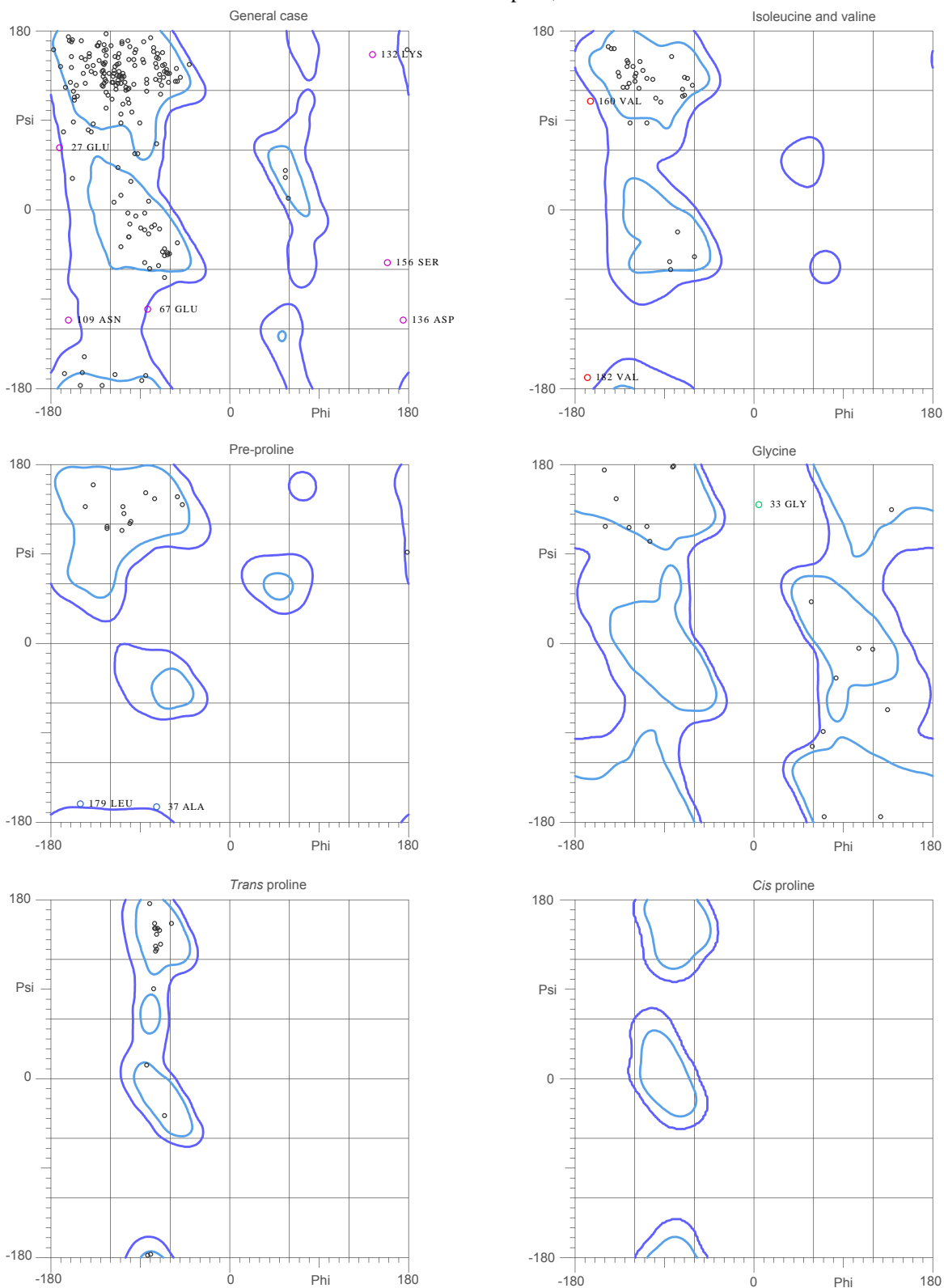


MolProbity Ramachandran analysis

1a02N.B99990008.pdb, model 1



86.7% (241/278) of all residues were in favored (98%) regions.
96.0% (267/278) of all residues were in allowed (>99.8%) regions.

There were 11 outliers (phi, psi):

27 GLU (-172.2, 63.8)
33 GLY (5.8, 140.3)
37 ALA (-74.7, -165.2)
67 GLU (-83.0, -100.2)
109 ASN (-163.5, -111.0)
132 LYS (144.9, 157.3)
136 ASP (175.3, -111.8)
156 SER (159.4, -53.4)
160 VAL (-165.7, 110.0)
179 LEU (-151.9, -162.9)
182 VAL (-168.0, -169.4)