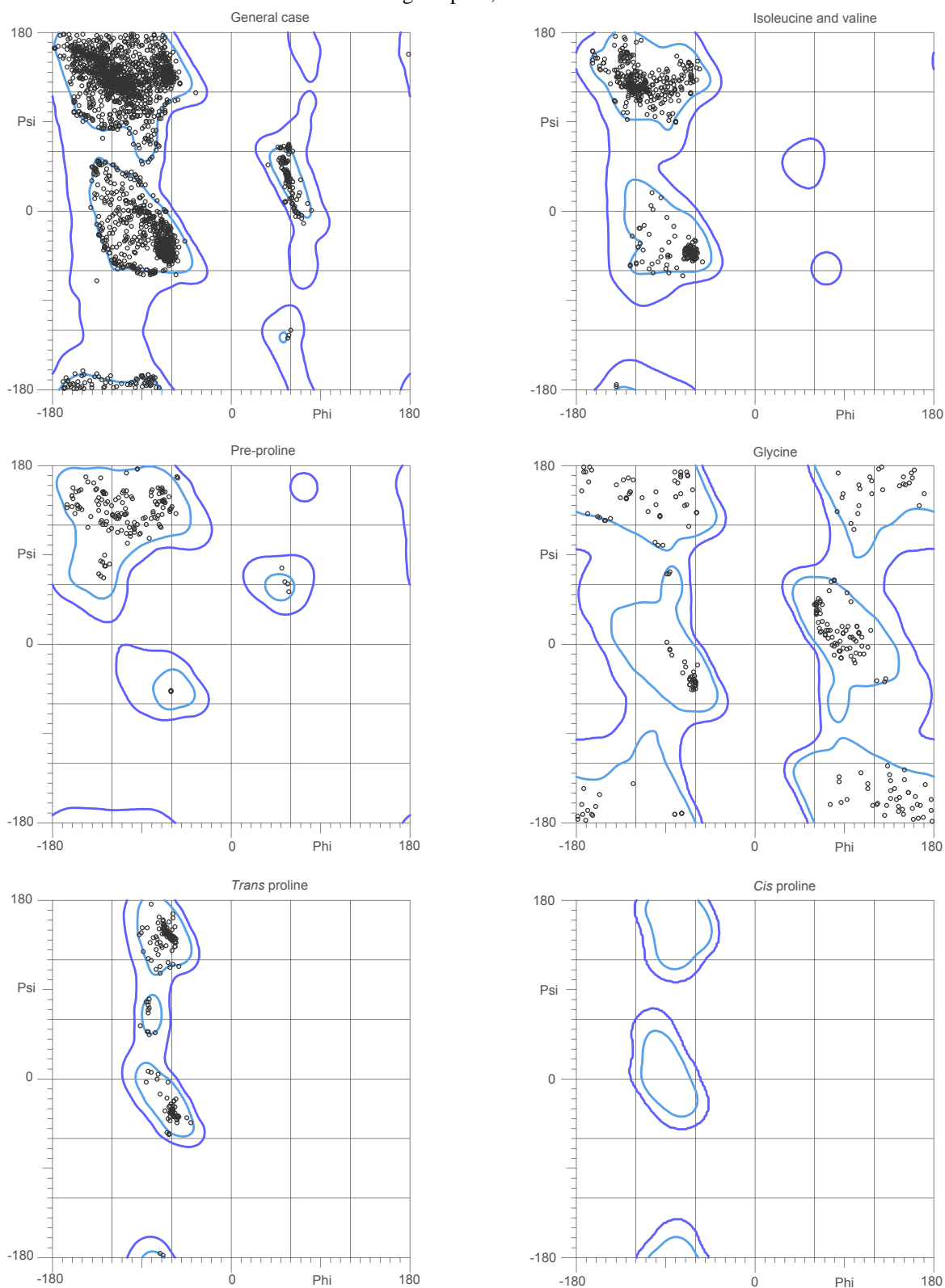


MolProbity Ramachandran analysis

7ksg.H.pdb, model 1



95.2% (3344/3514) of all residues were in favored (98%) regions.

100.0% (3514/3514) of all residues were in allowed (>99.8%) regions.

There were no outliers.