

# MolProbity Ramachandran analysis

5rf4.H.pdb, model 1



98.4% (309/314) of all residues were in favored (98%) regions.

99.7% (313/314) of all residues were in allowed (>99.8%) regions.

There were 1 outliers (phi, psi):

154 TYR (58.9, -82.2)