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All-Atom Contacts	Clashscore, all atoms:	104.16		0 th percentile* (N=1784, all resolutions)
	Clashscore is the number of serious steric overlaps (> 0.4 Å) per 1000 atoms.			
Protein Geometry	Poor rotamers	8	3.90%	Goal: <1%
	Ramachandran outliers	2	0.88%	Goal: <0.05%
	Ramachandran favored	221	97.36%	Goal: >98%
	MolProbity score [^]	3.06		21 st percentile* (N=27675, 0Å - 99Å)
	Cβ deviations >0.25Å	2	0.92%	Goal: 0
	Bad backbone bonds:	0 / 1906	0.00%	Goal: 0%
	Bad backbone angles:	33 / 2575	1.28%	Goal: <0.1%

In the two column results, the left column gives the raw count, right column gives the percentage.

* 100th percentile is the best among structures of comparable resolution; 0th percentile is the worst. For clashscore the comparative set of structures was selected in 2004, for MolProbity score in 2006.

[^] MolProbity score combines the clashscore, rotamer, and Ramachandran evaluations into a single score, normalized to be on the same scale as X-ray resolution.

#	Alt Res	High B	Clash > 0.4Å	Ramachandran	Rotamer	Cβ deviation	Bond lengths	Bond angles
		Avg: 83.62	Clashscore: 104.16	Outliers: 2 of 227	Poor rotamers: 8 of 205	Outliers: 2 of 218	Outliers: 0 of 229	Outliers: 31 of 229
20	ILE	23.64	0.653Å HD13 with 248 GLN HG2	-	87.9% (mt) chi angles: 299,168.2	0.087Å	-	-
21	LYS	50.69	0.799Å HB2 with 24 PRO HD2	Favored (49.3%) General / -64.5,149.0	60.3% (mtpt) chi angles: 295.8,165.7,66.4,179.3	0.081Å	-	-
22	ALA	19.75	0.91Å HA with 25 MET CE	Favored (95.54%) General / -61.6,-40.6	-	0.087Å	-	-
23	GLY	22.3	-	Favored (99.7%) Glycine / -62.5,-42.5	-	-	-	-

24	PRO	125.66	0.799Å HD2 with 21 LYS HB2	Favored (31.74%) Trans-proline / -70.8,-19.0	87.6% (<i>Cg_exo</i>) chi angles: 331.3	0.189Å	-	-
25	MET	46.45	0.91Å CE with 22 ALA HA	Favored (90.77%) Pre-proline / -73.0,155.0	69.6% (<i>mtm</i>) chi angles: 298.2,175.5,291.4	0.109Å	-	-
26	PRO	86.37	0.586Å HD2 with 244 ILE HG23	Favored (20.4%) Trans-proline / -73.0,133.8	10.2% (<i>Cg_endo</i>) chi angles: 13.3	0.098Å	-	-
27	LYS	38.12	-	Favored (70%) General / -70.9,-42.3	81.7% (<i>tttt</i>) chi angles: 181.7,166,180,179.8	0.091Å	-	-
28	HIS	56.4	-	Favored (40.2%) General / -129.7,127.3	10.3% (<i>t-160</i>) chi angles: 187,171.3	0.094Å	-	OUTLIER(S) worst is CD2- NE2-CE1: 4.417 σ
29	ILE	39.9	0.882Å HG12 with 202 ILE CG2	Favored (69.74%) Isoleucine or valine / -119.0,132.2	82.2% (<i>mt</i>) chi angles: 300.3,168.3	0.081Å	-	-
30	ALA	35.16	0.838Å HA with 74 THR HG22	Favored (51.92%) General / -111.8,135.8	-	0.115Å	-	-
31	PHE	57.83	0.702Å CE2 with 240 LEU HD13	Favored (52.2%) General / -123.2,130.0	83.1% (<i>m-85</i>) chi angles: 299.2,83.2	0.161Å	-	OUTLIER(S) worst is CA- CB-CG: 4.51 σ
32	ILE	159.94	0.696Å HG23 with 205 ARG HG3	Favored (22.09%) Isoleucine or valine / -105.4,107.2	0% chi angles: 185.6,293.7	0.053Å	-	-
33	MET	80.72	0.759Å HE3 with 75 VAL HG11	Favored (30.09%) General / -77.3,125.9	63.3% (<i>mtt</i>) chi angles: 299.2,171.9,179.9	0.106Å	-	-
34	ASP	73.54	-	Favored (43.05%) General / -148.8,156.9	38% (<i>t0</i>) chi angles: 197,2.4	0.086Å	-	-
35	GLY	38.43	0.665Å HA2 with 38 ARG HD3	Favored (44.26%) Glycine /	-	-	-	-

36	ASN	43.14	-	104.3,3.8 Favored (97.94%) General / -61.7,-43.9	97.8% (<i>m-20</i>) chi angles: 291.7,339.2	0.121Å	-	-
37	ARG	94.09	-	Favored (84.45%) General / -64.8,-45.9	79.1% (<i>ttt180</i>) chi angles: 179.9,182.2,179.9,184	0.087Å	-	-
38	ARG	107.27	0.666Å NH1 with 207 SER HB2	Favored (92.64%) General / -63.8,-38.9	2% (<i>mpt_?</i>) chi angles: 300.1,72.3,171.6,272	0.131Å	-	-
39	TYR	37.54	0.469Å HD1 with 234 GLU OE2	Favored (96.71%) General / -61.1,-44.9	89.9% (<i>t80</i>) chi angles: 176.2,79.5	0.076Å	-	-
#	Alt Res	High B	Clash > 0.4Å	Ramachandran	Rotamer	Cβ deviation	Bond lengths	Bond angles
		Avg: 83.62	Clashscore: 104.16	Outliers: 2 of 227	Poor rotamers: 8 of 205	Outliers: 2 of 218	Outliers: 0 of 229	Outliers: 31 of 229
40	ALA	27.59	-	Favored (96.07%) General / -64.7,-41.3	-	0.087Å	-	-
41	LYS	91.73	-	Favored (85.12%) General / -59.6,-39.7	58.9% (<i>tttm</i>) chi angles: 180,166.9,178.1,291	0.083Å	-	-
42	LYS	91.28	-	Favored (88.5%) General / -59.5,-40.9	31.1% (<i>ttmt</i>) chi angles: 180,167.4,277.7,180	0.101Å	-	-
43	CYS	58.04	0.523Å O with 44 GLN HB2	Favored (62.21%) General / -71.7,-17.7	71.6% (<i>m</i>) chi angles: 297.3	0.066Å	-	-
44	GLN	65.91	0.523Å HB2 with 43 CYS O	Favored (2.63%) General / 67.5,40.3	25% (<i>mt-30</i>) chi angles: 296.2,175.6,171.5	0.07Å	-	-
45	VAL	38.5	0.59Å HG12 with 46 GLU N	Favored (15.65%) Isoleucine or valine / -117.6,162.4	64% (<i>t</i>) chi angles: 179.9	0.071Å	-	-
			0.59Å	Favored	18.7% (<i>pt-20</i>)			

