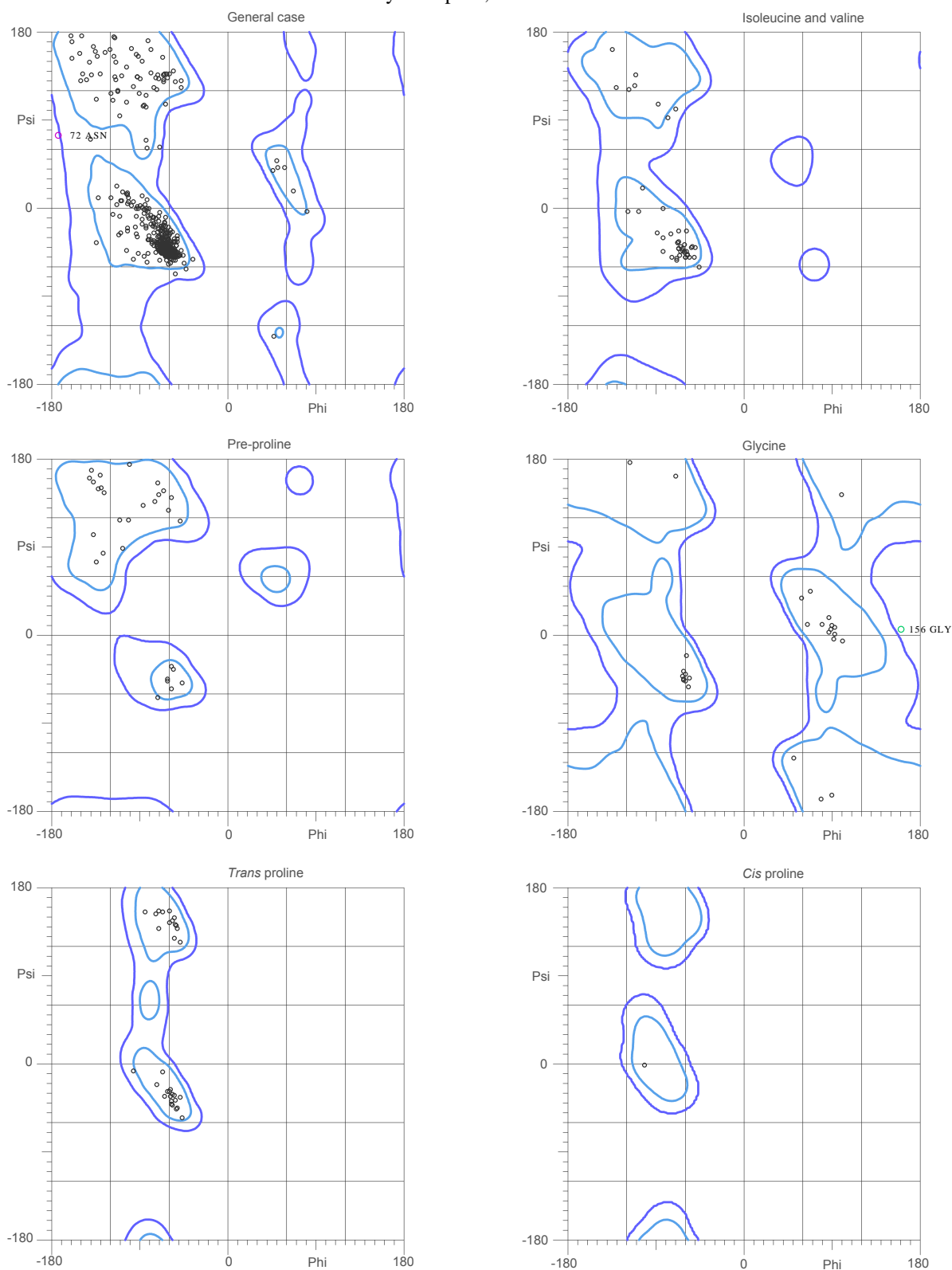


MolProbity Ramachandran analysis

2xy9.H.pdb, model 1



97.2% (561/577) of all residues were in favored (98%) regions.

99.7% (575/577) of all residues were in allowed (>99.8%) regions.

There were 2 outliers (ϕ , ψ):

72 ASN (-174.9, 75.4)

156 GLY (162.0, 6.1)