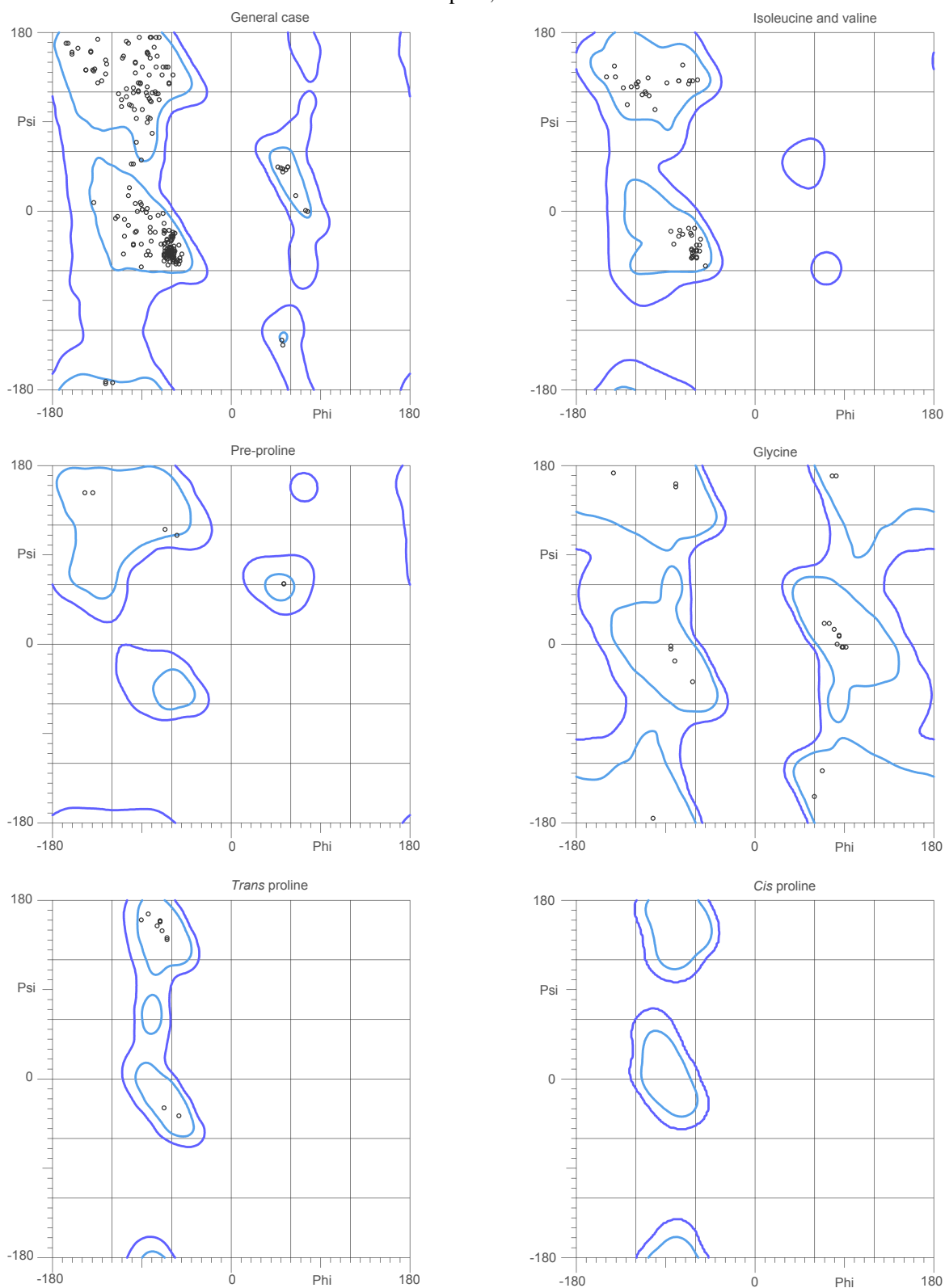


MolProbity Ramachandran analysis

7kr0.H.pdb, model 1



99.0% (305/308) of all residues were in favored (98%) regions.

100.0% (308/308) of all residues were in allowed (>99.8%) regions.

There were no outliers.