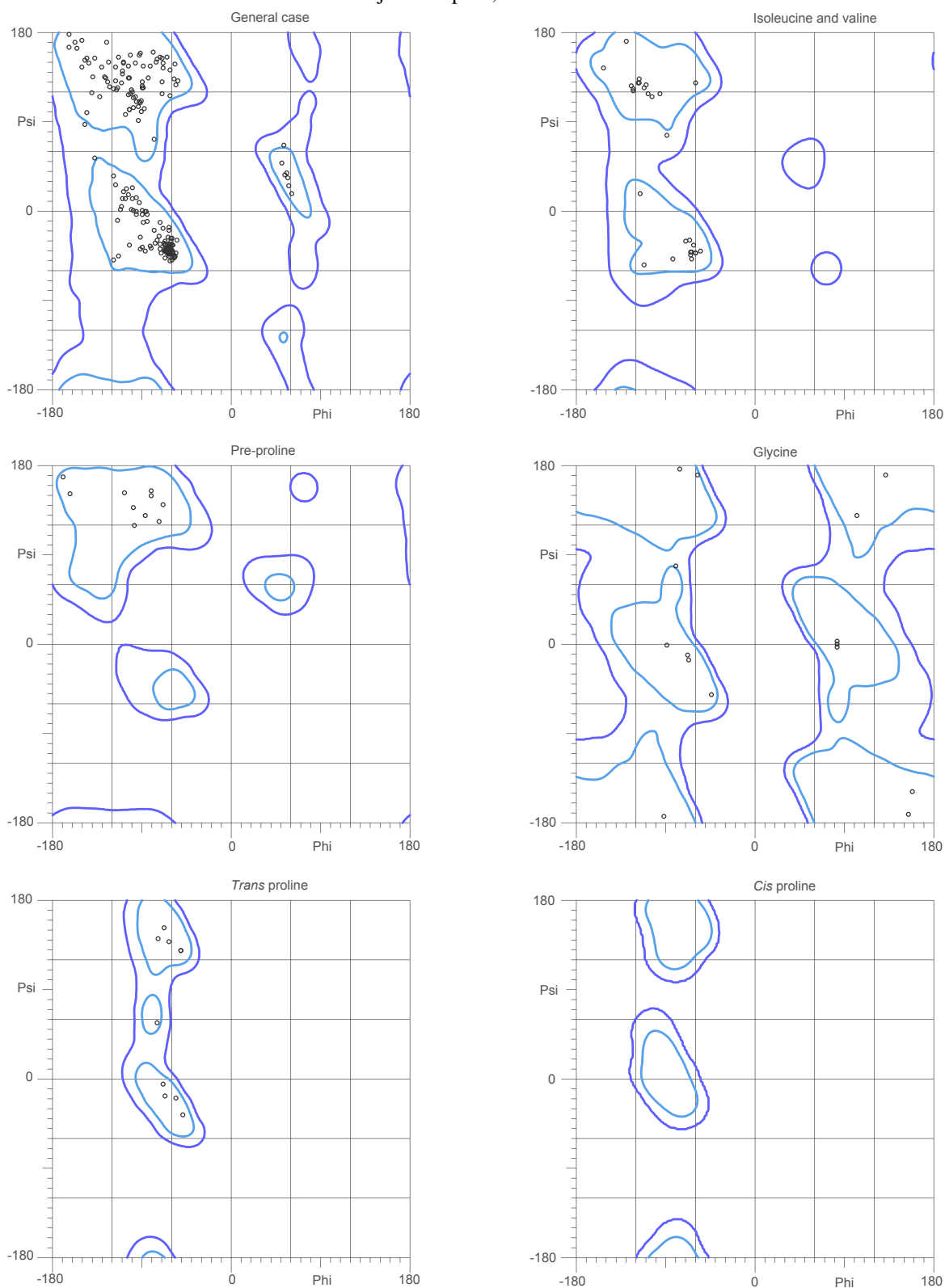


# MolProbity Ramachandran analysis

7jzm.H.pdb, model 1



98.0% (249/254) of all residues were in favored (98%) regions.

100.0% (254/254) of all residues were in allowed (>99.8%) regions.

There were no outliers.