1.021, 3.021, 10.333, 22.00 Introduction to Modeling and Simulation Spring 2018

# Basic molecular dynamics

Lecture 2

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### Content overview

#### I. Fundamentals of particle methods

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

Lectures 1-12 February/March

#### II. Advanced topics in particle methods

- 1. Quantum Weirdness: The Theory of Quantum Mechanics
- 2. The Many-Body Problem: From Many-Body to Single-Particle
- 3. Quantum modeling of materials
- 4. From Atoms to Solids
- 5. Basic properties of materials
- 6. Advanced properties of materials
- Materials Informatics

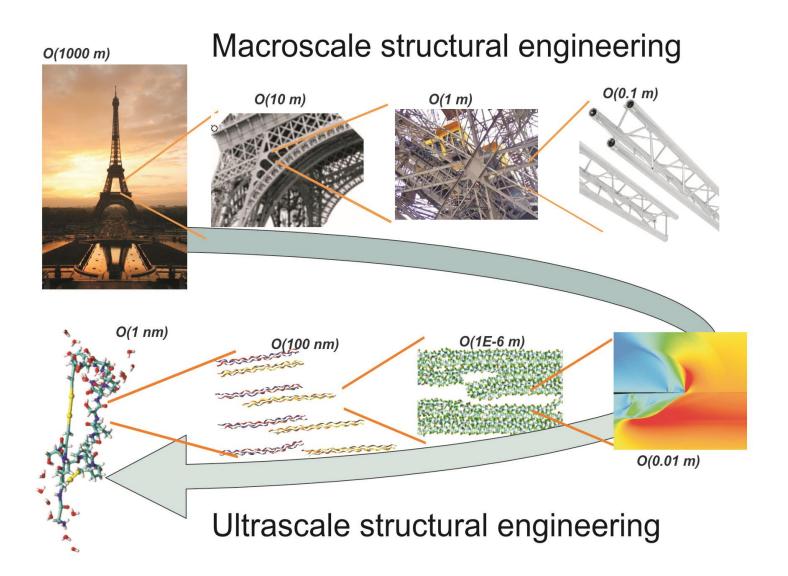
Lectures 13-24 March/April/May

# Goals of part I (particle methods)

You will be able to ...

- Carry out atomistic simulations of various processes (diffusion, deformation/stretching, materials failure)
   Nanowires, bulk metals, proteins, silicon crystals, etc.
- Analyze atomistic simulations (make sense of the numbers)
- Visualize atomistic/molecular data (bring data to life)
- Understand how to link atomistic simulation results with continuum models within a multi-scale scheme

## Multi-scale view of materials



# Lecture 2: Basic molecular dynamics

#### **Outline:**

- 1. Example application: Stiffness of materials
- 2. Introduction: Bottom-up vs. top-down
- 3. Case study: Diffusion
  - 3.1 Continuum model
  - 3.2 Atomistic model (to be continued)

#### Goals of today's lecture:

- Through case study of diffusion, illustrate the concepts of a continuum model and an atomistic model
- Develop appreciation for distinction of continuum and atomistic approach
- Develop equations/models for diffusion problem from both perspectives

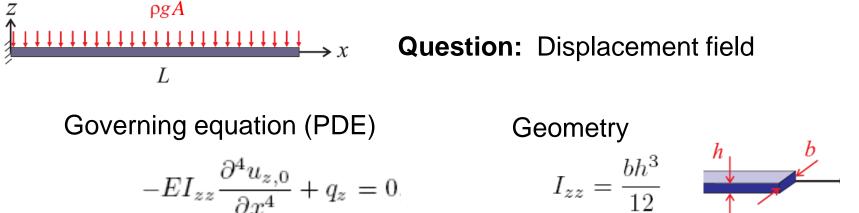
#### **Next:**

- Develop atomistic simulation approach (e.g. algorithm, pseudocode, etc.) and apply to describe diffusion (calculate diffusivity)
- Historical perspective on computer simulation with MD, examples from literature

# 1. Example application: Stiffness of materials (Young's modulus)

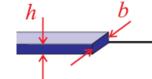
Objective: Illustrate the significance of multiple scales for material behavior and introduce multi-scale modeling paradigm

# Beam deformation problem – continuum model



Integration & BCs

$$I_{zz} = \frac{bh^3}{12}$$



BC - load:

$$\rho g A$$



$$u_z(x) = -\frac{\rho g A}{24 E I_{zz}} x^4$$

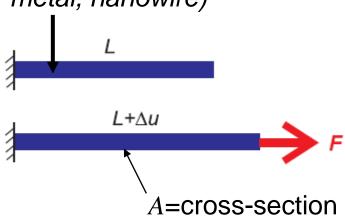
E = unknown parameter

E is parameter called "Young's modulus" that relates how force and deformation are related (captures properties of material)

# How to determine Young's modulus E?

#### Measurement (laboratory):

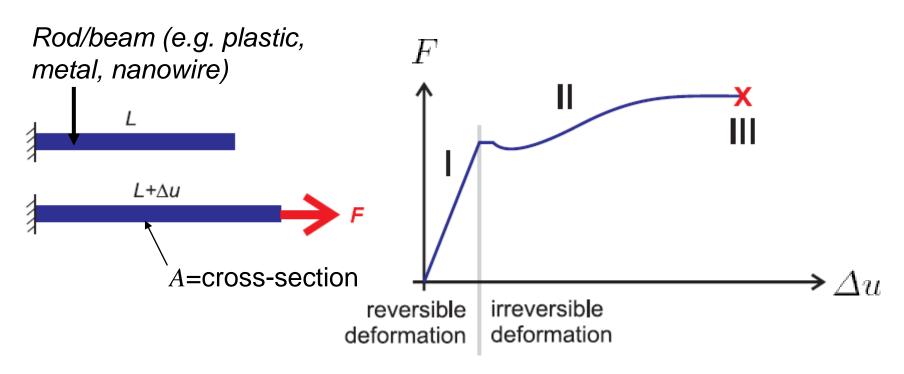
Rod/beam (e.g. plastic, metal, nanowire)





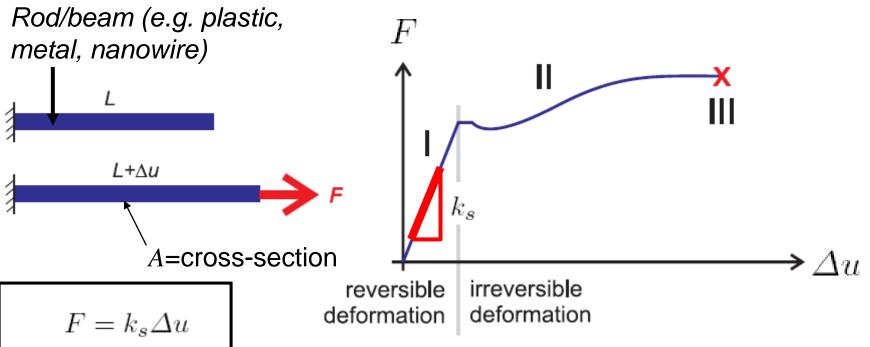
# How to determine Young's modulus E?

