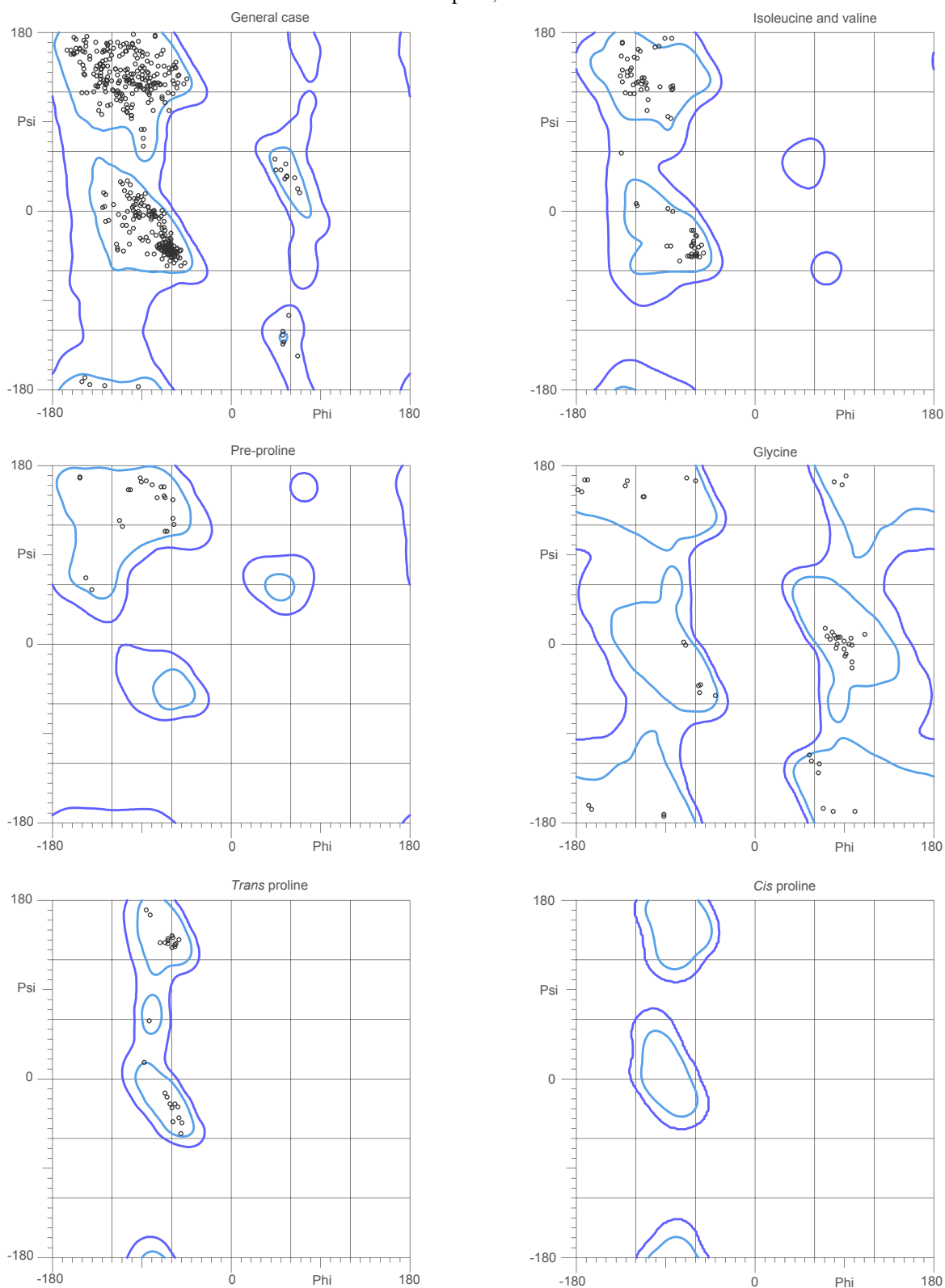


MolProbity Ramachandran analysis

7b3e.H.pdb, model 1



98.2% (594/605) of all residues were in favored (98%) regions.

100.0% (605/605) of all residues were in allowed (>99.8%) regions.

There were no outliers.