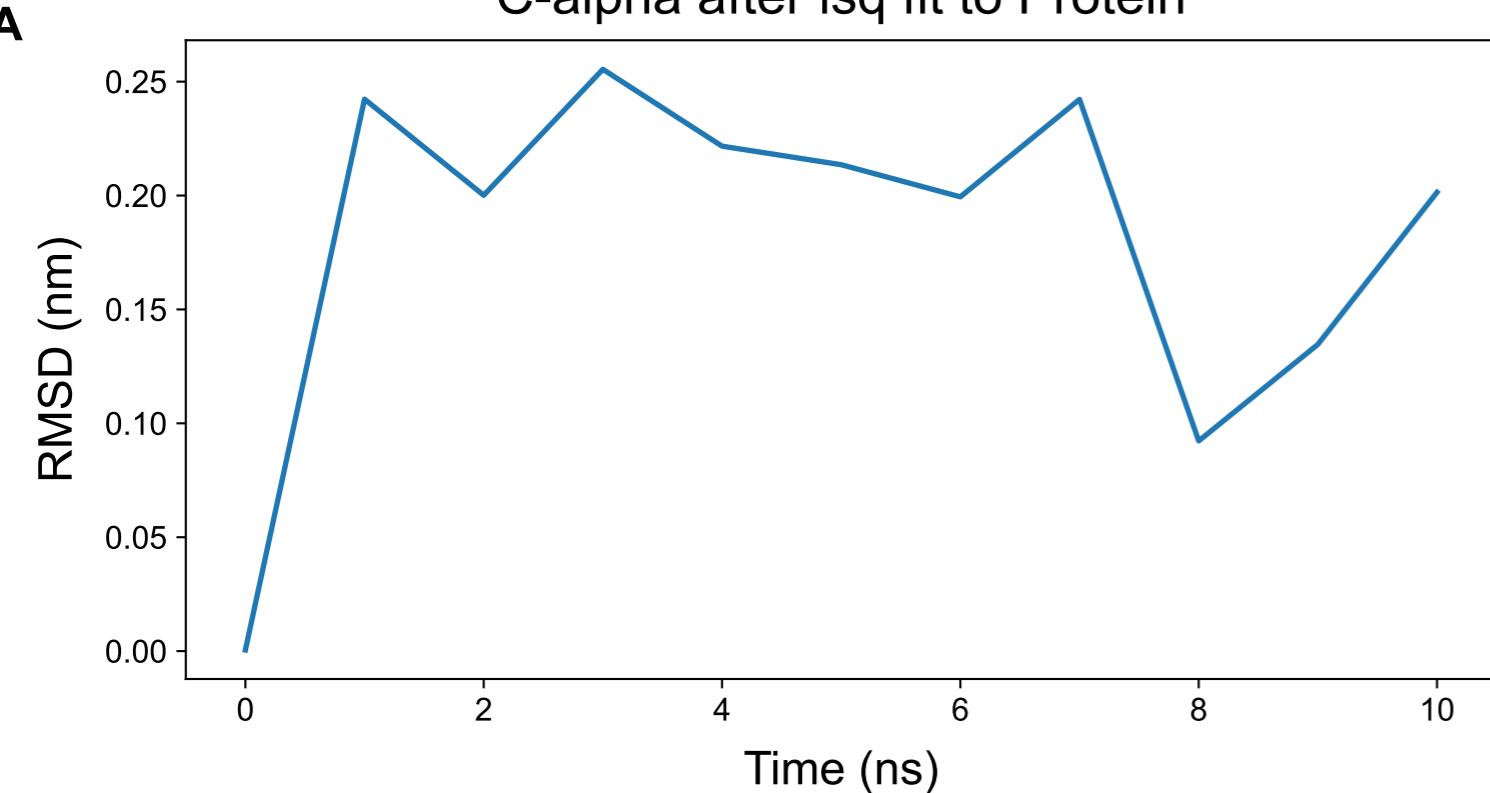
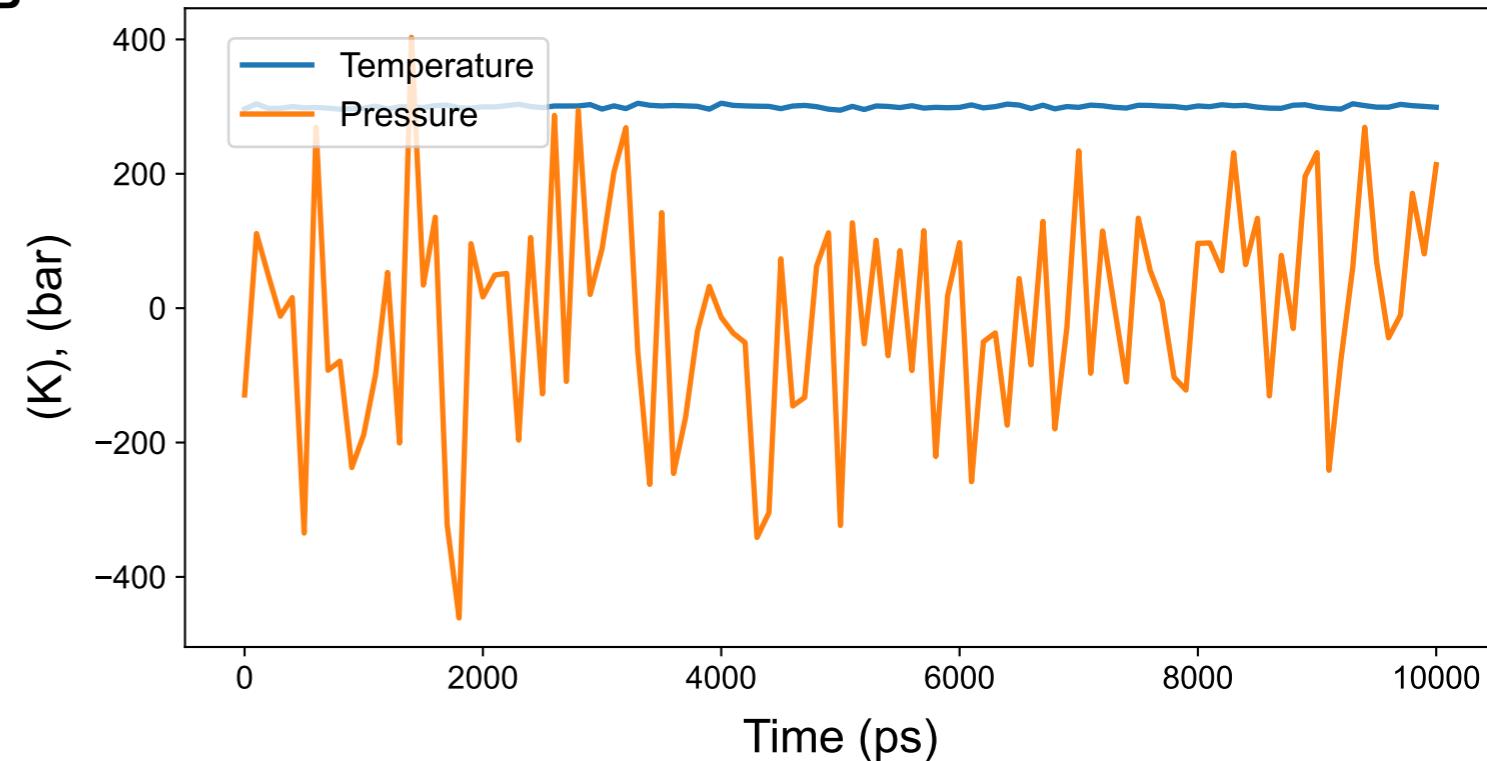


