## **Bayesian Methods**

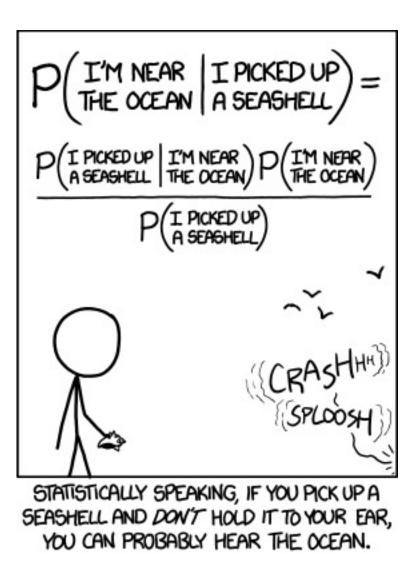
#### Quantitative Understanding in Biology

Thursday, 30 September 2021

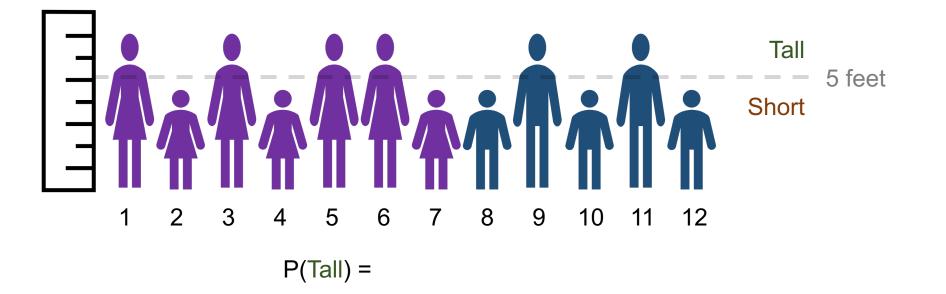
Lecture Notes by Jason Banfelder

Slide Compilation and Demonstratives by Noemi Linden

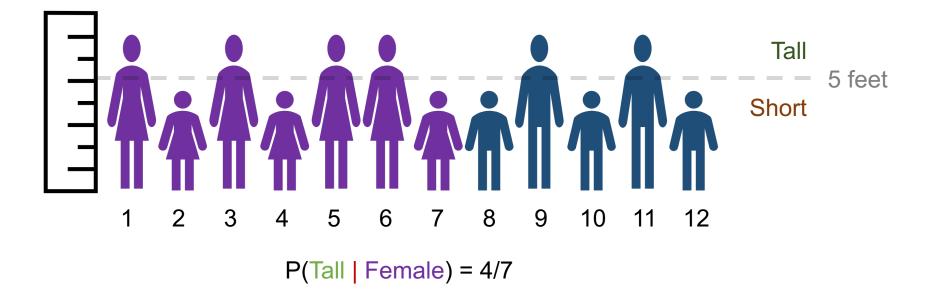
Based on slides from Ariana Clerkin



### Probability 7<sup>th</sup> grade classroom



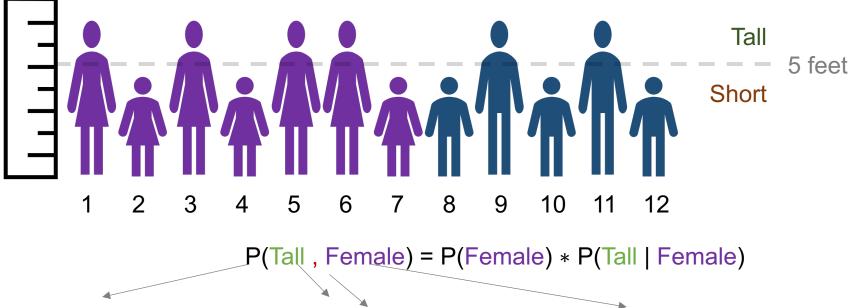
#### Conditional Probability 7<sup>th</sup> Grade Classroom



*Probability* that the student is tall *given that* the student is *female* (Conditional Probability)

We expect P(Tall | Female) > P(Tall) without taking any measurements of this particular class.

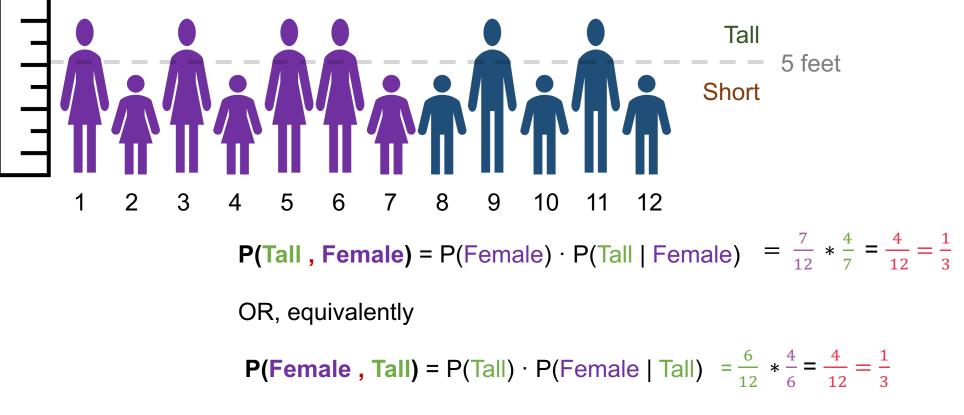
### Joint Probability 7<sup>th</sup> Grade Classroom



