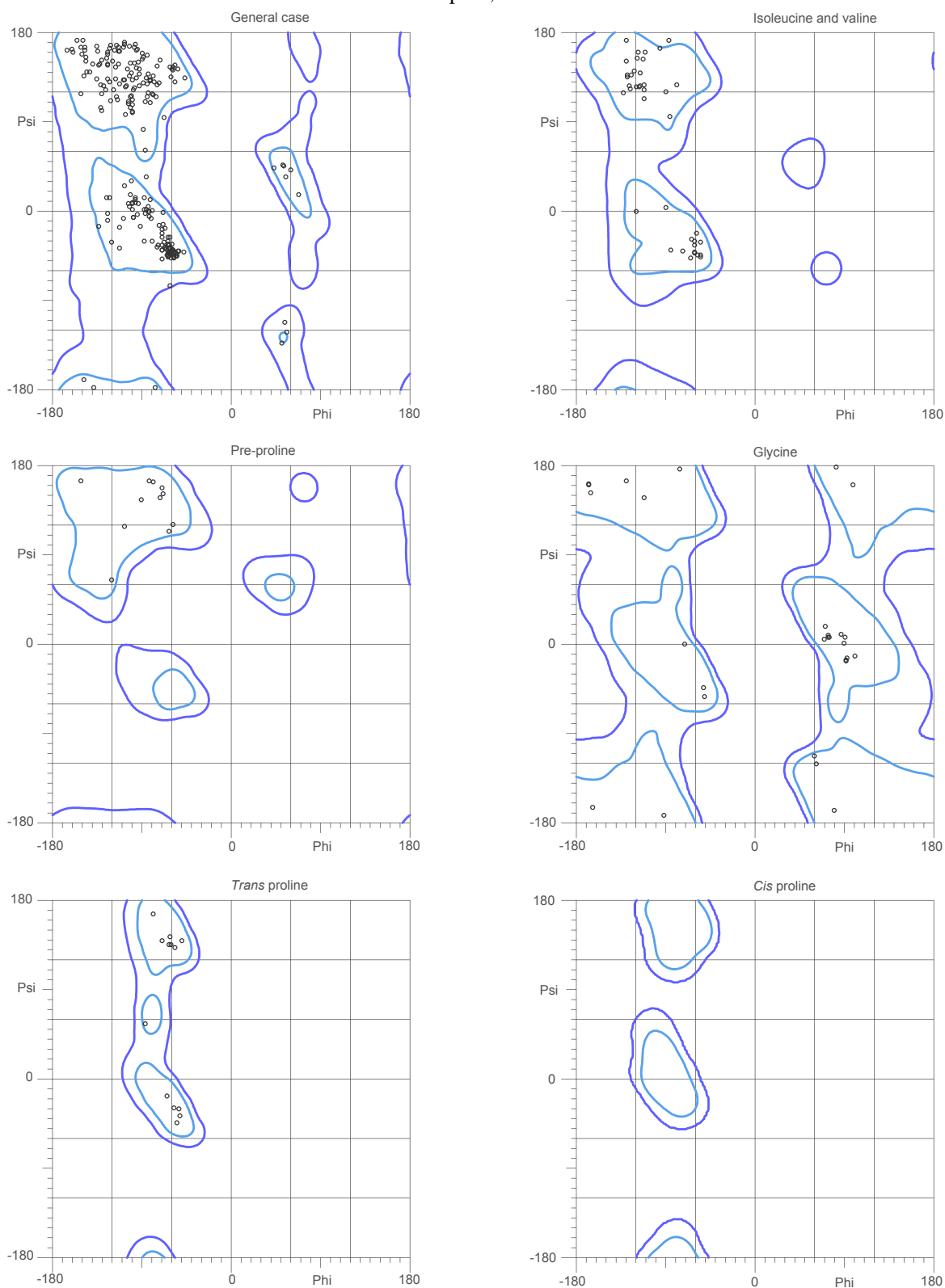


MolProbity Ramachandran analysis

715d.H.pdb, model 1



97.8% (318/325) of all residues were in favored (98%) regions.

100.0% (325/325) of all residues were in allowed (>99.8%) regions.

There were no outliers.