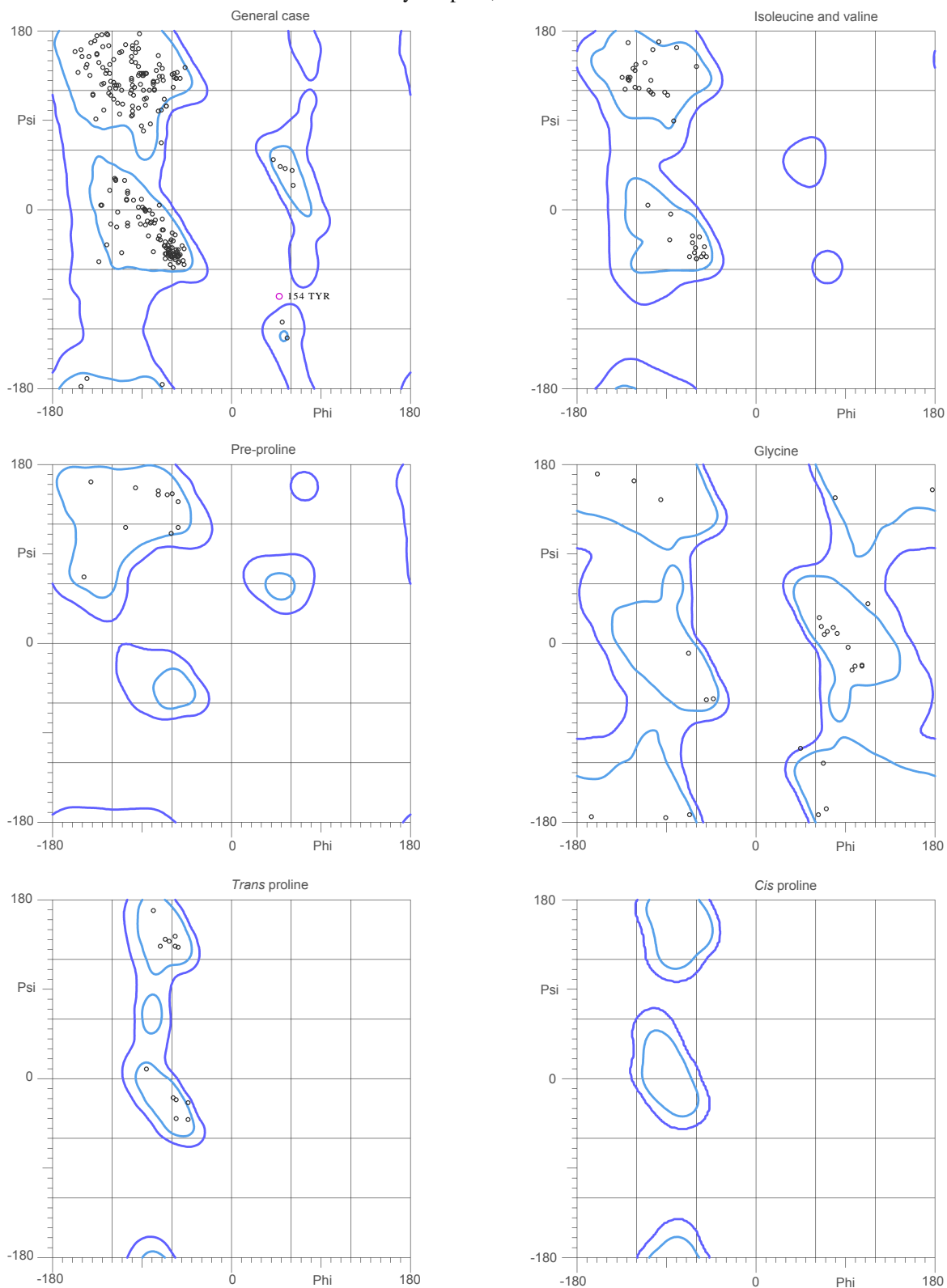


# MolProbity Ramachandran analysis

5rey.H.pdb, model 1



96.1% (299/311) of all residues were in favored (98%) regions.

99.7% (310/311) of all residues were in allowed (>99.8%) regions.

There were 1 outliers (phi, psi):

154 TYR (48.2, -87.6)