

Worksheet 2: Statistical Mechanics and Molecular Dynamics

April Cooper, Patrick Kreissl und Sebastian Weber

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University of Stuttgart

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1 Statistical Mechanics

1.1 Task 1

First we had to consider a system A consisting of two subsystems A_1 and A_2 with the related numbers of configurations $\Omega_1 = 10^{20}$ and $\Omega_2 = 10^{22}$.

The number of configurations available to the combined system is $\Omega = \Omega_1 \cdot \Omega_2 = 10^{42}$.

As the entropy is defined as $S = \ln(\Omega)k_B$ the entropies are:

$$S_1 = k_B \ln(\Omega_1) = k_B \ln(10^{20}), S_2 = k_B \ln(\Omega_2) = k_B \ln(10^{22}), S = k_B \ln(\Omega) = k_B \ln(10^{42}).$$

Then were asked to give the factor by which the number of available configurations increases in a system with given initial V_1, T, p_1 if V_1 was expanded isothermal by 0.001%.

With the help of the first law of thermodynamics it follows that for $T = \text{const}$ it is:

$dS = \frac{p}{T}dV$, which leads with the ideal gas equation to

$$\begin{aligned} \Delta S &= \int_{V_1}^{V_2} \frac{k_B N}{V} dV \\ &= k_B N \ln \left(\frac{V_2}{V_1} \right) \\ &= 10^{-7} k_B N \end{aligned} \tag{1}$$

If you consider now the definition of S: $\Delta S = \ln(\hat{\Delta}\Omega)k_B$ you get with equation (1):

$$\hat{\Delta}\Omega = \exp\left(\frac{\Delta S}{k_B}\right) = \exp(10^{-7}N) = \frac{\Omega_2}{\Omega_1}$$

Finally we were asked to give the factor by which the number of available configurations increases in a system with given initial N, V, T_1 when an energy of 150 kJ is added to the system at constant Volume.

Analogous to the task before it can be derived that $\Delta S = c_v N \ln \left(\frac{T_2}{T_1} \right)$. Using again that the factor by which the number of configurations is given by $\hat{\Delta}\Omega = \exp\left(\frac{S}{k_B}\right)$ it follows that:

$$\hat{\Delta}\Omega = \exp\left(\frac{c_v N \ln\left(\frac{T_2}{T_1}\right)}{k_B}\right) = \exp\left(\frac{c_v N}{k_B}\right) \frac{T_2}{T_1}$$

Using that $T_2 = \frac{\Delta Q}{c_v N} + T_1$ and plugging in the given values leads finally to:

$$\hat{\Delta}\Omega = \exp\left(\frac{2c_v}{k_B}\right) \left(\frac{78}{300c_v} - 1\right) = \frac{\Omega_2}{\Omega_1}$$

1.2 Task 2 - Thermodynamic Variables in the Canonical Ensemble

Given the Helmholtz free energy F we were asked to derive expressions for U, p, S . Using the Maxwell relations it follows:

$$S = -\frac{\partial F}{\partial T} = \ln(Q(N, V, T))k_B + \frac{k_B T}{Q(N, V, T)} \frac{\partial Q(N, V, T)}{\partial T}$$

$$p = -\frac{\partial F}{\partial V} = \frac{1}{\beta Q(N, V, T)} \frac{\partial Q(N, V, T)}{\partial V}$$

The derivation of the equation for U is as follows:

It is known from statistical mechanics that $Q(N, V, T) = \frac{1}{h^{3N} N!} \int d\Gamma \exp(-\beta H)$. The thermodynamic properties of the system can be obtained by $Q(N, V, T) = \exp(-\beta F(N, V, T))$. To justify this identification we show first that F is extensive and then that $F = U - TS$, where $U = \langle H \rangle$.

