## Simulation Methods in Physics I

# Worksheet 3: Molecular Dynamics 2 and Observables

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## 1 Command line parameters

For the simulation different command line options were defined. In the following code blocks you can see them. They will be explained during the report.

```
Code block 1: Commands for ljsim.py
                                                 script: src/ljsim.py
14 # command line arguments
15 parser = argparse.ArgumentParser()
16 parser.add_argument("--cont", type=double, help="continue
      calculation with for cont further time")
17 parser.add_argument("--time", type=double, help="How long
      do you want to run the simulation? | default time=10")
18 parser.add_argument("--tstat", type=double, help="Uses a
      thermostat with a given temperature")
19 parser.add_argument("--warm", type=double, help="Use the
     warming up | pass force")
20 parser.add_argument("--ctstat", type=double, help="
      continue for the simulation with tstat")
21 parser.add_argument("--cwarm", type=double, help="continue
       for the simulation with warm")
22
23 args = parser.parse_args()
8 # command line arguments
9 parser = argparse.ArgumentParser()
10 parser.add_argument("--M", type=int, default=10, help="
      window size | default: M=10")
11 parser.add_argument("--datafile", type=str, default='../
      dat/ljsim.dat', help="datafilename | default: '../dat/
      lisim.dat'")
12 parser.add_argument("--teq", type=double, help="
      equilibration time | calculates averages after
      equilibration time")
13 parser.add_argument("--tlim", type=double, nargs='+', help
     ="type in the time limits you want to plot")
14
15 args = parser.parse_args()
```

## 2 Restart the program where left it

First we want to change the programm in a way, that it is able to restart the simulation where it ended last time. The first thing that we have to do is to store the necessary information about the end state of the system, the position x and velocity v of each particle, into the file ljsim.dat. This happens in code block 3. Furthermore there are the new variables Ts and Ps. The represent the temperature and pressure of the system over time and will be explained in the next chapter.

Now this data must be read by ljsim.py. Therefore we define a command line option —cont which accepts as argument the time for which the simulation should be continued (code block 1). If the parameter is not used the simulation will start new. Furtehrmore there is another argument —time which takes the simulation time for a new simulation. The code in code block 4 does exactly this.

```
Code block 4: Continue simulation
```

```
script: src/ljsim.py
```

```
74
  # Import previous data
75
   if args.cont:
76
            # open datafile
77
            datafile = open(datafilename, 'r')
            ts, Es, Ts, Ps, x, v, hs = pickle.load(datafile)
78
79
            datafile.close()
80
81
            \# length of run
82
            t = ts[-1]
            tmax = t+args.cont
83
84
            step = 0
85
   else:
86
87
            \# length of run
            if args.time:
88
89
                     tmax = args.time
90
            else:
91
                     tmax = 10.0
```

## 3 Simple Observables

### 3.1 Expanding Energy Calculation

First the energy calculation should be changed in order to also get the kinetic and potential energy. This is done by changing the compute\_energy() function call and the parameter Es in ljsim.py (code block 5).

```
script: src/ljsim.py
        if step % measurement_stride == 0:
189
            E_pot, E_kin, E_tot = compute_energy(x, v)
190
            T = 2*E_kin/(3*N)
191
            P = compute_pressure(E_kin, x)
192
193
            print("t={}:\n\tE_pot={}\n\tE_kin={}\n\tE_tot={}\n
                \tT={}\n\tP={}\". format(t, E_pot, E_kin, E_tot,
               T, P))
             compute_distances(x, r)
194
            h = compute_histogram(r)
195
196
             ts.append(t)
197
198
            Es.append([E_pot,E_kin,E_tot])
199
            Ts.append(T)
200
            Ps.append(P)
201
            hs.append(h)
```

#### 3.2 Calculating the Temperature

The temperature of the system can be calculated from the kinetic Energy E\_kin like in code block 5 shown. This is a single scalar calculation and therefore can be done in python.

