



Full wwPDB NMR Structure Validation Report ⓘ

Mar 5, 2026 – 03:48 PM UTC

PDB ID : 7ATY / pdb_00007aty
BMRB ID : 34570
Title : Solution NMR structure of the SH3 domain of human Caskin1
Authors : Toke, O.; Koprivanacz, K.; Radnai, L.; Mero, B.; Juhasz, T.; Liliom, K.; Buday, L.
Deposited on : 2020-11-01

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 ⓘ symbol.

The types of validation reports are described at

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

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

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

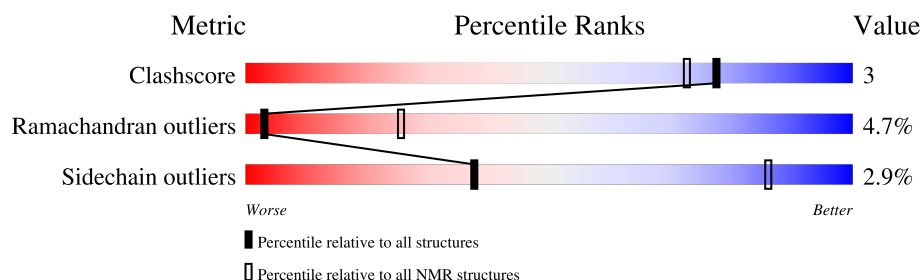
1 Overall quality at a glance

The following experimental techniques were used to determine the structure:

SOLUTION NMR

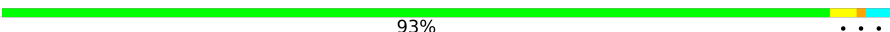
The overall completeness of chemical shifts assignment is 65%.

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 $\leq 5\%$.

Mol	Chain	Length	Quality of chain
1	A	67	 93% ...

2 Ensemble composition and analysis

This entry contains 30 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: *lowest energy*.

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:3-A:67 (65)	1.33	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 6 clusters and 4 single-model clusters were found.

Cluster number	Models
1	3, 4, 9, 10, 13, 16, 18, 20, 21, 23, 26, 27, 28
2	2, 5, 7, 11, 12
3	1, 24
4	6, 29
5	14, 15
6	19, 30
Single-model clusters	8; 17; 22; 25

3 Entry composition

There is only 1 type of molecule in this entry. The entry contains 1026 atoms, of which 503 are hydrogens and 0 are deuteriums.

- Molecule 1 is a protein called Caskin-1.

Mol	Chain	Residues	Atoms						Trace
1	A	67	Total	C	H	N	O	S	0
			1026	322	503	96	102	3	

There are 4 discrepancies between the modelled and reference sequences:

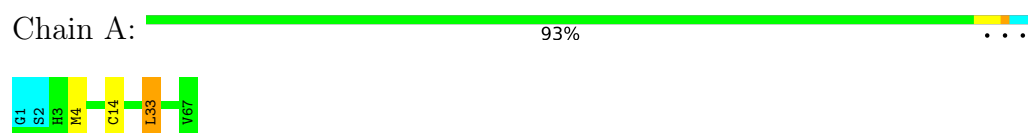
Chain	Residue	Modelled	Actual	Comment	Reference
A	1	GLY	-	expression tag	UNP Q8WXD9
A	2	SER	-	expression tag	UNP Q8WXD9
A	3	HIS	-	expression tag	UNP Q8WXD9
A	4	MET	-	expression tag	UNP Q8WXD9

4 Residue-property plots [i](#)

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: Caskin-1

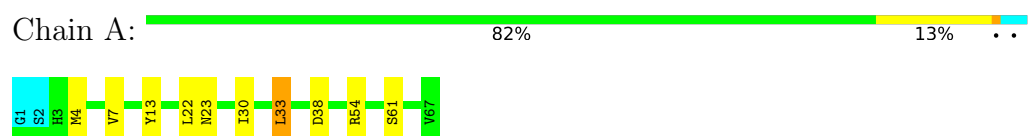


4.2 Scores per residue for each member of the ensemble

Colouring as in section [4.1](#) above.

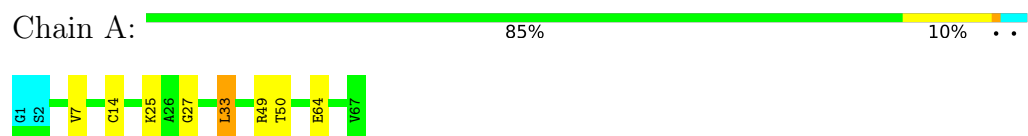
4.2.1 Score per residue for model 1

- Molecule 1: Caskin-1



4.2.2 Score per residue for model 2

- Molecule 1: Caskin-1



4.2.3 Score per residue for model 3

- Molecule 1: Caskin-1

Chain A:  84% 12% ..



4.2.4 Score per residue for model 4

- Molecule 1: Caskin-1

Chain A:  85% 9% ..



4.2.5 Score per residue for model 5

- Molecule 1: Caskin-1

Chain A:  87% 9% ..



4.2.6 Score per residue for model 6

- Molecule 1: Caskin-1

Chain A:  82% 13% ..



4.2.7 Score per residue for model 7

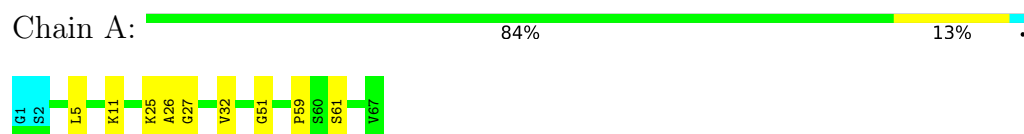
- Molecule 1: Caskin-1

Chain A:  88% 9% .



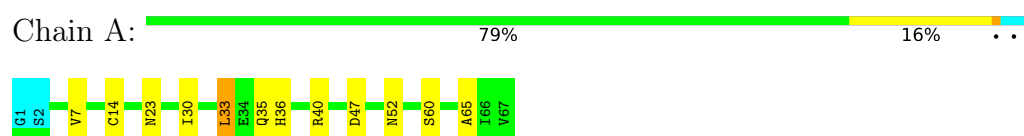
4.2.8 Score per residue for model 8

- Molecule 1: Caskin-1



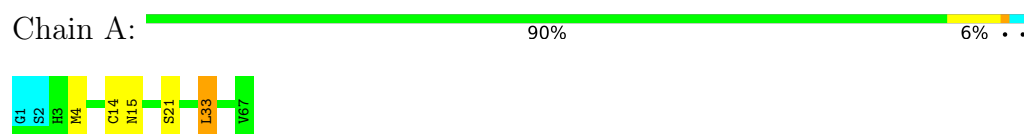
4.2.9 Score per residue for model 9

- Molecule 1: Caskin-1



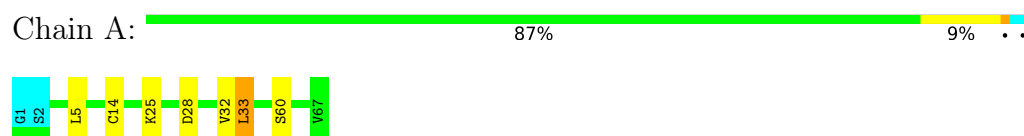
4.2.10 Score per residue for model 10

- Molecule 1: Caskin-1



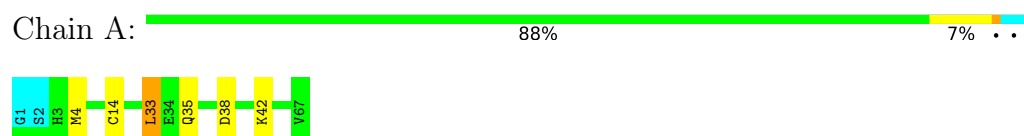
4.2.11 Score per residue for model 11

- Molecule 1: Caskin-1



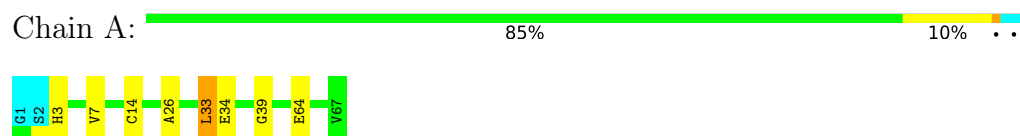
4.2.12 Score per residue for model 12

- Molecule 1: Caskin-1



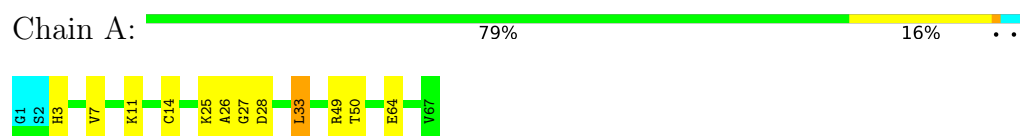
4.2.13 Score per residue for model 13 (medoid)

- Molecule 1: Caskin-1



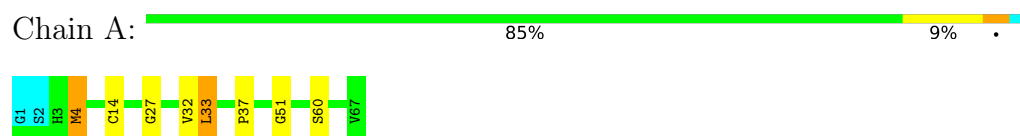
4.2.14 Score per residue for model 14

- Molecule 1: Caskin-1



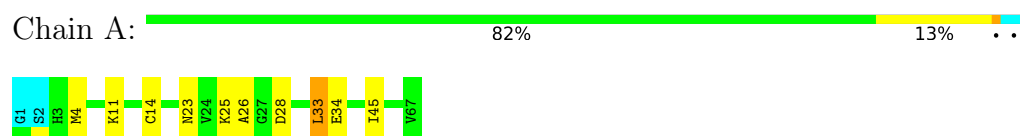
4.2.15 Score per residue for model 15

- Molecule 1: Caskin-1



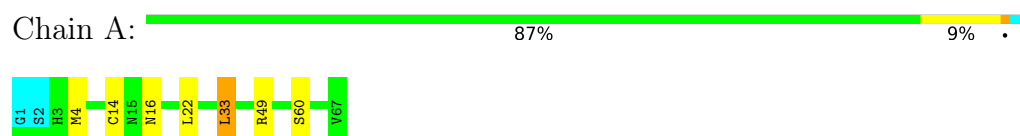
4.2.16 Score per residue for model 16

- Molecule 1: Caskin-1



4.2.17 Score per residue for model 17

- Molecule 1: Caskin-1



4.2.18 Score per residue for model 18

- Molecule 1: Caskin-1

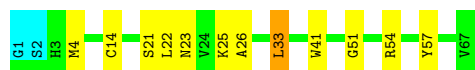
Chain A:  90% 6% ..



4.2.19 Score per residue for model 19

- Molecule 1: Caskin-1

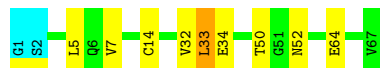
Chain A:  79% 16% ..



4.2.20 Score per residue for model 20

- Molecule 1: Caskin-1

Chain A:  84% 12% ..



4.2.21 Score per residue for model 21

- Molecule 1: Caskin-1

Chain A:  91% ..



4.2.22 Score per residue for model 22

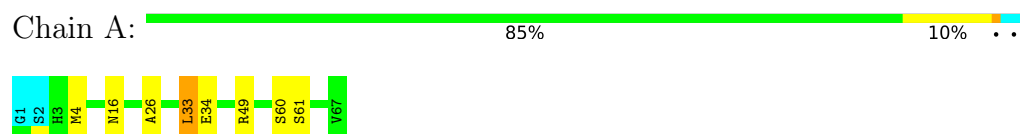
- Molecule 1: Caskin-1

Chain A:  91% ..



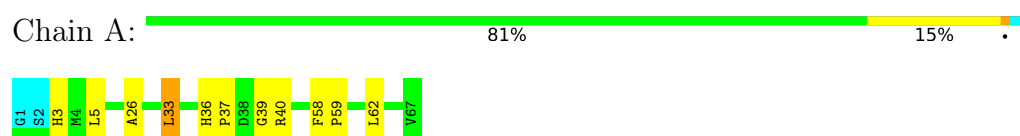
4.2.23 Score per residue for model 23

- Molecule 1: Caskin-1



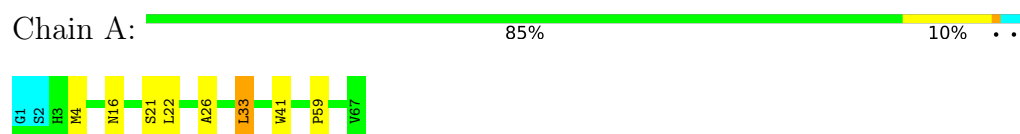
4.2.24 Score per residue for model 24

- Molecule 1: Caskin-1



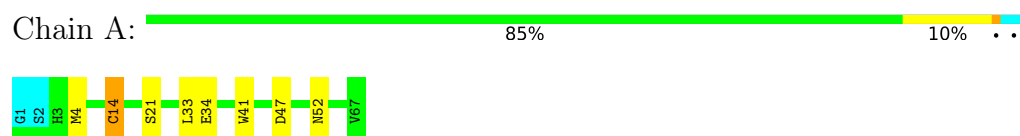
4.2.25 Score per residue for model 25

- Molecule 1: Caskin-1



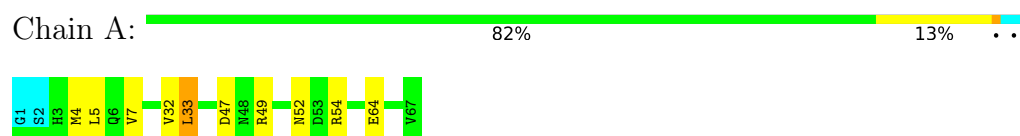
4.2.26 Score per residue for model 26

- Molecule 1: Caskin-1



4.2.27 Score per residue for model 27

- Molecule 1: Caskin-1



4.2.28 Score per residue for model 28

- Molecule 1: Caskin-1

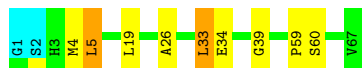
Chain A:  84% 12% ..



4.2.29 Score per residue for model 29

- Molecule 1: Caskin-1

Chain A:  84% 10% ..



4.2.30 Score per residue for model 30

- Molecule 1: Caskin-1

Chain A:  79% 16% ..



5 Refinement protocol and experimental data overview

The models were refined using the following method: *simulated annealing*.

Of the 100 calculated structures, 30 were deposited, based on the following criterion: *structures with the lowest energy*.

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

Software name	Classification	Version
ARIA	structure calculation	
ARIA	refinement	

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	570
Number of shifts mapped to atoms	570
Number of unparsed shifts	0
Number of shifts with mapping errors	0
Number of shifts with mapping warnings	0
Assignment completeness (well-defined parts)	65%

6 Model quality

6.1 Standard geometry

There are no covalent bond-length or bond-angle outliers.

There are no bond-length outliers.

There are no bond-angle outliers.

There are no chirality outliers.

There are no planarity outliers.

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	513	493	493	3±1
All	All	15390	14790	14790	89

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 3.

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

Atom-1	Atom-2	Clash(Å)	Distance(Å)	Models	
				Worst	Total
1:A:14:CYS:SG	1:A:21:SER:HB2	0.61	2.35	10	4
1:A:25:LYS:HG2	1:A:28:ASP:OD1	0.55	2.02	18	2
1:A:7:VAL:HA	1:A:64:GLU:O	0.54	2.03	3	6
1:A:54:ARG:HA	1:A:54:ARG:NE	0.54	2.18	27	1
1:A:25:LYS:HE3	1:A:28:ASP:OD1	0.52	2.04	16	1
1:A:4:MET:SD	1:A:33:LEU:HB3	0.52	2.44	22	1
1:A:11:LYS:N	1:A:26:ALA:HA	0.50	2.21	8	3
1:A:25:LYS:HG2	1:A:28:ASP:CG	0.50	2.32	11	1
1:A:39:GLY:O	1:A:59:PRO:HA	0.49	2.06	5	2
1:A:23:ASN:OD1	1:A:54:ARG:HD2	0.49	2.08	1	1
1:A:25:LYS:HE2	1:A:50:THR:O	0.48	2.08	2	1
1:A:36:HIS:CE1	1:A:39:GLY:HA2	0.47	2.45	6	2
1:A:23:ASN:N	1:A:45:ILE:HG13	0.47	2.25	16	1
1:A:35:GLN:OE1	1:A:42:LYS:HG2	0.46	2.10	12	1

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Atom-1	Atom-2	Clash(Å)	Distance(Å)	Models	
				Worst	Total
1:A:21:SER:HB3	1:A:57:TYR:O	0.45	2.10	19	1
1:A:33:LEU:HD13	1:A:33:LEU:H	0.45	1.70	20	2
1:A:11:LYS:HB3	1:A:13:TYR:CE1	0.45	2.47	22	1
1:A:7:VAL:CG2	1:A:30:ILE:HB	0.45	2.40	9	4
1:A:33:LEU:N	1:A:33:LEU:HD13	0.44	2.28	17	24
1:A:34:GLU:HG2	1:A:41:TRP:CZ2	0.44	2.46	26	1
1:A:5:LEU:HG	1:A:32:VAL:CG2	0.44	2.42	8	4
1:A:47:ASP:HB3	1:A:52:ASN:H	0.43	1.72	27	1
1:A:23:ASN:OD1	1:A:54:ARG:HG2	0.43	2.13	19	1
1:A:4:MET:HA	1:A:32:VAL:O	0.43	2.14	15	1
1:A:47:ASP:HB3	1:A:52:ASN:N	0.43	2.28	26	1
1:A:25:LYS:CE	1:A:51:GLY:HA3	0.42	2.43	8	1
1:A:5:LEU:HD13	1:A:5:LEU:N	0.42	2.28	29	1
1:A:22:LEU:H	1:A:56:GLY:HA3	0.42	1.74	30	1
1:A:47:ASP:OD2	1:A:49:ARG:HB3	0.42	2.14	27	1
1:A:13:TYR:HD2	1:A:22:LEU:O	0.42	1.97	1	1
1:A:21:SER:O	1:A:22:LEU:HB2	0.42	2.13	25	1
1:A:23:ASN:H	1:A:45:ILE:HG13	0.41	1.75	16	1
1:A:45:ILE:HB	1:A:54:ARG:HB2	0.41	1.92	30	1
1:A:47:ASP:CG	1:A:52:ASN:H	0.41	2.24	9	1
1:A:58:PHE:CD2	1:A:62:LEU:HB2	0.41	2.51	24	1
1:A:36:HIS:N	1:A:37:PRO:HD3	0.41	2.31	6	1
1:A:7:VAL:HG12	1:A:65:ALA:HA	0.41	1.93	9	1
1:A:36:HIS:HB2	1:A:40:ARG:O	0.41	2.15	9	1
1:A:33:LEU:O	1:A:41:TRP:HB3	0.41	2.16	19	2
1:A:37:PRO:HD2	1:A:40:ARG:O	0.41	2.16	24	1
1:A:33:LEU:HD13	1:A:33:LEU:N	0.41	2.30	29	1
1:A:11:LYS:C	1:A:26:ALA:HA	0.40	2.41	16	1
1:A:36:HIS:NE2	1:A:39:GLY:HA2	0.40	2.32	24	1
1:A:13:TYR:CE2	1:A:21:SER:HB3	0.40	2.51	6	1
1:A:5:LEU:N	1:A:5:LEU:HD13	0.40	2.32	28	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	64/67 (96%)	51±2 (79±3%)	10±2 (16±4%)	3±1 (5±2%)	3	25
All	All	1920/2010 (96%)	1522 (79%)	307 (16%)	91 (5%)	3	25

All 19 unique Ramachandran outliers are listed below. They are sorted by the frequency of occurrence in the ensemble.

Mol	Chain	Res	Type	Models (Total)
1	A	4	MET	17
1	A	14	CYS	16
1	A	60	SER	8
1	A	27	GLY	7
1	A	59	PRO	7
1	A	26	ALA	7
1	A	49	ARG	6
1	A	61	SER	4
1	A	51	GLY	3
1	A	3	HIS	3
1	A	38	ASP	2
1	A	52	ASN	2
1	A	22	LEU	2
1	A	50	THR	2
1	A	35	GLN	1
1	A	15	ASN	1
1	A	39	GLY	1
1	A	37	PRO	1
1	A	16	ASN	1

6.3.2 Protein sidechains ⓘ

In the following table, the Percentiles column shows the percent sidechain outliers of the chain as a percentile score with respect to all PDB entries followed by that with respect to all 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	56/57 (98%)	54±1 (97±2%)	2±1 (3±2%)	38	86
All	All	1680/1710 (98%)	1631 (97%)	49 (3%)	38	86

All 13 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	33	LEU	28
1	A	5	LEU	5
1	A	16	ASN	3
1	A	25	LYS	2
1	A	20	THR	2
1	A	23	ASN	2
1	A	28	ASP	1
1	A	34	GLU	1
1	A	40	ARG	1
1	A	50	THR	1
1	A	12	ASP	1
1	A	19	LEU	1
1	A	48	ASN	1

6.3.3 RNA [i](#)

There are no RNA molecules in this entry.

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

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

6.5 Carbohydrates [i](#)

There are no oligosaccharides in this entry.

6.6 Ligand geometry [i](#)

There are no ligands in this entry.

6.7 Other polymers [i](#)

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 65% for the well-defined parts and 64% for the entire structure.

7.1 Chemical shift list 1

File name: `working_cs.cif`

Chemical shift list name: *starch_output*

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	570
Number of shifts mapped to atoms	570
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

The following table shows the suggested chemical shift referencing corrections.

Nucleus	# values	Correction $\pm$ precision, ppm	Suggested action
$^{13}\text{C}_\alpha$	54	-0.25 ± 0.14	None needed (< 0.5 ppm)
$^{13}\text{C}_\beta$	47	0.19 ± 0.25	None needed (< 0.5 ppm)
$^{13}\text{C}'$	0	—	None (insufficient data)
^{15}N	61	-0.57 ± 0.76	None needed (imprecise)

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 65%, i.e. 565 atoms were assigned a chemical shift out of a possible 871. 0 out of 10 assigned methyl groups (LEU and VAL) were assigned stereospecifically.

	Total	^1H	^{13}C	^{15}N
Backbone	245/327 (75%)	131/134 (98%)	53/130 (41%)	61/63 (97%)
Sidechain	301/474 (64%)	248/306 (81%)	46/146 (32%)	7/22 (32%)

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	Total	¹ H	¹³ C	¹⁵ N
Aromatic	19/70 (27%)	18/35 (51%)	0/31 (0%)	1/4 (25%)
Overall	565/871 (65%)	397/475 (84%)	99/307 (32%)	69/89 (78%)

The following table shows the completeness of the chemical shift assignments for the full structure. The overall completeness is 64%, i.e. 570 atoms were assigned a chemical shift out of a possible 885. 0 out of 10 assigned methyl groups (LEU and VAL) were assigned stereospecifically.

	Total	¹ H	¹³ C	¹⁵ N
Backbone	247/338 (73%)	132/139 (95%)	54/134 (40%)	61/65 (94%)
Sidechain	304/477 (64%)	250/308 (81%)	47/147 (32%)	7/22 (32%)
Aromatic	19/70 (27%)	18/35 (51%)	0/31 (0%)	1/4 (25%)
Overall	570/885 (64%)	400/482 (83%)	101/312 (32%)	69/91 (76%)

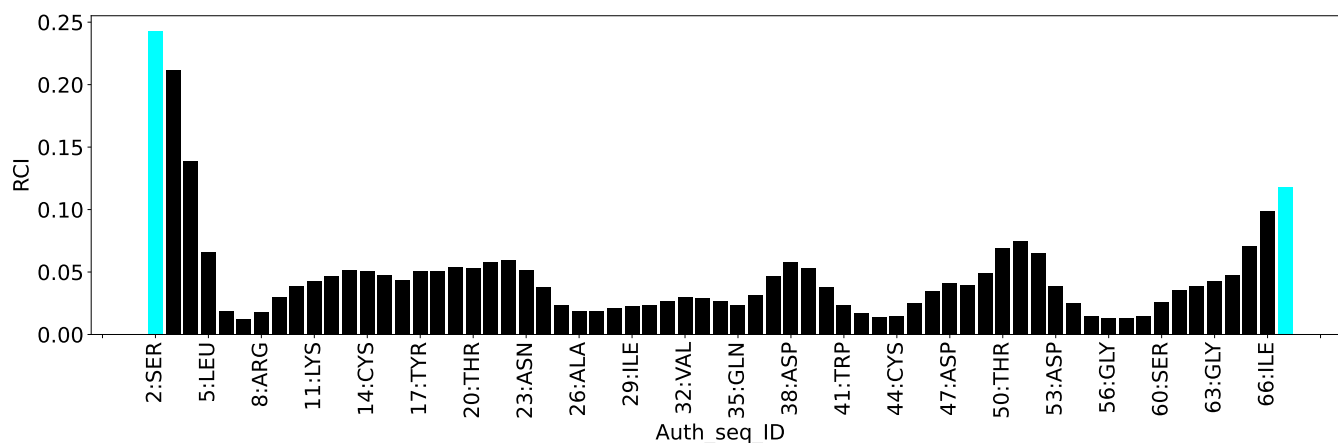
7.1.4 Statistically unusual chemical shifts [i](#)

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:



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	969
Intra-residue ($ i-j =0$)	355
Sequential ($ i-j =1$)	187
Medium range ($ i-j >1$ and $ i-j <5$)	89
Long range ($ i-j \geq 5$)	338
Inter-chain	0
Hydrogen bond restraints	0
Disulfide bond restraints	0
Total dihedral-angle restraints	0
Number of unmapped restraints	0
Number of restraints per residue	14.5
Number of long range restraints per residue ¹	5.0

¹Long 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)	25.1	0.2
0.2-0.5 (Medium)	64.1	0.5
>0.5 (Large)	107.2	7.57

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 ⓘ

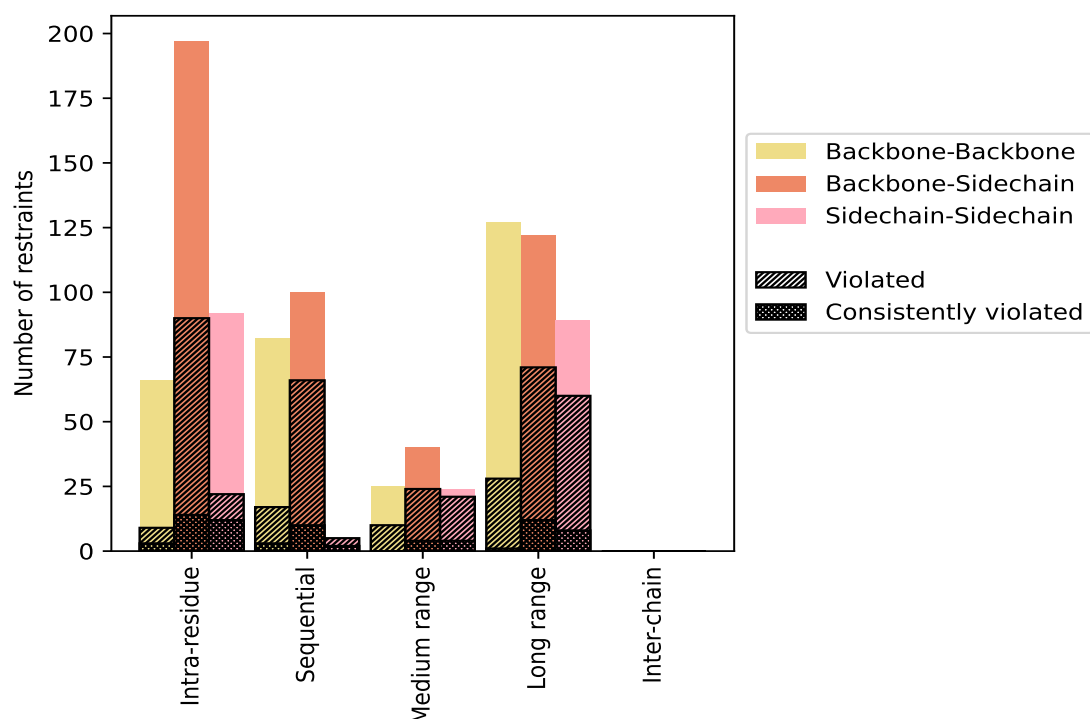
9.1 Summary of distance violations ⓘ

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 ($i-j =0$)	355	36.6	121	34.1	12.5	29	8.2	3.0
Backbone-Backbone	66	6.8	9	13.6	0.9	3	4.5	0.3
Backbone-Sidechain	197	20.3	90	45.7	9.3	14	7.1	1.4
Sidechain-Sidechain	92	9.5	22	23.9	2.3	12	13.0	1.2
Sequential ($i-j =1$)	187	19.3	88	47.1	9.1	15	8.0	1.5
Backbone-Backbone	82	8.5	17	20.7	1.8	3	3.7	0.3
Backbone-Sidechain	100	10.3	66	66.0	6.8	10	10.0	1.0
Sidechain-Sidechain	5	0.5	5	100.0	0.5	2	40.0	0.2
Medium range ($i-j >1$ & $i-j <5$)	89	9.2	55	61.8	5.7	8	9.0	0.8
Backbone-Backbone	25	2.6	10	40.0	1.0	0	0.0	0.0
Backbone-Sidechain	40	4.1	24	60.0	2.5	4	10.0	0.4
Sidechain-Sidechain	24	2.5	21	87.5	2.2	4	16.7	0.4
Long range ($i-j \geq 5$)	338	34.9	159	47.0	16.4	21	6.2	2.2
Backbone-Backbone	127	13.1	28	22.0	2.9	1	0.8	0.1
Backbone-Sidechain	122	12.6	71	58.2	7.3	12	9.8	1.2
Sidechain-Sidechain	89	9.2	60	67.4	6.2	8	9.0	0.8
Inter-chain	0	0.0	0	0.0	0.0	0	0.0	0.0
Backbone-Backbone	0	0.0	0	0.0	0.0	0	0.0	0.0
Backbone-Sidechain	0	0.0	0	0.0	0.0	0	0.0	0.0
Sidechain-Sidechain	0	0.0	0	0.0	0.0	0	0.0	0.0
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	969	100.0	423	43.7	43.7	73	7.5	7.5
Backbone-Backbone	300	31.0	64	21.3	6.6	7	2.3	0.7
Backbone-Sidechain	459	47.4	251	54.7	25.9	40	8.7	4.1
Sidechain-Sidechain	210	21.7	108	51.4	11.1	26	12.4	2.7

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 [i](#)



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 [i](#)

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	62	44	33	63	0	202	0.71	2.83	0.55	0.56
2	71	38	28	50	0	187	0.7	3.1	0.52	0.55
3	57	45	27	60	0	189	0.66	2.03	0.45	0.54
4	67	37	26	73	0	203	0.74	3.67	0.6	0.58
5	65	42	28	60	0	195	0.64	3.78	0.51	0.53
6	62	37	28	57	0	184	0.66	2.74	0.45	0.56
7	65	47	28	59	0	199	0.7	4.0	0.57	0.52
8	69	51	26	71	0	217	0.78	5.37	0.71	0.61
9	64	45	28	63	0	200	0.68	2.37	0.47	0.56
10	61	49	27	58	0	195	0.64	2.2	0.47	0.5

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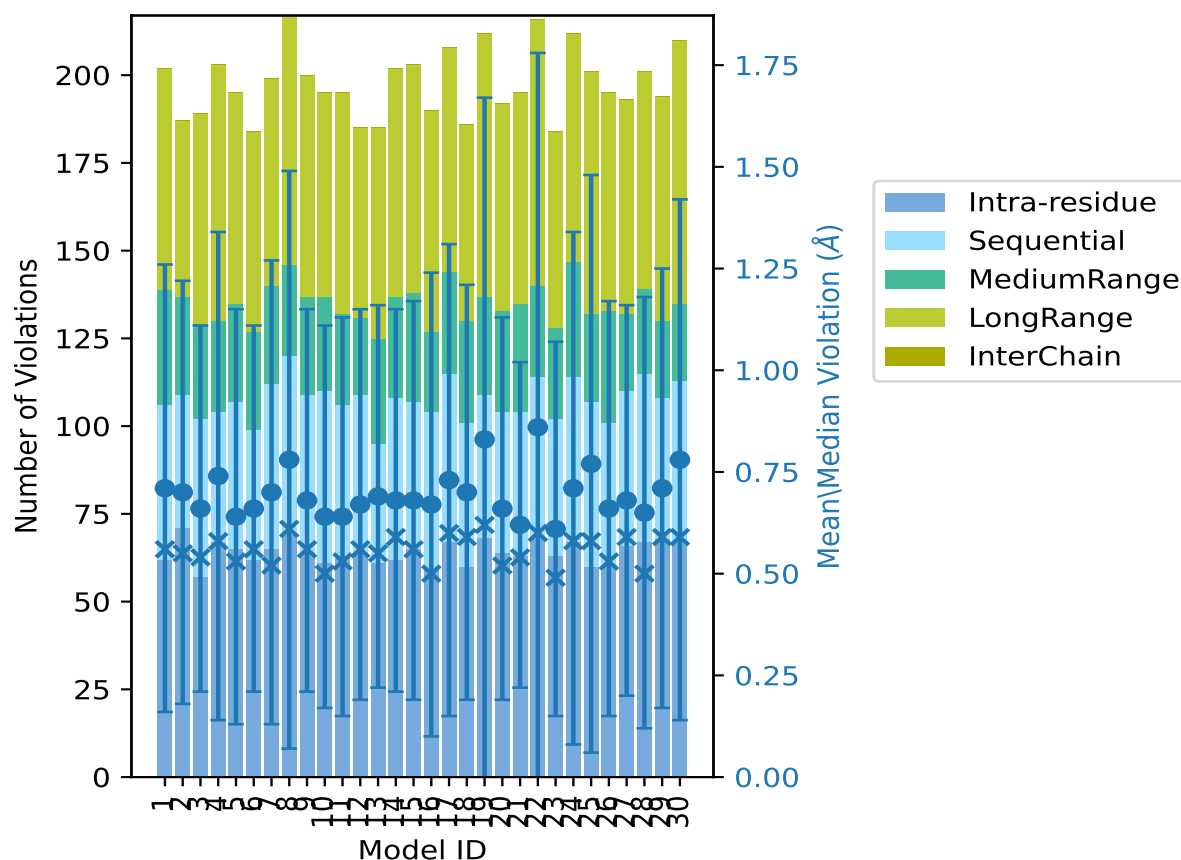
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Model ID	Number of violations						Mean (Å)	Max (Å)	SD ⁶ (Å)	Median (Å)
	IR ¹	SQ ²	MR ³	LR ⁴	IC ⁵	Total				
11	63	43	26	63	0	195	0.64	2.49	0.49	0.53
12	64	45	22	54	0	185	0.67	2.42	0.48	0.56
13	61	34	30	60	0	185	0.69	2.51	0.47	0.55
14	62	46	29	65	0	202	0.68	2.72	0.47	0.59
15	65	42	31	65	0	203	0.68	2.76	0.49	0.56
16	59	45	23	63	0	190	0.67	3.42	0.57	0.5
17	67	48	29	64	0	208	0.73	4.17	0.58	0.6
18	60	41	29	56	0	186	0.7	2.69	0.51	0.59
19	68	41	28	75	0	212	0.83	6.9	0.84	0.62
20	64	40	29	59	0	192	0.66	2.39	0.47	0.52
21	63	41	31	60	0	195	0.62	1.97	0.4	0.54
22	68	46	26	76	0	216	0.86	7.57	0.92	0.6
23	63	39	26	56	0	184	0.61	2.11	0.46	0.49
24	67	47	33	65	0	212	0.71	4.83	0.63	0.58
25	60	47	25	69	0	201	0.77	4.43	0.71	0.58
26	60	41	32	62	0	195	0.66	2.84	0.51	0.53
27	66	44	22	61	0	193	0.68	2.75	0.48	0.59
28	67	48	24	62	0	201	0.65	3.1	0.53	0.5
29	68	40	22	64	0	194	0.71	3.73	0.54	0.59
30	67	46	22	75	0	210	0.78	3.8	0.64	0.59

¹Intra-residue restraints, ²Sequential restraints, ³Medium range restraints, ⁴Long range restraints,

⁵Inter-chain restraints, ⁶Standard deviation

9.2.1 Bar graph : Distance Violation statistics for each model [i](#)



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, 546(IR:234, SQ:99, MR:34, LR:179, IC:0) 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	%
17	8	7	27	0	59	1	3.3
8	8	8	28	0	52	2	6.7
7	2	4	12	0	25	3	10.0
5	4	1	4	0	14	4	13.3
5	6	1	5	0	17	5	16.7
1	1	2	5	0	9	6	20.0

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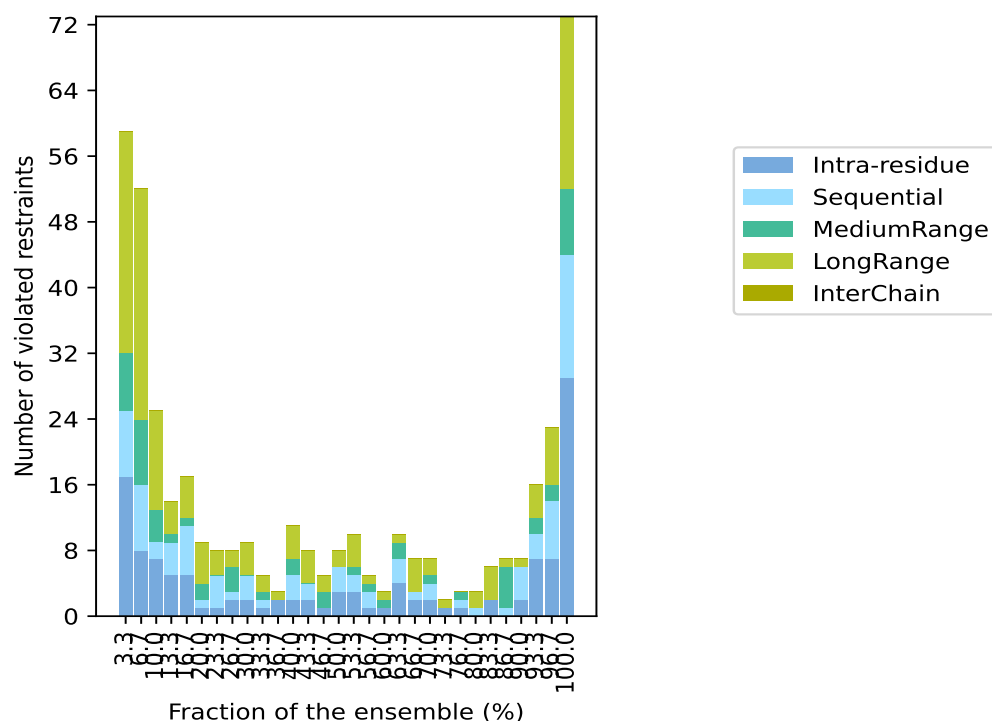
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Number of violated restraints						Fraction of the ensemble	
IR ¹	SQ ²	MR ³	LR ⁴	IC ⁵	Total	Count ⁶	%
1	4	0	3	0	8	7	23.3
2	1	3	2	0	8	8	26.7
2	3	0	4	0	9	9	30.0
1	1	1	2	0	5	10	33.3
2	0	0	1	0	3	11	36.7
2	3	2	4	0	11	12	40.0
2	2	0	4	0	8	13	43.3
1	0	2	2	0	5	14	46.7
3	3	0	2	0	8	15	50.0
3	2	1	4	0	10	16	53.3
1	2	1	1	0	5	17	56.7
1	0	1	1	0	3	18	60.0
4	3	2	1	0	10	19	63.3
2	1	0	4	0	7	20	66.7
2	2	1	2	0	7	21	70.0
1	0	0	1	0	2	22	73.3
1	1	1	0	0	3	23	76.7
0	1	0	2	0	3	24	80.0
2	0	0	4	0	6	25	83.3
0	1	5	1	0	7	26	86.7
2	4	0	1	0	7	27	90.0
7	3	2	4	0	16	28	93.3
7	7	2	7	0	23	29	96.7
29	15	8	21	0	73	30	100.0

¹Intra-residue restraints, ²Sequential restraints, ³Medium range restraints, ⁴Long range restraints,

⁵Inter-chain restraints, ⁶ Number of models with violations

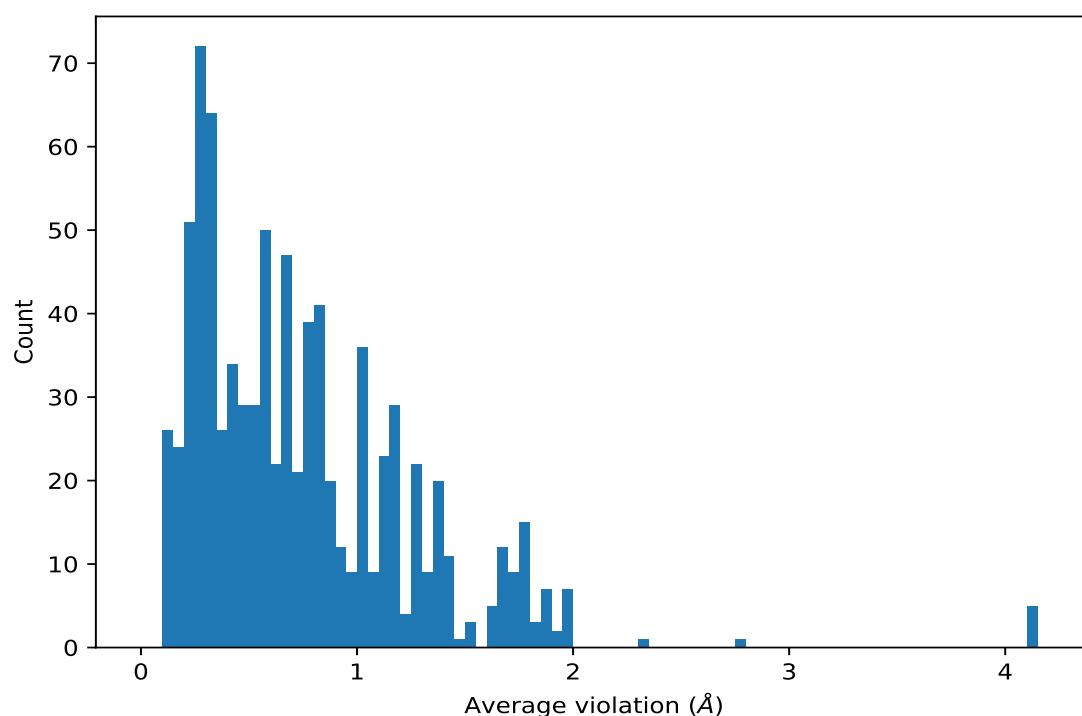
9.3.1 Bar graph : Distance violation statistics for the ensemble [i](#)



9.4 Most violated distance restraints in the ensemble [i](#)

9.4.1 Histogram : Distribution of mean distance violations [i](#)

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 [i](#)

The following table provides the mean and the standard deviation of the violation 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)
(1,916)	1:29:A:ILE:HG12	1:30:A:ILE:H	30	1.89	0.2	1.96
(1,916)	1:30:A:ILE:H	1:8:A:ARG:HB2	30	1.89	0.2	1.96
(1,916)	1:10:A:THR:H	1:62:A:LEU:HG	30	1.89	0.2	1.96
(1,916)	1:22:A:LEU:H	1:54:A:ARG:HB3	30	1.89	0.2	1.96
(1,916)	1:10:A:THR:H	1:11:A:LYS:HD2	30	1.89	0.2	1.96
(1,916)	1:22:A:LEU:H	1:54:A:ARG:HB2	30	1.89	0.2	1.96
(1,415)	1:5:A:LEU:HD11	1:7:A:VAL:HG23	30	1.76	0.46	1.66
(1,415)	1:5:A:LEU:HD11	1:7:A:VAL:HG21	30	1.76	0.46	1.66
(1,415)	1:5:A:LEU:HD13	1:7:A:VAL:HG23	30	1.76	0.46	1.66
(1,415)	1:5:A:LEU:HD12	1:7:A:VAL:HG21	30	1.76	0.46	1.66
(1,415)	1:5:A:LEU:HD12	1:7:A:VAL:HG22	30	1.76	0.46	1.66
(1,415)	1:5:A:LEU:HD12	1:7:A:VAL:HG23	30	1.76	0.46	1.66
(1,415)	1:5:A:LEU:HD13	1:7:A:VAL:HG22	30	1.76	0.46	1.66
(1,415)	1:5:A:LEU:HD11	1:7:A:VAL:HG22	30	1.76	0.46	1.66
(1,415)	1:5:A:LEU:HD13	1:7:A:VAL:HG21	30	1.76	0.46	1.66
(1,844)	1:22:A:LEU:HD13	1:13:A:TYR:HD2	30	1.73	0.31	1.7

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Key	Atom-1	Atom-2	Models ¹	Mean (Å)	SD ¹ (Å)	Median (Å)
(1,844)	1:22:A:LEU:HD12	1:13:A:TYR:HD2	30	1.73	0.31	1.7
(1,844)	1:22:A:LEU:HD11	1:13:A:TYR:HD2	30	1.73	0.31	1.7
(1,844)	1:32:A:VAL:HG23	1:41:A:TRP:HH2	30	1.73	0.31	1.7
(1,844)	1:32:A:VAL:HG21	1:41:A:TRP:HH2	30	1.73	0.31	1.7
(1,844)	1:32:A:VAL:HG22	1:41:A:TRP:HH2	30	1.73	0.31	1.7
(1,848)	1:16:A:ASN:HB2	1:18:A:ASP:H	30	1.68	0.73	1.65
(1,848)	1:16:A:ASN:HB2	1:17:A:TYR:HE1	30	1.68	0.73	1.65
(1,918)	1:13:A:TYR:H	1:11:A:LYS:HD3	30	1.43	0.48	1.41
(1,918)	1:13:A:TYR:H	1:11:A:LYS:HD2	30	1.43	0.48	1.41
(1,918)	1:13:A:TYR:H	1:25:A:LYS:HB3	30	1.43	0.48	1.41
(1,898)	1:22:A:LEU:HD11	1:44:A:CYS:HB3	30	1.4	0.37	1.41
(1,898)	1:32:A:VAL:HG22	1:60:A:SER:HA	30	1.4	0.37	1.41
(1,898)	1:32:A:VAL:HG21	1:60:A:SER:HA	30	1.4	0.37	1.41
(1,898)	1:32:A:VAL:HG23	1:60:A:SER:HA	30	1.4	0.37	1.41
(1,898)	1:22:A:LEU:HD13	1:23:A:ASN:HB3	30	1.4	0.37	1.41
(1,898)	1:22:A:LEU:HD12	1:44:A:CYS:HB2	30	1.4	0.37	1.41
(1,898)	1:22:A:LEU:HD12	1:23:A:ASN:HB2	30	1.4	0.37	1.41
(1,898)	1:22:A:LEU:HD13	1:23:A:ASN:HB2	30	1.4	0.37	1.41
(1,926)	1:6:A:GLN:H	1:65:A:ALA:HB3	30	1.39	0.44	1.4
(1,926)	1:6:A:GLN:H	1:65:A:ALA:HB2	30	1.39	0.44	1.4
(1,926)	1:28:A:ASP:H	1:25:A:LYS:HG2	30	1.39	0.44	1.4
(1,926)	1:28:A:ASP:H	1:25:A:LYS:HG3	30	1.39	0.44	1.4
(1,926)	1:6:A:GLN:H	1:65:A:ALA:HB1	30	1.39	0.44	1.4
(1,357)	1:65:A:ALA:HB3	1:5:A:LEU:HB3	30	1.37	0.37	1.4
(1,357)	1:65:A:ALA:HB2	1:5:A:LEU:HB3	30	1.37	0.37	1.4
(1,357)	1:65:A:ALA:HB1	1:5:A:LEU:HB3	30	1.37	0.37	1.4
(1,828)	1:23:A:ASN:H	1:22:A:LEU:HD22	30	1.35	0.1	1.37
(1,828)	1:23:A:ASN:H	1:22:A:LEU:HD23	30	1.35	0.1	1.37
(1,828)	1:23:A:ASN:H	1:22:A:LEU:HD21	30	1.35	0.1	1.37
(1,430)	1:22:A:LEU:HD11	1:45:A:ILE:HG21	30	1.31	0.13	1.31
(1,430)	1:22:A:LEU:HD13	1:45:A:ILE:HG22	30	1.31	0.13	1.31
(1,430)	1:22:A:LEU:HD11	1:45:A:ILE:HG22	30	1.31	0.13	1.31
(1,430)	1:22:A:LEU:HD13	1:45:A:ILE:HG23	30	1.31	0.13	1.31
(1,430)	1:22:A:LEU:HD12	1:45:A:ILE:HG22	30	1.31	0.13	1.31
(1,430)	1:22:A:LEU:HD12	1:45:A:ILE:HG21	30	1.31	0.13	1.31
(1,430)	1:22:A:LEU:HD11	1:45:A:ILE:HG23	30	1.31	0.13	1.31
(1,951)	1:8:A:ARG:H	1:7:A:VAL:HG22	30	1.29	0.11	1.3
(1,951)	1:13:A:TYR:H	1:24:A:VAL:HG21	30	1.29	0.11	1.3
(1,951)	1:13:A:TYR:H	1:24:A:VAL:HG13	30	1.29	0.11	1.3
(1,951)	1:8:A:ARG:H	1:7:A:VAL:HG23	30	1.29	0.11	1.3
(1,951)	1:13:A:TYR:H	1:24:A:VAL:HG11	30	1.29	0.11	1.3
(1,951)	1:8:A:ARG:H	1:7:A:VAL:HG21	30	1.29	0.11	1.3

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Key	Atom-1	Atom-2	Models ¹	Mean (Å)	SD ¹ (Å)	Median (Å)
(1,951)	1:13:A:TYR:H	1:24:A:VAL:HG12	30	1.29	0.11	1.3
(1,802)	1:26:A:ALA:HB2	1:12:A:ASP:HA	30	1.29	0.7	1.14
(1,802)	1:26:A:ALA:HB3	1:11:A:LYS:HA	30	1.29	0.7	1.14
(1,802)	1:26:A:ALA:HB3	1:12:A:ASP:HA	30	1.29	0.7	1.14
(1,802)	1:26:A:ALA:HB1	1:11:A:LYS:HA	30	1.29	0.7	1.14
(1,802)	1:26:A:ALA:HB2	1:11:A:LYS:HA	30	1.29	0.7	1.14
(1,802)	1:26:A:ALA:HB1	1:12:A:ASP:HA	30	1.29	0.7	1.14
(1,445)	1:30:A:ILE:HD12	1:45:A:ILE:HG21	30	1.2	0.45	1.31
(1,445)	1:30:A:ILE:HD13	1:45:A:ILE:HG21	30	1.2	0.45	1.31
(1,445)	1:30:A:ILE:HD12	1:45:A:ILE:HG22	30	1.2	0.45	1.31
(1,445)	1:30:A:ILE:HD13	1:45:A:ILE:HG22	30	1.2	0.45	1.31
(1,445)	1:30:A:ILE:HD11	1:45:A:ILE:HG23	30	1.2	0.45	1.31
(1,445)	1:30:A:ILE:HD11	1:45:A:ILE:HG22	30	1.2	0.45	1.31
(1,445)	1:30:A:ILE:HD12	1:45:A:ILE:HG23	30	1.2	0.45	1.31
(1,445)	1:30:A:ILE:HD13	1:45:A:ILE:HG23	30	1.2	0.45	1.31
(1,445)	1:30:A:ILE:HD11	1:45:A:ILE:HG21	30	1.2	0.45	1.31
(1,963)	1:30:A:ILE:HD13	1:22:A:LEU:H	30	1.2	0.24	1.25
(1,963)	1:10:A:THR:H	1:30:A:ILE:HD12	30	1.2	0.24	1.25
(1,963)	1:30:A:ILE:HD11	1:22:A:LEU:H	30	1.2	0.24	1.25
(1,963)	1:10:A:THR:H	1:30:A:ILE:HD11	30	1.2	0.24	1.25
(1,963)	1:30:A:ILE:HD12	1:22:A:LEU:H	30	1.2	0.24	1.25
(1,963)	1:10:A:THR:H	1:30:A:ILE:HD13	30	1.2	0.24	1.25
(1,813)	1:22:A:LEU:HB2	1:13:A:TYR:H	30	1.2	0.5	1.06
(1,27)	1:43:A:GLY:HA3	1:33:A:LEU:HD12	30	1.15	0.74	0.92
(1,27)	1:43:A:GLY:HA3	1:33:A:LEU:HD13	30	1.15	0.74	0.92
(1,27)	1:43:A:GLY:HA3	1:33:A:LEU:HD11	30	1.15	0.74	0.92
(1,812)	1:30:A:ILE:HG21	1:33:A:LEU:H	30	1.13	0.1	1.11
(1,812)	1:30:A:ILE:HG22	1:33:A:LEU:H	30	1.13	0.1	1.11
(1,812)	1:30:A:ILE:HG23	1:33:A:LEU:H	30	1.13	0.1	1.11
(1,950)	1:9:A:ALA:H	1:24:A:VAL:HG11	30	1.13	0.14	1.15
(1,950)	1:45:A:ILE:HD11	1:45:A:ILE:H	30	1.13	0.14	1.15
(1,950)	1:7:A:VAL:HG22	1:9:A:ALA:H	30	1.13	0.14	1.15
(1,950)	1:45:A:ILE:HD13	1:45:A:ILE:H	30	1.13	0.14	1.15
(1,950)	1:7:A:VAL:HG23	1:9:A:ALA:H	30	1.13	0.14	1.15
(1,950)	1:9:A:ALA:H	1:24:A:VAL:HG12	30	1.13	0.14	1.15
(1,950)	1:45:A:ILE:HD12	1:45:A:ILE:H	30	1.13	0.14	1.15
(1,950)	1:7:A:VAL:HG21	1:9:A:ALA:H	30	1.13	0.14	1.15
(1,950)	1:9:A:ALA:H	1:24:A:VAL:HG13	30	1.13	0.14	1.15
(1,468)	1:30:A:ILE:H	1:30:A:ILE:HG12	30	1.11	0.09	1.1
(1,959)	1:31:A:THR:H	1:22:A:LEU:HD11	30	1.11	0.27	1.08
(1,959)	1:31:A:THR:H	1:22:A:LEU:HD13	30	1.11	0.27	1.08
(1,959)	1:31:A:THR:H	1:22:A:LEU:HD12	30	1.11	0.27	1.08

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Key	Atom-1	Atom-2	Models ¹	Mean (Å)	SD ¹ (Å)	Median (Å)
(1,947)	1:7:A:VAL:HG22	1:9:A:ALA:H	30	1.07	0.13	1.08
(1,947)	1:45:A:ILE:HD11	1:45:A:ILE:H	30	1.07	0.13	1.08
(1,947)	1:45:A:ILE:HD13	1:45:A:ILE:H	30	1.07	0.13	1.08
(1,947)	1:7:A:VAL:HG23	1:9:A:ALA:H	30	1.07	0.13	1.08
(1,947)	1:45:A:ILE:HD12	1:45:A:ILE:H	30	1.07	0.13	1.08
(1,947)	1:7:A:VAL:HG21	1:9:A:ALA:H	30	1.07	0.13	1.08
(1,945)	1:41:A:TRP:HH2	1:5:A:LEU:HD12	30	1.04	0.46	1.03
(1,945)	1:41:A:TRP:HH2	1:5:A:LEU:HD11	30	1.04	0.46	1.03
(1,945)	1:41:A:TRP:HH2	1:5:A:LEU:HD13	30	1.04	0.46	1.03
(1,353)	1:22:A:LEU:HD13	1:22:A:LEU:HB2	30	1.03	0.02	1.03
(1,353)	1:22:A:LEU:HD11	1:22:A:LEU:HB2	30	1.03	0.02	1.03
(1,353)	1:22:A:LEU:HD12	1:22:A:LEU:HB2	30	1.03	0.02	1.03
(1,922)	1:66:A:ILE:H	1:6:A:GLN:HB3	30	1.03	0.32	1.02
(1,922)	1:23:A:ASN:H	1:22:A:LEU:HB2	30	1.03	0.32	1.02
(1,547)	1:6:A:GLN:H	1:5:A:LEU:HB3	30	1.03	0.1	1.02
(1,823)	1:28:A:ASP:H	1:9:A:ALA:HB3	30	1.0	0.36	0.98
(1,823)	1:28:A:ASP:H	1:25:A:LYS:HG2	30	1.0	0.36	0.98
(1,823)	1:28:A:ASP:H	1:9:A:ALA:HB1	30	1.0	0.36	0.98
(1,823)	1:28:A:ASP:H	1:25:A:LYS:HG3	30	1.0	0.36	0.98
(1,823)	1:28:A:ASP:H	1:9:A:ALA:HB2	30	1.0	0.36	0.98
(1,571)	1:66:A:ILE:H	1:65:A:ALA:HB3	30	0.97	0.29	1.02
(1,571)	1:66:A:ILE:H	1:65:A:ALA:HB2	30	0.97	0.29	1.02
(1,571)	1:66:A:ILE:H	1:65:A:ALA:HB1	30	0.97	0.29	1.02
(1,797)	1:7:A:VAL:HB	1:63:A:GLY:HA2	30	0.91	0.17	0.86
(1,797)	1:7:A:VAL:HB	1:64:A:GLU:HA	30	0.91	0.17	0.86
(1,322)	1:58:A:PHE:HE1	1:58:A:PHE:HB3	30	0.91	0.02	0.91
(1,440)	1:30:A:ILE:HG21	1:32:A:VAL:HG22	30	0.89	0.09	0.86
(1,440)	1:30:A:ILE:HG22	1:32:A:VAL:HG21	30	0.89	0.09	0.86
(1,440)	1:30:A:ILE:HG21	1:32:A:VAL:HG23	30	0.89	0.09	0.86
(1,440)	1:30:A:ILE:HG22	1:32:A:VAL:HG22	30	0.89	0.09	0.86
(1,440)	1:30:A:ILE:HG23	1:32:A:VAL:HG22	30	0.89	0.09	0.86
(1,440)	1:30:A:ILE:HG23	1:32:A:VAL:HG21	30	0.89	0.09	0.86
(1,440)	1:30:A:ILE:HG22	1:32:A:VAL:HG23	30	0.89	0.09	0.86
(1,440)	1:30:A:ILE:HG23	1:32:A:VAL:HG23	30	0.89	0.09	0.86
(1,440)	1:30:A:ILE:HG21	1:32:A:VAL:HG21	30	0.89	0.09	0.86
(1,529)	1:31:A:THR:H	1:30:A:ILE:HG13	30	0.88	0.03	0.88
(1,382)	1:65:A:ALA:H	1:5:A:LEU:HB3	30	0.85	0.37	0.92
(1,444)	1:30:A:ILE:HD11	1:32:A:VAL:HG12	30	0.84	0.28	0.84
(1,444)	1:30:A:ILE:HD13	1:32:A:VAL:HG12	30	0.84	0.28	0.84
(1,444)	1:30:A:ILE:HD11	1:32:A:VAL:HG13	30	0.84	0.28	0.84
(1,444)	1:30:A:ILE:HD12	1:32:A:VAL:HG12	30	0.84	0.28	0.84
(1,444)	1:30:A:ILE:HD12	1:32:A:VAL:HG13	30	0.84	0.28	0.84

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Key	Atom-1	Atom-2	Models ¹	Mean (Å)	SD ¹ (Å)	Median (Å)
(1,444)	1:30:A:ILE:HD13	1:32:A:VAL:HG11	30	0.84	0.28	0.84
(1,444)	1:30:A:ILE:HD11	1:32:A:VAL:HG11	30	0.84	0.28	0.84
(1,444)	1:30:A:ILE:HD12	1:32:A:VAL:HG11	30	0.84	0.28	0.84
(1,444)	1:30:A:ILE:HD13	1:32:A:VAL:HG13	30	0.84	0.28	0.84
(1,887)	1:29:A:ILE:HG22	1:31:A:THR:HB	30	0.82	0.25	0.79
(1,887)	1:29:A:ILE:HG21	1:31:A:THR:HB	30	0.82	0.25	0.79
(1,887)	1:29:A:ILE:HG23	1:31:A:THR:HB	30	0.82	0.25	0.79
(1,795)	1:17:A:TYR:HB3	1:18:A:ASP:H	30	0.79	0.39	0.68
(1,795)	1:15:A:ASN:HD22	1:15:A:ASN:HB3	30	0.79	0.39	0.68
(1,485)	1:46:A:HIS:H	1:45:A:ILE:HD11	30	0.79	0.16	0.8
(1,485)	1:46:A:HIS:H	1:45:A:ILE:HD13	30	0.79	0.16	0.8
(1,485)	1:46:A:HIS:H	1:45:A:ILE:HD12	30	0.79	0.16	0.8
(1,6)	1:55:A:VAL:HA	1:55:A:VAL:HG12	30	0.78	0.01	0.78
(1,6)	1:55:A:VAL:HA	1:55:A:VAL:HG11	30	0.78	0.01	0.78
(1,6)	1:55:A:VAL:HA	1:55:A:VAL:HG13	30	0.78	0.01	0.78
(1,939)	1:58:A:PHE:HD1	1:32:A:VAL:HG11	30	0.77	0.21	0.77
(1,939)	1:58:A:PHE:HD1	1:32:A:VAL:HG12	30	0.77	0.21	0.77
(1,939)	1:58:A:PHE:HD1	1:32:A:VAL:HG13	30	0.77	0.21	0.77
(1,939)	1:61:A:SER:H	1:32:A:VAL:HG11	30	0.77	0.21	0.77
(1,437)	1:24:A:VAL:HG21	1:30:A:ILE:HG22	30	0.75	0.29	0.74
(1,437)	1:24:A:VAL:HG22	1:30:A:ILE:HG23	30	0.75	0.29	0.74
(1,437)	1:24:A:VAL:HG21	1:30:A:ILE:HG23	30	0.75	0.29	0.74
(1,437)	1:24:A:VAL:HG22	1:30:A:ILE:HG21	30	0.75	0.29	0.74
(1,437)	1:24:A:VAL:HG22	1:30:A:ILE:HG22	30	0.75	0.29	0.74
(1,437)	1:24:A:VAL:HG23	1:30:A:ILE:HG21	30	0.75	0.29	0.74
(1,437)	1:24:A:VAL:HG23	1:30:A:ILE:HG23	30	0.75	0.29	0.74
(1,437)	1:24:A:VAL:HG21	1:30:A:ILE:HG21	30	0.75	0.29	0.74
(1,441)	1:30:A:ILE:HG23	1:45:A:ILE:HG21	30	0.67	0.18	0.68
(1,441)	1:30:A:ILE:HG21	1:45:A:ILE:HG21	30	0.67	0.18	0.68
(1,441)	1:30:A:ILE:HG21	1:45:A:ILE:HG22	30	0.67	0.18	0.68
(1,441)	1:30:A:ILE:HG22	1:45:A:ILE:HG22	30	0.67	0.18	0.68
(1,441)	1:30:A:ILE:HG23	1:45:A:ILE:HG22	30	0.67	0.18	0.68
(1,441)	1:30:A:ILE:HG22	1:45:A:ILE:HG23	30	0.67	0.18	0.68
(1,441)	1:30:A:ILE:HG23	1:45:A:ILE:HG23	30	0.67	0.18	0.68
(1,441)	1:30:A:ILE:HG21	1:45:A:ILE:HG23	30	0.67	0.18	0.68
(1,441)	1:30:A:ILE:HG22	1:45:A:ILE:HG21	30	0.67	0.18	0.68
(1,470)	1:30:A:ILE:H	1:30:A:ILE:HG21	30	0.63	0.02	0.63
(1,470)	1:30:A:ILE:H	1:30:A:ILE:HG22	30	0.63	0.02	0.63
(1,470)	1:30:A:ILE:H	1:30:A:ILE:HG23	30	0.63	0.02	0.63
(1,800)	1:44:A:CYS:HB3	1:55:A:VAL:HA	30	0.62	0.17	0.62
(1,800)	1:44:A:CYS:HB2	1:55:A:VAL:HA	30	0.62	0.17	0.62
(1,907)	1:11:A:LYS:HD2	1:11:A:LYS:HB2	30	0.61	0.03	0.62

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Key	Atom-1	Atom-2	Models ¹	Mean (Å)	SD ¹ (Å)	Median (Å)
(1,907)	1:11:A:LYS:HD3	1:11:A:LYS:HB3	30	0.61	0.03	0.62
(1,907)	1:11:A:LYS:HD2	1:11:A:LYS:HB3	30	0.61	0.03	0.62
(1,174)	1:30:A:ILE:HG12	1:9:A:ALA:H	30	0.58	0.15	0.55
(1,451)	1:25:A:LYS:H	1:24:A:VAL:HB	30	0.58	0.15	0.57
(1,448)	1:25:A:LYS:H	1:25:A:LYS:HB2	30	0.58	0.06	0.57
(1,496)	1:41:A:TRP:H	1:41:A:TRP:HB3	30	0.55	0.05	0.56
(1,659)	1:11:A:LYS:H	1:10:A:THR:H	30	0.54	0.04	0.53
(1,466)	1:30:A:ILE:H	1:29:A:ILE:HB	30	0.51	0.11	0.53
(1,923)	1:33:A:LEU:HD13	1:33:A:LEU:HB3	30	0.47	0.05	0.47
(1,923)	1:33:A:LEU:HD11	1:33:A:LEU:HB3	30	0.47	0.05	0.47
(1,923)	1:33:A:LEU:HD12	1:33:A:LEU:HB3	30	0.47	0.05	0.47
(1,923)	1:33:A:LEU:HD13	1:33:A:LEU:HB2	30	0.47	0.05	0.47
(1,399)	1:29:A:ILE:H	1:29:A:ILE:HD12	30	0.45	0.14	0.45
(1,399)	1:29:A:ILE:H	1:29:A:ILE:HD11	30	0.45	0.14	0.45
(1,399)	1:29:A:ILE:H	1:29:A:ILE:HD13	30	0.45	0.14	0.45
(1,480)	1:45:A:ILE:H	1:44:A:CYS:HB2	30	0.44	0.13	0.44
(1,803)	1:30:A:ILE:HG21	1:32:A:VAL:H	30	0.41	0.11	0.39
(1,803)	1:30:A:ILE:HG22	1:32:A:VAL:H	30	0.41	0.11	0.39
(1,803)	1:30:A:ILE:HG23	1:32:A:VAL:H	30	0.41	0.11	0.39
(1,964)	1:45:A:ILE:H	1:53:A:ASP:HA	30	0.38	0.16	0.36
(1,964)	1:45:A:ILE:H	1:55:A:VAL:HA	30	0.38	0.16	0.36
(1,71)	1:38:A:ASP:HB2	1:38:A:ASP:HA	30	0.38	0.28	0.22
(1,654)	1:17:A:TYR:H	1:17:A:TYR:HB3	30	0.38	0.16	0.31
(1,191)	1:6:A:GLN:H	1:6:A:GLN:HG2	30	0.37	0.07	0.37
(1,218)	1:41:A:TRP:HZ3	1:41:A:TRP:HE3	30	0.35	0.0	0.35
(1,60)	1:28:A:ASP:HB2	1:28:A:ASP:HA	30	0.34	0.17	0.29
(1,85)	1:7:A:VAL:HB	1:7:A:VAL:HG11	30	0.34	0.01	0.34
(1,85)	1:7:A:VAL:HB	1:7:A:VAL:HG13	30	0.34	0.01	0.34
(1,85)	1:7:A:VAL:HB	1:7:A:VAL:HG23	30	0.34	0.01	0.34
(1,85)	1:7:A:VAL:HB	1:7:A:VAL:HG12	30	0.34	0.01	0.34
(1,85)	1:7:A:VAL:HB	1:7:A:VAL:HG22	30	0.34	0.01	0.34
(1,372)	1:33:A:LEU:HD23	1:33:A:LEU:HG	30	0.34	0.01	0.34
(1,372)	1:33:A:LEU:HD22	1:33:A:LEU:HG	30	0.34	0.01	0.34
(1,372)	1:33:A:LEU:HD21	1:33:A:LEU:HG	30	0.34	0.01	0.34
(1,268)	1:30:A:ILE:HG13	1:30:A:ILE:HA	30	0.33	0.04	0.34
(1,943)	1:7:A:VAL:HB	1:7:A:VAL:HG11	30	0.33	0.01	0.33
(1,943)	1:7:A:VAL:HB	1:7:A:VAL:HG13	30	0.33	0.01	0.33
(1,943)	1:66:A:ILE:HG12	1:66:A:ILE:HD13	30	0.33	0.01	0.33
(1,943)	1:66:A:ILE:HB	1:66:A:ILE:HG23	30	0.33	0.01	0.33
(1,943)	1:66:A:ILE:HG12	1:66:A:ILE:HD11	30	0.33	0.01	0.33
(1,943)	1:7:A:VAL:HB	1:7:A:VAL:HG12	30	0.33	0.01	0.33
(1,943)	1:66:A:ILE:HB	1:66:A:ILE:HG21	30	0.33	0.01	0.33

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Key	Atom-1	Atom-2	Models ¹	Mean (Å)	SD ¹ (Å)	Median (Å)
(1,943)	1:7:A:VAL:HB	1:7:A:VAL:HG22	30	0.33	0.01	0.33
(1,667)	1:43:A:GLY:H	1:44:A:CYS:H	30	0.3	0.03	0.3
(1,882)	1:50:A:THR:H	1:49:A:ARG:HA	30	0.29	0.01	0.29
(1,882)	1:58:A:PHE:HD2	1:62:A:LEU:HA	30	0.29	0.01	0.29
(1,805)	1:30:A:ILE:H	1:30:A:ILE:HD11	30	0.28	0.07	0.28
(1,805)	1:30:A:ILE:H	1:30:A:ILE:HD12	30	0.28	0.07	0.28
(1,805)	1:30:A:ILE:H	1:30:A:ILE:HD13	30	0.28	0.07	0.28
(1,376)	1:32:A:VAL:HG23	1:32:A:VAL:HG12	30	0.25	0.02	0.26
(1,376)	1:32:A:VAL:HG22	1:32:A:VAL:HG13	30	0.25	0.02	0.26
(1,376)	1:32:A:VAL:HG21	1:32:A:VAL:HG13	30	0.25	0.02	0.26
(1,376)	1:32:A:VAL:HG23	1:32:A:VAL:HG11	30	0.25	0.02	0.26
(1,376)	1:32:A:VAL:HG23	1:32:A:VAL:HG13	30	0.25	0.02	0.26
(1,376)	1:32:A:VAL:HG22	1:32:A:VAL:HG11	30	0.25	0.02	0.26
(1,376)	1:32:A:VAL:HG22	1:32:A:VAL:HG12	30	0.25	0.02	0.26
(1,376)	1:32:A:VAL:HG21	1:32:A:VAL:HG12	30	0.25	0.02	0.26
(1,376)	1:32:A:VAL:HG21	1:32:A:VAL:HG11	30	0.25	0.02	0.26
(1,393)	1:33:A:LEU:HD22	1:33:A:LEU:HD11	30	0.25	0.03	0.24
(1,393)	1:33:A:LEU:HD23	1:33:A:LEU:HD12	30	0.25	0.03	0.24
(1,393)	1:33:A:LEU:HD23	1:33:A:LEU:HD11	30	0.25	0.03	0.24
(1,393)	1:33:A:LEU:HD23	1:33:A:LEU:HD13	30	0.25	0.03	0.24
(1,393)	1:33:A:LEU:HD22	1:33:A:LEU:HD13	30	0.25	0.03	0.24
(1,393)	1:33:A:LEU:HD21	1:33:A:LEU:HD11	30	0.25	0.03	0.24
(1,393)	1:33:A:LEU:HD21	1:33:A:LEU:HD12	30	0.25	0.03	0.24
(1,393)	1:33:A:LEU:HD22	1:33:A:LEU:HD12	30	0.25	0.03	0.24
(1,393)	1:33:A:LEU:HD21	1:33:A:LEU:HD13	30	0.25	0.03	0.24
(1,79)	1:17:A:TYR:HB2	1:17:A:TYR:HA	30	0.24	0.0	0.24
(1,221)	1:13:A:TYR:HE2	1:13:A:TYR:HD2	30	0.21	0.0	0.21
(1,221)	1:13:A:TYR:HE1	1:13:A:TYR:HD1	30	0.21	0.0	0.21
(1,649)	1:12:A:ASP:H	1:12:A:ASP:HA	30	0.2	0.02	0.2
(1,521)	1:5:A:LEU:H	1:5:A:LEU:HA	30	0.2	0.03	0.2
(1,559)	1:64:A:GLU:H	1:64:A:GLU:HA	30	0.18	0.03	0.17
(1,935)	1:58:A:PHE:HB3	1:40:A:ARG:HB2	29	1.66	0.6	1.54
(1,935)	1:40:A:ARG:HB3	1:38:A:ASP:HB3	29	1.66	0.6	1.54
(1,935)	1:12:A:ASP:HB3	1:11:A:LYS:HG3	29	1.66	0.6	1.54
(1,935)	1:13:A:TYR:HB3	1:11:A:LYS:HG3	29	1.66	0.6	1.54
(1,935)	1:40:A:ARG:HB2	1:38:A:ASP:HB3	29	1.66	0.6	1.54
(1,935)	1:13:A:TYR:HB3	1:9:A:ALA:HB3	29	1.66	0.6	1.54
(1,942)	1:32:A:VAL:HB	1:7:A:VAL:HG23	29	1.2	0.09	1.22
(1,942)	1:32:A:VAL:HB	1:7:A:VAL:HG21	29	1.2	0.09	1.22
(1,942)	1:32:A:VAL:HB	1:7:A:VAL:HG22	29	1.2	0.09	1.22
(1,942)	1:32:A:VAL:HB	1:5:A:LEU:HD12	29	1.2	0.09	1.22
(1,617)	1:21:A:SER:H	1:21:A:SER:HB3	29	0.93	0.13	0.99

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Key	Atom-1	Atom-2	Models ¹	Mean (Å)	SD ¹ (Å)	Median (Å)
(1,925)	1:11:A:LYS:HG2	1:13:A:TYR:H	29	0.87	0.49	0.76
(1,685)	1:27:A:GLY:H	1:24:A:VAL:HG22	29	0.85	0.63	0.83
(1,685)	1:27:A:GLY:H	1:24:A:VAL:HG23	29	0.85	0.63	0.83
(1,685)	1:27:A:GLY:H	1:24:A:VAL:HG21	29	0.85	0.63	0.83
(1,536)	1:23:A:ASN:H	1:22:A:LEU:HD13	29	0.82	0.2	0.88
(1,536)	1:23:A:ASN:H	1:22:A:LEU:HD12	29	0.82	0.2	0.88
(1,536)	1:23:A:ASN:H	1:22:A:LEU:HD11	29	0.82	0.2	0.88
(1,934)	1:26:A:ALA:H	1:25:A:LYS:HG2	29	0.79	0.25	0.83
(1,934)	1:26:A:ALA:H	1:25:A:LYS:HG3	29	0.79	0.25	0.83
(1,941)	1:33:A:LEU:H	1:33:A:LEU:HG	29	0.78	0.09	0.77
(1,941)	1:32:A:VAL:HG12	1:33:A:LEU:H	29	0.78	0.09	0.77
(1,941)	1:32:A:VAL:HG13	1:33:A:LEU:H	29	0.78	0.09	0.77
(1,912)	1:41:A:TRP:H	1:40:A:ARG:HG3	29	0.74	0.28	0.81
(1,912)	1:34:A:GLU:HB3	1:35:A:GLN:H	29	0.74	0.28	0.81
(1,121)	1:26:A:ALA:H	1:26:A:ALA:HB2	29	0.71	0.05	0.72
(1,121)	1:26:A:ALA:H	1:26:A:ALA:HB3	29	0.71	0.05	0.72
(1,121)	1:26:A:ALA:H	1:26:A:ALA:HB1	29	0.71	0.05	0.72
(1,863)	1:9:A:ALA:H	1:29:A:ILE:HA	29	0.69	0.37	0.72
(1,863)	1:25:A:LYS:H	1:12:A:ASP:HA	29	0.69	0.37	0.72
(1,687)	1:27:A:GLY:H	1:26:A:ALA:HB3	29	0.68	0.12	0.69
(1,687)	1:27:A:GLY:H	1:26:A:ALA:HB2	29	0.68	0.12	0.69
(1,687)	1:27:A:GLY:H	1:26:A:ALA:HB1	29	0.68	0.12	0.69
(1,650)	1:12:A:ASP:H	1:11:A:LYS:HB2	29	0.65	0.06	0.66
(1,840)	1:11:A:LYS:HD3	1:12:A:ASP:H	29	0.63	0.3	0.69
(1,840)	1:12:A:ASP:H	1:11:A:LYS:HD2	29	0.63	0.3	0.69
(1,420)	1:7:A:VAL:HG13	1:32:A:VAL:HG13	29	0.58	0.15	0.58
(1,420)	1:7:A:VAL:HG13	1:32:A:VAL:HG11	29	0.58	0.15	0.58
(1,420)	1:7:A:VAL:HG11	1:32:A:VAL:HG13	29	0.58	0.15	0.58
(1,420)	1:7:A:VAL:HG13	1:32:A:VAL:HG12	29	0.58	0.15	0.58
(1,420)	1:7:A:VAL:HG12	1:32:A:VAL:HG13	29	0.58	0.15	0.58
(1,420)	1:7:A:VAL:HG12	1:32:A:VAL:HG11	29	0.58	0.15	0.58
(1,420)	1:7:A:VAL:HG12	1:32:A:VAL:HG12	29	0.58	0.15	0.58
(1,420)	1:7:A:VAL:HG11	1:32:A:VAL:HG11	29	0.58	0.15	0.58
(1,420)	1:7:A:VAL:HG11	1:32:A:VAL:HG12	29	0.58	0.15	0.58
(1,946)	1:7:A:VAL:HG22	1:30:A:ILE:H	29	0.56	0.16	0.57
(1,946)	1:7:A:VAL:HG23	1:30:A:ILE:H	29	0.56	0.16	0.57
(1,946)	1:7:A:VAL:HG21	1:30:A:ILE:H	29	0.56	0.16	0.57
(1,946)	1:32:A:VAL:H	1:7:A:VAL:HG23	29	0.56	0.16	0.57
(1,109)	1:32:A:VAL:H	1:32:A:VAL:HB	29	0.53	0.02	0.52
(1,810)	1:23:A:ASN:H	1:45:A:ILE:H	29	0.49	0.23	0.44
(1,804)	1:22:A:LEU:HB3	1:22:A:LEU:H	29	0.34	0.16	0.29
(1,804)	1:22:A:LEU:HB3	1:24:A:VAL:H	29	0.34	0.16	0.29

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Key	Atom-1	Atom-2	Models ¹	Mean (Å)	SD ¹ (Å)	Median (Å)
(1,816)	1:5:A:LEU:HB2	1:7:A:VAL:H	29	0.33	0.12	0.32
(1,816)	1:57:A:TYR:H	1:22:A:LEU:HB2	29	0.33	0.12	0.32
(1,289)	1:32:A:VAL:HA	1:32:A:VAL:HG21	29	0.27	0.01	0.27
(1,289)	1:32:A:VAL:HA	1:32:A:VAL:HG23	29	0.27	0.01	0.27
(1,289)	1:32:A:VAL:HA	1:32:A:VAL:HG22	29	0.27	0.01	0.27
(1,581)	1:13:A:TYR:H	1:13:A:TYR:HA	29	0.16	0.02	0.16
(1,885)	1:62:A:LEU:HB3	1:62:A:LEU:HA	29	0.12	0.0	0.12
(1,885)	1:62:A:LEU:HG	1:62:A:LEU:HA	29	0.12	0.0	0.12
(1,850)	1:11:A:LYS:HG3	1:13:A:TYR:HE1	28	1.46	0.42	1.52
(1,958)	1:58:A:PHE:HD1	1:33:A:LEU:HD12	28	1.29	0.63	1.39
(1,958)	1:58:A:PHE:HD1	1:33:A:LEU:HD13	28	1.29	0.63	1.39
(1,958)	1:61:A:SER:H	1:5:A:LEU:HD23	28	1.29	0.63	1.39
(1,958)	1:61:A:SER:H	1:5:A:LEU:HD22	28	1.29	0.63	1.39
(1,958)	1:61:A:SER:H	1:5:A:LEU:HD21	28	1.29	0.63	1.39
(1,958)	1:58:A:PHE:HD1	1:33:A:LEU:HD11	28	1.29	0.63	1.39
(1,416)	1:5:A:LEU:HD21	1:7:A:VAL:HG13	28	1.02	0.21	1.06
(1,416)	1:5:A:LEU:HD22	1:7:A:VAL:HG13	28	1.02	0.21	1.06
(1,416)	1:5:A:LEU:HD23	1:7:A:VAL:HG12	28	1.02	0.21	1.06
(1,416)	1:5:A:LEU:HD23	1:7:A:VAL:HG13	28	1.02	0.21	1.06
(1,416)	1:5:A:LEU:HD21	1:7:A:VAL:HG12	28	1.02	0.21	1.06
(1,416)	1:5:A:LEU:HD23	1:7:A:VAL:HG11	28	1.02	0.21	1.06
(1,416)	1:5:A:LEU:HD22	1:7:A:VAL:HG12	28	1.02	0.21	1.06
(1,928)	1:9:A:ALA:HB2	1:11:A:LYS:H	28	0.95	0.16	0.99
(1,928)	1:9:A:ALA:HB3	1:11:A:LYS:H	28	0.95	0.16	0.99
(1,928)	1:60:A:SER:HB3	1:41:A:TRP:HH2	28	0.95	0.16	0.99
(1,928)	1:9:A:ALA:HB1	1:11:A:LYS:H	28	0.95	0.16	0.99
(1,928)	1:11:A:LYS:H	1:11:A:LYS:HG2	28	0.95	0.16	0.99
(1,528)	1:31:A:THR:H	1:31:A:THR:HG22	28	0.88	0.03	0.88
(1,528)	1:31:A:THR:H	1:31:A:THR:HG23	28	0.88	0.03	0.88
(1,528)	1:31:A:THR:H	1:31:A:THR:HG21	28	0.88	0.03	0.88
(1,220)	1:15:A:ASN:H	1:15:A:ASN:HD22	28	0.86	0.67	0.6
(1,892)	1:13:A:TYR:HD2	1:56:A:GLY:HA3	28	0.79	0.36	0.86
(1,892)	1:41:A:TRP:HH2	1:61:A:SER:HA	28	0.79	0.36	0.86
(1,892)	1:13:A:TYR:HD2	1:24:A:VAL:HA	28	0.79	0.36	0.86
(1,892)	1:13:A:TYR:HD1	1:24:A:VAL:HA	28	0.79	0.36	0.86
(1,956)	1:24:A:VAL:HG23	1:13:A:TYR:HD2	28	0.68	0.3	0.7
(1,956)	1:24:A:VAL:HG21	1:13:A:TYR:HD2	28	0.68	0.3	0.7
(1,956)	1:5:A:LEU:HD23	1:41:A:TRP:HH2	28	0.68	0.3	0.7
(1,956)	1:24:A:VAL:HG22	1:13:A:TYR:HD2	28	0.68	0.3	0.7
(1,956)	1:24:A:VAL:HG23	1:13:A:TYR:HD1	28	0.68	0.3	0.7
(1,956)	1:5:A:LEU:HD21	1:41:A:TRP:HH2	28	0.68	0.3	0.7
(1,507)	1:22:A:LEU:H	1:22:A:LEU:HD13	28	0.59	0.13	0.57

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Key	Atom-1	Atom-2	Models ¹	Mean (Å)	SD ¹ (Å)	Median (Å)
(1,507)	1:22:A:LEU:H	1:22:A:LEU:HD12	28	0.59	0.13	0.57
(1,507)	1:22:A:LEU:H	1:22:A:LEU:HD11	28	0.59	0.13	0.57
(1,403)	1:58:A:PHE:H	1:22:A:LEU:HD12	28	0.53	0.24	0.52
(1,403)	1:58:A:PHE:H	1:22:A:LEU:HD11	28	0.53	0.24	0.52
(1,403)	1:58:A:PHE:H	1:22:A:LEU:HD13	28	0.53	0.24	0.52
(1,814)	1:26:A:ALA:H	1:25:A:LYS:HB2	28	0.52	0.06	0.54
(1,467)	1:30:A:ILE:H	1:29:A:ILE:HG22	28	0.52	0.2	0.52
(1,467)	1:30:A:ILE:H	1:29:A:ILE:HG23	28	0.52	0.2	0.52
(1,467)	1:30:A:ILE:H	1:29:A:ILE:HG21	28	0.52	0.2	0.52
(1,627)	1:38:A:ASP:H	1:37:A:PRO:HB3	28	0.37	0.12	0.39
(1,107)	1:41:A:TRP:HB2	1:41:A:TRP:HD1	28	0.34	0.1	0.37
(1,54)	1:13:A:TYR:HB3	1:13:A:TYR:HA	28	0.2	0.01	0.21
(1,223)	1:48:A:ASN:HD22	1:48:A:ASN:HB2	28	0.12	0.01	0.12
(1,223)	1:48:A:ASN:HB3	1:48:A:ASN:HD22	28	0.12	0.01	0.12
(1,927)	1:41:A:TRP:HD1	1:40:A:ARG:HB2	27	1.6	0.44	1.73
(1,927)	1:41:A:TRP:HD1	1:40:A:ARG:HB3	27	1.6	0.44	1.73
(1,927)	1:48:A:ASN:HD22	1:25:A:LYS:HG3	27	1.6	0.44	1.73
(1,927)	1:48:A:ASN:HD22	1:25:A:LYS:HG2	27	1.6	0.44	1.73
(1,514)	1:65:A:ALA:H	1:64:A:GLU:HB2	27	0.73	0.09	0.73
(1,940)	1:44:A:CYS:H	1:32:A:VAL:HG11	27	0.42	0.15	0.41
(1,940)	1:7:A:VAL:H	1:32:A:VAL:HG11	27	0.42	0.15	0.41
(1,940)	1:44:A:CYS:H	1:32:A:VAL:HG13	27	0.42	0.15	0.41
(1,940)	1:44:A:CYS:H	1:32:A:VAL:HG12	27	0.42	0.15	0.41
(1,940)	1:7:A:VAL:H	1:32:A:VAL:HG12	27	0.42	0.15	0.41
(1,940)	1:7:A:VAL:H	1:32:A:VAL:HG13	27	0.42	0.15	0.41
(1,884)	1:62:A:LEU:H	1:62:A:LEU:HA	27	0.38	0.03	0.4
(1,330)	1:63:A:GLY:H	1:62:A:LEU:HB2	27	0.38	0.24	0.3
(1,279)	1:63:A:GLY:H	1:62:A:LEU:HA	27	0.25	0.05	0.27
(1,355)	1:33:A:LEU:HB3	1:33:A:LEU:H	27	0.25	0.05	0.24
(1,46)	1:13:A:TYR:HB2	1:11:A:LYS:HD2	26	1.03	0.91	0.74
(1,919)	1:23:A:ASN:HD21	1:54:A:ARG:HB2	26	1.01	0.64	0.76
(1,919)	1:23:A:ASN:HD21	1:25:A:LYS:HB3	26	1.01	0.64	0.76
(1,919)	1:23:A:ASN:HD21	1:54:A:ARG:HB3	26	1.01	0.64	0.76
(1,834)	1:51:A:GLY:HA3	1:47:A:ASP:H	26	0.84	0.38	0.77
(1,834)	1:39:A:GLY:HA3	1:41:A:TRP:H	26	0.84	0.38	0.77
(1,894)	1:23:A:ASN:H	1:54:A:ARG:HD2	26	0.83	0.72	0.5
(1,894)	1:23:A:ASN:H	1:54:A:ARG:HD3	26	0.83	0.72	0.5
(1,894)	1:53:A:ASP:H	1:54:A:ARG:HD2	26	0.83	0.72	0.5
(1,894)	1:53:A:ASP:H	1:54:A:ARG:HD3	26	0.83	0.72	0.5
(1,966)	1:8:A:ARG:H	1:64:A:GLU:HG3	26	0.73	0.34	0.72
(1,966)	1:8:A:ARG:H	1:64:A:GLU:HG2	26	0.73	0.34	0.72
(1,966)	1:8:A:ARG:H	1:6:A:GLN:HG2	26	0.73	0.34	0.72

