:TYR:HE2	9	2.45
(1,96)	1:6:A:VAL:HG13	2:20:B:TYR:HE1	9	2.45
(1,96)	1:6:A:VAL:HG13	2:20:B:TYR:HE2	9	2.45
(1,402)	1:10:A:PHE:HD1	2:14:B:GLY:HA3	8	2.44
(1,402)	1:10:A:PHE:HD2	2:14:B:GLY:HA3	8	2.44
(1,83)	1:10:A:PHE:HD1	2:14:B:GLY:HA3	8	2.44
(1,83)	1:10:A:PHE:HD2	2:14:B:GLY:HA3	8	2.44
(1,1073)	2:18:B:SER:HB3	2:38:D:ILE:HD11	5	2.43
(1,1073)	2:18:B:SER:HB3	2:38:D:ILE:HD12	5	2.43
(1,1073)	2:18:B:SER:HB3	2:38:D:ILE:HD13	5	2.43
(1,754)	2:18:B:SER:HB3	2:38:D:ILE:HD11	5	2.43
(1,754)	2:18:B:SER:HB3	2:38:D:ILE:HD12	5	2.43
(1,754)	2:18:B:SER:HB3	2:38:D:ILE:HD13	5	2.43
(1,437)	1:8:A:GLU:HG2	2:20:B:TYR:HB2	13	2.43
(1,437)	1:8:A:GLU:HG2	2:20:B:TYR:HB3	13	2.43
(1,118)	1:8:A:GLU:HG2	2:20:B:TYR:HB2	13	2.43
(1,118)	1:8:A:GLU:HG2	2:20:B:TYR:HB3	13	2.43
(1,1176)	2:19:B:PHE:HB3	2:38:D:ILE:HG13	17	2.42
(1,857)	2:19:B:PHE:HB3	2:38:D:ILE:HG13	17	2.42

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,420)	1:4:A:GLY:HA3	1:5:A:HIS:HD2	5	2.42
(1,101)	1:4:A:GLY:HA3	1:5:A:HIS:HD2	5	2.42
(1,664)	2:19:B:PHE:HB2	2:38:D:ILE:HG13	13	2.41
(1,17)	2:19:B:PHE:HB2	2:38:D:ILE:HG13	13	2.41
(1,1176)	2:19:B:PHE:HB3	2:38:D:ILE:HG13	19	2.4
(1,857)	2:19:B:PHE:HB3	2:38:D:ILE:HG13	19	2.4
(1,663)	2:19:B:PHE:HD1	2:38:D:ILE:HG13	20	2.4
(1,663)	2:19:B:PHE:HD2	2:38:D:ILE:HG13	20	2.4
(1,661)	2:19:B:PHE:H	2:38:D:ILE:HD11	9	2.4
(1,661)	2:19:B:PHE:H	2:38:D:ILE:HD12	9	2.4
(1,661)	2:19:B:PHE:H	2:38:D:ILE:HD13	9	2.4
(1,595)	1:12:A:ARG:HA	1:12:A:ARG:HE	15	2.4
(1,491)	1:7:A:PRO:HB2	1:10:A:PHE:HE1	19	2.4
(1,491)	1:7:A:PRO:HB2	1:10:A:PHE:HE2	19	2.4
(1,276)	1:12:A:ARG:HA	1:12:A:ARG:HE	15	2.4
(1,172)	1:7:A:PRO:HB2	1:10:A:PHE:HE1	19	2.4
(1,172)	1:7:A:PRO:HB2	1:10:A:PHE:HE2	19	2.4
(1,16)	2:19:B:PHE:HD1	2:38:D:ILE:HG13	20	2.4
(1,16)	2:19:B:PHE:HD2	2:38:D:ILE:HG13	20	2.4
(1,14)	2:19:B:PHE:H	2:38:D:ILE:HD11	9	2.4
(1,14)	2:19:B:PHE:H	2:38:D:ILE:HD12	9	2.4
(1,14)	2:19:B:PHE:H	2:38:D:ILE:HD13	9	2.4
(1,595)	1:12:A:ARG:HA	1:12:A:ARG:HE	18	2.39
(1,276)	1:12:A:ARG:HA	1:12:A:ARG:HE	18	2.39
(1,1040)	1:31:C:PHE:HE1	2:37:D:PRO:HD2	20	2.38
(1,1040)	1:31:C:PHE:HE2	2:37:D:PRO:HD2	20	2.38
(1,1017)	1:31:C:PHE:HE1	2:35:D:GLY:HA3	20	2.38
(1,1017)	1:31:C:PHE:HE2	2:35:D:GLY:HA3	20	2.38
(1,721)	1:31:C:PHE:HE1	2:37:D:PRO:HD2	20	2.38
(1,721)	1:31:C:PHE:HE2	2:37:D:PRO:HD2	20	2.38
(1,698)	1:31:C:PHE:HE1	2:35:D:GLY:HA3	20	2.38
(1,698)	1:31:C:PHE:HE2	2:35:D:GLY:HA3	20	2.38
(1,1169)	2:36:D:THR:HB	2:38:D:ILE:HD11	18	2.37
(1,1169)	2:36:D:THR:HB	2:38:D:ILE:HD12	18	2.37
(1,1169)	2:36:D:THR:HB	2:38:D:ILE:HD13	18	2.37
(1,850)	2:36:D:THR:HB	2:38:D:ILE:HD11	18	2.37
(1,850)	2:36:D:THR:HB	2:38:D:ILE:HD12	18	2.37
(1,850)	2:36:D:THR:HB	2:38:D:ILE:HD13	18	2.37
(1,1058)	1:31:C:PHE:HD1	2:35:D:GLY:HA3	18	2.36
(1,1058)	1:31:C:PHE:HD2	2:35:D:GLY:HA3	18	2.36
(1,739)	1:31:C:PHE:HD1	2:35:D:GLY:HA3	18	2.36
(1,739)	1:31:C:PHE:HD2	2:35:D:GLY:HA3	18	2.36

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,1251)	1:33:C:ARG:HA	1:33:C:ARG:HE	5	2.35
(1,1176)	2:19:B:PHE:HB3	2:38:D:ILE:HG13	8	2.35
(1,932)	1:33:C:ARG:HA	1:33:C:ARG:HE	5	2.35
(1,857)	2:19:B:PHE:HB3	2:38:D:ILE:HG13	8	2.35
(1,402)	1:10:A:PHE:HD1	2:14:B:GLY:HA3	15	2.35
(1,402)	1:10:A:PHE:HD2	2:14:B:GLY:HA3	15	2.35
(1,402)	1:10:A:PHE:HD1	2:14:B:GLY:HA3	18	2.35
(1,402)	1:10:A:PHE:HD2	2:14:B:GLY:HA3	18	2.35
(1,361)	1:10:A:PHE:HE1	2:14:B:GLY:HA3	15	2.35
(1,361)	1:10:A:PHE:HE2	2:14:B:GLY:HA3	15	2.35
(1,83)	1:10:A:PHE:HD1	2:14:B:GLY:HA3	15	2.35
(1,83)	1:10:A:PHE:HD2	2:14:B:GLY:HA3	15	2.35
(1,83)	1:10:A:PHE:HD1	2:14:B:GLY:HA3	18	2.35
(1,83)	1:10:A:PHE:HD2	2:14:B:GLY:HA3	18	2.35
(1,42)	1:10:A:PHE:HE1	2:14:B:GLY:HA3	15	2.35
(1,42)	1:10:A:PHE:HE2	2:14:B:GLY:HA3	15	2.35
(1,1171)	2:36:D:THR:HG21	2:38:D:ILE:HD11	10	2.33
(1,1171)	2:36:D:THR:HG21	2:38:D:ILE:HD12	10	2.33
(1,1171)	2:36:D:THR:HG21	2:38:D:ILE:HD13	10	2.33
(1,1171)	2:36:D:THR:HG22	2:38:D:ILE:HD11	10	2.33
(1,1171)	2:36:D:THR:HG22	2:38:D:ILE:HD12	10	2.33
(1,1171)	2:36:D:THR:HG22	2:38:D:ILE:HD13	10	2.33
(1,1171)	2:36:D:THR:HG23	2:38:D:ILE:HD11	10	2.33
(1,1171)	2:36:D:THR:HG23	2:38:D:ILE:HD12	10	2.33
(1,1171)	2:36:D:THR:HG23	2:38:D:ILE:HD13	10	2.33
(1,852)	2:36:D:THR:HG21	2:38:D:ILE:HD11	10	2.33
(1,852)	2:36:D:THR:HG21	2:38:D:ILE:HD12	10	2.33
(1,852)	2:36:D:THR:HG21	2:38:D:ILE:HD13	10	2.33
(1,852)	2:36:D:THR:HG22	2:38:D:ILE:HD11	10	2.33
(1,852)	2:36:D:THR:HG22	2:38:D:ILE:HD12	10	2.33
(1,852)	2:36:D:THR:HG22	2:38:D:ILE:HD13	10	2.33
(1,852)	2:36:D:THR:HG23	2:38:D:ILE:HD11	10	2.33
(1,852)	2:36:D:THR:HG23	2:38:D:ILE:HD12	10	2.33
(1,852)	2:36:D:THR:HG23	2:38:D:ILE:HD13	10	2.33
(1,662)	2:19:B:PHE:HE1	2:38:D:ILE:HG13	5	2.33
(1,662)	2:19:B:PHE:HE2	2:38:D:ILE:HG13	5	2.33
(1,661)	2:19:B:PHE:H	2:38:D:ILE:HD11	13	2.33
(1,661)	2:19:B:PHE:H	2:38:D:ILE:HD12	13	2.33
(1,661)	2:19:B:PHE:H	2:38:D:ILE:HD13	13	2.33
(1,361)	1:10:A:PHE:HE1	2:14:B:GLY:HA3	17	2.33
(1,361)	1:10:A:PHE:HE2	2:14:B:GLY:HA3	17	2.33
(1,42)	1:10:A:PHE:HE1	2:14:B:GLY:HA3	17	2.33

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,42)	1:10:A:PHE:HE2	2:14:B:GLY:HA3	17	2.33
(1,15)	2:19:B:PHE:HE1	2:38:D:ILE:HG13	5	2.33
(1,15)	2:19:B:PHE:HE2	2:38:D:ILE:HG13	5	2.33
(1,14)	2:19:B:PHE:H	2:38:D:ILE:HD11	13	2.33
(1,14)	2:19:B:PHE:H	2:38:D:ILE:HD12	13	2.33
(1,14)	2:19:B:PHE:H	2:38:D:ILE:HD13	13	2.33
(1,1159)	1:28:C:PRO:HB3	1:31:C:PHE:HE1	2	2.32
(1,1159)	1:28:C:PRO:HB3	1:31:C:PHE:HE2	2	2.32
(1,840)	1:28:C:PRO:HB3	1:31:C:PHE:HE1	2	2.32
(1,840)	1:28:C:PRO:HB3	1:31:C:PHE:HE2	2	2.32
(1,1058)	1:31:C:PHE:HD1	2:35:D:GLY:HA3	17	2.31
(1,1058)	1:31:C:PHE:HD2	2:35:D:GLY:HA3	17	2.31
(1,739)	1:31:C:PHE:HD1	2:35:D:GLY:HA3	17	2.31
(1,739)	1:31:C:PHE:HD2	2:35:D:GLY:HA3	17	2.31
(1,466)	1:4:A:GLY:HA2	1:5:A:HIS:HD2	10	2.31
(1,147)	1:4:A:GLY:HA2	1:5:A:HIS:HD2	10	2.31
(1,1058)	1:31:C:PHE:HD1	2:35:D:GLY:HA3	19	2.3
(1,1058)	1:31:C:PHE:HD2	2:35:D:GLY:HA3	19	2.3
(1,739)	1:31:C:PHE:HD1	2:35:D:GLY:HA3	19	2.3
(1,739)	1:31:C:PHE:HD2	2:35:D:GLY:HA3	19	2.3
(1,515)	2:15:B:THR:HG21	2:17:B:ILE:HD11	14	2.3
(1,515)	2:15:B:THR:HG21	2:17:B:ILE:HD12	14	2.3
(1,515)	2:15:B:THR:HG21	2:17:B:ILE:HD13	14	2.3
(1,515)	2:15:B:THR:HG22	2:17:B:ILE:HD11	14	2.3
(1,515)	2:15:B:THR:HG22	2:17:B:ILE:HD12	14	2.3
(1,515)	2:15:B:THR:HG22	2:17:B:ILE:HD13	14	2.3
(1,515)	2:15:B:THR:HG23	2:17:B:ILE:HD11	14	2.3
(1,515)	2:15:B:THR:HG23	2:17:B:ILE:HD12	14	2.3
(1,515)	2:15:B:THR:HG23	2:17:B:ILE:HD13	14	2.3
(1,491)	1:7:A:PRO:HB2	1:10:A:PHE:HE1	3	2.3
(1,491)	1:7:A:PRO:HB2	1:10:A:PHE:HE2	3	2.3
(1,196)	2:15:B:THR:HG21	2:17:B:ILE:HD11	14	2.3
(1,196)	2:15:B:THR:HG21	2:17:B:ILE:HD12	14	2.3
(1,196)	2:15:B:THR:HG21	2:17:B:ILE:HD13	14	2.3
(1,196)	2:15:B:THR:HG22	2:17:B:ILE:HD11	14	2.3
(1,196)	2:15:B:THR:HG22	2:17:B:ILE:HD12	14	2.3
(1,196)	2:15:B:THR:HG22	2:17:B:ILE:HD13	14	2.3
(1,196)	2:15:B:THR:HG23	2:17:B:ILE:HD11	14	2.3
(1,196)	2:15:B:THR:HG23	2:17:B:ILE:HD12	14	2.3
(1,196)	2:15:B:THR:HG23	2:17:B:ILE:HD13	14	2.3
(1,172)	1:7:A:PRO:HB2	1:10:A:PHE:HE1	3	2.3
(1,172)	1:7:A:PRO:HB2	1:10:A:PHE:HE2	3	2.3

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,1121)	1:26:C:HIS:HD2	1:27:C:VAL:HG11	13	2.29
(1,1121)	1:26:C:HIS:HD2	1:27:C:VAL:HG12	13	2.29
(1,1121)	1:26:C:HIS:HD2	1:27:C:VAL:HG13	13	2.29
(1,802)	1:26:C:HIS:HD2	1:27:C:VAL:HG11	13	2.29
(1,802)	1:26:C:HIS:HD2	1:27:C:VAL:HG12	13	2.29
(1,802)	1:26:C:HIS:HD2	1:27:C:VAL:HG13	13	2.29
(1,503)	1:7:A:PRO:HB3	1:10:A:PHE:HE1	3	2.29
(1,503)	1:7:A:PRO:HB3	1:10:A:PHE:HE2	3	2.29
(1,463)	1:5:A:HIS:HB2	1:6:A:VAL:HG11	18	2.29
(1,463)	1:5:A:HIS:HB2	1:6:A:VAL:HG12	18	2.29
(1,463)	1:5:A:HIS:HB2	1:6:A:VAL:HG13	18	2.29
(1,184)	1:7:A:PRO:HB3	1:10:A:PHE:HE1	3	2.29
(1,184)	1:7:A:PRO:HB3	1:10:A:PHE:HE2	3	2.29
(1,144)	1:5:A:HIS:HB2	1:6:A:VAL:HG11	18	2.29
(1,144)	1:5:A:HIS:HB2	1:6:A:VAL:HG12	18	2.29
(1,144)	1:5:A:HIS:HB2	1:6:A:VAL:HG13	18	2.29
(1,1073)	2:18:B:SER:HB3	2:38:D:ILE:HD11	6	2.28
(1,1073)	2:18:B:SER:HB3	2:38:D:ILE:HD12	6	2.28
(1,1073)	2:18:B:SER:HB3	2:38:D:ILE:HD13	6	2.28
(1,754)	2:18:B:SER:HB3	2:38:D:ILE:HD11	6	2.28
(1,754)	2:18:B:SER:HB3	2:38:D:ILE:HD12	6	2.28
(1,754)	2:18:B:SER:HB3	2:38:D:ILE:HD13	6	2.28
(1,663)	2:19:B:PHE:HD1	2:38:D:ILE:HG13	14	2.28
(1,663)	2:19:B:PHE:HD2	2:38:D:ILE:HG13	14	2.28
(1,662)	2:19:B:PHE:HE1	2:38:D:ILE:HG13	17	2.28
(1,662)	2:19:B:PHE:HE2	2:38:D:ILE:HG13	17	2.28
(1,361)	1:10:A:PHE:HE1	2:14:B:GLY:HA3	1	2.28
(1,361)	1:10:A:PHE:HE2	2:14:B:GLY:HA3	1	2.28
(1,42)	1:10:A:PHE:HE1	2:14:B:GLY:HA3	1	2.28
(1,42)	1:10:A:PHE:HE2	2:14:B:GLY:HA3	1	2.28
(1,16)	2:19:B:PHE:HD1	2:38:D:ILE:HG13	14	2.28
(1,16)	2:19:B:PHE:HD2	2:38:D:ILE:HG13	14	2.28
(1,15)	2:19:B:PHE:HE1	2:38:D:ILE:HG13	17	2.28
(1,15)	2:19:B:PHE:HE2	2:38:D:ILE:HG13	17	2.28
(1,1121)	1:26:C:HIS:HD2	1:27:C:VAL:HG11	8	2.27
(1,1121)	1:26:C:HIS:HD2	1:27:C:VAL:HG12	8	2.27
(1,1121)	1:26:C:HIS:HD2	1:27:C:VAL:HG13	8	2.27
(1,1093)	1:29:C:GLU:HG2	2:41:D:TYR:HB2	9	2.27
(1,1093)	1:29:C:GLU:HG2	2:41:D:TYR:HB3	9	2.27
(1,802)	1:26:C:HIS:HD2	1:27:C:VAL:HG11	8	2.27
(1,802)	1:26:C:HIS:HD2	1:27:C:VAL:HG12	8	2.27
(1,802)	1:26:C:HIS:HD2	1:27:C:VAL:HG13	8	2.27

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,774)	1:29:C:GLU:HG2	2:41:D:TYR:HB2	9	2.27
(1,774)	1:29:C:GLU:HG2	2:41:D:TYR:HB3	9	2.27
(1,515)	2:15:B:THR:HG21	2:17:B:ILE:HD11	6	2.27
(1,515)	2:15:B:THR:HG21	2:17:B:ILE:HD12	6	2.27
(1,515)	2:15:B:THR:HG21	2:17:B:ILE:HD13	6	2.27
(1,515)	2:15:B:THR:HG22	2:17:B:ILE:HD11	6	2.27
(1,515)	2:15:B:THR:HG22	2:17:B:ILE:HD12	6	2.27
(1,515)	2:15:B:THR:HG22	2:17:B:ILE:HD13	6	2.27
(1,515)	2:15:B:THR:HG23	2:17:B:ILE:HD11	6	2.27
(1,515)	2:15:B:THR:HG23	2:17:B:ILE:HD12	6	2.27
(1,515)	2:15:B:THR:HG23	2:17:B:ILE:HD13	6	2.27
(1,491)	1:7:A:PRO:HB2	1:10:A:PHE:HE1	10	2.27
(1,491)	1:7:A:PRO:HB2	1:10:A:PHE:HE2	10	2.27
(1,196)	2:15:B:THR:HG21	2:17:B:ILE:HD11	6	2.27
(1,196)	2:15:B:THR:HG21	2:17:B:ILE:HD12	6	2.27
(1,196)	2:15:B:THR:HG21	2:17:B:ILE:HD13	6	2.27
(1,196)	2:15:B:THR:HG22	2:17:B:ILE:HD11	6	2.27
(1,196)	2:15:B:THR:HG22	2:17:B:ILE:HD12	6	2.27
(1,196)	2:15:B:THR:HG22	2:17:B:ILE:HD13	6	2.27
(1,196)	2:15:B:THR:HG23	2:17:B:ILE:HD11	6	2.27
(1,196)	2:15:B:THR:HG23	2:17:B:ILE:HD12	6	2.27
(1,196)	2:15:B:THR:HG23	2:17:B:ILE:HD13	6	2.27
(1,172)	1:7:A:PRO:HB2	1:10:A:PHE:HE1	10	2.27
(1,172)	1:7:A:PRO:HB2	1:10:A:PHE:HE2	10	2.27
(1,1122)	1:25:C:GLY:HA2	1:26:C:HIS:HD2	7	2.26
(1,1071)	1:27:C:VAL:HG11	2:41:D:TYR:HE1	11	2.26
(1,1071)	1:27:C:VAL:HG11	2:41:D:TYR:HE2	11	2.26
(1,1071)	1:27:C:VAL:HG12	2:41:D:TYR:HE1	11	2.26
(1,1071)	1:27:C:VAL:HG12	2:41:D:TYR:HE2	11	2.26
(1,1071)	1:27:C:VAL:HG13	2:41:D:TYR:HE1	11	2.26
(1,1071)	1:27:C:VAL:HG13	2:41:D:TYR:HE2	11	2.26
(1,803)	1:25:C:GLY:HA2	1:26:C:HIS:HD2	7	2.26
(1,752)	1:27:C:VAL:HG11	2:41:D:TYR:HE1	11	2.26
(1,752)	1:27:C:VAL:HG11	2:41:D:TYR:HE2	11	2.26
(1,752)	1:27:C:VAL:HG12	2:41:D:TYR:HE1	11	2.26
(1,752)	1:27:C:VAL:HG12	2:41:D:TYR:HE2	11	2.26
(1,752)	1:27:C:VAL:HG13	2:41:D:TYR:HE1	11	2.26
(1,752)	1:27:C:VAL:HG13	2:41:D:TYR:HE2	11	2.26
(1,662)	2:19:B:PHE:HE1	2:38:D:ILE:HG13	20	2.26
(1,662)	2:19:B:PHE:HE2	2:38:D:ILE:HG13	20	2.26
(1,437)	1:8:A:GLU:HG2	2:20:B:TYR:HB2	18	2.26
(1,437)	1:8:A:GLU:HG2	2:20:B:TYR:HB3	18	2.26

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,436)	1:8:A:GLU:HG2	2:20:B:TYR:HD1	10	2.26
(1,436)	1:8:A:GLU:HG2	2:20:B:TYR:HD2	10	2.26
(1,415)	1:6:A:VAL:HG11	2:20:B:TYR:HE1	8	2.26
(1,415)	1:6:A:VAL:HG11	2:20:B:TYR:HE2	8	2.26
(1,415)	1:6:A:VAL:HG12	2:20:B:TYR:HE1	8	2.26
(1,415)	1:6:A:VAL:HG12	2:20:B:TYR:HE2	8	2.26
(1,415)	1:6:A:VAL:HG13	2:20:B:TYR:HE1	8	2.26
(1,415)	1:6:A:VAL:HG13	2:20:B:TYR:HE2	8	2.26
(1,118)	1:8:A:GLU:HG2	2:20:B:TYR:HB2	18	2.26
(1,118)	1:8:A:GLU:HG2	2:20:B:TYR:HB3	18	2.26
(1,117)	1:8:A:GLU:HG2	2:20:B:TYR:HD1	10	2.26
(1,117)	1:8:A:GLU:HG2	2:20:B:TYR:HD2	10	2.26
(1,96)	1:6:A:VAL:HG11	2:20:B:TYR:HE1	8	2.26
(1,96)	1:6:A:VAL:HG11	2:20:B:TYR:HE2	8	2.26
(1,96)	1:6:A:VAL:HG12	2:20:B:TYR:HE1	8	2.26
(1,96)	1:6:A:VAL:HG12	2:20:B:TYR:HE2	8	2.26
(1,96)	1:6:A:VAL:HG13	2:20:B:TYR:HE1	8	2.26
(1,96)	1:6:A:VAL:HG13	2:20:B:TYR:HE2	8	2.26
(1,15)	2:19:B:PHE:HE1	2:38:D:ILE:HG13	20	2.26
(1,15)	2:19:B:PHE:HE2	2:38:D:ILE:HG13	20	2.26
(1,670)	2:17:B:ILE:HD11	2:40:D:PHE:H	9	2.25
(1,670)	2:17:B:ILE:HD12	2:40:D:PHE:H	9	2.25
(1,670)	2:17:B:ILE:HD13	2:40:D:PHE:H	9	2.25
(1,657)	2:19:B:PHE:HD1	2:38:D:ILE:HD11	19	2.25
(1,657)	2:19:B:PHE:HD1	2:38:D:ILE:HD12	19	2.25
(1,657)	2:19:B:PHE:HD1	2:38:D:ILE:HD13	19	2.25
(1,657)	2:19:B:PHE:HD2	2:38:D:ILE:HD11	19	2.25
(1,657)	2:19:B:PHE:HD2	2:38:D:ILE:HD12	19	2.25
(1,657)	2:19:B:PHE:HD2	2:38:D:ILE:HD13	19	2.25
(1,10)	2:19:B:PHE:HD1	2:38:D:ILE:HD11	19	2.25
(1,10)	2:19:B:PHE:HD1	2:38:D:ILE:HD12	19	2.25
(1,10)	2:19:B:PHE:HD1	2:38:D:ILE:HD13	19	2.25
(1,10)	2:19:B:PHE:HD2	2:38:D:ILE:HD11	19	2.25
(1,10)	2:19:B:PHE:HD2	2:38:D:ILE:HD12	19	2.25
(1,10)	2:19:B:PHE:HD2	2:38:D:ILE:HD13	19	2.25
(1,5)	2:17:B:ILE:HD11	2:40:D:PHE:H	9	2.25
(1,5)	2:17:B:ILE:HD12	2:40:D:PHE:H	9	2.25
(1,5)	2:17:B:ILE:HD13	2:40:D:PHE:H	9	2.25
(1,1171)	2:36:D:THR:HG21	2:38:D:ILE:HD11	13	2.24
(1,1171)	2:36:D:THR:HG21	2:38:D:ILE:HD12	13	2.24
(1,1171)	2:36:D:THR:HG21	2:38:D:ILE:HD13	13	2.24
(1,1171)	2:36:D:THR:HG22	2:38:D:ILE:HD11	13	2.24

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,1171)	2:36:D:THR:HG22	2:38:D:ILE:HD12	13	2.24
(1,1171)	2:36:D:THR:HG22	2:38:D:ILE:HD13	13	2.24
(1,1171)	2:36:D:THR:HG23	2:38:D:ILE:HD11	13	2.24
(1,1171)	2:36:D:THR:HG23	2:38:D:ILE:HD12	13	2.24
(1,1171)	2:36:D:THR:HG23	2:38:D:ILE:HD13	13	2.24
(1,852)	2:36:D:THR:HG21	2:38:D:ILE:HD11	13	2.24
(1,852)	2:36:D:THR:HG21	2:38:D:ILE:HD12	13	2.24
(1,852)	2:36:D:THR:HG21	2:38:D:ILE:HD13	13	2.24
(1,852)	2:36:D:THR:HG22	2:38:D:ILE:HD11	13	2.24
(1,852)	2:36:D:THR:HG22	2:38:D:ILE:HD12	13	2.24
(1,852)	2:36:D:THR:HG22	2:38:D:ILE:HD13	13	2.24
(1,852)	2:36:D:THR:HG23	2:38:D:ILE:HD11	13	2.24
(1,852)	2:36:D:THR:HG23	2:38:D:ILE:HD12	13	2.24
(1,852)	2:36:D:THR:HG23	2:38:D:ILE:HD13	13	2.24
(1,420)	1:4:A:GLY:HA3	1:5:A:HIS:HD2	18	2.24
(1,101)	1:4:A:GLY:HA3	1:5:A:HIS:HD2	18	2.24
(1,1159)	1:28:C:PRO:HB3	1:31:C:PHE:HE1	9	2.23
(1,1159)	1:28:C:PRO:HB3	1:31:C:PHE:HE2	9	2.23
(1,1093)	1:29:C:GLU:HG2	2:41:D:TYR:HB2	20	2.23
(1,1093)	1:29:C:GLU:HG2	2:41:D:TYR:HB3	20	2.23
(1,840)	1:28:C:PRO:HB3	1:31:C:PHE:HE1	9	2.23
(1,840)	1:28:C:PRO:HB3	1:31:C:PHE:HE2	9	2.23
(1,774)	1:29:C:GLU:HG2	2:41:D:TYR:HB2	20	2.23
(1,774)	1:29:C:GLU:HG2	2:41:D:TYR:HB3	20	2.23
(1,513)	2:15:B:THR:HB	2:17:B:ILE:HD11	15	2.23
(1,513)	2:15:B:THR:HB	2:17:B:ILE:HD12	15	2.23
(1,513)	2:15:B:THR:HB	2:17:B:ILE:HD13	15	2.23
(1,194)	2:15:B:THR:HB	2:17:B:ILE:HD11	15	2.23
(1,194)	2:15:B:THR:HB	2:17:B:ILE:HD12	15	2.23
(1,194)	2:15:B:THR:HB	2:17:B:ILE:HD13	15	2.23
(1,1251)	1:33:C:ARG:HA	1:33:C:ARG:HE	14	2.22
(1,1071)	1:27:C:VAL:HG11	2:41:D:TYR:HE1	1	2.22
(1,1071)	1:27:C:VAL:HG11	2:41:D:TYR:HE2	1	2.22
(1,1071)	1:27:C:VAL:HG12	2:41:D:TYR:HE1	1	2.22
(1,1071)	1:27:C:VAL:HG12	2:41:D:TYR:HE2	1	2.22
(1,1071)	1:27:C:VAL:HG13	2:41:D:TYR:HE1	1	2.22
(1,1071)	1:27:C:VAL:HG13	2:41:D:TYR:HE2	1	2.22
(1,1071)	1:27:C:VAL:HG11	2:41:D:TYR:HE1	13	2.22
(1,1071)	1:27:C:VAL:HG11	2:41:D:TYR:HE2	13	2.22
(1,1071)	1:27:C:VAL:HG12	2:41:D:TYR:HE1	13	2.22
(1,1071)	1:27:C:VAL:HG12	2:41:D:TYR:HE2	13	2.22
(1,1071)	1:27:C:VAL:HG13	2:41:D:TYR:HE1	13	2.22

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,1071)	1:27:C:VAL:HG13	2:41:D:TYR:HE2	13	2.22
(1,932)	1:33:C:ARG:HA	1:33:C:ARG:HE	14	2.22
(1,752)	1:27:C:VAL:HG11	2:41:D:TYR:HE1	1	2.22
(1,752)	1:27:C:VAL:HG11	2:41:D:TYR:HE2	1	2.22
(1,752)	1:27:C:VAL:HG12	2:41:D:TYR:HE1	1	2.22
(1,752)	1:27:C:VAL:HG12	2:41:D:TYR:HE2	1	2.22
(1,752)	1:27:C:VAL:HG13	2:41:D:TYR:HE1	1	2.22
(1,752)	1:27:C:VAL:HG13	2:41:D:TYR:HE2	1	2.22
(1,752)	1:27:C:VAL:HG11	2:41:D:TYR:HE1	13	2.22
(1,752)	1:27:C:VAL:HG11	2:41:D:TYR:HE2	13	2.22
(1,752)	1:27:C:VAL:HG12	2:41:D:TYR:HE1	13	2.22
(1,752)	1:27:C:VAL:HG12	2:41:D:TYR:HE2	13	2.22
(1,752)	1:27:C:VAL:HG13	2:41:D:TYR:HE1	13	2.22
(1,752)	1:27:C:VAL:HG13	2:41:D:TYR:HE2	13	2.22
(1,513)	2:15:B:THR:HB	2:17:B:ILE:HD11	17	2.22
(1,513)	2:15:B:THR:HB	2:17:B:ILE:HD12	17	2.22
(1,513)	2:15:B:THR:HB	2:17:B:ILE:HD13	17	2.22
(1,194)	2:15:B:THR:HB	2:17:B:ILE:HD11	17	2.22
(1,194)	2:15:B:THR:HB	2:17:B:ILE:HD12	17	2.22
(1,194)	2:15:B:THR:HB	2:17:B:ILE:HD13	17	2.22
(1,1093)	1:29:C:GLU:HG2	2:41:D:TYR:HB2	6	2.21
(1,1093)	1:29:C:GLU:HG2	2:41:D:TYR:HB3	6	2.21
(1,774)	1:29:C:GLU:HG2	2:41:D:TYR:HB2	6	2.21
(1,774)	1:29:C:GLU:HG2	2:41:D:TYR:HB3	6	2.21
(1,437)	1:8:A:GLU:HG2	2:20:B:TYR:HB2	20	2.21
(1,437)	1:8:A:GLU:HG2	2:20:B:TYR:HB3	20	2.21
(1,415)	1:6:A:VAL:HG11	2:20:B:TYR:HE1	1	2.21
(1,415)	1:6:A:VAL:HG11	2:20:B:TYR:HE2	1	2.21
(1,415)	1:6:A:VAL:HG12	2:20:B:TYR:HE1	1	2.21
(1,415)	1:6:A:VAL:HG12	2:20:B:TYR:HE2	1	2.21
(1,415)	1:6:A:VAL:HG13	2:20:B:TYR:HE1	1	2.21
(1,415)	1:6:A:VAL:HG13	2:20:B:TYR:HE2	1	2.21
(1,118)	1:8:A:GLU:HG2	2:20:B:TYR:HB2	20	2.21
(1,118)	1:8:A:GLU:HG2	2:20:B:TYR:HB3	20	2.21
(1,96)	1:6:A:VAL:HG11	2:20:B:TYR:HE1	1	2.21
(1,96)	1:6:A:VAL:HG11	2:20:B:TYR:HE2	1	2.21
(1,96)	1:6:A:VAL:HG12	2:20:B:TYR:HE1	1	2.21
(1,96)	1:6:A:VAL:HG12	2:20:B:TYR:HE2	1	2.21
(1,96)	1:6:A:VAL:HG13	2:20:B:TYR:HE1	1	2.21
(1,96)	1:6:A:VAL:HG13	2:20:B:TYR:HE2	1	2.21
(1,515)	2:15:B:THR:HG21	2:17:B:ILE:HD11	10	2.2
(1,515)	2:15:B:THR:HG21	2:17:B:ILE:HD12	10	2.2

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,515)	2:15:B:THR:HG21	2:17:B:ILE:HD13	10	2.2
(1,515)	2:15:B:THR:HG22	2:17:B:ILE:HD11	10	2.2
(1,515)	2:15:B:THR:HG22	2:17:B:ILE:HD12	10	2.2
(1,515)	2:15:B:THR:HG22	2:17:B:ILE:HD13	10	2.2
(1,515)	2:15:B:THR:HG23	2:17:B:ILE:HD11	10	2.2
(1,515)	2:15:B:THR:HG23	2:17:B:ILE:HD12	10	2.2
(1,515)	2:15:B:THR:HG23	2:17:B:ILE:HD13	10	2.2
(1,196)	2:15:B:THR:HG21	2:17:B:ILE:HD11	10	2.2
(1,196)	2:15:B:THR:HG21	2:17:B:ILE:HD12	10	2.2
(1,196)	2:15:B:THR:HG21	2:17:B:ILE:HD13	10	2.2
(1,196)	2:15:B:THR:HG22	2:17:B:ILE:HD11	10	2.2
(1,196)	2:15:B:THR:HG22	2:17:B:ILE:HD12	10	2.2
(1,196)	2:15:B:THR:HG22	2:17:B:ILE:HD13	10	2.2
(1,196)	2:15:B:THR:HG23	2:17:B:ILE:HD11	10	2.2
(1,196)	2:15:B:THR:HG23	2:17:B:ILE:HD12	10	2.2
(1,196)	2:15:B:THR:HG23	2:17:B:ILE:HD13	10	2.2
(1,998)	1:22:C:GLY:HA3	2:39:D:SER:HB2	1	2.19
(1,995)	1:22:C:GLY:HA3	2:39:D:SER:HB2	1	2.19
(1,679)	1:22:C:GLY:HA3	2:39:D:SER:HB2	1	2.19
(1,676)	1:22:C:GLY:HA3	2:39:D:SER:HB2	1	2.19
(1,1092)	1:29:C:GLU:HG2	2:41:D:TYR:HD1	1	2.18
(1,1092)	1:29:C:GLU:HG2	2:41:D:TYR:HD2	1	2.18
(1,1076)	1:25:C:GLY:HA3	1:26:C:HIS:HD2	6	2.18
(1,773)	1:29:C:GLU:HG2	2:41:D:TYR:HD1	1	2.18
(1,773)	1:29:C:GLU:HG2	2:41:D:TYR:HD2	1	2.18
(1,757)	1:25:C:GLY:HA3	1:26:C:HIS:HD2	6	2.18
(1,520)	2:17:B:ILE:HG13	2:40:D:PHE:HB3	14	2.18
(1,201)	2:17:B:ILE:HG13	2:40:D:PHE:HB3	14	2.18
(1,1055)	1:28:C:PRO:HG2	1:31:C:PHE:HE1	2	2.17
(1,1055)	1:28:C:PRO:HG2	1:31:C:PHE:HE2	2	2.17
(1,1055)	1:28:C:PRO:HG3	1:31:C:PHE:HE1	2	2.17
(1,1055)	1:28:C:PRO:HG3	1:31:C:PHE:HE2	2	2.17
(1,1015)	1:30:C:TYR:HD1	2:39:D:SER:HA	16	2.17
(1,1015)	1:30:C:TYR:HD2	2:39:D:SER:HA	16	2.17
(1,736)	1:28:C:PRO:HG2	1:31:C:PHE:HE1	2	2.17
(1,736)	1:28:C:PRO:HG2	1:31:C:PHE:HE2	2	2.17
(1,736)	1:28:C:PRO:HG3	1:31:C:PHE:HE1	2	2.17
(1,736)	1:28:C:PRO:HG3	1:31:C:PHE:HE2	2	2.17
(1,696)	1:30:C:TYR:HD1	2:39:D:SER:HA	16	2.17
(1,696)	1:30:C:TYR:HD2	2:39:D:SER:HA	16	2.17
(1,1147)	1:28:C:PRO:HB2	1:31:C:PHE:HE1	13	2.16
(1,1147)	1:28:C:PRO:HB2	1:31:C:PHE:HE2	13	2.16

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,828)	1:28:C:PRO:HB2	1:31:C:PHE:HE1	13	2.16
(1,828)	1:28:C:PRO:HB2	1:31:C:PHE:HE2	13	2.16
(1,491)	1:7:A:PRO:HB2	1:10:A:PHE:HE1	14	2.16
(1,491)	1:7:A:PRO:HB2	1:10:A:PHE:HE2	14	2.16
(1,172)	1:7:A:PRO:HB2	1:10:A:PHE:HE1	14	2.16
(1,172)	1:7:A:PRO:HB2	1:10:A:PHE:HE2	14	2.16
(1,674)	2:17:B:ILE:HD11	2:40:D:PHE:HE1	8	2.15
(1,674)	2:17:B:ILE:HD11	2:40:D:PHE:HE2	8	2.15
(1,674)	2:17:B:ILE:HD12	2:40:D:PHE:HE1	8	2.15
(1,674)	2:17:B:ILE:HD12	2:40:D:PHE:HE2	8	2.15
(1,674)	2:17:B:ILE:HD13	2:40:D:PHE:HE1	8	2.15
(1,674)	2:17:B:ILE:HD13	2:40:D:PHE:HE2	8	2.15
(1,673)	2:17:B:ILE:HG13	2:40:D:PHE:HB2	10	2.15
(1,661)	2:19:B:PHE:H	2:38:D:ILE:HD11	5	2.15
(1,661)	2:19:B:PHE:H	2:38:D:ILE:HD12	5	2.15
(1,661)	2:19:B:PHE:H	2:38:D:ILE:HD13	5	2.15
(1,14)	2:19:B:PHE:H	2:38:D:ILE:HD11	5	2.15
(1,14)	2:19:B:PHE:H	2:38:D:ILE:HD12	5	2.15
(1,14)	2:19:B:PHE:H	2:38:D:ILE:HD13	5	2.15
(1,9)	2:17:B:ILE:HD11	2:40:D:PHE:HE1	8	2.15
(1,9)	2:17:B:ILE:HD11	2:40:D:PHE:HE2	8	2.15
(1,9)	2:17:B:ILE:HD12	2:40:D:PHE:HE1	8	2.15
(1,9)	2:17:B:ILE:HD12	2:40:D:PHE:HE2	8	2.15
(1,9)	2:17:B:ILE:HD13	2:40:D:PHE:HE1	8	2.15
(1,9)	2:17:B:ILE:HD13	2:40:D:PHE:HE2	8	2.15
(1,8)	2:17:B:ILE:HG13	2:40:D:PHE:HB2	10	2.15
(1,1046)	1:27:C:VAL:HB	2:41:D:TYR:HE1	18	2.14
(1,1046)	1:27:C:VAL:HB	2:41:D:TYR:HE2	18	2.14
(1,727)	1:27:C:VAL:HB	2:41:D:TYR:HE1	18	2.14
(1,727)	1:27:C:VAL:HB	2:41:D:TYR:HE2	18	2.14
(1,665)	2:19:B:PHE:HE1	2:38:D:ILE:HD11	19	2.14
(1,665)	2:19:B:PHE:HE1	2:38:D:ILE:HD12	19	2.14
(1,665)	2:19:B:PHE:HE1	2:38:D:ILE:HD13	19	2.14
(1,665)	2:19:B:PHE:HE2	2:38:D:ILE:HD11	19	2.14
(1,665)	2:19:B:PHE:HE2	2:38:D:ILE:HD12	19	2.14
(1,665)	2:19:B:PHE:HE2	2:38:D:ILE:HD13	19	2.14
(1,18)	2:19:B:PHE:HE1	2:38:D:ILE:HD11	19	2.14
(1,18)	2:19:B:PHE:HE1	2:38:D:ILE:HD12	19	2.14
(1,18)	2:19:B:PHE:HE1	2:38:D:ILE:HD13	19	2.14
(1,18)	2:19:B:PHE:HE2	2:38:D:ILE:HD11	19	2.14
(1,18)	2:19:B:PHE:HE2	2:38:D:ILE:HD12	19	2.14
(1,18)	2:19:B:PHE:HE2	2:38:D:ILE:HD13	19	2.14

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,1093)	1:29:C:GLU:HG2	2:41:D:TYR:HB2	17	2.13
(1,1093)	1:29:C:GLU:HG2	2:41:D:TYR:HB3	17	2.13
(1,774)	1:29:C:GLU:HG2	2:41:D:TYR:HB2	17	2.13
(1,774)	1:29:C:GLU:HG2	2:41:D:TYR:HB3	17	2.13
(1,673)	2:17:B:ILE:HG13	2:40:D:PHE:HB2	17	2.13
(1,657)	2:19:B:PHE:HD1	2:38:D:ILE:HD11	18	2.13
(1,657)	2:19:B:PHE:HD1	2:38:D:ILE:HD12	18	2.13
(1,657)	2:19:B:PHE:HD1	2:38:D:ILE:HD13	18	2.13
(1,657)	2:19:B:PHE:HD2	2:38:D:ILE:HD11	18	2.13
(1,657)	2:19:B:PHE:HD2	2:38:D:ILE:HD12	18	2.13
(1,657)	2:19:B:PHE:HD2	2:38:D:ILE:HD13	18	2.13
(1,436)	1:8:A:GLU:HG2	2:20:B:TYR:HD1	5	2.13
(1,436)	1:8:A:GLU:HG2	2:20:B:TYR:HD2	5	2.13
(1,117)	1:8:A:GLU:HG2	2:20:B:TYR:HD1	5	2.13
(1,117)	1:8:A:GLU:HG2	2:20:B:TYR:HD2	5	2.13
(1,10)	2:19:B:PHE:HD1	2:38:D:ILE:HD11	18	2.13
(1,10)	2:19:B:PHE:HD1	2:38:D:ILE:HD12	18	2.13
(1,10)	2:19:B:PHE:HD1	2:38:D:ILE:HD13	18	2.13
(1,10)	2:19:B:PHE:HD2	2:38:D:ILE:HD11	18	2.13
(1,10)	2:19:B:PHE:HD2	2:38:D:ILE:HD12	18	2.13
(1,10)	2:19:B:PHE:HD2	2:38:D:ILE:HD13	18	2.13
(1,8)	2:17:B:ILE:HG13	2:40:D:PHE:HB2	17	2.13
(1,1252)	1:28:C:PRO:HG2	1:31:C:PHE:HE1	2	2.12
(1,1252)	1:28:C:PRO:HG2	1:31:C:PHE:HE2	2	2.12
(1,1252)	1:28:C:PRO:HG3	1:31:C:PHE:HE1	2	2.12
(1,1252)	1:28:C:PRO:HG3	1:31:C:PHE:HE2	2	2.12
(1,933)	1:28:C:PRO:HG2	1:31:C:PHE:HE1	2	2.12
(1,933)	1:28:C:PRO:HG2	1:31:C:PHE:HE2	2	2.12
(1,933)	1:28:C:PRO:HG3	1:31:C:PHE:HE1	2	2.12
(1,933)	1:28:C:PRO:HG3	1:31:C:PHE:HE2	2	2.12
(1,415)	1:6:A:VAL:HG11	2:20:B:TYR:HE1	19	2.12
(1,415)	1:6:A:VAL:HG11	2:20:B:TYR:HE2	19	2.12
(1,415)	1:6:A:VAL:HG12	2:20:B:TYR:HE1	19	2.12
(1,415)	1:6:A:VAL:HG12	2:20:B:TYR:HE2	19	2.12
(1,415)	1:6:A:VAL:HG13	2:20:B:TYR:HE1	19	2.12
(1,415)	1:6:A:VAL:HG13	2:20:B:TYR:HE2	19	2.12
(1,96)	1:6:A:VAL:HG11	2:20:B:TYR:HE1	19	2.12
(1,96)	1:6:A:VAL:HG11	2:20:B:TYR:HE2	19	2.12
(1,96)	1:6:A:VAL:HG12	2:20:B:TYR:HE1	19	2.12
(1,96)	1:6:A:VAL:HG12	2:20:B:TYR:HE2	19	2.12
(1,96)	1:6:A:VAL:HG13	2:20:B:TYR:HE1	19	2.12
(1,96)	1:6:A:VAL:HG13	2:20:B:TYR:HE2	19	2.12

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,1176)	2:19:B:PHE:HB3	2:38:D:ILE:HG13	12	2.11
(1,857)	2:19:B:PHE:HB3	2:38:D:ILE:HG13	12	2.11
(1,670)	2:17:B:ILE:HD11	2:40:D:PHE:H	4	2.11
(1,670)	2:17:B:ILE:HD12	2:40:D:PHE:H	4	2.11
(1,670)	2:17:B:ILE:HD13	2:40:D:PHE:H	4	2.11
(1,5)	2:17:B:ILE:HD11	2:40:D:PHE:H	4	2.11
(1,5)	2:17:B:ILE:HD12	2:40:D:PHE:H	4	2.11
(1,5)	2:17:B:ILE:HD13	2:40:D:PHE:H	4	2.11
(1,1073)	2:18:B:SER:HB3	2:38:D:ILE:HD11	14	2.1
(1,1073)	2:18:B:SER:HB3	2:38:D:ILE:HD12	14	2.1
(1,1073)	2:18:B:SER:HB3	2:38:D:ILE:HD13	14	2.1
(1,1058)	1:31:C:PHE:HD1	2:35:D:GLY:HA3	10	2.1
(1,1058)	1:31:C:PHE:HD2	2:35:D:GLY:HA3	10	2.1
(1,754)	2:18:B:SER:HB3	2:38:D:ILE:HD11	14	2.1
(1,754)	2:18:B:SER:HB3	2:38:D:ILE:HD12	14	2.1
(1,754)	2:18:B:SER:HB3	2:38:D:ILE:HD13	14	2.1
(1,739)	1:31:C:PHE:HD1	2:35:D:GLY:HA3	10	2.1
(1,739)	1:31:C:PHE:HD2	2:35:D:GLY:HA3	10	2.1
(1,666)	2:17:B:ILE:HD11	2:40:D:PHE:HD1	14	2.09
(1,666)	2:17:B:ILE:HD11	2:40:D:PHE:HD2	14	2.09
(1,666)	2:17:B:ILE:HD12	2:40:D:PHE:HD1	14	2.09
(1,666)	2:17:B:ILE:HD12	2:40:D:PHE:HD2	14	2.09
(1,666)	2:17:B:ILE:HD13	2:40:D:PHE:HD1	14	2.09
(1,666)	2:17:B:ILE:HD13	2:40:D:PHE:HD2	14	2.09
(1,1)	2:17:B:ILE:HD11	2:40:D:PHE:HD1	14	2.09
(1,1)	2:17:B:ILE:HD11	2:40:D:PHE:HD2	14	2.09
(1,1)	2:17:B:ILE:HD12	2:40:D:PHE:HD1	14	2.09
(1,1)	2:17:B:ILE:HD12	2:40:D:PHE:HD2	14	2.09
(1,1)	2:17:B:ILE:HD13	2:40:D:PHE:HD1	14	2.09
(1,1)	2:17:B:ILE:HD13	2:40:D:PHE:HD2	14	2.09
(1,1251)	1:33:C:ARG:HA	1:33:C:ARG:HE	10	2.08
(1,1119)	1:26:C:HIS:HB2	1:27:C:VAL:HG11	17	2.08
(1,1119)	1:26:C:HIS:HB2	1:27:C:VAL:HG12	17	2.08
(1,1119)	1:26:C:HIS:HB2	1:27:C:VAL:HG13	17	2.08
(1,1002)	1:30:C:TYR:HE1	2:39:D:SER:HB2	16	2.08
(1,1002)	1:30:C:TYR:HE2	2:39:D:SER:HB2	16	2.08
(1,932)	1:33:C:ARG:HA	1:33:C:ARG:HE	10	2.08
(1,800)	1:26:C:HIS:HB2	1:27:C:VAL:HG11	17	2.08
(1,800)	1:26:C:HIS:HB2	1:27:C:VAL:HG12	17	2.08
(1,800)	1:26:C:HIS:HB2	1:27:C:VAL:HG13	17	2.08
(1,683)	1:30:C:TYR:HE1	2:39:D:SER:HB2	16	2.08
(1,683)	1:30:C:TYR:HE2	2:39:D:SER:HB2	16	2.08

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,520)	2:17:B:ILE:HG13	2:40:D:PHE:HB3	9	2.08
(1,491)	1:7:A:PRO:HB2	1:10:A:PHE:HE1	20	2.08
(1,491)	1:7:A:PRO:HB2	1:10:A:PHE:HE2	20	2.08
(1,201)	2:17:B:ILE:HG13	2:40:D:PHE:HB3	9	2.08
(1,172)	1:7:A:PRO:HB2	1:10:A:PHE:HE1	20	2.08
(1,172)	1:7:A:PRO:HB2	1:10:A:PHE:HE2	20	2.08
(1,466)	1:4:A:GLY:HA2	1:5:A:HIS:HD2	12	2.06
(1,399)	1:7:A:PRO:HG2	1:10:A:PHE:HE1	12	2.06
(1,399)	1:7:A:PRO:HG2	1:10:A:PHE:HE2	12	2.06
(1,399)	1:7:A:PRO:HG3	1:10:A:PHE:HE1	12	2.06
(1,399)	1:7:A:PRO:HG3	1:10:A:PHE:HE2	12	2.06
(1,351)	2:16:B:PRO:HA	2:17:B:ILE:HD11	15	2.06
(1,351)	2:16:B:PRO:HA	2:17:B:ILE:HD12	15	2.06
(1,351)	2:16:B:PRO:HA	2:17:B:ILE:HD13	15	2.06
(1,147)	1:4:A:GLY:HA2	1:5:A:HIS:HD2	12	2.06
(1,80)	1:7:A:PRO:HG2	1:10:A:PHE:HE1	12	2.06
(1,80)	1:7:A:PRO:HG2	1:10:A:PHE:HE2	12	2.06
(1,80)	1:7:A:PRO:HG3	1:10:A:PHE:HE1	12	2.06
(1,80)	1:7:A:PRO:HG3	1:10:A:PHE:HE2	12	2.06
(1,32)	2:16:B:PRO:HA	2:17:B:ILE:HD11	15	2.06
(1,32)	2:16:B:PRO:HA	2:17:B:ILE:HD12	15	2.06
(1,32)	2:16:B:PRO:HA	2:17:B:ILE:HD13	15	2.06
(1,1093)	1:29:C:GLU:HG2	2:41:D:TYR:HB2	12	2.05
(1,1093)	1:29:C:GLU:HG2	2:41:D:TYR:HB3	12	2.05
(1,1063)	1:22:C:GLY:H	2:39:D:SER:HA	1	2.05
(1,774)	1:29:C:GLU:HG2	2:41:D:TYR:HB2	12	2.05
(1,774)	1:29:C:GLU:HG2	2:41:D:TYR:HB3	12	2.05
(1,744)	1:22:C:GLY:H	2:39:D:SER:HA	1	2.05
(1,417)	2:17:B:ILE:HD11	2:39:D:SER:HB3	4	2.05
(1,417)	2:17:B:ILE:HD12	2:39:D:SER:HB3	4	2.05
(1,417)	2:17:B:ILE:HD13	2:39:D:SER:HB3	4	2.05
(1,98)	2:17:B:ILE:HD11	2:39:D:SER:HB3	4	2.05
(1,98)	2:17:B:ILE:HD12	2:39:D:SER:HB3	4	2.05
(1,98)	2:17:B:ILE:HD13	2:39:D:SER:HB3	4	2.05
(1,1159)	1:28:C:PRO:HB3	1:31:C:PHE:HE1	16	2.04
(1,1159)	1:28:C:PRO:HB3	1:31:C:PHE:HE2	16	2.04
(1,1012)	1:29:C:GLU:HG2	2:41:D:TYR:HE1	1	2.04
(1,1012)	1:29:C:GLU:HG2	2:41:D:TYR:HE2	1	2.04
(1,840)	1:28:C:PRO:HB3	1:31:C:PHE:HE1	16	2.04
(1,840)	1:28:C:PRO:HB3	1:31:C:PHE:HE2	16	2.04
(1,693)	1:29:C:GLU:HG2	2:41:D:TYR:HE1	1	2.04
(1,693)	1:29:C:GLU:HG2	2:41:D:TYR:HE2	1	2.04

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,520)	2:17:B:ILE:HG13	2:40:D:PHE:HB3	5	2.03
(1,436)	1:8:A:GLU:HG2	2:20:B:TYR:HD1	3	2.03
(1,436)	1:8:A:GLU:HG2	2:20:B:TYR:HD2	3	2.03
(1,415)	1:6:A:VAL:HG11	2:20:B:TYR:HE1	12	2.03
(1,415)	1:6:A:VAL:HG11	2:20:B:TYR:HE2	12	2.03
(1,415)	1:6:A:VAL:HG12	2:20:B:TYR:HE1	12	2.03
(1,415)	1:6:A:VAL:HG12	2:20:B:TYR:HE2	12	2.03
(1,415)	1:6:A:VAL:HG13	2:20:B:TYR:HE1	12	2.03
(1,415)	1:6:A:VAL:HG13	2:20:B:TYR:HE2	12	2.03
(1,201)	2:17:B:ILE:HG13	2:40:D:PHE:HB3	5	2.03
(1,117)	1:8:A:GLU:HG2	2:20:B:TYR:HD1	3	2.03
(1,117)	1:8:A:GLU:HG2	2:20:B:TYR:HD2	3	2.03
(1,96)	1:6:A:VAL:HG11	2:20:B:TYR:HE1	12	2.03
(1,96)	1:6:A:VAL:HG11	2:20:B:TYR:HE2	12	2.03
(1,96)	1:6:A:VAL:HG12	2:20:B:TYR:HE1	12	2.03
(1,96)	1:6:A:VAL:HG12	2:20:B:TYR:HE2	12	2.03
(1,96)	1:6:A:VAL:HG13	2:20:B:TYR:HE1	12	2.03
(1,96)	1:6:A:VAL:HG13	2:20:B:TYR:HE2	12	2.03
(1,1171)	2:36:D:THR:HG21	2:38:D:ILE:HD11	18	2.02
(1,1171)	2:36:D:THR:HG21	2:38:D:ILE:HD12	18	2.02
(1,1171)	2:36:D:THR:HG21	2:38:D:ILE:HD13	18	2.02
(1,1171)	2:36:D:THR:HG22	2:38:D:ILE:HD11	18	2.02
(1,1171)	2:36:D:THR:HG22	2:38:D:ILE:HD12	18	2.02
(1,1171)	2:36:D:THR:HG22	2:38:D:ILE:HD13	18	2.02
(1,1171)	2:36:D:THR:HG23	2:38:D:ILE:HD11	18	2.02
(1,1171)	2:36:D:THR:HG23	2:38:D:ILE:HD12	18	2.02
(1,1171)	2:36:D:THR:HG23	2:38:D:ILE:HD13	18	2.02
(1,852)	2:36:D:THR:HG21	2:38:D:ILE:HD11	18	2.02
(1,852)	2:36:D:THR:HG21	2:38:D:ILE:HD12	18	2.02
(1,852)	2:36:D:THR:HG21	2:38:D:ILE:HD13	18	2.02
(1,852)	2:36:D:THR:HG22	2:38:D:ILE:HD11	18	2.02
(1,852)	2:36:D:THR:HG22	2:38:D:ILE:HD12	18	2.02
(1,852)	2:36:D:THR:HG22	2:38:D:ILE:HD13	18	2.02
(1,852)	2:36:D:THR:HG23	2:38:D:ILE:HD11	18	2.02
(1,852)	2:36:D:THR:HG23	2:38:D:ILE:HD12	18	2.02
(1,852)	2:36:D:THR:HG23	2:38:D:ILE:HD13	18	2.02
(1,466)	1:4:A:GLY:HA2	1:5:A:HIS:HD2	11	2.02
(1,147)	1:4:A:GLY:HA2	1:5:A:HIS:HD2	11	2.02
(1,1176)	2:19:B:PHE:HB3	2:38:D:ILE:HG13	7	2.01
(1,1002)	1:30:C:TYR:HE1	2:39:D:SER:HB2	4	2.01
(1,1002)	1:30:C:TYR:HE2	2:39:D:SER:HB2	4	2.01
(1,857)	2:19:B:PHE:HB3	2:38:D:ILE:HG13	7	2.01

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,683)	1:30:C:TYR:HE1	2:39:D:SER:HB2	4	2.01
(1,683)	1:30:C:TYR:HE2	2:39:D:SER:HB2	4	2.01
(1,596)	1:7:A:PRO:HG2	1:10:A:PHE:HE1	12	2.01
(1,596)	1:7:A:PRO:HG2	1:10:A:PHE:HE2	12	2.01
(1,596)	1:7:A:PRO:HG3	1:10:A:PHE:HE1	12	2.01
(1,596)	1:7:A:PRO:HG3	1:10:A:PHE:HE2	12	2.01
(1,595)	1:12:A:ARG:HA	1:12:A:ARG:HE	1	2.01
(1,277)	1:7:A:PRO:HG2	1:10:A:PHE:HE1	12	2.01
(1,277)	1:7:A:PRO:HG2	1:10:A:PHE:HE2	12	2.01
(1,277)	1:7:A:PRO:HG3	1:10:A:PHE:HE1	12	2.01
(1,277)	1:7:A:PRO:HG3	1:10:A:PHE:HE2	12	2.01
(1,276)	1:12:A:ARG:HA	1:12:A:ARG:HE	1	2.01
(1,673)	2:17:B:ILE:HG13	2:40:D:PHE:HB2	18	2.0
(1,665)	2:19:B:PHE:HE1	2:38:D:ILE:HD11	16	2.0
(1,665)	2:19:B:PHE:HE1	2:38:D:ILE:HD12	16	2.0
(1,665)	2:19:B:PHE:HE1	2:38:D:ILE:HD13	16	2.0
(1,665)	2:19:B:PHE:HE2	2:38:D:ILE:HD11	16	2.0
(1,665)	2:19:B:PHE:HE2	2:38:D:ILE:HD12	16	2.0
(1,665)	2:19:B:PHE:HE2	2:38:D:ILE:HD13	16	2.0
(1,595)	1:12:A:ARG:HA	1:12:A:ARG:HE	14	2.0
(1,276)	1:12:A:ARG:HA	1:12:A:ARG:HE	14	2.0
(1,18)	2:19:B:PHE:HE1	2:38:D:ILE:HD11	16	2.0
(1,18)	2:19:B:PHE:HE1	2:38:D:ILE:HD12	16	2.0
(1,18)	2:19:B:PHE:HE1	2:38:D:ILE:HD13	16	2.0
(1,18)	2:19:B:PHE:HE2	2:38:D:ILE:HD11	16	2.0
(1,18)	2:19:B:PHE:HE2	2:38:D:ILE:HD12	16	2.0
(1,18)	2:19:B:PHE:HE2	2:38:D:ILE:HD13	16	2.0
(1,8)	2:17:B:ILE:HG13	2:40:D:PHE:HB2	18	2.0
(1,1169)	2:36:D:THR:HB	2:38:D:ILE:HD11	9	1.98
(1,1169)	2:36:D:THR:HB	2:38:D:ILE:HD12	9	1.98
(1,1169)	2:36:D:THR:HB	2:38:D:ILE:HD13	9	1.98
(1,850)	2:36:D:THR:HB	2:38:D:ILE:HD11	9	1.98
(1,850)	2:36:D:THR:HB	2:38:D:ILE:HD12	9	1.98
(1,850)	2:36:D:THR:HB	2:38:D:ILE:HD13	9	1.98
(1,664)	2:19:B:PHE:HB2	2:38:D:ILE:HG13	17	1.98
(1,17)	2:19:B:PHE:HB2	2:38:D:ILE:HG13	17	1.98
(1,1159)	1:28:C:PRO:HB3	1:31:C:PHE:HE1	10	1.97
(1,1159)	1:28:C:PRO:HB3	1:31:C:PHE:HE2	10	1.97
(1,840)	1:28:C:PRO:HB3	1:31:C:PHE:HE1	10	1.97
(1,840)	1:28:C:PRO:HB3	1:31:C:PHE:HE2	10	1.97
(1,674)	2:17:B:ILE:HD11	2:40:D:PHE:HE1	15	1.97
(1,674)	2:17:B:ILE:HD11	2:40:D:PHE:HE2	15	1.97

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,674)	2:17:B:ILE:HD12	2:40:D:PHE:HE1	15	1.97
(1,674)	2:17:B:ILE:HD12	2:40:D:PHE:HE2	15	1.97
(1,674)	2:17:B:ILE:HD13	2:40:D:PHE:HE1	15	1.97
(1,674)	2:17:B:ILE:HD13	2:40:D:PHE:HE2	15	1.97
(1,466)	1:4:A:GLY:HA2	1:5:A:HIS:HD2	5	1.97
(1,466)	1:4:A:GLY:HA2	1:5:A:HIS:HD2	18	1.97
(1,420)	1:4:A:GLY:HA3	1:5:A:HIS:HD2	15	1.97
(1,147)	1:4:A:GLY:HA2	1:5:A:HIS:HD2	5	1.97
(1,147)	1:4:A:GLY:HA2	1:5:A:HIS:HD2	18	1.97
(1,101)	1:4:A:GLY:HA3	1:5:A:HIS:HD2	15	1.97
(1,9)	2:17:B:ILE:HD11	2:40:D:PHE:HE1	15	1.97
(1,9)	2:17:B:ILE:HD11	2:40:D:PHE:HE2	15	1.97
(1,9)	2:17:B:ILE:HD12	2:40:D:PHE:HE1	15	1.97
(1,9)	2:17:B:ILE:HD12	2:40:D:PHE:HE2	15	1.97
(1,9)	2:17:B:ILE:HD13	2:40:D:PHE:HE1	15	1.97
(1,9)	2:17:B:ILE:HD13	2:40:D:PHE:HE2	15	1.97
(1,1251)	1:33:C:ARG:HA	1:33:C:ARG:HE	1	1.96
(1,932)	1:33:C:ARG:HA	1:33:C:ARG:HE	1	1.96
(1,674)	2:17:B:ILE:HD11	2:40:D:PHE:HE1	3	1.96
(1,674)	2:17:B:ILE:HD11	2:40:D:PHE:HE2	3	1.96
(1,674)	2:17:B:ILE:HD12	2:40:D:PHE:HE1	3	1.96
(1,674)	2:17:B:ILE:HD12	2:40:D:PHE:HE2	3	1.96
(1,674)	2:17:B:ILE:HD13	2:40:D:PHE:HE1	3	1.96
(1,674)	2:17:B:ILE:HD13	2:40:D:PHE:HE2	3	1.96
(1,466)	1:4:A:GLY:HA2	1:5:A:HIS:HD2	14	1.96
(1,393)	1:7:A:PRO:HG2	1:10:A:PHE:HD1	16	1.96
(1,393)	1:7:A:PRO:HG2	1:10:A:PHE:HD2	16	1.96
(1,393)	1:7:A:PRO:HG3	1:10:A:PHE:HD1	16	1.96
(1,393)	1:7:A:PRO:HG3	1:10:A:PHE:HD2	16	1.96
(1,147)	1:4:A:GLY:HA2	1:5:A:HIS:HD2	14	1.96
(1,74)	1:7:A:PRO:HG2	1:10:A:PHE:HD1	16	1.96
(1,74)	1:7:A:PRO:HG2	1:10:A:PHE:HD2	16	1.96
(1,74)	1:7:A:PRO:HG3	1:10:A:PHE:HD1	16	1.96
(1,74)	1:7:A:PRO:HG3	1:10:A:PHE:HD2	16	1.96
(1,9)	2:17:B:ILE:HD11	2:40:D:PHE:HE1	3	1.96
(1,9)	2:17:B:ILE:HD11	2:40:D:PHE:HE2	3	1.96
(1,9)	2:17:B:ILE:HD12	2:40:D:PHE:HE1	3	1.96
(1,9)	2:17:B:ILE:HD12	2:40:D:PHE:HE2	3	1.96
(1,9)	2:17:B:ILE:HD13	2:40:D:PHE:HE1	3	1.96
(1,9)	2:17:B:ILE:HD13	2:40:D:PHE:HE2	3	1.96
(1,1093)	1:29:C:GLU:HG2	2:41:D:TYR:HB2	2	1.95
(1,1093)	1:29:C:GLU:HG2	2:41:D:TYR:HB3	2	1.95

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,774)	1:29:C:GLU:HG2	2:41:D:TYR:HB2	2	1.95
(1,774)	1:29:C:GLU:HG2	2:41:D:TYR:HB3	2	1.95
(1,1017)	1:31:C:PHE:HE1	2:35:D:GLY:HA3	14	1.94
(1,1017)	1:31:C:PHE:HE2	2:35:D:GLY:HA3	14	1.94
(1,698)	1:31:C:PHE:HE1	2:35:D:GLY:HA3	14	1.94
(1,698)	1:31:C:PHE:HE2	2:35:D:GLY:HA3	14	1.94
(1,666)	2:17:B:ILE:HD11	2:40:D:PHE:HD1	9	1.94
(1,666)	2:17:B:ILE:HD11	2:40:D:PHE:HD2	9	1.94
(1,666)	2:17:B:ILE:HD12	2:40:D:PHE:HD1	9	1.94
(1,666)	2:17:B:ILE:HD12	2:40:D:PHE:HD2	9	1.94
(1,666)	2:17:B:ILE:HD13	2:40:D:PHE:HD1	9	1.94
(1,666)	2:17:B:ILE:HD13	2:40:D:PHE:HD2	9	1.94
(1,661)	2:19:B:PHE:H	2:38:D:ILE:HD11	18	1.94
(1,661)	2:19:B:PHE:H	2:38:D:ILE:HD12	18	1.94
(1,661)	2:19:B:PHE:H	2:38:D:ILE:HD13	18	1.94
(1,660)	1:4:A:GLY:HA2	2:34:D:CYS:HB3	10	1.94
(1,399)	1:7:A:PRO:HG2	1:10:A:PHE:HE1	13	1.94
(1,399)	1:7:A:PRO:HG2	1:10:A:PHE:HE2	13	1.94
(1,399)	1:7:A:PRO:HG3	1:10:A:PHE:HE1	13	1.94
(1,399)	1:7:A:PRO:HG3	1:10:A:PHE:HE2	13	1.94
(1,80)	1:7:A:PRO:HG2	1:10:A:PHE:HE1	13	1.94
(1,80)	1:7:A:PRO:HG2	1:10:A:PHE:HE2	13	1.94
(1,80)	1:7:A:PRO:HG3	1:10:A:PHE:HE1	13	1.94
(1,80)	1:7:A:PRO:HG3	1:10:A:PHE:HE2	13	1.94
(1,14)	2:19:B:PHE:H	2:38:D:ILE:HD11	18	1.94
(1,14)	2:19:B:PHE:H	2:38:D:ILE:HD12	18	1.94
(1,14)	2:19:B:PHE:H	2:38:D:ILE:HD13	18	1.94
(1,13)	1:4:A:GLY:HA2	2:34:D:CYS:HB3	10	1.94
(1,1)	2:17:B:ILE:HD11	2:40:D:PHE:HD1	9	1.94
(1,1)	2:17:B:ILE:HD11	2:40:D:PHE:HD2	9	1.94
(1,1)	2:17:B:ILE:HD12	2:40:D:PHE:HD1	9	1.94
(1,1)	2:17:B:ILE:HD12	2:40:D:PHE:HD2	9	1.94
(1,1)	2:17:B:ILE:HD13	2:40:D:PHE:HD1	9	1.94
(1,1)	2:17:B:ILE:HD13	2:40:D:PHE:HD2	9	1.94
(1,1076)	1:25:C:GLY:HA3	1:26:C:HIS:HD2	5	1.93
(1,757)	1:25:C:GLY:HA3	1:26:C:HIS:HD2	5	1.93
(1,666)	2:17:B:ILE:HD11	2:40:D:PHE:HD1	4	1.93
(1,666)	2:17:B:ILE:HD11	2:40:D:PHE:HD2	4	1.93
(1,666)	2:17:B:ILE:HD12	2:40:D:PHE:HD1	4	1.93
(1,666)	2:17:B:ILE:HD12	2:40:D:PHE:HD2	4	1.93
(1,666)	2:17:B:ILE:HD13	2:40:D:PHE:HD1	4	1.93
(1,666)	2:17:B:ILE:HD13	2:40:D:PHE:HD2	4	1.93

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,433)	1:8:A:GLU:HB2	2:20:B:TYR:HB2	11	1.93
(1,433)	1:8:A:GLU:HB2	2:20:B:TYR:HB3	11	1.93
(1,114)	1:8:A:GLU:HB2	2:20:B:TYR:HB2	11	1.93
(1,114)	1:8:A:GLU:HB2	2:20:B:TYR:HB3	11	1.93
(1,1)	2:17:B:ILE:HD11	2:40:D:PHE:HD1	4	1.93
(1,1)	2:17:B:ILE:HD11	2:40:D:PHE:HD2	4	1.93
(1,1)	2:17:B:ILE:HD12	2:40:D:PHE:HD1	4	1.93
(1,1)	2:17:B:ILE:HD12	2:40:D:PHE:HD2	4	1.93
(1,1)	2:17:B:ILE:HD13	2:40:D:PHE:HD1	4	1.93
(1,1)	2:17:B:ILE:HD13	2:40:D:PHE:HD2	4	1.93
(1,1017)	1:31:C:PHE:HE1	2:35:D:GLY:HA3	19	1.92
(1,1017)	1:31:C:PHE:HE2	2:35:D:GLY:HA3	19	1.92
(1,698)	1:31:C:PHE:HE1	2:35:D:GLY:HA3	19	1.92
(1,698)	1:31:C:PHE:HE2	2:35:D:GLY:HA3	19	1.92
(1,437)	1:8:A:GLU:HG2	2:20:B:TYR:HB2	10	1.92
(1,437)	1:8:A:GLU:HG2	2:20:B:TYR:HB3	10	1.92
(1,390)	1:6:A:VAL:HB	2:20:B:TYR:HE1	9	1.92
(1,390)	1:6:A:VAL:HB	2:20:B:TYR:HE2	9	1.92
(1,118)	1:8:A:GLU:HG2	2:20:B:TYR:HB2	10	1.92
(1,118)	1:8:A:GLU:HG2	2:20:B:TYR:HB3	10	1.92
(1,71)	1:6:A:VAL:HB	2:20:B:TYR:HE1	9	1.92
(1,71)	1:6:A:VAL:HB	2:20:B:TYR:HE2	9	1.92
(1,1015)	1:30:C:TYR:HD1	2:39:D:SER:HA	8	1.9
(1,1015)	1:30:C:TYR:HD2	2:39:D:SER:HA	8	1.9
(1,696)	1:30:C:TYR:HD1	2:39:D:SER:HA	8	1.9
(1,696)	1:30:C:TYR:HD2	2:39:D:SER:HA	8	1.9
(1,664)	2:19:B:PHE:HB2	2:38:D:ILE:HG13	8	1.9
(1,513)	2:15:B:THR:HB	2:17:B:ILE:HD11	12	1.9
(1,513)	2:15:B:THR:HB	2:17:B:ILE:HD12	12	1.9
(1,513)	2:15:B:THR:HB	2:17:B:ILE:HD13	12	1.9
(1,503)	1:7:A:PRO:HB3	1:10:A:PHE:HE1	19	1.9
(1,503)	1:7:A:PRO:HB3	1:10:A:PHE:HE2	19	1.9
(1,437)	1:8:A:GLU:HG2	2:20:B:TYR:HB2	5	1.9
(1,437)	1:8:A:GLU:HG2	2:20:B:TYR:HB3	5	1.9
(1,415)	1:6:A:VAL:HG11	2:20:B:TYR:HE1	16	1.9
(1,415)	1:6:A:VAL:HG11	2:20:B:TYR:HE2	16	1.9
(1,415)	1:6:A:VAL:HG12	2:20:B:TYR:HE1	16	1.9
(1,415)	1:6:A:VAL:HG12	2:20:B:TYR:HE2	16	1.9
(1,415)	1:6:A:VAL:HG13	2:20:B:TYR:HE1	16	1.9
(1,415)	1:6:A:VAL:HG13	2:20:B:TYR:HE2	16	1.9
(1,194)	2:15:B:THR:HB	2:17:B:ILE:HD11	12	1.9
(1,194)	2:15:B:THR:HB	2:17:B:ILE:HD12	12	1.9

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,194)	2:15:B:THR:HB	2:17:B:ILE:HD13	12	1.9
(1,184)	1:7:A:PRO:HB3	1:10:A:PHE:HE1	19	1.9
(1,184)	1:7:A:PRO:HB3	1:10:A:PHE:HE2	19	1.9
(1,118)	1:8:A:GLU:HG2	2:20:B:TYR:HB2	5	1.9
(1,118)	1:8:A:GLU:HG2	2:20:B:TYR:HB3	5	1.9
(1,96)	1:6:A:VAL:HG11	2:20:B:TYR:HE1	16	1.9
(1,96)	1:6:A:VAL:HG11	2:20:B:TYR:HE2	16	1.9
(1,96)	1:6:A:VAL:HG12	2:20:B:TYR:HE1	16	1.9
(1,96)	1:6:A:VAL:HG12	2:20:B:TYR:HE2	16	1.9
(1,96)	1:6:A:VAL:HG13	2:20:B:TYR:HE1	16	1.9
(1,96)	1:6:A:VAL:HG13	2:20:B:TYR:HE2	16	1.9
(1,17)	2:19:B:PHE:HB2	2:38:D:ILE:HG13	8	1.9
(1,1251)	1:33:C:ARG:HA	1:33:C:ARG:HE	3	1.89
(1,1171)	2:36:D:THR:HG21	2:38:D:ILE:HD11	3	1.89
(1,1171)	2:36:D:THR:HG21	2:38:D:ILE:HD12	3	1.89
(1,1171)	2:36:D:THR:HG21	2:38:D:ILE:HD13	3	1.89
(1,1171)	2:36:D:THR:HG22	2:38:D:ILE:HD11	3	1.89
(1,1171)	2:36:D:THR:HG22	2:38:D:ILE:HD12	3	1.89
(1,1171)	2:36:D:THR:HG22	2:38:D:ILE:HD13	3	1.89
(1,1171)	2:36:D:THR:HG23	2:38:D:ILE:HD11	3	1.89
(1,1171)	2:36:D:THR:HG23	2:38:D:ILE:HD12	3	1.89
(1,1171)	2:36:D:THR:HG23	2:38:D:ILE:HD13	3	1.89
(1,1147)	1:28:C:PRO:HB2	1:31:C:PHE:HE1	19	1.89
(1,1147)	1:28:C:PRO:HB2	1:31:C:PHE:HE2	19	1.89
(1,1093)	1:29:C:GLU:HG2	2:41:D:TYR:HB2	13	1.89
(1,1093)	1:29:C:GLU:HG2	2:41:D:TYR:HB3	13	1.89
(1,1055)	1:28:C:PRO:HG2	1:31:C:PHE:HE1	9	1.89
(1,1055)	1:28:C:PRO:HG2	1:31:C:PHE:HE2	9	1.89
(1,1055)	1:28:C:PRO:HG3	1:31:C:PHE:HE1	9	1.89
(1,1055)	1:28:C:PRO:HG3	1:31:C:PHE:HE2	9	1.89
(1,1002)	1:30:C:TYR:HE1	2:39:D:SER:HB2	8	1.89
(1,1002)	1:30:C:TYR:HE2	2:39:D:SER:HB2	8	1.89
(1,932)	1:33:C:ARG:HA	1:33:C:ARG:HE	3	1.89
(1,852)	2:36:D:THR:HG21	2:38:D:ILE:HD11	3	1.89
(1,852)	2:36:D:THR:HG21	2:38:D:ILE:HD12	3	1.89
(1,852)	2:36:D:THR:HG21	2:38:D:ILE:HD13	3	1.89
(1,852)	2:36:D:THR:HG22	2:38:D:ILE:HD11	3	1.89
(1,852)	2:36:D:THR:HG22	2:38:D:ILE:HD12	3	1.89
(1,852)	2:36:D:THR:HG22	2:38:D:ILE:HD13	3	1.89
(1,852)	2:36:D:THR:HG23	2:38:D:ILE:HD11	3	1.89
(1,852)	2:36:D:THR:HG23	2:38:D:ILE:HD12	3	1.89
(1,852)	2:36:D:THR:HG23	2:38:D:ILE:HD13	3	1.89

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,828)	1:28:C:PRO:HB2	1:31:C:PHE:HE1	19	1.89
(1,828)	1:28:C:PRO:HB2	1:31:C:PHE:HE2	19	1.89
(1,774)	1:29:C:GLU:HG2	2:41:D:TYR:HB2	13	1.89
(1,774)	1:29:C:GLU:HG2	2:41:D:TYR:HB3	13	1.89
(1,736)	1:28:C:PRO:HG2	1:31:C:PHE:HE1	9	1.89
(1,736)	1:28:C:PRO:HG2	1:31:C:PHE:HE2	9	1.89
(1,736)	1:28:C:PRO:HG3	1:31:C:PHE:HE1	9	1.89
(1,736)	1:28:C:PRO:HG3	1:31:C:PHE:HE2	9	1.89
(1,683)	1:30:C:TYR:HE1	2:39:D:SER:HB2	8	1.89
(1,683)	1:30:C:TYR:HE2	2:39:D:SER:HB2	8	1.89
(1,596)	1:7:A:PRO:HG2	1:10:A:PHE:HE1	13	1.89
(1,596)	1:7:A:PRO:HG2	1:10:A:PHE:HE2	13	1.89
(1,596)	1:7:A:PRO:HG3	1:10:A:PHE:HE1	13	1.89
(1,596)	1:7:A:PRO:HG3	1:10:A:PHE:HE2	13	1.89
(1,277)	1:7:A:PRO:HG2	1:10:A:PHE:HE1	13	1.89
(1,277)	1:7:A:PRO:HG2	1:10:A:PHE:HE2	13	1.89
(1,277)	1:7:A:PRO:HG3	1:10:A:PHE:HE1	13	1.89
(1,277)	1:7:A:PRO:HG3	1:10:A:PHE:HE2	13	1.89
(1,1002)	1:30:C:TYR:HE1	2:39:D:SER:HB2	14	1.88
(1,1002)	1:30:C:TYR:HE2	2:39:D:SER:HB2	14	1.88
(1,683)	1:30:C:TYR:HE1	2:39:D:SER:HB2	14	1.88
(1,683)	1:30:C:TYR:HE2	2:39:D:SER:HB2	14	1.88
(1,671)	2:17:B:ILE:HG13	2:40:D:PHE:HE1	18	1.88
(1,671)	2:17:B:ILE:HG13	2:40:D:PHE:HE2	18	1.88
(1,415)	1:6:A:VAL:HG11	2:20:B:TYR:HE1	3	1.88
(1,415)	1:6:A:VAL:HG11	2:20:B:TYR:HE2	3	1.88
(1,415)	1:6:A:VAL:HG12	2:20:B:TYR:HE1	3	1.88
(1,415)	1:6:A:VAL:HG12	2:20:B:TYR:HE2	3	1.88
(1,415)	1:6:A:VAL:HG13	2:20:B:TYR:HE1	3	1.88
(1,415)	1:6:A:VAL:HG13	2:20:B:TYR:HE2	3	1.88
(1,96)	1:6:A:VAL:HG11	2:20:B:TYR:HE1	3	1.88
(1,96)	1:6:A:VAL:HG11	2:20:B:TYR:HE2	3	1.88
(1,96)	1:6:A:VAL:HG12	2:20:B:TYR:HE1	3	1.88
(1,96)	1:6:A:VAL:HG12	2:20:B:TYR:HE2	3	1.88
(1,96)	1:6:A:VAL:HG13	2:20:B:TYR:HE1	3	1.88
(1,96)	1:6:A:VAL:HG13	2:20:B:TYR:HE2	3	1.88
(1,6)	2:17:B:ILE:HG13	2:40:D:PHE:HE1	18	1.88
(1,6)	2:17:B:ILE:HG13	2:40:D:PHE:HE2	18	1.88
(1,1169)	2:36:D:THR:HB	2:38:D:ILE:HD11	10	1.87
(1,1169)	2:36:D:THR:HB	2:38:D:ILE:HD12	10	1.87
(1,1169)	2:36:D:THR:HB	2:38:D:ILE:HD13	10	1.87
(1,1137)	1:22:C:GLY:H	2:40:D:PHE:H	1	1.87

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,850)	2:36:D:THR:HB	2:38:D:ILE:HD11	10	1.87
(1,850)	2:36:D:THR:HB	2:38:D:ILE:HD12	10	1.87
(1,850)	2:36:D:THR:HB	2:38:D:ILE:HD13	10	1.87
(1,818)	1:22:C:GLY:H	2:40:D:PHE:H	1	1.87
(1,671)	2:17:B:ILE:HG13	2:40:D:PHE:HE1	14	1.87
(1,671)	2:17:B:ILE:HG13	2:40:D:PHE:HE2	14	1.87
(1,520)	2:17:B:ILE:HG13	2:40:D:PHE:HB3	4	1.87
(1,491)	1:7:A:PRO:HB2	1:10:A:PHE:HE1	11	1.87
(1,491)	1:7:A:PRO:HB2	1:10:A:PHE:HE2	11	1.87
(1,415)	1:6:A:VAL:HG11	2:20:B:TYR:HE1	10	1.87
(1,415)	1:6:A:VAL:HG11	2:20:B:TYR:HE2	10	1.87
(1,415)	1:6:A:VAL:HG12	2:20:B:TYR:HE1	10	1.87
(1,415)	1:6:A:VAL:HG12	2:20:B:TYR:HE2	10	1.87
(1,415)	1:6:A:VAL:HG13	2:20:B:TYR:HE1	10	1.87
(1,415)	1:6:A:VAL:HG13	2:20:B:TYR:HE2	10	1.87
(1,201)	2:17:B:ILE:HG13	2:40:D:PHE:HB3	4	1.87
(1,172)	1:7:A:PRO:HB2	1:10:A:PHE:HE1	11	1.87
(1,172)	1:7:A:PRO:HB2	1:10:A:PHE:HE2	11	1.87
(1,96)	1:6:A:VAL:HG11	2:20:B:TYR:HE1	10	1.87
(1,96)	1:6:A:VAL:HG11	2:20:B:TYR:HE2	10	1.87
(1,96)	1:6:A:VAL:HG12	2:20:B:TYR:HE1	10	1.87
(1,96)	1:6:A:VAL:HG12	2:20:B:TYR:HE2	10	1.87
(1,96)	1:6:A:VAL:HG13	2:20:B:TYR:HE1	10	1.87
(1,96)	1:6:A:VAL:HG13	2:20:B:TYR:HE2	10	1.87
(1,6)	2:17:B:ILE:HG13	2:40:D:PHE:HE1	14	1.87
(1,6)	2:17:B:ILE:HG13	2:40:D:PHE:HE2	14	1.87
(1,1176)	2:19:B:PHE:HB3	2:38:D:ILE:HG13	16	1.86
(1,1076)	1:25:C:GLY:HA3	1:26:C:HIS:HD2	20	1.86
(1,857)	2:19:B:PHE:HB3	2:38:D:ILE:HG13	16	1.86
(1,757)	1:25:C:GLY:HA3	1:26:C:HIS:HD2	20	1.86
(1,520)	2:17:B:ILE:HG13	2:40:D:PHE:HB3	3	1.86
(1,201)	2:17:B:ILE:HG13	2:40:D:PHE:HB3	3	1.86
(1,1122)	1:25:C:GLY:HA2	1:26:C:HIS:HD2	20	1.85
(1,803)	1:25:C:GLY:HA2	1:26:C:HIS:HD2	20	1.85
(1,673)	2:17:B:ILE:HG13	2:40:D:PHE:HB2	9	1.85
(1,436)	1:8:A:GLU:HG2	2:20:B:TYR:HD1	4	1.85
(1,436)	1:8:A:GLU:HG2	2:20:B:TYR:HD2	4	1.85
(1,117)	1:8:A:GLU:HG2	2:20:B:TYR:HD1	4	1.85
(1,117)	1:8:A:GLU:HG2	2:20:B:TYR:HD2	4	1.85
(1,8)	2:17:B:ILE:HG13	2:40:D:PHE:HB2	9	1.85
(1,1252)	1:28:C:PRO:HG2	1:31:C:PHE:HE1	9	1.84
(1,1252)	1:28:C:PRO:HG2	1:31:C:PHE:HE2	9	1.84

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,1252)	1:28:C:PRO:HG3	1:31:C:PHE:HE1	9	1.84
(1,1252)	1:28:C:PRO:HG3	1:31:C:PHE:HE2	9	1.84
(1,1159)	1:28:C:PRO:HB3	1:31:C:PHE:HE1	13	1.84
(1,1159)	1:28:C:PRO:HB3	1:31:C:PHE:HE2	13	1.84
(1,933)	1:28:C:PRO:HG2	1:31:C:PHE:HE1	9	1.84
(1,933)	1:28:C:PRO:HG2	1:31:C:PHE:HE2	9	1.84
(1,933)	1:28:C:PRO:HG3	1:31:C:PHE:HE1	9	1.84
(1,933)	1:28:C:PRO:HG3	1:31:C:PHE:HE2	9	1.84
(1,840)	1:28:C:PRO:HB3	1:31:C:PHE:HE1	13	1.84
(1,840)	1:28:C:PRO:HB3	1:31:C:PHE:HE2	13	1.84
(1,515)	2:15:B:THR:HG21	2:17:B:ILE:HD11	9	1.84
(1,515)	2:15:B:THR:HG21	2:17:B:ILE:HD12	9	1.84
(1,515)	2:15:B:THR:HG21	2:17:B:ILE:HD13	9	1.84
(1,515)	2:15:B:THR:HG22	2:17:B:ILE:HD11	9	1.84
(1,515)	2:15:B:THR:HG22	2:17:B:ILE:HD12	9	1.84
(1,515)	2:15:B:THR:HG22	2:17:B:ILE:HD13	9	1.84
(1,515)	2:15:B:THR:HG23	2:17:B:ILE:HD11	9	1.84
(1,515)	2:15:B:THR:HG23	2:17:B:ILE:HD12	9	1.84
(1,515)	2:15:B:THR:HG23	2:17:B:ILE:HD13	9	1.84
(1,196)	2:15:B:THR:HG21	2:17:B:ILE:HD11	9	1.84
(1,196)	2:15:B:THR:HG21	2:17:B:ILE:HD12	9	1.84
(1,196)	2:15:B:THR:HG21	2:17:B:ILE:HD13	9	1.84
(1,196)	2:15:B:THR:HG22	2:17:B:ILE:HD11	9	1.84
(1,196)	2:15:B:THR:HG22	2:17:B:ILE:HD12	9	1.84
(1,196)	2:15:B:THR:HG22	2:17:B:ILE:HD13	9	1.84
(1,196)	2:15:B:THR:HG23	2:17:B:ILE:HD11	9	1.84
(1,196)	2:15:B:THR:HG23	2:17:B:ILE:HD12	9	1.84
(1,196)	2:15:B:THR:HG23	2:17:B:ILE:HD13	9	1.84
(1,1089)	1:29:C:GLU:HB2	2:41:D:TYR:HB2	7	1.83
(1,1089)	1:29:C:GLU:HB2	2:41:D:TYR:HB3	7	1.83
(1,770)	1:29:C:GLU:HB2	2:41:D:TYR:HB2	7	1.83
(1,770)	1:29:C:GLU:HB2	2:41:D:TYR:HB3	7	1.83
(1,466)	1:4:A:GLY:HA2	1:5:A:HIS:HD2	8	1.83
(1,437)	1:8:A:GLU:HG2	2:20:B:TYR:HB2	19	1.83
(1,437)	1:8:A:GLU:HG2	2:20:B:TYR:HB3	19	1.83
(1,147)	1:4:A:GLY:HA2	1:5:A:HIS:HD2	8	1.83
(1,118)	1:8:A:GLU:HG2	2:20:B:TYR:HB2	19	1.83
(1,118)	1:8:A:GLU:HG2	2:20:B:TYR:HB3	19	1.83
(1,1251)	1:33:C:ARG:HA	1:33:C:ARG:HE	17	1.82
(1,932)	1:33:C:ARG:HA	1:33:C:ARG:HE	17	1.82
(1,669)	2:13:B:CYS:HB3	1:25:C:GLY:HA2	12	1.82
(1,595)	1:12:A:ARG:HA	1:12:A:ARG:HE	6	1.82

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,276)	1:12:A:ARG:HA	1:12:A:ARG:HE	6	1.82
(1,4)	2:13:B:CYS:HB3	1:25:C:GLY:HA2	12	1.82
(1,1093)	1:29:C:GLU:HG2	2:41:D:TYR:HB2	19	1.81
(1,1093)	1:29:C:GLU:HG2	2:41:D:TYR:HB3	19	1.81
(1,1092)	1:29:C:GLU:HG2	2:41:D:TYR:HD1	5	1.81
(1,1092)	1:29:C:GLU:HG2	2:41:D:TYR:HD2	5	1.81
(1,1092)	1:29:C:GLU:HG2	2:41:D:TYR:HD1	13	1.81
(1,1092)	1:29:C:GLU:HG2	2:41:D:TYR:HD2	13	1.81
(1,774)	1:29:C:GLU:HG2	2:41:D:TYR:HB2	19	1.81
(1,774)	1:29:C:GLU:HG2	2:41:D:TYR:HB3	19	1.81
(1,773)	1:29:C:GLU:HG2	2:41:D:TYR:HD1	5	1.81
(1,773)	1:29:C:GLU:HG2	2:41:D:TYR:HD2	5	1.81
(1,773)	1:29:C:GLU:HG2	2:41:D:TYR:HD1	13	1.81
(1,773)	1:29:C:GLU:HG2	2:41:D:TYR:HD2	13	1.81
(1,420)	1:4:A:GLY:HA3	1:5:A:HIS:HD2	11	1.81
(1,101)	1:4:A:GLY:HA3	1:5:A:HIS:HD2	11	1.81
(1,1076)	1:25:C:GLY:HA3	1:26:C:HIS:HD2	1	1.8
(1,757)	1:25:C:GLY:HA3	1:26:C:HIS:HD2	1	1.8
(1,657)	2:19:B:PHE:HD1	2:38:D:ILE:HD11	7	1.8
(1,657)	2:19:B:PHE:HD1	2:38:D:ILE:HD12	7	1.8
(1,657)	2:19:B:PHE:HD1	2:38:D:ILE:HD13	7	1.8
(1,657)	2:19:B:PHE:HD2	2:38:D:ILE:HD11	7	1.8
(1,657)	2:19:B:PHE:HD2	2:38:D:ILE:HD12	7	1.8
(1,657)	2:19:B:PHE:HD2	2:38:D:ILE:HD13	7	1.8
(1,415)	1:6:A:VAL:HG11	2:20:B:TYR:HE1	11	1.8
(1,415)	1:6:A:VAL:HG11	2:20:B:TYR:HE2	11	1.8
(1,415)	1:6:A:VAL:HG12	2:20:B:TYR:HE1	11	1.8
(1,415)	1:6:A:VAL:HG12	2:20:B:TYR:HE2	11	1.8
(1,415)	1:6:A:VAL:HG13	2:20:B:TYR:HE1	11	1.8
(1,415)	1:6:A:VAL:HG13	2:20:B:TYR:HE2	11	1.8
(1,96)	1:6:A:VAL:HG11	2:20:B:TYR:HE1	11	1.8
(1,96)	1:6:A:VAL:HG11	2:20:B:TYR:HE2	11	1.8
(1,96)	1:6:A:VAL:HG12	2:20:B:TYR:HE1	11	1.8
(1,96)	1:6:A:VAL:HG12	2:20:B:TYR:HE2	11	1.8
(1,96)	1:6:A:VAL:HG13	2:20:B:TYR:HE1	11	1.8
(1,96)	1:6:A:VAL:HG13	2:20:B:TYR:HE2	11	1.8
(1,10)	2:19:B:PHE:HD1	2:38:D:ILE:HD11	7	1.8
(1,10)	2:19:B:PHE:HD1	2:38:D:ILE:HD12	7	1.8
(1,10)	2:19:B:PHE:HD1	2:38:D:ILE:HD13	7	1.8
(1,10)	2:19:B:PHE:HD2	2:38:D:ILE:HD11	7	1.8
(1,10)	2:19:B:PHE:HD2	2:38:D:ILE:HD12	7	1.8
(1,10)	2:19:B:PHE:HD2	2:38:D:ILE:HD13	7	1.8

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,1176)	2:19:B:PHE:HB3	2:38:D:ILE:HG13	1	1.79
(1,857)	2:19:B:PHE:HB3	2:38:D:ILE:HG13	1	1.79
(1,420)	1:4:A:GLY:HA3	1:5:A:HIS:HD2	12	1.79
(1,342)	1:1:A:GLY:HA3	2:18:B:SER:HB2	10	1.79
(1,339)	1:1:A:GLY:HA3	2:18:B:SER:HB2	10	1.79
(1,101)	1:4:A:GLY:HA3	1:5:A:HIS:HD2	12	1.79
(1,23)	1:1:A:GLY:HA3	2:18:B:SER:HB2	10	1.79
(1,20)	1:1:A:GLY:HA3	2:18:B:SER:HB2	10	1.79
(1,1015)	1:30:C:TYR:HD1	2:39:D:SER:HA	12	1.78
(1,1015)	1:30:C:TYR:HD2	2:39:D:SER:HA	12	1.78
(1,696)	1:30:C:TYR:HD1	2:39:D:SER:HA	12	1.78
(1,696)	1:30:C:TYR:HD2	2:39:D:SER:HA	12	1.78
(1,520)	2:17:B:ILE:HG13	2:40:D:PHE:HB3	7	1.78
(1,519)	1:3:A:ALA:HB1	2:19:B:PHE:HB3	9	1.78
(1,519)	1:3:A:ALA:HB2	2:19:B:PHE:HB3	9	1.78
(1,519)	1:3:A:ALA:HB3	2:19:B:PHE:HB3	9	1.78
(1,361)	1:10:A:PHE:HE1	2:14:B:GLY:HA3	8	1.78
(1,361)	1:10:A:PHE:HE2	2:14:B:GLY:HA3	8	1.78
(1,201)	2:17:B:ILE:HG13	2:40:D:PHE:HB3	7	1.78
(1,200)	1:3:A:ALA:HB1	2:19:B:PHE:HB3	9	1.78
(1,200)	1:3:A:ALA:HB2	2:19:B:PHE:HB3	9	1.78
(1,200)	1:3:A:ALA:HB3	2:19:B:PHE:HB3	9	1.78
(1,42)	1:10:A:PHE:HE1	2:14:B:GLY:HA3	8	1.78
(1,42)	1:10:A:PHE:HE2	2:14:B:GLY:HA3	8	1.78
(1,1251)	1:33:C:ARG:HA	1:33:C:ARG:HE	6	1.77
(1,932)	1:33:C:ARG:HA	1:33:C:ARG:HE	6	1.77
(1,437)	1:8:A:GLU:HG2	2:20:B:TYR:HB2	4	1.77
(1,437)	1:8:A:GLU:HG2	2:20:B:TYR:HB3	4	1.77
(1,118)	1:8:A:GLU:HG2	2:20:B:TYR:HB2	4	1.77
(1,118)	1:8:A:GLU:HG2	2:20:B:TYR:HB3	4	1.77
(1,662)	2:19:B:PHE:HE1	2:38:D:ILE:HG13	15	1.76
(1,662)	2:19:B:PHE:HE2	2:38:D:ILE:HG13	15	1.76
(1,520)	2:17:B:ILE:HG13	2:40:D:PHE:HB3	11	1.76
(1,402)	1:10:A:PHE:HD1	2:14:B:GLY:HA3	20	1.76
(1,402)	1:10:A:PHE:HD2	2:14:B:GLY:HA3	20	1.76
(1,201)	2:17:B:ILE:HG13	2:40:D:PHE:HB3	11	1.76
(1,83)	1:10:A:PHE:HD1	2:14:B:GLY:HA3	20	1.76
(1,83)	1:10:A:PHE:HD2	2:14:B:GLY:HA3	20	1.76
(1,15)	2:19:B:PHE:HE1	2:38:D:ILE:HG13	15	1.76
(1,15)	2:19:B:PHE:HE2	2:38:D:ILE:HG13	15	1.76
(1,1176)	2:19:B:PHE:HB3	2:38:D:ILE:HG13	18	1.75
(1,1076)	1:25:C:GLY:HA3	1:26:C:HIS:HD2	2	1.75

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,1071)	1:27:C:VAL:HG11	2:41:D:TYR:HE1	16	1.75
(1,1071)	1:27:C:VAL:HG11	2:41:D:TYR:HE2	16	1.75
(1,1071)	1:27:C:VAL:HG12	2:41:D:TYR:HE1	16	1.75
(1,1071)	1:27:C:VAL:HG12	2:41:D:TYR:HE2	16	1.75
(1,1071)	1:27:C:VAL:HG13	2:41:D:TYR:HE1	16	1.75
(1,1071)	1:27:C:VAL:HG13	2:41:D:TYR:HE2	16	1.75
(1,857)	2:19:B:PHE:HB3	2:38:D:ILE:HG13	18	1.75
(1,757)	1:25:C:GLY:HA3	1:26:C:HIS:HD2	2	1.75
(1,752)	1:27:C:VAL:HG11	2:41:D:TYR:HE1	16	1.75
(1,752)	1:27:C:VAL:HG11	2:41:D:TYR:HE2	16	1.75
(1,752)	1:27:C:VAL:HG12	2:41:D:TYR:HE1	16	1.75
(1,752)	1:27:C:VAL:HG12	2:41:D:TYR:HE2	16	1.75
(1,752)	1:27:C:VAL:HG13	2:41:D:TYR:HE1	16	1.75
(1,752)	1:27:C:VAL:HG13	2:41:D:TYR:HE2	16	1.75
(1,657)	2:19:B:PHE:HD1	2:38:D:ILE:HD11	8	1.75
(1,657)	2:19:B:PHE:HD1	2:38:D:ILE:HD12	8	1.75
(1,657)	2:19:B:PHE:HD1	2:38:D:ILE:HD13	8	1.75
(1,657)	2:19:B:PHE:HD2	2:38:D:ILE:HD11	8	1.75
(1,657)	2:19:B:PHE:HD2	2:38:D:ILE:HD12	8	1.75
(1,657)	2:19:B:PHE:HD2	2:38:D:ILE:HD13	8	1.75
(1,420)	1:4:A:GLY:HA3	1:5:A:HIS:HD2	10	1.75
(1,420)	1:4:A:GLY:HA3	1:5:A:HIS:HD2	14	1.75
(1,101)	1:4:A:GLY:HA3	1:5:A:HIS:HD2	10	1.75
(1,101)	1:4:A:GLY:HA3	1:5:A:HIS:HD2	14	1.75
(1,10)	2:19:B:PHE:HD1	2:38:D:ILE:HD11	8	1.75
(1,10)	2:19:B:PHE:HD1	2:38:D:ILE:HD12	8	1.75
(1,10)	2:19:B:PHE:HD1	2:38:D:ILE:HD13	8	1.75
(1,10)	2:19:B:PHE:HD2	2:38:D:ILE:HD11	8	1.75
(1,10)	2:19:B:PHE:HD2	2:38:D:ILE:HD12	8	1.75
(1,10)	2:19:B:PHE:HD2	2:38:D:ILE:HD13	8	1.75
(1,1122)	1:25:C:GLY:HA2	1:26:C:HIS:HD2	2	1.74
(1,1122)	1:25:C:GLY:HA2	1:26:C:HIS:HD2	6	1.74
(1,803)	1:25:C:GLY:HA2	1:26:C:HIS:HD2	2	1.74
(1,803)	1:25:C:GLY:HA2	1:26:C:HIS:HD2	6	1.74
(1,663)	2:19:B:PHE:HD1	2:38:D:ILE:HG13	9	1.74
(1,663)	2:19:B:PHE:HD2	2:38:D:ILE:HG13	9	1.74
(1,662)	2:19:B:PHE:HE1	2:38:D:ILE:HG13	8	1.74
(1,662)	2:19:B:PHE:HE2	2:38:D:ILE:HG13	8	1.74
(1,483)	1:1:A:GLY:H	2:20:B:TYR:HD1	15	1.74
(1,483)	1:1:A:GLY:H	2:20:B:TYR:HD2	15	1.74
(1,415)	1:6:A:VAL:HG11	2:20:B:TYR:HE1	15	1.74
(1,415)	1:6:A:VAL:HG11	2:20:B:TYR:HE2	15	1.74

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,415)	1:6:A:VAL:HG12	2:20:B:TYR:HE1	15	1.74
(1,415)	1:6:A:VAL:HG12	2:20:B:TYR:HE2	15	1.74
(1,415)	1:6:A:VAL:HG13	2:20:B:TYR:HE1	15	1.74
(1,415)	1:6:A:VAL:HG13	2:20:B:TYR:HE2	15	1.74
(1,164)	1:1:A:GLY:H	2:20:B:TYR:HD1	15	1.74
(1,164)	1:1:A:GLY:H	2:20:B:TYR:HD2	15	1.74
(1,96)	1:6:A:VAL:HG11	2:20:B:TYR:HE1	15	1.74
(1,96)	1:6:A:VAL:HG11	2:20:B:TYR:HE2	15	1.74
(1,96)	1:6:A:VAL:HG12	2:20:B:TYR:HE1	15	1.74
(1,96)	1:6:A:VAL:HG12	2:20:B:TYR:HE2	15	1.74
(1,96)	1:6:A:VAL:HG13	2:20:B:TYR:HE1	15	1.74
(1,96)	1:6:A:VAL:HG13	2:20:B:TYR:HE2	15	1.74
(1,16)	2:19:B:PHE:HD1	2:38:D:ILE:HG13	9	1.74
(1,16)	2:19:B:PHE:HD2	2:38:D:ILE:HG13	9	1.74
(1,15)	2:19:B:PHE:HE1	2:38:D:ILE:HG13	8	1.74
(1,15)	2:19:B:PHE:HE2	2:38:D:ILE:HG13	8	1.74
(1,1071)	1:27:C:VAL:HG11	2:41:D:TYR:HE1	5	1.73
(1,1071)	1:27:C:VAL:HG11	2:41:D:TYR:HE2	5	1.73
(1,1071)	1:27:C:VAL:HG12	2:41:D:TYR:HE1	5	1.73
(1,1071)	1:27:C:VAL:HG12	2:41:D:TYR:HE2	5	1.73
(1,1071)	1:27:C:VAL:HG13	2:41:D:TYR:HE1	5	1.73
(1,1071)	1:27:C:VAL:HG13	2:41:D:TYR:HE2	5	1.73
(1,752)	1:27:C:VAL:HG11	2:41:D:TYR:HE1	5	1.73
(1,752)	1:27:C:VAL:HG11	2:41:D:TYR:HE2	5	1.73
(1,752)	1:27:C:VAL:HG12	2:41:D:TYR:HE1	5	1.73
(1,752)	1:27:C:VAL:HG12	2:41:D:TYR:HE2	5	1.73
(1,752)	1:27:C:VAL:HG13	2:41:D:TYR:HE1	5	1.73
(1,752)	1:27:C:VAL:HG13	2:41:D:TYR:HE2	5	1.73
(1,361)	1:10:A:PHE:HE1	2:14:B:GLY:HA3	9	1.73
(1,361)	1:10:A:PHE:HE2	2:14:B:GLY:HA3	9	1.73
(1,344)	1:10:A:PHE:HE1	2:16:B:PRO:HD2	19	1.73
(1,344)	1:10:A:PHE:HE2	2:16:B:PRO:HD2	19	1.73
(1,42)	1:10:A:PHE:HE1	2:14:B:GLY:HA3	9	1.73
(1,42)	1:10:A:PHE:HE2	2:14:B:GLY:HA3	9	1.73
(1,25)	1:10:A:PHE:HE1	2:16:B:PRO:HD2	19	1.73
(1,25)	1:10:A:PHE:HE2	2:16:B:PRO:HD2	19	1.73
(1,1231)	1:31:C:PHE:HE1	2:36:D:THR:HA	20	1.72
(1,1231)	1:31:C:PHE:HE2	2:36:D:THR:HA	20	1.72
(1,1073)	2:18:B:SER:HB3	2:38:D:ILE:HD11	18	1.72
(1,1073)	2:18:B:SER:HB3	2:38:D:ILE:HD12	18	1.72
(1,1073)	2:18:B:SER:HB3	2:38:D:ILE:HD13	18	1.72
(1,912)	1:31:C:PHE:HE1	2:36:D:THR:HA	20	1.72

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,912)	1:31:C:PHE:HE2	2:36:D:THR:HA	20	1.72
(1,754)	2:18:B:SER:HB3	2:38:D:ILE:HD11	18	1.72
(1,754)	2:18:B:SER:HB3	2:38:D:ILE:HD12	18	1.72
(1,754)	2:18:B:SER:HB3	2:38:D:ILE:HD13	18	1.72
(1,436)	1:8:A:GLU:HG2	2:20:B:TYR:HD1	8	1.72
(1,436)	1:8:A:GLU:HG2	2:20:B:TYR:HD2	8	1.72
(1,117)	1:8:A:GLU:HG2	2:20:B:TYR:HD1	8	1.72
(1,117)	1:8:A:GLU:HG2	2:20:B:TYR:HD2	8	1.72
(1,1175)	1:24:C:ALA:HB1	2:40:D:PHE:HB3	16	1.71
(1,1175)	1:24:C:ALA:HB2	2:40:D:PHE:HB3	16	1.71
(1,1175)	1:24:C:ALA:HB3	2:40:D:PHE:HB3	16	1.71
(1,1017)	1:31:C:PHE:HE1	2:35:D:GLY:HA3	11	1.71
(1,1017)	1:31:C:PHE:HE2	2:35:D:GLY:HA3	11	1.71
(1,856)	1:24:C:ALA:HB1	2:40:D:PHE:HB3	16	1.71
(1,856)	1:24:C:ALA:HB2	2:40:D:PHE:HB3	16	1.71
(1,856)	1:24:C:ALA:HB3	2:40:D:PHE:HB3	16	1.71
(1,698)	1:31:C:PHE:HE1	2:35:D:GLY:HA3	11	1.71
(1,698)	1:31:C:PHE:HE2	2:35:D:GLY:HA3	11	1.71
(1,390)	1:6:A:VAL:HB	2:20:B:TYR:HE1	19	1.71
(1,390)	1:6:A:VAL:HB	2:20:B:TYR:HE2	19	1.71
(1,71)	1:6:A:VAL:HB	2:20:B:TYR:HE1	19	1.71
(1,71)	1:6:A:VAL:HB	2:20:B:TYR:HE2	19	1.71
(1,1145)	1:25:C:GLY:H	2:40:D:PHE:HE1	1	1.7
(1,1145)	1:25:C:GLY:H	2:40:D:PHE:HE2	1	1.7
(1,826)	1:25:C:GLY:H	2:40:D:PHE:HE1	1	1.7
(1,826)	1:25:C:GLY:H	2:40:D:PHE:HE2	1	1.7
(1,346)	1:9:A:TYR:HE1	2:18:B:SER:HB2	13	1.7
(1,346)	1:9:A:TYR:HE2	2:18:B:SER:HB2	13	1.7
(1,27)	1:9:A:TYR:HE1	2:18:B:SER:HB2	13	1.7
(1,27)	1:9:A:TYR:HE2	2:18:B:SER:HB2	13	1.7
(1,670)	2:17:B:ILE:HD11	2:40:D:PHE:H	2	1.69
(1,670)	2:17:B:ILE:HD12	2:40:D:PHE:H	2	1.69
(1,670)	2:17:B:ILE:HD13	2:40:D:PHE:H	2	1.69
(1,437)	1:8:A:GLU:HG2	2:20:B:TYR:HB2	3	1.69
(1,437)	1:8:A:GLU:HG2	2:20:B:TYR:HB3	3	1.69
(1,415)	1:6:A:VAL:HG11	2:20:B:TYR:HE1	5	1.69
(1,415)	1:6:A:VAL:HG11	2:20:B:TYR:HE2	5	1.69
(1,415)	1:6:A:VAL:HG12	2:20:B:TYR:HE1	5	1.69
(1,415)	1:6:A:VAL:HG12	2:20:B:TYR:HE2	5	1.69
(1,415)	1:6:A:VAL:HG13	2:20:B:TYR:HE1	5	1.69
(1,415)	1:6:A:VAL:HG13	2:20:B:TYR:HE2	5	1.69
(1,346)	1:9:A:TYR:HE1	2:18:B:SER:HB2	9	1.69

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,346)	1:9:A:TYR:HE2	2:18:B:SER:HB2	9	1.69
(1,118)	1:8:A:GLU:HG2	2:20:B:TYR:HB2	3	1.69
(1,118)	1:8:A:GLU:HG2	2:20:B:TYR:HB3	3	1.69
(1,96)	1:6:A:VAL:HG11	2:20:B:TYR:HE1	5	1.69
(1,96)	1:6:A:VAL:HG11	2:20:B:TYR:HE2	5	1.69
(1,96)	1:6:A:VAL:HG12	2:20:B:TYR:HE1	5	1.69
(1,96)	1:6:A:VAL:HG12	2:20:B:TYR:HE2	5	1.69
(1,96)	1:6:A:VAL:HG13	2:20:B:TYR:HE1	5	1.69
(1,96)	1:6:A:VAL:HG13	2:20:B:TYR:HE2	5	1.69
(1,27)	1:9:A:TYR:HE1	2:18:B:SER:HB2	9	1.69
(1,27)	1:9:A:TYR:HE2	2:18:B:SER:HB2	9	1.69
(1,5)	2:17:B:ILE:HD11	2:40:D:PHE:H	2	1.69
(1,5)	2:17:B:ILE:HD12	2:40:D:PHE:H	2	1.69
(1,5)	2:17:B:ILE:HD13	2:40:D:PHE:H	2	1.69
(1,1169)	2:36:D:THR:HB	2:38:D:ILE:HD11	17	1.68
(1,1169)	2:36:D:THR:HB	2:38:D:ILE:HD12	17	1.68
(1,1169)	2:36:D:THR:HB	2:38:D:ILE:HD13	17	1.68
(1,1061)	1:22:C:GLY:H	2:39:D:SER:HB2	4	1.68
(1,850)	2:36:D:THR:HB	2:38:D:ILE:HD11	17	1.68
(1,850)	2:36:D:THR:HB	2:38:D:ILE:HD12	17	1.68
(1,850)	2:36:D:THR:HB	2:38:D:ILE:HD13	17	1.68
(1,742)	1:22:C:GLY:H	2:39:D:SER:HB2	4	1.68
(1,436)	1:8:A:GLU:HG2	2:20:B:TYR:HD1	19	1.68
(1,436)	1:8:A:GLU:HG2	2:20:B:TYR:HD2	19	1.68
(1,117)	1:8:A:GLU:HG2	2:20:B:TYR:HD1	19	1.68
(1,117)	1:8:A:GLU:HG2	2:20:B:TYR:HD2	19	1.68
(1,1076)	1:25:C:GLY:HA3	1:26:C:HIS:HD2	17	1.67
(1,757)	1:25:C:GLY:HA3	1:26:C:HIS:HD2	17	1.67
(1,399)	1:7:A:PRO:HG2	1:10:A:PHE:HE1	17	1.67
(1,399)	1:7:A:PRO:HG2	1:10:A:PHE:HE2	17	1.67
(1,399)	1:7:A:PRO:HG3	1:10:A:PHE:HE1	17	1.67
(1,399)	1:7:A:PRO:HG3	1:10:A:PHE:HE2	17	1.67
(1,80)	1:7:A:PRO:HG2	1:10:A:PHE:HE1	17	1.67
(1,80)	1:7:A:PRO:HG2	1:10:A:PHE:HE2	17	1.67
(1,80)	1:7:A:PRO:HG3	1:10:A:PHE:HE1	17	1.67
(1,80)	1:7:A:PRO:HG3	1:10:A:PHE:HE2	17	1.67
(1,673)	2:17:B:ILE:HG13	2:40:D:PHE:HB2	14	1.66
(1,8)	2:17:B:ILE:HG13	2:40:D:PHE:HB2	14	1.66
(1,1092)	1:29:C:GLU:HG2	2:41:D:TYR:HD1	16	1.65
(1,1092)	1:29:C:GLU:HG2	2:41:D:TYR:HD2	16	1.65
(1,773)	1:29:C:GLU:HG2	2:41:D:TYR:HD1	16	1.65
(1,773)	1:29:C:GLU:HG2	2:41:D:TYR:HD2	16	1.65

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,673)	2:17:B:ILE:HG13	2:40:D:PHE:HB2	4	1.65
(1,595)	1:12:A:ARG:HA	1:12:A:ARG:HE	13	1.65
(1,491)	1:7:A:PRO:HB2	1:10:A:PHE:HE1	7	1.65
(1,491)	1:7:A:PRO:HB2	1:10:A:PHE:HE2	7	1.65
(1,436)	1:8:A:GLU:HG2	2:20:B:TYR:HD1	9	1.65
(1,436)	1:8:A:GLU:HG2	2:20:B:TYR:HD2	9	1.65
(1,276)	1:12:A:ARG:HA	1:12:A:ARG:HE	13	1.65
(1,172)	1:7:A:PRO:HB2	1:10:A:PHE:HE1	7	1.65
(1,172)	1:7:A:PRO:HB2	1:10:A:PHE:HE2	7	1.65
(1,117)	1:8:A:GLU:HG2	2:20:B:TYR:HD1	9	1.65
(1,117)	1:8:A:GLU:HG2	2:20:B:TYR:HD2	9	1.65
(1,8)	2:17:B:ILE:HG13	2:40:D:PHE:HB2	4	1.65
(1,668)	1:3:A:ALA:HB1	1:32:C:VAL:HA	1	1.64
(1,668)	1:3:A:ALA:HB2	1:32:C:VAL:HA	1	1.64
(1,668)	1:3:A:ALA:HB3	1:32:C:VAL:HA	1	1.64
(1,664)	2:19:B:PHE:HB2	2:38:D:ILE:HG13	7	1.64
(1,402)	1:10:A:PHE:HD1	2:14:B:GLY:HA3	11	1.64
(1,402)	1:10:A:PHE:HD2	2:14:B:GLY:HA3	11	1.64
(1,83)	1:10:A:PHE:HD1	2:14:B:GLY:HA3	11	1.64
(1,83)	1:10:A:PHE:HD2	2:14:B:GLY:HA3	11	1.64
(1,17)	2:19:B:PHE:HB2	2:38:D:ILE:HG13	7	1.64
(1,3)	1:3:A:ALA:HB1	1:32:C:VAL:HA	1	1.64
(1,3)	1:3:A:ALA:HB2	1:32:C:VAL:HA	1	1.64
(1,3)	1:3:A:ALA:HB3	1:32:C:VAL:HA	1	1.64
(1,1176)	2:19:B:PHE:HB3	2:38:D:ILE:HG13	11	1.63
(1,1147)	1:28:C:PRO:HB2	1:31:C:PHE:HE1	16	1.63
(1,1147)	1:28:C:PRO:HB2	1:31:C:PHE:HE2	16	1.63
(1,1122)	1:25:C:GLY:HA2	1:26:C:HIS:HD2	10	1.63
(1,1058)	1:31:C:PHE:HD1	2:35:D:GLY:HA3	3	1.63
(1,1058)	1:31:C:PHE:HD2	2:35:D:GLY:HA3	3	1.63
(1,857)	2:19:B:PHE:HB3	2:38:D:ILE:HG13	11	1.63
(1,828)	1:28:C:PRO:HB2	1:31:C:PHE:HE1	16	1.63
(1,828)	1:28:C:PRO:HB2	1:31:C:PHE:HE2	16	1.63
(1,803)	1:25:C:GLY:HA2	1:26:C:HIS:HD2	10	1.63
(1,739)	1:31:C:PHE:HD1	2:35:D:GLY:HA3	3	1.63
(1,739)	1:31:C:PHE:HD2	2:35:D:GLY:HA3	3	1.63
(1,673)	2:17:B:ILE:HG13	2:40:D:PHE:HB2	5	1.63
(1,664)	2:19:B:PHE:HB2	2:38:D:ILE:HG13	1	1.63
(1,501)	1:4:A:GLY:H	1:6:A:VAL:H	11	1.63
(1,415)	1:6:A:VAL:HG11	2:20:B:TYR:HE1	4	1.63
(1,415)	1:6:A:VAL:HG11	2:20:B:TYR:HE2	4	1.63
(1,415)	1:6:A:VAL:HG12	2:20:B:TYR:HE1	4	1.63

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,415)	1:6:A:VAL:HG12	2:20:B:TYR:HE2	4	1.63
(1,415)	1:6:A:VAL:HG13	2:20:B:TYR:HE1	4	1.63
(1,415)	1:6:A:VAL:HG13	2:20:B:TYR:HE2	4	1.63
(1,182)	1:4:A:GLY:H	1:6:A:VAL:H	11	1.63
(1,96)	1:6:A:VAL:HG11	2:20:B:TYR:HE1	4	1.63
(1,96)	1:6:A:VAL:HG11	2:20:B:TYR:HE2	4	1.63
(1,96)	1:6:A:VAL:HG12	2:20:B:TYR:HE1	4	1.63
(1,96)	1:6:A:VAL:HG12	2:20:B:TYR:HE2	4	1.63
(1,96)	1:6:A:VAL:HG13	2:20:B:TYR:HE1	4	1.63
(1,96)	1:6:A:VAL:HG13	2:20:B:TYR:HE2	4	1.63
(1,17)	2:19:B:PHE:HB2	2:38:D:ILE:HG13	1	1.63
(1,8)	2:17:B:ILE:HG13	2:40:D:PHE:HB2	5	1.63
(1,1007)	2:37:D:PRO:HA	2:38:D:ILE:HD11	10	1.62
(1,1007)	2:37:D:PRO:HA	2:38:D:ILE:HD12	10	1.62
(1,1007)	2:37:D:PRO:HA	2:38:D:ILE:HD13	10	1.62
(1,688)	2:37:D:PRO:HA	2:38:D:ILE:HD11	10	1.62
(1,688)	2:37:D:PRO:HA	2:38:D:ILE:HD12	10	1.62
(1,688)	2:37:D:PRO:HA	2:38:D:ILE:HD13	10	1.62
(1,596)	1:7:A:PRO:HG2	1:10:A:PHE:HE1	17	1.62
(1,596)	1:7:A:PRO:HG2	1:10:A:PHE:HE2	17	1.62
(1,596)	1:7:A:PRO:HG3	1:10:A:PHE:HE1	17	1.62
(1,596)	1:7:A:PRO:HG3	1:10:A:PHE:HE2	17	1.62
(1,595)	1:12:A:ARG:HA	1:12:A:ARG:HE	11	1.62
(1,503)	1:7:A:PRO:HB3	1:10:A:PHE:HE1	11	1.62
(1,503)	1:7:A:PRO:HB3	1:10:A:PHE:HE2	11	1.62
(1,277)	1:7:A:PRO:HG2	1:10:A:PHE:HE1	17	1.62
(1,277)	1:7:A:PRO:HG2	1:10:A:PHE:HE2	17	1.62
(1,277)	1:7:A:PRO:HG3	1:10:A:PHE:HE1	17	1.62
(1,277)	1:7:A:PRO:HG3	1:10:A:PHE:HE2	17	1.62
(1,276)	1:12:A:ARG:HA	1:12:A:ARG:HE	11	1.62
(1,184)	1:7:A:PRO:HB3	1:10:A:PHE:HE1	11	1.62
(1,184)	1:7:A:PRO:HB3	1:10:A:PHE:HE2	11	1.62
(1,674)	2:17:B:ILE:HD11	2:40:D:PHE:HE1	5	1.61
(1,674)	2:17:B:ILE:HD11	2:40:D:PHE:HE2	5	1.61
(1,674)	2:17:B:ILE:HD12	2:40:D:PHE:HE1	5	1.61
(1,674)	2:17:B:ILE:HD12	2:40:D:PHE:HE2	5	1.61
(1,674)	2:17:B:ILE:HD13	2:40:D:PHE:HE1	5	1.61
(1,674)	2:17:B:ILE:HD13	2:40:D:PHE:HE2	5	1.61
(1,491)	1:7:A:PRO:HB2	1:10:A:PHE:HE1	5	1.61
(1,491)	1:7:A:PRO:HB2	1:10:A:PHE:HE2	5	1.61
(1,384)	1:10:A:PHE:HE1	2:16:B:PRO:HD2	19	1.61
(1,384)	1:10:A:PHE:HE2	2:16:B:PRO:HD2	19	1.61

