# Analyze molecular interaction for mixture of argon and particles with different sizes

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Abstract. This article elaborates on molecular dynamics simulation, which is a technology that supports us in analyzing the structure and dynamics of materials and their properties at a microscopic level. In this article, the molecular dynamics simulation develops on Lennard-Jones potential. The simulation is based on the fundamental understanding between Argon-Argon molecules. The application for running molecular dynamics simulation is Moldy. The Argon particles' size is changed by altering van der Waals radius from Lennard-Jones potential parameters, and the mixture of Argon and different sizes' particles will be simulated by Moldy. The average potential energy is collected and compared. It discusses how the size of particles affect the average potential energy. Moreover, the trend of potential energy respects to timesteps for each molecule is depicted and compared graphicly by Gnuplot. It discusses how the potential energy changes according to timesteps for different sizes' particles. In the end, the result is collected and summarized. It discovers that potential energy increases slowly when particles' sizes increase, and the potential energy decreases significantly when particles' sizes decrease. In addition, it founds out that molecules with different sizes have a noticeable change initially. Oddly, there is no observation of significant changes in the potential energy of the original molecule. Moreover, the initial decrease for molecules that increase in size is not as significant as that for molecules that decrease in size.

Keywords: Moldy, Molecular Simulation, Argon-Argon, Lennard-Jones Potential, Gnuplot.

#### 1. Introduction

In material sciences and engineering, molecular dynamics simulation is one of the technologies that supports us in observing and measuring the structure and dynamics of materials and their properties at a microscopic level [1]. Lennard-Jones potential is widely used for molecular dynamics simulation, and it can describe the potential energy of interactions between atoms and molecules, like Argon [2]. Currently, we have a deep understanding on when and how to employ Lennard-Jones potential to analyze Argon, like calculating the thermodynamic properties and shear viscosity. Most of the simulations ran between Argon atoms (Ar-Ar) or between Argon and Nickel (Ar-Ni). There is no research discussed about the interaction between the mixture of Argon and Argon which have different particle sizes. This article will elaborate in this area. The molecular simulation for the mixture of Argon will be run by Moldy. In Moldy, the distance of two particles when intermolecular potential is zero from Lennard-Jones potential parameters is changed to reach the goal of different sizes of particles. It allows

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me to observe how the potential energy changes between different sizes of particle. The comparison of the trend of potential energy for different sizes of particles with respect to timesteps is compared by using Gnuplot. It can help us to have a better understanding of the change of potential energy respects to timesteps for different sizes of particles. The purpose of this study is to explore more on potential energy of interaction between molecules under a microscopic level via molecular dynamic aspect.

# 2. Methodology

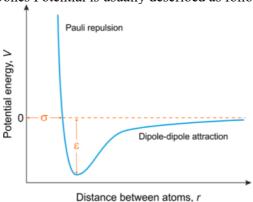
Lennard-Jones potential and its relative equation are employed in this article. Lennard-Jones potential is proposed by Sir John Edward Lennard-Jones. It is used to describe the potential energy for the interaction between molecules or atoms based on their separation. The Lennard-Jones potential equation is derived from the difference between attractive forces and repulsive forces. Lennard-Jones potential can be represented by following equation:

$$V(r) = 4\varepsilon \left[ \left( \frac{\sigma}{r} \right)^{12} - \left( \frac{\sigma}{r} \right)^{6} \right]$$
 (1)

Or

$$V(r) = \frac{A}{r^{12}} - \frac{B}{r^6}$$
 (2)

V is the intermolecular potential energy between two atoms or molecules;  $\varepsilon$  is the well depth, which measures how strongly the two atoms attract each other;  $\sigma$  is the distance between two particles where the intermolecular potential is zero, it is also known as van der Waals radius; r is the distance between two particles [3]. The Lennard-Jones Potential is usually described as following diagram (Figure 1).



