

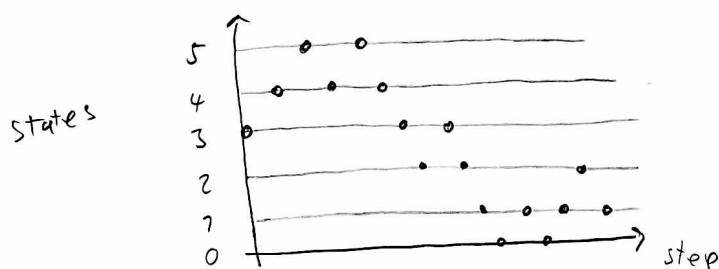
Markov Chain

(Andrey Markov)

- A random sequence in which the prob. for the next state only depends on the current state, not the history of states before that. (Markov property, memoryless)

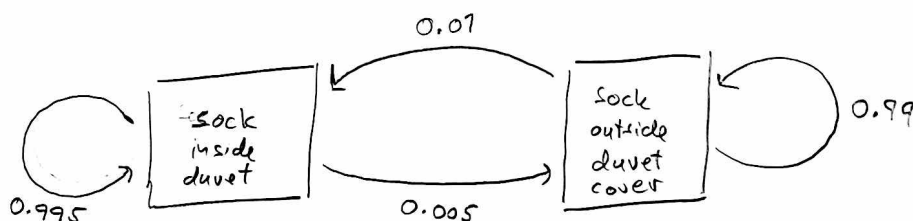
$$[p(x_{i+1} | x_i, x_{i-1}, \dots) = p(x_{i+1} | x_i)]$$

Examples: "Drunkards walk" : move +1 or -1 with equal prob.



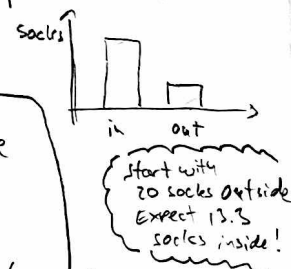
Probs. for next number dep. only on current number, not how the chain got there.

- "Socks in duvet cover"

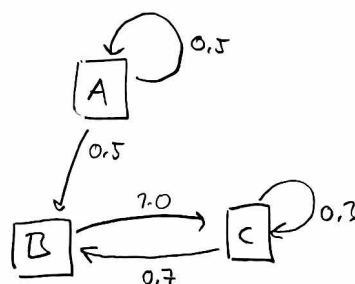


- Distribute walkers across states and iterate → evolve towards stationary / equilib. populations across states
- Important concept: Ergodicity

A chain is ergodic if any state can be reached from any other state with a finite number of steps



Non-ergodic chain:



Another example of Markov Chain: FH1's "agent-based models" for disease spread. Simulate a population of individuals.

Markov Chain Monte Carlo

A method that produces a Markov chain of samples / events / steps whose stationary distribution (long-run distribution) is a given prob distribution

In other words: Generate a set of samples $x_i \sim p(x)$ by recording the steps of a Markov chain in x -space.

General procedure:

- 1) Current state x_i
- 2) Generate a new candidate x' using a proposal pdf that only depends on current state
- 3) Apply acceptance rule. If accepted: $x_{i+1} = x'$
If rejected: $x_{i+1} = x_i$ ← Note: This is also a new sample!
- 4) Repeat

For the list of samples $\{x_1, x_2, x_3, \dots\}$ to end up as a set of samples distributed according to $p(x)$, the acceptance rule should ensure

• Ergodicity (can visit all states)

• Detailed balance: $p(x_i) \text{Prob}(x_i \rightarrow x') = p(x') \text{Prob}(x' \rightarrow x_i)$
can have: \uparrow small \uparrow large $= \uparrow$ large \uparrow small

This is where your chosen pdf appears!

- The Metropolis (or Metropolis-Hastings) algorithm is one example that ensures this. $\left[\text{Basis: } \text{Prob}(x \rightarrow x') = T(x \rightarrow x') A(x \rightarrow x') \right]$

- Generate candidate according to some proposal prob. $T(x_i \rightarrow x')$

- Calculate acceptance prob.

$$A(x_i \rightarrow x') = \min\left(1, \frac{p(x')}{p(x_i)} \frac{T(x' \rightarrow x_i)}{T(x_i \rightarrow x')}\right)$$

If proposal is symmetric: $T(x' \rightarrow x_i) = T(x_i \rightarrow x')$ (Metropolis algo.)

$$A(x_i \rightarrow x') = \min\left(1, \frac{p(x')}{p(x_i)}\right)$$

See explanation on next page

- Generate random number r from $U(0,1)$ (Uniform dist on $(0,1)$)
 - If $r \leq A(x_i \rightarrow x')$, then accept: $x_{i+1} = x'$
 - If $r > A(x_i \rightarrow x')$, then reject: $x_{i+1} = x_i$
-

• Note:

- Always accepts jump to state with higher prob. ($p(x') > p(x_i)$)
- Sometimes accepts jump to state with lower prob ($p(x') < p(x_i)$) (with prob. that ensures detailed balance.)

