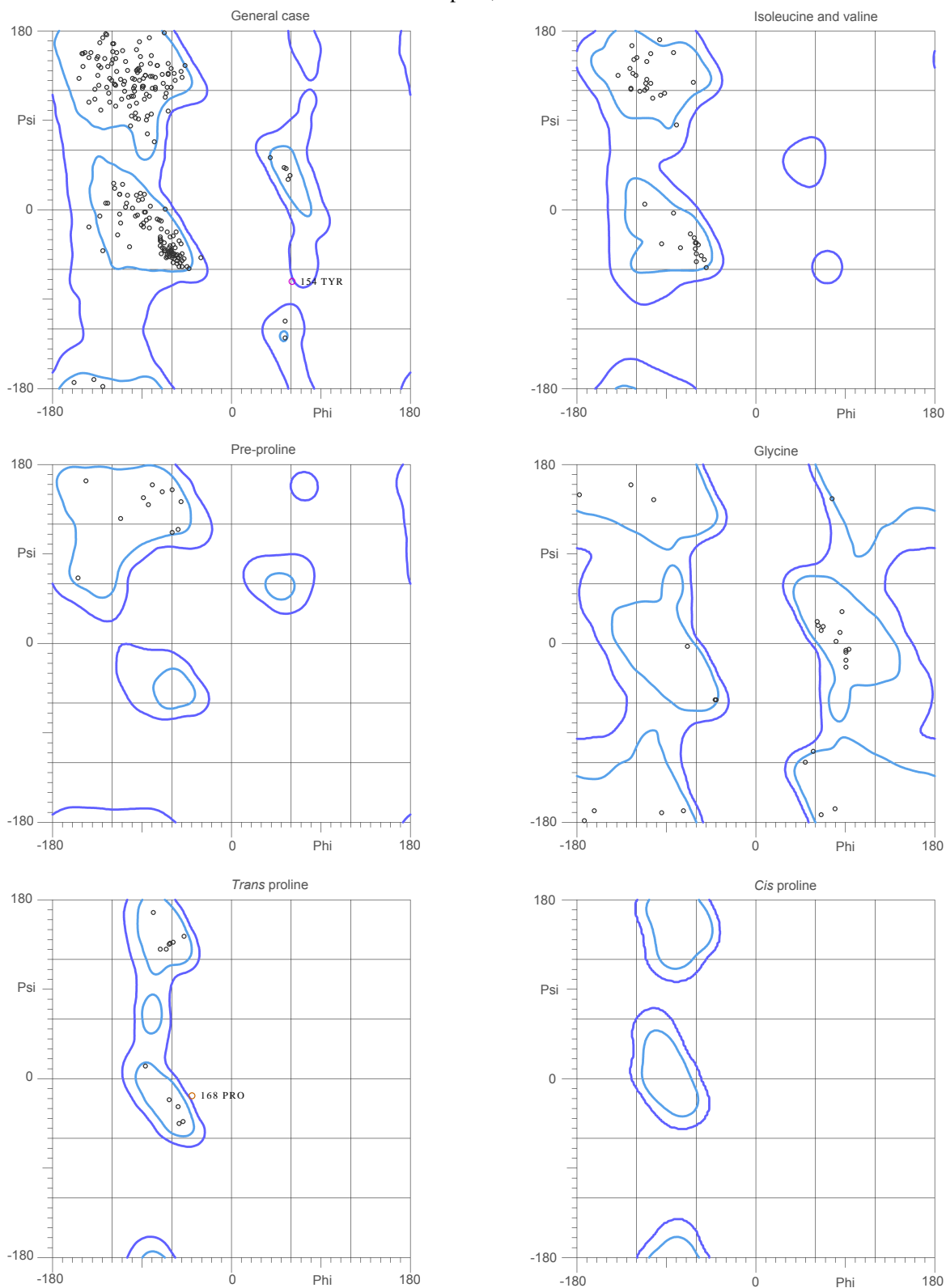


# MolProbity Ramachandran analysis

5rex.H.pdb, model 1



96.5% (300/311) of all residues were in favored (98%) regions.

99.4% (309/311) of all residues were in allowed (>99.8%) regions.

There were 2 outliers (phi, psi):

154 TYR (61.5, -72.2)

168 PRO (-40.6, -17.3)