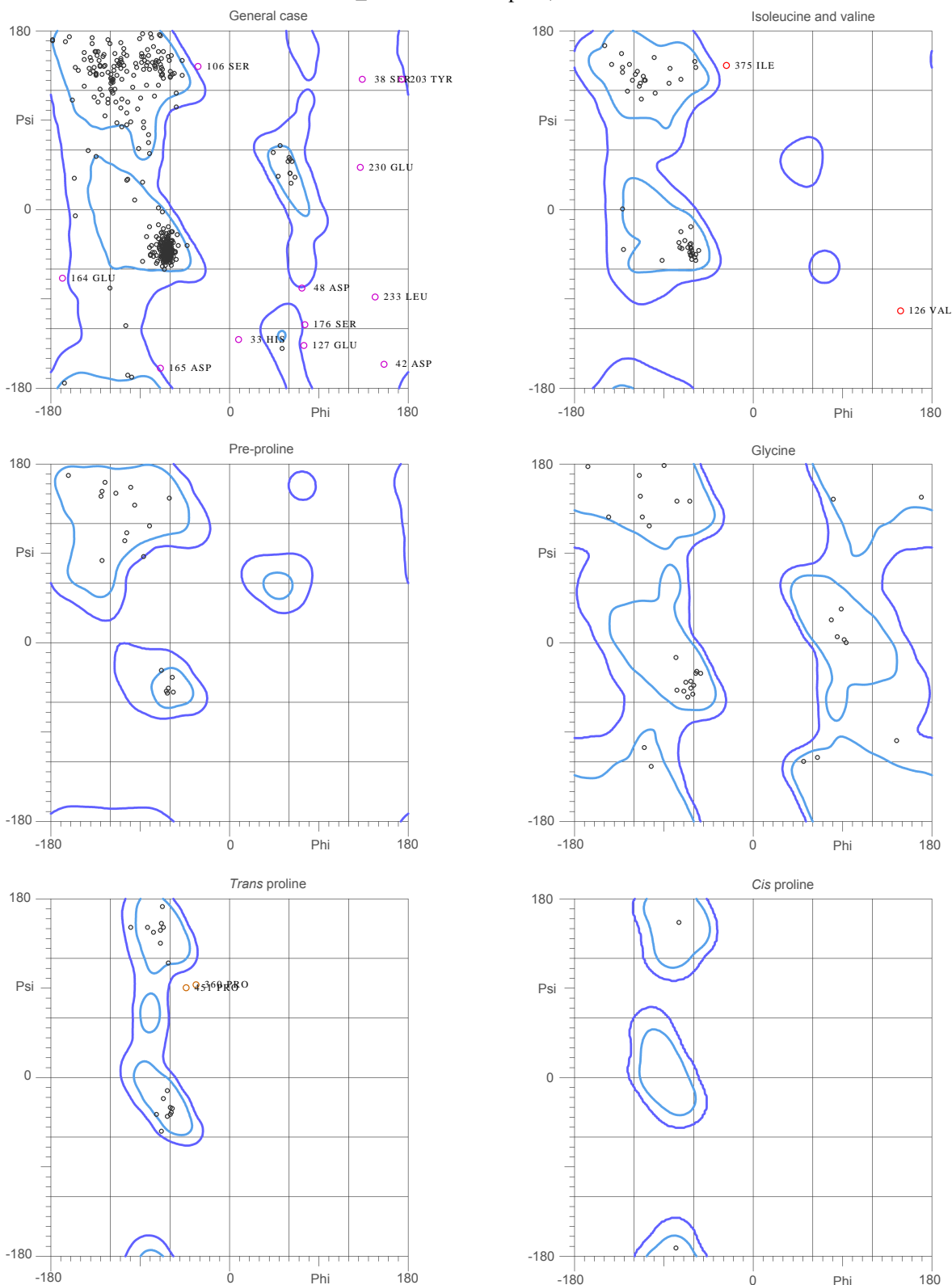# MolProbity Ramachandran analysis

1ac5_.B99990002.pdb, model 1

91.9% (442/481) of all residues were in favored (98%) regions.
96.7% (465/481) of all residues were in allowed (>99.8%) regions.

There were 16 outliers (phi, psi):
33 HIS (9.6, -131.7)
38 SER (134.6, 132.7)
42 ASP (156.6, -156.1)
48 ASP (73.3, -79.4)
106 SER (-32.1, 145.1)
126 VAL (149.2, -102.6)
127 GLU (75.1, -137.1)
164 GLU (-169.7, -70.0)
165 ASP (-70.8, -160.8)
176 SER (76.5, -116.1)
203 TYR (174.0, 132.3)
230 GLU (132.9, 43.4)
233 LEU (147.9, -88.7)
360 PRO (-34.1, 94.8)
375 ILE (-28.0, 146.3)
451 PRO (-44.3, 91.6)