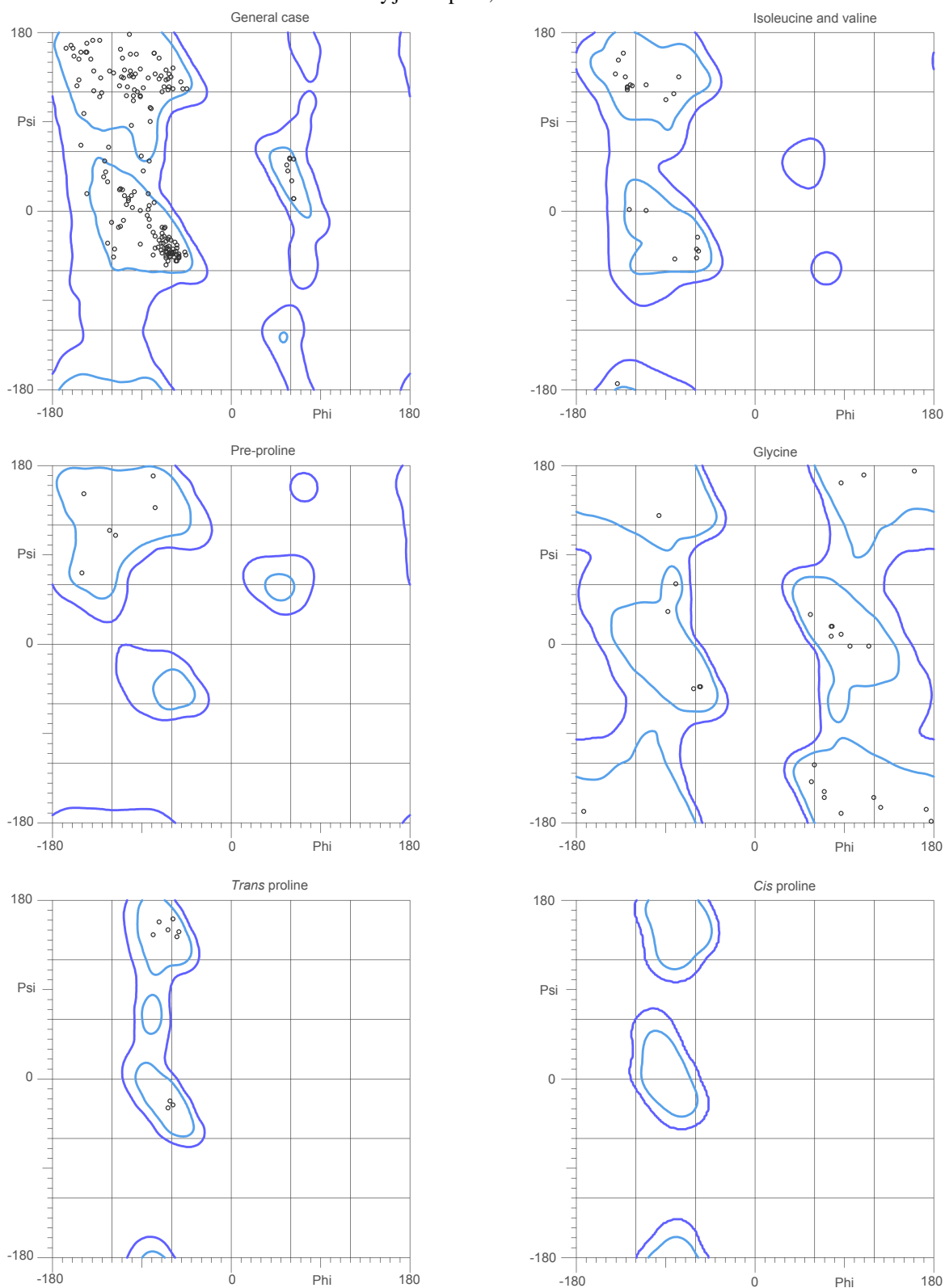


# MolProbity Ramachandran analysis

2yjb.H.pdb, model 1



97.0% (226/233) of all residues were in favored (98%) regions.

100.0% (233/233) of all residues were in allowed (>99.8%) regions.

There were no outliers.