

NAMD configuration file

#####

```
set nCARBONS      ;# 6,8,10,12,14
set nQUARTERS     ;# 0,1,2,3,4
set TIME          ;# total time at end of this simulation (ie: 06)

set MOLECULE       tsb35-c${nCARBONS}
set LATTICE        ${MOLECULE}_0${nQUARTERS}-quarter

set temperature   ;# new temperature (in Kelvin)

set RESTARTTIME   ;# total time of previous simulation (ie: 12)
set RESTARTTEMP   ;# previous temperature

set runSteps      ;# numsteps for this simulation (ie: 6000000)
```

#####

```
timestep          1.0
firsttimestep     0
stepspercycle     4

cutoff            10.0
switching         on
switchdist       8.0
pairlistdist     12.0
margin           1.0

coordinates       ../../${LATTICE}_simPDB.pdb
structure         ../../${LATTICE}_simPSF.psf
parameters        ../../../CHARMM-parameters/${MOLECULE}_charmm.params
paraTypeCharmm    on

wrapAll           on
wrapNearest       on

exclude          scaled1-4
1-4scaling        0.4

temperature       $temperature
rescaleFreq       10
rescaleTemp       $temperature

CoMmotion         yes
rigidBonds        all

fixedAtoms        on
fixedAtomsFile    ../../${LATTICE}_fixedPDB.pdb
fixedAtomsCol     B

outputname        ${LATTICE}_${temperature}K_${TIME}ns_OUTPUT
```

```
binaryoutput      no
outputEnergies    1000

restartname        ${LATTICE}_${temperature}K_${TIME}ns_RESTART
restartfreq        5000

DCDfile            ${LATTICE}_${temperature}K_${TIME}ns_DCD.dcd
DCDfreq            1000

if {1} {
set inputname
../../${RESTARTTEMP}K/${RESTARTTIME}ns/${LATTICE}_${RESTARTTEMP}K_${RESTARTTIME}ns_RESTART
binCoordinates     ${inputname}.coord
extendedSystem      ${inputname}.xsc
#binVelocities      ${inputname}.vel      ;# DO NOT USE temperature OR reinitvels WITH THIS
}

reinitvels         $temperature

run                $runSteps
```