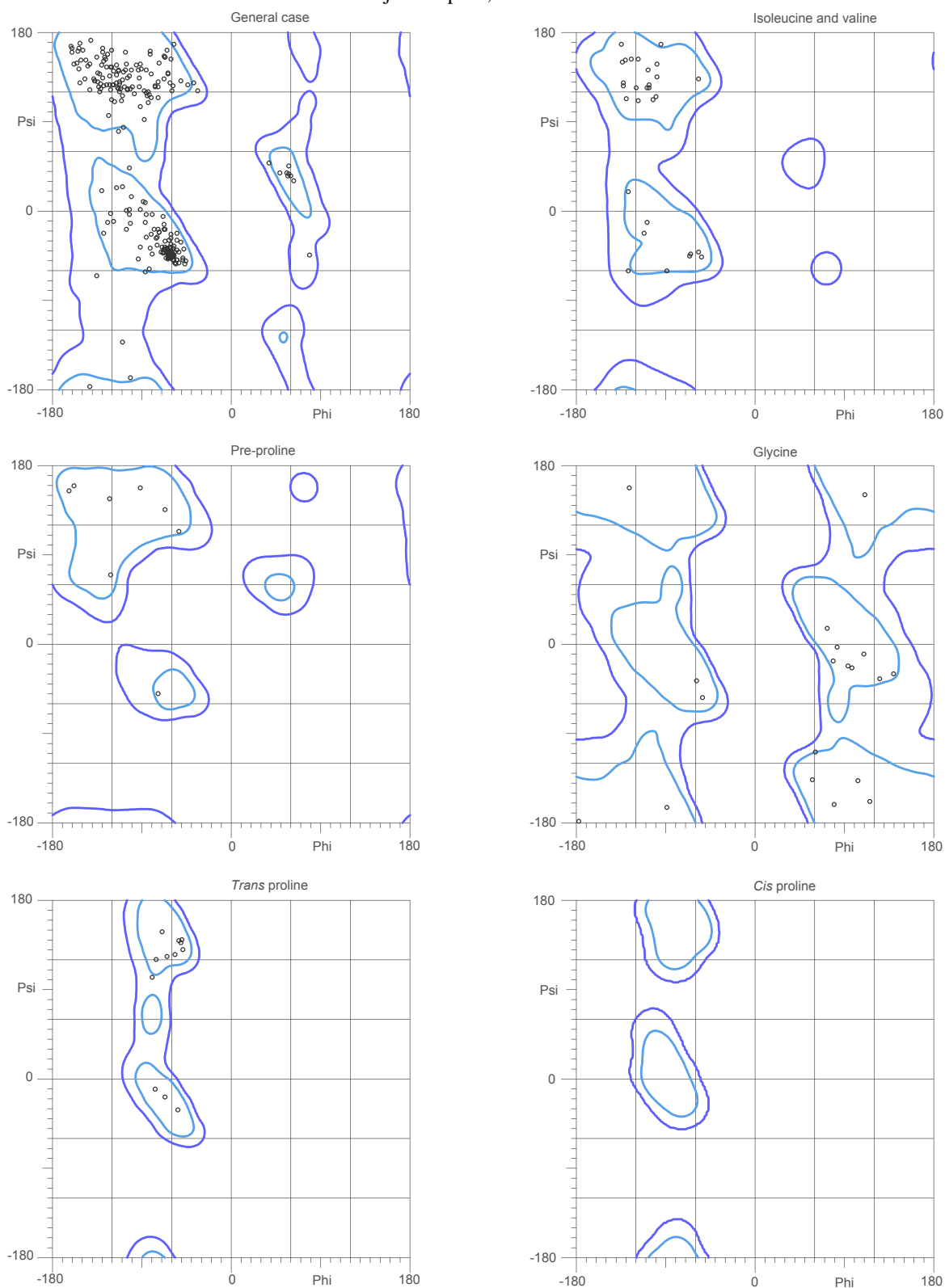


# MolProbity Ramachandran analysis

7jiv.H.pdb, model 1



95.5% (299/313) of all residues were in favored (98%) regions.

100.0% (313/313) of all residues were in allowed (>99.8%) regions.

There were no outliers.