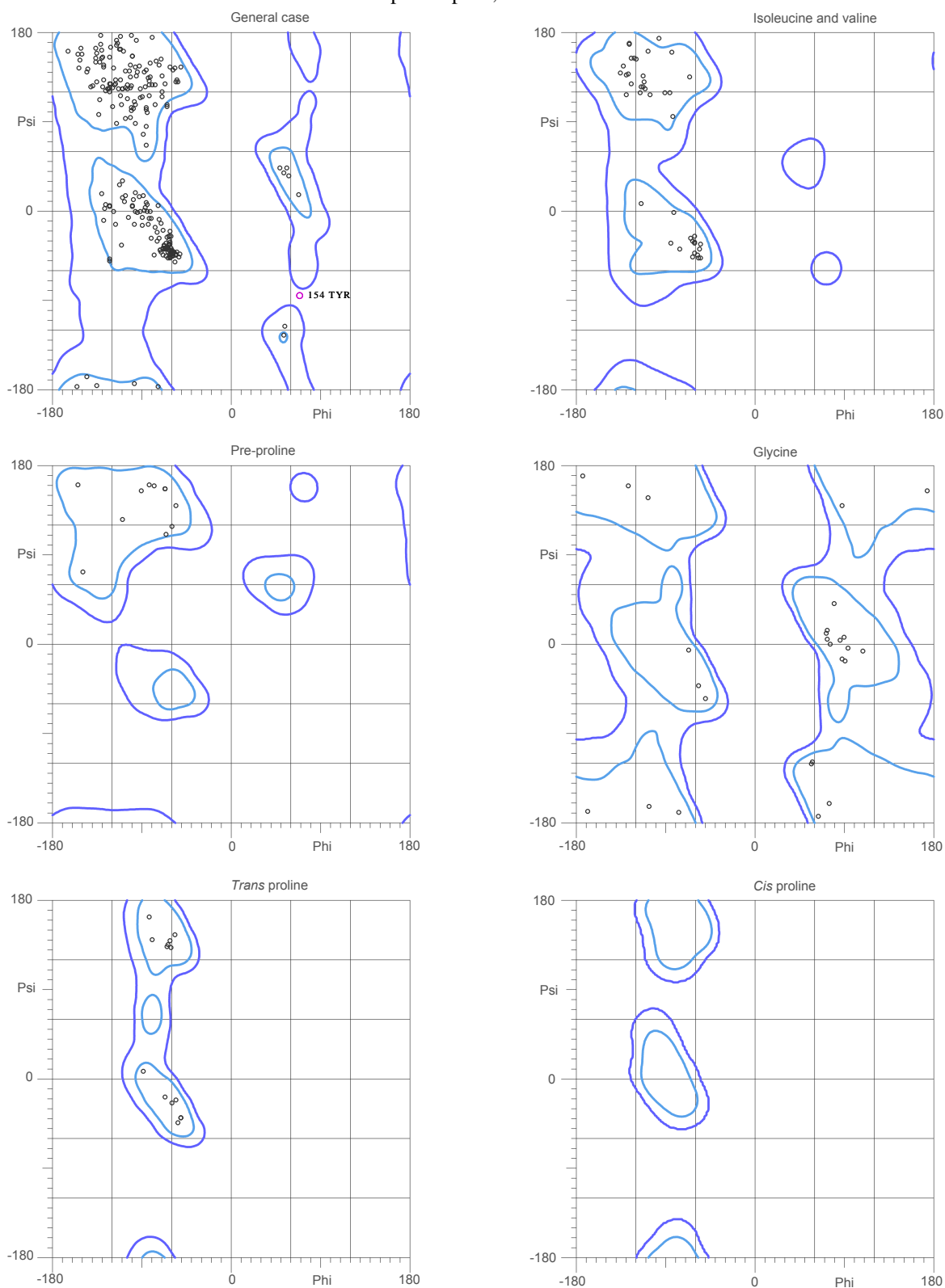


# MolProbity Ramachandran analysis

7ap6.H.pdb, model 1



97.6% (320/328) of all residues were in favored (98%) regions.

99.4% (326/328) of all residues were in allowed (>99.8%) regions.

There were 2 outliers (phi, psi):

154 TYR (69.6, -85.5)

154 TYR (69.6, -85.6)