

# Temperature Measurements in Optical Tweezer Experiments

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# 1 Introduction

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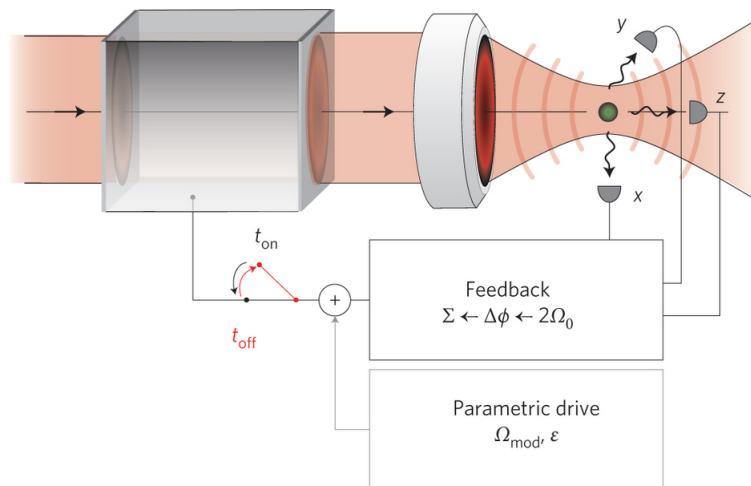
The starting point of this thesis is an experiment conducted by Gieseler et al [1]. It is an optical tweezer experiment, where the motion of a glass nanoparticle in a laser trap was used to investigate the fluctuation theorem[2].

### 3.1 Experimental setup

In the experiment, a silica nano particle with a radius of about 75 nm and mass of about  $3 \times 10^{-18}$ kg is trapped in a laser beam within a vacuum chamber. The trapping of the silica nano particle (which will be referred to as *glass particle*) is achieved by a gradient force of the laser beam acting on the particle. The experimental setup is depicted in fig. 1.

The particle fluctuates within the trap in all three spatial directions. These fluctuations can be approximated such that they are decoupled, which means that they can be described by a 1-dimensional Langevin equation:

$$\ddot{x} + \Gamma_0 \dot{x} + \Omega_0^2 x = \frac{1}{m} (F_{\text{fluct}} + F_{\text{ext}}) \quad (1)$$



**Figure 1:** Experimental setup of the optical tweezer experiment. A silica nano particle is trapped in a laser beam via gradient force in a vacuum. The feedback is used to cool down the particle and create a non-equilibrium steady state. In the first part of the experiment, the feedback is turned off and the motion of the particle towards an equilibrated state is observed. In the second part of the experiment, the steady state of the particle is modified by a parametric drive. Both the parametric drive and the feedback are turned off and – as in the first part – the motion of the particle towards an equilibrated state is observed.

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On the left hand side we have the friction coefficient  $\Gamma_0$  and the angular frequency that describes the fluctuation along the chosen axis. On the right hand side, there are two forces. The first one is  $F_{\text{fluct}}$ , which describes a stochastic force caused by interactions with the gas in the vacuum chamber. This force is given by

$$F_{\text{fluct}} = \sqrt{2m\Gamma_0 k_B T_0} \xi(t) \quad (2)$$

where  $T_0$  is the temperature of the heat bath (i.e. the surrounding gas in the vacuum chamber),  $k_B$  is the Boltzmann constant and  $\xi(t)$  is white noise, which obeys the equations  $\langle \xi(t) \rangle = 0$  and  $\langle \xi(t)\xi(t') \rangle = \delta(t - t')$ , which means that it is a random force. The term  $\Gamma_0$  appears in the formulat due to the fluctuation-dissipation theorem, which links the damping rate to the stochastic force.

The external force  $F_{\text{ext}}$  is part of the experimental setup, where the frequency of the fluctuation along an axis,  $\Omega_0$ , is measured and used to suppress the motion along said axis. This causes a decrease in the particle fluctuations and thus acts as a cooling mechanism for the particle in the trap. This process creates a non-equilibrium steady state  $\rho_{ss}(u, \alpha)$ , which is not known analytically. This state is the starting point of this thesis.

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## 4 Simulation

The problem at hand can be studied on an atomic level with the use of computer simulation. There is a variety of methods for computer simulations that are widely used, one of which being Molecular Dynamics (MD) simulations. The following section will give a brief overview of the concepts of this method, which is followed by the application to the simulation of the experiment.

### 4.1 Molecular Dynamics

Molecular Dynamics[3] simulations is a technique for simulating, as the name suggests, the dynamics of a classical many-body system. In this case, classical means, that the trajectories of the individual particles are calculated using classical mechanics rather than quantum mechanics. For relatively big atoms/molecules this is a very good approximation, whereas for systems consisting of hydrogen or helium the effects of quantum mechanics cannot be neglected and other methods (such as ab-initio simulation) has to be used.

The dynamics of the system are obtained by solving Newton's equations of motion for every particle.

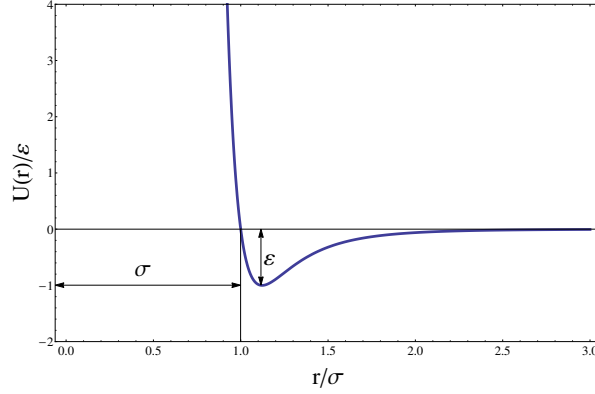
### 4.2 The Glass Particle

The glass particle from the experiment will be modeled as a system of particles interacting via a Lennard-Jones pair potential,

$$U(r) = 4\varepsilon \left[ \left( \frac{\sigma}{r} \right)^{12} - \left( \frac{\sigma}{r} \right)^6 \right] \quad (3)$$

where  $\varepsilon$  is the depth of the potential well (and thus its unit is energy) and  $\sigma$  is the distance at which the potential is zero. The form of the potential and the relation to the parameters is depicted in Fig. 2. Since  $\varepsilon$  and  $\sigma$  are crucial parameters for the simulation and do (usually) not change over time, it is practical to use them to define the dimensions of the system. This means that the unit of distance is  $\sigma$ , the unit of energy is  $\varepsilon$  and the unit of mass is the mass of the simulated particle. The so called *reduced units* can be constructed from these three parameters and put into relation to the original units. Here are some examples:

