Probability

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The Hot Hand

Basketball players who make several baskets in succession are described as having a *hot hand*. Fans and players have long believed in the hot hand phenomenon, which refutes the assumption that each shot is independent of the next. However, a 1985 paper by Gilovich, Vallone, and Tversky collected evidence that contradicted this belief and showed that successive shots are independent events. This paper started a great controversy that continues to this day, as you can see by Googling *hot hand basketball*.

We do not expect to resolve this controversy today. However, in this lab we'll apply one approach to answering questions like this. The goals for this lab are to (1) think about the effects of independent and dependent events, (2) learn how to simulate shooting streaks in R, and (3) to compare a simulation to actual data in order to determine if the hot hand phenomenon appears to be real.

Getting Started

Load packages

In this lab, we will explore and visualize the data using the tidyverse suite of packages. The data can be found in the companion package for OpenIntro labs, openintro.

Let's load the packages.

```
library(tidyverse)
library(openintro)
```

Data

Your investigation will focus on the performance of one player: Kobe Bryant of the Los Angeles Lakers. His performance against the Orlando Magic in the 2009 NBA Finals earned him the title *Most Valuable Player* and many spectators commented on how he appeared to show a hot hand. The data file we'll use is called kobe_basket.

glimpse(kobe_basket)

This data frame contains 133 observations and 6 variables, where every row records a shot taken by Kobe Bryant. The **shot** variable in this dataset indicates whether the shot was a hit (H) or a miss (M).

Just looking at the string of hits and misses, it can be difficult to gauge whether or not it seems like Kobe was shooting with a hot hand. One way we can approach this is by considering the belief that hot hand shooters tend to go on shooting streaks. For this lab, we define the length of a shooting streak to be the number of consecutive baskets made until a miss occurs.

For example, in Game 1 Kobe had the following sequence of hits and misses from his nine shot attempts in the first quarter:

$$HM \mid M \mid HHM \mid M \mid M \mid M$$

You can verify this by viewing the first 9 rows of the data in the data viewer.

Within the nine shot attempts, there are six streaks, which are separated by a "|" above. Their lengths are one, zero, two, zero, zero (in order of occurrence).

1. What does a streak length of 1 mean, i.e. how many hits and misses are in a streak of 1? What about a streak length of 0?

Insert your answer here

Using the example above with the first 9 shots, a streak of 1 means that Kobe (RIP) has made one shot and missed another. So there are two shots within a streak of 1.

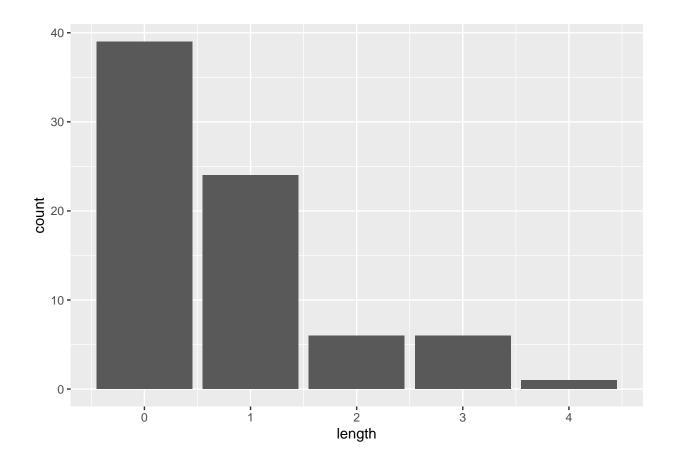
End of Answer

Counting streak lengths manually for all 133 shots would get tedious, so we'll use the custom function calc_streak to calculate them, and store the results in a data frame called kobe_streak as the length variable.

```
kobe_streak <- calc_streak(kobe_basket$shot)
```

We can then take a look at the distribution of these streak lengths.

