*Applied Statistical Mechanics Lecture Note - 12*



## **Advanced Topics in Molecular Monte Carlo Method**

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#### **Contents**



- I. Free Energy Calculations
- II. Gibbs Ensemble Method
- III. Biased Sampling Method

#### **Introduction**



- **Previous simulation methods: Properties of a single** isolated phase
- Molecular Simulation Techniques can be extended to multiple phases
	- **Thermodynamic Phase Coexistence : Two or more phases are** equally stable

**Condition of Phase Coexistence** 

$$
T^{\alpha} = T^{\beta}
$$
  
\n
$$
p^{\alpha} = p^{\beta}
$$
  
\n
$$
\mu_i^{\alpha} = \mu_i^{\beta} \qquad i = 1, ..., C
$$



### **Straightforward Simulation**



- ■ Change T or P for given model system and wait for a phase transformation occur
	- $\Box$  Major Drawbacks
		- Hysterisis : Irreversible phase transformation
		- Large free energy barrier at interface  $\rightarrow$  Depends on the size of the interface  $\rightarrow$  Depends on the choice of simulation system



# **EG 59** स्रध **Alternative Methods** 고려대학교 Gibbs Ensemble Method (Panagiotopoulos, 1987)  $\Box$  Gibbs – Duhem Integration Method (Kofke, 1993)  $\Box$

## **Why free energies are important in Phase Coexistence ?**



 From 2nd Law of Thermodynamics  $\Box$  At equilibrium, S is maximum for  $(N, V, E)$  $\Box$  (N,V,E) System : A is minimum  $A = U - TS$  (Helmholtz Free Energy)  $\Box$  (N,P,T) System : G is Minimum *G = H – TS* (*Gibbs Free Energy*) Equilibrium Condition

$$
A^{\alpha} = A^{\beta}
$$

$$
G^{\alpha} = G^{\beta}
$$

## **Free energy cannot be directly measured in simulation**



■ Helmholtz free energy

$$
A = -kT \ln Q(N, V, T) = -kT \left( \frac{\int dp^N dr^N \exp(-\beta H(p^N, r^N))}{\Lambda^{dN} N!} \right)
$$

 $\Box$  Not the form of canonical average over phase space  $\Box$  Depends directly on the available volume in the phase space  $\Box$ Cannot be directly measured in real experiment, too.

#### **Derivatives of free energy**

$$
\left(\frac{\partial A}{\partial V}\right)_{N,T} = -p \qquad \left(\frac{\partial A/T}{\partial 1/T}\right)_{V,T} = -E
$$

**Find a reversible path in V,T plane and perform integration**

## **Kirkwood's coupling parameter method**



٠ **Assume U** depends on the coupling parameter  $\lambda$  :

 $U(\lambda) = (1 - \lambda)U_I + \lambda U_{II} = U_I + \lambda (U_{II} - U_I)$ 

■ Partition function for potential energy function

$$
Q(N, V, T, \lambda) = \frac{1}{\Lambda^{3N} N!} \int dr^N \exp(-\beta U(\lambda))
$$

$$
\left(\frac{\partial A(\lambda)}{\partial \lambda}\right)_{N,V,T} = -\frac{1}{\beta} \frac{\partial}{\partial \lambda} \ln Q(N,V,T,\lambda) = -\frac{1}{\beta Q(N,V,T,\lambda)} \frac{\partial Q(N,V,T,\lambda)}{\partial \lambda}
$$

$$
= \frac{\int dr^N \left(\frac{\partial U(\lambda)}{\partial \lambda}\right) \exp(-\beta U(\lambda))}{\int dr^N \exp(-\beta U(\lambda))} = \left\langle \frac{\partial U(\lambda)}{\partial \lambda} \right\rangle_{\lambda} \longrightarrow \text{Ensemble average of with potential function } U(\lambda)
$$

## **Kirkwood's coupling parameter method**



**Firm Free energy difference** 

$$
A(\lambda = 1) - A(\lambda = 0) = \int_{\lambda = 0}^{\lambda = 1} d\lambda \left\langle \frac{\partial U(\lambda)}{\partial \lambda} \right\rangle
$$

- All free energy methods are based on calculation of free energy differences
- $\blacksquare$  Example :
	- $\Box$  Volume of R can be measured as a fraction of total volume
		- Sample the reference system
		- keep an average of the fraction of time occupying the target system

$$
\frac{\Omega_R}{\Omega_{\Gamma}} = \langle s(\Gamma) \rangle_{\Gamma} \qquad S_R - S_{\Gamma} = k \ln \left( \Omega_R / \Omega_{\Gamma} \right)
$$

### **Chemical Potentials**



#### **E** Chemical Potentials

$$
\mu_a = \left(\frac{\partial G}{\partial N_a}\right)_{T,P,N_{b\neq a}} = \left(\frac{\partial A}{\partial N_a}\right)_{T,V,N_{b\neq a}} = -T\left(\frac{\partial S}{\partial N_a}\right)_{V,E,N_{b\neq a}}
$$

$$
Q(N, V, T) = \frac{V^N}{\Lambda^{dN} N!} \int_0^1 \dots \int_0^1 ds^N \exp(-\beta U(s^N; L))
$$
  

$$
s^N = r^N / L
$$

$$
A(N, V, T) = -kT \ln Q = -kT \ln \left[ \frac{V^N}{\Lambda^{dN} N!} \right] - kT \ln \left\{ \int_0^1 ... \int_0^1 ds^N \exp(-\beta U(s^N; L)) \right\}
$$
  