#### 3.3 Calcating Pressure

#### Derivation of the pressure

The virial of a system is defined as:

$$G = \sum_{i}^{N} \boldsymbol{p}_{i} \cdot \boldsymbol{r}_{i} \tag{1}$$

In the case

$$0 = \left\langle \frac{\mathrm{d}G}{\mathrm{d}t} \right\rangle = \left\langle \sum_{i}^{N} \frac{\boldsymbol{p}_{i}^{2}}{m_{i}} \right\rangle + \sum_{i}^{N} \left\langle \boldsymbol{F}_{i} \cdot \boldsymbol{r}_{i} \right\rangle \tag{2}$$

the following equation can be derived:

$$-\sum_{i}^{N} \langle \mathbf{F}_{i} \cdot \mathbf{r}_{i} \rangle = \left\langle \sum_{i}^{N} \frac{\mathbf{p}_{i}^{2}}{m_{i}} \right\rangle = 2 \langle E_{\text{kin}} \rangle = 3Nk_{B}T \stackrel{id.gas}{=} 3PV$$
 (3)

For a lennard jones pair interaction (like in the simulation) let the forces  $f_{ij}$  and the distance vectors  $\mathbf{r}_{ij}$  be defined as follows. Thereby  $f^{lj}(|\mathbf{r}_{ij}|)$  is the scalar lennard jones force for the distance  $|\mathbf{r}_{ij}|$ , where a positive value means repulsion.

$$\boldsymbol{r}_{ij} = \boldsymbol{r}_j - \boldsymbol{r}_i \tag{4}$$

$$\mathbf{f}_{ij} = f^{\mathrm{lj}}(|\mathbf{r}_{ij}|) \cdot \frac{\mathbf{r}_{ij}}{|\mathbf{r}_{ij}|} \tag{5}$$

$$\mathbf{f}_{ij} = -\mathbf{f}_{ji} \quad \mathbf{r}_{ij} = -\mathbf{r}_{ji} \tag{6}$$

With this knowldege we can derive the pressure of the system by splitting up the force  $\mathbf{F}$  in equation (3) into an ideal gas part  $\mathbf{F}^{\text{id.gas}}$  and a lennard jones part  $\mathbf{F}^{\text{lj}}$ .

$$-\sum_{i}^{N} \langle \mathbf{F}_{i} \cdot \mathbf{r}_{i} \rangle = -\sum_{i}^{N} \left\langle \mathbf{F}_{i}^{\text{id.gas}} \cdot \mathbf{r}_{i} \right\rangle - \sum_{i}^{N} \left\langle \mathbf{F}_{i}^{\text{lj}} \cdot \mathbf{r}_{i} \right\rangle$$
(7)

$$\left\langle \sum_{i}^{N} \frac{\boldsymbol{p}_{i}^{2}}{m_{i}} \right\rangle = 3PV - \sum_{i}^{N} \left\langle \boldsymbol{F}_{i}^{lj} \cdot \boldsymbol{r}_{i} \right\rangle \tag{8}$$

$$P = \frac{Nk_BT}{V} + \frac{1}{3V} \sum_{i}^{N} \langle \mathbf{F}_{i,\text{interaction}} \cdot \mathbf{r}_i \rangle$$
 (9)

$$= \frac{1}{3V} \left[ \left\langle \sum_{i}^{N} \frac{\boldsymbol{p}_{i}^{2}}{m_{i}} \right\rangle + \left\langle \sum_{i,j\neq i}^{N} -\boldsymbol{f}_{ij} \cdot \boldsymbol{r}_{i} \right\rangle \right]$$
(10)

$$= \frac{1}{3V} \left[ \left\langle \sum_{i}^{N} \frac{\boldsymbol{p}_{i}^{2}}{m_{i}} \right\rangle + \left\langle \sum_{i,j>i}^{N} \boldsymbol{f}_{ij} \cdot \boldsymbol{r}_{j} - \boldsymbol{f}_{ij} \cdot \boldsymbol{r}_{i} \right\rangle \right]$$
(11)

$$= \frac{1}{3V} \left[ \left\langle \sum_{i}^{N} \frac{\boldsymbol{p}_{i}^{2}}{m_{i}} \right\rangle + \left\langle \sum_{i,j>i}^{N} \boldsymbol{f}_{ij} \cdot \boldsymbol{r}_{ij} \right\rangle \right]$$
(12)

$$= \frac{1}{3V} \left[ 2 \langle E_{\text{kin}} \rangle + \left\langle \sum_{i,j>i}^{N} \mathbf{f}_{ij} \cdot \mathbf{r}_{ij} \right\rangle \right]$$
 (13)

This works also for a non lennard jones pair interaction with analogous definition. In the programm we will use equation (13) but we use the actual state instead of calculating the expection values.

