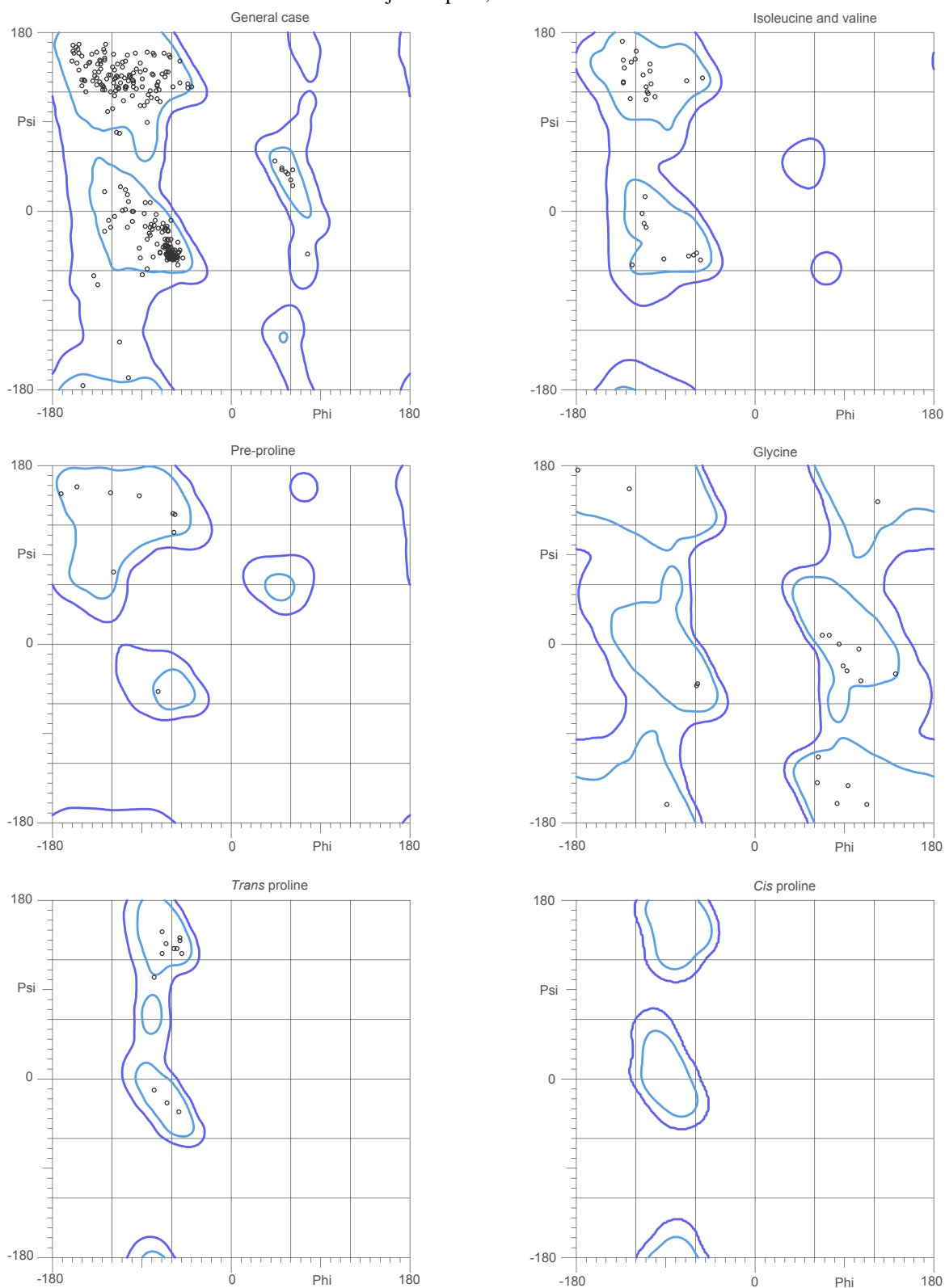


# MolProbity Ramachandran analysis

7jit.H.pdb, model 1



97.2% (315/324) of all residues were in favored (98%) regions.

100.0% (324/324) of all residues were in allowed (>99.8%) regions.

There were no outliers.