

# Molecular Dynamics Simulation of a Van der Waals Gas

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<b>1 Project-III: Simulation of a Van der Waals gas using Molecular Dynamics.</b>	<b>1</b>
1.1 Brief description of the project . . . . .	1
1.2 Team Members and Responsibilities . . . . .	1
1.3 Prerequisites . . . . .	1
1.4 How to . . . . .	2
1.4.1 Quick guide . . . . .	2
1.5 Help . . . . .	2
1.6 Contributors . . . . .	2
<b>2 Modules Index</b>	<b>5</b>
2.1 Modules List . . . . .	5
<b>3 Module Documentation</b>	<b>7</b>
3.1 forces Module Reference . . . . .	7
3.1.1 Detailed Description . . . . .	7
3.1.2 Function/Subroutine Documentation . . . . .	7
3.1.2.1 find_force_lj() [1/2] . . . . .	7
3.1.2.2 find_force_lj() [2/2] . . . . .	8
3.2 initialization Module Reference . . . . .	9
3.2.1 Detailed Description . . . . .	9
3.2.2 Function/Subroutine Documentation . . . . .	9
3.2.2.1 distribute_particles() . . . . .	9
3.2.2.2 initialize_positions() . . . . .	10
3.2.2.3 initialize_velocities() . . . . .	10
3.3 integrators Module Reference . . . . .	11
3.3.1 Detailed Description . . . . .	11
3.3.2 Function/Subroutine Documentation . . . . .	11
3.3.2.1 bm() . . . . .	11
3.3.2.2 inst_temp() . . . . .	12
3.3.2.3 kinetic_energy() . . . . .	12
3.3.2.4 momentum() . . . . .	13
3.3.2.5 time_step_euler_pbc() . . . . .	13
3.3.2.6 time_step_verlet() . . . . .	13
3.3.2.7 time_step_vverlet() [1/2] . . . . .	14
3.3.2.8 time_step_vverlet() [2/2] . . . . .	15
<b>Index</b>	<b>17</b>



# Chapter 1

## Project-III: Simulation of a Van der Waals gas using Molecular Dynamics.

### 1.1 Brief description of the project

This project involves the development of a Molecular Dynamics code to simulate a Van der Waals gas, performing calculations in both serial and parallel modes.

Additionally, this project requires collaborative work to ensure the proper functioning of the code.

### 1.2 Team Members and Responsibilities

Each team member has assigned tasks, which are indicated below:

1. **Anna Monclús (@anna-mr98)**: Initialize the configuration and define boundary conditions. Also coordinates the GitHub repository.
1. **Aina Gaya (@ainagaya)**: Integration Newton's equations.
2. **Albert Plazas (@Alplalo)**: Compute the forces for a Van der Waals interaction.
3. **Manel Serrano (@gluoon8)**: Post-processing of data, statistics, and visualization.

### 1.3 Prerequisites

To execute the program, there are some pre-requisites:

- Make: to execute the program ( <https://www.gnu.org/software/make/#download>)
- Gfortran: to run the MD simulation. ( <https://gcc.gnu.org/wiki/GFortran>)
- Python 3.x : to generate the plots after simulation
  - Numpy ( <https://numpy.org/install/>)
  - Matplotlib ( <https://matplotlib.org/stable/users/installing/index.html>)

## 1.4 How to

[!IMPORTANT] Current features are only available for serial code, parallel code is WIP!

1. Clone repository to your local host
2. Choose *serial* or *parallel* (WIP) folder with `cd serial` or `cd parallel`
3. Use `make` or `make help` to see the available commands.
4. Before starting a simulation, change your parameters in *namMD.nml* file
5. To carry out the simulation, use `make run` and the program will be compiled and run.
6. Data is generated in \*.dat\* files. If you want to generate figures, use `make plot`

### 1.4.1 Quick guide

To carry out a simulation after choosing parameters in *namMD.nml* file you can use:

```
make run
make plot
```

And files will appear in your main directory!

[!TIP] There are some ways to clean generated files, have a look at `make clean`, `make cleandata` and `make cleanplot`.

