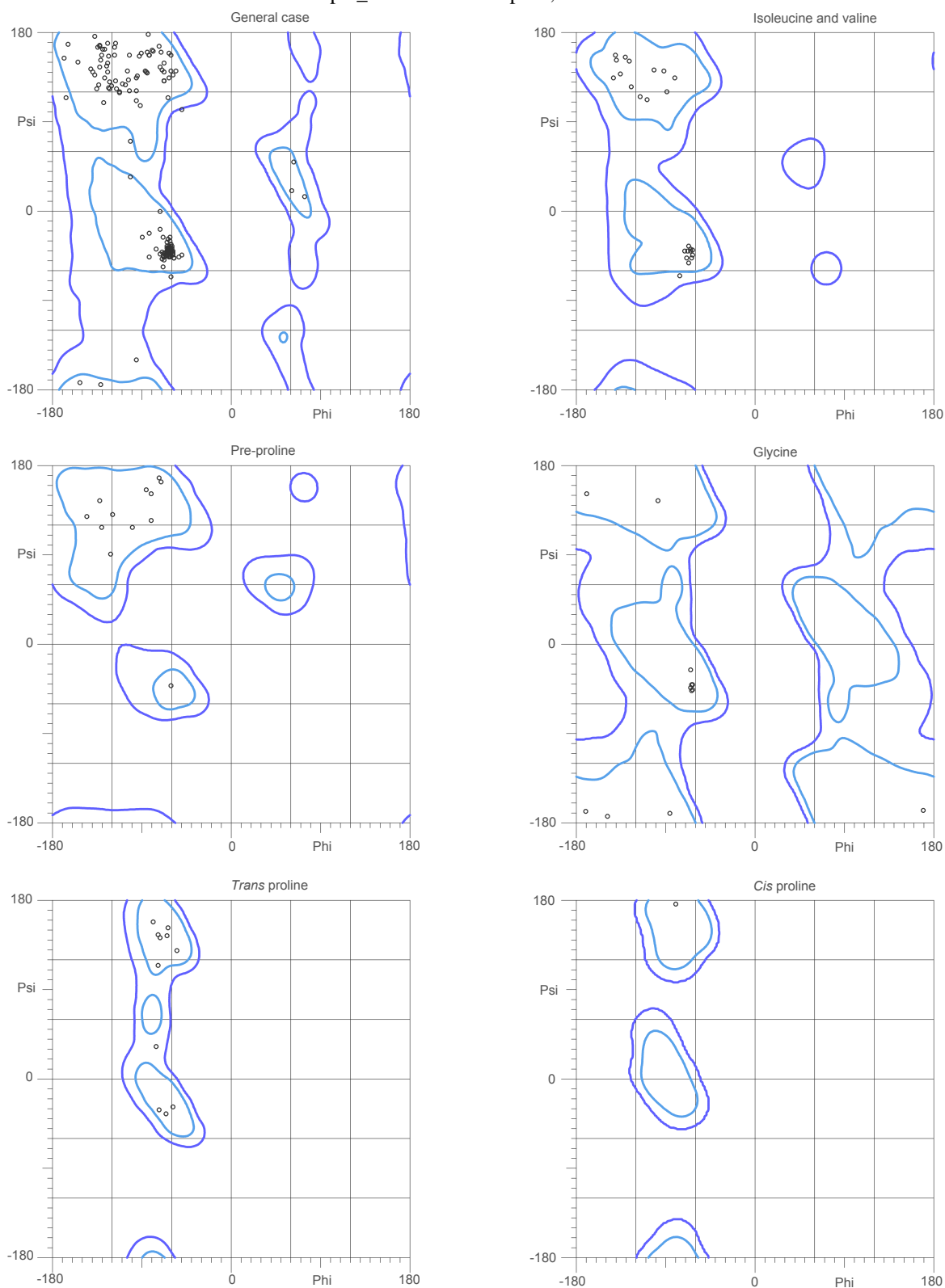


# MolProbity Ramachandran analysis

1ap8\_.B99990023.pdb, model 1



96,7% (204/211) of all residues were in favored (98%) regions.

100,0% (211/211) of all residues were in allowed (>99.8%) regions.

There were no outliers.