

## Lecture 12: Random walks, Markov chains, and how to analyse them

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Scribe:

Today we study random walks on graphs. When the graph is allowed to be directed and weighted, such a walk is also called a *Markov Chain*. These are ubiquitous in modeling many real-life settings.

EXAMPLE 1 (DRUNKARD'S WALK) There is a sequence of  $2n + 1$  pubs on a street. A drunkard starts at the middle house. At every time step, if he is at pub number  $i$ , then with probability  $1/2$  he goes to pub number  $i - 1$  and with probability  $1/2$  to pub  $i + 1$ . How many time steps does it take him to reach either the first or the last pub?

Thinking a bit, we quickly realize that the first  $m$  steps correspond to  $m$  coin tosses, and the distance from the starting point is simply the difference between the number of heads and the number of tails. We need this difference to be  $n$ . Recall that the number of heads is distributed like a normal distribution with mean  $m/2$  and standard deviation  $\sqrt{m}/2$ . Thus  $m$  needs to be of the order of  $n^2$  before there is a good chance of this random variable taking the value  $m + n/2$ .

Thus being drunk slows down the poor guy by a quadratic factor.

EXAMPLE 2 (EXERCISE) Suppose the drunkard does his random walk in a city that's designed like a grid. At each step he goes North/South/East/West by one block with probability  $1/4$ . How many steps does it take him to get to his intended address, which is  $n$  blocks north and  $n$  blocks east away?

Random walks in space are sometimes called *Brownian motion*, after botanist Robert Brown, who in 1826 peered at a drop of water using a microscope and observed tiny particles (such as pollen grains and other impurities) in it performing strange random-looking movements. He probably saw motion similar to the one in the above figure. Explaining this movement was a big open problem. In 1905, during his "miraculous year" (when he solved 3 famous open problems in physics) Einstein explained Brownian motion as a random walk in space caused by the little momentum being imparted to the pollen in random directions by the (invisible) molecules of water. This theoretical prediction was soon experimentally confirmed and seen as a "proof" of the existence of molecules. Today random walks and brownian motion are used to model the movements of many systems, including stock prices.

EXAMPLE 3 (RANDOM WALKS ON GRAPH) We can consider a random walk on a  $d$ -regular graph  $G = (V, E)$  instead of in physical space. The particle starts at some vertex  $v_0$  and at each step, if it is at a vertex  $u$ , it picks a random edge of  $u$  with probability  $1/d$  and then moves to the other vertex in that edge. There is also a *lazy* version of this walk where he stays at  $u$  with probability  $1/2$  and moves to a random neighbor with probability  $1/2d$ .

Thus the drunkard's walk can be viewed as a random walk on a line graph.

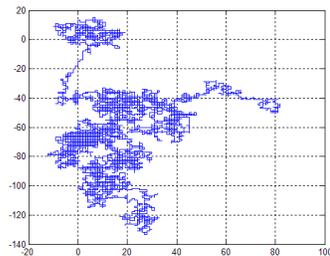


Figure 1: A 2D Random Walk

One can similarly consider random walks on directed graph (randomly pick an outgoing edge out of  $u$  to leave from) and walks on weighted graph (pick an edge with probability proportional to its weight). Walks on directed weighted graphs are called *Markov Chains*.

In a random walk, the next step does not depend upon the previous history of steps, only on the current position/state of the moving particle. In general, the term *markovian* refers to systems with a “memoryless” property. In an earlier lecture we encountered Markov Decision Processes, which also had this memoryless property.

EXAMPLE 4 (BIGRAM AND TRIGRAM MODELS IN SPEECH RECOGNITION) Language recognition systems work by constantly predicting what’s coming next. Having heard the first  $i$  words they try to generate a prediction of the  $i + 1$ th word<sup>1</sup>. This is a very complicated piece of software, and one underlying idea is to model language generation as a markov chain. (This is not an exact model; language is known to not be markovian, at least in the simple way described below.)