Probability that the student is tall and that the student is female (Joint Probability)

$$\frac{7}{12} * \frac{4}{7} = \frac{4}{12} = \frac{1}{3}$$

### Joint Probability 7<sup>th</sup> Grade Classroom



P(Tall, Female) = P(Female, Tall)

## **Deriving Bayes' Rule**

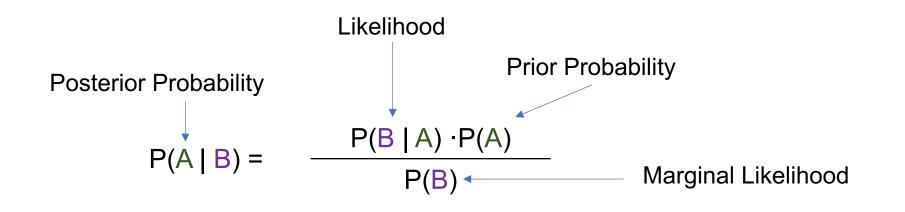
We have shown that:

 $P(Tall , Female) = P(Female) \cdot P(Tall | Female)$  $P(Tall , Female) = P(Tall) \cdot P(Female | Tall)$ 

Therefore:

 $P(Female) \cdot P(Tall | Female) = P(Tall) \cdot P(Female | Tall)$   $P(Tall | Female) = \frac{P(Female | Tall) \cdot P(Tall)}{P(Female)}$ Or generally, for generic events A & B, we have  $P(A | B) = \frac{P(B | A) \cdot P(A)}{P(B)}$ 

### Bayes' Rule: Terminology



#### Information:

- 1% of women in a given population have breast cancer
- If a woman has breast cancer, there is a 90% chance that a particular diagnostic test will return a positive result (10% false negative rate)
- If a woman does not have breast cancer, there is a 10% chance that this diagnostic test will return a positive result (10% false positive rate).

#### Question:

What is the probability that a woman with a positive test result actually has cancer?

## Multiple Choice:

Which notation shows the probability that a woman with a positive test result actually has cancer?

a.) P(Cancer | Positive Test)

b.) P(Cancer, Positive Test)

c.) P(Positive Test | Cancer)

d.) P(Positive Test  $\cap$  Cancer)

- 1% of women in a given population have breast cancer
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$$P(Cancer | Positive) = \frac{P(Positive | Cancer) \cdot P(Cancer)}{P(Positive)}$$

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```
P(Positive) = P(True Positive) + P(False Positive)
P(Positive) = P(Positive | Cancer) \cdot P(Cancer) + P(+ | Healthy) \cdot P(Healthy)
P(Positive) = 0.9 \cdot 0.01 + 0.1 \cdot (1 - 0.01)
P(Positive) = 0.108
```

### Now we can complete Bayes' Rule

 $P(Cancer | Positive) = \frac{P(Positive | Cancer) \cdot P(Cancer)}{P(Positive)}$  $P(Cancer | Positive) = \frac{0.9 \cdot 0.01}{0.108} = 0.083$ 

How can we apply Bayes' rule to estimating model parameters?

## Frequentist Coin Flip: 20 Flips; 13 Heads

Objective: Estimate the Coin's Bias with a 95% Confidence Interval

