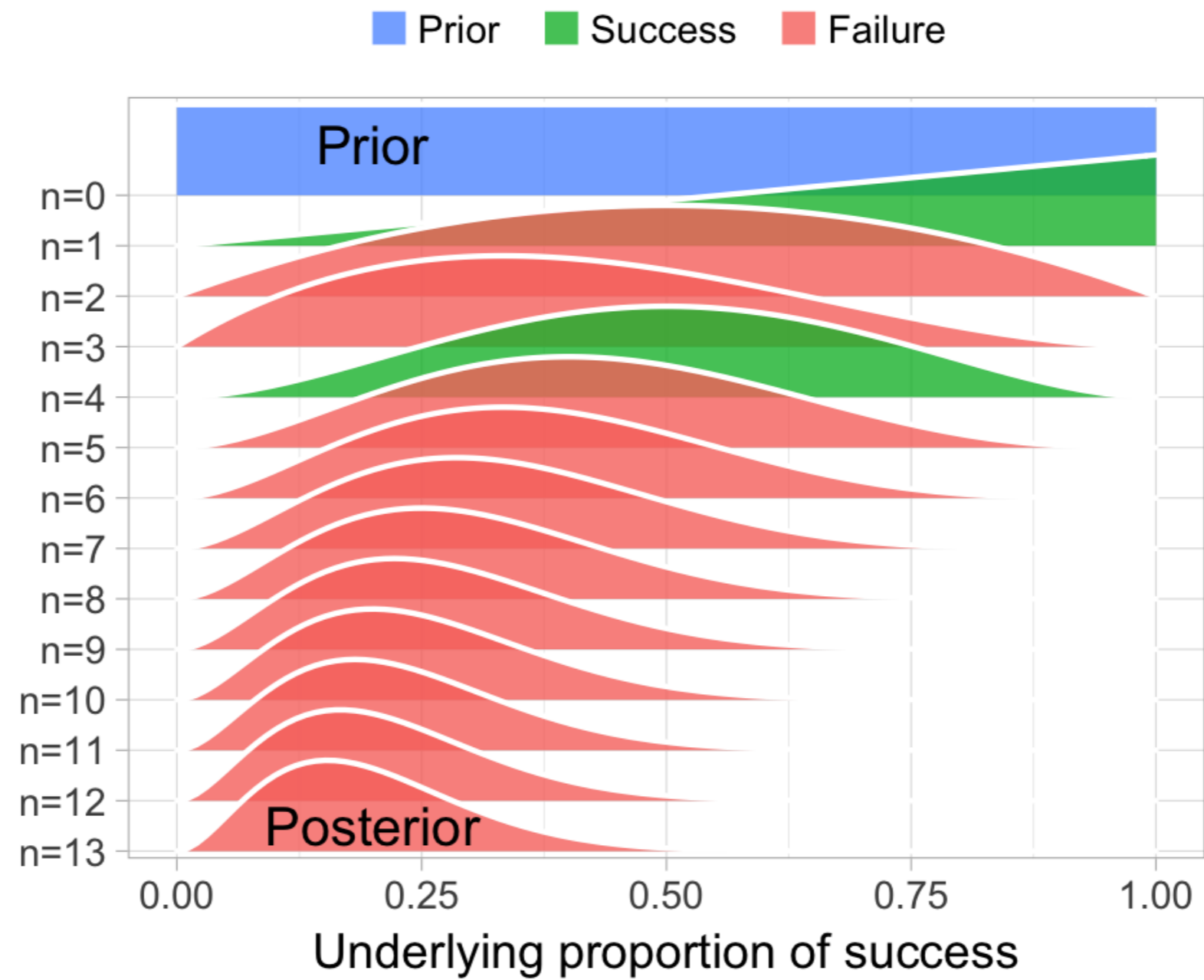


The parts needed for Bayesian inference

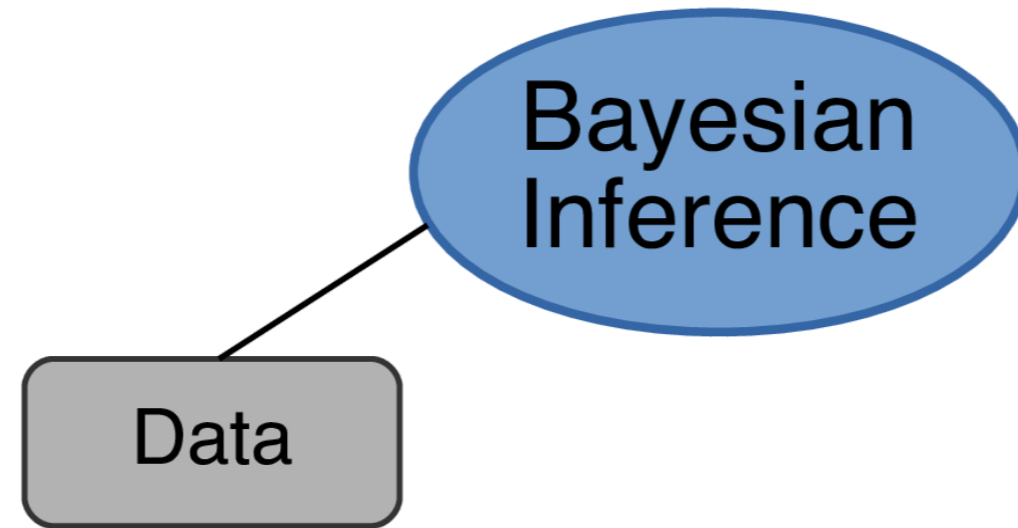
FUNDAMENTALS OF BAYESIAN DATA ANALYSIS IN R

