

# Stochastic Calculus Notes, Lecture 2

Last modified January 25, 2007

## 1 Forward and Backward Equations for Markov chains

**1.1.** Introduction: Forward and backward equations are useful ways to get answers to quantitative questions about Markov chains. The probabilities  $u(k, t) = P(X(t) = k)$  satisfy forward equations. These allow us to compute all the numbers  $u(k, t + 1)$  once all the numbers  $u(j, t)$  are known. The expected values  $f(k, t) = E[V(X(T)) | X(t) = k]$  (for  $t < T$ ) satisfy a backward equation that allows us to calculate the numbers  $f(k, t)$  once all the  $f(j, t + 1)$  are known. A *duality* relation allows us to infer the forward equation from the backward equation, or conversely. The transition matrix is the *generator* of both equations, though in different ways. There are many related problems that have solutions involving forward and backward equations. Two treated here are hitting probabilities and random compound interest.

**1.2.** Forward equation, functional version: Let  $u(k, t) = P(X(t) = k)$ . The law of total probability gives

$$\begin{aligned} u(k, t + 1) &= P(X(t + 1) = k) \\ &= \sum_j P(X(t + 1) = k | X(t) = j) \cdot P(X(t) = j). \end{aligned}$$

Therefore

$$u(k, t + 1) = \sum_j P_{jk} u(j, t). \quad (1)$$

This is the *forward equation* for probabilities. It is also called the *Kolmogorov* forward equation or the *Chapman Kolmogorov* equation. Once  $u(j, t)$  is known for all  $j \in \mathcal{S}$ , (1) gives  $u(k, t + 1)$  for any  $k$ . Thus, we can go forward in time from  $t = 0$  to  $t = 1$ , etc. and calculate all the numbers  $u(k, t)$ .

Note that if we just wanted one number, say  $u(17, 49)$ , still we would have to calculate many related quantities, all the  $u(j, t)$  for  $t < 49$ . If the state space is too large, this direct forward equation approach may be impractical.

**1.3.** Row and column vectors: If  $A$  is an  $n \times m$  matrix, and  $B$  is an  $m \times p$  matrix, then  $AB$  has dimensions  $n \times p$ . The matrices are compatible for multiplication because the number of columns of  $A$  is the same as the number of rows of  $B$ . A matrix with just one column is a *column vector*.<sup>1</sup> A matrix with just one row is a *row vector*. Matrix-vector multiplication is a special case

---

<sup>1</sup>The physicists' more sophisticated idea that a vector is a physical quantity with certain transformation properties is "inoperative" here.

of matrix-matrix multiplication. We often denote genuine matrices (more than one row and column) with capital letters and vectors, row or column, with lower case. In particular, if  $u$  is an  $n$  component row vector, a  $1 \times n$  matrix, and  $A$  is an  $n \times n$  matrix, then  $uA$  is another  $n$  component row vector. We do not write  $Au$  for this because that would be incompatible. Matrix multiplication is always associative. For example, if  $u$  is a row vector and  $A$  and  $B$  are square matrices, then  $(uA)B = u(AB)$ . We can compute the row vector  $uA$  then multiply by  $B$ , or we can compute the  $n \times n$  matrix  $AB$  then multiply by  $u$ .

If  $u$  is a row vector, we usually denote the  $k$ -th entry by  $u_k$  instead of  $u_{1k}$ . Similarly, the  $k$ -th entry of column vector  $f$  is  $f_k$  instead of  $f_{k1}$ . If both  $u$  and  $f$  have  $n$  components, then  $uf = \sum_{k=1}^n u_k f_k$  is a  $1 \times 1$  matrix, i.e. a number. Thus, treating row and column vectors as special kinds of matrices makes the product of a row with a column vector natural. This is not the case for the product of two row or two column vectors.

**1.4.** Forward equation, matrix version: The probabilities  $u(k, t)$  form the components of a row vector,  $u(t)$ , with components  $u_k(t) = u(k, t)$  (an abuse of notation). The forward equation (1) may be expressed (check this)

$$u(t+1) = u(t)P . \quad (2)$$

Because matrix multiplication is associative, we have

$$u(t) = u(t-1)P = (u(t-2)P)P = u(t-2)P^2 = \dots = u(0)P^t . \quad (3)$$

Tricks of matrix multiplication give information about the evolution of probabilities. For example, we can write a formula for  $u(t)$  in terms of the eigenvectors and eigenvalues of  $P$ . Also, we can save effort in computing  $u(t)$  for large  $t$  by repeated squaring:

$$P \rightarrow P^2 \rightarrow (P^2)^2 = P^4 \rightarrow \dots \rightarrow P^{2^k}$$

using just  $k$  matrix multiplications. For example, this computes  $P^{1024}$  using just ten matrix multiplies, instead of a thousand.

**1.5.** Backward equation, functional version: Suppose we run the Markov chain until time  $T$  then get a “reward”,  $V(X(T))$ . For  $t \leq T$ , define the conditional expectations

$$f(k, t) = E[V(X(T)) | X(t) = k] . \quad (4)$$

This expression is used so often it often is abbreviated

$$f(k, t) = E_{k,t}[V(X(T))] .$$

These numbers  $f(k, t)$  satisfy a *backward equation* that follows from the law of total probability:

$$f(k, t) = E[V(X(T)) | X(t) = k]$$

$$\begin{aligned}
&= \sum_{j \in \mathcal{S}} E[V(X(T)) \mid X(t) = k \text{ and } X(t+1) = j] \cdot P(X(t+1) = j \mid X(t) = k) \\
f(k, t) &= \sum_{j \in \mathcal{S}} f(j, t+1) P_{kj} .
\end{aligned} \tag{5}$$

The Markov property is used to infer that

$$E[V(X(T)) \mid X(t) = k \text{ and } X(t+1) = j] = E_{j, t+1}[V(X(T))] .$$

The dynamics (5) must be supplemented with the *final condition*

$$f(k, T) = V(k) . \tag{6}$$

Using these, we may compute all the numbers  $f(k, T-1)$ , then all the numbers  $f(k, T-2)$ , etc.

**1.6.** Backward equation using modern conditional expectation: As usual,  $\mathcal{F}_t$  denotes the  $\sigma$ -algebra generated by  $X(0), \dots, X(t)$ . Define  $F(t) = E[V(X(T)) \mid \mathcal{F}_t]$ . The left side is a random variable that is measurable in  $\mathcal{F}_t$ , which means that  $F(t)$  is a function of  $(X(0), \dots, X(t))$ . The Markov property implies that  $F(t)$  actually is measurable with respect to  $\mathcal{G}_t$ , the  $\sigma$ -algebra generated by  $X(t)$  alone. This means that  $F(t)$  is a function of  $X(t)$  alone, which is to say that there is a function  $f(k, t)$  so that  $F(t) = f(X(t), t)$ , and

$$f(X(t), t) = E[V(X(T)) \mid \mathcal{F}_t] = E[V(X(T)) \mid \mathcal{G}_t] .$$

Since  $\mathcal{G}_t$  is generated by the partition  $\{k\} = \{X(t) = k\}$ , this is the same definition (4). Moreover, because  $\mathcal{F}_t \subseteq \mathcal{F}_{t+1}$  and  $F(t+1) = E[V(X(T)) \mid \mathcal{F}_{t+1}]$ , the tower property gives

$$E[V(X(T)) \mid \mathcal{F}_t] = E[F(t+1) \mid \mathcal{F}_t] ,$$

so that, again using the Markov property,

$$F(t) = E[F(t+1) \mid \mathcal{G}_t] . \tag{7}$$

Note that this is a version of the tower property. On the event  $\{X(t) = k\}$ , the right side above takes the value

$$\sum_{j \in \mathcal{S}} f(j, t+1) \cdot P(x(t+1) = j \mid X(t) = k) .$$

Thus, (7) is the same as the backward equation (5). In the continuous time versions to come, (7) will be very handy.

