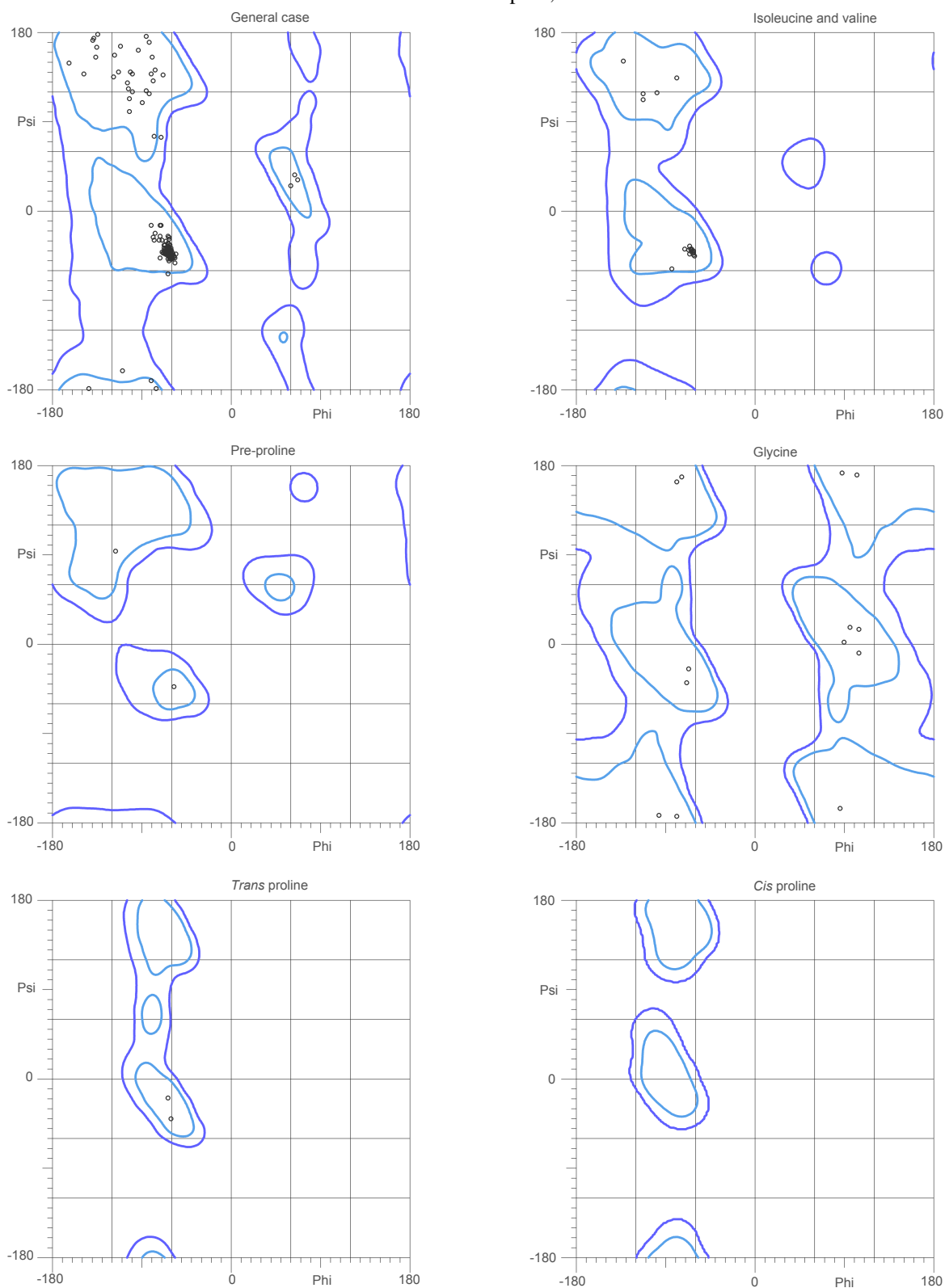


# MolProbity Ramachandran analysis

1alvA.B99990049.pdb, model 1



98,2% (168/171) of all residues were in favored (98%) regions.

100,0% (171/171) of all residues were in allowed (>99.8%) regions.

There were no outliers.