- distance:  $r^* = r/\sigma$
- potential energy:  $U^* = U/\varepsilon$
- temperature:  $T^* = k_B T/\varepsilon$



**Figure 2:** The Lennard-Jones 12-6 potential from (3). The x-axis is the particle distance divided by  $\sigma$  and the y-axis is the potential divided by the depth of the potential well.

- time:  $t^* = t\sqrt{\varepsilon/(m\sigma^2)}$
- pressure:  $P^* = P\sigma^3/\varepsilon$
- density:  $\rho^* = \rho\sigma^3$

One very popular choice for the simulated atoms is Argon because it is an inert gas and the atoms behave approximately like hard spheres which attract each other with weak van der Waals forces, which justifies the use of the Lennard-Jones potential. Argon has a mass of  $m = 6.69 \times 10^{-26}$  kg,  $\sigma = 3.4 \times 10^{-10}$  m and  $\varepsilon = 1.65 \times 10^{-21}$  J.

With the above introduced reduced units, the Lennard-Jones potential can be written as

$$U(r^*) = 4 [r^{*-12} - r^{*-6}]. \quad (4)$$

Since the reduced units will be used throughout the rest of this thesis, i will drop the asterisk henceforth.

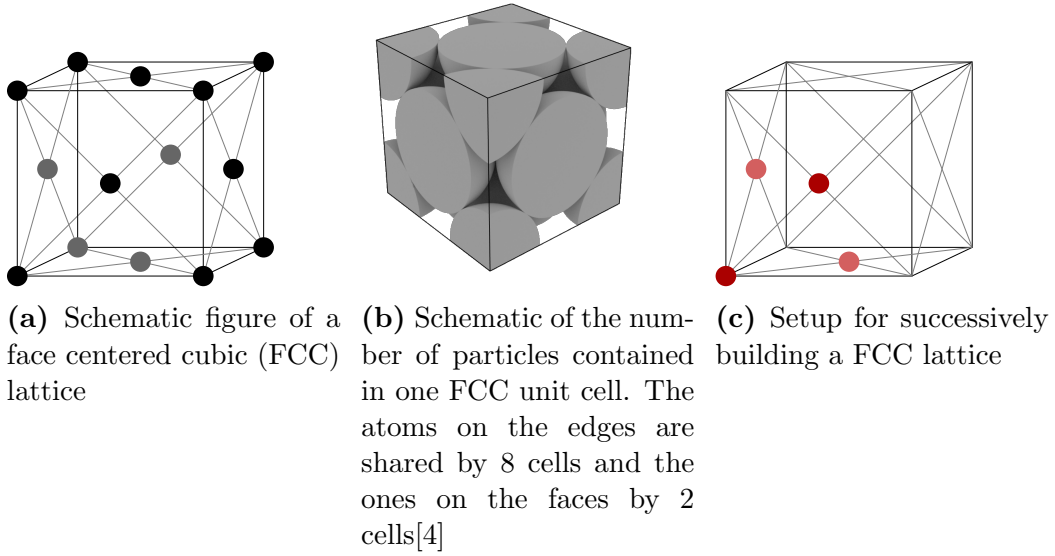
From the Lennard-Jones potential the corresponding force can be calculated by taking the derivative with respect to the direction of interest:

$$\begin{aligned} F_x &= -\frac{\partial}{\partial x} U(r) \\ &= -\frac{\partial}{\partial x} 4 [r^{-12} - r^{-6}] \\ &= -4 [(-12)r^{-13} - (-6)r^{-7}] \frac{\partial r}{\partial x} \\ &= 48 [r^{-13} - 0.5 r^{-7}] \frac{x}{r} \\ &= 48 [r^{-14} - 0.5 r^{-8}] x \end{aligned} \quad (5)$$

The force in the y and z direction can be calculated analogously.

The initial configuration of the particles is a face centered cubic (FCC) lattice. A schematic of the FCC lattice is depicted in fig. 3a. With the choice of FCC as initial configuration, there are optimal numbers for the numbers of the particles in the system. Since once FCC cell (as depicted) shares its atoms with its next neighbours, the number of atom per unit cell is 4, as depicted in fig. ???. The whole system of atoms is then created by repeating this cell structure. One convenient way is to arrange the unit cells in a cubic system, so if there are  $M$  FCC unit cells on one edge, the whole system consists of  $M^3$  cells. Since there are 4 particles per cell, there are ideal or so called *magic numbers* for atoms for which this setup works perfectly:  $N = 4M^3 = 4, 32, 108, 256, 500, 864, \dots$

There are several ways to achieve this initial configuration and the one used in



this thesis [5] was to create a kind of unit cell consisting of four atoms, as shown in fig. 3c, which can be described by a set of points

$$\begin{aligned} p_1 &= \{0, 0, 0\} \\ p_2 &= \{0.5, 0.5, 0\} \\ p_3 &= \{0.5, 0, 0.5\} \\ p_4 &= \{0, 0.5, 0.5\} \end{aligned}$$

From the particle number  $N$  and the number of FCC unit cells per edge  $M$  the lattice constant  $a$  can be calculated

$$a = \frac{L}{M} \tag{6}$$

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where  $L$  is the side length of the cube that is the whole system and it is calculated via the density of the system

$$L = \sqrt[3]{\frac{N}{\rho}} \quad (7)$$

With the lattice constant and the 4 points of the FCC cell, all the particles can be put into place.

