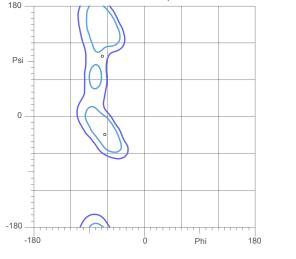
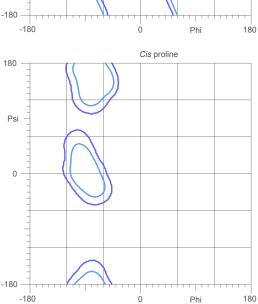
MolProbity Ramachandran analysis

1agg_.B99990015.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 Trans proline Cis proline 180 180 Psi Psi





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.

There were 1 outliers (phi, psi): 46 SER (-179,1, 67,3)