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()
```

Code block 2: Commands for ljanalyze.py — script: src/ljanalyze.py

```
 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/
      ljsim.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.

Code block 3: Storing data script: src/ljsim.py

```
221 # write out simulation data
222 print("Writing simulation data to {}.".format(datafilename
    ))
223 datafile = open(datafilename, 'w')
224 pickle.dump([ts, Es, Ts, Ps, x, v, hs], datafile)
225 datafile.close()
```

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')
78         ts, Es, Ts, Ps, x, v, hs = pickle.load(datafile)
79         datafile.close()
80
81         # length of run
82         t = ts[-1]
83         tmax = t+args.cont
84         step = 0
85
86 else:
87         # length of run
88         if args.time:
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).

Code block 5: $E_{\text{kin}} + E_{\text{pot}}$ script: src/ljsim.py

```
189     if step % measurement_stride == 0:
190         E_pot, E_kin, E_tot = compute_energy(x, v)
191         T = 2*E_kin/(3*N)
192         P = compute_pressure(E_kin, x)
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))
194         compute_distances(x, r)
195         h = compute_histogram(r)
196 
197         ts.append(t)
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

$$0 = \left\langle \frac{\mathrm{d}G}{\mathrm{d}t} \right\rangle = \left\langle \sum_i^N \frac{\boldsymbol{p}_i^2}{m_i} \right\rangle + \sum_i^N \langle \boldsymbol{F}_i \cdot \boldsymbol{r}_i \rangle \tag{2}$$

the following equation can be derived:

$$\left\langle \sum_i^N \frac{\boldsymbol{p}_i^2}{m_i} \right\rangle = -\sum_i^N \langle \boldsymbol{F}_i \cdot \boldsymbol{r}_i \rangle = 3Nk_BT \overset{id.gas}{=} 3PV \tag{3}$$

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

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

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

$$\boldsymbol{f}_{ij} = -\boldsymbol{f}_{ji} \tag{6}$$

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

$$-\sum_i^N \langle \boldsymbol{F}_i \cdot \boldsymbol{r}_i \rangle = -\sum_i^N \left\langle \boldsymbol{F}_i^{\mathrm{id.gas}} \cdot \boldsymbol{r}_i \right\rangle - \sum_i^N \left\langle \boldsymbol{F}_i^{\mathrm{lj}} \cdot \boldsymbol{r}_i \right\rangle \tag{7}$$

$$\left\langle \sum_i^N \frac{\boldsymbol{p}_i^2}{m_i} \right\rangle = 3PV - \sum_i^N \left\langle \boldsymbol{F}_i^{\mathrm{lj}} \cdot \boldsymbol{r}_i \right\rangle \tag{8}$$

$$P = \frac{Nk_BT}{V} + \frac{1}{3V} \sum_i^N \langle \boldsymbol{F}_{i,\mathrm{interaction}} \cdot \boldsymbol{r}_i \rangle \tag{9}$$

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

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

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

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

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){
254     double rij[3];
255     double fij[3];
256     double interaction = 0.0;
257
258     // for (int i = 1; i < N; i++)
259     //   for (int j = 0; j < i; j++) {
260
261     // add up fij*rij
262     vector<int>::iterator it = verlet_list.begin();
263     vector<int>::iterator end = verlet_list.end();
264     while (it != end) {
265       int i = *it;
266       ++it;
267       int j = *it;
268       ++it;
269
270       if (i!=j){
271         minimum_image(x, i, j, rij);
272         compute_lj_force(rij,fij);
273
274         for (int k = 0; k < 3; k++)
275           interaction += fij[k]*rij[k];
276       }
277     }
278     return (2.*E_kin+interaction)/(3*L*L*L);
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 Molecular Dynamics at a Desired Temperature

For the *velocity rescaling* thermostat we can derive the rescaling-factor f_{re} from equation (14).

$$\frac{3}{2}k_B T_0 = \frac{E_{\mathrm{kin},0}}{N} \tag{14}$$

$$= \frac{1}{N}\sum_{i=1}^{N}\frac{(f_{\mathrm{re}}\cdot \boldsymbol{v}^{(i)})^2}{2m} \tag{15}$$

$$= f_{\mathrm{re}}^2\frac{E_{\mathrm{kin}}}{N} \tag{16}$$

$$= f_{\mathrm{re}}^2\frac{3}{2}k_B T \tag{17}$$

$$f_{\mathrm{re}} = \sqrt{\frac{T_0}{T}} \tag{18}$$

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 7.

Code block 7: Velocity rescaling — script: src/c_lj.cpp

```
281  void c_velocity_rescaling(double T0, double T, double* v
        ){
282    double f = sqrt(T0/T);
283    for (int k = 0; k < 3*N; k++){
284        v[k] *= f;
285    }
286  }
```

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.

5 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 8: Force capping script: src/c_lj.cpp

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

The function in code block 8 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 9: Limiting forces script: src/ljsim.py

```
164    # compute new forces
165    # we assume that m=1 for all particles
166    f = compute_forces(x)
167    if args.warm:
168        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 10.

Code block 10: 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 11.

Code block 11: Random initial position script: src/ljsim.py

```
98          # Initialize particle position
99          if args.warm:
100                 # The warming up random positions
101                 x = L*random.random((3,N))
```

6 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

$$g(r) = \frac{1}{\rho 4\pi \mathbf{r}^2 dr} \sum_{i,j} \langle \delta(r - |\mathbf{r}_{ij}|) \rangle \tag{19}$$

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 12: Radial Distances script: src/c_lj.cpp

```
307     void c_compute_distances(double* x, double* r){
308        double rij[3];
309        int k = 0;
310        for ( int i = 0; i < N; i++){
311           for ( int j = i+1; j < N; j++){
312              minimum_image(x, i, j, rij);
313              r[k] = sqrt(rij[0]*rij[0]+rij[1]*rij[1]+rij
                    [2]*rij[2]);
314              k++;
315           }
316        }
317     }
```

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

Code block 13: Numpy Histogram script: src/ljsim.py

```
10 def compute_histogram(r):
11         h, bins = histogram(r, bins=100, range=(0.8, 5.),
               density=True)
12         return  h*vol/(4*pi*bins[:-1]*bins[:-1])
```

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

Code block 14: Compute Mean RDF script: src/ljanalyze.py

```
51 def compute_mean_rdf(O,keq):
52     N = O.shape[0]
53     Om = empty(O.shape[1])
54     for k in xrange(keq,N):
55         Om += O[k,:]
56     return Om/(N-keq)
```

and plot it.