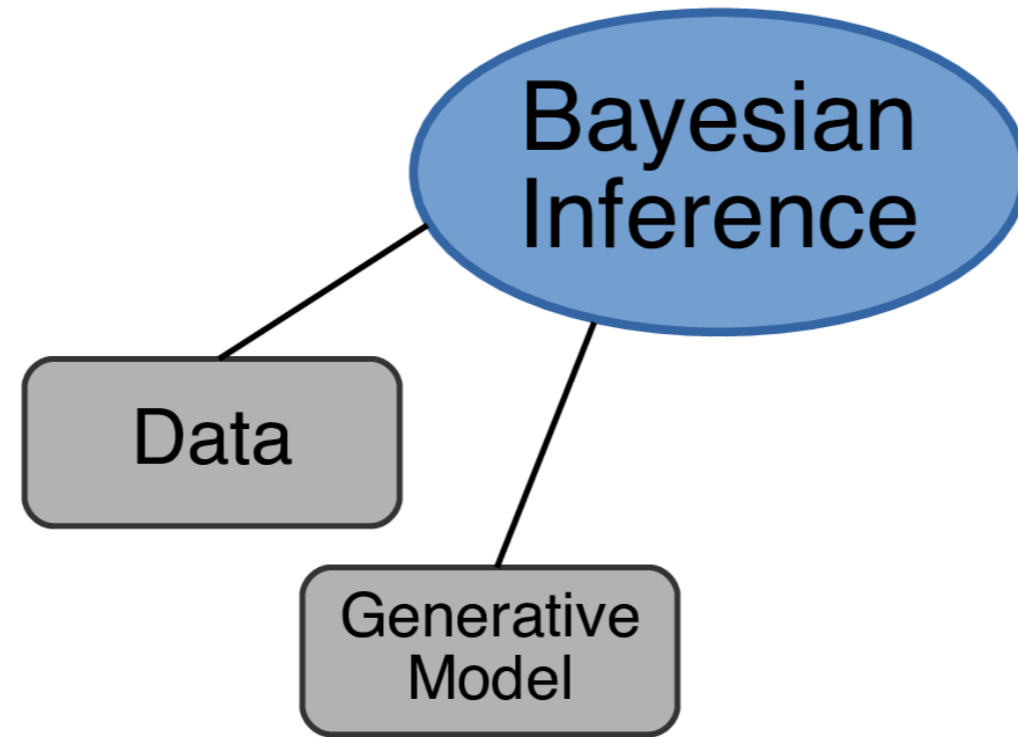


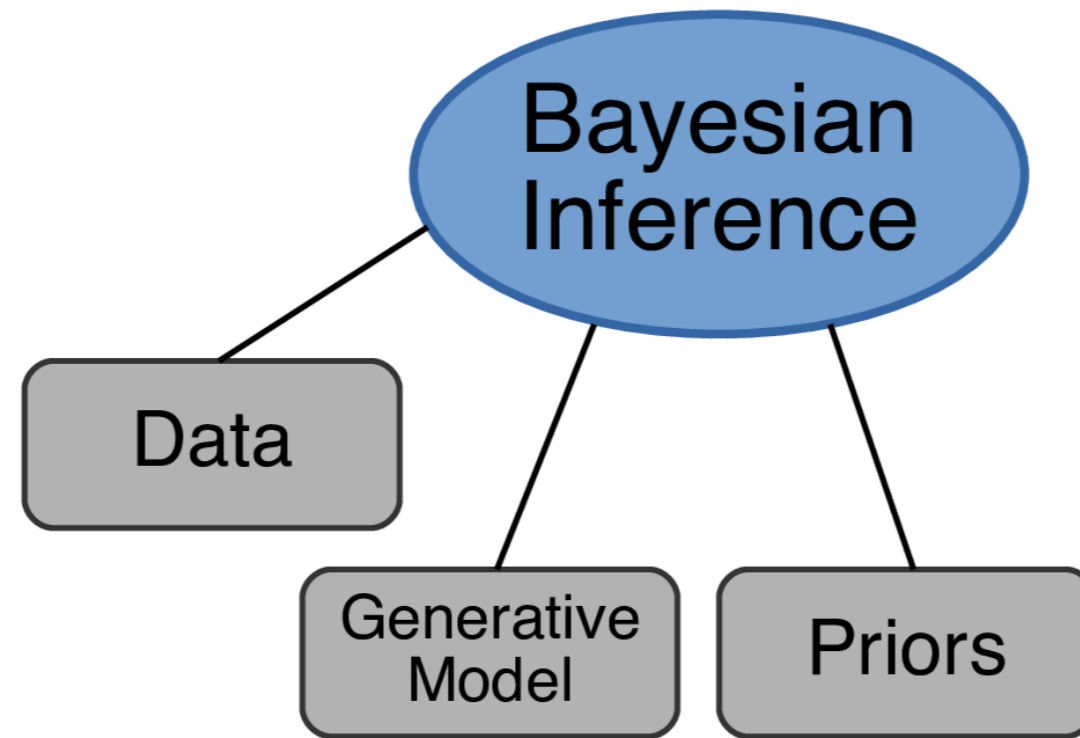
Rasmus Bååth
Data Scientist



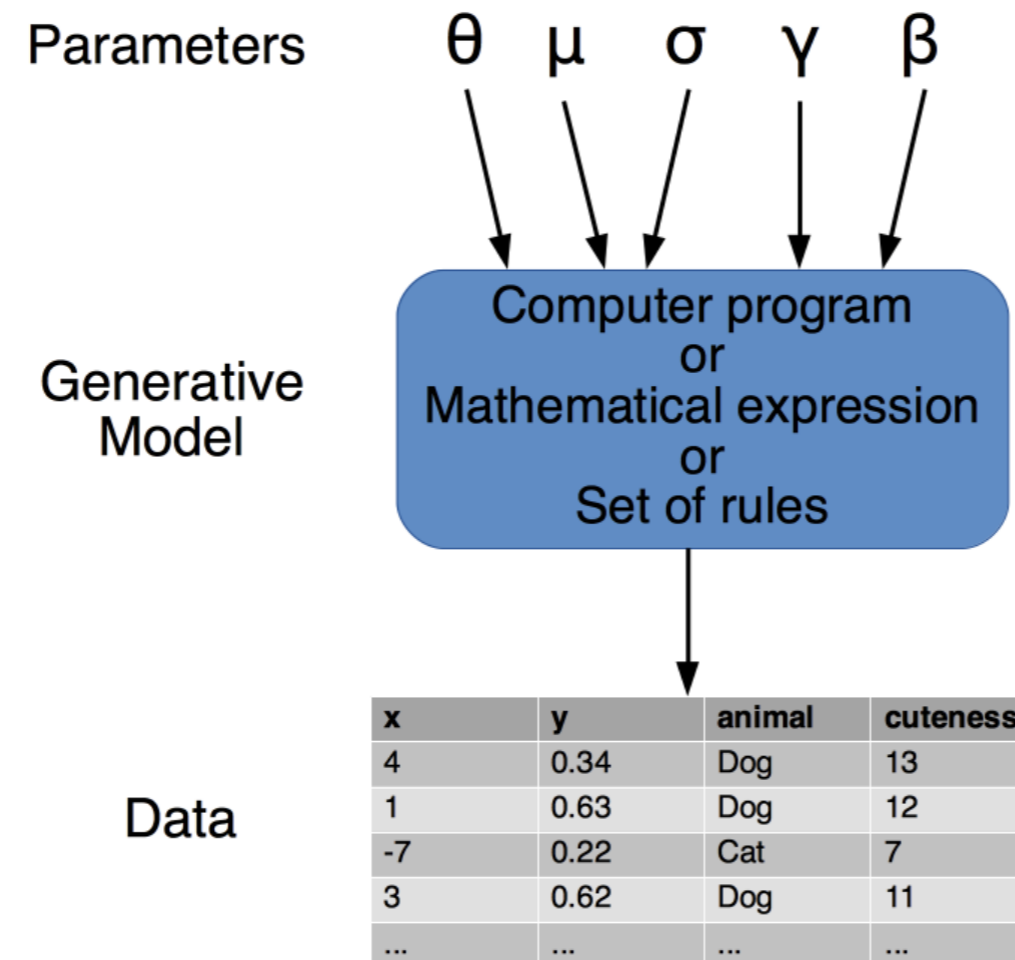
Bayesian Inference







What is a generative model?



Generative zombie drug model

Generative zombie drug model

```
# Parameters  
prop_success <- ???  
n_zombies <- ???
```

Generative zombie drug model

```
# Parameters  
prop_success <- 0.15  
n_zombies <- 13
```

Generative zombie drug model

```
# Parameters
prop_success <- 0.15
n_zombies <- 13
# Simulating data
data <- c()
for(zombie in 1:n_zombies) {
  data[zombie] <- ???
}
```

Generative zombie drug model

```
# Parameters
prop_success <- 0.15
n_zombies <- 13
# Simulating data
data <- c()
for(zombie in 1:n_zombies) {
  data[zombie] <- runif(1, min = 0, max = 1) < prop_success
}
```

Generative zombie drug model

```
# Parameters
prop_success <- 0.15
n_zombies <- 13
# Simulating data
data <- c()
for(zombie in 1:n_zombies) {
  data[zombie] <- runif(1, min = 0, max = 1) < prop_success}
```

```
data
```

```
FALSE FALSE FALSE TRUE FALSE FALSE FALSE TRUE FALSE TRUE FALSE FALSE
```

Generative zombie drug model

```
# Parameters
prop_success <- 0.15
n_zombies <- 13
# Simulating data
data <- c()
for(zombie in 1:n_zombies) {
  data[zombie] <- runif(1, min = 0, max = 1) < prop_success
}
data <- as.numeric(data)
```

Generative zombie drug model

```
# Parameters
prop_success <- 0.15
n_zombies <- 13
# Simulating data
data <- c()
for(zombie in 1:n_zombies) {
  data[zombie] <- runif(1, min = 0, max = 1) < prop_success
}
data <- as.numeric(data)
data
```

```
0 0 0 1 0 0 0 0 1 0 1 0 0
```

Generative zombie drug model

```
# Parameters
prop_success <- 0.15
n_zombies <- 13
# Simulating data
data <- c()
for(zombie in 1:n_zombies) {
  data[zombie] <- runif(1, min = 0, max = 1) < prop_success
}
data <- as.numeric(data)
data
```

```
0 0 1 0 0 0 0 0 0 0 0 0 0
```


Generative zombie drug model

```
# Parameters
prop_success <- 0.15
n_zombies <- 13
# Simulating data
data <- c()
for(zombie in 1:n_zombies) {
  data[zombie] <- runif(1, min = 0, max = 1) < prop_success
}
data <- as.numeric(data)
data
```

```
0 1 0 1 1 0 0 1 0 1 0 0 0
```

Generative zombie drug model

```
# Parameters
prop_success <- 0.15
n_zombies <- 13
# Simulating data
data <- c()
for(zombie in 1:n_zombies) {
  data[zombie] <- runif(1, min = 0, max = 1) < prop_success
}
data <- as.numeric(data)
data
```

```
0 0 0 0 0 0 0 1 0 0 0 0 0
```

Generative zombie drug model

```
# Parameters
prop_success <- 0.15
n_zombies <- 13
# Simulating data
data <- c()
for(zombie in 1:n_zombies) {
  data[zombie] <- runif(1, min = 0, max = 1) < prop_success
}
data <- as.numeric(data)
data
```

```
0 0 0 0 1 0 0 0 0 1 0 1 0
```

