

# Computational Material Science

## Part I

Ito Chao (趙奕娣)  
Institute of Chemistry  
Academia Sinica

## ➤ Simulation conditions

NVT : constant N, V, T (number of particles, volume, temperature)  
- the so-called **canonical ensemble** (often used in MC)

NVE : constant N, V, E (number of particles, volume, energy)  
- the so-called **microcanonical ensemble** (often used in MD)

NPT : constant N, P, T (number of particles, pressure, temperature)  
- the so-called **isothermal-isobaric ensemble**

$\mu$ VT : constant  $\mu$ , V, T (chemical potential, volume, temperature)  
- the so-called **grand canonical ensemble**

*What is chemical potential ,  $\mu$  ?*

$$\mu_i = \left( \frac{\partial A}{\partial N_i} \right)_{T, V, N_{j \neq i}}$$

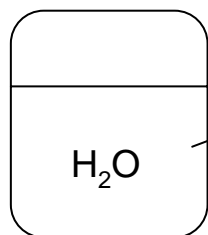
Under the condition of constant T and V, chemical potential  
is the partial derivative of Helmholtz free energy

- **Potential energy surface** can be based on
  - empirical energy function (MM)
  - quantum mechanical energy function (QM)
  - combined (QM/MM)
- **Initial configuration**: from experimental and/or optimized

Note: MD and MC are time-consuming when hundreds to thousands of atoms are handled. Do not start from a configuration that is far from your interested target.
- **Equilibration phase**: enable the system to evolve from the starting configuration to reach equilibrium (monitored by quantities such as E, T, P).
- **Averaging phase**: at this stage, data are accumulated for the calculation of interested quantities. (E.g., energy, heat capacity, free energy, radial distribution function, correlation function, diffusion constant, etc...)

- Boundaries

How to simulate the bulk with efficiency?



1 liter of water =  $3.3 \times 10^{25}$  molecules

The molecular diameter of water is  $2.8 \text{ \AA}$

If interactions with the walls extend up to 10 molecular diameters,  
Only  $2 \times 10^{19}$  molecules are interacting with the wall. (one in 1.5 million)

$\therefore$  Simulate bulk without considering the wall

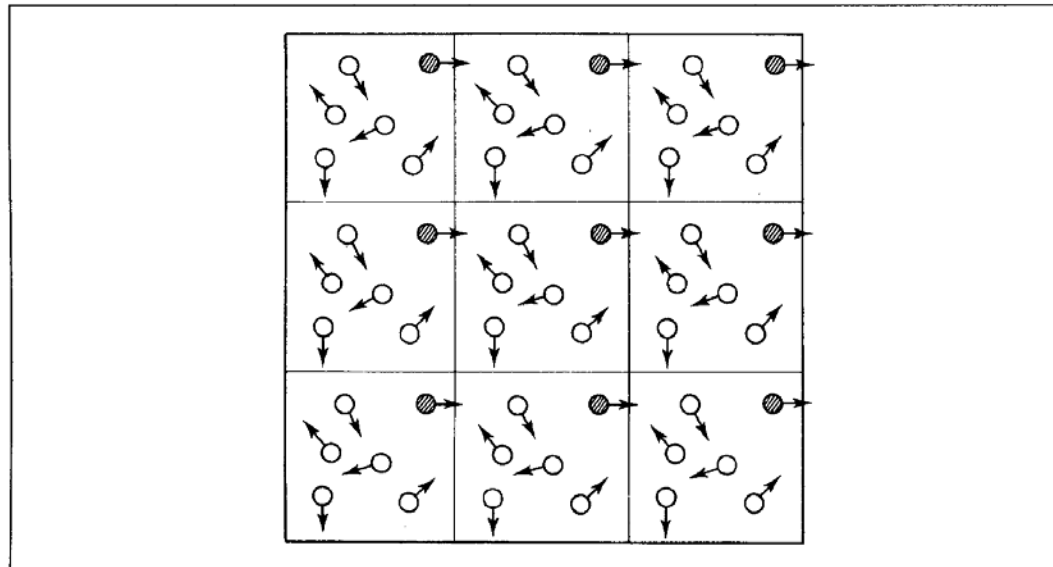
$\Rightarrow$  Periodic boundary condition

Examples: surfaces, bulk materials

$\Rightarrow$  Non-periodic boundary condition

Examples: hydrated enzyme, conformation analysis of a molecule  
or a molecular complex

⇒ Periodic boundary condition (PBC) : enable a simulation to be performed using a relatively small number of particles, in such a way that the particles experience forces as if they were in bulk fluid.



