Modelowanie Nanostruktur



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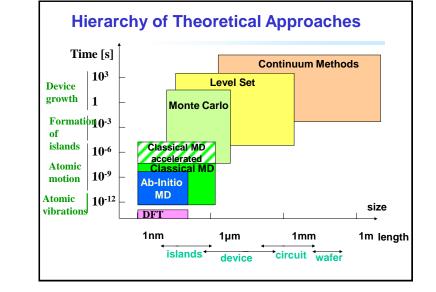
Semester Zimowy 2011/2012

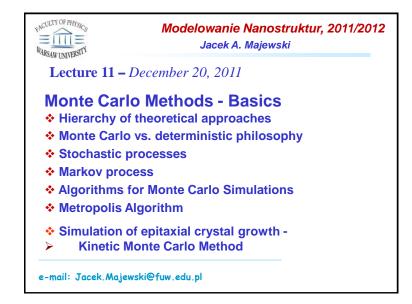
Wykład

Modelowanie Nanostruktur

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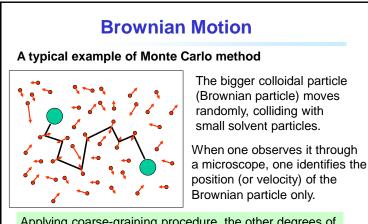
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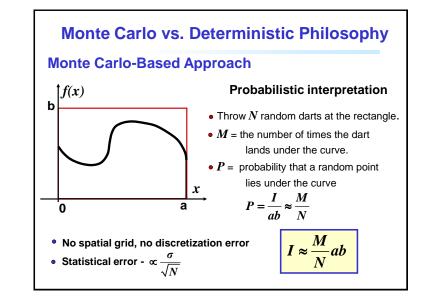


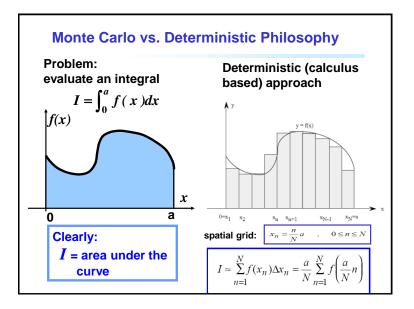
Basics of the Monte Carlo Method

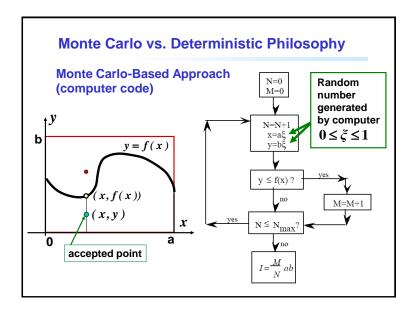
- In a macroscopic system, it is difficult to treat the motions of the all (microscopic) atoms or molecules
- Macroscopic properties of a systems (i.e., how the whole system behaves) are of interest
 - ➡ Coarse-graining necessary
- If the time evolution of the system is coarse-grained stochastically, one achieves one class of models, so-called stochastic models.
- Monte Carlo Method efficient method to realize this numerically on a computer
- Monte Carlo methods provide a powerful way to solve numerically the fluctuation or relaxation in a stochastic system

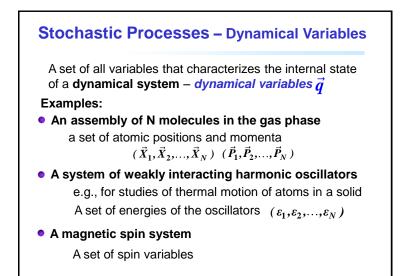


Applying coarse-graining procedure, the other degrees of freedom (e.g., the motion of small solvent particles) are removed and, finally they can be regarded as a **random force acting on the Brownian particle.**









Stochastic Processes – Dynamical Variables

Studies of the time evolution of the dynamic variables

- Suppose we have obtained successive data for the dynamical variables *q*₁, *q*₂,...,*q*_L by observing the system *L* times at different time steps *t*₁, *t*₂,...,*t*_L
- If these successive data seem to change stochastically with time step, we may call this a *stochastic process*.
- In the Monte Carlo methods which are used to the dynamics of a many-body system,
 - the time evolution is considered as a stochastic process,
 - the dynamical variables at each time step are updated by using random numbers.
- A sample obtained in one simulation $\vec{q}_1, \vec{q}_2, ..., \vec{q}_L$ The real Monte Carlo - simulations repeated many times and analyzed using standards of statistical physics

Stochastic Processes – Dynamical Variables at Equilibrium

One of the most important subjects in the *Monte Carlo* method distribution of dynamical variables at thermal equilibrium of the system ?

