1.021, 3.021, 10.333, 22.00 Introduction to Modeling and Simulation Spring 2011

Part I – Continuum and particle methods

How to model chemical interactions

Lecture 5

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Content overview

I. Particle and continuum methods

Lectures 1-13

- 1. Atoms, molecules, chemistry
- 2. Continuum modeling approaches and solution approaches
- Statistical mechanics
- 4. Molecular dynamics, Monte Carlo
- 5. Visualization and data analysis
- 6. Mechanical properties application: how things fail (and how to prevent it)
- 7. Multi-scale modeling paradigm
- 8. Biological systems (simulation in biophysics) how proteins work and how to model them

II. Quantum mechanical methods

- Lectures 14-26
- 1. It's A Quantum World: The Theory of Quantum Mechanics
- 2. Quantum Mechanics: Practice Makes Perfect
- 3. The Many-Body Problem: From Many-Body to Single-Particle
- 4. Quantum modeling of materials
- 5. From Atoms to Solids
- 6. Basic properties of materials
- 7. Advanced properties of materials
- 8. What else can we do?

Overview: Material covered so far...

- Lecture 1: Broad introduction to IM/S
- Lecture 2: Introduction to atomistic and continuum modeling (multi-scale modeling paradigm, difference between continuum and atomistic approach, case study: diffusion)
- Lecture 3: Basic statistical mechanics property calculation I (property calculation: microscopic states vs. macroscopic properties, ensembles, probability density and partition function)
- Lecture 4: Property calculation II (advanced property calculation, introduction to chemical interactions, Monte Carlo method)
- Lecture 5: How to model chemical interactions

Lecture 5: How to model chemical interactions

Outline:

- 1. Monte-Carlo (MC) approach: Metropolis-Hastings algorithm
- 2. How to model chemical interactions
 - 2.1 Pair potentials
 - 2.2 How to model metals: Multi-body potentials

Goals of today's lecture:

- Get to know basic methods to model chemical bonds (starting with simple "pair potentials")
- Learn how to identify parameters for models of chemical bonds (for pair potentials)
- Limitations of pair potentials and other, alternative methods

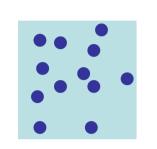
1. Monte-Carlo (MC) approach: Metropolis-Hastings algorithm

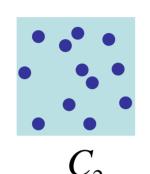
Averaging over the ensemble

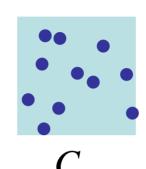
Property A₁



Property A₃

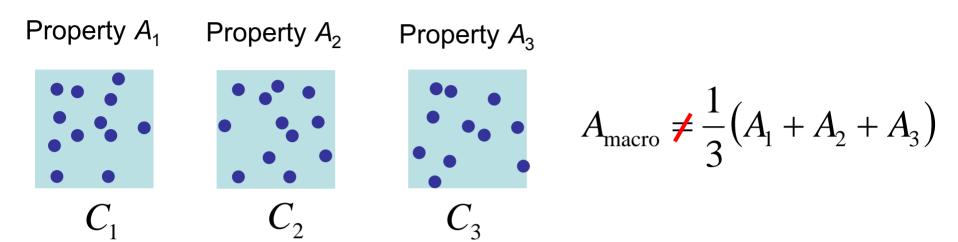






$$A_{\text{macro}} \neq \frac{1}{3} \left(A_1 + A_2 + A_3 \right)$$

Averaging over the ensemble



Instead, we must average with proper weights that represent the probability of a system in a particular microscopic state!

(I.e., not all microscopic states are equal)

$$A_{\text{macro}} = \rho_1 A_1 + \rho_2 A_2 + \rho_3 A_3 =$$

$$\rho_1(r_1, p_1) A_1(r_1, p_1) + \rho_2(r_2, p_2) A_2(r_2, p_2) + \rho_3(r_3, p_3) A_3(r_3, p_3)$$

Probability to find system in state C_1

How to solve...

$$\langle A \rangle = \iint_{p} A(p,r) \rho(p,r) dr dp$$

Probability density distribution

E.g.:
$$T(p) = \frac{1}{3} \frac{1}{Nk_B} \sum_{i=1}^{N} \frac{m_i^2 \vec{v}_i^2}{m_i} = A(p)$$

Virtually impossible to carry out analytically

Must know all possible configurations

Therefore: Require numerical simulation

Molecular dynamics OR Monte Carlo

Monte Carlo scheme

Method to carry out integration over "domain"

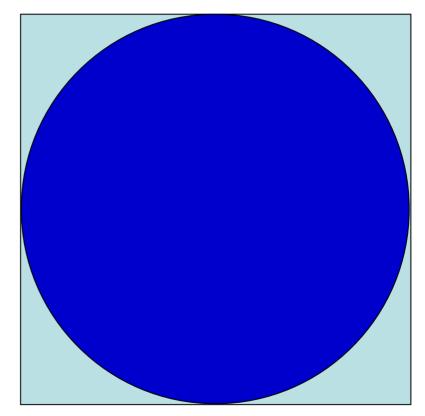
Want:

$$A = \int_{\Omega} f(\vec{x}) d\Omega$$

E.g.: Area of circle (= $\pi/4$ exact solution)

$$A_C = \frac{\pi d^2}{4} \qquad A_C = \frac{\pi}{4}$$

$$\pi = 4A_C$$

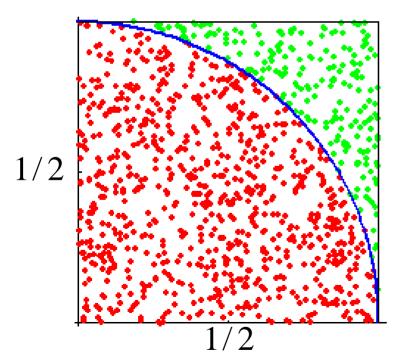


$$d = 1$$

$$f(\vec{x}) = \begin{cases} 1 & \text{inside} \\ 0 & \text{outside} \end{cases}$$

Monte Carlo scheme for integration

- **Step 1**: Pick random point \vec{x}_i in Ω
- Step 2: Accept/reject point based on criterion (e.g. if inside or outside of circle and if in area not yet counted)
- Step 3: If accepted, add $f(\vec{x}_i) = 1$ to the total sum



$$A_{C} = \int_{\Omega} f(\vec{x}) d\Omega \qquad A_{C} = \frac{\pi}{16}$$

$$N_{A}: \text{ Attenmade}$$

$$A_{C} = \frac{1}{N} \sum_{i} f(\vec{x}_{i})$$

$$A_C = \frac{\pi}{16}$$

 N_{A} : Attempts made

$$A_C = \frac{1}{N_A} \sum_{i} f(\vec{x}_i)$$

How to apply to ensemble average?

 Similar method can be used to apply to integrate the ensemble average

$$< A >= \iint_{p} A(p,r)\rho(p,r)drdp$$

$$\rho(p,r) = \frac{1}{Q} \exp\left[-\frac{H(p,r)}{k_B T}\right]$$

"discrete"

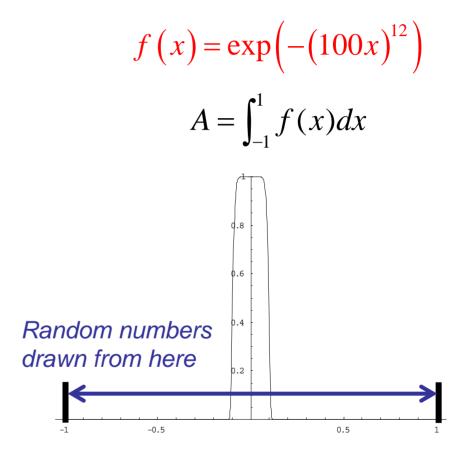
$$< A > = \sum_{i=1}^{N_A} \frac{A \exp(-H(r_A, p_A)/(k_B T))}{\sum_{i=1}^{N_A} \exp(-H(r_A, p_A)/(k_B T))}$$

Computationally inefficient: If states are created "randomly" that have low probability....

