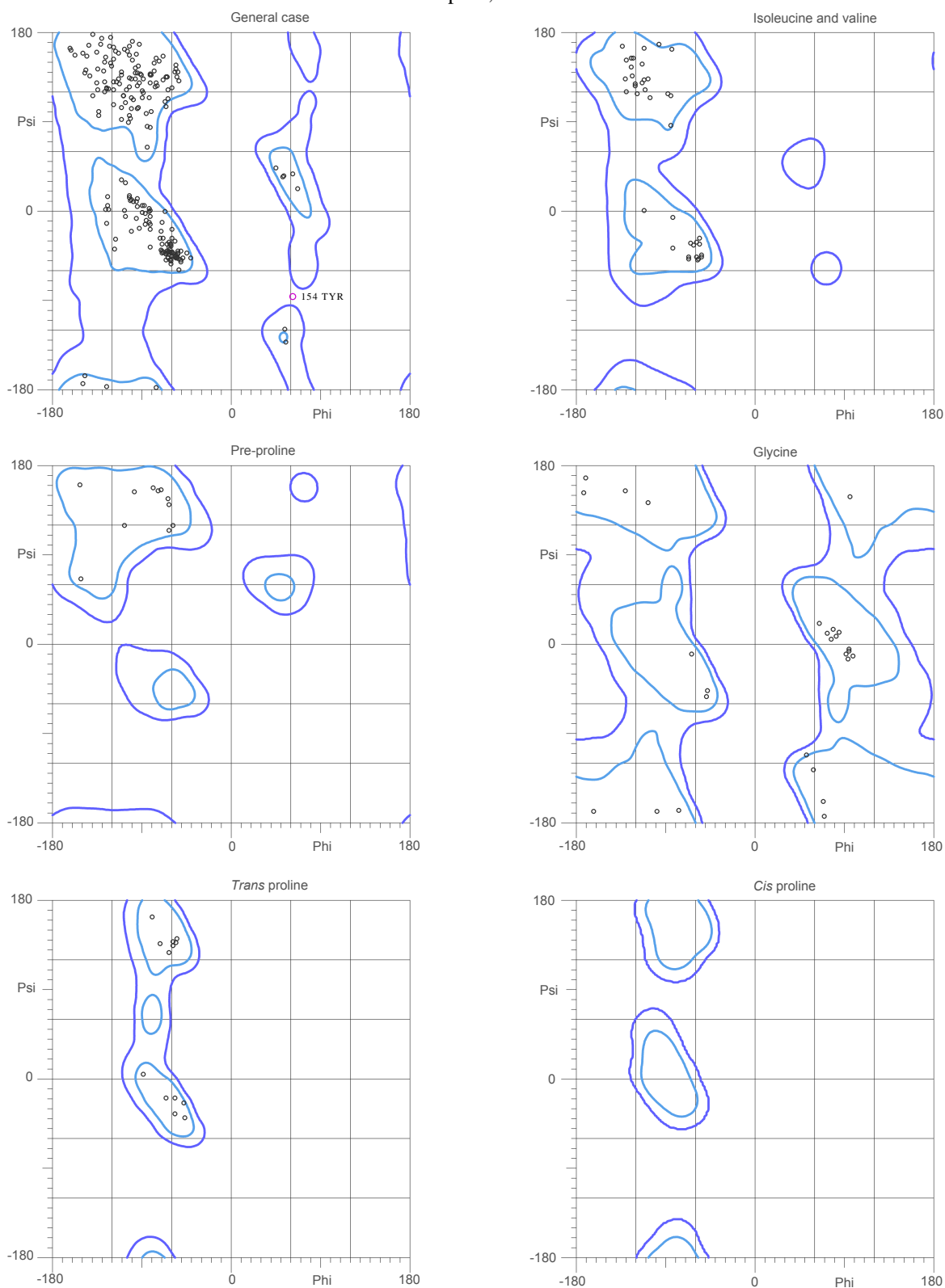


# MolProbity Ramachandran analysis

5rfa.H.pdb, model 1



98.4% (303/308) of all residues were in favored (98%) regions.

99.7% (307/308) of all residues were in allowed (>99.8%) regions.

There were 1 outliers (phi, psi):

154 TYR (62.7, -86.8)