#### Measurement (laboratory):



# How to determine Young's modulus E?

#### Measurement (laboratory):

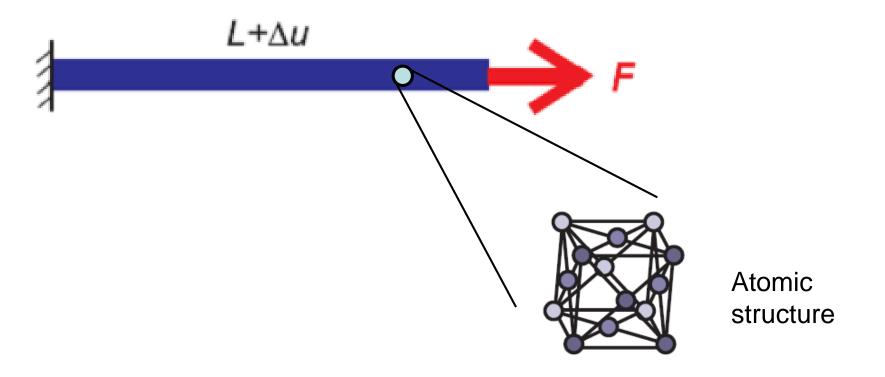


 $F = k_s \Delta u$   $E = \frac{Lk_s}{A}$ 

Young's modulus *E* (~stiffness=proportionality between force and displacement)

# How to determine E? - alternative approach to laboratory experiment

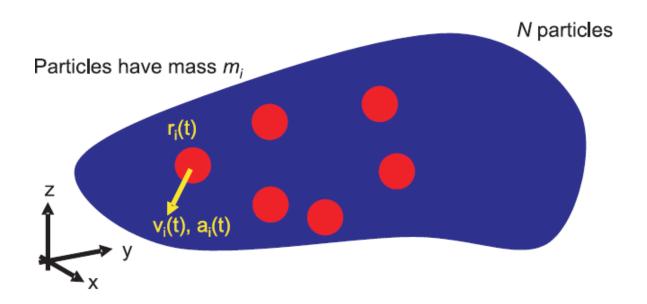
Atomistic simulation – new engineering paradigm



**Concept:** Consider the behavior of a collection of atoms inside the beam as deformation proceeds

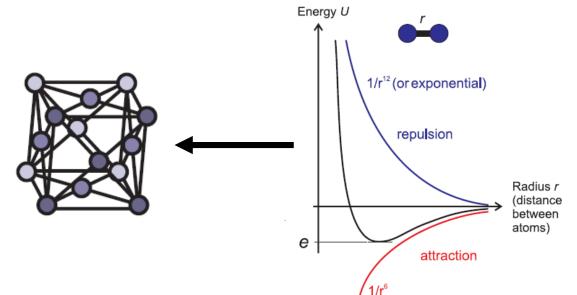
# Molecular dynamics simulation

- Newton's laws: F=ma
- Chemistry: Atomic interactions calculate interatomic forces from atomic interactions, that is, calculate F from energy landscape of atomic configuration (note that force and energy are related...)



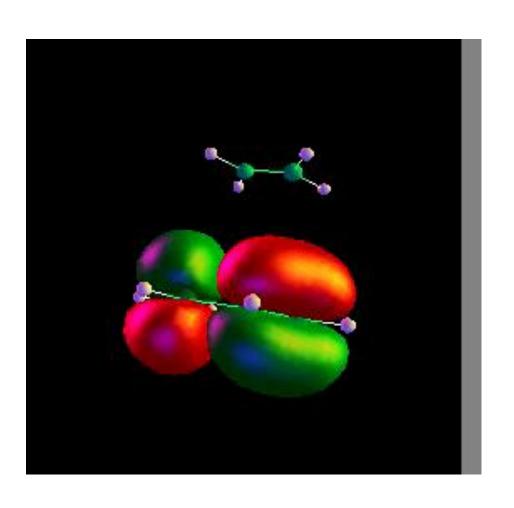
# Linking atomistic and continuum perspective

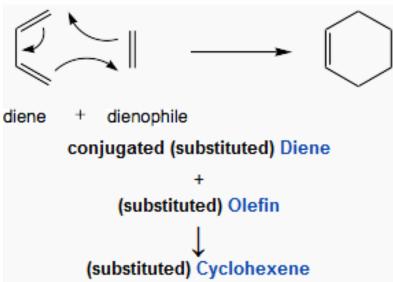
- Atomistic viewpoint enables us to calculate how force and deformation is related, that is, we can predict E once we know the atomic structure and the type of chemical bonds
- Example: In metals we have metallic bonding and crystal structures thus straightforward calculation of E
- Atomistic models provide fundamental perspective, and thereby a means to determine (solely from the atomistic / chemical structure of the material) important parameters to be used in continuum models



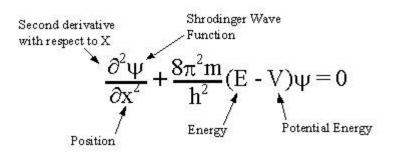
### Quantum mechanics

 Deals with fundamental view of chemical bonding, based on electrons in atoms



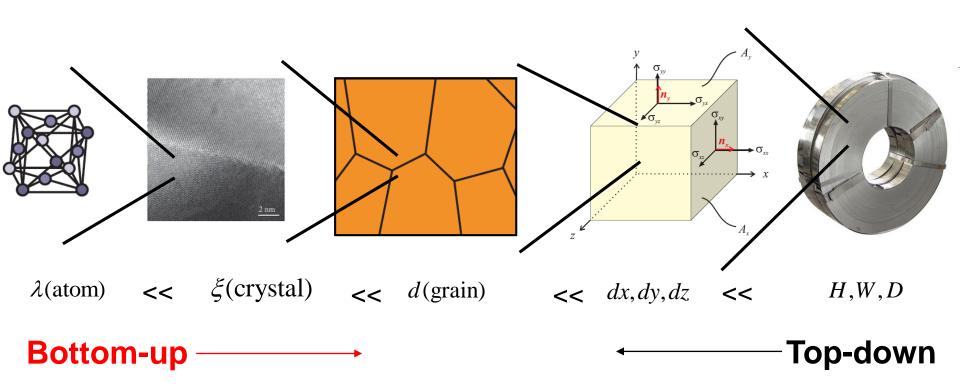


#### "Schroedinger equation"

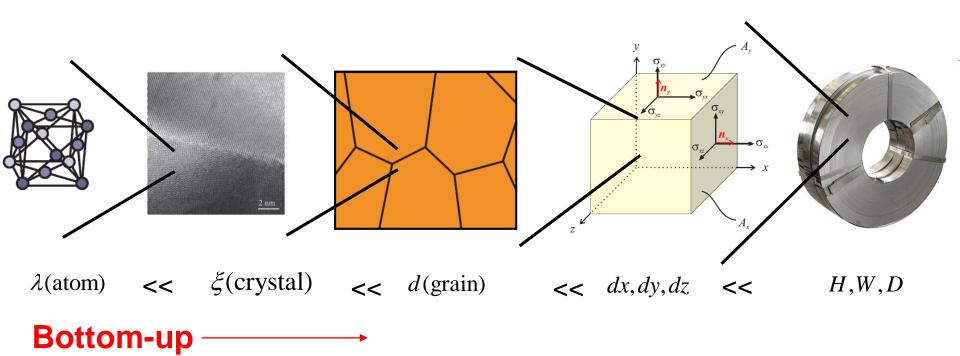


# 2. Introduction: Bottom-up vs. top-down

## Relevant scales in materials



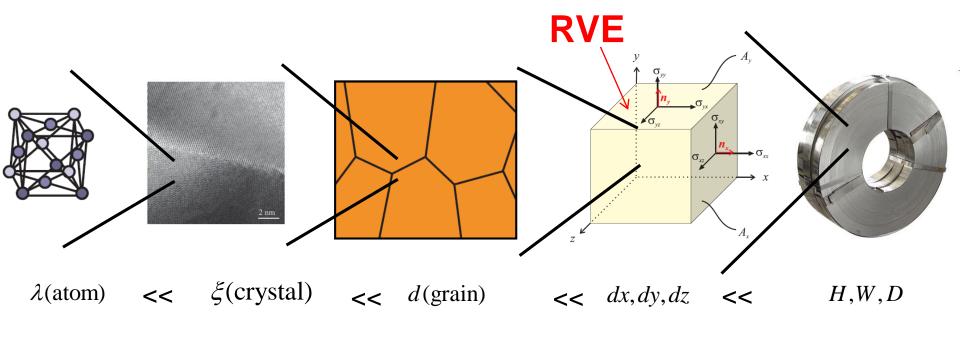
## Relevant scales in materials



#### Atomistic viewpoint:

- Explicitly consider discrete atomistic structure
- Solve for atomic trajectories and infer from these about material properties & behavior
- Features internal length scales (atomic distance)
- "Many-particle system with statistical properties"

