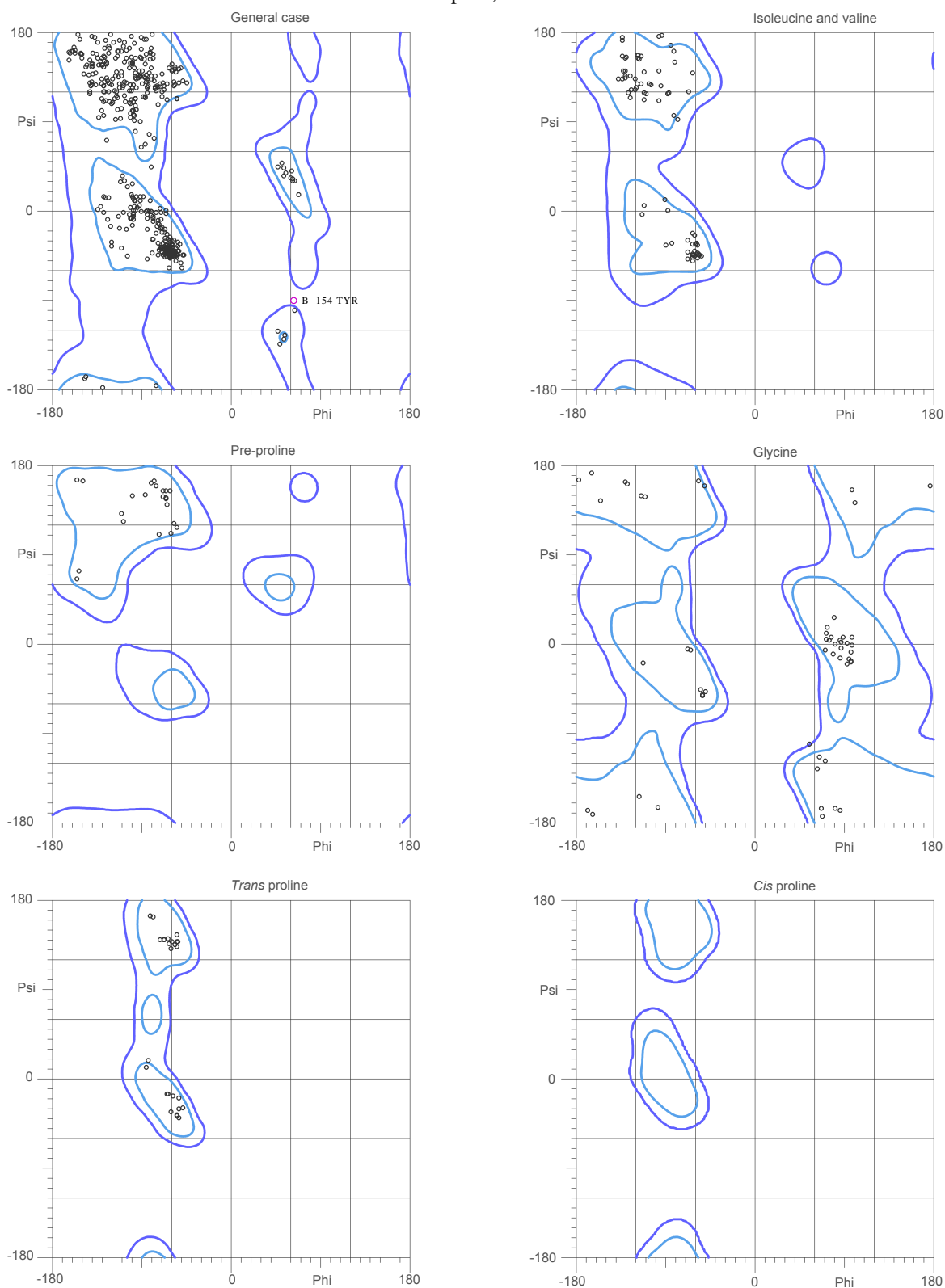


# MolProbity Ramachandran analysis

7avd.H.pdb, model 1



97.6% (600/615) of all residues were in favored (98%) regions.

99.8% (614/615) of all residues were in allowed (>99.8%) regions.

There were 1 outliers (phi, psi):

B 154 TYR (63.9, -90.3)