**1.7.** Backward equation, matrix version: For each  $t$ , we organize the numbers  $f(k, t)$  into a column vector  $f(t) = (f(1, t), f(2, t), \dots)^t$ . It is barely an abuse to write  $f(t)$  both for a function of  $k$  and a vector. After all, any computer

programmer knows that a vector really is a function of the index. The backward equation (5) then is equivalent to (check this)

$$f(t) = Pf(t+1) . \quad (8)$$

Again the associativity of matrix multiplication lets us write, for example,

$$f(t) = P^{T-t}V ,$$

writing  $V$  for the vector of values of  $V$ .

**1.8.** Invariant expectation value: We combine the conditional expectations (4) with the probabilities  $u(k, t)$  and the law of total probability to get, for any  $t$ ,

$$\begin{aligned} E[V(X(T))] &= \sum_{k \in \mathcal{S}} P(X(t) = k) \cdot E[V(X(T)) \mid X(t) = k] \\ &= \sum_{k \in \mathcal{S}} u(k, t) f(k, t) \\ &= u(t) f(t) . \end{aligned}$$

The last line is a natural example of an inner product between a row vector and a column vector. Note that the product  $E[V(X(T))] = u(t)f(t)$  does not depend on  $t$  even though  $u(t)$  and  $f(t)$  are different for different  $t$ . For this *invariance* to be possible, the forward evolution equation for  $u$  and the backward equation for  $f$  must be related.

**1.9.** Relationship between the forward and backward equations: It often is possible to derive the backward equation from the forward equation and conversely using the invariance of  $u(t)f(t)$ . For example, suppose we know that  $f(t) = Pf(t+1)$ . Then  $u(t+1)f(t+1) = u(t)f(t)$  may be rewritten  $u(t+1)f(t+1) = u(t)Pf(t+1)$ , which may be rearranged as (using rules of matrix multiplication)

$$(u(t+1) - u(t)P) f(t+1) = 0 .$$

If this is true for enough linearly independent vectors  $f(t+1)$ , then the vector  $w = u(t+1) - u(t)P$  must be zero, which is the matrix version of the forward equation (2). A theoretically minded reader can verify that enough  $f$  vectors are produced if the transition matrix is nonsingular and we choose a linearly independent family of “reward” vectors,  $V$ . In the same way, the backward evolution of  $f$  is a consequence of invariance and the forward evolution of  $u$ .

We now have two ways to evaluate  $E[V(X(T))]$ : (i) start with given  $u(0)$ , compute  $u(T) = u(0)P^T$ , evaluate  $u(T)V$ , or (ii) start with given  $V = f(T)$ , compute  $f(0) = P^T V$ , then evaluate  $u(0)f(0)$ . The former might be preferable, for example, if we had a number a number of different reward functions to evaluate. We could compute  $u(T)$  once then evaluate  $u(T)V$  for all our  $V$  vectors.

**1.10.** Duality: In its simplest form, *duality* is the relationship between a matrix and its transpose. The set of column vectors with  $n$  components is a vector space of dimension  $n$ . The set of  $n$  component row vectors is the *dual space*, which has the same dimension but may be considered to be a different space. We can combine an element of a vector space with an element of its dual to get a number: row vector  $u$  multiplied by column vector  $f$  yields the number  $uf$ . Any linear transformation on the vector space of column vectors is represented by an  $n \times n$  matrix,  $A$ . This matrix also defines a linear transformation, the *dual transformation*, on the dual space of row vectors, given by  $u \rightarrow uA$ . This is the sense in which the forward and backward equations are dual to each other.

Some people prefer not to use row vectors and instead think of organizing the probabilities  $u(k, t)$  into a column vector that is the transpose of what we called  $u(t)$ . For them, the forward equation would be written  $u(t+1) = P^t u(t)$  (note the notational problem: the  $t$  in  $P^t$  means “transpose” while the  $t$  in  $u(t)$  and  $f(t)$  refers to time.). The invariance relation for them would be  $u^t(t+1)f(t+1) = u^t(t)f(t)$ . The transpose of a matrix is often called its *dual*.

**1.11.** Comparison principles for the backward equation: There are some helpful theoretical properties of solutions of backward and forward equations. Suppose  $V_1(k)$  and  $V_2(k)$  are two payout functions, and  $f_j(k, t) = E_{k,t}[V_j(X(T))]$  (for  $j = 1$  and  $j = 2$ ). If  $V_1(k) \leq V_2(k)$  for all  $k \in \mathcal{S}$ , then clearly  $f_1(k, t) \leq f_2(k, t)$  for all  $k \in \mathcal{S}$  and all  $t \leq T$ . Similarly, if  $t_2 > t$  then  $f_j(k, t) = E_{k,t}[f_j(X(t_2), t_2)]$  (by the tower property). Therefore we have the *comparison principle*: if  $f_1(k, t_2) \leq f_2(k, t_2)$  for all  $k \in \mathcal{S}$ , then also  $f_1(k, t) \leq f_2(k, t)$  for all  $k \in \mathcal{S}$ . A related simple fact is the *maximum principle*: if  $t_2 \geq t$  then

$$f(k, t) \leq \max_{j \in \mathcal{S}} f(j, t_2) ,$$

as you move backward in time, the maximum of  $f$  cannot increase.

**1.12.**  $L^1$  contraction for the forward equation: The forward equation does not have a maximum principle, but there is something that cannot increase in time, the  $L^1$  norm of the solution. If  $w$  is an  $n$  component row vector, the  $L^1$  norm is

$$\|w\|_{L^1} = \sum_{j \in \mathcal{S}} |w_j| .$$

If the numbers  $u(j, t)$  satisfy the forward equation (1), then (proof just below)

$$\|u(t+1)\|_{L^1} \leq \|u(t)\|_{L^1} , \tag{9}$$

which can be put more explicitly as

$$\sum_{j \in \mathcal{S}} |u(j, t+1)| \leq \sum_{j \in \mathcal{S}} |u(j, t)| .$$

Of course, if  $u(j, t) = P(X(t) = j)$ , then  $u(j, t) \geq 0$  for all  $j$  and

$$\|u(t)\| = \sum_{j \in \mathcal{S}} u(j, t) = 1 .$$

This makes (9) true in the trivial sense that both sides are equal to one.

However, there are solutions to the forward equation (1) that are not probabilities, and the  $L^1$  contraction inequality (9) holds also for them. For example, suppose  $u_1(j, t)$  and  $u_2(j, t)$  are two probability distributions that satisfy (1), then the difference  $u(j, t) = u_2(j, t) - u_1(j, t)$  also satisfies (1) (the forward equation is linear). The  $L^1$  contraction principle implies that

$$\|u_2(t+1) - u_1(t+1)\|_{L^1} \leq \|u_2(t) - u_1(t)\| .$$

The  $L^1$  distance between two solutions decreases (contracts) at each time step. We will see in a future lecture that in nondegenerate cases,  $\|u_2(t) - u_1(t)\| \rightarrow 0$  as  $t \rightarrow \infty$ . Note that this  $u$  not only has  $\sum_j u(j, t) \neq 1$ , but probably  $u(j, t) < 0$  for some  $j$ .

