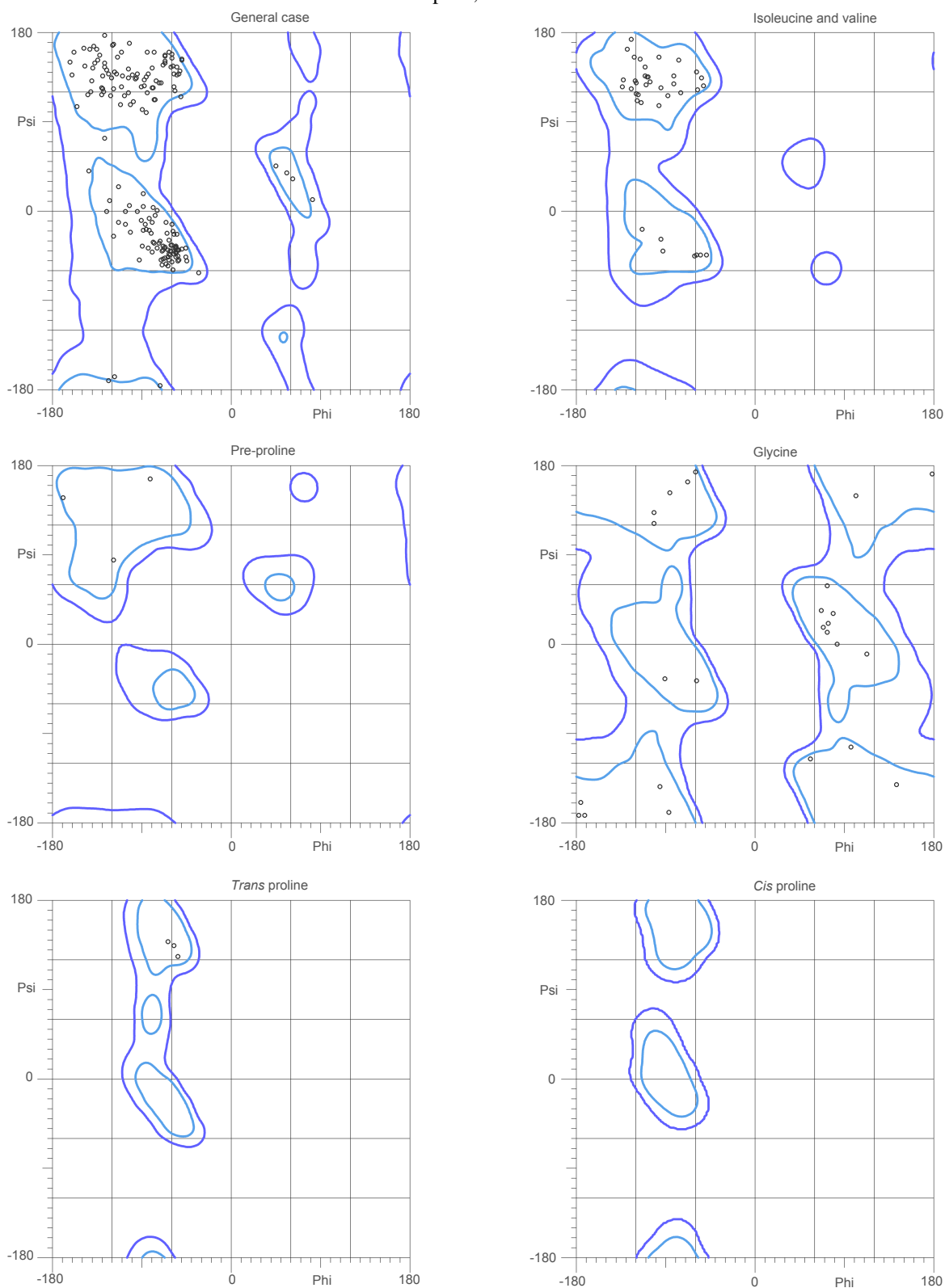


MolProbity Ramachandran analysis

630.pdb, model 1



96.3% (237/246) of all residues were in favored (98%) regions.

100.0% (246/246) of all residues were in allowed (>99.8%) regions.

There were no outliers.