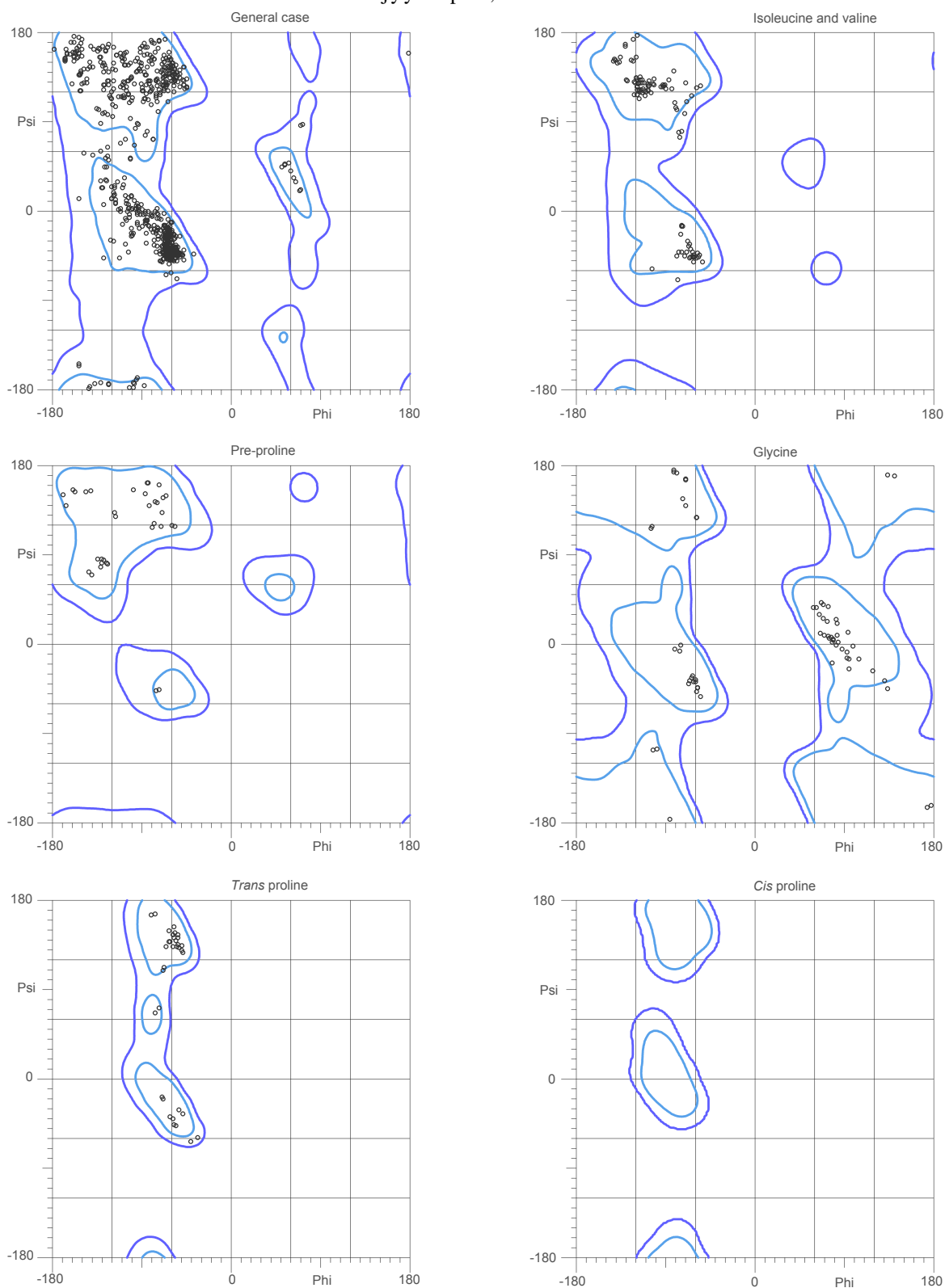


# MolProbity Ramachandran analysis

7jyy.H.pdb, model 1



96.3% (863/896) of all residues were in favored (98%) regions.

100.0% (896/896) of all residues were in allowed (>99.8%) regions.

There were no outliers.