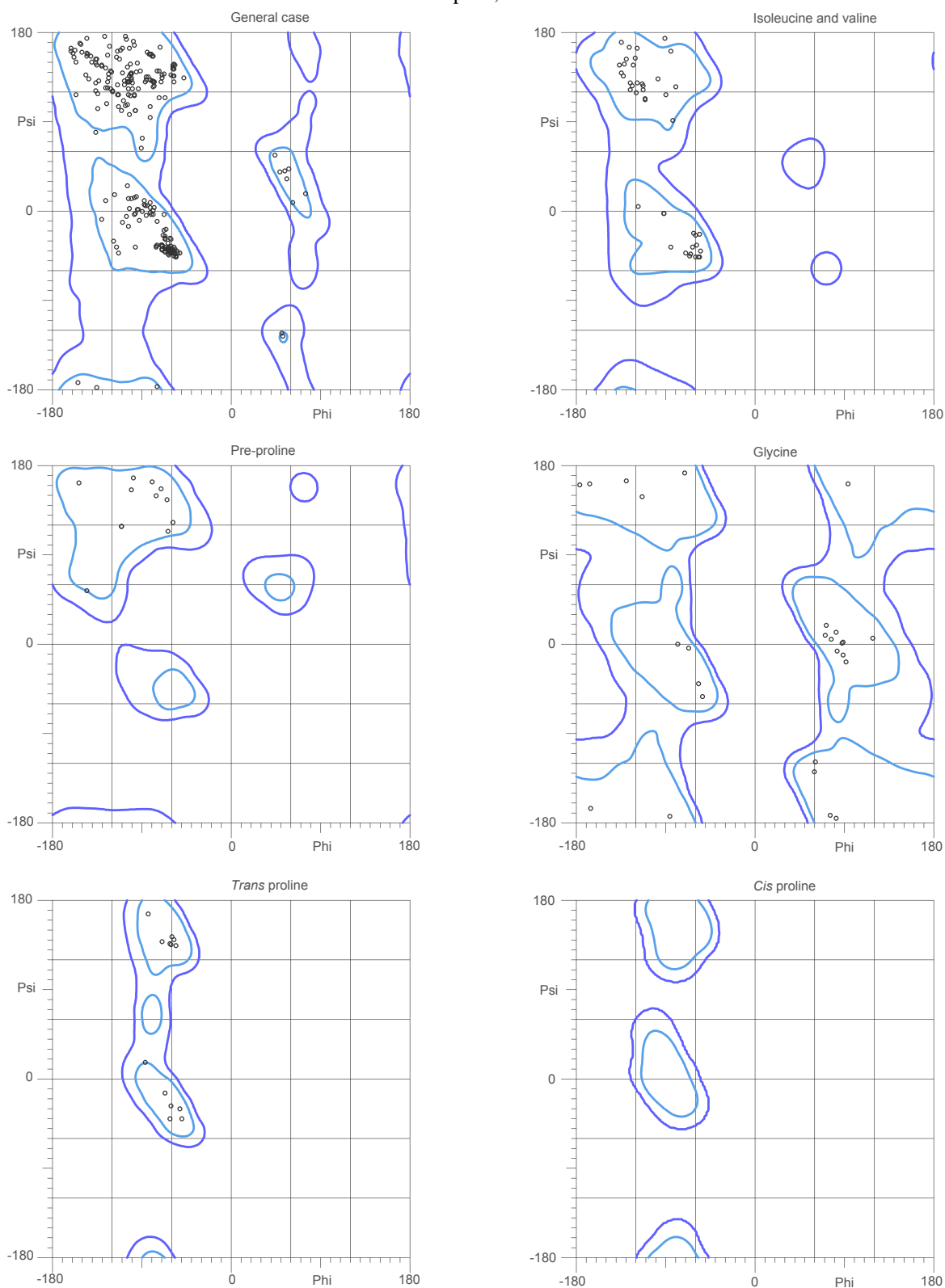


# MolProbity Ramachandran analysis

6w79.H.pdb, model 1



99.1% (333/336) of all residues were in favored (98%) regions.

100.0% (336/336) of all residues were in allowed (>99.8%) regions.

There were no outliers.