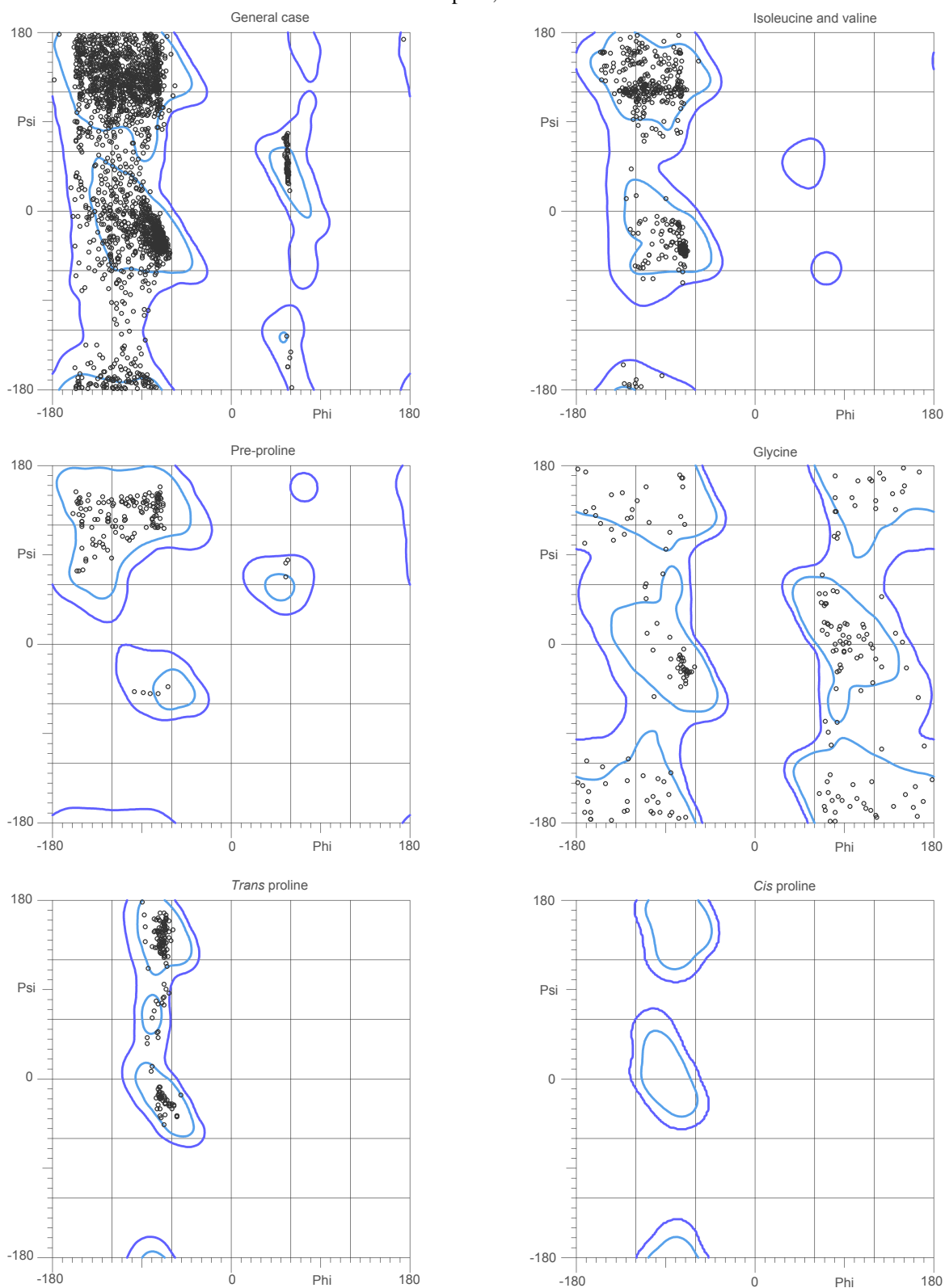


MolProbity Ramachandran analysis

7a93.H.pdb, model 1



90.1% (2870/3185) of all residues were in favored (98%) regions.

100.0% (3185/3185) of all residues were in allowed (>99.8%) regions.

There were no outliers.