

# MolProbity Ramachandran analysis

5rfw.H.pdb, model 1



98.4% (306/311) 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 (54.4, -84.3)