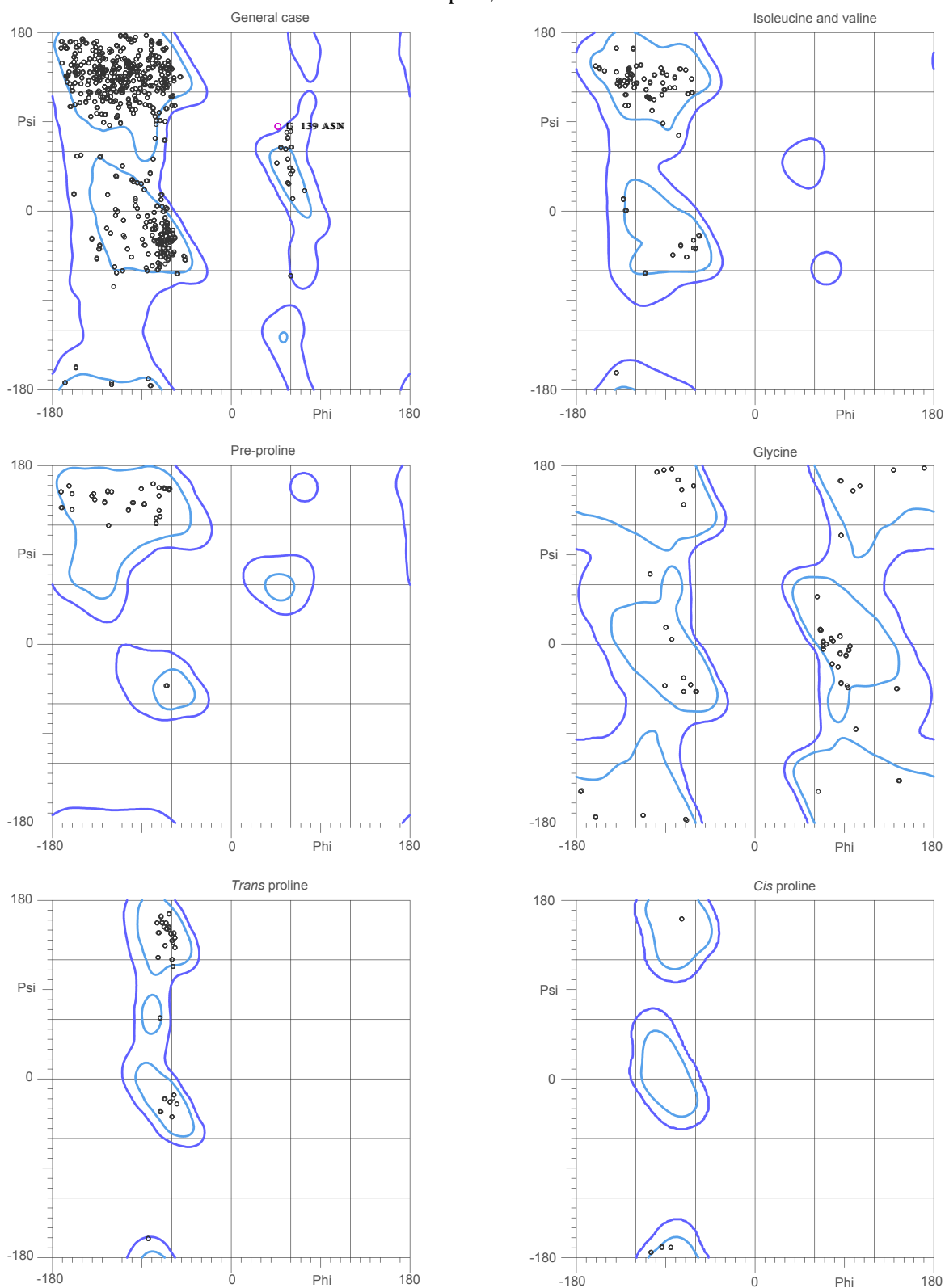


MolProbity Ramachandran analysis

6zer.H.pdb, model 1



92.5% (1732/1872) of all residues were in favored (98%) regions.

99.9% (1870/1872) of all residues were in allowed (>99.8%) regions.

There were 2 outliers (phi, psi):

L 139 ASN (47.3, 86.8)

G 139 ASN (47.6, 86.8)