That F is extensive can be derived directly from $Q(N, V, T) = \frac{1}{h^{3N} N!} \int d\Gamma \exp(-\beta H)$. When the system is split up into two systems with a very weak correlation, then is Q a product of two factors. To show the second equivalence we rewrite $F = U - TS$ to $U = \langle H \rangle = A - T \left(\frac{\partial A}{\partial T} \right)$. To show this we divide the two expressions of Q that we stated before and therefore get:

$$\frac{1}{h^{3N} N!} \int d\Gamma \exp(\beta(F - H)) = 1$$

Deriving both sides by β leads to:

$$\frac{1}{h^{3N} N!} \int d\Gamma \exp(\beta(F - H)) (F - H + \beta \left(\frac{\partial F}{\partial \beta} \right)) = 0$$

This is equivalent to $F - U - T \left(\frac{\partial F}{\partial T} \right)$, wherefore it is: $U = F + TS$.

1.3 Task 3 - Ideal Gas

Given the partition function $Q(N, V, T)$ we had to derive expressions for the free Helmholtz energy $F(N, V, T)$ and the pressure $p(N, V, T)$.

As we know from task 2 it is $F = -\frac{\ln(Q(N, V, T))}{\beta}$. Plugging in the $Q(N, V, T)$ given on the sheet and using the hint leads to:

$$\begin{aligned} F &= -\frac{\ln(Q(N, V, T))}{\beta} \\ &= -k_B T \ln \left(\frac{V^N}{\lambda^{3N} N!} \right) \\ &= -k_B T (N \ln(V) - N \ln(\lambda^3) - \ln(N!)) \\ &= -k_B T N (\ln(V) - \ln(\lambda^3) - \ln(N) + 1) \\ &= k_B T N \left(\ln \left(\frac{N \lambda^3}{V} \right) - 1 \right) \end{aligned} \tag{2}$$

Applying the Maxwell relation $\frac{-\partial F}{\partial V} = p$ on equation (1) it follows:

$$\begin{aligned}
 p &= \frac{-\partial F}{\partial V} \\
 &= \frac{-\partial k_B T N \left(\ln \left(\frac{N \lambda^3}{V} \right) - 1 \right)}{\partial V} \\
 &= \frac{k_B T N}{V}
 \end{aligned} \tag{3}$$

As it is obvious equation(3) leads to the ideal gas law : $pV = Nk_B T$.

2 Molecular Dynamics: Lennard-Jones Fluid

2.1 Implementation of the Lennard Jones Potential

The Lennard Jones Potential is $V_{LJ}(r) = 4\epsilon \left(\left(\frac{\sigma}{r} \right)^{12} - \left(\frac{\sigma}{r} \right)^6 \right)$ and it's straight forward implementation works as follows:

Listing 1: Function compute_lj_potential

```
def compute_lj_potential(rij , eps = EPS, sig = SIG):
    q = sig/np.linalg.norm(rij)
    return 4*eps*(q**12-q**6)
```

Consequently the force being the negative gradient of the potential V_{LJ} is computed by the following function:

