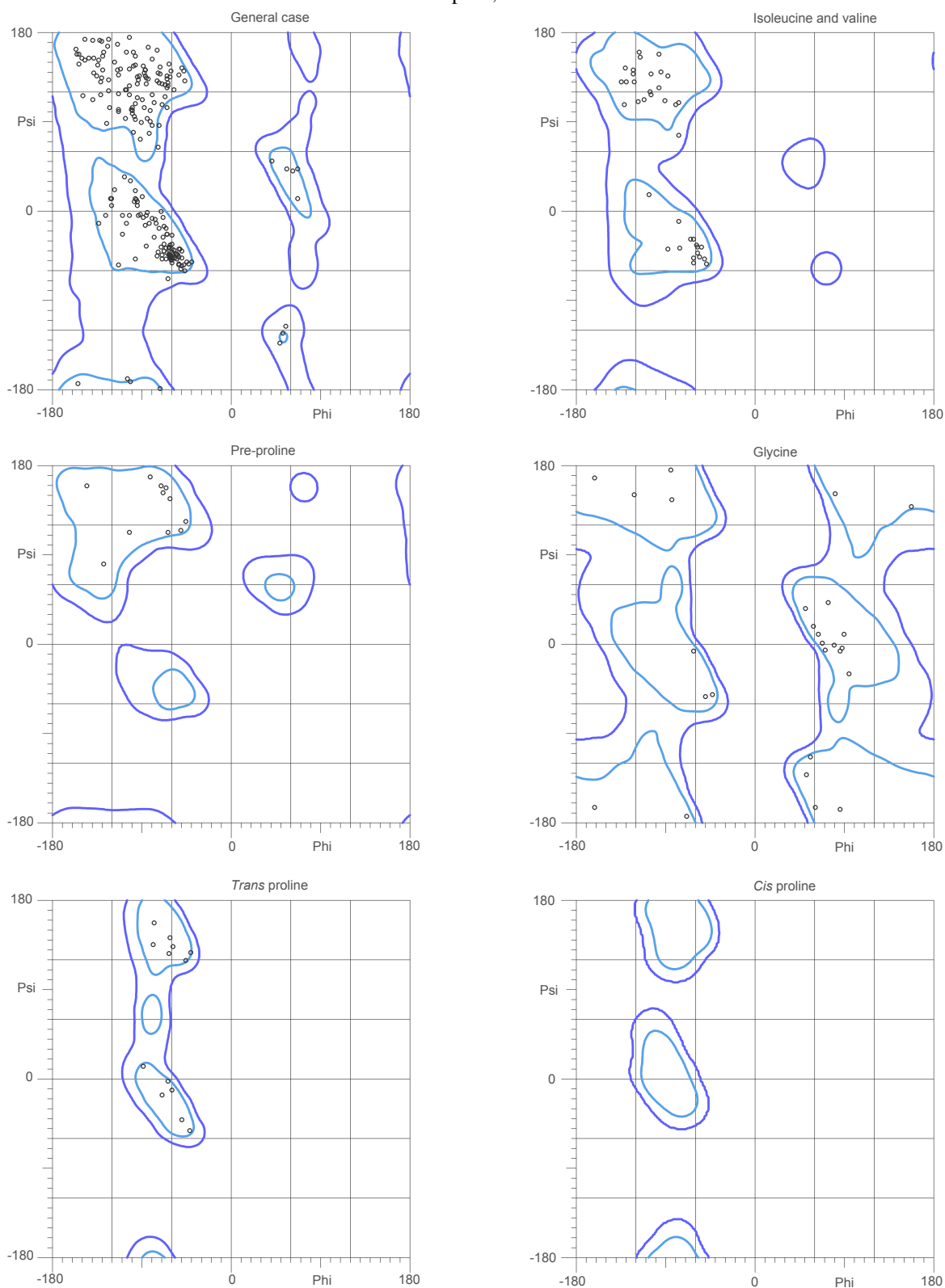


MolProbity Ramachandran analysis

7kx5.H.pdb, model 1



97.0% (292/301) of all residues were in favored (98%) regions.

100.0% (301/301) of all residues were in allowed (>99.8%) regions.

There were no outliers.