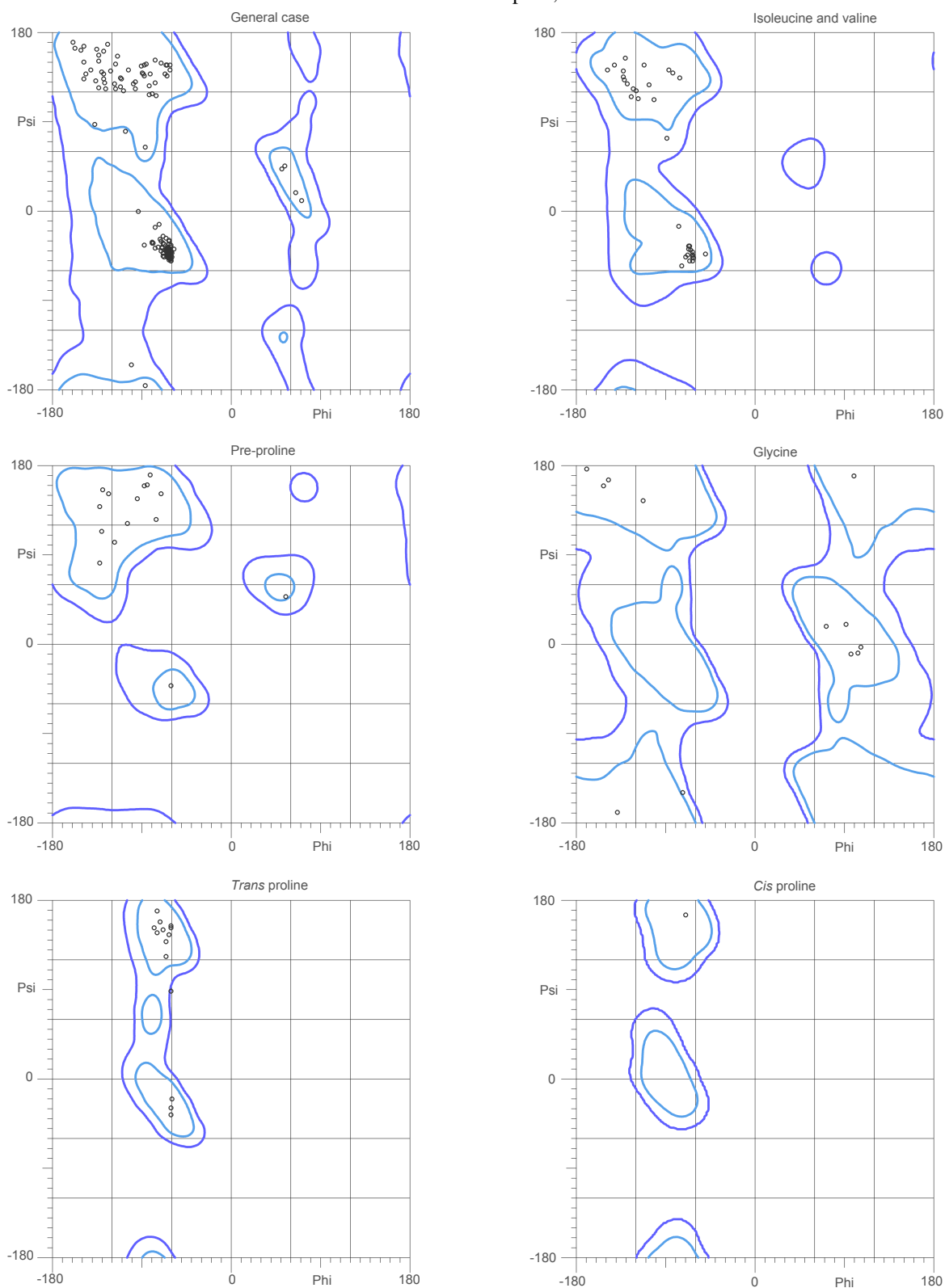


# MolProbity Ramachandran analysis

1a2zA.B99990013.pdb, model 1



97,7% (213/218) of all residues were in favored (98%) regions.

100,0% (218/218) of all residues were in allowed (>99.8%) regions.

There were no outliers.