 To be computationally more effective, need more complex iteration scheme (replace "random sampling" by "importance sampling")

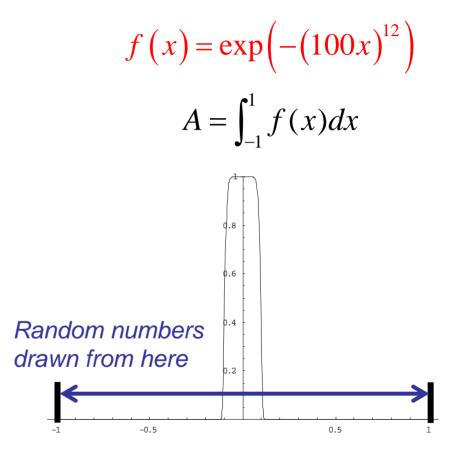
Challenge: sampling specific types of distributions

- We want to
 - Integrate a sharply-peaked function
 - Use Monte Carlo with uniformly-distributed random numbers (e.g. here from -1 to 1)



Challenge: sampling specific types of distributions

- We want to
 - Integrate a sharply-peaked function
 - Use Monte Carlo with uniformly-distributed random numbers (e.g. here from -1 to 1)
- What happens?
 - Very few points contribute to the integral (~9%)
 - Poor computational efficiency/convergence
- Solution: use a different distribution of random numbers to sample "importance sampling"



Importance sampling

 Core concept: Picking states with a biased probability: Importance sampling (sampling the "correct" way...)

$$\langle A \rangle = \iint_{p} A(p,r)\rho(p,r)drdp$$
 $\rho(p,r) = \frac{1}{Q} \exp \left[-\frac{H(p,r)}{k_B T} \right]$

Importance sampling

 Core concept: Picking states with a biased probability: Importance sampling (sampling the "correct" way...)

$$\langle A \rangle = \iint_{p} A(p,r)\rho(p,r)drdp$$
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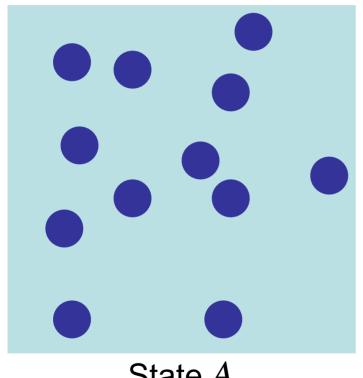
Notice: Probability (and thus importance) related to energy of state

Importance sampling: Metropolis algorithm

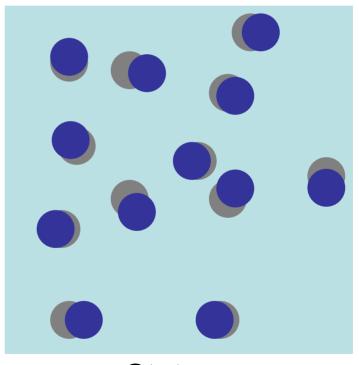
 Leads to an appropriate "chain" of states, visiting each state with correct probability

- Concept:
 - Pick random initial state
 - Move to trial states
 - Accept trial state with certain probability (based on knowledge about behavior of system, i.e., energy states)

Concept: Generate set of random microscopic configurations Accept or reject with certain scheme



State A



State B

Random move to new state B

Have: State A (initial state) + energy function H(A)

Step 1: Generate new state B (random move)

Have: State A (initial state) + energy function H(A)

Step 1: Generate new state B (random move)

Step 2: if H(B) < H(A) then a = 1

else

a = true[1]/false[0]
for acceptance

Draw random number 0

"Downhill" moves always accepted

Have: State A (initial state) + energy function H(A)

Step 1: Generate new state B (random move)

Step 2: if H(B) < H(A) then a = 1 else

a = true[1]/false[0]
for acceptance

Draw random number 0

"Downhill" moves always accepted, uphill moves with finite ("thermal") probability

if
$$p < \exp\left[-\frac{H(B) - H(A)}{k_B T}\right]$$
 $a = I$

$$a = 0$$

endif

a=variable either 0 or 1

Have: State A (initial state) + energy function H(A)

Step 1: Generate new state B (random move)

Step 2: if
$$H(B) < H(A)$$
 then $a = 1$ else

a = true[1]/false[0]
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Draw random number 0

if
$$p < \exp\left[-\frac{H(B) - H(A)}{k_B T}\right]$$
 $a = 1$

$$a = 0$$

endif

endif

a=variable either 0 or 1 (used to detect acceptance of state *B* when *a*=1)

Step 3: if a = 1 then accept state B endif

Have: State
$$A$$
 (initial state) + energy function $H(A)$

Step 1: Generate new state B (random move)

Step 2: if
$$H(B) < H(A)$$
 then $a = 1$

else

repeat N_A times

a = true[1]/false[0]for acceptance

Draw random number 0

Draw random number
$$0$$

if
$$p < \exp\left[-\frac{H(B) - H(A)}{k_B T}\right]$$
 $a = 1$

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endif

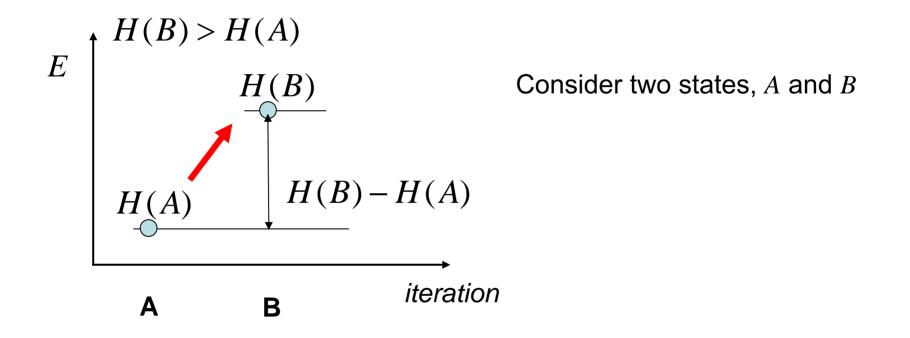
a=variable either 0 or 1 (used to detect acceptance of state B when a=1)

endif

Step 3: if (a = 1) then accept state Bendif

$$< A > = \frac{1}{N_A} \sum_{i=1..N_A} A(i)$$

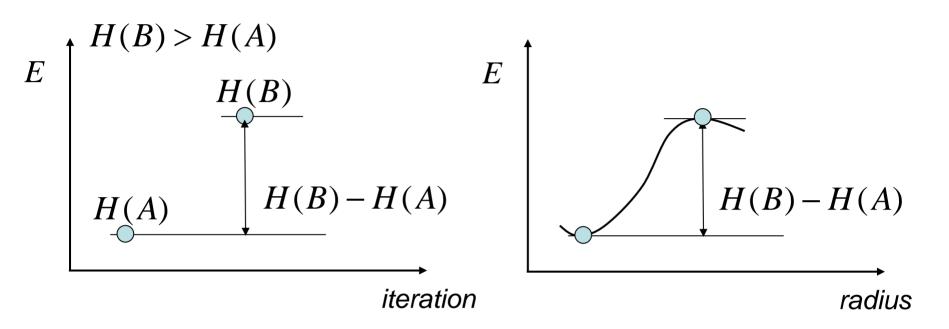
Arrhenius law - explanation



State B has higher energy than state A

Otherwise accepted anyway!

