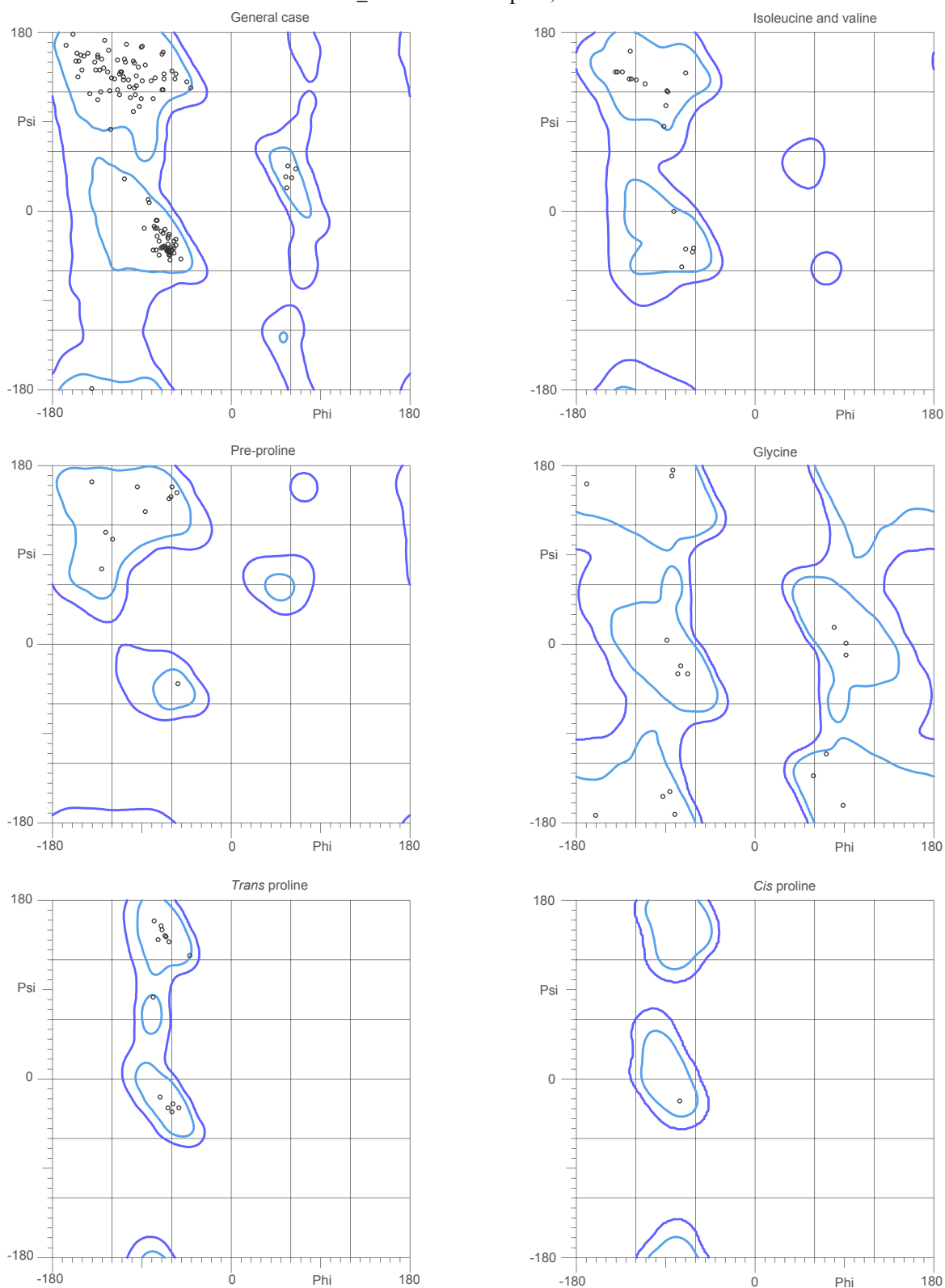


MolProbity Ramachandran analysis

1a44_.B99990008.pdb, model 1



100,0% (183/183) of all residues were in favored (98%) regions.

100,0% (183/183) of all residues were in allowed (>99.8%) regions.

There were no outliers.