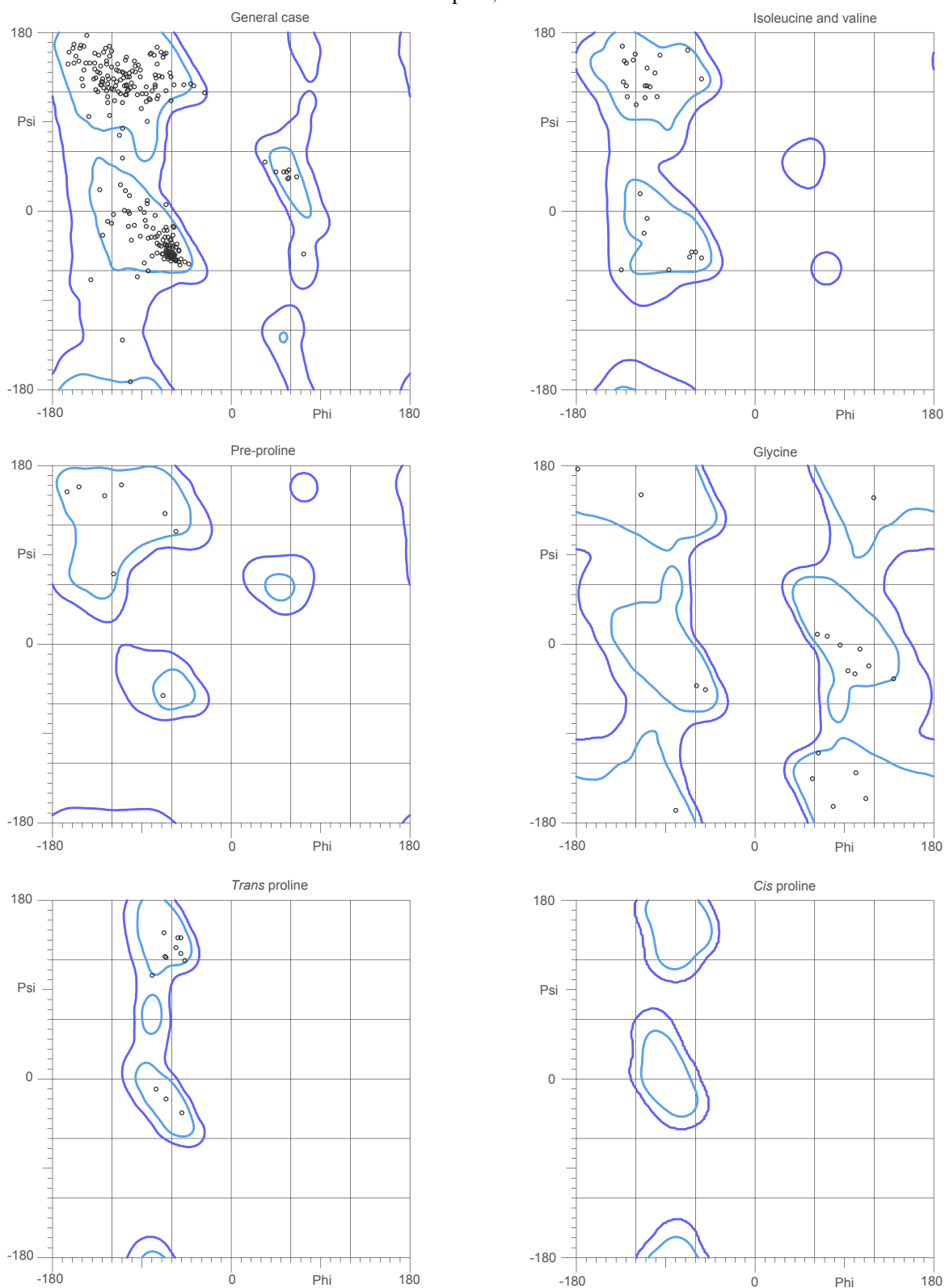


# MolProbity Ramachandran analysis

7kok.H.pdb, model 1



95.6% (304/318) of all residues were in favored (98%) regions.

100.0% (318/318) of all residues were in allowed (>99.8%) regions.

There were no outliers.