Random Walks

October 16, 2014

Convergence of Binomial Probabilities¹

We all intuitively understand that when a fair coin (50% heads, 50% tails) is flipped many times. the probability of heads versus tails should start to converge. For example, let's say we get 1 point for every head and o points for every tail. What will the average number of points we get at any point in time? Let's compute and plot this in R with n=5000 coin flips!

```
<sup>1</sup> These examples are based on:
http://nvenkataraman1.github.
io/random-walks/
```

```
n <- 5000
flips <- rbinom(n, 1, 0.5) # a fair coin follows a simple binomial distribution
means < - c(0)
for (i in 1:n) {
    means[i] <- mean(flips[1:i])</pre>
}
plot(means, type = "l", ylim = c(0, 1))
abline(h = 0.5, col = "red")
```

We see that the average number of points quickly converges to 0.5 (a half), just as we suspected. This is our common intuition about how probabilities average out over time.

0.8 9.0 9.0 0.2 0.0 1000 2000 3000 4000 5000

Figure 1: Coin flipping probabilities

0.

Now, instead of average probabilities, let's make a game where we gain 1 point for heads and lose one point for tails. How many total points would we make over time? Will it converge?

```
n <- 5000
flip <-2 * rbinom(n, 1, 0.5) - 1
total <- c(0)
for (i in 2:n) {
    total[i] <- total[i - 1] + flip[i]
}
plot(total, type = "l", ylim = c(-300, 300))
abline(h = 0)
```

When we plot out our total number of points, rather than converging, it seems to just randomly walk away! Random walks are one way of modeling many everyday phenomena that are affected by so many variables that they appear random in their progress: stock markets, weather fluctuations, ant foraging, and more².

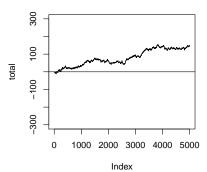


Figure 2: A random (binomial) walk ² http://en.wikipedia.org/wiki/ Random_walk

But how far from their starting points do random walks go on average? It isn't easy to tell, so let's simulate random walks and find out!

Random binomial walk function

```
random_binomial_walk <- function(steps) {</pre>
    flip <-2 * rbinom(n, 1, 0.5) - 1
    total <- c(0)
    for (i in 2:steps) {
        total[i] \leftarrow total[i - 1] + flip[i]
    }
    return(total)
}
# Simulate six random walks of 5000 steps each
walks <- data.frame(matrix(0, nrow = 5000, ncol = 6))</pre>
for (i in 1:6) {
    walks[i] <- random_binomial_walk(n)</pre>
}
# Plot out our six random walks
par(mfrow = c(2, 3)) # setup plotting to 2rows x 3 columns
for (i in 1:6) {
    plot(walks[[i]], type = "l", ylim = c(-200,
    abline(h = 0, col = "red")
    abline(h = 0, col = "red")
}
```

Even with a small number of random walks, we see they can go anywhere. It will be hard to plot a large number of walks. So let's simulate a large number of walks and see how their final distance from zero is distributed.

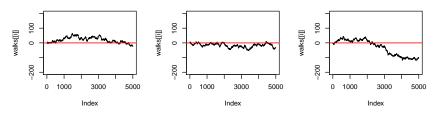
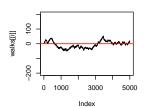
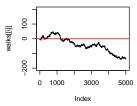
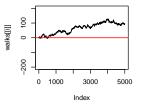


Figure 3: Six Binomial Walks







```
par(mfrow = c(1, 1)) # return graphics settings to 1 x 1
```

Even with a small number of random walks, we see they can go anywhere. It will be hard to plot a large number of walks. So let's simulate a large number of walks and see how their final distance from zero is distributed.

```
m < -3000
n <- 2000
walks <- data.frame(matrix(0, nrow = n, ncol = m))</pre>
for (i in 1:m) {
    walks[i] = random_binomial_walk(n)
}
distance <- c(0)
for (i in 1:m) {
    distance[i] <- walks[n, i]</pre>
}
hist(distance, breaks = 20, prob = TRUE)
lines(density(distance), lwd = 3, col = "blue")
```

Histogram of distance

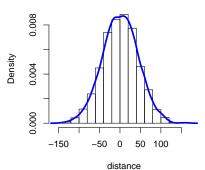


Figure 4: Distribution of binomial walks

Histogram of distance

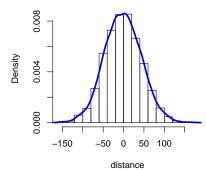


Figure 5: Distribution of gaussian walks