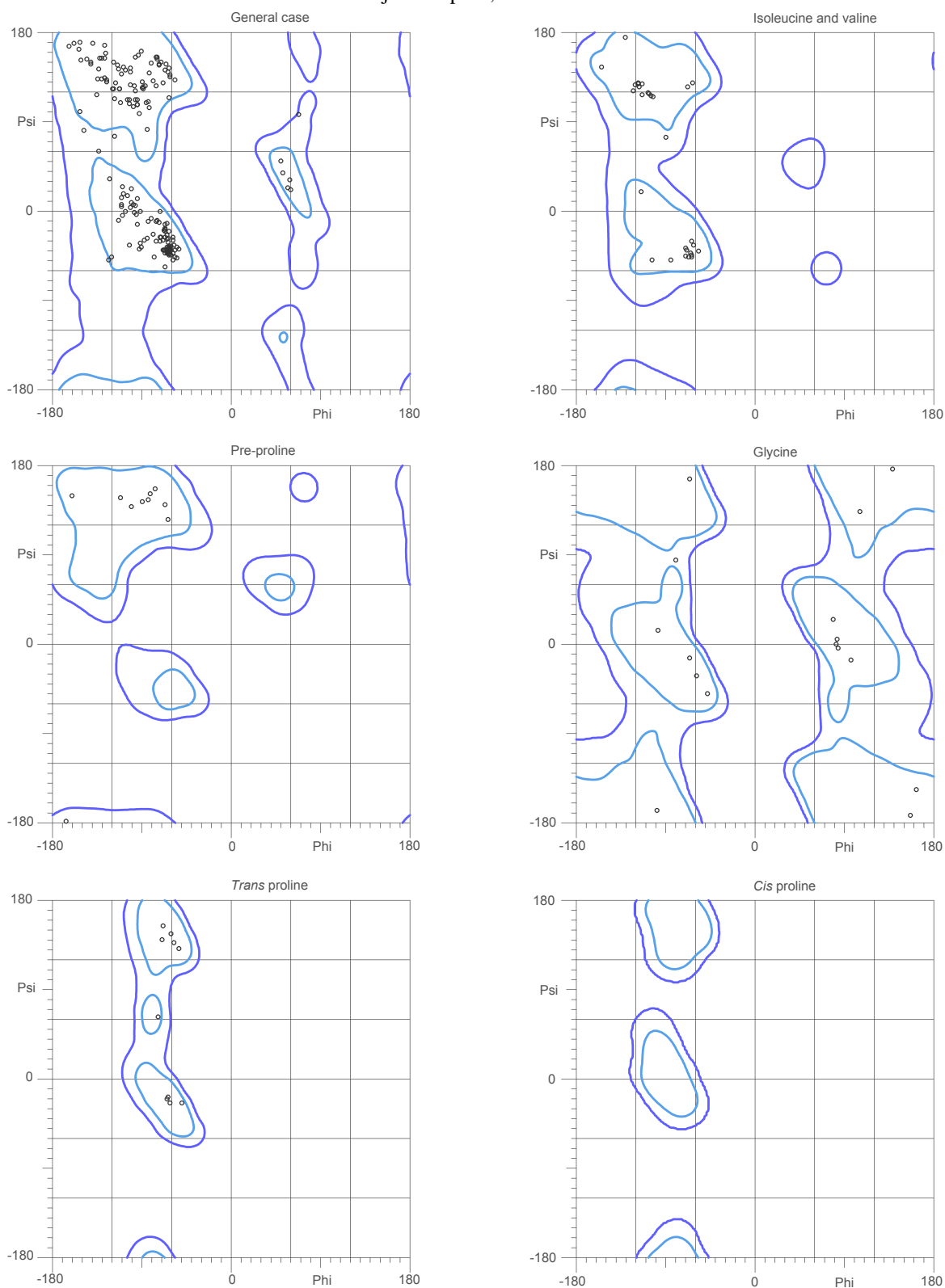


# MolProbity Ramachandran analysis

7jzu.H.pdb, model 1



96.7% (236/244) of all residues were in favored (98%) regions.

100.0% (244/244) of all residues were in allowed (>99.8%) regions.

There were no outliers.