Arrhenius law - explanation



Energy difference between states *A* and *B* ("uphill")

Probability of success of overcoming the barrier at temperature
$$T$$
 $\exp\left[-\frac{H(B)-H(A)}{k_BT}\right]$

Arrhenius law - explanation

Probability
of success
of overcoming the
barrier

$$\exp\left[-\frac{H(B) - H(A)}{k_B T}\right]$$

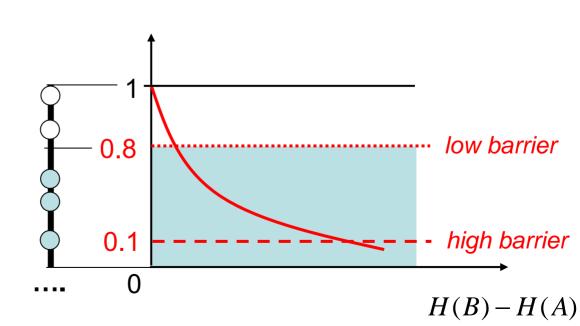
Random number 0

(equal probability to draw any number between 0 and 1)

Acceptance if:

$$p < \exp\left[-\frac{H(B) - H(A)}{k_B T}\right]$$

E.g. when $\exp(..) = 0.8$ most choices for p will be below, that is, higher chance for acceptance



Play "1D darts"

Summary: Metropolis-Hastings Algorithm

Have: State
$$A$$
 (initial state) + energy function $H(A)$

Step 1: Generate new state B (random move)

Step 2: if
$$H(B) < H(A)$$
 then $a = 1$

else

repeat N_A times

a = true[1]/false[0]
for acceptance

Draw random number 0

if
$$p < \exp \left[-\frac{H(B) - H(A)}{k_B T} \right]$$
 $a = D$

else

a = 0

endif

a=variable either 0 or 1 (used to detect acceptance of state B when a=1)

endif

Step 3: if (a = 1) then accept state B endif

 $< A > = \frac{1}{N_A} \sum_{i=1..N_A} A(i)$

Summary: MC scheme

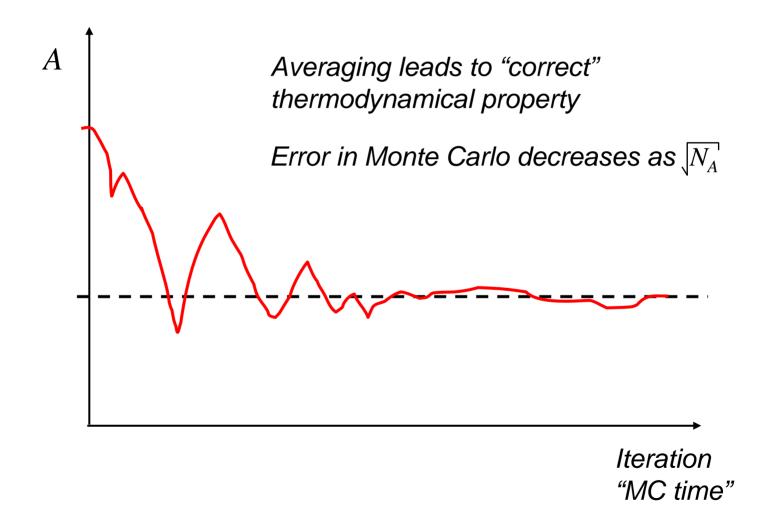
Have achieved:

$$\langle A \rangle = \iint_{p} A(p,r)\rho(p,r)drdp$$
 \longleftrightarrow $\langle A \rangle \frac{1}{N_A} \sum_{i=1..N_A} A_i$

Note:

- Do not need forces between atoms (for accelerations)
- Only valid for equilibrium processes

Property calculation with MC: example



Complex moves

 Move sets can be adapted for other cases, e.g. not just move of particles but also rotations of side chains (=rotamers), torsions, etc.

E.g. application in protein folding problem when we'd like to determine the 3D folded structure of a protein in thermal equilibrium

Possible Monte Carlo moves

Trial moves

- Rigid body translation
- Rigid body rotation
- Internal conformational changes (soft vs. stiff modes)
- Titration/electronic states
- ...

• Questions:

- How "big" a move should we take?
- Move one particle or many?

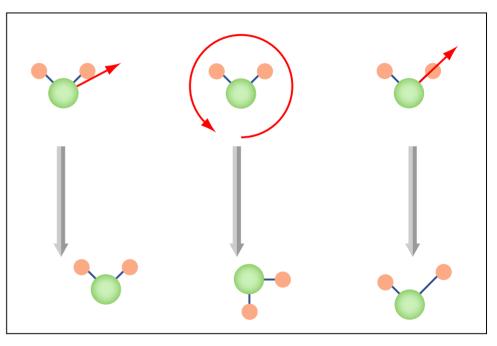
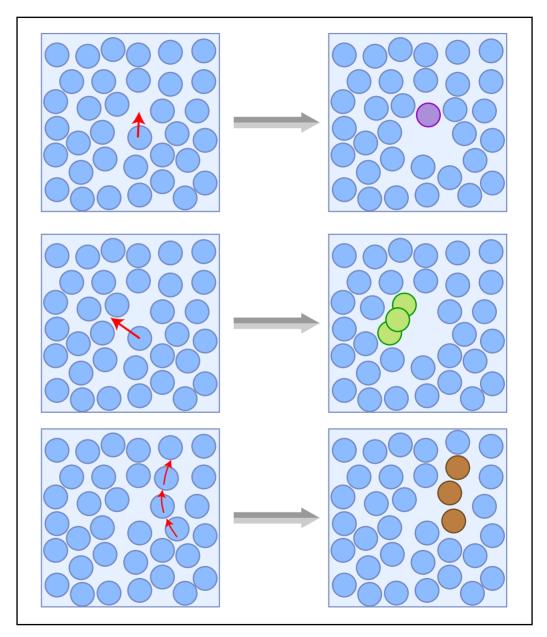


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Monte Carlo moves

- How "big" a move should we take?
 - Smaller moves: better acceptance rate, slower sampling
 - Bigger moves: faster sampling, poorer acceptance rate
- Move one particle or many?
 - Possible to achieve more efficient sampling with correct multiparticle moves
 - One-particle moves must choose particles at random



2. How to model chemical interactions

Atomic interactions – different types of chemical bonds

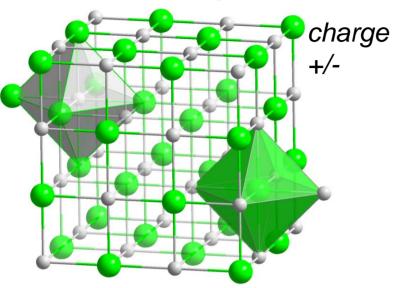
- Primary bonds ("strong")
 - Ionic (ceramics, quartz, feldspar rocks)
 - Covalent (silicon)
 - Metallic (copper, nickel, gold, silver) (high melting point, 1000-5,000K)
- Secondary bonds ("weak")
 - Van der Waals (wax, low melting point)
 - Hydrogen bonds (proteins, spider silk) (melting point 100-500K)

Atomic interactions – different types of chemical bonds

- Primary bonds ("strong")
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- Secondary bonds ("weak")
 - Van der Waals (wax, low melting point)
 - Hydrogen bonds (proteins, spider silk) (melting point 100-500K)
- Ionic: Non-directional (point charges interacting)
- Covalent: Directional (bond angles, torsions matter)
- Metallic: Non-directional (electron gas concept)

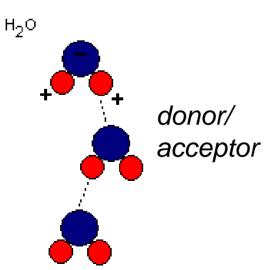
Difference of material properties originates from different atomic interactions

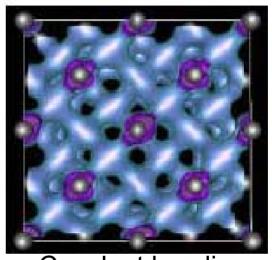
Types of bonding (illustrations)



Ionic bonding

Hydrogen bonding

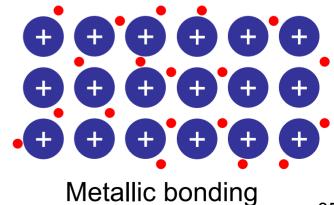




electron density (localized!)

