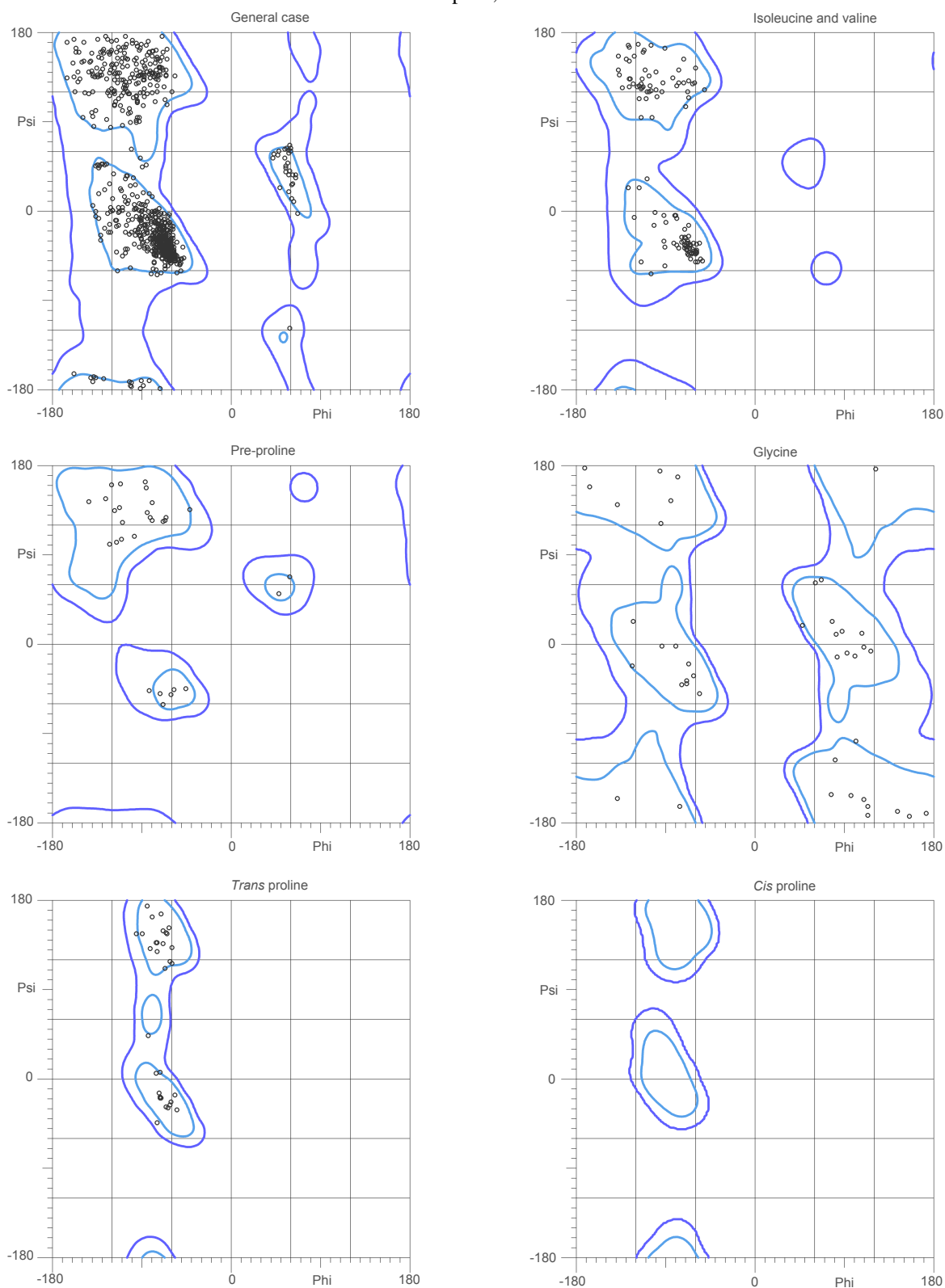


MolProbity Ramachandran analysis

7bv2.H.pdb, model 1



96.8% (962/994) of all residues were in favored (98%) regions.

100.0% (994/994) of all residues were in allowed (>99.8%) regions.

There were no outliers.