To prove (9), start with (1) and sum over  $k$ . Note that  $P_{jk} \geq 0$  for all  $j$  and  $k$ , and that  $\sum_{k \in \mathcal{S}} P_{jk} = 1$  (see previous lecture). We use the *triangle inequality* ( $|a + b + c + \dots| \leq |a| + |b| + |c| + \dots$ ):

$$\begin{aligned} \|u(t+1)\|_{L^1} &= \sum_{k \in \mathcal{S}} |u(k, t+1)| \\ &= \sum_{k \in \mathcal{S}} \left| \sum_{j \in \mathcal{S}} u(j, t) P_{jk} \right| \\ &\leq \sum_{k \in \mathcal{S}} \sum_{j \in \mathcal{S}} |u(j, t)| P_{jk} \\ &= \sum_{j \in \mathcal{S}} |u(j, t)| \left( \sum_{k \in \mathcal{S}} P_{jk} \right) \\ &= \sum_{j \in \mathcal{S}} |u(j, t)| . \end{aligned}$$

This is the proof of (9). There is a similar proof of the comparison and maximum principles for the backward equation. Solutions of backward equations generally do not satisfy the  $L^1$  contraction principle.

**1.13.** Hitting probabilities, backwards: The hitting probability for state  $j$  up to time  $T$  is

$$P(X(t) = j \text{ for some } t \in [0, T]) . \tag{10}$$

Here and below we write  $[a, b]$  for all the *integers* between  $a$  and  $b$ , including  $a$  and/or  $b$  if they are integers. Hitting probabilities can be computed using forward or backward equations, often by modifying  $P$  and adding *boundary*

*conditions.* One backward equation approach using a backward equation starts by defining

$$f(k, t) = P(X(t') = j \text{ for some } t' \in [t, T] \mid X(t) = k) . \quad (11)$$

Clearly,

$$f(j, t) = 1 \text{ for all } t, \quad (12)$$

and

$$f(k, T) = 0 \text{ for } k \neq j. \quad (13)$$

Moreover, if  $k \neq j$ , the law of total probabilities yields a backward relation

$$f(k, t) = \sum_{l \in \mathcal{S}} P_{kl} f(l, t+1) . \quad (14)$$

The difference between this and the plain backward equation (5) is that the relations (14) hold only for *interior* states  $k \neq j$ , while the boundary condition (12) supplies the values of  $f(j, t)$ . The sum on the right of (14) includes the term corresponding to state  $l = j$ .

**1.14.** Matrix formulation: The difference between the two backward equations (one for expected final payout, one for hitting probability) becomes clearer if we express them in matrix terms. Let  $\tilde{f}(t)$  be the column vector with  $n - 1$  components consisting of the numbers  $f(k, t)$  for interior states  $k \neq j$ . Let  $g$  be the  $n - 1$  component column vector whose entries are  $g_k = P_{kj}$ , for  $k \neq j$ . Since  $f(j, t) = 1$ , we may rewrite (14) as

$$f(k, t) = \sum_{l \neq k} P_{kl} f(l, t+1) + g_k .$$

If we define the  $(n - 1) \times (n - 1)$  matrix  $\tilde{P}$  to be  $P$  with row  $j$  and column  $j$  removed, this becomes

$$\tilde{f}(t) = \tilde{P}\tilde{f}(t+1) + g . \quad (15)$$

The *final conditions* for (15) are  $\tilde{f}(T) = 0$ , because the components of  $\tilde{f}(T)$  are the hitting probabilities for states  $k \neq j$ . This means  $\tilde{f}(T - 1) = g$ ,

$$\tilde{f}(T - 2) = \tilde{P}\tilde{f}(T - 1) + g = \tilde{P}^2 g + \tilde{P}g + g .$$

Clearly, the general formula contains a geometric series of matrices. We can compute the sum using the matrix version of the trick for ordinary geometric series, which is the calculation, for any square matrix  $A$ ,

$$(I - A)(A^k + A^{k-1} + \dots + A + I) = I - A^{k+1} .$$

In our case, if  $I - \tilde{P}$  is invertable, we have

$$\tilde{f}(0) = (I - \tilde{P})^{-1} (I - \tilde{P}^T) g . \quad (16)$$

The  $n \times n$  matrix  $P$  has different properties from the  $(n-1) \times (n-1)$  matrix  $\tilde{P}$ . At least one of the eigenvalues of  $P$  is equal to one (see below), so  $(I - P)^{-1}$  never exists. By contrast, it is the normal case<sup>2</sup> that all the eigenvalues of  $\tilde{P}$  are inside the unit circle, so (16) makes sense and  $\tilde{P}^T \rightarrow 0$  exponentially as  $T \rightarrow \infty$ .

**1.15.** Hitting probabilities, forward: We also can compute the hitting probabilities (10) using a forward equation approach. Define the *survival probabilities*

$$u(k, t) = P(X(t) = k \text{ and } X(t') \neq j \text{ for } t' \in [0, t]) . \quad (17)$$

These satisfy the obvious *boundary condition*

$$u(j, t) = 0 , \quad (18)$$

and initial condition

$$u(k, 0) = 1 \text{ for } k \neq j . \quad (19)$$

The forward equation is (as the reader should check), that for  $k \neq j$ ,

$$u(k, t+1) = \sum_{l \in \mathcal{S}} u(l, t) P_{lk} . \quad (20)$$

We may include or exclude the term with  $l = j$  on the right because  $u(j, t) = 0$ . Of course, (20) applies only at interior states  $k \neq j$ . The overall probability of survival up to time  $T$  is  $\sum_{k \in \mathcal{S}} u(k, T)$  and the hitting probability is the complementary  $1 - \sum_{k \in \mathcal{S}} u(k, T)$ .

The matrix vector formulation of this involves the  $n-1$  component row vector,  $\tilde{u}(t)$ , whose components are the numbers  $u(k, t)$  for  $k \neq j$ . Using the same  $\tilde{P}$  the forward equation (20) and boundary condition (18) are equivalent to the matrix equation

$$\tilde{u}(t+1) = \tilde{u}(t)\tilde{P} .$$

If the *spectral gap*,

$$\rho = 1 - |\lambda_{max}|(\tilde{P}) ,$$

is positive (the usual case), the survival probabilities converge exponentially to zero at a rate  $(1 - \rho)^t$ .

**1.16.** Absorbing boundaries: *Absorbing boundaries* are another way to think about hitting and survival probabilities. It is clear that we could treat the problem of hitting a set of states  $\mathcal{B} \subset \mathcal{S}$  rather than a single state,  $j$ . We call the states  $j \in \mathcal{B}$  *boundary states*. The vectors  $f$ ,  $\tilde{u}$ , and the matrix  $\tilde{P}$  leave out components relating to boundary states. The absorbing boundary Markov chain is the same as the original chain (same transition probabilities) as long as the state is not one of the boundary states. In the absorbing chain, the state never again changes after it visits an absorbing boundary point. If  $\bar{P}$  is the

<sup>2</sup>See the discussion of positive recurrent, null recurrent, and transient Markov chains in a future lecture.



transition matrix of the absorbing chain and  $P$  is the original transition matrix, this means that  $\overline{P}_{jk} = P_{lk}$  if  $l$  is not a boundary state, while  $\overline{P}_{jk} = 0$  if  $j$  is a boundary state and  $k \neq j$ , and  $\overline{P}_{jk} = 1$ . This means that the chain stops changing (is absorbed) the first time it visits a boundary state. The probabilities  $u(k, t)$  for the absorbing chain are the same as the survival probabilities (17) for the original chain, if  $k \notin \mathcal{B}$ .

