

Recap:

We have a very high-dimensional space

$$\text{ex: } \vec{S} = \{S_1, S_2, \dots, S_N\}$$

↑
State of spin 1 (± 1)

So the number of microstates is HUGE — we could never practically enumerate them all.

We are interested in the mean value of an order parameter, which is a function of \vec{S} , in this case $|M|(\vec{S})$

For any sample of possible spin configurations, $\vec{S}^{(1)} = \{S_1^{(1)}, S_2^{(1)}, S_3^{(1)}, \dots, S_N^{(1)}\}$, it is trivial to compute $M^{(1)} = \sum_i S_i^{(1)}$, but it is not trivial to compute $\langle M \rangle$, the average over all possible spin configurations.

If we could generate p independent samples $\vec{S}^{(1)}, \vec{S}^{(2)}, \dots, \vec{S}^{(p)}$, we could approximate the expectation value by

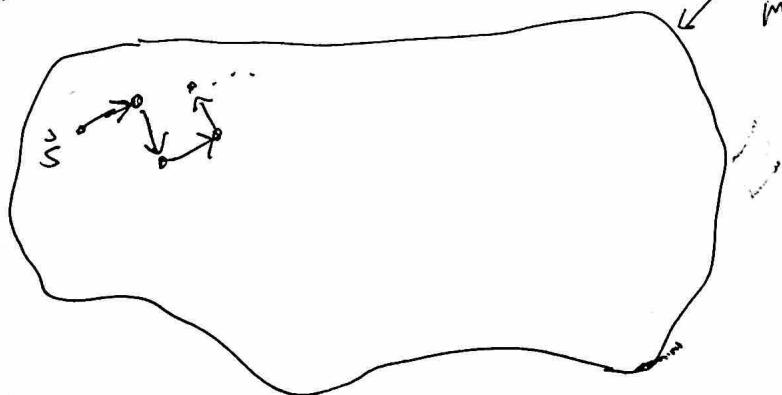
$$\langle M \rangle \approx [M(\vec{S}^{(1)}) + M(\vec{S}^{(2)}) + \dots + M(\vec{S}^{(p)})] \frac{1}{p}$$

provided the samples come from the canonical distribution

$$P(\vec{S}) \propto e^{-\beta E(\vec{S})} \xrightarrow{\text{After normalization}} P(\vec{S}) = \frac{e^{-\beta E(\vec{S})}}{Q}$$

How do we sample a configuration \vec{S} from this distribution when we have no hope of computing Q ?

Answer: We cook up a dynamics



Set of all microstates

Goal of the dynamics:
No matter where we start,
if you run the movie
long enough and pick out
a frame, it will have
probability $P(\vec{S}) = \frac{e^{-\beta E(\vec{S})}}{Q}$.

Imagine for a moment that we could list all the possible microstates
And we make a vector of the probability of being in each microstate at time t .

$$\vec{p}(t) = \begin{pmatrix} p(1,t) \\ p(2,t) \\ \vdots \\ p(n,t) \end{pmatrix}$$

After one step of our dynamics we are at time $t + \Delta t$ with

$$\vec{p}(t + \Delta t) = \begin{pmatrix} p(1, t + \Delta t) \\ p(2, t + \Delta t) \\ \vdots \end{pmatrix} = \begin{pmatrix} p(1, t) p(1 \rightarrow 1) + p(2, t) p(2 \rightarrow 1) + \dots \\ p(1, t) p(1 \rightarrow 2) + p(2, t) p(2 \rightarrow 2) + \dots \\ \vdots \end{pmatrix}$$

$$= \begin{pmatrix} p(1 \rightarrow 1) & p(2 \rightarrow 1) & \dots \\ p(1 \rightarrow 2) & p(2 \rightarrow 2) & \vdots \end{pmatrix} \vec{p}(t)$$

Call this matrix of all the transition probabilities the "transition matrix" T .

$$\Rightarrow \vec{p}(t + \Delta t) = T \vec{p}(t)$$

T defines the "dynamics" - it tells us how to transform from one microstate to another.

What is $\vec{p}(0)$? (For a simulation)

We'll start in one particular configuration, so

$$\vec{p}(0) = \begin{pmatrix} 1 \\ 0 \\ 0 \\ \vdots \\ 0 \\ 0 \end{pmatrix}$$

The index of this spot is the state we start in with probability 1

Pictorially



After n steps,

$$\vec{p}(n\Delta t) = T^n \vec{p}(0).$$

We want

$$\lim_{n \rightarrow \infty} \vec{p}(n\Delta t) = \lim_{n \rightarrow \infty} T^n \vec{p}(0) = \frac{1}{Q} \begin{pmatrix} e^{-\beta E(\vec{s}_1)} \\ e^{-\beta E(\vec{s}_2)} \\ \vdots \end{pmatrix} = \vec{\pi}$$

A stationary distribution
(not a function of time).

Once you reach this
distribution the prob. dist.
stops changing.

