

Lecture 8: Monte Carlo Sampling

@ 2pm, chemistry seminar,

Bin Zhang, MIT

in 2 weeks @ 2pm

Rob DiStasio, Cornell

Today: Nobel Prize in Physics 2021

Giorgio Parisi

Reminder $\langle O \rangle = \int d\vec{x} O(\vec{x}) P(\vec{x})$

$$\vec{X} = \{x_1, x_2, \dots, x_{3N}, p_1, p_2, \dots, p_{3N}\}$$

$$\vec{x} = \{x_1, \dots, x_{3N}\} \quad \vec{p} = \{p_1, \dots, p_{3N}\}$$

$$\vec{X} = \{\vec{x}, \vec{p}\}$$

$$\text{if } P(\vec{x}, \vec{p}) = P(\vec{x}) P(\vec{p})$$

$$\text{if } O(\vec{x}, \vec{p}) = O(\vec{x})$$

@ constant N, V, T

$$P(\vec{x}, \vec{p}) = e^{-\beta H(\vec{x}, \vec{p})} / Z$$

$$Z = \int dx \int dp e^{-\beta H(\vec{x}, \vec{p})}$$

$$H(\vec{x}, \vec{p}) = \sum p_i^2 / 2m_i + U(\vec{x})$$

$$P(\vec{x}, \vec{p}) = \frac{e^{-\beta(\text{KE})} \cdot e^{-\beta U(\vec{x})}}{Z_{\text{KE}} \cdot Z_{\text{PE}}}$$

$$P(\vec{x}) = \int d\vec{p} P(\vec{x}, \vec{p}) = \int d\vec{p} P(\vec{x}) P(\vec{p})$$

$$\int d\vec{p} P(\vec{p}) = 1$$

$$= e^{-\beta U(x)} / Z_{\text{[position]}}$$

Generate a set of configurations

$\{\vec{x}_t\}$ where these are distributed according to $P(\vec{x})$

$\{x_t\}$ want

$$P(x_j) / P(x_i) = \frac{e^{-\beta u(x_j)}}{e^{-\beta u(x_i)}}$$

$$= e^{-\beta \Delta u_{ij}}$$

$$\Delta u_{ij} = u(x_j) - u(x_i)$$

Algorithm: (wrt)

$$x_t \rightarrow x_{t+1} \quad \text{st}$$

$$P(x_{t+1})/P(x_t) = e^{-\beta \Delta U_{t+1,t}}$$

• Generate a Markov Chain

$$\vec{x}_1 \rightarrow \vec{x}_2 \rightarrow \vec{x}_3 \rightarrow \vec{x}_4 \dots \vec{x}_T$$

"Markovian": no memory

$P(\vec{x}_t \rightarrow \vec{x}_{t+1})$ only depends on \vec{x}_t

[not $\vec{x}_1 \dots \vec{x}_t$]

Eg Molecular Dynamics

$$\vec{x}_{t+\Delta t} = \vec{x}_t + \vec{v} \Delta t$$

$$[\vec{x}, \vec{p}] \quad \vec{v} = \vec{p}/m$$

$$\vec{p}_{t+\Delta t} = \vec{p}_t + \vec{a} \Delta t$$

Want:

$$P(x_{t+\Delta t}) / P(x_t) \longrightarrow e^{-\beta \Delta U_{t+\Delta t, t}}$$