## 1.5 Help

- Commands:

- `make run`: Compiles needed files and also runs the program.
- `make plot`: Plots the output data:
  - \* Epot, Ekin, Etot vs time
  - \* Momentum vs time
  - \* T vs time
  - \* Pressure vs time
- `make all`: Compiles the program and creates executable MD.exe
- `make clean`: Removes the modules, objects and executable
- `make cleandata`: Removes data files
- `make cleanplot`: Removes plot files

## 1.6 Contributors

Anna Monclús	Aina Gaya	Albert Plazas	Manel Serrano
anna-mr98	ainagaya	Alplalo	gluoon8

Work developed in the Advanced Informatic Tools subject from [Master of Atomistic and Multiscale Computational Modelling in Physics, Chemistry and Biochemistry](#).

Universitat de Barcelona	Universitat Politècnica de Catalunya





## Chapter 2

# Modules Index

### 2.1 Modules List

Here is a list of all documented modules with brief descriptions:

#### forces

This module contains the subroutine `find_force_LJ` which calculates the forces and potential energy between particles using the Lennard-Jones potential. The module provides a public interface for the `find_force_LJ` subroutine. The `find_force_LJ` subroutine takes in the positions of particles, box size, cutoff distance, and returns the forces acting on the particles and the total potential energy. The forces are calculated using periodic boundary conditions (PBC). The module also contains a private section for internal use . . . . .

7

#### initialization

Module containing initialization routines for molecular dynamics simulations . . . . .

9

#### integrators

Module containing various integrators for molecular dynamics simulations . . . . .

11



## Chapter 3

# Module Documentation

### 3.1 forces Module Reference

This module contains the subroutine `find_force_LJ` which calculates the forces and potential energy between particles using the Lennard-Jones potential. The module provides a public interface for the `find_force_LJ` subroutine. The `find_force_LJ` subroutine takes in the positions of particles, box size, cutoff distance, and returns the forces acting on the particles and the total potential energy. The forces are calculated using periodic boundary conditions (PBC). The module also contains a private section for internal use.

#### Functions/Subroutines

- subroutine, public [find\\_force\\_lj](#) (`r`, `n`, `l`, `cutoff`, `f`, `pot`, `ppot`, `nprocs`, `rank`, `counts_recv`, `displs_recv`, `imin`, `imax`)  
*Calculates the forces and potential energy between particles using the Lennard-Jones potential.*
- subroutine, public [find\\_force\\_lj](#) (`r`, `n`, `l`, `cutoff`, `f`, `pot`, `ppot`)  
*Calculates the forces and potential energy between particles using the Lennard-Jones potential.*

#### 3.1.1 Detailed Description

This module contains the subroutine `find_force_LJ` which calculates the forces and potential energy between particles using the Lennard-Jones potential. The module provides a public interface for the `find_force_LJ` subroutine. The `find_force_LJ` subroutine takes in the positions of particles, box size, cutoff distance, and returns the forces acting on the particles and the total potential energy. The forces are calculated using periodic boundary conditions (PBC). The module also contains a private section for internal use.

#### 3.1.2 Function/Subroutine Documentation

##### 3.1.2.1 `find_force_lj()` [1/2]

```
subroutine, public forces::find_force_lj (  
    real(8), dimension(n, 3), intent(in) r,  
    integer, intent(in) n,  
    real(8), intent(in) l,  
    real(8), intent(in) cutoff,  
    real(8), dimension(n, 3), intent(out) f,
```

```

real(8), intent(out) pot,
real(8), intent(out) ppot )

```

Calculates the forces and potential energy between particles using the Lennard-Jones potential.

This subroutine calculates the forces and potential energy between particles in a system using the Lennard-Jones potential. The Lennard-Jones potential is a pairwise potential that models the interaction between neutral atoms or molecules. It is commonly used to simulate the behavior of noble gases and other non-reactive systems.

Parameters: *r*: real(8), dimension(N, 3), intent(in)

- The positions of the particles in the system. *N*: integer, intent(in)
- The number of particles in the system. *L*: real(8), intent(in)
- The length of the simulation box. *cutoff*: real(8), intent(in)
- The cutoff distance for the Lennard-Jones potential. *F*: real(8), dimension(N, 3), intent(out)
- The forces acting on each particle. *pot*: real(8), intent(out)
- The potential energy of the system.