Covalent bonding

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Wax



Courtesy of Ruth Ruane, http://www.whitewitch.ie. Used with permission.

Rocks



Image courtesy of Wikimedia Commons.

Rocks and sand on Mars

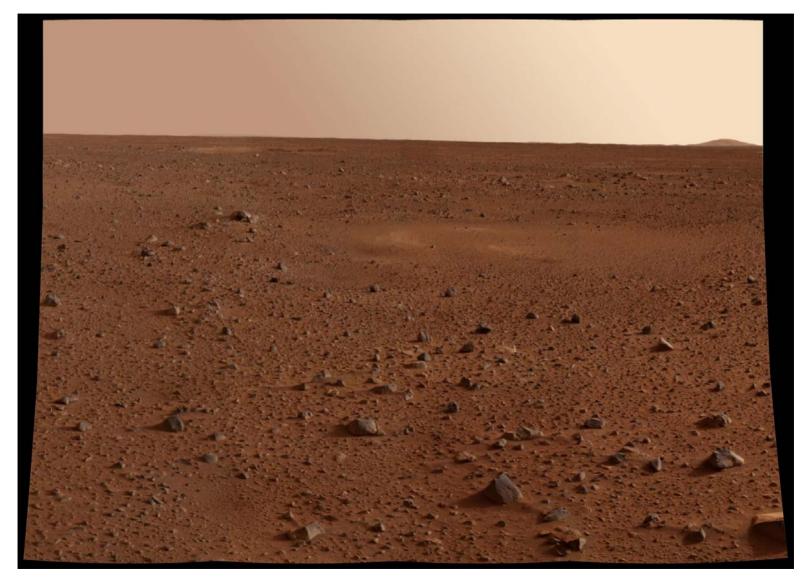


Image courtesy of NASA.

Gold



Image courtesy of Wikimedia Commons.

Silicon



Image courtesy of NASA.

Spider web



Image courtesy of U.S. Fish and Wildlife Service.

Tree's leaf



Image courtesy of Wikimedia Commons.

Particularly intriguing...brittle or ductile?

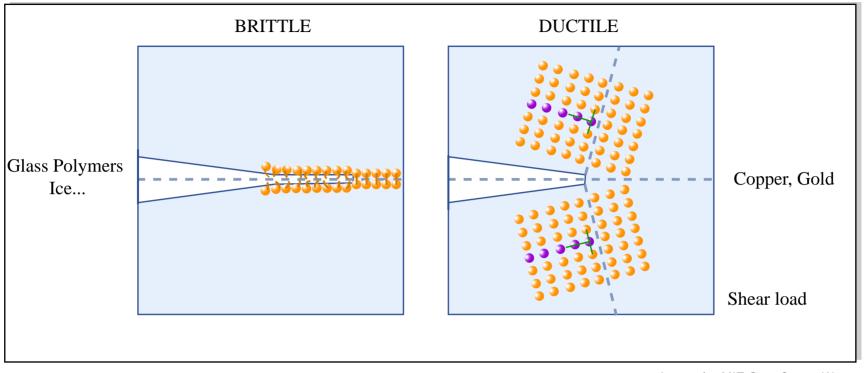


Image by MIT OpenCourseWare.

Image courtesy of quinn.anya.







Outline

 Goal: model chemical bonds with the objective to enable force calculation (see lecture 2, basic MD algorithm) or energy calculation (see lecture 4/5, MC)

Two-step approach:

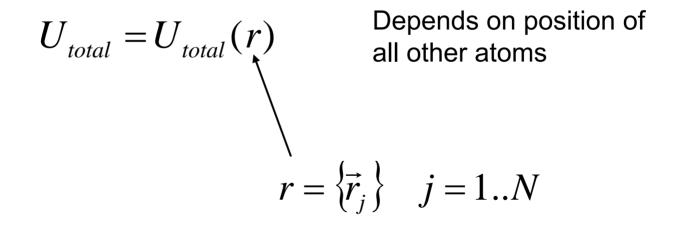
- 1. Define energy landscape, *i.e.* defines how distance between particles controls the energy stored in the bond
- 2. Then take derivatives to obtain forces, to be used in the MD algorithm

"Modeling and simulation" paradigm:

- First, develop mathematical expressions (modeling)
- Second, use model in numerical solution (simulation, =MD)

Models for atomic interactions

• Define interatomic potentials that describe the energy of a set of atoms as a function of their coordinates r:



Models for atomic interactions

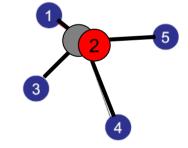
Define interatomic potentials that describe the energy of a set of atoms as a function of their coordinates r:

$$r = \left\{\vec{r}_j\right\} \quad j = 1..N$$

$$U_{total} = U_{total}(r)$$
 Depends on position of all other atoms

$$\vec{F}_i = -\nabla_{\vec{r}_i} U_{total}(r) \qquad i = 1..N$$

$$i = 1..N$$

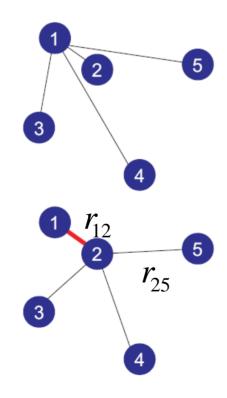


$$\nabla_{\vec{r_i}} = \left(\frac{\partial}{\partial r_{1,i}}, \frac{\partial}{\partial r_{2,i}}, \frac{\partial}{\partial r_{3,i}}\right)$$
 Change of potential energy due to change of position of particle *i* ("gradient")

2.1 Pair potentials

Pair potentials: energy calculation

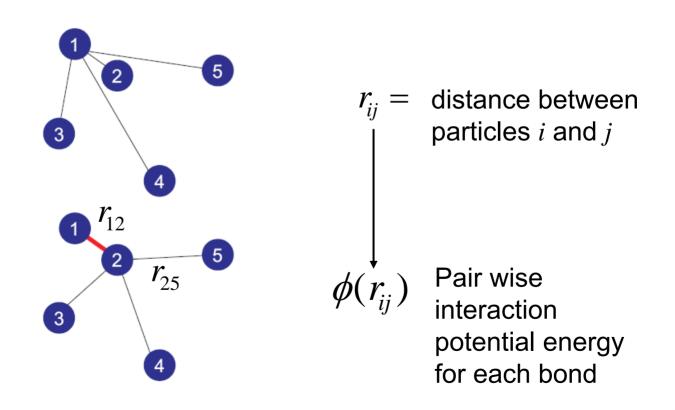
Simple approximation: Total energy is sum over the energy of all pairs of atoms in the system



 r_{ij} = distance between particles i and j

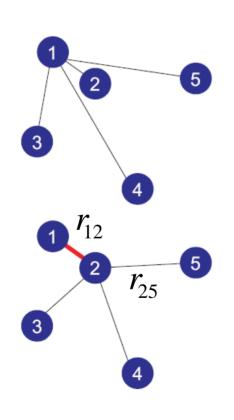
Pair potentials: energy calculation

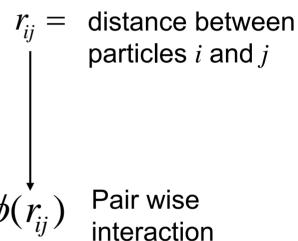
Simple approximation: Total energy is sum over the energy of all pairs of atoms in the system



