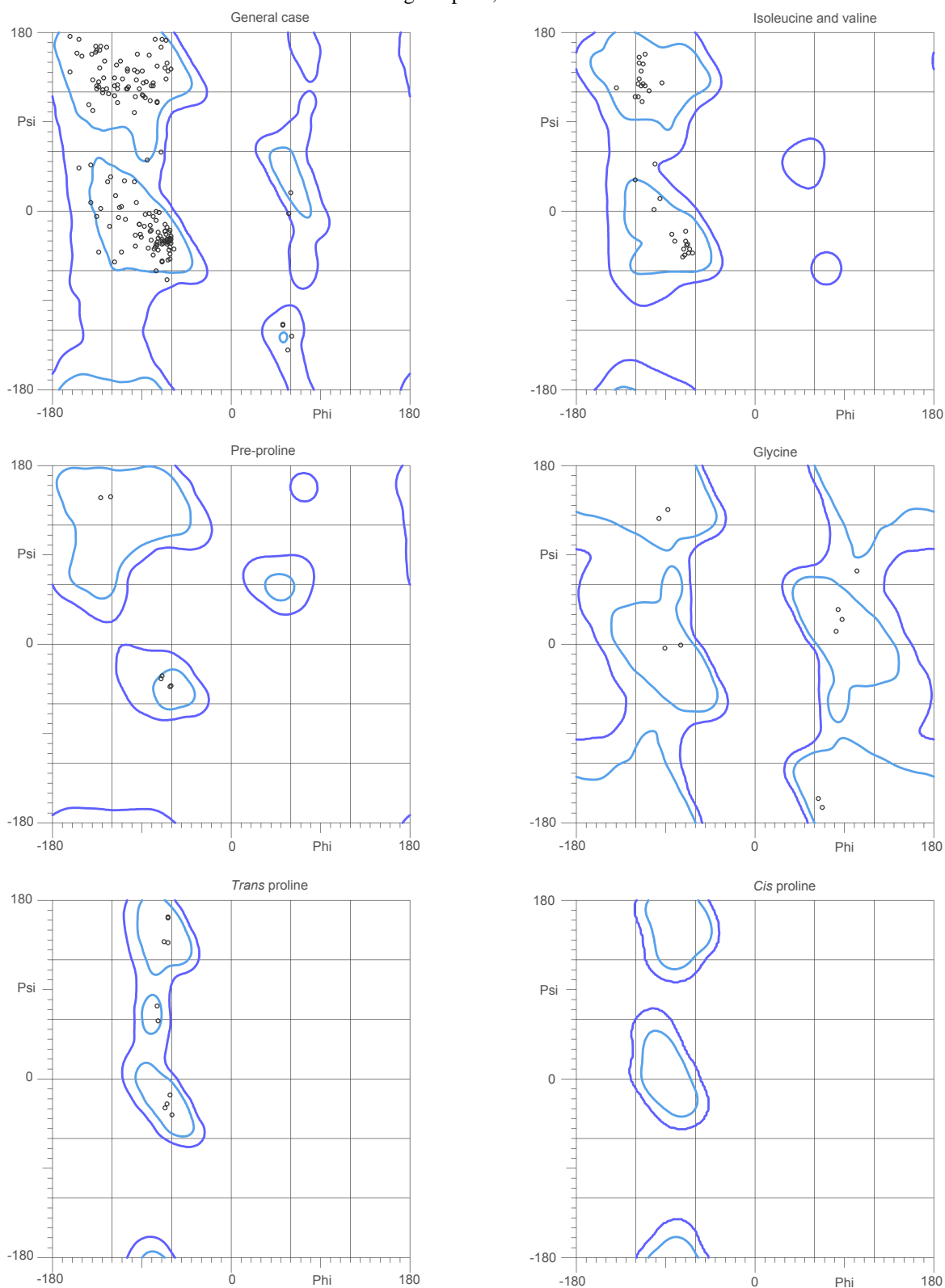


# MolProbity Ramachandran analysis

7kag.H.pdb, model 1



93.6% (204/218) 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.