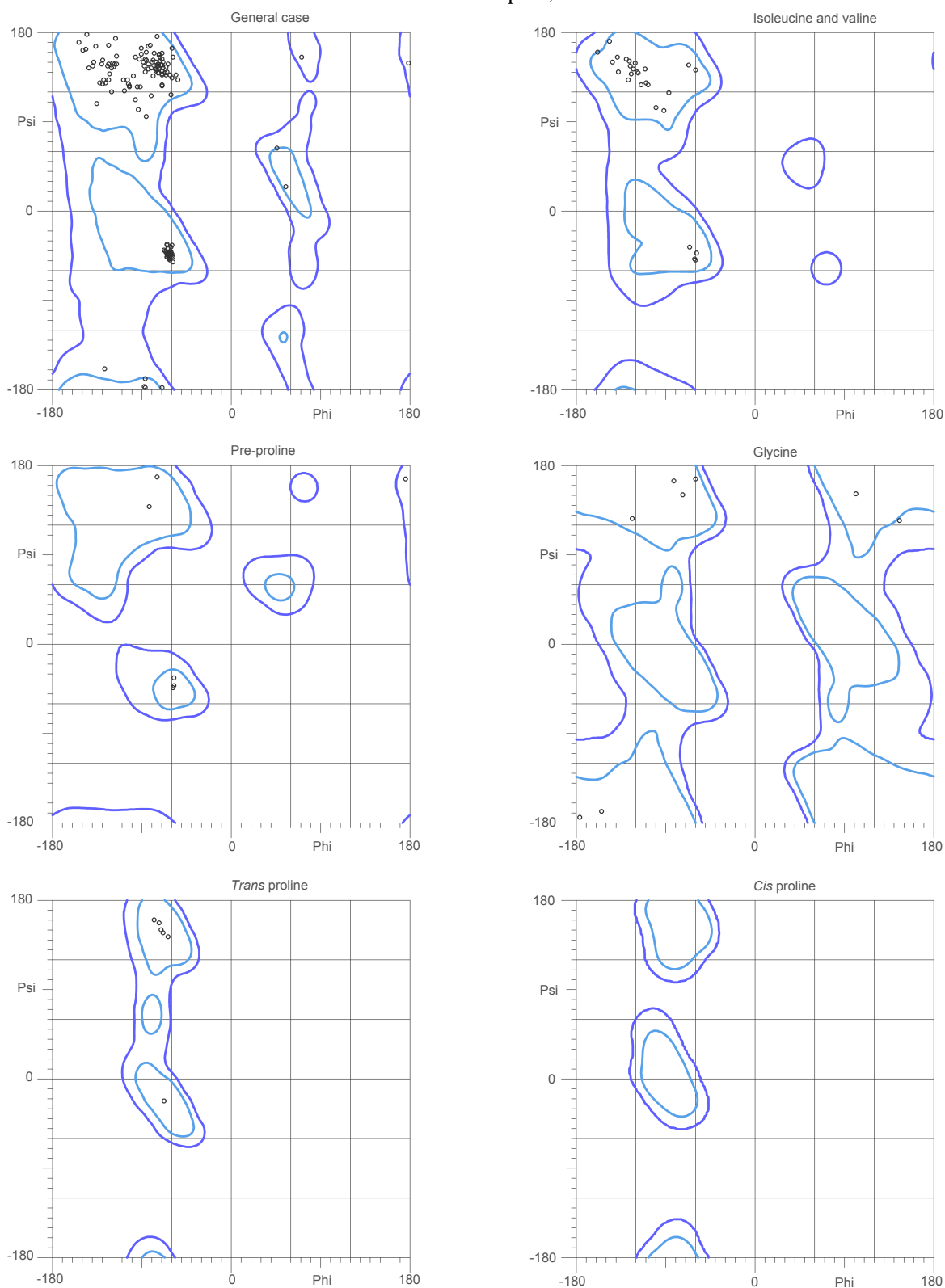


# MolProbity Ramachandran analysis

1aisB.B99990046.pdb, model 1



95,8% (183/191) of all residues were in favored (98%) regions.

100,0% (191/191) of all residues were in allowed (>99.8%) regions.

There were no outliers.