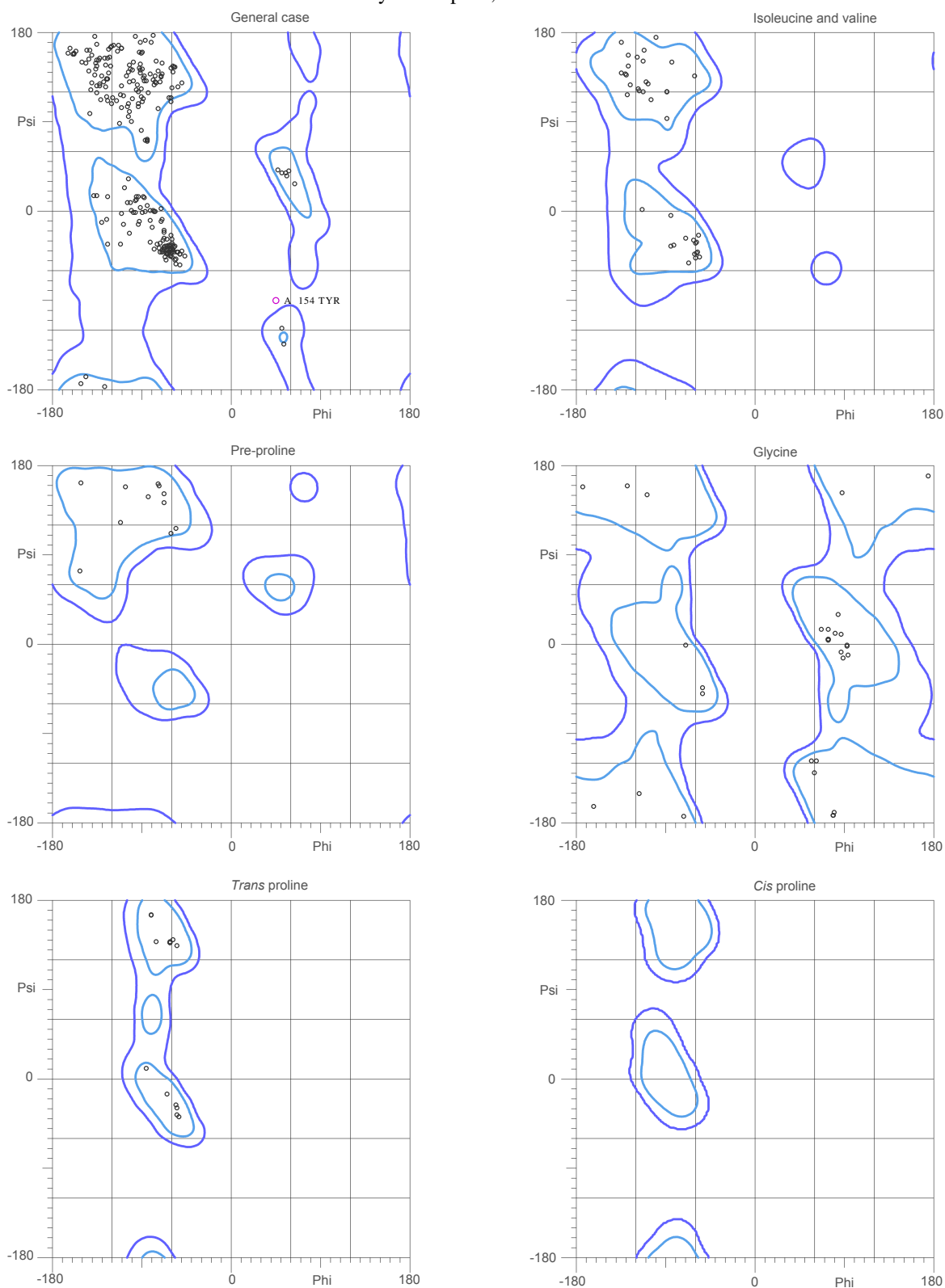


# MolProbity Ramachandran analysis

6yz6.H.pdb, model 1



98.5% (328/333) of all residues were in favored (98%) regions.

99.7% (332/333) of all residues were in allowed (>99.8%) regions.

There were 1 outliers (phi, psi):

A 154 TYR (45.3, -90.4)