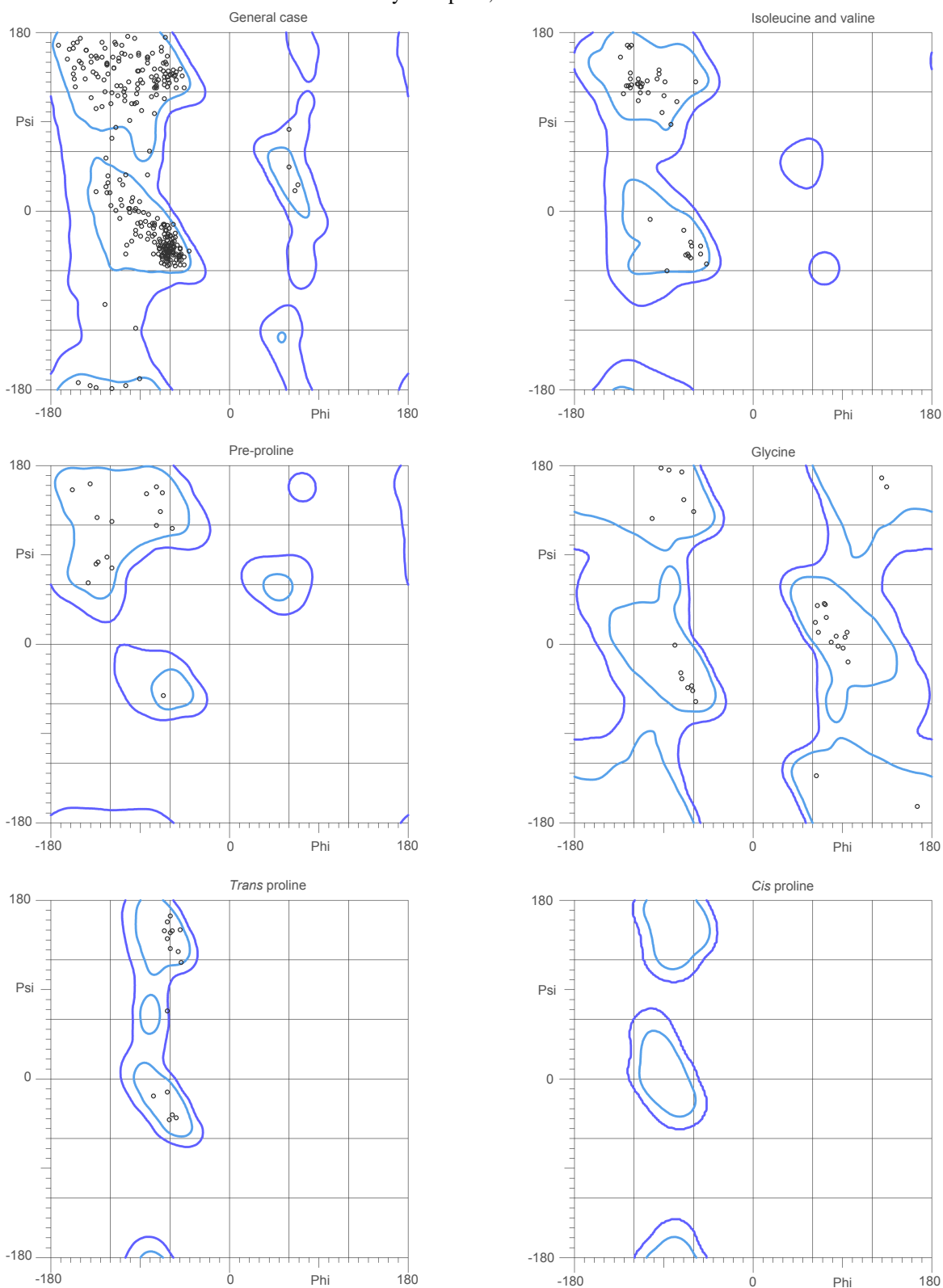


# MolProbity Ramachandran analysis

2xyr.H.pdb, model 1



97.7% (386/395) of all residues were in favored (98%) regions.

100.0% (395/395) of all residues were in allowed (>99.8%) regions.

There were no outliers.