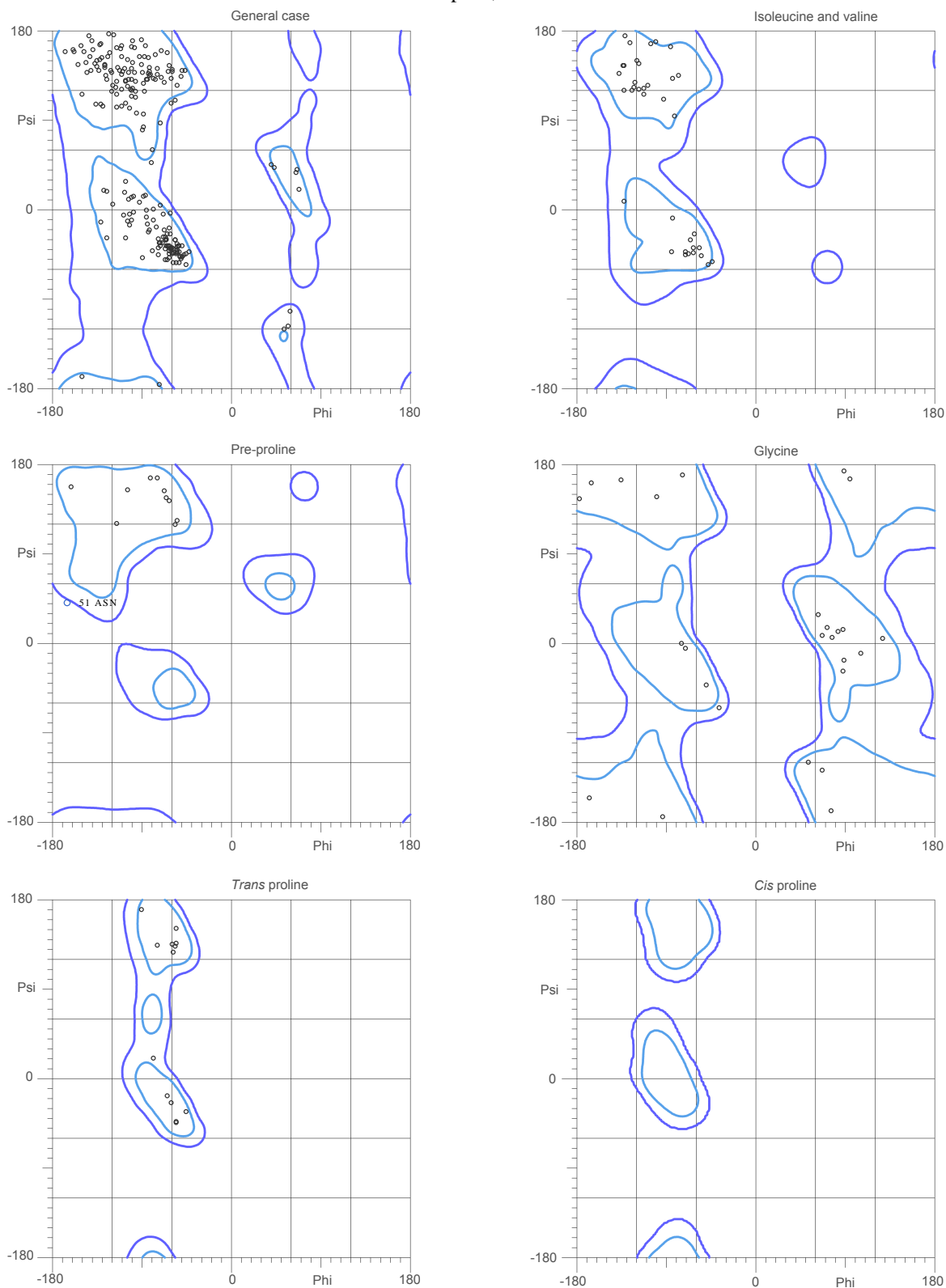


# MolProbity Ramachandran analysis

3sn8.H.pdb, model 1



95.4% (291/305) of all residues were in favored (98%) regions.

99.7% (304/305) of all residues were in allowed (>99.8%) regions.

There were 1 outliers (phi, psi):

51 ASN (-166.4, 41.3)