Pair potentials: energy calculation

Simple approximation: Total energy is sum over the energy of all pairs of atoms in the system





interaction potential energy for each bond

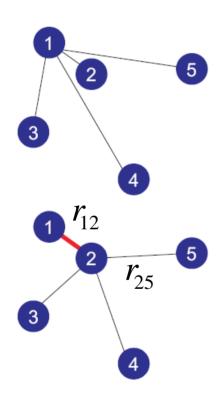
Energy of atom
$$i$$
 $U_i = \sum_{j=1}^N \phi(r_{ij})$

Overview - pair potentials: total energy calculation

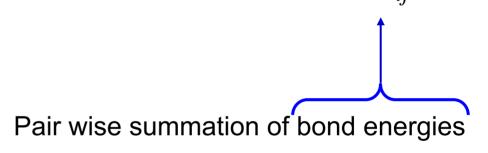
Simple approximation: Total energy is sum over the energy of all pairs of atoms in the system

Pair wise interaction potential

 $\phi(r_{ii})$



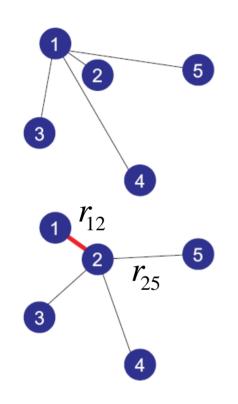
$$r_{ij}$$
 = distance between particles i and j



Energy of atom
$$i$$
 $U_i = \sum_{j=1}^N \phi(r_{ij})$

avoid double counting
$$U_{total} = \underbrace{\frac{1}{2} \sum_{i=1, i \neq j}^{N} \sum_{j=1}^{N} \phi(r_{ij})}_{}$$

Example: calculation of total energy



two "loops" over pairs of all particles

$$U_{total} = \frac{1}{2} \sum_{i=1, i \neq j}^{N} \sum_{j=1}^{N} \phi(r_{ij})$$

with
$$\phi_{ij} = \phi(r_{ij})$$

$$U_{total} = \frac{1}{2} \left(\phi_{12} + \phi_{13} + \phi_{14} + \phi_{1N} \dots + \phi_{21} + \phi_{23} + \dots + \phi_{2N} + \dots + \phi_{N-1,N} \right)$$

Interatomic pair potentials: examples

$$\phi(r_{ij}) = D \exp(-2\alpha(r_{ij} - r_0)) - 2D \exp(-\alpha(r_{ij} - r_0))$$

Morse potential

$$\phi(r_{ij}) = 4\varepsilon \left[\left(\frac{\sigma}{r_{ij}} \right)^{12} - \left(\frac{\sigma}{r_{ij}} \right)^{6} \right]$$

Lennard-Jones 12:6 potential (excellent model for noble Gases, Ar, Ne, Xe..)

$$\phi(r_{ij}) = A \exp\left(-\frac{r_{ij}}{\sigma}\right) - C\left(\frac{\sigma}{r_{ij}}\right)^{6}$$

Buckingham potential

$$\phi(r_{ij}) = a_0 + \frac{1}{2}k(r_{ij} - r_0)^2$$

Harmonic approximation (no bond breaking)

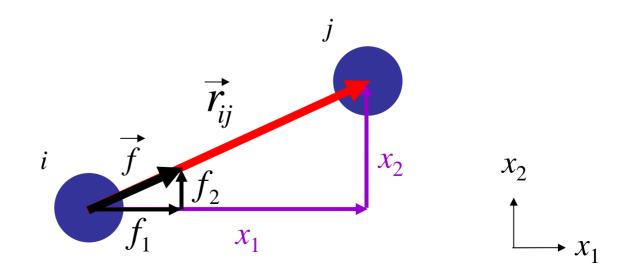
How to use a pair potential, e.g. LJ

Force calculation – pair potential

Forces on particles can be calculated by taking derivatives from the potential function & by considering all pairs of atoms

Start with **force magnitude (STEP 1):** Negative derivative of potential energy with respect to atomic distance

$$F = -\frac{d \phi(r)}{d r} \Big|_{r=r_{ij}} = -\frac{d \phi(r_{ij})}{d r_{ij}} = -\phi'(r_{ij})$$



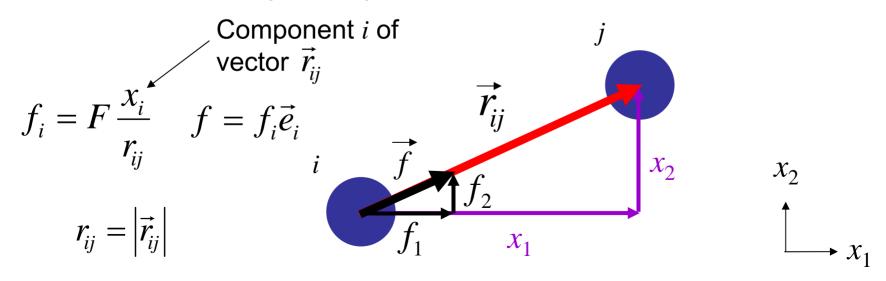
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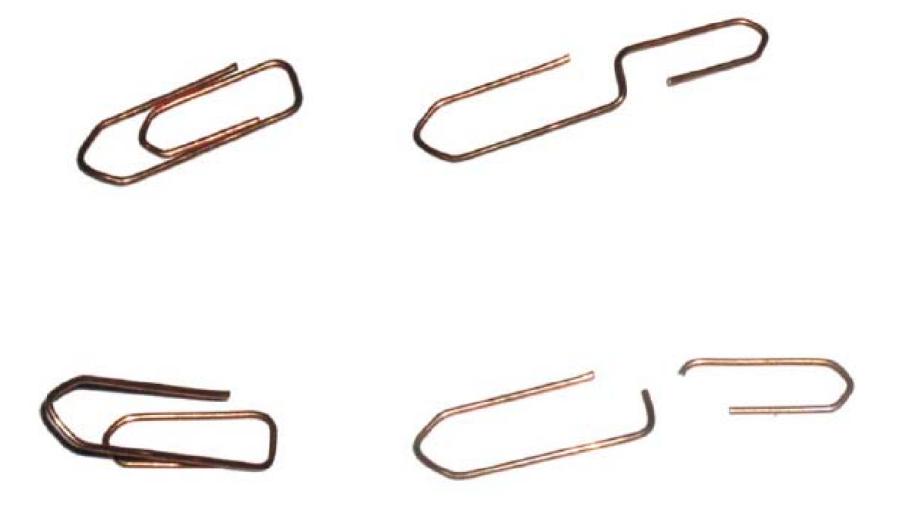
$$F = -\frac{d \phi(r)}{d r} \Big|_{r=r_{ij}} = -\frac{d \phi(r_{ij})}{d r_{ij}} = -\phi'(r_{ij})$$

Calculate force vector (STEP 2):