### 4.3 The Velocity-Verlet Algorithm

When we look at the system from a thermodynamical standpoint, we see that it follows some kind of path in the phase space as time progresses. Every point in this space corresponds to a set of positions and momenta and the connection between two points corresponds to the evolution of the system from one state to another. As mentioned above, this evolution (the dynamics of the system) is a crucial element to Molecular Dynamics. Since the equations of motion cannot be solved analytically in general, we need to approximate the solution.

The method used here is called finite difference approach. The trajectory of the system in the phase space is cut into finite pieces of length  $\Delta t$  and the equations of motion are solved for every segment separately (see fig. 4).

There are several ways to solve this kind of problem, but since we are interested in implementing it into a computer program the ideal solution should have some basic properties [6]:

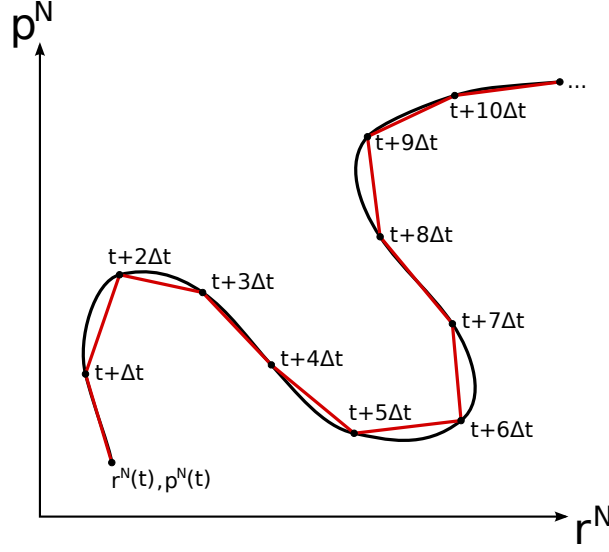
- It should be fast and require little memory
- It should permit the use of a large time step  $\Delta t$
- It should duplicate the classical trajectory as closely as possible
- It should satisfy known conservation laws
- It should be simple and easy to program

One algorithm that has all of the above mentioned features is the one proposed by Verlet [7]. In his paper, Verlet starts by Taylor expanding the coordinate  $\mathbf{r}_i$  for one particle after one time step  $\Delta t$ :

$$\mathbf{r}_i(t + \Delta t) = \mathbf{r}_i(t) + \dot{\mathbf{r}}_i \Delta t + \frac{1}{2} \ddot{\mathbf{r}}_i \Delta t^2 + \frac{1}{3!} \ddot{\mathbf{r}}_i \Delta t^3 + \mathcal{O}(\Delta t^4) \quad (8)$$

Since the first derivative of the coordinate  $\mathbf{r}_i$  is the velocity,  $\dot{\mathbf{r}}_i(t) = \mathbf{v}_i(t)$ , and the second derivative of the coordinate  $\mathbf{r}_i$  is the acceleration,  $\ddot{\mathbf{r}}_i(t) =$





**Figure 4:** Simplified graphical schematic of the finite difference approach. The evolution of the system from a point  $(r^N(t), p^N(t))$  in the phase space is approximated by slicing it up into pieces of length  $\Delta t$ . On every stop after the starting point ( $t + \Delta t, t + 2\Delta t, t + 3\Delta t, \dots$ ) the equations of motion can be solved numerically.

$\mathbf{a}_i(t)$ , using Newton's law  $\mathbf{F}_i(t) = m_i \mathbf{a}_i(t)$  equation (8) can be written as:

$$\mathbf{r}_i(t + \Delta t) = \mathbf{r}_i(t) + \mathbf{v}_i(t)\Delta t + \frac{1}{2m}\mathbf{F}_i(t)\Delta t^2 + \frac{1}{3!}\ddot{\mathbf{r}}_i(t)\Delta t^3 + \mathcal{O}(\Delta t^4) \quad (9)$$

The same calculation can be carried out for one time step before  $t$ :

$$\mathbf{r}_i(t - \Delta t) = \mathbf{r}_i(t) - \mathbf{v}_i(t)\Delta t + \frac{1}{2m}\mathbf{F}_i(t)\Delta t^2 - \frac{1}{3!}\ddot{\mathbf{r}}_i(t)\Delta t^3 + \mathcal{O}(\Delta t^4) \quad (10)$$

The sum of those two equations yields

$$\mathbf{r}_i(t + \Delta t) + \mathbf{r}_i(t - \Delta t) = 2\mathbf{r}_i(t) + \frac{1}{2}\mathbf{F}_i(t)\Delta t^2 + \mathcal{O}(\Delta t^4) \quad (11)$$

and from this we get the final form for the new coordinates:

$$\mathbf{r}_i(t + \Delta t) = 2\mathbf{r}_i(t) + \mathbf{r}_i(t - \Delta t) + \frac{1}{2}\mathbf{F}_i(t)\Delta t^2 \quad (12)$$

This means that the new coordinates can be calculated using the current coordinates, the current forces and the coordinates from the past time step.

From equation (12) one thing becomes clear: the velocities are not necessary to calculate the new positions and they are not computed in the process, but for

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the calculation of i.e. the kinetic energy the velocities are needed. Thus, the velocities have to be calculated with a combination of the Taylor expansions of the coordinate mathbf{r} at  $t + \Delta t$  and  $t - \Delta t$ :

$$\mathbf{r}_i(t + \Delta t) - \mathbf{r}_i(t - \Delta t) = 2\mathbf{v}_i(t)\Delta t + \mathcal{O}(\Delta t^3) \quad (13)$$

which can be rewritten as

$$\mathbf{v}_i(t) = \frac{\mathbf{r}_i(t + \Delta t) - \mathbf{r}_i(t - \Delta t)}{2\Delta t} + \mathcal{O}(\Delta t^2) \quad (14)$$

This approach has the advantage that it is fast, requires little memory and is reliable in the sense that there is no energy drift occuring during the simulation, which means that the energy is conserved. When we compare this to the list of desired properties this seems like a good algorithm.