**1.17.** Running cost: Suppose we have a *running cost* function,  $W(x)$ , and we want to calculate

$$f = E \left[ \sum_{t=0}^T W(X(t)) \right]. \quad (21)$$

Sums like this are called *path dependent* because their value depends on the whole path, not just the final state  $X(T)$ . We can calculate (21) with the forward equation using

$$\begin{aligned} f &= \sum_{t=0}^T E[W(X(t))] \\ &= \sum_{t=0}^T u(t)W. \end{aligned} \quad (22)$$

Here  $W$  is the column vector with components  $W_k = W(k)$ . We compute the probabilities that are the components of the  $u(t)$  using the standard forward equation (2) and sum the products (22).

One backward equation approach uses the quantities

$$f(k, t) = E_{k,t} \left[ \sum_{t'=t}^T W(X(t')) \right]. \quad (23)$$

These satisfy (check this):

$$f(t) = Pf(t+1) + W. \quad (24)$$

Starting with  $f(T) = W$ , we work backwards with (24) until we reach the desired  $f(0)$ .

**1.18.** Multiplicative functionals: For some reason, a function of a function is often called a *functional*. The path,  $X(t)$ , is a function of  $t$ , so a function,  $F(X)$ , that depends on the whole path is often called a functional. Some applications call for finding the expected value of a multiplicative functional:

$$f = E \left[ \prod_{t=0}^T V(X(t)) \right]. \quad (25)$$

For example,  $X(t)$  could represent the state of a financial market and  $V(k) = 1 + r(k)$  the interest rate for state  $k$ . Then (25) would be the expected total

interest. We also can write  $V(k) = e^{W(k)}$ , so that

$$\prod V(X(t)) = \exp \left[ \sum W(X(t)) \right] = e^Z ,$$

with

$$Z(x) = \sum_{t=0}^T W(x(t)) .$$

This does not solve the problem of evaluating (25) because  $E[e^Z] \neq e^{E(Z)}$ .

The backward equation approach uses the intermediate quantities

$$f(k, t) = E_{k, t} \left[ \prod_{t'=t}^T V(X(t')) \right] .$$

The  $t' = t$  term in the product has  $V(X(t)) = V(k)$ . The final condition is  $f(k, T) = V(k)$ . The backward evolution equation is derived more or less as before:

$$\begin{aligned} f(k, t) &= E_{k, t} \left[ V(k) \prod_{t' > t} V(X(t')) \right] \\ &= V(k) E_{k, t} \left[ \prod_{t'=t+1}^T V(X(t')) \right] \\ &= V(k) E_{k, t} [f(X(t+1), t+1)] \quad (\text{the tower property}) \\ f(k, t) &= V(k) (Pf(t+1))(k) . \end{aligned} \tag{26}$$

In the last line on the right,  $f(t+1)$  is the column vector with components  $f(k, t+1)$  and  $Pf(t+1)$  is the matrix vector product. We write  $(Pf(t+1))(k)$  for the  $k^{\text{th}}$  component of the column vector  $Pf(t+1)$ . We could express the whole thing in matrix terms using  $\text{diag}(V)$ , the diagonal matrix with  $V(k)$  in the  $(k, k)$  position:

$$f(t) = \text{diag}(V) Pf(t+1) .$$

A version of (26) for Brownian motion is called the Feynman-Kac formula.

**1.19.** Branching processes: One forward equation approach to (25) leads to a different interpretation of the answer. Let  $B(k, t)$  be the event  $\{X(t) = k\}$  and  $I(k, t, X)$  the indicator function of  $B(k, t)$ . That is  $I(k, t, X) = 1$  if  $X \in B(k, t)$  (i.e.  $X(t) = k$ ), and  $I(k, t, X) = 0$  otherwise. It is in keeping with the probabilists' habit of leaving out the arguments of functions when the argument is the underlying random outcome that we write  $I(k, t)$  in place of  $I(k, t, X)$ . The probabilities  $u(k, t) = P(X(t) = k)$  satisfy  $u(k, t) = E[I(k, t)]$ . The forward equation for the quantities

$$g(k, t) = E \left[ I(k, t) \prod_{t'=0}^t V(X(t')) \right] \tag{27}$$

is (see homework):

$$g(k, t) = (g(t-1)P)(k)V(k) . \tag{28}$$

In matrix terms, it would be  $g(t) = g(t-1)P\text{diag}(V)$ .

This is also the forward equation for a *branching process* with branching factors  $V(k)$ . At time  $t$ , the branching process has  $N(k, t)$  *particles*, or *walkers*, at state  $k$ . The numbers  $N(k, t)$  are random. A time step of the branching process has two parts. First, each particle takes one step of the Markov chain. A particle at state  $j$  goes to state  $k$  with probability  $P_{jk}$ . All steps for all particles are independent. Then, each particle at state  $k$  does a *branching* or *birth/death* step in which the particle is replaced by a random number of particles with expected number  $V(k)$ . For example, if  $V(k) = 1/2$ , we could delete the particle (death) with probability half. If  $V(k) = 2.8$ , we could keep the existing particle, one new one, then add a third with probability .8. All particles are treated independently. If there are  $m$  particles in state  $k$  before the birth/death step, the expected number after the birth/death step is  $V(k)m$ . The expected number of particles,  $g(k, t) = E[N(k, t)]$ , satisfies (28).

When  $V(k) = 1$  for all  $k$  there need be no birth or death. There will be just one particle, the path  $X(t)$ . The number of particles at state  $k$  at time  $t$ ,  $N(k, t)$ , will be zero if  $X(t) \neq k$  or one if  $X(t) = k$ . In fact,  $N(k, t) = I(k, t)(X)$ . The expected values will be  $g(k, t) = E[N(k, t)] = E[I(k, t)] = u(k, t)$ .

The branching process representation of (25) is possible when  $V(k) \geq 0$  for all  $k$ . Monte Carlo methods based on branching processes are more accurate than direct Monte Carlo in many cases.

## 2 Lattices, trees, and random walk

**2.1.** Introduction: Random walk on a lattice is an important example where the abstract theory of Markov chains is used. It is the simplest model of something randomly moving through space with none of the subtlety of Brownian motion, though random walk on a lattice is a useful approximation to Brownian motion, and vice versa. The forward and backward equations take a specific simple form for lattice random walk and it is often possible to calculate or approximate the solutions by hand. Boundary conditions will be applied at the boundaries of lattices, hence the name.

We pursue forward and backward equations for several reasons. First, they often are the best way to calculate expectations and hitting probabilities. Second, many theoretical qualitative properties of specific Markov chains are understood using backward or forward equations. Third, they help explain and motivate the partial differential equations that arise as backward and forward equations for diffusion processes.