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,172)	1:7:A:PRO:HB2	1:10:A:PHE:HE1	5	1.61
(1,172)	1:7:A:PRO:HB2	1:10:A:PHE:HE2	5	1.61
(1,65)	1:10:A:PHE:HE1	2:16:B:PRO:HD2	19	1.61
(1,65)	1:10:A:PHE:HE2	2:16:B:PRO:HD2	19	1.61
(1,9)	2:17:B:ILE:HD11	2:40:D:PHE:HE1	5	1.61
(1,9)	2:17:B:ILE:HD11	2:40:D:PHE:HE2	5	1.61
(1,9)	2:17:B:ILE:HD12	2:40:D:PHE:HE1	5	1.61
(1,9)	2:17:B:ILE:HD12	2:40:D:PHE:HE2	5	1.61
(1,9)	2:17:B:ILE:HD13	2:40:D:PHE:HE1	5	1.61
(1,9)	2:17:B:ILE:HD13	2:40:D:PHE:HE2	5	1.61
(1,1175)	1:24:C:ALA:HB1	2:40:D:PHE:HB3	15	1.6
(1,1175)	1:24:C:ALA:HB2	2:40:D:PHE:HB3	15	1.6
(1,1175)	1:24:C:ALA:HB3	2:40:D:PHE:HB3	15	1.6
(1,856)	1:24:C:ALA:HB1	2:40:D:PHE:HB3	15	1.6
(1,856)	1:24:C:ALA:HB2	2:40:D:PHE:HB3	15	1.6
(1,856)	1:24:C:ALA:HB3	2:40:D:PHE:HB3	15	1.6
(1,664)	2:19:B:PHE:HB2	2:38:D:ILE:HG13	12	1.6
(1,491)	1:7:A:PRO:HB2	1:10:A:PHE:HE1	4	1.6
(1,491)	1:7:A:PRO:HB2	1:10:A:PHE:HE2	4	1.6
(1,172)	1:7:A:PRO:HB2	1:10:A:PHE:HE1	4	1.6
(1,172)	1:7:A:PRO:HB2	1:10:A:PHE:HE2	4	1.6
(1,17)	2:19:B:PHE:HB2	2:38:D:ILE:HG13	12	1.6
(1,1147)	1:28:C:PRO:HB2	1:31:C:PHE:HE1	4	1.59
(1,1147)	1:28:C:PRO:HB2	1:31:C:PHE:HE2	4	1.59
(1,1076)	1:25:C:GLY:HA3	1:26:C:HIS:HD2	7	1.59
(1,828)	1:28:C:PRO:HB2	1:31:C:PHE:HE1	4	1.59
(1,828)	1:28:C:PRO:HB2	1:31:C:PHE:HE2	4	1.59
(1,757)	1:25:C:GLY:HA3	1:26:C:HIS:HD2	7	1.59
(1,673)	2:17:B:ILE:HG13	2:40:D:PHE:HB2	3	1.59
(1,8)	2:17:B:ILE:HG13	2:40:D:PHE:HB2	3	1.59
(1,1251)	1:33:C:ARG:HA	1:33:C:ARG:HE	20	1.58
(1,1171)	2:36:D:THR:HG21	2:38:D:ILE:HD11	20	1.58
(1,1171)	2:36:D:THR:HG21	2:38:D:ILE:HD12	20	1.58
(1,1171)	2:36:D:THR:HG21	2:38:D:ILE:HD13	20	1.58
(1,1171)	2:36:D:THR:HG22	2:38:D:ILE:HD11	20	1.58
(1,1171)	2:36:D:THR:HG22	2:38:D:ILE:HD12	20	1.58
(1,1171)	2:36:D:THR:HG22	2:38:D:ILE:HD13	20	1.58
(1,1171)	2:36:D:THR:HG23	2:38:D:ILE:HD11	20	1.58
(1,1171)	2:36:D:THR:HG23	2:38:D:ILE:HD12	20	1.58
(1,1171)	2:36:D:THR:HG23	2:38:D:ILE:HD13	20	1.58
(1,1122)	1:25:C:GLY:HA2	1:26:C:HIS:HD2	17	1.58
(1,1046)	1:27:C:VAL:HB	2:41:D:TYR:HE1	2	1.58

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,1046)	1:27:C:VAL:HB	2:41:D:TYR:HE2	2	1.58
(1,932)	1:33:C:ARG:HA	1:33:C:ARG:HE	20	1.58
(1,852)	2:36:D:THR:HG21	2:38:D:ILE:HD11	20	1.58
(1,852)	2:36:D:THR:HG21	2:38:D:ILE:HD12	20	1.58
(1,852)	2:36:D:THR:HG21	2:38:D:ILE:HD13	20	1.58
(1,852)	2:36:D:THR:HG22	2:38:D:ILE:HD11	20	1.58
(1,852)	2:36:D:THR:HG22	2:38:D:ILE:HD12	20	1.58
(1,852)	2:36:D:THR:HG22	2:38:D:ILE:HD13	20	1.58
(1,852)	2:36:D:THR:HG23	2:38:D:ILE:HD11	20	1.58
(1,852)	2:36:D:THR:HG23	2:38:D:ILE:HD12	20	1.58
(1,852)	2:36:D:THR:HG23	2:38:D:ILE:HD13	20	1.58
(1,803)	1:25:C:GLY:HA2	1:26:C:HIS:HD2	17	1.58
(1,727)	1:27:C:VAL:HB	2:41:D:TYR:HE1	2	1.58
(1,727)	1:27:C:VAL:HB	2:41:D:TYR:HE2	2	1.58
(1,466)	1:4:A:GLY:HA2	1:5:A:HIS:HD2	7	1.58
(1,394)	1:7:A:PRO:HG2	1:10:A:PHE:HB2	18	1.58
(1,394)	1:7:A:PRO:HG3	1:10:A:PHE:HB2	18	1.58
(1,360)	1:7:A:PRO:HG2	1:10:A:PHE:HB2	18	1.58
(1,360)	1:7:A:PRO:HG3	1:10:A:PHE:HB2	18	1.58
(1,147)	1:4:A:GLY:HA2	1:5:A:HIS:HD2	7	1.58
(1,75)	1:7:A:PRO:HG2	1:10:A:PHE:HB2	18	1.58
(1,75)	1:7:A:PRO:HG3	1:10:A:PHE:HB2	18	1.58
(1,41)	1:7:A:PRO:HG2	1:10:A:PHE:HB2	18	1.58
(1,41)	1:7:A:PRO:HG3	1:10:A:PHE:HB2	18	1.58
(1,1251)	1:33:C:ARG:HA	1:33:C:ARG:HE	7	1.57
(1,1157)	1:25:C:GLY:H	1:27:C:VAL:H	3	1.57
(1,1147)	1:28:C:PRO:HB2	1:31:C:PHE:HE1	10	1.57
(1,1147)	1:28:C:PRO:HB2	1:31:C:PHE:HE2	10	1.57
(1,1122)	1:25:C:GLY:HA2	1:26:C:HIS:HD2	12	1.57
(1,1073)	2:18:B:SER:HB3	2:38:D:ILE:HD11	13	1.57
(1,1073)	2:18:B:SER:HB3	2:38:D:ILE:HD12	13	1.57
(1,1073)	2:18:B:SER:HB3	2:38:D:ILE:HD13	13	1.57
(1,1071)	1:27:C:VAL:HG11	2:41:D:TYR:HE1	2	1.57
(1,1071)	1:27:C:VAL:HG11	2:41:D:TYR:HE2	2	1.57
(1,1071)	1:27:C:VAL:HG12	2:41:D:TYR:HE1	2	1.57
(1,1071)	1:27:C:VAL:HG12	2:41:D:TYR:HE2	2	1.57
(1,1071)	1:27:C:VAL:HG13	2:41:D:TYR:HE1	2	1.57
(1,1071)	1:27:C:VAL:HG13	2:41:D:TYR:HE2	2	1.57
(1,1004)	1:32:C:VAL:HA	1:33:C:ARG:HD2	18	1.57
(1,1004)	1:32:C:VAL:HA	1:33:C:ARG:HD3	18	1.57
(1,932)	1:33:C:ARG:HA	1:33:C:ARG:HE	7	1.57
(1,838)	1:25:C:GLY:H	1:27:C:VAL:H	3	1.57

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,828)	1:28:C:PRO:HB2	1:31:C:PHE:HE1	10	1.57
(1,828)	1:28:C:PRO:HB2	1:31:C:PHE:HE2	10	1.57
(1,803)	1:25:C:GLY:HA2	1:26:C:HIS:HD2	12	1.57
(1,754)	2:18:B:SER:HB3	2:38:D:ILE:HD11	13	1.57
(1,754)	2:18:B:SER:HB3	2:38:D:ILE:HD12	13	1.57
(1,754)	2:18:B:SER:HB3	2:38:D:ILE:HD13	13	1.57
(1,752)	1:27:C:VAL:HG11	2:41:D:TYR:HE1	2	1.57
(1,752)	1:27:C:VAL:HG11	2:41:D:TYR:HE2	2	1.57
(1,752)	1:27:C:VAL:HG12	2:41:D:TYR:HE1	2	1.57
(1,752)	1:27:C:VAL:HG12	2:41:D:TYR:HE2	2	1.57
(1,752)	1:27:C:VAL:HG13	2:41:D:TYR:HE1	2	1.57
(1,752)	1:27:C:VAL:HG13	2:41:D:TYR:HE2	2	1.57
(1,685)	1:32:C:VAL:HA	1:33:C:ARG:HD2	18	1.57
(1,685)	1:32:C:VAL:HA	1:33:C:ARG:HD3	18	1.57
(1,575)	1:10:A:PHE:HE1	2:15:B:THR:HA	1	1.57
(1,575)	1:10:A:PHE:HE2	2:15:B:THR:HA	1	1.57
(1,465)	1:5:A:HIS:HD2	1:6:A:VAL:HG11	13	1.57
(1,465)	1:5:A:HIS:HD2	1:6:A:VAL:HG12	13	1.57
(1,465)	1:5:A:HIS:HD2	1:6:A:VAL:HG13	13	1.57
(1,346)	1:9:A:TYR:HE1	2:18:B:SER:HB2	7	1.57
(1,346)	1:9:A:TYR:HE2	2:18:B:SER:HB2	7	1.57
(1,256)	1:10:A:PHE:HE1	2:15:B:THR:HA	1	1.57
(1,256)	1:10:A:PHE:HE2	2:15:B:THR:HA	1	1.57
(1,146)	1:5:A:HIS:HD2	1:6:A:VAL:HG11	13	1.57
(1,146)	1:5:A:HIS:HD2	1:6:A:VAL:HG12	13	1.57
(1,146)	1:5:A:HIS:HD2	1:6:A:VAL:HG13	13	1.57
(1,27)	1:9:A:TYR:HE1	2:18:B:SER:HB2	7	1.57
(1,27)	1:9:A:TYR:HE2	2:18:B:SER:HB2	7	1.57
(1,1121)	1:26:C:HIS:HD2	1:27:C:VAL:HG11	18	1.56
(1,1121)	1:26:C:HIS:HD2	1:27:C:VAL:HG12	18	1.56
(1,1121)	1:26:C:HIS:HD2	1:27:C:VAL:HG13	18	1.56
(1,1093)	1:29:C:GLU:HG2	2:41:D:TYR:HB2	8	1.56
(1,1093)	1:29:C:GLU:HG2	2:41:D:TYR:HB3	8	1.56
(1,802)	1:26:C:HIS:HD2	1:27:C:VAL:HG11	18	1.56
(1,802)	1:26:C:HIS:HD2	1:27:C:VAL:HG12	18	1.56
(1,802)	1:26:C:HIS:HD2	1:27:C:VAL:HG13	18	1.56
(1,774)	1:29:C:GLU:HG2	2:41:D:TYR:HB2	8	1.56
(1,774)	1:29:C:GLU:HG2	2:41:D:TYR:HB3	8	1.56
(1,465)	1:5:A:HIS:HD2	1:6:A:VAL:HG11	18	1.56
(1,465)	1:5:A:HIS:HD2	1:6:A:VAL:HG12	18	1.56
(1,465)	1:5:A:HIS:HD2	1:6:A:VAL:HG13	18	1.56
(1,390)	1:6:A:VAL:HB	2:20:B:TYR:HE1	1	1.56

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,390)	1:6:A:VAL:HB	2:20:B:TYR:HE2	1	1.56
(1,146)	1:5:A:HIS:HD2	1:6:A:VAL:HG11	18	1.56
(1,146)	1:5:A:HIS:HD2	1:6:A:VAL:HG12	18	1.56
(1,146)	1:5:A:HIS:HD2	1:6:A:VAL:HG13	18	1.56
(1,71)	1:6:A:VAL:HB	2:20:B:TYR:HE1	1	1.56
(1,71)	1:6:A:VAL:HB	2:20:B:TYR:HE2	1	1.56
(1,1159)	1:28:C:PRO:HB3	1:31:C:PHE:HE1	19	1.54
(1,1159)	1:28:C:PRO:HB3	1:31:C:PHE:HE2	19	1.54
(1,840)	1:28:C:PRO:HB3	1:31:C:PHE:HE1	19	1.54
(1,840)	1:28:C:PRO:HB3	1:31:C:PHE:HE2	19	1.54
(1,664)	2:19:B:PHE:HB2	2:38:D:ILE:HG13	16	1.54
(1,657)	2:19:B:PHE:HD1	2:38:D:ILE:HD11	6	1.54
(1,657)	2:19:B:PHE:HD1	2:38:D:ILE:HD12	6	1.54
(1,657)	2:19:B:PHE:HD1	2:38:D:ILE:HD13	6	1.54
(1,657)	2:19:B:PHE:HD2	2:38:D:ILE:HD11	6	1.54
(1,657)	2:19:B:PHE:HD2	2:38:D:ILE:HD12	6	1.54
(1,657)	2:19:B:PHE:HD2	2:38:D:ILE:HD13	6	1.54
(1,415)	1:6:A:VAL:HG11	2:20:B:TYR:HE1	2	1.54
(1,415)	1:6:A:VAL:HG11	2:20:B:TYR:HE2	2	1.54
(1,415)	1:6:A:VAL:HG12	2:20:B:TYR:HE1	2	1.54
(1,415)	1:6:A:VAL:HG12	2:20:B:TYR:HE2	2	1.54
(1,415)	1:6:A:VAL:HG13	2:20:B:TYR:HE1	2	1.54
(1,415)	1:6:A:VAL:HG13	2:20:B:TYR:HE2	2	1.54
(1,96)	1:6:A:VAL:HG11	2:20:B:TYR:HE1	2	1.54
(1,96)	1:6:A:VAL:HG11	2:20:B:TYR:HE2	2	1.54
(1,96)	1:6:A:VAL:HG12	2:20:B:TYR:HE1	2	1.54
(1,96)	1:6:A:VAL:HG12	2:20:B:TYR:HE2	2	1.54
(1,96)	1:6:A:VAL:HG13	2:20:B:TYR:HE1	2	1.54
(1,96)	1:6:A:VAL:HG13	2:20:B:TYR:HE2	2	1.54
(1,17)	2:19:B:PHE:HB2	2:38:D:ILE:HG13	16	1.54
(1,10)	2:19:B:PHE:HD1	2:38:D:ILE:HD11	6	1.54
(1,10)	2:19:B:PHE:HD1	2:38:D:ILE:HD12	6	1.54
(1,10)	2:19:B:PHE:HD1	2:38:D:ILE:HD13	6	1.54
(1,10)	2:19:B:PHE:HD2	2:38:D:ILE:HD11	6	1.54
(1,10)	2:19:B:PHE:HD2	2:38:D:ILE:HD12	6	1.54
(1,10)	2:19:B:PHE:HD2	2:38:D:ILE:HD13	6	1.54
(1,1171)	2:36:D:THR:HG21	2:38:D:ILE:HD11	12	1.53
(1,1171)	2:36:D:THR:HG21	2:38:D:ILE:HD12	12	1.53
(1,1171)	2:36:D:THR:HG21	2:38:D:ILE:HD13	12	1.53
(1,1171)	2:36:D:THR:HG22	2:38:D:ILE:HD11	12	1.53
(1,1171)	2:36:D:THR:HG22	2:38:D:ILE:HD12	12	1.53
(1,1171)	2:36:D:THR:HG22	2:38:D:ILE:HD13	12	1.53

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,1171)	2:36:D:THR:HG23	2:38:D:ILE:HD11	12	1.53
(1,1171)	2:36:D:THR:HG23	2:38:D:ILE:HD12	12	1.53
(1,1171)	2:36:D:THR:HG23	2:38:D:ILE:HD13	12	1.53
(1,1009)	1:27:C:VAL:HB	2:41:D:TYR:HB2	3	1.53
(1,1009)	1:27:C:VAL:HB	2:41:D:TYR:HB3	3	1.53
(1,852)	2:36:D:THR:HG21	2:38:D:ILE:HD11	12	1.53
(1,852)	2:36:D:THR:HG21	2:38:D:ILE:HD12	12	1.53
(1,852)	2:36:D:THR:HG21	2:38:D:ILE:HD13	12	1.53
(1,852)	2:36:D:THR:HG22	2:38:D:ILE:HD11	12	1.53
(1,852)	2:36:D:THR:HG22	2:38:D:ILE:HD12	12	1.53
(1,852)	2:36:D:THR:HG22	2:38:D:ILE:HD13	12	1.53
(1,852)	2:36:D:THR:HG23	2:38:D:ILE:HD11	12	1.53
(1,852)	2:36:D:THR:HG23	2:38:D:ILE:HD12	12	1.53
(1,852)	2:36:D:THR:HG23	2:38:D:ILE:HD13	12	1.53
(1,690)	1:27:C:VAL:HB	2:41:D:TYR:HB2	3	1.53
(1,690)	1:27:C:VAL:HB	2:41:D:TYR:HB3	3	1.53
(1,670)	2:17:B:ILE:HD11	2:40:D:PHE:H	8	1.53
(1,670)	2:17:B:ILE:HD12	2:40:D:PHE:H	8	1.53
(1,670)	2:17:B:ILE:HD13	2:40:D:PHE:H	8	1.53
(1,503)	1:7:A:PRO:HB3	1:10:A:PHE:HE1	7	1.53
(1,503)	1:7:A:PRO:HB3	1:10:A:PHE:HE2	7	1.53
(1,420)	1:4:A:GLY:HA3	1:5:A:HIS:HD2	7	1.53
(1,184)	1:7:A:PRO:HB3	1:10:A:PHE:HE1	7	1.53
(1,184)	1:7:A:PRO:HB3	1:10:A:PHE:HE2	7	1.53
(1,101)	1:4:A:GLY:HA3	1:5:A:HIS:HD2	7	1.53
(1,5)	2:17:B:ILE:HD11	2:40:D:PHE:H	8	1.53
(1,5)	2:17:B:ILE:HD12	2:40:D:PHE:H	8	1.53
(1,5)	2:17:B:ILE:HD13	2:40:D:PHE:H	8	1.53
(1,1171)	2:36:D:THR:HG21	2:38:D:ILE:HD11	14	1.52
(1,1171)	2:36:D:THR:HG21	2:38:D:ILE:HD12	14	1.52
(1,1171)	2:36:D:THR:HG21	2:38:D:ILE:HD13	14	1.52
(1,1171)	2:36:D:THR:HG22	2:38:D:ILE:HD11	14	1.52
(1,1171)	2:36:D:THR:HG22	2:38:D:ILE:HD12	14	1.52
(1,1171)	2:36:D:THR:HG22	2:38:D:ILE:HD13	14	1.52
(1,1171)	2:36:D:THR:HG23	2:38:D:ILE:HD11	14	1.52
(1,1171)	2:36:D:THR:HG23	2:38:D:ILE:HD12	14	1.52
(1,1171)	2:36:D:THR:HG23	2:38:D:ILE:HD13	14	1.52
(1,1093)	1:29:C:GLU:HG2	2:41:D:TYR:HB2	16	1.52
(1,1093)	1:29:C:GLU:HG2	2:41:D:TYR:HB3	16	1.52
(1,1076)	1:25:C:GLY:HA3	1:26:C:HIS:HD2	10	1.52
(1,852)	2:36:D:THR:HG21	2:38:D:ILE:HD11	14	1.52
(1,852)	2:36:D:THR:HG21	2:38:D:ILE:HD12	14	1.52

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,852)	2:36:D:THR:HG21	2:38:D:ILE:HD13	14	1.52
(1,852)	2:36:D:THR:HG22	2:38:D:ILE:HD11	14	1.52
(1,852)	2:36:D:THR:HG22	2:38:D:ILE:HD12	14	1.52
(1,852)	2:36:D:THR:HG22	2:38:D:ILE:HD13	14	1.52
(1,852)	2:36:D:THR:HG23	2:38:D:ILE:HD11	14	1.52
(1,852)	2:36:D:THR:HG23	2:38:D:ILE:HD12	14	1.52
(1,852)	2:36:D:THR:HG23	2:38:D:ILE:HD13	14	1.52
(1,774)	1:29:C:GLU:HG2	2:41:D:TYR:HB2	16	1.52
(1,774)	1:29:C:GLU:HG2	2:41:D:TYR:HB3	16	1.52
(1,757)	1:25:C:GLY:HA3	1:26:C:HIS:HD2	10	1.52
(1,513)	2:15:B:THR:HB	2:17:B:ILE:HD11	7	1.52
(1,513)	2:15:B:THR:HB	2:17:B:ILE:HD12	7	1.52
(1,513)	2:15:B:THR:HB	2:17:B:ILE:HD13	7	1.52
(1,466)	1:4:A:GLY:HA2	1:5:A:HIS:HD2	15	1.52
(1,436)	1:8:A:GLU:HG2	2:20:B:TYR:HD1	2	1.52
(1,436)	1:8:A:GLU:HG2	2:20:B:TYR:HD2	2	1.52
(1,344)	1:10:A:PHE:HE1	2:16:B:PRO:HD2	1	1.52
(1,344)	1:10:A:PHE:HE2	2:16:B:PRO:HD2	1	1.52
(1,194)	2:15:B:THR:HB	2:17:B:ILE:HD11	7	1.52
(1,194)	2:15:B:THR:HB	2:17:B:ILE:HD12	7	1.52
(1,194)	2:15:B:THR:HB	2:17:B:ILE:HD13	7	1.52
(1,147)	1:4:A:GLY:HA2	1:5:A:HIS:HD2	15	1.52
(1,117)	1:8:A:GLU:HG2	2:20:B:TYR:HD1	2	1.52
(1,117)	1:8:A:GLU:HG2	2:20:B:TYR:HD2	2	1.52
(1,25)	1:10:A:PHE:HE1	2:16:B:PRO:HD2	1	1.52
(1,25)	1:10:A:PHE:HE2	2:16:B:PRO:HD2	1	1.52
(1,1121)	1:26:C:HIS:HD2	1:27:C:VAL:HG11	17	1.51
(1,1121)	1:26:C:HIS:HD2	1:27:C:VAL:HG12	17	1.51
(1,1121)	1:26:C:HIS:HD2	1:27:C:VAL:HG13	17	1.51
(1,1061)	1:22:C:GLY:H	2:39:D:SER:HB2	9	1.51
(1,1058)	1:31:C:PHE:HD1	2:35:D:GLY:HA3	9	1.51
(1,1058)	1:31:C:PHE:HD2	2:35:D:GLY:HA3	9	1.51
(1,1046)	1:27:C:VAL:HB	2:41:D:TYR:HE1	7	1.51
(1,1046)	1:27:C:VAL:HB	2:41:D:TYR:HE2	7	1.51
(1,1007)	2:37:D:PRO:HA	2:38:D:ILE:HD11	20	1.51
(1,1007)	2:37:D:PRO:HA	2:38:D:ILE:HD12	20	1.51
(1,1007)	2:37:D:PRO:HA	2:38:D:ILE:HD13	20	1.51
(1,802)	1:26:C:HIS:HD2	1:27:C:VAL:HG11	17	1.51
(1,802)	1:26:C:HIS:HD2	1:27:C:VAL:HG12	17	1.51
(1,802)	1:26:C:HIS:HD2	1:27:C:VAL:HG13	17	1.51
(1,742)	1:22:C:GLY:H	2:39:D:SER:HB2	9	1.51
(1,739)	1:31:C:PHE:HD1	2:35:D:GLY:HA3	9	1.51

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,739)	1:31:C:PHE:HD2	2:35:D:GLY:HA3	9	1.51
(1,727)	1:27:C:VAL:HB	2:41:D:TYR:HE1	7	1.51
(1,727)	1:27:C:VAL:HB	2:41:D:TYR:HE2	7	1.51
(1,688)	2:37:D:PRO:HA	2:38:D:ILE:HD11	20	1.51
(1,688)	2:37:D:PRO:HA	2:38:D:ILE:HD12	20	1.51
(1,688)	2:37:D:PRO:HA	2:38:D:ILE:HD13	20	1.51
(1,491)	1:7:A:PRO:HB2	1:10:A:PHE:HE1	6	1.51
(1,491)	1:7:A:PRO:HB2	1:10:A:PHE:HE2	6	1.51
(1,172)	1:7:A:PRO:HB2	1:10:A:PHE:HE1	6	1.51
(1,172)	1:7:A:PRO:HB2	1:10:A:PHE:HE2	6	1.51
(1,1071)	1:27:C:VAL:HG11	2:41:D:TYR:HE1	7	1.5
(1,1071)	1:27:C:VAL:HG11	2:41:D:TYR:HE2	7	1.5
(1,1071)	1:27:C:VAL:HG12	2:41:D:TYR:HE1	7	1.5
(1,1071)	1:27:C:VAL:HG12	2:41:D:TYR:HE2	7	1.5
(1,1071)	1:27:C:VAL:HG13	2:41:D:TYR:HE1	7	1.5
(1,1071)	1:27:C:VAL:HG13	2:41:D:TYR:HE2	7	1.5
(1,1058)	1:31:C:PHE:HD1	2:35:D:GLY:HA3	13	1.5
(1,1058)	1:31:C:PHE:HD2	2:35:D:GLY:HA3	13	1.5
(1,752)	1:27:C:VAL:HG11	2:41:D:TYR:HE1	7	1.5
(1,752)	1:27:C:VAL:HG11	2:41:D:TYR:HE2	7	1.5
(1,752)	1:27:C:VAL:HG12	2:41:D:TYR:HE1	7	1.5
(1,752)	1:27:C:VAL:HG12	2:41:D:TYR:HE2	7	1.5
(1,752)	1:27:C:VAL:HG13	2:41:D:TYR:HE1	7	1.5
(1,752)	1:27:C:VAL:HG13	2:41:D:TYR:HE2	7	1.5
(1,739)	1:31:C:PHE:HD1	2:35:D:GLY:HA3	13	1.5
(1,739)	1:31:C:PHE:HD2	2:35:D:GLY:HA3	13	1.5
(1,1251)	1:33:C:ARG:HA	1:33:C:ARG:HE	8	1.49
(1,1159)	1:28:C:PRO:HB3	1:31:C:PHE:HE1	4	1.49
(1,1159)	1:28:C:PRO:HB3	1:31:C:PHE:HE2	4	1.49
(1,1076)	1:25:C:GLY:HA3	1:26:C:HIS:HD2	11	1.49
(1,932)	1:33:C:ARG:HA	1:33:C:ARG:HE	8	1.49
(1,840)	1:28:C:PRO:HB3	1:31:C:PHE:HE1	4	1.49
(1,840)	1:28:C:PRO:HB3	1:31:C:PHE:HE2	4	1.49
(1,757)	1:25:C:GLY:HA3	1:26:C:HIS:HD2	11	1.49
(1,1251)	1:33:C:ARG:HA	1:33:C:ARG:HE	4	1.48
(1,1147)	1:28:C:PRO:HB2	1:31:C:PHE:HE1	15	1.48
(1,1147)	1:28:C:PRO:HB2	1:31:C:PHE:HE2	15	1.48
(1,932)	1:33:C:ARG:HA	1:33:C:ARG:HE	4	1.48
(1,828)	1:28:C:PRO:HB2	1:31:C:PHE:HE1	15	1.48
(1,828)	1:28:C:PRO:HB2	1:31:C:PHE:HE2	15	1.48
(1,664)	2:19:B:PHE:HB2	2:38:D:ILE:HG13	11	1.48
(1,483)	1:1:A:GLY:H	2:20:B:TYR:HD1	6	1.48

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,483)	1:1:A:GLY:H	2:20:B:TYR:HD2	6	1.48
(1,164)	1:1:A:GLY:H	2:20:B:TYR:HD1	6	1.48
(1,164)	1:1:A:GLY:H	2:20:B:TYR:HD2	6	1.48
(1,17)	2:19:B:PHE:HB2	2:38:D:ILE:HG13	11	1.48
(1,1015)	1:30:C:TYR:HD1	2:39:D:SER:HA	4	1.47
(1,1015)	1:30:C:TYR:HD2	2:39:D:SER:HA	4	1.47
(1,1002)	1:30:C:TYR:HE1	2:39:D:SER:HB2	13	1.47
(1,1002)	1:30:C:TYR:HE2	2:39:D:SER:HB2	13	1.47
(1,1002)	1:30:C:TYR:HE1	2:39:D:SER:HB2	17	1.47
(1,1002)	1:30:C:TYR:HE2	2:39:D:SER:HB2	17	1.47
(1,696)	1:30:C:TYR:HD1	2:39:D:SER:HA	4	1.47
(1,696)	1:30:C:TYR:HD2	2:39:D:SER:HA	4	1.47
(1,683)	1:30:C:TYR:HE1	2:39:D:SER:HB2	13	1.47
(1,683)	1:30:C:TYR:HE2	2:39:D:SER:HB2	13	1.47
(1,683)	1:30:C:TYR:HE1	2:39:D:SER:HB2	17	1.47
(1,683)	1:30:C:TYR:HE2	2:39:D:SER:HB2	17	1.47
(1,673)	2:17:B:ILE:HG13	2:40:D:PHE:HB2	11	1.47
(1,595)	1:12:A:ARG:HA	1:12:A:ARG:HE	4	1.47
(1,483)	1:1:A:GLY:H	2:20:B:TYR:HD1	12	1.47
(1,483)	1:1:A:GLY:H	2:20:B:TYR:HD2	12	1.47
(1,443)	1:8:A:GLU:HG3	2:20:B:TYR:HD1	13	1.47
(1,443)	1:8:A:GLU:HG3	2:20:B:TYR:HD2	13	1.47
(1,276)	1:12:A:ARG:HA	1:12:A:ARG:HE	4	1.47
(1,164)	1:1:A:GLY:H	2:20:B:TYR:HD1	12	1.47
(1,164)	1:1:A:GLY:H	2:20:B:TYR:HD2	12	1.47
(1,124)	1:8:A:GLU:HG3	2:20:B:TYR:HD1	13	1.47
(1,124)	1:8:A:GLU:HG3	2:20:B:TYR:HD2	13	1.47
(1,8)	2:17:B:ILE:HG13	2:40:D:PHE:HB2	11	1.47
(1,1093)	1:29:C:GLU:HG2	2:41:D:TYR:HB2	3	1.46
(1,1093)	1:29:C:GLU:HG2	2:41:D:TYR:HB3	3	1.46
(1,1071)	1:27:C:VAL:HG11	2:41:D:TYR:HE1	8	1.46
(1,1071)	1:27:C:VAL:HG11	2:41:D:TYR:HE2	8	1.46
(1,1071)	1:27:C:VAL:HG12	2:41:D:TYR:HE1	8	1.46
(1,1071)	1:27:C:VAL:HG12	2:41:D:TYR:HE2	8	1.46
(1,1071)	1:27:C:VAL:HG13	2:41:D:TYR:HE1	8	1.46
(1,1071)	1:27:C:VAL:HG13	2:41:D:TYR:HE2	8	1.46
(1,774)	1:29:C:GLU:HG2	2:41:D:TYR:HB2	3	1.46
(1,774)	1:29:C:GLU:HG2	2:41:D:TYR:HB3	3	1.46
(1,752)	1:27:C:VAL:HG11	2:41:D:TYR:HE1	8	1.46
(1,752)	1:27:C:VAL:HG11	2:41:D:TYR:HE2	8	1.46
(1,752)	1:27:C:VAL:HG12	2:41:D:TYR:HE1	8	1.46
(1,752)	1:27:C:VAL:HG12	2:41:D:TYR:HE2	8	1.46

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,752)	1:27:C:VAL:HG13	2:41:D:TYR:HE1	8	1.46
(1,752)	1:27:C:VAL:HG13	2:41:D:TYR:HE2	8	1.46
(1,437)	1:8:A:GLU:HG2	2:20:B:TYR:HB2	8	1.46
(1,437)	1:8:A:GLU:HG2	2:20:B:TYR:HB3	8	1.46
(1,436)	1:8:A:GLU:HG2	2:20:B:TYR:HD1	17	1.46
(1,436)	1:8:A:GLU:HG2	2:20:B:TYR:HD2	17	1.46
(1,417)	2:17:B:ILE:HD11	2:39:D:SER:HB3	7	1.46
(1,417)	2:17:B:ILE:HD12	2:39:D:SER:HB3	7	1.46
(1,417)	2:17:B:ILE:HD13	2:39:D:SER:HB3	7	1.46
(1,390)	1:6:A:VAL:HB	2:20:B:TYR:HE1	8	1.46
(1,390)	1:6:A:VAL:HB	2:20:B:TYR:HE2	8	1.46
(1,118)	1:8:A:GLU:HG2	2:20:B:TYR:HB2	8	1.46
(1,118)	1:8:A:GLU:HG2	2:20:B:TYR:HB3	8	1.46
(1,117)	1:8:A:GLU:HG2	2:20:B:TYR:HD1	17	1.46
(1,117)	1:8:A:GLU:HG2	2:20:B:TYR:HD2	17	1.46
(1,98)	2:17:B:ILE:HD11	2:39:D:SER:HB3	7	1.46
(1,98)	2:17:B:ILE:HD12	2:39:D:SER:HB3	7	1.46
(1,98)	2:17:B:ILE:HD13	2:39:D:SER:HB3	7	1.46
(1,71)	1:6:A:VAL:HB	2:20:B:TYR:HE1	8	1.46
(1,71)	1:6:A:VAL:HB	2:20:B:TYR:HE2	8	1.46
(1,1171)	2:36:D:THR:HG21	2:38:D:ILE:HD11	19	1.45
(1,1171)	2:36:D:THR:HG21	2:38:D:ILE:HD12	19	1.45
(1,1171)	2:36:D:THR:HG21	2:38:D:ILE:HD13	19	1.45
(1,1171)	2:36:D:THR:HG22	2:38:D:ILE:HD11	19	1.45
(1,1171)	2:36:D:THR:HG22	2:38:D:ILE:HD12	19	1.45
(1,1171)	2:36:D:THR:HG22	2:38:D:ILE:HD13	19	1.45
(1,1171)	2:36:D:THR:HG23	2:38:D:ILE:HD11	19	1.45
(1,1171)	2:36:D:THR:HG23	2:38:D:ILE:HD12	19	1.45
(1,1171)	2:36:D:THR:HG23	2:38:D:ILE:HD13	19	1.45
(1,1076)	1:25:C:GLY:HA3	1:26:C:HIS:HD2	15	1.45
(1,852)	2:36:D:THR:HG21	2:38:D:ILE:HD11	19	1.45
(1,852)	2:36:D:THR:HG21	2:38:D:ILE:HD12	19	1.45
(1,852)	2:36:D:THR:HG21	2:38:D:ILE:HD13	19	1.45
(1,852)	2:36:D:THR:HG22	2:38:D:ILE:HD11	19	1.45
(1,852)	2:36:D:THR:HG22	2:38:D:ILE:HD12	19	1.45
(1,852)	2:36:D:THR:HG22	2:38:D:ILE:HD13	19	1.45
(1,852)	2:36:D:THR:HG23	2:38:D:ILE:HD11	19	1.45
(1,852)	2:36:D:THR:HG23	2:38:D:ILE:HD12	19	1.45
(1,852)	2:36:D:THR:HG23	2:38:D:ILE:HD13	19	1.45
(1,757)	1:25:C:GLY:HA3	1:26:C:HIS:HD2	15	1.45
(1,491)	1:7:A:PRO:HB2	1:10:A:PHE:HE1	8	1.45
(1,491)	1:7:A:PRO:HB2	1:10:A:PHE:HE2	8	1.45

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,172)	1:7:A:PRO:HB2	1:10:A:PHE:HE1	8	1.45
(1,172)	1:7:A:PRO:HB2	1:10:A:PHE:HE2	8	1.45
(1,666)	2:17:B:ILE:HD11	2:40:D:PHE:HD1	8	1.44
(1,666)	2:17:B:ILE:HD11	2:40:D:PHE:HD2	8	1.44
(1,666)	2:17:B:ILE:HD12	2:40:D:PHE:HD1	8	1.44
(1,666)	2:17:B:ILE:HD12	2:40:D:PHE:HD2	8	1.44
(1,666)	2:17:B:ILE:HD13	2:40:D:PHE:HD1	8	1.44
(1,666)	2:17:B:ILE:HD13	2:40:D:PHE:HD2	8	1.44
(1,662)	2:19:B:PHE:HE1	2:38:D:ILE:HG13	4	1.44
(1,662)	2:19:B:PHE:HE2	2:38:D:ILE:HG13	4	1.44
(1,515)	2:15:B:THR:HG21	2:17:B:ILE:HD11	2	1.44
(1,515)	2:15:B:THR:HG21	2:17:B:ILE:HD12	2	1.44
(1,515)	2:15:B:THR:HG21	2:17:B:ILE:HD13	2	1.44
(1,515)	2:15:B:THR:HG22	2:17:B:ILE:HD11	2	1.44
(1,515)	2:15:B:THR:HG22	2:17:B:ILE:HD12	2	1.44
(1,515)	2:15:B:THR:HG22	2:17:B:ILE:HD13	2	1.44
(1,515)	2:15:B:THR:HG23	2:17:B:ILE:HD11	2	1.44
(1,515)	2:15:B:THR:HG23	2:17:B:ILE:HD12	2	1.44
(1,515)	2:15:B:THR:HG23	2:17:B:ILE:HD13	2	1.44
(1,437)	1:8:A:GLU:HG2	2:20:B:TYR:HB2	14	1.44
(1,437)	1:8:A:GLU:HG2	2:20:B:TYR:HB3	14	1.44
(1,196)	2:15:B:THR:HG21	2:17:B:ILE:HD11	2	1.44
(1,196)	2:15:B:THR:HG21	2:17:B:ILE:HD12	2	1.44
(1,196)	2:15:B:THR:HG21	2:17:B:ILE:HD13	2	1.44
(1,196)	2:15:B:THR:HG22	2:17:B:ILE:HD11	2	1.44
(1,196)	2:15:B:THR:HG22	2:17:B:ILE:HD12	2	1.44
(1,196)	2:15:B:THR:HG22	2:17:B:ILE:HD13	2	1.44
(1,196)	2:15:B:THR:HG23	2:17:B:ILE:HD11	2	1.44
(1,196)	2:15:B:THR:HG23	2:17:B:ILE:HD12	2	1.44
(1,196)	2:15:B:THR:HG23	2:17:B:ILE:HD13	2	1.44
(1,118)	1:8:A:GLU:HG2	2:20:B:TYR:HB2	14	1.44
(1,118)	1:8:A:GLU:HG2	2:20:B:TYR:HB3	14	1.44
(1,15)	2:19:B:PHE:HE1	2:38:D:ILE:HG13	4	1.44
(1,15)	2:19:B:PHE:HE2	2:38:D:ILE:HG13	4	1.44
(1,1)	2:17:B:ILE:HD11	2:40:D:PHE:HD1	8	1.44
(1,1)	2:17:B:ILE:HD11	2:40:D:PHE:HD2	8	1.44
(1,1)	2:17:B:ILE:HD12	2:40:D:PHE:HD1	8	1.44
(1,1)	2:17:B:ILE:HD12	2:40:D:PHE:HD2	8	1.44
(1,1)	2:17:B:ILE:HD13	2:40:D:PHE:HD1	8	1.44
(1,1)	2:17:B:ILE:HD13	2:40:D:PHE:HD2	8	1.44
(1,1171)	2:36:D:THR:HG21	2:38:D:ILE:HD11	7	1.43
(1,1171)	2:36:D:THR:HG21	2:38:D:ILE:HD12	7	1.43

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,1171)	2:36:D:THR:HG21	2:38:D:ILE:HD13	7	1.43
(1,1171)	2:36:D:THR:HG22	2:38:D:ILE:HD11	7	1.43
(1,1171)	2:36:D:THR:HG22	2:38:D:ILE:HD12	7	1.43
(1,1171)	2:36:D:THR:HG22	2:38:D:ILE:HD13	7	1.43
(1,1171)	2:36:D:THR:HG23	2:38:D:ILE:HD11	7	1.43
(1,1171)	2:36:D:THR:HG23	2:38:D:ILE:HD12	7	1.43
(1,1171)	2:36:D:THR:HG23	2:38:D:ILE:HD13	7	1.43
(1,1046)	1:27:C:VAL:HB	2:41:D:TYR:HE1	13	1.43
(1,1046)	1:27:C:VAL:HB	2:41:D:TYR:HE2	13	1.43
(1,852)	2:36:D:THR:HG21	2:38:D:ILE:HD11	7	1.43
(1,852)	2:36:D:THR:HG21	2:38:D:ILE:HD12	7	1.43
(1,852)	2:36:D:THR:HG21	2:38:D:ILE:HD13	7	1.43
(1,852)	2:36:D:THR:HG22	2:38:D:ILE:HD11	7	1.43
(1,852)	2:36:D:THR:HG22	2:38:D:ILE:HD12	7	1.43
(1,852)	2:36:D:THR:HG22	2:38:D:ILE:HD13	7	1.43
(1,852)	2:36:D:THR:HG23	2:38:D:ILE:HD11	7	1.43
(1,852)	2:36:D:THR:HG23	2:38:D:ILE:HD12	7	1.43
(1,852)	2:36:D:THR:HG23	2:38:D:ILE:HD13	7	1.43
(1,727)	1:27:C:VAL:HB	2:41:D:TYR:HE1	13	1.43
(1,727)	1:27:C:VAL:HB	2:41:D:TYR:HE2	13	1.43
(1,1042)	1:25:C:GLY:H	2:40:D:PHE:HD1	1	1.42
(1,1042)	1:25:C:GLY:H	2:40:D:PHE:HD2	1	1.42
(1,723)	1:25:C:GLY:H	2:40:D:PHE:HD1	1	1.42
(1,723)	1:25:C:GLY:H	2:40:D:PHE:HD2	1	1.42
(1,674)	2:17:B:ILE:HD11	2:40:D:PHE:HE1	16	1.42
(1,674)	2:17:B:ILE:HD11	2:40:D:PHE:HE2	16	1.42
(1,674)	2:17:B:ILE:HD12	2:40:D:PHE:HE1	16	1.42
(1,674)	2:17:B:ILE:HD12	2:40:D:PHE:HE2	16	1.42
(1,674)	2:17:B:ILE:HD13	2:40:D:PHE:HE1	16	1.42
(1,674)	2:17:B:ILE:HD13	2:40:D:PHE:HE2	16	1.42
(1,665)	2:19:B:PHE:HE1	2:38:D:ILE:HD11	1	1.42
(1,665)	2:19:B:PHE:HE1	2:38:D:ILE:HD12	1	1.42
(1,665)	2:19:B:PHE:HE1	2:38:D:ILE:HD13	1	1.42
(1,665)	2:19:B:PHE:HE2	2:38:D:ILE:HD11	1	1.42
(1,665)	2:19:B:PHE:HE2	2:38:D:ILE:HD12	1	1.42
(1,665)	2:19:B:PHE:HE2	2:38:D:ILE:HD13	1	1.42
(1,664)	2:19:B:PHE:HB2	2:38:D:ILE:HG13	18	1.42
(1,513)	2:15:B:THR:HB	2:17:B:ILE:HD11	16	1.42
(1,513)	2:15:B:THR:HB	2:17:B:ILE:HD12	16	1.42
(1,513)	2:15:B:THR:HB	2:17:B:ILE:HD13	16	1.42
(1,437)	1:8:A:GLU:HG2	2:20:B:TYR:HB2	9	1.42
(1,437)	1:8:A:GLU:HG2	2:20:B:TYR:HB3	9	1.42

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,194)	2:15:B:THR:HB	2:17:B:ILE:HD11	16	1.42
(1,194)	2:15:B:THR:HB	2:17:B:ILE:HD12	16	1.42
(1,194)	2:15:B:THR:HB	2:17:B:ILE:HD13	16	1.42
(1,118)	1:8:A:GLU:HG2	2:20:B:TYR:HB2	9	1.42
(1,118)	1:8:A:GLU:HG2	2:20:B:TYR:HB3	9	1.42
(1,18)	2:19:B:PHE:HE1	2:38:D:ILE:HD11	1	1.42
(1,18)	2:19:B:PHE:HE1	2:38:D:ILE:HD12	1	1.42
(1,18)	2:19:B:PHE:HE1	2:38:D:ILE:HD13	1	1.42
(1,18)	2:19:B:PHE:HE2	2:38:D:ILE:HD11	1	1.42
(1,18)	2:19:B:PHE:HE2	2:38:D:ILE:HD12	1	1.42
(1,18)	2:19:B:PHE:HE2	2:38:D:ILE:HD13	1	1.42
(1,17)	2:19:B:PHE:HB2	2:38:D:ILE:HG13	18	1.42
(1,9)	2:17:B:ILE:HD11	2:40:D:PHE:HE1	16	1.42
(1,9)	2:17:B:ILE:HD11	2:40:D:PHE:HE2	16	1.42
(1,9)	2:17:B:ILE:HD12	2:40:D:PHE:HE1	16	1.42
(1,9)	2:17:B:ILE:HD12	2:40:D:PHE:HE2	16	1.42
(1,9)	2:17:B:ILE:HD13	2:40:D:PHE:HE1	16	1.42
(1,9)	2:17:B:ILE:HD13	2:40:D:PHE:HE2	16	1.42
(1,1102)	1:29:C:GLU:HB3	2:41:D:TYR:HD1	5	1.41
(1,1102)	1:29:C:GLU:HB3	2:41:D:TYR:HD2	5	1.41
(1,1045)	1:27:C:VAL:HB	2:41:D:TYR:HD1	1	1.41
(1,1045)	1:27:C:VAL:HB	2:41:D:TYR:HD2	1	1.41
(1,783)	1:29:C:GLU:HB3	2:41:D:TYR:HD1	5	1.41
(1,783)	1:29:C:GLU:HB3	2:41:D:TYR:HD2	5	1.41
(1,726)	1:27:C:VAL:HB	2:41:D:TYR:HD1	1	1.41
(1,726)	1:27:C:VAL:HB	2:41:D:TYR:HD2	1	1.41
(1,671)	2:17:B:ILE:HG13	2:40:D:PHE:HE1	10	1.41
(1,671)	2:17:B:ILE:HG13	2:40:D:PHE:HE2	10	1.41
(1,437)	1:8:A:GLU:HG2	2:20:B:TYR:HB2	2	1.41
(1,437)	1:8:A:GLU:HG2	2:20:B:TYR:HB3	2	1.41
(1,420)	1:4:A:GLY:HA3	1:5:A:HIS:HD2	8	1.41
(1,361)	1:10:A:PHE:HE1	2:14:B:GLY:HA3	20	1.41
(1,361)	1:10:A:PHE:HE2	2:14:B:GLY:HA3	20	1.41
(1,118)	1:8:A:GLU:HG2	2:20:B:TYR:HB2	2	1.41
(1,118)	1:8:A:GLU:HG2	2:20:B:TYR:HB3	2	1.41
(1,101)	1:4:A:GLY:HA3	1:5:A:HIS:HD2	8	1.41
(1,42)	1:10:A:PHE:HE1	2:14:B:GLY:HA3	20	1.41
(1,42)	1:10:A:PHE:HE2	2:14:B:GLY:HA3	20	1.41
(1,6)	2:17:B:ILE:HG13	2:40:D:PHE:HE1	10	1.41
(1,6)	2:17:B:ILE:HG13	2:40:D:PHE:HE2	10	1.41
(1,1176)	2:19:B:PHE:HB3	2:38:D:ILE:HG13	2	1.4
(1,1063)	1:22:C:GLY:H	2:39:D:SER:HA	4	1.4

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,1058)	1:31:C:PHE:HD1	2:35:D:GLY:HA3	14	1.4
(1,1058)	1:31:C:PHE:HD2	2:35:D:GLY:HA3	14	1.4
(1,1015)	1:30:C:TYR:HD1	2:39:D:SER:HA	17	1.4
(1,1015)	1:30:C:TYR:HD2	2:39:D:SER:HA	17	1.4
(1,857)	2:19:B:PHE:HB3	2:38:D:ILE:HG13	2	1.4
(1,744)	1:22:C:GLY:H	2:39:D:SER:HA	4	1.4
(1,739)	1:31:C:PHE:HD1	2:35:D:GLY:HA3	14	1.4
(1,739)	1:31:C:PHE:HD2	2:35:D:GLY:HA3	14	1.4
(1,696)	1:30:C:TYR:HD1	2:39:D:SER:HA	17	1.4
(1,696)	1:30:C:TYR:HD2	2:39:D:SER:HA	17	1.4
(1,671)	2:17:B:ILE:HG13	2:40:D:PHE:HE1	17	1.4
(1,671)	2:17:B:ILE:HG13	2:40:D:PHE:HE2	17	1.4
(1,446)	1:8:A:GLU:HB3	2:20:B:TYR:HD1	9	1.4
(1,446)	1:8:A:GLU:HB3	2:20:B:TYR:HD2	9	1.4
(1,437)	1:8:A:GLU:HG2	2:20:B:TYR:HB2	7	1.4
(1,437)	1:8:A:GLU:HG2	2:20:B:TYR:HB3	7	1.4
(1,384)	1:10:A:PHE:HE1	2:16:B:PRO:HD2	1	1.4
(1,384)	1:10:A:PHE:HE2	2:16:B:PRO:HD2	1	1.4
(1,344)	1:10:A:PHE:HE1	2:16:B:PRO:HD2	17	1.4
(1,344)	1:10:A:PHE:HE2	2:16:B:PRO:HD2	17	1.4
(1,127)	1:8:A:GLU:HB3	2:20:B:TYR:HD1	9	1.4
(1,127)	1:8:A:GLU:HB3	2:20:B:TYR:HD2	9	1.4
(1,118)	1:8:A:GLU:HG2	2:20:B:TYR:HB2	7	1.4
(1,118)	1:8:A:GLU:HG2	2:20:B:TYR:HB3	7	1.4
(1,65)	1:10:A:PHE:HE1	2:16:B:PRO:HD2	1	1.4
(1,65)	1:10:A:PHE:HE2	2:16:B:PRO:HD2	1	1.4
(1,25)	1:10:A:PHE:HE1	2:16:B:PRO:HD2	17	1.4
(1,25)	1:10:A:PHE:HE2	2:16:B:PRO:HD2	17	1.4
(1,6)	2:17:B:ILE:HG13	2:40:D:PHE:HE1	17	1.4
(1,6)	2:17:B:ILE:HG13	2:40:D:PHE:HE2	17	1.4
(1,1249)	1:33:C:ARG:HB2	1:33:C:ARG:HE	4	1.39
(1,1171)	2:36:D:THR:HG21	2:38:D:ILE:HD11	8	1.39
(1,1171)	2:36:D:THR:HG21	2:38:D:ILE:HD12	8	1.39
(1,1171)	2:36:D:THR:HG21	2:38:D:ILE:HD13	8	1.39
(1,1171)	2:36:D:THR:HG22	2:38:D:ILE:HD11	8	1.39
(1,1171)	2:36:D:THR:HG22	2:38:D:ILE:HD12	8	1.39
(1,1171)	2:36:D:THR:HG22	2:38:D:ILE:HD13	8	1.39
(1,1171)	2:36:D:THR:HG23	2:38:D:ILE:HD11	8	1.39
(1,1171)	2:36:D:THR:HG23	2:38:D:ILE:HD12	8	1.39
(1,1171)	2:36:D:THR:HG23	2:38:D:ILE:HD13	8	1.39
(1,1092)	1:29:C:GLU:HG2	2:41:D:TYR:HD1	11	1.39
(1,1092)	1:29:C:GLU:HG2	2:41:D:TYR:HD2	11	1.39

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,930)	1:33:C:ARG:HB2	1:33:C:ARG:HE	4	1.39
(1,852)	2:36:D:THR:HG21	2:38:D:ILE:HD11	8	1.39
(1,852)	2:36:D:THR:HG21	2:38:D:ILE:HD12	8	1.39
(1,852)	2:36:D:THR:HG21	2:38:D:ILE:HD13	8	1.39
(1,852)	2:36:D:THR:HG22	2:38:D:ILE:HD11	8	1.39
(1,852)	2:36:D:THR:HG22	2:38:D:ILE:HD12	8	1.39
(1,852)	2:36:D:THR:HG22	2:38:D:ILE:HD13	8	1.39
(1,852)	2:36:D:THR:HG23	2:38:D:ILE:HD11	8	1.39
(1,852)	2:36:D:THR:HG23	2:38:D:ILE:HD12	8	1.39
(1,852)	2:36:D:THR:HG23	2:38:D:ILE:HD13	8	1.39
(1,773)	1:29:C:GLU:HG2	2:41:D:TYR:HD1	11	1.39
(1,773)	1:29:C:GLU:HG2	2:41:D:TYR:HD2	11	1.39
(1,657)	2:19:B:PHE:HD1	2:38:D:ILE:HD11	12	1.39
(1,657)	2:19:B:PHE:HD1	2:38:D:ILE:HD12	12	1.39
(1,657)	2:19:B:PHE:HD1	2:38:D:ILE:HD13	12	1.39
(1,657)	2:19:B:PHE:HD2	2:38:D:ILE:HD11	12	1.39
(1,657)	2:19:B:PHE:HD2	2:38:D:ILE:HD12	12	1.39
(1,657)	2:19:B:PHE:HD2	2:38:D:ILE:HD13	12	1.39
(1,520)	2:17:B:ILE:HG13	2:40:D:PHE:HB3	6	1.39
(1,399)	1:7:A:PRO:HG2	1:10:A:PHE:HE1	3	1.39
(1,399)	1:7:A:PRO:HG2	1:10:A:PHE:HE2	3	1.39
(1,399)	1:7:A:PRO:HG3	1:10:A:PHE:HE1	3	1.39
(1,399)	1:7:A:PRO:HG3	1:10:A:PHE:HE2	3	1.39
(1,344)	1:10:A:PHE:HE1	2:16:B:PRO:HD2	5	1.39
(1,344)	1:10:A:PHE:HE2	2:16:B:PRO:HD2	5	1.39
(1,201)	2:17:B:ILE:HG13	2:40:D:PHE:HB3	6	1.39
(1,80)	1:7:A:PRO:HG2	1:10:A:PHE:HE1	3	1.39
(1,80)	1:7:A:PRO:HG2	1:10:A:PHE:HE2	3	1.39
(1,80)	1:7:A:PRO:HG3	1:10:A:PHE:HE1	3	1.39
(1,80)	1:7:A:PRO:HG3	1:10:A:PHE:HE2	3	1.39
(1,25)	1:10:A:PHE:HE1	2:16:B:PRO:HD2	5	1.39
(1,25)	1:10:A:PHE:HE2	2:16:B:PRO:HD2	5	1.39
(1,10)	2:19:B:PHE:HD1	2:38:D:ILE:HD11	12	1.39
(1,10)	2:19:B:PHE:HD1	2:38:D:ILE:HD12	12	1.39
(1,10)	2:19:B:PHE:HD1	2:38:D:ILE:HD13	12	1.39
(1,10)	2:19:B:PHE:HD2	2:38:D:ILE:HD11	12	1.39
(1,10)	2:19:B:PHE:HD2	2:38:D:ILE:HD12	12	1.39
(1,10)	2:19:B:PHE:HD2	2:38:D:ILE:HD13	12	1.39
(1,1147)	1:28:C:PRO:HB2	1:31:C:PHE:HE1	1	1.38
(1,1147)	1:28:C:PRO:HB2	1:31:C:PHE:HE2	1	1.38
(1,1122)	1:25:C:GLY:HA2	1:26:C:HIS:HD2	1	1.38
(1,1093)	1:29:C:GLU:HG2	2:41:D:TYR:HB2	11	1.38

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,1093)	1:29:C:GLU:HG2	2:41:D:TYR:HB3	11	1.38
(1,828)	1:28:C:PRO:HB2	1:31:C:PHE:HE1	1	1.38
(1,828)	1:28:C:PRO:HB2	1:31:C:PHE:HE2	1	1.38
(1,803)	1:25:C:GLY:HA2	1:26:C:HIS:HD2	1	1.38
(1,774)	1:29:C:GLU:HG2	2:41:D:TYR:HB2	11	1.38
(1,774)	1:29:C:GLU:HG2	2:41:D:TYR:HB3	11	1.38
(1,666)	2:17:B:ILE:HD11	2:40:D:PHE:HD1	6	1.38
(1,666)	2:17:B:ILE:HD11	2:40:D:PHE:HD2	6	1.38
(1,666)	2:17:B:ILE:HD12	2:40:D:PHE:HD1	6	1.38
(1,666)	2:17:B:ILE:HD12	2:40:D:PHE:HD2	6	1.38
(1,666)	2:17:B:ILE:HD13	2:40:D:PHE:HD1	6	1.38
(1,666)	2:17:B:ILE:HD13	2:40:D:PHE:HD2	6	1.38
(1,595)	1:12:A:ARG:HA	1:12:A:ARG:HE	7	1.38
(1,503)	1:7:A:PRO:HB3	1:10:A:PHE:HE1	14	1.38
(1,503)	1:7:A:PRO:HB3	1:10:A:PHE:HE2	14	1.38
(1,276)	1:12:A:ARG:HA	1:12:A:ARG:HE	7	1.38
(1,184)	1:7:A:PRO:HB3	1:10:A:PHE:HE1	14	1.38
(1,184)	1:7:A:PRO:HB3	1:10:A:PHE:HE2	14	1.38
(1,1)	2:17:B:ILE:HD11	2:40:D:PHE:HD1	6	1.38
(1,1)	2:17:B:ILE:HD11	2:40:D:PHE:HD2	6	1.38
(1,1)	2:17:B:ILE:HD12	2:40:D:PHE:HD1	6	1.38
(1,1)	2:17:B:ILE:HD12	2:40:D:PHE:HD2	6	1.38
(1,1)	2:17:B:ILE:HD13	2:40:D:PHE:HD1	6	1.38
(1,1)	2:17:B:ILE:HD13	2:40:D:PHE:HD2	6	1.38
(1,595)	1:12:A:ARG:HA	1:12:A:ARG:HE	16	1.37
(1,446)	1:8:A:GLU:HB3	2:20:B:TYR:HD1	11	1.37
(1,446)	1:8:A:GLU:HB3	2:20:B:TYR:HD2	11	1.37
(1,407)	1:1:A:GLY:H	2:18:B:SER:HA	12	1.37
(1,276)	1:12:A:ARG:HA	1:12:A:ARG:HE	16	1.37
(1,127)	1:8:A:GLU:HB3	2:20:B:TYR:HD1	11	1.37
(1,127)	1:8:A:GLU:HB3	2:20:B:TYR:HD2	11	1.37
(1,88)	1:1:A:GLY:H	2:18:B:SER:HA	12	1.37
(1,1157)	1:25:C:GLY:H	1:27:C:VAL:H	4	1.36
(1,1046)	1:27:C:VAL:HB	2:41:D:TYR:HE1	11	1.36
(1,1046)	1:27:C:VAL:HB	2:41:D:TYR:HE2	11	1.36
(1,838)	1:25:C:GLY:H	1:27:C:VAL:H	4	1.36
(1,727)	1:27:C:VAL:HB	2:41:D:TYR:HE1	11	1.36
(1,727)	1:27:C:VAL:HB	2:41:D:TYR:HE2	11	1.36
(1,629)	2:17:B:ILE:H	2:17:B:ILE:HD11	15	1.36
(1,629)	2:17:B:ILE:H	2:17:B:ILE:HD12	15	1.36
(1,629)	2:17:B:ILE:H	2:17:B:ILE:HD13	15	1.36
(1,595)	1:12:A:ARG:HA	1:12:A:ARG:HE	2	1.36

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,575)	1:10:A:PHE:HE1	2:15:B:THR:HA	18	1.36
(1,575)	1:10:A:PHE:HE2	2:15:B:THR:HA	18	1.36
(1,492)	1:7:A:PRO:HB2	1:10:A:PHE:HD1	16	1.36
(1,492)	1:7:A:PRO:HB2	1:10:A:PHE:HD2	16	1.36
(1,437)	1:8:A:GLU:HG2	2:20:B:TYR:HB2	17	1.36
(1,437)	1:8:A:GLU:HG2	2:20:B:TYR:HB3	17	1.36
(1,348)	1:11:A:VAL:HA	1:12:A:ARG:HD2	5	1.36
(1,348)	1:11:A:VAL:HA	1:12:A:ARG:HD3	5	1.36
(1,310)	2:17:B:ILE:H	2:17:B:ILE:HD11	15	1.36
(1,310)	2:17:B:ILE:H	2:17:B:ILE:HD12	15	1.36
(1,310)	2:17:B:ILE:H	2:17:B:ILE:HD13	15	1.36
(1,276)	1:12:A:ARG:HA	1:12:A:ARG:HE	2	1.36
(1,256)	1:10:A:PHE:HE1	2:15:B:THR:HA	18	1.36
(1,256)	1:10:A:PHE:HE2	2:15:B:THR:HA	18	1.36
(1,173)	1:7:A:PRO:HB2	1:10:A:PHE:HD1	16	1.36
(1,173)	1:7:A:PRO:HB2	1:10:A:PHE:HD2	16	1.36
(1,118)	1:8:A:GLU:HG2	2:20:B:TYR:HB2	17	1.36
(1,118)	1:8:A:GLU:HG2	2:20:B:TYR:HB3	17	1.36
(1,29)	1:11:A:VAL:HA	1:12:A:ARG:HD2	5	1.36
(1,29)	1:11:A:VAL:HA	1:12:A:ARG:HD3	5	1.36
(1,1251)	1:33:C:ARG:HA	1:33:C:ARG:HE	19	1.35
(1,1125)	1:31:C:PHE:HD1	1:32:C:VAL:HG11	18	1.35
(1,1125)	1:31:C:PHE:HD1	1:32:C:VAL:HG12	18	1.35
(1,1125)	1:31:C:PHE:HD1	1:32:C:VAL:HG13	18	1.35
(1,1125)	1:31:C:PHE:HD1	1:32:C:VAL:HG21	18	1.35
(1,1125)	1:31:C:PHE:HD1	1:32:C:VAL:HG22	18	1.35
(1,1125)	1:31:C:PHE:HD1	1:32:C:VAL:HG23	18	1.35
(1,1125)	1:31:C:PHE:HD2	1:32:C:VAL:HG11	18	1.35
(1,1125)	1:31:C:PHE:HD2	1:32:C:VAL:HG12	18	1.35
(1,1125)	1:31:C:PHE:HD2	1:32:C:VAL:HG13	18	1.35
(1,1125)	1:31:C:PHE:HD2	1:32:C:VAL:HG21	18	1.35
(1,1125)	1:31:C:PHE:HD2	1:32:C:VAL:HG22	18	1.35
(1,1125)	1:31:C:PHE:HD2	1:32:C:VAL:HG23	18	1.35
(1,1093)	1:29:C:GLU:HG2	2:41:D:TYR:HB2	4	1.35
(1,1093)	1:29:C:GLU:HG2	2:41:D:TYR:HB3	4	1.35
(1,1046)	1:27:C:VAL:HB	2:41:D:TYR:HE1	16	1.35
(1,1046)	1:27:C:VAL:HB	2:41:D:TYR:HE2	16	1.35
(1,932)	1:33:C:ARG:HA	1:33:C:ARG:HE	19	1.35
(1,806)	1:31:C:PHE:HD1	1:32:C:VAL:HG11	18	1.35
(1,806)	1:31:C:PHE:HD1	1:32:C:VAL:HG12	18	1.35
(1,806)	1:31:C:PHE:HD1	1:32:C:VAL:HG13	18	1.35
(1,806)	1:31:C:PHE:HD1	1:32:C:VAL:HG21	18	1.35

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,806)	1:31:C:PHE:HD1	1:32:C:VAL:HG22	18	1.35
(1,806)	1:31:C:PHE:HD1	1:32:C:VAL:HG23	18	1.35
(1,806)	1:31:C:PHE:HD2	1:32:C:VAL:HG11	18	1.35
(1,806)	1:31:C:PHE:HD2	1:32:C:VAL:HG12	18	1.35
(1,806)	1:31:C:PHE:HD2	1:32:C:VAL:HG13	18	1.35
(1,806)	1:31:C:PHE:HD2	1:32:C:VAL:HG21	18	1.35
(1,806)	1:31:C:PHE:HD2	1:32:C:VAL:HG22	18	1.35
(1,806)	1:31:C:PHE:HD2	1:32:C:VAL:HG23	18	1.35
(1,774)	1:29:C:GLU:HG2	2:41:D:TYR:HB2	4	1.35
(1,774)	1:29:C:GLU:HG2	2:41:D:TYR:HB3	4	1.35
(1,727)	1:27:C:VAL:HB	2:41:D:TYR:HE1	16	1.35
(1,727)	1:27:C:VAL:HB	2:41:D:TYR:HE2	16	1.35
(1,663)	2:19:B:PHE:HD1	2:38:D:ILE:HG13	15	1.35
(1,663)	2:19:B:PHE:HD2	2:38:D:ILE:HG13	15	1.35
(1,352)	2:17:B:ILE:HG13	2:39:D:SER:HA	16	1.35
(1,33)	2:17:B:ILE:HG13	2:39:D:SER:HA	16	1.35
(1,16)	2:19:B:PHE:HD1	2:38:D:ILE:HG13	15	1.35
(1,16)	2:19:B:PHE:HD2	2:38:D:ILE:HG13	15	1.35
(1,1251)	1:33:C:ARG:HA	1:33:C:ARG:HE	13	1.34
(1,1176)	2:19:B:PHE:HB3	2:38:D:ILE:HG13	6	1.34
(1,1092)	1:29:C:GLU:HG2	2:41:D:TYR:HD1	18	1.34
(1,1092)	1:29:C:GLU:HG2	2:41:D:TYR:HD2	18	1.34
(1,932)	1:33:C:ARG:HA	1:33:C:ARG:HE	13	1.34
(1,857)	2:19:B:PHE:HB3	2:38:D:ILE:HG13	6	1.34
(1,773)	1:29:C:GLU:HG2	2:41:D:TYR:HD1	18	1.34
(1,773)	1:29:C:GLU:HG2	2:41:D:TYR:HD2	18	1.34
(1,666)	2:17:B:ILE:HD11	2:40:D:PHE:HD1	12	1.34
(1,666)	2:17:B:ILE:HD11	2:40:D:PHE:HD2	12	1.34
(1,666)	2:17:B:ILE:HD12	2:40:D:PHE:HD1	12	1.34
(1,666)	2:17:B:ILE:HD12	2:40:D:PHE:HD2	12	1.34
(1,666)	2:17:B:ILE:HD13	2:40:D:PHE:HD1	12	1.34
(1,666)	2:17:B:ILE:HD13	2:40:D:PHE:HD2	12	1.34
(1,663)	2:19:B:PHE:HD1	2:38:D:ILE:HG13	17	1.34
(1,663)	2:19:B:PHE:HD2	2:38:D:ILE:HG13	17	1.34
(1,596)	1:7:A:PRO:HG2	1:10:A:PHE:HE1	3	1.34
(1,596)	1:7:A:PRO:HG2	1:10:A:PHE:HE2	3	1.34
(1,596)	1:7:A:PRO:HG3	1:10:A:PHE:HE1	3	1.34
(1,596)	1:7:A:PRO:HG3	1:10:A:PHE:HE2	3	1.34
(1,503)	1:7:A:PRO:HB3	1:10:A:PHE:HE1	5	1.34
(1,503)	1:7:A:PRO:HB3	1:10:A:PHE:HE2	5	1.34
(1,277)	1:7:A:PRO:HG2	1:10:A:PHE:HE1	3	1.34
(1,277)	1:7:A:PRO:HG2	1:10:A:PHE:HE2	3	1.34

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,277)	1:7:A:PRO:HG3	1:10:A:PHE:HE1	3	1.34
(1,277)	1:7:A:PRO:HG3	1:10:A:PHE:HE2	3	1.34
(1,184)	1:7:A:PRO:HB3	1:10:A:PHE:HE1	5	1.34
(1,184)	1:7:A:PRO:HB3	1:10:A:PHE:HE2	5	1.34
(1,16)	2:19:B:PHE:HD1	2:38:D:ILE:HG13	17	1.34
(1,16)	2:19:B:PHE:HD2	2:38:D:ILE:HG13	17	1.34
(1,1)	2:17:B:ILE:HD11	2:40:D:PHE:HD1	12	1.34
(1,1)	2:17:B:ILE:HD11	2:40:D:PHE:HD2	12	1.34
(1,1)	2:17:B:ILE:HD12	2:40:D:PHE:HD1	12	1.34
(1,1)	2:17:B:ILE:HD12	2:40:D:PHE:HD2	12	1.34
(1,1)	2:17:B:ILE:HD13	2:40:D:PHE:HD1	12	1.34
(1,1)	2:17:B:ILE:HD13	2:40:D:PHE:HD2	12	1.34
(1,1109)	1:31:C:PHE:HD1	2:36:D:THR:HG21	20	1.33
(1,1109)	1:31:C:PHE:HD1	2:36:D:THR:HG22	20	1.33
(1,1109)	1:31:C:PHE:HD1	2:36:D:THR:HG23	20	1.33
(1,1109)	1:31:C:PHE:HD2	2:36:D:THR:HG21	20	1.33
(1,1109)	1:31:C:PHE:HD2	2:36:D:THR:HG22	20	1.33
(1,1109)	1:31:C:PHE:HD2	2:36:D:THR:HG23	20	1.33
(1,1092)	1:29:C:GLU:HG2	2:41:D:TYR:HD1	19	1.33
(1,1092)	1:29:C:GLU:HG2	2:41:D:TYR:HD2	19	1.33
(1,1004)	1:32:C:VAL:HA	1:33:C:ARG:HD2	10	1.33
(1,1004)	1:32:C:VAL:HA	1:33:C:ARG:HD3	10	1.33
(1,790)	1:31:C:PHE:HD1	2:36:D:THR:HG21	20	1.33
(1,790)	1:31:C:PHE:HD1	2:36:D:THR:HG22	20	1.33
(1,790)	1:31:C:PHE:HD1	2:36:D:THR:HG23	20	1.33
(1,790)	1:31:C:PHE:HD2	2:36:D:THR:HG21	20	1.33
(1,790)	1:31:C:PHE:HD2	2:36:D:THR:HG22	20	1.33
(1,790)	1:31:C:PHE:HD2	2:36:D:THR:HG23	20	1.33
(1,773)	1:29:C:GLU:HG2	2:41:D:TYR:HD1	19	1.33
(1,773)	1:29:C:GLU:HG2	2:41:D:TYR:HD2	19	1.33
(1,685)	1:32:C:VAL:HA	1:33:C:ARG:HD2	10	1.33
(1,685)	1:32:C:VAL:HA	1:33:C:ARG:HD3	10	1.33
(1,520)	2:17:B:ILE:HG13	2:40:D:PHE:HB3	8	1.33
(1,503)	1:7:A:PRO:HB3	1:10:A:PHE:HE1	10	1.33
(1,503)	1:7:A:PRO:HB3	1:10:A:PHE:HE2	10	1.33
(1,436)	1:8:A:GLU:HG2	2:20:B:TYR:HD1	18	1.33
(1,436)	1:8:A:GLU:HG2	2:20:B:TYR:HD2	18	1.33
(1,201)	2:17:B:ILE:HG13	2:40:D:PHE:HB3	8	1.33
(1,184)	1:7:A:PRO:HB3	1:10:A:PHE:HE1	10	1.33
(1,184)	1:7:A:PRO:HB3	1:10:A:PHE:HE2	10	1.33
(1,117)	1:8:A:GLU:HG2	2:20:B:TYR:HD1	18	1.33
(1,117)	1:8:A:GLU:HG2	2:20:B:TYR:HD2	18	1.33

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,1093)	1:29:C:GLU:HG2	2:41:D:TYR:HB2	18	1.32
(1,1093)	1:29:C:GLU:HG2	2:41:D:TYR:HB3	18	1.32
(1,774)	1:29:C:GLU:HG2	2:41:D:TYR:HB2	18	1.32
(1,774)	1:29:C:GLU:HG2	2:41:D:TYR:HB3	18	1.32
(1,519)	1:3:A:ALA:HB1	2:19:B:PHE:HB3	11	1.32
(1,519)	1:3:A:ALA:HB2	2:19:B:PHE:HB3	11	1.32
(1,519)	1:3:A:ALA:HB3	2:19:B:PHE:HB3	11	1.32
(1,481)	1:1:A:GLY:H	2:19:B:PHE:H	6	1.32
(1,420)	1:4:A:GLY:HA3	1:5:A:HIS:HD2	1	1.32
(1,420)	1:4:A:GLY:HA3	1:5:A:HIS:HD2	20	1.32
(1,402)	1:10:A:PHE:HD1	2:14:B:GLY:HA3	5	1.32
(1,402)	1:10:A:PHE:HD2	2:14:B:GLY:HA3	5	1.32
(1,399)	1:7:A:PRO:HG2	1:10:A:PHE:HE1	19	1.32
(1,399)	1:7:A:PRO:HG2	1:10:A:PHE:HE2	19	1.32
(1,399)	1:7:A:PRO:HG3	1:10:A:PHE:HE1	19	1.32
(1,399)	1:7:A:PRO:HG3	1:10:A:PHE:HE2	19	1.32
(1,200)	1:3:A:ALA:HB1	2:19:B:PHE:HB3	11	1.32
(1,200)	1:3:A:ALA:HB2	2:19:B:PHE:HB3	11	1.32
(1,200)	1:3:A:ALA:HB3	2:19:B:PHE:HB3	11	1.32
(1,162)	1:1:A:GLY:H	2:19:B:PHE:H	6	1.32
(1,101)	1:4:A:GLY:HA3	1:5:A:HIS:HD2	1	1.32
(1,101)	1:4:A:GLY:HA3	1:5:A:HIS:HD2	20	1.32
(1,83)	1:10:A:PHE:HD1	2:14:B:GLY:HA3	5	1.32
(1,83)	1:10:A:PHE:HD2	2:14:B:GLY:HA3	5	1.32
(1,80)	1:7:A:PRO:HG2	1:10:A:PHE:HE1	19	1.32
(1,80)	1:7:A:PRO:HG2	1:10:A:PHE:HE2	19	1.32
(1,80)	1:7:A:PRO:HG3	1:10:A:PHE:HE1	19	1.32
(1,80)	1:7:A:PRO:HG3	1:10:A:PHE:HE2	19	1.32
(1,1093)	1:29:C:GLU:HG2	2:41:D:TYR:HB2	5	1.31
(1,1093)	1:29:C:GLU:HG2	2:41:D:TYR:HB3	5	1.31
(1,1076)	1:25:C:GLY:HA3	1:26:C:HIS:HD2	16	1.31
(1,1058)	1:31:C:PHE:HD1	2:35:D:GLY:HA3	20	1.31
(1,1058)	1:31:C:PHE:HD2	2:35:D:GLY:HA3	20	1.31
(1,774)	1:29:C:GLU:HG2	2:41:D:TYR:HB2	5	1.31
(1,774)	1:29:C:GLU:HG2	2:41:D:TYR:HB3	5	1.31
(1,757)	1:25:C:GLY:HA3	1:26:C:HIS:HD2	16	1.31
(1,739)	1:31:C:PHE:HD1	2:35:D:GLY:HA3	20	1.31
(1,739)	1:31:C:PHE:HD2	2:35:D:GLY:HA3	20	1.31
(1,466)	1:4:A:GLY:HA2	1:5:A:HIS:HD2	20	1.31
(1,402)	1:10:A:PHE:HD1	2:14:B:GLY:HA3	12	1.31
(1,402)	1:10:A:PHE:HD2	2:14:B:GLY:HA3	12	1.31
(1,351)	2:16:B:PRO:HA	2:17:B:ILE:HD11	10	1.31

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,351)	2:16:B:PRO:HA	2:17:B:ILE:HD12	10	1.31
(1,351)	2:16:B:PRO:HA	2:17:B:ILE:HD13	10	1.31
(1,147)	1:4:A:GLY:HA2	1:5:A:HIS:HD2	20	1.31
(1,83)	1:10:A:PHE:HD1	2:14:B:GLY:HA3	12	1.31
(1,83)	1:10:A:PHE:HD2	2:14:B:GLY:HA3	12	1.31
(1,32)	2:16:B:PRO:HA	2:17:B:ILE:HD11	10	1.31
(1,32)	2:16:B:PRO:HA	2:17:B:ILE:HD12	10	1.31
(1,32)	2:16:B:PRO:HA	2:17:B:ILE:HD13	10	1.31
(1,662)	2:19:B:PHE:HE1	2:38:D:ILE:HG13	18	1.3
(1,662)	2:19:B:PHE:HE2	2:38:D:ILE:HG13	18	1.3
(1,513)	2:15:B:THR:HB	2:17:B:ILE:HD11	3	1.3
(1,513)	2:15:B:THR:HB	2:17:B:ILE:HD12	3	1.3
(1,513)	2:15:B:THR:HB	2:17:B:ILE:HD13	3	1.3
(1,437)	1:8:A:GLU:HG2	2:20:B:TYR:HB2	6	1.3
(1,437)	1:8:A:GLU:HG2	2:20:B:TYR:HB3	6	1.3
(1,436)	1:8:A:GLU:HG2	2:20:B:TYR:HD1	20	1.3
(1,436)	1:8:A:GLU:HG2	2:20:B:TYR:HD2	20	1.3
(1,356)	1:8:A:GLU:HG2	2:20:B:TYR:HE1	13	1.3
(1,356)	1:8:A:GLU:HG2	2:20:B:TYR:HE2	13	1.3
(1,194)	2:15:B:THR:HB	2:17:B:ILE:HD11	3	1.3
(1,194)	2:15:B:THR:HB	2:17:B:ILE:HD12	3	1.3
(1,194)	2:15:B:THR:HB	2:17:B:ILE:HD13	3	1.3
(1,118)	1:8:A:GLU:HG2	2:20:B:TYR:HB2	6	1.3
(1,118)	1:8:A:GLU:HG2	2:20:B:TYR:HB3	6	1.3
(1,117)	1:8:A:GLU:HG2	2:20:B:TYR:HD1	20	1.3
(1,117)	1:8:A:GLU:HG2	2:20:B:TYR:HD2	20	1.3
(1,37)	1:8:A:GLU:HG2	2:20:B:TYR:HE1	13	1.3
(1,37)	1:8:A:GLU:HG2	2:20:B:TYR:HE2	13	1.3
(1,15)	2:19:B:PHE:HE1	2:38:D:ILE:HG13	18	1.3
(1,15)	2:19:B:PHE:HE2	2:38:D:ILE:HG13	18	1.3
(1,1122)	1:25:C:GLY:HA2	1:26:C:HIS:HD2	5	1.29
(1,1092)	1:29:C:GLU:HG2	2:41:D:TYR:HD1	9	1.29
(1,1092)	1:29:C:GLU:HG2	2:41:D:TYR:HD2	9	1.29
(1,1071)	1:27:C:VAL:HG11	2:41:D:TYR:HE1	9	1.29
(1,1071)	1:27:C:VAL:HG11	2:41:D:TYR:HE2	9	1.29
(1,1071)	1:27:C:VAL:HG12	2:41:D:TYR:HE1	9	1.29
(1,1071)	1:27:C:VAL:HG12	2:41:D:TYR:HE2	9	1.29
(1,1071)	1:27:C:VAL:HG13	2:41:D:TYR:HE1	9	1.29
(1,1071)	1:27:C:VAL:HG13	2:41:D:TYR:HE2	9	1.29
(1,803)	1:25:C:GLY:HA2	1:26:C:HIS:HD2	5	1.29
(1,773)	1:29:C:GLU:HG2	2:41:D:TYR:HD1	9	1.29
(1,773)	1:29:C:GLU:HG2	2:41:D:TYR:HD2	9	1.29

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,752)	1:27:C:VAL:HG11	2:41:D:TYR:HE1	9	1.29
(1,752)	1:27:C:VAL:HG11	2:41:D:TYR:HE2	9	1.29
(1,752)	1:27:C:VAL:HG12	2:41:D:TYR:HE1	9	1.29
(1,752)	1:27:C:VAL:HG12	2:41:D:TYR:HE2	9	1.29
(1,752)	1:27:C:VAL:HG13	2:41:D:TYR:HE1	9	1.29
(1,752)	1:27:C:VAL:HG13	2:41:D:TYR:HE2	9	1.29
(1,674)	2:17:B:ILE:HD11	2:40:D:PHE:HE1	2	1.29
(1,674)	2:17:B:ILE:HD11	2:40:D:PHE:HE2	2	1.29
(1,674)	2:17:B:ILE:HD12	2:40:D:PHE:HE1	2	1.29
(1,674)	2:17:B:ILE:HD12	2:40:D:PHE:HE2	2	1.29
(1,674)	2:17:B:ILE:HD13	2:40:D:PHE:HE1	2	1.29
(1,674)	2:17:B:ILE:HD13	2:40:D:PHE:HE2	2	1.29
(1,657)	2:19:B:PHE:HD1	2:38:D:ILE:HD11	16	1.29
(1,657)	2:19:B:PHE:HD1	2:38:D:ILE:HD12	16	1.29
(1,657)	2:19:B:PHE:HD1	2:38:D:ILE:HD13	16	1.29
(1,657)	2:19:B:PHE:HD2	2:38:D:ILE:HD11	16	1.29
(1,657)	2:19:B:PHE:HD2	2:38:D:ILE:HD12	16	1.29
(1,657)	2:19:B:PHE:HD2	2:38:D:ILE:HD13	16	1.29
(1,483)	1:1:A:GLY:H	2:20:B:TYR:HD1	9	1.29
(1,483)	1:1:A:GLY:H	2:20:B:TYR:HD2	9	1.29
(1,164)	1:1:A:GLY:H	2:20:B:TYR:HD1	9	1.29
(1,164)	1:1:A:GLY:H	2:20:B:TYR:HD2	9	1.29
(1,10)	2:19:B:PHE:HD1	2:38:D:ILE:HD11	16	1.29
(1,10)	2:19:B:PHE:HD1	2:38:D:ILE:HD12	16	1.29
(1,10)	2:19:B:PHE:HD1	2:38:D:ILE:HD13	16	1.29
(1,10)	2:19:B:PHE:HD2	2:38:D:ILE:HD11	16	1.29
(1,10)	2:19:B:PHE:HD2	2:38:D:ILE:HD12	16	1.29
(1,10)	2:19:B:PHE:HD2	2:38:D:ILE:HD13	16	1.29
(1,9)	2:17:B:ILE:HD11	2:40:D:PHE:HE1	2	1.29
(1,9)	2:17:B:ILE:HD11	2:40:D:PHE:HE2	2	1.29
(1,9)	2:17:B:ILE:HD12	2:40:D:PHE:HE1	2	1.29
(1,9)	2:17:B:ILE:HD12	2:40:D:PHE:HE2	2	1.29
(1,9)	2:17:B:ILE:HD13	2:40:D:PHE:HE1	2	1.29
(1,9)	2:17:B:ILE:HD13	2:40:D:PHE:HE2	2	1.29
(1,1157)	1:25:C:GLY:H	1:27:C:VAL:H	11	1.28
(1,1017)	1:31:C:PHE:HE1	2:35:D:GLY:HA3	3	1.28
(1,1017)	1:31:C:PHE:HE2	2:35:D:GLY:HA3	3	1.28
(1,838)	1:25:C:GLY:H	1:27:C:VAL:H	11	1.28
(1,698)	1:31:C:PHE:HE1	2:35:D:GLY:HA3	3	1.28
(1,698)	1:31:C:PHE:HE2	2:35:D:GLY:HA3	3	1.28
(1,520)	2:17:B:ILE:HG13	2:40:D:PHE:HB3	13	1.28
(1,466)	1:4:A:GLY:HA2	1:5:A:HIS:HD2	6	1.28

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,443)	1:8:A:GLU:HG3	2:20:B:TYR:HD1	10	1.28
(1,443)	1:8:A:GLU:HG3	2:20:B:TYR:HD2	10	1.28
(1,384)	1:10:A:PHE:HE1	2:16:B:PRO:HD2	17	1.28
(1,384)	1:10:A:PHE:HE2	2:16:B:PRO:HD2	17	1.28
(1,201)	2:17:B:ILE:HG13	2:40:D:PHE:HB3	13	1.28
(1,147)	1:4:A:GLY:HA2	1:5:A:HIS:HD2	6	1.28
(1,124)	1:8:A:GLU:HG3	2:20:B:TYR:HD1	10	1.28
(1,124)	1:8:A:GLU:HG3	2:20:B:TYR:HD2	10	1.28
(1,65)	1:10:A:PHE:HE1	2:16:B:PRO:HD2	17	1.28
(1,65)	1:10:A:PHE:HE2	2:16:B:PRO:HD2	17	1.28
(1,1102)	1:29:C:GLU:HB3	2:41:D:TYR:HD1	14	1.27
(1,1102)	1:29:C:GLU:HB3	2:41:D:TYR:HD2	14	1.27
(1,1093)	1:29:C:GLU:HG2	2:41:D:TYR:HB2	15	1.27
(1,1093)	1:29:C:GLU:HG2	2:41:D:TYR:HB3	15	1.27
(1,783)	1:29:C:GLU:HB3	2:41:D:TYR:HD1	14	1.27
(1,783)	1:29:C:GLU:HB3	2:41:D:TYR:HD2	14	1.27
(1,774)	1:29:C:GLU:HG2	2:41:D:TYR:HB2	15	1.27
(1,774)	1:29:C:GLU:HG2	2:41:D:TYR:HB3	15	1.27
(1,596)	1:7:A:PRO:HG2	1:10:A:PHE:HE1	19	1.27
(1,596)	1:7:A:PRO:HG2	1:10:A:PHE:HE2	19	1.27
(1,596)	1:7:A:PRO:HG3	1:10:A:PHE:HE1	19	1.27
(1,596)	1:7:A:PRO:HG3	1:10:A:PHE:HE2	19	1.27
(1,453)	1:10:A:PHE:HD1	2:15:B:THR:HG21	1	1.27
(1,453)	1:10:A:PHE:HD1	2:15:B:THR:HG22	1	1.27
(1,453)	1:10:A:PHE:HD1	2:15:B:THR:HG23	1	1.27
(1,453)	1:10:A:PHE:HD2	2:15:B:THR:HG21	1	1.27
(1,453)	1:10:A:PHE:HD2	2:15:B:THR:HG22	1	1.27
(1,453)	1:10:A:PHE:HD2	2:15:B:THR:HG23	1	1.27
(1,445)	1:4:A:GLY:HA2	1:6:A:VAL:H	11	1.27
(1,438)	1:8:A:GLU:HG2	2:19:B:PHE:HA	5	1.27
(1,436)	1:8:A:GLU:HG2	2:20:B:TYR:HD1	6	1.27
(1,436)	1:8:A:GLU:HG2	2:20:B:TYR:HD2	6	1.27
(1,393)	1:7:A:PRO:HG2	1:10:A:PHE:HD1	12	1.27
(1,393)	1:7:A:PRO:HG2	1:10:A:PHE:HD2	12	1.27
(1,393)	1:7:A:PRO:HG3	1:10:A:PHE:HD1	12	1.27
(1,393)	1:7:A:PRO:HG3	1:10:A:PHE:HD2	12	1.27
(1,384)	1:10:A:PHE:HE1	2:16:B:PRO:HD2	5	1.27
(1,384)	1:10:A:PHE:HE2	2:16:B:PRO:HD2	5	1.27
(1,277)	1:7:A:PRO:HG2	1:10:A:PHE:HE1	19	1.27
(1,277)	1:7:A:PRO:HG2	1:10:A:PHE:HE2	19	1.27
(1,277)	1:7:A:PRO:HG3	1:10:A:PHE:HE1	19	1.27
(1,277)	1:7:A:PRO:HG3	1:10:A:PHE:HE2	19	1.27

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,134)	1:10:A:PHE:HD1	2:15:B:THR:HG21	1	1.27
(1,134)	1:10:A:PHE:HD1	2:15:B:THR:HG22	1	1.27
(1,134)	1:10:A:PHE:HD1	2:15:B:THR:HG23	1	1.27
(1,134)	1:10:A:PHE:HD2	2:15:B:THR:HG21	1	1.27
(1,134)	1:10:A:PHE:HD2	2:15:B:THR:HG22	1	1.27
(1,134)	1:10:A:PHE:HD2	2:15:B:THR:HG23	1	1.27
(1,126)	1:4:A:GLY:HA2	1:6:A:VAL:H	11	1.27
(1,119)	1:8:A:GLU:HG2	2:19:B:PHE:HA	5	1.27
(1,117)	1:8:A:GLU:HG2	2:20:B:TYR:HD1	6	1.27
(1,117)	1:8:A:GLU:HG2	2:20:B:TYR:HD2	6	1.27
(1,74)	1:7:A:PRO:HG2	1:10:A:PHE:HD1	12	1.27
(1,74)	1:7:A:PRO:HG2	1:10:A:PHE:HD2	12	1.27
(1,74)	1:7:A:PRO:HG3	1:10:A:PHE:HD1	12	1.27
(1,74)	1:7:A:PRO:HG3	1:10:A:PHE:HD2	12	1.27
(1,65)	1:10:A:PHE:HE1	2:16:B:PRO:HD2	5	1.27
(1,65)	1:10:A:PHE:HE2	2:16:B:PRO:HD2	5	1.27
(1,1102)	1:29:C:GLU:HB3	2:41:D:TYR:HD1	3	1.26
(1,1102)	1:29:C:GLU:HB3	2:41:D:TYR:HD2	3	1.26
(1,1092)	1:29:C:GLU:HG2	2:41:D:TYR:HD1	20	1.26
(1,1092)	1:29:C:GLU:HG2	2:41:D:TYR:HD2	20	1.26
(1,1036)	2:34:D:CYS:HB2	2:35:D:GLY:HA3	8	1.26
(1,783)	1:29:C:GLU:HB3	2:41:D:TYR:HD1	3	1.26
(1,783)	1:29:C:GLU:HB3	2:41:D:TYR:HD2	3	1.26
(1,773)	1:29:C:GLU:HG2	2:41:D:TYR:HD1	20	1.26
(1,773)	1:29:C:GLU:HG2	2:41:D:TYR:HD2	20	1.26
(1,717)	2:34:D:CYS:HB2	2:35:D:GLY:HA3	8	1.26
(1,595)	1:12:A:ARG:HA	1:12:A:ARG:HE	9	1.26
(1,466)	1:4:A:GLY:HA2	1:5:A:HIS:HD2	4	1.26
(1,453)	1:10:A:PHE:HD1	2:15:B:THR:HG21	8	1.26
(1,453)	1:10:A:PHE:HD1	2:15:B:THR:HG22	8	1.26
(1,453)	1:10:A:PHE:HD1	2:15:B:THR:HG23	8	1.26
(1,453)	1:10:A:PHE:HD2	2:15:B:THR:HG21	8	1.26
(1,453)	1:10:A:PHE:HD2	2:15:B:THR:HG22	8	1.26
(1,453)	1:10:A:PHE:HD2	2:15:B:THR:HG23	8	1.26
(1,276)	1:12:A:ARG:HA	1:12:A:ARG:HE	9	1.26
(1,147)	1:4:A:GLY:HA2	1:5:A:HIS:HD2	4	1.26
(1,134)	1:10:A:PHE:HD1	2:15:B:THR:HG21	8	1.26
(1,134)	1:10:A:PHE:HD1	2:15:B:THR:HG22	8	1.26
(1,134)	1:10:A:PHE:HD1	2:15:B:THR:HG23	8	1.26
(1,134)	1:10:A:PHE:HD2	2:15:B:THR:HG21	8	1.26
(1,134)	1:10:A:PHE:HD2	2:15:B:THR:HG22	8	1.26
(1,134)	1:10:A:PHE:HD2	2:15:B:THR:HG23	8	1.26

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,1251)	1:33:C:ARG:HA	1:33:C:ARG:HE	9	1.25
(1,1101)	1:25:C:GLY:HA2	1:27:C:VAL:H	7	1.25
(1,1067)	1:26:C:HIS:HB3	1:27:C:VAL:HG21	1	1.25
(1,1067)	1:26:C:HIS:HB3	1:27:C:VAL:HG22	1	1.25
(1,1067)	1:26:C:HIS:HB3	1:27:C:VAL:HG23	1	1.25
(1,1067)	1:26:C:HIS:HB3	1:27:C:VAL:HG21	16	1.25
(1,1067)	1:26:C:HIS:HB3	1:27:C:VAL:HG22	16	1.25
(1,1067)	1:26:C:HIS:HB3	1:27:C:VAL:HG23	16	1.25
(1,932)	1:33:C:ARG:HA	1:33:C:ARG:HE	9	1.25
(1,782)	1:25:C:GLY:HA2	1:27:C:VAL:H	7	1.25
(1,748)	1:26:C:HIS:HB3	1:27:C:VAL:HG21	1	1.25
(1,748)	1:26:C:HIS:HB3	1:27:C:VAL:HG22	1	1.25
(1,748)	1:26:C:HIS:HB3	1:27:C:VAL:HG23	1	1.25
(1,748)	1:26:C:HIS:HB3	1:27:C:VAL:HG21	16	1.25
(1,748)	1:26:C:HIS:HB3	1:27:C:VAL:HG22	16	1.25
(1,748)	1:26:C:HIS:HB3	1:27:C:VAL:HG23	16	1.25
(1,673)	2:17:B:ILE:HG13	2:40:D:PHE:HB2	7	1.25
(1,661)	2:19:B:PHE:H	2:38:D:ILE:HD11	17	1.25
(1,661)	2:19:B:PHE:H	2:38:D:ILE:HD12	17	1.25
(1,661)	2:19:B:PHE:H	2:38:D:ILE:HD13	17	1.25
(1,405)	1:1:A:GLY:H	2:18:B:SER:HB2	10	1.25
(1,397)	2:18:B:SER:HB2	2:19:B:PHE:H	5	1.25
(1,348)	1:11:A:VAL:HA	1:12:A:ARG:HD2	16	1.25
(1,348)	1:11:A:VAL:HA	1:12:A:ARG:HD3	16	1.25
(1,86)	1:1:A:GLY:H	2:18:B:SER:HB2	10	1.25
(1,78)	2:18:B:SER:HB2	2:19:B:PHE:H	5	1.25
(1,29)	1:11:A:VAL:HA	1:12:A:ARG:HD2	16	1.25
(1,29)	1:11:A:VAL:HA	1:12:A:ARG:HD3	16	1.25
(1,14)	2:19:B:PHE:H	2:38:D:ILE:HD11	17	1.25
(1,14)	2:19:B:PHE:H	2:38:D:ILE:HD12	17	1.25
(1,14)	2:19:B:PHE:H	2:38:D:ILE:HD13	17	1.25
(1,8)	2:17:B:ILE:HG13	2:40:D:PHE:HB2	7	1.25
(1,1175)	1:24:C:ALA:HB1	2:40:D:PHE:HB3	9	1.24
(1,1175)	1:24:C:ALA:HB2	2:40:D:PHE:HB3	9	1.24
(1,1175)	1:24:C:ALA:HB3	2:40:D:PHE:HB3	9	1.24
(1,1102)	1:29:C:GLU:HB3	2:41:D:TYR:HD1	1	1.24
(1,1102)	1:29:C:GLU:HB3	2:41:D:TYR:HD2	1	1.24
(1,856)	1:24:C:ALA:HB1	2:40:D:PHE:HB3	9	1.24
(1,856)	1:24:C:ALA:HB2	2:40:D:PHE:HB3	9	1.24
(1,856)	1:24:C:ALA:HB3	2:40:D:PHE:HB3	9	1.24
(1,783)	1:29:C:GLU:HB3	2:41:D:TYR:HD1	1	1.24
(1,783)	1:29:C:GLU:HB3	2:41:D:TYR:HD2	1	1.24

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,653)	1:3:A:ALA:HA	2:19:B:PHE:HD1	4	1.24
(1,653)	1:3:A:ALA:HA	2:19:B:PHE:HD2	4	1.24
(1,520)	2:17:B:ILE:HG13	2:40:D:PHE:HB3	19	1.24
(1,501)	1:4:A:GLY:H	1:6:A:VAL:H	14	1.24
(1,391)	1:3:A:ALA:HA	2:19:B:PHE:HD1	4	1.24
(1,391)	1:3:A:ALA:HA	2:19:B:PHE:HD2	4	1.24
(1,334)	1:3:A:ALA:HA	2:19:B:PHE:HD1	4	1.24
(1,334)	1:3:A:ALA:HA	2:19:B:PHE:HD2	4	1.24
(1,201)	2:17:B:ILE:HG13	2:40:D:PHE:HB3	19	1.24
(1,182)	1:4:A:GLY:H	1:6:A:VAL:H	14	1.24
(1,72)	1:3:A:ALA:HA	2:19:B:PHE:HD1	4	1.24
(1,72)	1:3:A:ALA:HA	2:19:B:PHE:HD2	4	1.24
(1,1176)	2:19:B:PHE:HB3	2:38:D:ILE:HG13	4	1.23
(1,1071)	1:27:C:VAL:HG11	2:41:D:TYR:HE1	14	1.23
(1,1071)	1:27:C:VAL:HG11	2:41:D:TYR:HE2	14	1.23
(1,1071)	1:27:C:VAL:HG12	2:41:D:TYR:HE1	14	1.23
(1,1071)	1:27:C:VAL:HG12	2:41:D:TYR:HE2	14	1.23
(1,1071)	1:27:C:VAL:HG13	2:41:D:TYR:HE1	14	1.23
(1,1071)	1:27:C:VAL:HG13	2:41:D:TYR:HE2	14	1.23
(1,857)	2:19:B:PHE:HB3	2:38:D:ILE:HG13	4	1.23
(1,752)	1:27:C:VAL:HG11	2:41:D:TYR:HE1	14	1.23
(1,752)	1:27:C:VAL:HG11	2:41:D:TYR:HE2	14	1.23
(1,752)	1:27:C:VAL:HG12	2:41:D:TYR:HE1	14	1.23
(1,752)	1:27:C:VAL:HG12	2:41:D:TYR:HE2	14	1.23
(1,752)	1:27:C:VAL:HG13	2:41:D:TYR:HE1	14	1.23
(1,752)	1:27:C:VAL:HG13	2:41:D:TYR:HE2	14	1.23
(1,672)	2:17:B:ILE:HG13	2:40:D:PHE:HD1	16	1.23
(1,672)	2:17:B:ILE:HG13	2:40:D:PHE:HD2	16	1.23
(1,7)	2:17:B:ILE:HG13	2:40:D:PHE:HD1	16	1.23
(1,7)	2:17:B:ILE:HG13	2:40:D:PHE:HD2	16	1.23
(1,1122)	1:25:C:GLY:HA2	1:26:C:HIS:HD2	15	1.22
(1,1046)	1:27:C:VAL:HB	2:41:D:TYR:HE1	15	1.22
(1,1046)	1:27:C:VAL:HB	2:41:D:TYR:HE2	15	1.22
(1,803)	1:25:C:GLY:HA2	1:26:C:HIS:HD2	15	1.22
(1,727)	1:27:C:VAL:HB	2:41:D:TYR:HE1	15	1.22
(1,727)	1:27:C:VAL:HB	2:41:D:TYR:HE2	15	1.22
(1,669)	2:13:B:CYS:HB3	1:25:C:GLY:HA2	20	1.22
(1,503)	1:7:A:PRO:HB3	1:10:A:PHE:HE1	4	1.22
(1,503)	1:7:A:PRO:HB3	1:10:A:PHE:HE2	4	1.22
(1,446)	1:8:A:GLU:HB3	2:20:B:TYR:HD1	13	1.22
(1,446)	1:8:A:GLU:HB3	2:20:B:TYR:HD2	13	1.22
(1,422)	1:4:A:GLY:H	2:20:B:TYR:H	2	1.22

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,390)	1:6:A:VAL:HB	2:20:B:TYR:HE1	11	1.22
(1,390)	1:6:A:VAL:HB	2:20:B:TYR:HE2	11	1.22
(1,184)	1:7:A:PRO:HB3	1:10:A:PHE:HE1	4	1.22
(1,184)	1:7:A:PRO:HB3	1:10:A:PHE:HE2	4	1.22
(1,127)	1:8:A:GLU:HB3	2:20:B:TYR:HD1	13	1.22
(1,127)	1:8:A:GLU:HB3	2:20:B:TYR:HD2	13	1.22
(1,103)	1:4:A:GLY:H	2:20:B:TYR:H	2	1.22
(1,71)	1:6:A:VAL:HB	2:20:B:TYR:HE1	11	1.22
(1,71)	1:6:A:VAL:HB	2:20:B:TYR:HE2	11	1.22
(1,4)	2:13:B:CYS:HB3	1:25:C:GLY:HA2	20	1.22
(1,1251)	1:33:C:ARG:HA	1:33:C:ARG:HE	2	1.21
(1,1171)	2:36:D:THR:HG21	2:38:D:ILE:HD11	1	1.21
(1,1171)	2:36:D:THR:HG21	2:38:D:ILE:HD12	1	1.21
(1,1171)	2:36:D:THR:HG21	2:38:D:ILE:HD13	1	1.21
(1,1171)	2:36:D:THR:HG22	2:38:D:ILE:HD11	1	1.21
(1,1171)	2:36:D:THR:HG22	2:38:D:ILE:HD12	1	1.21
(1,1171)	2:36:D:THR:HG22	2:38:D:ILE:HD13	1	1.21
(1,1171)	2:36:D:THR:HG23	2:38:D:ILE:HD11	1	1.21
(1,1171)	2:36:D:THR:HG23	2:38:D:ILE:HD12	1	1.21
(1,1171)	2:36:D:THR:HG23	2:38:D:ILE:HD13	1	1.21
(1,1017)	1:31:C:PHE:HE1	2:35:D:GLY:HA3	10	1.21
(1,1017)	1:31:C:PHE:HE2	2:35:D:GLY:HA3	10	1.21
(1,932)	1:33:C:ARG:HA	1:33:C:ARG:HE	2	1.21
(1,852)	2:36:D:THR:HG21	2:38:D:ILE:HD11	1	1.21
(1,852)	2:36:D:THR:HG21	2:38:D:ILE:HD12	1	1.21
(1,852)	2:36:D:THR:HG21	2:38:D:ILE:HD13	1	1.21
(1,852)	2:36:D:THR:HG22	2:38:D:ILE:HD11	1	1.21
(1,852)	2:36:D:THR:HG22	2:38:D:ILE:HD12	1	1.21
(1,852)	2:36:D:THR:HG22	2:38:D:ILE:HD13	1	1.21
(1,852)	2:36:D:THR:HG23	2:38:D:ILE:HD11	1	1.21
(1,852)	2:36:D:THR:HG23	2:38:D:ILE:HD12	1	1.21
(1,852)	2:36:D:THR:HG23	2:38:D:ILE:HD13	1	1.21
(1,698)	1:31:C:PHE:HE1	2:35:D:GLY:HA3	10	1.21
(1,698)	1:31:C:PHE:HE2	2:35:D:GLY:HA3	10	1.21
(1,670)	2:17:B:ILE:HD11	2:40:D:PHE:H	7	1.21
(1,670)	2:17:B:ILE:HD12	2:40:D:PHE:H	7	1.21
(1,670)	2:17:B:ILE:HD13	2:40:D:PHE:H	7	1.21
(1,661)	2:19:B:PHE:H	2:38:D:ILE:HD11	6	1.21
(1,661)	2:19:B:PHE:H	2:38:D:ILE:HD12	6	1.21
(1,661)	2:19:B:PHE:H	2:38:D:ILE:HD13	6	1.21
(1,661)	2:19:B:PHE:H	2:38:D:ILE:HD11	16	1.21
(1,661)	2:19:B:PHE:H	2:38:D:ILE:HD12	16	1.21

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,661)	2:19:B:PHE:H	2:38:D:ILE:HD13	16	1.21
(1,593)	1:12:A:ARG:HB2	1:12:A:ARG:HE	5	1.21
(1,593)	1:12:A:ARG:HB2	1:12:A:ARG:HE	8	1.21
(1,520)	2:17:B:ILE:HG13	2:40:D:PHE:HB3	20	1.21
(1,438)	1:8:A:GLU:HG2	2:19:B:PHE:HA	6	1.21
(1,274)	1:12:A:ARG:HB2	1:12:A:ARG:HE	5	1.21
(1,274)	1:12:A:ARG:HB2	1:12:A:ARG:HE	8	1.21
(1,201)	2:17:B:ILE:HG13	2:40:D:PHE:HB3	20	1.21
(1,119)	1:8:A:GLU:HG2	2:19:B:PHE:HA	6	1.21
(1,14)	2:19:B:PHE:H	2:38:D:ILE:HD11	6	1.21
(1,14)	2:19:B:PHE:H	2:38:D:ILE:HD12	6	1.21
(1,14)	2:19:B:PHE:H	2:38:D:ILE:HD13	6	1.21
(1,14)	2:19:B:PHE:H	2:38:D:ILE:HD11	16	1.21
(1,14)	2:19:B:PHE:H	2:38:D:ILE:HD12	16	1.21
(1,14)	2:19:B:PHE:H	2:38:D:ILE:HD13	16	1.21
(1,5)	2:17:B:ILE:HD11	2:40:D:PHE:H	7	1.21
(1,5)	2:17:B:ILE:HD12	2:40:D:PHE:H	7	1.21
(1,5)	2:17:B:ILE:HD13	2:40:D:PHE:H	7	1.21
(1,663)	2:19:B:PHE:HD1	2:38:D:ILE:HG13	12	1.2
(1,663)	2:19:B:PHE:HD2	2:38:D:ILE:HG13	12	1.2
(1,16)	2:19:B:PHE:HD1	2:38:D:ILE:HG13	12	1.2
(1,16)	2:19:B:PHE:HD2	2:38:D:ILE:HG13	12	1.2
(1,1251)	1:33:C:ARG:HA	1:33:C:ARG:HE	16	1.19
(1,1093)	1:29:C:GLU:HG2	2:41:D:TYR:HB2	10	1.19
(1,1093)	1:29:C:GLU:HG2	2:41:D:TYR:HB3	10	1.19
(1,1093)	1:29:C:GLU:HG2	2:41:D:TYR:HB2	14	1.19
(1,1093)	1:29:C:GLU:HG2	2:41:D:TYR:HB3	14	1.19
(1,1092)	1:29:C:GLU:HG2	2:41:D:TYR:HD1	6	1.19
(1,1092)	1:29:C:GLU:HG2	2:41:D:TYR:HD2	6	1.19
(1,1071)	1:27:C:VAL:HG11	2:41:D:TYR:HE1	19	1.19
(1,1071)	1:27:C:VAL:HG11	2:41:D:TYR:HE2	19	1.19
(1,1071)	1:27:C:VAL:HG12	2:41:D:TYR:HE1	19	1.19
(1,1071)	1:27:C:VAL:HG12	2:41:D:TYR:HE2	19	1.19
(1,1071)	1:27:C:VAL:HG13	2:41:D:TYR:HE1	19	1.19
(1,1071)	1:27:C:VAL:HG13	2:41:D:TYR:HE2	19	1.19
(1,1050)	1:28:C:PRO:HG2	1:31:C:PHE:HB2	11	1.19
(1,1050)	1:28:C:PRO:HG3	1:31:C:PHE:HB2	11	1.19
(1,1016)	1:28:C:PRO:HG2	1:31:C:PHE:HB2	11	1.19
(1,1016)	1:28:C:PRO:HG3	1:31:C:PHE:HB2	11	1.19
(1,932)	1:33:C:ARG:HA	1:33:C:ARG:HE	16	1.19
(1,774)	1:29:C:GLU:HG2	2:41:D:TYR:HB2	10	1.19
(1,774)	1:29:C:GLU:HG2	2:41:D:TYR:HB3	10	1.19

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,774)	1:29:C:GLU:HG2	2:41:D:TYR:HB2	14	1.19
(1,774)	1:29:C:GLU:HG2	2:41:D:TYR:HB3	14	1.19
(1,773)	1:29:C:GLU:HG2	2:41:D:TYR:HD1	6	1.19
(1,773)	1:29:C:GLU:HG2	2:41:D:TYR:HD2	6	1.19
(1,752)	1:27:C:VAL:HG11	2:41:D:TYR:HE1	19	1.19
(1,752)	1:27:C:VAL:HG11	2:41:D:TYR:HE2	19	1.19
(1,752)	1:27:C:VAL:HG12	2:41:D:TYR:HE1	19	1.19
(1,752)	1:27:C:VAL:HG12	2:41:D:TYR:HE2	19	1.19
(1,752)	1:27:C:VAL:HG13	2:41:D:TYR:HE1	19	1.19
(1,752)	1:27:C:VAL:HG13	2:41:D:TYR:HE2	19	1.19
(1,731)	1:28:C:PRO:HG2	1:31:C:PHE:HB2	11	1.19
(1,731)	1:28:C:PRO:HG3	1:31:C:PHE:HB2	11	1.19
(1,697)	1:28:C:PRO:HG2	1:31:C:PHE:HB2	11	1.19
(1,697)	1:28:C:PRO:HG3	1:31:C:PHE:HB2	11	1.19
(1,1092)	1:29:C:GLU:HG2	2:41:D:TYR:HD1	3	1.18
(1,1092)	1:29:C:GLU:HG2	2:41:D:TYR:HD2	3	1.18
(1,1092)	1:29:C:GLU:HG2	2:41:D:TYR:HD1	8	1.18
(1,1092)	1:29:C:GLU:HG2	2:41:D:TYR:HD2	8	1.18
(1,1076)	1:25:C:GLY:HA3	1:26:C:HIS:HD2	12	1.18
(1,1008)	2:18:B:SER:HA	2:38:D:ILE:HG13	17	1.18
(1,1007)	2:37:D:PRO:HA	2:38:D:ILE:HD11	15	1.18
(1,1007)	2:37:D:PRO:HA	2:38:D:ILE:HD12	15	1.18
(1,1007)	2:37:D:PRO:HA	2:38:D:ILE:HD13	15	1.18
(1,773)	1:29:C:GLU:HG2	2:41:D:TYR:HD1	3	1.18
(1,773)	1:29:C:GLU:HG2	2:41:D:TYR:HD2	3	1.18
(1,773)	1:29:C:GLU:HG2	2:41:D:TYR:HD1	8	1.18
(1,773)	1:29:C:GLU:HG2	2:41:D:TYR:HD2	8	1.18
(1,757)	1:25:C:GLY:HA3	1:26:C:HIS:HD2	12	1.18
(1,689)	2:18:B:SER:HA	2:38:D:ILE:HG13	17	1.18
(1,688)	2:37:D:PRO:HA	2:38:D:ILE:HD11	15	1.18
(1,688)	2:37:D:PRO:HA	2:38:D:ILE:HD12	15	1.18
(1,688)	2:37:D:PRO:HA	2:38:D:ILE:HD13	15	1.18
(1,503)	1:7:A:PRO:HB3	1:10:A:PHE:HE1	20	1.18
(1,503)	1:7:A:PRO:HB3	1:10:A:PHE:HE2	20	1.18
(1,483)	1:1:A:GLY:H	2:20:B:TYR:HD1	7	1.18
(1,483)	1:1:A:GLY:H	2:20:B:TYR:HD2	7	1.18
(1,184)	1:7:A:PRO:HB3	1:10:A:PHE:HE1	20	1.18
(1,184)	1:7:A:PRO:HB3	1:10:A:PHE:HE2	20	1.18
(1,164)	1:1:A:GLY:H	2:20:B:TYR:HD1	7	1.18
(1,164)	1:1:A:GLY:H	2:20:B:TYR:HD2	7	1.18
(1,1074)	1:22:C:GLY:H	2:39:D:SER:HB3	1	1.17
(1,755)	1:22:C:GLY:H	2:39:D:SER:HB3	1	1.17

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,661)	2:19:B:PHE:H	2:38:D:ILE:HD11	7	1.17
(1,661)	2:19:B:PHE:H	2:38:D:ILE:HD12	7	1.17
(1,661)	2:19:B:PHE:H	2:38:D:ILE:HD13	7	1.17
(1,489)	1:4:A:GLY:H	2:19:B:PHE:HE1	3	1.17
(1,489)	1:4:A:GLY:H	2:19:B:PHE:HE2	3	1.17
(1,415)	1:6:A:VAL:HG11	2:20:B:TYR:HE1	17	1.17
(1,415)	1:6:A:VAL:HG11	2:20:B:TYR:HE2	17	1.17
(1,415)	1:6:A:VAL:HG12	2:20:B:TYR:HE1	17	1.17
(1,415)	1:6:A:VAL:HG12	2:20:B:TYR:HE2	17	1.17
(1,415)	1:6:A:VAL:HG13	2:20:B:TYR:HE1	17	1.17
(1,415)	1:6:A:VAL:HG13	2:20:B:TYR:HE2	17	1.17
(1,390)	1:6:A:VAL:HB	2:20:B:TYR:HE1	4	1.17
(1,390)	1:6:A:VAL:HB	2:20:B:TYR:HE2	4	1.17
(1,380)	2:13:B:CYS:HB2	2:14:B:GLY:HA3	17	1.17
(1,170)	1:4:A:GLY:H	2:19:B:PHE:HE1	3	1.17
(1,170)	1:4:A:GLY:H	2:19:B:PHE:HE2	3	1.17
(1,96)	1:6:A:VAL:HG11	2:20:B:TYR:HE1	17	1.17
(1,96)	1:6:A:VAL:HG11	2:20:B:TYR:HE2	17	1.17
(1,96)	1:6:A:VAL:HG12	2:20:B:TYR:HE1	17	1.17
(1,96)	1:6:A:VAL:HG12	2:20:B:TYR:HE2	17	1.17
(1,96)	1:6:A:VAL:HG13	2:20:B:TYR:HE1	17	1.17
(1,96)	1:6:A:VAL:HG13	2:20:B:TYR:HE2	17	1.17
(1,71)	1:6:A:VAL:HB	2:20:B:TYR:HE1	4	1.17
(1,71)	1:6:A:VAL:HB	2:20:B:TYR:HE2	4	1.17
(1,61)	2:13:B:CYS:HB2	2:14:B:GLY:HA3	17	1.17
(1,14)	2:19:B:PHE:H	2:38:D:ILE:HD11	7	1.17
(1,14)	2:19:B:PHE:H	2:38:D:ILE:HD12	7	1.17
(1,14)	2:19:B:PHE:H	2:38:D:ILE:HD13	7	1.17
(1,673)	2:17:B:ILE:HG13	2:40:D:PHE:HB2	8	1.16
(1,513)	2:15:B:THR:HB	2:17:B:ILE:HD11	6	1.16
(1,513)	2:15:B:THR:HB	2:17:B:ILE:HD12	6	1.16
(1,513)	2:15:B:THR:HB	2:17:B:ILE:HD13	6	1.16
(1,501)	1:4:A:GLY:H	1:6:A:VAL:H	2	1.16
(1,390)	1:6:A:VAL:HB	2:20:B:TYR:HE1	12	1.16
(1,390)	1:6:A:VAL:HB	2:20:B:TYR:HE2	12	1.16
(1,194)	2:15:B:THR:HB	2:17:B:ILE:HD11	6	1.16
(1,194)	2:15:B:THR:HB	2:17:B:ILE:HD12	6	1.16
(1,194)	2:15:B:THR:HB	2:17:B:ILE:HD13	6	1.16
(1,182)	1:4:A:GLY:H	1:6:A:VAL:H	2	1.16
(1,71)	1:6:A:VAL:HB	2:20:B:TYR:HE1	12	1.16
(1,71)	1:6:A:VAL:HB	2:20:B:TYR:HE2	12	1.16
(1,8)	2:17:B:ILE:HG13	2:40:D:PHE:HB2	8	1.16

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,1076)	1:25:C:GLY:HA3	1:26:C:HIS:HD2	4	1.15
(1,757)	1:25:C:GLY:HA3	1:26:C:HIS:HD2	4	1.15
(1,453)	1:10:A:PHE:HD1	2:15:B:THR:HG21	19	1.15
(1,453)	1:10:A:PHE:HD1	2:15:B:THR:HG22	19	1.15
(1,453)	1:10:A:PHE:HD1	2:15:B:THR:HG23	19	1.15
(1,453)	1:10:A:PHE:HD2	2:15:B:THR:HG21	19	1.15
(1,453)	1:10:A:PHE:HD2	2:15:B:THR:HG22	19	1.15
(1,453)	1:10:A:PHE:HD2	2:15:B:THR:HG23	19	1.15
(1,134)	1:10:A:PHE:HD1	2:15:B:THR:HG21	19	1.15
(1,134)	1:10:A:PHE:HD1	2:15:B:THR:HG22	19	1.15
(1,134)	1:10:A:PHE:HD1	2:15:B:THR:HG23	19	1.15
(1,134)	1:10:A:PHE:HD2	2:15:B:THR:HG21	19	1.15
(1,134)	1:10:A:PHE:HD2	2:15:B:THR:HG22	19	1.15
(1,134)	1:10:A:PHE:HD2	2:15:B:THR:HG23	19	1.15
(1,1102)	1:29:C:GLU:HB3	2:41:D:TYR:HD1	11	1.14
(1,1102)	1:29:C:GLU:HB3	2:41:D:TYR:HD2	11	1.14
(1,1045)	1:27:C:VAL:HB	2:41:D:TYR:HD1	15	1.14
(1,1045)	1:27:C:VAL:HB	2:41:D:TYR:HD2	15	1.14
(1,783)	1:29:C:GLU:HB3	2:41:D:TYR:HD1	11	1.14
(1,783)	1:29:C:GLU:HB3	2:41:D:TYR:HD2	11	1.14
(1,726)	1:27:C:VAL:HB	2:41:D:TYR:HD1	15	1.14
(1,726)	1:27:C:VAL:HB	2:41:D:TYR:HD2	15	1.14
(1,672)	2:17:B:ILE:HG13	2:40:D:PHE:HD1	7	1.14
(1,672)	2:17:B:ILE:HG13	2:40:D:PHE:HD2	7	1.14
(1,664)	2:19:B:PHE:HB2	2:38:D:ILE:HG13	2	1.14
(1,451)	2:15:B:THR:HG21	2:16:B:PRO:HB3	3	1.14
(1,451)	2:15:B:THR:HG22	2:16:B:PRO:HB3	3	1.14
(1,451)	2:15:B:THR:HG23	2:16:B:PRO:HB3	3	1.14
(1,415)	1:6:A:VAL:HG11	2:20:B:TYR:HE1	6	1.14
(1,415)	1:6:A:VAL:HG11	2:20:B:TYR:HE2	6	1.14
(1,415)	1:6:A:VAL:HG12	2:20:B:TYR:HE1	6	1.14
(1,415)	1:6:A:VAL:HG12	2:20:B:TYR:HE2	6	1.14
(1,415)	1:6:A:VAL:HG13	2:20:B:TYR:HE1	6	1.14
(1,415)	1:6:A:VAL:HG13	2:20:B:TYR:HE2	6	1.14
(1,390)	1:6:A:VAL:HB	2:20:B:TYR:HE1	10	1.14
(1,390)	1:6:A:VAL:HB	2:20:B:TYR:HE2	10	1.14
(1,132)	2:15:B:THR:HG21	2:16:B:PRO:HB3	3	1.14
(1,132)	2:15:B:THR:HG22	2:16:B:PRO:HB3	3	1.14
(1,132)	2:15:B:THR:HG23	2:16:B:PRO:HB3	3	1.14
(1,96)	1:6:A:VAL:HG11	2:20:B:TYR:HE1	6	1.14
(1,96)	1:6:A:VAL:HG11	2:20:B:TYR:HE2	6	1.14
(1,96)	1:6:A:VAL:HG12	2:20:B:TYR:HE1	6	1.14

