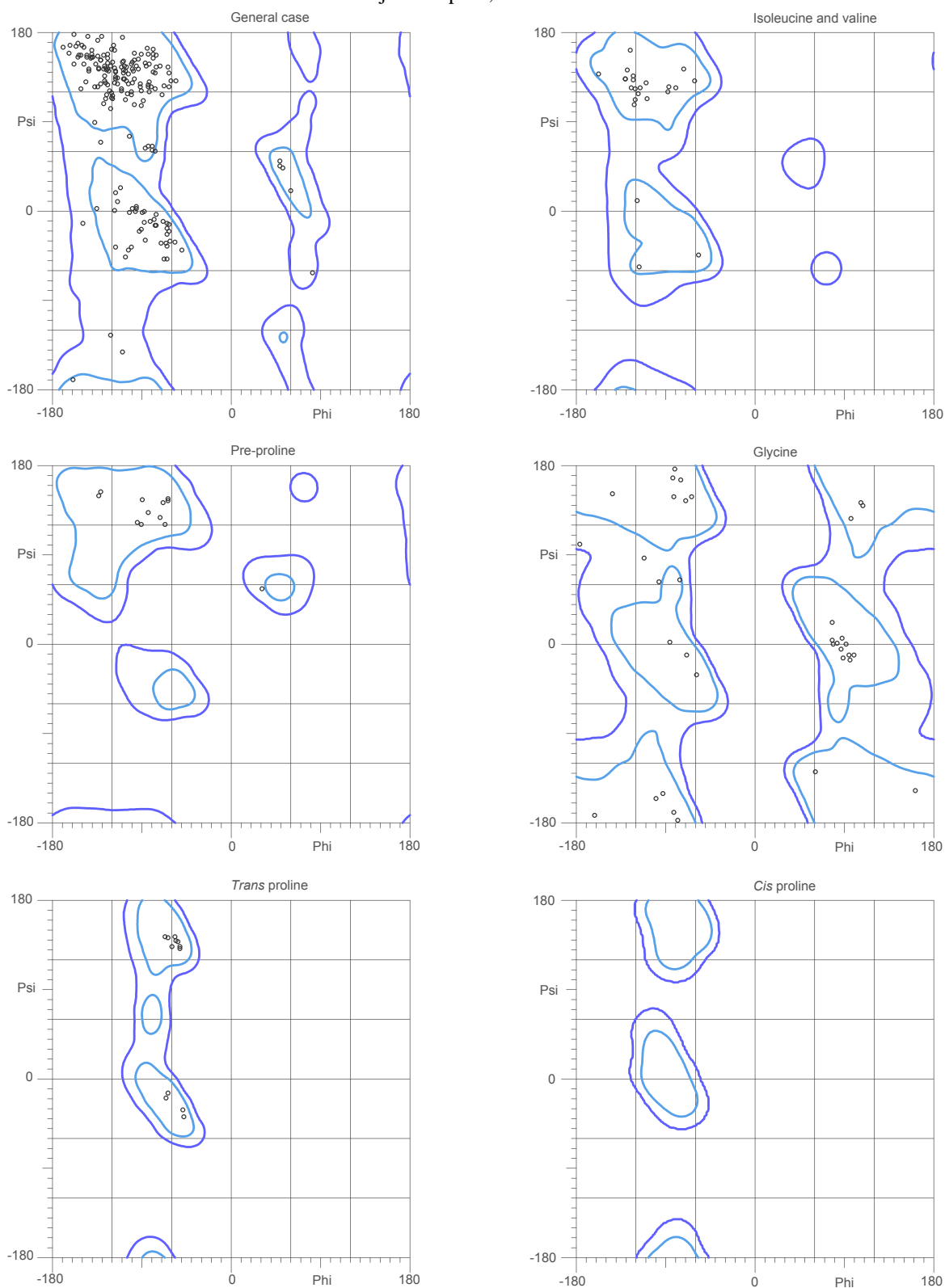


# MolProbity Ramachandran analysis

7jv2.H.pdb, model 1



96.4% (264/274) of all residues were in favored (98%) regions.

100.0% (274/274) of all residues were in allowed (>99.8%) regions.

There were no outliers.