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Key	Atom-1	Atom-2	Models ¹	Mean (Å)	SD ¹ (Å)	Median (Å)
(1,431)	1:22:A:LEU:HD11	1:24:A:VAL:HG11	26	0.65	0.24	0.67
(1,431)	1:22:A:LEU:HD11	1:24:A:VAL:HG12	26	0.65	0.24	0.67
(1,431)	1:22:A:LEU:HD13	1:24:A:VAL:HG11	26	0.65	0.24	0.67
(1,431)	1:22:A:LEU:HD11	1:24:A:VAL:HG13	26	0.65	0.24	0.67
(1,431)	1:22:A:LEU:HD13	1:24:A:VAL:HG13	26	0.65	0.24	0.67
(1,431)	1:22:A:LEU:HD12	1:24:A:VAL:HG13	26	0.65	0.24	0.67
(1,431)	1:22:A:LEU:HD12	1:24:A:VAL:HG12	26	0.65	0.24	0.67
(1,431)	1:22:A:LEU:HD12	1:24:A:VAL:HG11	26	0.65	0.24	0.67
(1,431)	1:22:A:LEU:HD13	1:24:A:VAL:HG12	26	0.65	0.24	0.67
(1,35)	1:27:A:GLY:HA3	1:28:A:ASP:H	26	0.23	0.04	0.25
(1,462)	1:67:A:VAL:H	1:67:A:VAL:HG13	25	1.54	0.05	1.56
(1,462)	1:67:A:VAL:H	1:67:A:VAL:HG11	25	1.54	0.05	1.56
(1,462)	1:67:A:VAL:H	1:67:A:VAL:HG12	25	1.54	0.05	1.56
(1,544)	1:40:A:ARG:H	1:40:A:ARG:HB2	25	0.65	0.11	0.67
(1,852)	1:13:A:TYR:HE2	1:24:A:VAL:H	25	0.62	0.3	0.63
(1,852)	1:25:A:LYS:H	1:48:A:ASN:HD22	25	0.62	0.3	0.63
(1,852)	1:13:A:TYR:HE1	1:24:A:VAL:H	25	0.62	0.3	0.63
(1,900)	1:23:A:ASN:HB3	1:24:A:VAL:H	25	0.46	0.34	0.3
(1,900)	1:14:A:CYS:HB3	1:24:A:VAL:H	25	0.46	0.34	0.3
(1,900)	1:24:A:VAL:H	1:14:A:CYS:HB2	25	0.46	0.34	0.3
(1,900)	1:24:A:VAL:H	1:23:A:ASN:HB2	25	0.46	0.34	0.3
(1,944)	1:30:A:ILE:HD11	1:7:A:VAL:HG22	25	0.35	0.16	0.38
(1,944)	1:30:A:ILE:HD13	1:45:A:ILE:HD11	25	0.35	0.16	0.38
(1,944)	1:30:A:ILE:HD12	1:7:A:VAL:HG23	25	0.35	0.16	0.38
(1,944)	1:30:A:ILE:HD11	1:7:A:VAL:HG23	25	0.35	0.16	0.38
(1,944)	1:30:A:ILE:HD11	1:7:A:VAL:HG21	25	0.35	0.16	0.38
(1,944)	1:30:A:ILE:HD12	1:7:A:VAL:HG22	25	0.35	0.16	0.38
(1,944)	1:30:A:ILE:HD13	1:7:A:VAL:HG22	25	0.35	0.16	0.38
(1,944)	1:30:A:ILE:HD13	1:7:A:VAL:HG21	25	0.35	0.16	0.38
(1,944)	1:30:A:ILE:HD12	1:7:A:VAL:HG21	25	0.35	0.16	0.38
(1,422)	1:7:A:VAL:HG23	1:32:A:VAL:HG13	25	0.21	0.1	0.2
(1,422)	1:7:A:VAL:HG21	1:32:A:VAL:HG12	25	0.21	0.1	0.2
(1,422)	1:7:A:VAL:HG21	1:32:A:VAL:HG11	25	0.21	0.1	0.2
(1,422)	1:7:A:VAL:HG23	1:32:A:VAL:HG11	25	0.21	0.1	0.2
(1,422)	1:7:A:VAL:HG23	1:32:A:VAL:HG12	25	0.21	0.1	0.2
(1,422)	1:7:A:VAL:HG22	1:32:A:VAL:HG12	25	0.21	0.1	0.2
(1,422)	1:7:A:VAL:HG21	1:32:A:VAL:HG13	25	0.21	0.1	0.2
(1,422)	1:7:A:VAL:HG22	1:32:A:VAL:HG13	25	0.21	0.1	0.2
(1,422)	1:7:A:VAL:HG22	1:32:A:VAL:HG11	25	0.21	0.1	0.2
(1,909)	1:41:A:TRP:HD1	1:40:A:ARG:HG2	24	0.55	0.3	0.46
(1,909)	1:41:A:TRP:HD1	1:60:A:SER:HB2	24	0.55	0.3	0.46
(1,968)	1:10:A:THR:H	1:63:A:GLY:HA3	24	0.43	0.23	0.4

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Key	Atom-1	Atom-2	Models ¹	Mean (Å)	SD ¹ (Å)	Median (Å)
(1,968)	1:10:A:THR:H	1:26:A:ALA:HA	24	0.43	0.23	0.4
(1,423)	1:7:A:VAL:HG23	1:32:A:VAL:HG22	24	0.33	0.39	0.24
(1,423)	1:7:A:VAL:HG21	1:32:A:VAL:HG22	24	0.33	0.39	0.24
(1,423)	1:7:A:VAL:HG22	1:32:A:VAL:HG22	24	0.33	0.39	0.24
(1,423)	1:7:A:VAL:HG23	1:32:A:VAL:HG23	24	0.33	0.39	0.24
(1,423)	1:7:A:VAL:HG23	1:32:A:VAL:HG21	24	0.33	0.39	0.24
(1,423)	1:7:A:VAL:HG21	1:32:A:VAL:HG23	24	0.33	0.39	0.24
(1,423)	1:7:A:VAL:HG22	1:32:A:VAL:HG23	24	0.33	0.39	0.24
(1,423)	1:7:A:VAL:HG21	1:32:A:VAL:HG21	24	0.33	0.39	0.24
(1,423)	1:7:A:VAL:HG22	1:32:A:VAL:HG21	24	0.33	0.39	0.24
(1,847)	1:8:A:ARG:HD3	1:6:A:GLN:HE21	23	1.77	0.93	1.56
(1,847)	1:36:A:HIS:HB2	1:41:A:TRP:HZ2	23	1.77	0.93	1.56
(1,493)	1:41:A:TRP:HE1	1:40:A:ARG:H	23	0.78	0.36	0.8
(1,917)	1:25:A:LYS:HB3	1:25:A:LYS:H	23	0.4	0.16	0.39
(1,917)	1:54:A:ARG:HB3	1:45:A:ILE:H	23	0.4	0.16	0.39
(1,917)	1:54:A:ARG:HB2	1:45:A:ILE:H	23	0.4	0.16	0.39
(1,948)	1:6:A:GLN:H	1:7:A:VAL:HG13	22	1.17	0.56	1.31
(1,948)	1:6:A:GLN:H	1:67:A:VAL:HG11	22	1.17	0.56	1.31
(1,948)	1:6:A:GLN:H	1:7:A:VAL:HG12	22	1.17	0.56	1.31
(1,948)	1:6:A:GLN:H	1:67:A:VAL:HG22	22	1.17	0.56	1.31
(1,948)	1:6:A:GLN:H	1:67:A:VAL:HG21	22	1.17	0.56	1.31
(1,948)	1:6:A:GLN:H	1:67:A:VAL:HG23	22	1.17	0.56	1.31
(1,948)	1:6:A:GLN:H	1:67:A:VAL:HG13	22	1.17	0.56	1.31
(1,799)	1:41:A:TRP:H	1:41:A:TRP:HB3	22	0.19	0.04	0.2
(1,878)	1:52:A:ASN:H	1:50:A:THR:HB	21	0.73	0.57	0.5
(1,878)	1:52:A:ASN:H	1:50:A:THR:HA	21	0.73	0.57	0.5
(1,541)	1:40:A:ARG:H	1:40:A:ARG:HD2	21	0.65	0.25	0.59
(1,870)	1:29:A:ILE:HG22	1:45:A:ILE:HA	21	0.52	0.31	0.45
(1,870)	1:29:A:ILE:HG21	1:45:A:ILE:HA	21	0.52	0.31	0.45
(1,870)	1:29:A:ILE:HG23	1:45:A:ILE:HA	21	0.52	0.31	0.45
(1,870)	1:29:A:ILE:HG21	1:28:A:ASP:HA	21	0.52	0.31	0.45
(1,395)	1:42:A:LYS:H	1:33:A:LEU:HD12	21	0.47	0.5	0.34
(1,395)	1:42:A:LYS:H	1:33:A:LEU:HD11	21	0.47	0.5	0.34
(1,395)	1:42:A:LYS:H	1:33:A:LEU:HD13	21	0.47	0.5	0.34
(1,798)	1:24:A:VAL:HB	1:12:A:ASP:HA	21	0.43	0.22	0.37
(1,798)	1:24:A:VAL:HB	1:13:A:TYR:HA	21	0.43	0.22	0.37
(1,954)	1:44:A:CYS:HA	1:45:A:ILE:HD11	21	0.28	0.13	0.24
(1,954)	1:44:A:CYS:HA	1:45:A:ILE:HD13	21	0.28	0.13	0.24
(1,954)	1:44:A:CYS:HA	1:45:A:ILE:HD12	21	0.28	0.13	0.24
(1,583)	1:55:A:VAL:H	1:55:A:VAL:HB	21	0.24	0.1	0.22
(1,427)	1:20:A:THR:HG22	1:55:A:VAL:HG21	20	1.98	2.26	0.94
(1,427)	1:20:A:THR:HG21	1:55:A:VAL:HG23	20	1.98	2.26	0.94

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Key	Atom-1	Atom-2	Models ¹	Mean (Å)	SD ¹ (Å)	Median (Å)
(1,427)	1:20:A:THR:HG21	1:55:A:VAL:HG21	20	1.98	2.26	0.94
(1,427)	1:20:A:THR:HG22	1:55:A:VAL:HG22	20	1.98	2.26	0.94
(1,427)	1:20:A:THR:HG23	1:55:A:VAL:HG21	20	1.98	2.26	0.94
(1,427)	1:20:A:THR:HG21	1:55:A:VAL:HG22	20	1.98	2.26	0.94
(1,427)	1:20:A:THR:HG23	1:55:A:VAL:HG22	20	1.98	2.26	0.94
(1,893)	1:41:A:TRP:HZ3	1:61:A:SER:HB2	20	1.41	0.56	1.53
(1,893)	1:41:A:TRP:HZ3	1:32:A:VAL:HA	20	1.41	0.56	1.53
(1,193)	1:20:A:THR:H	1:20:A:THR:HG22	20	1.0	0.03	1.01
(1,193)	1:20:A:THR:H	1:20:A:THR:HG23	20	1.0	0.03	1.01
(1,193)	1:20:A:THR:H	1:20:A:THR:HG21	20	1.0	0.03	1.01
(1,906)	1:6:A:GLN:H	1:6:A:GLN:HG2	20	0.62	0.28	0.6
(1,906)	1:34:A:GLU:HG3	1:3:A:HIS:H	20	0.62	0.28	0.6
(1,906)	1:34:A:GLU:HG2	1:3:A:HIS:H	20	0.62	0.28	0.6
(1,714)	1:15:A:ASN:HD22	1:15:A:ASN:HA	20	0.58	0.37	0.48
(1,434)	1:24:A:VAL:HG11	1:30:A:ILE:HD12	20	0.55	0.27	0.5
(1,434)	1:24:A:VAL:HG13	1:30:A:ILE:HD12	20	0.55	0.27	0.5
(1,434)	1:24:A:VAL:HG12	1:30:A:ILE:HD13	20	0.55	0.27	0.5
(1,434)	1:24:A:VAL:HG12	1:30:A:ILE:HD11	20	0.55	0.27	0.5
(1,434)	1:24:A:VAL:HG11	1:30:A:ILE:HD13	20	0.55	0.27	0.5
(1,434)	1:24:A:VAL:HG11	1:30:A:ILE:HD11	20	0.55	0.27	0.5
(1,434)	1:24:A:VAL:HG13	1:30:A:ILE:HD13	20	0.55	0.27	0.5
(1,434)	1:24:A:VAL:HG12	1:30:A:ILE:HD12	20	0.55	0.27	0.5
(1,624)	1:18:A:ASP:H	1:17:A:TYR:HA	20	0.13	0.01	0.13
(1,849)	1:13:A:TYR:H	1:15:A:ASN:HD22	19	1.69	1.17	1.36
(1,849)	1:13:A:TYR:H	1:13:A:TYR:HE1	19	1.69	1.17	1.36
(1,66)	1:28:A:ASP:HB3	1:25:A:LYS:HG2	19	0.7	0.49	0.62
(1,66)	1:28:A:ASP:HB3	1:25:A:LYS:HG3	19	0.7	0.49	0.62
(1,589)	1:42:A:LYS:H	1:42:A:LYS:HB3	19	0.6	0.19	0.67
(1,670)	1:43:A:GLY:H	1:42:A:LYS:HB2	19	0.55	0.13	0.59
(1,39)	1:54:A:ARG:H	1:53:A:ASP:HB2	19	0.54	0.31	0.57
(1,39)	1:53:A:ASP:HB3	1:54:A:ARG:H	19	0.54	0.31	0.57
(1,908)	1:52:A:ASN:H	1:49:A:ARG:HB2	19	0.51	0.26	0.48
(1,908)	1:34:A:GLU:H	1:35:A:GLN:HG2	19	0.51	0.26	0.48
(1,908)	1:34:A:GLU:H	1:35:A:GLN:HG3	19	0.51	0.26	0.48
(1,908)	1:52:A:ASN:H	1:49:A:ARG:HB3	19	0.51	0.26	0.48
(1,743)	1:63:A:GLY:H	1:62:A:LEU:HG	19	0.47	0.3	0.36
(1,560)	1:64:A:GLU:H	1:64:A:GLU:HG2	19	0.47	0.34	0.35
(1,203)	1:52:A:ASN:HD22	1:52:A:ASN:H	19	0.37	0.31	0.28
(1,858)	1:13:A:TYR:HD2	1:23:A:ASN:HA	19	0.3	0.28	0.24
(1,858)	1:13:A:TYR:HD2	1:21:A:SER:HA	19	0.3	0.28	0.24
(1,299)	1:50:A:THR:H	1:47:A:ASP:HB2	18	0.74	0.59	0.42
(1,859)	1:67:A:VAL:H	1:7:A:VAL:HA	18	0.36	0.24	0.28

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Key	Atom-1	Atom-2	Models ¹	Mean (Å)	SD ¹ (Å)	Median (Å)
(1,859)	1:63:A:GLY:H	1:7:A:VAL:HA	18	0.36	0.24	0.28
(1,791)	1:39:A:GLY:H	1:39:A:GLY:HA3	18	0.25	0.03	0.26
(1,610)	1:58:A:PHE:H	1:57:A:TYR:HB2	17	1.09	0.08	1.06
(1,62)	1:18:A:ASP:HB2	1:18:A:ASP:H	17	0.69	0.35	0.58
(1,211)	1:25:A:LYS:H	1:11:A:LYS:H	17	0.39	0.25	0.36
(1,967)	1:62:A:LEU:H	1:63:A:GLY:HA2	17	0.36	0.24	0.33
(1,432)	1:22:A:LEU:HD11	1:24:A:VAL:HG21	17	0.29	0.19	0.22
(1,432)	1:22:A:LEU:HD11	1:24:A:VAL:HG23	17	0.29	0.19	0.22
(1,432)	1:22:A:LEU:HD11	1:24:A:VAL:HG22	17	0.29	0.19	0.22
(1,432)	1:22:A:LEU:HD13	1:24:A:VAL:HG22	17	0.29	0.19	0.22
(1,432)	1:22:A:LEU:HD12	1:24:A:VAL:HG21	17	0.29	0.19	0.22
(1,432)	1:22:A:LEU:HD12	1:24:A:VAL:HG23	17	0.29	0.19	0.22
(1,432)	1:22:A:LEU:HD12	1:24:A:VAL:HG22	17	0.29	0.19	0.22
(1,432)	1:22:A:LEU:HD13	1:24:A:VAL:HG21	17	0.29	0.19	0.22
(1,432)	1:22:A:LEU:HD13	1:24:A:VAL:HG23	17	0.29	0.19	0.22
(1,888)	1:5:A:LEU:HB3	1:67:A:VAL:HA	16	1.64	1.04	1.47
(1,360)	1:27:A:GLY:H	1:25:A:LYS:HG3	16	0.85	0.49	0.75
(1,360)	1:27:A:GLY:H	1:25:A:LYS:HG2	16	0.85	0.49	0.75
(1,648)	1:12:A:ASP:H	1:12:A:ASP:HB3	16	0.75	0.16	0.79
(1,587)	1:53:A:ASP:H	1:53:A:ASP:HB2	16	0.65	0.33	0.66
(1,392)	1:43:A:GLY:H	1:33:A:LEU:HD12	16	0.56	0.75	0.32
(1,392)	1:43:A:GLY:H	1:33:A:LEU:HD11	16	0.56	0.75	0.32
(1,392)	1:43:A:GLY:H	1:33:A:LEU:HD13	16	0.56	0.75	0.32
(1,953)	1:23:A:ASN:HB2	1:45:A:ILE:HG23	16	0.46	0.2	0.42
(1,953)	1:23:A:ASN:HB3	1:45:A:ILE:HG21	16	0.46	0.2	0.42
(1,953)	1:23:A:ASN:HB3	1:45:A:ILE:HG22	16	0.46	0.2	0.42
(1,953)	1:23:A:ASN:HB2	1:45:A:ILE:HG21	16	0.46	0.2	0.42
(1,953)	1:23:A:ASN:HB2	1:45:A:ILE:HG22	16	0.46	0.2	0.42
(1,953)	1:23:A:ASN:HB3	1:45:A:ILE:HG23	16	0.46	0.2	0.42
(1,525)	1:15:A:ASN:H	1:14:A:CYS:HA	16	0.42	0.04	0.43
(1,815)	1:44:A:CYS:H	1:30:A:ILE:HG21	16	0.25	0.08	0.28
(1,815)	1:44:A:CYS:H	1:30:A:ILE:HG22	16	0.25	0.08	0.28
(1,815)	1:44:A:CYS:H	1:30:A:ILE:HG23	16	0.25	0.08	0.28
(1,367)	1:9:A:ALA:HB2	1:10:A:THR:H	16	0.19	0.08	0.16
(1,367)	1:9:A:ALA:HB3	1:10:A:THR:H	16	0.19	0.08	0.16
(1,367)	1:9:A:ALA:HB1	1:10:A:THR:H	16	0.19	0.08	0.16
(1,458)	1:9:A:ALA:H	1:9:A:ALA:HB3	16	0.15	0.04	0.14
(1,458)	1:9:A:ALA:H	1:9:A:ALA:HB1	16	0.15	0.04	0.14
(1,458)	1:9:A:ALA:H	1:9:A:ALA:HB2	16	0.15	0.04	0.14
(1,737)	1:50:A:THR:H	1:50:A:THR:HG22	15	0.89	0.07	0.86
(1,737)	1:50:A:THR:H	1:50:A:THR:HG23	15	0.89	0.07	0.86
(1,737)	1:50:A:THR:H	1:50:A:THR:HG21	15	0.89	0.07	0.86

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Key	Atom-1	Atom-2	Models ¹	Mean (Å)	SD ¹ (Å)	Median (Å)
(1,300)	1:4:A:MET:H	1:3:A:HIS:HB3	15	0.86	0.39	0.94
(1,300)	1:4:A:MET:H	1:3:A:HIS:HB2	15	0.86	0.39	0.94
(1,707)	1:35:A:GLN:HE22	1:35:A:GLN:HB3	15	0.7	0.28	0.6
(1,734)	1:39:A:GLY:H	1:40:A:ARG:HB2	15	0.58	0.31	0.62
(1,58)	1:28:A:ASP:H	1:28:A:ASP:HB2	15	0.39	0.29	0.29
(1,515)	1:14:A:CYS:H	1:13:A:TYR:HB3	15	0.35	0.17	0.32
(1,29)	1:43:A:GLY:HA3	1:30:A:ILE:HG21	15	0.28	0.1	0.29
(1,29)	1:43:A:GLY:HA3	1:30:A:ILE:HG22	15	0.28	0.1	0.29
(1,29)	1:43:A:GLY:HA3	1:30:A:ILE:HG23	15	0.28	0.1	0.29
(1,924)	1:7:A:VAL:HG11	1:65:A:ALA:HB2	15	0.2	0.11	0.16
(1,924)	1:65:A:ALA:HB1	1:5:A:LEU:HD13	15	0.2	0.11	0.16
(1,924)	1:65:A:ALA:HB1	1:5:A:LEU:HD11	15	0.2	0.11	0.16
(1,924)	1:65:A:ALA:HB3	1:5:A:LEU:HD12	15	0.2	0.11	0.16
(1,924)	1:65:A:ALA:HB2	1:5:A:LEU:HD11	15	0.2	0.11	0.16
(1,924)	1:65:A:ALA:HB3	1:5:A:LEU:HD13	15	0.2	0.11	0.16
(1,924)	1:65:A:ALA:HB1	1:5:A:LEU:HD12	15	0.2	0.11	0.16
(1,924)	1:65:A:ALA:HB2	1:5:A:LEU:HD13	15	0.2	0.11	0.16
(1,924)	1:7:A:VAL:HG11	1:65:A:ALA:HB1	15	0.2	0.11	0.16
(1,924)	1:7:A:VAL:HG13	1:65:A:ALA:HB1	15	0.2	0.11	0.16
(1,57)	1:28:A:ASP:HB2	1:25:A:LYS:HG2	14	1.01	0.76	0.74
(1,57)	1:28:A:ASP:HB2	1:25:A:LYS:HG3	14	1.01	0.76	0.74
(1,628)	1:38:A:ASP:H	1:36:A:HIS:HB2	14	0.84	0.75	0.44
(1,793)	1:28:A:ASP:HB3	1:24:A:VAL:HG12	14	0.76	0.7	0.48
(1,793)	1:28:A:ASP:HB3	1:45:A:ILE:HG21	14	0.76	0.7	0.48
(1,793)	1:28:A:ASP:HB3	1:45:A:ILE:HG22	14	0.76	0.7	0.48
(1,793)	1:28:A:ASP:HB3	1:45:A:ILE:HG23	14	0.76	0.7	0.48
(1,793)	1:28:A:ASP:HB3	1:24:A:VAL:HG11	14	0.76	0.7	0.48
(1,635)	1:52:A:ASN:H	1:52:A:ASN:HB2	14	0.63	0.06	0.64
(1,819)	1:58:A:PHE:HD1	1:42:A:LYS:H	14	0.32	0.16	0.31
(1,881)	1:13:A:TYR:HD2	1:21:A:SER:HB3	13	1.0	0.8	0.62
(1,881)	1:13:A:TYR:HD2	1:21:A:SER:HB2	13	1.0	0.8	0.62
(1,881)	1:13:A:TYR:HD1	1:21:A:SER:HB2	13	1.0	0.8	0.62
(1,899)	1:27:A:GLY:H	1:28:A:ASP:HB2	13	0.74	0.22	0.8
(1,899)	1:15:A:ASN:H	1:14:A:CYS:HB3	13	0.74	0.22	0.8
(1,228)	1:51:A:GLY:H	1:52:A:ASN:HD22	13	0.66	0.39	0.64
(1,873)	1:41:A:TRP:HB3	1:34:A:GLU:HA	13	0.6	0.35	0.62
(1,190)	1:34:A:GLU:HG2	1:3:A:HIS:H	13	0.58	0.35	0.55
(1,190)	1:34:A:GLU:HG3	1:3:A:HIS:H	13	0.58	0.35	0.55
(1,61)	1:18:A:ASP:HB2	1:18:A:ASP:HA	13	0.52	0.01	0.52
(1,937)	1:10:A:THR:HG22	1:10:A:THR:H	13	0.29	0.17	0.21
(1,937)	1:10:A:THR:HG23	1:10:A:THR:H	13	0.29	0.17	0.21
(1,937)	1:22:A:LEU:HB3	1:22:A:LEU:H	13	0.29	0.17	0.21

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Key	Atom-1	Atom-2	Models ¹	Mean (Å)	SD ¹ (Å)	Median (Å)
(1,937)	1:10:A:THR:HG21	1:10:A:THR:H	13	0.29	0.17	0.21
(1,232)	1:32:A:VAL:HG12	1:41:A:TRP:HZ3	13	0.23	0.09	0.24
(1,232)	1:32:A:VAL:HG13	1:41:A:TRP:HZ3	13	0.23	0.09	0.24
(1,232)	1:32:A:VAL:HG11	1:41:A:TRP:HZ3	13	0.23	0.09	0.24
(1,234)	1:8:A:ARG:HG3	1:6:A:GLN:HE22	12	1.0	0.9	0.52
(1,853)	1:29:A:ILE:HG12	1:48:A:ASN:HD22	12	0.9	0.65	0.92
(1,853)	1:25:A:LYS:HB2	1:48:A:ASN:HD22	12	0.9	0.65	0.92
(1,391)	1:34:A:GLU:H	1:5:A:LEU:HD21	12	0.83	0.85	0.56
(1,391)	1:34:A:GLU:H	1:5:A:LEU:HD23	12	0.83	0.85	0.56
(1,391)	1:34:A:GLU:H	1:5:A:LEU:HD22	12	0.83	0.85	0.56
(1,317)	1:39:A:GLY:H	1:38:A:ASP:HB3	12	0.58	0.2	0.56
(1,182)	1:25:A:LYS:H	1:27:A:GLY:H	12	0.48	0.17	0.46
(1,14)	1:27:A:GLY:H	1:26:A:ALA:HA	12	0.44	0.2	0.41
(1,69)	1:52:A:ASN:H	1:52:A:ASN:HB2	12	0.35	0.06	0.36
(1,952)	1:23:A:ASN:H	1:45:A:ILE:HD12	12	0.3	0.14	0.28
(1,952)	1:23:A:ASN:H	1:45:A:ILE:HD13	12	0.3	0.14	0.28
(1,952)	1:23:A:ASN:H	1:45:A:ILE:HD11	12	0.3	0.14	0.28
(1,106)	1:41:A:TRP:HB3	1:41:A:TRP:HZ3	12	0.22	0.06	0.2
(1,669)	1:43:A:GLY:H	1:55:A:VAL:HG23	12	0.21	0.1	0.18
(1,669)	1:43:A:GLY:H	1:55:A:VAL:HG21	12	0.21	0.1	0.18
(1,669)	1:43:A:GLY:H	1:55:A:VAL:HG22	12	0.21	0.1	0.18
(1,830)	1:9:A:ALA:H	1:29:A:ILE:H	12	0.2	0.09	0.16
(1,830)	1:23:A:ASN:H	1:45:A:ILE:H	12	0.2	0.09	0.16
(1,225)	1:35:A:GLN:HG2	1:41:A:TRP:HD1	11	0.85	0.65	0.51
(1,225)	1:35:A:GLN:HG3	1:41:A:TRP:HD1	11	0.85	0.65	0.51
(1,74)	1:15:A:ASN:HD22	1:15:A:ASN:HB2	11	0.37	0.11	0.42
(1,252)	1:15:A:ASN:H	1:15:A:ASN:HA	11	0.15	0.02	0.16
(1,346)	1:55:A:VAL:H	1:54:A:ARG:HB3	10	0.84	0.38	1.0
(1,226)	1:39:A:GLY:HA3	1:41:A:TRP:HD1	10	0.41	0.21	0.36
(1,176)	1:33:A:LEU:H	1:42:A:LYS:H	10	0.39	0.13	0.38
(1,55)	1:47:A:ASP:HB3	1:52:A:ASN:H	10	0.31	0.09	0.3
(1,453)	1:32:A:VAL:H	1:32:A:VAL:HG13	10	0.13	0.02	0.12
(1,453)	1:32:A:VAL:H	1:32:A:VAL:HG12	10	0.13	0.02	0.12
(1,453)	1:32:A:VAL:H	1:32:A:VAL:HG11	10	0.13	0.02	0.12
(1,901)	1:14:A:CYS:HB2	1:13:A:TYR:HD2	9	1.72	0.82	2.08
(1,901)	1:14:A:CYS:HB3	1:13:A:TYR:HD2	9	1.72	0.82	2.08
(1,901)	1:14:A:CYS:HB2	1:13:A:TYR:HD1	9	1.72	0.82	2.08
(1,620)	1:21:A:SER:H	1:14:A:CYS:HB2	9	1.7	1.38	1.06
(1,318)	1:44:A:CYS:HB3	1:45:A:ILE:H	9	0.6	0.11	0.61
(1,554)	1:54:A:ARG:H	1:54:A:ARG:HB2	9	0.52	0.29	0.46
(1,965)	1:26:A:ALA:H	1:12:A:ASP:HA	9	0.41	0.29	0.38
(1,660)	1:10:A:THR:H	1:11:A:LYS:HB2	9	0.35	0.5	0.17

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Key	Atom-1	Atom-2	Models ¹	Mean (Å)	SD ¹ (Å)	Median (Å)
(1,414)	1:5:A:LEU:HD12	1:32:A:VAL:HG12	9	0.34	0.31	0.25
(1,414)	1:5:A:LEU:HD12	1:32:A:VAL:HG11	9	0.34	0.31	0.25
(1,414)	1:5:A:LEU:HD13	1:32:A:VAL:HG13	9	0.34	0.31	0.25
(1,414)	1:5:A:LEU:HD13	1:32:A:VAL:HG12	9	0.34	0.31	0.25
(1,414)	1:5:A:LEU:HD12	1:32:A:VAL:HG13	9	0.34	0.31	0.25
(1,414)	1:5:A:LEU:HD11	1:32:A:VAL:HG13	9	0.34	0.31	0.25
(1,896)	1:48:A:ASN:H	1:47:A:ASP:HB2	9	0.32	0.17	0.22
(1,896)	1:48:A:ASN:H	1:25:A:LYS:HE3	9	0.32	0.17	0.22
(1,738)	1:50:A:THR:H	1:50:A:THR:HB	9	0.27	0.11	0.27
(1,305)	1:13:A:TYR:HD1	1:11:A:LYS:HE3	8	1.44	0.81	1.46
(1,305)	1:13:A:TYR:HD1	1:11:A:LYS:HE2	8	1.44	0.81	1.46
(1,305)	1:13:A:TYR:HD2	1:11:A:LYS:HE3	8	1.44	0.81	1.46
(1,931)	1:6:A:GLN:HE21	1:8:A:ARG:HG2	8	0.58	0.37	0.55
(1,931)	1:41:A:TRP:HZ2	1:60:A:SER:HB3	8	0.58	0.37	0.55
(1,775)	1:38:A:ASP:H	1:40:A:ARG:H	8	0.56	0.32	0.55
(1,929)	1:13:A:TYR:HD1	1:11:A:LYS:HG2	8	0.5	0.21	0.51
(1,929)	1:13:A:TYR:HD1	1:11:A:LYS:HG3	8	0.5	0.21	0.51
(1,929)	1:13:A:TYR:HD2	1:11:A:LYS:HG2	8	0.5	0.21	0.51
(1,56)	1:28:A:ASP:H	1:28:A:ASP:HB3	8	0.48	0.17	0.52
(1,664)	1:61:A:SER:H	1:61:A:SER:HB3	8	0.4	0.1	0.43
(1,598)	1:29:A:ILE:H	1:28:A:ASP:HA	8	0.19	0.04	0.18
(1,482)	1:46:A:HIS:H	1:30:A:ILE:HG13	8	0.18	0.05	0.18
(1,426)	1:20:A:THR:HG23	1:55:A:VAL:HG12	7	4.14	2.27	4.3
(1,426)	1:20:A:THR:HG22	1:55:A:VAL:HG11	7	4.14	2.27	4.3
(1,426)	1:20:A:THR:HG22	1:55:A:VAL:HG13	7	4.14	2.27	4.3
(1,426)	1:20:A:THR:HG21	1:55:A:VAL:HG12	7	4.14	2.27	4.3
(1,426)	1:20:A:THR:HG23	1:55:A:VAL:HG11	7	4.14	2.27	4.3
(1,168)	1:13:A:TYR:HD2	1:22:A:LEU:H	7	1.41	0.51	1.27
(1,249)	1:13:A:TYR:HE2	1:21:A:SER:HA	7	0.96	0.94	0.46
(1,249)	1:13:A:TYR:HE1	1:21:A:SER:HA	7	0.96	0.94	0.46
(1,324)	1:4:A:MET:H	1:4:A:MET:HG3	7	0.96	0.26	1.05
(1,358)	1:41:A:TRP:H	1:40:A:ARG:HB2	7	0.5	0.3	0.3
(1,358)	1:41:A:TRP:H	1:40:A:ARG:HB3	7	0.5	0.3	0.3
(1,872)	1:5:A:LEU:HD21	1:33:A:LEU:HA	7	0.4	0.34	0.21
(1,872)	1:5:A:LEU:HD21	1:4:A:MET:HA	7	0.4	0.34	0.21
(1,872)	1:5:A:LEU:HD22	1:33:A:LEU:HA	7	0.4	0.34	0.21
(1,872)	1:33:A:LEU:HD12	1:33:A:LEU:HA	7	0.4	0.34	0.21
(1,825)	1:34:A:GLU:HB2	1:3:A:HIS:H	7	0.24	0.1	0.23
(1,825)	1:19:A:LEU:HB3	1:20:A:THR:H	7	0.24	0.1	0.23
(1,450)	1:25:A:LYS:H	1:24:A:VAL:HG12	7	0.21	0.05	0.23
(1,450)	1:25:A:LYS:H	1:24:A:VAL:HG13	7	0.21	0.05	0.23
(1,151)	1:20:A:THR:HA	1:56:A:GLY:HA2	6	2.76	0.8	2.58

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Key	Atom-1	Atom-2	Models ¹	Mean (Å)	SD ¹ (Å)	Median (Å)
(1,761)	1:15:A:ASN:H	1:22:A:LEU:H	6	2.32	0.84	2.18
(1,796)	1:18:A:ASP:H	1:17:A:TYR:HB2	6	1.35	0.19	1.37
(1,545)	1:40:A:ARG:H	1:36:A:HIS:HB2	6	0.85	0.85	0.38
(1,390)	1:24:A:VAL:HG21	1:13:A:TYR:HB3	6	0.71	0.77	0.41
(1,390)	1:24:A:VAL:HG23	1:13:A:TYR:HB3	6	0.71	0.77	0.41
(1,390)	1:24:A:VAL:HG22	1:13:A:TYR:HB3	6	0.71	0.77	0.41
(1,921)	1:22:A:LEU:H	1:22:A:LEU:HB2	6	0.7	0.04	0.69
(1,705)	1:35:A:GLN:HE22	1:33:A:LEU:HD23	6	0.68	0.79	0.3
(1,705)	1:35:A:GLN:HE22	1:33:A:LEU:HD21	6	0.68	0.79	0.3
(1,705)	1:35:A:GLN:HE22	1:33:A:LEU:HD22	6	0.68	0.79	0.3
(1,369)	1:29:A:ILE:HG13	1:6:A:GLN:HE22	6	0.52	0.29	0.54
(1,43)	1:13:A:TYR:HB2	1:24:A:VAL:HG21	6	0.34	0.29	0.24
(1,43)	1:13:A:TYR:HB2	1:24:A:VAL:HG22	6	0.34	0.29	0.24
(1,43)	1:13:A:TYR:HB2	1:24:A:VAL:HG23	6	0.34	0.29	0.24
(1,364)	1:29:A:ILE:HD13	1:8:A:ARG:HG2	5	1.95	0.96	1.48
(1,364)	1:29:A:ILE:HD12	1:8:A:ARG:HG2	5	1.95	0.96	1.48
(1,851)	1:13:A:TYR:HE2	1:57:A:TYR:H	5	1.75	1.05	1.08
(1,851)	1:13:A:TYR:HE1	1:57:A:TYR:H	5	1.75	1.05	1.08
(1,347)	1:63:A:GLY:H	1:62:A:LEU:HB3	5	0.85	0.51	0.85
(1,196)	1:34:A:GLU:HB3	1:4:A:MET:H	5	0.8	0.53	0.64
(1,543)	1:40:A:ARG:H	1:40:A:ARG:HG2	5	0.8	0.21	0.83
(1,519)	1:14:A:CYS:H	1:13:A:TYR:HD1	5	0.77	0.48	0.68
(1,822)	1:36:A:HIS:HB3	1:38:A:ASP:H	5	0.72	0.32	0.75
(1,822)	1:54:A:ARG:HD3	1:55:A:VAL:H	5	0.72	0.32	0.75
(1,822)	1:26:A:ALA:HA	1:28:A:ASP:H	5	0.72	0.32	0.75
(1,389)	1:11:A:LYS:HB3	1:24:A:VAL:HG12	5	0.63	0.42	0.48
(1,389)	1:11:A:LYS:HB3	1:24:A:VAL:HG11	5	0.63	0.42	0.48
(1,902)	1:28:A:ASP:H	1:28:A:ASP:HB2	5	0.46	0.32	0.27
(1,902)	1:38:A:ASP:H	1:38:A:ASP:HB3	5	0.46	0.32	0.27
(1,673)	1:34:A:GLU:H	1:33:A:LEU:HB2	5	0.39	0.5	0.14
(1,394)	1:41:A:TRP:HZ3	1:5:A:LEU:HD23	5	0.37	0.21	0.24
(1,394)	1:41:A:TRP:HZ3	1:5:A:LEU:HD21	5	0.37	0.21	0.24
(1,574)	1:36:A:HIS:H	1:35:A:GLN:HB2	5	0.33	0.16	0.27
(1,461)	1:67:A:VAL:H	1:67:A:VAL:HA	5	0.26	0.05	0.28
(1,653)	1:17:A:TYR:H	1:16:A:ASN:HB2	5	0.18	0.07	0.15
(1,334)	1:49:A:ARG:H	1:49:A:ARG:HB3	5	0.17	0.06	0.17
(1,334)	1:49:A:ARG:H	1:49:A:ARG:HB2	5	0.17	0.06	0.17
(1,590)	1:42:A:LYS:H	1:41:A:TRP:HB2	5	0.14	0.04	0.11
(1,368)	1:8:A:ARG:HD2	1:8:A:ARG:HG2	5	0.1	0.0	0.1
(1,724)	1:20:A:THR:H	1:14:A:CYS:HB2	4	1.89	1.4	1.52
(1,281)	1:6:A:GLN:H	1:67:A:VAL:HA	4	1.8	0.28	1.7
(1,891)	1:55:A:VAL:H	1:56:A:GLY:HA2	4	1.76	0.11	1.74

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Key	Atom-1	Atom-2	Models ¹	Mean (Å)	SD ¹ (Å)	Median (Å)
(1,891)	1:6:A:GLN:H	1:67:A:VAL:HA	4	1.76	0.11	1.74
(1,720)	1:51:A:GLY:H	1:50:A:THR:H	4	0.98	0.51	1.24
(1,733)	1:39:A:GLY:H	1:36:A:HIS:HB3	4	0.9	0.45	0.97
(1,464)	1:67:A:VAL:H	1:66:A:ILE:HG12	4	0.61	0.16	0.58
(1,50)	1:47:A:ASP:H	1:47:A:ASP:HB2	4	0.52	0.02	0.51
(1,727)	1:39:A:GLY:H	1:40:A:ARG:HG2	4	0.27	0.08	0.3
(1,371)	1:28:A:ASP:H	1:10:A:THR:HG22	4	0.26	0.11	0.26
(1,371)	1:28:A:ASP:H	1:10:A:THR:HG21	4	0.26	0.11	0.26
(1,371)	1:28:A:ASP:H	1:10:A:THR:HG23	4	0.26	0.11	0.26
(1,512)	1:65:A:ALA:H	1:66:A:ILE:HG22	4	0.26	0.09	0.24
(1,512)	1:65:A:ALA:H	1:66:A:ILE:HG21	4	0.26	0.09	0.24
(1,691)	1:16:A:ASN:HD21	1:16:A:ASN:HA	4	0.22	0.04	0.22
(1,709)	1:35:A:GLN:HE21	1:35:A:GLN:HB3	4	0.12	0.02	0.12
(1,323)	1:15:A:ASN:H	1:15:A:ASN:HB2	4	0.12	0.01	0.13
(1,524)	1:15:A:ASN:H	1:15:A:ASN:HB3	4	0.12	0.0	0.12
(1,283)	1:13:A:TYR:HE2	1:56:A:GLY:HA3	3	1.22	0.61	1.38
(1,801)	1:65:A:ALA:HB1	1:5:A:LEU:HB3	3	1.21	0.74	1.39
(1,801)	1:65:A:ALA:HB2	1:5:A:LEU:HB3	3	1.21	0.74	1.39
(1,801)	1:65:A:ALA:HB3	1:5:A:LEU:HD12	3	1.21	0.74	1.39
(1,490)	1:41:A:TRP:HE1	1:33:A:LEU:HD21	3	1.16	0.73	1.36
(1,490)	1:41:A:TRP:HE1	1:33:A:LEU:HD22	3	1.16	0.73	1.36
(1,192)	1:67:A:VAL:H	1:6:A:GLN:H	3	1.13	0.98	0.51
(1,212)	1:52:A:ASN:HD21	1:52:A:ASN:H	3	1.09	0.39	1.34
(1,340)	1:63:A:GLY:H	1:7:A:VAL:HB	3	1.06	0.79	0.99
(1,418)	1:7:A:VAL:HG11	1:32:A:VAL:HG21	3	0.99	1.2	0.18
(1,418)	1:7:A:VAL:HG12	1:32:A:VAL:HG21	3	0.99	1.2	0.18
(1,572)	1:66:A:ILE:H	1:66:A:ILE:HG12	3	0.76	0.13	0.69
(1,605)	1:57:A:TYR:H	1:21:A:SER:HA	3	0.57	0.32	0.52
(1,341)	1:29:A:ILE:H	1:29:A:ILE:HG12	3	0.45	0.48	0.11
(1,597)	1:62:A:LEU:H	1:62:A:LEU:HB3	3	0.43	0.32	0.31
(1,735)	1:39:A:GLY:H	1:41:A:TRP:HD1	3	0.39	0.18	0.51
(1,866)	1:5:A:LEU:HD11	1:65:A:ALA:HA	3	0.37	0.16	0.38
(1,866)	1:66:A:ILE:HG21	1:65:A:ALA:HA	3	0.37	0.16	0.38
(1,866)	1:66:A:ILE:HG22	1:65:A:ALA:HA	3	0.37	0.16	0.38
(1,903)	1:28:A:ASP:HB2	1:25:A:LYS:HG3	3	0.31	0.09	0.37
(1,903)	1:60:A:SER:HB3	1:60:A:SER:HA	3	0.31	0.09	0.37
(1,446)	1:33:A:LEU:HD22	1:55:A:VAL:HG23	3	0.3	0.15	0.2
(1,446)	1:33:A:LEU:HD21	1:55:A:VAL:HG22	3	0.3	0.15	0.2
(1,595)	1:62:A:LEU:H	1:62:A:LEU:HB2	3	0.23	0.09	0.23
(1,435)	1:24:A:VAL:HG13	1:30:A:ILE:HG23	3	0.21	0.12	0.13
(1,435)	1:24:A:VAL:HG12	1:30:A:ILE:HG21	3	0.21	0.12	0.13
(1,435)	1:24:A:VAL:HG12	1:30:A:ILE:HG23	3	0.21	0.12	0.13

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Key	Atom-1	Atom-2	Models ¹	Mean (Å)	SD ¹ (Å)	Median (Å)
(1,876)	1:45:A:ILE:HD13	1:30:A:ILE:HA	3	0.21	0.03	0.23
(1,876)	1:45:A:ILE:HD11	1:30:A:ILE:HA	3	0.21	0.03	0.23
(1,593)	1:62:A:LEU:H	1:60:A:SER:HA	3	0.2	0.05	0.22
(1,386)	1:46:A:HIS:H	1:45:A:ILE:HG21	3	0.19	0.09	0.13
(1,386)	1:46:A:HIS:H	1:45:A:ILE:HG22	3	0.19	0.09	0.13
(1,836)	1:40:A:ARG:H	1:41:A:TRP:HD1	3	0.19	0.07	0.17
(1,774)	1:42:A:LYS:H	1:34:A:GLU:H	3	0.18	0.05	0.19
(1,567)	1:8:A:ARG:H	1:8:A:ARG:HD3	3	0.17	0.04	0.17
(1,855)	1:14:A:CYS:HB3	1:23:A:ASN:HD22	3	0.15	0.03	0.14
(1,855)	1:52:A:ASN:HB2	1:23:A:ASN:HD22	3	0.15	0.03	0.14
(1,97)	1:6:A:GLN:HB2	1:6:A:GLN:HG3	3	0.1	0.0	0.1
(1,883)	1:13:A:TYR:HE1	1:21:A:SER:HB2	2	1.84	1.0	1.84
(1,883)	1:13:A:TYR:HE2	1:21:A:SER:HB3	2	1.84	1.0	1.84
(1,169)	1:13:A:TYR:HE2	1:22:A:LEU:H	2	1.66	0.29	1.66
(1,579)	1:36:A:HIS:H	1:33:A:LEU:HD23	2	1.41	0.14	1.41
(1,579)	1:36:A:HIS:H	1:33:A:LEU:HD21	2	1.41	0.14	1.41
(1,397)	1:33:A:LEU:H	1:33:A:LEU:HD12	2	1.34	0.5	1.34
(1,397)	1:33:A:LEU:H	1:33:A:LEU:HD11	2	1.34	0.5	1.34
(1,387)	1:63:A:GLY:H	1:7:A:VAL:HG23	2	1.29	0.58	1.29
(1,387)	1:63:A:GLY:H	1:7:A:VAL:HG21	2	1.29	0.58	1.29
(1,328)	1:24:A:VAL:HB	1:26:A:ALA:H	2	1.27	0.56	1.27
(1,122)	1:12:A:ASP:H	1:26:A:ALA:HB2	2	1.15	0.85	1.15
(1,122)	1:12:A:ASP:H	1:26:A:ALA:HB3	2	1.15	0.85	1.15
(1,44)	1:13:A:TYR:HB3	1:24:A:VAL:HB	2	1.14	0.76	1.14
(1,396)	1:44:A:CYS:H	1:33:A:LEU:HD12	2	1.01	0.89	1.01
(1,396)	1:44:A:CYS:H	1:33:A:LEU:HD11	2	1.01	0.89	1.01
(1,113)	1:31:A:THR:HB	1:44:A:CYS:HB2	2	0.83	0.14	0.83
(1,210)	1:41:A:TRP:HZ3	1:61:A:SER:H	2	0.82	0.09	0.82
(1,838)	1:36:A:HIS:HB2	1:39:A:GLY:H	2	0.8	0.14	0.8
(1,45)	1:47:A:ASP:HB3	1:52:A:ASN:HA	2	0.75	0.28	0.75
(1,227)	1:29:A:ILE:H	1:48:A:ASN:HD22	2	0.74	0.6	0.74
(1,17)	1:26:A:ALA:HA	1:24:A:VAL:HG13	2	0.7	0.53	0.7
(1,17)	1:26:A:ALA:HA	1:24:A:VAL:HG12	2	0.7	0.53	0.7
(1,582)	1:13:A:TYR:H	1:13:A:TYR:HB3	2	0.66	0.02	0.66
(1,124)	1:26:A:ALA:HB2	1:11:A:LYS:H	2	0.65	0.54	0.65
(1,124)	1:26:A:ALA:HB3	1:11:A:LYS:H	2	0.65	0.54	0.65
(1,575)	1:36:A:HIS:H	1:41:A:TRP:HA	2	0.64	0.02	0.64
(1,152)	1:20:A:THR:HA	1:56:A:GLY:HA3	2	0.62	0.45	0.62
(1,25)	1:43:A:GLY:HA2	1:33:A:LEU:HD12	2	0.57	0.42	0.57
(1,25)	1:43:A:GLY:HA2	1:33:A:LEU:HD11	2	0.57	0.42	0.57
(1,373)	1:41:A:TRP:HE1	1:35:A:GLN:HB3	2	0.57	0.09	0.57
(1,373)	1:41:A:TRP:HE1	1:35:A:GLN:HB2	2	0.57	0.09	0.57

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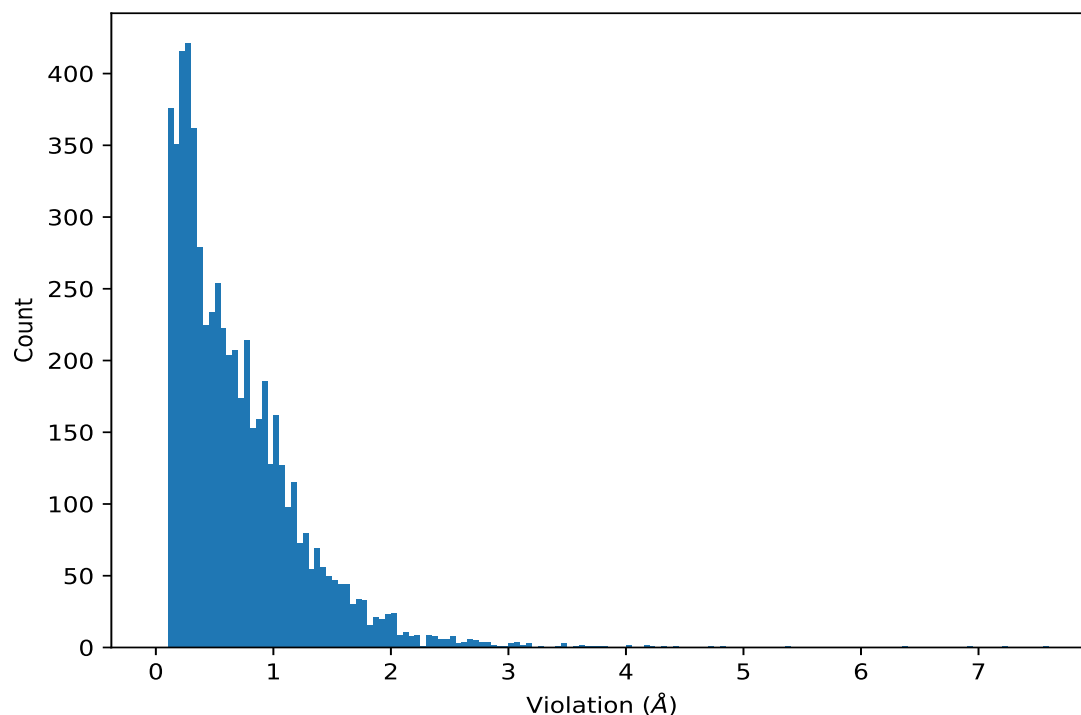
Key	Atom-1	Atom-2	Models ¹	Mean (Å)	SD ¹ (Å)	Median (Å)
(1,722)	1:51:A:GLY:H	1:49:A:ARG:HB2	2	0.46	0.05	0.46
(1,116)	1:35:A:GLN:HB3	1:36:A:HIS:H	2	0.43	0.24	0.43
(1,116)	1:36:A:HIS:H	1:35:A:GLN:HB2	2	0.43	0.24	0.43
(1,241)	1:4:A:MET:H	1:2:A:SER:HA	2	0.42	0.18	0.42
(1,303)	1:25:A:LYS:HE2	1:48:A:ASN:HD22	2	0.42	0.09	0.42
(1,135)	1:8:A:ARG:HA	1:27:A:GLY:HA3	2	0.4	0.24	0.4
(1,200)	1:14:A:CYS:H	1:13:A:TYR:HD1	2	0.4	0.1	0.4
(1,200)	1:13:A:TYR:HD2	1:14:A:CYS:H	2	0.4	0.1	0.4
(1,584)	1:3:A:HIS:H	1:34:A:GLU:HB3	2	0.38	0.14	0.38
(1,694)	1:52:A:ASN:HD22	1:50:A:THR:HB	2	0.37	0.03	0.37
(1,272)	1:17:A:TYR:HD2	1:16:A:ASN:HA	2	0.36	0.1	0.36
(1,820)	1:15:A:ASN:H	1:15:A:ASN:HB2	2	0.34	0.14	0.34
(1,65)	1:28:A:ASP:HB3	1:25:A:LYS:H	2	0.32	0.16	0.32
(1,534)	1:23:A:ASN:H	1:23:A:ASN:HB2	2	0.32	0.06	0.32
(1,897)	1:41:A:TRP:HB2	1:41:A:TRP:HE3	2	0.3	0.11	0.3
(1,314)	1:60:A:SER:HB2	1:60:A:SER:HA	2	0.3	0.01	0.3
(1,655)	1:18:A:ASP:H	1:17:A:TYR:H	2	0.29	0.05	0.29
(1,41)	1:13:A:TYR:HB2	1:24:A:VAL:HB	2	0.25	0.13	0.25
(1,549)	1:54:A:ARG:H	1:54:A:ARG:HD2	2	0.25	0.0	0.25
(1,20)	1:11:A:LYS:H	1:26:A:ALA:HA	2	0.23	0.11	0.23
(1,905)	1:53:A:ASP:H	1:52:A:ASN:HB3	2	0.2	0.0	0.2
(1,905)	1:23:A:ASN:HB3	1:23:A:ASN:H	2	0.2	0.0	0.2
(1,222)	1:22:A:LEU:HB2	1:13:A:TYR:HE2	2	0.19	0.01	0.19
(1,49)	1:13:A:TYR:HB2	1:13:A:TYR:HA	2	0.18	0.01	0.18
(1,171)	1:31:A:THR:HG23	1:32:A:VAL:H	2	0.18	0.04	0.18
(1,171)	1:31:A:THR:HG22	1:32:A:VAL:H	2	0.18	0.04	0.18
(1,755)	1:10:A:THR:H	1:63:A:GLY:H	2	0.18	0.04	0.18
(1,295)	1:41:A:TRP:HD1	1:36:A:HIS:HB2	2	0.16	0.02	0.16
(1,348)	1:41:A:TRP:HZ3	1:34:A:GLU:HB2	2	0.16	0.06	0.16
(1,527)	1:26:A:ALA:H	1:25:A:LYS:HA	2	0.16	0.03	0.16
(1,244)	1:30:A:ILE:HG12	1:8:A:ARG:HA	2	0.14	0.02	0.14
(1,552)	1:54:A:ARG:H	1:45:A:ILE:HG22	2	0.14	0.01	0.14
(1,552)	1:54:A:ARG:H	1:45:A:ILE:HG23	2	0.14	0.01	0.14
(1,137)	1:8:A:ARG:HA	1:29:A:ILE:HA	2	0.12	0.02	0.12
(1,792)	1:47:A:ASP:HB3	1:46:A:HIS:HA	2	0.12	0.01	0.12
(1,566)	1:8:A:ARG:H	1:7:A:VAL:HB	2	0.12	0.01	0.12

¹Number of violated models, ²Standard 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)
(1,426)	1:20:A:THR:HG21	1:55:A:VAL:HG12	22	7.57
(1,427)	1:20:A:THR:HG21	1:55:A:VAL:HG23	22	7.22
(1,427)	1:20:A:THR:HG22	1:55:A:VAL:HG21	19	6.9
(1,426)	1:20:A:THR:HG22	1:55:A:VAL:HG13	19	6.38
(1,427)	1:20:A:THR:HG23	1:55:A:VAL:HG21	8	5.37
(1,620)	1:21:A:SER:H	1:14:A:CYS:HB2	24	4.83
(1,426)	1:20:A:THR:HG23	1:55:A:VAL:HG12	8	4.7
(1,427)	1:20:A:THR:HG22	1:55:A:VAL:HG22	25	4.43
(1,426)	1:20:A:THR:HG22	1:55:A:VAL:HG13	25	4.3
(1,46)	1:13:A:TYR:HB2	1:11:A:LYS:HD2	25	4.2

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,427)	1:20:A:THR:HG22	1:55:A:VAL:HG22	17	4.17
(1,724)	1:20:A:THR:H	1:14:A:CYS:HB2	24	4.16
(1,151)	1:20:A:THR:HA	1:56:A:GLY:HA2	22	4.02
(1,27)	1:43:A:GLY:HA3	1:33:A:LEU:HD11	7	4.0
(1,761)	1:15:A:ASN:H	1:22:A:LEU:H	30	3.8
(1,364)	1:29:A:ILE:HD13	1:8:A:ARG:HG2	5	3.78
(1,415)	1:5:A:LEU:HD11	1:7:A:VAL:HG22	29	3.73
(1,802)	1:26:A:ALA:HB3	1:12:A:ASP:HA	4	3.67
(1,847)	1:36:A:HIS:HB2	1:41:A:TRP:HZ2	4	3.64
(1,426)	1:20:A:THR:HG22	1:55:A:VAL:HG11	17	3.63
(1,46)	1:13:A:TYR:HB2	1:11:A:LYS:HD2	22	3.55
(1,847)	1:36:A:HIS:HB2	1:41:A:TRP:HZ2	30	3.49
(1,888)	1:5:A:LEU:HB3	1:67:A:VAL:HA	4	3.48
(1,151)	1:20:A:THR:HA	1:56:A:GLY:HA2	8	3.46
(1,888)	1:5:A:LEU:HB3	1:67:A:VAL:HA	16	3.42
(1,888)	1:5:A:LEU:HB3	1:67:A:VAL:HA	24	3.27
(1,27)	1:43:A:GLY:HA3	1:33:A:LEU:HD12	8	3.19
(1,851)	1:13:A:TYR:HE1	1:57:A:TYR:H	25	3.17
(1,849)	1:13:A:TYR:H	1:13:A:TYR:HE1	29	3.16
(1,849)	1:13:A:TYR:H	1:13:A:TYR:HE1	2	3.1
(1,415)	1:5:A:LEU:HD12	1:7:A:VAL:HG21	28	3.1
(1,391)	1:34:A:GLU:H	1:5:A:LEU:HD22	28	3.08
(1,958)	1:61:A:SER:H	1:5:A:LEU:HD23	7	3.07
(1,849)	1:13:A:TYR:H	1:15:A:ASN:HD22	16	3.06
(1,847)	1:8:A:ARG:HD3	1:6:A:GLN:HE21	19	3.06
(1,392)	1:43:A:GLY:H	1:33:A:LEU:HD12	8	3.03
(1,849)	1:13:A:TYR:H	1:15:A:ASN:HD22	19	3.02
(1,849)	1:13:A:TYR:H	1:13:A:TYR:HE1	24	3.0
(1,305)	1:13:A:TYR:HD2	1:11:A:LYS:HE3	22	2.96
(1,847)	1:8:A:ARG:HD3	1:6:A:GLN:HE21	5	2.93
(1,851)	1:13:A:TYR:HE2	1:57:A:TYR:H	22	2.88
(1,849)	1:13:A:TYR:H	1:13:A:TYR:HE1	30	2.86
(1,883)	1:13:A:TYR:HE2	1:21:A:SER:HB3	22	2.84
(1,848)	1:16:A:ASN:HB2	1:18:A:ASP:H	26	2.84
(1,850)	1:11:A:LYS:HG3	1:13:A:TYR:HE1	1	2.83
(1,249)	1:13:A:TYR:HE2	1:21:A:SER:HA	22	2.8
(1,761)	1:15:A:ASN:H	1:22:A:LEU:H	17	2.77
(1,628)	1:38:A:ASP:H	1:36:A:HIS:HB2	15	2.76
(1,888)	1:5:A:LEU:HB3	1:67:A:VAL:HA	27	2.75
(1,848)	1:16:A:ASN:HB2	1:18:A:ASP:H	7	2.75
(1,894)	1:53:A:ASP:H	1:54:A:ARG:HD3	15	2.74
(1,849)	1:13:A:TYR:H	1:13:A:TYR:HE1	6	2.74

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,848)	1:16:A:ASN:HB2	1:18:A:ASP:H	14	2.72
(1,151)	1:20:A:THR:HA	1:56:A:GLY:HA2	25	2.72
(1,813)	1:22:A:LEU:HB2	1:13:A:TYR:H	30	2.71
(1,620)	1:21:A:SER:H	1:14:A:CYS:HB2	30	2.69
(1,418)	1:7:A:VAL:HG11	1:32:A:VAL:HG21	18	2.69
(1,848)	1:16:A:ASN:HB2	1:18:A:ASP:H	16	2.65
(1,847)	1:8:A:ARG:HD3	1:6:A:GLN:HE21	28	2.65
(1,844)	1:32:A:VAL:HG22	1:41:A:TRP:HH2	19	2.65
(1,761)	1:15:A:ASN:H	1:22:A:LEU:H	19	2.65
(1,918)	1:13:A:TYR:H	1:25:A:LYS:HB3	16	2.64
(1,427)	1:20:A:THR:HG23	1:55:A:VAL:HG21	30	2.63
(1,919)	1:23:A:ASN:HD21	1:25:A:LYS:HB3	30	2.61
(1,848)	1:16:A:ASN:HB2	1:18:A:ASP:H	2	2.6
(1,935)	1:58:A:PHE:HB3	1:40:A:ARG:HB2	1	2.59
(1,685)	1:27:A:GLY:H	1:24:A:VAL:HG22	4	2.58
(1,848)	1:16:A:ASN:HB2	1:18:A:ASP:H	5	2.56
(1,935)	1:40:A:ARG:HB3	1:38:A:ASP:HB3	2	2.54
(1,802)	1:26:A:ALA:HB2	1:12:A:ASP:HA	26	2.54
(1,793)	1:28:A:ASP:HB3	1:24:A:VAL:HG12	2	2.53
(1,234)	1:8:A:ARG:HG3	1:6:A:GLN:HE22	19	2.53
(1,192)	1:67:A:VAL:H	1:6:A:GLN:H	27	2.52
(1,935)	1:58:A:PHE:HB3	1:40:A:ARG:HB2	13	2.51
(1,935)	1:12:A:ASP:HB3	1:11:A:LYS:HG3	27	2.51
(1,901)	1:14:A:CYS:HB2	1:13:A:TYR:HD1	25	2.5
(1,847)	1:8:A:ARG:HD3	1:6:A:GLN:HE21	11	2.49
(1,395)	1:42:A:LYS:H	1:33:A:LEU:HD12	8	2.49
(1,848)	1:16:A:ASN:HB2	1:18:A:ASP:H	15	2.48
(1,849)	1:13:A:TYR:H	1:15:A:ASN:HD22	28	2.47
(1,901)	1:14:A:CYS:HB2	1:13:A:TYR:HD2	1	2.46
(1,844)	1:32:A:VAL:HG23	1:41:A:TRP:HH2	22	2.45
(1,151)	1:20:A:THR:HA	1:56:A:GLY:HA2	19	2.44
(1,57)	1:28:A:ASP:HB2	1:25:A:LYS:HG3	19	2.44
(1,813)	1:22:A:LEU:HB2	1:13:A:TYR:H	19	2.43
(1,234)	1:8:A:ARG:HG3	1:6:A:GLN:HE22	30	2.43
(1,848)	1:16:A:ASN:HB2	1:18:A:ASP:H	12	2.42
(1,151)	1:20:A:THR:HA	1:56:A:GLY:HA2	17	2.42
(1,847)	1:8:A:ARG:HD3	1:6:A:GLN:HE21	20	2.39
(1,849)	1:13:A:TYR:H	1:15:A:ASN:HD22	1	2.38
(1,390)	1:24:A:VAL:HG23	1:13:A:TYR:HB3	25	2.38
(1,926)	1:6:A:GLN:H	1:65:A:ALA:HB1	9	2.37
(1,853)	1:29:A:ILE:HG12	1:48:A:ASN:HD22	11	2.37
(1,847)	1:8:A:ARG:HD3	1:6:A:GLN:HE21	16	2.37

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,901)	1:14:A:CYS:HB2	1:13:A:TYR:HD2	24	2.35
(1,848)	1:16:A:ASN:HB2	1:18:A:ASP:H	18	2.35
(1,844)	1:32:A:VAL:HG23	1:41:A:TRP:HH2	11	2.34
(1,919)	1:23:A:ASN:HD21	1:25:A:LYS:HB3	2	2.33
(1,705)	1:35:A:GLN:HE22	1:33:A:LEU:HD22	18	2.32
(1,620)	1:21:A:SER:H	1:14:A:CYS:HB2	1	2.32
(1,935)	1:12:A:ASP:HB3	1:11:A:LYS:HG3	8	2.31
(1,935)	1:13:A:TYR:HB3	1:11:A:LYS:HG3	28	2.31
(1,545)	1:40:A:ARG:H	1:36:A:HIS:HB2	15	2.31
(1,234)	1:8:A:ARG:HG3	1:6:A:GLN:HE22	4	2.31
(1,220)	1:15:A:ASN:H	1:15:A:ASN:HD22	29	2.31
(1,281)	1:6:A:GLN:H	1:67:A:VAL:HA	24	2.27
(1,927)	1:41:A:TRP:HD1	1:40:A:ARG:HB2	8	2.24
(1,901)	1:14:A:CYS:HB2	1:13:A:TYR:HD2	22	2.24
(1,881)	1:13:A:TYR:HD2	1:21:A:SER:HB3	22	2.24
(1,918)	1:13:A:TYR:H	1:11:A:LYS:HD3	22	2.21
(1,927)	1:41:A:TRP:HD1	1:40:A:ARG:HB2	10	2.2
(1,893)	1:41:A:TRP:HZ3	1:32:A:VAL:HA	2	2.2
(1,893)	1:41:A:TRP:HZ3	1:32:A:VAL:HA	20	2.2
(1,620)	1:21:A:SER:H	1:14:A:CYS:HB2	19	2.2
(1,391)	1:34:A:GLU:H	1:5:A:LEU:HD23	29	2.2
(1,958)	1:58:A:PHE:HD1	1:33:A:LEU:HD12	12	2.19
(1,927)	1:41:A:TRP:HD1	1:40:A:ARG:HB2	22	2.19
(1,935)	1:12:A:ASP:HB3	1:11:A:LYS:HG3	25	2.17
(1,423)	1:7:A:VAL:HG21	1:32:A:VAL:HG21	18	2.17
(1,168)	1:13:A:TYR:HD2	1:22:A:LEU:H	25	2.17
(1,813)	1:22:A:LEU:HB2	1:13:A:TYR:H	17	2.16
(1,802)	1:26:A:ALA:HB3	1:12:A:ASP:HA	18	2.16
(1,225)	1:35:A:GLN:HG3	1:41:A:TRP:HD1	29	2.15
(1,935)	1:12:A:ASP:HB3	1:11:A:LYS:HG3	7	2.13
(1,918)	1:13:A:TYR:H	1:11:A:LYS:HD3	10	2.13
(1,426)	1:20:A:THR:HG23	1:55:A:VAL:HG11	30	2.13
(1,220)	1:15:A:ASN:H	1:15:A:ASN:HD22	30	2.13
(1,927)	1:41:A:TRP:HD1	1:40:A:ARG:HB2	25	2.12
(1,894)	1:53:A:ASP:H	1:54:A:ARG:HD2	8	2.12
(1,945)	1:41:A:TRP:HH2	1:5:A:LEU:HD11	23	2.11
(1,848)	1:16:A:ASN:HB2	1:18:A:ASP:H	20	2.11
(1,685)	1:27:A:GLY:H	1:24:A:VAL:HG23	13	2.11
(1,927)	1:41:A:TRP:HD1	1:40:A:ARG:HB2	7	2.1
(1,927)	1:41:A:TRP:HD1	1:40:A:ARG:HB2	23	2.1
(1,935)	1:12:A:ASP:HB3	1:11:A:LYS:HG3	6	2.09
(1,935)	1:12:A:ASP:HB3	1:11:A:LYS:HG3	17	2.09

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,850)	1:11:A:LYS:HG3	1:13:A:TYR:HE1	4	2.09
(1,793)	1:28:A:ASP:HB3	1:45:A:ILE:HG21	11	2.09
(1,360)	1:27:A:GLY:H	1:25:A:LYS:HG3	19	2.09
(1,901)	1:14:A:CYS:HB3	1:13:A:TYR:HD2	19	2.08
(1,802)	1:26:A:ALA:HB3	1:12:A:ASP:HA	9	2.06
(1,628)	1:38:A:ASP:H	1:36:A:HIS:HB2	24	2.06
(1,340)	1:63:A:GLY:H	1:7:A:VAL:HB	7	2.06
(1,919)	1:23:A:ASN:HD21	1:54:A:ARG:HB3	19	2.04
(1,893)	1:41:A:TRP:HZ3	1:61:A:SER:HB2	10	2.04
(1,364)	1:29:A:ILE:HD13	1:8:A:ARG:HG2	22	2.04
(1,220)	1:15:A:ASN:H	1:15:A:ASN:HD22	6	2.04
(1,935)	1:13:A:TYR:HB3	1:11:A:LYS:HG3	4	2.03
(1,918)	1:13:A:TYR:H	1:11:A:LYS:HD3	3	2.03
(1,916)	1:29:A:ILE:HG12	1:30:A:ILE:H	16	2.03
(1,916)	1:29:A:ILE:HG12	1:30:A:ILE:H	18	2.03
(1,878)	1:52:A:ASN:H	1:50:A:THR:HB	19	2.03
(1,916)	1:29:A:ILE:HG12	1:30:A:ILE:H	15	2.02
(1,916)	1:29:A:ILE:HG12	1:30:A:ILE:H	23	2.02
(1,894)	1:23:A:ASN:H	1:54:A:ARG:HD3	19	2.02
(1,305)	1:13:A:TYR:HD1	1:11:A:LYS:HE2	18	2.02
(1,220)	1:15:A:ASN:H	1:15:A:ASN:HD22	22	2.02
(1,916)	1:29:A:ILE:HG12	1:30:A:ILE:H	20	2.01
(1,916)	1:29:A:ILE:HG12	1:30:A:ILE:H	24	2.01
(1,916)	1:29:A:ILE:HG12	1:30:A:ILE:H	26	2.01
(1,888)	1:5:A:LEU:HB3	1:67:A:VAL:HA	5	2.01
(1,801)	1:65:A:ALA:HB2	1:5:A:LEU:HB3	28	2.01
(1,444)	1:30:A:ILE:HD12	1:32:A:VAL:HG13	18	2.01
(1,122)	1:12:A:ASP:H	1:26:A:ALA:HB3	4	2.01
(1,57)	1:28:A:ASP:HB2	1:25:A:LYS:HG2	8	2.01
(1,927)	1:41:A:TRP:HD1	1:40:A:ARG:HB2	12	2.0
(1,916)	1:29:A:ILE:HG12	1:30:A:ILE:H	13	2.0
(1,926)	1:28:A:ASP:H	1:25:A:LYS:HG2	3	1.99
(1,916)	1:29:A:ILE:HG12	1:30:A:ILE:H	11	1.99
(1,916)	1:29:A:ILE:HG12	1:30:A:ILE:H	12	1.99
(1,916)	1:29:A:ILE:HG12	1:30:A:ILE:H	27	1.99
(1,898)	1:32:A:VAL:HG23	1:60:A:SER:HA	16	1.99
(1,894)	1:23:A:ASN:H	1:54:A:ARG:HD3	3	1.99
(1,357)	1:65:A:ALA:HB1	1:5:A:LEU:HB3	3	1.99
(1,249)	1:13:A:TYR:HE1	1:21:A:SER:HA	25	1.99
(1,916)	1:29:A:ILE:HG12	1:30:A:ILE:H	14	1.98
(1,916)	1:29:A:ILE:HG12	1:30:A:ILE:H	25	1.98
(1,916)	1:29:A:ILE:HG12	1:30:A:ILE:H	29	1.98

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,878)	1:52:A:ASN:H	1:50:A:THR:HB	9	1.98
(1,844)	1:22:A:LEU:HD13	1:13:A:TYR:HD2	29	1.98
(1,926)	1:28:A:ASP:H	1:25:A:LYS:HG3	26	1.97
(1,916)	1:29:A:ILE:HG12	1:30:A:ILE:H	28	1.97
(1,881)	1:13:A:TYR:HD2	1:21:A:SER:HB2	30	1.97
(1,848)	1:16:A:ASN:HB2	1:18:A:ASP:H	21	1.97
(1,958)	1:58:A:PHE:HD1	1:33:A:LEU:HD13	26	1.96
(1,925)	1:11:A:LYS:HG2	1:13:A:TYR:H	10	1.95
(1,916)	1:29:A:ILE:HG12	1:30:A:ILE:H	1	1.95
(1,916)	1:29:A:ILE:HG12	1:30:A:ILE:H	2	1.95
(1,916)	1:29:A:ILE:HG12	1:30:A:ILE:H	21	1.95
(1,169)	1:13:A:TYR:HE2	1:22:A:LEU:H	22	1.95
(1,927)	1:41:A:TRP:HD1	1:40:A:ARG:HB2	11	1.94
(1,918)	1:13:A:TYR:H	1:11:A:LYS:HD2	12	1.94
(1,916)	1:29:A:ILE:HG12	1:30:A:ILE:H	3	1.94
(1,916)	1:29:A:ILE:HG12	1:30:A:ILE:H	7	1.94
(1,891)	1:55:A:VAL:H	1:56:A:GLY:HA2	24	1.94
(1,490)	1:41:A:TRP:HE1	1:33:A:LEU:HD21	8	1.94
(1,844)	1:22:A:LEU:HD13	1:13:A:TYR:HD2	25	1.93
(1,168)	1:13:A:TYR:HD2	1:22:A:LEU:H	30	1.93
(1,926)	1:6:A:GLN:H	1:65:A:ALA:HB1	8	1.92
(1,844)	1:32:A:VAL:HG22	1:41:A:TRP:HH2	30	1.92
(1,220)	1:15:A:ASN:H	1:15:A:ASN:HD22	16	1.92
(1,916)	1:30:A:ILE:H	1:8:A:ARG:HB2	19	1.91
(1,848)	1:16:A:ASN:HB2	1:17:A:TYR:HE1	8	1.91
(1,935)	1:13:A:TYR:HB3	1:9:A:ALA:HB3	26	1.9
(1,927)	1:41:A:TRP:HD1	1:40:A:ARG:HB3	28	1.9
(1,916)	1:10:A:THR:H	1:62:A:LEU:HG	6	1.9
(1,916)	1:29:A:ILE:HG12	1:30:A:ILE:H	9	1.9
(1,802)	1:26:A:ALA:HB2	1:12:A:ASP:HA	29	1.9
(1,396)	1:44:A:CYS:H	1:33:A:LEU:HD11	7	1.9
(1,357)	1:65:A:ALA:HB1	1:5:A:LEU:HB3	5	1.9
(1,927)	1:41:A:TRP:HD1	1:40:A:ARG:HB2	20	1.89
(1,844)	1:22:A:LEU:HD12	1:13:A:TYR:HD2	20	1.89
(1,445)	1:30:A:ILE:HD12	1:45:A:ILE:HG21	3	1.89
(1,44)	1:13:A:TYR:HB3	1:24:A:VAL:HB	25	1.89
(1,898)	1:22:A:LEU:HD12	1:44:A:CYS:HB2	24	1.88
(1,893)	1:41:A:TRP:HZ3	1:32:A:VAL:HA	12	1.88
(1,916)	1:29:A:ILE:HG12	1:30:A:ILE:H	22	1.87
(1,387)	1:63:A:GLY:H	1:7:A:VAL:HG21	7	1.87
(1,283)	1:13:A:TYR:HE2	1:56:A:GLY:HA3	22	1.87
(1,925)	1:11:A:LYS:HG2	1:13:A:TYR:H	25	1.86

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,898)	1:22:A:LEU:HD11	1:44:A:CYS:HB3	1	1.86
(1,898)	1:22:A:LEU:HD12	1:44:A:CYS:HB2	14	1.86
(1,850)	1:11:A:LYS:HG3	1:13:A:TYR:HE1	19	1.86
(1,848)	1:16:A:ASN:HB2	1:17:A:TYR:HE1	30	1.86
(1,415)	1:5:A:LEU:HD12	1:7:A:VAL:HG23	8	1.86
(1,415)	1:5:A:LEU:HD13	1:7:A:VAL:HG22	30	1.86
(1,305)	1:13:A:TYR:HD1	1:11:A:LYS:HE3	1	1.86
(1,57)	1:28:A:ASP:HB2	1:25:A:LYS:HG3	20	1.86
(1,894)	1:53:A:ASP:H	1:54:A:ARG:HD2	12	1.85
(1,802)	1:26:A:ALA:HB3	1:11:A:LYS:HA	13	1.85
(1,357)	1:65:A:ALA:HB1	1:5:A:LEU:HB3	30	1.85
(1,893)	1:41:A:TRP:HZ3	1:32:A:VAL:HA	15	1.84
(1,848)	1:16:A:ASN:HB2	1:18:A:ASP:H	1	1.84
(1,415)	1:5:A:LEU:HD12	1:7:A:VAL:HG21	5	1.84
(1,397)	1:33:A:LEU:H	1:33:A:LEU:HD11	7	1.84
(1,893)	1:41:A:TRP:HZ3	1:32:A:VAL:HA	27	1.83
(1,850)	1:11:A:LYS:HG3	1:13:A:TYR:HE1	26	1.83
(1,357)	1:65:A:ALA:HB2	1:5:A:LEU:HB3	23	1.83
(1,57)	1:28:A:ASP:HB2	1:25:A:LYS:HG2	14	1.83
(1,927)	1:41:A:TRP:HD1	1:40:A:ARG:HB3	19	1.82
(1,925)	1:11:A:LYS:HG2	1:13:A:TYR:H	16	1.82
(1,328)	1:24:A:VAL:HB	1:26:A:ALA:H	13	1.82
(1,844)	1:22:A:LEU:HD12	1:13:A:TYR:HD2	10	1.81
(1,66)	1:28:A:ASP:HB3	1:25:A:LYS:HG3	19	1.81
(1,46)	1:13:A:TYR:HB2	1:11:A:LYS:HD2	16	1.81
(1,935)	1:40:A:ARG:HB3	1:38:A:ASP:HB3	24	1.8
(1,898)	1:32:A:VAL:HG22	1:60:A:SER:HA	23	1.8
(1,948)	1:6:A:GLN:H	1:7:A:VAL:HG12	7	1.79
(1,926)	1:28:A:ASP:H	1:25:A:LYS:HG3	13	1.79
(1,916)	1:30:A:ILE:H	1:8:A:ARG:HB2	30	1.79
(1,844)	1:22:A:LEU:HD13	1:13:A:TYR:HD2	4	1.79
(1,415)	1:5:A:LEU:HD13	1:7:A:VAL:HG21	13	1.79
(1,299)	1:50:A:THR:H	1:47:A:ASP:HB2	14	1.79
(1,948)	1:6:A:GLN:H	1:7:A:VAL:HG12	8	1.78
(1,927)	1:48:A:ASN:HD22	1:25:A:LYS:HG3	9	1.78
(1,918)	1:13:A:TYR:H	1:11:A:LYS:HD3	14	1.78
(1,881)	1:13:A:TYR:HD2	1:21:A:SER:HB2	17	1.78
(1,847)	1:8:A:ARG:HD3	1:6:A:GLN:HE21	7	1.78
(1,392)	1:43:A:GLY:H	1:33:A:LEU:HD11	7	1.78
(1,948)	1:6:A:GLN:H	1:7:A:VAL:HG13	1	1.77
(1,948)	1:6:A:GLN:H	1:7:A:VAL:HG12	3	1.77
(1,926)	1:6:A:GLN:H	1:65:A:ALA:HB3	1	1.77

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,847)	1:8:A:ARG:HD3	1:6:A:GLN:HE21	9	1.77
(1,844)	1:22:A:LEU:HD13	1:13:A:TYR:HD2	1	1.77
(1,844)	1:22:A:LEU:HD12	1:13:A:TYR:HD2	16	1.77
(1,844)	1:22:A:LEU:HD13	1:13:A:TYR:HD2	18	1.77
(1,415)	1:5:A:LEU:HD13	1:7:A:VAL:HG23	3	1.77
(1,281)	1:6:A:GLN:H	1:67:A:VAL:HA	16	1.77
(1,958)	1:58:A:PHE:HD1	1:33:A:LEU:HD12	10	1.76
(1,918)	1:13:A:TYR:H	1:11:A:LYS:HD3	1	1.76
(1,898)	1:32:A:VAL:HG23	1:60:A:SER:HA	6	1.76
(1,844)	1:22:A:LEU:HD13	1:13:A:TYR:HD2	3	1.76
(1,802)	1:26:A:ALA:HB2	1:12:A:ASP:HA	1	1.76
(1,27)	1:43:A:GLY:HA3	1:33:A:LEU:HD12	27	1.76
(1,958)	1:58:A:PHE:HD1	1:33:A:LEU:HD13	15	1.75
(1,945)	1:41:A:TRP:HH2	1:5:A:LEU:HD12	16	1.75
(1,926)	1:6:A:GLN:H	1:65:A:ALA:HB3	12	1.75
(1,891)	1:55:A:VAL:H	1:56:A:GLY:HA2	16	1.75
(1,888)	1:5:A:LEU:HB3	1:67:A:VAL:HA	30	1.75
(1,660)	1:10:A:THR:H	1:11:A:LYS:HB2	10	1.75
(1,927)	1:41:A:TRP:HD1	1:40:A:ARG:HB2	1	1.74
(1,919)	1:23:A:ASN:HD21	1:54:A:ARG:HB2	1	1.74
(1,898)	1:32:A:VAL:HG23	1:60:A:SER:HA	9	1.74
(1,881)	1:13:A:TYR:HD2	1:21:A:SER:HB2	8	1.74
(1,415)	1:5:A:LEU:HD11	1:7:A:VAL:HG23	22	1.74
(1,357)	1:65:A:ALA:HB1	1:5:A:LEU:HB3	9	1.74
(1,927)	1:41:A:TRP:HD1	1:40:A:ARG:HB2	30	1.73
(1,893)	1:41:A:TRP:HZ3	1:61:A:SER:HB2	22	1.73
(1,891)	1:6:A:GLN:H	1:67:A:VAL:HA	4	1.73
(1,881)	1:13:A:TYR:HD1	1:21:A:SER:HB2	25	1.73
(1,844)	1:32:A:VAL:HG22	1:41:A:TRP:HH2	26	1.73
(1,637)	1:52:A:ASN:H	1:47:A:ASP:HB2	30	1.73
(1,545)	1:40:A:ARG:H	1:36:A:HIS:HB2	1	1.73
(1,57)	1:28:A:ASP:HB2	1:25:A:LYS:HG2	9	1.73
(1,918)	1:13:A:TYR:H	1:11:A:LYS:HD3	20	1.72
(1,893)	1:41:A:TRP:HZ3	1:32:A:VAL:HA	29	1.72
(1,445)	1:30:A:ILE:HD13	1:45:A:ILE:HG21	24	1.72
(1,415)	1:5:A:LEU:HD11	1:7:A:VAL:HG21	4	1.72
(1,415)	1:5:A:LEU:HD13	1:7:A:VAL:HG22	9	1.72
(1,945)	1:41:A:TRP:HH2	1:5:A:LEU:HD11	19	1.71
(1,926)	1:6:A:GLN:H	1:65:A:ALA:HB1	14	1.71
(1,925)	1:11:A:LYS:HG2	1:13:A:TYR:H	22	1.71
(1,850)	1:11:A:LYS:HG3	1:13:A:TYR:HE1	29	1.71
(1,844)	1:22:A:LEU:HD13	1:13:A:TYR:HD2	8	1.71

