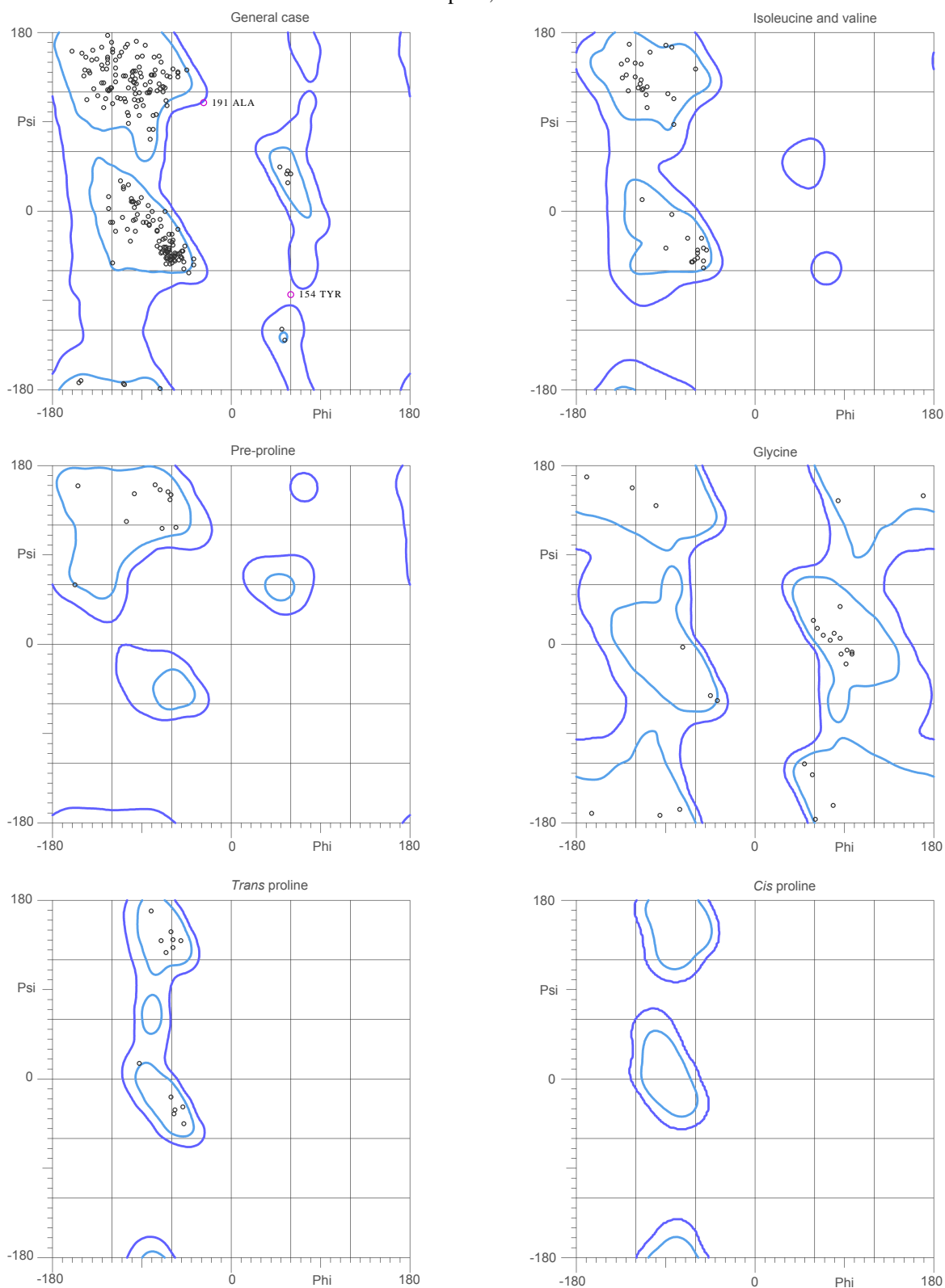


MolProbity Ramachandran analysis

5re8.H.pdb, model 1



96.1% (299/311) of all residues were in favored (98%) regions.

99.4% (309/311) of all residues were in allowed (>99.8%) regions.

There were 2 outliers (phi, psi):

154 TYR (60.9, -84.3)

191 ALA (-28.3, 110.4)