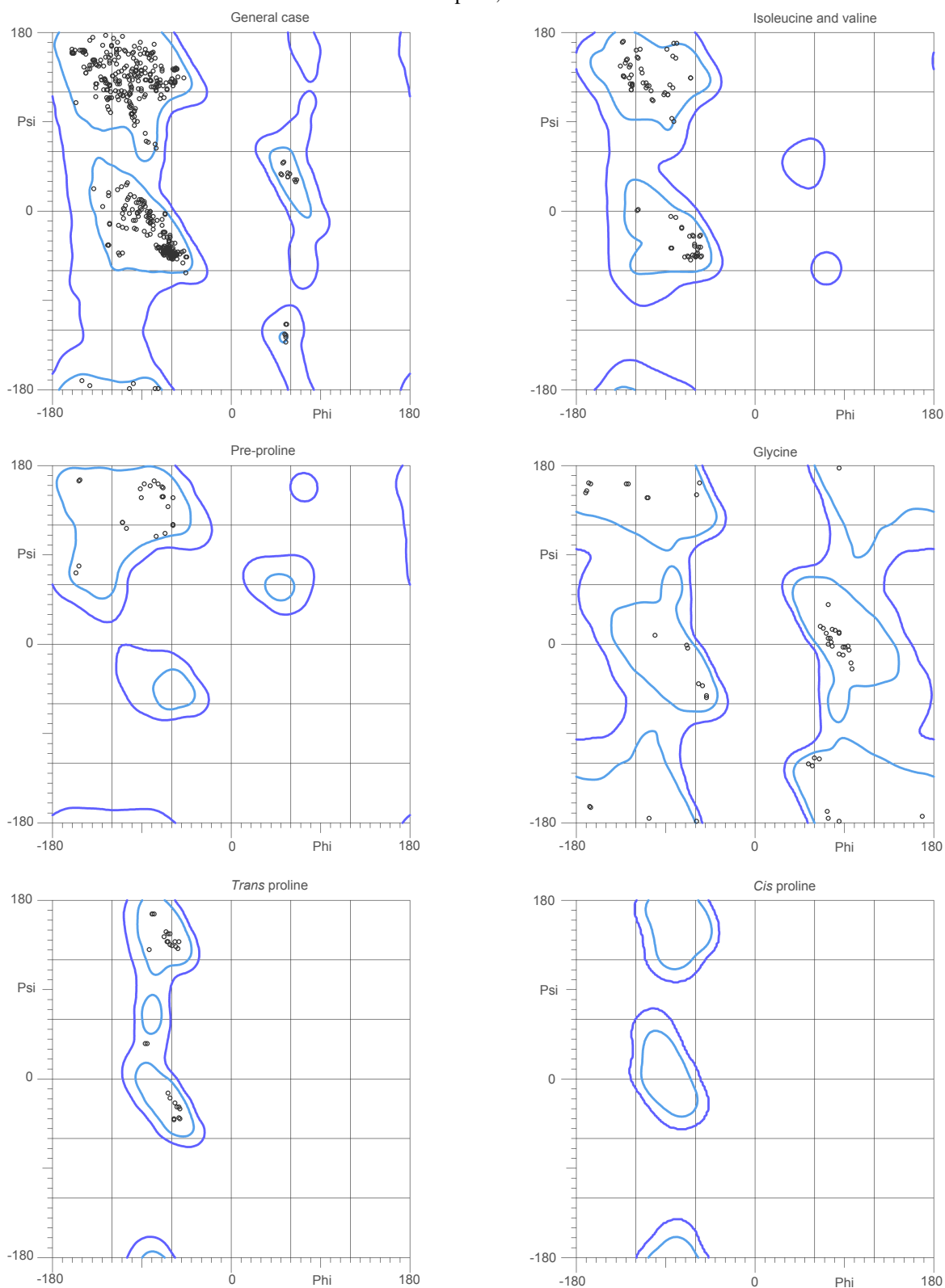


# MolProbity Ramachandran analysis

7d1m.H.pdb, model 1



98.6% (626/635) of all residues were in favored (98%) regions.

100.0% (635/635) of all residues were in allowed (>99.8%) regions.

There were no outliers.