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,761)	1:15:A:ASN:H	1:22:A:LEU:H	25	1.71
(1,685)	1:27:A:GLY:H	1:24:A:VAL:HG22	29	1.71
(1,445)	1:30:A:ILE:HD13	1:45:A:ILE:HG22	22	1.71
(1,415)	1:5:A:LEU:HD11	1:7:A:VAL:HG23	1	1.71
(1,958)	1:58:A:PHE:HD1	1:33:A:LEU:HD11	19	1.7
(1,948)	1:6:A:GLN:H	1:67:A:VAL:HG11	17	1.7
(1,948)	1:6:A:GLN:H	1:7:A:VAL:HG12	23	1.7
(1,926)	1:6:A:GLN:H	1:65:A:ALA:HB3	18	1.7
(1,415)	1:5:A:LEU:HD12	1:7:A:VAL:HG23	14	1.7
(1,357)	1:65:A:ALA:HB2	1:5:A:LEU:HB3	26	1.7
(1,958)	1:58:A:PHE:HD1	1:33:A:LEU:HD13	13	1.69
(1,945)	1:41:A:TRP:HH2	1:5:A:LEU:HD12	4	1.69
(1,850)	1:11:A:LYS:HG3	1:13:A:TYR:HE1	30	1.69
(1,848)	1:16:A:ASN:HB2	1:18:A:ASP:H	11	1.69
(1,844)	1:22:A:LEU:HD12	1:13:A:TYR:HD2	5	1.69
(1,685)	1:27:A:GLY:H	1:24:A:VAL:HG21	11	1.69
(1,357)	1:65:A:ALA:HB2	1:5:A:LEU:HB3	4	1.69
(1,919)	1:23:A:ASN:HD21	1:54:A:ARG:HB2	22	1.68
(1,898)	1:32:A:VAL:HG22	1:60:A:SER:HA	5	1.68
(1,893)	1:41:A:TRP:HZ3	1:32:A:VAL:HA	26	1.68
(1,840)	1:12:A:ASP:H	1:11:A:LYS:HD2	10	1.68
(1,802)	1:26:A:ALA:HB3	1:11:A:LYS:HA	2	1.68
(1,415)	1:5:A:LEU:HD13	1:7:A:VAL:HG21	18	1.68
(1,196)	1:34:A:GLU:HB3	1:4:A:MET:H	9	1.68
(1,935)	1:40:A:ARG:HB3	1:38:A:ASP:HB3	22	1.67
(1,918)	1:13:A:TYR:H	1:11:A:LYS:HD3	29	1.67
(1,850)	1:11:A:LYS:HG3	1:13:A:TYR:HE1	17	1.67
(1,847)	1:8:A:ARG:HD3	1:6:A:GLN:HE21	22	1.67
(1,813)	1:22:A:LEU:HB2	1:13:A:TYR:H	8	1.67
(1,415)	1:5:A:LEU:HD12	1:7:A:VAL:HG22	6	1.67
(1,415)	1:5:A:LEU:HD13	1:7:A:VAL:HG21	12	1.67
(1,299)	1:50:A:THR:H	1:47:A:ASP:HB2	22	1.67
(1,724)	1:20:A:THR:H	1:14:A:CYS:HB2	1	1.66
(1,445)	1:30:A:ILE:HD12	1:45:A:ILE:HG21	1	1.66
(1,415)	1:5:A:LEU:HD11	1:7:A:VAL:HG21	10	1.66
(1,415)	1:5:A:LEU:HD13	1:7:A:VAL:HG22	19	1.66
(1,415)	1:5:A:LEU:HD13	1:7:A:VAL:HG21	23	1.66
(1,959)	1:31:A:THR:H	1:22:A:LEU:HD13	8	1.65
(1,878)	1:52:A:ASN:H	1:50:A:THR:HB	3	1.65
(1,415)	1:5:A:LEU:HD11	1:7:A:VAL:HG23	7	1.65
(1,959)	1:31:A:THR:H	1:22:A:LEU:HD13	20	1.64
(1,850)	1:11:A:LYS:HG3	1:13:A:TYR:HE1	11	1.64

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,844)	1:22:A:LEU:HD12	1:13:A:TYR:HD2	23	1.64
(1,834)	1:39:A:GLY:HA3	1:41:A:TRP:H	15	1.64
(1,445)	1:30:A:ILE:HD13	1:45:A:ILE:HG22	12	1.64
(1,430)	1:22:A:LEU:HD12	1:45:A:ILE:HG22	12	1.64
(1,357)	1:65:A:ALA:HB3	1:5:A:LEU:HB3	13	1.64
(1,958)	1:58:A:PHE:HD1	1:33:A:LEU:HD12	1	1.63
(1,958)	1:58:A:PHE:HD1	1:33:A:LEU:HD13	14	1.63
(1,948)	1:6:A:GLN:H	1:67:A:VAL:HG13	26	1.63
(1,926)	1:6:A:GLN:H	1:65:A:ALA:HB2	28	1.63
(1,916)	1:10:A:THR:H	1:11:A:LYS:HD2	10	1.63
(1,891)	1:6:A:GLN:H	1:67:A:VAL:HA	27	1.63
(1,850)	1:11:A:LYS:HG3	1:13:A:TYR:HE1	28	1.63
(1,281)	1:6:A:GLN:H	1:67:A:VAL:HA	4	1.63
(1,959)	1:31:A:THR:H	1:22:A:LEU:HD11	4	1.62
(1,916)	1:30:A:ILE:H	1:8:A:ARG:HB2	4	1.62
(1,901)	1:14:A:CYS:HB2	1:13:A:TYR:HD2	17	1.62
(1,901)	1:14:A:CYS:HB2	1:13:A:TYR:HD2	30	1.62
(1,844)	1:22:A:LEU:HD13	1:13:A:TYR:HD2	2	1.62
(1,462)	1:67:A:VAL:H	1:67:A:VAL:HG11	7	1.62
(1,963)	1:10:A:THR:H	1:30:A:ILE:HD13	30	1.61
(1,958)	1:61:A:SER:H	1:5:A:LEU:HD21	27	1.61
(1,926)	1:6:A:GLN:H	1:65:A:ALA:HB1	22	1.61
(1,848)	1:16:A:ASN:HB2	1:17:A:TYR:HE1	17	1.61
(1,844)	1:22:A:LEU:HD11	1:13:A:TYR:HD2	9	1.61
(1,844)	1:22:A:LEU:HD11	1:13:A:TYR:HD2	12	1.61
(1,761)	1:15:A:ASN:H	1:22:A:LEU:H	22	1.61
(1,462)	1:67:A:VAL:H	1:67:A:VAL:HG13	1	1.61
(1,462)	1:67:A:VAL:H	1:67:A:VAL:HG13	18	1.61
(1,462)	1:67:A:VAL:H	1:67:A:VAL:HG11	22	1.61
(1,445)	1:30:A:ILE:HD11	1:45:A:ILE:HG21	15	1.61
(1,415)	1:5:A:LEU:HD12	1:7:A:VAL:HG22	17	1.61
(1,225)	1:35:A:GLN:HG2	1:41:A:TRP:HD1	10	1.61
(1,945)	1:41:A:TRP:HH2	1:5:A:LEU:HD11	9	1.6
(1,939)	1:61:A:SER:H	1:32:A:VAL:HG11	18	1.6
(1,916)	1:22:A:LEU:H	1:54:A:ARG:HB3	8	1.6
(1,802)	1:26:A:ALA:HB1	1:11:A:LYS:HA	23	1.6
(1,462)	1:67:A:VAL:H	1:67:A:VAL:HG11	13	1.6
(1,445)	1:30:A:ILE:HD13	1:45:A:ILE:HG23	18	1.6
(1,445)	1:30:A:ILE:HD11	1:45:A:ILE:HG23	26	1.6
(1,415)	1:5:A:LEU:HD11	1:7:A:VAL:HG21	2	1.6
(1,357)	1:65:A:ALA:HB3	1:5:A:LEU:HB3	24	1.6
(1,357)	1:65:A:ALA:HB2	1:5:A:LEU:HB3	28	1.6

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,925)	1:11:A:LYS:HG2	1:13:A:TYR:H	1	1.59
(1,918)	1:13:A:TYR:H	1:11:A:LYS:HD3	27	1.59
(1,850)	1:11:A:LYS:HG3	1:13:A:TYR:HE1	2	1.59
(1,437)	1:24:A:VAL:HG21	1:30:A:ILE:HG23	30	1.59
(1,415)	1:5:A:LEU:HD11	1:7:A:VAL:HG21	26	1.59
(1,168)	1:13:A:TYR:HD2	1:22:A:LEU:H	22	1.59
(1,926)	1:28:A:ASP:H	1:25:A:LYS:HG3	6	1.58
(1,916)	1:22:A:LEU:H	1:54:A:ARG:HB2	17	1.58
(1,888)	1:5:A:LEU:HB3	1:67:A:VAL:HA	11	1.58
(1,881)	1:13:A:TYR:HD2	1:21:A:SER:HB2	19	1.58
(1,850)	1:11:A:LYS:HG3	1:13:A:TYR:HE1	5	1.58
(1,823)	1:28:A:ASP:H	1:9:A:ALA:HB3	8	1.58
(1,823)	1:28:A:ASP:H	1:9:A:ALA:HB3	25	1.58
(1,445)	1:30:A:ILE:HD13	1:45:A:ILE:HG23	17	1.58
(1,415)	1:5:A:LEU:HD11	1:7:A:VAL:HG23	15	1.58
(1,46)	1:13:A:TYR:HB2	1:11:A:LYS:HD2	3	1.58
(1,850)	1:11:A:LYS:HG3	1:13:A:TYR:HE1	27	1.57
(1,844)	1:22:A:LEU:HD11	1:13:A:TYR:HD2	13	1.57
(1,844)	1:22:A:LEU:HD12	1:13:A:TYR:HD2	21	1.57
(1,462)	1:67:A:VAL:H	1:67:A:VAL:HG12	6	1.57
(1,462)	1:67:A:VAL:H	1:67:A:VAL:HG13	11	1.57
(1,462)	1:67:A:VAL:H	1:67:A:VAL:HG11	25	1.57
(1,948)	1:6:A:GLN:H	1:7:A:VAL:HG12	27	1.56
(1,927)	1:41:A:TRP:HD1	1:40:A:ARG:HB2	2	1.56
(1,898)	1:32:A:VAL:HG21	1:60:A:SER:HA	28	1.56
(1,893)	1:41:A:TRP:HZ3	1:61:A:SER:HB2	13	1.56
(1,863)	1:9:A:ALA:H	1:29:A:ILE:HA	24	1.56
(1,847)	1:36:A:HIS:HB2	1:41:A:TRP:HZ2	14	1.56
(1,844)	1:22:A:LEU:HD12	1:13:A:TYR:HD2	28	1.56
(1,796)	1:18:A:ASP:H	1:17:A:TYR:HB2	22	1.56
(1,462)	1:67:A:VAL:H	1:67:A:VAL:HG12	8	1.56
(1,462)	1:67:A:VAL:H	1:67:A:VAL:HG11	9	1.56
(1,462)	1:67:A:VAL:H	1:67:A:VAL:HG12	10	1.56
(1,462)	1:67:A:VAL:H	1:67:A:VAL:HG13	14	1.56
(1,462)	1:67:A:VAL:H	1:67:A:VAL:HG13	20	1.56
(1,415)	1:5:A:LEU:HD11	1:7:A:VAL:HG23	16	1.56
(1,220)	1:15:A:ASN:H	1:15:A:ASN:HD22	19	1.56
(1,958)	1:61:A:SER:H	1:5:A:LEU:HD22	17	1.55
(1,918)	1:13:A:TYR:H	1:11:A:LYS:HD3	21	1.55
(1,850)	1:11:A:LYS:HG3	1:13:A:TYR:HE1	9	1.55
(1,848)	1:16:A:ASN:HB2	1:18:A:ASP:H	9	1.55
(1,579)	1:36:A:HIS:H	1:33:A:LEU:HD23	8	1.55

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,415)	1:5:A:LEU:HD11	1:7:A:VAL:HG23	25	1.55
(1,220)	1:15:A:ASN:H	1:15:A:ASN:HD22	2	1.55
(1,935)	1:40:A:ARG:HB2	1:38:A:ASP:HB3	5	1.54
(1,918)	1:13:A:TYR:H	1:11:A:LYS:HD3	17	1.54
(1,918)	1:13:A:TYR:H	1:11:A:LYS:HD3	30	1.54
(1,888)	1:5:A:LEU:HB3	1:67:A:VAL:HA	19	1.54
(1,850)	1:11:A:LYS:HG3	1:13:A:TYR:HE1	18	1.54
(1,462)	1:67:A:VAL:H	1:67:A:VAL:HG12	23	1.54
(1,462)	1:67:A:VAL:H	1:67:A:VAL:HG13	26	1.54
(1,430)	1:22:A:LEU:HD11	1:45:A:ILE:HG21	4	1.54
(1,299)	1:50:A:THR:H	1:47:A:ASP:HB2	12	1.54
(1,963)	1:10:A:THR:H	1:30:A:ILE:HD13	22	1.53
(1,959)	1:31:A:THR:H	1:22:A:LEU:HD13	25	1.53
(1,926)	1:28:A:ASP:H	1:25:A:LYS:HG3	23	1.53
(1,834)	1:39:A:GLY:HA3	1:41:A:TRP:H	9	1.53
(1,796)	1:18:A:ASP:H	1:17:A:TYR:HB2	30	1.53
(1,462)	1:67:A:VAL:H	1:67:A:VAL:HG13	28	1.53
(1,430)	1:22:A:LEU:HD11	1:45:A:ILE:HG23	18	1.53
(1,347)	1:63:A:GLY:H	1:62:A:LEU:HB3	30	1.53
(1,281)	1:6:A:GLN:H	1:67:A:VAL:HA	27	1.53
(1,66)	1:28:A:ASP:HB3	1:25:A:LYS:HG2	20	1.53
(1,966)	1:8:A:ARG:H	1:6:A:GLN:HG2	21	1.52
(1,963)	1:30:A:ILE:HD11	1:22:A:LEU:H	17	1.52
(1,898)	1:22:A:LEU:HD12	1:23:A:ASN:HB2	22	1.52
(1,853)	1:29:A:ILE:HG12	1:48:A:ASN:HD22	3	1.52
(1,844)	1:22:A:LEU:HD11	1:13:A:TYR:HD2	15	1.52
(1,795)	1:15:A:ASN:HD22	1:15:A:ASN:HB3	30	1.52
(1,519)	1:14:A:CYS:H	1:13:A:TYR:HD1	24	1.52
(1,415)	1:5:A:LEU:HD12	1:7:A:VAL:HG23	24	1.52
(1,945)	1:41:A:TRP:HH2	1:5:A:LEU:HD11	30	1.51
(1,918)	1:13:A:TYR:H	1:11:A:LYS:HD3	15	1.51
(1,878)	1:52:A:ASN:H	1:50:A:THR:HB	15	1.51
(1,850)	1:11:A:LYS:HG3	1:13:A:TYR:HE1	8	1.51
(1,847)	1:8:A:ARG:HD3	1:6:A:GLN:HE21	2	1.51
(1,828)	1:23:A:ASN:H	1:22:A:LEU:HD23	17	1.51
(1,797)	1:7:A:VAL:HB	1:64:A:GLU:HA	7	1.51
(1,462)	1:67:A:VAL:H	1:67:A:VAL:HG13	15	1.51
(1,462)	1:67:A:VAL:H	1:67:A:VAL:HG11	17	1.51
(1,462)	1:67:A:VAL:H	1:67:A:VAL:HG11	21	1.51
(1,462)	1:67:A:VAL:H	1:67:A:VAL:HG11	29	1.51
(1,151)	1:20:A:THR:HA	1:56:A:GLY:HA2	30	1.51
(1,935)	1:40:A:ARG:HB3	1:38:A:ASP:HB3	21	1.5

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,935)	1:12:A:ASP:HB3	1:11:A:LYS:HG3	23	1.5
(1,898)	1:32:A:VAL:HG21	1:60:A:SER:HA	3	1.5
(1,893)	1:41:A:TRP:HZ3	1:32:A:VAL:HA	4	1.5
(1,813)	1:22:A:LEU:HB2	1:13:A:TYR:H	29	1.5
(1,795)	1:15:A:ASN:HD22	1:15:A:ASN:HB3	25	1.5
(1,462)	1:67:A:VAL:H	1:67:A:VAL:HG13	12	1.5
(1,357)	1:65:A:ALA:HB1	1:5:A:LEU:HB3	8	1.5
(1,958)	1:58:A:PHE:HD1	1:33:A:LEU:HD12	22	1.49
(1,918)	1:13:A:TYR:H	1:11:A:LYS:HD3	23	1.49
(1,898)	1:22:A:LEU:HD13	1:23:A:ASN:HB3	8	1.49
(1,813)	1:22:A:LEU:HB2	1:13:A:TYR:H	24	1.49
(1,795)	1:15:A:ASN:HD22	1:15:A:ASN:HB3	8	1.49
(1,795)	1:15:A:ASN:HD22	1:15:A:ASN:HB3	22	1.49
(1,382)	1:65:A:ALA:H	1:5:A:LEU:HB3	3	1.49
(1,27)	1:43:A:GLY:HA3	1:33:A:LEU:HD11	25	1.49
(1,948)	1:6:A:GLN:H	1:67:A:VAL:HG11	13	1.48
(1,945)	1:41:A:TRP:HH2	1:5:A:LEU:HD12	22	1.48
(1,919)	1:23:A:ASN:HD21	1:54:A:ARG:HB2	29	1.48
(1,850)	1:11:A:LYS:HG3	1:13:A:TYR:HE1	24	1.48
(1,844)	1:32:A:VAL:HG21	1:41:A:TRP:HH2	17	1.48
(1,828)	1:23:A:ASN:H	1:22:A:LEU:HD23	15	1.48
(1,823)	1:28:A:ASP:H	1:9:A:ALA:HB3	26	1.48
(1,796)	1:18:A:ASP:H	1:17:A:TYR:HB2	25	1.48
(1,628)	1:38:A:ASP:H	1:36:A:HIS:HB2	6	1.48
(1,462)	1:67:A:VAL:H	1:67:A:VAL:HG12	19	1.48
(1,430)	1:22:A:LEU:HD11	1:45:A:ILE:HG22	6	1.48
(1,364)	1:29:A:ILE:HD13	1:8:A:ARG:HG2	19	1.48
(1,357)	1:65:A:ALA:HB3	1:5:A:LEU:HB3	16	1.48
(1,963)	1:30:A:ILE:HD11	1:22:A:LEU:H	19	1.47
(1,926)	1:6:A:GLN:H	1:65:A:ALA:HB3	24	1.47
(1,919)	1:23:A:ASN:HD21	1:25:A:LYS:HB3	4	1.47
(1,898)	1:32:A:VAL:HG23	1:60:A:SER:HA	11	1.47
(1,850)	1:11:A:LYS:HG3	1:13:A:TYR:HE1	7	1.47
(1,795)	1:15:A:ASN:HD22	1:15:A:ASN:HB3	17	1.47
(1,462)	1:67:A:VAL:H	1:67:A:VAL:HG11	27	1.47
(1,360)	1:27:A:GLY:H	1:25:A:LYS:HG3	26	1.47
(1,357)	1:65:A:ALA:HB2	1:5:A:LEU:HB3	15	1.47
(1,305)	1:13:A:TYR:HD1	1:11:A:LYS:HE3	12	1.47
(1,951)	1:8:A:ARG:H	1:7:A:VAL:HG23	12	1.46
(1,935)	1:12:A:ASP:HB3	1:11:A:LYS:HG3	11	1.46
(1,927)	1:48:A:ASN:HD22	1:25:A:LYS:HG3	6	1.46
(1,898)	1:32:A:VAL:HG22	1:60:A:SER:HA	19	1.46

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,828)	1:23:A:ASN:H	1:22:A:LEU:HD23	6	1.46
(1,828)	1:23:A:ASN:H	1:22:A:LEU:HD21	12	1.46
(1,462)	1:67:A:VAL:H	1:67:A:VAL:HG13	30	1.46
(1,415)	1:5:A:LEU:HD11	1:7:A:VAL:HG22	21	1.46
(1,225)	1:35:A:GLN:HG3	1:41:A:TRP:HD1	14	1.46
(1,963)	1:10:A:THR:H	1:30:A:ILE:HD11	8	1.45
(1,858)	1:13:A:TYR:HD2	1:21:A:SER:HA	22	1.45
(1,847)	1:8:A:ARG:HD3	1:6:A:GLN:HE21	17	1.45
(1,828)	1:23:A:ASN:H	1:22:A:LEU:HD23	19	1.45
(1,813)	1:22:A:LEU:HB2	1:13:A:TYR:H	25	1.45
(1,462)	1:67:A:VAL:H	1:67:A:VAL:HG11	4	1.45
(1,430)	1:22:A:LEU:HD13	1:45:A:ILE:HG23	16	1.45
(1,415)	1:5:A:LEU:HD12	1:7:A:VAL:HG23	20	1.45
(1,415)	1:5:A:LEU:HD12	1:7:A:VAL:HG23	27	1.45
(1,305)	1:13:A:TYR:HD1	1:11:A:LYS:HE3	25	1.45
(1,951)	1:8:A:ARG:H	1:7:A:VAL:HG22	27	1.44
(1,927)	1:48:A:ASN:HD22	1:25:A:LYS:HG3	27	1.44
(1,922)	1:23:A:ASN:H	1:22:A:LEU:HB2	11	1.44
(1,828)	1:23:A:ASN:H	1:22:A:LEU:HD21	3	1.44
(1,828)	1:23:A:ASN:H	1:22:A:LEU:HD22	7	1.44
(1,300)	1:4:A:MET:H	1:3:A:HIS:HB3	10	1.44
(1,959)	1:31:A:THR:H	1:22:A:LEU:HD12	22	1.43
(1,951)	1:8:A:ARG:H	1:7:A:VAL:HG22	1	1.43
(1,863)	1:9:A:ALA:H	1:29:A:ILE:HA	26	1.43
(1,828)	1:23:A:ASN:H	1:22:A:LEU:HD21	20	1.43
(1,802)	1:26:A:ALA:HB2	1:12:A:ASP:HA	24	1.43
(1,571)	1:66:A:ILE:H	1:65:A:ALA:HB1	22	1.43
(1,27)	1:43:A:GLY:HA3	1:33:A:LEU:HD13	4	1.43
(1,27)	1:43:A:GLY:HA3	1:33:A:LEU:HD13	26	1.43
(1,951)	1:8:A:ARG:H	1:7:A:VAL:HG23	13	1.42
(1,922)	1:23:A:ASN:H	1:22:A:LEU:HB2	9	1.42
(1,922)	1:23:A:ASN:H	1:22:A:LEU:HB2	10	1.42
(1,922)	1:23:A:ASN:H	1:22:A:LEU:HB2	19	1.42
(1,898)	1:32:A:VAL:HG22	1:60:A:SER:HA	30	1.42
(1,848)	1:16:A:ASN:HB2	1:17:A:TYR:HE1	22	1.42
(1,823)	1:28:A:ASP:H	1:9:A:ALA:HB3	20	1.42
(1,733)	1:39:A:GLY:H	1:36:A:HIS:HB3	6	1.42
(1,685)	1:27:A:GLY:H	1:24:A:VAL:HG23	19	1.42
(1,493)	1:41:A:TRP:HE1	1:40:A:ARG:H	15	1.42
(1,430)	1:22:A:LEU:HD12	1:45:A:ILE:HG21	13	1.42
(1,357)	1:65:A:ALA:HB3	1:5:A:LEU:HB3	18	1.42
(1,959)	1:31:A:THR:H	1:22:A:LEU:HD11	27	1.41

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,951)	1:8:A:ARG:H	1:7:A:VAL:HG23	23	1.41
(1,945)	1:41:A:TRP:HH2	1:5:A:LEU:HD11	13	1.41
(1,926)	1:6:A:GLN:H	1:65:A:ALA:HB1	17	1.41
(1,922)	1:23:A:ASN:H	1:22:A:LEU:HB2	14	1.41
(1,873)	1:41:A:TRP:HB3	1:34:A:GLU:HA	24	1.41
(1,844)	1:32:A:VAL:HG23	1:41:A:TRP:HH2	24	1.41
(1,844)	1:22:A:LEU:HD13	1:13:A:TYR:HD2	27	1.41
(1,828)	1:23:A:ASN:H	1:22:A:LEU:HD21	9	1.41
(1,828)	1:23:A:ASN:H	1:22:A:LEU:HD21	18	1.41
(1,493)	1:41:A:TRP:HE1	1:40:A:ARG:H	14	1.41
(1,445)	1:30:A:ILE:HD13	1:45:A:ILE:HG23	16	1.41
(1,430)	1:22:A:LEU:HD13	1:45:A:ILE:HG23	21	1.41
(1,963)	1:30:A:ILE:HD13	1:22:A:LEU:H	13	1.4
(1,963)	1:30:A:ILE:HD11	1:22:A:LEU:H	16	1.4
(1,958)	1:58:A:PHE:HD1	1:33:A:LEU:HD13	2	1.4
(1,958)	1:61:A:SER:H	1:5:A:LEU:HD23	20	1.4
(1,951)	1:8:A:ARG:H	1:7:A:VAL:HG22	14	1.4
(1,922)	1:66:A:ILE:H	1:6:A:GLN:HB3	26	1.4
(1,898)	1:32:A:VAL:HG23	1:60:A:SER:HA	4	1.4
(1,893)	1:41:A:TRP:HZ3	1:32:A:VAL:HA	17	1.4
(1,893)	1:41:A:TRP:HZ3	1:61:A:SER:HB2	19	1.4
(1,888)	1:5:A:LEU:HB3	1:67:A:VAL:HA	20	1.4
(1,887)	1:29:A:ILE:HG22	1:31:A:THR:HB	19	1.4
(1,828)	1:23:A:ASN:H	1:22:A:LEU:HD22	14	1.4
(1,828)	1:23:A:ASN:H	1:22:A:LEU:HD21	21	1.4
(1,813)	1:22:A:LEU:HB2	1:13:A:TYR:H	1	1.4
(1,445)	1:30:A:ILE:HD12	1:45:A:ILE:HG22	29	1.4
(1,357)	1:65:A:ALA:HB1	1:5:A:LEU:HB3	6	1.4
(1,357)	1:65:A:ALA:HB1	1:5:A:LEU:HB3	22	1.4
(1,951)	1:13:A:TYR:H	1:24:A:VAL:HG21	4	1.39
(1,951)	1:13:A:TYR:H	1:24:A:VAL:HG11	6	1.39
(1,951)	1:8:A:ARG:H	1:7:A:VAL:HG22	7	1.39
(1,926)	1:6:A:GLN:H	1:65:A:ALA:HB1	29	1.39
(1,898)	1:32:A:VAL:HG22	1:60:A:SER:HA	26	1.39
(1,853)	1:29:A:ILE:HG12	1:48:A:ASN:HD22	4	1.39
(1,844)	1:22:A:LEU:HD13	1:13:A:TYR:HD2	7	1.39
(1,844)	1:22:A:LEU:HD13	1:13:A:TYR:HD2	14	1.39
(1,828)	1:23:A:ASN:H	1:22:A:LEU:HD22	28	1.39
(1,802)	1:26:A:ALA:HB2	1:12:A:ASP:HA	3	1.39
(1,801)	1:65:A:ALA:HB1	1:5:A:LEU:HB3	29	1.39
(1,685)	1:27:A:GLY:H	1:24:A:VAL:HG22	2	1.39
(1,571)	1:66:A:ILE:H	1:65:A:ALA:HB1	10	1.39

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,445)	1:30:A:ILE:HD13	1:45:A:ILE:HG23	14	1.39
(1,445)	1:30:A:ILE:HD12	1:45:A:ILE:HG23	25	1.39
(1,963)	1:10:A:THR:H	1:30:A:ILE:HD12	4	1.38
(1,963)	1:30:A:ILE:HD13	1:22:A:LEU:H	5	1.38
(1,963)	1:10:A:THR:H	1:30:A:ILE:HD13	11	1.38
(1,958)	1:61:A:SER:H	1:5:A:LEU:HD21	16	1.38
(1,926)	1:28:A:ASP:H	1:25:A:LYS:HG2	7	1.38
(1,813)	1:22:A:LEU:HB2	1:13:A:TYR:H	11	1.38
(1,673)	1:34:A:GLU:H	1:33:A:LEU:HB2	8	1.38
(1,415)	1:5:A:LEU:HD11	1:7:A:VAL:HG22	11	1.38
(1,357)	1:65:A:ALA:HB3	1:5:A:LEU:HB3	12	1.38
(1,357)	1:65:A:ALA:HB1	1:5:A:LEU:HB3	14	1.38
(1,357)	1:65:A:ALA:HB2	1:5:A:LEU:HB3	19	1.38
(1,283)	1:13:A:TYR:HE2	1:56:A:GLY:HA3	25	1.38
(1,212)	1:52:A:ASN:HD21	1:52:A:ASN:H	14	1.38
(1,169)	1:13:A:TYR:HE2	1:22:A:LEU:H	25	1.38
(1,46)	1:13:A:TYR:HB2	1:11:A:LYS:HD2	20	1.38
(1,966)	1:8:A:ARG:H	1:6:A:GLN:HG2	20	1.37
(1,963)	1:30:A:ILE:HD12	1:22:A:LEU:H	26	1.37
(1,951)	1:8:A:ARG:H	1:7:A:VAL:HG21	9	1.37
(1,951)	1:8:A:ARG:H	1:7:A:VAL:HG21	25	1.37
(1,951)	1:13:A:TYR:H	1:24:A:VAL:HG11	29	1.37
(1,948)	1:6:A:GLN:H	1:67:A:VAL:HG13	15	1.37
(1,919)	1:23:A:ASN:HD21	1:25:A:LYS:HB3	24	1.37
(1,898)	1:32:A:VAL:HG22	1:60:A:SER:HA	2	1.37
(1,892)	1:13:A:TYR:HD2	1:56:A:GLY:HA3	26	1.37
(1,828)	1:23:A:ASN:H	1:22:A:LEU:HD22	1	1.37
(1,828)	1:23:A:ASN:H	1:22:A:LEU:HD23	13	1.37
(1,828)	1:23:A:ASN:H	1:22:A:LEU:HD23	23	1.37
(1,828)	1:23:A:ASN:H	1:22:A:LEU:HD21	27	1.37
(1,823)	1:28:A:ASP:H	1:9:A:ALA:HB3	24	1.37
(1,802)	1:26:A:ALA:HB1	1:12:A:ASP:HA	19	1.37
(1,761)	1:15:A:ASN:H	1:22:A:LEU:H	8	1.37
(1,724)	1:20:A:THR:H	1:14:A:CYS:HB2	30	1.37
(1,416)	1:5:A:LEU:HD23	1:7:A:VAL:HG12	9	1.37
(1,963)	1:30:A:ILE:HD11	1:22:A:LEU:H	23	1.36
(1,948)	1:6:A:GLN:H	1:67:A:VAL:HG13	12	1.36
(1,849)	1:13:A:TYR:H	1:13:A:TYR:HE1	22	1.36
(1,847)	1:36:A:HIS:HB2	1:41:A:TRP:HZ2	29	1.36
(1,834)	1:39:A:GLY:HA3	1:41:A:TRP:H	8	1.36
(1,828)	1:23:A:ASN:H	1:22:A:LEU:HD23	29	1.36
(1,571)	1:66:A:ILE:H	1:65:A:ALA:HB1	14	1.36

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,490)	1:41:A:TRP:HE1	1:33:A:LEU:HD22	7	1.36
(1,445)	1:30:A:ILE:HD11	1:45:A:ILE:HG23	21	1.36
(1,431)	1:22:A:LEU:HD11	1:24:A:VAL:HG12	30	1.36
(1,324)	1:4:A:MET:H	1:4:A:MET:HG3	10	1.36
(1,963)	1:10:A:THR:H	1:30:A:ILE:HD12	20	1.35
(1,959)	1:31:A:THR:H	1:22:A:LEU:HD13	10	1.35
(1,951)	1:13:A:TYR:H	1:24:A:VAL:HG11	19	1.35
(1,947)	1:7:A:VAL:HG22	1:9:A:ALA:H	1	1.35
(1,945)	1:41:A:TRP:HH2	1:5:A:LEU:HD11	3	1.35
(1,908)	1:52:A:ASN:H	1:49:A:ARG:HB3	20	1.35
(1,834)	1:39:A:GLY:HA3	1:41:A:TRP:H	3	1.35
(1,828)	1:23:A:ASN:H	1:22:A:LEU:HD21	26	1.35
(1,430)	1:22:A:LEU:HD13	1:45:A:ILE:HG23	10	1.35
(1,430)	1:22:A:LEU:HD11	1:45:A:ILE:HG23	26	1.35
(1,950)	1:9:A:ALA:H	1:24:A:VAL:HG11	1	1.34
(1,945)	1:41:A:TRP:HH2	1:5:A:LEU:HD13	5	1.34
(1,892)	1:13:A:TYR:HD2	1:24:A:VAL:HA	19	1.34
(1,720)	1:51:A:GLY:H	1:50:A:THR:H	6	1.34
(1,468)	1:30:A:ILE:H	1:30:A:ILE:HG12	2	1.34
(1,430)	1:22:A:LEU:HD13	1:45:A:ILE:HG22	28	1.34
(1,357)	1:65:A:ALA:HB2	1:5:A:LEU:HB3	7	1.34
(1,227)	1:29:A:ILE:H	1:48:A:ASN:HD22	11	1.34
(1,212)	1:52:A:ASN:HD21	1:52:A:ASN:H	17	1.34
(1,950)	1:9:A:ALA:H	1:24:A:VAL:HG12	11	1.33
(1,935)	1:40:A:ARG:HB3	1:38:A:ASP:HB3	20	1.33
(1,918)	1:13:A:TYR:H	1:11:A:LYS:HD2	18	1.33
(1,918)	1:13:A:TYR:H	1:25:A:LYS:HB3	25	1.33
(1,823)	1:28:A:ASP:H	1:25:A:LYS:HG3	23	1.33
(1,571)	1:66:A:ILE:H	1:65:A:ALA:HB1	25	1.33
(1,430)	1:22:A:LEU:HD11	1:45:A:ILE:HG21	2	1.33
(1,430)	1:22:A:LEU:HD11	1:45:A:ILE:HG22	7	1.33
(1,234)	1:8:A:ARG:HG3	1:6:A:GLN:HE22	5	1.33
(1,27)	1:43:A:GLY:HA3	1:33:A:LEU:HD13	18	1.33
(1,951)	1:8:A:ARG:H	1:7:A:VAL:HG22	22	1.32
(1,942)	1:32:A:VAL:HB	1:7:A:VAL:HG23	7	1.32
(1,942)	1:32:A:VAL:HB	1:7:A:VAL:HG22	30	1.32
(1,828)	1:23:A:ASN:H	1:22:A:LEU:HD21	16	1.32
(1,828)	1:23:A:ASN:H	1:22:A:LEU:HD23	24	1.32
(1,812)	1:30:A:ILE:HG21	1:33:A:LEU:H	10	1.32
(1,812)	1:30:A:ILE:HG23	1:33:A:LEU:H	27	1.32
(1,802)	1:26:A:ALA:HB2	1:12:A:ASP:HA	28	1.32
(1,560)	1:64:A:GLU:H	1:64:A:GLU:HG2	29	1.32

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,445)	1:30:A:ILE:HD12	1:45:A:ILE:HG22	28	1.32
(1,430)	1:22:A:LEU:HD12	1:45:A:ILE:HG22	22	1.32
(1,430)	1:22:A:LEU:HD13	1:45:A:ILE:HG22	23	1.32
(1,225)	1:35:A:GLN:HG2	1:41:A:TRP:HD1	15	1.32
(1,927)	1:41:A:TRP:HD1	1:40:A:ARG:HB2	14	1.31
(1,927)	1:48:A:ASN:HD22	1:25:A:LYS:HG3	21	1.31
(1,922)	1:23:A:ASN:H	1:22:A:LEU:HB2	29	1.31
(1,918)	1:13:A:TYR:H	1:11:A:LYS:HD3	19	1.31
(1,812)	1:30:A:ILE:HG22	1:33:A:LEU:H	11	1.31
(1,720)	1:51:A:GLY:H	1:50:A:THR:H	2	1.31
(1,430)	1:22:A:LEU:HD13	1:45:A:ILE:HG23	25	1.31
(1,382)	1:65:A:ALA:H	1:5:A:LEU:HB3	26	1.31
(1,357)	1:65:A:ALA:HB3	1:5:A:LEU:HB3	1	1.31
(1,300)	1:4:A:MET:H	1:3:A:HIS:HB3	26	1.31
(1,27)	1:43:A:GLY:HA3	1:33:A:LEU:HD12	22	1.31
(1,951)	1:13:A:TYR:H	1:24:A:VAL:HG12	15	1.3
(1,950)	1:7:A:VAL:HG22	1:9:A:ALA:H	3	1.3
(1,950)	1:45:A:ILE:HD12	1:45:A:ILE:H	29	1.3
(1,947)	1:45:A:ILE:HD11	1:45:A:ILE:H	11	1.3
(1,922)	1:66:A:ILE:H	1:6:A:GLN:HB3	13	1.3
(1,898)	1:32:A:VAL:HG21	1:60:A:SER:HA	10	1.3
(1,848)	1:16:A:ASN:HB2	1:18:A:ASP:H	13	1.3
(1,828)	1:23:A:ASN:H	1:22:A:LEU:HD21	4	1.3
(1,430)	1:22:A:LEU:HD13	1:45:A:ILE:HG22	20	1.3
(1,382)	1:65:A:ALA:H	1:5:A:LEU:HB3	9	1.3
(1,382)	1:65:A:ALA:H	1:5:A:LEU:HB3	23	1.3
(1,299)	1:50:A:THR:H	1:47:A:ASP:HB2	8	1.3
(1,958)	1:61:A:SER:H	1:5:A:LEU:HD23	9	1.29
(1,951)	1:8:A:ARG:H	1:7:A:VAL:HG23	11	1.29
(1,942)	1:32:A:VAL:HB	1:7:A:VAL:HG21	4	1.29
(1,942)	1:32:A:VAL:HB	1:7:A:VAL:HG22	25	1.29
(1,935)	1:12:A:ASP:HB3	1:11:A:LYS:HG3	12	1.29
(1,934)	1:26:A:ALA:H	1:25:A:LYS:HG3	4	1.29
(1,918)	1:13:A:TYR:H	1:11:A:LYS:HD3	4	1.29
(1,898)	1:22:A:LEU:HD13	1:23:A:ASN:HB2	25	1.29
(1,707)	1:35:A:GLN:HE22	1:35:A:GLN:HB3	18	1.29
(1,493)	1:41:A:TRP:HE1	1:40:A:ARG:H	3	1.29
(1,445)	1:30:A:ILE:HD13	1:45:A:ILE:HG22	7	1.29
(1,430)	1:22:A:LEU:HD11	1:45:A:ILE:HG21	1	1.29
(1,427)	1:20:A:THR:HG22	1:55:A:VAL:HG22	5	1.29
(1,347)	1:63:A:GLY:H	1:62:A:LEU:HB3	7	1.29
(1,963)	1:10:A:THR:H	1:30:A:ILE:HD13	24	1.28

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,956)	1:24:A:VAL:HG21	1:13:A:TYR:HD2	14	1.28
(1,951)	1:13:A:TYR:H	1:24:A:VAL:HG13	3	1.28
(1,951)	1:8:A:ARG:H	1:7:A:VAL:HG23	10	1.28
(1,951)	1:13:A:TYR:H	1:24:A:VAL:HG13	18	1.28
(1,945)	1:41:A:TRP:HH2	1:5:A:LEU:HD12	15	1.28
(1,942)	1:32:A:VAL:HB	1:7:A:VAL:HG22	9	1.28
(1,942)	1:32:A:VAL:HB	1:7:A:VAL:HG21	10	1.28
(1,942)	1:32:A:VAL:HB	1:7:A:VAL:HG21	13	1.28
(1,942)	1:32:A:VAL:HB	1:7:A:VAL:HG23	20	1.28
(1,935)	1:40:A:ARG:HB3	1:38:A:ASP:HB3	18	1.28
(1,909)	1:41:A:TRP:HD1	1:40:A:ARG:HG2	4	1.28
(1,853)	1:29:A:ILE:HG12	1:48:A:ASN:HD22	9	1.28
(1,852)	1:13:A:TYR:HE2	1:24:A:VAL:H	14	1.28
(1,823)	1:28:A:ASP:H	1:25:A:LYS:HG2	2	1.28
(1,812)	1:30:A:ILE:HG22	1:33:A:LEU:H	26	1.28
(1,797)	1:7:A:VAL:HB	1:63:A:GLY:HA2	19	1.28
(1,571)	1:66:A:ILE:H	1:65:A:ALA:HB3	11	1.28
(1,445)	1:30:A:ILE:HD12	1:45:A:ILE:HG22	6	1.28
(1,430)	1:22:A:LEU:HD11	1:45:A:ILE:HG22	29	1.28
(1,416)	1:5:A:LEU:HD23	1:7:A:VAL:HG13	10	1.28
(1,300)	1:4:A:MET:H	1:3:A:HIS:HB3	1	1.28
(1,299)	1:50:A:THR:H	1:47:A:ASP:HB2	7	1.28
(1,46)	1:13:A:TYR:HB2	1:11:A:LYS:HD2	30	1.28
(1,27)	1:43:A:GLY:HA3	1:33:A:LEU:HD12	28	1.28
(1,963)	1:10:A:THR:H	1:30:A:ILE:HD11	10	1.27
(1,951)	1:13:A:TYR:H	1:24:A:VAL:HG13	28	1.27
(1,934)	1:26:A:ALA:H	1:25:A:LYS:HG3	26	1.27
(1,926)	1:6:A:GLN:H	1:65:A:ALA:HB1	25	1.27
(1,898)	1:32:A:VAL:HG21	1:60:A:SER:HA	7	1.27
(1,850)	1:11:A:LYS:HG3	1:13:A:TYR:HE1	15	1.27
(1,834)	1:39:A:GLY:HA3	1:41:A:TRP:H	11	1.27
(1,610)	1:58:A:PHE:H	1:57:A:TYR:HB2	22	1.27
(1,579)	1:36:A:HIS:H	1:33:A:LEU:HD21	7	1.27
(1,468)	1:30:A:ILE:H	1:30:A:ILE:HG12	10	1.27
(1,430)	1:22:A:LEU:HD11	1:45:A:ILE:HG23	27	1.27
(1,234)	1:8:A:ARG:HG3	1:6:A:GLN:HE22	22	1.27
(1,168)	1:13:A:TYR:HD2	1:22:A:LEU:H	19	1.27
(1,950)	1:45:A:ILE:HD13	1:45:A:ILE:H	19	1.26
(1,942)	1:32:A:VAL:HB	1:7:A:VAL:HG23	1	1.26
(1,942)	1:32:A:VAL:HB	1:7:A:VAL:HG21	11	1.26
(1,942)	1:32:A:VAL:HB	1:7:A:VAL:HG23	27	1.26
(1,887)	1:29:A:ILE:HG22	1:31:A:THR:HB	14	1.26

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,823)	1:28:A:ASP:H	1:9:A:ALA:HB1	13	1.26
(1,813)	1:22:A:LEU:HB2	1:13:A:TYR:H	16	1.26
(1,812)	1:30:A:ILE:HG23	1:33:A:LEU:H	17	1.26
(1,812)	1:30:A:ILE:HG23	1:33:A:LEU:H	20	1.26
(1,796)	1:18:A:ASP:H	1:17:A:TYR:HB2	8	1.26
(1,610)	1:58:A:PHE:H	1:57:A:TYR:HB2	1	1.26
(1,203)	1:52:A:ASN:HD22	1:52:A:ASN:H	14	1.26
(1,190)	1:34:A:GLU:HG2	1:3:A:HIS:H	15	1.26
(1,27)	1:43:A:GLY:HA3	1:33:A:LEU:HD12	17	1.26
(1,950)	1:45:A:ILE:HD13	1:45:A:ILE:H	9	1.25
(1,950)	1:7:A:VAL:HG21	1:9:A:ALA:H	17	1.25
(1,948)	1:6:A:GLN:H	1:67:A:VAL:HG23	11	1.25
(1,900)	1:23:A:ASN:HB3	1:24:A:VAL:H	24	1.25
(1,828)	1:23:A:ASN:H	1:22:A:LEU:HD21	5	1.25
(1,828)	1:23:A:ASN:H	1:22:A:LEU:HD22	11	1.25
(1,823)	1:28:A:ASP:H	1:9:A:ALA:HB2	14	1.25
(1,795)	1:15:A:ASN:HD22	1:15:A:ASN:HB3	27	1.25
(1,743)	1:63:A:GLY:H	1:62:A:LEU:HG	14	1.25
(1,468)	1:30:A:ILE:H	1:30:A:ILE:HG12	4	1.25
(1,382)	1:65:A:ALA:H	1:5:A:LEU:HB3	13	1.25
(1,357)	1:65:A:ALA:HB1	1:5:A:LEU:HB3	17	1.25
(1,66)	1:28:A:ASP:HB3	1:25:A:LYS:HG2	14	1.25
(1,27)	1:43:A:GLY:HA3	1:33:A:LEU:HD11	6	1.25
(1,966)	1:8:A:ARG:H	1:6:A:GLN:HG2	25	1.24
(1,963)	1:10:A:THR:H	1:30:A:ILE:HD12	27	1.24
(1,951)	1:8:A:ARG:H	1:7:A:VAL:HG22	20	1.24
(1,951)	1:8:A:ARG:H	1:7:A:VAL:HG22	24	1.24
(1,950)	1:45:A:ILE:HD12	1:45:A:ILE:H	13	1.24
(1,927)	1:48:A:ASN:HD22	1:25:A:LYS:HG3	26	1.24
(1,898)	1:32:A:VAL:HG21	1:60:A:SER:HA	12	1.24
(1,898)	1:32:A:VAL:HG22	1:60:A:SER:HA	21	1.24
(1,834)	1:39:A:GLY:HA3	1:41:A:TRP:H	7	1.24
(1,828)	1:23:A:ASN:H	1:22:A:LEU:HD21	10	1.24
(1,828)	1:23:A:ASN:H	1:22:A:LEU:HD21	22	1.24
(1,828)	1:23:A:ASN:H	1:22:A:LEU:HD23	30	1.24
(1,707)	1:35:A:GLN:HE22	1:35:A:GLN:HB3	21	1.24
(1,571)	1:66:A:ILE:H	1:65:A:ALA:HB1	29	1.24
(1,445)	1:30:A:ILE:HD12	1:45:A:ILE:HG22	5	1.24
(1,364)	1:29:A:ILE:HD13	1:8:A:ARG:HG2	4	1.24
(1,360)	1:27:A:GLY:H	1:25:A:LYS:HG2	18	1.24
(1,66)	1:28:A:ASP:HB3	1:25:A:LYS:HG2	8	1.24
(1,17)	1:26:A:ALA:HA	1:24:A:VAL:HG13	4	1.24

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,950)	1:45:A:ILE:HD12	1:45:A:ILE:H	20	1.23
(1,942)	1:32:A:VAL:HB	1:7:A:VAL:HG22	17	1.23
(1,942)	1:32:A:VAL:HB	1:7:A:VAL:HG22	29	1.23
(1,922)	1:66:A:ILE:H	1:6:A:GLN:HB3	20	1.23
(1,918)	1:13:A:TYR:H	1:11:A:LYS:HD3	11	1.23
(1,823)	1:28:A:ASP:H	1:9:A:ALA:HB3	16	1.23
(1,812)	1:30:A:ILE:HG21	1:33:A:LEU:H	1	1.23
(1,541)	1:40:A:ARG:H	1:40:A:ARG:HD2	5	1.23
(1,416)	1:5:A:LEU:HD21	1:7:A:VAL:HG12	8	1.23
(1,346)	1:55:A:VAL:H	1:54:A:ARG:HB3	4	1.23
(1,299)	1:50:A:THR:H	1:47:A:ASP:HB2	11	1.23
(1,168)	1:13:A:TYR:HD2	1:22:A:LEU:H	8	1.23
(1,959)	1:31:A:THR:H	1:22:A:LEU:HD12	13	1.22
(1,950)	1:7:A:VAL:HG22	1:9:A:ALA:H	15	1.22
(1,942)	1:32:A:VAL:HB	1:7:A:VAL:HG23	15	1.22
(1,942)	1:32:A:VAL:HB	1:7:A:VAL:HG22	19	1.22
(1,942)	1:32:A:VAL:HB	1:7:A:VAL:HG22	21	1.22
(1,828)	1:23:A:ASN:H	1:22:A:LEU:HD23	2	1.22
(1,802)	1:26:A:ALA:HB2	1:12:A:ASP:HA	14	1.22
(1,796)	1:18:A:ASP:H	1:17:A:TYR:HB2	17	1.22
(1,571)	1:66:A:ILE:H	1:65:A:ALA:HB1	9	1.22
(1,468)	1:30:A:ILE:H	1:30:A:ILE:HG12	20	1.22
(1,445)	1:30:A:ILE:HD11	1:45:A:ILE:HG22	9	1.22
(1,430)	1:22:A:LEU:HD12	1:45:A:ILE:HG21	15	1.22
(1,430)	1:22:A:LEU:HD13	1:45:A:ILE:HG23	17	1.22
(1,416)	1:5:A:LEU:HD23	1:7:A:VAL:HG12	13	1.22
(1,228)	1:51:A:GLY:H	1:52:A:ASN:HD22	14	1.22
(1,958)	1:61:A:SER:H	1:5:A:LEU:HD22	18	1.21
(1,951)	1:13:A:TYR:H	1:24:A:VAL:HG13	17	1.21
(1,927)	1:48:A:ASN:HD22	1:25:A:LYS:HG3	13	1.21
(1,898)	1:32:A:VAL:HG23	1:60:A:SER:HA	13	1.21
(1,887)	1:29:A:ILE:HG21	1:31:A:THR:HB	30	1.21
(1,878)	1:52:A:ASN:H	1:50:A:THR:HB	10	1.21
(1,813)	1:22:A:LEU:HB2	1:13:A:TYR:H	26	1.21
(1,430)	1:22:A:LEU:HD12	1:45:A:ILE:HG22	9	1.21
(1,416)	1:5:A:LEU:HD21	1:7:A:VAL:HG13	1	1.21
(1,416)	1:5:A:LEU:HD21	1:7:A:VAL:HG12	27	1.21
(1,360)	1:27:A:GLY:H	1:25:A:LYS:HG3	24	1.21
(1,346)	1:55:A:VAL:H	1:54:A:ARG:HB3	3	1.21
(1,58)	1:28:A:ASP:H	1:28:A:ASP:HB2	2	1.21
(1,947)	1:7:A:VAL:HG22	1:9:A:ALA:H	3	1.2
(1,947)	1:45:A:ILE:HD12	1:45:A:ILE:H	24	1.2

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,947)	1:45:A:ILE:HD12	1:45:A:ILE:H	29	1.2
(1,892)	1:41:A:TRP:HH2	1:61:A:SER:HA	10	1.2
(1,850)	1:11:A:LYS:HG3	1:13:A:TYR:HE1	13	1.2
(1,733)	1:39:A:GLY:H	1:36:A:HIS:HB3	15	1.2
(1,547)	1:6:A:GLN:H	1:5:A:LEU:HB3	1	1.2
(1,468)	1:30:A:ILE:H	1:30:A:ILE:HG12	13	1.2
(1,445)	1:30:A:ILE:HD13	1:45:A:ILE:HG21	11	1.2
(1,437)	1:24:A:VAL:HG22	1:30:A:ILE:HG21	14	1.2
(1,430)	1:22:A:LEU:HD13	1:45:A:ILE:HG23	8	1.2
(1,430)	1:22:A:LEU:HD11	1:45:A:ILE:HG21	24	1.2
(1,416)	1:5:A:LEU:HD23	1:7:A:VAL:HG12	7	1.2
(1,364)	1:29:A:ILE:HD12	1:8:A:ARG:HG2	30	1.2
(1,959)	1:31:A:THR:H	1:22:A:LEU:HD11	6	1.19
(1,959)	1:31:A:THR:H	1:22:A:LEU:HD11	19	1.19
(1,959)	1:31:A:THR:H	1:22:A:LEU:HD11	30	1.19
(1,947)	1:45:A:ILE:HD11	1:45:A:ILE:H	5	1.19
(1,947)	1:45:A:ILE:HD13	1:45:A:ILE:H	26	1.19
(1,945)	1:41:A:TRP:HH2	1:5:A:LEU:HD12	25	1.19
(1,942)	1:32:A:VAL:HB	1:7:A:VAL:HG23	8	1.19
(1,942)	1:32:A:VAL:HB	1:7:A:VAL:HG21	12	1.19
(1,935)	1:40:A:ARG:HB3	1:38:A:ASP:HB3	15	1.19
(1,935)	1:40:A:ARG:HB3	1:38:A:ASP:HB3	30	1.19
(1,931)	1:6:A:GLN:HE21	1:8:A:ARG:HG2	2	1.19
(1,909)	1:41:A:TRP:HD1	1:40:A:ARG:HG2	25	1.19
(1,894)	1:53:A:ASP:H	1:54:A:ARG:HD2	27	1.19
(1,844)	1:22:A:LEU:HD13	1:13:A:TYR:HD2	6	1.19
(1,828)	1:23:A:ASN:H	1:22:A:LEU:HD21	8	1.19
(1,823)	1:28:A:ASP:H	1:9:A:ALA:HB1	22	1.19
(1,685)	1:27:A:GLY:H	1:24:A:VAL:HG21	26	1.19
(1,571)	1:66:A:ILE:H	1:65:A:ALA:HB2	19	1.19
(1,571)	1:66:A:ILE:H	1:65:A:ALA:HB3	20	1.19
(1,468)	1:30:A:ILE:H	1:30:A:ILE:HG12	5	1.19
(1,437)	1:24:A:VAL:HG22	1:30:A:ILE:HG23	9	1.19
(1,430)	1:22:A:LEU:HD11	1:45:A:ILE:HG21	3	1.19
(1,427)	1:20:A:THR:HG23	1:55:A:VAL:HG22	28	1.19
(1,124)	1:26:A:ALA:HB2	1:11:A:LYS:H	4	1.19
(1,963)	1:30:A:ILE:HD13	1:22:A:LEU:H	2	1.18
(1,963)	1:10:A:THR:H	1:30:A:ILE:HD12	25	1.18
(1,951)	1:8:A:ARG:H	1:7:A:VAL:HG21	21	1.18
(1,950)	1:45:A:ILE:HD12	1:45:A:ILE:H	16	1.18
(1,950)	1:7:A:VAL:HG23	1:9:A:ALA:H	23	1.18
(1,950)	1:45:A:ILE:HD12	1:45:A:ILE:H	28	1.18

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,898)	1:32:A:VAL:HG23	1:60:A:SER:HA	20	1.18
(1,887)	1:29:A:ILE:HG22	1:31:A:THR:HB	16	1.18
(1,812)	1:30:A:ILE:HG22	1:33:A:LEU:H	12	1.18
(1,812)	1:30:A:ILE:HG23	1:33:A:LEU:H	24	1.18
(1,812)	1:30:A:ILE:HG21	1:33:A:LEU:H	28	1.18
(1,610)	1:58:A:PHE:H	1:57:A:TYR:HB2	21	1.18
(1,571)	1:66:A:ILE:H	1:65:A:ALA:HB1	3	1.18
(1,547)	1:6:A:GLN:H	1:5:A:LEU:HB3	11	1.18
(1,445)	1:30:A:ILE:HD12	1:45:A:ILE:HG21	13	1.18
(1,434)	1:24:A:VAL:HG12	1:30:A:ILE:HD13	30	1.18
(1,427)	1:20:A:THR:HG21	1:55:A:VAL:HG21	24	1.18
(1,416)	1:5:A:LEU:HD23	1:7:A:VAL:HG13	21	1.18
(1,168)	1:13:A:TYR:HD2	1:22:A:LEU:H	17	1.18
(1,951)	1:8:A:ARG:H	1:7:A:VAL:HG22	16	1.17
(1,950)	1:45:A:ILE:HD11	1:45:A:ILE:H	8	1.17
(1,945)	1:41:A:TRP:HH2	1:5:A:LEU:HD13	14	1.17
(1,935)	1:12:A:ASP:HB3	1:11:A:LYS:HG3	3	1.17
(1,925)	1:11:A:LYS:HG2	1:13:A:TYR:H	13	1.17
(1,918)	1:13:A:TYR:H	1:11:A:LYS:HD3	2	1.17
(1,892)	1:41:A:TRP:HH2	1:61:A:SER:HA	4	1.17
(1,870)	1:29:A:ILE:HG21	1:28:A:ASP:HA	19	1.17
(1,848)	1:16:A:ASN:HB2	1:17:A:TYR:HE1	19	1.17
(1,802)	1:26:A:ALA:HB1	1:12:A:ASP:HA	25	1.17
(1,793)	1:28:A:ASP:HB3	1:45:A:ILE:HG23	10	1.17
(1,720)	1:51:A:GLY:H	1:50:A:THR:H	17	1.17
(1,587)	1:53:A:ASP:H	1:53:A:ASP:HB2	13	1.17
(1,547)	1:6:A:GLN:H	1:5:A:LEU:HB3	18	1.17
(1,468)	1:30:A:ILE:H	1:30:A:ILE:HG12	14	1.17
(1,445)	1:30:A:ILE:HD13	1:45:A:ILE:HG22	23	1.17
(1,430)	1:22:A:LEU:HD13	1:45:A:ILE:HG22	5	1.17
(1,430)	1:22:A:LEU:HD11	1:45:A:ILE:HG22	30	1.17
(1,389)	1:11:A:LYS:HB3	1:24:A:VAL:HG12	22	1.17
(1,300)	1:4:A:MET:H	1:3:A:HIS:HB3	12	1.17
(1,62)	1:18:A:ASP:HB2	1:18:A:ASP:H	23	1.17
(1,951)	1:13:A:TYR:H	1:24:A:VAL:HG13	26	1.16
(1,948)	1:6:A:GLN:H	1:67:A:VAL:HG11	4	1.16
(1,947)	1:45:A:ILE:HD13	1:45:A:ILE:H	19	1.16
(1,942)	1:32:A:VAL:HB	1:7:A:VAL:HG23	16	1.16
(1,942)	1:32:A:VAL:HB	1:7:A:VAL:HG21	23	1.16
(1,927)	1:48:A:ASN:HD22	1:25:A:LYS:HG2	16	1.16
(1,927)	1:48:A:ASN:HD22	1:25:A:LYS:HG2	24	1.16
(1,926)	1:28:A:ASP:H	1:25:A:LYS:HG3	15	1.16

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,926)	1:6:A:GLN:H	1:65:A:ALA:HB2	19	1.16
(1,850)	1:11:A:LYS:HG3	1:13:A:TYR:HE1	6	1.16
(1,847)	1:8:A:ARG:HD3	1:6:A:GLN:HE21	27	1.16
(1,813)	1:22:A:LEU:HB2	1:13:A:TYR:H	22	1.16
(1,812)	1:30:A:ILE:HG21	1:33:A:LEU:H	21	1.16
(1,795)	1:17:A:TYR:HB3	1:18:A:ASP:H	6	1.16
(1,685)	1:27:A:GLY:H	1:24:A:VAL:HG22	1	1.16
(1,571)	1:66:A:ILE:H	1:65:A:ALA:HB2	23	1.16
(1,547)	1:6:A:GLN:H	1:5:A:LEU:HB3	2	1.16
(1,468)	1:30:A:ILE:H	1:30:A:ILE:HG12	19	1.16
(1,468)	1:30:A:ILE:H	1:30:A:ILE:HG12	21	1.16
(1,416)	1:5:A:LEU:HD22	1:7:A:VAL:HG13	15	1.16
(1,414)	1:5:A:LEU:HD11	1:32:A:VAL:HG13	29	1.16
(1,382)	1:65:A:ALA:H	1:5:A:LEU:HB3	28	1.16
(1,359)	1:12:A:ASP:H	1:11:A:LYS:HG2	10	1.16
(1,357)	1:65:A:ALA:HB1	1:5:A:LEU:HB3	10	1.16
(1,228)	1:51:A:GLY:H	1:52:A:ASN:HD22	8	1.16
(1,959)	1:31:A:THR:H	1:22:A:LEU:HD12	11	1.15
(1,950)	1:9:A:ALA:H	1:24:A:VAL:HG12	25	1.15
(1,950)	1:45:A:ILE:HD13	1:45:A:ILE:H	27	1.15
(1,948)	1:6:A:GLN:H	1:67:A:VAL:HG21	18	1.15
(1,948)	1:6:A:GLN:H	1:67:A:VAL:HG21	21	1.15
(1,947)	1:45:A:ILE:HD13	1:45:A:ILE:H	9	1.15
(1,947)	1:7:A:VAL:HG21	1:9:A:ALA:H	17	1.15
(1,942)	1:32:A:VAL:HB	1:5:A:LEU:HD12	28	1.15
(1,926)	1:6:A:GLN:H	1:65:A:ALA:HB2	4	1.15
(1,926)	1:28:A:ASP:H	1:25:A:LYS:HG2	30	1.15
(1,922)	1:66:A:ILE:H	1:6:A:GLN:HB3	12	1.15
(1,912)	1:34:A:GLU:HB3	1:35:A:GLN:H	28	1.15
(1,900)	1:24:A:VAL:H	1:23:A:ASN:HB2	25	1.15
(1,892)	1:13:A:TYR:HD2	1:56:A:GLY:HA3	18	1.15
(1,892)	1:13:A:TYR:HD2	1:24:A:VAL:HA	29	1.15
(1,812)	1:30:A:ILE:HG23	1:33:A:LEU:H	22	1.15
(1,685)	1:27:A:GLY:H	1:24:A:VAL:HG23	18	1.15
(1,610)	1:58:A:PHE:H	1:57:A:TYR:HB2	17	1.15
(1,493)	1:41:A:TRP:HE1	1:40:A:ARG:H	7	1.15
(1,485)	1:46:A:HIS:H	1:45:A:ILE:HD13	30	1.15
(1,444)	1:30:A:ILE:HD11	1:32:A:VAL:HG11	27	1.15
(1,430)	1:22:A:LEU:HD12	1:45:A:ILE:HG21	11	1.15
(1,430)	1:22:A:LEU:HD11	1:45:A:ILE:HG22	19	1.15
(1,203)	1:52:A:ASN:HD22	1:52:A:ASN:H	17	1.15
(1,62)	1:18:A:ASP:HB2	1:18:A:ASP:H	27	1.15

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,27)	1:43:A:GLY:HA3	1:33:A:LEU:HD12	29	1.15
(1,959)	1:31:A:THR:H	1:22:A:LEU:HD13	23	1.14
(1,947)	1:45:A:ILE:HD12	1:45:A:ILE:H	13	1.14
(1,942)	1:32:A:VAL:HB	1:7:A:VAL:HG21	2	1.14
(1,935)	1:40:A:ARG:HB3	1:38:A:ASP:HB3	16	1.14
(1,919)	1:23:A:ASN:HD21	1:25:A:LYS:HB3	14	1.14
(1,898)	1:32:A:VAL:HG22	1:60:A:SER:HA	29	1.14
(1,850)	1:11:A:LYS:HG3	1:13:A:TYR:HE1	23	1.14
(1,834)	1:39:A:GLY:HA3	1:41:A:TRP:H	2	1.14
(1,797)	1:7:A:VAL:HB	1:63:A:GLY:HA2	12	1.14
(1,441)	1:30:A:ILE:HG21	1:45:A:ILE:HG23	18	1.14
(1,403)	1:58:A:PHE:H	1:22:A:LEU:HD12	24	1.14
(1,382)	1:65:A:ALA:H	1:5:A:LEU:HB3	8	1.14
(1,360)	1:27:A:GLY:H	1:25:A:LYS:HG3	25	1.14
(1,66)	1:28:A:ASP:HB3	1:25:A:LYS:HG2	9	1.14
(1,27)	1:43:A:GLY:HA3	1:33:A:LEU:HD13	14	1.14
(1,963)	1:30:A:ILE:HD12	1:22:A:LEU:H	21	1.13
(1,959)	1:31:A:THR:H	1:22:A:LEU:HD13	21	1.13
(1,951)	1:8:A:ARG:H	1:7:A:VAL:HG23	5	1.13
(1,950)	1:45:A:ILE:HD11	1:45:A:ILE:H	2	1.13
(1,950)	1:9:A:ALA:H	1:24:A:VAL:HG11	5	1.13
(1,947)	1:45:A:ILE:HD12	1:45:A:ILE:H	20	1.13
(1,945)	1:41:A:TRP:HH2	1:5:A:LEU:HD12	21	1.13
(1,942)	1:32:A:VAL:HB	1:7:A:VAL:HG23	22	1.13
(1,922)	1:66:A:ILE:H	1:6:A:GLN:HB3	7	1.13
(1,918)	1:13:A:TYR:H	1:11:A:LYS:HD2	13	1.13
(1,834)	1:39:A:GLY:HA3	1:41:A:TRP:H	17	1.13
(1,823)	1:28:A:ASP:H	1:9:A:ALA:HB1	21	1.13
(1,714)	1:15:A:ASN:HD22	1:15:A:ASN:HA	2	1.13
(1,685)	1:27:A:GLY:H	1:24:A:VAL:HG22	23	1.13
(1,628)	1:38:A:ASP:H	1:36:A:HIS:HB2	1	1.13
(1,610)	1:58:A:PHE:H	1:57:A:TYR:HB2	9	1.13
(1,610)	1:58:A:PHE:H	1:57:A:TYR:HB2	25	1.13
(1,571)	1:66:A:ILE:H	1:65:A:ALA:HB3	13	1.13
(1,547)	1:6:A:GLN:H	1:5:A:LEU:HB3	8	1.13
(1,547)	1:6:A:GLN:H	1:5:A:LEU:HB3	17	1.13
(1,493)	1:41:A:TRP:HE1	1:40:A:ARG:H	28	1.13
(1,468)	1:30:A:ILE:H	1:30:A:ILE:HG12	7	1.13
(1,468)	1:30:A:ILE:H	1:30:A:ILE:HG12	22	1.13
(1,468)	1:30:A:ILE:H	1:30:A:ILE:HG12	23	1.13
(1,403)	1:58:A:PHE:H	1:22:A:LEU:HD12	3	1.13
(1,27)	1:43:A:GLY:HA3	1:33:A:LEU:HD12	11	1.13

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,965)	1:26:A:ALA:H	1:12:A:ASP:HA	13	1.12
(1,963)	1:30:A:ILE:HD11	1:22:A:LEU:H	18	1.12
(1,959)	1:31:A:THR:H	1:22:A:LEU:HD11	2	1.12
(1,951)	1:13:A:TYR:H	1:24:A:VAL:HG11	8	1.12
(1,951)	1:13:A:TYR:H	1:24:A:VAL:HG13	30	1.12
(1,947)	1:7:A:VAL:HG22	1:9:A:ALA:H	15	1.12
(1,942)	1:32:A:VAL:HB	1:7:A:VAL:HG21	26	1.12
(1,935)	1:40:A:ARG:HB3	1:38:A:ASP:HB3	9	1.12
(1,925)	1:11:A:LYS:HG2	1:13:A:TYR:H	3	1.12
(1,848)	1:16:A:ASN:HB2	1:18:A:ASP:H	29	1.12
(1,812)	1:30:A:ILE:HG22	1:33:A:LEU:H	25	1.12
(1,685)	1:27:A:GLY:H	1:24:A:VAL:HG22	10	1.12
(1,610)	1:58:A:PHE:H	1:57:A:TYR:HB2	12	1.12
(1,571)	1:66:A:ILE:H	1:65:A:ALA:HB1	30	1.12
(1,468)	1:30:A:ILE:H	1:30:A:ILE:HG12	1	1.12
(1,468)	1:30:A:ILE:H	1:30:A:ILE:HG12	28	1.12
(1,439)	1:30:A:ILE:HG22	1:32:A:VAL:HG13	18	1.12
(1,416)	1:5:A:LEU:HD21	1:7:A:VAL:HG12	22	1.12
(1,382)	1:65:A:ALA:H	1:5:A:LEU:HB3	5	1.12
(1,346)	1:55:A:VAL:H	1:54:A:ARG:HB3	12	1.12
(1,346)	1:55:A:VAL:H	1:54:A:ARG:HB3	19	1.12
(1,341)	1:29:A:ILE:H	1:29:A:ILE:HG12	5	1.12
(1,966)	1:8:A:ARG:H	1:6:A:GLN:HG2	10	1.11
(1,958)	1:61:A:SER:H	1:5:A:LEU:HD23	24	1.11
(1,956)	1:24:A:VAL:HG23	1:13:A:TYR:HD2	16	1.11
(1,813)	1:22:A:LEU:HB2	1:13:A:TYR:H	10	1.11
(1,812)	1:30:A:ILE:HG22	1:33:A:LEU:H	4	1.11
(1,812)	1:30:A:ILE:HG22	1:33:A:LEU:H	13	1.11
(1,812)	1:30:A:ILE:HG22	1:33:A:LEU:H	14	1.11
(1,812)	1:30:A:ILE:HG22	1:33:A:LEU:H	15	1.11
(1,812)	1:30:A:ILE:HG22	1:33:A:LEU:H	18	1.11
(1,543)	1:40:A:ARG:H	1:40:A:ARG:HG2	5	1.11
(1,468)	1:30:A:ILE:H	1:30:A:ILE:HG12	8	1.11
(1,444)	1:30:A:ILE:HD13	1:32:A:VAL:HG12	10	1.11
(1,416)	1:5:A:LEU:HD23	1:7:A:VAL:HG11	14	1.11
(1,300)	1:4:A:MET:H	1:3:A:HIS:HB2	28	1.11
(1,46)	1:13:A:TYR:HB2	1:11:A:LYS:HD2	14	1.11
(1,950)	1:7:A:VAL:HG22	1:9:A:ALA:H	14	1.1
(1,942)	1:32:A:VAL:HB	1:7:A:VAL:HG23	14	1.1
(1,928)	1:11:A:LYS:H	1:11:A:LYS:HG2	24	1.1
(1,928)	1:11:A:LYS:H	1:11:A:LYS:HG2	27	1.1
(1,928)	1:9:A:ALA:HB1	1:11:A:LYS:H	28	1.1

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,928)	1:9:A:ALA:HB3	1:11:A:LYS:H	30	1.1
(1,926)	1:6:A:GLN:H	1:65:A:ALA:HB2	2	1.1
(1,812)	1:30:A:ILE:HG22	1:33:A:LEU:H	2	1.1
(1,812)	1:30:A:ILE:HG21	1:33:A:LEU:H	29	1.1
(1,802)	1:26:A:ALA:HB3	1:12:A:ASP:HA	12	1.1
(1,797)	1:7:A:VAL:HB	1:63:A:GLY:HA2	10	1.1
(1,714)	1:15:A:ASN:HD22	1:15:A:ASN:HA	5	1.1
(1,714)	1:15:A:ASN:HD22	1:15:A:ASN:HA	15	1.1
(1,468)	1:30:A:ILE:H	1:30:A:ILE:HG12	9	1.1
(1,468)	1:30:A:ILE:H	1:30:A:ILE:HG12	11	1.1
(1,468)	1:30:A:ILE:H	1:30:A:ILE:HG12	27	1.1
(1,468)	1:30:A:ILE:H	1:30:A:ILE:HG12	29	1.1
(1,434)	1:24:A:VAL:HG12	1:30:A:ILE:HD11	9	1.1
(1,382)	1:65:A:ALA:H	1:5:A:LEU:HB3	15	1.1
(1,324)	1:4:A:MET:H	1:4:A:MET:HG3	29	1.1
(1,966)	1:8:A:ARG:H	1:64:A:GLU:HG3	16	1.09
(1,950)	1:45:A:ILE:HD13	1:45:A:ILE:H	22	1.09
(1,928)	1:9:A:ALA:HB2	1:11:A:LYS:H	26	1.09
(1,922)	1:66:A:ILE:H	1:6:A:GLN:HB3	23	1.09
(1,912)	1:34:A:GLU:HB3	1:35:A:GLN:H	29	1.09
(1,887)	1:29:A:ILE:HG22	1:31:A:THR:HB	6	1.09
(1,863)	1:25:A:LYS:H	1:12:A:ASP:HA	25	1.09
(1,850)	1:11:A:LYS:HG3	1:13:A:TYR:HE1	10	1.09
(1,848)	1:16:A:ASN:HB2	1:18:A:ASP:H	6	1.09
(1,822)	1:54:A:ARG:HD3	1:55:A:VAL:H	6	1.09
(1,813)	1:22:A:LEU:HB2	1:13:A:TYR:H	4	1.09
(1,812)	1:30:A:ILE:HG21	1:33:A:LEU:H	3	1.09
(1,775)	1:38:A:ASP:H	1:40:A:ARG:H	27	1.09
(1,714)	1:15:A:ASN:HD22	1:15:A:ASN:HA	3	1.09
(1,571)	1:66:A:ILE:H	1:65:A:ALA:HB3	12	1.09
(1,554)	1:54:A:ARG:H	1:54:A:ARG:HB2	21	1.09
(1,541)	1:40:A:ARG:H	1:40:A:ARG:HD2	28	1.09
(1,519)	1:14:A:CYS:H	1:13:A:TYR:HD1	22	1.09
(1,468)	1:30:A:ILE:H	1:30:A:ILE:HG12	6	1.09
(1,430)	1:22:A:LEU:HD11	1:45:A:ILE:HG23	14	1.09
(1,416)	1:5:A:LEU:HD23	1:7:A:VAL:HG12	12	1.09
(1,416)	1:5:A:LEU:HD22	1:7:A:VAL:HG13	17	1.09
(1,346)	1:55:A:VAL:H	1:54:A:ARG:HB3	9	1.09
(1,228)	1:51:A:GLY:H	1:52:A:ASN:HD22	30	1.09
(1,196)	1:34:A:GLU:HB3	1:4:A:MET:H	14	1.09
(1,62)	1:18:A:ASP:HB2	1:18:A:ASP:H	13	1.09
(1,947)	1:45:A:ILE:HD12	1:45:A:ILE:H	16	1.08

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,947)	1:7:A:VAL:HG23	1:9:A:ALA:H	23	1.08
(1,947)	1:45:A:ILE:HD12	1:45:A:ILE:H	28	1.08
(1,887)	1:29:A:ILE:HG22	1:31:A:THR:HB	2	1.08
(1,887)	1:29:A:ILE:HG22	1:31:A:THR:HB	4	1.08
(1,851)	1:13:A:TYR:HE2	1:57:A:TYR:H	24	1.08
(1,847)	1:8:A:ARG:HD3	1:6:A:GLN:HE21	13	1.08
(1,714)	1:15:A:ASN:HD22	1:15:A:ASN:HA	9	1.08
(1,714)	1:15:A:ASN:HD22	1:15:A:ASN:HA	28	1.08
(1,547)	1:6:A:GLN:H	1:5:A:LEU:HB3	6	1.08
(1,547)	1:6:A:GLN:H	1:5:A:LEU:HB3	10	1.08
(1,493)	1:41:A:TRP:HE1	1:40:A:ARG:H	26	1.08
(1,468)	1:30:A:ILE:H	1:30:A:ILE:HG12	24	1.08
(1,440)	1:30:A:ILE:HG22	1:32:A:VAL:HG22	18	1.08
(1,300)	1:4:A:MET:H	1:3:A:HIS:HB3	27	1.08
(1,966)	1:8:A:ARG:H	1:64:A:GLU:HG3	4	1.07
(1,950)	1:45:A:ILE:HD13	1:45:A:ILE:H	30	1.07
(1,947)	1:45:A:ILE:HD11	1:45:A:ILE:H	8	1.07
(1,934)	1:26:A:ALA:H	1:25:A:LYS:HG3	28	1.07
(1,922)	1:66:A:ILE:H	1:6:A:GLN:HB3	4	1.07
(1,916)	1:29:A:ILE:HG12	1:30:A:ILE:H	5	1.07
(1,898)	1:32:A:VAL:HG22	1:60:A:SER:HA	27	1.07
(1,892)	1:13:A:TYR:HD2	1:56:A:GLY:HA3	20	1.07
(1,850)	1:11:A:LYS:HG3	1:13:A:TYR:HE1	14	1.07
(1,828)	1:23:A:ASN:H	1:22:A:LEU:HD21	25	1.07
(1,813)	1:22:A:LEU:HB2	1:13:A:TYR:H	12	1.07
(1,812)	1:30:A:ILE:HG23	1:33:A:LEU:H	6	1.07
(1,610)	1:58:A:PHE:H	1:57:A:TYR:HB2	20	1.07
(1,560)	1:64:A:GLU:H	1:64:A:GLU:HG2	17	1.07
(1,547)	1:6:A:GLN:H	1:5:A:LEU:HB3	15	1.07
(1,547)	1:6:A:GLN:H	1:5:A:LEU:HB3	20	1.07
(1,468)	1:30:A:ILE:H	1:30:A:ILE:HG12	30	1.07
(1,437)	1:24:A:VAL:HG22	1:30:A:ILE:HG21	18	1.07
(1,437)	1:24:A:VAL:HG23	1:30:A:ILE:HG23	19	1.07
(1,389)	1:11:A:LYS:HB3	1:24:A:VAL:HG12	10	1.07
(1,299)	1:50:A:THR:H	1:47:A:ASP:HB2	5	1.07
(1,152)	1:20:A:THR:HA	1:56:A:GLY:HA3	22	1.07
(1,947)	1:45:A:ILE:HD12	1:45:A:ILE:H	25	1.06
(1,942)	1:32:A:VAL:HB	1:7:A:VAL:HG21	24	1.06
(1,939)	1:58:A:PHE:HD1	1:32:A:VAL:HG13	20	1.06
(1,934)	1:26:A:ALA:H	1:25:A:LYS:HG3	21	1.06
(1,919)	1:23:A:ASN:HD21	1:54:A:ARG:HB3	26	1.06
(1,900)	1:23:A:ASN:HB3	1:24:A:VAL:H	1	1.06

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,898)	1:32:A:VAL:HG21	1:60:A:SER:HA	17	1.06
(1,863)	1:9:A:ALA:H	1:29:A:ILE:HA	16	1.06
(1,850)	1:11:A:LYS:HG3	1:13:A:TYR:HE1	21	1.06
(1,849)	1:13:A:TYR:H	1:15:A:ASN:HD22	5	1.06
(1,823)	1:28:A:ASP:H	1:9:A:ALA:HB1	27	1.06
(1,813)	1:22:A:LEU:HB2	1:13:A:TYR:H	3	1.06
(1,810)	1:23:A:ASN:H	1:45:A:ILE:H	19	1.06
(1,620)	1:21:A:SER:H	1:14:A:CYS:HB2	11	1.06
(1,610)	1:58:A:PHE:H	1:57:A:TYR:HB2	13	1.06
(1,547)	1:6:A:GLN:H	1:5:A:LEU:HB3	14	1.06
(1,547)	1:6:A:GLN:H	1:5:A:LEU:HB3	25	1.06
(1,547)	1:6:A:GLN:H	1:5:A:LEU:HB3	26	1.06
(1,485)	1:46:A:HIS:H	1:45:A:ILE:HD11	4	1.06
(1,485)	1:46:A:HIS:H	1:45:A:ILE:HD11	10	1.06
(1,468)	1:30:A:ILE:H	1:30:A:ILE:HG12	16	1.06
(1,416)	1:5:A:LEU:HD23	1:7:A:VAL:HG13	20	1.06
(1,360)	1:27:A:GLY:H	1:25:A:LYS:HG3	29	1.06
(1,330)	1:63:A:GLY:H	1:62:A:LEU:HB2	30	1.06
(1,324)	1:4:A:MET:H	1:4:A:MET:HG3	7	1.06
(1,14)	1:27:A:GLY:H	1:26:A:ALA:HA	4	1.06
(1,968)	1:10:A:THR:H	1:63:A:GLY:HA3	22	1.05
(1,959)	1:31:A:THR:H	1:22:A:LEU:HD11	29	1.05
(1,950)	1:9:A:ALA:H	1:24:A:VAL:HG12	18	1.05
(1,947)	1:45:A:ILE:HD13	1:45:A:ILE:H	27	1.05
(1,942)	1:32:A:VAL:HB	1:7:A:VAL:HG23	3	1.05
(1,942)	1:32:A:VAL:HB	1:7:A:VAL:HG22	6	1.05
(1,928)	1:60:A:SER:HB3	1:41:A:TRP:HH2	4	1.05
(1,926)	1:28:A:ASP:H	1:25:A:LYS:HG3	5	1.05
(1,922)	1:66:A:ILE:H	1:6:A:GLN:HB3	24	1.05
(1,912)	1:34:A:GLU:HB3	1:35:A:GLN:H	2	1.05
(1,899)	1:27:A:GLY:H	1:28:A:ASP:HB2	29	1.05
(1,894)	1:23:A:ASN:H	1:54:A:ARG:HD3	28	1.05
(1,892)	1:13:A:TYR:HD2	1:56:A:GLY:HA3	2	1.05
(1,892)	1:13:A:TYR:HD2	1:56:A:GLY:HA3	5	1.05
(1,892)	1:41:A:TRP:HH2	1:61:A:SER:HA	17	1.05
(1,888)	1:5:A:LEU:HB3	1:67:A:VAL:HA	25	1.05
(1,848)	1:16:A:ASN:HB2	1:18:A:ASP:H	4	1.05
(1,813)	1:22:A:LEU:HB2	1:13:A:TYR:H	15	1.05
(1,810)	1:23:A:ASN:H	1:45:A:ILE:H	30	1.05
(1,798)	1:24:A:VAL:HB	1:12:A:ASP:HA	19	1.05
(1,797)	1:7:A:VAL:HB	1:63:A:GLY:HA2	6	1.05
(1,737)	1:50:A:THR:H	1:50:A:THR:HG23	2	1.05

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,610)	1:58:A:PHE:H	1:57:A:TYR:HB2	4	1.05
(1,610)	1:58:A:PHE:H	1:57:A:TYR:HB2	6	1.05
(1,547)	1:6:A:GLN:H	1:5:A:LEU:HB3	21	1.05
(1,468)	1:30:A:ILE:H	1:30:A:ILE:HG12	18	1.05
(1,416)	1:5:A:LEU:HD23	1:7:A:VAL:HG12	3	1.05
(1,382)	1:65:A:ALA:H	1:5:A:LEU:HB3	30	1.05
(1,358)	1:41:A:TRP:H	1:40:A:ARG:HB3	5	1.05
(1,353)	1:22:A:LEU:HD11	1:22:A:LEU:HB2	2	1.05
(1,353)	1:22:A:LEU:HD12	1:22:A:LEU:HB2	5	1.05
(1,353)	1:22:A:LEU:HD12	1:22:A:LEU:HB2	8	1.05
(1,353)	1:22:A:LEU:HD12	1:22:A:LEU:HB2	10	1.05
(1,353)	1:22:A:LEU:HD11	1:22:A:LEU:HB2	22	1.05
(1,353)	1:22:A:LEU:HD13	1:22:A:LEU:HB2	30	1.05
(1,324)	1:4:A:MET:H	1:4:A:MET:HG3	4	1.05
(1,220)	1:15:A:ASN:H	1:15:A:ASN:HD22	18	1.05
(1,193)	1:20:A:THR:H	1:20:A:THR:HG22	10	1.05
(1,959)	1:31:A:THR:H	1:22:A:LEU:HD13	16	1.04
(1,951)	1:13:A:TYR:H	1:24:A:VAL:HG21	2	1.04
(1,945)	1:41:A:TRP:HH2	1:5:A:LEU:HD13	8	1.04
(1,939)	1:58:A:PHE:HD1	1:32:A:VAL:HG13	17	1.04
(1,934)	1:26:A:ALA:H	1:25:A:LYS:HG3	25	1.04
(1,928)	1:9:A:ALA:HB2	1:11:A:LYS:H	12	1.04
(1,928)	1:9:A:ALA:HB3	1:11:A:LYS:H	22	1.04
(1,918)	1:13:A:TYR:H	1:11:A:LYS:HD3	6	1.04
(1,918)	1:13:A:TYR:H	1:11:A:LYS:HD3	7	1.04
(1,888)	1:5:A:LEU:HB3	1:67:A:VAL:HA	14	1.04
(1,812)	1:30:A:ILE:HG22	1:33:A:LEU:H	23	1.04
(1,796)	1:18:A:ASP:H	1:17:A:TYR:HB2	19	1.04
(1,685)	1:27:A:GLY:H	1:24:A:VAL:HG22	17	1.04
(1,571)	1:66:A:ILE:H	1:65:A:ALA:HB2	2	1.04
(1,560)	1:64:A:GLU:H	1:64:A:GLU:HG2	7	1.04
(1,547)	1:6:A:GLN:H	1:5:A:LEU:HB3	27	1.04
(1,468)	1:30:A:ILE:H	1:30:A:ILE:HG12	17	1.04
(1,468)	1:30:A:ILE:H	1:30:A:ILE:HG12	25	1.04
(1,444)	1:30:A:ILE:HD13	1:32:A:VAL:HG13	20	1.04
(1,440)	1:30:A:ILE:HG21	1:32:A:VAL:HG21	21	1.04
(1,382)	1:65:A:ALA:H	1:5:A:LEU:HB3	18	1.04
(1,353)	1:22:A:LEU:HD13	1:22:A:LEU:HB2	1	1.04
(1,353)	1:22:A:LEU:HD11	1:22:A:LEU:HB2	11	1.04
(1,353)	1:22:A:LEU:HD11	1:22:A:LEU:HB2	13	1.04
(1,353)	1:22:A:LEU:HD13	1:22:A:LEU:HB2	24	1.04
(1,353)	1:22:A:LEU:HD13	1:22:A:LEU:HB2	25	1.04

