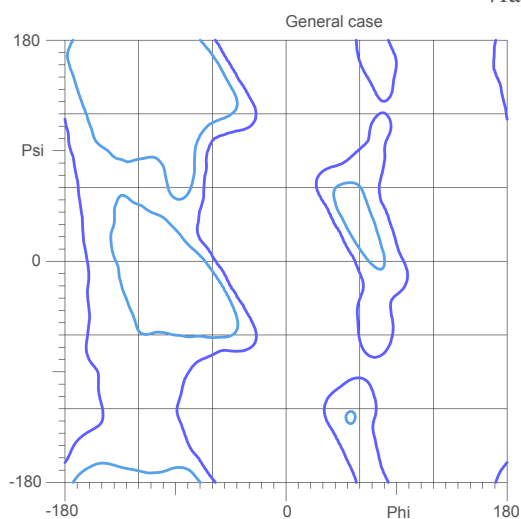


MolProbity Ramachandran analysis

7fay.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.