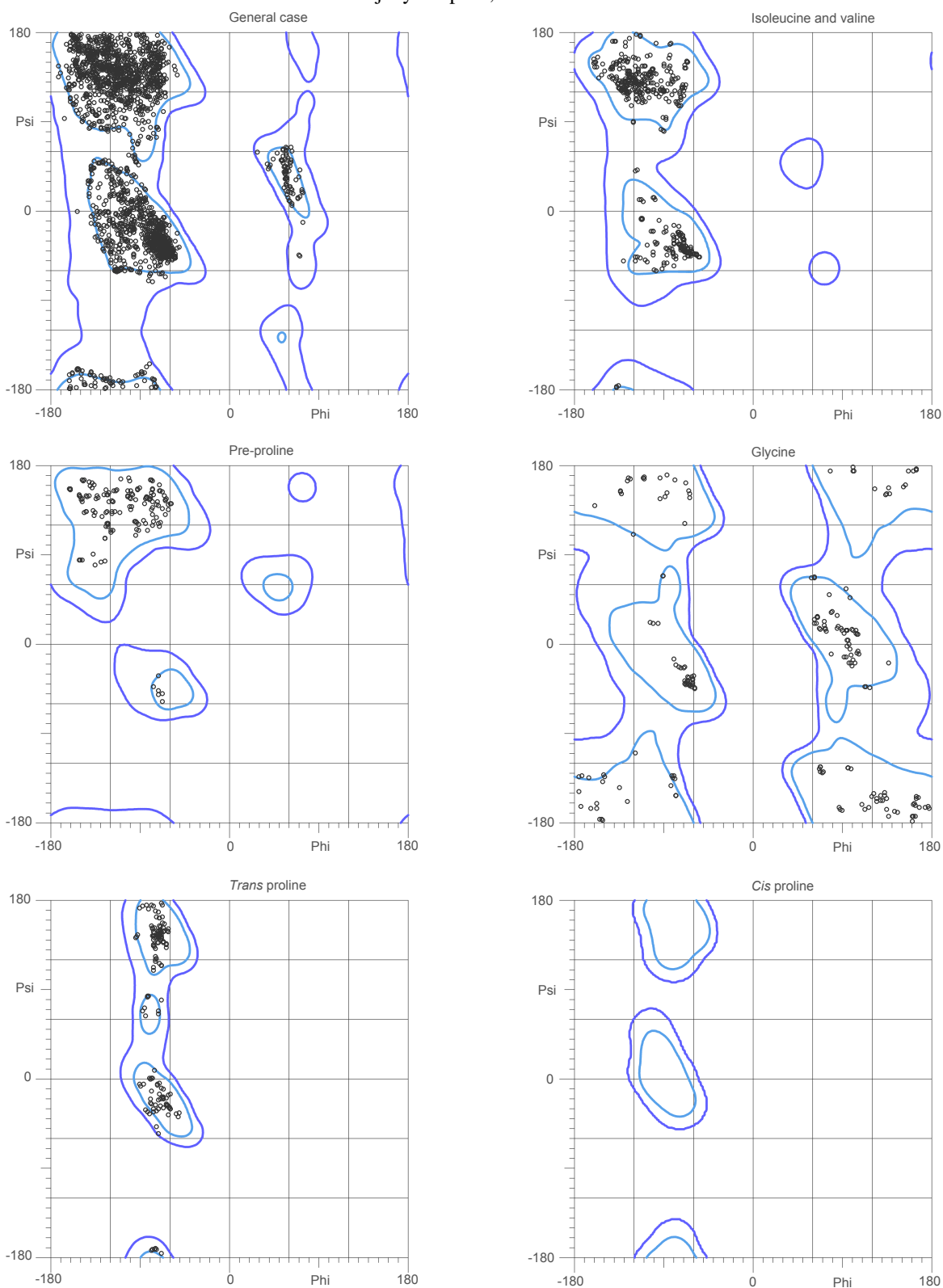


# MolProbity Ramachandran analysis

7jwy.H.pdb, model 1



93.9% (2943/3135) of all residues were in favored (98%) regions.

100.0% (3135/3135) of all residues were in allowed (>99.8%) regions.

There were no outliers.