46	GLU	78.33	N with 45 VAL HG12	(47.02%) General / -63.8,149.4 Favored	chi angles: 60.1,180.8,338.2	0.086Å	-	-
47	ARG	185.53	0.671Å HD2 with 89 GLU CD	(69.57%) General / -65.5,-29.0 Favored	5% (<i>ptp180</i>) chi angles: 60.8,189,89.2,141.8	0.144Å	-	-
48	GLN	47.19	0.935Å HA with 51 HIS HD2	(69.21%) General / -59.2,-31.6 Favored	89.4% (<i>mt-30</i>) chi angles: 296.3,180.1,353.3	0.125Å	-	-
49	GLU	108.83	0.544Å HG2 with 45 VAL HG11	(92.56%) General / -62.5,-39.2 Favored	11.9% (<i>tp10</i>) chi angles: 192.7,71.6,61.4	0.074Å	-	-
50	GLY	21.77	-	(96.17%) Glycine / -63.6,-38.4 Favored	-	-	-	-
51	HIS	66.26	0.935Å HD2 with 48 GLN HA	(99.52%) General / -62.9,-41.7 Favored	81.9% (<i>m-70</i>) chi angles: 304,301.7	0.082Å	-	OUTLIER(S) worst is CD2- NE2-CE1: 4.522 σ
52	SER	73.09	0.461Å O with 55 PHE HD2	(78.84%) General / -61.0,-36.2 Favored	69.2% (<i>m</i>) chi angles: 296	0.078Å	-	-
53	GLN	81.23	0.835Å HG2 with 233 PRO HB3	(70.76%) General / -64.8,-29.8 Favored	34.3% (<i>tp60</i>) chi angles: 180.2,75.2,8.8	0.083Å	-	-
54	GLY	44.78	-	(73.58%) Glycine / -61.6,-30.0 Favored	-	-	-	-
55	PHE	104.14	0.461Å HD2 with 52 SER O	(96.32%) General / -61.3,-45.0 Favored	11.9% (<i>p90</i>) chi angles: 79.3,84.6	0.144Å	-	-
56	ASN	65.29	0.46Å N with 55 PHE CG	(90.02%) General / -61.9,-38.9 Favored	32.7% (<i>t-20</i>) chi angles: 191,309	0.082Å	-	OUTLIER(S) worst is CA- CB-CG: 4.048 σ
57	LYS	132.64	0.643Å NZ with 61 THR HB	(97.43%) General / -62.2,-40.8 Favored	0.1% chi angles: 201.5,305.1,101.8,174.4	0.088Å	-	-

58	LEU	46.79	0.783Å O with 61 THR HG22	(91.9%) General / -60.9,-46.1	61.6% (<i>tp</i>) chi angles: 180,63.8	0.088Å	-	-
59	ALA	31.08	-	Favored (96.02%) General / -64.0,-40.3	-	0.086Å	-	-
#	Alt Res	High B	Clash > 0.4Å	Ramachandran	Rotamer	Cβ deviation	Bond lengths	Bond angles
		Avg: 83.62	Clashscore: 104.16	Outliers: 2 of 227	Poor rotamers: 8 of 205	Outliers: 2 of 218	Outliers: 0 of 229	Outliers: 31 of 229
60	GLU	96.63	-	Favored (99.26%) General / -62.5,-42.0	23.4% (<i>tp10</i>) chi angles: 189.3,76.3,352.9	0.087Å	-	-
61	THR	99.88	0.783Å HG22 with 58 LEU O	Favored (83.24%) General / -67.4,-37.5	8.3% (<i>t</i>) chi angles: 193.3	0.074Å	-	-
62	LEU	92.65	0.743Å HD13 with 58 LEU O	Favored (96.68%) General / -60.5,-43.7	8.3% (<i>mp</i>) chi angles: 276.1,51.4	0.099Å	-	-
63	ARG	96.87	-	Favored (98.53%) General / -62.8,-43.5	59.1% (<i>ttm-85</i>) chi angles: 182.3,184.4,300,271.3	0.092Å	-	-
64	TRP	141.47	1.038Å HB2 with 237 PHE HB2	Favored (89.85%) General / -66.0,-39.0	76.6% (<i>t-105</i>) chi angles: 179.9,254.1	0.141Å	-	-
65	CYS	29.36	-	Favored (93.96%) General / -62.7,-39.4	76.6% (<i>m</i>) chi angles: 296.4	0.09Å	-	-
66	LEU	119.74	0.741Å HB2 with 114 HIS CE1	Favored (94.91%) General / -63.5,-44.2	55.8% (<i>tp</i>) chi angles: 179.8,66.1	0.122Å	-	-
67	ASN	93.22	0.65Å OD1 with 64 TRP HA	Favored (91.94%) General / -64.8,-38.7	1.5% (<i>p-10</i>) chi angles: 64.5,258.7	0.114Å	-	-
68	LEU	108.71	0.786Å O with 68 LEU HD23	Favored (72.7%) General / -68.4,-32.6	6.1% (<i>tt</i>) chi angles: 192.7,168.6	0.079Å	-	-

69	GLY	38.41	-	Favored (13.98%) Glycine / 102.0,19.9	-	-	-	-
70	ILE	126.46	0.814Å HD11 with 244 ILE HD13	Favored (33.22%) Isoleucine or valine / -70.0,128.4	34% (<i>mm</i>) chi angles: 300.8,292.9	0.14Å	-	-
71	LEU	69.62	0.55Å HB2 with 72 GLU OE1	Favored (49.05%) General / -75.3,-43.9	87.3% (<i>mt</i>) chi angles: 297.9,180.8	0.09Å	-	-
72	GLU	81.06	0.55Å OE1 with 71 LEU HB2	Favored (46.3%) General / -133.5,138.9	58.8% (<i>mp0</i>) chi angles: 288.6,79.2,4.5	0.095Å	-	-
73	VAL	45.43	0.632Å HB with 150 LEU CD2	Favored (36.69%) Isoleucine or valine / -139.1,131.4	98.3% (<i>t</i>) chi angles: 176.6	0.08Å	-	-
74	THR	109.79	1.111Å HG21 with 216 LEU HD11	Favored (56.25%) General / -118.5,131.8	15.1% (<i>t</i>) chi angles: 188.1	0.063Å	-	-
75	VAL	46.1	0.759Å HG11 with 33 MET HE3	Favored (42.45%) Isoleucine or valine / -117.1,139.8	78.5% (<i>t</i>) chi angles: 172.4	0.086Å	-	-
76	TYR	169.54	0.553Å CD1 with 215 PHE CD1	Favored (2.52%) General / -103.4,85.7	3.3% (<i>t80</i>) chi angles: 198.7,38	0.084Å	-	-
77	ALA	66.17	-	Favored (71.06%) General / -70.6,-42.4	-	0.059Å	-	-
78	PHE	112.99	0.799Å HB2 with 154 PHE CE2	Favored (12.86%) General / -158.9,139.8	43.5% (<i>t80</i>) chi angles: 191.5,89.3	0.118Å	-	-
79	SER	132.05	0.427Å HB3 with 76 TYR OH	Favored (15.34%) General / -107.3,160.8	3.1% (<i>m</i>) chi angles: 317.7	0.278Å	-	OUTLIER(S) worst is CB- CA-C: 5.039 σ

