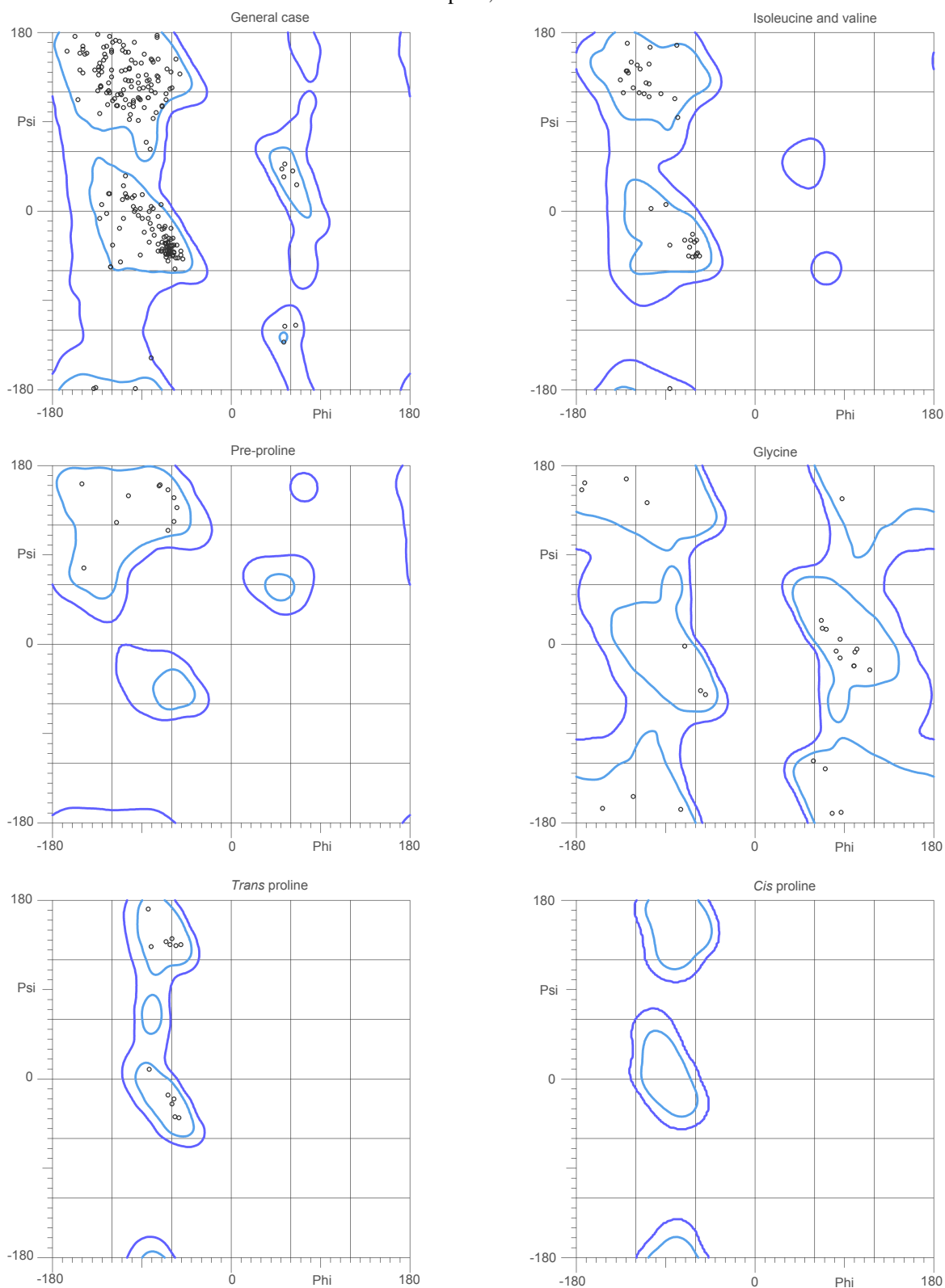


# MolProbity Ramachandran analysis

7arf.H.pdb, model 1



97.1% (302/311) of all residues were in favored (98%) regions.

100.0% (311/311) of all residues were in allowed (>99.8%) regions.

There were no outliers.