## Relevant scales in materials



#### Continuum viewpoint:

- Treat material as matter with no internal structure
- •Develop mathematical model (governing equation) based on representative volume element (RVE, contains "enough" material such that internal structure can be neglected

Top-down

•Features no characteristic length scales, provided RVE is large enough <sup>18</sup> "PDE with parameters"

# 3. Case study: Diffusion

Continuum and atomistic modeling

### Goal of this section

- Diffusion as example
- Continuum description (top-down approach), partial differential equation
- Atomistic description (bottom-up approach), based on dynamics of molecules, obtained via numerical simulation of the molecular dynamics

### Introduction: Diffusion

- Particles move from a domain with high concentration to an area of low concentration
- Macroscopically, diffusion measured by change in concentration
- Microscopically, diffusion is process of spontaneous net movement of particles

Result of random motion of particles ("Brownian motion")



$$c = m/V = c(\vec{x}, t)$$

# Ink droplet in water



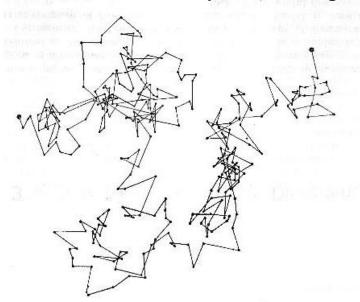
hot cold

(same time has passed)

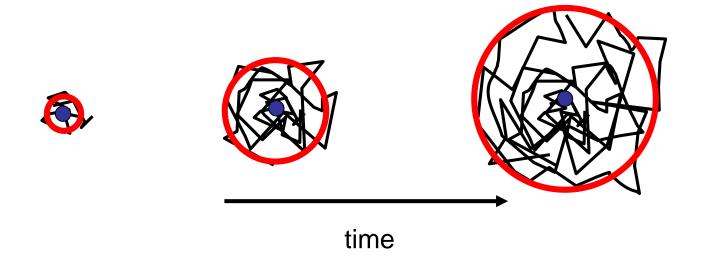
# Microscopic observation of diffusion

# Microscopic mechanism: "Random walk" – or Brownian motion

- Brownian motion was first observed (1827) by the British botanist Robert
   Brown (1773-1858) when studying pollen grains in water
- Initially thought to be sign of life, but later confirmed to be also present in inorganic particles
- The effect was finally explained in 1905 by Albert Einstein, who realized it was caused by water molecules randomly smacking into the particles.



# Brownian motion leads to net particle movement



Particle "slowly" moves away from its initial position

# Robert Brown's microscope

1827



http://www.brianjford.com/pbrownmica.jpg

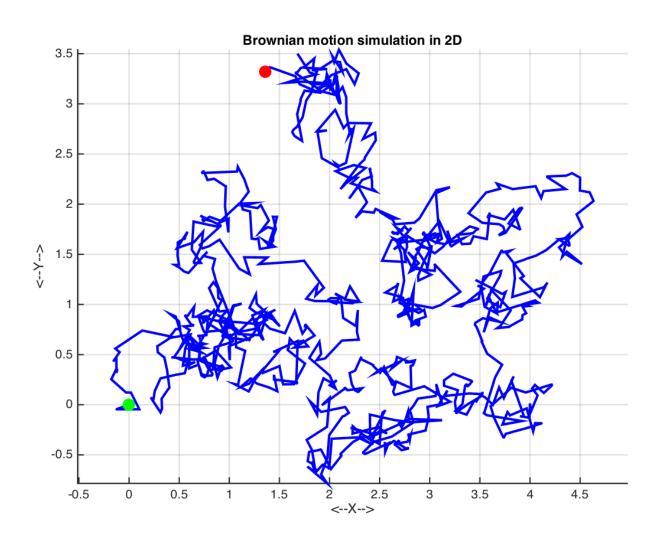
#### **Robert Brown's Microscope**

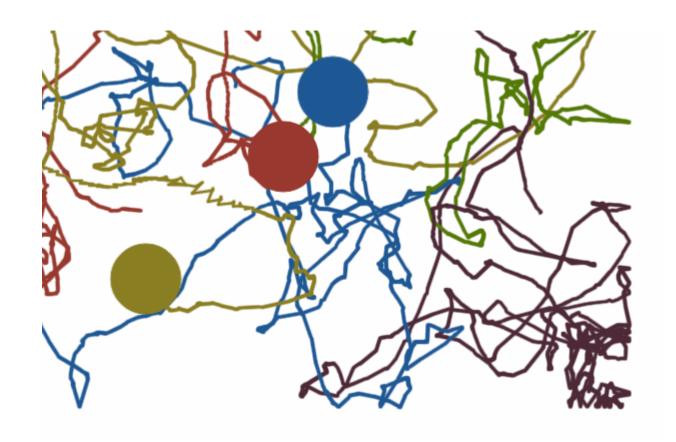
Instrument with which Robert Brown studied **Brownian motion** and which he used in his work on identifying the nucleus of the living cell

Instrument is preserved at the Linnean Society in London

It is made of brass and is mounted onto the lid of the box in which it can be stored