Clash >

Cβ

Bond

#	Alt Res	High B	0.4Å	Ramachandran	Rotamer	deviation	lengths	Bond angles
		Avg: 83.62	Clashscore: 104.16	Outliers: 2 of 227	Poor rotamers: 8 of 205	Outliers: 2 of 218	Outliers: 0 of 229	Outliers: 31 of 229
80	ILE	138.09	0.718Å HD13 with 157 THR CB	Favored (44.91%) Isoleucine or valine / -61.9,-28.7	2% (<i>mp</i>) chi angles: 286.1,67.7	0.146Å	-	-
81	GLU	170.75	-	Favored (88.69%) General / -65.1,-37.8	0.9% chi angles: 233.8,71.8,63.4	0.132Å	-	OUTLIER(S) worst is CB-CG-CD: 5.596 σ
82	ASN	216.87	-	Allowed (0.42%) General / -82.2,29.4	32.3% (<i>p-10</i>) chi angles: 54.2,352	0.275Å	-	-
83	PHE	181.42	-	Favored (64.03%) General / -71.5,-28.6	14.9% (<i>t80</i>) chi angles: 199.7,278.5	0.191Å	-	-
84	LYS	170.56	0.614Å HB2 with 84 LYS NZ	Favored (73.7%) General / -59.3,-35.4	0% chi angles: 204.4,77.5,65.1,289.6	0.126Å	-	-
85	ARG	171.09	0.934Å HH21 with 93 LEU HD11	Favored (3.46%) General / -64.5,171.0	27.5% (<i>mtm180</i>) chi angles: 292.2,211.3,287.4,158.1	0.134Å	-	OUTLIER(S) worst is CD-NE-CZ: 4.556 σ
86	SER	85.67	-	Favored (16.63%) General / -58.5,153.2	50.7% (<i>m</i>) chi angles: 301.6	0.17Å	-	-
87	LYS	125.34	0.561Å HA with 90 VAL HG22	Favored (75.3%) General / -65.3,-33.1	9.8% (<i>ptmt</i>) chi angles: 67.3,172,315.6,160.1	0.086Å	-	-
88	SER	32.87	-	Favored (96.37%) General / -61.8,-44.7	74.1% (<i>m</i>) chi angles: 294	0.082Å	-	-
89	GLU	163.12	0.671Å CD with 47 ARG HD2	Favored (85.31%) General / -64.4,-46.0	76.8% (<i>mm-40</i>) chi angles: 292.5,293.5,313.4	0.21Å	-	-
90	VAL	104.43	0.561Å HG22 with 87 LYS HA	Favored (91.24%) Isoleucine or valine / -65.0,-41.1	33.5% (<i>m</i>) chi angles: 297.2	0.078Å	-	-

91	ASP	69.22	N with 90 VAL HG23	0.532Å	Favored (98.04%) General / -61.5,-43.9	51.7% (<i>t0</i>) chi angles: 189.1,11.2	0.095Å	-	OUTLIER(S) worst is CA- CB-CG: 4.865 σ
92	GLY	22.82	O with 96 LEU HG	0.623Å	Favored (97.94%) Glycine / -63.9,-43.1	-	-	-	-
93	LEU	66.25	HD11 with 85 ARG HH21	0.934Å	Favored (95.26%) General / -64.0,-40.0	83.9% (<i>mt</i>) chi angles: 301.2,180	0.048Å	-	-
94	MET	109.79	HB3 with 128 PRO HG3	0.794Å	Favored (82.04%) General / -67.3,-43.3	65.9% (<i>mtt</i>) chi angles: 294.7,180,177.3	0.132Å	-	OUTLIER(S) worst is CG- SD-CE: 4.327 σ
95	ASP	72.33	-	-	Favored (90.21%) General / -66.2,-40.2	62.4% (<i>t0</i>) chi angles: 187.3,1.9	0.1Å	-	-
96	LEU	135.95	HG with 92 GLY O	0.623Å	Favored (95.48%) General / -63.7,-43.8	76.8% (<i>mt</i>) chi angles: 292.5,180.3	0.072Å	-	-
97	ALA	27.64	HA with 101 PHE HB3	0.429Å	Favored (97.77%) General / -63.8,-42.5	-	0.084Å	-	-
98	ARG	98.95	HG2 with 134 LEU HD12	0.983Å	Favored (78.68%) General / -63.5,-48.2	28.5% (<i>ttp180</i>) chi angles: 180,212.1,71.3,203.3	0.133Å	-	-
99	GLN	110.6	O with 105 MET HB3	0.548Å	Favored (88.32%) General / -63.5,-45.9	8.9% (<i>tt0</i>) chi angles: 186,175.2,185	0.059Å	-	OUTLIER(S) worst is C-N- CA: 5.634 σ
#	Alt Res	High B		Clash > 0.4Å	Ramachandran	Rotamer	Cβ deviation	Bond lengths	Bond angles
		Avg: 83.62		Clashscore: 104.16	Outliers: 2 of 227	Poor rotamers: 8 of 205	Outliers: 2 of 218	Outliers: 0 of 229	Outliers: 31 of 229
100	LYS	246.81	-	-	OUTLIER (0.02%) General / -169.8,-26.8	56% (<i>pttt</i>) chi angles: 59.5,168,180,180	0.087Å	-	-
101	PHE	166.33	HB3 with 97 ALA HA	0.429Å	Favored (9.32%) General /	51.4% (<i>t80</i>) chi angles: 182.3,274.5	0.061Å	-	OUTLIER(S) worst is CA- CB-CG: 5.189

