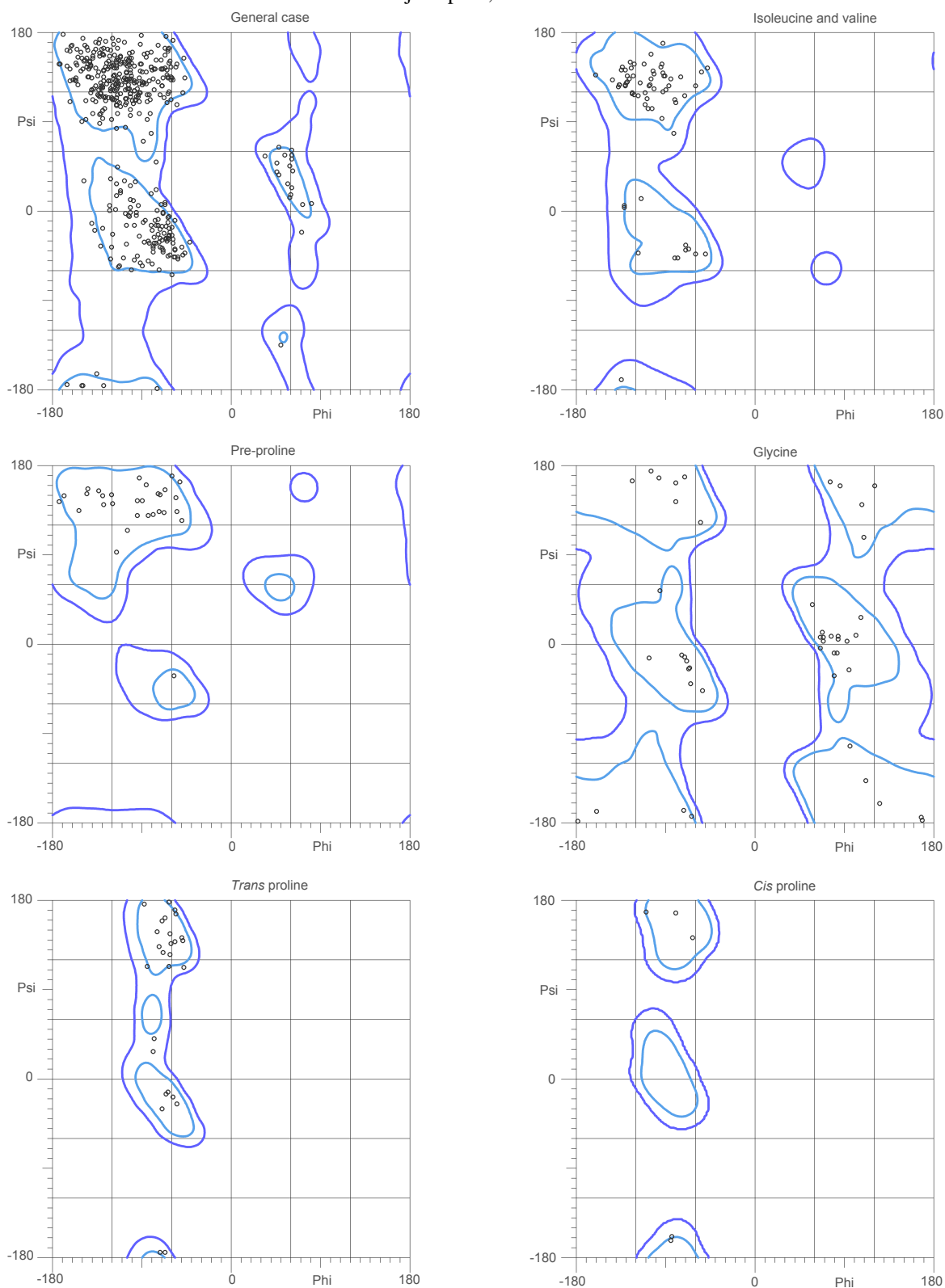


# MolProbity Ramachandran analysis

7cdj.H.pdb, model 1



91.7% (560/611) of all residues were in favored (98%) regions.

100.0% (611/611) of all residues were in allowed (>99.8%) regions.

There were no outliers.