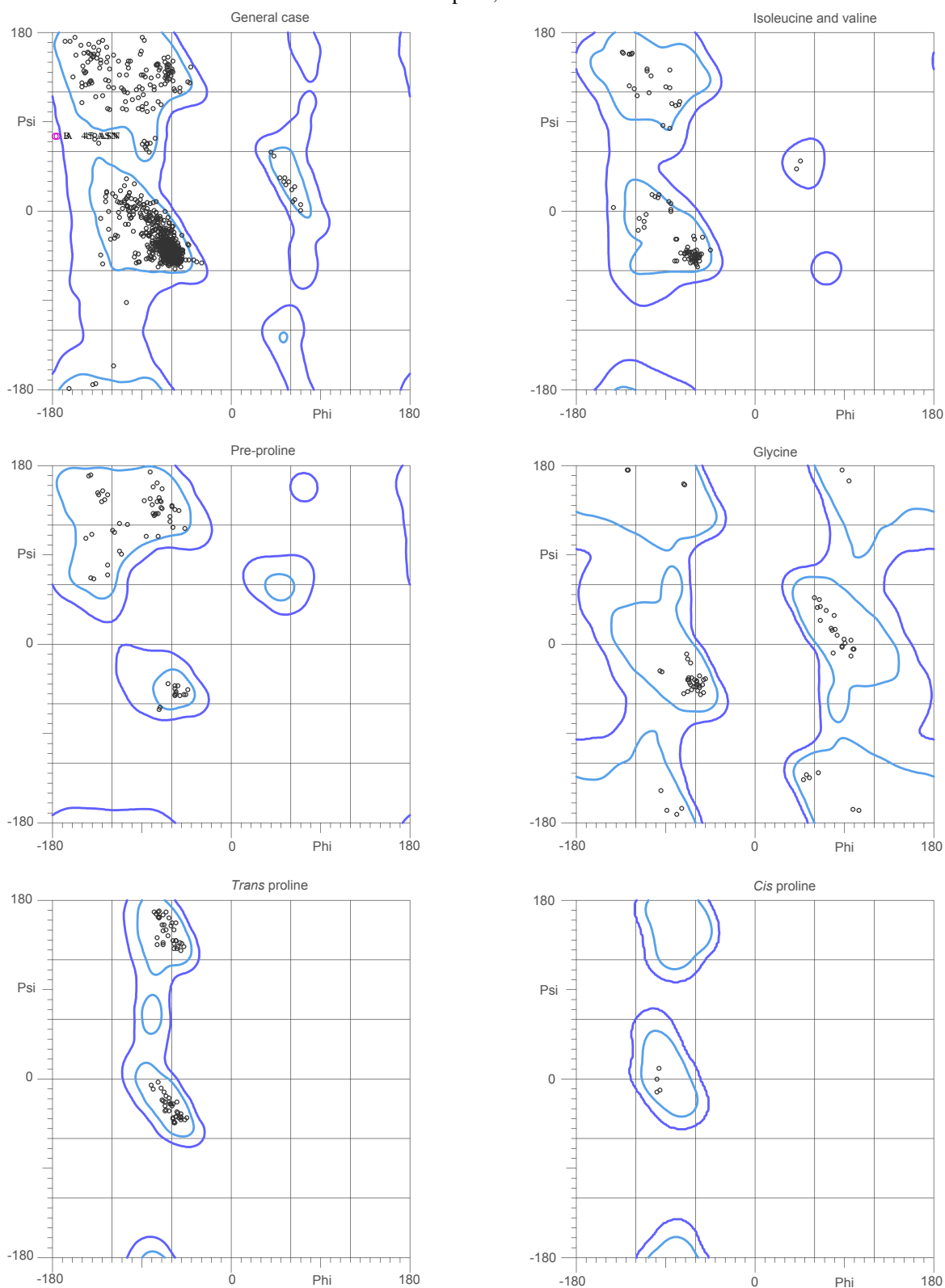


MolProbity Ramachandran analysis

5amc.H.pdb, model 1



98.1% (1187/1210) of all residues were in favored (98%) regions.

99.8% (1208/1210) of all residues were in allowed (>99.8%) regions.

There were 2 outliers (ϕ , ψ):

A 45 ASN (-176.3, 76.7)

B 45 ASN (-178.5, 76.3)