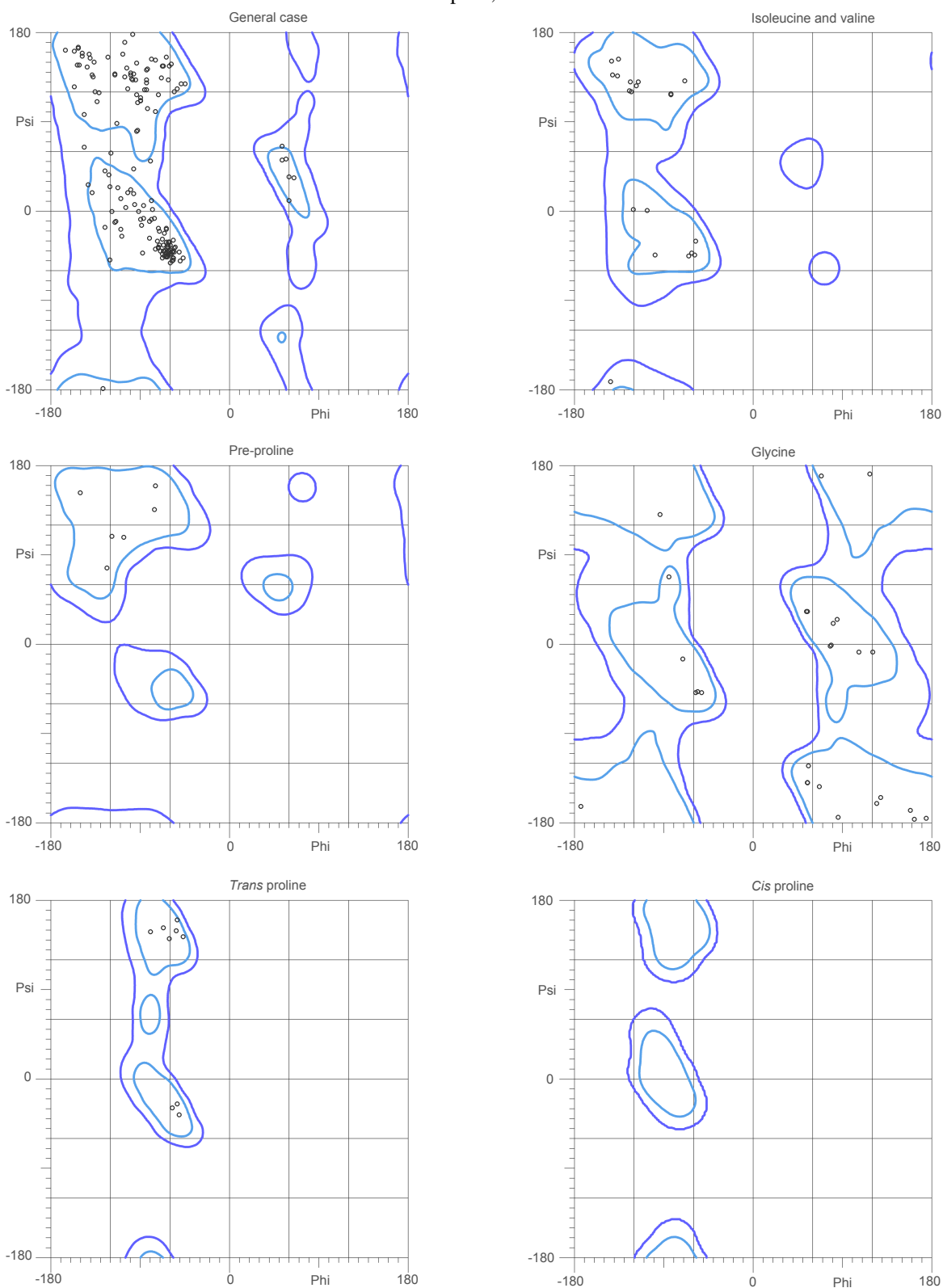


# MolProbity Ramachandran analysis

2xu5.H.pdb, model 1



96.8% (214/221) of all residues were in favored (98%) regions.

100.0% (221/221) of all residues were in allowed (>99.8%) regions.

There were no outliers.