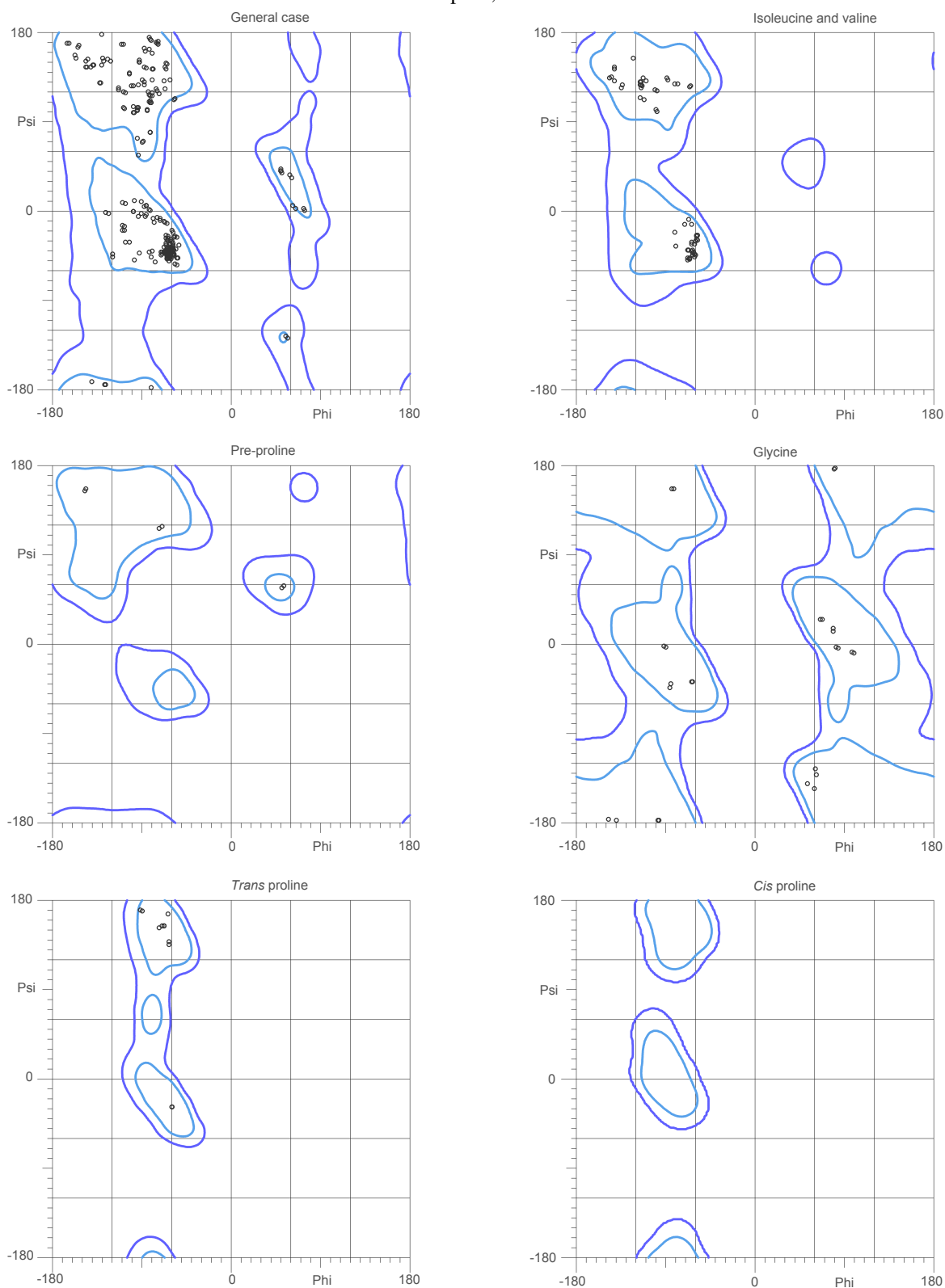


MolProbity Ramachandran analysis

5s2h.H.pdb, model 1



99.1% (341/344) of all residues were in favored (98%) regions.

100.0% (344/344) of all residues were in allowed (>99.8%) regions.

There were no outliers.