```
binom.test(13, 20)
##
## Exact binomial test
##
## data: 13 and 20
## number of successes = 13, number of trials = 20, p-value =
## 0.2632
## alternative hypothesis: true probability of success is not equal to 0.5
## 95 percent confidence interval:
## 0.4078115 0.8460908
## sample estimates:
## probability of success
## 0.65
```

#### Conclusion:

- Bias = 0.65
- 95% CI = (0.41, 0.85)

## We know how to compute the probability of any particular data outcome

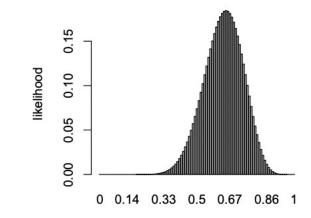
```
dbinom(13, size = 20, prob = 0.5)
```

## [1] 0.07392883

dbinom(13, size = 20, prob = 0.25)
## [1] 0.0001541923

# Computing the probability of getting the data that we observed at various values of the coin's bias

coin.bias <- seq(from = 0, to = 1, by = 0.01)
likelihood <- dbinom(13, 20, prob = coin.bias)
barplot(likelihood, names.arg = coin.bias, ylab = "likelihood")</pre>



## Imagine we have a pool of 101 coins each with a different bias (0.00, 0.001, 0.002,...)

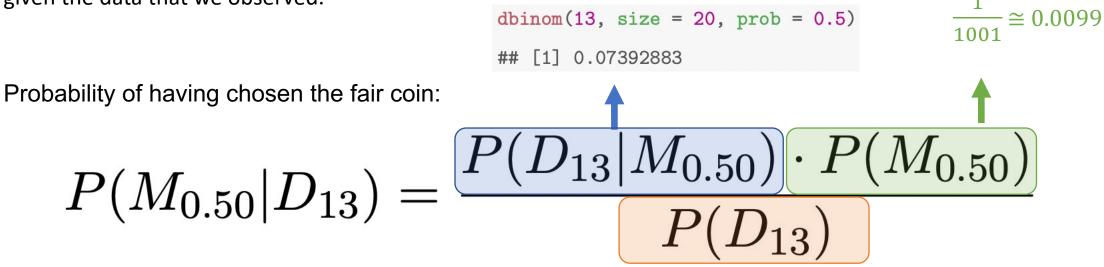
We can calculate the probability of each of the 101 coins being the one that we chose, given the data that we observed.

Probability of having chosen the fair coin:

$$P(M_{0.50}|D_{13}) = \frac{P(D_{13}|M_{0.50}) \cdot P(M_{0.50})}{P(D_{13})}$$

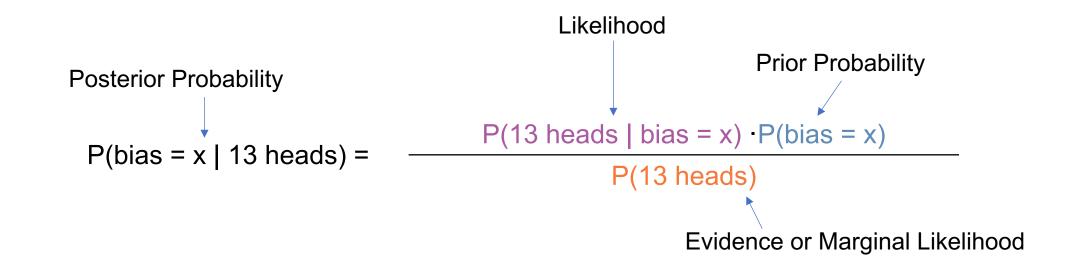
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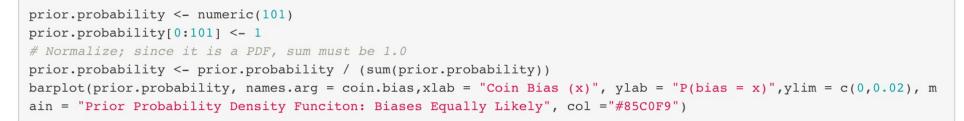


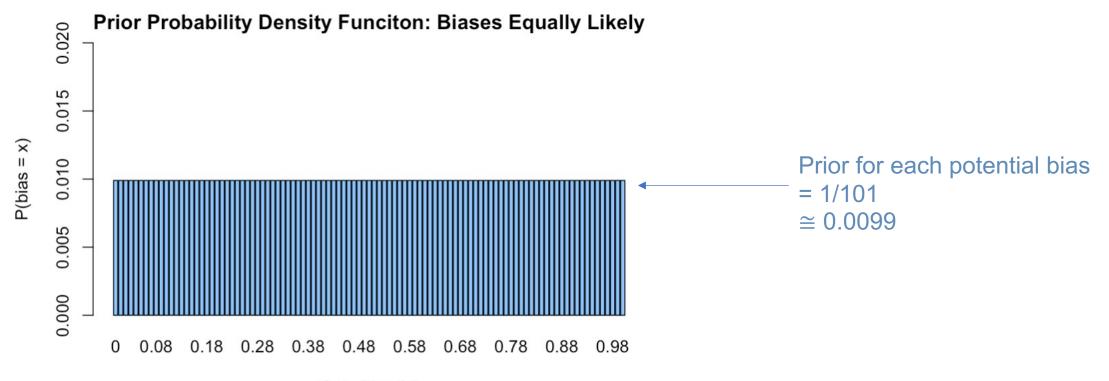
## Bayesian Coin Flip: 20 Flips; 13 Heads

Objective: Identify the bias (x) that yields the highest posterior probability. Given 13 heads were observed out of 20 flips



