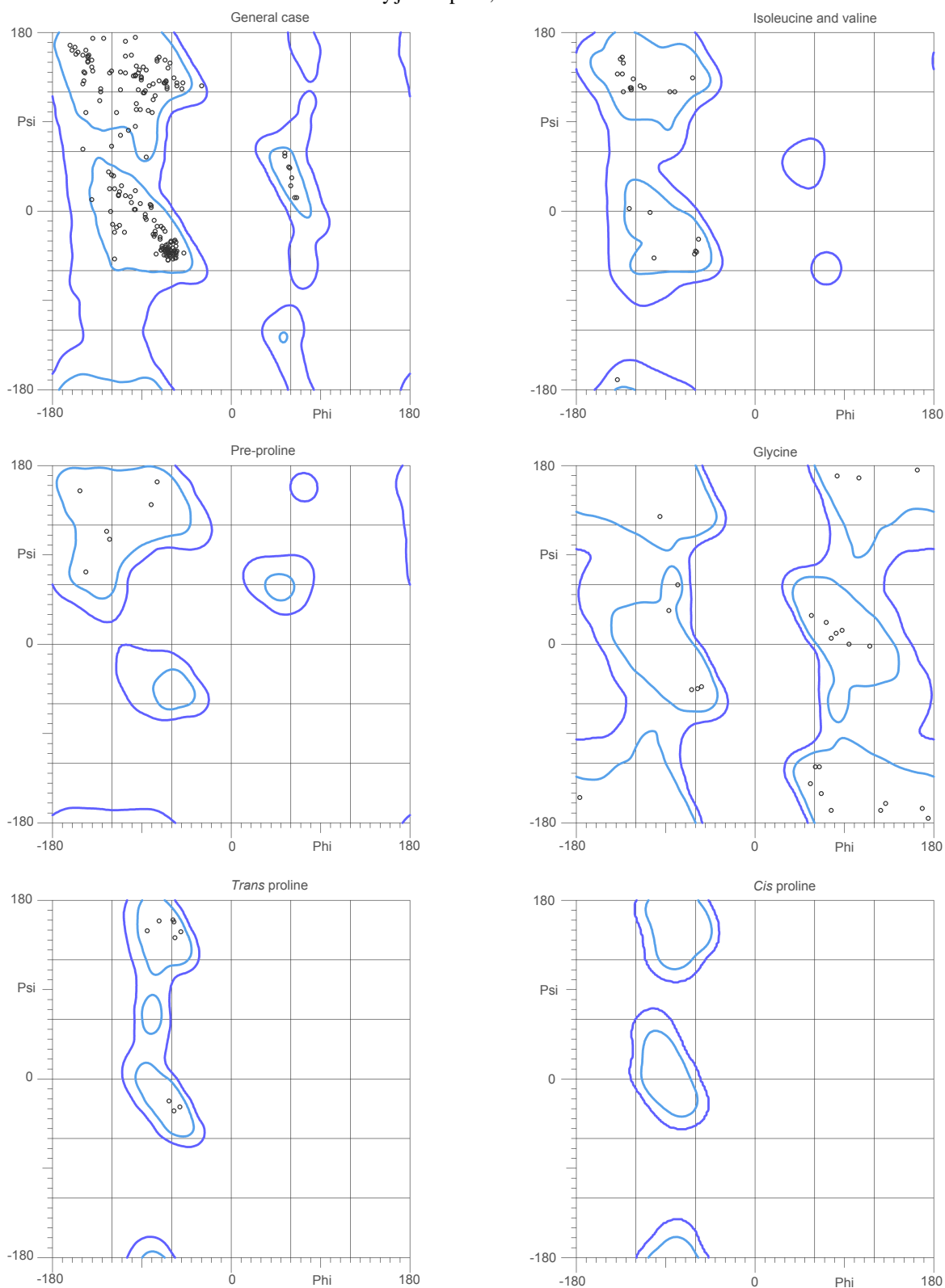


# MolProbity Ramachandran analysis

2yjc.H.pdb, model 1



97.6% (245/251) of all residues were in favored (98%) regions.

100.0% (251/251) of all residues were in allowed (>99.8%) regions.

There were no outliers.