

# MolProbity Ramachandran analysis

2xyv.H.pdb, model 1



97.2% (388/399) of all residues were in favored (98%) regions.

99.5% (397/399) of all residues were in allowed (>99.8%) regions.

There were 2 outliers (phi, psi):

A 33 ASN (-9.6, 101.7)

A 34 ALA (109.0, -61.0)