The simplest idea would be to model this as a markov chain on the words of a dictionary. Recall that everyday English has about 5,000 words. A simple markovian model consists of thinking of a piece of text as a random walk on a space with 5000 states (= words). A state corresponds to the last word that was just seen. For each word pair  $w_1, w_2$  there is a probability  $p_{w_1, w_2}$  of going from  $w_1$  to  $w_2$ . According to this Markovian model, the probability of generating a sentence with the words  $w_1, w_2, w_3, w_4$  is  $q_{w_1} p_{w_1 w_2} p_{w_2 w_3} p_{w_3 w_4}$  where  $q_{w_1}$  is the probability that the first word is  $w_1$ .

To actually fit such a model to real-life text data, we have to estimate 5,000 probabilities  $q_{w_1}$  for all words and  $(5,000)^2$  probabilities  $p_{w_1 w_2}$  for all word pairs. Here

$$p_{w_1 w_2} = \Pr[w_2 \mid w_1] = \frac{\Pr[w_2 w_1]}{\Pr[w_1]},$$

namely, the probability that word  $w_2$  is the next word given that the last word was  $w_1$ .

One can derive empirical values of these probabilities using a sufficiently large text corpus. (Realize that we have to estimate 25 million numbers, which requires either a very large text corpus or using some shortcuts.)

<sup>1</sup>You can see this in the typing box on smartphones, which always display their guesses of the next word you are going to type. This lets you save time by clicking the correct guess.

An even better model in practice is a *trigram model* which uses the previous two words to predict the next word. This involves a markov chain containing one state for every *pair* of words. Thus the model is specified by  $(5,000)^3$  numbers of the form  $\Pr[w_3 | w_2w_1]$ . Fitting such a model is beyond the reach of current computers but we won't discuss the shortcuts that need to be taken.

## 1 Recasting a random walk as linear algebra

A *Markov chain* is a discrete-time stochastic process on  $n$  states defined in terms of a transition probability matrix ( $M$ ) with rows  $i$  and columns  $j$ .

$$\mathbf{M} = (M_{ij})$$

where  $M_{ij}$  corresponds to the probability that the state at time step  $t + 1$  will be  $j$ , given that the state at time  $t$  is  $i$ . This process is *memoryless* in the sense that this transition probability does not depend upon the history of previous transitions.

Therefore, each row in the matrix  $\mathbf{M}$  is a distribution, implying  $M_{ij} \geq 0 \forall i, j \in S$  and  $\sum_j M_{ij} = 1$ . The bigram or trigram models are examples of Markov chains.

Using a slight twist in the viewpoint we can use linear algebra to analyse random walks. Instead of thinking of the drunkard as being at a specific point in the state space, we think of the vector that specifies his probability of being at point  $i \in S$ . Then the randomness goes away and this vector evolves according to deterministic rules. Let us understand this evolution.

Let the initial distribution be given by the row vector  $\mathbf{x} \in \mathfrak{R}^n$ ,  $x_i \geq 0$  and  $\sum_i x_i = 1$ . After one step, the probability of being at space  $i$  is  $\sum_j x_j M_{ji}$ , which corresponds to a new distribution  $\mathbf{xM}$ . It is easy to see that  $\mathbf{xM}$  is again a distribution.

Sometimes it is useful to think of  $x$  as describing the amount of *probability fluid* sitting at each node, such that the sum of the amounts is 1. After one step, the fluid sitting at node  $i$  distributes to its neighbors, such that  $M_{ij}$  fraction goes to  $j$ .

Suppose we take two steps in this Markov chain. The memoryless property implies that the probability of going from  $i$  to  $j$  is  $\sum_k M_{ik}M_{kj}$ , which is just the  $(i, j)$ th entry of the matrix  $M^2$ . In general taking  $t$  steps in the Markov chain corresponds to the matrix  $M^t$ , and the state at the end is  $\mathbf{xM}^t$ . Thus the

**DEFINITION 1** A distribution  $\pi$  for the Markov chain  $\mathbf{M}$  is a stationary distribution if  $\pi\mathbf{M} = \pi$ .

**EXAMPLE 5 (DRUNKARD'S WALK ON  $n$ -CYCLE)** Consider a Markov chain defined by the following random walk on the nodes of an  $n$ -cycle. At each step, stay at the same node with probability  $1/2$ . Go left with probability  $1/4$  and right with probability  $1/4$ .

The uniform distribution, which assigns probability  $1/n$  to each node, is a stationary distribution for this chain, since it is unchanged after applying one step of the chain.