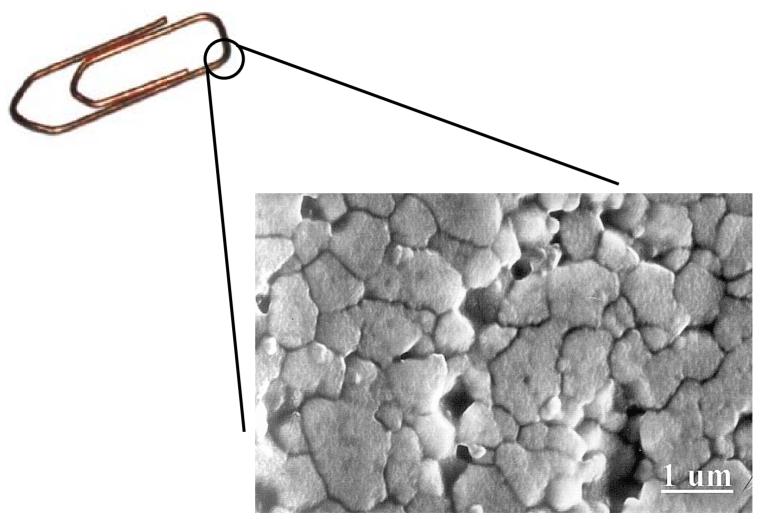


What can we do with this potential?

Bending a copper wire until it breaks



A closer look

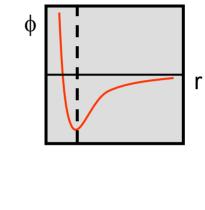


Courtesy of Goran Drazic. Used with permission.

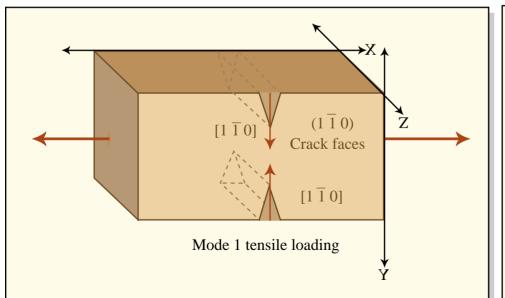
Case study: plasticity in a micrometer crystal of copper

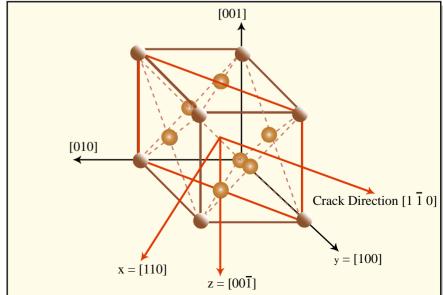
Simulation details

- 1,000,000,000 atoms (0.3 micrometer side length)
- 12:6 Lennard-Jones ductile material, for copper
- Visualization using energy filtering method (only show high energy atoms)



Generic features of atomic bonding: "repulsion vs. attraction"





A simulation with 1,000,000,000 particles Lennard-Jones - copper

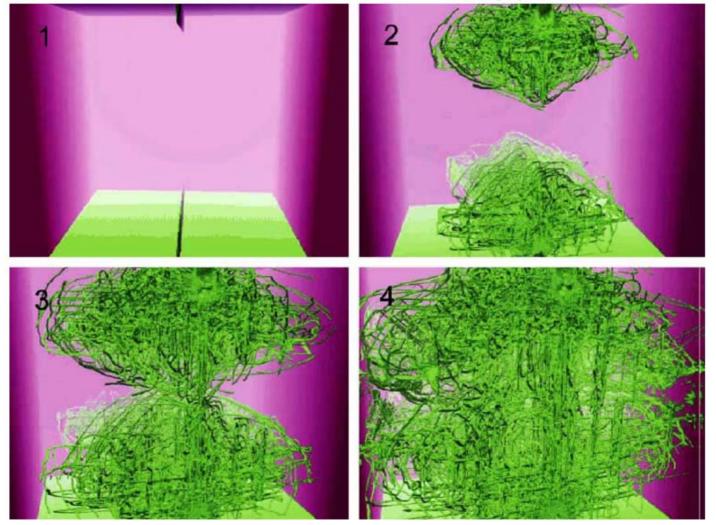
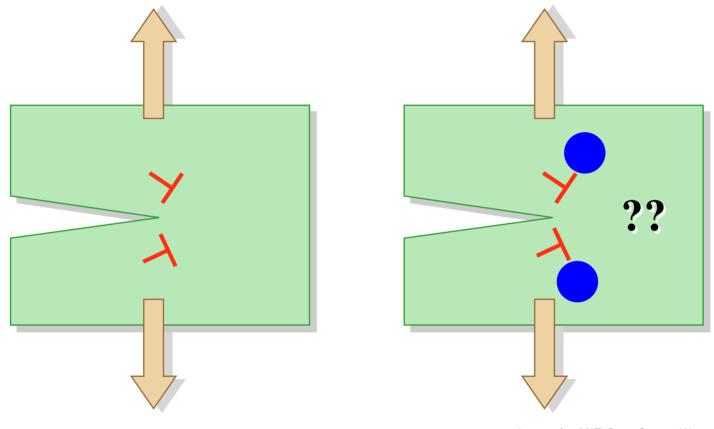
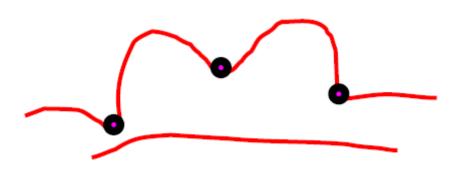


Fig. 1 c from Buehler, M., et al. "The Dynamical Complexity of Work-Hardening: A Large-Scale Molecular Dynamics Simulation." *Acta Mech Sinica* 21 (2005): 103-11.
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Strengthening mechanisms







Strengthening caused by hindering dislocation motion

If too difficult, ductile modes break down and material becomes brittle

Parameters for Morse potential

(for reference)

Morse potential parameters for various metals

N / T	1 - 4 4 : - 1	D	C 1	✓ N / - 4 - 1 -
- Morse E	orennai	Parameters	TOT I	b Metals

Metal	αa_0	β	L x 10 ⁻²² (eV)	$\alpha (A^{-1})$	$r_0(A)$	D (eV)	
Pb	2.921	83.02	7.073	1.1836	3.733	0.2348	
Ag	2.788	71.17	10.012	1.3690	3.115	0.3323	
Ni	2.500	51.78	12.667	1.4199	2.780	0.4205	
Cu	2.450	49.11	10.330	1.3588	2.866	0.3429	
Al	2.347	44.17	8.144	1.1646	3.253	0.2703	
Ca	2.238	39.63	4.888	0.80535	4.569	0.1623	
Sr	2.238	39.63	4.557	0.73776	4.988	0.1513	
Mo	2.368	88.91	24.197	1.5079	2.976	0.8032	
W	2.225	72.19	29.843	1.4116	3.032	0.9906	
Cr	2.260	75.92	13.297	1.5721	2.754	0.4414	
Fe	1.988	51.97	12.573	1.3885	2.845	0.4174	
Ba	1.650	34.12	4.266	0.65698	5.373	0.1416	
K	1.293	23.80	1.634	0.49767	6.369	0.05424	
Na	1.267	23.28	1.908	0.58993	5.336	0.06334	
Cs	1.260	23.14	1.351	0.41569	7.557	0.04485	
Rb	1.206	22.15	1.399	0.42981	7.207	0.04644	

Adapted from Table I in Girifalco, L. A., and V. G. Weizer. "Application of the Morse Potential Function to Cubic Metals." *Physical Review* 114 (May 1, 1959): 687-690.

