

Table S1. Details of the PDB file, contact distance, atoms involved and geometric parameters describing Trp-aromatic interaction

PDB Theta CODE (T-P)	Resolution Phi	R D	Dxy factor	TRP	Partner		Shortest Contact		
				Dz	Res	Res.No.	Trp Atom	Partner Atom	Distance (Angstrom)
				Res.No.	Res	Res.No.			
1531	1.60	0.15		84	PHE	91	CZ2	CD1	3.68
22.88	39.06	5.22	2.03	4.81					
1531	1.60	0.15		107	TYR	48	CH2	OH	3.71
42.97	88.17	7.39	5.04	-5.41					
1531	1.60	0.15		134	PHE	131	CE2	CG	3.57
34.74	29.76	4.26	2.43	-3.50					
1531	1.60	0.15		134	TYR	157	CZ3	CD1	3.99
19.74	99.72	5.67	1.91	5.34					
1931	1.33	0.18		108	TRP	28	CH2	CZ3	3.85
39.71	96.12	7.37	4.71	5.67					
1931	1.33	0.18		28	TRP	108	CZ3	CH2	3.85
69.19	114.63	7.37	6.89	2.62					
1931	1.33	0.18		111	TYR	23	CZ2	CE1	3.80
36.88	35.57	5.95	3.57	-4.76					
1931	1.33	0.18		123	PHE	34	CZ2	CZ	3.73
56.32	31.11	6.00	4.99	-3.33					
lade	2.00	0.20		309 A	TYR	372 A	CD1	CE1	3.38
64.05	-87.82	5.72	5.14	-2.51					
lade	2.00	0.20		365 A	TYR	347 A	CZ3	OH	3.91
42.30	102.10	6.45	4.34	4.77					
lamp	1.80	0.16		13	PHE	137	CZ3	CD1	3.63
84.09	105.50	7.07	7.03	-0.73					
lamp	1.80	0.16		48	PHE	32	CG	CZ	3.95
33.16	-136.72	5.91	3.23	-4.95					
lamp	1.80	0.16		233	TYR	148	NE1	OH	2.91
79.10	-21.25	6.30	6.19	1.19					
lamp	1.80	0.16		233	TYR	238	CE3	CD2	3.59
85.81	149.10	6.48	6.46	-0.47					
laoz	1.90	0.20		8 A	TYR	6 A	NE1	CD2	3.97
50.49	-1.98	5.80	4.47	-3.69					
laoz	1.90	0.20		8 A	TRP	61 A	CZ3	CH2	3.95
64.53	103.81	8.41	7.59	-3.62					
laoz	1.90	0.20		61 A	TRP	8 A	CH2	CZ3	3.95
87.09	84.84	8.41	8.40	-0.43					
laoz	1.90	0.20		8 A	TYR	103 A	CH2	CZ	3.51
68.40	81.43	6.07	5.64	2.24					
laoz	1.90	0.20		15 A	PHE	31 A	NE1	CD2	3.56
45.92	-66.43	5.87	4.22	4.08					
laoz	1.90	0.20		61 A	TYR	91 A	NE1	CD1	3.70
84.32	-2.66	5.17	5.14	0.51					
laoz	1.90	0.20		61 A	PHE	93 A	CE2	CE2	3.67
20.55	-109.74	4.69	1.64	-4.39					
laoz	1.90	0.20		144 A	HIS	146 A	NE1	CD2	3.72
70.92	-55.38	6.36	6.01	-2.08					
laoz	1.90	0.20		277 A	TYR	186 A	NE1	OH	3.09
73.01	-16.62	5.85	5.59	-1.71					
laoz	1.90	0.20		362 A	HIS	445 A	CH2	CE1	3.48
44.58	96.33	5.54	3.89	-3.94					

laoz	1.90	0.20	362	A	HIS	512	A	NE1	CD2	3.71
30.36	-10.53	4.37	2.20	-3.77						
laoz	1.90	0.20	447	A	PHE	505	A	CE3	CZ	3.73
70.34	157.37	5.15	4.85	-1.73						
laoz	1.90	0.20	490	A	PHE	464	A	CD2	CD1	3.81
13.31	113.05	5.08	1.15	4.95						
laoz	1.90	0.20	503	A	HIS	452	A	CD1	CE1	3.47
32.66	-81.30	5.29	2.85	-4.45						
larb	1.20	0.15	73		PHE	52		CH2	CE1	3.76
58.40	68.11	6.82	5.81	3.57						
larb	1.20	0.15	73		PHE	127		CD1	CD2	3.31
65.99	-85.35	5.99	5.47	-2.44						
larb	1.20	0.15	134		TRP	234		CH2	CH2	3.60
38.42	92.61	6.62	4.11	5.19						
larb	1.20	0.15	234		TRP	134		CH2	CH2	3.60
68.07	97.20	6.62	6.14	2.47						
larb	1.20	0.15	169		HIS	210		CE2	NE2	3.45
8.52	119.79	3.53	0.52	3.49						
larb	1.20	0.15	182		TYR	227		CZ2	OH	3.51
71.44	36.48	7.17	6.80	2.28						
larb	1.20	0.15	247		TYR	51		CE3	OH	3.80
63.46	-175.98	6.46	5.78	-2.89						
lasu	1.70	0.15	105		HIS	104		CD1	CD2	3.70
48.09	-82.70	6.04	4.49	-4.04						
lasu	1.70	0.15	105		PHE	126		CH2	CD1	3.67
60.65	41.34	6.45	5.62	-3.16						
lasu	1.70	0.15	105		TRP	134		CE3	CZ3	3.97
82.93	165.88	6.91	6.86	0.85						
lasu	1.70	0.15	134		TRP	105		CZ3	CE3	3.97
53.46	115.01	6.91	5.55	-4.11						
lasu	1.70	0.15	134		TRP	138		CZ2	NE1	3.40
35.85	58.10	5.53	3.24	4.48						
lasu	1.70	0.15	138		TRP	134		NE1	CZ2	3.40
71.69	-64.59	5.53	5.25	-1.74						
lasu#	1.70	0.15	138		TYR	194		CZ3	CE2	3.78
83.20	95.74	6.53	6.48	-0.77						
latl	1.80	0.16	59	A	TRP	81	A	CZ2	CH2	3.80
74.07	14.85	5.62	5.40	1.54						
latl	1.80	0.16	81	A	TRP	59	A	CH2	CZ2	3.80
30.51	45.46	5.62	2.85	-4.84						
latl	1.80	0.16	81	A	PHE	78	A	CZ3	CE1	3.76
70.83	145.35	6.41	6.05	-2.10						
lbbp	2.00	0.20	20	D	PHE	18	D	CE3	CE1	3.78
65.53	156.63	6.57	5.98	-2.72						
lbbp	2.00	0.20	20	D	TYR	23	D	CH2	CD1	3.65
36.98	96.00	5.62	3.38	-4.49						
lbbp	2.00	0.20	27	D	TYR	48	D	CD1	CD2	3.42
58.48	-75.66	6.01	5.12	-3.14						
lbbp	2.00	0.20	28	D	PHE	170	D	CE3	CE1	3.47
46.08	170.87	5.42	3.90	-3.76						
lbbp	2.00	0.20	45	D	TYR	60	D	NE1	CD2	3.32
46.50	6.07	4.66	3.38	3.21						
lbbp	2.00	0.20	133	D	PHE	103	D	CE3	CE1	3.92
58.19	150.28	6.68	5.68	3.52						
lbbp	2.00	0.20	133	D	PHE	131	D	CH2	CD2	3.54
60.77	59.17	6.44	5.62	-3.14						
lbam	1.95	0.19	38		TRP	206		CE3	CZ2	3.44
83.37	138.46	5.99	5.95	0.69						
lbam	1.95	0.19	206		TRP	38		CZ2	CE3	3.44
40.01	20.19	5.99	3.85	-4.59						

1bdm	1.80	0.17		183 B	TYR	277 B	CH2	CE2	3.63
80.12	76.80	6.16	6.07	-1.06					
1bdm	1.80	0.17		183 B	PHE	191 B	CD1	CE2	3.85
61.56	-83.33	6.34	5.58	-3.02					
1bdm	1.80	0.17		183 B	PHE	287 B	CZ3	CD1	3.72
74.02	131.14	7.04	6.77	1.94					
1bdm	1.80	0.17		256 B	TYR	296 B	NE1	CD2	3.60
36.48	-23.55	5.54	3.29	-4.45					
1bec	1.70	0.20		34	PHE	75	CD1	CD2	3.40
73.95	-76.07	5.80	5.57	1.60					
1bec	1.70	0.20		203	PHE	210	CE3	CE2	3.43
81.93	143.75	5.91	5.85	0.83					
1bri	1.90	0.16		71 A	TYR	97 A	CH2	CD1	3.69
65.19	70.31	6.81	6.18	-2.86					
1bri	1.90	0.16		94 A	HIS	18 A	CE2	CD2	3.59
26.84	-14.77	4.71	2.13	-4.20					
1byb	1.90	0.15		55	TYR	72	CD1	OH	3.36
64.76	-93.70	7.95	7.19	3.39					
1byb	1.90	0.15		55	HIS	93	CZ3	NE2	3.48
69.55	103.23	6.06	5.68	2.12					
1byb	1.90	0.15		56	PHE	92	CG	CD1	3.78
29.01	-159.94	5.59	2.71	-4.89					
1byb	1.90	0.15		56	TYR	157	CD1	CE2	3.65
61.39	-100.76	6.11	5.36	-2.92					
1byb	1.90	0.15		69	TYR	157	CH2	CZ	3.99
74.22	62.04	6.42	6.18	1.75					
1byb	1.90	0.15		69	PHE	161	CZ3	CD1	3.63
42.36	98.03	6.29	4.24	-4.65					
1byb	1.90	0.15		110	HIS	146	CE3	CD2	3.80
62.50	147.46	6.46	5.73	2.98					
1byb	1.90	0.15		198	TRP	301	CZ2	CZ2	3.60
80.47	25.48	6.10	6.02	-1.01					
1byb	1.90	0.15		301	TRP	198	CZ2	CZ2	3.60
57.25	-5.37	6.10	5.13	-3.30					
1byb	1.90	0.15		229	PHE	261	CZ3	CG	3.77
85.53	124.27	6.57	6.55	0.51					
1byb	1.90	0.15		265	PHE	217	CD2	CD1	3.80
16.99	137.81	5.25	1.52	-5.02					
1byb	1.90	0.15		265	PHE	262	CZ3	CE1	3.75
77.46	144.27	6.45	6.30	-1.40					
1byb	1.90	0.15		301	HIS	300	CE3	NE2	3.53
86.79	153.22	5.38	5.37	-0.30					
1byb	1.90	0.15		301	TRP	302	CD1	CD1	3.74
83.33	-101.15	7.33	7.28	0.85					
1byb	1.90	0.15		302	TRP	301	CD1	CD1	3.74
57.73	-73.56	7.33	6.20	3.92					
1byb	1.90	0.15		371	TRP	485	CZ3	CH2	3.90
85.04	103.27	8.13	8.10	-0.70					
1byb	1.90	0.15		485	TRP	371	CH2	CZ3	3.90
56.84	71.21	8.13	6.80	-4.45					
lccr	1.50	0.19		67	TYR	75	CH2	CG	3.59
62.38	70.11	5.64	5.00	-2.62					
lcel	1.81	0.18		16 B	PHE	120 B	CZ3	CE2	3.85
35.08	78.96	5.87	3.37	4.81					
lcel	1.81	0.18		38 B	TYR	82 B	CD1	CE1	3.92
71.92	-107.15	6.80	6.46	2.11					
lcel	1.81	0.18		40 B	HIS	42 B	CD2	CD2	3.47
19.39	76.21	4.35	1.45	4.10					
lcel	1.81	0.18		56 B	TYR	51 B	CD2	CD1	3.81
7.30	-39.00	5.03	0.62	4.99					

