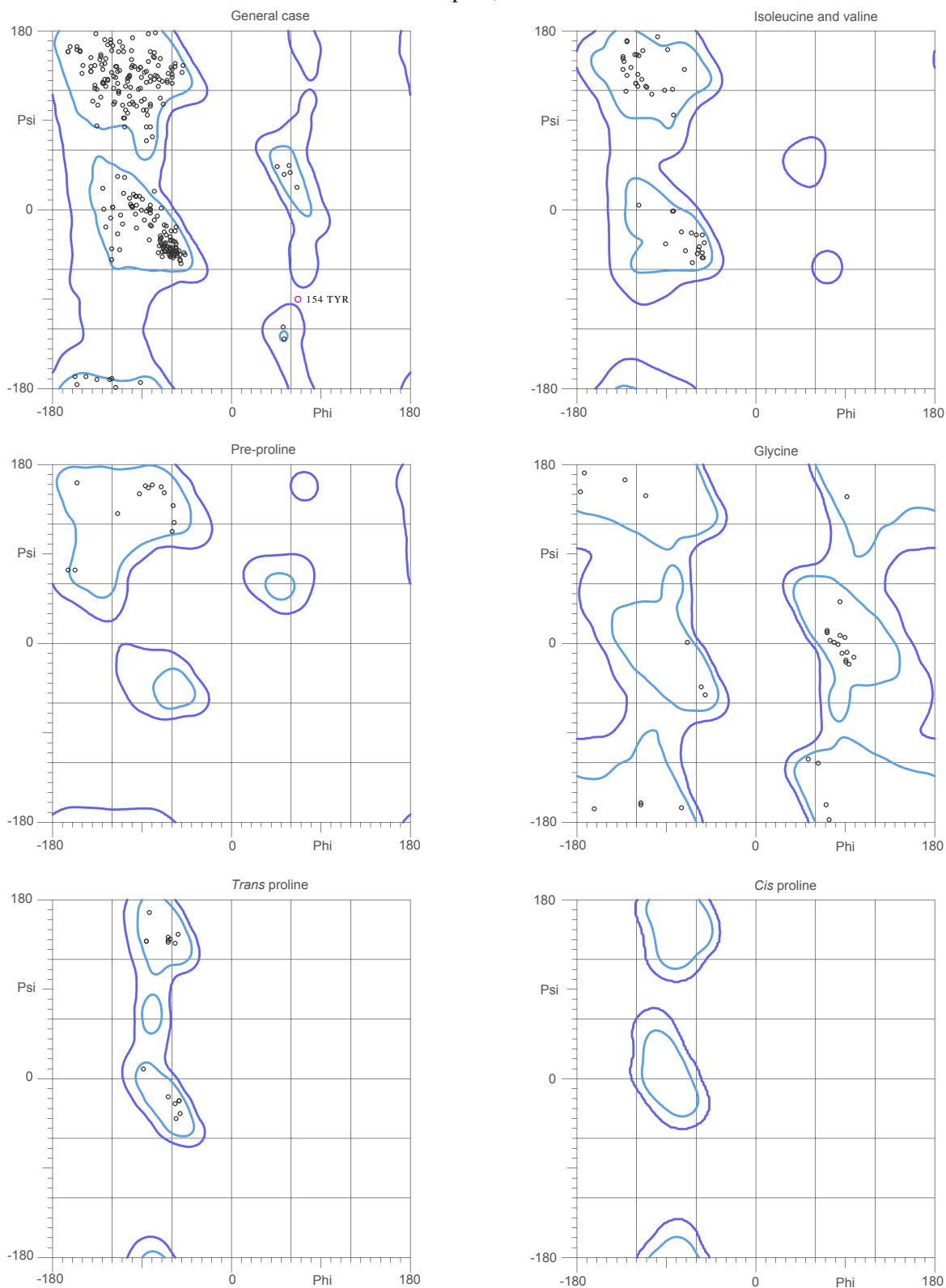


MolProbity Ramachandran analysis

7aws.H.pdb, model 1



98.3% (357/363) of all residues were in favored (98%) regions.

99.7% (362/363) of all residues were in allowed (>99.8%) regions.

There were 1 outliers (phi, psi):

154 TYR (67.5, -90.9)