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,96)	1:6:A:VAL:HG12	2:20:B:TYR:HE2	6	1.14
(1,96)	1:6:A:VAL:HG13	2:20:B:TYR:HE1	6	1.14
(1,96)	1:6:A:VAL:HG13	2:20:B:TYR:HE2	6	1.14
(1,71)	1:6:A:VAL:HB	2:20:B:TYR:HE1	10	1.14
(1,71)	1:6:A:VAL:HB	2:20:B:TYR:HE2	10	1.14
(1,17)	2:19:B:PHE:HB2	2:38:D:ILE:HG13	2	1.14
(1,7)	2:17:B:ILE:HG13	2:40:D:PHE:HD1	7	1.14
(1,7)	2:17:B:ILE:HG13	2:40:D:PHE:HD2	7	1.14
(1,1008)	2:18:B:SER:HA	2:38:D:ILE:HG13	19	1.13
(1,689)	2:18:B:SER:HA	2:38:D:ILE:HG13	19	1.13
(1,513)	2:15:B:THR:HB	2:17:B:ILE:HD11	8	1.13
(1,513)	2:15:B:THR:HB	2:17:B:ILE:HD12	8	1.13
(1,513)	2:15:B:THR:HB	2:17:B:ILE:HD13	8	1.13
(1,359)	1:9:A:TYR:HD1	2:18:B:SER:HA	13	1.13
(1,359)	1:9:A:TYR:HD2	2:18:B:SER:HA	13	1.13
(1,194)	2:15:B:THR:HB	2:17:B:ILE:HD11	8	1.13
(1,194)	2:15:B:THR:HB	2:17:B:ILE:HD12	8	1.13
(1,194)	2:15:B:THR:HB	2:17:B:ILE:HD13	8	1.13
(1,40)	1:9:A:TYR:HD1	2:18:B:SER:HA	13	1.13
(1,40)	1:9:A:TYR:HD2	2:18:B:SER:HA	13	1.13
(1,1137)	1:22:C:GLY:H	2:40:D:PHE:H	14	1.12
(1,1004)	1:32:C:VAL:HA	1:33:C:ARG:HD2	13	1.12
(1,1004)	1:32:C:VAL:HA	1:33:C:ARG:HD3	13	1.12
(1,1001)	1:30:C:TYR:HD1	2:39:D:SER:HB2	12	1.12
(1,1001)	1:30:C:TYR:HD2	2:39:D:SER:HB2	12	1.12
(1,818)	1:22:C:GLY:H	2:40:D:PHE:H	14	1.12
(1,685)	1:32:C:VAL:HA	1:33:C:ARG:HD2	13	1.12
(1,685)	1:32:C:VAL:HA	1:33:C:ARG:HD3	13	1.12
(1,682)	1:30:C:TYR:HD1	2:39:D:SER:HB2	12	1.12
(1,682)	1:30:C:TYR:HD2	2:39:D:SER:HB2	12	1.12
(1,519)	1:3:A:ALA:HB1	2:19:B:PHE:HB3	19	1.12
(1,519)	1:3:A:ALA:HB2	2:19:B:PHE:HB3	19	1.12
(1,519)	1:3:A:ALA:HB3	2:19:B:PHE:HB3	19	1.12
(1,443)	1:8:A:GLU:HG3	2:20:B:TYR:HD1	5	1.12
(1,443)	1:8:A:GLU:HG3	2:20:B:TYR:HD2	5	1.12
(1,342)	1:1:A:GLY:HA3	2:18:B:SER:HB2	17	1.12
(1,339)	1:1:A:GLY:HA3	2:18:B:SER:HB2	17	1.12
(1,200)	1:3:A:ALA:HB1	2:19:B:PHE:HB3	19	1.12
(1,200)	1:3:A:ALA:HB2	2:19:B:PHE:HB3	19	1.12
(1,200)	1:3:A:ALA:HB3	2:19:B:PHE:HB3	19	1.12
(1,124)	1:8:A:GLU:HG3	2:20:B:TYR:HD1	5	1.12
(1,124)	1:8:A:GLU:HG3	2:20:B:TYR:HD2	5	1.12

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,23)	1:1:A:GLY:HA3	2:18:B:SER:HB2	17	1.12
(1,20)	1:1:A:GLY:HA3	2:18:B:SER:HB2	17	1.12
(1,1157)	1:25:C:GLY:H	1:27:C:VAL:H	15	1.11
(1,1122)	1:25:C:GLY:HA2	1:26:C:HIS:HD2	19	1.11
(1,1061)	1:22:C:GLY:H	2:39:D:SER:HB2	19	1.11
(1,1046)	1:27:C:VAL:HB	2:41:D:TYR:HE1	9	1.11
(1,1046)	1:27:C:VAL:HB	2:41:D:TYR:HE2	9	1.11
(1,1025)	1:27:C:VAL:HG21	2:41:D:TYR:HE1	11	1.11
(1,1025)	1:27:C:VAL:HG21	2:41:D:TYR:HE2	11	1.11
(1,1025)	1:27:C:VAL:HG22	2:41:D:TYR:HE1	11	1.11
(1,1025)	1:27:C:VAL:HG22	2:41:D:TYR:HE2	11	1.11
(1,1025)	1:27:C:VAL:HG23	2:41:D:TYR:HE1	11	1.11
(1,1025)	1:27:C:VAL:HG23	2:41:D:TYR:HE2	11	1.11
(1,838)	1:25:C:GLY:H	1:27:C:VAL:H	15	1.11
(1,803)	1:25:C:GLY:HA2	1:26:C:HIS:HD2	19	1.11
(1,742)	1:22:C:GLY:H	2:39:D:SER:HB2	19	1.11
(1,727)	1:27:C:VAL:HB	2:41:D:TYR:HE1	9	1.11
(1,727)	1:27:C:VAL:HB	2:41:D:TYR:HE2	9	1.11
(1,706)	1:27:C:VAL:HG21	2:41:D:TYR:HE1	11	1.11
(1,706)	1:27:C:VAL:HG21	2:41:D:TYR:HE2	11	1.11
(1,706)	1:27:C:VAL:HG22	2:41:D:TYR:HE1	11	1.11
(1,706)	1:27:C:VAL:HG22	2:41:D:TYR:HE2	11	1.11
(1,706)	1:27:C:VAL:HG23	2:41:D:TYR:HE1	11	1.11
(1,706)	1:27:C:VAL:HG23	2:41:D:TYR:HE2	11	1.11
(1,587)	1:12:A:ARG:H	1:12:A:ARG:HE	12	1.11
(1,501)	1:4:A:GLY:H	1:6:A:VAL:H	1	1.11
(1,268)	1:12:A:ARG:H	1:12:A:ARG:HE	12	1.11
(1,182)	1:4:A:GLY:H	1:6:A:VAL:H	1	1.11
(1,1171)	2:36:D:THR:HG21	2:38:D:ILE:HD11	16	1.1
(1,1171)	2:36:D:THR:HG21	2:38:D:ILE:HD12	16	1.1
(1,1171)	2:36:D:THR:HG21	2:38:D:ILE:HD13	16	1.1
(1,1171)	2:36:D:THR:HG22	2:38:D:ILE:HD11	16	1.1
(1,1171)	2:36:D:THR:HG22	2:38:D:ILE:HD12	16	1.1
(1,1171)	2:36:D:THR:HG22	2:38:D:ILE:HD13	16	1.1
(1,1171)	2:36:D:THR:HG23	2:38:D:ILE:HD11	16	1.1
(1,1171)	2:36:D:THR:HG23	2:38:D:ILE:HD12	16	1.1
(1,1171)	2:36:D:THR:HG23	2:38:D:ILE:HD13	16	1.1
(1,1159)	1:28:C:PRO:HB3	1:31:C:PHE:HE1	15	1.1
(1,1159)	1:28:C:PRO:HB3	1:31:C:PHE:HE2	15	1.1
(1,1043)	1:27:C:VAL:HG11	1:28:C:PRO:HG2	16	1.1
(1,1043)	1:27:C:VAL:HG11	1:28:C:PRO:HG3	16	1.1
(1,1043)	1:27:C:VAL:HG12	1:28:C:PRO:HG2	16	1.1

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,1043)	1:27:C:VAL:HG12	1:28:C:PRO:HG3	16	1.1
(1,1043)	1:27:C:VAL:HG13	1:28:C:PRO:HG2	16	1.1
(1,1043)	1:27:C:VAL:HG13	1:28:C:PRO:HG3	16	1.1
(1,1036)	2:34:D:CYS:HB2	2:35:D:GLY:HA3	2	1.1
(1,1004)	1:32:C:VAL:HA	1:33:C:ARG:HD2	19	1.1
(1,1004)	1:32:C:VAL:HA	1:33:C:ARG:HD3	19	1.1
(1,852)	2:36:D:THR:HG21	2:38:D:ILE:HD11	16	1.1
(1,852)	2:36:D:THR:HG21	2:38:D:ILE:HD12	16	1.1
(1,852)	2:36:D:THR:HG21	2:38:D:ILE:HD13	16	1.1
(1,852)	2:36:D:THR:HG22	2:38:D:ILE:HD11	16	1.1
(1,852)	2:36:D:THR:HG22	2:38:D:ILE:HD12	16	1.1
(1,852)	2:36:D:THR:HG22	2:38:D:ILE:HD13	16	1.1
(1,852)	2:36:D:THR:HG23	2:38:D:ILE:HD11	16	1.1
(1,852)	2:36:D:THR:HG23	2:38:D:ILE:HD12	16	1.1
(1,852)	2:36:D:THR:HG23	2:38:D:ILE:HD13	16	1.1
(1,840)	1:28:C:PRO:HB3	1:31:C:PHE:HE1	15	1.1
(1,840)	1:28:C:PRO:HB3	1:31:C:PHE:HE2	15	1.1
(1,724)	1:27:C:VAL:HG11	1:28:C:PRO:HG2	16	1.1
(1,724)	1:27:C:VAL:HG11	1:28:C:PRO:HG3	16	1.1
(1,724)	1:27:C:VAL:HG12	1:28:C:PRO:HG2	16	1.1
(1,724)	1:27:C:VAL:HG12	1:28:C:PRO:HG3	16	1.1
(1,724)	1:27:C:VAL:HG13	1:28:C:PRO:HG2	16	1.1
(1,724)	1:27:C:VAL:HG13	1:28:C:PRO:HG3	16	1.1
(1,717)	2:34:D:CYS:HB2	2:35:D:GLY:HA3	2	1.1
(1,685)	1:32:C:VAL:HA	1:33:C:ARG:HD2	19	1.1
(1,685)	1:32:C:VAL:HA	1:33:C:ARG:HD3	19	1.1
(1,669)	2:13:B:CYS:HB3	1:25:C:GLY:HA2	5	1.1
(1,513)	2:15:B:THR:HB	2:17:B:ILE:HD11	9	1.1
(1,513)	2:15:B:THR:HB	2:17:B:ILE:HD12	9	1.1
(1,513)	2:15:B:THR:HB	2:17:B:ILE:HD13	9	1.1
(1,194)	2:15:B:THR:HB	2:17:B:ILE:HD11	9	1.1
(1,194)	2:15:B:THR:HB	2:17:B:ILE:HD12	9	1.1
(1,194)	2:15:B:THR:HB	2:17:B:ILE:HD13	9	1.1
(1,4)	2:13:B:CYS:HB3	1:25:C:GLY:HA2	5	1.1
(1,1139)	1:22:C:GLY:H	2:41:D:TYR:HD1	8	1.09
(1,1139)	1:22:C:GLY:H	2:41:D:TYR:HD2	8	1.09
(1,1100)	2:40:D:PHE:HA	2:41:D:TYR:HD1	1	1.09
(1,1100)	2:40:D:PHE:HA	2:41:D:TYR:HD2	1	1.09
(1,1092)	1:29:C:GLU:HG2	2:41:D:TYR:HD1	12	1.09
(1,1092)	1:29:C:GLU:HG2	2:41:D:TYR:HD2	12	1.09
(1,1067)	1:26:C:HIS:HB3	1:27:C:VAL:HG21	15	1.09
(1,1067)	1:26:C:HIS:HB3	1:27:C:VAL:HG22	15	1.09

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,1067)	1:26:C:HIS:HB3	1:27:C:VAL:HG23	15	1.09
(1,820)	1:22:C:GLY:H	2:41:D:TYR:HD1	8	1.09
(1,820)	1:22:C:GLY:H	2:41:D:TYR:HD2	8	1.09
(1,781)	2:40:D:PHE:HA	2:41:D:TYR:HD1	1	1.09
(1,781)	2:40:D:PHE:HA	2:41:D:TYR:HD2	1	1.09
(1,773)	1:29:C:GLU:HG2	2:41:D:TYR:HD1	12	1.09
(1,773)	1:29:C:GLU:HG2	2:41:D:TYR:HD2	12	1.09
(1,748)	1:26:C:HIS:HB3	1:27:C:VAL:HG21	15	1.09
(1,748)	1:26:C:HIS:HB3	1:27:C:VAL:HG22	15	1.09
(1,748)	1:26:C:HIS:HB3	1:27:C:VAL:HG23	15	1.09
(1,671)	2:17:B:ILE:HG13	2:40:D:PHE:HE1	6	1.09
(1,671)	2:17:B:ILE:HG13	2:40:D:PHE:HE2	6	1.09
(1,481)	1:1:A:GLY:H	2:19:B:PHE:H	12	1.09
(1,441)	1:8:A:GLU:HG3	2:20:B:TYR:HB2	18	1.09
(1,441)	1:8:A:GLU:HG3	2:20:B:TYR:HB3	18	1.09
(1,417)	2:17:B:ILE:HD11	2:39:D:SER:HB3	8	1.09
(1,417)	2:17:B:ILE:HD12	2:39:D:SER:HB3	8	1.09
(1,417)	2:17:B:ILE:HD13	2:39:D:SER:HB3	8	1.09
(1,402)	1:10:A:PHE:HD1	2:14:B:GLY:HA3	10	1.09
(1,402)	1:10:A:PHE:HD2	2:14:B:GLY:HA3	10	1.09
(1,162)	1:1:A:GLY:H	2:19:B:PHE:H	12	1.09
(1,122)	1:8:A:GLU:HG3	2:20:B:TYR:HB2	18	1.09
(1,122)	1:8:A:GLU:HG3	2:20:B:TYR:HB3	18	1.09
(1,98)	2:17:B:ILE:HD11	2:39:D:SER:HB3	8	1.09
(1,98)	2:17:B:ILE:HD12	2:39:D:SER:HB3	8	1.09
(1,98)	2:17:B:ILE:HD13	2:39:D:SER:HB3	8	1.09
(1,83)	1:10:A:PHE:HD1	2:14:B:GLY:HA3	10	1.09
(1,83)	1:10:A:PHE:HD2	2:14:B:GLY:HA3	10	1.09
(1,6)	2:17:B:ILE:HG13	2:40:D:PHE:HE1	6	1.09
(1,6)	2:17:B:ILE:HG13	2:40:D:PHE:HE2	6	1.09
(1,1157)	1:25:C:GLY:H	1:27:C:VAL:H	5	1.08
(1,1055)	1:28:C:PRO:HG2	1:31:C:PHE:HE1	3	1.08
(1,1055)	1:28:C:PRO:HG2	1:31:C:PHE:HE2	3	1.08
(1,1055)	1:28:C:PRO:HG3	1:31:C:PHE:HE1	3	1.08
(1,1055)	1:28:C:PRO:HG3	1:31:C:PHE:HE2	3	1.08
(1,838)	1:25:C:GLY:H	1:27:C:VAL:H	5	1.08
(1,736)	1:28:C:PRO:HG2	1:31:C:PHE:HE1	3	1.08
(1,736)	1:28:C:PRO:HG2	1:31:C:PHE:HE2	3	1.08
(1,736)	1:28:C:PRO:HG3	1:31:C:PHE:HE1	3	1.08
(1,736)	1:28:C:PRO:HG3	1:31:C:PHE:HE2	3	1.08
(1,593)	1:12:A:ARG:HB2	1:12:A:ARG:HE	7	1.08
(1,436)	1:8:A:GLU:HG2	2:20:B:TYR:HD1	12	1.08

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,436)	1:8:A:GLU:HG2	2:20:B:TYR:HD2	12	1.08
(1,380)	2:13:B:CYS:HB2	2:14:B:GLY:HA3	2	1.08
(1,356)	1:8:A:GLU:HG2	2:20:B:TYR:HE1	10	1.08
(1,356)	1:8:A:GLU:HG2	2:20:B:TYR:HE2	10	1.08
(1,274)	1:12:A:ARG:HB2	1:12:A:ARG:HE	7	1.08
(1,117)	1:8:A:GLU:HG2	2:20:B:TYR:HD1	12	1.08
(1,117)	1:8:A:GLU:HG2	2:20:B:TYR:HD2	12	1.08
(1,61)	2:13:B:CYS:HB2	2:14:B:GLY:HA3	2	1.08
(1,37)	1:8:A:GLU:HG2	2:20:B:TYR:HE1	10	1.08
(1,37)	1:8:A:GLU:HG2	2:20:B:TYR:HE2	10	1.08
(1,1309)	1:24:C:ALA:HA	2:40:D:PHE:HD1	7	1.07
(1,1309)	1:24:C:ALA:HA	2:40:D:PHE:HD2	7	1.07
(1,1071)	1:27:C:VAL:HG11	2:41:D:TYR:HE1	3	1.07
(1,1071)	1:27:C:VAL:HG11	2:41:D:TYR:HE2	3	1.07
(1,1071)	1:27:C:VAL:HG12	2:41:D:TYR:HE1	3	1.07
(1,1071)	1:27:C:VAL:HG12	2:41:D:TYR:HE2	3	1.07
(1,1071)	1:27:C:VAL:HG13	2:41:D:TYR:HE1	3	1.07
(1,1071)	1:27:C:VAL:HG13	2:41:D:TYR:HE2	3	1.07
(1,1047)	1:24:C:ALA:HA	2:40:D:PHE:HD1	7	1.07
(1,1047)	1:24:C:ALA:HA	2:40:D:PHE:HD2	7	1.07
(1,1046)	1:27:C:VAL:HB	2:41:D:TYR:HE1	12	1.07
(1,1046)	1:27:C:VAL:HB	2:41:D:TYR:HE2	12	1.07
(1,1008)	2:18:B:SER:HA	2:38:D:ILE:HG13	9	1.07
(1,990)	1:24:C:ALA:HA	2:40:D:PHE:HD1	7	1.07
(1,990)	1:24:C:ALA:HA	2:40:D:PHE:HD2	7	1.07
(1,752)	1:27:C:VAL:HG11	2:41:D:TYR:HE1	3	1.07
(1,752)	1:27:C:VAL:HG11	2:41:D:TYR:HE2	3	1.07
(1,752)	1:27:C:VAL:HG12	2:41:D:TYR:HE1	3	1.07
(1,752)	1:27:C:VAL:HG12	2:41:D:TYR:HE2	3	1.07
(1,752)	1:27:C:VAL:HG13	2:41:D:TYR:HE1	3	1.07
(1,752)	1:27:C:VAL:HG13	2:41:D:TYR:HE2	3	1.07
(1,728)	1:24:C:ALA:HA	2:40:D:PHE:HD1	7	1.07
(1,728)	1:24:C:ALA:HA	2:40:D:PHE:HD2	7	1.07
(1,727)	1:27:C:VAL:HB	2:41:D:TYR:HE1	12	1.07
(1,727)	1:27:C:VAL:HB	2:41:D:TYR:HE2	12	1.07
(1,689)	2:18:B:SER:HA	2:38:D:ILE:HG13	9	1.07
(1,665)	2:19:B:PHE:HE1	2:38:D:ILE:HD11	20	1.07
(1,665)	2:19:B:PHE:HE1	2:38:D:ILE:HD12	20	1.07
(1,665)	2:19:B:PHE:HE1	2:38:D:ILE:HD13	20	1.07
(1,665)	2:19:B:PHE:HE2	2:38:D:ILE:HD11	20	1.07
(1,665)	2:19:B:PHE:HE2	2:38:D:ILE:HD12	20	1.07
(1,665)	2:19:B:PHE:HE2	2:38:D:ILE:HD13	20	1.07

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,662)	2:19:B:PHE:HE1	2:38:D:ILE:HG13	9	1.07
(1,662)	2:19:B:PHE:HE2	2:38:D:ILE:HG13	9	1.07
(1,520)	2:17:B:ILE:HG13	2:40:D:PHE:HB3	12	1.07
(1,404)	1:10:A:PHE:HD1	2:16:B:PRO:HA	5	1.07
(1,404)	1:10:A:PHE:HD2	2:16:B:PRO:HA	5	1.07
(1,201)	2:17:B:ILE:HG13	2:40:D:PHE:HB3	12	1.07
(1,85)	1:10:A:PHE:HD1	2:16:B:PRO:HA	5	1.07
(1,85)	1:10:A:PHE:HD2	2:16:B:PRO:HA	5	1.07
(1,18)	2:19:B:PHE:HE1	2:38:D:ILE:HD11	20	1.07
(1,18)	2:19:B:PHE:HE1	2:38:D:ILE:HD12	20	1.07
(1,18)	2:19:B:PHE:HE1	2:38:D:ILE:HD13	20	1.07
(1,18)	2:19:B:PHE:HE2	2:38:D:ILE:HD11	20	1.07
(1,18)	2:19:B:PHE:HE2	2:38:D:ILE:HD12	20	1.07
(1,18)	2:19:B:PHE:HE2	2:38:D:ILE:HD13	20	1.07
(1,15)	2:19:B:PHE:HE1	2:38:D:ILE:HG13	9	1.07
(1,15)	2:19:B:PHE:HE2	2:38:D:ILE:HG13	9	1.07
(1,1092)	1:29:C:GLU:HG2	2:41:D:TYR:HD1	17	1.06
(1,1092)	1:29:C:GLU:HG2	2:41:D:TYR:HD2	17	1.06
(1,1038)	1:31:C:PHE:HD1	2:37:D:PRO:HD2	20	1.06
(1,1038)	1:31:C:PHE:HD2	2:37:D:PRO:HD2	20	1.06
(1,773)	1:29:C:GLU:HG2	2:41:D:TYR:HD1	17	1.06
(1,773)	1:29:C:GLU:HG2	2:41:D:TYR:HD2	17	1.06
(1,719)	1:31:C:PHE:HD1	2:37:D:PRO:HD2	20	1.06
(1,719)	1:31:C:PHE:HD2	2:37:D:PRO:HD2	20	1.06
(1,420)	1:4:A:GLY:HA3	1:5:A:HIS:HD2	4	1.06
(1,407)	1:1:A:GLY:H	2:18:B:SER:HA	1	1.06
(1,398)	1:7:A:PRO:HA	2:19:B:PHE:HD1	5	1.06
(1,398)	1:7:A:PRO:HA	2:19:B:PHE:HD2	5	1.06
(1,390)	1:6:A:VAL:HB	2:20:B:TYR:HE1	16	1.06
(1,390)	1:6:A:VAL:HB	2:20:B:TYR:HE2	16	1.06
(1,101)	1:4:A:GLY:HA3	1:5:A:HIS:HD2	4	1.06
(1,88)	1:1:A:GLY:H	2:18:B:SER:HA	1	1.06
(1,79)	1:7:A:PRO:HA	2:19:B:PHE:HD1	5	1.06
(1,79)	1:7:A:PRO:HA	2:19:B:PHE:HD2	5	1.06
(1,71)	1:6:A:VAL:HB	2:20:B:TYR:HE1	16	1.06
(1,71)	1:6:A:VAL:HB	2:20:B:TYR:HE2	16	1.06
(1,1092)	1:29:C:GLU:HG2	2:41:D:TYR:HD1	14	1.05
(1,1092)	1:29:C:GLU:HG2	2:41:D:TYR:HD2	14	1.05
(1,1058)	1:31:C:PHE:HD1	2:35:D:GLY:HA3	15	1.05
(1,1058)	1:31:C:PHE:HD2	2:35:D:GLY:HA3	15	1.05
(1,1036)	2:34:D:CYS:HB2	2:35:D:GLY:HA3	11	1.05
(1,773)	1:29:C:GLU:HG2	2:41:D:TYR:HD1	14	1.05

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,773)	1:29:C:GLU:HG2	2:41:D:TYR:HD2	14	1.05
(1,739)	1:31:C:PHE:HD1	2:35:D:GLY:HA3	15	1.05
(1,739)	1:31:C:PHE:HD2	2:35:D:GLY:HA3	15	1.05
(1,717)	2:34:D:CYS:HB2	2:35:D:GLY:HA3	11	1.05
(1,663)	2:19:B:PHE:HD1	2:38:D:ILE:HG13	10	1.05
(1,663)	2:19:B:PHE:HD2	2:38:D:ILE:HG13	10	1.05
(1,452)	2:15:B:THR:HG21	2:16:B:PRO:HD3	7	1.05
(1,452)	2:15:B:THR:HG22	2:16:B:PRO:HD3	7	1.05
(1,452)	2:15:B:THR:HG23	2:16:B:PRO:HD3	7	1.05
(1,451)	2:15:B:THR:HG21	2:16:B:PRO:HB3	15	1.05
(1,451)	2:15:B:THR:HG22	2:16:B:PRO:HB3	15	1.05
(1,451)	2:15:B:THR:HG23	2:16:B:PRO:HB3	15	1.05
(1,399)	1:7:A:PRO:HG2	1:10:A:PHE:HE1	14	1.05
(1,399)	1:7:A:PRO:HG2	1:10:A:PHE:HE2	14	1.05
(1,399)	1:7:A:PRO:HG3	1:10:A:PHE:HE1	14	1.05
(1,399)	1:7:A:PRO:HG3	1:10:A:PHE:HE2	14	1.05
(1,133)	2:15:B:THR:HG21	2:16:B:PRO:HD3	7	1.05
(1,133)	2:15:B:THR:HG22	2:16:B:PRO:HD3	7	1.05
(1,133)	2:15:B:THR:HG23	2:16:B:PRO:HD3	7	1.05
(1,132)	2:15:B:THR:HG21	2:16:B:PRO:HB3	15	1.05
(1,132)	2:15:B:THR:HG22	2:16:B:PRO:HB3	15	1.05
(1,132)	2:15:B:THR:HG23	2:16:B:PRO:HB3	15	1.05
(1,80)	1:7:A:PRO:HG2	1:10:A:PHE:HE1	14	1.05
(1,80)	1:7:A:PRO:HG2	1:10:A:PHE:HE2	14	1.05
(1,80)	1:7:A:PRO:HG3	1:10:A:PHE:HE1	14	1.05
(1,80)	1:7:A:PRO:HG3	1:10:A:PHE:HE2	14	1.05
(1,16)	2:19:B:PHE:HD1	2:38:D:ILE:HG13	10	1.05
(1,16)	2:19:B:PHE:HD2	2:38:D:ILE:HG13	10	1.05
(1,1145)	1:25:C:GLY:H	2:40:D:PHE:HE1	7	1.04
(1,1145)	1:25:C:GLY:H	2:40:D:PHE:HE2	7	1.04
(1,1049)	1:28:C:PRO:HG2	1:31:C:PHE:HD1	2	1.04
(1,1049)	1:28:C:PRO:HG2	1:31:C:PHE:HD2	2	1.04
(1,1049)	1:28:C:PRO:HG3	1:31:C:PHE:HD1	2	1.04
(1,1049)	1:28:C:PRO:HG3	1:31:C:PHE:HD2	2	1.04
(1,826)	1:25:C:GLY:H	2:40:D:PHE:HE1	7	1.04
(1,826)	1:25:C:GLY:H	2:40:D:PHE:HE2	7	1.04
(1,730)	1:28:C:PRO:HG2	1:31:C:PHE:HD1	2	1.04
(1,730)	1:28:C:PRO:HG2	1:31:C:PHE:HD2	2	1.04
(1,730)	1:28:C:PRO:HG3	1:31:C:PHE:HD1	2	1.04
(1,730)	1:28:C:PRO:HG3	1:31:C:PHE:HD2	2	1.04
(1,380)	2:13:B:CYS:HB2	2:14:B:GLY:HA3	19	1.04
(1,61)	2:13:B:CYS:HB2	2:14:B:GLY:HA3	19	1.04

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,1252)	1:28:C:PRO:HG2	1:31:C:PHE:HE1	3	1.03
(1,1252)	1:28:C:PRO:HG2	1:31:C:PHE:HE2	3	1.03
(1,1252)	1:28:C:PRO:HG3	1:31:C:PHE:HE1	3	1.03
(1,1252)	1:28:C:PRO:HG3	1:31:C:PHE:HE2	3	1.03
(1,1251)	1:33:C:ARG:HA	1:33:C:ARG:HE	18	1.03
(1,1171)	2:36:D:THR:HG21	2:38:D:ILE:HD11	6	1.03
(1,1171)	2:36:D:THR:HG21	2:38:D:ILE:HD12	6	1.03
(1,1171)	2:36:D:THR:HG21	2:38:D:ILE:HD13	6	1.03
(1,1171)	2:36:D:THR:HG22	2:38:D:ILE:HD11	6	1.03
(1,1171)	2:36:D:THR:HG22	2:38:D:ILE:HD12	6	1.03
(1,1171)	2:36:D:THR:HG22	2:38:D:ILE:HD13	6	1.03
(1,1171)	2:36:D:THR:HG23	2:38:D:ILE:HD11	6	1.03
(1,1171)	2:36:D:THR:HG23	2:38:D:ILE:HD12	6	1.03
(1,1171)	2:36:D:THR:HG23	2:38:D:ILE:HD13	6	1.03
(1,1036)	2:34:D:CYS:HB2	2:35:D:GLY:HA3	3	1.03
(1,933)	1:28:C:PRO:HG2	1:31:C:PHE:HE1	3	1.03
(1,933)	1:28:C:PRO:HG2	1:31:C:PHE:HE2	3	1.03
(1,933)	1:28:C:PRO:HG3	1:31:C:PHE:HE1	3	1.03
(1,933)	1:28:C:PRO:HG3	1:31:C:PHE:HE2	3	1.03
(1,932)	1:33:C:ARG:HA	1:33:C:ARG:HE	18	1.03
(1,852)	2:36:D:THR:HG21	2:38:D:ILE:HD11	6	1.03
(1,852)	2:36:D:THR:HG21	2:38:D:ILE:HD12	6	1.03
(1,852)	2:36:D:THR:HG21	2:38:D:ILE:HD13	6	1.03
(1,852)	2:36:D:THR:HG22	2:38:D:ILE:HD11	6	1.03
(1,852)	2:36:D:THR:HG22	2:38:D:ILE:HD12	6	1.03
(1,852)	2:36:D:THR:HG22	2:38:D:ILE:HD13	6	1.03
(1,852)	2:36:D:THR:HG23	2:38:D:ILE:HD11	6	1.03
(1,852)	2:36:D:THR:HG23	2:38:D:ILE:HD12	6	1.03
(1,852)	2:36:D:THR:HG23	2:38:D:ILE:HD13	6	1.03
(1,717)	2:34:D:CYS:HB2	2:35:D:GLY:HA3	3	1.03
(1,407)	1:1:A:GLY:H	2:18:B:SER:HA	17	1.03
(1,88)	1:1:A:GLY:H	2:18:B:SER:HA	17	1.03
(1,1102)	1:29:C:GLU:HB3	2:41:D:TYR:HD1	13	1.02
(1,1102)	1:29:C:GLU:HB3	2:41:D:TYR:HD2	13	1.02
(1,1046)	1:27:C:VAL:HB	2:41:D:TYR:HE1	8	1.02
(1,1046)	1:27:C:VAL:HB	2:41:D:TYR:HE2	8	1.02
(1,1036)	2:34:D:CYS:HB2	2:35:D:GLY:HA3	19	1.02
(1,783)	1:29:C:GLU:HB3	2:41:D:TYR:HD1	13	1.02
(1,783)	1:29:C:GLU:HB3	2:41:D:TYR:HD2	13	1.02
(1,727)	1:27:C:VAL:HB	2:41:D:TYR:HE1	8	1.02
(1,727)	1:27:C:VAL:HB	2:41:D:TYR:HE2	8	1.02
(1,717)	2:34:D:CYS:HB2	2:35:D:GLY:HA3	19	1.02

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,483)	1:1:A:GLY:H	2:20:B:TYR:HD1	3	1.02
(1,483)	1:1:A:GLY:H	2:20:B:TYR:HD2	3	1.02
(1,452)	2:15:B:THR:HG21	2:16:B:PRO:HD3	8	1.02
(1,452)	2:15:B:THR:HG22	2:16:B:PRO:HD3	8	1.02
(1,452)	2:15:B:THR:HG23	2:16:B:PRO:HD3	8	1.02
(1,437)	1:8:A:GLU:HG2	2:20:B:TYR:HB2	15	1.02
(1,437)	1:8:A:GLU:HG2	2:20:B:TYR:HB3	15	1.02
(1,436)	1:8:A:GLU:HG2	2:20:B:TYR:HD1	16	1.02
(1,436)	1:8:A:GLU:HG2	2:20:B:TYR:HD2	16	1.02
(1,402)	1:10:A:PHE:HD1	2:14:B:GLY:HA3	4	1.02
(1,402)	1:10:A:PHE:HD2	2:14:B:GLY:HA3	4	1.02
(1,352)	2:17:B:ILE:HG13	2:39:D:SER:HA	17	1.02
(1,164)	1:1:A:GLY:H	2:20:B:TYR:HD1	3	1.02
(1,164)	1:1:A:GLY:H	2:20:B:TYR:HD2	3	1.02
(1,133)	2:15:B:THR:HG21	2:16:B:PRO:HD3	8	1.02
(1,133)	2:15:B:THR:HG22	2:16:B:PRO:HD3	8	1.02
(1,133)	2:15:B:THR:HG23	2:16:B:PRO:HD3	8	1.02
(1,118)	1:8:A:GLU:HG2	2:20:B:TYR:HB2	15	1.02
(1,118)	1:8:A:GLU:HG2	2:20:B:TYR:HB3	15	1.02
(1,117)	1:8:A:GLU:HG2	2:20:B:TYR:HD1	16	1.02
(1,117)	1:8:A:GLU:HG2	2:20:B:TYR:HD2	16	1.02
(1,83)	1:10:A:PHE:HD1	2:14:B:GLY:HA3	4	1.02
(1,83)	1:10:A:PHE:HD2	2:14:B:GLY:HA3	4	1.02
(1,33)	2:17:B:ILE:HG13	2:39:D:SER:HA	17	1.02
(1,1092)	1:29:C:GLU:HG2	2:41:D:TYR:HD1	10	1.01
(1,1092)	1:29:C:GLU:HG2	2:41:D:TYR:HD2	10	1.01
(1,1055)	1:28:C:PRO:HG2	1:31:C:PHE:HE1	16	1.01
(1,1055)	1:28:C:PRO:HG2	1:31:C:PHE:HE2	16	1.01
(1,1055)	1:28:C:PRO:HG3	1:31:C:PHE:HE1	16	1.01
(1,1055)	1:28:C:PRO:HG3	1:31:C:PHE:HE2	16	1.01
(1,773)	1:29:C:GLU:HG2	2:41:D:TYR:HD1	10	1.01
(1,773)	1:29:C:GLU:HG2	2:41:D:TYR:HD2	10	1.01
(1,736)	1:28:C:PRO:HG2	1:31:C:PHE:HE1	16	1.01
(1,736)	1:28:C:PRO:HG2	1:31:C:PHE:HE2	16	1.01
(1,736)	1:28:C:PRO:HG3	1:31:C:PHE:HE1	16	1.01
(1,736)	1:28:C:PRO:HG3	1:31:C:PHE:HE2	16	1.01
(1,595)	1:12:A:ARG:HA	1:12:A:ARG:HE	19	1.01
(1,489)	1:4:A:GLY:H	2:19:B:PHE:HE1	4	1.01
(1,489)	1:4:A:GLY:H	2:19:B:PHE:HE2	4	1.01
(1,446)	1:8:A:GLU:HB3	2:20:B:TYR:HD1	3	1.01
(1,446)	1:8:A:GLU:HB3	2:20:B:TYR:HD2	3	1.01
(1,436)	1:8:A:GLU:HG2	2:20:B:TYR:HD1	7	1.01

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,436)	1:8:A:GLU:HG2	2:20:B:TYR:HD2	7	1.01
(1,380)	2:13:B:CYS:HB2	2:14:B:GLY:HA3	13	1.01
(1,276)	1:12:A:ARG:HA	1:12:A:ARG:HE	19	1.01
(1,170)	1:4:A:GLY:H	2:19:B:PHE:HE1	4	1.01
(1,170)	1:4:A:GLY:H	2:19:B:PHE:HE2	4	1.01
(1,127)	1:8:A:GLU:HB3	2:20:B:TYR:HD1	3	1.01
(1,127)	1:8:A:GLU:HB3	2:20:B:TYR:HD2	3	1.01
(1,117)	1:8:A:GLU:HG2	2:20:B:TYR:HD1	7	1.01
(1,117)	1:8:A:GLU:HG2	2:20:B:TYR:HD2	7	1.01
(1,61)	2:13:B:CYS:HB2	2:14:B:GLY:HA3	13	1.01
(1,1157)	1:25:C:GLY:H	1:27:C:VAL:H	13	1.0
(1,1092)	1:29:C:GLU:HG2	2:41:D:TYR:HD1	2	1.0
(1,1092)	1:29:C:GLU:HG2	2:41:D:TYR:HD2	2	1.0
(1,1045)	1:27:C:VAL:HB	2:41:D:TYR:HD1	7	1.0
(1,1045)	1:27:C:VAL:HB	2:41:D:TYR:HD2	7	1.0
(1,1025)	1:27:C:VAL:HG21	2:41:D:TYR:HE1	18	1.0
(1,1025)	1:27:C:VAL:HG21	2:41:D:TYR:HE2	18	1.0
(1,1025)	1:27:C:VAL:HG22	2:41:D:TYR:HE1	18	1.0
(1,1025)	1:27:C:VAL:HG22	2:41:D:TYR:HE2	18	1.0
(1,1025)	1:27:C:VAL:HG23	2:41:D:TYR:HE1	18	1.0
(1,1025)	1:27:C:VAL:HG23	2:41:D:TYR:HE2	18	1.0
(1,1004)	1:32:C:VAL:HA	1:33:C:ARG:HD2	3	1.0
(1,1004)	1:32:C:VAL:HA	1:33:C:ARG:HD3	3	1.0
(1,838)	1:25:C:GLY:H	1:27:C:VAL:H	13	1.0
(1,773)	1:29:C:GLU:HG2	2:41:D:TYR:HD1	2	1.0
(1,773)	1:29:C:GLU:HG2	2:41:D:TYR:HD2	2	1.0
(1,726)	1:27:C:VAL:HB	2:41:D:TYR:HD1	7	1.0
(1,726)	1:27:C:VAL:HB	2:41:D:TYR:HD2	7	1.0
(1,706)	1:27:C:VAL:HG21	2:41:D:TYR:HE1	18	1.0
(1,706)	1:27:C:VAL:HG21	2:41:D:TYR:HE2	18	1.0
(1,706)	1:27:C:VAL:HG22	2:41:D:TYR:HE1	18	1.0
(1,706)	1:27:C:VAL:HG22	2:41:D:TYR:HE2	18	1.0
(1,706)	1:27:C:VAL:HG23	2:41:D:TYR:HE1	18	1.0
(1,706)	1:27:C:VAL:HG23	2:41:D:TYR:HE2	18	1.0
(1,685)	1:32:C:VAL:HA	1:33:C:ARG:HD2	3	1.0
(1,685)	1:32:C:VAL:HA	1:33:C:ARG:HD3	3	1.0
(1,673)	2:17:B:ILE:HG13	2:40:D:PHE:HB2	19	1.0
(1,673)	2:17:B:ILE:HG13	2:40:D:PHE:HB2	20	1.0
(1,596)	1:7:A:PRO:HG2	1:10:A:PHE:HE1	14	1.0
(1,596)	1:7:A:PRO:HG2	1:10:A:PHE:HE2	14	1.0
(1,596)	1:7:A:PRO:HG3	1:10:A:PHE:HE1	14	1.0
(1,596)	1:7:A:PRO:HG3	1:10:A:PHE:HE2	14	1.0

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,593)	1:12:A:ARG:HB2	1:12:A:ARG:HE	6	1.0
(1,380)	2:13:B:CYS:HB2	2:14:B:GLY:HA3	4	1.0
(1,277)	1:7:A:PRO:HG2	1:10:A:PHE:HE1	14	1.0
(1,277)	1:7:A:PRO:HG2	1:10:A:PHE:HE2	14	1.0
(1,277)	1:7:A:PRO:HG3	1:10:A:PHE:HE1	14	1.0
(1,277)	1:7:A:PRO:HG3	1:10:A:PHE:HE2	14	1.0
(1,274)	1:12:A:ARG:HB2	1:12:A:ARG:HE	6	1.0
(1,61)	2:13:B:CYS:HB2	2:14:B:GLY:HA3	4	1.0
(1,8)	2:17:B:ILE:HG13	2:40:D:PHE:HB2	19	1.0
(1,8)	2:17:B:ILE:HG13	2:40:D:PHE:HB2	20	1.0
(1,1285)	2:38:D:ILE:H	2:38:D:ILE:HD11	10	0.99
(1,1285)	2:38:D:ILE:H	2:38:D:ILE:HD12	10	0.99
(1,1285)	2:38:D:ILE:H	2:38:D:ILE:HD13	10	0.99
(1,1157)	1:25:C:GLY:H	1:27:C:VAL:H	18	0.99
(1,1043)	1:27:C:VAL:HG11	1:28:C:PRO:HG2	15	0.99
(1,1043)	1:27:C:VAL:HG11	1:28:C:PRO:HG3	15	0.99
(1,1043)	1:27:C:VAL:HG12	1:28:C:PRO:HG2	15	0.99
(1,1043)	1:27:C:VAL:HG12	1:28:C:PRO:HG3	15	0.99
(1,1043)	1:27:C:VAL:HG13	1:28:C:PRO:HG2	15	0.99
(1,1043)	1:27:C:VAL:HG13	1:28:C:PRO:HG3	15	0.99
(1,966)	2:38:D:ILE:H	2:38:D:ILE:HD11	10	0.99
(1,966)	2:38:D:ILE:H	2:38:D:ILE:HD12	10	0.99
(1,966)	2:38:D:ILE:H	2:38:D:ILE:HD13	10	0.99
(1,838)	1:25:C:GLY:H	1:27:C:VAL:H	18	0.99
(1,724)	1:27:C:VAL:HG11	1:28:C:PRO:HG2	15	0.99
(1,724)	1:27:C:VAL:HG11	1:28:C:PRO:HG3	15	0.99
(1,724)	1:27:C:VAL:HG12	1:28:C:PRO:HG2	15	0.99
(1,724)	1:27:C:VAL:HG12	1:28:C:PRO:HG3	15	0.99
(1,724)	1:27:C:VAL:HG13	1:28:C:PRO:HG2	15	0.99
(1,724)	1:27:C:VAL:HG13	1:28:C:PRO:HG3	15	0.99
(1,666)	2:17:B:ILE:HD11	2:40:D:PHE:HD1	2	0.99
(1,666)	2:17:B:ILE:HD11	2:40:D:PHE:HD2	2	0.99
(1,666)	2:17:B:ILE:HD12	2:40:D:PHE:HD1	2	0.99
(1,666)	2:17:B:ILE:HD12	2:40:D:PHE:HD2	2	0.99
(1,666)	2:17:B:ILE:HD13	2:40:D:PHE:HD1	2	0.99
(1,666)	2:17:B:ILE:HD13	2:40:D:PHE:HD2	2	0.99
(1,593)	1:12:A:ARG:HB2	1:12:A:ARG:HE	2	0.99
(1,415)	1:6:A:VAL:HG11	2:20:B:TYR:HE1	14	0.99
(1,415)	1:6:A:VAL:HG11	2:20:B:TYR:HE2	14	0.99
(1,415)	1:6:A:VAL:HG12	2:20:B:TYR:HE1	14	0.99
(1,415)	1:6:A:VAL:HG12	2:20:B:TYR:HE2	14	0.99
(1,415)	1:6:A:VAL:HG13	2:20:B:TYR:HE1	14	0.99

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,415)	1:6:A:VAL:HG13	2:20:B:TYR:HE2	14	0.99
(1,404)	1:10:A:PHE:HD1	2:16:B:PRO:HA	19	0.99
(1,404)	1:10:A:PHE:HD2	2:16:B:PRO:HA	19	0.99
(1,274)	1:12:A:ARG:HB2	1:12:A:ARG:HE	2	0.99
(1,96)	1:6:A:VAL:HG11	2:20:B:TYR:HE1	14	0.99
(1,96)	1:6:A:VAL:HG11	2:20:B:TYR:HE2	14	0.99
(1,96)	1:6:A:VAL:HG12	2:20:B:TYR:HE1	14	0.99
(1,96)	1:6:A:VAL:HG12	2:20:B:TYR:HE2	14	0.99
(1,96)	1:6:A:VAL:HG13	2:20:B:TYR:HE1	14	0.99
(1,96)	1:6:A:VAL:HG13	2:20:B:TYR:HE2	14	0.99
(1,85)	1:10:A:PHE:HD1	2:16:B:PRO:HA	19	0.99
(1,85)	1:10:A:PHE:HD2	2:16:B:PRO:HA	19	0.99
(1,1)	2:17:B:ILE:HD11	2:40:D:PHE:HD1	2	0.99
(1,1)	2:17:B:ILE:HD11	2:40:D:PHE:HD2	2	0.99
(1,1)	2:17:B:ILE:HD12	2:40:D:PHE:HD1	2	0.99
(1,1)	2:17:B:ILE:HD12	2:40:D:PHE:HD2	2	0.99
(1,1)	2:17:B:ILE:HD13	2:40:D:PHE:HD1	2	0.99
(1,1)	2:17:B:ILE:HD13	2:40:D:PHE:HD2	2	0.99
(1,1285)	2:38:D:ILE:H	2:38:D:ILE:HD11	20	0.98
(1,1285)	2:38:D:ILE:H	2:38:D:ILE:HD12	20	0.98
(1,1285)	2:38:D:ILE:H	2:38:D:ILE:HD13	20	0.98
(1,1122)	1:25:C:GLY:HA2	1:26:C:HIS:HD2	11	0.98
(1,1078)	1:25:C:GLY:H	2:41:D:TYR:H	4	0.98
(1,1071)	1:27:C:VAL:HG11	2:41:D:TYR:HE1	10	0.98
(1,1071)	1:27:C:VAL:HG11	2:41:D:TYR:HE2	10	0.98
(1,1071)	1:27:C:VAL:HG12	2:41:D:TYR:HE1	10	0.98
(1,1071)	1:27:C:VAL:HG12	2:41:D:TYR:HE2	10	0.98
(1,1071)	1:27:C:VAL:HG13	2:41:D:TYR:HE1	10	0.98
(1,1071)	1:27:C:VAL:HG13	2:41:D:TYR:HE2	10	0.98
(1,1071)	1:27:C:VAL:HG11	2:41:D:TYR:HE1	20	0.98
(1,1071)	1:27:C:VAL:HG11	2:41:D:TYR:HE2	20	0.98
(1,1071)	1:27:C:VAL:HG12	2:41:D:TYR:HE1	20	0.98
(1,1071)	1:27:C:VAL:HG12	2:41:D:TYR:HE2	20	0.98
(1,1071)	1:27:C:VAL:HG13	2:41:D:TYR:HE1	20	0.98
(1,1071)	1:27:C:VAL:HG13	2:41:D:TYR:HE2	20	0.98
(1,966)	2:38:D:ILE:H	2:38:D:ILE:HD11	20	0.98
(1,966)	2:38:D:ILE:H	2:38:D:ILE:HD12	20	0.98
(1,966)	2:38:D:ILE:H	2:38:D:ILE:HD13	20	0.98
(1,803)	1:25:C:GLY:HA2	1:26:C:HIS:HD2	11	0.98
(1,759)	1:25:C:GLY:H	2:41:D:TYR:H	4	0.98
(1,752)	1:27:C:VAL:HG11	2:41:D:TYR:HE1	10	0.98
(1,752)	1:27:C:VAL:HG11	2:41:D:TYR:HE2	10	0.98

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,752)	1:27:C:VAL:HG12	2:41:D:TYR:HE1	10	0.98
(1,752)	1:27:C:VAL:HG12	2:41:D:TYR:HE2	10	0.98
(1,752)	1:27:C:VAL:HG13	2:41:D:TYR:HE1	10	0.98
(1,752)	1:27:C:VAL:HG13	2:41:D:TYR:HE2	10	0.98
(1,752)	1:27:C:VAL:HG11	2:41:D:TYR:HE1	20	0.98
(1,752)	1:27:C:VAL:HG11	2:41:D:TYR:HE2	20	0.98
(1,752)	1:27:C:VAL:HG12	2:41:D:TYR:HE1	20	0.98
(1,752)	1:27:C:VAL:HG12	2:41:D:TYR:HE2	20	0.98
(1,752)	1:27:C:VAL:HG13	2:41:D:TYR:HE1	20	0.98
(1,752)	1:27:C:VAL:HG13	2:41:D:TYR:HE2	20	0.98
(1,520)	2:17:B:ILE:HG13	2:40:D:PHE:HB3	15	0.98
(1,436)	1:8:A:GLU:HG2	2:20:B:TYR:HD1	15	0.98
(1,436)	1:8:A:GLU:HG2	2:20:B:TYR:HD2	15	0.98
(1,420)	1:4:A:GLY:HA3	1:5:A:HIS:HD2	3	0.98
(1,411)	1:5:A:HIS:HB3	1:6:A:VAL:HG21	1	0.98
(1,411)	1:5:A:HIS:HB3	1:6:A:VAL:HG22	1	0.98
(1,411)	1:5:A:HIS:HB3	1:6:A:VAL:HG23	1	0.98
(1,367)	2:15:B:THR:HG21	2:17:B:ILE:HA	15	0.98
(1,367)	2:15:B:THR:HG22	2:17:B:ILE:HA	15	0.98
(1,367)	2:15:B:THR:HG23	2:17:B:ILE:HA	15	0.98
(1,201)	2:17:B:ILE:HG13	2:40:D:PHE:HB3	15	0.98
(1,117)	1:8:A:GLU:HG2	2:20:B:TYR:HD1	15	0.98
(1,117)	1:8:A:GLU:HG2	2:20:B:TYR:HD2	15	0.98
(1,101)	1:4:A:GLY:HA3	1:5:A:HIS:HD2	3	0.98
(1,92)	1:5:A:HIS:HB3	1:6:A:VAL:HG21	1	0.98
(1,92)	1:5:A:HIS:HB3	1:6:A:VAL:HG22	1	0.98
(1,92)	1:5:A:HIS:HB3	1:6:A:VAL:HG23	1	0.98
(1,48)	2:15:B:THR:HG21	2:17:B:ILE:HA	15	0.98
(1,48)	2:15:B:THR:HG22	2:17:B:ILE:HA	15	0.98
(1,48)	2:15:B:THR:HG23	2:17:B:ILE:HA	15	0.98
(1,513)	2:15:B:THR:HB	2:17:B:ILE:HD11	14	0.97
(1,513)	2:15:B:THR:HB	2:17:B:ILE:HD12	14	0.97
(1,513)	2:15:B:THR:HB	2:17:B:ILE:HD13	14	0.97
(1,491)	1:7:A:PRO:HB2	1:10:A:PHE:HE1	15	0.97
(1,491)	1:7:A:PRO:HB2	1:10:A:PHE:HE2	15	0.97
(1,390)	1:6:A:VAL:HB	2:20:B:TYR:HE1	3	0.97
(1,390)	1:6:A:VAL:HB	2:20:B:TYR:HE2	3	0.97
(1,194)	2:15:B:THR:HB	2:17:B:ILE:HD11	14	0.97
(1,194)	2:15:B:THR:HB	2:17:B:ILE:HD12	14	0.97
(1,194)	2:15:B:THR:HB	2:17:B:ILE:HD13	14	0.97
(1,172)	1:7:A:PRO:HB2	1:10:A:PHE:HE1	15	0.97
(1,172)	1:7:A:PRO:HB2	1:10:A:PHE:HE2	15	0.97

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,71)	1:6:A:VAL:HB	2:20:B:TYR:HE1	3	0.97
(1,71)	1:6:A:VAL:HB	2:20:B:TYR:HE2	3	0.97
(1,1252)	1:28:C:PRO:HG2	1:31:C:PHE:HE1	16	0.96
(1,1252)	1:28:C:PRO:HG2	1:31:C:PHE:HE2	16	0.96
(1,1252)	1:28:C:PRO:HG3	1:31:C:PHE:HE1	16	0.96
(1,1252)	1:28:C:PRO:HG3	1:31:C:PHE:HE2	16	0.96
(1,1169)	2:36:D:THR:HB	2:38:D:ILE:HD11	3	0.96
(1,1169)	2:36:D:THR:HB	2:38:D:ILE:HD12	3	0.96
(1,1169)	2:36:D:THR:HB	2:38:D:ILE:HD13	3	0.96
(1,1076)	1:25:C:GLY:HA3	1:26:C:HIS:HD2	9	0.96
(1,1053)	2:39:D:SER:HB2	2:40:D:PHE:H	9	0.96
(1,933)	1:28:C:PRO:HG2	1:31:C:PHE:HE1	16	0.96
(1,933)	1:28:C:PRO:HG2	1:31:C:PHE:HE2	16	0.96
(1,933)	1:28:C:PRO:HG3	1:31:C:PHE:HE1	16	0.96
(1,933)	1:28:C:PRO:HG3	1:31:C:PHE:HE2	16	0.96
(1,850)	2:36:D:THR:HB	2:38:D:ILE:HD11	3	0.96
(1,850)	2:36:D:THR:HB	2:38:D:ILE:HD12	3	0.96
(1,850)	2:36:D:THR:HB	2:38:D:ILE:HD13	3	0.96
(1,757)	1:25:C:GLY:HA3	1:26:C:HIS:HD2	9	0.96
(1,734)	2:39:D:SER:HB2	2:40:D:PHE:H	9	0.96
(1,665)	2:19:B:PHE:HE1	2:38:D:ILE:HD11	15	0.96
(1,665)	2:19:B:PHE:HE1	2:38:D:ILE:HD12	15	0.96
(1,665)	2:19:B:PHE:HE1	2:38:D:ILE:HD13	15	0.96
(1,665)	2:19:B:PHE:HE2	2:38:D:ILE:HD11	15	0.96
(1,665)	2:19:B:PHE:HE2	2:38:D:ILE:HD12	15	0.96
(1,665)	2:19:B:PHE:HE2	2:38:D:ILE:HD13	15	0.96
(1,513)	2:15:B:THR:HB	2:17:B:ILE:HD11	10	0.96
(1,513)	2:15:B:THR:HB	2:17:B:ILE:HD12	10	0.96
(1,513)	2:15:B:THR:HB	2:17:B:ILE:HD13	10	0.96
(1,411)	1:5:A:HIS:HB3	1:6:A:VAL:HG21	19	0.96
(1,411)	1:5:A:HIS:HB3	1:6:A:VAL:HG22	19	0.96
(1,411)	1:5:A:HIS:HB3	1:6:A:VAL:HG23	19	0.96
(1,399)	1:7:A:PRO:HG2	1:10:A:PHE:HE1	4	0.96
(1,399)	1:7:A:PRO:HG2	1:10:A:PHE:HE2	4	0.96
(1,399)	1:7:A:PRO:HG3	1:10:A:PHE:HE1	4	0.96
(1,399)	1:7:A:PRO:HG3	1:10:A:PHE:HE2	4	0.96
(1,355)	1:1:A:GLY:H	2:20:B:TYR:HA	15	0.96
(1,344)	1:10:A:PHE:HE1	2:16:B:PRO:HD2	8	0.96
(1,344)	1:10:A:PHE:HE2	2:16:B:PRO:HD2	8	0.96
(1,342)	1:1:A:GLY:HA3	2:18:B:SER:HB2	20	0.96
(1,339)	1:1:A:GLY:HA3	2:18:B:SER:HB2	20	0.96
(1,194)	2:15:B:THR:HB	2:17:B:ILE:HD11	10	0.96

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,194)	2:15:B:THR:HB	2:17:B:ILE:HD12	10	0.96
(1,194)	2:15:B:THR:HB	2:17:B:ILE:HD13	10	0.96
(1,92)	1:5:A:HIS:HB3	1:6:A:VAL:HG21	19	0.96
(1,92)	1:5:A:HIS:HB3	1:6:A:VAL:HG22	19	0.96
(1,92)	1:5:A:HIS:HB3	1:6:A:VAL:HG23	19	0.96
(1,80)	1:7:A:PRO:HG2	1:10:A:PHE:HE1	4	0.96
(1,80)	1:7:A:PRO:HG2	1:10:A:PHE:HE2	4	0.96
(1,80)	1:7:A:PRO:HG3	1:10:A:PHE:HE1	4	0.96
(1,80)	1:7:A:PRO:HG3	1:10:A:PHE:HE2	4	0.96
(1,36)	1:1:A:GLY:H	2:20:B:TYR:HA	15	0.96
(1,25)	1:10:A:PHE:HE1	2:16:B:PRO:HD2	8	0.96
(1,25)	1:10:A:PHE:HE2	2:16:B:PRO:HD2	8	0.96
(1,23)	1:1:A:GLY:HA3	2:18:B:SER:HB2	20	0.96
(1,20)	1:1:A:GLY:HA3	2:18:B:SER:HB2	20	0.96
(1,18)	2:19:B:PHE:HE1	2:38:D:ILE:HD11	15	0.96
(1,18)	2:19:B:PHE:HE1	2:38:D:ILE:HD12	15	0.96
(1,18)	2:19:B:PHE:HE1	2:38:D:ILE:HD13	15	0.96
(1,18)	2:19:B:PHE:HE2	2:38:D:ILE:HD11	15	0.96
(1,18)	2:19:B:PHE:HE2	2:38:D:ILE:HD12	15	0.96
(1,18)	2:19:B:PHE:HE2	2:38:D:ILE:HD13	15	0.96
(1,1249)	1:33:C:ARG:HB2	1:33:C:ARG:HE	11	0.95
(1,1175)	1:24:C:ALA:HB1	2:40:D:PHE:HB3	3	0.95
(1,1175)	1:24:C:ALA:HB2	2:40:D:PHE:HB3	3	0.95
(1,1175)	1:24:C:ALA:HB3	2:40:D:PHE:HB3	3	0.95
(1,1071)	1:27:C:VAL:HG11	2:41:D:TYR:HE1	15	0.95
(1,1071)	1:27:C:VAL:HG11	2:41:D:TYR:HE2	15	0.95
(1,1071)	1:27:C:VAL:HG12	2:41:D:TYR:HE1	15	0.95
(1,1071)	1:27:C:VAL:HG12	2:41:D:TYR:HE2	15	0.95
(1,1071)	1:27:C:VAL:HG13	2:41:D:TYR:HE1	15	0.95
(1,1071)	1:27:C:VAL:HG13	2:41:D:TYR:HE2	15	0.95
(1,930)	1:33:C:ARG:HB2	1:33:C:ARG:HE	11	0.95
(1,856)	1:24:C:ALA:HB1	2:40:D:PHE:HB3	3	0.95
(1,856)	1:24:C:ALA:HB2	2:40:D:PHE:HB3	3	0.95
(1,856)	1:24:C:ALA:HB3	2:40:D:PHE:HB3	3	0.95
(1,752)	1:27:C:VAL:HG11	2:41:D:TYR:HE1	15	0.95
(1,752)	1:27:C:VAL:HG11	2:41:D:TYR:HE2	15	0.95
(1,752)	1:27:C:VAL:HG12	2:41:D:TYR:HE1	15	0.95
(1,752)	1:27:C:VAL:HG12	2:41:D:TYR:HE2	15	0.95
(1,752)	1:27:C:VAL:HG13	2:41:D:TYR:HE1	15	0.95
(1,752)	1:27:C:VAL:HG13	2:41:D:TYR:HE2	15	0.95
(1,663)	2:19:B:PHE:HD1	2:38:D:ILE:HG13	13	0.95
(1,663)	2:19:B:PHE:HD2	2:38:D:ILE:HG13	13	0.95

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,622)	2:16:B:PRO:HB3	2:16:B:PRO:HD3	17	0.95
(1,595)	1:12:A:ARG:HA	1:12:A:ARG:HE	3	0.95
(1,483)	1:1:A:GLY:H	2:20:B:TYR:HD1	10	0.95
(1,483)	1:1:A:GLY:H	2:20:B:TYR:HD2	10	0.95
(1,452)	2:15:B:THR:HG21	2:16:B:PRO:HD3	11	0.95
(1,452)	2:15:B:THR:HG22	2:16:B:PRO:HD3	11	0.95
(1,452)	2:15:B:THR:HG23	2:16:B:PRO:HD3	11	0.95
(1,445)	1:4:A:GLY:HA2	1:6:A:VAL:H	14	0.95
(1,442)	1:8:A:GLU:HG3	2:19:B:PHE:HA	5	0.95
(1,397)	2:18:B:SER:HB2	2:19:B:PHE:H	6	0.95
(1,390)	1:6:A:VAL:HB	2:20:B:TYR:HE1	14	0.95
(1,390)	1:6:A:VAL:HB	2:20:B:TYR:HE2	14	0.95
(1,380)	2:13:B:CYS:HB2	2:14:B:GLY:HA3	7	0.95
(1,303)	2:16:B:PRO:HB3	2:16:B:PRO:HD3	17	0.95
(1,276)	1:12:A:ARG:HA	1:12:A:ARG:HE	3	0.95
(1,164)	1:1:A:GLY:H	2:20:B:TYR:HD1	10	0.95
(1,164)	1:1:A:GLY:H	2:20:B:TYR:HD2	10	0.95
(1,133)	2:15:B:THR:HG21	2:16:B:PRO:HD3	11	0.95
(1,133)	2:15:B:THR:HG22	2:16:B:PRO:HD3	11	0.95
(1,133)	2:15:B:THR:HG23	2:16:B:PRO:HD3	11	0.95
(1,126)	1:4:A:GLY:HA2	1:6:A:VAL:H	14	0.95
(1,123)	1:8:A:GLU:HG3	2:19:B:PHE:HA	5	0.95
(1,78)	2:18:B:SER:HB2	2:19:B:PHE:H	6	0.95
(1,71)	1:6:A:VAL:HB	2:20:B:TYR:HE1	14	0.95
(1,71)	1:6:A:VAL:HB	2:20:B:TYR:HE2	14	0.95
(1,61)	2:13:B:CYS:HB2	2:14:B:GLY:HA3	7	0.95
(1,16)	2:19:B:PHE:HD1	2:38:D:ILE:HG13	13	0.95
(1,16)	2:19:B:PHE:HD2	2:38:D:ILE:HG13	13	0.95
(1,1159)	1:28:C:PRO:HB3	1:31:C:PHE:HE1	18	0.94
(1,1159)	1:28:C:PRO:HB3	1:31:C:PHE:HE2	18	0.94
(1,1122)	1:25:C:GLY:HA2	1:26:C:HIS:HD2	9	0.94
(1,1045)	1:27:C:VAL:HB	2:41:D:TYR:HD1	12	0.94
(1,1045)	1:27:C:VAL:HB	2:41:D:TYR:HD2	12	0.94
(1,1043)	1:27:C:VAL:HG11	1:28:C:PRO:HG2	1	0.94
(1,1043)	1:27:C:VAL:HG11	1:28:C:PRO:HG3	1	0.94
(1,1043)	1:27:C:VAL:HG12	1:28:C:PRO:HG2	1	0.94
(1,1043)	1:27:C:VAL:HG12	1:28:C:PRO:HG3	1	0.94
(1,1043)	1:27:C:VAL:HG13	1:28:C:PRO:HG2	1	0.94
(1,1043)	1:27:C:VAL:HG13	1:28:C:PRO:HG3	1	0.94
(1,840)	1:28:C:PRO:HB3	1:31:C:PHE:HE1	18	0.94
(1,840)	1:28:C:PRO:HB3	1:31:C:PHE:HE2	18	0.94
(1,803)	1:25:C:GLY:HA2	1:26:C:HIS:HD2	9	0.94

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,726)	1:27:C:VAL:HB	2:41:D:TYR:HD1	12	0.94
(1,726)	1:27:C:VAL:HB	2:41:D:TYR:HD2	12	0.94
(1,724)	1:27:C:VAL:HG11	1:28:C:PRO:HG2	1	0.94
(1,724)	1:27:C:VAL:HG11	1:28:C:PRO:HG3	1	0.94
(1,724)	1:27:C:VAL:HG12	1:28:C:PRO:HG2	1	0.94
(1,724)	1:27:C:VAL:HG12	1:28:C:PRO:HG3	1	0.94
(1,724)	1:27:C:VAL:HG13	1:28:C:PRO:HG2	1	0.94
(1,724)	1:27:C:VAL:HG13	1:28:C:PRO:HG3	1	0.94
(1,668)	1:3:A:ALA:HB1	1:32:C:VAL:HA	19	0.94
(1,668)	1:3:A:ALA:HB2	1:32:C:VAL:HA	19	0.94
(1,668)	1:3:A:ALA:HB3	1:32:C:VAL:HA	19	0.94
(1,661)	2:19:B:PHE:H	2:38:D:ILE:HD11	8	0.94
(1,661)	2:19:B:PHE:H	2:38:D:ILE:HD12	8	0.94
(1,661)	2:19:B:PHE:H	2:38:D:ILE:HD13	8	0.94
(1,631)	2:17:B:ILE:HA	2:17:B:ILE:HD11	16	0.94
(1,631)	2:17:B:ILE:HA	2:17:B:ILE:HD12	16	0.94
(1,631)	2:17:B:ILE:HA	2:17:B:ILE:HD13	16	0.94
(1,474)	1:6:A:VAL:H	2:19:B:PHE:HB2	17	0.94
(1,445)	1:4:A:GLY:HA2	1:6:A:VAL:H	10	0.94
(1,380)	2:13:B:CYS:HB2	2:14:B:GLY:HA3	6	0.94
(1,367)	2:15:B:THR:HG21	2:17:B:ILE:HA	3	0.94
(1,367)	2:15:B:THR:HG22	2:17:B:ILE:HA	3	0.94
(1,367)	2:15:B:THR:HG23	2:17:B:ILE:HA	3	0.94
(1,312)	2:17:B:ILE:HA	2:17:B:ILE:HD11	16	0.94
(1,312)	2:17:B:ILE:HA	2:17:B:ILE:HD12	16	0.94
(1,312)	2:17:B:ILE:HA	2:17:B:ILE:HD13	16	0.94
(1,155)	1:6:A:VAL:H	2:19:B:PHE:HB2	17	0.94
(1,126)	1:4:A:GLY:HA2	1:6:A:VAL:H	10	0.94
(1,61)	2:13:B:CYS:HB2	2:14:B:GLY:HA3	6	0.94
(1,48)	2:15:B:THR:HG21	2:17:B:ILE:HA	3	0.94
(1,48)	2:15:B:THR:HG22	2:17:B:ILE:HA	3	0.94
(1,48)	2:15:B:THR:HG23	2:17:B:ILE:HA	3	0.94
(1,14)	2:19:B:PHE:H	2:38:D:ILE:HD11	8	0.94
(1,14)	2:19:B:PHE:H	2:38:D:ILE:HD12	8	0.94
(1,14)	2:19:B:PHE:H	2:38:D:ILE:HD13	8	0.94
(1,3)	1:3:A:ALA:HB1	1:32:C:VAL:HA	19	0.94
(1,3)	1:3:A:ALA:HB2	1:32:C:VAL:HA	19	0.94
(1,3)	1:3:A:ALA:HB3	1:32:C:VAL:HA	19	0.94
(1,1166)	1:24:C:ALA:H	2:40:D:PHE:HD1	7	0.93
(1,1166)	1:24:C:ALA:H	2:40:D:PHE:HD2	7	0.93
(1,1051)	2:34:D:CYS:HB2	2:35:D:GLY:H	3	0.93
(1,1051)	2:34:D:CYS:HB2	2:35:D:GLY:H	8	0.93

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,1036)	2:34:D:CYS:HB2	2:35:D:GLY:HA3	7	0.93
(1,1027)	1:24:C:ALA:H	2:40:D:PHE:HD1	7	0.93
(1,1027)	1:24:C:ALA:H	2:40:D:PHE:HD2	7	0.93
(1,847)	1:24:C:ALA:H	2:40:D:PHE:HD1	7	0.93
(1,847)	1:24:C:ALA:H	2:40:D:PHE:HD2	7	0.93
(1,732)	2:34:D:CYS:HB2	2:35:D:GLY:H	3	0.93
(1,732)	2:34:D:CYS:HB2	2:35:D:GLY:H	8	0.93
(1,717)	2:34:D:CYS:HB2	2:35:D:GLY:HA3	7	0.93
(1,708)	1:24:C:ALA:H	2:40:D:PHE:HD1	7	0.93
(1,708)	1:24:C:ALA:H	2:40:D:PHE:HD2	7	0.93
(1,519)	1:3:A:ALA:HB1	2:19:B:PHE:HB3	3	0.93
(1,519)	1:3:A:ALA:HB2	2:19:B:PHE:HB3	3	0.93
(1,519)	1:3:A:ALA:HB3	2:19:B:PHE:HB3	3	0.93
(1,443)	1:8:A:GLU:HG3	2:20:B:TYR:HD1	4	0.93
(1,443)	1:8:A:GLU:HG3	2:20:B:TYR:HD2	4	0.93
(1,380)	2:13:B:CYS:HB2	2:14:B:GLY:HA3	5	0.93
(1,200)	1:3:A:ALA:HB1	2:19:B:PHE:HB3	3	0.93
(1,200)	1:3:A:ALA:HB2	2:19:B:PHE:HB3	3	0.93
(1,200)	1:3:A:ALA:HB3	2:19:B:PHE:HB3	3	0.93
(1,124)	1:8:A:GLU:HG3	2:20:B:TYR:HD1	4	0.93
(1,124)	1:8:A:GLU:HG3	2:20:B:TYR:HD2	4	0.93
(1,61)	2:13:B:CYS:HB2	2:14:B:GLY:HA3	5	0.93
(1,1251)	1:33:C:ARG:HA	1:33:C:ARG:HE	15	0.92
(1,1157)	1:25:C:GLY:H	1:27:C:VAL:H	6	0.92
(1,1102)	1:29:C:GLU:HB3	2:41:D:TYR:HD1	7	0.92
(1,1102)	1:29:C:GLU:HB3	2:41:D:TYR:HD2	7	0.92
(1,1093)	1:29:C:GLU:HG2	2:41:D:TYR:HB2	1	0.92
(1,1093)	1:29:C:GLU:HG2	2:41:D:TYR:HB3	1	0.92
(1,932)	1:33:C:ARG:HA	1:33:C:ARG:HE	15	0.92
(1,838)	1:25:C:GLY:H	1:27:C:VAL:H	6	0.92
(1,783)	1:29:C:GLU:HB3	2:41:D:TYR:HD1	7	0.92
(1,783)	1:29:C:GLU:HB3	2:41:D:TYR:HD2	7	0.92
(1,774)	1:29:C:GLU:HG2	2:41:D:TYR:HB2	1	0.92
(1,774)	1:29:C:GLU:HG2	2:41:D:TYR:HB3	1	0.92
(1,672)	2:17:B:ILE:HG13	2:40:D:PHE:HD1	18	0.92
(1,672)	2:17:B:ILE:HG13	2:40:D:PHE:HD2	18	0.92
(1,670)	2:17:B:ILE:HD11	2:40:D:PHE:H	12	0.92
(1,670)	2:17:B:ILE:HD12	2:40:D:PHE:H	12	0.92
(1,670)	2:17:B:ILE:HD13	2:40:D:PHE:H	12	0.92
(1,466)	1:4:A:GLY:HA2	1:5:A:HIS:HD2	3	0.92
(1,395)	2:13:B:CYS:HB2	2:14:B:GLY:H	10	0.92
(1,147)	1:4:A:GLY:HA2	1:5:A:HIS:HD2	3	0.92

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,76)	2:13:B:CYS:HB2	2:14:B:GLY:H	10	0.92
(1,7)	2:17:B:ILE:HG13	2:40:D:PHE:HD1	18	0.92
(1,7)	2:17:B:ILE:HG13	2:40:D:PHE:HD2	18	0.92
(1,5)	2:17:B:ILE:HD11	2:40:D:PHE:H	12	0.92
(1,5)	2:17:B:ILE:HD12	2:40:D:PHE:H	12	0.92
(1,5)	2:17:B:ILE:HD13	2:40:D:PHE:H	12	0.92
(1,1159)	1:28:C:PRO:HB3	1:31:C:PHE:HE1	8	0.91
(1,1159)	1:28:C:PRO:HB3	1:31:C:PHE:HE2	8	0.91
(1,1053)	2:39:D:SER:HB2	2:40:D:PHE:H	1	0.91
(1,840)	1:28:C:PRO:HB3	1:31:C:PHE:HE1	8	0.91
(1,840)	1:28:C:PRO:HB3	1:31:C:PHE:HE2	8	0.91
(1,734)	2:39:D:SER:HB2	2:40:D:PHE:H	1	0.91
(1,673)	2:17:B:ILE:HG13	2:40:D:PHE:HB2	6	0.91
(1,664)	2:19:B:PHE:HB2	2:38:D:ILE:HG13	6	0.91
(1,596)	1:7:A:PRO:HG2	1:10:A:PHE:HE1	4	0.91
(1,596)	1:7:A:PRO:HG2	1:10:A:PHE:HE2	4	0.91
(1,596)	1:7:A:PRO:HG3	1:10:A:PHE:HE1	4	0.91
(1,596)	1:7:A:PRO:HG3	1:10:A:PHE:HE2	4	0.91
(1,483)	1:1:A:GLY:H	2:20:B:TYR:HD1	8	0.91
(1,483)	1:1:A:GLY:H	2:20:B:TYR:HD2	8	0.91
(1,424)	1:4:A:GLY:H	2:20:B:TYR:H	2	0.91
(1,422)	1:4:A:GLY:H	2:20:B:TYR:H	1	0.91
(1,417)	2:17:B:ILE:HD11	2:39:D:SER:HB3	1	0.91
(1,417)	2:17:B:ILE:HD12	2:39:D:SER:HB3	1	0.91
(1,417)	2:17:B:ILE:HD13	2:39:D:SER:HB3	1	0.91
(1,398)	1:7:A:PRO:HA	2:19:B:PHE:HD1	13	0.91
(1,398)	1:7:A:PRO:HA	2:19:B:PHE:HD2	13	0.91
(1,395)	2:13:B:CYS:HB2	2:14:B:GLY:H	20	0.91
(1,380)	2:13:B:CYS:HB2	2:14:B:GLY:HA3	10	0.91
(1,277)	1:7:A:PRO:HG2	1:10:A:PHE:HE1	4	0.91
(1,277)	1:7:A:PRO:HG2	1:10:A:PHE:HE2	4	0.91
(1,277)	1:7:A:PRO:HG3	1:10:A:PHE:HE1	4	0.91
(1,277)	1:7:A:PRO:HG3	1:10:A:PHE:HE2	4	0.91
(1,164)	1:1:A:GLY:H	2:20:B:TYR:HD1	8	0.91
(1,164)	1:1:A:GLY:H	2:20:B:TYR:HD2	8	0.91
(1,105)	1:4:A:GLY:H	2:20:B:TYR:H	2	0.91
(1,103)	1:4:A:GLY:H	2:20:B:TYR:H	1	0.91
(1,98)	2:17:B:ILE:HD11	2:39:D:SER:HB3	1	0.91
(1,98)	2:17:B:ILE:HD12	2:39:D:SER:HB3	1	0.91
(1,98)	2:17:B:ILE:HD13	2:39:D:SER:HB3	1	0.91
(1,79)	1:7:A:PRO:HA	2:19:B:PHE:HD1	13	0.91
(1,79)	1:7:A:PRO:HA	2:19:B:PHE:HD2	13	0.91

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,76)	2:13:B:CYS:HB2	2:14:B:GLY:H	20	0.91
(1,61)	2:13:B:CYS:HB2	2:14:B:GLY:HA3	10	0.91
(1,17)	2:19:B:PHE:HB2	2:38:D:ILE:HG13	6	0.91
(1,8)	2:17:B:ILE:HG13	2:40:D:PHE:HB2	6	0.91
(1,1175)	1:24:C:ALA:HB1	2:40:D:PHE:HB3	14	0.9
(1,1175)	1:24:C:ALA:HB2	2:40:D:PHE:HB3	14	0.9
(1,1175)	1:24:C:ALA:HB3	2:40:D:PHE:HB3	14	0.9
(1,1139)	1:22:C:GLY:H	2:41:D:TYR:HD1	14	0.9
(1,1139)	1:22:C:GLY:H	2:41:D:TYR:HD2	14	0.9
(1,1117)	1:27:C:VAL:HG11	1:28:C:PRO:HB3	1	0.9
(1,1117)	1:27:C:VAL:HG12	1:28:C:PRO:HB3	1	0.9
(1,1117)	1:27:C:VAL:HG13	1:28:C:PRO:HB3	1	0.9
(1,1051)	2:34:D:CYS:HB2	2:35:D:GLY:H	2	0.9
(1,1051)	2:34:D:CYS:HB2	2:35:D:GLY:H	17	0.9
(1,1051)	2:34:D:CYS:HB2	2:35:D:GLY:H	19	0.9
(1,1036)	2:34:D:CYS:HB2	2:35:D:GLY:HA3	12	0.9
(1,856)	1:24:C:ALA:HB1	2:40:D:PHE:HB3	14	0.9
(1,856)	1:24:C:ALA:HB2	2:40:D:PHE:HB3	14	0.9
(1,856)	1:24:C:ALA:HB3	2:40:D:PHE:HB3	14	0.9
(1,820)	1:22:C:GLY:H	2:41:D:TYR:HD1	14	0.9
(1,820)	1:22:C:GLY:H	2:41:D:TYR:HD2	14	0.9
(1,798)	1:27:C:VAL:HG11	1:28:C:PRO:HB3	1	0.9
(1,798)	1:27:C:VAL:HG12	1:28:C:PRO:HB3	1	0.9
(1,798)	1:27:C:VAL:HG13	1:28:C:PRO:HB3	1	0.9
(1,732)	2:34:D:CYS:HB2	2:35:D:GLY:H	2	0.9
(1,732)	2:34:D:CYS:HB2	2:35:D:GLY:H	17	0.9
(1,732)	2:34:D:CYS:HB2	2:35:D:GLY:H	19	0.9
(1,717)	2:34:D:CYS:HB2	2:35:D:GLY:HA3	12	0.9
(1,666)	2:17:B:ILE:HD11	2:40:D:PHE:HD1	10	0.9
(1,666)	2:17:B:ILE:HD11	2:40:D:PHE:HD2	10	0.9
(1,666)	2:17:B:ILE:HD12	2:40:D:PHE:HD1	10	0.9
(1,666)	2:17:B:ILE:HD12	2:40:D:PHE:HD2	10	0.9
(1,666)	2:17:B:ILE:HD13	2:40:D:PHE:HD1	10	0.9
(1,666)	2:17:B:ILE:HD13	2:40:D:PHE:HD2	10	0.9
(1,483)	1:1:A:GLY:H	2:20:B:TYR:HD1	5	0.9
(1,483)	1:1:A:GLY:H	2:20:B:TYR:HD2	5	0.9
(1,351)	2:16:B:PRO:HA	2:17:B:ILE:HD11	17	0.9
(1,351)	2:16:B:PRO:HA	2:17:B:ILE:HD12	17	0.9
(1,351)	2:16:B:PRO:HA	2:17:B:ILE:HD13	17	0.9
(1,348)	1:11:A:VAL:HA	1:12:A:ARG:HD2	6	0.9
(1,348)	1:11:A:VAL:HA	1:12:A:ARG:HD3	6	0.9
(1,348)	1:11:A:VAL:HA	1:12:A:ARG:HD2	15	0.9

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,348)	1:11:A:VAL:HA	1:12:A:ARG:HD3	15	0.9
(1,344)	1:10:A:PHE:HE1	2:16:B:PRO:HD2	18	0.9
(1,344)	1:10:A:PHE:HE2	2:16:B:PRO:HD2	18	0.9
(1,164)	1:1:A:GLY:H	2:20:B:TYR:HD1	5	0.9
(1,164)	1:1:A:GLY:H	2:20:B:TYR:HD2	5	0.9
(1,32)	2:16:B:PRO:HA	2:17:B:ILE:HD11	17	0.9
(1,32)	2:16:B:PRO:HA	2:17:B:ILE:HD12	17	0.9
(1,32)	2:16:B:PRO:HA	2:17:B:ILE:HD13	17	0.9
(1,29)	1:11:A:VAL:HA	1:12:A:ARG:HD2	6	0.9
(1,29)	1:11:A:VAL:HA	1:12:A:ARG:HD3	6	0.9
(1,29)	1:11:A:VAL:HA	1:12:A:ARG:HD2	15	0.9
(1,29)	1:11:A:VAL:HA	1:12:A:ARG:HD3	15	0.9
(1,25)	1:10:A:PHE:HE1	2:16:B:PRO:HD2	18	0.9
(1,25)	1:10:A:PHE:HE2	2:16:B:PRO:HD2	18	0.9
(1,1)	2:17:B:ILE:HD11	2:40:D:PHE:HD1	10	0.9
(1,1)	2:17:B:ILE:HD11	2:40:D:PHE:HD2	10	0.9
(1,1)	2:17:B:ILE:HD12	2:40:D:PHE:HD1	10	0.9
(1,1)	2:17:B:ILE:HD12	2:40:D:PHE:HD2	10	0.9
(1,1)	2:17:B:ILE:HD13	2:40:D:PHE:HD1	10	0.9
(1,1)	2:17:B:ILE:HD13	2:40:D:PHE:HD2	10	0.9
(1,1175)	1:24:C:ALA:HB1	2:40:D:PHE:HB3	11	0.89
(1,1175)	1:24:C:ALA:HB2	2:40:D:PHE:HB3	11	0.89
(1,1175)	1:24:C:ALA:HB3	2:40:D:PHE:HB3	11	0.89
(1,1145)	1:25:C:GLY:H	2:40:D:PHE:HE1	14	0.89
(1,1145)	1:25:C:GLY:H	2:40:D:PHE:HE2	14	0.89
(1,1125)	1:31:C:PHE:HD1	1:32:C:VAL:HG11	11	0.89
(1,1125)	1:31:C:PHE:HD1	1:32:C:VAL:HG12	11	0.89
(1,1125)	1:31:C:PHE:HD1	1:32:C:VAL:HG13	11	0.89
(1,1125)	1:31:C:PHE:HD1	1:32:C:VAL:HG21	11	0.89
(1,1125)	1:31:C:PHE:HD1	1:32:C:VAL:HG22	11	0.89
(1,1125)	1:31:C:PHE:HD1	1:32:C:VAL:HG23	11	0.89
(1,1125)	1:31:C:PHE:HD2	1:32:C:VAL:HG11	11	0.89
(1,1125)	1:31:C:PHE:HD2	1:32:C:VAL:HG12	11	0.89
(1,1125)	1:31:C:PHE:HD2	1:32:C:VAL:HG13	11	0.89
(1,1125)	1:31:C:PHE:HD2	1:32:C:VAL:HG21	11	0.89
(1,1125)	1:31:C:PHE:HD2	1:32:C:VAL:HG22	11	0.89
(1,1125)	1:31:C:PHE:HD2	1:32:C:VAL:HG23	11	0.89
(1,1023)	2:36:D:THR:HG21	2:38:D:ILE:HA	15	0.89
(1,1023)	2:36:D:THR:HG22	2:38:D:ILE:HA	15	0.89
(1,1023)	2:36:D:THR:HG23	2:38:D:ILE:HA	15	0.89
(1,856)	1:24:C:ALA:HB1	2:40:D:PHE:HB3	11	0.89
(1,856)	1:24:C:ALA:HB2	2:40:D:PHE:HB3	11	0.89

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,856)	1:24:C:ALA:HB3	2:40:D:PHE:HB3	11	0.89
(1,826)	1:25:C:GLY:H	2:40:D:PHE:HE1	14	0.89
(1,826)	1:25:C:GLY:H	2:40:D:PHE:HE2	14	0.89
(1,806)	1:31:C:PHE:HD1	1:32:C:VAL:HG11	11	0.89
(1,806)	1:31:C:PHE:HD1	1:32:C:VAL:HG12	11	0.89
(1,806)	1:31:C:PHE:HD1	1:32:C:VAL:HG13	11	0.89
(1,806)	1:31:C:PHE:HD1	1:32:C:VAL:HG21	11	0.89
(1,806)	1:31:C:PHE:HD1	1:32:C:VAL:HG22	11	0.89
(1,806)	1:31:C:PHE:HD1	1:32:C:VAL:HG23	11	0.89
(1,806)	1:31:C:PHE:HD2	1:32:C:VAL:HG11	11	0.89
(1,806)	1:31:C:PHE:HD2	1:32:C:VAL:HG12	11	0.89
(1,806)	1:31:C:PHE:HD2	1:32:C:VAL:HG13	11	0.89
(1,806)	1:31:C:PHE:HD2	1:32:C:VAL:HG21	11	0.89
(1,806)	1:31:C:PHE:HD2	1:32:C:VAL:HG22	11	0.89
(1,806)	1:31:C:PHE:HD2	1:32:C:VAL:HG23	11	0.89
(1,704)	2:36:D:THR:HG21	2:38:D:ILE:HA	15	0.89
(1,704)	2:36:D:THR:HG22	2:38:D:ILE:HA	15	0.89
(1,704)	2:36:D:THR:HG23	2:38:D:ILE:HA	15	0.89
(1,622)	2:16:B:PRO:HB3	2:16:B:PRO:HD3	6	0.89
(1,622)	2:16:B:PRO:HB3	2:16:B:PRO:HD3	12	0.89
(1,483)	1:1:A:GLY:H	2:20:B:TYR:HD1	11	0.89
(1,483)	1:1:A:GLY:H	2:20:B:TYR:HD2	11	0.89
(1,446)	1:8:A:GLU:HB3	2:20:B:TYR:HD1	17	0.89
(1,446)	1:8:A:GLU:HB3	2:20:B:TYR:HD2	17	0.89
(1,390)	1:6:A:VAL:HB	2:20:B:TYR:HE1	5	0.89
(1,390)	1:6:A:VAL:HB	2:20:B:TYR:HE2	5	0.89
(1,303)	2:16:B:PRO:HB3	2:16:B:PRO:HD3	6	0.89
(1,303)	2:16:B:PRO:HB3	2:16:B:PRO:HD3	12	0.89
(1,164)	1:1:A:GLY:H	2:20:B:TYR:HD1	11	0.89
(1,164)	1:1:A:GLY:H	2:20:B:TYR:HD2	11	0.89
(1,127)	1:8:A:GLU:HB3	2:20:B:TYR:HD1	17	0.89
(1,127)	1:8:A:GLU:HB3	2:20:B:TYR:HD2	17	0.89
(1,71)	1:6:A:VAL:HB	2:20:B:TYR:HE1	5	0.89
(1,71)	1:6:A:VAL:HB	2:20:B:TYR:HE2	5	0.89
(1,1278)	2:37:D:PRO:HB3	2:37:D:PRO:HD3	10	0.88
(1,1117)	1:27:C:VAL:HG11	1:28:C:PRO:HB3	16	0.88
(1,1117)	1:27:C:VAL:HG12	1:28:C:PRO:HB3	16	0.88
(1,1117)	1:27:C:VAL:HG13	1:28:C:PRO:HB3	16	0.88
(1,1089)	1:29:C:GLU:HB2	2:41:D:TYR:HB2	17	0.88
(1,1089)	1:29:C:GLU:HB2	2:41:D:TYR:HB3	17	0.88
(1,1061)	1:22:C:GLY:H	2:39:D:SER:HB2	6	0.88
(1,959)	2:37:D:PRO:HB3	2:37:D:PRO:HD3	10	0.88

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,798)	1:27:C:VAL:HG11	1:28:C:PRO:HB3	16	0.88
(1,798)	1:27:C:VAL:HG12	1:28:C:PRO:HB3	16	0.88
(1,798)	1:27:C:VAL:HG13	1:28:C:PRO:HB3	16	0.88
(1,770)	1:29:C:GLU:HB2	2:41:D:TYR:HB2	17	0.88
(1,770)	1:29:C:GLU:HB2	2:41:D:TYR:HB3	17	0.88
(1,742)	1:22:C:GLY:H	2:39:D:SER:HB2	6	0.88
(1,622)	2:16:B:PRO:HB3	2:16:B:PRO:HD3	5	0.88
(1,595)	1:12:A:ARG:HA	1:12:A:ARG:HE	20	0.88
(1,587)	1:12:A:ARG:H	1:12:A:ARG:HE	15	0.88
(1,453)	1:10:A:PHE:HD1	2:15:B:THR:HG21	15	0.88
(1,453)	1:10:A:PHE:HD1	2:15:B:THR:HG22	15	0.88
(1,453)	1:10:A:PHE:HD1	2:15:B:THR:HG23	15	0.88
(1,453)	1:10:A:PHE:HD2	2:15:B:THR:HG21	15	0.88
(1,453)	1:10:A:PHE:HD2	2:15:B:THR:HG22	15	0.88
(1,453)	1:10:A:PHE:HD2	2:15:B:THR:HG23	15	0.88
(1,397)	2:18:B:SER:HB2	2:19:B:PHE:H	19	0.88
(1,361)	1:10:A:PHE:HE1	2:14:B:GLY:HA3	5	0.88
(1,361)	1:10:A:PHE:HE2	2:14:B:GLY:HA3	5	0.88
(1,361)	1:10:A:PHE:HE1	2:14:B:GLY:HA3	11	0.88
(1,361)	1:10:A:PHE:HE2	2:14:B:GLY:HA3	11	0.88
(1,348)	1:11:A:VAL:HA	1:12:A:ARG:HD2	13	0.88
(1,348)	1:11:A:VAL:HA	1:12:A:ARG:HD3	13	0.88
(1,303)	2:16:B:PRO:HB3	2:16:B:PRO:HD3	5	0.88
(1,276)	1:12:A:ARG:HA	1:12:A:ARG:HE	20	0.88
(1,268)	1:12:A:ARG:H	1:12:A:ARG:HE	15	0.88
(1,134)	1:10:A:PHE:HD1	2:15:B:THR:HG21	15	0.88
(1,134)	1:10:A:PHE:HD1	2:15:B:THR:HG22	15	0.88
(1,134)	1:10:A:PHE:HD1	2:15:B:THR:HG23	15	0.88
(1,134)	1:10:A:PHE:HD2	2:15:B:THR:HG21	15	0.88
(1,134)	1:10:A:PHE:HD2	2:15:B:THR:HG22	15	0.88
(1,134)	1:10:A:PHE:HD2	2:15:B:THR:HG23	15	0.88
(1,78)	2:18:B:SER:HB2	2:19:B:PHE:H	19	0.88
(1,42)	1:10:A:PHE:HE1	2:14:B:GLY:HA3	5	0.88
(1,42)	1:10:A:PHE:HE2	2:14:B:GLY:HA3	5	0.88
(1,42)	1:10:A:PHE:HE1	2:14:B:GLY:HA3	11	0.88
(1,42)	1:10:A:PHE:HE2	2:14:B:GLY:HA3	11	0.88
(1,29)	1:11:A:VAL:HA	1:12:A:ARG:HD2	13	0.88
(1,29)	1:11:A:VAL:HA	1:12:A:ARG:HD3	13	0.88
(1,1251)	1:33:C:ARG:HA	1:33:C:ARG:HE	11	0.87
(1,1145)	1:25:C:GLY:H	2:40:D:PHE:HE1	18	0.87
(1,1145)	1:25:C:GLY:H	2:40:D:PHE:HE2	18	0.87
(1,1101)	1:25:C:GLY:HA2	1:27:C:VAL:H	8	0.87

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,1051)	2:34:D:CYS:HB2	2:35:D:GLY:H	4	0.87
(1,1007)	2:37:D:PRO:HA	2:38:D:ILE:HD11	5	0.87
(1,1007)	2:37:D:PRO:HA	2:38:D:ILE:HD12	5	0.87
(1,1007)	2:37:D:PRO:HA	2:38:D:ILE:HD13	5	0.87
(1,932)	1:33:C:ARG:HA	1:33:C:ARG:HE	11	0.87
(1,826)	1:25:C:GLY:H	2:40:D:PHE:HE1	18	0.87
(1,826)	1:25:C:GLY:H	2:40:D:PHE:HE2	18	0.87
(1,782)	1:25:C:GLY:HA2	1:27:C:VAL:H	8	0.87
(1,732)	2:34:D:CYS:HB2	2:35:D:GLY:H	4	0.87
(1,688)	2:37:D:PRO:HA	2:38:D:ILE:HD11	5	0.87
(1,688)	2:37:D:PRO:HA	2:38:D:ILE:HD12	5	0.87
(1,688)	2:37:D:PRO:HA	2:38:D:ILE:HD13	5	0.87
(1,672)	2:17:B:ILE:HG13	2:40:D:PHE:HD1	17	0.87
(1,672)	2:17:B:ILE:HG13	2:40:D:PHE:HD2	17	0.87
(1,668)	1:3:A:ALA:HB1	1:32:C:VAL:HA	14	0.87
(1,668)	1:3:A:ALA:HB2	1:32:C:VAL:HA	14	0.87
(1,668)	1:3:A:ALA:HB3	1:32:C:VAL:HA	14	0.87
(1,663)	2:19:B:PHE:HD1	2:38:D:ILE:HG13	3	0.87
(1,663)	2:19:B:PHE:HD2	2:38:D:ILE:HG13	3	0.87
(1,504)	1:11:A:VAL:HA	1:12:A:ARG:HB3	8	0.87
(1,386)	1:4:A:GLY:H	2:19:B:PHE:HD1	3	0.87
(1,386)	1:4:A:GLY:H	2:19:B:PHE:HD2	3	0.87
(1,369)	1:6:A:VAL:HG21	2:20:B:TYR:HE1	8	0.87
(1,369)	1:6:A:VAL:HG21	2:20:B:TYR:HE2	8	0.87
(1,369)	1:6:A:VAL:HG22	2:20:B:TYR:HE1	8	0.87
(1,369)	1:6:A:VAL:HG22	2:20:B:TYR:HE2	8	0.87
(1,369)	1:6:A:VAL:HG23	2:20:B:TYR:HE1	8	0.87
(1,369)	1:6:A:VAL:HG23	2:20:B:TYR:HE2	8	0.87
(1,185)	1:11:A:VAL:HA	1:12:A:ARG:HB3	8	0.87
(1,67)	1:4:A:GLY:H	2:19:B:PHE:HD1	3	0.87
(1,67)	1:4:A:GLY:H	2:19:B:PHE:HD2	3	0.87
(1,50)	1:6:A:VAL:HG21	2:20:B:TYR:HE1	8	0.87
(1,50)	1:6:A:VAL:HG21	2:20:B:TYR:HE2	8	0.87
(1,50)	1:6:A:VAL:HG22	2:20:B:TYR:HE1	8	0.87
(1,50)	1:6:A:VAL:HG22	2:20:B:TYR:HE2	8	0.87
(1,50)	1:6:A:VAL:HG23	2:20:B:TYR:HE1	8	0.87
(1,50)	1:6:A:VAL:HG23	2:20:B:TYR:HE2	8	0.87
(1,16)	2:19:B:PHE:HD1	2:38:D:ILE:HG13	3	0.87
(1,16)	2:19:B:PHE:HD2	2:38:D:ILE:HG13	3	0.87
(1,7)	2:17:B:ILE:HG13	2:40:D:PHE:HD1	17	0.87
(1,7)	2:17:B:ILE:HG13	2:40:D:PHE:HD2	17	0.87
(1,3)	1:3:A:ALA:HB1	1:32:C:VAL:HA	14	0.87

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,3)	1:3:A:ALA:HB2	1:32:C:VAL:HA	14	0.87
(1,3)	1:3:A:ALA:HB3	1:32:C:VAL:HA	14	0.87
(1,1175)	1:24:C:ALA:HB1	2:40:D:PHE:HB3	4	0.86
(1,1175)	1:24:C:ALA:HB2	2:40:D:PHE:HB3	4	0.86
(1,1175)	1:24:C:ALA:HB3	2:40:D:PHE:HB3	4	0.86
(1,1144)	1:27:C:VAL:H	2:40:D:PHE:HD1	2	0.86
(1,1144)	1:27:C:VAL:H	2:40:D:PHE:HD2	2	0.86
(1,1036)	2:34:D:CYS:HB2	2:35:D:GLY:HA3	6	0.86
(1,856)	1:24:C:ALA:HB1	2:40:D:PHE:HB3	4	0.86
(1,856)	1:24:C:ALA:HB2	2:40:D:PHE:HB3	4	0.86
(1,856)	1:24:C:ALA:HB3	2:40:D:PHE:HB3	4	0.86
(1,825)	1:27:C:VAL:H	2:40:D:PHE:HD1	2	0.86
(1,825)	1:27:C:VAL:H	2:40:D:PHE:HD2	2	0.86
(1,717)	2:34:D:CYS:HB2	2:35:D:GLY:HA3	6	0.86
(1,631)	2:17:B:ILE:HA	2:17:B:ILE:HD11	10	0.86
(1,631)	2:17:B:ILE:HA	2:17:B:ILE:HD12	10	0.86
(1,631)	2:17:B:ILE:HA	2:17:B:ILE:HD13	10	0.86
(1,595)	1:12:A:ARG:HA	1:12:A:ARG:HE	10	0.86
(1,452)	2:15:B:THR:HG21	2:16:B:PRO:HD3	12	0.86
(1,452)	2:15:B:THR:HG22	2:16:B:PRO:HD3	12	0.86
(1,452)	2:15:B:THR:HG23	2:16:B:PRO:HD3	12	0.86
(1,451)	2:15:B:THR:HG21	2:16:B:PRO:HB3	12	0.86
(1,451)	2:15:B:THR:HG22	2:16:B:PRO:HB3	12	0.86
(1,451)	2:15:B:THR:HG23	2:16:B:PRO:HB3	12	0.86
(1,312)	2:17:B:ILE:HA	2:17:B:ILE:HD11	10	0.86
(1,312)	2:17:B:ILE:HA	2:17:B:ILE:HD12	10	0.86
(1,312)	2:17:B:ILE:HA	2:17:B:ILE:HD13	10	0.86
(1,276)	1:12:A:ARG:HA	1:12:A:ARG:HE	10	0.86
(1,133)	2:15:B:THR:HG21	2:16:B:PRO:HD3	12	0.86
(1,133)	2:15:B:THR:HG22	2:16:B:PRO:HD3	12	0.86
(1,133)	2:15:B:THR:HG23	2:16:B:PRO:HD3	12	0.86
(1,132)	2:15:B:THR:HG21	2:16:B:PRO:HB3	12	0.86
(1,132)	2:15:B:THR:HG22	2:16:B:PRO:HB3	12	0.86
(1,132)	2:15:B:THR:HG23	2:16:B:PRO:HB3	12	0.86
(1,1169)	2:36:D:THR:HB	2:38:D:ILE:HD11	8	0.85
(1,1169)	2:36:D:THR:HB	2:38:D:ILE:HD12	8	0.85
(1,1169)	2:36:D:THR:HB	2:38:D:ILE:HD13	8	0.85
(1,1130)	1:27:C:VAL:H	2:40:D:PHE:HB2	1	0.85
(1,1058)	1:31:C:PHE:HD1	2:35:D:GLY:HA3	2	0.85
(1,1058)	1:31:C:PHE:HD2	2:35:D:GLY:HA3	2	0.85
(1,850)	2:36:D:THR:HB	2:38:D:ILE:HD11	8	0.85
(1,850)	2:36:D:THR:HB	2:38:D:ILE:HD12	8	0.85

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,850)	2:36:D:THR:HB	2:38:D:ILE:HD13	8	0.85
(1,811)	1:27:C:VAL:H	2:40:D:PHE:HB2	1	0.85
(1,739)	1:31:C:PHE:HD1	2:35:D:GLY:HA3	2	0.85
(1,739)	1:31:C:PHE:HD2	2:35:D:GLY:HA3	2	0.85
(1,673)	2:17:B:ILE:HG13	2:40:D:PHE:HB2	2	0.85
(1,671)	2:17:B:ILE:HG13	2:40:D:PHE:HE1	5	0.85
(1,671)	2:17:B:ILE:HG13	2:40:D:PHE:HE2	5	0.85
(1,666)	2:17:B:ILE:HD11	2:40:D:PHE:HD1	13	0.85
(1,666)	2:17:B:ILE:HD11	2:40:D:PHE:HD2	13	0.85
(1,666)	2:17:B:ILE:HD12	2:40:D:PHE:HD1	13	0.85
(1,666)	2:17:B:ILE:HD12	2:40:D:PHE:HD2	13	0.85
(1,666)	2:17:B:ILE:HD13	2:40:D:PHE:HD1	13	0.85
(1,666)	2:17:B:ILE:HD13	2:40:D:PHE:HD2	13	0.85
(1,622)	2:16:B:PRO:HB3	2:16:B:PRO:HD3	16	0.85
(1,470)	1:2:A:GLY:H	2:19:B:PHE:H	17	0.85
(1,397)	2:18:B:SER:HB2	2:19:B:PHE:H	10	0.85
(1,395)	2:13:B:CYS:HB2	2:14:B:GLY:H	14	0.85
(1,395)	2:13:B:CYS:HB2	2:14:B:GLY:H	17	0.85
(1,303)	2:16:B:PRO:HB3	2:16:B:PRO:HD3	16	0.85
(1,151)	1:2:A:GLY:H	2:19:B:PHE:H	17	0.85
(1,78)	2:18:B:SER:HB2	2:19:B:PHE:H	10	0.85
(1,76)	2:13:B:CYS:HB2	2:14:B:GLY:H	14	0.85
(1,76)	2:13:B:CYS:HB2	2:14:B:GLY:H	17	0.85
(1,8)	2:17:B:ILE:HG13	2:40:D:PHE:HB2	2	0.85
(1,6)	2:17:B:ILE:HG13	2:40:D:PHE:HE1	5	0.85
(1,6)	2:17:B:ILE:HG13	2:40:D:PHE:HE2	5	0.85
(1,1)	2:17:B:ILE:HD11	2:40:D:PHE:HD1	13	0.85
(1,1)	2:17:B:ILE:HD11	2:40:D:PHE:HD2	13	0.85
(1,1)	2:17:B:ILE:HD12	2:40:D:PHE:HD1	13	0.85
(1,1)	2:17:B:ILE:HD12	2:40:D:PHE:HD2	13	0.85
(1,1)	2:17:B:ILE:HD13	2:40:D:PHE:HD1	13	0.85
(1,1)	2:17:B:ILE:HD13	2:40:D:PHE:HD2	13	0.85
(1,1139)	1:22:C:GLY:H	2:41:D:TYR:HD1	5	0.84
(1,1139)	1:22:C:GLY:H	2:41:D:TYR:HD2	5	0.84
(1,1073)	2:18:B:SER:HB3	2:38:D:ILE:HD11	8	0.84
(1,1073)	2:18:B:SER:HB3	2:38:D:ILE:HD12	8	0.84
(1,1073)	2:18:B:SER:HB3	2:38:D:ILE:HD13	8	0.84
(1,1053)	2:39:D:SER:HB2	2:40:D:PHE:H	2	0.84
(1,1051)	2:34:D:CYS:HB2	2:35:D:GLY:H	14	0.84
(1,1025)	1:27:C:VAL:HG21	2:41:D:TYR:HE1	10	0.84
(1,1025)	1:27:C:VAL:HG21	2:41:D:TYR:HE2	10	0.84
(1,1025)	1:27:C:VAL:HG22	2:41:D:TYR:HE1	10	0.84

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,1025)	1:27:C:VAL:HG22	2:41:D:TYR:HE2	10	0.84
(1,1025)	1:27:C:VAL:HG23	2:41:D:TYR:HE1	10	0.84
(1,1025)	1:27:C:VAL:HG23	2:41:D:TYR:HE2	10	0.84
(1,1025)	1:27:C:VAL:HG21	2:41:D:TYR:HE1	13	0.84
(1,1025)	1:27:C:VAL:HG21	2:41:D:TYR:HE2	13	0.84
(1,1025)	1:27:C:VAL:HG22	2:41:D:TYR:HE1	13	0.84
(1,1025)	1:27:C:VAL:HG22	2:41:D:TYR:HE2	13	0.84
(1,1025)	1:27:C:VAL:HG23	2:41:D:TYR:HE1	13	0.84
(1,1025)	1:27:C:VAL:HG23	2:41:D:TYR:HE2	13	0.84
(1,1017)	1:31:C:PHE:HE1	2:35:D:GLY:HA3	13	0.84
(1,1017)	1:31:C:PHE:HE2	2:35:D:GLY:HA3	13	0.84
(1,1004)	1:32:C:VAL:HA	1:33:C:ARG:HD2	8	0.84
(1,1004)	1:32:C:VAL:HA	1:33:C:ARG:HD3	8	0.84
(1,999)	1:28:C:PRO:HD3	2:40:D:PHE:HD1	2	0.84
(1,999)	1:28:C:PRO:HD3	2:40:D:PHE:HD2	2	0.84
(1,820)	1:22:C:GLY:H	2:41:D:TYR:HD1	5	0.84
(1,820)	1:22:C:GLY:H	2:41:D:TYR:HD2	5	0.84
(1,754)	2:18:B:SER:HB3	2:38:D:ILE:HD11	8	0.84
(1,754)	2:18:B:SER:HB3	2:38:D:ILE:HD12	8	0.84
(1,754)	2:18:B:SER:HB3	2:38:D:ILE:HD13	8	0.84
(1,734)	2:39:D:SER:HB2	2:40:D:PHE:H	2	0.84
(1,732)	2:34:D:CYS:HB2	2:35:D:GLY:H	14	0.84
(1,706)	1:27:C:VAL:HG21	2:41:D:TYR:HE1	10	0.84
(1,706)	1:27:C:VAL:HG21	2:41:D:TYR:HE2	10	0.84
(1,706)	1:27:C:VAL:HG22	2:41:D:TYR:HE1	10	0.84
(1,706)	1:27:C:VAL:HG22	2:41:D:TYR:HE2	10	0.84
(1,706)	1:27:C:VAL:HG23	2:41:D:TYR:HE1	10	0.84
(1,706)	1:27:C:VAL:HG23	2:41:D:TYR:HE2	10	0.84
(1,706)	1:27:C:VAL:HG21	2:41:D:TYR:HE1	13	0.84
(1,706)	1:27:C:VAL:HG21	2:41:D:TYR:HE2	13	0.84
(1,706)	1:27:C:VAL:HG22	2:41:D:TYR:HE1	13	0.84
(1,706)	1:27:C:VAL:HG22	2:41:D:TYR:HE2	13	0.84
(1,706)	1:27:C:VAL:HG23	2:41:D:TYR:HE1	13	0.84
(1,706)	1:27:C:VAL:HG23	2:41:D:TYR:HE2	13	0.84
(1,698)	1:31:C:PHE:HE1	2:35:D:GLY:HA3	13	0.84
(1,698)	1:31:C:PHE:HE2	2:35:D:GLY:HA3	13	0.84
(1,685)	1:32:C:VAL:HA	1:33:C:ARG:HD2	8	0.84
(1,685)	1:32:C:VAL:HA	1:33:C:ARG:HD3	8	0.84
(1,680)	1:28:C:PRO:HD3	2:40:D:PHE:HD1	2	0.84
(1,680)	1:28:C:PRO:HD3	2:40:D:PHE:HD2	2	0.84
(1,503)	1:7:A:PRO:HB3	1:10:A:PHE:HE1	6	0.84
(1,503)	1:7:A:PRO:HB3	1:10:A:PHE:HE2	6	0.84

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,403)	1:7:A:PRO:HG2	2:19:B:PHE:HD1	13	0.84
(1,403)	1:7:A:PRO:HG2	2:19:B:PHE:HD2	13	0.84
(1,403)	1:7:A:PRO:HG3	2:19:B:PHE:HD1	13	0.84
(1,403)	1:7:A:PRO:HG3	2:19:B:PHE:HD2	13	0.84
(1,384)	1:10:A:PHE:HE1	2:16:B:PRO:HD2	8	0.84
(1,384)	1:10:A:PHE:HE2	2:16:B:PRO:HD2	8	0.84
(1,380)	2:13:B:CYS:HB2	2:14:B:GLY:HA3	14	0.84
(1,184)	1:7:A:PRO:HB3	1:10:A:PHE:HE1	6	0.84
(1,184)	1:7:A:PRO:HB3	1:10:A:PHE:HE2	6	0.84
(1,84)	1:7:A:PRO:HG2	2:19:B:PHE:HD1	13	0.84
(1,84)	1:7:A:PRO:HG2	2:19:B:PHE:HD2	13	0.84
(1,84)	1:7:A:PRO:HG3	2:19:B:PHE:HD1	13	0.84
(1,84)	1:7:A:PRO:HG3	2:19:B:PHE:HD2	13	0.84
(1,65)	1:10:A:PHE:HE1	2:16:B:PRO:HD2	8	0.84
(1,65)	1:10:A:PHE:HE2	2:16:B:PRO:HD2	8	0.84
(1,61)	2:13:B:CYS:HB2	2:14:B:GLY:HA3	14	0.84
(1,1278)	2:37:D:PRO:HB3	2:37:D:PRO:HD3	18	0.83
(1,1120)	1:27:C:VAL:HG11	2:41:D:TYR:HB2	3	0.83
(1,1120)	1:27:C:VAL:HG11	2:41:D:TYR:HB3	3	0.83
(1,1120)	1:27:C:VAL:HG12	2:41:D:TYR:HB2	3	0.83
(1,1120)	1:27:C:VAL:HG12	2:41:D:TYR:HB3	3	0.83
(1,1120)	1:27:C:VAL:HG13	2:41:D:TYR:HB2	3	0.83
(1,1120)	1:27:C:VAL:HG13	2:41:D:TYR:HB3	3	0.83
(1,959)	2:37:D:PRO:HB3	2:37:D:PRO:HD3	18	0.83
(1,801)	1:27:C:VAL:HG11	2:41:D:TYR:HB2	3	0.83
(1,801)	1:27:C:VAL:HG11	2:41:D:TYR:HB3	3	0.83
(1,801)	1:27:C:VAL:HG12	2:41:D:TYR:HB2	3	0.83
(1,801)	1:27:C:VAL:HG12	2:41:D:TYR:HB3	3	0.83
(1,801)	1:27:C:VAL:HG13	2:41:D:TYR:HB2	3	0.83
(1,801)	1:27:C:VAL:HG13	2:41:D:TYR:HB3	3	0.83
(1,629)	2:17:B:ILE:H	2:17:B:ILE:HD11	10	0.83
(1,629)	2:17:B:ILE:H	2:17:B:ILE:HD12	10	0.83
(1,629)	2:17:B:ILE:H	2:17:B:ILE:HD13	10	0.83
(1,492)	1:7:A:PRO:HB2	1:10:A:PHE:HD1	17	0.83
(1,492)	1:7:A:PRO:HB2	1:10:A:PHE:HD2	17	0.83
(1,489)	1:4:A:GLY:H	2:19:B:PHE:HE1	19	0.83
(1,489)	1:4:A:GLY:H	2:19:B:PHE:HE2	19	0.83
(1,469)	1:10:A:PHE:HD1	1:11:A:VAL:HG11	18	0.83
(1,469)	1:10:A:PHE:HD1	1:11:A:VAL:HG12	18	0.83
(1,469)	1:10:A:PHE:HD1	1:11:A:VAL:HG13	18	0.83
(1,469)	1:10:A:PHE:HD1	1:11:A:VAL:HG21	18	0.83
(1,469)	1:10:A:PHE:HD1	1:11:A:VAL:HG22	18	0.83

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,469)	1:10:A:PHE:HD1	1:11:A:VAL:HG23	18	0.83
(1,469)	1:10:A:PHE:HD2	1:11:A:VAL:HG11	18	0.83
(1,469)	1:10:A:PHE:HD2	1:11:A:VAL:HG12	18	0.83
(1,469)	1:10:A:PHE:HD2	1:11:A:VAL:HG13	18	0.83
(1,469)	1:10:A:PHE:HD2	1:11:A:VAL:HG21	18	0.83
(1,469)	1:10:A:PHE:HD2	1:11:A:VAL:HG22	18	0.83
(1,469)	1:10:A:PHE:HD2	1:11:A:VAL:HG23	18	0.83
(1,437)	1:8:A:GLU:HG2	2:20:B:TYR:HB2	16	0.83
(1,437)	1:8:A:GLU:HG2	2:20:B:TYR:HB3	16	0.83
(1,393)	1:7:A:PRO:HG2	1:10:A:PHE:HD1	13	0.83
(1,393)	1:7:A:PRO:HG2	1:10:A:PHE:HD2	13	0.83
(1,393)	1:7:A:PRO:HG3	1:10:A:PHE:HD1	13	0.83
(1,393)	1:7:A:PRO:HG3	1:10:A:PHE:HD2	13	0.83
(1,386)	1:4:A:GLY:H	2:19:B:PHE:HD1	2	0.83
(1,386)	1:4:A:GLY:H	2:19:B:PHE:HD2	2	0.83
(1,351)	2:16:B:PRO:HA	2:17:B:ILE:HD11	3	0.83
(1,351)	2:16:B:PRO:HA	2:17:B:ILE:HD12	3	0.83
(1,351)	2:16:B:PRO:HA	2:17:B:ILE:HD13	3	0.83
(1,310)	2:17:B:ILE:H	2:17:B:ILE:HD11	10	0.83
(1,310)	2:17:B:ILE:H	2:17:B:ILE:HD12	10	0.83
(1,310)	2:17:B:ILE:H	2:17:B:ILE:HD13	10	0.83
(1,173)	1:7:A:PRO:HB2	1:10:A:PHE:HD1	17	0.83
(1,173)	1:7:A:PRO:HB2	1:10:A:PHE:HD2	17	0.83
(1,170)	1:4:A:GLY:H	2:19:B:PHE:HE1	19	0.83
(1,170)	1:4:A:GLY:H	2:19:B:PHE:HE2	19	0.83
(1,150)	1:10:A:PHE:HD1	1:11:A:VAL:HG11	18	0.83
(1,150)	1:10:A:PHE:HD1	1:11:A:VAL:HG12	18	0.83
(1,150)	1:10:A:PHE:HD1	1:11:A:VAL:HG13	18	0.83
(1,150)	1:10:A:PHE:HD1	1:11:A:VAL:HG21	18	0.83
(1,150)	1:10:A:PHE:HD1	1:11:A:VAL:HG22	18	0.83
(1,150)	1:10:A:PHE:HD1	1:11:A:VAL:HG23	18	0.83
(1,150)	1:10:A:PHE:HD2	1:11:A:VAL:HG11	18	0.83
(1,150)	1:10:A:PHE:HD2	1:11:A:VAL:HG12	18	0.83
(1,150)	1:10:A:PHE:HD2	1:11:A:VAL:HG13	18	0.83
(1,150)	1:10:A:PHE:HD2	1:11:A:VAL:HG21	18	0.83
(1,150)	1:10:A:PHE:HD2	1:11:A:VAL:HG22	18	0.83
(1,150)	1:10:A:PHE:HD2	1:11:A:VAL:HG23	18	0.83
(1,118)	1:8:A:GLU:HG2	2:20:B:TYR:HB2	16	0.83
(1,118)	1:8:A:GLU:HG2	2:20:B:TYR:HB3	16	0.83
(1,74)	1:7:A:PRO:HG2	1:10:A:PHE:HD1	13	0.83
(1,74)	1:7:A:PRO:HG2	1:10:A:PHE:HD2	13	0.83
(1,74)	1:7:A:PRO:HG3	1:10:A:PHE:HD1	13	0.83

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,74)	1:7:A:PRO:HG3	1:10:A:PHE:HD2	13	0.83
(1,67)	1:4:A:GLY:H	2:19:B:PHE:HD1	2	0.83
(1,67)	1:4:A:GLY:H	2:19:B:PHE:HD2	2	0.83
(1,32)	2:16:B:PRO:HA	2:17:B:ILE:HD11	3	0.83
(1,32)	2:16:B:PRO:HA	2:17:B:ILE:HD12	3	0.83
(1,32)	2:16:B:PRO:HA	2:17:B:ILE:HD13	3	0.83
(1,1285)	2:38:D:ILE:H	2:38:D:ILE:HD11	15	0.82
(1,1285)	2:38:D:ILE:H	2:38:D:ILE:HD12	15	0.82
(1,1285)	2:38:D:ILE:H	2:38:D:ILE:HD13	15	0.82
(1,1058)	1:31:C:PHE:HD1	2:35:D:GLY:HA3	16	0.82
(1,1058)	1:31:C:PHE:HD2	2:35:D:GLY:HA3	16	0.82
(1,1036)	2:34:D:CYS:HB2	2:35:D:GLY:HA3	17	0.82
(1,1002)	1:30:C:TYR:HE1	2:39:D:SER:HB2	5	0.82
(1,1002)	1:30:C:TYR:HE2	2:39:D:SER:HB2	5	0.82
(1,966)	2:38:D:ILE:H	2:38:D:ILE:HD11	15	0.82
(1,966)	2:38:D:ILE:H	2:38:D:ILE:HD12	15	0.82
(1,966)	2:38:D:ILE:H	2:38:D:ILE:HD13	15	0.82
(1,739)	1:31:C:PHE:HD1	2:35:D:GLY:HA3	16	0.82
(1,739)	1:31:C:PHE:HD2	2:35:D:GLY:HA3	16	0.82
(1,717)	2:34:D:CYS:HB2	2:35:D:GLY:HA3	17	0.82
(1,683)	1:30:C:TYR:HE1	2:39:D:SER:HB2	5	0.82
(1,683)	1:30:C:TYR:HE2	2:39:D:SER:HB2	5	0.82
(1,653)	1:3:A:ALA:HA	2:19:B:PHE:HD1	17	0.82
(1,653)	1:3:A:ALA:HA	2:19:B:PHE:HD2	17	0.82
(1,519)	1:3:A:ALA:HB1	2:19:B:PHE:HB3	1	0.82
(1,519)	1:3:A:ALA:HB2	2:19:B:PHE:HB3	1	0.82
(1,519)	1:3:A:ALA:HB3	2:19:B:PHE:HB3	1	0.82
(1,519)	1:3:A:ALA:HB1	2:19:B:PHE:HB3	15	0.82
(1,519)	1:3:A:ALA:HB2	2:19:B:PHE:HB3	15	0.82
(1,519)	1:3:A:ALA:HB3	2:19:B:PHE:HB3	15	0.82
(1,515)	2:15:B:THR:HG21	2:17:B:ILE:HD11	13	0.82
(1,515)	2:15:B:THR:HG21	2:17:B:ILE:HD12	13	0.82
(1,515)	2:15:B:THR:HG21	2:17:B:ILE:HD13	13	0.82
(1,515)	2:15:B:THR:HG22	2:17:B:ILE:HD11	13	0.82
(1,515)	2:15:B:THR:HG22	2:17:B:ILE:HD12	13	0.82
(1,515)	2:15:B:THR:HG22	2:17:B:ILE:HD13	13	0.82
(1,515)	2:15:B:THR:HG23	2:17:B:ILE:HD11	13	0.82
(1,515)	2:15:B:THR:HG23	2:17:B:ILE:HD12	13	0.82
(1,515)	2:15:B:THR:HG23	2:17:B:ILE:HD13	13	0.82
(1,515)	2:15:B:THR:HG21	2:17:B:ILE:HD11	18	0.82
(1,515)	2:15:B:THR:HG21	2:17:B:ILE:HD12	18	0.82
(1,515)	2:15:B:THR:HG21	2:17:B:ILE:HD13	18	0.82