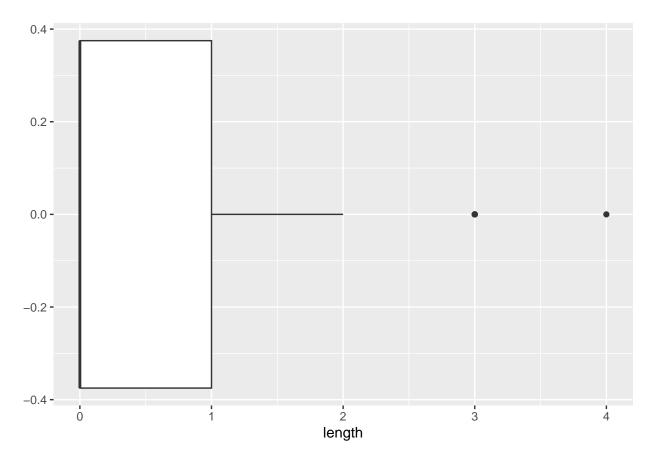
```
ggplot(data = kobe_streak, aes(x = length)) +
geom_bar()
```



2. Describe the distribution of Kobe's streak lengths from the 2009 NBA finals. What was his typical streak length? How long was his longest streak of baskets? Make sure to include the accompanying plot in your answer.

Insert your answer here

```
ggplot(data = kobe_streak, aes(x = length)) +
geom_boxplot()
```



According to the box and whisker plot above, we can see that the median streak is of 0. Meaning that half of his streaks were missing two shots in a row and his longest streak of baskets were 4.

End of Answer

Compared to What?

We've shown that Kobe had some long shooting streaks, but are they long enough to support the belief that he had a hot hand? What can we compare them to?

To answer these questions, let's return to the idea of *independence*. Two processes are independent if the outcome of one process doesn't effect the outcome of the second. If each shot that a player takes is an independent process, having made or missed your first shot will not affect the probability that you will make or miss your second shot.

A shooter with a hot hand will have shots that are *not* independent of one another. Specifically, if the shooter makes his first shot, the hot hand model says he will have a *higher* probability of making his second shot.

Let's suppose for a moment that the hot hand model is valid for Kobe. During his career, the percentage of time Kobe makes a basket (i.e. his shooting percentage) is about 45%, or in probability notation,

$$P(\text{shot } 1 = \text{H}) = 0.45$$

If he makes the first shot and has a hot hand (*not* independent shots), then the probability that he makes his second shot would go up to, let's say, 60%,

$$P(\text{shot } 2 = \text{H} | \text{shot } 1 = \text{H}) = 0.60$$

As a result of these increased probabilities, you'd expect Kobe to have longer streaks. Compare this to the skeptical perspective where Kobe does *not* have a hot hand, where each shot is independent of the next. If he hit his first shot, the probability that he makes the second is still 0.45.

$$P(\text{shot } 2 = H | \text{shot } 1 = H) = 0.45$$

In other words, making the first shot did nothing to effect the probability that he'd make his second shot. If Kobe's shots are independent, then he'd have the same probability of hitting every shot regardless of his past shots: 45%.

Now that we've phrased the situation in terms of independent shots, let's return to the question: how do we tell if Kobe's shooting streaks are long enough to indicate that he has a hot hand? We can compare his streak lengths to someone without a hot hand: an independent shooter.

Simulations in R

While we don't have any data from a shooter we know to have independent shots, that sort of data is very easy to simulate in R. In a simulation, you set the ground rules of a random process and then the computer uses random numbers to generate an outcome that adheres to those rules. As a simple example, you can simulate flipping a fair coin with the following.

```
coin_outcomes <- c("heads", "tails")
sample(coin_outcomes, size = 1, replace = TRUE)</pre>
```

```
## [1] "tails"
```

The vector coin_outcomes can be thought of as a hat with two slips of paper in it: one slip says heads and the other says tails. The function sample draws one slip from the hat and tells us if it was a head or a tail.

Run the second command listed above several times. Just like when flipping a coin, sometimes you'll get a heads, sometimes you'll get a tails, but in the long run, you'd expect to get roughly equal numbers of each.

