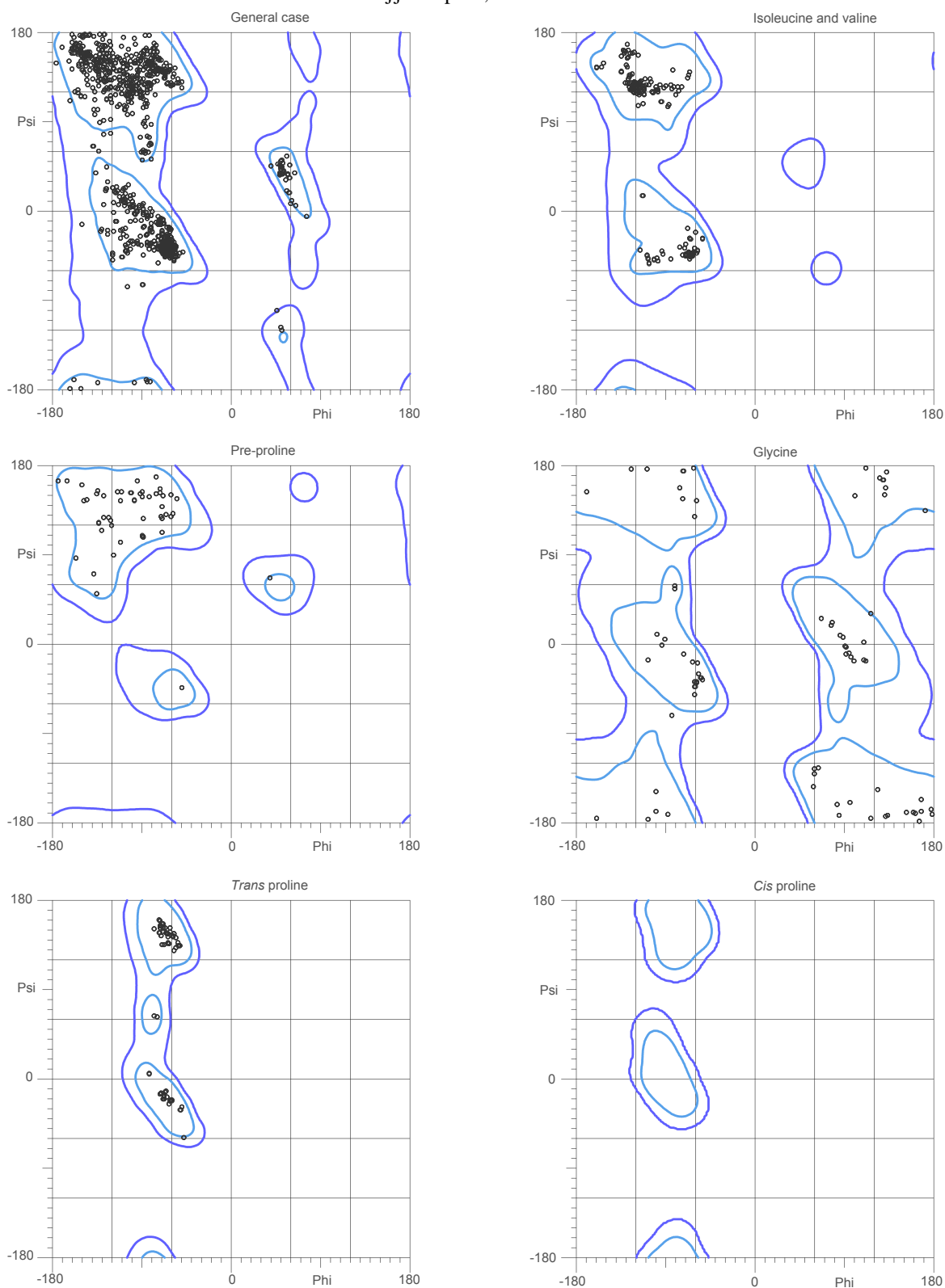


# MolProbity Ramachandran analysis

7jji.H.pdb, model 1



98.4% (3255/3309) of all residues were in favored (98%) regions.

100.0% (3309/3309) of all residues were in allowed (>99.8%) regions.

There were no outliers.