- It is not necessary to examine the trajectory of the dynamical variable according to some deterministic equation.
- It is important to discuss the value of the dynamical variable at a certain place and a certain time

Stochastic Processes – Distribution Function

• **Transition probability** that the system with a dynamic variable q_i at time t_i moves to the state between $q_i + \delta q_i$ and q_j at a later time t_i

$$\varPhi(q_i,t_i \,|\, q_j,t_j) \delta q_j$$

• *Distribution function of q* - probability p(q,t) that the system is in the state defined by q at time t

$$p(q,t) = \int dq_0 \ p(q_0,t_0) \Phi(q_0,t_0 | q,t)$$

• Transition probability should be normalized

$$\int \Phi(q_i, t_i | q_j, t_j) \, dq_j = 1$$

for any state q_i at time $t_i < t_j$



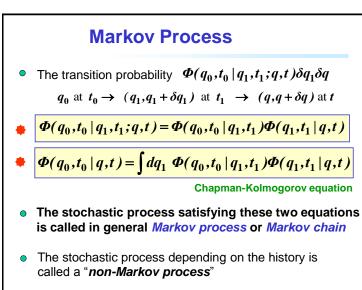
 Most algorithms used in simulating a realistic system by Monte Carlo methods, are based on the following Markov process.

• $\Phi(q_0, t_0 | q, t)$ is independent of any information about any time *t*' before t_0

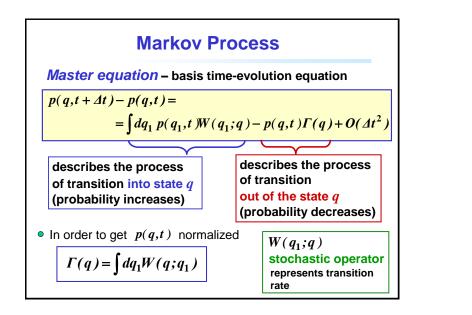
• All the history before time t_0 is contracted into the single piece of information that the system has the dynamic variable q_0 at the time t_0 .

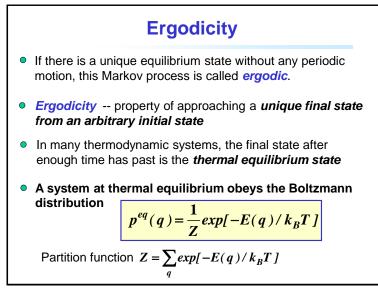
Markov Process

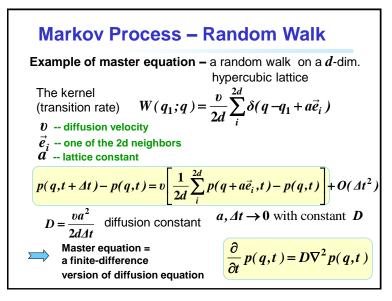
- By introducing a kind of random updating of the dynamical variable in compensation for neglecting the microscopic details of the real system, one may arrive at a Markov process.
- Replacement of the neglected details with a random updating process is identical to the introduction of a *heat bath*
- Detailed motion of atoms on the microscopic scale can be seen as heat on the macroscopic scale.
- The form and amount of the updating probability are directly related to the temperature of the equilibrium state.



Markov ProcessMarkov Processnew variables $\Gamma(q)$ and $W(q_1;q)$ Transition amplitude $\Phi(q_1,t|q,t+\Delta t) = [1-\Gamma(q)]\delta(q-q_1) + W(q_1;q) + O(\Delta t^2)$ Change of the Transition amplitude $\Phi(q_0,t_0|q,t+\Delta t) - \Phi(q_0,t_0|q,t) =$ $= \int dq_1 \Phi(q_0,t_0|q_1,t) W(q_1;q) - \Phi(q_0,t_0|q,t) \Gamma(q) + O(\Delta t^2)$ Change of the distribution function $p(q,t+\Delta t) - p(q,t) =$







Algoritms for Monte Carlo Simulations

The most basic algorithm of the Monte Carlo method:

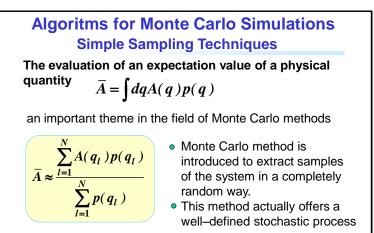
- (1) Generate a random number
- (2) Take or do not take a new random step, depending on the generated random number
- (3) Repeat trial

