

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

**Part I – Continuum and particle methods**

# **Basic molecular dynamics**

*Lecture 2*

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**Massachusetts Institute of Technology**

# Content overview

## I. Particle and continuum methods

Lectures 2-13

1. Atoms, molecules, chemistry
2. Continuum modeling approaches and solution approaches
3. 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?

# Goals of part I (particle methods)

You will be able to ...

- **Carry out atomistic simulations** of various processes (diffusion, deformation/stretching, materials failure)

*Carbon nanotubes, nanowires, bulk metals, proteins, silicon crystals, etc.*

- **Analyze** atomistic simulations (make sense of all 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

# Lecture 2: Basic molecular dynamics

## Outline:

1. Introduction
2. Case study: Diffusion
  - 2.1 Continuum model
  - 2.2 Atomistic model
3. Additional remarks – historical perspective

## 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
- 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. Introduction

# Relevant scales in materials

$\lambda(\text{atom}) \ll \xi(\text{crystal}) \ll d(\text{grain}) \ll dx, dy, dz \ll H, W, D$

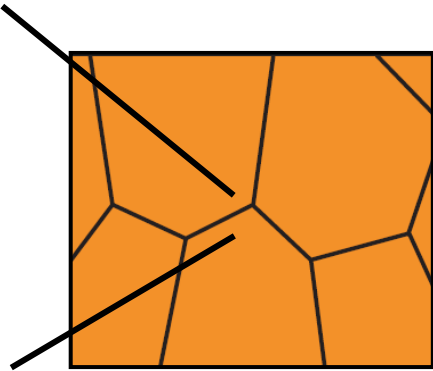
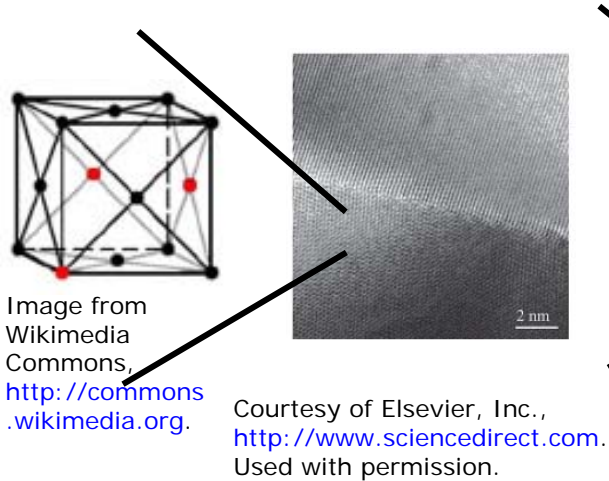
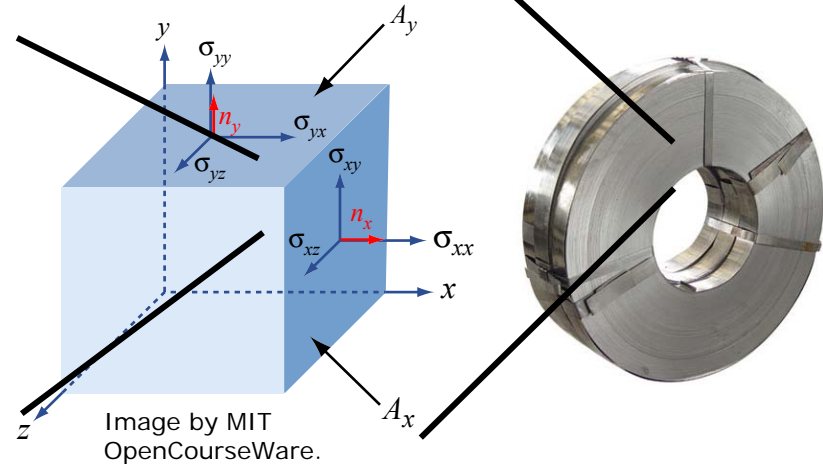


Fig. 8.7 in: Buehler, M. *Atomistic Modeling of Materials Failure*. Springer, 2008. © Springer. All rights reserved. This content is excluded from our Creative Commons license. For more information, see <http://ocw.mit.edu/fairuse>.



**Bottom-up** →

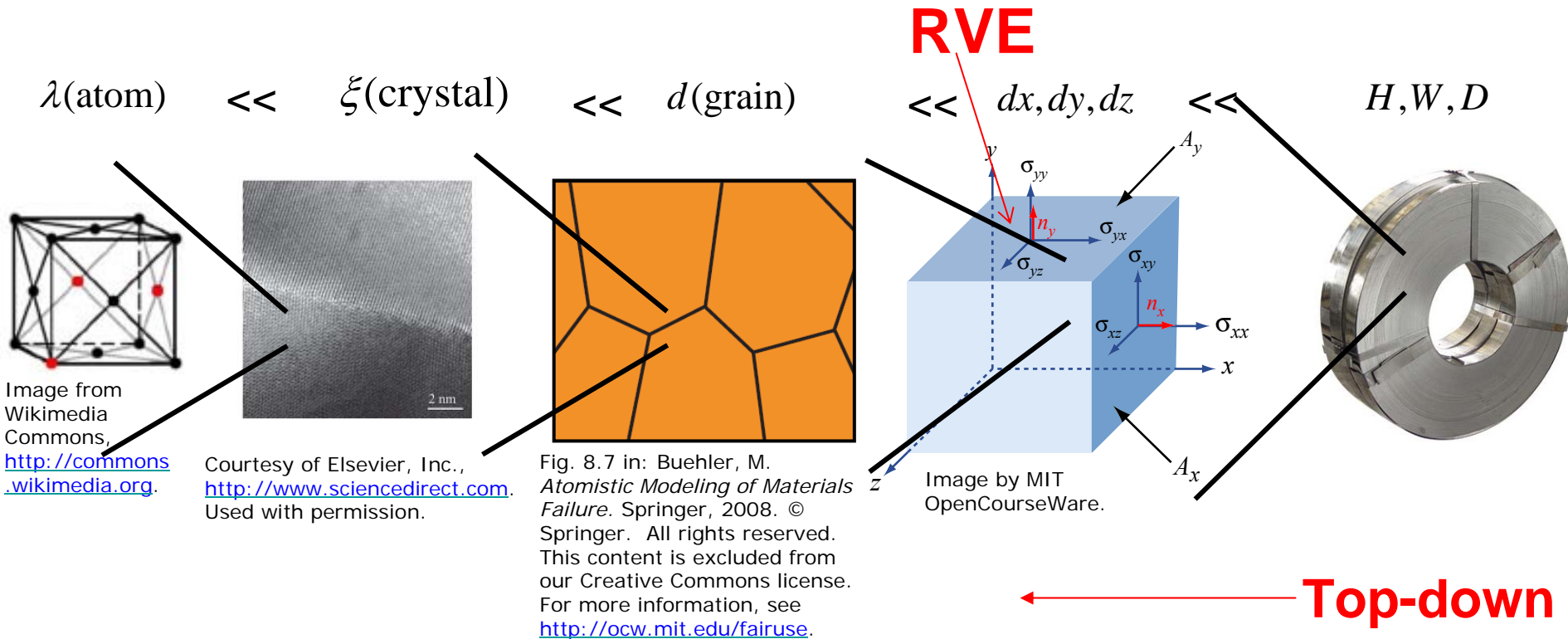
← **Top-down**

## 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)
- Features no characteristic length scales, provided RVE is large enough <sup>7</sup>

“PDE with parameters”

## 2. Case study: Diffusion

*Continuum and atomistic modeling*



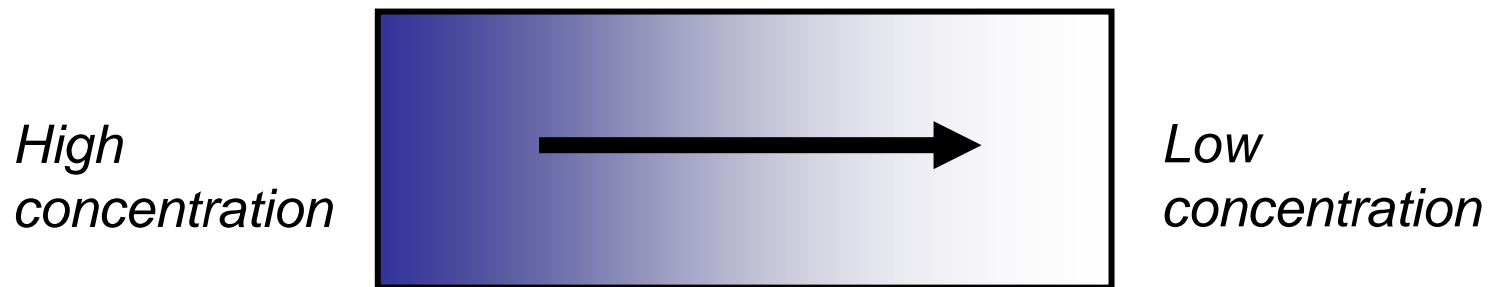
# Goal of this section

- 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*

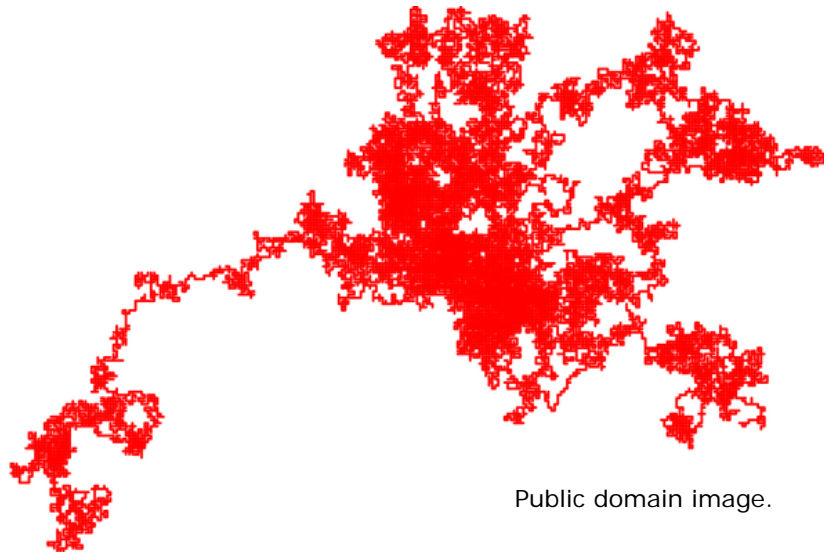
*cold*

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# *Microscopic observation of diffusion*

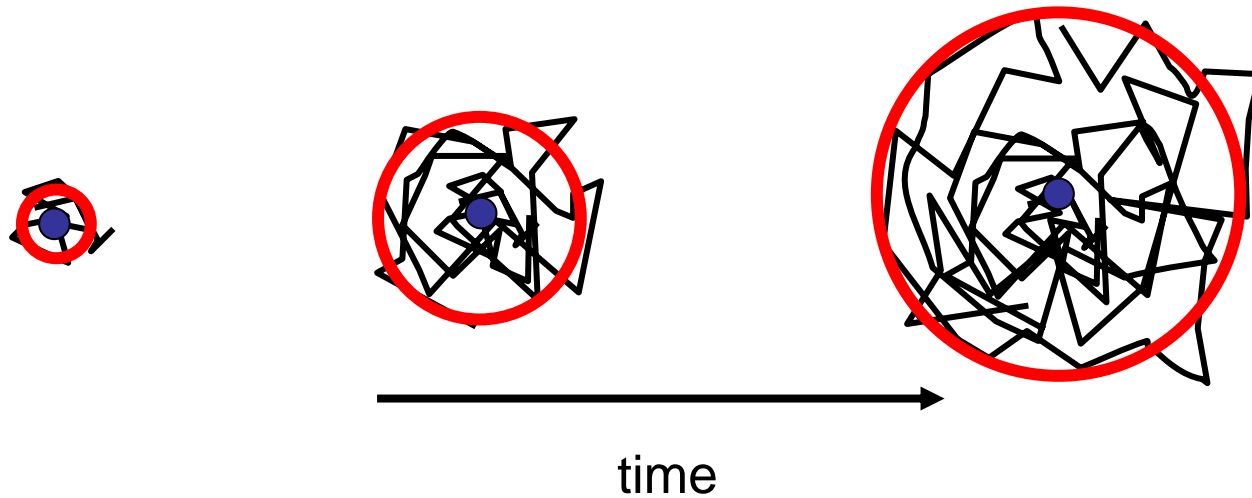
# Microscopic mechanism: “Random walk” – or Brownian motion

- 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.



