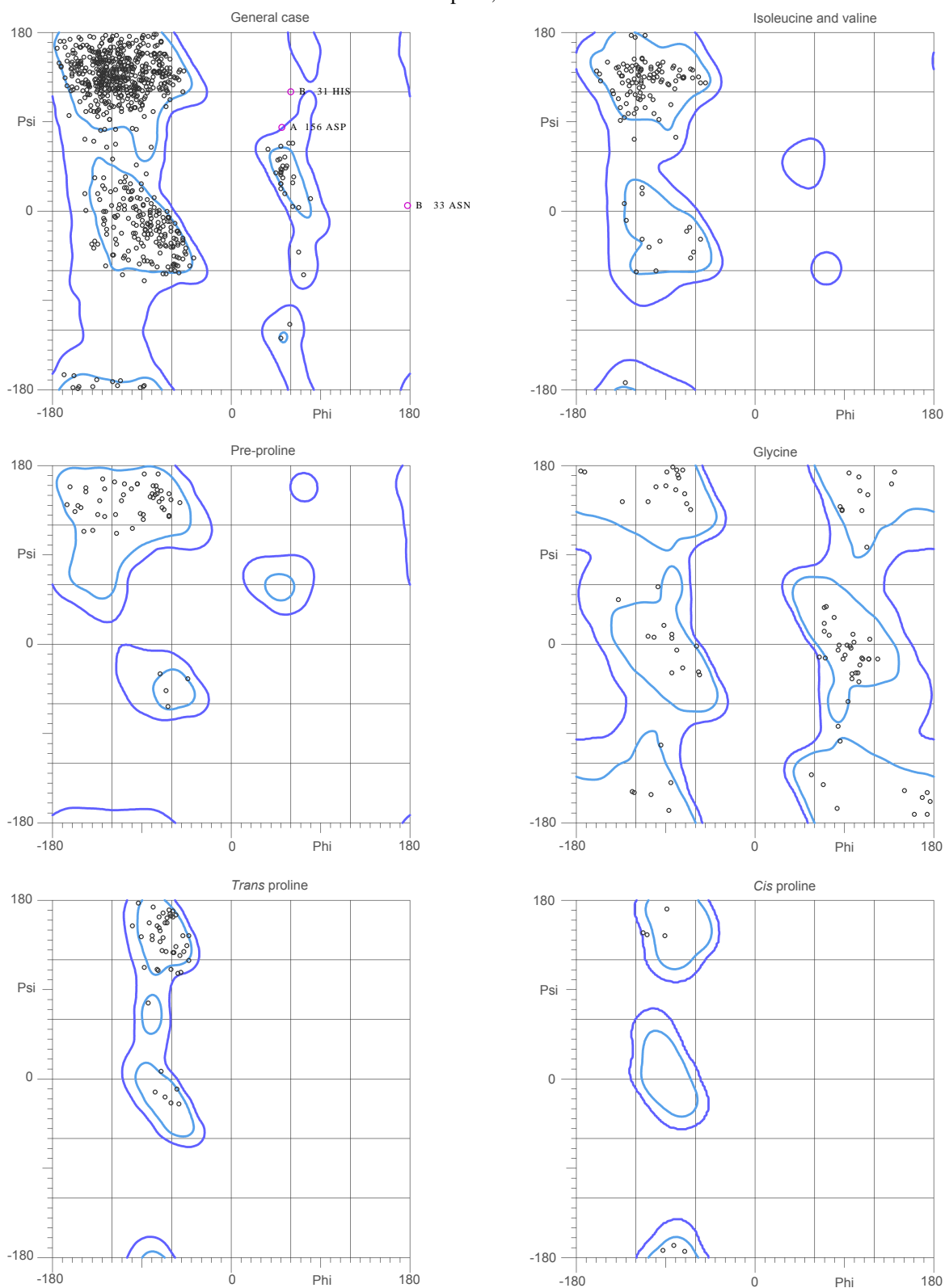


# MolProbity Ramachandran analysis

7che.H.pdb, model 1



93.1% (960/1031) of all residues were in favored (98%) regions.  
99.7% (1028/1031) of all residues were in allowed (>99.8%) regions.

There were 3 outliers (phi, psi):

- A 156 ASP (51.9, 85.1)
- B 31 HIS (60.9, 121.9)
- B 33 ASN (178.7, 6.5)