This algorithm however has two significant disadvantages. The first one is the calculation of the velocities. As can be seen in equation (14), the accuracy of the calculation is only  $\mathcal{O}(\Delta t^2)$  while the positions can be calculated with an error of order  $\Delta t^4$ . The other big disadvantage is the first step of the algorithm. Since the calculation of the new positions requires the current positions and the ones from one time step before, which technically do not exist.

The solutions to this problem is to include the stepwise calculation of the velocities [8]. For this we start again with the Taylor expansion of the coordinates mathbf{r} (9) and instead of Taylor expanding  $\mathbf{r}_i(t - \Delta t)$ , we write it as

$$\mathbf{r}_i(t) = \mathbf{r}_i(t + \Delta t) - \mathbf{v}_i(t + \Delta t)\Delta t + \frac{1}{2m}\mathbf{F}_i(t + \Delta t)\Delta t^2 \quad (15)$$

Using (9) in the above equation we get

$$\begin{aligned} \mathbf{r}_i(t) &= \mathbf{r}_i(t) + \mathbf{v}_i(t)\Delta t + \frac{1}{2m}\mathbf{F}_i(t)\Delta t^2 \\ &\quad - \mathbf{v}_i(t + \Delta t)\Delta t + \frac{1}{2m}\mathbf{F}_i(t + \Delta t)\Delta t^2 \end{aligned} \quad (16)$$

and thus

$$\mathbf{v}_i(t + \Delta t) = \mathbf{v}_i(t) + \frac{1}{2m}(\mathbf{F}_i(t) + \mathbf{F}_i(t + \Delta t))\Delta t \quad (17)$$

With the addition of this equation this algorithm is called the Velocity-Verlet algorithm, which is summarized in the following two equations:

$$\begin{aligned} \mathbf{r}_i(t + \Delta t) &= \mathbf{r}_i(t) + \mathbf{v}_i(t)\Delta t + \frac{1}{2m}\mathbf{F}_i(t)\Delta t^2 \\ \mathbf{v}_i(t + \Delta t) &= \mathbf{v}_i(t) + \frac{1}{2m}(\mathbf{F}_i(t) + \mathbf{F}_i(t + \Delta t))\Delta t \end{aligned} \quad (18)$$

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This algorithm is self starting, uses a small amount of memory, conserves the energy and gives a very good approximation to the original trajectory in the phase space. To program this algorithm the following steps are needed:

1. Calculate all the forces between the particles (for the first run only)
2. Calculate the new positions with current velocity and forces
3. Calculate first half of the new velocities with current forces
4. Calculate all the forces for the new position
5. Use new forces to calculate second half of new velocities

The first point only has to be carried out for the first run, because the forces have not been calculated at this point. As the forces are calculated for the new positions in step 4, they can be used for the next time step. For the first time step the velocities have to be chosen randomly and are calculated with this algorithm from that point forward.

#### **4.4 The Laser Beam - Energy influx**

The glass sphere is trapped in the laser beam. While the motion of the center of mass is suppressed, which acts as a cooling mechanism, the individual atoms that make up the glass sphere absorb the energy from the laser which increases their velocity.

To simulate this kind behaviour, thermostat algorithms [9] such as the Nosé-Hoover [10, 11] or the Berendsen [12] algorithms.

The problem with these kind of algorithms is, that a target temperature has to be fixed which will be reached at some point. In order to simulate the influx of energy from the laser it would be better to have an algorithm that supplies the system with a certain amount of energy continuously. Fortunately, such an algorithm exists.

The algorithm is called Heat Exchange Algorithm (HEX) [13]. Its intended purpose is the use in non-equilibrium molecular dynamics (NEMD) to study transport phenomena and determine transport coefficients. The algorithm works by introducing two regions in the system, one serving as a heat source and the other as a heat sink. A specific amount of heat is then exchanged between those two reservoirs. As it turns out however, this algorithm introduces an energy shift for longer simulation times. This led Wirnsberger et al [14] to revisit the algorithm and identify the cause of this energy drift, to create a more suitable algorithm, which they called eHEX.

As in the HEX algorithm, regions are introduced to the system which act either

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as heat sinks or heat sources. These are labelled with  $\Gamma_k$ , where  $k > 0$ , and have corresponding amount of exchanged heat  $\Delta Q_{\Gamma_k}$ . If  $\Delta Q_{\Gamma_k}$  is negative, heat is subtracted from the system and vice versa. Regions that neither act as heat source or heat sink are labelled with  $\Gamma_0$ , which are also called Hamiltonian regions (see fig. 5). The simulation box, denoted by  $\Omega$ , and the regions  $\Gamma_k$  are assumed to be moving with velocities  $v_\Omega$  and  $v_{\Gamma_k}$  respectively.

The change of the energy in a region  $\Gamma_k$  is achieved by rescaling the velocities by a factor  $\xi_k$  and shifted by the velocity of the corresponding region:

$$\mathbf{v}_i \rightarrow \bar{\mathbf{v}}_i = \xi_k \mathbf{v}_i + (1 - \xi_k) \mathbf{v}_{\Gamma_k} \quad (19)$$

The bar over a quantity denotes the value after the exchange of heat.

