

# MolProbity Ramachandran analysis

5rha.H.pdb, model 1



99.0% (301/304) of all residues were in favored (98%) regions.

99.7% (303/304) of all residues were in allowed (>99.8%) regions.

There were 1 outliers (phi, psi):

154 TYR (60.3, -81.4)