Distribution of Random Walks

October 17, 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.

1000 2000 4000 5000 3000 Figure 1: Coin flipping probabilities

0. 0.8 9.0

0.4

0.2

0.0

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, brownian motion, and more².

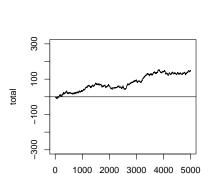


Figure 2: A random (binomial) walk ² http://en.wikipedia.org/wiki/ Random_walk

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] <- total[i - 1] + flip[i]</pre>
    }
    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(5000)</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")
}
```

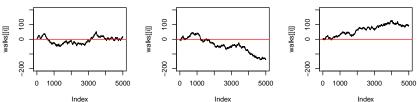
We first create a 'function' that creates binomial walks for us. It reuses the code we saw earlier. This function takes the number of 'steps' to walk as a parameter.

We first create a 'data.frame' that is the size of a 2x3 matrix. The we use a loop to run our random binomial walk function six times. The result of our six function calls are stored in the data.frame

We then plot the six walks. The 'mfrow' function lets us put plots sideby-side in a grid pattern! At the end, we have to be careful to call mfrow again to return the graphics setting to plot 1x1 figures at a time.

Figure 3: Six Random Binomial Walks

```
walks[[i]]
                                             valks[[i]]
-200
                                                  200
                                                                                                    200
     0 1000
                     3000
                                  5000
                                                        0 1000
                                                                        3000
                                                                                    5000
                                                                                                          0 1000
                                                                                                                           3000
                                                                                                                                       5000
                   Index
```



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 what distance from zero points each walk ended up. We will use m=3000 walks, each having n=2000 steps.

```
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)
}
final.points <- c(0)
for (i in 1:m) {
    final.points[i] <- walks[n, i]</pre>
}
```

We can now plot the distribution of how many points away from zero (positive or negative) each each of the 3000 walks finally got to, after 2000 steps.

```
hist(final.points, breaks = 20, prob = TRUE)
lines(density(final.points), lwd = 3, col = "blue")
```

Does this bell-shaped symmetric distribution look familiar? It is our old friend the normal distribution again!

Histogram of final.points

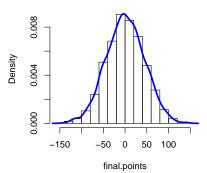


Figure 4: Distribution of binomial walks