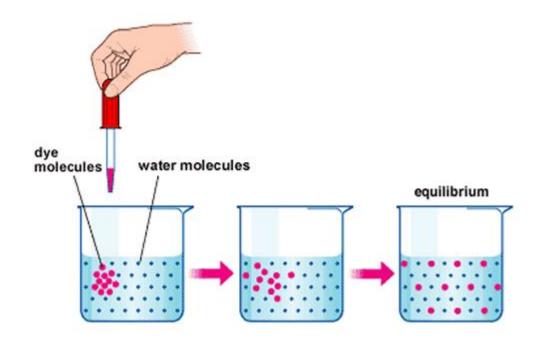
## Simulation of Brownian motion



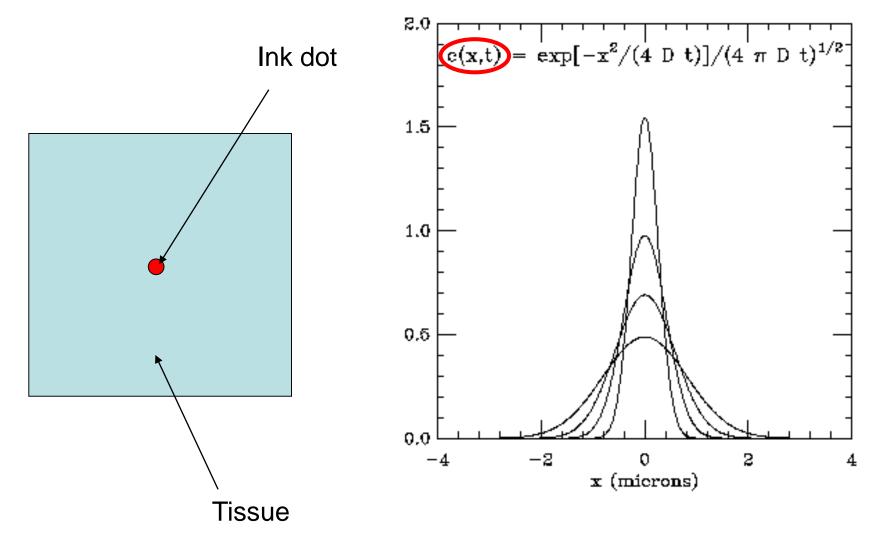




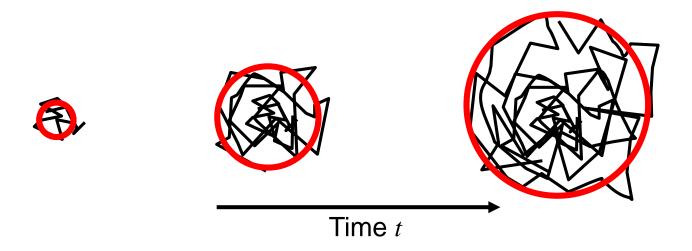
# Macroscopic observation of diffusion

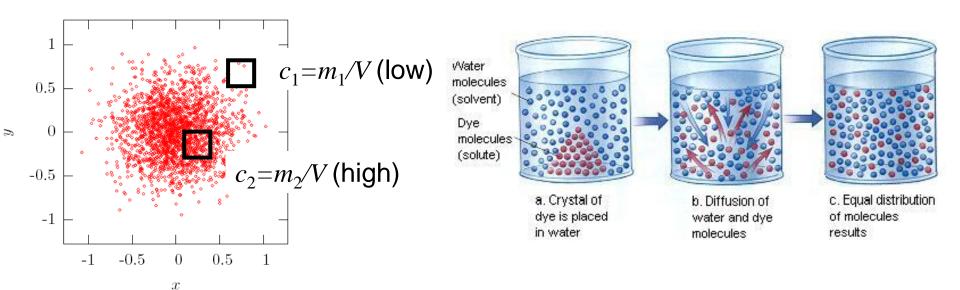


## Macroscopic observation: concentration change



## Brownian motion leads to net particle movement

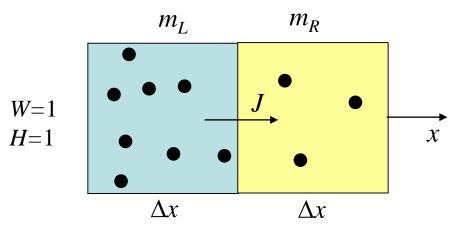




#### 3.1 Continuum model

How to build a continuum model to describe the physical phenomena of diffusion?

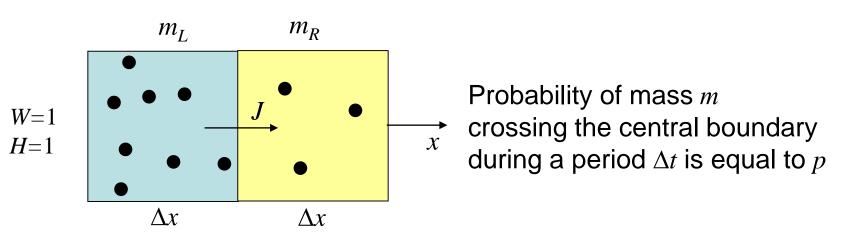
 Develop differential equation based on differential element



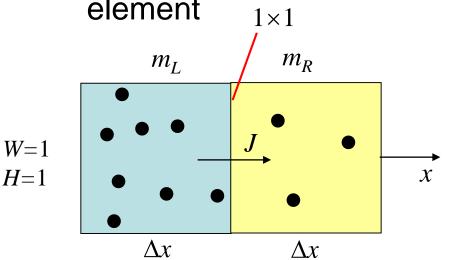
J: Mass flux (mass per unit time per unit area)

**Concept:** Balance mass [here], force etc. in a differential volume element; much greater in dimension than inhomogeneities ("sufficiently large RVE")

Develop differential equation based on differential element



Develop differential equation based on differential element



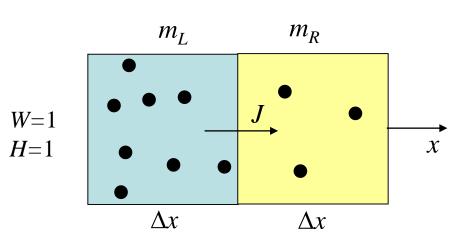
Probability of mass m crossing the central boundary during a period  $\Delta t$  is equal to p

$$J_L = \frac{1}{1 \times 1} \frac{p}{\Delta t} m_L$$
 Mass flux from left to right

$$J_R = \frac{1}{1 \times 1} \frac{p}{\Delta t} m_R$$
 Mass flux from right to left

[J] = mass per unit time per unit area

Develop differential equation based on differential element



Probability of mass mcrossing the central boundary during a period  $\Delta t$  is equal to p

$$J_L = \frac{1}{1 \times 1} \frac{p}{\Delta t} m_L$$
 Mass flux from left to right

$$J_R = \frac{1}{1 \times 1} \frac{p}{\Delta t} m_R$$
 Mass flux from right to left

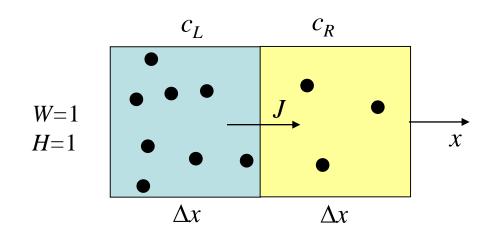
Effective mass flux

$$J = \frac{1}{1 \times 1} \frac{p}{\Delta t} (m_L - m_R)$$

More mass, more flux ( $m_L$  is ~ to number of particles)

Express in terms of mass concentrations

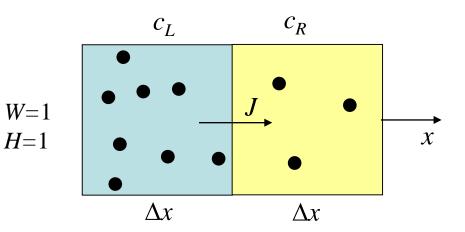
$$c = \frac{m}{V}$$
  $m = cV$   $J = \frac{p}{\Delta t} (m_L - m_R)$ 



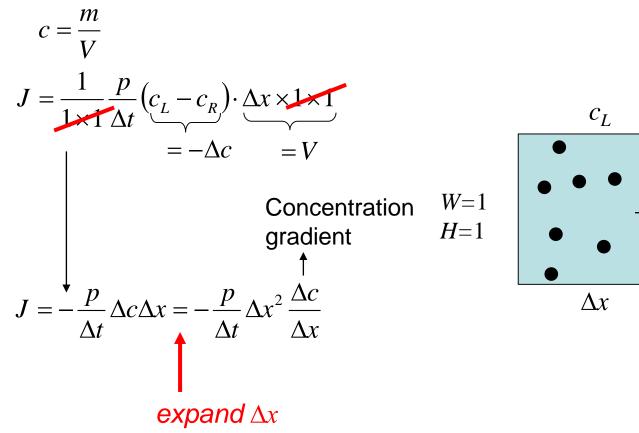
#### Express in terms of mass concentrations

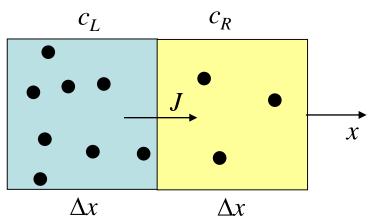
$$c = \frac{m}{V}$$
  $m = cV$   $J = \frac{p}{\Delta t} (m_L - m_R)$ 

$$J = \frac{1}{1 \times 1} \frac{p}{\Delta t} \left( \underbrace{c_L - c_R} \right) \cdot \underbrace{\Delta x \times 1 \times 1}_{= -\Delta c}$$

