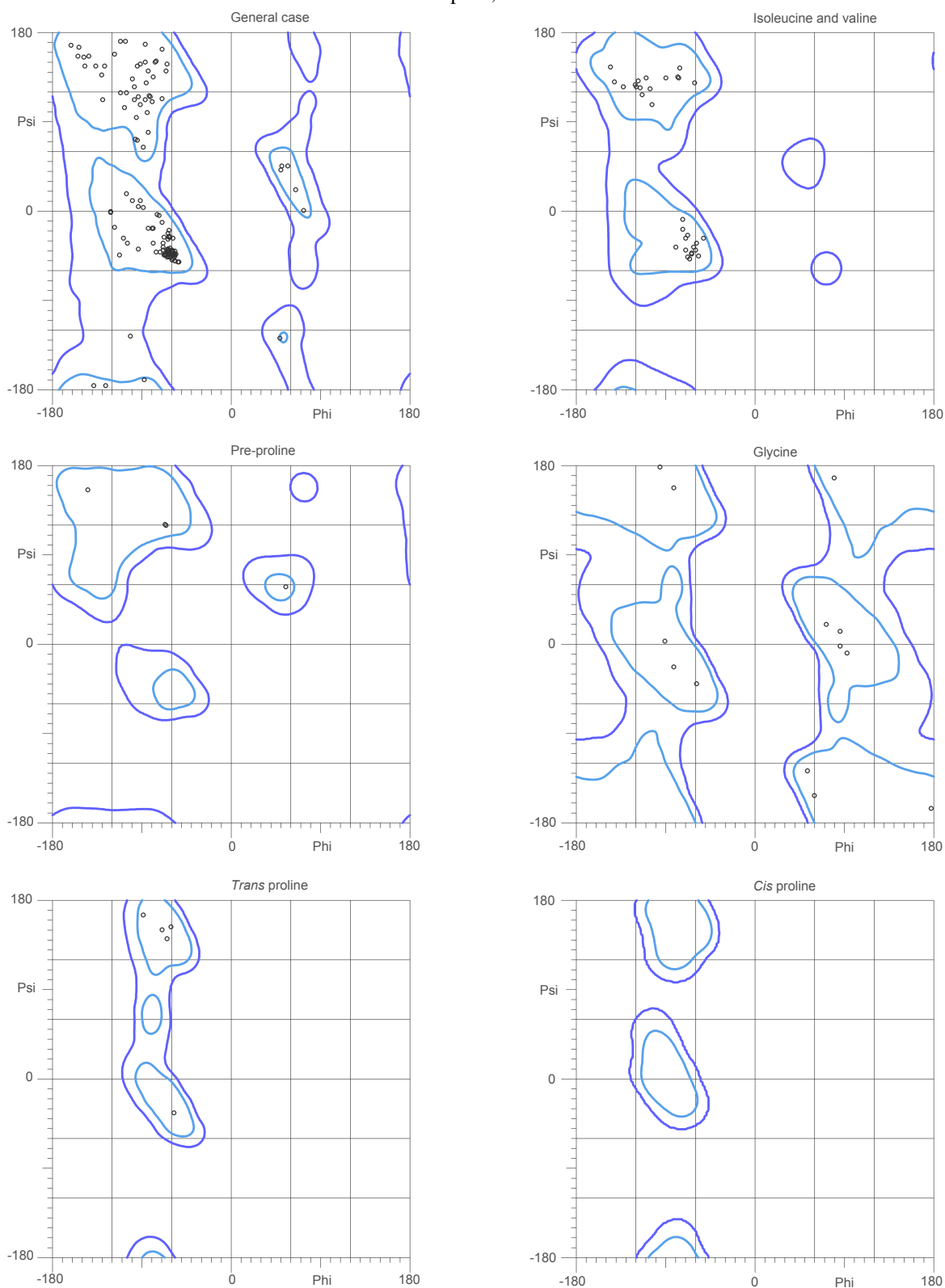


MolProbity Ramachandran analysis

7bf4.H.pdb, model 1



99.4% (178/179) of all residues were in favored (98%) regions.

100.0% (179/179) of all residues were in allowed (>99.8%) regions.

There were no outliers.