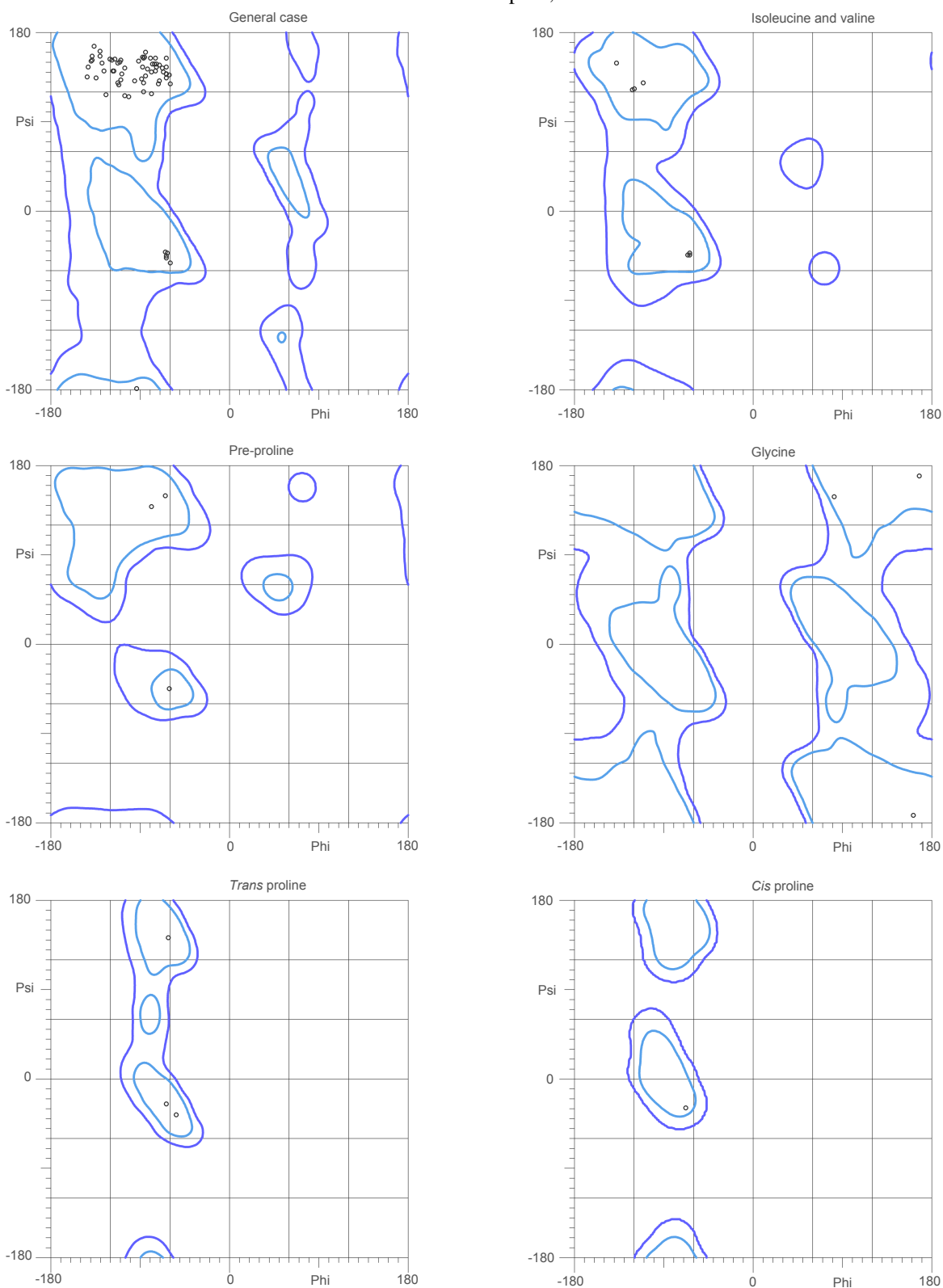


# MolProbity Ramachandran analysis

1aliA.B99990003.pdb, model 1



100,0% (83/83) of all residues were in favored (98%) regions.

100,0% (83/83) of all residues were in allowed (>99.8%) regions.

There were no outliers.