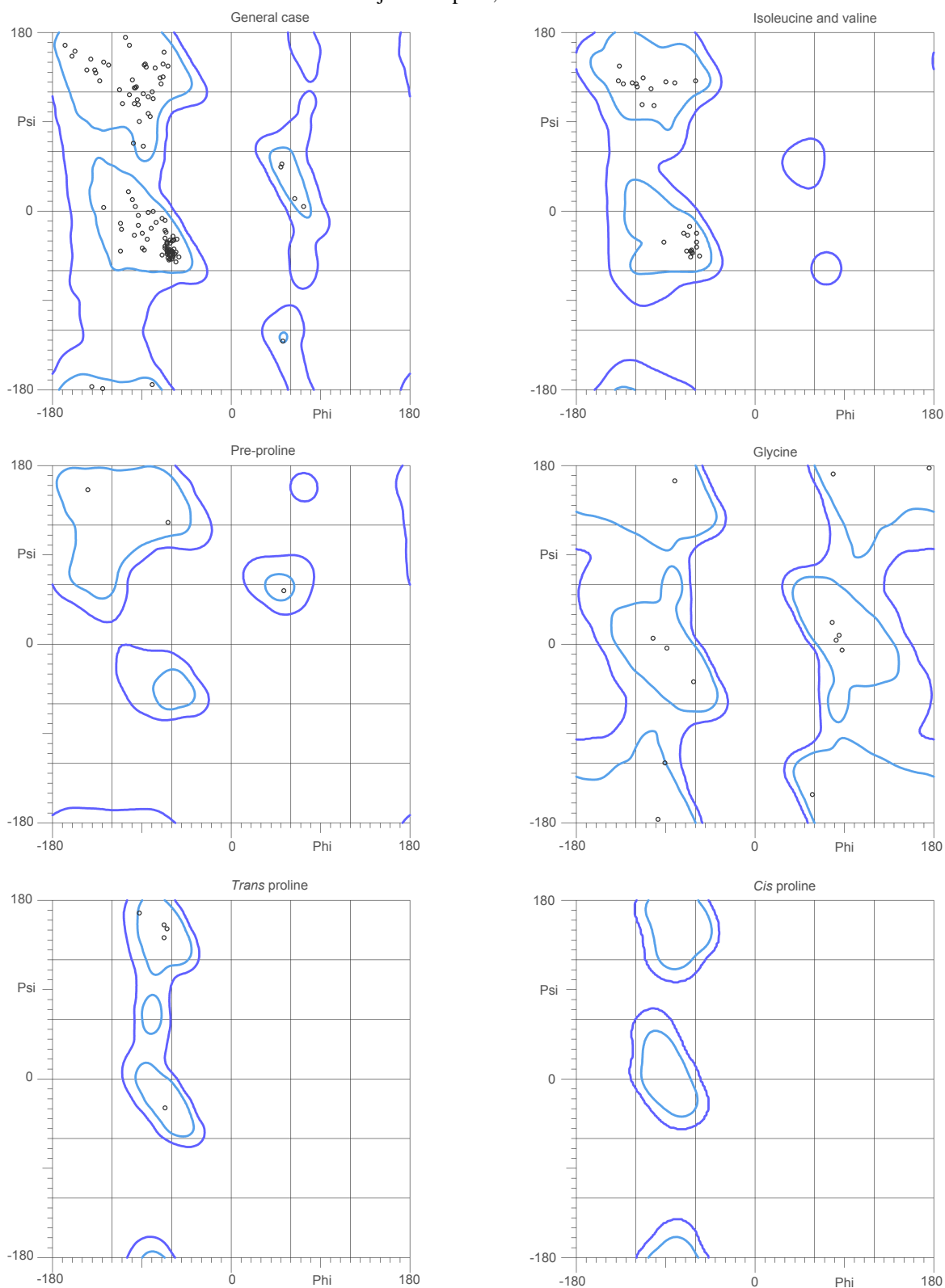


# MolProbity Ramachandran analysis

7jme.H.pdb, model 1



99.4% (165/166) of all residues were in favored (98%) regions.

100.0% (166/166) of all residues were in allowed (>99.8%) regions.

There were no outliers.