**2.2.** Simple random walk: The state space for simple random walk is the integers, positive and negative. At each time, the walker has three choices: (A) move one unit to the right, (B) do not move, (C) move one unit to the left. The

probabilities are  $P(A) = P(k \rightarrow k + 1) = a$ ,  $P(B) = P(X(t + 1) = X(t)) = b$ , and  $P(X(t + 1) = X(t) - 1) = c$ . Naturally, we need  $a$ ,  $b$ , and  $c$  to be non-negative and  $a + b + c = 1$ . The transition matrix<sup>3</sup> has  $b$  on the diagonal ( $P_{kk} = b$  for all  $k$ ),  $a$  on the *super-diagonal* ( $P_{k,k+1} = a$  for all  $k$ ), and  $c$  on the *sub diagonal*. All other matrix elements  $P_{jk}$  are zero.

This Markov chain is *homogeneous* or *translation invariant*: The probabilities of moving up or down are independent of  $X(t)$ . A *translation* by  $k$  is a shift of everything by  $k$  (I do not know why this is called “translation”). Translation invariance implies, among other things, that the probability of going from  $m$  to  $l$  in  $s$  steps is the same as the probability of going from  $m + k$  to  $l + k$  in  $s$  steps:  $P(X(t + s) = l \mid X(t) = m) = P(X(t + s) = l + k \mid X(t) = m + k)$ . It is common to simplify general discussions by choosing  $k$  so that  $X(0) = 0$ . Mathematicians often say “without loss of generality” or “w.l.o.g.” when doing so.

Often, particularly when discussing multidimensional random walk, we use  $x$ ,  $y$ , etc. instead of  $j$ ,  $k$ , etc. to denote lattice points (states of the Markov chain). Thus, we might write  $P_{xy} = P(X(t + 1) = x \mid X(t) = y)$ . As an exercise in definition unwrapping, review Lecture 1 and check that this is the same as  $P_{X(t),x} = P(X(t + 1) = x \mid \mathcal{F}_t)$ .

**2.3.** Gaussian approximation, drift, and volatility: We can write  $X(t + 1) = X(t) + Y(t)$ , where  $P(Y(t) = 1) = a$ ,  $P(Y(t) = 0) = b$ , and  $P(Y(t) = -1) = c$ . The random variables  $Y(t)$  are independent of each other because of the Markov property and homogeneity. Assuming (without loss of generality) that  $X(0) = 0$ , we have

$$X(t) = \sum_{s=0}^{t-1} Y(s) , \tag{29}$$

which expresses  $X(t)$  as a sum of *iid* (independent and identically distributed) random variables. The central limit theorem then tells us that for large  $t$ ,  $X(t)$  is approximately Gaussian with mean  $\mu t$  and variance  $\sigma^2 t$ , where  $\mu = E[Y(t)] = a - c$  and  $\sigma^2 = \text{var}[Y(t)] = a + c - (a - c)^2$ . These are called *drift* and *volatility*<sup>4</sup> respectively. The mean and variance of  $X(t)$  grow linearly in time with rate  $\mu$  and  $\sigma^2$  respectively. Figure 1 shows some probability distributions for simple random walk. The central limit theorem Gaussian approximation is somewhat accurate after 8 steps and rather more accurate after 60 steps. The effect of *skewness* is clearly visible in the top plot. Values just to the left of the mean are more likely than the central limit theorem approximation predicts and values just to the right are less likely.

**2.4.** Forward and backward equations: The probabilities  $u(x, t) = P(X(t) =$

<sup>3</sup>This “matrix” is infinite when the state space is infinite. Matrix multiplication is still defined. For example, the  $k$  component of  $uP$  is given by  $(uP)_k = \sum_j u_j P_{jk}$ . This possibly infinite sum has only three nonzero terms when  $P$  is tridiagonal.

<sup>4</sup>People use the term *volatility* in two distinct ways. In the Black Scholes theory, volatility means something else.

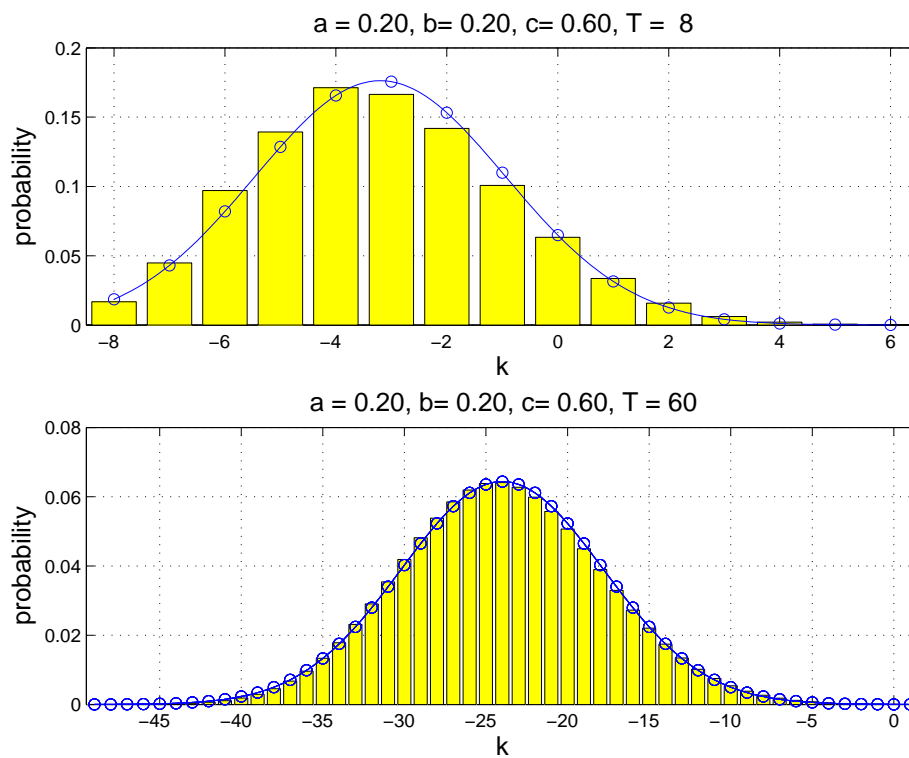


Figure 1: The probability distributions after  $T = 8$  (top) and  $T = 60$  (bottom) steps for simple random walk. The smooth curve and circles represent the central limit theorem Gaussian approximation. The plots have different probability and  $k$  scales. Values not shown have very small probability.

$x$ ) satisfy the forward equation (Check that these agree with the general theory of Lecture 1):

$$u(x, t + 1) = au(x - 1, t) + bu(x, t) + cu(x + 1, t) . \quad (30)$$

The expected values  $f(x, t) = E_{x,t}[V(X(T))]$  satisfy the backward equation

$$f(x, t) = cf(x - 1, t + 1) + b(f(x, t + 1) + af(x + 1, t + 1)) . \quad (31)$$

These equations have several differences. Note that  $a$  multiplies the value from the left in (30) and the value from the right in (31). Of course, (30) moves forward in time, computing the probabilities  $u$  at time  $t+1$  from the probabilities at time  $t$ . The backward equation (31) computes expected values at time  $t$  from those at time  $t + 1$ .

The backward equation has a variety of explicit solutions. Polynomial solutions such as,  $f(x, t) = p_2(t)x^2 + p_1(t)x + p_0(t)$  can be found by plugging into (31) and calculating formulas for  $p_2(t)$ ,  $p_1(t)$ , and  $p_0(t)$  in terms of the same coefficients at time  $t + 1$ . There also are exponential solutions  $f(x, t) = m(t)z^x$ , where  $m(t) = (cz^{-1} + b + az)m(t + 1)$ .