lcel	1.81	0.18		56 B	TRP	192 B	NE1	CD2	3.45
77.39	-46.31	5.19	5.06	1.13					
lcel	1.81	0.18		192 B	TYR	51 B	CZ2	OH	3.57
47.48	13.15	5.88	4.34	3.97					
lcel	1.81	0.18		192 B	TRP	56 B	CD2	NE1	3.45
9.12	143.01	5.19	0.82	5.12					
lcel	1.81	0.18		216 B	PHE	129 B	CG	CZ	3.65
24.76	-134.58	5.45	2.28	4.95					
lcel	1.81	0.18		216 B	TYR	274 B	CZ2	OH	3.37
65.16	58.26	6.79	6.16	-2.85					
lcel	1.81	0.18		263 B	TYR	321 B	CH2	OH	3.46
72.67	87.96	7.20	6.87	-2.14					
lcel	1.81	0.18		263 B	TYR	330 B	CZ3	CD1	3.52
54.63	139.83	4.86	3.96	-2.81					
lcel	1.81	0.18		367 B	TRP	376 B	CH2	CZ2	3.97
64.62	58.75	7.46	6.74	-3.20					
lcel	1.81	0.18		376 B	TRP	367 B	CZ2	CH2	3.97
85.90	15.16	7.46	7.44	0.53					
lcfb	2.00	0.20		634	TYR	647	NE1	CZ	3.38
64.35	-24.94	4.62	4.16	-2.00					
lcfb	2.00	0.20		681	PHE	655	CE3	CZ	3.65
38.31	120.11	5.81	3.60	4.56					
lcfb	2.00	0.20		734	TYR	749	NE1	CE2	3.44
61.71	-29.33	4.82	4.24	-2.29					
lcfb	2.00	0.20		770	HIS	742	CG	CE1	3.30
31.11	-152.47	4.92	2.55	4.21					
lchm	1.90	0.18		219 B	PHE	62 A	CZ2	CE2	3.72
85.57	32.19	6.69	6.67	-0.52					
lchm	1.90	0.18		219 B	TRP	221 B	NE1	CZ2	3.17
84.49	-46.27	4.81	4.79	-0.46					
lchm	1.90	0.18		221 B	TRP	219 B	CZ2	NE1	3.17
15.81	58.48	4.81	1.31	-4.63					
lchm	1.90	0.18		219 B	PHE	251 B	NE1	CD2	3.40
28.99	-0.45	4.27	2.06	3.74					
lchm	1.90	0.18		219 B	TYR	258 B	CH2	CE1	3.91
60.06	65.07	7.16	6.20	3.57					
lchm	1.90	0.18		219 B	HIS	331 B	CH2	CD2	3.87
23.89	61.71	5.34	2.17	-4.88					
lchm	1.90	0.18		221 B	PHE	62 A	CH2	CE2	3.63
76.52	83.39	7.39	7.19	-1.72					
lchm	1.90	0.18		221 B	HIS	232 B	CZ3	CD2	3.37
66.24	90.09	6.27	5.74	2.53					
lchm	1.90	0.18		221 B	PHE	251 B	CE3	CE2	3.43
47.21	143.79	5.57	4.09	-3.79					
lchm	1.90	0.18		278 B	PHE	266 B	CZ3	CD1	3.65
43.93	133.24	5.95	4.13	-4.29					
lclc	1.90	0.20		220	PHE	244	CE3	CD2	3.81
68.39	165.96	6.27	5.83	2.31					
lclc	1.90	0.20		220	TYR	316	CZ2	OH	3.60
65.91	18.62	6.83	6.24	2.79					
lclc	1.90	0.20		529	HIS	170	CH2	CD2	3.85
65.78	75.40	7.03	6.41	-2.88					
lclc	1.90	0.20		544	TYR	499	CE3	CD2	3.93
89.97	164.51	5.48	5.48	0.00					
lcmb	1.80	0.20		102 A	TYR	104 A	CD1	CD1	3.72
75.96	-78.78	6.90	6.69	-1.67					
lcns	1.91	0.19		82 A	HIS	17 A	NE1	ND1	3.39
33.67	5.72	4.39	2.43	3.66					
lcns	1.91	0.19		158 A	PHE	44 A	CD1	CD1	3.76
41.97	-76.68	5.96	3.98	4.43					

1ede	1.90	0.16	125	PHE	128	CD2	CD2	3.35
8.59	-120.28	4.65	0.71	4.59				
1ede	1.90	0.16	125	PHE	222	CD2	CE1	3.43
23.97	-132.36	4.03	1.64	-3.68				
1ede	1.90	0.16	175	PHE	172	CD1	CE1	3.80
60.97	-91.59	6.32	5.53	3.07				
1ede	1.90	0.16	175	PHE	190	CZ3	CZ	3.67
24.94	114.59	5.43	2.29	-4.92				
1ede	1.90	0.16	175	TRP	194	CE3	CD2	3.82
72.56	157.74	5.54	5.29	-1.66				
1ede	1.90	0.16	194	TRP	175	CD2	CE3	3.82
6.53	127.65	5.54	0.67	-5.50				
1ede	1.90	0.16	175	HIS	289	CH2	CD2	3.56
48.91	77.63	5.64	4.25	3.71				
1ede	1.90	0.16	175	PHE	290	CH2	CZ	3.47
87.66	81.06	6.99	6.98	-0.28				
1ede	1.90	0.16	194	PHE	190	CD1	CE1	3.47
55.43	-82.25	5.11	4.21	-2.90				
1ede	1.90	0.16	244	PHE	243	CD2	CD2	3.81
2.82	167.95	5.05	0.14	5.05				
1ede	1.90	0.16	244	TRP	248	CZ3	CD2	3.42
72.96	120.36	5.57	5.32	1.63				
1ede	1.90	0.16	248	TRP	244	CD2	CZ3	3.42
23.39	-158.88	5.57	2.20	5.12				
1ede	1.90	0.16	248	PHE	243	CE3	CE2	3.73
70.75	173.76	6.19	5.84	2.04				
1fba	1.90	0.18	147 B	TYR	173 B	CE3	CD1	3.60
47.11	-160.35	4.70	3.44	3.20				
1fba	1.90	0.18	313 B	TYR	58 B	CE3	CD1	3.82
44.18	132.43	6.15	4.29	-4.41				
1fkj	1.70	0.16	59	TYR	26	CH2	CG	3.94
69.09	43.13	6.32	5.90	-2.26				
1fkj	1.70	0.16	59	PHE	48	CZ3	CE1	3.75
82.13	147.93	6.08	6.02	-0.83				
1fkj	1.70	0.16	59	PHE	99	NE1	CG	3.55
85.89	-23.10	4.90	4.89	-0.35				
1fnc	1.70	0.15	182	PHE	186	CE2	CD2	3.58
9.24	-30.89	4.93	0.79	4.87				
1fnc	1.70	0.15	199	PHE	201	CE2	CZ	3.54
32.59	-35.72	4.62	2.48	3.89				
1fnc	1.70	0.15	199	TYR	254	CZ2	CD1	3.74
85.37	23.77	5.55	5.53	0.45				
1fnc	1.70	0.15	259	TYR	299	CH2	CD2	3.73
58.36	69.81	6.43	5.47	-3.37				
1fnc	1.70	0.15	296	TYR	299	CZ3	CD2	3.61
68.64	129.48	6.84	6.37	-2.49				
1fnc	1.70	0.15	296	TRP	309	CH2	CZ3	3.64
79.37	36.36	6.24	6.13	-1.15				
1fnc	1.70	0.15	309	TRP	296	CZ3	CH2	3.64
29.31	64.64	6.24	3.06	5.44				
1gky	2.00	0.17	70	PHE	67	CD1	CD2	3.87
53.86	-120.45	5.92	4.78	3.49				
1gof	1.70	0.18	105	HIS	85	CD2	CE1	3.38
8.49	85.41	3.67	0.51	-3.63				
1gof	1.70	0.18	161	PHE	453	CZ2	CZ	3.88
54.17	32.23	6.64	5.38	3.89				
1gof	1.70	0.18	161	PHE	528	NE1	CD1	3.54
21.02	-15.13	4.95	1.78	4.62				
1gof	1.70	0.18	207	PHE	508	CE3	CE1	3.63
55.47	145.54	6.01	4.95	-3.40				

lgof	1.70	0.18		290	PHE	194	CZ2	CE1	3.94
80.86	29.35	6.59	6.51	-1.05					
lgof	1.70	0.18		290	PHE	227	CH2	CD2	3.68
47.65	56.91	6.26	4.63	-4.22					
lgof	1.70	0.18		290	TYR	272	CD1	CE2	3.40
32.01	-114.83	4.08	2.17	-3.46					
lgof	1.70	0.18		290	TYR	329	CD1	OH	3.38
58.53	-90.64	6.41	5.47	3.35					
lgox	2.00	0.19		33	TYR	261	CG	CE1	3.45
36.65	-161.20	4.93	2.94	-3.96					
lgpl	2.00	0.17		148 A	PHE	77 A	CH2	CE1	3.82
51.11	100.38	6.42	5.00	-4.03					
lgpl	2.00	0.17		158 A	PHE	145 A	CH2	CD2	3.89
75.32	81.49	6.42	6.21	1.63					
lgpb	1.90	0.19		189	TYR	52	NE1	OH	3.67
64.88	-7.10	7.30	6.61	3.10					
lgpb	1.90	0.19		215	TYR	297	CZ3	OH	3.77
74.39	82.87	7.60	7.32	-2.05					
lgpb	1.90	0.19		244	TYR	157	NE1	CD1	3.38
67.50	-11.74	5.80	5.36	2.22					
lgpb	1.90	0.19		825	TYR	820	CZ2	CD1	3.72
42.52	33.28	5.66	3.83	-4.17					
lgse	2.00	0.20		21 B	TYR	165 B	CH2	CD1	4.00
84.35	46.85	6.81	6.78	-0.67					
lgse	2.00	0.20		21 B	PHE	197 B	CD1	CE2	3.29
21.66	-101.76	4.90	1.81	4.55					
lhan	1.90	0.16		19	PHE	11	CH2	CD1	3.88
58.93	78.39	5.87	5.03	-3.03					
lhan	1.90	0.16		48	TYR	8	CH2	CG	3.42
57.85	51.22	5.39	4.56	-2.87					
lhan	1.90	0.16		272	HIS	194	CH2	CD2	3.71
51.52	46.07	6.01	4.70	3.74					
lhan#	1.90	0.16		283	PHE	223	CD1	CE1	3.96
73.35	-88.66	5.77	5.53	1.65					
lhbq	1.70	0.20		24	PHE	20	NE1	CE2	3.72
50.68	-20.85	4.56	3.53	-2.89					
lhbq	1.70	0.20		24	PHE	45	CD1	CD1	3.78
64.57	-77.89	6.54	5.91	-2.81					
lhle	1.95	0.18		160 A	TYR	119 A	CG	OH	3.40
33.38	-170.61	6.43	3.54	-5.37					
lhle	1.95	0.18		194 A	TYR	244 A	CH2	OH	3.43
48.02	125.14	5.24	3.90	3.50					
lhpm	1.70	0.20		90	PHE	68	CD1	CE1	3.84
57.20	-75.26	6.22	5.23	3.37					
lhpm	1.70	0.20		90	PHE	92	CG	CZ	3.45
36.39	-151.34	4.77	2.83	3.84					
lhsl	1.89	0.20		130 A	PHE	125 A	CE2	CE1	3.79
12.66	-18.98	5.26	1.16	-5.13					
lhuw	2.00	0.19		86	PHE	166	CD2	CD1	3.69
17.78	116.80	5.01	1.52	-4.78					
lhxn	1.80	0.18		353	TRP	365	CG	CE3	3.67
22.50	-149.35	5.63	2.15	5.20					
lhxn	1.80	0.18		365	TRP	353	CE3	CG	3.67
80.77	156.23	5.63	5.56	-0.90					
lhxn	1.80	0.18		370	HIS	372	CE3	CG	3.94
46.14	140.75	5.75	4.14	-3.99					
lilb#	2.00	0.19		120	PHE	150	CZ2	CZ	3.91
50.99	-2.57	5.69	4.42	-3.58					
liae	1.83	0.14		10	TYR	194	CD1	OH	3.52
87.65	-111.12	7.95	7.94	-0.32					

liae	1.83	0.14	65	PHE	58	CZ2	CZ	3.91
46.43	29.83	6.34	4.59	-4.37				
lisc	1.80	0.19	71 A	PHE	75 A	CH2	CD1	3.44
58.50	81.16	5.97	5.09	3.12				
lisc	1.80	0.19	77 A	TYR	9 A	CE3	OH	3.75
59.66	178.03	7.10	6.13	-3.59				
lisc	1.80	0.19	77 A	TYR	27 A	NE1	CD1	3.53
32.93	-34.47	5.61	3.05	-4.71				
lisc	1.80	0.19	77 A	HIS	31 A	NE1	ND1	3.06
84.64	-41.31	5.78	5.75	0.54				
lisc	1.80	0.19	77 A	TYR	76 A	CZ3	CE2	3.59
29.62	123.81	5.38	2.66	4.68				
lisc	1.80	0.19	122 A	TRP	158 A	NE1	NE1	3.74
49.53	-22.64	6.35	4.83	4.12				
lisc	1.80	0.19	158 A	TRP	122 A	NE1	NE1	3.74
66.36	-56.75	6.35	5.82	-2.55				
lisc	1.80	0.19	158 A	PHE	118 B	CH2	CZ	3.82
89.00	53.92	6.40	6.40	-0.11				
lisc	1.80	0.19	124 A	PHE	75 A	CZ3	CD2	3.97
84.66	121.07	6.94	6.91	0.65				
lisc	1.80	0.19	178 A	PHE	106 A	CH2	CE2	3.70
70.43	75.57	6.85	6.45	-2.29				
lisc	1.80	0.19	178 A	PHE	110 A	CZ2	CD1	3.56
64.23	50.62	6.15	5.54	2.67				
lisc	1.80	0.19	183 A	PHE	103 A	CZ2	CE1	3.68
57.55	30.87	6.24	5.27	-3.35				
1knb	1.70	0.16	478	PHE	461	NE1	CE1	3.95
58.19	-9.79	6.30	5.35	3.32				
1knb	1.70	0.16	478	TYR	491	NE1	OH	3.01
58.50	-59.71	6.33	5.40	-3.31				
1knb	1.70	0.16	556	PHE	568	CG	CZ	3.66
25.67	-150.83	5.50	2.38	4.96				
1knb	1.70	0.16	558	TYR	563	CD1	CD2	3.63
47.69	-111.67	5.94	4.39	-4.00				
1knb	1.70	0.16	558	PHE	568	CZ3	CE1	3.54
57.97	97.25	6.11	5.18	-3.24				
1lcp	1.65	0.16	207 A	PHE	229 A	CZ3	CG	3.75
72.65	97.58	6.49	6.19	-1.94				
1lcp	1.65	0.16	207 A	TYR	345 A	CZ2	OH	3.44
85.20	49.34	6.08	6.06	0.51				
1lcp	1.65	0.16	381 A	PHE	482 A	CH2	CE2	3.79
87.57	55.08	7.39	7.38	-0.31				
1lcp	1.65	0.16	383 A	PHE	387 A	CZ2	CD1	3.64
52.78	14.28	5.51	4.39	-3.34				
1lcp	1.65	0.16	398 A	PHE	403 B	CZ3	CZ	3.94
52.82	129.29	6.42	5.12	3.88				
1lct	2.00	0.18	8	TRP	22	CD2	CZ3	3.73
25.60	170.99	6.45	2.79	5.82				
1lct	2.00	0.18	22	TRP	8	CZ3	CD2	3.73
88.62	101.03	6.45	6.45	0.16				
1lct	2.00	0.18	8	PHE	300	CZ2	CE1	3.62
39.49	16.69	5.45	3.47	4.20				
1lct	2.00	0.18	22	PHE	286	CD1	CE2	3.75
55.23	-108.39	6.17	5.07	-3.52				
1lct	2.00	0.18	22	PHE	300	CH2	CZ	3.71
43.17	83.74	6.31	4.32	-4.60				
1lis	1.90	0.19	34	HIS	38	CE2	CD2	3.35
25.50	-7.13	4.31	1.86	3.89				
1lis	1.90	0.19	34	TYR	102	CG	CD2	3.87
20.17	-178.93	5.25	1.80	-4.93				

