

Random Walk and Markov Chains

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1 Preliminaries

2 MCMC & Methods: Metropolis-Hastings and Gibbs

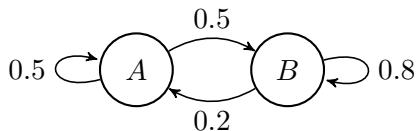
- Metropolis-Hastings Algorithm
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3 Mixing Time

Toy Example: A Two-state Random Walk

Example

Consider the two-state random walk below:



Notations

- **p**: **probability vector**. A row vector with nonnegative components that sum up to one. Each component specifies the probability mass of a vertex.
- **p_t**: **probability vector at time t**, specifying the probability masses of vertices at time *t*.
- **P = (p_{ij})**: **transition matrix**. Entry *p_{ij}* is the probability of the walk at vertex *i* selecting the edge to vertex *j*.
- The defining relationship of a random walk is

$$\mathbf{p}_t P = \mathbf{p}_{t+1}$$

Toy Example(Cont'd): Observations

- In our two-state random walk example:

$$P = \begin{pmatrix} 0.5 & 0.5 \\ 0.2 & 0.8 \end{pmatrix}$$

- We observe that, given initial distribution \mathbf{p}_1 , we can compute \mathbf{p}_t by the recursive formula:

$$\mathbf{p}_t = \mathbf{p}_{t-1}P = \cdots = \mathbf{p}_1P^{t-1}$$

- We call P^k the k -step transition matrix.

Long Term Behavior: A Computational View

- We are interested in the asymptotic behavior of \mathbf{p}_t , namely when $t \rightarrow \infty$.
- We observe that P can be diagonalized as

$$P = Q^{-1} \Lambda Q = \begin{pmatrix} 2/7 & 5/7 \\ 1 & -1 \end{pmatrix}^{-1} \begin{pmatrix} 1 & 0 \\ 0 & 0.3 \end{pmatrix} \begin{pmatrix} 2/7 & 5/7 \\ 1 & -1 \end{pmatrix}$$

- As $t \rightarrow \infty$,

$$\lim_{t \rightarrow \infty} P^{t-1} = \begin{pmatrix} 2/7 & 5/7 \\ 2/7 & 5/7 \end{pmatrix}.$$

- We can verify that, given arbitrary initial probability vector \mathbf{p}_1 , \mathbf{p}_t will converge to $(2/7, 5/7)$ after sufficiently long time.

Natural Questions Arise...

- We have seen an example of a random walk whose probability vector converges to equilibrium despite the initial probability vector.
- It is natural for us to ask whether every random walk has this property.
- Also, can two different initial distributions converge to different limits?
- Both answers are **NO**.
- However, we will make certain assumptions of the random walk and instead focus on an another distribution other than \mathbf{p}_t .

(Discrete time) Markov Chains

- In statistical literature, a concept of **Markov chains** is usually regarded as the synonym of random walks.

Definition (Markov Chain)

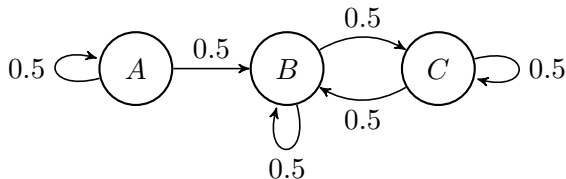
A *Markov chain* is a *stochastic process* in which future states are independent of past states given the present state.

- Consider a sequence of random variables X_1, X_2, \dots, X_t , where X_i is the state at time i . If the random variables form a Markov chain, the state at time $t + 1$ only depends on the state at time t , not on any of the past states.
- This is the **Markov property**:

$$\mathbb{P}(X_{t+1} | X_1, X_2, \dots, X_t) = \mathbb{P}(X_{t+1} | X_t)$$

Basic Assumption: Connected/Irreducible

- We say a Markov chain is **connected/irreducible** if the underlying graph is strongly connected.
- In other words, there exists a directed path from every vertex to every other vertex.
- Here is an example of a not connected Markov chain/random walk:

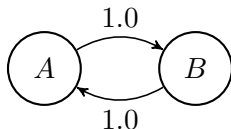


- State B cannot reach state A , thus it is not connected.

