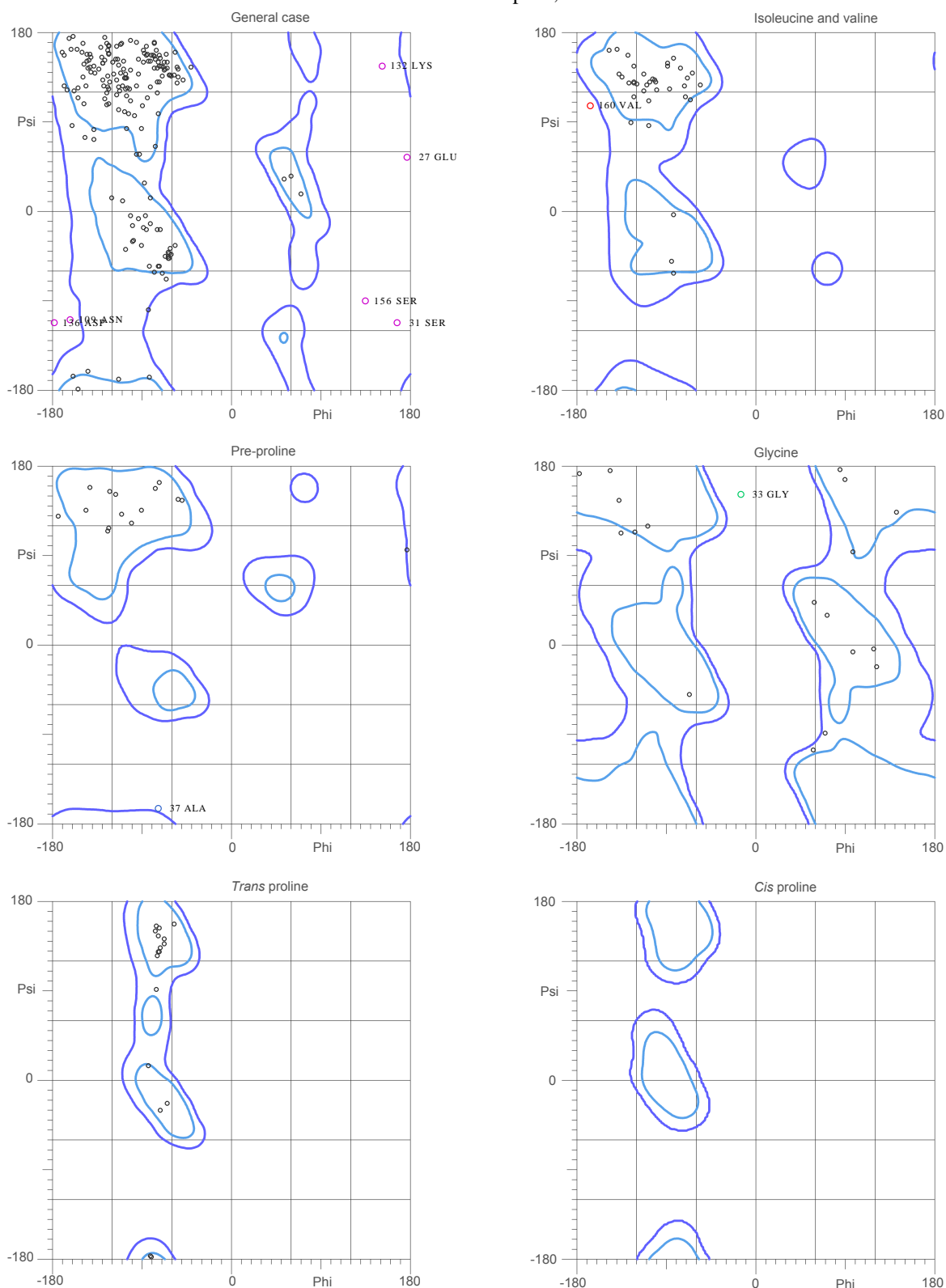# MolProbity Ramachandran analysis

1a02N.B99990020.pdb, model 1

87.4% (243/278) of all residues were in favored (98%) regions.
96.8% (269/278) of all residues were in allowed (>99.8%) regions.

There were 9 outliers (phi, psi):
    27 GLU (177.6, 55.3)
    31 SER (167.3, -112.8)
    33 GLY (-16.0, 153.0)
    37 ALA (-74.5, -165.0)
    109 ASN (-163.6, -109.5)
    132 LYS (152.4, 147.2)
    136 ASP (-179.6, -112.0)
    156 SER (135.5, -90.5)
    160 VAL (-167.8, 107.6)