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,193)	1:20:A:THR:H	1:20:A:THR:HG23	6	1.04
(1,193)	1:20:A:THR:H	1:20:A:THR:HG22	13	1.04
(1,959)	1:31:A:THR:H	1:22:A:LEU:HD11	18	1.03
(1,956)	1:24:A:VAL:HG21	1:13:A:TYR:HD2	9	1.03
(1,956)	1:24:A:VAL:HG21	1:13:A:TYR:HD2	18	1.03
(1,950)	1:45:A:ILE:HD11	1:45:A:ILE:H	6	1.03
(1,950)	1:45:A:ILE:HD11	1:45:A:ILE:H	7	1.03
(1,947)	1:45:A:ILE:HD11	1:45:A:ILE:H	2	1.03
(1,928)	1:9:A:ALA:HB1	1:11:A:LYS:H	17	1.03
(1,909)	1:41:A:TRP:HD1	1:40:A:ARG:HG2	8	1.03
(1,894)	1:53:A:ASP:H	1:54:A:ARG:HD2	22	1.03
(1,812)	1:30:A:ILE:HG21	1:33:A:LEU:H	7	1.03
(1,812)	1:30:A:ILE:HG21	1:33:A:LEU:H	9	1.03
(1,802)	1:26:A:ALA:HB1	1:12:A:ASP:HA	20	1.03
(1,734)	1:39:A:GLY:H	1:40:A:ARG:HB2	17	1.03
(1,617)	1:21:A:SER:H	1:21:A:SER:HB3	8	1.03
(1,610)	1:58:A:PHE:H	1:57:A:TYR:HB2	29	1.03
(1,468)	1:30:A:ILE:H	1:30:A:ILE:HG12	12	1.03
(1,440)	1:30:A:ILE:HG23	1:32:A:VAL:HG21	22	1.03
(1,440)	1:30:A:ILE:HG21	1:32:A:VAL:HG23	28	1.03
(1,416)	1:5:A:LEU:HD22	1:7:A:VAL:HG13	2	1.03
(1,382)	1:65:A:ALA:H	1:5:A:LEU:HB3	12	1.03
(1,382)	1:65:A:ALA:H	1:5:A:LEU:HB3	22	1.03
(1,353)	1:22:A:LEU:HD13	1:22:A:LEU:HB2	3	1.03
(1,353)	1:22:A:LEU:HD12	1:22:A:LEU:HB2	16	1.03
(1,353)	1:22:A:LEU:HD13	1:22:A:LEU:HB2	18	1.03
(1,353)	1:22:A:LEU:HD12	1:22:A:LEU:HB2	23	1.03
(1,353)	1:22:A:LEU:HD13	1:22:A:LEU:HB2	26	1.03
(1,353)	1:22:A:LEU:HD13	1:22:A:LEU:HB2	29	1.03
(1,193)	1:20:A:THR:H	1:20:A:THR:HG22	26	1.03
(1,190)	1:34:A:GLU:HG2	1:3:A:HIS:H	1	1.03
(1,45)	1:47:A:ASP:HB3	1:52:A:ASN:HA	15	1.03
(1,963)	1:30:A:ILE:HD12	1:22:A:LEU:H	15	1.02
(1,963)	1:30:A:ILE:HD13	1:22:A:LEU:H	29	1.02
(1,958)	1:61:A:SER:H	1:5:A:LEU:HD21	29	1.02
(1,950)	1:45:A:ILE:HD11	1:45:A:ILE:H	4	1.02
(1,945)	1:41:A:TRP:HH2	1:5:A:LEU:HD13	6	1.02
(1,925)	1:11:A:LYS:HG2	1:13:A:TYR:H	12	1.02
(1,922)	1:66:A:ILE:H	1:6:A:GLN:HB3	3	1.02
(1,918)	1:13:A:TYR:H	1:11:A:LYS:HD3	5	1.02
(1,906)	1:6:A:GLN:H	1:6:A:GLN:HG2	1	1.02
(1,906)	1:6:A:GLN:H	1:6:A:GLN:HG2	8	1.02

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,887)	1:29:A:ILE:HG21	1:31:A:THR:HB	5	1.02
(1,863)	1:9:A:ALA:H	1:29:A:ILE:HA	23	1.02
(1,850)	1:11:A:LYS:HG3	1:13:A:TYR:HE1	20	1.02
(1,797)	1:7:A:VAL:HB	1:63:A:GLY:HA2	13	1.02
(1,617)	1:21:A:SER:H	1:21:A:SER:HB3	10	1.02
(1,617)	1:21:A:SER:H	1:21:A:SER:HB3	20	1.02
(1,617)	1:21:A:SER:H	1:21:A:SER:HB3	26	1.02
(1,610)	1:58:A:PHE:H	1:57:A:TYR:HB2	15	1.02
(1,610)	1:58:A:PHE:H	1:57:A:TYR:HB2	24	1.02
(1,444)	1:30:A:ILE:HD12	1:32:A:VAL:HG12	4	1.02
(1,416)	1:5:A:LEU:HD21	1:7:A:VAL:HG13	11	1.02
(1,416)	1:5:A:LEU:HD23	1:7:A:VAL:HG13	26	1.02
(1,353)	1:22:A:LEU:HD13	1:22:A:LEU:HB2	4	1.02
(1,353)	1:22:A:LEU:HD13	1:22:A:LEU:HB2	7	1.02
(1,353)	1:22:A:LEU:HD11	1:22:A:LEU:HB2	12	1.02
(1,353)	1:22:A:LEU:HD13	1:22:A:LEU:HB2	14	1.02
(1,353)	1:22:A:LEU:HD12	1:22:A:LEU:HB2	21	1.02
(1,228)	1:51:A:GLY:H	1:52:A:ASN:HD22	5	1.02
(1,193)	1:20:A:THR:H	1:20:A:THR:HG22	3	1.02
(1,193)	1:20:A:THR:H	1:20:A:THR:HG22	17	1.02
(1,193)	1:20:A:THR:H	1:20:A:THR:HG22	20	1.02
(1,62)	1:18:A:ASP:HB2	1:18:A:ASP:H	2	1.02
(1,46)	1:13:A:TYR:HB2	1:11:A:LYS:HD2	27	1.02
(1,958)	1:61:A:SER:H	1:5:A:LEU:HD23	5	1.01
(1,956)	1:5:A:LEU:HD21	1:41:A:TRP:HH2	30	1.01
(1,950)	1:45:A:ILE:HD13	1:45:A:ILE:H	12	1.01
(1,947)	1:45:A:ILE:HD12	1:45:A:ILE:H	18	1.01
(1,931)	1:6:A:GLN:HE21	1:8:A:ARG:HG2	27	1.01
(1,928)	1:9:A:ALA:HB1	1:11:A:LYS:H	9	1.01
(1,928)	1:9:A:ALA:HB2	1:11:A:LYS:H	11	1.01
(1,922)	1:66:A:ILE:H	1:6:A:GLN:HB3	6	1.01
(1,918)	1:13:A:TYR:H	1:11:A:LYS:HD3	8	1.01
(1,912)	1:34:A:GLU:HB3	1:35:A:GLN:H	18	1.01
(1,899)	1:27:A:GLY:H	1:28:A:ASP:HB2	19	1.01
(1,850)	1:11:A:LYS:HG3	1:13:A:TYR:HE1	16	1.01
(1,848)	1:16:A:ASN:HB2	1:18:A:ASP:H	25	1.01
(1,823)	1:28:A:ASP:H	1:9:A:ALA:HB2	28	1.01
(1,822)	1:26:A:ALA:HA	1:28:A:ASP:H	14	1.01
(1,813)	1:22:A:LEU:HB2	1:13:A:TYR:H	9	1.01
(1,812)	1:30:A:ILE:HG23	1:33:A:LEU:H	8	1.01
(1,617)	1:21:A:SER:H	1:21:A:SER:HB3	3	1.01
(1,617)	1:21:A:SER:H	1:21:A:SER:HB3	14	1.01

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,610)	1:58:A:PHE:H	1:57:A:TYR:HB2	7	1.01
(1,610)	1:58:A:PHE:H	1:57:A:TYR:HB2	27	1.01
(1,547)	1:6:A:GLN:H	1:5:A:LEU:HB3	3	1.01
(1,547)	1:6:A:GLN:H	1:5:A:LEU:HB3	4	1.01
(1,547)	1:6:A:GLN:H	1:5:A:LEU:HB3	12	1.01
(1,468)	1:30:A:ILE:H	1:30:A:ILE:HG12	26	1.01
(1,431)	1:22:A:LEU:HD12	1:24:A:VAL:HG13	9	1.01
(1,353)	1:22:A:LEU:HD13	1:22:A:LEU:HB2	6	1.01
(1,353)	1:22:A:LEU:HD11	1:22:A:LEU:HB2	9	1.01
(1,353)	1:22:A:LEU:HD11	1:22:A:LEU:HB2	15	1.01
(1,353)	1:22:A:LEU:HD12	1:22:A:LEU:HB2	20	1.01
(1,353)	1:22:A:LEU:HD13	1:22:A:LEU:HB2	27	1.01
(1,353)	1:22:A:LEU:HD12	1:22:A:LEU:HB2	28	1.01
(1,193)	1:20:A:THR:H	1:20:A:THR:HG22	2	1.01
(1,193)	1:20:A:THR:H	1:20:A:THR:HG22	12	1.01
(1,193)	1:20:A:THR:H	1:20:A:THR:HG23	21	1.01
(1,193)	1:20:A:THR:H	1:20:A:THR:HG23	30	1.01
(1,62)	1:18:A:ASP:HB2	1:18:A:ASP:H	15	1.01
(1,963)	1:30:A:ILE:HD13	1:22:A:LEU:H	28	1.0
(1,947)	1:7:A:VAL:HG22	1:9:A:ALA:H	14	1.0
(1,934)	1:26:A:ALA:H	1:25:A:LYS:HG3	24	1.0
(1,928)	1:9:A:ALA:HB1	1:11:A:LYS:H	29	1.0
(1,926)	1:6:A:GLN:H	1:65:A:ALA:HB3	20	1.0
(1,922)	1:66:A:ILE:H	1:6:A:GLN:HB3	28	1.0
(1,912)	1:34:A:GLU:HB3	1:35:A:GLN:H	20	1.0
(1,893)	1:41:A:TRP:HZ3	1:61:A:SER:HB2	1	1.0
(1,892)	1:13:A:TYR:HD2	1:56:A:GLY:HA3	15	1.0
(1,888)	1:5:A:LEU:HB3	1:67:A:VAL:HA	2	1.0
(1,887)	1:29:A:ILE:HG22	1:31:A:THR:HB	8	1.0
(1,863)	1:9:A:ALA:H	1:29:A:ILE:HA	10	1.0
(1,852)	1:13:A:TYR:HE2	1:24:A:VAL:H	2	1.0
(1,812)	1:30:A:ILE:HG22	1:33:A:LEU:H	5	1.0
(1,812)	1:30:A:ILE:HG22	1:33:A:LEU:H	19	1.0
(1,802)	1:26:A:ALA:HB1	1:11:A:LYS:HA	8	1.0
(1,743)	1:63:A:GLY:H	1:62:A:LEU:HG	9	1.0
(1,734)	1:39:A:GLY:H	1:40:A:ARG:HB2	27	1.0
(1,617)	1:21:A:SER:H	1:21:A:SER:HB3	2	1.0
(1,617)	1:21:A:SER:H	1:21:A:SER:HB3	5	1.0
(1,617)	1:21:A:SER:H	1:21:A:SER:HB3	9	1.0
(1,617)	1:21:A:SER:H	1:21:A:SER:HB3	12	1.0
(1,617)	1:21:A:SER:H	1:21:A:SER:HB3	13	1.0
(1,617)	1:21:A:SER:H	1:21:A:SER:HB3	19	1.0

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,617)	1:21:A:SER:H	1:21:A:SER:HB3	30	1.0
(1,547)	1:6:A:GLN:H	1:5:A:LEU:HB3	28	1.0
(1,416)	1:5:A:LEU:HD23	1:7:A:VAL:HG12	24	1.0
(1,353)	1:22:A:LEU:HD13	1:22:A:LEU:HB2	19	1.0
(1,224)	1:35:A:GLN:HB2	1:41:A:TRP:HD1	29	1.0
(1,211)	1:25:A:LYS:H	1:11:A:LYS:H	22	1.0
(1,193)	1:20:A:THR:H	1:20:A:THR:HG21	7	1.0
(1,193)	1:20:A:THR:H	1:20:A:THR:HG23	8	1.0
(1,193)	1:20:A:THR:H	1:20:A:THR:HG23	15	1.0
(1,140)	1:9:A:ALA:HA	1:26:A:ALA:HA	4	1.0
(1,959)	1:31:A:THR:H	1:22:A:LEU:HD13	28	0.99
(1,947)	1:45:A:ILE:HD13	1:45:A:ILE:H	22	0.99
(1,934)	1:26:A:ALA:H	1:25:A:LYS:HG3	29	0.99
(1,928)	1:9:A:ALA:HB1	1:11:A:LYS:H	14	0.99
(1,928)	1:60:A:SER:HB3	1:41:A:TRP:HH2	21	0.99
(1,928)	1:9:A:ALA:HB3	1:11:A:LYS:H	23	0.99
(1,922)	1:66:A:ILE:H	1:6:A:GLN:HB3	1	0.99
(1,894)	1:23:A:ASN:H	1:54:A:ARG:HD2	4	0.99
(1,847)	1:8:A:ARG:HD3	1:6:A:GLN:HE21	21	0.99
(1,813)	1:22:A:LEU:HB2	1:13:A:TYR:H	5	0.99
(1,812)	1:30:A:ILE:HG21	1:33:A:LEU:H	16	0.99
(1,743)	1:63:A:GLY:H	1:62:A:LEU:HG	8	0.99
(1,617)	1:21:A:SER:H	1:21:A:SER:HB3	15	0.99
(1,617)	1:21:A:SER:H	1:21:A:SER:HB3	16	0.99
(1,605)	1:57:A:TYR:H	1:21:A:SER:HA	22	0.99
(1,571)	1:66:A:ILE:H	1:65:A:ALA:HB2	26	0.99
(1,547)	1:6:A:GLN:H	1:5:A:LEU:HB3	30	0.99
(1,444)	1:30:A:ILE:HD13	1:32:A:VAL:HG11	25	0.99
(1,340)	1:63:A:GLY:H	1:7:A:VAL:HB	30	0.99
(1,228)	1:51:A:GLY:H	1:52:A:ASN:HD22	29	0.99
(1,193)	1:20:A:THR:H	1:20:A:THR:HG22	18	0.99
(1,25)	1:43:A:GLY:HA2	1:33:A:LEU:HD11	7	0.99
(1,942)	1:32:A:VAL:HB	1:7:A:VAL:HG21	5	0.98
(1,926)	1:6:A:GLN:H	1:65:A:ALA:HB3	21	0.98
(1,925)	1:11:A:LYS:HG2	1:13:A:TYR:H	15	0.98
(1,919)	1:23:A:ASN:HD21	1:25:A:LYS:HB3	12	0.98
(1,870)	1:29:A:ILE:HG21	1:45:A:ILE:HA	11	0.98
(1,823)	1:28:A:ASP:H	1:9:A:ALA:HB3	12	0.98
(1,813)	1:22:A:LEU:HB2	1:13:A:TYR:H	28	0.98
(1,797)	1:7:A:VAL:HB	1:63:A:GLY:HA2	4	0.98
(1,775)	1:38:A:ASP:H	1:40:A:ARG:H	17	0.98
(1,737)	1:50:A:THR:H	1:50:A:THR:HG22	15	0.98

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,617)	1:21:A:SER:H	1:21:A:SER:HB3	4	0.98
(1,617)	1:21:A:SER:H	1:21:A:SER:HB3	7	0.98
(1,617)	1:21:A:SER:H	1:21:A:SER:HB3	17	0.98
(1,617)	1:21:A:SER:H	1:21:A:SER:HB3	21	0.98
(1,617)	1:21:A:SER:H	1:21:A:SER:HB3	28	0.98
(1,610)	1:58:A:PHE:H	1:57:A:TYR:HB2	14	0.98
(1,547)	1:6:A:GLN:H	1:5:A:LEU:HB3	13	0.98
(1,536)	1:23:A:ASN:H	1:22:A:LEU:HD11	9	0.98
(1,440)	1:30:A:ILE:HG23	1:32:A:VAL:HG22	6	0.98
(1,427)	1:20:A:THR:HG22	1:55:A:VAL:HG21	1	0.98
(1,357)	1:65:A:ALA:HB1	1:5:A:LEU:HB3	29	0.98
(1,353)	1:22:A:LEU:HD12	1:22:A:LEU:HB2	17	0.98
(1,300)	1:4:A:MET:H	1:3:A:HIS:HB3	17	0.98
(1,226)	1:39:A:GLY:HA3	1:41:A:TRP:HD1	6	0.98
(1,62)	1:18:A:ASP:HB2	1:18:A:ASP:H	3	0.98
(1,62)	1:18:A:ASP:HB2	1:18:A:ASP:H	9	0.98
(1,956)	1:24:A:VAL:HG23	1:13:A:TYR:HD2	3	0.97
(1,956)	1:24:A:VAL:HG22	1:13:A:TYR:HD2	12	0.97
(1,948)	1:6:A:GLN:H	1:67:A:VAL:HG11	9	0.97
(1,947)	1:45:A:ILE:HD13	1:45:A:ILE:H	30	0.97
(1,928)	1:9:A:ALA:HB3	1:11:A:LYS:H	2	0.97
(1,927)	1:41:A:TRP:HD1	1:40:A:ARG:HB3	5	0.97
(1,926)	1:6:A:GLN:H	1:65:A:ALA:HB1	27	0.97
(1,922)	1:66:A:ILE:H	1:6:A:GLN:HB3	5	0.97
(1,912)	1:34:A:GLU:HB3	1:35:A:GLN:H	4	0.97
(1,900)	1:14:A:CYS:HB3	1:24:A:VAL:H	22	0.97
(1,894)	1:23:A:ASN:H	1:54:A:ARG:HD2	1	0.97
(1,852)	1:13:A:TYR:HE2	1:24:A:VAL:H	13	0.97
(1,852)	1:13:A:TYR:HE2	1:24:A:VAL:H	20	0.97
(1,823)	1:28:A:ASP:H	1:9:A:ALA:HB3	1	0.97
(1,823)	1:28:A:ASP:H	1:25:A:LYS:HG3	15	0.97
(1,800)	1:44:A:CYS:HB3	1:55:A:VAL:HA	13	0.97
(1,793)	1:28:A:ASP:HB3	1:45:A:ILE:HG22	20	0.97
(1,705)	1:35:A:GLN:HE22	1:33:A:LEU:HD23	21	0.97
(1,547)	1:6:A:GLN:H	1:5:A:LEU:HB3	5	0.97
(1,536)	1:23:A:ASN:H	1:22:A:LEU:HD13	1	0.97
(1,536)	1:23:A:ASN:H	1:22:A:LEU:HD12	17	0.97
(1,536)	1:23:A:ASN:H	1:22:A:LEU:HD13	18	0.97
(1,536)	1:23:A:ASN:H	1:22:A:LEU:HD12	20	0.97
(1,440)	1:30:A:ILE:HG21	1:32:A:VAL:HG23	7	0.97
(1,369)	1:29:A:ILE:HG13	1:6:A:GLN:HE22	29	0.97
(1,357)	1:65:A:ALA:HB1	1:5:A:LEU:HB3	25	0.97

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,299)	1:50:A:THR:H	1:47:A:ASP:HB2	29	0.97
(1,193)	1:20:A:THR:H	1:20:A:THR:HG23	9	0.97
(1,193)	1:20:A:THR:H	1:20:A:THR:HG22	11	0.97
(1,113)	1:31:A:THR:HB	1:44:A:CYS:HB2	16	0.97
(1,43)	1:13:A:TYR:HB2	1:24:A:VAL:HG22	19	0.97
(1,959)	1:31:A:THR:H	1:22:A:LEU:HD12	9	0.96
(1,941)	1:33:A:LEU:H	1:33:A:LEU:HG	8	0.96
(1,893)	1:41:A:TRP:HZ3	1:61:A:SER:HB2	18	0.96
(1,852)	1:25:A:LYS:H	1:48:A:ASN:HD22	6	0.96
(1,823)	1:28:A:ASP:H	1:9:A:ALA:HB1	4	0.96
(1,823)	1:28:A:ASP:H	1:9:A:ALA:HB3	6	0.96
(1,802)	1:26:A:ALA:HB3	1:12:A:ASP:HA	17	0.96
(1,797)	1:7:A:VAL:HB	1:63:A:GLY:HA2	5	0.96
(1,587)	1:53:A:ASP:H	1:53:A:ASP:HB2	9	0.96
(1,587)	1:53:A:ASP:H	1:53:A:ASP:HB2	15	0.96
(1,547)	1:6:A:GLN:H	1:5:A:LEU:HB3	9	0.96
(1,547)	1:6:A:GLN:H	1:5:A:LEU:HB3	22	0.96
(1,444)	1:30:A:ILE:HD13	1:32:A:VAL:HG12	2	0.96
(1,440)	1:30:A:ILE:HG21	1:32:A:VAL:HG21	29	0.96
(1,399)	1:29:A:ILE:H	1:29:A:ILE:HD12	5	0.96
(1,322)	1:58:A:PHE:HE1	1:58:A:PHE:HB3	12	0.96
(1,322)	1:58:A:PHE:HE1	1:58:A:PHE:HB3	26	0.96
(1,317)	1:39:A:GLY:H	1:38:A:ASP:HB3	17	0.96
(1,193)	1:20:A:THR:H	1:20:A:THR:HG21	23	0.96
(1,190)	1:34:A:GLU:HG2	1:3:A:HIS:H	22	0.96
(1,46)	1:13:A:TYR:HB2	1:11:A:LYS:HD2	21	0.96
(1,967)	1:62:A:LEU:H	1:63:A:GLY:HA2	30	0.95
(1,958)	1:61:A:SER:H	1:5:A:LEU:HD23	21	0.95
(1,928)	1:60:A:SER:HB3	1:41:A:TRP:HH2	6	0.95
(1,922)	1:23:A:ASN:H	1:22:A:LEU:HB2	22	0.95
(1,912)	1:34:A:GLU:HB3	1:35:A:GLN:H	6	0.95
(1,872)	1:33:A:LEU:HD12	1:33:A:LEU:HA	29	0.95
(1,863)	1:9:A:ALA:H	1:29:A:ILE:HA	21	0.95
(1,813)	1:22:A:LEU:HB2	1:13:A:TYR:H	23	0.95
(1,802)	1:26:A:ALA:HB3	1:12:A:ASP:HA	5	0.95
(1,797)	1:7:A:VAL:HB	1:64:A:GLU:HA	16	0.95
(1,587)	1:53:A:ASP:H	1:53:A:ASP:HB2	8	0.95
(1,571)	1:66:A:ILE:H	1:65:A:ALA:HB1	6	0.95
(1,554)	1:54:A:ARG:H	1:54:A:ARG:HB2	2	0.95
(1,547)	1:6:A:GLN:H	1:5:A:LEU:HB3	7	0.95
(1,536)	1:23:A:ASN:H	1:22:A:LEU:HD13	14	0.95
(1,493)	1:41:A:TRP:HE1	1:40:A:ARG:H	8	0.95

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,444)	1:30:A:ILE:HD11	1:32:A:VAL:HG13	19	0.95
(1,440)	1:30:A:ILE:HG21	1:32:A:VAL:HG23	3	0.95
(1,440)	1:30:A:ILE:HG23	1:32:A:VAL:HG23	17	0.95
(1,440)	1:30:A:ILE:HG22	1:32:A:VAL:HG21	19	0.95
(1,432)	1:22:A:LEU:HD11	1:24:A:VAL:HG21	30	0.95
(1,395)	1:42:A:LYS:H	1:33:A:LEU:HD13	15	0.95
(1,382)	1:65:A:ALA:H	1:5:A:LEU:HB3	7	0.95
(1,382)	1:65:A:ALA:H	1:5:A:LEU:HB3	14	0.95
(1,193)	1:20:A:THR:H	1:20:A:THR:HG23	27	0.95
(1,66)	1:28:A:ASP:HB3	1:25:A:LYS:HG2	4	0.95
(1,46)	1:13:A:TYR:HB2	1:11:A:LYS:HD2	1	0.95
(1,27)	1:43:A:GLY:HA3	1:33:A:LEU:HD12	20	0.95
(1,963)	1:30:A:ILE:HD13	1:22:A:LEU:H	6	0.94
(1,959)	1:31:A:THR:H	1:22:A:LEU:HD11	26	0.94
(1,958)	1:61:A:SER:H	1:5:A:LEU:HD21	8	0.94
(1,956)	1:24:A:VAL:HG21	1:13:A:TYR:HD2	4	0.94
(1,947)	1:45:A:ILE:HD11	1:45:A:ILE:H	6	0.94
(1,929)	1:13:A:TYR:HD1	1:11:A:LYS:HG2	1	0.94
(1,928)	1:9:A:ALA:HB1	1:11:A:LYS:H	7	0.94
(1,928)	1:9:A:ALA:HB2	1:11:A:LYS:H	8	0.94
(1,928)	1:9:A:ALA:HB3	1:11:A:LYS:H	13	0.94
(1,928)	1:9:A:ALA:HB1	1:11:A:LYS:H	18	0.94
(1,925)	1:11:A:LYS:HG2	1:13:A:TYR:H	14	0.94
(1,912)	1:41:A:TRP:H	1:40:A:ARG:HG3	10	0.94
(1,912)	1:34:A:GLU:HB3	1:35:A:GLN:H	23	0.94
(1,893)	1:41:A:TRP:HZ3	1:32:A:VAL:HA	24	0.94
(1,872)	1:33:A:LEU:HD12	1:33:A:LEU:HA	28	0.94
(1,853)	1:29:A:ILE:HG12	1:48:A:ASN:HD22	13	0.94
(1,853)	1:25:A:LYS:HB2	1:48:A:ASN:HD22	18	0.94
(1,851)	1:13:A:TYR:HE2	1:57:A:TYR:H	8	0.94
(1,838)	1:36:A:HIS:HB2	1:39:A:GLY:H	15	0.94
(1,834)	1:39:A:GLY:HA3	1:41:A:TRP:H	5	0.94
(1,834)	1:51:A:GLY:HA3	1:47:A:ASP:H	21	0.94
(1,795)	1:15:A:ASN:HD22	1:15:A:ASN:HB3	13	0.94
(1,737)	1:50:A:THR:H	1:50:A:THR:HG22	9	0.94
(1,737)	1:50:A:THR:H	1:50:A:THR:HG21	21	0.94
(1,617)	1:21:A:SER:H	1:21:A:SER:HB3	23	0.94
(1,587)	1:53:A:ASP:H	1:53:A:ASP:HB2	5	0.94
(1,587)	1:53:A:ASP:H	1:53:A:ASP:HB2	21	0.94
(1,572)	1:66:A:ILE:H	1:66:A:ILE:HG12	7	0.94
(1,547)	1:6:A:GLN:H	1:5:A:LEU:HB3	16	0.94
(1,547)	1:6:A:GLN:H	1:5:A:LEU:HB3	24	0.94

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,536)	1:23:A:ASN:H	1:22:A:LEU:HD11	12	0.94
(1,528)	1:31:A:THR:H	1:31:A:THR:HG23	11	0.94
(1,493)	1:41:A:TRP:HE1	1:40:A:ARG:H	13	0.94
(1,493)	1:41:A:TRP:HE1	1:40:A:ARG:H	17	0.94
(1,485)	1:46:A:HIS:H	1:45:A:ILE:HD13	27	0.94
(1,420)	1:7:A:VAL:HG12	1:32:A:VAL:HG11	30	0.94
(1,416)	1:5:A:LEU:HD21	1:7:A:VAL:HG12	19	0.94
(1,360)	1:27:A:GLY:H	1:25:A:LYS:HG3	23	0.94
(1,322)	1:58:A:PHE:HE1	1:58:A:PHE:HB3	13	0.94
(1,300)	1:4:A:MET:H	1:3:A:HIS:HB3	24	0.94
(1,66)	1:28:A:ASP:HB3	1:25:A:LYS:HG2	18	0.94
(1,966)	1:8:A:ARG:H	1:64:A:GLU:HG3	6	0.93
(1,959)	1:31:A:THR:H	1:22:A:LEU:HD12	15	0.93
(1,950)	1:7:A:VAL:HG21	1:9:A:ALA:H	21	0.93
(1,947)	1:45:A:ILE:HD11	1:45:A:ILE:H	7	0.93
(1,934)	1:26:A:ALA:H	1:25:A:LYS:HG2	18	0.93
(1,925)	1:11:A:LYS:HG2	1:13:A:TYR:H	23	0.93
(1,912)	1:34:A:GLU:HB3	1:35:A:GLN:H	16	0.93
(1,892)	1:13:A:TYR:HD2	1:56:A:GLY:HA3	9	0.93
(1,859)	1:63:A:GLY:H	1:7:A:VAL:HA	27	0.93
(1,848)	1:16:A:ASN:HB2	1:18:A:ASP:H	3	0.93
(1,823)	1:28:A:ASP:H	1:9:A:ALA:HB1	3	0.93
(1,797)	1:7:A:VAL:HB	1:63:A:GLY:HA2	9	0.93
(1,797)	1:7:A:VAL:HB	1:63:A:GLY:HA2	23	0.93
(1,707)	1:35:A:GLN:HE22	1:35:A:GLN:HB3	27	0.93
(1,587)	1:53:A:ASP:H	1:53:A:ASP:HB2	7	0.93
(1,571)	1:66:A:ILE:H	1:65:A:ALA:HB1	27	0.93
(1,547)	1:6:A:GLN:H	1:5:A:LEU:HB3	19	0.93
(1,536)	1:23:A:ASN:H	1:22:A:LEU:HD13	24	0.93
(1,528)	1:31:A:THR:H	1:31:A:THR:HG22	17	0.93
(1,528)	1:31:A:THR:H	1:31:A:THR:HG23	19	0.93
(1,485)	1:46:A:HIS:H	1:45:A:ILE:HD12	25	0.93
(1,468)	1:30:A:ILE:H	1:30:A:ILE:HG12	15	0.93
(1,467)	1:30:A:ILE:H	1:29:A:ILE:HG22	18	0.93
(1,444)	1:30:A:ILE:HD11	1:32:A:VAL:HG11	11	0.93
(1,440)	1:30:A:ILE:HG21	1:32:A:VAL:HG22	9	0.93
(1,437)	1:24:A:VAL:HG22	1:30:A:ILE:HG21	6	0.93
(1,322)	1:58:A:PHE:HE1	1:58:A:PHE:HB3	2	0.93
(1,322)	1:58:A:PHE:HE1	1:58:A:PHE:HB3	27	0.93
(1,193)	1:20:A:THR:H	1:20:A:THR:HG23	29	0.93
(1,959)	1:31:A:THR:H	1:22:A:LEU:HD13	17	0.92
(1,947)	1:45:A:ILE:HD11	1:45:A:ILE:H	4	0.92

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,941)	1:32:A:VAL:HG13	1:33:A:LEU:H	9	0.92
(1,934)	1:26:A:ALA:H	1:25:A:LYS:HG2	1	0.92
(1,928)	1:9:A:ALA:HB2	1:11:A:LYS:H	1	0.92
(1,912)	1:41:A:TRP:H	1:40:A:ARG:HG3	22	0.92
(1,909)	1:41:A:TRP:HD1	1:40:A:ARG:HG2	21	0.92
(1,906)	1:34:A:GLU:HG2	1:3:A:HIS:H	19	0.92
(1,870)	1:29:A:ILE:HG22	1:45:A:ILE:HA	8	0.92
(1,863)	1:9:A:ALA:H	1:29:A:ILE:HA	18	0.92
(1,848)	1:16:A:ASN:HB2	1:18:A:ASP:H	27	0.92
(1,812)	1:30:A:ILE:HG22	1:33:A:LEU:H	30	0.92
(1,795)	1:15:A:ASN:HD22	1:15:A:ASN:HB3	19	0.92
(1,737)	1:50:A:THR:H	1:50:A:THR:HG23	16	0.92
(1,617)	1:21:A:SER:H	1:21:A:SER:HB3	29	0.92
(1,529)	1:31:A:THR:H	1:30:A:ILE:HG13	15	0.92
(1,529)	1:31:A:THR:H	1:30:A:ILE:HG13	16	0.92
(1,529)	1:31:A:THR:H	1:30:A:ILE:HG13	28	0.92
(1,528)	1:31:A:THR:H	1:31:A:THR:HG23	29	0.92
(1,485)	1:46:A:HIS:H	1:45:A:ILE:HD11	7	0.92
(1,468)	1:30:A:ILE:H	1:30:A:ILE:HG12	3	0.92
(1,444)	1:30:A:ILE:HD12	1:32:A:VAL:HG13	23	0.92
(1,440)	1:30:A:ILE:HG22	1:32:A:VAL:HG23	12	0.92
(1,437)	1:24:A:VAL:HG22	1:30:A:ILE:HG22	7	0.92
(1,437)	1:24:A:VAL:HG21	1:30:A:ILE:HG21	17	0.92
(1,330)	1:63:A:GLY:H	1:62:A:LEU:HB2	14	0.92
(1,322)	1:58:A:PHE:HE1	1:58:A:PHE:HB3	23	0.92
(1,322)	1:58:A:PHE:HE1	1:58:A:PHE:HB3	25	0.92
(1,322)	1:58:A:PHE:HE1	1:58:A:PHE:HB3	29	0.92
(1,62)	1:18:A:ASP:HB2	1:18:A:ASP:H	1	0.92
(1,39)	1:53:A:ASP:HB3	1:54:A:ARG:H	17	0.92
(1,963)	1:30:A:ILE:HD12	1:22:A:LEU:H	9	0.91
(1,950)	1:7:A:VAL:HG23	1:9:A:ALA:H	10	0.91
(1,947)	1:45:A:ILE:HD13	1:45:A:ILE:H	12	0.91
(1,941)	1:33:A:LEU:H	1:33:A:LEU:HG	2	0.91
(1,934)	1:26:A:ALA:H	1:25:A:LYS:HG3	19	0.91
(1,927)	1:48:A:ASN:HD22	1:25:A:LYS:HG2	15	0.91
(1,926)	1:6:A:GLN:H	1:65:A:ALA:HB3	16	0.91
(1,925)	1:11:A:LYS:HG2	1:13:A:TYR:H	2	0.91
(1,922)	1:66:A:ILE:H	1:6:A:GLN:HB3	2	0.91
(1,922)	1:66:A:ILE:H	1:6:A:GLN:HB3	17	0.91
(1,912)	1:34:A:GLU:HB3	1:35:A:GLN:H	13	0.91
(1,899)	1:27:A:GLY:H	1:28:A:ASP:HB2	22	0.91
(1,893)	1:41:A:TRP:HZ3	1:61:A:SER:HB2	5	0.91

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,870)	1:29:A:ILE:HG21	1:45:A:ILE:HA	20	0.91
(1,849)	1:13:A:TYR:H	1:15:A:ASN:HD22	17	0.91
(1,834)	1:51:A:GLY:HA3	1:47:A:ASP:H	10	0.91
(1,797)	1:7:A:VAL:HB	1:63:A:GLY:HA2	18	0.91
(1,795)	1:17:A:TYR:HB3	1:18:A:ASP:H	24	0.91
(1,687)	1:27:A:GLY:H	1:26:A:ALA:HB2	21	0.91
(1,617)	1:21:A:SER:H	1:21:A:SER:HB3	1	0.91
(1,617)	1:21:A:SER:H	1:21:A:SER:HB3	6	0.91
(1,571)	1:66:A:ILE:H	1:65:A:ALA:HB1	5	0.91
(1,547)	1:6:A:GLN:H	1:5:A:LEU:HB3	23	0.91
(1,541)	1:40:A:ARG:H	1:40:A:ARG:HD2	2	0.91
(1,536)	1:23:A:ASN:H	1:22:A:LEU:HD13	4	0.91
(1,536)	1:23:A:ASN:H	1:22:A:LEU:HD12	21	0.91
(1,536)	1:23:A:ASN:H	1:22:A:LEU:HD12	28	0.91
(1,529)	1:31:A:THR:H	1:30:A:ILE:HG13	3	0.91
(1,529)	1:31:A:THR:H	1:30:A:ILE:HG13	17	0.91
(1,529)	1:31:A:THR:H	1:30:A:ILE:HG13	26	0.91
(1,528)	1:31:A:THR:H	1:31:A:THR:HG23	2	0.91
(1,528)	1:31:A:THR:H	1:31:A:THR:HG21	4	0.91
(1,528)	1:31:A:THR:H	1:31:A:THR:HG23	6	0.91
(1,528)	1:31:A:THR:H	1:31:A:THR:HG22	7	0.91
(1,528)	1:31:A:THR:H	1:31:A:THR:HG22	12	0.91
(1,528)	1:31:A:THR:H	1:31:A:THR:HG21	22	0.91
(1,514)	1:65:A:ALA:H	1:64:A:GLU:HB2	10	0.91
(1,514)	1:65:A:ALA:H	1:64:A:GLU:HB2	29	0.91
(1,444)	1:30:A:ILE:HD12	1:32:A:VAL:HG13	8	0.91
(1,437)	1:24:A:VAL:HG22	1:30:A:ILE:HG22	28	0.91
(1,427)	1:20:A:THR:HG21	1:55:A:VAL:HG23	16	0.91
(1,346)	1:55:A:VAL:H	1:54:A:ARG:HB3	15	0.91
(1,322)	1:58:A:PHE:HE1	1:58:A:PHE:HB3	1	0.91
(1,322)	1:58:A:PHE:HE1	1:58:A:PHE:HB3	4	0.91
(1,322)	1:58:A:PHE:HE1	1:58:A:PHE:HB3	5	0.91
(1,322)	1:58:A:PHE:HE1	1:58:A:PHE:HB3	7	0.91
(1,322)	1:58:A:PHE:HE1	1:58:A:PHE:HB3	15	0.91
(1,322)	1:58:A:PHE:HE1	1:58:A:PHE:HB3	18	0.91
(1,322)	1:58:A:PHE:HE1	1:58:A:PHE:HB3	19	0.91
(1,322)	1:58:A:PHE:HE1	1:58:A:PHE:HB3	20	0.91
(1,322)	1:58:A:PHE:HE1	1:58:A:PHE:HB3	30	0.91
(1,210)	1:41:A:TRP:HZ3	1:61:A:SER:H	15	0.91
(1,58)	1:28:A:ASP:H	1:28:A:ASP:HB2	11	0.91
(1,39)	1:53:A:ASP:HB3	1:54:A:ARG:H	16	0.91
(1,963)	1:30:A:ILE:HD13	1:22:A:LEU:H	1	0.9

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,939)	1:58:A:PHE:HD1	1:32:A:VAL:HG12	2	0.9
(1,927)	1:48:A:ASN:HD22	1:25:A:LYS:HG2	17	0.9
(1,906)	1:6:A:GLN:H	1:6:A:GLN:HG2	15	0.9
(1,899)	1:27:A:GLY:H	1:28:A:ASP:HB2	1	0.9
(1,892)	1:41:A:TRP:HH2	1:61:A:SER:HA	3	0.9
(1,873)	1:41:A:TRP:HB3	1:34:A:GLU:HA	26	0.9
(1,853)	1:29:A:ILE:HG12	1:48:A:ASN:HD22	21	0.9
(1,840)	1:12:A:ASP:H	1:11:A:LYS:HD2	17	0.9
(1,797)	1:7:A:VAL:HB	1:63:A:GLY:HA2	15	0.9
(1,795)	1:17:A:TYR:HB3	1:18:A:ASP:H	28	0.9
(1,737)	1:50:A:THR:H	1:50:A:THR:HG22	1	0.9
(1,737)	1:50:A:THR:H	1:50:A:THR:HG21	26	0.9
(1,707)	1:35:A:GLN:HE22	1:35:A:GLN:HB3	26	0.9
(1,648)	1:12:A:ASP:H	1:12:A:ASP:HB3	4	0.9
(1,543)	1:40:A:ARG:H	1:40:A:ARG:HG2	19	0.9
(1,536)	1:23:A:ASN:H	1:22:A:LEU:HD13	7	0.9
(1,529)	1:31:A:THR:H	1:30:A:ILE:HG13	1	0.9
(1,529)	1:31:A:THR:H	1:30:A:ILE:HG13	9	0.9
(1,529)	1:31:A:THR:H	1:30:A:ILE:HG13	11	0.9
(1,529)	1:31:A:THR:H	1:30:A:ILE:HG13	25	0.9
(1,528)	1:31:A:THR:H	1:31:A:THR:HG23	30	0.9
(1,441)	1:30:A:ILE:HG22	1:45:A:ILE:HG23	17	0.9
(1,440)	1:30:A:ILE:HG21	1:32:A:VAL:HG22	16	0.9
(1,382)	1:65:A:ALA:H	1:5:A:LEU:HB3	1	0.9
(1,322)	1:58:A:PHE:HE1	1:58:A:PHE:HB3	3	0.9
(1,322)	1:58:A:PHE:HE1	1:58:A:PHE:HB3	10	0.9
(1,322)	1:58:A:PHE:HE1	1:58:A:PHE:HB3	14	0.9
(1,322)	1:58:A:PHE:HE1	1:58:A:PHE:HB3	16	0.9
(1,322)	1:58:A:PHE:HE1	1:58:A:PHE:HB3	28	0.9
(1,27)	1:43:A:GLY:HA3	1:33:A:LEU:HD13	13	0.9
(1,934)	1:26:A:ALA:H	1:25:A:LYS:HG2	9	0.89
(1,928)	1:9:A:ALA:HB3	1:11:A:LYS:H	3	0.89
(1,925)	1:11:A:LYS:HG2	1:13:A:TYR:H	4	0.89
(1,922)	1:23:A:ASN:H	1:22:A:LEU:HB2	25	0.89
(1,918)	1:13:A:TYR:H	1:11:A:LYS:HD3	9	0.89
(1,906)	1:6:A:GLN:H	1:6:A:GLN:HG2	14	0.89
(1,900)	1:14:A:CYS:HB3	1:24:A:VAL:H	11	0.89
(1,899)	1:27:A:GLY:H	1:28:A:ASP:HB2	4	0.89
(1,893)	1:41:A:TRP:HZ3	1:61:A:SER:HB2	7	0.89
(1,892)	1:13:A:TYR:HD2	1:56:A:GLY:HA3	27	0.89
(1,887)	1:29:A:ILE:HG23	1:31:A:THR:HB	7	0.89
(1,802)	1:26:A:ALA:HB3	1:11:A:LYS:HA	10	0.89

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,800)	1:44:A:CYS:HB3	1:55:A:VAL:HA	19	0.89
(1,628)	1:38:A:ASP:H	1:36:A:HIS:HB2	26	0.89
(1,536)	1:23:A:ASN:H	1:22:A:LEU:HD13	19	0.89
(1,536)	1:23:A:ASN:H	1:22:A:LEU:HD13	26	0.89
(1,529)	1:31:A:THR:H	1:30:A:ILE:HG13	6	0.89
(1,529)	1:31:A:THR:H	1:30:A:ILE:HG13	21	0.89
(1,529)	1:31:A:THR:H	1:30:A:ILE:HG13	29	0.89
(1,528)	1:31:A:THR:H	1:31:A:THR:HG22	10	0.89
(1,528)	1:31:A:THR:H	1:31:A:THR:HG21	21	0.89
(1,528)	1:31:A:THR:H	1:31:A:THR:HG21	28	0.89
(1,485)	1:46:A:HIS:H	1:45:A:ILE:HD13	12	0.89
(1,440)	1:30:A:ILE:HG22	1:32:A:VAL:HG21	26	0.89
(1,431)	1:22:A:LEU:HD11	1:24:A:VAL:HG11	1	0.89
(1,431)	1:22:A:LEU:HD11	1:24:A:VAL:HG12	14	0.89
(1,322)	1:58:A:PHE:HE1	1:58:A:PHE:HB3	8	0.89
(1,322)	1:58:A:PHE:HE1	1:58:A:PHE:HB3	17	0.89
(1,322)	1:58:A:PHE:HE1	1:58:A:PHE:HB3	22	0.89
(1,963)	1:30:A:ILE:HD11	1:22:A:LEU:H	12	0.88
(1,956)	1:5:A:LEU:HD23	1:41:A:TRP:HH2	24	0.88
(1,934)	1:26:A:ALA:H	1:25:A:LYS:HG3	23	0.88
(1,919)	1:23:A:ASN:HD21	1:25:A:LYS:HB3	6	0.88
(1,887)	1:29:A:ILE:HG23	1:31:A:THR:HB	13	0.88
(1,802)	1:26:A:ALA:HB2	1:11:A:LYS:HA	11	0.88
(1,734)	1:39:A:GLY:H	1:40:A:ARG:HB2	25	0.88
(1,536)	1:23:A:ASN:H	1:22:A:LEU:HD11	15	0.88
(1,536)	1:23:A:ASN:H	1:22:A:LEU:HD13	27	0.88
(1,529)	1:31:A:THR:H	1:30:A:ILE:HG13	18	0.88
(1,529)	1:31:A:THR:H	1:30:A:ILE:HG13	24	0.88
(1,529)	1:31:A:THR:H	1:30:A:ILE:HG13	30	0.88
(1,528)	1:31:A:THR:H	1:31:A:THR:HG21	9	0.88
(1,528)	1:31:A:THR:H	1:31:A:THR:HG21	18	0.88
(1,528)	1:31:A:THR:H	1:31:A:THR:HG22	25	0.88
(1,485)	1:46:A:HIS:H	1:45:A:ILE:HD13	22	0.88
(1,437)	1:24:A:VAL:HG22	1:30:A:ILE:HG21	8	0.88
(1,324)	1:4:A:MET:H	1:4:A:MET:HG3	18	0.88
(1,322)	1:58:A:PHE:HE1	1:58:A:PHE:HB3	11	0.88
(1,322)	1:58:A:PHE:HE1	1:58:A:PHE:HB3	21	0.88
(1,39)	1:53:A:ASP:HB3	1:54:A:ARG:H	22	0.88
(1,27)	1:43:A:GLY:HA3	1:33:A:LEU:HD12	12	0.88
(1,956)	1:24:A:VAL:HG23	1:13:A:TYR:HD2	21	0.87
(1,941)	1:33:A:LEU:H	1:33:A:LEU:HG	1	0.87
(1,939)	1:58:A:PHE:HD1	1:32:A:VAL:HG13	24	0.87

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,928)	1:60:A:SER:HB3	1:41:A:TRP:HH2	20	0.87
(1,922)	1:66:A:ILE:H	1:6:A:GLN:HB3	15	0.87
(1,906)	1:6:A:GLN:H	1:6:A:GLN:HG2	22	0.87
(1,888)	1:5:A:LEU:HB3	1:67:A:VAL:HA	22	0.87
(1,863)	1:9:A:ALA:H	1:29:A:ILE:HA	11	0.87
(1,863)	1:9:A:ALA:H	1:29:A:ILE:HA	20	0.87
(1,859)	1:67:A:VAL:H	1:7:A:VAL:HA	16	0.87
(1,840)	1:11:A:LYS:HD3	1:12:A:ASP:H	21	0.87
(1,797)	1:7:A:VAL:HB	1:63:A:GLY:HA2	8	0.87
(1,685)	1:27:A:GLY:H	1:24:A:VAL:HG22	8	0.87
(1,529)	1:31:A:THR:H	1:30:A:ILE:HG13	12	0.87
(1,529)	1:31:A:THR:H	1:30:A:ILE:HG13	13	0.87
(1,529)	1:31:A:THR:H	1:30:A:ILE:HG13	14	0.87
(1,529)	1:31:A:THR:H	1:30:A:ILE:HG13	22	0.87
(1,529)	1:31:A:THR:H	1:30:A:ILE:HG13	23	0.87
(1,529)	1:31:A:THR:H	1:30:A:ILE:HG13	27	0.87
(1,528)	1:31:A:THR:H	1:31:A:THR:HG21	23	0.87
(1,528)	1:31:A:THR:H	1:31:A:THR:HG23	26	0.87
(1,444)	1:30:A:ILE:HD13	1:32:A:VAL:HG11	13	0.87
(1,434)	1:24:A:VAL:HG11	1:30:A:ILE:HD13	14	0.87
(1,420)	1:7:A:VAL:HG13	1:32:A:VAL:HG12	4	0.87
(1,416)	1:5:A:LEU:HD22	1:7:A:VAL:HG12	18	0.87
(1,382)	1:65:A:ALA:H	1:5:A:LEU:HB3	6	0.87
(1,358)	1:41:A:TRP:H	1:40:A:ARG:HB2	19	0.87
(1,357)	1:65:A:ALA:HB3	1:5:A:LEU:HB3	11	0.87
(1,346)	1:55:A:VAL:H	1:54:A:ARG:HB3	8	0.87
(1,322)	1:58:A:PHE:HE1	1:58:A:PHE:HB3	6	0.87
(1,322)	1:58:A:PHE:HE1	1:58:A:PHE:HB3	24	0.87
(1,300)	1:4:A:MET:H	1:3:A:HIS:HB3	3	0.87
(1,225)	1:35:A:GLN:HG2	1:41:A:TRP:HD1	22	0.87
(1,39)	1:53:A:ASP:HB3	1:54:A:ARG:H	28	0.87
(1,959)	1:31:A:THR:H	1:22:A:LEU:HD11	3	0.86
(1,941)	1:33:A:LEU:H	1:33:A:LEU:HG	3	0.86
(1,941)	1:33:A:LEU:H	1:33:A:LEU:HG	4	0.86
(1,941)	1:33:A:LEU:H	1:33:A:LEU:HG	13	0.86
(1,939)	1:58:A:PHE:HD1	1:32:A:VAL:HG11	28	0.86
(1,928)	1:9:A:ALA:HB2	1:11:A:LYS:H	5	0.86
(1,927)	1:41:A:TRP:HD1	1:40:A:ARG:HB2	4	0.86
(1,922)	1:66:A:ILE:H	1:6:A:GLN:HB3	16	0.86
(1,906)	1:6:A:GLN:H	1:6:A:GLN:HG2	12	0.86
(1,906)	1:6:A:GLN:H	1:6:A:GLN:HG2	13	0.86
(1,887)	1:29:A:ILE:HG21	1:31:A:THR:HB	11	0.86

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,887)	1:29:A:ILE:HG21	1:31:A:THR:HB	29	0.86
(1,847)	1:8:A:ARG:HD3	1:6:A:GLN:HE21	26	0.86
(1,797)	1:7:A:VAL:HB	1:63:A:GLY:HA2	22	0.86
(1,737)	1:50:A:THR:H	1:50:A:THR:HG23	18	0.86
(1,737)	1:50:A:THR:H	1:50:A:THR:HG23	23	0.86
(1,687)	1:27:A:GLY:H	1:26:A:ALA:HB2	12	0.86
(1,597)	1:62:A:LEU:H	1:62:A:LEU:HB3	3	0.86
(1,571)	1:66:A:ILE:H	1:65:A:ALA:HB3	18	0.86
(1,541)	1:40:A:ARG:H	1:40:A:ARG:HD2	8	0.86
(1,536)	1:23:A:ASN:H	1:22:A:LEU:HD13	3	0.86
(1,536)	1:23:A:ASN:H	1:22:A:LEU:HD12	10	0.86
(1,529)	1:31:A:THR:H	1:30:A:ILE:HG13	7	0.86
(1,529)	1:31:A:THR:H	1:30:A:ILE:HG13	8	0.86
(1,528)	1:31:A:THR:H	1:31:A:THR:HG23	20	0.86
(1,444)	1:30:A:ILE:HD12	1:32:A:VAL:HG13	21	0.86
(1,440)	1:30:A:ILE:HG22	1:32:A:VAL:HG21	25	0.86
(1,322)	1:58:A:PHE:HE1	1:58:A:PHE:HB3	9	0.86
(1,305)	1:13:A:TYR:HD1	1:11:A:LYS:HE3	24	0.86
(1,39)	1:53:A:ASP:HB3	1:54:A:ARG:H	2	0.86
(1,953)	1:23:A:ASN:HB3	1:45:A:ILE:HG21	30	0.85
(1,941)	1:32:A:VAL:HG12	1:33:A:LEU:H	5	0.85
(1,934)	1:26:A:ALA:H	1:25:A:LYS:HG3	12	0.85
(1,934)	1:26:A:ALA:H	1:25:A:LYS:HG3	22	0.85
(1,925)	1:11:A:LYS:HG2	1:13:A:TYR:H	29	0.85
(1,912)	1:34:A:GLU:HB3	1:35:A:GLN:H	30	0.85
(1,902)	1:28:A:ASP:H	1:28:A:ASP:HB2	11	0.85
(1,852)	1:13:A:TYR:HE2	1:24:A:VAL:H	4	0.85
(1,823)	1:28:A:ASP:H	1:25:A:LYS:HG3	5	0.85
(1,802)	1:26:A:ALA:HB1	1:12:A:ASP:HA	16	0.85
(1,797)	1:7:A:VAL:HB	1:63:A:GLY:HA2	1	0.85
(1,797)	1:7:A:VAL:HB	1:63:A:GLY:HA2	3	0.85
(1,737)	1:50:A:THR:H	1:50:A:THR:HG23	4	0.85
(1,737)	1:50:A:THR:H	1:50:A:THR:HG23	13	0.85
(1,737)	1:50:A:THR:H	1:50:A:THR:HG22	28	0.85
(1,707)	1:35:A:GLN:HE22	1:35:A:GLN:HB3	8	0.85
(1,544)	1:40:A:ARG:H	1:40:A:ARG:HB2	6	0.85
(1,536)	1:23:A:ASN:H	1:22:A:LEU:HD13	6	0.85
(1,536)	1:23:A:ASN:H	1:22:A:LEU:HD12	16	0.85
(1,529)	1:31:A:THR:H	1:30:A:ILE:HG13	4	0.85
(1,529)	1:31:A:THR:H	1:30:A:ILE:HG13	5	0.85
(1,529)	1:31:A:THR:H	1:30:A:ILE:HG13	19	0.85
(1,529)	1:31:A:THR:H	1:30:A:ILE:HG13	20	0.85

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,528)	1:31:A:THR:H	1:31:A:THR:HG22	1	0.85
(1,528)	1:31:A:THR:H	1:31:A:THR:HG21	5	0.85
(1,528)	1:31:A:THR:H	1:31:A:THR:HG21	14	0.85
(1,528)	1:31:A:THR:H	1:31:A:THR:HG22	24	0.85
(1,514)	1:65:A:ALA:H	1:64:A:GLU:HB2	9	0.85
(1,493)	1:41:A:TRP:HE1	1:40:A:ARG:H	1	0.85
(1,485)	1:46:A:HIS:H	1:45:A:ILE:HD11	6	0.85
(1,467)	1:30:A:ILE:H	1:29:A:ILE:HG23	15	0.85
(1,444)	1:30:A:ILE:HD11	1:32:A:VAL:HG12	28	0.85
(1,440)	1:30:A:ILE:HG21	1:32:A:VAL:HG22	1	0.85
(1,440)	1:30:A:ILE:HG22	1:32:A:VAL:HG21	2	0.85
(1,416)	1:5:A:LEU:HD21	1:7:A:VAL:HG12	23	0.85
(1,403)	1:58:A:PHE:H	1:22:A:LEU:HD11	17	0.85
(1,397)	1:33:A:LEU:H	1:33:A:LEU:HD12	8	0.85
(1,360)	1:27:A:GLY:H	1:25:A:LYS:HG2	9	0.85
(1,357)	1:65:A:ALA:HB3	1:5:A:LEU:HB3	21	0.85
(1,347)	1:63:A:GLY:H	1:62:A:LEU:HB3	14	0.85
(1,174)	1:30:A:ILE:HG12	1:9:A:ALA:H	2	0.85
(1,174)	1:30:A:ILE:HG12	1:9:A:ALA:H	13	0.85
(1,39)	1:54:A:ARG:H	1:53:A:ASP:HB2	1	0.85
(1,27)	1:43:A:GLY:HA3	1:33:A:LEU:HD12	10	0.85
(1,966)	1:8:A:ARG:H	1:64:A:GLU:HG3	1	0.84
(1,966)	1:8:A:ARG:H	1:64:A:GLU:HG3	24	0.84
(1,959)	1:31:A:THR:H	1:22:A:LEU:HD13	5	0.84
(1,953)	1:23:A:ASN:HB2	1:45:A:ILE:HG21	19	0.84
(1,941)	1:33:A:LEU:H	1:33:A:LEU:HG	14	0.84
(1,922)	1:66:A:ILE:H	1:6:A:GLN:HB3	21	0.84
(1,918)	1:13:A:TYR:H	1:11:A:LYS:HD3	28	0.84
(1,912)	1:34:A:GLU:HB3	1:35:A:GLN:H	21	0.84
(1,870)	1:29:A:ILE:HG22	1:45:A:ILE:HA	2	0.84
(1,849)	1:13:A:TYR:H	1:15:A:ASN:HD22	21	0.84
(1,840)	1:11:A:LYS:HD3	1:12:A:ASP:H	7	0.84
(1,823)	1:28:A:ASP:H	1:9:A:ALA:HB2	29	0.84
(1,800)	1:44:A:CYS:HB3	1:55:A:VAL:HA	4	0.84
(1,797)	1:7:A:VAL:HB	1:63:A:GLY:HA2	24	0.84
(1,734)	1:39:A:GLY:H	1:40:A:ARG:HB2	7	0.84
(1,541)	1:40:A:ARG:H	1:40:A:ARG:HD2	11	0.84
(1,528)	1:31:A:THR:H	1:31:A:THR:HG22	3	0.84
(1,528)	1:31:A:THR:H	1:31:A:THR:HG23	15	0.84
(1,514)	1:65:A:ALA:H	1:64:A:GLU:HB2	8	0.84
(1,493)	1:41:A:TRP:HE1	1:40:A:ARG:H	27	0.84
(1,485)	1:46:A:HIS:H	1:45:A:ILE:HD11	2	0.84

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,485)	1:46:A:HIS:H	1:45:A:ILE:HD11	8	0.84
(1,485)	1:46:A:HIS:H	1:45:A:ILE:HD12	18	0.84
(1,485)	1:46:A:HIS:H	1:45:A:ILE:HD12	20	0.84
(1,464)	1:67:A:VAL:H	1:66:A:ILE:HG12	7	0.84
(1,444)	1:30:A:ILE:HD13	1:32:A:VAL:HG13	26	0.84
(1,441)	1:30:A:ILE:HG21	1:45:A:ILE:HG22	12	0.84
(1,441)	1:30:A:ILE:HG22	1:45:A:ILE:HG22	22	0.84
(1,440)	1:30:A:ILE:HG23	1:32:A:VAL:HG21	8	0.84
(1,440)	1:30:A:ILE:HG21	1:32:A:VAL:HG23	10	0.84
(1,440)	1:30:A:ILE:HG22	1:32:A:VAL:HG22	11	0.84
(1,382)	1:65:A:ALA:H	1:5:A:LEU:HB3	24	0.84
(1,350)	1:4:A:MET:H	1:34:A:GLU:HB2	9	0.84
(1,300)	1:4:A:MET:H	1:3:A:HIS:HB3	14	0.84
(1,27)	1:43:A:GLY:HA3	1:33:A:LEU:HD11	21	0.84
(1,950)	1:9:A:ALA:H	1:24:A:VAL:HG13	24	0.83
(1,947)	1:7:A:VAL:HG21	1:9:A:ALA:H	21	0.83
(1,946)	1:7:A:VAL:HG21	1:30:A:ILE:H	25	0.83
(1,945)	1:41:A:TRP:HH2	1:5:A:LEU:HD13	27	0.83
(1,941)	1:33:A:LEU:H	1:33:A:LEU:HG	15	0.83
(1,939)	1:58:A:PHE:HD1	1:32:A:VAL:HG12	12	0.83
(1,935)	1:40:A:ARG:HB3	1:38:A:ASP:HB3	14	0.83
(1,934)	1:26:A:ALA:H	1:25:A:LYS:HG2	3	0.83
(1,908)	1:52:A:ASN:H	1:49:A:ARG:HB3	10	0.83
(1,902)	1:38:A:ASP:H	1:38:A:ASP:HB3	2	0.83
(1,883)	1:13:A:TYR:HE1	1:21:A:SER:HB2	25	0.83
(1,873)	1:41:A:TRP:HB3	1:34:A:GLU:HA	28	0.83
(1,863)	1:9:A:ALA:H	1:29:A:ILE:HA	15	0.83
(1,852)	1:13:A:TYR:HE1	1:24:A:VAL:H	22	0.83
(1,840)	1:11:A:LYS:HD3	1:12:A:ASP:H	16	0.83
(1,834)	1:39:A:GLY:HA3	1:41:A:TRP:H	29	0.83
(1,813)	1:22:A:LEU:HB2	1:13:A:TYR:H	2	0.83
(1,813)	1:22:A:LEU:HB2	1:13:A:TYR:H	13	0.83
(1,804)	1:22:A:LEU:HB3	1:22:A:LEU:H	19	0.83
(1,797)	1:7:A:VAL:HB	1:63:A:GLY:HA2	20	0.83
(1,797)	1:7:A:VAL:HB	1:63:A:GLY:HA2	27	0.83
(1,687)	1:27:A:GLY:H	1:26:A:ALA:HB2	2	0.83
(1,685)	1:27:A:GLY:H	1:24:A:VAL:HG22	5	0.83
(1,685)	1:27:A:GLY:H	1:24:A:VAL:HG23	24	0.83
(1,543)	1:40:A:ARG:H	1:40:A:ARG:HG2	28	0.83
(1,536)	1:23:A:ASN:H	1:22:A:LEU:HD11	11	0.83
(1,528)	1:31:A:THR:H	1:31:A:THR:HG23	8	0.83
(1,485)	1:46:A:HIS:H	1:45:A:ILE:HD13	19	0.83

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,444)	1:30:A:ILE:HD12	1:32:A:VAL:HG13	12	0.83
(1,441)	1:30:A:ILE:HG23	1:45:A:ILE:HG21	3	0.83
(1,441)	1:30:A:ILE:HG23	1:45:A:ILE:HG22	28	0.83
(1,440)	1:30:A:ILE:HG22	1:32:A:VAL:HG23	14	0.83
(1,440)	1:30:A:ILE:HG22	1:32:A:VAL:HG22	15	0.83
(1,437)	1:24:A:VAL:HG22	1:30:A:ILE:HG21	15	0.83
(1,431)	1:22:A:LEU:HD11	1:24:A:VAL:HG13	19	0.83
(1,416)	1:5:A:LEU:HD23	1:7:A:VAL:HG11	25	0.83
(1,403)	1:58:A:PHE:H	1:22:A:LEU:HD12	14	0.83
(1,317)	1:39:A:GLY:H	1:38:A:ASP:HB3	24	0.83
(1,220)	1:15:A:ASN:H	1:15:A:ASN:HD22	5	0.83
(1,121)	1:26:A:ALA:H	1:26:A:ALA:HB3	4	0.83
(1,967)	1:62:A:LEU:H	1:63:A:GLY:HA2	7	0.82
(1,939)	1:58:A:PHE:HD1	1:32:A:VAL:HG12	10	0.82
(1,939)	1:58:A:PHE:HD1	1:32:A:VAL:HG11	13	0.82
(1,939)	1:58:A:PHE:HD1	1:32:A:VAL:HG12	26	0.82
(1,928)	1:60:A:SER:HB3	1:41:A:TRP:HH2	15	0.82
(1,899)	1:27:A:GLY:H	1:28:A:ASP:HB2	17	0.82
(1,892)	1:13:A:TYR:HD2	1:56:A:GLY:HA3	7	0.82
(1,878)	1:52:A:ASN:H	1:50:A:THR:HB	18	0.82
(1,863)	1:9:A:ALA:H	1:29:A:ILE:HA	28	0.82
(1,819)	1:58:A:PHE:HD1	1:42:A:LYS:H	18	0.82
(1,813)	1:22:A:LEU:HB2	1:13:A:TYR:H	27	0.82
(1,800)	1:44:A:CYS:HB3	1:55:A:VAL:HA	23	0.82
(1,797)	1:7:A:VAL:HB	1:63:A:GLY:HA2	21	0.82
(1,737)	1:50:A:THR:H	1:50:A:THR:HG21	24	0.82
(1,648)	1:12:A:ASP:H	1:12:A:ASP:HB3	2	0.82
(1,628)	1:38:A:ASP:H	1:36:A:HIS:HB2	13	0.82
(1,529)	1:31:A:THR:H	1:30:A:ILE:HG13	2	0.82
(1,528)	1:31:A:THR:H	1:31:A:THR:HG22	27	0.82
(1,514)	1:65:A:ALA:H	1:64:A:GLU:HB2	23	0.82
(1,507)	1:22:A:LEU:H	1:22:A:LEU:HD13	2	0.82
(1,467)	1:30:A:ILE:H	1:29:A:ILE:HG21	17	0.82
(1,440)	1:30:A:ILE:HG23	1:32:A:VAL:HG21	27	0.82
(1,382)	1:65:A:ALA:H	1:5:A:LEU:HB3	10	0.82
(1,324)	1:4:A:MET:H	1:4:A:MET:HG3	15	0.82
(1,211)	1:25:A:LYS:H	1:11:A:LYS:H	21	0.82
(1,963)	1:30:A:ILE:HD13	1:22:A:LEU:H	3	0.81
(1,956)	1:24:A:VAL:HG22	1:13:A:TYR:HD2	13	0.81
(1,947)	1:7:A:VAL:HG23	1:9:A:ALA:H	10	0.81
(1,919)	1:23:A:ASN:HD21	1:25:A:LYS:HB3	20	0.81
(1,912)	1:41:A:TRP:H	1:40:A:ARG:HG3	9	0.81

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,892)	1:13:A:TYR:HD2	1:56:A:GLY:HA3	21	0.81
(1,873)	1:41:A:TRP:HB3	1:34:A:GLU:HA	27	0.81
(1,863)	1:9:A:ALA:H	1:29:A:ILE:HA	17	0.81
(1,823)	1:28:A:ASP:H	1:9:A:ALA:HB3	19	0.81
(1,797)	1:7:A:VAL:HB	1:63:A:GLY:HA2	11	0.81
(1,737)	1:50:A:THR:H	1:50:A:THR:HG21	25	0.81
(1,687)	1:27:A:GLY:H	1:26:A:ALA:HB1	9	0.81
(1,571)	1:66:A:ILE:H	1:65:A:ALA:HB2	15	0.81
(1,536)	1:23:A:ASN:H	1:22:A:LEU:HD11	13	0.81
(1,536)	1:23:A:ASN:H	1:22:A:LEU:HD12	23	0.81
(1,529)	1:31:A:THR:H	1:30:A:ILE:HG13	10	0.81
(1,514)	1:65:A:ALA:H	1:64:A:GLU:HB2	14	0.81
(1,485)	1:46:A:HIS:H	1:45:A:ILE:HD13	23	0.81
(1,444)	1:30:A:ILE:HD12	1:32:A:VAL:HG11	17	0.81
(1,441)	1:30:A:ILE:HG21	1:45:A:ILE:HG23	14	0.81
(1,440)	1:30:A:ILE:HG22	1:32:A:VAL:HG21	23	0.81
(1,330)	1:63:A:GLY:H	1:62:A:LEU:HB2	7	0.81
(1,46)	1:13:A:TYR:HB2	1:11:A:LYS:HD2	23	0.81
(1,966)	1:8:A:ARG:H	1:64:A:GLU:HG3	5	0.8
(1,941)	1:33:A:LEU:H	1:33:A:LEU:HG	10	0.8
(1,941)	1:33:A:LEU:H	1:33:A:LEU:HG	16	0.8
(1,939)	1:58:A:PHE:HD1	1:32:A:VAL:HG12	3	0.8
(1,909)	1:41:A:TRP:HD1	1:60:A:SER:HB2	29	0.8
(1,899)	1:27:A:GLY:H	1:28:A:ASP:HB2	21	0.8
(1,892)	1:13:A:TYR:HD2	1:56:A:GLY:HA3	12	0.8
(1,887)	1:29:A:ILE:HG21	1:31:A:THR:HB	20	0.8
(1,873)	1:41:A:TRP:HB3	1:34:A:GLU:HA	15	0.8
(1,852)	1:13:A:TYR:HE2	1:24:A:VAL:H	7	0.8
(1,847)	1:8:A:ARG:HD3	1:6:A:GLN:HE21	23	0.8
(1,840)	1:11:A:LYS:HD3	1:12:A:ASP:H	25	0.8
(1,834)	1:51:A:GLY:HA3	1:47:A:ASP:H	24	0.8
(1,797)	1:7:A:VAL:HB	1:63:A:GLY:HA2	28	0.8
(1,797)	1:7:A:VAL:HB	1:63:A:GLY:HA2	30	0.8
(1,734)	1:39:A:GLY:H	1:40:A:ARG:HB2	11	0.8
(1,734)	1:39:A:GLY:H	1:40:A:ARG:HB2	12	0.8
(1,687)	1:27:A:GLY:H	1:26:A:ALA:HB1	7	0.8
(1,648)	1:12:A:ASP:H	1:12:A:ASP:HB3	1	0.8
(1,648)	1:12:A:ASP:H	1:12:A:ASP:HB3	13	0.8
(1,648)	1:12:A:ASP:H	1:12:A:ASP:HB3	19	0.8
(1,648)	1:12:A:ASP:H	1:12:A:ASP:HB3	27	0.8
(1,544)	1:40:A:ARG:H	1:40:A:ARG:HB2	3	0.8
(1,536)	1:23:A:ASN:H	1:22:A:LEU:HD13	29	0.8

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,493)	1:41:A:TRP:HE1	1:40:A:ARG:H	22	0.8
(1,441)	1:30:A:ILE:HG22	1:45:A:ILE:HG21	24	0.8
(1,437)	1:24:A:VAL:HG22	1:30:A:ILE:HG22	29	0.8
(1,391)	1:34:A:GLU:H	1:5:A:LEU:HD22	19	0.8
(1,382)	1:65:A:ALA:H	1:5:A:LEU:HB3	19	0.8
(1,174)	1:30:A:ILE:HG12	1:9:A:ALA:H	28	0.8
(1,966)	1:8:A:ARG:H	1:64:A:GLU:HG3	29	0.79
(1,963)	1:30:A:ILE:HD11	1:22:A:LEU:H	14	0.79
(1,941)	1:33:A:LEU:H	1:33:A:LEU:HG	22	0.79
(1,939)	1:58:A:PHE:HD1	1:32:A:VAL:HG12	6	0.79
(1,939)	1:58:A:PHE:HD1	1:32:A:VAL:HG11	29	0.79
(1,939)	1:58:A:PHE:HD1	1:32:A:VAL:HG12	30	0.79
(1,912)	1:34:A:GLU:HB3	1:35:A:GLN:H	3	0.79
(1,887)	1:29:A:ILE:HG21	1:31:A:THR:HB	9	0.79
(1,887)	1:29:A:ILE:HG23	1:31:A:THR:HB	28	0.79
(1,852)	1:25:A:LYS:H	1:48:A:ASN:HD22	21	0.79
(1,840)	1:11:A:LYS:HD3	1:12:A:ASP:H	15	0.79
(1,816)	1:5:A:LEU:HB2	1:7:A:VAL:H	28	0.79
(1,813)	1:22:A:LEU:HB2	1:13:A:TYR:H	7	0.79
(1,804)	1:22:A:LEU:HB3	1:22:A:LEU:H	30	0.79
(1,802)	1:26:A:ALA:HB2	1:12:A:ASP:HA	6	0.79
(1,800)	1:44:A:CYS:HB3	1:55:A:VAL:HA	6	0.79
(1,797)	1:7:A:VAL:HB	1:63:A:GLY:HA2	17	0.79
(1,737)	1:50:A:THR:H	1:50:A:THR:HG22	27	0.79
(1,687)	1:27:A:GLY:H	1:26:A:ALA:HB1	28	0.79
(1,648)	1:12:A:ASP:H	1:12:A:ASP:HB3	5	0.79
(1,648)	1:12:A:ASP:H	1:12:A:ASP:HB3	9	0.79
(1,648)	1:12:A:ASP:H	1:12:A:ASP:HB3	14	0.79
(1,648)	1:12:A:ASP:H	1:12:A:ASP:HB3	20	0.79
(1,648)	1:12:A:ASP:H	1:12:A:ASP:HB3	24	0.79
(1,571)	1:66:A:ILE:H	1:65:A:ALA:HB2	4	0.79
(1,544)	1:40:A:ARG:H	1:40:A:ARG:HB2	12	0.79
(1,541)	1:40:A:ARG:H	1:40:A:ARG:HD2	12	0.79
(1,514)	1:65:A:ALA:H	1:64:A:GLU:HB2	1	0.79
(1,507)	1:22:A:LEU:H	1:22:A:LEU:HD11	11	0.79
(1,493)	1:41:A:TRP:HE1	1:40:A:ARG:H	25	0.79
(1,451)	1:25:A:LYS:H	1:24:A:VAL:HB	26	0.79
(1,441)	1:30:A:ILE:HG23	1:45:A:ILE:HG22	7	0.79
(1,441)	1:30:A:ILE:HG21	1:45:A:ILE:HG23	26	0.79
(1,440)	1:30:A:ILE:HG22	1:32:A:VAL:HG22	13	0.79
(1,437)	1:24:A:VAL:HG21	1:30:A:ILE:HG23	3	0.79
(1,437)	1:24:A:VAL:HG21	1:30:A:ILE:HG21	20	0.79

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,416)	1:5:A:LEU:HD23	1:7:A:VAL:HG12	6	0.79
(1,357)	1:65:A:ALA:HB2	1:5:A:LEU:HB3	2	0.79
(1,220)	1:15:A:ASN:H	1:15:A:ASN:HD22	13	0.79
(1,121)	1:26:A:ALA:H	1:26:A:ALA:HB1	19	0.79
(1,57)	1:28:A:ASP:HB2	1:25:A:LYS:HG2	4	0.79
(1,6)	1:55:A:VAL:HA	1:55:A:VAL:HG12	3	0.79
(1,6)	1:55:A:VAL:HA	1:55:A:VAL:HG12	6	0.79
(1,6)	1:55:A:VAL:HA	1:55:A:VAL:HG13	7	0.79
(1,6)	1:55:A:VAL:HA	1:55:A:VAL:HG13	13	0.79
(1,6)	1:55:A:VAL:HA	1:55:A:VAL:HG11	14	0.79
(1,6)	1:55:A:VAL:HA	1:55:A:VAL:HG13	15	0.79
(1,6)	1:55:A:VAL:HA	1:55:A:VAL:HG11	17	0.79
(1,6)	1:55:A:VAL:HA	1:55:A:VAL:HG13	18	0.79
(1,6)	1:55:A:VAL:HA	1:55:A:VAL:HG13	20	0.79
(1,6)	1:55:A:VAL:HA	1:55:A:VAL:HG11	25	0.79
(1,6)	1:55:A:VAL:HA	1:55:A:VAL:HG11	26	0.79
(1,968)	1:10:A:THR:H	1:63:A:GLY:HA3	30	0.78
(1,966)	1:8:A:ARG:H	1:64:A:GLU:HG3	11	0.78
(1,966)	1:8:A:ARG:H	1:64:A:GLU:HG3	28	0.78
(1,959)	1:31:A:THR:H	1:22:A:LEU:HD11	7	0.78
(1,948)	1:6:A:GLN:H	1:67:A:VAL:HG22	6	0.78
(1,946)	1:7:A:VAL:HG22	1:30:A:ILE:H	20	0.78
(1,946)	1:7:A:VAL:HG23	1:30:A:ILE:H	26	0.78
(1,939)	1:58:A:PHE:HD1	1:32:A:VAL:HG13	27	0.78
(1,934)	1:26:A:ALA:H	1:25:A:LYS:HG3	16	0.78
(1,909)	1:41:A:TRP:HD1	1:40:A:ARG:HG2	17	0.78
(1,908)	1:52:A:ASN:H	1:49:A:ARG:HB2	24	0.78
(1,899)	1:15:A:ASN:H	1:14:A:CYS:HB3	24	0.78
(1,813)	1:22:A:LEU:HB2	1:13:A:TYR:H	21	0.78
(1,810)	1:23:A:ASN:H	1:45:A:ILE:H	14	0.78
(1,800)	1:44:A:CYS:HB3	1:55:A:VAL:HA	17	0.78
(1,800)	1:44:A:CYS:HB3	1:55:A:VAL:HA	20	0.78
(1,798)	1:24:A:VAL:HB	1:12:A:ASP:HA	24	0.78
(1,798)	1:24:A:VAL:HB	1:13:A:TYR:HA	30	0.78
(1,795)	1:17:A:TYR:HB3	1:18:A:ASP:H	12	0.78
(1,654)	1:17:A:TYR:H	1:17:A:TYR:HB3	8	0.78
(1,648)	1:12:A:ASP:H	1:12:A:ASP:HB3	21	0.78
(1,648)	1:12:A:ASP:H	1:12:A:ASP:HB3	30	0.78
(1,544)	1:40:A:ARG:H	1:40:A:ARG:HB2	20	0.78
(1,544)	1:40:A:ARG:H	1:40:A:ARG:HB2	24	0.78
(1,485)	1:46:A:HIS:H	1:45:A:ILE:HD12	16	0.78
(1,440)	1:30:A:ILE:HG22	1:32:A:VAL:HG22	4	0.78

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,440)	1:30:A:ILE:HG22	1:32:A:VAL:HG21	5	0.78
(1,440)	1:30:A:ILE:HG23	1:32:A:VAL:HG21	24	0.78
(1,434)	1:24:A:VAL:HG11	1:30:A:ILE:HD12	3	0.78
(1,420)	1:7:A:VAL:HG12	1:32:A:VAL:HG13	5	0.78
(1,382)	1:65:A:ALA:H	1:5:A:LEU:HB3	4	0.78
(1,229)	1:36:A:HIS:HB2	1:36:A:HIS:H	14	0.78
(1,121)	1:26:A:ALA:H	1:26:A:ALA:HB3	9	0.78
(1,6)	1:55:A:VAL:HA	1:55:A:VAL:HG12	1	0.78
(1,6)	1:55:A:VAL:HA	1:55:A:VAL:HG12	4	0.78
(1,6)	1:55:A:VAL:HA	1:55:A:VAL:HG11	5	0.78
(1,6)	1:55:A:VAL:HA	1:55:A:VAL:HG13	9	0.78
(1,6)	1:55:A:VAL:HA	1:55:A:VAL:HG11	10	0.78
(1,6)	1:55:A:VAL:HA	1:55:A:VAL:HG13	11	0.78
(1,6)	1:55:A:VAL:HA	1:55:A:VAL:HG11	12	0.78
(1,6)	1:55:A:VAL:HA	1:55:A:VAL:HG12	16	0.78
(1,6)	1:55:A:VAL:HA	1:55:A:VAL:HG11	21	0.78
(1,6)	1:55:A:VAL:HA	1:55:A:VAL:HG12	22	0.78
(1,6)	1:55:A:VAL:HA	1:55:A:VAL:HG11	23	0.78
(1,6)	1:55:A:VAL:HA	1:55:A:VAL:HG12	24	0.78
(1,6)	1:55:A:VAL:HA	1:55:A:VAL:HG11	27	0.78
(1,6)	1:55:A:VAL:HA	1:55:A:VAL:HG13	28	0.78
(1,6)	1:55:A:VAL:HA	1:55:A:VAL:HG11	30	0.78
(1,959)	1:31:A:THR:H	1:22:A:LEU:HD11	1	0.77
(1,956)	1:24:A:VAL:HG23	1:13:A:TYR:HD2	23	0.77
(1,941)	1:33:A:LEU:H	1:33:A:LEU:HG	11	0.77
(1,941)	1:33:A:LEU:H	1:33:A:LEU:HG	28	0.77
(1,934)	1:26:A:ALA:H	1:25:A:LYS:HG3	27	0.77
(1,931)	1:41:A:TRP:HZ2	1:60:A:SER:HB3	19	0.77
(1,922)	1:66:A:ILE:H	1:6:A:GLN:HB3	18	0.77
(1,873)	1:41:A:TRP:HB3	1:34:A:GLU:HA	14	0.77
(1,850)	1:11:A:LYS:HG3	1:13:A:TYR:HE1	12	0.77
(1,840)	1:11:A:LYS:HD3	1:12:A:ASP:H	14	0.77
(1,823)	1:28:A:ASP:H	1:9:A:ALA:HB2	18	0.77
(1,813)	1:22:A:LEU:HB2	1:13:A:TYR:H	18	0.77
(1,802)	1:26:A:ALA:HB1	1:12:A:ASP:HA	21	0.77
(1,802)	1:26:A:ALA:HB2	1:12:A:ASP:HA	27	0.77
(1,800)	1:44:A:CYS:HB3	1:55:A:VAL:HA	21	0.77
(1,687)	1:27:A:GLY:H	1:26:A:ALA:HB3	3	0.77
(1,670)	1:43:A:GLY:H	1:42:A:LYS:HB2	7	0.77
(1,654)	1:17:A:TYR:H	1:17:A:TYR:HB3	17	0.77
(1,654)	1:17:A:TYR:H	1:17:A:TYR:HB3	19	0.77
(1,536)	1:23:A:ASN:H	1:22:A:LEU:HD13	2	0.77

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,536)	1:23:A:ASN:H	1:22:A:LEU:HD12	5	0.77
(1,514)	1:65:A:ALA:H	1:64:A:GLU:HB2	6	0.77
(1,514)	1:65:A:ALA:H	1:64:A:GLU:HB2	17	0.77
(1,514)	1:65:A:ALA:H	1:64:A:GLU:HB2	20	0.77
(1,507)	1:22:A:LEU:H	1:22:A:LEU:HD12	5	0.77
(1,451)	1:25:A:LYS:H	1:24:A:VAL:HB	15	0.77
(1,451)	1:25:A:LYS:H	1:24:A:VAL:HB	30	0.77
(1,441)	1:30:A:ILE:HG21	1:45:A:ILE:HG23	25	0.77
(1,431)	1:22:A:LEU:HD11	1:24:A:VAL:HG12	3	0.77
(1,318)	1:44:A:CYS:HB3	1:45:A:ILE:H	30	0.77
(1,317)	1:39:A:GLY:H	1:38:A:ASP:HB3	26	0.77
(1,305)	1:13:A:TYR:HD1	1:11:A:LYS:HE3	13	0.77
(1,228)	1:51:A:GLY:H	1:52:A:ASN:HD22	15	0.77
(1,220)	1:15:A:ASN:H	1:15:A:ASN:HD22	3	0.77
(1,121)	1:26:A:ALA:H	1:26:A:ALA:HB1	30	0.77
(1,71)	1:38:A:ASP:HB2	1:38:A:ASP:HA	2	0.77
(1,71)	1:38:A:ASP:HB2	1:38:A:ASP:HA	4	0.77
(1,71)	1:38:A:ASP:HB2	1:38:A:ASP:HA	5	0.77
(1,71)	1:38:A:ASP:HB2	1:38:A:ASP:HA	6	0.77
(1,71)	1:38:A:ASP:HB2	1:38:A:ASP:HA	8	0.77
(1,71)	1:38:A:ASP:HB2	1:38:A:ASP:HA	10	0.77
(1,71)	1:38:A:ASP:HB2	1:38:A:ASP:HA	13	0.77
(1,71)	1:38:A:ASP:HB2	1:38:A:ASP:HA	26	0.77
(1,39)	1:54:A:ARG:H	1:53:A:ASP:HB2	6	0.77
(1,39)	1:54:A:ARG:H	1:53:A:ASP:HB2	27	0.77
(1,6)	1:55:A:VAL:HA	1:55:A:VAL:HG12	2	0.77
(1,6)	1:55:A:VAL:HA	1:55:A:VAL:HG12	8	0.77
(1,6)	1:55:A:VAL:HA	1:55:A:VAL:HG11	19	0.77
(1,968)	1:10:A:THR:H	1:63:A:GLY:HA3	16	0.76
(1,959)	1:31:A:THR:H	1:22:A:LEU:HD12	12	0.76
(1,959)	1:31:A:THR:H	1:22:A:LEU:HD11	14	0.76
(1,956)	1:24:A:VAL:HG23	1:13:A:TYR:HD2	11	0.76
(1,941)	1:33:A:LEU:H	1:33:A:LEU:HG	23	0.76
(1,941)	1:33:A:LEU:H	1:33:A:LEU:HG	26	0.76
(1,940)	1:7:A:VAL:H	1:32:A:VAL:HG12	24	0.76
(1,939)	1:58:A:PHE:HD1	1:32:A:VAL:HG13	16	0.76
(1,925)	1:11:A:LYS:HG2	1:13:A:TYR:H	20	0.76
(1,912)	1:41:A:TRP:H	1:40:A:ARG:HG3	27	0.76
(1,852)	1:13:A:TYR:HE2	1:24:A:VAL:H	12	0.76
(1,848)	1:16:A:ASN:HB2	1:18:A:ASP:H	10	0.76
(1,795)	1:17:A:TYR:HB3	1:18:A:ASP:H	20	0.76
(1,687)	1:27:A:GLY:H	1:26:A:ALA:HB3	1	0.76

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,687)	1:27:A:GLY:H	1:26:A:ALA:HB2	20	0.76
(1,670)	1:43:A:GLY:H	1:42:A:LYS:HB2	18	0.76
(1,654)	1:17:A:TYR:H	1:17:A:TYR:HB3	30	0.76
(1,651)	1:12:A:ASP:H	1:11:A:LYS:HG3	10	0.76
(1,648)	1:12:A:ASP:H	1:12:A:ASP:HB3	28	0.76
(1,514)	1:65:A:ALA:H	1:64:A:GLU:HB2	18	0.76
(1,514)	1:65:A:ALA:H	1:64:A:GLU:HB2	27	0.76
(1,434)	1:24:A:VAL:HG12	1:30:A:ILE:HD13	18	0.76
(1,431)	1:22:A:LEU:HD13	1:24:A:VAL:HG11	20	0.76
(1,220)	1:15:A:ASN:H	1:15:A:ASN:HD22	9	0.76
(1,174)	1:30:A:ILE:HG12	1:9:A:ALA:H	22	0.76
(1,121)	1:26:A:ALA:H	1:26:A:ALA:HB2	14	0.76
(1,71)	1:38:A:ASP:HB2	1:38:A:ASP:HA	11	0.76
(1,71)	1:38:A:ASP:HB2	1:38:A:ASP:HA	24	0.76
(1,46)	1:13:A:TYR:HB2	1:11:A:LYS:HD2	6	0.76
(1,39)	1:54:A:ARG:H	1:53:A:ASP:HB2	24	0.76
(1,6)	1:55:A:VAL:HA	1:55:A:VAL:HG12	29	0.76
(1,956)	1:5:A:LEU:HD23	1:41:A:TRP:HH2	20	0.75
(1,941)	1:33:A:LEU:H	1:33:A:LEU:HG	24	0.75
(1,939)	1:58:A:PHE:HD1	1:32:A:VAL:HG11	1	0.75
(1,934)	1:26:A:ALA:H	1:25:A:LYS:HG3	17	0.75
(1,925)	1:11:A:LYS:HG2	1:13:A:TYR:H	27	0.75
(1,921)	1:22:A:LEU:H	1:22:A:LEU:HB2	17	0.75
(1,887)	1:29:A:ILE:HG22	1:31:A:THR:HB	27	0.75
(1,849)	1:13:A:TYR:H	1:15:A:ASN:HD22	13	0.75
(1,840)	1:11:A:LYS:HD3	1:12:A:ASP:H	30	0.75
(1,822)	1:36:A:HIS:HB3	1:38:A:ASP:H	1	0.75
(1,810)	1:23:A:ASN:H	1:45:A:ILE:H	8	0.75
(1,804)	1:22:A:LEU:HB3	1:24:A:VAL:H	8	0.75
(1,800)	1:44:A:CYS:HB3	1:55:A:VAL:HA	25	0.75
(1,797)	1:7:A:VAL:HB	1:63:A:GLY:HA2	29	0.75
(1,670)	1:43:A:GLY:H	1:42:A:LYS:HB2	5	0.75
(1,635)	1:52:A:ASN:H	1:52:A:ASN:HB2	15	0.75
(1,544)	1:40:A:ARG:H	1:40:A:ARG:HB2	30	0.75
(1,514)	1:65:A:ALA:H	1:64:A:GLU:HB2	22	0.75
(1,467)	1:30:A:ILE:H	1:29:A:ILE:HG23	12	0.75
(1,467)	1:30:A:ILE:H	1:29:A:ILE:HG21	24	0.75
(1,467)	1:30:A:ILE:H	1:29:A:ILE:HG22	25	0.75
(1,451)	1:25:A:LYS:H	1:24:A:VAL:HB	16	0.75
(1,451)	1:25:A:LYS:H	1:24:A:VAL:HB	27	0.75
(1,444)	1:30:A:ILE:HD12	1:32:A:VAL:HG11	22	0.75
(1,441)	1:30:A:ILE:HG23	1:45:A:ILE:HG23	16	0.75

