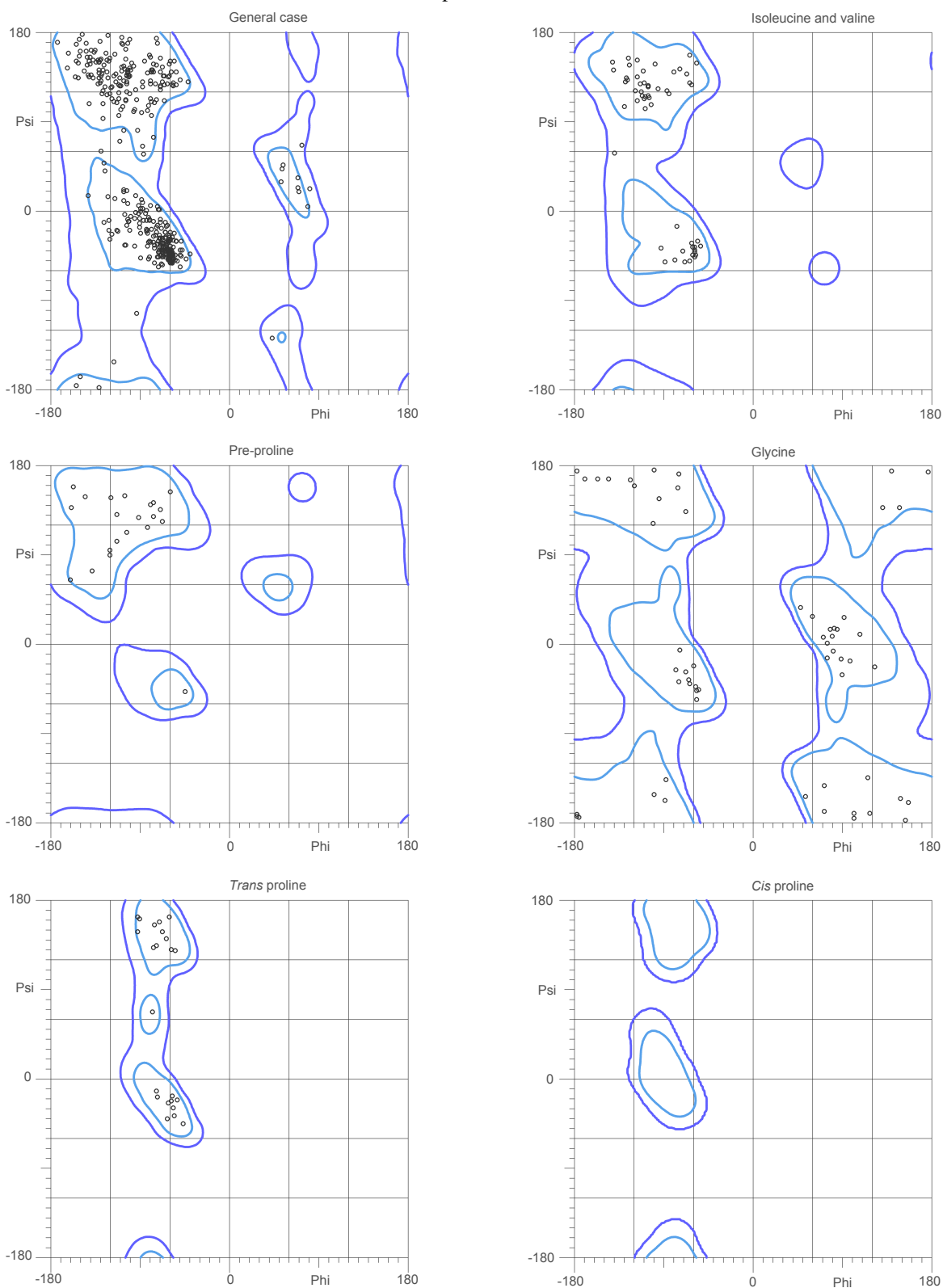


MolProbity Ramachandran analysis

1ctn.pdb, model 1



97.6% (523/536) of all residues were in favored (98%) regions.
100.0% (536/536) of all residues were in allowed (>99.8%) regions.

There were no outliers.