Limiting Distribution Does Not Exist

Example

We now consider a case where the probability vector does not necessarily converge. The transition diagram is given by:



- We consider $\mathbf{p}_1 = (1, 0)$, i.e., all the probability mass is at state A initially.
- It is straightforward to see $\mathbf{p}_{2k} = (0, 1)$, $\mathbf{p}_{2k+1} = (1, 0)$, for all $k \in \mathbb{N}$.
- This implies $\lim_{t \rightarrow \infty} \mathbf{p}_t$ does not exist.

Limit of the Long Term Avg. is Invariant

- However, if we consider the **long-term average probability distribution** \mathbf{a}_t given by

$$\mathbf{a}_t = \frac{1}{t}(\mathbf{p}_1 + \mathbf{p}_2 + \cdots + \mathbf{p}_t),$$

We observe that this distribution converges:

$$\lim_{t \rightarrow \infty} \mathbf{a}_t = (0.5, 0.5).$$

- We also observe that:

$$\mathbf{a}_t P = \mathbf{a}_t,$$

where P is the transition matrix

$$P = \begin{pmatrix} 0 & 1 \\ 1 & 0 \end{pmatrix}$$

Stationary Distribution

Theorem (FT of Markov Chains)

Let P be the transition probability matrix for a connected Markov chain, \mathbf{p}_t be the probability distribution at time t , and \mathbf{a}_t be the long term average probability distribution. Then there is a **unique** probability vector $\boldsymbol{\pi}$ satisfying $\boldsymbol{\pi}P = \boldsymbol{\pi}$. Moreover, for **any** starting distribution, $\lim_{t \rightarrow \infty} \mathbf{a}_t$ exists and equals $\boldsymbol{\pi}$.

- By $\boldsymbol{\pi}P = \boldsymbol{\pi}$, we have $\boldsymbol{\pi}P^k = \boldsymbol{\pi}$ for all $k \in \mathbb{N}$, which indicates running any number of steps of the Markov Chain starting with $\boldsymbol{\pi}$ leaves the distribution unchanged.
- For this reason, we call $\boldsymbol{\pi}$ the **stationary distribution**.

Proof

Proposition

For two probability distribution \mathbf{p} and \mathbf{q} :

$$\|\mathbf{p} - \mathbf{q}\|_1 = 2 \sum_i (p_i - q_i)^+ = 2 \sum_i (q_i - p_i)^+,$$

where x^+ is defined to be $\max(x, 0)$.

Lemma

The $n \times (n + 1)$ matrix $A = [P - I, \mathbf{1}_n]$ has rank n if P is the transition matrix for a connected Markov chain.

Detailed Balance

- We end this part with a sufficient condition for stationary distributions which will be of great use in the following part:

Lemma (Detailed Balance)

For a random walk on a strongly connected graph with probabilities on the edges, if the vector $\boldsymbol{\pi}$ satisfies $\pi_x p_{xy} = \pi_y p_{yx}$ for all x and y , and $\sum_x \pi_x = 1$, then $\boldsymbol{\pi}$ is the stationary distribution of the walk.

- **Proof:** Sum both sides of the detailed balance equation over y , we get $\pi_x = \sum_y \pi_y p_{yx}$. This is equivalent to $\boldsymbol{\pi}P = \boldsymbol{\pi}$, which indicates $\boldsymbol{\pi}$ is a stationary distribution.

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What is Monte Carlo?

- **Monte Carlo:** A fancy name of **simulation**.

