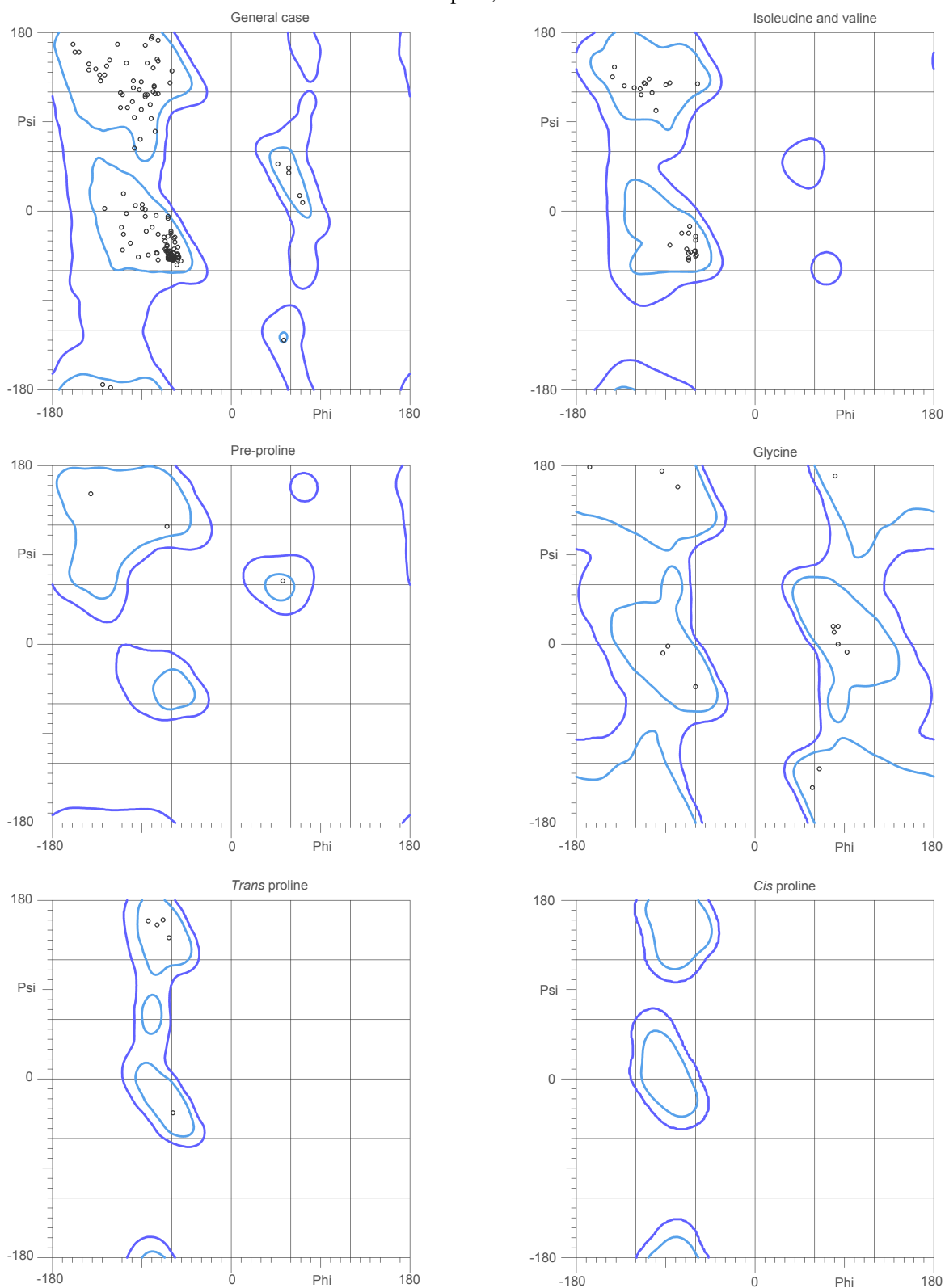


# MolProbity Ramachandran analysis

5rvk.H.pdb, model 1



99.5% (183/184) of all residues were in favored (98%) regions.

100.0% (184/184) of all residues were in allowed (>99.8%) regions.

There were no outliers.