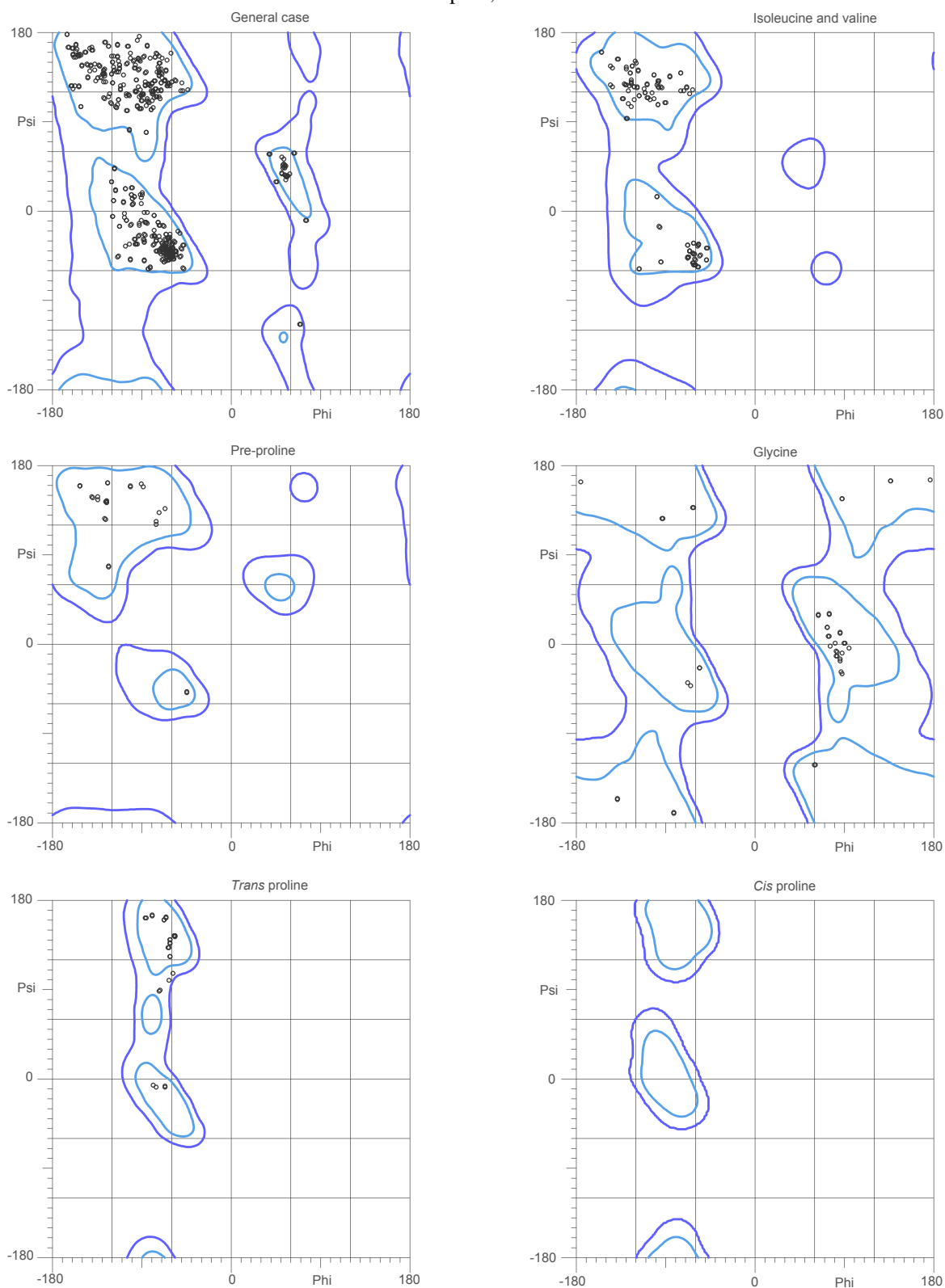


# MolProbity Ramachandran analysis

5s6x.H.pdb, model 1



98.0% (688/702) of all residues were in favored (98%) regions.

100.0% (702/702) of all residues were in allowed (>99.8%) regions.

There were no outliers.