Notes:

- The positions of the particles are given as a 3D array, where each row represents the position of a particle in Cartesian coordinates.
- The forces acting on each particle are calculated and stored in the *F* array.
- The potential energy of the system is calculated and stored in the *pot* variable.
- The subroutine uses periodic boundary conditions to handle particles that are close to the edges of the simulation box.
- The subroutine assumes that the *pbcc* subroutine is defined elsewhere in the code, which handles the periodic boundary conditions.
- The subroutine assumes that the *isnan* function is available to check for NaN values.

### 3.1.2.2 find\_force\_lj() [2/2]

```

subroutine, public forces::find_force_lj (
    real(8), dimension(n, 3), intent(in) r,
    integer, intent(in) n,
    real(8), intent(in) l,
    real(8), intent(in) cutoff,
    real(8), dimension(n, 3), intent(out) f,
    real(8), intent(out) pot,
    real(8), intent(out) ppot,
    integer, intent(in) nprocs,
    integer, intent(in) rank,
    integer, dimension(:) counts_recv,
    integer, dimension(:) displs_recv,
    integer imin,
    integer imax )

```

Calculates the forces and potential energy between particles using the Lennard-Jones potential.

This subroutine calculates the forces and potential energy between particles in a system using the Lennard-Jones potential. The Lennard-Jones potential is a pairwise potential that models the interaction between neutral atoms or molecules. It is commonly used to simulate the behavior of noble gases and other non-reactive systems.

Parameters: *r*: real(8), dimension(N, 3), intent(in)

- The positions of the particles in the system. `N`: integer, intent(in)
- The number of particles in the system. `L`: real(8), intent(in)
- The length of the simulation box. `cutoff`: real(8), intent(in)
- The cutoff distance for the Lennard-Jones potential. `F`: real(8), dimension(`N`, 3), intent(out)
- The forces acting on each particle. `pot`: real(8), intent(out)
- The potential energy of the system.

Notes:

- The positions of the particles are given as a 3D array, where each row represents the position of a particle in Cartesian coordinates.
- The forces acting on each particle are calculated and stored in the `F` array.
- The potential energy of the system is calculated and stored in the `pot` variable.
- The subroutine uses periodic boundary conditions to handle particles that are close to the edges of the simulation box.
- The subroutine assumes that the `pbcc` subroutine is defined elsewhere in the code, which handles the periodic boundary conditions.
- The subroutine assumes that the `isnan` function is available to check for NaN values.

## 3.2 initialization Module Reference

Module containing initialization routines for molecular dynamics simulations.

### Functions/Subroutines

- subroutine [initialize\\_positions](#) (`n`, `rho`, `r`)  
*Initializes the positions of  $N$  particles in a cubic box with density  $\rho$ . The box is centered at the origin. The positions are stored in the array  $r$ . The length of the box is calculated as  $L = (N/\rho)^{1/3}$ . The box is divided in  $M = N^{1/3}$  cells. The length of each cell is  $a = L/M$ .*
- subroutine [initialize\\_velocities](#) (`n`, `v_ini`, `v`)  
*Initializes the velocities of  $N$  particles with a given velocity  $v_{ini}$ . The velocities are stored in the array  $v$ . The velocities are chosen randomly from a bimodal distribution.*
- subroutine [distribute\\_particles](#) (`n`, `rank`, `nprocs`, `imin`, `imax`)  
*Distributes the particles among the processors.*

### 3.2.1 Detailed Description

Module containing initialization routines for molecular dynamics simulations.

### 3.2.2 Function/Subroutine Documentation

#### 3.2.2.1 [distribute\\_particles\(\)](#)

```
subroutine initialization::distribute_particles (
    integer, intent(in) n,
    integer, intent(in) rank,
    integer, intent(in) nprocs,
    integer imin,
    integer imax )
```

Distributes the particles among the processors.

## Parameters

<i>N</i>	Number of particles.
<i>rank</i>	Rank of the processor.
<i>nprocs</i>	Number of processors.
<i>imin,imax</i>	Indexes of the particles to be handled by the processor.
<i>chunklength</i>	Number of particles to be handled by each processor.
<i>uneven_parts</i>	Number of processors that will handle an extra particle.
<i>i</i>	Loop variable.

