

MolProbity Ramachandran analysis

5rfg.H.pdb, model 1



95.0% (302/318) of all residues were in favored (98%) regions.

99.7% (317/318) of all residues were in allowed (>99.8%) regions.

There were 1 outliers (phi, psi):

154 TYR (58.8, -82.2)