

# MolProbity Ramachandran analysis

3atw.H.pdb, model 1



97.5% (585/600) of all residues were in favored (98%) regions.

99.8% (599/600) of all residues were in allowed (>99.8%) regions.

There were 1 outliers (phi, psi):

A 154 TYR (58.6, -89.4)