

# MolProbity Ramachandran analysis

5rhe.H.pdb, model 1



98.7% (303/307) of all residues were in favored (98%) regions.

99.7% (306/307) of all residues were in allowed (>99.8%) regions.

There were 1 outliers (phi, psi):

154 TYR (61.7, -85.1)