=  $A_{id}(N, V, T) + A_{ex}(N, V, T)$ 



## **Widom's Test Particle Insertion Method**



- Widom (1963)
- A "ghost particle" is randomly inserted into the ensemble and calculating the energy of its interaction
- The test particle is a "ghost", it does not affect the properties of real molecule
- In principle , this method can be used in any simulation system for the calculation of chemical potential

## **Implementation of Widom's Method**



- Carry out conventional NVT or NPT Monte Carlo Simulation of N particles
- At frequent interval during simulation, randomly generate a coordinate,  $s_{N+1}$  uniformly over unit cube
- For given  $s_{N+1}$ , compute :

$$
\mu_{ex} = -kT \ln \langle \exp(-\beta \Delta U) \rangle \longrightarrow \text{NVT ensemble}
$$

$$
\mu_{ex} = -kT \ln \left[ \frac{\langle V \exp(-\beta \Delta U) \rangle}{\langle V \rangle} \right] \longrightarrow \text{NPT ensemble}
$$

 $\blacksquare$  Average  $\rightarrow$  chemical potential

#### **Note on Widom's Method**



- Widom's method can be used as a verification that equilibrium has been attained
- At high density, some difficulties are encountered  $\Box$ Difficult to insert a particle at given location



- Proposed by Panagiotopoulos (1987,1988)
- Simulation method without interface
- Thermodynamic contact without physical contact





٠ MC simulation includes moves that couples two simulation volumes

> Particle exchange equilibrates chemical potential

Volume exchange equilibrates pressure



∩

Incidentally, the coupled moves enforce mass and volume balance





Box 1:  $N_1$ ,  $V_1$ , T



Box 2:  $N_2$ ,  $V_2$ , T



(2) Attempt volume fluctuation

Box 1:  $N_1$ ,  $V_1 + \Delta V$ , T



Box 1:  $N_1 + 1$ ,  $V_1$ , T











Acceptance of three moves  $\rightarrow$  Governed by Pseudo Boltzman Factor

Displacement movement

$$
\Delta Y_{disp} = \Delta E_{\alpha} + \Delta E_{\beta}
$$

**U** Volume change

$$
\Delta Y_{disp} = \Delta E_{\alpha} + \Delta E_{\beta} - N_{\alpha} kT \ln \frac{V_{\alpha} + \Delta V_{\alpha}}{V_{\alpha}} - N_{\beta} kT \ln \frac{V_{\beta} + \Delta V_{\beta}}{V_{\beta}} + P(\Delta V_{\alpha} + \Delta V_{\beta})
$$

**Nolecular transfer** 

$$
\Delta Y_{disp} = \Delta E_{\alpha} + \Delta E_{\beta} - kT \ln \frac{V_{\beta}(N_{\alpha}+1)}{V_{\alpha}N_{\beta}}
$$

## **Gibbs Ensemble - Algorithm**





### **DO 60** 四印 **Gibbs Ensemble – Result**  고려대학교  $\blacksquare$  Water + Methanol Mixture

# $\circ$   $\circ \bullet$   $\circ$  $O^{\circ}$  $\Omega$  $\circ$  $\circ$  $\circ$

•*Strauch and Cummings, Fluid Phase Equilibria, 86 (1993) 147-172;* • *Chialvo and Cummings, Molecular Simulation, 11 (1993) 163-175.* 

#### **Gibbs Ensemble – Result**



#### ■ Panagiotopoulos Group



#### **Gibbs Ensemble Limitation**



#### ■ Limitation arise from particle-exchange requirements



**Dense phase or Complex molecules**



**Solid phases**

### **Gibbs-Duhem Integration**



- $\Box$ David Kofke (1993)
- Basis : Numerical Integration of Cluisus Clapeyron Equation

$$
\mu_{\alpha} - \mu_{\beta} = -(S_{\alpha} - S_{\beta})dT - (V_{\alpha} - V_{\beta})dP
$$

$$
\frac{dP}{dT} = \frac{S_{\alpha} - S_{\beta}}{V_{\alpha} - V_{\beta}} = \frac{\Delta H}{T\Delta V}
$$

$$
\frac{d \ln P}{d1/T} = -\frac{\Delta H}{P\Delta V/T} \qquad \longrightarrow \qquad \left(\frac{\partial \ln p}{\partial \beta}\right)_{\sigma} = -\frac{\Delta h}{\Delta Z}
$$

#### **Gibbs – Duhem Integration**



GE equation

$$
\left(\frac{\partial \ln p}{\partial \beta}\right)_{\sigma} = -\frac{\Delta h}{\Delta Z}
$$

n Treat as nonlinear first order ODE  $\Box$  Use (NPT) simulation to obtain  $\Delta H / \Delta z$ 



## **Predictor-Corrector Algorithm Implementation**

Given initial condition and slope  $(=-\Delta h/\Delta Z)$ , predict new (p,T) pair.

Evaluate slope at new state condition…

 …and use to correct estimate of new (p,T) pair



### **Gibbs Duhem Integration**



- ٠ Potentially very efficient algorithm
	- □ Coexistence curves for solid-liquid systems
	- $\Box$ Coexistence curves for complex molecules
- Algorithm is not robust
- No built-in diagnostics
- Additional free energy calculation may be required to check the result