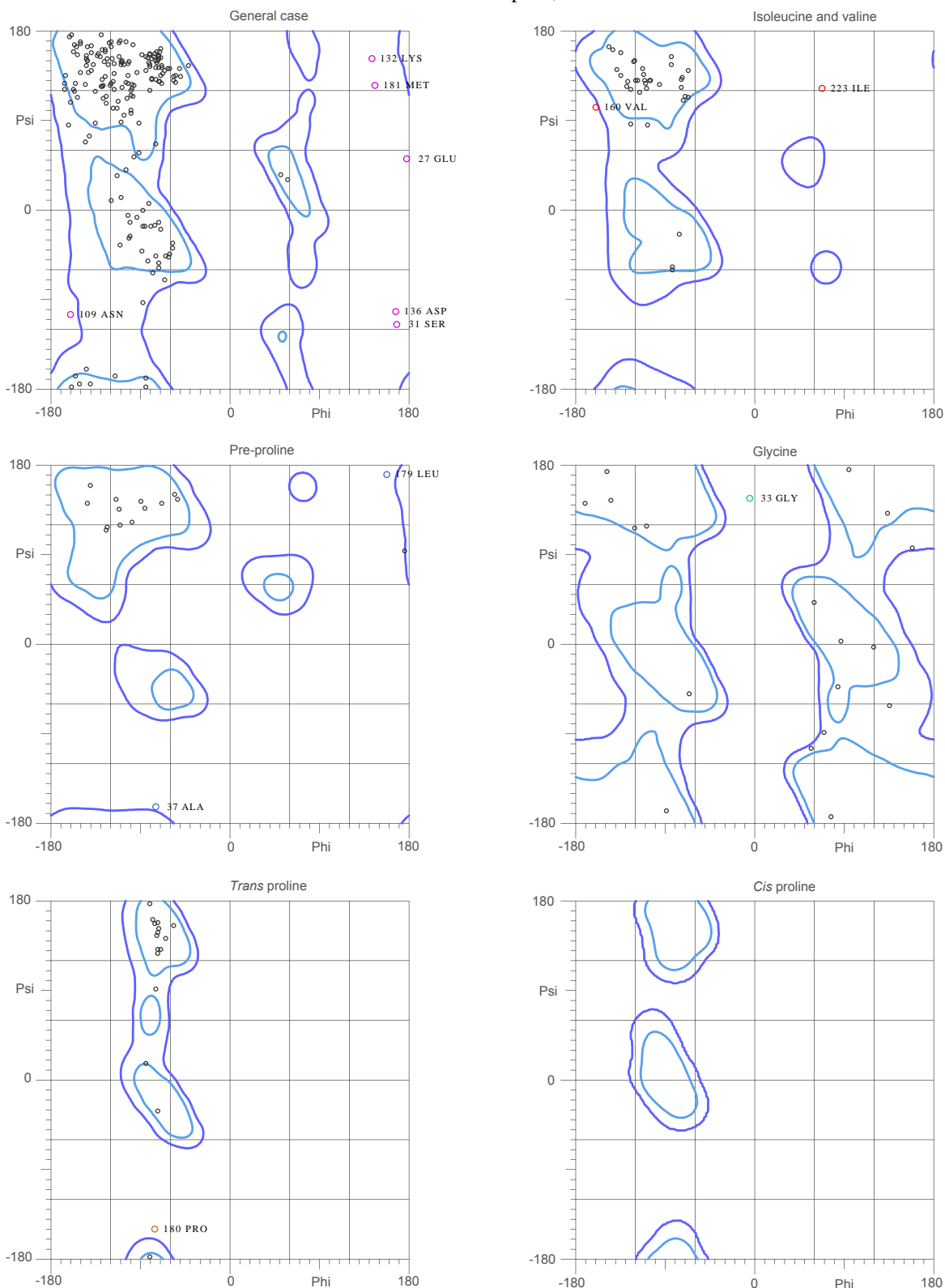


MolProbity Ramachandran analysis

1a02N.B99990019.pdb, model 1



87.1% (242/278) 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 (ϕ , ψ):

27 GLU (178.5, 52.3)
31 SER (168.7, -115.8)
33 GLY (-6.0, 147.8)
37 ALA (-75.8, -164.3)
109 ASN (-161.8, -105.2)
132 LYS (143.1, 153.8)
136 ASP (167.3, -102.5)
160 VAL (-160.8, 104.3)
179 LEU (158.9, 171.1)
180 PRO (-76.1, -150.4)
181 MET (146.6, 126.4)
223 ILE (69.0, 123.9)