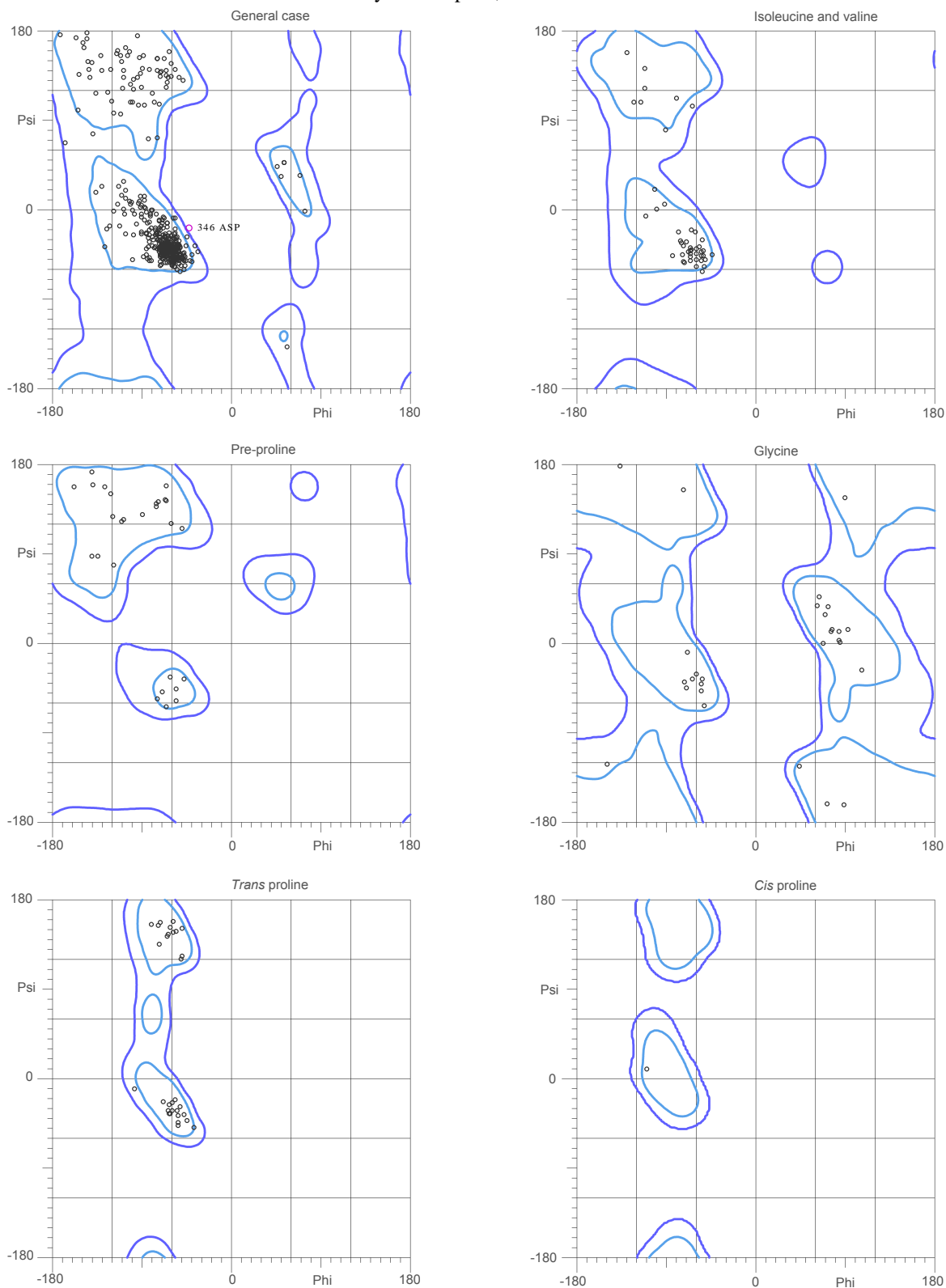


# MolProbity Ramachandran analysis

2ydm.H.pdb, model 1



97.0% (554/571) of all residues were in favored (98%) regions.

99.8% (570/571) of all residues were in allowed (>99.8%) regions.

There were 1 outliers (phi, psi):

346 ASP (-43.8, -18.4)