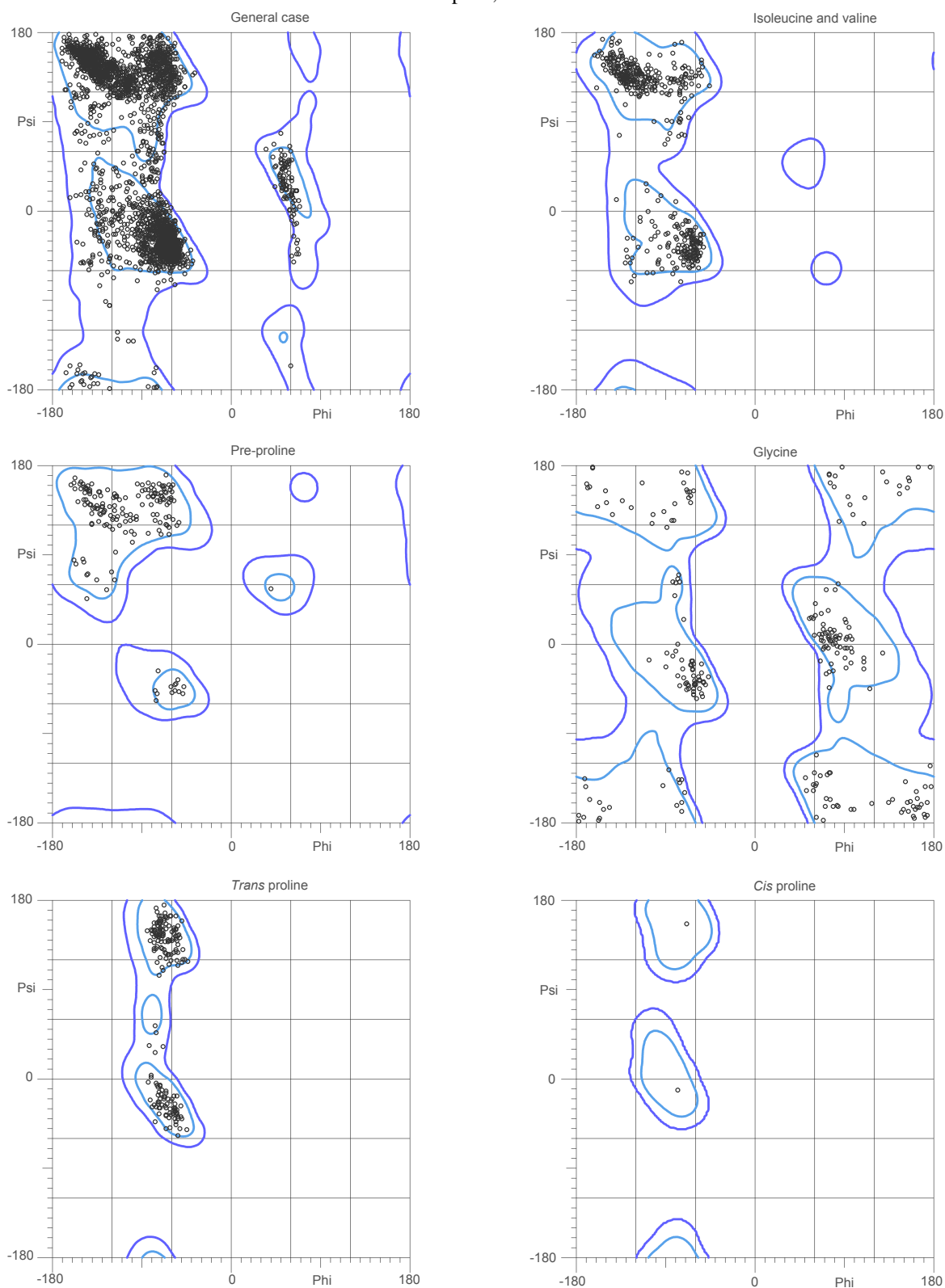


# MolProbity Ramachandran analysis

7a94.H.pdb, model 1



93.2% (3526/3785) of all residues were in favored (98%) regions.

100.0% (3785/3785) of all residues were in allowed (>99.8%) regions.

There were no outliers.