DATA606 - Intro to Bayesian Analysis

Jason Bryer, Ph.D. December 5, 2018

Meetup Presentations

- · Vishal Arora (7.5)
- · Corey Arnouts (7.35)
- · Simon Ustoyev (8.3) http://rpubs.com/simon63/chapter8_3
- · Charles Meyers

by

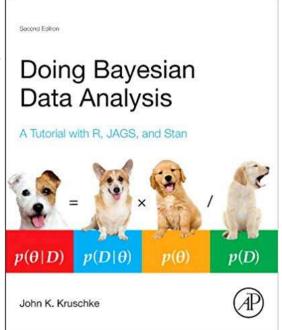
Bayesian Analysis

Kruschke's videos are an excelent introduction to Bayesian Analysis https://www.youtube.com/watch?v=YyohWpjl6KU!

Doing Bayesian Data Analysis, Second Edition: A Tutorial with R, JAGS, and Stan

Sharon Bertsch McGrayne

Video series by Rasmus Baath Part 1, Part 2, Part 3



Bayes Theorem

$$P(A|B) = \frac{P(B|A)P(A)}{P(B|A)P(A) + P(B|A')P(A')}$$

Consider the following data from a cancer test:

- 1% of women have breast cancer (and therefore 99% do not).
- · 80% of mammograms detect breast cancer when it is there (and therefore 20% miss it).
- 9.6% of mammograms detect breast cancer when it's not there (and therefore 90.4% correctly return a negative result).

	Cancer (1%)	No Cancer (99%)
Test postive	80%	9.6%
Test negative	20%	90.4%

How accurate is the test?

Now suppose you get a positive test result. What are the chances you have cancer? 80%? 99%? 1%?

- · Ok, we got a positive result. It means we're somewhere in the top row of our table. Let's not assume anything it could be a true positive or a false positive.
- The chances of a true positive = chance you have cancer * chance test caught it = 1% * 80% = .008
- The chances of a false positive = chance you don't have cancer * chance test caught it anyway = 99% * 9.6% = 0.09504

	Cancer (1%)	No Cancer (99%)
Test postive	True +: 1% * 80%	False +: 99% * 9.6%
Test negative	False -: 1% * 20%	True -: 99% * 90.4%

How accurate is the test?

$$Probability = \frac{desired \quad event}{all \quad possibilities}$$

The chance of getting a real, positive result is .008. The chance of getting any type of positive result is the chance of a true positive plus the chance of a false positive (.008 + 0.09504 = .10304).

So, our chance of cancer is .008/.10304 = 0.0776, or about 7.8%.

Bayes Formula

It all comes down to the chance of a true positive result divided by the chance of any positive result. We can simplify the equation to:

$$P(A|B) = \frac{P(B|A)P(A)}{P(B)}$$

How many fish are in the lake?

- · Catch them all, count them. Not practical (or even possible)!
- · We can sample some fish.

Our strategy:

- 1. Catch some fish.
- 2. Mark them.
- 3. Return the fish to the pond. Let them get mixed up (i.e. wait a while).
- 4. Catch some more fish.
- 5. Count how many are marked.

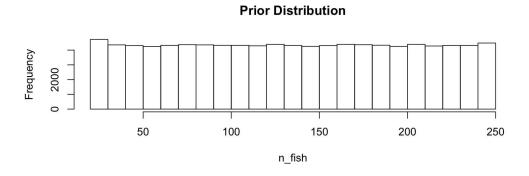
For example, we initially caught 20 fish, marked them, returned them to the pond. We then caught another 20 fish and 5 of them were marked (i.e they were caught the first time).

 $Adopted from Rasmath \ B\aa \$th use R! \ 2015 \ workshop: \ http://www.sumsar.net/files/academia/user_2015_tutorial_bayesian_data_analysis_short_version.pdf$

Strategy for fitting a model

Step 1: Define Prior Distribution. Draw a lot of random samples from the "prior" probability distribution on the parameters.

```
n_draw <- 100000
n_fish <- sample(20:250, n_draw, replace = TRUE)
head(n_fish, n=10)
## [1] 209 130 107 81 81 166 81 128 186 98
hist(n_fish, main="Prior Distribution")</pre>
```



Strategy for fitting a model

Step 2: Plug in each draw into the generative model which generates "fake" data.

```
pick_fish <- function(n_fish) { # The generative model
    fish <- rep(0:1, c(n_fish - 20, 20))
    sum(sample(fish, 20))
}
n_marked <- rep(NA, n_draw)
for(i in 1:n_draw) {
    n_marked[i] <- pick_fish(n_fish[i])
}
head(n_marked, n=10)

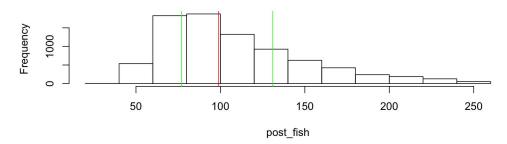
## [1] 1 3 3 2 2 2 4 2 2 7</pre>
```

Strategy for fitting a model

Step 3: Keep only those parameter values that generated the data that was actually observed (in this case, 5).

```
post_fish <- n_fish[n_marked == 5]
hist(post_fish, main='Posterior Distribution')
abline(v=median(post_fish), col='red')
abline(v=quantile(post_fish, probs=c(.25, .75)), col='green')</pre>
```

Posterior Distribution

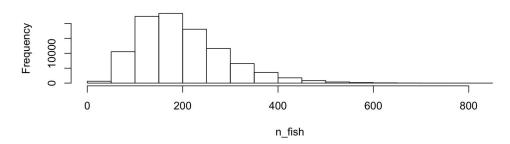


What if we have better prior information?

An "expert" believes there are around 200 fish in the pond. Insteand of a uniform distribution, we can use a binomial distribution to define our "prior" distribution.

```
n_fish <- rnbinom(n_draw, mu = 200 - 20, size = 4) + 20
hist(n_fish, main='Prior Distribution')</pre>
```

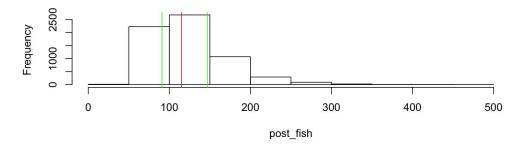




What if we have better prior information?

```
n_marked <- rep(NA, n_draw)
for(i in 1:n_draw) {
    n_marked[i] <- pick_fish(n_fish[i])
}
post_fish <- n_fish[n_marked == 5]
hist(post_fish, main='Posterior Distribution')
abline(v=median(post_fish), col='red')
abline(v=quantile(post_fish, probs=c(.25, .75)), col='green')</pre>
```

Posterior Distribution



Bayes Billiards Balls

Consider a pool table of length one. An 8-ball is thrown such that the likelihood of its stopping point is uniform across the entire table (i.e. the table is perfectly level). The location of the 8-ball is recorded, but not known to the observer. Subsequent balls are thrown one at a time and all that is reported is whether the ball stopped to the left or right of the 8-ball. Given only this information, what is the position of the 8-ball? How does the estimate change as more balls are thrown and recorded?

shiny_demo('BayesBilliards', package='DATA606')

See also: http://www.bryer.org/post/2016-02-21-bayes_billiards_shiny/