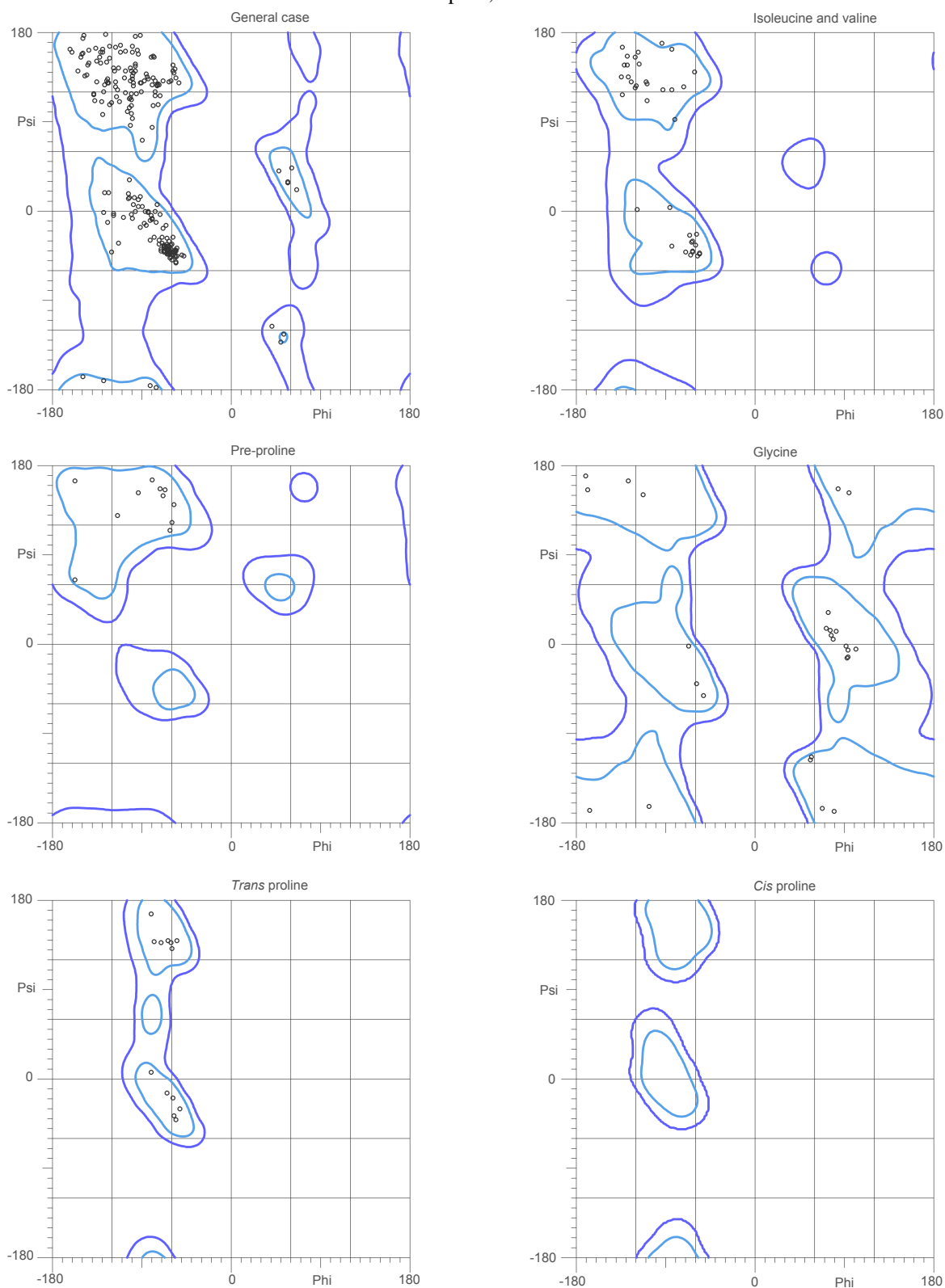


# MolProbity Ramachandran analysis

7alh.H.pdb, model 1



99.0% (306/309) of all residues were in favored (98%) regions.

100.0% (309/309) of all residues were in allowed (>99.8%) regions.

There were no outliers.