Public domain image.

# Brownian motion leads to net particle movement



*Particle “slowly” moves away from its initial position*

# Robert Brown's microscope

**Robert Brown's Microscope** 1827

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*

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

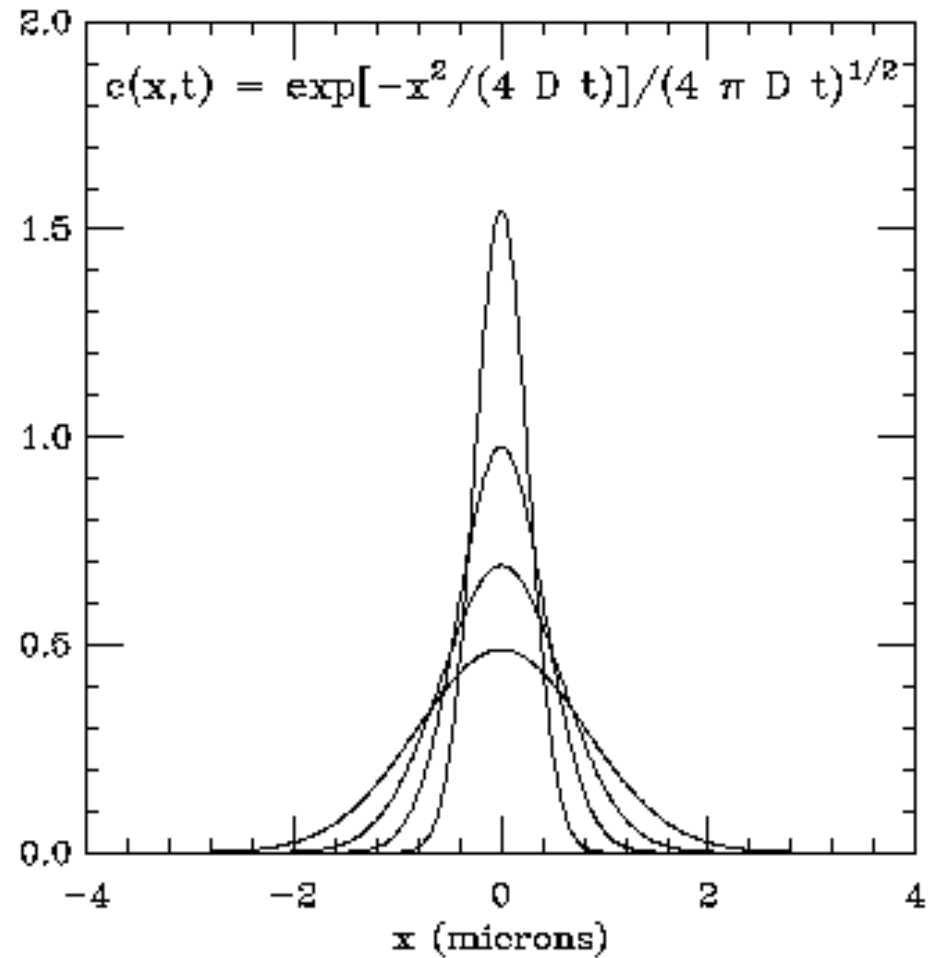
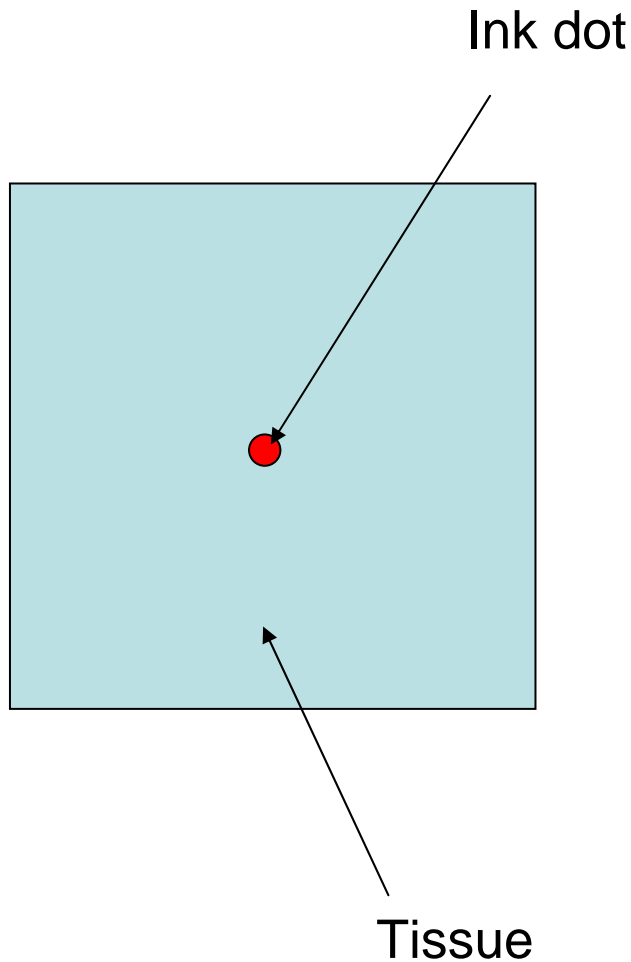
# Simulation of Brownian motion

- [http://www.scienceisart.com/A\\_Diffus/Jav1\\_2.html](http://www.scienceisart.com/A_Diffus/Jav1_2.html)



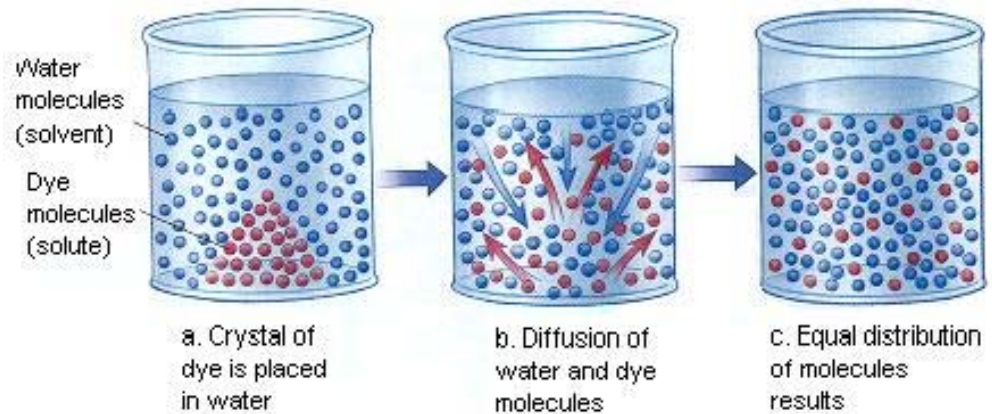
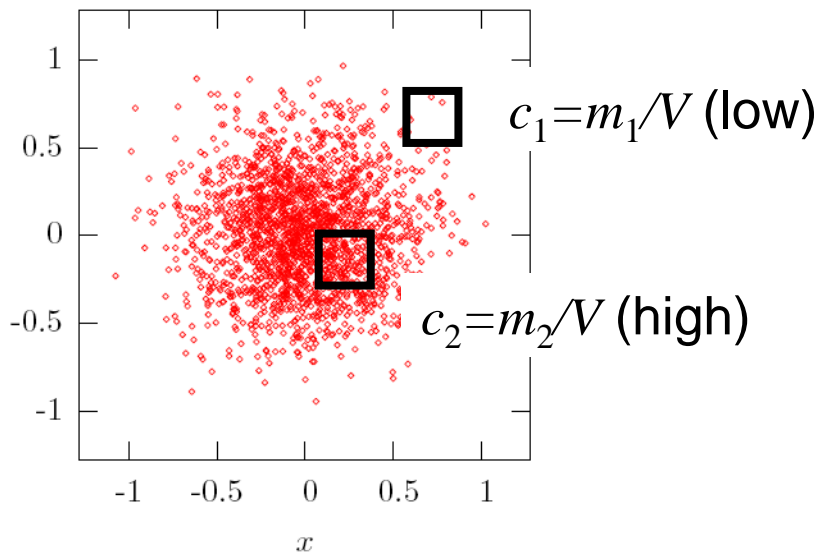
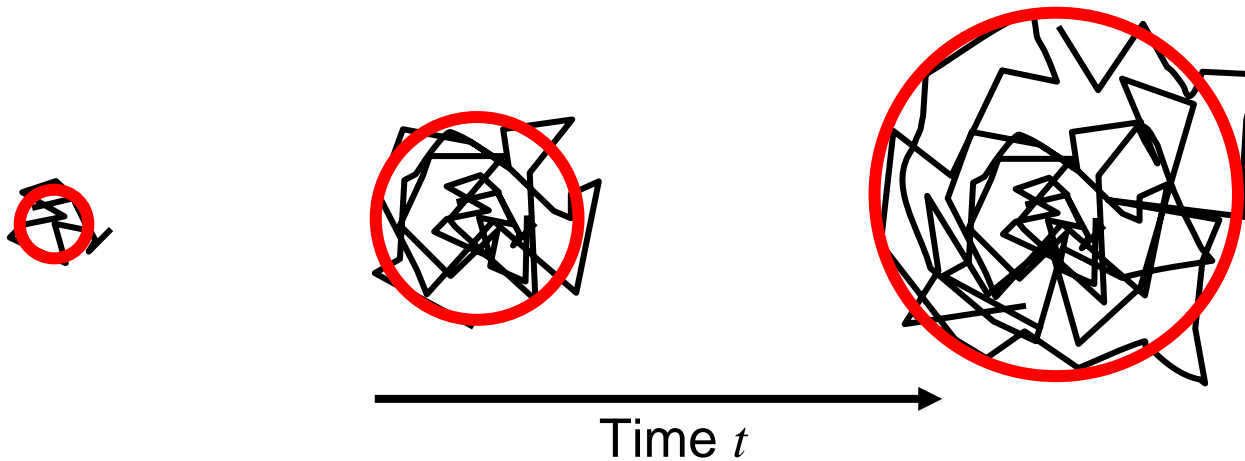
# *Macroscopic observation of diffusion*

# Macroscopic observation: concentration change



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# Brownian motion leads to net particle movement



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# Diffusion in biology

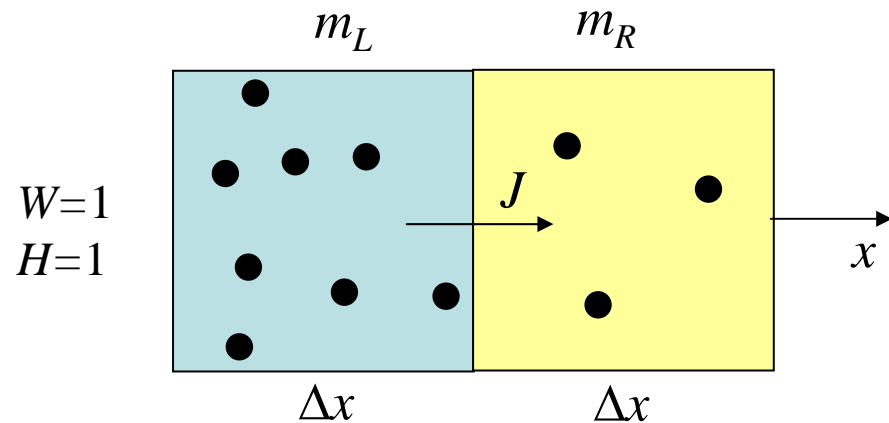
Image removed due to copyright restrictions. See the image now: <http://www.biog1105-1106.org/demos/105/unit9/media/diffusion-after-act-pot.jpg>.