*Note: other shapes are possible in PBC*

Fig. 6.4: Periodic boundary conditions in two dimensions.

In 2-dimensional case, each box is surrounded by 8 neighbours;  
3-dimensional 26

- Coordinates of the particles in the image boxed can be computed by adding or subtracting integral multiples of the box sides.
- A particle leaving the box is replaced by an image particle that enters from the opposite side (so N is constant).

- Cutoff distances and the minimum image convention

In MM force fields, the numbers of bond-stretching, angle-bending, and torsion are proportional  $N$  (the number of atoms), but the number of **non-bonded terms increase as  $N^2$**  in a pairwise model.

∴ calc. of nonbonded  $\Rightarrow$  time consuming!  $\Rightarrow$  beyond a cutoff distance the energy is set to zero or calculated differently

The Lennard-Jones potential (for vdW interaction) falls off very rapidly with distance

$\Rightarrow$  at  $2.5\sigma$ , the Lennard-Jones potential has just 1% of its value at  $\sigma$ !

The electrostatic interaction ( $1/r$ ) falls off far less rapidly than Lennard-Jones interaction

$\Rightarrow$  larger cutoff distance is needed or special treatment  
(for details, see chap 6.8 of Molecular Modelling: Principles and Applications)

## *Rule of thumb for cutoff distance?*

- At least 10 Å
- The minimum image convention: each atom “sees” at most just one image of every other atom in the system; that is, should not see its own image or the same atom twice. => the cutoff distance cannot be larger than half of the cell length

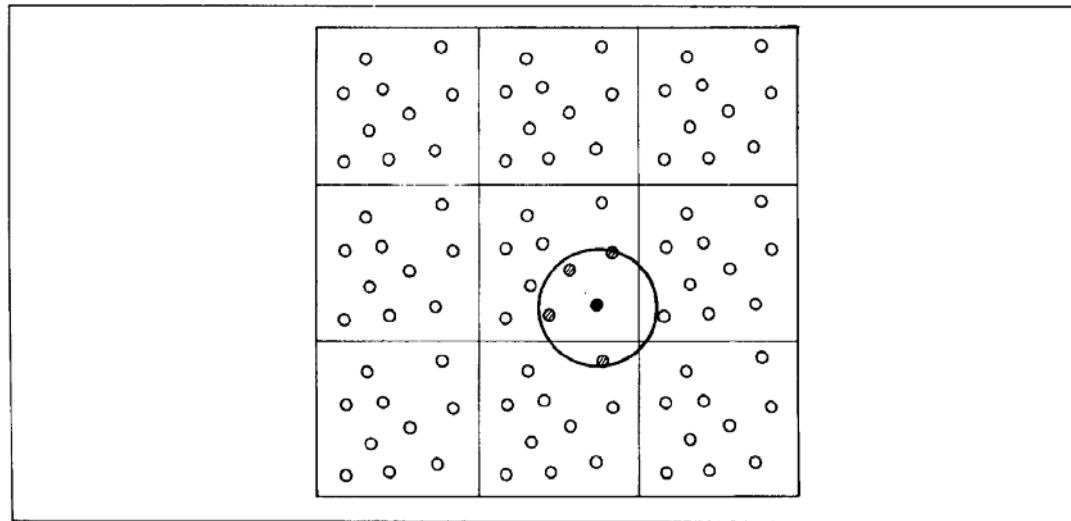
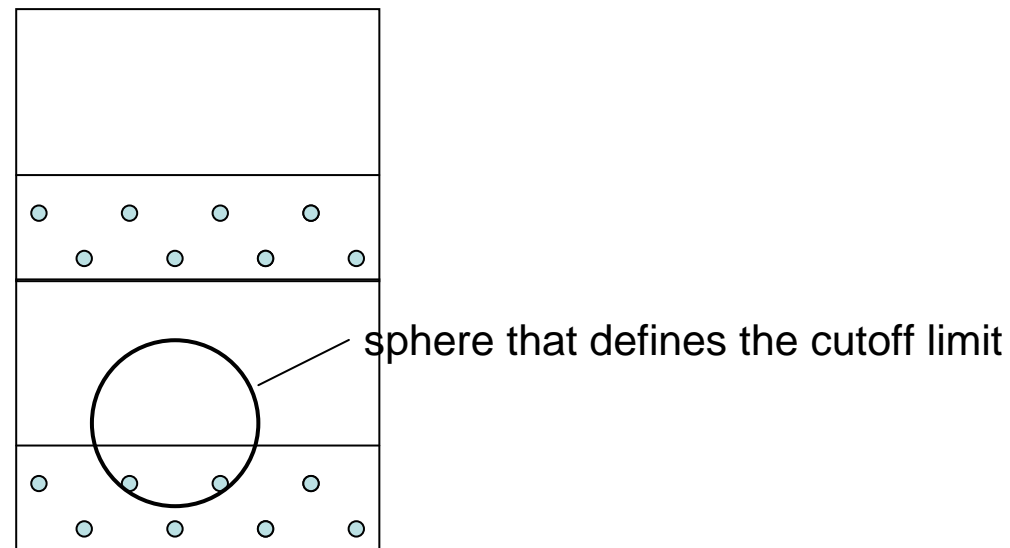