Listing 2: Function compute_lj_force

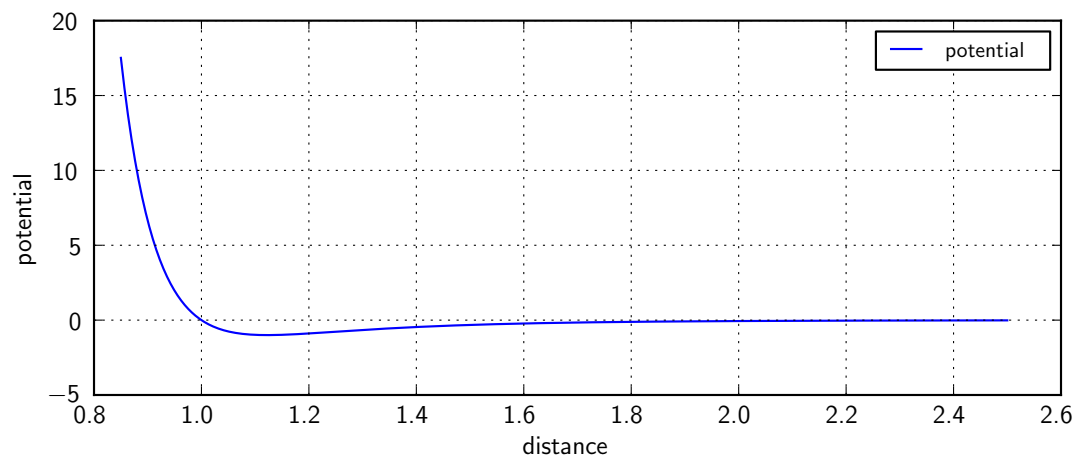
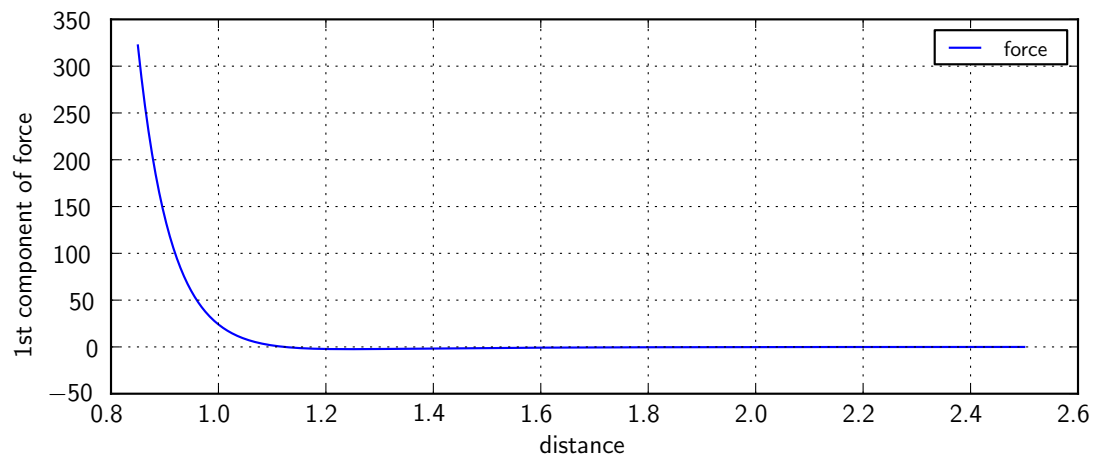
```
def compute_lj_force(rij , eps = EPS, sig = SIG):
    norm = np.linalg.norm(rij)
    q = sig/norm
    return 4*eps*(12*q**11-6*q**5)*q/norm**2*rij
```

To compute the plot of the potential and the first component of the force against d the following routine was implemented:

Listing 3: Computation of the potential and force

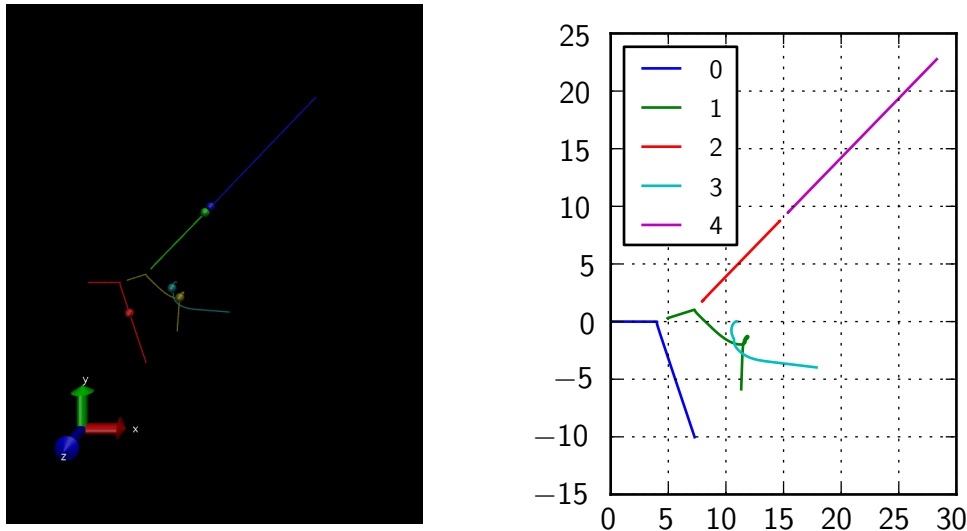
```
d = np.zeros((NParticles,3))
d[:,0] = np.linspace(0.85,2.5,NParticles)
potential = np.array(map(compute_lj_potential , d))
force = np.array(map(compute_lj_force , d))
```

Both, the plot of the potential and the force, show a steep increase for $x \rightarrow 0$ and become constant 0 for $x \rightarrow \infty$. This simply reflects the fact that the force and potential increase with declining distances for already small distances and that the interaction of two particles being far apart is negligible.

