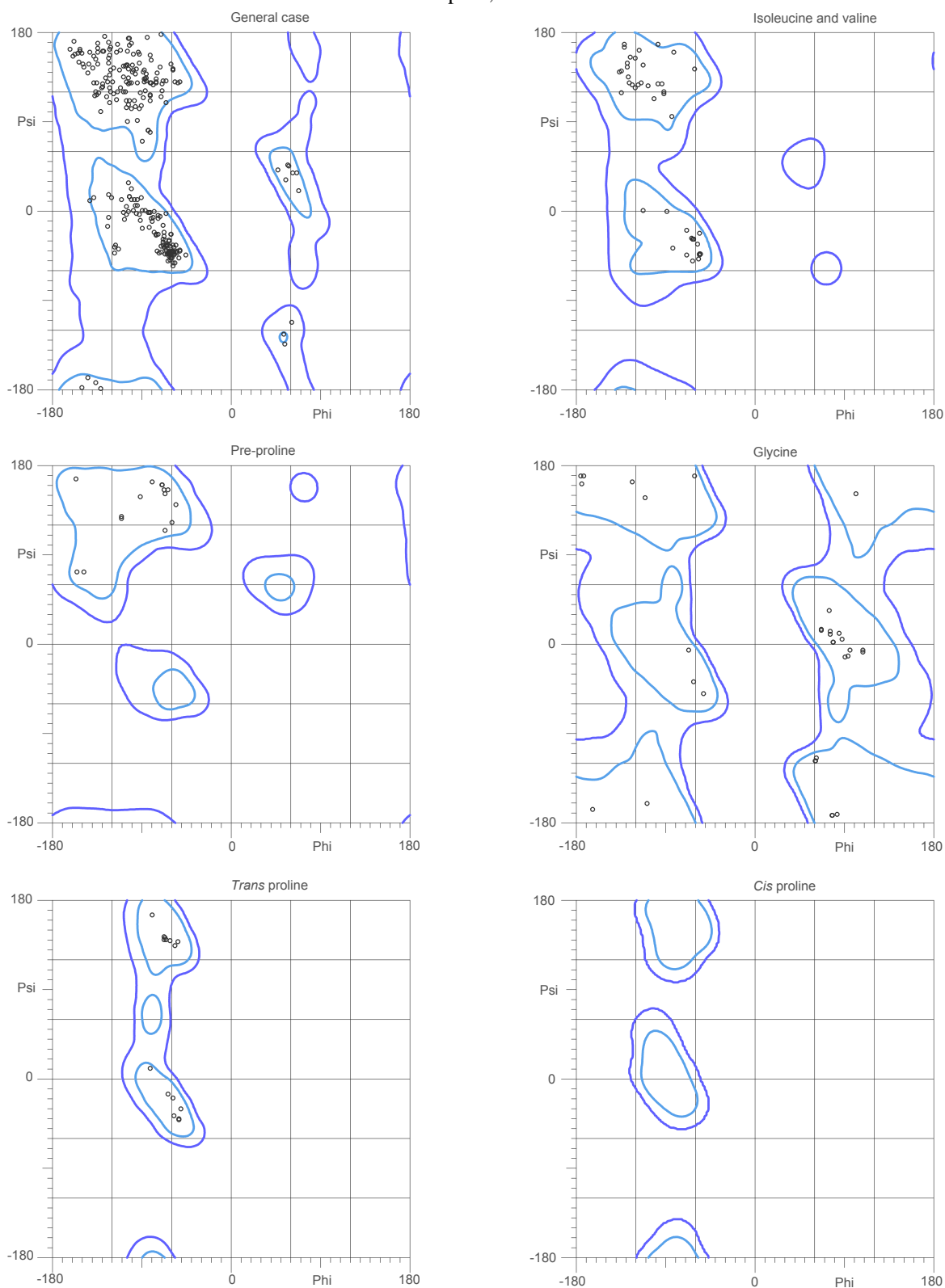


MolProbity Ramachandran analysis

7ar5.H.pdb, model 1



98.9% (368/372) of all residues were in favored (98%) regions.

100.0% (372/372) of all residues were in allowed (>99.8%) regions.

There were no outliers.