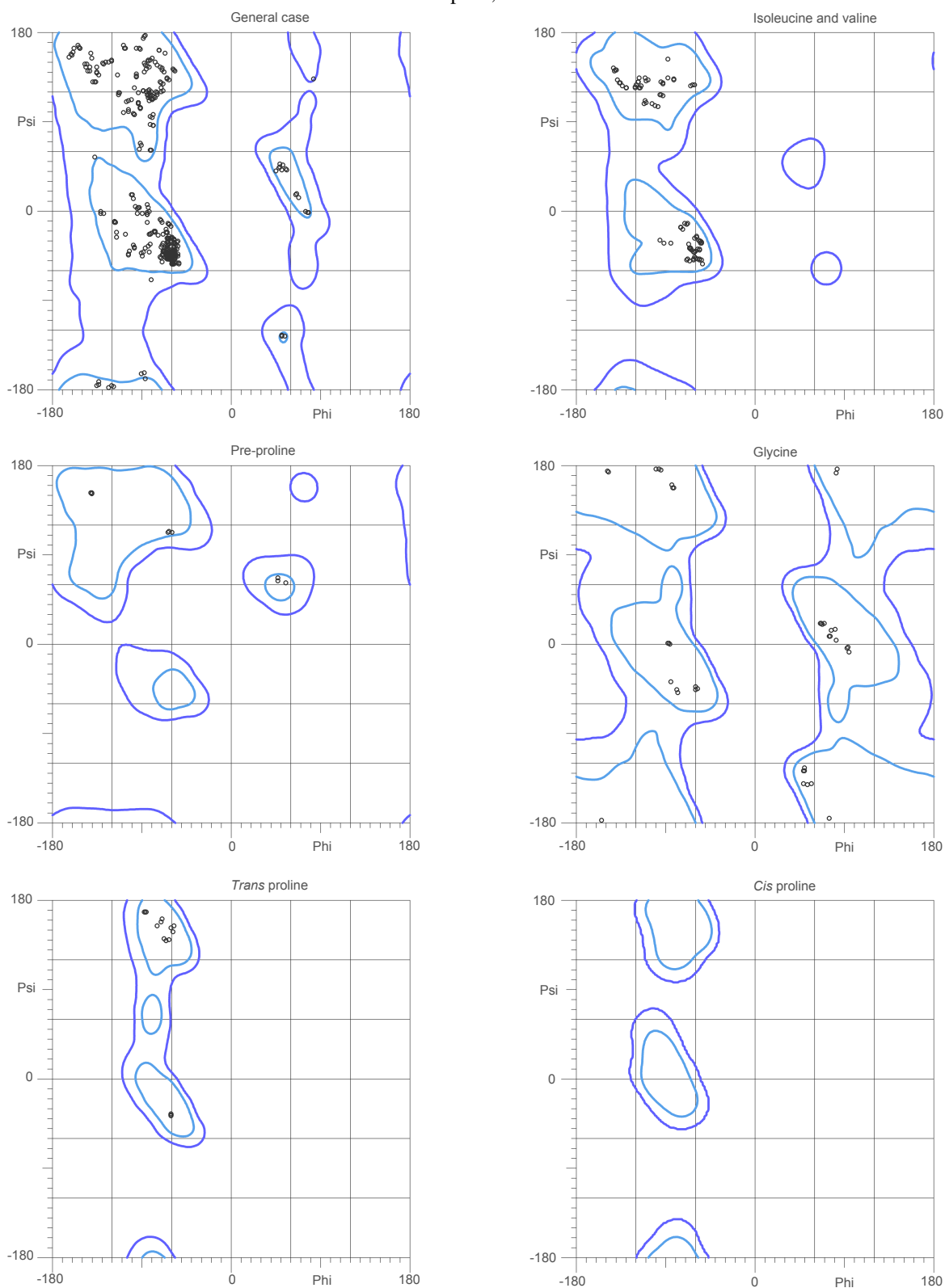


MolProbity Ramachandran analysis

7bf6.H.pdb, model 1



99.0% (496/501) of all residues were in favored (98%) regions.

100.0% (501/501) of all residues were in allowed (>99.8%) regions.

There were no outliers.