



What is a random walk?

The original statement of a random walk was formulated in the context of a drunken sailor. If drunkard begins at the lamp post and takes N steps of equal length in random directions, how far will the drunkard be from the lamp post? The result is related to the diffusion!

There are very many versions of random walks

Random walks have multiple applications in

- Science: physics, chemistry, biology, ...
- Medicine (in particular, spread of inflectional diseases and effects of immunization)
- Engineering
- Economics
- Sociology

· 3





4

Some of random walks

We will consider some of random walks (in one and/or two dimensions) with multiple applications

- 1) A simple random walk (all directions are equal)
- 2) A persistent random walk (probability depends on the previous step)
- 3) A self-avoiding random walk (the same site cannot be occupied twice)
- 4) A restricted random walk (walls or traps)
- 5) Correlated random walks (a connection between walkers)

1) A Simple random walk

A simple random walk is a sequence of unit steps where each step is taken in the direction of one of the coordinate axis, and each possible direction has equal probability of being chosen.

In one dimension 1D random walk there are two possible directions (left and right)

In two dimensions 2D there are four possible directions, e.g., a single step starting at the point with integer coordinates (x, y) would be equally likely to move to any of one of the four neighbors (x + 1, y), (x - 1, y), (x, y + 1) or (x, y - 1).



7

1D Random simple walk and diffusion

The one-dimensional diffusion equation can be written as

$$\frac{\partial p(x,t)}{\partial t} = D \frac{\partial^2 p(x,t)}{\partial x^2}$$

where *D* is the self-diffusion coefficient, and p(x, t)dx is the probability of a particle being in the interval between *x* and x + dx at time *t*.

The solution gives

 $\langle x^2(t) \rangle = 2Dt$

We can see that the random walk method gives the same time dependence.

While the diffusion equation can be solved numerically (e.g. Crank-Nicholson method), it can be very challenging to treat complicated boundary conditions.

Formulating the diffusion problem as a random walk is straightforward to incorporate various boundary conditions.

9



1D Random simple walk

A particle (the walker) starts at the origin (x = 0), and then steps (same length) are chosen randomly left or right with the same probability.

After N steps a position can be recorded as a function of N.

Evaluating the average distance form the starting after many trials would give (the result can easily be derived using that each step is random and it is independent from a previous step)

 $\langle x \rangle \approx 0$ and $\langle x^2 \rangle \sim N$.

In many physical processes (such as the motion of a molecule in solution), the time between steps is approximately a constant, so that number of steps is roughly a proportional to time, then we can write

 $< x^2 > \sim Dt$,

where the factor D is the diffusion constant.

8

2D Random walks

 Type1: Simple random walk on a lattice: Four directions are possible left, right, up and down with equal probability 1/4.

Same step-size (one random number is needed)

 b) Type 2: Random directions but fixed step-size Choose a random angle θ in [0, 2π], and set x = h cos θ, y = h sin θ where h is a fixed step size and θ is a variable angle (one random number is needed θ_i



c) Type 3: Random x_i and random y_i : Random step-size $\sqrt{x_i^2 + y_i^2}$ and random direction (two random numbers are needed)

10

Average distance traveled

The means square distance traveled from the starting point after *N* steps (averaged over K trials)

$$< R^{2}(N) > = \frac{1}{K} \sum_{k=1}^{K} R_{k}^{2}(N)$$

Root-mean-square distance for a constant step size

$$R_{rms} = \sqrt{R^2(N)} \approx \sqrt{N}$$

Root-mean-square distance for a variable step size $r_i^2 = x_i^2 + y_i^2$

$$R_{rms} \approx \sqrt{Nr_{rms}}$$

where $r_{rms} = \sqrt{\langle r^2 \rangle}$ is the root-mean-square step size

where



N	R_rms	R_rms/sqrt(N)	
2	1.189395	0.841029	
4	1.731610	0.865805	
8	2.501420	0.884386	
16	3.542654	0.885663	
32	4.955683	0.876049	
54	7.098192	0.887274	-
28	10.003223	0.884168	$R_{rms} \approx \sqrt{N}r_{rms}$
56	14.184277	0.886517	
L2	20.260120	0.895379	
24	28.338716	0.885585	
18	40.154944	0.887307	
96	56.679889	0.885623	

15



14



16



2) Persistent random walk In a persistent random walk, the transition probability depends on the previous step.

One of the earliest applications of a persistent random walk what to the study of diffusion in chromatographic column.

