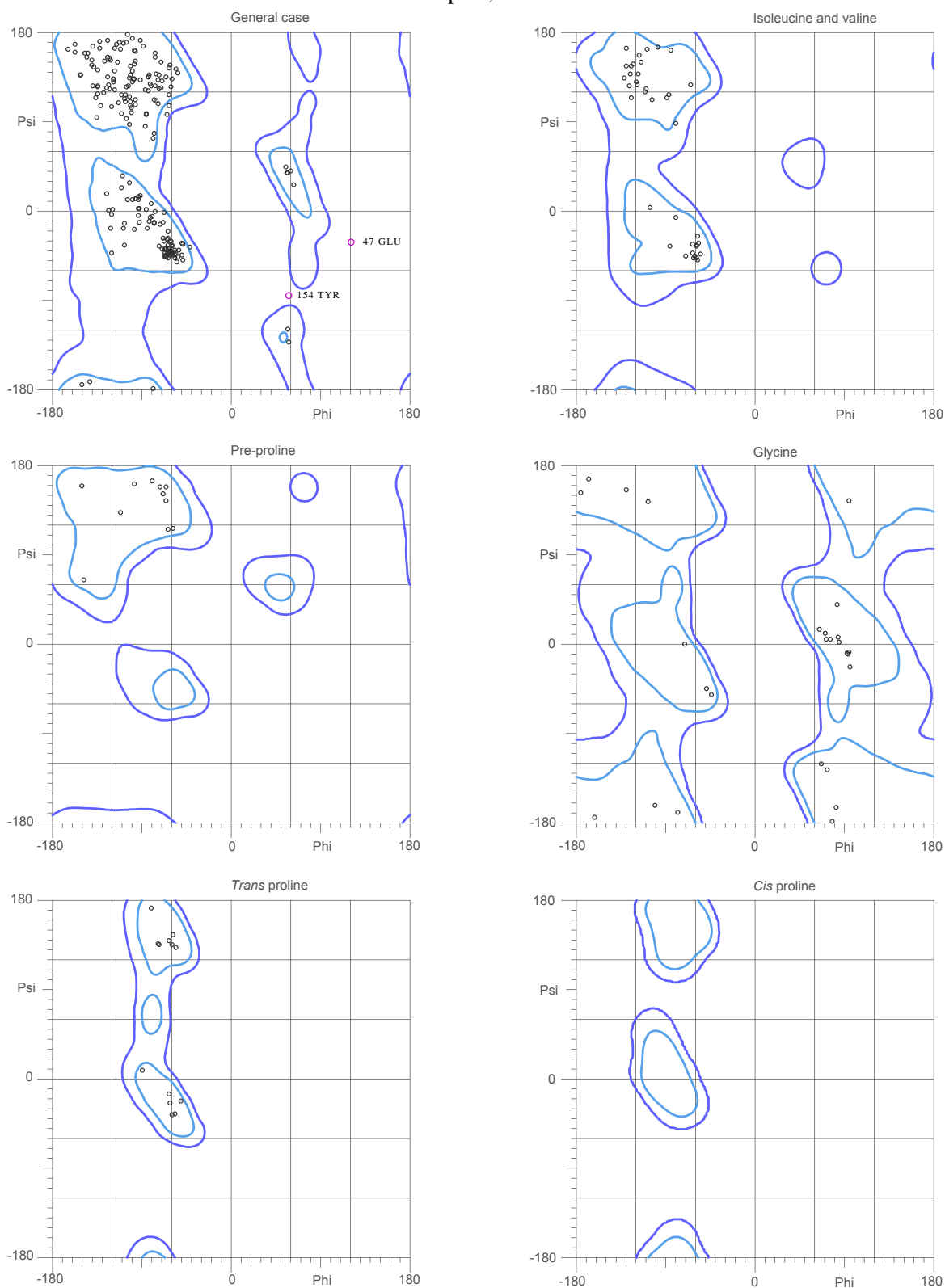


MolProbity Ramachandran analysis

5r11.H.pdb, model 1



97.4% (296/304) of all residues were in favored (98%) regions.

99.3% (302/304) of all residues were in allowed (>99.8%) regions.

There were 2 outliers (phi, psi):

47 GLU (121.3, -31.9)

154 TYR (58.4, -86.0)