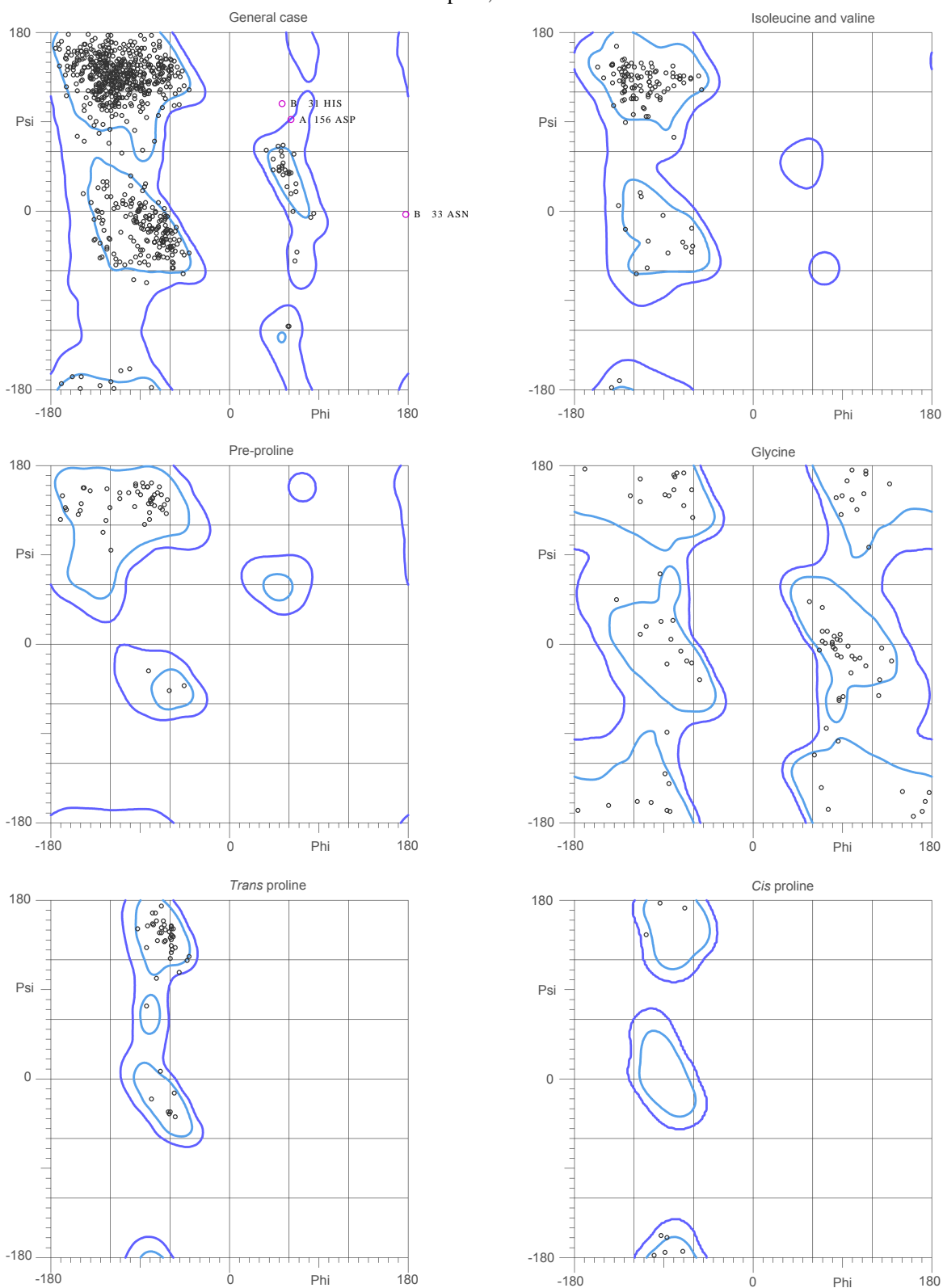


# MolProbity Ramachandran analysis

7chf.H.pdb, model 1



93.7% (973/1038) of all residues were in favored (98%) regions.  
99.7% (1035/1038) of all residues were in allowed (>99.8%) regions.

There were 3 outliers (phi, psi):

- A 156 ASP (62.3, 93.6)
- B 31 HIS (53.3, 109.8)
- B 33 ASN (178.6, -3.6)