			-51.4,-57.3				σ
102	SER 100.55	0.629Å HB2 with 152 VAL CG1	Favored (5.93%) General / -88.5,61.1	56.5% (<i>p</i>) chi angles: 55.4	0.056Å	-	-
103	ARG 318.05	0.797Å CD with 118 ILE HD13	Favored (2.06%) General / 49.9,26.7	0% chi angles: 55.3,68.2,79.3,208.5	0.061Å	-	OUTLIER(S) worst is C-N- CA: 6.093 σ
104	LEU 248.73	0.908Å HD21 with 135 ILE HD11	OUTLIER (0.01%) General / -68.0,-89.4	3.2% (<i>mm?</i>) chi angles: 268.8,284.8	0.154Å	-	-
105	MET 160.58	0.631Å HE2 with 134 LEU CD1	Favored (35.39%) General / -72.3,-49.3	58.3% (<i>tpp</i>) chi angles: 181.9,70.3,68.8	0.081Å	-	-
106	GLU 83.44	-	Favored (65.37%) General / -70.0,-46.3	32.5% (<i>tp10</i>) chi angles: 191.2,76.2,3.7	0.08Å	-	-
107	GLU 71.44	0.791Å HB2 with 103 ARG CA	Favored (58.02%) General / -75.8,-27.5	51.3% (<i>mt-10</i>) chi angles: 291.8,183.3,57.5	0.11Å	-	-
108	LYS 148.79	0.775Å HB3 with 103 ARG NH1	Favored (72.58%) General / -55.9,-40.2	5.7% (<i>ptmm?</i>) chi angles: 54.1,179.4,269.3,278.8	0.117Å	-	-
109	GLU 75.86	-	Favored (99.02%) General / -62.8,-43.4	89.4% (<i>tt0</i>) chi angles: 180.2,180,357.1	0.081Å	-	-
110	LYS 42.43	0.615Å O with 113 LYS HG2	Favored (88.94%) General / -66.3,-42.0	93.1% (<i>mttt</i>) chi angles: 290,168.9,179.9,180	0.067Å	-	-
111	LEU 98.76	0.704Å HD11 with 103 ARG HB2	Favored (89.36%) General / -62.1,-38.6	6.9% (<i>mp</i>) chi angles: 288.8,76.9	0.158Å	-	-
112	GLN 84.21	0.559Å HG3 with 108 LYS O	Favored (98.05%) General / -60.8,-42.7	76.1% (<i>mt-30</i>) chi angles: 298.4,176.8,43.6	0.079Å	-	-
113	LYS 101.02	0.615Å HG2 with 110	Favored (74.5%) General /	20.7% (<i>ptmt</i>) chi angles:	0.062Å	-	-

			LYS O	-65.4,-32.7	65.4,170.5,298.8,178.4				
114	HIS	100.14	0.741Å CE1 with 66 LEU HB2	Favored (66.46%) General / -67.0,-25.7	51.2% (<i>t</i> -80) chi angles: 180.2,278.1	0.064Å	-	OUTLIER(S) worst is CD2- NE2-CE1: 4.504 σ	
115	GLY	29.32	0.783Å HA2 with 145 TYR CD2	Favored (49.39%) Glycine / 98.3,11.2	-	-	-	-	
116	VAL	54.56	0.648Å HG23 with 114 HIS ND1	Favored (14.76%) Isoleucine or valine / -80.1,141.8	77.1% (<i>t</i>) chi angles: 172.3	0.088Å	-	-	
117	CYS	52.86	0.734Å HA with 142 THR OG1	Favored (32.25%) General / -107.2,116.6	52.5% (<i>t</i>) chi angles: 180.1	0.113Å	-	-	
118	ILE	71.89	0.797Å HD13 with 103 ARG CD	Favored (36.42%) Isoleucine or valine / -129.3,145.0	14.3% (<i>tt</i>) chi angles: 195.1,160.2	0.05Å	-	-	
119	ARG	174.03	0.97Å HD2 with 149 PHE CZ	Favored (38.34%) General / -157.8,163.1	0% chi angles: 63.1,332,108,78.2	0.112Å	-	-	
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			Avg: 83.62	Clashscore: 104.16	Outliers: 2 of 227	Poor rotamers: 8 of 205	Outliers: 2 of 218	Outliers: 0 of 229	Outliers: 31 of 229
120	VAL	99.09	1.048Å HB with 135 ILE HG21	Favored (14.9%) Isoleucine or valine / -82.5,141.5	14.2% (<i>p</i>) chi angles: 63.1	0.071Å	-	-	
121	LEU	115.08	0.827Å HD12 with 156 TYR CD1	Favored (31.05%) General / -129.8,123.5	1.8% (<i>tm</i> ?) chi angles: 193.7,296.8	0.095Å	-	-	
122	GLY	27	0.634Å N with 121 LEU HD13	Favored (42.36%) Glycine / 167.9,-177.1	-	-	-	-	
123	ASP	83.59	-	Favored (36.29%) General /	53.8% (<i>t0</i>) chi angles: 188.1,10.7	0.094Å	-	-	

				-98.3,138.0					
124	LEU	101.71	0.675Å O with 124 LEU HD23	Favored (78.19%) General / -68.4,-43.0	4.1% (<i>tt</i>) chi angles: 190.5,177.2	0.032Å	-	-	
125	HIS	85.86	0.514Å HB3 with 126 LEU CD2	Favored (55.74%) General / -62.5,134.0	96.3% (<i>m-70</i>) chi angles: 297.4,276.3	0.09Å	-		OUTLIER(S) worst is CD2- NE2-CE1: 4.408 σ
126	LEU	245.68	0.549Å N with 126 LEU HD22	Allowed (0.74%) General / 75.7,-14.8	2% (<i>mm?</i>) chi angles: 292.9,293.9	0.08Å	-	-	
127	LEU	76.51	1.12Å HD12 with 128 PRO HD2	Favored (79.54%) Pre-proline / -74.5,143.6	56.7% (<i>tp</i>) chi angles: 180,59.2	0.068Å	-	-	
128	PRO	88.69	1.12Å HD2 with 127 LEU HD12	Favored (45.64%) Trans-proline / -67.9,138.9	9.3% (<i>Cg_endo</i>) chi angles: 10.9	0.135Å	-	-	
129	LEU	85.18	0.692Å O with 129 LEU HD23	Favored (86.51%) General / -63.1,-37.3	3.3% (<i>tt</i>) chi angles: 188.8,180	0.059Å	-	-	
130	ASP	89.82	0.764Å OD2 with 98 ARG HD3	Favored (79.22%) General / -60.6,-36.6	26.1% (<i>p-10</i>) chi angles: 62.3,331.5	0.073Å	-	-	
131	LEU	38.13	0.503Å HD23 with 127 LEU HD11	Favored (86.35%) General / -66.9,-42.0	59.4% (<i>tp</i>) chi angles: 180.2,64.5	0.082Å	-	-	
132	GLN	43.97	0.525Å HE22 with 124 LEU HG	Favored (97.85%) General / -63.8,-41.4	87.6% (<i>mt-30</i>) chi angles: 295.7,179.9,0.5	0.115Å	-	-	
133	GLU	69.96	-	Favored (94.95%) General / -62.8,-44.9	87.2% (<i>tt0</i>) chi angles: 182.5,180,7.4	0.07Å	-	-	
134	LEU	53.28	0.983Å HD12 with 98 ARG HG2	Favored (86.85%) General / -66.5,-38.2	36.2% (<i>mt</i>) chi angles: 293.4,188.1	0.084Å	-	-	
135	ILE	91.94	1.048Å HG21 with	Favored (83.14%)	34.4% (<i>mm</i>)	0.097Å	-	-	

