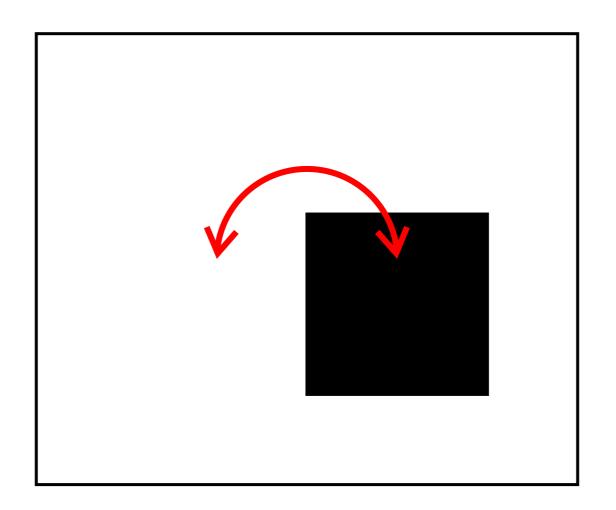
# Advanced Molecular Dynamics

**Ensembles** 

May 11, 2021

# Sub system



Exchange of "things"

#### "Pure" Molecular Dynamics

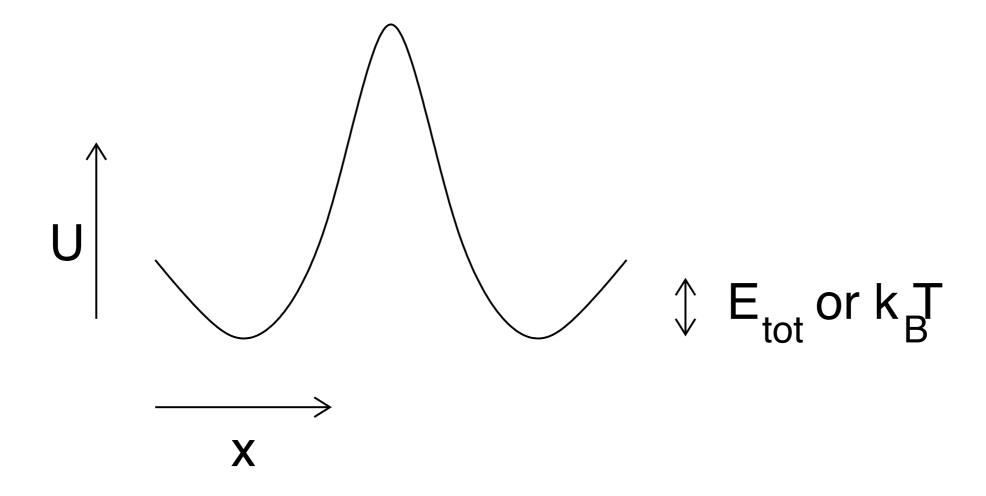
$$m_i \frac{d^2 \mathbf{x}_i}{dt^2} = -\nabla_i V(\mathbf{x}) , \quad i = 1, \dots, N$$

• Plain MD conserves  $E_{tot}=E_{pot}+E_{kin}=U+K$ 

 Plain MD generates a constant NVE, or microcanonical ensemble

This results in reliable dynamics

# Crossing a barrier?



## Thermal equilibrium

- In practice systems are in thermal equilibrium with the environment
- Simplest constant-T ensemble:
  - constant NVT or canonical ensemble

$$\rho(\mathbf{x}, \mathbf{v}) = \frac{1}{Z} \exp \left[ -\frac{U(\mathbf{x}) + K(\mathbf{v})}{k_B T} \right]$$

partition function: 
$$Z = \int \exp \left[ -\frac{U(\mathbf{x}) + K(\mathbf{v})}{k_B T} \right] d\mathbf{x} d\mathbf{v}$$

$$\rho(\mathbf{x}) = \frac{1}{Z} \exp\left[-\frac{U(\mathbf{x})}{k_B T}\right]$$

#### Ensemble averages

What you measure in experiment are ensemble averages

$$\langle g(\mathbf{x}) \rangle = \int g(\mathbf{x}) \rho(\mathbf{x}) d\mathbf{x}$$

For the canonical ensemble:

$$\langle g(\mathbf{x}) \rangle = \int g(\mathbf{x}) \frac{1}{Z} \exp \left[ -\frac{U(\mathbf{x})}{k_B T} \right] d\mathbf{x}$$

#### Different ensembles

- In the limit of large systems all ensembles (or ensemble averages) are identical
  - But how large is large?
    - MD systems are often "small"
      - so you need to think about which ensemble to use