### **Bayesian Coin Flip: Define Priors**





Coin Bias (x)

#### P(13 heads)

+ 
$$P(13 \text{ heads} | \text{ bias} = 0.50) \cdot P(\text{bias} = 0.50)$$

+ P(13 heads | bias = 
$$0.99$$
) · P(bias =  $0.99$ )  
+ P(13 heads | bias =  $1.00$ ) · P(bias =  $1.00$ )

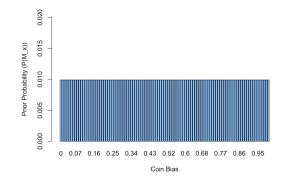
#### P(13 heads)

. . .

=  $P(13 \text{ heads } | \text{ bias } = 0.00) \cdot P(\text{bias } = 0.00)$ +  $P(13 \text{ heads } | \text{ bias } = 0.01) \cdot P(\text{bias } = 0.01)$ +  $P(13 \text{ heads } | \text{ bias } = 0.02) \cdot P(\text{bias } = 0.02)$ 

+ P(13 heads | bias = 
$$0.50$$
) · P(bias =  $0.50$ )

+ P(13 heads | bias = 
$$0.99$$
) · P(bias =  $0.99$ )  
+ P(13 heads | bias =  $1.00$ ) · P(bias =  $1.00$ )



#### P(13 heads)

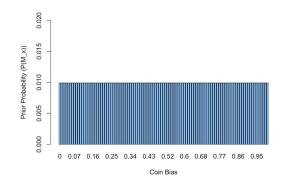
. . .

= P(13 heads | bias = 0.00) · 0.0099 + P(13 heads | bias = 0.01) · 0.0099 + P(13 heads | bias = 0.02) · 0.0099

+ P(13 heads | bias = 
$$0.50$$
) ·  $0.0099$ 

... ⊥ D/12 h

+  $P(13 \text{ heads} | \text{ bias} = 0.99) \cdot 0.0099$ +  $P(13 \text{ heads} | \text{ bias} = 1.00) \cdot 0.0099$ 

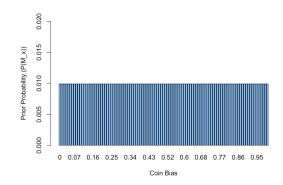


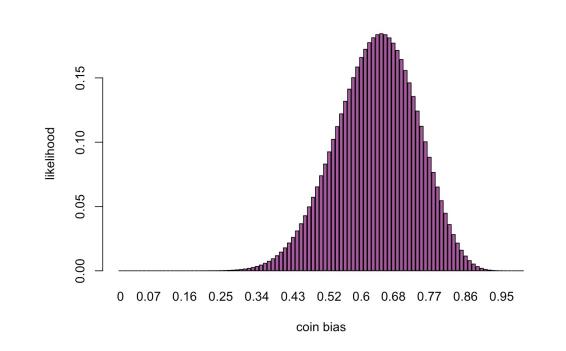
#### P(13 heads)

= P(13 heads | bias = 0.00) · 0.0099 + P(13 heads | bias = 0.01) · 0.0099 + P(13 heads | bias = 0.02) · 0.0099

+  $P(13 \text{ heads} | \text{ bias} = 0.50) \cdot 0.0099$ 

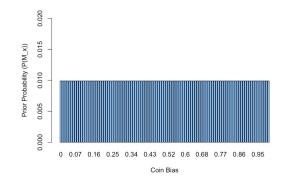
+ P(13 heads | bias = 0.99) · 0.0099 + P(13 heads | bias = 1.00) · 0.0099

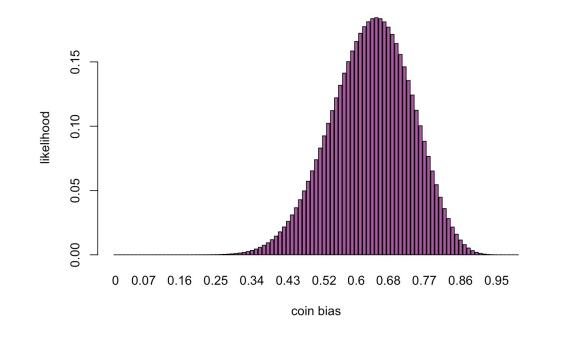




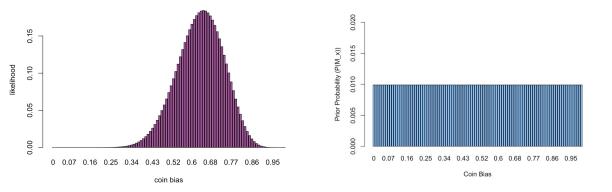
#### P(13 heads)

- = 0.0· 0.0099
  - + 7.2e-22· 0.0099
  - + 5.5e-18 · 0.0099
  - • •
  - + 0.07392883 0.0099
