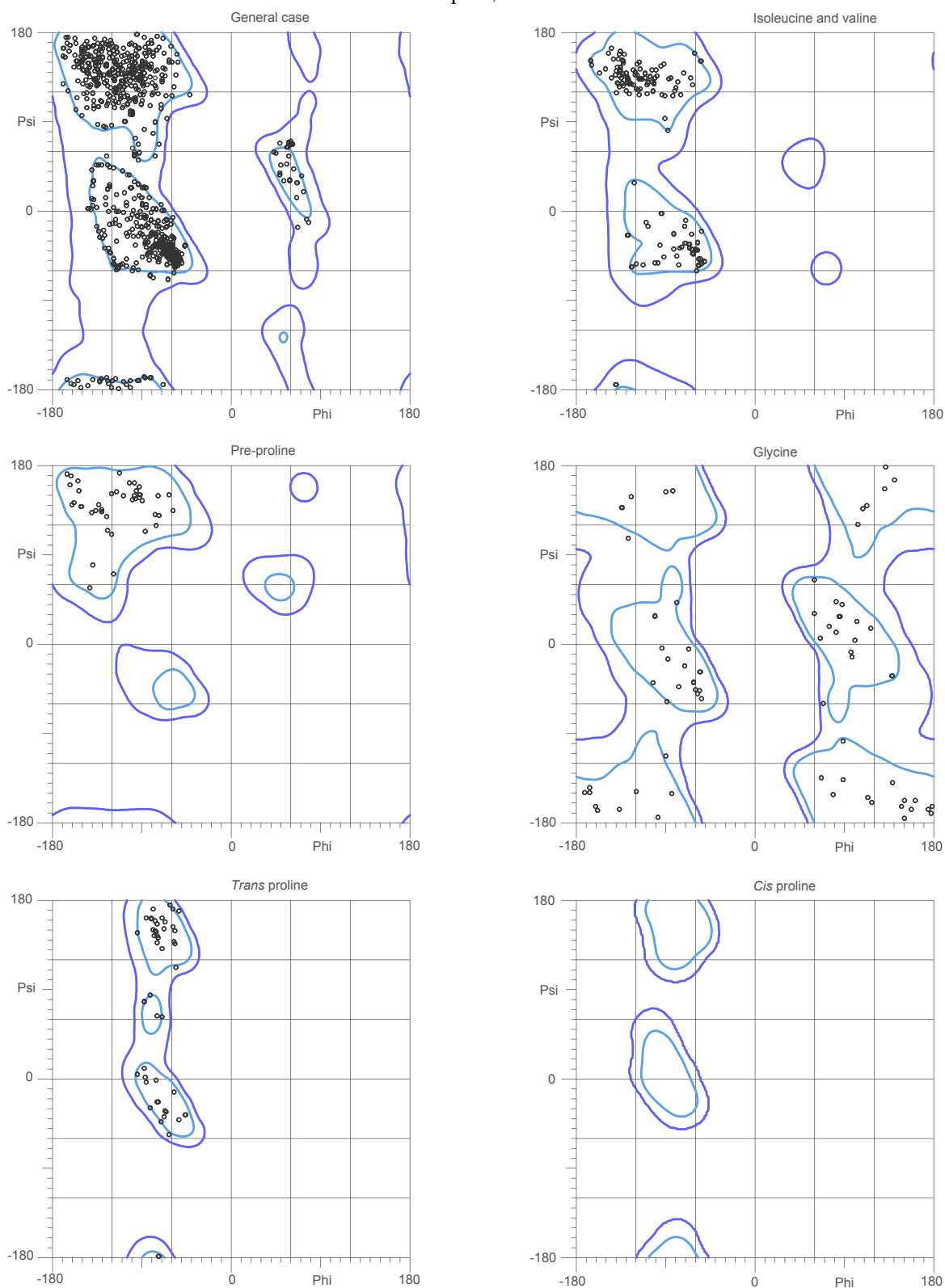


# MolProbity Ramachandran analysis

6zb5.H.pdb, model 1



92.4% (2795/3024) of all residues were in favored (98%) regions.

100.0% (3024/3024) of all residues were in allowed (>99.8%) regions.

There were no outliers.