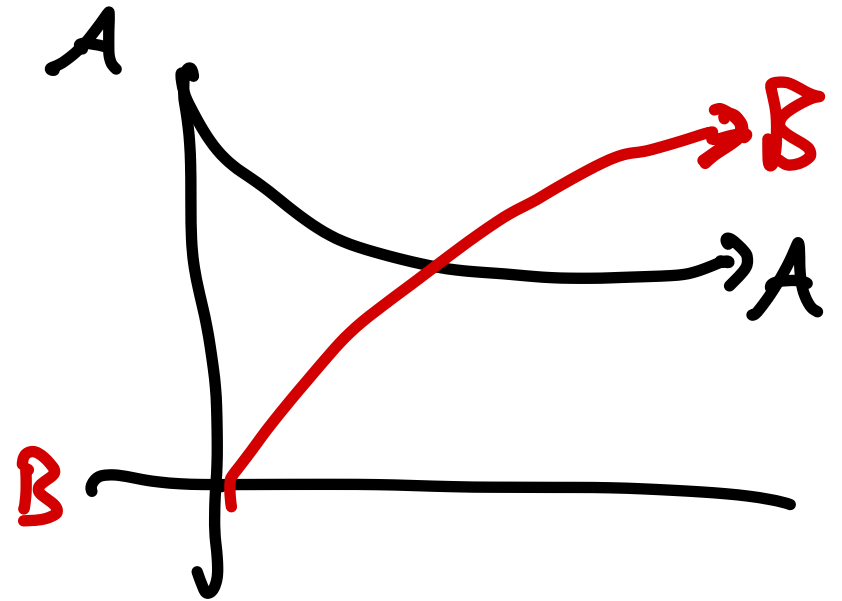
converges? as $t \rightarrow \infty$

(1) Markovian

(2) Detailed Balance *

Detailed Balance

$$A \rightleftharpoons B$$



DB:

$$P(A) \cdot P(A \rightarrow B) = P(B) \cdot P(B \rightarrow A)$$

$$X_t \rightarrow X_{t+1} \rightarrow X_{t+2}$$



$$P(X_t) \underbrace{P(X_t \rightarrow X_{t+1})}_{\text{Choosing a rule}} = P(X_{t+1}) P(X_{t+1} \rightarrow X_t)$$

Choosing a rule

$$P(x \rightarrow y)$$

$$\hookrightarrow P_{\text{gen}}(x \rightarrow y) P_{\text{acc}}(x \rightarrow y)$$

Generation, make new config y

Accept / reject going to y

$$P(x) P_{\text{gen}}(x \rightarrow y) P_{\text{acc}}(x \rightarrow y) =$$

$$P(y) P_{\text{gen}}(y \rightarrow x) P_{\text{acc}}(y \rightarrow x)$$

$$P_{\text{acc}}(x \rightarrow y) = \left[\frac{P(y)}{P(x)} \cdot \frac{P_{\text{gen}}(y \rightarrow x)}{P_{\text{gen}}(x \rightarrow y)} \right] P_{\text{acc}}(y \rightarrow x)$$

"rate"

$r(x \rightarrow y)$



Metropolis Monte Carlo (1953)

MTTRR → Manhattan Project
Los Alamos Computing Center

$$P_{acc}(x \rightarrow y) = \min(1, r(x \rightarrow y))$$

$$P_{acc}(x \rightarrow y) = \left[\frac{P(y)}{P(x)} \cdot \frac{P_{gen}(y \rightarrow x)}{P_{gen}(x \rightarrow y)} \right] P_{acc}(y \rightarrow x)$$

$$P_{acc}(x \rightarrow y) = r(x \rightarrow y) P_{acc}(y \rightarrow x)$$

$$P_{acc}(y \rightarrow x) = r(y \rightarrow x) P_{acc}(x \rightarrow y)$$

$$r(x \rightarrow y) = \frac{1}{r(y \rightarrow x)}$$

$$a) \Gamma(x \rightarrow y) > 1 \quad \text{st} \quad \Gamma(y \rightarrow x) < 1$$

$$b) \Gamma(x \rightarrow y) < 1 \quad \text{st} \quad \Gamma(y \rightarrow x) > 1$$

$$P_{acc} = \min[1, \Gamma(x \rightarrow y)] \quad \leftarrow$$

$$\frac{P_{acc}(x \rightarrow y)}{P_{acc}(y \rightarrow x)} = \Gamma(x \rightarrow y)$$

$$P_{acc}(y \rightarrow x)$$

$$a) \quad P_{acc}(x \rightarrow y) = 1$$

$$P_{acc}(y \rightarrow x) = \frac{1}{\Gamma(x \rightarrow y)}$$

$$b) \quad P_{acc}(x \rightarrow y) = \Gamma(x \rightarrow y) < 1$$

$$P_{acc}(y \rightarrow x) = 1$$

Algorithm: Start at config \vec{x}_t

① Propose x_{t+1} w/ prob $P_{\text{gen}}(x_t \rightarrow x_{t+1})$
generate ~~rand~~ ^{uniform} number "a" $[0, 1)$

② if $a < \min[1, r(x_t \rightarrow x_{t+1})] \leftarrow$
accept, move to x_{t+1}

else:

$$x_{t+1} = x_t$$

③ go back to 1

$$r(x \rightarrow y) = \frac{P_{\text{gen}}(y \rightarrow x)}{P_{\text{gen}}(x \rightarrow y)} \frac{P(y)}{P(x)}$$

