

# MolProbity Ramachandran analysis

5rh9.H.pdb, model 1



96.4% (293/304) of all residues were in favored (98%) regions.

99.3% (302/304) of all residues were in allowed (>99.8%) regions.

There were 2 outliers (phi, psi):

48 ASP (-169.6, 32.6)

154 TYR (61.2, -87.0)