

# MolProbity Ramachandran analysis

5rgq.H.pdb, model 1



96.0% (309/322) of all residues were in favored (98%) regions.

99.4% (320/322) of all residues were in allowed (>99.8%) regions.

There were 2 outliers (phi, psi):

154 TYR (57.2, -81.4)

191 ALA (-55.4, 94.8)