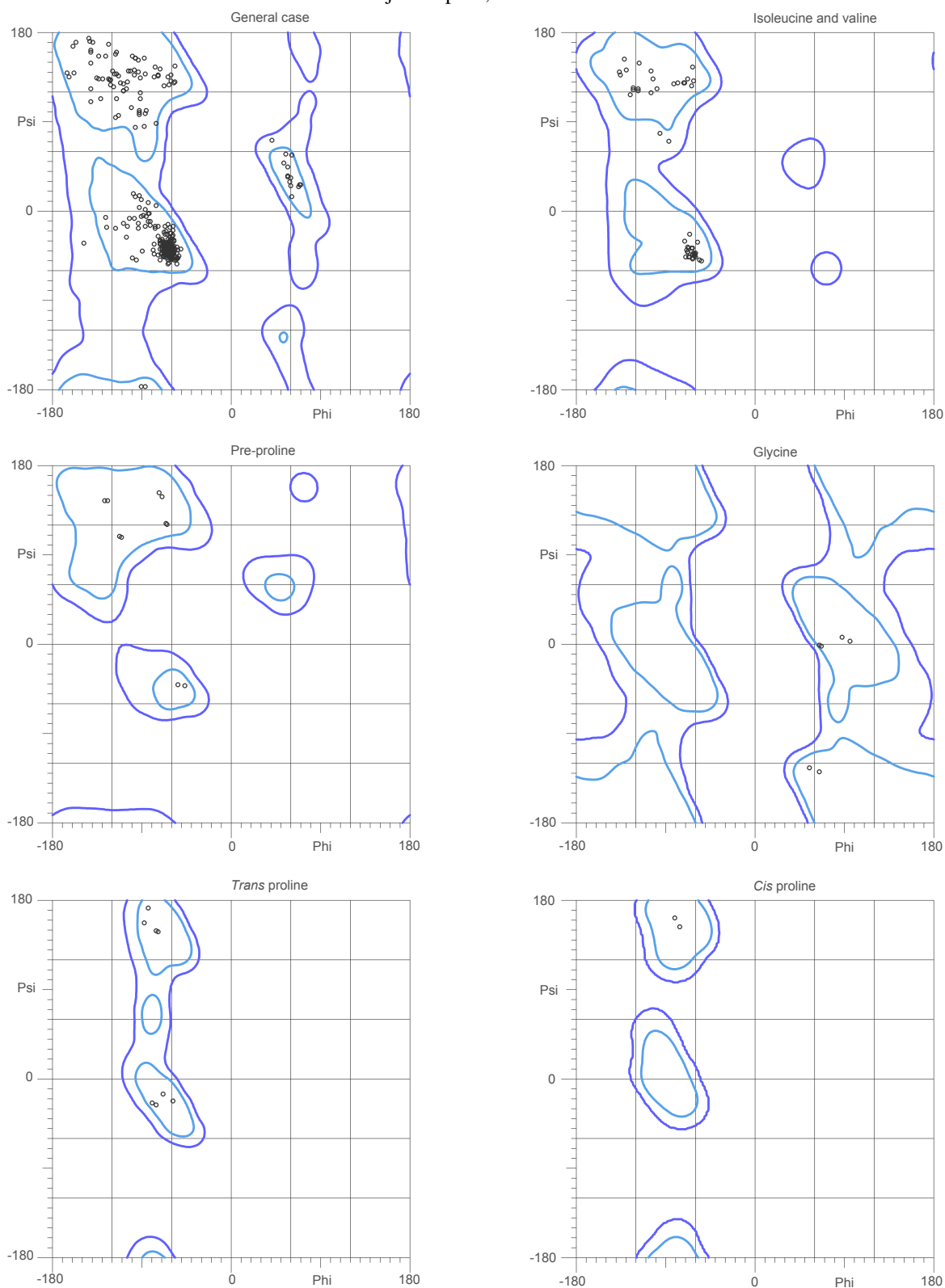


# MolProbity Ramachandran analysis

7jlt.H.pdb, model 1



99.0% (381/385) of all residues were in favored (98%) regions.

100.0% (385/385) of all residues were in allowed (>99.8%) regions.

There were no outliers.