**Figure 1.** Diagram of Lennard-Jones Potential [4].

In this article, the simulation is performed by altering the van der Waals radius from the Lennard-Jones Potential equation to change particles' sizes. The particles' size is bigger when the van der Waals radius decreases. It means that the particles are closer to each other. The particles' size is smaller when the van der Waals radius increases. It means that the particles are farther away from each other.

## 3. Analysis

This section first introduces how the molecular simulation runs based on Moldy and how the data is analyzed using Gnuplot. Then, it discusses the data that is abstracted from Gnuplot and interprets the correlation between the data.

# 3.1. Framework and background of applied algorithm and technology

In this article, there are two applications that are used, which are Moldy and VMD. Moldy is used to run the molecular-dynamics simulation based on the Lennard-Jones Potential. VMD is used to illustrate the interaction between molecules visually, based on the result from Moldy.

- 3.1.1. Moldy. Moldy is the C program for running molecular-dynamics simulations of molecules or atoms. The model system contains a mixture of atoms, ions, or molecules in a run-time input file. Rigid-molecule approximation is used to treat molecules or ions, and quaternion methods are used to model their rotational motion. Beeman algorithm is used to present the equations of motion [5]. In this article, the molecular-dynamics simulation runs based on Lennard-Jones Potential for mixtures of Argon. The file for illustrating Argon's basic information includes Argon's atomic mass and Lennard-Jones Potential values. The value for the depth of attractive well is 3.984 and interparticle distance is 3.41 for Argon-Argon molecule. There is another file needed to run the simulation for the mixture of Argons. It contains how the simulation performs, like temperature, density, and scale interval.
- 3.1.2. Gnuplot. Gnuplot is a command-driven program. It can present mathematical functions and data visually [6]. In this article, Gnuplot is used to demonstrate the potential energy for each simulation step for Argon-Argon molecule with different particles' sizes. It allows us to see the trend of fluctuating potential energy, and the trend for each molecule is compared. In the end, we can have a deep understanding of the molecular interaction between each molecule.

# 3.2. Data and explanation

3.2.1. Constant parameters. The constant parameters are the variables that do not change during the simulation. In Moldy, the van der Waals radius ( $\sigma$ ) from the file contains basic information that will be changed, and the rest of the information is the same. In Figure 2, 3.41 is the initial value for van der Waals radius, and this value will be modified for further simulation.

```
# LJ Argon - about as simple as you can get
# Parameters from Allen and Tildesley Table 1.1
Argon 108
1 0 0 0 39.948 0 Ar
end
Lennard-Jones
1 1 3.984 3.41
```

**Figure 2.** File of Argon's Basic Information.

Another file contains the code for running simulation for Argon is shown in Figure 3. The information from this file is the same through the entire simulation, which can be defined as independent variables.

3.2.2. Average potential energy respected with variance in van der Waals radius. The van der Waals radius ( $\sigma$ ) is changed in this section. The initial value is 3.41 and change the value by increasing or decreasing. The average potential energy is collected. The Table 1 below illustrates how the average potential energy is changed based on the difference in van der Waals radius. The initial value is highlighted in yellow.

Based on Table 1, it is evident to see that the average potential energy increases when van der Waals radius increases, and the average potential energy increases when van der Waals decreases. In other words, the average potential energy is higher when the particles' size decreases, and the average potential energy is higher when the particles' size increases. Moldy is unable to process with 5.41 Angstrom for van der Waals radius because the particles are too big, which makes them too close to each other. The graph of average potential energy versus van der Waals radius is portrayed in Figure 4. According to the graph, the change in average potential energy increases slightly when van der Waals radius decreases, and the change in average potential energy increases significantly when van der Waals radius increases from 3.91 to 4.91.

```
This sets up conditions at the triple point
sys-spec-file=argon.in.4
density=1.428
temperature=84
scale-interval=5
scale-end=10000
nsteps=15000
print-interval=100
roll-interval=100
begin-average=10001
average-interval=5000
step=0.01
subcell=2
strict-cutoff=1
cutoff=8.5125 #2.5 * sigma
begin-rdf=10001
rdf-interval=50
rdf-out=5000
begin-dump=0
dump-interval=5
dump-file=argon-dmp
dump-level=1
ndumps=10000
```

Figure 3. File of Running Simulation for Argon.

