## Data Science Course Group: Simulation Exercise 8th Session Lecture Materials

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## 1. Lecture Schedule

- This exercise will be conducted during the Spring 1st Term.
- Lecture materials will be uploaded by 11:00 AM on the scheduled day of each lecture.
- Schedule
  - 4/15: 1st Session
  - 4/22: 2nd Session
  - 3 4/30 (Tue): 3rd Session
  - 4 5/13: 4th Session (Midterm Report Assignment Released)
  - 5 5/20: 5th Session
  - 6 5/27: 6th Session (Midterm Report Submission Deadline)
  - 7 6/03: No Class
  - 8 6/10: 7th Session
  - 9 6/17: 8th Session (Final Report Assignment Released)
  - 6/24: 9th Session (Make-up Class)

# 1.1. Syllabus

The following topics are planned to be covered in this exercise (subject to change depending on progress):

- Introduction
  - Usage of C(C++) (mainly for numerical computation)
  - Usage of Python (for data analysis and plotting)
  - Principles of numerical computation
  - Loss of significance
  - Nondimensionalization in scientific computation
- Numerical solutions of ordinary differential equations: Examples of damped oscillations and harmonic oscillators
  - Numerical integration of differential equations
  - Euler method
- Brownian motion of a single particle
  - Langevin equation (stochastic differential equation)
  - Method of generating normal random numbers
  - Euler-Maruyama method
  - Time average and ensemble average
  - Calculation of the diffusion coefficient
- Brownian motion of multi-particle systems
  - Method of calculating interaction forces
  - Simulation of nonequilibrium systems: Example of phase separation
- 5 Molecular dynamics simulation of multi-particle systems
  - Position Verlet method and velocity Verlet method
  - Conservation laws in multi-particle systems (momentum, energy, angular momentum)
- 6 Monte Carlo method
  - Review of statistical mechanics
  - Markov chain Monte Carlo method

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# 2. Explanation of the 7th Assignment

### Assignment 7 Implementation of Molecular Dynamics Simulation (Classical MD)

Consider a 2D system confined within a periodic boundary of length L=40a on each side, consisting of identical disk particles with a number of particles N = 1024, diameter a, and mass m. The interparticle potential used here is a repulsive-only potential

$$U(r_{jk}) = \epsilon \left(\frac{a_{jk}}{r_{jk}}\right)^{12} + C_{jk} \quad (r_{jk} < a_{\text{cut}})$$

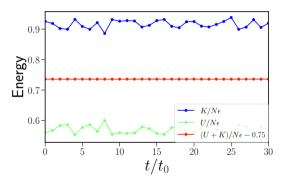
where the cutoff length is  $a_{\rm cut} = 3.0a$ . Let the units of length be  $a_{\rm cut}$ , energy be  $\epsilon_{\rm cut}$ , and time be  $t_0 = \sqrt{ma^2/\epsilon}$ . Answer the following auestions:

- Using the Langevin heat bath constructed in Assignment 6, obtain the position coordinates  $\{\mathbf{r}_i\}$  and velocities  $\{\mathbf{v}_i\}$  of each particle in the thermally equilibrated state at a dimensionless temperature  $T^* = k_B T/\epsilon = 0.9$ .
- Using the coordinates and velocities obtained in (1) as the initial conditions, execute a molecular dynamics simulation, and show that the mechanical energy U + K, where K is the kinetic energy and U is the potential energy is conserved over time.
- (Advanced Assignment Optimization) Accelerate the interaction calculation using the list (ledger) introduced in Assignment 5. The strategy is to create a list of particles within the radius  $r_{cut} + r_{skin}$ , and keep using this list until the maximum displacement of any particle exceeds  $r_{skin}/2$ .

#### **Explanation**

# 2. Explanation of the 7th Assignment (2)

The sample program for Assignment 7 (which does not include the optimization using lists), "md.cpp", is shown in List 1 below and can be obtained from the GitHub repository at **[Link]**.



☑ 1: Example of the solution for Assignment 7(2). It can be seen that the mechanical energy is conserved. Here, the energy per particle is shown.