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,440)	1:30:A:ILE:HG23	1:32:A:VAL:HG22	20	0.75
(1,420)	1:7:A:VAL:HG12	1:32:A:VAL:HG12	24	0.75
(1,416)	1:5:A:LEU:HD22	1:7:A:VAL:HG13	4	0.75
(1,190)	1:34:A:GLU:HG2	1:3:A:HIS:H	14	0.75
(1,174)	1:30:A:ILE:HG12	1:9:A:ALA:H	7	0.75
(1,174)	1:30:A:ILE:HG12	1:9:A:ALA:H	19	0.75
(1,121)	1:26:A:ALA:H	1:26:A:ALA:HB1	8	0.75
(1,121)	1:26:A:ALA:H	1:26:A:ALA:HB3	10	0.75
(1,121)	1:26:A:ALA:H	1:26:A:ALA:HB2	11	0.75
(1,57)	1:28:A:ASP:HB2	1:25:A:LYS:HG2	30	0.75
(1,963)	1:30:A:ILE:HD11	1:22:A:LEU:H	7	0.74
(1,950)	1:9:A:ALA:H	1:24:A:VAL:HG12	26	0.74
(1,945)	1:41:A:TRP:HH2	1:5:A:LEU:HD12	1	0.74
(1,945)	1:41:A:TRP:HH2	1:5:A:LEU:HD13	24	0.74
(1,941)	1:33:A:LEU:H	1:33:A:LEU:HG	17	0.74
(1,939)	1:58:A:PHE:HD1	1:32:A:VAL:HG13	4	0.74
(1,939)	1:58:A:PHE:HD1	1:32:A:VAL:HG11	25	0.74
(1,921)	1:22:A:LEU:H	1:22:A:LEU:HB2	19	0.74
(1,894)	1:23:A:ASN:H	1:54:A:ARG:HD2	5	0.74
(1,834)	1:39:A:GLY:HA3	1:41:A:TRP:H	18	0.74
(1,797)	1:7:A:VAL:HB	1:63:A:GLY:HA2	25	0.74
(1,797)	1:7:A:VAL:HB	1:63:A:GLY:HA2	26	0.74
(1,733)	1:39:A:GLY:H	1:36:A:HIS:HB3	1	0.74
(1,687)	1:27:A:GLY:H	1:26:A:ALA:HB1	22	0.74
(1,650)	1:12:A:ASP:H	1:11:A:LYS:HB2	15	0.74
(1,650)	1:12:A:ASP:H	1:11:A:LYS:HB2	21	0.74
(1,571)	1:66:A:ILE:H	1:65:A:ALA:HB1	8	0.74
(1,571)	1:66:A:ILE:H	1:65:A:ALA:HB3	24	0.74
(1,541)	1:40:A:ARG:H	1:40:A:ARG:HD2	24	0.74
(1,485)	1:46:A:HIS:H	1:45:A:ILE:HD13	17	0.74
(1,485)	1:46:A:HIS:H	1:45:A:ILE:HD12	28	0.74
(1,444)	1:30:A:ILE:HD12	1:32:A:VAL:HG12	24	0.74
(1,440)	1:30:A:ILE:HG22	1:32:A:VAL:HG21	30	0.74
(1,437)	1:24:A:VAL:HG21	1:30:A:ILE:HG23	16	0.74
(1,437)	1:24:A:VAL:HG23	1:30:A:ILE:HG21	27	0.74
(1,431)	1:22:A:LEU:HD13	1:24:A:VAL:HG13	8	0.74
(1,431)	1:22:A:LEU:HD11	1:24:A:VAL:HG12	18	0.74
(1,420)	1:7:A:VAL:HG11	1:32:A:VAL:HG13	25	0.74
(1,394)	1:41:A:TRP:HZ3	1:5:A:LEU:HD21	8	0.74
(1,391)	1:34:A:GLU:H	1:5:A:LEU:HD22	30	0.74
(1,174)	1:30:A:ILE:HG12	1:9:A:ALA:H	20	0.74
(1,121)	1:26:A:ALA:H	1:26:A:ALA:HB2	6	0.74

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,121)	1:26:A:ALA:H	1:26:A:ALA:HB1	16	0.74
(1,121)	1:26:A:ALA:H	1:26:A:ALA:HB1	20	0.74
(1,46)	1:13:A:TYR:HB2	1:11:A:LYS:HD2	17	0.74
(1,946)	1:7:A:VAL:HG23	1:30:A:ILE:H	10	0.73
(1,946)	1:7:A:VAL:HG22	1:30:A:ILE:H	16	0.73
(1,941)	1:33:A:LEU:H	1:33:A:LEU:HG	12	0.73
(1,941)	1:33:A:LEU:H	1:33:A:LEU:HG	25	0.73
(1,934)	1:26:A:ALA:H	1:25:A:LYS:HG3	13	0.73
(1,918)	1:13:A:TYR:H	1:11:A:LYS:HD2	24	0.73
(1,887)	1:29:A:ILE:HG21	1:31:A:THR:HB	3	0.73
(1,887)	1:29:A:ILE:HG22	1:31:A:THR:HB	17	0.73
(1,870)	1:29:A:ILE:HG22	1:45:A:ILE:HA	14	0.73
(1,863)	1:9:A:ALA:H	1:29:A:ILE:HA	22	0.73
(1,850)	1:11:A:LYS:HG3	1:13:A:TYR:HE1	3	0.73
(1,840)	1:11:A:LYS:HD3	1:12:A:ASP:H	27	0.73
(1,810)	1:23:A:ASN:H	1:45:A:ILE:H	2	0.73
(1,800)	1:44:A:CYS:HB3	1:55:A:VAL:HA	27	0.73
(1,795)	1:17:A:TYR:HB3	1:18:A:ASP:H	10	0.73
(1,752)	1:9:A:ALA:H	1:27:A:GLY:H	4	0.73
(1,687)	1:27:A:GLY:H	1:26:A:ALA:HB3	8	0.73
(1,687)	1:27:A:GLY:H	1:26:A:ALA:HB1	17	0.73
(1,620)	1:21:A:SER:H	1:14:A:CYS:HB2	25	0.73
(1,589)	1:42:A:LYS:H	1:42:A:LYS:HB3	30	0.73
(1,587)	1:53:A:ASP:H	1:53:A:ASP:HB2	20	0.73
(1,544)	1:40:A:ARG:H	1:40:A:ARG:HB2	2	0.73
(1,514)	1:65:A:ALA:H	1:64:A:GLU:HB2	3	0.73
(1,514)	1:65:A:ALA:H	1:64:A:GLU:HB2	4	0.73
(1,507)	1:22:A:LEU:H	1:22:A:LEU:HD12	10	0.73
(1,485)	1:46:A:HIS:H	1:45:A:ILE:HD13	26	0.73
(1,451)	1:25:A:LYS:H	1:24:A:VAL:HB	21	0.73
(1,444)	1:30:A:ILE:HD11	1:32:A:VAL:HG13	16	0.73
(1,403)	1:58:A:PHE:H	1:22:A:LEU:HD12	19	0.73
(1,210)	1:41:A:TRP:HZ3	1:61:A:SER:H	7	0.73
(1,190)	1:34:A:GLU:HG2	1:3:A:HIS:H	8	0.73
(1,178)	1:58:A:PHE:HE1	1:33:A:LEU:H	18	0.73
(1,121)	1:26:A:ALA:H	1:26:A:ALA:HB3	12	0.73
(1,57)	1:28:A:ASP:HB2	1:25:A:LYS:HG2	2	0.73
(1,46)	1:13:A:TYR:HB2	1:11:A:LYS:HD2	10	0.73
(1,27)	1:43:A:GLY:HA3	1:33:A:LEU:HD12	3	0.73
(1,27)	1:43:A:GLY:HA3	1:33:A:LEU:HD11	23	0.73
(1,946)	1:7:A:VAL:HG23	1:30:A:ILE:H	11	0.72
(1,946)	1:7:A:VAL:HG23	1:30:A:ILE:H	23	0.72

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,941)	1:33:A:LEU:H	1:33:A:LEU:HG	6	0.72
(1,941)	1:33:A:LEU:H	1:33:A:LEU:HG	19	0.72
(1,941)	1:33:A:LEU:H	1:33:A:LEU:HG	30	0.72
(1,939)	1:58:A:PHE:HD1	1:32:A:VAL:HG11	14	0.72
(1,863)	1:9:A:ALA:H	1:29:A:ILE:HA	6	0.72
(1,863)	1:9:A:ALA:H	1:29:A:ILE:HA	12	0.72
(1,852)	1:13:A:TYR:HE2	1:24:A:VAL:H	23	0.72
(1,848)	1:16:A:ASN:HB2	1:18:A:ASP:H	28	0.72
(1,840)	1:11:A:LYS:HD3	1:12:A:ASP:H	11	0.72
(1,840)	1:12:A:ASP:H	1:11:A:LYS:HD2	20	0.72
(1,793)	1:28:A:ASP:HB3	1:45:A:ILE:HG23	14	0.72
(1,650)	1:12:A:ASP:H	1:11:A:LYS:HB2	16	0.72
(1,650)	1:12:A:ASP:H	1:11:A:LYS:HB2	17	0.72
(1,650)	1:12:A:ASP:H	1:11:A:LYS:HB2	25	0.72
(1,589)	1:42:A:LYS:H	1:42:A:LYS:HB3	29	0.72
(1,571)	1:66:A:ILE:H	1:65:A:ALA:HB3	1	0.72
(1,571)	1:66:A:ILE:H	1:65:A:ALA:HB2	28	0.72
(1,544)	1:40:A:ARG:H	1:40:A:ARG:HB2	8	0.72
(1,544)	1:40:A:ARG:H	1:40:A:ARG:HB2	26	0.72
(1,444)	1:30:A:ILE:HD13	1:32:A:VAL:HG12	15	0.72
(1,431)	1:22:A:LEU:HD12	1:24:A:VAL:HG12	12	0.72
(1,431)	1:22:A:LEU:HD12	1:24:A:VAL:HG11	15	0.72
(1,420)	1:7:A:VAL:HG13	1:32:A:VAL:HG13	15	0.72
(1,369)	1:29:A:ILE:HG13	1:6:A:GLN:HE22	5	0.72
(1,318)	1:44:A:CYS:HB3	1:45:A:ILE:H	26	0.72
(1,300)	1:4:A:MET:H	1:3:A:HIS:HB3	13	0.72
(1,174)	1:30:A:ILE:HG12	1:9:A:ALA:H	29	0.72
(1,121)	1:26:A:ALA:H	1:26:A:ALA:HB2	1	0.72
(1,121)	1:26:A:ALA:H	1:26:A:ALA:HB2	15	0.72
(1,121)	1:26:A:ALA:H	1:26:A:ALA:HB3	18	0.72
(1,121)	1:26:A:ALA:H	1:26:A:ALA:HB2	27	0.72
(1,121)	1:26:A:ALA:H	1:26:A:ALA:HB2	28	0.72
(1,941)	1:33:A:LEU:H	1:33:A:LEU:HG	21	0.71
(1,939)	1:58:A:PHE:HD1	1:32:A:VAL:HG13	21	0.71
(1,921)	1:22:A:LEU:H	1:22:A:LEU:HB2	22	0.71
(1,870)	1:29:A:ILE:HG22	1:45:A:ILE:HA	27	0.71
(1,863)	1:9:A:ALA:H	1:29:A:ILE:HA	27	0.71
(1,813)	1:22:A:LEU:HB2	1:13:A:TYR:H	20	0.71
(1,798)	1:24:A:VAL:HB	1:12:A:ASP:HA	17	0.71
(1,743)	1:63:A:GLY:H	1:62:A:LEU:HG	16	0.71
(1,635)	1:52:A:ASN:H	1:52:A:ASN:HB2	29	0.71
(1,617)	1:21:A:SER:H	1:21:A:SER:HB3	18	0.71

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,617)	1:21:A:SER:H	1:21:A:SER:HB3	25	0.71
(1,589)	1:42:A:LYS:H	1:42:A:LYS:HB3	16	0.71
(1,544)	1:40:A:ARG:H	1:40:A:ARG:HB2	11	0.71
(1,541)	1:40:A:ARG:H	1:40:A:ARG:HD2	29	0.71
(1,507)	1:22:A:LEU:H	1:22:A:LEU:HD13	24	0.71
(1,507)	1:22:A:LEU:H	1:22:A:LEU:HD12	25	0.71
(1,485)	1:46:A:HIS:H	1:45:A:ILE:HD11	5	0.71
(1,451)	1:25:A:LYS:H	1:24:A:VAL:HB	18	0.71
(1,444)	1:30:A:ILE:HD13	1:32:A:VAL:HG12	6	0.71
(1,441)	1:30:A:ILE:HG23	1:45:A:ILE:HG22	9	0.71
(1,431)	1:22:A:LEU:HD11	1:24:A:VAL:HG13	6	0.71
(1,420)	1:7:A:VAL:HG13	1:32:A:VAL:HG13	1	0.71
(1,395)	1:42:A:LYS:H	1:33:A:LEU:HD13	18	0.71
(1,387)	1:63:A:GLY:H	1:7:A:VAL:HG23	30	0.71
(1,328)	1:24:A:VAL:HB	1:26:A:ALA:H	4	0.71
(1,220)	1:15:A:ASN:H	1:15:A:ASN:HD22	15	0.71
(1,182)	1:25:A:LYS:H	1:27:A:GLY:H	24	0.71
(1,174)	1:30:A:ILE:HG12	1:9:A:ALA:H	1	0.71
(1,121)	1:26:A:ALA:H	1:26:A:ALA:HB2	7	0.71
(1,57)	1:28:A:ASP:HB2	1:25:A:LYS:HG2	11	0.71
(1,968)	1:10:A:THR:H	1:63:A:GLY:HA3	28	0.7
(1,958)	1:61:A:SER:H	1:5:A:LEU:HD21	23	0.7
(1,945)	1:41:A:TRP:HH2	1:5:A:LEU:HD12	10	0.7
(1,945)	1:41:A:TRP:HH2	1:5:A:LEU:HD13	29	0.7
(1,939)	1:58:A:PHE:HD1	1:32:A:VAL:HG11	11	0.7
(1,928)	1:60:A:SER:HB3	1:41:A:TRP:HH2	25	0.7
(1,925)	1:11:A:LYS:HG2	1:13:A:TYR:H	21	0.7
(1,919)	1:23:A:ASN:HD21	1:54:A:ARG:HB2	13	0.7
(1,912)	1:41:A:TRP:H	1:40:A:ARG:HG3	26	0.7
(1,906)	1:34:A:GLU:HG2	1:3:A:HIS:H	6	0.7
(1,851)	1:13:A:TYR:HE2	1:57:A:TYR:H	11	0.7
(1,840)	1:11:A:LYS:HD3	1:12:A:ASP:H	3	0.7
(1,840)	1:11:A:LYS:HD3	1:12:A:ASP:H	6	0.7
(1,810)	1:23:A:ASN:H	1:45:A:ILE:H	22	0.7
(1,800)	1:44:A:CYS:HB2	1:55:A:VAL:HA	26	0.7
(1,797)	1:7:A:VAL:HB	1:63:A:GLY:HA2	14	0.7
(1,670)	1:43:A:GLY:H	1:42:A:LYS:HB2	26	0.7
(1,650)	1:12:A:ASP:H	1:11:A:LYS:HB2	5	0.7
(1,589)	1:42:A:LYS:H	1:42:A:LYS:HB3	2	0.7
(1,589)	1:42:A:LYS:H	1:42:A:LYS:HB3	19	0.7
(1,544)	1:40:A:ARG:H	1:40:A:ARG:HB2	25	0.7
(1,514)	1:65:A:ALA:H	1:64:A:GLU:HB2	7	0.7

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,514)	1:65:A:ALA:H	1:64:A:GLU:HB2	15	0.7
(1,514)	1:65:A:ALA:H	1:64:A:GLU:HB2	25	0.7
(1,514)	1:65:A:ALA:H	1:64:A:GLU:HB2	26	0.7
(1,485)	1:46:A:HIS:H	1:45:A:ILE:HD13	9	0.7
(1,485)	1:46:A:HIS:H	1:45:A:ILE:HD12	13	0.7
(1,480)	1:45:A:ILE:H	1:44:A:CYS:HB2	26	0.7
(1,451)	1:25:A:LYS:H	1:24:A:VAL:HB	5	0.7
(1,448)	1:25:A:LYS:H	1:25:A:LYS:HB2	2	0.7
(1,448)	1:25:A:LYS:H	1:25:A:LYS:HB2	9	0.7
(1,448)	1:25:A:LYS:H	1:25:A:LYS:HB2	11	0.7
(1,444)	1:30:A:ILE:HD11	1:32:A:VAL:HG12	29	0.7
(1,441)	1:30:A:ILE:HG23	1:45:A:ILE:HG22	29	0.7
(1,390)	1:24:A:VAL:HG22	1:13:A:TYR:HB3	19	0.7
(1,369)	1:29:A:ILE:HG13	1:6:A:GLN:HE22	16	0.7
(1,182)	1:25:A:LYS:H	1:27:A:GLY:H	5	0.7
(1,121)	1:26:A:ALA:H	1:26:A:ALA:HB1	21	0.7
(1,946)	1:32:A:VAL:H	1:7:A:VAL:HG23	24	0.69
(1,939)	1:58:A:PHE:HD1	1:32:A:VAL:HG11	15	0.69
(1,934)	1:26:A:ALA:H	1:25:A:LYS:HG3	5	0.69
(1,925)	1:11:A:LYS:HG2	1:13:A:TYR:H	30	0.69
(1,840)	1:11:A:LYS:HD3	1:12:A:ASP:H	8	0.69
(1,800)	1:44:A:CYS:HB3	1:55:A:VAL:HA	8	0.69
(1,795)	1:17:A:TYR:HB3	1:18:A:ASP:H	23	0.69
(1,687)	1:27:A:GLY:H	1:26:A:ALA:HB3	11	0.69
(1,687)	1:27:A:GLY:H	1:26:A:ALA:HB1	15	0.69
(1,687)	1:27:A:GLY:H	1:26:A:ALA:HB3	26	0.69
(1,650)	1:12:A:ASP:H	1:11:A:LYS:HB2	6	0.69
(1,650)	1:12:A:ASP:H	1:11:A:LYS:HB2	7	0.69
(1,650)	1:12:A:ASP:H	1:11:A:LYS:HB2	24	0.69
(1,648)	1:12:A:ASP:H	1:12:A:ASP:HB3	18	0.69
(1,620)	1:21:A:SER:H	1:14:A:CYS:HB2	17	0.69
(1,572)	1:66:A:ILE:H	1:66:A:ILE:HG12	15	0.69
(1,507)	1:22:A:LEU:H	1:22:A:LEU:HD13	29	0.69
(1,485)	1:46:A:HIS:H	1:45:A:ILE:HD11	21	0.69
(1,445)	1:30:A:ILE:HD13	1:45:A:ILE:HG22	30	0.69
(1,444)	1:30:A:ILE:HD11	1:32:A:VAL:HG12	30	0.69
(1,441)	1:30:A:ILE:HG22	1:45:A:ILE:HG22	6	0.69
(1,437)	1:24:A:VAL:HG21	1:30:A:ILE:HG22	1	0.69
(1,437)	1:24:A:VAL:HG22	1:30:A:ILE:HG23	4	0.69
(1,403)	1:58:A:PHE:H	1:22:A:LEU:HD11	25	0.69
(1,395)	1:42:A:LYS:H	1:33:A:LEU:HD11	21	0.69
(1,391)	1:34:A:GLU:H	1:5:A:LEU:HD21	25	0.69

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,382)	1:65:A:ALA:H	1:5:A:LEU:HB3	17	0.69
(1,317)	1:39:A:GLY:H	1:38:A:ASP:HB3	28	0.69
(1,113)	1:31:A:THR:HB	1:44:A:CYS:HB2	13	0.69
(1,66)	1:28:A:ASP:HB3	1:25:A:LYS:HG3	25	0.69
(1,27)	1:43:A:GLY:HA3	1:33:A:LEU:HD12	1	0.69
(1,961)	1:44:A:CYS:H	1:22:A:LEU:HD12	18	0.68
(1,959)	1:31:A:THR:H	1:22:A:LEU:HD11	24	0.68
(1,908)	1:34:A:GLU:H	1:35:A:GLN:HG2	2	0.68
(1,899)	1:27:A:GLY:H	1:28:A:ASP:HB2	8	0.68
(1,898)	1:32:A:VAL:HG23	1:60:A:SER:HA	15	0.68
(1,887)	1:29:A:ILE:HG23	1:31:A:THR:HB	22	0.68
(1,878)	1:52:A:ASN:H	1:50:A:THR:HA	22	0.68
(1,878)	1:52:A:ASN:H	1:50:A:THR:HA	30	0.68
(1,870)	1:29:A:ILE:HG21	1:45:A:ILE:HA	10	0.68
(1,840)	1:12:A:ASP:H	1:11:A:LYS:HD2	19	0.68
(1,687)	1:27:A:GLY:H	1:26:A:ALA:HB1	6	0.68
(1,687)	1:27:A:GLY:H	1:26:A:ALA:HB1	14	0.68
(1,617)	1:21:A:SER:H	1:21:A:SER:HB3	27	0.68
(1,589)	1:42:A:LYS:H	1:42:A:LYS:HB3	4	0.68
(1,589)	1:42:A:LYS:H	1:42:A:LYS:HB3	22	0.68
(1,589)	1:42:A:LYS:H	1:42:A:LYS:HB3	23	0.68
(1,589)	1:42:A:LYS:H	1:42:A:LYS:HB3	25	0.68
(1,582)	1:13:A:TYR:H	1:13:A:TYR:HB3	25	0.68
(1,547)	1:6:A:GLN:H	1:5:A:LEU:HB3	29	0.68
(1,544)	1:40:A:ARG:H	1:40:A:ARG:HB2	27	0.68
(1,519)	1:14:A:CYS:H	1:13:A:TYR:HD1	30	0.68
(1,467)	1:30:A:ILE:H	1:29:A:ILE:HG23	26	0.68
(1,466)	1:30:A:ILE:H	1:29:A:ILE:HB	18	0.68
(1,451)	1:25:A:LYS:H	1:24:A:VAL:HB	20	0.68
(1,444)	1:30:A:ILE:HD11	1:32:A:VAL:HG12	7	0.68
(1,441)	1:30:A:ILE:HG21	1:45:A:ILE:HG22	19	0.68
(1,420)	1:7:A:VAL:HG12	1:32:A:VAL:HG12	16	0.68
(1,357)	1:65:A:ALA:HB3	1:5:A:LEU:HB3	20	0.68
(1,174)	1:30:A:ILE:HG12	1:9:A:ALA:H	9	0.68
(1,46)	1:13:A:TYR:HB2	1:11:A:LYS:HD2	29	0.68
(1,964)	1:45:A:ILE:H	1:55:A:VAL:HA	13	0.67
(1,941)	1:33:A:LEU:H	1:33:A:LEU:HG	29	0.67
(1,921)	1:22:A:LEU:H	1:22:A:LEU:HB2	8	0.67
(1,921)	1:22:A:LEU:H	1:22:A:LEU:HB2	25	0.67
(1,917)	1:25:A:LYS:HB3	1:25:A:LYS:H	19	0.67
(1,909)	1:41:A:TRP:HD1	1:60:A:SER:HB2	28	0.67
(1,907)	1:11:A:LYS:HD2	1:11:A:LYS:HB2	28	0.67

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,906)	1:34:A:GLU:HG3	1:3:A:HIS:H	26	0.67
(1,892)	1:13:A:TYR:HD2	1:56:A:GLY:HA3	13	0.67
(1,887)	1:29:A:ILE:HG21	1:31:A:THR:HB	24	0.67
(1,859)	1:67:A:VAL:H	1:7:A:VAL:HA	3	0.67
(1,853)	1:29:A:ILE:HG12	1:48:A:ASN:HD22	26	0.67
(1,800)	1:44:A:CYS:HB3	1:55:A:VAL:HA	1	0.67
(1,800)	1:44:A:CYS:HB3	1:55:A:VAL:HA	29	0.67
(1,797)	1:7:A:VAL:HB	1:63:A:GLY:HA2	2	0.67
(1,734)	1:39:A:GLY:H	1:40:A:ARG:HB2	3	0.67
(1,714)	1:15:A:ASN:HD22	1:15:A:ASN:HA	29	0.67
(1,650)	1:12:A:ASP:H	1:11:A:LYS:HB2	12	0.67
(1,650)	1:12:A:ASP:H	1:11:A:LYS:HB2	22	0.67
(1,650)	1:12:A:ASP:H	1:11:A:LYS:HB2	30	0.67
(1,635)	1:52:A:ASN:H	1:52:A:ASN:HB2	5	0.67
(1,635)	1:52:A:ASN:H	1:52:A:ASN:HB2	20	0.67
(1,617)	1:21:A:SER:H	1:21:A:SER:HB3	11	0.67
(1,589)	1:42:A:LYS:H	1:42:A:LYS:HB3	6	0.67
(1,589)	1:42:A:LYS:H	1:42:A:LYS:HB3	10	0.67
(1,589)	1:42:A:LYS:H	1:42:A:LYS:HB3	18	0.67
(1,589)	1:42:A:LYS:H	1:42:A:LYS:HB3	27	0.67
(1,589)	1:42:A:LYS:H	1:42:A:LYS:HB3	28	0.67
(1,544)	1:40:A:ARG:H	1:40:A:ARG:HB2	13	0.67
(1,543)	1:40:A:ARG:H	1:40:A:ARG:HG2	4	0.67
(1,541)	1:40:A:ARG:H	1:40:A:ARG:HD2	26	0.67
(1,514)	1:65:A:ALA:H	1:64:A:GLU:HB2	21	0.67
(1,507)	1:22:A:LEU:H	1:22:A:LEU:HD13	26	0.67
(1,485)	1:46:A:HIS:H	1:45:A:ILE:HD13	15	0.67
(1,470)	1:30:A:ILE:H	1:30:A:ILE:HG21	1	0.67
(1,470)	1:30:A:ILE:H	1:30:A:ILE:HG21	3	0.67
(1,470)	1:30:A:ILE:H	1:30:A:ILE:HG22	26	0.67
(1,451)	1:25:A:LYS:H	1:24:A:VAL:HB	7	0.67
(1,448)	1:25:A:LYS:H	1:25:A:LYS:HB2	17	0.67
(1,437)	1:24:A:VAL:HG21	1:30:A:ILE:HG23	21	0.67
(1,431)	1:22:A:LEU:HD11	1:24:A:VAL:HG12	7	0.67
(1,431)	1:22:A:LEU:HD13	1:24:A:VAL:HG12	28	0.67
(1,420)	1:7:A:VAL:HG12	1:32:A:VAL:HG12	27	0.67
(1,391)	1:34:A:GLU:H	1:5:A:LEU:HD23	4	0.67
(1,330)	1:63:A:GLY:H	1:62:A:LEU:HB2	8	0.67
(1,318)	1:44:A:CYS:HB3	1:45:A:ILE:H	11	0.67
(1,121)	1:26:A:ALA:H	1:26:A:ALA:HB3	22	0.67
(1,121)	1:26:A:ALA:H	1:26:A:ALA:HB2	24	0.67
(1,121)	1:26:A:ALA:H	1:26:A:ALA:HB1	25	0.67

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,116)	1:36:A:HIS:H	1:35:A:GLN:HB2	29	0.67
(1,60)	1:28:A:ASP:HB2	1:28:A:ASP:HA	8	0.67
(1,60)	1:28:A:ASP:HB2	1:28:A:ASP:HA	14	0.67
(1,56)	1:28:A:ASP:H	1:28:A:ASP:HB3	14	0.67
(1,946)	1:7:A:VAL:HG23	1:30:A:ILE:H	28	0.66
(1,939)	1:58:A:PHE:HD1	1:32:A:VAL:HG13	8	0.66
(1,939)	1:58:A:PHE:HD1	1:32:A:VAL:HG11	23	0.66
(1,934)	1:26:A:ALA:H	1:25:A:LYS:HG3	7	0.66
(1,922)	1:66:A:ILE:H	1:6:A:GLN:HB3	27	0.66
(1,919)	1:23:A:ASN:HD21	1:25:A:LYS:HB3	21	0.66
(1,917)	1:54:A:ARG:HB3	1:45:A:ILE:H	4	0.66
(1,907)	1:11:A:LYS:HD2	1:11:A:LYS:HB2	19	0.66
(1,887)	1:29:A:ILE:HG22	1:31:A:THR:HB	1	0.66
(1,887)	1:29:A:ILE:HG21	1:31:A:THR:HB	10	0.66
(1,848)	1:16:A:ASN:HB2	1:18:A:ASP:H	24	0.66
(1,834)	1:51:A:GLY:HA3	1:47:A:ASP:H	28	0.66
(1,823)	1:28:A:ASP:H	1:9:A:ALA:HB2	7	0.66
(1,795)	1:17:A:TYR:HB3	1:18:A:ASP:H	4	0.66
(1,793)	1:28:A:ASP:HB3	1:24:A:VAL:HG11	13	0.66
(1,650)	1:12:A:ASP:H	1:11:A:LYS:HB2	8	0.66
(1,650)	1:12:A:ASP:H	1:11:A:LYS:HB2	9	0.66
(1,650)	1:12:A:ASP:H	1:11:A:LYS:HB2	11	0.66
(1,650)	1:12:A:ASP:H	1:11:A:LYS:HB2	14	0.66
(1,635)	1:52:A:ASN:H	1:52:A:ASN:HB2	8	0.66
(1,589)	1:42:A:LYS:H	1:42:A:LYS:HB3	11	0.66
(1,589)	1:42:A:LYS:H	1:42:A:LYS:HB3	14	0.66
(1,560)	1:64:A:GLU:H	1:64:A:GLU:HG2	20	0.66
(1,514)	1:65:A:ALA:H	1:64:A:GLU:HB2	11	0.66
(1,485)	1:46:A:HIS:H	1:45:A:ILE:HD12	24	0.66
(1,485)	1:46:A:HIS:H	1:45:A:ILE:HD12	29	0.66
(1,470)	1:30:A:ILE:H	1:30:A:ILE:HG22	5	0.66
(1,470)	1:30:A:ILE:H	1:30:A:ILE:HG21	16	0.66
(1,470)	1:30:A:ILE:H	1:30:A:ILE:HG21	28	0.66
(1,466)	1:30:A:ILE:H	1:29:A:ILE:HB	17	0.66
(1,445)	1:30:A:ILE:HD11	1:45:A:ILE:HG23	8	0.66
(1,373)	1:41:A:TRP:HE1	1:35:A:GLN:HB2	29	0.66
(1,182)	1:25:A:LYS:H	1:27:A:GLY:H	19	0.66
(1,174)	1:30:A:ILE:HG12	1:9:A:ALA:H	18	0.66
(1,121)	1:26:A:ALA:H	1:26:A:ALA:HB1	23	0.66
(1,60)	1:28:A:ASP:HB2	1:28:A:ASP:HA	9	0.66
(1,60)	1:28:A:ASP:HB2	1:28:A:ASP:HA	20	0.66
(1,27)	1:43:A:GLY:HA3	1:33:A:LEU:HD13	16	0.66

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,27)	1:43:A:GLY:HA3	1:33:A:LEU:HD11	30	0.66
(1,966)	1:8:A:ARG:H	1:64:A:GLU:HG3	26	0.65
(1,964)	1:45:A:ILE:H	1:55:A:VAL:HA	19	0.65
(1,956)	1:24:A:VAL:HG21	1:13:A:TYR:HD2	2	0.65
(1,956)	1:24:A:VAL:HG22	1:13:A:TYR:HD2	27	0.65
(1,953)	1:23:A:ASN:HB2	1:45:A:ILE:HG22	25	0.65
(1,946)	1:7:A:VAL:HG23	1:30:A:ILE:H	4	0.65
(1,944)	1:30:A:ILE:HD13	1:45:A:ILE:HD11	2	0.65
(1,921)	1:22:A:LEU:H	1:22:A:LEU:HB2	30	0.65
(1,919)	1:23:A:ASN:HD21	1:54:A:ARG:HB3	15	0.65
(1,919)	1:23:A:ASN:HD21	1:54:A:ARG:HB3	25	0.65
(1,907)	1:11:A:LYS:HD2	1:11:A:LYS:HB2	14	0.65
(1,907)	1:11:A:LYS:HD2	1:11:A:LYS:HB3	18	0.65
(1,907)	1:11:A:LYS:HD3	1:11:A:LYS:HB3	22	0.65
(1,900)	1:14:A:CYS:HB3	1:24:A:VAL:H	9	0.65
(1,894)	1:23:A:ASN:H	1:54:A:ARG:HD3	20	0.65
(1,887)	1:29:A:ILE:HG21	1:31:A:THR:HB	12	0.65
(1,870)	1:29:A:ILE:HG22	1:45:A:ILE:HA	4	0.65
(1,847)	1:8:A:ARG:HD3	1:6:A:GLN:HE21	12	0.65
(1,838)	1:36:A:HIS:HB2	1:39:A:GLY:H	1	0.65
(1,834)	1:51:A:GLY:HA3	1:47:A:ASP:H	16	0.65
(1,813)	1:22:A:LEU:HB2	1:13:A:TYR:H	14	0.65
(1,800)	1:44:A:CYS:HB2	1:55:A:VAL:HA	3	0.65
(1,707)	1:35:A:GLN:HE22	1:35:A:GLN:HB3	14	0.65
(1,687)	1:27:A:GLY:H	1:26:A:ALA:HB3	16	0.65
(1,659)	1:11:A:LYS:H	1:10:A:THR:H	4	0.65
(1,650)	1:12:A:ASP:H	1:11:A:LYS:HB2	3	0.65
(1,650)	1:12:A:ASP:H	1:11:A:LYS:HB2	27	0.65
(1,582)	1:13:A:TYR:H	1:13:A:TYR:HB3	22	0.65
(1,575)	1:36:A:HIS:H	1:41:A:TRP:HA	27	0.65
(1,572)	1:66:A:ILE:H	1:66:A:ILE:HG12	12	0.65
(1,544)	1:40:A:ARG:H	1:40:A:ARG:HB2	9	0.65
(1,493)	1:41:A:TRP:HE1	1:40:A:ARG:H	24	0.65
(1,480)	1:45:A:ILE:H	1:44:A:CYS:HB2	11	0.65
(1,480)	1:45:A:ILE:H	1:44:A:CYS:HB2	30	0.65
(1,470)	1:30:A:ILE:H	1:30:A:ILE:HG21	29	0.65
(1,467)	1:30:A:ILE:H	1:29:A:ILE:HG22	6	0.65
(1,466)	1:30:A:ILE:H	1:29:A:ILE:HB	15	0.65
(1,464)	1:67:A:VAL:H	1:66:A:ILE:HG12	27	0.65
(1,441)	1:30:A:ILE:HG23	1:45:A:ILE:HG23	21	0.65
(1,441)	1:30:A:ILE:HG21	1:45:A:ILE:HG22	23	0.65
(1,434)	1:24:A:VAL:HG12	1:30:A:ILE:HD13	7	0.65

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,420)	1:7:A:VAL:HG12	1:32:A:VAL:HG12	19	0.65
(1,420)	1:7:A:VAL:HG13	1:32:A:VAL:HG11	26	0.65
(1,399)	1:29:A:ILE:H	1:29:A:ILE:HD11	7	0.65
(1,382)	1:65:A:ALA:H	1:5:A:LEU:HB3	29	0.65
(1,360)	1:27:A:GLY:H	1:25:A:LYS:HG3	1	0.65
(1,174)	1:30:A:ILE:HG12	1:9:A:ALA:H	14	0.65
(1,135)	1:8:A:ARG:HA	1:27:A:GLY:HA3	8	0.65
(1,121)	1:26:A:ALA:H	1:26:A:ALA:HB3	5	0.65
(1,121)	1:26:A:ALA:H	1:26:A:ALA:HB3	17	0.65
(1,121)	1:26:A:ALA:H	1:26:A:ALA:HB2	26	0.65
(1,60)	1:28:A:ASP:HB2	1:28:A:ASP:HA	19	0.65
(1,56)	1:28:A:ASP:H	1:28:A:ASP:HB3	8	0.65
(1,27)	1:43:A:GLY:HA3	1:33:A:LEU:HD13	2	0.65
(1,964)	1:45:A:ILE:H	1:55:A:VAL:HA	22	0.64
(1,956)	1:24:A:VAL:HG21	1:13:A:TYR:HD2	6	0.64
(1,953)	1:23:A:ASN:HB2	1:45:A:ILE:HG21	9	0.64
(1,953)	1:23:A:ASN:HB2	1:45:A:ILE:HG21	22	0.64
(1,941)	1:33:A:LEU:H	1:33:A:LEU:HG	20	0.64
(1,940)	1:7:A:VAL:H	1:32:A:VAL:HG12	9	0.64
(1,923)	1:33:A:LEU:HD13	1:33:A:LEU:HB3	8	0.64
(1,917)	1:25:A:LYS:HB3	1:25:A:LYS:H	29	0.64
(1,912)	1:41:A:TRP:H	1:40:A:ARG:HG3	1	0.64
(1,907)	1:11:A:LYS:HD2	1:11:A:LYS:HB2	7	0.64
(1,907)	1:11:A:LYS:HD2	1:11:A:LYS:HB2	11	0.64
(1,907)	1:11:A:LYS:HD2	1:11:A:LYS:HB3	12	0.64
(1,907)	1:11:A:LYS:HD2	1:11:A:LYS:HB3	24	0.64
(1,900)	1:14:A:CYS:HB3	1:24:A:VAL:H	23	0.64
(1,896)	1:48:A:ASN:H	1:47:A:ASP:HB2	14	0.64
(1,859)	1:67:A:VAL:H	1:7:A:VAL:HA	24	0.64
(1,823)	1:28:A:ASP:H	1:9:A:ALA:HB2	17	0.64
(1,795)	1:17:A:TYR:HB3	1:18:A:ASP:H	3	0.64
(1,714)	1:15:A:ASN:HD22	1:15:A:ASN:HA	24	0.64
(1,659)	1:11:A:LYS:H	1:10:A:THR:H	9	0.64
(1,650)	1:12:A:ASP:H	1:11:A:LYS:HB2	23	0.64
(1,635)	1:52:A:ASN:H	1:52:A:ASN:HB2	7	0.64
(1,635)	1:52:A:ASN:H	1:52:A:ASN:HB2	11	0.64
(1,635)	1:52:A:ASN:H	1:52:A:ASN:HB2	30	0.64
(1,627)	1:38:A:ASP:H	1:37:A:PRO:HB3	12	0.64
(1,544)	1:40:A:ARG:H	1:40:A:ARG:HB2	7	0.64
(1,514)	1:65:A:ALA:H	1:64:A:GLU:HB2	24	0.64
(1,507)	1:22:A:LEU:H	1:22:A:LEU:HD11	13	0.64
(1,493)	1:41:A:TRP:HE1	1:40:A:ARG:H	11	0.64

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,470)	1:30:A:ILE:H	1:30:A:ILE:HG21	9	0.64
(1,470)	1:30:A:ILE:H	1:30:A:ILE:HG22	11	0.64
(1,470)	1:30:A:ILE:H	1:30:A:ILE:HG22	15	0.64
(1,470)	1:30:A:ILE:H	1:30:A:ILE:HG23	17	0.64
(1,470)	1:30:A:ILE:H	1:30:A:ILE:HG23	24	0.64
(1,451)	1:25:A:LYS:H	1:24:A:VAL:HB	6	0.64
(1,448)	1:25:A:LYS:H	1:25:A:LYS:HB2	1	0.64
(1,448)	1:25:A:LYS:H	1:25:A:LYS:HB2	4	0.64
(1,445)	1:30:A:ILE:HD13	1:45:A:ILE:HG22	19	0.64
(1,445)	1:30:A:ILE:HD12	1:45:A:ILE:HG23	27	0.64
(1,427)	1:20:A:THR:HG21	1:55:A:VAL:HG23	14	0.64
(1,392)	1:43:A:GLY:H	1:33:A:LEU:HD13	18	0.64
(1,360)	1:27:A:GLY:H	1:25:A:LYS:HG3	5	0.64
(1,228)	1:51:A:GLY:H	1:52:A:ASN:HD22	11	0.64
(1,196)	1:34:A:GLU:HB3	1:4:A:MET:H	3	0.64
(1,182)	1:25:A:LYS:H	1:27:A:GLY:H	26	0.64
(1,121)	1:26:A:ALA:H	1:26:A:ALA:HB3	2	0.64
(1,121)	1:26:A:ALA:H	1:26:A:ALA:HB2	3	0.64
(1,121)	1:26:A:ALA:H	1:26:A:ALA:HB2	29	0.64
(1,56)	1:28:A:ASP:H	1:28:A:ASP:HB3	20	0.64
(1,27)	1:43:A:GLY:HA3	1:33:A:LEU:HD11	9	0.64
(1,968)	1:10:A:THR:H	1:26:A:ALA:HA	26	0.63
(1,934)	1:26:A:ALA:H	1:25:A:LYS:HG3	6	0.63
(1,917)	1:25:A:LYS:HB3	1:25:A:LYS:H	13	0.63
(1,912)	1:41:A:TRP:H	1:40:A:ARG:HG3	17	0.63
(1,908)	1:34:A:GLU:H	1:35:A:GLN:HG3	12	0.63
(1,907)	1:11:A:LYS:HD2	1:11:A:LYS:HB2	6	0.63
(1,907)	1:11:A:LYS:HD3	1:11:A:LYS:HB3	15	0.63
(1,907)	1:11:A:LYS:HD2	1:11:A:LYS:HB2	17	0.63
(1,907)	1:11:A:LYS:HD2	1:11:A:LYS:HB2	20	0.63
(1,907)	1:11:A:LYS:HD2	1:11:A:LYS:HB2	23	0.63
(1,892)	1:41:A:TRP:HH2	1:61:A:SER:HA	23	0.63
(1,852)	1:25:A:LYS:H	1:48:A:ASN:HD22	16	0.63
(1,852)	1:13:A:TYR:HE2	1:24:A:VAL:H	27	0.63
(1,834)	1:51:A:GLY:HA3	1:47:A:ASP:H	23	0.63
(1,810)	1:23:A:ASN:H	1:45:A:ILE:H	1	0.63
(1,810)	1:23:A:ASN:H	1:45:A:ILE:H	29	0.63
(1,802)	1:26:A:ALA:HB2	1:12:A:ASP:HA	15	0.63
(1,574)	1:36:A:HIS:H	1:35:A:GLN:HB2	10	0.63
(1,541)	1:40:A:ARG:H	1:40:A:ARG:HD2	19	0.63
(1,514)	1:65:A:ALA:H	1:64:A:GLU:HB2	30	0.63
(1,507)	1:22:A:LEU:H	1:22:A:LEU:HD13	18	0.63

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,507)	1:22:A:LEU:H	1:22:A:LEU:HD12	23	0.63
(1,470)	1:30:A:ILE:H	1:30:A:ILE:HG21	7	0.63
(1,470)	1:30:A:ILE:H	1:30:A:ILE:HG22	12	0.63
(1,470)	1:30:A:ILE:H	1:30:A:ILE:HG22	14	0.63
(1,470)	1:30:A:ILE:H	1:30:A:ILE:HG23	22	0.63
(1,470)	1:30:A:ILE:H	1:30:A:ILE:HG22	23	0.63
(1,470)	1:30:A:ILE:H	1:30:A:ILE:HG22	25	0.63
(1,470)	1:30:A:ILE:H	1:30:A:ILE:HG22	30	0.63
(1,466)	1:30:A:ILE:H	1:29:A:ILE:HB	12	0.63
(1,451)	1:25:A:LYS:H	1:24:A:VAL:HB	24	0.63
(1,448)	1:25:A:LYS:H	1:25:A:LYS:HB2	8	0.63
(1,448)	1:25:A:LYS:H	1:25:A:LYS:HB2	18	0.63
(1,448)	1:25:A:LYS:H	1:25:A:LYS:HB2	19	0.63
(1,444)	1:30:A:ILE:HD11	1:32:A:VAL:HG12	1	0.63
(1,434)	1:24:A:VAL:HG12	1:30:A:ILE:HD12	28	0.63
(1,427)	1:20:A:THR:HG23	1:55:A:VAL:HG21	29	0.63
(1,420)	1:7:A:VAL:HG12	1:32:A:VAL:HG11	12	0.63
(1,416)	1:5:A:LEU:HD21	1:7:A:VAL:HG12	16	0.63
(1,403)	1:58:A:PHE:H	1:22:A:LEU:HD13	15	0.63
(1,392)	1:43:A:GLY:H	1:33:A:LEU:HD12	17	0.63
(1,174)	1:30:A:ILE:HG12	1:9:A:ALA:H	8	0.63
(1,66)	1:28:A:ASP:HB3	1:25:A:LYS:HG2	30	0.63
(1,966)	1:8:A:ARG:H	1:64:A:GLU:HG2	15	0.62
(1,954)	1:44:A:CYS:HA	1:45:A:ILE:HD13	3	0.62
(1,946)	1:7:A:VAL:HG21	1:30:A:ILE:H	6	0.62
(1,946)	1:7:A:VAL:HG23	1:30:A:ILE:H	18	0.62
(1,945)	1:41:A:TRP:HH2	1:5:A:LEU:HD11	28	0.62
(1,944)	1:30:A:ILE:HD12	1:7:A:VAL:HG21	19	0.62
(1,935)	1:40:A:ARG:HB3	1:38:A:ASP:HB3	19	0.62
(1,909)	1:41:A:TRP:HD1	1:40:A:ARG:HG2	23	0.62
(1,907)	1:11:A:LYS:HD2	1:11:A:LYS:HB2	2	0.62
(1,907)	1:11:A:LYS:HD2	1:11:A:LYS:HB2	5	0.62
(1,907)	1:11:A:LYS:HD2	1:11:A:LYS:HB2	27	0.62
(1,881)	1:13:A:TYR:HD2	1:21:A:SER:HB2	24	0.62
(1,873)	1:41:A:TRP:HB3	1:34:A:GLU:HA	29	0.62
(1,840)	1:11:A:LYS:HD3	1:12:A:ASP:H	5	0.62
(1,834)	1:51:A:GLY:HA3	1:47:A:ASP:H	27	0.62
(1,814)	1:26:A:ALA:H	1:25:A:LYS:HB2	30	0.62
(1,810)	1:23:A:ASN:H	1:45:A:ILE:H	17	0.62
(1,803)	1:30:A:ILE:HG22	1:32:A:VAL:H	11	0.62
(1,775)	1:38:A:ASP:H	1:40:A:ARG:H	28	0.62
(1,734)	1:39:A:GLY:H	1:40:A:ARG:HB2	8	0.62

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,734)	1:39:A:GLY:H	1:40:A:ARG:HB2	26	0.62
(1,707)	1:35:A:GLN:HE22	1:35:A:GLN:HB3	15	0.62
(1,670)	1:43:A:GLY:H	1:42:A:LYS:HB2	4	0.62
(1,650)	1:12:A:ASP:H	1:11:A:LYS:HB2	1	0.62
(1,650)	1:12:A:ASP:H	1:11:A:LYS:HB2	20	0.62
(1,650)	1:12:A:ASP:H	1:11:A:LYS:HB2	28	0.62
(1,635)	1:52:A:ASN:H	1:52:A:ASN:HB2	2	0.62
(1,575)	1:36:A:HIS:H	1:41:A:TRP:HA	29	0.62
(1,562)	1:9:A:ALA:H	1:28:A:ASP:H	11	0.62
(1,514)	1:65:A:ALA:H	1:64:A:GLU:HB2	2	0.62
(1,514)	1:65:A:ALA:H	1:64:A:GLU:HB2	28	0.62
(1,507)	1:22:A:LEU:H	1:22:A:LEU:HD13	3	0.62
(1,496)	1:41:A:TRP:H	1:41:A:TRP:HB3	20	0.62
(1,480)	1:45:A:ILE:H	1:44:A:CYS:HB2	10	0.62
(1,470)	1:30:A:ILE:H	1:30:A:ILE:HG23	6	0.62
(1,470)	1:30:A:ILE:H	1:30:A:ILE:HG23	8	0.62
(1,470)	1:30:A:ILE:H	1:30:A:ILE:HG22	13	0.62
(1,470)	1:30:A:ILE:H	1:30:A:ILE:HG21	21	0.62
(1,467)	1:30:A:ILE:H	1:29:A:ILE:HG23	29	0.62
(1,466)	1:30:A:ILE:H	1:29:A:ILE:HB	25	0.62
(1,451)	1:25:A:LYS:H	1:24:A:VAL:HB	3	0.62
(1,447)	1:25:A:LYS:H	1:25:A:LYS:HG2	19	0.62
(1,441)	1:30:A:ILE:HG22	1:45:A:ILE:HG23	8	0.62
(1,318)	1:44:A:CYS:HB3	1:45:A:ILE:H	3	0.62
(1,220)	1:15:A:ASN:H	1:15:A:ASN:HD22	24	0.62
(1,66)	1:28:A:ASP:HB3	1:25:A:LYS:HG2	3	0.62
(1,46)	1:13:A:TYR:HB2	1:11:A:LYS:HD2	7	0.62
(1,46)	1:13:A:TYR:HB2	1:11:A:LYS:HD2	11	0.62
(1,966)	1:8:A:ARG:H	1:64:A:GLU:HG3	8	0.61
(1,946)	1:7:A:VAL:HG22	1:30:A:ILE:H	27	0.61
(1,945)	1:41:A:TRP:HH2	1:5:A:LEU:HD12	11	0.61
(1,945)	1:41:A:TRP:HH2	1:5:A:LEU:HD12	26	0.61
(1,940)	1:7:A:VAL:H	1:32:A:VAL:HG13	14	0.61
(1,912)	1:41:A:TRP:H	1:40:A:ARG:HG3	12	0.61
(1,912)	1:41:A:TRP:H	1:40:A:ARG:HG3	24	0.61
(1,907)	1:11:A:LYS:HD2	1:11:A:LYS:HB2	16	0.61
(1,907)	1:11:A:LYS:HD2	1:11:A:LYS:HB2	25	0.61
(1,907)	1:11:A:LYS:HD2	1:11:A:LYS:HB2	26	0.61
(1,840)	1:11:A:LYS:HD3	1:12:A:ASP:H	28	0.61
(1,834)	1:51:A:GLY:HA3	1:47:A:ASP:H	1	0.61
(1,685)	1:27:A:GLY:H	1:24:A:VAL:HG23	9	0.61
(1,670)	1:43:A:GLY:H	1:42:A:LYS:HB2	16	0.61

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,650)	1:12:A:ASP:H	1:11:A:LYS:HB2	2	0.61
(1,635)	1:52:A:ASN:H	1:52:A:ASN:HB2	14	0.61
(1,515)	1:14:A:CYS:H	1:13:A:TYR:HB3	1	0.61
(1,507)	1:22:A:LEU:H	1:22:A:LEU:HD13	4	0.61
(1,496)	1:41:A:TRP:H	1:41:A:TRP:HB3	9	0.61
(1,470)	1:30:A:ILE:H	1:30:A:ILE:HG22	4	0.61
(1,470)	1:30:A:ILE:H	1:30:A:ILE:HG23	20	0.61
(1,466)	1:30:A:ILE:H	1:29:A:ILE:HB	24	0.61
(1,451)	1:25:A:LYS:H	1:24:A:VAL:HB	22	0.61
(1,448)	1:25:A:LYS:H	1:25:A:LYS:HB2	23	0.61
(1,437)	1:24:A:VAL:HG23	1:30:A:ILE:HG21	12	0.61
(1,330)	1:63:A:GLY:H	1:62:A:LEU:HB2	9	0.61
(1,318)	1:44:A:CYS:HB3	1:45:A:ILE:H	10	0.61
(1,234)	1:8:A:ARG:HG3	1:6:A:GLN:HE22	21	0.61
(1,968)	1:10:A:THR:H	1:26:A:ALA:HA	24	0.6
(1,964)	1:45:A:ILE:H	1:53:A:ASP:HA	17	0.6
(1,931)	1:6:A:GLN:HE21	1:8:A:ARG:HG2	10	0.6
(1,909)	1:41:A:TRP:HD1	1:40:A:ARG:HG2	1	0.6
(1,907)	1:11:A:LYS:HD2	1:11:A:LYS:HB2	3	0.6
(1,907)	1:11:A:LYS:HD2	1:11:A:LYS:HB2	4	0.6
(1,852)	1:25:A:LYS:H	1:48:A:ASN:HD22	9	0.6
(1,834)	1:51:A:GLY:HA3	1:47:A:ASP:H	25	0.6
(1,814)	1:26:A:ALA:H	1:25:A:LYS:HB2	7	0.6
(1,803)	1:30:A:ILE:HG23	1:32:A:VAL:H	17	0.6
(1,775)	1:38:A:ASP:H	1:40:A:ARG:H	11	0.6
(1,714)	1:15:A:ASN:HD22	1:15:A:ASN:HA	18	0.6
(1,707)	1:35:A:GLN:HE22	1:35:A:GLN:HB3	13	0.6
(1,687)	1:27:A:GLY:H	1:26:A:ALA:HB1	18	0.6
(1,687)	1:27:A:GLY:H	1:26:A:ALA:HB2	23	0.6
(1,687)	1:27:A:GLY:H	1:26:A:ALA:HB2	25	0.6
(1,687)	1:27:A:GLY:H	1:26:A:ALA:HB1	27	0.6
(1,670)	1:43:A:GLY:H	1:42:A:LYS:HB2	14	0.6
(1,670)	1:43:A:GLY:H	1:42:A:LYS:HB2	22	0.6
(1,496)	1:41:A:TRP:H	1:41:A:TRP:HB3	12	0.6
(1,496)	1:41:A:TRP:H	1:41:A:TRP:HB3	21	0.6
(1,485)	1:46:A:HIS:H	1:45:A:ILE:HD11	11	0.6
(1,480)	1:45:A:ILE:H	1:44:A:CYS:HB2	3	0.6
(1,470)	1:30:A:ILE:H	1:30:A:ILE:HG22	18	0.6
(1,470)	1:30:A:ILE:H	1:30:A:ILE:HG23	27	0.6
(1,466)	1:30:A:ILE:H	1:29:A:ILE:HB	26	0.6
(1,441)	1:30:A:ILE:HG21	1:45:A:ILE:HG21	15	0.6
(1,422)	1:7:A:VAL:HG21	1:32:A:VAL:HG13	18	0.6

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,399)	1:29:A:ILE:H	1:29:A:ILE:HD13	22	0.6
(1,318)	1:44:A:CYS:HB3	1:45:A:ILE:H	14	0.6
(1,241)	1:4:A:MET:H	1:2:A:SER:HA	9	0.6
(1,190)	1:34:A:GLU:HG2	1:3:A:HIS:H	12	0.6
(1,174)	1:30:A:ILE:HG12	1:9:A:ALA:H	17	0.6
(1,60)	1:28:A:ASP:HB2	1:28:A:ASP:HA	30	0.6
(1,57)	1:28:A:ASP:HB2	1:25:A:LYS:HG2	18	0.6
(1,941)	1:33:A:LEU:H	1:33:A:LEU:HG	27	0.59
(1,940)	1:7:A:VAL:H	1:32:A:VAL:HG11	30	0.59
(1,939)	1:58:A:PHE:HD1	1:32:A:VAL:HG12	7	0.59
(1,907)	1:11:A:LYS:HD2	1:11:A:LYS:HB2	29	0.59
(1,863)	1:9:A:ALA:H	1:29:A:ILE:HA	5	0.59
(1,840)	1:11:A:LYS:HD3	1:12:A:ASP:H	12	0.59
(1,814)	1:26:A:ALA:H	1:25:A:LYS:HB2	10	0.59
(1,800)	1:44:A:CYS:HB3	1:55:A:VAL:HA	22	0.59
(1,710)	1:6:A:GLN:HE21	1:29:A:ILE:HG12	29	0.59
(1,707)	1:35:A:GLN:HE22	1:35:A:GLN:HB3	19	0.59
(1,670)	1:43:A:GLY:H	1:42:A:LYS:HB2	25	0.59
(1,670)	1:43:A:GLY:H	1:42:A:LYS:HB2	27	0.59
(1,659)	1:11:A:LYS:H	1:10:A:THR:H	21	0.59
(1,659)	1:11:A:LYS:H	1:10:A:THR:H	24	0.59
(1,659)	1:11:A:LYS:H	1:10:A:THR:H	26	0.59
(1,659)	1:11:A:LYS:H	1:10:A:THR:H	28	0.59
(1,650)	1:12:A:ASP:H	1:11:A:LYS:HB2	19	0.59
(1,635)	1:52:A:ASN:H	1:52:A:ASN:HB2	6	0.59
(1,587)	1:53:A:ASP:H	1:53:A:ASP:HB2	27	0.59
(1,560)	1:64:A:GLU:H	1:64:A:GLU:HG2	10	0.59
(1,541)	1:40:A:ARG:H	1:40:A:ARG:HD2	20	0.59
(1,515)	1:14:A:CYS:H	1:13:A:TYR:HB3	24	0.59
(1,514)	1:65:A:ALA:H	1:64:A:GLU:HB2	16	0.59
(1,496)	1:41:A:TRP:H	1:41:A:TRP:HB3	1	0.59
(1,496)	1:41:A:TRP:H	1:41:A:TRP:HB3	19	0.59
(1,485)	1:46:A:HIS:H	1:45:A:ILE:HD12	14	0.59
(1,480)	1:45:A:ILE:H	1:44:A:CYS:HB2	14	0.59
(1,470)	1:30:A:ILE:H	1:30:A:ILE:HG22	19	0.59
(1,466)	1:30:A:ILE:H	1:29:A:ILE:HB	7	0.59
(1,448)	1:25:A:LYS:H	1:25:A:LYS:HB2	10	0.59
(1,448)	1:25:A:LYS:H	1:25:A:LYS:HB2	13	0.59
(1,431)	1:22:A:LEU:HD11	1:24:A:VAL:HG13	29	0.59
(1,420)	1:7:A:VAL:HG13	1:32:A:VAL:HG12	17	0.59
(1,416)	1:5:A:LEU:HD21	1:7:A:VAL:HG12	30	0.59
(1,403)	1:58:A:PHE:H	1:22:A:LEU:HD12	30	0.59

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,399)	1:29:A:ILE:H	1:29:A:ILE:HD12	9	0.59
(1,390)	1:24:A:VAL:HG21	1:13:A:TYR:HB3	22	0.59
(1,176)	1:33:A:LEU:H	1:42:A:LYS:H	21	0.59
(1,46)	1:13:A:TYR:HB2	1:11:A:LYS:HD2	5	0.59
(1,964)	1:45:A:ILE:H	1:53:A:ASP:HA	1	0.58
(1,948)	1:6:A:GLN:H	1:67:A:VAL:HG21	10	0.58
(1,945)	1:41:A:TRP:HH2	1:5:A:LEU:HD12	2	0.58
(1,944)	1:30:A:ILE:HD12	1:7:A:VAL:HG22	14	0.58
(1,934)	1:26:A:ALA:H	1:25:A:LYS:HG3	30	0.58
(1,919)	1:23:A:ASN:HD21	1:54:A:ARG:HB2	8	0.58
(1,907)	1:11:A:LYS:HD2	1:11:A:LYS:HB3	13	0.58
(1,863)	1:9:A:ALA:H	1:29:A:ILE:HA	3	0.58
(1,803)	1:30:A:ILE:HG23	1:32:A:VAL:H	27	0.58
(1,800)	1:44:A:CYS:HB2	1:55:A:VAL:HA	11	0.58
(1,687)	1:27:A:GLY:H	1:26:A:ALA:HB3	30	0.58
(1,659)	1:11:A:LYS:H	1:10:A:THR:H	12	0.58
(1,560)	1:64:A:GLU:H	1:64:A:GLU:HG2	21	0.58
(1,560)	1:64:A:GLU:H	1:64:A:GLU:HG2	25	0.58
(1,541)	1:40:A:ARG:H	1:40:A:ARG:HD2	21	0.58
(1,541)	1:40:A:ARG:H	1:40:A:ARG:HD2	25	0.58
(1,515)	1:14:A:CYS:H	1:13:A:TYR:HB3	19	0.58
(1,507)	1:22:A:LEU:H	1:22:A:LEU:HD12	16	0.58
(1,496)	1:41:A:TRP:H	1:41:A:TRP:HB3	2	0.58
(1,496)	1:41:A:TRP:H	1:41:A:TRP:HB3	16	0.58
(1,496)	1:41:A:TRP:H	1:41:A:TRP:HB3	18	0.58
(1,496)	1:41:A:TRP:H	1:41:A:TRP:HB3	25	0.58
(1,496)	1:41:A:TRP:H	1:41:A:TRP:HB3	30	0.58
(1,493)	1:41:A:TRP:HE1	1:40:A:ARG:H	6	0.58
(1,480)	1:45:A:ILE:H	1:44:A:CYS:HB2	9	0.58
(1,470)	1:30:A:ILE:H	1:30:A:ILE:HG22	2	0.58
(1,470)	1:30:A:ILE:H	1:30:A:ILE:HG21	10	0.58
(1,467)	1:30:A:ILE:H	1:29:A:ILE:HG23	21	0.58
(1,467)	1:30:A:ILE:H	1:29:A:ILE:HG22	27	0.58
(1,451)	1:25:A:LYS:H	1:24:A:VAL:HB	19	0.58
(1,448)	1:25:A:LYS:H	1:25:A:LYS:HB2	24	0.58
(1,441)	1:30:A:ILE:HG23	1:45:A:ILE:HG21	1	0.58
(1,441)	1:30:A:ILE:HG21	1:45:A:ILE:HG21	11	0.58
(1,431)	1:22:A:LEU:HD11	1:24:A:VAL:HG12	26	0.58
(1,420)	1:7:A:VAL:HG13	1:32:A:VAL:HG12	11	0.58
(1,420)	1:7:A:VAL:HG13	1:32:A:VAL:HG12	14	0.58
(1,318)	1:44:A:CYS:HB3	1:45:A:ILE:H	9	0.58
(1,317)	1:39:A:GLY:H	1:38:A:ASP:HB3	4	0.58

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,317)	1:39:A:GLY:H	1:38:A:ASP:HB3	8	0.58
(1,220)	1:15:A:ASN:H	1:15:A:ASN:HD22	8	0.58
(1,211)	1:25:A:LYS:H	1:11:A:LYS:H	5	0.58
(1,62)	1:18:A:ASP:HB2	1:18:A:ASP:H	10	0.58
(1,46)	1:13:A:TYR:HB2	1:11:A:LYS:HD2	15	0.58
(1,967)	1:62:A:LEU:H	1:63:A:GLY:HA2	24	0.57
(1,964)	1:45:A:ILE:H	1:55:A:VAL:HA	15	0.57
(1,946)	1:7:A:VAL:HG23	1:30:A:ILE:H	5	0.57
(1,946)	1:7:A:VAL:HG21	1:30:A:ILE:H	21	0.57
(1,945)	1:41:A:TRP:HH2	1:5:A:LEU:HD11	12	0.57
(1,940)	1:7:A:VAL:H	1:32:A:VAL:HG13	13	0.57
(1,929)	1:13:A:TYR:HD1	1:11:A:LYS:HG2	4	0.57
(1,925)	1:11:A:LYS:HG2	1:13:A:TYR:H	18	0.57
(1,919)	1:23:A:ASN:HD21	1:25:A:LYS:HB3	18	0.57
(1,908)	1:34:A:GLU:H	1:35:A:GLN:HG2	11	0.57
(1,907)	1:11:A:LYS:HD2	1:11:A:LYS:HB3	10	0.57
(1,899)	1:15:A:ASN:H	1:14:A:CYS:HB3	25	0.57
(1,814)	1:26:A:ALA:H	1:25:A:LYS:HB2	27	0.57
(1,813)	1:22:A:LEU:HB2	1:13:A:TYR:H	6	0.57
(1,800)	1:44:A:CYS:HB3	1:55:A:VAL:HA	2	0.57
(1,743)	1:63:A:GLY:H	1:62:A:LEU:HG	6	0.57
(1,687)	1:27:A:GLY:H	1:26:A:ALA:HB1	10	0.57
(1,687)	1:27:A:GLY:H	1:26:A:ALA:HB3	29	0.57
(1,635)	1:52:A:ASN:H	1:52:A:ASN:HB2	17	0.57
(1,541)	1:40:A:ARG:H	1:40:A:ARG:HD2	10	0.57
(1,507)	1:22:A:LEU:H	1:22:A:LEU:HD13	6	0.57
(1,507)	1:22:A:LEU:H	1:22:A:LEU:HD13	27	0.57
(1,496)	1:41:A:TRP:H	1:41:A:TRP:HB3	17	0.57
(1,496)	1:41:A:TRP:H	1:41:A:TRP:HB3	23	0.57
(1,466)	1:30:A:ILE:H	1:29:A:ILE:HB	29	0.57
(1,451)	1:25:A:LYS:H	1:24:A:VAL:HB	10	0.57
(1,448)	1:25:A:LYS:H	1:25:A:LYS:HB2	12	0.57
(1,448)	1:25:A:LYS:H	1:25:A:LYS:HB2	26	0.57
(1,448)	1:25:A:LYS:H	1:25:A:LYS:HB2	28	0.57
(1,434)	1:24:A:VAL:HG11	1:30:A:ILE:HD11	15	0.57
(1,431)	1:22:A:LEU:HD11	1:24:A:VAL:HG12	4	0.57
(1,420)	1:7:A:VAL:HG13	1:32:A:VAL:HG11	20	0.57
(1,403)	1:58:A:PHE:H	1:22:A:LEU:HD12	6	0.57
(1,357)	1:65:A:ALA:HB1	1:5:A:LEU:HB3	27	0.57
(1,299)	1:50:A:THR:H	1:47:A:ASP:HB2	3	0.57
(1,176)	1:33:A:LEU:H	1:42:A:LYS:H	9	0.57
(1,109)	1:32:A:VAL:H	1:32:A:VAL:HB	1	0.57

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,109)	1:32:A:VAL:H	1:32:A:VAL:HB	11	0.57
(1,56)	1:28:A:ASP:H	1:28:A:ASP:HB3	9	0.57
(1,39)	1:54:A:ARG:H	1:53:A:ASP:HB2	20	0.57
(1,966)	1:8:A:ARG:H	1:64:A:GLU:HG2	9	0.56
(1,966)	1:8:A:ARG:H	1:64:A:GLU:HG3	30	0.56
(1,965)	1:26:A:ALA:H	1:12:A:ASP:HA	10	0.56
(1,946)	1:7:A:VAL:HG23	1:30:A:ILE:H	12	0.56
(1,945)	1:41:A:TRP:HH2	1:5:A:LEU:HD12	7	0.56
(1,934)	1:26:A:ALA:H	1:25:A:LYS:HG3	8	0.56
(1,925)	1:11:A:LYS:HG2	1:13:A:TYR:H	5	0.56
(1,925)	1:11:A:LYS:HG2	1:13:A:TYR:H	9	0.56
(1,907)	1:11:A:LYS:HD2	1:11:A:LYS:HB2	9	0.56
(1,907)	1:11:A:LYS:HD2	1:11:A:LYS:HB2	21	0.56
(1,907)	1:11:A:LYS:HD2	1:11:A:LYS:HB2	30	0.56
(1,878)	1:52:A:ASN:H	1:50:A:THR:HA	12	0.56
(1,878)	1:52:A:ASN:H	1:50:A:THR:HA	14	0.56
(1,866)	1:66:A:ILE:HG22	1:65:A:ALA:HA	29	0.56
(1,847)	1:8:A:ARG:HD3	1:6:A:GLN:HE21	25	0.56
(1,840)	1:11:A:LYS:HD3	1:12:A:ASP:H	23	0.56
(1,814)	1:26:A:ALA:H	1:25:A:LYS:HB2	2	0.56
(1,814)	1:26:A:ALA:H	1:25:A:LYS:HB2	8	0.56
(1,814)	1:26:A:ALA:H	1:25:A:LYS:HB2	11	0.56
(1,814)	1:26:A:ALA:H	1:25:A:LYS:HB2	20	0.56
(1,803)	1:30:A:ILE:HG21	1:32:A:VAL:H	1	0.56
(1,800)	1:44:A:CYS:HB3	1:55:A:VAL:HA	15	0.56
(1,795)	1:17:A:TYR:HB3	1:18:A:ASP:H	29	0.56
(1,793)	1:28:A:ASP:HB3	1:24:A:VAL:HG12	26	0.56
(1,687)	1:27:A:GLY:H	1:26:A:ALA:HB3	24	0.56
(1,659)	1:11:A:LYS:H	1:10:A:THR:H	16	0.56
(1,659)	1:11:A:LYS:H	1:10:A:THR:H	18	0.56
(1,659)	1:11:A:LYS:H	1:10:A:THR:H	27	0.56
(1,650)	1:12:A:ASP:H	1:11:A:LYS:HB2	13	0.56
(1,650)	1:12:A:ASP:H	1:11:A:LYS:HB2	18	0.56
(1,650)	1:12:A:ASP:H	1:11:A:LYS:HB2	29	0.56
(1,635)	1:52:A:ASN:H	1:52:A:ASN:HB2	12	0.56
(1,544)	1:40:A:ARG:H	1:40:A:ARG:HB2	17	0.56
(1,544)	1:40:A:ARG:H	1:40:A:ARG:HB2	21	0.56
(1,496)	1:41:A:TRP:H	1:41:A:TRP:HB3	5	0.56
(1,496)	1:41:A:TRP:H	1:41:A:TRP:HB3	6	0.56
(1,496)	1:41:A:TRP:H	1:41:A:TRP:HB3	11	0.56
(1,480)	1:45:A:ILE:H	1:44:A:CYS:HB2	18	0.56
(1,467)	1:30:A:ILE:H	1:29:A:ILE:HG21	23	0.56

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,466)	1:30:A:ILE:H	1:29:A:ILE:HB	3	0.56
(1,466)	1:30:A:ILE:H	1:29:A:ILE:HB	6	0.56
(1,466)	1:30:A:ILE:H	1:29:A:ILE:HB	22	0.56
(1,451)	1:25:A:LYS:H	1:24:A:VAL:HB	29	0.56
(1,403)	1:58:A:PHE:H	1:22:A:LEU:HD11	8	0.56
(1,395)	1:42:A:LYS:H	1:33:A:LEU:HD11	9	0.56
(1,109)	1:32:A:VAL:H	1:32:A:VAL:HB	17	0.56
(1,109)	1:32:A:VAL:H	1:32:A:VAL:HB	20	0.56
(1,109)	1:32:A:VAL:H	1:32:A:VAL:HB	27	0.56
(1,27)	1:43:A:GLY:HA3	1:33:A:LEU:HD13	15	0.56
(1,964)	1:45:A:ILE:H	1:55:A:VAL:HA	7	0.55
(1,958)	1:61:A:SER:H	1:5:A:LEU:HD23	28	0.55
(1,946)	1:7:A:VAL:HG22	1:30:A:ILE:H	8	0.55
(1,945)	1:41:A:TRP:HH2	1:5:A:LEU:HD11	18	0.55
(1,944)	1:30:A:ILE:HD13	1:7:A:VAL:HG21	9	0.55
(1,941)	1:33:A:LEU:H	1:33:A:LEU:HG	18	0.55
(1,940)	1:7:A:VAL:H	1:32:A:VAL:HG11	5	0.55
(1,937)	1:22:A:LEU:HB3	1:22:A:LEU:H	20	0.55
(1,934)	1:26:A:ALA:H	1:25:A:LYS:HG3	14	0.55
(1,929)	1:13:A:TYR:HD1	1:11:A:LYS:HG2	24	0.55
(1,919)	1:23:A:ASN:HD21	1:25:A:LYS:HB3	16	0.55
(1,907)	1:11:A:LYS:HD2	1:11:A:LYS:HB2	1	0.55
(1,907)	1:11:A:LYS:HD3	1:11:A:LYS:HB3	8	0.55
(1,852)	1:13:A:TYR:HE2	1:24:A:VAL:H	10	0.55
(1,814)	1:26:A:ALA:H	1:25:A:LYS:HB2	1	0.55
(1,814)	1:26:A:ALA:H	1:25:A:LYS:HB2	6	0.55
(1,814)	1:26:A:ALA:H	1:25:A:LYS:HB2	9	0.55
(1,814)	1:26:A:ALA:H	1:25:A:LYS:HB2	14	0.55
(1,814)	1:26:A:ALA:H	1:25:A:LYS:HB2	15	0.55
(1,803)	1:30:A:ILE:HG22	1:32:A:VAL:H	26	0.55
(1,803)	1:30:A:ILE:HG21	1:32:A:VAL:H	28	0.55
(1,800)	1:44:A:CYS:HB2	1:55:A:VAL:HA	14	0.55
(1,800)	1:44:A:CYS:HB3	1:55:A:VAL:HA	28	0.55
(1,800)	1:44:A:CYS:HB2	1:55:A:VAL:HA	30	0.55
(1,795)	1:17:A:TYR:HB3	1:18:A:ASP:H	18	0.55
(1,685)	1:27:A:GLY:H	1:24:A:VAL:HG22	25	0.55
(1,659)	1:11:A:LYS:H	1:10:A:THR:H	7	0.55
(1,659)	1:11:A:LYS:H	1:10:A:THR:H	30	0.55
(1,514)	1:65:A:ALA:H	1:64:A:GLU:HB2	5	0.55
(1,507)	1:22:A:LEU:H	1:22:A:LEU:HD13	1	0.55
(1,496)	1:41:A:TRP:H	1:41:A:TRP:HB3	8	0.55
(1,496)	1:41:A:TRP:H	1:41:A:TRP:HB3	10	0.55

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,466)	1:30:A:ILE:H	1:29:A:ILE:HB	23	0.55
(1,451)	1:25:A:LYS:H	1:24:A:VAL:HB	12	0.55
(1,451)	1:25:A:LYS:H	1:24:A:VAL:HB	13	0.55
(1,451)	1:25:A:LYS:H	1:24:A:VAL:HB	28	0.55
(1,448)	1:25:A:LYS:H	1:25:A:LYS:HB2	3	0.55
(1,448)	1:25:A:LYS:H	1:25:A:LYS:HB2	14	0.55
(1,448)	1:25:A:LYS:H	1:25:A:LYS:HB2	29	0.55
(1,441)	1:30:A:ILE:HG21	1:45:A:ILE:HG21	13	0.55
(1,434)	1:24:A:VAL:HG11	1:30:A:ILE:HD13	16	0.55
(1,420)	1:7:A:VAL:HG12	1:32:A:VAL:HG11	6	0.55
(1,399)	1:29:A:ILE:H	1:29:A:ILE:HD12	18	0.55
(1,395)	1:42:A:LYS:H	1:33:A:LEU:HD12	3	0.55
(1,330)	1:63:A:GLY:H	1:62:A:LEU:HB2	16	0.55
(1,317)	1:39:A:GLY:H	1:38:A:ASP:HB3	5	0.55
(1,211)	1:25:A:LYS:H	1:11:A:LYS:H	27	0.55
(1,190)	1:34:A:GLU:HG2	1:3:A:HIS:H	19	0.55
(1,174)	1:30:A:ILE:HG12	1:9:A:ALA:H	30	0.55
(1,109)	1:32:A:VAL:H	1:32:A:VAL:HB	10	0.55
(1,109)	1:32:A:VAL:H	1:32:A:VAL:HB	28	0.55
(1,50)	1:47:A:ASP:H	1:47:A:ASP:HB2	2	0.55
(1,967)	1:62:A:LEU:H	1:63:A:GLY:HA2	3	0.54
(1,966)	1:8:A:ARG:H	1:6:A:GLN:HG2	18	0.54
(1,958)	1:61:A:SER:H	1:5:A:LEU:HD23	3	0.54
(1,956)	1:24:A:VAL:HG22	1:13:A:TYR:HD2	19	0.54
(1,952)	1:23:A:ASN:H	1:45:A:ILE:HD13	11	0.54
(1,937)	1:22:A:LEU:HB3	1:22:A:LEU:H	26	0.54
(1,934)	1:26:A:ALA:H	1:25:A:LYS:HG3	15	0.54
(1,925)	1:11:A:LYS:HG2	1:13:A:TYR:H	8	0.54
(1,908)	1:34:A:GLU:H	1:35:A:GLN:HG3	5	0.54
(1,906)	1:34:A:GLU:HG3	1:3:A:HIS:H	17	0.54
(1,906)	1:34:A:GLU:HG3	1:3:A:HIS:H	24	0.54
(1,900)	1:14:A:CYS:HB3	1:24:A:VAL:H	16	0.54
(1,887)	1:29:A:ILE:HG23	1:31:A:THR:HB	21	0.54
(1,863)	1:9:A:ALA:H	1:29:A:ILE:HA	13	0.54
(1,823)	1:28:A:ASP:H	1:9:A:ALA:HB1	30	0.54
(1,814)	1:26:A:ALA:H	1:25:A:LYS:HB2	5	0.54
(1,814)	1:26:A:ALA:H	1:25:A:LYS:HB2	16	0.54
(1,798)	1:24:A:VAL:HB	1:12:A:ASP:HA	3	0.54
(1,795)	1:17:A:TYR:HB3	1:18:A:ASP:H	5	0.54
(1,743)	1:63:A:GLY:H	1:62:A:LEU:HG	30	0.54
(1,707)	1:35:A:GLN:HE22	1:35:A:GLN:HB3	24	0.54
(1,687)	1:27:A:GLY:H	1:26:A:ALA:HB1	5	0.54

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,670)	1:43:A:GLY:H	1:42:A:LYS:HB2	29	0.54
(1,659)	1:11:A:LYS:H	1:10:A:THR:H	2	0.54
(1,659)	1:11:A:LYS:H	1:10:A:THR:H	13	0.54
(1,620)	1:21:A:SER:H	1:14:A:CYS:HB2	22	0.54
(1,544)	1:40:A:ARG:H	1:40:A:ARG:HB2	10	0.54
(1,544)	1:40:A:ARG:H	1:40:A:ARG:HB2	22	0.54
(1,541)	1:40:A:ARG:H	1:40:A:ARG:HD2	30	0.54
(1,507)	1:22:A:LEU:H	1:22:A:LEU:HD11	15	0.54
(1,507)	1:22:A:LEU:H	1:22:A:LEU:HD12	21	0.54
(1,496)	1:41:A:TRP:H	1:41:A:TRP:HB3	4	0.54
(1,467)	1:30:A:ILE:H	1:29:A:ILE:HG22	7	0.54
(1,466)	1:30:A:ILE:H	1:29:A:ILE:HB	21	0.54
(1,451)	1:25:A:LYS:H	1:24:A:VAL:HB	14	0.54
(1,448)	1:25:A:LYS:H	1:25:A:LYS:HB2	20	0.54
(1,448)	1:25:A:LYS:H	1:25:A:LYS:HB2	21	0.54
(1,448)	1:25:A:LYS:H	1:25:A:LYS:HB2	22	0.54
(1,431)	1:22:A:LEU:HD13	1:24:A:VAL:HG12	21	0.54
(1,420)	1:7:A:VAL:HG11	1:32:A:VAL:HG12	13	0.54
(1,420)	1:7:A:VAL:HG11	1:32:A:VAL:HG12	23	0.54
(1,420)	1:7:A:VAL:HG12	1:32:A:VAL:HG13	28	0.54
(1,416)	1:5:A:LEU:HD23	1:7:A:VAL:HG13	5	0.54
(1,212)	1:52:A:ASN:HD21	1:52:A:ASN:H	15	0.54
(1,211)	1:25:A:LYS:H	1:11:A:LYS:H	16	0.54
(1,190)	1:34:A:GLU:HG3	1:3:A:HIS:H	13	0.54
(1,174)	1:30:A:ILE:HG12	1:9:A:ALA:H	4	0.54
(1,109)	1:32:A:VAL:H	1:32:A:VAL:HB	26	0.54
(1,109)	1:32:A:VAL:H	1:32:A:VAL:HB	29	0.54
(1,66)	1:28:A:ASP:HB3	1:25:A:LYS:HG2	6	0.54
(1,62)	1:18:A:ASP:HB2	1:18:A:ASP:H	29	0.54
(1,954)	1:44:A:CYS:HA	1:45:A:ILE:HD11	11	0.53
(1,946)	1:7:A:VAL:HG22	1:30:A:ILE:H	15	0.53
(1,937)	1:22:A:LEU:HB3	1:22:A:LEU:H	10	0.53
(1,909)	1:41:A:TRP:HD1	1:40:A:ARG:HG2	18	0.53
(1,900)	1:24:A:VAL:H	1:14:A:CYS:HB2	17	0.53
(1,887)	1:29:A:ILE:HG23	1:31:A:THR:HB	15	0.53
(1,887)	1:29:A:ILE:HG22	1:31:A:THR:HB	23	0.53
(1,840)	1:11:A:LYS:HD3	1:12:A:ASP:H	29	0.53
(1,810)	1:23:A:ASN:H	1:45:A:ILE:H	9	0.53
(1,714)	1:15:A:ASN:HD22	1:15:A:ASN:HA	17	0.53
(1,685)	1:27:A:GLY:H	1:24:A:VAL:HG22	3	0.53
(1,670)	1:43:A:GLY:H	1:42:A:LYS:HB2	10	0.53
(1,659)	1:11:A:LYS:H	1:10:A:THR:H	8	0.53

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,659)	1:11:A:LYS:H	1:10:A:THR:H	11	0.53
(1,659)	1:11:A:LYS:H	1:10:A:THR:H	15	0.53
(1,659)	1:11:A:LYS:H	1:10:A:THR:H	17	0.53
(1,659)	1:11:A:LYS:H	1:10:A:THR:H	23	0.53
(1,650)	1:12:A:ASP:H	1:11:A:LYS:HB2	26	0.53
(1,496)	1:41:A:TRP:H	1:41:A:TRP:HB3	13	0.53
(1,485)	1:46:A:HIS:H	1:45:A:ILE:HD11	1	0.53
(1,466)	1:30:A:ILE:H	1:29:A:ILE:HB	27	0.53
(1,466)	1:30:A:ILE:H	1:29:A:ILE:HB	28	0.53
(1,448)	1:25:A:LYS:H	1:25:A:LYS:HB2	5	0.53
(1,441)	1:30:A:ILE:HG22	1:45:A:ILE:HG23	27	0.53
(1,437)	1:24:A:VAL:HG21	1:30:A:ILE:HG23	26	0.53
(1,431)	1:22:A:LEU:HD11	1:24:A:VAL:HG11	27	0.53
(1,420)	1:7:A:VAL:HG11	1:32:A:VAL:HG13	29	0.53
(1,403)	1:58:A:PHE:H	1:22:A:LEU:HD12	2	0.53
(1,403)	1:58:A:PHE:H	1:22:A:LEU:HD12	7	0.53
(1,399)	1:29:A:ILE:H	1:29:A:ILE:HD12	12	0.53
(1,228)	1:51:A:GLY:H	1:52:A:ASN:HD22	23	0.53
(1,170)	1:32:A:VAL:HG23	1:32:A:VAL:H	18	0.53
(1,109)	1:32:A:VAL:H	1:32:A:VAL:HB	2	0.53
(1,109)	1:32:A:VAL:H	1:32:A:VAL:HB	5	0.53
(1,109)	1:32:A:VAL:H	1:32:A:VAL:HB	12	0.53
(1,109)	1:32:A:VAL:H	1:32:A:VAL:HB	14	0.53
(1,109)	1:32:A:VAL:H	1:32:A:VAL:HB	21	0.53
(1,62)	1:18:A:ASP:HB2	1:18:A:ASP:H	24	0.53
(1,61)	1:18:A:ASP:HB2	1:18:A:ASP:HA	4	0.53
(1,61)	1:18:A:ASP:HB2	1:18:A:ASP:HA	17	0.53
(1,61)	1:18:A:ASP:HB2	1:18:A:ASP:HA	19	0.53
(1,61)	1:18:A:ASP:HB2	1:18:A:ASP:HA	22	0.53
(1,27)	1:43:A:GLY:HA3	1:33:A:LEU:HD11	5	0.53
(1,27)	1:43:A:GLY:HA3	1:33:A:LEU:HD11	19	0.53
(1,968)	1:10:A:THR:H	1:26:A:ALA:HA	14	0.52
(1,968)	1:10:A:THR:H	1:63:A:GLY:HA3	25	0.52
(1,956)	1:24:A:VAL:HG23	1:13:A:TYR:HD2	5	0.52
(1,940)	1:7:A:VAL:H	1:32:A:VAL:HG13	23	0.52
(1,937)	1:22:A:LEU:HB3	1:22:A:LEU:H	23	0.52
(1,929)	1:13:A:TYR:HD2	1:11:A:LYS:HG2	25	0.52
(1,925)	1:11:A:LYS:HG2	1:13:A:TYR:H	17	0.52
(1,917)	1:54:A:ARG:HB2	1:45:A:ILE:H	25	0.52
(1,912)	1:41:A:TRP:H	1:40:A:ARG:HG3	7	0.52
(1,900)	1:14:A:CYS:HB3	1:24:A:VAL:H	2	0.52
(1,894)	1:53:A:ASP:H	1:54:A:ARG:HD2	7	0.52