			120 VAL HB	Isoleucine or valine / -63.7,-38.8	chi angles: 295,294.7			
136	ALA	24.6	-	Favored (97.16%) General / -60.9,-42.0	-	0.086Å	-	-
137	GLN	64.47	-	Favored (96.03%) General / -64.6,-41.7	81.8% (<i>mm</i> -40) chi angles: 292,289.2,328.7	0.093Å	-	-
138	ALA	37.75	0.549Å HB2 with 103 ARG HH12	Favored (95.77%) General / -62.4,-40.2	-	0.075Å	-	-
139	VAL	30.64	0.718Å O with 143 LYS HG3	Favored (98.34%) Isoleucine or valine / -63.8,-44.6	97.5% (<i>t</i>) chi angles: 176.2	0.071Å	-	-
#	Alt Res	High B	Clash > 0.4Å	Ramachandran	Rotamer	Cβ deviation	Bond lengths	Bond angles
		Avg: 83.62	Clashscore: 104.16	Outliers: 2 of 227	Poor rotamers: 8 of 205	Outliers: 2 of 218	Outliers: 0 of 229	Outliers: 31 of 229
140	GLN	89.19	0.591Å HA with 143 LYS HE2	Favored (86.72%) General / -65.4,-44.7	1.1% (<i>tm</i> 0?) chi angles: 195.7,288.6,128	0.071Å	-	-
141	ALA	57.16	0.438Å HB1 with 112 GLN OE1	Favored (89.52%) General / -64.1,-38.0	-	0.081Å	-	OUTLIER(S) worst is C-N- CA: 4.198 σ
142	THR	160.99	0.734Å OG1 with 117 CYS HA	Favored (54.63%) General / -82.1,-15.1	6.1% (<i>t</i>) chi angles: 195.6	0.053Å	-	-
143	LYS	97.47	0.718Å HG3 with 139 VAL O	Favored (62.78%) General / -53.9,-36.5	49.1% (<i>mtmt</i>) chi angles: 295.4,170.4,292.9,180	0.083Å	-	-
144	ASN	52.24	-	Favored (82.31%) General / -68.1,-38.4	81.4% (<i>m</i> -20) chi angles: 293.1,309.1	0.082Å	-	OUTLIER(S) worst is CA- CB-CG: 4.278 σ
145	TYR	81.1	0.783Å CD2 with 115 GLY HA2	Favored (19.5%) General / -55.4,146.4	96.7% (<i>m</i> -85) chi angles: 295.9,88.2	0.089Å	-	OUTLIER(S) worst is CA- CB-CG: 4.281 σ

146	ASN	185.82	0.49Å H with 147 LYS NZ	Allowed (1.96%) General / -143.0,20.9	32% (<i>t</i> -20) chi angles: 198.8,358.4	0.072Å	-	OUTLIER(S) worst is CA- CB-CG: 4.917 σ
147	LYS	116.48	0.727Å HE3 with 147 LYS N	Favored (68.45%) General / -68.4,-47.0	0.9% chi angles: 289.9,67.1,271.2,177.7	0.086Å	-	-
148	CYS	78.16	-	Favored (31.16%) General / -153.6,152.7	52.1% (<i>t</i>) chi angles: 180	0.113Å	-	-
149	PHE	56.99	0.97Å CZ with 119 ARG HD2	Favored (55.98%) General / -113.5,127.1	41.6% (<i>t</i> 80) chi angles: 193.5,82.7	0.127Å	-	-
150	LEU	54.8	0.7Å HD11 with 62 LEU HG	Favored (45.38%) General / -117.4,123.0	88.7% (<i>mt</i>) chi angles: 298.5,180.1	0.088Å	-	-
151	ASN	60.46	0.417Å OD1 with 149 PHE HE2	Favored (22.11%) General / -93.8,110.5	79.6% (<i>m</i> -20) chi angles: 301.5,317.5	0.142Å	-	OUTLIER(S) worst is CA- CB-CG: 4.161 σ
152	VAL	53.8	0.787Å CG2 with 118 ILE HD11	Favored (18.38%) Isoleucine or valine / -93.8,106.7	76.6% (<i>t</i>) chi angles: 174.2	0.096Å	-	-
153	CYS	52.84	-	Favored (23.99%) General / -96.9,112.1	80.1% (<i>m</i>) chi angles: 300.9	0.072Å	-	-
154	PHE	142.34	0.799Å CE2 with 78 PHE HB2	Favored (50.65%) General / -126.7,134.5	15.1% (<i>t</i> 80) chi angles: 201.6,271	0.088Å	-	OUTLIER(S) worst is CA- CB-CG: 6.105 σ
155	ALA	51.37	0.518Å O with 80 ILE HD11	Favored (20.83%) General / 58.8,41.8	-	0.07Å	-	-
156	TYR	75.18	0.827Å CD1 with 121 LEU HD12	Favored (45.74%) General / -99.8,131.6	71.1% (<i>t</i> 80) chi angles: 175.4,87.1	0.122Å	-	-
157	THR	67.95	0.718Å CB with 80 ILE HD13	Favored (28.86%) General /	37.5% (<i>m</i>) chi angles: 292.9	0.148Å	-	-