**3.2.2.2 initialize\_positions()**

```
subroutine initialization::initialize_positions (
    integer, intent(in) n,
    real(8), intent(in) rho,
    real(8), dimension(n, 3), intent(out) r )
```

Initializes the positions of  $N$  particles in a cubic box with density  $\rho$ . The box is centered at the origin. The positions are stored in the array  $r$ . The length of the box is calculated as  $L = (N/\rho)^{1/3}$ . The box is divided in  $M = N^{1/3}$  cells. The length of each cell is  $a = L/M$ .

## Parameters

<i>N</i>	Number of particles.
<i>rho</i>	Density of the system.
<i>r</i>	Array containing the positions of the particles.
<i>L</i>	Length of the box.
<i>a</i>	Length of each cell.
<i>x,y,z</i>	Coordinates of the particle.
<i>ini</i>	Initial position of the box.
<i>M</i>	Number of cells in each direction.
<i>i,j,k</i>	Loop variables.
<i>particle</i>	Index of the particle.

**3.2.2.3 initialize\_velocities()**

```
subroutine initialization::initialize_velocities (
    integer, intent(in) n,
    real(8) v_ini,
    real(8), dimension(n, 3), intent(out) v )
```

Initializes the velocities of  $N$  particles with a given velocity  $v_{ini}$ . The velocities are stored in the array  $v$ . The velocities are chosen randomly from a bimodal distribution.

## Parameters

<i>N</i>	Number of particles.
<i>v_ini</i>	Initial velocity of the particles.

## Parameters

<i>v</i>	Array containing the velocities of the particles.
<i>v_i</i>	Velocity of the particle.
<i>rand</i>	Random number.
<i>particle</i>	Index of the particle.
<i>i</i>	Loop variable.

## 3.3 integrators Module Reference

Module containing various integrators for molecular dynamics simulations.

### Functions/Subroutines

- subroutine, public [time\\_step\\_vverlet](#) (*r*, *vel*, *pot*, *n*, *l*, *cutoff*, *dt*, *ppot*, *nprocs*, *rank*, *counts\_recv*, *displs\_recv*, *imin*, *imax*)  
*Perform a time step using the velocity Verlet integration method. Calculates new positions and velocities for particles based on forces and previous positions/velocities.*
- subroutine, public [bm](#) (*ndat*, *xnums*, *sigma*)  
*Generate random numbers following a Box-Muller transformation. Generates normally distributed random numbers using the Box-Muller transformation.*
- subroutine, public [kinetic\\_energy](#) (*vel*, *k\_energy*, *n*)
- real(8) function, public [inst\\_temp](#) (*n*, *k\_energy*)  
*Calculate the instantaneous temperature of the system.*
- subroutine, public [momentum](#) (*vel*, *p*, *n*)
- subroutine, public [therm\\_andersen](#) (*vel*, *nu*, *sigma\_gaussian*, *n*)
- subroutine [time\\_step\\_vverlet](#) (*r*, *vel*, *pot*, *n*, *l*, *cutoff*, *dt*, *ppot*)  
*Perform a time step using the velocity Verlet integration method. Calculates new positions and velocities for particles based on forces and previous positions/velocities.*
- subroutine [time\\_step\\_euler\\_pbc](#) (*r\_in*, *r\_out*, *vel*, *n*, *l*, *cutoff*, *dt*, *pot*)  
*Perform a time step using the Euler method with periodic boundary conditions. Calculates new positions and velocities for particles based on forces and previous positions/velocities.*
- subroutine [time\\_step\\_verlet](#) (*r*, *rold*, *vel*, *n*, *l*, *cutoff*, *dt*, *pot*)  
*Perform a time step using the Verlet integration method. Calculates new positions and velocities for particles based on forces and previous positions/velocities.*

### 3.3.1 Detailed Description

Module containing various integrators for molecular dynamics simulations.

