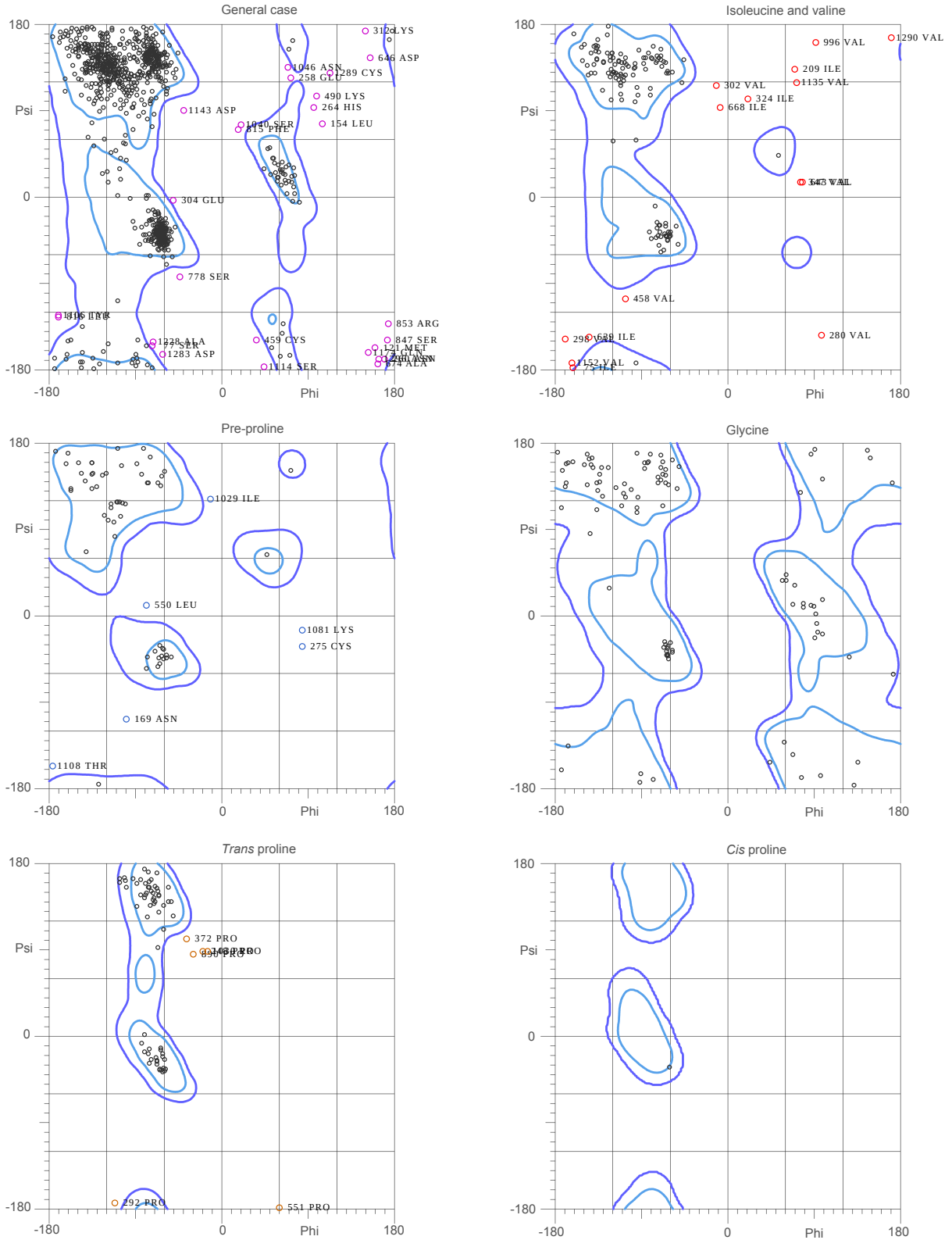
MolProbity Ramachandran analysis

6abu.B99990013.pdb, model 1



89.9% (1171/1302) of all residues were in favored (98%) regions.
95.9% (1248/1302) of all residues were in allowed (>99.8%) regions.

There were 54 outliers (pki, pri):
75 ILE (-162.8, -178.3)
77 SER (-74.9, -151.1)
121 MET (160.1, -157.8)
154 LEU (105.4, 77.4)
169 ASN (-160.9, -148.4)
209 ILE (70.1, 134.2)
248 PRO (-20.5, 89.7)
258 GLU (72.4, 125.6)
264 ASN (170.1, -170.4)
264 HIS (56.1, 94.7)
275 CYS (84.1, -32.3)
280 VAL (98.8, -164.7)
292 PRO (-112.5, -174.4)
298 VAL (-170.7, -148.4)
302 VAL (-12.4, 117.9)
304 GLU (-51.6, -3.6)
312 LYS (151.0, 178.7)
324 ILE (21.2, 103.2)
347 VAL (76.8, 56.9)
372 PRO (-57.0, 102.4)
458 VAL (-117.6, -106.3)
459 CYS (16.6, -149.8)
490 LYS (99.1, 106.7)
550 LEU (-79.5, 11.1)
551 PRO (65.2, -179.9)
639 ILE (-165.7, -146.6)
643 VAL (78.2, 16.1)
646 ASP (133.9, 147.0)
668 ILE (-8.3, 94.4)
674 ALA (163.4, -174.3)
778 SER (-64.9, -81.1)
815 PHE (17.4, 71.2)
816 LEU (-171.9, -125.4)
847 SER (173.9, -149.2)
853 ARG (174.2, -132.5)
890 PRO (-30.3, 86.1)
996 VAL (92.0, 162.3)
1029 ILE (-12.9, -132.9)
1030 PRO (-15.5, 89.8)
1046 ASN (69.6, 136.5)
1081 LYS (66.6, -15.5)
1106 TYR (-171.8, -123.8)
1108 THR (-177.8, -157.4)
1114 SER (44.8, -177.7)
1135 VAL (72.1, 125.8)
1143 ASP (-69.6, 91.6)
1152 VAL (-161.8, -175.3)
1174 GLN (153.5, -162.2)
1228 ALA (-72.1, -151.5)
1283 ASP (-62.3, -164.2)
1289 CYS (113.4, 130.6)
1290 VAL (171.0, 168.8)
1294 ASN (164.1, -169.6)