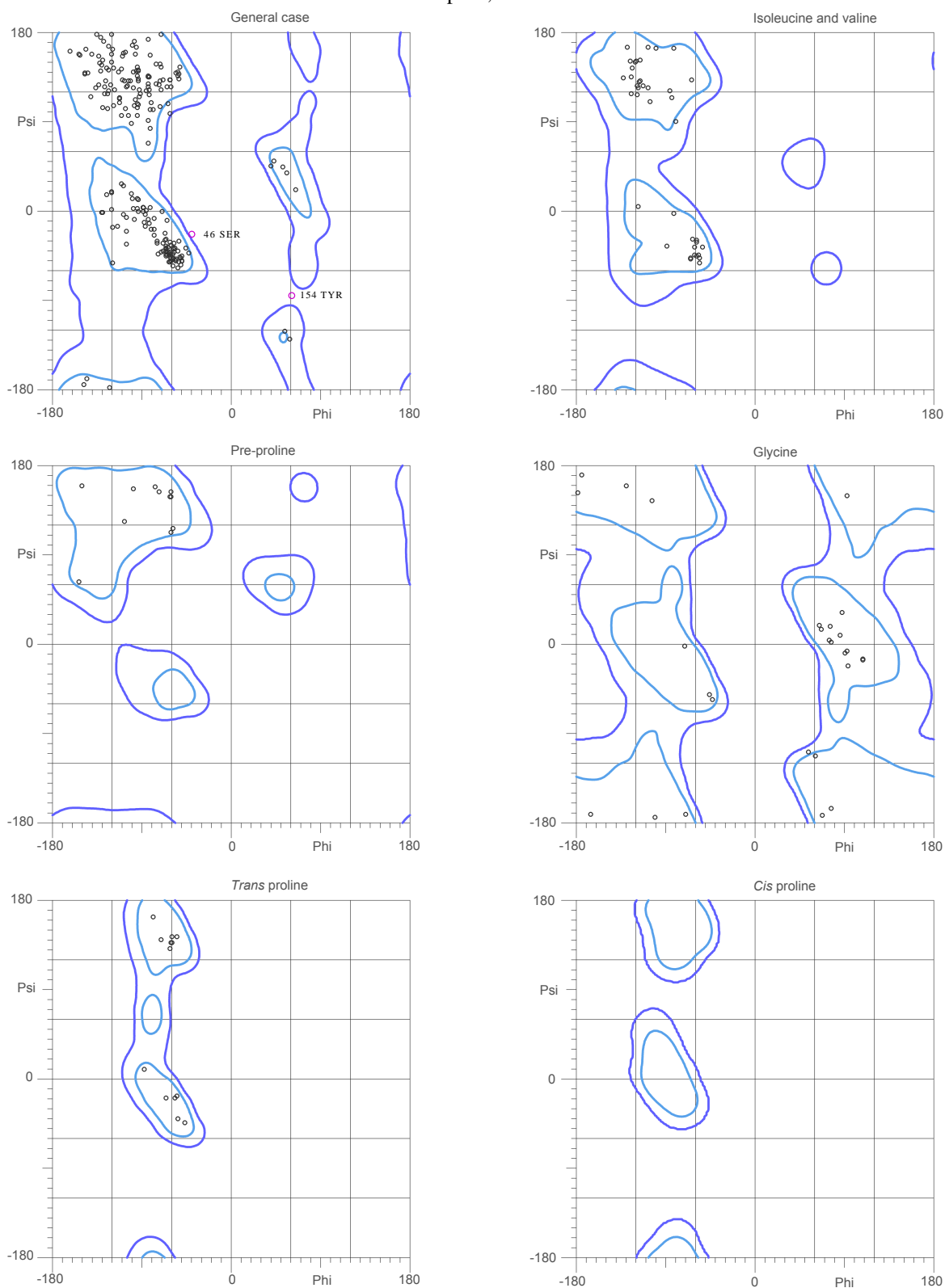


MolProbity Ramachandran analysis

5rei.H.pdb, model 1



97.4% (303/311) 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 (Φ , Ψ):

46 SER (-40.4, -23.9)

154 TYR (61.2, -85.1)