#### Pressure in Cython

Becaus of the vectorial calculations for each particle pair the calculation has to be done in C if it should be fast. Therefor the function c\_compute\_pressure() is written in c\_lj.cpp.

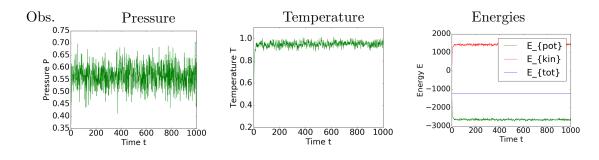
```
Code block 6: Pressure calculation in C
                                                       script: src/c_lj.cpp
253
      double c_compute_pressure(double E_kin, double* x){
        double rij[3];
254
        double fij[3];
255
        double interaction = 0.0;
256
257
258
        // \ for \ (int \ i = 1; \ i < N; \ i++)
           for (int j = 0; j < i; j++) 
259
260
        // add up fij*rij
261
262
        vector<int>::iterator it = verlet_list.begin();
263
        vector<int>::iterator end = verlet_list.end();
        while (it != end) {
264
           int i = *it;
265
266
           ++it;
           int j = *it;
267
           ++it;
268
269
           if (i!=j){
270
271
             minimum_image(x, i, j, rij);
272
             compute_lj_force(rij,fij);
273
274
             for (int k = 0; k < 3; k++)
               interaction += fij[k]*rij[k];
275
276
           }
277
        return (2.*E_kin+interaction)/(3*L*L*L);
278
279
```

As you can see the first part is the same like in c\_compute\_forces(), but the last part differs. In order to save run time the function takes the kinetic energy, which is already calculated, as an argument.

## 4 Equilibration

It is important that we allow the system to equilibrate before we can trust our data. We can tell that our simulation has equilibrated when there is no longer a time drift in any of our observables. In the following figures, we plot our observables over 1000 timesteps in order to see the cessation in drift in our observables.

In order to smooth out these plots we would like to compute the running average of



the observables. This is given by

$$\hat{O}_i = \frac{1}{M+1} \sum_{i-M/2}^{i+M/2} O_i \tag{14}$$

where M is the window size of the average. We implemented the running average in python in the file ljanalyze.py.

```
Code block 7: Running Average
```

script: src/ljanalyze.py

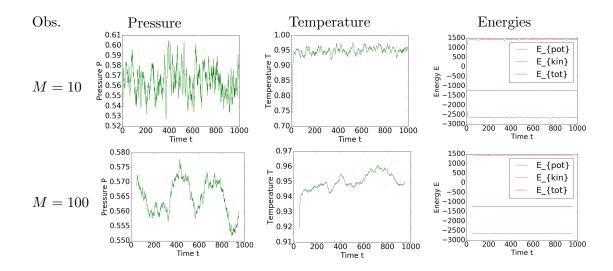
```
33
34
   def compute_running_average(0,M):
           m = M/2
35
           N = len(0)
36
            0a = zeros(len(0))
37
                                                                 #
               new empty array for average
38
            Oa[:m] = nan
                                                                 #
               set values to NaN
            0a[-m:] = nan
39
            for k in xrange(m,N-m):
40
                                                                 #
               loop over all observables
41
                     for 1 in xrange(k-m,k+m+1):
                                                                 #
                        loop over window values
42
                             0a[k] += 0[1]
```

The following figures show the running averages with two different window sizes M. All three observables seem to equilibrate within 100 timesteps. The pressure starts close to equilibrium so that its fluctuations are larger than the initial equilibration. The temperature also quickly equilibrates but changes from a much different nonequilibrium state. The energies quickly exchange between kinetic and potential before coming to equilibrium. Our plots make it clear that 200 timesteps is plenty of time to allow the system to equilibrate.

Finally, we extend ljanalyze.py to compute the average value of the observables after equilibration. This is done with the following function.