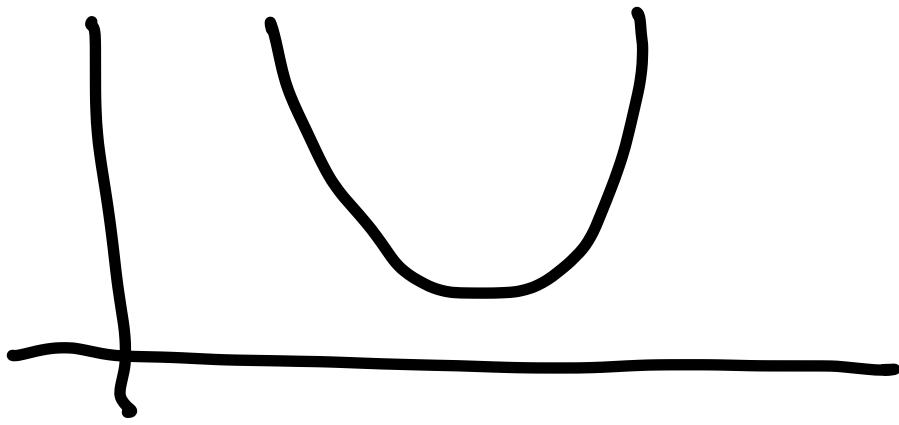
gen part
prob part

For canonical ensemble

$$P(y)/P(x) = e^{-\beta[U(y) - U(x)]}$$

$$e^{-\beta[H(\vec{x}_2, \vec{p}_2) - H(\vec{x}_1, \vec{p}_1)]}$$

Make generation symmetric



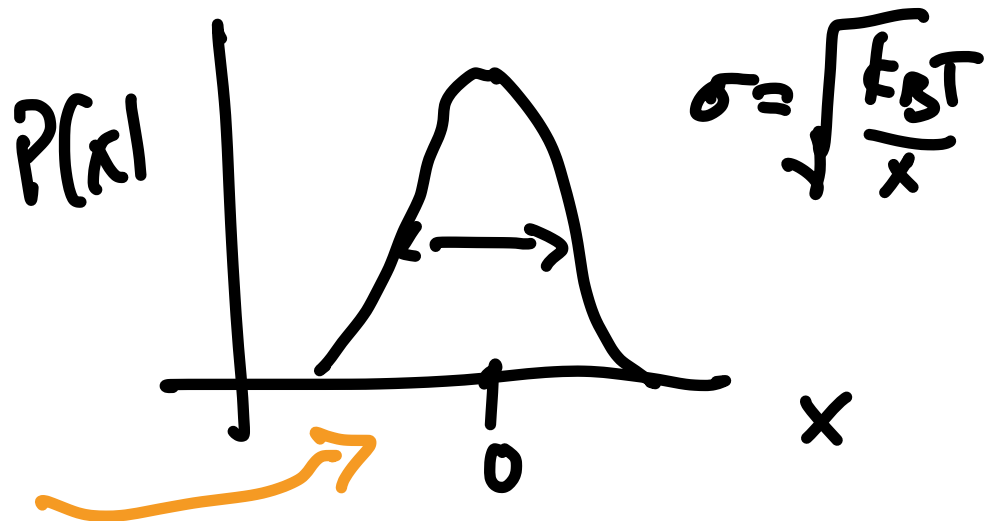
$$\mathcal{H}(x, p) = \frac{p^2}{2m} + \frac{1}{2} kx^2$$

sample

$$P(x) = \frac{e^{-\beta U(x)}}{\int e^{-\beta U(x)} dx} =$$

$$-\beta \frac{1}{2} kx^2$$

$$\sqrt{\frac{2\pi k_B T}{k}}$$



Move rule, that is symmetric
propose:

$$X_{t+1} = X_t + \alpha_2 \cdot \xi_t \quad \checkmark$$

$\alpha_2 \in (-1, 1)$ uniform random

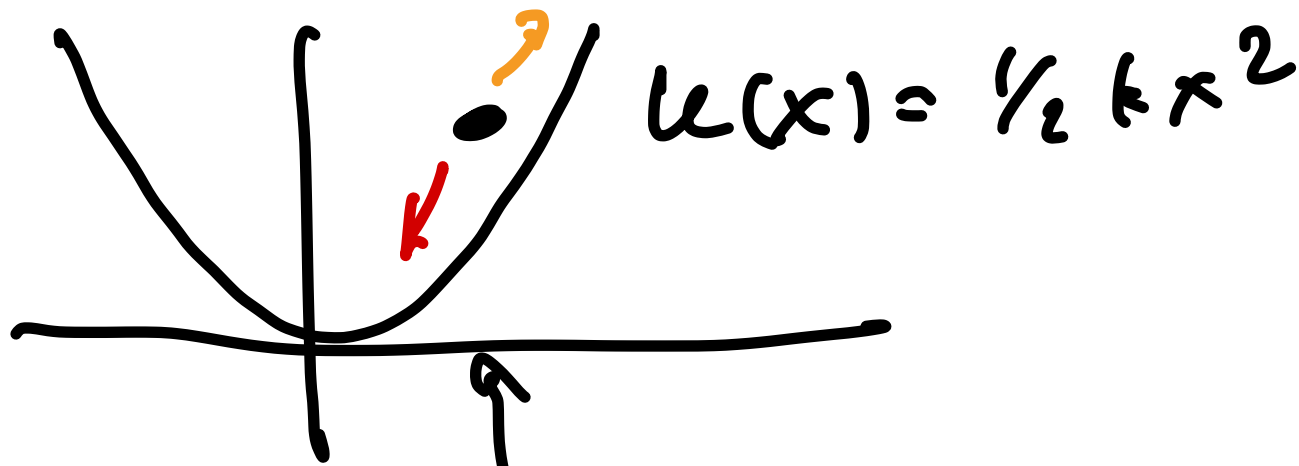
ξ_t biggest possible move

$$X_{t+2} = X_{t+1} + \alpha_2 \cdot \xi_{t+1}$$

$$\frac{P_{\text{gen}}(y \rightarrow x)}{P_{\text{gen}}(x \rightarrow y)} = 1 \quad r(x \rightarrow y) = e^{-\beta \Delta u}$$

$$P_{\text{acc}}(X_t \rightarrow X_{t+1}) = \min(1, e^{-\beta \Delta u})$$

$$P_{acc}(x_t \rightarrow x_{t+1}) = \min(1, e^{-\beta \Delta u})$$



how does Δu
compare to $k_B T$

current pos

move 1, energy goes down

$U(x_2) - U(x_1) < 0, e^{-\beta \Delta u} > 1$, always accept

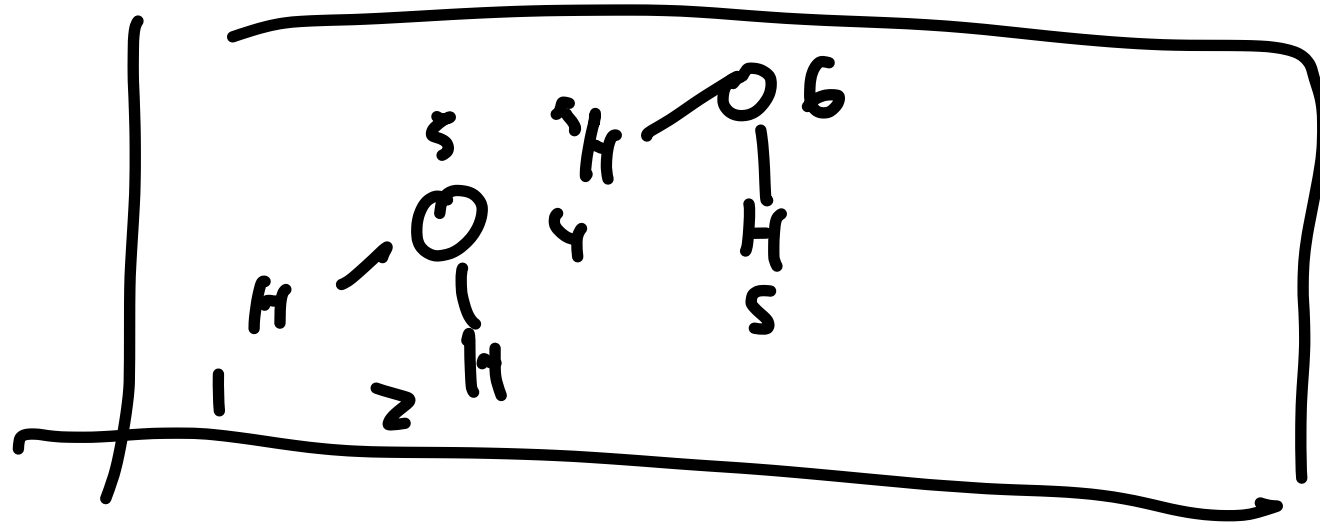
move 2 $U(x_2) - U(x_1) > 0, e^{-\beta \Delta u} < 1$, accept prob $e^{-\beta \Delta u}$

