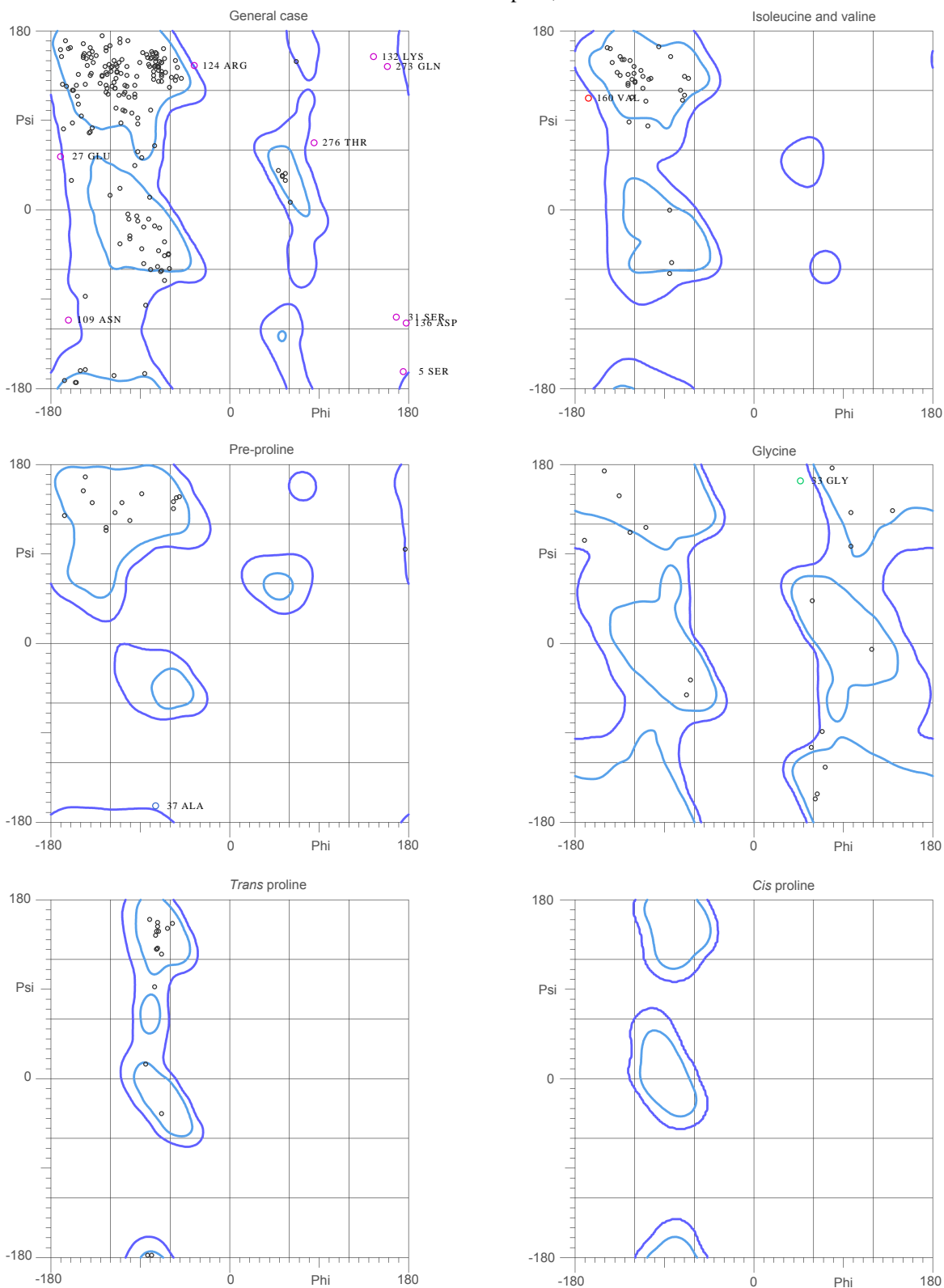


MolProbity Ramachandran analysis

1a02N.B99990039.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):

5 SER (175.2, -143.7)
27 GLU (-171.7, 54.9)
31 SER (168.4, -108.7)
33 GLY (47.6, 164.8)
37 ALA (-75.5, -164.5)
109 ASN (-163.8, -111.2)
124 ARG (-36.8, 146.7)
132 LYS (145.9, 155.8)
136 ASP (178.3, -114.6)
160 VAL (-167.5, 113.6)
273 GLN (159.9, 145.4)
276 THR (85.1, 68.1)