**DEFINITION 2** A Markov chain  $\mathbf{M}$  is ergodic if there exists a unique stationary distribution  $\pi$  and for every (initial) distribution  $\mathbf{x}$  the limit  $\lim_{t \rightarrow \infty} \mathbf{xM}^t = \pi$ .

In other words, no matter what initial distribution you choose, if you let it evolve long enough the distribution converges to the stationary distribution. Some basic questions are when stationary distributions exist, whether or not they are unique, and how fast the Markov chain converges to the stationary distribution.

Does Definition 1 remind you of something? Almost all of you know about eigenvalues, and you can see that the definition requires  $\pi$  to be an eigenvector which has all nonnegative coordinates and whose corresponding eigenvalue is 1.

In today's lecture we will be interested in Markov chains corresponding to undirected  $d$ -regular graphs, where the math is easier because the underlying matrix is *symmetric*:  $M_{ij} = M_{ji}$ .

**Eigenvalues.** Recall that if  $M \in \mathbb{R}^{n \times n}$  is a square symmetric matrix of  $n$  rows and columns then an **eigenvalue** of  $M$  is a scalar  $\lambda \in \mathbb{R}$  such that exists a vector  $x \in \mathbb{R}^n$  for which  $M \cdot x = \lambda \cdot x$ . The vector  $x$  is called the **eigenvector** corresponding to the eigenvalue  $\lambda$ .  $M$  has  $n$  real eigenvalues denoted  $\lambda_1 \leq \dots \leq \lambda_n$ . (The multiset of eigenvalues is called the *spectrum*.) The eigenvectors associated with these eigenvalues form an orthogonal basis for the vector space  $\mathbb{R}^n$  (for any two such vectors the inner product is zero and all vectors are linear independent). The word *eigenvector* comes from German, and it means “one’s own vector.” The eigenvectors are  $n$  preferred directions  $u_1, u_2, \dots, u_n$  for the matrix, such that applying the matrix on these directions amounts to simple scaling by the corresponding eigenvalue. Furthermore these eigenvectors span  $\mathbb{R}^n$  so every vector  $x$  can be written as a linear combination of these.

EXAMPLE 6 We show that every eigenvalue  $\lambda$  of  $M$  is at most 1. Suppose  $\vec{e}$  is the corresponding eigenvector. Say the largest coordinate is  $i$ . Then  $\lambda e_i = \sum_{j:\{i,j\} \in E} \frac{1}{d} e_j$  by definition. If  $\lambda > 1$  then at least one of the neighbors must have  $e_j > e_i$ , which is a contradiction. By similar argument we conclude that every eigenvalue of  $M$  is at most  $-1$  in absolute value.

## 1.1 Mixing Time

Informally, the *mixing time* of a Markov chain is the time it takes to reach “nearly stationary” distribution from any arbitrary starting distribution.

DEFINITION 3 *The mixing time of an ergodic Markov chain  $M$  is  $t$  if for every starting distribution  $x$ , the distribution  $xM^t$  satisfies  $|xM^t - \pi|_1 \leq 1/4$ . (Here  $|\cdot|_1$  denotes the  $\ell_1$  norm and the constant “1/4” is arbitrary.)*

EXAMPLE 7 (MIXING TIME OF DRUNKARD’S WALK ON A CYCLE) Let us consider the mixing time of the walk in Example 5. Suppose the initial distribution concentrates all probability at state 0. Then  $2t$  steps correspond to about  $t$  random steps (= coin tosses) since with probability  $1/2$  the drunk does not move. Thus the location of the drunk is

$$(\#(\text{Heads}) - \#(\text{Tails})) \pmod{n}.$$

As argued earlier, it takes  $\Omega(n^2)$  steps for the walk to reach the other half of the circle with any reasonable probability, which implies that the mixing time is  $\Omega(n^2)$ . We will soon see that this lowerbound is fairly tight; the walk takes about  $O(n^2 \log n)$  steps to mix well.

## 2 Upper bounding the mixing time (undirected $d$ -regular graphs)

For simplicity we restrict attention to random walks on regular graphs. Let  $M$  be a Markov chain on a  $d$ -regular undirected graph with an adjacency matrix  $A$ . Then, clearly  $M = \frac{1}{d}A$ .

