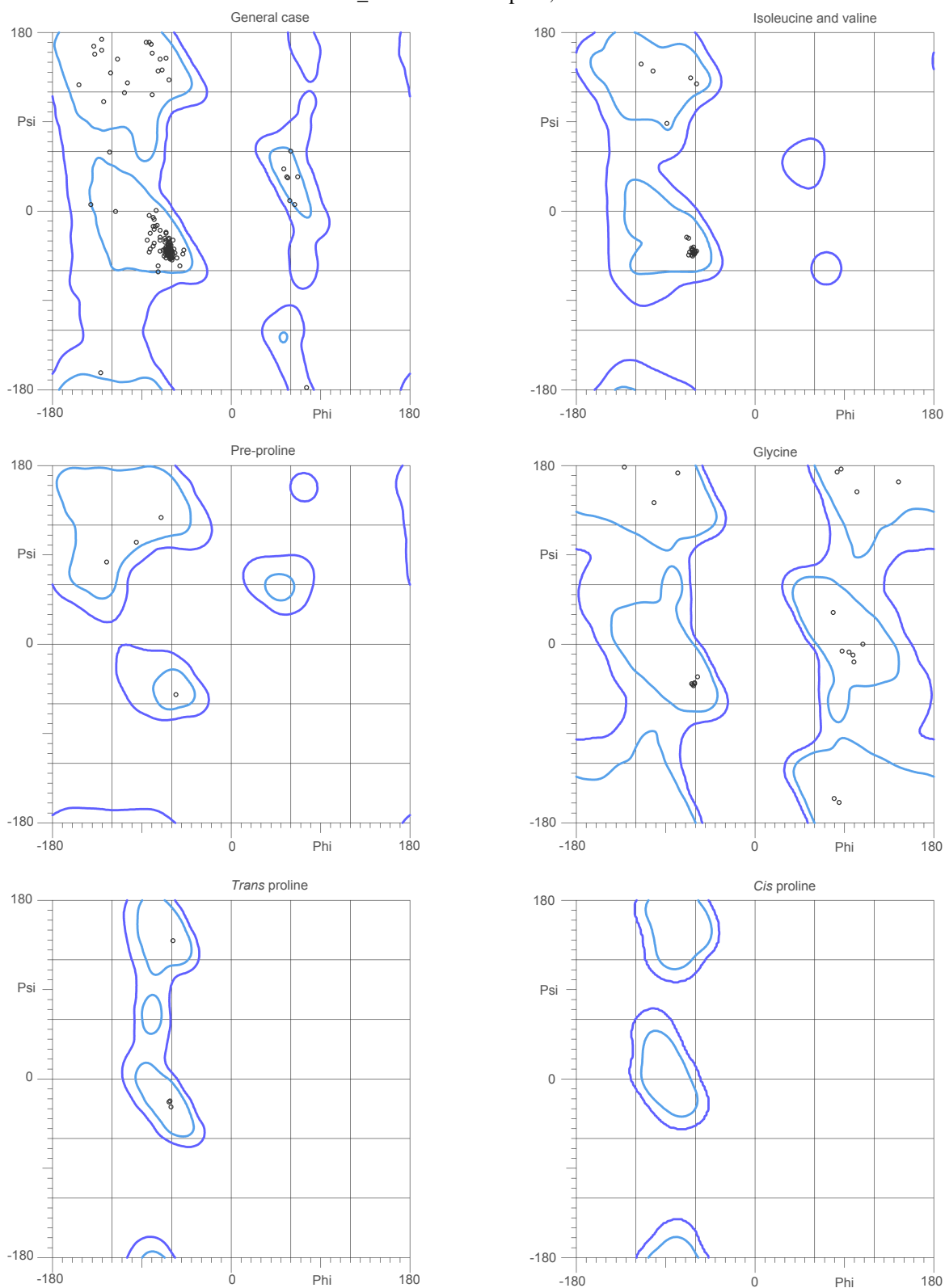


# MolProbity Ramachandran analysis

1531\_.B99990030.pdb, model 1



96,7% (177/183) of all residues were in favored (98%) regions.

100,0% (183/183) of all residues were in allowed (>99.8%) regions.

There were no outliers.