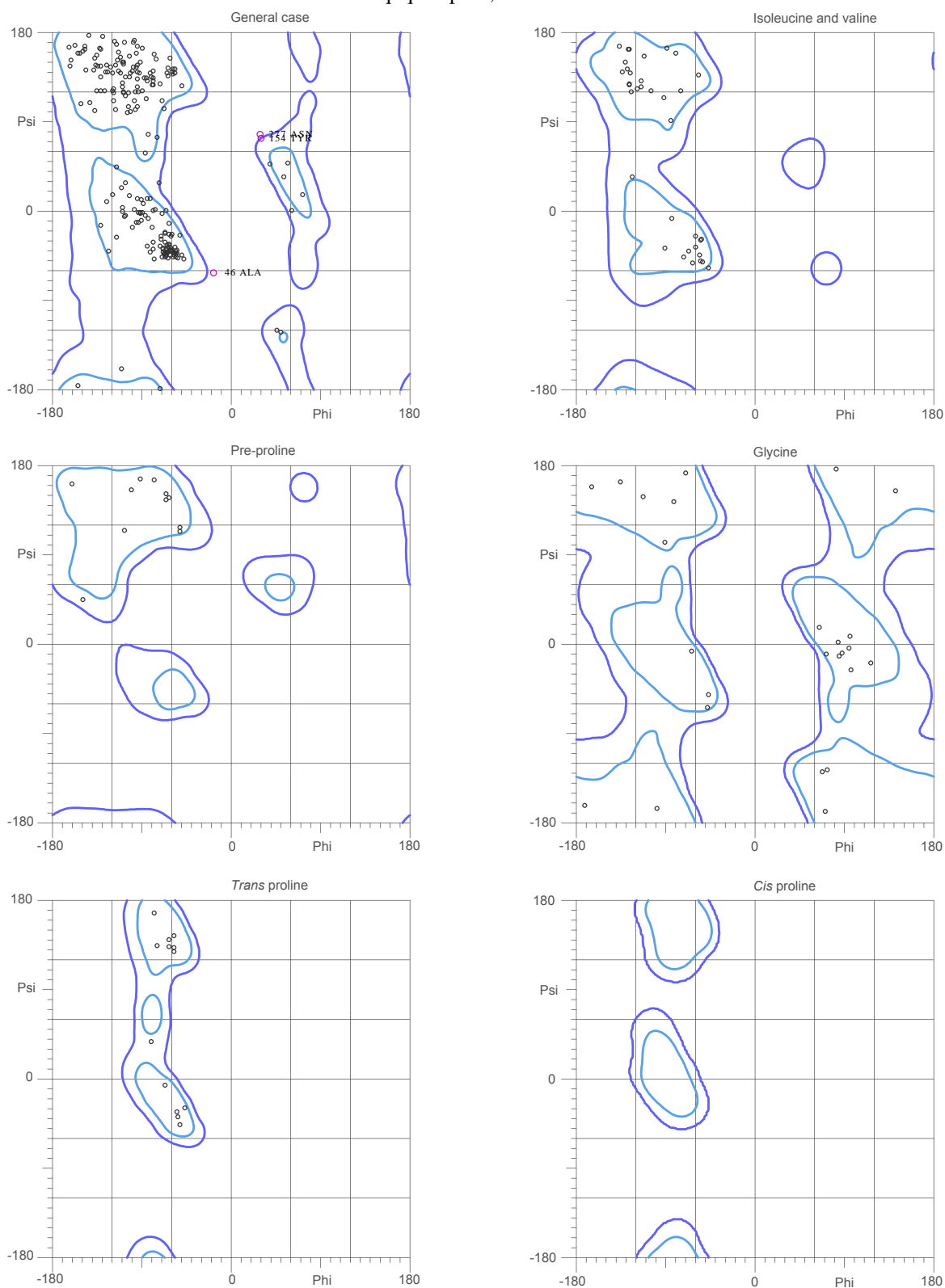


# MolProbity Ramachandran analysis

2qiq.H.pdb, model 1



94.6% (283/299) of all residues were in favored (98%) regions.  
99.0% (296/299) of all residues were in allowed (>99.8%) regions.

There were 3 outliers (phi, psi):

46 ALA (-18.9, -62.9)

154 TYR (30.6, 74.8)

277 ASN (29.8, 78.2)