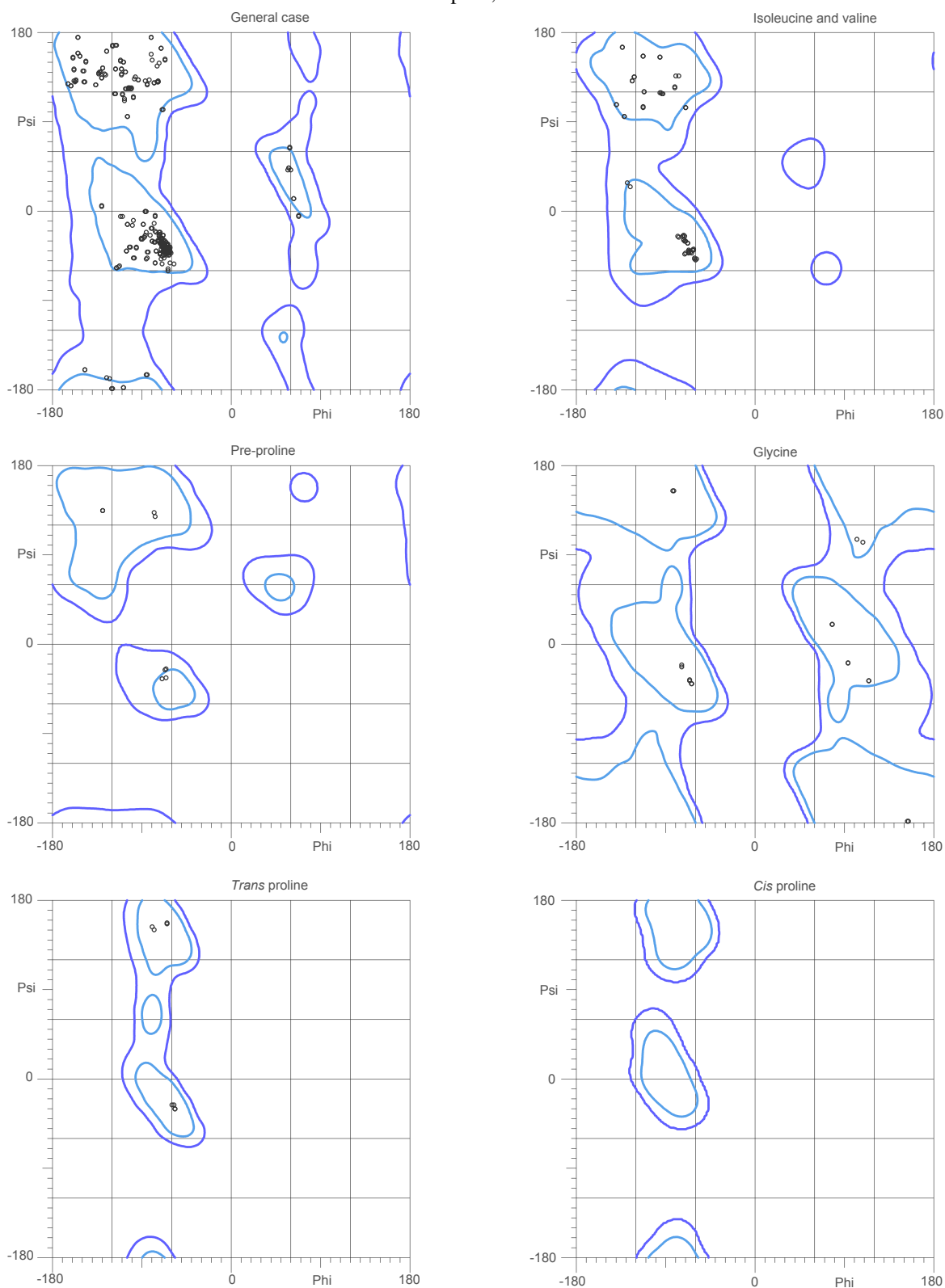


# MolProbity Ramachandran analysis

6xdc.H.pdb, model 1



96.0% (363/378) of all residues were in favored (98%) regions.

100.0% (378/378) of all residues were in allowed (>99.8%) regions.

There were no outliers.