*Figure 1: LJ Potential**Figure 2: LJ Force*

2.2 Lennard-Jones Billards

In this task we first had to insert the already programmed routines into the given templates. After that we had to run the program and to visualize the system. The visualization via VMD and the resulting trajectories in the xy-plane are:



As it can be seen in this picture ball 4 has already been moved so that it hits ball 2. The position needed, that this occurs is: $x[:, 4] = [15.4324, 9.51146, 0.0]$

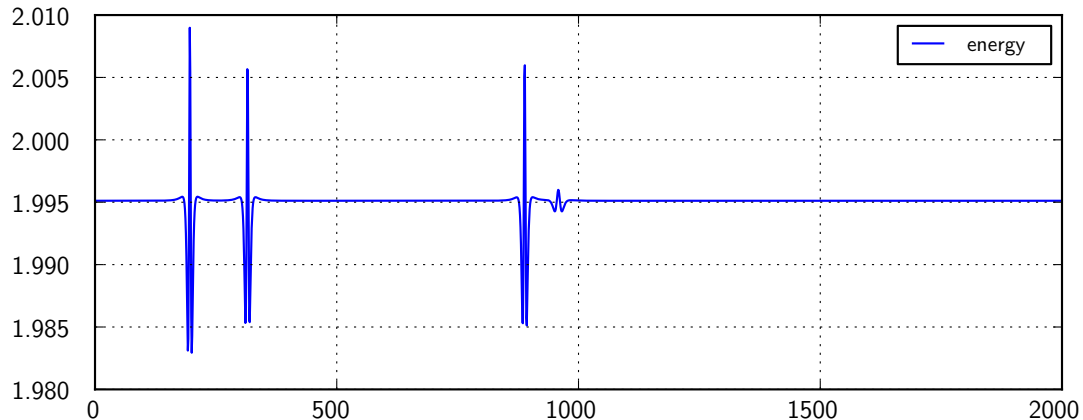


Figure 3: Energy

The energy is constant except from the times when the balls hit each other. In these cases occur relatively small variations in the energy. As it can be seen in Figure 3 the particles 0, 2 and 4 act just like normal billards balls. However the trajectories of the particles 1 and 3 show that the interaction is not based on a hard sphere potential, which would be exact for a billards ball, but on a potential with an attractive part for certain distances. This is obvious as the trajectories show that particle 1 influences particle 3 without hitting it directly which results in a curved trajectory of these two balls. After the distance between the balls becomes big enough the attraction is not acting anymore and the particles follow straight trajectories afterwards.

2.3 Minimum Image Convention and PBC

First we had to implement the potential and the force such that it implements the truncated LJ potential for $r_{cutoff} = 2.5$. The straight forward implementation is analogous to the function implemented before:

Listing 4: Function `compute_lj_potential`

```
def compute_lj_potential(rij , eps = EPS, sig = SIG, rcoeff = RCOFF,\
                        vcoeff = VCOFF):
    norm = np.linalg.norm(rij)
    if norm > rcoeff: return 0
    q = sig/norm
    return 4*eps*(q**12-q**6) - vcoeff
```

Listing 5: Function `compute_lj_force`

```
def compute_lj_force(rij , eps = EPS, sig = SIG, rcoeff = RCOFF):
    norm = np.linalg.norm(rij)
    if norm > rcoeff: return np.zeros(3)
    q = sig/norm
    return 4*eps*(12*q**11-6*q**5)*q/norm**2*rij
```

In order to implement PBC some of the routines have to be changed. First of all the function `compute_forces` had to be updated in order to fulfill the minimum image convention.

