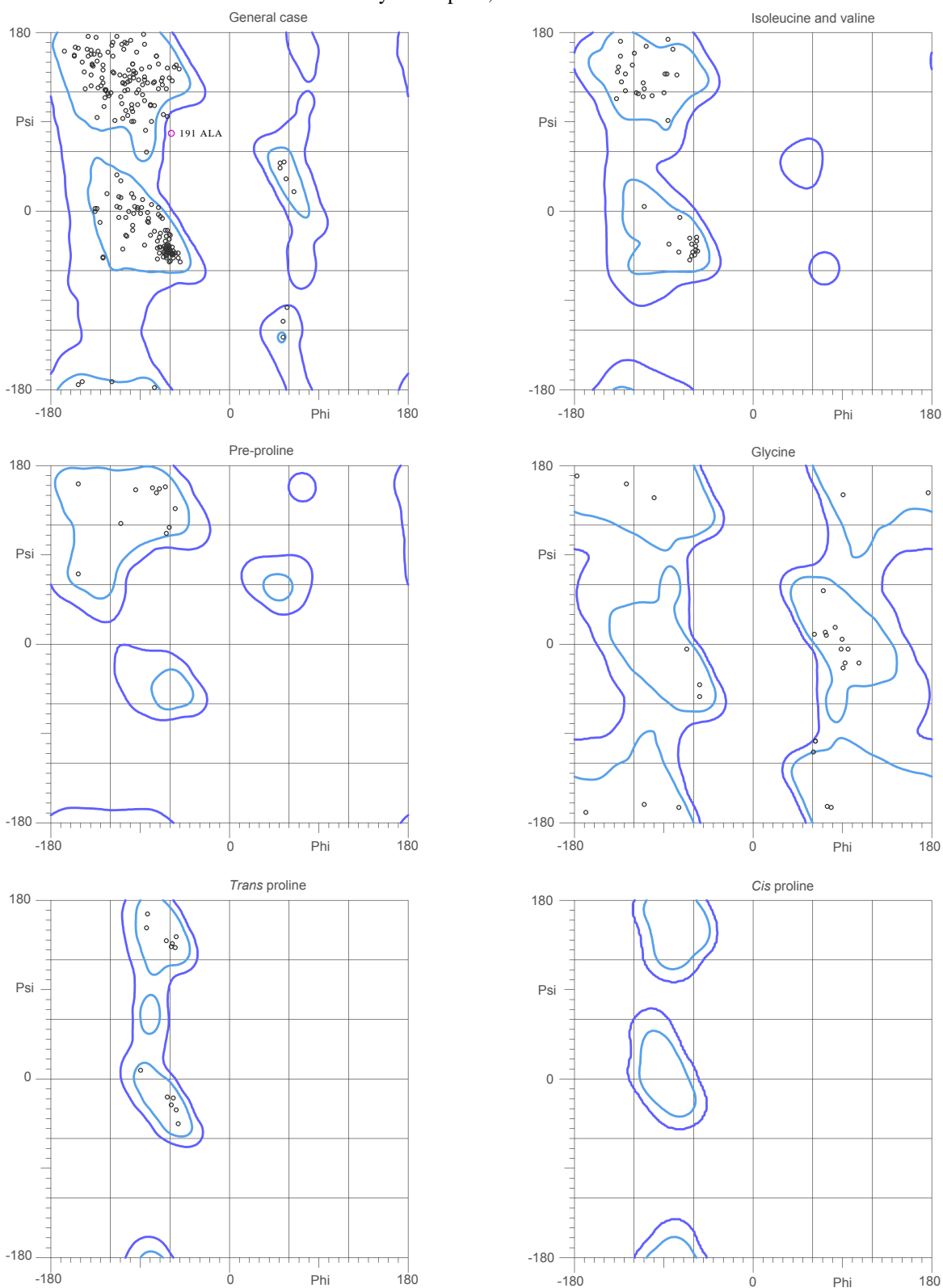


# MolProbity Ramachandran analysis

6yt8.H.pdb, model 1



96.8% (306/316) of all residues were in favored (98%) regions.

99.7% (315/316) of all residues were in allowed (>99.8%) regions.

There were 1 outliers (phi, psi):

191 ALA (-59.1, 79.6)