  - • •
  - + 6.8e-10 · 0.0099 + 0.0 · 0.0099





= 0.04714757



coin.bias  $\leq seq(from = 0, to = 1, by = 0.01)$ 

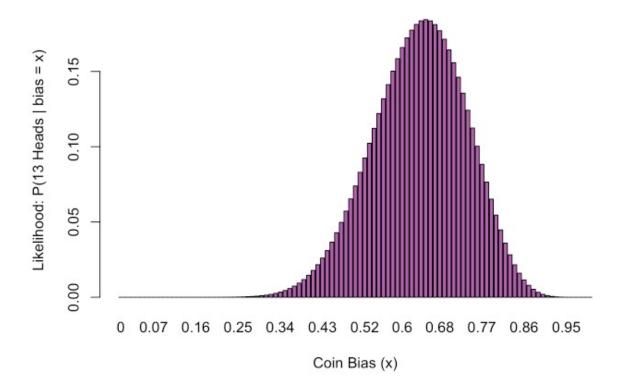
(p.d13 <- sum(dbinom(13, 20, coin.bias) \* (1 / 101)))

## [1] 0.04714757

= 0.04714757

### **Bayesian Coin Flip: Likelihood**

coin.bias <- seq(from = 0, to = 1, by = 0.01)
likelihood <- dbinom(13, 20, prob = coin.bias)
barplot(likelihood, names.arg = coin.bias, ylab = "Likelihood: P(13 Heads | bias = x)", xlab = "Coin Bias (x)", c
ol = "#A95AA1") #Color-blindess friendly purple</pre>

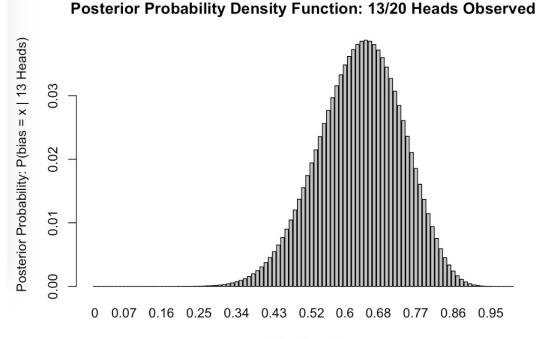


### **Posterior Probability**

posterior.probability <- dbinom(13, 20, coin.bias) \* (1 / 101) / p.d13
sum(posterior.probability)</pre>

#### ## [1] 1

barplot(posterior.probability, names.arg = coin.bias, xlab = "Coin Bias (x)", y lab = "Posterior Probability: P(bias = x | 13 Heads)", main = "Posterior Probab ility Density Function: 13/20 Heads Observed")

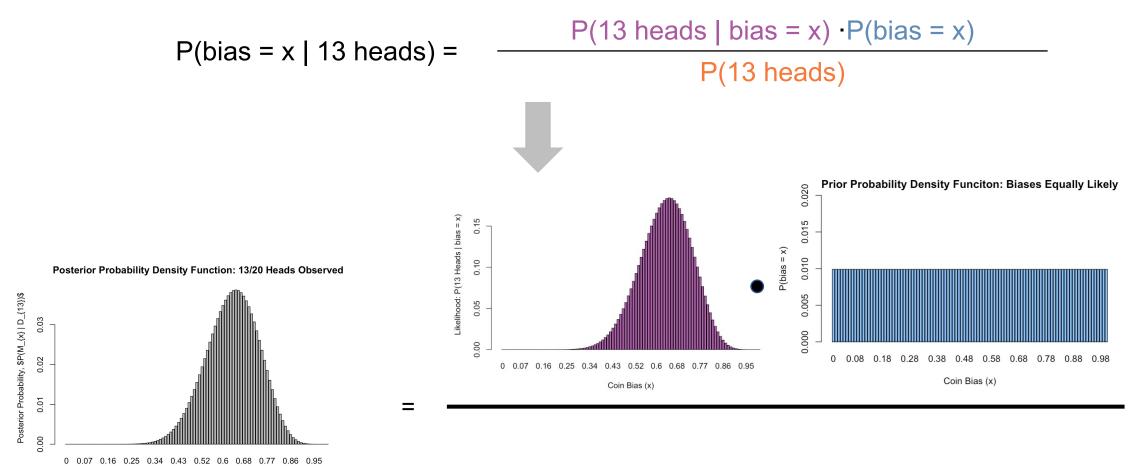


#### Recall Frequentist Conclusion:

- Bias = 0.65
- 95% CI = (0.41, 0.85)

Coin Bias (x)

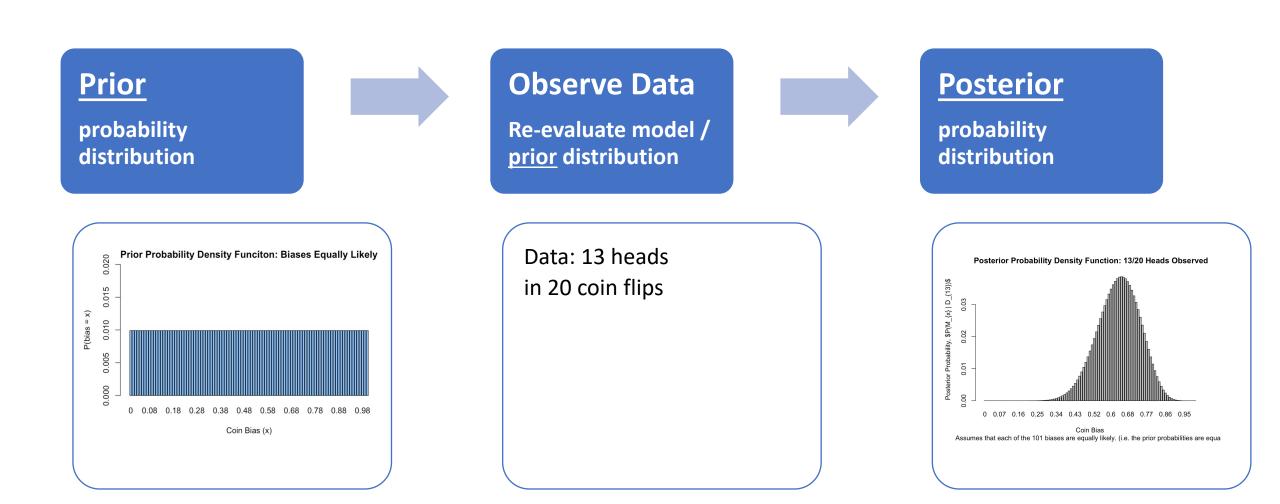
## Summary: Flipping a Coin with No expectations of fairness



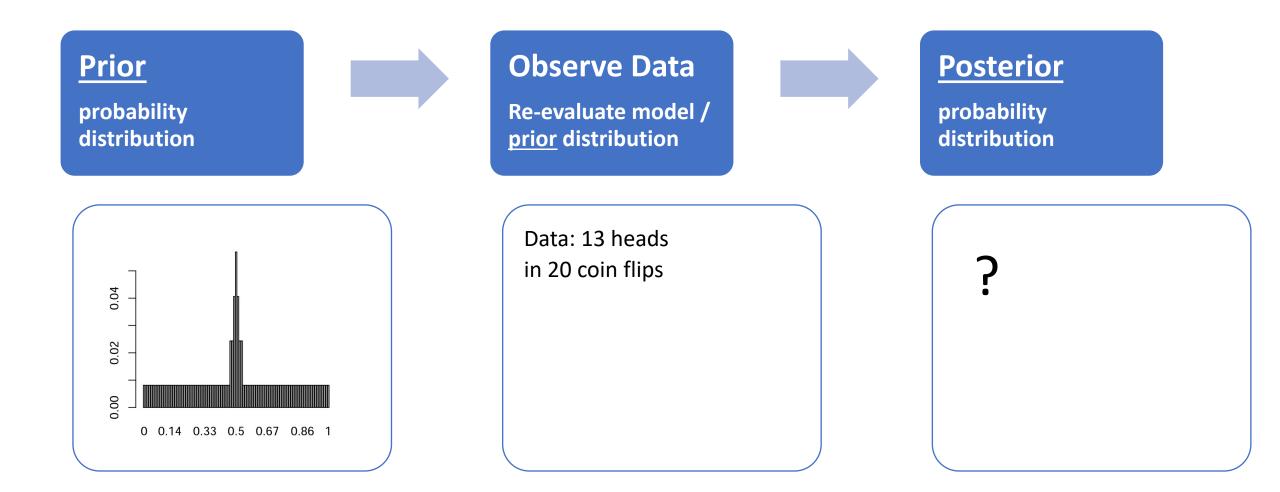
Coin Bias Assumes that each of the 101 biases are equally likely. (i.e. the prior probabilities are equa

#### P(13 Heads) = 0.04714757

## Summary of Bayes' method

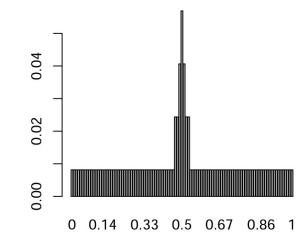


## What if I assume there is a good chance of the coin having a certain "bias"?



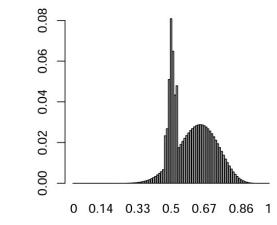
## Our <u>prior</u> will reflect our assumption that our friend is honest

