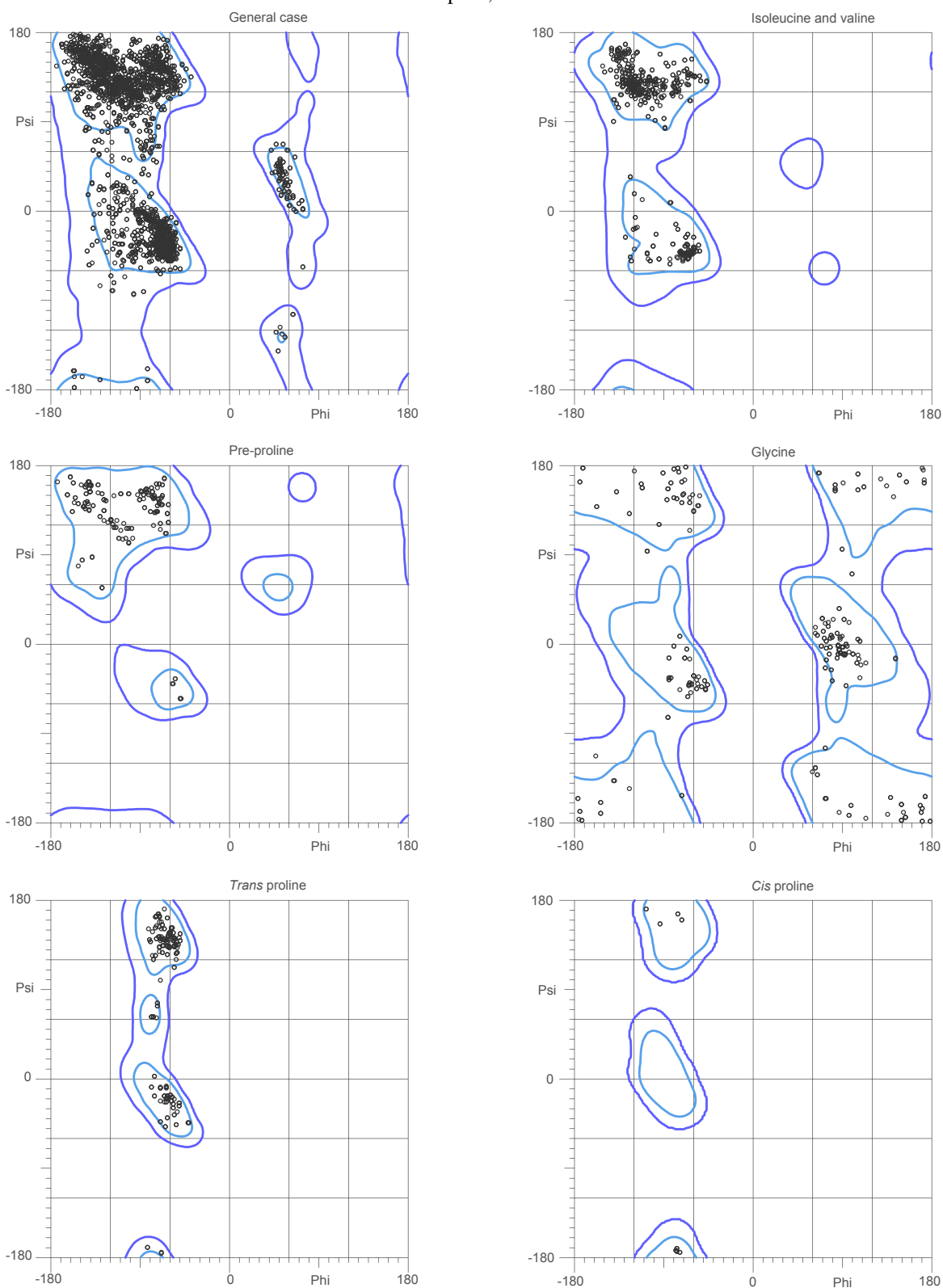


# MolProbity Ramachandran analysis

7106.H.pdb, model 1



96.2% (4368/4540) of all residues were in favored (98%) regions.

100.0% (4540/4540) of all residues were in allowed (>99.8%) regions.

There were no outliers.