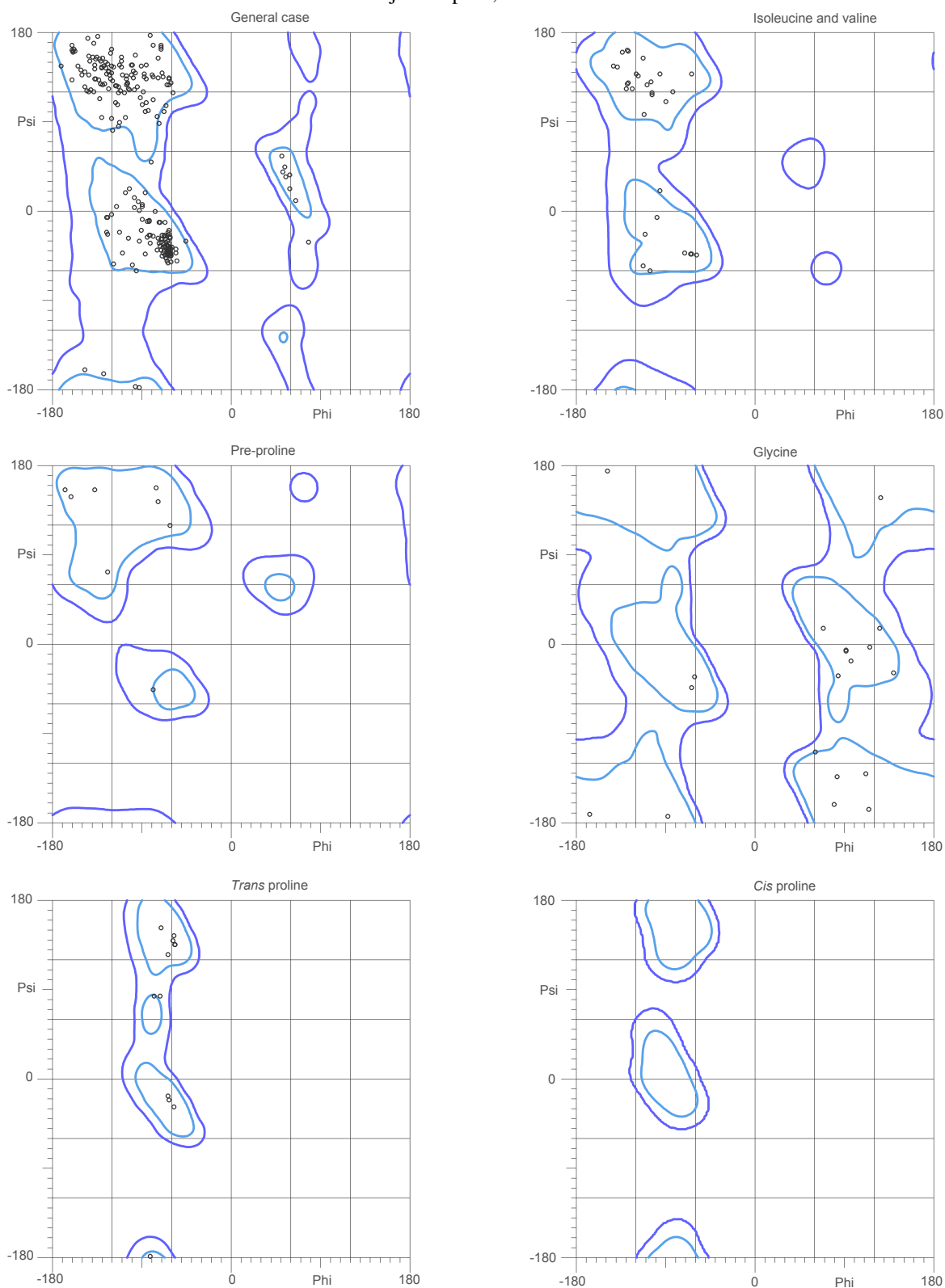


# MolProbity Ramachandran analysis

7cjm.H.pdb, model 1



95.5% (300/314) of all residues were in favored (98%) regions.

100.0% (314/314) of all residues were in allowed (>99.8%) regions.

There were no outliers.