# 2. Explanation of the 7th Assignment (3)

#### リスト 1: Sample program of the 7th Assignment"md.cpp". GitHub [Link].

```
#include <stdio.h>
    #include <stdlib.h>
    #include <math.h>
    #include <iomanip>
    #include <iostream>
    #include <fstream>
    #include <cfloat>
    #include "BM.h"
    #define Np 1024
    #define I. 40.0
    #define teg 100
    #define tmax 30
    #define dtmd 0.001
    #define dtbd 0.01
    #define temp 0.9
17
    #define dim 2
    #define cut 3.0
18
19
    #define polydispersity 0.0
20
    void ini_coord_square(double (*x)[dim]){
22
      int num_x = (int)sqrt(Np)+1;
23
      int num_y = (int)sqrt(Np)+1;
24
      int i.i.k=0:
25
      for(i=0:i<num_v:i++){
26
        for(i=0:i<num x:i++){
```

# 2. Explanation of the 7th Assignment (4)

```
27
            x[i+num_x*j][0] = i*L/(double)num_x;
28
            x[i+num \ x*i][1] = i*L/(double)num v:
29
            k++:
30
            if(k>=Np)
31
              break:
32
33
          if(k>=Np)
34
            break:
35
36
37
38
     void set diameter(double *a){
39
       for(int i=0:i<Np:i++)</pre>
40
          a[i]=1.0+polvdispersity*gaussian_rand():
41
42
43
     void p_boundary(double (*x)[dim]){
44
       for(int i=0:i<Np:i++)</pre>
45
          for(int i=0:i<dim:i++)</pre>
46
            x[i][i]-=L*floor(x[i][i]/L):
47
48
49
     void ini_arrav(double (*x)[dim]){
50
       for(int i=0:i<Np:i++)</pre>
51
          for(int j=0; j < dim; j++)</pre>
52
            x[i][i]=0.0:
53
```

# 2. Explanation of the 7th Assignment (5)

```
54
     void calc force(double (*x)[dim].double (*f)[dim].double *a.double *U){
56
       double dx, dy, dr2, dUr, w2, w6, w12, aij;
57
       double Ucut=1./pow(cut,12);
58
       ini arrav(f):
59
       *U=0:
60
       for(int i=0;i<Np;i++)</pre>
61
         for(int j=0; j<Np; j++) {</pre>
62
           if(i<i){
           dx=x[i][0]-x[i][0]:
64
           dv=x[i][1]-x[i][1]:
65
           dx = L * floor((dx+0.5*L)/L):
66
           dv = L * floor((dv + 0.5*L)/L);
67
           dr2=dx*dx+dv*dv:
68
           if(dr2<cut*cut){
69
             aii=0.5*(a[i]+a[i]):
70
             w2=aii*aii/dr2:
71
             w6=w2*w2*w2:
72
             w12=w6*w6:
73
             dUr = -12.*w12/dr2:
74
             f[i][0]-=dUr*dx:
75
             f[i][0]+=dUr*dx:
76
             f[i][1]-=dUr*dv:
77
             f[i][1]+=dUr*dv:
78
             *U+=w12-Ucut:
79
80
```

# 2. Explanation of the 7th Assignment (6)

```
81
82
83
84
     void eom_langevin(double (*v)[dim],double (*x)[dim],double (*f)[dim],double *a,double *U,double dt,double
           temp0){
85
        double zeta=1.0:
86
        double fluc=sqrt(2.*zeta*temp0*dt);
87
88
        calc force(x.f.a.&(*U)):
        for (int i=0:i<Np:i++)
90
          for(int i=0:i<dim:i++){</pre>
            v[i][i]+=-zeta*v[i][i]*dt+f[i][i]*dt+fluc*gaussian rand():
92
            x[i][i]+=v[i][i]*dt:
93
94
       p_boundary(x);
95
96
97
     void eom_md(double (*v)[dim].double (*x)[dim].double (*f)[dim].double *a.double *U.double dt){
98
        for(int i=0:i<Np:i++)</pre>
99
          for(int i=0:i<dim:i++){</pre>
100
            x[i][i]+=v[i][j]*dt+0.5*f[i][j]*dt*dt;
101
            v[i][i]+=0.5*f[i][i]*dt:
102
103
        calc_force(x.f.a.&(*U)):
104
        for(int i=0;i<Np;i++)</pre>
105
          for(int i=0:i<dim:i++){</pre>
106
            v[i][j]+=0.5*f[i][j]*dt;
```

