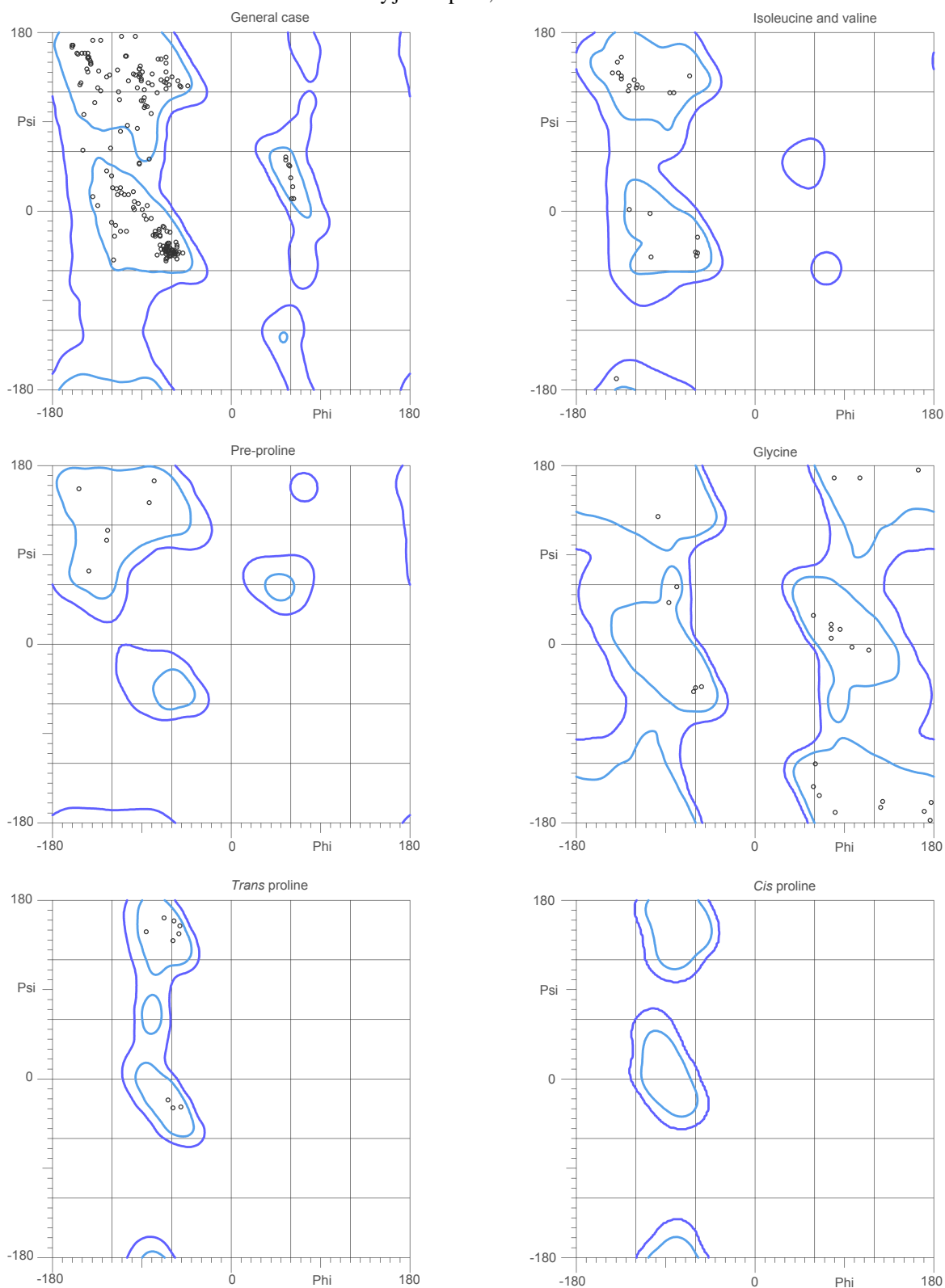


# MolProbity Ramachandran analysis

2yj2.H.pdb, model 1



97.5% (237/243) of all residues were in favored (98%) regions.

100.0% (243/243) of all residues were in allowed (>99.8%) regions.

There were no outliers.