Listing 6: Function `compute_forces`

```
def compute_forces(x, L = L):
    """Compute and return the forces acting onto the particles ,
    depending on the positions x."""
    global epsilon , sigma
    _, N = x.shape
    f = np.zeros_like(x)
    for i in range(1,N):
        for j in range(i):
            # distance vector
            rij = x[:,j] - x[:,i]
            rij -= np rint(rij/L)*L #BECAUSE OF PBC
            fij = compute_lj_force(rij)
            f[:,i] -= fij
            f[:,j] += fij
    return f
```

In this case the function `rint()` takes an array and rounds the entries to the closest integer. The same modification had to be done in the function `compute_energy`, meaning that `rij` had again to be rounded with the help of `rint`. As no other things are changed we ask the reader to have a look in the source code in the appendix in order to see the whole function. Another change had to be done in the main loop. As not all particles considered in the box the positions `x` had to be folded back into the central box. This is done as follows:

Listing 7: Folding x back into the central box

```
xfolded = x.copy()
xfolded -= np.floor(x/L)*L #BECAUSE OF PBC
```

Finally also the trajectories had to be folded back before being plotted. As it works just as the back folding of x we again ask the reader to have a closer look at the source code for further information. Now the implemented PBC were tested using a simulation of LJ billiards balls. The main loop of the program is implemented as

Listing 8: Test of the PBC using ljbiliards

```
# main loop
while t < tmax:
    x, v, f = step_vv(x, v, f, dt)
    t += dt

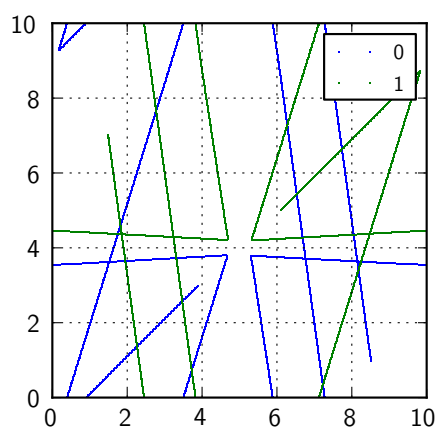
    traj.append(x.copy())
    Es.append(compute_energy(x, v))

    # write out that a new timestep starts
    vtffile.write('timestep\n')
    # write out the coordinates of the particles
    for i in range(N):
        vtffile.write("%s %s %s\n" % (x[0,i], x[1,i], x[2,i]))

vtffile.close()

traj = np.array(traj)
```

The resulting trajectories are:

*Figure 4: Trajectories of the two particles for 20 timesteps*

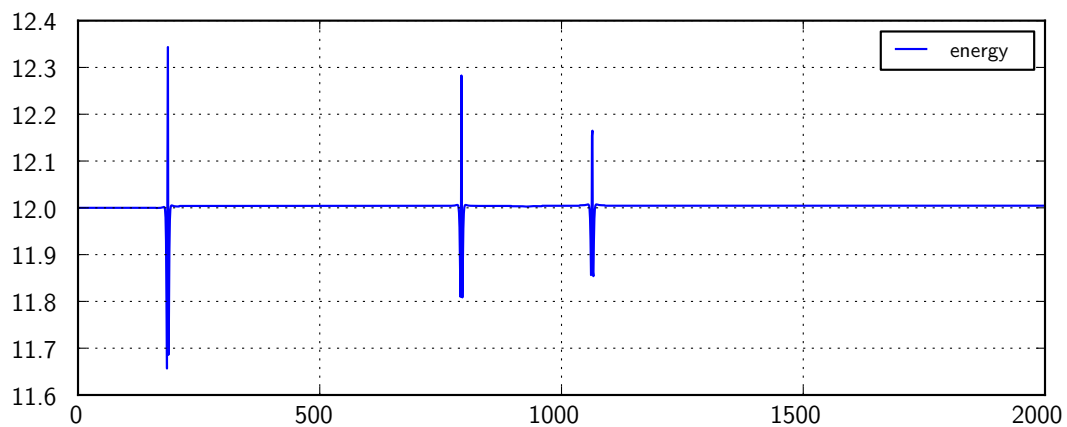


Figure 5: Energy - truncated potential

In the plot of the trajectories it becomes obvious that the PBC work, as when a particle leaves the box on one side the trajectory seems, after being folded back into the central box, to continue on the opposite side. Again the energy stays constant and shows that the particles hit each other thrice during the simulation time, which can be seen at the sharp energy peaks that have already been described before.

