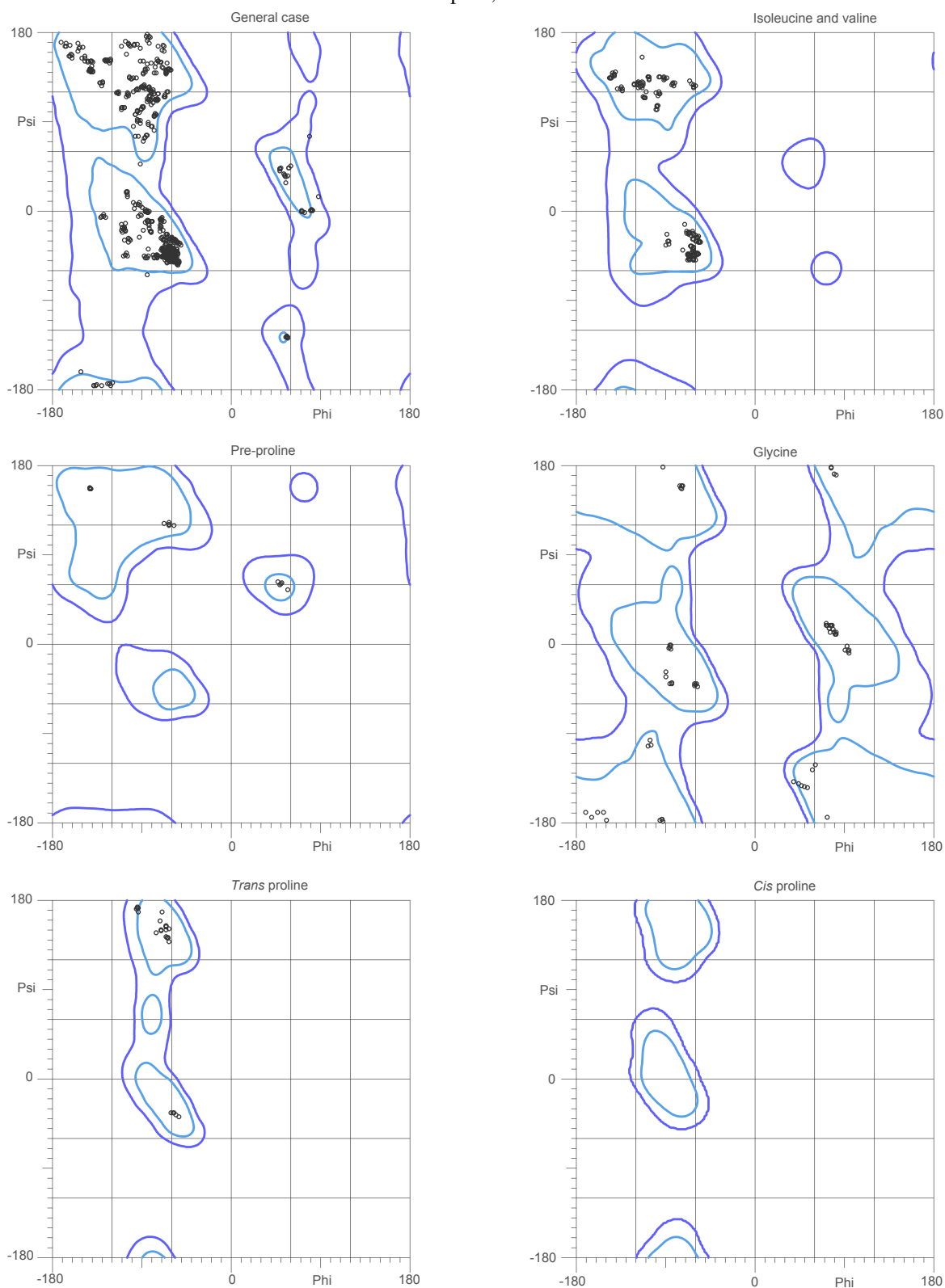


MolProbity Ramachandran analysis

7bf5.H.pdb, model 1



97.6% (831/851) of all residues were in favored (98%) regions.

100.0% (851/851) of all residues were in allowed (>99.8%) regions.

There were no outliers.