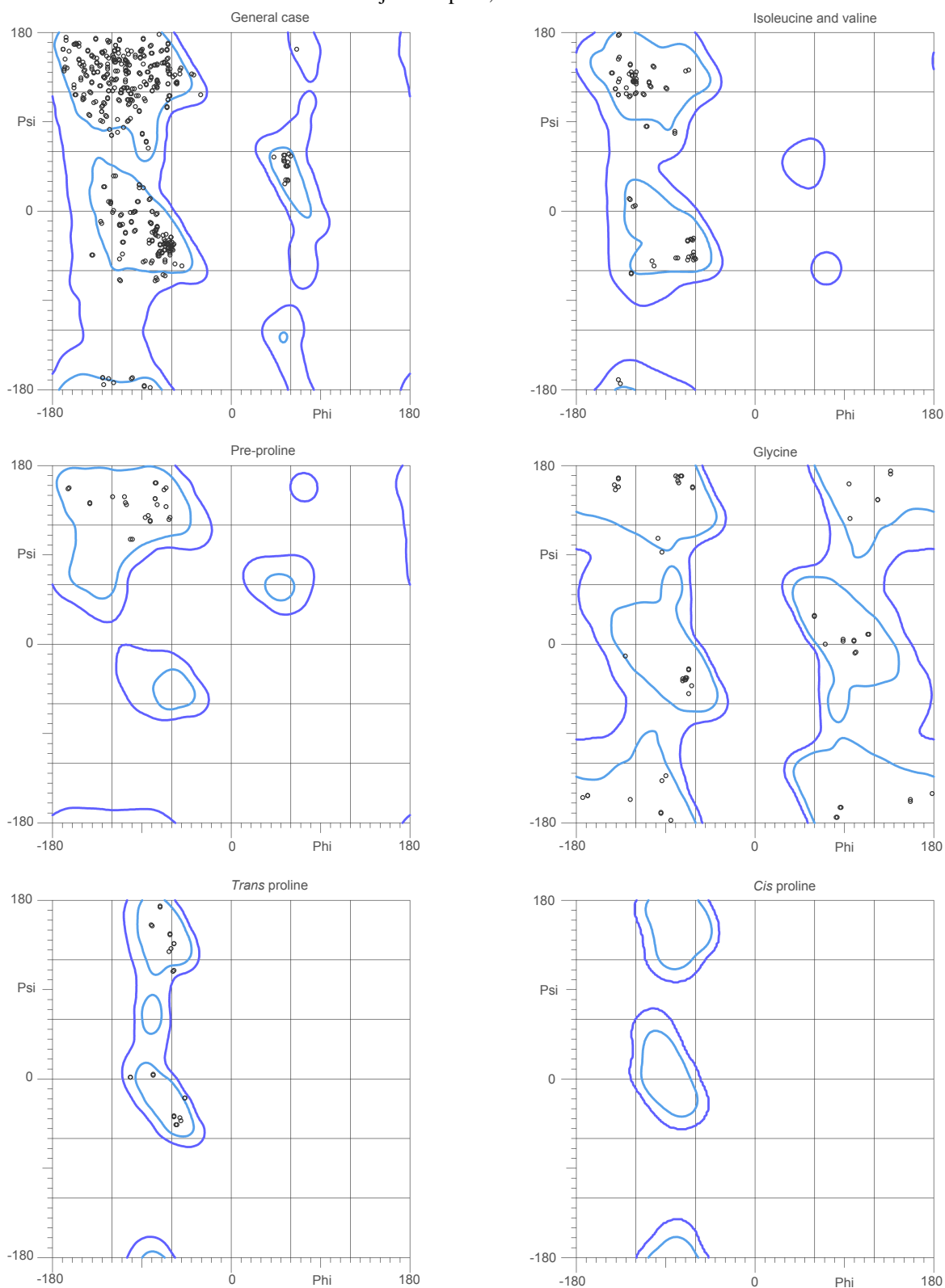


# MolProbity Ramachandran analysis

7jvb.H.pdb, model 1



94.0% (576/613) of all residues were in favored (98%) regions.

100.0% (613/613) of all residues were in allowed (>99.8%) regions.

There were no outliers.