11is	1.90	0.19	62	TYR	133	NE1	CE2	3.49
83.90	-26.96	5.23	5.20	0.56				
1mml	1.80	0.20	185	PHE	156	CZ3	CZ	3.84
65.75	122.49	6.77	6.17	-2.78				
1mml	1.80	0.20	185	PHE	170	NE1	CE1	3.65
48.99	-20.27	5.89	4.44	3.87				
1mml#	1.80	0.20	185	TYR	271	CZ3	OH	3.36
86.89	124.83	8.06	8.05	-0.44				
1mpp	2.00	0.16	39	TYR	75	NE1	OH	2.68
83.75	-17.80	6.43	6.39	0.70				
1mpp	2.00	0.16	190	TYR	125	CZ3	OH	3.49
17.94	52.80	4.86	1.50	4.62				
1mpp	2.00	0.16	190	TYR	313	CD2	CD1	3.68
4.61	-119.39	5.01	0.39	-4.99				
1mrj	1.60	0.17	192	TYR	164	CD1	OH	3.76
62.22	-95.12	6.39	5.65	-2.98				
1msc	2.00	0.20	32	PHE	4	CZ2	CD1	3.87
10.23	19.10	5.22	0.94	-5.14				
1msc	2.00	0.20	32	PHE	7	NE1	CD1	3.61
22.89	-64.33	5.02	1.94	4.63				
1nar	1.80	0.16	58	TYR	116	NE1	OH	2.99
84.02	4.68	6.15	6.12	-0.64				
1nar	1.80	0.16	101	PHE	93	CZ2	CE1	3.69
25.63	39.33	5.43	2.35	-4.90				
1nar	1.80	0.16	101	TYR	131	CZ2	OH	3.70
71.27	26.36	7.44	7.05	-2.39				
1nar	1.80	0.16	261	TYR	8	CE3	CG	3.64
83.81	147.05	5.69	5.66	-0.61				
1nar	1.80	0.16	261	PHE	39	CH2	CE2	3.69
86.80	86.28	7.07	7.06	0.39				
1nba	2.00	0.19	56 A	PHE	63 A	NE1	CD2	3.52
50.92	-5.13	5.54	4.30	-3.49				
1nba	2.00	0.19	111 A	TYR	96 A	CE3	OH	3.34
73.62	159.69	7.67	7.36	2.16				
1nba	2.00	0.19	126 A	HIS	61 A	CD2	CD2	3.51
17.65	-161.90	4.61	1.40	-4.39				
1nba	2.00	0.19	126 A	TYR	125 A	CD2	CD2	3.83
16.83	135.67	5.26	1.54	5.03				
1nba	2.00	0.19	209 A	PHE	23 B	CH2	CE1	3.65
54.52	65.00	6.74	5.49	-3.91				
1nba	2.00	0.19	209 A	TYR	212 B	CZ2	CD2	3.88
77.78	32.54	6.67	6.52	-1.41				
1nfp	1.60	0.17	4	PHE	194	CE3	CE2	3.79
43.94	158.12	6.13	4.26	4.41				
1nfp	1.60	0.17	96	PHE	9	CD1	CZ	3.56
46.67	-87.40	5.81	4.23	3.99				
1nfp	1.60	0.17	96	PHE	131	CD2	CE1	3.50
16.76	131.97	4.82	1.38	4.62				
1nfp	1.60	0.17	96	TYR	152	CH2	CE1	3.36
79.26	37.73	5.78	5.68	-1.08				
1nif	1.60	0.17	265	PHE	270	CD2	CD1	3.79
9.26	-162.44	5.01	0.83	4.94				
1onc	1.70	0.18	3	PHE	6	CZ3	CD2	3.67
76.51	130.93	6.84	6.65	-1.60				
1ora	2.00	0.18	35	TYR	107	CE3	CE1	3.77
53.06	-176.68	5.28	4.22	-3.17				
1ora	2.00	0.18	275	TRP	278	CZ3	CD1	4.00
64.28	133.57	7.72	6.95	-3.35				
1ora	2.00	0.18	278	TRP	275	CD1	CZ3	4.00
76.00	-103.22	7.72	7.49	1.87				

lora	2.00	0.18	278	TYR	164	CE3	CD1	3.34	
49.50	124.69	5.62	4.27	-3.65	TYR	271	CE3	OH	3.52
lora	2.00	0.18	278	PHE	101	CE3	CE1	3.71	
65.68	161.52	6.50	5.92	2.67	PHE	284	CZ3	CE1	3.51
loyc	2.00	0.17	97	PHE	74	CE3	CE1	3.41	
61.10	180.00	5.59	4.89	-2.70	TYR	82	CH2	CE2	3.95
loyc	2.00	0.17	112	HIS	191	CD1	ND1	3.38	
33.65	129.40	5.61	3.11	-4.67	TYR	196	NE1	CD2	3.33
loyc	2.00	0.17	116	PHE	208	CZ2	CE2	3.43	
46.77	148.64	5.44	3.96	3.73	TYR	222	CE3	CD1	3.51
loyc	2.00	0.17	116	HIS	351	NE1	NE2	2.83	
50.63	62.77	6.73	5.20	4.27	TYR	221	CH2	CD2	3.47
loyc	2.00	0.17	116	PHE	177	CZ2	CG	3.69	
53.73	-63.51	5.61	4.52	-3.32	TYR	181	CD1	OH	3.68
loyc	2.00	0.17	116	TYR	371	CZ2	CD1	3.68	
82.10	-17.66	4.75	4.70	0.65	HIS	20	NE1	CE1	3.61
1pbe	1.90	0.16	185	PHE	48	CZ3	CD1	3.87	
38.09	0.63	4.99	3.08	3.92	TYR	5	CE3	CD2	3.40
1pbe	1.90	0.16	185	TYR	16	CZ3	CE1	3.94	
69.33	146.18	5.32	4.98	1.88	TRP	168	CH2	CZ3	3.56
1pbe	1.90	0.16	185	TRP	156	CZ3	CH2	3.56	
82.31	-28.91	5.48	5.43	0.73	TYR	104	CZ3	CD2	3.80
1pbe	1.90	0.16	240	TYR	193	CZ2	OH	3.31	
84.30	95.78	6.49	6.46	0.64	PHE	224	CD1	CE1	3.49
1pbe	1.90	0.16	337	TYR	23	CD2	CD1	3.34	
76.21	20.90	5.63	5.47	-1.34	PHE	70	CD1	CE2	3.53
1pbe	1.90	0.16	344	PHE	72	CZ2	CE2	3.70	
77.87	-89.55	7.85	7.67	1.65	TYR	281	CH2	CE2	3.76
1pbe	1.90	0.16	345	TYR	321	CE3	CE1	3.71	
51.91	15.49	5.10	4.02	-3.14	TYR	22	CH2	CD1	3.77
1pbn	2.00	0.20	16	HIS	20	NE1	CE1	3.61	
15.58	8.06	4.17	1.13	4.01	PHE	48	CZ3	CD1	3.87
1pbn	2.00	0.20	16	TYR	5	CE3	CD2	3.40	
70.79	101.03	6.75	6.37	2.22	TYR	16	CZ3	CE1	3.94
1pbn	2.00	0.20	94	TRP	168	CH2	CZ3	3.56	
85.07	134.77	6.14	6.12	0.53	TRP	156	CZ3	CH2	3.56
1pbb	1.90	0.15	19	TYR	104	CZ3	CD2	3.80	
57.92	131.52	6.65	5.63	-3.53	TYR	193	CZ2	OH	3.31
1pbb	1.90	0.15	156	PHE	224	CD1	CE1	3.49	
86.91	44.58	7.19	7.18	-0.39	TYR	23	CD2	CD1	3.34
1pbb	1.90	0.15	168	PHE	70	CD1	CE2	3.53	
48.55	119.00	7.19	5.39	4.76	PHE	72	CZ2	CE2	3.70
1pbb	1.90	0.15	168	TYR	281	CH2	CE2	3.76	
67.54	121.12	7.04	6.51	-2.69	TYR	321	CE3	CE1	3.71
1pbb	1.90	0.15	168	TYR	22	CH2	CD1	3.77	
57.09	62.51	6.75	5.67	-3.67	PHE	224	CD1	CE1	3.49
1pbb	1.90	0.15	250	TYR	23	CD2	CD1	3.34	
40.70	-106.92	5.73	3.74	4.34	PHE	70	CD1	CE2	3.53
1pbb	1.90	0.15	279	PHE	72	CZ2	CE2	3.70	
21.47	-175.70	4.68	1.72	-4.35	TYR	281	CH2	CE2	3.76
1pbb	1.90	0.15	308	TYR	321	CE3	CE1	3.71	
75.77	-74.00	6.52	6.32	-1.60	TYR	22	CH2	CD1	3.77
1pbb	1.90	0.15	308	PHE	224	CD1	CE1	3.49	
85.32	-12.65	4.98	4.96	-0.41	PHE	70	CD1	CE2	3.53
1pbb	1.90	0.15	308	PHE	72	CZ2	CE2	3.70	
60.29	53.67	6.86	5.96	-3.40	TYR	281	CH2	CE2	3.76
1pbb	1.90	0.15	308	TYR	321	CE3	CE1	3.71	
76.76	157.45	5.28	5.14	1.21	TYR	22	CH2	CD1	3.77
1pda	1.76	0.19	18	PHE	224	CD1	CE1	3.49	
63.34	87.24	6.48	5.79	2.91	TYR	23	CD2	CD1	3.34

lpgs	1.80	0.17	191	PHE	292	CE3	CD2	3.79
57.45	146.28	6.54	5.51	-3.52				
lpgs	1.80	0.17	282	TYR	161	CE3	CD2	3.78
59.86	170.94	5.37	4.65	-2.69				
lphg	1.60	0.19	42	PHE	26	CD2	CZ	3.72
9.55	171.45	4.85	0.82	4.78				
lphg	1.60	0.19	55	TRP	63	CZ3	NE1	3.34
80.59	148.36	5.12	5.05	0.84				
lphg	1.60	0.19	63	TRP	55	NE1	CZ3	3.34
6.41	15.86	5.12	0.61	5.08				
lphg	1.60	0.19	374	PHE	150	CZ3	CE1	3.58
38.46	101.37	5.96	3.71	4.67				
lphg	1.60	0.19	374	PHE	381	NE1	CE1	3.46
65.60	-35.86	5.16	4.70	2.13				
lpii	2.00	0.17	15	PHE	97	CH2	CE2	3.56
65.25	62.37	6.67	6.06	-2.79				
lpii	2.00	0.17	356	TYR	278	CZ2	CE1	3.63
85.71	49.60	5.67	5.65	-0.42				
lpii	2.00	0.17	391	PHE	389	CD1	CE1	3.73
88.12	-67.17	5.85	5.85	0.19				
lpne	2.00	0.17	3	TYR	6	CE3	CD2	3.52
49.42	131.68	5.42	4.12	3.52				
lpne	2.00	0.17	3	TRP	31	CZ2	CZ3	3.79
9.38	-14.66	5.40	0.88	-5.33				
lpne	2.00	0.17	31	TRP	3	CZ3	CZ2	3.79
66.99	154.33	5.40	4.97	2.11				
lpne	2.00	0.17	3	TYR	139	CZ2	CE1	3.63
84.54	42.80	5.70	5.67	0.54				
lpoc	2.00	0.19	128	TYR	96	CZ3	CD2	3.59
47.84	89.74	6.41	4.75	4.30				
lppn	1.60	0.16	69	TYR	67	NE1	CD1	3.57
48.57	14.67	4.89	3.67	-3.23				
lppn	1.60	0.16	69	PHE	207	CD1	CE1	3.52
64.33	-117.88	5.61	5.05	2.43				
lppn	1.60	0.16	177	PHE	141	CH2	CE1	3.72
56.33	86.00	6.69	5.57	-3.71				
lppn	1.60	0.16	177	HIS	159	NE1	CE1	3.36
48.90	-4.70	4.74	3.57	-3.11				
lppn	1.60	0.16	177	TRP	181	CE3	NE1	3.53
71.39	149.03	5.36	5.08	-1.71				
lppn	1.60	0.16	181	TRP	177	NE1	CE3	3.53
8.85	-80.97	5.36	0.84	-5.29				
lppn	1.60	0.16	181	PHE	141	CZ2	CE1	3.78
69.25	29.43	5.93	5.55	-2.10				
lppn	1.60	0.16	181	TYR	144	CZ2	CD2	3.71
36.72	48.87	5.79	3.46	4.64				
lrcf	1.40	0.14	120	TYR	125	NE1	OH	2.95
85.49	-49.83	6.80	6.78	-0.53				
lrcf	1.40	0.14	120	TRP	159	CE3	CD1	3.74
38.27	159.14	6.49	4.02	5.09				
lrcf	1.40	0.14	159	TRP	120	CD1	CE3	3.74
52.47	-97.68	6.49	5.15	-3.95				
lrci	2.00	0.19	89	TYR	36	CD1	CG	3.97
81.29	-110.32	6.58	6.50	1.00				
lrec	1.90	0.19	31	PHE	23	CZ3	CE2	3.58
65.93	135.16	5.74	5.24	-2.34				
lrec	1.90	0.19	31	PHE	83	CG	CZ	3.80
15.56	-122.02	5.19	1.38	5.00				
lrec	1.90	0.19	31	TYR	86	NE1	OH	2.98
54.30	1.99	5.44	4.42	3.17				

