

# MolProbity Ramachandran analysis

5red.H.pdb, model 1



97.8% (350/358) of all residues were in favored (98%) regions.

99.7% (357/358) of all residues were in allowed (>99.8%) regions.

There were 1 outliers (phi, psi):

154 TYR (55.9, -85.3)