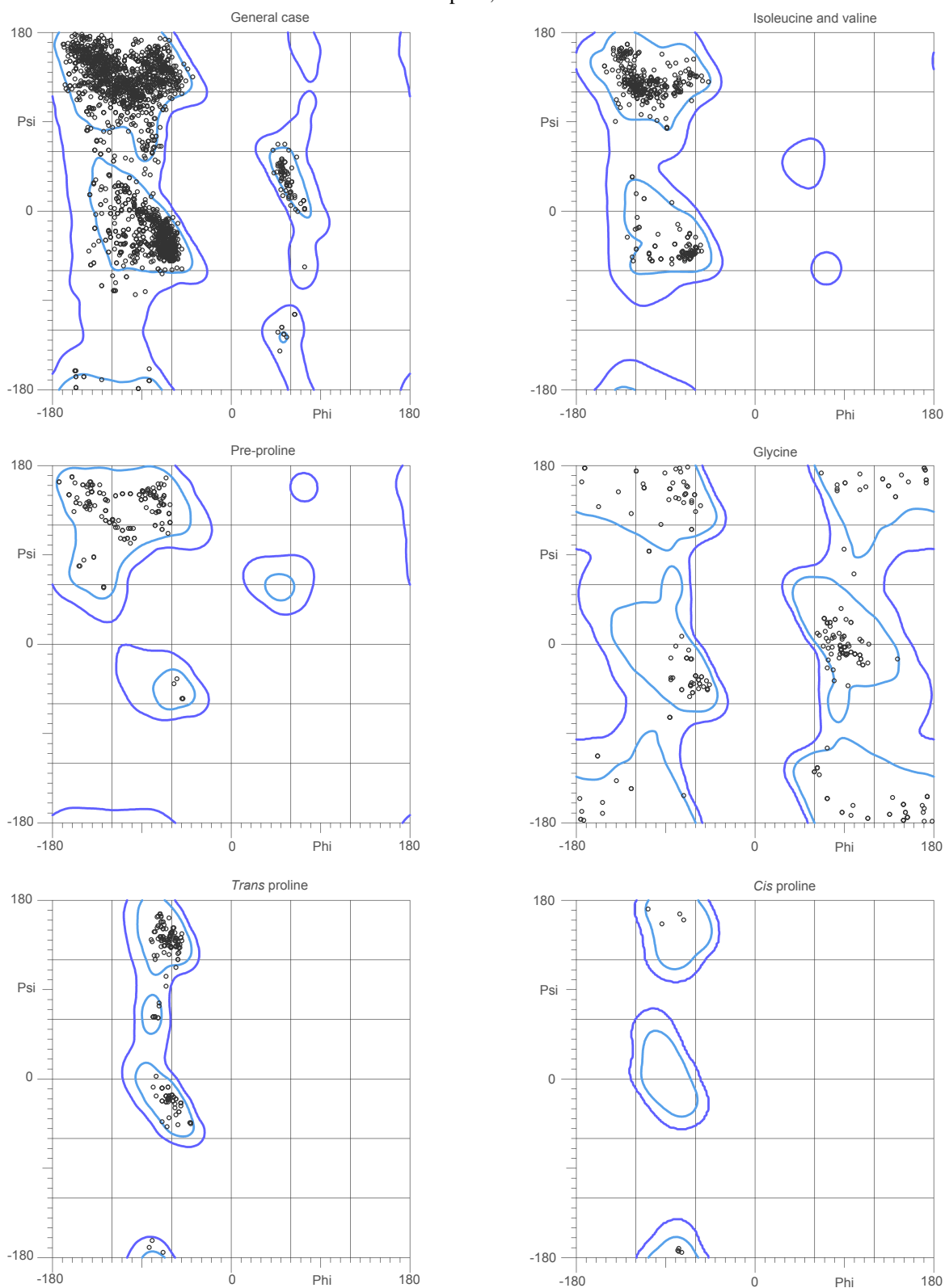


# MolProbity Ramachandran analysis

7109.H.pdb, model 1



96.7% (3569/3692) of all residues were in favored (98%) regions.

100.0% (3692/3692) of all residues were in allowed (>99.8%) regions.

There were no outliers.