lrec	1.90	0.19	104	PHE	73	NE1	CZ	3.72
71.22	-36.64	5.79	5.48	-1.86				
lrec	1.90	0.19	104	PHE	188	CZ3	CE1	3.39
87.86	120.88	6.12	6.12	0.23				
lrva	2.00	0.16	216 A	TYR	110 A	CZ3	CD2	3.62
77.24	137.41	5.98	5.83	-1.32				
lrva	2.00	0.16	216 A	PHE	206 A	NE1	CZ	3.84
70.38	-53.27	6.50	6.12	2.18				
lrva	2.00	0.16	216 A	PHE	212 A	CD1	CE1	3.65
67.87	-88.48	6.37	5.90	-2.40				
lrva	2.00	0.16	216 A	TYR	215 A	NE1	CD1	3.61
16.89	-36.06	4.96	1.45	4.74				
lrva	2.00	0.16	216 A	TYR	219 A	CZ3	CD2	3.44
55.55	105.08	5.28	4.36	2.98				
lrva	2.00	0.16	239 A	TYR	230 A	CE3	CE2	3.71
27.78	145.72	5.27	2.46	-4.66				
lrva	2.00	0.16	239 A	TYR	236 A	CE3	CD1	3.93
77.20	147.06	6.57	6.41	-1.45				
lsac	2.00	0.18	98 A	PHE	33 A	CE3	CE2	3.92
74.53	-178.79	6.19	5.97	-1.65				
lsac	2.00	0.18	185 A	PHE	9 A	CH2	CE2	3.40
68.86	67.61	6.18	5.76	-2.23				
lsac	2.00	0.18	203 A	TYR	40 A	CG	CD2	3.60
18.57	-144.90	4.93	1.58	4.67				
lsat	1.75	0.19	46	TYR	246	CD1	OH	3.39
85.22	-104.93	7.60	7.57	-0.63				
lsat	1.75	0.19	153	TYR	65	CH2	CZ	3.34
69.93	71.20	5.69	5.34	1.95				
lsat	1.75	0.19	153	TYR	135	CD1	CE1	3.46
64.22	-121.93	5.88	5.29	-2.56				
lsat	1.75	0.19	217	TYR	190	NE1	OH	3.72
70.06	-3.91	5.12	4.81	1.75				
lsat	1.75	0.19	217	TYR	199	CD1	CD1	3.76
45.77	-109.66	6.28	4.50	-4.38				
lsbp	1.70	0.18	28	PHE	24	CE3	CZ	3.78
47.35	-179.81	5.55	4.08	3.76				
lsbp	1.70	0.18	112	PHE	99	CD2	CZ	3.56
9.70	109.82	4.82	0.81	4.75				
lsbp	1.70	0.18	112	TYR	143	CD1	CD2	3.37
68.00	-116.42	5.62	5.21	-2.10				
lsbp	1.70	0.18	135	TRP	192	CD1	CH2	3.42
61.82	-86.34	6.90	6.08	-3.26				
lsbp	1.70	0.18	141	TYR	137	CZ3	OH	3.66
69.35	155.56	6.30	5.90	2.22				
lsbp	1.70	0.18	141	PHE	287	CD1	CE1	3.64
50.82	-100.13	6.38	4.95	4.03				
lsbp	1.70	0.18	192	TYR	10	CZ2	CE2	3.77
39.03	27.19	5.60	3.53	-4.35				
lsbp	1.70	0.18	192	TRP	135	CH2	CD1	3.42
57.13	77.06	6.90	5.80	3.74				
lsbp	1.70	0.18	135	TRP	290	CH2	NE1	3.49
32.30	59.97	5.47	2.92	4.62				
lsbp	1.70	0.18	290	TRP	135	NE1	CH2	3.49
88.90	12.50	5.47	5.47	0.10				
lsbp	1.70	0.18	141	HIS	297	CE2	CE1	3.70
10.39	110.70	4.64	0.86	4.56				
lscs	1.60	0.18	40	TYR	67	CZ2	OH	3.54
71.13	-12.06	6.13	5.80	1.98				
lscs	1.60	0.18	40	PHE	197	CZ2	CE2	3.91
69.31	29.10	6.59	6.16	-2.33				

lscs	1.60	0.18		88	HIS	180	CE2	CD2	3.51
29.48	18.47	3.93	1.93	3.42					
lscs	1.60	0.18		109	PHE	128	CZ2	CZ	3.74
10.81	57.63	5.00	0.95	4.91					
lscs#	1.60	0.18		182	TYR	12	CZ2	OH	3.32
86.24	26.77	6.67	6.66	0.44					
lscs#	1.60	0.18		182	TYR	100	NE1	OH	2.98
82.50	-16.25	6.30	6.25	-0.82					
lscs#	1.60	0.18		182	HIS	205	CE2	NE2	3.43
18.68	25.22	3.67	1.18	-3.47					
lslt	1.90	0.17		68 A	HIS	44 A	CH2	CG	3.38
68.19	77.51	5.43	5.04	-2.02					
lsri	1.65	0.17		21 A	PHE	29 A	CD1	CD2	3.89
44.45	-96.99	6.05	4.24	-4.32					
lsri	1.65	0.17		21 A	TYR	96 A	CH2	CE1	3.76
76.94	50.81	7.01	6.83	-1.59					
lsri	1.65	0.17		75 A	PHE	29 A	CE2	CE1	3.66
25.47	41.72	4.28	1.84	-3.86					
lsri	1.65	0.17		75 A	TRP	92 A	NE1	CE3	3.05
47.68	-11.20	4.55	3.37	3.06					
lsri	1.65	0.17		92 A	TRP	75 A	CE3	NE1	3.05
43.22	-177.39	4.55	3.12	3.31					
lsri	1.65	0.17		75 A	PHE	130 A	CH2	CE2	3.83
85.53	39.30	6.34	6.32	-0.49					
lsri	1.65	0.17		79 A	TYR	43 A	CH2	CD1	3.63
77.83	64.12	6.90	6.75	1.45					
lsri	1.65	0.17		79 A	TRP	92 A	CZ2	CH2	3.89
74.81	20.49	7.88	7.60	2.06					
lsri	1.65	0.17		92 A	TRP	79 A	CH2	CZ2	3.89
74.64	71.27	7.88	7.60	-2.09					
lsri	1.65	0.17		79 A	TYR	54 A	CZ3	CD1	3.69
78.61	150.28	5.46	5.35	1.08					
lsri	1.65	0.17		92 A	TYR	43 A	CZ2	OH	3.44
49.38	50.93	5.42	4.11	3.53					
lsri	1.65	0.17		92 A	TRP	108 A	NE1	CD1	3.84
38.76	-35.59	6.14	3.85	-4.79					
lsri	1.65	0.17		108 A	TRP	92 A	CD1	NE1	3.84
47.81	-121.26	6.14	4.55	4.12					
lsri	1.65	0.17		92 A	PHE	130 A	NE1	CZ	3.86
40.18	-54.09	6.07	3.91	4.64					
ltca	1.55	0.16		52	PHE	131	CZ2	CZ	3.62
45.79	-16.33	5.16	3.70	-3.60					
ltca	1.55	0.16		52	TYR	234	CH2	CD1	3.76
51.57	68.36	6.63	5.19	4.12					
ltca	1.55	0.16		104	HIS	224	CZ2	CD2	3.73
50.90	-13.20	4.97	3.86	3.13					
ltca	1.55	0.16		113	PHE	117	CE2	CD2	3.63
16.18	47.45	4.56	1.27	-4.38					
ltca	1.55	0.16		113	TYR	300	NE1	OH	3.00
64.61	-12.17	6.10	5.51	-2.61					
ltml	1.80	0.18		16	PHE	274	CG	CE2	3.91
37.39	-142.57	5.80	3.52	-4.61					
ltml	1.80	0.18		41	TYR	73	CZ3	CG	3.92
88.82	130.35	6.20	6.20	0.13					
ltml	1.80	0.18		97	TYR	94	CE3	CE1	3.72
72.53	150.89	6.15	5.87	-1.85					
ltml	1.80	0.18		97	PHE	101	CH2	CD2	3.90
42.15	68.03	6.28	4.22	4.65					
ltml	1.80	0.18		162	HIS	159	CE3	NE2	3.36
63.02	161.41	4.97	4.43	-2.25					

1tml	1.80	0.18	162	HIS	163	NE1	NE2	3.55		
28.86	-63.66	4.40	2.13	3.85						
1ton	1.80	0.20	51	PHE	89	CZ3	CE1	3.38		
85.18	93.50	6.87	6.85	0.58						
1ton	1.80	0.20	51	HIS	107	CE2	CD2	3.42		
35.62	26.68	4.06	2.36	3.30						
1ton	1.80	0.20	237	HIS	91	CD2	CD2	3.56		
7.66	-174.47	4.61	0.58	-4.57						
1ton#	1.80	0.20	237	TYR	172	CD1	CE1	3.60		
68.15	-78.10	6.64	6.16	2.47						
1tph	1.80	0.19	90	1	PHE	7	CZ2	CE2	3.86	
29.11	58.90	5.84	2.84	-5.10						
1tph	1.80	0.19	191	1	PHE	144	1	CD1	CG	3.49
66.69	-97.71	5.54	5.09	-2.19						
1tph	1.80	0.19	191	1	HIS	195	1	CD1	CE1	3.35
22.81	-38.31	3.75	1.46	3.45						
1trk	2.00	0.16	45	B	PHE	63	B	CZ2	CD2	3.59
72.94	-11.67	4.95	4.73	1.45						
1trk	2.00	0.16	45	B	TYR	83	B	CZ3	CE1	3.96
79.64	106.85	7.46	7.34	1.34						
1trk	2.00	0.16	57	B	TRP	312	B	NE1	CG	3.41
86.34	-34.67	5.47	5.46	0.35						
1trk	2.00	0.16	312	B	TRP	57	B	CG	NE1	3.41
33.41	-143.26	5.47	3.01	-4.57						
1trk	2.00	0.16	212	B	TYR	207	B	CD2	CD2	3.58
6.67	168.03	4.87	0.59	-4.83						
1trk	2.00	0.16	312	B	PHE	316	B	CH2	CD2	3.89
55.36	72.77	6.21	5.11	3.53						
1trk	2.00	0.16	623	B	PHE	591	B	CH2	CE1	3.75
22.91	77.90	5.23	2.03	4.82						
1tsp	2.00	0.18	207	TYR	328	CH2	CE1	3.61		
72.54	49.91	6.90	6.58	-2.07						
1tsp	2.00	0.18	315	TYR	377	CZ3	CD2	3.69		
55.96	89.41	6.72	5.57	-3.76						
1tys	1.80	0.18	61	PHE	62	CZ3	CE1	3.53		
66.20	145.76	6.04	5.53	2.44						
1tys	1.80	0.18	61	TYR	71	CD1	CD2	3.75		
67.45	-106.64	6.59	6.09	-2.53						
1tys	1.80	0.18	61	TRP	80	CZ3	NE1	3.65		
73.63	90.48	6.67	6.40	-1.88						
1tys	1.80	0.18	80	TRP	61	NE1	CZ3	3.65		
40.38	-77.00	6.67	4.32	-5.08						
1tys	1.80	0.18	98	PHE	62	CH2	CD2	3.83		
33.42	65.00	5.76	3.18	-4.80						
1tys	1.80	0.18	98	PHE	150	CZ2	CE1	3.71		
34.81	11.81	5.45	3.11	4.47						
1tys	1.80	0.18	98	TYR	181	CZ2	OH	3.59		
75.33	36.87	7.16	6.93	1.81						
1tys#	1.80	0.18	133	PHE	149	CG	CE2	3.77		
46.87	119.21	7.25	5.29	-4.95						
1tys	1.80	0.18	201	HIS	32	CZ3	CD2	3.99		
84.40	125.92	7.33	7.30	-0.71						
1udg	1.75	0.20	19	PHE	22	CZ3	CE2	3.88		
45.37	120.83	6.25	4.45	-4.39						
1udg	1.75	0.20	32	TRP	71	CZ3	CZ3	3.79		
16.15	83.90	6.34	1.77	-6.09						
1udg	1.75	0.20	71	TRP	32	CZ3	CZ3	3.79		
69.14	125.83	6.34	5.93	-2.26						
1udg	1.75	0.20	71	PHE	69	CZ2	CD1	3.57		
55.62	-3.06	5.74	4.74	-3.24						

