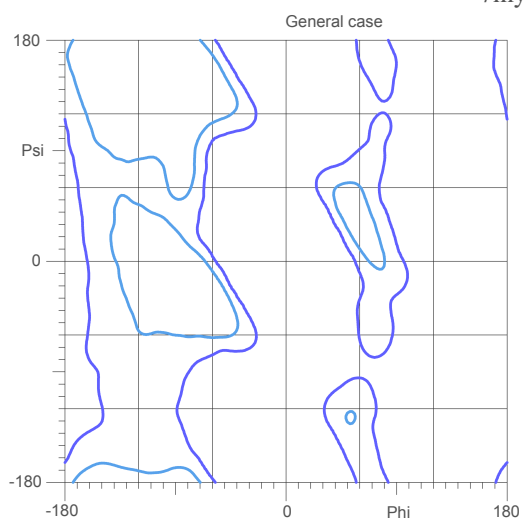


# MolProbity Ramachandran analysis

7my2.H.pdb, model 1



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

There were no outliers.