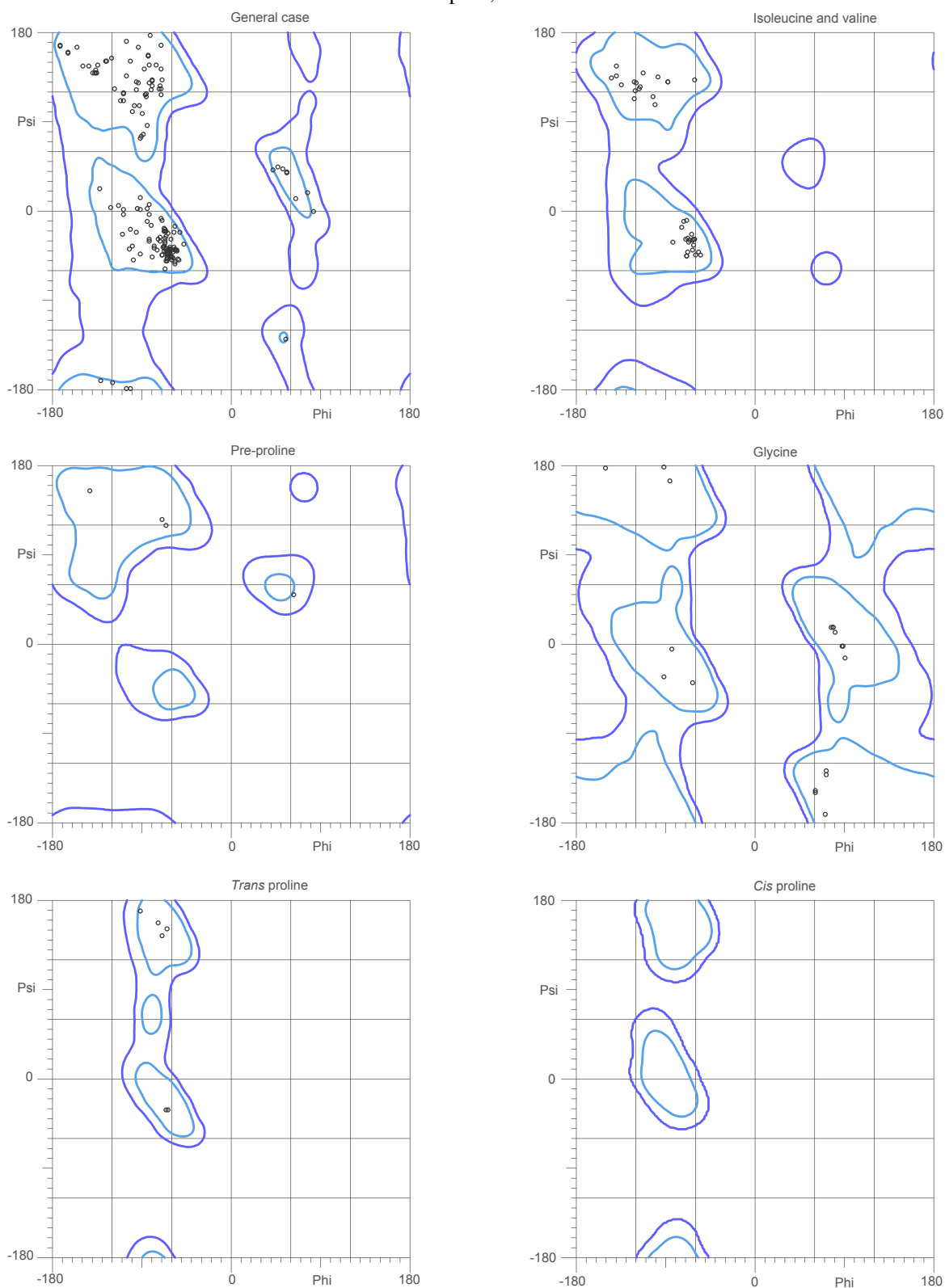


# MolProbity Ramachandran analysis

7kr1.H.pdb, model 1



98.7% (229/232) of all residues were in favored (98%) regions.

100.0% (232/232) of all residues were in allowed (>99.8%) regions.

There were no outliers.