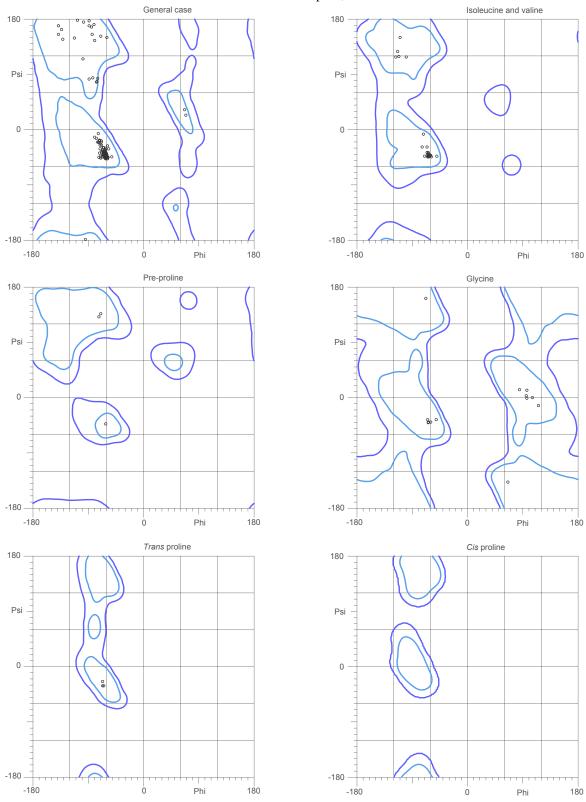
MolProbity Ramachandran analysis

1auiB.B99990008.pdb, model 1



100,0% (163/163) of all residues were in favored (98%) regions. 100,0% (163/163) of all residues were in allowed (>99.8%) regions.

There were no outliers.