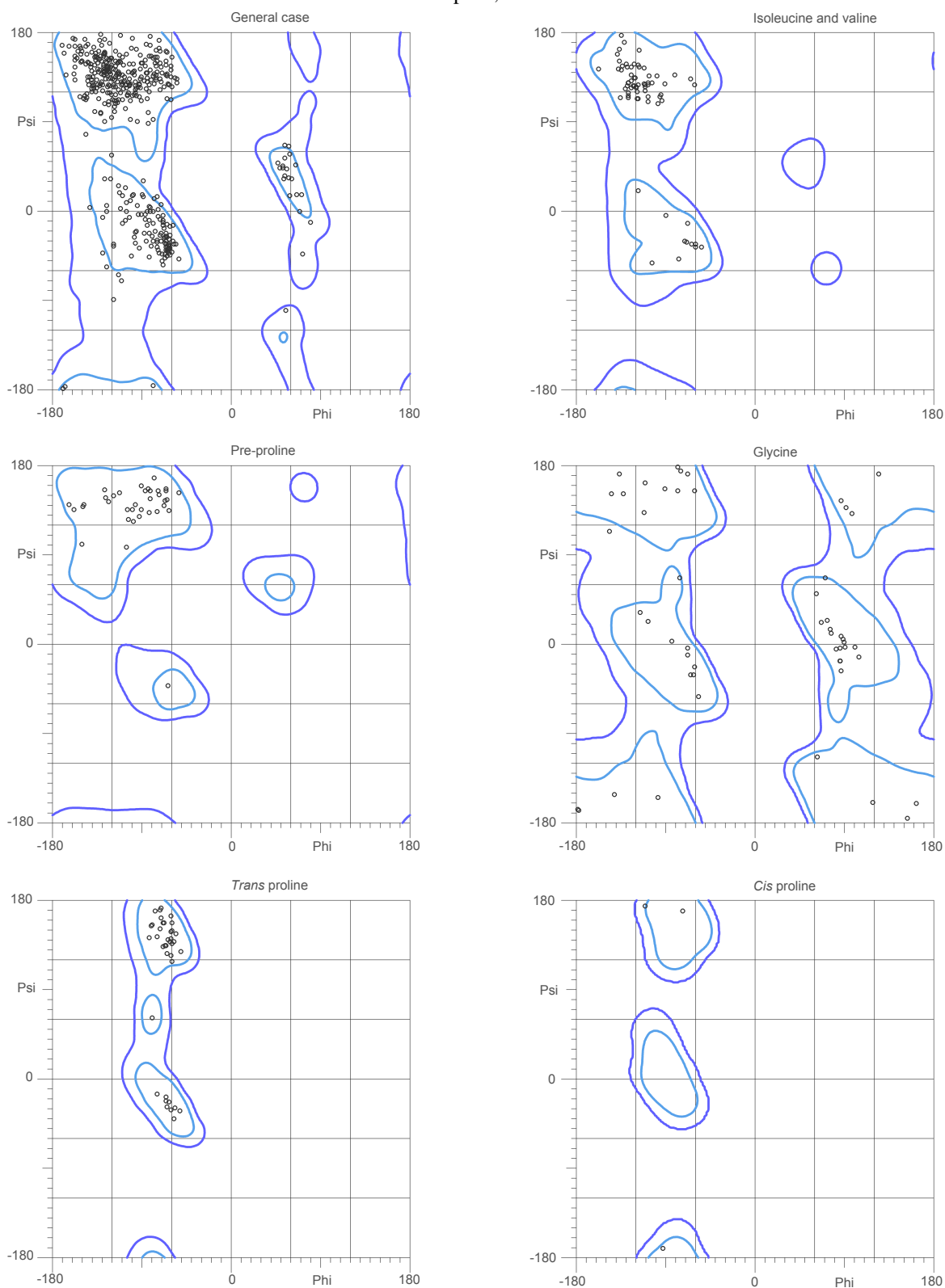


# MolProbity Ramachandran analysis

7cm4.H.pdb, model 1



96.8% (607/627) of all residues were in favored (98%) regions.

100.0% (627/627) of all residues were in allowed (>99.8%) regions.

There were no outliers.