Example for a walk on a lattice: A persistent random walk in 2 dimensions in a city with $n \times n$ blocks. Condition: the walker can not step back Goal: find average number of steps to get out the city. Is it different from a simple random walk?

3) Self avoiding random walk

Example: using random walk for studying protein growth

Note: A protein is a large biological molecule made up of molecular chains (the residues of amino acids). These chains are formed from monomers, that is, molecules that bind chemically with other molecules.

Random walk is perfectly suited for modelling protein grows.

However, the walk is restricted such that the only positions available after each step are the three neighboring sites (if random walk on a lattice), with the already-occupied sites excluded

This is why this technique is known as a self-avoiding random walk.

Attention: the walk stops when there are no empty neighboring sites available.



20

19



21

4) Restricted random walk

Consider a one-dimensional lattice with traps sites at x = 0 and x = L (L > 0). A walker begins at a site x_0 and takes unit steps to the left and right with equal probability.

When the walker arrives at the trap side, it can no longer move.

Do a Monte Carlo simulation and verify that the mean number of steps τ for the particle to be trapped is given by

 $\tau = (2D)^{-1} x_0 (L - x_0)$

where D is the self-diffusion coefficient in the absence of the traps, and the average is over all possible walks.

The problem is relevant to condense-matter physics (energy transport in solids)



Protein chains consist of (H) and (P) monomers. The actual structure of a protein results from a folding process in which random coils of chains rearrange themselves into a configuration of minimum energy.

Simulation: At each step, you randomly choose an H or a P monomer and drop it on the lattice, with your choice weighted such that H monomers are more likely than P ones.

The goal of the simulation is to find the lowest energy state of an HP sequence of various lengths.

The energy of a chain is defined as

 $E = -\epsilon k$

where ϵ is a positive constant and k is the number of H–H neighbor *not* connected directly (P–P and H–P bonds do not count at lowering the energy).

22

4) Restricted random walk (more)

Suppose that the trap sites are distributed it random on one dimensional lattice with density ρ . For example, if $\rho = 0.01$, the probability that a site is a trap site is 1%.

This site is a trap site if $r < \rho$ where, as usual, r is uniformly distributed in the interval $0 \le r \le 1$.

If a walker is placed at random at any non-trapping site, determine its mean survival time τ , that is, the mean number of steps before a trap site is reached.

Of the major complication is that it is necessary to perform three averages: the distribution of traps, the origin of the walker, and the different walks for a given trap distribution and origin.



5) Synchronized random walk

Randomly please two walkers on a one-dimensional lattice of L site, so that both walkers are not at the same site.

It each time step randomly choose whether the walkers move to the left or to the right. Both walkers move in the same direction.

If a walker cannot move into choosing direction because it is at the boundary, then this walker remains at the same side for this time step.

The trail ends when both walkers are the same site. Find the mean time for two walkers to reach the same side.

This model is relevant to a method of doing cryptography using neural networks.

25

Areas

Stochastic optimization
 or the problem of local minima.

Swarm intelligence
 or the ant colony optimization

- Genetic algorithms
 Use Darwinian evolution of a gene pool
 to find the fittest genes
- Simulated annealing
- and many more ...



27





26

Example of Problems Using Stochastic Search and Optimization

- · Place sensors in manner to maximize useful information
- · Minimize the costs of shipping from production facilities to warehouses
- Maximize the probability of detecting an incoming warhead (vs. decoy) in a missile defense system
- Determine the times to administer a sequence of drugs for maximum therapeutic effect
- · Find the best red-yellow-green signal timings in an urban traffic network
- Determine the best schedule for use of laboratory facilities to serve an organization's overall interests

28

Example 1

Initial (random) configuration.

Find a configuration of 10 particles (interacting by Lenard-Jones potential) that has the lowest energy

Final configuration



30





Part : 4 Problems for a curious student



32



34

1. Buffon's needle

The French naturalist and mathematician Comte de Buffon showed that the probability that a needle of length L thrown randomly onto a grid of parallel lines with distance D > L apart intersects a line is $2L/(D * \pi)$

```
A part of a code ...
```

```
c*** loop over trials
    hit = 0
    do it=1,itests
    x0 = float(N)*D*rand()
    k = int(x0/D)
    x1 = x0 - D*float(k)
    x2 = D - x1
    x = min(x1,x2)
    dx = 0.5*abs(L*cos(1.0*pi*rand()))
    if(dx.ge.x) hit = hit + 1
    end do
c*** average number of hits
    ahit = float(hit)/float(itests)
    buffon = (2*L)/(pi*D)
```

2. Conditional probability

Suppose that many people in the community tested at random for Covid. The accuracy of the test is 87%, and the incidence of the disease in the general population, independent of any test, is 1%.