# 2. Explanation of the 7th Assignment (7)

```
107
108
       p boundarv(x):
109
110
111
     void output(int k.double (*v)[dim].double U){
112
        char filename[128]:
113
        double K=0.0:
114
115
        std::ofstream file:
116
        sprintf(filename, "energy.dat");
117
        file.open(filename.std::ios::app): //append
118
        for(int i=0:i<Np:i++)
119
          for(int j=0:j<dim:j++)</pre>
120
            K+=0.5*v[i][i]*v[i][i]:
121
122
        std::cout<< std::setprecision(6)<<k*dtmd<<"\t"<<K/Np<<"\t"<<U/Np<<"\t"<<(K+U)/Np<<std::endl:
123
       file<< std::setprecision(6)<<k*dtmd<<"\t"<<K/Np<<"\t"<<U/Np<<"\t"<<(K+U)/Np<<std::endl:
124
        file.close():
125
126
127
     int main(){
128
        double x[Np][dim],v[Np][dim],f[Np][dim],a[Np];
129
       double tout=0.0.U:
130
       int j=0;
131
        set diameter(a):
132
        ini_coord_square(x):
133
        ini_array(v);
```

# 2. Explanation of the 7th Assignment (8)

```
134
135
        while(j*dtbd < 10.){</pre>
136
          1++:
137
          eom_langevin(v,x,f,a,&U,dtbd,5.0);
138
139
140
        i=0:
141
        while(j*dtbd < teq){</pre>
142
          1++:
143
          eom_langevin(v,x,f,a,&U,dtbd,temp);
144
145
        i=0:
146
        while(j*dtmd < tmax){</pre>
147
          j++;
148
          eom_md(v.x.f.a.&U.dtmd):
149
          if(j*dtmd >= tout){
150
             output(j,v,U);
151
             tout+=1.:
152
153
154
        return 0:
155
```

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# 3 Monte Carlo Method in Particle Systems

#### Monte Carlo Method in Particle Systems

- The Monte Carlo method (MC method) in particle systems is a method used to determine the distribution of particles in thermal equilibrium and the associated statistical quantities by using random numbers.
- In MD (Molecular Dynamics) calculations, the calculation of interparticle forces is crucial, while in the MC method, the calculation of energy is important.
- Except for this part, there are many overlapping aspects between the computational methods learned in Brownian dynamics (Langevin heat bath method) and molecular dynamics.
- The MC method primarily generates ensembles such as NVT, NPT, and μVT based on statistical mechanics.
- Below, the basic principles of the MC method are explained.

## 3.1 Markov Chain

## Markov Chain [?, ?, ?]

- Suppose that in  $\ell$  steps, the system transitions from state i to state k, and after another m steps, it reaches state i.
- If the transition probability between these states does not depend on the history leading to that state, this process is called a Markov chain.
- In particular, in the limit as *m* becomes large, the **probability of entering each state** becomes **independent of the initial state**. This is known as the **Markov chain's limit theorem**.

#### **Markov Chain Limit Theorem and Stationary State**

■ In a Markov chain, the probability of transitioning from state i to state j in  $\ell$  steps is denoted by  $P_{ij}(\ell)$ , and the following relationship holds (the **Chapman-Kolmogorov equation/Smoluchowski equation**):

$$P_{ij}(\ell+m) = \sum_{k=1}^{M} P_{ik}(\ell) P_{kj}(m)$$
(1)

## 3.1 Markov Chain (2)

In particular, since a thermal equilibrium state is obtained after a long time (many steps), it does not depend on the initial state. This means that

$$P_{ij}(\infty) = \sum_{k=1}^{M} P_{ik}(\infty) P_{kj}(1)$$
 (2)

Now, if we consider

$$P_{ij}(\infty) \to \Pi_j$$
 (3)

$$P_{ik}(\infty) \to \Pi_k \tag{4}$$

then  $\Pi_j$  represents the probability of the system being in state j in the stationary state. In a Markov chain,  $P_{kj}(1)$  does not depend on the previous state transition path, so it can be written as  $P_{kj}(1) = P_{kj}$ . Therefore.

$$\Pi_j = \sum_{k=1}^M \Pi_k P_{kj} \tag{5}$$

holds. Here.

$$\sum_{k=1}^{M} P_{jk} = 1$$
(6)

## 3.1 Markov Chain (3)

Therefore, equation (5) shows that

$$\left| \sum_{k=1}^{M} \Pi_{j} P_{jk} = \sum_{k=1}^{M} \Pi_{k} P_{kj} \right| \tag{7}$$

indicating that the inflow and outflow of probability into state j are balanced.

Moreover, this stationary state relationship corresponds to the master equation, which describes the time evolution of the probability of being in state j:

$$\frac{\partial \Pi_j}{\partial t} = \sum_{k=1}^M (\Pi_k P_{kj} - \Pi_j P_{jk}) \tag{8}$$

where  $\frac{\partial \Pi_j}{\partial t} = 0$  in the stationary state.

