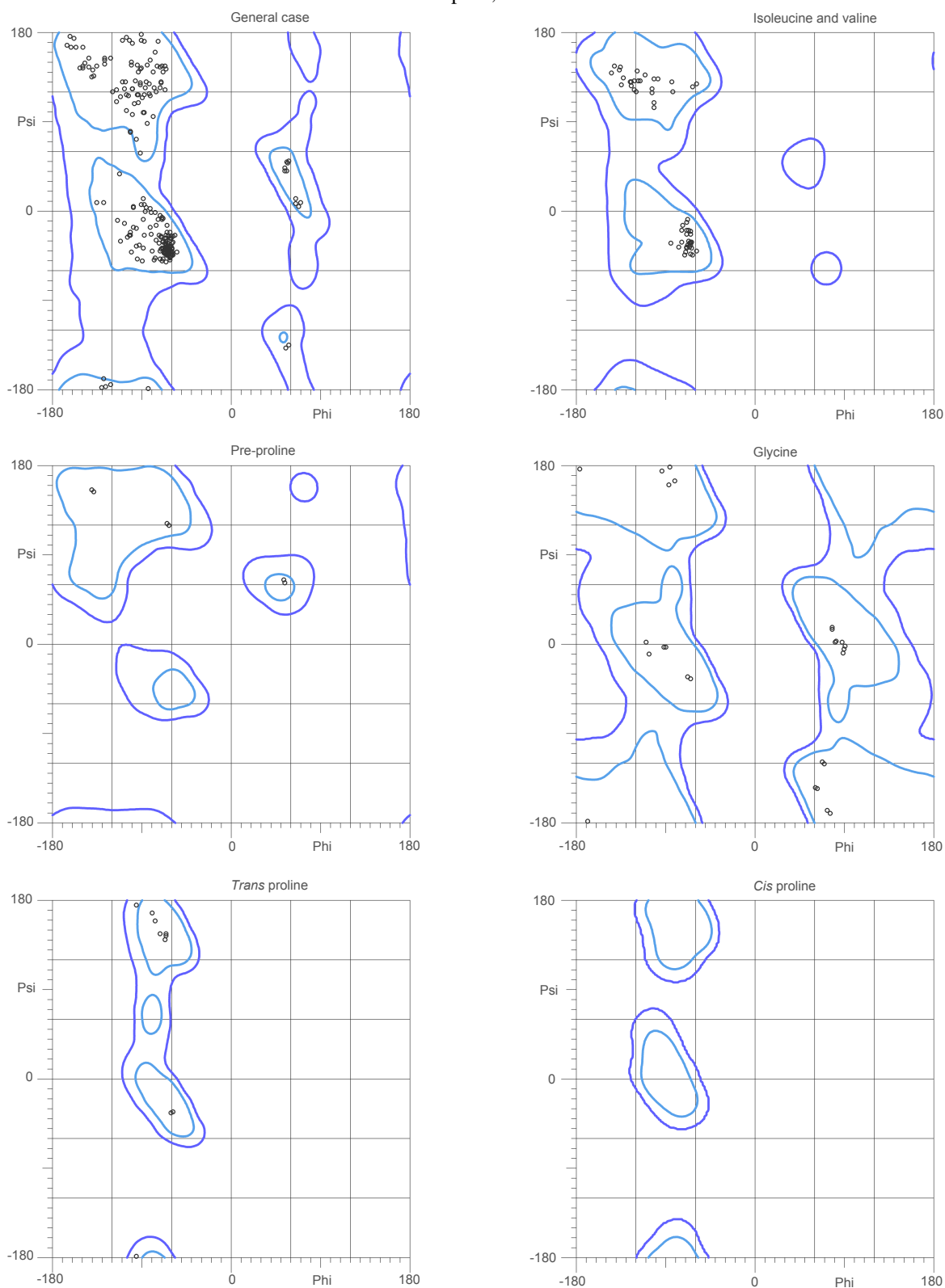


# MolProbity Ramachandran analysis

6vxs.H.pdb, model 1



98.5% (325/330) of all residues were in favored (98%) regions.

100.0% (330/330) of all residues were in allowed (>99.8%) regions.

There were no outliers.