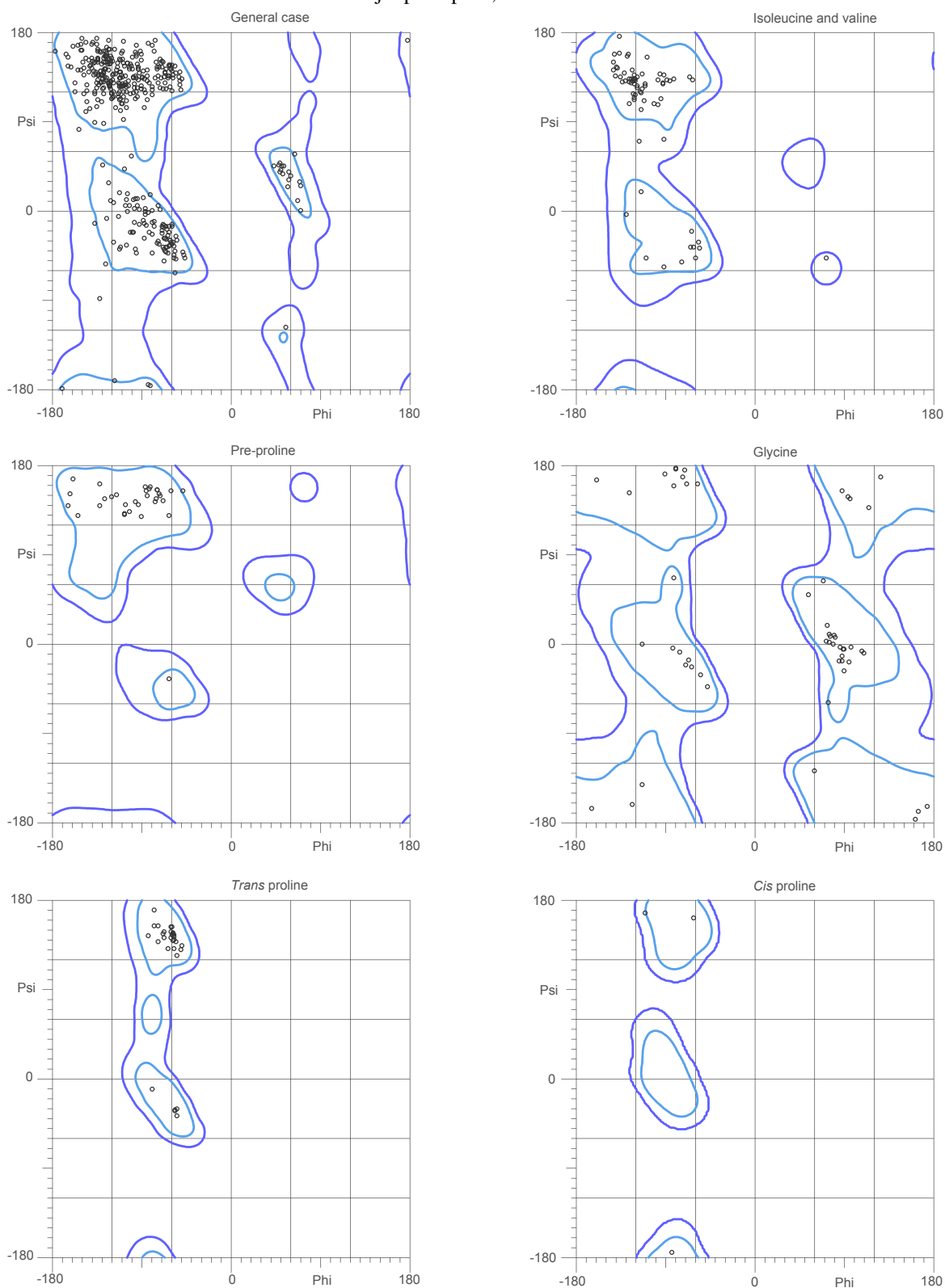


# MolProbity Ramachandran analysis

7jmp.H.pdb, model 1



97.1% (571/588) of all residues were in favored (98%) regions.

100.0% (588/588) of all residues were in allowed (>99.8%) regions.

There were no outliers.