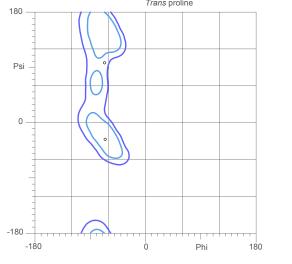
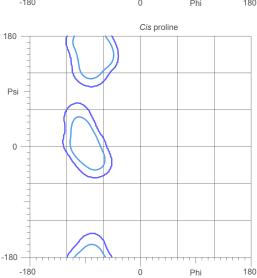
MolProbity Ramachandran analysis

1agg_.B99990007.pdb, model 1 180 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





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 (-174,4, 66,8)