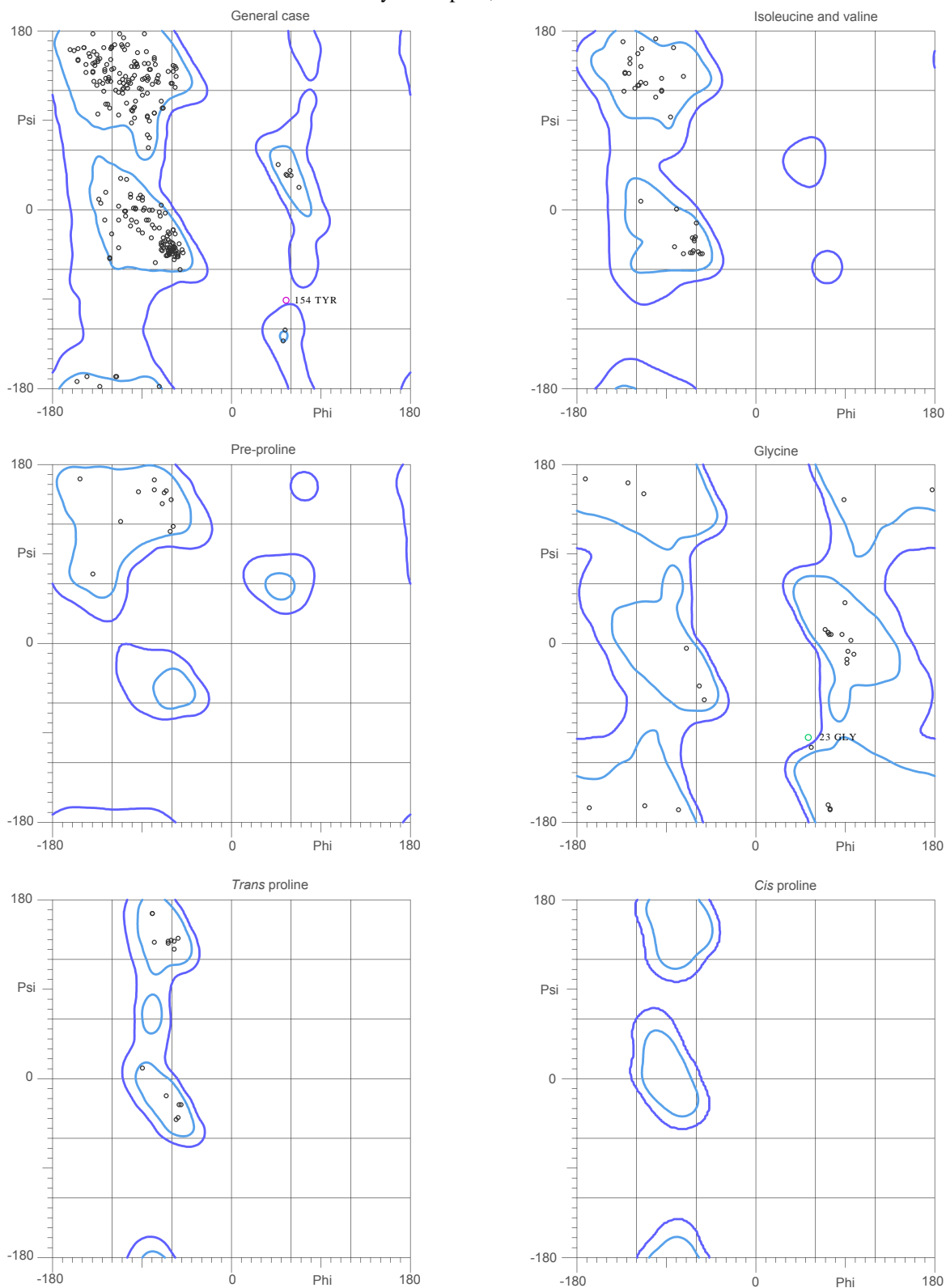


MolProbity Ramachandran analysis

6yvf.H.pdb, model 1



96.8% (329/340) of all residues were in favored (98%) regions.

99.4% (338/340) of all residues were in allowed (>99.8%) regions.

There were 2 outliers (phi, psi):

23 GLY (53.1, -95.8)

154 TYR (55.2, -91.5)