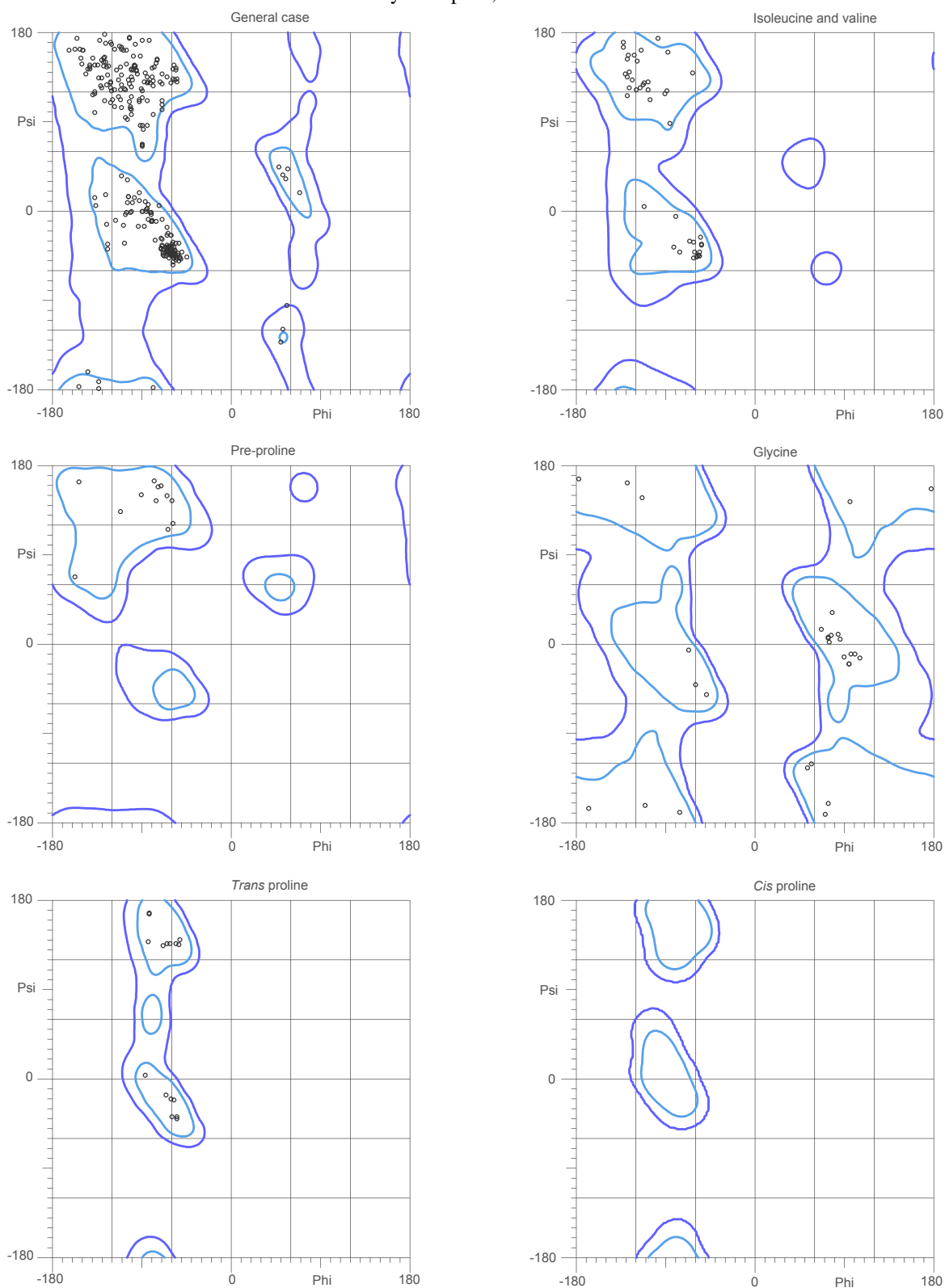


# MolProbity Ramachandran analysis

7ay7.H.pdb, model 1



98.6% (355/360) of all residues were in favored (98%) regions.

100.0% (360/360) of all residues were in allowed (>99.8%) regions.

There were no outliers.