#### Code block 8: Compute Mean

script: src/ljanalyze.py



We pass the number of timesteps needed for equilibration in with a command line argument and our code returns the mean values. In our case we measured  $E_{\rm pot}=-2652.8$ ,  $E_{\rm kin}=1427.8$ ,  $E_{\rm tot}=-1225.0$ , T=0.952, and P=0.568.

## 5 Molecular Dynamics at a Desired Temperature

#### **Programming**

For the velocity rescaling thermostat we can derive the rescaling-factor  $f_{re}$  from equation (15).

$$\frac{3}{2}k_B T_0 = \frac{E_{\text{kin},0}}{N} \tag{15}$$

$$= \frac{1}{N} \sum_{i=1}^{N} \frac{(f_{\text{re}} \cdot \boldsymbol{v}^{(i)})^2}{2m}$$
 (16)

$$= f_{\rm re}^2 \frac{E_{\rm kin}}{N} \tag{17}$$

$$= f_{\rm re}^2 \frac{3}{2} k_B T \tag{18}$$

$$f_{\rm re} = \sqrt{\frac{T_0}{T}} \tag{19}$$

This rescaling is implemented in C. In fact the implementation in C is not much faster than in python, but the thermostat in python did some crazy stuff. The function c\_velocity\_rescaling can be seen in code block 9.

```
Code block 9: Velocity rescaling

void c_velocity_rescaling(double T0, double T, double* v
) {
    double f = sqrt(T0/T);
    for (int k = 0; k < 3*N; k++) {
        v[k] *= f;
}
</pre>

Code block 9: Velocity rescaling

double T, double* v
) {
    double T, double* v
) {
        void c_velocity_rescaling(double T0, double T, double* v
) {
        void c_velocity_rescaling(double T0, double T, double* v
) {
        void c_velocity_rescaling(double T0, double T, double* v
) {
        void c_velocity_rescaling(double T0, double T, double* v
) {
        void c_velocity_rescaling(double T0, double T, double* v
) {
        void c_velocity_rescaling(double T0, double T, double* v
) {
        void c_velocity_rescaling(double T0, double T, double* v
) {
        void c_velocity_rescaling(double T0, double T, double* v
) {
        void c_velocity_rescaling(double T0, double T, double* v
) {
        void c_velocity_rescaling(double T0, double T, double* v
) {
        void c_velocity_rescaling(double T0, double T, double* v
) {
        void c_velocity_rescaling(double T0, double T, double* v
) {
        void c_velocity_rescaling(double T0, double T, double* v
) {
        void c_velocity_rescaling(double T0, double T, double* v
) {
        void c_velocity_rescaling(double T0, double T, double* v
) {
        void c_velocity_rescaling(double T0, double T, double* v
) {
        void c_velocity_rescaling(double T0, double T, double* v
) {
        void c_velocity_rescaling(double T0, double T) {
        void c_velocity_r
```

This function is used in the main loop in ljsim.py every time after meausuring the observalbles.

To start the thermostat you can use the command line option ——tstat which accepts the desired temperature as argument (see code block 1). The option ——ctstat is there for continuing the simulation for the interrupted simulation with temperature T thermostat, but now with deactivated thermostat. This is necessary because the programm saves the results of the simulation with different names for different temperatures.

#### Measurement

Now we can test the simulation for desired temperatures of  $T\epsilon \{0.3, 1.0, 2.0\}$ . Therefor we must type into the command line:

>> python ljsim.py --tstat T0 --time 1000.

Figure 3 shows the results respectively.

We can easily see the system reach the desired temperature and the energies reach equilibrium. The pressure is a bit harder to see as the fluctuations are so large and take large amounts of time.

As you can see the equilibration time seems to be highest for  $T_O = 0.3$  and lowest for  $T_O = 2.0$ . This is because the temperature  $T_0 = 0.3$  is much lower than the initial temperature. Therefor the total energy of the system must decrease. But the positive kinetic energy is much lower, than the necessary energy difference. If the temperature should be T = 0.3 the potential energy of the system must decrease, but it is not directly effected by the velocity rescaling. By contrast for  $T_0 = 2.0$  the necessary energy can simply be reached by increasing the kinetic energy because it is much higher than the initial energy. For  $T_0 = 1.0$  the necessary energy is not much different from the initial one and therefor the other effects on equilibration have a higher influence.