## 2.1 Continuum model

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

# Approach 1: Continuum model

- Develop differential equation based on differential element

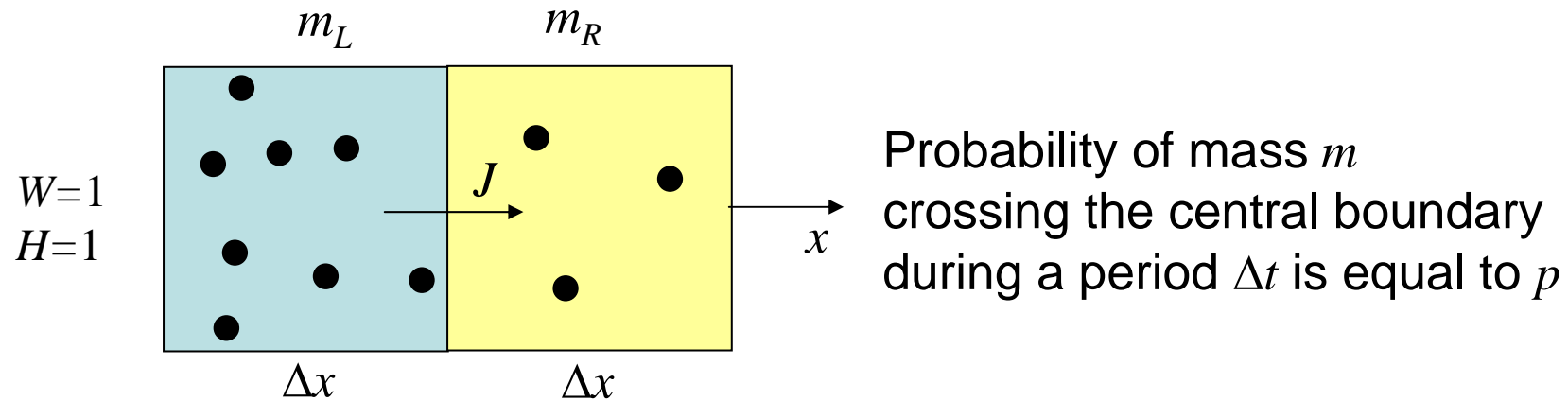


$J$ : Mass flux (mass per unit time per unit area)

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

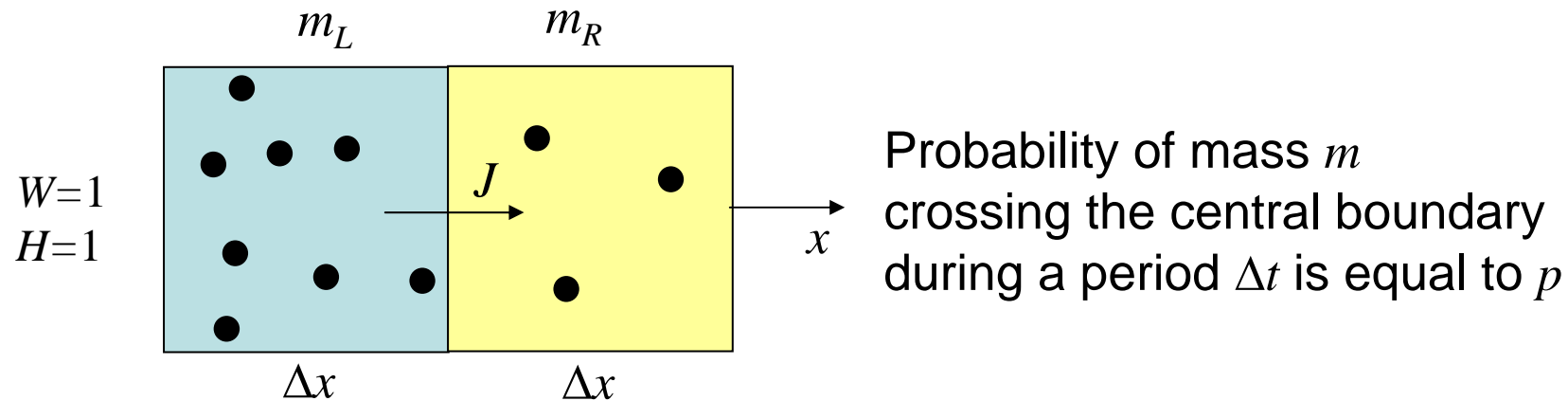
# Approach 1: Continuum model

- Develop differential equation based on differential element



# Approach 1: Continuum model

- Develop differential equation based on differential element



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

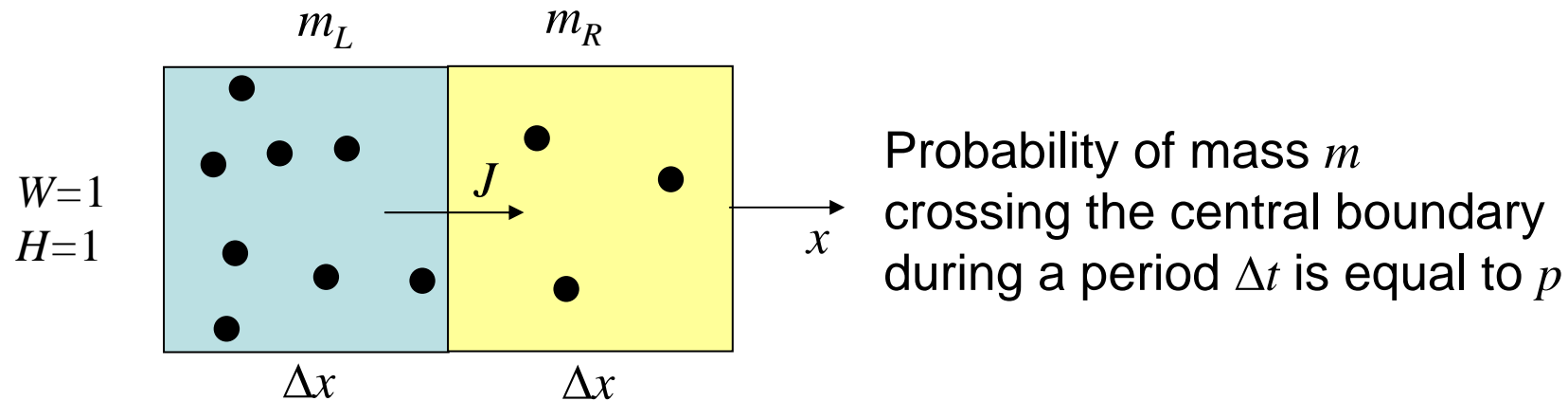
$$J_R = \frac{1}{1 \times 1} \frac{p}{\Delta t} m_R \quad \text{Mass flux from right to left}$$

$[J]$  = mass per unit time per unit area



# Approach 1: Continuum model

- Develop differential equation based on differential element



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

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

Effective mass flux

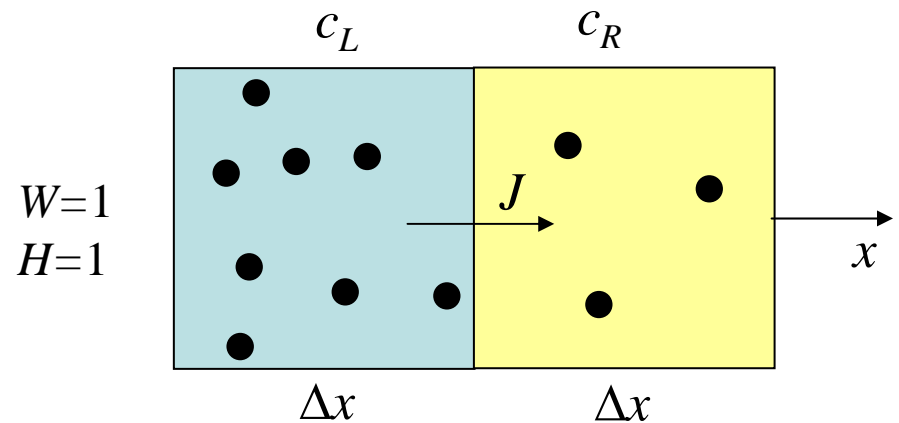
$$J = \frac{1}{1 \times 1} \frac{p}{\Delta t} (m_L - m_R)$$

*More mass, more flux ( $m_L$  is  $\sim$  to number of particles)*

# Continuum model of diffusion

Express in terms of mass concentrations

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

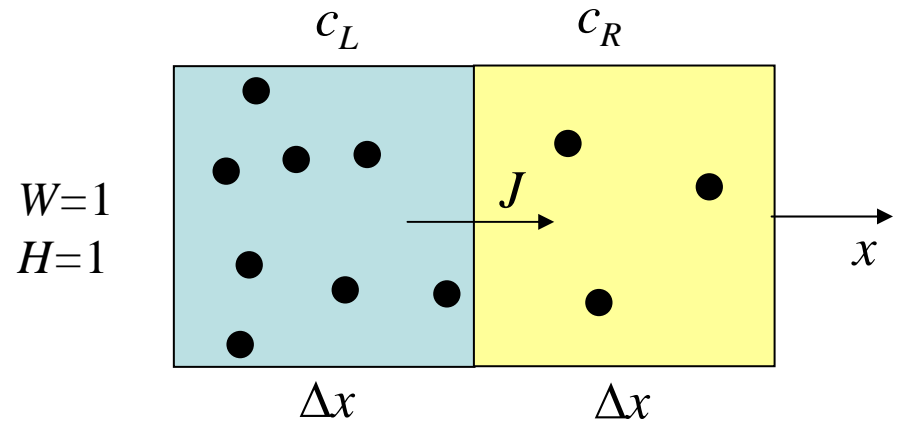


# Continuum model of diffusion

Express in terms of mass concentrations

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

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



# Continuum model of diffusion

Express in terms of mass concentrations

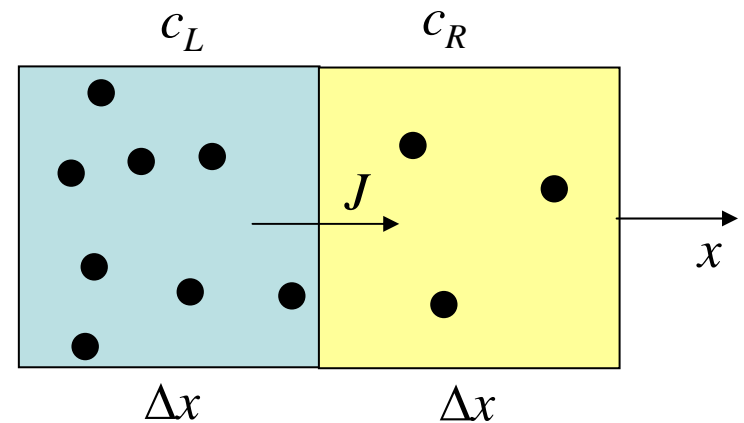
$$c = \frac{m}{V}$$

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

$$J = -\frac{p}{\Delta t} \Delta c \Delta x = -\frac{p}{\Delta t} \Delta x^2 \frac{\Delta c}{\Delta x}$$

Concentration gradient

*expand  $\Delta x$*



$W=1$   
 $H=1$

# Continuum model of diffusion

Express in terms of mass concentrations

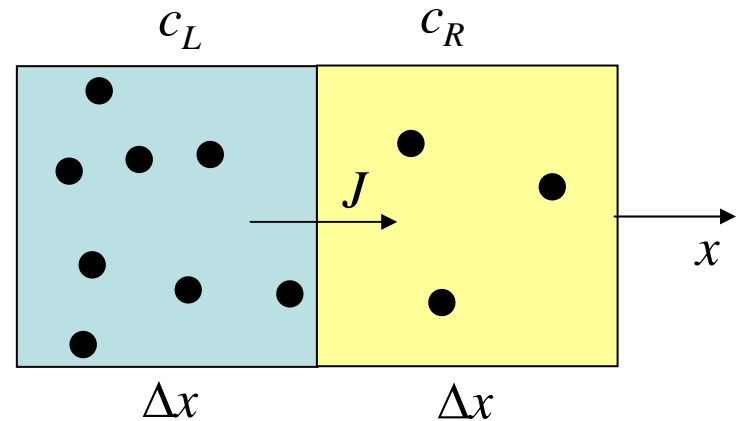
$$c = \frac{m}{V}$$

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

$$J = -\frac{p}{\Delta t} \Delta c \Delta x = -\frac{p}{\Delta t} \Delta x^2 \frac{\Delta c}{\Delta x}$$

