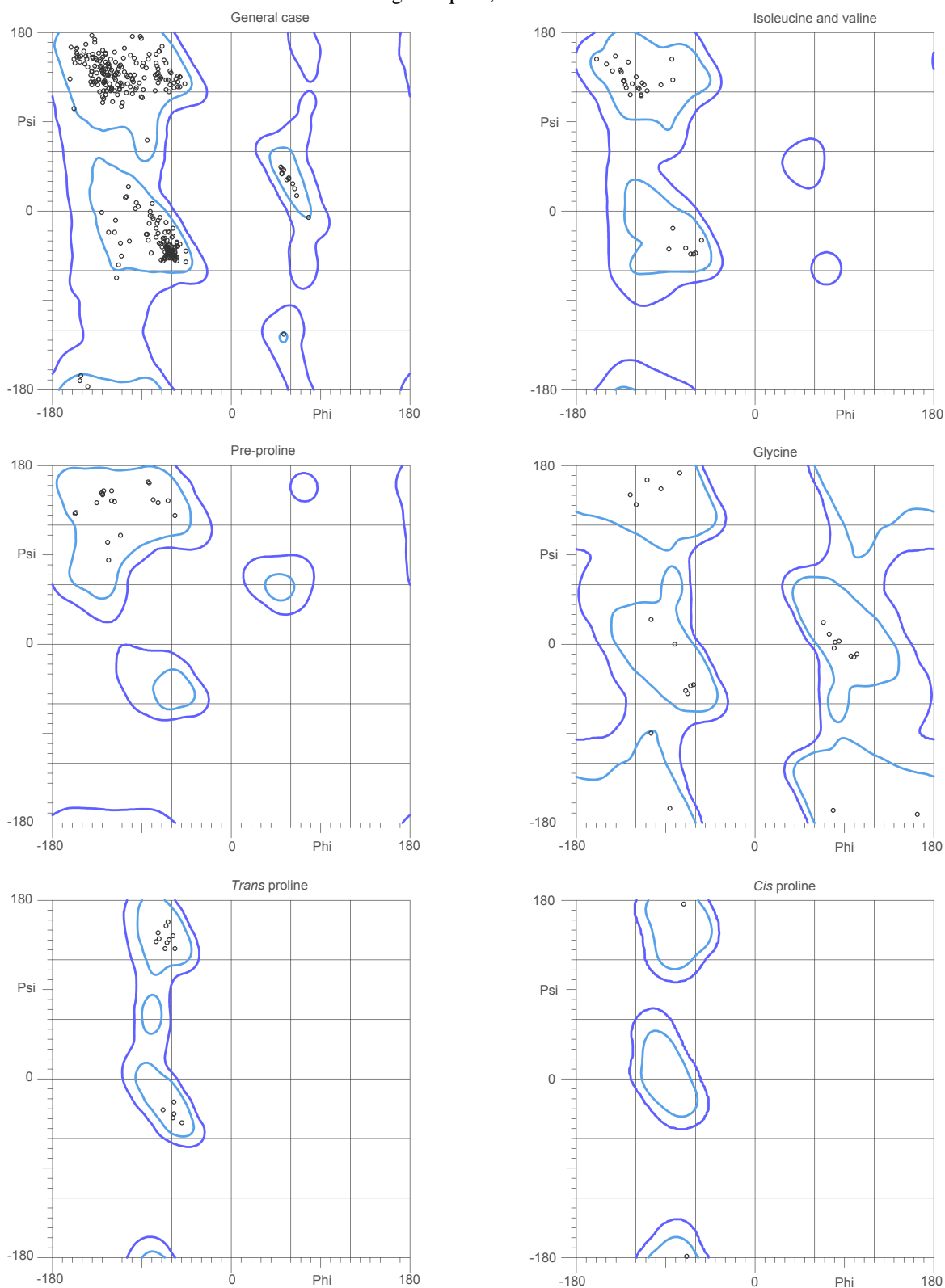


MolProbity Ramachandran analysis

7kgr.H.pdb, model 1



99.2% (391/394) of all residues were in favored (98%) regions.

100.0% (394/394) of all residues were in allowed (>99.8%) regions.

There were no outliers.