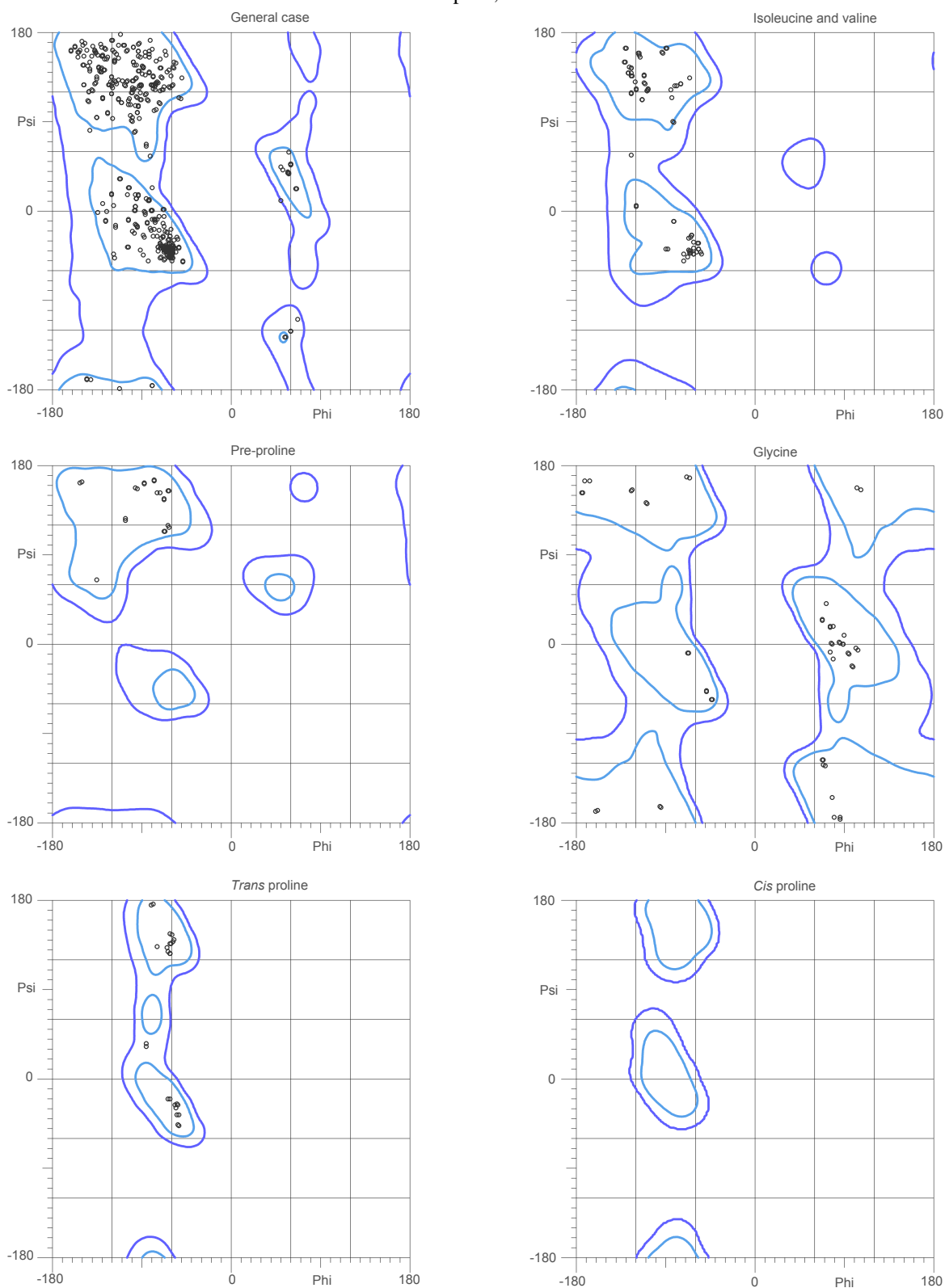


# MolProbity Ramachandran analysis

7com.H.pdb, model 1



98.1% (583/594) of all residues were in favored (98%) regions.

100.0% (594/594) of all residues were in allowed (>99.8%) regions.

There were no outliers.