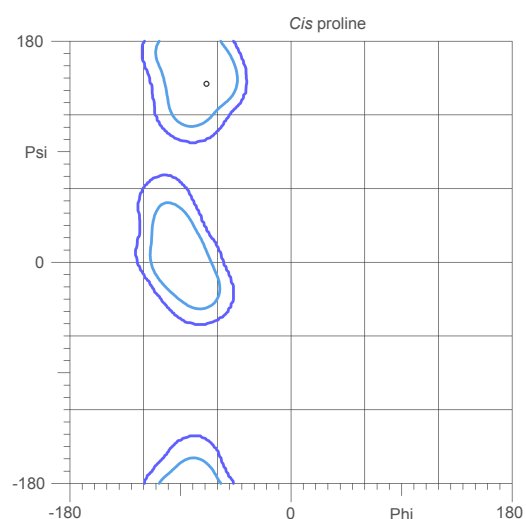
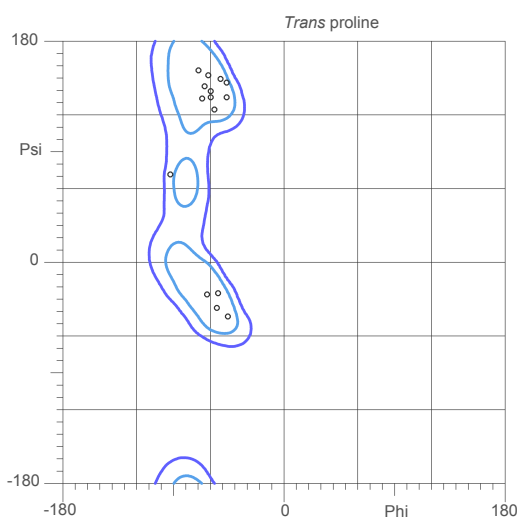
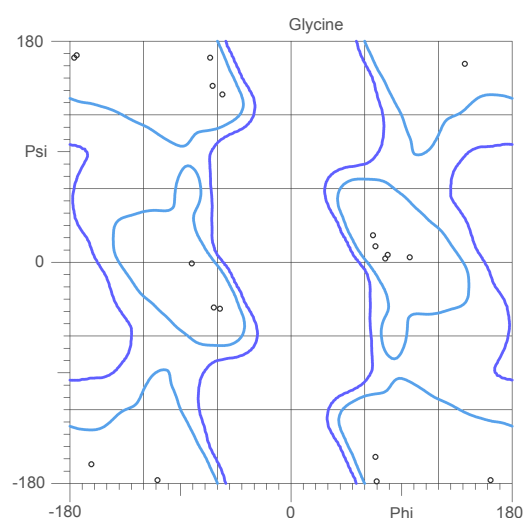
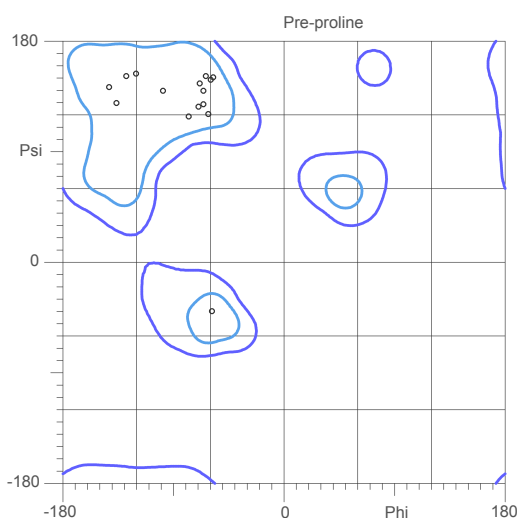
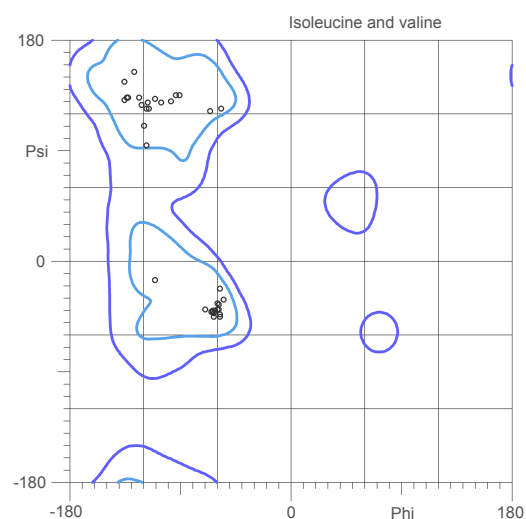
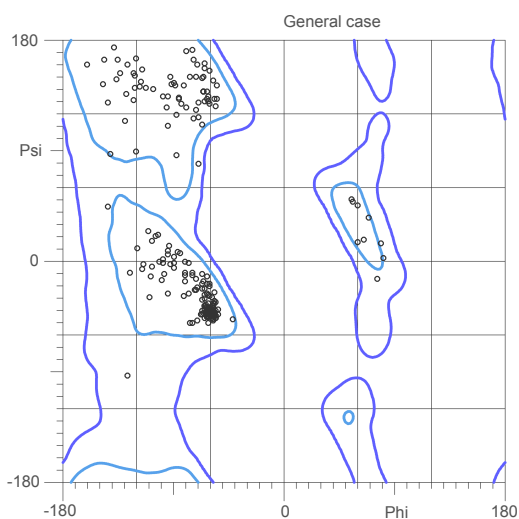


# MolProbity Ramachandran analysis

model\_01.pdb, model 1



97.6% (288/295) of all residues were in favored (98%) regions.  
100.0% (295/295) of all residues were in allowed (>99.8%) regions.

There were no outliers.