3 Lennard- Jones Fluid

3.1 Pure Python

First we had to extend the program `ljfluid.py` to set up the particles on a cubic lattice given the Number n of particles per side and their density ρ . The implementation of this is the following:

Listing 9: Placing Particles on a Cubic Lattice

```
# total number of particles
N = n*n*n

# particle positions on a cubic lattice
x = np.zeros((3,N))

# compute system size
L = n/density**(1/3)

# - set up n*n*n particles on a cubic lattice
positions = np.arange(0.5, n + 0.5)/density**(1/3)
for a in range(n):
    for b in range(n):
        for c in range(n):
            x[0,(a*n*n)+(b*n)+c] = positions[a]
            x[1,(a*n*n)+(b*n)+c] = positions[b]
            x[2,(a*n*n)+(b*n)+c] = positions[c]
```

3.1.1 Timings

The measured timings are given in the following table:

| n | time [s] |
|---|----------|
| 3 | 2.628 |
| 4 | 13.081 |
| 5 | 44.11 |

It can be seen that the time needed to do the simulations is basically proportional to n^3 which would be expected as we have three nested for loops over n in our computation of the positions on the cubic lattice and no other function has a higher complexity.

3.2 Pure C/C++

First we had to implement the function `compute_energy()` in C:

Listing 10: Computing the energy

```
double compute_energy(double *x, double *v) {
    double E_pot = 0;
    double E_kin = 0;
    double rij[3];

    // sum up potential energies
    for (int i=1; i < N; i++) {
        for (int j = 0; j < i; j++) {
            // distance vector
            minimum_image(x, i, j, rij);
            E_pot += compute_lj_potential(rij);
        }
    }

    // sum up kinetic energy
    for (int i=0; i < N; i++) {
        double dot = 0;
        dot += v[i]*v[i];
        dot += v[i+N]*v[i+N];
        dot += v[i+2*N]*v[i+2*N];
        E_kin += 0.5 * dot;
    }
    return E_pot + E_kin;
}
```

The total energy is put in the file *ljfluid.c.dat*. From the values in the file it can be seen that the energy fluctuates during the simulation time about 5%. This is supposedly because the system size L is quite small.

3.2.1 Timing

The measured timings are given in the following table:

| n | time [s] |
|----|----------|
| 5 | 0.148 |
| 6 | 0.342 |
| 7 | 0.584 |
| 8 | 1.052 |
| 9 | 1.892 |
| 10 | 3.11 |
| 11 | 5.11 |
| 12 | 8.121 |

Compared to the timings in Python measured for $n \leq 5$ the timings of the C code are much lower even for $n > 5$. The comparison of the timings for $n = 5$ shows the great difference between the run times of these two codes. As the Python code was running 44.11 s the C code only ran 0.148 s which is significantly lesser. Further more it can be observed that the time is not scaling with n^3 as in the Python code but seems to increase somehow quadratic. This can also be seen in the plot:

