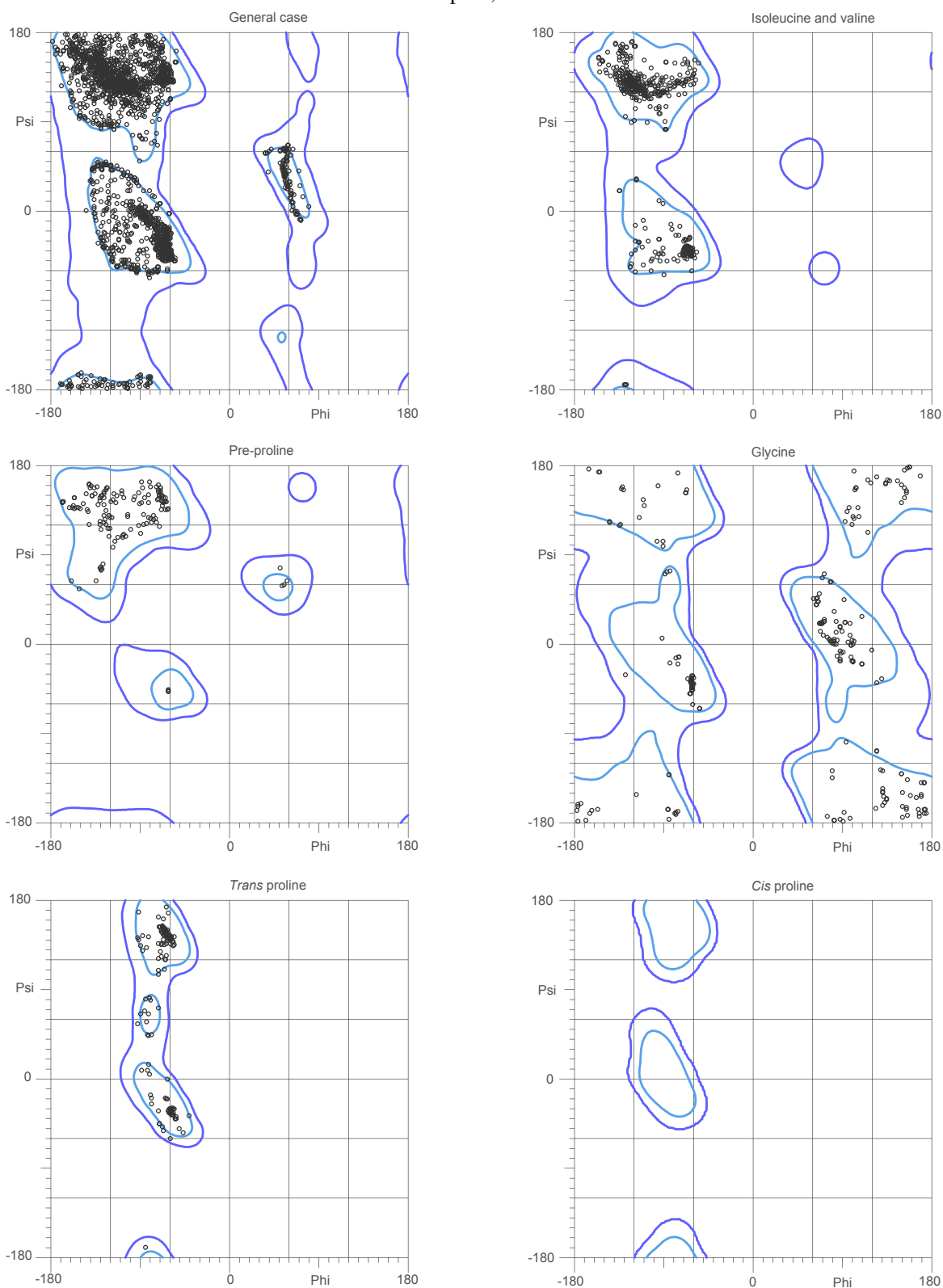


MolProbity Ramachandran analysis

7a25.H.pdb, model 1



95.8% (3294/3440) of all residues were in favored (98%) regions.

100.0% (3440/3440) of all residues were in allowed (>99.8%) regions.

There were no outliers.