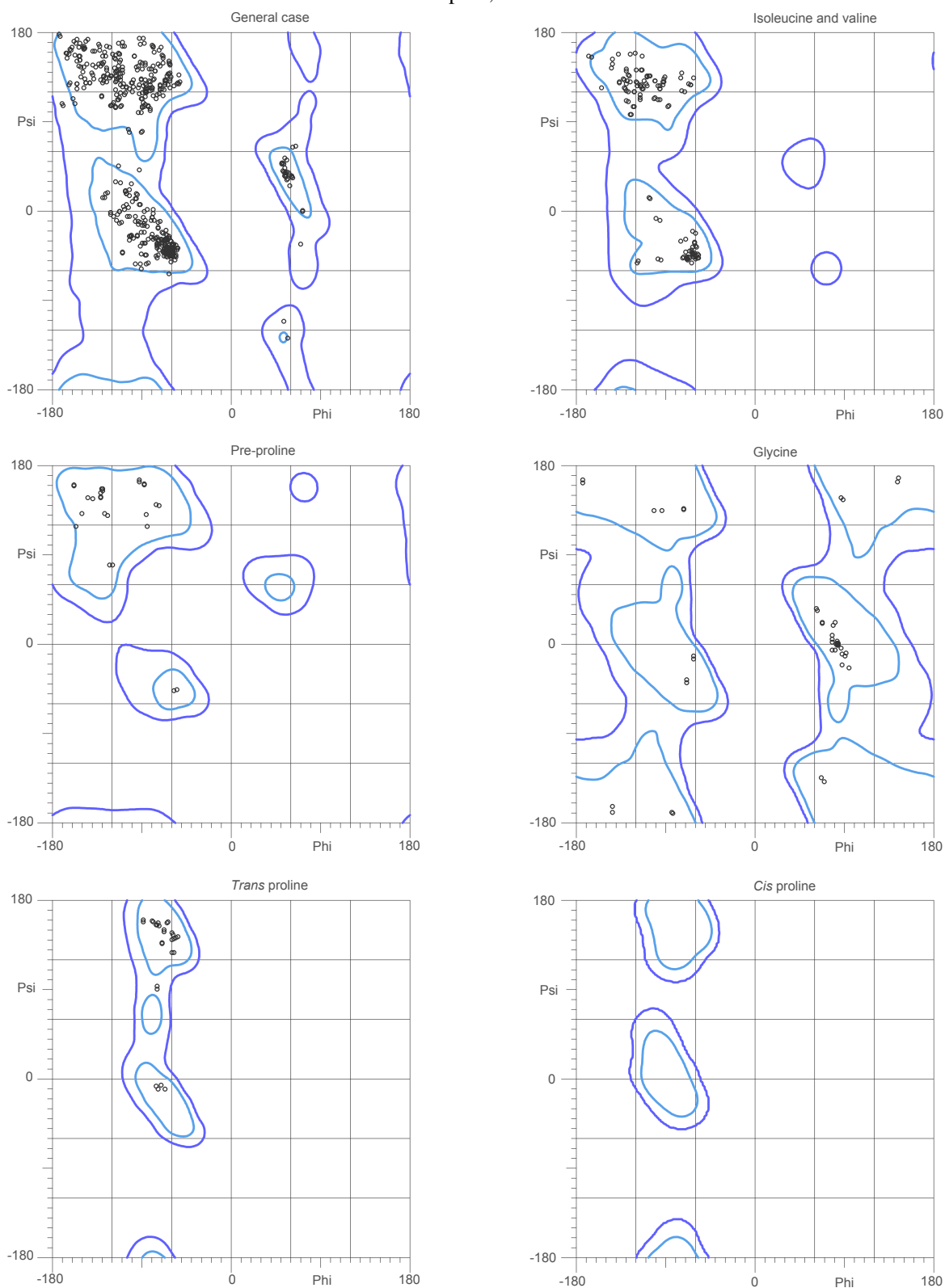


# MolProbity Ramachandran analysis

6x4i.H.pdb, model 1



97.7% (715/732) of all residues were in favored (98%) regions.

100.0% (732/732) of all residues were in allowed (>99.8%) regions.

There were no outliers.