**Take this model for a
spin!**

FUNDAMENTALS OF BAYESIAN DATA ANALYSIS IN R

Using a generative model

FUNDAMENTALS OF BAYESIAN DATA ANALYSIS IN R

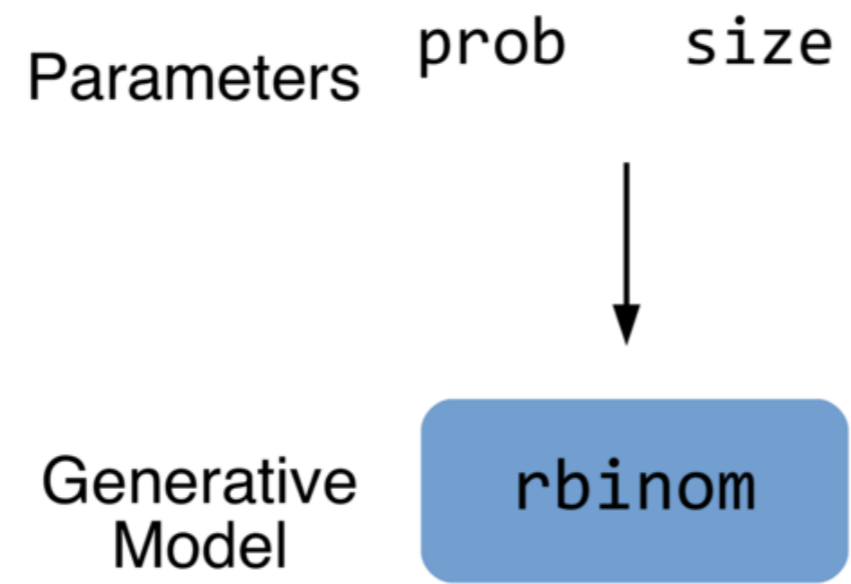


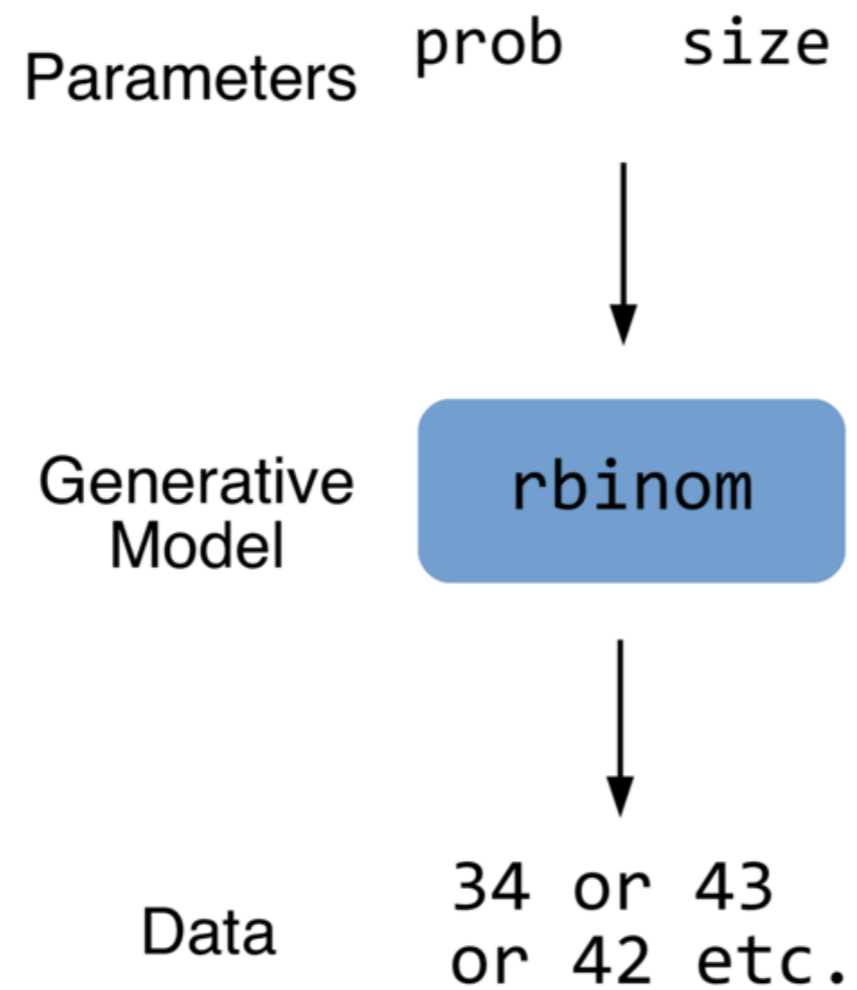
Rasmus Bååth
Data Scientist

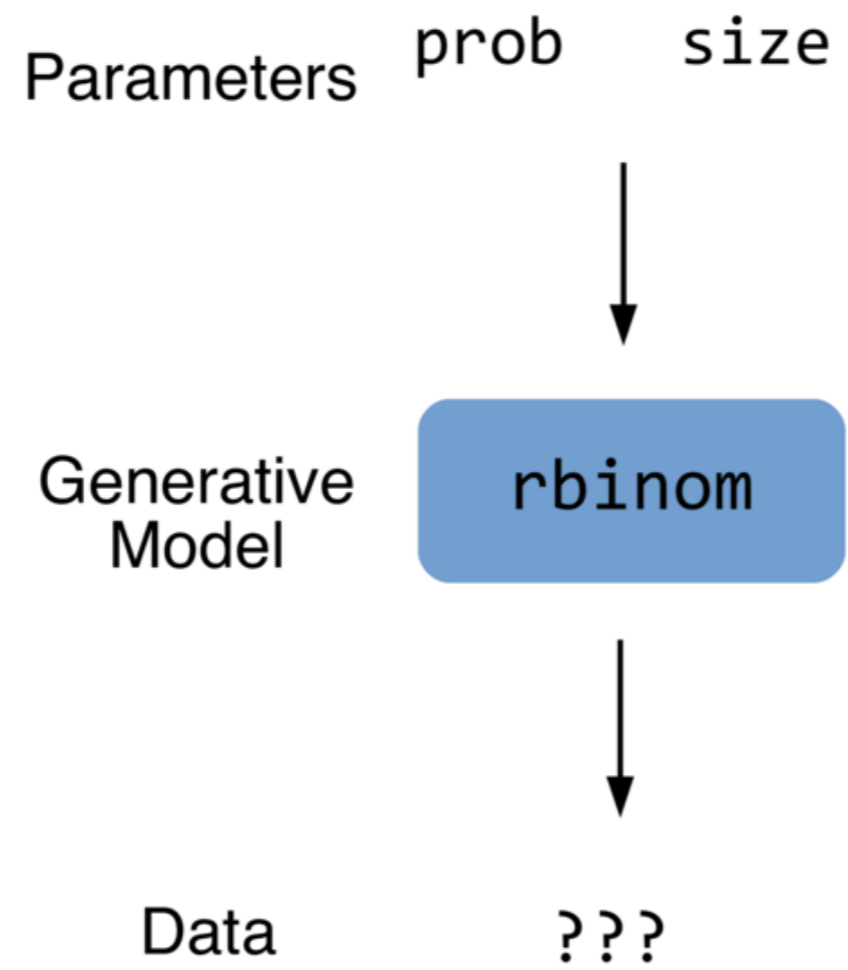
```
rbinom(n, size, prob)
```

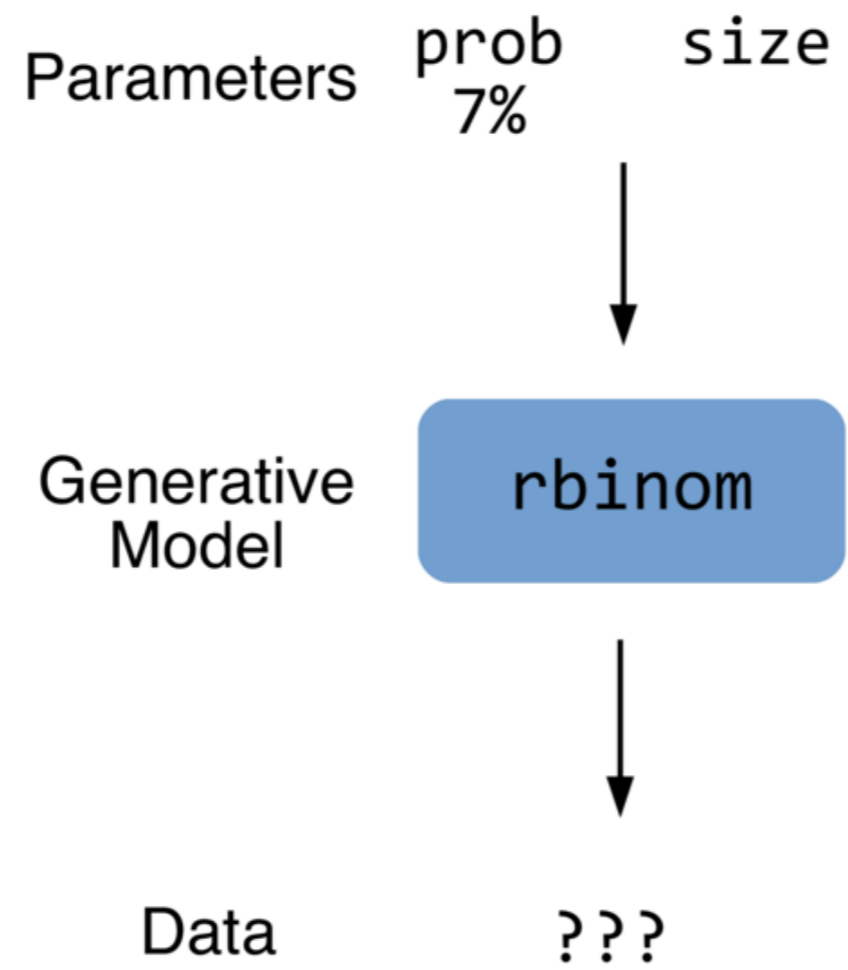
