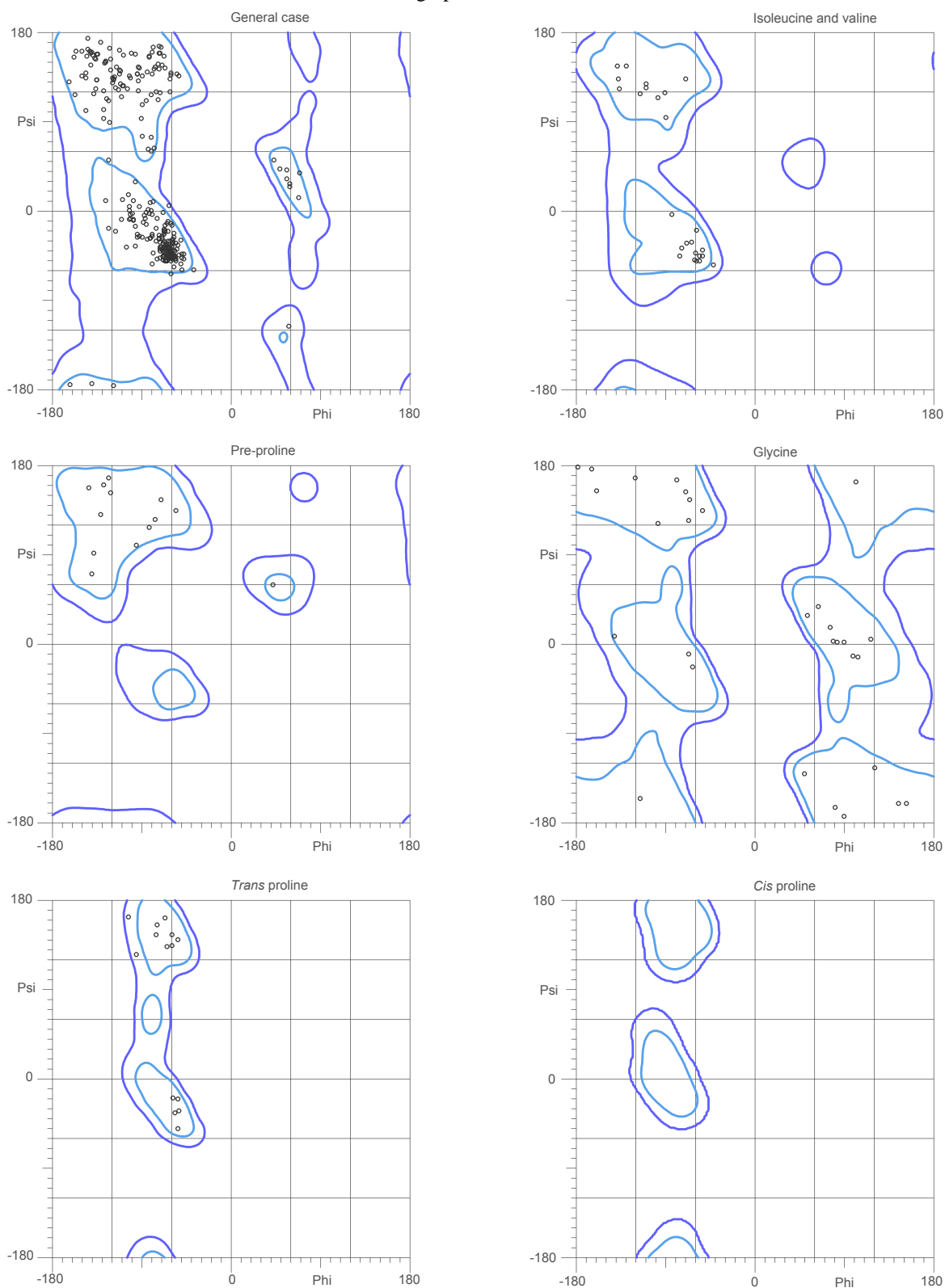


MolProbity Ramachandran analysis

1lg2.pdb, model 1



97.5% (346/355) of all residues were in favored (98%) regions.
100.0% (355/355) of all residues were in allowed (>99.8%) regions.

There were no outliers.