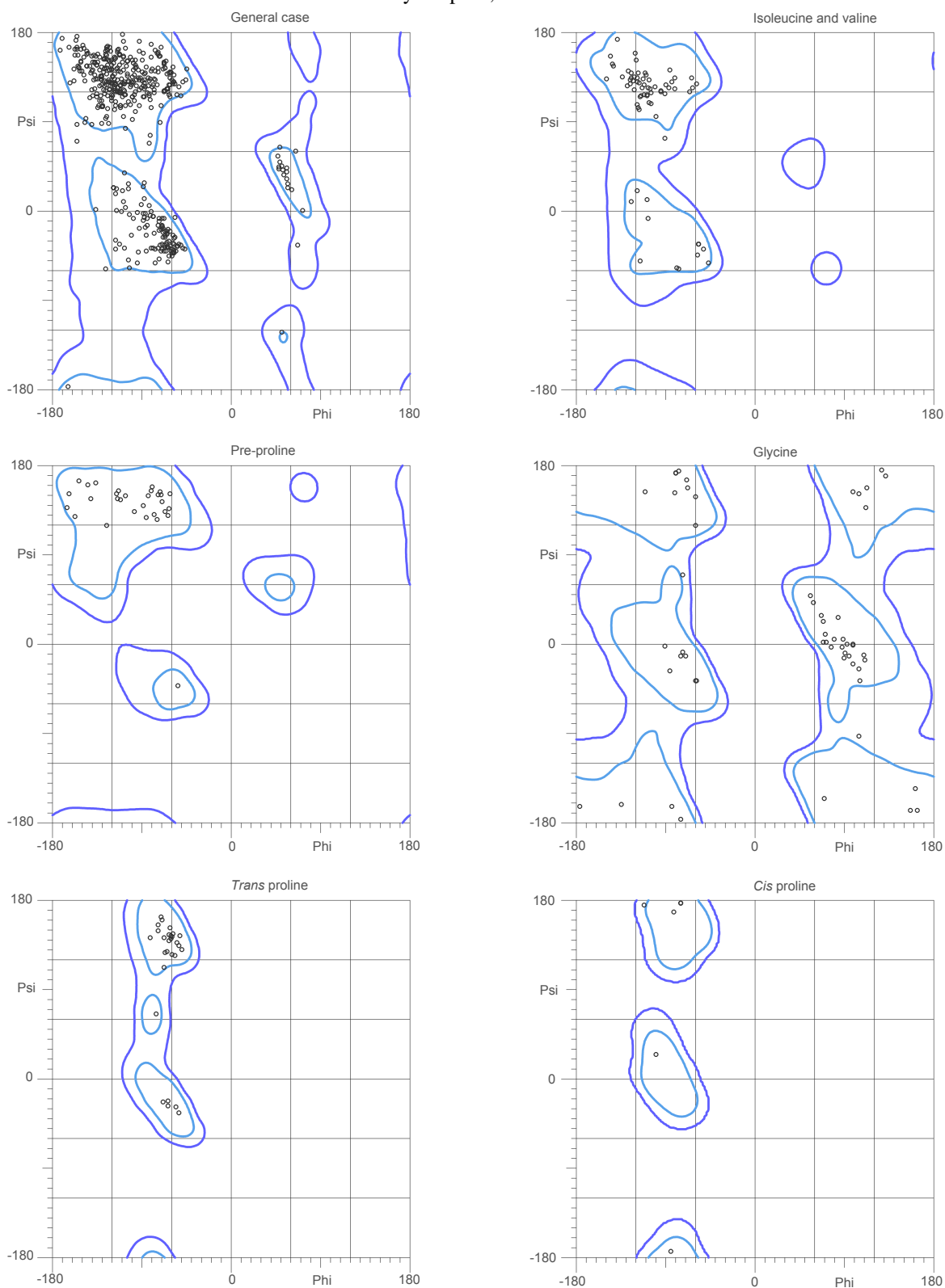


# MolProbity Ramachandran analysis

7kfy.H.pdb, model 1



97.6% (623/638) of all residues were in favored (98%) regions.

100.0% (638/638) of all residues were in allowed (>99.8%) regions.

There were no outliers.