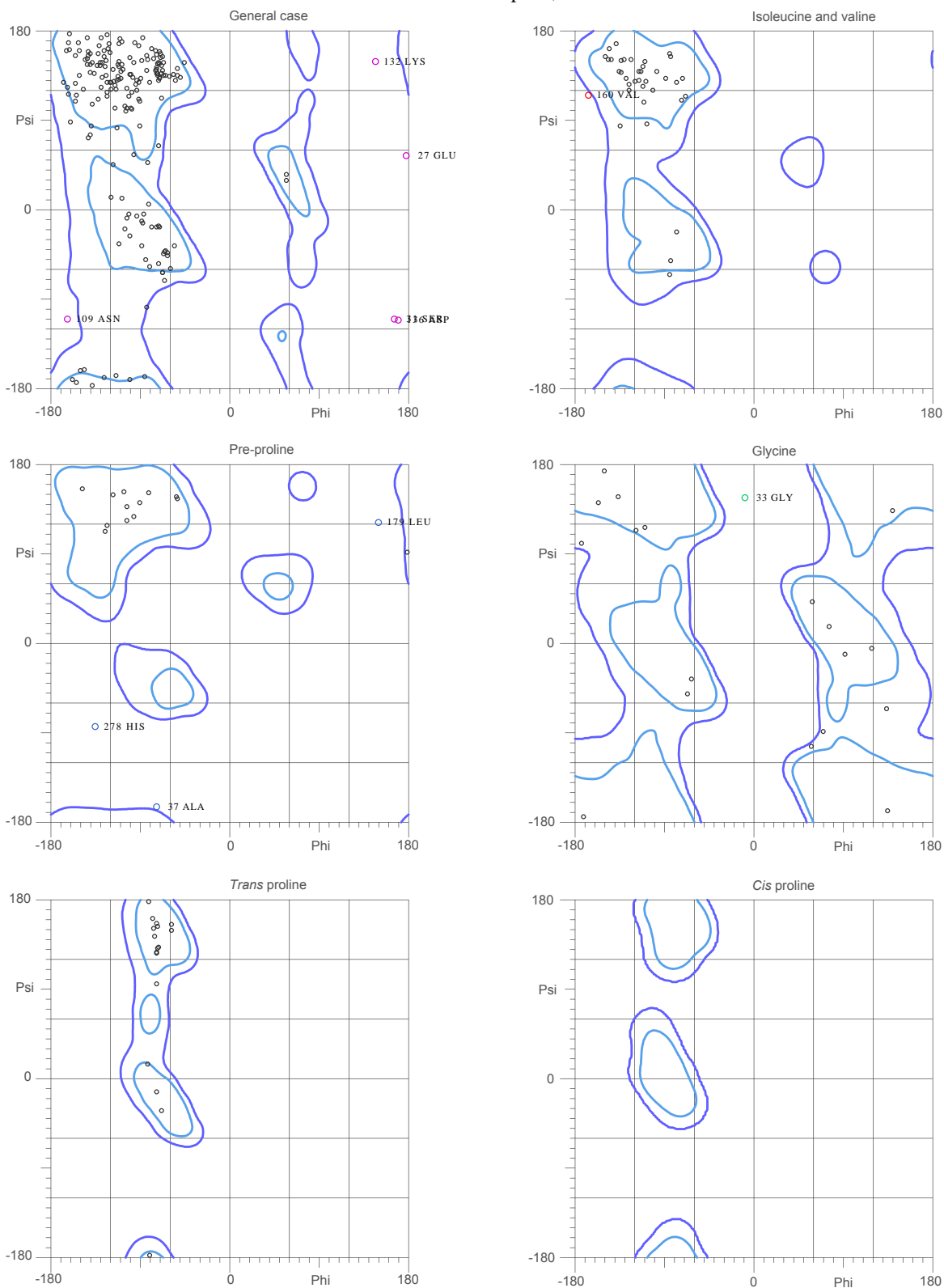


MolProbity Ramachandran analysis

1a02N.B99990028.pdb, model 1



86.0% (239/278) of all residues were in favored (98%) regions.
96.4% (268/278) of all residues were in allowed (>99.8%) regions.

There were 10 outliers (phi, psi):

27 GLU (178.1, 55.6)
31 SER (166.2, -110.1)
33 GLY (-9.8, 148.0)
37 ALA (-74.4, -165.0)
109 ASN (-164.3, -110.1)
132 LYS (147.9, 150.6)
136 ASP (170.2, -111.4)
160 VAL (-167.6, 116.2)
179 LEU (150.2, 122.4)
278 HIS (-136.6, -85.0)