1udg	1.75	0.20	138	TRP	242	CZ2	CE3	3.82	
20.85	19.75	5.71	2.03	5.34	TRP	138	CE3	CZ2	3.82
1udg	1.75	0.20	242	PHE	69	CZ3	CZ	3.96	
83.05	153.55	5.71	5.67	0.69	PHE	101	CE3	CZ	3.79
1udg	1.75	0.20	188	PHE	219	CE2	CZ	3.61	
59.97	95.20	7.11	6.15	-3.56	HIS	224	NE1	ND1	3.07
1udg	1.75	0.20	188	PHE	225	CD2	CD1	3.89	
83.43	133.54	6.33	6.29	-0.72	PHE	74	CH2	CE1	3.68
1udg	1.75	0.20	188	PHE	48	NE1	CD2	3.67	
3.92	-145.03	4.86	0.33	4.85	TYR	175	CH2	CG	3.68
1udg	1.75	0.20	188	HIS	183	CZ3	CD2	3.62	
84.68	-43.41	5.77	5.75	0.54	PHE	92 A	CD1	CD1	3.68
1udg	1.75	0.20	242	PHE	411 B	CE3	CG	3.41	
12.52	141.97	5.19	1.14	-5.06	TYR	32 A	NE1	CE1	3.62
1vhh	1.70	0.19	129	PHE	122 A	CD1	CZ	3.53	
51.72	78.79	6.52	5.12	-4.04	PHE	130 A	NE1	CE1	3.23
1vhh	1.70	0.19	173	TYR	133 A	CH2	CD2	3.60	
67.35	3.98	5.51	5.09	2.12	PHE	189 A	CZ3	CE1	3.90
1vhh	1.70	0.19	173	TYR	203 A	CG	CD2	3.26	
60.83	72.67	6.24	5.45	3.04	TYR	147 A	NE1	OH	3.52
1vhh	1.70	0.19	173	HIS	151 A	CH2	OH	3.59	
87.91	131.46	6.55	6.55	0.24	PHE	414 B	CG	CD1	3.41
1wht	2.00	0.18	49 A	PHE	380 B	NE1	CG	3.23	
59.14	-111.33	6.37	5.47	3.27	TRP	369 B	CG	NE1	3.23
1wht	2.00	0.18	84 A	TYR	348 B	CH2	CZ	3.61	
42.98	159.87	4.23	2.89	3.09	TYR	26	CZ3	CG	3.97
1wht	2.00	0.18	126 A	PHE	36	CD2	CZ	3.50	
83.06	-46.96	5.17	5.13	-0.62	TYR	69	NE1	OH	3.01
1wht	2.00	0.18	126 A	PHE	125	CH2	CE1	3.66	
56.81	-114.57	5.86	4.90	-3.21	TYR	166	NE1	OH	3.34
1wht	2.00	0.18	126 A	TRP	242	CZ2	CE3	3.82	
35.53	11.58	4.11	2.39	3.34	TRP	138	CE3	CZ2	3.82
1wht	2.00	0.18	126 A	PHE	69	CZ3	CZ	3.96	
82.43	75.69	6.60	6.54	0.87	PHE	101	CE3	CZ	3.79
1wht	2.00	0.18	192 A	PHE	219	CE2	CZ	3.61	
33.84	121.29	5.93	3.30	-4.92	HIS	224	NE1	ND1	3.07
1wht	2.00	0.18	193 A	PHE	225	CD2	CD1	3.89	
41.10	-150.43	4.74	3.11	-3.57	PHE	74	CH2	CE1	3.68
1wht	2.00	0.18	308 B	PHE	48	NE1	CD2	3.67	
27.06	-22.59	5.58	2.54	4.97	TYR	175	CH2	CG	3.68
1wht	2.00	0.18	308 B	HIS	183	CZ3	CD2	3.62	
56.02	42.57	6.81	5.65	3.80	PHE	92 A	CD1	CD1	3.68
1wht	2.00	0.18	331 B	PHE	411 B	CE3	CG	3.41	
26.14	-141.51	5.26	2.32	-4.72	TYR	32 A	NE1	CE1	3.62
1wht	2.00	0.18	369 B	PHE	122 A	CD1	CZ	3.53	
65.00	-31.93	5.69	5.16	-2.41	PHE	130 A	NE1	CE1	3.23
1wht	2.00	0.18	380 B	TYR	133 A	CH2	CD2	3.60	
56.18	-126.75	5.69	4.73	3.17	PHE	189 A	CZ3	CE1	3.90
1wht#	2.00	0.18	193 A	TYR	203 A	CG	CD2	3.26	
83.87	78.65	6.16	6.12	-0.66	TYR	147 A	NE1	OH	3.52
1xnb	1.49	0.17	6	TYR	151 A	CH2	OH	3.59	
82.94	109.90	7.22	7.17	-0.89	PHE	414 B	CG	CD1	3.41
1xnb	1.49	0.17	30	TRP	380 B	NE1	CG	3.23	
21.49	-129.51	4.84	1.78	-4.50	TRP	369 B	CG	NE1	3.23
1xnb	1.49	0.17	71	TYR	348 B	CH2	CZ	3.61	
54.93	-46.17	5.85	4.79	3.36	TYR	26	CZ3	CG	3.97
1xnb	1.49	0.17	71	PHE	36	CD2	CZ	3.50	
77.15	81.70	6.16	6.01	1.37	TYR	69	NE1	OH	3.01
1xnb	1.49	0.17	71	PHE	125	CH2	CE1	3.66	
74.48	-81.56	6.27	6.04	-1.68	TYR	166	NE1	OH	3.34

1xnb	1.49	0.17	129	TYR	80	CZ3	CG	3.62
38.22	94.87	4.57	2.83	3.59				
1xnb	1.49	0.17	153	TYR	128	CE3	CD2	3.95
34.50	159.44	5.91	3.34	4.87				
1xnb	1.49	0.17	153	PHE	146	CH2	CE1	3.97
48.26	93.12	6.69	4.99	-4.45				
1xnb	1.49	0.17	164	TYR	79	CZ3	OH	3.77
63.97	110.57	7.11	6.39	3.12				
1xnb	1.49	0.17	164	PHE	146	CZ3	CE2	3.95
81.36	89.12	7.61	7.52	-1.14				
1xyl	1.80	0.18	15 A	HIS	53 A	CE3	ND1	3.35
24.50	139.76	4.55	1.89	-4.14				
1xyl	1.80	0.18	15 A	HIS	284 A	CH2	CE1	3.80
58.99	46.84	6.35	5.44	3.27				
1xyl	1.80	0.18	136 A	PHE	93 A	CE3	CE2	3.47
77.49	127.92	5.59	5.46	1.21				
1xyl	1.80	0.18	136 A	PHE	525 B	CH2	CD1	3.67
73.74	68.27	6.27	6.02	-1.75				
1xyl#	1.80	0.18	236 A	PHE	201 A	CZ3	CD1	3.69
61.77	80.08	6.62	5.83	-3.13				
1xyl	1.80	0.18	269 A	PHE	268 A	CD2	CD2	3.74
13.14	125.66	5.01	1.14	4.88				
1xyz	1.40	0.18	600 B	HIS	601 B	CD1	CD2	3.47
51.73	-59.52	5.30	4.16	3.28				
1xyz	1.40	0.18	607 B	PHE	564 B	CZ2	CE1	3.74
52.95	28.19	6.43	5.13	-3.87				
1xyz	1.40	0.18	659 B	PHE	564 B	CZ2	CE2	3.81
86.70	31.94	6.43	6.42	-0.37				
1xyz	1.40	0.18	795 B	TRP	803 B	CD1	NE1	3.73
70.32	-74.90	6.40	6.03	2.15				
1xyz	1.40	0.18	803 B	TRP	795 B	NE1	CD1	3.73
29.20	-53.75	6.40	3.12	5.59				
1xyz	1.40	0.18	803 B	HIS	723 B	CZ2	CE1	3.41
73.11	13.19	4.96	4.75	1.44				
2abk	1.85	0.19	178	PHE	171	CZ3	CD1	3.74
71.74	93.39	7.27	6.90	2.28				
2acq	1.76	0.17	20	TYR	48	CD1	CD1	3.73
63.97	-121.08	5.72	5.14	-2.51				
2acq	1.76	0.17	20	TRP	219	CH2	CH2	3.30
89.42	52.28	7.37	7.37	0.07				
2acq	1.76	0.17	219	TRP	20	CH2	CH2	3.30
60.07	74.67	7.37	6.39	3.68				
2acq	1.76	0.17	79	HIS	110	CE3	CD2	3.59
87.84	135.46	5.96	5.96	0.22				
2acq	1.76	0.17	79	TRP	111	CZ3	CD1	3.58
66.40	94.58	7.52	6.89	-3.01				
2acq	1.76	0.17	111	TRP	79	CD1	CZ3	3.58
70.31	-78.95	7.52	7.08	-2.54				
2acq	1.76	0.17	79	PHE	115	CZ2	CE2	3.90
43.63	3.83	5.97	4.12	-4.32				
2acq	1.76	0.17	79	PHE	121	CD1	CD1	3.84
33.95	-72.08	5.58	3.11	4.63				
2acq	1.76	0.17	79	PHE	122	CZ2	CE1	3.58
83.97	23.27	5.84	5.81	0.61				
2acq	1.76	0.17	111	HIS	110	NE1	ND1	3.60
62.17	-58.36	5.60	4.95	2.62				
2acq	1.76	0.17	111	PHE	311	CZ3	CE2	3.71
85.06	108.68	7.17	7.14	0.62				
2acq	1.76	0.17	295	HIS	187	CE3	NE2	3.93
72.14	144.35	6.55	6.23	2.01				

2acq	1.76	0.17	295	TYR	189	CE3	CD2	3.87
49.52	156.23	6.42	4.88	-4.17				
2ak3	1.85	0.19	203 B	HIS	28 B	CZ2	CD2	3.69
74.04	20.39	5.66	5.44	1.56				
2ak3	1.85	0.19	203 B	TYR	207 B	CZ2	CD1	3.82
48.44	13.51	5.75	4.30	-3.82				
2alp	1.70	0.13	105	TYR	238	CD1	CG	3.49
73.43	-75.26	5.62	5.39	1.60				
2ayh	1.60	0.14	18	TYR	64	CG	CE2	3.42
46.79	-164.50	5.04	3.67	-3.45				
2ayh	1.60	0.14	34	TYR	24	CD1	CE2	3.39
79.32	-81.85	5.83	5.73	-1.08				
2ayh	1.60	0.14	34	PHE	30	CZ2	CD2	3.81
72.99	27.08	6.32	6.04	1.85				
2ayh	1.60	0.14	103	TYR	94	NE1	CE2	3.54
49.62	-68.13	5.85	4.45	-3.79				
2ayh	1.60	0.14	103	TYR	123	CZ2	CD2	3.48
45.36	19.90	4.55	3.24	3.19				
2ayh	1.60	0.14	151	TYR	72	NE1	CE1	3.52
20.85	-26.31	4.93	1.74	4.61				
2ayh	1.60	0.14	151	TYR	75	CZ3	OH	3.63
85.12	158.06	6.53	6.51	-0.56				
2ayh	1.60	0.14	158	PHE	120	CD1	CD1	3.26
35.43	-113.50	4.33	2.50	3.53				
2ayh	1.60	0.14	158	HIS	166	CZ3	CD2	3.43
53.97	129.47	5.91	4.78	3.48				
2ayh	1.60	0.14	184	PHE	30	CD2	CE2	3.44
11.77	115.93	4.13	0.83	4.05				
2ayh	1.60	0.14	184	TRP	192	CH2	CZ3	3.89
65.59	54.05	8.15	7.42	-3.37				
2ayh	1.60	0.14	192	TRP	184	CZ3	CH2	3.89
62.82	89.58	8.15	7.25	3.72				
2ayh	1.60	0.14	208	PHE	7	CE3	CE1	3.77
39.49	-168.92	5.54	3.52	4.28				
2aza	1.80	0.16	48 A	TYR	15 A	CD1	OH	3.90
81.10	-89.24	6.84	6.76	1.06				
2aza	1.80	0.16	48 A	PHE	97 A	CH2	CE1	3.92
80.18	79.63	7.41	7.30	-1.26				
2aza	1.80	0.16	48 A	TYR	110 A	NE1	CG	3.18
70.55	-34.04	5.20	4.90	1.73				
2bbk	1.75	0.17	127 H	PHE	172 H	CG	CE1	3.69
39.78	-152.96	5.66	3.62	-4.35				
2bbk	1.75	0.17	127 H	TYR	220 H	CH2	CD2	3.78
25.86	80.58	5.62	2.46	-5.05				
2bbk	1.75	0.17	229 H	PHE	172 H	CZ3	CE2	3.64
54.47	100.42	6.38	5.19	-3.71				
2bbk	1.75	0.17	229 H	HIS	182 H	CH2	CG	3.79
38.30	63.96	5.51	3.42	4.32				
2bbk	1.75	0.17	264 H	PHE	299 H	CZ3	CD1	3.87
85.13	79.47	6.31	6.29	0.54				
2bbk	1.75	0.17	269 H	HIS	217 H	NE1	NE2	3.10
15.78	-68.06	4.15	1.12	-3.99				
2bbk	1.75	0.17	291 H	TYR	232 H	CZ3	CE2	3.92
60.75	115.89	7.25	6.33	3.54				
2bbk	1.75	0.17	291 H	HIS	293 H	CE3	CE1	3.87
78.54	151.91	6.41	6.28	1.27				
2bbk	1.75	0.17	26 L	TYR	25 L	NE1	OH	3.65
48.92	-82.29	5.50	4.15	-3.61				
2bbk	1.75	0.17	26 L	TYR	70 L	CH2	CD1	3.41
62.96	68.15	6.30	5.61	2.87				