Image by MIT OpenCourseWare.

$$\phi(r_{ij}) = D \exp(-2\alpha(r_{ij} - r_0)) - 2D \exp(-\alpha(r_{ij} - r_0))$$

Morse potential: application example (nanowire)

Source: Komanduri, R., et al. "Molecular Dynamics (MD) Simulation of Uniaxial Tension of Some Single-Crystal Cubic Metals at Nanolevel." *International Journal of Mechanical Sciences* 43, no. 10 (2001): 2237-60.

Further Morse potential parameters:

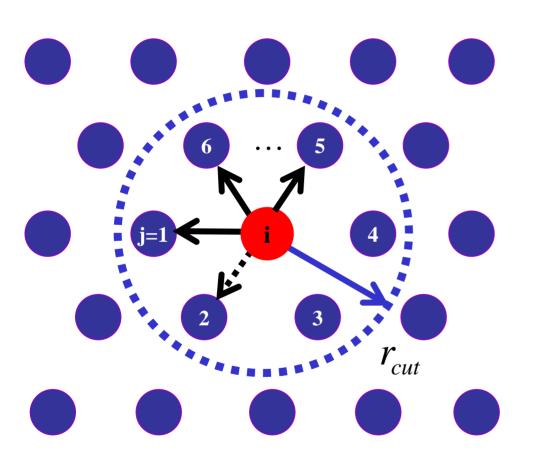
Table 3 Morse potential parameters used in MD simulation of uniaxial tensile loading [24]

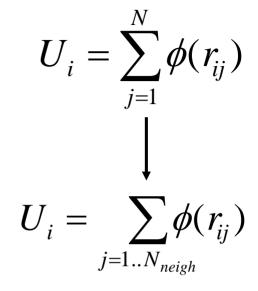
Material	Crystal structure	Dissociation energy, D (eV)	Equilibrium radius, r_0 (Å)	α -parameter (\mathring{A}^{-1})	Lattice constant (Å)
Aluminium	FCC	0.2703	3.253	1.1650	4.05
Copper	FCC	0.3429	2.866	1.3590	3.62
Nickel	FCC	0.4205	2.780	1.4199	3.52
Iron	BCC	0.4172	2.845	1.3890	2.87
Chromium	BCC	0.4414	2.754	1.5721	2.89
Tungsten	BCC	0.9906	3.032	1.4116	3.17

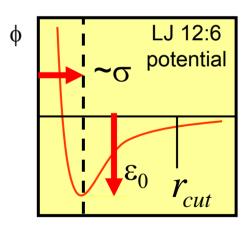
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Cutoff-radius: saving time

Cutoff radius







1

Cutoff radius = considering interactions only to a certain distance Basis: Force contribution negligible (slope)

Derivative of LJ potential ~ force

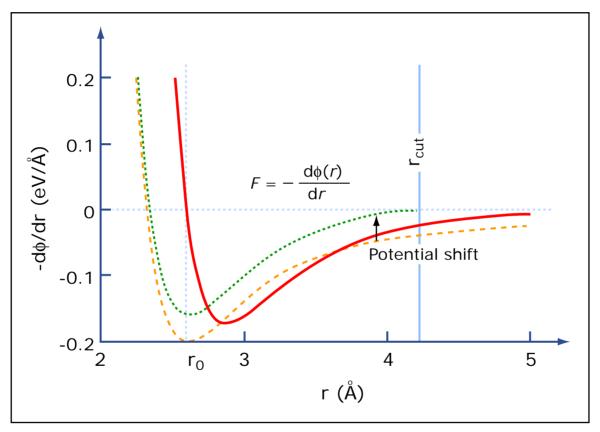


Image by MIT OpenCourseWare.

Beyond cutoff: Changes in energy (and thus forces) small

Putting it all together...

MD updating scheme: Complete

(1) Updating method (integration scheme)

$$r_i(t_0 + \Delta t) = -r_i(t_0 - \Delta t) + 2r_i(t_0)\Delta t + a_i(t_0)(\Delta t)^2 + \dots$$

Positions at t_0 - Δt

Positions at t_0

Accelerations at t_0

(2) Obtain accelerations from forces

$$f_i = ma_i$$
 $a_i = f_i / m$

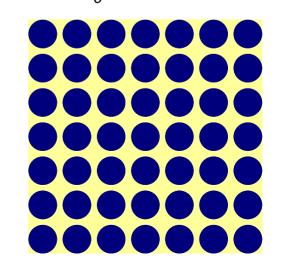
(4) Crystal (initial conditions) Positions at t_0

(3) Obtain forces from potential

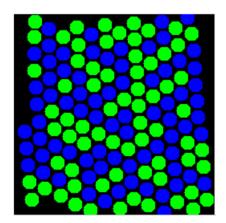
$$F = -\frac{\mathrm{d}\,\phi(r)}{\mathrm{d}\,r} \qquad f_i = F\frac{x_i}{r}$$

Potential

$$\phi(r) = 4\varepsilon \left(\left[\frac{\sigma}{r} \right]^{12} - \left[\frac{\sigma}{r} \right]^{6} \right)$$



2.2 How to model metals: Multi-body potentials



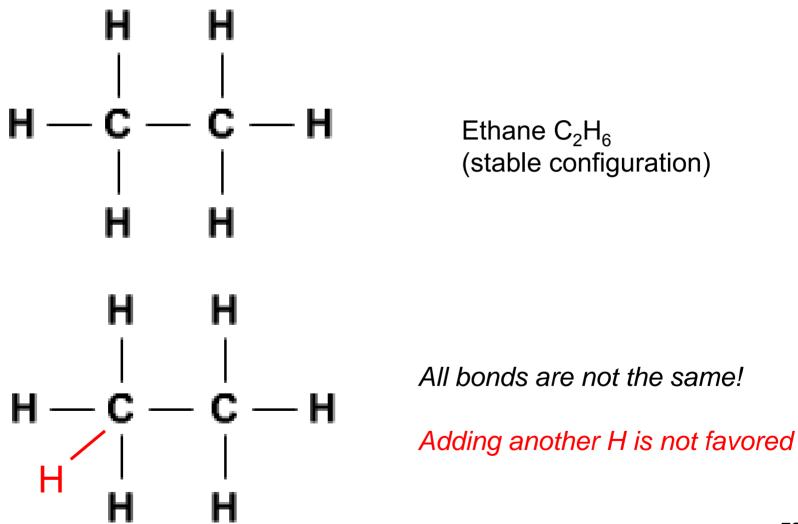
Courtesy of the Center for Polymer Studies at Boston University. Used with permission.

Pair potential: Total energy sum of all pairs of bonds Individual bond contribution does not depend on other atoms

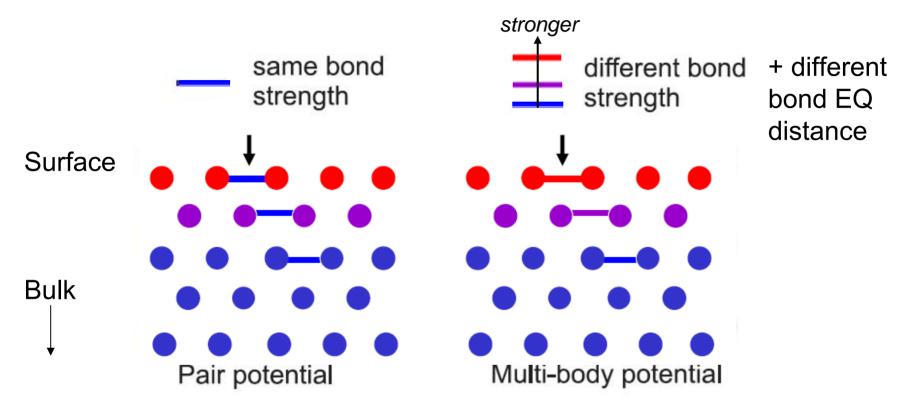
"all bonds are the same"

$$U_{total} = \frac{1}{2} \sum_{i=1, i \neq j}^{N} \sum_{j=1}^{N} \phi(r_{ij})$$

Are all bonds the same? - valency in hydrocarbons



Are all bonds the same? – metallic systems



Pair potentials: All bonds are equal!

