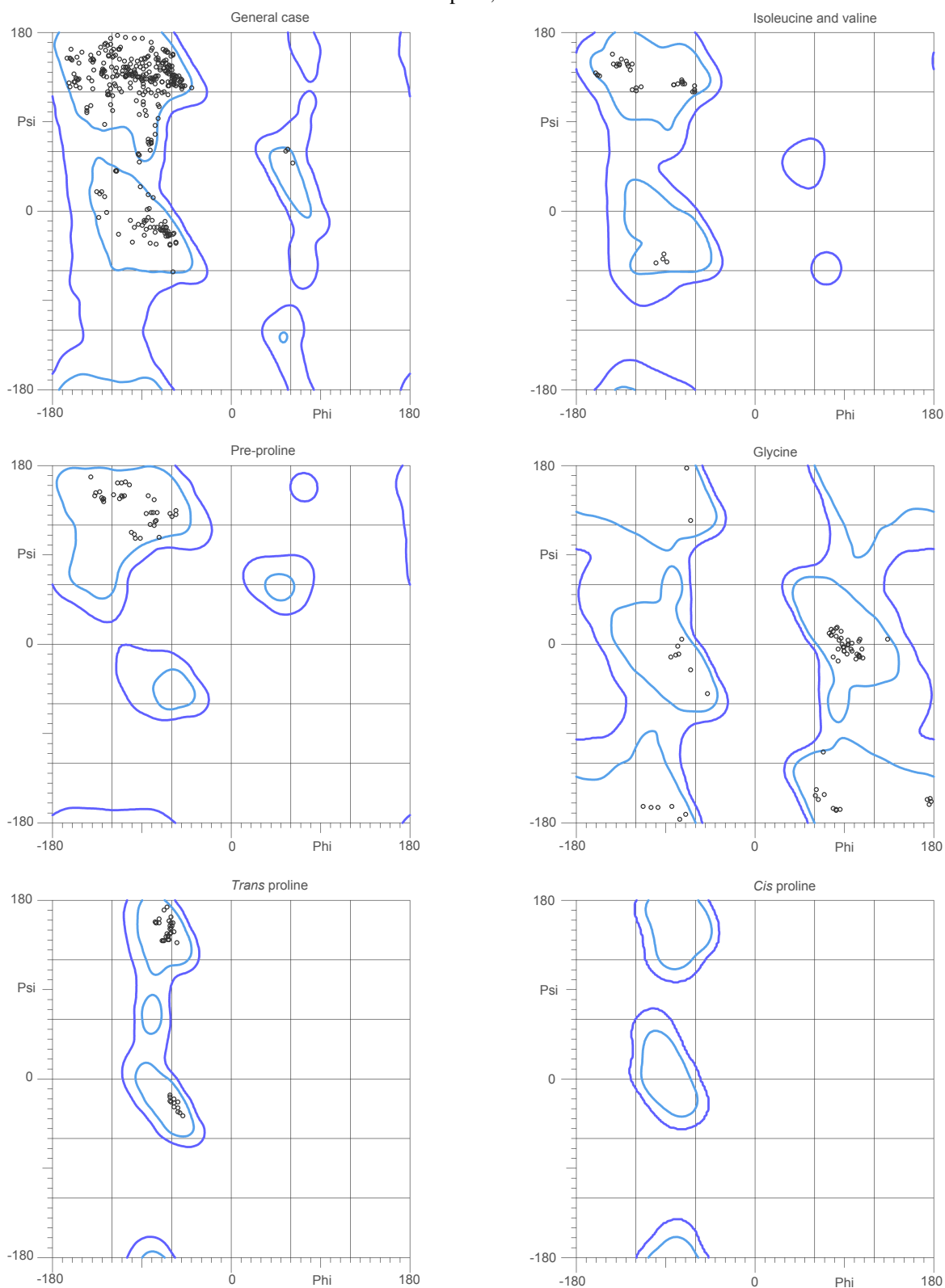


MolProbity Ramachandran analysis

7cdz.H.pdb, model 1



99.4% (497/500) of all residues were in favored (98%) regions.

100.0% (500/500) of all residues were in allowed (>99.8%) regions.

There were no outliers.