Concentration gradient

$W=1$   
 $H=1$



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

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

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

# Diffusion constant & 1<sup>st</sup> 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)

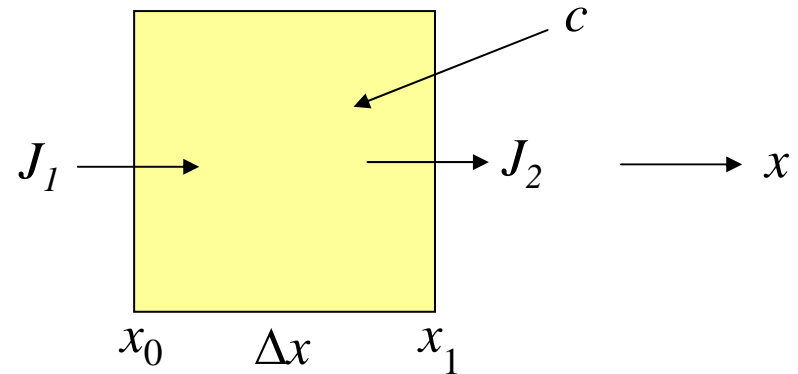
$$D = p \frac{\Delta x^2}{\Delta t}$$

$$D \sim p$$

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)

$$J = -D \frac{dc}{dx} \quad \text{1<sup>st</sup> Fick law}$$



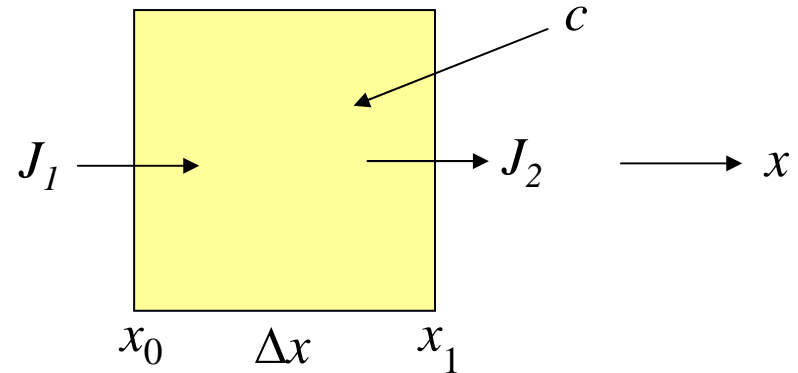
$$\frac{\Delta c}{\Delta t} = \frac{(J_1 - J_2) \times \cancel{1 \times 1}}{\Delta x \times \cancel{1 \times 1}}$$

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

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

$$J = -D \frac{dc}{dx}$$

1<sup>st</sup> Fick law



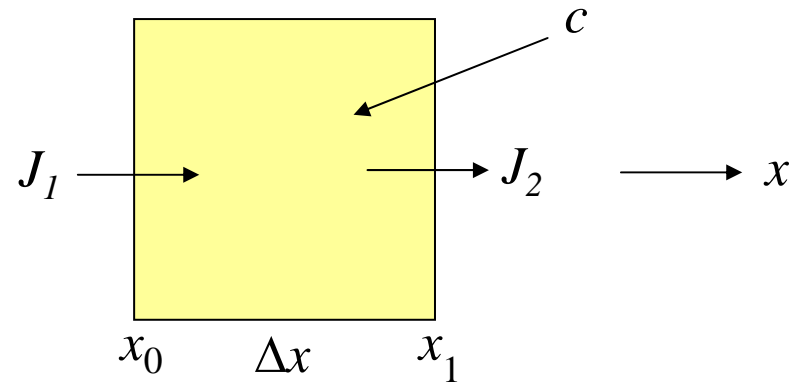
$$\frac{\Delta c}{\Delta t} = \frac{J_1 - J_2}{\Delta x} = \frac{1}{\Delta x} \left( -D \frac{dc}{dx} \Big|_{x=x_0} - \left[ -D \frac{dc}{dx} \Big|_{x=x_1} \right] \right)$$

$$J_1 = J(x = x_0) = -D \frac{dc}{dx} \Big|_{x=x_0}$$



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

$$J = -D \frac{dc}{dx} \quad \text{1<sup>st</sup> Fick law}$$



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

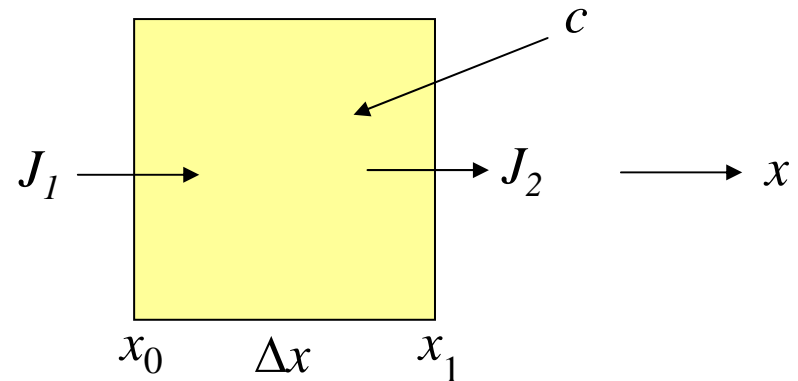
$$\downarrow$$

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

Change of concentration in time equals change of flux with  $x$  (mass balance)

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

$$J = -D \frac{dc}{dx} \quad \text{1<sup>st</sup> Fick law}$$



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

$$\downarrow$$

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

Change of concentration in time equals change of flux with  $x$  (mass balance)

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

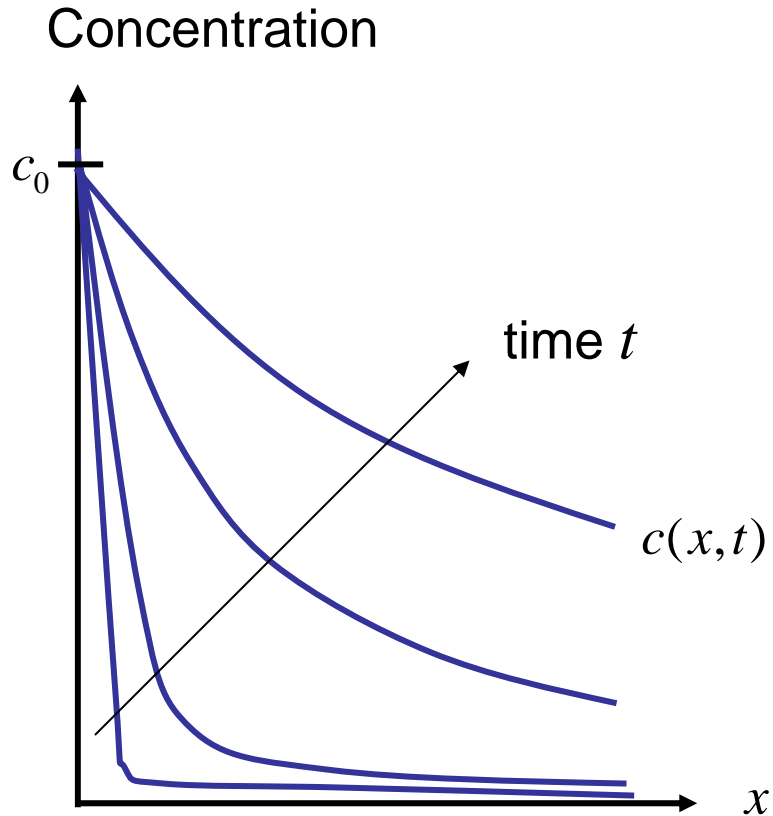
2<sup>nd</sup> Fick law

PDE

Solve by applying ICs and BCs...

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

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



BC:  $c(x=0) = c_0$

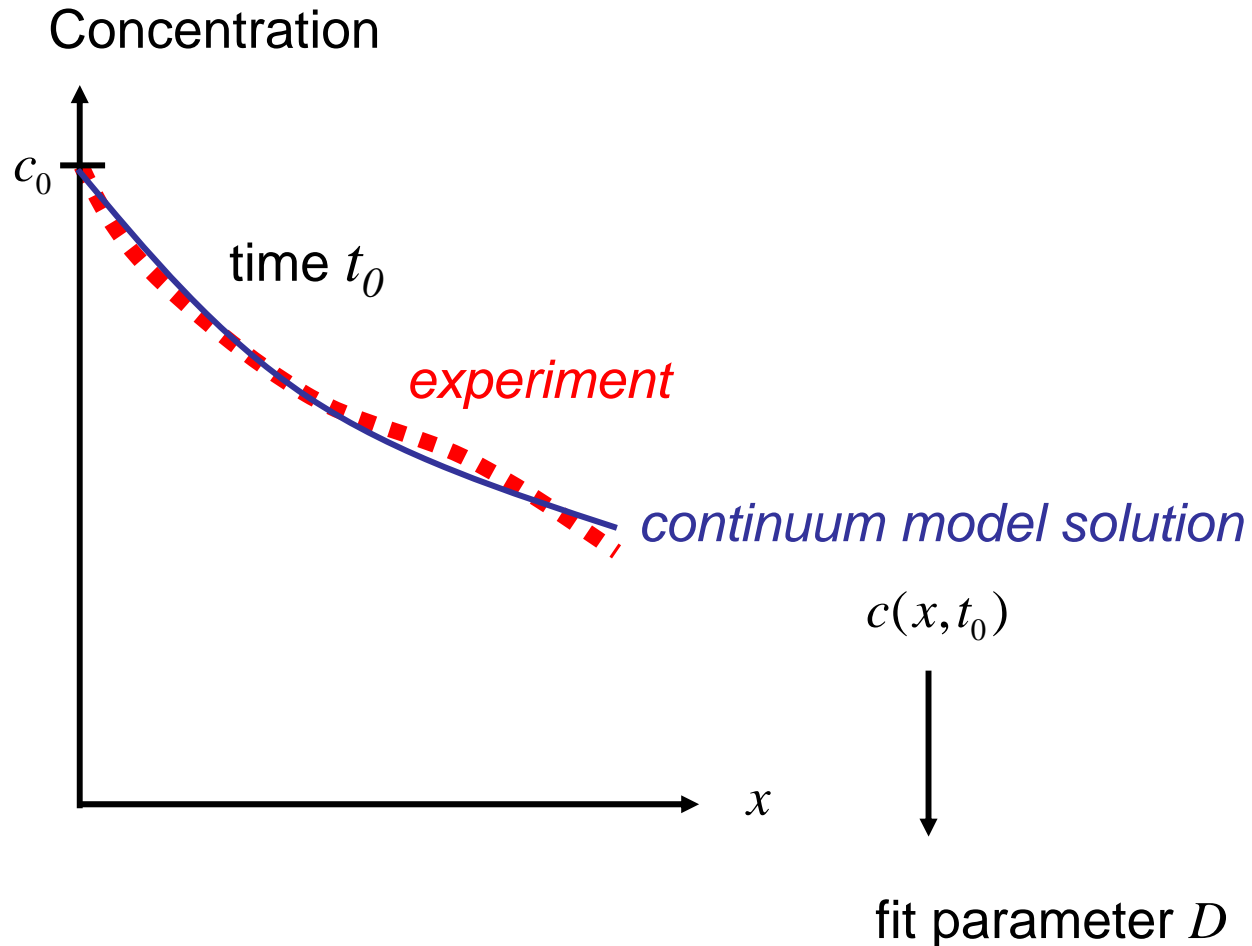
IC:  $c(x > 0, t = 0) = 0$

***Need diffusion coefficient to solve for distribution!***

# How to obtain diffusion coefficient?

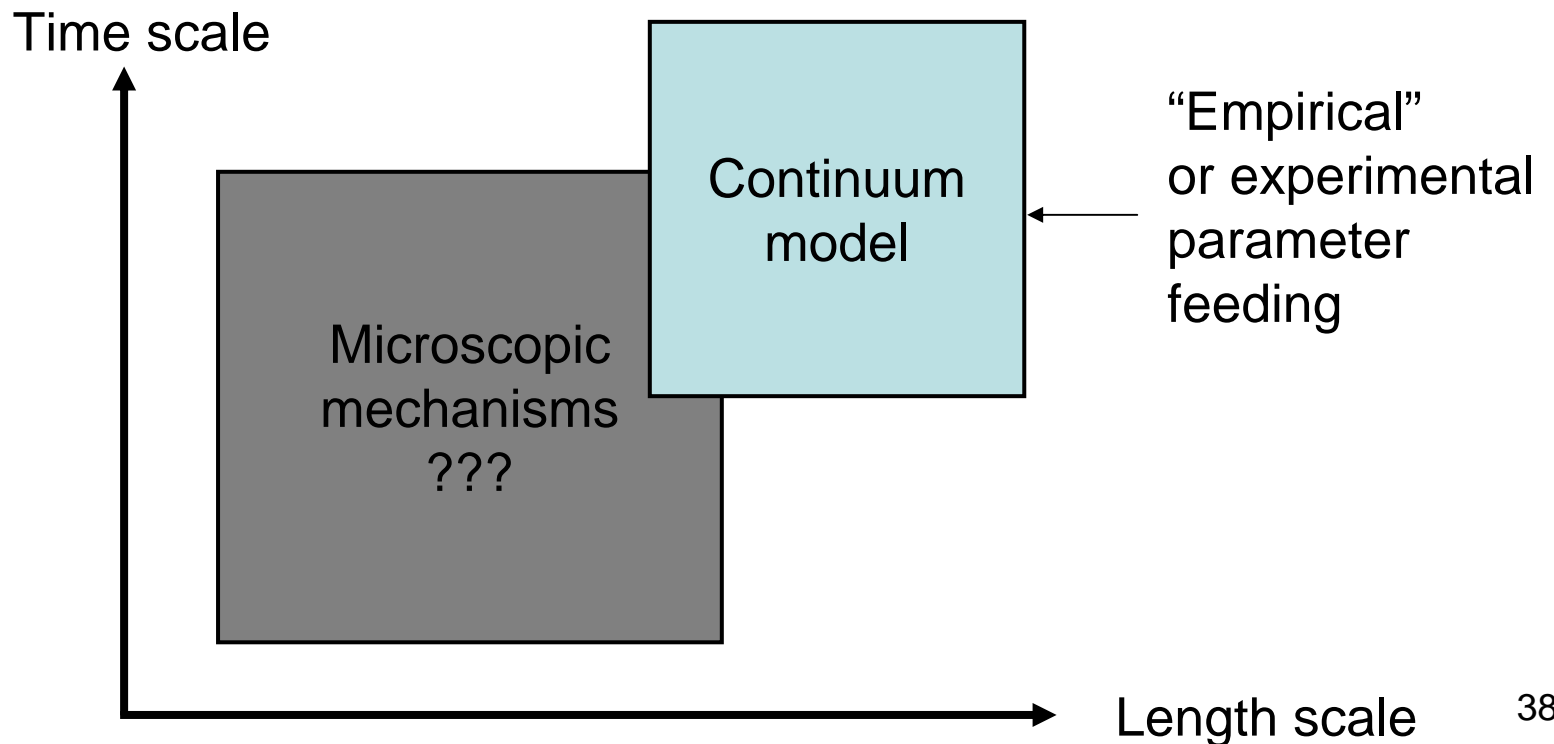
- Laboratory experiment
- Study distribution of concentrations (previous slide)
- Then “fit” the appropriate diffusion coefficient so that the solution matches
- Approach can then be used to solve for more complex geometries, situations etc. for which no lab experiment exists
- “Top down approach”

