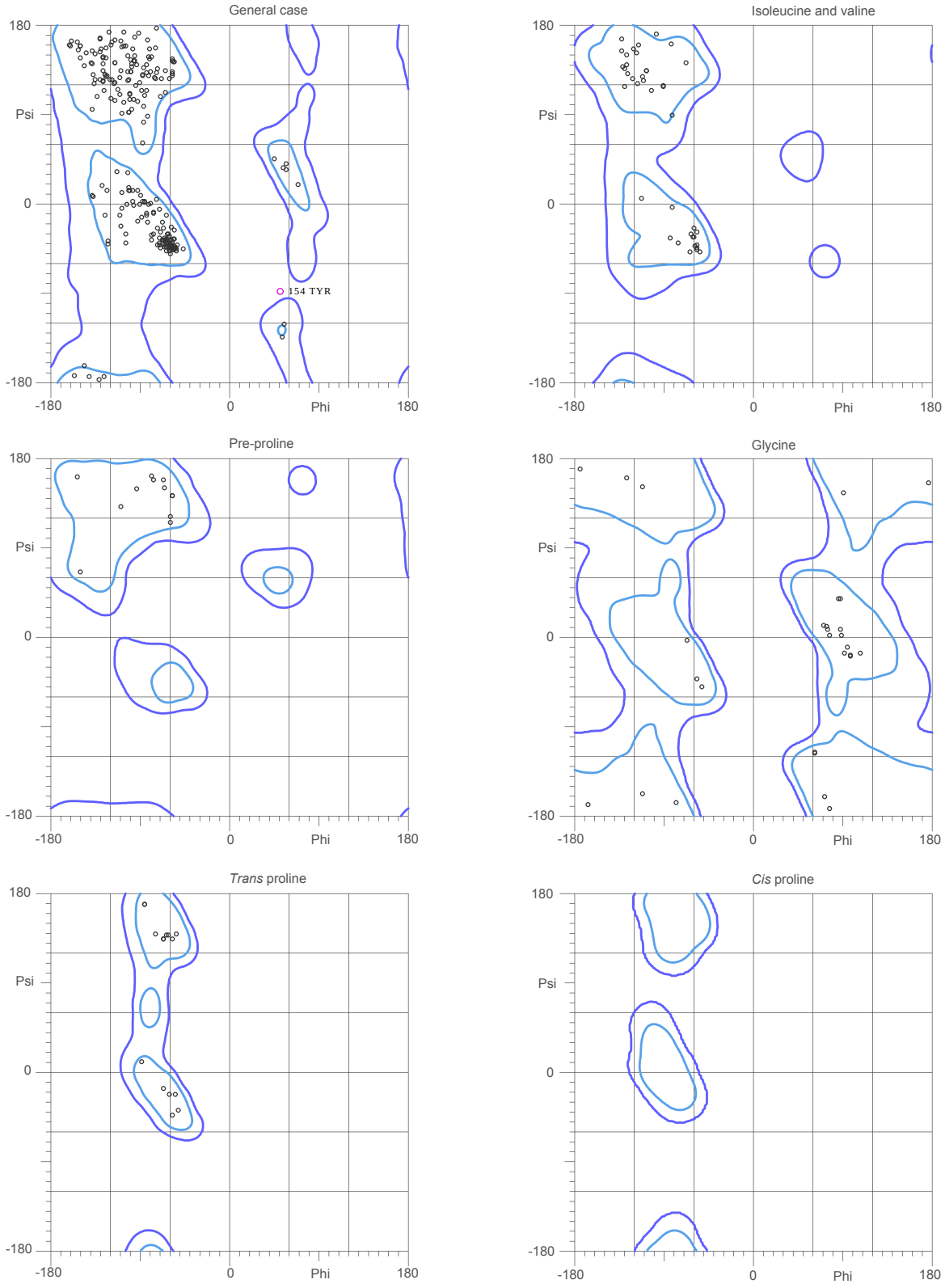
MolProbity Ramachandran analysis

7amj.H.pdb, model 1



98.5% (327/332) of all residues were in favored (98%) regions.
99.7% (331/332) of all residues were in allowed (>99.8%) regions.

There were 1 outliers (phi, psi):
154 TYR (51.2, -88.1)