Random numbers

- The "random numbers" generated on computer are not mathematically ideal random numbers
- **pseudo-random numbers** uniformly distributed numbers in the interval [0,1] having long but finite period
- For 32-bit processor,

the **period** is $M = 2^{31} - 1 = 2147483647$

Modelowanie Nanostruktur



Such a method of Monte Carlo sampling is called a **Simple Sampling Technique**

Algoritms for Monte Carlo Simulations

Simple Sampling Technique for thermodynamic ensamble

The one-particle energy distribution function $p(\varepsilon)$ shows a peak of height \sqrt{N} and width $1/\sqrt{N}$ around its expectation value $\langle E \rangle / N$

$$p(\varepsilon) \propto exp\left[-\frac{(\varepsilon - \langle E \rangle / N)^2}{2Ck_B T^2}N\right]$$
 Note, Gaussian form
for large N

Probability to generate states with one particle energy close to < E > / N becomes exponentially small for large N

Simple Sampling Algorithm becomes very inefficient

Algoritms for Monte Carlo Simulations Simple Sampling Technique

Handling a thermodynamic ensemble based on a stochastic model by the simple sampling technique

$$\overline{A} \approx \frac{\sum_{l=1}^{N} A(q_l) exp[-E(q)/k_B T]}{\sum_{l=1}^{N} exp[-E(q)/k_B T]}$$

Note, all samples *l* are selected here completely randomly, irrespective of the Boltzmann weight

Now assume $A(q_l) = E(q_l)$

The relative width of the energy fluctuations decreases with the number of particles N, in the system

$$\frac{\langle E^2 \rangle - \langle E \rangle^2}{\langle E^2 \rangle} \propto \frac{1}{N}$$

Algoritms for Monte Carlo Simulations Importance Sampling Technique

- Simple Sampling Technique offers an efficient algorithm to evaluate the average of a physical quantity A(q), only if the distribution function p(q) resembles a more or less uniform distribution
- In contrast, if the distribution function has a big value only at an isolated point, it becomes more efficient to choose the integration points with the same probability as that given by the distribution function p(q).
- Such a biased sampling with a probabilistic weight is called *Importance Sampling Technique* N. Metropolis, A.W. Rosenbluth, M.N. Rosenbluth, A. M. Teller, and E. Teller, J. Chem. Phys. 21, 1087 (1953).
 - Commonly called Metropolis Algorithm

Algoritms for Monte Carlo Simulations Importance Sampling Technique

• If the sample integration points are chosen with the same probability as the distribution function p(q), the expression for the expectation value can be replaced simply by -1 $\frac{N}{2}$

$$\overline{A} = \frac{1}{N} \sum_{l=1}^{N} A(q_l)$$

• **Metropolis et al.** considered a Markov process that generates a descendant sample *l*' from the present sample *l* according to the transition rate $W(q_l, q_{l'})$

They showed that by suitable choice of the transition rate, it is possible to bring the distribution of the dynamic variable to the expected (prescribed) distribution p(q) in the limit $N \rightarrow \infty$

Algoritms for Monte Carlo Simulations Importance Sampling Technique

- In IST one updates the particle position with the probability min[*x*,1] $x = \frac{p(q')}{p(q)}$
- After repeating this process enough times, the distribution of this random walker approaches p(q)

Algoritms for Monte Carlo Simulations Importance Sampling Technique

Markov chain

We consider one particle in the multidimensional space of the dynamic variable q and move it step by step according to random numbers. $q \rightarrow q'$

- The movement obeys following rule
 - Calculate the value of the distribution function p(q')
 - Compare p(q') to the present value of the distribution function p(q)
 - If p(q') > p(q) move the particle to the new position
 - If p(q') < p(q) generate random number $\xi \in [0,1]$
 - If $\xi < \frac{p(q')}{p(q)}$ move the particle to the new position
 - If $\xi \ge \frac{p(q')}{p(q)}$ leave the particle at the same position

Algoritms for Monte Carlo Simulations Importance Sampling Technique

Back to the specific problem of the thermodynamic ensembles

• We assume that the samples are chosen with the same probability as the distribution function $p(q_l)$

$$\overline{A} \approx \frac{\sum_{l=1}^{N} A(q_l) exp[-E(q)/k_B T] / p(q_l)}{\sum_{l=1}^{N} exp[-E(q)/k_B T] / p(q_l)}$$

