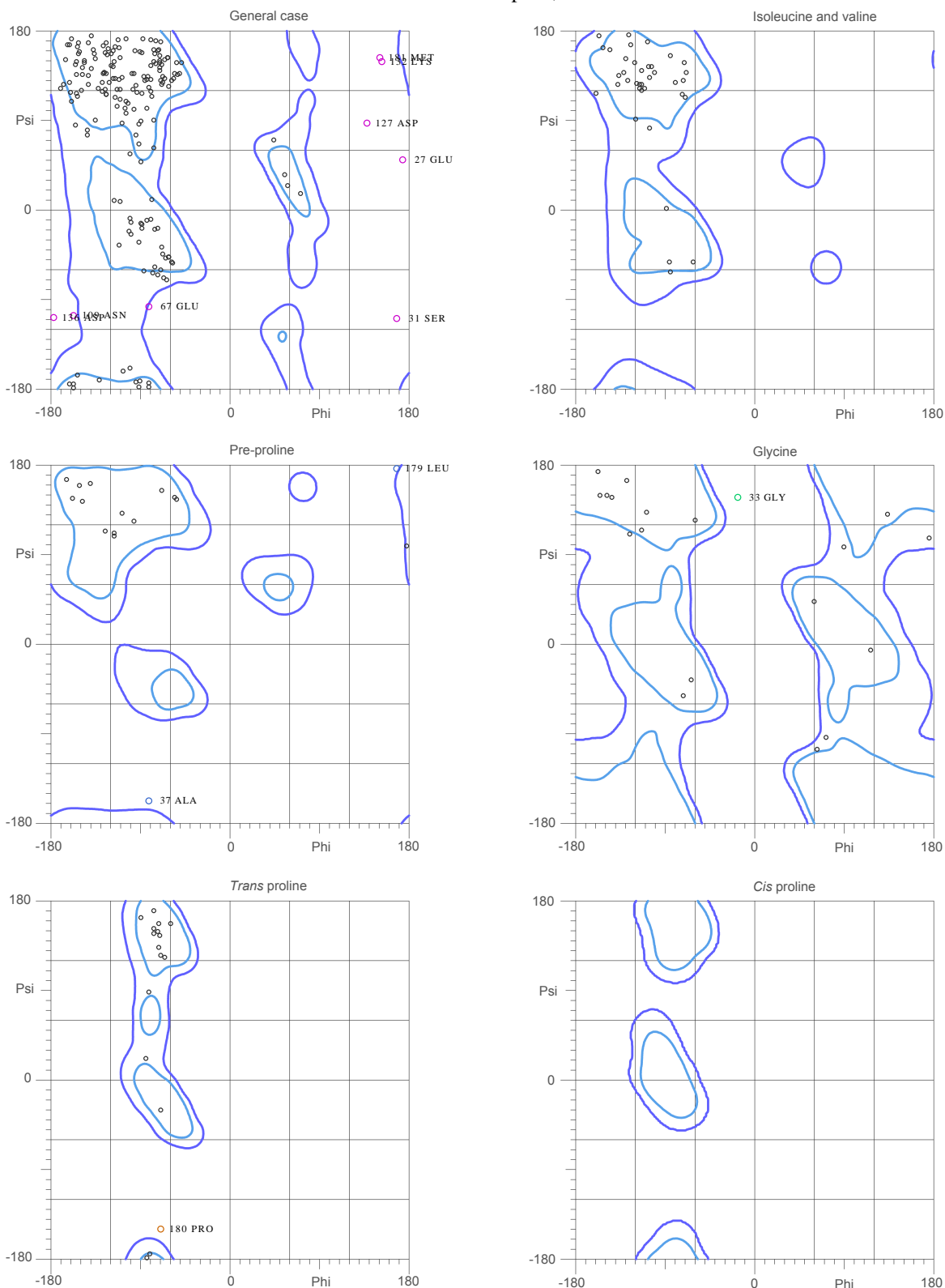


MolProbity Ramachandran analysis

1a02N.B99990033.pdb, model 1



84.9% (236/278) of all residues were in favored (98%) regions.
95.7% (266/278) of all residues were in allowed (>99.8%) regions.

There were 12 outliers (phi, psi):

27 GLU (174.5, 51.9)
31 SER (168.7, -109.2)
33 GLY (-17.7, 148.7)
37 ALA (-82.3, -158.4)
67 GLU (-82.3, -97.5)
109 ASN (-158.4, -106.5)
127 ASP (136.8, 88.6)
132 LYS (153.4, 151.0)
136 ASP (-178.3, -108.6)
179 LEU (168.9, 177.9)
180 PRO (-70.8, -150.8)
181 MET (151.8, 154.5)