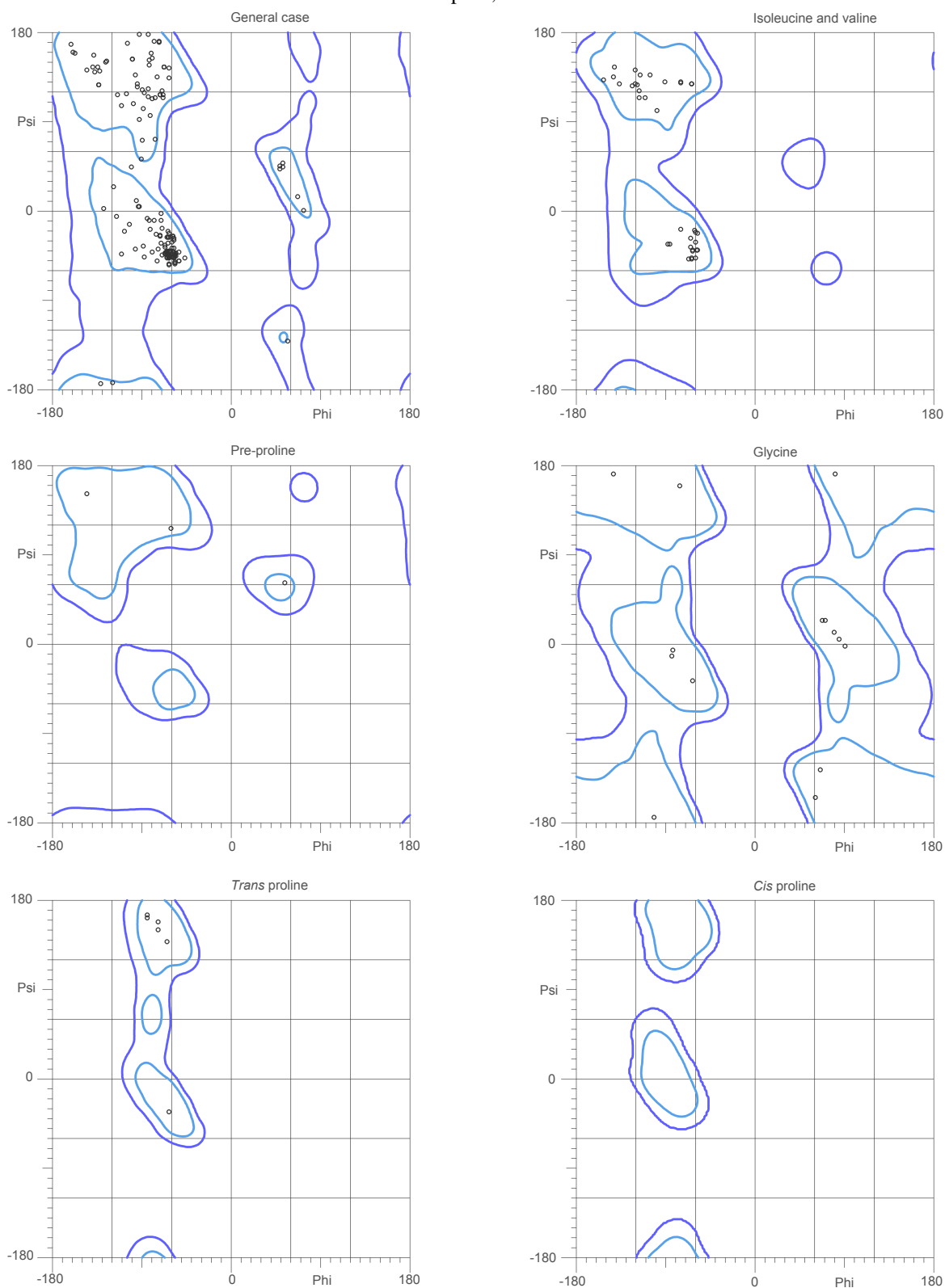


# MolProbity Ramachandran analysis

5rvu.H.pdb, model 1



99.0% (208/210) of all residues were in favored (98%) regions.

100.0% (210/210) of all residues were in allowed (>99.8%) regions.

There were no outliers.