- How to choose the distribution function $p(q_l)$?
- The simplest and most natural choice is the Boltzmann distribution $p(q_1) \propto exp[-E(q)/k_BT]$



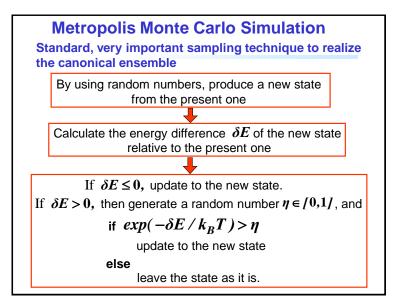
• Using Metropolis's idea of importance sampling, one may bring the distribution function $p(q_l)$ close to the thermal-equilibrium distribution

$$p(q_l) = \frac{1}{Z} exp[-E(q)/k_BT]$$

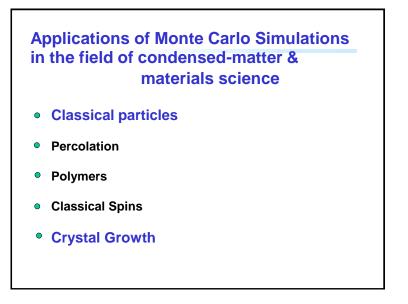
• A sufficient condition to achieve the equilibrium distribution is to impose *the principle of detailed balance*

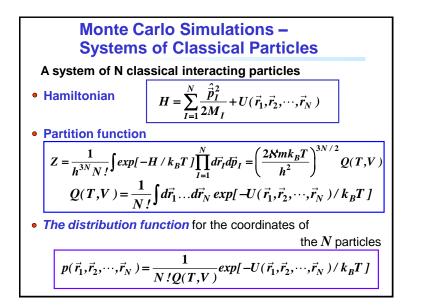
$$p^{eq}(q_l)W(q_l,q_{l'}) = p^{eq}(q_{l'})W(q_{l'},q_{l'})$$

However, the principle of detailed balance does not uniquely determine the transition rate $W(q_I, q_{I'})$



Algoritms for Monte Carlo Simulations Importance Sampling Technique Two often used choices of transition rate • Heat bath method (Glauber algorithm) R. J. Glauber, J. Math. Phys. 4, 294 (1963) $W(q_l,q_{l'}) = \frac{1}{\tau} \left(1 - tanh \left(\frac{\delta E}{2k_B T} \right) \right) = \frac{1}{\tau} \frac{exp(-\delta E / k_B T)}{[1 + exp(-\delta E / k_B T)]}$ $\delta E = E(q_{l'}) - E(q_l)$ • Metropolis Algorithm $W(q_l,q_{l'}) = \begin{cases} \frac{1}{\tau} exp(-\delta E / k_B T) & \delta E > 0\\ \frac{1}{\tau} & \text{otherwise} \end{cases}$







Procedure

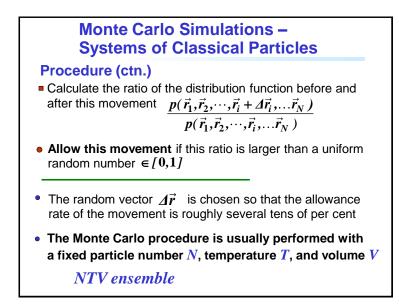
- One chooses one particle among N particles randomly (or one may select them one by one from 1 to N)
- Move the position $\vec{r_i}$ of the selected particle to the new position $\vec{r_i} + \Delta \vec{r_i}$ $\vec{r_i} \rightarrow \vec{r_i} + \Delta \vec{r}$

a random vector

Metropolis algorithm

Check whether the new configuration is energetically more stable than the original configuration

- If YES allow this movement
- If NO further procedure



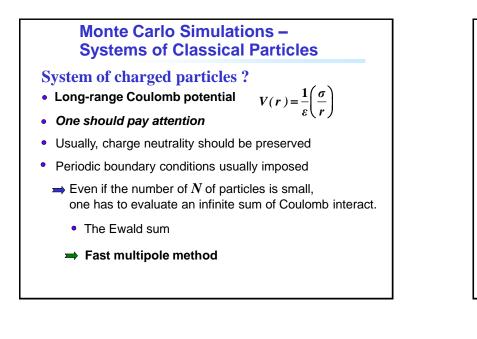
Monte Carlo Simulations – Systems of Classical Particles

Form of potential U?