**2.5. Qualitative properties:** Solutions of the forward and backward equations (30) and (31) have special qualitative properties not shared by forward and backward equation solutions for general Markov chains. For random walk, both the forward and backward equation solutions have maximum and comparison principles and the  $L^1$  contraction property. For example,

$$\max_x u(x, t + 1) \leq \max_x u(x, t) .$$

The proof is a simple application of the facts that  $a + b + c = 1$  and  $a \geq 0$ ,  $b \geq 0$ , and  $c \geq 0$ . Similarly

$$\sum_x |f(x, t)| \leq \sum_x |f(x, t + 1)| .$$

We will see that in the urn model, neither of these extra inequalities is satisfied.

Another qualitative property is the *smoothing property*. If the random walk is nondegenerate ( $a > 0$ ,  $b > 0$ ,  $c > 0$ ), then solutions to forward or backward equations become smoother over time. The reader can check that

$$\begin{aligned} \sum_x u(x, t + 1)^2 &= \sum_x u(x, t)^2 \\ &- (ab + bc) \left( \sum_x (u(x + 1, t) - u(x, t))^2 \right) \\ &- ac \left( \sum_x (u(x + 1, t) - u(x - 1, t))^2 \right) . \end{aligned}$$

**2.6.** Trees: Simple random walk can be thought of as a sequence of decisions. At each time you decide: right ( $A$ ), stay ( $B$ ), or left ( $C$ ). A more general sequence of decisions is a *decision tree*. In a general decision tree, making choice  $A$  at time 0 then  $B$  at time one would have a different result than choosing first  $B$  then  $A$ . After  $t$  decisions, there could be  $3^t$  different decision paths and results.

The simple random walk decision tree is *recombining*, which means that many different decision paths lead to the same  $X(t)$ . For example, if we start with  $X(0) = 0$ , the paths  $ABB$ ,  $CAA$ ,  $BBA$ , etc. all lead to  $X(3) = 1$ . A recombining tree is much smaller than a general decision tree. For simple random walk, after  $t$  steps there are  $2t + 1$  possible states, instead of up to  $3^t$ . For  $t = 10$ , this is 21 instead of about 60 thousand.

**2.7.** Urn models: Urn models illustrate several features of more general random walks. Unlike simple random walk, urn models are *mean reverting* and have *steady state probabilities* that determine their large time behavior. We will come back to them when we discuss *scaling* in future lectures.

The simple urn contains  $n$  balls that are identical except for their color. There are  $k$  red balls and  $n - k$  green ones. At each state, someone chooses one of the balls at random with each ball equally likely to be chosen. He or she replaces the chosen ball with a fresh ball that is red with probability  $p$  and green with probability  $1 - p$ . All choices are independent. The number of red balls decreases by one if he or she removes a red ball and returns a green one. This happens with probability  $(k/n) \cdot (1 - p)$ . Similarly, the  $k \rightarrow k + 1$  probability is  $((n - k)/n) \cdot p$ . There are two ways to get a  $k \rightarrow k$  transition, pick red and replace with red (prob =  $(k/n) \cdot p$ ) or pick green and replace with green (prob =  $((n - k)/n) \cdot (1 - p)$ ). In formal terms, the state space is the integers from 0 to  $n$  and the transition probabilities are

$$P_{k,k-1} = \frac{k(1-p)}{n}, \quad P_{kk} = \frac{(2p-1)k + (1-p)n}{n}, \quad P_{k,k+1} = \frac{(n-k)p}{n},$$

$$P_{jk} = 0 \text{ otherwise.}$$

If these formulas are right, then  $P_{k,k-1} + P_{kk} + P_{k,k+1} = 1$ . (check this)

**2.8.** Urn model steady state: For the simple urn model, the probabilities  $u(k, t) = P(X(t) = k)$  converge to steady state probabilities,  $v(k)$ , as  $t \rightarrow \infty$ . This is illustrated in Figure (2). The steady state probabilities are

$$v(k) = \binom{n}{k} p^k (1-p)^{n-k}.$$

The steady state probabilities have the property that if  $u(k, t) = v(k)$  for all  $k$ , then  $u(k, t + 1) = v(k)$  also for all  $k$ . This is *statistical steady state* because the probabilities have reached steady state values though the states themselves keep changing, as in Figure (3). In matrix vector notation, we can form the row vector,  $v$ , with entries  $v(k)$ . Then  $v$  is a statistical steady state if  $vP = v$ . It is

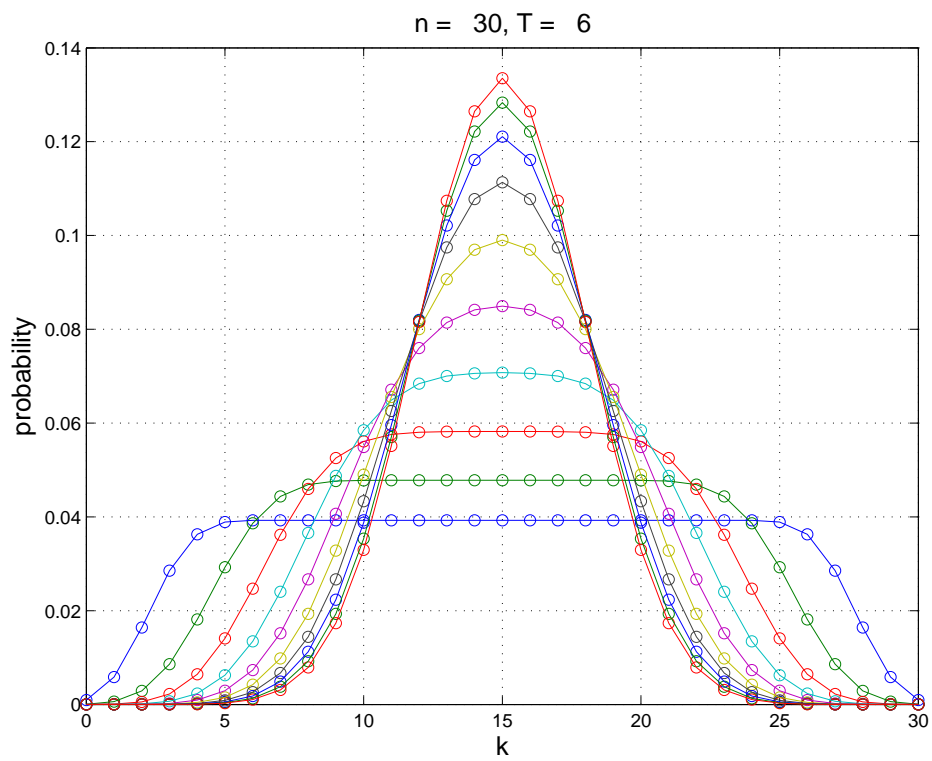


Figure 2: The probability distributions for the simple urn model plotted every  $T$  time steps. The first curve is blue, low, and flat. The last one is red and most peaked in the center. The computation starts with each state being equally likely. Over time, states near the edges become less likely.



