

# MolProbity Ramachandran analysis

2ghv.H.pdb, model 1



94.8% (343/362) of all residues were in favored (98%) regions.

99.4% (360/362) of all residues were in allowed (>99.8%) regions.

There were 2 outliers ( $\phi$ ,  $\psi$ ):

E 321 ASN (63.1, 103.5)

C 321 ASN (66.5, 115.4)