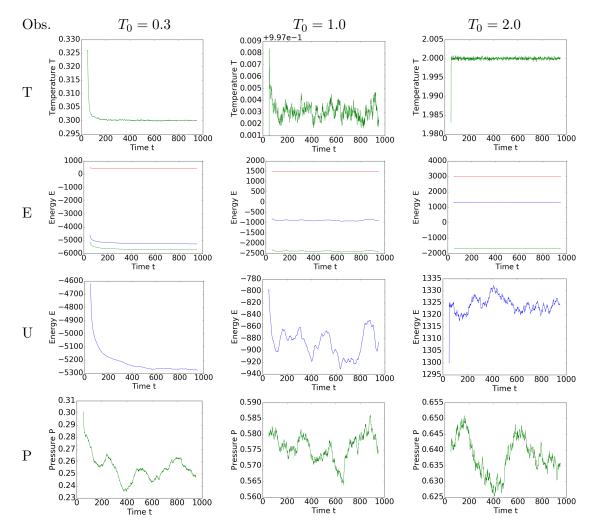


Fig. 3: Measured Observables (Obs.) for a thermostat temperature  $T_O$ . T is the real temperature of the system, E are the energies (green: potential, red: kinetic, blue: total).

U is the total energy of the system alone.

P is the pressure of the system.

## 6 Warming up the System

When warming up the system you want to state a initial maximum force. Therefor the command line option ——warm can be used (code block 1). The option ——cwarm works analougus to ——ctstat.

The first thing needed is the function c\_force\_capping() which is implemented in C. The reason for this is because you have to do many if requests and calculations for each particle.

```
Code block 10: Force capping
                                                       script: src/c_lj.cpp
      double c_force_capping(double* f, double fcap){
288
        double fcap2 = fcap*fcap;
289
        double test_cap = true; // if true: no force is capped
290
             any more
291
        for (int k = 0; k < N; k++) {
           double fabs2 = f[k]*f[k] + f[k+1]*f[k+1] + f[k+2]*f[
292
              k+2];
           if (fabs2 > fcap2) {
293
             test_cap = false;
294
             double ff = sqrt(fcap2/fabs2);
295
             f[k] *= ff;
296
297
             f[k+N] *= ff;
             f[k+2*N] *= ff;
298
           }
299
300
301
        if (test_cap){
302
           fcap = 0;
303
304
      return fcap;
305
```

The function in code block 10 limits the forces to a given maximum force fcap, but conserves the direction if the forces. Although this process is not physically correct. Furthermore the function tests if no force is capped any more. When this is the case it sets fcap to zero. The function is called in the vv\_step() function in ljsim.py.

```
Code block 11: Limiting forces

# compute new forces

# we assume that m=1 for all particles

f = compute_forces(x)

if args.warm:

force_capping(f,args.warm)
```

As you can see, the function will not be activated any more if fcap = args.warm is set to zero.

At least the fcap should be increased by ten percent every time the measurements are done. This is done in code block 12.

```
Code block 12: Increase maximum force
                                                               script: src/ljsim.py
213
               \# rescale fcap
214
               if args.warm:
215
                         args.warm *= 1.1
```

But all the capping is useless without random initial conditions. The velocity is already random, so only the initial position has to be changed. You can do this with a similar command like in 13.

```
Code block 13: Random initial position
                                                          script: src/ljsim.py
98
              # Initialize particle position
              if args.warm:
99
100
                       # The warming up random positions
101
                       x = L*random.random((3,N))
```

#### 7 Radial Distribution Function

An important observable for our simulation is the radial distribution function. It is the probability to find a particle at a given radial distance away from another particle compared to an ideal gas. Mathematically it is written as

$$g(r) = \frac{1}{\rho 4\pi \mathbf{r}^2 dr} \sum_{i,j} \langle \delta(r - |\mathbf{r}_{ij}|) \rangle$$
 (20)

For an ideal gas we expect g(r) = 1 if we normalize correctly.