Generative
Model

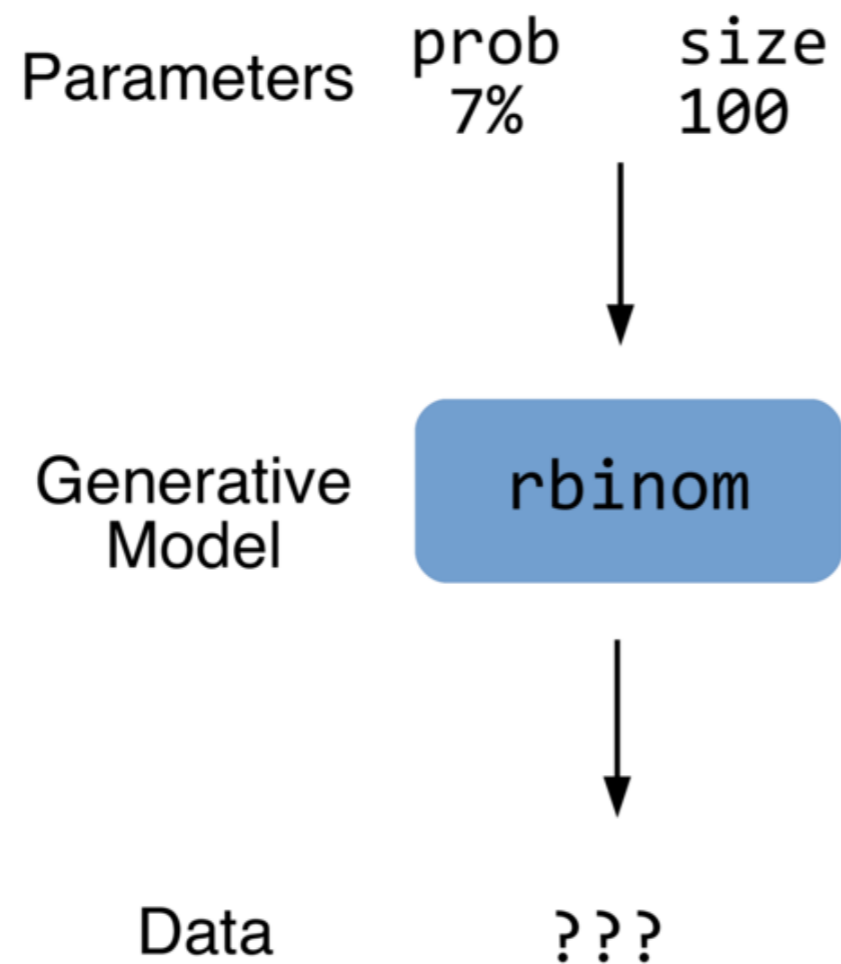
`rbinom`











¹ https://commons.wikimedia.org/wiki/File:The_Hoard_III_-_Flickr_-_SoulStealer.co.uk.jpg

```
cured_zombies <- rbinom(n = 1000000, size = 100, prob = 0.07)
```

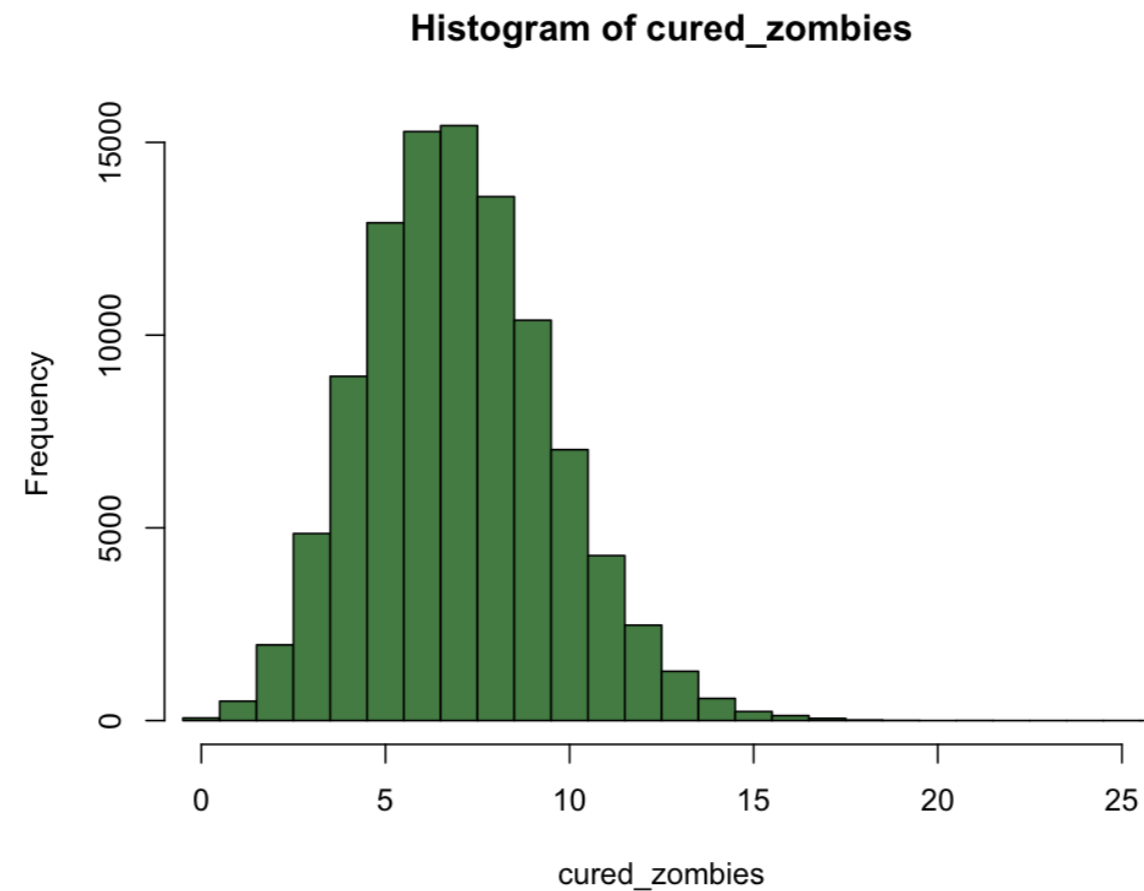
```
cured_zombies <- rbinom(n = 1000000, size = 100, prob = 0.07)
```