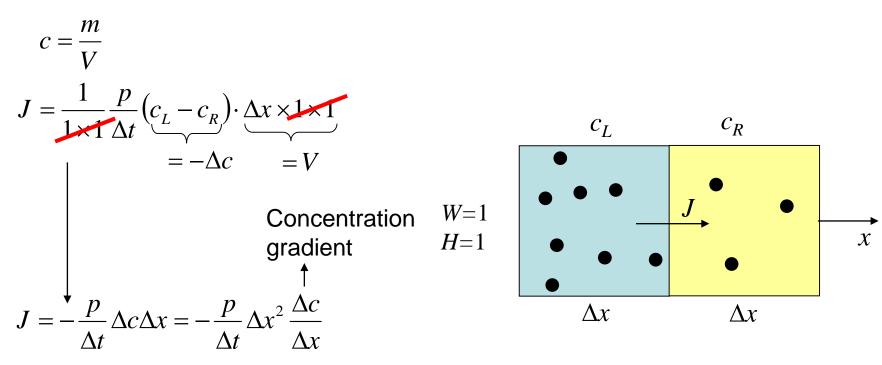


Express in terms of mass concentrations





Express in terms of mass concentrations



$$J = -\frac{p}{\Delta t} \Delta x^{2} \frac{dc}{dx} = -D \frac{dc}{dx}$$

$$D = p \frac{\Delta x^{2}}{\Delta t}$$

Parameter that measures how "fast" mass moves (in square of distance per unit time)

#### Diffusion constant & 1st Fick law

Reiterate: Diffusion constant *D* describes the how much mass moves per unit time

Movement of mass characterized by square of displacement from initial position

Flux
$$J = -\frac{p}{\Delta t} \Delta x^2 \frac{dc}{dx} = -D \frac{dc}{dx}$$

1<sup>st</sup> Fick law (Adolph Fick, 1829-1901)

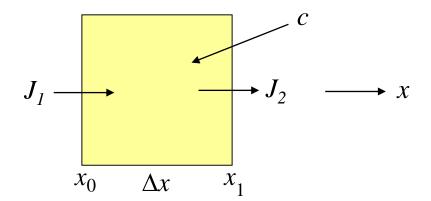
$$D = p \frac{\Delta x^2}{\Delta t}$$

$$D \sim p$$

Diffusion constant relates to the ability of mass to move a distance  $\Delta x^2$  over a time  $\Delta t$  (strongly temperature dependent, e.g. Arrhenius)

## 2<sup>nd</sup> Fick law (time dependence)

$$J = -D\frac{dc}{dx}$$
 1st Fick law

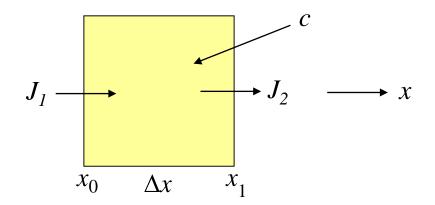


$$\frac{\Delta c}{\Delta t} = \frac{\left(J_1 - J_2\right) \times 1 \times 1}{\Delta x \times 1 \times 1}$$

J: Mass flux (mass per unit time per unit area)

## 2<sup>nd</sup> Fick law (time dependence)

$$J = -D\frac{dc}{dx}$$
 1st Fick law



$$\frac{\Delta c}{\Delta t} = \frac{J_1 - J_2}{\Delta x} = \frac{1}{\Delta x} \left( -D \frac{dc}{dx} \Big|_{x=x_0} - \left[ -D \frac{dc}{dx} \Big|_{x=x_1} \right] \right)$$

$$\int_1 = J(x = x_0) = -D \frac{dc}{dx} \Big|_{x=x_0}$$

# 2<sup>nd</sup> Fick law (time dependence)

$$J = -D\frac{dc}{dx}$$
 1st Fick law

$$J_1 \longrightarrow J_2 \longrightarrow x$$

$$x_0 \quad \Delta x \quad x_1$$

$$\frac{\Delta J}{\Delta t} = \frac{1}{\Delta x} \left( -D \frac{dc}{dx} \Big|_{x=x_0} + D \frac{dc}{dx} \Big|_{x=x_1} \right)$$

$$\frac{\partial c}{\partial t} = -\frac{d}{dx}(J) = -\frac{d}{dx}\left(-D\frac{dc}{dx}\right)$$

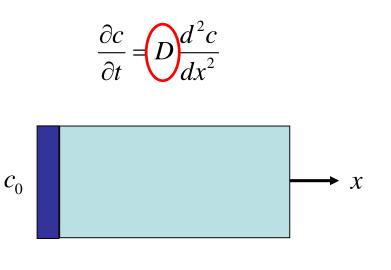
Change of concentration in  $\frac{\partial c}{\partial t} = -\frac{d}{dx}(J) = -\frac{d}{dx}\left(-D\frac{dc}{dx}\right)$  Unange of concentration in time equals change of flux with x (mass balance)

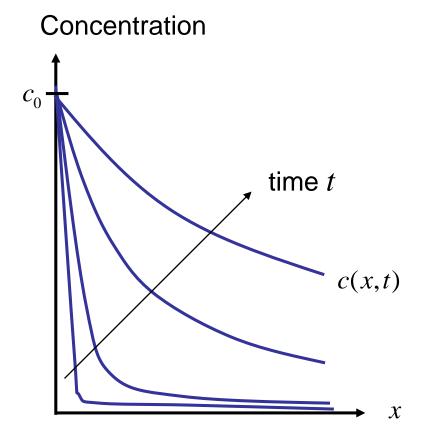
$$\frac{\partial c}{\partial t} = D \frac{d^2 c}{dx^2}$$

2<sup>nd</sup> Fick law

PDE

## Example solution – 2<sup>nd</sup> Fick's law



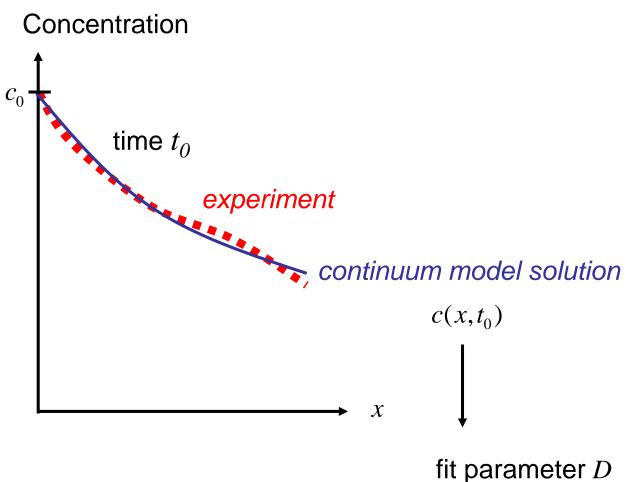


Boundary condition (BC): c (x = 0) =  $c_0$ Initial condition (IC): c (x > 0, t = 0) = 0

#### How to obtain diffusion coefficient?

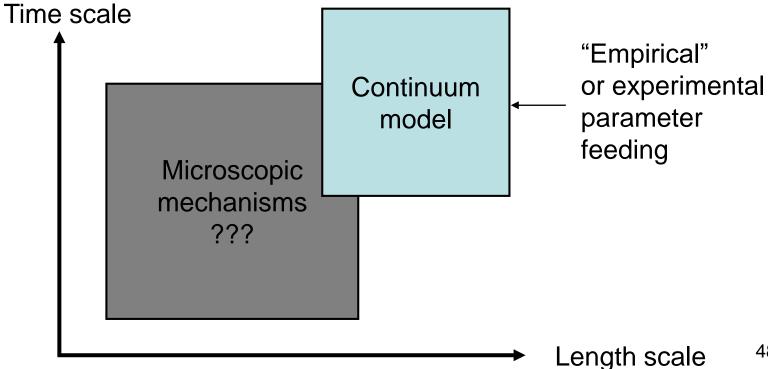
- Laboratory experiment
- Study distribution of concentrations (previous slide)
- Then "fit" the appropriate diffusion coefficient so that the solution matches
- Approach can then be used to solve for more complex geometries, situations etc. for which no lab experiment exists
- "Top down approach"

## Matching with experiment (parameter identification)



## Summary

- Continuum model requires parameter that describes microscopic processes inside the material
- Typically need experimental measurements to calibrate



#### 3.2 Atomistic model

How to build an atomistic bottom-up model to describe the physical phenomena of diffusion?