### 3.3.2 Function/Subroutine Documentation

#### 3.3.2.1 [bm\(\)](#)

```
subroutine integrators::bm (
    integer, intent(in) ndat,
    real(8), dimension(ndat), intent(out) xnums,
    real(8), intent(in) sigma )
```

Generate random numbers following a Box-Muller transformation. Generates normally distributed random numbers using the Box-Muller transformation.

**Parameters**

	<i>ndat</i>	Number of data points (must be even).
	<i>xnums</i>	Output array containing the generated random numbers.
	<i>sigma</i>	Standard deviation of the normal distribution.
in	<i>ndat</i>	Number of data points (must be even)
out	<i>xnums</i>	Output array containing generated random numbers
in	<i>sigma</i>	Standard deviation of the normal distribution

**3.3.2.2 inst\_temp()**

```
real(8) function, public integrators::inst_temp (
    integer, intent(in) n,
    real(8), intent(in) k_energy )
```

Calculate the instantaneous temperature of the system.

**Parameters**

<i>N</i>	Number of particles.
<i>K_energy</i>	Kinetic energy of the particles.

**Returns**

Instantaneous temperature of the system.

**Parameters**

in	<i>n</i>	Number of particles
in	<i>k_energy</i>	Kinetic energy of the particles

**Returns**

Instantaneous temperature of the system

**3.3.2.3 kinetic\_energy()**

```
subroutine integrators::kinetic_energy (
    real(8), dimension(n, 3), intent(in) vel,
    real(8), intent(out) k_energy,
    integer, intent(in) n )
```

**Parameters**

in	<i>n</i>	Number of particles
in	<i>vel</i>	Array containing particle velocities
out	<i>k_energy</i>	Output variable for the kinetic energy



### 3.3.2.4 momentum()

```
subroutine integrators::momentum (
    real(8), dimension(n, 3), intent(in) vel,
    real(8), intent(out) p,
    integer, intent(in) n )
```

#### Parameters

in	<i>n</i>	Number of particles
in	<i>vel</i>	Array containing particle velocities
out	<i>p</i>	Output variable for the total momentum

### 3.3.2.5 time\_step\_euler\_pbc()

```
subroutine integrators::time_step_euler_pbc (
    real(8), dimension(n, 3), intent(in) r_in,
    real(8), dimension(n, 3), intent(out) r_out,
    real(8), dimension(n, 3) vel,
    integer, intent(in) n,
    real(8), intent(in) l,
    real(8), intent(in) cutoff,
    real(8), intent(in) dt,
    real(8) pot )
```

Perform a time step using the Euler method with periodic boundary conditions. Calculates new positions and velocities for particles based on forces and previous positions/velocities.

#### Parameters

	<i>r_in</i>	Input array containing initial particle positions.
	<i>r_out</i>	Output array containing updated particle positions.
	<i>vel</i>	Input/Output array containing particle velocities.
	<i>N</i>	Number of particles.
	<i>L</i>	Box size.
	<i>cutoff</i>	Cutoff distance for LJ potential.
	<i>dt</i>	Time step size.
	<i>pot</i>	Output potential energy.
in	<i>n</i>	Number of particles
in	<i>r_in</i>	Initial particle positions
out	<i>r_out</i>	Updated particle positions
	<i>vel</i>	Particle velocities
in	<i>cutoff</i>	Box size, time step size, cutoff distance

### 3.3.2.6 time\_step\_verlet()

```
subroutine integrators::time_step_verlet (
    real(8), dimension(n, 3), intent(inout) r,
```

```

real(8), dimension(n, 3), intent(inout) rold,
real(8), dimension(n, 3), intent(inout) vel,
integer, intent(in) n,
real(8), intent(in) l,
real(8), intent(in) cutoff,
real(8), intent(in) dt,
real(8) pot )

```

Perform a time step using the Verlet integration method. Calculates new positions and velocities for particles based on forces and previous positions/velocities.