A person tested positive for Covid, what is the probability that this person really has Covid?

Comment: the answer is much less than 87%.

37

<pre>write (*,*)'enter numbers of tests, money and goal' read (*,*) itests, money1, money2 c*** loop over trials total = 0 wins = 0 </pre>
d0 1T=1,1Tests x=monev1
games=0
<pre>do while(x.gt.0.and.x.lt.money2) games = games + 1</pre>
luck = 1 if(rand() le 0 5) luck=-1
x = x+luck
<pre>total = total+games if(x.gt.0) wins = wins+1 end do</pre>
<pre>c*** average number of games and wins agames = float(total)/float(itests) awins = float(wins)/float(itests) aloose = 1.0-awins write (*,100) itests, money1, money2 write (*,101) awins, aloose, azames</pre>

39

If a chance to win in each bet 49/51 The gambler's ruin problem. Chances to reach certain goal enter numbers of tests, money and goal 10000 10

tests: 100000 initial: 10 goal: 100 win = 9.44000E-03 loose = 9.90560E-01 games = 4.51806E+02

chance to win is about 0.9%

3. The gambler's ruin problem.

Suppose that a person decides to try to increase the amount of money in his/her pocket by participating in some gambling. Initially, the gambler begins with \$m in capital. The gambler decides that he/she will gamble until a certain goal, \$n (n>m), is achieved or there is no money left (credit is not allowed). On each throw of a coin (roll of the dice, etc.) the gambler either win \$1 or lose \$1. If the gambler achieves the goal, he/she will stop playing. If the gambler ends up with no money he/she is ruined.

What are chances for the gambler to achieve the goal as a function of $k, \ensuremath{\mathsf{where}}\xspace\, k=n/m?$

How long on average will it take to play to achieve the goal or to be ruined?

38

If a chance to win in each bet 50/50

The gambler's ruin problem. Chances to reach certain goal enter numbers of tests, money and goal 10000 10 100 tests: 10000

initial: 10 goal: 100 win = 1.026E-01 loose = 8.974E-01 games = 9.019E+02

chance to win is about 10%

40

4. Cooking burgers

An industrious physics major finds a job at a local fast food restaurant to help him pay his way through college. His task is to cook 20 hamburgers on a grill at any one time. When a hamburger is cooked, he is supposed to replace it with uncooked hamburger. However, our physics major does not pay attention to whether the hamburger is cooked or not. His method is to choose a hamburger at random and replace it by an uncooked one. He does not check if the hamburger that he removes from the grill is ready.

What is the distribution of cooking times of the hamburgers that he removes? What is a chance for a customer to get a well cooked hamburger if it takes 5 minutes to cook a hamburger.

Does the answers to the first two questions change if he cooks 40 hamburgers at any one time?

Comment: For simplicity, assume that he replaces a hamburger at a regular interval of 30 seconds and there is an indefinite supply of uncooked hamburgers.

41





45







60

50

40

0 L

30 subscription su

44

Investigate a simple problem that generated much attention several years ago and for which many mathematicians obtained an incorrect solution. The problem was the analysis of the optimal strategy in a television game show popular at the time. The show was Let's Make a Deal with host Monty Hall. At some point in the show, a contestant was given a choice of selecting one of three possible items, each concealed behind one of three closed doors. The items varied considerably in value. After the contestant made a choice but before the chosen door was opened, the host, who knew where the most valuable item was, would open one of the doors not selected and reveal a worthless item. The host would then offer to let the contestant select a different door from what was originally selected. The question, of course, is should the contestant switch? A popular magazine writer Marilyn vos Savant concluded that the optimal strategy is to switch. This strategy is counterintuitive to many mathematicians, who would say that there is nothing to be gained by switching; that is, that the probability of improving the selection is 0.5. Study this problem by Monte Carlo methods. What is the probability of improving the selection by switching? Be careful to understand all of the assumptions, and then work the problem analytically also. (A Monte Carlo study is no substitute for analytic study.)

grill for 20 burgers

૿ૻૡ૾ૻ૱૿ૡ૾ૺ૱

10 20 30 40 50 60 70 80 90 100 cooking time (min)

burgers cooked = 1,000

പ്പംഷം പ്

