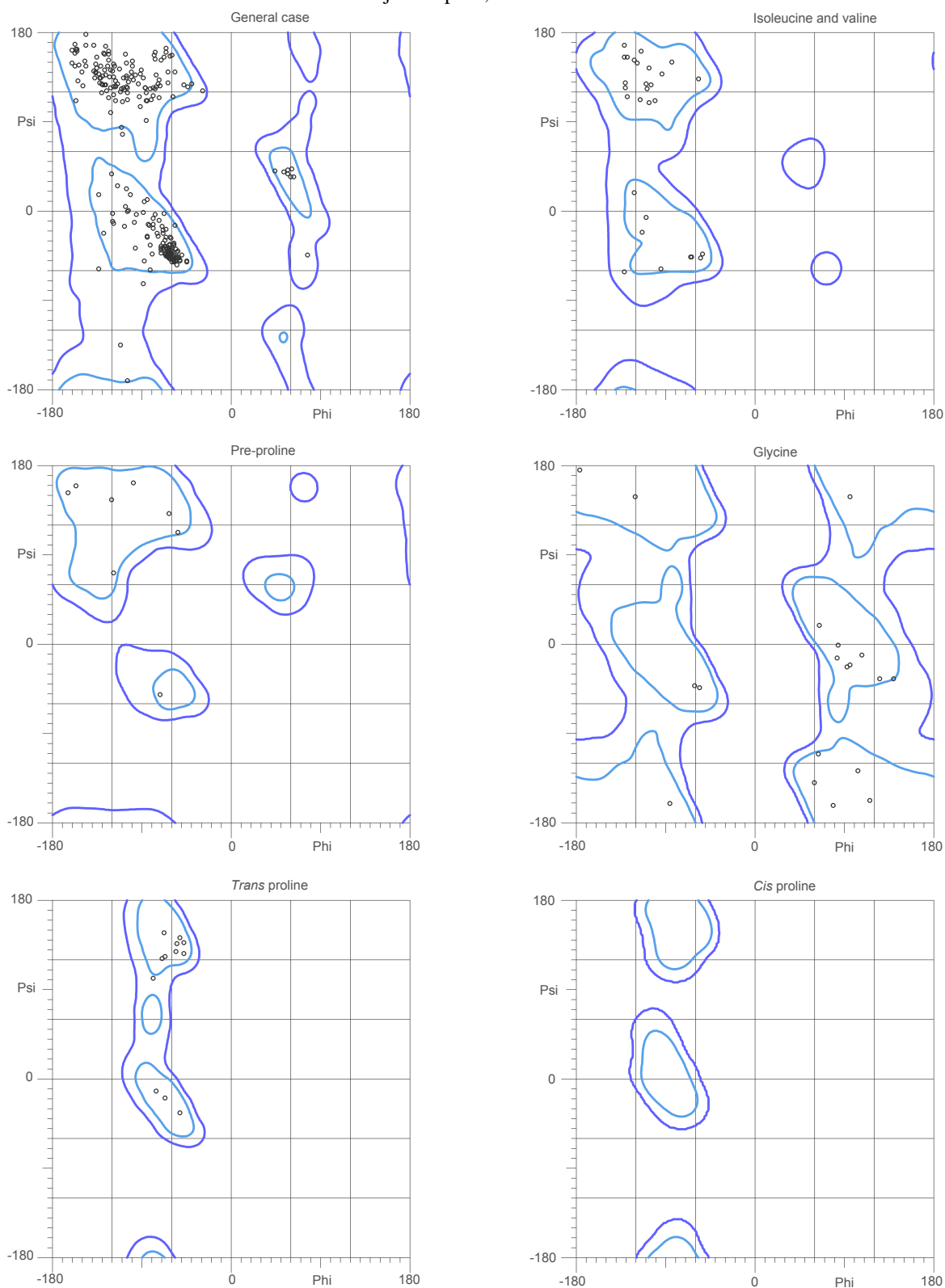


# MolProbity Ramachandran analysis

7jir.H.pdb, model 1



96.8% (303/313) 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.