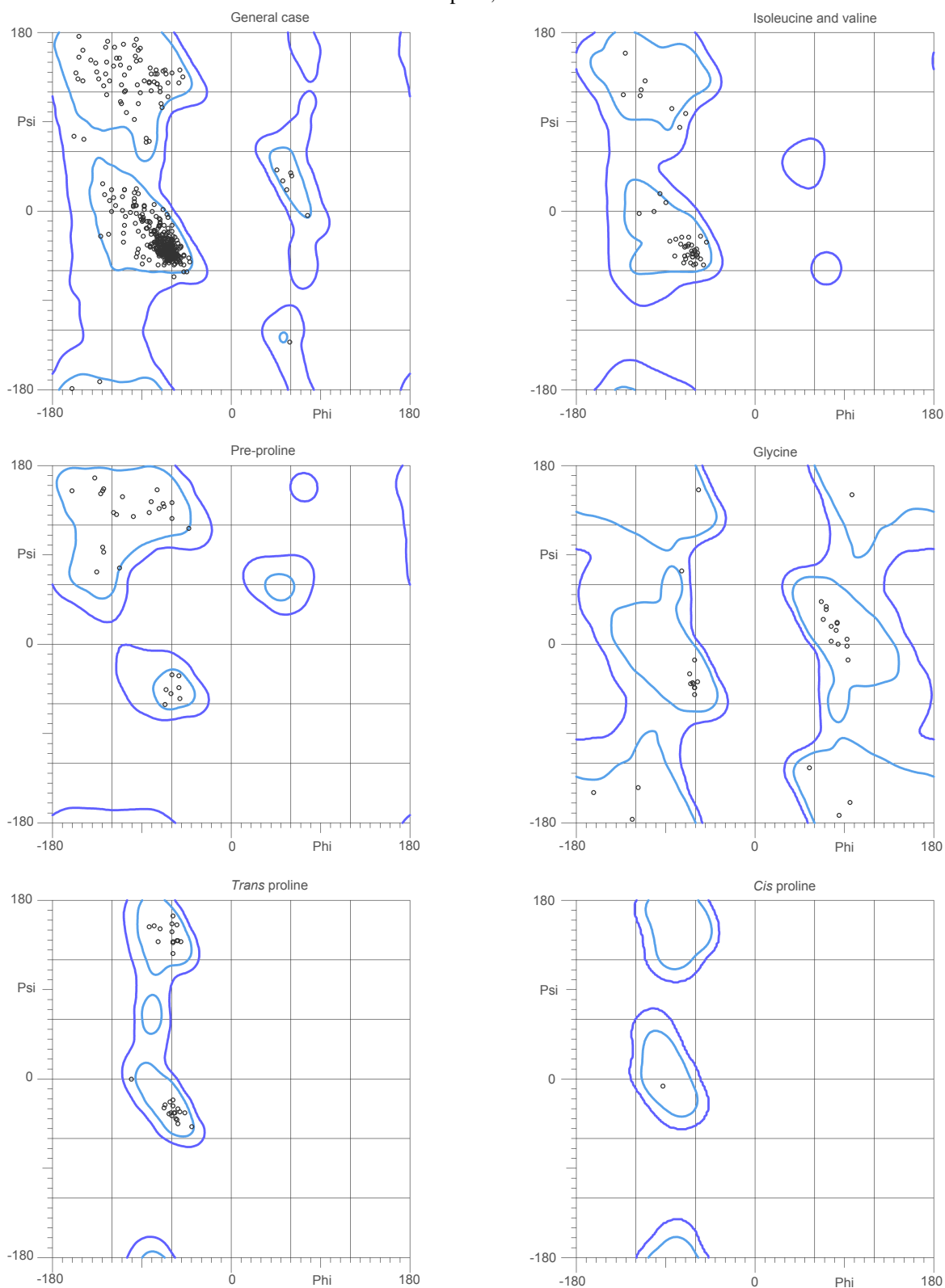


# MolProbity Ramachandran analysis

4c2r.H.pdb, model 1



98.0% (583/595) of all residues were in favored (98%) regions.

100.0% (595/595) of all residues were in allowed (>99.8%) regions.

There were no outliers.