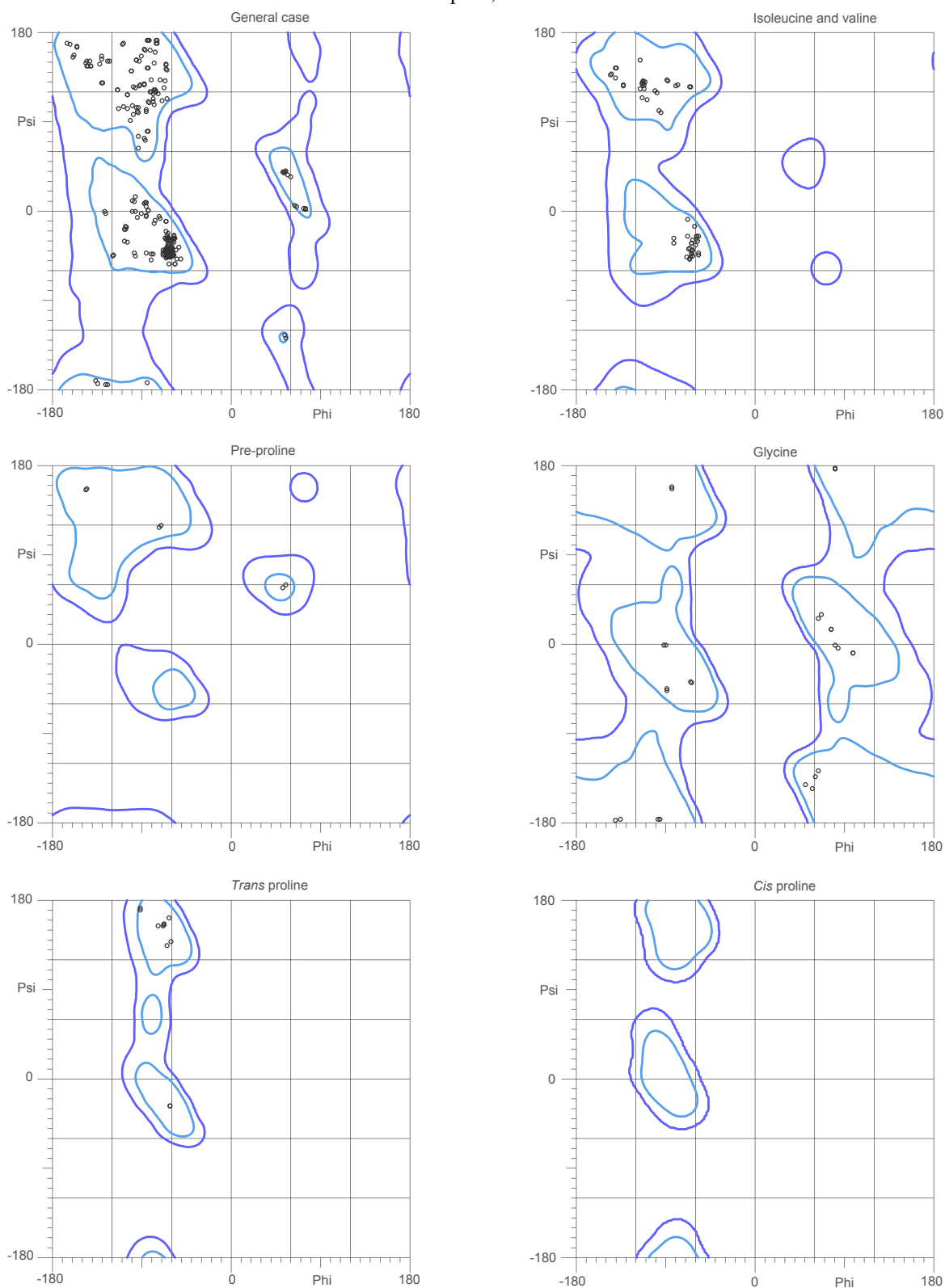# MolProbity Ramachandran analysis

5s42.H.pdb, model 1

100.0% (347/347) of all residues were in favored (98%) regions.

100.0% (347/347) of all residues were in allowed (>99.8%) regions.

There were no outliers.