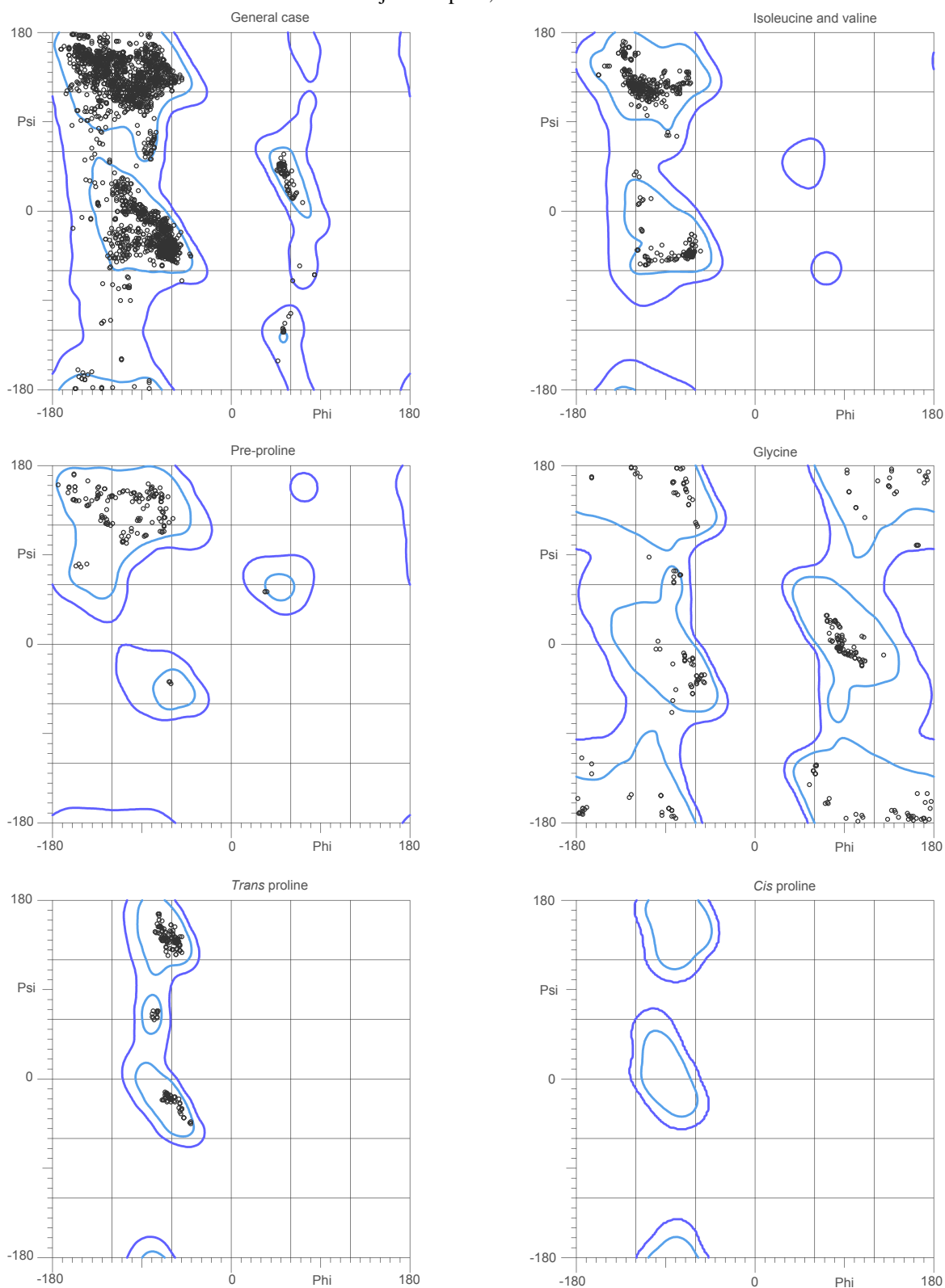


# MolProbity Ramachandran analysis

7jv4.H.pdb, model 1



97.5% (3386/3472) of all residues were in favored (98%) regions.

100.0% (3472/3472) of all residues were in allowed (>99.8%) regions.

There were no outliers.