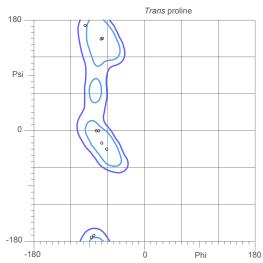
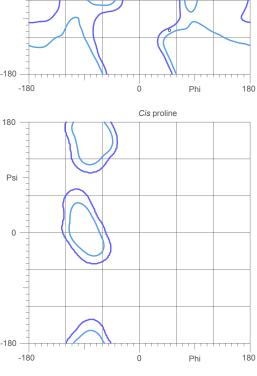
## MolProbity Ramachandran analysis

1avqA.B99990007.pdb, model 1 180 Psi -180 -180 -180 Phi 180 -180 Phi 180 Glycine 180 180 Psi Psi 0 -180 Phi -180 -180 Phi Trans proline Cis proline 180 180





95,1% (215/226) of all residues were in favored (98%) regions. 100,0% (226/226) of all residues were in allowed (>99.8%) regions.

There were no outliers.