## Approach 2: Atomistic model

- Atomistic model provides an alternative approach to describe diffusion
- Enables us to directly calculate the diffusion constant from the trajectory of atoms ("microscopic definition")
- Approach: Consider set of atoms/molecules



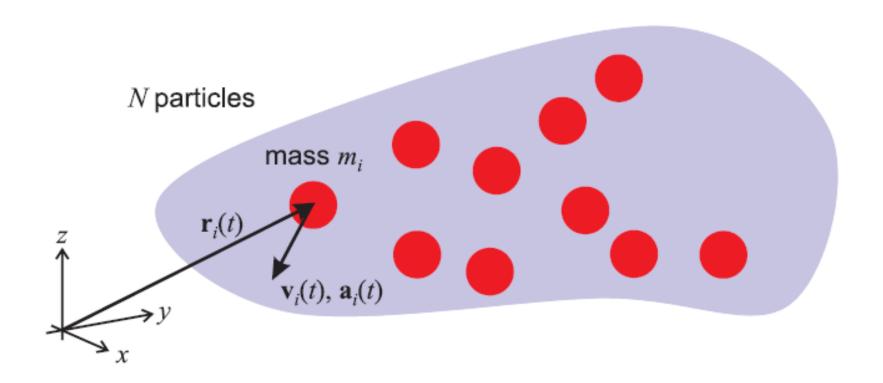
Follow their trajectory and calculate how fast atoms leave their initial position
 Follow this quantity over

time

$$D = p \frac{\Delta x^2}{\Delta t}$$

**Recall**: Diffusion constant relates to the "ability" of particle to move a distance  $\Delta x^2$  over a time  $\Delta t$ 

## Molecular dynamics – simulate trajectory of atoms



**Goal:** Need an algorithm to predict positions, velocities, accelerations as function of time

## Solving the equations: What we want

To solve those equations: Discretize in time (n steps),  $\Delta t$  time step:

$$r_i(t_0) \rightarrow r_i(t_0 + \Delta t) \rightarrow r_i(t_0 + 2\Delta t) \rightarrow r_i(t_0 + 3\Delta t) \rightarrow \dots \rightarrow r_i(t_0 + n\Delta t)$$

## Solving the equations

To solve those equations: Discretize in time (n steps),  $\Delta t$  time step:

$$r_i(t_0) \rightarrow r_i(t_0 + \Delta t) \rightarrow r_i(t_0 + 2\Delta t) \rightarrow r_i(t_0 + 3\Delta t) \rightarrow \dots \rightarrow r_i(t_0 + n\Delta t)$$

**Recall**: Taylor expansion of function *f* around point *a* 

$$f(x) = f(a) + f'(a)(x - a) + \frac{f''(a)}{2!}(x - a)^2 + \frac{f^{(3)}(a)}{3!}(x - a)^3 + \dots + \frac{f^{(n)}(a)}{n!}(x - a)^n + \dots$$

## Solving the equations

To solve those equations: Discretize in time (n steps),  $\Delta t$  time step:

$$r_i(t_0) \rightarrow r_i(t_0 + \Delta t) \rightarrow r_i(t_0 + 2\Delta t) \rightarrow r_i(t_0 + 3\Delta t) \rightarrow \dots \rightarrow r_i(t_0 + n\Delta t)$$

**Recall**: Taylor series expansion of function *f* around point *a* 

$$f(x) = f(a) + f'(a)(x - a) + \frac{f''(a)}{2!}(x - a)^2 + \frac{f^{(3)}(a)}{3!}(x - a)^3 + \dots + \frac{f^{(n)}(a)}{n!}(x - a)^n + \dots$$

Taylor series expansion  $r_i(t)$  around  $a = t_0$   $x = t_0 + \Delta t$ 

$$a = t_0 x = t_0 + \Delta t$$

$$x - a = t_0 + \Delta t - t_0 = \Delta t$$

## Solving the equations

To solve those equations: Discretize in time (n steps),  $\Delta t$  time step:

$$r_i(t_0) \rightarrow r_i(t_0 + \Delta t) \rightarrow r_i(t_0 + 2\Delta t) \rightarrow r_i(t_0 + 3\Delta t) \rightarrow \dots \rightarrow r_i(t_0 + n\Delta t)$$

**Recall**: Taylor series expansion of function f around point a

$$f(x) = f(a) + f'(a)(x - a) + \frac{f''(a)}{2!}(x - a)^2 + \frac{f^{(3)}(a)}{3!}(x - a)^3 + \dots + \frac{f^{(n)}(a)}{n!}(x - a)^n + \dots$$

Taylor series expansion 
$$r_i(t)$$
 around 
$$a = t_0 x = t_0 + \Delta t$$
$$x - a = t_0 + \Delta t - t_0 = \Delta t$$

$$r_i(t_0 + \Delta t) = r_i(t_0) + v_i(t_0)\Delta t + \frac{1}{2}a_i(t_0)\Delta t^2 + \dots$$

# Taylor expansion of $r_i(t)$

$$r_i(t_0 + \Delta t) = r_i(t_0) + v_i(t_0)\Delta t + \frac{1}{2}a_i(t_0)\Delta t^2 + \dots$$
$$r_i(t_0 - \Delta t) = r_i(t_0) - v_i(t_0)\Delta t + \frac{1}{2}a_i(t_0)\Delta t^2 + \dots$$

$$a = t_0$$
  $x = t_0 + \Delta t$   $x - a = t_0 + \Delta t - t_0 = \Delta t$   $a = t_0$   $x = t_0 - \Delta t$   $x - a = t_0 - \Delta t - t_0 = -\Delta t$ 

## Taylor expansion of $r_i(t)$

$$r_{i}(t_{0} + \Delta t) = r_{i}(t_{0}) + v_{i}(t_{0})\Delta t + \frac{1}{2}a_{i}(t_{0})\Delta t^{2} + \dots$$

$$+ \left[r_{i}(t_{0} - \Delta t) = r_{i}(t_{0}) - v_{i}(t_{0})\Delta t + \frac{1}{2}a_{i}(t_{0})\Delta t^{2} + \dots\right]$$

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

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

Positions Positions Accelerations at  $t_0$  at  $t_0$ - $\Delta t$  at  $t_0$ 

## Physics of particle interactions

#### Laws of Motion of Isaac Newton (1642 – 1727):

- Every body continues in its state of rest, or of uniform motion in a right line, unless it is compelled to change that state by forces impressed upon it.
- 2. The change of motion is proportional to the motive force impresses, and is made in the direction of the right line in which that force is impressed.
- 3. To every action there is always opposed an equal reaction: or, the mutual action of two bodies upon each other are always equal, and directed to contrary parts.

$$f = m\frac{d^2x}{dt^2} = ma$$
 2<sup>nd</sup> law

#### Verlet central difference method

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

How to obtain accelerations?

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

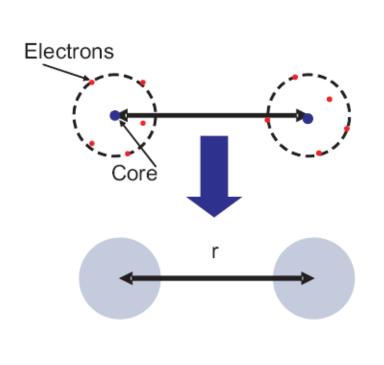
Need forces on atoms!

