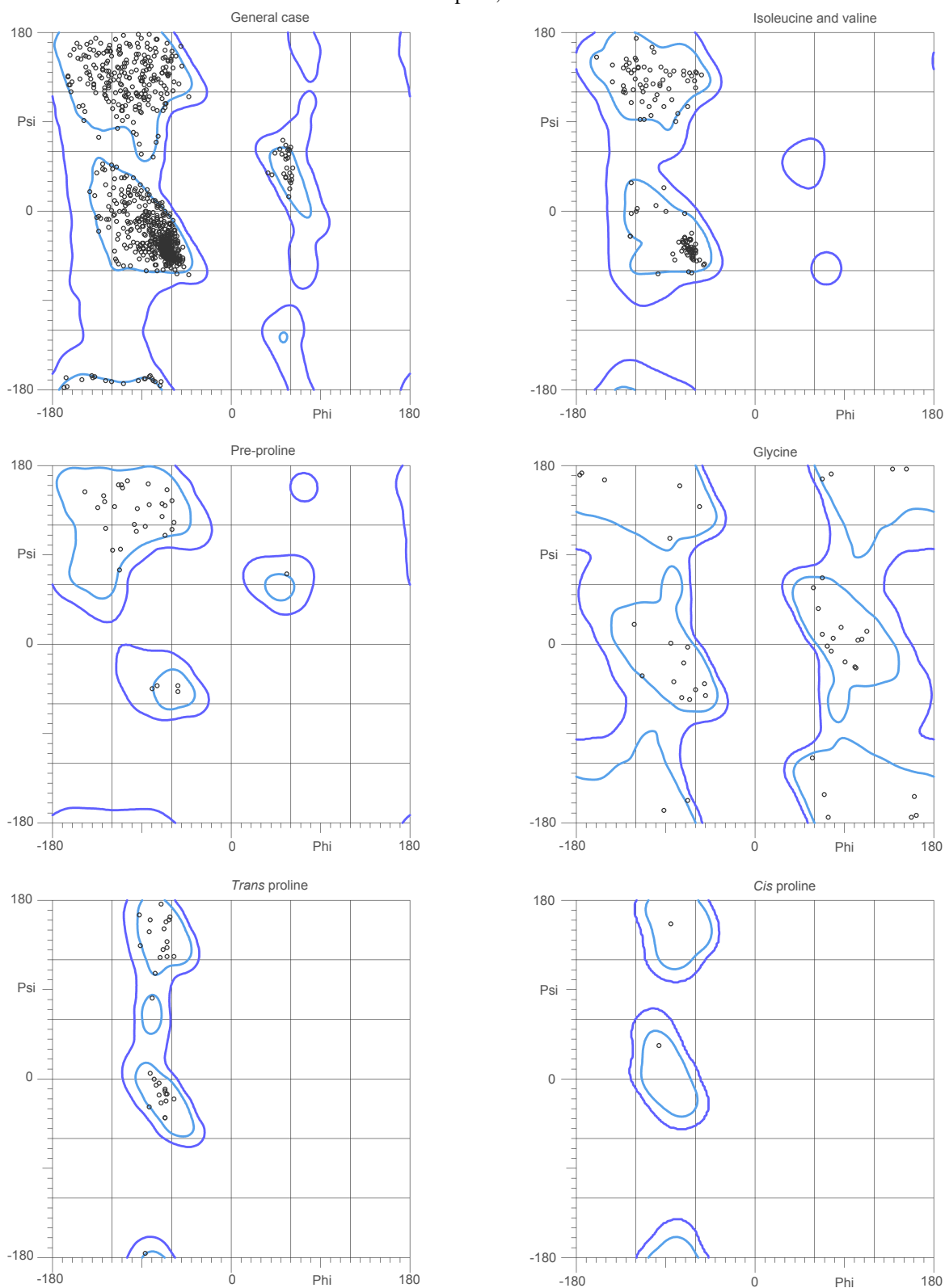


# MolProbity Ramachandran analysis

6m71.H.pdb, model 1



95.8% (1020/1065) of all residues were in favored (98%) regions.

100.0% (1065/1065) of all residues were in allowed (>99.8%) regions.

There were no outliers.