

# MolProbity Ramachandran analysis

3snb.H.pdb, model 1



95.2% (296/311) of all residues were in favored (98%) regions.

99.4% (309/311) of all residues were in allowed (>99.8%) regions.

There were 2 outliers (phi, psi):

A 5 LYS (-60.4, 96.3)

A 154 TYR (77.2, -77.2)