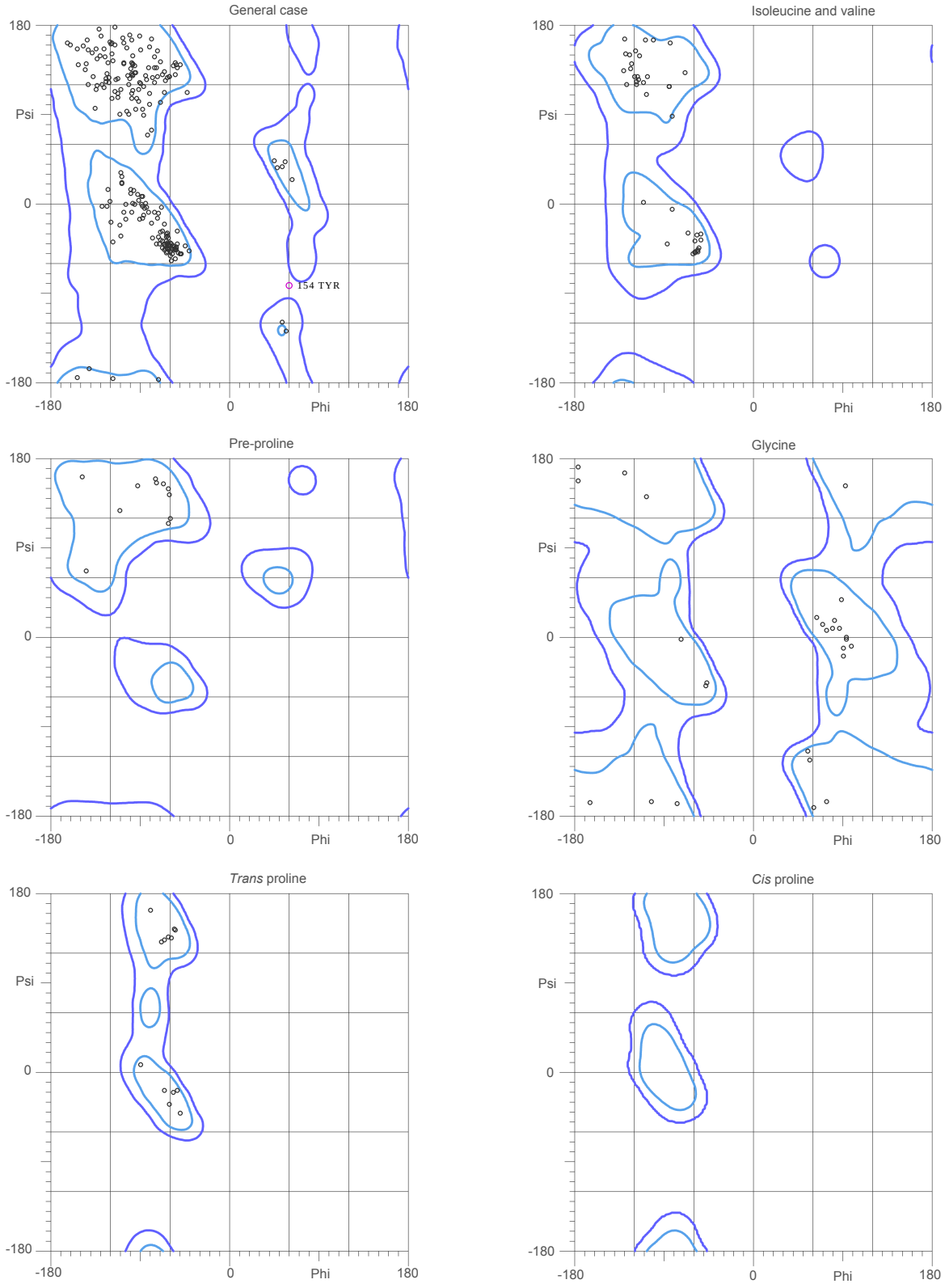
MolProbity Ramachandran analysis

5reb.H.pdb, model 1



98.7% (307/311) of all residues were in favored (98%) regions.
99.7% (310/311) of all residues were in allowed (>99.8%) regions.

There were 1 outliers (phi, psi):
154 TYR (60.3, -82.3)