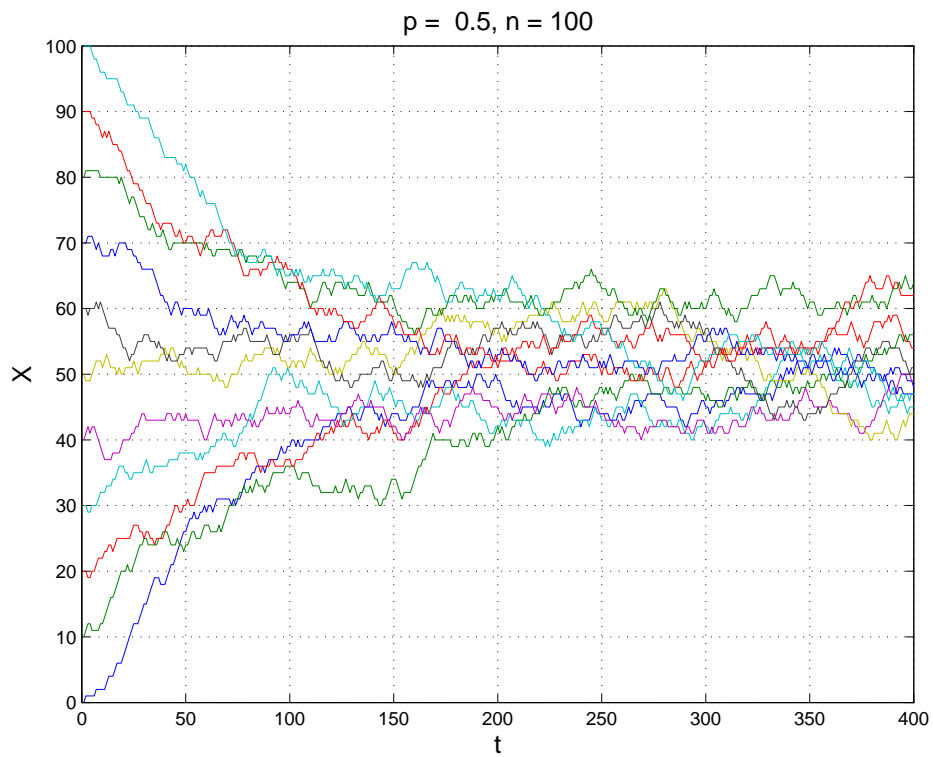


Figure 3: A Monte-Carlo sampling of 11 paths from the simple urn model. At time  $t = 0$  (the left edge), the paths are evenly spaced within the state space.

no coincidence that  $v(k)$  is the probability of getting  $k$  red balls in  $n$  independent trials with probability  $p$  for each trial. The steady state expected number of red balls is

$$E_v[X] = np ,$$

where the notation  $E_v[\cdot]$  refers to expectation in probability distribution  $v$ .

**2.9.** Urn model mean reversion: If we let  $m(t)$  be the expected value if  $X(t)$ , then a calculation using the transition probabilities gives the relation

$$m(t+1) = m(t) + \frac{1}{n} (np - m(t)) . \quad (32)$$

This relation shows not only that  $m(t) = np$  is a steady state value ( $m(t) = np$  implies  $m(t+1) = np$ ), but also that  $m(t) \rightarrow np$  as  $t \rightarrow \infty$  (if  $r(t) = m(t) - np$ , then  $r(t+1) = \alpha r(t)$  with  $|\alpha| = |1 - \frac{1}{n}| < 1$ ).

Another way of expressing mean reversion will be useful in discussing stochastic differential equations later. Because the urn Model is a Markov chain,

$$E[X(t+1) | \mathcal{F}_t] = E[X(t+1) | X(t)]$$

Again using the transition probabilities, we get

$$E[X(t+1) | \mathcal{F}_t] = X(t) + \frac{1}{n} (np - X(t)) . \quad (33)$$

If  $X(t) > np$ , we have

$$E[\Delta X(t)] = E[X(t+1) - X(t)] = \frac{1}{n} (np - X(t)) ,$$

is negative. If  $X(t) < np$ , it is positive.

**2.10.** Boundaries: The terms *boundary*, *interior*, *region*, etc. as used in the general discussion of Markov chain hitting probabilities come from applications in lattice Markov chains such as simple random walk. For example, the region  $x > \beta$  has boundary  $x = \beta$ . The quantities

$$u(x, t) = P(X(t) = x \text{ and } X(s) > \beta \text{ for } 0 \leq s \leq t)$$

satisfy the forward equation (just (1) in this special case)

$$u(x, t+1) = au(x-1, t) + bu(x, t) + cu(x+1, t)$$

for  $x > \beta$  together with the *absorbing boundary condition*  $u(\beta, t) = 0$ . We could create a finite state space Markov chain by considering a region  $\beta < x < \gamma$  with simple random walk in the *interior* together with absorbing boundaries at  $x = \beta$  and  $x = \gamma$ . Absorbing boundary conditions are also called *Dirichlet boundary conditions*.

Another way to create a finite state space Markov chain is to put *reflecting boundaries* at  $x = \beta$  and  $x = \gamma$ . This chain has the same transition probabilities as ordinary random walk in the interior ( $\beta < x < \gamma$ ). However, transitions from  $\beta$  to  $\beta - 1$  are disallowed and replaced by transitions from  $\beta$  to  $\beta + 1$ . This means changing the transition probabilities starting from  $x = \beta$  to

$$P(\beta \rightarrow \beta - 1) = P_{\beta, \beta - 1} = 0, \quad P(\beta \rightarrow \beta) = P_{\beta\beta} = b, \quad P(\beta \rightarrow \beta + 1) = P_{\beta, \beta + 1} = a + c.$$

The transition rules at  $x = \gamma$  are similarly changed to block  $\gamma \rightarrow \gamma + 1$  transitions. There is some freedom in defining the reflection rules at the boundaries. We could, for example, make  $P(\beta \rightarrow \beta) = b + c$  and  $P(\beta \rightarrow \beta + 1) = a$ , which changes the blocked transition to standing still rather than moving right. We return to this point in discussing *oblique reflection* in multidimensional random walks and diffusions.

**2.11.** Some boundary value problems: To be concrete, let us put absorbing boundaries at  $\beta = 0$  and  $\gamma = L$ . We assume the random walk starts at a state  $X(0) = x \in [0, L]$  and ask which of the boundary points is hit first. We seek to calculate

$$w(x) = P(X(t) = 0 \text{ before } X(t) = L \mid X(0) = x). \quad (34)$$

(We make the technical assumption that  $a \neq 0$  and  $c \neq 0$ , so that hitting 0 first and hitting  $L$  first both are possible.) Clearly, this satisfies *boundary conditions*  $w(0) = 1$  and  $w(L) = 0$ . For  $x \in [1, L - 1]$ , we can use the law of total probability to find equations satisfied by  $w$ . Let  $A$  be the event  $\{X(t) = 0 \text{ before } X(t) = L\}$ . Then, reasoning as before (e.g.  $P(A \mid X(0) = x \text{ and } X(1) = x - 1) = w(x - 1)$ , by the Markov property),

$$\begin{aligned} w(x) &= P(A \mid X(0) = x) \\ &= P(A \mid X(0) = x \text{ and } X(1) = x - 1) \cdot P(x \rightarrow x - 1) \\ &+ P(A \mid X(0) = x \text{ and } X(1) = x) \cdot P(x \rightarrow x) \\ &+ P(A \mid X(0) = x \text{ and } X(1) = x + 1) \cdot P(x \rightarrow x + 1) \\ &= cw(x - 1) + bw(x) + aw(x + 1). \end{aligned}$$

To summarize, we can calculate the probabilities (34) by solving the *boundary value problem*, which consists of an equation satisfied in the interior:

$$cw(x - 1) + (b - 1)w(x) + aw(x + 1) = 0, \quad (35)$$

together with the boundary conditions given above.