Clearly,  $\frac{1}{n}\vec{1}$  is a stationary distribution, which means it is an eigenvector of  $M$ . What is the mixing time? In other words if we start in the initial distribution  $\mathbf{x}$  then how fast does  $\mathbf{x}M^t$  converge to  $\vec{1}$ ?

First, let's identify two hurdles that would prevent such convergence, and in fact prevent the graph from having a unique stationary distribution. (a) *Being disconnected*: if the walk starts in a vertex in one connected component, it never visits another component, and vice versa. So two walks starting in the two components cannot converge to the same distribution, no longer how long we run them. (b) *Being bipartite*: This means the graph consists of two sets  $A, B$  such that there are no edges within  $A$  and within  $B$ ; all edges go between  $A$  and  $B$ . Then the walk starting in  $A$  will bounce back and forth between  $A$  and  $B$  and thus not converge.

EXAMPLE 8 (Exercise: ) Show that if the graph is connected, then every eigenvalue of  $M$  apart from the first one is strictly less than 1. However, the value  $-1$  is still possible. Show that if  $-1$  is an eigenvalue then the graph is bipartite.

Note that if  $\mathbf{x}$  is a distribution,  $\mathbf{x}$  can be written as

$$\mathbf{x} = \frac{1}{n}\vec{1} + \sum_{i=2}^n \alpha_i \mathbf{e}_i$$

where  $\mathbf{e}_i$  are the eigenvectors of  $M$  which form an orthogonal basis and  $\vec{1}$  is the first eigenvector with eigenvalue 1. (Clearly,  $\mathbf{x}$  can be written as a combination of the eigenvectors; the observation here is that the coefficient in front of the first eigenvector  $\vec{1}$  is  $\vec{1} \cdot \mathbf{x} / \|\vec{1}\|_2^2$  which is  $\frac{1}{n} \sum_i x_i = \frac{1}{n}$ .)

$$\begin{aligned} M^t \mathbf{x} &= M^{t-1}(M\mathbf{x}) \\ &= M^{t-1}\left(\frac{1}{n}\vec{1} + \sum_{i=2}^n \alpha_i \lambda_i \mathbf{e}_i\right) \\ &= M^{t-2}\left(M\left(\frac{1}{n}\vec{1} + \sum_{i=2}^n \alpha_i \lambda_i \mathbf{e}_i\right)\right) \\ &\quad \dots \\ &= \frac{1}{n}\vec{1} + \sum_{i=2}^n \alpha_i \lambda_i^t \mathbf{e}_i \end{aligned}$$

Also

$$\left\| \sum_{i=2}^n \alpha_i \lambda_i^t \mathbf{e}_i \right\|_2 \leq \lambda_{max}^t$$

where  $\lambda_{max}$  is the second largest eigenvalue of  $M$  in absolute value. (Note that we are using the fact that the total  $\ell_2$  norm of any distribution is  $\sum_i x_i^2 \leq \sum_i x_i = 1$ .)

Thus we have proved  $\|M^t \mathbf{x} - \frac{1}{n} \mathbf{1}\|_2 \leq \lambda_{max}^t$ . Mixing times were defined using  $\ell_1$  distance, but Cauchy Schwartz inequality relates the  $\ell_2$  and  $\ell_1$  distances:  $\|p\|_1 \leq \sqrt{n} \|p\|_2$ . So we have proved:

**THEOREM 1**

*The mixing time is at most  $O(\frac{\log n}{\lambda_{max}})$ .*

Note also that if we let the Markov chain run for  $O(k \log n / \lambda_{max})$  steps then the distance to uniform distribution drops to  $\exp(-k)$ . This is why we were not very fussy about the constant  $1/4$  in the definition of the mixing time earlier.

**Remark:** What if  $\lambda_{max}$  is 1 (i.e.,  $-1$  is an eigenvalue)? This breaks the proof and in fact the walk may not be ergodic. However, we can get around this problem by modifying the random walk to be *lazy*, by adding a self-loop at each node that ensures that the walk stays at a node with probability  $1/2$ . Then the matrix describing the new walk is  $\frac{1}{2}(I + M)$ , and its eigenvalues are  $\frac{1}{2}(1 + \lambda_i)$ . Now all eigenvalues are less than 1 in absolute value. This is a general technique for making walks *ergodic*.

