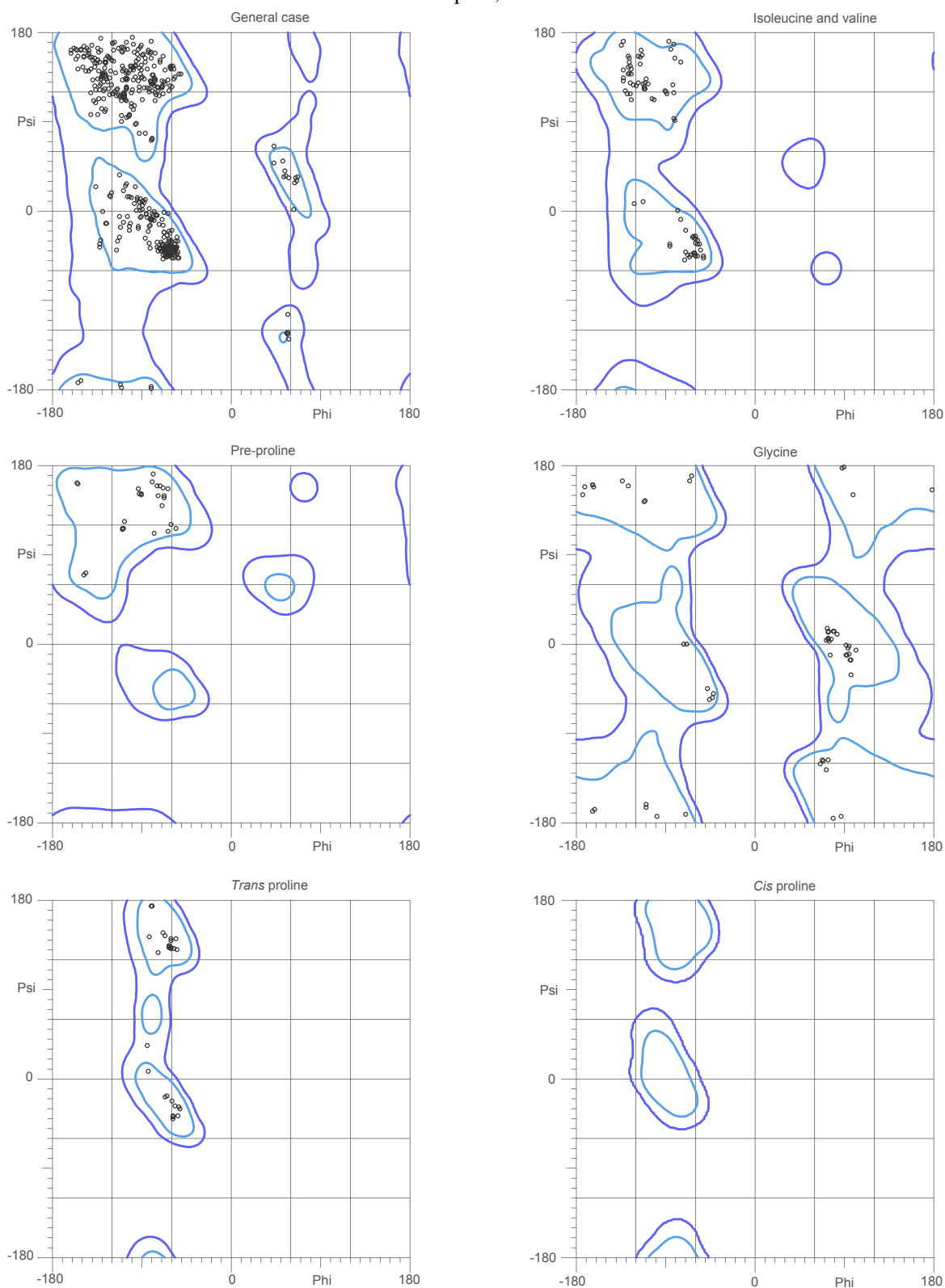


# MolProbity Ramachandran analysis

6xhm.H.pdb, model 1



98.0% (632/645) of all residues were in favored (98%) regions.

100.0% (645/645) of all residues were in allowed (>99.8%) regions.

There were no outliers.