The factor  $\xi_k$  is given by

$$\xi_k = \sqrt{1 + \frac{\Delta Q_{\Gamma_k}}{\mathcal{K}_{\Gamma_k}}} \quad (20)$$

where  $\Delta Q_{\Gamma_k}$  is the exchanged heat in the region  $\Gamma_k$  and  $\mathcal{K}_{\Gamma_k}$  is the non-translational kinetic energy of the region  $\Gamma_k$  and is given by

$$\mathcal{K}_{\Gamma_k} = \sum_{i \in \gamma_k} \frac{m_i v_i^2}{2} - \frac{m_{\Gamma_k} v_{\Gamma_k}^2}{2} \quad (21)$$

The sum is taken over all indices in  $\gamma_k$  which is the set of indices of particles in the region  $\Gamma_k$ .

For the final version of the eHEX algorithm there are three more quantities needed. The first one is the heat flux per time step, denoted by  $\mathcal{F}_{\Gamma_k}$ :

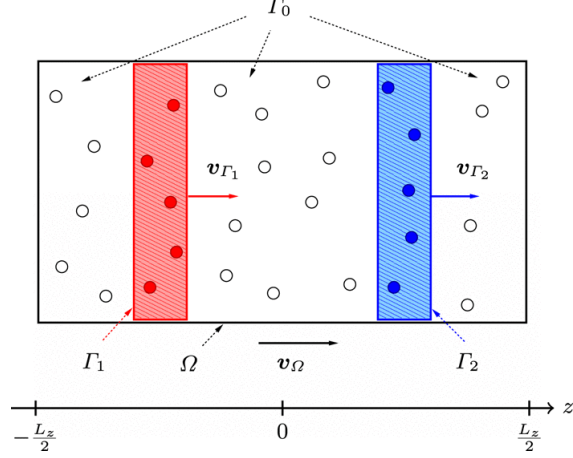
$$\mathcal{F}_{\Gamma_k} = \frac{\Delta Q_{\Gamma_k}}{\Delta t} \quad (22)$$

The second one is the thermostating force  $\boldsymbol{\eta}_i$ , which is defined as

$$\boldsymbol{\eta}_i = \begin{cases} m_i \frac{\mathcal{F}_{\Gamma_{k(\mathbf{r}_i)}}}{2\mathcal{K}_{\Gamma_{k(\mathbf{r}_i)}}} \left( \mathbf{v}_i - \mathbf{v}_{\Gamma_{k(\mathbf{r}_i)}} \right) & \text{if } k(\mathbf{r}_i) > 0 \\ 0 & \text{otherwise} \end{cases} \quad (23)$$

where  $k(\mathbf{r}_i)$  is the index of the region in which particle  $i$  located, i.e.  $k(\mathbf{r}_i) = 0$  means that the particle is in a Hamiltonian region and  $k(\mathbf{r}_i) > 0$  denotes the heat sinks and sources.

The last quantity is the one that corrects the long term energy drift of the HEX algorithm, denoted by  $\mathcal{E}r_{i,\alpha}$ . The analysis and derivation of this term is given in



**Figure 5:** Setup for the use of the HEX and eHEX algorithms. The simulation box  $\Omega$ , which is moving with a velocity of  $v_\Omega$ , contains a heat source (red),  $\Gamma_1$ , which is moving with velocity  $v_{\Gamma_1}$  and a heat sink (blue),  $\Gamma_2$ , which is moving with velocity  $v_{\Gamma_2}$ . The regions that are neither heat sources or heat sinks are denoted by  $\Gamma_0$ .

[14].

$$\begin{aligned} \mathcal{E}r_{i,\alpha} = & \frac{\eta_{i,\alpha}}{m_i \mathcal{K}_{\Gamma_k(\mathbf{r}_i)}} \left[ \frac{\mathcal{F}_{\Gamma_k(\mathbf{r}_i)}}{48} + \frac{1}{6} \sum_{j \in \gamma_k(\mathbf{r}_i)} \mathbf{f}_j \cdot (\mathbf{v}_j - \mathbf{v}_{\Gamma_k(\mathbf{r}_i)}) \right] \\ & - \frac{\mathcal{F}_{\Gamma_k(\mathbf{r}_i)}}{12 \mathcal{K}_{\Gamma_k(\mathbf{r}_i)}} \left[ \frac{f_{i,\alpha}}{m_i} - \frac{1}{m_{\Gamma_k(\mathbf{r}_i)}} \sum_{j \in \gamma_k(\mathbf{r}_i)} f_{j,\alpha} \right] \end{aligned} \quad (24)$$

The  $\mathbf{f}$  in the above equation denotes the force corresponding to the chosen inter-molecular potential  $U(\mathbf{r})$ ,

$$\mathbf{f} = -\nabla_{\mathbf{r}_i} U(\mathbf{r}_i) \quad (25)$$

With all the necessary quantities established, the updating sequence of the eHEX algorithm can be written down:

$$\bar{\mathbf{v}}_i^n = \xi_{k(\mathbf{r}_i)}^n \mathbf{v}_i^n + (1 - \xi_{k(\mathbf{r}_i)}^n) \mathbf{v}_{\Gamma_k(\mathbf{r}_i)}^n \quad (26a)$$

$$\bar{\mathbf{v}}_i^{n+\frac{1}{2}} = \bar{\mathbf{v}}_i^n + \frac{\Delta t}{2m_i} \mathbf{f}_i^n \quad (26b)$$

$$\bar{\mathbf{r}}_i^{n+1} = \mathbf{r}_i^n + \Delta t \bar{\mathbf{v}}_i^{n+\frac{1}{2}} \quad (26c)$$

$$\mathbf{f}_i^{n+1} = -\nabla_{\mathbf{r}_i} U(\mathbf{r})|_{\mathbf{r}=\bar{\mathbf{r}}_i^{n+1}} \quad (26d)$$

$$\bar{\mathbf{v}}_i^{n+1} = \bar{\mathbf{v}}_i^{n+\frac{1}{2}} + \frac{\Delta t}{2m_i} \mathbf{f}_i^{n+1} \quad (26e)$$

$$\mathbf{v}_i^{n+1} = \bar{\xi}_{k(\bar{\mathbf{r}}_i)}^{n+1} \bar{\mathbf{v}}_i^{n+1} + (1 - \bar{\xi}_{k(\bar{\mathbf{r}}_i)}^{n+1}) \bar{\mathbf{v}}_{\Gamma_k(\bar{\mathbf{r}}_i)}^{n+1} \quad (26f)$$

$$\mathbf{r}_i^{n+1} = \bar{\mathbf{r}}_i^{n+1} - \Delta t^3 \mathcal{E} \bar{\mathbf{r}}_i^{n+1} \quad (26g)$$

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This algorithm can be applied to the problem of the levitating nano sphere in the laser beam by adjusting the setup and some of the parameters.