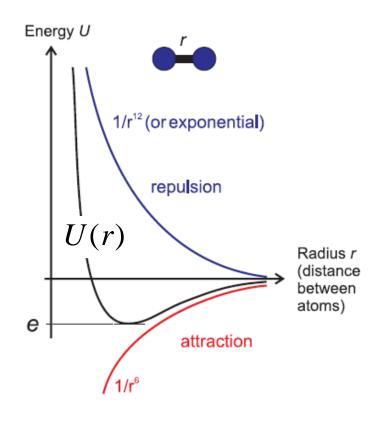
#### Verlet central difference method

$$r_i(t_0 + \Delta t) = 2r_i(t_0) - r_i(t_0 - \Delta t) + f_i(t_0) / m\Delta t^2 + \dots$$
Positions at  $t_0$  Positions at  $t_0$  at  $t_0$ 
Lecture 3

#### Preview: Forces on atoms

Consider energy landscape due to chemical bonds





Attraction: Formation of chemical bond by sharing of electrons

**Repulsion:** Pauli exclusion (too many electrons in small volume)

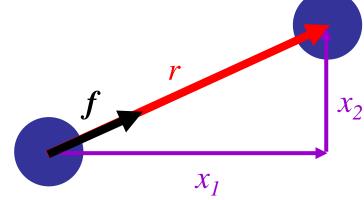
#### How are forces calculated?

Force magnitude: Derivative of potential energy with respect to atomic distance

$$f = -\frac{\mathrm{d}U(r)}{\mathrm{d}r}$$

To obtain force vector  $f_i$ , take projections into the three axial directions

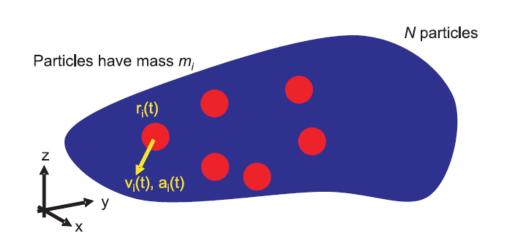
$$f_i = f \frac{x_i}{r}$$



#### Note on force calculation

- Forces can be obtained from a variety of models for interatomic energy, e.g.
  - Pair potentials (e.g. LJ, Morse, Buckingham)
  - Multi-body potentials (e.g. EAM, CHARMM, UFF, DREIDING)
  - Reactive potentials (e.g. ReaxFF)
  - Quantum mechanics (e.g. DFT) part II
  - Tight-binding
  - **.** . . .
- ...will be discussed in next lectures

## Molecular dynamics



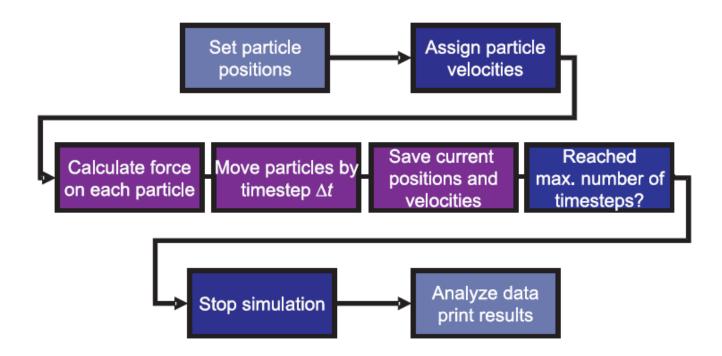
Follow trajectories of atoms

"Verlet central difference method"

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

# Summary: Atomistic simulation – numerical approach "molecular dynamics – MD"

- Atomistic model; requires atomistic microstructure and atomic position at beginning
- Step through time by integration scheme
- Repeated force calculation of atomic forces
- Explicit notion of chemical bonds captured in interatomic potential



#### Pseudocode

```
Set particle positions (e.g. crystal lattice)
Assign initial velocities
For (all time steps):
   Calculate force on each particle (subroutine)
   Move particle by time step \Delta t
   Save particle position, velocity, acceleration
Save results
                                                             Assign particle
                                             Set particle
Stop simulation
                                                              velocities
                                              positions
                                                          Save current
                                                                       Reached
                                   Calculate force Move particles by
                                                          positions and
                                                                     max, number of
                                   on each particle
                                               timestep ∆t
                                                           velocities
                                                                      timesteps?
```

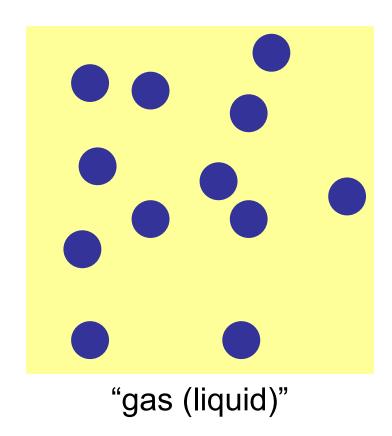
Stop simulation

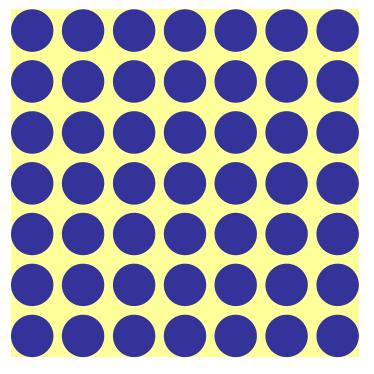
Analyze data

print results

# Atomic positions (initial conditions)

 Typically, have cubical cell in which particles are places in a regular or irregular manner





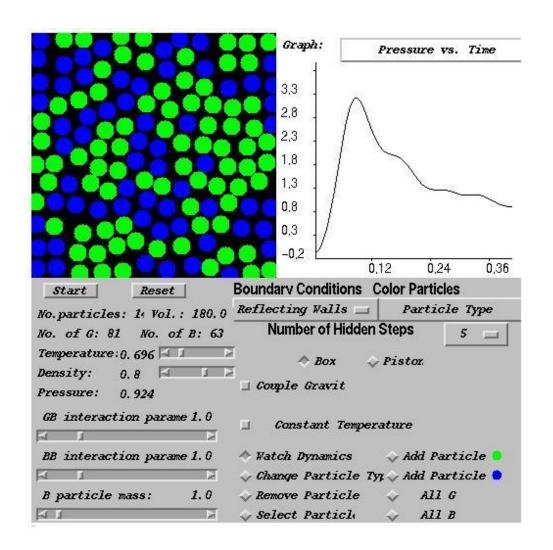
"solid - crystal"

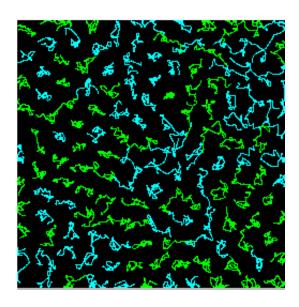
## Atomistic description

Back to the application of diffusion problem...

