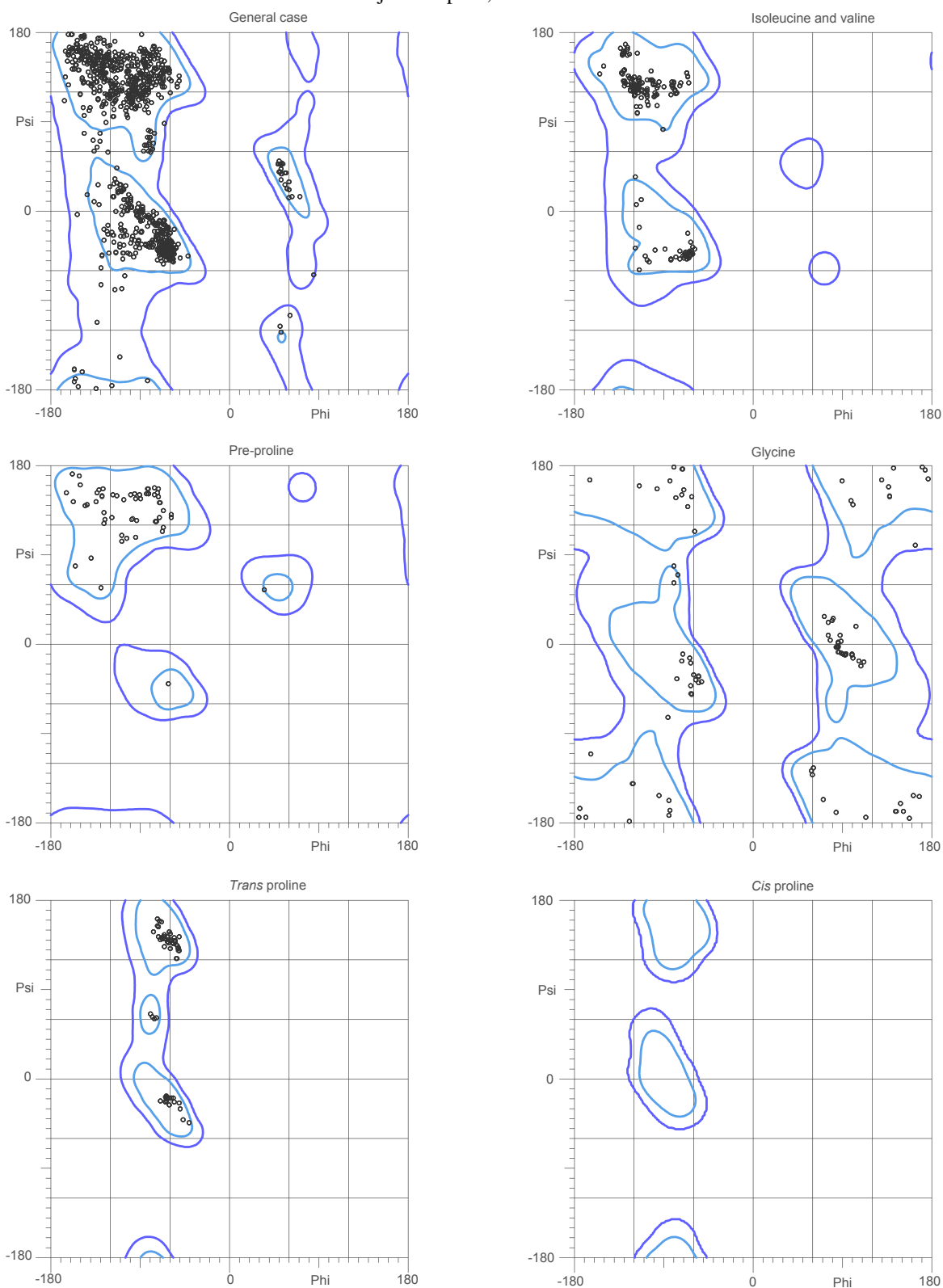


# MolProbity Ramachandran analysis

7jv6.H.pdb, model 1



97.2% (3429/3528) of all residues were in favored (98%) regions.

100.0% (3528/3528) of all residues were in allowed (>99.8%) regions.

There were no outliers.