

MolProbity Ramachandran analysis

5rh2.H.pdb, model 1



98.4% (299/304) of all residues were in favored (98%) regions.

99.7% (303/304) of all residues were in allowed (>99.8%) regions.

There were 1 outliers (ϕ , ψ):

154 TYR (57.3, -90.1)