# Matching with experiment (parameter identification)



# Summary

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

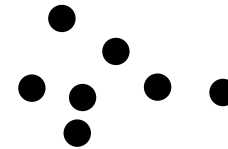


## 2.2 Atomistic model

*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



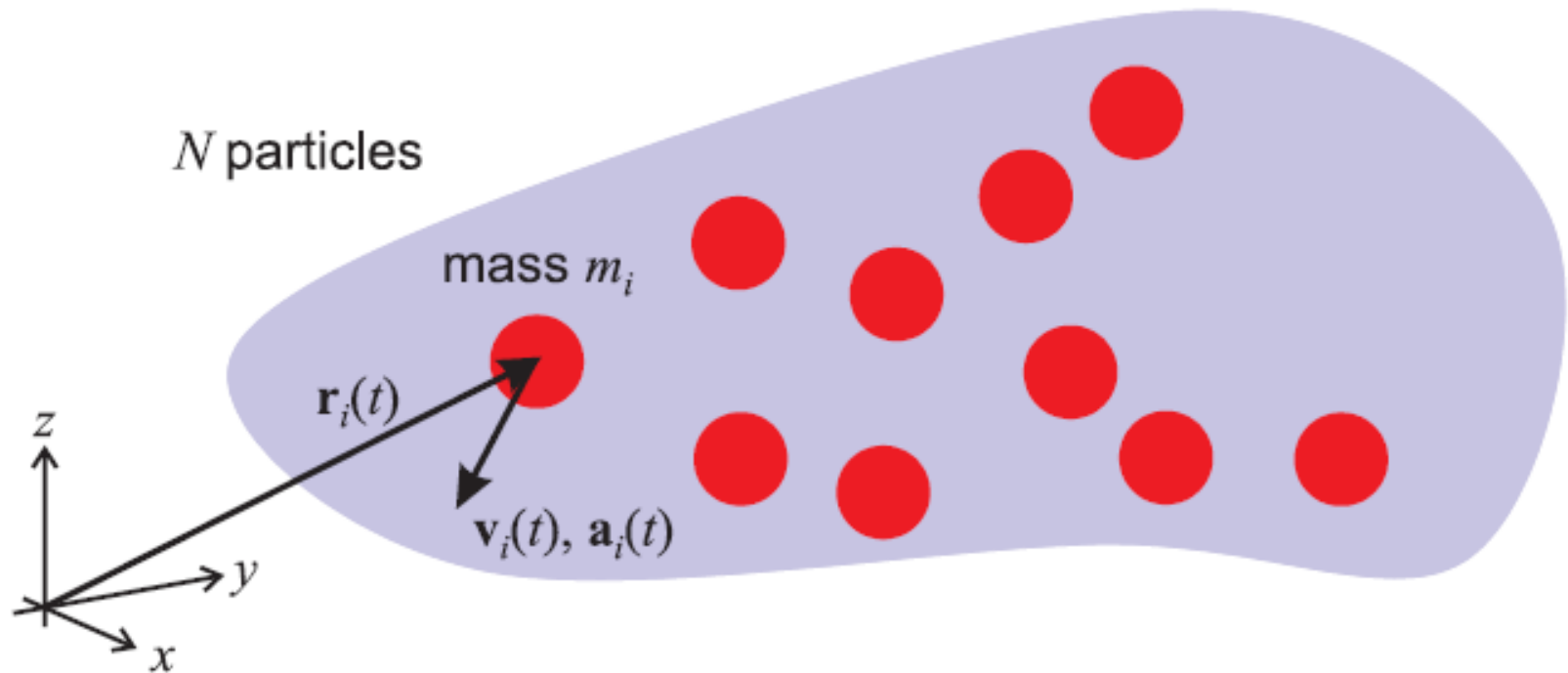
Follow this quantity over time

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

**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:

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Taylor series expansion  $r_i(t)$  around

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

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

# Solving the equations

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

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

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Taylor series expansion  $r_i(t)$  around

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

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

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

# Taylor expansion of $r_i(t)$

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

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

---

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

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

# Taylor expansion of $r_i(t)$

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

---



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

# Taylor expansion of $r_i(t)$

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

---



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



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



# Taylor expansion of $r_i(t)$

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

---



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



$$r_i(t_0 + \Delta t) = \underbrace{2r_i(t_0)}_{\text{Positions at } t_0} - \underbrace{r_i(t_0 - \Delta t)}_{\text{Positions at } t_0 - \Delta t} + \underbrace{a_i(t_0)\Delta t^2}_{\text{Accelerations at } t_0} + \dots$$

Positions  
at  $t_0$

Positions  
at  $t_0 - \Delta t$

Accelerations  
at  $t_0$

# Physics of particle interactions

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

1. 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.

$$f = m \frac{d^2 x}{dt^2} = ma$$

2<sup>nd</sup> law

# Verlet central difference method

$$r_i(t_0 + \Delta t) = \underbrace{2r_i(t_0)}_{\substack{\text{Positions} \\ \text{at } t_0}} - \underbrace{r_i(t_0 - \Delta t)}_{\substack{\text{Positions} \\ \text{at } t_0 - \Delta t}} + \underbrace{a_i(t_0)\Delta t^2}_{\substack{\text{Accelerations} \\ \text{at } t_0}} + \dots$$

How to obtain  
accelerations?

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

Need forces on atoms!

# Verlet central difference method

$$r_i(t_0 + \Delta t) = \underbrace{2r_i(t_0)}_{\substack{\text{Positions} \\ \text{at } t_0}} - \underbrace{r_i(t_0 - \Delta t)}_{\substack{\text{Positions} \\ \text{at } t_0 - \Delta t}} + \underbrace{f_i(t_0) / m\Delta t^2}_{\substack{\text{Forces} \\ \text{at } t_0}} + \dots$$

# Forces on atoms

- Consider energy landscape due to chemical bonds

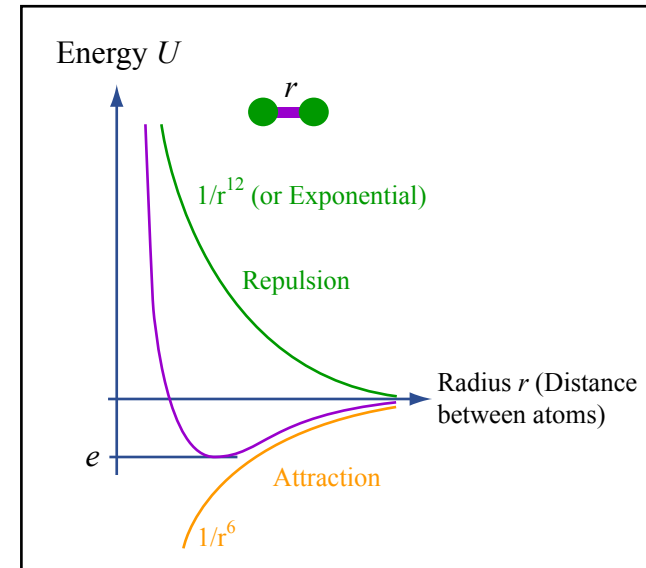
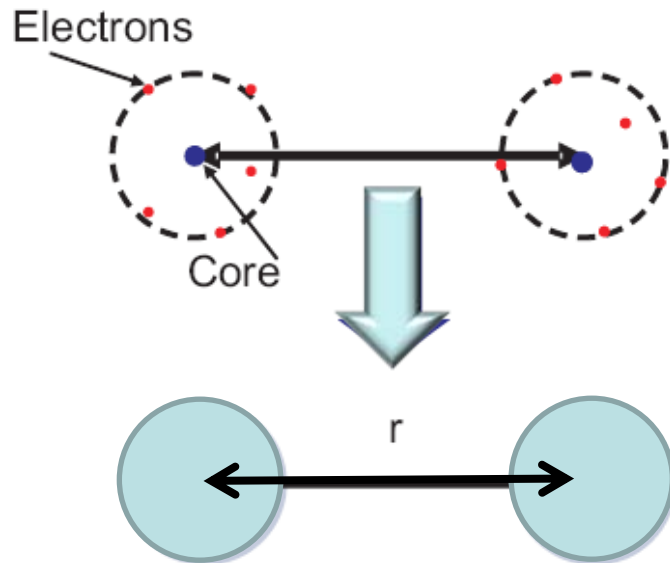


Image by MIT OpenCourseWare.

**Attraction:** Formation of chemical bond by sharing of electrons

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

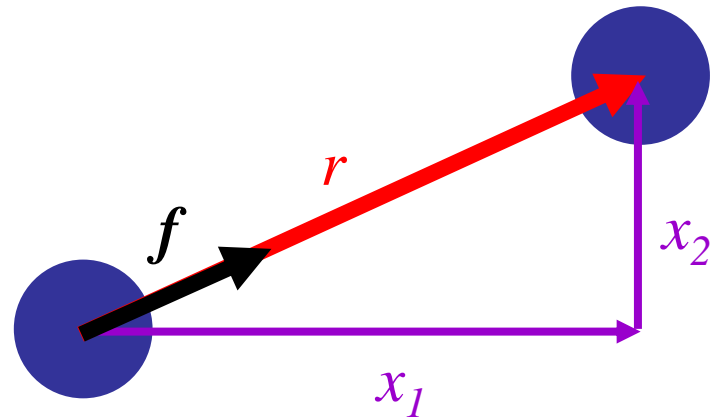
# How are forces calculated?