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,887)	1:29:A:ILE:HG23	1:31:A:THR:HB	26	0.52
(1,814)	1:26:A:ALA:H	1:25:A:LYS:HB2	17	0.52
(1,814)	1:26:A:ALA:H	1:25:A:LYS:HB2	22	0.52
(1,810)	1:23:A:ASN:H	1:45:A:ILE:H	26	0.52
(1,800)	1:44:A:CYS:HB2	1:55:A:VAL:HA	10	0.52
(1,735)	1:39:A:GLY:H	1:41:A:TRP:HD1	6	0.52
(1,659)	1:11:A:LYS:H	1:10:A:THR:H	20	0.52
(1,605)	1:57:A:TYR:H	1:21:A:SER:HA	8	0.52
(1,584)	1:3:A:HIS:H	1:34:A:GLU:HB3	3	0.52
(1,507)	1:22:A:LEU:H	1:22:A:LEU:HD13	7	0.52
(1,507)	1:22:A:LEU:H	1:22:A:LEU:HD11	22	0.52
(1,496)	1:41:A:TRP:H	1:41:A:TRP:HB3	15	0.52
(1,493)	1:41:A:TRP:HE1	1:40:A:ARG:H	10	0.52
(1,480)	1:45:A:ILE:H	1:44:A:CYS:HB2	4	0.52
(1,467)	1:30:A:ILE:H	1:29:A:ILE:HG23	3	0.52
(1,464)	1:67:A:VAL:H	1:66:A:ILE:HG12	12	0.52
(1,445)	1:30:A:ILE:HD12	1:45:A:ILE:HG22	20	0.52
(1,444)	1:30:A:ILE:HD11	1:32:A:VAL:HG13	3	0.52
(1,437)	1:24:A:VAL:HG22	1:30:A:ILE:HG23	2	0.52
(1,437)	1:24:A:VAL:HG23	1:30:A:ILE:HG23	13	0.52
(1,434)	1:24:A:VAL:HG11	1:30:A:ILE:HD13	12	0.52
(1,403)	1:58:A:PHE:H	1:22:A:LEU:HD11	5	0.52
(1,403)	1:58:A:PHE:H	1:22:A:LEU:HD13	11	0.52
(1,403)	1:58:A:PHE:H	1:22:A:LEU:HD12	29	0.52
(1,399)	1:29:A:ILE:H	1:29:A:ILE:HD11	15	0.52
(1,317)	1:39:A:GLY:H	1:38:A:ASP:HB3	11	0.52
(1,182)	1:25:A:LYS:H	1:27:A:GLY:H	18	0.52
(1,109)	1:32:A:VAL:H	1:32:A:VAL:HB	6	0.52
(1,109)	1:32:A:VAL:H	1:32:A:VAL:HB	7	0.52
(1,109)	1:32:A:VAL:H	1:32:A:VAL:HB	8	0.52
(1,109)	1:32:A:VAL:H	1:32:A:VAL:HB	9	0.52
(1,109)	1:32:A:VAL:H	1:32:A:VAL:HB	25	0.52
(1,61)	1:18:A:ASP:HB2	1:18:A:ASP:HA	6	0.52
(1,61)	1:18:A:ASP:HB2	1:18:A:ASP:HA	8	0.52
(1,61)	1:18:A:ASP:HB2	1:18:A:ASP:HA	11	0.52
(1,61)	1:18:A:ASP:HB2	1:18:A:ASP:HA	12	0.52
(1,61)	1:18:A:ASP:HB2	1:18:A:ASP:HA	25	0.52
(1,61)	1:18:A:ASP:HB2	1:18:A:ASP:HA	26	0.52
(1,61)	1:18:A:ASP:HB2	1:18:A:ASP:HA	30	0.52
(1,46)	1:13:A:TYR:HB2	1:11:A:LYS:HD2	8	0.52
(1,964)	1:45:A:ILE:H	1:55:A:VAL:HA	2	0.51
(1,946)	1:7:A:VAL:HG23	1:30:A:ILE:H	2	0.51

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,940)	1:44:A:CYS:H	1:32:A:VAL:HG12	26	0.51
(1,939)	1:58:A:PHE:HD1	1:32:A:VAL:HG13	19	0.51
(1,931)	1:6:A:GLN:HE21	1:8:A:ARG:HG2	26	0.51
(1,923)	1:33:A:LEU:HD11	1:33:A:LEU:HB3	2	0.51
(1,923)	1:33:A:LEU:HD11	1:33:A:LEU:HB3	4	0.51
(1,917)	1:25:A:LYS:HB3	1:25:A:LYS:H	23	0.51
(1,908)	1:52:A:ASN:H	1:49:A:ARG:HB3	28	0.51
(1,899)	1:27:A:GLY:H	1:28:A:ASP:HB2	30	0.51
(1,896)	1:48:A:ASN:H	1:47:A:ASP:HB2	12	0.51
(1,892)	1:41:A:TRP:HH2	1:61:A:SER:HA	24	0.51
(1,863)	1:9:A:ALA:H	1:29:A:ILE:HA	29	0.51
(1,835)	1:5:A:LEU:H	1:4:A:MET:H	9	0.51
(1,822)	1:26:A:ALA:HA	1:28:A:ASP:H	13	0.51
(1,814)	1:26:A:ALA:H	1:25:A:LYS:HB2	12	0.51
(1,814)	1:26:A:ALA:H	1:25:A:LYS:HB2	18	0.51
(1,814)	1:26:A:ALA:H	1:25:A:LYS:HB2	19	0.51
(1,810)	1:23:A:ASN:H	1:45:A:ILE:H	4	0.51
(1,803)	1:30:A:ILE:HG23	1:32:A:VAL:H	22	0.51
(1,775)	1:38:A:ASP:H	1:40:A:ARG:H	7	0.51
(1,735)	1:39:A:GLY:H	1:41:A:TRP:HD1	24	0.51
(1,722)	1:51:A:GLY:H	1:49:A:ARG:HB2	2	0.51
(1,707)	1:35:A:GLN:HE22	1:35:A:GLN:HB3	16	0.51
(1,659)	1:11:A:LYS:H	1:10:A:THR:H	3	0.51
(1,659)	1:11:A:LYS:H	1:10:A:THR:H	5	0.51
(1,659)	1:11:A:LYS:H	1:10:A:THR:H	29	0.51
(1,654)	1:17:A:TYR:H	1:17:A:TYR:HB3	22	0.51
(1,628)	1:38:A:ASP:H	1:36:A:HIS:HB2	8	0.51
(1,617)	1:21:A:SER:H	1:21:A:SER:HB3	24	0.51
(1,544)	1:40:A:ARG:H	1:40:A:ARG:HB2	1	0.51
(1,541)	1:40:A:ARG:H	1:40:A:ARG:HD2	13	0.51
(1,507)	1:22:A:LEU:H	1:22:A:LEU:HD11	12	0.51
(1,496)	1:41:A:TRP:H	1:41:A:TRP:HB3	7	0.51
(1,467)	1:30:A:ILE:H	1:29:A:ILE:HG22	16	0.51
(1,467)	1:30:A:ILE:H	1:29:A:ILE:HG23	28	0.51
(1,466)	1:30:A:ILE:H	1:29:A:ILE:HB	9	0.51
(1,448)	1:25:A:LYS:H	1:25:A:LYS:HB2	16	0.51
(1,448)	1:25:A:LYS:H	1:25:A:LYS:HB2	27	0.51
(1,446)	1:33:A:LEU:HD21	1:55:A:VAL:HG22	26	0.51
(1,420)	1:7:A:VAL:HG13	1:32:A:VAL:HG11	2	0.51
(1,303)	1:25:A:LYS:HE2	1:48:A:ASN:HD22	1	0.51
(1,225)	1:35:A:GLN:HG2	1:41:A:TRP:HD1	24	0.51
(1,211)	1:25:A:LYS:H	1:11:A:LYS:H	24	0.51

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,192)	1:67:A:VAL:H	1:6:A:GLN:H	16	0.51
(1,174)	1:30:A:ILE:HG12	1:9:A:ALA:H	10	0.51
(1,174)	1:30:A:ILE:HG12	1:9:A:ALA:H	11	0.51
(1,109)	1:32:A:VAL:H	1:32:A:VAL:HB	3	0.51
(1,109)	1:32:A:VAL:H	1:32:A:VAL:HB	4	0.51
(1,109)	1:32:A:VAL:H	1:32:A:VAL:HB	16	0.51
(1,109)	1:32:A:VAL:H	1:32:A:VAL:HB	22	0.51
(1,109)	1:32:A:VAL:H	1:32:A:VAL:HB	24	0.51
(1,61)	1:18:A:ASP:HB2	1:18:A:ASP:HA	21	0.51
(1,50)	1:47:A:ASP:H	1:47:A:ASP:HB2	6	0.51
(1,50)	1:47:A:ASP:H	1:47:A:ASP:HB2	9	0.51
(1,50)	1:47:A:ASP:H	1:47:A:ASP:HB2	17	0.51
(1,14)	1:27:A:GLY:H	1:26:A:ALA:HA	30	0.51
(1,964)	1:45:A:ILE:H	1:55:A:VAL:HA	24	0.5
(1,954)	1:44:A:CYS:HA	1:45:A:ILE:HD11	1	0.5
(1,952)	1:23:A:ASN:H	1:45:A:ILE:HD11	20	0.5
(1,944)	1:30:A:ILE:HD12	1:7:A:VAL:HG23	4	0.5
(1,940)	1:44:A:CYS:H	1:32:A:VAL:HG11	15	0.5
(1,929)	1:13:A:TYR:HD1	1:11:A:LYS:HG3	10	0.5
(1,924)	1:65:A:ALA:HB3	1:5:A:LEU:HD12	5	0.5
(1,923)	1:33:A:LEU:HD11	1:33:A:LEU:HB3	16	0.5
(1,887)	1:29:A:ILE:HG23	1:31:A:THR:HB	18	0.5
(1,878)	1:52:A:ASN:H	1:50:A:THR:HA	20	0.5
(1,852)	1:13:A:TYR:HE2	1:24:A:VAL:H	5	0.5
(1,814)	1:26:A:ALA:H	1:25:A:LYS:HB2	23	0.5
(1,802)	1:26:A:ALA:HB3	1:12:A:ASP:HA	22	0.5
(1,800)	1:44:A:CYS:HB2	1:55:A:VAL:HA	9	0.5
(1,670)	1:43:A:GLY:H	1:42:A:LYS:HB2	28	0.5
(1,659)	1:11:A:LYS:H	1:10:A:THR:H	22	0.5
(1,659)	1:11:A:LYS:H	1:10:A:THR:H	25	0.5
(1,650)	1:12:A:ASP:H	1:11:A:LYS:HB2	4	0.5
(1,635)	1:52:A:ASN:H	1:52:A:ASN:HB2	22	0.5
(1,627)	1:38:A:ASP:H	1:37:A:PRO:HB3	22	0.5
(1,587)	1:53:A:ASP:H	1:53:A:ASP:HB2	16	0.5
(1,571)	1:66:A:ILE:H	1:65:A:ALA:HB3	21	0.5
(1,544)	1:40:A:ARG:H	1:40:A:ARG:HB2	14	0.5
(1,544)	1:40:A:ARG:H	1:40:A:ARG:HB2	15	0.5
(1,544)	1:40:A:ARG:H	1:40:A:ARG:HB2	16	0.5
(1,544)	1:40:A:ARG:H	1:40:A:ARG:HB2	23	0.5
(1,515)	1:14:A:CYS:H	1:13:A:TYR:HB3	17	0.5
(1,507)	1:22:A:LEU:H	1:22:A:LEU:HD11	9	0.5
(1,496)	1:41:A:TRP:H	1:41:A:TRP:HB3	3	0.5

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,496)	1:41:A:TRP:H	1:41:A:TRP:HB3	14	0.5
(1,480)	1:45:A:ILE:H	1:44:A:CYS:HB2	28	0.5
(1,466)	1:30:A:ILE:H	1:29:A:ILE:HB	16	0.5
(1,448)	1:25:A:LYS:H	1:25:A:LYS:HB2	6	0.5
(1,448)	1:25:A:LYS:H	1:25:A:LYS:HB2	25	0.5
(1,448)	1:25:A:LYS:H	1:25:A:LYS:HB2	30	0.5
(1,444)	1:30:A:ILE:HD12	1:32:A:VAL:HG12	14	0.5
(1,431)	1:22:A:LEU:HD13	1:24:A:VAL:HG12	17	0.5
(1,423)	1:7:A:VAL:HG23	1:32:A:VAL:HG23	7	0.5
(1,399)	1:29:A:ILE:H	1:29:A:ILE:HD13	3	0.5
(1,399)	1:29:A:ILE:H	1:29:A:ILE:HD12	29	0.5
(1,226)	1:39:A:GLY:HA3	1:41:A:TRP:HD1	20	0.5
(1,225)	1:35:A:GLN:HG2	1:41:A:TRP:HD1	16	0.5
(1,220)	1:15:A:ASN:H	1:15:A:ASN:HD22	21	0.5
(1,176)	1:33:A:LEU:H	1:42:A:LYS:H	15	0.5
(1,168)	1:13:A:TYR:HD2	1:22:A:LEU:H	24	0.5
(1,109)	1:32:A:VAL:H	1:32:A:VAL:HB	13	0.5
(1,109)	1:32:A:VAL:H	1:32:A:VAL:HB	15	0.5
(1,109)	1:32:A:VAL:H	1:32:A:VAL:HB	30	0.5
(1,62)	1:18:A:ASP:HB2	1:18:A:ASP:H	18	0.5
(1,46)	1:13:A:TYR:HB2	1:11:A:LYS:HD2	2	0.5
(1,46)	1:13:A:TYR:HB2	1:11:A:LYS:HD2	4	0.5
(1,968)	1:10:A:THR:H	1:63:A:GLY:HA3	1	0.49
(1,964)	1:45:A:ILE:H	1:55:A:VAL:HA	14	0.49
(1,926)	1:28:A:ASP:H	1:25:A:LYS:HG3	10	0.49
(1,923)	1:33:A:LEU:HD12	1:33:A:LEU:HB3	5	0.49
(1,923)	1:33:A:LEU:HD13	1:33:A:LEU:HB3	10	0.49
(1,870)	1:29:A:ILE:HG22	1:45:A:ILE:HA	6	0.49
(1,840)	1:11:A:LYS:HD3	1:12:A:ASP:H	22	0.49
(1,834)	1:51:A:GLY:HA3	1:47:A:ASP:H	4	0.49
(1,816)	1:5:A:LEU:HB2	1:7:A:VAL:H	5	0.49
(1,816)	1:57:A:TYR:H	1:22:A:LEU:HB2	30	0.49
(1,814)	1:26:A:ALA:H	1:25:A:LYS:HB2	29	0.49
(1,805)	1:30:A:ILE:H	1:30:A:ILE:HD11	2	0.49
(1,798)	1:24:A:VAL:HB	1:12:A:ASP:HA	26	0.49
(1,795)	1:17:A:TYR:HB3	1:18:A:ASP:H	14	0.49
(1,743)	1:63:A:GLY:H	1:62:A:LEU:HG	12	0.49
(1,743)	1:63:A:GLY:H	1:62:A:LEU:HG	18	0.49
(1,659)	1:11:A:LYS:H	1:10:A:THR:H	1	0.49
(1,659)	1:11:A:LYS:H	1:10:A:THR:H	6	0.49
(1,627)	1:38:A:ASP:H	1:37:A:PRO:HB3	16	0.49
(1,627)	1:38:A:ASP:H	1:37:A:PRO:HB3	20	0.49

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,544)	1:40:A:ARG:H	1:40:A:ARG:HB2	18	0.49
(1,515)	1:14:A:CYS:H	1:13:A:TYR:HB3	30	0.49
(1,496)	1:41:A:TRP:H	1:41:A:TRP:HB3	22	0.49
(1,496)	1:41:A:TRP:H	1:41:A:TRP:HB3	27	0.49
(1,480)	1:45:A:ILE:H	1:44:A:CYS:HB2	20	0.49
(1,480)	1:45:A:ILE:H	1:44:A:CYS:HB2	27	0.49
(1,444)	1:30:A:ILE:HD11	1:32:A:VAL:HG13	5	0.49
(1,441)	1:30:A:ILE:HG21	1:45:A:ILE:HG22	5	0.49
(1,437)	1:24:A:VAL:HG21	1:30:A:ILE:HG23	23	0.49
(1,434)	1:24:A:VAL:HG13	1:30:A:ILE:HD13	19	0.49
(1,399)	1:29:A:ILE:H	1:29:A:ILE:HD12	17	0.49
(1,399)	1:29:A:ILE:H	1:29:A:ILE:HD11	25	0.49
(1,395)	1:42:A:LYS:H	1:33:A:LEU:HD12	13	0.49
(1,373)	1:41:A:TRP:HE1	1:35:A:GLN:HB3	10	0.49
(1,318)	1:44:A:CYS:HB3	1:45:A:ILE:H	18	0.49
(1,249)	1:13:A:TYR:HE2	1:21:A:SER:HA	8	0.49
(1,200)	1:13:A:TYR:HD2	1:14:A:CYS:H	30	0.49
(1,191)	1:6:A:GLN:H	1:6:A:GLN:HG2	27	0.49
(1,182)	1:25:A:LYS:H	1:27:A:GLY:H	13	0.49
(1,174)	1:30:A:ILE:HG12	1:9:A:ALA:H	6	0.49
(1,174)	1:30:A:ILE:HG12	1:9:A:ALA:H	27	0.49
(1,109)	1:32:A:VAL:H	1:32:A:VAL:HB	19	0.49
(1,109)	1:32:A:VAL:H	1:32:A:VAL:HB	23	0.49
(1,61)	1:18:A:ASP:HB2	1:18:A:ASP:HA	7	0.49
(1,965)	1:26:A:ALA:H	1:12:A:ASP:HA	19	0.48
(1,946)	1:7:A:VAL:HG22	1:30:A:ILE:H	3	0.48
(1,940)	1:7:A:VAL:H	1:32:A:VAL:HG11	3	0.48
(1,923)	1:33:A:LEU:HD13	1:33:A:LEU:HB3	1	0.48
(1,923)	1:33:A:LEU:HD13	1:33:A:LEU:HB3	3	0.48
(1,923)	1:33:A:LEU:HD12	1:33:A:LEU:HB3	19	0.48
(1,923)	1:33:A:LEU:HD12	1:33:A:LEU:HB3	23	0.48
(1,923)	1:33:A:LEU:HD11	1:33:A:LEU:HB3	24	0.48
(1,923)	1:33:A:LEU:HD12	1:33:A:LEU:HB3	30	0.48
(1,917)	1:25:A:LYS:HB3	1:25:A:LYS:H	2	0.48
(1,917)	1:25:A:LYS:HB3	1:25:A:LYS:H	26	0.48
(1,909)	1:41:A:TRP:HD1	1:40:A:ARG:HG2	3	0.48
(1,908)	1:52:A:ASN:H	1:49:A:ARG:HB2	9	0.48
(1,908)	1:34:A:GLU:H	1:35:A:GLN:HG2	25	0.48
(1,899)	1:27:A:GLY:H	1:28:A:ASP:HB2	28	0.48
(1,894)	1:23:A:ASN:H	1:54:A:ARG:HD2	18	0.48
(1,892)	1:13:A:TYR:HD1	1:24:A:VAL:HA	22	0.48
(1,878)	1:52:A:ASN:H	1:50:A:THR:HB	27	0.48

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,852)	1:13:A:TYR:HE2	1:24:A:VAL:H	8	0.48
(1,820)	1:15:A:ASN:H	1:15:A:ASN:HB2	2	0.48
(1,814)	1:26:A:ALA:H	1:25:A:LYS:HB2	3	0.48
(1,814)	1:26:A:ALA:H	1:25:A:LYS:HB2	21	0.48
(1,814)	1:26:A:ALA:H	1:25:A:LYS:HB2	24	0.48
(1,795)	1:17:A:TYR:HB3	1:18:A:ASP:H	21	0.48
(1,795)	1:17:A:TYR:HB3	1:18:A:ASP:H	26	0.48
(1,659)	1:11:A:LYS:H	1:10:A:THR:H	10	0.48
(1,659)	1:11:A:LYS:H	1:10:A:THR:H	14	0.48
(1,571)	1:66:A:ILE:H	1:65:A:ALA:HB1	17	0.48
(1,554)	1:54:A:ARG:H	1:54:A:ARG:HB2	3	0.48
(1,543)	1:40:A:ARG:H	1:40:A:ARG:HG2	29	0.48
(1,496)	1:41:A:TRP:H	1:41:A:TRP:HB3	29	0.48
(1,451)	1:25:A:LYS:H	1:24:A:VAL:HB	1	0.48
(1,451)	1:25:A:LYS:H	1:24:A:VAL:HB	9	0.48
(1,448)	1:25:A:LYS:H	1:25:A:LYS:HB2	7	0.48
(1,448)	1:25:A:LYS:H	1:25:A:LYS:HB2	15	0.48
(1,445)	1:30:A:ILE:HD12	1:45:A:ILE:HG21	2	0.48
(1,441)	1:30:A:ILE:HG21	1:45:A:ILE:HG21	4	0.48
(1,403)	1:58:A:PHE:H	1:22:A:LEU:HD13	12	0.48
(1,403)	1:58:A:PHE:H	1:22:A:LEU:HD11	16	0.48
(1,403)	1:58:A:PHE:H	1:22:A:LEU:HD12	26	0.48
(1,399)	1:29:A:ILE:H	1:29:A:ILE:HD12	28	0.48
(1,395)	1:42:A:LYS:H	1:33:A:LEU:HD12	12	0.48
(1,389)	1:11:A:LYS:HB3	1:24:A:VAL:HG12	25	0.48
(1,330)	1:63:A:GLY:H	1:62:A:LEU:HB2	18	0.48
(1,220)	1:15:A:ASN:H	1:15:A:ASN:HD22	4	0.48
(1,203)	1:52:A:ASN:HD22	1:52:A:ASN:H	8	0.48
(1,174)	1:30:A:ILE:HG12	1:9:A:ALA:H	12	0.48
(1,174)	1:30:A:ILE:HG12	1:9:A:ALA:H	23	0.48
(1,65)	1:28:A:ASP:HB3	1:25:A:LYS:H	20	0.48
(1,60)	1:28:A:ASP:HB2	1:28:A:ASP:HA	4	0.48
(1,27)	1:43:A:GLY:HA3	1:33:A:LEU:HD13	24	0.48
(1,967)	1:62:A:LEU:H	1:63:A:GLY:HA2	28	0.47
(1,953)	1:23:A:ASN:HB3	1:45:A:ILE:HG22	8	0.47
(1,946)	1:7:A:VAL:HG23	1:30:A:ILE:H	13	0.47
(1,940)	1:7:A:VAL:H	1:32:A:VAL:HG12	22	0.47
(1,925)	1:11:A:LYS:HG2	1:13:A:TYR:H	11	0.47
(1,923)	1:33:A:LEU:HD12	1:33:A:LEU:HB3	9	0.47
(1,923)	1:33:A:LEU:HD13	1:33:A:LEU:HB3	11	0.47
(1,923)	1:33:A:LEU:HD11	1:33:A:LEU:HB3	13	0.47
(1,923)	1:33:A:LEU:HD11	1:33:A:LEU:HB3	15	0.47

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,923)	1:33:A:LEU:HD12	1:33:A:LEU:HB3	21	0.47
(1,923)	1:33:A:LEU:HD12	1:33:A:LEU:HB3	25	0.47
(1,919)	1:23:A:ASN:HD21	1:54:A:ARG:HB2	27	0.47
(1,906)	1:34:A:GLU:HG2	1:3:A:HIS:H	20	0.47
(1,863)	1:9:A:ALA:H	1:29:A:ILE:HA	4	0.47
(1,814)	1:26:A:ALA:H	1:25:A:LYS:HB2	25	0.47
(1,810)	1:23:A:ASN:H	1:45:A:ILE:H	11	0.47
(1,800)	1:44:A:CYS:HB2	1:55:A:VAL:HA	18	0.47
(1,798)	1:24:A:VAL:HB	1:12:A:ASP:HA	27	0.47
(1,795)	1:17:A:TYR:HB3	1:18:A:ASP:H	16	0.47
(1,687)	1:27:A:GLY:H	1:26:A:ALA:HB1	13	0.47
(1,664)	1:61:A:SER:H	1:61:A:SER:HB3	22	0.47
(1,627)	1:38:A:ASP:H	1:37:A:PRO:HB3	9	0.47
(1,627)	1:38:A:ASP:H	1:37:A:PRO:HB3	21	0.47
(1,507)	1:22:A:LEU:H	1:22:A:LEU:HD12	20	0.47
(1,496)	1:41:A:TRP:H	1:41:A:TRP:HB3	24	0.47
(1,496)	1:41:A:TRP:H	1:41:A:TRP:HB3	28	0.47
(1,480)	1:45:A:ILE:H	1:44:A:CYS:HB2	8	0.47
(1,480)	1:45:A:ILE:H	1:44:A:CYS:HB2	25	0.47
(1,466)	1:30:A:ILE:H	1:29:A:ILE:HB	13	0.47
(1,466)	1:30:A:ILE:H	1:29:A:ILE:HB	30	0.47
(1,451)	1:25:A:LYS:H	1:24:A:VAL:HB	8	0.47
(1,451)	1:25:A:LYS:H	1:24:A:VAL:HB	23	0.47
(1,431)	1:22:A:LEU:HD13	1:24:A:VAL:HG11	16	0.47
(1,403)	1:58:A:PHE:H	1:22:A:LEU:HD12	1	0.47
(1,394)	1:41:A:TRP:HZ3	1:5:A:LEU:HD23	3	0.47
(1,382)	1:65:A:ALA:H	1:5:A:LEU:HB3	25	0.47
(1,360)	1:27:A:GLY:H	1:25:A:LYS:HG3	22	0.47
(1,358)	1:41:A:TRP:H	1:40:A:ARG:HB3	4	0.47
(1,309)	1:40:A:ARG:HB3	1:40:A:ARG:HD2	6	0.47
(1,203)	1:52:A:ASN:HD22	1:52:A:ASN:H	3	0.47
(1,203)	1:52:A:ASN:HD22	1:52:A:ASN:H	23	0.47
(1,191)	1:6:A:GLN:H	1:6:A:GLN:HG2	4	0.47
(1,191)	1:6:A:GLN:H	1:6:A:GLN:HG2	11	0.47
(1,56)	1:28:A:ASP:H	1:28:A:ASP:HB3	19	0.47
(1,45)	1:47:A:ASP:HB3	1:52:A:ASN:HA	3	0.47
(1,29)	1:43:A:GLY:HA3	1:30:A:ILE:HG21	1	0.47
(1,29)	1:43:A:GLY:HA3	1:30:A:ILE:HG21	10	0.47
(1,968)	1:10:A:THR:H	1:26:A:ALA:HA	20	0.46
(1,964)	1:45:A:ILE:H	1:55:A:VAL:HA	3	0.46
(1,953)	1:23:A:ASN:HB2	1:45:A:ILE:HG23	11	0.46
(1,952)	1:23:A:ASN:H	1:45:A:ILE:HD12	3	0.46

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,944)	1:30:A:ILE:HD12	1:7:A:VAL:HG22	7	0.46
(1,939)	1:58:A:PHE:HD1	1:32:A:VAL:HG12	5	0.46
(1,934)	1:26:A:ALA:H	1:25:A:LYS:HG2	2	0.46
(1,923)	1:33:A:LEU:HD13	1:33:A:LEU:HB3	12	0.46
(1,923)	1:33:A:LEU:HD11	1:33:A:LEU:HB3	14	0.46
(1,923)	1:33:A:LEU:HD13	1:33:A:LEU:HB3	17	0.46
(1,923)	1:33:A:LEU:HD11	1:33:A:LEU:HB3	18	0.46
(1,923)	1:33:A:LEU:HD13	1:33:A:LEU:HB3	22	0.46
(1,917)	1:25:A:LYS:HB3	1:25:A:LYS:H	3	0.46
(1,912)	1:41:A:TRP:H	1:40:A:ARG:HG3	11	0.46
(1,909)	1:41:A:TRP:HD1	1:40:A:ARG:HG2	15	0.46
(1,878)	1:52:A:ASN:H	1:50:A:THR:HB	21	0.46
(1,859)	1:67:A:VAL:H	1:7:A:VAL:HA	8	0.46
(1,814)	1:26:A:ALA:H	1:25:A:LYS:HB2	28	0.46
(1,810)	1:23:A:ASN:H	1:45:A:ILE:H	3	0.46
(1,800)	1:44:A:CYS:HB3	1:55:A:VAL:HA	16	0.46
(1,795)	1:17:A:TYR:HB3	1:18:A:ASP:H	2	0.46
(1,734)	1:39:A:GLY:H	1:40:A:ARG:HB2	13	0.46
(1,707)	1:35:A:GLN:HE22	1:35:A:GLN:HB3	17	0.46
(1,670)	1:43:A:GLY:H	1:42:A:LYS:HB2	11	0.46
(1,659)	1:11:A:LYS:H	1:10:A:THR:H	19	0.46
(1,627)	1:38:A:ASP:H	1:37:A:PRO:HB3	10	0.46
(1,627)	1:38:A:ASP:H	1:37:A:PRO:HB3	23	0.46
(1,587)	1:53:A:ASP:H	1:53:A:ASP:HB2	24	0.46
(1,560)	1:64:A:GLU:H	1:64:A:GLU:HG2	6	0.46
(1,554)	1:54:A:ARG:H	1:54:A:ARG:HB2	4	0.46
(1,554)	1:54:A:ARG:H	1:54:A:ARG:HB2	9	0.46
(1,496)	1:41:A:TRP:H	1:41:A:TRP:HB3	26	0.46
(1,437)	1:24:A:VAL:HG22	1:30:A:ILE:HG22	22	0.46
(1,434)	1:24:A:VAL:HG11	1:30:A:ILE:HD11	21	0.46
(1,432)	1:22:A:LEU:HD11	1:24:A:VAL:HG21	1	0.46
(1,427)	1:20:A:THR:HG21	1:55:A:VAL:HG23	2	0.46
(1,403)	1:58:A:PHE:H	1:22:A:LEU:HD11	23	0.46
(1,399)	1:29:A:ILE:H	1:29:A:ILE:HD12	1	0.46
(1,399)	1:29:A:ILE:H	1:29:A:ILE:HD12	21	0.46
(1,360)	1:27:A:GLY:H	1:25:A:LYS:HG3	17	0.46
(1,272)	1:17:A:TYR:HD2	1:16:A:ASN:HA	28	0.46
(1,249)	1:13:A:TYR:HE2	1:21:A:SER:HA	30	0.46
(1,191)	1:6:A:GLN:H	1:6:A:GLN:HG2	21	0.46
(1,176)	1:33:A:LEU:H	1:42:A:LYS:H	1	0.46
(1,107)	1:41:A:TRP:HB2	1:41:A:TRP:HD1	19	0.46
(1,107)	1:41:A:TRP:HB2	1:41:A:TRP:HD1	21	0.46

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,55)	1:47:A:ASP:HB3	1:52:A:ASN:H	8	0.46
(1,14)	1:27:A:GLY:H	1:26:A:ALA:HA	14	0.46
(1,14)	1:27:A:GLY:H	1:26:A:ALA:HA	27	0.46
(1,4)	1:32:A:VAL:HA	1:32:A:VAL:HG13	18	0.46
(1,964)	1:45:A:ILE:H	1:55:A:VAL:HA	30	0.45
(1,956)	1:24:A:VAL:HG23	1:13:A:TYR:HD2	10	0.45
(1,946)	1:7:A:VAL:HG21	1:30:A:ILE:H	17	0.45
(1,944)	1:30:A:ILE:HD11	1:7:A:VAL:HG23	10	0.45
(1,923)	1:33:A:LEU:HD13	1:33:A:LEU:HB3	29	0.45
(1,909)	1:41:A:TRP:HD1	1:40:A:ARG:HG2	19	0.45
(1,878)	1:52:A:ASN:H	1:50:A:THR:HB	1	0.45
(1,870)	1:29:A:ILE:HG22	1:45:A:ILE:HA	1	0.45
(1,858)	1:13:A:TYR:HD2	1:23:A:ASN:HA	15	0.45
(1,852)	1:13:A:TYR:HE2	1:24:A:VAL:H	28	0.45
(1,803)	1:30:A:ILE:HG22	1:32:A:VAL:H	12	0.45
(1,798)	1:24:A:VAL:HB	1:12:A:ASP:HA	15	0.45
(1,734)	1:39:A:GLY:H	1:40:A:ARG:HB2	24	0.45
(1,669)	1:43:A:GLY:H	1:55:A:VAL:HG23	30	0.45
(1,664)	1:61:A:SER:H	1:61:A:SER:HB3	29	0.45
(1,627)	1:38:A:ASP:H	1:37:A:PRO:HB3	2	0.45
(1,627)	1:38:A:ASP:H	1:37:A:PRO:HB3	13	0.45
(1,583)	1:55:A:VAL:H	1:55:A:VAL:HB	25	0.45
(1,525)	1:15:A:ASN:H	1:14:A:CYS:HA	5	0.45
(1,525)	1:15:A:ASN:H	1:14:A:CYS:HA	13	0.45
(1,515)	1:14:A:CYS:H	1:13:A:TYR:HB3	8	0.45
(1,507)	1:22:A:LEU:H	1:22:A:LEU:HD13	14	0.45
(1,480)	1:45:A:ILE:H	1:44:A:CYS:HB2	12	0.45
(1,467)	1:30:A:ILE:H	1:29:A:ILE:HG21	30	0.45
(1,451)	1:25:A:LYS:H	1:24:A:VAL:HB	4	0.45
(1,451)	1:25:A:LYS:H	1:24:A:VAL:HB	17	0.45
(1,437)	1:24:A:VAL:HG22	1:30:A:ILE:HG22	24	0.45
(1,434)	1:24:A:VAL:HG13	1:30:A:ILE:HD12	6	0.45
(1,399)	1:29:A:ILE:H	1:29:A:ILE:HD11	8	0.45
(1,399)	1:29:A:ILE:H	1:29:A:ILE:HD11	24	0.45
(1,399)	1:29:A:ILE:H	1:29:A:ILE:HD12	26	0.45
(1,391)	1:34:A:GLU:H	1:5:A:LEU:HD21	3	0.45
(1,330)	1:63:A:GLY:H	1:62:A:LEU:HB2	6	0.45
(1,324)	1:4:A:MET:H	1:4:A:MET:HG3	23	0.45
(1,220)	1:15:A:ASN:H	1:15:A:ASN:HD22	17	0.45
(1,211)	1:25:A:LYS:H	1:11:A:LYS:H	20	0.45
(1,191)	1:6:A:GLN:H	1:6:A:GLN:HG2	19	0.45
(1,174)	1:30:A:ILE:HG12	1:9:A:ALA:H	16	0.45

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,174)	1:30:A:ILE:HG12	1:9:A:ALA:H	21	0.45
(1,107)	1:41:A:TRP:HB2	1:41:A:TRP:HD1	9	0.45
(1,107)	1:41:A:TRP:HB2	1:41:A:TRP:HD1	18	0.45
(1,58)	1:28:A:ASP:H	1:28:A:ASP:HB2	16	0.45
(1,953)	1:23:A:ASN:HB3	1:45:A:ILE:HG21	12	0.44
(1,944)	1:30:A:ILE:HD12	1:7:A:VAL:HG21	30	0.44
(1,940)	1:7:A:VAL:H	1:32:A:VAL:HG11	7	0.44
(1,929)	1:13:A:TYR:HD1	1:11:A:LYS:HG2	22	0.44
(1,923)	1:33:A:LEU:HD12	1:33:A:LEU:HB3	6	0.44
(1,923)	1:33:A:LEU:HD11	1:33:A:LEU:HB3	26	0.44
(1,919)	1:23:A:ASN:HD21	1:25:A:LYS:HB3	5	0.44
(1,917)	1:25:A:LYS:HB3	1:25:A:LYS:H	17	0.44
(1,908)	1:52:A:ASN:H	1:49:A:ARG:HB2	1	0.44
(1,896)	1:48:A:ASN:H	1:47:A:ASP:HB2	7	0.44
(1,840)	1:11:A:LYS:HD3	1:12:A:ASP:H	9	0.44
(1,840)	1:11:A:LYS:HD3	1:12:A:ASP:H	24	0.44
(1,816)	1:5:A:LEU:HB2	1:7:A:VAL:H	1	0.44
(1,810)	1:23:A:ASN:H	1:45:A:ILE:H	25	0.44
(1,798)	1:24:A:VAL:HB	1:12:A:ASP:HA	28	0.44
(1,707)	1:35:A:GLN:HE22	1:35:A:GLN:HB3	22	0.44
(1,664)	1:61:A:SER:H	1:61:A:SER:HB3	2	0.44
(1,664)	1:61:A:SER:H	1:61:A:SER:HB3	20	0.44
(1,627)	1:38:A:ASP:H	1:37:A:PRO:HB3	1	0.44
(1,541)	1:40:A:ARG:H	1:40:A:ARG:HD2	7	0.44
(1,525)	1:15:A:ASN:H	1:14:A:CYS:HA	2	0.44
(1,525)	1:15:A:ASN:H	1:14:A:CYS:HA	9	0.44
(1,525)	1:15:A:ASN:H	1:14:A:CYS:HA	14	0.44
(1,525)	1:15:A:ASN:H	1:14:A:CYS:HA	18	0.44
(1,507)	1:22:A:LEU:H	1:22:A:LEU:HD12	28	0.44
(1,480)	1:45:A:ILE:H	1:44:A:CYS:HB2	23	0.44
(1,467)	1:30:A:ILE:H	1:29:A:ILE:HG22	19	0.44
(1,467)	1:30:A:ILE:H	1:29:A:ILE:HG22	22	0.44
(1,466)	1:30:A:ILE:H	1:29:A:ILE:HB	10	0.44
(1,444)	1:30:A:ILE:HD13	1:32:A:VAL:HG11	9	0.44
(1,431)	1:22:A:LEU:HD11	1:24:A:VAL:HG11	24	0.44
(1,226)	1:39:A:GLY:HA3	1:41:A:TRP:HD1	9	0.44
(1,203)	1:52:A:ASN:HD22	1:52:A:ASN:H	7	0.44
(1,182)	1:25:A:LYS:H	1:27:A:GLY:H	22	0.44
(1,74)	1:15:A:ASN:HD22	1:15:A:ASN:HB2	13	0.44
(1,74)	1:15:A:ASN:HD22	1:15:A:ASN:HB2	14	0.44
(1,69)	1:52:A:ASN:H	1:52:A:ASN:HB2	29	0.44
(1,956)	1:24:A:VAL:HG23	1:13:A:TYR:HD2	26	0.43

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,956)	1:24:A:VAL:HG21	1:13:A:TYR:HD2	28	0.43
(1,939)	1:58:A:PHE:HD1	1:32:A:VAL:HG13	9	0.43
(1,923)	1:33:A:LEU:HD13	1:33:A:LEU:HB3	27	0.43
(1,919)	1:23:A:ASN:HD21	1:54:A:ARG:HB2	28	0.43
(1,917)	1:25:A:LYS:HB3	1:25:A:LYS:H	11	0.43
(1,878)	1:52:A:ASN:H	1:50:A:THR:HB	26	0.43
(1,825)	1:19:A:LEU:HB3	1:20:A:THR:H	15	0.43
(1,810)	1:23:A:ASN:H	1:45:A:ILE:H	24	0.43
(1,803)	1:30:A:ILE:HG22	1:32:A:VAL:H	5	0.43
(1,803)	1:30:A:ILE:HG21	1:32:A:VAL:H	21	0.43
(1,803)	1:30:A:ILE:HG21	1:32:A:VAL:H	29	0.43
(1,802)	1:26:A:ALA:HB2	1:12:A:ASP:HA	7	0.43
(1,800)	1:44:A:CYS:HB3	1:55:A:VAL:HA	24	0.43
(1,795)	1:17:A:TYR:HB3	1:18:A:ASP:H	15	0.43
(1,670)	1:43:A:GLY:H	1:42:A:LYS:HB2	2	0.43
(1,627)	1:38:A:ASP:H	1:37:A:PRO:HB3	3	0.43
(1,554)	1:54:A:ARG:H	1:54:A:ARG:HB2	15	0.43
(1,545)	1:40:A:ARG:H	1:36:A:HIS:HB2	6	0.43
(1,525)	1:15:A:ASN:H	1:14:A:CYS:HA	12	0.43
(1,525)	1:15:A:ASN:H	1:14:A:CYS:HA	15	0.43
(1,525)	1:15:A:ASN:H	1:14:A:CYS:HA	16	0.43
(1,525)	1:15:A:ASN:H	1:14:A:CYS:HA	26	0.43
(1,493)	1:41:A:TRP:HE1	1:40:A:ARG:H	29	0.43
(1,431)	1:22:A:LEU:HD12	1:24:A:VAL:HG11	13	0.43
(1,420)	1:7:A:VAL:HG11	1:32:A:VAL:HG13	3	0.43
(1,382)	1:65:A:ALA:H	1:5:A:LEU:HB3	16	0.43
(1,191)	1:6:A:GLN:H	1:6:A:GLN:HG2	1	0.43
(1,191)	1:6:A:GLN:H	1:6:A:GLN:HG2	8	0.43
(1,191)	1:6:A:GLN:H	1:6:A:GLN:HG2	25	0.43
(1,182)	1:25:A:LYS:H	1:27:A:GLY:H	23	0.43
(1,176)	1:33:A:LEU:H	1:42:A:LYS:H	13	0.43
(1,174)	1:30:A:ILE:HG12	1:9:A:ALA:H	26	0.43
(1,107)	1:41:A:TRP:HB2	1:41:A:TRP:HD1	12	0.43
(1,107)	1:41:A:TRP:HB2	1:41:A:TRP:HD1	25	0.43
(1,74)	1:15:A:ASN:HD22	1:15:A:ASN:HB2	3	0.43
(1,74)	1:15:A:ASN:HD22	1:15:A:ASN:HB2	27	0.43
(1,62)	1:18:A:ASP:HB2	1:18:A:ASP:H	28	0.43
(1,58)	1:28:A:ASP:H	1:28:A:ASP:HB2	28	0.43
(1,14)	1:27:A:GLY:H	1:26:A:ALA:HA	16	0.43
(1,968)	1:10:A:THR:H	1:63:A:GLY:HA3	15	0.42
(1,968)	1:10:A:THR:H	1:26:A:ALA:HA	17	0.42
(1,967)	1:62:A:LEU:H	1:63:A:GLY:HA2	6	0.42

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,967)	1:62:A:LEU:H	1:63:A:GLY:HA2	11	0.42
(1,944)	1:30:A:ILE:HD11	1:7:A:VAL:HG21	6	0.42
(1,940)	1:7:A:VAL:H	1:32:A:VAL:HG13	28	0.42
(1,923)	1:33:A:LEU:HD13	1:33:A:LEU:HB3	28	0.42
(1,908)	1:34:A:GLU:H	1:35:A:GLN:HG2	3	0.42
(1,900)	1:14:A:CYS:HB3	1:24:A:VAL:H	13	0.42
(1,894)	1:23:A:ASN:H	1:54:A:ARG:HD2	16	0.42
(1,894)	1:23:A:ASN:H	1:54:A:ARG:HD2	30	0.42
(1,816)	1:5:A:LEU:HB2	1:7:A:VAL:H	6	0.42
(1,816)	1:5:A:LEU:HB2	1:7:A:VAL:H	9	0.42
(1,810)	1:23:A:ASN:H	1:45:A:ILE:H	23	0.42
(1,803)	1:30:A:ILE:HG21	1:32:A:VAL:H	3	0.42
(1,714)	1:15:A:ASN:HD22	1:15:A:ASN:HA	1	0.42
(1,670)	1:43:A:GLY:H	1:42:A:LYS:HB2	23	0.42
(1,664)	1:61:A:SER:H	1:61:A:SER:HB3	12	0.42
(1,466)	1:30:A:ILE:H	1:29:A:ILE:HB	1	0.42
(1,466)	1:30:A:ILE:H	1:29:A:ILE:HB	4	0.42
(1,466)	1:30:A:ILE:H	1:29:A:ILE:HB	8	0.42
(1,466)	1:30:A:ILE:H	1:29:A:ILE:HB	19	0.42
(1,464)	1:67:A:VAL:H	1:66:A:ILE:HG12	15	0.42
(1,441)	1:30:A:ILE:HG22	1:45:A:ILE:HG22	20	0.42
(1,434)	1:24:A:VAL:HG13	1:30:A:ILE:HD12	20	0.42
(1,420)	1:7:A:VAL:HG13	1:32:A:VAL:HG11	10	0.42
(1,330)	1:63:A:GLY:H	1:62:A:LEU:HB2	17	0.42
(1,317)	1:39:A:GLY:H	1:38:A:ASP:HB3	13	0.42
(1,268)	1:30:A:ILE:HG13	1:30:A:ILE:HA	3	0.42
(1,234)	1:8:A:ARG:HG3	1:6:A:GLN:HE22	2	0.42
(1,226)	1:39:A:GLY:HA3	1:41:A:TRP:HD1	24	0.42
(1,191)	1:6:A:GLN:H	1:6:A:GLN:HG2	28	0.42
(1,191)	1:6:A:GLN:H	1:6:A:GLN:HG2	30	0.42
(1,174)	1:30:A:ILE:HG12	1:9:A:ALA:H	5	0.42
(1,107)	1:41:A:TRP:HB2	1:41:A:TRP:HD1	6	0.42
(1,107)	1:41:A:TRP:HB2	1:41:A:TRP:HD1	23	0.42
(1,74)	1:15:A:ASN:HD22	1:15:A:ASN:HB2	5	0.42
(1,74)	1:15:A:ASN:HD22	1:15:A:ASN:HB2	9	0.42
(1,74)	1:15:A:ASN:HD22	1:15:A:ASN:HB2	15	0.42
(1,74)	1:15:A:ASN:HD22	1:15:A:ASN:HB2	19	0.42
(1,46)	1:13:A:TYR:HB2	1:11:A:LYS:HD2	28	0.42
(1,966)	1:8:A:ARG:H	1:64:A:GLU:HG2	7	0.41
(1,966)	1:8:A:ARG:H	1:64:A:GLU:HG2	17	0.41
(1,965)	1:26:A:ALA:H	1:12:A:ASP:HA	2	0.41
(1,944)	1:30:A:ILE:HD12	1:7:A:VAL:HG21	17	0.41

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,940)	1:44:A:CYS:H	1:32:A:VAL:HG13	4	0.41
(1,939)	1:58:A:PHE:HD1	1:32:A:VAL:HG13	22	0.41
(1,924)	1:7:A:VAL:HG13	1:65:A:ALA:HB1	28	0.41
(1,908)	1:34:A:GLU:H	1:35:A:GLN:HG3	4	0.41
(1,884)	1:62:A:LEU:H	1:62:A:LEU:HA	2	0.41
(1,884)	1:62:A:LEU:H	1:62:A:LEU:HA	8	0.41
(1,884)	1:62:A:LEU:H	1:62:A:LEU:HA	9	0.41
(1,884)	1:62:A:LEU:H	1:62:A:LEU:HA	12	0.41
(1,884)	1:62:A:LEU:H	1:62:A:LEU:HA	13	0.41
(1,884)	1:62:A:LEU:H	1:62:A:LEU:HA	14	0.41
(1,884)	1:62:A:LEU:H	1:62:A:LEU:HA	16	0.41
(1,884)	1:62:A:LEU:H	1:62:A:LEU:HA	22	0.41
(1,884)	1:62:A:LEU:H	1:62:A:LEU:HA	26	0.41
(1,884)	1:62:A:LEU:H	1:62:A:LEU:HA	27	0.41
(1,884)	1:62:A:LEU:H	1:62:A:LEU:HA	29	0.41
(1,870)	1:29:A:ILE:HG21	1:45:A:ILE:HA	5	0.41
(1,849)	1:13:A:TYR:H	1:15:A:ASN:HD22	27	0.41
(1,834)	1:51:A:GLY:HA3	1:47:A:ASP:H	13	0.41
(1,816)	1:5:A:LEU:HB2	1:7:A:VAL:H	3	0.41
(1,805)	1:30:A:ILE:H	1:30:A:ILE:HD13	9	0.41
(1,805)	1:30:A:ILE:H	1:30:A:ILE:HD12	14	0.41
(1,804)	1:22:A:LEU:HB3	1:24:A:VAL:H	17	0.41
(1,803)	1:30:A:ILE:HG21	1:32:A:VAL:H	10	0.41
(1,800)	1:44:A:CYS:HB3	1:55:A:VAL:HA	7	0.41
(1,738)	1:50:A:THR:H	1:50:A:THR:HB	19	0.41
(1,722)	1:51:A:GLY:H	1:49:A:ARG:HB2	17	0.41
(1,707)	1:35:A:GLN:HE22	1:35:A:GLN:HB3	29	0.41
(1,627)	1:38:A:ASP:H	1:37:A:PRO:HB3	30	0.41
(1,541)	1:40:A:ARG:H	1:40:A:ARG:HD2	9	0.41
(1,525)	1:15:A:ASN:H	1:14:A:CYS:HA	20	0.41
(1,485)	1:46:A:HIS:H	1:45:A:ILE:HD13	3	0.41
(1,480)	1:45:A:ILE:H	1:44:A:CYS:HB2	17	0.41
(1,480)	1:45:A:ILE:H	1:44:A:CYS:HB2	29	0.41
(1,441)	1:30:A:ILE:HG21	1:45:A:ILE:HG21	2	0.41
(1,434)	1:24:A:VAL:HG11	1:30:A:ILE:HD12	1	0.41
(1,432)	1:22:A:LEU:HD12	1:24:A:VAL:HG21	9	0.41
(1,423)	1:7:A:VAL:HG22	1:32:A:VAL:HG21	21	0.41
(1,414)	1:5:A:LEU:HD11	1:32:A:VAL:HG13	30	0.41
(1,317)	1:39:A:GLY:H	1:38:A:ASP:HB3	2	0.41
(1,228)	1:51:A:GLY:H	1:52:A:ASN:HD22	7	0.41
(1,107)	1:41:A:TRP:HB2	1:41:A:TRP:HD1	3	0.41
(1,107)	1:41:A:TRP:HB2	1:41:A:TRP:HD1	8	0.41

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,107)	1:41:A:TRP:HB2	1:41:A:TRP:HD1	10	0.41
(1,107)	1:41:A:TRP:HB2	1:41:A:TRP:HD1	16	0.41
(1,74)	1:15:A:ASN:HD22	1:15:A:ASN:HB2	28	0.41
(1,66)	1:28:A:ASP:HB3	1:25:A:LYS:HG2	12	0.41
(1,58)	1:28:A:ASP:H	1:28:A:ASP:HB2	21	0.41
(1,58)	1:28:A:ASP:H	1:28:A:ASP:HB2	27	0.41
(1,46)	1:13:A:TYR:HB2	1:11:A:LYS:HD2	19	0.41
(1,14)	1:27:A:GLY:H	1:26:A:ALA:HA	6	0.41
(1,14)	1:27:A:GLY:H	1:26:A:ALA:HA	15	0.41
(1,966)	1:8:A:ARG:H	1:64:A:GLU:HG3	23	0.4
(1,964)	1:45:A:ILE:H	1:55:A:VAL:HA	21	0.4
(1,946)	1:7:A:VAL:HG21	1:30:A:ILE:H	19	0.4
(1,945)	1:41:A:TRP:HH2	1:5:A:LEU:HD13	20	0.4
(1,909)	1:41:A:TRP:HD1	1:40:A:ARG:HG2	9	0.4
(1,909)	1:41:A:TRP:HD1	1:60:A:SER:HB2	13	0.4
(1,897)	1:41:A:TRP:HB2	1:41:A:TRP:HE3	24	0.4
(1,896)	1:48:A:ASN:H	1:25:A:LYS:HE3	10	0.4
(1,894)	1:53:A:ASP:H	1:54:A:ARG:HD3	26	0.4
(1,892)	1:41:A:TRP:HH2	1:61:A:SER:HA	8	0.4
(1,888)	1:5:A:LEU:HB3	1:67:A:VAL:HA	6	0.4
(1,888)	1:5:A:LEU:HB3	1:67:A:VAL:HA	10	0.4
(1,884)	1:62:A:LEU:H	1:62:A:LEU:HA	1	0.4
(1,884)	1:62:A:LEU:H	1:62:A:LEU:HA	5	0.4
(1,884)	1:62:A:LEU:H	1:62:A:LEU:HA	17	0.4
(1,847)	1:8:A:ARG:HD3	1:6:A:GLN:HE21	10	0.4
(1,823)	1:28:A:ASP:H	1:9:A:ALA:HB2	9	0.4
(1,819)	1:58:A:PHE:HD1	1:42:A:LYS:H	21	0.4
(1,803)	1:30:A:ILE:HG22	1:32:A:VAL:H	19	0.4
(1,803)	1:30:A:ILE:HG23	1:32:A:VAL:H	20	0.4
(1,798)	1:24:A:VAL:HB	1:12:A:ASP:HA	20	0.4
(1,795)	1:17:A:TYR:HB3	1:18:A:ASP:H	7	0.4
(1,795)	1:17:A:TYR:HB3	1:18:A:ASP:H	9	0.4
(1,738)	1:50:A:THR:H	1:50:A:THR:HB	12	0.4
(1,707)	1:35:A:GLN:HE22	1:35:A:GLN:HB3	6	0.4
(1,694)	1:52:A:ASN:HD22	1:50:A:THR:HB	25	0.4
(1,664)	1:61:A:SER:H	1:61:A:SER:HB3	24	0.4
(1,664)	1:61:A:SER:H	1:61:A:SER:HB3	28	0.4
(1,627)	1:38:A:ASP:H	1:37:A:PRO:HB3	5	0.4
(1,560)	1:64:A:GLU:H	1:64:A:GLU:HG2	4	0.4
(1,525)	1:15:A:ASN:H	1:14:A:CYS:HA	11	0.4
(1,493)	1:41:A:TRP:HE1	1:40:A:ARG:H	23	0.4
(1,467)	1:30:A:ILE:H	1:29:A:ILE:HG23	13	0.4

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,437)	1:24:A:VAL:HG21	1:30:A:ILE:HG23	5	0.4
(1,429)	1:22:A:LEU:HD13	1:30:A:ILE:HG22	8	0.4
(1,427)	1:20:A:THR:HG21	1:55:A:VAL:HG21	3	0.4
(1,399)	1:29:A:ILE:H	1:29:A:ILE:HD11	16	0.4
(1,399)	1:29:A:ILE:H	1:29:A:ILE:HD11	27	0.4
(1,360)	1:27:A:GLY:H	1:25:A:LYS:HG3	21	0.4
(1,300)	1:4:A:MET:H	1:3:A:HIS:HB2	18	0.4
(1,283)	1:13:A:TYR:HE2	1:56:A:GLY:HA3	30	0.4
(1,249)	1:13:A:TYR:HE2	1:21:A:SER:HA	17	0.4
(1,249)	1:13:A:TYR:HE2	1:21:A:SER:HA	24	0.4
(1,234)	1:8:A:ARG:HG3	1:6:A:GLN:HE22	27	0.4
(1,191)	1:6:A:GLN:H	1:6:A:GLN:HG2	6	0.4
(1,174)	1:30:A:ILE:HG12	1:9:A:ALA:H	15	0.4
(1,174)	1:30:A:ILE:HG12	1:9:A:ALA:H	24	0.4
(1,69)	1:52:A:ASN:H	1:52:A:ASN:HB2	5	0.4
(1,69)	1:52:A:ASN:H	1:52:A:ASN:HB2	20	0.4
(1,55)	1:47:A:ASP:HB3	1:52:A:ASN:H	7	0.4
(1,39)	1:53:A:ASP:HB3	1:54:A:ARG:H	30	0.4
(1,14)	1:27:A:GLY:H	1:26:A:ALA:HA	8	0.4
(1,968)	1:10:A:THR:H	1:63:A:GLY:HA3	5	0.39
(1,953)	1:23:A:ASN:HB2	1:45:A:ILE:HG22	10	0.39
(1,952)	1:23:A:ASN:H	1:45:A:ILE:HD11	24	0.39
(1,940)	1:7:A:VAL:H	1:32:A:VAL:HG11	2	0.39
(1,940)	1:44:A:CYS:H	1:32:A:VAL:HG11	29	0.39
(1,934)	1:26:A:ALA:H	1:25:A:LYS:HG3	20	0.39
(1,917)	1:25:A:LYS:HB3	1:25:A:LYS:H	9	0.39
(1,892)	1:41:A:TRP:HH2	1:61:A:SER:HA	28	0.39
(1,884)	1:62:A:LEU:H	1:62:A:LEU:HA	15	0.39
(1,884)	1:62:A:LEU:H	1:62:A:LEU:HA	20	0.39
(1,884)	1:62:A:LEU:H	1:62:A:LEU:HA	25	0.39
(1,881)	1:13:A:TYR:HD2	1:21:A:SER:HB2	11	0.39
(1,834)	1:51:A:GLY:HA3	1:47:A:ASP:H	6	0.39
(1,830)	1:23:A:ASN:H	1:45:A:ILE:H	22	0.39
(1,819)	1:58:A:PHE:HD1	1:42:A:LYS:H	6	0.39
(1,819)	1:58:A:PHE:HD1	1:42:A:LYS:H	10	0.39
(1,793)	1:28:A:ASP:HB3	1:45:A:ILE:HG21	4	0.39
(1,793)	1:28:A:ASP:HB3	1:45:A:ILE:HG22	9	0.39
(1,670)	1:43:A:GLY:H	1:42:A:LYS:HB2	6	0.39
(1,670)	1:43:A:GLY:H	1:42:A:LYS:HB2	19	0.39
(1,571)	1:66:A:ILE:H	1:65:A:ALA:HB2	7	0.39
(1,525)	1:15:A:ASN:H	1:14:A:CYS:HA	3	0.39
(1,525)	1:15:A:ASN:H	1:14:A:CYS:HA	7	0.39

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,512)	1:65:A:ALA:H	1:66:A:ILE:HG21	21	0.39
(1,493)	1:41:A:TRP:HE1	1:40:A:ARG:H	21	0.39
(1,466)	1:30:A:ILE:H	1:29:A:ILE:HB	14	0.39
(1,435)	1:24:A:VAL:HG12	1:30:A:ILE:HG23	30	0.39
(1,399)	1:29:A:ILE:H	1:29:A:ILE:HD11	14	0.39
(1,371)	1:28:A:ASP:H	1:10:A:THR:HG23	20	0.39
(1,369)	1:29:A:ILE:HG13	1:6:A:GLN:HE22	6	0.39
(1,367)	1:9:A:ALA:HB1	1:10:A:THR:H	9	0.39
(1,360)	1:27:A:GLY:H	1:25:A:LYS:HG3	13	0.39
(1,268)	1:30:A:ILE:HG13	1:30:A:ILE:HA	9	0.39
(1,191)	1:6:A:GLN:H	1:6:A:GLN:HG2	20	0.39
(1,191)	1:6:A:GLN:H	1:6:A:GLN:HG2	24	0.39
(1,191)	1:6:A:GLN:H	1:6:A:GLN:HG2	26	0.39
(1,69)	1:52:A:ASN:H	1:52:A:ASN:HB2	8	0.39
(1,55)	1:47:A:ASP:HB3	1:52:A:ASN:H	5	0.39
(1,966)	1:8:A:ARG:H	1:64:A:GLU:HG3	22	0.38
(1,965)	1:26:A:ALA:H	1:12:A:ASP:HA	11	0.38
(1,964)	1:45:A:ILE:H	1:55:A:VAL:HA	20	0.38
(1,953)	1:23:A:ASN:HB3	1:45:A:ILE:HG22	14	0.38
(1,944)	1:30:A:ILE:HD13	1:7:A:VAL:HG22	8	0.38
(1,944)	1:30:A:ILE:HD11	1:7:A:VAL:HG22	27	0.38
(1,944)	1:30:A:ILE:HD11	1:7:A:VAL:HG21	29	0.38
(1,940)	1:7:A:VAL:H	1:32:A:VAL:HG12	8	0.38
(1,940)	1:44:A:CYS:H	1:32:A:VAL:HG12	10	0.38
(1,940)	1:7:A:VAL:H	1:32:A:VAL:HG11	12	0.38
(1,925)	1:11:A:LYS:HG2	1:13:A:TYR:H	19	0.38
(1,919)	1:23:A:ASN:HD21	1:54:A:ARG:HB2	7	0.38
(1,917)	1:25:A:LYS:HB3	1:25:A:LYS:H	1	0.38
(1,906)	1:34:A:GLU:HG2	1:3:A:HIS:H	10	0.38
(1,887)	1:29:A:ILE:HG23	1:31:A:THR:HB	25	0.38
(1,884)	1:62:A:LEU:H	1:62:A:LEU:HA	18	0.38
(1,873)	1:41:A:TRP:HB3	1:34:A:GLU:HA	22	0.38
(1,866)	1:66:A:ILE:HG21	1:65:A:ALA:HA	28	0.38
(1,810)	1:23:A:ASN:H	1:45:A:ILE:H	28	0.38
(1,803)	1:30:A:ILE:HG22	1:32:A:VAL:H	25	0.38
(1,685)	1:27:A:GLY:H	1:24:A:VAL:HG23	28	0.38
(1,628)	1:38:A:ASP:H	1:36:A:HIS:HB2	14	0.38
(1,627)	1:38:A:ASP:H	1:37:A:PRO:HB3	4	0.38
(1,626)	1:38:A:ASP:H	1:38:A:ASP:HB2	1	0.38
(1,583)	1:55:A:VAL:H	1:55:A:VAL:HB	30	0.38
(1,534)	1:23:A:ASN:H	1:23:A:ASN:HB2	8	0.38
(1,525)	1:15:A:ASN:H	1:14:A:CYS:HA	10	0.38

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,466)	1:30:A:ILE:H	1:29:A:ILE:HB	20	0.38
(1,432)	1:22:A:LEU:HD11	1:24:A:VAL:HG23	19	0.38
(1,420)	1:7:A:VAL:HG12	1:32:A:VAL:HG11	7	0.38
(1,399)	1:29:A:ILE:H	1:29:A:ILE:HD11	10	0.38
(1,360)	1:27:A:GLY:H	1:25:A:LYS:HG2	3	0.38
(1,354)	1:13:A:TYR:HD2	1:22:A:LEU:HB2	30	0.38
(1,318)	1:44:A:CYS:HB3	1:45:A:ILE:H	12	0.38
(1,268)	1:30:A:ILE:HG13	1:30:A:ILE:HA	15	0.38
(1,268)	1:30:A:ILE:HG13	1:30:A:ILE:HA	26	0.38
(1,232)	1:32:A:VAL:HG11	1:41:A:TRP:HZ3	11	0.38
(1,232)	1:32:A:VAL:HG11	1:41:A:TRP:HZ3	19	0.38
(1,232)	1:32:A:VAL:HG11	1:41:A:TRP:HZ3	22	0.38
(1,226)	1:39:A:GLY:HA3	1:41:A:TRP:HD1	13	0.38
(1,107)	1:41:A:TRP:HB2	1:41:A:TRP:HD1	20	0.38
(1,66)	1:28:A:ASP:HB3	1:25:A:LYS:HG2	17	0.38
(1,44)	1:13:A:TYR:HB3	1:24:A:VAL:HB	22	0.38
(1,41)	1:13:A:TYR:HB2	1:24:A:VAL:HB	25	0.38
(1,29)	1:43:A:GLY:HA3	1:30:A:ILE:HG23	20	0.38
(1,964)	1:45:A:ILE:H	1:53:A:ASP:HA	12	0.37
(1,946)	1:7:A:VAL:HG22	1:30:A:ILE:H	14	0.37
(1,946)	1:7:A:VAL:HG22	1:30:A:ILE:H	22	0.37
(1,946)	1:7:A:VAL:HG21	1:30:A:ILE:H	29	0.37
(1,940)	1:7:A:VAL:H	1:32:A:VAL:HG12	16	0.37
(1,926)	1:28:A:ASP:H	1:25:A:LYS:HG2	11	0.37
(1,925)	1:11:A:LYS:HG2	1:13:A:TYR:H	6	0.37
(1,925)	1:11:A:LYS:HG2	1:13:A:TYR:H	24	0.37
(1,923)	1:33:A:LEU:HD13	1:33:A:LEU:HB3	20	0.37
(1,917)	1:25:A:LYS:HB3	1:25:A:LYS:H	8	0.37
(1,909)	1:41:A:TRP:HD1	1:60:A:SER:HB2	12	0.37
(1,909)	1:41:A:TRP:HD1	1:60:A:SER:HB2	14	0.37
(1,903)	1:60:A:SER:HB3	1:60:A:SER:HA	2	0.37
(1,903)	1:60:A:SER:HB3	1:60:A:SER:HA	18	0.37
(1,892)	1:41:A:TRP:HH2	1:61:A:SER:HA	30	0.37
(1,884)	1:62:A:LEU:H	1:62:A:LEU:HA	4	0.37
(1,884)	1:62:A:LEU:H	1:62:A:LEU:HA	10	0.37
(1,884)	1:62:A:LEU:H	1:62:A:LEU:HA	19	0.37
(1,884)	1:62:A:LEU:H	1:62:A:LEU:HA	21	0.37
(1,884)	1:62:A:LEU:H	1:62:A:LEU:HA	23	0.37
(1,878)	1:52:A:ASN:H	1:50:A:THR:HA	2	0.37
(1,873)	1:41:A:TRP:HB3	1:34:A:GLU:HA	20	0.37
(1,858)	1:13:A:TYR:HD2	1:23:A:ASN:HA	2	0.37
(1,852)	1:13:A:TYR:HE2	1:24:A:VAL:H	11	0.37

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,849)	1:13:A:TYR:H	1:15:A:ASN:HD22	9	0.37
(1,837)	1:36:A:HIS:HB2	1:39:A:GLY:H	15	0.37
(1,816)	1:5:A:LEU:HB2	1:7:A:VAL:H	4	0.37
(1,815)	1:44:A:CYS:H	1:30:A:ILE:HG21	10	0.37
(1,810)	1:23:A:ASN:H	1:45:A:ILE:H	10	0.37
(1,803)	1:30:A:ILE:HG22	1:32:A:VAL:H	4	0.37
(1,803)	1:30:A:ILE:HG21	1:32:A:VAL:H	9	0.37
(1,798)	1:24:A:VAL:HB	1:12:A:ASP:HA	8	0.37
(1,743)	1:63:A:GLY:H	1:62:A:LEU:HG	29	0.37
(1,583)	1:55:A:VAL:H	1:55:A:VAL:HB	27	0.37
(1,571)	1:66:A:ILE:H	1:65:A:ALA:HB3	16	0.37
(1,541)	1:40:A:ARG:H	1:40:A:ARG:HD2	4	0.37
(1,507)	1:22:A:LEU:H	1:22:A:LEU:HD12	8	0.37
(1,480)	1:45:A:ILE:H	1:44:A:CYS:HB2	13	0.37
(1,467)	1:30:A:ILE:H	1:29:A:ILE:HG21	10	0.37
(1,466)	1:30:A:ILE:H	1:29:A:ILE:HB	5	0.37
(1,437)	1:24:A:VAL:HG21	1:30:A:ILE:HG22	10	0.37
(1,420)	1:7:A:VAL:HG12	1:32:A:VAL:HG12	8	0.37
(1,395)	1:42:A:LYS:H	1:33:A:LEU:HD11	17	0.37
(1,391)	1:34:A:GLU:H	1:5:A:LEU:HD22	16	0.37
(1,268)	1:30:A:ILE:HG13	1:30:A:ILE:HA	1	0.37
(1,268)	1:30:A:ILE:HG13	1:30:A:ILE:HA	16	0.37
(1,268)	1:30:A:ILE:HG13	1:30:A:ILE:HA	17	0.37
(1,268)	1:30:A:ILE:HG13	1:30:A:ILE:HA	28	0.37
(1,225)	1:35:A:GLN:HG3	1:41:A:TRP:HD1	18	0.37
(1,191)	1:6:A:GLN:H	1:6:A:GLN:HG2	2	0.37
(1,191)	1:6:A:GLN:H	1:6:A:GLN:HG2	10	0.37
(1,107)	1:41:A:TRP:HB2	1:41:A:TRP:HD1	13	0.37
(1,107)	1:41:A:TRP:HB2	1:41:A:TRP:HD1	17	0.37
(1,69)	1:52:A:ASN:H	1:52:A:ASN:HB2	7	0.37
(1,69)	1:52:A:ASN:H	1:52:A:ASN:HB2	11	0.37
(1,55)	1:47:A:ASP:HB3	1:52:A:ASN:H	14	0.37
(1,964)	1:45:A:ILE:H	1:55:A:VAL:HA	18	0.36
(1,956)	1:24:A:VAL:HG23	1:13:A:TYR:HD2	1	0.36
(1,956)	1:5:A:LEU:HD23	1:41:A:TRP:HH2	7	0.36
(1,954)	1:44:A:CYS:HA	1:45:A:ILE:HD13	26	0.36
(1,946)	1:7:A:VAL:HG21	1:30:A:ILE:H	30	0.36
(1,918)	1:13:A:TYR:H	1:11:A:LYS:HD3	26	0.36
(1,909)	1:41:A:TRP:HD1	1:40:A:ARG:HG2	6	0.36
(1,884)	1:62:A:LEU:H	1:62:A:LEU:HA	11	0.36
(1,858)	1:13:A:TYR:HD2	1:23:A:ASN:HA	3	0.36
(1,848)	1:16:A:ASN:HB2	1:18:A:ASP:H	23	0.36

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,834)	1:51:A:GLY:HA3	1:47:A:ASP:H	19	0.36
(1,819)	1:58:A:PHE:HD1	1:42:A:LYS:H	9	0.36
(1,815)	1:44:A:CYS:H	1:30:A:ILE:HG22	15	0.36
(1,810)	1:23:A:ASN:H	1:45:A:ILE:H	13	0.36
(1,810)	1:23:A:ASN:H	1:45:A:ILE:H	27	0.36
(1,798)	1:24:A:VAL:HB	1:12:A:ASP:HA	29	0.36
(1,793)	1:28:A:ASP:HB3	1:45:A:ILE:HG23	27	0.36
(1,743)	1:63:A:GLY:H	1:62:A:LEU:HG	13	0.36
(1,724)	1:20:A:THR:H	1:14:A:CYS:HB2	19	0.36
(1,667)	1:43:A:GLY:H	1:44:A:CYS:H	21	0.36
(1,654)	1:17:A:TYR:H	1:17:A:TYR:HB3	10	0.36
(1,628)	1:38:A:ASP:H	1:36:A:HIS:HB2	7	0.36
(1,627)	1:38:A:ASP:H	1:37:A:PRO:HB3	7	0.36
(1,627)	1:38:A:ASP:H	1:37:A:PRO:HB3	18	0.36
(1,627)	1:38:A:ASP:H	1:37:A:PRO:HB3	27	0.36
(1,587)	1:53:A:ASP:H	1:53:A:ASP:HB2	6	0.36
(1,583)	1:55:A:VAL:H	1:55:A:VAL:HB	16	0.36
(1,536)	1:23:A:ASN:H	1:22:A:LEU:HD11	30	0.36
(1,493)	1:41:A:TRP:HE1	1:40:A:ARG:H	12	0.36
(1,480)	1:45:A:ILE:H	1:44:A:CYS:HB2	21	0.36
(1,467)	1:30:A:ILE:H	1:29:A:ILE:HG23	9	0.36
(1,441)	1:30:A:ILE:HG21	1:45:A:ILE:HG22	30	0.36
(1,437)	1:24:A:VAL:HG21	1:30:A:ILE:HG23	11	0.36
(1,420)	1:7:A:VAL:HG13	1:32:A:VAL:HG12	21	0.36
(1,403)	1:58:A:PHE:H	1:22:A:LEU:HD13	22	0.36
(1,399)	1:29:A:ILE:H	1:29:A:ILE:HD12	6	0.36
(1,399)	1:29:A:ILE:H	1:29:A:ILE:HD12	30	0.36
(1,395)	1:42:A:LYS:H	1:33:A:LEU:HD13	16	0.36
(1,392)	1:43:A:GLY:H	1:33:A:LEU:HD13	15	0.36
(1,392)	1:43:A:GLY:H	1:33:A:LEU:HD11	21	0.36
(1,372)	1:33:A:LEU:HD22	1:33:A:LEU:HG	7	0.36
(1,326)	1:41:A:TRP:HZ2	1:34:A:GLU:HG2	28	0.36
(1,268)	1:30:A:ILE:HG13	1:30:A:ILE:HA	29	0.36
(1,220)	1:15:A:ASN:H	1:15:A:ASN:HD22	1	0.36
(1,220)	1:15:A:ASN:H	1:15:A:ASN:HD22	10	0.36
(1,213)	1:12:A:ASP:HB3	1:15:A:ASN:HD21	1	0.36
(1,211)	1:25:A:LYS:H	1:11:A:LYS:H	15	0.36
(1,211)	1:25:A:LYS:H	1:11:A:LYS:H	30	0.36
(1,203)	1:52:A:ASN:HD22	1:52:A:ASN:H	21	0.36
(1,192)	1:67:A:VAL:H	1:6:A:GLN:H	24	0.36
(1,182)	1:25:A:LYS:H	1:27:A:GLY:H	10	0.36
(1,182)	1:25:A:LYS:H	1:27:A:GLY:H	25	0.36

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,174)	1:30:A:ILE:HG12	1:9:A:ALA:H	25	0.36
(1,85)	1:7:A:VAL:HB	1:7:A:VAL:HG13	16	0.36
(1,85)	1:7:A:VAL:HB	1:7:A:VAL:HG13	30	0.36
(1,60)	1:28:A:ASP:HB2	1:28:A:ASP:HA	6	0.36
(1,58)	1:28:A:ASP:H	1:28:A:ASP:HB2	12	0.36
(1,946)	1:7:A:VAL:HG22	1:30:A:ILE:H	1	0.35
(1,945)	1:41:A:TRP:HH2	1:5:A:LEU:HD13	17	0.35
(1,940)	1:44:A:CYS:H	1:32:A:VAL:HG13	17	0.35
(1,870)	1:29:A:ILE:HG21	1:45:A:ILE:HA	30	0.35
(1,858)	1:13:A:TYR:HD2	1:23:A:ASN:HA	16	0.35
(1,830)	1:9:A:ALA:H	1:29:A:ILE:H	8	0.35
(1,803)	1:30:A:ILE:HG23	1:32:A:VAL:H	6	0.35
(1,803)	1:30:A:ILE:HG23	1:32:A:VAL:H	8	0.35
(1,803)	1:30:A:ILE:HG21	1:32:A:VAL:H	16	0.35
(1,798)	1:24:A:VAL:HB	1:12:A:ASP:HA	21	0.35
(1,743)	1:63:A:GLY:H	1:62:A:LEU:HG	17	0.35
(1,738)	1:50:A:THR:H	1:50:A:THR:HB	22	0.35
(1,727)	1:39:A:GLY:H	1:40:A:ARG:HG2	5	0.35
(1,685)	1:27:A:GLY:H	1:24:A:VAL:HG22	20	0.35
(1,669)	1:43:A:GLY:H	1:55:A:VAL:HG22	26	0.35
(1,667)	1:43:A:GLY:H	1:44:A:CYS:H	17	0.35
(1,667)	1:43:A:GLY:H	1:44:A:CYS:H	18	0.35
(1,654)	1:17:A:TYR:H	1:17:A:TYR:HB3	4	0.35
(1,560)	1:64:A:GLU:H	1:64:A:GLU:HG2	16	0.35
(1,554)	1:54:A:ARG:H	1:54:A:ARG:HB2	8	0.35
(1,480)	1:45:A:ILE:H	1:44:A:CYS:HB2	16	0.35
(1,467)	1:30:A:ILE:H	1:29:A:ILE:HG22	4	0.35
(1,445)	1:30:A:ILE:HD13	1:45:A:ILE:HG21	4	0.35
(1,432)	1:22:A:LEU:HD11	1:24:A:VAL:HG21	14	0.35
(1,420)	1:7:A:VAL:HG11	1:32:A:VAL:HG11	22	0.35
(1,399)	1:29:A:ILE:H	1:29:A:ILE:HD12	13	0.35
(1,392)	1:43:A:GLY:H	1:33:A:LEU:HD12	12	0.35
(1,392)	1:43:A:GLY:H	1:33:A:LEU:HD11	27	0.35
(1,372)	1:33:A:LEU:HD22	1:33:A:LEU:HG	6	0.35
(1,372)	1:33:A:LEU:HD23	1:33:A:LEU:HG	8	0.35
(1,372)	1:33:A:LEU:HD21	1:33:A:LEU:HG	17	0.35
(1,372)	1:33:A:LEU:HD21	1:33:A:LEU:HG	18	0.35
(1,372)	1:33:A:LEU:HD21	1:33:A:LEU:HG	24	0.35
(1,372)	1:33:A:LEU:HD21	1:33:A:LEU:HG	26	0.35
(1,372)	1:33:A:LEU:HD23	1:33:A:LEU:HG	27	0.35
(1,371)	1:28:A:ASP:H	1:10:A:THR:HG21	26	0.35
(1,317)	1:39:A:GLY:H	1:38:A:ASP:HB3	10	0.35

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,300)	1:4:A:MET:H	1:3:A:HIS:HB2	2	0.35
(1,268)	1:30:A:ILE:HG13	1:30:A:ILE:HA	14	0.35
(1,268)	1:30:A:ILE:HG13	1:30:A:ILE:HA	30	0.35
(1,218)	1:41:A:TRP:HZ3	1:41:A:TRP:HE3	1	0.35
(1,218)	1:41:A:TRP:HZ3	1:41:A:TRP:HE3	2	0.35
(1,218)	1:41:A:TRP:HZ3	1:41:A:TRP:HE3	3	0.35
(1,218)	1:41:A:TRP:HZ3	1:41:A:TRP:HE3	4	0.35
(1,218)	1:41:A:TRP:HZ3	1:41:A:TRP:HE3	5	0.35
(1,218)	1:41:A:TRP:HZ3	1:41:A:TRP:HE3	6	0.35
(1,218)	1:41:A:TRP:HZ3	1:41:A:TRP:HE3	7	0.35
(1,218)	1:41:A:TRP:HZ3	1:41:A:TRP:HE3	8	0.35
(1,218)	1:41:A:TRP:HZ3	1:41:A:TRP:HE3	9	0.35
(1,218)	1:41:A:TRP:HZ3	1:41:A:TRP:HE3	10	0.35
(1,218)	1:41:A:TRP:HZ3	1:41:A:TRP:HE3	11	0.35
(1,218)	1:41:A:TRP:HZ3	1:41:A:TRP:HE3	12	0.35
(1,218)	1:41:A:TRP:HZ3	1:41:A:TRP:HE3	13	0.35
(1,218)	1:41:A:TRP:HZ3	1:41:A:TRP:HE3	14	0.35
(1,218)	1:41:A:TRP:HZ3	1:41:A:TRP:HE3	15	0.35
(1,218)	1:41:A:TRP:HZ3	1:41:A:TRP:HE3	16	0.35
(1,218)	1:41:A:TRP:HZ3	1:41:A:TRP:HE3	17	0.35
(1,218)	1:41:A:TRP:HZ3	1:41:A:TRP:HE3	18	0.35
(1,218)	1:41:A:TRP:HZ3	1:41:A:TRP:HE3	19	0.35
(1,218)	1:41:A:TRP:HZ3	1:41:A:TRP:HE3	20	0.35
(1,218)	1:41:A:TRP:HZ3	1:41:A:TRP:HE3	21	0.35
(1,218)	1:41:A:TRP:HZ3	1:41:A:TRP:HE3	22	0.35
(1,218)	1:41:A:TRP:HZ3	1:41:A:TRP:HE3	23	0.35
(1,218)	1:41:A:TRP:HZ3	1:41:A:TRP:HE3	24	0.35
(1,218)	1:41:A:TRP:HZ3	1:41:A:TRP:HE3	25	0.35
(1,218)	1:41:A:TRP:HZ3	1:41:A:TRP:HE3	26	0.35
(1,218)	1:41:A:TRP:HZ3	1:41:A:TRP:HE3	27	0.35
(1,218)	1:41:A:TRP:HZ3	1:41:A:TRP:HE3	28	0.35
(1,218)	1:41:A:TRP:HZ3	1:41:A:TRP:HE3	29	0.35
(1,218)	1:41:A:TRP:HZ3	1:41:A:TRP:HE3	30	0.35
(1,196)	1:34:A:GLU:HB3	1:4:A:MET:H	22	0.35
(1,191)	1:6:A:GLN:H	1:6:A:GLN:HG2	23	0.35
(1,107)	1:41:A:TRP:HB2	1:41:A:TRP:HD1	1	0.35
(1,107)	1:41:A:TRP:HB2	1:41:A:TRP:HD1	11	0.35
(1,85)	1:7:A:VAL:HB	1:7:A:VAL:HG11	2	0.35
(1,85)	1:7:A:VAL:HB	1:7:A:VAL:HG11	4	0.35
(1,85)	1:7:A:VAL:HB	1:7:A:VAL:HG13	6	0.35
(1,85)	1:7:A:VAL:HB	1:7:A:VAL:HG13	19	0.35
(1,85)	1:7:A:VAL:HB	1:7:A:VAL:HG11	21	0.35

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,85)	1:7:A:VAL:HB	1:7:A:VAL:HG12	29	0.35
(1,69)	1:52:A:ASN:H	1:52:A:ASN:HB2	2	0.35
(1,62)	1:18:A:ASP:HB2	1:18:A:ASP:H	20	0.35
(1,46)	1:13:A:TYR:HB2	1:11:A:LYS:HD2	26	0.35
(1,968)	1:10:A:THR:H	1:63:A:GLY:HA3	21	0.34
(1,967)	1:62:A:LEU:H	1:63:A:GLY:HA2	4	0.34
(1,954)	1:44:A:CYS:HA	1:45:A:ILE:HD12	24	0.34
(1,952)	1:23:A:ASN:H	1:45:A:ILE:HD12	26	0.34
(1,943)	1:7:A:VAL:HB	1:7:A:VAL:HG11	2	0.34
(1,943)	1:7:A:VAL:HB	1:7:A:VAL:HG11	4	0.34
(1,943)	1:66:A:ILE:HG12	1:66:A:ILE:HD13	6	0.34
(1,943)	1:66:A:ILE:HG12	1:66:A:ILE:HD13	21	0.34
(1,943)	1:7:A:VAL:HB	1:7:A:VAL:HG12	29	0.34
(1,943)	1:66:A:ILE:HB	1:66:A:ILE:HG21	30	0.34
(1,909)	1:41:A:TRP:HD1	1:40:A:ARG:HG2	16	0.34
(1,908)	1:34:A:GLU:H	1:35:A:GLN:HG3	30	0.34
(1,906)	1:34:A:GLU:HG2	1:3:A:HIS:H	23	0.34
(1,900)	1:14:A:CYS:HB3	1:24:A:VAL:H	10	0.34
(1,859)	1:67:A:VAL:H	1:7:A:VAL:HA	5	0.34
(1,840)	1:11:A:LYS:HD3	1:12:A:ASP:H	2	0.34
(1,819)	1:58:A:PHE:HD1	1:42:A:LYS:H	11	0.34
(1,805)	1:30:A:ILE:H	1:30:A:ILE:HD11	29	0.34
(1,803)	1:30:A:ILE:HG21	1:32:A:VAL:H	7	0.34
(1,803)	1:30:A:ILE:HG22	1:32:A:VAL:H	15	0.34
(1,803)	1:30:A:ILE:HG22	1:32:A:VAL:H	23	0.34
(1,743)	1:63:A:GLY:H	1:62:A:LEU:HG	5	0.34
(1,727)	1:39:A:GLY:H	1:40:A:ARG:HG2	3	0.34
(1,705)	1:35:A:GLN:HE22	1:33:A:LEU:HD21	7	0.34
(1,694)	1:52:A:ASN:HD22	1:50:A:THR:HB	28	0.34
(1,667)	1:43:A:GLY:H	1:44:A:CYS:H	12	0.34
(1,667)	1:43:A:GLY:H	1:44:A:CYS:H	20	0.34
(1,655)	1:18:A:ASP:H	1:17:A:TYR:H	25	0.34
(1,628)	1:38:A:ASP:H	1:36:A:HIS:HB2	25	0.34
(1,627)	1:38:A:ASP:H	1:37:A:PRO:HB3	29	0.34
(1,595)	1:62:A:LEU:H	1:62:A:LEU:HB2	7	0.34
(1,545)	1:40:A:ARG:H	1:36:A:HIS:HB2	24	0.34
(1,515)	1:14:A:CYS:H	1:13:A:TYR:HB3	11	0.34
(1,480)	1:45:A:ILE:H	1:44:A:CYS:HB2	19	0.34
(1,423)	1:7:A:VAL:HG21	1:32:A:VAL:HG23	10	0.34
(1,423)	1:7:A:VAL:HG22	1:32:A:VAL:HG21	19	0.34
(1,423)	1:7:A:VAL:HG23	1:32:A:VAL:HG22	20	0.34
(1,395)	1:42:A:LYS:H	1:33:A:LEU:HD11	19	0.34

