MolProbity Ramachandran analysis

1agg_.B99990049.pdb, model 1 General case 180 Psi Psi 0 46 SER -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

78,3% (36/46) of all residues were in favored (98%) regions. 97,8% (45/46) of all residues were in allowed (>99.8%) regions.

Phi

There were 1 outliers (phi, psi): 46 SER (175,5, 54,1)

-180

Phi