Firstly, the setup of the heat sinks and sources has to be changed. Since there is only energy pumped into the system from the outside, the region acting as a heat sink vanishes and the region acting as a heat source spans over the whole simulation box. This means that, using the notation of fig. 5,  $\Omega = \Gamma_1$ . Furthermore, neither the simulation box nor the heat source are moving, i.e.  $\mathbf{v}_\Omega = \mathbf{v}_{\Gamma_1} = 0$ . This affects the non-translational kinetic energy term  $\mathcal{K}_{\Gamma_1}$ , the thermostating force  $\boldsymbol{\eta}$  and the correction term  $\mathcal{E}\mathbf{r}$ . Since there is only one region acting as a heat sink, the terms  $\Delta Q$ ,  $\mathcal{K}$  and  $\mathcal{F}$  don't need an index. The summation index in (21) and (24) can be changed to the number of particles in the system,  $N$ , since all the particles are within the heat exchanging region. The masses are set to 1, i.e.  $m_i = 1$  with the chosen reduced units and with this the total mass of the region (in the term  $1/m_{\Gamma_k(\mathbf{r}_i)}$  in (24)) is equal to the number of particles in the system. With these changes, the quantities can be written down as:

$$\mathcal{K} = \sum_N \frac{v_i^2}{2} \quad (27)$$

$$\xi = \sqrt{1 + \frac{\Delta Q}{\mathcal{K}}} \quad (28)$$

$$\mathcal{F} = \frac{\Delta Q}{\Delta t} \quad (29)$$

$$\boldsymbol{\eta}_i = \frac{\mathcal{F}}{2\mathcal{K}} \mathbf{v}_i \quad (30)$$

$$\begin{aligned} \mathcal{E}r_{i,\alpha} &= \frac{\eta_{i,\alpha}}{\mathcal{K}} \left[ \frac{\mathcal{F}}{48} + \frac{1}{6} \sum_N \mathbf{f}_j \cdot \mathbf{v}_j \right] \\ &- \frac{\mathcal{F}}{12\mathcal{K}} \left[ f_{i,\alpha} - \frac{1}{N} \sum_N f_{j,\alpha} \right] \end{aligned} \quad (31)$$

This means that the laser is modeled to be pumping energy into the system, which increases the velocities of the particles in the system over time, while the center of mass motion is not affected by this.

## 4.5 The Laser Beam - Trapping

As mentioned above, the glass particle is trapped in the laser beam and the motion of the particle is suppressed in every direction. This behaviour has to be modeled as well.

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The approximation of the original paper [1], where the movements in the three spatial directions are decoupled will be used in the model as well. As to the model of the trapping force itself, the most straightforward approach is to use a harmonic oscillator potential. The force and the corresponding potential can be written as

$$\mathbf{F} = -kx \quad (32)$$

$$U = \frac{1}{2}kx^2 \quad (33)$$

The trap is implemented by using the velocity Verlet algorithm with the above force to propagate the glass particle. Since the trap is acting on the whole system, the force is acting on the center of mass position and velocity. To calculate the center of mass positions and velocities, the positions and velocities of all particles have to be summed up and divided by the particle number:

$$\mathbf{r}_{\text{COM}} = \sum_{i=1}^N \frac{\mathbf{r}_i}{N} \quad (34)$$

$$\mathbf{v}_{\text{COM}} = \sum_{i=1}^N \frac{\mathbf{v}_i}{N} \quad (35)$$

The position of the center of mass and the center of mass velocity are then propagated by using the same procedure as introduced in section 4.3.

## 4.6 Surrounding Gas - Barostat

Without any equilibrating mechanism the setup described by now would lead to the system melting, which is not desirable. In the next step we will introduce the surrounding gas of the gas chamber that will absorb some of the energy in the system, leading to the final state.

Generally, pressure is introduced to the system by surrounding the object of interest (in this case the glass nano sphere) with a pressure medium. There are two main requirements for the choice of such a pressure medium: the exerted pressure must be hydrostatic and the computation of the interaction between the pressure medium and the object of interest must not take up a lot of resources.

The model used in this thesis was developed by Grünwald and Dellago [15] and uses an ideal gas of non-interacting particles as pressure medium. The particles of this pressure medium flow into the simulation from an outside volume, whose geometry is based on the form of the object of interest (this will be referred to as the minimal volume of cells). This barostat also acts as a thermostat and thus is suitable for the application in this thesis.

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The gas particles interact with the object via a soft-sphere potential of the form

$$U(r) = \varepsilon \left( \frac{\sigma}{r} \right)^{12} \quad (36)$$

where  $\varepsilon$  is the interaction strength and  $\sigma$  is the interaction range and  $r$  is the distance between the gas particle and the interacting particle. The gas particles do not interact with one another in this model.

In order to increase the efficiency of the computing process,  $\sigma$  should be chosen carefully. For larger  $\sigma$ , the number of interaction partners increases, which increases the number of force calculations which are a very time consuming part. For smaller  $\sigma$  the possibility for gas particles reaching the inside of the nano crystal increases, which is not desirable. So  $\sigma$  should be chosen small enough to keep the force calculations at a minimum and large enough for the particle to stay on the outside of the crystal.

