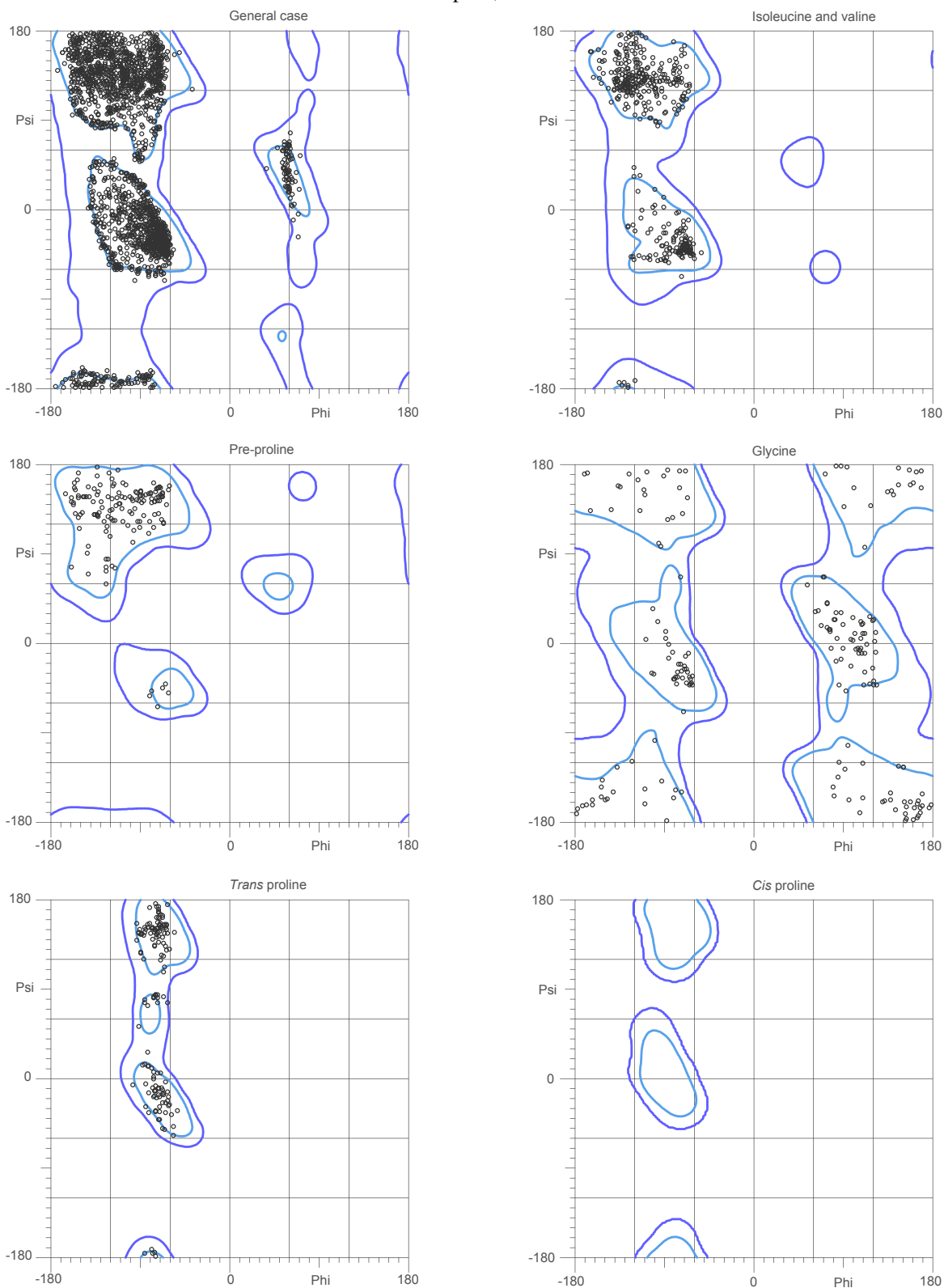


# MolProbity Ramachandran analysis

6xm5.H.pdb, model 1



93.2% (2893/3105) of all residues were in favored (98%) regions.

100.0% (3105/3105) of all residues were in allowed (>99.8%) regions.

There were no outliers.