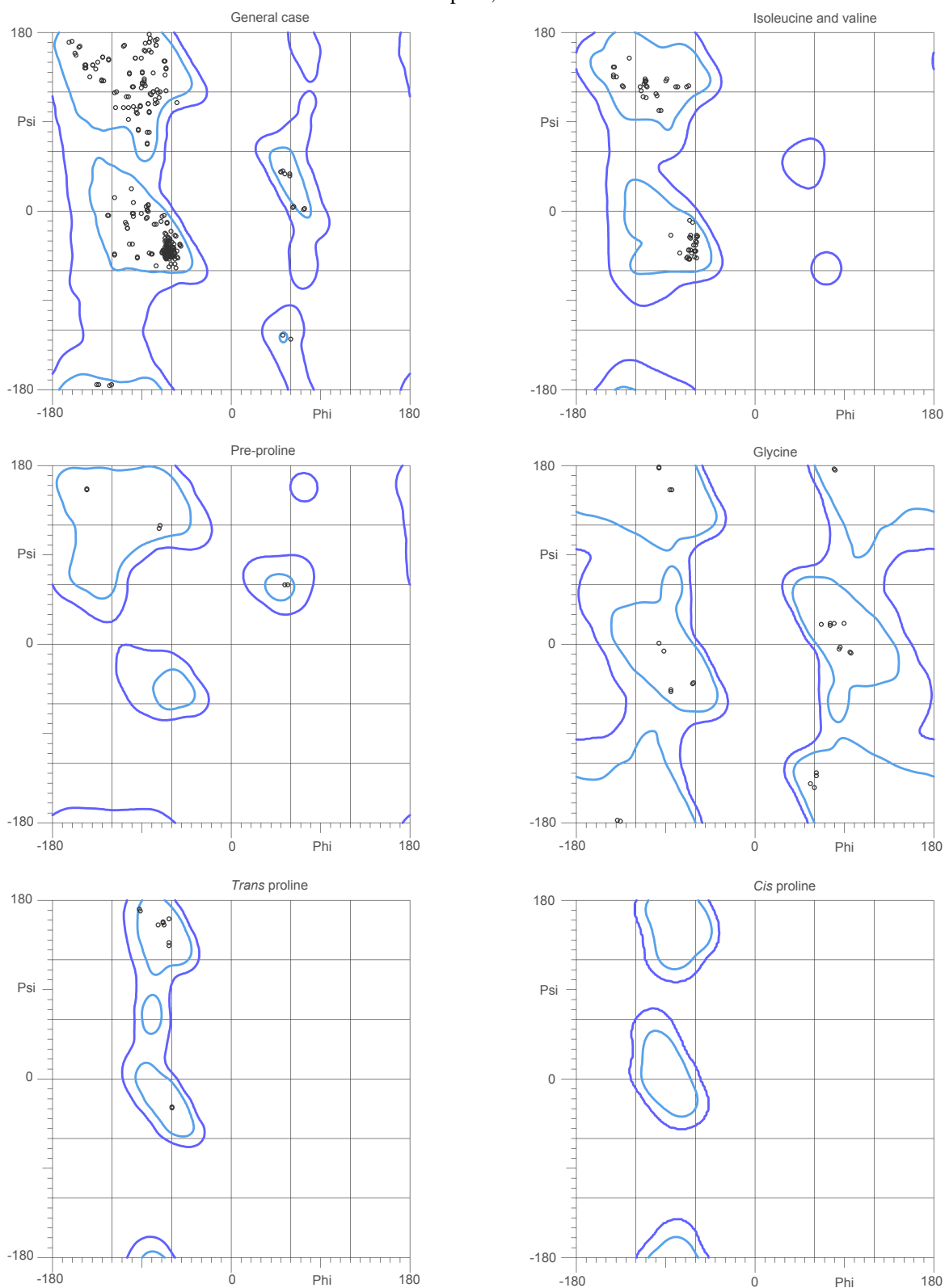


# MolProbity Ramachandran analysis

5s22.H.pdb, model 1



98.9% (344/348) of all residues were in favored (98%) regions.

100.0% (348/348) of all residues were in allowed (>99.8%) regions.

There were no outliers.