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,515)	2:15:B:THR:HG22	2:17:B:ILE:HD11	18	0.82
(1,515)	2:15:B:THR:HG22	2:17:B:ILE:HD12	18	0.82
(1,515)	2:15:B:THR:HG22	2:17:B:ILE:HD13	18	0.82
(1,515)	2:15:B:THR:HG23	2:17:B:ILE:HD11	18	0.82
(1,515)	2:15:B:THR:HG23	2:17:B:ILE:HD12	18	0.82
(1,515)	2:15:B:THR:HG23	2:17:B:ILE:HD13	18	0.82
(1,501)	1:4:A:GLY:H	1:6:A:VAL:H	15	0.82
(1,391)	1:3:A:ALA:HA	2:19:B:PHE:HD1	17	0.82
(1,391)	1:3:A:ALA:HA	2:19:B:PHE:HD2	17	0.82
(1,389)	1:6:A:VAL:HB	2:20:B:TYR:HD1	14	0.82
(1,389)	1:6:A:VAL:HB	2:20:B:TYR:HD2	14	0.82
(1,387)	1:6:A:VAL:HG11	1:7:A:PRO:HG2	14	0.82
(1,387)	1:6:A:VAL:HG11	1:7:A:PRO:HG3	14	0.82
(1,387)	1:6:A:VAL:HG12	1:7:A:PRO:HG2	14	0.82
(1,387)	1:6:A:VAL:HG12	1:7:A:PRO:HG3	14	0.82
(1,387)	1:6:A:VAL:HG13	1:7:A:PRO:HG2	14	0.82
(1,387)	1:6:A:VAL:HG13	1:7:A:PRO:HG3	14	0.82
(1,348)	1:11:A:VAL:HA	1:12:A:ARG:HD2	14	0.82
(1,348)	1:11:A:VAL:HA	1:12:A:ARG:HD3	14	0.82
(1,334)	1:3:A:ALA:HA	2:19:B:PHE:HD1	17	0.82
(1,334)	1:3:A:ALA:HA	2:19:B:PHE:HD2	17	0.82
(1,200)	1:3:A:ALA:HB1	2:19:B:PHE:HB3	1	0.82
(1,200)	1:3:A:ALA:HB2	2:19:B:PHE:HB3	1	0.82
(1,200)	1:3:A:ALA:HB3	2:19:B:PHE:HB3	1	0.82
(1,200)	1:3:A:ALA:HB1	2:19:B:PHE:HB3	15	0.82
(1,200)	1:3:A:ALA:HB2	2:19:B:PHE:HB3	15	0.82
(1,200)	1:3:A:ALA:HB3	2:19:B:PHE:HB3	15	0.82
(1,196)	2:15:B:THR:HG21	2:17:B:ILE:HD11	13	0.82
(1,196)	2:15:B:THR:HG21	2:17:B:ILE:HD12	13	0.82
(1,196)	2:15:B:THR:HG21	2:17:B:ILE:HD13	13	0.82
(1,196)	2:15:B:THR:HG22	2:17:B:ILE:HD11	13	0.82
(1,196)	2:15:B:THR:HG22	2:17:B:ILE:HD12	13	0.82
(1,196)	2:15:B:THR:HG22	2:17:B:ILE:HD13	13	0.82
(1,196)	2:15:B:THR:HG23	2:17:B:ILE:HD11	13	0.82
(1,196)	2:15:B:THR:HG23	2:17:B:ILE:HD12	13	0.82
(1,196)	2:15:B:THR:HG23	2:17:B:ILE:HD13	13	0.82
(1,196)	2:15:B:THR:HG21	2:17:B:ILE:HD11	18	0.82
(1,196)	2:15:B:THR:HG21	2:17:B:ILE:HD12	18	0.82
(1,196)	2:15:B:THR:HG21	2:17:B:ILE:HD13	18	0.82
(1,196)	2:15:B:THR:HG22	2:17:B:ILE:HD11	18	0.82
(1,196)	2:15:B:THR:HG22	2:17:B:ILE:HD12	18	0.82
(1,196)	2:15:B:THR:HG22	2:17:B:ILE:HD13	18	0.82

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,196)	2:15:B:THR:HG23	2:17:B:ILE:HD11	18	0.82
(1,196)	2:15:B:THR:HG23	2:17:B:ILE:HD12	18	0.82
(1,196)	2:15:B:THR:HG23	2:17:B:ILE:HD13	18	0.82
(1,182)	1:4:A:GLY:H	1:6:A:VAL:H	15	0.82
(1,72)	1:3:A:ALA:HA	2:19:B:PHE:HD1	17	0.82
(1,72)	1:3:A:ALA:HA	2:19:B:PHE:HD2	17	0.82
(1,70)	1:6:A:VAL:HB	2:20:B:TYR:HD1	14	0.82
(1,70)	1:6:A:VAL:HB	2:20:B:TYR:HD2	14	0.82
(1,68)	1:6:A:VAL:HG11	1:7:A:PRO:HG2	14	0.82
(1,68)	1:6:A:VAL:HG11	1:7:A:PRO:HG3	14	0.82
(1,68)	1:6:A:VAL:HG12	1:7:A:PRO:HG2	14	0.82
(1,68)	1:6:A:VAL:HG12	1:7:A:PRO:HG3	14	0.82
(1,68)	1:6:A:VAL:HG13	1:7:A:PRO:HG2	14	0.82
(1,68)	1:6:A:VAL:HG13	1:7:A:PRO:HG3	14	0.82
(1,29)	1:11:A:VAL:HA	1:12:A:ARG:HD2	14	0.82
(1,29)	1:11:A:VAL:HA	1:12:A:ARG:HD3	14	0.82
(1,1117)	1:27:C:VAL:HG11	1:28:C:PRO:HB3	15	0.81
(1,1117)	1:27:C:VAL:HG12	1:28:C:PRO:HB3	15	0.81
(1,1117)	1:27:C:VAL:HG13	1:28:C:PRO:HB3	15	0.81
(1,1051)	2:34:D:CYS:HB2	2:35:D:GLY:H	15	0.81
(1,1046)	1:27:C:VAL:HB	2:41:D:TYR:HE1	5	0.81
(1,1046)	1:27:C:VAL:HB	2:41:D:TYR:HE2	5	0.81
(1,1036)	2:34:D:CYS:HB2	2:35:D:GLY:HA3	15	0.81
(1,998)	1:22:C:GLY:HA3	2:39:D:SER:HB2	4	0.81
(1,995)	1:22:C:GLY:HA3	2:39:D:SER:HB2	4	0.81
(1,798)	1:27:C:VAL:HG11	1:28:C:PRO:HB3	15	0.81
(1,798)	1:27:C:VAL:HG12	1:28:C:PRO:HB3	15	0.81
(1,798)	1:27:C:VAL:HG13	1:28:C:PRO:HB3	15	0.81
(1,732)	2:34:D:CYS:HB2	2:35:D:GLY:H	15	0.81
(1,727)	1:27:C:VAL:HB	2:41:D:TYR:HE1	5	0.81
(1,727)	1:27:C:VAL:HB	2:41:D:TYR:HE2	5	0.81
(1,717)	2:34:D:CYS:HB2	2:35:D:GLY:HA3	15	0.81
(1,679)	1:22:C:GLY:HA3	2:39:D:SER:HB2	4	0.81
(1,676)	1:22:C:GLY:HA3	2:39:D:SER:HB2	4	0.81
(1,631)	2:17:B:ILE:HA	2:17:B:ILE:HD11	14	0.81
(1,631)	2:17:B:ILE:HA	2:17:B:ILE:HD12	14	0.81
(1,631)	2:17:B:ILE:HA	2:17:B:ILE:HD13	14	0.81
(1,510)	1:3:A:ALA:H	2:19:B:PHE:HD1	4	0.81
(1,510)	1:3:A:ALA:H	2:19:B:PHE:HD2	4	0.81
(1,503)	1:7:A:PRO:HB3	1:10:A:PHE:HE1	8	0.81
(1,503)	1:7:A:PRO:HB3	1:10:A:PHE:HE2	8	0.81
(1,492)	1:7:A:PRO:HB2	1:10:A:PHE:HD1	12	0.81

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,492)	1:7:A:PRO:HB2	1:10:A:PHE:HD2	12	0.81
(1,491)	1:7:A:PRO:HB2	1:10:A:PHE:HE1	1	0.81
(1,491)	1:7:A:PRO:HB2	1:10:A:PHE:HE2	1	0.81
(1,483)	1:1:A:GLY:H	2:20:B:TYR:HD1	13	0.81
(1,483)	1:1:A:GLY:H	2:20:B:TYR:HD2	13	0.81
(1,437)	1:8:A:GLU:HG2	2:20:B:TYR:HB2	12	0.81
(1,437)	1:8:A:GLU:HG2	2:20:B:TYR:HB3	12	0.81
(1,418)	1:1:A:GLY:H	2:18:B:SER:HB3	13	0.81
(1,395)	2:13:B:CYS:HB2	2:14:B:GLY:H	2	0.81
(1,380)	2:13:B:CYS:HB2	2:14:B:GLY:HA3	1	0.81
(1,371)	1:3:A:ALA:H	2:19:B:PHE:HD1	4	0.81
(1,371)	1:3:A:ALA:H	2:19:B:PHE:HD2	4	0.81
(1,312)	2:17:B:ILE:HA	2:17:B:ILE:HD11	14	0.81
(1,312)	2:17:B:ILE:HA	2:17:B:ILE:HD12	14	0.81
(1,312)	2:17:B:ILE:HA	2:17:B:ILE:HD13	14	0.81
(1,191)	1:3:A:ALA:H	2:19:B:PHE:HD1	4	0.81
(1,191)	1:3:A:ALA:H	2:19:B:PHE:HD2	4	0.81
(1,184)	1:7:A:PRO:HB3	1:10:A:PHE:HE1	8	0.81
(1,184)	1:7:A:PRO:HB3	1:10:A:PHE:HE2	8	0.81
(1,173)	1:7:A:PRO:HB2	1:10:A:PHE:HD1	12	0.81
(1,173)	1:7:A:PRO:HB2	1:10:A:PHE:HD2	12	0.81
(1,172)	1:7:A:PRO:HB2	1:10:A:PHE:HE1	1	0.81
(1,172)	1:7:A:PRO:HB2	1:10:A:PHE:HE2	1	0.81
(1,164)	1:1:A:GLY:H	2:20:B:TYR:HD1	13	0.81
(1,164)	1:1:A:GLY:H	2:20:B:TYR:HD2	13	0.81
(1,118)	1:8:A:GLU:HG2	2:20:B:TYR:HB2	12	0.81
(1,118)	1:8:A:GLU:HG2	2:20:B:TYR:HB3	12	0.81
(1,99)	1:1:A:GLY:H	2:18:B:SER:HB3	13	0.81
(1,76)	2:13:B:CYS:HB2	2:14:B:GLY:H	2	0.81
(1,61)	2:13:B:CYS:HB2	2:14:B:GLY:HA3	1	0.81
(1,52)	1:3:A:ALA:H	2:19:B:PHE:HD1	4	0.81
(1,52)	1:3:A:ALA:H	2:19:B:PHE:HD2	4	0.81
(1,1155)	1:27:C:VAL:HG11	1:28:C:PRO:HD3	1	0.8
(1,1155)	1:27:C:VAL:HG12	1:28:C:PRO:HD3	1	0.8
(1,1155)	1:27:C:VAL:HG13	1:28:C:PRO:HD3	1	0.8
(1,1036)	2:34:D:CYS:HB2	2:35:D:GLY:HA3	20	0.8
(1,1025)	1:27:C:VAL:HG21	2:41:D:TYR:HE1	1	0.8
(1,1025)	1:27:C:VAL:HG21	2:41:D:TYR:HE2	1	0.8
(1,1025)	1:27:C:VAL:HG22	2:41:D:TYR:HE1	1	0.8
(1,1025)	1:27:C:VAL:HG22	2:41:D:TYR:HE2	1	0.8
(1,1025)	1:27:C:VAL:HG23	2:41:D:TYR:HE1	1	0.8
(1,1025)	1:27:C:VAL:HG23	2:41:D:TYR:HE2	1	0.8

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,1004)	1:32:C:VAL:HA	1:33:C:ARG:HD2	5	0.8
(1,1004)	1:32:C:VAL:HA	1:33:C:ARG:HD3	5	0.8
(1,1000)	1:31:C:PHE:HE1	2:37:D:PRO:HD2	10	0.8
(1,1000)	1:31:C:PHE:HE2	2:37:D:PRO:HD2	10	0.8
(1,836)	1:27:C:VAL:HG11	1:28:C:PRO:HD3	1	0.8
(1,836)	1:27:C:VAL:HG12	1:28:C:PRO:HD3	1	0.8
(1,836)	1:27:C:VAL:HG13	1:28:C:PRO:HD3	1	0.8
(1,717)	2:34:D:CYS:HB2	2:35:D:GLY:HA3	20	0.8
(1,706)	1:27:C:VAL:HG21	2:41:D:TYR:HE1	1	0.8
(1,706)	1:27:C:VAL:HG21	2:41:D:TYR:HE2	1	0.8
(1,706)	1:27:C:VAL:HG22	2:41:D:TYR:HE1	1	0.8
(1,706)	1:27:C:VAL:HG22	2:41:D:TYR:HE2	1	0.8
(1,706)	1:27:C:VAL:HG23	2:41:D:TYR:HE1	1	0.8
(1,706)	1:27:C:VAL:HG23	2:41:D:TYR:HE2	1	0.8
(1,685)	1:32:C:VAL:HA	1:33:C:ARG:HD2	5	0.8
(1,685)	1:32:C:VAL:HA	1:33:C:ARG:HD3	5	0.8
(1,681)	1:31:C:PHE:HE1	2:37:D:PRO:HD2	10	0.8
(1,681)	1:31:C:PHE:HE2	2:37:D:PRO:HD2	10	0.8
(1,673)	2:17:B:ILE:HG13	2:40:D:PHE:HB2	13	0.8
(1,671)	2:17:B:ILE:HG13	2:40:D:PHE:HE1	13	0.8
(1,671)	2:17:B:ILE:HG13	2:40:D:PHE:HE2	13	0.8
(1,666)	2:17:B:ILE:HD11	2:40:D:PHE:HD1	17	0.8
(1,666)	2:17:B:ILE:HD11	2:40:D:PHE:HD2	17	0.8
(1,666)	2:17:B:ILE:HD12	2:40:D:PHE:HD1	17	0.8
(1,666)	2:17:B:ILE:HD12	2:40:D:PHE:HD2	17	0.8
(1,666)	2:17:B:ILE:HD13	2:40:D:PHE:HD1	17	0.8
(1,666)	2:17:B:ILE:HD13	2:40:D:PHE:HD2	17	0.8
(1,415)	1:6:A:VAL:HG11	2:20:B:TYR:HE1	7	0.8
(1,415)	1:6:A:VAL:HG11	2:20:B:TYR:HE2	7	0.8
(1,415)	1:6:A:VAL:HG12	2:20:B:TYR:HE1	7	0.8
(1,415)	1:6:A:VAL:HG12	2:20:B:TYR:HE2	7	0.8
(1,415)	1:6:A:VAL:HG13	2:20:B:TYR:HE1	7	0.8
(1,415)	1:6:A:VAL:HG13	2:20:B:TYR:HE2	7	0.8
(1,96)	1:6:A:VAL:HG11	2:20:B:TYR:HE1	7	0.8
(1,96)	1:6:A:VAL:HG11	2:20:B:TYR:HE2	7	0.8
(1,96)	1:6:A:VAL:HG12	2:20:B:TYR:HE1	7	0.8
(1,96)	1:6:A:VAL:HG12	2:20:B:TYR:HE2	7	0.8
(1,96)	1:6:A:VAL:HG13	2:20:B:TYR:HE1	7	0.8
(1,96)	1:6:A:VAL:HG13	2:20:B:TYR:HE2	7	0.8
(1,8)	2:17:B:ILE:HG13	2:40:D:PHE:HB2	13	0.8
(1,6)	2:17:B:ILE:HG13	2:40:D:PHE:HE1	13	0.8
(1,6)	2:17:B:ILE:HG13	2:40:D:PHE:HE2	13	0.8

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,1)	2:17:B:ILE:HD11	2:40:D:PHE:HD1	17	0.8
(1,1)	2:17:B:ILE:HD11	2:40:D:PHE:HD2	17	0.8
(1,1)	2:17:B:ILE:HD12	2:40:D:PHE:HD1	17	0.8
(1,1)	2:17:B:ILE:HD12	2:40:D:PHE:HD2	17	0.8
(1,1)	2:17:B:ILE:HD13	2:40:D:PHE:HD1	17	0.8
(1,1)	2:17:B:ILE:HD13	2:40:D:PHE:HD2	17	0.8
(1,1175)	1:24:C:ALA:HB1	2:40:D:PHE:HB3	2	0.79
(1,1175)	1:24:C:ALA:HB2	2:40:D:PHE:HB3	2	0.79
(1,1175)	1:24:C:ALA:HB3	2:40:D:PHE:HB3	2	0.79
(1,1168)	1:27:C:VAL:HG11	2:41:D:TYR:HB2	3	0.79
(1,1168)	1:27:C:VAL:HG11	2:41:D:TYR:HB3	3	0.79
(1,1168)	1:27:C:VAL:HG12	2:41:D:TYR:HB2	3	0.79
(1,1168)	1:27:C:VAL:HG12	2:41:D:TYR:HB3	3	0.79
(1,1168)	1:27:C:VAL:HG13	2:41:D:TYR:HB2	3	0.79
(1,1168)	1:27:C:VAL:HG13	2:41:D:TYR:HB3	3	0.79
(1,1099)	1:29:C:GLU:HG3	2:41:D:TYR:HD1	1	0.79
(1,1099)	1:29:C:GLU:HG3	2:41:D:TYR:HD2	1	0.79
(1,1043)	1:27:C:VAL:HG11	1:28:C:PRO:HG2	2	0.79
(1,1043)	1:27:C:VAL:HG11	1:28:C:PRO:HG3	2	0.79
(1,1043)	1:27:C:VAL:HG12	1:28:C:PRO:HG2	2	0.79
(1,1043)	1:27:C:VAL:HG12	1:28:C:PRO:HG3	2	0.79
(1,1043)	1:27:C:VAL:HG13	1:28:C:PRO:HG2	2	0.79
(1,1043)	1:27:C:VAL:HG13	1:28:C:PRO:HG3	2	0.79
(1,856)	1:24:C:ALA:HB1	2:40:D:PHE:HB3	2	0.79
(1,856)	1:24:C:ALA:HB2	2:40:D:PHE:HB3	2	0.79
(1,856)	1:24:C:ALA:HB3	2:40:D:PHE:HB3	2	0.79
(1,849)	1:27:C:VAL:HG11	2:41:D:TYR:HB2	3	0.79
(1,849)	1:27:C:VAL:HG11	2:41:D:TYR:HB3	3	0.79
(1,849)	1:27:C:VAL:HG12	2:41:D:TYR:HB2	3	0.79
(1,849)	1:27:C:VAL:HG12	2:41:D:TYR:HB3	3	0.79
(1,849)	1:27:C:VAL:HG13	2:41:D:TYR:HB2	3	0.79
(1,849)	1:27:C:VAL:HG13	2:41:D:TYR:HB3	3	0.79
(1,780)	1:29:C:GLU:HG3	2:41:D:TYR:HD1	1	0.79
(1,780)	1:29:C:GLU:HG3	2:41:D:TYR:HD2	1	0.79
(1,724)	1:27:C:VAL:HG11	1:28:C:PRO:HG2	2	0.79
(1,724)	1:27:C:VAL:HG11	1:28:C:PRO:HG3	2	0.79
(1,724)	1:27:C:VAL:HG12	1:28:C:PRO:HG2	2	0.79
(1,724)	1:27:C:VAL:HG12	1:28:C:PRO:HG3	2	0.79
(1,724)	1:27:C:VAL:HG13	1:28:C:PRO:HG2	2	0.79
(1,724)	1:27:C:VAL:HG13	1:28:C:PRO:HG3	2	0.79
(1,664)	2:19:B:PHE:HB2	2:38:D:ILE:HG13	4	0.79
(1,515)	2:15:B:THR:HG21	2:17:B:ILE:HD11	12	0.79

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,515)	2:15:B:THR:HG21	2:17:B:ILE:HD12	12	0.79
(1,515)	2:15:B:THR:HG21	2:17:B:ILE:HD13	12	0.79
(1,515)	2:15:B:THR:HG22	2:17:B:ILE:HD11	12	0.79
(1,515)	2:15:B:THR:HG22	2:17:B:ILE:HD12	12	0.79
(1,515)	2:15:B:THR:HG22	2:17:B:ILE:HD13	12	0.79
(1,515)	2:15:B:THR:HG23	2:17:B:ILE:HD11	12	0.79
(1,515)	2:15:B:THR:HG23	2:17:B:ILE:HD12	12	0.79
(1,515)	2:15:B:THR:HG23	2:17:B:ILE:HD13	12	0.79
(1,489)	1:4:A:GLY:H	2:19:B:PHE:HE1	12	0.79
(1,489)	1:4:A:GLY:H	2:19:B:PHE:HE2	12	0.79
(1,474)	1:6:A:VAL:H	2:19:B:PHE:HB2	11	0.79
(1,466)	1:4:A:GLY:HA2	1:5:A:HIS:HD2	1	0.79
(1,446)	1:8:A:GLU:HB3	2:20:B:TYR:HD1	6	0.79
(1,446)	1:8:A:GLU:HB3	2:20:B:TYR:HD2	6	0.79
(1,443)	1:8:A:GLU:HG3	2:20:B:TYR:HD1	3	0.79
(1,443)	1:8:A:GLU:HG3	2:20:B:TYR:HD2	3	0.79
(1,438)	1:8:A:GLU:HG2	2:19:B:PHE:HA	17	0.79
(1,348)	1:11:A:VAL:HA	1:12:A:ARG:HD2	8	0.79
(1,348)	1:11:A:VAL:HA	1:12:A:ARG:HD3	8	0.79
(1,196)	2:15:B:THR:HG21	2:17:B:ILE:HD11	12	0.79
(1,196)	2:15:B:THR:HG21	2:17:B:ILE:HD12	12	0.79
(1,196)	2:15:B:THR:HG21	2:17:B:ILE:HD13	12	0.79
(1,196)	2:15:B:THR:HG22	2:17:B:ILE:HD11	12	0.79
(1,196)	2:15:B:THR:HG22	2:17:B:ILE:HD12	12	0.79
(1,196)	2:15:B:THR:HG22	2:17:B:ILE:HD13	12	0.79
(1,196)	2:15:B:THR:HG23	2:17:B:ILE:HD11	12	0.79
(1,196)	2:15:B:THR:HG23	2:17:B:ILE:HD12	12	0.79
(1,196)	2:15:B:THR:HG23	2:17:B:ILE:HD13	12	0.79
(1,170)	1:4:A:GLY:H	2:19:B:PHE:HE1	12	0.79
(1,170)	1:4:A:GLY:H	2:19:B:PHE:HE2	12	0.79
(1,155)	1:6:A:VAL:H	2:19:B:PHE:HB2	11	0.79
(1,147)	1:4:A:GLY:HA2	1:5:A:HIS:HD2	1	0.79
(1,127)	1:8:A:GLU:HB3	2:20:B:TYR:HD1	6	0.79
(1,127)	1:8:A:GLU:HB3	2:20:B:TYR:HD2	6	0.79
(1,124)	1:8:A:GLU:HG3	2:20:B:TYR:HD1	3	0.79
(1,124)	1:8:A:GLU:HG3	2:20:B:TYR:HD2	3	0.79
(1,119)	1:8:A:GLU:HG2	2:19:B:PHE:HA	17	0.79
(1,29)	1:11:A:VAL:HA	1:12:A:ARG:HD2	8	0.79
(1,29)	1:11:A:VAL:HA	1:12:A:ARG:HD3	8	0.79
(1,17)	2:19:B:PHE:HB2	2:38:D:ILE:HG13	4	0.79
(1,1102)	1:29:C:GLU:HB3	2:41:D:TYR:HD1	17	0.78
(1,1102)	1:29:C:GLU:HB3	2:41:D:TYR:HD2	17	0.78

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,1051)	2:34:D:CYS:HB2	2:35:D:GLY:H	10	0.78
(1,1045)	1:27:C:VAL:HB	2:41:D:TYR:HD1	2	0.78
(1,1045)	1:27:C:VAL:HB	2:41:D:TYR:HD2	2	0.78
(1,1008)	2:18:B:SER:HA	2:38:D:ILE:HG13	3	0.78
(1,783)	1:29:C:GLU:HB3	2:41:D:TYR:HD1	17	0.78
(1,783)	1:29:C:GLU:HB3	2:41:D:TYR:HD2	17	0.78
(1,732)	2:34:D:CYS:HB2	2:35:D:GLY:H	10	0.78
(1,726)	1:27:C:VAL:HB	2:41:D:TYR:HD1	2	0.78
(1,726)	1:27:C:VAL:HB	2:41:D:TYR:HD2	2	0.78
(1,689)	2:18:B:SER:HA	2:38:D:ILE:HG13	3	0.78
(1,674)	2:17:B:ILE:HD11	2:40:D:PHE:HE1	11	0.78
(1,674)	2:17:B:ILE:HD11	2:40:D:PHE:HE2	11	0.78
(1,674)	2:17:B:ILE:HD12	2:40:D:PHE:HE1	11	0.78
(1,674)	2:17:B:ILE:HD12	2:40:D:PHE:HE2	11	0.78
(1,674)	2:17:B:ILE:HD13	2:40:D:PHE:HE1	11	0.78
(1,674)	2:17:B:ILE:HD13	2:40:D:PHE:HE2	11	0.78
(1,446)	1:8:A:GLU:HB3	2:20:B:TYR:HD1	8	0.78
(1,446)	1:8:A:GLU:HB3	2:20:B:TYR:HD2	8	0.78
(1,384)	1:10:A:PHE:HE1	2:16:B:PRO:HD2	18	0.78
(1,384)	1:10:A:PHE:HE2	2:16:B:PRO:HD2	18	0.78
(1,382)	1:10:A:PHE:HD1	2:16:B:PRO:HD2	1	0.78
(1,382)	1:10:A:PHE:HD2	2:16:B:PRO:HD2	1	0.78
(1,127)	1:8:A:GLU:HB3	2:20:B:TYR:HD1	8	0.78
(1,127)	1:8:A:GLU:HB3	2:20:B:TYR:HD2	8	0.78
(1,65)	1:10:A:PHE:HE1	2:16:B:PRO:HD2	18	0.78
(1,65)	1:10:A:PHE:HE2	2:16:B:PRO:HD2	18	0.78
(1,63)	1:10:A:PHE:HD1	2:16:B:PRO:HD2	1	0.78
(1,63)	1:10:A:PHE:HD2	2:16:B:PRO:HD2	1	0.78
(1,9)	2:17:B:ILE:HD11	2:40:D:PHE:HE1	11	0.78
(1,9)	2:17:B:ILE:HD11	2:40:D:PHE:HE2	11	0.78
(1,9)	2:17:B:ILE:HD12	2:40:D:PHE:HE1	11	0.78
(1,9)	2:17:B:ILE:HD12	2:40:D:PHE:HE2	11	0.78
(1,9)	2:17:B:ILE:HD13	2:40:D:PHE:HE1	11	0.78
(1,9)	2:17:B:ILE:HD13	2:40:D:PHE:HE2	11	0.78
(1,1278)	2:37:D:PRO:HB3	2:37:D:PRO:HD3	13	0.77
(1,1175)	1:24:C:ALA:HB1	2:40:D:PHE:HB3	10	0.77
(1,1175)	1:24:C:ALA:HB2	2:40:D:PHE:HB3	10	0.77
(1,1175)	1:24:C:ALA:HB3	2:40:D:PHE:HB3	10	0.77
(1,1157)	1:25:C:GLY:H	1:27:C:VAL:H	9	0.77
(1,1155)	1:27:C:VAL:HG11	1:28:C:PRO:HD3	16	0.77
(1,1155)	1:27:C:VAL:HG12	1:28:C:PRO:HD3	16	0.77
(1,1155)	1:27:C:VAL:HG13	1:28:C:PRO:HD3	16	0.77

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,1105)	1:24:C:ALA:HB1	1:25:C:GLY:H	7	0.77
(1,1105)	1:24:C:ALA:HB2	1:25:C:GLY:H	7	0.77
(1,1105)	1:24:C:ALA:HB3	1:25:C:GLY:H	7	0.77
(1,1094)	1:29:C:GLU:HG2	2:40:D:PHE:HA	10	0.77
(1,1094)	1:29:C:GLU:HG2	2:40:D:PHE:HA	20	0.77
(1,1073)	2:18:B:SER:HB3	2:38:D:ILE:HD11	16	0.77
(1,1073)	2:18:B:SER:HB3	2:38:D:ILE:HD12	16	0.77
(1,1073)	2:18:B:SER:HB3	2:38:D:ILE:HD13	16	0.77
(1,959)	2:37:D:PRO:HB3	2:37:D:PRO:HD3	13	0.77
(1,856)	1:24:C:ALA:HB1	2:40:D:PHE:HB3	10	0.77
(1,856)	1:24:C:ALA:HB2	2:40:D:PHE:HB3	10	0.77
(1,856)	1:24:C:ALA:HB3	2:40:D:PHE:HB3	10	0.77
(1,838)	1:25:C:GLY:H	1:27:C:VAL:H	9	0.77
(1,836)	1:27:C:VAL:HG11	1:28:C:PRO:HD3	16	0.77
(1,836)	1:27:C:VAL:HG12	1:28:C:PRO:HD3	16	0.77
(1,836)	1:27:C:VAL:HG13	1:28:C:PRO:HD3	16	0.77
(1,786)	1:24:C:ALA:HB1	1:25:C:GLY:H	7	0.77
(1,786)	1:24:C:ALA:HB2	1:25:C:GLY:H	7	0.77
(1,786)	1:24:C:ALA:HB3	1:25:C:GLY:H	7	0.77
(1,775)	1:29:C:GLU:HG2	2:40:D:PHE:HA	10	0.77
(1,775)	1:29:C:GLU:HG2	2:40:D:PHE:HA	20	0.77
(1,754)	2:18:B:SER:HB3	2:38:D:ILE:HD11	16	0.77
(1,754)	2:18:B:SER:HB3	2:38:D:ILE:HD12	16	0.77
(1,754)	2:18:B:SER:HB3	2:38:D:ILE:HD13	16	0.77
(1,631)	2:17:B:ILE:HA	2:17:B:ILE:HD11	3	0.77
(1,631)	2:17:B:ILE:HA	2:17:B:ILE:HD12	3	0.77
(1,631)	2:17:B:ILE:HA	2:17:B:ILE:HD13	3	0.77
(1,501)	1:4:A:GLY:H	1:6:A:VAL:H	9	0.77
(1,483)	1:1:A:GLY:H	2:20:B:TYR:HD1	19	0.77
(1,483)	1:1:A:GLY:H	2:20:B:TYR:HD2	19	0.77
(1,449)	1:3:A:ALA:HB1	1:4:A:GLY:H	9	0.77
(1,449)	1:3:A:ALA:HB2	1:4:A:GLY:H	9	0.77
(1,449)	1:3:A:ALA:HB3	1:4:A:GLY:H	9	0.77
(1,405)	1:1:A:GLY:H	2:18:B:SER:HB2	1	0.77
(1,390)	1:6:A:VAL:HB	2:20:B:TYR:HE1	15	0.77
(1,390)	1:6:A:VAL:HB	2:20:B:TYR:HE2	15	0.77
(1,354)	1:10:A:PHE:HD1	2:16:B:PRO:HB3	5	0.77
(1,354)	1:10:A:PHE:HD2	2:16:B:PRO:HB3	5	0.77
(1,312)	2:17:B:ILE:HA	2:17:B:ILE:HD11	3	0.77
(1,312)	2:17:B:ILE:HA	2:17:B:ILE:HD12	3	0.77
(1,312)	2:17:B:ILE:HA	2:17:B:ILE:HD13	3	0.77
(1,182)	1:4:A:GLY:H	1:6:A:VAL:H	9	0.77

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,164)	1:1:A:GLY:H	2:20:B:TYR:HD1	19	0.77
(1,164)	1:1:A:GLY:H	2:20:B:TYR:HD2	19	0.77
(1,130)	1:3:A:ALA:HB1	1:4:A:GLY:H	9	0.77
(1,130)	1:3:A:ALA:HB2	1:4:A:GLY:H	9	0.77
(1,130)	1:3:A:ALA:HB3	1:4:A:GLY:H	9	0.77
(1,86)	1:1:A:GLY:H	2:18:B:SER:HB2	1	0.77
(1,71)	1:6:A:VAL:HB	2:20:B:TYR:HE1	15	0.77
(1,71)	1:6:A:VAL:HB	2:20:B:TYR:HE2	15	0.77
(1,35)	1:10:A:PHE:HD1	2:16:B:PRO:HB3	5	0.77
(1,35)	1:10:A:PHE:HD2	2:16:B:PRO:HB3	5	0.77
(1,1076)	1:25:C:GLY:HA3	1:26:C:HIS:HD2	19	0.76
(1,1071)	1:27:C:VAL:HG11	2:41:D:TYR:HE1	6	0.76
(1,1071)	1:27:C:VAL:HG11	2:41:D:TYR:HE2	6	0.76
(1,1071)	1:27:C:VAL:HG12	2:41:D:TYR:HE1	6	0.76
(1,1071)	1:27:C:VAL:HG12	2:41:D:TYR:HE2	6	0.76
(1,1071)	1:27:C:VAL:HG13	2:41:D:TYR:HE1	6	0.76
(1,1071)	1:27:C:VAL:HG13	2:41:D:TYR:HE2	6	0.76
(1,1063)	1:22:C:GLY:H	2:39:D:SER:HA	14	0.76
(1,1055)	1:28:C:PRO:HG2	1:31:C:PHE:HE1	19	0.76
(1,1055)	1:28:C:PRO:HG2	1:31:C:PHE:HE2	19	0.76
(1,1055)	1:28:C:PRO:HG3	1:31:C:PHE:HE1	19	0.76
(1,1055)	1:28:C:PRO:HG3	1:31:C:PHE:HE2	19	0.76
(1,1008)	2:18:B:SER:HA	2:38:D:ILE:HG13	20	0.76
(1,757)	1:25:C:GLY:HA3	1:26:C:HIS:HD2	19	0.76
(1,752)	1:27:C:VAL:HG11	2:41:D:TYR:HE1	6	0.76
(1,752)	1:27:C:VAL:HG11	2:41:D:TYR:HE2	6	0.76
(1,752)	1:27:C:VAL:HG12	2:41:D:TYR:HE1	6	0.76
(1,752)	1:27:C:VAL:HG12	2:41:D:TYR:HE2	6	0.76
(1,752)	1:27:C:VAL:HG13	2:41:D:TYR:HE1	6	0.76
(1,752)	1:27:C:VAL:HG13	2:41:D:TYR:HE2	6	0.76
(1,744)	1:22:C:GLY:H	2:39:D:SER:HA	14	0.76
(1,736)	1:28:C:PRO:HG2	1:31:C:PHE:HE1	19	0.76
(1,736)	1:28:C:PRO:HG2	1:31:C:PHE:HE2	19	0.76
(1,736)	1:28:C:PRO:HG3	1:31:C:PHE:HE1	19	0.76
(1,736)	1:28:C:PRO:HG3	1:31:C:PHE:HE2	19	0.76
(1,689)	2:18:B:SER:HA	2:38:D:ILE:HG13	20	0.76
(1,667)	2:15:B:THR:HG21	2:40:D:PHE:HE1	15	0.76
(1,667)	2:15:B:THR:HG21	2:40:D:PHE:HE2	15	0.76
(1,667)	2:15:B:THR:HG22	2:40:D:PHE:HE1	15	0.76
(1,667)	2:15:B:THR:HG22	2:40:D:PHE:HE2	15	0.76
(1,667)	2:15:B:THR:HG23	2:40:D:PHE:HE1	15	0.76
(1,667)	2:15:B:THR:HG23	2:40:D:PHE:HE2	15	0.76

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,631)	2:17:B:ILE:HA	2:17:B:ILE:HD11	17	0.76
(1,631)	2:17:B:ILE:HA	2:17:B:ILE:HD12	17	0.76
(1,631)	2:17:B:ILE:HA	2:17:B:ILE:HD13	17	0.76
(1,595)	1:12:A:ARG:HA	1:12:A:ARG:HE	17	0.76
(1,519)	1:3:A:ALA:HB1	2:19:B:PHE:HB3	2	0.76
(1,519)	1:3:A:ALA:HB2	2:19:B:PHE:HB3	2	0.76
(1,519)	1:3:A:ALA:HB3	2:19:B:PHE:HB3	2	0.76
(1,441)	1:8:A:GLU:HG3	2:20:B:TYR:HB2	13	0.76
(1,441)	1:8:A:GLU:HG3	2:20:B:TYR:HB3	13	0.76
(1,438)	1:8:A:GLU:HG2	2:19:B:PHE:HA	8	0.76
(1,395)	2:13:B:CYS:HB2	2:14:B:GLY:H	19	0.76
(1,359)	1:9:A:TYR:HD1	2:18:B:SER:HA	9	0.76
(1,359)	1:9:A:TYR:HD2	2:18:B:SER:HA	9	0.76
(1,312)	2:17:B:ILE:HA	2:17:B:ILE:HD11	17	0.76
(1,312)	2:17:B:ILE:HA	2:17:B:ILE:HD12	17	0.76
(1,312)	2:17:B:ILE:HA	2:17:B:ILE:HD13	17	0.76
(1,276)	1:12:A:ARG:HA	1:12:A:ARG:HE	17	0.76
(1,200)	1:3:A:ALA:HB1	2:19:B:PHE:HB3	2	0.76
(1,200)	1:3:A:ALA:HB2	2:19:B:PHE:HB3	2	0.76
(1,200)	1:3:A:ALA:HB3	2:19:B:PHE:HB3	2	0.76
(1,122)	1:8:A:GLU:HG3	2:20:B:TYR:HB2	13	0.76
(1,122)	1:8:A:GLU:HG3	2:20:B:TYR:HB3	13	0.76
(1,119)	1:8:A:GLU:HG2	2:19:B:PHE:HA	8	0.76
(1,76)	2:13:B:CYS:HB2	2:14:B:GLY:H	19	0.76
(1,40)	1:9:A:TYR:HD1	2:18:B:SER:HA	9	0.76
(1,40)	1:9:A:TYR:HD2	2:18:B:SER:HA	9	0.76
(1,2)	2:15:B:THR:HG21	2:40:D:PHE:HE1	15	0.76
(1,2)	2:15:B:THR:HG21	2:40:D:PHE:HE2	15	0.76
(1,2)	2:15:B:THR:HG22	2:40:D:PHE:HE1	15	0.76
(1,2)	2:15:B:THR:HG22	2:40:D:PHE:HE2	15	0.76
(1,2)	2:15:B:THR:HG23	2:40:D:PHE:HE1	15	0.76
(1,2)	2:15:B:THR:HG23	2:40:D:PHE:HE2	15	0.76
(1,1096)	1:31:C:PHE:HD1	2:37:D:PRO:HG2	20	0.75
(1,1096)	1:31:C:PHE:HD1	2:37:D:PRO:HG3	20	0.75
(1,1096)	1:31:C:PHE:HD2	2:37:D:PRO:HG2	20	0.75
(1,1096)	1:31:C:PHE:HD2	2:37:D:PRO:HG3	20	0.75
(1,1042)	1:25:C:GLY:H	2:40:D:PHE:HD1	18	0.75
(1,1042)	1:25:C:GLY:H	2:40:D:PHE:HD2	18	0.75
(1,777)	1:31:C:PHE:HD1	2:37:D:PRO:HG2	20	0.75
(1,777)	1:31:C:PHE:HD1	2:37:D:PRO:HG3	20	0.75
(1,777)	1:31:C:PHE:HD2	2:37:D:PRO:HG2	20	0.75
(1,777)	1:31:C:PHE:HD2	2:37:D:PRO:HG3	20	0.75

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,723)	1:25:C:GLY:H	2:40:D:PHE:HD1	18	0.75
(1,723)	1:25:C:GLY:H	2:40:D:PHE:HD2	18	0.75
(1,668)	1:3:A:ALA:HB1	1:32:C:VAL:HA	16	0.75
(1,668)	1:3:A:ALA:HB2	1:32:C:VAL:HA	16	0.75
(1,668)	1:3:A:ALA:HB3	1:32:C:VAL:HA	16	0.75
(1,663)	2:19:B:PHE:HD1	2:38:D:ILE:HG13	8	0.75
(1,663)	2:19:B:PHE:HD2	2:38:D:ILE:HG13	8	0.75
(1,449)	1:3:A:ALA:HB1	1:4:A:GLY:H	6	0.75
(1,449)	1:3:A:ALA:HB2	1:4:A:GLY:H	6	0.75
(1,449)	1:3:A:ALA:HB3	1:4:A:GLY:H	6	0.75
(1,449)	1:3:A:ALA:HB1	1:4:A:GLY:H	12	0.75
(1,449)	1:3:A:ALA:HB2	1:4:A:GLY:H	12	0.75
(1,449)	1:3:A:ALA:HB3	1:4:A:GLY:H	12	0.75
(1,446)	1:8:A:GLU:HB3	2:20:B:TYR:HD1	5	0.75
(1,446)	1:8:A:GLU:HB3	2:20:B:TYR:HD2	5	0.75
(1,438)	1:8:A:GLU:HG2	2:19:B:PHE:HA	3	0.75
(1,438)	1:8:A:GLU:HG2	2:19:B:PHE:HA	13	0.75
(1,402)	1:10:A:PHE:HD1	2:14:B:GLY:HA3	14	0.75
(1,402)	1:10:A:PHE:HD2	2:14:B:GLY:HA3	14	0.75
(1,359)	1:9:A:TYR:HD1	2:18:B:SER:HA	19	0.75
(1,359)	1:9:A:TYR:HD2	2:18:B:SER:HA	19	0.75
(1,130)	1:3:A:ALA:HB1	1:4:A:GLY:H	6	0.75
(1,130)	1:3:A:ALA:HB2	1:4:A:GLY:H	6	0.75
(1,130)	1:3:A:ALA:HB3	1:4:A:GLY:H	6	0.75
(1,130)	1:3:A:ALA:HB1	1:4:A:GLY:H	12	0.75
(1,130)	1:3:A:ALA:HB2	1:4:A:GLY:H	12	0.75
(1,130)	1:3:A:ALA:HB3	1:4:A:GLY:H	12	0.75
(1,127)	1:8:A:GLU:HB3	2:20:B:TYR:HD1	5	0.75
(1,127)	1:8:A:GLU:HB3	2:20:B:TYR:HD2	5	0.75
(1,119)	1:8:A:GLU:HG2	2:19:B:PHE:HA	3	0.75
(1,119)	1:8:A:GLU:HG2	2:19:B:PHE:HA	13	0.75
(1,83)	1:10:A:PHE:HD1	2:14:B:GLY:HA3	14	0.75
(1,83)	1:10:A:PHE:HD2	2:14:B:GLY:HA3	14	0.75
(1,40)	1:9:A:TYR:HD1	2:18:B:SER:HA	19	0.75
(1,40)	1:9:A:TYR:HD2	2:18:B:SER:HA	19	0.75
(1,16)	2:19:B:PHE:HD1	2:38:D:ILE:HG13	8	0.75
(1,16)	2:19:B:PHE:HD2	2:38:D:ILE:HG13	8	0.75
(1,3)	1:3:A:ALA:HB1	1:32:C:VAL:HA	16	0.75
(1,3)	1:3:A:ALA:HB2	1:32:C:VAL:HA	16	0.75
(1,3)	1:3:A:ALA:HB3	1:32:C:VAL:HA	16	0.75
(1,1155)	1:27:C:VAL:HG11	1:28:C:PRO:HD3	15	0.74
(1,1155)	1:27:C:VAL:HG12	1:28:C:PRO:HD3	15	0.74

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,1155)	1:27:C:VAL:HG13	1:28:C:PRO:HD3	15	0.74
(1,1063)	1:22:C:GLY:H	2:39:D:SER:HA	18	0.74
(1,1051)	2:34:D:CYS:HB2	2:35:D:GLY:H	9	0.74
(1,836)	1:27:C:VAL:HG11	1:28:C:PRO:HD3	15	0.74
(1,836)	1:27:C:VAL:HG12	1:28:C:PRO:HD3	15	0.74
(1,836)	1:27:C:VAL:HG13	1:28:C:PRO:HD3	15	0.74
(1,744)	1:22:C:GLY:H	2:39:D:SER:HA	18	0.74
(1,732)	2:34:D:CYS:HB2	2:35:D:GLY:H	9	0.74
(1,631)	2:17:B:ILE:HA	2:17:B:ILE:HD11	6	0.74
(1,631)	2:17:B:ILE:HA	2:17:B:ILE:HD12	6	0.74
(1,631)	2:17:B:ILE:HA	2:17:B:ILE:HD13	6	0.74
(1,453)	1:10:A:PHE:HD1	2:15:B:THR:HG21	5	0.74
(1,453)	1:10:A:PHE:HD1	2:15:B:THR:HG22	5	0.74
(1,453)	1:10:A:PHE:HD1	2:15:B:THR:HG23	5	0.74
(1,453)	1:10:A:PHE:HD2	2:15:B:THR:HG21	5	0.74
(1,453)	1:10:A:PHE:HD2	2:15:B:THR:HG22	5	0.74
(1,453)	1:10:A:PHE:HD2	2:15:B:THR:HG23	5	0.74
(1,415)	1:6:A:VAL:HG11	2:20:B:TYR:HE1	13	0.74
(1,415)	1:6:A:VAL:HG11	2:20:B:TYR:HE2	13	0.74
(1,415)	1:6:A:VAL:HG12	2:20:B:TYR:HE1	13	0.74
(1,415)	1:6:A:VAL:HG12	2:20:B:TYR:HE2	13	0.74
(1,415)	1:6:A:VAL:HG13	2:20:B:TYR:HE1	13	0.74
(1,415)	1:6:A:VAL:HG13	2:20:B:TYR:HE2	13	0.74
(1,312)	2:17:B:ILE:HA	2:17:B:ILE:HD11	6	0.74
(1,312)	2:17:B:ILE:HA	2:17:B:ILE:HD12	6	0.74
(1,312)	2:17:B:ILE:HA	2:17:B:ILE:HD13	6	0.74
(1,134)	1:10:A:PHE:HD1	2:15:B:THR:HG21	5	0.74
(1,134)	1:10:A:PHE:HD1	2:15:B:THR:HG22	5	0.74
(1,134)	1:10:A:PHE:HD1	2:15:B:THR:HG23	5	0.74
(1,134)	1:10:A:PHE:HD2	2:15:B:THR:HG21	5	0.74
(1,134)	1:10:A:PHE:HD2	2:15:B:THR:HG22	5	0.74
(1,134)	1:10:A:PHE:HD2	2:15:B:THR:HG23	5	0.74
(1,96)	1:6:A:VAL:HG11	2:20:B:TYR:HE1	13	0.74
(1,96)	1:6:A:VAL:HG11	2:20:B:TYR:HE2	13	0.74
(1,96)	1:6:A:VAL:HG12	2:20:B:TYR:HE1	13	0.74
(1,96)	1:6:A:VAL:HG12	2:20:B:TYR:HE2	13	0.74
(1,96)	1:6:A:VAL:HG13	2:20:B:TYR:HE1	13	0.74
(1,96)	1:6:A:VAL:HG13	2:20:B:TYR:HE2	13	0.74
(1,1071)	1:27:C:VAL:HG11	2:41:D:TYR:HE1	12	0.73
(1,1071)	1:27:C:VAL:HG11	2:41:D:TYR:HE2	12	0.73
(1,1071)	1:27:C:VAL:HG12	2:41:D:TYR:HE1	12	0.73
(1,1071)	1:27:C:VAL:HG12	2:41:D:TYR:HE2	12	0.73

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,1071)	1:27:C:VAL:HG13	2:41:D:TYR:HE1	12	0.73
(1,1071)	1:27:C:VAL:HG13	2:41:D:TYR:HE2	12	0.73
(1,1049)	1:28:C:PRO:HG2	1:31:C:PHE:HD1	11	0.73
(1,1049)	1:28:C:PRO:HG2	1:31:C:PHE:HD2	11	0.73
(1,1049)	1:28:C:PRO:HG3	1:31:C:PHE:HD1	11	0.73
(1,1049)	1:28:C:PRO:HG3	1:31:C:PHE:HD2	11	0.73
(1,1004)	1:32:C:VAL:HA	1:33:C:ARG:HD2	2	0.73
(1,1004)	1:32:C:VAL:HA	1:33:C:ARG:HD3	2	0.73
(1,752)	1:27:C:VAL:HG11	2:41:D:TYR:HE1	12	0.73
(1,752)	1:27:C:VAL:HG11	2:41:D:TYR:HE2	12	0.73
(1,752)	1:27:C:VAL:HG12	2:41:D:TYR:HE1	12	0.73
(1,752)	1:27:C:VAL:HG12	2:41:D:TYR:HE2	12	0.73
(1,752)	1:27:C:VAL:HG13	2:41:D:TYR:HE1	12	0.73
(1,752)	1:27:C:VAL:HG13	2:41:D:TYR:HE2	12	0.73
(1,730)	1:28:C:PRO:HG2	1:31:C:PHE:HD1	11	0.73
(1,730)	1:28:C:PRO:HG2	1:31:C:PHE:HD2	11	0.73
(1,730)	1:28:C:PRO:HG3	1:31:C:PHE:HD1	11	0.73
(1,730)	1:28:C:PRO:HG3	1:31:C:PHE:HD2	11	0.73
(1,685)	1:32:C:VAL:HA	1:33:C:ARG:HD2	2	0.73
(1,685)	1:32:C:VAL:HA	1:33:C:ARG:HD3	2	0.73
(1,674)	2:17:B:ILE:HD11	2:40:D:PHE:HE1	1	0.73
(1,674)	2:17:B:ILE:HD11	2:40:D:PHE:HE2	1	0.73
(1,674)	2:17:B:ILE:HD12	2:40:D:PHE:HE1	1	0.73
(1,674)	2:17:B:ILE:HD12	2:40:D:PHE:HE2	1	0.73
(1,674)	2:17:B:ILE:HD13	2:40:D:PHE:HE1	1	0.73
(1,674)	2:17:B:ILE:HD13	2:40:D:PHE:HE2	1	0.73
(1,673)	2:17:B:ILE:HG13	2:40:D:PHE:HB2	15	0.73
(1,672)	2:17:B:ILE:HG13	2:40:D:PHE:HD1	10	0.73
(1,672)	2:17:B:ILE:HG13	2:40:D:PHE:HD2	10	0.73
(1,662)	2:19:B:PHE:HE1	2:38:D:ILE:HG13	13	0.73
(1,662)	2:19:B:PHE:HE2	2:38:D:ILE:HG13	13	0.73
(1,621)	1:6:A:VAL:HB	1:7:A:PRO:HD2	18	0.73
(1,461)	1:6:A:VAL:HG11	1:7:A:PRO:HB3	10	0.73
(1,461)	1:6:A:VAL:HG12	1:7:A:PRO:HB3	10	0.73
(1,461)	1:6:A:VAL:HG13	1:7:A:PRO:HB3	10	0.73
(1,446)	1:8:A:GLU:HB3	2:20:B:TYR:HD1	10	0.73
(1,446)	1:8:A:GLU:HB3	2:20:B:TYR:HD2	10	0.73
(1,395)	2:13:B:CYS:HB2	2:14:B:GLY:H	7	0.73
(1,386)	1:4:A:GLY:H	2:19:B:PHE:HD1	4	0.73
(1,386)	1:4:A:GLY:H	2:19:B:PHE:HD2	4	0.73
(1,302)	1:6:A:VAL:HB	1:7:A:PRO:HD2	18	0.73
(1,142)	1:6:A:VAL:HG11	1:7:A:PRO:HB3	10	0.73

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,142)	1:6:A:VAL:HG12	1:7:A:PRO:HB3	10	0.73
(1,142)	1:6:A:VAL:HG13	1:7:A:PRO:HB3	10	0.73
(1,127)	1:8:A:GLU:HB3	2:20:B:TYR:HD1	10	0.73
(1,127)	1:8:A:GLU:HB3	2:20:B:TYR:HD2	10	0.73
(1,76)	2:13:B:CYS:HB2	2:14:B:GLY:H	7	0.73
(1,67)	1:4:A:GLY:H	2:19:B:PHE:HD1	4	0.73
(1,67)	1:4:A:GLY:H	2:19:B:PHE:HD2	4	0.73
(1,15)	2:19:B:PHE:HE1	2:38:D:ILE:HG13	13	0.73
(1,15)	2:19:B:PHE:HE2	2:38:D:ILE:HG13	13	0.73
(1,9)	2:17:B:ILE:HD11	2:40:D:PHE:HE1	1	0.73
(1,9)	2:17:B:ILE:HD11	2:40:D:PHE:HE2	1	0.73
(1,9)	2:17:B:ILE:HD12	2:40:D:PHE:HE1	1	0.73
(1,9)	2:17:B:ILE:HD12	2:40:D:PHE:HE2	1	0.73
(1,9)	2:17:B:ILE:HD13	2:40:D:PHE:HE1	1	0.73
(1,9)	2:17:B:ILE:HD13	2:40:D:PHE:HE2	1	0.73
(1,8)	2:17:B:ILE:HG13	2:40:D:PHE:HB2	15	0.73
(1,7)	2:17:B:ILE:HG13	2:40:D:PHE:HD1	10	0.73
(1,7)	2:17:B:ILE:HG13	2:40:D:PHE:HD2	10	0.73
(1,1147)	1:28:C:PRO:HB2	1:31:C:PHE:HE1	8	0.72
(1,1147)	1:28:C:PRO:HB2	1:31:C:PHE:HE2	8	0.72
(1,1121)	1:26:C:HIS:HD2	1:27:C:VAL:HG11	14	0.72
(1,1121)	1:26:C:HIS:HD2	1:27:C:VAL:HG12	14	0.72
(1,1121)	1:26:C:HIS:HD2	1:27:C:VAL:HG13	14	0.72
(1,1116)	1:26:C:HIS:HD2	1:27:C:VAL:HG21	8	0.72
(1,1116)	1:26:C:HIS:HD2	1:27:C:VAL:HG22	8	0.72
(1,1116)	1:26:C:HIS:HD2	1:27:C:VAL:HG23	8	0.72
(1,1063)	1:22:C:GLY:H	2:39:D:SER:HA	19	0.72
(1,1015)	1:30:C:TYR:HD1	2:39:D:SER:HA	3	0.72
(1,1015)	1:30:C:TYR:HD2	2:39:D:SER:HA	3	0.72
(1,1015)	1:30:C:TYR:HD1	2:39:D:SER:HA	5	0.72
(1,1015)	1:30:C:TYR:HD2	2:39:D:SER:HA	5	0.72
(1,998)	1:22:C:GLY:HA3	2:39:D:SER:HB2	6	0.72
(1,995)	1:22:C:GLY:HA3	2:39:D:SER:HB2	6	0.72
(1,828)	1:28:C:PRO:HB2	1:31:C:PHE:HE1	8	0.72
(1,828)	1:28:C:PRO:HB2	1:31:C:PHE:HE2	8	0.72
(1,802)	1:26:C:HIS:HD2	1:27:C:VAL:HG11	14	0.72
(1,802)	1:26:C:HIS:HD2	1:27:C:VAL:HG12	14	0.72
(1,802)	1:26:C:HIS:HD2	1:27:C:VAL:HG13	14	0.72
(1,797)	1:26:C:HIS:HD2	1:27:C:VAL:HG21	8	0.72
(1,797)	1:26:C:HIS:HD2	1:27:C:VAL:HG22	8	0.72
(1,797)	1:26:C:HIS:HD2	1:27:C:VAL:HG23	8	0.72
(1,744)	1:22:C:GLY:H	2:39:D:SER:HA	19	0.72

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,696)	1:30:C:TYR:HD1	2:39:D:SER:HA	3	0.72
(1,696)	1:30:C:TYR:HD2	2:39:D:SER:HA	3	0.72
(1,696)	1:30:C:TYR:HD1	2:39:D:SER:HA	5	0.72
(1,696)	1:30:C:TYR:HD2	2:39:D:SER:HA	5	0.72
(1,679)	1:22:C:GLY:HA3	2:39:D:SER:HB2	6	0.72
(1,676)	1:22:C:GLY:HA3	2:39:D:SER:HB2	6	0.72
(1,448)	1:4:A:GLY:H	2:19:B:PHE:H	2	0.72
(1,393)	1:7:A:PRO:HG2	1:10:A:PHE:HD1	18	0.72
(1,393)	1:7:A:PRO:HG2	1:10:A:PHE:HD2	18	0.72
(1,393)	1:7:A:PRO:HG3	1:10:A:PHE:HD1	18	0.72
(1,393)	1:7:A:PRO:HG3	1:10:A:PHE:HD2	18	0.72
(1,387)	1:6:A:VAL:HG11	1:7:A:PRO:HG2	9	0.72
(1,387)	1:6:A:VAL:HG11	1:7:A:PRO:HG3	9	0.72
(1,387)	1:6:A:VAL:HG12	1:7:A:PRO:HG2	9	0.72
(1,387)	1:6:A:VAL:HG12	1:7:A:PRO:HG3	9	0.72
(1,387)	1:6:A:VAL:HG13	1:7:A:PRO:HG2	9	0.72
(1,387)	1:6:A:VAL:HG13	1:7:A:PRO:HG3	9	0.72
(1,129)	1:4:A:GLY:H	2:19:B:PHE:H	2	0.72
(1,74)	1:7:A:PRO:HG2	1:10:A:PHE:HD1	18	0.72
(1,74)	1:7:A:PRO:HG2	1:10:A:PHE:HD2	18	0.72
(1,74)	1:7:A:PRO:HG3	1:10:A:PHE:HD1	18	0.72
(1,74)	1:7:A:PRO:HG3	1:10:A:PHE:HD2	18	0.72
(1,68)	1:6:A:VAL:HG11	1:7:A:PRO:HG2	9	0.72
(1,68)	1:6:A:VAL:HG11	1:7:A:PRO:HG3	9	0.72
(1,68)	1:6:A:VAL:HG12	1:7:A:PRO:HG2	9	0.72
(1,68)	1:6:A:VAL:HG12	1:7:A:PRO:HG3	9	0.72
(1,68)	1:6:A:VAL:HG13	1:7:A:PRO:HG2	9	0.72
(1,68)	1:6:A:VAL:HG13	1:7:A:PRO:HG3	9	0.72
(1,1277)	1:27:C:VAL:HB	1:28:C:PRO:HD2	17	0.71
(1,1252)	1:28:C:PRO:HG2	1:31:C:PHE:HE1	19	0.71
(1,1252)	1:28:C:PRO:HG2	1:31:C:PHE:HE2	19	0.71
(1,1252)	1:28:C:PRO:HG3	1:31:C:PHE:HE1	19	0.71
(1,1252)	1:28:C:PRO:HG3	1:31:C:PHE:HE2	19	0.71
(1,1175)	1:24:C:ALA:HB1	2:40:D:PHE:HB3	5	0.71
(1,1175)	1:24:C:ALA:HB2	2:40:D:PHE:HB3	5	0.71
(1,1175)	1:24:C:ALA:HB3	2:40:D:PHE:HB3	5	0.71
(1,1169)	2:36:D:THR:HB	2:38:D:ILE:HD11	20	0.71
(1,1169)	2:36:D:THR:HB	2:38:D:ILE:HD12	20	0.71
(1,1169)	2:36:D:THR:HB	2:38:D:ILE:HD13	20	0.71
(1,1105)	1:24:C:ALA:HB1	1:25:C:GLY:H	16	0.71
(1,1105)	1:24:C:ALA:HB2	1:25:C:GLY:H	16	0.71
(1,1105)	1:24:C:ALA:HB3	1:25:C:GLY:H	16	0.71

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,1051)	2:34:D:CYS:HB2	2:35:D:GLY:H	7	0.71
(1,1036)	2:34:D:CYS:HB2	2:35:D:GLY:HA3	4	0.71
(1,1036)	2:34:D:CYS:HB2	2:35:D:GLY:HA3	18	0.71
(1,1008)	2:18:B:SER:HA	2:38:D:ILE:HG13	13	0.71
(1,1004)	1:32:C:VAL:HA	1:33:C:ARG:HD2	17	0.71
(1,1004)	1:32:C:VAL:HA	1:33:C:ARG:HD3	17	0.71
(1,1003)	1:28:C:PRO:HD2	2:41:D:TYR:HD1	12	0.71
(1,1003)	1:28:C:PRO:HD2	2:41:D:TYR:HD2	12	0.71
(1,958)	1:27:C:VAL:HB	1:28:C:PRO:HD2	17	0.71
(1,933)	1:28:C:PRO:HG2	1:31:C:PHE:HE1	19	0.71
(1,933)	1:28:C:PRO:HG2	1:31:C:PHE:HE2	19	0.71
(1,933)	1:28:C:PRO:HG3	1:31:C:PHE:HE1	19	0.71
(1,933)	1:28:C:PRO:HG3	1:31:C:PHE:HE2	19	0.71
(1,856)	1:24:C:ALA:HB1	2:40:D:PHE:HB3	5	0.71
(1,856)	1:24:C:ALA:HB2	2:40:D:PHE:HB3	5	0.71
(1,856)	1:24:C:ALA:HB3	2:40:D:PHE:HB3	5	0.71
(1,850)	2:36:D:THR:HB	2:38:D:ILE:HD11	20	0.71
(1,850)	2:36:D:THR:HB	2:38:D:ILE:HD12	20	0.71
(1,850)	2:36:D:THR:HB	2:38:D:ILE:HD13	20	0.71
(1,786)	1:24:C:ALA:HB1	1:25:C:GLY:H	16	0.71
(1,786)	1:24:C:ALA:HB2	1:25:C:GLY:H	16	0.71
(1,786)	1:24:C:ALA:HB3	1:25:C:GLY:H	16	0.71
(1,732)	2:34:D:CYS:HB2	2:35:D:GLY:H	7	0.71
(1,717)	2:34:D:CYS:HB2	2:35:D:GLY:HA3	4	0.71
(1,717)	2:34:D:CYS:HB2	2:35:D:GLY:HA3	18	0.71
(1,689)	2:18:B:SER:HA	2:38:D:ILE:HG13	13	0.71
(1,685)	1:32:C:VAL:HA	1:33:C:ARG:HD2	17	0.71
(1,685)	1:32:C:VAL:HA	1:33:C:ARG:HD3	17	0.71
(1,684)	1:28:C:PRO:HD2	2:41:D:TYR:HD1	12	0.71
(1,684)	1:28:C:PRO:HD2	2:41:D:TYR:HD2	12	0.71
(1,520)	2:17:B:ILE:HG13	2:40:D:PHE:HB3	2	0.71
(1,519)	1:3:A:ALA:HB1	2:19:B:PHE:HB3	10	0.71
(1,519)	1:3:A:ALA:HB2	2:19:B:PHE:HB3	10	0.71
(1,519)	1:3:A:ALA:HB3	2:19:B:PHE:HB3	10	0.71
(1,469)	1:10:A:PHE:HD1	1:11:A:VAL:HG11	9	0.71
(1,469)	1:10:A:PHE:HD1	1:11:A:VAL:HG12	9	0.71
(1,469)	1:10:A:PHE:HD1	1:11:A:VAL:HG13	9	0.71
(1,469)	1:10:A:PHE:HD1	1:11:A:VAL:HG21	9	0.71
(1,469)	1:10:A:PHE:HD1	1:11:A:VAL:HG22	9	0.71
(1,469)	1:10:A:PHE:HD1	1:11:A:VAL:HG23	9	0.71
(1,469)	1:10:A:PHE:HD2	1:11:A:VAL:HG11	9	0.71
(1,469)	1:10:A:PHE:HD2	1:11:A:VAL:HG12	9	0.71

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,469)	1:10:A:PHE:HD2	1:11:A:VAL:HG13	9	0.71
(1,469)	1:10:A:PHE:HD2	1:11:A:VAL:HG21	9	0.71
(1,469)	1:10:A:PHE:HD2	1:11:A:VAL:HG22	9	0.71
(1,469)	1:10:A:PHE:HD2	1:11:A:VAL:HG23	9	0.71
(1,463)	1:5:A:HIS:HB2	1:6:A:VAL:HG11	13	0.71
(1,463)	1:5:A:HIS:HB2	1:6:A:VAL:HG12	13	0.71
(1,463)	1:5:A:HIS:HB2	1:6:A:VAL:HG13	13	0.71
(1,449)	1:3:A:ALA:HB1	1:4:A:GLY:H	4	0.71
(1,449)	1:3:A:ALA:HB2	1:4:A:GLY:H	4	0.71
(1,449)	1:3:A:ALA:HB3	1:4:A:GLY:H	4	0.71
(1,403)	1:7:A:PRO:HG2	2:19:B:PHE:HD1	5	0.71
(1,403)	1:7:A:PRO:HG2	2:19:B:PHE:HD2	5	0.71
(1,403)	1:7:A:PRO:HG3	2:19:B:PHE:HD1	5	0.71
(1,403)	1:7:A:PRO:HG3	2:19:B:PHE:HD2	5	0.71
(1,369)	1:6:A:VAL:HG21	2:20:B:TYR:HE1	7	0.71
(1,369)	1:6:A:VAL:HG21	2:20:B:TYR:HE2	7	0.71
(1,369)	1:6:A:VAL:HG22	2:20:B:TYR:HE1	7	0.71
(1,369)	1:6:A:VAL:HG22	2:20:B:TYR:HE2	7	0.71
(1,369)	1:6:A:VAL:HG23	2:20:B:TYR:HE1	7	0.71
(1,369)	1:6:A:VAL:HG23	2:20:B:TYR:HE2	7	0.71
(1,354)	1:10:A:PHE:HD1	2:16:B:PRO:HB3	6	0.71
(1,354)	1:10:A:PHE:HD2	2:16:B:PRO:HB3	6	0.71
(1,201)	2:17:B:ILE:HG13	2:40:D:PHE:HB3	2	0.71
(1,200)	1:3:A:ALA:HB1	2:19:B:PHE:HB3	10	0.71
(1,200)	1:3:A:ALA:HB2	2:19:B:PHE:HB3	10	0.71
(1,200)	1:3:A:ALA:HB3	2:19:B:PHE:HB3	10	0.71
(1,150)	1:10:A:PHE:HD1	1:11:A:VAL:HG11	9	0.71
(1,150)	1:10:A:PHE:HD1	1:11:A:VAL:HG12	9	0.71
(1,150)	1:10:A:PHE:HD1	1:11:A:VAL:HG13	9	0.71
(1,150)	1:10:A:PHE:HD1	1:11:A:VAL:HG21	9	0.71
(1,150)	1:10:A:PHE:HD1	1:11:A:VAL:HG22	9	0.71
(1,150)	1:10:A:PHE:HD1	1:11:A:VAL:HG23	9	0.71
(1,150)	1:10:A:PHE:HD2	1:11:A:VAL:HG11	9	0.71
(1,150)	1:10:A:PHE:HD2	1:11:A:VAL:HG12	9	0.71
(1,150)	1:10:A:PHE:HD2	1:11:A:VAL:HG13	9	0.71
(1,150)	1:10:A:PHE:HD2	1:11:A:VAL:HG21	9	0.71
(1,150)	1:10:A:PHE:HD2	1:11:A:VAL:HG22	9	0.71
(1,150)	1:10:A:PHE:HD2	1:11:A:VAL:HG23	9	0.71
(1,144)	1:5:A:HIS:HB2	1:6:A:VAL:HG11	13	0.71
(1,144)	1:5:A:HIS:HB2	1:6:A:VAL:HG12	13	0.71
(1,144)	1:5:A:HIS:HB2	1:6:A:VAL:HG13	13	0.71
(1,130)	1:3:A:ALA:HB1	1:4:A:GLY:H	4	0.71

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,130)	1:3:A:ALA:HB2	1:4:A:GLY:H	4	0.71
(1,130)	1:3:A:ALA:HB3	1:4:A:GLY:H	4	0.71
(1,84)	1:7:A:PRO:HG2	2:19:B:PHE:HD1	5	0.71
(1,84)	1:7:A:PRO:HG2	2:19:B:PHE:HD2	5	0.71
(1,84)	1:7:A:PRO:HG3	2:19:B:PHE:HD1	5	0.71
(1,84)	1:7:A:PRO:HG3	2:19:B:PHE:HD2	5	0.71
(1,50)	1:6:A:VAL:HG21	2:20:B:TYR:HE1	7	0.71
(1,50)	1:6:A:VAL:HG21	2:20:B:TYR:HE2	7	0.71
(1,50)	1:6:A:VAL:HG22	2:20:B:TYR:HE1	7	0.71
(1,50)	1:6:A:VAL:HG22	2:20:B:TYR:HE2	7	0.71
(1,50)	1:6:A:VAL:HG23	2:20:B:TYR:HE1	7	0.71
(1,50)	1:6:A:VAL:HG23	2:20:B:TYR:HE2	7	0.71
(1,35)	1:10:A:PHE:HD1	2:16:B:PRO:HB3	6	0.71
(1,35)	1:10:A:PHE:HD2	2:16:B:PRO:HB3	6	0.71
(1,1303)	2:41:D:TYR:H	2:41:D:TYR:HD1	1	0.7
(1,1303)	2:41:D:TYR:H	2:41:D:TYR:HD2	1	0.7
(1,1169)	2:36:D:THR:HB	2:38:D:ILE:HD11	15	0.7
(1,1169)	2:36:D:THR:HB	2:38:D:ILE:HD12	15	0.7
(1,1169)	2:36:D:THR:HB	2:38:D:ILE:HD13	15	0.7
(1,1148)	1:28:C:PRO:HB2	1:31:C:PHE:HD1	2	0.7
(1,1148)	1:28:C:PRO:HB2	1:31:C:PHE:HD2	2	0.7
(1,1137)	1:22:C:GLY:H	2:40:D:PHE:H	4	0.7
(1,1122)	1:25:C:GLY:HA2	1:26:C:HIS:HD2	16	0.7
(1,1105)	1:24:C:ALA:HB1	1:25:C:GLY:H	2	0.7
(1,1105)	1:24:C:ALA:HB2	1:25:C:GLY:H	2	0.7
(1,1105)	1:24:C:ALA:HB3	1:25:C:GLY:H	2	0.7
(1,1102)	1:29:C:GLU:HB3	2:41:D:TYR:HD1	19	0.7
(1,1102)	1:29:C:GLU:HB3	2:41:D:TYR:HD2	19	0.7
(1,1094)	1:29:C:GLU:HG2	2:40:D:PHE:HA	14	0.7
(1,1063)	1:22:C:GLY:H	2:39:D:SER:HA	17	0.7
(1,984)	2:41:D:TYR:H	2:41:D:TYR:HD1	1	0.7
(1,984)	2:41:D:TYR:H	2:41:D:TYR:HD2	1	0.7
(1,850)	2:36:D:THR:HB	2:38:D:ILE:HD11	15	0.7
(1,850)	2:36:D:THR:HB	2:38:D:ILE:HD12	15	0.7
(1,850)	2:36:D:THR:HB	2:38:D:ILE:HD13	15	0.7
(1,829)	1:28:C:PRO:HB2	1:31:C:PHE:HD1	2	0.7
(1,829)	1:28:C:PRO:HB2	1:31:C:PHE:HD2	2	0.7
(1,818)	1:22:C:GLY:H	2:40:D:PHE:H	4	0.7
(1,803)	1:25:C:GLY:HA2	1:26:C:HIS:HD2	16	0.7
(1,786)	1:24:C:ALA:HB1	1:25:C:GLY:H	2	0.7
(1,786)	1:24:C:ALA:HB2	1:25:C:GLY:H	2	0.7
(1,786)	1:24:C:ALA:HB3	1:25:C:GLY:H	2	0.7

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,783)	1:29:C:GLU:HB3	2:41:D:TYR:HD1	19	0.7
(1,783)	1:29:C:GLU:HB3	2:41:D:TYR:HD2	19	0.7
(1,775)	1:29:C:GLU:HG2	2:40:D:PHE:HA	14	0.7
(1,744)	1:22:C:GLY:H	2:39:D:SER:HA	17	0.7
(1,461)	1:6:A:VAL:HG11	1:7:A:PRO:HB3	19	0.7
(1,461)	1:6:A:VAL:HG12	1:7:A:PRO:HB3	19	0.7
(1,461)	1:6:A:VAL:HG13	1:7:A:PRO:HB3	19	0.7
(1,445)	1:4:A:GLY:HA2	1:6:A:VAL:H	6	0.7
(1,442)	1:8:A:GLU:HG3	2:19:B:PHE:HA	6	0.7
(1,395)	2:13:B:CYS:HB2	2:14:B:GLY:H	6	0.7
(1,395)	2:13:B:CYS:HB2	2:14:B:GLY:H	11	0.7
(1,142)	1:6:A:VAL:HG11	1:7:A:PRO:HB3	19	0.7
(1,142)	1:6:A:VAL:HG12	1:7:A:PRO:HB3	19	0.7
(1,142)	1:6:A:VAL:HG13	1:7:A:PRO:HB3	19	0.7
(1,126)	1:4:A:GLY:HA2	1:6:A:VAL:H	6	0.7
(1,123)	1:8:A:GLU:HG3	2:19:B:PHE:HA	6	0.7
(1,76)	2:13:B:CYS:HB2	2:14:B:GLY:H	6	0.7
(1,76)	2:13:B:CYS:HB2	2:14:B:GLY:H	11	0.7
(1,1122)	1:25:C:GLY:HA2	1:26:C:HIS:HD2	4	0.69
(1,1058)	1:31:C:PHE:HD1	2:35:D:GLY:HA3	7	0.69
(1,1058)	1:31:C:PHE:HD2	2:35:D:GLY:HA3	7	0.69
(1,1043)	1:27:C:VAL:HG11	1:28:C:PRO:HG2	7	0.69
(1,1043)	1:27:C:VAL:HG11	1:28:C:PRO:HG3	7	0.69
(1,1043)	1:27:C:VAL:HG12	1:28:C:PRO:HG2	7	0.69
(1,1043)	1:27:C:VAL:HG12	1:28:C:PRO:HG3	7	0.69
(1,1043)	1:27:C:VAL:HG13	1:28:C:PRO:HG2	7	0.69
(1,1043)	1:27:C:VAL:HG13	1:28:C:PRO:HG3	7	0.69
(1,803)	1:25:C:GLY:HA2	1:26:C:HIS:HD2	4	0.69
(1,739)	1:31:C:PHE:HD1	2:35:D:GLY:HA3	7	0.69
(1,739)	1:31:C:PHE:HD2	2:35:D:GLY:HA3	7	0.69
(1,724)	1:27:C:VAL:HG11	1:28:C:PRO:HG2	7	0.69
(1,724)	1:27:C:VAL:HG11	1:28:C:PRO:HG3	7	0.69
(1,724)	1:27:C:VAL:HG12	1:28:C:PRO:HG2	7	0.69
(1,724)	1:27:C:VAL:HG12	1:28:C:PRO:HG3	7	0.69
(1,724)	1:27:C:VAL:HG13	1:28:C:PRO:HG2	7	0.69
(1,724)	1:27:C:VAL:HG13	1:28:C:PRO:HG3	7	0.69
(1,672)	2:17:B:ILE:HG13	2:40:D:PHE:HD1	14	0.69
(1,672)	2:17:B:ILE:HG13	2:40:D:PHE:HD2	14	0.69
(1,662)	2:19:B:PHE:HE1	2:38:D:ILE:HG13	3	0.69
(1,662)	2:19:B:PHE:HE2	2:38:D:ILE:HG13	3	0.69
(1,356)	1:8:A:GLU:HG2	2:20:B:TYR:HE1	4	0.69
(1,356)	1:8:A:GLU:HG2	2:20:B:TYR:HE2	4	0.69

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,347)	1:7:A:PRO:HD2	2:20:B:TYR:HD1	18	0.69
(1,347)	1:7:A:PRO:HD2	2:20:B:TYR:HD2	18	0.69
(1,37)	1:8:A:GLU:HG2	2:20:B:TYR:HE1	4	0.69
(1,37)	1:8:A:GLU:HG2	2:20:B:TYR:HE2	4	0.69
(1,28)	1:7:A:PRO:HD2	2:20:B:TYR:HD1	18	0.69
(1,28)	1:7:A:PRO:HD2	2:20:B:TYR:HD2	18	0.69
(1,15)	2:19:B:PHE:HE1	2:38:D:ILE:HG13	3	0.69
(1,15)	2:19:B:PHE:HE2	2:38:D:ILE:HG13	3	0.69
(1,7)	2:17:B:ILE:HG13	2:40:D:PHE:HD1	14	0.69
(1,7)	2:17:B:ILE:HG13	2:40:D:PHE:HD2	14	0.69
(1,1287)	2:38:D:ILE:HA	2:38:D:ILE:HD11	17	0.68
(1,1287)	2:38:D:ILE:HA	2:38:D:ILE:HD12	17	0.68
(1,1287)	2:38:D:ILE:HA	2:38:D:ILE:HD13	17	0.68
(1,1102)	1:29:C:GLU:HB3	2:41:D:TYR:HD1	10	0.68
(1,1102)	1:29:C:GLU:HB3	2:41:D:TYR:HD2	10	0.68
(1,1042)	1:25:C:GLY:H	2:40:D:PHE:HD1	13	0.68
(1,1042)	1:25:C:GLY:H	2:40:D:PHE:HD2	13	0.68
(1,1040)	1:31:C:PHE:HE1	2:37:D:PRO:HD2	10	0.68
(1,1040)	1:31:C:PHE:HE2	2:37:D:PRO:HD2	10	0.68
(1,968)	2:38:D:ILE:HA	2:38:D:ILE:HD11	17	0.68
(1,968)	2:38:D:ILE:HA	2:38:D:ILE:HD12	17	0.68
(1,968)	2:38:D:ILE:HA	2:38:D:ILE:HD13	17	0.68
(1,783)	1:29:C:GLU:HB3	2:41:D:TYR:HD1	10	0.68
(1,783)	1:29:C:GLU:HB3	2:41:D:TYR:HD2	10	0.68
(1,723)	1:25:C:GLY:H	2:40:D:PHE:HD1	13	0.68
(1,723)	1:25:C:GLY:H	2:40:D:PHE:HD2	13	0.68
(1,721)	1:31:C:PHE:HE1	2:37:D:PRO:HD2	10	0.68
(1,721)	1:31:C:PHE:HE2	2:37:D:PRO:HD2	10	0.68
(1,671)	2:17:B:ILE:HG13	2:40:D:PHE:HE1	12	0.68
(1,671)	2:17:B:ILE:HG13	2:40:D:PHE:HE2	12	0.68
(1,629)	2:17:B:ILE:H	2:17:B:ILE:HD11	17	0.68
(1,629)	2:17:B:ILE:H	2:17:B:ILE:HD12	17	0.68
(1,629)	2:17:B:ILE:H	2:17:B:ILE:HD13	17	0.68
(1,483)	1:1:A:GLY:H	2:20:B:TYR:HD1	4	0.68
(1,483)	1:1:A:GLY:H	2:20:B:TYR:HD2	4	0.68
(1,466)	1:4:A:GLY:HA2	1:5:A:HIS:HD2	9	0.68
(1,397)	2:18:B:SER:HB2	2:19:B:PHE:H	20	0.68
(1,387)	1:6:A:VAL:HG11	1:7:A:PRO:HG2	11	0.68
(1,387)	1:6:A:VAL:HG11	1:7:A:PRO:HG3	11	0.68
(1,387)	1:6:A:VAL:HG12	1:7:A:PRO:HG2	11	0.68
(1,387)	1:6:A:VAL:HG12	1:7:A:PRO:HG3	11	0.68
(1,387)	1:6:A:VAL:HG13	1:7:A:PRO:HG2	11	0.68

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,387)	1:6:A:VAL:HG13	1:7:A:PRO:HG3	11	0.68
(1,310)	2:17:B:ILE:H	2:17:B:ILE:HD11	17	0.68
(1,310)	2:17:B:ILE:H	2:17:B:ILE:HD12	17	0.68
(1,310)	2:17:B:ILE:H	2:17:B:ILE:HD13	17	0.68
(1,164)	1:1:A:GLY:H	2:20:B:TYR:HD1	4	0.68
(1,164)	1:1:A:GLY:H	2:20:B:TYR:HD2	4	0.68
(1,147)	1:4:A:GLY:HA2	1:5:A:HIS:HD2	9	0.68
(1,78)	2:18:B:SER:HB2	2:19:B:PHE:H	20	0.68
(1,68)	1:6:A:VAL:HG11	1:7:A:PRO:HG2	11	0.68
(1,68)	1:6:A:VAL:HG11	1:7:A:PRO:HG3	11	0.68
(1,68)	1:6:A:VAL:HG12	1:7:A:PRO:HG2	11	0.68
(1,68)	1:6:A:VAL:HG12	1:7:A:PRO:HG3	11	0.68
(1,68)	1:6:A:VAL:HG13	1:7:A:PRO:HG2	11	0.68
(1,68)	1:6:A:VAL:HG13	1:7:A:PRO:HG3	11	0.68
(1,6)	2:17:B:ILE:HG13	2:40:D:PHE:HE1	12	0.68
(1,6)	2:17:B:ILE:HG13	2:40:D:PHE:HE2	12	0.68
(1,1287)	2:38:D:ILE:HA	2:38:D:ILE:HD11	13	0.67
(1,1287)	2:38:D:ILE:HA	2:38:D:ILE:HD12	13	0.67
(1,1287)	2:38:D:ILE:HA	2:38:D:ILE:HD13	13	0.67
(1,1243)	1:33:C:ARG:H	1:33:C:ARG:HE	1	0.67
(1,1147)	1:28:C:PRO:HB2	1:31:C:PHE:HE1	12	0.67
(1,1147)	1:28:C:PRO:HB2	1:31:C:PHE:HE2	12	0.67
(1,1105)	1:24:C:ALA:HB1	1:25:C:GLY:H	15	0.67
(1,1105)	1:24:C:ALA:HB2	1:25:C:GLY:H	15	0.67
(1,1105)	1:24:C:ALA:HB3	1:25:C:GLY:H	15	0.67
(1,1080)	1:25:C:GLY:H	2:41:D:TYR:H	4	0.67
(1,1045)	1:27:C:VAL:HB	2:41:D:TYR:HD1	9	0.67
(1,1045)	1:27:C:VAL:HB	2:41:D:TYR:HD2	9	0.67
(1,1010)	1:31:C:PHE:HD1	2:37:D:PRO:HB3	18	0.67
(1,1010)	1:31:C:PHE:HD2	2:37:D:PRO:HB3	18	0.67
(1,968)	2:38:D:ILE:HA	2:38:D:ILE:HD11	13	0.67
(1,968)	2:38:D:ILE:HA	2:38:D:ILE:HD12	13	0.67
(1,968)	2:38:D:ILE:HA	2:38:D:ILE:HD13	13	0.67
(1,924)	1:33:C:ARG:H	1:33:C:ARG:HE	1	0.67
(1,828)	1:28:C:PRO:HB2	1:31:C:PHE:HE1	12	0.67
(1,828)	1:28:C:PRO:HB2	1:31:C:PHE:HE2	12	0.67
(1,786)	1:24:C:ALA:HB1	1:25:C:GLY:H	15	0.67
(1,786)	1:24:C:ALA:HB2	1:25:C:GLY:H	15	0.67
(1,786)	1:24:C:ALA:HB3	1:25:C:GLY:H	15	0.67
(1,761)	1:25:C:GLY:H	2:41:D:TYR:H	4	0.67
(1,726)	1:27:C:VAL:HB	2:41:D:TYR:HD1	9	0.67
(1,726)	1:27:C:VAL:HB	2:41:D:TYR:HD2	9	0.67

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,691)	1:31:C:PHE:HD1	2:37:D:PRO:HB3	18	0.67
(1,691)	1:31:C:PHE:HD2	2:37:D:PRO:HB3	18	0.67
(1,665)	2:19:B:PHE:HE1	2:38:D:ILE:HD11	4	0.67
(1,665)	2:19:B:PHE:HE1	2:38:D:ILE:HD12	4	0.67
(1,665)	2:19:B:PHE:HE1	2:38:D:ILE:HD13	4	0.67
(1,665)	2:19:B:PHE:HE2	2:38:D:ILE:HD11	4	0.67
(1,665)	2:19:B:PHE:HE2	2:38:D:ILE:HD12	4	0.67
(1,665)	2:19:B:PHE:HE2	2:38:D:ILE:HD13	4	0.67
(1,575)	1:10:A:PHE:HE1	2:15:B:THR:HA	17	0.67
(1,575)	1:10:A:PHE:HE2	2:15:B:THR:HA	17	0.67
(1,461)	1:6:A:VAL:HG11	1:7:A:PRO:HB3	14	0.67
(1,461)	1:6:A:VAL:HG12	1:7:A:PRO:HB3	14	0.67
(1,461)	1:6:A:VAL:HG13	1:7:A:PRO:HB3	14	0.67
(1,452)	2:15:B:THR:HG21	2:16:B:PRO:HD3	15	0.67
(1,452)	2:15:B:THR:HG22	2:16:B:PRO:HD3	15	0.67
(1,452)	2:15:B:THR:HG23	2:16:B:PRO:HD3	15	0.67
(1,446)	1:8:A:GLU:HB3	2:20:B:TYR:HD1	2	0.67
(1,446)	1:8:A:GLU:HB3	2:20:B:TYR:HD2	2	0.67
(1,446)	1:8:A:GLU:HB3	2:20:B:TYR:HD1	12	0.67
(1,446)	1:8:A:GLU:HB3	2:20:B:TYR:HD2	12	0.67
(1,420)	1:4:A:GLY:HA3	1:5:A:HIS:HD2	6	0.67
(1,405)	1:1:A:GLY:H	2:18:B:SER:HB2	17	0.67
(1,401)	1:10:A:PHE:HD1	1:11:A:VAL:HA	18	0.67
(1,401)	1:10:A:PHE:HD2	1:11:A:VAL:HA	18	0.67
(1,398)	1:7:A:PRO:HA	2:19:B:PHE:HD1	12	0.67
(1,398)	1:7:A:PRO:HA	2:19:B:PHE:HD2	12	0.67
(1,380)	2:13:B:CYS:HB2	2:14:B:GLY:HA3	20	0.67
(1,352)	2:17:B:ILE:HG13	2:39:D:SER:HA	2	0.67
(1,352)	2:17:B:ILE:HG13	2:39:D:SER:HA	9	0.67
(1,256)	1:10:A:PHE:HE1	2:15:B:THR:HA	17	0.67
(1,256)	1:10:A:PHE:HE2	2:15:B:THR:HA	17	0.67
(1,142)	1:6:A:VAL:HG11	1:7:A:PRO:HB3	14	0.67
(1,142)	1:6:A:VAL:HG12	1:7:A:PRO:HB3	14	0.67
(1,142)	1:6:A:VAL:HG13	1:7:A:PRO:HB3	14	0.67
(1,133)	2:15:B:THR:HG21	2:16:B:PRO:HD3	15	0.67
(1,133)	2:15:B:THR:HG22	2:16:B:PRO:HD3	15	0.67
(1,133)	2:15:B:THR:HG23	2:16:B:PRO:HD3	15	0.67
(1,127)	1:8:A:GLU:HB3	2:20:B:TYR:HD1	2	0.67
(1,127)	1:8:A:GLU:HB3	2:20:B:TYR:HD2	2	0.67
(1,127)	1:8:A:GLU:HB3	2:20:B:TYR:HD1	12	0.67
(1,127)	1:8:A:GLU:HB3	2:20:B:TYR:HD2	12	0.67
(1,101)	1:4:A:GLY:HA3	1:5:A:HIS:HD2	6	0.67

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,86)	1:1:A:GLY:H	2:18:B:SER:HB2	17	0.67
(1,82)	1:10:A:PHE:HD1	1:11:A:VAL:HA	18	0.67
(1,82)	1:10:A:PHE:HD2	1:11:A:VAL:HA	18	0.67
(1,79)	1:7:A:PRO:HA	2:19:B:PHE:HD1	12	0.67
(1,79)	1:7:A:PRO:HA	2:19:B:PHE:HD2	12	0.67
(1,61)	2:13:B:CYS:HB2	2:14:B:GLY:HA3	20	0.67
(1,33)	2:17:B:ILE:HG13	2:39:D:SER:HA	2	0.67
(1,33)	2:17:B:ILE:HG13	2:39:D:SER:HA	9	0.67
(1,18)	2:19:B:PHE:HE1	2:38:D:ILE:HD11	4	0.67
(1,18)	2:19:B:PHE:HE1	2:38:D:ILE:HD12	4	0.67
(1,18)	2:19:B:PHE:HE1	2:38:D:ILE:HD13	4	0.67
(1,18)	2:19:B:PHE:HE2	2:38:D:ILE:HD11	4	0.67
(1,18)	2:19:B:PHE:HE2	2:38:D:ILE:HD12	4	0.67
(1,18)	2:19:B:PHE:HE2	2:38:D:ILE:HD13	4	0.67
(1,1287)	2:38:D:ILE:HA	2:38:D:ILE:HD11	1	0.66
(1,1287)	2:38:D:ILE:HA	2:38:D:ILE:HD12	1	0.66
(1,1287)	2:38:D:ILE:HA	2:38:D:ILE:HD13	1	0.66
(1,1094)	1:29:C:GLU:HG2	2:40:D:PHE:HA	17	0.66
(1,1051)	2:34:D:CYS:HB2	2:35:D:GLY:H	6	0.66
(1,1025)	1:27:C:VAL:HG21	2:41:D:TYR:HE1	5	0.66
(1,1025)	1:27:C:VAL:HG21	2:41:D:TYR:HE2	5	0.66
(1,1025)	1:27:C:VAL:HG22	2:41:D:TYR:HE1	5	0.66
(1,1025)	1:27:C:VAL:HG22	2:41:D:TYR:HE2	5	0.66
(1,1025)	1:27:C:VAL:HG23	2:41:D:TYR:HE1	5	0.66
(1,1025)	1:27:C:VAL:HG23	2:41:D:TYR:HE2	5	0.66
(1,968)	2:38:D:ILE:HA	2:38:D:ILE:HD11	1	0.66
(1,968)	2:38:D:ILE:HA	2:38:D:ILE:HD12	1	0.66
(1,968)	2:38:D:ILE:HA	2:38:D:ILE:HD13	1	0.66
(1,775)	1:29:C:GLU:HG2	2:40:D:PHE:HA	17	0.66
(1,732)	2:34:D:CYS:HB2	2:35:D:GLY:H	6	0.66
(1,706)	1:27:C:VAL:HG21	2:41:D:TYR:HE1	5	0.66
(1,706)	1:27:C:VAL:HG21	2:41:D:TYR:HE2	5	0.66
(1,706)	1:27:C:VAL:HG22	2:41:D:TYR:HE1	5	0.66
(1,706)	1:27:C:VAL:HG22	2:41:D:TYR:HE2	5	0.66
(1,706)	1:27:C:VAL:HG23	2:41:D:TYR:HE1	5	0.66
(1,706)	1:27:C:VAL:HG23	2:41:D:TYR:HE2	5	0.66
(1,395)	2:13:B:CYS:HB2	2:14:B:GLY:H	1	0.66
(1,369)	1:6:A:VAL:HG21	2:20:B:TYR:HE1	1	0.66
(1,369)	1:6:A:VAL:HG21	2:20:B:TYR:HE2	1	0.66
(1,369)	1:6:A:VAL:HG22	2:20:B:TYR:HE1	1	0.66
(1,369)	1:6:A:VAL:HG22	2:20:B:TYR:HE2	1	0.66
(1,369)	1:6:A:VAL:HG23	2:20:B:TYR:HE1	1	0.66

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,369)	1:6:A:VAL:HG23	2:20:B:TYR:HE2	1	0.66
(1,76)	2:13:B:CYS:HB2	2:14:B:GLY:H	1	0.66
(1,50)	1:6:A:VAL:HG21	2:20:B:TYR:HE1	1	0.66
(1,50)	1:6:A:VAL:HG21	2:20:B:TYR:HE2	1	0.66
(1,50)	1:6:A:VAL:HG22	2:20:B:TYR:HE1	1	0.66
(1,50)	1:6:A:VAL:HG22	2:20:B:TYR:HE2	1	0.66
(1,50)	1:6:A:VAL:HG23	2:20:B:TYR:HE1	1	0.66
(1,50)	1:6:A:VAL:HG23	2:20:B:TYR:HE2	1	0.66
(1,1175)	1:24:C:ALA:HB1	2:40:D:PHE:HB3	19	0.65
(1,1175)	1:24:C:ALA:HB2	2:40:D:PHE:HB3	19	0.65
(1,1175)	1:24:C:ALA:HB3	2:40:D:PHE:HB3	19	0.65
(1,1109)	1:31:C:PHE:HD1	2:36:D:THR:HG21	10	0.65
(1,1109)	1:31:C:PHE:HD1	2:36:D:THR:HG22	10	0.65
(1,1109)	1:31:C:PHE:HD1	2:36:D:THR:HG23	10	0.65
(1,1109)	1:31:C:PHE:HD2	2:36:D:THR:HG21	10	0.65
(1,1109)	1:31:C:PHE:HD2	2:36:D:THR:HG22	10	0.65
(1,1109)	1:31:C:PHE:HD2	2:36:D:THR:HG23	10	0.65
(1,1105)	1:24:C:ALA:HB1	1:25:C:GLY:H	14	0.65
(1,1105)	1:24:C:ALA:HB2	1:25:C:GLY:H	14	0.65
(1,1105)	1:24:C:ALA:HB3	1:25:C:GLY:H	14	0.65
(1,1094)	1:29:C:GLU:HG2	2:40:D:PHE:HA	13	0.65
(1,1043)	1:27:C:VAL:HG11	1:28:C:PRO:HG2	9	0.65
(1,1043)	1:27:C:VAL:HG11	1:28:C:PRO:HG3	9	0.65
(1,1043)	1:27:C:VAL:HG12	1:28:C:PRO:HG2	9	0.65
(1,1043)	1:27:C:VAL:HG12	1:28:C:PRO:HG3	9	0.65
(1,1043)	1:27:C:VAL:HG13	1:28:C:PRO:HG2	9	0.65
(1,1043)	1:27:C:VAL:HG13	1:28:C:PRO:HG3	9	0.65
(1,856)	1:24:C:ALA:HB1	2:40:D:PHE:HB3	19	0.65
(1,856)	1:24:C:ALA:HB2	2:40:D:PHE:HB3	19	0.65
(1,856)	1:24:C:ALA:HB3	2:40:D:PHE:HB3	19	0.65
(1,790)	1:31:C:PHE:HD1	2:36:D:THR:HG21	10	0.65
(1,790)	1:31:C:PHE:HD1	2:36:D:THR:HG22	10	0.65
(1,790)	1:31:C:PHE:HD1	2:36:D:THR:HG23	10	0.65
(1,790)	1:31:C:PHE:HD2	2:36:D:THR:HG21	10	0.65
(1,790)	1:31:C:PHE:HD2	2:36:D:THR:HG22	10	0.65
(1,790)	1:31:C:PHE:HD2	2:36:D:THR:HG23	10	0.65
(1,786)	1:24:C:ALA:HB1	1:25:C:GLY:H	14	0.65
(1,786)	1:24:C:ALA:HB2	1:25:C:GLY:H	14	0.65
(1,786)	1:24:C:ALA:HB3	1:25:C:GLY:H	14	0.65
(1,775)	1:29:C:GLU:HG2	2:40:D:PHE:HA	13	0.65
(1,724)	1:27:C:VAL:HG11	1:28:C:PRO:HG2	9	0.65
(1,724)	1:27:C:VAL:HG11	1:28:C:PRO:HG3	9	0.65

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,724)	1:27:C:VAL:HG12	1:28:C:PRO:HG2	9	0.65
(1,724)	1:27:C:VAL:HG12	1:28:C:PRO:HG3	9	0.65
(1,724)	1:27:C:VAL:HG13	1:28:C:PRO:HG2	9	0.65
(1,724)	1:27:C:VAL:HG13	1:28:C:PRO:HG3	9	0.65
(1,453)	1:10:A:PHE:HD1	2:15:B:THR:HG21	7	0.65
(1,453)	1:10:A:PHE:HD1	2:15:B:THR:HG22	7	0.65
(1,453)	1:10:A:PHE:HD1	2:15:B:THR:HG23	7	0.65
(1,453)	1:10:A:PHE:HD2	2:15:B:THR:HG21	7	0.65
(1,453)	1:10:A:PHE:HD2	2:15:B:THR:HG22	7	0.65
(1,453)	1:10:A:PHE:HD2	2:15:B:THR:HG23	7	0.65
(1,436)	1:8:A:GLU:HG2	2:20:B:TYR:HD1	1	0.65
(1,436)	1:8:A:GLU:HG2	2:20:B:TYR:HD2	1	0.65
(1,398)	1:7:A:PRO:HA	2:19:B:PHE:HD1	6	0.65
(1,398)	1:7:A:PRO:HA	2:19:B:PHE:HD2	6	0.65
(1,353)	1:6:A:VAL:HB	2:20:B:TYR:HB2	14	0.65
(1,353)	1:6:A:VAL:HB	2:20:B:TYR:HB3	14	0.65
(1,134)	1:10:A:PHE:HD1	2:15:B:THR:HG21	7	0.65
(1,134)	1:10:A:PHE:HD1	2:15:B:THR:HG22	7	0.65
(1,134)	1:10:A:PHE:HD1	2:15:B:THR:HG23	7	0.65
(1,134)	1:10:A:PHE:HD2	2:15:B:THR:HG21	7	0.65
(1,134)	1:10:A:PHE:HD2	2:15:B:THR:HG22	7	0.65
(1,134)	1:10:A:PHE:HD2	2:15:B:THR:HG23	7	0.65
(1,117)	1:8:A:GLU:HG2	2:20:B:TYR:HD1	1	0.65
(1,117)	1:8:A:GLU:HG2	2:20:B:TYR:HD2	1	0.65
(1,79)	1:7:A:PRO:HA	2:19:B:PHE:HD1	6	0.65
(1,79)	1:7:A:PRO:HA	2:19:B:PHE:HD2	6	0.65
(1,34)	1:6:A:VAL:HB	2:20:B:TYR:HB2	14	0.65
(1,34)	1:6:A:VAL:HB	2:20:B:TYR:HB3	14	0.65
(1,1309)	1:24:C:ALA:HA	2:40:D:PHE:HD1	14	0.64
(1,1309)	1:24:C:ALA:HA	2:40:D:PHE:HD2	14	0.64
(1,1249)	1:33:C:ARG:HB2	1:33:C:ARG:HE	3	0.64
(1,1105)	1:24:C:ALA:HB1	1:25:C:GLY:H	19	0.64
(1,1105)	1:24:C:ALA:HB2	1:25:C:GLY:H	19	0.64
(1,1105)	1:24:C:ALA:HB3	1:25:C:GLY:H	19	0.64
(1,1097)	1:29:C:GLU:HG3	2:41:D:TYR:HB2	9	0.64
(1,1097)	1:29:C:GLU:HG3	2:41:D:TYR:HB3	9	0.64
(1,1047)	1:24:C:ALA:HA	2:40:D:PHE:HD1	14	0.64
(1,1047)	1:24:C:ALA:HA	2:40:D:PHE:HD2	14	0.64
(1,1042)	1:25:C:GLY:H	2:40:D:PHE:HD1	7	0.64
(1,1042)	1:25:C:GLY:H	2:40:D:PHE:HD2	7	0.64
(1,1036)	2:34:D:CYS:HB2	2:35:D:GLY:HA3	13	0.64
(1,1036)	2:34:D:CYS:HB2	2:35:D:GLY:HA3	16	0.64

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,998)	1:22:C:GLY:HA3	2:39:D:SER:HB2	9	0.64
(1,995)	1:22:C:GLY:HA3	2:39:D:SER:HB2	9	0.64
(1,990)	1:24:C:ALA:HA	2:40:D:PHE:HD1	14	0.64
(1,990)	1:24:C:ALA:HA	2:40:D:PHE:HD2	14	0.64
(1,930)	1:33:C:ARG:HB2	1:33:C:ARG:HE	3	0.64
(1,786)	1:24:C:ALA:HB1	1:25:C:GLY:H	19	0.64
(1,786)	1:24:C:ALA:HB2	1:25:C:GLY:H	19	0.64
(1,786)	1:24:C:ALA:HB3	1:25:C:GLY:H	19	0.64
(1,778)	1:29:C:GLU:HG3	2:41:D:TYR:HB2	9	0.64
(1,778)	1:29:C:GLU:HG3	2:41:D:TYR:HB3	9	0.64
(1,728)	1:24:C:ALA:HA	2:40:D:PHE:HD1	14	0.64
(1,728)	1:24:C:ALA:HA	2:40:D:PHE:HD2	14	0.64
(1,723)	1:25:C:GLY:H	2:40:D:PHE:HD1	7	0.64
(1,723)	1:25:C:GLY:H	2:40:D:PHE:HD2	7	0.64
(1,717)	2:34:D:CYS:HB2	2:35:D:GLY:HA3	13	0.64
(1,717)	2:34:D:CYS:HB2	2:35:D:GLY:HA3	16	0.64
(1,679)	1:22:C:GLY:HA3	2:39:D:SER:HB2	9	0.64
(1,676)	1:22:C:GLY:HA3	2:39:D:SER:HB2	9	0.64
(1,499)	1:6:A:VAL:HG11	1:7:A:PRO:HD3	19	0.64
(1,499)	1:6:A:VAL:HG12	1:7:A:PRO:HD3	19	0.64
(1,499)	1:6:A:VAL:HG13	1:7:A:PRO:HD3	19	0.64
(1,445)	1:4:A:GLY:HA2	1:6:A:VAL:H	8	0.64
(1,405)	1:1:A:GLY:H	2:18:B:SER:HB2	20	0.64
(1,380)	2:13:B:CYS:HB2	2:14:B:GLY:HA3	8	0.64
(1,362)	1:7:A:PRO:HG2	2:19:B:PHE:HD1	13	0.64
(1,362)	1:7:A:PRO:HG2	2:19:B:PHE:HD2	13	0.64
(1,362)	1:7:A:PRO:HG3	2:19:B:PHE:HD1	13	0.64
(1,362)	1:7:A:PRO:HG3	2:19:B:PHE:HD2	13	0.64
(1,348)	1:11:A:VAL:HA	1:12:A:ARG:HD2	12	0.64
(1,348)	1:11:A:VAL:HA	1:12:A:ARG:HD3	12	0.64
(1,180)	1:6:A:VAL:HG11	1:7:A:PRO:HD3	19	0.64
(1,180)	1:6:A:VAL:HG12	1:7:A:PRO:HD3	19	0.64
(1,180)	1:6:A:VAL:HG13	1:7:A:PRO:HD3	19	0.64
(1,126)	1:4:A:GLY:HA2	1:6:A:VAL:H	8	0.64
(1,86)	1:1:A:GLY:H	2:18:B:SER:HB2	20	0.64
(1,61)	2:13:B:CYS:HB2	2:14:B:GLY:HA3	8	0.64
(1,43)	1:7:A:PRO:HG2	2:19:B:PHE:HD1	13	0.64
(1,43)	1:7:A:PRO:HG2	2:19:B:PHE:HD2	13	0.64
(1,43)	1:7:A:PRO:HG3	2:19:B:PHE:HD1	13	0.64
(1,43)	1:7:A:PRO:HG3	2:19:B:PHE:HD2	13	0.64
(1,29)	1:11:A:VAL:HA	1:12:A:ARG:HD2	12	0.64
(1,29)	1:11:A:VAL:HA	1:12:A:ARG:HD3	12	0.64

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,1147)	1:28:C:PRO:HB2	1:31:C:PHE:HE1	7	0.63
(1,1147)	1:28:C:PRO:HB2	1:31:C:PHE:HE2	7	0.63
(1,1117)	1:27:C:VAL:HG11	1:28:C:PRO:HB3	2	0.63
(1,1117)	1:27:C:VAL:HG12	1:28:C:PRO:HB3	2	0.63
(1,1117)	1:27:C:VAL:HG13	1:28:C:PRO:HB3	2	0.63
(1,1109)	1:31:C:PHE:HD1	2:36:D:THR:HG21	3	0.63
(1,1109)	1:31:C:PHE:HD1	2:36:D:THR:HG22	3	0.63
(1,1109)	1:31:C:PHE:HD1	2:36:D:THR:HG23	3	0.63
(1,1109)	1:31:C:PHE:HD2	2:36:D:THR:HG21	3	0.63
(1,1109)	1:31:C:PHE:HD2	2:36:D:THR:HG22	3	0.63
(1,1109)	1:31:C:PHE:HD2	2:36:D:THR:HG23	3	0.63
(1,1078)	1:25:C:GLY:H	2:41:D:TYR:H	11	0.63
(1,1058)	1:31:C:PHE:HD1	2:35:D:GLY:HA3	8	0.63
(1,1058)	1:31:C:PHE:HD2	2:35:D:GLY:HA3	8	0.63
(1,1035)	1:22:C:GLY:H	2:41:D:TYR:HE1	8	0.63
(1,1035)	1:22:C:GLY:H	2:41:D:TYR:HE2	8	0.63
(1,1015)	1:30:C:TYR:HD1	2:39:D:SER:HA	14	0.63
(1,1015)	1:30:C:TYR:HD2	2:39:D:SER:HA	14	0.63
(1,828)	1:28:C:PRO:HB2	1:31:C:PHE:HE1	7	0.63
(1,828)	1:28:C:PRO:HB2	1:31:C:PHE:HE2	7	0.63
(1,798)	1:27:C:VAL:HG11	1:28:C:PRO:HB3	2	0.63
(1,798)	1:27:C:VAL:HG12	1:28:C:PRO:HB3	2	0.63
(1,798)	1:27:C:VAL:HG13	1:28:C:PRO:HB3	2	0.63
(1,790)	1:31:C:PHE:HD1	2:36:D:THR:HG21	3	0.63
(1,790)	1:31:C:PHE:HD1	2:36:D:THR:HG22	3	0.63
(1,790)	1:31:C:PHE:HD1	2:36:D:THR:HG23	3	0.63
(1,790)	1:31:C:PHE:HD2	2:36:D:THR:HG21	3	0.63
(1,790)	1:31:C:PHE:HD2	2:36:D:THR:HG22	3	0.63
(1,790)	1:31:C:PHE:HD2	2:36:D:THR:HG23	3	0.63
(1,759)	1:25:C:GLY:H	2:41:D:TYR:H	11	0.63
(1,739)	1:31:C:PHE:HD1	2:35:D:GLY:HA3	8	0.63
(1,739)	1:31:C:PHE:HD2	2:35:D:GLY:HA3	8	0.63
(1,716)	1:22:C:GLY:H	2:41:D:TYR:HE1	8	0.63
(1,716)	1:22:C:GLY:H	2:41:D:TYR:HE2	8	0.63
(1,696)	1:30:C:TYR:HD1	2:39:D:SER:HA	14	0.63
(1,696)	1:30:C:TYR:HD2	2:39:D:SER:HA	14	0.63
(1,519)	1:3:A:ALA:HB1	2:19:B:PHE:HB3	7	0.63
(1,519)	1:3:A:ALA:HB2	2:19:B:PHE:HB3	7	0.63
(1,519)	1:3:A:ALA:HB3	2:19:B:PHE:HB3	7	0.63
(1,489)	1:4:A:GLY:H	2:19:B:PHE:HE1	2	0.63
(1,489)	1:4:A:GLY:H	2:19:B:PHE:HE2	2	0.63
(1,461)	1:6:A:VAL:HG11	1:7:A:PRO:HB3	8	0.63

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,461)	1:6:A:VAL:HG12	1:7:A:PRO:HB3	8	0.63
(1,461)	1:6:A:VAL:HG13	1:7:A:PRO:HB3	8	0.63
(1,449)	1:3:A:ALA:HB1	1:4:A:GLY:H	11	0.63
(1,449)	1:3:A:ALA:HB2	1:4:A:GLY:H	11	0.63
(1,449)	1:3:A:ALA:HB3	1:4:A:GLY:H	11	0.63
(1,446)	1:8:A:GLU:HB3	2:20:B:TYR:HD1	15	0.63
(1,446)	1:8:A:GLU:HB3	2:20:B:TYR:HD2	15	0.63
(1,407)	1:1:A:GLY:H	2:18:B:SER:HA	14	0.63
(1,200)	1:3:A:ALA:HB1	2:19:B:PHE:HB3	7	0.63
(1,200)	1:3:A:ALA:HB2	2:19:B:PHE:HB3	7	0.63
(1,200)	1:3:A:ALA:HB3	2:19:B:PHE:HB3	7	0.63
(1,170)	1:4:A:GLY:H	2:19:B:PHE:HE1	2	0.63
(1,170)	1:4:A:GLY:H	2:19:B:PHE:HE2	2	0.63
(1,142)	1:6:A:VAL:HG11	1:7:A:PRO:HB3	8	0.63
(1,142)	1:6:A:VAL:HG12	1:7:A:PRO:HB3	8	0.63
(1,142)	1:6:A:VAL:HG13	1:7:A:PRO:HB3	8	0.63
(1,130)	1:3:A:ALA:HB1	1:4:A:GLY:H	11	0.63
(1,130)	1:3:A:ALA:HB2	1:4:A:GLY:H	11	0.63
(1,130)	1:3:A:ALA:HB3	1:4:A:GLY:H	11	0.63
(1,127)	1:8:A:GLU:HB3	2:20:B:TYR:HD1	15	0.63
(1,127)	1:8:A:GLU:HB3	2:20:B:TYR:HD2	15	0.63
(1,88)	1:1:A:GLY:H	2:18:B:SER:HA	14	0.63
(1,1175)	1:24:C:ALA:HB1	2:40:D:PHE:HB3	12	0.62
(1,1175)	1:24:C:ALA:HB2	2:40:D:PHE:HB3	12	0.62
(1,1175)	1:24:C:ALA:HB3	2:40:D:PHE:HB3	12	0.62
(1,1130)	1:27:C:VAL:H	2:40:D:PHE:HB2	18	0.62
(1,1094)	1:29:C:GLU:HG2	2:40:D:PHE:HA	8	0.62
(1,1067)	1:26:C:HIS:HB3	1:27:C:VAL:HG21	11	0.62
(1,1067)	1:26:C:HIS:HB3	1:27:C:VAL:HG22	11	0.62
(1,1067)	1:26:C:HIS:HB3	1:27:C:VAL:HG23	11	0.62
(1,1046)	1:27:C:VAL:HB	2:41:D:TYR:HE1	19	0.62
(1,1046)	1:27:C:VAL:HB	2:41:D:TYR:HE2	19	0.62
(1,1043)	1:27:C:VAL:HG11	1:28:C:PRO:HG2	10	0.62
(1,1043)	1:27:C:VAL:HG11	1:28:C:PRO:HG3	10	0.62
(1,1043)	1:27:C:VAL:HG12	1:28:C:PRO:HG2	10	0.62
(1,1043)	1:27:C:VAL:HG12	1:28:C:PRO:HG3	10	0.62
(1,1043)	1:27:C:VAL:HG13	1:28:C:PRO:HG2	10	0.62
(1,1043)	1:27:C:VAL:HG13	1:28:C:PRO:HG3	10	0.62
(1,1034)	2:38:D:ILE:H	2:38:D:ILE:HG13	4	0.62
(1,1000)	1:31:C:PHE:HE1	2:37:D:PRO:HD2	9	0.62
(1,1000)	1:31:C:PHE:HE2	2:37:D:PRO:HD2	9	0.62
(1,856)	1:24:C:ALA:HB1	2:40:D:PHE:HB3	12	0.62

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,856)	1:24:C:ALA:HB2	2:40:D:PHE:HB3	12	0.62
(1,856)	1:24:C:ALA:HB3	2:40:D:PHE:HB3	12	0.62
(1,811)	1:27:C:VAL:H	2:40:D:PHE:HB2	18	0.62
(1,775)	1:29:C:GLU:HG2	2:40:D:PHE:HA	8	0.62
(1,748)	1:26:C:HIS:HB3	1:27:C:VAL:HG21	11	0.62
(1,748)	1:26:C:HIS:HB3	1:27:C:VAL:HG22	11	0.62
(1,748)	1:26:C:HIS:HB3	1:27:C:VAL:HG23	11	0.62
(1,727)	1:27:C:VAL:HB	2:41:D:TYR:HE1	19	0.62
(1,727)	1:27:C:VAL:HB	2:41:D:TYR:HE2	19	0.62
(1,724)	1:27:C:VAL:HG11	1:28:C:PRO:HG2	10	0.62
(1,724)	1:27:C:VAL:HG11	1:28:C:PRO:HG3	10	0.62
(1,724)	1:27:C:VAL:HG12	1:28:C:PRO:HG2	10	0.62
(1,724)	1:27:C:VAL:HG12	1:28:C:PRO:HG3	10	0.62
(1,724)	1:27:C:VAL:HG13	1:28:C:PRO:HG2	10	0.62
(1,724)	1:27:C:VAL:HG13	1:28:C:PRO:HG3	10	0.62
(1,715)	2:38:D:ILE:H	2:38:D:ILE:HG13	4	0.62
(1,681)	1:31:C:PHE:HE1	2:37:D:PRO:HD2	9	0.62
(1,681)	1:31:C:PHE:HE2	2:37:D:PRO:HD2	9	0.62
(1,673)	2:17:B:ILE:HG13	2:40:D:PHE:HB2	12	0.62
(1,483)	1:1:A:GLY:H	2:20:B:TYR:HD1	16	0.62
(1,483)	1:1:A:GLY:H	2:20:B:TYR:HD2	16	0.62
(1,405)	1:1:A:GLY:H	2:18:B:SER:HB2	19	0.62
(1,380)	2:13:B:CYS:HB2	2:14:B:GLY:HA3	12	0.62
(1,369)	1:6:A:VAL:HG21	2:20:B:TYR:HE1	15	0.62
(1,369)	1:6:A:VAL:HG21	2:20:B:TYR:HE2	15	0.62
(1,369)	1:6:A:VAL:HG22	2:20:B:TYR:HE1	15	0.62
(1,369)	1:6:A:VAL:HG22	2:20:B:TYR:HE2	15	0.62
(1,369)	1:6:A:VAL:HG23	2:20:B:TYR:HE1	15	0.62
(1,369)	1:6:A:VAL:HG23	2:20:B:TYR:HE2	15	0.62
(1,164)	1:1:A:GLY:H	2:20:B:TYR:HD1	16	0.62
(1,164)	1:1:A:GLY:H	2:20:B:TYR:HD2	16	0.62
(1,86)	1:1:A:GLY:H	2:18:B:SER:HB2	19	0.62
(1,61)	2:13:B:CYS:HB2	2:14:B:GLY:HA3	12	0.62
(1,50)	1:6:A:VAL:HG21	2:20:B:TYR:HE1	15	0.62
(1,50)	1:6:A:VAL:HG21	2:20:B:TYR:HE2	15	0.62
(1,50)	1:6:A:VAL:HG22	2:20:B:TYR:HE1	15	0.62
(1,50)	1:6:A:VAL:HG22	2:20:B:TYR:HE2	15	0.62
(1,50)	1:6:A:VAL:HG23	2:20:B:TYR:HE1	15	0.62
(1,50)	1:6:A:VAL:HG23	2:20:B:TYR:HE2	15	0.62
(1,8)	2:17:B:ILE:HG13	2:40:D:PHE:HB2	12	0.62
(1,1159)	1:28:C:PRO:HB3	1:31:C:PHE:HE1	7	0.61
(1,1159)	1:28:C:PRO:HB3	1:31:C:PHE:HE2	7	0.61

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,1094)	1:29:C:GLU:HG2	2:40:D:PHE:HA	2	0.61
(1,1094)	1:29:C:GLU:HG2	2:40:D:PHE:HA	11	0.61
(1,1036)	2:34:D:CYS:HB2	2:35:D:GLY:HA3	9	0.61
(1,1000)	1:31:C:PHE:HE1	2:37:D:PRO:HD2	19	0.61
(1,1000)	1:31:C:PHE:HE2	2:37:D:PRO:HD2	19	0.61
(1,840)	1:28:C:PRO:HB3	1:31:C:PHE:HE1	7	0.61
(1,840)	1:28:C:PRO:HB3	1:31:C:PHE:HE2	7	0.61
(1,775)	1:29:C:GLU:HG2	2:40:D:PHE:HA	2	0.61
(1,775)	1:29:C:GLU:HG2	2:40:D:PHE:HA	11	0.61
(1,717)	2:34:D:CYS:HB2	2:35:D:GLY:HA3	9	0.61
(1,681)	1:31:C:PHE:HE1	2:37:D:PRO:HD2	19	0.61
(1,681)	1:31:C:PHE:HE2	2:37:D:PRO:HD2	19	0.61
(1,491)	1:7:A:PRO:HB2	1:10:A:PHE:HE1	2	0.61
(1,491)	1:7:A:PRO:HB2	1:10:A:PHE:HE2	2	0.61
(1,411)	1:5:A:HIS:HB3	1:6:A:VAL:HG21	17	0.61
(1,411)	1:5:A:HIS:HB3	1:6:A:VAL:HG22	17	0.61
(1,411)	1:5:A:HIS:HB3	1:6:A:VAL:HG23	17	0.61
(1,387)	1:6:A:VAL:HG11	1:7:A:PRO:HG2	1	0.61
(1,387)	1:6:A:VAL:HG11	1:7:A:PRO:HG3	1	0.61
(1,387)	1:6:A:VAL:HG12	1:7:A:PRO:HG2	1	0.61
(1,387)	1:6:A:VAL:HG12	1:7:A:PRO:HG3	1	0.61
(1,387)	1:6:A:VAL:HG13	1:7:A:PRO:HG2	1	0.61
(1,387)	1:6:A:VAL:HG13	1:7:A:PRO:HG3	1	0.61
(1,386)	1:4:A:GLY:H	2:19:B:PHE:HD1	12	0.61
(1,386)	1:4:A:GLY:H	2:19:B:PHE:HD2	12	0.61
(1,382)	1:10:A:PHE:HD1	2:16:B:PRO:HD2	8	0.61
(1,382)	1:10:A:PHE:HD2	2:16:B:PRO:HD2	8	0.61
(1,380)	2:13:B:CYS:HB2	2:14:B:GLY:HA3	11	0.61
(1,172)	1:7:A:PRO:HB2	1:10:A:PHE:HE1	2	0.61
(1,172)	1:7:A:PRO:HB2	1:10:A:PHE:HE2	2	0.61
(1,92)	1:5:A:HIS:HB3	1:6:A:VAL:HG21	17	0.61
(1,92)	1:5:A:HIS:HB3	1:6:A:VAL:HG22	17	0.61
(1,92)	1:5:A:HIS:HB3	1:6:A:VAL:HG23	17	0.61
(1,68)	1:6:A:VAL:HG11	1:7:A:PRO:HG2	1	0.61
(1,68)	1:6:A:VAL:HG11	1:7:A:PRO:HG3	1	0.61
(1,68)	1:6:A:VAL:HG12	1:7:A:PRO:HG2	1	0.61
(1,68)	1:6:A:VAL:HG12	1:7:A:PRO:HG3	1	0.61
(1,68)	1:6:A:VAL:HG13	1:7:A:PRO:HG2	1	0.61
(1,68)	1:6:A:VAL:HG13	1:7:A:PRO:HG3	1	0.61
(1,67)	1:4:A:GLY:H	2:19:B:PHE:HD1	12	0.61
(1,67)	1:4:A:GLY:H	2:19:B:PHE:HD2	12	0.61
(1,63)	1:10:A:PHE:HD1	2:16:B:PRO:HD2	8	0.61

