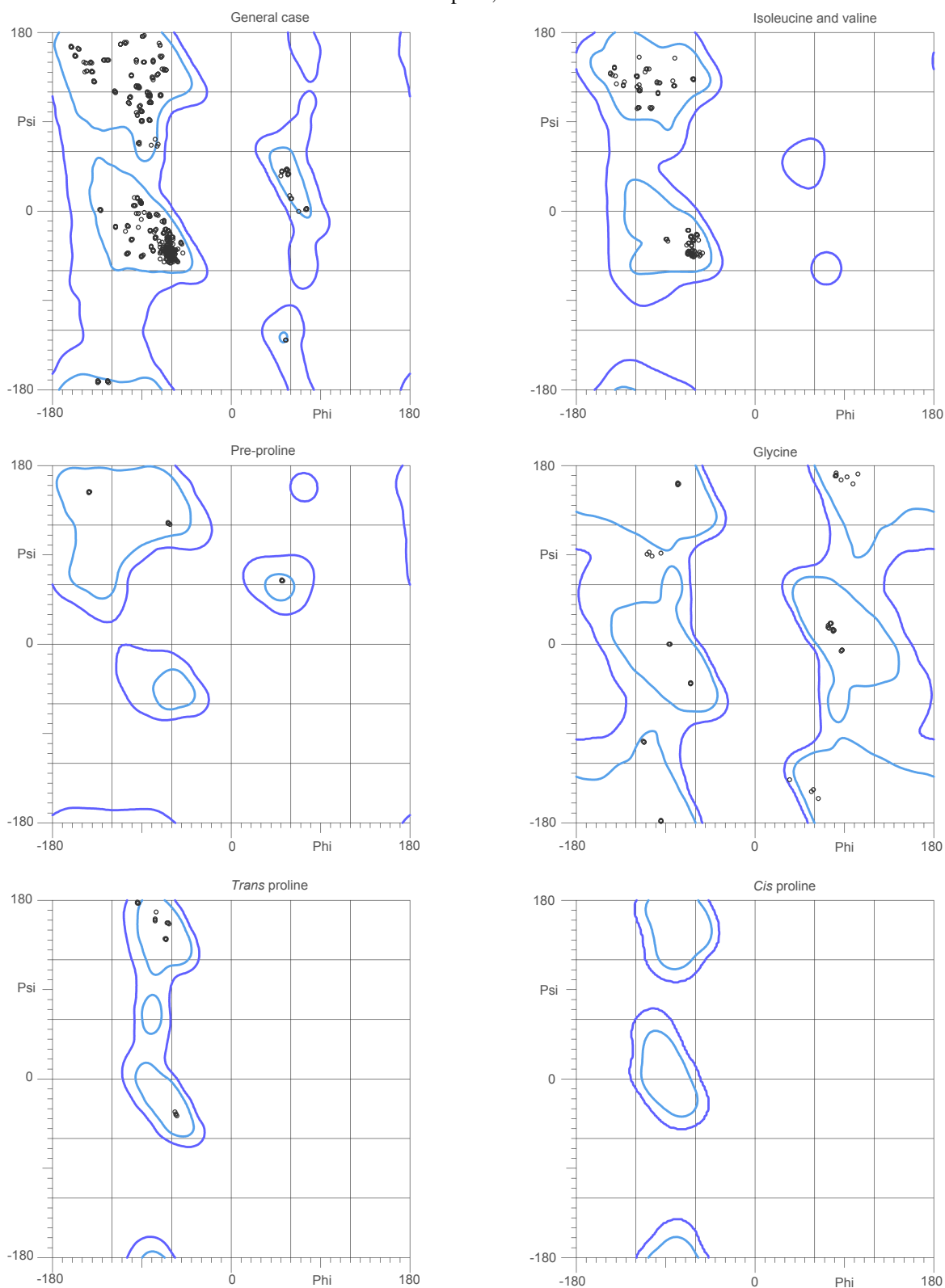


# MolProbity Ramachandran analysis

6z72.H.pdb, model 1



98.1% (674/687) of all residues were in favored (98%) regions.

100.0% (687/687) of all residues were in allowed (>99.8%) regions.

There were no outliers.