

# MolProbity Ramachandran analysis

1wyy.H.pdb, model 1



97.1% (235/242) of all residues were in favored (98%) regions.

99.6% (241/242) of all residues were in allowed (>99.8%) regions.

There were 1 outliers (phi, psi):

A\_892 GLY (139.0, 63.3)