Reality: Have environment effects; it matter that there is a free surface!

rree surrace!

Are all bonds the same?

Bonding energy of red atom in is six times bonding energy in •••





This is in contradiction with both experiments and more accurate quantum mechanical calculations on many materials

Bonding energy of atom
$$i$$
 $U_i = \sum_{j=1}^N \phi(r_{ij})$



$$U_i = \sum_{j=1}^6 \phi(r_{ij}) \qquad \longleftarrow \qquad U_i = \phi(r_{ij})$$



Are all bonds the same?

Bonding energy of red atom in is six times bonding energy in •••





This is in contradiction with both experiments and more accurate quantum mechanical calculations on many materials

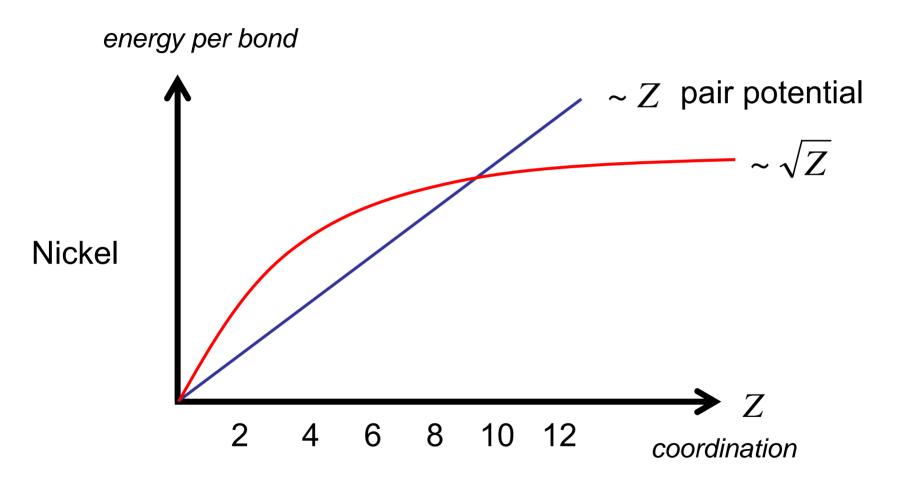
For pair potentials $\sim Z$

For metals $\sim \sqrt{Z}$

Z: Coordination = how many immediate neighbors an atom has

Bonds get "weaker" as more atoms are added to central atom

Bond strength depends on coordination



Transferability of pair potentials

Pair potentials have limited transferability:

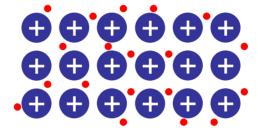
Parameters determined for molecules can not be used for crystals, parameters for specific types of crystals can not be used to describe range of crystal structures

 E.g. difference between FCC and BCC can not be captured using a pair potential

Metallic bonding: multi-body effects

 Need to consider more details of chemical bonding to understand environmental effects

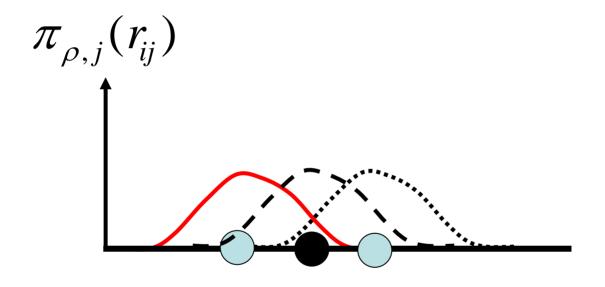
- Electron (q=-1)
- lon core (q=+N)



Delocalized valence electrons moving between nuclei generate a binding force to hold the atoms together: **Electron gas model** (positive ions in a sea of electrons)

Mostly non-directional bonding, but the bond strength indeed depends on the environment of an atom, precisely the electron density imposed by other atoms

Concept: include electron density effects



Each atom features a particular distribution of electron density

Concept: include electron density effects

Electron density at atom
$$i$$
 $\rho_i = \sum_{j=1..N_{neigh}} \pi_{\rho,j}(r_{ij})$ Atomic electron density of atom j

Contribution to electron density at site i due to electron density of atom j evaluated at correct distance (r_{ii})

Concept: include electron density effects

$$\phi_i = \sum_{j=1..N_{neigh}} \frac{1}{2} \phi(r_{ij}) + F(\rho_i)$$
Embedding term F
(how local electron density contributes to potential energy)

Electron density at atom i $\rho_i = \sum_{j=1..N_{neigh}} \pi_{\rho,j}(r_{ij})$ $\pi_{\rho,j}(r_{ij})$ Atomic electron density of atom i

Embedded-atom method (EAM)

Atomic energy

$$\phi_i = \sum_{j=1..N_{neigh}} \frac{1}{2} \phi(r_{ij}) + F(\rho_i)$$

Pair potential energy

Embedding energy as a function of electron density

Total energy

$$U_{total} = \sum_{i=1}^{N} \phi_i$$

 ρ_i Electron density at atom i based on a "pair potential":

$$\rho_i = \sum_{j=1..N_{neigh}} \pi_{\rho,j}(r_{ij})$$

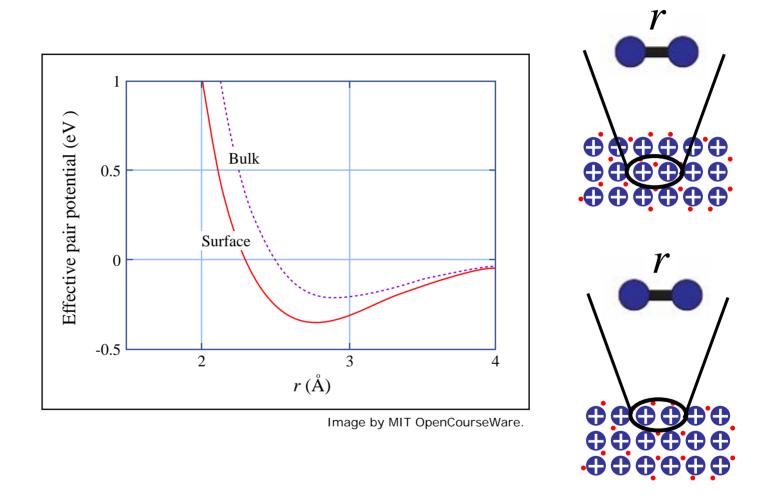
Physical concept: EAM potential

Describes bonding energy due to electron delocalization

As electrons get more states to spread out over their kinetic energy decreases

- When an impurity is put into a metal its energy is lowered because the electrons from the impurity can delocalize into the solid.
- The embedding density (electron density at the embedding site) is a measure of the number of states available to delocalize onto.
- Inherently MANY BODY effect!

Effective pair interactions



Can describe differences between bulk and surface

Summary: EAM method

- State of the art approach to model metals
- Very good potentials available for Ni, Cu, Al since late 1990s, 2000s
- Numerically efficient, can treat billions of particles
- Not much more expensive than pair potential (approximately three times), but describes physics much better
- Strongly recommended for use!



 $3.021 J\,/\,1.021 J\,/\,10.333 J\,/\,18.361 J\,/\,22.00 J$ Introduction to Modeling and Simulation Spring 2012

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