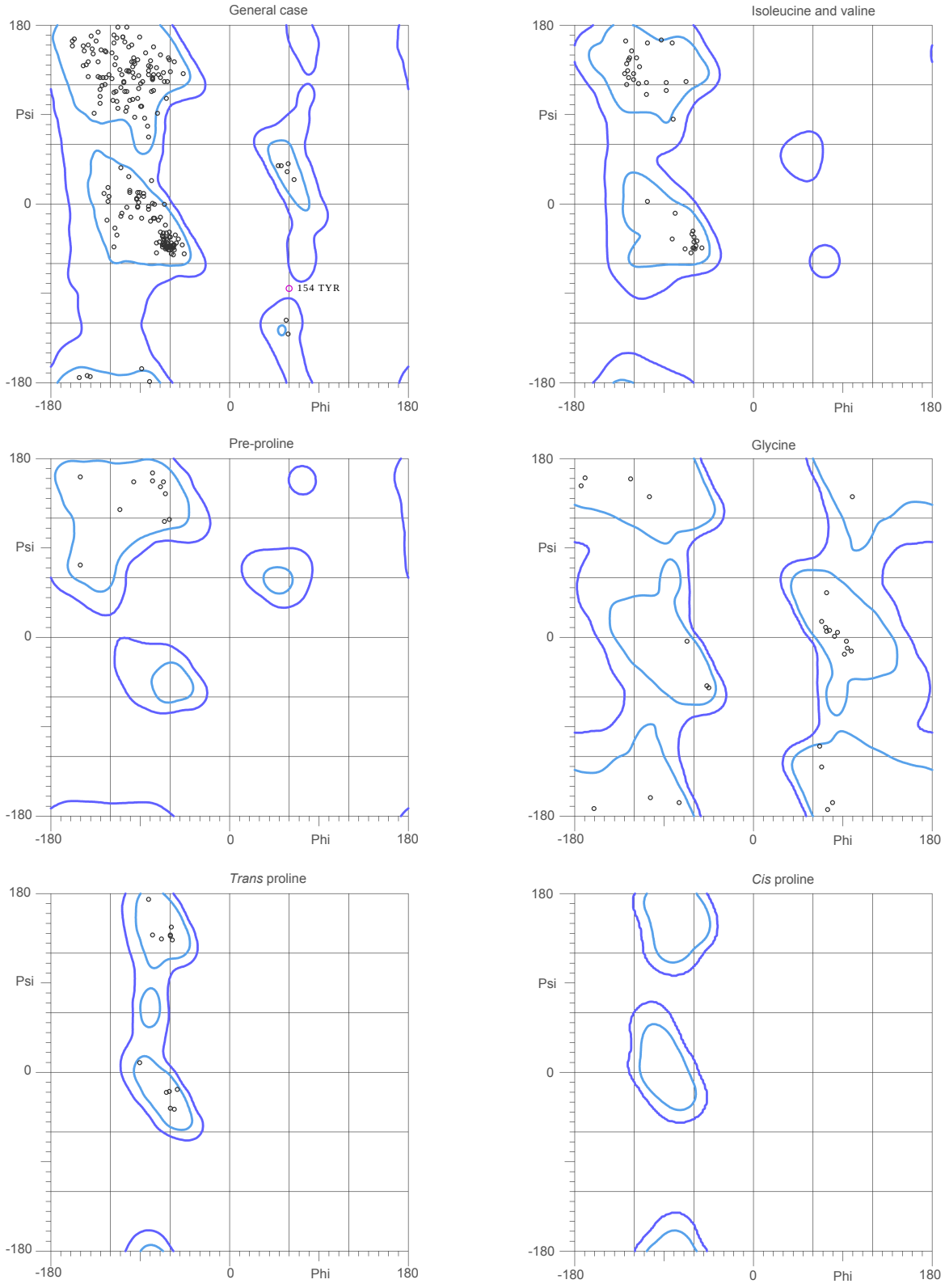
MolProbity Ramachandran analysis

5rh6.H.pdb, model 1



97.4% (296/304) of all residues were in favored (98%) regions.
99.7% (303/304) of all residues were in allowed (>99.8%) regions.

There were 1 outliers (phi, psi):
154 TYR (60.0, -85.9)