2bbk	1.75	0.17		108 L	PHE	110 L	CZ3	CE2	3.63
58.60	135.06	6.28	5.36	3.27					
2cba	1.54	0.15		5	TYR	7	CZ3	CE2	3.92
84.71	121.09	6.97	6.94	-0.64					
2cba	1.54	0.15		5	TRP	16	CE3	CZ2	3.66
53.80	163.52	5.52	4.45	3.26					
2cba	1.54	0.15		16	TRP	5	CZ2	CE3	3.66
26.08	-10.23	5.52	2.42	4.96					
2cba	1.54	0.15		16	TYR	7	CH2	CE1	3.80
86.87	29.46	6.13	6.12	-0.33					
2cba	1.54	0.15		97	PHE	66	CH2	CD1	3.94
58.21	86.61	6.80	5.78	-3.58					
2cba	1.54	0.15		97	PHE	226	CE3	CD2	3.74
19.30	153.56	5.19	1.71	-4.90					
2cba	1.54	0.15		123	PHE	131	CD1	CD1	3.92
46.60	-108.01	6.53	4.74	4.49					
2cba	1.54	0.15		192	PHE	260	CE2	CZ	3.67
30.89	-21.90	5.27	2.71	-4.52					
2cba	1.54	0.15		209	HIS	107	CE3	CE1	3.60
21.23	137.81	4.90	1.77	-4.57					
2cba	1.54	0.15		209	HIS	119	CH2	NE2	3.54
84.89	82.82	5.97	5.95	-0.53					
2cba	1.54	0.15		209	TYR	194	CD1	OH	3.57
55.03	-114.38	5.67	4.64	-3.25					
2ccy	1.67	0.19		58 A	PHE	75 A	CD1	CE2	3.92
88.58	-94.32	6.67	6.67	-0.17					
2ccy	1.67	0.19		58 A	PHE	82 A	NE1	CZ	3.59
18.43	-95.20	4.94	1.57	4.68					
2ccy	1.67	0.19		58 A	TRP	86 A	CH2	CZ2	3.49
41.19	108.46	5.82	3.83	4.38					
2ccy	1.67	0.19		86 A	TRP	58 A	CZ2	CH2	3.49
73.63	-3.79	5.82	5.58	-1.64					
2ccy	1.67	0.19		86 A	PHE	82 A	NE1	CE2	3.46
66.90	-39.03	5.31	4.88	2.08					
2cpl	1.63	0.18		121	PHE	60	CZ2	CE1	3.84
81.81	15.44	6.14	6.08	0.88					
2ctc	1.40	0.16		63	PHE	192	CD1	CE2	3.33
54.04	-81.91	4.97	4.02	2.92					
2ctc	1.40	0.16		73	TYR	12	CE3	CG	3.81
79.07	167.50	5.75	5.65	1.09					
2ctc	1.40	0.16		73	PHE	116	NE1	CE1	3.62
36.40	-44.82	5.55	3.30	4.46					
2ctc	1.40	0.16		73	HIS	120	CH2	CE1	3.49
50.50	35.18	4.83	3.72	3.08					
2ctc	1.40	0.16		81	PHE	82	CE3	CD1	3.70
83.88	148.79	6.84	6.80	-0.73					
2ctc	1.40	0.16		126	PHE	118	CG	CD2	3.66
20.53	-167.26	5.04	1.78	4.72					
2ctc	1.40	0.16		147	TRP	257	CZ3	CE3	3.66
84.12	147.33	6.00	5.97	-0.61					
2ctc	1.40	0.16		257	TRP	147	CE3	CZ3	3.66
37.00	-178.37	6.00	3.61	-4.79					
2ctc	1.40	0.16		294	PHE	86	CZ3	CE1	3.24
55.78	95.37	6.42	5.31	3.61					
2ctc	1.40	0.16		294	PHE	96	CH2	CD1	3.65
78.15	44.74	6.22	6.09	1.28					
2cyp	1.70	0.20		51	HIS	52	NE1	CD2	3.24
39.85	-50.14	5.05	3.23	-3.88					
2cyp	1.70	0.20		51	PHE	158	CH2	CZ	3.97
58.50	77.43	6.90	5.88	3.61					

2cyp	1.70	0.20		101	PHE	19	CZ3	CE2	3.54
84.83	91.82	6.46	6.43	0.58					
2cyp	1.70	0.20		101	PHE	99	NE1	CD2	3.67
57.09	-9.38	5.94	4.99	-3.22					
2cyp	1.70	0.20		126	PHE	19	CZ3	CD2	3.79
86.56	96.25	6.52	6.51	-0.39					
2cyp	1.70	0.20		126	PHE	284	CE3	CD1	3.94
78.00	143.78	7.08	6.93	1.47					
2cyp	1.70	0.20		126	PHE	286	CE3	CE2	3.70
51.08	162.41	6.00	4.67	-3.77					
2cyp	1.70	0.20		191	HIS	175	CZ2	CG	3.42
41.09	30.95	4.41	2.90	3.32					
2cyp	1.70	0.20		223	TYR	236	NE1	CG	3.51
7.16	7.93	3.67	0.45	-3.64					
2dnj	2.00	0.17		181 A	TRP	194 A	CZ3	CH2	3.76
56.68	104.61	7.25	6.06	-3.98					
2dnj	2.00	0.17		194 A	TRP	181 A	CH2	CZ3	3.76
40.35	91.19	7.25	4.69	-5.53					
2end#	1.45	0.16		128	HIS	107	CD2	CD2	3.18
7.63	-170.12	4.29	0.58	4.25					
2er7	1.60	0.14		39 E	TYR	75 E	NE1	OH	2.83
82.01	-26.55	6.62	6.56	0.92					
2er7	1.60	0.14		39 E	TRP	71 E	CG	CH2	3.61
45.47	-140.66	6.50	4.64	4.56					
2er7	1.60	0.14		71 E	TRP	39 E	CH2	CG	3.61
70.15	75.49	6.50	6.11	2.21					
2er7	1.60	0.14		39 E	PHE	111 E	CZ2	CZ	3.71
87.36	31.82	5.43	5.42	0.25					
2er7	1.60	0.14		190 E	PHE	125 E	CD2	CE1	3.82
13.73	14.48	4.60	1.07	4.47					
2er7	1.60	0.14		190 E	PHE	314 E	CD2	CD2	3.61
6.89	-130.52	4.97	0.61	-4.93					
2er7	1.60	0.14		192 E	PHE	259 E	CH2	CD1	3.91
84.21	50.99	7.01	6.97	0.71					
2er7	1.60	0.14		192 E	PHE	302 E	CZ2	CE2	3.55
60.44	14.37	4.72	4.11	-2.33					
2er7	1.60	0.14		192 E	PHE	323 E	CZ2	CZ	3.79
28.61	51.47	5.28	2.54	4.63					
2er7	1.60	0.14		232 E	TYR	231 E	CD2	CD1	3.65
5.73	87.43	4.86	0.51	4.83					
2er7	1.60	0.14		232 E	TYR	246 E	CD1	CG	3.36
51.36	-94.54	4.85	3.79	-3.03					
2er7	1.60	0.14		232 E	PHE	248 E	CZ3	CD1	3.61
72.53	92.03	6.55	6.25	-1.97					
2fal	1.80	0.15		14	PHE	18	CZ2	CD1	3.43
57.33	-14.26	5.16	4.34	-2.79					
2fal	1.80	0.15		14	PHE	68	CH2	CD2	3.83
69.79	68.46	5.81	5.45	-2.01					
2fal	1.80	0.15		14	PHE	112	CZ3	CZ	3.86
68.28	128.20	7.25	6.73	2.68					
2fal	1.80	0.15		14	TRP	130	CE3	CH2	4.00
52.89	172.17	6.74	5.38	4.06					
2fal	1.80	0.15		130	TRP	14	CH2	CE3	4.00
76.90	49.84	6.74	6.56	-1.53					
2fal	1.80	0.15		130	PHE	112	CE3	CE2	3.61
22.92	132.82	4.90	1.90	-4.52					
2gbp	1.90	0.15		183	TYR	10	CD1	OH	3.91
58.94	-74.35	7.62	6.53	-3.93					
2gbp	1.90	0.15		183	HIS	152	NE1	CD2	3.36
74.89	4.39	4.71	4.55	-1.23					

2gdm	1.70	0.16	15	TYR	72	CD2	CD2	3.63
33.01	175.44	4.47	2.44	-3.75				
2gst	1.80	0.16	7 A	TRP	45 A	CZ2	CD1	3.97
37.05	29.57	6.49	3.91	5.18				
2gst	1.80	0.16	45 A	TRP	7 A	CD1	CZ2	3.97
42.12	-109.50	6.49	4.35	4.82				
2gst	1.80	0.16	45 A	TYR	32 A	CE3	CZ	3.86
66.32	166.30	5.53	5.06	2.22				
2gst	1.80	0.16	45 A	TYR	61 A	CZ3	CE2	3.67
65.25	100.27	7.05	6.40	2.95				
2gst	1.80	0.16	146 A	PHE	183 A	CE3	CE1	3.61
68.07	137.35	6.52	6.05	2.44				
2gst	1.80	0.16	214 A	PHE	169 A	CE3	CD1	3.77
7.94	88.96	4.95	0.72	-4.90				
2gst	1.80	0.16	214 A	PHE	208 A	CE3	CE2	3.98
77.20	145.62	7.00	6.83	1.55				
2hbg	1.50	0.13	14	TRP	130	CE3	CH2	3.95
75.12	165.05	7.31	7.06	1.88				
2hbg	1.50	0.13	130	TRP	14	CH2	CE3	3.95
77.42	43.90	7.31	7.13	-1.59				
2hft	1.69	0.20	14	PHE	100	CD2	CG	3.90
24.92	176.75	4.75	2.00	-4.31				
2hft	1.69	0.20	25	TYR	34	NE1	CE2	3.37
77.07	-30.66	4.86	4.74	-1.09				
2hmz	1.66	0.18	10 A	PHE	29 A	CD1	CE1	3.64
75.08	-111.06	6.35	6.14	1.63				
2hmz	1.66	0.18	97 A	PHE	55 A	CH2	CZ	3.57
68.27	75.26	7.08	6.58	-2.62				
2hmz	1.66	0.18	97 A	PHE	80 A	CD2	CD1	3.41
30.12	147.46	4.16	2.08	-3.60				
2hmz	1.66	0.18	97 A	TYR	93 A	CD1	CZ	3.96
66.71	-113.75	6.47	5.94	-2.56				
2hmz	1.66	0.18	97 A	HIS	101 A	CH2	CG	3.40
56.27	60.74	5.38	4.48	2.98				
2hpd#	2.00	0.17	90 A	PHE	11 B	NE1	CD1	3.28
89.40	-28.67	5.10	5.10	0.05				
2hpd	2.00	0.17	96 A	HIS	100 A	CZ2	ND1	3.41
54.96	55.81	5.24	4.29	3.01				
2hpd	2.00	0.17	130 A	PHE	421 A	CE3	CE2	3.98
85.56	173.30	5.83	5.81	0.45				
2hpd	2.00	0.17	130 A	PHE	423 A	CZ3	CE2	3.94
61.21	120.63	7.11	6.23	3.43				
2hpd	2.00	0.17	325 A	PHE	275 A	CD1	CE1	3.75
51.32	-120.84	5.34	4.17	-3.34				
2hpd	2.00	0.17	325 A	PHE	279 A	CD1	CZ	3.83
83.80	-72.56	7.09	7.05	0.77				
2hts#	1.83	0.19	203	HIS	259	CH2	CD2	3.91
83.15	95.14	6.97	6.92	-0.83				
2hts	1.83	0.19	217	TRP	258	CZ3	CH2	3.93
62.71	86.14	8.09	7.19	3.71				
2hts	1.83	0.19	258	TRP	217	CH2	CZ3	3.93
58.30	58.85	8.09	6.88	-4.25				
2hts#	1.83	0.19	217	HIS	259	CH2	ND1	3.62
23.64	36.03	4.02	1.60	-3.69				
2hts	1.83	0.19	277	PHE	246	CD2	CE2	3.58
7.45	-150.01	4.88	0.62	-4.84				
2hts	1.83	0.19	277	PHE	279	CE3	CZ	3.46
82.22	167.46	6.02	5.96	-0.81				
2kau	2.00	0.19	79 C	HIS	78 C	CD1	NE2	3.99
81.24	-101.71	6.11	6.04	-0.93				

