

# MolProbity Ramachandran analysis

5ree.H.pdb, model 1



97.1% (302/311) of all residues were in favored (98%) regions.

99.7% (310/311) of all residues were in allowed (>99.8%) regions.

There were 1 outliers ( $\Phi$ ,  $\Psi$ ):

154 TYR (65.6, -78.4)