```
prior.probability <- numeric(101)
prior.probability[0:101] <- 1
prior.probability[48:54] <- 3
prior.probability[50:52] <- 5
prior.probability[51] <- 7
# Normalize; since it is a PDF, sum must be 1.0
prior.probability <- prior.probability / (sum(prior.probability))
barplot(prior.probability, names.arg = coin.bias)</pre>
```

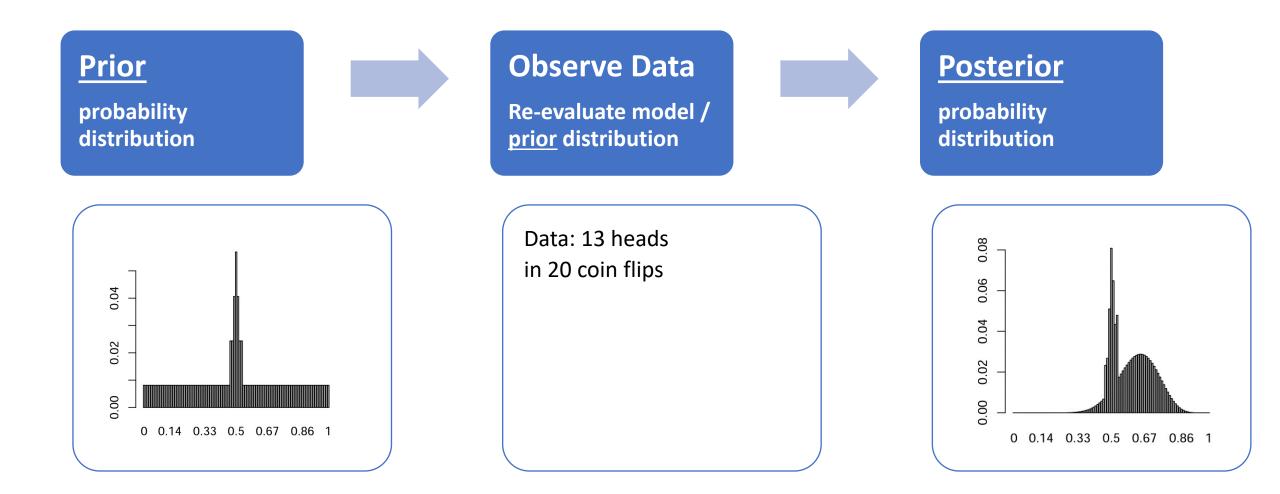


Our <u>posterior</u> probability distribution reflects a complex interplay between the prior and the data

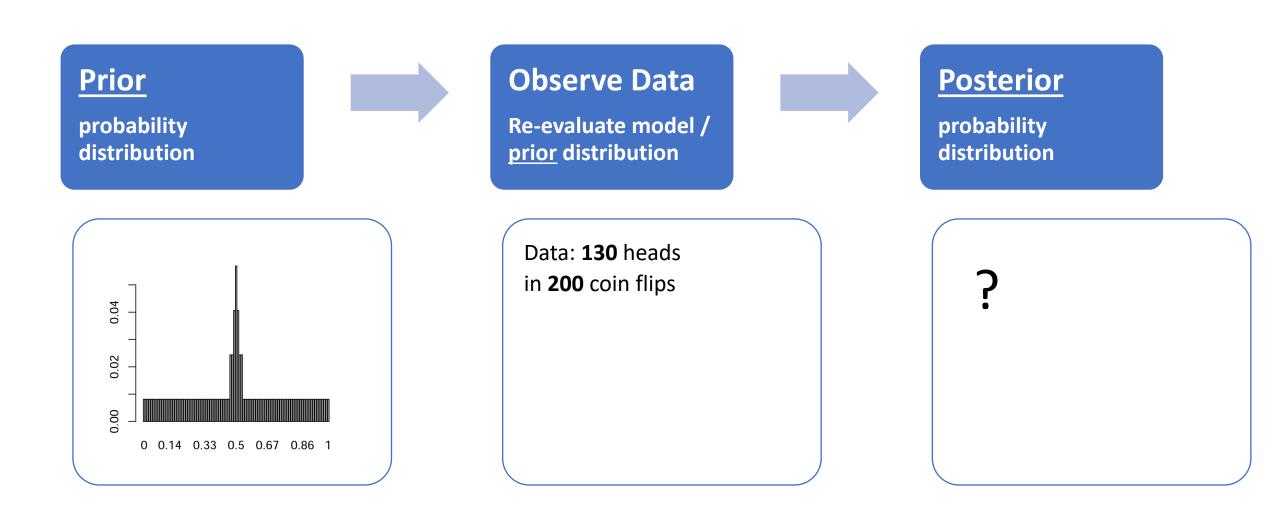
```
posterior.probability <-
   dbinom(13, 20, coin.bias) * prior.probability / p.d13
barplot(posterior.probability, names.arg = coin.bias)</pre>
```



## What if I assume there is a good chance of the coin having a certain "bias"?

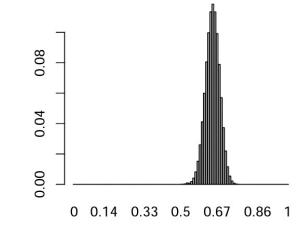


### What if we collect more data?

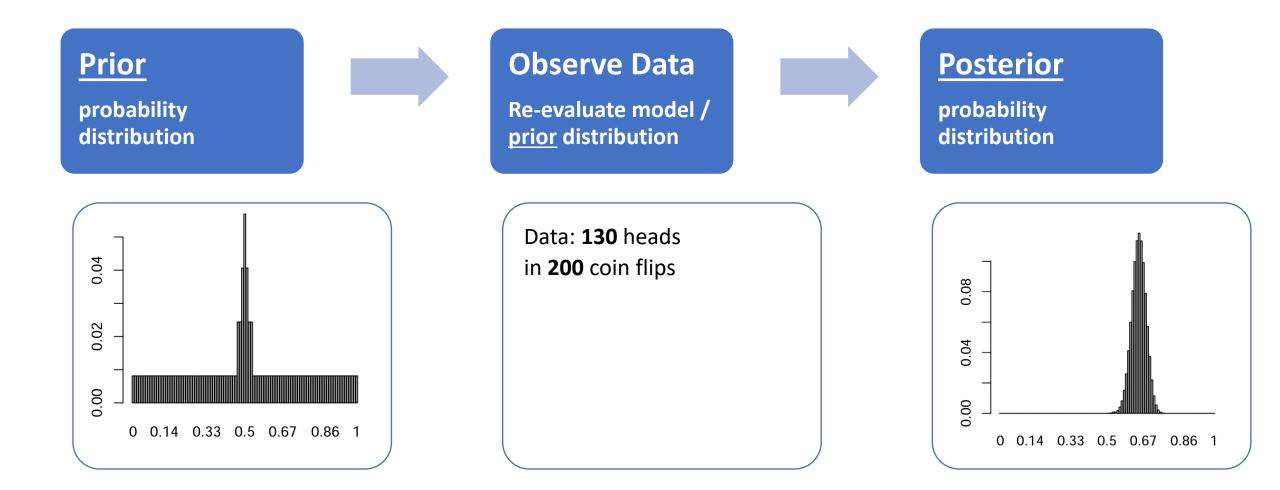


### Now in our <u>posterior</u> probability, the data "overwrites" our prior

