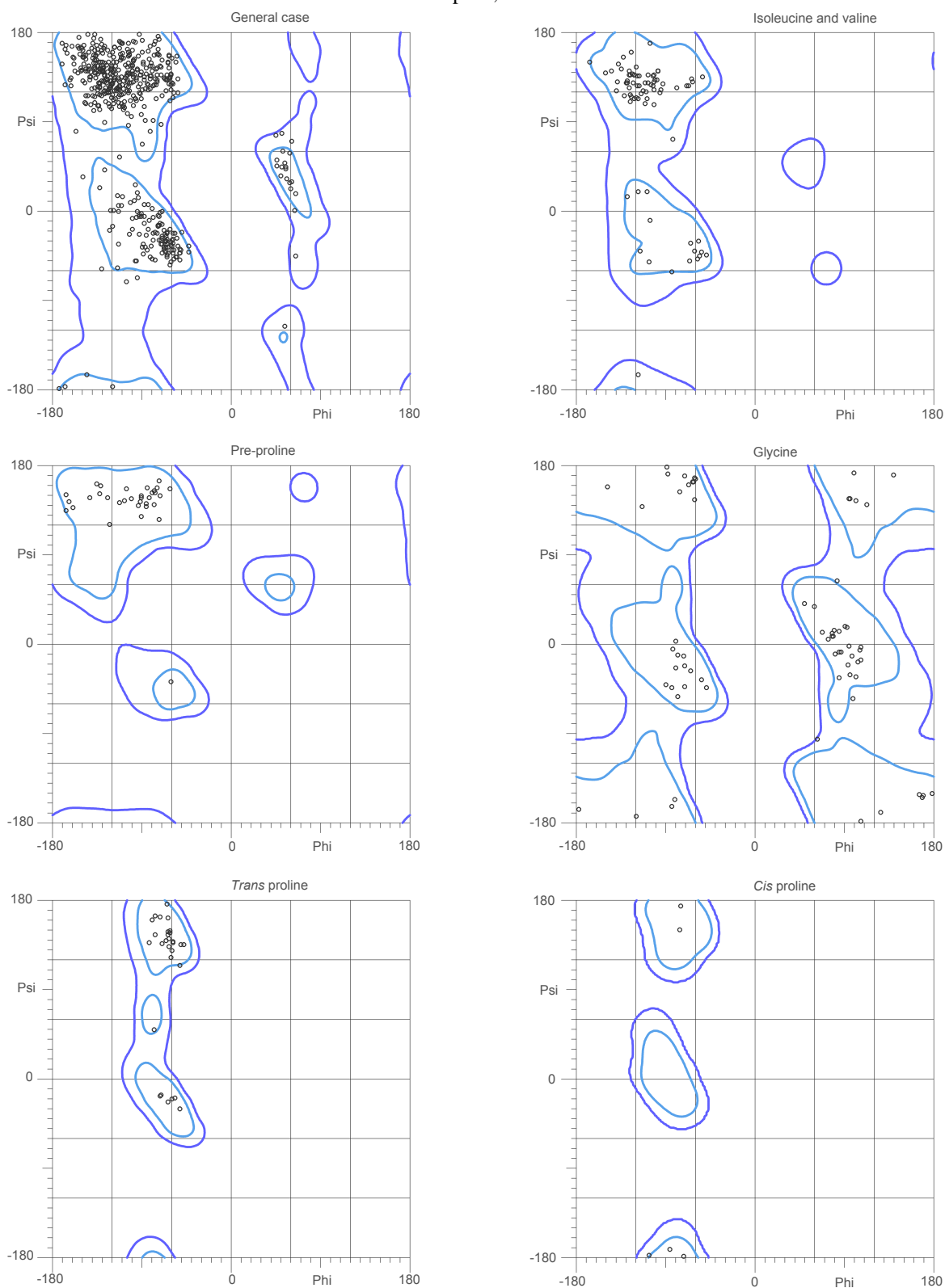


# MolProbity Ramachandran analysis

7kn6.H.pdb, model 1



96.5% (696/721) of all residues were in favored (98%) regions.

100.0% (721/721) of all residues were in allowed (>99.8%) regions.

There were no outliers.