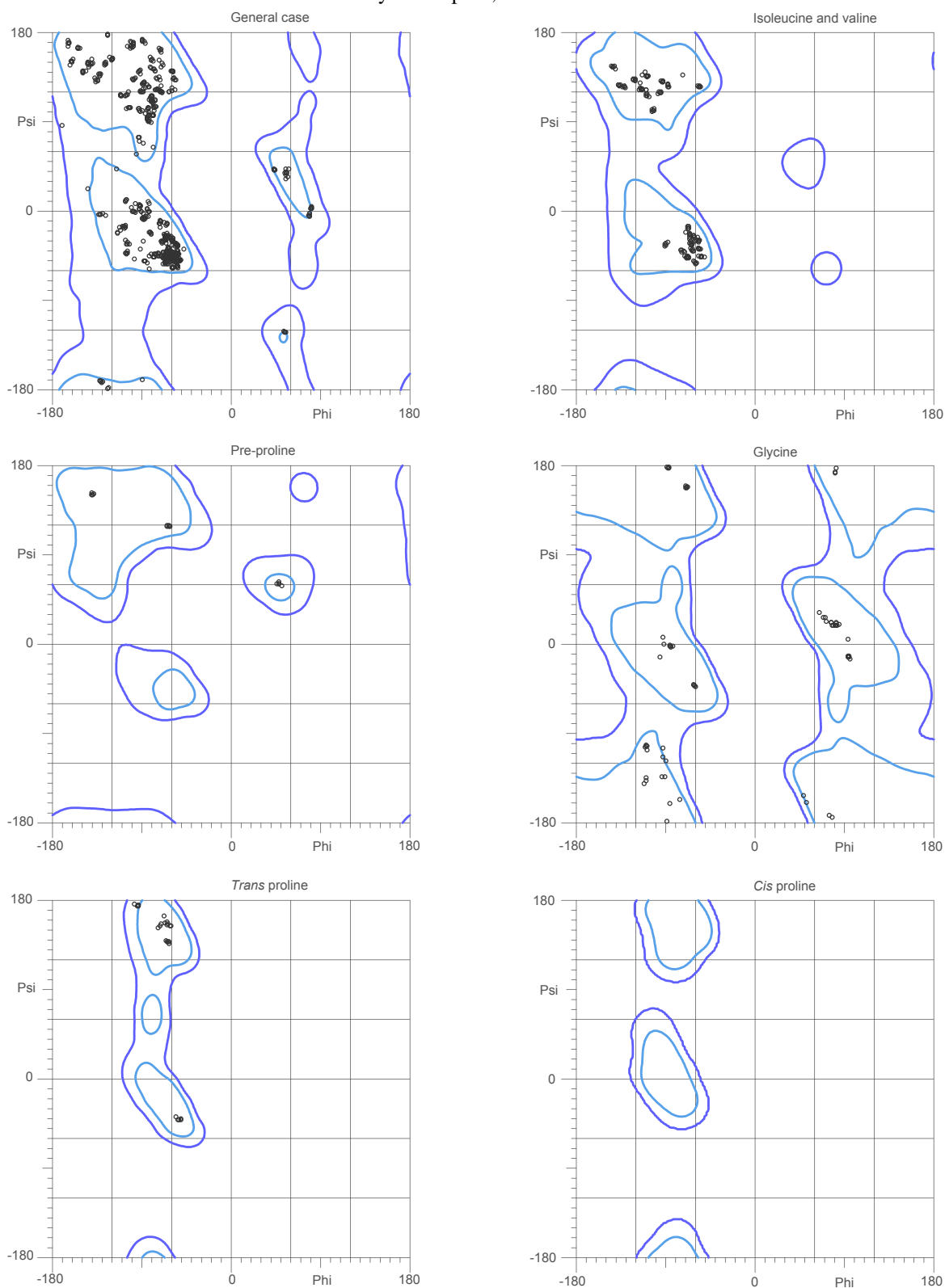


# MolProbity Ramachandran analysis

6yw1.H.pdb, model 1



98.1% (831/847) of all residues were in favored (98%) regions.

100.0% (847/847) of all residues were in allowed (>99.8%) regions.

There were no outliers.