Fig. 6.11: The spherical cutoff and the minimum image convention.

PBC conditions are normally 3D. *How to model surface?*

- Increase the cell dimension for the direction perpendicular to the surface
- Choose a cell length that is longer than the cutoff distance, so one slab do not see the next slab





- Structure analysis : Radial distribution function (rdf)

Consider a spherical shell of thickness  $\delta r$  at a distance  $r$  from a chosen atom, the pair distribution function,  $g(r)$ , gives the probability of finding an atom in that shell compared to the ideal gas distribution.

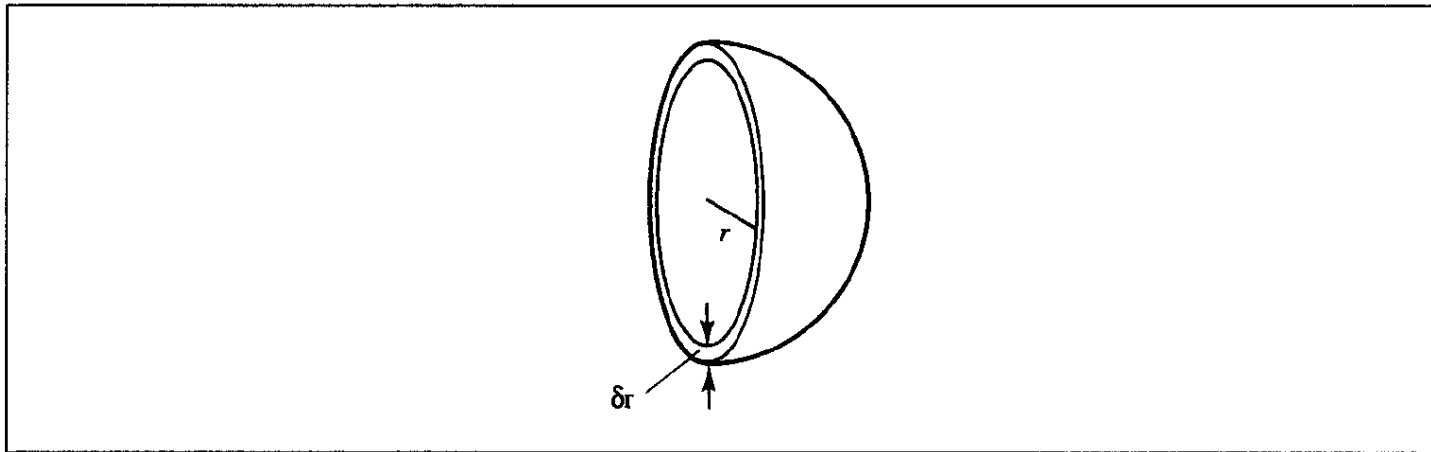


Fig. 6.1: Radial distribution functions use a spherical shell of thickness  $\delta r$ .