The solution to the boundary value problem takes different forms depending on whether  $a = c$  (no drift), or  $a \neq c$  (drift). In the case of no drift, it is easy to check<sup>5</sup> that any solution of the recurrence relation (35) takes the form  $w(x) = \alpha + \beta x$ . The boundary conditions determine  $\alpha$  and  $\beta$ . First,

<sup>5</sup>Choose  $\alpha$  and  $\beta$  so that  $w(x) = \alpha + \beta x$  for  $x = 1$  and  $x = 2$ . Since  $c \neq 0$ ,  $w(3)$  is determined by  $w(1)$  and  $w(2)$ . The value  $w(3) = \alpha + 3\beta$  satisfies  $w(3) = \frac{-1}{a}(cw(1) + (b - 1)w(2))$ .

$w(0) = 1 = \alpha + \beta \cdot 0$  gives  $\alpha = 1$ . Then,  $w(L) = 0 = \alpha + \beta L$  gives  $\beta = 1/L$ . Altogether, we have

$$P(X(t) = 0 \text{ before } X(t) = L \mid X(0) = x) = 1 - \frac{x}{L}. \quad (36)$$

The reader should make a final check that this satisfies the equation (35) and boundary conditions. Other checks are that starting close to 0 makes is more likely to hit 0 than to hit  $L$  first, and that if  $L$  is even then  $w(L/2) = 1/2$ , so it is equally likely to hit either boundary point.

In the case of drift, the solution of the recurrence relation is  $w(x) = \alpha + \beta \left(\frac{c}{a}\right)^x$ . For definiteness, we assume that the drift is to the left,  $c > a$ . The boundary conditions  $w(0) = 1$  and  $w(L) = 0$  give a system of two linear equations that determine  $\alpha$  and  $\beta$ :

$$\begin{aligned} \alpha + \beta &= 1 \\ \alpha + \beta \left(\frac{c}{a}\right)^L &= 0. \end{aligned}$$

If  $L$  is large, then  $(c/a)^L$  is exponentially large, and we have

$$\beta = \frac{1}{1 + \left(\frac{c}{a}\right)^L} \approx \left(\frac{a}{c}\right)^L.$$

and  $\alpha \approx 1$ . For  $x$  not too large, this implies that

$$w(x) \approx 1 - \left(\frac{a}{c}\right)^{L-x}.$$

This implies that if the drift is to the left and you start anywhere near the left boundary, then it is exponentially unlikely to hit the right boundary before the left one.

**2.12.** Hitting times: The *hitting time* will be

$$\tau = \min \{t \text{ with } X(0) = 0 \text{ or } X(t) = L\}.$$

The function  $f(x) = E[\tau \mid X(0) = x]$  may be found by solving another boundary value problem. The boundary conditions are  $f(0) = f(L) = 0$ . The equation to be satisfied for  $x \in [1, L-1]$  comes from the law of total probability much as before, except that  $f(x) = E[\tau \mid X(1) = x] + 1$ :

$$\begin{aligned} f(x) &= E[\tau \mid X(0) = x] \\ &= E[\tau \mid X(0) = x \text{ and } X(1) = x-1] \cdot P(x \rightarrow x-1) \\ &+ E[\tau \mid X(0) = x \text{ and } X(1) = x] \cdot P(x \rightarrow x) \\ &+ E[\tau \mid X(0) = x \text{ and } X(1) = x+1] \cdot P(x \rightarrow x+1) \\ &= cf(x-1) + bf(x) + af(x+1) + 1. \end{aligned}$$

The equation satisfied in the interior is

$$cf(x-1) + (b-1)f(x) + af(x+1) = -1. \quad (37)$$

We discuss the solution only in the zero drift case  $a = c \neq 0$ . In differential equations, you learn that the general solution to an inhomogeneous equation is the sum of the general solution to the homogeneous equation plus any particular solution to the inhomogeneous equation. In this case, the general solution to the homogeneous equation is  $\alpha + \beta x$ , as above. We guess that there is an inhomogeneous solution of the form  $f(x) = \gamma x^2$  and verify by substitution into (37)

$$\gamma(a(x-1)^2 + (b-1)x^2 + a(x+1)^2) = -1,$$

which gives  $\gamma = -1/2a$ . We now determine which of the general solutions to the inhomogeneous problem satisfies the boundary conditions:  $\alpha + \beta x - \frac{1}{2a}x^2 = 0$  when  $x = 0$  and  $x = L$ . The  $x = 0$  condition gives  $\alpha = 0$ . The  $x = L$  condition then gives

$$\beta L - \frac{1}{2a}L^2 = 0,$$

which gives  $\beta = L/(2a)$ . A little calculation involving completing the square puts the resulting solution in a helpful form:

$$\begin{aligned} f(x) &= \frac{L}{2a}x - \frac{1}{2a}x^2 \\ &= \frac{1}{2a} \left\{ -x^2 + Lx - \frac{1}{4}L^2 + \frac{1}{4}L^2 \right\} \\ f(x) &= \frac{1}{2a} \left\{ \frac{1}{4}L^2 - \left( x - \frac{L}{2} \right)^2 \right\}. \end{aligned} \quad (38)$$

This form of the answer makes it clear that the expected hitting time is symmetric about the center of the interval  $[0, L]$  and that the largest hitting times come from the center of the interval.

In particular,  $f(\frac{1}{2}L) = \frac{1}{4}L^2$ . This says that it takes on the order of  $L^2$  steps for a symmetric random walk to go a distance  $L$  from the center to the boundary. We will understand this result better when we discuss the approximation of random walk by Brownian motion in future lectures.

Perhaps more surprising is the result  $f(1) = \frac{1}{2a}(L-1)$ . You start with  $X(0) = 1$ , perform a symmetric random walk, and ask: What is the expected time until  $X(t) = 0$  or  $X(t) = L$ ? You might expect that the expected time might not be very large since you are starting next to a boundary point. The calculation shows this is wrong. The hitting time, even starting from  $x = 1$  grows linearly with  $L$ . Now suppose we set the right endpoint at infinity and calculate

$$f = E[\min\{t \text{ with } X(t) = 0\} \mid X(0) = 1].$$

The answer is  $f = \infty$  (proof:  $f \geq \frac{1}{2a}(L-1)$  for any  $L$ ). Although a symmetric random walk will eventually come to zero, the expected time to do so is infinite.

A more detailed calculation would show that for large  $t$ ,  $g(t) = P(\tau = t) \approx \text{Const} \cdot t^{-3/2}$ , so that  $E[\tau] = \sum_{t>0} tg(t)$  has a *tail* (the terms with large  $t$ ) that is approximately  $\text{Const} \cdot t^{-1/2}$ , so the sum is infinite.

**2.13.** Multidimensional lattice: The unit square lattice in  $d$  dimensions is the set of  $d$ -tuples of integers (the set of integers is called  $Z$ ):

$$x = (x_1, \dots, x_d) \text{ with } x_j \in Z \text{ for } 1 \leq j \leq d .$$

The scaled square lattice, with *lattice spacing*  $h > 0$ , is the set of points  $hx = (hx_1, \dots, hx_d)$ , where  $x$  are integer lattice points. In the present discussion, the scaling is irrelevant, so we use the unit lattice. We say that lattice points  $x$  and  $y$  are *neighbors* if

$$|x_j - y_j| \leq 1 \text{ for all coordinates } j = 1, \dots, d .$$