Example: Monte Carlo Integration

We have a distribution $p(x)$ that we want to take quantities of interest from (e.g., mean, variance). To derive it analytically, we have to take integrals:

$$I = \int_{\mathbb{R}} g(x)p(x)dx$$

where $g(x)$ is some function of x (e.g. $g(x) = x$ for the mean). We can approximate the integrals via Monte Carlo, where each $x^{(i)}$ is simulated from $p(x)$:

$$\hat{I}_M = \frac{1}{M} \sum_{i=1}^M g(x^{(i)})$$

Why Monte Carlo works?

- Intuitive as Monte Carlo may seem, how can one justify that $\hat{I}_M = I$ as $M \rightarrow \infty$?

Theorem (Strong Law of Large Numbers)

Let X_1, X_2, \dots, X_M be a sequence of **independent and identically distributed (i.i.d.)** random variables, each having a finite mean $\mu = \mathbb{E}(X_i)$.

Then with probability 1,

$$\frac{X_1 + X_2 + \dots + X_M}{M} \rightarrow \mu \text{ as } M \rightarrow \infty$$

- Recall our last example, every sample point $x^{(i)}$ is simulated independently.
- What if we can't generate **independent** draws?

Sampling with a Markov Chain

- In Bayesian framework, it is essential to sample from the posterior distribution as it allows Monte Carlo estimation of all posterior quantities of interest.
- Typically, it is not possible to sample directly from a posterior. For example, we may not know the normalizing constant.
- However, we can generate *slightly dependent* draws using a **Markov chain**.
- Under certain conditions, we can still find these quantities of interest from those draws.

Ergodic Theorem

Theorem (Ergodic Theorem)

Let $x^{(1)}, x^{(2)}, \dots, x^{(M)}$ be M values from a Markov chain that is aperiodic, irreducible, and positive recurrent (then the chain is ergodic), and $\mathbb{E}[g(x)] < \infty$. Then with probability 1,

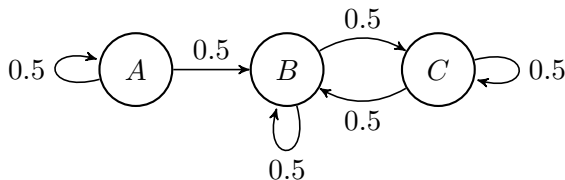
$$\frac{1}{M} \sum_{i=1}^M g[x^{(i)}] \rightarrow \sum_{\mathcal{X}} g(x) \pi(x) \text{ as } M \rightarrow \infty$$

where π is the stationary distribution.

- Note that by letting g be the indicator function $\mathbb{I}_{\{x=s\}}$, one can interpret the stationary distribution as the long-run fraction of time spent in each state.

Recap: Connected/Irreducible

- We say a Markov chain is **connected/irreducible** if the underlying graph is strongly connected.
- In other words, there exists a directed path from every vertex to every other vertex.
- Here is an example of a not connected Markov chain/random walk:



- State B cannot reach state A , thus it is not connected.

Technical Condition: Positive Recurrence

Definition (Recurrence)

A Markov chain is **recurrent** if for any given state i , if the chain starts at i , it will eventually return to i with probability 1.

Definition (Positive Recurrence)

A Markov chain is **positive recurrent** if the expected return time to state i is finite; otherwise it is **null recurrent**.

- The *simple symmetric random walk* on \mathbb{Z} is null recurrent.

Identifying Positive Recurrence

Theorem (Positive Recurrence & Stationary Distribution)

Suppose $\{X_n\}$ is an irreducible Markov chain with transition matrix P . Then $\{X_n\}$ is positive recurrent if and only if there exists a (non-negative, summing to 1) solution π , to the set of linear equations $\pi = \pi P$.

Moreover, the stationary distribution π is given by:

$$\pi_i = \frac{1}{\mathbb{E}(T_{ii})} > 0,$$

where $\mathbb{E}(T_{ii})$ is the expected return time to state i .

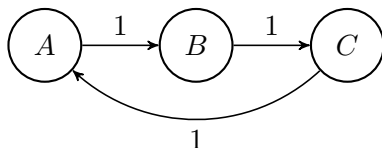
- Intuition: On average, the chain visits state i once every $\mathbb{E}(T_{ii})$ amount of time.