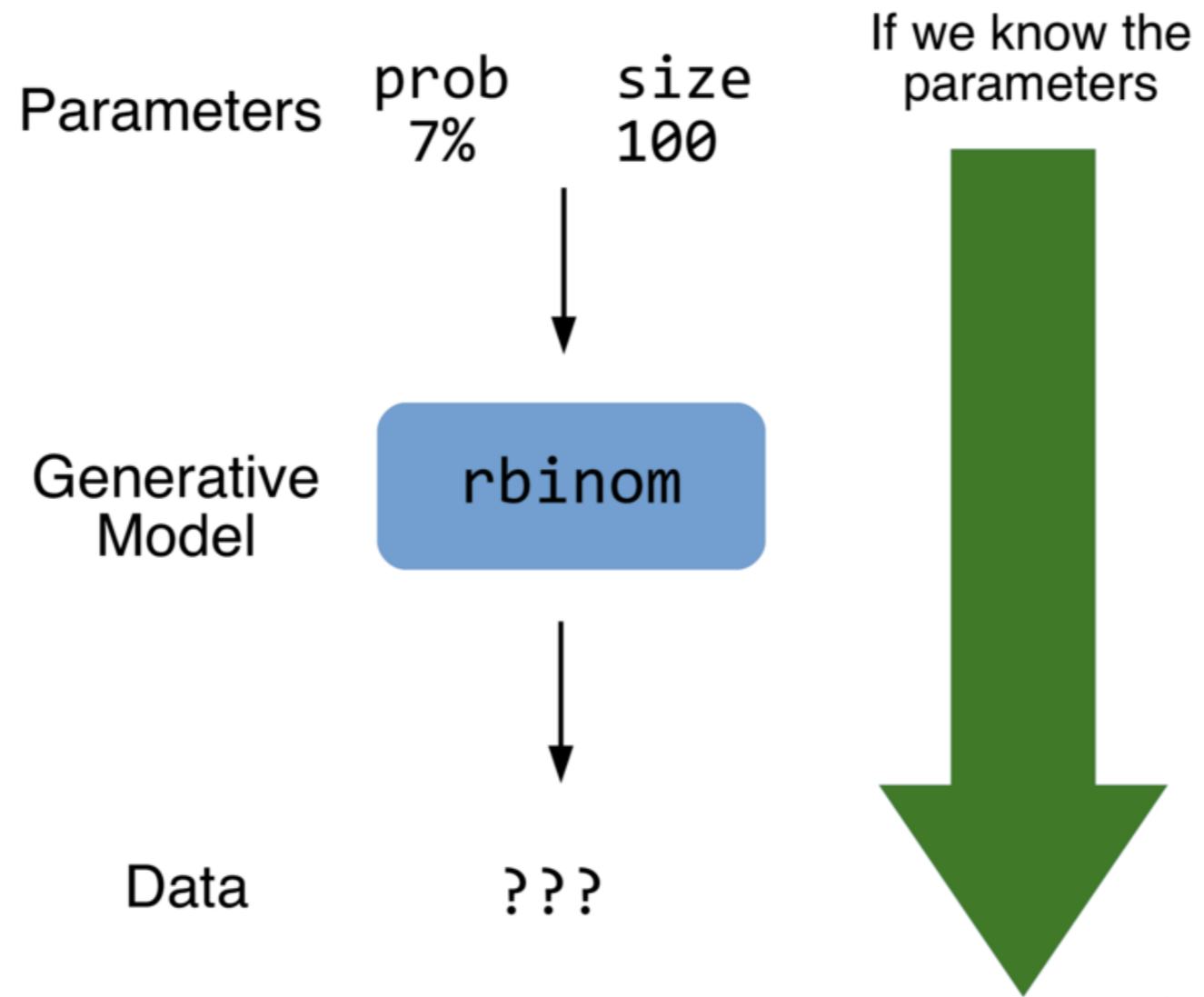
```
cured_zombies
```

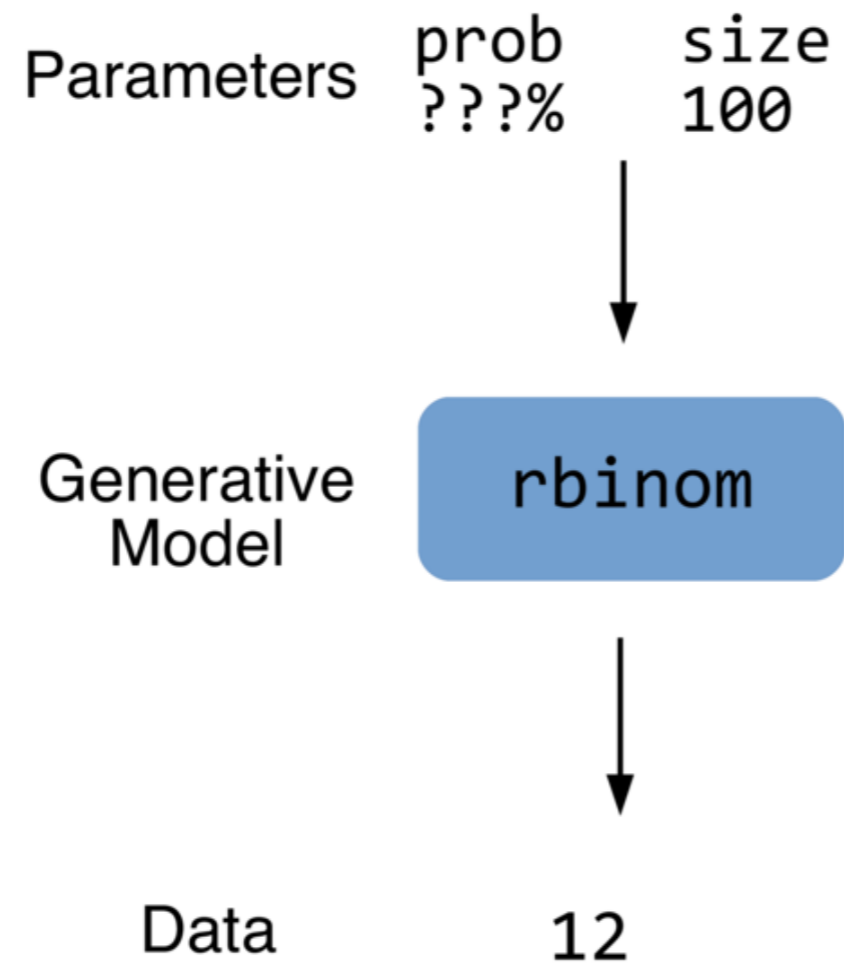
```
8  6  5  8 11  5  7  7  5  8  7 11  6  8  8  
9  9  4  7  4  8  7  5  5  6  9 14  4  4  8  
...
```

```
cured_zombies <- rbinom(n = 100000, size = 100, prob = 0.07)
```

