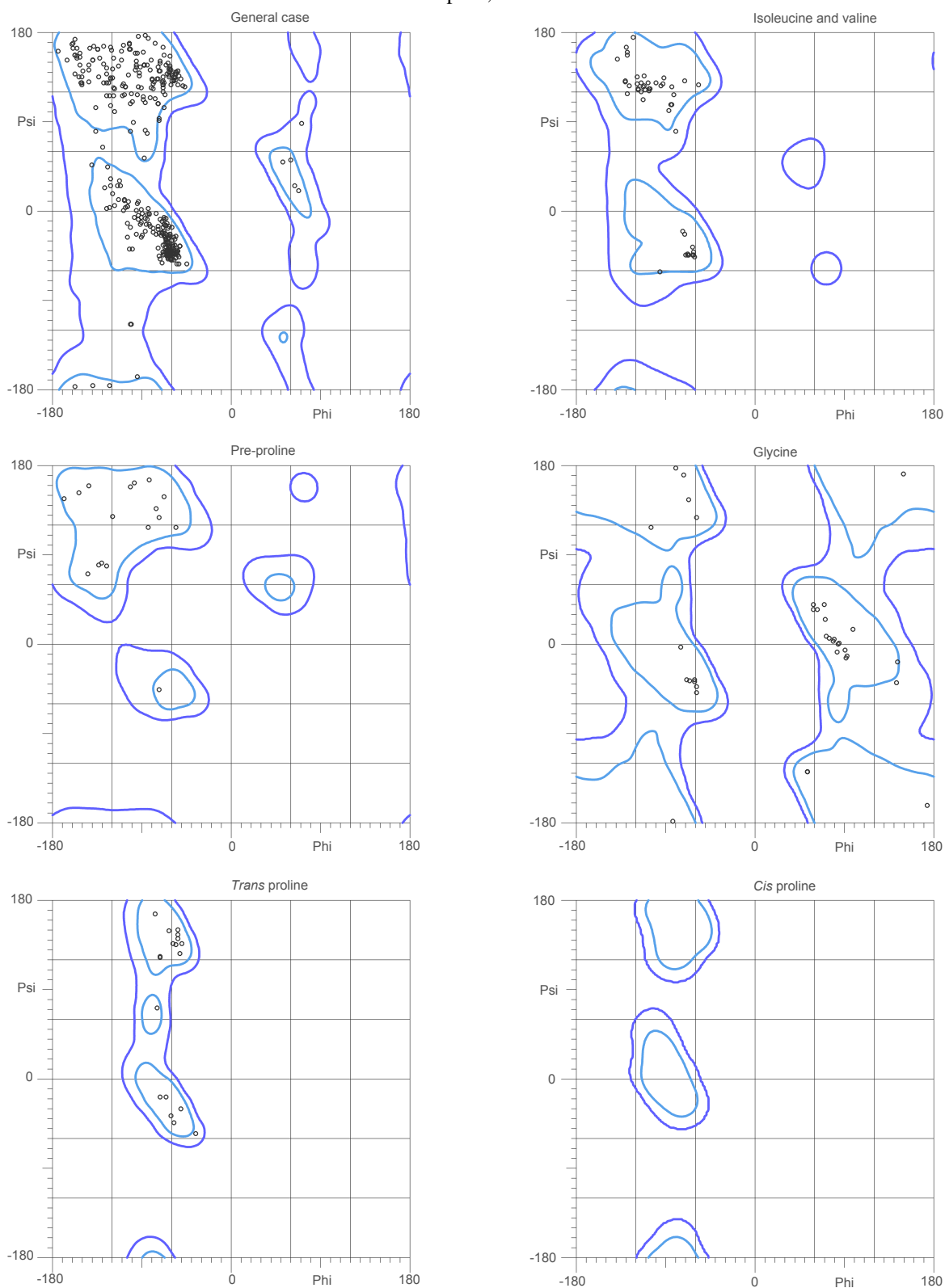


# MolProbity Ramachandran analysis

716t.H.pdb, model 1



97.4% (452/464) of all residues were in favored (98%) regions.

100.0% (464/464) of all residues were in allowed (>99.8%) regions.

There were no outliers.