2kau#	2.00	0.19		137 C	PHE	477 C	CE3	CZ	3.82
52.05	179.20	6.00	4.73	-3.69					
2kau	2.00	0.19		175 C	TYR	176 C	CZ3	CE2	3.86
89.82	139.90	5.89	5.89	-0.02					
2kau#	2.00	0.19		175 C	HIS	486 C	NE1	NE2	3.89
54.99	-10.16	5.74	4.70	3.29					
2kau#	2.00	0.19		222 C	PHE	45 C	CE2	CE1	3.67
24.43	68.37	4.91	2.02	4.47					
2kau	2.00	0.19		375 C	TYR	407 C	NE1	OH	2.99
53.59	-22.18	5.94	4.78	3.53					
2kau	2.00	0.19		375 C	PHE	440 C	NE1	CE2	3.46
27.69	-29.78	5.12	2.37	-4.54					
2kau	2.00	0.19		435 C	PHE	440 C	CG	CD1	3.95
13.16	-96.57	5.04	1.16	-4.91					
2mnr	1.90	0.16		219	TYR	137	CH2	CE2	3.72
60.56	42.37	6.06	5.28	2.98					
2mnr	1.90	0.16		219	TRP	315	CZ3	CD1	3.79
61.30	103.63	7.35	6.45	3.53					
2mnr	1.90	0.16		315	TRP	219	CD1	CZ3	3.79
41.00	-81.95	7.35	4.82	-5.55					
2mnr	1.90	0.16		249	TYR	229	CD1	OH	3.89
73.57	-43.08	6.43	6.17	1.82					
2mnr	1.90	0.16		315	HIS	314	CD1	CD2	3.41
73.72	-69.90	5.27	5.06	1.48					
2mnr	1.90	0.16		349	HIS	48	NE1	ND1	3.72
42.36	-37.34	4.63	3.12	3.42					
2nac	1.80	0.15		237 B	TYR	219 B	CE3	CZ	3.43
87.76	154.20	5.50	5.50	0.22					
2nac	1.80	0.15		310 B	HIS	319 B	CE3	ND1	3.72
88.32	150.20	5.56	5.56	-0.16					
2nac	1.80	0.15		321 B	TYR	305 B	CH2	OH	3.41
20.56	86.30	5.06	1.77	4.74					
2nac	1.80	0.15		321 B	HIS	319 B	CG	CE1	3.41
30.39	-151.65	5.01	2.54	-4.32					
2olb	1.40	0.18		64 A	PHE	74 A	CG	CE1	3.88
45.89	-144.29	5.80	4.16	-4.04					
2olb	1.40	0.18		72 A	PHE	155 A	CG	CD2	3.62
22.90	-93.12	5.00	1.94	-4.61					
2olb	1.40	0.18		72 A	TYR	156 A	CH2	CD1	3.96
63.10	92.53	7.21	6.43	-3.26					
2olb	1.40	0.18		97 A	PHE	93 A	NE1	CD1	3.84
50.10	-31.12	6.34	4.86	-4.07					
2olb	1.40	0.18		97 A	PHE	144 A	CH2	CE2	3.92
11.80	97.84	5.16	1.08	-5.05					
2olb	1.40	0.18		179 A	PHE	44 A	CH2	CZ	3.49
80.59	75.91	6.88	6.79	-1.12					
2olb	1.40	0.18		211 A	TYR	210 A	NE1	CE1	3.41
33.35	-73.12	4.89	2.69	-4.08					
2olb	1.40	0.18		332 A	PHE	333 A	CZ3	CD1	3.72
45.21	103.64	6.54	4.64	-4.61					
2olb	1.40	0.18		382 A	PHE	361 A	CZ3	CZ	3.83
50.27	126.93	6.58	5.06	4.20					
2olb	1.40	0.18		397 A	PHE	400 A	CE3	CD2	3.91
60.64	136.36	6.67	5.81	-3.27					
2olb	1.40	0.18		416 A	TYR	109 A	CH2	OH	3.13
77.49	59.68	7.13	6.96	-1.54					
2olb	1.40	0.18		416 A	TYR	317 A	CG	OH	3.57
46.42	-115.60	6.44	4.67	4.44					
2olb	1.40	0.18		416 A	PHE	426 A	CD1	CD2	3.51
88.42	-84.33	5.58	5.58	0.15					

2olb	1.40	0.18	494	A	HIS	517	A	CG	ND1	3.44
17.61	-177.79	3.62	1.09	-3.45						
2pgd	2.00	0.20	149		TRP	172		CE3	CE3	3.55
35.03	-152.58	5.21	2.99	4.26						
2pgd	2.00	0.20	172		TRP	149		CE3	CE3	3.55
66.51	176.42	5.21	4.78	2.08						
2pgd	2.00	0.20	350		PHE	181		CZ3	CG	3.88
84.20	93.70	6.78	6.75	-0.69						
2pgd	2.00	0.20	361		PHE	338		CZ3	CE1	3.96
63.25	111.67	6.77	6.05	-3.05						
2pgd	2.00	0.20	405		TYR	429		CZ3	CG	3.98
74.63	100.50	6.75	6.51	1.79						
2phy	1.40	0.19	119		PHE	121		CG	CZ	3.42
43.76	-162.99	5.17	3.58	-3.73						
2pia	2.00	0.19	164		PHE	141		CZ3	CZ	3.88
84.08	123.42	7.44	7.40	0.77						
2pia	2.00	0.19	184		HIS	214		CZ3	CE1	3.60
81.34	92.34	6.65	6.57	-1.00						
2pia	2.00	0.19	215		PHE	187		CZ2	CD1	3.90
35.21	3.29	5.88	3.39	-4.80						
2pia	2.00	0.19	215		HIS	214		CD1	CE1	3.52
77.52	-73.44	5.64	5.51	1.22						
2prk	1.50	0.17	212		HIS	72		CE3	CD2	3.85
72.32	140.11	6.63	6.56	0.96						
2rn2	1.48	0.20	81		TYR	73		CZ3	CE1	3.63
87.08	104.12	6.81	6.80	-0.35						
2rn2	1.48	0.20	81		TRP	85		CE3	CH2	3.41
56.50	-162.84	5.66	4.72	3.12						
2rn2	1.48	0.20	85		TRP	81		CH2	CE3	3.41
46.06	42.22	5.66	4.08	-3.92						
2rn2	1.48	0.20	85		TRP	90		CE3	CZ3	3.68
60.47	-173.99	6.66	5.80	3.28						
2rn2	1.48	0.20	90		TRP	85		CZ3	CE3	3.68
66.04	140.49	6.66	6.09	2.70						
2rn2	1.48	0.20	85		TRP	104		CZ3	CD1	3.62
87.43	105.01	6.21	6.20	0.28						
2rn2	1.48	0.20	104		TRP	85		CD1	CZ3	3.62
20.43	-82.49	6.21	2.16	-5.82						
2rn2	1.48	0.20	104		TYR	73		NE1	OH	3.07
72.82	2.33	5.37	5.13	1.59						
2rn2	1.48	0.20	118		TRP	120		CE3	CH2	3.36
54.16	-169.65	5.74	4.65	-3.36						
2rn2	1.48	0.20	120		TRP	118		CH2	CE3	3.36
56.38	59.68	5.74	4.78	3.18						
2scp	2.00	0.18	170	A	TYR	11	A	CZ2	CE1	3.35
21.31	-14.41	4.50	1.64	-4.19						
2scp	2.00	0.18	170	A	PHE	99	A	CH2	CE2	3.62
81.14	66.99	6.14	6.07	-0.95						
2scp	2.00	0.18	170	A	PHE	119	A	CZ3	CZ	3.62
60.89	114.28	6.90	6.03	3.36						
2scp	2.00	0.18	170	A	PHE	157	A	CD1	CE2	3.40
88.06	-76.54	5.57	5.57	-0.19						
2scp	2.00	0.18	170	A	PHE	169	A	NE1	CD2	3.41
26.78	-26.29	5.10	2.30	4.55						
2sil	1.60	0.17	80		PHE	368		CZ2	CG	3.78
86.64	28.30	6.00	5.99	0.35						
2sil	1.60	0.17	121		TRP	128		CZ2	CE3	3.89
68.84	42.28	6.61	6.16	2.38						
2sil	1.60	0.17	128		TRP	121		CE3	CZ2	3.89
58.85	-173.64	6.61	5.66	3.42						

2sil	1.60	0.17	121	TRP	140	CZ2	CD1	3.74
33.78	8.72	6.47	3.60	-5.38				
2sil	1.60	0.17	140	TRP	121	CD1	CZ2	3.74
86.86	-99.48	6.47	6.46	0.35				
2sil	1.60	0.17	140	HIS	162	CZ3	CD2	3.76
66.03	119.58	6.37	5.82	2.59				
2sil	1.60	0.17	218	PHE	191	CH2	CD2	3.45
52.56	74.87	6.36	5.05	-3.87				
2sil	1.60	0.17	263	PHE	209	CZ2	CE2	3.93
71.96	29.90	6.73	6.40	2.08				
2tgi#	1.80	0.17	32	HIS	58	NE1	NE2	2.64
41.03	-21.08	3.62	2.37	-2.73				
2tgi	1.80	0.17	52	TYR	6	CH2	OH	3.43
69.73	37.30	7.09	6.65	-2.46				
3bcl#	1.90	0.19	173	TYR	118	CG	OH	3.92
57.28	177.45	6.66	5.60	3.60				
3bcl#	1.90	0.19	173	PHE	139	CH2	CE1	3.88
53.88	75.80	6.93	5.60	4.08				
3bcl	1.90	0.19	178	PHE	179	CE2	CZ	3.45
21.82	-94.77	4.09	1.52	3.80				
3bcl	1.90	0.19	178	PHE	217	CH2	CG	3.91
88.38	56.26	6.01	6.01	-0.17				
3bcl	1.90	0.19	215	PHE	217	CZ3	CD1	3.60
65.23	108.20	6.98	6.34	-2.92				
3bcl	1.90	0.19	332	HIS	352	NE1	CE1	3.59
30.27	-34.18	5.26	2.65	4.54				
3bcl	1.90	0.19	340	TYR	337	CE3	CE1	3.83
68.16	149.52	6.34	5.89	-2.36				
3bcl	1.90	0.19	340	HIS	352	CZ3	CD2	3.84
19.16	95.23	5.14	1.68	-4.86				
3cla	1.75	0.16	16	PHE	11	CD2	CE2	3.39
5.01	-111.97	4.65	0.41	-4.63				
3cla	1.75	0.16	16	PHE	22	CZ3	CG	3.62
85.74	106.28	6.67	6.65	0.50				
3cla	1.75	0.16	86	TYR	8	CZ3	CD2	3.77
50.64	117.81	6.46	4.99	4.10				
3cla	1.75	0.16	152	PHE	96	CZ2	CG	3.60
64.40	15.60	5.10	4.60	2.20				
3cox	1.80	0.16	46	PHE	56	CE3	CE1	3.61
48.88	174.28	5.23	3.94	-3.44				
3cox	1.80	0.16	164	TYR	171	CZ3	CE2	3.94
64.52	104.54	7.27	6.56	3.13				
3cox	1.80	0.16	318	PHE	388	CD1	CE2	3.40
38.89	-104.21	5.49	3.45	-4.27				
3cox	1.80	0.16	318	TRP	401	CZ3	CD2	3.99
86.45	114.89	6.77	6.76	0.42				
3cox	1.80	0.16	401	TRP	318	CD2	CZ3	3.99
38.73	176.63	6.77	4.24	5.28				
3cox	1.80	0.16	333	HIS	331	CZ2	CD2	3.56
47.41	46.81	5.55	4.08	3.76				
3cox	1.80	0.16	351	TYR	130	CZ2	CE2	3.64
77.99	-9.42	4.89	4.78	1.02				
3cox	1.80	0.16	440	TYR	428	CZ2	OH	3.64
50.27	27.92	5.43	4.17	3.47				
3dfr	1.70	0.15	5	TRP	133	CH2	CH2	3.61
89.99	39.22	7.60	7.60	0.00				
3dfr	1.70	0.15	21	PHE	122	CE3	CD1	3.59
73.72	165.03	5.48	5.26	-1.54				
3dfr	1.70	0.15	133	TRP	5	CH2	CH2	3.61
74.79	46.26	7.60	7.33	1.99				

