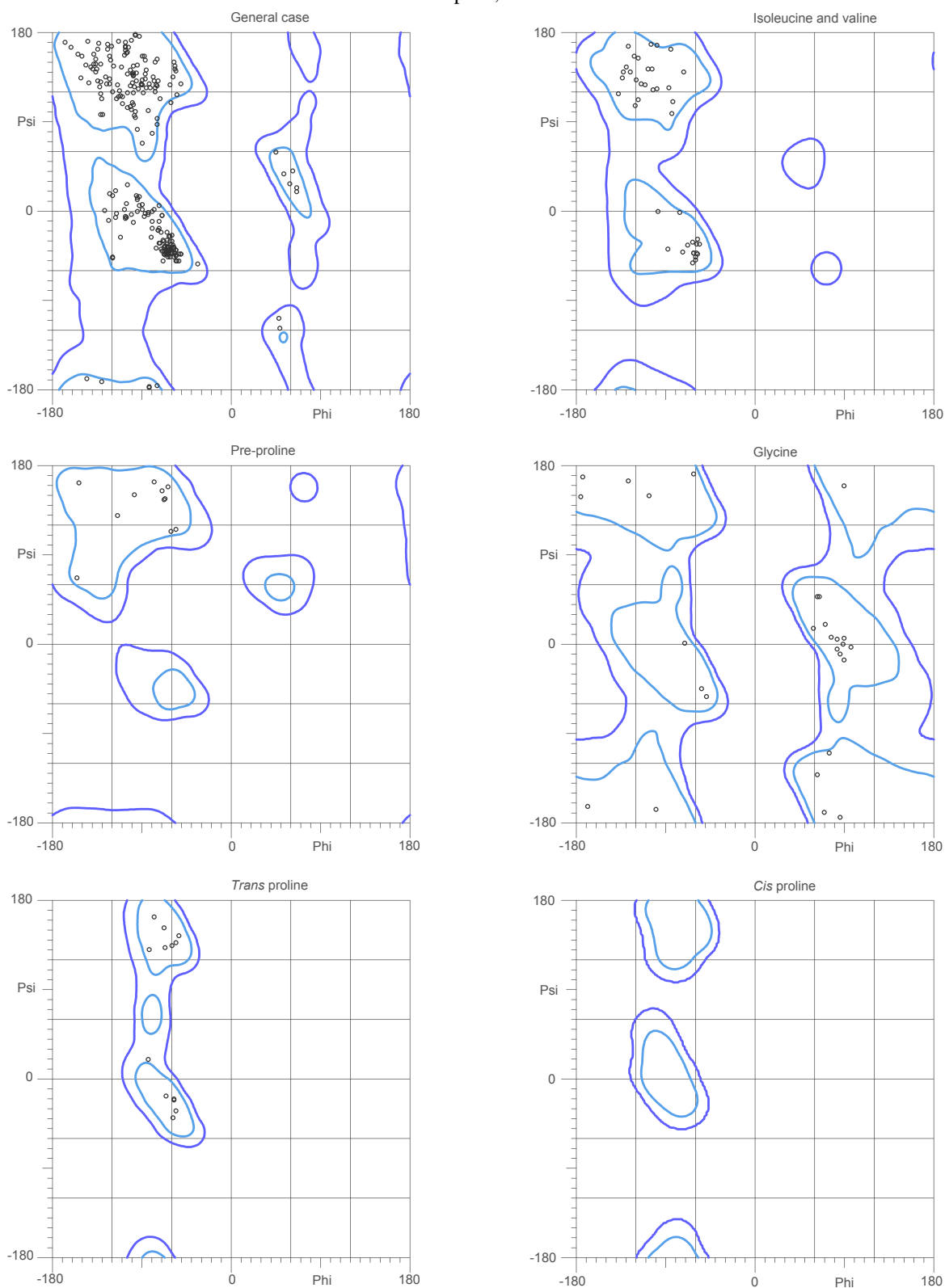


# MolProbity Ramachandran analysis

7cwb.H.pdb, model 1



98.2% (326/332) of all residues were in favored (98%) regions.

100.0% (332/332) of all residues were in allowed (>99.8%) regions.

There were no outliers.