Technical Condition: Aperiodicity

Definition (Aperiodicity)

A Markov chain is **aperiodic** if for any set A , the number of steps required to return to A must not always be a multiple of some value k .

- Here is an example of a periodic Markov chain:



- It always takes $3k$ ($k \in \mathbb{N}$) steps for the chain to return to state A . Thus the chain is periodic.

Goal Revisited

- **Goal:** We want to generate **slightly dependent** samples from a known distribution using a **Markov chain** in order to use **Ergodic Theorem**.
- Now we will introduce two algorithms to do it.

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Metropolis-Hastings Algorithm: Overview

- Given a target distribution p over the states, the *Metropolis-Hastings algorithm* is as follows:
 - Pick an initial state: $X^{(0)} = x$.
 - At iteration t , suppose $X^{(t)} = y$, propose a move to z with probability $q(z|y)$.
 - Compute the acceptance ratio:

$$r(z|y) = \frac{p(z)q(y|z)}{p(y)q(z|y)}$$

- Accept the proposed move (i.e., $X^{(t+1)} = z$) with probability

$$\alpha(z|y) = \min\{1, r(z|y)\}.$$

Otherwise, $X^{(t+1)} = X^{(t)} = y$

- Repeat 2 ~ 4.

Justification of M-H Algorithm

Theorem (Target distribution is stationary)

The Markov chain with transition probabilities arising from the *Metropolis-Hastings algorithm* has the distribution p as a stationary distribution.

Justification of M-H Algorithm

- **Proof:** The transition probability from state i to j of this chain constructed by M-H algorithm is given by $q(j|i)\alpha(j|i)$.
- Without loss of generality, assume $p(j)q(i|j) < p(i)q(j|i)$, then

$$\begin{aligned} p(i)q(j|i)\alpha(j|i) &= p(i)q(j|i) \cdot \frac{p(j)q(i|j)}{p(i)q(j|i)} \\ &= p(j)q(i|j) \cdot 1 \\ &= p(j)q(i|j)\alpha(i|j), \end{aligned}$$

which is the **detailed balance equation**.

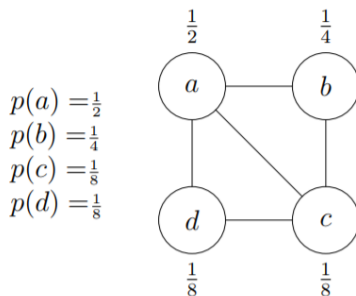
- By the previous lemma, p is the stationary distribution indeed.

Random Walk Metropolis Sampling

- To simplify things, we can have a *symmetric* proposal distribution, i.e., $q(y|x) = q(x|y)$, the acceptance ratio is simply $r(y|x) = p(y)/p(x)$. We call this **random walk Metropolis sampling**.
- If $p > 0$, it is not difficult to establish the ergodicity of this chain.
- This chain favors “heavier” states (with higher p_x), since heavier states have relatively low acceptance rates. .

Example

- We consider the example given in the textbook. The target distribution p is given below.
- We further assume that choosing any edge at a vertex has equal probability.



Python Code Implementation

```
import numpy as np
import matplotlib.pyplot as plt

n = 200000
path = [0]
pr = [1/2, 1/4, 1/8, 1/8]
proposal = [[0, 1/3, 1/3, 1/3], [1/2, 0, 1/2, 0], [1/3, 1/3, 0, 1/3], [1/2, 0, 1/2, 0]]
count = [1, 0, 0, 0]

for i in range(n-1):
    now = path[i]
    new = np.random.choice([0, 1, 2, 3], p=proposal[now])
    r = pr[new]*proposal[new][now]/(pr[now]*proposal[now][new])
    accept = min(1, r)
    gen = np.random.uniform()
    if accept > gen:
        path.append(new)
        count[new] += 1
    else:
        path.append(now)
        count[now] += 1

freq = [i/200000 for i in count]
print(freq)

plt.plot(path[100000::500], lw=1)
plt.show()
```