- The simplest one "Hard sphere system" liquid – solid phase transition
 - B. J. Alder & T. E. Wainwright, J. Chem. Phys. **27**, 1208 (1957) **Molecular dynamics**
 - W.W. Wood & J.D. Jacobson, J. Chem. Phys. 27, 1207 (1957)

Monte Carlo Simulations

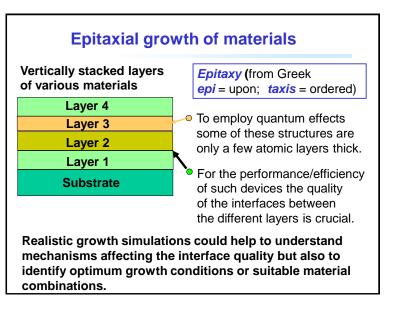
- Soft-core, pair potentials $V(r) = \frac{1}{\varepsilon} \left(\frac{\sigma}{r} \right)^n$
- Lennard-Jones potential $V_{ij}(R_{ij}) = U_0 \left[\left(\frac{R_0}{R_{ij}} \right)^{12} 2 \left(\frac{R_0}{R_{ij}} \right)^6 \right]$

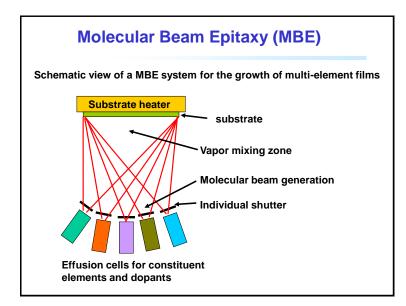


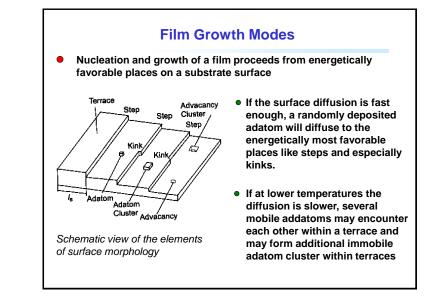
Kinetic Monte Carlo Simulations – An approach to perform epitaxial growth simulations

- *Epitaxial growth* is a key technique in fabricating semiconductor-based electronic and optoelectronic devices such as
 - light-emitting diodes (LED's),
 - laser diodes (LDs), or
 - high electron mobility transistors.
- These devices consist of vertically stacked thin films that differ by the material, alloy composition, or doping.

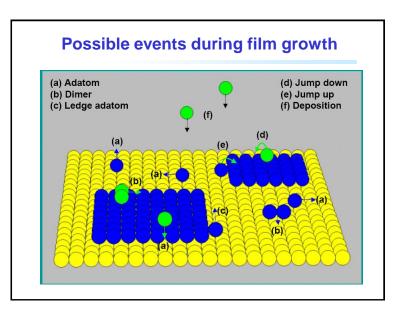












Lecture 11

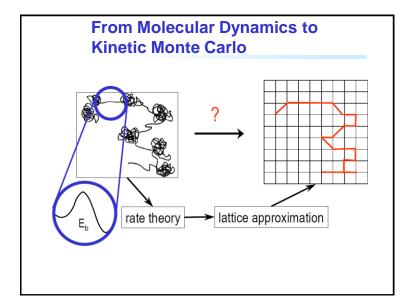


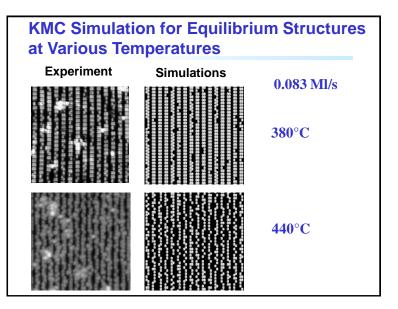
- A challenge to perform such growth simulations is the large range of relevant length and time scales.
- The features interesting for device design (interface morphology, formation of nanostructures) are of the order of 100–1000 nm and the time to grow these structures is of the order of seconds.
- *The origin of these effects*, however, lies in the atomic processes on the surface (adatom adsorption, desorption, nucleation,etc.).

This requires a resolution in the *length scale* 0.1 nm and in the *time scale* of 10^{-13} s.

Simulation of growth processes – Kinetic Monte Carlo (KMC)

- Modeling crystal growth with the KMC method allows one to cover experimentally relevant growth times and system sizes, since each event on the surface is just described by a single quantity—the transition rate rather than by modeling the full reaction path including atomic geometries and energies
 - ⇒ Bridging of length and time scales





Modelowanie Nanostruktur