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

$$f = -\frac{dU(r)}{dr}$$

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

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

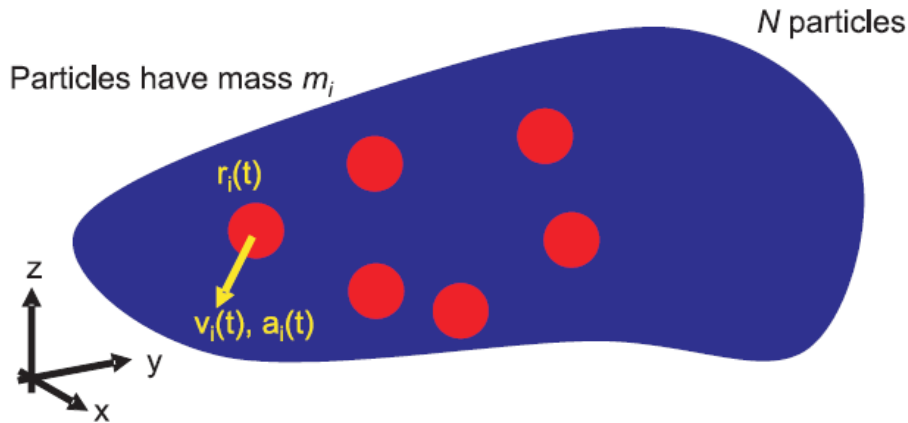


Often: Assume pair-wise interaction between atoms

# 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”*

$$r_i(t_0 + \Delta t) = \underbrace{2r_i(t_0)}_{\text{Positions at } t_0} - \underbrace{r_i(t_0 - \Delta t)}_{\text{Positions at } t_0 - \Delta t} + \underbrace{a_i(t_0)\Delta t^2}_{\text{Accelerations at } t_0} + \dots$$

Positions  
at  $t_0$

Positions  
at  $t_0 - \Delta t$

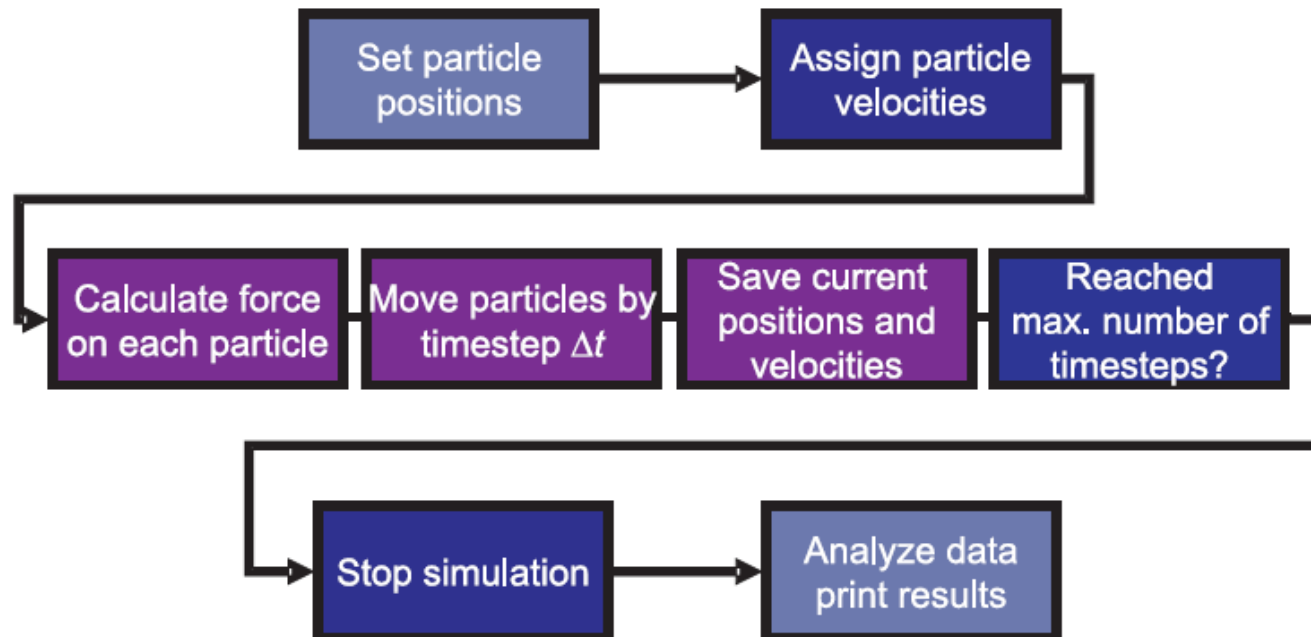
Accelerations  
at  $t_0$

$$a_i = f_i / m$$



# 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):**

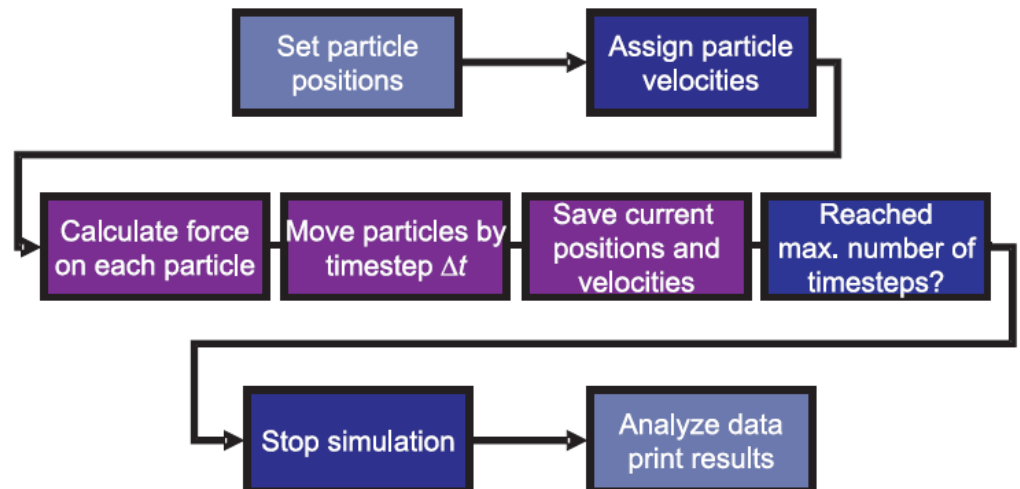
**Calculate force on each particle (subroutine)**

**Move particle by time step  $\Delta t$**

**Save particle position, velocity, acceleration**

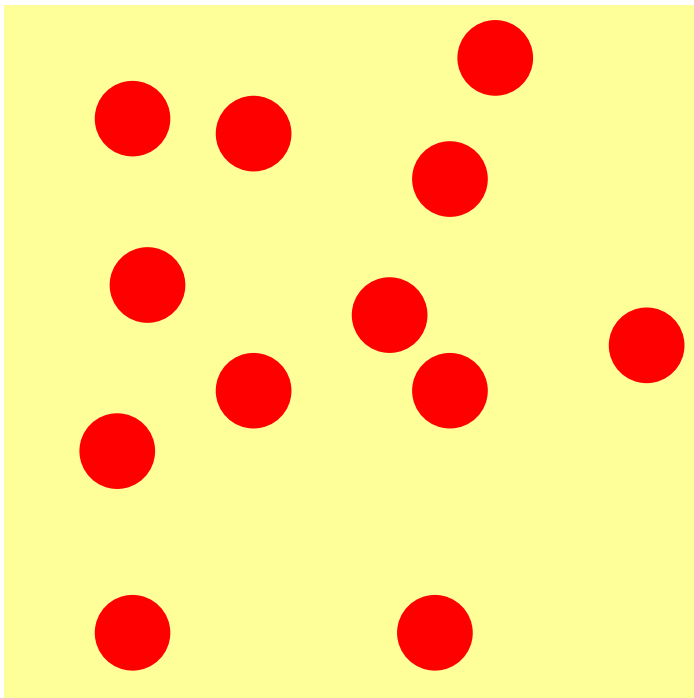
**Save results**

**Stop simulation**

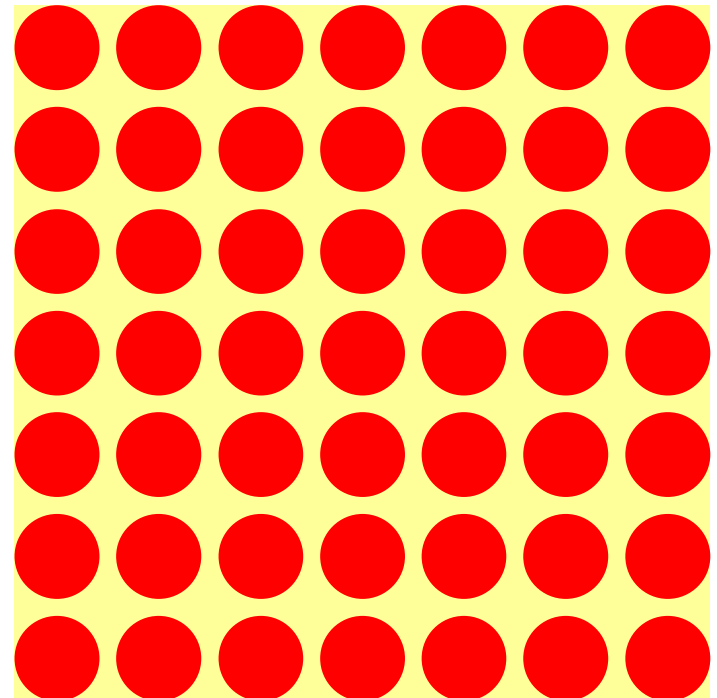


# Atomic positions (initial conditions)

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



“gas (liquid)”



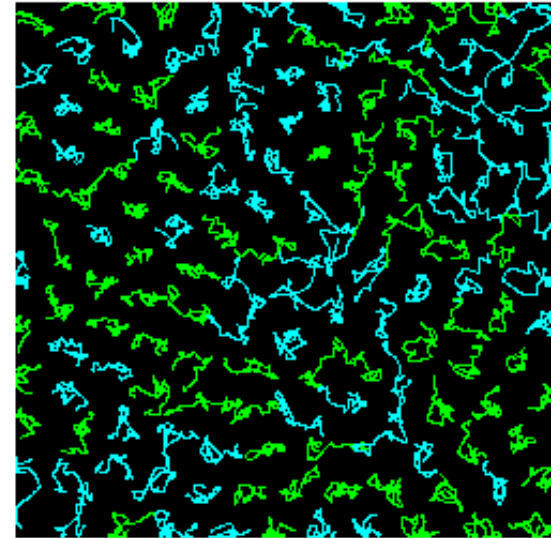
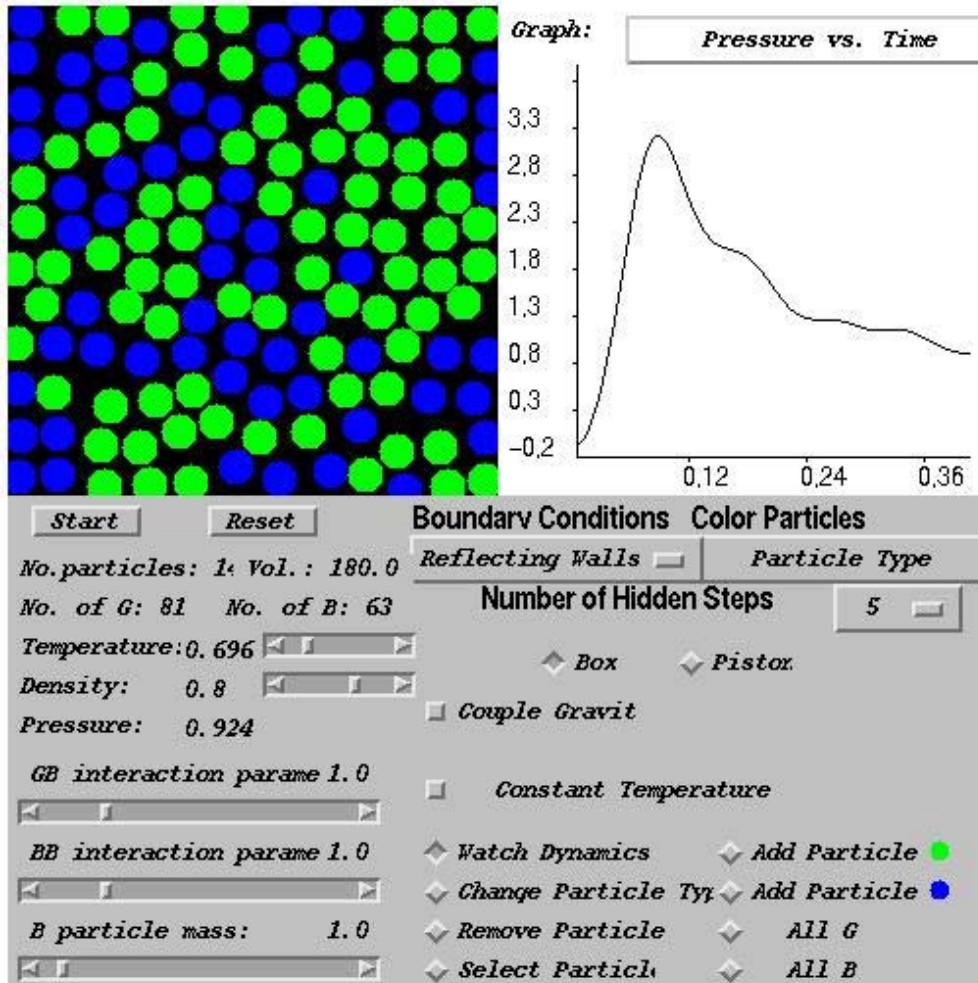
“solid - crystal”

# 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



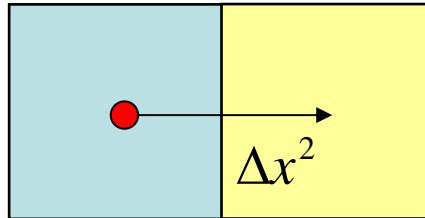
Courtesy of the Center for Polymer Studies at Boston University. Used with permission.