$\left[\text{Wait long enough} \Rightarrow \text{all states should be explored (theoretically). Freedom in choosing proposal prob. dist.} \right]$

- Only depends on ratio $\frac{p(x')}{p(x)}$, so normalization constants (like the partition function) cancel! (Very important in proj. 4!)

• Why $A(x_i \rightarrow x') = \min\left(1, \frac{p(x')}{p(x_i)}\right)$?

• Using our freedom to choose one of the A 's. Detailed balance is just a requirement on the ratio

$$\frac{A(x_i \rightarrow x')}{A(x' \rightarrow x_i)} = \frac{p(x')}{p(x_i)} \quad \left(\text{Let } T(x' \rightarrow x_i) = T(x_i \rightarrow x')\right)$$

• If $p(x') > p(x_i)$

$$\Rightarrow \frac{A(x_i \rightarrow x')}{A(x' \rightarrow x_i)} = \frac{p(x')}{p(x_i)} > 1$$

Choose $\boxed{A(x_i \rightarrow x') = 1}$

• If $p(x') < p(x_i)$

$$\Rightarrow \frac{A(x_i \rightarrow x')}{A(x' \rightarrow x_i)} = \frac{p(x')}{p(x_i)} < 1$$

Choose $A(x' \rightarrow x_i) = 1$

$$\Rightarrow \boxed{A(x_i \rightarrow x') = \frac{p(x')}{p(x_i)}} < 1$$

• In combination: $\boxed{A(x_i \rightarrow x') = \min\left(1, \frac{p(x')}{p(x_i)}\right)}$

• Setting $A(x_i \rightarrow x')$ this way

– satisfies detailed balance

– lets us move to a new state as often as we can (while being consistent with detailed balance)

• Suggested algorithm for Project 4 (Ising model):

• Generate candidate state \bar{S}'

- Pick random spin from lattice
- Flip it!

This is using a uniform distr. for $T(\bar{S}_i \rightarrow \bar{S}')$ for all "neighbour states" \bar{S}' of \bar{S}_i

• Find value of $\frac{p(\bar{S}')}{p(\bar{S}_i)}$

In an efficient way!
Don't evaluate $p(\bar{S}')$ and $p(\bar{S})$ separately, and don't evaluate $\exp(\dots)$ all the time!

• Generate random number $r \sim U(0,1)$

and carry out accept/reject step: accept if $r < \frac{p(\bar{S}')}{p(\bar{S}_i)}$

Repeat
N times
= "one MC cycle"

↓
due to strong
correlations
between subseq.
samples in
the state space.

• Then use the new state \bar{S}_{it+1} to compute energy, magnetization, expectation values after this number of cycles, etc.

• Store relevant numbers

• Repeat

Two views on the MCMC in Project 4

① As model of (discretized) time evolution of system

- Each time step there are $\sim N$ random spin-environment interactions that can flip the spin. So the Markov chain

$$\bar{S}_0 \rightarrow \bar{S}_1 \rightarrow \bar{S}_2 \rightarrow \bar{S}_3 \rightarrow \dots$$

effectively means

$$\bar{S}(t_0) \rightarrow \bar{S}(t_1) \rightarrow \bar{S}(t_2) \rightarrow \bar{S}(t_3) \rightarrow \dots$$

- If system starts in state \bar{S}_0 that has low prob. under equilibrium assumption (Boltzmann), then letting it evolve will "equilibrate the system", i.e. take \bar{S} into regions of state space that are more likely in equilibrium.

↳ Connects to MCMC "burn-in"

↳ Some unfortunate terminology in the literature, which makes it sound like $p(\bar{S})$ is changing, when it's just the chain of \bar{S} samples that is converging towards a good approx to $p(\bar{S})$.

② As a method for sampling from $p(\bar{S})$

- Just some way to draw a set of samples $\bar{S}_1, \bar{S}_2, \bar{S}_3, \dots$ such that the distribution of samples (in the long run) corresponds to $p(\bar{S})$.
- Not directly related to the physics — equally applicable for any pdf, regardless of topic.
- The physics of equilibrium was already contained in our choice to use the Boltzmann dist. for $p(\bar{S})$

Burn-in

- Samples in MCMC are correlated, i.e. not "independent and identically dist. draws" (iid)

$$p(x_{i+1} | x_i) \neq p(x_{i+1})$$

- o In the limit $n \rightarrow \infty$, we will have $\frac{n_{x \in [x, x+dx]}}{n} = p(x)dx$
- ↑
number of
samples

- But we cannot do $n \rightarrow \infty$ in practice!

\Rightarrow our resulting distribution of samples (and hence our estimated expectation values)

depends on starting point, which can be unreasonably
improbable compared to the number of samples we draw!

- o Usual solution :

- Throw away first part of a chain of samples (burn-in)
- Not use every new sample (even after burn-in)
- Run many chains from different starting points and combine results.