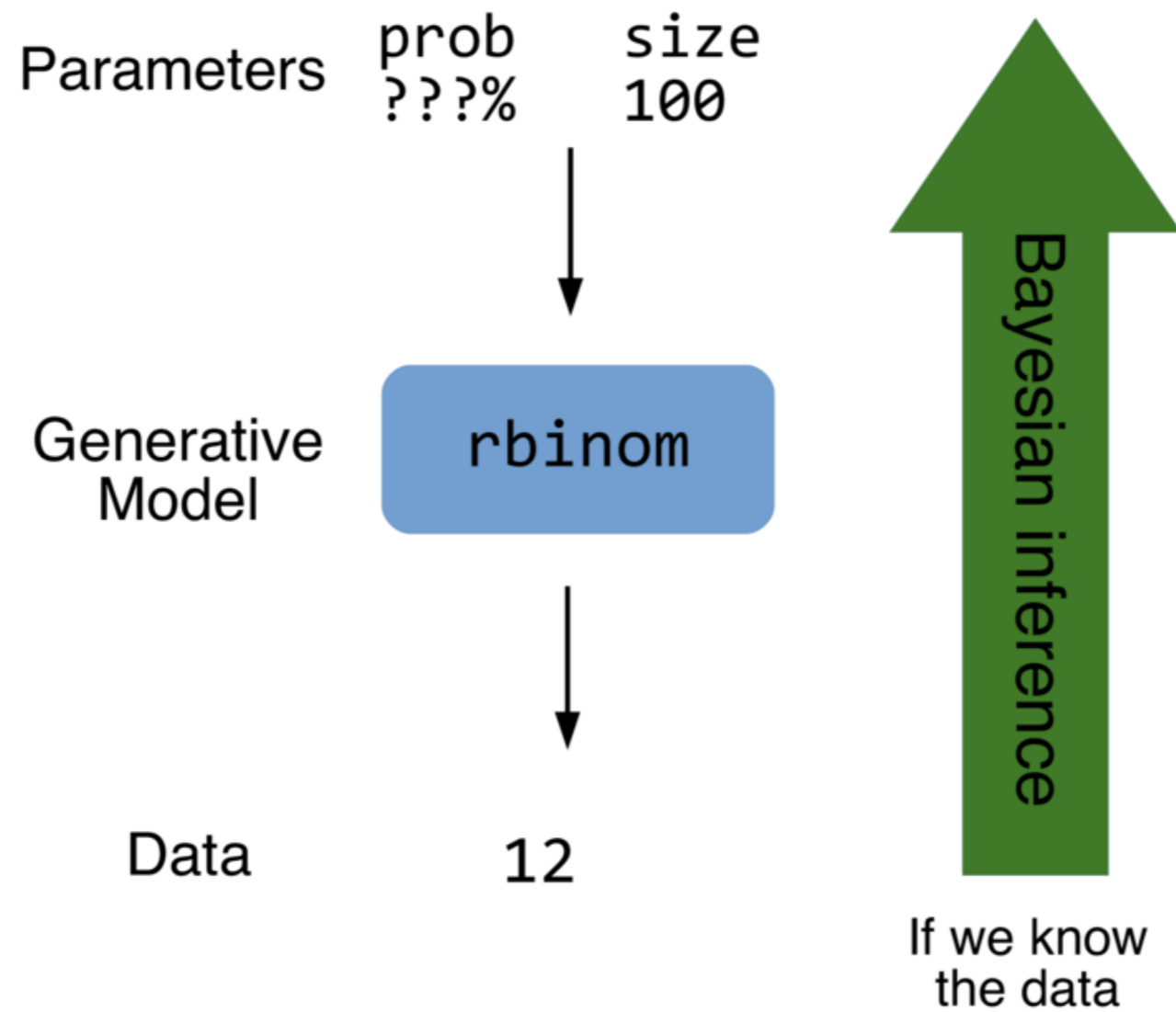
```
hist(cured_zombies)
```

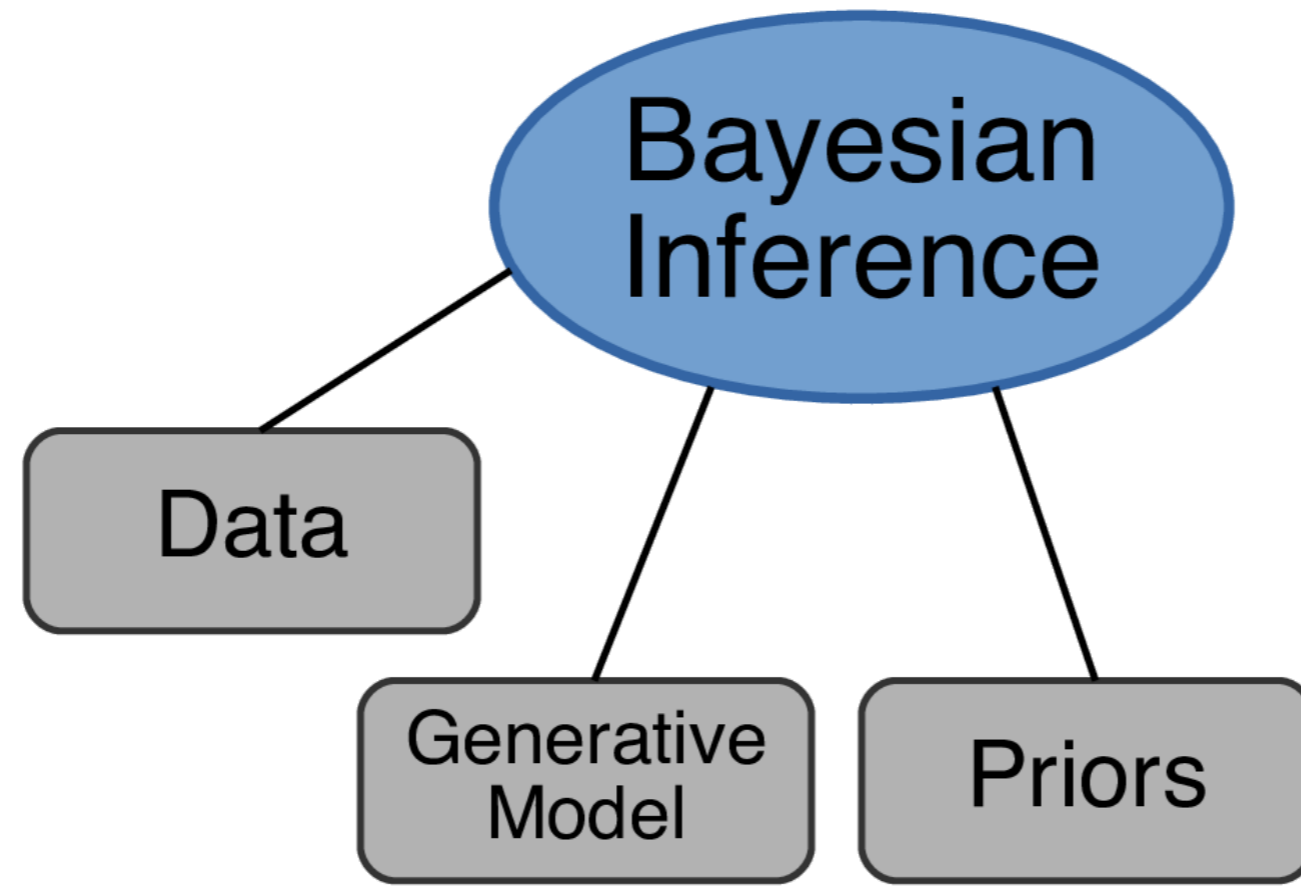


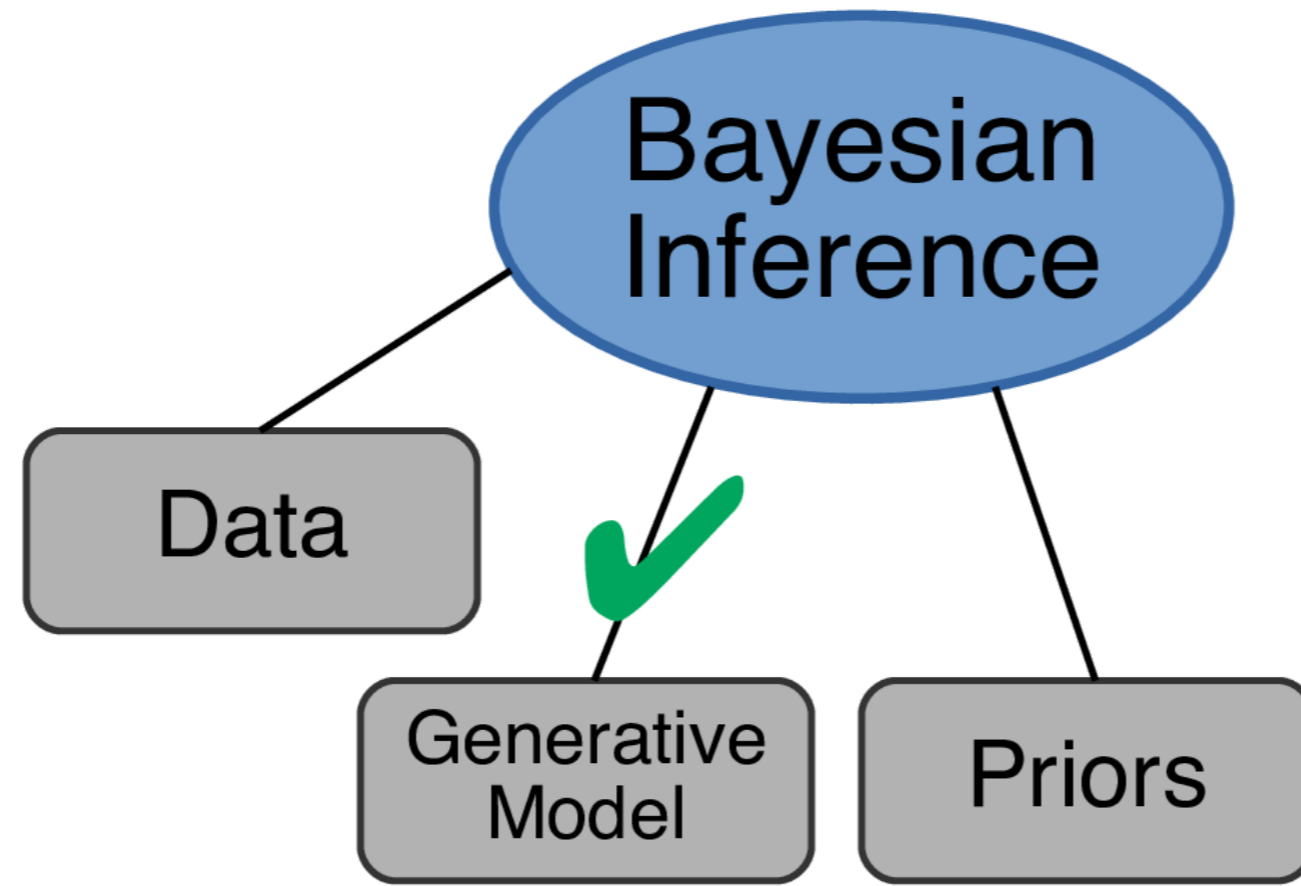




If we know
the data









¹ https://commons.wikimedia.org/wiki/File:The_Hoard_III_-_Flickr_-_SoulStealer.co.uk.jpg





How many visitors?

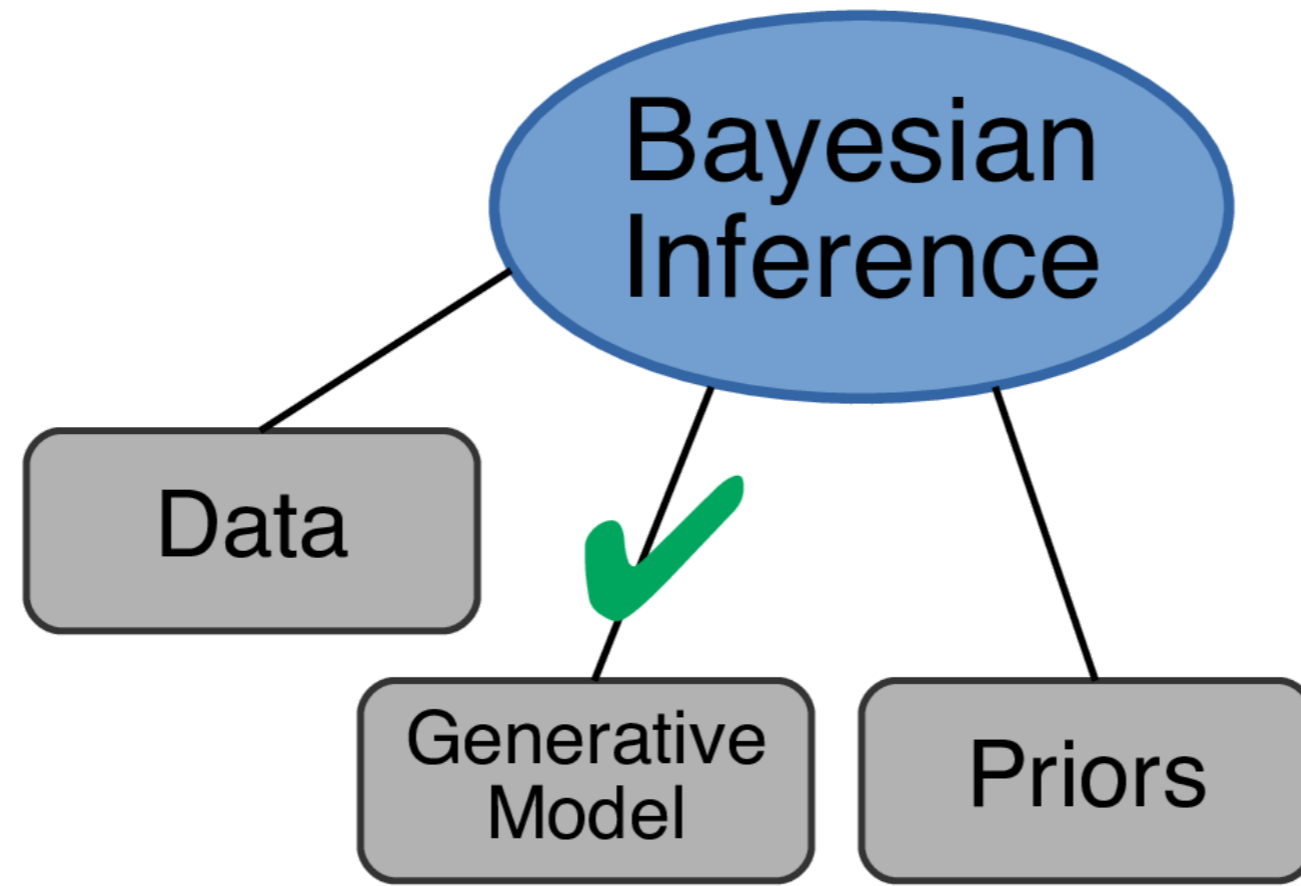
FUNDAMENTALS OF BAYESIAN DATA ANALYSIS IN R