URL <http://polymer.bu.edu/java/java/LJ/index.html>

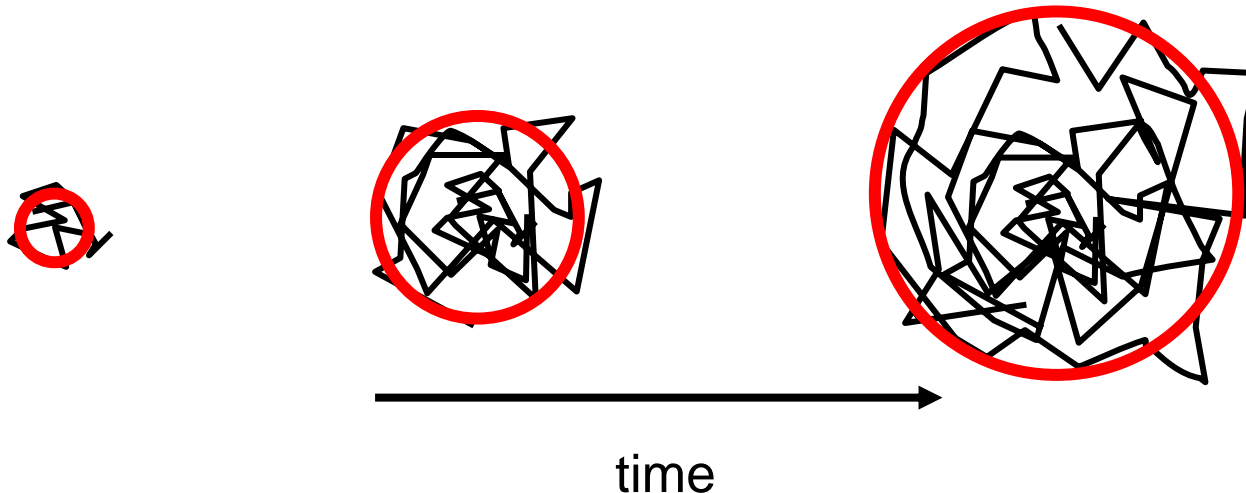
# Link atomistic trajectory with diffusion constant (1D)

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

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



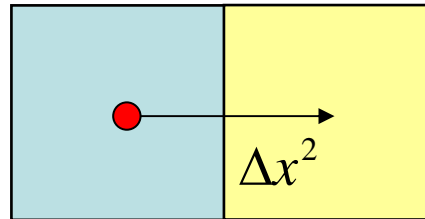
**Idea** – Use MD simulation to measure square of displacement from initial position of particles,  $\Delta r^2(t)$  :



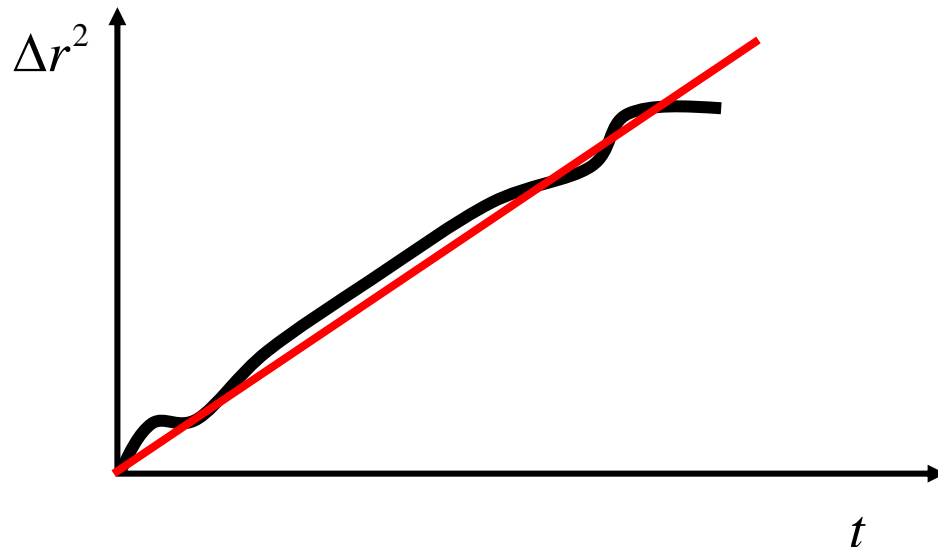
# Link atomistic trajectory with diffusion constant (1D)

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

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



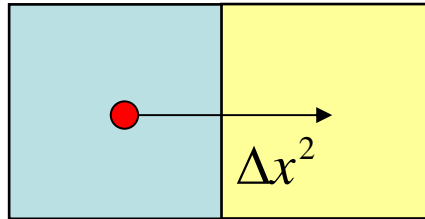
**MD simulation:** Measure square of displacement from initial position of particles,  $\Delta r^2(t)$  :



# Link atomistic trajectory with diffusion constant (1D)

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

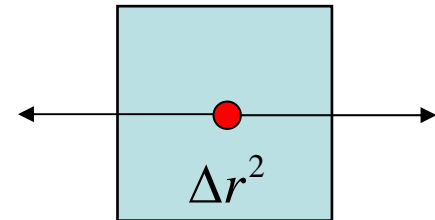
$$D = p \frac{\Delta x^2}{\Delta t}$$



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

Replace

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

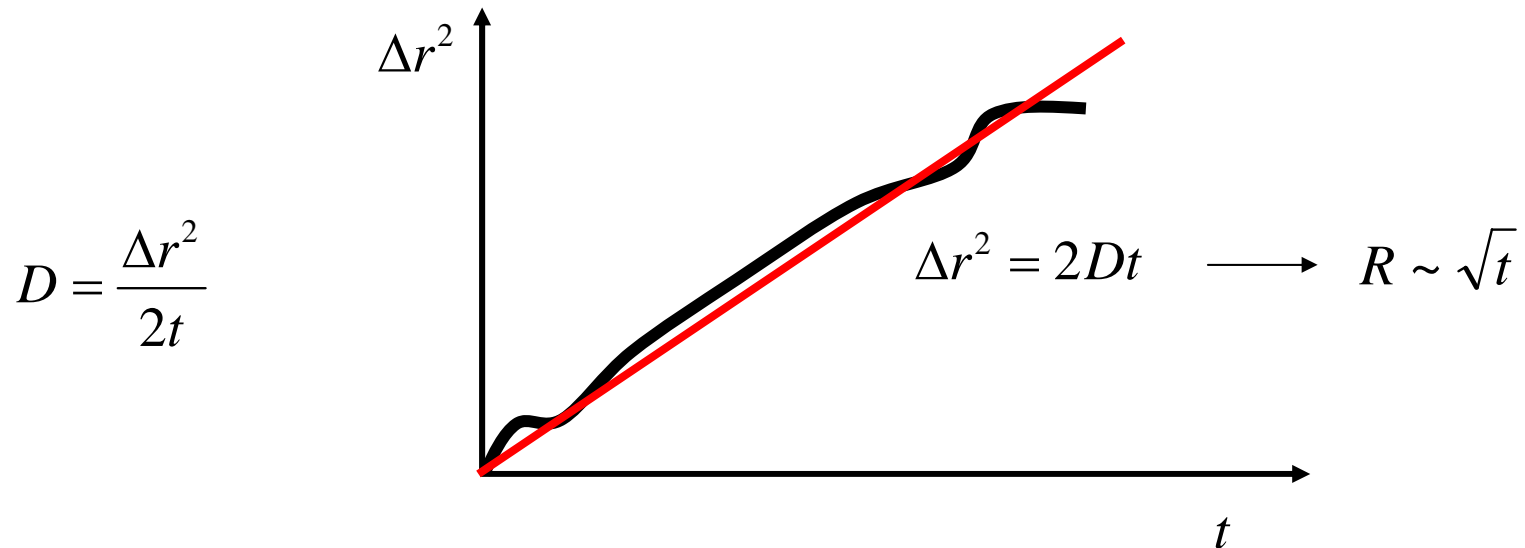


Factor 1/2 = no directionality in (equal probability to move forth or back)



# Link atomistic trajectory with diffusion constant (1D)

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



# Link atomistic trajectory with diffusion constant (2D/3D)

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



*Higher dimensions*

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

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

Factor  $d = 1, 2, \text{ or } 3$  due to 1D, 2D, 3D (dimensionality)

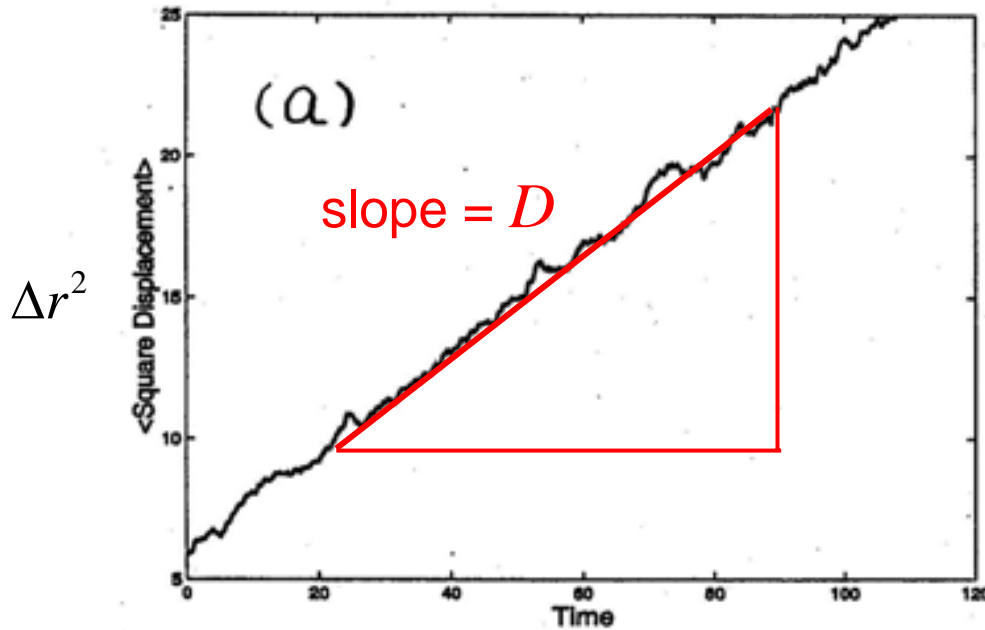
Since:

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

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

$C = \text{constant}$  (does not affect  $D$ )

# Example: MD simulation



source: S. Yip, lecture notes

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

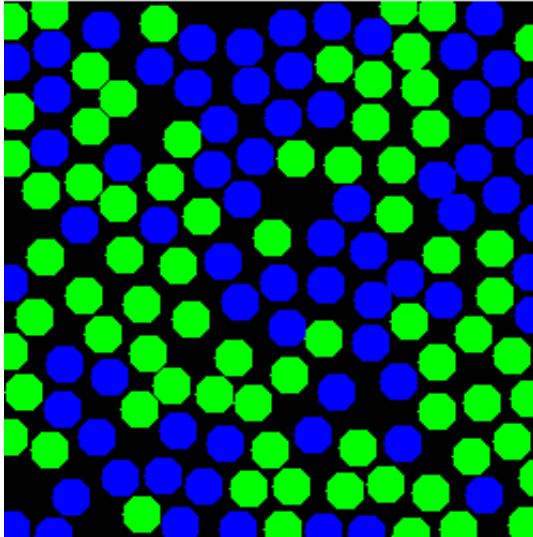
↑  
1D=1, 2D=2, 3D=3

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

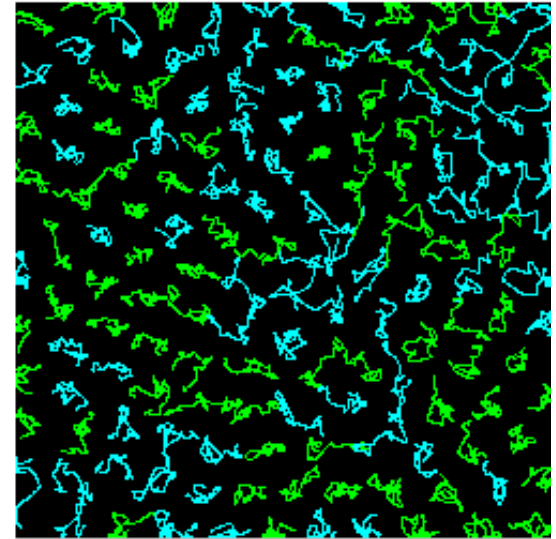
$\langle .. \rangle$  = average over all particles