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,63)	1:10:A:PHE:HD2	2:16:B:PRO:HD2	8	0.61
(1,61)	2:13:B:CYS:HB2	2:14:B:GLY:HA3	11	0.61
(1,1061)	1:22:C:GLY:H	2:39:D:SER:HB2	2	0.6
(1,1053)	2:39:D:SER:HB2	2:40:D:PHE:H	6	0.6
(1,1053)	2:39:D:SER:HB2	2:40:D:PHE:H	20	0.6
(1,1051)	2:34:D:CYS:HB2	2:35:D:GLY:H	12	0.6
(1,1042)	1:25:C:GLY:H	2:40:D:PHE:HD1	6	0.6
(1,1042)	1:25:C:GLY:H	2:40:D:PHE:HD2	6	0.6
(1,1009)	1:27:C:VAL:HB	2:41:D:TYR:HB2	16	0.6
(1,1009)	1:27:C:VAL:HB	2:41:D:TYR:HB3	16	0.6
(1,1003)	1:28:C:PRO:HD2	2:41:D:TYR:HD1	17	0.6
(1,1003)	1:28:C:PRO:HD2	2:41:D:TYR:HD2	17	0.6
(1,742)	1:22:C:GLY:H	2:39:D:SER:HB2	2	0.6
(1,734)	2:39:D:SER:HB2	2:40:D:PHE:H	6	0.6
(1,734)	2:39:D:SER:HB2	2:40:D:PHE:H	20	0.6
(1,732)	2:34:D:CYS:HB2	2:35:D:GLY:H	12	0.6
(1,723)	1:25:C:GLY:H	2:40:D:PHE:HD1	6	0.6
(1,723)	1:25:C:GLY:H	2:40:D:PHE:HD2	6	0.6
(1,690)	1:27:C:VAL:HB	2:41:D:TYR:HB2	16	0.6
(1,690)	1:27:C:VAL:HB	2:41:D:TYR:HB3	16	0.6
(1,684)	1:28:C:PRO:HD2	2:41:D:TYR:HD1	17	0.6
(1,684)	1:28:C:PRO:HD2	2:41:D:TYR:HD2	17	0.6
(1,510)	1:3:A:ALA:H	2:19:B:PHE:HD1	17	0.6
(1,510)	1:3:A:ALA:H	2:19:B:PHE:HD2	17	0.6
(1,501)	1:4:A:GLY:H	1:6:A:VAL:H	3	0.6
(1,453)	1:10:A:PHE:HD1	2:15:B:THR:HG21	2	0.6
(1,453)	1:10:A:PHE:HD1	2:15:B:THR:HG22	2	0.6
(1,453)	1:10:A:PHE:HD1	2:15:B:THR:HG23	2	0.6
(1,453)	1:10:A:PHE:HD2	2:15:B:THR:HG21	2	0.6
(1,453)	1:10:A:PHE:HD2	2:15:B:THR:HG22	2	0.6
(1,453)	1:10:A:PHE:HD2	2:15:B:THR:HG23	2	0.6
(1,453)	1:10:A:PHE:HD1	2:15:B:THR:HG21	17	0.6
(1,453)	1:10:A:PHE:HD1	2:15:B:THR:HG22	17	0.6
(1,453)	1:10:A:PHE:HD1	2:15:B:THR:HG23	17	0.6
(1,453)	1:10:A:PHE:HD2	2:15:B:THR:HG21	17	0.6
(1,453)	1:10:A:PHE:HD2	2:15:B:THR:HG22	17	0.6
(1,453)	1:10:A:PHE:HD2	2:15:B:THR:HG23	17	0.6
(1,438)	1:8:A:GLU:HG2	2:19:B:PHE:HA	15	0.6
(1,424)	1:4:A:GLY:H	2:20:B:TYR:H	1	0.6
(1,392)	1:11:A:VAL:HA	1:12:A:ARG:HG2	2	0.6
(1,392)	1:11:A:VAL:HA	1:12:A:ARG:HG3	2	0.6
(1,371)	1:3:A:ALA:H	2:19:B:PHE:HD1	17	0.6

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,371)	1:3:A:ALA:H	2:19:B:PHE:HD2	17	0.6
(1,191)	1:3:A:ALA:H	2:19:B:PHE:HD1	17	0.6
(1,191)	1:3:A:ALA:H	2:19:B:PHE:HD2	17	0.6
(1,182)	1:4:A:GLY:H	1:6:A:VAL:H	3	0.6
(1,134)	1:10:A:PHE:HD1	2:15:B:THR:HG21	2	0.6
(1,134)	1:10:A:PHE:HD1	2:15:B:THR:HG22	2	0.6
(1,134)	1:10:A:PHE:HD1	2:15:B:THR:HG23	2	0.6
(1,134)	1:10:A:PHE:HD2	2:15:B:THR:HG21	2	0.6
(1,134)	1:10:A:PHE:HD2	2:15:B:THR:HG22	2	0.6
(1,134)	1:10:A:PHE:HD2	2:15:B:THR:HG23	2	0.6
(1,134)	1:10:A:PHE:HD1	2:15:B:THR:HG21	17	0.6
(1,134)	1:10:A:PHE:HD1	2:15:B:THR:HG22	17	0.6
(1,134)	1:10:A:PHE:HD1	2:15:B:THR:HG23	17	0.6
(1,134)	1:10:A:PHE:HD2	2:15:B:THR:HG21	17	0.6
(1,134)	1:10:A:PHE:HD2	2:15:B:THR:HG22	17	0.6
(1,134)	1:10:A:PHE:HD2	2:15:B:THR:HG23	17	0.6
(1,119)	1:8:A:GLU:HG2	2:19:B:PHE:HA	15	0.6
(1,105)	1:4:A:GLY:H	2:20:B:TYR:H	1	0.6
(1,73)	1:11:A:VAL:HA	1:12:A:ARG:HG2	2	0.6
(1,73)	1:11:A:VAL:HA	1:12:A:ARG:HG3	2	0.6
(1,52)	1:3:A:ALA:H	2:19:B:PHE:HD1	17	0.6
(1,52)	1:3:A:ALA:H	2:19:B:PHE:HD2	17	0.6
(1,1145)	1:25:C:GLY:H	2:40:D:PHE:HE1	13	0.59
(1,1145)	1:25:C:GLY:H	2:40:D:PHE:HE2	13	0.59
(1,1105)	1:24:C:ALA:HB1	1:25:C:GLY:H	10	0.59
(1,1105)	1:24:C:ALA:HB2	1:25:C:GLY:H	10	0.59
(1,1105)	1:24:C:ALA:HB3	1:25:C:GLY:H	10	0.59
(1,1105)	1:24:C:ALA:HB1	1:25:C:GLY:H	20	0.59
(1,1105)	1:24:C:ALA:HB2	1:25:C:GLY:H	20	0.59
(1,1105)	1:24:C:ALA:HB3	1:25:C:GLY:H	20	0.59
(1,1094)	1:29:C:GLU:HG2	2:40:D:PHE:HA	3	0.59
(1,1094)	1:29:C:GLU:HG2	2:40:D:PHE:HA	9	0.59
(1,1005)	2:34:D:CYS:HB3	2:35:D:GLY:HA3	16	0.59
(1,826)	1:25:C:GLY:H	2:40:D:PHE:HE1	13	0.59
(1,826)	1:25:C:GLY:H	2:40:D:PHE:HE2	13	0.59
(1,786)	1:24:C:ALA:HB1	1:25:C:GLY:H	10	0.59
(1,786)	1:24:C:ALA:HB2	1:25:C:GLY:H	10	0.59
(1,786)	1:24:C:ALA:HB3	1:25:C:GLY:H	10	0.59
(1,786)	1:24:C:ALA:HB1	1:25:C:GLY:H	20	0.59
(1,786)	1:24:C:ALA:HB2	1:25:C:GLY:H	20	0.59
(1,786)	1:24:C:ALA:HB3	1:25:C:GLY:H	20	0.59
(1,775)	1:29:C:GLU:HG2	2:40:D:PHE:HA	3	0.59

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,775)	1:29:C:GLU:HG2	2:40:D:PHE:HA	9	0.59
(1,686)	2:34:D:CYS:HB3	2:35:D:GLY:HA3	16	0.59
(1,662)	2:19:B:PHE:HE1	2:38:D:ILE:HG13	7	0.59
(1,662)	2:19:B:PHE:HE2	2:38:D:ILE:HG13	7	0.59
(1,453)	1:10:A:PHE:HD1	2:15:B:THR:HG21	11	0.59
(1,453)	1:10:A:PHE:HD1	2:15:B:THR:HG22	11	0.59
(1,453)	1:10:A:PHE:HD1	2:15:B:THR:HG23	11	0.59
(1,453)	1:10:A:PHE:HD2	2:15:B:THR:HG21	11	0.59
(1,453)	1:10:A:PHE:HD2	2:15:B:THR:HG22	11	0.59
(1,453)	1:10:A:PHE:HD2	2:15:B:THR:HG23	11	0.59
(1,449)	1:3:A:ALA:HB1	1:4:A:GLY:H	19	0.59
(1,449)	1:3:A:ALA:HB2	1:4:A:GLY:H	19	0.59
(1,449)	1:3:A:ALA:HB3	1:4:A:GLY:H	19	0.59
(1,445)	1:4:A:GLY:HA2	1:6:A:VAL:H	13	0.59
(1,395)	2:13:B:CYS:HB2	2:14:B:GLY:H	4	0.59
(1,395)	2:13:B:CYS:HB2	2:14:B:GLY:H	13	0.59
(1,369)	1:6:A:VAL:HG21	2:20:B:TYR:HE1	2	0.59
(1,369)	1:6:A:VAL:HG21	2:20:B:TYR:HE2	2	0.59
(1,369)	1:6:A:VAL:HG22	2:20:B:TYR:HE1	2	0.59
(1,369)	1:6:A:VAL:HG22	2:20:B:TYR:HE2	2	0.59
(1,369)	1:6:A:VAL:HG23	2:20:B:TYR:HE1	2	0.59
(1,369)	1:6:A:VAL:HG23	2:20:B:TYR:HE2	2	0.59
(1,134)	1:10:A:PHE:HD1	2:15:B:THR:HG21	11	0.59
(1,134)	1:10:A:PHE:HD1	2:15:B:THR:HG22	11	0.59
(1,134)	1:10:A:PHE:HD1	2:15:B:THR:HG23	11	0.59
(1,134)	1:10:A:PHE:HD2	2:15:B:THR:HG21	11	0.59
(1,134)	1:10:A:PHE:HD2	2:15:B:THR:HG22	11	0.59
(1,134)	1:10:A:PHE:HD2	2:15:B:THR:HG23	11	0.59
(1,130)	1:3:A:ALA:HB1	1:4:A:GLY:H	19	0.59
(1,130)	1:3:A:ALA:HB2	1:4:A:GLY:H	19	0.59
(1,130)	1:3:A:ALA:HB3	1:4:A:GLY:H	19	0.59
(1,126)	1:4:A:GLY:HA2	1:6:A:VAL:H	13	0.59
(1,76)	2:13:B:CYS:HB2	2:14:B:GLY:H	4	0.59
(1,76)	2:13:B:CYS:HB2	2:14:B:GLY:H	13	0.59
(1,50)	1:6:A:VAL:HG21	2:20:B:TYR:HE1	2	0.59
(1,50)	1:6:A:VAL:HG21	2:20:B:TYR:HE2	2	0.59
(1,50)	1:6:A:VAL:HG22	2:20:B:TYR:HE1	2	0.59
(1,50)	1:6:A:VAL:HG22	2:20:B:TYR:HE2	2	0.59
(1,50)	1:6:A:VAL:HG23	2:20:B:TYR:HE1	2	0.59
(1,50)	1:6:A:VAL:HG23	2:20:B:TYR:HE2	2	0.59
(1,15)	2:19:B:PHE:HE1	2:38:D:ILE:HG13	7	0.59
(1,15)	2:19:B:PHE:HE2	2:38:D:ILE:HG13	7	0.59

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,1287)	2:38:D:ILE:HA	2:38:D:ILE:HD11	5	0.58
(1,1287)	2:38:D:ILE:HA	2:38:D:ILE:HD12	5	0.58
(1,1287)	2:38:D:ILE:HA	2:38:D:ILE:HD13	5	0.58
(1,1094)	1:29:C:GLU:HG2	2:40:D:PHE:HA	12	0.58
(1,1057)	1:31:C:PHE:HD1	1:32:C:VAL:HA	18	0.58
(1,1057)	1:31:C:PHE:HD2	1:32:C:VAL:HA	18	0.58
(1,1036)	2:34:D:CYS:HB2	2:35:D:GLY:HA3	10	0.58
(1,1008)	2:18:B:SER:HA	2:38:D:ILE:HG13	18	0.58
(1,968)	2:38:D:ILE:HA	2:38:D:ILE:HD11	5	0.58
(1,968)	2:38:D:ILE:HA	2:38:D:ILE:HD12	5	0.58
(1,968)	2:38:D:ILE:HA	2:38:D:ILE:HD13	5	0.58
(1,775)	1:29:C:GLU:HG2	2:40:D:PHE:HA	12	0.58
(1,738)	1:31:C:PHE:HD1	1:32:C:VAL:HA	18	0.58
(1,738)	1:31:C:PHE:HD2	1:32:C:VAL:HA	18	0.58
(1,717)	2:34:D:CYS:HB2	2:35:D:GLY:HA3	10	0.58
(1,689)	2:18:B:SER:HA	2:38:D:ILE:HG13	18	0.58
(1,461)	1:6:A:VAL:HG11	1:7:A:PRO:HB3	17	0.58
(1,461)	1:6:A:VAL:HG12	1:7:A:PRO:HB3	17	0.58
(1,461)	1:6:A:VAL:HG13	1:7:A:PRO:HB3	17	0.58
(1,451)	2:15:B:THR:HG21	2:16:B:PRO:HB3	7	0.58
(1,451)	2:15:B:THR:HG22	2:16:B:PRO:HB3	7	0.58
(1,451)	2:15:B:THR:HG23	2:16:B:PRO:HB3	7	0.58
(1,418)	1:1:A:GLY:H	2:18:B:SER:HB3	17	0.58
(1,411)	1:5:A:HIS:HB3	1:6:A:VAL:HG21	3	0.58
(1,411)	1:5:A:HIS:HB3	1:6:A:VAL:HG22	3	0.58
(1,411)	1:5:A:HIS:HB3	1:6:A:VAL:HG23	3	0.58
(1,411)	1:5:A:HIS:HB3	1:6:A:VAL:HG21	4	0.58
(1,411)	1:5:A:HIS:HB3	1:6:A:VAL:HG22	4	0.58
(1,411)	1:5:A:HIS:HB3	1:6:A:VAL:HG23	4	0.58
(1,347)	1:7:A:PRO:HD2	2:20:B:TYR:HD1	14	0.58
(1,347)	1:7:A:PRO:HD2	2:20:B:TYR:HD2	14	0.58
(1,346)	1:9:A:TYR:HE1	2:18:B:SER:HB2	18	0.58
(1,346)	1:9:A:TYR:HE2	2:18:B:SER:HB2	18	0.58
(1,142)	1:6:A:VAL:HG11	1:7:A:PRO:HB3	17	0.58
(1,142)	1:6:A:VAL:HG12	1:7:A:PRO:HB3	17	0.58
(1,142)	1:6:A:VAL:HG13	1:7:A:PRO:HB3	17	0.58
(1,132)	2:15:B:THR:HG21	2:16:B:PRO:HB3	7	0.58
(1,132)	2:15:B:THR:HG22	2:16:B:PRO:HB3	7	0.58
(1,132)	2:15:B:THR:HG23	2:16:B:PRO:HB3	7	0.58
(1,99)	1:1:A:GLY:H	2:18:B:SER:HB3	17	0.58
(1,92)	1:5:A:HIS:HB3	1:6:A:VAL:HG21	3	0.58
(1,92)	1:5:A:HIS:HB3	1:6:A:VAL:HG22	3	0.58

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,92)	1:5:A:HIS:HB3	1:6:A:VAL:HG23	3	0.58
(1,92)	1:5:A:HIS:HB3	1:6:A:VAL:HG21	4	0.58
(1,92)	1:5:A:HIS:HB3	1:6:A:VAL:HG22	4	0.58
(1,92)	1:5:A:HIS:HB3	1:6:A:VAL:HG23	4	0.58
(1,28)	1:7:A:PRO:HD2	2:20:B:TYR:HD1	14	0.58
(1,28)	1:7:A:PRO:HD2	2:20:B:TYR:HD2	14	0.58
(1,27)	1:9:A:TYR:HE1	2:18:B:SER:HB2	18	0.58
(1,27)	1:9:A:TYR:HE2	2:18:B:SER:HB2	18	0.58
(1,1166)	1:24:C:ALA:H	2:40:D:PHE:HD1	14	0.57
(1,1166)	1:24:C:ALA:H	2:40:D:PHE:HD2	14	0.57
(1,1117)	1:27:C:VAL:HG11	1:28:C:PRO:HB3	7	0.57
(1,1117)	1:27:C:VAL:HG12	1:28:C:PRO:HB3	7	0.57
(1,1117)	1:27:C:VAL:HG13	1:28:C:PRO:HB3	7	0.57
(1,1102)	1:29:C:GLU:HB3	2:41:D:TYR:HD1	18	0.57
(1,1102)	1:29:C:GLU:HB3	2:41:D:TYR:HD2	18	0.57
(1,1097)	1:29:C:GLU:HG3	2:41:D:TYR:HB2	20	0.57
(1,1097)	1:29:C:GLU:HG3	2:41:D:TYR:HB3	20	0.57
(1,1036)	2:34:D:CYS:HB2	2:35:D:GLY:HA3	14	0.57
(1,1027)	1:24:C:ALA:H	2:40:D:PHE:HD1	14	0.57
(1,1027)	1:24:C:ALA:H	2:40:D:PHE:HD2	14	0.57
(1,847)	1:24:C:ALA:H	2:40:D:PHE:HD1	14	0.57
(1,847)	1:24:C:ALA:H	2:40:D:PHE:HD2	14	0.57
(1,798)	1:27:C:VAL:HG11	1:28:C:PRO:HB3	7	0.57
(1,798)	1:27:C:VAL:HG12	1:28:C:PRO:HB3	7	0.57
(1,798)	1:27:C:VAL:HG13	1:28:C:PRO:HB3	7	0.57
(1,783)	1:29:C:GLU:HB3	2:41:D:TYR:HD1	18	0.57
(1,783)	1:29:C:GLU:HB3	2:41:D:TYR:HD2	18	0.57
(1,778)	1:29:C:GLU:HG3	2:41:D:TYR:HB2	20	0.57
(1,778)	1:29:C:GLU:HG3	2:41:D:TYR:HB3	20	0.57
(1,717)	2:34:D:CYS:HB2	2:35:D:GLY:HA3	14	0.57
(1,708)	1:24:C:ALA:H	2:40:D:PHE:HD1	14	0.57
(1,708)	1:24:C:ALA:H	2:40:D:PHE:HD2	14	0.57
(1,513)	2:15:B:THR:HB	2:17:B:ILE:HD11	11	0.57
(1,513)	2:15:B:THR:HB	2:17:B:ILE:HD12	11	0.57
(1,513)	2:15:B:THR:HB	2:17:B:ILE:HD13	11	0.57
(1,446)	1:8:A:GLU:HB3	2:20:B:TYR:HD1	19	0.57
(1,446)	1:8:A:GLU:HB3	2:20:B:TYR:HD2	19	0.57
(1,394)	1:7:A:PRO:HG2	1:10:A:PHE:HB2	9	0.57
(1,394)	1:7:A:PRO:HG3	1:10:A:PHE:HB2	9	0.57
(1,360)	1:7:A:PRO:HG2	1:10:A:PHE:HB2	9	0.57
(1,360)	1:7:A:PRO:HG3	1:10:A:PHE:HB2	9	0.57
(1,347)	1:7:A:PRO:HD2	2:20:B:TYR:HD1	20	0.57

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,347)	1:7:A:PRO:HD2	2:20:B:TYR:HD2	20	0.57
(1,194)	2:15:B:THR:HB	2:17:B:ILE:HD11	11	0.57
(1,194)	2:15:B:THR:HB	2:17:B:ILE:HD12	11	0.57
(1,194)	2:15:B:THR:HB	2:17:B:ILE:HD13	11	0.57
(1,127)	1:8:A:GLU:HB3	2:20:B:TYR:HD1	19	0.57
(1,127)	1:8:A:GLU:HB3	2:20:B:TYR:HD2	19	0.57
(1,75)	1:7:A:PRO:HG2	1:10:A:PHE:HB2	9	0.57
(1,75)	1:7:A:PRO:HG3	1:10:A:PHE:HB2	9	0.57
(1,41)	1:7:A:PRO:HG2	1:10:A:PHE:HB2	9	0.57
(1,41)	1:7:A:PRO:HG3	1:10:A:PHE:HB2	9	0.57
(1,28)	1:7:A:PRO:HD2	2:20:B:TYR:HD1	20	0.57
(1,28)	1:7:A:PRO:HD2	2:20:B:TYR:HD2	20	0.57
(1,1163)	1:23:C:GLY:HA2	1:24:C:ALA:HB1	5	0.56
(1,1163)	1:23:C:GLY:HA2	1:24:C:ALA:HB2	5	0.56
(1,1163)	1:23:C:GLY:HA2	1:24:C:ALA:HB3	5	0.56
(1,1117)	1:27:C:VAL:HG11	1:28:C:PRO:HB3	10	0.56
(1,1117)	1:27:C:VAL:HG12	1:28:C:PRO:HB3	10	0.56
(1,1117)	1:27:C:VAL:HG13	1:28:C:PRO:HB3	10	0.56
(1,1099)	1:29:C:GLU:HG3	2:41:D:TYR:HD1	5	0.56
(1,1099)	1:29:C:GLU:HG3	2:41:D:TYR:HD2	5	0.56
(1,1053)	2:39:D:SER:HB2	2:40:D:PHE:H	11	0.56
(1,1042)	1:25:C:GLY:H	2:40:D:PHE:HD1	14	0.56
(1,1042)	1:25:C:GLY:H	2:40:D:PHE:HD2	14	0.56
(1,1025)	1:27:C:VAL:HG21	2:41:D:TYR:HE1	14	0.56
(1,1025)	1:27:C:VAL:HG21	2:41:D:TYR:HE2	14	0.56
(1,1025)	1:27:C:VAL:HG22	2:41:D:TYR:HE1	14	0.56
(1,1025)	1:27:C:VAL:HG22	2:41:D:TYR:HE2	14	0.56
(1,1025)	1:27:C:VAL:HG23	2:41:D:TYR:HE1	14	0.56
(1,1025)	1:27:C:VAL:HG23	2:41:D:TYR:HE2	14	0.56
(1,844)	1:23:C:GLY:HA2	1:24:C:ALA:HB1	5	0.56
(1,844)	1:23:C:GLY:HA2	1:24:C:ALA:HB2	5	0.56
(1,844)	1:23:C:GLY:HA2	1:24:C:ALA:HB3	5	0.56
(1,798)	1:27:C:VAL:HG11	1:28:C:PRO:HB3	10	0.56
(1,798)	1:27:C:VAL:HG12	1:28:C:PRO:HB3	10	0.56
(1,798)	1:27:C:VAL:HG13	1:28:C:PRO:HB3	10	0.56
(1,780)	1:29:C:GLU:HG3	2:41:D:TYR:HD1	5	0.56
(1,780)	1:29:C:GLU:HG3	2:41:D:TYR:HD2	5	0.56
(1,734)	2:39:D:SER:HB2	2:40:D:PHE:H	11	0.56
(1,723)	1:25:C:GLY:H	2:40:D:PHE:HD1	14	0.56
(1,723)	1:25:C:GLY:H	2:40:D:PHE:HD2	14	0.56
(1,706)	1:27:C:VAL:HG21	2:41:D:TYR:HE1	14	0.56
(1,706)	1:27:C:VAL:HG21	2:41:D:TYR:HE2	14	0.56

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,706)	1:27:C:VAL:HG22	2:41:D:TYR:HE1	14	0.56
(1,706)	1:27:C:VAL:HG22	2:41:D:TYR:HE2	14	0.56
(1,706)	1:27:C:VAL:HG23	2:41:D:TYR:HE1	14	0.56
(1,706)	1:27:C:VAL:HG23	2:41:D:TYR:HE2	14	0.56
(1,666)	2:17:B:ILE:HD11	2:40:D:PHE:HD1	18	0.56
(1,666)	2:17:B:ILE:HD11	2:40:D:PHE:HD2	18	0.56
(1,666)	2:17:B:ILE:HD12	2:40:D:PHE:HD1	18	0.56
(1,666)	2:17:B:ILE:HD12	2:40:D:PHE:HD2	18	0.56
(1,666)	2:17:B:ILE:HD13	2:40:D:PHE:HD1	18	0.56
(1,666)	2:17:B:ILE:HD13	2:40:D:PHE:HD2	18	0.56
(1,631)	2:17:B:ILE:HA	2:17:B:ILE:HD11	15	0.56
(1,631)	2:17:B:ILE:HA	2:17:B:ILE:HD12	15	0.56
(1,631)	2:17:B:ILE:HA	2:17:B:ILE:HD13	15	0.56
(1,503)	1:7:A:PRO:HB3	1:10:A:PHE:HE1	1	0.56
(1,503)	1:7:A:PRO:HB3	1:10:A:PHE:HE2	1	0.56
(1,489)	1:4:A:GLY:H	2:19:B:PHE:HE1	6	0.56
(1,489)	1:4:A:GLY:H	2:19:B:PHE:HE2	6	0.56
(1,461)	1:6:A:VAL:HG11	1:7:A:PRO:HB3	11	0.56
(1,461)	1:6:A:VAL:HG12	1:7:A:PRO:HB3	11	0.56
(1,461)	1:6:A:VAL:HG13	1:7:A:PRO:HB3	11	0.56
(1,441)	1:8:A:GLU:HG3	2:20:B:TYR:HB2	4	0.56
(1,441)	1:8:A:GLU:HG3	2:20:B:TYR:HB3	4	0.56
(1,411)	1:5:A:HIS:HB3	1:6:A:VAL:HG21	10	0.56
(1,411)	1:5:A:HIS:HB3	1:6:A:VAL:HG22	10	0.56
(1,411)	1:5:A:HIS:HB3	1:6:A:VAL:HG23	10	0.56
(1,395)	2:13:B:CYS:HB2	2:14:B:GLY:H	3	0.56
(1,390)	1:6:A:VAL:HB	2:20:B:TYR:HE1	2	0.56
(1,390)	1:6:A:VAL:HB	2:20:B:TYR:HE2	2	0.56
(1,387)	1:6:A:VAL:HG11	1:7:A:PRO:HG2	19	0.56
(1,387)	1:6:A:VAL:HG11	1:7:A:PRO:HG3	19	0.56
(1,387)	1:6:A:VAL:HG12	1:7:A:PRO:HG2	19	0.56
(1,387)	1:6:A:VAL:HG12	1:7:A:PRO:HG3	19	0.56
(1,387)	1:6:A:VAL:HG13	1:7:A:PRO:HG2	19	0.56
(1,387)	1:6:A:VAL:HG13	1:7:A:PRO:HG3	19	0.56
(1,380)	2:13:B:CYS:HB2	2:14:B:GLY:HA3	3	0.56
(1,380)	2:13:B:CYS:HB2	2:14:B:GLY:HA3	18	0.56
(1,312)	2:17:B:ILE:HA	2:17:B:ILE:HD11	15	0.56
(1,312)	2:17:B:ILE:HA	2:17:B:ILE:HD12	15	0.56
(1,312)	2:17:B:ILE:HA	2:17:B:ILE:HD13	15	0.56
(1,184)	1:7:A:PRO:HB3	1:10:A:PHE:HE1	1	0.56
(1,184)	1:7:A:PRO:HB3	1:10:A:PHE:HE2	1	0.56
(1,170)	1:4:A:GLY:H	2:19:B:PHE:HE1	6	0.56

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,170)	1:4:A:GLY:H	2:19:B:PHE:HE2	6	0.56
(1,142)	1:6:A:VAL:HG11	1:7:A:PRO:HB3	11	0.56
(1,142)	1:6:A:VAL:HG12	1:7:A:PRO:HB3	11	0.56
(1,142)	1:6:A:VAL:HG13	1:7:A:PRO:HB3	11	0.56
(1,122)	1:8:A:GLU:HG3	2:20:B:TYR:HB2	4	0.56
(1,122)	1:8:A:GLU:HG3	2:20:B:TYR:HB3	4	0.56
(1,92)	1:5:A:HIS:HB3	1:6:A:VAL:HG21	10	0.56
(1,92)	1:5:A:HIS:HB3	1:6:A:VAL:HG22	10	0.56
(1,92)	1:5:A:HIS:HB3	1:6:A:VAL:HG23	10	0.56
(1,76)	2:13:B:CYS:HB2	2:14:B:GLY:H	3	0.56
(1,71)	1:6:A:VAL:HB	2:20:B:TYR:HE1	2	0.56
(1,71)	1:6:A:VAL:HB	2:20:B:TYR:HE2	2	0.56
(1,68)	1:6:A:VAL:HG11	1:7:A:PRO:HG2	19	0.56
(1,68)	1:6:A:VAL:HG11	1:7:A:PRO:HG3	19	0.56
(1,68)	1:6:A:VAL:HG12	1:7:A:PRO:HG2	19	0.56
(1,68)	1:6:A:VAL:HG12	1:7:A:PRO:HG3	19	0.56
(1,68)	1:6:A:VAL:HG13	1:7:A:PRO:HG2	19	0.56
(1,68)	1:6:A:VAL:HG13	1:7:A:PRO:HG3	19	0.56
(1,61)	2:13:B:CYS:HB2	2:14:B:GLY:HA3	3	0.56
(1,61)	2:13:B:CYS:HB2	2:14:B:GLY:HA3	18	0.56
(1,1)	2:17:B:ILE:HD11	2:40:D:PHE:HD1	18	0.56
(1,1)	2:17:B:ILE:HD11	2:40:D:PHE:HD2	18	0.56
(1,1)	2:17:B:ILE:HD12	2:40:D:PHE:HD1	18	0.56
(1,1)	2:17:B:ILE:HD12	2:40:D:PHE:HD2	18	0.56
(1,1)	2:17:B:ILE:HD13	2:40:D:PHE:HD1	18	0.56
(1,1)	2:17:B:ILE:HD13	2:40:D:PHE:HD2	18	0.56
(1,1175)	1:24:C:ALA:HB1	2:40:D:PHE:HB3	20	0.55
(1,1175)	1:24:C:ALA:HB2	2:40:D:PHE:HB3	20	0.55
(1,1175)	1:24:C:ALA:HB3	2:40:D:PHE:HB3	20	0.55
(1,1147)	1:28:C:PRO:HB2	1:31:C:PHE:HE1	5	0.55
(1,1147)	1:28:C:PRO:HB2	1:31:C:PHE:HE2	5	0.55
(1,1105)	1:24:C:ALA:HB1	1:25:C:GLY:H	12	0.55
(1,1105)	1:24:C:ALA:HB2	1:25:C:GLY:H	12	0.55
(1,1105)	1:24:C:ALA:HB3	1:25:C:GLY:H	12	0.55
(1,1094)	1:29:C:GLU:HG2	2:40:D:PHE:HA	6	0.55
(1,1067)	1:26:C:HIS:HB3	1:27:C:VAL:HG21	2	0.55
(1,1067)	1:26:C:HIS:HB3	1:27:C:VAL:HG22	2	0.55
(1,1067)	1:26:C:HIS:HB3	1:27:C:VAL:HG23	2	0.55
(1,1067)	1:26:C:HIS:HB3	1:27:C:VAL:HG21	8	0.55
(1,1067)	1:26:C:HIS:HB3	1:27:C:VAL:HG22	8	0.55
(1,1067)	1:26:C:HIS:HB3	1:27:C:VAL:HG23	8	0.55
(1,1043)	1:27:C:VAL:HG11	1:28:C:PRO:HG2	8	0.55

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,1043)	1:27:C:VAL:HG11	1:28:C:PRO:HG3	8	0.55
(1,1043)	1:27:C:VAL:HG12	1:28:C:PRO:HG2	8	0.55
(1,1043)	1:27:C:VAL:HG12	1:28:C:PRO:HG3	8	0.55
(1,1043)	1:27:C:VAL:HG13	1:28:C:PRO:HG2	8	0.55
(1,1043)	1:27:C:VAL:HG13	1:28:C:PRO:HG3	8	0.55
(1,1000)	1:31:C:PHE:HE1	2:37:D:PRO:HD2	5	0.55
(1,1000)	1:31:C:PHE:HE2	2:37:D:PRO:HD2	5	0.55
(1,856)	1:24:C:ALA:HB1	2:40:D:PHE:HB3	20	0.55
(1,856)	1:24:C:ALA:HB2	2:40:D:PHE:HB3	20	0.55
(1,856)	1:24:C:ALA:HB3	2:40:D:PHE:HB3	20	0.55
(1,828)	1:28:C:PRO:HB2	1:31:C:PHE:HE1	5	0.55
(1,828)	1:28:C:PRO:HB2	1:31:C:PHE:HE2	5	0.55
(1,786)	1:24:C:ALA:HB1	1:25:C:GLY:H	12	0.55
(1,786)	1:24:C:ALA:HB2	1:25:C:GLY:H	12	0.55
(1,786)	1:24:C:ALA:HB3	1:25:C:GLY:H	12	0.55
(1,775)	1:29:C:GLU:HG2	2:40:D:PHE:HA	6	0.55
(1,748)	1:26:C:HIS:HB3	1:27:C:VAL:HG21	2	0.55
(1,748)	1:26:C:HIS:HB3	1:27:C:VAL:HG22	2	0.55
(1,748)	1:26:C:HIS:HB3	1:27:C:VAL:HG23	2	0.55
(1,748)	1:26:C:HIS:HB3	1:27:C:VAL:HG21	8	0.55
(1,748)	1:26:C:HIS:HB3	1:27:C:VAL:HG22	8	0.55
(1,748)	1:26:C:HIS:HB3	1:27:C:VAL:HG23	8	0.55
(1,724)	1:27:C:VAL:HG11	1:28:C:PRO:HG2	8	0.55
(1,724)	1:27:C:VAL:HG11	1:28:C:PRO:HG3	8	0.55
(1,724)	1:27:C:VAL:HG12	1:28:C:PRO:HG2	8	0.55
(1,724)	1:27:C:VAL:HG12	1:28:C:PRO:HG3	8	0.55
(1,724)	1:27:C:VAL:HG13	1:28:C:PRO:HG2	8	0.55
(1,724)	1:27:C:VAL:HG13	1:28:C:PRO:HG3	8	0.55
(1,681)	1:31:C:PHE:HE1	2:37:D:PRO:HD2	5	0.55
(1,681)	1:31:C:PHE:HE2	2:37:D:PRO:HD2	5	0.55
(1,587)	1:12:A:ARG:H	1:12:A:ARG:HE	14	0.55
(1,468)	1:11:A:VAL:HG11	1:12:A:ARG:H	4	0.55
(1,468)	1:11:A:VAL:HG12	1:12:A:ARG:H	4	0.55
(1,468)	1:11:A:VAL:HG13	1:12:A:ARG:H	4	0.55
(1,468)	1:11:A:VAL:HG21	1:12:A:ARG:H	4	0.55
(1,468)	1:11:A:VAL:HG22	1:12:A:ARG:H	4	0.55
(1,468)	1:11:A:VAL:HG23	1:12:A:ARG:H	4	0.55
(1,461)	1:6:A:VAL:HG11	1:7:A:PRO:HB3	4	0.55
(1,461)	1:6:A:VAL:HG12	1:7:A:PRO:HB3	4	0.55
(1,461)	1:6:A:VAL:HG13	1:7:A:PRO:HB3	4	0.55
(1,449)	1:3:A:ALA:HB1	1:4:A:GLY:H	3	0.55
(1,449)	1:3:A:ALA:HB2	1:4:A:GLY:H	3	0.55

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,449)	1:3:A:ALA:HB3	1:4:A:GLY:H	3	0.55
(1,438)	1:8:A:GLU:HG2	2:19:B:PHE:HA	19	0.55
(1,387)	1:6:A:VAL:HG11	1:7:A:PRO:HG2	10	0.55
(1,387)	1:6:A:VAL:HG11	1:7:A:PRO:HG3	10	0.55
(1,387)	1:6:A:VAL:HG12	1:7:A:PRO:HG2	10	0.55
(1,387)	1:6:A:VAL:HG12	1:7:A:PRO:HG3	10	0.55
(1,387)	1:6:A:VAL:HG13	1:7:A:PRO:HG2	10	0.55
(1,387)	1:6:A:VAL:HG13	1:7:A:PRO:HG3	10	0.55
(1,268)	1:12:A:ARG:H	1:12:A:ARG:HE	14	0.55
(1,149)	1:11:A:VAL:HG11	1:12:A:ARG:H	4	0.55
(1,149)	1:11:A:VAL:HG12	1:12:A:ARG:H	4	0.55
(1,149)	1:11:A:VAL:HG13	1:12:A:ARG:H	4	0.55
(1,149)	1:11:A:VAL:HG21	1:12:A:ARG:H	4	0.55
(1,149)	1:11:A:VAL:HG22	1:12:A:ARG:H	4	0.55
(1,149)	1:11:A:VAL:HG23	1:12:A:ARG:H	4	0.55
(1,142)	1:6:A:VAL:HG11	1:7:A:PRO:HB3	4	0.55
(1,142)	1:6:A:VAL:HG12	1:7:A:PRO:HB3	4	0.55
(1,142)	1:6:A:VAL:HG13	1:7:A:PRO:HB3	4	0.55
(1,130)	1:3:A:ALA:HB1	1:4:A:GLY:H	3	0.55
(1,130)	1:3:A:ALA:HB2	1:4:A:GLY:H	3	0.55
(1,130)	1:3:A:ALA:HB3	1:4:A:GLY:H	3	0.55
(1,119)	1:8:A:GLU:HG2	2:19:B:PHE:HA	19	0.55
(1,68)	1:6:A:VAL:HG11	1:7:A:PRO:HG2	10	0.55
(1,68)	1:6:A:VAL:HG11	1:7:A:PRO:HG3	10	0.55
(1,68)	1:6:A:VAL:HG12	1:7:A:PRO:HG2	10	0.55
(1,68)	1:6:A:VAL:HG12	1:7:A:PRO:HG3	10	0.55
(1,68)	1:6:A:VAL:HG13	1:7:A:PRO:HG2	10	0.55
(1,68)	1:6:A:VAL:HG13	1:7:A:PRO:HG3	10	0.55
(1,1116)	1:26:C:HIS:HD2	1:27:C:VAL:HG21	1	0.54
(1,1116)	1:26:C:HIS:HD2	1:27:C:VAL:HG22	1	0.54
(1,1116)	1:26:C:HIS:HD2	1:27:C:VAL:HG23	1	0.54
(1,1102)	1:29:C:GLU:HB3	2:41:D:TYR:HD1	16	0.54
(1,1102)	1:29:C:GLU:HB3	2:41:D:TYR:HD2	16	0.54
(1,1063)	1:22:C:GLY:H	2:39:D:SER:HA	8	0.54
(1,797)	1:26:C:HIS:HD2	1:27:C:VAL:HG21	1	0.54
(1,797)	1:26:C:HIS:HD2	1:27:C:VAL:HG22	1	0.54
(1,797)	1:26:C:HIS:HD2	1:27:C:VAL:HG23	1	0.54
(1,783)	1:29:C:GLU:HB3	2:41:D:TYR:HD1	16	0.54
(1,783)	1:29:C:GLU:HB3	2:41:D:TYR:HD2	16	0.54
(1,744)	1:22:C:GLY:H	2:39:D:SER:HA	8	0.54
(1,663)	2:19:B:PHE:HD1	2:38:D:ILE:HG13	19	0.54
(1,663)	2:19:B:PHE:HD2	2:38:D:ILE:HG13	19	0.54

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,519)	1:3:A:ALA:HB1	2:19:B:PHE:HB3	14	0.54
(1,519)	1:3:A:ALA:HB2	2:19:B:PHE:HB3	14	0.54
(1,519)	1:3:A:ALA:HB3	2:19:B:PHE:HB3	14	0.54
(1,452)	2:15:B:THR:HG21	2:16:B:PRO:HD3	4	0.54
(1,452)	2:15:B:THR:HG22	2:16:B:PRO:HD3	4	0.54
(1,452)	2:15:B:THR:HG23	2:16:B:PRO:HD3	4	0.54
(1,438)	1:8:A:GLU:HG2	2:19:B:PHE:HA	10	0.54
(1,398)	1:7:A:PRO:HA	2:19:B:PHE:HD1	17	0.54
(1,398)	1:7:A:PRO:HA	2:19:B:PHE:HD2	17	0.54
(1,369)	1:6:A:VAL:HG21	2:20:B:TYR:HE1	3	0.54
(1,369)	1:6:A:VAL:HG21	2:20:B:TYR:HE2	3	0.54
(1,369)	1:6:A:VAL:HG22	2:20:B:TYR:HE1	3	0.54
(1,369)	1:6:A:VAL:HG22	2:20:B:TYR:HE2	3	0.54
(1,369)	1:6:A:VAL:HG23	2:20:B:TYR:HE1	3	0.54
(1,369)	1:6:A:VAL:HG23	2:20:B:TYR:HE2	3	0.54
(1,351)	2:16:B:PRO:HA	2:17:B:ILE:HD11	6	0.54
(1,351)	2:16:B:PRO:HA	2:17:B:ILE:HD12	6	0.54
(1,351)	2:16:B:PRO:HA	2:17:B:ILE:HD13	6	0.54
(1,200)	1:3:A:ALA:HB1	2:19:B:PHE:HB3	14	0.54
(1,200)	1:3:A:ALA:HB2	2:19:B:PHE:HB3	14	0.54
(1,200)	1:3:A:ALA:HB3	2:19:B:PHE:HB3	14	0.54
(1,133)	2:15:B:THR:HG21	2:16:B:PRO:HD3	4	0.54
(1,133)	2:15:B:THR:HG22	2:16:B:PRO:HD3	4	0.54
(1,133)	2:15:B:THR:HG23	2:16:B:PRO:HD3	4	0.54
(1,119)	1:8:A:GLU:HG2	2:19:B:PHE:HA	10	0.54
(1,79)	1:7:A:PRO:HA	2:19:B:PHE:HD1	17	0.54
(1,79)	1:7:A:PRO:HA	2:19:B:PHE:HD2	17	0.54
(1,50)	1:6:A:VAL:HG21	2:20:B:TYR:HE1	3	0.54
(1,50)	1:6:A:VAL:HG21	2:20:B:TYR:HE2	3	0.54
(1,50)	1:6:A:VAL:HG22	2:20:B:TYR:HE1	3	0.54
(1,50)	1:6:A:VAL:HG22	2:20:B:TYR:HE2	3	0.54
(1,50)	1:6:A:VAL:HG23	2:20:B:TYR:HE1	3	0.54
(1,50)	1:6:A:VAL:HG23	2:20:B:TYR:HE2	3	0.54
(1,32)	2:16:B:PRO:HA	2:17:B:ILE:HD11	6	0.54
(1,32)	2:16:B:PRO:HA	2:17:B:ILE:HD12	6	0.54
(1,32)	2:16:B:PRO:HA	2:17:B:ILE:HD13	6	0.54
(1,16)	2:19:B:PHE:HD1	2:38:D:ILE:HG13	19	0.54
(1,16)	2:19:B:PHE:HD2	2:38:D:ILE:HG13	19	0.54
(1,1309)	1:24:C:ALA:HA	2:40:D:PHE:HD1	15	0.53
(1,1309)	1:24:C:ALA:HA	2:40:D:PHE:HD2	15	0.53
(1,1117)	1:27:C:VAL:HG11	1:28:C:PRO:HB3	9	0.53
(1,1117)	1:27:C:VAL:HG12	1:28:C:PRO:HB3	9	0.53

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,1117)	1:27:C:VAL:HG13	1:28:C:PRO:HB3	9	0.53
(1,1117)	1:27:C:VAL:HG11	1:28:C:PRO:HB3	12	0.53
(1,1117)	1:27:C:VAL:HG12	1:28:C:PRO:HB3	12	0.53
(1,1117)	1:27:C:VAL:HG13	1:28:C:PRO:HB3	12	0.53
(1,1105)	1:24:C:ALA:HB1	1:25:C:GLY:H	1	0.53
(1,1105)	1:24:C:ALA:HB2	1:25:C:GLY:H	1	0.53
(1,1105)	1:24:C:ALA:HB3	1:25:C:GLY:H	1	0.53
(1,1105)	1:24:C:ALA:HB1	1:25:C:GLY:H	17	0.53
(1,1105)	1:24:C:ALA:HB2	1:25:C:GLY:H	17	0.53
(1,1105)	1:24:C:ALA:HB3	1:25:C:GLY:H	17	0.53
(1,1097)	1:29:C:GLU:HG3	2:41:D:TYR:HB2	6	0.53
(1,1097)	1:29:C:GLU:HG3	2:41:D:TYR:HB3	6	0.53
(1,1047)	1:24:C:ALA:HA	2:40:D:PHE:HD1	15	0.53
(1,1047)	1:24:C:ALA:HA	2:40:D:PHE:HD2	15	0.53
(1,1005)	2:34:D:CYS:HB3	2:35:D:GLY:HA3	19	0.53
(1,990)	1:24:C:ALA:HA	2:40:D:PHE:HD1	15	0.53
(1,990)	1:24:C:ALA:HA	2:40:D:PHE:HD2	15	0.53
(1,798)	1:27:C:VAL:HG11	1:28:C:PRO:HB3	9	0.53
(1,798)	1:27:C:VAL:HG12	1:28:C:PRO:HB3	9	0.53
(1,798)	1:27:C:VAL:HG13	1:28:C:PRO:HB3	9	0.53
(1,798)	1:27:C:VAL:HG11	1:28:C:PRO:HB3	12	0.53
(1,798)	1:27:C:VAL:HG12	1:28:C:PRO:HB3	12	0.53
(1,798)	1:27:C:VAL:HG13	1:28:C:PRO:HB3	12	0.53
(1,786)	1:24:C:ALA:HB1	1:25:C:GLY:H	1	0.53
(1,786)	1:24:C:ALA:HB2	1:25:C:GLY:H	1	0.53
(1,786)	1:24:C:ALA:HB3	1:25:C:GLY:H	1	0.53
(1,786)	1:24:C:ALA:HB1	1:25:C:GLY:H	17	0.53
(1,786)	1:24:C:ALA:HB2	1:25:C:GLY:H	17	0.53
(1,786)	1:24:C:ALA:HB3	1:25:C:GLY:H	17	0.53
(1,778)	1:29:C:GLU:HG3	2:41:D:TYR:HB2	6	0.53
(1,778)	1:29:C:GLU:HG3	2:41:D:TYR:HB3	6	0.53
(1,728)	1:24:C:ALA:HA	2:40:D:PHE:HD1	15	0.53
(1,728)	1:24:C:ALA:HA	2:40:D:PHE:HD2	15	0.53
(1,686)	2:34:D:CYS:HB3	2:35:D:GLY:HA3	19	0.53
(1,662)	2:19:B:PHE:HE1	2:38:D:ILE:HG13	6	0.53
(1,662)	2:19:B:PHE:HE2	2:38:D:ILE:HG13	6	0.53
(1,659)	1:11:A:VAL:HA	1:24:C:ALA:HB1	18	0.53
(1,659)	1:11:A:VAL:HA	1:24:C:ALA:HB2	18	0.53
(1,659)	1:11:A:VAL:HA	1:24:C:ALA:HB3	18	0.53
(1,481)	1:1:A:GLY:H	2:19:B:PHE:H	15	0.53
(1,387)	1:6:A:VAL:HG11	1:7:A:PRO:HG2	5	0.53
(1,387)	1:6:A:VAL:HG11	1:7:A:PRO:HG3	5	0.53

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,387)	1:6:A:VAL:HG12	1:7:A:PRO:HG2	5	0.53
(1,387)	1:6:A:VAL:HG12	1:7:A:PRO:HG3	5	0.53
(1,387)	1:6:A:VAL:HG13	1:7:A:PRO:HG2	5	0.53
(1,387)	1:6:A:VAL:HG13	1:7:A:PRO:HG3	5	0.53
(1,387)	1:6:A:VAL:HG11	1:7:A:PRO:HG2	15	0.53
(1,387)	1:6:A:VAL:HG11	1:7:A:PRO:HG3	15	0.53
(1,387)	1:6:A:VAL:HG12	1:7:A:PRO:HG2	15	0.53
(1,387)	1:6:A:VAL:HG12	1:7:A:PRO:HG3	15	0.53
(1,387)	1:6:A:VAL:HG13	1:7:A:PRO:HG2	15	0.53
(1,387)	1:6:A:VAL:HG13	1:7:A:PRO:HG3	15	0.53
(1,369)	1:6:A:VAL:HG21	2:20:B:TYR:HE1	10	0.53
(1,369)	1:6:A:VAL:HG21	2:20:B:TYR:HE2	10	0.53
(1,369)	1:6:A:VAL:HG22	2:20:B:TYR:HE1	10	0.53
(1,369)	1:6:A:VAL:HG22	2:20:B:TYR:HE2	10	0.53
(1,369)	1:6:A:VAL:HG23	2:20:B:TYR:HE1	10	0.53
(1,369)	1:6:A:VAL:HG23	2:20:B:TYR:HE2	10	0.53
(1,369)	1:6:A:VAL:HG21	2:20:B:TYR:HE1	12	0.53
(1,369)	1:6:A:VAL:HG21	2:20:B:TYR:HE2	12	0.53
(1,369)	1:6:A:VAL:HG22	2:20:B:TYR:HE1	12	0.53
(1,369)	1:6:A:VAL:HG22	2:20:B:TYR:HE2	12	0.53
(1,369)	1:6:A:VAL:HG23	2:20:B:TYR:HE1	12	0.53
(1,369)	1:6:A:VAL:HG23	2:20:B:TYR:HE2	12	0.53
(1,162)	1:1:A:GLY:H	2:19:B:PHE:H	15	0.53
(1,68)	1:6:A:VAL:HG11	1:7:A:PRO:HG2	5	0.53
(1,68)	1:6:A:VAL:HG11	1:7:A:PRO:HG3	5	0.53
(1,68)	1:6:A:VAL:HG12	1:7:A:PRO:HG2	5	0.53
(1,68)	1:6:A:VAL:HG12	1:7:A:PRO:HG3	5	0.53
(1,68)	1:6:A:VAL:HG13	1:7:A:PRO:HG2	5	0.53
(1,68)	1:6:A:VAL:HG13	1:7:A:PRO:HG3	5	0.53
(1,68)	1:6:A:VAL:HG11	1:7:A:PRO:HG2	15	0.53
(1,68)	1:6:A:VAL:HG11	1:7:A:PRO:HG3	15	0.53
(1,68)	1:6:A:VAL:HG12	1:7:A:PRO:HG2	15	0.53
(1,68)	1:6:A:VAL:HG12	1:7:A:PRO:HG3	15	0.53
(1,68)	1:6:A:VAL:HG13	1:7:A:PRO:HG2	15	0.53
(1,68)	1:6:A:VAL:HG13	1:7:A:PRO:HG3	15	0.53
(1,50)	1:6:A:VAL:HG21	2:20:B:TYR:HE1	10	0.53
(1,50)	1:6:A:VAL:HG21	2:20:B:TYR:HE2	10	0.53
(1,50)	1:6:A:VAL:HG22	2:20:B:TYR:HE1	10	0.53
(1,50)	1:6:A:VAL:HG22	2:20:B:TYR:HE2	10	0.53
(1,50)	1:6:A:VAL:HG23	2:20:B:TYR:HE1	10	0.53
(1,50)	1:6:A:VAL:HG23	2:20:B:TYR:HE2	10	0.53
(1,50)	1:6:A:VAL:HG21	2:20:B:TYR:HE1	12	0.53

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,50)	1:6:A:VAL:HG21	2:20:B:TYR:HE2	12	0.53
(1,50)	1:6:A:VAL:HG22	2:20:B:TYR:HE1	12	0.53
(1,50)	1:6:A:VAL:HG22	2:20:B:TYR:HE2	12	0.53
(1,50)	1:6:A:VAL:HG23	2:20:B:TYR:HE1	12	0.53
(1,50)	1:6:A:VAL:HG23	2:20:B:TYR:HE2	12	0.53
(1,15)	2:19:B:PHE:HE1	2:38:D:ILE:HG13	6	0.53
(1,15)	2:19:B:PHE:HE2	2:38:D:ILE:HG13	6	0.53
(1,12)	1:11:A:VAL:HA	1:24:C:ALA:HB1	18	0.53
(1,12)	1:11:A:VAL:HA	1:24:C:ALA:HB2	18	0.53
(1,12)	1:11:A:VAL:HA	1:24:C:ALA:HB3	18	0.53
(1,1159)	1:28:C:PRO:HB3	1:31:C:PHE:HE1	20	0.52
(1,1159)	1:28:C:PRO:HB3	1:31:C:PHE:HE2	20	0.52
(1,1134)	2:39:D:SER:HB3	2:40:D:PHE:H	9	0.52
(1,1132)	2:39:D:SER:HB3	2:40:D:PHE:H	9	0.52
(1,1071)	1:27:C:VAL:HG11	2:41:D:TYR:HE1	4	0.52
(1,1071)	1:27:C:VAL:HG11	2:41:D:TYR:HE2	4	0.52
(1,1071)	1:27:C:VAL:HG12	2:41:D:TYR:HE1	4	0.52
(1,1071)	1:27:C:VAL:HG12	2:41:D:TYR:HE2	4	0.52
(1,1071)	1:27:C:VAL:HG13	2:41:D:TYR:HE1	4	0.52
(1,1071)	1:27:C:VAL:HG13	2:41:D:TYR:HE2	4	0.52
(1,1005)	2:34:D:CYS:HB3	2:35:D:GLY:HA3	2	0.52
(1,840)	1:28:C:PRO:HB3	1:31:C:PHE:HE1	20	0.52
(1,840)	1:28:C:PRO:HB3	1:31:C:PHE:HE2	20	0.52
(1,815)	2:39:D:SER:HB3	2:40:D:PHE:H	9	0.52
(1,813)	2:39:D:SER:HB3	2:40:D:PHE:H	9	0.52
(1,752)	1:27:C:VAL:HG11	2:41:D:TYR:HE1	4	0.52
(1,752)	1:27:C:VAL:HG11	2:41:D:TYR:HE2	4	0.52
(1,752)	1:27:C:VAL:HG12	2:41:D:TYR:HE1	4	0.52
(1,752)	1:27:C:VAL:HG12	2:41:D:TYR:HE2	4	0.52
(1,752)	1:27:C:VAL:HG13	2:41:D:TYR:HE1	4	0.52
(1,752)	1:27:C:VAL:HG13	2:41:D:TYR:HE2	4	0.52
(1,686)	2:34:D:CYS:HB3	2:35:D:GLY:HA3	2	0.52
(1,519)	1:3:A:ALA:HB1	2:19:B:PHE:HB3	5	0.52
(1,519)	1:3:A:ALA:HB2	2:19:B:PHE:HB3	5	0.52
(1,519)	1:3:A:ALA:HB3	2:19:B:PHE:HB3	5	0.52
(1,519)	1:3:A:ALA:HB1	2:19:B:PHE:HB3	6	0.52
(1,519)	1:3:A:ALA:HB2	2:19:B:PHE:HB3	6	0.52
(1,519)	1:3:A:ALA:HB3	2:19:B:PHE:HB3	6	0.52
(1,519)	1:3:A:ALA:HB1	2:19:B:PHE:HB3	13	0.52
(1,519)	1:3:A:ALA:HB2	2:19:B:PHE:HB3	13	0.52
(1,519)	1:3:A:ALA:HB3	2:19:B:PHE:HB3	13	0.52
(1,510)	1:3:A:ALA:H	2:19:B:PHE:HD1	12	0.52

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,510)	1:3:A:ALA:H	2:19:B:PHE:HD2	12	0.52
(1,499)	1:6:A:VAL:HG11	1:7:A:PRO:HD3	10	0.52
(1,499)	1:6:A:VAL:HG12	1:7:A:PRO:HD3	10	0.52
(1,499)	1:6:A:VAL:HG13	1:7:A:PRO:HD3	10	0.52
(1,465)	1:5:A:HIS:HD2	1:6:A:VAL:HG11	2	0.52
(1,465)	1:5:A:HIS:HD2	1:6:A:VAL:HG12	2	0.52
(1,465)	1:5:A:HIS:HD2	1:6:A:VAL:HG13	2	0.52
(1,446)	1:8:A:GLU:HB3	2:20:B:TYR:HD1	16	0.52
(1,446)	1:8:A:GLU:HB3	2:20:B:TYR:HD2	16	0.52
(1,443)	1:8:A:GLU:HG3	2:20:B:TYR:HD1	19	0.52
(1,443)	1:8:A:GLU:HG3	2:20:B:TYR:HD2	19	0.52
(1,438)	1:8:A:GLU:HG2	2:19:B:PHE:HA	20	0.52
(1,395)	2:13:B:CYS:HB2	2:14:B:GLY:H	16	0.52
(1,387)	1:6:A:VAL:HG11	1:7:A:PRO:HG2	8	0.52
(1,387)	1:6:A:VAL:HG11	1:7:A:PRO:HG3	8	0.52
(1,387)	1:6:A:VAL:HG12	1:7:A:PRO:HG2	8	0.52
(1,387)	1:6:A:VAL:HG12	1:7:A:PRO:HG3	8	0.52
(1,387)	1:6:A:VAL:HG13	1:7:A:PRO:HG2	8	0.52
(1,387)	1:6:A:VAL:HG13	1:7:A:PRO:HG3	8	0.52
(1,371)	1:3:A:ALA:H	2:19:B:PHE:HD1	12	0.52
(1,371)	1:3:A:ALA:H	2:19:B:PHE:HD2	12	0.52
(1,369)	1:6:A:VAL:HG21	2:20:B:TYR:HE1	9	0.52
(1,369)	1:6:A:VAL:HG21	2:20:B:TYR:HE2	9	0.52
(1,369)	1:6:A:VAL:HG22	2:20:B:TYR:HE1	9	0.52
(1,369)	1:6:A:VAL:HG22	2:20:B:TYR:HE2	9	0.52
(1,369)	1:6:A:VAL:HG23	2:20:B:TYR:HE1	9	0.52
(1,369)	1:6:A:VAL:HG23	2:20:B:TYR:HE2	9	0.52
(1,356)	1:8:A:GLU:HG2	2:20:B:TYR:HE1	5	0.52
(1,356)	1:8:A:GLU:HG2	2:20:B:TYR:HE2	5	0.52
(1,200)	1:3:A:ALA:HB1	2:19:B:PHE:HB3	5	0.52
(1,200)	1:3:A:ALA:HB2	2:19:B:PHE:HB3	5	0.52
(1,200)	1:3:A:ALA:HB3	2:19:B:PHE:HB3	5	0.52
(1,200)	1:3:A:ALA:HB1	2:19:B:PHE:HB3	6	0.52
(1,200)	1:3:A:ALA:HB2	2:19:B:PHE:HB3	6	0.52
(1,200)	1:3:A:ALA:HB3	2:19:B:PHE:HB3	6	0.52
(1,200)	1:3:A:ALA:HB1	2:19:B:PHE:HB3	13	0.52
(1,200)	1:3:A:ALA:HB2	2:19:B:PHE:HB3	13	0.52
(1,200)	1:3:A:ALA:HB3	2:19:B:PHE:HB3	13	0.52
(1,191)	1:3:A:ALA:H	2:19:B:PHE:HD1	12	0.52
(1,191)	1:3:A:ALA:H	2:19:B:PHE:HD2	12	0.52
(1,180)	1:6:A:VAL:HG11	1:7:A:PRO:HD3	10	0.52
(1,180)	1:6:A:VAL:HG12	1:7:A:PRO:HD3	10	0.52

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,180)	1:6:A:VAL:HG13	1:7:A:PRO:HD3	10	0.52
(1,146)	1:5:A:HIS:HD2	1:6:A:VAL:HG11	2	0.52
(1,146)	1:5:A:HIS:HD2	1:6:A:VAL:HG12	2	0.52
(1,146)	1:5:A:HIS:HD2	1:6:A:VAL:HG13	2	0.52
(1,127)	1:8:A:GLU:HB3	2:20:B:TYR:HD1	16	0.52
(1,127)	1:8:A:GLU:HB3	2:20:B:TYR:HD2	16	0.52
(1,124)	1:8:A:GLU:HG3	2:20:B:TYR:HD1	19	0.52
(1,124)	1:8:A:GLU:HG3	2:20:B:TYR:HD2	19	0.52
(1,119)	1:8:A:GLU:HG2	2:19:B:PHE:HA	20	0.52
(1,76)	2:13:B:CYS:HB2	2:14:B:GLY:H	16	0.52
(1,68)	1:6:A:VAL:HG11	1:7:A:PRO:HG2	8	0.52
(1,68)	1:6:A:VAL:HG11	1:7:A:PRO:HG3	8	0.52
(1,68)	1:6:A:VAL:HG12	1:7:A:PRO:HG2	8	0.52
(1,68)	1:6:A:VAL:HG12	1:7:A:PRO:HG3	8	0.52
(1,68)	1:6:A:VAL:HG13	1:7:A:PRO:HG2	8	0.52
(1,68)	1:6:A:VAL:HG13	1:7:A:PRO:HG3	8	0.52
(1,52)	1:3:A:ALA:H	2:19:B:PHE:HD1	12	0.52
(1,52)	1:3:A:ALA:H	2:19:B:PHE:HD2	12	0.52
(1,50)	1:6:A:VAL:HG21	2:20:B:TYR:HE1	9	0.52
(1,50)	1:6:A:VAL:HG21	2:20:B:TYR:HE2	9	0.52
(1,50)	1:6:A:VAL:HG22	2:20:B:TYR:HE1	9	0.52
(1,50)	1:6:A:VAL:HG22	2:20:B:TYR:HE2	9	0.52
(1,50)	1:6:A:VAL:HG23	2:20:B:TYR:HE1	9	0.52
(1,50)	1:6:A:VAL:HG23	2:20:B:TYR:HE2	9	0.52
(1,37)	1:8:A:GLU:HG2	2:20:B:TYR:HE1	5	0.52
(1,37)	1:8:A:GLU:HG2	2:20:B:TYR:HE2	5	0.52
(1,1155)	1:27:C:VAL:HG11	1:28:C:PRO:HD3	2	0.51
(1,1155)	1:27:C:VAL:HG12	1:28:C:PRO:HD3	2	0.51
(1,1155)	1:27:C:VAL:HG13	1:28:C:PRO:HD3	2	0.51
(1,1051)	2:34:D:CYS:HB2	2:35:D:GLY:H	13	0.51
(1,836)	1:27:C:VAL:HG11	1:28:C:PRO:HD3	2	0.51
(1,836)	1:27:C:VAL:HG12	1:28:C:PRO:HD3	2	0.51
(1,836)	1:27:C:VAL:HG13	1:28:C:PRO:HD3	2	0.51
(1,732)	2:34:D:CYS:HB2	2:35:D:GLY:H	13	0.51
(1,663)	2:19:B:PHE:HD1	2:38:D:ILE:HG13	18	0.51
(1,663)	2:19:B:PHE:HD2	2:38:D:ILE:HG13	18	0.51
(1,483)	1:1:A:GLY:H	2:20:B:TYR:HD1	1	0.51
(1,483)	1:1:A:GLY:H	2:20:B:TYR:HD2	1	0.51
(1,446)	1:8:A:GLU:HB3	2:20:B:TYR:HD1	7	0.51
(1,446)	1:8:A:GLU:HB3	2:20:B:TYR:HD2	7	0.51
(1,411)	1:5:A:HIS:HB3	1:6:A:VAL:HG21	9	0.51
(1,411)	1:5:A:HIS:HB3	1:6:A:VAL:HG22	9	0.51

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,411)	1:5:A:HIS:HB3	1:6:A:VAL:HG23	9	0.51
(1,403)	1:7:A:PRO:HG2	2:19:B:PHE:HD1	12	0.51
(1,403)	1:7:A:PRO:HG2	2:19:B:PHE:HD2	12	0.51
(1,403)	1:7:A:PRO:HG3	2:19:B:PHE:HD1	12	0.51
(1,403)	1:7:A:PRO:HG3	2:19:B:PHE:HD2	12	0.51
(1,398)	1:7:A:PRO:HA	2:19:B:PHE:HD1	4	0.51
(1,398)	1:7:A:PRO:HA	2:19:B:PHE:HD2	4	0.51
(1,387)	1:6:A:VAL:HG11	1:7:A:PRO:HG2	7	0.51
(1,387)	1:6:A:VAL:HG11	1:7:A:PRO:HG3	7	0.51
(1,387)	1:6:A:VAL:HG12	1:7:A:PRO:HG2	7	0.51
(1,387)	1:6:A:VAL:HG12	1:7:A:PRO:HG3	7	0.51
(1,387)	1:6:A:VAL:HG13	1:7:A:PRO:HG2	7	0.51
(1,387)	1:6:A:VAL:HG13	1:7:A:PRO:HG3	7	0.51
(1,369)	1:6:A:VAL:HG21	2:20:B:TYR:HE1	17	0.51
(1,369)	1:6:A:VAL:HG21	2:20:B:TYR:HE2	17	0.51
(1,369)	1:6:A:VAL:HG22	2:20:B:TYR:HE1	17	0.51
(1,369)	1:6:A:VAL:HG22	2:20:B:TYR:HE2	17	0.51
(1,369)	1:6:A:VAL:HG23	2:20:B:TYR:HE1	17	0.51
(1,369)	1:6:A:VAL:HG23	2:20:B:TYR:HE2	17	0.51
(1,362)	1:7:A:PRO:HG2	2:19:B:PHE:HD1	5	0.51
(1,362)	1:7:A:PRO:HG2	2:19:B:PHE:HD2	5	0.51
(1,362)	1:7:A:PRO:HG3	2:19:B:PHE:HD1	5	0.51
(1,362)	1:7:A:PRO:HG3	2:19:B:PHE:HD2	5	0.51
(1,351)	2:16:B:PRO:HA	2:17:B:ILE:HD11	14	0.51
(1,351)	2:16:B:PRO:HA	2:17:B:ILE:HD12	14	0.51
(1,351)	2:16:B:PRO:HA	2:17:B:ILE:HD13	14	0.51
(1,164)	1:1:A:GLY:H	2:20:B:TYR:HD1	1	0.51
(1,164)	1:1:A:GLY:H	2:20:B:TYR:HD2	1	0.51
(1,127)	1:8:A:GLU:HB3	2:20:B:TYR:HD1	7	0.51
(1,127)	1:8:A:GLU:HB3	2:20:B:TYR:HD2	7	0.51
(1,92)	1:5:A:HIS:HB3	1:6:A:VAL:HG21	9	0.51
(1,92)	1:5:A:HIS:HB3	1:6:A:VAL:HG22	9	0.51
(1,92)	1:5:A:HIS:HB3	1:6:A:VAL:HG23	9	0.51
(1,84)	1:7:A:PRO:HG2	2:19:B:PHE:HD1	12	0.51
(1,84)	1:7:A:PRO:HG2	2:19:B:PHE:HD2	12	0.51
(1,84)	1:7:A:PRO:HG3	2:19:B:PHE:HD1	12	0.51
(1,84)	1:7:A:PRO:HG3	2:19:B:PHE:HD2	12	0.51
(1,79)	1:7:A:PRO:HA	2:19:B:PHE:HD1	4	0.51
(1,79)	1:7:A:PRO:HA	2:19:B:PHE:HD2	4	0.51
(1,68)	1:6:A:VAL:HG11	1:7:A:PRO:HG2	7	0.51
(1,68)	1:6:A:VAL:HG11	1:7:A:PRO:HG3	7	0.51
(1,68)	1:6:A:VAL:HG12	1:7:A:PRO:HG2	7	0.51

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,68)	1:6:A:VAL:HG12	1:7:A:PRO:HG3	7	0.51
(1,68)	1:6:A:VAL:HG13	1:7:A:PRO:HG2	7	0.51
(1,68)	1:6:A:VAL:HG13	1:7:A:PRO:HG3	7	0.51
(1,50)	1:6:A:VAL:HG21	2:20:B:TYR:HE1	17	0.51
(1,50)	1:6:A:VAL:HG21	2:20:B:TYR:HE2	17	0.51
(1,50)	1:6:A:VAL:HG22	2:20:B:TYR:HE1	17	0.51
(1,50)	1:6:A:VAL:HG22	2:20:B:TYR:HE2	17	0.51
(1,50)	1:6:A:VAL:HG23	2:20:B:TYR:HE1	17	0.51
(1,50)	1:6:A:VAL:HG23	2:20:B:TYR:HE2	17	0.51
(1,43)	1:7:A:PRO:HG2	2:19:B:PHE:HD1	5	0.51
(1,43)	1:7:A:PRO:HG2	2:19:B:PHE:HD2	5	0.51
(1,43)	1:7:A:PRO:HG3	2:19:B:PHE:HD1	5	0.51
(1,43)	1:7:A:PRO:HG3	2:19:B:PHE:HD2	5	0.51
(1,32)	2:16:B:PRO:HA	2:17:B:ILE:HD11	14	0.51
(1,32)	2:16:B:PRO:HA	2:17:B:ILE:HD12	14	0.51
(1,32)	2:16:B:PRO:HA	2:17:B:ILE:HD13	14	0.51
(1,16)	2:19:B:PHE:HD1	2:38:D:ILE:HG13	18	0.51
(1,16)	2:19:B:PHE:HD2	2:38:D:ILE:HG13	18	0.51
(1,1175)	1:24:C:ALA:HB1	2:40:D:PHE:HB3	1	0.5
(1,1175)	1:24:C:ALA:HB2	2:40:D:PHE:HB3	1	0.5
(1,1175)	1:24:C:ALA:HB3	2:40:D:PHE:HB3	1	0.5
(1,1167)	1:24:C:ALA:H	1:25:C:GLY:HA3	11	0.5
(1,1147)	1:28:C:PRO:HB2	1:31:C:PHE:HE1	17	0.5
(1,1147)	1:28:C:PRO:HB2	1:31:C:PHE:HE2	17	0.5
(1,1117)	1:27:C:VAL:HG11	1:28:C:PRO:HB3	8	0.5
(1,1117)	1:27:C:VAL:HG12	1:28:C:PRO:HB3	8	0.5
(1,1117)	1:27:C:VAL:HG13	1:28:C:PRO:HB3	8	0.5
(1,1099)	1:29:C:GLU:HG3	2:41:D:TYR:HD1	8	0.5
(1,1099)	1:29:C:GLU:HG3	2:41:D:TYR:HD2	8	0.5
(1,1073)	2:18:B:SER:HB3	2:38:D:ILE:HD11	7	0.5
(1,1073)	2:18:B:SER:HB3	2:38:D:ILE:HD12	7	0.5
(1,1073)	2:18:B:SER:HB3	2:38:D:ILE:HD13	7	0.5
(1,1057)	1:31:C:PHE:HD1	1:32:C:VAL:HA	11	0.5
(1,1057)	1:31:C:PHE:HD2	1:32:C:VAL:HA	11	0.5
(1,1054)	1:28:C:PRO:HA	2:40:D:PHE:HD1	7	0.5
(1,1054)	1:28:C:PRO:HA	2:40:D:PHE:HD2	7	0.5
(1,1051)	2:34:D:CYS:HB2	2:35:D:GLY:H	11	0.5
(1,1045)	1:27:C:VAL:HB	2:41:D:TYR:HD1	8	0.5
(1,1045)	1:27:C:VAL:HB	2:41:D:TYR:HD2	8	0.5
(1,1040)	1:31:C:PHE:HE1	2:37:D:PRO:HD2	9	0.5
(1,1040)	1:31:C:PHE:HE2	2:37:D:PRO:HD2	9	0.5
(1,1015)	1:30:C:TYR:HD1	2:39:D:SER:HA	2	0.5

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,1015)	1:30:C:TYR:HD2	2:39:D:SER:HA	2	0.5
(1,856)	1:24:C:ALA:HB1	2:40:D:PHE:HB3	1	0.5
(1,856)	1:24:C:ALA:HB2	2:40:D:PHE:HB3	1	0.5
(1,856)	1:24:C:ALA:HB3	2:40:D:PHE:HB3	1	0.5
(1,848)	1:24:C:ALA:H	1:25:C:GLY:HA3	11	0.5
(1,828)	1:28:C:PRO:HB2	1:31:C:PHE:HE1	17	0.5
(1,828)	1:28:C:PRO:HB2	1:31:C:PHE:HE2	17	0.5
(1,798)	1:27:C:VAL:HG11	1:28:C:PRO:HB3	8	0.5
(1,798)	1:27:C:VAL:HG12	1:28:C:PRO:HB3	8	0.5
(1,798)	1:27:C:VAL:HG13	1:28:C:PRO:HB3	8	0.5
(1,780)	1:29:C:GLU:HG3	2:41:D:TYR:HD1	8	0.5
(1,780)	1:29:C:GLU:HG3	2:41:D:TYR:HD2	8	0.5
(1,754)	2:18:B:SER:HB3	2:38:D:ILE:HD11	7	0.5
(1,754)	2:18:B:SER:HB3	2:38:D:ILE:HD12	7	0.5
(1,754)	2:18:B:SER:HB3	2:38:D:ILE:HD13	7	0.5
(1,738)	1:31:C:PHE:HD1	1:32:C:VAL:HA	11	0.5
(1,738)	1:31:C:PHE:HD2	1:32:C:VAL:HA	11	0.5
(1,735)	1:28:C:PRO:HA	2:40:D:PHE:HD1	7	0.5
(1,735)	1:28:C:PRO:HA	2:40:D:PHE:HD2	7	0.5
(1,732)	2:34:D:CYS:HB2	2:35:D:GLY:H	11	0.5
(1,726)	1:27:C:VAL:HB	2:41:D:TYR:HD1	8	0.5
(1,726)	1:27:C:VAL:HB	2:41:D:TYR:HD2	8	0.5
(1,721)	1:31:C:PHE:HE1	2:37:D:PRO:HD2	9	0.5
(1,721)	1:31:C:PHE:HE2	2:37:D:PRO:HD2	9	0.5
(1,696)	1:30:C:TYR:HD1	2:39:D:SER:HA	2	0.5
(1,696)	1:30:C:TYR:HD2	2:39:D:SER:HA	2	0.5
(1,519)	1:3:A:ALA:HB1	2:19:B:PHE:HB3	16	0.5
(1,519)	1:3:A:ALA:HB2	2:19:B:PHE:HB3	16	0.5
(1,519)	1:3:A:ALA:HB3	2:19:B:PHE:HB3	16	0.5
(1,481)	1:1:A:GLY:H	2:19:B:PHE:H	1	0.5
(1,474)	1:6:A:VAL:H	2:19:B:PHE:HB2	14	0.5
(1,461)	1:6:A:VAL:HG11	1:7:A:PRO:HB3	16	0.5
(1,461)	1:6:A:VAL:HG12	1:7:A:PRO:HB3	16	0.5
(1,461)	1:6:A:VAL:HG13	1:7:A:PRO:HB3	16	0.5
(1,445)	1:4:A:GLY:HA2	1:6:A:VAL:H	2	0.5
(1,439)	1:3:A:ALA:HA	1:4:A:GLY:H	19	0.5
(1,433)	1:8:A:GLU:HB2	2:20:B:TYR:HB2	13	0.5
(1,433)	1:8:A:GLU:HB2	2:20:B:TYR:HB3	13	0.5
(1,433)	1:8:A:GLU:HB2	2:20:B:TYR:HB2	18	0.5
(1,433)	1:8:A:GLU:HB2	2:20:B:TYR:HB3	18	0.5
(1,422)	1:4:A:GLY:H	2:20:B:TYR:H	14	0.5
(1,399)	1:7:A:PRO:HG2	1:10:A:PHE:HE1	20	0.5

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,399)	1:7:A:PRO:HG2	1:10:A:PHE:HE2	20	0.5
(1,399)	1:7:A:PRO:HG3	1:10:A:PHE:HE1	20	0.5
(1,399)	1:7:A:PRO:HG3	1:10:A:PHE:HE2	20	0.5
(1,379)	1:1:A:GLY:H	2:20:B:TYR:HE1	6	0.5
(1,379)	1:1:A:GLY:H	2:20:B:TYR:HE2	6	0.5
(1,369)	1:6:A:VAL:HG21	2:20:B:TYR:HE1	16	0.5
(1,369)	1:6:A:VAL:HG21	2:20:B:TYR:HE2	16	0.5
(1,369)	1:6:A:VAL:HG22	2:20:B:TYR:HE1	16	0.5
(1,369)	1:6:A:VAL:HG22	2:20:B:TYR:HE2	16	0.5
(1,369)	1:6:A:VAL:HG23	2:20:B:TYR:HE1	16	0.5
(1,369)	1:6:A:VAL:HG23	2:20:B:TYR:HE2	16	0.5
(1,200)	1:3:A:ALA:HB1	2:19:B:PHE:HB3	16	0.5
(1,200)	1:3:A:ALA:HB2	2:19:B:PHE:HB3	16	0.5
(1,200)	1:3:A:ALA:HB3	2:19:B:PHE:HB3	16	0.5
(1,162)	1:1:A:GLY:H	2:19:B:PHE:H	1	0.5
(1,155)	1:6:A:VAL:H	2:19:B:PHE:HB2	14	0.5
(1,142)	1:6:A:VAL:HG11	1:7:A:PRO:HB3	16	0.5
(1,142)	1:6:A:VAL:HG12	1:7:A:PRO:HB3	16	0.5
(1,142)	1:6:A:VAL:HG13	1:7:A:PRO:HB3	16	0.5
(1,126)	1:4:A:GLY:HA2	1:6:A:VAL:H	2	0.5
(1,120)	1:3:A:ALA:HA	1:4:A:GLY:H	19	0.5
(1,114)	1:8:A:GLU:HB2	2:20:B:TYR:HB2	13	0.5
(1,114)	1:8:A:GLU:HB2	2:20:B:TYR:HB3	13	0.5
(1,114)	1:8:A:GLU:HB2	2:20:B:TYR:HB2	18	0.5
(1,114)	1:8:A:GLU:HB2	2:20:B:TYR:HB3	18	0.5
(1,103)	1:4:A:GLY:H	2:20:B:TYR:H	14	0.5
(1,80)	1:7:A:PRO:HG2	1:10:A:PHE:HE1	20	0.5
(1,80)	1:7:A:PRO:HG2	1:10:A:PHE:HE2	20	0.5
(1,80)	1:7:A:PRO:HG3	1:10:A:PHE:HE1	20	0.5
(1,80)	1:7:A:PRO:HG3	1:10:A:PHE:HE2	20	0.5
(1,60)	1:1:A:GLY:H	2:20:B:TYR:HE1	6	0.5
(1,60)	1:1:A:GLY:H	2:20:B:TYR:HE2	6	0.5
(1,50)	1:6:A:VAL:HG21	2:20:B:TYR:HE1	16	0.5
(1,50)	1:6:A:VAL:HG21	2:20:B:TYR:HE2	16	0.5
(1,50)	1:6:A:VAL:HG22	2:20:B:TYR:HE1	16	0.5
(1,50)	1:6:A:VAL:HG22	2:20:B:TYR:HE2	16	0.5
(1,50)	1:6:A:VAL:HG23	2:20:B:TYR:HE1	16	0.5
(1,50)	1:6:A:VAL:HG23	2:20:B:TYR:HE2	16	0.5
(1,1309)	1:24:C:ALA:HA	2:40:D:PHE:HD1	1	0.49
(1,1309)	1:24:C:ALA:HA	2:40:D:PHE:HD2	1	0.49
(1,1287)	2:38:D:ILE:HA	2:38:D:ILE:HD11	3	0.49
(1,1287)	2:38:D:ILE:HA	2:38:D:ILE:HD12	3	0.49

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,1287)	2:38:D:ILE:HA	2:38:D:ILE:HD13	3	0.49
(1,1287)	2:38:D:ILE:HA	2:38:D:ILE:HD11	18	0.49
(1,1287)	2:38:D:ILE:HA	2:38:D:ILE:HD12	18	0.49
(1,1287)	2:38:D:ILE:HA	2:38:D:ILE:HD13	18	0.49
(1,1285)	2:38:D:ILE:H	2:38:D:ILE:HD11	4	0.49
(1,1285)	2:38:D:ILE:H	2:38:D:ILE:HD12	4	0.49
(1,1285)	2:38:D:ILE:H	2:38:D:ILE:HD13	4	0.49
(1,1175)	1:24:C:ALA:HB1	2:40:D:PHE:HB3	17	0.49
(1,1175)	1:24:C:ALA:HB2	2:40:D:PHE:HB3	17	0.49
(1,1175)	1:24:C:ALA:HB3	2:40:D:PHE:HB3	17	0.49
(1,1169)	2:36:D:THR:HB	2:38:D:ILE:HD11	7	0.49
(1,1169)	2:36:D:THR:HB	2:38:D:ILE:HD12	7	0.49
(1,1169)	2:36:D:THR:HB	2:38:D:ILE:HD13	7	0.49
(1,1105)	1:24:C:ALA:HB1	1:25:C:GLY:H	9	0.49
(1,1105)	1:24:C:ALA:HB2	1:25:C:GLY:H	9	0.49
(1,1105)	1:24:C:ALA:HB3	1:25:C:GLY:H	9	0.49
(1,1094)	1:29:C:GLU:HG2	2:40:D:PHE:HA	19	0.49
(1,1073)	2:18:B:SER:HB3	2:38:D:ILE:HD11	17	0.49
(1,1073)	2:18:B:SER:HB3	2:38:D:ILE:HD12	17	0.49
(1,1073)	2:18:B:SER:HB3	2:38:D:ILE:HD13	17	0.49
(1,1051)	2:34:D:CYS:HB2	2:35:D:GLY:H	18	0.49
(1,1047)	1:24:C:ALA:HA	2:40:D:PHE:HD1	1	0.49
(1,1047)	1:24:C:ALA:HA	2:40:D:PHE:HD2	1	0.49
(1,1045)	1:27:C:VAL:HB	2:41:D:TYR:HD1	3	0.49
(1,1045)	1:27:C:VAL:HB	2:41:D:TYR:HD2	3	0.49
(1,1045)	1:27:C:VAL:HB	2:41:D:TYR:HD1	16	0.49
(1,1045)	1:27:C:VAL:HB	2:41:D:TYR:HD2	16	0.49
(1,1040)	1:31:C:PHE:HE1	2:37:D:PRO:HD2	19	0.49
(1,1040)	1:31:C:PHE:HE2	2:37:D:PRO:HD2	19	0.49
(1,1034)	2:38:D:ILE:H	2:38:D:ILE:HG13	15	0.49
(1,1034)	2:38:D:ILE:H	2:38:D:ILE:HG13	20	0.49
(1,1026)	1:27:C:VAL:HG11	1:28:C:PRO:HD3	1	0.49
(1,1026)	1:27:C:VAL:HG12	1:28:C:PRO:HD3	1	0.49
(1,1026)	1:27:C:VAL:HG13	1:28:C:PRO:HD3	1	0.49
(1,1000)	1:31:C:PHE:HE1	2:37:D:PRO:HD2	3	0.49
(1,1000)	1:31:C:PHE:HE2	2:37:D:PRO:HD2	3	0.49
(1,990)	1:24:C:ALA:HA	2:40:D:PHE:HD1	1	0.49
(1,990)	1:24:C:ALA:HA	2:40:D:PHE:HD2	1	0.49
(1,968)	2:38:D:ILE:HA	2:38:D:ILE:HD11	3	0.49
(1,968)	2:38:D:ILE:HA	2:38:D:ILE:HD12	3	0.49
(1,968)	2:38:D:ILE:HA	2:38:D:ILE:HD13	3	0.49
(1,968)	2:38:D:ILE:HA	2:38:D:ILE:HD11	18	0.49

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,968)	2:38:D:ILE:HA	2:38:D:ILE:HD12	18	0.49
(1,968)	2:38:D:ILE:HA	2:38:D:ILE:HD13	18	0.49
(1,966)	2:38:D:ILE:H	2:38:D:ILE:HD11	4	0.49
(1,966)	2:38:D:ILE:H	2:38:D:ILE:HD12	4	0.49
(1,966)	2:38:D:ILE:H	2:38:D:ILE:HD13	4	0.49
(1,856)	1:24:C:ALA:HB1	2:40:D:PHE:HB3	17	0.49
(1,856)	1:24:C:ALA:HB2	2:40:D:PHE:HB3	17	0.49
(1,856)	1:24:C:ALA:HB3	2:40:D:PHE:HB3	17	0.49
(1,850)	2:36:D:THR:HB	2:38:D:ILE:HD11	7	0.49
(1,850)	2:36:D:THR:HB	2:38:D:ILE:HD12	7	0.49
(1,850)	2:36:D:THR:HB	2:38:D:ILE:HD13	7	0.49
(1,786)	1:24:C:ALA:HB1	1:25:C:GLY:H	9	0.49
(1,786)	1:24:C:ALA:HB2	1:25:C:GLY:H	9	0.49
(1,786)	1:24:C:ALA:HB3	1:25:C:GLY:H	9	0.49
(1,775)	1:29:C:GLU:HG2	2:40:D:PHE:HA	19	0.49
(1,754)	2:18:B:SER:HB3	2:38:D:ILE:HD11	17	0.49
(1,754)	2:18:B:SER:HB3	2:38:D:ILE:HD12	17	0.49
(1,754)	2:18:B:SER:HB3	2:38:D:ILE:HD13	17	0.49
(1,732)	2:34:D:CYS:HB2	2:35:D:GLY:H	18	0.49
(1,728)	1:24:C:ALA:HA	2:40:D:PHE:HD1	1	0.49
(1,728)	1:24:C:ALA:HA	2:40:D:PHE:HD2	1	0.49
(1,726)	1:27:C:VAL:HB	2:41:D:TYR:HD1	3	0.49
(1,726)	1:27:C:VAL:HB	2:41:D:TYR:HD2	3	0.49
(1,726)	1:27:C:VAL:HB	2:41:D:TYR:HD1	16	0.49
(1,726)	1:27:C:VAL:HB	2:41:D:TYR:HD2	16	0.49
(1,721)	1:31:C:PHE:HE1	2:37:D:PRO:HD2	19	0.49
(1,721)	1:31:C:PHE:HE2	2:37:D:PRO:HD2	19	0.49
(1,715)	2:38:D:ILE:H	2:38:D:ILE:HG13	15	0.49
(1,715)	2:38:D:ILE:H	2:38:D:ILE:HG13	20	0.49
(1,707)	1:27:C:VAL:HG11	1:28:C:PRO:HD3	1	0.49
(1,707)	1:27:C:VAL:HG12	1:28:C:PRO:HD3	1	0.49
(1,707)	1:27:C:VAL:HG13	1:28:C:PRO:HD3	1	0.49
(1,681)	1:31:C:PHE:HE1	2:37:D:PRO:HD2	3	0.49
(1,681)	1:31:C:PHE:HE2	2:37:D:PRO:HD2	3	0.49
(1,666)	2:17:B:ILE:HD11	2:40:D:PHE:HD1	16	0.49
(1,666)	2:17:B:ILE:HD11	2:40:D:PHE:HD2	16	0.49
(1,666)	2:17:B:ILE:HD12	2:40:D:PHE:HD1	16	0.49
(1,666)	2:17:B:ILE:HD12	2:40:D:PHE:HD2	16	0.49
(1,666)	2:17:B:ILE:HD13	2:40:D:PHE:HD1	16	0.49
(1,666)	2:17:B:ILE:HD13	2:40:D:PHE:HD2	16	0.49
(1,499)	1:6:A:VAL:HG11	1:7:A:PRO:HD3	8	0.49
(1,499)	1:6:A:VAL:HG12	1:7:A:PRO:HD3	8	0.49

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,499)	1:6:A:VAL:HG13	1:7:A:PRO:HD3	8	0.49
(1,485)	1:4:A:GLY:HA3	1:6:A:VAL:H	11	0.49
(1,467)	1:10:A:PHE:HB3	1:11:A:VAL:HG11	11	0.49
(1,467)	1:10:A:PHE:HB3	1:11:A:VAL:HG12	11	0.49
(1,467)	1:10:A:PHE:HB3	1:11:A:VAL:HG13	11	0.49
(1,467)	1:10:A:PHE:HB3	1:11:A:VAL:HG21	11	0.49
(1,467)	1:10:A:PHE:HB3	1:11:A:VAL:HG22	11	0.49
(1,467)	1:10:A:PHE:HB3	1:11:A:VAL:HG23	11	0.49
(1,461)	1:6:A:VAL:HG11	1:7:A:PRO:HB3	1	0.49
(1,461)	1:6:A:VAL:HG12	1:7:A:PRO:HB3	1	0.49
(1,461)	1:6:A:VAL:HG13	1:7:A:PRO:HB3	1	0.49
(1,445)	1:4:A:GLY:HA2	1:6:A:VAL:H	19	0.49
(1,407)	1:1:A:GLY:H	2:18:B:SER:HA	15	0.49
(1,395)	2:13:B:CYS:HB2	2:14:B:GLY:H	12	0.49
(1,386)	1:4:A:GLY:H	2:19:B:PHE:HD1	19	0.49
(1,386)	1:4:A:GLY:H	2:19:B:PHE:HD2	19	0.49
(1,180)	1:6:A:VAL:HG11	1:7:A:PRO:HD3	8	0.49
(1,180)	1:6:A:VAL:HG12	1:7:A:PRO:HD3	8	0.49
(1,180)	1:6:A:VAL:HG13	1:7:A:PRO:HD3	8	0.49
(1,166)	1:4:A:GLY:HA3	1:6:A:VAL:H	11	0.49
(1,148)	1:10:A:PHE:HB3	1:11:A:VAL:HG11	11	0.49
(1,148)	1:10:A:PHE:HB3	1:11:A:VAL:HG12	11	0.49
(1,148)	1:10:A:PHE:HB3	1:11:A:VAL:HG13	11	0.49
(1,148)	1:10:A:PHE:HB3	1:11:A:VAL:HG21	11	0.49
(1,148)	1:10:A:PHE:HB3	1:11:A:VAL:HG22	11	0.49
(1,148)	1:10:A:PHE:HB3	1:11:A:VAL:HG23	11	0.49
(1,142)	1:6:A:VAL:HG11	1:7:A:PRO:HB3	1	0.49
(1,142)	1:6:A:VAL:HG12	1:7:A:PRO:HB3	1	0.49
(1,142)	1:6:A:VAL:HG13	1:7:A:PRO:HB3	1	0.49
(1,126)	1:4:A:GLY:HA2	1:6:A:VAL:H	19	0.49
(1,88)	1:1:A:GLY:H	2:18:B:SER:HA	15	0.49
(1,76)	2:13:B:CYS:HB2	2:14:B:GLY:H	12	0.49
(1,67)	1:4:A:GLY:H	2:19:B:PHE:HD1	19	0.49
(1,67)	1:4:A:GLY:H	2:19:B:PHE:HD2	19	0.49
(1,1)	2:17:B:ILE:HD11	2:40:D:PHE:HD1	16	0.49
(1,1)	2:17:B:ILE:HD11	2:40:D:PHE:HD2	16	0.49
(1,1)	2:17:B:ILE:HD12	2:40:D:PHE:HD1	16	0.49
(1,1)	2:17:B:ILE:HD12	2:40:D:PHE:HD2	16	0.49
(1,1)	2:17:B:ILE:HD13	2:40:D:PHE:HD1	16	0.49
(1,1)	2:17:B:ILE:HD13	2:40:D:PHE:HD2	16	0.49
(1,1231)	1:31:C:PHE:HE1	2:36:D:THR:HA	5	0.48
(1,1231)	1:31:C:PHE:HE2	2:36:D:THR:HA	5	0.48

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,1130)	1:27:C:VAL:H	2:40:D:PHE:HB2	7	0.48
(1,1053)	2:39:D:SER:HB2	2:40:D:PHE:H	19	0.48
(1,1043)	1:27:C:VAL:HG11	1:28:C:PRO:HG2	11	0.48
(1,1043)	1:27:C:VAL:HG11	1:28:C:PRO:HG3	11	0.48
(1,1043)	1:27:C:VAL:HG12	1:28:C:PRO:HG2	11	0.48
(1,1043)	1:27:C:VAL:HG12	1:28:C:PRO:HG3	11	0.48
(1,1043)	1:27:C:VAL:HG13	1:28:C:PRO:HG2	11	0.48
(1,1043)	1:27:C:VAL:HG13	1:28:C:PRO:HG3	11	0.48
(1,1007)	2:37:D:PRO:HA	2:38:D:ILE:HD11	17	0.48
(1,1007)	2:37:D:PRO:HA	2:38:D:ILE:HD12	17	0.48
(1,1007)	2:37:D:PRO:HA	2:38:D:ILE:HD13	17	0.48
(1,1002)	1:30:C:TYR:HE1	2:39:D:SER:HB2	3	0.48
(1,1002)	1:30:C:TYR:HE2	2:39:D:SER:HB2	3	0.48
(1,912)	1:31:C:PHE:HE1	2:36:D:THR:HA	5	0.48
(1,912)	1:31:C:PHE:HE2	2:36:D:THR:HA	5	0.48
(1,811)	1:27:C:VAL:H	2:40:D:PHE:HB2	7	0.48
(1,734)	2:39:D:SER:HB2	2:40:D:PHE:H	19	0.48
(1,724)	1:27:C:VAL:HG11	1:28:C:PRO:HG2	11	0.48
(1,724)	1:27:C:VAL:HG11	1:28:C:PRO:HG3	11	0.48
(1,724)	1:27:C:VAL:HG12	1:28:C:PRO:HG2	11	0.48
(1,724)	1:27:C:VAL:HG12	1:28:C:PRO:HG3	11	0.48
(1,724)	1:27:C:VAL:HG13	1:28:C:PRO:HG2	11	0.48
(1,724)	1:27:C:VAL:HG13	1:28:C:PRO:HG3	11	0.48
(1,688)	2:37:D:PRO:HA	2:38:D:ILE:HD11	17	0.48
(1,688)	2:37:D:PRO:HA	2:38:D:ILE:HD12	17	0.48
(1,688)	2:37:D:PRO:HA	2:38:D:ILE:HD13	17	0.48
(1,683)	1:30:C:TYR:HE1	2:39:D:SER:HB2	3	0.48
(1,683)	1:30:C:TYR:HE2	2:39:D:SER:HB2	3	0.48
(1,515)	2:15:B:THR:HG21	2:17:B:ILE:HD11	7	0.48
(1,515)	2:15:B:THR:HG21	2:17:B:ILE:HD12	7	0.48
(1,515)	2:15:B:THR:HG21	2:17:B:ILE:HD13	7	0.48
(1,515)	2:15:B:THR:HG22	2:17:B:ILE:HD11	7	0.48
(1,515)	2:15:B:THR:HG22	2:17:B:ILE:HD12	7	0.48
(1,515)	2:15:B:THR:HG22	2:17:B:ILE:HD13	7	0.48
(1,515)	2:15:B:THR:HG23	2:17:B:ILE:HD11	7	0.48
(1,515)	2:15:B:THR:HG23	2:17:B:ILE:HD12	7	0.48
(1,515)	2:15:B:THR:HG23	2:17:B:ILE:HD13	7	0.48
(1,499)	1:6:A:VAL:HG11	1:7:A:PRO:HD3	17	0.48
(1,499)	1:6:A:VAL:HG12	1:7:A:PRO:HD3	17	0.48
(1,499)	1:6:A:VAL:HG13	1:7:A:PRO:HD3	17	0.48
(1,483)	1:1:A:GLY:H	2:20:B:TYR:HD1	18	0.48
(1,483)	1:1:A:GLY:H	2:20:B:TYR:HD2	18	0.48

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,452)	2:15:B:THR:HG21	2:16:B:PRO:HD3	3	0.48
(1,452)	2:15:B:THR:HG22	2:16:B:PRO:HD3	3	0.48
(1,452)	2:15:B:THR:HG23	2:16:B:PRO:HD3	3	0.48
(1,438)	1:8:A:GLU:HG2	2:19:B:PHE:HA	14	0.48
(1,387)	1:6:A:VAL:HG11	1:7:A:PRO:HG2	4	0.48
(1,387)	1:6:A:VAL:HG11	1:7:A:PRO:HG3	4	0.48
(1,387)	1:6:A:VAL:HG12	1:7:A:PRO:HG2	4	0.48
(1,387)	1:6:A:VAL:HG12	1:7:A:PRO:HG3	4	0.48
(1,387)	1:6:A:VAL:HG13	1:7:A:PRO:HG2	4	0.48
(1,387)	1:6:A:VAL:HG13	1:7:A:PRO:HG3	4	0.48
(1,344)	1:10:A:PHE:HE1	2:16:B:PRO:HD2	13	0.48
(1,344)	1:10:A:PHE:HE2	2:16:B:PRO:HD2	13	0.48
(1,196)	2:15:B:THR:HG21	2:17:B:ILE:HD11	7	0.48
(1,196)	2:15:B:THR:HG21	2:17:B:ILE:HD12	7	0.48
(1,196)	2:15:B:THR:HG21	2:17:B:ILE:HD13	7	0.48
(1,196)	2:15:B:THR:HG22	2:17:B:ILE:HD11	7	0.48
(1,196)	2:15:B:THR:HG22	2:17:B:ILE:HD12	7	0.48
(1,196)	2:15:B:THR:HG22	2:17:B:ILE:HD13	7	0.48
(1,196)	2:15:B:THR:HG23	2:17:B:ILE:HD11	7	0.48
(1,196)	2:15:B:THR:HG23	2:17:B:ILE:HD12	7	0.48
(1,196)	2:15:B:THR:HG23	2:17:B:ILE:HD13	7	0.48
(1,180)	1:6:A:VAL:HG11	1:7:A:PRO:HD3	17	0.48
(1,180)	1:6:A:VAL:HG12	1:7:A:PRO:HD3	17	0.48
(1,180)	1:6:A:VAL:HG13	1:7:A:PRO:HD3	17	0.48
(1,164)	1:1:A:GLY:H	2:20:B:TYR:HD1	18	0.48
(1,164)	1:1:A:GLY:H	2:20:B:TYR:HD2	18	0.48
(1,133)	2:15:B:THR:HG21	2:16:B:PRO:HD3	3	0.48
(1,133)	2:15:B:THR:HG22	2:16:B:PRO:HD3	3	0.48
(1,133)	2:15:B:THR:HG23	2:16:B:PRO:HD3	3	0.48
(1,119)	1:8:A:GLU:HG2	2:19:B:PHE:HA	14	0.48
(1,68)	1:6:A:VAL:HG11	1:7:A:PRO:HG2	4	0.48
(1,68)	1:6:A:VAL:HG11	1:7:A:PRO:HG3	4	0.48
(1,68)	1:6:A:VAL:HG12	1:7:A:PRO:HG2	4	0.48
(1,68)	1:6:A:VAL:HG12	1:7:A:PRO:HG3	4	0.48
(1,68)	1:6:A:VAL:HG13	1:7:A:PRO:HG2	4	0.48
(1,68)	1:6:A:VAL:HG13	1:7:A:PRO:HG3	4	0.48
(1,25)	1:10:A:PHE:HE1	2:16:B:PRO:HD2	13	0.48
(1,25)	1:10:A:PHE:HE2	2:16:B:PRO:HD2	13	0.48
(1,1175)	1:24:C:ALA:HB1	2:40:D:PHE:HB3	13	0.47
(1,1175)	1:24:C:ALA:HB2	2:40:D:PHE:HB3	13	0.47
(1,1175)	1:24:C:ALA:HB3	2:40:D:PHE:HB3	13	0.47
(1,1130)	1:27:C:VAL:H	2:40:D:PHE:HB2	6	0.47

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,1104)	1:25:C:GLY:H	2:40:D:PHE:H	11	0.47
(1,1094)	1:29:C:GLU:HG2	2:40:D:PHE:HA	18	0.47
(1,1074)	1:22:C:GLY:H	2:39:D:SER:HB3	13	0.47
(1,1003)	1:28:C:PRO:HD2	2:41:D:TYR:HD1	15	0.47
(1,1003)	1:28:C:PRO:HD2	2:41:D:TYR:HD2	15	0.47
(1,856)	1:24:C:ALA:HB1	2:40:D:PHE:HB3	13	0.47
(1,856)	1:24:C:ALA:HB2	2:40:D:PHE:HB3	13	0.47
(1,856)	1:24:C:ALA:HB3	2:40:D:PHE:HB3	13	0.47
(1,811)	1:27:C:VAL:H	2:40:D:PHE:HB2	6	0.47
(1,785)	1:25:C:GLY:H	2:40:D:PHE:H	11	0.47
(1,775)	1:29:C:GLU:HG2	2:40:D:PHE:HA	18	0.47
(1,755)	1:22:C:GLY:H	2:39:D:SER:HB3	13	0.47
(1,684)	1:28:C:PRO:HD2	2:41:D:TYR:HD1	15	0.47
(1,684)	1:28:C:PRO:HD2	2:41:D:TYR:HD2	15	0.47
(1,666)	2:17:B:ILE:HD11	2:40:D:PHE:HD1	3	0.47
(1,666)	2:17:B:ILE:HD11	2:40:D:PHE:HD2	3	0.47
(1,666)	2:17:B:ILE:HD12	2:40:D:PHE:HD1	3	0.47
(1,666)	2:17:B:ILE:HD12	2:40:D:PHE:HD2	3	0.47
(1,666)	2:17:B:ILE:HD13	2:40:D:PHE:HD1	3	0.47
(1,666)	2:17:B:ILE:HD13	2:40:D:PHE:HD2	3	0.47
(1,631)	2:17:B:ILE:HA	2:17:B:ILE:HD11	4	0.47
(1,631)	2:17:B:ILE:HA	2:17:B:ILE:HD12	4	0.47
(1,631)	2:17:B:ILE:HA	2:17:B:ILE:HD13	4	0.47
(1,438)	1:8:A:GLU:HG2	2:19:B:PHE:HA	1	0.47
(1,438)	1:8:A:GLU:HG2	2:19:B:PHE:HA	4	0.47
(1,438)	1:8:A:GLU:HG2	2:19:B:PHE:HA	12	0.47
(1,405)	1:1:A:GLY:H	2:18:B:SER:HB2	14	0.47
(1,403)	1:7:A:PRO:HG2	2:19:B:PHE:HD1	17	0.47
(1,403)	1:7:A:PRO:HG2	2:19:B:PHE:HD2	17	0.47
(1,403)	1:7:A:PRO:HG3	2:19:B:PHE:HD1	17	0.47
(1,403)	1:7:A:PRO:HG3	2:19:B:PHE:HD2	17	0.47
(1,393)	1:7:A:PRO:HG2	1:10:A:PHE:HD1	4	0.47
(1,393)	1:7:A:PRO:HG2	1:10:A:PHE:HD2	4	0.47
(1,393)	1:7:A:PRO:HG3	1:10:A:PHE:HD1	4	0.47
(1,393)	1:7:A:PRO:HG3	1:10:A:PHE:HD2	4	0.47
(1,389)	1:6:A:VAL:HB	2:20:B:TYR:HD1	20	0.47
(1,389)	1:6:A:VAL:HB	2:20:B:TYR:HD2	20	0.47
(1,359)	1:9:A:TYR:HD1	2:18:B:SER:HA	7	0.47
(1,359)	1:9:A:TYR:HD2	2:18:B:SER:HA	7	0.47
(1,312)	2:17:B:ILE:HA	2:17:B:ILE:HD11	4	0.47
(1,312)	2:17:B:ILE:HA	2:17:B:ILE:HD12	4	0.47
(1,312)	2:17:B:ILE:HA	2:17:B:ILE:HD13	4	0.47

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,119)	1:8:A:GLU:HG2	2:19:B:PHE:HA	1	0.47
(1,119)	1:8:A:GLU:HG2	2:19:B:PHE:HA	4	0.47
(1,119)	1:8:A:GLU:HG2	2:19:B:PHE:HA	12	0.47
(1,86)	1:1:A:GLY:H	2:18:B:SER:HB2	14	0.47
(1,84)	1:7:A:PRO:HG2	2:19:B:PHE:HD1	17	0.47
(1,84)	1:7:A:PRO:HG2	2:19:B:PHE:HD2	17	0.47
(1,84)	1:7:A:PRO:HG3	2:19:B:PHE:HD1	17	0.47
(1,84)	1:7:A:PRO:HG3	2:19:B:PHE:HD2	17	0.47
(1,74)	1:7:A:PRO:HG2	1:10:A:PHE:HD1	4	0.47
(1,74)	1:7:A:PRO:HG2	1:10:A:PHE:HD2	4	0.47
(1,74)	1:7:A:PRO:HG3	1:10:A:PHE:HD1	4	0.47
(1,74)	1:7:A:PRO:HG3	1:10:A:PHE:HD2	4	0.47
(1,70)	1:6:A:VAL:HB	2:20:B:TYR:HD1	20	0.47
(1,70)	1:6:A:VAL:HB	2:20:B:TYR:HD2	20	0.47
(1,40)	1:9:A:TYR:HD1	2:18:B:SER:HA	7	0.47
(1,40)	1:9:A:TYR:HD2	2:18:B:SER:HA	7	0.47
(1,1)	2:17:B:ILE:HD11	2:40:D:PHE:HD1	3	0.47
(1,1)	2:17:B:ILE:HD11	2:40:D:PHE:HD2	3	0.47
(1,1)	2:17:B:ILE:HD12	2:40:D:PHE:HD1	3	0.47
(1,1)	2:17:B:ILE:HD12	2:40:D:PHE:HD2	3	0.47
(1,1)	2:17:B:ILE:HD13	2:40:D:PHE:HD1	3	0.47
(1,1)	2:17:B:ILE:HD13	2:40:D:PHE:HD2	3	0.47
(1,1223)	1:31:C:PHE:HB3	1:32:C:VAL:H	18	0.46
(1,1157)	1:25:C:GLY:H	1:27:C:VAL:H	16	0.46
(1,1117)	1:27:C:VAL:HG11	1:28:C:PRO:HB3	19	0.46
(1,1117)	1:27:C:VAL:HG12	1:28:C:PRO:HB3	19	0.46
(1,1117)	1:27:C:VAL:HG13	1:28:C:PRO:HB3	19	0.46
(1,1102)	1:29:C:GLU:HB3	2:41:D:TYR:HD1	2	0.46
(1,1102)	1:29:C:GLU:HB3	2:41:D:TYR:HD2	2	0.46
(1,1051)	2:34:D:CYS:HB2	2:35:D:GLY:H	16	0.46
(1,1026)	1:27:C:VAL:HG11	1:28:C:PRO:HD3	16	0.46
(1,1026)	1:27:C:VAL:HG12	1:28:C:PRO:HD3	16	0.46
(1,1026)	1:27:C:VAL:HG13	1:28:C:PRO:HD3	16	0.46
(1,1020)	1:27:C:VAL:HB	2:41:D:TYR:H	3	0.46
(1,904)	1:31:C:PHE:HB3	1:32:C:VAL:H	18	0.46
(1,838)	1:25:C:GLY:H	1:27:C:VAL:H	16	0.46
(1,798)	1:27:C:VAL:HG11	1:28:C:PRO:HB3	19	0.46
(1,798)	1:27:C:VAL:HG12	1:28:C:PRO:HB3	19	0.46
(1,798)	1:27:C:VAL:HG13	1:28:C:PRO:HB3	19	0.46
(1,783)	1:29:C:GLU:HB3	2:41:D:TYR:HD1	2	0.46
(1,783)	1:29:C:GLU:HB3	2:41:D:TYR:HD2	2	0.46
(1,732)	2:34:D:CYS:HB2	2:35:D:GLY:H	16	0.46

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,707)	1:27:C:VAL:HG11	1:28:C:PRO:HD3	16	0.46
(1,707)	1:27:C:VAL:HG12	1:28:C:PRO:HD3	16	0.46
(1,707)	1:27:C:VAL:HG13	1:28:C:PRO:HD3	16	0.46
(1,701)	1:27:C:VAL:HB	2:41:D:TYR:H	3	0.46
(1,665)	2:19:B:PHE:HE1	2:38:D:ILE:HD11	2	0.46
(1,665)	2:19:B:PHE:HE1	2:38:D:ILE:HD12	2	0.46
(1,665)	2:19:B:PHE:HE1	2:38:D:ILE:HD13	2	0.46
(1,665)	2:19:B:PHE:HE2	2:38:D:ILE:HD11	2	0.46
(1,665)	2:19:B:PHE:HE2	2:38:D:ILE:HD12	2	0.46
(1,665)	2:19:B:PHE:HE2	2:38:D:ILE:HD13	2	0.46
(1,587)	1:12:A:ARG:H	1:12:A:ARG:HE	13	0.46
(1,575)	1:10:A:PHE:HE1	2:15:B:THR:HA	19	0.46
(1,575)	1:10:A:PHE:HE2	2:15:B:THR:HA	19	0.46
(1,519)	1:3:A:ALA:HB1	2:19:B:PHE:HB3	4	0.46
(1,519)	1:3:A:ALA:HB2	2:19:B:PHE:HB3	4	0.46
(1,519)	1:3:A:ALA:HB3	2:19:B:PHE:HB3	4	0.46
(1,519)	1:3:A:ALA:HB1	2:19:B:PHE:HB3	8	0.46
(1,519)	1:3:A:ALA:HB2	2:19:B:PHE:HB3	8	0.46
(1,519)	1:3:A:ALA:HB3	2:19:B:PHE:HB3	8	0.46
(1,443)	1:8:A:GLU:HG3	2:20:B:TYR:HD1	8	0.46
(1,443)	1:8:A:GLU:HG3	2:20:B:TYR:HD2	8	0.46
(1,443)	1:8:A:GLU:HG3	2:20:B:TYR:HD1	18	0.46
(1,443)	1:8:A:GLU:HG3	2:20:B:TYR:HD2	18	0.46
(1,438)	1:8:A:GLU:HG2	2:19:B:PHE:HA	2	0.46
(1,417)	2:17:B:ILE:HD11	2:39:D:SER:HB3	12	0.46
(1,417)	2:17:B:ILE:HD12	2:39:D:SER:HB3	12	0.46
(1,417)	2:17:B:ILE:HD13	2:39:D:SER:HB3	12	0.46
(1,399)	1:7:A:PRO:HG2	1:10:A:PHE:HE1	10	0.46
(1,399)	1:7:A:PRO:HG2	1:10:A:PHE:HE2	10	0.46
(1,399)	1:7:A:PRO:HG3	1:10:A:PHE:HE1	10	0.46
(1,399)	1:7:A:PRO:HG3	1:10:A:PHE:HE2	10	0.46
(1,351)	2:16:B:PRO:HA	2:17:B:ILE:HD11	16	0.46
(1,351)	2:16:B:PRO:HA	2:17:B:ILE:HD12	16	0.46
(1,351)	2:16:B:PRO:HA	2:17:B:ILE:HD13	16	0.46
(1,268)	1:12:A:ARG:H	1:12:A:ARG:HE	13	0.46
(1,256)	1:10:A:PHE:HE1	2:15:B:THR:HA	19	0.46
(1,256)	1:10:A:PHE:HE2	2:15:B:THR:HA	19	0.46
(1,200)	1:3:A:ALA:HB1	2:19:B:PHE:HB3	4	0.46
(1,200)	1:3:A:ALA:HB2	2:19:B:PHE:HB3	4	0.46
(1,200)	1:3:A:ALA:HB3	2:19:B:PHE:HB3	4	0.46
(1,200)	1:3:A:ALA:HB1	2:19:B:PHE:HB3	8	0.46
(1,200)	1:3:A:ALA:HB2	2:19:B:PHE:HB3	8	0.46

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,200)	1:3:A:ALA:HB3	2:19:B:PHE:HB3	8	0.46
(1,124)	1:8:A:GLU:HG3	2:20:B:TYR:HD1	8	0.46
(1,124)	1:8:A:GLU:HG3	2:20:B:TYR:HD2	8	0.46
(1,124)	1:8:A:GLU:HG3	2:20:B:TYR:HD1	18	0.46
(1,124)	1:8:A:GLU:HG3	2:20:B:TYR:HD2	18	0.46
(1,119)	1:8:A:GLU:HG2	2:19:B:PHE:HA	2	0.46
(1,98)	2:17:B:ILE:HD11	2:39:D:SER:HB3	12	0.46
(1,98)	2:17:B:ILE:HD12	2:39:D:SER:HB3	12	0.46
(1,98)	2:17:B:ILE:HD13	2:39:D:SER:HB3	12	0.46
(1,80)	1:7:A:PRO:HG2	1:10:A:PHE:HE1	10	0.46
(1,80)	1:7:A:PRO:HG2	1:10:A:PHE:HE2	10	0.46
(1,80)	1:7:A:PRO:HG3	1:10:A:PHE:HE1	10	0.46
(1,80)	1:7:A:PRO:HG3	1:10:A:PHE:HE2	10	0.46
(1,32)	2:16:B:PRO:HA	2:17:B:ILE:HD11	16	0.46
(1,32)	2:16:B:PRO:HA	2:17:B:ILE:HD12	16	0.46
(1,32)	2:16:B:PRO:HA	2:17:B:ILE:HD13	16	0.46
(1,18)	2:19:B:PHE:HE1	2:38:D:ILE:HD11	2	0.46
(1,18)	2:19:B:PHE:HE1	2:38:D:ILE:HD12	2	0.46
(1,18)	2:19:B:PHE:HE1	2:38:D:ILE:HD13	2	0.46
(1,18)	2:19:B:PHE:HE2	2:38:D:ILE:HD11	2	0.46
(1,18)	2:19:B:PHE:HE2	2:38:D:ILE:HD12	2	0.46
(1,18)	2:19:B:PHE:HE2	2:38:D:ILE:HD13	2	0.46
(1,1287)	2:38:D:ILE:HA	2:38:D:ILE:HD11	9	0.45
(1,1287)	2:38:D:ILE:HA	2:38:D:ILE:HD12	9	0.45
(1,1287)	2:38:D:ILE:HA	2:38:D:ILE:HD13	9	0.45
(1,1004)	1:32:C:VAL:HA	1:33:C:ARG:HD2	1	0.45
(1,1004)	1:32:C:VAL:HA	1:33:C:ARG:HD3	1	0.45
(1,968)	2:38:D:ILE:HA	2:38:D:ILE:HD11	9	0.45
(1,968)	2:38:D:ILE:HA	2:38:D:ILE:HD12	9	0.45
(1,968)	2:38:D:ILE:HA	2:38:D:ILE:HD13	9	0.45
(1,685)	1:32:C:VAL:HA	1:33:C:ARG:HD2	1	0.45
(1,685)	1:32:C:VAL:HA	1:33:C:ARG:HD3	1	0.45
(1,596)	1:7:A:PRO:HG2	1:10:A:PHE:HE1	20	0.45
(1,596)	1:7:A:PRO:HG2	1:10:A:PHE:HE2	20	0.45
(1,596)	1:7:A:PRO:HG3	1:10:A:PHE:HE1	20	0.45
(1,596)	1:7:A:PRO:HG3	1:10:A:PHE:HE2	20	0.45
(1,519)	1:3:A:ALA:HB1	2:19:B:PHE:HB3	20	0.45
(1,519)	1:3:A:ALA:HB2	2:19:B:PHE:HB3	20	0.45
(1,519)	1:3:A:ALA:HB3	2:19:B:PHE:HB3	20	0.45
(1,515)	2:15:B:THR:HG21	2:17:B:ILE:HD11	5	0.45
(1,515)	2:15:B:THR:HG21	2:17:B:ILE:HD12	5	0.45
(1,515)	2:15:B:THR:HG21	2:17:B:ILE:HD13	5	0.45

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,515)	2:15:B:THR:HG22	2:17:B:ILE:HD11	5	0.45
(1,515)	2:15:B:THR:HG22	2:17:B:ILE:HD12	5	0.45
(1,515)	2:15:B:THR:HG22	2:17:B:ILE:HD13	5	0.45
(1,515)	2:15:B:THR:HG23	2:17:B:ILE:HD11	5	0.45
(1,515)	2:15:B:THR:HG23	2:17:B:ILE:HD12	5	0.45
(1,515)	2:15:B:THR:HG23	2:17:B:ILE:HD13	5	0.45
(1,514)	1:10:A:PHE:HB2	1:11:A:VAL:HG11	1	0.45
(1,514)	1:10:A:PHE:HB2	1:11:A:VAL:HG12	1	0.45
(1,514)	1:10:A:PHE:HB2	1:11:A:VAL:HG13	1	0.45
(1,514)	1:10:A:PHE:HB2	1:11:A:VAL:HG21	1	0.45
(1,514)	1:10:A:PHE:HB2	1:11:A:VAL:HG22	1	0.45
(1,514)	1:10:A:PHE:HB2	1:11:A:VAL:HG23	1	0.45
(1,514)	1:10:A:PHE:HB2	1:11:A:VAL:HG11	15	0.45
(1,514)	1:10:A:PHE:HB2	1:11:A:VAL:HG12	15	0.45
(1,514)	1:10:A:PHE:HB2	1:11:A:VAL:HG13	15	0.45
(1,514)	1:10:A:PHE:HB2	1:11:A:VAL:HG21	15	0.45
(1,514)	1:10:A:PHE:HB2	1:11:A:VAL:HG22	15	0.45
(1,514)	1:10:A:PHE:HB2	1:11:A:VAL:HG23	15	0.45
(1,503)	1:7:A:PRO:HB3	1:10:A:PHE:HE1	9	0.45
(1,503)	1:7:A:PRO:HB3	1:10:A:PHE:HE2	9	0.45
(1,483)	1:1:A:GLY:H	2:20:B:TYR:HD1	2	0.45
(1,483)	1:1:A:GLY:H	2:20:B:TYR:HD2	2	0.45
(1,474)	1:6:A:VAL:H	2:19:B:PHE:HB2	4	0.45
(1,461)	1:6:A:VAL:HG11	1:7:A:PRO:HB3	7	0.45
(1,461)	1:6:A:VAL:HG12	1:7:A:PRO:HB3	7	0.45
(1,461)	1:6:A:VAL:HG13	1:7:A:PRO:HB3	7	0.45
(1,437)	1:8:A:GLU:HG2	2:20:B:TYR:HB2	1	0.45
(1,437)	1:8:A:GLU:HG2	2:20:B:TYR:HB3	1	0.45
(1,342)	1:1:A:GLY:HA3	2:18:B:SER:HB2	4	0.45
(1,339)	1:1:A:GLY:HA3	2:18:B:SER:HB2	4	0.45
(1,277)	1:7:A:PRO:HG2	1:10:A:PHE:HE1	20	0.45
(1,277)	1:7:A:PRO:HG2	1:10:A:PHE:HE2	20	0.45
(1,277)	1:7:A:PRO:HG3	1:10:A:PHE:HE1	20	0.45
(1,277)	1:7:A:PRO:HG3	1:10:A:PHE:HE2	20	0.45
(1,200)	1:3:A:ALA:HB1	2:19:B:PHE:HB3	20	0.45
(1,200)	1:3:A:ALA:HB2	2:19:B:PHE:HB3	20	0.45
(1,200)	1:3:A:ALA:HB3	2:19:B:PHE:HB3	20	0.45
(1,196)	2:15:B:THR:HG21	2:17:B:ILE:HD11	5	0.45
(1,196)	2:15:B:THR:HG21	2:17:B:ILE:HD12	5	0.45
(1,196)	2:15:B:THR:HG21	2:17:B:ILE:HD13	5	0.45
(1,196)	2:15:B:THR:HG22	2:17:B:ILE:HD11	5	0.45
(1,196)	2:15:B:THR:HG22	2:17:B:ILE:HD12	5	0.45