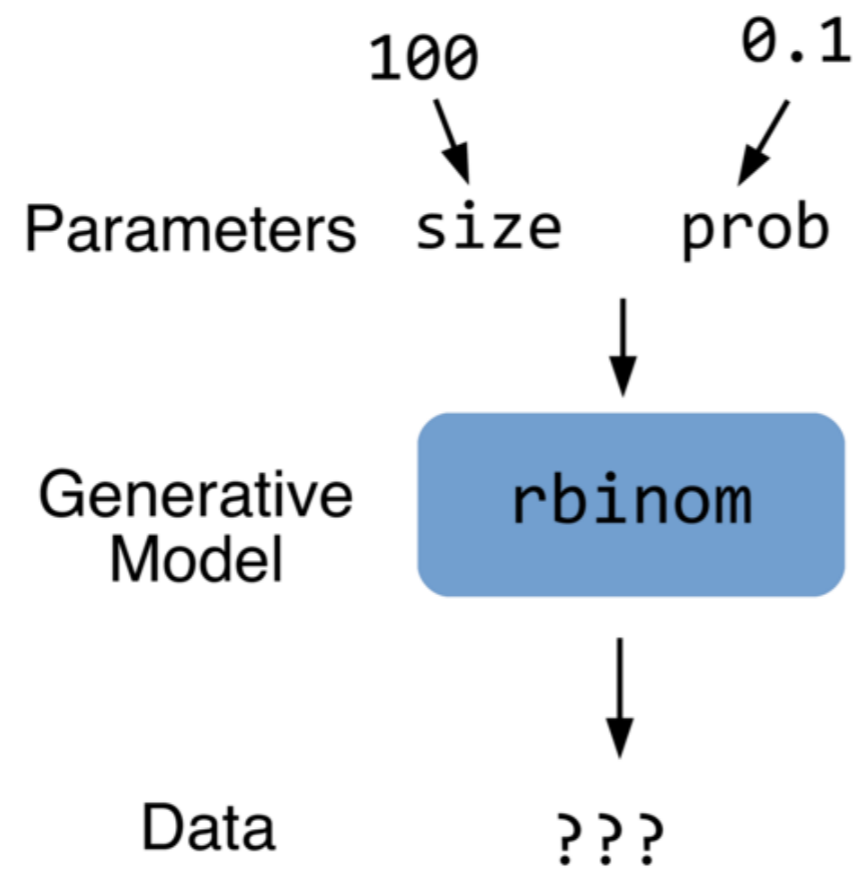
One more part

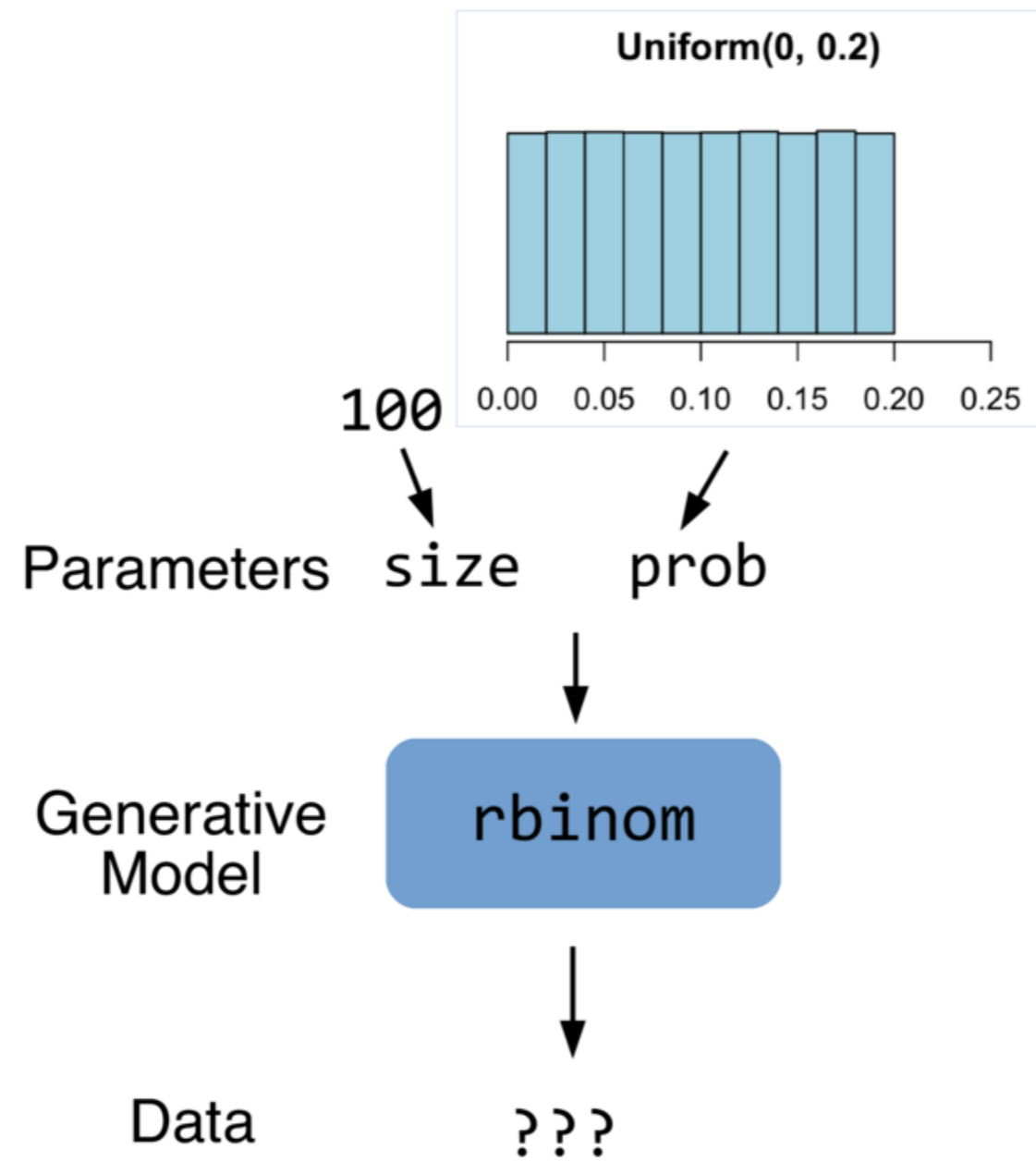
FUNDAMENTALS OF BAYESIAN DATA ANALYSIS IN R



Rasmus Bååth
Data Scientist







rbinom Bayesian model 2

```
n_samples <- 1000000
n_ads_shown <- 100
proportion_clicks <- 0.1
n_visitors <- rbinom(n_samples, size = n_ads_shown,
                    prob = proportion_clicks)
```

runif - Random Uniform samples

runif - Random Uniform samples

```
proportion_clicks <- runif(n = 6, min = 0.0, max = 1.0)
```

```
proportion_clicks
```

```
0.05 0.58 0.21 0.61 0.69 0.39
```

runif - Random Uniform samples

```
proportion_clicks <- runif(n = 6, min = 0.0, max = 1.0)  
n_clicks <- rbinom(n = 6, size = 100, proportion_clicks)
```

```
proportion_clicks
```

```
0.05  0.58  0.21  0.61  0.69  0.39
```


runif - Random Uniform samples

```
proportion_clicks <- runif(n = 6, min = 0.0, max = 1.0)
n_clicks <- rbinom(n = 6, size = 100, proportion_clicks)
```

```
proportion_clicks
```

```
0.05  0.58  0.21  0.61  0.69  0.39
```

```
n_clicks
```

```
7    59    13    63    67    29
```

Try this in practice!

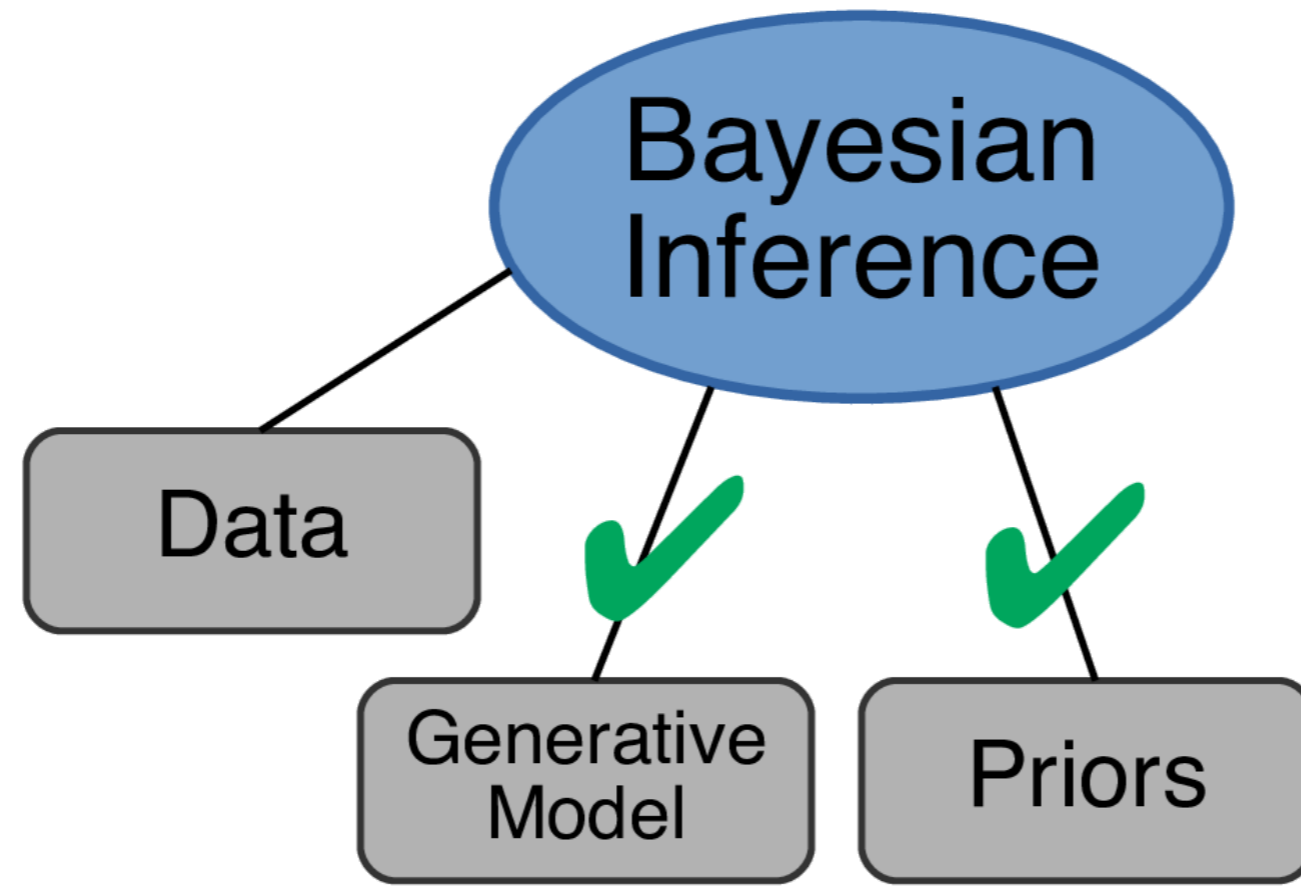
FUNDAMENTALS OF BAYESIAN DATA ANALYSIS IN R

