# HW9 Write-up

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An algorithm produced a simulation of the trajectories of 10 particles subject to the Lennard-Jones potential.

#### 1. INTRODUCTION

In this assignment, we simulated the motion of 10 identical particles subject only to the Lennard-Jones force exerted by neighboring particles. The Lennard-Jones force exerted on a particle by another particle is given by

$$\vec{F} = 24 \frac{\epsilon}{r} \left[ 2 \left( \frac{\sigma}{r} \right)^{12} - \left( \frac{\sigma}{r} \right)^{6} \right] \frac{\vec{r}}{r} \tag{1}$$

where  $\epsilon$  is the potential energy at the equilibrium bond length,  $\sigma$  is the distance at which the potential energy is zero,  $\vec{r}$  is the displacement vector between particles, and r is the distance between particles. The value used for  $\epsilon$  was 0.0103eV and the value used for  $\sigma$  was 3.4Å. The distance between two particles was calculated using the equation

$$r = \sqrt{r_x^2 + r_y^2} = \sqrt{(x_j - x_i)^2 + (y_j - y_i)^2}$$
 (2)

where i and j denote the particles and x and y denote the components of the position vectors of the particles. Using the Lennard-Jones potential allows for the simulation of collisions because the potential increases drastically when particles get too close to each other, producing a repulsive force.[1] However, it also accounts for the attractive effects of Van der Waals interactions of induced dipole moments of particles.[1] These characteristics make studying the effects of systems subject only to a Lennard-Jones potential interesting.

## 2. EXPERIMENT

As particles move in a system, the Lennard-Jones force exerted on each particle by all the other particles in the system changes, influencing the evolution of the system in time. Two-dimensional kinematics equations were used to calculate the trajectories of the particles in two dimensions. Figure 1 shows the plan for adding motion in a second dimension to molecular dynamics simulation.

In this experiment, 10 particles were created. They were given identical initial x- and y- coordinates of that

FIG. 1: Pseudocode to convert the algorithm for a molecular dynamics simulation in one-dimension into a simulation in two-dimensions.

were evenly spaced. Initial velocities were assigned according to

$$v = n\sqrt{\frac{k_B T}{m}} \tag{3}$$

with Boltzmann constant  $k_B$ , temperature of the system T and mass m. **Rand**, a random number generator in the **Numpy** package, produced an array of 10 random numbers n that were scaled according to Equation 3 to produce a system at the desired temperature of 300K. The mass of argon was used as the mass m of the simulation particles. This process was done twice to produce arrays with the initial component of the velocities in the x- and y-dimension.

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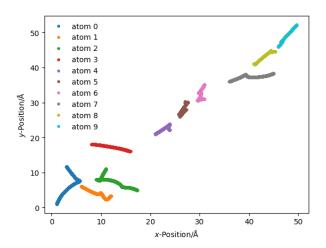


FIG. 2: Trajectories of simulation particles in two dimensions at 300 K.

The accelerations of the particle were found using the Lennard-Jones force (Eq. 1 on each particle with Equation 6 at each time-step, which are then used to update the velocity and position vectors:

$$\vec{v} = \vec{v_0} + (\vec{a_0} + \vec{a}) dt \tag{4}$$

$$\vec{x} = \vec{x_0} + \vec{v_0}dt + 0.5\vec{a_0}dt^2 \tag{5}$$

 $\vec{a_0}$  is the acceleration of the previous time-step and  $\vec{a}$  is

the acceleration of the current time-step.

$$\vec{a} = \frac{\vec{F}}{m} \tag{6}$$

The simulation ran for 10000 time-steps of 0.1 seconds, and the positions of the particles were saved every 100 time-steps to generate the frames of the video simulation. Visual Molecular Dynamics (VMD)[2] was used to generate an animation of the particles trajectories in time. In this video, collisions of the particles could observed as the two-dimensional positions of particles naturally evolved in time.

### 3. RESULTS

The resulting simulation produced trajectories (Fig. 2) that did not overlap. This is important because it means that the Lennard-Jones forces did not allow particles to occupy the same space, which would be an unphysical representation. Jagged edges in trajectories indicate the collisions are taking place.

## 4. CONCLUSIONS

A simulation was produced that showed twodimensional motion of 10 particles.

- C. Kittel, Introduction to Solid State Physics (2005), eighth ed., ISBN 0-4-41526-X.
- [2] W. Humphrey, A. Dalke, and K. Schulten, Journal of Molecular Graphics 14, 33 (1996).

### Appendix A: Statement of Collaborators

I collaborated with Melanie on this assignment.

#### Appendix B: Survey Question

I spent approximately 4 hours coding for this assignment, 2 hours visualizing using VMD and making the

video, and 5 hours on the write-up. I most enjoyed making the visualization, but it was frustrating to try to figure out a way to save the visualization that I had made because the Make Movie feature required an external download that I was struggling to find. Because of this I ended up not using VMD to do so, and instead I took screenshots of the frames and compiled them in iMovie. I thought writing the code for this project was pretty simple but helped me to engage with the theory a little bit more, and the most valuable thing I learned from this assignment was probably gaining some familiarity with VMD. The length of this assignment was just right for me.