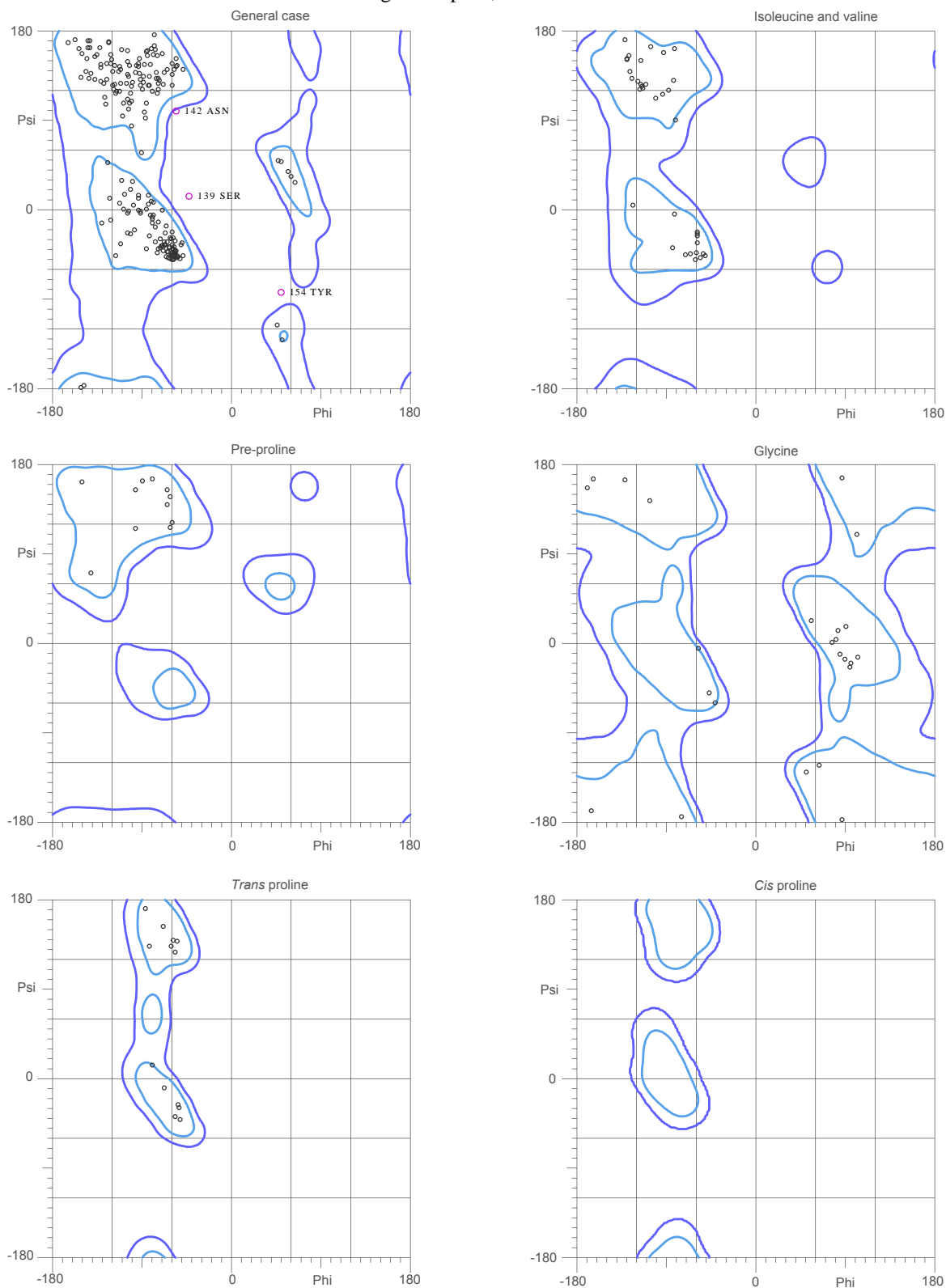


MolProbity Ramachandran analysis

2gt8.H.pdb, model 1



97.6% (289/296) of all residues were in favored (98%) regions.
99.0% (293/296) of all residues were in allowed (>99.8%) regions.

There were 3 outliers (phi, psi):

139 SER (-43.9, 14.8)
142 ASN (-56.0, 100.2)
154 TYR (50.2, -83.7)