The algorithm can be performed by following these steps:

1. Randomly draw the number of particles that are created on a single side of the minimal volume of cells,  $N_{\text{fac}}$ , from the distribution

$$\langle N_{\text{fac}} \rangle = \Delta t L^2 P \sqrt{\frac{1}{2\pi m k_B T}} \quad (37)$$

where  $\Delta t$  is the time step of the simulation,  $L$  is the side length of the cell in which the particle is created,  $P$  is the desired pressure,  $m$  is the mass of the gas particle,  $k_B$  is the Boltzmann constant (which will be set to 1 in reduced units) and  $T$  is the desired temperature. This chosen number of particles is then equally distributed over the face of the cell on which they are created. The velocities of the created particles are drawn from two different random number distributions. The component of the particle perpendicular to the surface is drawn from a Rayleigh distribution of the form

$$p(v_i) = \frac{m}{k_B T} v_i e^{-\frac{m v_i^2}{2 k_B T}} \quad (38)$$

The other components of the particles' velocity are drawn from a Maxwell-Boltzmann distribution.

2. Perform the first step of the velocity Verlet algorithm to propagate the particle positions by one time step.
3. Check if any gas particles have left the minimal volume of cells and remove those which have.



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4. Check if the geometry of the crystal and with it the minimal volume of cells has changed. If it has, remove all gas particles in the cells that are no longer needed. Then insert new gas particles to the created cells with a number drawn from a Poisson distribution with mean value

$$\langle N_{\text{ins}} \rangle = \frac{PL^3}{k_B T} \quad (39)$$

If the length of the cell  $L$  is chosen to be equal to the cut-off radius  $r_c$ , this insertion should only be carried out with a probability of

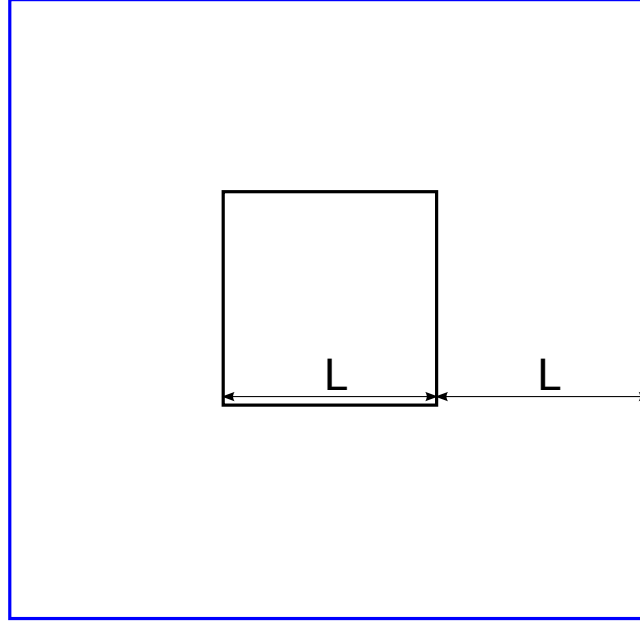
$$P_{\text{ins}} = e^{-\frac{U}{k_B T}} \quad (40)$$

(where  $U$  is the interaction energy between the gas particle and the crystal) because it is possible that the inserted particle is within the interaction range of the crystal.

5. Compute all forces.
6. Perform the second step of the velocity Verlet and propagate the particle velocities by one time step.

This algorithm is formulated in a general fashion, so that a wide range of crystals and geometries can be used. The setup used in this thesis is a FCC lattice, which has a nice symmetry that is not influenced by the surrounding gas. The algorithm will therefore not be carried out using cell-lists and a minimal volume of cells that is changing over the course of the simulation, but rather a fixed setup of the volume surrounding the nano crystal.

Since the crystal itself is a cube a straightforward approach is surrounding it by a bigger cube. The distance between the cube face and the nano crystal is chosen to be the side length of the nano crystal, so the setup will properly scale for different numbers of particles. The schematic setup for this is depicted in fig.6.



**Figure 6:** Schematic setup for the barostat. The outer volume (blue) is one side length  $L$  away from the nano crystal (black). The gas particles are created along the blue lines of the outer volume. Since the geometry of the nano crystal does not change, the geometry of the outer volume will be constant throughout the simulation.

## 5 Results

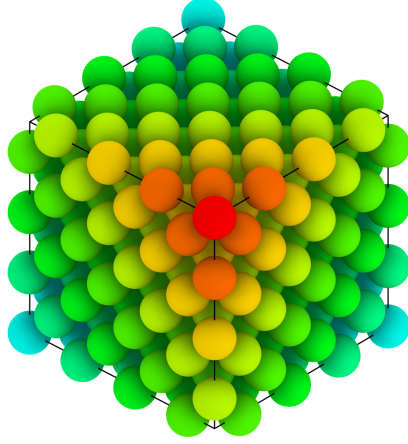
Since the simulation consists of various parts that have to work together, we first need to check if every part itself works.

### 5.1 The Crystal

The first element of the simulation is the setup of the FCC lattice on which the particles are placed on. To check if this is done correctly, we can use a visualisation and rendering software such as VMD[16] or OVITO[17]. The rendered image of the nano crystal can be seen in fig. 7. The chosen perspective shows the structure of the FCC lattice, where 3 layers are forming where the atoms take the position on the gap between the atoms from the layer below.

### 5.2 Velocity Verlet

The simulation is taking place in a NVE ensemble from a thermodynamic point of view. This means that the particle number  $N$ , the volume  $V$  and the total energy



**Figure 7:** A rendered image of the FCC crystal. The chosen color-coding along the (1,1,1) Miller-index highlights the layers for this perspective. The perspective is chosen to see the layering of the FCC crystal with its 3 layer structure. The black box acts as a reference point to outline the geometry of the system.

$E$  are constant. In contrast to the NVT ensemble used in Monte-Carlo simulations this means, that the velocities of the particle can not be chosen directly. The initial velocities have to be drawn from a random number generator. This means that the velocities have to be adjusted after they have been drawn.

