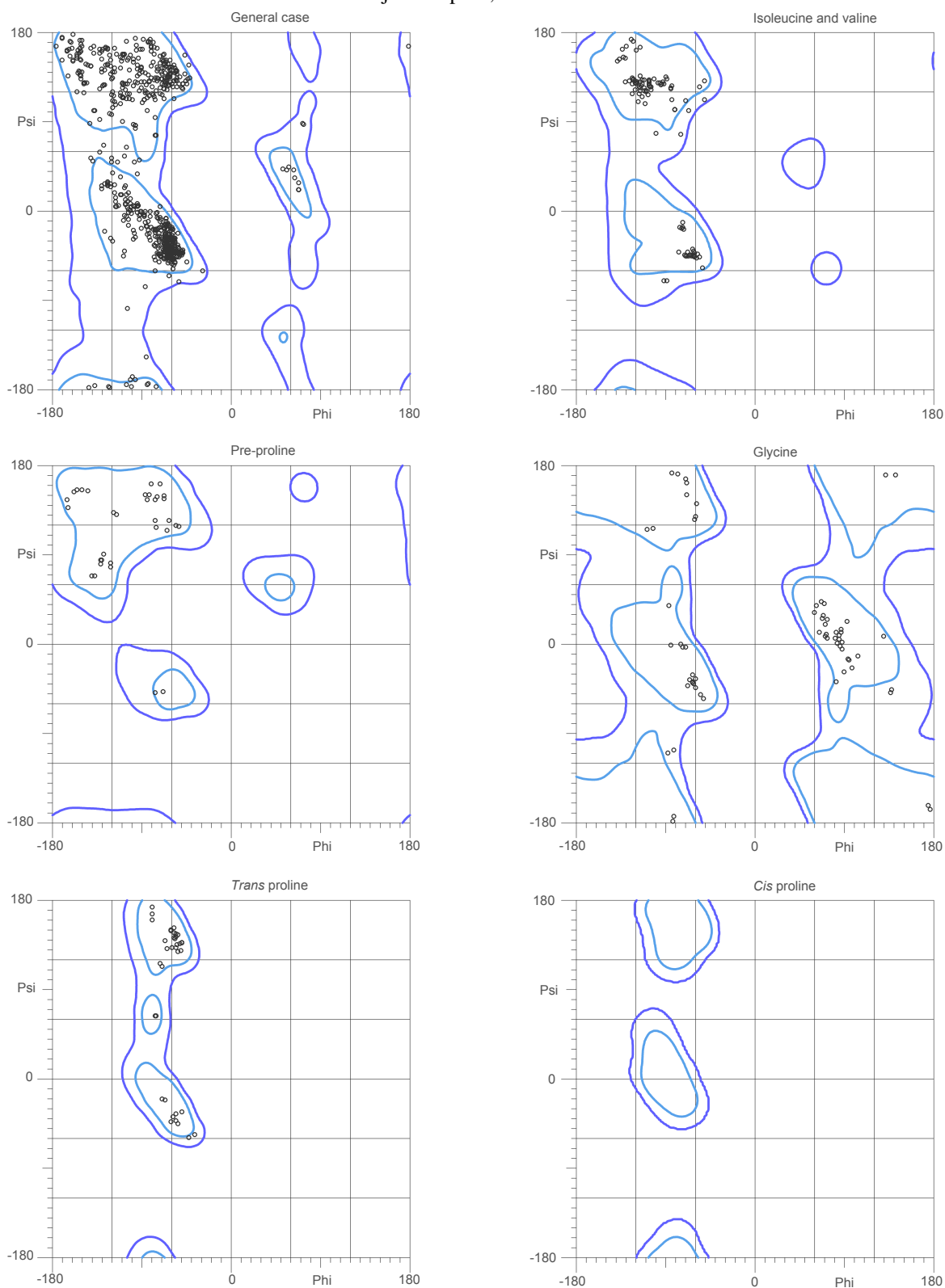


# MolProbity Ramachandran analysis

7jz0.H.pdb, model 1



95.8% (835/872) of all residues were in favored (98%) regions.

100.0% (872/872) of all residues were in allowed (>99.8%) regions.

There were no outliers.