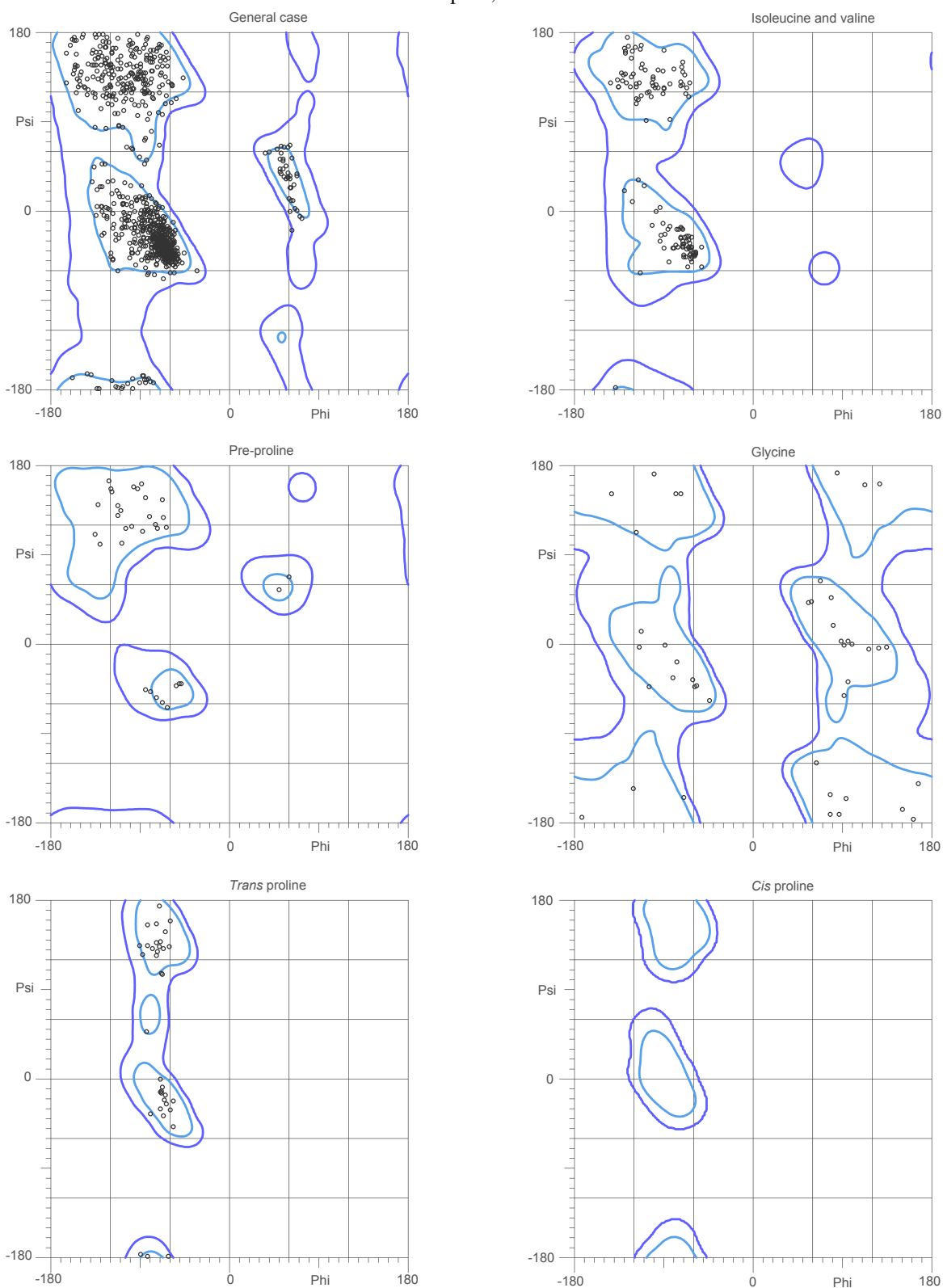


MolProbity Ramachandran analysis

7bv1.H.pdb, model 1



95.4% (1034/1084) of all residues were in favored (98%) regions.

100.0% (1084/1084) of all residues were in allowed (>99.8%) regions.

There were no outliers.