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,389)	1:11:A:LYS:HB3	1:24:A:VAL:HG11	14	0.34
(1,372)	1:33:A:LEU:HD23	1:33:A:LEU:HG	1	0.34
(1,372)	1:33:A:LEU:HD23	1:33:A:LEU:HG	3	0.34
(1,372)	1:33:A:LEU:HD21	1:33:A:LEU:HG	9	0.34
(1,372)	1:33:A:LEU:HD23	1:33:A:LEU:HG	10	0.34
(1,372)	1:33:A:LEU:HD22	1:33:A:LEU:HG	11	0.34
(1,372)	1:33:A:LEU:HD21	1:33:A:LEU:HG	12	0.34
(1,372)	1:33:A:LEU:HD23	1:33:A:LEU:HG	13	0.34
(1,372)	1:33:A:LEU:HD23	1:33:A:LEU:HG	16	0.34
(1,372)	1:33:A:LEU:HD22	1:33:A:LEU:HG	19	0.34
(1,372)	1:33:A:LEU:HD22	1:33:A:LEU:HG	20	0.34
(1,372)	1:33:A:LEU:HD23	1:33:A:LEU:HG	21	0.34
(1,372)	1:33:A:LEU:HD22	1:33:A:LEU:HG	22	0.34
(1,372)	1:33:A:LEU:HD22	1:33:A:LEU:HG	30	0.34
(1,346)	1:55:A:VAL:H	1:54:A:ARG:HB3	13	0.34
(1,330)	1:63:A:GLY:H	1:62:A:LEU:HB2	15	0.34
(1,268)	1:30:A:ILE:HG13	1:30:A:ILE:HA	7	0.34
(1,268)	1:30:A:ILE:HG13	1:30:A:ILE:HA	11	0.34
(1,268)	1:30:A:ILE:HG13	1:30:A:ILE:HA	12	0.34
(1,268)	1:30:A:ILE:HG13	1:30:A:ILE:HA	24	0.34
(1,191)	1:6:A:GLN:H	1:6:A:GLN:HG2	5	0.34
(1,190)	1:34:A:GLU:HG2	1:3:A:HIS:H	6	0.34
(1,106)	1:41:A:TRP:HB3	1:41:A:TRP:HZ3	21	0.34
(1,85)	1:7:A:VAL:HB	1:7:A:VAL:HG13	7	0.34
(1,85)	1:7:A:VAL:HB	1:7:A:VAL:HG23	11	0.34
(1,85)	1:7:A:VAL:HB	1:7:A:VAL:HG13	12	0.34
(1,85)	1:7:A:VAL:HB	1:7:A:VAL:HG11	15	0.34
(1,85)	1:7:A:VAL:HB	1:7:A:VAL:HG11	17	0.34
(1,85)	1:7:A:VAL:HB	1:7:A:VAL:HG13	18	0.34
(1,85)	1:7:A:VAL:HB	1:7:A:VAL:HG11	20	0.34
(1,85)	1:7:A:VAL:HB	1:7:A:VAL:HG13	22	0.34
(1,85)	1:7:A:VAL:HB	1:7:A:VAL:HG13	23	0.34
(1,85)	1:7:A:VAL:HB	1:7:A:VAL:HG22	24	0.34
(1,85)	1:7:A:VAL:HB	1:7:A:VAL:HG12	25	0.34
(1,85)	1:7:A:VAL:HB	1:7:A:VAL:HG13	27	0.34
(1,69)	1:52:A:ASN:H	1:52:A:ASN:HB2	14	0.34
(1,56)	1:28:A:ASP:H	1:28:A:ASP:HB3	30	0.34
(1,43)	1:13:A:TYR:HB2	1:24:A:VAL:HG23	17	0.34
(1,20)	1:11:A:LYS:H	1:26:A:ALA:HA	4	0.34
(1,967)	1:62:A:LEU:H	1:63:A:GLY:HA2	21	0.33
(1,966)	1:8:A:ARG:H	1:64:A:GLU:HG2	14	0.33
(1,964)	1:45:A:ILE:H	1:55:A:VAL:HA	16	0.33

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,964)	1:45:A:ILE:H	1:55:A:VAL:HA	23	0.33
(1,943)	1:66:A:ILE:HG12	1:66:A:ILE:HD11	11	0.33
(1,943)	1:7:A:VAL:HB	1:7:A:VAL:HG13	12	0.33
(1,943)	1:66:A:ILE:HG12	1:66:A:ILE:HD13	15	0.33
(1,943)	1:66:A:ILE:HB	1:66:A:ILE:HG23	16	0.33
(1,943)	1:66:A:ILE:HB	1:66:A:ILE:HG21	17	0.33
(1,943)	1:7:A:VAL:HB	1:7:A:VAL:HG13	18	0.33
(1,943)	1:7:A:VAL:HB	1:7:A:VAL:HG11	20	0.33
(1,943)	1:7:A:VAL:HB	1:7:A:VAL:HG13	22	0.33
(1,943)	1:7:A:VAL:HB	1:7:A:VAL:HG13	23	0.33
(1,943)	1:7:A:VAL:HB	1:7:A:VAL:HG22	24	0.33
(1,943)	1:66:A:ILE:HG12	1:66:A:ILE:HD11	25	0.33
(1,943)	1:7:A:VAL:HB	1:7:A:VAL:HG13	27	0.33
(1,937)	1:10:A:THR:HG21	1:10:A:THR:H	19	0.33
(1,917)	1:25:A:LYS:HB3	1:25:A:LYS:H	24	0.33
(1,912)	1:41:A:TRP:H	1:40:A:ARG:HG3	25	0.33
(1,909)	1:41:A:TRP:HD1	1:60:A:SER:HB2	11	0.33
(1,908)	1:34:A:GLU:H	1:35:A:GLN:HG3	19	0.33
(1,906)	1:34:A:GLU:HG3	1:3:A:HIS:H	4	0.33
(1,852)	1:13:A:TYR:HE2	1:24:A:VAL:H	26	0.33
(1,816)	1:5:A:LEU:HB2	1:7:A:VAL:H	2	0.33
(1,816)	1:5:A:LEU:HB2	1:7:A:VAL:H	23	0.33
(1,815)	1:44:A:CYS:H	1:30:A:ILE:HG22	2	0.33
(1,810)	1:23:A:ASN:H	1:45:A:ILE:H	21	0.33
(1,805)	1:30:A:ILE:H	1:30:A:ILE:HD11	1	0.33
(1,805)	1:30:A:ILE:H	1:30:A:ILE:HD12	7	0.33
(1,805)	1:30:A:ILE:H	1:30:A:ILE:HD11	10	0.33
(1,805)	1:30:A:ILE:H	1:30:A:ILE:HD11	13	0.33
(1,793)	1:28:A:ASP:HB3	1:24:A:VAL:HG12	21	0.33
(1,738)	1:50:A:THR:H	1:50:A:THR:HB	14	0.33
(1,669)	1:43:A:GLY:H	1:55:A:VAL:HG23	8	0.33
(1,667)	1:43:A:GLY:H	1:44:A:CYS:H	1	0.33
(1,667)	1:43:A:GLY:H	1:44:A:CYS:H	7	0.33
(1,667)	1:43:A:GLY:H	1:44:A:CYS:H	15	0.33
(1,667)	1:43:A:GLY:H	1:44:A:CYS:H	25	0.33
(1,654)	1:17:A:TYR:H	1:17:A:TYR:HB3	1	0.33
(1,654)	1:17:A:TYR:H	1:17:A:TYR:HB3	12	0.33
(1,654)	1:17:A:TYR:H	1:17:A:TYR:HB3	24	0.33
(1,627)	1:38:A:ASP:H	1:37:A:PRO:HB3	19	0.33
(1,480)	1:45:A:ILE:H	1:44:A:CYS:HB2	2	0.33
(1,480)	1:45:A:ILE:H	1:44:A:CYS:HB2	24	0.33
(1,467)	1:30:A:ILE:H	1:29:A:ILE:HG22	8	0.33

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,434)	1:24:A:VAL:HG12	1:30:A:ILE:HD13	22	0.33
(1,422)	1:7:A:VAL:HG22	1:32:A:VAL:HG11	30	0.33
(1,420)	1:7:A:VAL:HG11	1:32:A:VAL:HG11	9	0.33
(1,414)	1:5:A:LEU:HD13	1:32:A:VAL:HG13	5	0.33
(1,414)	1:5:A:LEU:HD12	1:32:A:VAL:HG13	28	0.33
(1,399)	1:29:A:ILE:H	1:29:A:ILE:HD11	2	0.33
(1,399)	1:29:A:ILE:H	1:29:A:ILE:HD13	4	0.33
(1,372)	1:33:A:LEU:HD22	1:33:A:LEU:HG	15	0.33
(1,372)	1:33:A:LEU:HD23	1:33:A:LEU:HG	23	0.33
(1,355)	1:33:A:LEU:HB3	1:33:A:LEU:H	9	0.33
(1,355)	1:33:A:LEU:HB3	1:33:A:LEU:H	14	0.33
(1,355)	1:33:A:LEU:HB3	1:33:A:LEU:H	15	0.33
(1,346)	1:55:A:VAL:H	1:54:A:ARG:HB3	24	0.33
(1,330)	1:63:A:GLY:H	1:62:A:LEU:HB2	12	0.33
(1,303)	1:25:A:LYS:HE2	1:48:A:ASN:HD22	11	0.33
(1,268)	1:30:A:ILE:HG13	1:30:A:ILE:HA	6	0.33
(1,268)	1:30:A:ILE:HG13	1:30:A:ILE:HA	25	0.33
(1,226)	1:39:A:GLY:HA3	1:41:A:TRP:HD1	3	0.33
(1,203)	1:52:A:ASN:HD22	1:52:A:ASN:H	12	0.33
(1,174)	1:30:A:ILE:HG12	1:9:A:ALA:H	3	0.33
(1,85)	1:7:A:VAL:HB	1:7:A:VAL:HG11	1	0.33
(1,85)	1:7:A:VAL:HB	1:7:A:VAL:HG11	5	0.33
(1,85)	1:7:A:VAL:HB	1:7:A:VAL:HG13	8	0.33
(1,85)	1:7:A:VAL:HB	1:7:A:VAL:HG11	10	0.33
(1,85)	1:7:A:VAL:HB	1:7:A:VAL:HG12	14	0.33
(1,85)	1:7:A:VAL:HB	1:7:A:VAL:HG11	26	0.33
(1,85)	1:7:A:VAL:HB	1:7:A:VAL:HG13	28	0.33
(1,60)	1:28:A:ASP:HB2	1:28:A:ASP:HA	18	0.33
(1,954)	1:44:A:CYS:HA	1:45:A:ILE:HD13	15	0.32
(1,954)	1:44:A:CYS:HA	1:45:A:ILE:HD13	17	0.32
(1,953)	1:23:A:ASN:HB2	1:45:A:ILE:HG23	3	0.32
(1,953)	1:23:A:ASN:HB3	1:45:A:ILE:HG21	7	0.32
(1,943)	1:7:A:VAL:HB	1:7:A:VAL:HG11	1	0.32
(1,943)	1:7:A:VAL:HB	1:7:A:VAL:HG11	5	0.32
(1,943)	1:66:A:ILE:HB	1:66:A:ILE:HG23	7	0.32
(1,943)	1:7:A:VAL:HB	1:7:A:VAL:HG13	8	0.32
(1,943)	1:7:A:VAL:HB	1:7:A:VAL:HG11	10	0.32
(1,943)	1:7:A:VAL:HB	1:7:A:VAL:HG12	14	0.32
(1,943)	1:7:A:VAL:HB	1:7:A:VAL:HG11	26	0.32
(1,943)	1:7:A:VAL:HB	1:7:A:VAL:HG13	28	0.32
(1,935)	1:40:A:ARG:HB2	1:38:A:ASP:HB3	29	0.32
(1,923)	1:33:A:LEU:HD13	1:33:A:LEU:HB2	7	0.32

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,909)	1:41:A:TRP:HD1	1:60:A:SER:HB2	10	0.32
(1,906)	1:34:A:GLU:HG2	1:3:A:HIS:H	27	0.32
(1,894)	1:23:A:ASN:H	1:54:A:ARG:HD3	11	0.32
(1,884)	1:62:A:LEU:H	1:62:A:LEU:HA	28	0.32
(1,853)	1:29:A:ILE:HG12	1:48:A:ASN:HD22	30	0.32
(1,834)	1:51:A:GLY:HA3	1:47:A:ASP:H	14	0.32
(1,819)	1:58:A:PHE:HD1	1:42:A:LYS:H	28	0.32
(1,816)	1:5:A:LEU:HB2	1:7:A:VAL:H	13	0.32
(1,816)	1:5:A:LEU:HB2	1:7:A:VAL:H	14	0.32
(1,816)	1:5:A:LEU:HB2	1:7:A:VAL:H	15	0.32
(1,816)	1:5:A:LEU:HB2	1:7:A:VAL:H	18	0.32
(1,816)	1:5:A:LEU:HB2	1:7:A:VAL:H	19	0.32
(1,816)	1:5:A:LEU:HB2	1:7:A:VAL:H	29	0.32
(1,814)	1:26:A:ALA:H	1:25:A:LYS:HB2	26	0.32
(1,810)	1:23:A:ASN:H	1:45:A:ILE:H	18	0.32
(1,803)	1:30:A:ILE:HG22	1:32:A:VAL:H	2	0.32
(1,803)	1:30:A:ILE:HG22	1:32:A:VAL:H	14	0.32
(1,798)	1:24:A:VAL:HB	1:12:A:ASP:HA	9	0.32
(1,798)	1:24:A:VAL:HB	1:12:A:ASP:HA	22	0.32
(1,795)	1:17:A:TYR:HB3	1:18:A:ASP:H	1	0.32
(1,795)	1:17:A:TYR:HB3	1:18:A:ASP:H	11	0.32
(1,714)	1:15:A:ASN:HD22	1:15:A:ASN:HA	11	0.32
(1,687)	1:27:A:GLY:H	1:26:A:ALA:HB2	19	0.32
(1,667)	1:43:A:GLY:H	1:44:A:CYS:H	3	0.32
(1,667)	1:43:A:GLY:H	1:44:A:CYS:H	8	0.32
(1,667)	1:43:A:GLY:H	1:44:A:CYS:H	9	0.32
(1,667)	1:43:A:GLY:H	1:44:A:CYS:H	13	0.32
(1,654)	1:17:A:TYR:H	1:17:A:TYR:HB3	11	0.32
(1,654)	1:17:A:TYR:H	1:17:A:TYR:HB3	13	0.32
(1,654)	1:17:A:TYR:H	1:17:A:TYR:HB3	25	0.32
(1,654)	1:17:A:TYR:H	1:17:A:TYR:HB3	28	0.32
(1,587)	1:53:A:ASP:H	1:53:A:ASP:HB2	28	0.32
(1,554)	1:54:A:ARG:H	1:54:A:ARG:HB2	19	0.32
(1,515)	1:14:A:CYS:H	1:13:A:TYR:HB3	9	0.32
(1,451)	1:25:A:LYS:H	1:24:A:VAL:HB	11	0.32
(1,431)	1:22:A:LEU:HD13	1:24:A:VAL:HG12	23	0.32
(1,399)	1:29:A:ILE:H	1:29:A:ILE:HD12	23	0.32
(1,391)	1:34:A:GLU:H	1:5:A:LEU:HD21	5	0.32
(1,391)	1:34:A:GLU:H	1:5:A:LEU:HD21	7	0.32
(1,372)	1:33:A:LEU:HD21	1:33:A:LEU:HG	14	0.32
(1,372)	1:33:A:LEU:HD23	1:33:A:LEU:HG	25	0.32
(1,372)	1:33:A:LEU:HD21	1:33:A:LEU:HG	28	0.32

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,330)	1:63:A:GLY:H	1:62:A:LEU:HB2	29	0.32
(1,268)	1:30:A:ILE:HG13	1:30:A:ILE:HA	5	0.32
(1,268)	1:30:A:ILE:HG13	1:30:A:ILE:HA	21	0.32
(1,268)	1:30:A:ILE:HG13	1:30:A:ILE:HA	22	0.32
(1,191)	1:6:A:GLN:H	1:6:A:GLN:HG2	16	0.32
(1,182)	1:25:A:LYS:H	1:27:A:GLY:H	29	0.32
(1,176)	1:33:A:LEU:H	1:42:A:LYS:H	17	0.32
(1,85)	1:7:A:VAL:HB	1:7:A:VAL:HG13	13	0.32
(1,69)	1:52:A:ASN:H	1:52:A:ASN:HB2	6	0.32
(1,60)	1:28:A:ASP:HB2	1:28:A:ASP:HA	10	0.32
(1,55)	1:47:A:ASP:HB3	1:52:A:ASN:H	15	0.32
(1,43)	1:13:A:TYR:HB2	1:24:A:VAL:HG21	9	0.32
(1,29)	1:43:A:GLY:HA3	1:30:A:ILE:HG22	2	0.32
(1,16)	1:26:A:ALA:HA	1:28:A:ASP:H	4	0.32
(1,944)	1:30:A:ILE:HD11	1:7:A:VAL:HG23	13	0.31
(1,943)	1:7:A:VAL:HB	1:7:A:VAL:HG13	13	0.31
(1,943)	1:66:A:ILE:HG12	1:66:A:ILE:HD11	19	0.31
(1,931)	1:41:A:TRP:HZ2	1:60:A:SER:HB3	28	0.31
(1,917)	1:25:A:LYS:HB3	1:25:A:LYS:H	28	0.31
(1,912)	1:41:A:TRP:H	1:40:A:ARG:HG3	5	0.31
(1,894)	1:23:A:ASN:H	1:54:A:ARG:HD2	9	0.31
(1,884)	1:62:A:LEU:H	1:62:A:LEU:HA	6	0.31
(1,830)	1:9:A:ALA:H	1:29:A:ILE:H	14	0.31
(1,815)	1:44:A:CYS:H	1:30:A:ILE:HG22	13	0.31
(1,815)	1:44:A:CYS:H	1:30:A:ILE:HG23	20	0.31
(1,805)	1:30:A:ILE:H	1:30:A:ILE:HD12	4	0.31
(1,805)	1:30:A:ILE:H	1:30:A:ILE:HD12	19	0.31
(1,804)	1:22:A:LEU:HB3	1:22:A:LEU:H	18	0.31
(1,804)	1:22:A:LEU:HB3	1:22:A:LEU:H	20	0.31
(1,803)	1:30:A:ILE:HG22	1:32:A:VAL:H	13	0.31
(1,743)	1:63:A:GLY:H	1:62:A:LEU:HG	20	0.31
(1,743)	1:63:A:GLY:H	1:62:A:LEU:HG	28	0.31
(1,714)	1:15:A:ASN:HD22	1:15:A:ASN:HA	21	0.31
(1,667)	1:43:A:GLY:H	1:44:A:CYS:H	6	0.31
(1,667)	1:43:A:GLY:H	1:44:A:CYS:H	19	0.31
(1,654)	1:17:A:TYR:H	1:17:A:TYR:HB3	3	0.31
(1,654)	1:17:A:TYR:H	1:17:A:TYR:HB3	9	0.31
(1,654)	1:17:A:TYR:H	1:17:A:TYR:HB3	21	0.31
(1,654)	1:17:A:TYR:H	1:17:A:TYR:HB3	29	0.31
(1,597)	1:62:A:LEU:H	1:62:A:LEU:HB3	7	0.31
(1,480)	1:45:A:ILE:H	1:44:A:CYS:HB2	6	0.31
(1,422)	1:7:A:VAL:HG23	1:32:A:VAL:HG13	1	0.31

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,403)	1:58:A:PHE:H	1:22:A:LEU:HD12	27	0.31
(1,403)	1:58:A:PHE:H	1:22:A:LEU:HD11	28	0.31
(1,386)	1:46:A:HIS:H	1:45:A:ILE:HG21	27	0.31
(1,372)	1:33:A:LEU:HD23	1:33:A:LEU:HG	2	0.31
(1,372)	1:33:A:LEU:HD23	1:33:A:LEU:HG	4	0.31
(1,372)	1:33:A:LEU:HD23	1:33:A:LEU:HG	5	0.31
(1,355)	1:33:A:LEU:HB3	1:33:A:LEU:H	5	0.31
(1,355)	1:33:A:LEU:HB3	1:33:A:LEU:H	13	0.31
(1,330)	1:63:A:GLY:H	1:62:A:LEU:HB2	3	0.31
(1,268)	1:30:A:ILE:HG13	1:30:A:ILE:HA	8	0.31
(1,268)	1:30:A:ILE:HG13	1:30:A:ILE:HA	13	0.31
(1,268)	1:30:A:ILE:HG13	1:30:A:ILE:HA	18	0.31
(1,220)	1:15:A:ASN:H	1:15:A:ASN:HD22	7	0.31
(1,211)	1:25:A:LYS:H	1:11:A:LYS:H	7	0.31
(1,191)	1:6:A:GLN:H	1:6:A:GLN:HG2	15	0.31
(1,191)	1:6:A:GLN:H	1:6:A:GLN:HG2	29	0.31
(1,190)	1:34:A:GLU:HG3	1:3:A:HIS:H	26	0.31
(1,85)	1:7:A:VAL:HB	1:7:A:VAL:HG13	3	0.31
(1,60)	1:28:A:ASP:HB2	1:28:A:ASP:HA	7	0.31
(1,60)	1:28:A:ASP:HB2	1:28:A:ASP:HA	21	0.31
(1,60)	1:28:A:ASP:HB2	1:28:A:ASP:HA	27	0.31
(1,53)	1:13:A:TYR:HB3	1:24:A:VAL:H	25	0.31
(1,29)	1:43:A:GLY:HA3	1:30:A:ILE:HG22	11	0.31
(1,956)	1:24:A:VAL:HG21	1:13:A:TYR:HD2	15	0.3
(1,954)	1:44:A:CYS:HA	1:45:A:ILE:HD12	14	0.3
(1,954)	1:44:A:CYS:HA	1:45:A:ILE:HD12	29	0.3
(1,953)	1:23:A:ASN:HB3	1:45:A:ILE:HG21	5	0.3
(1,952)	1:23:A:ASN:H	1:45:A:ILE:HD12	27	0.3
(1,943)	1:7:A:VAL:HB	1:7:A:VAL:HG13	3	0.3
(1,929)	1:13:A:TYR:HD1	1:11:A:LYS:HG2	26	0.3
(1,917)	1:25:A:LYS:HB3	1:25:A:LYS:H	18	0.3
(1,906)	1:34:A:GLU:HG2	1:3:A:HIS:H	28	0.3
(1,901)	1:14:A:CYS:HB2	1:13:A:TYR:HD2	6	0.3
(1,900)	1:14:A:CYS:HB3	1:24:A:VAL:H	4	0.3
(1,894)	1:23:A:ASN:H	1:54:A:ARG:HD3	13	0.3
(1,893)	1:41:A:TRP:HZ3	1:61:A:SER:HB2	21	0.3
(1,882)	1:50:A:THR:H	1:49:A:ARG:HA	3	0.3
(1,882)	1:50:A:THR:H	1:49:A:ARG:HA	4	0.3
(1,882)	1:50:A:THR:H	1:49:A:ARG:HA	7	0.3
(1,882)	1:50:A:THR:H	1:49:A:ARG:HA	9	0.3
(1,882)	1:50:A:THR:H	1:49:A:ARG:HA	11	0.3
(1,882)	1:50:A:THR:H	1:49:A:ARG:HA	13	0.3

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,882)	1:50:A:THR:H	1:49:A:ARG:HA	15	0.3
(1,882)	1:50:A:THR:H	1:49:A:ARG:HA	19	0.3
(1,882)	1:50:A:THR:H	1:49:A:ARG:HA	20	0.3
(1,882)	1:50:A:THR:H	1:49:A:ARG:HA	30	0.3
(1,859)	1:67:A:VAL:H	1:7:A:VAL:HA	1	0.3
(1,859)	1:67:A:VAL:H	1:7:A:VAL:HA	17	0.3
(1,849)	1:13:A:TYR:H	1:15:A:ASN:HD22	18	0.3
(1,823)	1:28:A:ASP:H	1:25:A:LYS:HG3	10	0.3
(1,819)	1:58:A:PHE:HD1	1:42:A:LYS:H	23	0.3
(1,815)	1:44:A:CYS:H	1:30:A:ILE:HG22	18	0.3
(1,810)	1:23:A:ASN:H	1:45:A:ILE:H	6	0.3
(1,805)	1:30:A:ILE:H	1:30:A:ILE:HD11	28	0.3
(1,804)	1:22:A:LEU:HB3	1:22:A:LEU:H	4	0.3
(1,804)	1:22:A:LEU:HB3	1:22:A:LEU:H	9	0.3
(1,804)	1:22:A:LEU:HB3	1:22:A:LEU:H	10	0.3
(1,804)	1:22:A:LEU:HB3	1:22:A:LEU:H	26	0.3
(1,804)	1:22:A:LEU:HB3	1:22:A:LEU:H	28	0.3
(1,803)	1:30:A:ILE:HG23	1:32:A:VAL:H	24	0.3
(1,667)	1:43:A:GLY:H	1:44:A:CYS:H	16	0.3
(1,667)	1:43:A:GLY:H	1:44:A:CYS:H	24	0.3
(1,667)	1:43:A:GLY:H	1:44:A:CYS:H	27	0.3
(1,667)	1:43:A:GLY:H	1:44:A:CYS:H	29	0.3
(1,667)	1:43:A:GLY:H	1:44:A:CYS:H	30	0.3
(1,654)	1:17:A:TYR:H	1:17:A:TYR:HB3	5	0.3
(1,654)	1:17:A:TYR:H	1:17:A:TYR:HB3	7	0.3
(1,654)	1:17:A:TYR:H	1:17:A:TYR:HB3	14	0.3
(1,654)	1:17:A:TYR:H	1:17:A:TYR:HB3	15	0.3
(1,654)	1:17:A:TYR:H	1:17:A:TYR:HB3	16	0.3
(1,654)	1:17:A:TYR:H	1:17:A:TYR:HB3	18	0.3
(1,654)	1:17:A:TYR:H	1:17:A:TYR:HB3	20	0.3
(1,654)	1:17:A:TYR:H	1:17:A:TYR:HB3	26	0.3
(1,654)	1:17:A:TYR:H	1:17:A:TYR:HB3	27	0.3
(1,628)	1:38:A:ASP:H	1:36:A:HIS:HB2	11	0.3
(1,628)	1:38:A:ASP:H	1:36:A:HIS:HB2	29	0.3
(1,587)	1:53:A:ASP:H	1:53:A:ASP:HB2	1	0.3
(1,583)	1:55:A:VAL:H	1:55:A:VAL:HB	26	0.3
(1,583)	1:55:A:VAL:H	1:55:A:VAL:HB	29	0.3
(1,541)	1:40:A:ARG:H	1:40:A:ARG:HD2	17	0.3
(1,525)	1:15:A:ASN:H	1:14:A:CYS:HA	23	0.3
(1,427)	1:20:A:THR:HG23	1:55:A:VAL:HG21	6	0.3
(1,426)	1:20:A:THR:HG22	1:55:A:VAL:HG11	29	0.3
(1,423)	1:7:A:VAL:HG22	1:32:A:VAL:HG22	9	0.3

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,422)	1:7:A:VAL:HG21	1:32:A:VAL:HG12	4	0.3
(1,399)	1:29:A:ILE:H	1:29:A:ILE:HD13	20	0.3
(1,393)	1:33:A:LEU:HD21	1:33:A:LEU:HD12	29	0.3
(1,382)	1:65:A:ALA:H	1:5:A:LEU:HB3	21	0.3
(1,372)	1:33:A:LEU:HD22	1:33:A:LEU:HG	29	0.3
(1,367)	1:9:A:ALA:HB3	1:10:A:THR:H	30	0.3
(1,358)	1:41:A:TRP:H	1:40:A:ARG:HB2	2	0.3
(1,355)	1:33:A:LEU:HB3	1:33:A:LEU:H	3	0.3
(1,347)	1:63:A:GLY:H	1:62:A:LEU:HB3	8	0.3
(1,330)	1:63:A:GLY:H	1:62:A:LEU:HB2	5	0.3
(1,314)	1:60:A:SER:HB2	1:60:A:SER:HA	19	0.3
(1,279)	1:63:A:GLY:H	1:62:A:LEU:HA	4	0.3
(1,268)	1:30:A:ILE:HG13	1:30:A:ILE:HA	23	0.3
(1,268)	1:30:A:ILE:HG13	1:30:A:ILE:HA	27	0.3
(1,200)	1:14:A:CYS:H	1:13:A:TYR:HD1	17	0.3
(1,191)	1:6:A:GLN:H	1:6:A:GLN:HG2	3	0.3
(1,191)	1:6:A:GLN:H	1:6:A:GLN:HG2	9	0.3
(1,191)	1:6:A:GLN:H	1:6:A:GLN:HG2	14	0.3
(1,191)	1:6:A:GLN:H	1:6:A:GLN:HG2	18	0.3
(1,176)	1:33:A:LEU:H	1:42:A:LYS:H	12	0.3
(1,176)	1:33:A:LEU:H	1:42:A:LYS:H	20	0.3
(1,122)	1:12:A:ASP:H	1:26:A:ALA:HB2	26	0.3
(1,107)	1:41:A:TRP:HB2	1:41:A:TRP:HD1	7	0.3
(1,107)	1:41:A:TRP:HB2	1:41:A:TRP:HD1	22	0.3
(1,69)	1:52:A:ASN:H	1:52:A:ASN:HB2	17	0.3
(1,39)	1:53:A:ASP:HB3	1:54:A:ARG:H	10	0.3
(1,29)	1:43:A:GLY:HA3	1:30:A:ILE:HG22	15	0.3
(1,29)	1:43:A:GLY:HA3	1:30:A:ILE:HG23	27	0.3
(1,14)	1:27:A:GLY:H	1:26:A:ALA:HA	2	0.3
(1,14)	1:27:A:GLY:H	1:26:A:ALA:HA	28	0.3
(1,958)	1:61:A:SER:H	1:5:A:LEU:HD23	6	0.29
(1,948)	1:6:A:GLN:H	1:67:A:VAL:HG21	25	0.29
(1,944)	1:30:A:ILE:HD12	1:7:A:VAL:HG22	22	0.29
(1,940)	1:7:A:VAL:H	1:32:A:VAL:HG12	19	0.29
(1,919)	1:23:A:ASN:HD21	1:54:A:ARG:HB2	11	0.29
(1,901)	1:14:A:CYS:HB2	1:13:A:TYR:HD2	9	0.29
(1,884)	1:62:A:LEU:H	1:62:A:LEU:HA	24	0.29
(1,882)	1:50:A:THR:H	1:49:A:ARG:HA	1	0.29
(1,882)	1:50:A:THR:H	1:49:A:ARG:HA	6	0.29
(1,882)	1:50:A:THR:H	1:49:A:ARG:HA	10	0.29
(1,882)	1:50:A:THR:H	1:49:A:ARG:HA	16	0.29
(1,882)	1:50:A:THR:H	1:49:A:ARG:HA	18	0.29

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,882)	1:50:A:THR:H	1:49:A:ARG:HA	22	0.29
(1,882)	1:50:A:THR:H	1:49:A:ARG:HA	25	0.29
(1,882)	1:50:A:THR:H	1:49:A:ARG:HA	26	0.29
(1,882)	1:50:A:THR:H	1:49:A:ARG:HA	27	0.29
(1,881)	1:13:A:TYR:HD2	1:21:A:SER:HB3	26	0.29
(1,873)	1:41:A:TRP:HB3	1:34:A:GLU:HA	3	0.29
(1,873)	1:41:A:TRP:HB3	1:34:A:GLU:HA	7	0.29
(1,870)	1:29:A:ILE:HG23	1:45:A:ILE:HA	13	0.29
(1,863)	1:9:A:ALA:H	1:29:A:ILE:HA	14	0.29
(1,859)	1:67:A:VAL:H	1:7:A:VAL:HA	2	0.29
(1,852)	1:13:A:TYR:HE2	1:24:A:VAL:H	1	0.29
(1,825)	1:19:A:LEU:HB3	1:20:A:THR:H	9	0.29
(1,816)	1:5:A:LEU:HB2	1:7:A:VAL:H	8	0.29
(1,816)	1:5:A:LEU:HB2	1:7:A:VAL:H	10	0.29
(1,816)	1:5:A:LEU:HB2	1:7:A:VAL:H	17	0.29
(1,815)	1:44:A:CYS:H	1:30:A:ILE:HG23	24	0.29
(1,810)	1:23:A:ASN:H	1:45:A:ILE:H	7	0.29
(1,810)	1:23:A:ASN:H	1:45:A:ILE:H	20	0.29
(1,805)	1:30:A:ILE:H	1:30:A:ILE:HD11	20	0.29
(1,805)	1:30:A:ILE:H	1:30:A:ILE:HD12	30	0.29
(1,804)	1:22:A:LEU:HB3	1:22:A:LEU:H	2	0.29
(1,804)	1:22:A:LEU:HB3	1:22:A:LEU:H	6	0.29
(1,804)	1:22:A:LEU:HB3	1:22:A:LEU:H	7	0.29
(1,804)	1:22:A:LEU:HB3	1:22:A:LEU:H	12	0.29
(1,804)	1:22:A:LEU:HB3	1:22:A:LEU:H	13	0.29
(1,804)	1:22:A:LEU:HB3	1:22:A:LEU:H	14	0.29
(1,804)	1:22:A:LEU:HB3	1:22:A:LEU:H	15	0.29
(1,804)	1:22:A:LEU:HB3	1:22:A:LEU:H	16	0.29
(1,804)	1:22:A:LEU:HB3	1:22:A:LEU:H	21	0.29
(1,804)	1:22:A:LEU:HB3	1:22:A:LEU:H	23	0.29
(1,804)	1:22:A:LEU:HB3	1:22:A:LEU:H	24	0.29
(1,804)	1:22:A:LEU:HB3	1:22:A:LEU:H	27	0.29
(1,804)	1:22:A:LEU:HB3	1:22:A:LEU:H	29	0.29
(1,775)	1:38:A:ASP:H	1:40:A:ARG:H	25	0.29
(1,714)	1:15:A:ASN:HD22	1:15:A:ASN:HA	8	0.29
(1,685)	1:27:A:GLY:H	1:24:A:VAL:HG22	6	0.29
(1,685)	1:27:A:GLY:H	1:24:A:VAL:HG22	7	0.29
(1,670)	1:43:A:GLY:H	1:42:A:LYS:HB2	30	0.29
(1,667)	1:43:A:GLY:H	1:44:A:CYS:H	22	0.29
(1,660)	1:10:A:THR:H	1:11:A:LYS:HB2	17	0.29
(1,654)	1:17:A:TYR:H	1:17:A:TYR:HB3	2	0.29
(1,654)	1:17:A:TYR:H	1:17:A:TYR:HB3	6	0.29

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,654)	1:17:A:TYR:H	1:17:A:TYR:HB3	23	0.29
(1,583)	1:55:A:VAL:H	1:55:A:VAL:HB	14	0.29
(1,583)	1:55:A:VAL:H	1:55:A:VAL:HB	23	0.29
(1,574)	1:36:A:HIS:H	1:35:A:GLN:HB2	21	0.29
(1,519)	1:14:A:CYS:H	1:13:A:TYR:HD1	20	0.29
(1,512)	1:65:A:ALA:H	1:66:A:ILE:HG21	24	0.29
(1,480)	1:45:A:ILE:H	1:44:A:CYS:HB2	1	0.29
(1,480)	1:45:A:ILE:H	1:44:A:CYS:HB2	15	0.29
(1,461)	1:67:A:VAL:H	1:67:A:VAL:HA	4	0.29
(1,461)	1:67:A:VAL:H	1:67:A:VAL:HA	24	0.29
(1,434)	1:24:A:VAL:HG12	1:30:A:ILE:HD11	8	0.29
(1,422)	1:7:A:VAL:HG23	1:32:A:VAL:HG12	27	0.29
(1,403)	1:58:A:PHE:H	1:22:A:LEU:HD12	18	0.29
(1,393)	1:33:A:LEU:HD23	1:33:A:LEU:HD12	4	0.29
(1,393)	1:33:A:LEU:HD21	1:33:A:LEU:HD11	28	0.29
(1,376)	1:32:A:VAL:HG22	1:32:A:VAL:HG12	11	0.29
(1,376)	1:32:A:VAL:HG23	1:32:A:VAL:HG13	20	0.29
(1,355)	1:33:A:LEU:HB3	1:33:A:LEU:H	26	0.29
(1,330)	1:63:A:GLY:H	1:62:A:LEU:HB2	13	0.29
(1,330)	1:63:A:GLY:H	1:62:A:LEU:HB2	20	0.29
(1,314)	1:60:A:SER:HB2	1:60:A:SER:HA	16	0.29
(1,289)	1:32:A:VAL:HA	1:32:A:VAL:HG23	5	0.29
(1,289)	1:32:A:VAL:HA	1:32:A:VAL:HG23	8	0.29
(1,289)	1:32:A:VAL:HA	1:32:A:VAL:HG23	19	0.29
(1,289)	1:32:A:VAL:HA	1:32:A:VAL:HG23	23	0.29
(1,289)	1:32:A:VAL:HA	1:32:A:VAL:HG23	24	0.29
(1,289)	1:32:A:VAL:HA	1:32:A:VAL:HG23	30	0.29
(1,279)	1:63:A:GLY:H	1:62:A:LEU:HA	2	0.29
(1,279)	1:63:A:GLY:H	1:62:A:LEU:HA	27	0.29
(1,234)	1:8:A:ARG:HG3	1:6:A:GLN:HE22	6	0.29
(1,228)	1:51:A:GLY:H	1:52:A:ASN:HD22	20	0.29
(1,226)	1:39:A:GLY:HA3	1:41:A:TRP:HD1	2	0.29
(1,220)	1:15:A:ASN:H	1:15:A:ASN:HD22	12	0.29
(1,203)	1:52:A:ASN:HD22	1:52:A:ASN:H	27	0.29
(1,191)	1:6:A:GLN:H	1:6:A:GLN:HG2	7	0.29
(1,107)	1:41:A:TRP:HB2	1:41:A:TRP:HD1	30	0.29
(1,106)	1:41:A:TRP:HB3	1:41:A:TRP:HZ3	19	0.29
(1,85)	1:7:A:VAL:HB	1:7:A:VAL:HG13	9	0.29
(1,69)	1:52:A:ASN:H	1:52:A:ASN:HB2	12	0.29
(1,60)	1:28:A:ASP:HB2	1:28:A:ASP:HA	22	0.29
(1,60)	1:28:A:ASP:HB2	1:28:A:ASP:HA	25	0.29
(1,58)	1:28:A:ASP:H	1:28:A:ASP:HB2	22	0.29

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,29)	1:43:A:GLY:HA3	1:30:A:ILE:HG22	13	0.29
(1,7)	1:61:A:SER:HA	1:62:A:LEU:H	3	0.29
(1,964)	1:45:A:ILE:H	1:55:A:VAL:HA	25	0.28
(1,964)	1:45:A:ILE:H	1:55:A:VAL:HA	29	0.28
(1,956)	1:24:A:VAL:HG23	1:13:A:TYR:HD2	17	0.28
(1,954)	1:44:A:CYS:HA	1:45:A:ILE:HD11	5	0.28
(1,943)	1:7:A:VAL:HB	1:7:A:VAL:HG13	9	0.28
(1,900)	1:14:A:CYS:HB3	1:24:A:VAL:H	3	0.28
(1,900)	1:14:A:CYS:HB3	1:24:A:VAL:H	12	0.28
(1,882)	1:50:A:THR:H	1:49:A:ARG:HA	2	0.28
(1,882)	1:50:A:THR:H	1:49:A:ARG:HA	5	0.28
(1,882)	1:50:A:THR:H	1:49:A:ARG:HA	8	0.28
(1,882)	1:50:A:THR:H	1:49:A:ARG:HA	12	0.28
(1,882)	1:50:A:THR:H	1:49:A:ARG:HA	21	0.28
(1,882)	1:50:A:THR:H	1:49:A:ARG:HA	23	0.28
(1,882)	1:50:A:THR:H	1:49:A:ARG:HA	28	0.28
(1,882)	1:50:A:THR:H	1:49:A:ARG:HA	29	0.28
(1,858)	1:13:A:TYR:HD2	1:23:A:ASN:HA	21	0.28
(1,852)	1:25:A:LYS:H	1:48:A:ASN:HD22	24	0.28
(1,836)	1:40:A:ARG:H	1:41:A:TRP:HD1	3	0.28
(1,819)	1:58:A:PHE:HD1	1:42:A:LYS:H	12	0.28
(1,815)	1:44:A:CYS:H	1:30:A:ILE:HG23	8	0.28
(1,815)	1:44:A:CYS:H	1:30:A:ILE:HG21	16	0.28
(1,805)	1:30:A:ILE:H	1:30:A:ILE:HD11	5	0.28
(1,805)	1:30:A:ILE:H	1:30:A:ILE:HD13	8	0.28
(1,805)	1:30:A:ILE:H	1:30:A:ILE:HD12	16	0.28
(1,805)	1:30:A:ILE:H	1:30:A:ILE:HD12	17	0.28
(1,805)	1:30:A:ILE:H	1:30:A:ILE:HD11	27	0.28
(1,804)	1:22:A:LEU:HB3	1:22:A:LEU:H	1	0.28
(1,804)	1:22:A:LEU:HB3	1:22:A:LEU:H	5	0.28
(1,667)	1:43:A:GLY:H	1:44:A:CYS:H	10	0.28
(1,667)	1:43:A:GLY:H	1:44:A:CYS:H	23	0.28
(1,667)	1:43:A:GLY:H	1:44:A:CYS:H	26	0.28
(1,653)	1:17:A:TYR:H	1:16:A:ASN:HB2	30	0.28
(1,620)	1:21:A:SER:H	1:14:A:CYS:HB2	9	0.28
(1,598)	1:29:A:ILE:H	1:28:A:ASP:HA	21	0.28
(1,560)	1:64:A:GLU:H	1:64:A:GLU:HG2	1	0.28
(1,466)	1:30:A:ILE:H	1:29:A:ILE:HB	2	0.28
(1,461)	1:67:A:VAL:H	1:67:A:VAL:HA	27	0.28
(1,450)	1:25:A:LYS:H	1:24:A:VAL:HG12	15	0.28
(1,423)	1:7:A:VAL:HG22	1:32:A:VAL:HG21	25	0.28
(1,410)	1:22:A:LEU:HD13	1:30:A:ILE:HD12	8	0.28

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,395)	1:42:A:LYS:H	1:33:A:LEU:HD11	23	0.28
(1,393)	1:33:A:LEU:HD23	1:33:A:LEU:HD13	5	0.28
(1,393)	1:33:A:LEU:HD22	1:33:A:LEU:HD13	6	0.28
(1,393)	1:33:A:LEU:HD21	1:33:A:LEU:HD12	20	0.28
(1,393)	1:33:A:LEU:HD23	1:33:A:LEU:HD13	25	0.28
(1,392)	1:43:A:GLY:H	1:33:A:LEU:HD13	13	0.28
(1,382)	1:65:A:ALA:H	1:5:A:LEU:HB3	11	0.28
(1,376)	1:32:A:VAL:HG23	1:32:A:VAL:HG11	4	0.28
(1,376)	1:32:A:VAL:HG21	1:32:A:VAL:HG13	10	0.28
(1,358)	1:41:A:TRP:H	1:40:A:ARG:HB2	18	0.28
(1,355)	1:33:A:LEU:HB3	1:33:A:LEU:H	2	0.28
(1,355)	1:33:A:LEU:HB3	1:33:A:LEU:H	17	0.28
(1,334)	1:49:A:ARG:H	1:49:A:ARG:HB2	22	0.28
(1,289)	1:32:A:VAL:HA	1:32:A:VAL:HG21	4	0.28
(1,289)	1:32:A:VAL:HA	1:32:A:VAL:HG21	13	0.28
(1,289)	1:32:A:VAL:HA	1:32:A:VAL:HG22	14	0.28
(1,279)	1:63:A:GLY:H	1:62:A:LEU:HA	1	0.28
(1,279)	1:63:A:GLY:H	1:62:A:LEU:HA	3	0.28
(1,279)	1:63:A:GLY:H	1:62:A:LEU:HA	11	0.28
(1,279)	1:63:A:GLY:H	1:62:A:LEU:HA	21	0.28
(1,279)	1:63:A:GLY:H	1:62:A:LEU:HA	22	0.28
(1,279)	1:63:A:GLY:H	1:62:A:LEU:HA	23	0.28
(1,279)	1:63:A:GLY:H	1:62:A:LEU:HA	24	0.28
(1,279)	1:63:A:GLY:H	1:62:A:LEU:HA	26	0.28
(1,268)	1:30:A:ILE:HG13	1:30:A:ILE:HA	19	0.28
(1,220)	1:15:A:ASN:H	1:15:A:ASN:HD22	25	0.28
(1,220)	1:15:A:ASN:H	1:15:A:ASN:HD22	27	0.28
(1,203)	1:52:A:ASN:HD22	1:52:A:ASN:H	4	0.28
(1,191)	1:6:A:GLN:H	1:6:A:GLN:HG2	22	0.28
(1,62)	1:18:A:ASP:HB2	1:18:A:ASP:H	16	0.28
(1,60)	1:28:A:ASP:HB2	1:28:A:ASP:HA	26	0.28
(1,56)	1:28:A:ASP:H	1:28:A:ASP:HB3	2	0.28
(1,55)	1:47:A:ASP:HB3	1:52:A:ASN:H	22	0.28
(1,39)	1:53:A:ASP:HB3	1:54:A:ARG:H	29	0.28
(1,968)	1:10:A:THR:H	1:63:A:GLY:HA3	2	0.27
(1,968)	1:10:A:THR:H	1:26:A:ALA:HA	8	0.27
(1,968)	1:10:A:THR:H	1:63:A:GLY:HA3	29	0.27
(1,964)	1:45:A:ILE:H	1:55:A:VAL:HA	4	0.27
(1,964)	1:45:A:ILE:H	1:55:A:VAL:HA	26	0.27
(1,909)	1:41:A:TRP:HD1	1:40:A:ARG:HG2	22	0.27
(1,902)	1:38:A:ASP:H	1:38:A:ASP:HB3	12	0.27
(1,899)	1:27:A:GLY:H	1:28:A:ASP:HB2	27	0.27

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,893)	1:41:A:TRP:HZ3	1:61:A:SER:HB2	14	0.27
(1,892)	1:41:A:TRP:HH2	1:61:A:SER:HA	14	0.27
(1,882)	1:50:A:THR:H	1:49:A:ARG:HA	14	0.27
(1,859)	1:67:A:VAL:H	1:7:A:VAL:HA	23	0.27
(1,858)	1:13:A:TYR:HD2	1:23:A:ASN:HA	27	0.27
(1,825)	1:34:A:GLU:HB2	1:3:A:HIS:H	22	0.27
(1,805)	1:30:A:ILE:H	1:30:A:ILE:HD13	21	0.27
(1,799)	1:41:A:TRP:H	1:41:A:TRP:HB3	9	0.27
(1,791)	1:39:A:GLY:H	1:39:A:GLY:HA3	2	0.27
(1,791)	1:39:A:GLY:H	1:39:A:GLY:HA3	4	0.27
(1,791)	1:39:A:GLY:H	1:39:A:GLY:HA3	9	0.27
(1,791)	1:39:A:GLY:H	1:39:A:GLY:HA3	10	0.27
(1,791)	1:39:A:GLY:H	1:39:A:GLY:HA3	14	0.27
(1,791)	1:39:A:GLY:H	1:39:A:GLY:HA3	16	0.27
(1,791)	1:39:A:GLY:H	1:39:A:GLY:HA3	21	0.27
(1,791)	1:39:A:GLY:H	1:39:A:GLY:HA3	23	0.27
(1,754)	1:10:A:THR:H	1:27:A:GLY:H	4	0.27
(1,743)	1:63:A:GLY:H	1:62:A:LEU:HG	15	0.27
(1,738)	1:50:A:THR:H	1:50:A:THR:HB	3	0.27
(1,738)	1:50:A:THR:H	1:50:A:THR:HB	10	0.27
(1,685)	1:27:A:GLY:H	1:24:A:VAL:HG21	12	0.27
(1,667)	1:43:A:GLY:H	1:44:A:CYS:H	5	0.27
(1,627)	1:38:A:ASP:H	1:37:A:PRO:HB3	26	0.27
(1,583)	1:55:A:VAL:H	1:55:A:VAL:HB	10	0.27
(1,574)	1:36:A:HIS:H	1:35:A:GLN:HB2	18	0.27
(1,480)	1:45:A:ILE:H	1:44:A:CYS:HB2	5	0.27
(1,467)	1:30:A:ILE:H	1:29:A:ILE:HG22	1	0.27
(1,461)	1:67:A:VAL:H	1:67:A:VAL:HA	16	0.27
(1,423)	1:7:A:VAL:HG21	1:32:A:VAL:HG23	12	0.27
(1,393)	1:33:A:LEU:HD23	1:33:A:LEU:HD12	2	0.27
(1,392)	1:43:A:GLY:H	1:33:A:LEU:HD11	9	0.27
(1,376)	1:32:A:VAL:HG23	1:32:A:VAL:HG12	13	0.27
(1,376)	1:32:A:VAL:HG23	1:32:A:VAL:HG12	15	0.27
(1,376)	1:32:A:VAL:HG21	1:32:A:VAL:HG11	17	0.27
(1,367)	1:9:A:ALA:HB1	1:10:A:THR:H	7	0.27
(1,358)	1:41:A:TRP:H	1:40:A:ARG:HB3	29	0.27
(1,299)	1:50:A:THR:H	1:47:A:ASP:HB2	19	0.27
(1,289)	1:32:A:VAL:HA	1:32:A:VAL:HG23	2	0.27
(1,289)	1:32:A:VAL:HA	1:32:A:VAL:HG22	3	0.27
(1,289)	1:32:A:VAL:HA	1:32:A:VAL:HG21	9	0.27
(1,289)	1:32:A:VAL:HA	1:32:A:VAL:HG22	12	0.27
(1,289)	1:32:A:VAL:HA	1:32:A:VAL:HG21	16	0.27

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,289)	1:32:A:VAL:HA	1:32:A:VAL:HG23	22	0.27
(1,289)	1:32:A:VAL:HA	1:32:A:VAL:HG23	25	0.27
(1,289)	1:32:A:VAL:HA	1:32:A:VAL:HG23	26	0.27
(1,289)	1:32:A:VAL:HA	1:32:A:VAL:HG22	28	0.27
(1,279)	1:63:A:GLY:H	1:62:A:LEU:HA	15	0.27
(1,279)	1:63:A:GLY:H	1:62:A:LEU:HA	19	0.27
(1,279)	1:63:A:GLY:H	1:62:A:LEU:HA	20	0.27
(1,279)	1:63:A:GLY:H	1:62:A:LEU:HA	25	0.27
(1,279)	1:63:A:GLY:H	1:62:A:LEU:HA	29	0.27
(1,268)	1:30:A:ILE:HG13	1:30:A:ILE:HA	4	0.27
(1,268)	1:30:A:ILE:HG13	1:30:A:ILE:HA	20	0.27
(1,220)	1:15:A:ASN:H	1:15:A:ASN:HD22	23	0.27
(1,191)	1:6:A:GLN:H	1:6:A:GLN:HG2	12	0.27
(1,191)	1:6:A:GLN:H	1:6:A:GLN:HG2	13	0.27
(1,191)	1:6:A:GLN:H	1:6:A:GLN:HG2	17	0.27
(1,107)	1:41:A:TRP:HB2	1:41:A:TRP:HD1	27	0.27
(1,106)	1:41:A:TRP:HB3	1:41:A:TRP:HZ3	9	0.27
(1,71)	1:38:A:ASP:HB2	1:38:A:ASP:HA	21	0.27
(1,66)	1:28:A:ASP:HB3	1:25:A:LYS:HG2	21	0.27
(1,60)	1:28:A:ASP:HB2	1:28:A:ASP:HA	13	0.27
(1,35)	1:27:A:GLY:HA3	1:28:A:ASP:H	8	0.27
(1,35)	1:27:A:GLY:HA3	1:28:A:ASP:H	13	0.27
(1,35)	1:27:A:GLY:HA3	1:28:A:ASP:H	14	0.27
(1,966)	1:8:A:ARG:H	1:64:A:GLU:HG3	27	0.26
(1,952)	1:23:A:ASN:H	1:45:A:ILE:HD11	28	0.26
(1,944)	1:30:A:ILE:HD13	1:7:A:VAL:HG22	15	0.26
(1,944)	1:30:A:ILE:HD13	1:7:A:VAL:HG21	21	0.26
(1,912)	1:41:A:TRP:H	1:40:A:ARG:HG3	8	0.26
(1,908)	1:34:A:GLU:H	1:35:A:GLN:HG3	7	0.26
(1,908)	1:34:A:GLU:H	1:35:A:GLN:HG2	17	0.26
(1,882)	1:50:A:THR:H	1:49:A:ARG:HA	17	0.26
(1,873)	1:41:A:TRP:HB3	1:34:A:GLU:HA	17	0.26
(1,816)	1:5:A:LEU:HB2	1:7:A:VAL:H	11	0.26
(1,810)	1:23:A:ASN:H	1:45:A:ILE:H	15	0.26
(1,805)	1:30:A:ILE:H	1:30:A:ILE:HD11	6	0.26
(1,804)	1:22:A:LEU:HB3	1:22:A:LEU:H	3	0.26
(1,799)	1:41:A:TRP:H	1:41:A:TRP:HB3	21	0.26
(1,798)	1:24:A:VAL:HB	1:12:A:ASP:HA	12	0.26
(1,791)	1:39:A:GLY:H	1:39:A:GLY:HA3	5	0.26
(1,791)	1:39:A:GLY:H	1:39:A:GLY:HA3	18	0.26
(1,791)	1:39:A:GLY:H	1:39:A:GLY:HA3	19	0.26
(1,791)	1:39:A:GLY:H	1:39:A:GLY:HA3	20	0.26

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,791)	1:39:A:GLY:H	1:39:A:GLY:HA3	29	0.26
(1,791)	1:39:A:GLY:H	1:39:A:GLY:HA3	30	0.26
(1,775)	1:38:A:ASP:H	1:40:A:ARG:H	8	0.26
(1,727)	1:39:A:GLY:H	1:40:A:ARG:HG2	28	0.26
(1,714)	1:15:A:ASN:HD22	1:15:A:ASN:HA	10	0.26
(1,714)	1:15:A:ASN:HD22	1:15:A:ASN:HA	20	0.26
(1,691)	1:16:A:ASN:HD21	1:16:A:ASN:HA	30	0.26
(1,685)	1:27:A:GLY:H	1:24:A:VAL:HG23	22	0.26
(1,667)	1:43:A:GLY:H	1:44:A:CYS:H	11	0.26
(1,667)	1:43:A:GLY:H	1:44:A:CYS:H	14	0.26
(1,667)	1:43:A:GLY:H	1:44:A:CYS:H	28	0.26
(1,627)	1:38:A:ASP:H	1:37:A:PRO:HB3	28	0.26
(1,583)	1:55:A:VAL:H	1:55:A:VAL:HB	4	0.26
(1,536)	1:23:A:ASN:H	1:22:A:LEU:HD12	22	0.26
(1,519)	1:14:A:CYS:H	1:13:A:TYR:HD1	17	0.26
(1,480)	1:45:A:ILE:H	1:44:A:CYS:HB2	22	0.26
(1,467)	1:30:A:ILE:H	1:29:A:ILE:HG21	20	0.26
(1,441)	1:30:A:ILE:HG23	1:45:A:ILE:HG23	10	0.26
(1,423)	1:7:A:VAL:HG21	1:32:A:VAL:HG22	13	0.26
(1,395)	1:42:A:LYS:H	1:33:A:LEU:HD12	29	0.26
(1,395)	1:42:A:LYS:H	1:33:A:LEU:HD11	30	0.26
(1,393)	1:33:A:LEU:HD21	1:33:A:LEU:HD12	14	0.26
(1,376)	1:32:A:VAL:HG22	1:32:A:VAL:HG13	5	0.26
(1,376)	1:32:A:VAL:HG22	1:32:A:VAL:HG11	21	0.26
(1,376)	1:32:A:VAL:HG22	1:32:A:VAL:HG11	22	0.26
(1,376)	1:32:A:VAL:HG22	1:32:A:VAL:HG12	23	0.26
(1,376)	1:32:A:VAL:HG22	1:32:A:VAL:HG11	24	0.26
(1,376)	1:32:A:VAL:HG22	1:32:A:VAL:HG11	27	0.26
(1,376)	1:32:A:VAL:HG22	1:32:A:VAL:HG12	29	0.26
(1,376)	1:32:A:VAL:HG22	1:32:A:VAL:HG13	30	0.26
(1,367)	1:9:A:ALA:HB2	1:10:A:THR:H	8	0.26
(1,358)	1:41:A:TRP:H	1:40:A:ARG:HB2	30	0.26
(1,355)	1:33:A:LEU:HB3	1:33:A:LEU:H	16	0.26
(1,355)	1:33:A:LEU:HB3	1:33:A:LEU:H	22	0.26
(1,347)	1:63:A:GLY:H	1:62:A:LEU:HB3	9	0.26
(1,317)	1:39:A:GLY:H	1:38:A:ASP:HB3	1	0.26
(1,289)	1:32:A:VAL:HA	1:32:A:VAL:HG21	1	0.26
(1,289)	1:32:A:VAL:HA	1:32:A:VAL:HG21	6	0.26
(1,289)	1:32:A:VAL:HA	1:32:A:VAL:HG22	7	0.26
(1,289)	1:32:A:VAL:HA	1:32:A:VAL:HG22	10	0.26
(1,289)	1:32:A:VAL:HA	1:32:A:VAL:HG21	15	0.26
(1,289)	1:32:A:VAL:HA	1:32:A:VAL:HG22	17	0.26

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,289)	1:32:A:VAL:HA	1:32:A:VAL:HG23	27	0.26
(1,279)	1:63:A:GLY:H	1:62:A:LEU:HA	5	0.26
(1,279)	1:63:A:GLY:H	1:62:A:LEU:HA	10	0.26
(1,279)	1:63:A:GLY:H	1:62:A:LEU:HA	13	0.26
(1,279)	1:63:A:GLY:H	1:62:A:LEU:HA	28	0.26
(1,272)	1:17:A:TYR:HD2	1:16:A:ASN:HA	21	0.26
(1,232)	1:32:A:VAL:HG12	1:41:A:TRP:HZ3	15	0.26
(1,226)	1:39:A:GLY:HA3	1:41:A:TRP:HD1	19	0.26
(1,220)	1:15:A:ASN:H	1:15:A:ASN:HD22	11	0.26
(1,203)	1:52:A:ASN:HD22	1:52:A:ASN:H	15	0.26
(1,123)	1:26:A:ALA:HB1	1:10:A:THR:HA	4	0.26
(1,107)	1:41:A:TRP:HB2	1:41:A:TRP:HD1	2	0.26
(1,106)	1:41:A:TRP:HB3	1:41:A:TRP:HZ3	18	0.26
(1,60)	1:28:A:ASP:HB2	1:28:A:ASP:HA	5	0.26
(1,60)	1:28:A:ASP:HB2	1:28:A:ASP:HA	12	0.26
(1,60)	1:28:A:ASP:HB2	1:28:A:ASP:HA	16	0.26
(1,57)	1:28:A:ASP:HB2	1:25:A:LYS:HG2	21	0.26
(1,55)	1:47:A:ASP:HB3	1:52:A:ASN:H	29	0.26
(1,35)	1:27:A:GLY:HA3	1:28:A:ASP:H	7	0.26
(1,35)	1:27:A:GLY:HA3	1:28:A:ASP:H	9	0.26
(1,35)	1:27:A:GLY:HA3	1:28:A:ASP:H	15	0.26
(1,35)	1:27:A:GLY:HA3	1:28:A:ASP:H	20	0.26
(1,35)	1:27:A:GLY:HA3	1:28:A:ASP:H	22	0.26
(1,35)	1:27:A:GLY:HA3	1:28:A:ASP:H	30	0.26
(1,29)	1:43:A:GLY:HA3	1:30:A:ILE:HG22	4	0.26
(1,14)	1:27:A:GLY:H	1:26:A:ALA:HA	7	0.26
(1,953)	1:23:A:ASN:HB3	1:45:A:ILE:HG23	24	0.25
(1,928)	1:60:A:SER:HB3	1:41:A:TRP:HH2	19	0.25
(1,924)	1:65:A:ALA:HB2	1:5:A:LEU:HD13	23	0.25
(1,917)	1:25:A:LYS:HB3	1:25:A:LYS:H	5	0.25
(1,894)	1:53:A:ASP:H	1:54:A:ARG:HD3	17	0.25
(1,878)	1:52:A:ASN:H	1:50:A:THR:HB	25	0.25
(1,870)	1:29:A:ILE:HG23	1:45:A:ILE:HA	21	0.25
(1,858)	1:13:A:TYR:HD2	1:23:A:ASN:HA	7	0.25
(1,858)	1:13:A:TYR:HD2	1:23:A:ASN:HA	29	0.25
(1,834)	1:51:A:GLY:HA3	1:47:A:ASP:H	26	0.25
(1,816)	1:5:A:LEU:HB2	1:7:A:VAL:H	25	0.25
(1,816)	1:5:A:LEU:HB2	1:7:A:VAL:H	26	0.25
(1,811)	1:11:A:LYS:HG2	1:13:A:TYR:H	22	0.25
(1,800)	1:44:A:CYS:HB3	1:55:A:VAL:HA	5	0.25
(1,800)	1:44:A:CYS:HB2	1:55:A:VAL:HA	12	0.25
(1,791)	1:39:A:GLY:H	1:39:A:GLY:HA3	22	0.25

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,733)	1:39:A:GLY:H	1:36:A:HIS:HB3	24	0.25
(1,714)	1:15:A:ASN:HD22	1:15:A:ASN:HA	12	0.25
(1,705)	1:35:A:GLN:HE22	1:33:A:LEU:HD23	22	0.25
(1,667)	1:43:A:GLY:H	1:44:A:CYS:H	2	0.25
(1,667)	1:43:A:GLY:H	1:44:A:CYS:H	4	0.25
(1,660)	1:10:A:THR:H	1:11:A:LYS:HB2	22	0.25
(1,593)	1:62:A:LEU:H	1:60:A:SER:HA	23	0.25
(1,584)	1:3:A:HIS:H	1:34:A:GLU:HB3	15	0.25
(1,574)	1:36:A:HIS:H	1:35:A:GLN:HB2	29	0.25
(1,560)	1:64:A:GLU:H	1:64:A:GLU:HG2	26	0.25
(1,549)	1:54:A:ARG:H	1:54:A:ARG:HD2	4	0.25
(1,549)	1:54:A:ARG:H	1:54:A:ARG:HD2	15	0.25
(1,534)	1:23:A:ASN:H	1:23:A:ASN:HB2	30	0.25
(1,515)	1:14:A:CYS:H	1:13:A:TYR:HB3	26	0.25
(1,507)	1:22:A:LEU:H	1:22:A:LEU:HD13	30	0.25
(1,482)	1:46:A:HIS:H	1:30:A:ILE:HG13	15	0.25
(1,480)	1:45:A:ILE:H	1:44:A:CYS:HB2	7	0.25
(1,451)	1:25:A:LYS:H	1:24:A:VAL:HB	2	0.25
(1,450)	1:25:A:LYS:H	1:24:A:VAL:HG12	16	0.25
(1,434)	1:24:A:VAL:HG12	1:30:A:ILE:HD13	17	0.25
(1,431)	1:22:A:LEU:HD11	1:24:A:VAL:HG12	2	0.25
(1,423)	1:7:A:VAL:HG23	1:32:A:VAL:HG21	8	0.25
(1,422)	1:7:A:VAL:HG22	1:32:A:VAL:HG12	19	0.25
(1,414)	1:5:A:LEU:HD12	1:32:A:VAL:HG11	4	0.25
(1,393)	1:33:A:LEU:HD22	1:33:A:LEU:HD11	11	0.25
(1,393)	1:33:A:LEU:HD22	1:33:A:LEU:HD12	15	0.25
(1,393)	1:33:A:LEU:HD22	1:33:A:LEU:HD11	22	0.25
(1,393)	1:33:A:LEU:HD23	1:33:A:LEU:HD13	23	0.25
(1,376)	1:32:A:VAL:HG23	1:32:A:VAL:HG12	1	0.25
(1,376)	1:32:A:VAL:HG21	1:32:A:VAL:HG13	3	0.25
(1,376)	1:32:A:VAL:HG23	1:32:A:VAL:HG11	9	0.25
(1,376)	1:32:A:VAL:HG23	1:32:A:VAL:HG11	16	0.25
(1,376)	1:32:A:VAL:HG22	1:32:A:VAL:HG12	25	0.25
(1,376)	1:32:A:VAL:HG22	1:32:A:VAL:HG13	26	0.25
(1,330)	1:63:A:GLY:H	1:62:A:LEU:HB2	10	0.25
(1,330)	1:63:A:GLY:H	1:62:A:LEU:HB2	25	0.25
(1,289)	1:32:A:VAL:HA	1:32:A:VAL:HG21	11	0.25
(1,289)	1:32:A:VAL:HA	1:32:A:VAL:HG21	20	0.25
(1,289)	1:32:A:VAL:HA	1:32:A:VAL:HG23	29	0.25
(1,241)	1:4:A:MET:H	1:2:A:SER:HA	6	0.25
(1,232)	1:32:A:VAL:HG12	1:41:A:TRP:HZ3	1	0.25
(1,226)	1:39:A:GLY:HA3	1:41:A:TRP:HD1	8	0.25

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,220)	1:15:A:ASN:H	1:15:A:ASN:HD22	20	0.25
(1,215)	1:60:A:SER:HB2	1:41:A:TRP:HZ2	19	0.25
(1,196)	1:34:A:GLU:HB3	1:4:A:MET:H	15	0.25
(1,107)	1:41:A:TRP:HB2	1:41:A:TRP:HD1	26	0.25
(1,79)	1:17:A:TYR:HB2	1:17:A:TYR:HA	1	0.25
(1,79)	1:17:A:TYR:HB2	1:17:A:TYR:HA	4	0.25
(1,79)	1:17:A:TYR:HB2	1:17:A:TYR:HA	7	0.25
(1,79)	1:17:A:TYR:HB2	1:17:A:TYR:HA	13	0.25
(1,79)	1:17:A:TYR:HB2	1:17:A:TYR:HA	17	0.25
(1,79)	1:17:A:TYR:HB2	1:17:A:TYR:HA	18	0.25
(1,79)	1:17:A:TYR:HB2	1:17:A:TYR:HA	22	0.25
(1,79)	1:17:A:TYR:HB2	1:17:A:TYR:HA	24	0.25
(1,79)	1:17:A:TYR:HB2	1:17:A:TYR:HA	26	0.25
(1,79)	1:17:A:TYR:HB2	1:17:A:TYR:HA	28	0.25
(1,79)	1:17:A:TYR:HB2	1:17:A:TYR:HA	29	0.25
(1,79)	1:17:A:TYR:HB2	1:17:A:TYR:HA	30	0.25
(1,60)	1:28:A:ASP:HB2	1:28:A:ASP:HA	15	0.25
(1,60)	1:28:A:ASP:HB2	1:28:A:ASP:HA	24	0.25
(1,60)	1:28:A:ASP:HB2	1:28:A:ASP:HA	28	0.25
(1,58)	1:28:A:ASP:H	1:28:A:ASP:HB2	3	0.25
(1,55)	1:47:A:ASP:HB3	1:52:A:ASN:H	30	0.25
(1,39)	1:53:A:ASP:HB3	1:54:A:ARG:H	14	0.25
(1,35)	1:27:A:GLY:HA3	1:28:A:ASP:H	1	0.25
(1,35)	1:27:A:GLY:HA3	1:28:A:ASP:H	2	0.25
(1,35)	1:27:A:GLY:HA3	1:28:A:ASP:H	3	0.25
(1,35)	1:27:A:GLY:HA3	1:28:A:ASP:H	6	0.25
(1,35)	1:27:A:GLY:HA3	1:28:A:ASP:H	17	0.25
(1,35)	1:27:A:GLY:HA3	1:28:A:ASP:H	24	0.25
(1,35)	1:27:A:GLY:HA3	1:28:A:ASP:H	25	0.25
(1,29)	1:43:A:GLY:HA3	1:30:A:ILE:HG23	24	0.25
(1,14)	1:27:A:GLY:H	1:26:A:ALA:HA	12	0.25
(1,968)	1:10:A:THR:H	1:63:A:GLY:HA3	27	0.24
(1,965)	1:26:A:ALA:H	1:12:A:ASP:HA	26	0.24
(1,964)	1:45:A:ILE:H	1:55:A:VAL:HA	9	0.24
(1,964)	1:45:A:ILE:H	1:55:A:VAL:HA	28	0.24
(1,954)	1:44:A:CYS:HA	1:45:A:ILE:HD13	9	0.24
(1,940)	1:7:A:VAL:H	1:32:A:VAL:HG12	21	0.24
(1,924)	1:65:A:ALA:HB3	1:5:A:LEU:HD13	30	0.24
(1,882)	1:58:A:PHE:HD2	1:62:A:LEU:HA	24	0.24
(1,876)	1:45:A:ILE:HD11	1:30:A:ILE:HA	2	0.24
(1,872)	1:5:A:LEU:HD21	1:33:A:LEU:HA	13	0.24
(1,858)	1:13:A:TYR:HD2	1:21:A:SER:HA	14	0.24

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,816)	1:5:A:LEU:HB2	1:7:A:VAL:H	16	0.24
(1,816)	1:5:A:LEU:HB2	1:7:A:VAL:H	22	0.24
(1,805)	1:30:A:ILE:H	1:30:A:ILE:HD11	3	0.24
(1,805)	1:30:A:ILE:H	1:30:A:ILE:HD12	11	0.24
(1,803)	1:30:A:ILE:HG22	1:32:A:VAL:H	30	0.24
(1,799)	1:41:A:TRP:H	1:41:A:TRP:HB3	16	0.24
(1,799)	1:41:A:TRP:H	1:41:A:TRP:HB3	18	0.24
(1,799)	1:41:A:TRP:H	1:41:A:TRP:HB3	30	0.24
(1,798)	1:24:A:VAL:HB	1:12:A:ASP:HA	5	0.24
(1,791)	1:39:A:GLY:H	1:39:A:GLY:HA3	15	0.24
(1,655)	1:18:A:ASP:H	1:17:A:TYR:H	22	0.24
(1,649)	1:12:A:ASP:H	1:12:A:ASP:HA	2	0.24
(1,649)	1:12:A:ASP:H	1:12:A:ASP:HA	4	0.24
(1,560)	1:64:A:GLU:H	1:64:A:GLU:HG2	28	0.24
(1,559)	1:64:A:GLU:H	1:64:A:GLU:HA	19	0.24
(1,521)	1:5:A:LEU:H	1:5:A:LEU:HA	8	0.24
(1,521)	1:5:A:LEU:H	1:5:A:LEU:HA	14	0.24
(1,521)	1:5:A:LEU:H	1:5:A:LEU:HA	15	0.24
(1,521)	1:5:A:LEU:H	1:5:A:LEU:HA	22	0.24
(1,515)	1:14:A:CYS:H	1:13:A:TYR:HB3	5	0.24
(1,482)	1:46:A:HIS:H	1:30:A:ILE:HG13	12	0.24
(1,458)	1:9:A:ALA:H	1:9:A:ALA:HB3	20	0.24
(1,437)	1:24:A:VAL:HG21	1:30:A:ILE:HG21	25	0.24
(1,432)	1:22:A:LEU:HD13	1:24:A:VAL:HG22	8	0.24
(1,423)	1:7:A:VAL:HG22	1:32:A:VAL:HG22	6	0.24
(1,423)	1:7:A:VAL:HG21	1:32:A:VAL:HG23	28	0.24
(1,423)	1:7:A:VAL:HG22	1:32:A:VAL:HG21	29	0.24
(1,394)	1:41:A:TRP:HZ3	1:5:A:LEU:HD21	23	0.24
(1,393)	1:33:A:LEU:HD22	1:33:A:LEU:HD11	1	0.24
(1,393)	1:33:A:LEU:HD23	1:33:A:LEU:HD11	3	0.24
(1,393)	1:33:A:LEU:HD23	1:33:A:LEU:HD12	13	0.24
(1,393)	1:33:A:LEU:HD21	1:33:A:LEU:HD12	26	0.24
(1,393)	1:33:A:LEU:HD22	1:33:A:LEU:HD11	27	0.24
(1,392)	1:43:A:GLY:H	1:33:A:LEU:HD12	26	0.24
(1,376)	1:32:A:VAL:HG21	1:32:A:VAL:HG13	7	0.24
(1,376)	1:32:A:VAL:HG22	1:32:A:VAL:HG11	8	0.24
(1,376)	1:32:A:VAL:HG21	1:32:A:VAL:HG13	12	0.24
(1,376)	1:32:A:VAL:HG22	1:32:A:VAL:HG11	19	0.24
(1,376)	1:32:A:VAL:HG21	1:32:A:VAL:HG12	28	0.24
(1,355)	1:33:A:LEU:HB3	1:33:A:LEU:H	4	0.24
(1,355)	1:33:A:LEU:HB3	1:33:A:LEU:H	10	0.24
(1,355)	1:33:A:LEU:HB3	1:33:A:LEU:H	24	0.24

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,355)	1:33:A:LEU:HB3	1:33:A:LEU:H	28	0.24
(1,355)	1:33:A:LEU:HB3	1:33:A:LEU:H	30	0.24
(1,300)	1:4:A:MET:H	1:3:A:HIS:HB2	20	0.24
(1,299)	1:50:A:THR:H	1:47:A:ASP:HB2	28	0.24
(1,289)	1:32:A:VAL:HA	1:32:A:VAL:HG23	21	0.24
(1,279)	1:63:A:GLY:H	1:62:A:LEU:HA	12	0.24
(1,268)	1:30:A:ILE:HG13	1:30:A:ILE:HA	2	0.24
(1,268)	1:30:A:ILE:HG13	1:30:A:ILE:HA	10	0.24
(1,232)	1:32:A:VAL:HG13	1:41:A:TRP:HZ3	3	0.24
(1,232)	1:32:A:VAL:HG13	1:41:A:TRP:HZ3	5	0.24
(1,107)	1:41:A:TRP:HB2	1:41:A:TRP:HD1	5	0.24
(1,79)	1:17:A:TYR:HB2	1:17:A:TYR:HA	2	0.24
(1,79)	1:17:A:TYR:HB2	1:17:A:TYR:HA	3	0.24
(1,79)	1:17:A:TYR:HB2	1:17:A:TYR:HA	5	0.24
(1,79)	1:17:A:TYR:HB2	1:17:A:TYR:HA	6	0.24
(1,79)	1:17:A:TYR:HB2	1:17:A:TYR:HA	8	0.24
(1,79)	1:17:A:TYR:HB2	1:17:A:TYR:HA	9	0.24
(1,79)	1:17:A:TYR:HB2	1:17:A:TYR:HA	10	0.24
(1,79)	1:17:A:TYR:HB2	1:17:A:TYR:HA	11	0.24
(1,79)	1:17:A:TYR:HB2	1:17:A:TYR:HA	12	0.24
(1,79)	1:17:A:TYR:HB2	1:17:A:TYR:HA	14	0.24
(1,79)	1:17:A:TYR:HB2	1:17:A:TYR:HA	15	0.24
(1,79)	1:17:A:TYR:HB2	1:17:A:TYR:HA	16	0.24
(1,79)	1:17:A:TYR:HB2	1:17:A:TYR:HA	19	0.24
(1,79)	1:17:A:TYR:HB2	1:17:A:TYR:HA	20	0.24
(1,79)	1:17:A:TYR:HB2	1:17:A:TYR:HA	21	0.24
(1,79)	1:17:A:TYR:HB2	1:17:A:TYR:HA	23	0.24
(1,79)	1:17:A:TYR:HB2	1:17:A:TYR:HA	25	0.24
(1,79)	1:17:A:TYR:HB2	1:17:A:TYR:HA	27	0.24
(1,71)	1:38:A:ASP:HB2	1:38:A:ASP:HA	17	0.24
(1,60)	1:28:A:ASP:HB2	1:28:A:ASP:HA	23	0.24
(1,58)	1:28:A:ASP:H	1:28:A:ASP:HB2	1	0.24
(1,39)	1:53:A:ASP:HB3	1:54:A:ARG:H	12	0.24
(1,35)	1:27:A:GLY:HA3	1:28:A:ASP:H	18	0.24
(1,965)	1:26:A:ALA:H	1:12:A:ASP:HA	29	0.23
(1,953)	1:23:A:ASN:HB2	1:45:A:ILE:HG23	1	0.23
(1,944)	1:30:A:ILE:HD11	1:7:A:VAL:HG23	28	0.23
(1,908)	1:34:A:GLU:H	1:35:A:GLN:HG3	8	0.23
(1,906)	1:34:A:GLU:HG3	1:3:A:HIS:H	21	0.23
(1,894)	1:53:A:ASP:H	1:54:A:ARG:HD3	10	0.23
(1,876)	1:45:A:ILE:HD11	1:30:A:ILE:HA	10	0.23
(1,870)	1:29:A:ILE:HG21	1:45:A:ILE:HA	3	0.23

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,863)	1:9:A:ALA:H	1:29:A:ILE:HA	1	0.23
(1,863)	1:9:A:ALA:H	1:29:A:ILE:HA	8	0.23
(1,858)	1:13:A:TYR:HD2	1:23:A:ASN:HA	28	0.23
(1,825)	1:19:A:LEU:HB3	1:20:A:THR:H	28	0.23
(1,816)	1:5:A:LEU:HB2	1:7:A:VAL:H	7	0.23
(1,816)	1:5:A:LEU:HB2	1:7:A:VAL:H	12	0.23
(1,816)	1:5:A:LEU:HB2	1:7:A:VAL:H	24	0.23
(1,815)	1:44:A:CYS:H	1:30:A:ILE:HG22	11	0.23
(1,801)	1:65:A:ALA:HB3	1:5:A:LEU:HD12	5	0.23
(1,799)	1:41:A:TRP:H	1:41:A:TRP:HB3	17	0.23
(1,799)	1:41:A:TRP:H	1:41:A:TRP:HB3	23	0.23
(1,798)	1:24:A:VAL:HB	1:12:A:ASP:HA	6	0.23
(1,774)	1:42:A:LYS:H	1:34:A:GLU:H	21	0.23
(1,691)	1:16:A:ASN:HD21	1:16:A:ASN:HA	17	0.23
(1,685)	1:27:A:GLY:H	1:24:A:VAL:HG22	15	0.23
(1,674)	1:34:A:GLU:H	1:2:A:SER:HA	20	0.23
(1,653)	1:17:A:TYR:H	1:16:A:ASN:HB2	17	0.23
(1,649)	1:12:A:ASP:H	1:12:A:ASP:HA	1	0.23
(1,649)	1:12:A:ASP:H	1:12:A:ASP:HA	9	0.23
(1,649)	1:12:A:ASP:H	1:12:A:ASP:HA	13	0.23
(1,649)	1:12:A:ASP:H	1:12:A:ASP:HA	18	0.23
(1,649)	1:12:A:ASP:H	1:12:A:ASP:HA	26	0.23
(1,627)	1:38:A:ASP:H	1:37:A:PRO:HB3	6	0.23
(1,627)	1:38:A:ASP:H	1:37:A:PRO:HB3	25	0.23
(1,595)	1:62:A:LEU:H	1:62:A:LEU:HB2	23	0.23
(1,560)	1:64:A:GLU:H	1:64:A:GLU:HG2	30	0.23
(1,559)	1:64:A:GLU:H	1:64:A:GLU:HA	7	0.23
(1,521)	1:5:A:LEU:H	1:5:A:LEU:HA	3	0.23
(1,521)	1:5:A:LEU:H	1:5:A:LEU:HA	24	0.23
(1,466)	1:30:A:ILE:H	1:29:A:ILE:HB	11	0.23
(1,451)	1:25:A:LYS:H	1:24:A:VAL:HB	25	0.23
(1,450)	1:25:A:LYS:H	1:24:A:VAL:HG13	26	0.23
(1,450)	1:25:A:LYS:H	1:24:A:VAL:HG13	30	0.23
(1,434)	1:24:A:VAL:HG11	1:30:A:ILE:HD12	27	0.23
(1,432)	1:22:A:LEU:HD11	1:24:A:VAL:HG22	18	0.23
(1,427)	1:20:A:THR:HG21	1:55:A:VAL:HG21	4	0.23
(1,423)	1:7:A:VAL:HG23	1:32:A:VAL:HG22	16	0.23
(1,423)	1:7:A:VAL:HG23	1:32:A:VAL:HG21	27	0.23
(1,422)	1:7:A:VAL:HG23	1:32:A:VAL:HG11	7	0.23
(1,422)	1:7:A:VAL:HG22	1:32:A:VAL:HG13	25	0.23
(1,399)	1:29:A:ILE:H	1:29:A:ILE:HD12	11	0.23
(1,399)	1:29:A:ILE:H	1:29:A:ILE:HD13	19	0.23

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,393)	1:33:A:LEU:HD21	1:33:A:LEU:HD11	7	0.23
(1,393)	1:33:A:LEU:HD22	1:33:A:LEU:HD11	8	0.23
(1,393)	1:33:A:LEU:HD23	1:33:A:LEU:HD11	10	0.23
(1,393)	1:33:A:LEU:HD22	1:33:A:LEU:HD12	16	0.23
(1,393)	1:33:A:LEU:HD21	1:33:A:LEU:HD12	18	0.23
(1,390)	1:24:A:VAL:HG23	1:13:A:TYR:HB3	17	0.23
(1,376)	1:32:A:VAL:HG22	1:32:A:VAL:HG13	2	0.23
(1,376)	1:32:A:VAL:HG23	1:32:A:VAL:HG13	6	0.23
(1,376)	1:32:A:VAL:HG21	1:32:A:VAL:HG12	14	0.23
(1,355)	1:33:A:LEU:HB3	1:33:A:LEU:H	1	0.23
(1,355)	1:33:A:LEU:HB3	1:33:A:LEU:H	19	0.23
(1,355)	1:33:A:LEU:HB3	1:33:A:LEU:H	23	0.23
(1,319)	1:13:A:TYR:HB3	1:13:A:TYR:HD2	24	0.23
(1,299)	1:50:A:THR:H	1:47:A:ASP:HB2	17	0.23
(1,279)	1:63:A:GLY:H	1:62:A:LEU:HA	17	0.23
(1,203)	1:52:A:ASN:HD22	1:52:A:ASN:H	9	0.23
(1,102)	1:64:A:GLU:HB3	1:64:A:GLU:H	19	0.23
(1,69)	1:52:A:ASN:H	1:52:A:ASN:HB2	22	0.23
(1,66)	1:28:A:ASP:HB3	1:25:A:LYS:HG2	26	0.23
(1,60)	1:28:A:ASP:HB2	1:28:A:ASP:HA	1	0.23
(1,58)	1:28:A:ASP:H	1:28:A:ASP:HB2	25	0.23
(1,56)	1:28:A:ASP:H	1:28:A:ASP:HB3	4	0.23
(1,35)	1:27:A:GLY:HA3	1:28:A:ASP:H	12	0.23
(1,968)	1:10:A:THR:H	1:26:A:ALA:HA	11	0.22
(1,964)	1:45:A:ILE:H	1:55:A:VAL:HA	10	0.22
(1,954)	1:44:A:CYS:HA	1:45:A:ILE:HD12	20	0.22
(1,954)	1:44:A:CYS:HA	1:45:A:ILE:HD11	21	0.22
(1,952)	1:23:A:ASN:H	1:45:A:ILE:HD11	16	0.22
(1,917)	1:25:A:LYS:HB3	1:25:A:LYS:H	14	0.22
(1,912)	1:41:A:TRP:H	1:40:A:ARG:HG3	14	0.22
(1,908)	1:34:A:GLU:H	1:35:A:GLN:HG2	27	0.22
(1,900)	1:14:A:CYS:HB3	1:24:A:VAL:H	18	0.22
(1,896)	1:48:A:ASN:H	1:47:A:ASP:HB2	4	0.22
(1,892)	1:41:A:TRP:HH2	1:61:A:SER:HA	6	0.22
(1,888)	1:5:A:LEU:HB3	1:67:A:VAL:HA	29	0.22
(1,881)	1:13:A:TYR:HD2	1:21:A:SER:HB3	10	0.22
(1,878)	1:52:A:ASN:H	1:50:A:THR:HB	13	0.22
(1,870)	1:29:A:ILE:HG21	1:45:A:ILE:HA	9	0.22
(1,859)	1:67:A:VAL:H	1:7:A:VAL:HA	7	0.22
(1,859)	1:67:A:VAL:H	1:7:A:VAL:HA	28	0.22
(1,822)	1:54:A:ARG:HD3	1:55:A:VAL:H	24	0.22
(1,819)	1:58:A:PHE:HD1	1:42:A:LYS:H	1	0.22

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,819)	1:58:A:PHE:HD1	1:42:A:LYS:H	20	0.22
(1,816)	1:5:A:LEU:HB2	1:7:A:VAL:H	20	0.22
(1,815)	1:44:A:CYS:H	1:30:A:ILE:HG21	1	0.22
(1,804)	1:22:A:LEU:HB3	1:24:A:VAL:H	22	0.22
(1,799)	1:41:A:TRP:H	1:41:A:TRP:HB3	5	0.22
(1,755)	1:10:A:THR:H	1:63:A:GLY:H	8	0.22
(1,691)	1:16:A:ASN:HD21	1:16:A:ASN:HA	22	0.22
(1,649)	1:12:A:ASP:H	1:12:A:ASP:HA	11	0.22
(1,649)	1:12:A:ASP:H	1:12:A:ASP:HA	23	0.22
(1,649)	1:12:A:ASP:H	1:12:A:ASP:HA	25	0.22
(1,649)	1:12:A:ASP:H	1:12:A:ASP:HA	29	0.22
(1,628)	1:38:A:ASP:H	1:36:A:HIS:HB2	28	0.22
(1,627)	1:38:A:ASP:H	1:37:A:PRO:HB3	17	0.22
(1,593)	1:62:A:LEU:H	1:60:A:SER:HA	7	0.22
(1,589)	1:42:A:LYS:H	1:42:A:LYS:HB3	24	0.22
(1,583)	1:55:A:VAL:H	1:55:A:VAL:HB	18	0.22
(1,567)	1:8:A:ARG:H	1:8:A:ARG:HD3	5	0.22
(1,560)	1:64:A:GLU:H	1:64:A:GLU:HG2	24	0.22
(1,559)	1:64:A:GLU:H	1:64:A:GLU:HA	4	0.22
(1,559)	1:64:A:GLU:H	1:64:A:GLU:HA	30	0.22
(1,521)	1:5:A:LEU:H	1:5:A:LEU:HA	21	0.22
(1,521)	1:5:A:LEU:H	1:5:A:LEU:HA	23	0.22
(1,515)	1:14:A:CYS:H	1:13:A:TYR:HB3	10	0.22
(1,482)	1:46:A:HIS:H	1:30:A:ILE:HG13	18	0.22
(1,458)	1:9:A:ALA:H	1:9:A:ALA:HB2	28	0.22
(1,450)	1:25:A:LYS:H	1:24:A:VAL:HG12	27	0.22
(1,432)	1:22:A:LEU:HD11	1:24:A:VAL:HG23	3	0.22
(1,432)	1:22:A:LEU:HD12	1:24:A:VAL:HG23	12	0.22
(1,432)	1:22:A:LEU:HD11	1:24:A:VAL:HG22	29	0.22
(1,422)	1:7:A:VAL:HG21	1:32:A:VAL:HG11	12	0.22
(1,403)	1:58:A:PHE:H	1:22:A:LEU:HD13	9	0.22
(1,403)	1:58:A:PHE:H	1:22:A:LEU:HD13	13	0.22
(1,395)	1:42:A:LYS:H	1:33:A:LEU:HD12	1	0.22
(1,395)	1:42:A:LYS:H	1:33:A:LEU:HD13	24	0.22
(1,393)	1:33:A:LEU:HD23	1:33:A:LEU:HD13	9	0.22
(1,393)	1:33:A:LEU:HD23	1:33:A:LEU:HD11	17	0.22
(1,393)	1:33:A:LEU:HD22	1:33:A:LEU:HD13	21	0.22
(1,393)	1:33:A:LEU:HD23	1:33:A:LEU:HD12	24	0.22
(1,392)	1:43:A:GLY:H	1:33:A:LEU:HD12	3	0.22
(1,390)	1:24:A:VAL:HG23	1:13:A:TYR:HB3	30	0.22
(1,376)	1:32:A:VAL:HG21	1:32:A:VAL:HG11	18	0.22
(1,367)	1:9:A:ALA:HB3	1:10:A:THR:H	3	0.22

