

# MolProbity Ramachandran analysis

3sne.H.pdb, model 1



95.8% (296/309) of all residues were in favored (98%) regions.

99.7% (308/309) of all residues were in allowed (>99.8%) regions.

There were 1 outliers (phi, psi):

A 154 TYR (83.2, -96.0)