That means our dynamics must satisfy

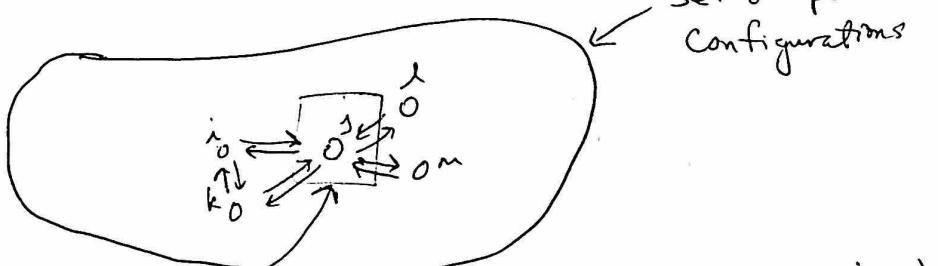
$$T\vec{\pi} = \vec{\pi}$$

Result of
another step of
dynamics

Exactly the
same dist.
we started with

Balance Condition

Why?



The total probability leaving site j : $\pi(j) [p(j \rightarrow l) + p(j \rightarrow m) + \dots]$
must balance with the total probability ~~entering~~ entering site j : $\pi(i)p(i \rightarrow j) + \pi(k)p(k \rightarrow j)$
 $+ \dots$

If the out flows did not balance the in flows, then the prob. of being in site j
would change from one step to the next.

One way (not the only way) to ensure balance is to require every
pair of states to have balanced probabilities

$$i \xleftrightarrow{j} j \quad \pi(i)p(i \rightarrow j) = \pi(j)p(j \rightarrow i)$$

Forcing all pairs to be balanced is called detailed balance — it is a stronger condition
that ensures $T\vec{\pi} = \vec{\pi}$.

For every pair of configurations $i \neq j$, we need a rule for picking transition probabilities $p(i \rightarrow j) + p(j \rightarrow i)$ such that

$$\pi(i) p(i \rightarrow j) = \pi(j) p(j \rightarrow i)$$

$$\Rightarrow \frac{p(j \rightarrow i)}{p(i \rightarrow j)} = \frac{\pi(i)}{\pi(j)} = \frac{e^{-\beta E(i)}/Q}{e^{-\beta E(j)}/Q} = e^{-\beta(E(i) - E(j))}$$

↑
The partition function has dropped out!

One choice you could make is

$$p(j \rightarrow i) = e^{-\beta E(i)} \quad \text{and} \quad p(i \rightarrow j) = e^{-\beta E(j)}. \quad (*)$$

~~This~~ This choice would satisfy $\frac{p(j \rightarrow i)}{p(i \rightarrow j)} = e^{-\beta(E(i) - E(j))}$, but there is a problem. We require that

$$\sum_j p(i \rightarrow j) = 1. \quad (\text{Normalization})$$

Why? Given that you start in i , you must go to some new conf's with probability 1. The new config. could be any j (including $i=j$), but normalization must be imposed for the random walker.

Our choice above (*) could be adapted to make that normalization work out:

$$p(j \rightarrow i) = e^{-\beta E(i)} \quad \text{and} \quad p(i \rightarrow j) = e^{-\beta E(j)} \quad \text{for } i \neq j$$

$$p(i \rightarrow i) = 1 - \sum_{i \neq j} C e^{-\beta E(i)} \quad \text{with } C \text{ chosen such that } 0 \leq p(i \rightarrow i) \leq 1.$$

But computing $C + p(i \rightarrow i)$ requires me to sum over $i \neq j$, a huge # of microstates.
Not looking great!

But there is another way

Conceptually we could split $p(i \rightarrow j)$ into two steps.

1. Propose a new j with some probability that you choose

$p_{\text{gen}}(i \rightarrow j)$: generation probability

2. Decide whether to accept or reject that proposal with probability $p_{\text{acc}}(i \rightarrow j)$

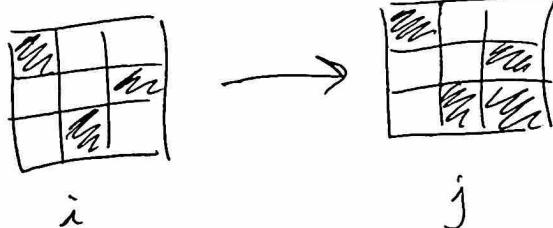
If you reject, stay in i .

The total probability of going from i to j is

$$p(i \rightarrow j) = p_{\text{gen}}(i \rightarrow j) p_{\text{acc}}(i \rightarrow j)$$

$p_{\text{gen}}(i \rightarrow j)$ can be (almost) anything you want as long as you compute an appropriate ~~other~~ p_{acc} . We'll see how to do that.

Ex:



1. Pick a spin at random. We pick the spin w/ probability $\frac{1}{N}$.
2. Flip that spin to produce config. j .

Hence this "MC move" has $p_{\text{gen}}(i \rightarrow j) = \frac{1}{N}$.

Can $p_{\text{acc}}(i \rightarrow j)$ be anything I want like $p_{\text{gen}}(i \rightarrow j)$ could? NO!

We choose $p_{\text{acc}}(i \rightarrow j) + p_{\text{acc}}(j \rightarrow i)$ in such a way to ensure the detailed balance condition holds:

$$\pi(i) p_{\text{gen}}(i \rightarrow j) p_{\text{acc}}(i \rightarrow j) = \pi(j) p_{\text{gen}}(j \rightarrow i) p_{\text{acc}}(j \rightarrow i)$$

There are many choices of pace that work. The most popular are the Metropolis choice + the Glauber choice

Metropolis:

$$p_{\text{acc}}(i \rightarrow j) = \min \left[1, \frac{\pi(j) p_{\text{gen}}(j \rightarrow i)}{\pi(i) p_{\text{gen}}(i \rightarrow j)} \right]$$

Next class we'll confirm that this works.