158	SER	84.63	0.784Å HB2 with 217 LEU HB3	-140.4,133.1 Favored (89.93%) General / -63.5,-45.6	94.9% (<i>p</i>) chi angles: 64.9	0.075Å	-	-
159	ARG	80.19	0.423Å HG2 with 157 THR CG2	Favored (94.36%) General / -65.1,-40.4	29.1% (<i>ptt-85</i>) chi angles: 60.8,174.4,184.2,264.6	0.067Å	-	-
#	Alt Res	High B	Clash > 0.4Å	Ramachandran	Rotamer	Cβ deviation	Bond lengths	Bond angles
		Avg: 83.62	Clashscore: 104.16	Outliers: 2 of 227	Poor rotamers: 8 of 205	Outliers: 2 of 218	Outliers: 0 of 229	Outliers: 31 of 229
160	HIS	69.86	-	Favored (92.77%) General / -65.5,-40.1	64.8% (<i>m80</i>) chi angles: 284.6,85.8	0.084Å	-	OUTLIER(S) worst is CD2- NE2-CE1: 4.497 σ
161	GLU	40.66	0.438Å HG2 with 192 LEU HD22	Favored (98.81%) General / -63.4,-43.0	45% (<i>tt0</i>) chi angles: 180.2,179.7,53.4	0.074Å	-	-
162	ILE	114.64	0.421Å HG12 with 159 ARG O	Favored (80.46%) Isoleucine or valine / -65.7,-37.6	40.7% (<i>pt</i>) chi angles: 60,175.5	0.111Å	-	-
163	SER	25.6	0.789Å O with 166 VAL HG12	Favored (87.2%) General / -64.4,-37.4	66.7% (<i>m</i>) chi angles: 296.4	0.086Å	-	-
164	ASN	25.29	0.54Å HA with 167 ARG NE	Favored (96.88%) General / -64.2,-40.8	93.9% (<i>m-20</i>) chi angles: 292.7,344.4	0.085Å	-	OUTLIER(S) worst is CA- CB-CG: 4.641 σ
165	ALA	27.2	0.486Å HB2 with 192 LEU HD23	Favored (99.26%) General / -63.2,-41.9	-	0.082Å	-	-
166	VAL	77.38	0.789Å HG12 with 163 SER O	Favored (81.4%) Isoleucine or valine / -64.4,-38.0	9.8% (<i>p</i>) chi angles: 72.1	0.077Å	-	-
167	ARG	101.62	0.54Å NE with 164 ASN HA	Favored (95.29%) General / -61.7,-40.5	32.2% (<i>mtm105</i>) chi angles: 294.7,180,297.6,87.5	0.092Å	-	-
168	GLU	31.59	0.549Å OE1 with 191	Favored (96.82%)	43.4% (<i>tt0</i>) chi angles:	0.102Å	-	-

			CYS HA	General / -62.1,-44.4	183.4,179.9,62.8				
169	MET	27.87	0.535Å O with 173 VAL HG23	Favored (95.11%) General / -62.7,-39.8	98% (<i>mtp</i>) chi angles: 294.5,179.9,68.3	0.107Å	-	-	
170	ALA	22.97	-	Favored (95.69%) General / -61.9,-40.5	-	0.086Å	-	-	
171	TRP	32	0.403Å CD1 with 167 ARG O	Favored (95.57%) General / -62.1,-44.9	98.7% (<i>m95</i>) chi angles: 291.7,97.3	0.118Å	-	-	
172	GLY	15.12	-	Favored (99.28%) Glycine / -62.2,-40.7	-	-	-	-	
173	VAL	25.04	0.535Å HG23 with 169 MET O	Favored (94.81%) Isoleucine or valine / -61.7,-42.6	74.5% (<i>t</i>) chi angles: 174	0.076Å	-	-	
174	GLU	58.08	-	Favored (91.56%) General / -62.6,-38.8	87.9% (<i>tt0</i>) chi angles: 185.1,176,358.5	0.071Å	-	-	
175	GLN	74.23	0.44Å OE1 with 177 LEU HD11	Favored (77.99%) General / -64.9,-34.3	77.5% (<i>mt-30</i>) chi angles: 296.4,180,51.5	0.081Å	-	-	
176	GLY	18.73	-	Favored (50.12%) Glycine / 102.4,-0.6	-	-	-	-	
177	LEU	46.99	0.44Å HD11 with 175 GLN OE1	Favored (82.44%) General / -65.8,-36.1	82.4% (<i>mt</i>) chi angles: 295.7,180.9	0.087Å	-	-	
178	LEU	56.54	-	Favored (43.06%) General / -116.3,145.6	59.2% (<i>tp</i>) chi angles: 179.8,60	0.059Å	-	-	
179	ASP	44.92	-	Favored (32.59%) Pre-proline / -100.3,127.6	64.5% (<i>t0</i>) chi angles: 186,348.5	0.094Å	-	-	
#	Alt	Res	High B	Clash > 0.4Å	Ramachandran	Rotamer	Cβ deviation	Bond lengths	Bond angles

		Avg: 83.62	Clashscore: 104.16	Outliers: 2 of 227	Poor rotamers: 8 of 205	Outliers: 2 of 218	Outliers: 0 of 229	Outliers: 31 of 229
180	PRO	65.4	-	Favored (70.33%) Trans-proline / -63.5,-27.6	87.1% (Cg_exo) chi angles: 331.3	0.158Å	-	-
181	SER	67.26	-	Favored (67.06%) General / -69.4,-29.4	47.5% (t) chi angles: 180	0.086Å	-	-
182	ASP	51.31	-	Favored (67.4%) General / -68.9,-29.2	82.9% (m-20) chi angles: 292.1,353.1	0.081Å	-	OUTLIER(S) worst is CA- CB-CG: 4.489 σ
183	ILE	75.08	1.071Å HA with 187 LEU HD12	Favored (2.78%) Isoleucine or valine / -46.0,132.8	89.9% (mt) chi angles: 298.6,168.2	0.059Å	-	-
184	SER	79.03	-	Favored (13.58%) General / -137.0,172.1	43.2% (t) chi angles: 178.1	0.122Å	-	-
185	GLU	72.66	-	Favored (79.25%) General / -62.6,-35.4	25.8% (pt-20) chi angles: 61.2,180,4.3	0.071Å	-	-
186	SER	70.87	-	Favored (72.5%) General / -63.9,-31.4	97.5% (p) chi angles: 64	0.075Å	-	-
187	LEU	140.02	1.071Å HD12 with 183 ILE HA	Favored (93.52%) General / -63.7,-39.1	27.7% (mt) chi angles: 292,189.8	0.07Å	-	-
188	LEU	72.96	0.829Å HD22 with 218 TRP CH2	Favored (95.53%) General / -64.7,-42.1	5.5% (mp) chi angles: 293.8,64.6	0.079Å	-	-
189	ASP	35.81	-	Favored (94.02%) General / -60.4,-41.5	60% (m-20) chi angles: 287.6,313.1	0.113Å	-	-
190	LYS	68.33	-	Favored (78.96%) General / -64.2,-34.8	61.7% (mttm) chi angles: 297.9,166.2,185.8,295.5	0.074Å	-	-