**EXAMPLE 9 (EXERCISE)** Compute the eigenvalues of the drunkard's walk on the  $n$ -cycle and show that its mixing time is  $O(n^2 \log n)$ .

### 3 Analysis of Mixing Time for General Markov Chains

*We did not do this in class; this is extra reading for those who are interested.*

In the class we only analysed random walks on  $d$ -regular graphs and showed that they converge exponentially fast with rate given by the second largest eigenvalue of the transition matrix. Here, we prove the same fact for general ergodic Markov chains.

**THEOREM 2**

*The following are necessary and sufficient conditions for ergodicity:*

1. *connectivity:*  $\forall i, j : \mathbf{M}^t(i, j) > 0$  for some  $t$ .
2. *aperiodicity:*  $\forall i : \gcd\{t : \mathbf{M}^t(i, j) > 0\} = 1$ .

**REMARK 1** Clearly, these conditions are necessary. If the Markov chain is disconnected it cannot have a unique stationary distribution —there is a different stationary distribution for each connected component. Similarly, a bipartite graph does not have a unique distribution: if the initial distribution places all probability on one side of the bipartite graph, then the distribution at time  $t$  oscillates between the two sides depending on whether  $t$  is odd or even. Note that in a bipartite graph  $\gcd\{t : \mathbf{M}^t(i, j) > 0\} \geq 2$ . The sufficiency of these conditions is proved using eigenvalue techniques (for inspiration see the analysis of mixing time later on).

Both conditions are easily satisfied in practice. In particular, any Markov chain can be made aperiodic by adding self-loops assigned probability  $1/2$ .

DEFINITION 4 *An ergodic Markov chain is reversible if the stationary distribution  $\pi$  satisfies for all  $i, j$ ,  $\pi_i \mathbf{P}_{ij} = \pi_j \mathbf{P}_{ji}$ .*

We need a lemma first.

LEMMA 3

*Let  $M$  be the transition matrix of an ergodic Markov chain with stationary distribution  $\pi$  and eigenvalues  $\lambda_1 (= 1) \geq \lambda_2 \geq \dots \geq \lambda_n$ , corresponding to eigenvectors  $v_1 (= \pi), v_2, \dots, v_n$ . Then for any  $k \geq 2$ ,*

$$v_k \vec{\mathbf{1}} = 0.$$

PROOF: We have  $v_k M = \lambda_k v_k$ . Multiplying by  $\vec{\mathbf{1}}$  and noting that  $M \vec{\mathbf{1}} = \vec{\mathbf{1}}$ , we get

$$v_k \vec{\mathbf{1}} = \lambda_k v_k \vec{\mathbf{1}}.$$

Since the Markov chain is ergodic,  $\lambda_k \neq 1$ , so  $v_k \vec{\mathbf{1}} = 0$  as required.  $\square$

We are now ready to prove the main result concerning the exponentially fast convergence of a general ergodic Markov chain:

THEOREM 4

*In the setup of the lemma above, let  $\lambda = \max \{|\lambda_2|, |\lambda_n|\}$ . Then for any initial distribution  $x$ , we have*

$$\|xM^t - \pi\|_2 \leq \lambda^t \|x\|_2.$$

PROOF: Write  $x$  in terms of  $v_1, v_2, \dots, v_n$  as

$$x = \alpha_1 \pi + \sum_{i=2}^n \alpha_i v_i.$$

Multiplying the above equation by  $\vec{\mathbf{1}}$ , we get  $\alpha_1 = 1$  (since  $x \vec{\mathbf{1}} = \pi \vec{\mathbf{1}} = 1$ ). Therefore  $xM^t = \pi + \sum_{i=2}^n \alpha_i \lambda_i^t v_i$ , and hence

$$\|xM^t - \pi\|_2 \leq \left\| \sum_{i=2}^n \alpha_i \lambda_i^t v_i \right\|_2 \tag{1}$$

$$\leq \lambda^t \sqrt{\alpha_2^2 + \dots + \alpha_n^2} \tag{2}$$

$$\leq \lambda^t \|x\|_2, \tag{3}$$

as needed.  $\square$