```
p.d130 <- sum(dbinom(130, 200, coin.bias) * prior.probability)
posterior.probability <-
    dbinom(130, 200, coin.bias) * prior.probability / p.d130
barplot(posterior.probability, names.arg = coin.bias)</pre>
```

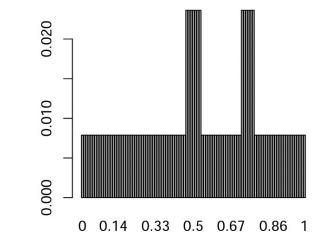


## When we collect more data, the data "overwrites" our prior



## Another example – imagine there is a magic shop around the corner...

```
prior.probability <- numeric(101)
prior.probability[0:101] <- 1
prior.probability[48:54] <- 3
prior.probability[73:78] <- 3
# Normalize; since it is a PDF, sum must be 1.0
prior.probability <- prior.probability / (sum(prior.probability))
barplot(prior.probability, names.arg = coin.bias)</pre>
```

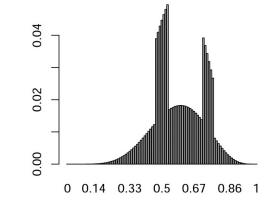


## If we assume we are talking to a swindler, our posterior will reflect that

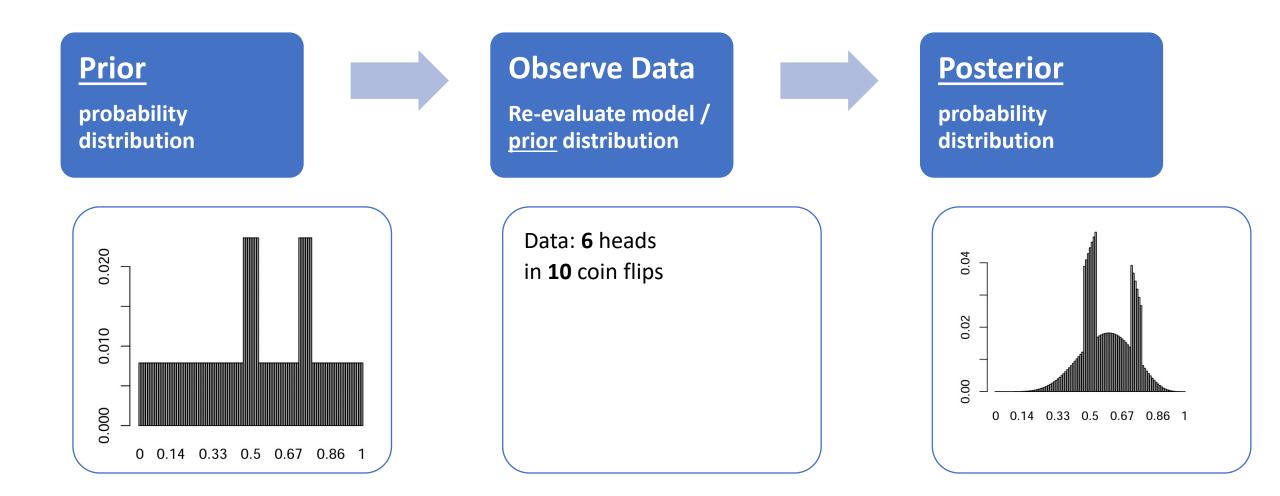
```
(p.d6 <- sum(dbinom(6, 10, coin.bias) * prior.probability))</pre>
```