In our simulation we implement the rdf using both C and python. First we calculate all the pair distances for our particles in C.

```
Code block 14: Radial Distances
                                                        script: src/c_lj.cpp
       void c_compute_distances(double* x, double* r) {
307
308
           double rij[3];
309
           int k = 0;
           for ( int i = 0; i < N; i++) {
310
              for ( int j = i+1; j < N; j++) {
311
                  minimum_image(x, i, j, rij);
312
                  r[k] = sqrt(rij[0]*rij[0]+rij[1]*rij[1]+rij
313
                     [2]*rij[2]);
```

We then send this to an array in python and compute a histogram using numpy and normalize it in the correct way

Finally we implement an average RDF after equilibrium in ljanalyze.py. We compute this using the following function

```
Code block 16: Compute Mean RDF

script: src/ljanalyze.py

def compute_mean_rdf(0,keq):

N = 0.shape[0]

Om = zeros(0.shape[1])

for k in xrange(keq,N):

Om += 0[k,:]
```

and plot it.

## 8 Measuring Equilibrium Mean Values of the Observables

We now run the simulation for the same three temperatures 0.3, 1.0, 2.0 including random particle positions and the previously described warm up. We run the simulations for 1200 time steps. Using our command line arguments an example looks like:

```
python ljsim.py --warm 20 --tstat 0.3 --time 1200
```

This sets an inital force cap of 20. The equilibration of the observables is plotted in the following figures with the first 100 timesteps omitted to preserve plot scaling. The command to produce these plots with our command line arguments looks like:

```
python ljanalyze.py --datafile ../dat/ljsim_T2d0_F20d0.dat --tlim 100 1199
--M 100
```

Here we used a window of M = 100.

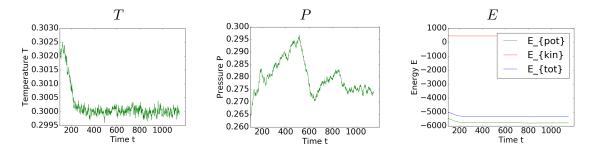


Fig. 4: Desired Temperature:  $T_0 = 0.3$ . Average values of temperature, pressure and energies from the left to the right.

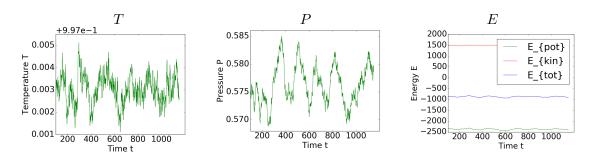


Fig. 5: Desired Temperature:  $T_0 = 1.0$ . Average values of temperature, pressure and energies from the left to the right.

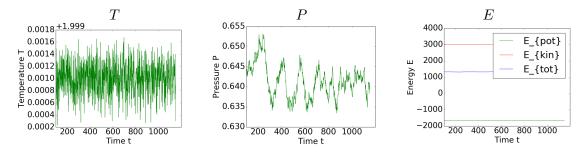
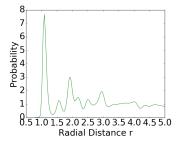


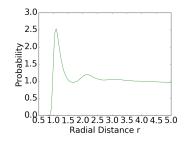
Fig. 6: Desired Temperature:  $T_0 = 2.0$ . Average values of temperature, pressure and energies from the left to the right.

The commands only differed in the thermostat temperature and the datafilename to be analyzed. Finally we plot the average RDF for the three different temperatures after equilibrium and compute the mean equilibrium values of the observables. The command to perform this analysis is:

Here we have assumed that our system has equilibrated by timestep 400. We measured the folloing average values:

$T_0$	$E_{ m pot}$	$E_{ m kin}$	$E_{ m tot}$	T	P
0.3	-5754.9	450.0	-5304.9	0.300	0.281
1.0	-2367.3	1500.1	-867.3	1.000	0.577
2.0	-1679.0	3000.0	1321.0	2.000	0.642





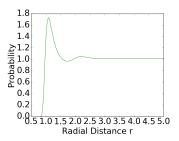


Fig. 7: Radial distribution function (RDF) in equilibrium for desired temperature of  $T_0 = \{0.3, 1.0, 2.0\}$  from the left to the right.