MolProbity Ramachandran analysis

1avqA.B99990019.pdb, model 1 180 Psi Psi -180 -180 -180 Phi 180 Phi 180 -180 Glycine 180 180 Psi Psi 0 -180 Phi -180 -180 Phi 180 Trans proline Cis proline 180 180 Psi Psi

94,7% (214/226) of all residues were in favored (98%) regions. 100,0% (226/226) of all residues were in allowed (>99.8%) regions.

Phi

180

-180

There were no outliers.

-180

-180

Phi