```
## [1] 0.1083962
```

```
posterior.probability <- dbinom(6, 10, coin.bias) * prior.probability / p.d6
barplot(posterior.probability, names.arg = coin.bias)
```



## If there is a magic shop around the corner, we conclude the coin may be biased



## Conclusion

#### **Bayesian statistics**

- Start with our understanding of how something works/ what is likely to happen
- We then update our belief based on our data
- Possible to perform multiple rounds of formulation of prior, evaluation of prior based on data and formulation of posterior.
- Does not rely on the notion of a finding "as or more inconsistent with our H<sub>0</sub>"

#### **Frequentist approaches**

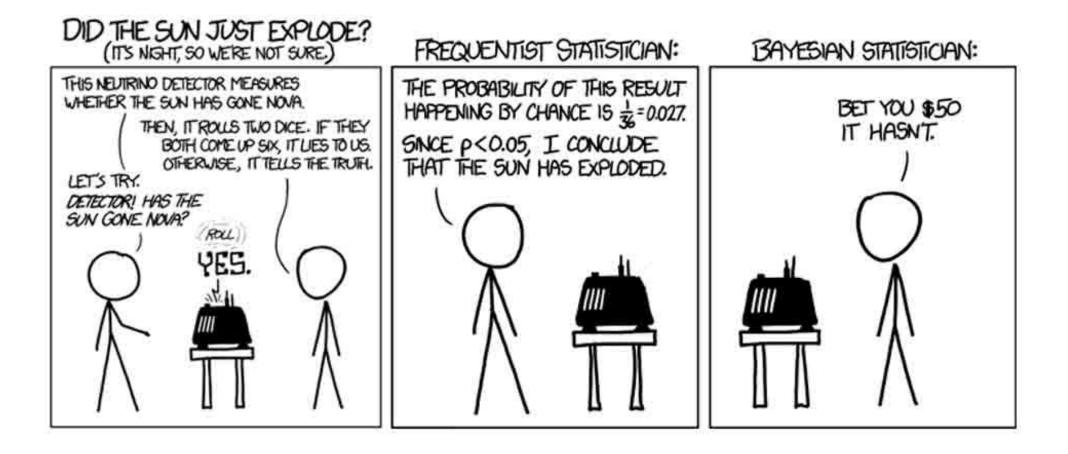
- Do not assign probabilities to a hypothesis (no prior, posterior)
- Usually less computationally intensive
- Lower risk of bias

### References

- Banfelder, J. Quantitative Understanding in Biology 1.7 Bayesian Methods (<u>https://physiology.med.cornell.edu/people/banfelder/qbio/lecture\_notes/1.7\_bayesian.pdf</u>)
- Orloff, J. and Bloom, J."Comparison of frequentist and Bayesian inference." 2014 (<u>https://ocw.mit.edu/courses/mathematics/18-05-introduction-to-probability-and-statistics-spring-2014/readings/MIT18\_05S14\_Reading20.pdf</u>)

Further interesting materials on this topic:

- Kruschke, J. "Doing Bayesian Data Analysis"
- <u>https://boyangzhao.github.io/posts/vaccine\_efficacy\_bayesian</u> (advanced blog post about how Bayesian statistics were used to determine COVID-19 vaccine efficacy)
- <u>https://youtu.be/9TDjifpGj-k</u> (fun crash course on the basics of Bayesian statistics)



https://www.cs.ubc.ca/~murphyk/Bayes/bayesrule.html