

MolProbity Ramachandran analysis

5rgi.H.pdb, model 1



97.5% (306/314) 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 (ϕ , ψ):

154 TYR (60.8, -83.2)