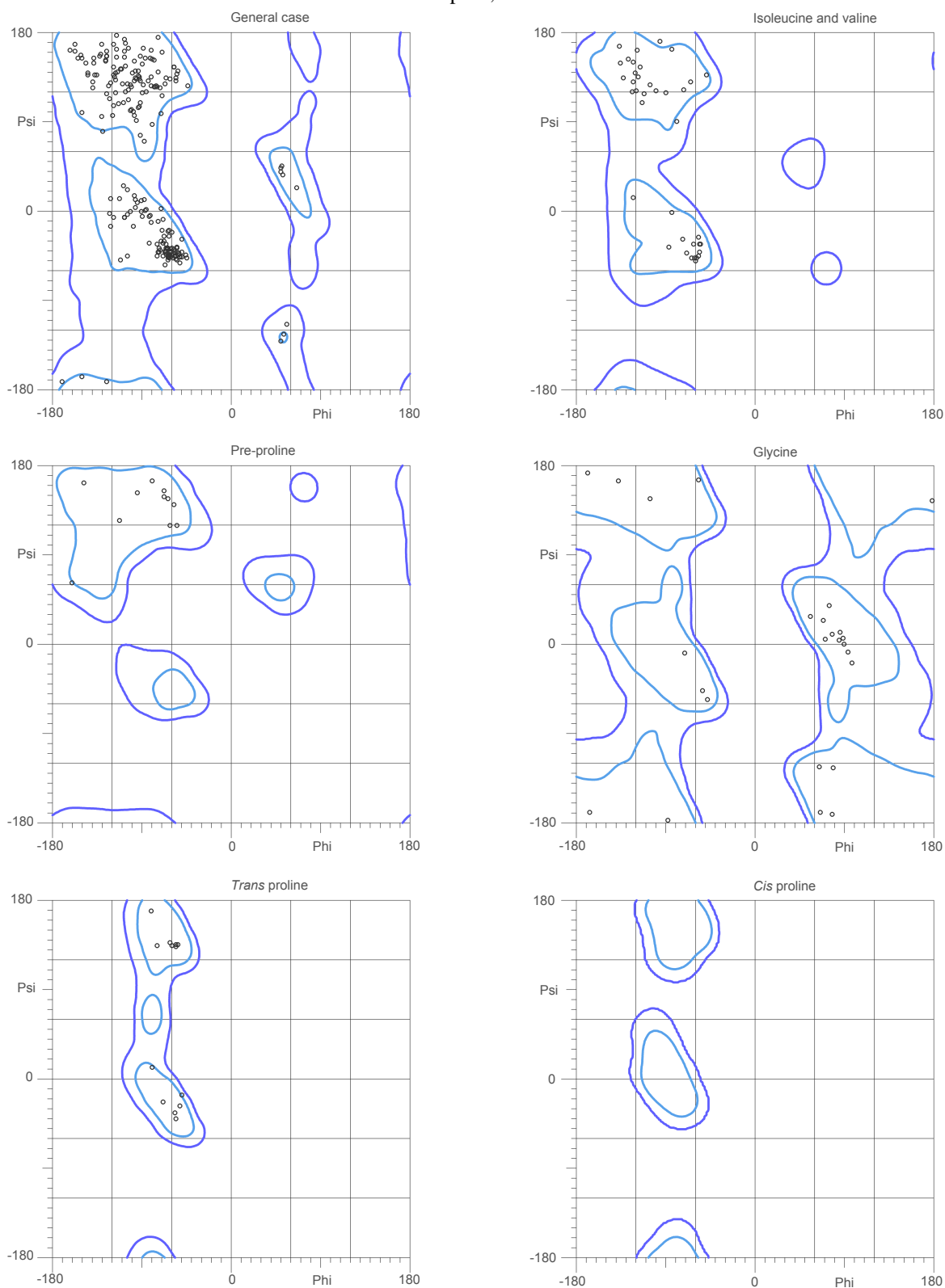


# MolProbity Ramachandran analysis

6xfn.H.pdb, model 1



98.4% (301/306) of all residues were in favored (98%) regions.

100.0% (306/306) of all residues were in allowed (>99.8%) regions.

There were no outliers.