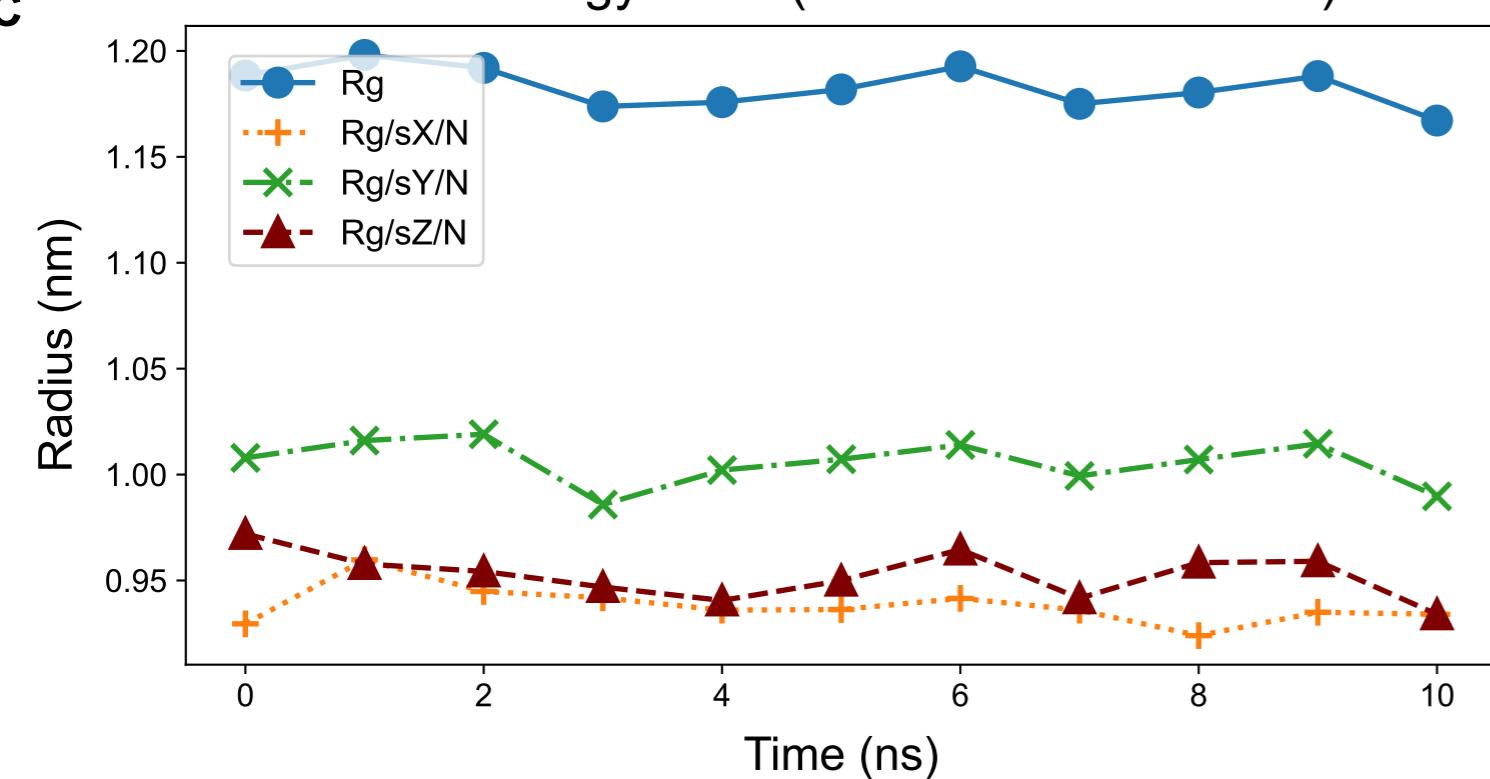
**RMSD**  
C-alpha after lsq fit to Protein



**GROMACS Energies**



**Radius of gyration (total and around axes)**



**2D projection of trajectory**

