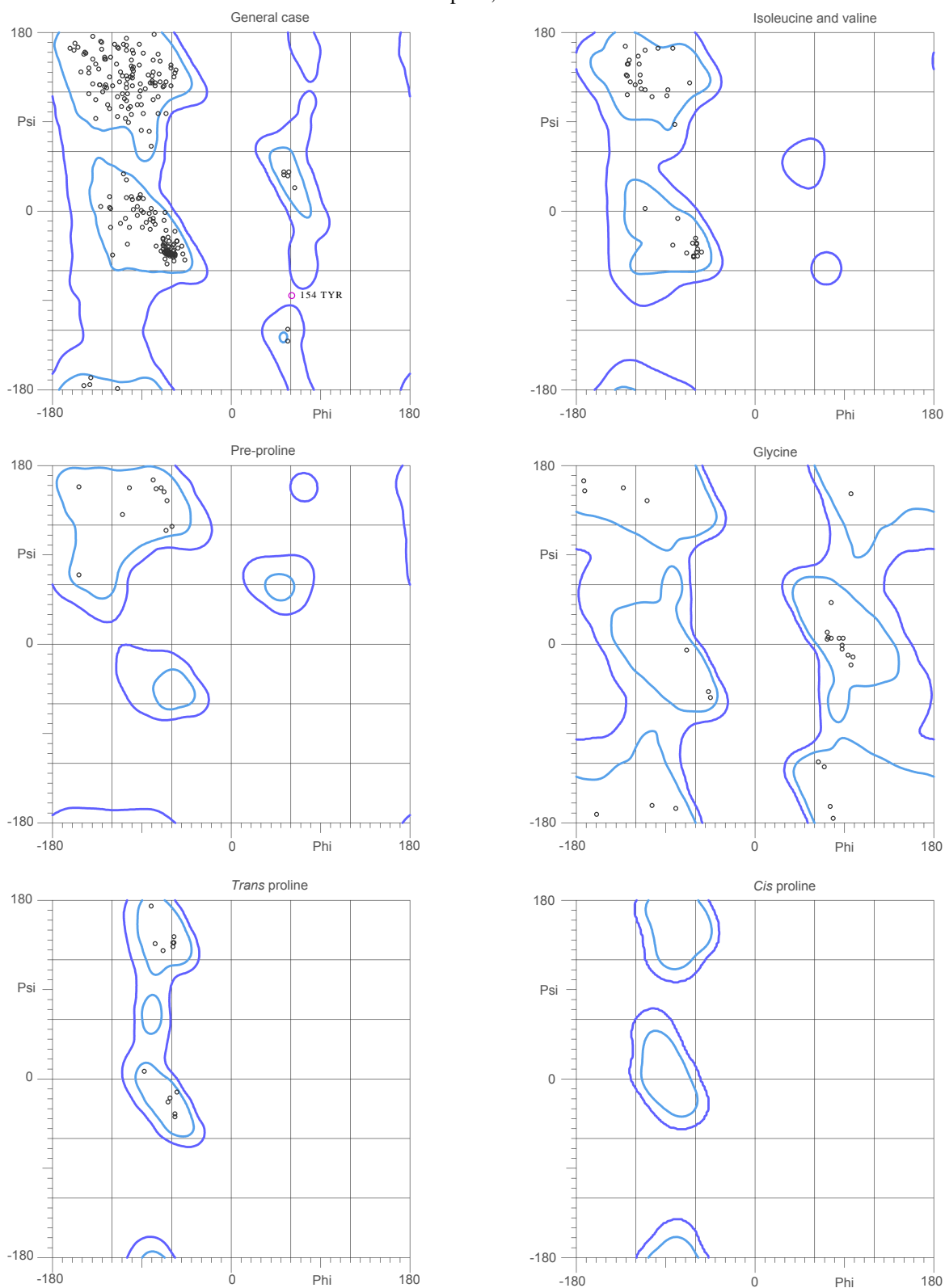


# MolProbity Ramachandran analysis

5rhc.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 (61.4, -85.0)