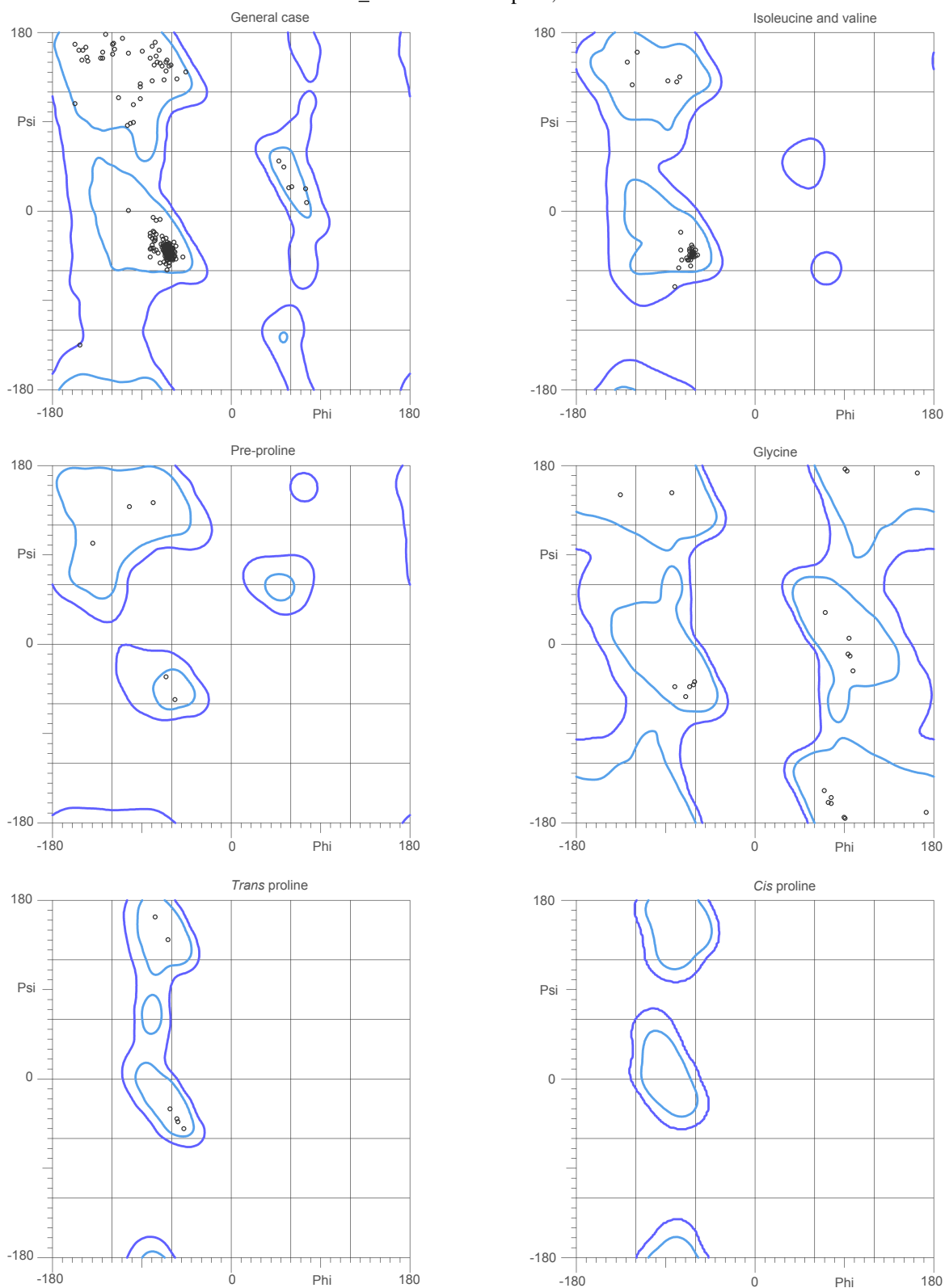


# MolProbity Ramachandran analysis

1axn\_.B99990006.pdb, model 1



99,1% (318/321) of all residues were in favored (98%) regions.

100,0% (321/321) of all residues were in allowed (>99.8%) regions.

There were no outliers.