If you wanted to simulate flipping a fair coin 100 times, you could either run the function 100 times or, more simply, adjust the size argument, which governs how many samples to draw (the replace = TRUE argument indicates we put the slip of paper back in the hat before drawing again). Save the resulting vector of heads and tails in a new object called sim_fair_coin.

```
sim_fair_coin <- sample(coin_outcomes, size = 100, replace = TRUE)</pre>
```

To view the results of this simulation, type the name of the object and then use table to count up the number of heads and tails.

```
sim_fair_coin
```

```
## [1] "heads" "tails" "heads" "tails" "tails" "tails" "tails" "heads"
## [10] "tails" "heads" "tails" "heads" "tails" "heads" "tails" "heads"
## [19] "tails" "tails" "tails" "heads" "heads" "tails" "tails" "heads"
## [28] "tails" "tails" "tails" "heads" "heads" "tails" "heads" "tails" "heads"
## [37] "heads" "tails" "tails" "heads" "heads" "tails" "tails" "heads"
```

```
## [46] "tails" "tails" "tails" "heads" "heads" "heads" "tails" "tails" "heads"
## [55] "tails" "tails" "heads" "tails" "heads" "tails" "heads" "tails" "heads"
## [64] "heads" "tails" "heads" "tails" "heads" "heads" "tails" "tails"
## [73] "heads" "heads" "tails" "tails" "tails" "heads" "heads" "tails" "tails"
## [82] "heads" "tails" "heads" "heads" "tails" "heads" "tails" "heads" "tails"
## [91] "tails" "heads" "tails" "heads" "heads" "heads" "tails"
## [100] "heads"
```

```
table(sim_fair_coin)
```

```
## sim_fair_coin
## heads tails
## 45 55
```

Since there are only two elements in coin_outcomes, the probability that we "flip" a coin and it lands heads is 0.5. Say we're trying to simulate an unfair coin that we know only lands heads 20% of the time. We can adjust for this by adding an argument called prob, which provides a vector of two probability weights.

prob=c(0.2, 0.8) indicates that for the two elements in the outcomes vector, we want to select the first one, heads, with probability 0.2 and the second one, tails with probability 0.8. Another way of thinking about this is to think of the outcome space as a bag of 10 chips, where 2 chips are labeled "head" and 8 chips "tail". Therefore at each draw, the probability of drawing a chip that says "head" is 20%, and "tail" is 80%.

3. In your simulation of flipping the unfair coin 100 times, how many flips came up heads? Include the code for sampling the unfair coin in your response. Since the markdown file will run the code, and generate a new sample each time you *Knit* it, you should also "set a seed" **before** you sample. Read more about setting a seed below.

Insert your answer here

```
## sim_8020_coin
## heads tails
## 21 79
```

Setting the seed to a fixed value should keep the result of successive simulations the same. With this seed, there were 21 heads and 79 tails, which is very close to the 20% to 80% split.

End of Answer

A note on setting a seed: Setting a seed will cause R to select the same sample each time you knit your document. This will make sure your results don't change each time you knit, and it will also ensure reproducibility of your work (by setting the same seed it will be possible to reproduce your results). You can set a seed like this:

```
set.seed(35797) # make sure to change the seed
```

The number above is completely arbitraty. If you need inspiration, you can use your ID, birthday, or just a random string of numbers. The important thing is that you use each seed only once in a document. Remember to do this **before** you sample in the exercise above.

In a sense, we've shrunken the size of the slip of paper that says "heads", making it less likely to be drawn, and we've increased the size of the slip of paper saying "tails", making it more likely to be drawn. When you simulated the fair coin, both slips of paper were the same size. This happens by default if you don't provide a prob argument; all elements in the outcomes vector have an equal probability of being drawn.

If you want to learn more about sample or any other function, recall that you can always check out its help file

```
?sample
```

Simulating the Independent Shooter

Simulating a basketball player who has independent shots uses the same mechanism that you used to simulate a coin flip. To simulate a single shot from an independent shooter with a shooting percentage of 50% you can type

```
shot_outcomes <- c("H", "M")
sim_basket <- sample(shot_outcomes, size = 1, replace = TRUE)</pre>
```

To make a valid comparison between Kobe and your simulated independent shooter, you need to align both their shooting percentage and the number of attempted shots.

4. What change needs to be made to the sample function so that it reflects a shooting percentage of 45%? Make this adjustment, then run a simulation to sample 133 shots. Assign the output of this simulation to a new object called sim_basket.

