

# MolProbity Ramachandran analysis

3ebn.H.pdb, model 1



96.4% (380/394) of all residues were in favored (98%) regions.

99.5% (392/394) of all residues were in allowed (>99.8%) regions.

There were 2 outliers ( $\Phi$ ,  $\Psi$ ):

B 222 ARG (-76.5, -153.9)

D 222 ARG (-80.5, -141.9)