3dfr	1.70	0.15	133	TRP	158	CZ3	CE2	3.55		
77.57	131.38	5.59	5.46	-1.20	TRP	133	CE2	CZ3	3.55	
3dfr	1.70	0.15	158	TYR	137	CZ2	CG	3.91		
19.23	75.04	5.59	1.84	-5.28	HIS	200	CE3	NE2	3.47	
3est	1.65	0.17	27	HIS	40	CE3	CE1	3.82		
72.68	35.91	6.28	6.00	-1.87	PHE	215	CH2	CE2	3.89	
3est	1.65	0.17	27	HIS	91	CD2	CD2	3.80		
50.83	151.04	4.52	3.50	-2.86	PHE	372	CD2	CD1	3.84	
3est	1.65	0.17	141	HIS	374	NE1	CE1	3.57		
58.84	178.99	5.64	4.83	2.92	PHE	94	CE3	CE2	3.71	
3est	1.65	0.17	172	PHE	200	4	CE2	CG	3.46	
27.40	67.98	5.76	2.65	-5.11	HIS	306	1	CD1	CG	3.85
3est	1.65	0.17	237	PHE	120	CE3	CE2	3.70		
8.73	-49.61	4.60	0.68	-4.55	TYR	296	CZ2	CD1	3.39	
3grs	1.54	0.19	70	PHE	218	L	CZ3	CD2	3.93	
9.80	138.63	5.09	0.89	-5.01	PHE	256	L	CZ3	CD2	3.84
3grs	1.54	0.19	70	TRP	462	L	CG	CD1	3.30	
57.30	-50.43	5.17	4.35	2.79	TRP	385	L	CD1	CG	3.30
3grs	1.54	0.19	96	TRP	135	B	CE3	CH2	3.62	
61.51	176.21	6.22	5.47	-2.97	TRP	22	B	CH2	CE3	3.62
3pga	2.00	0.17	186	1	TYR	91	E	CH2	CD2	3.90
30.70	29.61	3.88	1.98	-3.34	HIS	238	E	CZ2	NE2	3.34
3pga	2.00	0.17	335	1	TYR	11	NE1	OH	2.75	
73.07	-77.10	5.98	5.72	1.74	TYR	55	CD1	CG	3.75	
3pte	1.60	0.15	233	TYR	144	CD1	CG	4.00		
7.01	132.78	4.88	0.57	4.85	PHE	228	B	CZ2	CD1	3.88
3pte	1.60	0.15	271	PHE	380	B	CG	CD2	3.99	
65.30	21.60	6.05	5.50	-2.53	PHE	17	CD1	CE2	3.77	
3rub	2.00	0.19	214	L	HIS	259	CZ2	ND1	3.64	
82.24	151.48	5.50	5.45	0.74	PHE	7	CD1	CD1	3.60	
3rub	2.00	0.19	214	L						
64.69	99.22	7.09	6.41	-3.03						
3rub	2.00	0.19	385	L						
46.13	-163.34	4.66	3.36	-3.23						
3rub	2.00	0.19	462	L						
42.94	-102.59	4.66	3.17	-3.41						
3sdh	1.40	0.16	22	B						
65.36	160.49	6.99	6.35	2.91						
3sdh	1.40	0.16	135	B						
81.43	47.37	6.99	6.91	-1.04						
3sic	1.80	0.18	113	E						
83.97	70.46	7.43	7.39	0.78						
3sic	1.80	0.18	241	E						
28.57	9.58	3.76	1.79	-3.31						
4enl	1.90	0.15	56							
66.60	-40.80	6.10	5.60	2.42						
4gcr	1.47	0.18	68							
83.67	-69.99	5.92	5.88	-0.65						
4gcr	1.47	0.18	157							
88.61	-78.19	6.50	6.50	-0.16						
5rub	1.70	0.18	184	B						
71.26	44.68	6.80	6.44	-2.18						
5rub	1.70	0.18	417	B						
21.78	169.40	5.33	1.99	-4.95						
8abp	1.49	0.17	16							
88.45	-83.26	5.70	5.70	0.15						
8abp	1.49	0.17	16							
78.24	24.58	5.75	5.63	-1.17						
8abp	1.49	0.17	21							
78.25	-73.01	6.39	6.26	-1.30						

8abp	1.49	0.17	130	HIS	198	CD2	CG	3.70
21.47	-118.28	4.37	1.59	4.07				
8abp	1.49	0.17	199	PHE	221	NE1	CZ	3.38
86.48	-64.36	5.76	5.75	-0.35				
8acn	2.00	0.16	139	PHE	10	CZ2	CE2	3.73
65.36	32.47	5.87	5.34	2.45				
8acn	2.00	0.16	139	HIS	98	CD1	CE1	3.43
47.26	-90.62	5.21	3.83	-3.54				
8acn	2.00	0.16	139	TYR	154	NE1	OH	3.07
67.91	-6.53	5.93	5.50	-2.23				
8acn	2.00	0.16	139	TYR	517	NE1	CD1	3.30
27.78	-11.95	4.78	2.22	4.23				
8acn	2.00	0.16	346	HIS	332	CH2	CE1	3.69
77.12	92.88	6.59	6.42	1.47				
8acn	2.00	0.16	346	HIS	460	CD1	CD2	3.70
58.63	-98.40	6.35	5.42	3.31				
8acn	2.00	0.16	346	PHE	462	CE3	CZ	3.63
41.35	159.80	5.72	3.78	4.29				
8acn	2.00	0.16	547	PHE	739	CD1	CZ	3.74
22.46	-123.98	4.91	1.87	-4.54				
8acn	2.00	0.16	576	PHE	579	CE3	CD2	3.89
58.62	146.95	6.70	5.72	3.49				
8acn	2.00	0.16	630	TYR	623	CD1	CD2	4.00
50.81	-97.65	6.70	5.19	4.24				
8fab	1.80	0.17	185 C	TYR	191 C	CE3	CD1	3.69
82.00	149.96	6.27	6.21	-0.87				
8fab	1.80	0.17	47 D	HIS	35 D	NE1	ND1	3.03
75.39	-37.89	5.70	5.52	-1.44				
8fab	1.80	0.17	47 D	PHE	106 D	CZ2	CE2	3.98
37.50	11.87	5.61	3.41	-4.45				
8fab	1.80	0.17	47 D	PHE	108 D	CD1	CZ	3.80
67.63	-82.74	6.78	6.27	-2.58				
8fab	1.80	0.17	52 D	TYR	53 D	CD1	CE1	3.85
64.13	-84.86	6.63	5.97	-2.89				
8fab	1.80	0.17	52 D	TYR	59 D	CZ3	CE2	3.90
56.47	116.77	6.61	5.51	3.65				
8fab	1.80	0.17	111 D	TYR	35 C	NE1	CE2	3.81
76.26	4.93	5.51	5.35	-1.31				
8fab	1.80	0.17	111 D	TYR	95 D	CE3	CG	3.94
75.01	148.35	6.22	6.01	-1.61				
8fab	1.80	0.17	111 D	PHE	108 D	CZ2	CG	3.95
45.26	33.27	6.00	4.26	4.22				
8tln	1.60	0.17	55 E	PHE	62 E	NE1	CE2	3.93
57.44	-55.18	6.56	5.53	3.53				
8tln	1.60	0.17	55 E	TYR	66 E	CH2	CE2	3.54
68.89	61.67	6.54	6.10	-2.36				
8tln	1.60	0.17	55 E	HIS	105 E	CZ3	CD2	3.99
89.56	85.58	7.26	7.26	0.06				
8tln	1.60	0.17	186 E	PHE	178 E	NE1	CZ	3.62
31.04	-19.28	5.26	2.72	4.50				
8tln	1.60	0.17	186 E	TYR	242 E	CD2	CD1	3.83
13.59	-142.77	5.23	1.24	-5.08				
9rnt	1.50	0.14	59	TYR	68	CH2	CD2	3.88
35.26	75.71	6.27	3.61	-5.12				

#, the symmetry-related-residues is indicated by this sign

Table S2. Number of cases with various values of sequence-difference (absolute value) between TRP and the partner residue.

Residue	Sequence Difference													
	1	2	3	4	5	6	7	8	9	10	11	12	13	14
PHE(302) 213	9	12	13	18	5	3	9	8	0	5	1	2	3	1
TYR(222) 163	8	6	8	5	6	1	5	2	5	3	1	1	7	1
TRP(124) 80	2	4	2	8	2	0	2	6	2	0	4	4	6	2
HIS(93) 63	9	6	2	5	0	0	1	1	1	1	1	2	0	1

In parenthesis the number indicates the total number of interacting pairs. In case of TRP there are 62 residues giving rise to 124 interacting pairs.

Table S3. TYR residues with the hydroxyl group close to TRP ring.

PDB	Residue Number		Closest	Distance
file	TRP	TYR	TRP atom	(A)
(A) OH...pi (O atom on the face of the Trp ring)				
153l	107	48	CH2	3.71
laoz	8 A	103 A	CH2	3.57
ldsb	126 A	122 A	CE3	3.43
lgof	290	272	NE1	3.50
lgof	290	329	CD1	3.38
lhle	160 A	119 A	CG	3.40
lhle	194 A	244 A	CH2	3.43
lmpp	190	125	CZ3	3.49
lpbp	168	193	CZ2	3.31
lsat	153	65	CH2	3.74
lsat	153	135	CG	3.93
lsat	217	190	NE1	3.72
lwht	308 B	147 A	NE1	3.52
lwht	308 B	151 A	CH2	3.59
2ayh	18	64	CG	3.87
2cba	209	194	CD1	3.57
2nac	237 B	219 B	CE3	3.67
2nac	321 B	305 B	CH2	3.41
2o1b	211 A	210 A	NE1	3.60
2o1b	416 A	109 A	CH2	3.13
2o1b	416 A	317 A	CG	3.57
3bcl#	173	118	CG	3.92
(B) O atom on the edge of Trp ring				
lade	309 A	372 A	CD1	3.46
lade	365 A	347 A	CZ3	3.91
lamp	233	148	NE1	2.91

1aoz	277 A	186 A	NE1	3.09
1arb	182	227	CZ2	3.51
1arb	247	51	CE3	3.80
1byb	55	72	CD1	3.36
1cel	192 B	51 B	CZ2	3.57
1cel	216 B	274 B	CZ2	3.37
1cel	263 B	321 B	CH2	3.46
1cfb	634	647	NE1	3.90
1clc	220	316	CZ2	3.60
1csh	109	172	CH2	3.56
1ede	59	18	CZ3	3.44
1gpb	189	52	NE1	3.67
1gpb	215	297	CZ3	3.77
1iae	10	194	CD1	3.52
1isc	77 A	9 A	CE3	3.75
1knb	478	491	NE1	3.01
1lcp	207 A	345 A	CZ2	3.44
1lis	62	133	CZ2	3.73
1mml#	185	271	CZ3	3.36
1mpp	39	75	NE1	2.68
1mrj	192	164	CD1	3.76
1nar	58	116	NE1	2.99
1nar	101	131	CZ2	3.70
1nba	111 A	96 A	CE3	3.34
1nfp	96	152	CZ2	3.83
1ora	278	271	CE3	3.52
1oyc	116	196	CZ2	3.74
1pbe	344	181	CD1	3.68
1pii	356	278	CZ2	3.93
1rcf	120	125	NE1	2.95
1rec	31	86	NE1	2.98
1sat	46	246	CD1	3.39
1sbp	141	137	CZ3	3.66
1scs	40	67	CZ2	3.54
1scs#	182	12	CZ2	3.32
1scs#	182	100	NE1	2.98
1sri	92 A	43 A	CZ2	3.44
1tca	113	300	NE1	3.00
1ton#	237	172	CD1	3.70
1tys	98	181	CZ2	3.59
1wht#	193 A	348 B	CZ3	3.68
1xnb	71	69	NE1	3.01
1xnb	71	166	NE1	3.34
1xnb	164	79	CZ3	3.77
2ayh	151	75	CZ3	3.63
2aza	48 A	15 A	CD1	3.90
2bbk	26 L	25 L	NE1	3.65
2er7	39 E	75 E	NE1	2.83
2gbp	183	10	CD1	3.91
2kau	375 C	407 C	NE1	2.99
2mnr	219	137	CZ2	3.82
2mnr	249	229	CD1	3.89
2rn2	104	73	NE1	3.07
2tgi	52	6	CH2	3.43
3cox	351	130	CZ2	3.92
3cox	440	428	CZ2	3.64
4enl	56	11	NE1	2.75
8acn	139	154	NE1	3.07

(C) Hydrogen bonding (O atom on the edge
as in (B) + at <3.5A from NE1)

lamp	233	148	NE1	2.91
laoz	277 A	186 A	NE1	3.09
1knb	478	491	NE1	3.01
1mpp	39	75	NE1	2.68
1nar	58	116	NE1	2.99
1rcf	120	125	NE1	2.95
1rec	31	86	NE1	2.98
1scs#	182	100	NE1	2.98
1tca	113	300	NE1	3.00
1xnb	71	69	NE1	3.01
1xnb	71	166	NE1	3.34
2er7	39 E	75 E	NE1	2.83
2kau	375 C	407 C	NE1	2.99
2rn2	104	73	NE1	3.07
4en1	56	11	NE1	2.75
8acn	139	154	NE1	3.07

Interaction at crystallographic-symmetry-related position