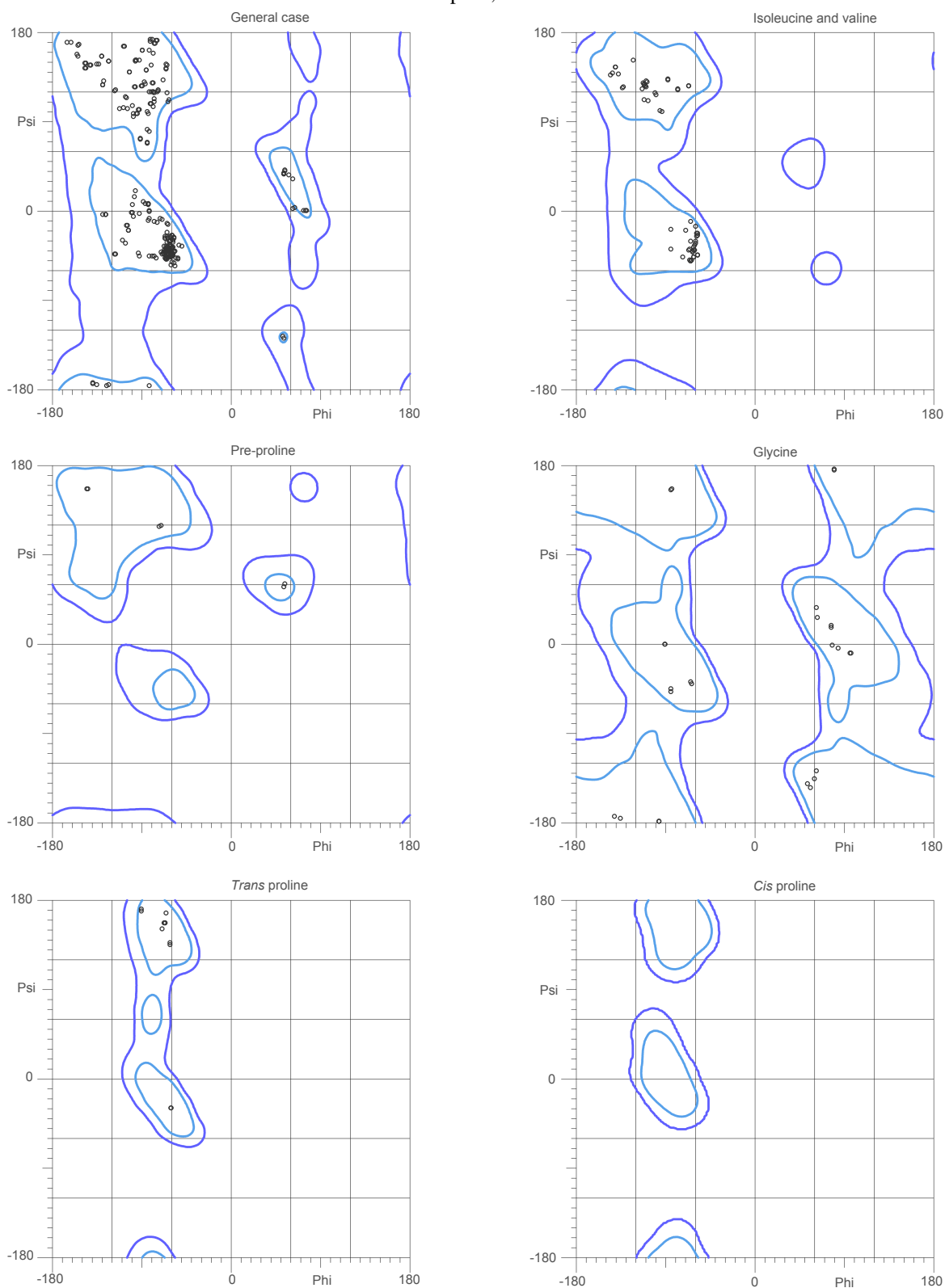


# MolProbity Ramachandran analysis

5s46.H.pdb, model 1



99.7% (351/352) of all residues were in favored (98%) regions.

100.0% (352/352) of all residues were in allowed (>99.8%) regions.

There were no outliers.