191	CYS	75.51	0.549Å HA with 168 GLU OE1	Favored (60.92%) General / -73.0,-20.8	99.5% (<i>m</i>) chi angles: 294.5	0.08Å	-	-
192	LEU	55.65	0.486Å HD23 with 165 ALA HB2	Favored (14.84%) General / -62.5,159.8	86.8% (<i>mt</i>) chi angles: 295.9,180	0.066Å	-	-
193	TYR	45.28	-	Favored (83.16%) General / -58.0,-47.4	88% (<i>t80</i>) chi angles: 178.7,82.5	0.088Å	-	-
194	THR	135.08	0.652Å HG22 with 121 LEU CD2	Favored (5.54%) General / -117.5,96.7	9.4% (<i>t</i>) chi angles: 180.3	0.051Å	-	-
195	ASN	90.8	-	Favored (71.36%) General / -65.3,-30.5	18.5% (<i>p-10</i>) chi angles: 64.5,307.7	0.086Å	-	OUTLIER(S) worst is CA- CB-CG: 4.615 σ
196	ARG	115.13	0.411Å O with 198 PRO HD3	Favored (71.95%) General / -64.1,-30.9	89% (<i>mtm180</i>) chi angles: 295.1,168.6,293.2,168.2	0.077Å	-	-
197	SER	90.23	0.44Å HA with 198 PRO HD2	Favored (71.63%) Pre-proline / -56.0,128.5	47.4% (<i>t</i>) chi angles: 180.1	0.081Å	-	-
198	PRO	72.55	0.44Å HD2 with 197 SER HA	Favored (17.27%) Trans-proline / -82.0,165.7	15.9% (<i>Cg_endo</i>) chi angles: 16.7	0.096Å	-	-
199	HIS	89.01	-	Favored (96.43%) Pre-proline / -67.5,142.5	80% (<i>m80</i>) chi angles: 291.8,85.4	0.082Å	-	OUTLIER(S) worst is CD2- NE2-CE1: 4.413 σ
<div> <div>#</div> <div>Alt Res</div> <div>High B</div> <div>Clash > 0.4Å</div> <div>Ramachandran</div> <div>Rotamer</div> <div>Cβ deviation</div> <div>Bond lengths</div> <div>Bond angles</div> </div>								
Avg: 83.62			Clashscore: 104.16	Outliers: 2 of 227	Poor rotamers: 8 of 205	Outliers: 2 of 218	Outliers: 0 of 229	Outliers: 31 of 229
200	PRO	122.94	0.793Å CD with 219 GLN HB3	Favored (10.4%) Trans-proline / -71.5,124.5	81.3% (<i>Cg_exo</i>) chi angles: 330.4	0.2Å	-	-
201	ASP	63.69	-	Favored (19.14%) General /	68.9% (<i>m-20</i>) chi angles: 292,311.2	0.067Å	-	OUTLIER(S) worst is CA- CB-CG: 5.514

				-78.8,-46.2				σ
202	ILE	93.63	0.882Å CG2 with 29 ILE HG12	Favored (63.67%) Isoleucine or valine / -131.0,130.1	21.1% (<i>tt</i>) chi angles: 188.7,168.3	0.059Å	-	-
203	LEU	99.6	0.754Å CD1 with 212 LEU HD22	Favored (56.4%) General / -113.2,128.2	3.3% (<i>tt</i>) chi angles: 189.2,180.1	0.056Å	-	-
204	ILE	106.77	0.685Å N with 203 LEU HD23	Favored (38.5%) Isoleucine or valine / -117.6,141.0	33.6% (<i>mm</i>) chi angles: 309.5,298.6	0.188Å	-	-
205	ARG	93.68	0.696Å HG3 with 32 ILE HG23	Favored (30.33%) General / -140.2,133.8	26.2% (<i>ttt180</i>) chi angles: 180.1,164.3,182.3,140.1	0.123Å	-	-
206	THR	47.29	0.516Å HB with 230 VAL O	Favored (5.02%) General / -99.8,177.8	73.6% (<i>p</i>) chi angles: 61.8	0.065Å	-	-
207	SER	80.64	0.666Å HB2 with 38 ARG NH1	Favored (3.4%) General / 76.4,1.8	27.7% (<i>p</i>) chi angles: 77.8	0.128Å	-	-
208	GLY	29.06	-	Favored (78.97%) Glycine / 92.0,-5.4	-	-	-	-
209	GLU	85.58	0.492Å HB2 with 205 ARG HE	Favored (37.19%) General / -79.1,141.7	75% (<i>mm-40</i>) chi angles: 300.8,294.9,345	0.086Å	-	-
210	VAL	112.57	0.914Å HG23 with 227 PHE CZ	Favored (2.19%) Isoleucine or valine / -112.4,92.8	3.2% (<i>p</i>) chi angles: 51.4	0.037Å	-	-
211	ARG	177.53	-	Favored (9.42%) General / -162.3,141.7	62.4% (<i>mtt180</i>) chi angles: 307.8,171.1,185,141.1	0.235Å	-	-
212	LEU	66.55	0.754Å HD22 with 203 LEU CD1	Favored (61.77%) General / -75.1,-36.0	82.3% (<i>mt</i>) chi angles: 299.6,181.6	0.098Å	-	-
				Favored (22.69%)	75.8% (<i>m</i>)			

213	SER	66.43	-	General / 54.8,47.3	chi angles: 294.5	0.081Å	-	-
214	ASP	106.44	0.426Å O with 158 SER HB3	Favored (19.26%) General / 59.6,29.7	48.3% (<i>m</i> -20) chi angles: 284,317.4	0.106Å	-	-
215	PHE	80.77	0.745Å HE2 with 30 ALA HB1	Favored (48.72%) General / -102.0,130.8	61.8% (<i>t</i> 80) chi angles: 180.1,64.5	0.045Å	-	OUTLIER(S) worst is CA- CB-CG: 4.439 σ
216	LEU	71.62	1.111Å HD11 with 74 THR HG21	Favored (21.92%) General / 49.2,47.6	75.8% (<i>mt</i>) chi angles: 295,182	0.082Å	-	-
217	LEU	74.59	0.784Å HB3 with 158 SER HB2	Favored (78.09%) General / -62.2,-48.9	60.5% (<i>tp</i>) chi angles: 180,60.7	0.144Å	-	-
218	TRP	68.25	0.829Å CH2 with 188 LEU HD22	Favored (81.67%) General / -66.0,-45.5	47.5% (<i>t</i> -105) chi angles: 187.7,272.8	0.056Å	-	-
219	GLN	101.42	0.793Å HB3 with 200 PRO CD	Favored (81.36%) General / -67.2,-36.2	28.7% (<i>mp</i> 0) chi angles: 295.6,82.6,358.1	0.034Å	-	-