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,196)	2:15:B:THR:HG22	2:17:B:ILE:HD13	5	0.45
(1,196)	2:15:B:THR:HG23	2:17:B:ILE:HD11	5	0.45
(1,196)	2:15:B:THR:HG23	2:17:B:ILE:HD12	5	0.45
(1,196)	2:15:B:THR:HG23	2:17:B:ILE:HD13	5	0.45
(1,195)	1:10:A:PHE:HB2	1:11:A:VAL:HG11	1	0.45
(1,195)	1:10:A:PHE:HB2	1:11:A:VAL:HG12	1	0.45
(1,195)	1:10:A:PHE:HB2	1:11:A:VAL:HG13	1	0.45
(1,195)	1:10:A:PHE:HB2	1:11:A:VAL:HG21	1	0.45
(1,195)	1:10:A:PHE:HB2	1:11:A:VAL:HG22	1	0.45
(1,195)	1:10:A:PHE:HB2	1:11:A:VAL:HG23	1	0.45
(1,195)	1:10:A:PHE:HB2	1:11:A:VAL:HG11	15	0.45
(1,195)	1:10:A:PHE:HB2	1:11:A:VAL:HG12	15	0.45
(1,195)	1:10:A:PHE:HB2	1:11:A:VAL:HG13	15	0.45
(1,195)	1:10:A:PHE:HB2	1:11:A:VAL:HG21	15	0.45
(1,195)	1:10:A:PHE:HB2	1:11:A:VAL:HG22	15	0.45
(1,195)	1:10:A:PHE:HB2	1:11:A:VAL:HG23	15	0.45
(1,184)	1:7:A:PRO:HB3	1:10:A:PHE:HE1	9	0.45
(1,184)	1:7:A:PRO:HB3	1:10:A:PHE:HE2	9	0.45
(1,164)	1:1:A:GLY:H	2:20:B:TYR:HD1	2	0.45
(1,164)	1:1:A:GLY:H	2:20:B:TYR:HD2	2	0.45
(1,155)	1:6:A:VAL:H	2:19:B:PHE:HB2	4	0.45
(1,142)	1:6:A:VAL:HG11	1:7:A:PRO:HB3	7	0.45
(1,142)	1:6:A:VAL:HG12	1:7:A:PRO:HB3	7	0.45
(1,142)	1:6:A:VAL:HG13	1:7:A:PRO:HB3	7	0.45
(1,118)	1:8:A:GLU:HG2	2:20:B:TYR:HB2	1	0.45
(1,118)	1:8:A:GLU:HG2	2:20:B:TYR:HB3	1	0.45
(1,23)	1:1:A:GLY:HA3	2:18:B:SER:HB2	4	0.45
(1,20)	1:1:A:GLY:HA3	2:18:B:SER:HB2	4	0.45
(1,1175)	1:24:C:ALA:HB1	2:40:D:PHE:HB3	18	0.44
(1,1175)	1:24:C:ALA:HB2	2:40:D:PHE:HB3	18	0.44
(1,1175)	1:24:C:ALA:HB3	2:40:D:PHE:HB3	18	0.44
(1,1155)	1:27:C:VAL:HG11	1:28:C:PRO:HD3	12	0.44
(1,1155)	1:27:C:VAL:HG12	1:28:C:PRO:HD3	12	0.44
(1,1155)	1:27:C:VAL:HG13	1:28:C:PRO:HD3	12	0.44
(1,1141)	1:25:C:GLY:HA3	1:27:C:VAL:H	7	0.44
(1,1095)	1:24:C:ALA:HA	1:25:C:GLY:H	1	0.44
(1,1089)	1:29:C:GLU:HB2	2:41:D:TYR:HB2	12	0.44
(1,1089)	1:29:C:GLU:HB2	2:41:D:TYR:HB3	12	0.44
(1,1078)	1:25:C:GLY:H	2:41:D:TYR:H	3	0.44
(1,1053)	2:39:D:SER:HB2	2:40:D:PHE:H	4	0.44
(1,1053)	2:39:D:SER:HB2	2:40:D:PHE:H	7	0.44
(1,1043)	1:27:C:VAL:HG11	1:28:C:PRO:HG2	3	0.44

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,1043)	1:27:C:VAL:HG11	1:28:C:PRO:HG3	3	0.44
(1,1043)	1:27:C:VAL:HG12	1:28:C:PRO:HG2	3	0.44
(1,1043)	1:27:C:VAL:HG12	1:28:C:PRO:HG3	3	0.44
(1,1043)	1:27:C:VAL:HG13	1:28:C:PRO:HG2	3	0.44
(1,1043)	1:27:C:VAL:HG13	1:28:C:PRO:HG3	3	0.44
(1,1042)	1:25:C:GLY:H	2:40:D:PHE:HD1	11	0.44
(1,1042)	1:25:C:GLY:H	2:40:D:PHE:HD2	11	0.44
(1,856)	1:24:C:ALA:HB1	2:40:D:PHE:HB3	18	0.44
(1,856)	1:24:C:ALA:HB2	2:40:D:PHE:HB3	18	0.44
(1,856)	1:24:C:ALA:HB3	2:40:D:PHE:HB3	18	0.44
(1,836)	1:27:C:VAL:HG11	1:28:C:PRO:HD3	12	0.44
(1,836)	1:27:C:VAL:HG12	1:28:C:PRO:HD3	12	0.44
(1,836)	1:27:C:VAL:HG13	1:28:C:PRO:HD3	12	0.44
(1,822)	1:25:C:GLY:HA3	1:27:C:VAL:H	7	0.44
(1,776)	1:24:C:ALA:HA	1:25:C:GLY:H	1	0.44
(1,770)	1:29:C:GLU:HB2	2:41:D:TYR:HB2	12	0.44
(1,770)	1:29:C:GLU:HB2	2:41:D:TYR:HB3	12	0.44
(1,759)	1:25:C:GLY:H	2:41:D:TYR:H	3	0.44
(1,734)	2:39:D:SER:HB2	2:40:D:PHE:H	4	0.44
(1,734)	2:39:D:SER:HB2	2:40:D:PHE:H	7	0.44
(1,724)	1:27:C:VAL:HG11	1:28:C:PRO:HG2	3	0.44
(1,724)	1:27:C:VAL:HG11	1:28:C:PRO:HG3	3	0.44
(1,724)	1:27:C:VAL:HG12	1:28:C:PRO:HG2	3	0.44
(1,724)	1:27:C:VAL:HG12	1:28:C:PRO:HG3	3	0.44
(1,724)	1:27:C:VAL:HG13	1:28:C:PRO:HG2	3	0.44
(1,724)	1:27:C:VAL:HG13	1:28:C:PRO:HG3	3	0.44
(1,723)	1:25:C:GLY:H	2:40:D:PHE:HD1	11	0.44
(1,723)	1:25:C:GLY:H	2:40:D:PHE:HD2	11	0.44
(1,661)	2:19:B:PHE:H	2:38:D:ILE:HD11	12	0.44
(1,661)	2:19:B:PHE:H	2:38:D:ILE:HD12	12	0.44
(1,661)	2:19:B:PHE:H	2:38:D:ILE:HD13	12	0.44
(1,568)	1:10:A:PHE:H	1:10:A:PHE:HD1	12	0.44
(1,568)	1:10:A:PHE:H	1:10:A:PHE:HD2	12	0.44
(1,517)	2:15:B:THR:HG21	2:40:D:PHE:HE1	15	0.44
(1,517)	2:15:B:THR:HG21	2:40:D:PHE:HE2	15	0.44
(1,517)	2:15:B:THR:HG22	2:40:D:PHE:HE1	15	0.44
(1,517)	2:15:B:THR:HG22	2:40:D:PHE:HE2	15	0.44
(1,517)	2:15:B:THR:HG23	2:40:D:PHE:HE1	15	0.44
(1,517)	2:15:B:THR:HG23	2:40:D:PHE:HE2	15	0.44
(1,501)	1:4:A:GLY:H	1:6:A:VAL:H	8	0.44
(1,461)	1:6:A:VAL:HG11	1:7:A:PRO:HB3	9	0.44
(1,461)	1:6:A:VAL:HG12	1:7:A:PRO:HB3	9	0.44

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,461)	1:6:A:VAL:HG13	1:7:A:PRO:HB3	9	0.44
(1,445)	1:4:A:GLY:HA2	1:6:A:VAL:H	12	0.44
(1,441)	1:8:A:GLU:HG3	2:20:B:TYR:HB2	10	0.44
(1,441)	1:8:A:GLU:HG3	2:20:B:TYR:HB3	10	0.44
(1,441)	1:8:A:GLU:HG3	2:20:B:TYR:HB2	20	0.44
(1,441)	1:8:A:GLU:HG3	2:20:B:TYR:HB3	20	0.44
(1,422)	1:4:A:GLY:H	2:20:B:TYR:H	10	0.44
(1,407)	1:1:A:GLY:H	2:18:B:SER:HA	13	0.44
(1,403)	1:7:A:PRO:HG2	2:19:B:PHE:HD1	6	0.44
(1,403)	1:7:A:PRO:HG2	2:19:B:PHE:HD2	6	0.44
(1,403)	1:7:A:PRO:HG3	2:19:B:PHE:HD1	6	0.44
(1,403)	1:7:A:PRO:HG3	2:19:B:PHE:HD2	6	0.44
(1,402)	1:10:A:PHE:HD1	2:14:B:GLY:HA3	9	0.44
(1,402)	1:10:A:PHE:HD2	2:14:B:GLY:HA3	9	0.44
(1,398)	1:7:A:PRO:HA	2:19:B:PHE:HD1	8	0.44
(1,398)	1:7:A:PRO:HA	2:19:B:PHE:HD2	8	0.44
(1,397)	2:18:B:SER:HB2	2:19:B:PHE:H	15	0.44
(1,378)	2:17:B:ILE:H	2:17:B:ILE:HG13	15	0.44
(1,345)	1:9:A:TYR:HD1	2:18:B:SER:HB2	13	0.44
(1,345)	1:9:A:TYR:HD2	2:18:B:SER:HB2	13	0.44
(1,249)	1:10:A:PHE:H	1:10:A:PHE:HD1	12	0.44
(1,249)	1:10:A:PHE:H	1:10:A:PHE:HD2	12	0.44
(1,198)	2:15:B:THR:HG21	2:40:D:PHE:HE1	15	0.44
(1,198)	2:15:B:THR:HG21	2:40:D:PHE:HE2	15	0.44
(1,198)	2:15:B:THR:HG22	2:40:D:PHE:HE1	15	0.44
(1,198)	2:15:B:THR:HG22	2:40:D:PHE:HE2	15	0.44
(1,198)	2:15:B:THR:HG23	2:40:D:PHE:HE1	15	0.44
(1,198)	2:15:B:THR:HG23	2:40:D:PHE:HE2	15	0.44
(1,182)	1:4:A:GLY:H	1:6:A:VAL:H	8	0.44
(1,142)	1:6:A:VAL:HG11	1:7:A:PRO:HB3	9	0.44
(1,142)	1:6:A:VAL:HG12	1:7:A:PRO:HB3	9	0.44
(1,142)	1:6:A:VAL:HG13	1:7:A:PRO:HB3	9	0.44
(1,126)	1:4:A:GLY:HA2	1:6:A:VAL:H	12	0.44
(1,122)	1:8:A:GLU:HG3	2:20:B:TYR:HB2	10	0.44
(1,122)	1:8:A:GLU:HG3	2:20:B:TYR:HB3	10	0.44
(1,122)	1:8:A:GLU:HG3	2:20:B:TYR:HB2	20	0.44
(1,122)	1:8:A:GLU:HG3	2:20:B:TYR:HB3	20	0.44
(1,103)	1:4:A:GLY:H	2:20:B:TYR:H	10	0.44
(1,88)	1:1:A:GLY:H	2:18:B:SER:HA	13	0.44
(1,84)	1:7:A:PRO:HG2	2:19:B:PHE:HD1	6	0.44
(1,84)	1:7:A:PRO:HG2	2:19:B:PHE:HD2	6	0.44
(1,84)	1:7:A:PRO:HG3	2:19:B:PHE:HD1	6	0.44

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,84)	1:7:A:PRO:HG3	2:19:B:PHE:HD2	6	0.44
(1,83)	1:10:A:PHE:HD1	2:14:B:GLY:HA3	9	0.44
(1,83)	1:10:A:PHE:HD2	2:14:B:GLY:HA3	9	0.44
(1,79)	1:7:A:PRO:HA	2:19:B:PHE:HD1	8	0.44
(1,79)	1:7:A:PRO:HA	2:19:B:PHE:HD2	8	0.44
(1,78)	2:18:B:SER:HB2	2:19:B:PHE:H	15	0.44
(1,59)	2:17:B:ILE:H	2:17:B:ILE:HG13	15	0.44
(1,26)	1:9:A:TYR:HD1	2:18:B:SER:HB2	13	0.44
(1,26)	1:9:A:TYR:HD2	2:18:B:SER:HB2	13	0.44
(1,14)	2:19:B:PHE:H	2:38:D:ILE:HD11	12	0.44
(1,14)	2:19:B:PHE:H	2:38:D:ILE:HD12	12	0.44
(1,14)	2:19:B:PHE:H	2:38:D:ILE:HD13	12	0.44
(1,1243)	1:33:C:ARG:H	1:33:C:ARG:HE	12	0.43
(1,1163)	1:23:C:GLY:HA2	1:24:C:ALA:HB1	18	0.43
(1,1163)	1:23:C:GLY:HA2	1:24:C:ALA:HB2	18	0.43
(1,1163)	1:23:C:GLY:HA2	1:24:C:ALA:HB3	18	0.43
(1,1159)	1:28:C:PRO:HB3	1:31:C:PHE:HE1	14	0.43
(1,1159)	1:28:C:PRO:HB3	1:31:C:PHE:HE2	14	0.43
(1,1157)	1:25:C:GLY:H	1:27:C:VAL:H	1	0.43
(1,1125)	1:31:C:PHE:HD1	1:32:C:VAL:HG11	8	0.43
(1,1125)	1:31:C:PHE:HD1	1:32:C:VAL:HG12	8	0.43
(1,1125)	1:31:C:PHE:HD1	1:32:C:VAL:HG13	8	0.43
(1,1125)	1:31:C:PHE:HD1	1:32:C:VAL:HG21	8	0.43
(1,1125)	1:31:C:PHE:HD1	1:32:C:VAL:HG22	8	0.43
(1,1125)	1:31:C:PHE:HD1	1:32:C:VAL:HG23	8	0.43
(1,1125)	1:31:C:PHE:HD2	1:32:C:VAL:HG11	8	0.43
(1,1125)	1:31:C:PHE:HD2	1:32:C:VAL:HG12	8	0.43
(1,1125)	1:31:C:PHE:HD2	1:32:C:VAL:HG13	8	0.43
(1,1125)	1:31:C:PHE:HD2	1:32:C:VAL:HG21	8	0.43
(1,1125)	1:31:C:PHE:HD2	1:32:C:VAL:HG22	8	0.43
(1,1125)	1:31:C:PHE:HD2	1:32:C:VAL:HG23	8	0.43
(1,1109)	1:31:C:PHE:HD1	2:36:D:THR:HG21	16	0.43
(1,1109)	1:31:C:PHE:HD1	2:36:D:THR:HG22	16	0.43
(1,1109)	1:31:C:PHE:HD1	2:36:D:THR:HG23	16	0.43
(1,1109)	1:31:C:PHE:HD2	2:36:D:THR:HG21	16	0.43
(1,1109)	1:31:C:PHE:HD2	2:36:D:THR:HG22	16	0.43
(1,1109)	1:31:C:PHE:HD2	2:36:D:THR:HG23	16	0.43
(1,1084)	1:25:C:GLY:H	2:40:D:PHE:HB3	4	0.43
(1,1050)	1:28:C:PRO:HG2	1:31:C:PHE:HB2	18	0.43
(1,1050)	1:28:C:PRO:HG3	1:31:C:PHE:HB2	18	0.43
(1,1040)	1:31:C:PHE:HE1	2:37:D:PRO:HD2	5	0.43
(1,1040)	1:31:C:PHE:HE2	2:37:D:PRO:HD2	5	0.43

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,1026)	1:27:C:VAL:HG11	1:28:C:PRO:HD3	15	0.43
(1,1026)	1:27:C:VAL:HG12	1:28:C:PRO:HD3	15	0.43
(1,1026)	1:27:C:VAL:HG13	1:28:C:PRO:HD3	15	0.43
(1,1016)	1:28:C:PRO:HG2	1:31:C:PHE:HB2	18	0.43
(1,1016)	1:28:C:PRO:HG3	1:31:C:PHE:HB2	18	0.43
(1,1008)	2:18:B:SER:HA	2:38:D:ILE:HG13	14	0.43
(1,1005)	2:34:D:CYS:HB3	2:35:D:GLY:HA3	9	0.43
(1,924)	1:33:C:ARG:H	1:33:C:ARG:HE	12	0.43
(1,844)	1:23:C:GLY:HA2	1:24:C:ALA:HB1	18	0.43
(1,844)	1:23:C:GLY:HA2	1:24:C:ALA:HB2	18	0.43
(1,844)	1:23:C:GLY:HA2	1:24:C:ALA:HB3	18	0.43
(1,840)	1:28:C:PRO:HB3	1:31:C:PHE:HE1	14	0.43
(1,840)	1:28:C:PRO:HB3	1:31:C:PHE:HE2	14	0.43
(1,838)	1:25:C:GLY:H	1:27:C:VAL:H	1	0.43
(1,806)	1:31:C:PHE:HD1	1:32:C:VAL:HG11	8	0.43
(1,806)	1:31:C:PHE:HD1	1:32:C:VAL:HG12	8	0.43
(1,806)	1:31:C:PHE:HD1	1:32:C:VAL:HG13	8	0.43
(1,806)	1:31:C:PHE:HD1	1:32:C:VAL:HG21	8	0.43
(1,806)	1:31:C:PHE:HD1	1:32:C:VAL:HG22	8	0.43
(1,806)	1:31:C:PHE:HD1	1:32:C:VAL:HG23	8	0.43
(1,806)	1:31:C:PHE:HD2	1:32:C:VAL:HG11	8	0.43
(1,806)	1:31:C:PHE:HD2	1:32:C:VAL:HG12	8	0.43
(1,806)	1:31:C:PHE:HD2	1:32:C:VAL:HG13	8	0.43
(1,806)	1:31:C:PHE:HD2	1:32:C:VAL:HG21	8	0.43
(1,806)	1:31:C:PHE:HD2	1:32:C:VAL:HG22	8	0.43
(1,806)	1:31:C:PHE:HD2	1:32:C:VAL:HG23	8	0.43
(1,790)	1:31:C:PHE:HD1	2:36:D:THR:HG21	16	0.43
(1,790)	1:31:C:PHE:HD1	2:36:D:THR:HG22	16	0.43
(1,790)	1:31:C:PHE:HD1	2:36:D:THR:HG23	16	0.43
(1,790)	1:31:C:PHE:HD2	2:36:D:THR:HG21	16	0.43
(1,790)	1:31:C:PHE:HD2	2:36:D:THR:HG22	16	0.43
(1,790)	1:31:C:PHE:HD2	2:36:D:THR:HG23	16	0.43
(1,765)	1:25:C:GLY:H	2:40:D:PHE:HB3	4	0.43
(1,731)	1:28:C:PRO:HG2	1:31:C:PHE:HB2	18	0.43
(1,731)	1:28:C:PRO:HG3	1:31:C:PHE:HB2	18	0.43
(1,721)	1:31:C:PHE:HE1	2:37:D:PRO:HD2	5	0.43
(1,721)	1:31:C:PHE:HE2	2:37:D:PRO:HD2	5	0.43
(1,707)	1:27:C:VAL:HG11	1:28:C:PRO:HD3	15	0.43
(1,707)	1:27:C:VAL:HG12	1:28:C:PRO:HD3	15	0.43
(1,707)	1:27:C:VAL:HG13	1:28:C:PRO:HD3	15	0.43
(1,697)	1:28:C:PRO:HG2	1:31:C:PHE:HB2	18	0.43
(1,697)	1:28:C:PRO:HG3	1:31:C:PHE:HB2	18	0.43

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,689)	2:18:B:SER:HA	2:38:D:ILE:HG13	14	0.43
(1,686)	2:34:D:CYS:HB3	2:35:D:GLY:HA3	9	0.43
(1,442)	1:8:A:GLU:HG3	2:19:B:PHE:HA	17	0.43
(1,436)	1:8:A:GLU:HG2	2:20:B:TYR:HD1	14	0.43
(1,436)	1:8:A:GLU:HG2	2:20:B:TYR:HD2	14	0.43
(1,431)	1:7:A:PRO:HB3	1:10:A:PHE:HD1	16	0.43
(1,431)	1:7:A:PRO:HB3	1:10:A:PHE:HD2	16	0.43
(1,420)	1:4:A:GLY:HA3	1:5:A:HIS:HD2	9	0.43
(1,387)	1:6:A:VAL:HG11	1:7:A:PRO:HG2	3	0.43
(1,387)	1:6:A:VAL:HG11	1:7:A:PRO:HG3	3	0.43
(1,387)	1:6:A:VAL:HG12	1:7:A:PRO:HG2	3	0.43
(1,387)	1:6:A:VAL:HG12	1:7:A:PRO:HG3	3	0.43
(1,387)	1:6:A:VAL:HG13	1:7:A:PRO:HG2	3	0.43
(1,387)	1:6:A:VAL:HG13	1:7:A:PRO:HG3	3	0.43
(1,387)	1:6:A:VAL:HG11	1:7:A:PRO:HG2	12	0.43
(1,387)	1:6:A:VAL:HG11	1:7:A:PRO:HG3	12	0.43
(1,387)	1:6:A:VAL:HG12	1:7:A:PRO:HG2	12	0.43
(1,387)	1:6:A:VAL:HG12	1:7:A:PRO:HG3	12	0.43
(1,387)	1:6:A:VAL:HG13	1:7:A:PRO:HG2	12	0.43
(1,387)	1:6:A:VAL:HG13	1:7:A:PRO:HG3	12	0.43
(1,346)	1:9:A:TYR:HE1	2:18:B:SER:HB2	19	0.43
(1,346)	1:9:A:TYR:HE2	2:18:B:SER:HB2	19	0.43
(1,123)	1:8:A:GLU:HG3	2:19:B:PHE:HA	17	0.43
(1,117)	1:8:A:GLU:HG2	2:20:B:TYR:HD1	14	0.43
(1,117)	1:8:A:GLU:HG2	2:20:B:TYR:HD2	14	0.43
(1,112)	1:7:A:PRO:HB3	1:10:A:PHE:HD1	16	0.43
(1,112)	1:7:A:PRO:HB3	1:10:A:PHE:HD2	16	0.43
(1,101)	1:4:A:GLY:HA3	1:5:A:HIS:HD2	9	0.43
(1,68)	1:6:A:VAL:HG11	1:7:A:PRO:HG2	3	0.43
(1,68)	1:6:A:VAL:HG11	1:7:A:PRO:HG3	3	0.43
(1,68)	1:6:A:VAL:HG12	1:7:A:PRO:HG2	3	0.43
(1,68)	1:6:A:VAL:HG12	1:7:A:PRO:HG3	3	0.43
(1,68)	1:6:A:VAL:HG13	1:7:A:PRO:HG2	3	0.43
(1,68)	1:6:A:VAL:HG13	1:7:A:PRO:HG3	3	0.43
(1,68)	1:6:A:VAL:HG11	1:7:A:PRO:HG2	12	0.43
(1,68)	1:6:A:VAL:HG11	1:7:A:PRO:HG3	12	0.43
(1,68)	1:6:A:VAL:HG12	1:7:A:PRO:HG2	12	0.43
(1,68)	1:6:A:VAL:HG12	1:7:A:PRO:HG3	12	0.43
(1,68)	1:6:A:VAL:HG13	1:7:A:PRO:HG2	12	0.43
(1,68)	1:6:A:VAL:HG13	1:7:A:PRO:HG3	12	0.43
(1,27)	1:9:A:TYR:HE1	2:18:B:SER:HB2	19	0.43
(1,27)	1:9:A:TYR:HE2	2:18:B:SER:HB2	19	0.43

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,1139)	1:22:C:GLY:H	2:41:D:TYR:HD1	10	0.42
(1,1139)	1:22:C:GLY:H	2:41:D:TYR:HD2	10	0.42
(1,1134)	2:39:D:SER:HB3	2:40:D:PHE:H	14	0.42
(1,1132)	2:39:D:SER:HB3	2:40:D:PHE:H	14	0.42
(1,1125)	1:31:C:PHE:HD1	1:32:C:VAL:HG11	12	0.42
(1,1125)	1:31:C:PHE:HD1	1:32:C:VAL:HG12	12	0.42
(1,1125)	1:31:C:PHE:HD1	1:32:C:VAL:HG13	12	0.42
(1,1125)	1:31:C:PHE:HD1	1:32:C:VAL:HG21	12	0.42
(1,1125)	1:31:C:PHE:HD1	1:32:C:VAL:HG22	12	0.42
(1,1125)	1:31:C:PHE:HD1	1:32:C:VAL:HG23	12	0.42
(1,1125)	1:31:C:PHE:HD2	1:32:C:VAL:HG11	12	0.42
(1,1125)	1:31:C:PHE:HD2	1:32:C:VAL:HG12	12	0.42
(1,1125)	1:31:C:PHE:HD2	1:32:C:VAL:HG13	12	0.42
(1,1125)	1:31:C:PHE:HD2	1:32:C:VAL:HG21	12	0.42
(1,1125)	1:31:C:PHE:HD2	1:32:C:VAL:HG22	12	0.42
(1,1125)	1:31:C:PHE:HD2	1:32:C:VAL:HG23	12	0.42
(1,1092)	1:29:C:GLU:HG2	2:41:D:TYR:HD1	4	0.42
(1,1092)	1:29:C:GLU:HG2	2:41:D:TYR:HD2	4	0.42
(1,1063)	1:22:C:GLY:H	2:39:D:SER:HA	6	0.42
(1,1053)	2:39:D:SER:HB2	2:40:D:PHE:H	15	0.42
(1,1009)	1:27:C:VAL:HB	2:41:D:TYR:HB2	15	0.42
(1,1009)	1:27:C:VAL:HB	2:41:D:TYR:HB3	15	0.42
(1,820)	1:22:C:GLY:H	2:41:D:TYR:HD1	10	0.42
(1,820)	1:22:C:GLY:H	2:41:D:TYR:HD2	10	0.42
(1,815)	2:39:D:SER:HB3	2:40:D:PHE:H	14	0.42
(1,813)	2:39:D:SER:HB3	2:40:D:PHE:H	14	0.42
(1,806)	1:31:C:PHE:HD1	1:32:C:VAL:HG11	12	0.42
(1,806)	1:31:C:PHE:HD1	1:32:C:VAL:HG12	12	0.42
(1,806)	1:31:C:PHE:HD1	1:32:C:VAL:HG13	12	0.42
(1,806)	1:31:C:PHE:HD1	1:32:C:VAL:HG21	12	0.42
(1,806)	1:31:C:PHE:HD1	1:32:C:VAL:HG22	12	0.42
(1,806)	1:31:C:PHE:HD1	1:32:C:VAL:HG23	12	0.42
(1,806)	1:31:C:PHE:HD2	1:32:C:VAL:HG11	12	0.42
(1,806)	1:31:C:PHE:HD2	1:32:C:VAL:HG12	12	0.42
(1,806)	1:31:C:PHE:HD2	1:32:C:VAL:HG13	12	0.42
(1,806)	1:31:C:PHE:HD2	1:32:C:VAL:HG21	12	0.42
(1,806)	1:31:C:PHE:HD2	1:32:C:VAL:HG22	12	0.42
(1,806)	1:31:C:PHE:HD2	1:32:C:VAL:HG23	12	0.42
(1,773)	1:29:C:GLU:HG2	2:41:D:TYR:HD1	4	0.42
(1,773)	1:29:C:GLU:HG2	2:41:D:TYR:HD2	4	0.42
(1,744)	1:22:C:GLY:H	2:39:D:SER:HA	6	0.42
(1,734)	2:39:D:SER:HB2	2:40:D:PHE:H	15	0.42

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,690)	1:27:C:VAL:HB	2:41:D:TYR:HB2	15	0.42
(1,690)	1:27:C:VAL:HB	2:41:D:TYR:HB3	15	0.42
(1,662)	2:19:B:PHE:HE1	2:38:D:ILE:HG13	10	0.42
(1,662)	2:19:B:PHE:HE2	2:38:D:ILE:HG13	10	0.42
(1,631)	2:17:B:ILE:HA	2:17:B:ILE:HD11	2	0.42
(1,631)	2:17:B:ILE:HA	2:17:B:ILE:HD12	2	0.42
(1,631)	2:17:B:ILE:HA	2:17:B:ILE:HD13	2	0.42
(1,499)	1:6:A:VAL:HG11	1:7:A:PRO:HD3	1	0.42
(1,499)	1:6:A:VAL:HG12	1:7:A:PRO:HD3	1	0.42
(1,499)	1:6:A:VAL:HG13	1:7:A:PRO:HD3	1	0.42
(1,481)	1:1:A:GLY:H	2:19:B:PHE:H	14	0.42
(1,474)	1:6:A:VAL:H	2:19:B:PHE:HB2	12	0.42
(1,433)	1:8:A:GLU:HB2	2:20:B:TYR:HB2	20	0.42
(1,433)	1:8:A:GLU:HB2	2:20:B:TYR:HB3	20	0.42
(1,428)	1:4:A:GLY:H	2:19:B:PHE:HB3	2	0.42
(1,397)	2:18:B:SER:HB2	2:19:B:PHE:H	3	0.42
(1,387)	1:6:A:VAL:HG11	1:7:A:PRO:HG2	16	0.42
(1,387)	1:6:A:VAL:HG11	1:7:A:PRO:HG3	16	0.42
(1,387)	1:6:A:VAL:HG12	1:7:A:PRO:HG2	16	0.42
(1,387)	1:6:A:VAL:HG12	1:7:A:PRO:HG3	16	0.42
(1,387)	1:6:A:VAL:HG13	1:7:A:PRO:HG2	16	0.42
(1,387)	1:6:A:VAL:HG13	1:7:A:PRO:HG3	16	0.42
(1,379)	1:1:A:GLY:H	2:20:B:TYR:HE1	10	0.42
(1,379)	1:1:A:GLY:H	2:20:B:TYR:HE2	10	0.42
(1,312)	2:17:B:ILE:HA	2:17:B:ILE:HD11	2	0.42
(1,312)	2:17:B:ILE:HA	2:17:B:ILE:HD12	2	0.42
(1,312)	2:17:B:ILE:HA	2:17:B:ILE:HD13	2	0.42
(1,180)	1:6:A:VAL:HG11	1:7:A:PRO:HD3	1	0.42
(1,180)	1:6:A:VAL:HG12	1:7:A:PRO:HD3	1	0.42
(1,180)	1:6:A:VAL:HG13	1:7:A:PRO:HD3	1	0.42
(1,162)	1:1:A:GLY:H	2:19:B:PHE:H	14	0.42
(1,155)	1:6:A:VAL:H	2:19:B:PHE:HB2	12	0.42
(1,114)	1:8:A:GLU:HB2	2:20:B:TYR:HB2	20	0.42
(1,114)	1:8:A:GLU:HB2	2:20:B:TYR:HB3	20	0.42
(1,109)	1:4:A:GLY:H	2:19:B:PHE:HB3	2	0.42
(1,78)	2:18:B:SER:HB2	2:19:B:PHE:H	3	0.42
(1,68)	1:6:A:VAL:HG11	1:7:A:PRO:HG2	16	0.42
(1,68)	1:6:A:VAL:HG11	1:7:A:PRO:HG3	16	0.42
(1,68)	1:6:A:VAL:HG12	1:7:A:PRO:HG2	16	0.42
(1,68)	1:6:A:VAL:HG12	1:7:A:PRO:HG3	16	0.42
(1,68)	1:6:A:VAL:HG13	1:7:A:PRO:HG2	16	0.42
(1,68)	1:6:A:VAL:HG13	1:7:A:PRO:HG3	16	0.42

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,60)	1:1:A:GLY:H	2:20:B:TYR:HE1	10	0.42
(1,60)	1:1:A:GLY:H	2:20:B:TYR:HE2	10	0.42
(1,15)	2:19:B:PHE:HE1	2:38:D:ILE:HG13	10	0.42
(1,15)	2:19:B:PHE:HE2	2:38:D:ILE:HG13	10	0.42
(1,1117)	1:27:C:VAL:HG11	1:28:C:PRO:HB3	3	0.41
(1,1117)	1:27:C:VAL:HG12	1:28:C:PRO:HB3	3	0.41
(1,1117)	1:27:C:VAL:HG13	1:28:C:PRO:HB3	3	0.41
(1,1117)	1:27:C:VAL:HG11	1:28:C:PRO:HB3	18	0.41
(1,1117)	1:27:C:VAL:HG12	1:28:C:PRO:HB3	18	0.41
(1,1117)	1:27:C:VAL:HG13	1:28:C:PRO:HB3	18	0.41
(1,1104)	1:25:C:GLY:H	2:40:D:PHE:H	4	0.41
(1,1089)	1:29:C:GLU:HB2	2:41:D:TYR:HB2	9	0.41
(1,1089)	1:29:C:GLU:HB2	2:41:D:TYR:HB3	9	0.41
(1,1043)	1:27:C:VAL:HG11	1:28:C:PRO:HG2	19	0.41
(1,1043)	1:27:C:VAL:HG11	1:28:C:PRO:HG3	19	0.41
(1,1043)	1:27:C:VAL:HG12	1:28:C:PRO:HG2	19	0.41
(1,1043)	1:27:C:VAL:HG12	1:28:C:PRO:HG3	19	0.41
(1,1043)	1:27:C:VAL:HG13	1:28:C:PRO:HG2	19	0.41
(1,1043)	1:27:C:VAL:HG13	1:28:C:PRO:HG3	19	0.41
(1,1042)	1:25:C:GLY:H	2:40:D:PHE:HD1	5	0.41
(1,1042)	1:25:C:GLY:H	2:40:D:PHE:HD2	5	0.41
(1,1003)	1:28:C:PRO:HD2	2:41:D:TYR:HD1	20	0.41
(1,1003)	1:28:C:PRO:HD2	2:41:D:TYR:HD2	20	0.41
(1,798)	1:27:C:VAL:HG11	1:28:C:PRO:HB3	3	0.41
(1,798)	1:27:C:VAL:HG12	1:28:C:PRO:HB3	3	0.41
(1,798)	1:27:C:VAL:HG13	1:28:C:PRO:HB3	3	0.41
(1,798)	1:27:C:VAL:HG11	1:28:C:PRO:HB3	18	0.41
(1,798)	1:27:C:VAL:HG12	1:28:C:PRO:HB3	18	0.41
(1,798)	1:27:C:VAL:HG13	1:28:C:PRO:HB3	18	0.41
(1,785)	1:25:C:GLY:H	2:40:D:PHE:H	4	0.41
(1,770)	1:29:C:GLU:HB2	2:41:D:TYR:HB2	9	0.41
(1,770)	1:29:C:GLU:HB2	2:41:D:TYR:HB3	9	0.41
(1,724)	1:27:C:VAL:HG11	1:28:C:PRO:HG2	19	0.41
(1,724)	1:27:C:VAL:HG11	1:28:C:PRO:HG3	19	0.41
(1,724)	1:27:C:VAL:HG12	1:28:C:PRO:HG2	19	0.41
(1,724)	1:27:C:VAL:HG12	1:28:C:PRO:HG3	19	0.41
(1,724)	1:27:C:VAL:HG13	1:28:C:PRO:HG2	19	0.41
(1,724)	1:27:C:VAL:HG13	1:28:C:PRO:HG3	19	0.41
(1,723)	1:25:C:GLY:H	2:40:D:PHE:HD1	5	0.41
(1,723)	1:25:C:GLY:H	2:40:D:PHE:HD2	5	0.41
(1,684)	1:28:C:PRO:HD2	2:41:D:TYR:HD1	20	0.41
(1,684)	1:28:C:PRO:HD2	2:41:D:TYR:HD2	20	0.41

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,653)	1:3:A:ALA:HA	2:19:B:PHE:HD1	12	0.41
(1,653)	1:3:A:ALA:HA	2:19:B:PHE:HD2	12	0.41
(1,596)	1:7:A:PRO:HG2	1:10:A:PHE:HE1	10	0.41
(1,596)	1:7:A:PRO:HG2	1:10:A:PHE:HE2	10	0.41
(1,596)	1:7:A:PRO:HG3	1:10:A:PHE:HE1	10	0.41
(1,596)	1:7:A:PRO:HG3	1:10:A:PHE:HE2	10	0.41
(1,519)	1:3:A:ALA:HB1	2:19:B:PHE:HB3	12	0.41
(1,519)	1:3:A:ALA:HB2	2:19:B:PHE:HB3	12	0.41
(1,519)	1:3:A:ALA:HB3	2:19:B:PHE:HB3	12	0.41
(1,501)	1:4:A:GLY:H	1:6:A:VAL:H	10	0.41
(1,466)	1:4:A:GLY:HA2	1:5:A:HIS:HD2	17	0.41
(1,461)	1:6:A:VAL:HG11	1:7:A:PRO:HB3	12	0.41
(1,461)	1:6:A:VAL:HG12	1:7:A:PRO:HB3	12	0.41
(1,461)	1:6:A:VAL:HG13	1:7:A:PRO:HB3	12	0.41
(1,451)	2:15:B:THR:HG21	2:16:B:PRO:HB3	11	0.41
(1,451)	2:15:B:THR:HG22	2:16:B:PRO:HB3	11	0.41
(1,451)	2:15:B:THR:HG23	2:16:B:PRO:HB3	11	0.41
(1,442)	1:8:A:GLU:HG3	2:19:B:PHE:HA	3	0.41
(1,428)	1:4:A:GLY:H	2:19:B:PHE:HB3	1	0.41
(1,405)	1:1:A:GLY:H	2:18:B:SER:HB2	6	0.41
(1,403)	1:7:A:PRO:HG2	2:19:B:PHE:HD1	1	0.41
(1,403)	1:7:A:PRO:HG2	2:19:B:PHE:HD2	1	0.41
(1,403)	1:7:A:PRO:HG3	2:19:B:PHE:HD1	1	0.41
(1,403)	1:7:A:PRO:HG3	2:19:B:PHE:HD2	1	0.41
(1,391)	1:3:A:ALA:HA	2:19:B:PHE:HD1	12	0.41
(1,391)	1:3:A:ALA:HA	2:19:B:PHE:HD2	12	0.41
(1,349)	2:13:B:CYS:HB3	2:14:B:GLY:HA3	17	0.41
(1,334)	1:3:A:ALA:HA	2:19:B:PHE:HD1	12	0.41
(1,334)	1:3:A:ALA:HA	2:19:B:PHE:HD2	12	0.41
(1,277)	1:7:A:PRO:HG2	1:10:A:PHE:HE1	10	0.41
(1,277)	1:7:A:PRO:HG2	1:10:A:PHE:HE2	10	0.41
(1,277)	1:7:A:PRO:HG3	1:10:A:PHE:HE1	10	0.41
(1,277)	1:7:A:PRO:HG3	1:10:A:PHE:HE2	10	0.41
(1,200)	1:3:A:ALA:HB1	2:19:B:PHE:HB3	12	0.41
(1,200)	1:3:A:ALA:HB2	2:19:B:PHE:HB3	12	0.41
(1,200)	1:3:A:ALA:HB3	2:19:B:PHE:HB3	12	0.41
(1,182)	1:4:A:GLY:H	1:6:A:VAL:H	10	0.41
(1,147)	1:4:A:GLY:HA2	1:5:A:HIS:HD2	17	0.41
(1,142)	1:6:A:VAL:HG11	1:7:A:PRO:HB3	12	0.41
(1,142)	1:6:A:VAL:HG12	1:7:A:PRO:HB3	12	0.41
(1,142)	1:6:A:VAL:HG13	1:7:A:PRO:HB3	12	0.41
(1,132)	2:15:B:THR:HG21	2:16:B:PRO:HB3	11	0.41

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,132)	2:15:B:THR:HG22	2:16:B:PRO:HB3	11	0.41
(1,132)	2:15:B:THR:HG23	2:16:B:PRO:HB3	11	0.41
(1,123)	1:8:A:GLU:HG3	2:19:B:PHE:HA	3	0.41
(1,109)	1:4:A:GLY:H	2:19:B:PHE:HB3	1	0.41
(1,86)	1:1:A:GLY:H	2:18:B:SER:HB2	6	0.41
(1,84)	1:7:A:PRO:HG2	2:19:B:PHE:HD1	1	0.41
(1,84)	1:7:A:PRO:HG2	2:19:B:PHE:HD2	1	0.41
(1,84)	1:7:A:PRO:HG3	2:19:B:PHE:HD1	1	0.41
(1,84)	1:7:A:PRO:HG3	2:19:B:PHE:HD2	1	0.41
(1,72)	1:3:A:ALA:HA	2:19:B:PHE:HD1	12	0.41
(1,72)	1:3:A:ALA:HA	2:19:B:PHE:HD2	12	0.41
(1,30)	2:13:B:CYS:HB3	2:14:B:GLY:HA3	17	0.41
(1,1243)	1:33:C:ARG:H	1:33:C:ARG:HE	2	0.4
(1,1231)	1:31:C:PHE:HE1	2:36:D:THR:HA	19	0.4
(1,1231)	1:31:C:PHE:HE2	2:36:D:THR:HA	19	0.4
(1,1175)	1:24:C:ALA:HB1	2:40:D:PHE:HB3	8	0.4
(1,1175)	1:24:C:ALA:HB2	2:40:D:PHE:HB3	8	0.4
(1,1175)	1:24:C:ALA:HB3	2:40:D:PHE:HB3	8	0.4
(1,1145)	1:25:C:GLY:H	2:40:D:PHE:HE1	5	0.4
(1,1145)	1:25:C:GLY:H	2:40:D:PHE:HE2	5	0.4
(1,1099)	1:29:C:GLU:HG3	2:41:D:TYR:HD1	16	0.4
(1,1099)	1:29:C:GLU:HG3	2:41:D:TYR:HD2	16	0.4
(1,1094)	1:29:C:GLU:HG2	2:40:D:PHE:HA	5	0.4
(1,1061)	1:22:C:GLY:H	2:39:D:SER:HB2	5	0.4
(1,1053)	2:39:D:SER:HB2	2:40:D:PHE:H	5	0.4
(1,1051)	2:34:D:CYS:HB2	2:35:D:GLY:H	1	0.4
(1,1046)	1:27:C:VAL:HB	2:41:D:TYR:HE1	20	0.4
(1,1046)	1:27:C:VAL:HB	2:41:D:TYR:HE2	20	0.4
(1,1003)	1:28:C:PRO:HD2	2:41:D:TYR:HD1	3	0.4
(1,1003)	1:28:C:PRO:HD2	2:41:D:TYR:HD2	3	0.4
(1,924)	1:33:C:ARG:H	1:33:C:ARG:HE	2	0.4
(1,912)	1:31:C:PHE:HE1	2:36:D:THR:HA	19	0.4
(1,912)	1:31:C:PHE:HE2	2:36:D:THR:HA	19	0.4
(1,856)	1:24:C:ALA:HB1	2:40:D:PHE:HB3	8	0.4
(1,856)	1:24:C:ALA:HB2	2:40:D:PHE:HB3	8	0.4
(1,856)	1:24:C:ALA:HB3	2:40:D:PHE:HB3	8	0.4
(1,826)	1:25:C:GLY:H	2:40:D:PHE:HE1	5	0.4
(1,826)	1:25:C:GLY:H	2:40:D:PHE:HE2	5	0.4
(1,780)	1:29:C:GLU:HG3	2:41:D:TYR:HD1	16	0.4
(1,780)	1:29:C:GLU:HG3	2:41:D:TYR:HD2	16	0.4
(1,775)	1:29:C:GLU:HG2	2:40:D:PHE:HA	5	0.4
(1,742)	1:22:C:GLY:H	2:39:D:SER:HB2	5	0.4

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,734)	2:39:D:SER:HB2	2:40:D:PHE:H	5	0.4
(1,732)	2:34:D:CYS:HB2	2:35:D:GLY:H	1	0.4
(1,727)	1:27:C:VAL:HB	2:41:D:TYR:HE1	20	0.4
(1,727)	1:27:C:VAL:HB	2:41:D:TYR:HE2	20	0.4
(1,684)	1:28:C:PRO:HD2	2:41:D:TYR:HD1	3	0.4
(1,684)	1:28:C:PRO:HD2	2:41:D:TYR:HD2	3	0.4
(1,631)	2:17:B:ILE:HA	2:17:B:ILE:HD11	8	0.4
(1,631)	2:17:B:ILE:HA	2:17:B:ILE:HD12	8	0.4
(1,631)	2:17:B:ILE:HA	2:17:B:ILE:HD13	8	0.4
(1,501)	1:4:A:GLY:H	1:6:A:VAL:H	7	0.4
(1,463)	1:5:A:HIS:HB2	1:6:A:VAL:HG11	17	0.4
(1,463)	1:5:A:HIS:HB2	1:6:A:VAL:HG12	17	0.4
(1,463)	1:5:A:HIS:HB2	1:6:A:VAL:HG13	17	0.4
(1,452)	2:15:B:THR:HG21	2:16:B:PRO:HD3	19	0.4
(1,452)	2:15:B:THR:HG22	2:16:B:PRO:HD3	19	0.4
(1,452)	2:15:B:THR:HG23	2:16:B:PRO:HD3	19	0.4
(1,397)	2:18:B:SER:HB2	2:19:B:PHE:H	8	0.4
(1,389)	1:6:A:VAL:HB	2:20:B:TYR:HD1	19	0.4
(1,389)	1:6:A:VAL:HB	2:20:B:TYR:HD2	19	0.4
(1,312)	2:17:B:ILE:HA	2:17:B:ILE:HD11	8	0.4
(1,312)	2:17:B:ILE:HA	2:17:B:ILE:HD12	8	0.4
(1,312)	2:17:B:ILE:HA	2:17:B:ILE:HD13	8	0.4
(1,182)	1:4:A:GLY:H	1:6:A:VAL:H	7	0.4
(1,144)	1:5:A:HIS:HB2	1:6:A:VAL:HG11	17	0.4
(1,144)	1:5:A:HIS:HB2	1:6:A:VAL:HG12	17	0.4
(1,144)	1:5:A:HIS:HB2	1:6:A:VAL:HG13	17	0.4
(1,133)	2:15:B:THR:HG21	2:16:B:PRO:HD3	19	0.4
(1,133)	2:15:B:THR:HG22	2:16:B:PRO:HD3	19	0.4
(1,133)	2:15:B:THR:HG23	2:16:B:PRO:HD3	19	0.4
(1,78)	2:18:B:SER:HB2	2:19:B:PHE:H	8	0.4
(1,70)	1:6:A:VAL:HB	2:20:B:TYR:HD1	19	0.4
(1,70)	1:6:A:VAL:HB	2:20:B:TYR:HD2	19	0.4
(1,1287)	2:38:D:ILE:HA	2:38:D:ILE:HD11	19	0.39
(1,1287)	2:38:D:ILE:HA	2:38:D:ILE:HD12	19	0.39
(1,1287)	2:38:D:ILE:HA	2:38:D:ILE:HD13	19	0.39
(1,1118)	1:27:C:VAL:HG11	2:41:D:TYR:H	3	0.39
(1,1118)	1:27:C:VAL:HG12	2:41:D:TYR:H	3	0.39
(1,1118)	1:27:C:VAL:HG13	2:41:D:TYR:H	3	0.39
(1,1094)	1:29:C:GLU:HG2	2:40:D:PHE:HA	16	0.39
(1,1036)	2:34:D:CYS:HB2	2:35:D:GLY:HA3	1	0.39
(1,1004)	1:32:C:VAL:HA	1:33:C:ARG:HD2	14	0.39
(1,1004)	1:32:C:VAL:HA	1:33:C:ARG:HD3	14	0.39

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,998)	1:22:C:GLY:HA3	2:39:D:SER:HB2	5	0.39
(1,995)	1:22:C:GLY:HA3	2:39:D:SER:HB2	5	0.39
(1,968)	2:38:D:ILE:HA	2:38:D:ILE:HD11	19	0.39
(1,968)	2:38:D:ILE:HA	2:38:D:ILE:HD12	19	0.39
(1,968)	2:38:D:ILE:HA	2:38:D:ILE:HD13	19	0.39
(1,799)	1:27:C:VAL:HG11	2:41:D:TYR:H	3	0.39
(1,799)	1:27:C:VAL:HG12	2:41:D:TYR:H	3	0.39
(1,799)	1:27:C:VAL:HG13	2:41:D:TYR:H	3	0.39
(1,775)	1:29:C:GLU:HG2	2:40:D:PHE:HA	16	0.39
(1,717)	2:34:D:CYS:HB2	2:35:D:GLY:HA3	1	0.39
(1,685)	1:32:C:VAL:HA	1:33:C:ARG:HD2	14	0.39
(1,685)	1:32:C:VAL:HA	1:33:C:ARG:HD3	14	0.39
(1,679)	1:22:C:GLY:HA3	2:39:D:SER:HB2	5	0.39
(1,676)	1:22:C:GLY:HA3	2:39:D:SER:HB2	5	0.39
(1,629)	2:17:B:ILE:H	2:17:B:ILE:HD11	14	0.39
(1,629)	2:17:B:ILE:H	2:17:B:ILE:HD12	14	0.39
(1,629)	2:17:B:ILE:H	2:17:B:ILE:HD13	14	0.39
(1,499)	1:6:A:VAL:HG11	1:7:A:PRO:HD3	14	0.39
(1,499)	1:6:A:VAL:HG12	1:7:A:PRO:HD3	14	0.39
(1,499)	1:6:A:VAL:HG13	1:7:A:PRO:HD3	14	0.39
(1,451)	2:15:B:THR:HG21	2:16:B:PRO:HB3	8	0.39
(1,451)	2:15:B:THR:HG22	2:16:B:PRO:HB3	8	0.39
(1,451)	2:15:B:THR:HG23	2:16:B:PRO:HB3	8	0.39
(1,449)	1:3:A:ALA:HB1	1:4:A:GLY:H	5	0.39
(1,449)	1:3:A:ALA:HB2	1:4:A:GLY:H	5	0.39
(1,449)	1:3:A:ALA:HB3	1:4:A:GLY:H	5	0.39
(1,449)	1:3:A:ALA:HB1	1:4:A:GLY:H	18	0.39
(1,449)	1:3:A:ALA:HB2	1:4:A:GLY:H	18	0.39
(1,449)	1:3:A:ALA:HB3	1:4:A:GLY:H	18	0.39
(1,442)	1:8:A:GLU:HG3	2:19:B:PHE:HA	4	0.39
(1,397)	2:18:B:SER:HB2	2:19:B:PHE:H	14	0.39
(1,387)	1:6:A:VAL:HG11	1:7:A:PRO:HG2	17	0.39
(1,387)	1:6:A:VAL:HG11	1:7:A:PRO:HG3	17	0.39
(1,387)	1:6:A:VAL:HG12	1:7:A:PRO:HG2	17	0.39
(1,387)	1:6:A:VAL:HG12	1:7:A:PRO:HG3	17	0.39
(1,387)	1:6:A:VAL:HG13	1:7:A:PRO:HG2	17	0.39
(1,387)	1:6:A:VAL:HG13	1:7:A:PRO:HG3	17	0.39
(1,346)	1:9:A:TYR:HE1	2:18:B:SER:HB2	15	0.39
(1,346)	1:9:A:TYR:HE2	2:18:B:SER:HB2	15	0.39
(1,342)	1:1:A:GLY:HA3	2:18:B:SER:HB2	15	0.39
(1,339)	1:1:A:GLY:HA3	2:18:B:SER:HB2	15	0.39
(1,310)	2:17:B:ILE:H	2:17:B:ILE:HD11	14	0.39

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,310)	2:17:B:ILE:H	2:17:B:ILE:HD12	14	0.39
(1,310)	2:17:B:ILE:H	2:17:B:ILE:HD13	14	0.39
(1,180)	1:6:A:VAL:HG11	1:7:A:PRO:HD3	14	0.39
(1,180)	1:6:A:VAL:HG12	1:7:A:PRO:HD3	14	0.39
(1,180)	1:6:A:VAL:HG13	1:7:A:PRO:HD3	14	0.39
(1,132)	2:15:B:THR:HG21	2:16:B:PRO:HB3	8	0.39
(1,132)	2:15:B:THR:HG22	2:16:B:PRO:HB3	8	0.39
(1,132)	2:15:B:THR:HG23	2:16:B:PRO:HB3	8	0.39
(1,130)	1:3:A:ALA:HB1	1:4:A:GLY:H	5	0.39
(1,130)	1:3:A:ALA:HB2	1:4:A:GLY:H	5	0.39
(1,130)	1:3:A:ALA:HB3	1:4:A:GLY:H	5	0.39
(1,130)	1:3:A:ALA:HB1	1:4:A:GLY:H	18	0.39
(1,130)	1:3:A:ALA:HB2	1:4:A:GLY:H	18	0.39
(1,130)	1:3:A:ALA:HB3	1:4:A:GLY:H	18	0.39
(1,123)	1:8:A:GLU:HG3	2:19:B:PHE:HA	4	0.39
(1,78)	2:18:B:SER:HB2	2:19:B:PHE:H	14	0.39
(1,68)	1:6:A:VAL:HG11	1:7:A:PRO:HG2	17	0.39
(1,68)	1:6:A:VAL:HG11	1:7:A:PRO:HG3	17	0.39
(1,68)	1:6:A:VAL:HG12	1:7:A:PRO:HG2	17	0.39
(1,68)	1:6:A:VAL:HG12	1:7:A:PRO:HG3	17	0.39
(1,68)	1:6:A:VAL:HG13	1:7:A:PRO:HG2	17	0.39
(1,68)	1:6:A:VAL:HG13	1:7:A:PRO:HG3	17	0.39
(1,27)	1:9:A:TYR:HE1	2:18:B:SER:HB2	15	0.39
(1,27)	1:9:A:TYR:HE2	2:18:B:SER:HB2	15	0.39
(1,23)	1:1:A:GLY:HA3	2:18:B:SER:HB2	15	0.39
(1,20)	1:1:A:GLY:HA3	2:18:B:SER:HB2	15	0.39
(1,1309)	1:24:C:ALA:HA	2:40:D:PHE:HD1	13	0.38
(1,1309)	1:24:C:ALA:HA	2:40:D:PHE:HD2	13	0.38
(1,1155)	1:27:C:VAL:HG11	1:28:C:PRO:HD3	19	0.38
(1,1155)	1:27:C:VAL:HG12	1:28:C:PRO:HD3	19	0.38
(1,1155)	1:27:C:VAL:HG13	1:28:C:PRO:HD3	19	0.38
(1,1129)	2:40:D:PHE:HB2	2:41:D:TYR:H	14	0.38
(1,1109)	1:31:C:PHE:HD1	2:36:D:THR:HG21	5	0.38
(1,1109)	1:31:C:PHE:HD1	2:36:D:THR:HG22	5	0.38
(1,1109)	1:31:C:PHE:HD1	2:36:D:THR:HG23	5	0.38
(1,1109)	1:31:C:PHE:HD2	2:36:D:THR:HG21	5	0.38
(1,1109)	1:31:C:PHE:HD2	2:36:D:THR:HG22	5	0.38
(1,1109)	1:31:C:PHE:HD2	2:36:D:THR:HG23	5	0.38
(1,1092)	1:29:C:GLU:HG2	2:41:D:TYR:HD1	15	0.38
(1,1092)	1:29:C:GLU:HG2	2:41:D:TYR:HD2	15	0.38
(1,1062)	1:32:C:VAL:HG11	1:33:C:ARG:HA	14	0.38
(1,1062)	1:32:C:VAL:HG12	1:33:C:ARG:HA	14	0.38

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,1062)	1:32:C:VAL:HG13	1:33:C:ARG:HA	14	0.38
(1,1062)	1:32:C:VAL:HG21	1:33:C:ARG:HA	14	0.38
(1,1062)	1:32:C:VAL:HG22	1:33:C:ARG:HA	14	0.38
(1,1062)	1:32:C:VAL:HG23	1:33:C:ARG:HA	14	0.38
(1,1054)	1:28:C:PRO:HA	2:40:D:PHE:HD1	3	0.38
(1,1054)	1:28:C:PRO:HA	2:40:D:PHE:HD2	3	0.38
(1,1047)	1:24:C:ALA:HA	2:40:D:PHE:HD1	13	0.38
(1,1047)	1:24:C:ALA:HA	2:40:D:PHE:HD2	13	0.38
(1,1043)	1:27:C:VAL:HG11	1:28:C:PRO:HG2	20	0.38
(1,1043)	1:27:C:VAL:HG11	1:28:C:PRO:HG3	20	0.38
(1,1043)	1:27:C:VAL:HG12	1:28:C:PRO:HG2	20	0.38
(1,1043)	1:27:C:VAL:HG12	1:28:C:PRO:HG3	20	0.38
(1,1043)	1:27:C:VAL:HG13	1:28:C:PRO:HG2	20	0.38
(1,1043)	1:27:C:VAL:HG13	1:28:C:PRO:HG3	20	0.38
(1,1010)	1:31:C:PHE:HD1	2:37:D:PRO:HB3	13	0.38
(1,1010)	1:31:C:PHE:HD2	2:37:D:PRO:HB3	13	0.38
(1,1005)	2:34:D:CYS:HB3	2:35:D:GLY:HA3	3	0.38
(1,990)	1:24:C:ALA:HA	2:40:D:PHE:HD1	13	0.38
(1,990)	1:24:C:ALA:HA	2:40:D:PHE:HD2	13	0.38
(1,836)	1:27:C:VAL:HG11	1:28:C:PRO:HD3	19	0.38
(1,836)	1:27:C:VAL:HG12	1:28:C:PRO:HD3	19	0.38
(1,836)	1:27:C:VAL:HG13	1:28:C:PRO:HD3	19	0.38
(1,810)	2:40:D:PHE:HB2	2:41:D:TYR:H	14	0.38
(1,790)	1:31:C:PHE:HD1	2:36:D:THR:HG21	5	0.38
(1,790)	1:31:C:PHE:HD1	2:36:D:THR:HG22	5	0.38
(1,790)	1:31:C:PHE:HD1	2:36:D:THR:HG23	5	0.38
(1,790)	1:31:C:PHE:HD2	2:36:D:THR:HG21	5	0.38
(1,790)	1:31:C:PHE:HD2	2:36:D:THR:HG22	5	0.38
(1,790)	1:31:C:PHE:HD2	2:36:D:THR:HG23	5	0.38
(1,773)	1:29:C:GLU:HG2	2:41:D:TYR:HD1	15	0.38
(1,773)	1:29:C:GLU:HG2	2:41:D:TYR:HD2	15	0.38
(1,743)	1:32:C:VAL:HG11	1:33:C:ARG:HA	14	0.38
(1,743)	1:32:C:VAL:HG12	1:33:C:ARG:HA	14	0.38
(1,743)	1:32:C:VAL:HG13	1:33:C:ARG:HA	14	0.38
(1,743)	1:32:C:VAL:HG21	1:33:C:ARG:HA	14	0.38
(1,743)	1:32:C:VAL:HG22	1:33:C:ARG:HA	14	0.38
(1,743)	1:32:C:VAL:HG23	1:33:C:ARG:HA	14	0.38
(1,735)	1:28:C:PRO:HA	2:40:D:PHE:HD1	3	0.38
(1,735)	1:28:C:PRO:HA	2:40:D:PHE:HD2	3	0.38
(1,728)	1:24:C:ALA:HA	2:40:D:PHE:HD1	13	0.38
(1,728)	1:24:C:ALA:HA	2:40:D:PHE:HD2	13	0.38
(1,724)	1:27:C:VAL:HG11	1:28:C:PRO:HG2	20	0.38

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,724)	1:27:C:VAL:HG11	1:28:C:PRO:HG3	20	0.38
(1,724)	1:27:C:VAL:HG12	1:28:C:PRO:HG2	20	0.38
(1,724)	1:27:C:VAL:HG12	1:28:C:PRO:HG3	20	0.38
(1,724)	1:27:C:VAL:HG13	1:28:C:PRO:HG2	20	0.38
(1,724)	1:27:C:VAL:HG13	1:28:C:PRO:HG3	20	0.38
(1,691)	1:31:C:PHE:HD1	2:37:D:PRO:HB3	13	0.38
(1,691)	1:31:C:PHE:HD2	2:37:D:PRO:HB3	13	0.38
(1,686)	2:34:D:CYS:HB3	2:35:D:GLY:HA3	3	0.38
(1,466)	1:4:A:GLY:HA2	1:5:A:HIS:HD2	19	0.38
(1,449)	1:3:A:ALA:HB1	1:4:A:GLY:H	16	0.38
(1,449)	1:3:A:ALA:HB2	1:4:A:GLY:H	16	0.38
(1,449)	1:3:A:ALA:HB3	1:4:A:GLY:H	16	0.38
(1,417)	2:17:B:ILE:HD11	2:39:D:SER:HB3	16	0.38
(1,417)	2:17:B:ILE:HD12	2:39:D:SER:HB3	16	0.38
(1,417)	2:17:B:ILE:HD13	2:39:D:SER:HB3	16	0.38
(1,386)	1:4:A:GLY:H	2:19:B:PHE:HD1	6	0.38
(1,386)	1:4:A:GLY:H	2:19:B:PHE:HD2	6	0.38
(1,349)	2:13:B:CYS:HB3	2:14:B:GLY:HA3	1	0.38
(1,147)	1:4:A:GLY:HA2	1:5:A:HIS:HD2	19	0.38
(1,130)	1:3:A:ALA:HB1	1:4:A:GLY:H	16	0.38
(1,130)	1:3:A:ALA:HB2	1:4:A:GLY:H	16	0.38
(1,130)	1:3:A:ALA:HB3	1:4:A:GLY:H	16	0.38
(1,98)	2:17:B:ILE:HD11	2:39:D:SER:HB3	16	0.38
(1,98)	2:17:B:ILE:HD12	2:39:D:SER:HB3	16	0.38
(1,98)	2:17:B:ILE:HD13	2:39:D:SER:HB3	16	0.38
(1,67)	1:4:A:GLY:H	2:19:B:PHE:HD1	6	0.38
(1,67)	1:4:A:GLY:H	2:19:B:PHE:HD2	6	0.38
(1,30)	2:13:B:CYS:HB3	2:14:B:GLY:HA3	1	0.38
(1,1287)	2:38:D:ILE:HA	2:38:D:ILE:HD11	6	0.37
(1,1287)	2:38:D:ILE:HA	2:38:D:ILE:HD12	6	0.37
(1,1287)	2:38:D:ILE:HA	2:38:D:ILE:HD13	6	0.37
(1,1283)	2:38:D:ILE:H	2:38:D:ILE:HB	5	0.37
(1,1170)	1:31:C:PHE:HB2	1:32:C:VAL:HG11	14	0.37
(1,1170)	1:31:C:PHE:HB2	1:32:C:VAL:HG12	14	0.37
(1,1170)	1:31:C:PHE:HB2	1:32:C:VAL:HG13	14	0.37
(1,1170)	1:31:C:PHE:HB2	1:32:C:VAL:HG21	14	0.37
(1,1170)	1:31:C:PHE:HB2	1:32:C:VAL:HG22	14	0.37
(1,1170)	1:31:C:PHE:HB2	1:32:C:VAL:HG23	14	0.37
(1,1159)	1:28:C:PRO:HB3	1:31:C:PHE:HE1	12	0.37
(1,1159)	1:28:C:PRO:HB3	1:31:C:PHE:HE2	12	0.37
(1,1130)	1:27:C:VAL:H	2:40:D:PHE:HB2	13	0.37
(1,1100)	2:40:D:PHE:HA	2:41:D:TYR:HD1	19	0.37

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,1100)	2:40:D:PHE:HA	2:41:D:TYR:HD2	19	0.37
(1,1099)	1:29:C:GLU:HG3	2:41:D:TYR:HD1	13	0.37
(1,1099)	1:29:C:GLU:HG3	2:41:D:TYR:HD2	13	0.37
(1,1098)	1:29:C:GLU:HG3	2:40:D:PHE:HA	10	0.37
(1,1084)	1:25:C:GLY:H	2:40:D:PHE:HB3	3	0.37
(1,1074)	1:22:C:GLY:H	2:39:D:SER:HB3	12	0.37
(1,1074)	1:22:C:GLY:H	2:39:D:SER:HB3	18	0.37
(1,1055)	1:28:C:PRO:HG2	1:31:C:PHE:HE1	4	0.37
(1,1055)	1:28:C:PRO:HG2	1:31:C:PHE:HE2	4	0.37
(1,1055)	1:28:C:PRO:HG3	1:31:C:PHE:HE1	4	0.37
(1,1055)	1:28:C:PRO:HG3	1:31:C:PHE:HE2	4	0.37
(1,1043)	1:27:C:VAL:HG11	1:28:C:PRO:HG2	12	0.37
(1,1043)	1:27:C:VAL:HG11	1:28:C:PRO:HG3	12	0.37
(1,1043)	1:27:C:VAL:HG12	1:28:C:PRO:HG2	12	0.37
(1,1043)	1:27:C:VAL:HG12	1:28:C:PRO:HG3	12	0.37
(1,1043)	1:27:C:VAL:HG13	1:28:C:PRO:HG2	12	0.37
(1,1043)	1:27:C:VAL:HG13	1:28:C:PRO:HG3	12	0.37
(1,1040)	1:31:C:PHE:HE1	2:37:D:PRO:HD2	3	0.37
(1,1040)	1:31:C:PHE:HE2	2:37:D:PRO:HD2	3	0.37
(1,1034)	2:38:D:ILE:H	2:38:D:ILE:HG13	10	0.37
(1,1000)	1:31:C:PHE:HE1	2:37:D:PRO:HD2	4	0.37
(1,1000)	1:31:C:PHE:HE2	2:37:D:PRO:HD2	4	0.37
(1,968)	2:38:D:ILE:HA	2:38:D:ILE:HD11	6	0.37
(1,968)	2:38:D:ILE:HA	2:38:D:ILE:HD12	6	0.37
(1,968)	2:38:D:ILE:HA	2:38:D:ILE:HD13	6	0.37
(1,964)	2:38:D:ILE:H	2:38:D:ILE:HB	5	0.37
(1,851)	1:31:C:PHE:HB2	1:32:C:VAL:HG11	14	0.37
(1,851)	1:31:C:PHE:HB2	1:32:C:VAL:HG12	14	0.37
(1,851)	1:31:C:PHE:HB2	1:32:C:VAL:HG13	14	0.37
(1,851)	1:31:C:PHE:HB2	1:32:C:VAL:HG21	14	0.37
(1,851)	1:31:C:PHE:HB2	1:32:C:VAL:HG22	14	0.37
(1,851)	1:31:C:PHE:HB2	1:32:C:VAL:HG23	14	0.37
(1,840)	1:28:C:PRO:HB3	1:31:C:PHE:HE1	12	0.37
(1,840)	1:28:C:PRO:HB3	1:31:C:PHE:HE2	12	0.37
(1,811)	1:27:C:VAL:H	2:40:D:PHE:HB2	13	0.37
(1,781)	2:40:D:PHE:HA	2:41:D:TYR:HD1	19	0.37
(1,781)	2:40:D:PHE:HA	2:41:D:TYR:HD2	19	0.37
(1,780)	1:29:C:GLU:HG3	2:41:D:TYR:HD1	13	0.37
(1,780)	1:29:C:GLU:HG3	2:41:D:TYR:HD2	13	0.37
(1,779)	1:29:C:GLU:HG3	2:40:D:PHE:HA	10	0.37
(1,765)	1:25:C:GLY:H	2:40:D:PHE:HB3	3	0.37
(1,755)	1:22:C:GLY:H	2:39:D:SER:HB3	12	0.37

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,755)	1:22:C:GLY:H	2:39:D:SER:HB3	18	0.37
(1,736)	1:28:C:PRO:HG2	1:31:C:PHE:HE1	4	0.37
(1,736)	1:28:C:PRO:HG2	1:31:C:PHE:HE2	4	0.37
(1,736)	1:28:C:PRO:HG3	1:31:C:PHE:HE1	4	0.37
(1,736)	1:28:C:PRO:HG3	1:31:C:PHE:HE2	4	0.37
(1,724)	1:27:C:VAL:HG11	1:28:C:PRO:HG2	12	0.37
(1,724)	1:27:C:VAL:HG11	1:28:C:PRO:HG3	12	0.37
(1,724)	1:27:C:VAL:HG12	1:28:C:PRO:HG2	12	0.37
(1,724)	1:27:C:VAL:HG12	1:28:C:PRO:HG3	12	0.37
(1,724)	1:27:C:VAL:HG13	1:28:C:PRO:HG2	12	0.37
(1,724)	1:27:C:VAL:HG13	1:28:C:PRO:HG3	12	0.37
(1,721)	1:31:C:PHE:HE1	2:37:D:PRO:HD2	3	0.37
(1,721)	1:31:C:PHE:HE2	2:37:D:PRO:HD2	3	0.37
(1,715)	2:38:D:ILE:H	2:38:D:ILE:HG13	10	0.37
(1,681)	1:31:C:PHE:HE1	2:37:D:PRO:HD2	4	0.37
(1,681)	1:31:C:PHE:HE2	2:37:D:PRO:HD2	4	0.37
(1,631)	2:17:B:ILE:HA	2:17:B:ILE:HD11	7	0.37
(1,631)	2:17:B:ILE:HA	2:17:B:ILE:HD12	7	0.37
(1,631)	2:17:B:ILE:HA	2:17:B:ILE:HD13	7	0.37
(1,590)	1:12:A:ARG:H	1:12:A:ARG:HG2	2	0.37
(1,590)	1:12:A:ARG:H	1:12:A:ARG:HG3	2	0.37
(1,514)	1:10:A:PHE:HB2	1:11:A:VAL:HG11	10	0.37
(1,514)	1:10:A:PHE:HB2	1:11:A:VAL:HG12	10	0.37
(1,514)	1:10:A:PHE:HB2	1:11:A:VAL:HG13	10	0.37
(1,514)	1:10:A:PHE:HB2	1:11:A:VAL:HG21	10	0.37
(1,514)	1:10:A:PHE:HB2	1:11:A:VAL:HG22	10	0.37
(1,514)	1:10:A:PHE:HB2	1:11:A:VAL:HG23	10	0.37
(1,499)	1:6:A:VAL:HG11	1:7:A:PRO:HD3	4	0.37
(1,499)	1:6:A:VAL:HG12	1:7:A:PRO:HD3	4	0.37
(1,499)	1:6:A:VAL:HG13	1:7:A:PRO:HD3	4	0.37
(1,463)	1:5:A:HIS:HB2	1:6:A:VAL:HG11	1	0.37
(1,463)	1:5:A:HIS:HB2	1:6:A:VAL:HG12	1	0.37
(1,463)	1:5:A:HIS:HB2	1:6:A:VAL:HG13	1	0.37
(1,461)	1:6:A:VAL:HG11	1:7:A:PRO:HB3	5	0.37
(1,461)	1:6:A:VAL:HG12	1:7:A:PRO:HB3	5	0.37
(1,461)	1:6:A:VAL:HG13	1:7:A:PRO:HB3	5	0.37
(1,449)	1:3:A:ALA:HB1	1:4:A:GLY:H	7	0.37
(1,449)	1:3:A:ALA:HB2	1:4:A:GLY:H	7	0.37
(1,449)	1:3:A:ALA:HB3	1:4:A:GLY:H	7	0.37
(1,449)	1:3:A:ALA:HB1	1:4:A:GLY:H	10	0.37
(1,449)	1:3:A:ALA:HB2	1:4:A:GLY:H	10	0.37
(1,449)	1:3:A:ALA:HB3	1:4:A:GLY:H	10	0.37

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,442)	1:8:A:GLU:HG3	2:19:B:PHE:HA	10	0.37
(1,403)	1:7:A:PRO:HG2	2:19:B:PHE:HD1	20	0.37
(1,403)	1:7:A:PRO:HG2	2:19:B:PHE:HD2	20	0.37
(1,403)	1:7:A:PRO:HG3	2:19:B:PHE:HD1	20	0.37
(1,403)	1:7:A:PRO:HG3	2:19:B:PHE:HD2	20	0.37
(1,399)	1:7:A:PRO:HG2	1:10:A:PHE:HE1	11	0.37
(1,399)	1:7:A:PRO:HG2	1:10:A:PHE:HE2	11	0.37
(1,399)	1:7:A:PRO:HG3	1:10:A:PHE:HE1	11	0.37
(1,399)	1:7:A:PRO:HG3	1:10:A:PHE:HE2	11	0.37
(1,398)	1:7:A:PRO:HA	2:19:B:PHE:HD1	1	0.37
(1,398)	1:7:A:PRO:HA	2:19:B:PHE:HD2	1	0.37
(1,352)	2:17:B:ILE:HG13	2:39:D:SER:HA	4	0.37
(1,342)	1:1:A:GLY:HA3	2:18:B:SER:HB2	19	0.37
(1,339)	1:1:A:GLY:HA3	2:18:B:SER:HB2	19	0.37
(1,312)	2:17:B:ILE:HA	2:17:B:ILE:HD11	7	0.37
(1,312)	2:17:B:ILE:HA	2:17:B:ILE:HD12	7	0.37
(1,312)	2:17:B:ILE:HA	2:17:B:ILE:HD13	7	0.37
(1,271)	1:12:A:ARG:H	1:12:A:ARG:HG2	2	0.37
(1,271)	1:12:A:ARG:H	1:12:A:ARG:HG3	2	0.37
(1,195)	1:10:A:PHE:HB2	1:11:A:VAL:HG11	10	0.37
(1,195)	1:10:A:PHE:HB2	1:11:A:VAL:HG12	10	0.37
(1,195)	1:10:A:PHE:HB2	1:11:A:VAL:HG13	10	0.37
(1,195)	1:10:A:PHE:HB2	1:11:A:VAL:HG21	10	0.37
(1,195)	1:10:A:PHE:HB2	1:11:A:VAL:HG22	10	0.37
(1,195)	1:10:A:PHE:HB2	1:11:A:VAL:HG23	10	0.37
(1,180)	1:6:A:VAL:HG11	1:7:A:PRO:HD3	4	0.37
(1,180)	1:6:A:VAL:HG12	1:7:A:PRO:HD3	4	0.37
(1,180)	1:6:A:VAL:HG13	1:7:A:PRO:HD3	4	0.37
(1,144)	1:5:A:HIS:HB2	1:6:A:VAL:HG11	1	0.37
(1,144)	1:5:A:HIS:HB2	1:6:A:VAL:HG12	1	0.37
(1,144)	1:5:A:HIS:HB2	1:6:A:VAL:HG13	1	0.37
(1,142)	1:6:A:VAL:HG11	1:7:A:PRO:HB3	5	0.37
(1,142)	1:6:A:VAL:HG12	1:7:A:PRO:HB3	5	0.37
(1,142)	1:6:A:VAL:HG13	1:7:A:PRO:HB3	5	0.37
(1,130)	1:3:A:ALA:HB1	1:4:A:GLY:H	7	0.37
(1,130)	1:3:A:ALA:HB2	1:4:A:GLY:H	7	0.37
(1,130)	1:3:A:ALA:HB3	1:4:A:GLY:H	7	0.37
(1,130)	1:3:A:ALA:HB1	1:4:A:GLY:H	10	0.37
(1,130)	1:3:A:ALA:HB2	1:4:A:GLY:H	10	0.37
(1,130)	1:3:A:ALA:HB3	1:4:A:GLY:H	10	0.37
(1,123)	1:8:A:GLU:HG3	2:19:B:PHE:HA	10	0.37
(1,84)	1:7:A:PRO:HG2	2:19:B:PHE:HD1	20	0.37

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,84)	1:7:A:PRO:HG2	2:19:B:PHE:HD2	20	0.37
(1,84)	1:7:A:PRO:HG3	2:19:B:PHE:HD1	20	0.37
(1,84)	1:7:A:PRO:HG3	2:19:B:PHE:HD2	20	0.37
(1,80)	1:7:A:PRO:HG2	1:10:A:PHE:HE1	11	0.37
(1,80)	1:7:A:PRO:HG2	1:10:A:PHE:HE2	11	0.37
(1,80)	1:7:A:PRO:HG3	1:10:A:PHE:HE1	11	0.37
(1,80)	1:7:A:PRO:HG3	1:10:A:PHE:HE2	11	0.37
(1,79)	1:7:A:PRO:HA	2:19:B:PHE:HD1	1	0.37
(1,79)	1:7:A:PRO:HA	2:19:B:PHE:HD2	1	0.37
(1,33)	2:17:B:ILE:HG13	2:39:D:SER:HA	4	0.37
(1,23)	1:1:A:GLY:HA3	2:18:B:SER:HB2	19	0.37
(1,20)	1:1:A:GLY:HA3	2:18:B:SER:HB2	19	0.37
(1,1186)	1:26:C:HIS:HA	1:26:C:HIS:HD2	10	0.36
(1,1102)	1:29:C:GLU:HB3	2:41:D:TYR:HD1	12	0.36
(1,1102)	1:29:C:GLU:HB3	2:41:D:TYR:HD2	12	0.36
(1,1101)	1:25:C:GLY:HA2	1:27:C:VAL:H	3	0.36
(1,1051)	2:34:D:CYS:HB2	2:35:D:GLY:H	20	0.36
(1,1025)	1:27:C:VAL:HG21	2:41:D:TYR:HE1	19	0.36
(1,1025)	1:27:C:VAL:HG21	2:41:D:TYR:HE2	19	0.36
(1,1025)	1:27:C:VAL:HG22	2:41:D:TYR:HE1	19	0.36
(1,1025)	1:27:C:VAL:HG22	2:41:D:TYR:HE2	19	0.36
(1,1025)	1:27:C:VAL:HG23	2:41:D:TYR:HE1	19	0.36
(1,1025)	1:27:C:VAL:HG23	2:41:D:TYR:HE2	19	0.36
(1,1017)	1:31:C:PHE:HE1	2:35:D:GLY:HA3	15	0.36
(1,1017)	1:31:C:PHE:HE2	2:35:D:GLY:HA3	15	0.36
(1,1007)	2:37:D:PRO:HA	2:38:D:ILE:HD11	4	0.36
(1,1007)	2:37:D:PRO:HA	2:38:D:ILE:HD12	4	0.36
(1,1007)	2:37:D:PRO:HA	2:38:D:ILE:HD13	4	0.36
(1,1005)	2:34:D:CYS:HB3	2:35:D:GLY:HA3	14	0.36
(1,1005)	2:34:D:CYS:HB3	2:35:D:GLY:HA3	15	0.36
(1,1000)	1:31:C:PHE:HE1	2:37:D:PRO:HD2	18	0.36
(1,1000)	1:31:C:PHE:HE2	2:37:D:PRO:HD2	18	0.36
(1,867)	1:26:C:HIS:HA	1:26:C:HIS:HD2	10	0.36
(1,783)	1:29:C:GLU:HB3	2:41:D:TYR:HD1	12	0.36
(1,783)	1:29:C:GLU:HB3	2:41:D:TYR:HD2	12	0.36
(1,782)	1:25:C:GLY:HA2	1:27:C:VAL:H	3	0.36
(1,732)	2:34:D:CYS:HB2	2:35:D:GLY:H	20	0.36
(1,706)	1:27:C:VAL:HG21	2:41:D:TYR:HE1	19	0.36
(1,706)	1:27:C:VAL:HG21	2:41:D:TYR:HE2	19	0.36
(1,706)	1:27:C:VAL:HG22	2:41:D:TYR:HE1	19	0.36
(1,706)	1:27:C:VAL:HG22	2:41:D:TYR:HE2	19	0.36
(1,706)	1:27:C:VAL:HG23	2:41:D:TYR:HE1	19	0.36

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,706)	1:27:C:VAL:HG23	2:41:D:TYR:HE2	19	0.36
(1,698)	1:31:C:PHE:HE1	2:35:D:GLY:HA3	15	0.36
(1,698)	1:31:C:PHE:HE2	2:35:D:GLY:HA3	15	0.36
(1,688)	2:37:D:PRO:HA	2:38:D:ILE:HD11	4	0.36
(1,688)	2:37:D:PRO:HA	2:38:D:ILE:HD12	4	0.36
(1,688)	2:37:D:PRO:HA	2:38:D:ILE:HD13	4	0.36
(1,686)	2:34:D:CYS:HB3	2:35:D:GLY:HA3	14	0.36
(1,686)	2:34:D:CYS:HB3	2:35:D:GLY:HA3	15	0.36
(1,681)	1:31:C:PHE:HE1	2:37:D:PRO:HD2	18	0.36
(1,681)	1:31:C:PHE:HE2	2:37:D:PRO:HD2	18	0.36
(1,631)	2:17:B:ILE:HA	2:17:B:ILE:HD11	12	0.36
(1,631)	2:17:B:ILE:HA	2:17:B:ILE:HD12	12	0.36
(1,631)	2:17:B:ILE:HA	2:17:B:ILE:HD13	12	0.36
(1,593)	1:12:A:ARG:HB2	1:12:A:ARG:HE	12	0.36
(1,520)	2:17:B:ILE:HG13	2:40:D:PHE:HB3	1	0.36
(1,516)	2:15:B:THR:HB	2:16:B:PRO:HD2	4	0.36
(1,501)	1:4:A:GLY:H	1:6:A:VAL:H	5	0.36
(1,493)	1:11:A:VAL:HB	1:12:A:ARG:H	9	0.36
(1,492)	1:7:A:PRO:HB2	1:10:A:PHE:HD1	13	0.36
(1,492)	1:7:A:PRO:HB2	1:10:A:PHE:HD2	13	0.36
(1,474)	1:6:A:VAL:H	2:19:B:PHE:HB2	7	0.36
(1,463)	1:5:A:HIS:HB2	1:6:A:VAL:HG11	19	0.36
(1,463)	1:5:A:HIS:HB2	1:6:A:VAL:HG12	19	0.36
(1,463)	1:5:A:HIS:HB2	1:6:A:VAL:HG13	19	0.36
(1,446)	1:8:A:GLU:HB3	2:20:B:TYR:HD1	1	0.36
(1,446)	1:8:A:GLU:HB3	2:20:B:TYR:HD2	1	0.36
(1,406)	1:11:A:VAL:HG11	1:12:A:ARG:HA	4	0.36
(1,406)	1:11:A:VAL:HG12	1:12:A:ARG:HA	4	0.36
(1,406)	1:11:A:VAL:HG13	1:12:A:ARG:HA	4	0.36
(1,406)	1:11:A:VAL:HG21	1:12:A:ARG:HA	4	0.36
(1,406)	1:11:A:VAL:HG22	1:12:A:ARG:HA	4	0.36
(1,406)	1:11:A:VAL:HG23	1:12:A:ARG:HA	4	0.36
(1,397)	2:18:B:SER:HB2	2:19:B:PHE:H	11	0.36
(1,384)	1:10:A:PHE:HE1	2:16:B:PRO:HD2	13	0.36
(1,384)	1:10:A:PHE:HE2	2:16:B:PRO:HD2	13	0.36
(1,378)	2:17:B:ILE:H	2:17:B:ILE:HG13	20	0.36
(1,369)	1:6:A:VAL:HG21	2:20:B:TYR:HE1	19	0.36
(1,369)	1:6:A:VAL:HG21	2:20:B:TYR:HE2	19	0.36
(1,369)	1:6:A:VAL:HG22	2:20:B:TYR:HE1	19	0.36
(1,369)	1:6:A:VAL:HG22	2:20:B:TYR:HE2	19	0.36
(1,369)	1:6:A:VAL:HG23	2:20:B:TYR:HE1	19	0.36
(1,369)	1:6:A:VAL:HG23	2:20:B:TYR:HE2	19	0.36

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,312)	2:17:B:ILE:HA	2:17:B:ILE:HD11	12	0.36
(1,312)	2:17:B:ILE:HA	2:17:B:ILE:HD12	12	0.36
(1,312)	2:17:B:ILE:HA	2:17:B:ILE:HD13	12	0.36
(1,274)	1:12:A:ARG:HB2	1:12:A:ARG:HE	12	0.36
(1,201)	2:17:B:ILE:HG13	2:40:D:PHE:HB3	1	0.36
(1,197)	2:15:B:THR:HB	2:16:B:PRO:HD2	4	0.36
(1,182)	1:4:A:GLY:H	1:6:A:VAL:H	5	0.36
(1,174)	1:11:A:VAL:HB	1:12:A:ARG:H	9	0.36
(1,173)	1:7:A:PRO:HB2	1:10:A:PHE:HD1	13	0.36
(1,173)	1:7:A:PRO:HB2	1:10:A:PHE:HD2	13	0.36
(1,155)	1:6:A:VAL:H	2:19:B:PHE:HB2	7	0.36
(1,144)	1:5:A:HIS:HB2	1:6:A:VAL:HG11	19	0.36
(1,144)	1:5:A:HIS:HB2	1:6:A:VAL:HG12	19	0.36
(1,144)	1:5:A:HIS:HB2	1:6:A:VAL:HG13	19	0.36
(1,127)	1:8:A:GLU:HB3	2:20:B:TYR:HD1	1	0.36
(1,127)	1:8:A:GLU:HB3	2:20:B:TYR:HD2	1	0.36
(1,87)	1:11:A:VAL:HG11	1:12:A:ARG:HA	4	0.36
(1,87)	1:11:A:VAL:HG12	1:12:A:ARG:HA	4	0.36
(1,87)	1:11:A:VAL:HG13	1:12:A:ARG:HA	4	0.36
(1,87)	1:11:A:VAL:HG21	1:12:A:ARG:HA	4	0.36
(1,87)	1:11:A:VAL:HG22	1:12:A:ARG:HA	4	0.36
(1,87)	1:11:A:VAL:HG23	1:12:A:ARG:HA	4	0.36
(1,78)	2:18:B:SER:HB2	2:19:B:PHE:H	11	0.36
(1,65)	1:10:A:PHE:HE1	2:16:B:PRO:HD2	13	0.36
(1,65)	1:10:A:PHE:HE2	2:16:B:PRO:HD2	13	0.36
(1,59)	2:17:B:ILE:H	2:17:B:ILE:HG13	20	0.36
(1,50)	1:6:A:VAL:HG21	2:20:B:TYR:HE1	19	0.36
(1,50)	1:6:A:VAL:HG21	2:20:B:TYR:HE2	19	0.36
(1,50)	1:6:A:VAL:HG22	2:20:B:TYR:HE1	19	0.36
(1,50)	1:6:A:VAL:HG22	2:20:B:TYR:HE2	19	0.36
(1,50)	1:6:A:VAL:HG23	2:20:B:TYR:HE1	19	0.36
(1,50)	1:6:A:VAL:HG23	2:20:B:TYR:HE2	19	0.36
(1,1243)	1:33:C:ARG:H	1:33:C:ARG:HE	10	0.35
(1,1175)	1:24:C:ALA:HB1	2:40:D:PHE:HB3	7	0.35
(1,1175)	1:24:C:ALA:HB2	2:40:D:PHE:HB3	7	0.35
(1,1175)	1:24:C:ALA:HB3	2:40:D:PHE:HB3	7	0.35
(1,1170)	1:31:C:PHE:HB2	1:32:C:VAL:HG11	13	0.35
(1,1170)	1:31:C:PHE:HB2	1:32:C:VAL:HG12	13	0.35
(1,1170)	1:31:C:PHE:HB2	1:32:C:VAL:HG13	13	0.35
(1,1170)	1:31:C:PHE:HB2	1:32:C:VAL:HG21	13	0.35
(1,1170)	1:31:C:PHE:HB2	1:32:C:VAL:HG22	13	0.35
(1,1170)	1:31:C:PHE:HB2	1:32:C:VAL:HG23	13	0.35

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,1155)	1:27:C:VAL:HG11	1:28:C:PRO:HD3	9	0.35
(1,1155)	1:27:C:VAL:HG12	1:28:C:PRO:HD3	9	0.35
(1,1155)	1:27:C:VAL:HG13	1:28:C:PRO:HD3	9	0.35
(1,1098)	1:29:C:GLU:HG3	2:40:D:PHE:HA	20	0.35
(1,1090)	1:24:C:ALA:HB1	1:25:C:GLY:HA3	15	0.35
(1,1090)	1:24:C:ALA:HB2	1:25:C:GLY:HA3	15	0.35
(1,1090)	1:24:C:ALA:HB3	1:25:C:GLY:HA3	15	0.35
(1,1078)	1:25:C:GLY:H	2:41:D:TYR:H	9	0.35
(1,1030)	1:24:C:ALA:HB1	1:25:C:GLY:HA3	15	0.35
(1,1030)	1:24:C:ALA:HB2	1:25:C:GLY:HA3	15	0.35
(1,1030)	1:24:C:ALA:HB3	1:25:C:GLY:HA3	15	0.35
(1,1002)	1:30:C:TYR:HE1	2:39:D:SER:HB2	10	0.35
(1,1002)	1:30:C:TYR:HE2	2:39:D:SER:HB2	10	0.35
(1,924)	1:33:C:ARG:H	1:33:C:ARG:HE	10	0.35
(1,856)	1:24:C:ALA:HB1	2:40:D:PHE:HB3	7	0.35
(1,856)	1:24:C:ALA:HB2	2:40:D:PHE:HB3	7	0.35
(1,856)	1:24:C:ALA:HB3	2:40:D:PHE:HB3	7	0.35
(1,851)	1:31:C:PHE:HB2	1:32:C:VAL:HG11	13	0.35
(1,851)	1:31:C:PHE:HB2	1:32:C:VAL:HG12	13	0.35
(1,851)	1:31:C:PHE:HB2	1:32:C:VAL:HG13	13	0.35
(1,851)	1:31:C:PHE:HB2	1:32:C:VAL:HG21	13	0.35
(1,851)	1:31:C:PHE:HB2	1:32:C:VAL:HG22	13	0.35
(1,851)	1:31:C:PHE:HB2	1:32:C:VAL:HG23	13	0.35
(1,836)	1:27:C:VAL:HG11	1:28:C:PRO:HD3	9	0.35
(1,836)	1:27:C:VAL:HG12	1:28:C:PRO:HD3	9	0.35
(1,836)	1:27:C:VAL:HG13	1:28:C:PRO:HD3	9	0.35
(1,779)	1:29:C:GLU:HG3	2:40:D:PHE:HA	20	0.35
(1,771)	1:24:C:ALA:HB1	1:25:C:GLY:HA3	15	0.35
(1,771)	1:24:C:ALA:HB2	1:25:C:GLY:HA3	15	0.35
(1,771)	1:24:C:ALA:HB3	1:25:C:GLY:HA3	15	0.35
(1,759)	1:25:C:GLY:H	2:41:D:TYR:H	9	0.35
(1,711)	1:24:C:ALA:HB1	1:25:C:GLY:HA3	15	0.35
(1,711)	1:24:C:ALA:HB2	1:25:C:GLY:HA3	15	0.35
(1,711)	1:24:C:ALA:HB3	1:25:C:GLY:HA3	15	0.35
(1,683)	1:30:C:TYR:HE1	2:39:D:SER:HB2	10	0.35
(1,683)	1:30:C:TYR:HE2	2:39:D:SER:HB2	10	0.35
(1,587)	1:12:A:ARG:H	1:12:A:ARG:HE	16	0.35
(1,575)	1:10:A:PHE:HE1	2:15:B:THR:HA	8	0.35
(1,575)	1:10:A:PHE:HE2	2:15:B:THR:HA	8	0.35
(1,478)	2:18:B:SER:HB3	2:19:B:PHE:H	13	0.35
(1,476)	2:18:B:SER:HB3	2:19:B:PHE:H	13	0.35
(1,461)	1:6:A:VAL:HG11	1:7:A:PRO:HB3	15	0.35

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,461)	1:6:A:VAL:HG12	1:7:A:PRO:HB3	15	0.35
(1,461)	1:6:A:VAL:HG13	1:7:A:PRO:HB3	15	0.35
(1,443)	1:8:A:GLU:HG3	2:20:B:TYR:HD1	9	0.35
(1,443)	1:8:A:GLU:HG3	2:20:B:TYR:HD2	9	0.35
(1,442)	1:8:A:GLU:HG3	2:19:B:PHE:HA	2	0.35
(1,406)	1:11:A:VAL:HG11	1:12:A:ARG:HA	7	0.35
(1,406)	1:11:A:VAL:HG12	1:12:A:ARG:HA	7	0.35
(1,406)	1:11:A:VAL:HG13	1:12:A:ARG:HA	7	0.35
(1,406)	1:11:A:VAL:HG21	1:12:A:ARG:HA	7	0.35
(1,406)	1:11:A:VAL:HG22	1:12:A:ARG:HA	7	0.35
(1,406)	1:11:A:VAL:HG23	1:12:A:ARG:HA	7	0.35
(1,404)	1:10:A:PHE:HD1	2:16:B:PRO:HA	10	0.35
(1,404)	1:10:A:PHE:HD2	2:16:B:PRO:HA	10	0.35
(1,386)	1:4:A:GLY:H	2:19:B:PHE:HD1	1	0.35
(1,386)	1:4:A:GLY:H	2:19:B:PHE:HD2	1	0.35
(1,378)	2:17:B:ILE:H	2:17:B:ILE:HG13	1	0.35
(1,369)	1:6:A:VAL:HG21	2:20:B:TYR:HE1	6	0.35
(1,369)	1:6:A:VAL:HG21	2:20:B:TYR:HE2	6	0.35
(1,369)	1:6:A:VAL:HG22	2:20:B:TYR:HE1	6	0.35
(1,369)	1:6:A:VAL:HG22	2:20:B:TYR:HE2	6	0.35
(1,369)	1:6:A:VAL:HG23	2:20:B:TYR:HE1	6	0.35
(1,369)	1:6:A:VAL:HG23	2:20:B:TYR:HE2	6	0.35
(1,268)	1:12:A:ARG:H	1:12:A:ARG:HE	16	0.35
(1,256)	1:10:A:PHE:HE1	2:15:B:THR:HA	8	0.35
(1,256)	1:10:A:PHE:HE2	2:15:B:THR:HA	8	0.35
(1,159)	2:18:B:SER:HB3	2:19:B:PHE:H	13	0.35
(1,157)	2:18:B:SER:HB3	2:19:B:PHE:H	13	0.35
(1,142)	1:6:A:VAL:HG11	1:7:A:PRO:HB3	15	0.35
(1,142)	1:6:A:VAL:HG12	1:7:A:PRO:HB3	15	0.35
(1,142)	1:6:A:VAL:HG13	1:7:A:PRO:HB3	15	0.35
(1,124)	1:8:A:GLU:HG3	2:20:B:TYR:HD1	9	0.35
(1,124)	1:8:A:GLU:HG3	2:20:B:TYR:HD2	9	0.35
(1,123)	1:8:A:GLU:HG3	2:19:B:PHE:HA	2	0.35
(1,87)	1:11:A:VAL:HG11	1:12:A:ARG:HA	7	0.35
(1,87)	1:11:A:VAL:HG12	1:12:A:ARG:HA	7	0.35
(1,87)	1:11:A:VAL:HG13	1:12:A:ARG:HA	7	0.35
(1,87)	1:11:A:VAL:HG21	1:12:A:ARG:HA	7	0.35
(1,87)	1:11:A:VAL:HG22	1:12:A:ARG:HA	7	0.35
(1,87)	1:11:A:VAL:HG23	1:12:A:ARG:HA	7	0.35
(1,85)	1:10:A:PHE:HD1	2:16:B:PRO:HA	10	0.35
(1,85)	1:10:A:PHE:HD2	2:16:B:PRO:HA	10	0.35
(1,67)	1:4:A:GLY:H	2:19:B:PHE:HD1	1	0.35

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,67)	1:4:A:GLY:H	2:19:B:PHE:HD2	1	0.35
(1,59)	2:17:B:ILE:H	2:17:B:ILE:HG13	1	0.35
(1,50)	1:6:A:VAL:HG21	2:20:B:TYR:HE1	6	0.35
(1,50)	1:6:A:VAL:HG21	2:20:B:TYR:HE2	6	0.35
(1,50)	1:6:A:VAL:HG22	2:20:B:TYR:HE1	6	0.35
(1,50)	1:6:A:VAL:HG22	2:20:B:TYR:HE2	6	0.35
(1,50)	1:6:A:VAL:HG23	2:20:B:TYR:HE1	6	0.35
(1,50)	1:6:A:VAL:HG23	2:20:B:TYR:HE2	6	0.35
(1,1287)	2:38:D:ILE:HA	2:38:D:ILE:HD11	7	0.34
(1,1287)	2:38:D:ILE:HA	2:38:D:ILE:HD12	7	0.34
(1,1287)	2:38:D:ILE:HA	2:38:D:ILE:HD13	7	0.34
(1,1249)	1:33:C:ARG:HB2	1:33:C:ARG:HE	1	0.34
(1,1243)	1:33:C:ARG:H	1:33:C:ARG:HE	5	0.34
(1,1170)	1:31:C:PHE:HB2	1:32:C:VAL:HG11	17	0.34
(1,1170)	1:31:C:PHE:HB2	1:32:C:VAL:HG12	17	0.34
(1,1170)	1:31:C:PHE:HB2	1:32:C:VAL:HG13	17	0.34
(1,1170)	1:31:C:PHE:HB2	1:32:C:VAL:HG21	17	0.34
(1,1170)	1:31:C:PHE:HB2	1:32:C:VAL:HG22	17	0.34
(1,1170)	1:31:C:PHE:HB2	1:32:C:VAL:HG23	17	0.34
(1,1155)	1:27:C:VAL:HG11	1:28:C:PRO:HD3	8	0.34
(1,1155)	1:27:C:VAL:HG12	1:28:C:PRO:HD3	8	0.34
(1,1155)	1:27:C:VAL:HG13	1:28:C:PRO:HD3	8	0.34
(1,1145)	1:25:C:GLY:H	2:40:D:PHE:HE1	11	0.34
(1,1145)	1:25:C:GLY:H	2:40:D:PHE:HE2	11	0.34
(1,1145)	1:25:C:GLY:H	2:40:D:PHE:HE1	15	0.34
(1,1145)	1:25:C:GLY:H	2:40:D:PHE:HE2	15	0.34
(1,1089)	1:29:C:GLU:HB2	2:41:D:TYR:HB2	3	0.34
(1,1089)	1:29:C:GLU:HB2	2:41:D:TYR:HB3	3	0.34
(1,1089)	1:29:C:GLU:HB2	2:41:D:TYR:HB2	6	0.34
(1,1089)	1:29:C:GLU:HB2	2:41:D:TYR:HB3	6	0.34
(1,1062)	1:32:C:VAL:HG11	1:33:C:ARG:HA	19	0.34
(1,1062)	1:32:C:VAL:HG12	1:33:C:ARG:HA	19	0.34
(1,1062)	1:32:C:VAL:HG13	1:33:C:ARG:HA	19	0.34
(1,1062)	1:32:C:VAL:HG21	1:33:C:ARG:HA	19	0.34
(1,1062)	1:32:C:VAL:HG22	1:33:C:ARG:HA	19	0.34
(1,1062)	1:32:C:VAL:HG23	1:33:C:ARG:HA	19	0.34
(1,1050)	1:28:C:PRO:HG2	1:31:C:PHE:HB2	20	0.34
(1,1050)	1:28:C:PRO:HG3	1:31:C:PHE:HB2	20	0.34
(1,1045)	1:27:C:VAL:HB	2:41:D:TYR:HD1	18	0.34
(1,1045)	1:27:C:VAL:HB	2:41:D:TYR:HD2	18	0.34
(1,1016)	1:28:C:PRO:HG2	1:31:C:PHE:HB2	20	0.34
(1,1016)	1:28:C:PRO:HG3	1:31:C:PHE:HB2	20	0.34

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,968)	2:38:D:ILE:HA	2:38:D:ILE:HD11	7	0.34
(1,968)	2:38:D:ILE:HA	2:38:D:ILE:HD12	7	0.34
(1,968)	2:38:D:ILE:HA	2:38:D:ILE:HD13	7	0.34
(1,930)	1:33:C:ARG:HB2	1:33:C:ARG:HE	1	0.34
(1,924)	1:33:C:ARG:H	1:33:C:ARG:HE	5	0.34
(1,851)	1:31:C:PHE:HB2	1:32:C:VAL:HG11	17	0.34
(1,851)	1:31:C:PHE:HB2	1:32:C:VAL:HG12	17	0.34
(1,851)	1:31:C:PHE:HB2	1:32:C:VAL:HG13	17	0.34
(1,851)	1:31:C:PHE:HB2	1:32:C:VAL:HG21	17	0.34
(1,851)	1:31:C:PHE:HB2	1:32:C:VAL:HG22	17	0.34
(1,851)	1:31:C:PHE:HB2	1:32:C:VAL:HG23	17	0.34
(1,836)	1:27:C:VAL:HG11	1:28:C:PRO:HD3	8	0.34
(1,836)	1:27:C:VAL:HG12	1:28:C:PRO:HD3	8	0.34
(1,836)	1:27:C:VAL:HG13	1:28:C:PRO:HD3	8	0.34
(1,826)	1:25:C:GLY:H	2:40:D:PHE:HE1	11	0.34
(1,826)	1:25:C:GLY:H	2:40:D:PHE:HE2	11	0.34
(1,826)	1:25:C:GLY:H	2:40:D:PHE:HE1	15	0.34
(1,826)	1:25:C:GLY:H	2:40:D:PHE:HE2	15	0.34
(1,770)	1:29:C:GLU:HB2	2:41:D:TYR:HB2	3	0.34
(1,770)	1:29:C:GLU:HB2	2:41:D:TYR:HB3	3	0.34
(1,770)	1:29:C:GLU:HB2	2:41:D:TYR:HB2	6	0.34
(1,770)	1:29:C:GLU:HB2	2:41:D:TYR:HB3	6	0.34
(1,743)	1:32:C:VAL:HG11	1:33:C:ARG:HA	19	0.34
(1,743)	1:32:C:VAL:HG12	1:33:C:ARG:HA	19	0.34
(1,743)	1:32:C:VAL:HG13	1:33:C:ARG:HA	19	0.34
(1,743)	1:32:C:VAL:HG21	1:33:C:ARG:HA	19	0.34
(1,743)	1:32:C:VAL:HG22	1:33:C:ARG:HA	19	0.34
(1,743)	1:32:C:VAL:HG23	1:33:C:ARG:HA	19	0.34
(1,731)	1:28:C:PRO:HG2	1:31:C:PHE:HB2	20	0.34
(1,731)	1:28:C:PRO:HG3	1:31:C:PHE:HB2	20	0.34
(1,726)	1:27:C:VAL:HB	2:41:D:TYR:HD1	18	0.34
(1,726)	1:27:C:VAL:HB	2:41:D:TYR:HD2	18	0.34
(1,697)	1:28:C:PRO:HG2	1:31:C:PHE:HB2	20	0.34
(1,697)	1:28:C:PRO:HG3	1:31:C:PHE:HB2	20	0.34
(1,567)	1:10:A:PHE:HB3	1:11:A:VAL:H	9	0.34
(1,514)	1:10:A:PHE:HB2	1:11:A:VAL:HG11	11	0.34
(1,514)	1:10:A:PHE:HB2	1:11:A:VAL:HG12	11	0.34
(1,514)	1:10:A:PHE:HB2	1:11:A:VAL:HG13	11	0.34
(1,514)	1:10:A:PHE:HB2	1:11:A:VAL:HG21	11	0.34
(1,514)	1:10:A:PHE:HB2	1:11:A:VAL:HG22	11	0.34
(1,514)	1:10:A:PHE:HB2	1:11:A:VAL:HG23	11	0.34
(1,501)	1:4:A:GLY:H	1:6:A:VAL:H	19	0.34

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,445)	1:4:A:GLY:HA2	1:6:A:VAL:H	9	0.34
(1,417)	2:17:B:ILE:HD11	2:39:D:SER:HB3	11	0.34
(1,417)	2:17:B:ILE:HD12	2:39:D:SER:HB3	11	0.34
(1,417)	2:17:B:ILE:HD13	2:39:D:SER:HB3	11	0.34
(1,403)	1:7:A:PRO:HG2	2:19:B:PHE:HD1	4	0.34
(1,403)	1:7:A:PRO:HG2	2:19:B:PHE:HD2	4	0.34
(1,403)	1:7:A:PRO:HG3	2:19:B:PHE:HD1	4	0.34
(1,403)	1:7:A:PRO:HG3	2:19:B:PHE:HD2	4	0.34
(1,402)	1:10:A:PHE:HD1	2:14:B:GLY:HA3	13	0.34
(1,402)	1:10:A:PHE:HD2	2:14:B:GLY:HA3	13	0.34
(1,343)	1:7:A:PRO:HD3	2:19:B:PHE:HD1	13	0.34
(1,343)	1:7:A:PRO:HD3	2:19:B:PHE:HD2	13	0.34
(1,248)	1:10:A:PHE:HB3	1:11:A:VAL:H	9	0.34
(1,195)	1:10:A:PHE:HB2	1:11:A:VAL:HG11	11	0.34
(1,195)	1:10:A:PHE:HB2	1:11:A:VAL:HG12	11	0.34
(1,195)	1:10:A:PHE:HB2	1:11:A:VAL:HG13	11	0.34
(1,195)	1:10:A:PHE:HB2	1:11:A:VAL:HG21	11	0.34
(1,195)	1:10:A:PHE:HB2	1:11:A:VAL:HG22	11	0.34
(1,195)	1:10:A:PHE:HB2	1:11:A:VAL:HG23	11	0.34
(1,182)	1:4:A:GLY:H	1:6:A:VAL:H	19	0.34
(1,126)	1:4:A:GLY:HA2	1:6:A:VAL:H	9	0.34
(1,98)	2:17:B:ILE:HD11	2:39:D:SER:HB3	11	0.34
(1,98)	2:17:B:ILE:HD12	2:39:D:SER:HB3	11	0.34
(1,98)	2:17:B:ILE:HD13	2:39:D:SER:HB3	11	0.34
(1,84)	1:7:A:PRO:HG2	2:19:B:PHE:HD1	4	0.34
(1,84)	1:7:A:PRO:HG2	2:19:B:PHE:HD2	4	0.34
(1,84)	1:7:A:PRO:HG3	2:19:B:PHE:HD1	4	0.34
(1,84)	1:7:A:PRO:HG3	2:19:B:PHE:HD2	4	0.34
(1,83)	1:10:A:PHE:HD1	2:14:B:GLY:HA3	13	0.34
(1,83)	1:10:A:PHE:HD2	2:14:B:GLY:HA3	13	0.34
(1,24)	1:7:A:PRO:HD3	2:19:B:PHE:HD1	13	0.34
(1,24)	1:7:A:PRO:HD3	2:19:B:PHE:HD2	13	0.34
(1,1159)	1:28:C:PRO:HB3	1:31:C:PHE:HE1	17	0.33
(1,1159)	1:28:C:PRO:HB3	1:31:C:PHE:HE2	17	0.33
(1,1148)	1:28:C:PRO:HB2	1:31:C:PHE:HD1	9	0.33
(1,1148)	1:28:C:PRO:HB2	1:31:C:PHE:HD2	9	0.33
(1,1126)	1:23:C:GLY:H	2:40:D:PHE:H	18	0.33
(1,1109)	1:31:C:PHE:HD1	2:36:D:THR:HG21	8	0.33
(1,1109)	1:31:C:PHE:HD1	2:36:D:THR:HG22	8	0.33
(1,1109)	1:31:C:PHE:HD1	2:36:D:THR:HG23	8	0.33
(1,1109)	1:31:C:PHE:HD2	2:36:D:THR:HG21	8	0.33
(1,1109)	1:31:C:PHE:HD2	2:36:D:THR:HG22	8	0.33

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,1109)	1:31:C:PHE:HD2	2:36:D:THR:HG23	8	0.33
(1,1102)	1:29:C:GLU:HB3	2:41:D:TYR:HD1	9	0.33
(1,1102)	1:29:C:GLU:HB3	2:41:D:TYR:HD2	9	0.33
(1,1089)	1:29:C:GLU:HB2	2:41:D:TYR:HB2	2	0.33
(1,1089)	1:29:C:GLU:HB2	2:41:D:TYR:HB3	2	0.33
(1,1089)	1:29:C:GLU:HB2	2:41:D:TYR:HB2	20	0.33
(1,1089)	1:29:C:GLU:HB2	2:41:D:TYR:HB3	20	0.33
(1,1084)	1:25:C:GLY:H	2:40:D:PHE:HB3	11	0.33
(1,1067)	1:26:C:HIS:HB3	1:27:C:VAL:HG21	19	0.33
(1,1067)	1:26:C:HIS:HB3	1:27:C:VAL:HG22	19	0.33
(1,1067)	1:26:C:HIS:HB3	1:27:C:VAL:HG23	19	0.33
(1,1015)	1:30:C:TYR:HD1	2:39:D:SER:HA	13	0.33
(1,1015)	1:30:C:TYR:HD2	2:39:D:SER:HA	13	0.33
(1,1011)	1:22:C:GLY:H	2:41:D:TYR:HA	14	0.33
(1,1008)	2:18:B:SER:HA	2:38:D:ILE:HG13	10	0.33
(1,840)	1:28:C:PRO:HB3	1:31:C:PHE:HE1	17	0.33
(1,840)	1:28:C:PRO:HB3	1:31:C:PHE:HE2	17	0.33
(1,829)	1:28:C:PRO:HB2	1:31:C:PHE:HD1	9	0.33
(1,829)	1:28:C:PRO:HB2	1:31:C:PHE:HD2	9	0.33
(1,807)	1:23:C:GLY:H	2:40:D:PHE:H	18	0.33
(1,790)	1:31:C:PHE:HD1	2:36:D:THR:HG21	8	0.33
(1,790)	1:31:C:PHE:HD1	2:36:D:THR:HG22	8	0.33
(1,790)	1:31:C:PHE:HD1	2:36:D:THR:HG23	8	0.33
(1,790)	1:31:C:PHE:HD2	2:36:D:THR:HG21	8	0.33
(1,790)	1:31:C:PHE:HD2	2:36:D:THR:HG22	8	0.33
(1,790)	1:31:C:PHE:HD2	2:36:D:THR:HG23	8	0.33
(1,783)	1:29:C:GLU:HB3	2:41:D:TYR:HD1	9	0.33
(1,783)	1:29:C:GLU:HB3	2:41:D:TYR:HD2	9	0.33
(1,770)	1:29:C:GLU:HB2	2:41:D:TYR:HB2	2	0.33
(1,770)	1:29:C:GLU:HB2	2:41:D:TYR:HB3	2	0.33
(1,770)	1:29:C:GLU:HB2	2:41:D:TYR:HB2	20	0.33
(1,770)	1:29:C:GLU:HB2	2:41:D:TYR:HB3	20	0.33
(1,765)	1:25:C:GLY:H	2:40:D:PHE:HB3	11	0.33
(1,748)	1:26:C:HIS:HB3	1:27:C:VAL:HG21	19	0.33
(1,748)	1:26:C:HIS:HB3	1:27:C:VAL:HG22	19	0.33
(1,748)	1:26:C:HIS:HB3	1:27:C:VAL:HG23	19	0.33
(1,696)	1:30:C:TYR:HD1	2:39:D:SER:HA	13	0.33
(1,696)	1:30:C:TYR:HD2	2:39:D:SER:HA	13	0.33
(1,692)	1:22:C:GLY:H	2:41:D:TYR:HA	14	0.33
(1,689)	2:18:B:SER:HA	2:38:D:ILE:HG13	10	0.33
(1,609)	2:15:B:THR:H	2:15:B:THR:HB	3	0.33
(1,485)	1:4:A:GLY:HA3	1:6:A:VAL:H	10	0.33

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,441)	1:8:A:GLU:HG3	2:20:B:TYR:HB2	5	0.33
(1,441)	1:8:A:GLU:HG3	2:20:B:TYR:HB3	5	0.33
(1,406)	1:11:A:VAL:HG11	1:12:A:ARG:HA	2	0.33
(1,406)	1:11:A:VAL:HG12	1:12:A:ARG:HA	2	0.33
(1,406)	1:11:A:VAL:HG13	1:12:A:ARG:HA	2	0.33
(1,406)	1:11:A:VAL:HG21	1:12:A:ARG:HA	2	0.33
(1,406)	1:11:A:VAL:HG22	1:12:A:ARG:HA	2	0.33
(1,406)	1:11:A:VAL:HG23	1:12:A:ARG:HA	2	0.33
(1,389)	1:6:A:VAL:HB	2:20:B:TYR:HD1	11	0.33
(1,389)	1:6:A:VAL:HB	2:20:B:TYR:HD2	11	0.33
(1,370)	1:6:A:VAL:HG11	1:7:A:PRO:HD3	19	0.33
(1,370)	1:6:A:VAL:HG12	1:7:A:PRO:HD3	19	0.33
(1,370)	1:6:A:VAL:HG13	1:7:A:PRO:HD3	19	0.33
(1,359)	1:9:A:TYR:HD1	2:18:B:SER:HA	15	0.33
(1,359)	1:9:A:TYR:HD2	2:18:B:SER:HA	15	0.33
(1,349)	2:13:B:CYS:HB3	2:14:B:GLY:HA3	14	0.33
(1,290)	2:15:B:THR:H	2:15:B:THR:HB	3	0.33
(1,166)	1:4:A:GLY:HA3	1:6:A:VAL:H	10	0.33
(1,122)	1:8:A:GLU:HG3	2:20:B:TYR:HB2	5	0.33
(1,122)	1:8:A:GLU:HG3	2:20:B:TYR:HB3	5	0.33
(1,87)	1:11:A:VAL:HG11	1:12:A:ARG:HA	2	0.33
(1,87)	1:11:A:VAL:HG12	1:12:A:ARG:HA	2	0.33
(1,87)	1:11:A:VAL:HG13	1:12:A:ARG:HA	2	0.33
(1,87)	1:11:A:VAL:HG21	1:12:A:ARG:HA	2	0.33
(1,87)	1:11:A:VAL:HG22	1:12:A:ARG:HA	2	0.33
(1,87)	1:11:A:VAL:HG23	1:12:A:ARG:HA	2	0.33
(1,70)	1:6:A:VAL:HB	2:20:B:TYR:HD1	11	0.33
(1,70)	1:6:A:VAL:HB	2:20:B:TYR:HD2	11	0.33
(1,51)	1:6:A:VAL:HG11	1:7:A:PRO:HD3	19	0.33
(1,51)	1:6:A:VAL:HG12	1:7:A:PRO:HD3	19	0.33
(1,51)	1:6:A:VAL:HG13	1:7:A:PRO:HD3	19	0.33
(1,40)	1:9:A:TYR:HD1	2:18:B:SER:HA	15	0.33
(1,40)	1:9:A:TYR:HD2	2:18:B:SER:HA	15	0.33
(1,30)	2:13:B:CYS:HB3	2:14:B:GLY:HA3	14	0.33
(1,1252)	1:28:C:PRO:HG2	1:31:C:PHE:HE1	4	0.32
(1,1252)	1:28:C:PRO:HG2	1:31:C:PHE:HE2	4	0.32
(1,1252)	1:28:C:PRO:HG3	1:31:C:PHE:HE1	4	0.32
(1,1252)	1:28:C:PRO:HG3	1:31:C:PHE:HE2	4	0.32
(1,1130)	1:27:C:VAL:H	2:40:D:PHE:HB2	12	0.32
(1,1117)	1:27:C:VAL:HG11	1:28:C:PRO:HB3	20	0.32
(1,1117)	1:27:C:VAL:HG12	1:28:C:PRO:HB3	20	0.32
(1,1117)	1:27:C:VAL:HG13	1:28:C:PRO:HB3	20	0.32

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,1096)	1:31:C:PHE:HD1	2:37:D:PRO:HG2	6	0.32
(1,1096)	1:31:C:PHE:HD1	2:37:D:PRO:HG3	6	0.32
(1,1096)	1:31:C:PHE:HD2	2:37:D:PRO:HG2	6	0.32
(1,1096)	1:31:C:PHE:HD2	2:37:D:PRO:HG3	6	0.32
(1,1080)	1:25:C:GLY:H	2:41:D:TYR:H	11	0.32
(1,1053)	2:39:D:SER:HB2	2:40:D:PHE:H	16	0.32
(1,1046)	1:27:C:VAL:HB	2:41:D:TYR:HE1	6	0.32
(1,1046)	1:27:C:VAL:HB	2:41:D:TYR:HE2	6	0.32
(1,1003)	1:28:C:PRO:HD2	2:41:D:TYR:HD1	9	0.32
(1,1003)	1:28:C:PRO:HD2	2:41:D:TYR:HD2	9	0.32
(1,933)	1:28:C:PRO:HG2	1:31:C:PHE:HE1	4	0.32
(1,933)	1:28:C:PRO:HG2	1:31:C:PHE:HE2	4	0.32
(1,933)	1:28:C:PRO:HG3	1:31:C:PHE:HE1	4	0.32
(1,933)	1:28:C:PRO:HG3	1:31:C:PHE:HE2	4	0.32
(1,811)	1:27:C:VAL:H	2:40:D:PHE:HB2	12	0.32
(1,798)	1:27:C:VAL:HG11	1:28:C:PRO:HB3	20	0.32
(1,798)	1:27:C:VAL:HG12	1:28:C:PRO:HB3	20	0.32
(1,798)	1:27:C:VAL:HG13	1:28:C:PRO:HB3	20	0.32
(1,777)	1:31:C:PHE:HD1	2:37:D:PRO:HG2	6	0.32
(1,777)	1:31:C:PHE:HD1	2:37:D:PRO:HG3	6	0.32
(1,777)	1:31:C:PHE:HD2	2:37:D:PRO:HG2	6	0.32
(1,777)	1:31:C:PHE:HD2	2:37:D:PRO:HG3	6	0.32
(1,761)	1:25:C:GLY:H	2:41:D:TYR:H	11	0.32
(1,734)	2:39:D:SER:HB2	2:40:D:PHE:H	16	0.32
(1,727)	1:27:C:VAL:HB	2:41:D:TYR:HE1	6	0.32
(1,727)	1:27:C:VAL:HB	2:41:D:TYR:HE2	6	0.32
(1,684)	1:28:C:PRO:HD2	2:41:D:TYR:HD1	9	0.32
(1,684)	1:28:C:PRO:HD2	2:41:D:TYR:HD2	9	0.32
(1,609)	2:15:B:THR:H	2:15:B:THR:HB	17	0.32
(1,596)	1:7:A:PRO:HG2	1:10:A:PHE:HE1	11	0.32
(1,596)	1:7:A:PRO:HG2	1:10:A:PHE:HE2	11	0.32
(1,596)	1:7:A:PRO:HG3	1:10:A:PHE:HE1	11	0.32
(1,596)	1:7:A:PRO:HG3	1:10:A:PHE:HE2	11	0.32
(1,461)	1:6:A:VAL:HG11	1:7:A:PRO:HB3	3	0.32
(1,461)	1:6:A:VAL:HG12	1:7:A:PRO:HB3	3	0.32
(1,461)	1:6:A:VAL:HG13	1:7:A:PRO:HB3	3	0.32
(1,449)	1:3:A:ALA:HB1	1:4:A:GLY:H	13	0.32
(1,449)	1:3:A:ALA:HB2	1:4:A:GLY:H	13	0.32
(1,449)	1:3:A:ALA:HB3	1:4:A:GLY:H	13	0.32
(1,397)	2:18:B:SER:HB2	2:19:B:PHE:H	2	0.32
(1,389)	1:6:A:VAL:HB	2:20:B:TYR:HD1	4	0.32
(1,389)	1:6:A:VAL:HB	2:20:B:TYR:HD2	4	0.32

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,345)	1:9:A:TYR:HD1	2:18:B:SER:HB2	9	0.32
(1,345)	1:9:A:TYR:HD2	2:18:B:SER:HB2	9	0.32
(1,290)	2:15:B:THR:H	2:15:B:THR:HB	17	0.32
(1,277)	1:7:A:PRO:HG2	1:10:A:PHE:HE1	11	0.32
(1,277)	1:7:A:PRO:HG2	1:10:A:PHE:HE2	11	0.32
(1,277)	1:7:A:PRO:HG3	1:10:A:PHE:HE1	11	0.32
(1,277)	1:7:A:PRO:HG3	1:10:A:PHE:HE2	11	0.32
(1,142)	1:6:A:VAL:HG11	1:7:A:PRO:HB3	3	0.32
(1,142)	1:6:A:VAL:HG12	1:7:A:PRO:HB3	3	0.32
(1,142)	1:6:A:VAL:HG13	1:7:A:PRO:HB3	3	0.32
(1,130)	1:3:A:ALA:HB1	1:4:A:GLY:H	13	0.32
(1,130)	1:3:A:ALA:HB2	1:4:A:GLY:H	13	0.32
(1,130)	1:3:A:ALA:HB3	1:4:A:GLY:H	13	0.32
(1,78)	2:18:B:SER:HB2	2:19:B:PHE:H	2	0.32
(1,70)	1:6:A:VAL:HB	2:20:B:TYR:HD1	4	0.32
(1,70)	1:6:A:VAL:HB	2:20:B:TYR:HD2	4	0.32
(1,26)	1:9:A:TYR:HD1	2:18:B:SER:HB2	9	0.32
(1,26)	1:9:A:TYR:HD2	2:18:B:SER:HB2	9	0.32
(1,1159)	1:28:C:PRO:HB3	1:31:C:PHE:HE1	11	0.31
(1,1159)	1:28:C:PRO:HB3	1:31:C:PHE:HE2	11	0.31
(1,1155)	1:27:C:VAL:HG11	1:28:C:PRO:HD3	18	0.31
(1,1155)	1:27:C:VAL:HG12	1:28:C:PRO:HD3	18	0.31
(1,1155)	1:27:C:VAL:HG13	1:28:C:PRO:HD3	18	0.31
(1,1149)	1:32:C:VAL:HB	1:33:C:ARG:H	19	0.31
(1,1105)	1:24:C:ALA:HB1	1:25:C:GLY:H	8	0.31
(1,1105)	1:24:C:ALA:HB2	1:25:C:GLY:H	8	0.31
(1,1105)	1:24:C:ALA:HB3	1:25:C:GLY:H	8	0.31
(1,1059)	1:28:C:PRO:HG2	2:40:D:PHE:HD1	7	0.31
(1,1059)	1:28:C:PRO:HG2	2:40:D:PHE:HD2	7	0.31
(1,1059)	1:28:C:PRO:HG3	2:40:D:PHE:HD1	7	0.31
(1,1059)	1:28:C:PRO:HG3	2:40:D:PHE:HD2	7	0.31
(1,1049)	1:28:C:PRO:HG2	1:31:C:PHE:HD1	12	0.31
(1,1049)	1:28:C:PRO:HG2	1:31:C:PHE:HD2	12	0.31
(1,1049)	1:28:C:PRO:HG3	1:31:C:PHE:HD1	12	0.31
(1,1049)	1:28:C:PRO:HG3	1:31:C:PHE:HD2	12	0.31
(1,1043)	1:27:C:VAL:HG11	1:28:C:PRO:HG2	4	0.31
(1,1043)	1:27:C:VAL:HG11	1:28:C:PRO:HG3	4	0.31
(1,1043)	1:27:C:VAL:HG12	1:28:C:PRO:HG2	4	0.31
(1,1043)	1:27:C:VAL:HG12	1:28:C:PRO:HG3	4	0.31
(1,1043)	1:27:C:VAL:HG13	1:28:C:PRO:HG2	4	0.31
(1,1043)	1:27:C:VAL:HG13	1:28:C:PRO:HG3	4	0.31
(1,1034)	2:38:D:ILE:H	2:38:D:ILE:HG13	11	0.31