Insert your answer here

In order to simulate an independent shooter, the sample function will need to be updated to change the size to 133 and the probabilities to reflect a 55% miss chance and a 45% hit chance.

I would also like to state that we switched the seed in the cell above to 35797 as well.

Reusing the shot_outcomes variable from before:

```
## sim_basket
## H M
## 62 71
```

From this simulation, we can see that there are 62 hits and 71 misses.

End of Answer

Note that we've named the new vector sim_basket, the same name that we gave to the previous vector reflecting a shooting percentage of 50%. In this situation, R overwrites the old object with the new one, so always make sure that you don't need the information in an old vector before reassigning its name.

With the results of the simulation saved as sim_basket, you have the data necessary to compare Kobe to our independent shooter.

Both data sets represent the results of 133 shot attempts, each with the same shooting percentage of 45%. We know that our simulated data is from a shooter that has independent shots. That is, we know the simulated shooter does not have a hot hand.

More Practice

Comparing Kobe Bryant to the Independent Shooter

5. Using calc_streak, compute the streak lengths of sim_basket, and save the results in a data frame called sim_streak.

Insert your answer here

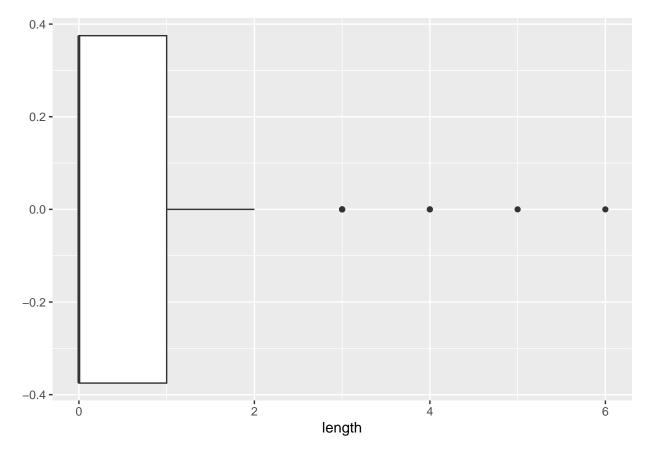
```
sim_streak <- calc_streak(sim_basket)</pre>
```

End of Answer

6. Describe the distribution of streak lengths. What is the typical streak length for this simulated independent shooter with a 45% shooting percentage? How long is the player's longest streak of baskets in 133 shots? Make sure to include a plot in your answer.

Insert your answer here

```
ggplot(data = sim_streak, aes(x = length)) +
geom_boxplot()
```



Looking at the box and whisker plot, we can see that the typical streak is still 0 although this independent player's longest streak is 6 shots.

Given that there is a 45% chance of making any individual shot, then we can calculate the chances of having a streak of 6 by this:

$$P(4 \text{ hits in a row}) = (0.45)^4 = 0.0410 = 4.10 P(6 \text{ hits in a row}) = (0.45)^6 = 0.0083 = 0.83$$

With a 0.83% chance, it's a rare occurance for 6 hits in a row but the 4 hits in a row that Kobe had isn't nearly as rare.

End of Answer

7. If you were to run the simulation of the independent shooter a second time, how would you expect its streak distribution to compare to the distribution from the question above? Exactly the same? Somewhat similar? Totally different? Explain your reasoning.

Insert your answer here

I expect it to be somewhat similar as there are a large number of shots being simulated and because of that, there is a relatively low chance that the distribution of data would be vastly different.

End of Answer

8. How does Kobe Bryant's distribution of streak lengths compare to the distribution of streak lengths for the simulated shooter? Using this comparison, do you have evidence that the hot hand model fits Kobe's shooting patterns? Explain.

Insert your answer here

Comparing our simulated shooter's streak length to Kobe Bryant's we see a very similar pattern. Both had a similar distribution of streaks, being that the majority of streaks are 0 and quickly taper off with 2 being the far edge of the whisker. With an IQR from 0 to 1, it is also farily uncommon that there are streaks greater than 1.

This simulation appears to be evidence against the hot hand model as it seems that the streak lengths are more random. If the hot hand model was true, I would expect to see more streaks and longer streaks.

End of Answer			