Figure: Python code for M-H algorithm

Simulation Results

[0.50136, 0.24929, 0.12413, 0.12522]

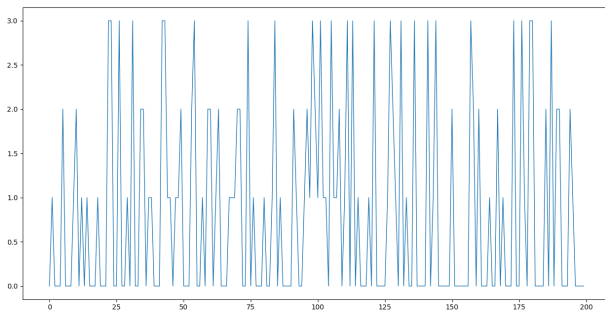


Figure: Simulated stationary distribution & Trace plot

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Gibbs Sampling: Idea

- *Gibbs sampling* is a technique to sample from **multivariate distributions**.
- The basic idea is to split the multidimensional vector into scalars.
- The beauty of this technique lies in that it simplifies a complex, high-dimensional problem by breaking it down into simple, low-dimensional problems.
- **Note:** We can only use Gibbs sampling if we know the **full conditional** distributions of the variables.

Full Conditional Distribution

- Suppose we have a joint distribution $p(x_1, x_2, \dots, x_d)$.
- The full conditional distribution of variable x_j is: $p(x_j | x_{-j})$, where x_{-j} denotes all variables except x_j .
- When the joint distribution is known, it is not difficult to find the full conditionals.

Gibbs Sampling: Algorithm

- To generate samples of $\mathbf{x} = (x_1, x_2, \dots, x_d)$ given a target distribution $p(\mathbf{x})$, do the following steps:
 - 1 Pick an initial state $\mathbf{x}^{(0)}$.
 - 2 At iteration t , current state $\mathbf{x}^{(t)} = (x_1^{(t)}, x_2^{(t)}, \dots, x_d^{(t)})$. Randomly choose a coordinate x_i to update, while leaving the rest to be unchanged. WLOG, let the coordinate be the first: x_1 .
 - 3 Draw $x_1^{(t+1)}$ from $p(x_1 | x_2, \dots, x_d)$
 - 4 Then $\mathbf{x}^{(t+1)} = (x_1^{(t+1)}, x_2^{(t+1)}, \dots, x_d^{(t+1)}) = (x_1^{(t+1)}, x_2^{(t)}, \dots, x_d^{(t)})$
 - 5 Repeat 2 \sim 4.

Selecting the Coordinate

- Randomly picking a coordinate to update is not the only scheme to choose the coordinate to update. Another option is to sequentially scan the coordinates from x_1 to x_d .

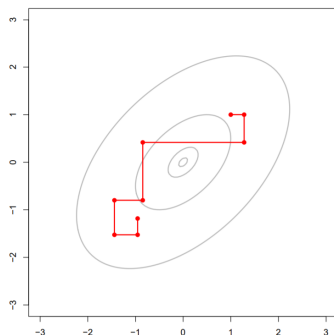


Figure: An illustration of the sequentially scanning scheme

Justification of Gibbs Sampling

- Let \mathbf{x} and \mathbf{y} be two states that differ in only one coordinate, say the first coordinate.
- Then the transition probability from \mathbf{x} to \mathbf{y} is given by:

$$p_{\mathbf{xy}} = \frac{1}{d} p(y_1 | x_2, \dots, x_d).$$

- Note that the normalizing constant is $1/d$ because $\sum_{y_1} p(y_1 | x_2, \dots, x_d) = 1$, and there are totally d directions to move towards.