# 3.2 Microscopic Reversibility (Detailed Balance) Equation

#### Microscopic Reversibility (Detailed Balance) Equation

Next, we consider a special thermal equilibrium state within the stationary state. In thermal equilibrium, the detailed balance (microscopic reversibility) equation, which is the balance relation for all microscopic states. holds:

$$\Pi_i P_{ij} = \Pi_j P_{ji} \tag{9}$$

Of course, this detailed balance equation ensures the stationary state relationship discussed in the previous section.

• According to statistical mechanics,  $\Pi_i$ , which generates an NVT ensemble, is given by the canonical distribution:

$$\Pi_i = \frac{\exp(-\beta U_N(\mathbf{r}^N; i))}{Z_N} \tag{10}$$

where  $Z_N$  is the partition function:

$$Z_N = \sum_{i} \exp[-\beta \{U_N(\mathbf{r}^N; i)\}]$$
 (11)

# 3.2 Microscopic Reversibility (Detailed Balance) Equation (2)

■ In an NPT ensemble where pressure and temperature are constant, the probability  $\Pi_i$  is given by

$$\Pi_i = \frac{\exp[-\beta \{U_N(\mathbf{r}^N; i) + PV(\mathbf{r}^N; i)\}]}{Y_N}$$
(12)

where  $Y_N$  is the partition function for this case:

$$Y_N = \sum_{i} \exp[-\beta \{U_N(\mathbf{r}^N; i) + PV(\mathbf{r}^N; i)\}]$$
(13)

# 3.3 Metropolis Method

## Metropolis Method [?]

The Metropolis method is a technique in Monte Carlo simulations that provides statistical mechanics criteria for accepting or rejecting new states generated by random numbers. This weighted sampling method allows for the approximate calculation of distribution functions.

For various  $P_{ij}$ , it is possible to determine the probability  $\Pi_i$ . In this section, by acknowledging the principle of detailed balance, the transition probability  $P_{ij}$  that most simply achieves thermal equilibrium is given by

$$P_{ij} = \begin{cases} \frac{1}{M} & (\Pi_j > \Pi_i) \quad \text{accept} \\ \frac{1}{M} & \Pi_i & (\Pi_j \leq \Pi_i) \quad \text{accept} \\ \frac{1}{M} & (1 - \frac{\Pi_j}{\Pi_i}) & (\Pi_j \leq \Pi_i) \quad \text{reject} \end{cases} , \tag{14}$$

This rule is known as the **Metropolis Method** [?].

# 3.3 Metropolis Method (2)

In the Metropolis method, when Π<sub>j</sub> > Π<sub>i</sub>, as shown in Figure 2, the transition from a state with higher energy to a state with lower energy always occurs. That is,

$$P_{ij}=\frac{1}{M},$$

is used.

On the other hand, when  $\Pi_j < \Pi_i$ , the transition probability from a state j to a state i corresponding to a transition from a higher energy state to a lower energy state is

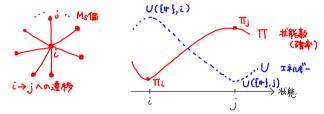
$$P_{ji}=\frac{1}{M},$$

Therefore, from the detailed balance equation, the transition probability  $P_{ij}$  is

$$P_{ij} = \frac{\Pi_j}{M\Pi_i},$$

This means that the transition from a lower energy state to a higher energy state  $i \to j$  occurs with probability  $\frac{\Pi_j}{\Pi_i}$  (it does not always occur). Conversely, the complementary event  $1 - \frac{\Pi_j}{\Pi_i}$  results in the rejection of the transition from i to j.

# 3.3 Metropolis Method (3)



 $\ensuremath{\mathbb{Z}}$  2: Transition from state i to state j. The number of possible transition states j (including i) is M. The transition  $i \to i$  indicates staying in the same state. "Rejection" in the Metropolis method corresponds to this. (Right) The relationship between probability  $\Pi$  and potential energy U. The higher the U, the lower the probability.

# 3.3 Metropolis Method

#### **Markov Chain Monte Carlo Method**

- Here, we will implement a Markov chain using the Metropolis method. The implementation of the Metropolis method involves trials using uniformly distributed random numbers, which constitute the Monte Carlo simulation. In general, such simulations are referred to as Markov Chain Monte Carlo (MCMC) Method.
- Now, the probability that the state  $\nu$  is generated under constant temperature and volume conditions, following the canonical distribution in statistical mechanics, is given by

$$\Pi_{\nu} \propto \exp(-\beta U_N(\mathbf{r}^N(\nu))),$$
 (15)

- Next, consider the state transition from {r<sup>N</sup>(ν)} to {r<sup>N</sup><sub>trial</sub>}. Here, each particle is randomly displaced individually.
- $\blacksquare$  Let k be the particle to be displaced, and the displaced state  $\mathbf{r}_{\mathrm{trial}}^k$

$$\mathbf{r}_{\text{trial}}^{k} = \mathbf{r}^{k}(\nu) + \Delta r \mathbf{R},\tag{16}$$

where **R** is a uniformly distributed random number in the range [-1,1].