#### Parameters

	<i>r</i>	Input/Output array containing current particle positions.
	<i>rold</i>	Input/Output array containing previous particle positions.
	<i>vel</i>	Input/Output array containing particle velocities.
	<i>N</i>	Number of particles.
	<i>L</i>	Box size.
	<i>cutoff</i>	Cutoff distance for LJ potential.
	<i>dt</i>	Time step size.
	<i>pot</i>	Output potential energy.
in	<i>n</i>	Number of particles
in, out	<i>r</i>	Current particle positions
in, out	<i>rold</i>	Previous particle positions
in, out	<i>vel</i>	Particle velocities
in	<i>l</i>	Time step size, cutoff distance, box size

#### 3.3.2.7 time\_step\_vverlet() [1/2]

```

subroutine integrators::time_step_vverlet (
    real(8), dimension(n, 3), intent(inout) r,
    real(8), dimension(n, 3), intent(inout) vel,
    real(8), intent(out) pot,
    integer, intent(in) n,
    real(8), intent(in) l,
    real(8), intent(in) cutoff,
    real(8), intent(in) dt,
    real(8), intent(out) ppot )

```

Perform a time step using the velocity Verlet integration method. Calculates new positions and velocities for particles based on forces and previous positions/velocities.

#### Parameters

	<i>r</i>	Input/Output array containing particle positions.
	<i>vel</i>	Input/Output array containing particle velocities.
	<i>pot</i>	Output potential energy.
	<i>N</i>	Number of particles.
	<i>L</i>	Box size.
	<i>cutoff</i>	Cutoff distance for LJ potential.
	<i>dt</i>	Time step size.
in	<i>n</i>	Number of particles

## Parameters

in, out	<i>r</i>	Particle positions
in, out	<i>vel</i>	Particle velocities
out	<i>ppot</i>	Potential energy
in	<i>cutoff</i>	Time step size, box size, cutoff distance

## 3.3.2.8 time\_step\_vverlet() [2/2]

```

subroutine, public integrators::time_step_vverlet (
    real(8), dimension(n, 3), intent(inout) r,
    real(8), dimension(n, 3), intent(inout) vel,
    real(8), intent(out) pot,
    integer, intent(in) n,
    real(8), intent(in) l,
    real(8), intent(in) cutoff,
    real(8), intent(in) dt,
    real(8), intent(out) ppot,
    integer nprocs,
    integer rank,
    integer, dimension(:) counts_recv,
    integer, dimension(:) displs_recv,
    integer imin,
    integer imax )

```

Perform a time step using the velocity Verlet integration method. Calculates new positions and velocities for particles based on forces and previous positions/velocities.

## Parameters

	<i>r</i>	Input/Output array containing particle positions.
	<i>vel</i>	Input/Output array containing particle velocities.
	<i>pot</i>	Output potential energy.
	<i>N</i>	Number of particles.
	<i>L</i>	Box size.
	<i>cutoff</i>	Cutoff distance for LJ potential.
	<i>dt</i>	Time step size.
in	<i>n</i>	Number of particles
in, out	<i>r</i>	Particle positions
in, out	<i>vel</i>	Particle velocities
out	<i>ppot</i>	Potential energy
in	<i>cutoff</i>	Time step size, box size, cutoff distance



# Index

- bm
  - integrators, [11](#)
- distribute\_particles
  - initialization, [9](#)
- find\_force\_lj
  - forces, [7](#), [8](#)
- forces, [7](#)
  - find\_force\_lj, [7](#), [8](#)
- initialization, [9](#)
  - distribute\_particles, [9](#)
  - initialize\_positions, [10](#)
  - initialize\_velocities, [10](#)
- initialize\_positions
  - initialization, [10](#)
- initialize\_velocities
  - initialization, [10](#)
- inst\_temp
  - integrators, [12](#)
- integrators, [11](#)
  - bm, [11](#)
  - inst\_temp, [12](#)
  - kinetic\_energy, [12](#)
  - momentum, [13](#)
  - time\_step\_euler\_pbc, [13](#)
  - time\_step\_verlet, [13](#)
  - time\_step\_vverlet, [14](#), [15](#)
- kinetic\_energy
  - integrators, [12](#)
- momentum
  - integrators, [13](#)
- Project-III: Simulation of a Van der Waals gas using  
Molecular Dynamics., [1](#)
- time\_step\_euler\_pbc
  - integrators, [13](#)
- time\_step\_verlet
  - integrators, [13](#)
- time\_step\_vverlet
  - integrators, [14](#), [15](#)