Justification of Gibbs Sampling

- Similarly,

$$\begin{aligned}p_{\mathbf{y}\mathbf{x}} &= \frac{1}{d} p(x_1 | y_2, \dots, y_d) \\ &= \frac{1}{d} p(x_1 | x_2, \dots, x_d)\end{aligned}$$

since the algorithm changes only one coordinate at a time.

- By **Law of Total Probability**,

$$p_{\mathbf{x}\mathbf{y}} = \frac{1}{d} \frac{p(\mathbf{x})}{p(x_2, \dots, x_d)}, \quad p_{\mathbf{y}\mathbf{x}} = \frac{1}{d} \frac{p(\mathbf{y})}{p(x_2, \dots, x_d)}$$

Justification of Gibbs Sampling

- This is just

$$p(\mathbf{x})p_{\mathbf{xy}} = p(\mathbf{y})p_{\mathbf{yx}},$$

which is again the **detailed balance** equation, indicating that p is the stationary distribution.

Gibbs Sampling: Metropolis-Hastings in Disguise

- *Gibbs Sampling* is actually a special case of *Metropolis-Hastings algorithm*, although they look quite different.
- We can see $p_{\mathbf{xy}}$ as the proposal distribution in M-H algorithm.
- The acceptance ratio of any move is 1, i.e. all moves that are proposed are accepted.
- Recall acceptance ratio:

$$r(y|x) = \frac{p(y)q(x|y)}{p(x)q(y|x)}.$$

Summary

- **Metropolis-Hastings Algorithm:** Constructing a Markov chain with target distribution in an accept-reject manner.
- **Gibbs Sampling:** A special form of **Metropolis-Hastings Algorithm** that converts a high-dimensional problem into low-dimensional (usually 1) problems.
- Both algorithm works when the state space is a continuum, where $p(x)$ is changed to the density.

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Motivation

- We can regard both the M-H algorithm and Gibbs sampling as random walks.
- We have demonstrated that no matter what initial state is picked, the walk will eventually converge.
- However, it is also intuitive that the first few states will be highly dependent on initial state.
- A natural question will be **how fast the walk starts to reflect the stationary probability?**

We will assume our Markov chain is connected in the following part.

Random Walks on Edge-weighted Undirected Graphs

- We exploit one nice property of the random walks involved in the M-H algorithm and Gibbs sampling: they are random walks on edge-weighted undirected graphs.
- These Markov chains are derived from electrical networks.

Conductance: A Notion from Electrical Networks

- Given a network of resistors, the *conductance* of edge (x, y) is denoted c_{xy} and the normalizing constant c_x equals $\sum_y c_{xy}$.
- The Markov chain has transition probabilities proportional to edge conductances, i.e.,

$$p_{xy} = \frac{c_{xy}}{c_x}$$

- Since $c_{xy} = c_{yx}$, we have

$$c_x p_{xy} = c_{xy} = c_{yx} = \frac{c_y}{c_{yx}} = c_y p_{yx},$$

we have from the detailed balance equation that the stationary distribution π is given by $\pi_i = c_i / \sum_x c_x$.

Time-reversibility

- A Markov chain satisfying the detailed balance equation is said to be **time-reversible**.
- The name comes from the fact that for a particular path (i_1, i_2, \dots, i_k) , the probability of observing the path is the same as observing its reversal:

$$\pi_{i_1} p_{i_1, i_2} p_{i_2, i_3} \cdots p_{i_{k-1}, i_k} = \pi_{i_k} p_{i_k, i_{k-1}} p_{i_{k-1}, i_{k-2}} \cdots p_{i_2, i_1}$$

- Given a sequence of states seen, one cannot tell whether the time runs forward or backward.

Slowly Mixing Random Walks

- In general, there are certain random walks that takes a long time to converge.

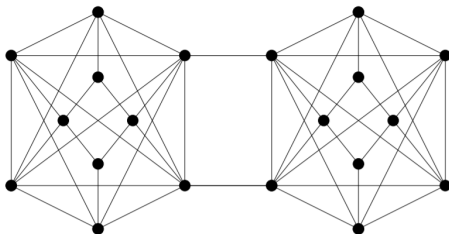


Figure: A network with a constriction

- The rapid mixing of a random walk on this graph is restricted by the narrow passage between two big components.

