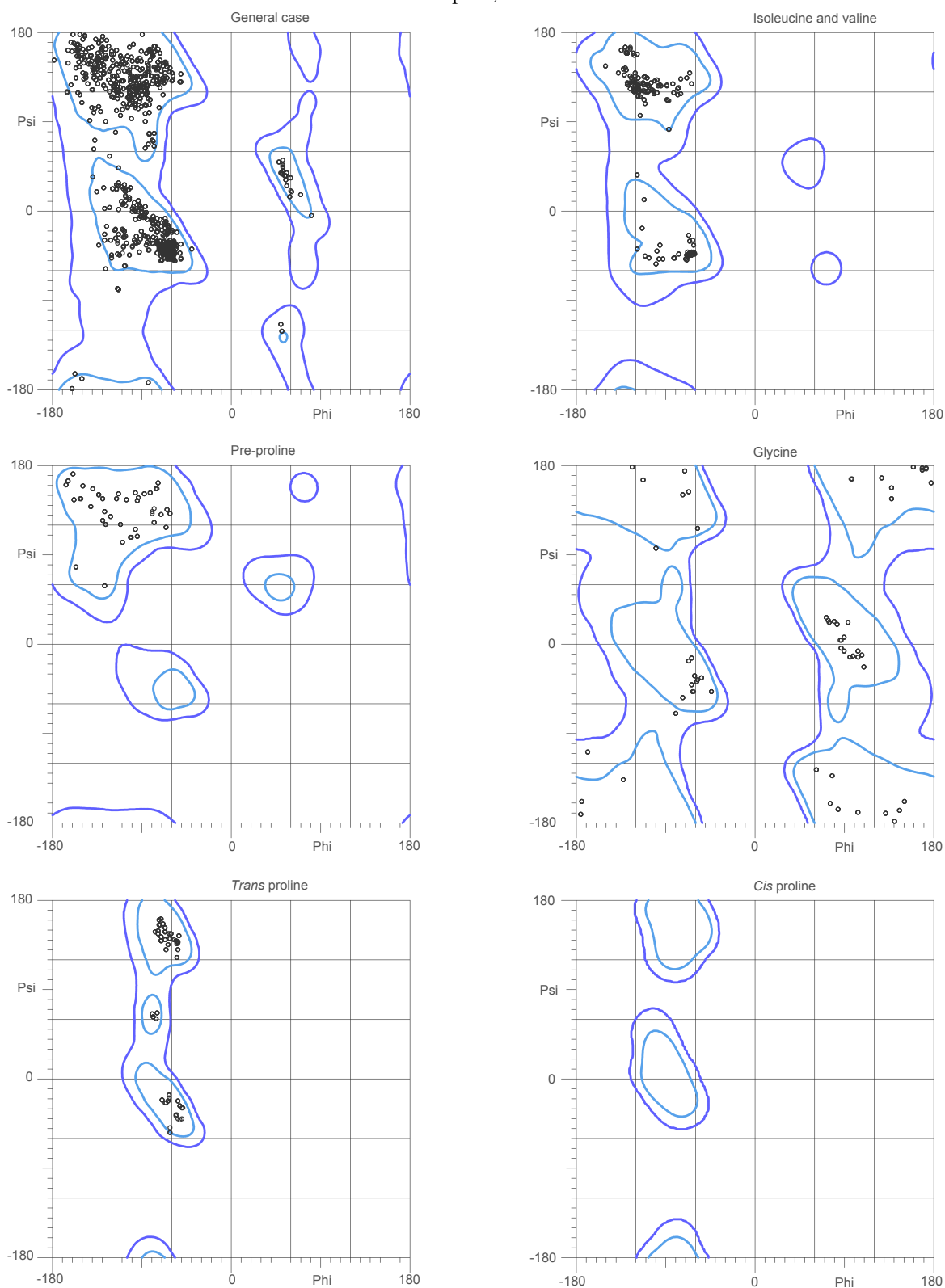


MolProbity Ramachandran analysis

6x79.H.pdb, model 1



97.5% (2692/2760) of all residues were in favored (98%) regions.

100.0% (2760/2760) of all residues were in allowed (>99.8%) regions.

There were no outliers.