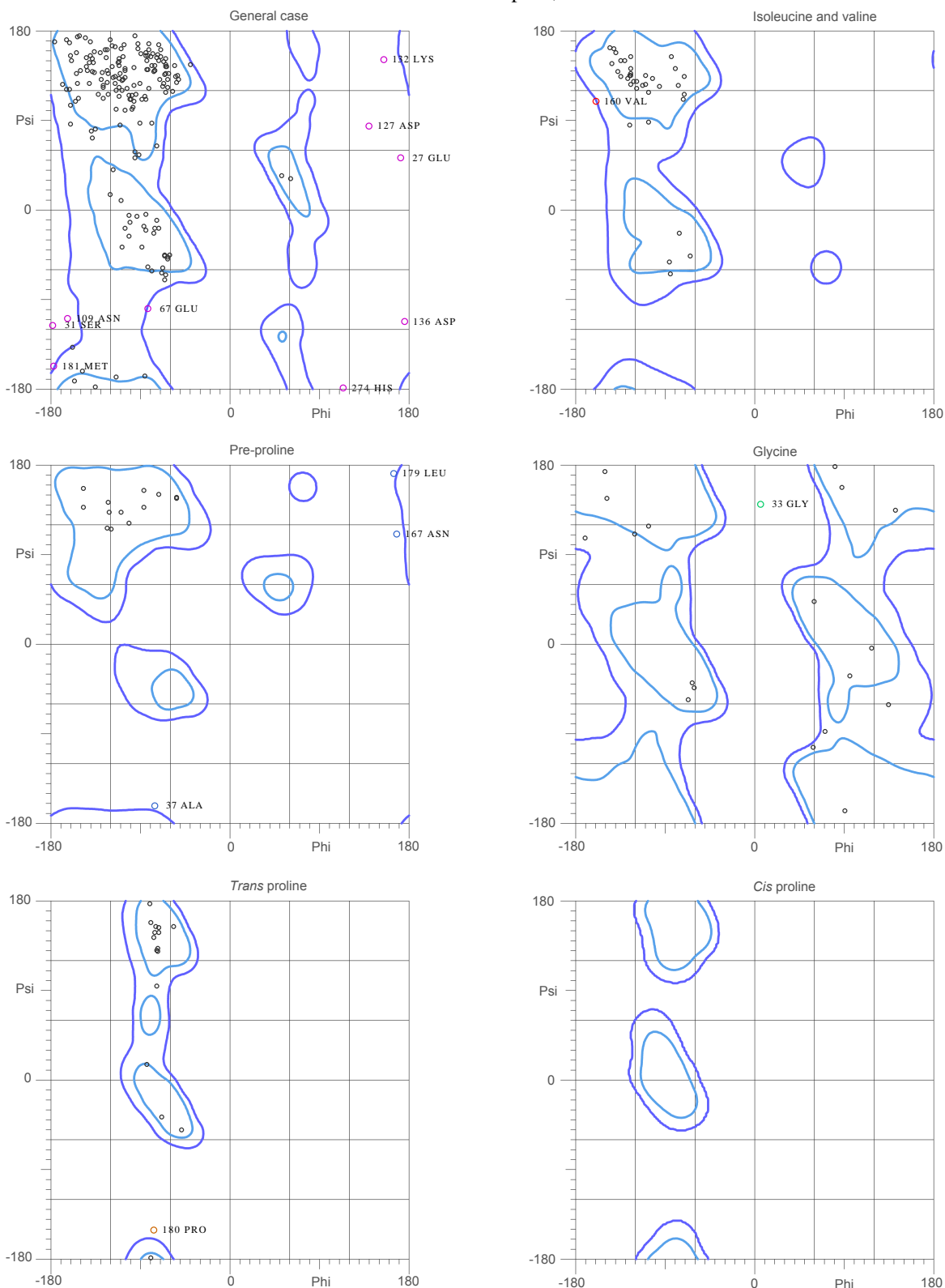


MolProbity Ramachandran analysis

1a02N.B99990014.pdb, model 1



84.2% (234/278) of all residues were in favored (98%) regions.
94.6% (263/278) of all residues were in allowed (>99.8%) regions.

There were 15 outliers (phi, psi):

27 GLU (172.4, 53.3)
31 SER (179.7, -116.2)
33 GLY (6.8, 141.3)
37 ALA (-76.4, -160.8)
67 GLU (-83.7, -99.6)
109 ASN (-166.5, -109.3)
127 ASP (140.5, 85.8)
132 LYS (155.3, 152.0)
136 ASP (176.2, -112.1)
160 VAL (-160.9, -110.8)
167 ASN (168.7, 111.2)
179 LEU (165.7, 172.7)
180 PRO (-72.1, -151.9)
181 MET (-178.1, -157.3)
274 HIS (114.2, -179.1)