Table 1. Change in Average Potential Energy based on van der Waals Radius.

van der Waals Radius (A)	Average Potential Energy (kJ/mol)
1.41	-296.35
2.41	-375.21
3.41	-666.83
3.91	-591.14
4.41	2828
4.91	19224
5.41	/

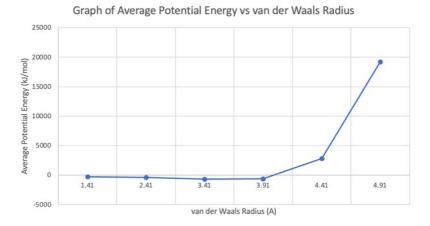


Figure 4. Graph of Average Potential Energy vs van der Waals Radius.

3.2.3. Analysis of trend of potential energy. The potential energy respects to each timestep are depicted graphically by using Gnuplot. The trend for each van der Waals radius is shown in Figure 5. Overall, the trend of potential energy respects to timesteps is similar for the particles with 4.91 Angstrom and 4.41 Angstrom. The trends for the rest are not clear due to large range of potential energy between van der Waals radius of 4.91 Angstrom and 1.41 Angstrom.

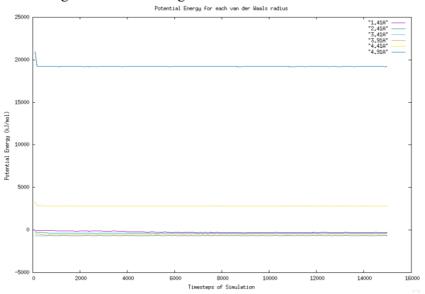


Figure 5. Graph of Potential Energy vs Timesteps For each van der Waals radius.

The trend comparison for particles with 4.91 Angstrom and 4.41 Angstrom are drawn in Figure 6. From the graph, we know that both start with the highest potential energy, and the potential energy initially decreases until it reaches a point, then the rest seems constant.

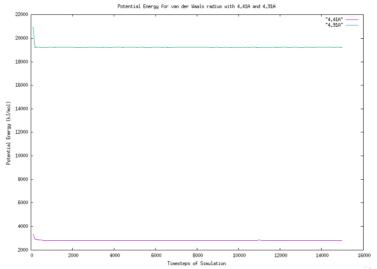
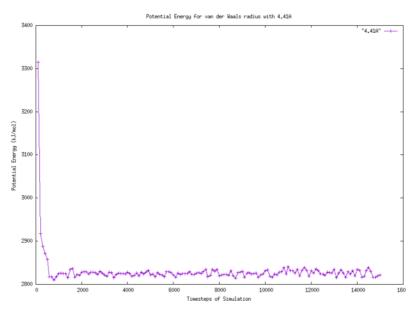


Figure 6. Graph of Potential Energy For van der Waals radius with 4.41A and 4.91A.

The range of potential energy between van der Waals radius of 4.41 Angstrom and 4.91 Angstrom is large. When both stop decreasing, the trends for potential energy are not very clear and easy to observe. The graph of potential energy with 4.41 Angstrom van der Waals radius is drawn in Figure 7.



**Figure 7.** Graph of Potential Energy for van der Waals radius with 4.41A.

According to Figure 7, it takes approximately 900 timesteps to decrease the potential energy from 3400 kJ/mol to 2830 kJ/mol. The oscillation of potential energy after 900 timesteps follows similar pattern without any outstanding number. The graph of potential energy with 4.91 Angstrom van der Waals radius is drawn in Figure 8.

