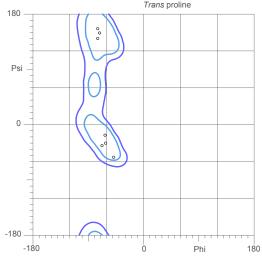
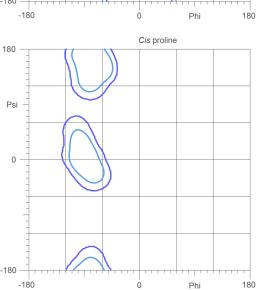
MolProbity Ramachandran analysis

1aac_.B99990025.pdb, model 1 General case 180 Psi Psi -180 -180 -180 Phi 180 -180 Phi 180 Glycine 180 180 Psi Psi 0 -180 Phi 180 -180 -180 Phi Trans proline Cis proline 180 180





98,1% (101/103) of all residues were in favored (98%) regions. 100,0% (103/103) of all residues were in allowed (>99.8%) regions.

There were no outliers.