Tune our moves, here ϵ
st average acceptance rate \sim
 $0.25 \rightarrow 0.5$

Trade off between efficiency &
exploration

if ϵ is large, each accepted
move will go far, but most
moves will be rejected

ϵ is really small, almost always accept
but stay close to starting point



Doesn't have to be this simple

① pick an atom [atom 4]

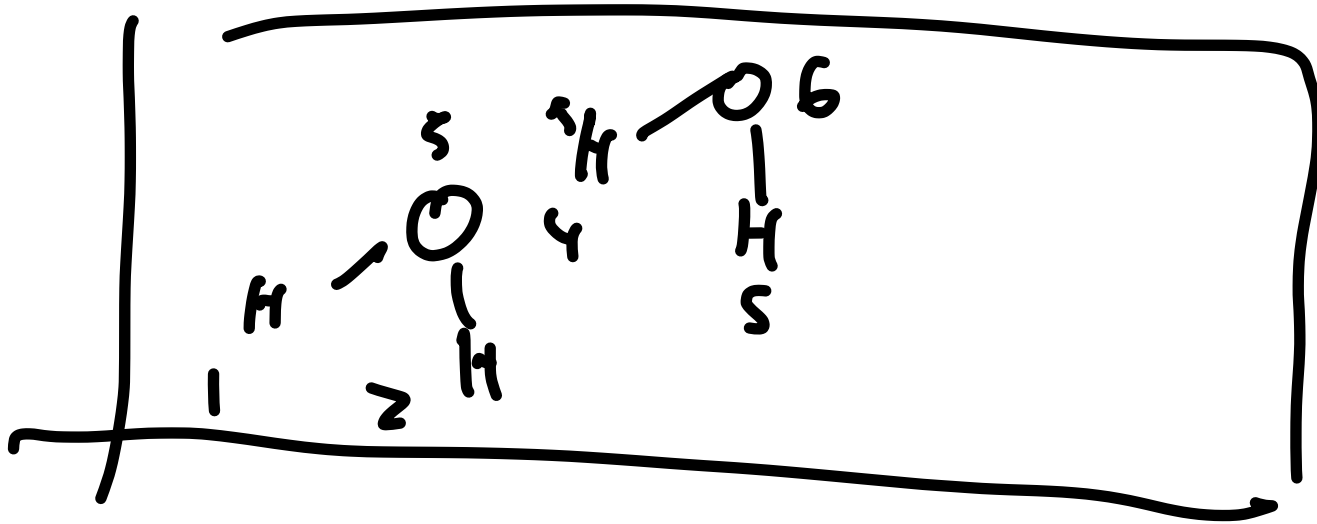
② generate a random move

$$\vec{x}_4(t+1) = \vec{x}_4(t) + \text{random}$$

$$\vec{x}_4 = \begin{bmatrix} x_4 \\ y_4 \\ z_4 \end{bmatrix} + \begin{bmatrix} \text{random} \cdot \xi \\ \text{random} \cdot \xi \\ \text{random} \cdot \xi \end{bmatrix}$$

$\xi \in (-1, 1)$

③ then calc $e^{-\beta \Delta U}$ for whole system



Random move

(1) move COM of molecule

$$X_{\text{com}} = X_{\text{com}} + \vec{a} \xi$$

(2) rotate molecule by a random angle

Why MC & why not:

① easy

② can choose very smart
types of moves, jump over
energy barriers st. explanation
is very fast

Why not:

① not real dynamics [give static properties]

