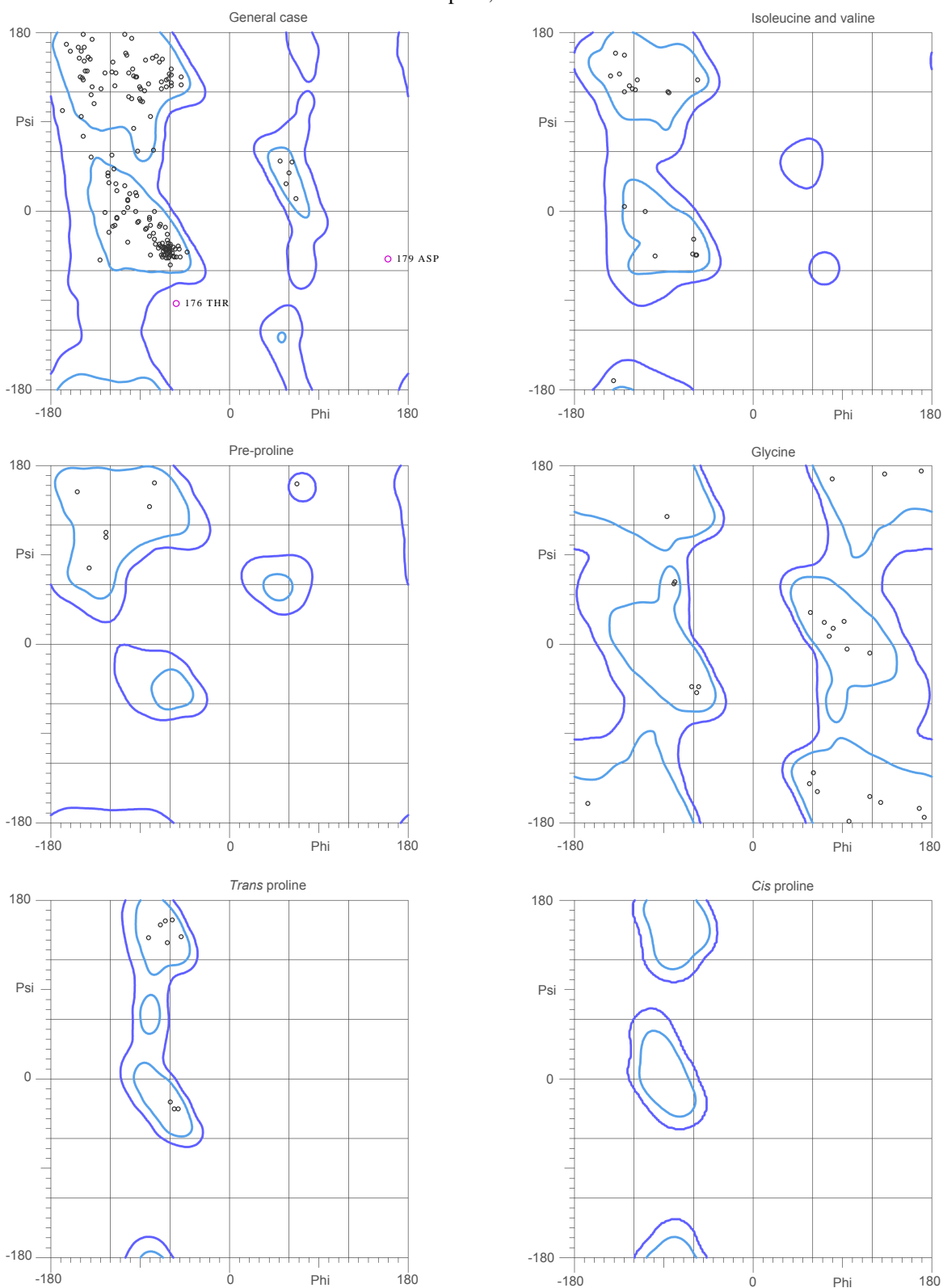


# MolProbity Ramachandran analysis

3of9.H.pdb, model 1



95.9% (210/219) of all residues were in favored (98%) regions.

99.1% (217/219) of all residues were in allowed (>99.8%) regions.

There were 2 outliers (phi, psi):

176 THR (-54.2, -93.1)

179 ASP (160.8, -48.2)