# Example molecular dynamics

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



Particles

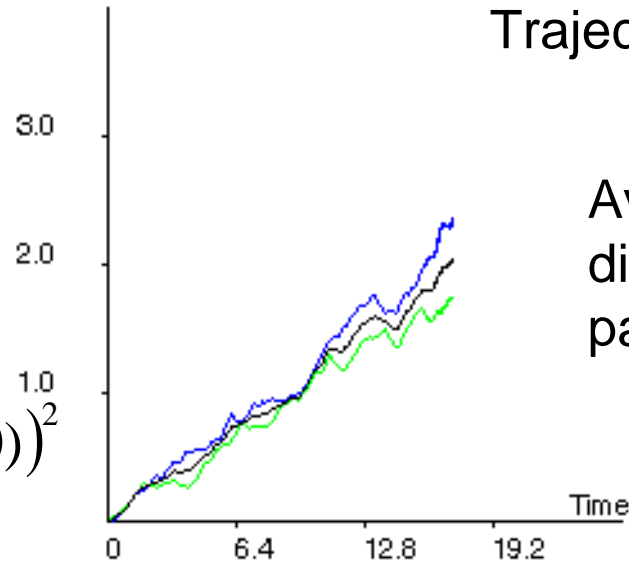


Trajectories

**Mean Square Displacement function**

$$\langle \Delta r^2(t) \rangle = \frac{1}{N} \sum_i (\vec{r}_i(t) - \vec{r}_i(t=0))^2$$

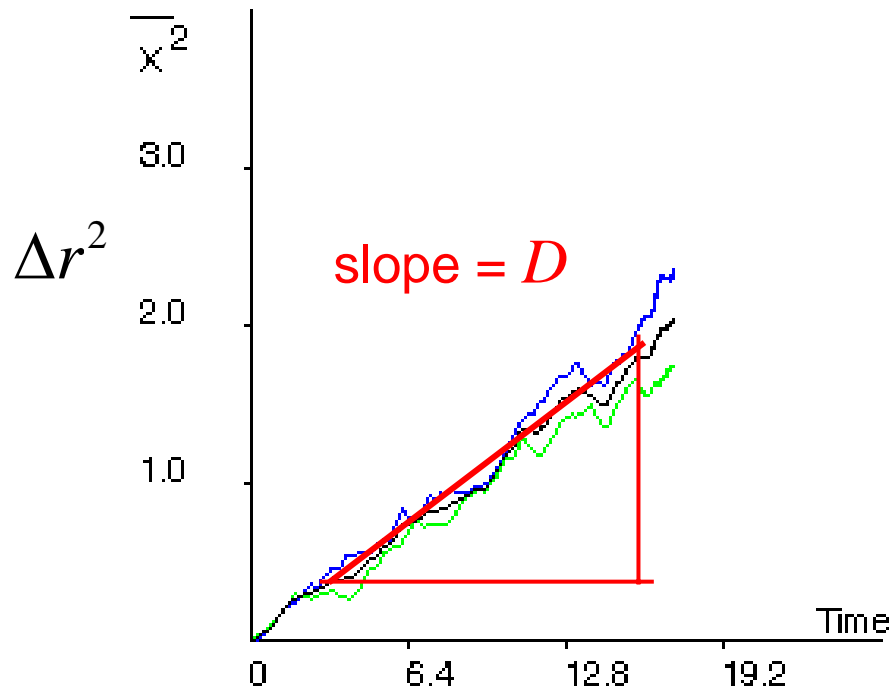
$\Delta r^2$



Average square of displacement of all particles

# Example calculation of diffusion coefficient

$$\langle \Delta r^2(t) \rangle = \frac{1}{N} \sum_i \underbrace{(r_i(t))}_{\text{Position of atom } i \text{ at time } t} - \underbrace{r_i(t=0)}_{\text{Position of atom } i \text{ at time } t=0}$$

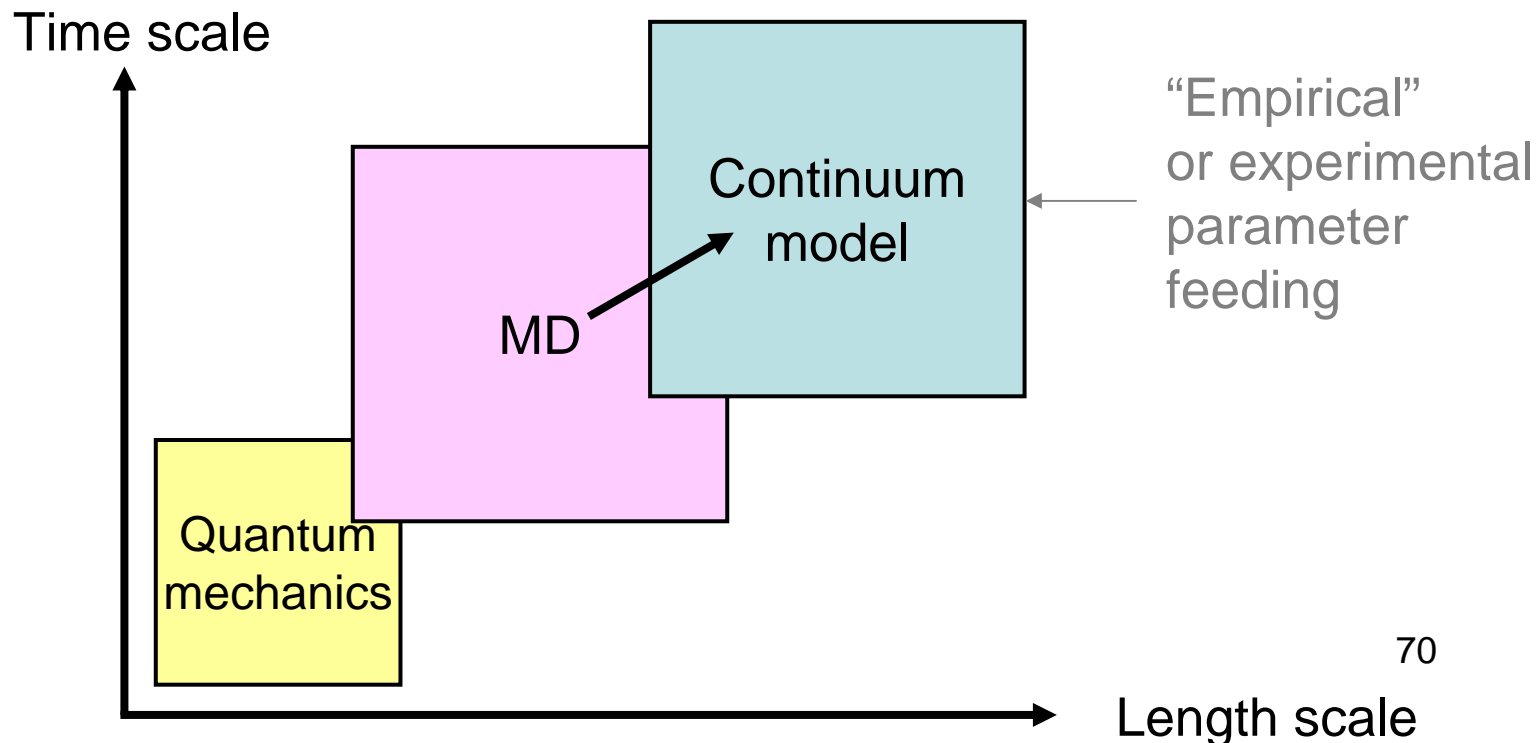


$$D = \frac{1}{2d} \lim_{t \rightarrow \infty} \frac{d}{dt} \langle \Delta r^2 \rangle$$

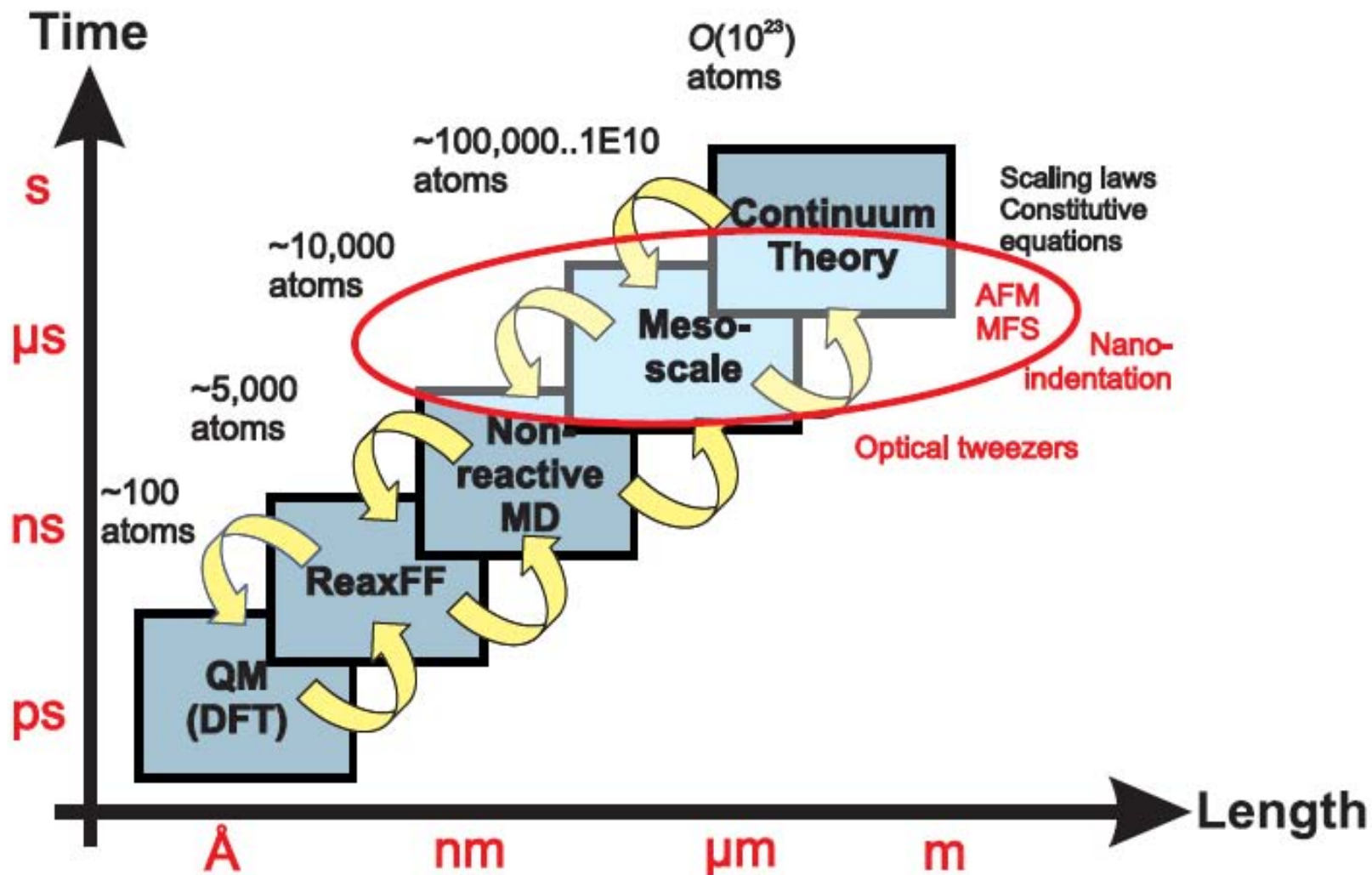
↑  
1D=1, 2D=2, 3D=3

# Summary

- 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



# Multi-scale simulation paradigm



## 3. Additional remarks



# Historical development of computer simulation

- Began as tool to exploit computing machines developed during World War II
- MANIAC (1952) at Los Alamos used for computer simulations
- Metropolis, Rosenbluth, Teller (1953): Metropolis Monte Carlo method
- Alder and Wainwright (Livermore National Lab, 1956/1957): dynamics of hard spheres
- Vineyard (Brookhaven 1959-60): dynamics of radiation damage in copper
- Rahman (Argonne 1964): liquid argon
- Application to more complex fluids (e.g. water) in 1970s
- Car and Parrinello (1985 and following): *ab-initio* MD
- Since 1980s: Many applications, including:
  - Karplus, Goddard et al.: Applications to polymers/biopolymers, proteins since 1980s
  - Applications to fracture since mid 1990s to 2000
  - Other engineering applications (nanotechnology, e.g. CNTs, nanowires etc.) since mid 1990s-2000

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Spring 2011

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