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,1028)	2:39:D:SER:HA	2:40:D:PHE:HD1	2	0.31
(1,1028)	2:39:D:SER:HA	2:40:D:PHE:HD2	2	0.31
(1,840)	1:28:C:PRO:HB3	1:31:C:PHE:HE1	11	0.31
(1,840)	1:28:C:PRO:HB3	1:31:C:PHE:HE2	11	0.31
(1,836)	1:27:C:VAL:HG11	1:28:C:PRO:HD3	18	0.31
(1,836)	1:27:C:VAL:HG12	1:28:C:PRO:HD3	18	0.31
(1,836)	1:27:C:VAL:HG13	1:28:C:PRO:HD3	18	0.31
(1,830)	1:32:C:VAL:HB	1:33:C:ARG:H	19	0.31
(1,786)	1:24:C:ALA:HB1	1:25:C:GLY:H	8	0.31
(1,786)	1:24:C:ALA:HB2	1:25:C:GLY:H	8	0.31
(1,786)	1:24:C:ALA:HB3	1:25:C:GLY:H	8	0.31
(1,740)	1:28:C:PRO:HG2	2:40:D:PHE:HD1	7	0.31
(1,740)	1:28:C:PRO:HG2	2:40:D:PHE:HD2	7	0.31
(1,740)	1:28:C:PRO:HG3	2:40:D:PHE:HD1	7	0.31
(1,740)	1:28:C:PRO:HG3	2:40:D:PHE:HD2	7	0.31
(1,730)	1:28:C:PRO:HG2	1:31:C:PHE:HD1	12	0.31
(1,730)	1:28:C:PRO:HG2	1:31:C:PHE:HD2	12	0.31
(1,730)	1:28:C:PRO:HG3	1:31:C:PHE:HD1	12	0.31
(1,730)	1:28:C:PRO:HG3	1:31:C:PHE:HD2	12	0.31
(1,724)	1:27:C:VAL:HG11	1:28:C:PRO:HG2	4	0.31
(1,724)	1:27:C:VAL:HG11	1:28:C:PRO:HG3	4	0.31
(1,724)	1:27:C:VAL:HG12	1:28:C:PRO:HG2	4	0.31
(1,724)	1:27:C:VAL:HG12	1:28:C:PRO:HG3	4	0.31
(1,724)	1:27:C:VAL:HG13	1:28:C:PRO:HG2	4	0.31
(1,724)	1:27:C:VAL:HG13	1:28:C:PRO:HG3	4	0.31
(1,715)	2:38:D:ILE:H	2:38:D:ILE:HG13	11	0.31
(1,709)	2:39:D:SER:HA	2:40:D:PHE:HD1	2	0.31
(1,709)	2:39:D:SER:HA	2:40:D:PHE:HD2	2	0.31
(1,668)	1:3:A:ALA:HB1	1:32:C:VAL:HA	8	0.31
(1,668)	1:3:A:ALA:HB2	1:32:C:VAL:HA	8	0.31
(1,668)	1:3:A:ALA:HB3	1:32:C:VAL:HA	8	0.31
(1,629)	2:17:B:ILE:H	2:17:B:ILE:HD11	6	0.31
(1,629)	2:17:B:ILE:H	2:17:B:ILE:HD12	6	0.31
(1,629)	2:17:B:ILE:H	2:17:B:ILE:HD13	6	0.31
(1,514)	1:10:A:PHE:HB2	1:11:A:VAL:HG11	13	0.31
(1,514)	1:10:A:PHE:HB2	1:11:A:VAL:HG12	13	0.31
(1,514)	1:10:A:PHE:HB2	1:11:A:VAL:HG13	13	0.31
(1,514)	1:10:A:PHE:HB2	1:11:A:VAL:HG21	13	0.31
(1,514)	1:10:A:PHE:HB2	1:11:A:VAL:HG22	13	0.31
(1,514)	1:10:A:PHE:HB2	1:11:A:VAL:HG23	13	0.31
(1,469)	1:10:A:PHE:HD1	1:11:A:VAL:HG11	8	0.31
(1,469)	1:10:A:PHE:HD1	1:11:A:VAL:HG12	8	0.31

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,469)	1:10:A:PHE:HD1	1:11:A:VAL:HG13	8	0.31
(1,469)	1:10:A:PHE:HD1	1:11:A:VAL:HG21	8	0.31
(1,469)	1:10:A:PHE:HD1	1:11:A:VAL:HG22	8	0.31
(1,469)	1:10:A:PHE:HD1	1:11:A:VAL:HG23	8	0.31
(1,469)	1:10:A:PHE:HD2	1:11:A:VAL:HG11	8	0.31
(1,469)	1:10:A:PHE:HD2	1:11:A:VAL:HG12	8	0.31
(1,469)	1:10:A:PHE:HD2	1:11:A:VAL:HG13	8	0.31
(1,469)	1:10:A:PHE:HD2	1:11:A:VAL:HG21	8	0.31
(1,469)	1:10:A:PHE:HD2	1:11:A:VAL:HG22	8	0.31
(1,469)	1:10:A:PHE:HD2	1:11:A:VAL:HG23	8	0.31
(1,467)	1:10:A:PHE:HB3	1:11:A:VAL:HG11	12	0.31
(1,467)	1:10:A:PHE:HB3	1:11:A:VAL:HG12	12	0.31
(1,467)	1:10:A:PHE:HB3	1:11:A:VAL:HG13	12	0.31
(1,467)	1:10:A:PHE:HB3	1:11:A:VAL:HG21	12	0.31
(1,467)	1:10:A:PHE:HB3	1:11:A:VAL:HG22	12	0.31
(1,467)	1:10:A:PHE:HB3	1:11:A:VAL:HG23	12	0.31
(1,451)	2:15:B:THR:HG21	2:16:B:PRO:HB3	9	0.31
(1,451)	2:15:B:THR:HG22	2:16:B:PRO:HB3	9	0.31
(1,451)	2:15:B:THR:HG23	2:16:B:PRO:HB3	9	0.31
(1,438)	1:8:A:GLU:HG2	2:19:B:PHE:HA	9	0.31
(1,438)	1:8:A:GLU:HG2	2:19:B:PHE:HA	16	0.31
(1,434)	1:3:A:ALA:HB1	1:4:A:GLY:HA3	4	0.31
(1,434)	1:3:A:ALA:HB2	1:4:A:GLY:HA3	4	0.31
(1,434)	1:3:A:ALA:HB3	1:4:A:GLY:HA3	4	0.31
(1,411)	1:5:A:HIS:HB3	1:6:A:VAL:HG21	14	0.31
(1,411)	1:5:A:HIS:HB3	1:6:A:VAL:HG22	14	0.31
(1,411)	1:5:A:HIS:HB3	1:6:A:VAL:HG23	14	0.31
(1,374)	1:3:A:ALA:HB1	1:4:A:GLY:HA3	4	0.31
(1,374)	1:3:A:ALA:HB2	1:4:A:GLY:HA3	4	0.31
(1,374)	1:3:A:ALA:HB3	1:4:A:GLY:HA3	4	0.31
(1,362)	1:7:A:PRO:HG2	2:19:B:PHE:HD1	12	0.31
(1,362)	1:7:A:PRO:HG2	2:19:B:PHE:HD2	12	0.31
(1,362)	1:7:A:PRO:HG3	2:19:B:PHE:HD1	12	0.31
(1,362)	1:7:A:PRO:HG3	2:19:B:PHE:HD2	12	0.31
(1,310)	2:17:B:ILE:H	2:17:B:ILE:HD11	6	0.31
(1,310)	2:17:B:ILE:H	2:17:B:ILE:HD12	6	0.31
(1,310)	2:17:B:ILE:H	2:17:B:ILE:HD13	6	0.31
(1,195)	1:10:A:PHE:HB2	1:11:A:VAL:HG11	13	0.31
(1,195)	1:10:A:PHE:HB2	1:11:A:VAL:HG12	13	0.31
(1,195)	1:10:A:PHE:HB2	1:11:A:VAL:HG13	13	0.31
(1,195)	1:10:A:PHE:HB2	1:11:A:VAL:HG21	13	0.31
(1,195)	1:10:A:PHE:HB2	1:11:A:VAL:HG22	13	0.31

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,195)	1:10:A:PHE:HB2	1:11:A:VAL:HG23	13	0.31
(1,150)	1:10:A:PHE:HD1	1:11:A:VAL:HG11	8	0.31
(1,150)	1:10:A:PHE:HD1	1:11:A:VAL:HG12	8	0.31
(1,150)	1:10:A:PHE:HD1	1:11:A:VAL:HG13	8	0.31
(1,150)	1:10:A:PHE:HD1	1:11:A:VAL:HG21	8	0.31
(1,150)	1:10:A:PHE:HD1	1:11:A:VAL:HG22	8	0.31
(1,150)	1:10:A:PHE:HD1	1:11:A:VAL:HG23	8	0.31
(1,150)	1:10:A:PHE:HD2	1:11:A:VAL:HG11	8	0.31
(1,150)	1:10:A:PHE:HD2	1:11:A:VAL:HG12	8	0.31
(1,150)	1:10:A:PHE:HD2	1:11:A:VAL:HG13	8	0.31
(1,150)	1:10:A:PHE:HD2	1:11:A:VAL:HG21	8	0.31
(1,150)	1:10:A:PHE:HD2	1:11:A:VAL:HG22	8	0.31
(1,150)	1:10:A:PHE:HD2	1:11:A:VAL:HG23	8	0.31
(1,148)	1:10:A:PHE:HB3	1:11:A:VAL:HG11	12	0.31
(1,148)	1:10:A:PHE:HB3	1:11:A:VAL:HG12	12	0.31
(1,148)	1:10:A:PHE:HB3	1:11:A:VAL:HG13	12	0.31
(1,148)	1:10:A:PHE:HB3	1:11:A:VAL:HG21	12	0.31
(1,148)	1:10:A:PHE:HB3	1:11:A:VAL:HG22	12	0.31
(1,148)	1:10:A:PHE:HB3	1:11:A:VAL:HG23	12	0.31
(1,132)	2:15:B:THR:HG21	2:16:B:PRO:HB3	9	0.31
(1,132)	2:15:B:THR:HG22	2:16:B:PRO:HB3	9	0.31
(1,132)	2:15:B:THR:HG23	2:16:B:PRO:HB3	9	0.31
(1,119)	1:8:A:GLU:HG2	2:19:B:PHE:HA	9	0.31
(1,119)	1:8:A:GLU:HG2	2:19:B:PHE:HA	16	0.31
(1,115)	1:3:A:ALA:HB1	1:4:A:GLY:HA3	4	0.31
(1,115)	1:3:A:ALA:HB2	1:4:A:GLY:HA3	4	0.31
(1,115)	1:3:A:ALA:HB3	1:4:A:GLY:HA3	4	0.31
(1,92)	1:5:A:HIS:HB3	1:6:A:VAL:HG21	14	0.31
(1,92)	1:5:A:HIS:HB3	1:6:A:VAL:HG22	14	0.31
(1,92)	1:5:A:HIS:HB3	1:6:A:VAL:HG23	14	0.31
(1,55)	1:3:A:ALA:HB1	1:4:A:GLY:HA3	4	0.31
(1,55)	1:3:A:ALA:HB2	1:4:A:GLY:HA3	4	0.31
(1,55)	1:3:A:ALA:HB3	1:4:A:GLY:HA3	4	0.31
(1,43)	1:7:A:PRO:HG2	2:19:B:PHE:HD1	12	0.31
(1,43)	1:7:A:PRO:HG2	2:19:B:PHE:HD2	12	0.31
(1,43)	1:7:A:PRO:HG3	2:19:B:PHE:HD1	12	0.31
(1,43)	1:7:A:PRO:HG3	2:19:B:PHE:HD2	12	0.31
(1,3)	1:3:A:ALA:HB1	1:32:C:VAL:HA	8	0.31
(1,3)	1:3:A:ALA:HB2	1:32:C:VAL:HA	8	0.31
(1,3)	1:3:A:ALA:HB3	1:32:C:VAL:HA	8	0.31
(1,1231)	1:31:C:PHE:HE1	2:36:D:THR:HA	18	0.3
(1,1231)	1:31:C:PHE:HE2	2:36:D:THR:HA	18	0.3

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,1170)	1:31:C:PHE:HB2	1:32:C:VAL:HG11	1	0.3
(1,1170)	1:31:C:PHE:HB2	1:32:C:VAL:HG12	1	0.3
(1,1170)	1:31:C:PHE:HB2	1:32:C:VAL:HG13	1	0.3
(1,1170)	1:31:C:PHE:HB2	1:32:C:VAL:HG21	1	0.3
(1,1170)	1:31:C:PHE:HB2	1:32:C:VAL:HG22	1	0.3
(1,1170)	1:31:C:PHE:HB2	1:32:C:VAL:HG23	1	0.3
(1,1169)	2:36:D:THR:HB	2:38:D:ILE:HD11	14	0.3
(1,1169)	2:36:D:THR:HB	2:38:D:ILE:HD12	14	0.3
(1,1169)	2:36:D:THR:HB	2:38:D:ILE:HD13	14	0.3
(1,1155)	1:27:C:VAL:HG11	1:28:C:PRO:HD3	7	0.3
(1,1155)	1:27:C:VAL:HG12	1:28:C:PRO:HD3	7	0.3
(1,1155)	1:27:C:VAL:HG13	1:28:C:PRO:HD3	7	0.3
(1,1139)	1:22:C:GLY:H	2:41:D:TYR:HD1	7	0.3
(1,1139)	1:22:C:GLY:H	2:41:D:TYR:HD2	7	0.3
(1,1130)	1:27:C:VAL:H	2:40:D:PHE:HB2	17	0.3
(1,1117)	1:27:C:VAL:HG11	1:28:C:PRO:HB3	14	0.3
(1,1117)	1:27:C:VAL:HG12	1:28:C:PRO:HB3	14	0.3
(1,1117)	1:27:C:VAL:HG13	1:28:C:PRO:HB3	14	0.3
(1,1109)	1:31:C:PHE:HD1	2:36:D:THR:HG21	19	0.3
(1,1109)	1:31:C:PHE:HD1	2:36:D:THR:HG22	19	0.3
(1,1109)	1:31:C:PHE:HD1	2:36:D:THR:HG23	19	0.3
(1,1109)	1:31:C:PHE:HD2	2:36:D:THR:HG21	19	0.3
(1,1109)	1:31:C:PHE:HD2	2:36:D:THR:HG22	19	0.3
(1,1109)	1:31:C:PHE:HD2	2:36:D:THR:HG23	19	0.3
(1,1102)	1:29:C:GLU:HB3	2:41:D:TYR:HD1	20	0.3
(1,1102)	1:29:C:GLU:HB3	2:41:D:TYR:HD2	20	0.3
(1,1098)	1:29:C:GLU:HG3	2:40:D:PHE:HA	18	0.3
(1,1094)	1:29:C:GLU:HG2	2:40:D:PHE:HA	4	0.3
(1,1094)	1:29:C:GLU:HG2	2:40:D:PHE:HA	15	0.3
(1,1078)	1:25:C:GLY:H	2:41:D:TYR:H	18	0.3
(1,1058)	1:31:C:PHE:HD1	2:35:D:GLY:HA3	4	0.3
(1,1058)	1:31:C:PHE:HD2	2:35:D:GLY:HA3	4	0.3
(1,1043)	1:27:C:VAL:HG11	1:28:C:PRO:HG2	14	0.3
(1,1043)	1:27:C:VAL:HG11	1:28:C:PRO:HG3	14	0.3
(1,1043)	1:27:C:VAL:HG12	1:28:C:PRO:HG2	14	0.3
(1,1043)	1:27:C:VAL:HG12	1:28:C:PRO:HG3	14	0.3
(1,1043)	1:27:C:VAL:HG13	1:28:C:PRO:HG2	14	0.3
(1,1043)	1:27:C:VAL:HG13	1:28:C:PRO:HG3	14	0.3
(1,1005)	2:34:D:CYS:HB3	2:35:D:GLY:HA3	7	0.3
(1,1005)	2:34:D:CYS:HB3	2:35:D:GLY:HA3	8	0.3
(1,912)	1:31:C:PHE:HE1	2:36:D:THR:HA	18	0.3
(1,912)	1:31:C:PHE:HE2	2:36:D:THR:HA	18	0.3

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,851)	1:31:C:PHE:HB2	1:32:C:VAL:HG11	1	0.3
(1,851)	1:31:C:PHE:HB2	1:32:C:VAL:HG12	1	0.3
(1,851)	1:31:C:PHE:HB2	1:32:C:VAL:HG13	1	0.3
(1,851)	1:31:C:PHE:HB2	1:32:C:VAL:HG21	1	0.3
(1,851)	1:31:C:PHE:HB2	1:32:C:VAL:HG22	1	0.3
(1,851)	1:31:C:PHE:HB2	1:32:C:VAL:HG23	1	0.3
(1,850)	2:36:D:THR:HB	2:38:D:ILE:HD11	14	0.3
(1,850)	2:36:D:THR:HB	2:38:D:ILE:HD12	14	0.3
(1,850)	2:36:D:THR:HB	2:38:D:ILE:HD13	14	0.3
(1,836)	1:27:C:VAL:HG11	1:28:C:PRO:HD3	7	0.3
(1,836)	1:27:C:VAL:HG12	1:28:C:PRO:HD3	7	0.3
(1,836)	1:27:C:VAL:HG13	1:28:C:PRO:HD3	7	0.3
(1,820)	1:22:C:GLY:H	2:41:D:TYR:HD1	7	0.3
(1,820)	1:22:C:GLY:H	2:41:D:TYR:HD2	7	0.3
(1,811)	1:27:C:VAL:H	2:40:D:PHE:HB2	17	0.3
(1,798)	1:27:C:VAL:HG11	1:28:C:PRO:HB3	14	0.3
(1,798)	1:27:C:VAL:HG12	1:28:C:PRO:HB3	14	0.3
(1,798)	1:27:C:VAL:HG13	1:28:C:PRO:HB3	14	0.3
(1,790)	1:31:C:PHE:HD1	2:36:D:THR:HG21	19	0.3
(1,790)	1:31:C:PHE:HD1	2:36:D:THR:HG22	19	0.3
(1,790)	1:31:C:PHE:HD1	2:36:D:THR:HG23	19	0.3
(1,790)	1:31:C:PHE:HD2	2:36:D:THR:HG21	19	0.3
(1,790)	1:31:C:PHE:HD2	2:36:D:THR:HG22	19	0.3
(1,790)	1:31:C:PHE:HD2	2:36:D:THR:HG23	19	0.3
(1,783)	1:29:C:GLU:HB3	2:41:D:TYR:HD1	20	0.3
(1,783)	1:29:C:GLU:HB3	2:41:D:TYR:HD2	20	0.3
(1,779)	1:29:C:GLU:HG3	2:40:D:PHE:HA	18	0.3
(1,775)	1:29:C:GLU:HG2	2:40:D:PHE:HA	4	0.3
(1,775)	1:29:C:GLU:HG2	2:40:D:PHE:HA	15	0.3
(1,759)	1:25:C:GLY:H	2:41:D:TYR:H	18	0.3
(1,739)	1:31:C:PHE:HD1	2:35:D:GLY:HA3	4	0.3
(1,739)	1:31:C:PHE:HD2	2:35:D:GLY:HA3	4	0.3
(1,724)	1:27:C:VAL:HG11	1:28:C:PRO:HG2	14	0.3
(1,724)	1:27:C:VAL:HG11	1:28:C:PRO:HG3	14	0.3
(1,724)	1:27:C:VAL:HG12	1:28:C:PRO:HG2	14	0.3
(1,724)	1:27:C:VAL:HG12	1:28:C:PRO:HG3	14	0.3
(1,724)	1:27:C:VAL:HG13	1:28:C:PRO:HG2	14	0.3
(1,724)	1:27:C:VAL:HG13	1:28:C:PRO:HG3	14	0.3
(1,686)	2:34:D:CYS:HB3	2:35:D:GLY:HA3	7	0.3
(1,686)	2:34:D:CYS:HB3	2:35:D:GLY:HA3	8	0.3
(1,348)	1:11:A:VAL:HA	1:12:A:ARG:HD2	7	0.3
(1,348)	1:11:A:VAL:HA	1:12:A:ARG:HD3	7	0.3

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,29)	1:11:A:VAL:HA	1:12:A:ARG:HD2	7	0.3
(1,29)	1:11:A:VAL:HA	1:12:A:ARG:HD3	7	0.3
(1,1293)	2:40:D:PHE:H	2:40:D:PHE:HE1	7	0.29
(1,1293)	2:40:D:PHE:H	2:40:D:PHE:HE2	7	0.29
(1,1170)	1:31:C:PHE:HB2	1:32:C:VAL:HG11	19	0.29
(1,1170)	1:31:C:PHE:HB2	1:32:C:VAL:HG12	19	0.29
(1,1170)	1:31:C:PHE:HB2	1:32:C:VAL:HG13	19	0.29
(1,1170)	1:31:C:PHE:HB2	1:32:C:VAL:HG21	19	0.29
(1,1170)	1:31:C:PHE:HB2	1:32:C:VAL:HG22	19	0.29
(1,1170)	1:31:C:PHE:HB2	1:32:C:VAL:HG23	19	0.29
(1,1166)	1:24:C:ALA:H	2:40:D:PHE:HD1	1	0.29
(1,1166)	1:24:C:ALA:H	2:40:D:PHE:HD2	1	0.29
(1,1154)	2:40:D:PHE:H	2:40:D:PHE:HE1	7	0.29
(1,1154)	2:40:D:PHE:H	2:40:D:PHE:HE2	7	0.29
(1,1134)	2:39:D:SER:HB3	2:40:D:PHE:H	13	0.29
(1,1132)	2:39:D:SER:HB3	2:40:D:PHE:H	13	0.29
(1,1109)	1:31:C:PHE:HD1	2:36:D:THR:HG21	13	0.29
(1,1109)	1:31:C:PHE:HD1	2:36:D:THR:HG22	13	0.29
(1,1109)	1:31:C:PHE:HD1	2:36:D:THR:HG23	13	0.29
(1,1109)	1:31:C:PHE:HD2	2:36:D:THR:HG21	13	0.29
(1,1109)	1:31:C:PHE:HD2	2:36:D:THR:HG22	13	0.29
(1,1109)	1:31:C:PHE:HD2	2:36:D:THR:HG23	13	0.29
(1,1062)	1:32:C:VAL:HG11	1:33:C:ARG:HA	17	0.29
(1,1062)	1:32:C:VAL:HG12	1:33:C:ARG:HA	17	0.29
(1,1062)	1:32:C:VAL:HG13	1:33:C:ARG:HA	17	0.29
(1,1062)	1:32:C:VAL:HG21	1:33:C:ARG:HA	17	0.29
(1,1062)	1:32:C:VAL:HG22	1:33:C:ARG:HA	17	0.29
(1,1062)	1:32:C:VAL:HG23	1:33:C:ARG:HA	17	0.29
(1,1058)	1:31:C:PHE:HD1	2:35:D:GLY:HA3	11	0.29
(1,1058)	1:31:C:PHE:HD2	2:35:D:GLY:HA3	11	0.29
(1,1055)	1:28:C:PRO:HG2	1:31:C:PHE:HE1	13	0.29
(1,1055)	1:28:C:PRO:HG2	1:31:C:PHE:HE2	13	0.29
(1,1055)	1:28:C:PRO:HG3	1:31:C:PHE:HE1	13	0.29
(1,1055)	1:28:C:PRO:HG3	1:31:C:PHE:HE2	13	0.29
(1,1045)	1:27:C:VAL:HB	2:41:D:TYR:HD1	6	0.29
(1,1045)	1:27:C:VAL:HB	2:41:D:TYR:HD2	6	0.29
(1,1027)	1:24:C:ALA:H	2:40:D:PHE:HD1	1	0.29
(1,1027)	1:24:C:ALA:H	2:40:D:PHE:HD2	1	0.29
(1,1000)	1:31:C:PHE:HE1	2:37:D:PRO:HD2	11	0.29
(1,1000)	1:31:C:PHE:HE2	2:37:D:PRO:HD2	11	0.29
(1,974)	2:40:D:PHE:H	2:40:D:PHE:HE1	7	0.29
(1,974)	2:40:D:PHE:H	2:40:D:PHE:HE2	7	0.29

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,851)	1:31:C:PHE:HB2	1:32:C:VAL:HG11	19	0.29
(1,851)	1:31:C:PHE:HB2	1:32:C:VAL:HG12	19	0.29
(1,851)	1:31:C:PHE:HB2	1:32:C:VAL:HG13	19	0.29
(1,851)	1:31:C:PHE:HB2	1:32:C:VAL:HG21	19	0.29
(1,851)	1:31:C:PHE:HB2	1:32:C:VAL:HG22	19	0.29
(1,851)	1:31:C:PHE:HB2	1:32:C:VAL:HG23	19	0.29
(1,847)	1:24:C:ALA:H	2:40:D:PHE:HD1	1	0.29
(1,847)	1:24:C:ALA:H	2:40:D:PHE:HD2	1	0.29
(1,835)	2:40:D:PHE:H	2:40:D:PHE:HE1	7	0.29
(1,835)	2:40:D:PHE:H	2:40:D:PHE:HE2	7	0.29
(1,815)	2:39:D:SER:HB3	2:40:D:PHE:H	13	0.29
(1,813)	2:39:D:SER:HB3	2:40:D:PHE:H	13	0.29
(1,790)	1:31:C:PHE:HD1	2:36:D:THR:HG21	13	0.29
(1,790)	1:31:C:PHE:HD1	2:36:D:THR:HG22	13	0.29
(1,790)	1:31:C:PHE:HD1	2:36:D:THR:HG23	13	0.29
(1,790)	1:31:C:PHE:HD2	2:36:D:THR:HG21	13	0.29
(1,790)	1:31:C:PHE:HD2	2:36:D:THR:HG22	13	0.29
(1,790)	1:31:C:PHE:HD2	2:36:D:THR:HG23	13	0.29
(1,743)	1:32:C:VAL:HG11	1:33:C:ARG:HA	17	0.29
(1,743)	1:32:C:VAL:HG12	1:33:C:ARG:HA	17	0.29
(1,743)	1:32:C:VAL:HG13	1:33:C:ARG:HA	17	0.29
(1,743)	1:32:C:VAL:HG21	1:33:C:ARG:HA	17	0.29
(1,743)	1:32:C:VAL:HG22	1:33:C:ARG:HA	17	0.29
(1,743)	1:32:C:VAL:HG23	1:33:C:ARG:HA	17	0.29
(1,739)	1:31:C:PHE:HD1	2:35:D:GLY:HA3	11	0.29
(1,739)	1:31:C:PHE:HD2	2:35:D:GLY:HA3	11	0.29
(1,736)	1:28:C:PRO:HG2	1:31:C:PHE:HE1	13	0.29
(1,736)	1:28:C:PRO:HG2	1:31:C:PHE:HE2	13	0.29
(1,736)	1:28:C:PRO:HG3	1:31:C:PHE:HE1	13	0.29
(1,736)	1:28:C:PRO:HG3	1:31:C:PHE:HE2	13	0.29
(1,726)	1:27:C:VAL:HB	2:41:D:TYR:HD1	6	0.29
(1,726)	1:27:C:VAL:HB	2:41:D:TYR:HD2	6	0.29
(1,708)	1:24:C:ALA:H	2:40:D:PHE:HD1	1	0.29
(1,708)	1:24:C:ALA:H	2:40:D:PHE:HD2	1	0.29
(1,681)	1:31:C:PHE:HE1	2:37:D:PRO:HD2	11	0.29
(1,681)	1:31:C:PHE:HE2	2:37:D:PRO:HD2	11	0.29
(1,671)	2:17:B:ILE:HG13	2:40:D:PHE:HE1	9	0.29
(1,671)	2:17:B:ILE:HG13	2:40:D:PHE:HE2	9	0.29
(1,568)	1:10:A:PHE:H	1:10:A:PHE:HD1	14	0.29
(1,568)	1:10:A:PHE:H	1:10:A:PHE:HD2	14	0.29
(1,453)	1:10:A:PHE:HD1	2:15:B:THR:HG21	20	0.29
(1,453)	1:10:A:PHE:HD1	2:15:B:THR:HG22	20	0.29

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,453)	1:10:A:PHE:HD1	2:15:B:THR:HG23	20	0.29
(1,453)	1:10:A:PHE:HD2	2:15:B:THR:HG21	20	0.29
(1,453)	1:10:A:PHE:HD2	2:15:B:THR:HG22	20	0.29
(1,453)	1:10:A:PHE:HD2	2:15:B:THR:HG23	20	0.29
(1,452)	2:15:B:THR:HG21	2:16:B:PRO:HD3	20	0.29
(1,452)	2:15:B:THR:HG22	2:16:B:PRO:HD3	20	0.29
(1,452)	2:15:B:THR:HG23	2:16:B:PRO:HD3	20	0.29
(1,434)	1:3:A:ALA:HB1	1:4:A:GLY:HA3	6	0.29
(1,434)	1:3:A:ALA:HB2	1:4:A:GLY:HA3	6	0.29
(1,434)	1:3:A:ALA:HB3	1:4:A:GLY:HA3	6	0.29
(1,434)	1:3:A:ALA:HB1	1:4:A:GLY:HA3	12	0.29
(1,434)	1:3:A:ALA:HB2	1:4:A:GLY:HA3	12	0.29
(1,434)	1:3:A:ALA:HB3	1:4:A:GLY:HA3	12	0.29
(1,406)	1:11:A:VAL:HG11	1:12:A:ARG:HA	11	0.29
(1,406)	1:11:A:VAL:HG12	1:12:A:ARG:HA	11	0.29
(1,406)	1:11:A:VAL:HG13	1:12:A:ARG:HA	11	0.29
(1,406)	1:11:A:VAL:HG21	1:12:A:ARG:HA	11	0.29
(1,406)	1:11:A:VAL:HG22	1:12:A:ARG:HA	11	0.29
(1,406)	1:11:A:VAL:HG23	1:12:A:ARG:HA	11	0.29
(1,397)	2:18:B:SER:HB2	2:19:B:PHE:H	1	0.29
(1,374)	1:3:A:ALA:HB1	1:4:A:GLY:HA3	6	0.29
(1,374)	1:3:A:ALA:HB2	1:4:A:GLY:HA3	6	0.29
(1,374)	1:3:A:ALA:HB3	1:4:A:GLY:HA3	6	0.29
(1,374)	1:3:A:ALA:HB1	1:4:A:GLY:HA3	12	0.29
(1,374)	1:3:A:ALA:HB2	1:4:A:GLY:HA3	12	0.29
(1,374)	1:3:A:ALA:HB3	1:4:A:GLY:HA3	12	0.29
(1,349)	2:13:B:CYS:HB3	2:14:B:GLY:HA3	2	0.29
(1,249)	1:10:A:PHE:H	1:10:A:PHE:HD1	14	0.29
(1,249)	1:10:A:PHE:H	1:10:A:PHE:HD2	14	0.29
(1,134)	1:10:A:PHE:HD1	2:15:B:THR:HG21	20	0.29
(1,134)	1:10:A:PHE:HD1	2:15:B:THR:HG22	20	0.29
(1,134)	1:10:A:PHE:HD1	2:15:B:THR:HG23	20	0.29
(1,134)	1:10:A:PHE:HD2	2:15:B:THR:HG21	20	0.29
(1,134)	1:10:A:PHE:HD2	2:15:B:THR:HG22	20	0.29
(1,134)	1:10:A:PHE:HD2	2:15:B:THR:HG23	20	0.29
(1,133)	2:15:B:THR:HG21	2:16:B:PRO:HD3	20	0.29
(1,133)	2:15:B:THR:HG22	2:16:B:PRO:HD3	20	0.29
(1,133)	2:15:B:THR:HG23	2:16:B:PRO:HD3	20	0.29
(1,115)	1:3:A:ALA:HB1	1:4:A:GLY:HA3	6	0.29
(1,115)	1:3:A:ALA:HB2	1:4:A:GLY:HA3	6	0.29
(1,115)	1:3:A:ALA:HB3	1:4:A:GLY:HA3	6	0.29
(1,115)	1:3:A:ALA:HB1	1:4:A:GLY:HA3	12	0.29

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,115)	1:3:A:ALA:HB2	1:4:A:GLY:HA3	12	0.29
(1,115)	1:3:A:ALA:HB3	1:4:A:GLY:HA3	12	0.29
(1,87)	1:11:A:VAL:HG11	1:12:A:ARG:HA	11	0.29
(1,87)	1:11:A:VAL:HG12	1:12:A:ARG:HA	11	0.29
(1,87)	1:11:A:VAL:HG13	1:12:A:ARG:HA	11	0.29
(1,87)	1:11:A:VAL:HG21	1:12:A:ARG:HA	11	0.29
(1,87)	1:11:A:VAL:HG22	1:12:A:ARG:HA	11	0.29
(1,87)	1:11:A:VAL:HG23	1:12:A:ARG:HA	11	0.29
(1,78)	2:18:B:SER:HB2	2:19:B:PHE:H	1	0.29
(1,55)	1:3:A:ALA:HB1	1:4:A:GLY:HA3	6	0.29
(1,55)	1:3:A:ALA:HB2	1:4:A:GLY:HA3	6	0.29
(1,55)	1:3:A:ALA:HB3	1:4:A:GLY:HA3	6	0.29
(1,55)	1:3:A:ALA:HB1	1:4:A:GLY:HA3	12	0.29
(1,55)	1:3:A:ALA:HB2	1:4:A:GLY:HA3	12	0.29
(1,55)	1:3:A:ALA:HB3	1:4:A:GLY:HA3	12	0.29
(1,30)	2:13:B:CYS:HB3	2:14:B:GLY:HA3	2	0.29
(1,6)	2:17:B:ILE:HG13	2:40:D:PHE:HE1	9	0.29
(1,6)	2:17:B:ILE:HG13	2:40:D:PHE:HE2	9	0.29
(1,1169)	2:36:D:THR:HB	2:38:D:ILE:HD11	19	0.28
(1,1169)	2:36:D:THR:HB	2:38:D:ILE:HD12	19	0.28
(1,1169)	2:36:D:THR:HB	2:38:D:ILE:HD13	19	0.28
(1,1130)	1:27:C:VAL:H	2:40:D:PHE:HB2	20	0.28
(1,1117)	1:27:C:VAL:HG11	1:28:C:PRO:HB3	4	0.28
(1,1117)	1:27:C:VAL:HG12	1:28:C:PRO:HB3	4	0.28
(1,1117)	1:27:C:VAL:HG13	1:28:C:PRO:HB3	4	0.28
(1,1055)	1:28:C:PRO:HG2	1:31:C:PHE:HE1	10	0.28
(1,1055)	1:28:C:PRO:HG2	1:31:C:PHE:HE2	10	0.28
(1,1055)	1:28:C:PRO:HG3	1:31:C:PHE:HE1	10	0.28
(1,1055)	1:28:C:PRO:HG3	1:31:C:PHE:HE2	10	0.28
(1,1055)	1:28:C:PRO:HG2	1:31:C:PHE:HE1	12	0.28
(1,1055)	1:28:C:PRO:HG2	1:31:C:PHE:HE2	12	0.28
(1,1055)	1:28:C:PRO:HG3	1:31:C:PHE:HE1	12	0.28
(1,1055)	1:28:C:PRO:HG3	1:31:C:PHE:HE2	12	0.28
(1,1020)	1:27:C:VAL:HB	2:41:D:TYR:H	16	0.28
(1,1000)	1:31:C:PHE:HE1	2:37:D:PRO:HD2	13	0.28
(1,1000)	1:31:C:PHE:HE2	2:37:D:PRO:HD2	13	0.28
(1,850)	2:36:D:THR:HB	2:38:D:ILE:HD11	19	0.28
(1,850)	2:36:D:THR:HB	2:38:D:ILE:HD12	19	0.28
(1,850)	2:36:D:THR:HB	2:38:D:ILE:HD13	19	0.28
(1,811)	1:27:C:VAL:H	2:40:D:PHE:HB2	20	0.28
(1,798)	1:27:C:VAL:HG11	1:28:C:PRO:HB3	4	0.28
(1,798)	1:27:C:VAL:HG12	1:28:C:PRO:HB3	4	0.28

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,798)	1:27:C:VAL:HG13	1:28:C:PRO:HB3	4	0.28
(1,736)	1:28:C:PRO:HG2	1:31:C:PHE:HE1	10	0.28
(1,736)	1:28:C:PRO:HG2	1:31:C:PHE:HE2	10	0.28
(1,736)	1:28:C:PRO:HG3	1:31:C:PHE:HE1	10	0.28
(1,736)	1:28:C:PRO:HG3	1:31:C:PHE:HE2	10	0.28
(1,736)	1:28:C:PRO:HG2	1:31:C:PHE:HE1	12	0.28
(1,736)	1:28:C:PRO:HG2	1:31:C:PHE:HE2	12	0.28
(1,736)	1:28:C:PRO:HG3	1:31:C:PHE:HE1	12	0.28
(1,736)	1:28:C:PRO:HG3	1:31:C:PHE:HE2	12	0.28
(1,701)	1:27:C:VAL:HB	2:41:D:TYR:H	16	0.28
(1,681)	1:31:C:PHE:HE1	2:37:D:PRO:HD2	13	0.28
(1,681)	1:31:C:PHE:HE2	2:37:D:PRO:HD2	13	0.28
(1,587)	1:12:A:ARG:H	1:12:A:ARG:HE	1	0.28
(1,569)	1:10:A:PHE:HD1	1:11:A:VAL:H	9	0.28
(1,569)	1:10:A:PHE:HD2	1:11:A:VAL:H	9	0.28
(1,562)	1:9:A:TYR:HA	1:9:A:TYR:HE1	10	0.28
(1,562)	1:9:A:TYR:HA	1:9:A:TYR:HE2	10	0.28
(1,514)	1:10:A:PHE:HB2	1:11:A:VAL:HG11	5	0.28
(1,514)	1:10:A:PHE:HB2	1:11:A:VAL:HG12	5	0.28
(1,514)	1:10:A:PHE:HB2	1:11:A:VAL:HG13	5	0.28
(1,514)	1:10:A:PHE:HB2	1:11:A:VAL:HG21	5	0.28
(1,514)	1:10:A:PHE:HB2	1:11:A:VAL:HG22	5	0.28
(1,514)	1:10:A:PHE:HB2	1:11:A:VAL:HG23	5	0.28
(1,499)	1:6:A:VAL:HG11	1:7:A:PRO:HD3	7	0.28
(1,499)	1:6:A:VAL:HG12	1:7:A:PRO:HD3	7	0.28
(1,499)	1:6:A:VAL:HG13	1:7:A:PRO:HD3	7	0.28
(1,499)	1:6:A:VAL:HG11	1:7:A:PRO:HD3	11	0.28
(1,499)	1:6:A:VAL:HG12	1:7:A:PRO:HD3	11	0.28
(1,499)	1:6:A:VAL:HG13	1:7:A:PRO:HD3	11	0.28
(1,478)	2:18:B:SER:HB3	2:19:B:PHE:H	5	0.28
(1,476)	2:18:B:SER:HB3	2:19:B:PHE:H	5	0.28
(1,463)	1:5:A:HIS:HB2	1:6:A:VAL:HG11	20	0.28
(1,463)	1:5:A:HIS:HB2	1:6:A:VAL:HG12	20	0.28
(1,463)	1:5:A:HIS:HB2	1:6:A:VAL:HG13	20	0.28
(1,446)	1:8:A:GLU:HB3	2:20:B:TYR:HD1	20	0.28
(1,446)	1:8:A:GLU:HB3	2:20:B:TYR:HD2	20	0.28
(1,439)	1:3:A:ALA:HA	1:4:A:GLY:H	3	0.28
(1,436)	1:8:A:GLU:HG2	2:20:B:TYR:HD1	11	0.28
(1,436)	1:8:A:GLU:HG2	2:20:B:TYR:HD2	11	0.28
(1,434)	1:3:A:ALA:HB1	1:4:A:GLY:HA3	3	0.28
(1,434)	1:3:A:ALA:HB2	1:4:A:GLY:HA3	3	0.28
(1,434)	1:3:A:ALA:HB3	1:4:A:GLY:HA3	3	0.28

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,411)	1:5:A:HIS:HB3	1:6:A:VAL:HG21	8	0.28
(1,411)	1:5:A:HIS:HB3	1:6:A:VAL:HG22	8	0.28
(1,411)	1:5:A:HIS:HB3	1:6:A:VAL:HG23	8	0.28
(1,403)	1:7:A:PRO:HG2	2:19:B:PHE:HD1	19	0.28
(1,403)	1:7:A:PRO:HG2	2:19:B:PHE:HD2	19	0.28
(1,403)	1:7:A:PRO:HG3	2:19:B:PHE:HD1	19	0.28
(1,403)	1:7:A:PRO:HG3	2:19:B:PHE:HD2	19	0.28
(1,398)	1:7:A:PRO:HA	2:19:B:PHE:HD1	20	0.28
(1,398)	1:7:A:PRO:HA	2:19:B:PHE:HD2	20	0.28
(1,379)	1:1:A:GLY:H	2:20:B:TYR:HE1	7	0.28
(1,379)	1:1:A:GLY:H	2:20:B:TYR:HE2	7	0.28
(1,374)	1:3:A:ALA:HB1	1:4:A:GLY:HA3	3	0.28
(1,374)	1:3:A:ALA:HB2	1:4:A:GLY:HA3	3	0.28
(1,374)	1:3:A:ALA:HB3	1:4:A:GLY:HA3	3	0.28
(1,268)	1:12:A:ARG:H	1:12:A:ARG:HE	1	0.28
(1,250)	1:10:A:PHE:HD1	1:11:A:VAL:H	9	0.28
(1,250)	1:10:A:PHE:HD2	1:11:A:VAL:H	9	0.28
(1,243)	1:9:A:TYR:HA	1:9:A:TYR:HE1	10	0.28
(1,243)	1:9:A:TYR:HA	1:9:A:TYR:HE2	10	0.28
(1,195)	1:10:A:PHE:HB2	1:11:A:VAL:HG11	5	0.28
(1,195)	1:10:A:PHE:HB2	1:11:A:VAL:HG12	5	0.28
(1,195)	1:10:A:PHE:HB2	1:11:A:VAL:HG13	5	0.28
(1,195)	1:10:A:PHE:HB2	1:11:A:VAL:HG21	5	0.28
(1,195)	1:10:A:PHE:HB2	1:11:A:VAL:HG22	5	0.28
(1,195)	1:10:A:PHE:HB2	1:11:A:VAL:HG23	5	0.28
(1,180)	1:6:A:VAL:HG11	1:7:A:PRO:HD3	7	0.28
(1,180)	1:6:A:VAL:HG12	1:7:A:PRO:HD3	7	0.28
(1,180)	1:6:A:VAL:HG13	1:7:A:PRO:HD3	7	0.28
(1,180)	1:6:A:VAL:HG11	1:7:A:PRO:HD3	11	0.28
(1,180)	1:6:A:VAL:HG12	1:7:A:PRO:HD3	11	0.28
(1,180)	1:6:A:VAL:HG13	1:7:A:PRO:HD3	11	0.28
(1,159)	2:18:B:SER:HB3	2:19:B:PHE:H	5	0.28
(1,157)	2:18:B:SER:HB3	2:19:B:PHE:H	5	0.28
(1,144)	1:5:A:HIS:HB2	1:6:A:VAL:HG11	20	0.28
(1,144)	1:5:A:HIS:HB2	1:6:A:VAL:HG12	20	0.28
(1,144)	1:5:A:HIS:HB2	1:6:A:VAL:HG13	20	0.28
(1,127)	1:8:A:GLU:HB3	2:20:B:TYR:HD1	20	0.28
(1,127)	1:8:A:GLU:HB3	2:20:B:TYR:HD2	20	0.28
(1,120)	1:3:A:ALA:HA	1:4:A:GLY:H	3	0.28
(1,117)	1:8:A:GLU:HG2	2:20:B:TYR:HD1	11	0.28
(1,117)	1:8:A:GLU:HG2	2:20:B:TYR:HD2	11	0.28
(1,115)	1:3:A:ALA:HB1	1:4:A:GLY:HA3	3	0.28

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,115)	1:3:A:ALA:HB2	1:4:A:GLY:HA3	3	0.28
(1,115)	1:3:A:ALA:HB3	1:4:A:GLY:HA3	3	0.28
(1,92)	1:5:A:HIS:HB3	1:6:A:VAL:HG21	8	0.28
(1,92)	1:5:A:HIS:HB3	1:6:A:VAL:HG22	8	0.28
(1,92)	1:5:A:HIS:HB3	1:6:A:VAL:HG23	8	0.28
(1,84)	1:7:A:PRO:HG2	2:19:B:PHE:HD1	19	0.28
(1,84)	1:7:A:PRO:HG2	2:19:B:PHE:HD2	19	0.28
(1,84)	1:7:A:PRO:HG3	2:19:B:PHE:HD1	19	0.28
(1,84)	1:7:A:PRO:HG3	2:19:B:PHE:HD2	19	0.28
(1,79)	1:7:A:PRO:HA	2:19:B:PHE:HD1	20	0.28
(1,79)	1:7:A:PRO:HA	2:19:B:PHE:HD2	20	0.28
(1,60)	1:1:A:GLY:H	2:20:B:TYR:HE1	7	0.28
(1,60)	1:1:A:GLY:H	2:20:B:TYR:HE2	7	0.28
(1,55)	1:3:A:ALA:HB1	1:4:A:GLY:HA3	3	0.28
(1,55)	1:3:A:ALA:HB2	1:4:A:GLY:HA3	3	0.28
(1,55)	1:3:A:ALA:HB3	1:4:A:GLY:HA3	3	0.28
(1,1283)	2:38:D:ILE:H	2:38:D:ILE:HB	9	0.27
(1,1249)	1:33:C:ARG:HB2	1:33:C:ARG:HE	12	0.27
(1,1243)	1:33:C:ARG:H	1:33:C:ARG:HE	8	0.27
(1,1242)	1:33:C:ARG:H	1:33:C:ARG:HB2	8	0.27
(1,1242)	1:33:C:ARG:H	1:33:C:ARG:HB3	8	0.27
(1,1242)	1:33:C:ARG:H	1:33:C:ARG:HB2	9	0.27
(1,1242)	1:33:C:ARG:H	1:33:C:ARG:HB3	9	0.27
(1,1218)	1:30:C:TYR:HA	1:30:C:TYR:HE1	6	0.27
(1,1218)	1:30:C:TYR:HA	1:30:C:TYR:HE2	6	0.27
(1,1155)	1:27:C:VAL:HG11	1:28:C:PRO:HD3	10	0.27
(1,1155)	1:27:C:VAL:HG12	1:28:C:PRO:HD3	10	0.27
(1,1155)	1:27:C:VAL:HG13	1:28:C:PRO:HD3	10	0.27
(1,1130)	1:27:C:VAL:H	2:40:D:PHE:HB2	15	0.27
(1,1109)	1:31:C:PHE:HD1	2:36:D:THR:HG21	12	0.27
(1,1109)	1:31:C:PHE:HD1	2:36:D:THR:HG22	12	0.27
(1,1109)	1:31:C:PHE:HD1	2:36:D:THR:HG23	12	0.27
(1,1109)	1:31:C:PHE:HD2	2:36:D:THR:HG21	12	0.27
(1,1109)	1:31:C:PHE:HD2	2:36:D:THR:HG22	12	0.27
(1,1109)	1:31:C:PHE:HD2	2:36:D:THR:HG23	12	0.27
(1,1100)	2:40:D:PHE:HA	2:41:D:TYR:HD1	18	0.27
(1,1100)	2:40:D:PHE:HA	2:41:D:TYR:HD2	18	0.27
(1,1090)	1:24:C:ALA:HB1	1:25:C:GLY:HA3	7	0.27
(1,1090)	1:24:C:ALA:HB2	1:25:C:GLY:HA3	7	0.27
(1,1090)	1:24:C:ALA:HB3	1:25:C:GLY:HA3	7	0.27
(1,1067)	1:26:C:HIS:HB3	1:27:C:VAL:HG21	4	0.27
(1,1067)	1:26:C:HIS:HB3	1:27:C:VAL:HG22	4	0.27

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,1067)	1:26:C:HIS:HB3	1:27:C:VAL:HG23	4	0.27
(1,1061)	1:22:C:GLY:H	2:39:D:SER:HB2	18	0.27
(1,1059)	1:28:C:PRO:HG2	2:40:D:PHE:HD1	9	0.27
(1,1059)	1:28:C:PRO:HG2	2:40:D:PHE:HD2	9	0.27
(1,1059)	1:28:C:PRO:HG3	2:40:D:PHE:HD1	9	0.27
(1,1059)	1:28:C:PRO:HG3	2:40:D:PHE:HD2	9	0.27
(1,1046)	1:27:C:VAL:HB	2:41:D:TYR:HE1	14	0.27
(1,1046)	1:27:C:VAL:HB	2:41:D:TYR:HE2	14	0.27
(1,1042)	1:25:C:GLY:H	2:40:D:PHE:HD1	4	0.27
(1,1042)	1:25:C:GLY:H	2:40:D:PHE:HD2	4	0.27
(1,1030)	1:24:C:ALA:HB1	1:25:C:GLY:HA3	7	0.27
(1,1030)	1:24:C:ALA:HB2	1:25:C:GLY:HA3	7	0.27
(1,1030)	1:24:C:ALA:HB3	1:25:C:GLY:HA3	7	0.27
(1,1013)	1:32:C:VAL:HA	1:33:C:ARG:H	1	0.27
(1,1008)	2:18:B:SER:HA	2:38:D:ILE:HG13	6	0.27
(1,1000)	1:31:C:PHE:HE1	2:37:D:PRO:HD2	16	0.27
(1,1000)	1:31:C:PHE:HE2	2:37:D:PRO:HD2	16	0.27
(1,964)	2:38:D:ILE:H	2:38:D:ILE:HB	9	0.27
(1,930)	1:33:C:ARG:HB2	1:33:C:ARG:HE	12	0.27
(1,924)	1:33:C:ARG:H	1:33:C:ARG:HE	8	0.27
(1,923)	1:33:C:ARG:H	1:33:C:ARG:HB2	8	0.27
(1,923)	1:33:C:ARG:H	1:33:C:ARG:HB3	8	0.27
(1,923)	1:33:C:ARG:H	1:33:C:ARG:HB2	9	0.27
(1,923)	1:33:C:ARG:H	1:33:C:ARG:HB3	9	0.27
(1,899)	1:30:C:TYR:HA	1:30:C:TYR:HE1	6	0.27
(1,899)	1:30:C:TYR:HA	1:30:C:TYR:HE2	6	0.27
(1,836)	1:27:C:VAL:HG11	1:28:C:PRO:HD3	10	0.27
(1,836)	1:27:C:VAL:HG12	1:28:C:PRO:HD3	10	0.27
(1,836)	1:27:C:VAL:HG13	1:28:C:PRO:HD3	10	0.27
(1,811)	1:27:C:VAL:H	2:40:D:PHE:HB2	15	0.27
(1,790)	1:31:C:PHE:HD1	2:36:D:THR:HG21	12	0.27
(1,790)	1:31:C:PHE:HD1	2:36:D:THR:HG22	12	0.27
(1,790)	1:31:C:PHE:HD1	2:36:D:THR:HG23	12	0.27
(1,790)	1:31:C:PHE:HD2	2:36:D:THR:HG21	12	0.27
(1,790)	1:31:C:PHE:HD2	2:36:D:THR:HG22	12	0.27
(1,790)	1:31:C:PHE:HD2	2:36:D:THR:HG23	12	0.27
(1,781)	2:40:D:PHE:HA	2:41:D:TYR:HD1	18	0.27
(1,781)	2:40:D:PHE:HA	2:41:D:TYR:HD2	18	0.27
(1,771)	1:24:C:ALA:HB1	1:25:C:GLY:HA3	7	0.27
(1,771)	1:24:C:ALA:HB2	1:25:C:GLY:HA3	7	0.27
(1,771)	1:24:C:ALA:HB3	1:25:C:GLY:HA3	7	0.27
(1,748)	1:26:C:HIS:HB3	1:27:C:VAL:HG21	4	0.27

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,748)	1:26:C:HIS:HB3	1:27:C:VAL:HG22	4	0.27
(1,748)	1:26:C:HIS:HB3	1:27:C:VAL:HG23	4	0.27
(1,742)	1:22:C:GLY:H	2:39:D:SER:HB2	18	0.27
(1,740)	1:28:C:PRO:HG2	2:40:D:PHE:HD1	9	0.27
(1,740)	1:28:C:PRO:HG2	2:40:D:PHE:HD2	9	0.27
(1,740)	1:28:C:PRO:HG3	2:40:D:PHE:HD1	9	0.27
(1,740)	1:28:C:PRO:HG3	2:40:D:PHE:HD2	9	0.27
(1,727)	1:27:C:VAL:HB	2:41:D:TYR:HE1	14	0.27
(1,727)	1:27:C:VAL:HB	2:41:D:TYR:HE2	14	0.27
(1,723)	1:25:C:GLY:H	2:40:D:PHE:HD1	4	0.27
(1,723)	1:25:C:GLY:H	2:40:D:PHE:HD2	4	0.27
(1,711)	1:24:C:ALA:HB1	1:25:C:GLY:HA3	7	0.27
(1,711)	1:24:C:ALA:HB2	1:25:C:GLY:HA3	7	0.27
(1,711)	1:24:C:ALA:HB3	1:25:C:GLY:HA3	7	0.27
(1,694)	1:32:C:VAL:HA	1:33:C:ARG:H	1	0.27
(1,689)	2:18:B:SER:HA	2:38:D:ILE:HG13	6	0.27
(1,681)	1:31:C:PHE:HE1	2:37:D:PRO:HD2	16	0.27
(1,681)	1:31:C:PHE:HE2	2:37:D:PRO:HD2	16	0.27
(1,631)	2:17:B:ILE:HA	2:17:B:ILE:HD11	9	0.27
(1,631)	2:17:B:ILE:HA	2:17:B:ILE:HD12	9	0.27
(1,631)	2:17:B:ILE:HA	2:17:B:ILE:HD13	9	0.27
(1,474)	1:6:A:VAL:H	2:19:B:PHE:HB2	5	0.27
(1,465)	1:5:A:HIS:HD2	1:6:A:VAL:HG11	16	0.27
(1,465)	1:5:A:HIS:HD2	1:6:A:VAL:HG12	16	0.27
(1,465)	1:5:A:HIS:HD2	1:6:A:VAL:HG13	16	0.27
(1,411)	1:5:A:HIS:HB3	1:6:A:VAL:HG21	6	0.27
(1,411)	1:5:A:HIS:HB3	1:6:A:VAL:HG22	6	0.27
(1,411)	1:5:A:HIS:HB3	1:6:A:VAL:HG23	6	0.27
(1,369)	1:6:A:VAL:HG21	2:20:B:TYR:HE1	13	0.27
(1,369)	1:6:A:VAL:HG21	2:20:B:TYR:HE2	13	0.27
(1,369)	1:6:A:VAL:HG22	2:20:B:TYR:HE1	13	0.27
(1,369)	1:6:A:VAL:HG22	2:20:B:TYR:HE2	13	0.27
(1,369)	1:6:A:VAL:HG23	2:20:B:TYR:HE1	13	0.27
(1,369)	1:6:A:VAL:HG23	2:20:B:TYR:HE2	13	0.27
(1,362)	1:7:A:PRO:HG2	2:19:B:PHE:HD1	17	0.27
(1,362)	1:7:A:PRO:HG2	2:19:B:PHE:HD2	17	0.27
(1,362)	1:7:A:PRO:HG3	2:19:B:PHE:HD1	17	0.27
(1,362)	1:7:A:PRO:HG3	2:19:B:PHE:HD2	17	0.27
(1,351)	2:16:B:PRO:HA	2:17:B:ILE:HD11	18	0.27
(1,351)	2:16:B:PRO:HA	2:17:B:ILE:HD12	18	0.27
(1,351)	2:16:B:PRO:HA	2:17:B:ILE:HD13	18	0.27
(1,346)	1:9:A:TYR:HE1	2:18:B:SER:HB2	6	0.27

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,346)	1:9:A:TYR:HE2	2:18:B:SER:HB2	6	0.27
(1,312)	2:17:B:ILE:HA	2:17:B:ILE:HD11	9	0.27
(1,312)	2:17:B:ILE:HA	2:17:B:ILE:HD12	9	0.27
(1,312)	2:17:B:ILE:HA	2:17:B:ILE:HD13	9	0.27
(1,155)	1:6:A:VAL:H	2:19:B:PHE:HB2	5	0.27
(1,146)	1:5:A:HIS:HD2	1:6:A:VAL:HG11	16	0.27
(1,146)	1:5:A:HIS:HD2	1:6:A:VAL:HG12	16	0.27
(1,146)	1:5:A:HIS:HD2	1:6:A:VAL:HG13	16	0.27
(1,92)	1:5:A:HIS:HB3	1:6:A:VAL:HG21	6	0.27
(1,92)	1:5:A:HIS:HB3	1:6:A:VAL:HG22	6	0.27
(1,92)	1:5:A:HIS:HB3	1:6:A:VAL:HG23	6	0.27
(1,50)	1:6:A:VAL:HG21	2:20:B:TYR:HE1	13	0.27
(1,50)	1:6:A:VAL:HG21	2:20:B:TYR:HE2	13	0.27
(1,50)	1:6:A:VAL:HG22	2:20:B:TYR:HE1	13	0.27
(1,50)	1:6:A:VAL:HG22	2:20:B:TYR:HE2	13	0.27
(1,50)	1:6:A:VAL:HG23	2:20:B:TYR:HE1	13	0.27
(1,50)	1:6:A:VAL:HG23	2:20:B:TYR:HE2	13	0.27
(1,43)	1:7:A:PRO:HG2	2:19:B:PHE:HD1	17	0.27
(1,43)	1:7:A:PRO:HG2	2:19:B:PHE:HD2	17	0.27
(1,43)	1:7:A:PRO:HG3	2:19:B:PHE:HD1	17	0.27
(1,43)	1:7:A:PRO:HG3	2:19:B:PHE:HD2	17	0.27
(1,32)	2:16:B:PRO:HA	2:17:B:ILE:HD11	18	0.27
(1,32)	2:16:B:PRO:HA	2:17:B:ILE:HD12	18	0.27
(1,32)	2:16:B:PRO:HA	2:17:B:ILE:HD13	18	0.27
(1,27)	1:9:A:TYR:HE1	2:18:B:SER:HB2	6	0.27
(1,27)	1:9:A:TYR:HE2	2:18:B:SER:HB2	6	0.27
(1,1186)	1:26:C:HIS:HA	1:26:C:HIS:HD2	1	0.26
(1,1186)	1:26:C:HIS:HA	1:26:C:HIS:HD2	8	0.26
(1,1139)	1:22:C:GLY:H	2:41:D:TYR:HD1	1	0.26
(1,1139)	1:22:C:GLY:H	2:41:D:TYR:HD2	1	0.26
(1,1137)	1:22:C:GLY:H	2:40:D:PHE:H	19	0.26
(1,1134)	2:39:D:SER:HB3	2:40:D:PHE:H	12	0.26
(1,1132)	2:39:D:SER:HB3	2:40:D:PHE:H	12	0.26
(1,1130)	1:27:C:VAL:H	2:40:D:PHE:HB2	2	0.26
(1,1109)	1:31:C:PHE:HD1	2:36:D:THR:HG21	4	0.26
(1,1109)	1:31:C:PHE:HD1	2:36:D:THR:HG22	4	0.26
(1,1109)	1:31:C:PHE:HD1	2:36:D:THR:HG23	4	0.26
(1,1109)	1:31:C:PHE:HD2	2:36:D:THR:HG21	4	0.26
(1,1109)	1:31:C:PHE:HD2	2:36:D:THR:HG22	4	0.26
(1,1109)	1:31:C:PHE:HD2	2:36:D:THR:HG23	4	0.26
(1,1104)	1:25:C:GLY:H	2:40:D:PHE:H	6	0.26
(1,1089)	1:29:C:GLU:HB2	2:41:D:TYR:HB2	14	0.26

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,1089)	1:29:C:GLU:HB2	2:41:D:TYR:HB3	14	0.26
(1,1061)	1:22:C:GLY:H	2:39:D:SER:HB2	20	0.26
(1,1060)	1:31:C:PHE:HD1	2:37:D:PRO:HA	20	0.26
(1,1060)	1:31:C:PHE:HD2	2:37:D:PRO:HA	20	0.26
(1,867)	1:26:C:HIS:HA	1:26:C:HIS:HD2	1	0.26
(1,867)	1:26:C:HIS:HA	1:26:C:HIS:HD2	8	0.26
(1,820)	1:22:C:GLY:H	2:41:D:TYR:HD1	1	0.26
(1,820)	1:22:C:GLY:H	2:41:D:TYR:HD2	1	0.26
(1,818)	1:22:C:GLY:H	2:40:D:PHE:H	19	0.26
(1,815)	2:39:D:SER:HB3	2:40:D:PHE:H	12	0.26
(1,813)	2:39:D:SER:HB3	2:40:D:PHE:H	12	0.26
(1,811)	1:27:C:VAL:H	2:40:D:PHE:HB2	2	0.26
(1,790)	1:31:C:PHE:HD1	2:36:D:THR:HG21	4	0.26
(1,790)	1:31:C:PHE:HD1	2:36:D:THR:HG22	4	0.26
(1,790)	1:31:C:PHE:HD1	2:36:D:THR:HG23	4	0.26
(1,790)	1:31:C:PHE:HD2	2:36:D:THR:HG21	4	0.26
(1,790)	1:31:C:PHE:HD2	2:36:D:THR:HG22	4	0.26
(1,790)	1:31:C:PHE:HD2	2:36:D:THR:HG23	4	0.26
(1,785)	1:25:C:GLY:H	2:40:D:PHE:H	6	0.26
(1,770)	1:29:C:GLU:HB2	2:41:D:TYR:HB2	14	0.26
(1,770)	1:29:C:GLU:HB2	2:41:D:TYR:HB3	14	0.26
(1,742)	1:22:C:GLY:H	2:39:D:SER:HB2	20	0.26
(1,741)	1:31:C:PHE:HD1	2:37:D:PRO:HA	20	0.26
(1,741)	1:31:C:PHE:HD2	2:37:D:PRO:HA	20	0.26
(1,575)	1:10:A:PHE:HE1	2:15:B:THR:HA	5	0.26
(1,575)	1:10:A:PHE:HE2	2:15:B:THR:HA	5	0.26
(1,493)	1:11:A:VAL:HB	1:12:A:ARG:H	12	0.26
(1,474)	1:6:A:VAL:H	2:19:B:PHE:HB2	16	0.26
(1,438)	1:8:A:GLU:HG2	2:19:B:PHE:HA	18	0.26
(1,401)	1:10:A:PHE:HD1	1:11:A:VAL:HA	9	0.26
(1,401)	1:10:A:PHE:HD2	1:11:A:VAL:HA	9	0.26
(1,397)	2:18:B:SER:HB2	2:19:B:PHE:H	4	0.26
(1,382)	1:10:A:PHE:HD1	2:16:B:PRO:HD2	5	0.26
(1,382)	1:10:A:PHE:HD2	2:16:B:PRO:HD2	5	0.26
(1,372)	2:18:B:SER:HA	2:19:B:PHE:HD1	6	0.26
(1,372)	2:18:B:SER:HA	2:19:B:PHE:HD2	6	0.26
(1,344)	1:10:A:PHE:HE1	2:16:B:PRO:HD2	4	0.26
(1,344)	1:10:A:PHE:HE2	2:16:B:PRO:HD2	4	0.26
(1,256)	1:10:A:PHE:HE1	2:15:B:THR:HA	5	0.26
(1,256)	1:10:A:PHE:HE2	2:15:B:THR:HA	5	0.26
(1,174)	1:11:A:VAL:HB	1:12:A:ARG:H	12	0.26
(1,155)	1:6:A:VAL:H	2:19:B:PHE:HB2	16	0.26

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,119)	1:8:A:GLU:HG2	2:19:B:PHE:HA	18	0.26
(1,82)	1:10:A:PHE:HD1	1:11:A:VAL:HA	9	0.26
(1,82)	1:10:A:PHE:HD2	1:11:A:VAL:HA	9	0.26
(1,78)	2:18:B:SER:HB2	2:19:B:PHE:H	4	0.26
(1,63)	1:10:A:PHE:HD1	2:16:B:PRO:HD2	5	0.26
(1,63)	1:10:A:PHE:HD2	2:16:B:PRO:HD2	5	0.26
(1,53)	2:18:B:SER:HA	2:19:B:PHE:HD1	6	0.26
(1,53)	2:18:B:SER:HA	2:19:B:PHE:HD2	6	0.26
(1,25)	1:10:A:PHE:HE1	2:16:B:PRO:HD2	4	0.26
(1,25)	1:10:A:PHE:HE2	2:16:B:PRO:HD2	4	0.26
(1,1175)	1:24:C:ALA:HB1	2:40:D:PHE:HB3	6	0.25
(1,1175)	1:24:C:ALA:HB2	2:40:D:PHE:HB3	6	0.25
(1,1175)	1:24:C:ALA:HB3	2:40:D:PHE:HB3	6	0.25
(1,1157)	1:25:C:GLY:H	1:27:C:VAL:H	7	0.25
(1,1139)	1:22:C:GLY:H	2:41:D:TYR:HD1	16	0.25
(1,1139)	1:22:C:GLY:H	2:41:D:TYR:HD2	16	0.25
(1,1109)	1:31:C:PHE:HD1	2:36:D:THR:HG21	7	0.25
(1,1109)	1:31:C:PHE:HD1	2:36:D:THR:HG22	7	0.25
(1,1109)	1:31:C:PHE:HD1	2:36:D:THR:HG23	7	0.25
(1,1109)	1:31:C:PHE:HD2	2:36:D:THR:HG21	7	0.25
(1,1109)	1:31:C:PHE:HD2	2:36:D:THR:HG22	7	0.25
(1,1109)	1:31:C:PHE:HD2	2:36:D:THR:HG23	7	0.25
(1,1088)	1:23:C:GLY:HA2	1:24:C:ALA:H	18	0.25
(1,1053)	2:39:D:SER:HB2	2:40:D:PHE:H	3	0.25
(1,1043)	1:27:C:VAL:HG11	1:28:C:PRO:HG2	6	0.25
(1,1043)	1:27:C:VAL:HG11	1:28:C:PRO:HG3	6	0.25
(1,1043)	1:27:C:VAL:HG12	1:28:C:PRO:HG2	6	0.25
(1,1043)	1:27:C:VAL:HG12	1:28:C:PRO:HG3	6	0.25
(1,1043)	1:27:C:VAL:HG13	1:28:C:PRO:HG2	6	0.25
(1,1043)	1:27:C:VAL:HG13	1:28:C:PRO:HG3	6	0.25
(1,1040)	1:31:C:PHE:HE1	2:37:D:PRO:HD2	4	0.25
(1,1040)	1:31:C:PHE:HE2	2:37:D:PRO:HD2	4	0.25
(1,1015)	1:30:C:TYR:HD1	2:39:D:SER:HA	6	0.25
(1,1015)	1:30:C:TYR:HD2	2:39:D:SER:HA	6	0.25
(1,1007)	2:37:D:PRO:HA	2:38:D:ILE:HD11	1	0.25
(1,1007)	2:37:D:PRO:HA	2:38:D:ILE:HD12	1	0.25
(1,1007)	2:37:D:PRO:HA	2:38:D:ILE:HD13	1	0.25
(1,856)	1:24:C:ALA:HB1	2:40:D:PHE:HB3	6	0.25
(1,856)	1:24:C:ALA:HB2	2:40:D:PHE:HB3	6	0.25
(1,856)	1:24:C:ALA:HB3	2:40:D:PHE:HB3	6	0.25
(1,838)	1:25:C:GLY:H	1:27:C:VAL:H	7	0.25
(1,820)	1:22:C:GLY:H	2:41:D:TYR:HD1	16	0.25

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,820)	1:22:C:GLY:H	2:41:D:TYR:HD2	16	0.25
(1,790)	1:31:C:PHE:HD1	2:36:D:THR:HG21	7	0.25
(1,790)	1:31:C:PHE:HD1	2:36:D:THR:HG22	7	0.25
(1,790)	1:31:C:PHE:HD1	2:36:D:THR:HG23	7	0.25
(1,790)	1:31:C:PHE:HD2	2:36:D:THR:HG21	7	0.25
(1,790)	1:31:C:PHE:HD2	2:36:D:THR:HG22	7	0.25
(1,790)	1:31:C:PHE:HD2	2:36:D:THR:HG23	7	0.25
(1,769)	1:23:C:GLY:HA2	1:24:C:ALA:H	18	0.25
(1,734)	2:39:D:SER:HB2	2:40:D:PHE:H	3	0.25
(1,724)	1:27:C:VAL:HG11	1:28:C:PRO:HG2	6	0.25
(1,724)	1:27:C:VAL:HG11	1:28:C:PRO:HG3	6	0.25
(1,724)	1:27:C:VAL:HG12	1:28:C:PRO:HG2	6	0.25
(1,724)	1:27:C:VAL:HG12	1:28:C:PRO:HG3	6	0.25
(1,724)	1:27:C:VAL:HG13	1:28:C:PRO:HG2	6	0.25
(1,724)	1:27:C:VAL:HG13	1:28:C:PRO:HG3	6	0.25
(1,721)	1:31:C:PHE:HE1	2:37:D:PRO:HD2	4	0.25
(1,721)	1:31:C:PHE:HE2	2:37:D:PRO:HD2	4	0.25
(1,696)	1:30:C:TYR:HD1	2:39:D:SER:HA	6	0.25
(1,696)	1:30:C:TYR:HD2	2:39:D:SER:HA	6	0.25
(1,688)	2:37:D:PRO:HA	2:38:D:ILE:HD11	1	0.25
(1,688)	2:37:D:PRO:HA	2:38:D:ILE:HD12	1	0.25
(1,688)	2:37:D:PRO:HA	2:38:D:ILE:HD13	1	0.25
(1,629)	2:17:B:ILE:H	2:17:B:ILE:HD11	16	0.25
(1,629)	2:17:B:ILE:H	2:17:B:ILE:HD12	16	0.25
(1,629)	2:17:B:ILE:H	2:17:B:ILE:HD13	16	0.25
(1,519)	1:3:A:ALA:HB1	2:19:B:PHE:HB3	18	0.25
(1,519)	1:3:A:ALA:HB2	2:19:B:PHE:HB3	18	0.25
(1,519)	1:3:A:ALA:HB3	2:19:B:PHE:HB3	18	0.25
(1,499)	1:6:A:VAL:HG11	1:7:A:PRO:HD3	9	0.25
(1,499)	1:6:A:VAL:HG12	1:7:A:PRO:HD3	9	0.25
(1,499)	1:6:A:VAL:HG13	1:7:A:PRO:HD3	9	0.25
(1,493)	1:11:A:VAL:HB	1:12:A:ARG:H	4	0.25
(1,478)	2:18:B:SER:HB3	2:19:B:PHE:H	7	0.25
(1,476)	2:18:B:SER:HB3	2:19:B:PHE:H	7	0.25
(1,470)	1:2:A:GLY:H	2:19:B:PHE:H	12	0.25
(1,452)	2:15:B:THR:HG21	2:16:B:PRO:HD3	18	0.25
(1,452)	2:15:B:THR:HG22	2:16:B:PRO:HD3	18	0.25
(1,452)	2:15:B:THR:HG23	2:16:B:PRO:HD3	18	0.25
(1,410)	1:2:A:GLY:HA3	1:3:A:ALA:HB1	13	0.25
(1,410)	1:2:A:GLY:HA3	1:3:A:ALA:HB2	13	0.25
(1,410)	1:2:A:GLY:HA3	1:3:A:ALA:HB3	13	0.25
(1,404)	1:10:A:PHE:HD1	2:16:B:PRO:HA	17	0.25

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,404)	1:10:A:PHE:HD2	2:16:B:PRO:HA	17	0.25
(1,359)	1:9:A:TYR:HD1	2:18:B:SER:HA	18	0.25
(1,359)	1:9:A:TYR:HD2	2:18:B:SER:HA	18	0.25
(1,310)	2:17:B:ILE:H	2:17:B:ILE:HD11	16	0.25
(1,310)	2:17:B:ILE:H	2:17:B:ILE:HD12	16	0.25
(1,310)	2:17:B:ILE:H	2:17:B:ILE:HD13	16	0.25
(1,200)	1:3:A:ALA:HB1	2:19:B:PHE:HB3	18	0.25
(1,200)	1:3:A:ALA:HB2	2:19:B:PHE:HB3	18	0.25
(1,200)	1:3:A:ALA:HB3	2:19:B:PHE:HB3	18	0.25
(1,180)	1:6:A:VAL:HG11	1:7:A:PRO:HD3	9	0.25
(1,180)	1:6:A:VAL:HG12	1:7:A:PRO:HD3	9	0.25
(1,180)	1:6:A:VAL:HG13	1:7:A:PRO:HD3	9	0.25
(1,174)	1:11:A:VAL:HB	1:12:A:ARG:H	4	0.25
(1,159)	2:18:B:SER:HB3	2:19:B:PHE:H	7	0.25
(1,157)	2:18:B:SER:HB3	2:19:B:PHE:H	7	0.25
(1,151)	1:2:A:GLY:H	2:19:B:PHE:H	12	0.25
(1,133)	2:15:B:THR:HG21	2:16:B:PRO:HD3	18	0.25
(1,133)	2:15:B:THR:HG22	2:16:B:PRO:HD3	18	0.25
(1,133)	2:15:B:THR:HG23	2:16:B:PRO:HD3	18	0.25
(1,91)	1:2:A:GLY:HA3	1:3:A:ALA:HB1	13	0.25
(1,91)	1:2:A:GLY:HA3	1:3:A:ALA:HB2	13	0.25
(1,91)	1:2:A:GLY:HA3	1:3:A:ALA:HB3	13	0.25
(1,85)	1:10:A:PHE:HD1	2:16:B:PRO:HA	17	0.25
(1,85)	1:10:A:PHE:HD2	2:16:B:PRO:HA	17	0.25
(1,40)	1:9:A:TYR:HD1	2:18:B:SER:HA	18	0.25
(1,40)	1:9:A:TYR:HD2	2:18:B:SER:HA	18	0.25
(1,1252)	1:28:C:PRO:HG2	1:31:C:PHE:HE1	13	0.24
(1,1252)	1:28:C:PRO:HG2	1:31:C:PHE:HE2	13	0.24
(1,1252)	1:28:C:PRO:HG3	1:31:C:PHE:HE1	13	0.24
(1,1252)	1:28:C:PRO:HG3	1:31:C:PHE:HE2	13	0.24
(1,1218)	1:30:C:TYR:HA	1:30:C:TYR:HE1	11	0.24
(1,1218)	1:30:C:TYR:HA	1:30:C:TYR:HE2	11	0.24
(1,1170)	1:31:C:PHE:HB2	1:32:C:VAL:HG11	10	0.24
(1,1170)	1:31:C:PHE:HB2	1:32:C:VAL:HG12	10	0.24
(1,1170)	1:31:C:PHE:HB2	1:32:C:VAL:HG13	10	0.24
(1,1170)	1:31:C:PHE:HB2	1:32:C:VAL:HG21	10	0.24
(1,1170)	1:31:C:PHE:HB2	1:32:C:VAL:HG22	10	0.24
(1,1170)	1:31:C:PHE:HB2	1:32:C:VAL:HG23	10	0.24
(1,1157)	1:25:C:GLY:H	1:27:C:VAL:H	8	0.24
(1,1130)	1:27:C:VAL:H	2:40:D:PHE:HB2	8	0.24
(1,1121)	1:26:C:HIS:HD2	1:27:C:VAL:HG11	3	0.24
(1,1121)	1:26:C:HIS:HD2	1:27:C:VAL:HG12	3	0.24

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,1121)	1:26:C:HIS:HD2	1:27:C:VAL:HG13	3	0.24
(1,1101)	1:25:C:GLY:HA2	1:27:C:VAL:H	9	0.24
(1,1089)	1:29:C:GLU:HB2	2:41:D:TYR:HB2	15	0.24
(1,1089)	1:29:C:GLU:HB2	2:41:D:TYR:HB3	15	0.24
(1,1088)	1:23:C:GLY:HA2	1:24:C:ALA:H	5	0.24
(1,1043)	1:27:C:VAL:HG11	1:28:C:PRO:HG2	5	0.24
(1,1043)	1:27:C:VAL:HG11	1:28:C:PRO:HG3	5	0.24
(1,1043)	1:27:C:VAL:HG12	1:28:C:PRO:HG2	5	0.24
(1,1043)	1:27:C:VAL:HG12	1:28:C:PRO:HG3	5	0.24
(1,1043)	1:27:C:VAL:HG13	1:28:C:PRO:HG2	5	0.24
(1,1043)	1:27:C:VAL:HG13	1:28:C:PRO:HG3	5	0.24
(1,1040)	1:31:C:PHE:HE1	2:37:D:PRO:HD2	18	0.24
(1,1040)	1:31:C:PHE:HE2	2:37:D:PRO:HD2	18	0.24
(1,1012)	1:29:C:GLU:HG2	2:41:D:TYR:HE1	5	0.24
(1,1012)	1:29:C:GLU:HG2	2:41:D:TYR:HE2	5	0.24
(1,1000)	1:31:C:PHE:HE1	2:37:D:PRO:HD2	6	0.24
(1,1000)	1:31:C:PHE:HE2	2:37:D:PRO:HD2	6	0.24
(1,933)	1:28:C:PRO:HG2	1:31:C:PHE:HE1	13	0.24
(1,933)	1:28:C:PRO:HG2	1:31:C:PHE:HE2	13	0.24
(1,933)	1:28:C:PRO:HG3	1:31:C:PHE:HE1	13	0.24
(1,933)	1:28:C:PRO:HG3	1:31:C:PHE:HE2	13	0.24
(1,899)	1:30:C:TYR:HA	1:30:C:TYR:HE1	11	0.24
(1,899)	1:30:C:TYR:HA	1:30:C:TYR:HE2	11	0.24
(1,851)	1:31:C:PHE:HB2	1:32:C:VAL:HG11	10	0.24
(1,851)	1:31:C:PHE:HB2	1:32:C:VAL:HG12	10	0.24
(1,851)	1:31:C:PHE:HB2	1:32:C:VAL:HG13	10	0.24
(1,851)	1:31:C:PHE:HB2	1:32:C:VAL:HG21	10	0.24
(1,851)	1:31:C:PHE:HB2	1:32:C:VAL:HG22	10	0.24
(1,851)	1:31:C:PHE:HB2	1:32:C:VAL:HG23	10	0.24
(1,838)	1:25:C:GLY:H	1:27:C:VAL:H	8	0.24
(1,811)	1:27:C:VAL:H	2:40:D:PHE:HB2	8	0.24
(1,802)	1:26:C:HIS:HD2	1:27:C:VAL:HG11	3	0.24
(1,802)	1:26:C:HIS:HD2	1:27:C:VAL:HG12	3	0.24
(1,802)	1:26:C:HIS:HD2	1:27:C:VAL:HG13	3	0.24
(1,782)	1:25:C:GLY:HA2	1:27:C:VAL:H	9	0.24
(1,770)	1:29:C:GLU:HB2	2:41:D:TYR:HB2	15	0.24
(1,770)	1:29:C:GLU:HB2	2:41:D:TYR:HB3	15	0.24
(1,769)	1:23:C:GLY:HA2	1:24:C:ALA:H	5	0.24
(1,724)	1:27:C:VAL:HG11	1:28:C:PRO:HG2	5	0.24
(1,724)	1:27:C:VAL:HG11	1:28:C:PRO:HG3	5	0.24
(1,724)	1:27:C:VAL:HG12	1:28:C:PRO:HG2	5	0.24
(1,724)	1:27:C:VAL:HG12	1:28:C:PRO:HG3	5	0.24

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,724)	1:27:C:VAL:HG13	1:28:C:PRO:HG2	5	0.24
(1,724)	1:27:C:VAL:HG13	1:28:C:PRO:HG3	5	0.24
(1,721)	1:31:C:PHE:HE1	2:37:D:PRO:HD2	18	0.24
(1,721)	1:31:C:PHE:HE2	2:37:D:PRO:HD2	18	0.24
(1,693)	1:29:C:GLU:HG2	2:41:D:TYR:HE1	5	0.24
(1,693)	1:29:C:GLU:HG2	2:41:D:TYR:HE2	5	0.24
(1,681)	1:31:C:PHE:HE1	2:37:D:PRO:HD2	6	0.24
(1,681)	1:31:C:PHE:HE2	2:37:D:PRO:HD2	6	0.24
(1,586)	1:12:A:ARG:H	1:12:A:ARG:HB2	10	0.24
(1,586)	1:12:A:ARG:H	1:12:A:ARG:HB3	10	0.24
(1,586)	1:12:A:ARG:H	1:12:A:ARG:HB2	12	0.24
(1,586)	1:12:A:ARG:H	1:12:A:ARG:HB3	12	0.24
(1,586)	1:12:A:ARG:H	1:12:A:ARG:HB2	17	0.24
(1,586)	1:12:A:ARG:H	1:12:A:ARG:HB3	17	0.24
(1,530)	1:5:A:HIS:HA	1:5:A:HIS:HD2	14	0.24
(1,478)	2:18:B:SER:HB3	2:19:B:PHE:H	9	0.24
(1,476)	2:18:B:SER:HB3	2:19:B:PHE:H	9	0.24
(1,461)	1:6:A:VAL:HG11	1:7:A:PRO:HB3	6	0.24
(1,461)	1:6:A:VAL:HG12	1:7:A:PRO:HB3	6	0.24
(1,461)	1:6:A:VAL:HG13	1:7:A:PRO:HB3	6	0.24
(1,448)	1:4:A:GLY:H	2:19:B:PHE:H	1	0.24
(1,437)	1:8:A:GLU:HG2	2:20:B:TYR:HB2	11	0.24
(1,437)	1:8:A:GLU:HG2	2:20:B:TYR:HB3	11	0.24
(1,422)	1:4:A:GLY:H	2:20:B:TYR:H	15	0.24
(1,380)	2:13:B:CYS:HB2	2:14:B:GLY:HA3	9	0.24
(1,362)	1:7:A:PRO:HG2	2:19:B:PHE:HD1	6	0.24
(1,362)	1:7:A:PRO:HG2	2:19:B:PHE:HD2	6	0.24
(1,362)	1:7:A:PRO:HG3	2:19:B:PHE:HD1	6	0.24
(1,362)	1:7:A:PRO:HG3	2:19:B:PHE:HD2	6	0.24
(1,356)	1:8:A:GLU:HG2	2:20:B:TYR:HE1	3	0.24
(1,356)	1:8:A:GLU:HG2	2:20:B:TYR:HE2	3	0.24
(1,344)	1:10:A:PHE:HE1	2:16:B:PRO:HD2	2	0.24
(1,344)	1:10:A:PHE:HE2	2:16:B:PRO:HD2	2	0.24
(1,267)	1:12:A:ARG:H	1:12:A:ARG:HB2	10	0.24
(1,267)	1:12:A:ARG:H	1:12:A:ARG:HB3	10	0.24
(1,267)	1:12:A:ARG:H	1:12:A:ARG:HB2	12	0.24
(1,267)	1:12:A:ARG:H	1:12:A:ARG:HB3	12	0.24
(1,267)	1:12:A:ARG:H	1:12:A:ARG:HB2	17	0.24
(1,267)	1:12:A:ARG:H	1:12:A:ARG:HB3	17	0.24
(1,211)	1:5:A:HIS:HA	1:5:A:HIS:HD2	14	0.24
(1,159)	2:18:B:SER:HB3	2:19:B:PHE:H	9	0.24
(1,157)	2:18:B:SER:HB3	2:19:B:PHE:H	9	0.24

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,142)	1:6:A:VAL:HG11	1:7:A:PRO:HB3	6	0.24
(1,142)	1:6:A:VAL:HG12	1:7:A:PRO:HB3	6	0.24
(1,142)	1:6:A:VAL:HG13	1:7:A:PRO:HB3	6	0.24
(1,129)	1:4:A:GLY:H	2:19:B:PHE:H	1	0.24
(1,118)	1:8:A:GLU:HG2	2:20:B:TYR:HB2	11	0.24
(1,118)	1:8:A:GLU:HG2	2:20:B:TYR:HB3	11	0.24
(1,103)	1:4:A:GLY:H	2:20:B:TYR:H	15	0.24
(1,61)	2:13:B:CYS:HB2	2:14:B:GLY:HA3	9	0.24
(1,43)	1:7:A:PRO:HG2	2:19:B:PHE:HD1	6	0.24
(1,43)	1:7:A:PRO:HG2	2:19:B:PHE:HD2	6	0.24
(1,43)	1:7:A:PRO:HG3	2:19:B:PHE:HD1	6	0.24
(1,43)	1:7:A:PRO:HG3	2:19:B:PHE:HD2	6	0.24
(1,37)	1:8:A:GLU:HG2	2:20:B:TYR:HE1	3	0.24
(1,37)	1:8:A:GLU:HG2	2:20:B:TYR:HE2	3	0.24
(1,25)	1:10:A:PHE:HE1	2:16:B:PRO:HD2	2	0.24
(1,25)	1:10:A:PHE:HE2	2:16:B:PRO:HD2	2	0.24
(1,1252)	1:28:C:PRO:HG2	1:31:C:PHE:HE1	10	0.23
(1,1252)	1:28:C:PRO:HG2	1:31:C:PHE:HE2	10	0.23
(1,1252)	1:28:C:PRO:HG3	1:31:C:PHE:HE1	10	0.23
(1,1252)	1:28:C:PRO:HG3	1:31:C:PHE:HE2	10	0.23
(1,1252)	1:28:C:PRO:HG2	1:31:C:PHE:HE1	12	0.23
(1,1252)	1:28:C:PRO:HG2	1:31:C:PHE:HE2	12	0.23
(1,1252)	1:28:C:PRO:HG3	1:31:C:PHE:HE1	12	0.23
(1,1252)	1:28:C:PRO:HG3	1:31:C:PHE:HE2	12	0.23
(1,1186)	1:26:C:HIS:HA	1:26:C:HIS:HD2	2	0.23
(1,1186)	1:26:C:HIS:HA	1:26:C:HIS:HD2	20	0.23
(1,1156)	2:34:D:CYS:HB3	2:35:D:GLY:H	14	0.23
(1,1098)	1:29:C:GLU:HG3	2:40:D:PHE:HA	9	0.23
(1,1090)	1:24:C:ALA:HB1	1:25:C:GLY:HA3	16	0.23
(1,1090)	1:24:C:ALA:HB2	1:25:C:GLY:HA3	16	0.23
(1,1090)	1:24:C:ALA:HB3	1:25:C:GLY:HA3	16	0.23
(1,1073)	2:18:B:SER:HB3	2:38:D:ILE:HD11	11	0.23
(1,1073)	2:18:B:SER:HB3	2:38:D:ILE:HD12	11	0.23
(1,1073)	2:18:B:SER:HB3	2:38:D:ILE:HD13	11	0.23
(1,1066)	1:23:C:GLY:HA3	1:24:C:ALA:HB1	9	0.23
(1,1066)	1:23:C:GLY:HA3	1:24:C:ALA:HB2	9	0.23
(1,1066)	1:23:C:GLY:HA3	1:24:C:ALA:HB3	9	0.23
(1,1043)	1:27:C:VAL:HG11	1:28:C:PRO:HG2	18	0.23
(1,1043)	1:27:C:VAL:HG11	1:28:C:PRO:HG3	18	0.23
(1,1043)	1:27:C:VAL:HG12	1:28:C:PRO:HG2	18	0.23
(1,1043)	1:27:C:VAL:HG12	1:28:C:PRO:HG3	18	0.23
(1,1043)	1:27:C:VAL:HG13	1:28:C:PRO:HG2	18	0.23

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,1043)	1:27:C:VAL:HG13	1:28:C:PRO:HG3	18	0.23
(1,1030)	1:24:C:ALA:HB1	1:25:C:GLY:HA3	16	0.23
(1,1030)	1:24:C:ALA:HB2	1:25:C:GLY:HA3	16	0.23
(1,1030)	1:24:C:ALA:HB3	1:25:C:GLY:HA3	16	0.23
(1,1003)	1:28:C:PRO:HD2	2:41:D:TYR:HD1	6	0.23
(1,1003)	1:28:C:PRO:HD2	2:41:D:TYR:HD2	6	0.23
(1,933)	1:28:C:PRO:HG2	1:31:C:PHE:HE1	10	0.23
(1,933)	1:28:C:PRO:HG2	1:31:C:PHE:HE2	10	0.23
(1,933)	1:28:C:PRO:HG3	1:31:C:PHE:HE1	10	0.23
(1,933)	1:28:C:PRO:HG3	1:31:C:PHE:HE2	10	0.23
(1,933)	1:28:C:PRO:HG2	1:31:C:PHE:HE1	12	0.23
(1,933)	1:28:C:PRO:HG2	1:31:C:PHE:HE2	12	0.23
(1,933)	1:28:C:PRO:HG3	1:31:C:PHE:HE1	12	0.23
(1,933)	1:28:C:PRO:HG3	1:31:C:PHE:HE2	12	0.23
(1,867)	1:26:C:HIS:HA	1:26:C:HIS:HD2	2	0.23
(1,867)	1:26:C:HIS:HA	1:26:C:HIS:HD2	20	0.23
(1,837)	2:34:D:CYS:HB3	2:35:D:GLY:H	14	0.23
(1,779)	1:29:C:GLU:HG3	2:40:D:PHE:HA	9	0.23
(1,771)	1:24:C:ALA:HB1	1:25:C:GLY:HA3	16	0.23
(1,771)	1:24:C:ALA:HB2	1:25:C:GLY:HA3	16	0.23
(1,771)	1:24:C:ALA:HB3	1:25:C:GLY:HA3	16	0.23
(1,754)	2:18:B:SER:HB3	2:38:D:ILE:HD11	11	0.23
(1,754)	2:18:B:SER:HB3	2:38:D:ILE:HD12	11	0.23
(1,754)	2:18:B:SER:HB3	2:38:D:ILE:HD13	11	0.23
(1,747)	1:23:C:GLY:HA3	1:24:C:ALA:HB1	9	0.23
(1,747)	1:23:C:GLY:HA3	1:24:C:ALA:HB2	9	0.23
(1,747)	1:23:C:GLY:HA3	1:24:C:ALA:HB3	9	0.23
(1,724)	1:27:C:VAL:HG11	1:28:C:PRO:HG2	18	0.23
(1,724)	1:27:C:VAL:HG11	1:28:C:PRO:HG3	18	0.23
(1,724)	1:27:C:VAL:HG12	1:28:C:PRO:HG2	18	0.23
(1,724)	1:27:C:VAL:HG12	1:28:C:PRO:HG3	18	0.23
(1,724)	1:27:C:VAL:HG13	1:28:C:PRO:HG2	18	0.23
(1,724)	1:27:C:VAL:HG13	1:28:C:PRO:HG3	18	0.23
(1,711)	1:24:C:ALA:HB1	1:25:C:GLY:HA3	16	0.23
(1,711)	1:24:C:ALA:HB2	1:25:C:GLY:HA3	16	0.23
(1,711)	1:24:C:ALA:HB3	1:25:C:GLY:HA3	16	0.23
(1,684)	1:28:C:PRO:HD2	2:41:D:TYR:HD1	6	0.23
(1,684)	1:28:C:PRO:HD2	2:41:D:TYR:HD2	6	0.23
(1,672)	2:17:B:ILE:HG13	2:40:D:PHE:HD1	9	0.23
(1,672)	2:17:B:ILE:HG13	2:40:D:PHE:HD2	9	0.23
(1,659)	1:11:A:VAL:HA	1:24:C:ALA:HB1	1	0.23
(1,659)	1:11:A:VAL:HA	1:24:C:ALA:HB2	1	0.23

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,659)	1:11:A:VAL:HA	1:24:C:ALA:HB3	1	0.23
(1,503)	1:7:A:PRO:HB3	1:10:A:PHE:HE1	15	0.23
(1,503)	1:7:A:PRO:HB3	1:10:A:PHE:HE2	15	0.23
(1,478)	2:18:B:SER:HB3	2:19:B:PHE:H	18	0.23
(1,476)	2:18:B:SER:HB3	2:19:B:PHE:H	18	0.23
(1,442)	1:8:A:GLU:HG3	2:19:B:PHE:HA	13	0.23
(1,434)	1:3:A:ALA:HB1	1:4:A:GLY:HA3	9	0.23
(1,434)	1:3:A:ALA:HB2	1:4:A:GLY:HA3	9	0.23
(1,434)	1:3:A:ALA:HB3	1:4:A:GLY:HA3	9	0.23
(1,405)	1:1:A:GLY:H	2:18:B:SER:HB2	15	0.23
(1,390)	1:6:A:VAL:HB	2:20:B:TYR:HE1	17	0.23
(1,390)	1:6:A:VAL:HB	2:20:B:TYR:HE2	17	0.23
(1,389)	1:6:A:VAL:HB	2:20:B:TYR:HD1	9	0.23
(1,389)	1:6:A:VAL:HB	2:20:B:TYR:HD2	9	0.23
(1,374)	1:3:A:ALA:HB1	1:4:A:GLY:HA3	9	0.23
(1,374)	1:3:A:ALA:HB2	1:4:A:GLY:HA3	9	0.23
(1,374)	1:3:A:ALA:HB3	1:4:A:GLY:HA3	9	0.23
(1,349)	2:13:B:CYS:HB3	2:14:B:GLY:HA3	10	0.23
(1,184)	1:7:A:PRO:HB3	1:10:A:PHE:HE1	15	0.23
(1,184)	1:7:A:PRO:HB3	1:10:A:PHE:HE2	15	0.23
(1,159)	2:18:B:SER:HB3	2:19:B:PHE:H	18	0.23
(1,157)	2:18:B:SER:HB3	2:19:B:PHE:H	18	0.23
(1,123)	1:8:A:GLU:HG3	2:19:B:PHE:HA	13	0.23
(1,115)	1:3:A:ALA:HB1	1:4:A:GLY:HA3	9	0.23
(1,115)	1:3:A:ALA:HB2	1:4:A:GLY:HA3	9	0.23
(1,115)	1:3:A:ALA:HB3	1:4:A:GLY:HA3	9	0.23
(1,86)	1:1:A:GLY:H	2:18:B:SER:HB2	15	0.23
(1,71)	1:6:A:VAL:HB	2:20:B:TYR:HE1	17	0.23
(1,71)	1:6:A:VAL:HB	2:20:B:TYR:HE2	17	0.23
(1,70)	1:6:A:VAL:HB	2:20:B:TYR:HD1	9	0.23
(1,70)	1:6:A:VAL:HB	2:20:B:TYR:HD2	9	0.23
(1,55)	1:3:A:ALA:HB1	1:4:A:GLY:HA3	9	0.23
(1,55)	1:3:A:ALA:HB2	1:4:A:GLY:HA3	9	0.23
(1,55)	1:3:A:ALA:HB3	1:4:A:GLY:HA3	9	0.23
(1,30)	2:13:B:CYS:HB3	2:14:B:GLY:HA3	10	0.23
(1,12)	1:11:A:VAL:HA	1:24:C:ALA:HB1	1	0.23
(1,12)	1:11:A:VAL:HA	1:24:C:ALA:HB2	1	0.23
(1,12)	1:11:A:VAL:HA	1:24:C:ALA:HB3	1	0.23
(1,7)	2:17:B:ILE:HG13	2:40:D:PHE:HD1	9	0.23
(1,7)	2:17:B:ILE:HG13	2:40:D:PHE:HD2	9	0.23
(1,1242)	1:33:C:ARG:H	1:33:C:ARG:HB2	3	0.22
(1,1242)	1:33:C:ARG:H	1:33:C:ARG:HB3	3	0.22

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,1166)	1:24:C:ALA:H	2:40:D:PHE:HD1	15	0.22
(1,1166)	1:24:C:ALA:H	2:40:D:PHE:HD2	15	0.22
(1,1159)	1:28:C:PRO:HB3	1:31:C:PHE:HE1	5	0.22
(1,1159)	1:28:C:PRO:HB3	1:31:C:PHE:HE2	5	0.22
(1,1104)	1:25:C:GLY:H	2:40:D:PHE:H	3	0.22
(1,1098)	1:29:C:GLU:HG3	2:40:D:PHE:HA	13	0.22
(1,1097)	1:29:C:GLU:HG3	2:41:D:TYR:HB2	12	0.22
(1,1097)	1:29:C:GLU:HG3	2:41:D:TYR:HB3	12	0.22
(1,1050)	1:28:C:PRO:HG2	1:31:C:PHE:HB2	12	0.22
(1,1050)	1:28:C:PRO:HG3	1:31:C:PHE:HB2	12	0.22
(1,1034)	2:38:D:ILE:H	2:38:D:ILE:HG13	2	0.22
(1,1027)	1:24:C:ALA:H	2:40:D:PHE:HD1	15	0.22
(1,1027)	1:24:C:ALA:H	2:40:D:PHE:HD2	15	0.22
(1,1017)	1:31:C:PHE:HE1	2:35:D:GLY:HA3	9	0.22
(1,1017)	1:31:C:PHE:HE2	2:35:D:GLY:HA3	9	0.22
(1,1016)	1:28:C:PRO:HG2	1:31:C:PHE:HB2	12	0.22
(1,1016)	1:28:C:PRO:HG3	1:31:C:PHE:HB2	12	0.22
(1,1003)	1:28:C:PRO:HD2	2:41:D:TYR:HD1	19	0.22
(1,1003)	1:28:C:PRO:HD2	2:41:D:TYR:HD2	19	0.22
(1,923)	1:33:C:ARG:H	1:33:C:ARG:HB2	3	0.22
(1,923)	1:33:C:ARG:H	1:33:C:ARG:HB3	3	0.22
(1,847)	1:24:C:ALA:H	2:40:D:PHE:HD1	15	0.22
(1,847)	1:24:C:ALA:H	2:40:D:PHE:HD2	15	0.22
(1,840)	1:28:C:PRO:HB3	1:31:C:PHE:HE1	5	0.22
(1,840)	1:28:C:PRO:HB3	1:31:C:PHE:HE2	5	0.22
(1,785)	1:25:C:GLY:H	2:40:D:PHE:H	3	0.22
(1,779)	1:29:C:GLU:HG3	2:40:D:PHE:HA	13	0.22
(1,778)	1:29:C:GLU:HG3	2:41:D:TYR:HB2	12	0.22
(1,778)	1:29:C:GLU:HG3	2:41:D:TYR:HB3	12	0.22
(1,731)	1:28:C:PRO:HG2	1:31:C:PHE:HB2	12	0.22
(1,731)	1:28:C:PRO:HG3	1:31:C:PHE:HB2	12	0.22
(1,715)	2:38:D:ILE:H	2:38:D:ILE:HG13	2	0.22
(1,708)	1:24:C:ALA:H	2:40:D:PHE:HD1	15	0.22
(1,708)	1:24:C:ALA:H	2:40:D:PHE:HD2	15	0.22
(1,698)	1:31:C:PHE:HE1	2:35:D:GLY:HA3	9	0.22
(1,698)	1:31:C:PHE:HE2	2:35:D:GLY:HA3	9	0.22
(1,697)	1:28:C:PRO:HG2	1:31:C:PHE:HB2	12	0.22
(1,697)	1:28:C:PRO:HG3	1:31:C:PHE:HB2	12	0.22
(1,684)	1:28:C:PRO:HD2	2:41:D:TYR:HD1	19	0.22
(1,684)	1:28:C:PRO:HD2	2:41:D:TYR:HD2	19	0.22
(1,657)	2:19:B:PHE:HD1	2:38:D:ILE:HD11	1	0.22
(1,657)	2:19:B:PHE:HD1	2:38:D:ILE:HD12	1	0.22

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,657)	2:19:B:PHE:HD1	2:38:D:ILE:HD13	1	0.22
(1,657)	2:19:B:PHE:HD2	2:38:D:ILE:HD11	1	0.22
(1,657)	2:19:B:PHE:HD2	2:38:D:ILE:HD12	1	0.22
(1,657)	2:19:B:PHE:HD2	2:38:D:ILE:HD13	1	0.22
(1,516)	2:15:B:THR:HB	2:16:B:PRO:HD2	19	0.22
(1,514)	1:10:A:PHE:HB2	1:11:A:VAL:HG11	19	0.22
(1,514)	1:10:A:PHE:HB2	1:11:A:VAL:HG12	19	0.22
(1,514)	1:10:A:PHE:HB2	1:11:A:VAL:HG13	19	0.22
(1,514)	1:10:A:PHE:HB2	1:11:A:VAL:HG21	19	0.22
(1,514)	1:10:A:PHE:HB2	1:11:A:VAL:HG22	19	0.22
(1,514)	1:10:A:PHE:HB2	1:11:A:VAL:HG23	19	0.22
(1,493)	1:11:A:VAL:HB	1:12:A:ARG:H	1	0.22
(1,451)	2:15:B:THR:HG21	2:16:B:PRO:HB3	1	0.22
(1,451)	2:15:B:THR:HG22	2:16:B:PRO:HB3	1	0.22
(1,451)	2:15:B:THR:HG23	2:16:B:PRO:HB3	1	0.22
(1,449)	1:3:A:ALA:HB1	1:4:A:GLY:H	20	0.22
(1,449)	1:3:A:ALA:HB2	1:4:A:GLY:H	20	0.22
(1,449)	1:3:A:ALA:HB3	1:4:A:GLY:H	20	0.22
(1,445)	1:4:A:GLY:HA2	1:6:A:VAL:H	17	0.22
(1,444)	2:19:B:PHE:HA	2:20:B:TYR:HD1	7	0.22
(1,444)	2:19:B:PHE:HA	2:20:B:TYR:HD2	7	0.22
(1,422)	1:4:A:GLY:H	2:20:B:TYR:H	7	0.22
(1,418)	1:1:A:GLY:H	2:18:B:SER:HB3	12	0.22
(1,408)	2:15:B:THR:HG21	2:16:B:PRO:HA	15	0.22
(1,408)	2:15:B:THR:HG22	2:16:B:PRO:HA	15	0.22
(1,408)	2:15:B:THR:HG23	2:16:B:PRO:HA	15	0.22
(1,406)	1:11:A:VAL:HG11	1:12:A:ARG:HA	9	0.22
(1,406)	1:11:A:VAL:HG12	1:12:A:ARG:HA	9	0.22
(1,406)	1:11:A:VAL:HG13	1:12:A:ARG:HA	9	0.22
(1,406)	1:11:A:VAL:HG21	1:12:A:ARG:HA	9	0.22
(1,406)	1:11:A:VAL:HG22	1:12:A:ARG:HA	9	0.22
(1,406)	1:11:A:VAL:HG23	1:12:A:ARG:HA	9	0.22
(1,403)	1:7:A:PRO:HG2	2:19:B:PHE:HD1	8	0.22
(1,403)	1:7:A:PRO:HG2	2:19:B:PHE:HD2	8	0.22
(1,403)	1:7:A:PRO:HG3	2:19:B:PHE:HD1	8	0.22
(1,403)	1:7:A:PRO:HG3	2:19:B:PHE:HD2	8	0.22
(1,379)	1:1:A:GLY:H	2:20:B:TYR:HE1	11	0.22
(1,379)	1:1:A:GLY:H	2:20:B:TYR:HE2	11	0.22
(1,378)	2:17:B:ILE:H	2:17:B:ILE:HG13	18	0.22
(1,361)	1:10:A:PHE:HE1	2:14:B:GLY:HA3	12	0.22
(1,361)	1:10:A:PHE:HE2	2:14:B:GLY:HA3	12	0.22
(1,197)	2:15:B:THR:HB	2:16:B:PRO:HD2	19	0.22

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,195)	1:10:A:PHE:HB2	1:11:A:VAL:HG11	19	0.22
(1,195)	1:10:A:PHE:HB2	1:11:A:VAL:HG12	19	0.22
(1,195)	1:10:A:PHE:HB2	1:11:A:VAL:HG13	19	0.22
(1,195)	1:10:A:PHE:HB2	1:11:A:VAL:HG21	19	0.22
(1,195)	1:10:A:PHE:HB2	1:11:A:VAL:HG22	19	0.22
(1,195)	1:10:A:PHE:HB2	1:11:A:VAL:HG23	19	0.22
(1,174)	1:11:A:VAL:HB	1:12:A:ARG:H	1	0.22
(1,132)	2:15:B:THR:HG21	2:16:B:PRO:HB3	1	0.22
(1,132)	2:15:B:THR:HG22	2:16:B:PRO:HB3	1	0.22
(1,132)	2:15:B:THR:HG23	2:16:B:PRO:HB3	1	0.22
(1,130)	1:3:A:ALA:HB1	1:4:A:GLY:H	20	0.22
(1,130)	1:3:A:ALA:HB2	1:4:A:GLY:H	20	0.22
(1,130)	1:3:A:ALA:HB3	1:4:A:GLY:H	20	0.22
(1,126)	1:4:A:GLY:HA2	1:6:A:VAL:H	17	0.22
(1,125)	2:19:B:PHE:HA	2:20:B:TYR:HD1	7	0.22
(1,125)	2:19:B:PHE:HA	2:20:B:TYR:HD2	7	0.22
(1,103)	1:4:A:GLY:H	2:20:B:TYR:H	7	0.22
(1,99)	1:1:A:GLY:H	2:18:B:SER:HB3	12	0.22
(1,89)	2:15:B:THR:HG21	2:16:B:PRO:HA	15	0.22
(1,89)	2:15:B:THR:HG22	2:16:B:PRO:HA	15	0.22
(1,89)	2:15:B:THR:HG23	2:16:B:PRO:HA	15	0.22
(1,87)	1:11:A:VAL:HG11	1:12:A:ARG:HA	9	0.22
(1,87)	1:11:A:VAL:HG12	1:12:A:ARG:HA	9	0.22
(1,87)	1:11:A:VAL:HG13	1:12:A:ARG:HA	9	0.22
(1,87)	1:11:A:VAL:HG21	1:12:A:ARG:HA	9	0.22
(1,87)	1:11:A:VAL:HG22	1:12:A:ARG:HA	9	0.22
(1,87)	1:11:A:VAL:HG23	1:12:A:ARG:HA	9	0.22
(1,84)	1:7:A:PRO:HG2	2:19:B:PHE:HD1	8	0.22
(1,84)	1:7:A:PRO:HG2	2:19:B:PHE:HD2	8	0.22
(1,84)	1:7:A:PRO:HG3	2:19:B:PHE:HD1	8	0.22
(1,84)	1:7:A:PRO:HG3	2:19:B:PHE:HD2	8	0.22
(1,60)	1:1:A:GLY:H	2:20:B:TYR:HE1	11	0.22
(1,60)	1:1:A:GLY:H	2:20:B:TYR:HE2	11	0.22
(1,59)	2:17:B:ILE:H	2:17:B:ILE:HG13	18	0.22
(1,42)	1:10:A:PHE:HE1	2:14:B:GLY:HA3	12	0.22
(1,42)	1:10:A:PHE:HE2	2:14:B:GLY:HA3	12	0.22
(1,10)	2:19:B:PHE:HD1	2:38:D:ILE:HD11	1	0.22
(1,10)	2:19:B:PHE:HD1	2:38:D:ILE:HD12	1	0.22
(1,10)	2:19:B:PHE:HD1	2:38:D:ILE:HD13	1	0.22
(1,10)	2:19:B:PHE:HD2	2:38:D:ILE:HD11	1	0.22
(1,10)	2:19:B:PHE:HD2	2:38:D:ILE:HD12	1	0.22
(1,10)	2:19:B:PHE:HD2	2:38:D:ILE:HD13	1	0.22

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,1285)	2:38:D:ILE:H	2:38:D:ILE:HD11	1	0.21
(1,1285)	2:38:D:ILE:H	2:38:D:ILE:HD12	1	0.21
(1,1285)	2:38:D:ILE:H	2:38:D:ILE:HD13	1	0.21
(1,1243)	1:33:C:ARG:H	1:33:C:ARG:HE	18	0.21
(1,1242)	1:33:C:ARG:H	1:33:C:ARG:HB2	15	0.21
(1,1242)	1:33:C:ARG:H	1:33:C:ARG:HB3	15	0.21
(1,1231)	1:31:C:PHE:HE1	2:36:D:THR:HA	3	0.21
(1,1231)	1:31:C:PHE:HE2	2:36:D:THR:HA	3	0.21
(1,1224)	1:31:C:PHE:H	1:31:C:PHE:HD1	3	0.21
(1,1224)	1:31:C:PHE:H	1:31:C:PHE:HD2	3	0.21
(1,1156)	2:34:D:CYS:HB3	2:35:D:GLY:H	19	0.21
(1,1146)	1:28:C:PRO:HA	2:41:D:TYR:H	3	0.21
(1,1117)	1:27:C:VAL:HG11	1:28:C:PRO:HB3	11	0.21
(1,1117)	1:27:C:VAL:HG12	1:28:C:PRO:HB3	11	0.21
(1,1117)	1:27:C:VAL:HG13	1:28:C:PRO:HB3	11	0.21
(1,1097)	1:29:C:GLU:HG3	2:41:D:TYR:HB2	13	0.21
(1,1097)	1:29:C:GLU:HG3	2:41:D:TYR:HB3	13	0.21
(1,1090)	1:24:C:ALA:HB1	1:25:C:GLY:HA3	1	0.21
(1,1090)	1:24:C:ALA:HB2	1:25:C:GLY:HA3	1	0.21
(1,1090)	1:24:C:ALA:HB3	1:25:C:GLY:HA3	1	0.21
(1,1030)	1:24:C:ALA:HB1	1:25:C:GLY:HA3	1	0.21
(1,1030)	1:24:C:ALA:HB2	1:25:C:GLY:HA3	1	0.21
(1,1030)	1:24:C:ALA:HB3	1:25:C:GLY:HA3	1	0.21
(1,1005)	2:34:D:CYS:HB3	2:35:D:GLY:HA3	12	0.21
(1,1003)	1:28:C:PRO:HD2	2:41:D:TYR:HD1	2	0.21
(1,1003)	1:28:C:PRO:HD2	2:41:D:TYR:HD2	2	0.21
(1,966)	2:38:D:ILE:H	2:38:D:ILE:HD11	1	0.21
(1,966)	2:38:D:ILE:H	2:38:D:ILE:HD12	1	0.21
(1,966)	2:38:D:ILE:H	2:38:D:ILE:HD13	1	0.21
(1,924)	1:33:C:ARG:H	1:33:C:ARG:HE	18	0.21
(1,923)	1:33:C:ARG:H	1:33:C:ARG:HB2	15	0.21
(1,923)	1:33:C:ARG:H	1:33:C:ARG:HB3	15	0.21
(1,912)	1:31:C:PHE:HE1	2:36:D:THR:HA	3	0.21
(1,912)	1:31:C:PHE:HE2	2:36:D:THR:HA	3	0.21
(1,905)	1:31:C:PHE:H	1:31:C:PHE:HD1	3	0.21
(1,905)	1:31:C:PHE:H	1:31:C:PHE:HD2	3	0.21
(1,837)	2:34:D:CYS:HB3	2:35:D:GLY:H	19	0.21
(1,827)	1:28:C:PRO:HA	2:41:D:TYR:H	3	0.21
(1,798)	1:27:C:VAL:HG11	1:28:C:PRO:HB3	11	0.21
(1,798)	1:27:C:VAL:HG12	1:28:C:PRO:HB3	11	0.21
(1,798)	1:27:C:VAL:HG13	1:28:C:PRO:HB3	11	0.21
(1,778)	1:29:C:GLU:HG3	2:41:D:TYR:HB2	13	0.21

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,778)	1:29:C:GLU:HG3	2:41:D:TYR:HB3	13	0.21
(1,771)	1:24:C:ALA:HB1	1:25:C:GLY:HA3	1	0.21
(1,771)	1:24:C:ALA:HB2	1:25:C:GLY:HA3	1	0.21
(1,771)	1:24:C:ALA:HB3	1:25:C:GLY:HA3	1	0.21
(1,711)	1:24:C:ALA:HB1	1:25:C:GLY:HA3	1	0.21
(1,711)	1:24:C:ALA:HB2	1:25:C:GLY:HA3	1	0.21
(1,711)	1:24:C:ALA:HB3	1:25:C:GLY:HA3	1	0.21
(1,686)	2:34:D:CYS:HB3	2:35:D:GLY:HA3	12	0.21
(1,684)	1:28:C:PRO:HD2	2:41:D:TYR:HD1	2	0.21
(1,684)	1:28:C:PRO:HD2	2:41:D:TYR:HD2	2	0.21
(1,674)	2:17:B:ILE:HD11	2:40:D:PHE:HE1	20	0.21
(1,674)	2:17:B:ILE:HD11	2:40:D:PHE:HE2	20	0.21
(1,674)	2:17:B:ILE:HD12	2:40:D:PHE:HE1	20	0.21
(1,674)	2:17:B:ILE:HD12	2:40:D:PHE:HE2	20	0.21
(1,674)	2:17:B:ILE:HD13	2:40:D:PHE:HE1	20	0.21
(1,674)	2:17:B:ILE:HD13	2:40:D:PHE:HE2	20	0.21
(1,586)	1:12:A:ARG:H	1:12:A:ARG:HB2	15	0.21
(1,586)	1:12:A:ARG:H	1:12:A:ARG:HB3	15	0.21
(1,562)	1:9:A:TYR:HA	1:9:A:TYR:HE1	12	0.21
(1,562)	1:9:A:TYR:HA	1:9:A:TYR:HE2	12	0.21
(1,514)	1:10:A:PHE:HB2	1:11:A:VAL:HG11	14	0.21
(1,514)	1:10:A:PHE:HB2	1:11:A:VAL:HG12	14	0.21
(1,514)	1:10:A:PHE:HB2	1:11:A:VAL:HG13	14	0.21
(1,514)	1:10:A:PHE:HB2	1:11:A:VAL:HG21	14	0.21
(1,514)	1:10:A:PHE:HB2	1:11:A:VAL:HG22	14	0.21
(1,514)	1:10:A:PHE:HB2	1:11:A:VAL:HG23	14	0.21
(1,445)	1:4:A:GLY:HA2	1:6:A:VAL:H	7	0.21
(1,411)	1:5:A:HIS:HB3	1:6:A:VAL:HG21	5	0.21
(1,411)	1:5:A:HIS:HB3	1:6:A:VAL:HG22	5	0.21
(1,411)	1:5:A:HIS:HB3	1:6:A:VAL:HG23	5	0.21
(1,410)	1:2:A:GLY:HA3	1:3:A:ALA:HB1	20	0.21
(1,410)	1:2:A:GLY:HA3	1:3:A:ALA:HB2	20	0.21
(1,410)	1:2:A:GLY:HA3	1:3:A:ALA:HB3	20	0.21
(1,405)	1:1:A:GLY:H	2:18:B:SER:HB2	5	0.21
(1,370)	1:6:A:VAL:HG11	1:7:A:PRO:HD3	10	0.21
(1,370)	1:6:A:VAL:HG12	1:7:A:PRO:HD3	10	0.21
(1,370)	1:6:A:VAL:HG13	1:7:A:PRO:HD3	10	0.21
(1,362)	1:7:A:PRO:HG2	2:19:B:PHE:HD1	1	0.21
(1,362)	1:7:A:PRO:HG2	2:19:B:PHE:HD2	1	0.21
(1,362)	1:7:A:PRO:HG3	2:19:B:PHE:HD1	1	0.21
(1,362)	1:7:A:PRO:HG3	2:19:B:PHE:HD2	1	0.21
(1,353)	1:6:A:VAL:HB	2:20:B:TYR:HB2	20	0.21

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,353)	1:6:A:VAL:HB	2:20:B:TYR:HB3	20	0.21
(1,267)	1:12:A:ARG:H	1:12:A:ARG:HB2	15	0.21
(1,267)	1:12:A:ARG:H	1:12:A:ARG:HB3	15	0.21
(1,243)	1:9:A:TYR:HA	1:9:A:TYR:HE1	12	0.21
(1,243)	1:9:A:TYR:HA	1:9:A:TYR:HE2	12	0.21
(1,195)	1:10:A:PHE:HB2	1:11:A:VAL:HG11	14	0.21
(1,195)	1:10:A:PHE:HB2	1:11:A:VAL:HG12	14	0.21
(1,195)	1:10:A:PHE:HB2	1:11:A:VAL:HG13	14	0.21
(1,195)	1:10:A:PHE:HB2	1:11:A:VAL:HG21	14	0.21
(1,195)	1:10:A:PHE:HB2	1:11:A:VAL:HG22	14	0.21
(1,195)	1:10:A:PHE:HB2	1:11:A:VAL:HG23	14	0.21
(1,126)	1:4:A:GLY:HA2	1:6:A:VAL:H	7	0.21
(1,92)	1:5:A:HIS:HB3	1:6:A:VAL:HG21	5	0.21
(1,92)	1:5:A:HIS:HB3	1:6:A:VAL:HG22	5	0.21
(1,92)	1:5:A:HIS:HB3	1:6:A:VAL:HG23	5	0.21
(1,91)	1:2:A:GLY:HA3	1:3:A:ALA:HB1	20	0.21
(1,91)	1:2:A:GLY:HA3	1:3:A:ALA:HB2	20	0.21
(1,91)	1:2:A:GLY:HA3	1:3:A:ALA:HB3	20	0.21
(1,86)	1:1:A:GLY:H	2:18:B:SER:HB2	5	0.21
(1,51)	1:6:A:VAL:HG11	1:7:A:PRO:HD3	10	0.21
(1,51)	1:6:A:VAL:HG12	1:7:A:PRO:HD3	10	0.21
(1,51)	1:6:A:VAL:HG13	1:7:A:PRO:HD3	10	0.21
(1,43)	1:7:A:PRO:HG2	2:19:B:PHE:HD1	1	0.21
(1,43)	1:7:A:PRO:HG2	2:19:B:PHE:HD2	1	0.21
(1,43)	1:7:A:PRO:HG3	2:19:B:PHE:HD1	1	0.21
(1,43)	1:7:A:PRO:HG3	2:19:B:PHE:HD2	1	0.21
(1,34)	1:6:A:VAL:HB	2:20:B:TYR:HB2	20	0.21
(1,34)	1:6:A:VAL:HB	2:20:B:TYR:HB3	20	0.21
(1,9)	2:17:B:ILE:HD11	2:40:D:PHE:HE1	20	0.21
(1,9)	2:17:B:ILE:HD11	2:40:D:PHE:HE2	20	0.21
(1,9)	2:17:B:ILE:HD12	2:40:D:PHE:HE1	20	0.21
(1,9)	2:17:B:ILE:HD12	2:40:D:PHE:HE2	20	0.21
(1,9)	2:17:B:ILE:HD13	2:40:D:PHE:HE1	20	0.21
(1,9)	2:17:B:ILE:HD13	2:40:D:PHE:HE2	20	0.21
(1,1221)	1:31:C:PHE:HA	1:32:C:VAL:H	12	0.2
(1,1218)	1:30:C:TYR:HA	1:30:C:TYR:HE1	18	0.2
(1,1218)	1:30:C:TYR:HA	1:30:C:TYR:HE2	18	0.2
(1,1135)	1:22:C:GLY:H	1:29:C:GLU:HB3	14	0.2
(1,1135)	1:22:C:GLY:H	1:29:C:GLU:HB2	14	0.2
(1,1130)	1:27:C:VAL:H	2:40:D:PHE:HB2	3	0.2
(1,1130)	1:27:C:VAL:H	2:40:D:PHE:HB2	19	0.2
(1,1117)	1:27:C:VAL:HG11	1:28:C:PRO:HB3	13	0.2

