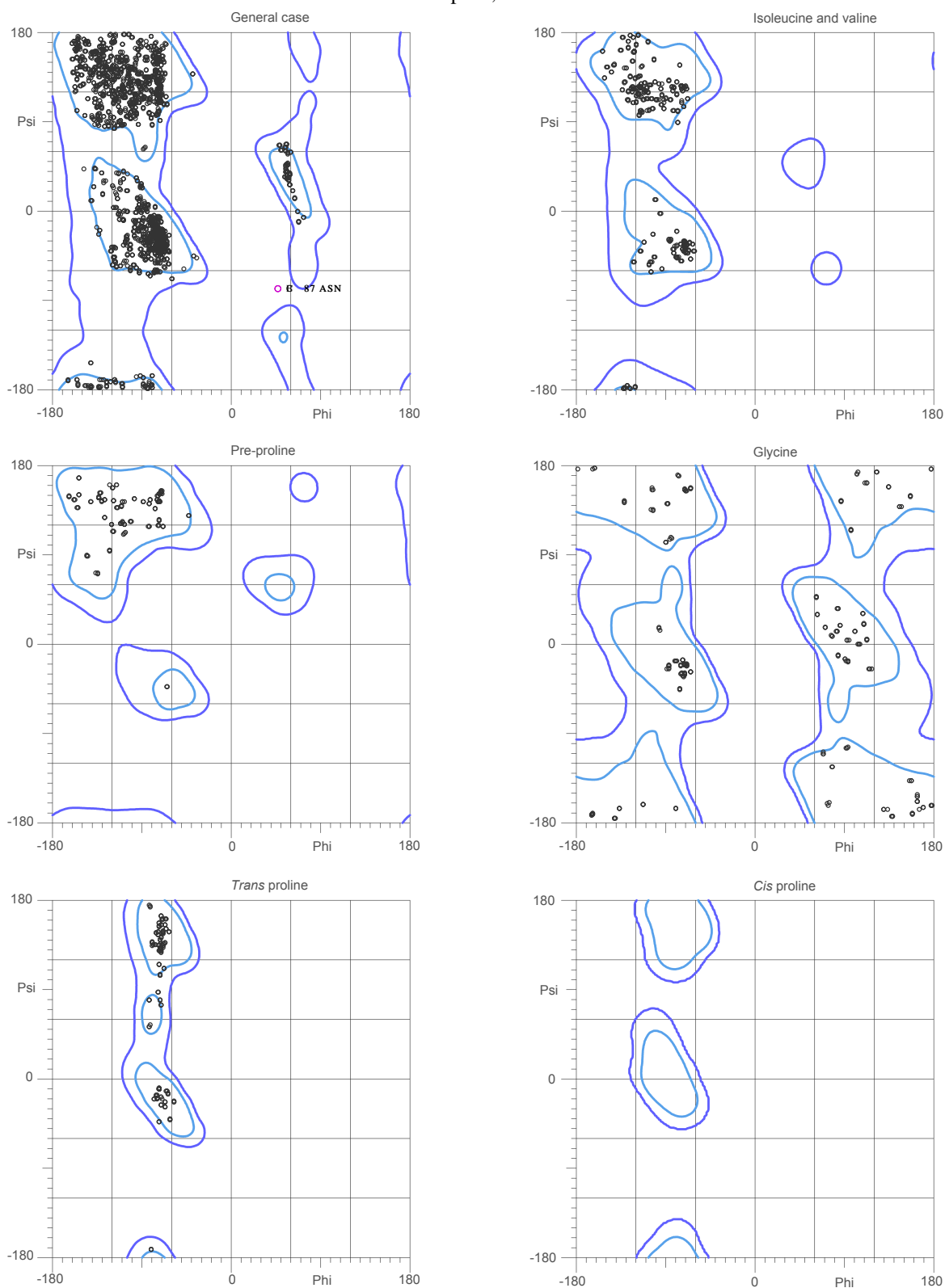


MolProbity Ramachandran analysis

7ddd.H.pdb, model 1



96.4% (3116/3234) of all residues were in favored (98%) regions.

99.9% (3232/3234) of all residues were in allowed (>99.8%) regions.

There were 2 outliers (phi, psi):

B 87 ASN (47.1, -78.4)

C 87 ASN (47.2, -78.3)