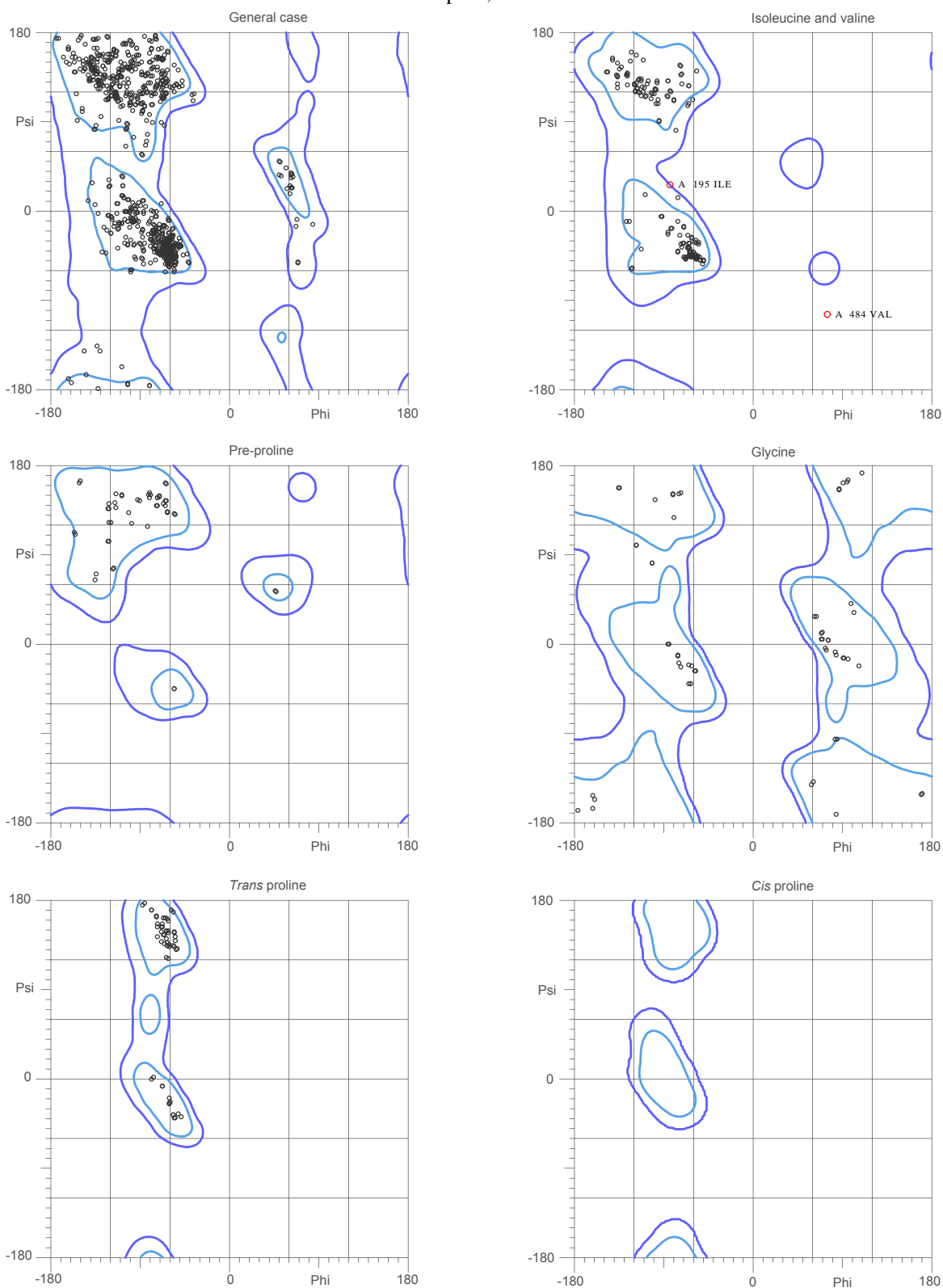


MolProbity Ramachandran analysis

5r19.H.pdb, model 1



96.7% (1108/1146) of all residues were in favored (98%) regions.

99.8% (1144/1146) of all residues were in allowed (>99.8%) regions.

There were 2 outliers (Φ , Ψ):

A 195 ILE (-84.1, 27.9)

A 484 VAL (75.4, -104.2)