■ The value of  $\Delta r$  is empirically found to be O(0.1).

# 3.3 Metropolis Method (2)

Intuitively, if  $\Delta r$  is too large, many transitions will be rejected in the following Metropolis criteria, reducing computational efficiency. This value must be finely adjusted depending on the problem.

Next, for the trial coordinates obtained by displacing particle k, perform a Metropolis criterion so that the canonical distribution is satisfied:

$$\begin{split} \frac{\Pi_{\text{trial}}}{\Pi_{\nu}} &= \frac{\exp[-\beta U_N(\{\mathbf{r}_{\text{trial}}^N\})]}{\exp[-\beta U_N(\{\mathbf{r}^N(\nu)\})]} \\ &= \exp[-\beta (U_N(\{\mathbf{r}_{\text{trial}}^N\}) - U_N(\{\mathbf{r}^N(\nu)\}))] \\ &= \exp[-\beta (\Delta U)] \end{split} \tag{17}$$

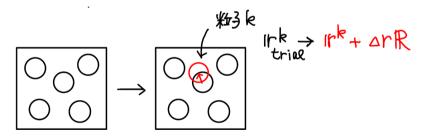
where  $\Delta U$  represents the change in energy.

# 3.3 Metropolis Method (3)

Using this probability, the following Metropolis criterion can be performed:

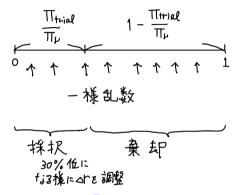
#### **Metropolis Criterion**

- (1) If  $\frac{\Pi_{\text{trial}}}{\Pi_{\text{tr}}} > 1$ : Always accept.
- (2) If  $\frac{\Pi_{trial}}{\Pi_{\nu}} < 1$ : Accept with probability  $\frac{\Pi_{trial}}{\Pi_{\nu}}$ . Reject with probability  $1 \frac{\Pi_{trial}}{\Pi_{\nu}}$ .



 $\boxtimes$  3: The random displacement of particle k is given by  $\mathbf{r}_{\text{trial}}^k = \mathbf{r}^k(\nu) + \Delta r \mathbf{R}$ . Here,  $\mathbf{R}$  is a uniformly distributed random number in the range [-1,1].

# 3.3 Metropolis Method (4)



 $\boxtimes$  4: Trial method for the Metropolis criterion  $\frac{\Pi_{\text{trial}}}{\Pi_{\nu}} < 1$ . Using a uniformly distributed random number R in the range [0,1], accept the trial if  $0 < R < \frac{\Pi_{\text{trial}}}{\Pi_{\nu}}$  is satisfied.

# 3.3 Metropolis Method (5)

Finally, we handle a special case. For a hard-sphere potential, the Metropolis criterion simplifies as follows:

## Metropolis Criterion for Hard Spheres

- (1) If  $\frac{\Pi_{trial}}{\Pi_{\nu}} = 1$ : Accept.
- (2) If  $\frac{\Pi_{trial}}{\Pi_{\nu}}$  < 1: Reject.

Calculations for hard spheres, which are challenging in MD simulations learned previously, are relatively easy using the Monte Carlo method.

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# 4. 8th Assignment

## 8th Assignment Implementation of Markov Chain Monte Carlo Method

Reproduce the phase separation phenomenon from Assignment 6 using the Markov Chain Monte Carlo method.

The sample program for Assignment 8 (Monte Carlo method) "mc.cpp" (which does not include optimization using lists) is available in the GitHub repository at [Link].

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## **References and Websites**

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- [2] Frenkel D, Smit B (2001) <u>Understanding Molecular Simulation: From Algorithms to Applications</u>. (Elsevier).
- [3] Allen MP, Tildesley DJ (2017) <u>Computer Simulation of Liquids: Second Edition</u>. (Oxford University Press).
- [4] Metropolis N, Rosenbluth AW, Rosenbluth MN, Teller AH, Teller E (1953) Equation of State Calculations by Fast Computing Machines. The Journal of Chemical Physics 21(6):1087–1092.