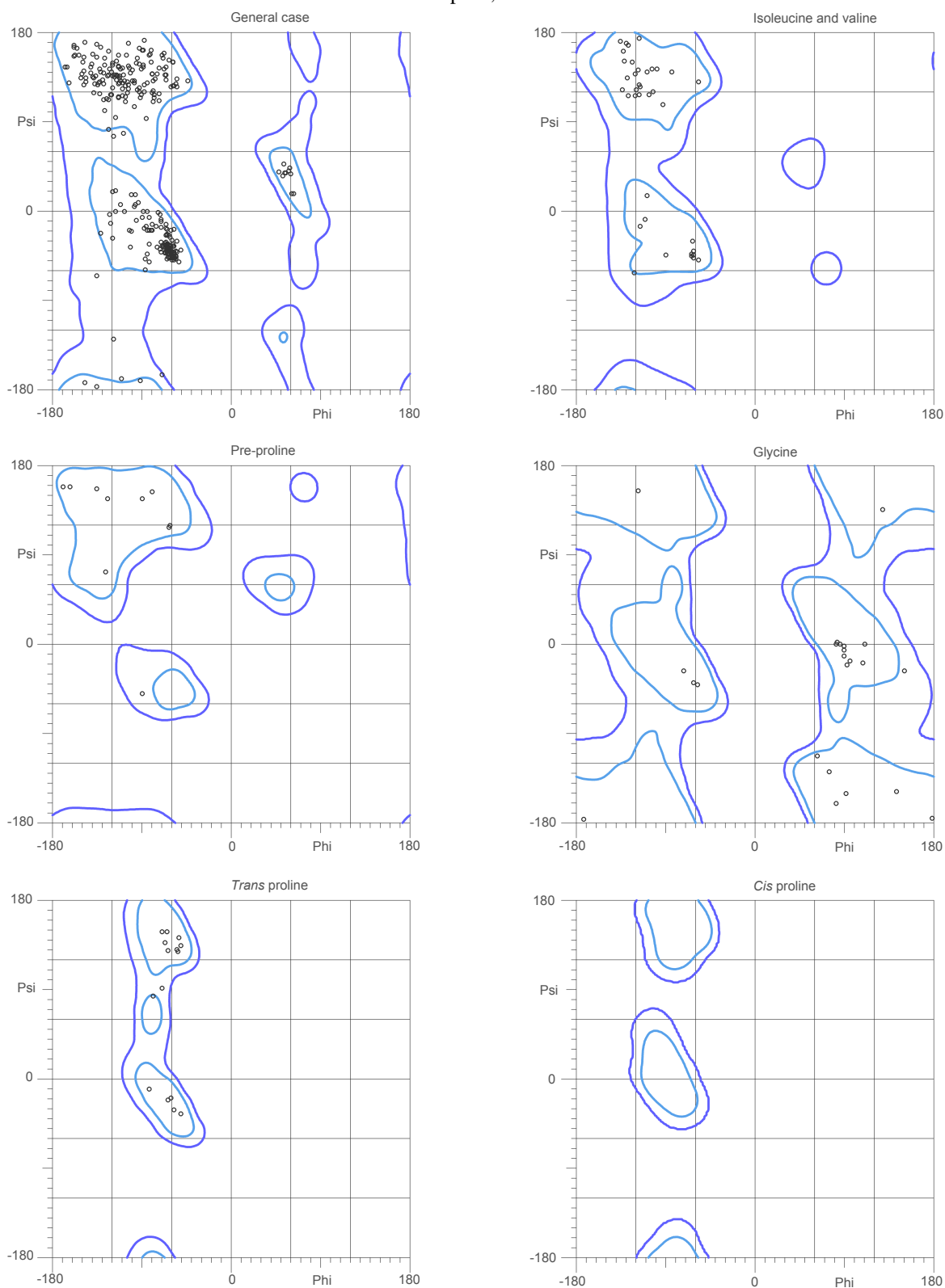


# MolProbity Ramachandran analysis

6xaa.H.pdb, model 1



97.4% (373/383) of all residues were in favored (98%) regions.

100.0% (383/383) of all residues were in allowed (>99.8%) regions.

There were no outliers.