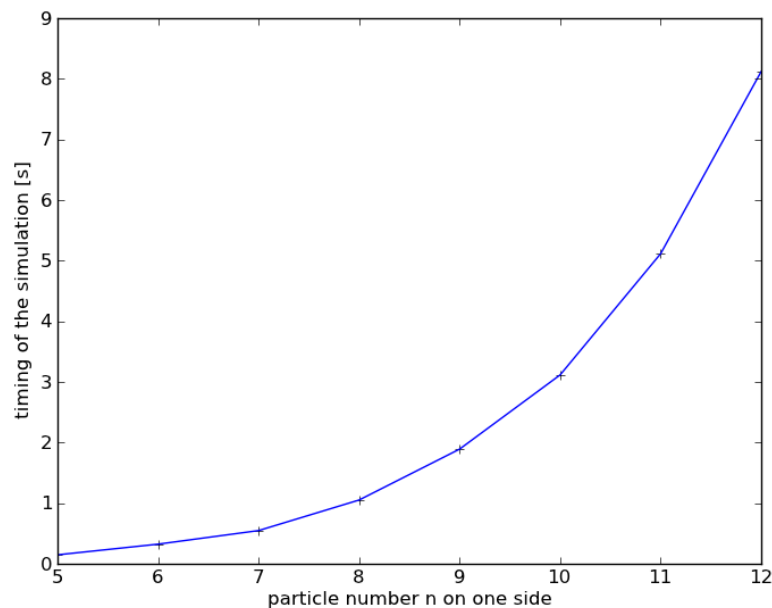


Figure 6: Timing of the simulation evolving with n

3.3 Mixing C/C++ and Python

In this section we mixed c and Python code. Our task was it to use the C code to measure the energy simulated with the code mixture. The resulting energy is:

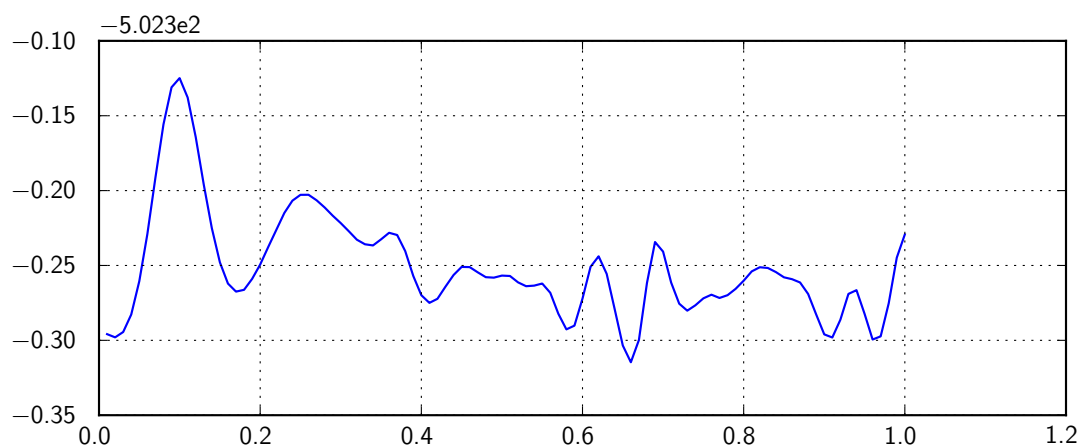


Figure 7: Energy of the system

Again it can be seen that the energy fluctuates a bit, but as the fluctuations are less than 1%, it's basically constant.

3.3.1 Timing

| n | time C [s] | tim Cython[s] |
|----|------------|---------------|
| 5 | 0.148 | 0.416 |
| 10 | 3.11 | 3.896 |

As it can be seen the timings for Cython are a bit greater than those for plain C, but Cython outperforms pure Python by far. As the variation in these two timings is quite small it can be said, that it is useful to use Cython as it's much more convenient than c and not significantly slower.

3.4 Neighborhood Lists

Using Cython we have up to now the problem that the scaling of the timings is still unfavorable. In order to improve this we introduced neighborhood lists. Which great effect these have on the simulation time can be seen in the following paragraph.

3.4.1 Timings

| n | time [s] |
|----|----------|
| 5 | 0.376 |
| 10 | 1.352 |
| 15 | 4.088 |
| 20 | 9.777 |

If you compare the timings of the simulation with and without neighborhood lists for $n = 5$ and $n = 10$ it becomes clear that introducing neighborhood lists reduced the timing approximately by a factor 4 respectively 3 which is quite a lot. Therefore it is indeed useful and worthwhile to introduce neighborhood lists even though this means some more implementation.