# Ergodicity

time average = average over many instances

#### How to generate a canonical ensemble?

- Add a thermostat
  - for instance a Nose-Hoover thermostat:

$$\frac{d^2\mathbf{r}_i}{dt^2} = \frac{\mathbf{F}_i}{m_i} - \frac{p_\xi}{Q} \frac{d\mathbf{r}_i}{dt}$$
 Q: mass parameter, 
$$\frac{dp_\xi}{dt} = (T - T_0)$$
 determines coupling strength (time)

determines coupling strength (time)

Conserved quantity:

$$H = \sum_{i=1}^{N} \frac{\mathbf{p}_i}{2m_i} + U\left(\mathbf{r}_1, \mathbf{r}_2, \dots, \mathbf{r}_N\right) + \frac{p_{\xi}^2}{2Q} + N_f kT\xi$$

# Simple velocity scaling

- Nose-Hoover is correct, but second order coupling, this can cause oscillations
- Simple first order velocity scaling (Berendsen thermostat):

$$\frac{dT}{dt} = \frac{T_0 - T}{\tau}$$

velocity scaling factor:

$$\lambda = \left[1 + \frac{\Delta t}{\tau_T} \left\{ \frac{T_0}{T(t - \frac{1}{2}\Delta t)} - 1 \right\} \right]^{1/2}$$

issue: no proper ensemble

# Corrected velocity scaling

• Add a stochastic term: Bussi thermostat (JCP 126:014101,2007)

$$dK = (K_0 - K)\frac{dt}{\tau_T} + 2\sqrt{\frac{KK_0}{N_f}}\frac{dW}{\sqrt{\tau_T}}$$

W is a Wiener process:

a random process with
expectation zero, which can be
constructed from a random walk

Advantage: nice first order decay of deviations

# Stochastic dynamics

 Another option: add friction and noise to every degree of freedom:

friction coefficient 
$$m_i \frac{d^2 \mathbf{r}_i}{dt^2} = -m_i \xi_i \frac{d \mathbf{r}_i}{dt} + \mathbf{F}_i(\mathbf{r}) + \mathbf{w}_i \quad \text{Wiener process}$$
$$\langle \mathbf{w}(t) \mathbf{w}(t+s) \rangle = 2m_i \xi_i k_B T \delta(s) \delta_{ij}$$

- Advantage: good ergodicity/equilibration
- Disadvantage: loss of dynamics/inertia

## Dissipative particle dynamics

- Stochastic dynamics on velocity differences between particle pairs
  - advantage: preserves hydrodynamics
  - adds some friction (viscosity)
  - you can solve Navier-Stokes equations using particles
  - but pair interactions are more complex and expensive to calculate

#### Constant pressure

- Constant NPT ensemble
  - one of the most popular ensembles
  - Use a barostat
    - for instance Parrinello-Rahman:

$$\frac{d\mathbf{b}^2}{dt^2} = V\mathbf{W}^{-1}\mathbf{b}'^{-1}\left(\mathbf{P} - \mathbf{P}_{ref}\right)$$
b: box vectors W: mass parameter

 disadvantage second order coupling: volume oscillations

#### Berendsen pressure coupling

- For equilibration you can use first order Berendsen pressure coupling:
  - scale box elements with:

$$\mu_{ij} = \delta_{ij} - \frac{\Delta t}{3\,\tau_p} \beta_{ij} \{P_{0ij} - P_{ij}(t)\}$$
 coupling time

- disadvantage: no proper ensemble and fluctuations
- Conclusion: use Berendsen for equilibration and Parrinello-Rahman for productions

# Stochastic cell rescaling

 To get the correct ensemble, again add fluctuations to Berendsen coupling:

Use the strain:  $\epsilon = \log(V/V_0)$ 

$$\mathrm{d}\epsilon = -\frac{\beta}{\tau_p}(P_0 - P)\mathrm{d}t + \sqrt{\frac{2k_B T\beta}{V\tau_p}}\mathrm{d}W$$

- Velocities are scaled reciprocal to the coordinates
- First order coupling and the correct ensemble
- Bernetti, M. and Bussi, G., "Pressure control using stochastic cell rescaling", J. Chem. Phys., 153, 114107 (2020)

#### Constant chemical potential

- Constant mu-PT ensemble
  - you can use test particle insertion or particle deletion to determine chemical potential and remove/add particles
    - but this disrupts dynamics
    - and can be complex to implement and use

# Reading

• Read Frenkel&Smit II.6