② usually only tiny changes accepted

Do we have to use Metropolis rule

Glauber Rule

$$P_{acc}(x \rightarrow y) = \frac{e^{-\beta \Delta U / 2}}{e^{-\beta \Delta U / 2} + e^{+\beta \Delta U / 2}} \quad \leftarrow$$

why? $\Delta U_{xy} = U(y) - U(x)$

$$\Delta U_{yx} = U(x) - U(y) = -\Delta U_{xy}$$

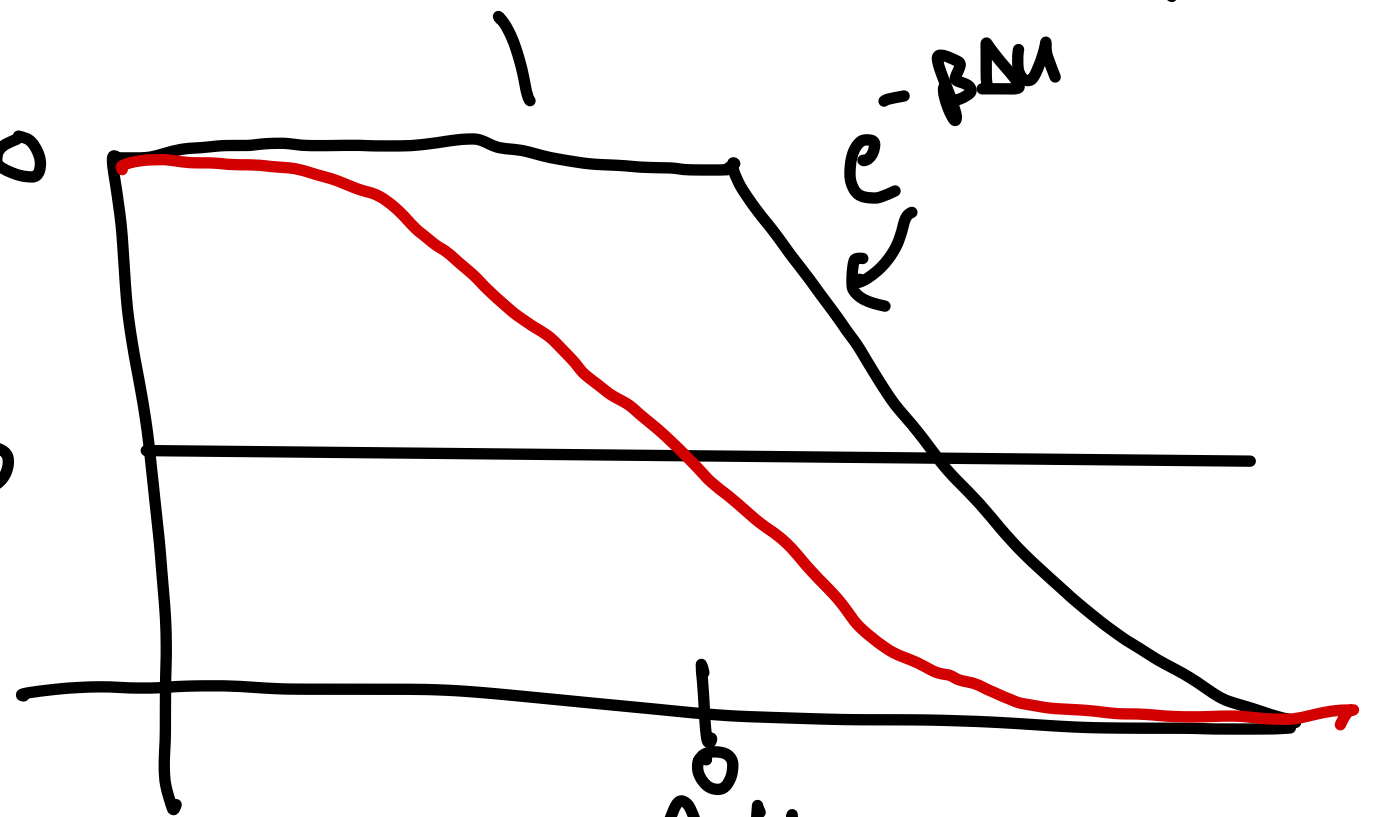
$$\frac{P_{acc}(x \rightarrow y)}{P_{acc}(y \rightarrow x)} = e^{-\beta \Delta U_{xy}} \quad \hat{=} \quad \checkmark$$

accept
rate

metropolis

100%

50%



$e^{-\beta\Delta U}$

ΔU

$-\infty$

$+\infty$