#	Alt Res	High B	Clash > 0.4Å	Ramachandran	Rotamer	Cβ deviation	Bond lengths	Bond angles
		Avg: 83.62	Clashscore: 104.16	Outliers: 2 of 227	Poor rotamers: 8 of 205	Outliers: 2 of 218	Outliers: 0 of 229	Outliers: 31 of 229
220	THR	167.95	0.738Å HG22 with 216 LEU O	Allowed (1.73%) General / -97.4,34.8	15.2% (<i>t</i>) chi angles: 187.9	0.077Å	-	-
221	SER	89.87	-	Favored (88.32%) General / -58.3,-43.8	47.5% (<i>t</i>) chi angles: 180	0.084Å	-	-
222	HIS	138.29	-	Favored (2.83%) General / -120.7,43.2	80.3% (<i>m</i> 80) chi angles: 294,83.8	0.092Å	-	OUTLIER(S) worst is CD2- NE2-CE1: 4.422 σ
223	SER	80.7	0.474Å HB3 with 220 THR OG1	Favored (19.95%) General / -85.3,160.0	47.3% (<i>t</i>) chi angles: 180.1	0.1Å	-	-

224	CYS	70.71	-	Favored (27.55%) General / -94.4,142.3	52.8% (<i>t</i>) chi angles: 180.2	0.107Å	-	-
225	LEU	83.49	0.715Å HD21 with 212 LEU HD21	Favored (50.24%) General / -112.1,137.4	3.4% (<i>mp</i>) chi angles: 299.2,64	0.107Å	-	-
226	VAL	99.51	0.596Å HG12 with 204 ILE HD13	Favored (66.09%) Isoleucine or valine / -129.9,129.7	14.1% (<i>p</i>) chi angles: 64.7	0.07Å	-	-
227	PHE	47.73	0.914Å CZ with 210 VAL HG23	Favored (55.09%) General / -111.1,131.4	84.6% (<i>m-85</i>) chi angles: 298.7,286.2	0.097Å	-	OUTLIER(S) worst is CA- CB-CG: 6.002 σ
228	GLN	50.74	-	Favored (23%) Pre-proline / -116.6,133.6	78% (<i>mt-30</i>) chi angles: 295.4,169.5,296.8	0.06Å	-	-
229	PRO	124.45	-	Favored (32.12%) Trans-proline / -70.6,-19.3	13.8% (<i>Cg_endo</i>) chi angles: 16	0.187Å	-	-
230	VAL	90.33	0.672Å CG2 with 234 GLU HB2	Favored (13.78%) Isoleucine or valine / -67.9,145.2	14.9% (<i>p</i>) chi angles: 63.7	0.063Å	-	-
231	LEU	99.8	0.686Å HB2 with 234 GLU OE1	Favored (30.27%) General / -78.4,154.9	5.3% (<i>mp</i>) chi angles: 296.2,71.7	0.099Å	-	-
232	TRP	71.59	0.71Å HB3 with 233 PRO HD3	Favored (95.2%) Pre-proline / -58.9,-42.6	76.9% (<i>t-105</i>) chi angles: 180.4,253.5	0.074Å	-	-
233	PRO	76.42	0.835Å HB3 with 53 GLN HG2	Favored (69.52%) Trans-proline / -63.6,-24.5	84.3% (<i>Cg_exo</i>) chi angles: 331	0.15Å	-	-
234	GLU	91.99	0.686Å OE1 with 231 LEU HB2	Favored (51.39%) General / -77.5,-26.7	54.3% (<i>mp0</i>) chi angles: 288.5,83.8,352.7	0.095Å	-	-
235	TYR	92.64	0.455Å HD2 with 57 LYS HD2	Favored (55.77%) General /	16.7% (<i>t80</i>) chi angles: 201.7,82.3	0.052Å	-	-

236	THR	48.68	-	-60.7,142.5 Favored (8.86%) General / -140.0,176.3	67.8% (<i>p</i>) chi angles: 63.1	0.084Å	-	-
237	PHE	154.88	1.038Å HB2 with 64 TRP HB2	Favored (72.74%) General / -60.6,-33.4	39.4% (<i>p90</i>) chi angles: 59.3,82.1	0.043Å	-	-
238	TRP	56.58	0.76Å HA with 241 PHE CD2	Favored (88.7%) General / -59.0,-41.9	87.1% (<i>m95</i>) chi angles: 295.1,86	0.192Å	-	-
239	ASN	33.61	-	Favored (92.45%) General / -64.2,-38.8	95.8% (<i>m-20</i>) chi angles: 290.9,332.8	0.068Å	-	OUTLIER(S) worst is CA- CB-CG: 4.86 σ
#	Alt Res	High B	Clash > 0.4Å	Ramachandran	Rotamer	Cβ deviation	Bond lengths	Bond angles
		Avg: 83.62	Clashscore: 104.16	Outliers: 2 of 227	Poor rotamers: 8 of 205	Outliers: 2 of 218	Outliers: 0 of 229	Outliers: 31 of 229
240	LEU	49.42	0.802Å HD23 with 64 TRP HD1	Favored (99.05%) General / -62.5,-43.3	45.3% (<i>tp</i>) chi angles: 171.1,60.7	0.116Å	-	-
241	PHE	166.79	0.825Å HB3 with 64 TRP CE2	Favored (90.31%) General / -65.5,-38.7	7.8% (<i>p90</i>) chi angles: 73.6,74.5	0.17Å	-	-
242	GLU	67.85	0.416Å N with 241 PHE CD1	Favored (90.12%) General / -58.6,-44.7	83.8% (<i>tt0</i>) chi angles: 180.1,179.9,346.7	0.07Å	-	-
243	ALA	28.23	-	Favored (96.19%) General / -64.4,-42.4	-	0.076Å	-	-
244	ILE	40.47	0.814Å HD13 with 70 ILE HD11	Favored (84.23%) Isoleucine or valine / -64.8,-39.0	95.6% (<i>mt</i>) chi angles: 296.4,168.4	0.051Å	-	-
245	LEU	39.09	0.617Å O with 245 LEU HD23	Favored (99.94%) General / -62.8,-43.0	3.3% (<i>tt</i>) chi angles: 189.1,179.9	0.07Å	-	-
				Favored	65.8% (<i>tt0</i>)			

246	GLN	60.03	-	(99.53%) General / -62.5,-42.5	chi angles: 180.2,179.9,356.4	0.09Å	-	-
247	PHE	36.59	0.536Å HB2 with 202 ILE HD12	Favored (79.4%) General / -63.1,-48.3	81.9% (<i>t80</i>) chi angles: 177.4,85	0.091Å	-	OUTLIER(S) worst is CA- CB-CG: 5.21 σ
248	GLN	108.76	0.653Å HG2 with 20 ILE HD13	-	0% chi angles: 63,95.8,180.6	0.053Å	-	-

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