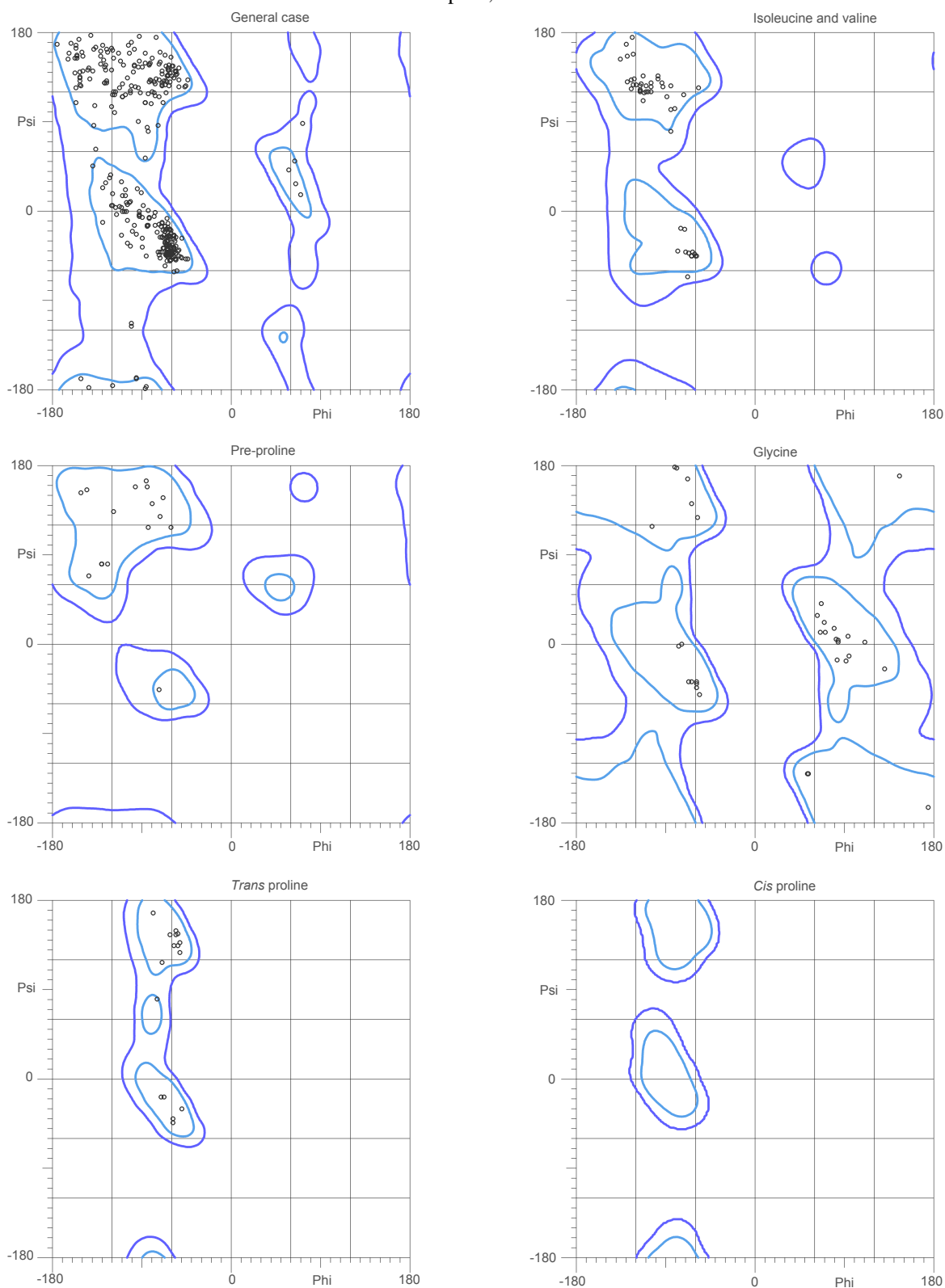


MolProbity Ramachandran analysis

716r.H.pdb, model 1



97.8% (440/450) of all residues were in favored (98%) regions.

100.0% (450/450) of all residues were in allowed (>99.8%) regions.

There were no outliers.