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,355)	1:33:A:LEU:HB3	1:33:A:LEU:H	12	0.22
(1,348)	1:41:A:TRP:HZ3	1:34:A:GLU:HB2	3	0.22
(1,330)	1:63:A:GLY:H	1:62:A:LEU:HB2	28	0.22
(1,299)	1:50:A:THR:H	1:47:A:ASP:HB2	16	0.22
(1,226)	1:39:A:GLY:HA3	1:41:A:TRP:HD1	21	0.22
(1,225)	1:35:A:GLN:HG3	1:41:A:TRP:HD1	21	0.22
(1,220)	1:15:A:ASN:H	1:15:A:ASN:HD22	26	0.22
(1,176)	1:33:A:LEU:H	1:42:A:LYS:H	18	0.22
(1,171)	1:31:A:THR:HG23	1:32:A:VAL:H	16	0.22
(1,106)	1:41:A:TRP:HB3	1:41:A:TRP:HZ3	25	0.22
(1,71)	1:38:A:ASP:HB2	1:38:A:ASP:HA	14	0.22
(1,71)	1:38:A:ASP:HB2	1:38:A:ASP:HA	16	0.22
(1,71)	1:38:A:ASP:HB2	1:38:A:ASP:HA	18	0.22
(1,71)	1:38:A:ASP:HB2	1:38:A:ASP:HA	30	0.22
(1,54)	1:13:A:TYR:HB3	1:13:A:TYR:HA	1	0.22
(1,54)	1:13:A:TYR:HB3	1:13:A:TYR:HA	4	0.22
(1,54)	1:13:A:TYR:HB3	1:13:A:TYR:HA	11	0.22
(1,54)	1:13:A:TYR:HB3	1:13:A:TYR:HA	17	0.22
(1,54)	1:13:A:TYR:HB3	1:13:A:TYR:HA	29	0.22
(1,35)	1:27:A:GLY:HA3	1:28:A:ASP:H	19	0.22
(1,968)	1:10:A:THR:H	1:26:A:ALA:HA	23	0.21
(1,956)	1:24:A:VAL:HG23	1:13:A:TYR:HD1	22	0.21
(1,952)	1:23:A:ASN:H	1:45:A:ILE:HD12	9	0.21
(1,944)	1:30:A:ILE:HD11	1:7:A:VAL:HG22	20	0.21
(1,940)	1:44:A:CYS:H	1:32:A:VAL:HG12	6	0.21
(1,937)	1:10:A:THR:HG22	1:10:A:THR:H	25	0.21
(1,937)	1:10:A:THR:HG22	1:10:A:THR:H	30	0.21
(1,924)	1:65:A:ALA:HB1	1:5:A:LEU:HD13	19	0.21
(1,919)	1:23:A:ASN:HD21	1:54:A:ARG:HB3	10	0.21
(1,917)	1:25:A:LYS:HB3	1:25:A:LYS:H	12	0.21
(1,900)	1:14:A:CYS:HB3	1:24:A:VAL:H	5	0.21
(1,872)	1:5:A:LEU:HD21	1:4:A:MET:HA	24	0.21
(1,859)	1:67:A:VAL:H	1:7:A:VAL:HA	15	0.21
(1,859)	1:67:A:VAL:H	1:7:A:VAL:HA	18	0.21
(1,858)	1:13:A:TYR:HD2	1:21:A:SER:HA	13	0.21
(1,853)	1:25:A:LYS:HB2	1:48:A:ASN:HD22	29	0.21
(1,830)	1:23:A:ASN:H	1:45:A:ILE:H	26	0.21
(1,815)	1:44:A:CYS:H	1:30:A:ILE:HG22	23	0.21
(1,805)	1:30:A:ILE:H	1:30:A:ILE:HD12	23	0.21
(1,804)	1:22:A:LEU:HB3	1:24:A:VAL:H	11	0.21
(1,799)	1:41:A:TRP:H	1:41:A:TRP:HB3	8	0.21
(1,799)	1:41:A:TRP:H	1:41:A:TRP:HB3	10	0.21

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,798)	1:24:A:VAL:HB	1:12:A:ASP:HA	7	0.21
(1,798)	1:24:A:VAL:HB	1:12:A:ASP:HA	16	0.21
(1,793)	1:28:A:ASP:HB3	1:24:A:VAL:HG11	16	0.21
(1,743)	1:63:A:GLY:H	1:62:A:LEU:HG	10	0.21
(1,649)	1:12:A:ASP:H	1:12:A:ASP:HA	19	0.21
(1,649)	1:12:A:ASP:H	1:12:A:ASP:HA	24	0.21
(1,628)	1:38:A:ASP:H	1:36:A:HIS:HB2	17	0.21
(1,605)	1:57:A:TYR:H	1:21:A:SER:HA	17	0.21
(1,598)	1:29:A:ILE:H	1:28:A:ASP:HA	10	0.21
(1,598)	1:29:A:ILE:H	1:28:A:ASP:HA	24	0.21
(1,559)	1:64:A:GLU:H	1:64:A:GLU:HA	16	0.21
(1,559)	1:64:A:GLU:H	1:64:A:GLU:HA	28	0.21
(1,536)	1:23:A:ASN:H	1:22:A:LEU:HD13	8	0.21
(1,521)	1:5:A:LEU:H	1:5:A:LEU:HA	7	0.21
(1,521)	1:5:A:LEU:H	1:5:A:LEU:HA	9	0.21
(1,521)	1:5:A:LEU:H	1:5:A:LEU:HA	13	0.21
(1,521)	1:5:A:LEU:H	1:5:A:LEU:HA	17	0.21
(1,521)	1:5:A:LEU:H	1:5:A:LEU:HA	19	0.21
(1,521)	1:5:A:LEU:H	1:5:A:LEU:HA	27	0.21
(1,467)	1:30:A:ILE:H	1:29:A:ILE:HG22	14	0.21
(1,442)	1:30:A:ILE:HD11	1:45:A:ILE:HD13	22	0.21
(1,432)	1:22:A:LEU:HD11	1:24:A:VAL:HG22	7	0.21
(1,432)	1:22:A:LEU:HD12	1:24:A:VAL:HG22	15	0.21
(1,422)	1:7:A:VAL:HG23	1:32:A:VAL:HG13	15	0.21
(1,422)	1:7:A:VAL:HG22	1:32:A:VAL:HG12	17	0.21
(1,393)	1:33:A:LEU:HD23	1:33:A:LEU:HD11	12	0.21
(1,369)	1:29:A:ILE:HG13	1:6:A:GLN:HE22	30	0.21
(1,367)	1:9:A:ALA:HB3	1:10:A:THR:H	22	0.21
(1,355)	1:33:A:LEU:HB3	1:33:A:LEU:H	11	0.21
(1,345)	1:41:A:TRP:HH2	1:34:A:GLU:HB2	19	0.21
(1,330)	1:63:A:GLY:H	1:62:A:LEU:HB2	19	0.21
(1,300)	1:4:A:MET:H	1:3:A:HIS:HB3	9	0.21
(1,299)	1:50:A:THR:H	1:47:A:ASP:HB2	21	0.21
(1,299)	1:50:A:THR:H	1:47:A:ASP:HB2	26	0.21
(1,225)	1:35:A:GLN:HG3	1:41:A:TRP:HD1	23	0.21
(1,221)	1:13:A:TYR:HE2	1:13:A:TYR:HD2	1	0.21
(1,221)	1:13:A:TYR:HE2	1:13:A:TYR:HD2	2	0.21
(1,221)	1:13:A:TYR:HE2	1:13:A:TYR:HD2	3	0.21
(1,221)	1:13:A:TYR:HE1	1:13:A:TYR:HD1	4	0.21
(1,221)	1:13:A:TYR:HE2	1:13:A:TYR:HD2	5	0.21
(1,221)	1:13:A:TYR:HE2	1:13:A:TYR:HD2	6	0.21
(1,221)	1:13:A:TYR:HE2	1:13:A:TYR:HD2	7	0.21

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,221)	1:13:A:TYR:HE1	1:13:A:TYR:HD1	8	0.21
(1,221)	1:13:A:TYR:HE1	1:13:A:TYR:HD1	9	0.21
(1,221)	1:13:A:TYR:HE2	1:13:A:TYR:HD2	10	0.21
(1,221)	1:13:A:TYR:HE1	1:13:A:TYR:HD1	11	0.21
(1,221)	1:13:A:TYR:HE1	1:13:A:TYR:HD1	12	0.21
(1,221)	1:13:A:TYR:HE1	1:13:A:TYR:HD1	13	0.21
(1,221)	1:13:A:TYR:HE1	1:13:A:TYR:HD1	14	0.21
(1,221)	1:13:A:TYR:HE2	1:13:A:TYR:HD2	15	0.21
(1,221)	1:13:A:TYR:HE2	1:13:A:TYR:HD2	16	0.21
(1,221)	1:13:A:TYR:HE2	1:13:A:TYR:HD2	17	0.21
(1,221)	1:13:A:TYR:HE1	1:13:A:TYR:HD1	18	0.21
(1,221)	1:13:A:TYR:HE1	1:13:A:TYR:HD1	19	0.21
(1,221)	1:13:A:TYR:HE2	1:13:A:TYR:HD2	20	0.21
(1,221)	1:13:A:TYR:HE1	1:13:A:TYR:HD1	21	0.21
(1,221)	1:13:A:TYR:HE2	1:13:A:TYR:HD2	22	0.21
(1,221)	1:13:A:TYR:HE1	1:13:A:TYR:HD1	23	0.21
(1,221)	1:13:A:TYR:HE2	1:13:A:TYR:HD2	24	0.21
(1,221)	1:13:A:TYR:HE1	1:13:A:TYR:HD1	25	0.21
(1,221)	1:13:A:TYR:HE2	1:13:A:TYR:HD2	26	0.21
(1,221)	1:13:A:TYR:HE1	1:13:A:TYR:HD1	27	0.21
(1,221)	1:13:A:TYR:HE2	1:13:A:TYR:HD2	28	0.21
(1,221)	1:13:A:TYR:HE2	1:13:A:TYR:HD2	29	0.21
(1,221)	1:13:A:TYR:HE1	1:13:A:TYR:HD1	30	0.21
(1,211)	1:25:A:LYS:H	1:11:A:LYS:H	6	0.21
(1,176)	1:33:A:LEU:H	1:42:A:LYS:H	3	0.21
(1,107)	1:41:A:TRP:HB2	1:41:A:TRP:HD1	28	0.21
(1,106)	1:41:A:TRP:HB3	1:41:A:TRP:HZ3	12	0.21
(1,71)	1:38:A:ASP:HB2	1:38:A:ASP:HA	20	0.21
(1,66)	1:28:A:ASP:HB3	1:25:A:LYS:HG2	16	0.21
(1,60)	1:28:A:ASP:HB2	1:28:A:ASP:HA	17	0.21
(1,58)	1:28:A:ASP:H	1:28:A:ASP:HB2	13	0.21
(1,55)	1:47:A:ASP:HB3	1:52:A:ASN:H	11	0.21
(1,54)	1:13:A:TYR:HB3	1:13:A:TYR:HA	5	0.21
(1,54)	1:13:A:TYR:HB3	1:13:A:TYR:HA	7	0.21
(1,54)	1:13:A:TYR:HB3	1:13:A:TYR:HA	8	0.21
(1,54)	1:13:A:TYR:HB3	1:13:A:TYR:HA	9	0.21
(1,54)	1:13:A:TYR:HB3	1:13:A:TYR:HA	10	0.21
(1,54)	1:13:A:TYR:HB3	1:13:A:TYR:HA	15	0.21
(1,54)	1:13:A:TYR:HB3	1:13:A:TYR:HA	18	0.21
(1,54)	1:13:A:TYR:HB3	1:13:A:TYR:HA	19	0.21
(1,54)	1:13:A:TYR:HB3	1:13:A:TYR:HA	23	0.21
(1,54)	1:13:A:TYR:HB3	1:13:A:TYR:HA	26	0.21

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,54)	1:13:A:TYR:HB3	1:13:A:TYR:HA	28	0.21
(1,35)	1:27:A:GLY:HA3	1:28:A:ASP:H	29	0.21
(1,29)	1:43:A:GLY:HA3	1:30:A:ILE:HG22	5	0.21
(1,968)	1:10:A:THR:H	1:63:A:GLY:HA3	18	0.2
(1,967)	1:62:A:LEU:H	1:63:A:GLY:HA2	19	0.2
(1,964)	1:45:A:ILE:H	1:55:A:VAL:HA	5	0.2
(1,954)	1:44:A:CYS:HA	1:45:A:ILE:HD12	13	0.2
(1,954)	1:44:A:CYS:HA	1:45:A:ILE:HD13	23	0.2
(1,940)	1:7:A:VAL:H	1:32:A:VAL:HG13	18	0.2
(1,937)	1:10:A:THR:HG23	1:10:A:THR:H	29	0.2
(1,906)	1:34:A:GLU:HG2	1:3:A:HIS:H	9	0.2
(1,905)	1:53:A:ASP:H	1:52:A:ASN:HB3	8	0.2
(1,905)	1:23:A:ASN:HB3	1:23:A:ASN:H	30	0.2
(1,896)	1:48:A:ASN:H	1:47:A:ASP:HB2	13	0.2
(1,892)	1:41:A:TRP:HH2	1:61:A:SER:HA	16	0.2
(1,878)	1:52:A:ASN:H	1:50:A:THR:HA	11	0.2
(1,872)	1:5:A:LEU:HD22	1:33:A:LEU:HA	22	0.2
(1,863)	1:9:A:ALA:H	1:29:A:ILE:HA	7	0.2
(1,858)	1:13:A:TYR:HD2	1:23:A:ASN:HA	25	0.2
(1,852)	1:25:A:LYS:H	1:48:A:ASN:HD22	15	0.2
(1,830)	1:23:A:ASN:H	1:45:A:ILE:H	4	0.2
(1,819)	1:58:A:PHE:HD1	1:42:A:LYS:H	25	0.2
(1,810)	1:23:A:ASN:H	1:45:A:ILE:H	5	0.2
(1,805)	1:30:A:ILE:H	1:30:A:ILE:HD12	12	0.2
(1,805)	1:30:A:ILE:H	1:30:A:ILE:HD13	15	0.2
(1,805)	1:30:A:ILE:H	1:30:A:ILE:HD12	22	0.2
(1,805)	1:30:A:ILE:H	1:30:A:ILE:HD12	24	0.2
(1,799)	1:41:A:TRP:H	1:41:A:TRP:HB3	4	0.2
(1,669)	1:43:A:GLY:H	1:55:A:VAL:HG23	4	0.2
(1,669)	1:43:A:GLY:H	1:55:A:VAL:HG21	5	0.2
(1,649)	1:12:A:ASP:H	1:12:A:ASP:HA	10	0.2
(1,649)	1:12:A:ASP:H	1:12:A:ASP:HA	12	0.2
(1,590)	1:42:A:LYS:H	1:41:A:TRP:HB2	28	0.2
(1,521)	1:5:A:LEU:H	1:5:A:LEU:HA	4	0.2
(1,521)	1:5:A:LEU:H	1:5:A:LEU:HA	5	0.2
(1,521)	1:5:A:LEU:H	1:5:A:LEU:HA	25	0.2
(1,482)	1:46:A:HIS:H	1:30:A:ILE:HG13	26	0.2
(1,446)	1:33:A:LEU:HD22	1:55:A:VAL:HG23	1	0.2
(1,445)	1:30:A:ILE:HD12	1:45:A:ILE:HG23	10	0.2
(1,432)	1:22:A:LEU:HD11	1:24:A:VAL:HG22	6	0.2
(1,427)	1:20:A:THR:HG21	1:55:A:VAL:HG22	13	0.2
(1,427)	1:20:A:THR:HG23	1:55:A:VAL:HG22	15	0.2

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,422)	1:7:A:VAL:HG21	1:32:A:VAL:HG13	13	0.2
(1,422)	1:7:A:VAL:HG23	1:32:A:VAL:HG11	20	0.2
(1,403)	1:58:A:PHE:H	1:22:A:LEU:HD11	20	0.2
(1,394)	1:41:A:TRP:HZ3	1:5:A:LEU:HD21	19	0.2
(1,393)	1:33:A:LEU:HD21	1:33:A:LEU:HD13	19	0.2
(1,393)	1:33:A:LEU:HD21	1:33:A:LEU:HD13	30	0.2
(1,391)	1:34:A:GLU:H	1:5:A:LEU:HD22	23	0.2
(1,355)	1:33:A:LEU:HB3	1:33:A:LEU:H	27	0.2
(1,330)	1:63:A:GLY:H	1:62:A:LEU:HB2	26	0.2
(1,279)	1:63:A:GLY:H	1:62:A:LEU:HA	6	0.2
(1,249)	1:13:A:TYR:HE2	1:21:A:SER:HA	19	0.2
(1,234)	1:8:A:ARG:HG3	1:6:A:GLN:HE22	1	0.2
(1,232)	1:32:A:VAL:HG12	1:41:A:TRP:HZ3	23	0.2
(1,228)	1:51:A:GLY:H	1:52:A:ASN:HD22	22	0.2
(1,222)	1:22:A:LEU:HB2	1:13:A:TYR:HE2	30	0.2
(1,90)	1:13:A:TYR:H	1:24:A:VAL:HB	25	0.2
(1,57)	1:28:A:ASP:HB2	1:25:A:LYS:HG3	25	0.2
(1,54)	1:13:A:TYR:HB3	1:13:A:TYR:HA	2	0.2
(1,54)	1:13:A:TYR:HB3	1:13:A:TYR:HA	6	0.2
(1,54)	1:13:A:TYR:HB3	1:13:A:TYR:HA	12	0.2
(1,54)	1:13:A:TYR:HB3	1:13:A:TYR:HA	13	0.2
(1,54)	1:13:A:TYR:HB3	1:13:A:TYR:HA	14	0.2
(1,54)	1:13:A:TYR:HB3	1:13:A:TYR:HA	21	0.2
(1,54)	1:13:A:TYR:HB3	1:13:A:TYR:HA	27	0.2
(1,39)	1:53:A:ASP:HB3	1:54:A:ARG:H	18	0.2
(1,35)	1:27:A:GLY:HA3	1:28:A:ASP:H	26	0.2
(1,29)	1:43:A:GLY:HA3	1:30:A:ILE:HG23	8	0.2
(1,967)	1:62:A:LEU:H	1:63:A:GLY:HA2	15	0.19
(1,944)	1:30:A:ILE:HD12	1:7:A:VAL:HG23	11	0.19
(1,929)	1:13:A:TYR:HD1	1:11:A:LYS:HG2	19	0.19
(1,900)	1:14:A:CYS:HB3	1:24:A:VAL:H	19	0.19
(1,897)	1:41:A:TRP:HB2	1:41:A:TRP:HE3	4	0.19
(1,878)	1:52:A:ASN:H	1:50:A:THR:HA	7	0.19
(1,858)	1:13:A:TYR:HD2	1:23:A:ASN:HA	10	0.19
(1,858)	1:13:A:TYR:HD2	1:23:A:ASN:HA	20	0.19
(1,855)	1:14:A:CYS:HB3	1:23:A:ASN:HD22	20	0.19
(1,840)	1:11:A:LYS:HD3	1:12:A:ASP:H	18	0.19
(1,820)	1:15:A:ASN:H	1:15:A:ASN:HB2	28	0.19
(1,805)	1:30:A:ILE:H	1:30:A:ILE:HD13	26	0.19
(1,799)	1:41:A:TRP:H	1:41:A:TRP:HB3	13	0.19
(1,791)	1:39:A:GLY:H	1:39:A:GLY:HA3	1	0.19
(1,774)	1:42:A:LYS:H	1:34:A:GLU:H	15	0.19

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,685)	1:27:A:GLY:H	1:24:A:VAL:HG21	30	0.19
(1,673)	1:34:A:GLU:H	1:33:A:LEU:HB2	18	0.19
(1,669)	1:43:A:GLY:H	1:55:A:VAL:HG22	23	0.19
(1,649)	1:12:A:ASP:H	1:12:A:ASP:HA	3	0.19
(1,649)	1:12:A:ASP:H	1:12:A:ASP:HA	5	0.19
(1,649)	1:12:A:ASP:H	1:12:A:ASP:HA	14	0.19
(1,649)	1:12:A:ASP:H	1:12:A:ASP:HA	20	0.19
(1,649)	1:12:A:ASP:H	1:12:A:ASP:HA	28	0.19
(1,627)	1:38:A:ASP:H	1:37:A:PRO:HB3	8	0.19
(1,627)	1:38:A:ASP:H	1:37:A:PRO:HB3	11	0.19
(1,598)	1:29:A:ILE:H	1:28:A:ASP:HA	22	0.19
(1,583)	1:55:A:VAL:H	1:55:A:VAL:HB	2	0.19
(1,574)	1:36:A:HIS:H	1:35:A:GLN:HB2	9	0.19
(1,559)	1:64:A:GLU:H	1:64:A:GLU:HA	11	0.19
(1,559)	1:64:A:GLU:H	1:64:A:GLU:HA	15	0.19
(1,559)	1:64:A:GLU:H	1:64:A:GLU:HA	17	0.19
(1,559)	1:64:A:GLU:H	1:64:A:GLU:HA	24	0.19
(1,559)	1:64:A:GLU:H	1:64:A:GLU:HA	29	0.19
(1,541)	1:40:A:ARG:H	1:40:A:ARG:HD2	1	0.19
(1,527)	1:26:A:ALA:H	1:25:A:LYS:HA	10	0.19
(1,521)	1:5:A:LEU:H	1:5:A:LEU:HA	1	0.19
(1,521)	1:5:A:LEU:H	1:5:A:LEU:HA	10	0.19
(1,521)	1:5:A:LEU:H	1:5:A:LEU:HA	12	0.19
(1,521)	1:5:A:LEU:H	1:5:A:LEU:HA	18	0.19
(1,521)	1:5:A:LEU:H	1:5:A:LEU:HA	26	0.19
(1,521)	1:5:A:LEU:H	1:5:A:LEU:HA	28	0.19
(1,490)	1:41:A:TRP:HE1	1:33:A:LEU:HD21	16	0.19
(1,423)	1:7:A:VAL:HG23	1:32:A:VAL:HG22	1	0.19
(1,422)	1:7:A:VAL:HG22	1:32:A:VAL:HG12	9	0.19
(1,422)	1:7:A:VAL:HG21	1:32:A:VAL:HG12	11	0.19
(1,395)	1:42:A:LYS:H	1:33:A:LEU:HD12	10	0.19
(1,382)	1:65:A:ALA:H	1:5:A:LEU:HB3	2	0.19
(1,355)	1:33:A:LEU:HB3	1:33:A:LEU:H	21	0.19
(1,346)	1:55:A:VAL:H	1:54:A:ARG:HB3	20	0.19
(1,299)	1:50:A:THR:H	1:47:A:ASP:HB2	24	0.19
(1,295)	1:41:A:TRP:HD1	1:36:A:HIS:HB2	21	0.19
(1,279)	1:63:A:GLY:H	1:62:A:LEU:HA	18	0.19
(1,228)	1:51:A:GLY:H	1:52:A:ASN:HD22	4	0.19
(1,211)	1:25:A:LYS:H	1:11:A:LYS:H	14	0.19
(1,116)	1:35:A:GLN:HB3	1:36:A:HIS:H	10	0.19
(1,71)	1:38:A:ASP:HB2	1:38:A:ASP:HA	9	0.19
(1,71)	1:38:A:ASP:HB2	1:38:A:ASP:HA	12	0.19

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,71)	1:38:A:ASP:HB2	1:38:A:ASP:HA	22	0.19
(1,54)	1:13:A:TYR:HB3	1:13:A:TYR:HA	24	0.19
(1,54)	1:13:A:TYR:HB3	1:13:A:TYR:HA	30	0.19
(1,49)	1:13:A:TYR:HB2	1:13:A:TYR:HA	25	0.19
(1,38)	1:63:A:GLY:HA3	1:7:A:VAL:HB	7	0.19
(1,968)	1:10:A:THR:H	1:26:A:ALA:HA	13	0.18
(1,964)	1:45:A:ILE:H	1:55:A:VAL:HA	8	0.18
(1,964)	1:45:A:ILE:H	1:55:A:VAL:HA	27	0.18
(1,953)	1:23:A:ASN:HB2	1:45:A:ILE:HG21	23	0.18
(1,944)	1:30:A:ILE:HD11	1:7:A:VAL:HG22	1	0.18
(1,940)	1:44:A:CYS:H	1:32:A:VAL:HG11	25	0.18
(1,924)	1:7:A:VAL:HG11	1:65:A:ALA:HB2	1	0.18
(1,903)	1:28:A:ASP:HB2	1:25:A:LYS:HG3	26	0.18
(1,902)	1:28:A:ASP:H	1:28:A:ASP:HB2	1	0.18
(1,900)	1:14:A:CYS:HB3	1:24:A:VAL:H	14	0.18
(1,900)	1:24:A:VAL:H	1:14:A:CYS:HB2	29	0.18
(1,896)	1:48:A:ASN:H	1:47:A:ASP:HB2	25	0.18
(1,866)	1:5:A:LEU:HD11	1:65:A:ALA:HA	11	0.18
(1,859)	1:67:A:VAL:H	1:7:A:VAL:HA	21	0.18
(1,859)	1:67:A:VAL:H	1:7:A:VAL:HA	26	0.18
(1,819)	1:58:A:PHE:HD1	1:42:A:LYS:H	16	0.18
(1,805)	1:30:A:ILE:H	1:30:A:ILE:HD12	18	0.18
(1,803)	1:30:A:ILE:HG22	1:32:A:VAL:H	18	0.18
(1,799)	1:41:A:TRP:H	1:41:A:TRP:HB3	15	0.18
(1,734)	1:39:A:GLY:H	1:40:A:ARG:HB2	28	0.18
(1,660)	1:10:A:THR:H	1:11:A:LYS:HB2	11	0.18
(1,649)	1:12:A:ASP:H	1:12:A:ASP:HA	6	0.18
(1,649)	1:12:A:ASP:H	1:12:A:ASP:HA	7	0.18
(1,649)	1:12:A:ASP:H	1:12:A:ASP:HA	15	0.18
(1,649)	1:12:A:ASP:H	1:12:A:ASP:HA	17	0.18
(1,649)	1:12:A:ASP:H	1:12:A:ASP:HA	21	0.18
(1,649)	1:12:A:ASP:H	1:12:A:ASP:HA	27	0.18
(1,581)	1:13:A:TYR:H	1:13:A:TYR:HA	1	0.18
(1,581)	1:13:A:TYR:H	1:13:A:TYR:HA	2	0.18
(1,581)	1:13:A:TYR:H	1:13:A:TYR:HA	4	0.18
(1,581)	1:13:A:TYR:H	1:13:A:TYR:HA	10	0.18
(1,581)	1:13:A:TYR:H	1:13:A:TYR:HA	12	0.18
(1,581)	1:13:A:TYR:H	1:13:A:TYR:HA	13	0.18
(1,581)	1:13:A:TYR:H	1:13:A:TYR:HA	15	0.18
(1,581)	1:13:A:TYR:H	1:13:A:TYR:HA	25	0.18
(1,581)	1:13:A:TYR:H	1:13:A:TYR:HA	29	0.18
(1,559)	1:64:A:GLU:H	1:64:A:GLU:HA	13	0.18

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,521)	1:5:A:LEU:H	1:5:A:LEU:HA	16	0.18
(1,521)	1:5:A:LEU:H	1:5:A:LEU:HA	29	0.18
(1,515)	1:14:A:CYS:H	1:13:A:TYR:HB3	28	0.18
(1,512)	1:65:A:ALA:H	1:66:A:ILE:HG22	16	0.18
(1,458)	1:9:A:ALA:H	1:9:A:ALA:HB3	8	0.18
(1,458)	1:9:A:ALA:H	1:9:A:ALA:HB1	13	0.18
(1,458)	1:9:A:ALA:H	1:9:A:ALA:HB3	16	0.18
(1,453)	1:32:A:VAL:H	1:32:A:VAL:HG11	5	0.18
(1,450)	1:25:A:LYS:H	1:24:A:VAL:HG13	21	0.18
(1,446)	1:33:A:LEU:HD21	1:55:A:VAL:HG22	14	0.18
(1,432)	1:22:A:LEU:HD13	1:24:A:VAL:HG22	28	0.18
(1,427)	1:20:A:THR:HG21	1:55:A:VAL:HG21	11	0.18
(1,418)	1:7:A:VAL:HG12	1:32:A:VAL:HG21	30	0.18
(1,394)	1:41:A:TRP:HZ3	1:5:A:LEU:HD21	16	0.18
(1,367)	1:9:A:ALA:HB1	1:10:A:THR:H	17	0.18
(1,355)	1:33:A:LEU:HB3	1:33:A:LEU:H	6	0.18
(1,355)	1:33:A:LEU:HB3	1:33:A:LEU:H	25	0.18
(1,330)	1:63:A:GLY:H	1:62:A:LEU:HB2	1	0.18
(1,330)	1:63:A:GLY:H	1:62:A:LEU:HB2	24	0.18
(1,299)	1:50:A:THR:H	1:47:A:ASP:HB2	1	0.18
(1,232)	1:32:A:VAL:HG11	1:41:A:TRP:HZ3	9	0.18
(1,222)	1:22:A:LEU:HB2	1:13:A:TYR:HE2	22	0.18
(1,211)	1:25:A:LYS:H	1:11:A:LYS:H	26	0.18
(1,203)	1:52:A:ASN:HD22	1:52:A:ASN:H	24	0.18
(1,190)	1:34:A:GLU:HG3	1:3:A:HIS:H	24	0.18
(1,152)	1:20:A:THR:HA	1:56:A:GLY:HA3	8	0.18
(1,106)	1:41:A:TRP:HB3	1:41:A:TRP:HZ3	6	0.18
(1,106)	1:41:A:TRP:HB3	1:41:A:TRP:HZ3	23	0.18
(1,71)	1:38:A:ASP:HB2	1:38:A:ASP:HA	3	0.18
(1,71)	1:38:A:ASP:HB2	1:38:A:ASP:HA	27	0.18
(1,71)	1:38:A:ASP:HB2	1:38:A:ASP:HA	28	0.18
(1,71)	1:38:A:ASP:HB2	1:38:A:ASP:HA	29	0.18
(1,66)	1:28:A:ASP:HB3	1:25:A:LYS:HG2	24	0.18
(1,60)	1:28:A:ASP:HB2	1:28:A:ASP:HA	3	0.18
(1,58)	1:28:A:ASP:H	1:28:A:ASP:HB2	7	0.18
(1,55)	1:47:A:ASP:HB3	1:52:A:ASN:H	27	0.18
(1,54)	1:13:A:TYR:HB3	1:13:A:TYR:HA	16	0.18
(1,54)	1:13:A:TYR:HB3	1:13:A:TYR:HA	20	0.18
(1,49)	1:13:A:TYR:HB2	1:13:A:TYR:HA	22	0.18
(1,968)	1:10:A:THR:H	1:63:A:GLY:HA3	3	0.17
(1,968)	1:10:A:THR:H	1:63:A:GLY:HA3	4	0.17
(1,954)	1:44:A:CYS:HA	1:45:A:ILE:HD11	8	0.17

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,948)	1:6:A:GLN:H	1:67:A:VAL:HG11	2	0.17
(1,944)	1:30:A:ILE:HD12	1:7:A:VAL:HG22	16	0.17
(1,925)	1:11:A:LYS:HG2	1:13:A:TYR:H	7	0.17
(1,924)	1:7:A:VAL:HG11	1:65:A:ALA:HB1	26	0.17
(1,917)	1:25:A:LYS:HB3	1:25:A:LYS:H	20	0.17
(1,898)	1:32:A:VAL:HG21	1:60:A:SER:HA	18	0.17
(1,896)	1:48:A:ASN:H	1:47:A:ASP:HB2	1	0.17
(1,876)	1:45:A:ILE:HD13	1:30:A:ILE:HA	19	0.17
(1,870)	1:29:A:ILE:HG22	1:45:A:ILE:HA	16	0.17
(1,870)	1:29:A:ILE:HG22	1:45:A:ILE:HA	23	0.17
(1,870)	1:29:A:ILE:HG23	1:45:A:ILE:HA	25	0.17
(1,863)	1:9:A:ALA:H	1:29:A:ILE:HA	30	0.17
(1,836)	1:40:A:ARG:H	1:41:A:TRP:HD1	8	0.17
(1,830)	1:23:A:ASN:H	1:45:A:ILE:H	11	0.17
(1,825)	1:34:A:GLU:HB2	1:3:A:HIS:H	4	0.17
(1,823)	1:28:A:ASP:H	1:25:A:LYS:HG2	11	0.17
(1,815)	1:44:A:CYS:H	1:30:A:ILE:HG21	21	0.17
(1,805)	1:30:A:ILE:H	1:30:A:ILE:HD11	25	0.17
(1,799)	1:41:A:TRP:H	1:41:A:TRP:HB3	7	0.17
(1,793)	1:28:A:ASP:HB3	1:45:A:ILE:HG22	19	0.17
(1,787)	1:61:A:SER:H	1:63:A:GLY:H	7	0.17
(1,743)	1:63:A:GLY:H	1:62:A:LEU:HG	25	0.17
(1,685)	1:27:A:GLY:H	1:24:A:VAL:HG21	16	0.17
(1,660)	1:10:A:THR:H	1:11:A:LYS:HB2	25	0.17
(1,649)	1:12:A:ASP:H	1:12:A:ASP:HA	8	0.17
(1,649)	1:12:A:ASP:H	1:12:A:ASP:HA	22	0.17
(1,649)	1:12:A:ASP:H	1:12:A:ASP:HA	30	0.17
(1,598)	1:29:A:ILE:H	1:28:A:ASP:HA	16	0.17
(1,590)	1:42:A:LYS:H	1:41:A:TRP:HB2	24	0.17
(1,583)	1:55:A:VAL:H	1:55:A:VAL:HB	8	0.17
(1,583)	1:55:A:VAL:H	1:55:A:VAL:HB	19	0.17
(1,583)	1:55:A:VAL:H	1:55:A:VAL:HB	22	0.17
(1,581)	1:13:A:TYR:H	1:13:A:TYR:HA	3	0.17
(1,581)	1:13:A:TYR:H	1:13:A:TYR:HA	14	0.17
(1,581)	1:13:A:TYR:H	1:13:A:TYR:HA	19	0.17
(1,581)	1:13:A:TYR:H	1:13:A:TYR:HA	22	0.17
(1,581)	1:13:A:TYR:H	1:13:A:TYR:HA	23	0.17
(1,567)	1:8:A:ARG:H	1:8:A:ARG:HD3	19	0.17
(1,559)	1:64:A:GLU:H	1:64:A:GLU:HA	1	0.17
(1,559)	1:64:A:GLU:H	1:64:A:GLU:HA	5	0.17
(1,559)	1:64:A:GLU:H	1:64:A:GLU:HA	6	0.17
(1,559)	1:64:A:GLU:H	1:64:A:GLU:HA	8	0.17

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,559)	1:64:A:GLU:H	1:64:A:GLU:HA	20	0.17
(1,515)	1:14:A:CYS:H	1:13:A:TYR:HB3	20	0.17
(1,512)	1:65:A:ALA:H	1:66:A:ILE:HG21	15	0.17
(1,482)	1:46:A:HIS:H	1:30:A:ILE:HG13	16	0.17
(1,458)	1:9:A:ALA:H	1:9:A:ALA:HB3	11	0.17
(1,423)	1:7:A:VAL:HG21	1:32:A:VAL:HG22	11	0.17
(1,423)	1:7:A:VAL:HG23	1:32:A:VAL:HG22	15	0.17
(1,414)	1:5:A:LEU:HD12	1:32:A:VAL:HG11	16	0.17
(1,392)	1:43:A:GLY:H	1:33:A:LEU:HD11	25	0.17
(1,360)	1:27:A:GLY:H	1:25:A:LYS:HG2	11	0.17
(1,334)	1:49:A:ARG:H	1:49:A:ARG:HB3	2	0.17
(1,334)	1:49:A:ARG:H	1:49:A:ARG:HB3	17	0.17
(1,332)	1:11:A:LYS:HB3	1:11:A:LYS:H	10	0.17
(1,330)	1:63:A:GLY:H	1:62:A:LEU:HB2	2	0.17
(1,252)	1:15:A:ASN:H	1:15:A:ASN:HA	1	0.17
(1,252)	1:15:A:ASN:H	1:15:A:ASN:HA	8	0.17
(1,252)	1:15:A:ASN:H	1:15:A:ASN:HA	17	0.17
(1,252)	1:15:A:ASN:H	1:15:A:ASN:HA	19	0.17
(1,252)	1:15:A:ASN:H	1:15:A:ASN:HA	21	0.17
(1,203)	1:52:A:ASN:HD22	1:52:A:ASN:H	6	0.17
(1,190)	1:34:A:GLU:HG3	1:3:A:HIS:H	17	0.17
(1,107)	1:41:A:TRP:HB2	1:41:A:TRP:HD1	15	0.17
(1,107)	1:41:A:TRP:HB2	1:41:A:TRP:HD1	29	0.17
(1,106)	1:41:A:TRP:HB3	1:41:A:TRP:HZ3	8	0.17
(1,106)	1:41:A:TRP:HB3	1:41:A:TRP:HZ3	16	0.17
(1,74)	1:15:A:ASN:HD22	1:15:A:ASN:HB2	29	0.17
(1,71)	1:38:A:ASP:HB2	1:38:A:ASP:HA	7	0.17
(1,71)	1:38:A:ASP:HB2	1:38:A:ASP:HA	23	0.17
(1,66)	1:28:A:ASP:HB3	1:25:A:LYS:HG2	22	0.17
(1,65)	1:28:A:ASP:HB3	1:25:A:LYS:H	14	0.17
(1,60)	1:28:A:ASP:HB2	1:28:A:ASP:HA	29	0.17
(1,54)	1:13:A:TYR:HB3	1:13:A:TYR:HA	3	0.17
(1,40)	1:53:A:ASP:HB3	1:53:A:ASP:H	13	0.17
(1,35)	1:27:A:GLY:HA3	1:28:A:ASP:H	21	0.17
(1,35)	1:27:A:GLY:HA3	1:28:A:ASP:H	28	0.17
(1,17)	1:26:A:ALA:HA	1:24:A:VAL:HG12	13	0.17
(1,967)	1:62:A:LEU:H	1:63:A:GLY:HA2	22	0.16
(1,967)	1:62:A:LEU:H	1:63:A:GLY:HA2	25	0.16
(1,964)	1:45:A:ILE:H	1:55:A:VAL:HA	11	0.16
(1,958)	1:61:A:SER:H	1:5:A:LEU:HD23	25	0.16
(1,956)	1:24:A:VAL:HG21	1:13:A:TYR:HD2	8	0.16
(1,954)	1:44:A:CYS:HA	1:45:A:ILE:HD12	16	0.16

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,954)	1:44:A:CYS:HA	1:45:A:ILE:HD13	19	0.16
(1,940)	1:44:A:CYS:H	1:32:A:VAL:HG11	1	0.16
(1,937)	1:10:A:THR:HG23	1:10:A:THR:H	2	0.16
(1,924)	1:65:A:ALA:HB1	1:5:A:LEU:HD13	3	0.16
(1,924)	1:65:A:ALA:HB1	1:5:A:LEU:HD12	14	0.16
(1,922)	1:23:A:ASN:H	1:22:A:LEU:HB2	30	0.16
(1,913)	1:6:A:GLN:HE21	1:66:A:ILE:HG12	9	0.16
(1,912)	1:41:A:TRP:H	1:40:A:ARG:HG3	15	0.16
(1,900)	1:14:A:CYS:HB3	1:24:A:VAL:H	20	0.16
(1,900)	1:14:A:CYS:HB3	1:24:A:VAL:H	27	0.16
(1,894)	1:23:A:ASN:H	1:54:A:ARG:HD2	29	0.16
(1,881)	1:13:A:TYR:HD2	1:21:A:SER:HB3	4	0.16
(1,881)	1:13:A:TYR:HD2	1:21:A:SER:HB3	29	0.16
(1,872)	1:5:A:LEU:HD21	1:33:A:LEU:HA	3	0.16
(1,870)	1:29:A:ILE:HG21	1:45:A:ILE:HA	29	0.16
(1,799)	1:41:A:TRP:H	1:41:A:TRP:HB3	3	0.16
(1,799)	1:41:A:TRP:H	1:41:A:TRP:HB3	14	0.16
(1,734)	1:39:A:GLY:H	1:40:A:ARG:HB2	10	0.16
(1,734)	1:39:A:GLY:H	1:40:A:ARG:HB2	14	0.16
(1,714)	1:15:A:ASN:HD22	1:15:A:ASN:HA	25	0.16
(1,691)	1:16:A:ASN:HD21	1:16:A:ASN:HA	11	0.16
(1,685)	1:27:A:GLY:H	1:24:A:VAL:HG23	27	0.16
(1,669)	1:43:A:GLY:H	1:55:A:VAL:HG22	14	0.16
(1,649)	1:12:A:ASP:H	1:12:A:ASP:HA	16	0.16
(1,647)	1:12:A:ASP:H	1:13:A:TYR:H	16	0.16
(1,598)	1:29:A:ILE:H	1:28:A:ASP:HA	28	0.16
(1,589)	1:42:A:LYS:H	1:42:A:LYS:HB3	20	0.16
(1,581)	1:13:A:TYR:H	1:13:A:TYR:HA	9	0.16
(1,581)	1:13:A:TYR:H	1:13:A:TYR:HA	16	0.16
(1,581)	1:13:A:TYR:H	1:13:A:TYR:HA	17	0.16
(1,581)	1:13:A:TYR:H	1:13:A:TYR:HA	18	0.16
(1,581)	1:13:A:TYR:H	1:13:A:TYR:HA	27	0.16
(1,581)	1:13:A:TYR:H	1:13:A:TYR:HA	30	0.16
(1,560)	1:64:A:GLU:H	1:64:A:GLU:HG2	5	0.16
(1,559)	1:64:A:GLU:H	1:64:A:GLU:HA	2	0.16
(1,559)	1:64:A:GLU:H	1:64:A:GLU:HA	12	0.16
(1,559)	1:64:A:GLU:H	1:64:A:GLU:HA	14	0.16
(1,559)	1:64:A:GLU:H	1:64:A:GLU:HA	18	0.16
(1,559)	1:64:A:GLU:H	1:64:A:GLU:HA	22	0.16
(1,559)	1:64:A:GLU:H	1:64:A:GLU:HA	25	0.16
(1,559)	1:64:A:GLU:H	1:64:A:GLU:HA	26	0.16
(1,559)	1:64:A:GLU:H	1:64:A:GLU:HA	27	0.16

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,521)	1:5:A:LEU:H	1:5:A:LEU:HA	6	0.16
(1,521)	1:5:A:LEU:H	1:5:A:LEU:HA	11	0.16
(1,521)	1:5:A:LEU:H	1:5:A:LEU:HA	30	0.16
(1,511)	1:65:A:ALA:H	1:64:A:GLU:HG2	10	0.16
(1,461)	1:67:A:VAL:H	1:67:A:VAL:HA	5	0.16
(1,434)	1:24:A:VAL:HG13	1:30:A:ILE:HD13	24	0.16
(1,431)	1:22:A:LEU:HD13	1:24:A:VAL:HG11	5	0.16
(1,423)	1:7:A:VAL:HG22	1:32:A:VAL:HG23	17	0.16
(1,422)	1:7:A:VAL:HG21	1:32:A:VAL:HG11	26	0.16
(1,422)	1:7:A:VAL:HG22	1:32:A:VAL:HG13	29	0.16
(1,403)	1:58:A:PHE:H	1:22:A:LEU:HD11	21	0.16
(1,371)	1:28:A:ASP:H	1:10:A:THR:HG22	10	0.16
(1,369)	1:29:A:ILE:HG13	1:6:A:GLN:HE22	20	0.16
(1,367)	1:9:A:ALA:HB2	1:10:A:THR:H	1	0.16
(1,305)	1:13:A:TYR:HD1	1:11:A:LYS:HE3	5	0.16
(1,279)	1:63:A:GLY:H	1:62:A:LEU:HA	16	0.16
(1,252)	1:15:A:ASN:H	1:15:A:ASN:HA	29	0.16
(1,244)	1:30:A:ILE:HG12	1:8:A:ARG:HA	7	0.16
(1,234)	1:8:A:ARG:HG3	1:6:A:GLN:HE22	20	0.16
(1,211)	1:25:A:LYS:H	1:11:A:LYS:H	25	0.16
(1,203)	1:52:A:ASN:HD22	1:52:A:ASN:H	22	0.16
(1,135)	1:8:A:ARG:HA	1:27:A:GLY:HA3	2	0.16
(1,106)	1:41:A:TRP:HB3	1:41:A:TRP:HZ3	3	0.16
(1,106)	1:41:A:TRP:HB3	1:41:A:TRP:HZ3	10	0.16
(1,60)	1:28:A:ASP:HB2	1:28:A:ASP:HA	2	0.16
(1,46)	1:13:A:TYR:HB2	1:11:A:LYS:HD2	9	0.16
(1,35)	1:27:A:GLY:HA3	1:28:A:ASP:H	23	0.16
(1,29)	1:43:A:GLY:HA3	1:30:A:ILE:HG22	23	0.16
(1,25)	1:43:A:GLY:HA2	1:33:A:LEU:HD12	8	0.16
(1,967)	1:62:A:LEU:H	1:63:A:GLY:HA2	18	0.15
(1,954)	1:44:A:CYS:HA	1:45:A:ILE:HD13	27	0.15
(1,952)	1:23:A:ASN:H	1:45:A:ILE:HD13	21	0.15
(1,948)	1:6:A:GLN:H	1:67:A:VAL:HG23	22	0.15
(1,946)	1:7:A:VAL:HG22	1:30:A:ILE:H	7	0.15
(1,934)	1:26:A:ALA:H	1:25:A:LYS:HG2	11	0.15
(1,902)	1:28:A:ASP:H	1:28:A:ASP:HB2	13	0.15
(1,878)	1:52:A:ASN:H	1:50:A:THR:HA	17	0.15
(1,859)	1:67:A:VAL:H	1:7:A:VAL:HA	11	0.15
(1,849)	1:13:A:TYR:H	1:15:A:ASN:HD22	15	0.15
(1,830)	1:23:A:ASN:H	1:45:A:ILE:H	3	0.15
(1,799)	1:41:A:TRP:H	1:41:A:TRP:HB3	22	0.15
(1,799)	1:41:A:TRP:H	1:41:A:TRP:HB3	27	0.15

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,791)	1:39:A:GLY:H	1:39:A:GLY:HA3	24	0.15
(1,743)	1:63:A:GLY:H	1:62:A:LEU:HG	3	0.15
(1,714)	1:15:A:ASN:HD22	1:15:A:ASN:HA	23	0.15
(1,709)	1:35:A:GLN:HE21	1:35:A:GLN:HB3	27	0.15
(1,669)	1:43:A:GLY:H	1:55:A:VAL:HG21	25	0.15
(1,664)	1:61:A:SER:H	1:61:A:SER:HB3	9	0.15
(1,653)	1:17:A:TYR:H	1:16:A:ASN:HB2	8	0.15
(1,624)	1:18:A:ASP:H	1:17:A:TYR:HA	5	0.15
(1,624)	1:18:A:ASP:H	1:17:A:TYR:HA	14	0.15
(1,598)	1:29:A:ILE:H	1:28:A:ASP:HA	12	0.15
(1,583)	1:55:A:VAL:H	1:55:A:VAL:HB	17	0.15
(1,583)	1:55:A:VAL:H	1:55:A:VAL:HB	24	0.15
(1,581)	1:13:A:TYR:H	1:13:A:TYR:HA	8	0.15
(1,581)	1:13:A:TYR:H	1:13:A:TYR:HA	20	0.15
(1,581)	1:13:A:TYR:H	1:13:A:TYR:HA	21	0.15
(1,573)	1:36:A:HIS:H	1:39:A:GLY:H	24	0.15
(1,560)	1:64:A:GLU:H	1:64:A:GLU:HG2	11	0.15
(1,560)	1:64:A:GLU:H	1:64:A:GLU:HG2	23	0.15
(1,559)	1:64:A:GLU:H	1:64:A:GLU:HA	9	0.15
(1,559)	1:64:A:GLU:H	1:64:A:GLU:HA	21	0.15
(1,521)	1:5:A:LEU:H	1:5:A:LEU:HA	2	0.15
(1,521)	1:5:A:LEU:H	1:5:A:LEU:HA	20	0.15
(1,458)	1:9:A:ALA:H	1:9:A:ALA:HB3	25	0.15
(1,423)	1:7:A:VAL:HG23	1:32:A:VAL:HG21	22	0.15
(1,422)	1:7:A:VAL:HG23	1:32:A:VAL:HG13	14	0.15
(1,422)	1:7:A:VAL:HG23	1:32:A:VAL:HG12	16	0.15
(1,422)	1:7:A:VAL:HG23	1:32:A:VAL:HG12	24	0.15
(1,395)	1:42:A:LYS:H	1:33:A:LEU:HD13	14	0.15
(1,390)	1:24:A:VAL:HG21	1:13:A:TYR:HB3	9	0.15
(1,382)	1:65:A:ALA:H	1:5:A:LEU:HB3	27	0.15
(1,367)	1:9:A:ALA:HB3	1:10:A:THR:H	10	0.15
(1,367)	1:9:A:ALA:HB2	1:10:A:THR:H	16	0.15
(1,355)	1:33:A:LEU:HB3	1:33:A:LEU:H	18	0.15
(1,355)	1:33:A:LEU:HB3	1:33:A:LEU:H	29	0.15
(1,330)	1:63:A:GLY:H	1:62:A:LEU:HB2	27	0.15
(1,252)	1:15:A:ASN:H	1:15:A:ASN:HA	4	0.15
(1,252)	1:15:A:ASN:H	1:15:A:ASN:HA	28	0.15
(1,171)	1:31:A:THR:HG22	1:32:A:VAL:H	13	0.15
(1,57)	1:28:A:ASP:HB2	1:25:A:LYS:HG3	24	0.15
(1,43)	1:13:A:TYR:HB2	1:24:A:VAL:HG21	4	0.15
(1,43)	1:13:A:TYR:HB2	1:24:A:VAL:HG22	12	0.15
(1,35)	1:27:A:GLY:HA3	1:28:A:ASP:H	16	0.15

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,967)	1:62:A:LEU:H	1:63:A:GLY:HA2	1	0.14
(1,948)	1:6:A:GLN:H	1:67:A:VAL:HG22	5	0.14
(1,931)	1:41:A:TRP:HZ2	1:60:A:SER:HB3	11	0.14
(1,924)	1:65:A:ALA:HB2	1:5:A:LEU:HD11	7	0.14
(1,924)	1:65:A:ALA:HB1	1:5:A:LEU:HD13	9	0.14
(1,922)	1:23:A:ASN:H	1:22:A:LEU:HB2	8	0.14
(1,909)	1:41:A:TRP:HD1	1:40:A:ARG:HG2	5	0.14
(1,900)	1:14:A:CYS:HB3	1:24:A:VAL:H	15	0.14
(1,894)	1:53:A:ASP:H	1:54:A:ARG:HD3	24	0.14
(1,855)	1:52:A:ASN:HB2	1:23:A:ASN:HD22	6	0.14
(1,852)	1:13:A:TYR:HE2	1:24:A:VAL:H	19	0.14
(1,852)	1:13:A:TYR:HE2	1:24:A:VAL:H	30	0.14
(1,840)	1:11:A:LYS:HD3	1:12:A:ASP:H	4	0.14
(1,830)	1:9:A:ALA:H	1:29:A:ILE:H	1	0.14
(1,830)	1:23:A:ASN:H	1:45:A:ILE:H	25	0.14
(1,825)	1:19:A:LEU:HB3	1:20:A:THR:H	21	0.14
(1,799)	1:41:A:TRP:H	1:41:A:TRP:HB3	29	0.14
(1,775)	1:38:A:ASP:H	1:40:A:ARG:H	26	0.14
(1,743)	1:63:A:GLY:H	1:62:A:LEU:HG	19	0.14
(1,735)	1:39:A:GLY:H	1:41:A:TRP:HD1	3	0.14
(1,727)	1:39:A:GLY:H	1:40:A:ARG:HG2	19	0.14
(1,714)	1:15:A:ASN:HD22	1:15:A:ASN:HA	4	0.14
(1,673)	1:34:A:GLU:H	1:33:A:LEU:HB2	16	0.14
(1,669)	1:43:A:GLY:H	1:55:A:VAL:HG23	7	0.14
(1,660)	1:10:A:THR:H	1:11:A:LYS:HB2	5	0.14
(1,627)	1:38:A:ASP:H	1:37:A:PRO:HB3	14	0.14
(1,624)	1:18:A:ASP:H	1:17:A:TYR:HA	2	0.14
(1,624)	1:18:A:ASP:H	1:17:A:TYR:HA	3	0.14
(1,624)	1:18:A:ASP:H	1:17:A:TYR:HA	12	0.14
(1,624)	1:18:A:ASP:H	1:17:A:TYR:HA	20	0.14
(1,624)	1:18:A:ASP:H	1:17:A:TYR:HA	26	0.14
(1,594)	1:29:A:ILE:H	1:28:A:ASP:HB3	11	0.14
(1,593)	1:62:A:LEU:H	1:60:A:SER:HA	8	0.14
(1,587)	1:53:A:ASP:H	1:53:A:ASP:HB2	22	0.14
(1,581)	1:13:A:TYR:H	1:13:A:TYR:HA	11	0.14
(1,559)	1:64:A:GLU:H	1:64:A:GLU:HA	23	0.14
(1,552)	1:54:A:ARG:H	1:45:A:ILE:HG22	4	0.14
(1,545)	1:40:A:ARG:H	1:36:A:HIS:HB2	27	0.14
(1,515)	1:14:A:CYS:H	1:13:A:TYR:HB3	3	0.14
(1,493)	1:41:A:TRP:HE1	1:40:A:ARG:H	16	0.14
(1,493)	1:41:A:TRP:HE1	1:40:A:ARG:H	19	0.14
(1,458)	1:9:A:ALA:H	1:9:A:ALA:HB3	1	0.14

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,458)	1:9:A:ALA:H	1:9:A:ALA:HB1	22	0.14
(1,453)	1:32:A:VAL:H	1:32:A:VAL:HG12	27	0.14
(1,432)	1:22:A:LEU:HD13	1:24:A:VAL:HG21	17	0.14
(1,423)	1:7:A:VAL:HG21	1:32:A:VAL:HG21	23	0.14
(1,423)	1:7:A:VAL:HG22	1:32:A:VAL:HG21	30	0.14
(1,422)	1:7:A:VAL:HG21	1:32:A:VAL:HG11	10	0.14
(1,422)	1:7:A:VAL:HG22	1:32:A:VAL:HG12	21	0.14
(1,414)	1:5:A:LEU:HD13	1:32:A:VAL:HG13	6	0.14
(1,414)	1:5:A:LEU:HD13	1:32:A:VAL:HG12	14	0.14
(1,398)	1:5:A:LEU:H	1:5:A:LEU:HD22	28	0.14
(1,395)	1:42:A:LYS:H	1:33:A:LEU:HD11	25	0.14
(1,392)	1:43:A:GLY:H	1:33:A:LEU:HD12	22	0.14
(1,383)	1:41:A:TRP:HH2	1:5:A:LEU:HD11	23	0.14
(1,340)	1:63:A:GLY:H	1:7:A:VAL:HB	14	0.14
(1,330)	1:63:A:GLY:H	1:62:A:LEU:HB2	23	0.14
(1,299)	1:50:A:THR:H	1:47:A:ASP:HB2	20	0.14
(1,295)	1:41:A:TRP:HD1	1:36:A:HIS:HB2	29	0.14
(1,252)	1:15:A:ASN:H	1:15:A:ASN:HA	25	0.14
(1,232)	1:32:A:VAL:HG11	1:41:A:TRP:HZ3	21	0.14
(1,223)	1:48:A:ASN:HB3	1:48:A:ASN:HD22	4	0.14
(1,223)	1:48:A:ASN:HB3	1:48:A:ASN:HD22	13	0.14
(1,223)	1:48:A:ASN:HB3	1:48:A:ASN:HD22	15	0.14
(1,223)	1:48:A:ASN:HB3	1:48:A:ASN:HD22	17	0.14
(1,203)	1:52:A:ASN:HD22	1:52:A:ASN:H	5	0.14
(1,203)	1:52:A:ASN:HD22	1:52:A:ASN:H	11	0.14
(1,137)	1:8:A:ARG:HA	1:29:A:ILE:HA	24	0.14
(1,71)	1:38:A:ASP:HB2	1:38:A:ASP:HA	1	0.14
(1,71)	1:38:A:ASP:HB2	1:38:A:ASP:HA	15	0.14
(1,71)	1:38:A:ASP:HB2	1:38:A:ASP:HA	25	0.14
(1,68)	1:52:A:ASN:HB3	1:52:A:ASN:HA	22	0.14
(1,62)	1:18:A:ASP:HB2	1:18:A:ASP:H	5	0.14
(1,62)	1:18:A:ASP:HB2	1:18:A:ASP:H	14	0.14
(1,60)	1:28:A:ASP:HB2	1:28:A:ASP:HA	11	0.14
(1,39)	1:54:A:ARG:H	1:53:A:ASP:HB2	7	0.14
(1,29)	1:43:A:GLY:HA3	1:30:A:ILE:HG22	14	0.14
(1,29)	1:43:A:GLY:HA3	1:30:A:ILE:HG22	25	0.14
(1,966)	1:8:A:ARG:H	1:64:A:GLU:HG3	2	0.13
(1,958)	1:61:A:SER:H	1:5:A:LEU:HD22	4	0.13
(1,954)	1:44:A:CYS:HA	1:45:A:ILE:HD12	25	0.13
(1,954)	1:44:A:CYS:HA	1:45:A:ILE:HD12	28	0.13
(1,944)	1:30:A:ILE:HD11	1:7:A:VAL:HG23	5	0.13
(1,944)	1:30:A:ILE:HD12	1:7:A:VAL:HG23	12	0.13

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,944)	1:30:A:ILE:HD12	1:7:A:VAL:HG23	23	0.13
(1,937)	1:10:A:THR:HG22	1:10:A:THR:H	1	0.13
(1,937)	1:10:A:THR:HG23	1:10:A:THR:H	8	0.13
(1,924)	1:65:A:ALA:HB3	1:5:A:LEU:HD13	13	0.13
(1,917)	1:25:A:LYS:HB3	1:25:A:LYS:H	6	0.13
(1,917)	1:25:A:LYS:HB3	1:25:A:LYS:H	21	0.13
(1,896)	1:48:A:ASN:H	1:47:A:ASP:HB2	18	0.13
(1,894)	1:23:A:ASN:H	1:54:A:ARG:HD2	6	0.13
(1,892)	1:41:A:TRP:HH2	1:61:A:SER:HA	11	0.13
(1,885)	1:62:A:LEU:HB3	1:62:A:LEU:HA	1	0.13
(1,885)	1:62:A:LEU:HB3	1:62:A:LEU:HA	7	0.13
(1,885)	1:62:A:LEU:HB3	1:62:A:LEU:HA	14	0.13
(1,885)	1:62:A:LEU:HB3	1:62:A:LEU:HA	16	0.13
(1,885)	1:62:A:LEU:HB3	1:62:A:LEU:HA	21	0.13
(1,885)	1:62:A:LEU:HB3	1:62:A:LEU:HA	30	0.13
(1,863)	1:9:A:ALA:H	1:29:A:ILE:HA	19	0.13
(1,859)	1:67:A:VAL:H	1:7:A:VAL:HA	12	0.13
(1,858)	1:13:A:TYR:HD2	1:23:A:ASN:HA	23	0.13
(1,858)	1:13:A:TYR:HD2	1:23:A:ASN:HA	26	0.13
(1,849)	1:13:A:TYR:H	1:15:A:ASN:HD22	23	0.13
(1,840)	1:11:A:LYS:HD3	1:12:A:ASP:H	26	0.13
(1,830)	1:9:A:ALA:H	1:29:A:ILE:H	2	0.13
(1,830)	1:23:A:ASN:H	1:45:A:ILE:H	24	0.13
(1,825)	1:34:A:GLU:HB2	1:3:A:HIS:H	1	0.13
(1,819)	1:58:A:PHE:HD1	1:42:A:LYS:H	26	0.13
(1,816)	1:5:A:LEU:HB2	1:7:A:VAL:H	21	0.13
(1,799)	1:41:A:TRP:H	1:41:A:TRP:HB3	24	0.13
(1,799)	1:41:A:TRP:H	1:41:A:TRP:HB3	28	0.13
(1,798)	1:24:A:VAL:HB	1:12:A:ASP:HA	13	0.13
(1,792)	1:47:A:ASP:HB3	1:46:A:HIS:HA	14	0.13
(1,755)	1:10:A:THR:H	1:63:A:GLY:H	11	0.13
(1,709)	1:35:A:GLN:HE21	1:35:A:GLN:HB3	26	0.13
(1,685)	1:27:A:GLY:H	1:24:A:VAL:HG22	14	0.13
(1,673)	1:34:A:GLU:H	1:33:A:LEU:HB2	23	0.13
(1,669)	1:43:A:GLY:H	1:55:A:VAL:HG23	27	0.13
(1,660)	1:10:A:THR:H	1:11:A:LYS:HB2	28	0.13
(1,648)	1:12:A:ASP:H	1:12:A:ASP:HB3	26	0.13
(1,624)	1:18:A:ASP:H	1:17:A:TYR:HA	4	0.13
(1,624)	1:18:A:ASP:H	1:17:A:TYR:HA	7	0.13
(1,624)	1:18:A:ASP:H	1:17:A:TYR:HA	9	0.13
(1,624)	1:18:A:ASP:H	1:17:A:TYR:HA	18	0.13
(1,624)	1:18:A:ASP:H	1:17:A:TYR:HA	23	0.13

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,624)	1:18:A:ASP:H	1:17:A:TYR:HA	29	0.13
(1,598)	1:29:A:ILE:H	1:28:A:ASP:HA	26	0.13
(1,587)	1:53:A:ASP:H	1:53:A:ASP:HB2	17	0.13
(1,583)	1:55:A:VAL:H	1:55:A:VAL:HB	28	0.13
(1,581)	1:13:A:TYR:H	1:13:A:TYR:HA	5	0.13
(1,581)	1:13:A:TYR:H	1:13:A:TYR:HA	24	0.13
(1,566)	1:8:A:ARG:H	1:7:A:VAL:HB	8	0.13
(1,559)	1:64:A:GLU:H	1:64:A:GLU:HA	10	0.13
(1,552)	1:54:A:ARG:H	1:45:A:ILE:HG23	30	0.13
(1,545)	1:40:A:ARG:H	1:36:A:HIS:HB2	13	0.13
(1,527)	1:26:A:ALA:H	1:25:A:LYS:HA	11	0.13
(1,482)	1:46:A:HIS:H	1:30:A:ILE:HG13	17	0.13
(1,458)	1:9:A:ALA:H	1:9:A:ALA:HB1	2	0.13
(1,458)	1:9:A:ALA:H	1:9:A:ALA:HB1	27	0.13
(1,453)	1:32:A:VAL:H	1:32:A:VAL:HG13	1	0.13
(1,453)	1:32:A:VAL:H	1:32:A:VAL:HG12	11	0.13
(1,435)	1:24:A:VAL:HG13	1:30:A:ILE:HG23	9	0.13
(1,432)	1:22:A:LEU:HD11	1:24:A:VAL:HG22	4	0.13
(1,432)	1:22:A:LEU:HD13	1:24:A:VAL:HG23	20	0.13
(1,423)	1:7:A:VAL:HG21	1:32:A:VAL:HG22	4	0.13
(1,395)	1:42:A:LYS:H	1:33:A:LEU:HD11	7	0.13
(1,386)	1:46:A:HIS:H	1:45:A:ILE:HG21	25	0.13
(1,371)	1:28:A:ASP:H	1:10:A:THR:HG22	23	0.13
(1,330)	1:63:A:GLY:H	1:62:A:LEU:HB2	4	0.13
(1,323)	1:15:A:ASN:H	1:15:A:ASN:HB2	2	0.13
(1,323)	1:15:A:ASN:H	1:15:A:ASN:HB2	5	0.13
(1,323)	1:15:A:ASN:H	1:15:A:ASN:HB2	9	0.13
(1,279)	1:63:A:GLY:H	1:62:A:LEU:HA	8	0.13
(1,275)	1:22:A:LEU:HD23	1:22:A:LEU:HA	17	0.13
(1,252)	1:15:A:ASN:H	1:15:A:ASN:HA	24	0.13
(1,232)	1:32:A:VAL:HG13	1:41:A:TRP:HZ3	6	0.13
(1,227)	1:29:A:ILE:H	1:48:A:ASN:HD22	18	0.13
(1,225)	1:35:A:GLN:HG2	1:41:A:TRP:HD1	9	0.13
(1,223)	1:48:A:ASN:HD22	1:48:A:ASN:HB2	1	0.13
(1,223)	1:48:A:ASN:HD22	1:48:A:ASN:HB2	11	0.13
(1,223)	1:48:A:ASN:HB3	1:48:A:ASN:HD22	16	0.13
(1,223)	1:48:A:ASN:HD22	1:48:A:ASN:HB2	19	0.13
(1,223)	1:48:A:ASN:HD22	1:48:A:ASN:HB2	25	0.13
(1,203)	1:52:A:ASN:HD22	1:52:A:ASN:H	10	0.13
(1,182)	1:25:A:LYS:H	1:27:A:GLY:H	20	0.13
(1,58)	1:28:A:ASP:H	1:28:A:ASP:HB2	15	0.13
(1,43)	1:13:A:TYR:HB2	1:24:A:VAL:HG23	28	0.13

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,39)	1:53:A:ASP:HB3	1:54:A:ARG:H	25	0.13
(1,39)	1:53:A:ASP:HB3	1:54:A:ARG:H	26	0.13
(1,967)	1:62:A:LEU:H	1:63:A:GLY:HA2	10	0.12
(1,965)	1:26:A:ALA:H	1:12:A:ASP:HA	9	0.12
(1,952)	1:23:A:ASN:H	1:45:A:ILE:HD11	18	0.12
(1,937)	1:10:A:THR:HG21	1:10:A:THR:H	21	0.12
(1,900)	1:14:A:CYS:HB3	1:24:A:VAL:H	6	0.12
(1,885)	1:62:A:LEU:HB3	1:62:A:LEU:HA	2	0.12
(1,885)	1:62:A:LEU:HB3	1:62:A:LEU:HA	3	0.12
(1,885)	1:62:A:LEU:HB3	1:62:A:LEU:HA	4	0.12
(1,885)	1:62:A:LEU:HB3	1:62:A:LEU:HA	5	0.12
(1,885)	1:62:A:LEU:HB3	1:62:A:LEU:HA	6	0.12
(1,885)	1:62:A:LEU:HB3	1:62:A:LEU:HA	8	0.12
(1,885)	1:62:A:LEU:HB3	1:62:A:LEU:HA	10	0.12
(1,885)	1:62:A:LEU:HB3	1:62:A:LEU:HA	11	0.12
(1,885)	1:62:A:LEU:HB3	1:62:A:LEU:HA	12	0.12
(1,885)	1:62:A:LEU:HB3	1:62:A:LEU:HA	13	0.12
(1,885)	1:62:A:LEU:HB3	1:62:A:LEU:HA	15	0.12
(1,885)	1:62:A:LEU:HB3	1:62:A:LEU:HA	17	0.12
(1,885)	1:62:A:LEU:HB3	1:62:A:LEU:HA	18	0.12
(1,885)	1:62:A:LEU:HB3	1:62:A:LEU:HA	19	0.12
(1,885)	1:62:A:LEU:HB3	1:62:A:LEU:HA	20	0.12
(1,885)	1:62:A:LEU:HB3	1:62:A:LEU:HA	22	0.12
(1,885)	1:62:A:LEU:HB3	1:62:A:LEU:HA	23	0.12
(1,885)	1:62:A:LEU:HB3	1:62:A:LEU:HA	24	0.12
(1,885)	1:62:A:LEU:HB3	1:62:A:LEU:HA	25	0.12
(1,885)	1:62:A:LEU:HB3	1:62:A:LEU:HA	26	0.12
(1,885)	1:62:A:LEU:HB3	1:62:A:LEU:HA	27	0.12
(1,885)	1:62:A:LEU:HG	1:62:A:LEU:HA	28	0.12
(1,885)	1:62:A:LEU:HB3	1:62:A:LEU:HA	29	0.12
(1,881)	1:13:A:TYR:HD2	1:21:A:SER:HB3	28	0.12
(1,873)	1:41:A:TRP:HB3	1:34:A:GLU:HA	18	0.12
(1,872)	1:5:A:LEU:HD21	1:4:A:MET:HA	10	0.12
(1,853)	1:29:A:ILE:HG12	1:48:A:ASN:HD22	5	0.12
(1,840)	1:11:A:LYS:HD3	1:12:A:ASP:H	1	0.12
(1,815)	1:44:A:CYS:H	1:30:A:ILE:HG22	12	0.12
(1,802)	1:26:A:ALA:HB1	1:12:A:ASP:HA	30	0.12
(1,799)	1:41:A:TRP:H	1:41:A:TRP:HB3	26	0.12
(1,792)	1:47:A:ASP:HB3	1:46:A:HIS:HA	22	0.12
(1,774)	1:42:A:LYS:H	1:34:A:GLU:H	9	0.12
(1,738)	1:50:A:THR:H	1:50:A:THR:HB	17	0.12
(1,738)	1:50:A:THR:H	1:50:A:THR:HB	20	0.12

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,738)	1:50:A:THR:H	1:50:A:THR:HB	30	0.12
(1,705)	1:35:A:GLN:HE22	1:33:A:LEU:HD21	10	0.12
(1,660)	1:10:A:THR:H	1:11:A:LYS:HB2	16	0.12
(1,653)	1:17:A:TYR:H	1:16:A:ASN:HB2	2	0.12
(1,624)	1:18:A:ASP:H	1:17:A:TYR:HA	10	0.12
(1,624)	1:18:A:ASP:H	1:17:A:TYR:HA	11	0.12
(1,624)	1:18:A:ASP:H	1:17:A:TYR:HA	15	0.12
(1,624)	1:18:A:ASP:H	1:17:A:TYR:HA	16	0.12
(1,624)	1:18:A:ASP:H	1:17:A:TYR:HA	21	0.12
(1,624)	1:18:A:ASP:H	1:17:A:TYR:HA	28	0.12
(1,595)	1:62:A:LEU:H	1:62:A:LEU:HB2	3	0.12
(1,583)	1:55:A:VAL:H	1:55:A:VAL:HB	7	0.12
(1,583)	1:55:A:VAL:H	1:55:A:VAL:HB	11	0.12
(1,581)	1:13:A:TYR:H	1:13:A:TYR:HA	7	0.12
(1,560)	1:64:A:GLU:H	1:64:A:GLU:HG2	8	0.12
(1,524)	1:15:A:ASN:H	1:15:A:ASN:HB3	7	0.12
(1,524)	1:15:A:ASN:H	1:15:A:ASN:HB3	10	0.12
(1,482)	1:46:A:HIS:H	1:30:A:ILE:HG13	25	0.12
(1,467)	1:30:A:ILE:H	1:29:A:ILE:HG23	5	0.12
(1,453)	1:32:A:VAL:H	1:32:A:VAL:HG12	4	0.12
(1,453)	1:32:A:VAL:H	1:32:A:VAL:HG13	14	0.12
(1,453)	1:32:A:VAL:H	1:32:A:VAL:HG12	24	0.12
(1,435)	1:24:A:VAL:HG12	1:30:A:ILE:HG21	14	0.12
(1,422)	1:7:A:VAL:HG21	1:32:A:VAL:HG11	5	0.12
(1,414)	1:5:A:LEU:HD12	1:32:A:VAL:HG12	1	0.12
(1,396)	1:44:A:CYS:H	1:33:A:LEU:HD12	8	0.12
(1,395)	1:42:A:LYS:H	1:33:A:LEU:HD12	22	0.12
(1,392)	1:43:A:GLY:H	1:33:A:LEU:HD12	1	0.12
(1,391)	1:34:A:GLU:H	1:5:A:LEU:HD21	13	0.12
(1,386)	1:46:A:HIS:H	1:45:A:ILE:HG22	4	0.12
(1,377)	1:6:A:GLN:HA	1:32:A:VAL:HG12	24	0.12
(1,367)	1:9:A:ALA:HB3	1:10:A:THR:H	2	0.12
(1,367)	1:9:A:ALA:HB2	1:10:A:THR:H	19	0.12
(1,252)	1:15:A:ASN:H	1:15:A:ASN:HA	27	0.12
(1,232)	1:32:A:VAL:HG11	1:41:A:TRP:HZ3	27	0.12
(1,231)	1:33:A:LEU:H	1:41:A:TRP:HZ3	19	0.12
(1,228)	1:51:A:GLY:H	1:52:A:ASN:HD22	24	0.12
(1,223)	1:48:A:ASN:HB3	1:48:A:ASN:HD22	3	0.12
(1,223)	1:48:A:ASN:HD22	1:48:A:ASN:HB2	5	0.12
(1,223)	1:48:A:ASN:HD22	1:48:A:ASN:HB2	7	0.12
(1,223)	1:48:A:ASN:HB3	1:48:A:ASN:HD22	9	0.12
(1,223)	1:48:A:ASN:HD22	1:48:A:ASN:HB2	10	0.12

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,223)	1:48:A:ASN:HD22	1:48:A:ASN:HB2	20	0.12
(1,223)	1:48:A:ASN:HB3	1:48:A:ASN:HD22	24	0.12
(1,223)	1:48:A:ASN:HB3	1:48:A:ASN:HD22	27	0.12
(1,223)	1:48:A:ASN:HD22	1:48:A:ASN:HB2	28	0.12
(1,223)	1:48:A:ASN:HB3	1:48:A:ASN:HD22	29	0.12
(1,74)	1:15:A:ASN:HD22	1:15:A:ASN:HB2	30	0.12
(1,66)	1:28:A:ASP:HB3	1:25:A:LYS:HG2	15	0.12
(1,57)	1:28:A:ASP:HB2	1:25:A:LYS:HG3	26	0.12
(1,41)	1:13:A:TYR:HB2	1:24:A:VAL:HB	4	0.12
(1,20)	1:11:A:LYS:H	1:26:A:ALA:HA	10	0.12
(1,967)	1:62:A:LEU:H	1:63:A:GLY:HA2	17	0.11
(1,965)	1:26:A:ALA:H	1:12:A:ASP:HA	23	0.11
(1,937)	1:10:A:THR:HG22	1:10:A:THR:H	7	0.11
(1,931)	1:41:A:TRP:HZ2	1:60:A:SER:HB3	30	0.11
(1,925)	1:11:A:LYS:HG2	1:13:A:TYR:H	26	0.11
(1,924)	1:65:A:ALA:HB3	1:5:A:LEU:HD12	6	0.11
(1,924)	1:65:A:ALA:HB3	1:5:A:LEU:HD13	12	0.11
(1,909)	1:41:A:TRP:HD1	1:40:A:ARG:HG2	20	0.11
(1,900)	1:14:A:CYS:HB3	1:24:A:VAL:H	28	0.11
(1,894)	1:53:A:ASP:H	1:54:A:ARG:HD2	23	0.11
(1,858)	1:13:A:TYR:HD2	1:23:A:ASN:HA	5	0.11
(1,858)	1:13:A:TYR:HD2	1:23:A:ASN:HA	19	0.11
(1,855)	1:14:A:CYS:HB3	1:23:A:ASN:HD22	14	0.11
(1,836)	1:40:A:ARG:H	1:41:A:TRP:HD1	15	0.11
(1,830)	1:23:A:ASN:H	1:45:A:ILE:H	23	0.11
(1,815)	1:44:A:CYS:H	1:30:A:ILE:HG22	5	0.11
(1,810)	1:23:A:ASN:H	1:45:A:ILE:H	12	0.11
(1,793)	1:28:A:ASP:HB3	1:24:A:VAL:HG11	15	0.11
(1,709)	1:35:A:GLN:HE21	1:35:A:GLN:HB3	8	0.11
(1,709)	1:35:A:GLN:HE21	1:35:A:GLN:HB3	12	0.11
(1,673)	1:34:A:GLU:H	1:33:A:LEU:HB2	27	0.11
(1,669)	1:43:A:GLY:H	1:55:A:VAL:HG22	10	0.11
(1,669)	1:43:A:GLY:H	1:55:A:VAL:HG21	18	0.11
(1,660)	1:10:A:THR:H	1:11:A:LYS:HB2	7	0.11
(1,633)	1:52:A:ASN:H	1:51:A:GLY:H	30	0.11
(1,597)	1:62:A:LEU:H	1:62:A:LEU:HB3	23	0.11
(1,590)	1:42:A:LYS:H	1:41:A:TRP:HB2	14	0.11
(1,589)	1:42:A:LYS:H	1:42:A:LYS:HB3	15	0.11
(1,581)	1:13:A:TYR:H	1:13:A:TYR:HA	28	0.11
(1,567)	1:8:A:ARG:H	1:8:A:ARG:HD3	11	0.11
(1,566)	1:8:A:ARG:H	1:7:A:VAL:HB	18	0.11
(1,559)	1:64:A:GLU:H	1:64:A:GLU:HA	3	0.11

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,554)	1:54:A:ARG:H	1:54:A:ARG:HB2	12	0.11
(1,524)	1:15:A:ASN:H	1:15:A:ASN:HB3	23	0.11
(1,524)	1:15:A:ASN:H	1:15:A:ASN:HB3	26	0.11
(1,515)	1:14:A:CYS:H	1:13:A:TYR:HB3	18	0.11
(1,505)	1:22:A:LEU:H	1:22:A:LEU:HB2	17	0.11
(1,458)	1:9:A:ALA:H	1:9:A:ALA:HB2	7	0.11
(1,458)	1:9:A:ALA:H	1:9:A:ALA:HB3	24	0.11
(1,458)	1:9:A:ALA:H	1:9:A:ALA:HB3	26	0.11
(1,453)	1:32:A:VAL:H	1:32:A:VAL:HG12	17	0.11
(1,453)	1:32:A:VAL:H	1:32:A:VAL:HG11	26	0.11
(1,453)	1:32:A:VAL:H	1:32:A:VAL:HG11	30	0.11
(1,422)	1:7:A:VAL:HG21	1:32:A:VAL:HG13	28	0.11
(1,418)	1:7:A:VAL:HG11	1:32:A:VAL:HG21	25	0.11
(1,368)	1:8:A:ARG:HD2	1:8:A:ARG:HG2	11	0.11
(1,367)	1:9:A:ALA:HB2	1:10:A:THR:H	11	0.11
(1,367)	1:9:A:ALA:HB3	1:10:A:THR:H	21	0.11
(1,367)	1:9:A:ALA:HB3	1:10:A:THR:H	27	0.11
(1,367)	1:9:A:ALA:HB1	1:10:A:THR:H	28	0.11
(1,341)	1:29:A:ILE:H	1:29:A:ILE:HG12	7	0.11
(1,341)	1:29:A:ILE:H	1:29:A:ILE:HG12	22	0.11
(1,334)	1:49:A:ARG:H	1:49:A:ARG:HB3	28	0.11
(1,279)	1:63:A:GLY:H	1:62:A:LEU:HA	9	0.11
(1,244)	1:30:A:ILE:HG12	1:8:A:ARG:HA	9	0.11
(1,232)	1:32:A:VAL:HG11	1:41:A:TRP:HZ3	8	0.11
(1,223)	1:48:A:ASN:HB3	1:48:A:ASN:HD22	2	0.11
(1,223)	1:48:A:ASN:HB3	1:48:A:ASN:HD22	6	0.11
(1,223)	1:48:A:ASN:HD22	1:48:A:ASN:HB2	12	0.11
(1,223)	1:48:A:ASN:HB3	1:48:A:ASN:HD22	14	0.11
(1,223)	1:48:A:ASN:HB3	1:48:A:ASN:HD22	18	0.11
(1,223)	1:48:A:ASN:HB3	1:48:A:ASN:HD22	21	0.11
(1,223)	1:48:A:ASN:HD22	1:48:A:ASN:HB2	22	0.11
(1,223)	1:48:A:ASN:HD22	1:48:A:ASN:HB2	23	0.11
(1,211)	1:25:A:LYS:H	1:11:A:LYS:H	3	0.11
(1,211)	1:25:A:LYS:H	1:11:A:LYS:H	10	0.11
(1,211)	1:25:A:LYS:H	1:11:A:LYS:H	12	0.11
(1,203)	1:52:A:ASN:HD22	1:52:A:ASN:H	29	0.11
(1,137)	1:8:A:ARG:HA	1:29:A:ILE:HA	26	0.11
(1,124)	1:26:A:ALA:HB3	1:11:A:LYS:H	28	0.11
(1,97)	1:6:A:GLN:HB2	1:6:A:GLN:HG3	6	0.11
(1,71)	1:38:A:ASP:HB2	1:38:A:ASP:HA	19	0.11
(1,58)	1:28:A:ASP:H	1:28:A:ASP:HB2	14	0.11
(1,35)	1:27:A:GLY:HA3	1:28:A:ASP:H	27	0.11

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Key	Atom-1	Atom-2	Model ID	Violation (Å)
(1,964)	1:45:A:ILE:H	1:53:A:ASP:HA	6	0.1
(1,952)	1:23:A:ASN:H	1:45:A:ILE:HD12	23	0.1
(1,932)	1:9:A:ALA:H	1:9:A:ALA:HB3	20	0.1
(1,924)	1:65:A:ALA:HB1	1:5:A:LEU:HD11	4	0.1
(1,863)	1:9:A:ALA:H	1:29:A:ILE:HA	2	0.1
(1,853)	1:29:A:ILE:HG12	1:48:A:ASN:HD22	28	0.1
(1,845)	1:9:A:ALA:HB3	1:11:A:LYS:H	24	0.1
(1,815)	1:44:A:CYS:H	1:30:A:ILE:HG22	19	0.1
(1,734)	1:39:A:GLY:H	1:40:A:ARG:HB2	21	0.1
(1,720)	1:51:A:GLY:H	1:50:A:THR:H	12	0.1
(1,705)	1:35:A:GLN:HE22	1:33:A:LEU:HD23	6	0.1
(1,653)	1:17:A:TYR:H	1:16:A:ASN:HB2	14	0.1
(1,624)	1:18:A:ASP:H	1:17:A:TYR:HA	1	0.1
(1,590)	1:42:A:LYS:H	1:41:A:TRP:HB2	7	0.1
(1,590)	1:42:A:LYS:H	1:41:A:TRP:HB2	26	0.1
(1,583)	1:55:A:VAL:H	1:55:A:VAL:HB	5	0.1
(1,581)	1:13:A:TYR:H	1:13:A:TYR:HA	6	0.1
(1,482)	1:46:A:HIS:H	1:30:A:ILE:HG13	3	0.1
(1,458)	1:9:A:ALA:H	1:9:A:ALA:HB3	5	0.1
(1,458)	1:9:A:ALA:H	1:9:A:ALA:HB2	15	0.1
(1,450)	1:25:A:LYS:H	1:24:A:VAL:HG12	5	0.1
(1,422)	1:7:A:VAL:HG23	1:32:A:VAL:HG12	8	0.1
(1,392)	1:43:A:GLY:H	1:33:A:LEU:HD12	14	0.1
(1,389)	1:11:A:LYS:HB3	1:24:A:VAL:HG11	21	0.1
(1,382)	1:65:A:ALA:H	1:5:A:LEU:HB3	20	0.1
(1,368)	1:8:A:ARG:HD2	1:8:A:ARG:HG2	5	0.1
(1,368)	1:8:A:ARG:HD2	1:8:A:ARG:HG2	7	0.1
(1,368)	1:8:A:ARG:HD2	1:8:A:ARG:HG2	9	0.1
(1,368)	1:8:A:ARG:HD2	1:8:A:ARG:HG2	30	0.1
(1,348)	1:41:A:TRP:HZ3	1:34:A:GLU:HB2	19	0.1
(1,334)	1:49:A:ARG:H	1:49:A:ARG:HB3	23	0.1
(1,323)	1:15:A:ASN:H	1:15:A:ASN:HB2	3	0.1
(1,237)	1:43:A:GLY:H	1:57:A:TYR:HA	6	0.1
(1,234)	1:8:A:ARG:HG3	1:6:A:GLN:HE22	16	0.1
(1,223)	1:48:A:ASN:HB3	1:48:A:ASN:HD22	26	0.1
(1,190)	1:34:A:GLU:HG2	1:3:A:HIS:H	20	0.1
(1,107)	1:41:A:TRP:HB2	1:41:A:TRP:HD1	4	0.1
(1,97)	1:6:A:GLN:HB2	1:6:A:GLN:HG3	19	0.1
(1,97)	1:6:A:GLN:HB2	1:6:A:GLN:HG3	28	0.1
(1,22)	1:6:A:GLN:H	1:6:A:GLN:HA	4	0.1

10 Dihedral-angle violation analysis ⓘ

No dihedral-angle restraints found