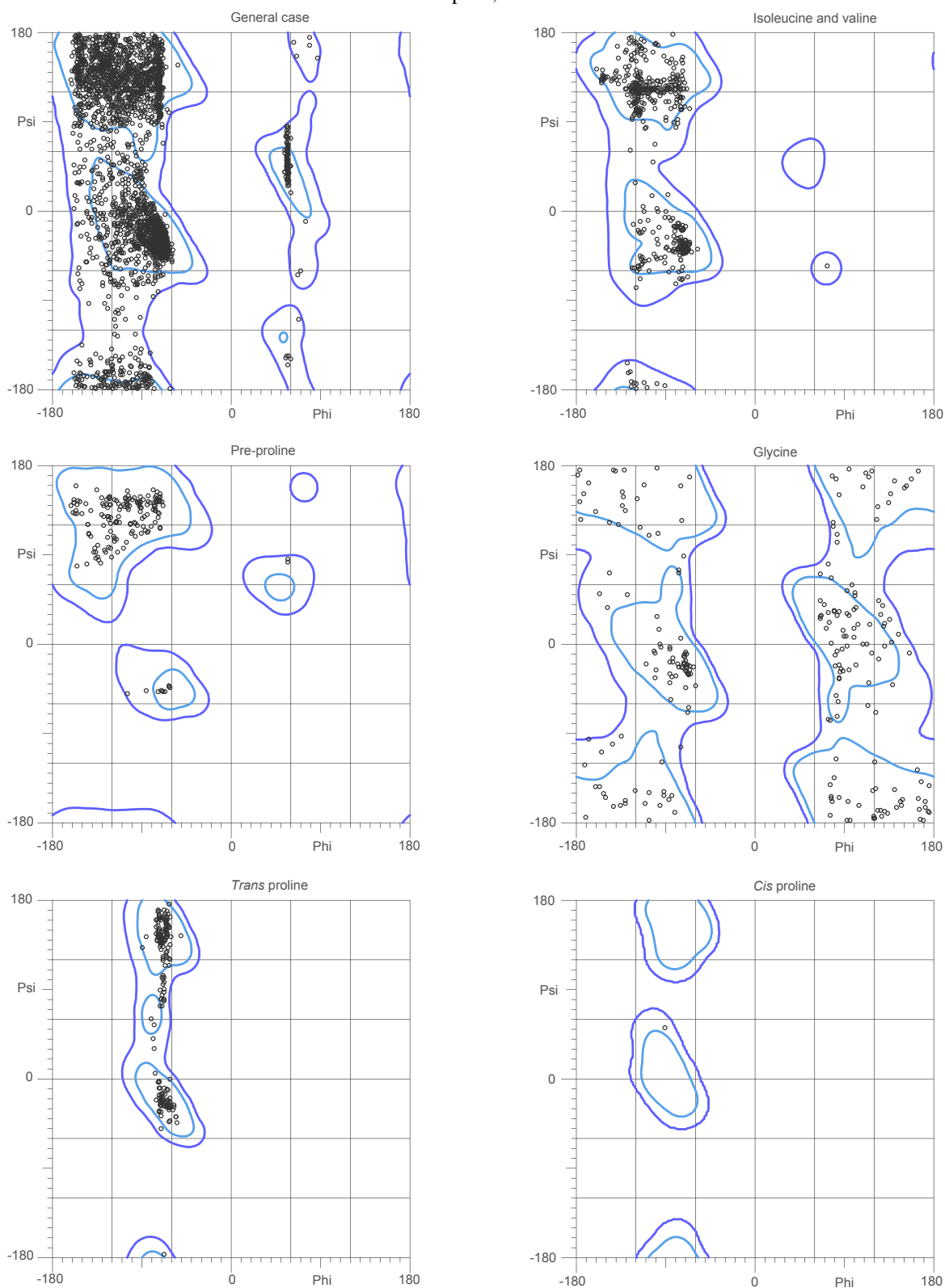


MolProbity Ramachandran analysis

7a95.H.pdb, model 1



89.0% (3340/3754) of all residues were in favored (98%) regions.

100.0% (3754/3754) of all residues were in allowed (>99.8%) regions.

There were no outliers.