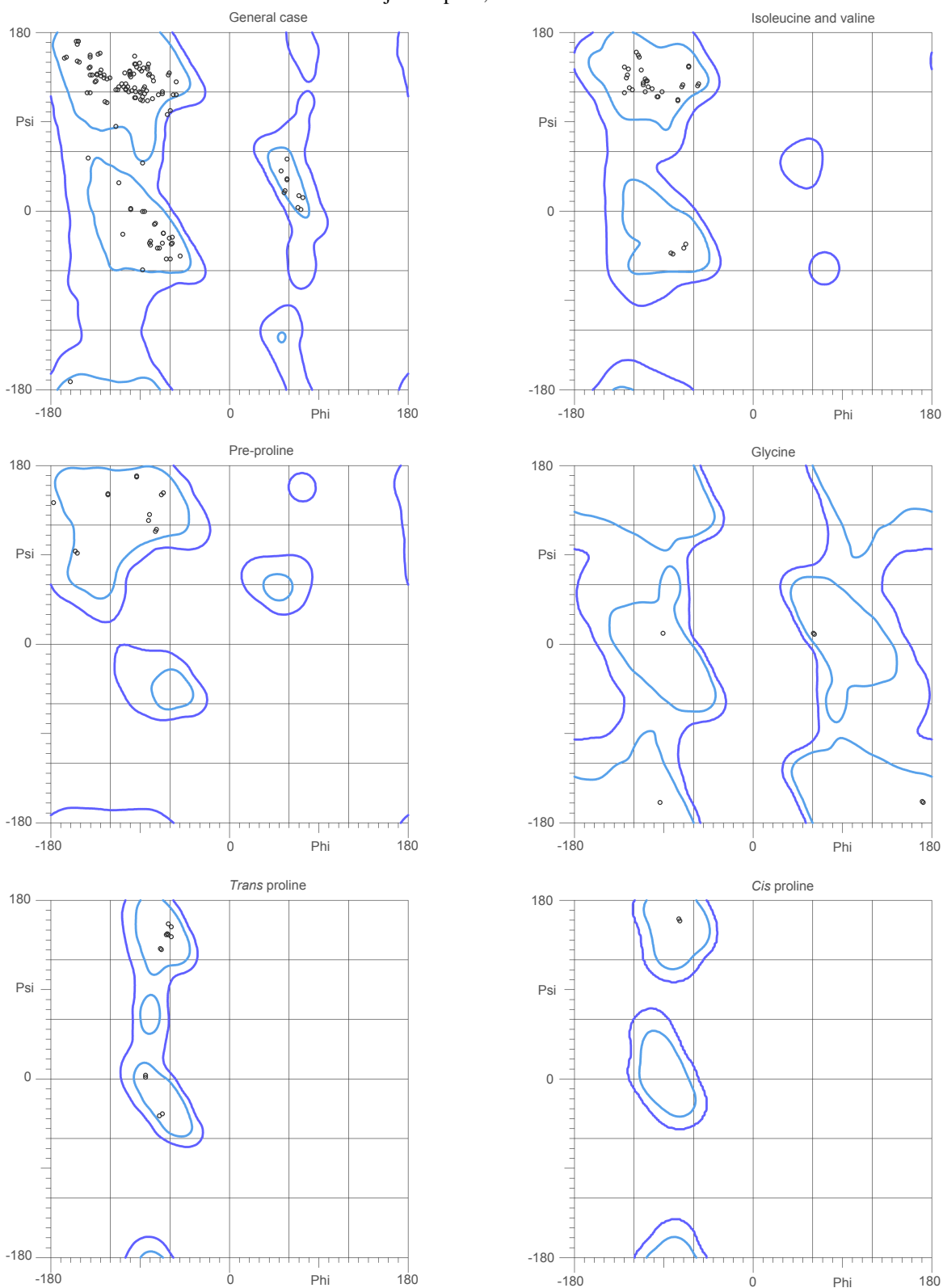


# MolProbity Ramachandran analysis

7jtl.H.pdb, model 1



97.4% (189/194) of all residues were in favored (98%) regions.

100.0% (194/194) of all residues were in allowed (>99.8%) regions.

There were no outliers.