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,1117)	1:27:C:VAL:HG12	1:28:C:PRO:HB3	13	0.2
(1,1117)	1:27:C:VAL:HG13	1:28:C:PRO:HB3	13	0.2
(1,1107)	2:36:D:THR:HG21	2:37:D:PRO:HB3	2	0.2
(1,1107)	2:36:D:THR:HG22	2:37:D:PRO:HB3	2	0.2
(1,1107)	2:36:D:THR:HG23	2:37:D:PRO:HB3	2	0.2
(1,1093)	1:29:C:GLU:HG2	2:41:D:TYR:HB2	7	0.2
(1,1093)	1:29:C:GLU:HG2	2:41:D:TYR:HB3	7	0.2
(1,1078)	1:25:C:GLY:H	2:41:D:TYR:H	6	0.2
(1,1059)	1:28:C:PRO:HG2	2:40:D:PHE:HD1	2	0.2
(1,1059)	1:28:C:PRO:HG2	2:40:D:PHE:HD2	2	0.2
(1,1059)	1:28:C:PRO:HG3	2:40:D:PHE:HD1	2	0.2
(1,1059)	1:28:C:PRO:HG3	2:40:D:PHE:HD2	2	0.2
(1,1053)	2:39:D:SER:HB2	2:40:D:PHE:H	10	0.2
(1,1038)	1:31:C:PHE:HD1	2:37:D:PRO:HD2	12	0.2
(1,1038)	1:31:C:PHE:HD2	2:37:D:PRO:HD2	12	0.2
(1,1026)	1:27:C:VAL:HG11	1:28:C:PRO:HD3	2	0.2
(1,1026)	1:27:C:VAL:HG12	1:28:C:PRO:HD3	2	0.2
(1,1026)	1:27:C:VAL:HG13	1:28:C:PRO:HD3	2	0.2
(1,902)	1:31:C:PHE:HA	1:32:C:VAL:H	12	0.2
(1,899)	1:30:C:TYR:HA	1:30:C:TYR:HE1	18	0.2
(1,899)	1:30:C:TYR:HA	1:30:C:TYR:HE2	18	0.2
(1,816)	1:22:C:GLY:H	1:29:C:GLU:HB3	14	0.2
(1,816)	1:22:C:GLY:H	1:29:C:GLU:HB2	14	0.2
(1,811)	1:27:C:VAL:H	2:40:D:PHE:HB2	3	0.2
(1,811)	1:27:C:VAL:H	2:40:D:PHE:HB2	19	0.2
(1,798)	1:27:C:VAL:HG11	1:28:C:PRO:HB3	13	0.2
(1,798)	1:27:C:VAL:HG12	1:28:C:PRO:HB3	13	0.2
(1,798)	1:27:C:VAL:HG13	1:28:C:PRO:HB3	13	0.2
(1,788)	2:36:D:THR:HG21	2:37:D:PRO:HB3	2	0.2
(1,788)	2:36:D:THR:HG22	2:37:D:PRO:HB3	2	0.2
(1,788)	2:36:D:THR:HG23	2:37:D:PRO:HB3	2	0.2
(1,774)	1:29:C:GLU:HG2	2:41:D:TYR:HB2	7	0.2
(1,774)	1:29:C:GLU:HG2	2:41:D:TYR:HB3	7	0.2
(1,759)	1:25:C:GLY:H	2:41:D:TYR:H	6	0.2
(1,740)	1:28:C:PRO:HG2	2:40:D:PHE:HD1	2	0.2
(1,740)	1:28:C:PRO:HG2	2:40:D:PHE:HD2	2	0.2
(1,740)	1:28:C:PRO:HG3	2:40:D:PHE:HD1	2	0.2
(1,740)	1:28:C:PRO:HG3	2:40:D:PHE:HD2	2	0.2
(1,734)	2:39:D:SER:HB2	2:40:D:PHE:H	10	0.2
(1,719)	1:31:C:PHE:HD1	2:37:D:PRO:HD2	12	0.2
(1,719)	1:31:C:PHE:HD2	2:37:D:PRO:HD2	12	0.2
(1,707)	1:27:C:VAL:HG11	1:28:C:PRO:HD3	2	0.2

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,707)	1:27:C:VAL:HG12	1:28:C:PRO:HD3	2	0.2
(1,707)	1:27:C:VAL:HG13	1:28:C:PRO:HD3	2	0.2
(1,609)	2:15:B:THR:H	2:15:B:THR:HB	15	0.2
(1,586)	1:12:A:ARG:H	1:12:A:ARG:HB2	19	0.2
(1,586)	1:12:A:ARG:H	1:12:A:ARG:HB3	19	0.2
(1,481)	1:1:A:GLY:H	2:19:B:PHE:H	5	0.2
(1,463)	1:5:A:HIS:HB2	1:6:A:VAL:HG11	3	0.2
(1,463)	1:5:A:HIS:HB2	1:6:A:VAL:HG12	3	0.2
(1,463)	1:5:A:HIS:HB2	1:6:A:VAL:HG13	3	0.2
(1,443)	1:8:A:GLU:HG3	2:20:B:TYR:HD1	2	0.2
(1,443)	1:8:A:GLU:HG3	2:20:B:TYR:HD2	2	0.2
(1,442)	1:8:A:GLU:HG3	2:19:B:PHE:HA	18	0.2
(1,406)	1:11:A:VAL:HG11	1:12:A:ARG:HA	13	0.2
(1,406)	1:11:A:VAL:HG12	1:12:A:ARG:HA	13	0.2
(1,406)	1:11:A:VAL:HG13	1:12:A:ARG:HA	13	0.2
(1,406)	1:11:A:VAL:HG21	1:12:A:ARG:HA	13	0.2
(1,406)	1:11:A:VAL:HG22	1:12:A:ARG:HA	13	0.2
(1,406)	1:11:A:VAL:HG23	1:12:A:ARG:HA	13	0.2
(1,393)	1:7:A:PRO:HG2	1:10:A:PHE:HD1	3	0.2
(1,393)	1:7:A:PRO:HG2	1:10:A:PHE:HD2	3	0.2
(1,393)	1:7:A:PRO:HG3	1:10:A:PHE:HD1	3	0.2
(1,393)	1:7:A:PRO:HG3	1:10:A:PHE:HD2	3	0.2
(1,382)	1:10:A:PHE:HD1	2:16:B:PRO:HD2	19	0.2
(1,382)	1:10:A:PHE:HD2	2:16:B:PRO:HD2	19	0.2
(1,352)	2:17:B:ILE:HG13	2:39:D:SER:HA	10	0.2
(1,342)	1:1:A:GLY:HA3	2:18:B:SER:HB2	5	0.2
(1,339)	1:1:A:GLY:HA3	2:18:B:SER:HB2	5	0.2
(1,290)	2:15:B:THR:H	2:15:B:THR:HB	15	0.2
(1,267)	1:12:A:ARG:H	1:12:A:ARG:HB2	19	0.2
(1,267)	1:12:A:ARG:H	1:12:A:ARG:HB3	19	0.2
(1,162)	1:1:A:GLY:H	2:19:B:PHE:H	5	0.2
(1,144)	1:5:A:HIS:HB2	1:6:A:VAL:HG11	3	0.2
(1,144)	1:5:A:HIS:HB2	1:6:A:VAL:HG12	3	0.2
(1,144)	1:5:A:HIS:HB2	1:6:A:VAL:HG13	3	0.2
(1,124)	1:8:A:GLU:HG3	2:20:B:TYR:HD1	2	0.2
(1,124)	1:8:A:GLU:HG3	2:20:B:TYR:HD2	2	0.2
(1,123)	1:8:A:GLU:HG3	2:19:B:PHE:HA	18	0.2
(1,87)	1:11:A:VAL:HG11	1:12:A:ARG:HA	13	0.2
(1,87)	1:11:A:VAL:HG12	1:12:A:ARG:HA	13	0.2
(1,87)	1:11:A:VAL:HG13	1:12:A:ARG:HA	13	0.2
(1,87)	1:11:A:VAL:HG21	1:12:A:ARG:HA	13	0.2
(1,87)	1:11:A:VAL:HG22	1:12:A:ARG:HA	13	0.2

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,87)	1:11:A:VAL:HG23	1:12:A:ARG:HA	13	0.2
(1,74)	1:7:A:PRO:HG2	1:10:A:PHE:HD1	3	0.2
(1,74)	1:7:A:PRO:HG2	1:10:A:PHE:HD2	3	0.2
(1,74)	1:7:A:PRO:HG3	1:10:A:PHE:HD1	3	0.2
(1,74)	1:7:A:PRO:HG3	1:10:A:PHE:HD2	3	0.2
(1,63)	1:10:A:PHE:HD1	2:16:B:PRO:HD2	19	0.2
(1,63)	1:10:A:PHE:HD2	2:16:B:PRO:HD2	19	0.2
(1,33)	2:17:B:ILE:HG13	2:39:D:SER:HA	10	0.2
(1,23)	1:1:A:GLY:HA3	2:18:B:SER:HB2	5	0.2
(1,20)	1:1:A:GLY:HA3	2:18:B:SER:HB2	5	0.2
(1,1304)	2:40:D:PHE:HD1	2:41:D:TYR:H	14	0.19
(1,1304)	2:40:D:PHE:HD2	2:41:D:TYR:H	14	0.19
(1,1249)	1:33:C:ARG:HB2	1:33:C:ARG:HE	7	0.19
(1,1221)	1:31:C:PHE:HA	1:32:C:VAL:H	4	0.19
(1,1218)	1:30:C:TYR:HA	1:30:C:TYR:HE1	9	0.19
(1,1218)	1:30:C:TYR:HA	1:30:C:TYR:HE2	9	0.19
(1,1218)	1:30:C:TYR:HA	1:30:C:TYR:HE1	17	0.19
(1,1218)	1:30:C:TYR:HA	1:30:C:TYR:HE2	17	0.19
(1,1143)	1:27:C:VAL:H	2:41:D:TYR:HD1	1	0.19
(1,1143)	1:27:C:VAL:H	2:41:D:TYR:HD2	1	0.19
(1,1110)	1:26:C:HIS:HA	1:27:C:VAL:HG21	1	0.19
(1,1110)	1:26:C:HIS:HA	1:27:C:VAL:HG22	1	0.19
(1,1110)	1:26:C:HIS:HA	1:27:C:VAL:HG23	1	0.19
(1,1089)	1:29:C:GLU:HB2	2:41:D:TYR:HB2	16	0.19
(1,1089)	1:29:C:GLU:HB2	2:41:D:TYR:HB3	16	0.19
(1,1074)	1:22:C:GLY:H	2:39:D:SER:HB3	14	0.19
(1,1067)	1:26:C:HIS:HB3	1:27:C:VAL:HG21	7	0.19
(1,1067)	1:26:C:HIS:HB3	1:27:C:VAL:HG22	7	0.19
(1,1067)	1:26:C:HIS:HB3	1:27:C:VAL:HG23	7	0.19
(1,1066)	1:23:C:GLY:HA3	1:24:C:ALA:HB1	8	0.19
(1,1066)	1:23:C:GLY:HA3	1:24:C:ALA:HB2	8	0.19
(1,1066)	1:23:C:GLY:HA3	1:24:C:ALA:HB3	8	0.19
(1,1049)	1:28:C:PRO:HG2	1:31:C:PHE:HD1	16	0.19
(1,1049)	1:28:C:PRO:HG2	1:31:C:PHE:HD2	16	0.19
(1,1049)	1:28:C:PRO:HG3	1:31:C:PHE:HD1	16	0.19
(1,1049)	1:28:C:PRO:HG3	1:31:C:PHE:HD2	16	0.19
(1,1009)	1:27:C:VAL:HB	2:41:D:TYR:HB2	12	0.19
(1,1009)	1:27:C:VAL:HB	2:41:D:TYR:HB3	12	0.19
(1,985)	2:40:D:PHE:HD1	2:41:D:TYR:H	14	0.19
(1,985)	2:40:D:PHE:HD2	2:41:D:TYR:H	14	0.19
(1,930)	1:33:C:ARG:HB2	1:33:C:ARG:HE	7	0.19
(1,902)	1:31:C:PHE:HA	1:32:C:VAL:H	4	0.19

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,899)	1:30:C:TYR:HA	1:30:C:TYR:HE1	9	0.19
(1,899)	1:30:C:TYR:HA	1:30:C:TYR:HE2	9	0.19
(1,899)	1:30:C:TYR:HA	1:30:C:TYR:HE1	17	0.19
(1,899)	1:30:C:TYR:HA	1:30:C:TYR:HE2	17	0.19
(1,824)	1:27:C:VAL:H	2:41:D:TYR:HD1	1	0.19
(1,824)	1:27:C:VAL:H	2:41:D:TYR:HD2	1	0.19
(1,791)	1:26:C:HIS:HA	1:27:C:VAL:HG21	1	0.19
(1,791)	1:26:C:HIS:HA	1:27:C:VAL:HG22	1	0.19
(1,791)	1:26:C:HIS:HA	1:27:C:VAL:HG23	1	0.19
(1,770)	1:29:C:GLU:HB2	2:41:D:TYR:HB2	16	0.19
(1,770)	1:29:C:GLU:HB2	2:41:D:TYR:HB3	16	0.19
(1,755)	1:22:C:GLY:H	2:39:D:SER:HB3	14	0.19
(1,748)	1:26:C:HIS:HB3	1:27:C:VAL:HG21	7	0.19
(1,748)	1:26:C:HIS:HB3	1:27:C:VAL:HG22	7	0.19
(1,748)	1:26:C:HIS:HB3	1:27:C:VAL:HG23	7	0.19
(1,747)	1:23:C:GLY:HA3	1:24:C:ALA:HB1	8	0.19
(1,747)	1:23:C:GLY:HA3	1:24:C:ALA:HB2	8	0.19
(1,747)	1:23:C:GLY:HA3	1:24:C:ALA:HB3	8	0.19
(1,730)	1:28:C:PRO:HG2	1:31:C:PHE:HD1	16	0.19
(1,730)	1:28:C:PRO:HG2	1:31:C:PHE:HD2	16	0.19
(1,730)	1:28:C:PRO:HG3	1:31:C:PHE:HD1	16	0.19
(1,730)	1:28:C:PRO:HG3	1:31:C:PHE:HD2	16	0.19
(1,690)	1:27:C:VAL:HB	2:41:D:TYR:HB2	12	0.19
(1,690)	1:27:C:VAL:HB	2:41:D:TYR:HB3	12	0.19
(1,562)	1:9:A:TYR:HA	1:9:A:TYR:HE1	7	0.19
(1,562)	1:9:A:TYR:HA	1:9:A:TYR:HE2	7	0.19
(1,481)	1:1:A:GLY:H	2:19:B:PHE:H	17	0.19
(1,445)	1:4:A:GLY:HA2	1:6:A:VAL:H	4	0.19
(1,424)	1:4:A:GLY:H	2:20:B:TYR:H	14	0.19
(1,409)	2:15:B:THR:HG21	2:16:B:PRO:HD2	12	0.19
(1,409)	2:15:B:THR:HG22	2:16:B:PRO:HD2	12	0.19
(1,409)	2:15:B:THR:HG23	2:16:B:PRO:HD2	12	0.19
(1,387)	1:6:A:VAL:HG11	1:7:A:PRO:HG2	20	0.19
(1,387)	1:6:A:VAL:HG11	1:7:A:PRO:HG3	20	0.19
(1,387)	1:6:A:VAL:HG12	1:7:A:PRO:HG2	20	0.19
(1,387)	1:6:A:VAL:HG12	1:7:A:PRO:HG3	20	0.19
(1,387)	1:6:A:VAL:HG13	1:7:A:PRO:HG2	20	0.19
(1,387)	1:6:A:VAL:HG13	1:7:A:PRO:HG3	20	0.19
(1,243)	1:9:A:TYR:HA	1:9:A:TYR:HE1	7	0.19
(1,243)	1:9:A:TYR:HA	1:9:A:TYR:HE2	7	0.19
(1,162)	1:1:A:GLY:H	2:19:B:PHE:H	17	0.19
(1,126)	1:4:A:GLY:HA2	1:6:A:VAL:H	4	0.19

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,105)	1:4:A:GLY:H	2:20:B:TYR:H	14	0.19
(1,90)	2:15:B:THR:HG21	2:16:B:PRO:HD2	12	0.19
(1,90)	2:15:B:THR:HG22	2:16:B:PRO:HD2	12	0.19
(1,90)	2:15:B:THR:HG23	2:16:B:PRO:HD2	12	0.19
(1,68)	1:6:A:VAL:HG11	1:7:A:PRO:HG2	20	0.19
(1,68)	1:6:A:VAL:HG11	1:7:A:PRO:HG3	20	0.19
(1,68)	1:6:A:VAL:HG12	1:7:A:PRO:HG2	20	0.19
(1,68)	1:6:A:VAL:HG12	1:7:A:PRO:HG3	20	0.19
(1,68)	1:6:A:VAL:HG13	1:7:A:PRO:HG2	20	0.19
(1,68)	1:6:A:VAL:HG13	1:7:A:PRO:HG3	20	0.19
(1,1231)	1:31:C:PHE:HE1	2:36:D:THR:HA	17	0.18
(1,1231)	1:31:C:PHE:HE2	2:36:D:THR:HA	17	0.18
(1,1218)	1:30:C:TYR:HA	1:30:C:TYR:HE1	3	0.18
(1,1218)	1:30:C:TYR:HA	1:30:C:TYR:HE2	3	0.18
(1,1186)	1:26:C:HIS:HA	1:26:C:HIS:HD2	16	0.18
(1,1147)	1:28:C:PRO:HB2	1:31:C:PHE:HE1	14	0.18
(1,1147)	1:28:C:PRO:HB2	1:31:C:PHE:HE2	14	0.18
(1,1145)	1:25:C:GLY:H	2:40:D:PHE:HE1	6	0.18
(1,1145)	1:25:C:GLY:H	2:40:D:PHE:HE2	6	0.18
(1,1138)	1:22:C:GLY:H	1:23:C:GLY:H	19	0.18
(1,1110)	1:26:C:HIS:HA	1:27:C:VAL:HG21	15	0.18
(1,1110)	1:26:C:HIS:HA	1:27:C:VAL:HG22	15	0.18
(1,1110)	1:26:C:HIS:HA	1:27:C:VAL:HG23	15	0.18
(1,1109)	1:31:C:PHE:HD1	2:36:D:THR:HG21	14	0.18
(1,1109)	1:31:C:PHE:HD1	2:36:D:THR:HG22	14	0.18
(1,1109)	1:31:C:PHE:HD1	2:36:D:THR:HG23	14	0.18
(1,1109)	1:31:C:PHE:HD2	2:36:D:THR:HG21	14	0.18
(1,1109)	1:31:C:PHE:HD2	2:36:D:THR:HG22	14	0.18
(1,1109)	1:31:C:PHE:HD2	2:36:D:THR:HG23	14	0.18
(1,1101)	1:25:C:GLY:HA2	1:27:C:VAL:H	15	0.18
(1,1100)	2:40:D:PHE:HA	2:41:D:TYR:HD1	13	0.18
(1,1100)	2:40:D:PHE:HA	2:41:D:TYR:HD2	13	0.18
(1,1097)	1:29:C:GLU:HG3	2:41:D:TYR:HB2	17	0.18
(1,1097)	1:29:C:GLU:HG3	2:41:D:TYR:HB3	17	0.18
(1,1067)	1:26:C:HIS:HB3	1:27:C:VAL:HG21	9	0.18
(1,1067)	1:26:C:HIS:HB3	1:27:C:VAL:HG22	9	0.18
(1,1067)	1:26:C:HIS:HB3	1:27:C:VAL:HG23	9	0.18
(1,1062)	1:32:C:VAL:HG11	1:33:C:ARG:HA	5	0.18
(1,1062)	1:32:C:VAL:HG12	1:33:C:ARG:HA	5	0.18
(1,1062)	1:32:C:VAL:HG13	1:33:C:ARG:HA	5	0.18
(1,1062)	1:32:C:VAL:HG21	1:33:C:ARG:HA	5	0.18
(1,1062)	1:32:C:VAL:HG22	1:33:C:ARG:HA	5	0.18

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,1062)	1:32:C:VAL:HG23	1:33:C:ARG:HA	5	0.18
(1,1055)	1:28:C:PRO:HG2	1:31:C:PHE:HE1	1	0.18
(1,1055)	1:28:C:PRO:HG2	1:31:C:PHE:HE2	1	0.18
(1,1055)	1:28:C:PRO:HG3	1:31:C:PHE:HE1	1	0.18
(1,1055)	1:28:C:PRO:HG3	1:31:C:PHE:HE2	1	0.18
(1,1043)	1:27:C:VAL:HG11	1:28:C:PRO:HG2	13	0.18
(1,1043)	1:27:C:VAL:HG11	1:28:C:PRO:HG3	13	0.18
(1,1043)	1:27:C:VAL:HG12	1:28:C:PRO:HG2	13	0.18
(1,1043)	1:27:C:VAL:HG12	1:28:C:PRO:HG3	13	0.18
(1,1043)	1:27:C:VAL:HG13	1:28:C:PRO:HG2	13	0.18
(1,1043)	1:27:C:VAL:HG13	1:28:C:PRO:HG3	13	0.18
(1,912)	1:31:C:PHE:HE1	2:36:D:THR:HA	17	0.18
(1,912)	1:31:C:PHE:HE2	2:36:D:THR:HA	17	0.18
(1,899)	1:30:C:TYR:HA	1:30:C:TYR:HE1	3	0.18
(1,899)	1:30:C:TYR:HA	1:30:C:TYR:HE2	3	0.18
(1,867)	1:26:C:HIS:HA	1:26:C:HIS:HD2	16	0.18
(1,828)	1:28:C:PRO:HB2	1:31:C:PHE:HE1	14	0.18
(1,828)	1:28:C:PRO:HB2	1:31:C:PHE:HE2	14	0.18
(1,826)	1:25:C:GLY:H	2:40:D:PHE:HE1	6	0.18
(1,826)	1:25:C:GLY:H	2:40:D:PHE:HE2	6	0.18
(1,819)	1:22:C:GLY:H	1:23:C:GLY:H	19	0.18
(1,791)	1:26:C:HIS:HA	1:27:C:VAL:HG21	15	0.18
(1,791)	1:26:C:HIS:HA	1:27:C:VAL:HG22	15	0.18
(1,791)	1:26:C:HIS:HA	1:27:C:VAL:HG23	15	0.18
(1,790)	1:31:C:PHE:HD1	2:36:D:THR:HG21	14	0.18
(1,790)	1:31:C:PHE:HD1	2:36:D:THR:HG22	14	0.18
(1,790)	1:31:C:PHE:HD1	2:36:D:THR:HG23	14	0.18
(1,790)	1:31:C:PHE:HD2	2:36:D:THR:HG21	14	0.18
(1,790)	1:31:C:PHE:HD2	2:36:D:THR:HG22	14	0.18
(1,790)	1:31:C:PHE:HD2	2:36:D:THR:HG23	14	0.18
(1,782)	1:25:C:GLY:HA2	1:27:C:VAL:H	15	0.18
(1,781)	2:40:D:PHE:HA	2:41:D:TYR:HD1	13	0.18
(1,781)	2:40:D:PHE:HA	2:41:D:TYR:HD2	13	0.18
(1,778)	1:29:C:GLU:HG3	2:41:D:TYR:HB2	17	0.18
(1,778)	1:29:C:GLU:HG3	2:41:D:TYR:HB3	17	0.18
(1,748)	1:26:C:HIS:HB3	1:27:C:VAL:HG21	9	0.18
(1,748)	1:26:C:HIS:HB3	1:27:C:VAL:HG22	9	0.18
(1,748)	1:26:C:HIS:HB3	1:27:C:VAL:HG23	9	0.18
(1,743)	1:32:C:VAL:HG11	1:33:C:ARG:HA	5	0.18
(1,743)	1:32:C:VAL:HG12	1:33:C:ARG:HA	5	0.18
(1,743)	1:32:C:VAL:HG13	1:33:C:ARG:HA	5	0.18
(1,743)	1:32:C:VAL:HG21	1:33:C:ARG:HA	5	0.18

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,743)	1:32:C:VAL:HG22	1:33:C:ARG:HA	5	0.18
(1,743)	1:32:C:VAL:HG23	1:33:C:ARG:HA	5	0.18
(1,736)	1:28:C:PRO:HG2	1:31:C:PHE:HE1	1	0.18
(1,736)	1:28:C:PRO:HG2	1:31:C:PHE:HE2	1	0.18
(1,736)	1:28:C:PRO:HG3	1:31:C:PHE:HE1	1	0.18
(1,736)	1:28:C:PRO:HG3	1:31:C:PHE:HE2	1	0.18
(1,724)	1:27:C:VAL:HG11	1:28:C:PRO:HG2	13	0.18
(1,724)	1:27:C:VAL:HG11	1:28:C:PRO:HG3	13	0.18
(1,724)	1:27:C:VAL:HG12	1:28:C:PRO:HG2	13	0.18
(1,724)	1:27:C:VAL:HG12	1:28:C:PRO:HG3	13	0.18
(1,724)	1:27:C:VAL:HG13	1:28:C:PRO:HG2	13	0.18
(1,724)	1:27:C:VAL:HG13	1:28:C:PRO:HG3	13	0.18
(1,659)	1:11:A:VAL:HA	1:24:C:ALA:HB1	8	0.18
(1,659)	1:11:A:VAL:HA	1:24:C:ALA:HB2	8	0.18
(1,659)	1:11:A:VAL:HA	1:24:C:ALA:HB3	8	0.18
(1,650)	2:20:B:TYR:HA	2:20:B:TYR:HD1	13	0.18
(1,650)	2:20:B:TYR:HA	2:20:B:TYR:HD2	13	0.18
(1,629)	2:17:B:ILE:H	2:17:B:ILE:HD11	1	0.18
(1,629)	2:17:B:ILE:H	2:17:B:ILE:HD12	1	0.18
(1,629)	2:17:B:ILE:H	2:17:B:ILE:HD13	1	0.18
(1,479)	1:1:A:GLY:H	1:8:A:GLU:HB3	12	0.18
(1,479)	1:1:A:GLY:H	1:8:A:GLU:HB2	12	0.18
(1,463)	1:5:A:HIS:HB2	1:6:A:VAL:HG11	5	0.18
(1,463)	1:5:A:HIS:HB2	1:6:A:VAL:HG12	5	0.18
(1,463)	1:5:A:HIS:HB2	1:6:A:VAL:HG13	5	0.18
(1,399)	1:7:A:PRO:HG2	1:10:A:PHE:HE1	7	0.18
(1,399)	1:7:A:PRO:HG2	1:10:A:PHE:HE2	7	0.18
(1,399)	1:7:A:PRO:HG3	1:10:A:PHE:HE1	7	0.18
(1,399)	1:7:A:PRO:HG3	1:10:A:PHE:HE2	7	0.18
(1,389)	1:6:A:VAL:HB	2:20:B:TYR:HD1	10	0.18
(1,389)	1:6:A:VAL:HB	2:20:B:TYR:HD2	10	0.18
(1,379)	1:1:A:GLY:H	2:20:B:TYR:HE1	15	0.18
(1,379)	1:1:A:GLY:H	2:20:B:TYR:HE2	15	0.18
(1,370)	1:6:A:VAL:HG11	1:7:A:PRO:HD3	8	0.18
(1,370)	1:6:A:VAL:HG12	1:7:A:PRO:HD3	8	0.18
(1,370)	1:6:A:VAL:HG13	1:7:A:PRO:HD3	8	0.18
(1,349)	2:13:B:CYS:HB3	2:14:B:GLY:HA3	7	0.18
(1,346)	1:9:A:TYR:HE1	2:18:B:SER:HB2	3	0.18
(1,346)	1:9:A:TYR:HE2	2:18:B:SER:HB2	3	0.18
(1,331)	2:20:B:TYR:HA	2:20:B:TYR:HD1	13	0.18
(1,331)	2:20:B:TYR:HA	2:20:B:TYR:HD2	13	0.18
(1,310)	2:17:B:ILE:H	2:17:B:ILE:HD11	1	0.18

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,310)	2:17:B:ILE:H	2:17:B:ILE:HD12	1	0.18
(1,310)	2:17:B:ILE:H	2:17:B:ILE:HD13	1	0.18
(1,160)	1:1:A:GLY:H	1:8:A:GLU:HB3	12	0.18
(1,160)	1:1:A:GLY:H	1:8:A:GLU:HB2	12	0.18
(1,144)	1:5:A:HIS:HB2	1:6:A:VAL:HG11	5	0.18
(1,144)	1:5:A:HIS:HB2	1:6:A:VAL:HG12	5	0.18
(1,144)	1:5:A:HIS:HB2	1:6:A:VAL:HG13	5	0.18
(1,80)	1:7:A:PRO:HG2	1:10:A:PHE:HE1	7	0.18
(1,80)	1:7:A:PRO:HG2	1:10:A:PHE:HE2	7	0.18
(1,80)	1:7:A:PRO:HG3	1:10:A:PHE:HE1	7	0.18
(1,80)	1:7:A:PRO:HG3	1:10:A:PHE:HE2	7	0.18
(1,70)	1:6:A:VAL:HB	2:20:B:TYR:HD1	10	0.18
(1,70)	1:6:A:VAL:HB	2:20:B:TYR:HD2	10	0.18
(1,60)	1:1:A:GLY:H	2:20:B:TYR:HE1	15	0.18
(1,60)	1:1:A:GLY:H	2:20:B:TYR:HE2	15	0.18
(1,51)	1:6:A:VAL:HG11	1:7:A:PRO:HD3	8	0.18
(1,51)	1:6:A:VAL:HG12	1:7:A:PRO:HD3	8	0.18
(1,51)	1:6:A:VAL:HG13	1:7:A:PRO:HD3	8	0.18
(1,30)	2:13:B:CYS:HB3	2:14:B:GLY:HA3	7	0.18
(1,27)	1:9:A:TYR:HE1	2:18:B:SER:HB2	3	0.18
(1,27)	1:9:A:TYR:HE2	2:18:B:SER:HB2	3	0.18
(1,12)	1:11:A:VAL:HA	1:24:C:ALA:HB1	8	0.18
(1,12)	1:11:A:VAL:HA	1:24:C:ALA:HB2	8	0.18
(1,12)	1:11:A:VAL:HA	1:24:C:ALA:HB3	8	0.18
(1,1306)	2:41:D:TYR:HA	2:41:D:TYR:HD1	8	0.17
(1,1306)	2:41:D:TYR:HA	2:41:D:TYR:HD2	8	0.17
(1,1303)	2:41:D:TYR:H	2:41:D:TYR:HD1	13	0.17
(1,1303)	2:41:D:TYR:H	2:41:D:TYR:HD2	13	0.17
(1,1287)	2:38:D:ILE:HA	2:38:D:ILE:HD11	14	0.17
(1,1287)	2:38:D:ILE:HA	2:38:D:ILE:HD12	14	0.17
(1,1287)	2:38:D:ILE:HA	2:38:D:ILE:HD13	14	0.17
(1,1285)	2:38:D:ILE:H	2:38:D:ILE:HD11	17	0.17
(1,1285)	2:38:D:ILE:H	2:38:D:ILE:HD12	17	0.17
(1,1285)	2:38:D:ILE:H	2:38:D:ILE:HD13	17	0.17
(1,1224)	1:31:C:PHE:H	1:31:C:PHE:HD1	2	0.17
(1,1224)	1:31:C:PHE:H	1:31:C:PHE:HD2	2	0.17
(1,1221)	1:31:C:PHE:HA	1:32:C:VAL:H	1	0.17
(1,1170)	1:31:C:PHE:HB2	1:32:C:VAL:HG11	5	0.17
(1,1170)	1:31:C:PHE:HB2	1:32:C:VAL:HG12	5	0.17
(1,1170)	1:31:C:PHE:HB2	1:32:C:VAL:HG13	5	0.17
(1,1170)	1:31:C:PHE:HB2	1:32:C:VAL:HG21	5	0.17
(1,1170)	1:31:C:PHE:HB2	1:32:C:VAL:HG22	5	0.17

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,1170)	1:31:C:PHE:HB2	1:32:C:VAL:HG23	5	0.17
(1,1169)	2:36:D:THR:HB	2:38:D:ILE:HD11	12	0.17
(1,1169)	2:36:D:THR:HB	2:38:D:ILE:HD12	12	0.17
(1,1169)	2:36:D:THR:HB	2:38:D:ILE:HD13	12	0.17
(1,1167)	1:24:C:ALA:H	1:25:C:GLY:HA3	5	0.17
(1,1119)	1:26:C:HIS:HB2	1:27:C:VAL:HG11	4	0.17
(1,1119)	1:26:C:HIS:HB2	1:27:C:VAL:HG12	4	0.17
(1,1119)	1:26:C:HIS:HB2	1:27:C:VAL:HG13	4	0.17
(1,1078)	1:25:C:GLY:H	2:41:D:TYR:H	5	0.17
(1,1054)	1:28:C:PRO:HA	2:40:D:PHE:HD1	6	0.17
(1,1054)	1:28:C:PRO:HA	2:40:D:PHE:HD2	6	0.17
(1,1040)	1:31:C:PHE:HE1	2:37:D:PRO:HD2	11	0.17
(1,1040)	1:31:C:PHE:HE2	2:37:D:PRO:HD2	11	0.17
(1,987)	2:41:D:TYR:HA	2:41:D:TYR:HD1	8	0.17
(1,987)	2:41:D:TYR:HA	2:41:D:TYR:HD2	8	0.17
(1,984)	2:41:D:TYR:H	2:41:D:TYR:HD1	13	0.17
(1,984)	2:41:D:TYR:H	2:41:D:TYR:HD2	13	0.17
(1,968)	2:38:D:ILE:HA	2:38:D:ILE:HD11	14	0.17
(1,968)	2:38:D:ILE:HA	2:38:D:ILE:HD12	14	0.17
(1,968)	2:38:D:ILE:HA	2:38:D:ILE:HD13	14	0.17
(1,966)	2:38:D:ILE:H	2:38:D:ILE:HD11	17	0.17
(1,966)	2:38:D:ILE:H	2:38:D:ILE:HD12	17	0.17
(1,966)	2:38:D:ILE:H	2:38:D:ILE:HD13	17	0.17
(1,905)	1:31:C:PHE:H	1:31:C:PHE:HD1	2	0.17
(1,905)	1:31:C:PHE:H	1:31:C:PHE:HD2	2	0.17
(1,902)	1:31:C:PHE:HA	1:32:C:VAL:H	1	0.17
(1,851)	1:31:C:PHE:HB2	1:32:C:VAL:HG11	5	0.17
(1,851)	1:31:C:PHE:HB2	1:32:C:VAL:HG12	5	0.17
(1,851)	1:31:C:PHE:HB2	1:32:C:VAL:HG13	5	0.17
(1,851)	1:31:C:PHE:HB2	1:32:C:VAL:HG21	5	0.17
(1,851)	1:31:C:PHE:HB2	1:32:C:VAL:HG22	5	0.17
(1,851)	1:31:C:PHE:HB2	1:32:C:VAL:HG23	5	0.17
(1,850)	2:36:D:THR:HB	2:38:D:ILE:HD11	12	0.17
(1,850)	2:36:D:THR:HB	2:38:D:ILE:HD12	12	0.17
(1,850)	2:36:D:THR:HB	2:38:D:ILE:HD13	12	0.17
(1,848)	1:24:C:ALA:H	1:25:C:GLY:HA3	5	0.17
(1,800)	1:26:C:HIS:HB2	1:27:C:VAL:HG11	4	0.17
(1,800)	1:26:C:HIS:HB2	1:27:C:VAL:HG12	4	0.17
(1,800)	1:26:C:HIS:HB2	1:27:C:VAL:HG13	4	0.17
(1,759)	1:25:C:GLY:H	2:41:D:TYR:H	5	0.17
(1,735)	1:28:C:PRO:HA	2:40:D:PHE:HD1	6	0.17
(1,735)	1:28:C:PRO:HA	2:40:D:PHE:HD2	6	0.17

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,721)	1:31:C:PHE:HE1	2:37:D:PRO:HD2	11	0.17
(1,721)	1:31:C:PHE:HE2	2:37:D:PRO:HD2	11	0.17
(1,672)	2:17:B:ILE:HG13	2:40:D:PHE:HD1	5	0.17
(1,672)	2:17:B:ILE:HG13	2:40:D:PHE:HD2	5	0.17
(1,593)	1:12:A:ARG:HB2	1:12:A:ARG:HE	15	0.17
(1,562)	1:9:A:TYR:HA	1:9:A:TYR:HE1	11	0.17
(1,562)	1:9:A:TYR:HA	1:9:A:TYR:HE2	11	0.17
(1,562)	1:9:A:TYR:HA	1:9:A:TYR:HE1	17	0.17
(1,562)	1:9:A:TYR:HA	1:9:A:TYR:HE2	17	0.17
(1,501)	1:4:A:GLY:H	1:6:A:VAL:H	13	0.17
(1,499)	1:6:A:VAL:HG11	1:7:A:PRO:HD3	12	0.17
(1,499)	1:6:A:VAL:HG12	1:7:A:PRO:HD3	12	0.17
(1,499)	1:6:A:VAL:HG13	1:7:A:PRO:HD3	12	0.17
(1,499)	1:6:A:VAL:HG11	1:7:A:PRO:HD3	18	0.17
(1,499)	1:6:A:VAL:HG12	1:7:A:PRO:HD3	18	0.17
(1,499)	1:6:A:VAL:HG13	1:7:A:PRO:HD3	18	0.17
(1,472)	1:4:A:GLY:HA3	2:19:B:PHE:HB2	17	0.17
(1,472)	1:4:A:GLY:HA2	2:19:B:PHE:HB2	17	0.17
(1,461)	1:6:A:VAL:HG11	1:7:A:PRO:HB3	2	0.17
(1,461)	1:6:A:VAL:HG12	1:7:A:PRO:HB3	2	0.17
(1,461)	1:6:A:VAL:HG13	1:7:A:PRO:HB3	2	0.17
(1,453)	1:10:A:PHE:HD1	2:15:B:THR:HG21	3	0.17
(1,453)	1:10:A:PHE:HD1	2:15:B:THR:HG22	3	0.17
(1,453)	1:10:A:PHE:HD1	2:15:B:THR:HG23	3	0.17
(1,453)	1:10:A:PHE:HD2	2:15:B:THR:HG21	3	0.17
(1,453)	1:10:A:PHE:HD2	2:15:B:THR:HG22	3	0.17
(1,453)	1:10:A:PHE:HD2	2:15:B:THR:HG23	3	0.17
(1,451)	2:15:B:THR:HG21	2:16:B:PRO:HB3	4	0.17
(1,451)	2:15:B:THR:HG22	2:16:B:PRO:HB3	4	0.17
(1,451)	2:15:B:THR:HG23	2:16:B:PRO:HB3	4	0.17
(1,445)	1:4:A:GLY:HA2	1:6:A:VAL:H	3	0.17
(1,445)	1:4:A:GLY:HA2	1:6:A:VAL:H	5	0.17
(1,444)	2:19:B:PHE:HA	2:20:B:TYR:HD1	14	0.17
(1,444)	2:19:B:PHE:HA	2:20:B:TYR:HD2	14	0.17
(1,442)	1:8:A:GLU:HG3	2:19:B:PHE:HA	8	0.17
(1,422)	1:4:A:GLY:H	2:20:B:TYR:H	8	0.17
(1,410)	1:2:A:GLY:HA3	1:3:A:ALA:HB1	8	0.17
(1,410)	1:2:A:GLY:HA3	1:3:A:ALA:HB2	8	0.17
(1,410)	1:2:A:GLY:HA3	1:3:A:ALA:HB3	8	0.17
(1,410)	1:2:A:GLY:HA3	1:3:A:ALA:HB1	14	0.17
(1,410)	1:2:A:GLY:HA3	1:3:A:ALA:HB2	14	0.17
(1,410)	1:2:A:GLY:HA3	1:3:A:ALA:HB3	14	0.17

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,408)	2:15:B:THR:HG21	2:16:B:PRO:HA	3	0.17
(1,408)	2:15:B:THR:HG22	2:16:B:PRO:HA	3	0.17
(1,408)	2:15:B:THR:HG23	2:16:B:PRO:HA	3	0.17
(1,405)	1:1:A:GLY:H	2:18:B:SER:HB2	12	0.17
(1,403)	1:7:A:PRO:HG2	2:19:B:PHE:HD1	14	0.17
(1,403)	1:7:A:PRO:HG2	2:19:B:PHE:HD2	14	0.17
(1,403)	1:7:A:PRO:HG3	2:19:B:PHE:HD1	14	0.17
(1,403)	1:7:A:PRO:HG3	2:19:B:PHE:HD2	14	0.17
(1,379)	1:1:A:GLY:H	2:20:B:TYR:HE1	4	0.17
(1,379)	1:1:A:GLY:H	2:20:B:TYR:HE2	4	0.17
(1,370)	1:6:A:VAL:HG11	1:7:A:PRO:HD3	17	0.17
(1,370)	1:6:A:VAL:HG12	1:7:A:PRO:HD3	17	0.17
(1,370)	1:6:A:VAL:HG13	1:7:A:PRO:HD3	17	0.17
(1,362)	1:7:A:PRO:HG2	2:19:B:PHE:HD1	20	0.17
(1,362)	1:7:A:PRO:HG2	2:19:B:PHE:HD2	20	0.17
(1,362)	1:7:A:PRO:HG3	2:19:B:PHE:HD1	20	0.17
(1,362)	1:7:A:PRO:HG3	2:19:B:PHE:HD2	20	0.17
(1,274)	1:12:A:ARG:HB2	1:12:A:ARG:HE	15	0.17
(1,243)	1:9:A:TYR:HA	1:9:A:TYR:HE1	11	0.17
(1,243)	1:9:A:TYR:HA	1:9:A:TYR:HE2	11	0.17
(1,243)	1:9:A:TYR:HA	1:9:A:TYR:HE1	17	0.17
(1,243)	1:9:A:TYR:HA	1:9:A:TYR:HE2	17	0.17
(1,182)	1:4:A:GLY:H	1:6:A:VAL:H	13	0.17
(1,180)	1:6:A:VAL:HG11	1:7:A:PRO:HD3	12	0.17
(1,180)	1:6:A:VAL:HG12	1:7:A:PRO:HD3	12	0.17
(1,180)	1:6:A:VAL:HG13	1:7:A:PRO:HD3	12	0.17
(1,180)	1:6:A:VAL:HG11	1:7:A:PRO:HD3	18	0.17
(1,180)	1:6:A:VAL:HG12	1:7:A:PRO:HD3	18	0.17
(1,180)	1:6:A:VAL:HG13	1:7:A:PRO:HD3	18	0.17
(1,153)	1:4:A:GLY:HA3	2:19:B:PHE:HB2	17	0.17
(1,153)	1:4:A:GLY:HA2	2:19:B:PHE:HB2	17	0.17
(1,142)	1:6:A:VAL:HG11	1:7:A:PRO:HB3	2	0.17
(1,142)	1:6:A:VAL:HG12	1:7:A:PRO:HB3	2	0.17
(1,142)	1:6:A:VAL:HG13	1:7:A:PRO:HB3	2	0.17
(1,134)	1:10:A:PHE:HD1	2:15:B:THR:HG21	3	0.17
(1,134)	1:10:A:PHE:HD1	2:15:B:THR:HG22	3	0.17
(1,134)	1:10:A:PHE:HD1	2:15:B:THR:HG23	3	0.17
(1,134)	1:10:A:PHE:HD2	2:15:B:THR:HG21	3	0.17
(1,134)	1:10:A:PHE:HD2	2:15:B:THR:HG22	3	0.17
(1,134)	1:10:A:PHE:HD2	2:15:B:THR:HG23	3	0.17
(1,132)	2:15:B:THR:HG21	2:16:B:PRO:HB3	4	0.17
(1,132)	2:15:B:THR:HG22	2:16:B:PRO:HB3	4	0.17

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,132)	2:15:B:THR:HG23	2:16:B:PRO:HB3	4	0.17
(1,126)	1:4:A:GLY:HA2	1:6:A:VAL:H	3	0.17
(1,126)	1:4:A:GLY:HA2	1:6:A:VAL:H	5	0.17
(1,125)	2:19:B:PHE:HA	2:20:B:TYR:HD1	14	0.17
(1,125)	2:19:B:PHE:HA	2:20:B:TYR:HD2	14	0.17
(1,123)	1:8:A:GLU:HG3	2:19:B:PHE:HA	8	0.17
(1,103)	1:4:A:GLY:H	2:20:B:TYR:H	8	0.17
(1,91)	1:2:A:GLY:HA3	1:3:A:ALA:HB1	8	0.17
(1,91)	1:2:A:GLY:HA3	1:3:A:ALA:HB2	8	0.17
(1,91)	1:2:A:GLY:HA3	1:3:A:ALA:HB3	8	0.17
(1,91)	1:2:A:GLY:HA3	1:3:A:ALA:HB1	14	0.17
(1,91)	1:2:A:GLY:HA3	1:3:A:ALA:HB2	14	0.17
(1,91)	1:2:A:GLY:HA3	1:3:A:ALA:HB3	14	0.17
(1,89)	2:15:B:THR:HG21	2:16:B:PRO:HA	3	0.17
(1,89)	2:15:B:THR:HG22	2:16:B:PRO:HA	3	0.17
(1,89)	2:15:B:THR:HG23	2:16:B:PRO:HA	3	0.17
(1,86)	1:1:A:GLY:H	2:18:B:SER:HB2	12	0.17
(1,84)	1:7:A:PRO:HG2	2:19:B:PHE:HD1	14	0.17
(1,84)	1:7:A:PRO:HG2	2:19:B:PHE:HD2	14	0.17
(1,84)	1:7:A:PRO:HG3	2:19:B:PHE:HD1	14	0.17
(1,84)	1:7:A:PRO:HG3	2:19:B:PHE:HD2	14	0.17
(1,60)	1:1:A:GLY:H	2:20:B:TYR:HE1	4	0.17
(1,60)	1:1:A:GLY:H	2:20:B:TYR:HE2	4	0.17
(1,51)	1:6:A:VAL:HG11	1:7:A:PRO:HD3	17	0.17
(1,51)	1:6:A:VAL:HG12	1:7:A:PRO:HD3	17	0.17
(1,51)	1:6:A:VAL:HG13	1:7:A:PRO:HD3	17	0.17
(1,43)	1:7:A:PRO:HG2	2:19:B:PHE:HD1	20	0.17
(1,43)	1:7:A:PRO:HG2	2:19:B:PHE:HD2	20	0.17
(1,43)	1:7:A:PRO:HG3	2:19:B:PHE:HD1	20	0.17
(1,43)	1:7:A:PRO:HG3	2:19:B:PHE:HD2	20	0.17
(1,7)	2:17:B:ILE:HG13	2:40:D:PHE:HD1	5	0.17
(1,7)	2:17:B:ILE:HG13	2:40:D:PHE:HD2	5	0.17
(1,1303)	2:41:D:TYR:H	2:41:D:TYR:HD1	11	0.16
(1,1303)	2:41:D:TYR:H	2:41:D:TYR:HD2	11	0.16
(1,1242)	1:33:C:ARG:H	1:33:C:ARG:HB2	7	0.16
(1,1242)	1:33:C:ARG:H	1:33:C:ARG:HB3	7	0.16
(1,1242)	1:33:C:ARG:H	1:33:C:ARG:HB2	12	0.16
(1,1242)	1:33:C:ARG:H	1:33:C:ARG:HB3	12	0.16
(1,1224)	1:31:C:PHE:H	1:31:C:PHE:HD1	4	0.16
(1,1224)	1:31:C:PHE:H	1:31:C:PHE:HD2	4	0.16
(1,1224)	1:31:C:PHE:H	1:31:C:PHE:HD1	12	0.16
(1,1224)	1:31:C:PHE:H	1:31:C:PHE:HD2	12	0.16

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,1218)	1:30:C:TYR:HA	1:30:C:TYR:HE1	8	0.16
(1,1218)	1:30:C:TYR:HA	1:30:C:TYR:HE2	8	0.16
(1,1149)	1:32:C:VAL:HB	1:33:C:ARG:H	16	0.16
(1,1117)	1:27:C:VAL:HG11	1:28:C:PRO:HB3	5	0.16
(1,1117)	1:27:C:VAL:HG12	1:28:C:PRO:HB3	5	0.16
(1,1117)	1:27:C:VAL:HG13	1:28:C:PRO:HB3	5	0.16
(1,1108)	2:36:D:THR:HG21	2:37:D:PRO:HD3	3	0.16
(1,1108)	2:36:D:THR:HG22	2:37:D:PRO:HD3	3	0.16
(1,1108)	2:36:D:THR:HG23	2:37:D:PRO:HD3	3	0.16
(1,1102)	1:29:C:GLU:HB3	2:41:D:TYR:HD1	4	0.16
(1,1102)	1:29:C:GLU:HB3	2:41:D:TYR:HD2	4	0.16
(1,1101)	1:25:C:GLY:HA2	1:27:C:VAL:H	2	0.16
(1,1049)	1:28:C:PRO:HG2	1:31:C:PHE:HD1	18	0.16
(1,1049)	1:28:C:PRO:HG2	1:31:C:PHE:HD2	18	0.16
(1,1049)	1:28:C:PRO:HG3	1:31:C:PHE:HD1	18	0.16
(1,1049)	1:28:C:PRO:HG3	1:31:C:PHE:HD2	18	0.16
(1,1048)	1:32:C:VAL:HA	1:33:C:ARG:HG2	18	0.16
(1,1048)	1:32:C:VAL:HA	1:33:C:ARG:HG3	18	0.16
(1,1040)	1:31:C:PHE:HE1	2:37:D:PRO:HD2	13	0.16
(1,1040)	1:31:C:PHE:HE2	2:37:D:PRO:HD2	13	0.16
(1,1002)	1:30:C:TYR:HE1	2:39:D:SER:HB2	2	0.16
(1,1002)	1:30:C:TYR:HE2	2:39:D:SER:HB2	2	0.16
(1,1000)	1:31:C:PHE:HE1	2:37:D:PRO:HD2	1	0.16
(1,1000)	1:31:C:PHE:HE2	2:37:D:PRO:HD2	1	0.16
(1,984)	2:41:D:TYR:H	2:41:D:TYR:HD1	11	0.16
(1,984)	2:41:D:TYR:H	2:41:D:TYR:HD2	11	0.16
(1,923)	1:33:C:ARG:H	1:33:C:ARG:HB2	7	0.16
(1,923)	1:33:C:ARG:H	1:33:C:ARG:HB3	7	0.16
(1,923)	1:33:C:ARG:H	1:33:C:ARG:HB2	12	0.16
(1,923)	1:33:C:ARG:H	1:33:C:ARG:HB3	12	0.16
(1,905)	1:31:C:PHE:H	1:31:C:PHE:HD1	4	0.16
(1,905)	1:31:C:PHE:H	1:31:C:PHE:HD2	4	0.16
(1,905)	1:31:C:PHE:H	1:31:C:PHE:HD1	12	0.16
(1,905)	1:31:C:PHE:H	1:31:C:PHE:HD2	12	0.16
(1,899)	1:30:C:TYR:HA	1:30:C:TYR:HE1	8	0.16
(1,899)	1:30:C:TYR:HA	1:30:C:TYR:HE2	8	0.16
(1,830)	1:32:C:VAL:HB	1:33:C:ARG:H	16	0.16
(1,798)	1:27:C:VAL:HG11	1:28:C:PRO:HB3	5	0.16
(1,798)	1:27:C:VAL:HG12	1:28:C:PRO:HB3	5	0.16
(1,798)	1:27:C:VAL:HG13	1:28:C:PRO:HB3	5	0.16
(1,789)	2:36:D:THR:HG21	2:37:D:PRO:HD3	3	0.16
(1,789)	2:36:D:THR:HG22	2:37:D:PRO:HD3	3	0.16

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,789)	2:36:D:THR:HG23	2:37:D:PRO:HD3	3	0.16
(1,783)	1:29:C:GLU:HB3	2:41:D:TYR:HD1	4	0.16
(1,783)	1:29:C:GLU:HB3	2:41:D:TYR:HD2	4	0.16
(1,782)	1:25:C:GLY:HA2	1:27:C:VAL:H	2	0.16
(1,730)	1:28:C:PRO:HG2	1:31:C:PHE:HD1	18	0.16
(1,730)	1:28:C:PRO:HG2	1:31:C:PHE:HD2	18	0.16
(1,730)	1:28:C:PRO:HG3	1:31:C:PHE:HD1	18	0.16
(1,730)	1:28:C:PRO:HG3	1:31:C:PHE:HD2	18	0.16
(1,729)	1:32:C:VAL:HA	1:33:C:ARG:HG2	18	0.16
(1,729)	1:32:C:VAL:HA	1:33:C:ARG:HG3	18	0.16
(1,721)	1:31:C:PHE:HE1	2:37:D:PRO:HD2	13	0.16
(1,721)	1:31:C:PHE:HE2	2:37:D:PRO:HD2	13	0.16
(1,683)	1:30:C:TYR:HE1	2:39:D:SER:HB2	2	0.16
(1,683)	1:30:C:TYR:HE2	2:39:D:SER:HB2	2	0.16
(1,681)	1:31:C:PHE:HE1	2:37:D:PRO:HD2	1	0.16
(1,681)	1:31:C:PHE:HE2	2:37:D:PRO:HD2	1	0.16
(1,568)	1:10:A:PHE:H	1:10:A:PHE:HD1	7	0.16
(1,568)	1:10:A:PHE:H	1:10:A:PHE:HD2	7	0.16
(1,565)	1:10:A:PHE:HA	1:11:A:VAL:H	6	0.16
(1,562)	1:9:A:TYR:HA	1:9:A:TYR:HE1	5	0.16
(1,562)	1:9:A:TYR:HA	1:9:A:TYR:HE2	5	0.16
(1,562)	1:9:A:TYR:HA	1:9:A:TYR:HE1	13	0.16
(1,562)	1:9:A:TYR:HA	1:9:A:TYR:HE2	13	0.16
(1,562)	1:9:A:TYR:HA	1:9:A:TYR:HE1	18	0.16
(1,562)	1:9:A:TYR:HA	1:9:A:TYR:HE2	18	0.16
(1,530)	1:5:A:HIS:HA	1:5:A:HIS:HD2	11	0.16
(1,515)	2:15:B:THR:HG21	2:17:B:ILE:HD11	8	0.16
(1,515)	2:15:B:THR:HG21	2:17:B:ILE:HD12	8	0.16
(1,515)	2:15:B:THR:HG21	2:17:B:ILE:HD13	8	0.16
(1,515)	2:15:B:THR:HG22	2:17:B:ILE:HD11	8	0.16
(1,515)	2:15:B:THR:HG22	2:17:B:ILE:HD12	8	0.16
(1,515)	2:15:B:THR:HG22	2:17:B:ILE:HD13	8	0.16
(1,515)	2:15:B:THR:HG23	2:17:B:ILE:HD11	8	0.16
(1,515)	2:15:B:THR:HG23	2:17:B:ILE:HD12	8	0.16
(1,515)	2:15:B:THR:HG23	2:17:B:ILE:HD13	8	0.16
(1,514)	1:10:A:PHE:HB2	1:11:A:VAL:HG11	12	0.16
(1,514)	1:10:A:PHE:HB2	1:11:A:VAL:HG12	12	0.16
(1,514)	1:10:A:PHE:HB2	1:11:A:VAL:HG13	12	0.16
(1,514)	1:10:A:PHE:HB2	1:11:A:VAL:HG21	12	0.16
(1,514)	1:10:A:PHE:HB2	1:11:A:VAL:HG22	12	0.16
(1,514)	1:10:A:PHE:HB2	1:11:A:VAL:HG23	12	0.16
(1,469)	1:10:A:PHE:HD1	1:11:A:VAL:HG11	19	0.16

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,469)	1:10:A:PHE:HD1	1:11:A:VAL:HG12	19	0.16
(1,469)	1:10:A:PHE:HD1	1:11:A:VAL:HG13	19	0.16
(1,469)	1:10:A:PHE:HD1	1:11:A:VAL:HG21	19	0.16
(1,469)	1:10:A:PHE:HD1	1:11:A:VAL:HG22	19	0.16
(1,469)	1:10:A:PHE:HD1	1:11:A:VAL:HG23	19	0.16
(1,469)	1:10:A:PHE:HD2	1:11:A:VAL:HG11	19	0.16
(1,469)	1:10:A:PHE:HD2	1:11:A:VAL:HG12	19	0.16
(1,469)	1:10:A:PHE:HD2	1:11:A:VAL:HG13	19	0.16
(1,469)	1:10:A:PHE:HD2	1:11:A:VAL:HG21	19	0.16
(1,469)	1:10:A:PHE:HD2	1:11:A:VAL:HG22	19	0.16
(1,469)	1:10:A:PHE:HD2	1:11:A:VAL:HG23	19	0.16
(1,453)	1:10:A:PHE:HD1	2:15:B:THR:HG21	18	0.16
(1,453)	1:10:A:PHE:HD1	2:15:B:THR:HG22	18	0.16
(1,453)	1:10:A:PHE:HD1	2:15:B:THR:HG23	18	0.16
(1,453)	1:10:A:PHE:HD2	2:15:B:THR:HG21	18	0.16
(1,453)	1:10:A:PHE:HD2	2:15:B:THR:HG22	18	0.16
(1,453)	1:10:A:PHE:HD2	2:15:B:THR:HG23	18	0.16
(1,446)	1:8:A:GLU:HB3	2:20:B:TYR:HD1	4	0.16
(1,446)	1:8:A:GLU:HB3	2:20:B:TYR:HD2	4	0.16
(1,393)	1:7:A:PRO:HG2	1:10:A:PHE:HD1	9	0.16
(1,393)	1:7:A:PRO:HG2	1:10:A:PHE:HD2	9	0.16
(1,393)	1:7:A:PRO:HG3	1:10:A:PHE:HD1	9	0.16
(1,393)	1:7:A:PRO:HG3	1:10:A:PHE:HD2	9	0.16
(1,387)	1:6:A:VAL:HG11	1:7:A:PRO:HG2	6	0.16
(1,387)	1:6:A:VAL:HG11	1:7:A:PRO:HG3	6	0.16
(1,387)	1:6:A:VAL:HG12	1:7:A:PRO:HG2	6	0.16
(1,387)	1:6:A:VAL:HG12	1:7:A:PRO:HG3	6	0.16
(1,387)	1:6:A:VAL:HG13	1:7:A:PRO:HG2	6	0.16
(1,387)	1:6:A:VAL:HG13	1:7:A:PRO:HG3	6	0.16
(1,379)	1:1:A:GLY:H	2:20:B:TYR:HE1	12	0.16
(1,379)	1:1:A:GLY:H	2:20:B:TYR:HE2	12	0.16
(1,249)	1:10:A:PHE:H	1:10:A:PHE:HD1	7	0.16
(1,249)	1:10:A:PHE:H	1:10:A:PHE:HD2	7	0.16
(1,246)	1:10:A:PHE:HA	1:11:A:VAL:H	6	0.16
(1,243)	1:9:A:TYR:HA	1:9:A:TYR:HE1	5	0.16
(1,243)	1:9:A:TYR:HA	1:9:A:TYR:HE2	5	0.16
(1,243)	1:9:A:TYR:HA	1:9:A:TYR:HE1	13	0.16
(1,243)	1:9:A:TYR:HA	1:9:A:TYR:HE2	13	0.16
(1,243)	1:9:A:TYR:HA	1:9:A:TYR:HE1	18	0.16
(1,243)	1:9:A:TYR:HA	1:9:A:TYR:HE2	18	0.16
(1,211)	1:5:A:HIS:HA	1:5:A:HIS:HD2	11	0.16
(1,196)	2:15:B:THR:HG21	2:17:B:ILE:HD11	8	0.16

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,196)	2:15:B:THR:HG21	2:17:B:ILE:HD12	8	0.16
(1,196)	2:15:B:THR:HG21	2:17:B:ILE:HD13	8	0.16
(1,196)	2:15:B:THR:HG22	2:17:B:ILE:HD11	8	0.16
(1,196)	2:15:B:THR:HG22	2:17:B:ILE:HD12	8	0.16
(1,196)	2:15:B:THR:HG22	2:17:B:ILE:HD13	8	0.16
(1,196)	2:15:B:THR:HG23	2:17:B:ILE:HD11	8	0.16
(1,196)	2:15:B:THR:HG23	2:17:B:ILE:HD12	8	0.16
(1,196)	2:15:B:THR:HG23	2:17:B:ILE:HD13	8	0.16
(1,195)	1:10:A:PHE:HB2	1:11:A:VAL:HG11	12	0.16
(1,195)	1:10:A:PHE:HB2	1:11:A:VAL:HG12	12	0.16
(1,195)	1:10:A:PHE:HB2	1:11:A:VAL:HG13	12	0.16
(1,195)	1:10:A:PHE:HB2	1:11:A:VAL:HG21	12	0.16
(1,195)	1:10:A:PHE:HB2	1:11:A:VAL:HG22	12	0.16
(1,195)	1:10:A:PHE:HB2	1:11:A:VAL:HG23	12	0.16
(1,150)	1:10:A:PHE:HD1	1:11:A:VAL:HG11	19	0.16
(1,150)	1:10:A:PHE:HD1	1:11:A:VAL:HG12	19	0.16
(1,150)	1:10:A:PHE:HD1	1:11:A:VAL:HG13	19	0.16
(1,150)	1:10:A:PHE:HD1	1:11:A:VAL:HG21	19	0.16
(1,150)	1:10:A:PHE:HD1	1:11:A:VAL:HG22	19	0.16
(1,150)	1:10:A:PHE:HD1	1:11:A:VAL:HG23	19	0.16
(1,150)	1:10:A:PHE:HD2	1:11:A:VAL:HG11	19	0.16
(1,150)	1:10:A:PHE:HD2	1:11:A:VAL:HG12	19	0.16
(1,150)	1:10:A:PHE:HD2	1:11:A:VAL:HG13	19	0.16
(1,150)	1:10:A:PHE:HD2	1:11:A:VAL:HG21	19	0.16
(1,150)	1:10:A:PHE:HD2	1:11:A:VAL:HG22	19	0.16
(1,150)	1:10:A:PHE:HD2	1:11:A:VAL:HG23	19	0.16
(1,134)	1:10:A:PHE:HD1	2:15:B:THR:HG21	18	0.16
(1,134)	1:10:A:PHE:HD1	2:15:B:THR:HG22	18	0.16
(1,134)	1:10:A:PHE:HD1	2:15:B:THR:HG23	18	0.16
(1,134)	1:10:A:PHE:HD2	2:15:B:THR:HG21	18	0.16
(1,134)	1:10:A:PHE:HD2	2:15:B:THR:HG22	18	0.16
(1,134)	1:10:A:PHE:HD2	2:15:B:THR:HG23	18	0.16
(1,127)	1:8:A:GLU:HB3	2:20:B:TYR:HD1	4	0.16
(1,127)	1:8:A:GLU:HB3	2:20:B:TYR:HD2	4	0.16
(1,74)	1:7:A:PRO:HG2	1:10:A:PHE:HD1	9	0.16
(1,74)	1:7:A:PRO:HG2	1:10:A:PHE:HD2	9	0.16
(1,74)	1:7:A:PRO:HG3	1:10:A:PHE:HD1	9	0.16
(1,74)	1:7:A:PRO:HG3	1:10:A:PHE:HD2	9	0.16
(1,68)	1:6:A:VAL:HG11	1:7:A:PRO:HG2	6	0.16
(1,68)	1:6:A:VAL:HG11	1:7:A:PRO:HG3	6	0.16
(1,68)	1:6:A:VAL:HG12	1:7:A:PRO:HG2	6	0.16
(1,68)	1:6:A:VAL:HG12	1:7:A:PRO:HG3	6	0.16

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,68)	1:6:A:VAL:HG13	1:7:A:PRO:HG2	6	0.16
(1,68)	1:6:A:VAL:HG13	1:7:A:PRO:HG3	6	0.16
(1,60)	1:1:A:GLY:H	2:20:B:TYR:HE1	12	0.16
(1,60)	1:1:A:GLY:H	2:20:B:TYR:HE2	12	0.16
(1,1242)	1:33:C:ARG:H	1:33:C:ARG:HB2	2	0.15
(1,1242)	1:33:C:ARG:H	1:33:C:ARG:HB3	2	0.15
(1,1109)	1:31:C:PHE:HD1	2:36:D:THR:HG21	9	0.15
(1,1109)	1:31:C:PHE:HD1	2:36:D:THR:HG22	9	0.15
(1,1109)	1:31:C:PHE:HD1	2:36:D:THR:HG23	9	0.15
(1,1109)	1:31:C:PHE:HD2	2:36:D:THR:HG21	9	0.15
(1,1109)	1:31:C:PHE:HD2	2:36:D:THR:HG22	9	0.15
(1,1109)	1:31:C:PHE:HD2	2:36:D:THR:HG23	9	0.15
(1,1108)	2:36:D:THR:HG21	2:37:D:PRO:HD3	14	0.15
(1,1108)	2:36:D:THR:HG22	2:37:D:PRO:HD3	14	0.15
(1,1108)	2:36:D:THR:HG23	2:37:D:PRO:HD3	14	0.15
(1,1104)	1:25:C:GLY:H	2:40:D:PHE:H	13	0.15
(1,1104)	1:25:C:GLY:H	2:40:D:PHE:H	18	0.15
(1,1091)	1:24:C:ALA:H	1:25:C:GLY:H	11	0.15
(1,1040)	1:31:C:PHE:HE1	2:37:D:PRO:HD2	16	0.15
(1,1040)	1:31:C:PHE:HE2	2:37:D:PRO:HD2	16	0.15
(1,1005)	2:34:D:CYS:HB3	2:35:D:GLY:HA3	6	0.15
(1,1002)	1:30:C:TYR:HE1	2:39:D:SER:HB2	1	0.15
(1,1002)	1:30:C:TYR:HE2	2:39:D:SER:HB2	1	0.15
(1,1001)	1:30:C:TYR:HD1	2:39:D:SER:HB2	14	0.15
(1,1001)	1:30:C:TYR:HD2	2:39:D:SER:HB2	14	0.15
(1,923)	1:33:C:ARG:H	1:33:C:ARG:HB2	2	0.15
(1,923)	1:33:C:ARG:H	1:33:C:ARG:HB3	2	0.15
(1,790)	1:31:C:PHE:HD1	2:36:D:THR:HG21	9	0.15
(1,790)	1:31:C:PHE:HD1	2:36:D:THR:HG22	9	0.15
(1,790)	1:31:C:PHE:HD1	2:36:D:THR:HG23	9	0.15
(1,790)	1:31:C:PHE:HD2	2:36:D:THR:HG21	9	0.15
(1,790)	1:31:C:PHE:HD2	2:36:D:THR:HG22	9	0.15
(1,790)	1:31:C:PHE:HD2	2:36:D:THR:HG23	9	0.15
(1,789)	2:36:D:THR:HG21	2:37:D:PRO:HD3	14	0.15
(1,789)	2:36:D:THR:HG22	2:37:D:PRO:HD3	14	0.15
(1,789)	2:36:D:THR:HG23	2:37:D:PRO:HD3	14	0.15
(1,785)	1:25:C:GLY:H	2:40:D:PHE:H	13	0.15
(1,785)	1:25:C:GLY:H	2:40:D:PHE:H	18	0.15
(1,772)	1:24:C:ALA:H	1:25:C:GLY:H	11	0.15
(1,721)	1:31:C:PHE:HE1	2:37:D:PRO:HD2	16	0.15
(1,721)	1:31:C:PHE:HE2	2:37:D:PRO:HD2	16	0.15
(1,686)	2:34:D:CYS:HB3	2:35:D:GLY:HA3	6	0.15

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,683)	1:30:C:TYR:HE1	2:39:D:SER:HB2	1	0.15
(1,683)	1:30:C:TYR:HE2	2:39:D:SER:HB2	1	0.15
(1,682)	1:30:C:TYR:HD1	2:39:D:SER:HB2	14	0.15
(1,682)	1:30:C:TYR:HD2	2:39:D:SER:HB2	14	0.15
(1,670)	2:17:B:ILE:HD11	2:40:D:PHE:H	16	0.15
(1,670)	2:17:B:ILE:HD12	2:40:D:PHE:H	16	0.15
(1,670)	2:17:B:ILE:HD13	2:40:D:PHE:H	16	0.15
(1,568)	1:10:A:PHE:H	1:10:A:PHE:HD1	16	0.15
(1,568)	1:10:A:PHE:H	1:10:A:PHE:HD2	16	0.15
(1,511)	1:3:A:ALA:H	1:4:A:GLY:HA3	1	0.15
(1,474)	1:6:A:VAL:H	2:19:B:PHE:HB2	20	0.15
(1,407)	1:1:A:GLY:H	2:18:B:SER:HA	16	0.15
(1,398)	1:7:A:PRO:HA	2:19:B:PHE:HD1	3	0.15
(1,398)	1:7:A:PRO:HA	2:19:B:PHE:HD2	3	0.15
(1,387)	1:6:A:VAL:HG11	1:7:A:PRO:HG2	2	0.15
(1,387)	1:6:A:VAL:HG11	1:7:A:PRO:HG3	2	0.15
(1,387)	1:6:A:VAL:HG12	1:7:A:PRO:HG2	2	0.15
(1,387)	1:6:A:VAL:HG12	1:7:A:PRO:HG3	2	0.15
(1,387)	1:6:A:VAL:HG13	1:7:A:PRO:HG2	2	0.15
(1,387)	1:6:A:VAL:HG13	1:7:A:PRO:HG3	2	0.15
(1,344)	1:10:A:PHE:HE1	2:16:B:PRO:HD2	12	0.15
(1,344)	1:10:A:PHE:HE2	2:16:B:PRO:HD2	12	0.15
(1,344)	1:10:A:PHE:HE1	2:16:B:PRO:HD2	14	0.15
(1,344)	1:10:A:PHE:HE2	2:16:B:PRO:HD2	14	0.15
(1,249)	1:10:A:PHE:H	1:10:A:PHE:HD1	16	0.15
(1,249)	1:10:A:PHE:H	1:10:A:PHE:HD2	16	0.15
(1,192)	1:3:A:ALA:H	1:4:A:GLY:HA3	1	0.15
(1,155)	1:6:A:VAL:H	2:19:B:PHE:HB2	20	0.15
(1,88)	1:1:A:GLY:H	2:18:B:SER:HA	16	0.15
(1,79)	1:7:A:PRO:HA	2:19:B:PHE:HD1	3	0.15
(1,79)	1:7:A:PRO:HA	2:19:B:PHE:HD2	3	0.15
(1,68)	1:6:A:VAL:HG11	1:7:A:PRO:HG2	2	0.15
(1,68)	1:6:A:VAL:HG11	1:7:A:PRO:HG3	2	0.15
(1,68)	1:6:A:VAL:HG12	1:7:A:PRO:HG2	2	0.15
(1,68)	1:6:A:VAL:HG12	1:7:A:PRO:HG3	2	0.15
(1,68)	1:6:A:VAL:HG13	1:7:A:PRO:HG2	2	0.15
(1,68)	1:6:A:VAL:HG13	1:7:A:PRO:HG3	2	0.15
(1,25)	1:10:A:PHE:HE1	2:16:B:PRO:HD2	12	0.15
(1,25)	1:10:A:PHE:HE2	2:16:B:PRO:HD2	12	0.15
(1,25)	1:10:A:PHE:HE1	2:16:B:PRO:HD2	14	0.15
(1,25)	1:10:A:PHE:HE2	2:16:B:PRO:HD2	14	0.15
(1,5)	2:17:B:ILE:HD11	2:40:D:PHE:H	16	0.15

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,5)	2:17:B:ILE:HD12	2:40:D:PHE:H	16	0.15
(1,5)	2:17:B:ILE:HD13	2:40:D:PHE:H	16	0.15
(1,1186)	1:26:C:HIS:HA	1:26:C:HIS:HD2	5	0.14
(1,1166)	1:24:C:ALA:H	2:40:D:PHE:HD1	13	0.14
(1,1166)	1:24:C:ALA:H	2:40:D:PHE:HD2	13	0.14
(1,1159)	1:28:C:PRO:HB3	1:31:C:PHE:HE1	1	0.14
(1,1159)	1:28:C:PRO:HB3	1:31:C:PHE:HE2	1	0.14
(1,1156)	2:34:D:CYS:HB3	2:35:D:GLY:H	2	0.14
(1,1139)	1:22:C:GLY:H	2:41:D:TYR:HD1	11	0.14
(1,1139)	1:22:C:GLY:H	2:41:D:TYR:HD2	11	0.14
(1,1124)	1:32:C:VAL:HG11	1:33:C:ARG:H	19	0.14
(1,1124)	1:32:C:VAL:HG12	1:33:C:ARG:H	19	0.14
(1,1124)	1:32:C:VAL:HG13	1:33:C:ARG:H	19	0.14
(1,1124)	1:32:C:VAL:HG21	1:33:C:ARG:H	19	0.14
(1,1124)	1:32:C:VAL:HG22	1:33:C:ARG:H	19	0.14
(1,1124)	1:32:C:VAL:HG23	1:33:C:ARG:H	19	0.14
(1,1104)	1:25:C:GLY:H	2:40:D:PHE:H	5	0.14
(1,1097)	1:29:C:GLU:HG3	2:41:D:TYR:HB2	2	0.14
(1,1097)	1:29:C:GLU:HG3	2:41:D:TYR:HB3	2	0.14
(1,1059)	1:28:C:PRO:HG2	2:40:D:PHE:HD1	3	0.14
(1,1059)	1:28:C:PRO:HG2	2:40:D:PHE:HD2	3	0.14
(1,1059)	1:28:C:PRO:HG3	2:40:D:PHE:HD1	3	0.14
(1,1059)	1:28:C:PRO:HG3	2:40:D:PHE:HD2	3	0.14
(1,1049)	1:28:C:PRO:HG2	1:31:C:PHE:HD1	9	0.14
(1,1049)	1:28:C:PRO:HG2	1:31:C:PHE:HD2	9	0.14
(1,1049)	1:28:C:PRO:HG3	1:31:C:PHE:HD1	9	0.14
(1,1049)	1:28:C:PRO:HG3	1:31:C:PHE:HD2	9	0.14
(1,1035)	1:22:C:GLY:H	2:41:D:TYR:HE1	10	0.14
(1,1035)	1:22:C:GLY:H	2:41:D:TYR:HE2	10	0.14
(1,1027)	1:24:C:ALA:H	2:40:D:PHE:HD1	13	0.14
(1,1027)	1:24:C:ALA:H	2:40:D:PHE:HD2	13	0.14
(1,1020)	1:27:C:VAL:HB	2:41:D:TYR:H	1	0.14
(1,1015)	1:30:C:TYR:HD1	2:39:D:SER:HA	10	0.14
(1,1015)	1:30:C:TYR:HD2	2:39:D:SER:HA	10	0.14
(1,1005)	2:34:D:CYS:HB3	2:35:D:GLY:HA3	18	0.14
(1,867)	1:26:C:HIS:HA	1:26:C:HIS:HD2	5	0.14
(1,847)	1:24:C:ALA:H	2:40:D:PHE:HD1	13	0.14
(1,847)	1:24:C:ALA:H	2:40:D:PHE:HD2	13	0.14
(1,840)	1:28:C:PRO:HB3	1:31:C:PHE:HE1	1	0.14
(1,840)	1:28:C:PRO:HB3	1:31:C:PHE:HE2	1	0.14
(1,837)	2:34:D:CYS:HB3	2:35:D:GLY:H	2	0.14
(1,820)	1:22:C:GLY:H	2:41:D:TYR:HD1	11	0.14

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,820)	1:22:C:GLY:H	2:41:D:TYR:HD2	11	0.14
(1,805)	1:32:C:VAL:HG11	1:33:C:ARG:H	19	0.14
(1,805)	1:32:C:VAL:HG12	1:33:C:ARG:H	19	0.14
(1,805)	1:32:C:VAL:HG13	1:33:C:ARG:H	19	0.14
(1,805)	1:32:C:VAL:HG21	1:33:C:ARG:H	19	0.14
(1,805)	1:32:C:VAL:HG22	1:33:C:ARG:H	19	0.14
(1,805)	1:32:C:VAL:HG23	1:33:C:ARG:H	19	0.14
(1,785)	1:25:C:GLY:H	2:40:D:PHE:H	5	0.14
(1,778)	1:29:C:GLU:HG3	2:41:D:TYR:HB2	2	0.14
(1,778)	1:29:C:GLU:HG3	2:41:D:TYR:HB3	2	0.14
(1,740)	1:28:C:PRO:HG2	2:40:D:PHE:HD1	3	0.14
(1,740)	1:28:C:PRO:HG2	2:40:D:PHE:HD2	3	0.14
(1,740)	1:28:C:PRO:HG3	2:40:D:PHE:HD1	3	0.14
(1,740)	1:28:C:PRO:HG3	2:40:D:PHE:HD2	3	0.14
(1,730)	1:28:C:PRO:HG2	1:31:C:PHE:HD1	9	0.14
(1,730)	1:28:C:PRO:HG2	1:31:C:PHE:HD2	9	0.14
(1,730)	1:28:C:PRO:HG3	1:31:C:PHE:HD1	9	0.14
(1,730)	1:28:C:PRO:HG3	1:31:C:PHE:HD2	9	0.14
(1,716)	1:22:C:GLY:H	2:41:D:TYR:HE1	10	0.14
(1,716)	1:22:C:GLY:H	2:41:D:TYR:HE2	10	0.14
(1,708)	1:24:C:ALA:H	2:40:D:PHE:HD1	13	0.14
(1,708)	1:24:C:ALA:H	2:40:D:PHE:HD2	13	0.14
(1,701)	1:27:C:VAL:HB	2:41:D:TYR:H	1	0.14
(1,696)	1:30:C:TYR:HD1	2:39:D:SER:HA	10	0.14
(1,696)	1:30:C:TYR:HD2	2:39:D:SER:HA	10	0.14
(1,686)	2:34:D:CYS:HB3	2:35:D:GLY:HA3	18	0.14
(1,514)	1:10:A:PHE:HB2	1:11:A:VAL:HG11	2	0.14
(1,514)	1:10:A:PHE:HB2	1:11:A:VAL:HG12	2	0.14
(1,514)	1:10:A:PHE:HB2	1:11:A:VAL:HG13	2	0.14
(1,514)	1:10:A:PHE:HB2	1:11:A:VAL:HG21	2	0.14
(1,514)	1:10:A:PHE:HB2	1:11:A:VAL:HG22	2	0.14
(1,514)	1:10:A:PHE:HB2	1:11:A:VAL:HG23	2	0.14
(1,474)	1:6:A:VAL:H	2:19:B:PHE:HB2	18	0.14
(1,461)	1:6:A:VAL:HG11	1:7:A:PRO:HB3	20	0.14
(1,461)	1:6:A:VAL:HG12	1:7:A:PRO:HB3	20	0.14
(1,461)	1:6:A:VAL:HG13	1:7:A:PRO:HB3	20	0.14
(1,433)	1:8:A:GLU:HB2	2:20:B:TYR:HB2	9	0.14
(1,433)	1:8:A:GLU:HB2	2:20:B:TYR:HB3	9	0.14
(1,395)	2:13:B:CYS:HB2	2:14:B:GLY:H	15	0.14
(1,394)	1:7:A:PRO:HG2	1:10:A:PHE:HB2	16	0.14
(1,394)	1:7:A:PRO:HG3	1:10:A:PHE:HB2	16	0.14
(1,384)	1:10:A:PHE:HE1	2:16:B:PRO:HD2	4	0.14

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,384)	1:10:A:PHE:HE2	2:16:B:PRO:HD2	4	0.14
(1,369)	1:6:A:VAL:HG21	2:20:B:TYR:HE1	11	0.14
(1,369)	1:6:A:VAL:HG21	2:20:B:TYR:HE2	11	0.14
(1,369)	1:6:A:VAL:HG22	2:20:B:TYR:HE1	11	0.14
(1,369)	1:6:A:VAL:HG22	2:20:B:TYR:HE2	11	0.14
(1,369)	1:6:A:VAL:HG23	2:20:B:TYR:HE1	11	0.14
(1,369)	1:6:A:VAL:HG23	2:20:B:TYR:HE2	11	0.14
(1,362)	1:7:A:PRO:HG2	2:19:B:PHE:HD1	4	0.14
(1,362)	1:7:A:PRO:HG2	2:19:B:PHE:HD2	4	0.14
(1,362)	1:7:A:PRO:HG3	2:19:B:PHE:HD1	4	0.14
(1,362)	1:7:A:PRO:HG3	2:19:B:PHE:HD2	4	0.14
(1,360)	1:7:A:PRO:HG2	1:10:A:PHE:HB2	16	0.14
(1,360)	1:7:A:PRO:HG3	1:10:A:PHE:HB2	16	0.14
(1,349)	2:13:B:CYS:HB3	2:14:B:GLY:HA3	13	0.14
(1,344)	1:10:A:PHE:HE1	2:16:B:PRO:HD2	15	0.14
(1,344)	1:10:A:PHE:HE2	2:16:B:PRO:HD2	15	0.14
(1,195)	1:10:A:PHE:HB2	1:11:A:VAL:HG11	2	0.14
(1,195)	1:10:A:PHE:HB2	1:11:A:VAL:HG12	2	0.14
(1,195)	1:10:A:PHE:HB2	1:11:A:VAL:HG13	2	0.14
(1,195)	1:10:A:PHE:HB2	1:11:A:VAL:HG21	2	0.14
(1,195)	1:10:A:PHE:HB2	1:11:A:VAL:HG22	2	0.14
(1,195)	1:10:A:PHE:HB2	1:11:A:VAL:HG23	2	0.14
(1,155)	1:6:A:VAL:H	2:19:B:PHE:HB2	18	0.14
(1,142)	1:6:A:VAL:HG11	1:7:A:PRO:HB3	20	0.14
(1,142)	1:6:A:VAL:HG12	1:7:A:PRO:HB3	20	0.14
(1,142)	1:6:A:VAL:HG13	1:7:A:PRO:HB3	20	0.14
(1,114)	1:8:A:GLU:HB2	2:20:B:TYR:HB2	9	0.14
(1,114)	1:8:A:GLU:HB2	2:20:B:TYR:HB3	9	0.14
(1,76)	2:13:B:CYS:HB2	2:14:B:GLY:H	15	0.14
(1,75)	1:7:A:PRO:HG2	1:10:A:PHE:HB2	16	0.14
(1,75)	1:7:A:PRO:HG3	1:10:A:PHE:HB2	16	0.14
(1,65)	1:10:A:PHE:HE1	2:16:B:PRO:HD2	4	0.14
(1,65)	1:10:A:PHE:HE2	2:16:B:PRO:HD2	4	0.14
(1,50)	1:6:A:VAL:HG21	2:20:B:TYR:HE1	11	0.14
(1,50)	1:6:A:VAL:HG21	2:20:B:TYR:HE2	11	0.14
(1,50)	1:6:A:VAL:HG22	2:20:B:TYR:HE1	11	0.14
(1,50)	1:6:A:VAL:HG22	2:20:B:TYR:HE2	11	0.14
(1,50)	1:6:A:VAL:HG23	2:20:B:TYR:HE1	11	0.14
(1,50)	1:6:A:VAL:HG23	2:20:B:TYR:HE2	11	0.14
(1,43)	1:7:A:PRO:HG2	2:19:B:PHE:HD1	4	0.14
(1,43)	1:7:A:PRO:HG2	2:19:B:PHE:HD2	4	0.14
(1,43)	1:7:A:PRO:HG3	2:19:B:PHE:HD1	4	0.14

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,43)	1:7:A:PRO:HG3	2:19:B:PHE:HD2	4	0.14
(1,41)	1:7:A:PRO:HG2	1:10:A:PHE:HB2	16	0.14
(1,41)	1:7:A:PRO:HG3	1:10:A:PHE:HB2	16	0.14
(1,30)	2:13:B:CYS:HB3	2:14:B:GLY:HA3	13	0.14
(1,25)	1:10:A:PHE:HE1	2:16:B:PRO:HD2	15	0.14
(1,25)	1:10:A:PHE:HE2	2:16:B:PRO:HD2	15	0.14
(1,1303)	2:41:D:TYR:H	2:41:D:TYR:HD1	18	0.13
(1,1303)	2:41:D:TYR:H	2:41:D:TYR:HD2	18	0.13
(1,1303)	2:41:D:TYR:H	2:41:D:TYR:HD1	19	0.13
(1,1303)	2:41:D:TYR:H	2:41:D:TYR:HD2	19	0.13
(1,1252)	1:28:C:PRO:HG2	1:31:C:PHE:HE1	1	0.13
(1,1252)	1:28:C:PRO:HG2	1:31:C:PHE:HE2	1	0.13
(1,1252)	1:28:C:PRO:HG3	1:31:C:PHE:HE1	1	0.13
(1,1252)	1:28:C:PRO:HG3	1:31:C:PHE:HE2	1	0.13
(1,1242)	1:33:C:ARG:H	1:33:C:ARG:HB2	16	0.13
(1,1242)	1:33:C:ARG:H	1:33:C:ARG:HB3	16	0.13
(1,1218)	1:30:C:TYR:HA	1:30:C:TYR:HE1	14	0.13
(1,1218)	1:30:C:TYR:HA	1:30:C:TYR:HE2	14	0.13
(1,1167)	1:24:C:ALA:H	1:25:C:GLY:HA3	13	0.13
(1,1149)	1:32:C:VAL:HB	1:33:C:ARG:H	2	0.13
(1,1149)	1:32:C:VAL:HB	1:33:C:ARG:H	12	0.13
(1,1108)	2:36:D:THR:HG21	2:37:D:PRO:HD3	11	0.13
(1,1108)	2:36:D:THR:HG22	2:37:D:PRO:HD3	11	0.13
(1,1108)	2:36:D:THR:HG23	2:37:D:PRO:HD3	11	0.13
(1,1080)	1:25:C:GLY:H	2:41:D:TYR:H	3	0.13
(1,1078)	1:25:C:GLY:H	2:41:D:TYR:H	13	0.13
(1,1066)	1:23:C:GLY:HA3	1:24:C:ALA:HB1	16	0.13
(1,1066)	1:23:C:GLY:HA3	1:24:C:ALA:HB2	16	0.13
(1,1066)	1:23:C:GLY:HA3	1:24:C:ALA:HB3	16	0.13
(1,1054)	1:28:C:PRO:HA	2:40:D:PHE:HD1	8	0.13
(1,1054)	1:28:C:PRO:HA	2:40:D:PHE:HD2	8	0.13
(1,1026)	1:27:C:VAL:HG11	1:28:C:PRO:HD3	12	0.13
(1,1026)	1:27:C:VAL:HG12	1:28:C:PRO:HD3	12	0.13
(1,1026)	1:27:C:VAL:HG13	1:28:C:PRO:HD3	12	0.13
(1,1000)	1:31:C:PHE:HE1	2:37:D:PRO:HD2	8	0.13
(1,1000)	1:31:C:PHE:HE2	2:37:D:PRO:HD2	8	0.13
(1,984)	2:41:D:TYR:H	2:41:D:TYR:HD1	18	0.13
(1,984)	2:41:D:TYR:H	2:41:D:TYR:HD2	18	0.13
(1,984)	2:41:D:TYR:H	2:41:D:TYR:HD1	19	0.13
(1,984)	2:41:D:TYR:H	2:41:D:TYR:HD2	19	0.13
(1,933)	1:28:C:PRO:HG2	1:31:C:PHE:HE1	1	0.13
(1,933)	1:28:C:PRO:HG2	1:31:C:PHE:HE2	1	0.13

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,933)	1:28:C:PRO:HG3	1:31:C:PHE:HE1	1	0.13
(1,933)	1:28:C:PRO:HG3	1:31:C:PHE:HE2	1	0.13
(1,923)	1:33:C:ARG:H	1:33:C:ARG:HB2	16	0.13
(1,923)	1:33:C:ARG:H	1:33:C:ARG:HB3	16	0.13
(1,899)	1:30:C:TYR:HA	1:30:C:TYR:HE1	14	0.13
(1,899)	1:30:C:TYR:HA	1:30:C:TYR:HE2	14	0.13
(1,848)	1:24:C:ALA:H	1:25:C:GLY:HA3	13	0.13
(1,830)	1:32:C:VAL:HB	1:33:C:ARG:H	2	0.13
(1,830)	1:32:C:VAL:HB	1:33:C:ARG:H	12	0.13
(1,789)	2:36:D:THR:HG21	2:37:D:PRO:HD3	11	0.13
(1,789)	2:36:D:THR:HG22	2:37:D:PRO:HD3	11	0.13
(1,789)	2:36:D:THR:HG23	2:37:D:PRO:HD3	11	0.13
(1,761)	1:25:C:GLY:H	2:41:D:TYR:H	3	0.13
(1,759)	1:25:C:GLY:H	2:41:D:TYR:H	13	0.13
(1,747)	1:23:C:GLY:HA3	1:24:C:ALA:HB1	16	0.13
(1,747)	1:23:C:GLY:HA3	1:24:C:ALA:HB2	16	0.13
(1,747)	1:23:C:GLY:HA3	1:24:C:ALA:HB3	16	0.13
(1,735)	1:28:C:PRO:HA	2:40:D:PHE:HD1	8	0.13
(1,735)	1:28:C:PRO:HA	2:40:D:PHE:HD2	8	0.13
(1,707)	1:27:C:VAL:HG11	1:28:C:PRO:HD3	12	0.13
(1,707)	1:27:C:VAL:HG12	1:28:C:PRO:HD3	12	0.13
(1,707)	1:27:C:VAL:HG13	1:28:C:PRO:HD3	12	0.13
(1,681)	1:31:C:PHE:HE1	2:37:D:PRO:HD2	8	0.13
(1,681)	1:31:C:PHE:HE2	2:37:D:PRO:HD2	8	0.13
(1,596)	1:7:A:PRO:HG2	1:10:A:PHE:HE1	7	0.13
(1,596)	1:7:A:PRO:HG2	1:10:A:PHE:HE2	7	0.13
(1,596)	1:7:A:PRO:HG3	1:10:A:PHE:HE1	7	0.13
(1,596)	1:7:A:PRO:HG3	1:10:A:PHE:HE2	7	0.13
(1,568)	1:10:A:PHE:H	1:10:A:PHE:HD1	15	0.13
(1,568)	1:10:A:PHE:H	1:10:A:PHE:HD2	15	0.13
(1,565)	1:10:A:PHE:HA	1:11:A:VAL:H	9	0.13
(1,530)	1:5:A:HIS:HA	1:5:A:HIS:HD2	7	0.13
(1,503)	1:7:A:PRO:HB3	1:10:A:PHE:HE1	2	0.13
(1,503)	1:7:A:PRO:HB3	1:10:A:PHE:HE2	2	0.13
(1,499)	1:6:A:VAL:HG11	1:7:A:PRO:HD3	5	0.13
(1,499)	1:6:A:VAL:HG12	1:7:A:PRO:HD3	5	0.13
(1,499)	1:6:A:VAL:HG13	1:7:A:PRO:HD3	5	0.13
(1,474)	1:6:A:VAL:H	2:19:B:PHE:HB2	6	0.13
(1,468)	1:11:A:VAL:HG11	1:12:A:ARG:H	2	0.13
(1,468)	1:11:A:VAL:HG12	1:12:A:ARG:H	2	0.13
(1,468)	1:11:A:VAL:HG13	1:12:A:ARG:H	2	0.13
(1,468)	1:11:A:VAL:HG21	1:12:A:ARG:H	2	0.13

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,468)	1:11:A:VAL:HG22	1:12:A:ARG:H	2	0.13
(1,468)	1:11:A:VAL:HG23	1:12:A:ARG:H	2	0.13
(1,451)	2:15:B:THR:HG21	2:16:B:PRO:HB3	19	0.13
(1,451)	2:15:B:THR:HG22	2:16:B:PRO:HB3	19	0.13
(1,451)	2:15:B:THR:HG23	2:16:B:PRO:HB3	19	0.13
(1,435)	1:3:A:ALA:H	1:4:A:GLY:H	15	0.13
(1,424)	1:4:A:GLY:H	2:20:B:TYR:H	10	0.13
(1,398)	1:7:A:PRO:HA	2:19:B:PHE:HD1	14	0.13
(1,398)	1:7:A:PRO:HA	2:19:B:PHE:HD2	14	0.13
(1,388)	1:11:A:VAL:HG11	1:12:A:ARG:HG2	4	0.13
(1,388)	1:11:A:VAL:HG11	1:12:A:ARG:HG3	4	0.13
(1,388)	1:11:A:VAL:HG12	1:12:A:ARG:HG2	4	0.13
(1,388)	1:11:A:VAL:HG12	1:12:A:ARG:HG3	4	0.13
(1,388)	1:11:A:VAL:HG13	1:12:A:ARG:HG2	4	0.13
(1,388)	1:11:A:VAL:HG13	1:12:A:ARG:HG3	4	0.13
(1,388)	1:11:A:VAL:HG21	1:12:A:ARG:HG2	4	0.13
(1,388)	1:11:A:VAL:HG21	1:12:A:ARG:HG3	4	0.13
(1,388)	1:11:A:VAL:HG22	1:12:A:ARG:HG2	4	0.13
(1,388)	1:11:A:VAL:HG22	1:12:A:ARG:HG3	4	0.13
(1,388)	1:11:A:VAL:HG23	1:12:A:ARG:HG2	4	0.13
(1,388)	1:11:A:VAL:HG23	1:12:A:ARG:HG3	4	0.13
(1,388)	1:11:A:VAL:HG11	1:12:A:ARG:HG2	12	0.13
(1,388)	1:11:A:VAL:HG11	1:12:A:ARG:HG3	12	0.13
(1,388)	1:11:A:VAL:HG12	1:12:A:ARG:HG2	12	0.13
(1,388)	1:11:A:VAL:HG12	1:12:A:ARG:HG3	12	0.13
(1,388)	1:11:A:VAL:HG13	1:12:A:ARG:HG2	12	0.13
(1,388)	1:11:A:VAL:HG13	1:12:A:ARG:HG3	12	0.13
(1,388)	1:11:A:VAL:HG21	1:12:A:ARG:HG2	12	0.13
(1,388)	1:11:A:VAL:HG21	1:12:A:ARG:HG3	12	0.13
(1,388)	1:11:A:VAL:HG22	1:12:A:ARG:HG2	12	0.13
(1,388)	1:11:A:VAL:HG22	1:12:A:ARG:HG3	12	0.13
(1,388)	1:11:A:VAL:HG23	1:12:A:ARG:HG2	12	0.13
(1,388)	1:11:A:VAL:HG23	1:12:A:ARG:HG3	12	0.13
(1,369)	1:6:A:VAL:HG21	2:20:B:TYR:HE1	5	0.13
(1,369)	1:6:A:VAL:HG21	2:20:B:TYR:HE2	5	0.13
(1,369)	1:6:A:VAL:HG22	2:20:B:TYR:HE1	5	0.13
(1,369)	1:6:A:VAL:HG22	2:20:B:TYR:HE2	5	0.13
(1,369)	1:6:A:VAL:HG23	2:20:B:TYR:HE1	5	0.13
(1,369)	1:6:A:VAL:HG23	2:20:B:TYR:HE2	5	0.13
(1,277)	1:7:A:PRO:HG2	1:10:A:PHE:HE1	7	0.13
(1,277)	1:7:A:PRO:HG2	1:10:A:PHE:HE2	7	0.13
(1,277)	1:7:A:PRO:HG3	1:10:A:PHE:HE1	7	0.13

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,277)	1:7:A:PRO:HG3	1:10:A:PHE:HE2	7	0.13
(1,249)	1:10:A:PHE:H	1:10:A:PHE:HD1	15	0.13
(1,249)	1:10:A:PHE:H	1:10:A:PHE:HD2	15	0.13
(1,246)	1:10:A:PHE:HA	1:11:A:VAL:H	9	0.13
(1,211)	1:5:A:HIS:HA	1:5:A:HIS:HD2	7	0.13
(1,184)	1:7:A:PRO:HB3	1:10:A:PHE:HE1	2	0.13
(1,184)	1:7:A:PRO:HB3	1:10:A:PHE:HE2	2	0.13
(1,180)	1:6:A:VAL:HG11	1:7:A:PRO:HD3	5	0.13
(1,180)	1:6:A:VAL:HG12	1:7:A:PRO:HD3	5	0.13
(1,180)	1:6:A:VAL:HG13	1:7:A:PRO:HD3	5	0.13
(1,155)	1:6:A:VAL:H	2:19:B:PHE:HB2	6	0.13
(1,149)	1:11:A:VAL:HG11	1:12:A:ARG:H	2	0.13
(1,149)	1:11:A:VAL:HG12	1:12:A:ARG:H	2	0.13
(1,149)	1:11:A:VAL:HG13	1:12:A:ARG:H	2	0.13
(1,149)	1:11:A:VAL:HG21	1:12:A:ARG:H	2	0.13
(1,149)	1:11:A:VAL:HG22	1:12:A:ARG:H	2	0.13
(1,149)	1:11:A:VAL:HG23	1:12:A:ARG:H	2	0.13
(1,132)	2:15:B:THR:HG21	2:16:B:PRO:HB3	19	0.13
(1,132)	2:15:B:THR:HG22	2:16:B:PRO:HB3	19	0.13
(1,132)	2:15:B:THR:HG23	2:16:B:PRO:HB3	19	0.13
(1,116)	1:3:A:ALA:H	1:4:A:GLY:H	15	0.13
(1,105)	1:4:A:GLY:H	2:20:B:TYR:H	10	0.13
(1,79)	1:7:A:PRO:HA	2:19:B:PHE:HD1	14	0.13
(1,79)	1:7:A:PRO:HA	2:19:B:PHE:HD2	14	0.13
(1,69)	1:11:A:VAL:HG11	1:12:A:ARG:HG2	4	0.13
(1,69)	1:11:A:VAL:HG11	1:12:A:ARG:HG3	4	0.13
(1,69)	1:11:A:VAL:HG12	1:12:A:ARG:HG2	4	0.13
(1,69)	1:11:A:VAL:HG12	1:12:A:ARG:HG3	4	0.13
(1,69)	1:11:A:VAL:HG13	1:12:A:ARG:HG2	4	0.13
(1,69)	1:11:A:VAL:HG13	1:12:A:ARG:HG3	4	0.13
(1,69)	1:11:A:VAL:HG21	1:12:A:ARG:HG2	4	0.13
(1,69)	1:11:A:VAL:HG21	1:12:A:ARG:HG3	4	0.13
(1,69)	1:11:A:VAL:HG22	1:12:A:ARG:HG2	4	0.13
(1,69)	1:11:A:VAL:HG22	1:12:A:ARG:HG3	4	0.13
(1,69)	1:11:A:VAL:HG23	1:12:A:ARG:HG2	4	0.13
(1,69)	1:11:A:VAL:HG23	1:12:A:ARG:HG3	4	0.13
(1,69)	1:11:A:VAL:HG11	1:12:A:ARG:HG2	12	0.13
(1,69)	1:11:A:VAL:HG11	1:12:A:ARG:HG3	12	0.13
(1,69)	1:11:A:VAL:HG12	1:12:A:ARG:HG2	12	0.13
(1,69)	1:11:A:VAL:HG12	1:12:A:ARG:HG3	12	0.13
(1,69)	1:11:A:VAL:HG13	1:12:A:ARG:HG2	12	0.13
(1,69)	1:11:A:VAL:HG13	1:12:A:ARG:HG3	12	0.13

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,69)	1:11:A:VAL:HG21	1:12:A:ARG:HG2	12	0.13
(1,69)	1:11:A:VAL:HG21	1:12:A:ARG:HG3	12	0.13
(1,69)	1:11:A:VAL:HG22	1:12:A:ARG:HG2	12	0.13
(1,69)	1:11:A:VAL:HG22	1:12:A:ARG:HG3	12	0.13
(1,69)	1:11:A:VAL:HG23	1:12:A:ARG:HG2	12	0.13
(1,69)	1:11:A:VAL:HG23	1:12:A:ARG:HG3	12	0.13
(1,50)	1:6:A:VAL:HG21	2:20:B:TYR:HE1	5	0.13
(1,50)	1:6:A:VAL:HG21	2:20:B:TYR:HE2	5	0.13
(1,50)	1:6:A:VAL:HG22	2:20:B:TYR:HE1	5	0.13
(1,50)	1:6:A:VAL:HG22	2:20:B:TYR:HE2	5	0.13
(1,50)	1:6:A:VAL:HG23	2:20:B:TYR:HE1	5	0.13
(1,50)	1:6:A:VAL:HG23	2:20:B:TYR:HE2	5	0.13
(1,1249)	1:33:C:ARG:HB2	1:33:C:ARG:HE	6	0.12
(1,1231)	1:31:C:PHE:HE1	2:36:D:THR:HA	1	0.12
(1,1231)	1:31:C:PHE:HE2	2:36:D:THR:HA	1	0.12
(1,1228)	1:31:C:PHE:HA	1:31:C:PHE:HD1	8	0.12
(1,1228)	1:31:C:PHE:HA	1:31:C:PHE:HD2	8	0.12
(1,1221)	1:31:C:PHE:HA	1:32:C:VAL:H	15	0.12
(1,1186)	1:26:C:HIS:HA	1:26:C:HIS:HD2	7	0.12
(1,1186)	1:26:C:HIS:HA	1:26:C:HIS:HD2	9	0.12
(1,1167)	1:24:C:ALA:H	1:25:C:GLY:HA3	3	0.12
(1,1162)	1:27:C:VAL:H	2:40:D:PHE:HA	1	0.12
(1,1143)	1:27:C:VAL:H	2:41:D:TYR:HD1	19	0.12
(1,1143)	1:27:C:VAL:H	2:41:D:TYR:HD2	19	0.12
(1,1133)	2:34:D:CYS:HA	2:35:D:GLY:H	5	0.12
(1,1130)	1:27:C:VAL:H	2:40:D:PHE:HB2	5	0.12
(1,1091)	1:24:C:ALA:H	1:25:C:GLY:H	13	0.12
(1,1089)	1:29:C:GLU:HB2	2:41:D:TYR:HB2	19	0.12
(1,1089)	1:29:C:GLU:HB2	2:41:D:TYR:HB3	19	0.12
(1,1066)	1:23:C:GLY:HA3	1:24:C:ALA:HB1	15	0.12
(1,1066)	1:23:C:GLY:HA3	1:24:C:ALA:HB2	15	0.12
(1,1066)	1:23:C:GLY:HA3	1:24:C:ALA:HB3	15	0.12
(1,1045)	1:27:C:VAL:HB	2:41:D:TYR:HD1	20	0.12
(1,1045)	1:27:C:VAL:HB	2:41:D:TYR:HD2	20	0.12
(1,1040)	1:31:C:PHE:HE1	2:37:D:PRO:HD2	6	0.12
(1,1040)	1:31:C:PHE:HE2	2:37:D:PRO:HD2	6	0.12
(1,1035)	1:22:C:GLY:H	2:41:D:TYR:HE1	7	0.12
(1,1035)	1:22:C:GLY:H	2:41:D:TYR:HE2	7	0.12
(1,1003)	1:28:C:PRO:HD2	2:41:D:TYR:HD1	4	0.12
(1,1003)	1:28:C:PRO:HD2	2:41:D:TYR:HD2	4	0.12
(1,930)	1:33:C:ARG:HB2	1:33:C:ARG:HE	6	0.12
(1,912)	1:31:C:PHE:HE1	2:36:D:THR:HA	1	0.12

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,912)	1:31:C:PHE:HE2	2:36:D:THR:HA	1	0.12
(1,909)	1:31:C:PHE:HA	1:31:C:PHE:HD1	8	0.12
(1,909)	1:31:C:PHE:HA	1:31:C:PHE:HD2	8	0.12
(1,902)	1:31:C:PHE:HA	1:32:C:VAL:H	15	0.12
(1,867)	1:26:C:HIS:HA	1:26:C:HIS:HD2	7	0.12
(1,867)	1:26:C:HIS:HA	1:26:C:HIS:HD2	9	0.12
(1,848)	1:24:C:ALA:H	1:25:C:GLY:HA3	3	0.12
(1,843)	1:27:C:VAL:H	2:40:D:PHE:HA	1	0.12
(1,824)	1:27:C:VAL:H	2:41:D:TYR:HD1	19	0.12
(1,824)	1:27:C:VAL:H	2:41:D:TYR:HD2	19	0.12
(1,814)	2:34:D:CYS:HA	2:35:D:GLY:H	5	0.12
(1,811)	1:27:C:VAL:H	2:40:D:PHE:HB2	5	0.12
(1,772)	1:24:C:ALA:H	1:25:C:GLY:H	13	0.12
(1,770)	1:29:C:GLU:HB2	2:41:D:TYR:HB2	19	0.12
(1,770)	1:29:C:GLU:HB2	2:41:D:TYR:HB3	19	0.12
(1,747)	1:23:C:GLY:HA3	1:24:C:ALA:HB1	15	0.12
(1,747)	1:23:C:GLY:HA3	1:24:C:ALA:HB2	15	0.12
(1,747)	1:23:C:GLY:HA3	1:24:C:ALA:HB3	15	0.12
(1,726)	1:27:C:VAL:HB	2:41:D:TYR:HD1	20	0.12
(1,726)	1:27:C:VAL:HB	2:41:D:TYR:HD2	20	0.12
(1,721)	1:31:C:PHE:HE1	2:37:D:PRO:HD2	6	0.12
(1,721)	1:31:C:PHE:HE2	2:37:D:PRO:HD2	6	0.12
(1,716)	1:22:C:GLY:H	2:41:D:TYR:HE1	7	0.12
(1,716)	1:22:C:GLY:H	2:41:D:TYR:HE2	7	0.12
(1,684)	1:28:C:PRO:HD2	2:41:D:TYR:HD1	4	0.12
(1,684)	1:28:C:PRO:HD2	2:41:D:TYR:HD2	4	0.12
(1,673)	2:17:B:ILE:HG13	2:40:D:PHE:HB2	1	0.12
(1,562)	1:9:A:TYR:HA	1:9:A:TYR:HE1	20	0.12
(1,562)	1:9:A:TYR:HA	1:9:A:TYR:HE2	20	0.12
(1,559)	1:9:A:TYR:HA	1:9:A:TYR:HD1	17	0.12
(1,559)	1:9:A:TYR:HA	1:9:A:TYR:HD2	17	0.12
(1,530)	1:5:A:HIS:HA	1:5:A:HIS:HD2	12	0.12
(1,492)	1:7:A:PRO:HB2	1:10:A:PHE:HD1	18	0.12
(1,492)	1:7:A:PRO:HB2	1:10:A:PHE:HD2	18	0.12
(1,463)	1:5:A:HIS:HB2	1:6:A:VAL:HG11	6	0.12
(1,463)	1:5:A:HIS:HB2	1:6:A:VAL:HG12	6	0.12
(1,463)	1:5:A:HIS:HB2	1:6:A:VAL:HG13	6	0.12
(1,415)	1:6:A:VAL:HG11	2:20:B:TYR:HE1	20	0.12
(1,415)	1:6:A:VAL:HG11	2:20:B:TYR:HE2	20	0.12
(1,415)	1:6:A:VAL:HG12	2:20:B:TYR:HE1	20	0.12
(1,415)	1:6:A:VAL:HG12	2:20:B:TYR:HE2	20	0.12
(1,415)	1:6:A:VAL:HG13	2:20:B:TYR:HE1	20	0.12

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,415)	1:6:A:VAL:HG13	2:20:B:TYR:HE2	20	0.12
(1,410)	1:2:A:GLY:HA3	1:3:A:ALA:HB1	9	0.12
(1,410)	1:2:A:GLY:HA3	1:3:A:ALA:HB2	9	0.12
(1,410)	1:2:A:GLY:HA3	1:3:A:ALA:HB3	9	0.12
(1,384)	1:10:A:PHE:HE1	2:16:B:PRO:HD2	2	0.12
(1,384)	1:10:A:PHE:HE2	2:16:B:PRO:HD2	2	0.12
(1,343)	1:7:A:PRO:HD3	2:19:B:PHE:HD1	5	0.12
(1,343)	1:7:A:PRO:HD3	2:19:B:PHE:HD2	5	0.12
(1,243)	1:9:A:TYR:HA	1:9:A:TYR:HE1	20	0.12
(1,243)	1:9:A:TYR:HA	1:9:A:TYR:HE2	20	0.12
(1,240)	1:9:A:TYR:HA	1:9:A:TYR:HD1	17	0.12
(1,240)	1:9:A:TYR:HA	1:9:A:TYR:HD2	17	0.12
(1,211)	1:5:A:HIS:HA	1:5:A:HIS:HD2	12	0.12
(1,173)	1:7:A:PRO:HB2	1:10:A:PHE:HD1	18	0.12
(1,173)	1:7:A:PRO:HB2	1:10:A:PHE:HD2	18	0.12
(1,144)	1:5:A:HIS:HB2	1:6:A:VAL:HG11	6	0.12
(1,144)	1:5:A:HIS:HB2	1:6:A:VAL:HG12	6	0.12
(1,144)	1:5:A:HIS:HB2	1:6:A:VAL:HG13	6	0.12
(1,96)	1:6:A:VAL:HG11	2:20:B:TYR:HE1	20	0.12
(1,96)	1:6:A:VAL:HG11	2:20:B:TYR:HE2	20	0.12
(1,96)	1:6:A:VAL:HG12	2:20:B:TYR:HE1	20	0.12
(1,96)	1:6:A:VAL:HG12	2:20:B:TYR:HE2	20	0.12
(1,96)	1:6:A:VAL:HG13	2:20:B:TYR:HE1	20	0.12
(1,96)	1:6:A:VAL:HG13	2:20:B:TYR:HE2	20	0.12
(1,91)	1:2:A:GLY:HA3	1:3:A:ALA:HB1	9	0.12
(1,91)	1:2:A:GLY:HA3	1:3:A:ALA:HB2	9	0.12
(1,91)	1:2:A:GLY:HA3	1:3:A:ALA:HB3	9	0.12
(1,65)	1:10:A:PHE:HE1	2:16:B:PRO:HD2	2	0.12
(1,65)	1:10:A:PHE:HE2	2:16:B:PRO:HD2	2	0.12
(1,24)	1:7:A:PRO:HD3	2:19:B:PHE:HD1	5	0.12
(1,24)	1:7:A:PRO:HD3	2:19:B:PHE:HD2	5	0.12
(1,8)	2:17:B:ILE:HG13	2:40:D:PHE:HB2	1	0.12
(1,1242)	1:33:C:ARG:H	1:33:C:ARG:HB2	6	0.11
(1,1242)	1:33:C:ARG:H	1:33:C:ARG:HB3	6	0.11
(1,1186)	1:26:C:HIS:HA	1:26:C:HIS:HD2	12	0.11
(1,1156)	2:34:D:CYS:HB3	2:35:D:GLY:H	15	0.11
(1,1155)	1:27:C:VAL:HG11	1:28:C:PRO:HD3	3	0.11
(1,1155)	1:27:C:VAL:HG12	1:28:C:PRO:HD3	3	0.11
(1,1155)	1:27:C:VAL:HG13	1:28:C:PRO:HD3	3	0.11
(1,1148)	1:28:C:PRO:HB2	1:31:C:PHE:HD1	3	0.11
(1,1148)	1:28:C:PRO:HB2	1:31:C:PHE:HD2	3	0.11
(1,1108)	2:36:D:THR:HG21	2:37:D:PRO:HD3	4	0.11

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,1108)	2:36:D:THR:HG22	2:37:D:PRO:HD3	4	0.11
(1,1108)	2:36:D:THR:HG23	2:37:D:PRO:HD3	4	0.11
(1,1094)	1:29:C:GLU:HG2	2:40:D:PHE:HA	1	0.11
(1,1091)	1:24:C:ALA:H	1:25:C:GLY:H	5	0.11
(1,1042)	1:25:C:GLY:H	2:40:D:PHE:HD1	3	0.11
(1,1042)	1:25:C:GLY:H	2:40:D:PHE:HD2	3	0.11
(1,1018)	1:28:C:PRO:HG2	2:40:D:PHE:HD1	7	0.11
(1,1018)	1:28:C:PRO:HG2	2:40:D:PHE:HD2	7	0.11
(1,1018)	1:28:C:PRO:HG3	2:40:D:PHE:HD1	7	0.11
(1,1018)	1:28:C:PRO:HG3	2:40:D:PHE:HD2	7	0.11
(1,923)	1:33:C:ARG:H	1:33:C:ARG:HB2	6	0.11
(1,923)	1:33:C:ARG:H	1:33:C:ARG:HB3	6	0.11
(1,867)	1:26:C:HIS:HA	1:26:C:HIS:HD2	12	0.11
(1,837)	2:34:D:CYS:HB3	2:35:D:GLY:H	15	0.11
(1,836)	1:27:C:VAL:HG11	1:28:C:PRO:HD3	3	0.11
(1,836)	1:27:C:VAL:HG12	1:28:C:PRO:HD3	3	0.11
(1,836)	1:27:C:VAL:HG13	1:28:C:PRO:HD3	3	0.11
(1,829)	1:28:C:PRO:HB2	1:31:C:PHE:HD1	3	0.11
(1,829)	1:28:C:PRO:HB2	1:31:C:PHE:HD2	3	0.11
(1,789)	2:36:D:THR:HG21	2:37:D:PRO:HD3	4	0.11
(1,789)	2:36:D:THR:HG22	2:37:D:PRO:HD3	4	0.11
(1,789)	2:36:D:THR:HG23	2:37:D:PRO:HD3	4	0.11
(1,775)	1:29:C:GLU:HG2	2:40:D:PHE:HA	1	0.11
(1,772)	1:24:C:ALA:H	1:25:C:GLY:H	5	0.11
(1,723)	1:25:C:GLY:H	2:40:D:PHE:HD1	3	0.11
(1,723)	1:25:C:GLY:H	2:40:D:PHE:HD2	3	0.11
(1,699)	1:28:C:PRO:HG2	2:40:D:PHE:HD1	7	0.11
(1,699)	1:28:C:PRO:HG2	2:40:D:PHE:HD2	7	0.11
(1,699)	1:28:C:PRO:HG3	2:40:D:PHE:HD1	7	0.11
(1,699)	1:28:C:PRO:HG3	2:40:D:PHE:HD2	7	0.11
(1,586)	1:12:A:ARG:H	1:12:A:ARG:HB2	1	0.11
(1,586)	1:12:A:ARG:H	1:12:A:ARG:HB3	1	0.11
(1,567)	1:10:A:PHE:HB3	1:11:A:VAL:H	11	0.11
(1,501)	1:4:A:GLY:H	1:6:A:VAL:H	4	0.11
(1,500)	2:13:B:CYS:HB3	2:14:B:GLY:H	14	0.11
(1,493)	1:11:A:VAL:HB	1:12:A:ARG:H	18	0.11
(1,465)	1:5:A:HIS:HD2	1:6:A:VAL:HG11	5	0.11
(1,465)	1:5:A:HIS:HD2	1:6:A:VAL:HG12	5	0.11
(1,465)	1:5:A:HIS:HD2	1:6:A:VAL:HG13	5	0.11
(1,438)	1:8:A:GLU:HG2	2:19:B:PHE:HA	7	0.11
(1,405)	1:1:A:GLY:H	2:18:B:SER:HB2	16	0.11
(1,389)	1:6:A:VAL:HB	2:20:B:TYR:HD1	1	0.11

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,389)	1:6:A:VAL:HB	2:20:B:TYR:HD2	1	0.11
(1,379)	1:1:A:GLY:H	2:20:B:TYR:HE1	13	0.11
(1,379)	1:1:A:GLY:H	2:20:B:TYR:HE2	13	0.11
(1,370)	1:6:A:VAL:HG11	1:7:A:PRO:HD3	1	0.11
(1,370)	1:6:A:VAL:HG12	1:7:A:PRO:HD3	1	0.11
(1,370)	1:6:A:VAL:HG13	1:7:A:PRO:HD3	1	0.11
(1,267)	1:12:A:ARG:H	1:12:A:ARG:HB2	1	0.11
(1,267)	1:12:A:ARG:H	1:12:A:ARG:HB3	1	0.11
(1,248)	1:10:A:PHE:HB3	1:11:A:VAL:H	11	0.11
(1,182)	1:4:A:GLY:H	1:6:A:VAL:H	4	0.11
(1,181)	2:13:B:CYS:HB3	2:14:B:GLY:H	14	0.11
(1,174)	1:11:A:VAL:HB	1:12:A:ARG:H	18	0.11
(1,146)	1:5:A:HIS:HD2	1:6:A:VAL:HG11	5	0.11
(1,146)	1:5:A:HIS:HD2	1:6:A:VAL:HG12	5	0.11
(1,146)	1:5:A:HIS:HD2	1:6:A:VAL:HG13	5	0.11
(1,119)	1:8:A:GLU:HG2	2:19:B:PHE:HA	7	0.11
(1,86)	1:1:A:GLY:H	2:18:B:SER:HB2	16	0.11
(1,70)	1:6:A:VAL:HB	2:20:B:TYR:HD1	1	0.11
(1,70)	1:6:A:VAL:HB	2:20:B:TYR:HD2	1	0.11
(1,60)	1:1:A:GLY:H	2:20:B:TYR:HE1	13	0.11
(1,60)	1:1:A:GLY:H	2:20:B:TYR:HE2	13	0.11
(1,51)	1:6:A:VAL:HG11	1:7:A:PRO:HD3	1	0.11
(1,51)	1:6:A:VAL:HG12	1:7:A:PRO:HD3	1	0.11
(1,51)	1:6:A:VAL:HG13	1:7:A:PRO:HD3	1	0.11
(1,1228)	1:31:C:PHE:HA	1:31:C:PHE:HD1	4	0.1
(1,1228)	1:31:C:PHE:HA	1:31:C:PHE:HD2	4	0.1
(1,1224)	1:31:C:PHE:H	1:31:C:PHE:HD1	9	0.1
(1,1224)	1:31:C:PHE:H	1:31:C:PHE:HD2	9	0.1
(1,1221)	1:31:C:PHE:HA	1:32:C:VAL:H	16	0.1
(1,1156)	2:34:D:CYS:HB3	2:35:D:GLY:H	9	0.1
(1,1074)	1:22:C:GLY:H	2:39:D:SER:HB3	4	0.1
(1,1035)	1:22:C:GLY:H	2:41:D:TYR:HE1	5	0.1
(1,1035)	1:22:C:GLY:H	2:41:D:TYR:HE2	5	0.1
(1,1004)	1:32:C:VAL:HA	1:33:C:ARG:HD2	11	0.1
(1,1004)	1:32:C:VAL:HA	1:33:C:ARG:HD3	11	0.1
(1,909)	1:31:C:PHE:HA	1:31:C:PHE:HD1	4	0.1
(1,909)	1:31:C:PHE:HA	1:31:C:PHE:HD2	4	0.1
(1,905)	1:31:C:PHE:H	1:31:C:PHE:HD1	9	0.1
(1,905)	1:31:C:PHE:H	1:31:C:PHE:HD2	9	0.1
(1,902)	1:31:C:PHE:HA	1:32:C:VAL:H	16	0.1
(1,837)	2:34:D:CYS:HB3	2:35:D:GLY:H	9	0.1
(1,755)	1:22:C:GLY:H	2:39:D:SER:HB3	4	0.1

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,716)	1:22:C:GLY:H	2:41:D:TYR:HE1	5	0.1
(1,716)	1:22:C:GLY:H	2:41:D:TYR:HE2	5	0.1
(1,685)	1:32:C:VAL:HA	1:33:C:ARG:HD2	11	0.1
(1,685)	1:32:C:VAL:HA	1:33:C:ARG:HD3	11	0.1
(1,572)	1:10:A:PHE:HA	1:10:A:PHE:HD1	7	0.1
(1,572)	1:10:A:PHE:HA	1:10:A:PHE:HD2	7	0.1
(1,565)	1:10:A:PHE:HA	1:11:A:VAL:H	7	0.1
(1,565)	1:10:A:PHE:HA	1:11:A:VAL:H	17	0.1
(1,565)	1:10:A:PHE:HA	1:11:A:VAL:H	19	0.1
(1,493)	1:11:A:VAL:HB	1:12:A:ARG:H	16	0.1
(1,253)	1:10:A:PHE:HA	1:10:A:PHE:HD1	7	0.1
(1,253)	1:10:A:PHE:HA	1:10:A:PHE:HD2	7	0.1
(1,246)	1:10:A:PHE:HA	1:11:A:VAL:H	7	0.1
(1,246)	1:10:A:PHE:HA	1:11:A:VAL:H	17	0.1
(1,246)	1:10:A:PHE:HA	1:11:A:VAL:H	19	0.1
(1,174)	1:11:A:VAL:HB	1:12:A:ARG:H	16	0.1

10 Dihedral-angle violation analysis (i)

No dihedral-angle restraints found