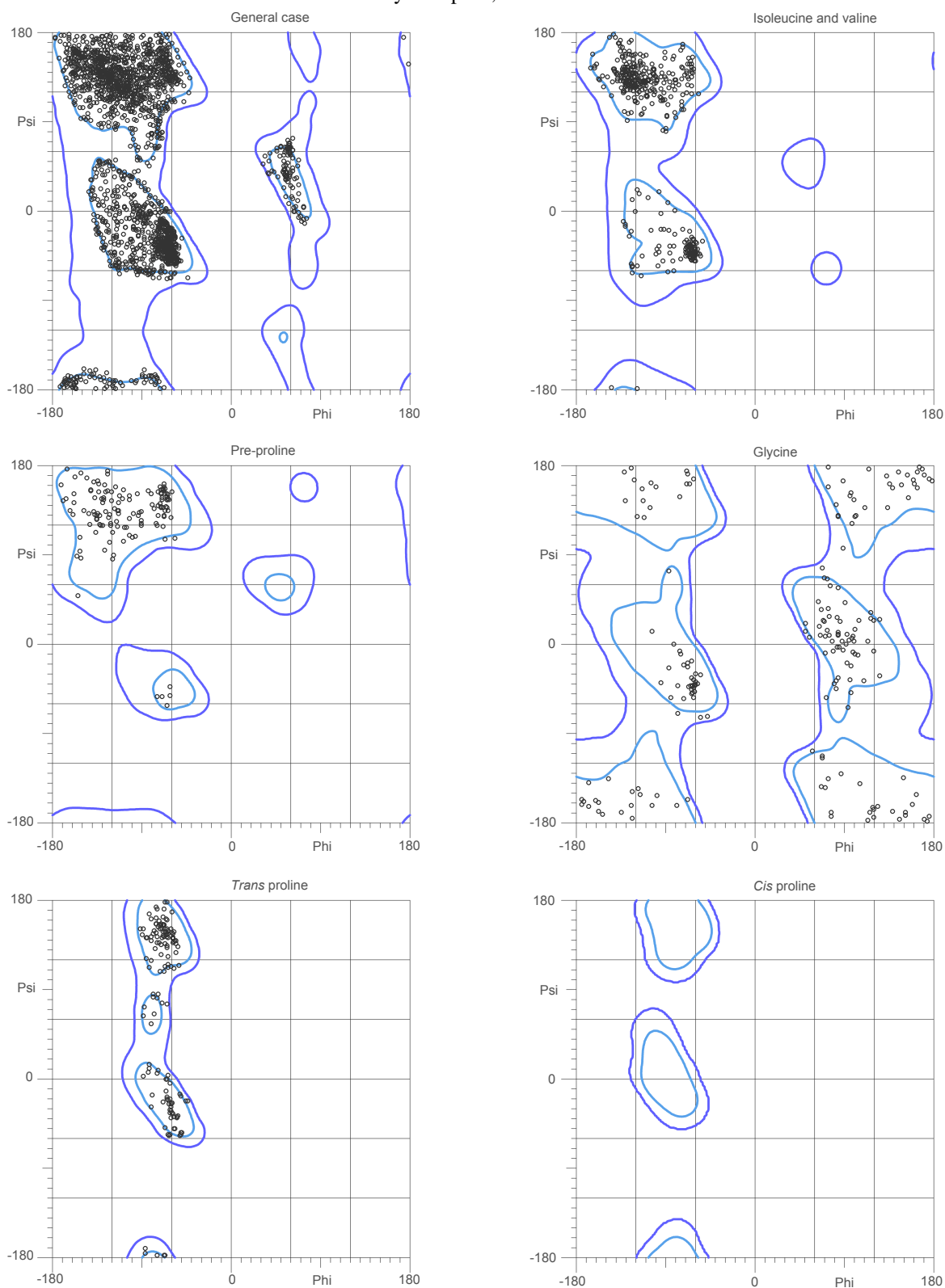


# MolProbity Ramachandran analysis

7byr.H.pdb, model 1



91.6% (2795/3051) of all residues were in favored (98%) regions.

100.0% (3051/3051) of all residues were in allowed (>99.8%) regions.

There were no outliers.