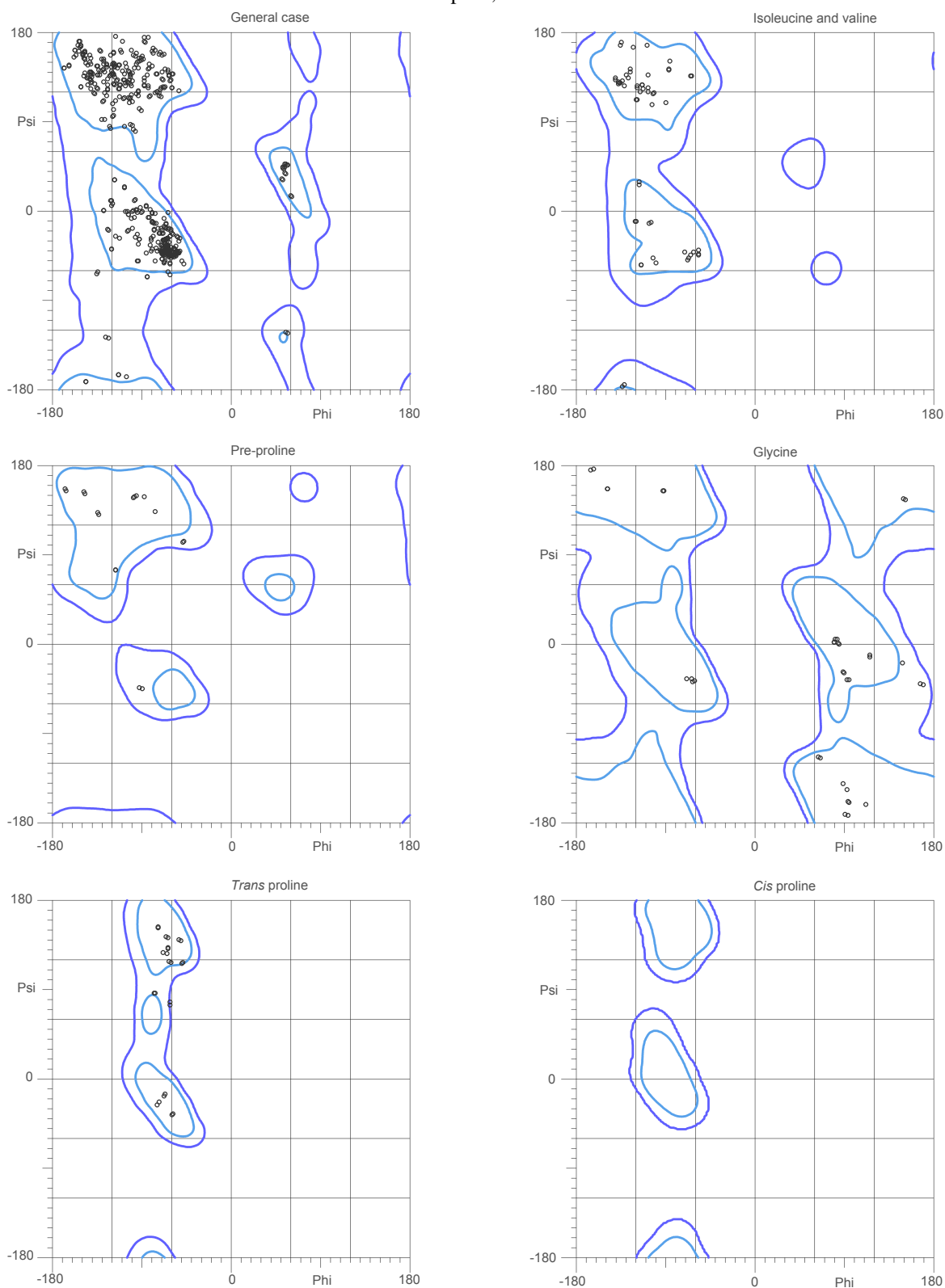


# MolProbity Ramachandran analysis

7d47.H.pdb, model 1



95.9% (582/607) of all residues were in favored (98%) regions.

100.0% (607/607) of all residues were in allowed (>99.8%) regions.

There were no outliers.