MolProbity Ramachandran analysis

1aru_.B99990011.pdb, model 1 General case 180 Psi -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

-180

96,1% (321/334) of all residues were in favored (98%) regions. 99,7% (333/334) of all residues were in allowed (>99.8%) regions.

Phi

There were 1 outliers (phi, psi): 294 SER (131,4, 142,6)

-180

Phi