The first adjustment that needs to be made is making sure that the system is staying in place and not floating away. To do this, the velocity vectors of all the atoms in the system have to be added to get the resulting velocity vector for the whole system that is thought to be acting on the center of mass. This resulting vector is then divided by the number of particles in the system and the resulting vector is then subtracted from every velocity vector of every atom in the system.

$$\mathbf{v}_{\text{tot}} = \sum_{i=1}^N \mathbf{v}_i \quad (41)$$

$$\mathbf{v}_{\text{cut}} = \frac{\mathbf{v}_{\text{tot}}}{N} \quad (42)$$

$$\mathbf{v}_i^{\text{new}} = \mathbf{v}_i - \mathbf{v}_{\text{cut}} \quad (43)$$

The next adjustment is a small one that makes a difference in the long run. When choosing random velocities, the system not only tends to have a resulting velocity vector that causes the whole system to float away, but also a resulting angular momentum that causes the system to rotate. The angular momentum can be

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defined via the moment of inertia  $\mathbf{I}$

$$\mathbf{L} = \mathbf{I}\boldsymbol{\omega} \quad (44)$$

$$\boldsymbol{\omega} = \frac{\mathbf{r} \times \mathbf{v}}{r^2} \quad (45)$$

where  $\boldsymbol{\omega}$  is the angular velocity of the system.

The velocities of the particle can roughly be chosen by adjusting the range of numbers the random number generator can choose from. However, the resulting velocities (and therefore the temperature of the system) are generally not precisely how they should be. If the system should have a specific starting temperature  $T$ , the velocities need to be scaled accordingly. This rescaling is done by summing the square of the velocities (this equals calculating the kinetic energy since the masses are set to 1) and rescale them by a factor  $\lambda$ :

$$E = \sum_{i=1}^N \mathbf{v}_i^2 \quad (46)$$

$$\lambda = \sqrt{\frac{3(N-1)T}{E}} \quad (47)$$

$$\mathbf{v}_i^{\text{new}} = \lambda \mathbf{v}_i \quad (48)$$

This process has to be done a couple of times, until the system equilibrates to the desired temperature. The problem with this procedure is the fact that it contradicts the constant energy property of the ensemble. If we look at the energy values in fig. ?? we can see the irregularities before the system equilibrates to a constant value.

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## 6 Conclusion

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## References

- [1] Jan Gieseler, Romain Quidant, Christoph Dellago, and Lukas Novotny. Dynamic relaxation of a levitated nanoparticle from a non-equilibrium steady state. *Nat Nano*, 9(5):358–364, May 2014.
- [2] Gavin E. Crooks. Entropy production fluctuation theorem and the nonequilibrium work relation for free energy differences. *Phys. Rev. E*, 60:2721–2726, Sep 1999.
- [3] D. Frenkel and B. Smit. *Understanding Molecular Simulation: From Algorithms to Applications*. Computational science series. Elsevier Science, 2001.
- [4] Image taken from [https://upload.wikimedia.org/wikipedia/commons/2/27/Elementarzelle\\_einer\\_kubisch\\_raumzentrierten\\_Elementarzelle.png](https://upload.wikimedia.org/wikipedia/commons/2/27/Elementarzelle_einer_kubisch_raumzentrierten_Elementarzelle.png).
- [5] Idea taken from the Molecular Dynamics part of this lecture: <http://www.physics.buffalo.edu/phy411-506/>.
- [6] M. P. Allen and D. J. Tildesley. *Computer Simulation of Liquids*. Clarendon Press, New York, NY, USA, 1989.
- [7] Loup Verlet. Computer "experiments" on classical fluids. i. thermodynamical properties of lennard-jones molecules. *Phys. Rev.*, 159:98–103, Jul 1967.
- [8] William C. Swope, Hans C. Andersen, Peter H. Berens, and Kent R. Wilson. A computer simulation method for the calculation of equilibrium constants for the formation of physical clusters of molecules: Application to small water clusters. *The Journal of Chemical Physics*, 76(1):637–649, 1982.
- [9] Philippe H. Hünenberger. *Thermostat Algorithms for Molecular Dynamics Simulations*, pages 105–149. Springer Berlin Heidelberg, Berlin, Heidelberg, 2005.
- [10] Shuichi Nosé. A unified formulation of the constant temperature molecular dynamics methods. *The Journal of Chemical Physics*, 81(1):511–519, 1984.
- [11] William G. Hoover. Canonical dynamics: Equilibrium phase-space distributions. *Phys. Rev. A*, 31:1695–1697, Mar 1985.
- [12] H. J. C. Berendsen, J. P. M. Postma, W. F. van Gunsteren, A. DiNola, and J. R. Haak. Molecular dynamics with coupling to an external bath. *The Journal of Chemical Physics*, 81(8):3684–3690, 1984.

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- [13] B. Hafskjold, T. Ikeshoji, and S. Kjelstrup Ratkje. On the molecular mechanism of thermal diffusion in liquids. *Molecular Physics*, 80:1389–1412, December 1993.
  - [14] P. Wirnsberger, D. Frenkel, and C. Dellago. An enhanced version of the heat exchange algorithm with excellent energy conservation properties. *The Journal of Chemical Physics*, 143(12), 2015.
  - [15] M. Grünwald and C. Dellago. Ideal gas pressure bath: a method for applying hydrostatic pressure in the computer simulation of nanoparticles. *Molecular Physics*, 104(22-24):3709–3715, 2006.
  - [16] William Humphrey, Andrew Dalke, and Klaus Schulten. VMD – Visual Molecular Dynamics. *Journal of Molecular Graphics*, 14:33–38, 1996.
  - [17] Alexander Stukowski. Visualization and analysis of atomistic simulation data with ovito—the open visualization tool. *Modelling and Simulation in Materials Science and Engineering*, 18(1):015012, 2010.