- Atomistic description provides alternative way to predict D
- Simple solve equation of motion
- Follow the trajectory of an atom
- Relate the average distance as function of time from initial point to diffusivity
- Goal: Calculate how particles move "randomly", away from initial position

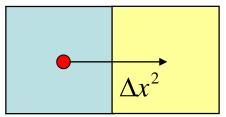
## JAVA applet





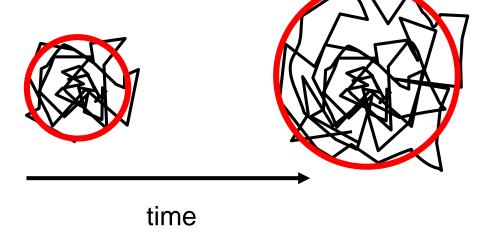
Diffusion constant relates to the "ability" of a particle to move a distance  $\Delta x^2$  (from left to right) over a time  $\Delta t$ 

$$D = p \frac{\Delta x^2}{\Delta t}$$



**Idea** – Use MD simulation to measure square of displacement from initial position of particles,  $\Delta r^2(t)$ :

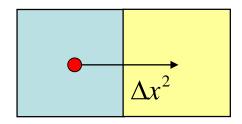




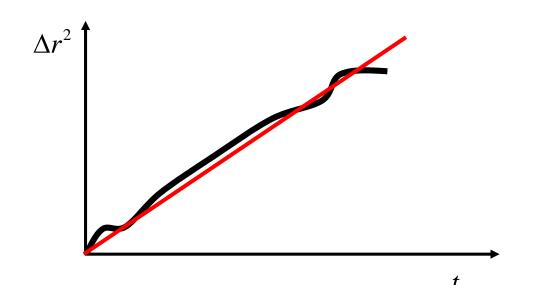
70

Diffusion constant relates to the "ability" of a particle to move a distance  $\Delta x^2$  (from left to right) over a time  $\Delta t$ 

$$D = p \frac{\Delta x^2}{\Delta t}$$



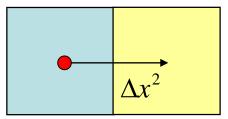
**MD simulation:** Measure square of displacement from initial position of particles,  $\Delta r^2(t)$ :



71

Diffusion constant relates to the "ability" of a particle to move a distance  $\Delta x^2$  (from left to right) over a time  $\Delta t$ 

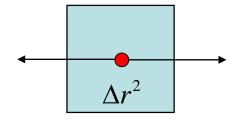
$$D = p \frac{\Delta x^2}{\Delta t}$$



**MD simulation:** Measure square of displacement from initial position of particles,  $\Delta r^2(t)$  and not  $\Delta x^2(t)$  ....

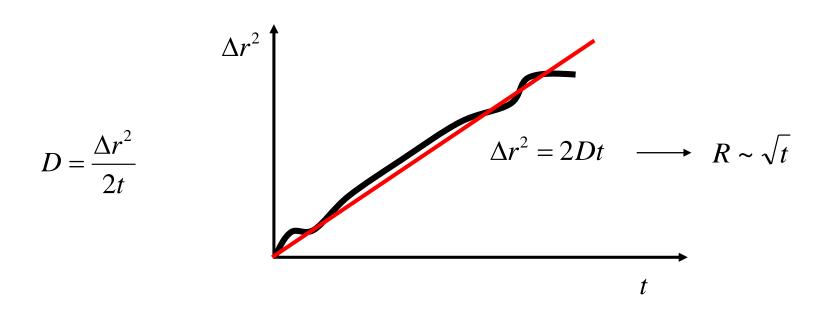
Replace

$$D = p \frac{\Delta x^2}{\Delta t} \longrightarrow D = \frac{1}{2} \underbrace{\Delta r^2}_{\Delta t}$$



Factor 1/2 = no directionality in (equal probability to move forth or back)

**MD simulation:** Measure square of displacement from initial position of particles,  $\Delta r^2(t)$ :



$$D = p \frac{\Delta x^2}{\Delta t}$$
Higher dimensions

$$D = \frac{1}{2} \frac{1}{d} \frac{\Delta r^2}{\Delta t}$$

Factor 1/2 = no directionality in (forth/back)

Factor d = 1, 2, or 3 due to 1D, 2D, 3D (dimensionality)

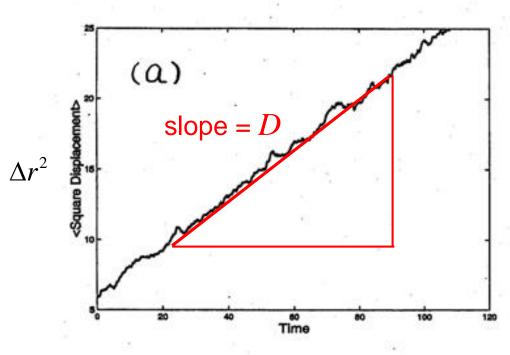
Since:

$$2dD\Delta t \sim \Delta r^2$$

$$2dD\Delta t + C = \Delta r^2$$

C = constant (does not affect D)

## Example: MD simulation



$$D = \frac{1}{2d} \lim_{t \to \infty} \frac{d}{dt} (\Delta r^2)$$

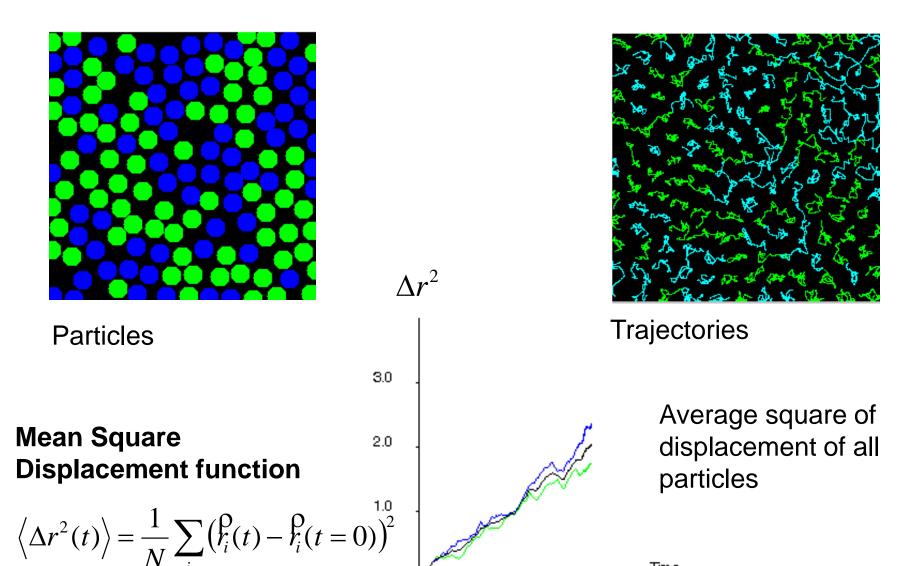
$$\uparrow$$

$$1D=1, 2D=2, 3D=3$$

$$D = \frac{1}{2d} \lim_{t \to \infty} \frac{\mathrm{d}}{\mathrm{d}t} \left\langle \Delta r^2 \right\rangle$$

 $\langle ... \rangle$  = average over all particles

## Example molecular dynamics



6.4

12.8

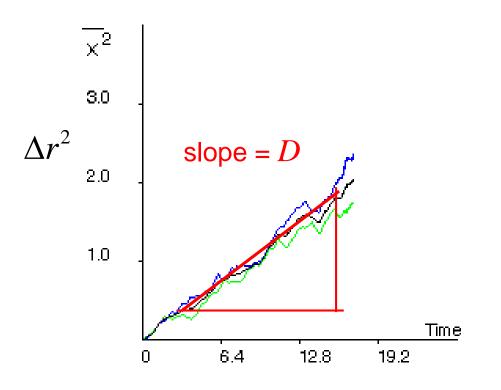
Time

19.2

76

## Example calculation of diffusion coefficient

$$\left\langle \Delta r^2(t) \right\rangle = \frac{1}{N} \sum_{i} \left( r_i(t) - r_i(t=0) \right)^2$$
Position of Position of atom *i* at time t atom *i* at time t=0



$$D = \frac{1}{2d} \lim_{t \to \infty} \frac{d}{dt} \langle \Delta r^2 \rangle$$

$$1D=1, 2D=2, 3D=3$$

## Summary: "Multiscale simulation paradigm"

- Molecular dynamics provides a powerful approach to relate the diffusion constant that appears in continuum models to atomistic trajectories
- Outlines multi-scale approach: Feed parameters from atomistic simulations to continuum models

