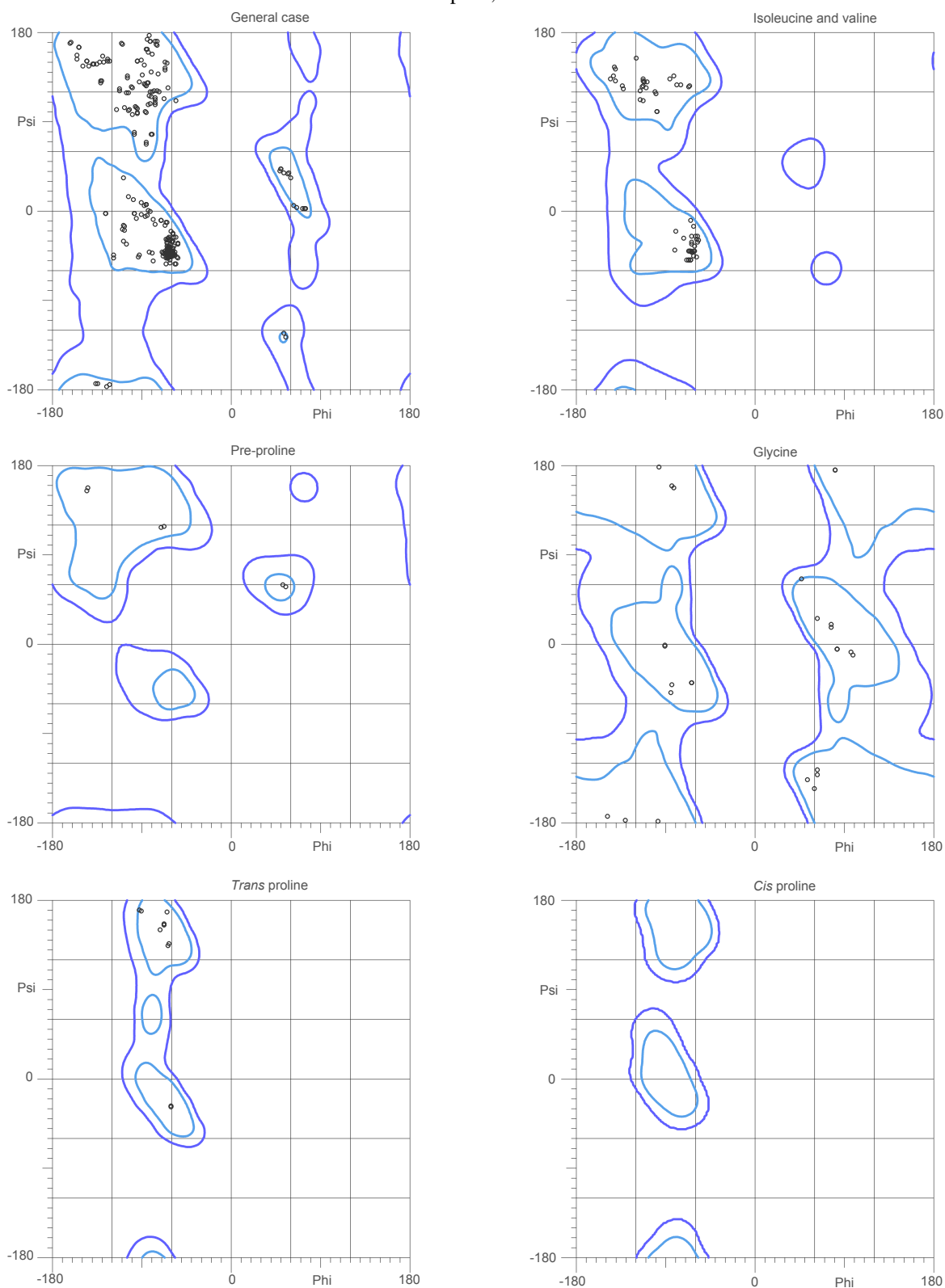


# MolProbity Ramachandran analysis

5s31.H.pdb, model 1



99.4% (347/349) of all residues were in favored (98%) regions.

100.0% (349/349) of all residues were in allowed (>99.8%) regions.

There were no outliers.