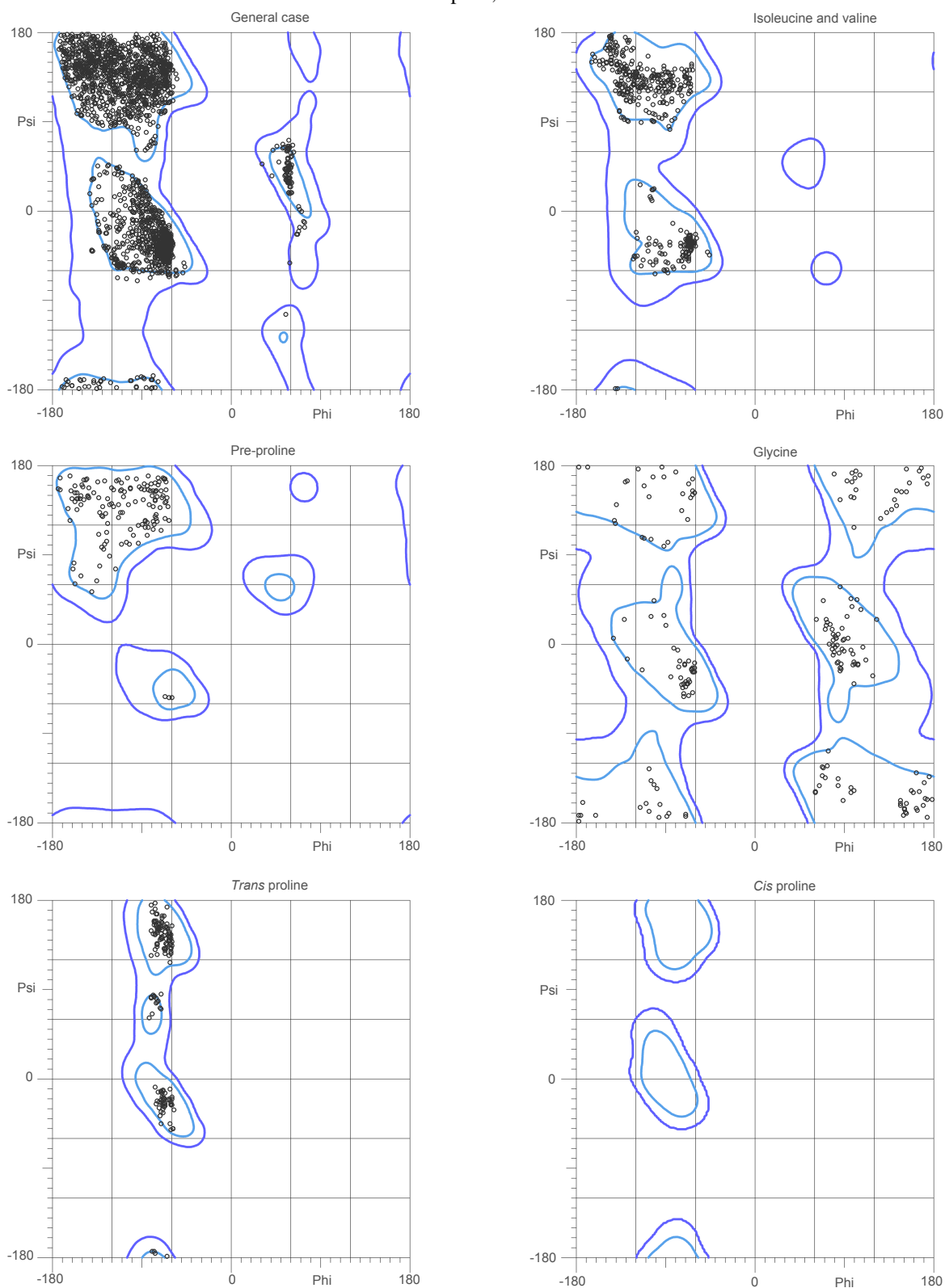


# MolProbity Ramachandran analysis

7ddn.H.pdb, model 1



96.4% (3053/3166) of all residues were in favored (98%) regions.

100.0% (3166/3166) of all residues were in allowed (>99.8%) regions.

There were no outliers.