ϵ -mixing Time

Definition (ϵ -mixing Time)

Fix $\epsilon > 0$. The ϵ -mixing time of a Markov chain is the minimum integer t such that for any starting distribution \mathbf{p}_0 , the 1-norm distance between the t -step running average probability distribution \mathbf{a}_t and the stationary distribution $\boldsymbol{\pi}$ is at most ϵ .

Normalized Conductance

Definition (Normalized Conductance)

For a subset S of vertices, let $\pi(S)$ denote $\sum_{x \in S} \pi_x$. The normalized conductance $\Phi(S)$ of set S is

$$\Phi(S) = \frac{\sum_{(x,y) \in (S, \bar{S})} \pi_x p_{xy}}{\min(\pi(S), \pi(\bar{S}))}$$

- Observe that conductance is symmetric, i.e., $\Phi(S) = \Phi(\bar{S})$

Interpreting the Normalized Conductance

- Suppose WLOG that $\pi(S) \leq \pi(\bar{S})$. Then we can write $\Phi(S)$ as:

$$\Phi(S) = \sum_{x \in S} \frac{\pi_x}{\pi(S)} \sum_{y \in \bar{S}} p_{xy}.$$

- The red term is the probability that the walk is in state x given that the walk is in set S .
- The blue term is the probability of stepping from x to \bar{S} in one step.
- Since the red terms sum to 1, it can be seen as a distribution. $\Phi(S)$ is thus the overall probability of stepping to \bar{S} from S in one step.

Interpreting the Normalized Conductance

- Since the number of step needed to get into \bar{S} follows $Geo(\Phi(S))$, the expected number of steps needed to get into \bar{S} is $1/\Phi(S)$.
- Clearly, to be close to the stationary distribution, we must at least get to \bar{S} once.
- Hence, $1/\Phi(S)$ is a lower bound of mixing time.
- And since we can choose any S to start with, mixing time is lower bounded by the minimum over all S of $\Phi(S)$.

Normalized Conductance of the Markov Chain

Definition (Normalized Conductance of the Markov Chain)

The normalized conductance of the Markov chain, denoted Φ , is defined by

$$\Phi = \min_S \Phi(S).$$

Finding the ϵ -mixing time

Theorem (Mixing Time for Undirected Graph)

The ϵ -mixing time of a random walk on an undirected graph is

$$O\left(\frac{\ln(1/\pi_{\min})}{\Phi^2 \epsilon^3}\right)$$

where π_{\min} is the minimum stationary probability of any state.

Proof

Lemma

Suppose G_1, \dots, G_r and u_1, \dots, u_{r+1} defined as before, then

$$\pi(G_1 \cup G_2 \cup \dots \cup G_k)(u_k - u_{k+1}) \leq \frac{8}{t\Phi\epsilon} + \frac{2}{t}$$

Example: Mixing Time of 1-D Lattice

- Consider a random walk on an undirected graph consisting of an $2n$ -vertex path with self-loops at the both ends. Conductance of each edge (self-loops included) is the same.
- The stationary distribution is thus uniform over all states.
- The set with minimum normalized conductance is the set with probability $\pi \leq 1/2$ and the maximum number of vertices with the minimum number of edges leaving it.
- This is just the set with the first n vertices. (Why?)

Example: Mixing Time of 1-D Lattice (Cont'd)

- The conductance from S to \bar{S} is $\pi_n p_{n,n+1} = \frac{1}{4n} = O(\frac{1}{n})$.
- $\pi(S) = \pi(\bar{S}) = \frac{1}{2}$
- Hence, $\Phi = 2\pi_n p_{n,n+1} = O(\frac{1}{n})$
- The mixing time is thus $O(\frac{n^2 \ln n}{\epsilon})$.