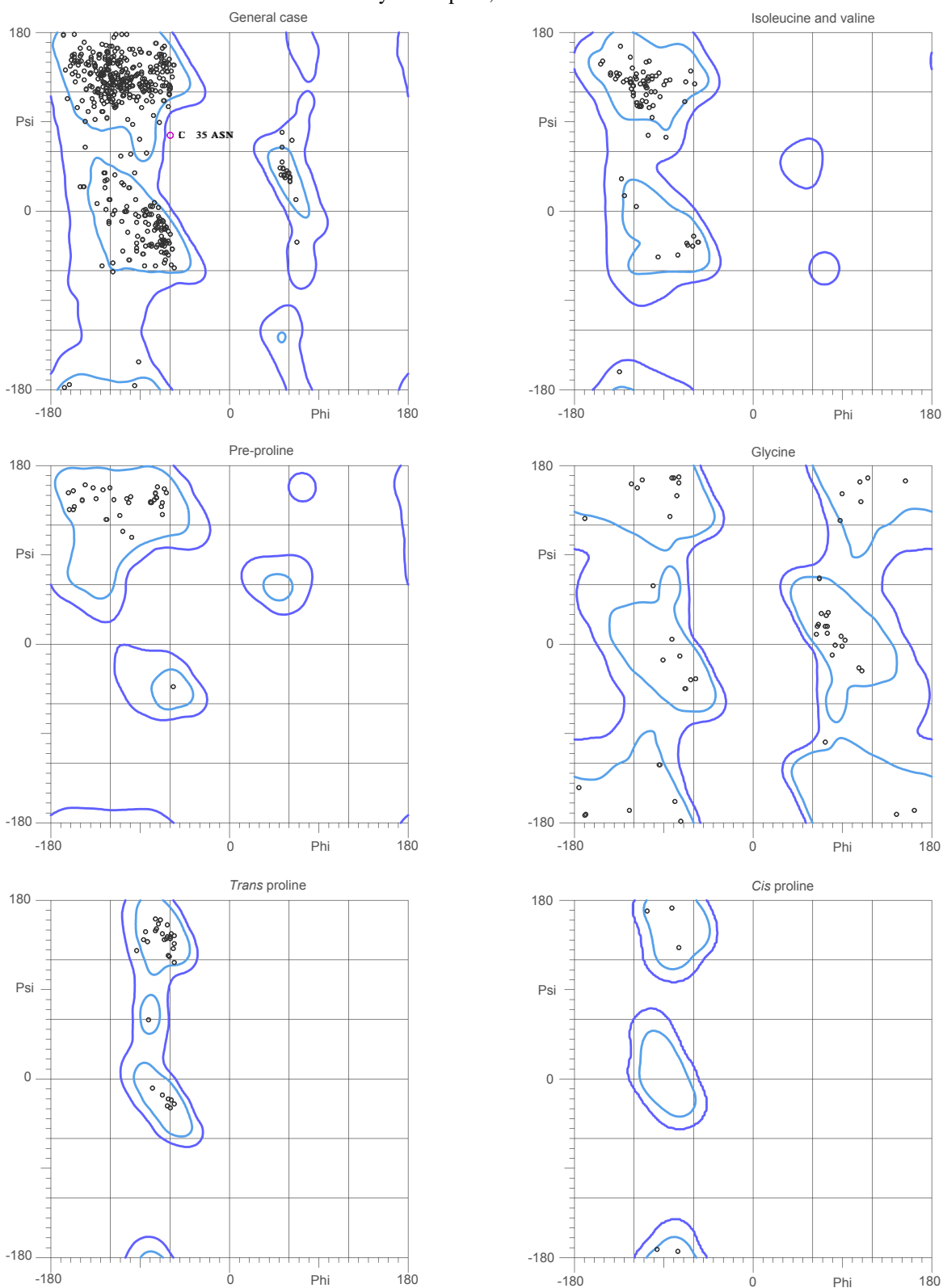


MolProbity Ramachandran analysis

6yor.H.pdb, model 1



96.0% (1186/1236) of all residues were in favored (98%) regions.

99.8% (1234/1236) of all residues were in allowed (>99.8%) regions.

There were 2 outliers (ϕ , ψ):

L 35 ASN (-60.4, 77.1)

C 35 ASN (-60.4, 77.1)