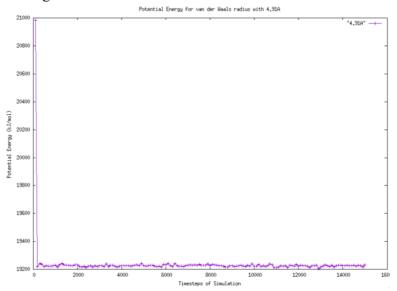


Figure 8. Graph of Potential Energy for van der Waals radius with 4.91A.

From Figure 8, it is evident to see the potential energy suddenly decreases from the highest potential energy which is about 21000 kJ/mol to approximately 19250 kJ/mol in 100 timesteps. Comparing with Figure 7, the overall trend is similar that the potential energy decreases from the highest potential energy and then the potential energy changes in the similar pattern. The potential energy with 1.41 Angstrom, 2.41 Angstrom, 3.41 Angstrom and 3.91 Angstrom van der Waals radius are compared in Figure 9. To clarify, the data relates to 3.41 Angstrom van der Waals radius is the original one.

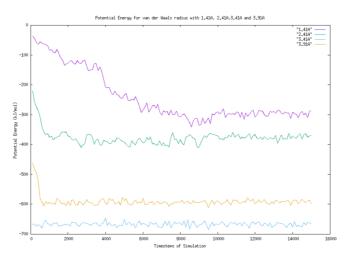


Figure 9. Graph of Potential Energy for van der Waals radius with 1.41A, 2.41A, 3.41A and 3.91A. According to Figure 8, there is no sudden change for the molecules with 3.41 Angstrom van der Waals radius. For the molecules with 3.91 Angstrom van der radius, the potential energy decreases significantly in first 300 timesteps from -480 kJ/mol to -600 kJ/mol. Comparing the trend of potential energy with molecules with 4.41 Angstrom and 4.91 Angstrom van der Waals radius, the overall trend is similar which is a sudden decreasing from the highest potential energy to a point where the change of potential energy is insignificant. These three trends have one thing in common that their sizes are all smaller than the original molecules. In addition, we can notice that there is a decreasing for 1.41 Angstrom and 2.41 Angstrom van der Waals radius from the highest potential energy like what happens for particles with 3.91 Angstrom van der Waals radius. However, the decreasing is not sharp. It can be concluded that the change in potential energy at initial timesteps for the molecules that have a bigger size than the original molecule is slow. Vice versa.

#### 4. Conclusion

This article discusses how the potential energy changes with respect to different particle sizes by using Moldy and Gnuplot. Firstly, the article discusses the methodology that is employed, which is Lennard-Jones Potential Function. Then, it introduces the application to run molecular dynamics simulation which is Moldy, and the application to analyze the results from simulations, which is Gnuplot. The analysis is based on Lennard-Jones Potential, and van deer Waals radius is changed to reach the goal. To conclude the result, the average potential energy for each molecule is discovered from Moldy. It is surprising found that the potential energy increases slowly when particles' sizes increase, and the potential energy decreases significantly when particles' sizes decrease. The trend of the change of potential energy for each molecule is depicted and compared graphicly from Gnuplot. It discovers that the molecules with changed the sizes have a noticeable change initially, while it does not observe in the original molecule. In addition, the initial decreasing for molecules that increases the size is not as significant as the molecules that decreases the size.

This article has some drawbacks. One drawback is that it does not show visually how the difference in potential energy affects molecules' movements. In the future, we can try to visualize the molecular movement respect to different potential energy. It can help readers to have a better understanding on molecular interaction for different sizes of particles. Another drawback is that this article does not further discuss why the potential energy increases when particles' size increases and decreases, and it does not expand on why there is a sudden decreasing for the particles with smaller sizes than origin at initial timesteps. The future can be spent researching those questions.

## 5. Acknowledgment

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to analyze the result from Moldy. He also helped me to target the research topic. In addition, I also want to thank Shuai Yuan who is a graduate student from University of Southern California. He provided a lot of help on coding the Moldy and answered my questions during the simulation. Without their help, this research will not be completed. Finally, I want to thank my parents who offer me this opportunity to study the molecular simulation. It is hard to complete this project without their support.

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