$$\begin{aligned} V &= \frac{4}{3}\pi(r + \delta r)^3 - \frac{4}{3}\pi r^3 \\ &= 4\pi r^2 \delta r + 4\pi r \delta r^2 + \frac{4}{3}\pi \delta r^3 \approx 4\pi r^2 \delta r \end{aligned}$$

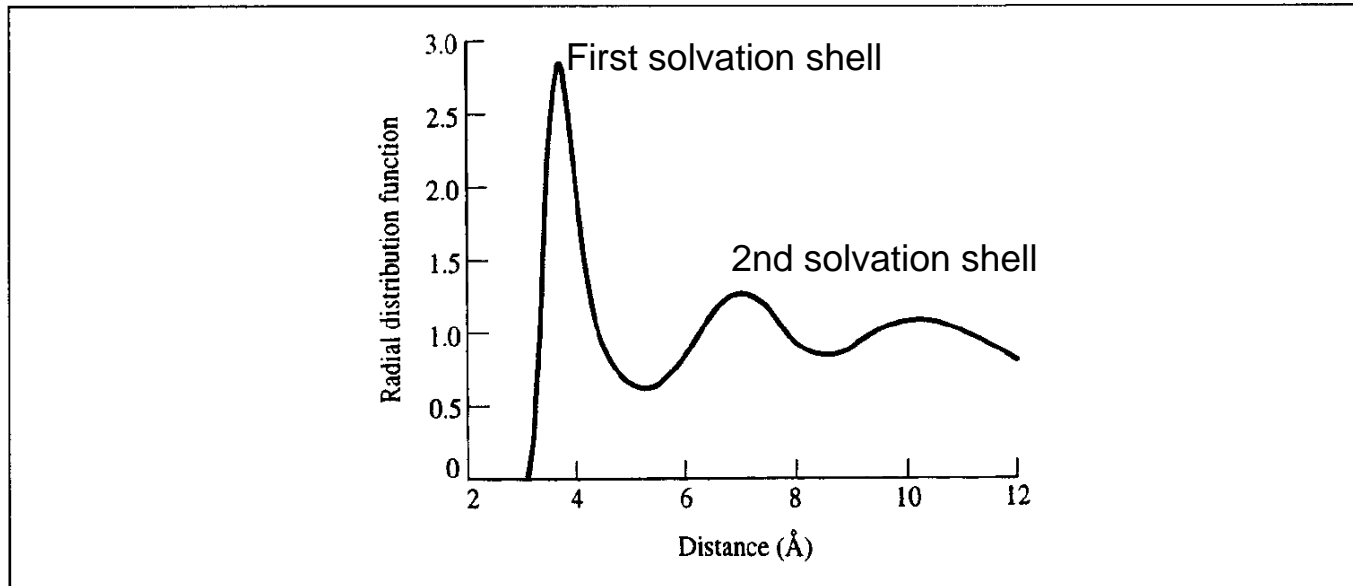


Fig. 6.2: Radial distribution function determined from a 100 ps molecular dynamics simulation of liquid argon at a temperature of 100 K and a density of  $1.396 \text{ g cm}^{-3}$ .

Experimentally, radial distribution functions can be measured using X-ray diffraction. So, experimental and theoretical results can be compared to know the quality of a simulation

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