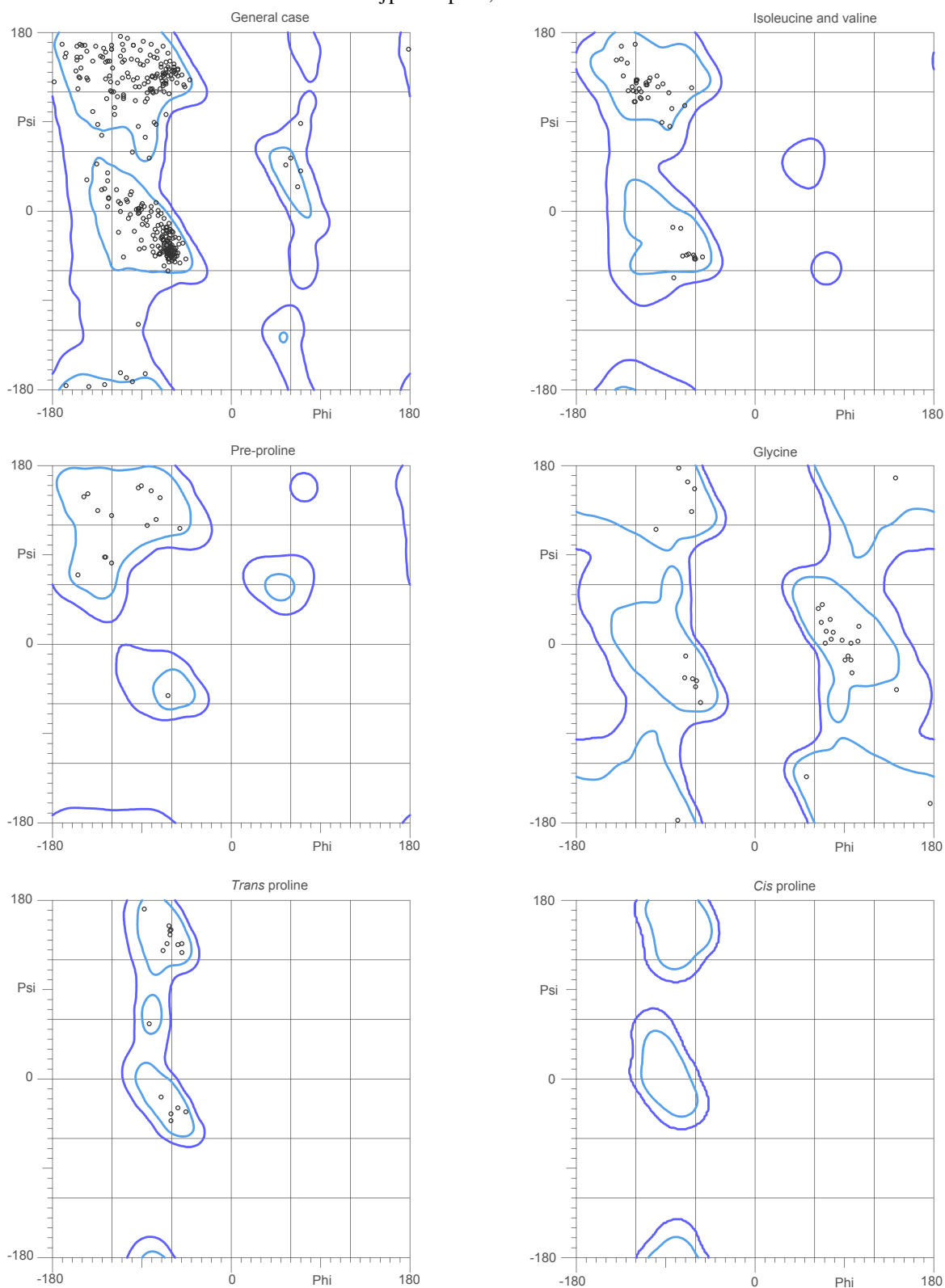


# MolProbity Ramachandran analysis

7jpe.H.pdb, model 1



96.6% (402/416) of all residues were in favored (98%) regions.

100.0% (416/416) of all residues were in allowed (>99.8%) regions.

There were no outliers.