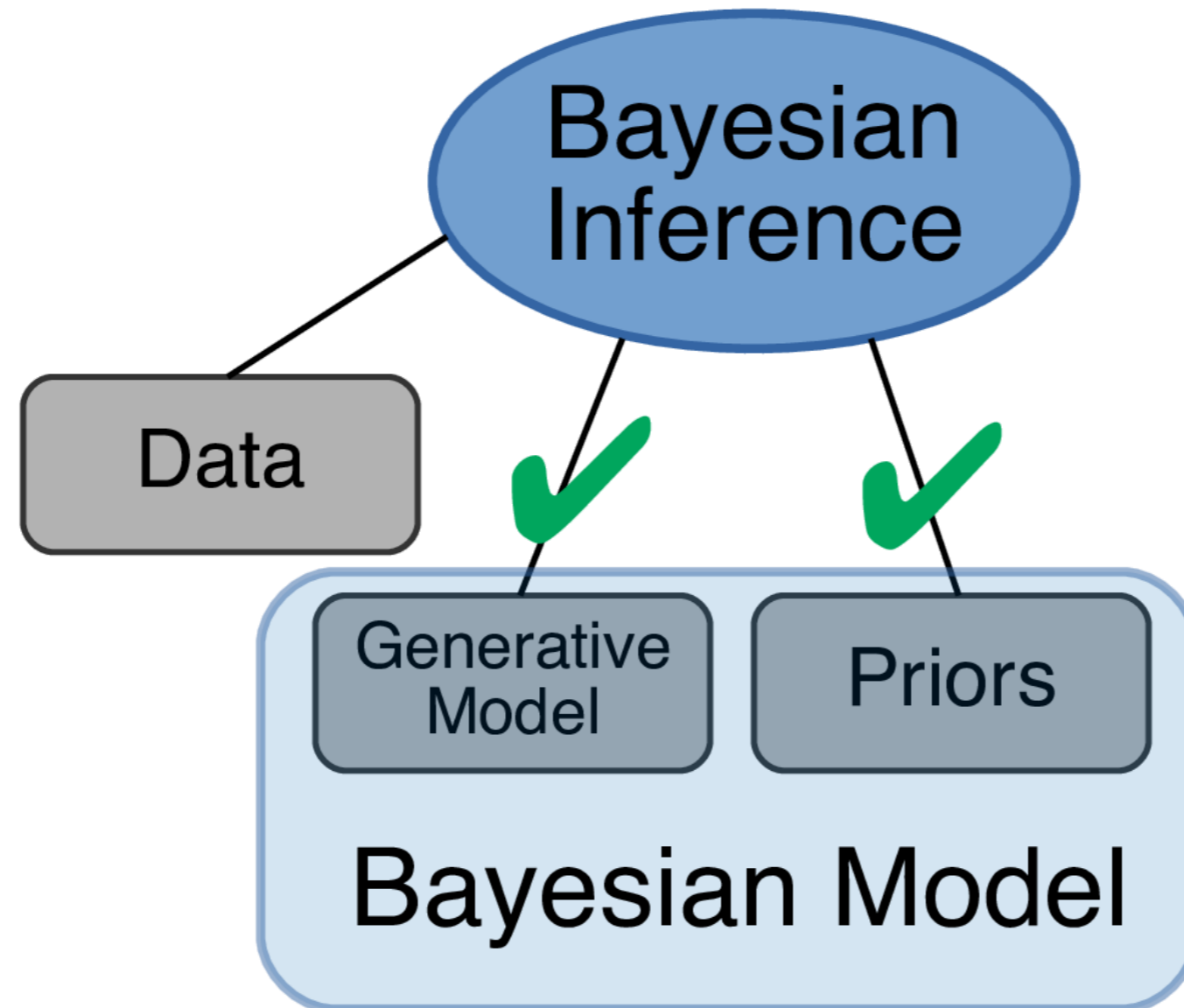
Bayesian models and conditioning

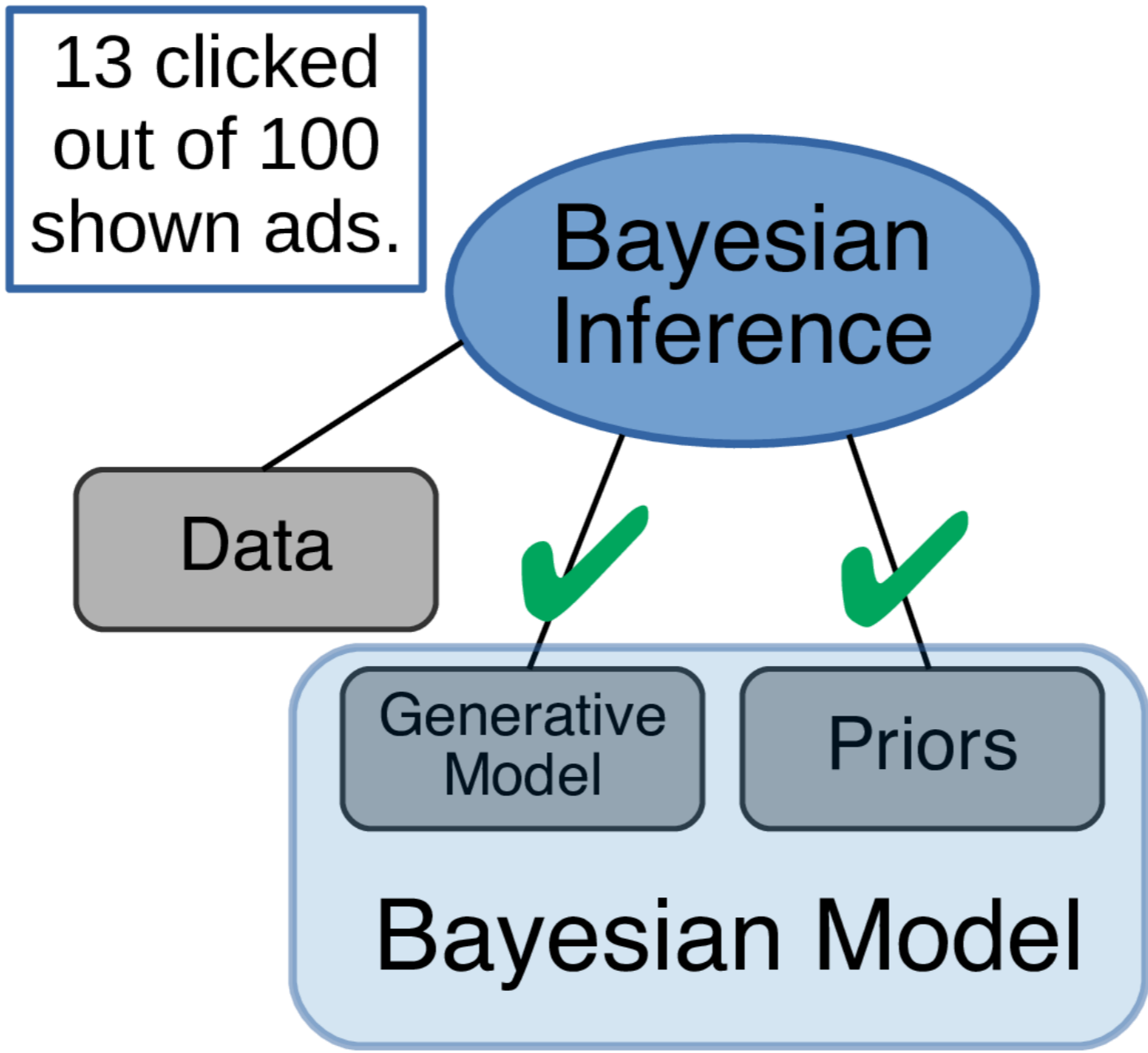
FUNDAMENTALS OF BAYESIAN DATA ANALYSIS IN R

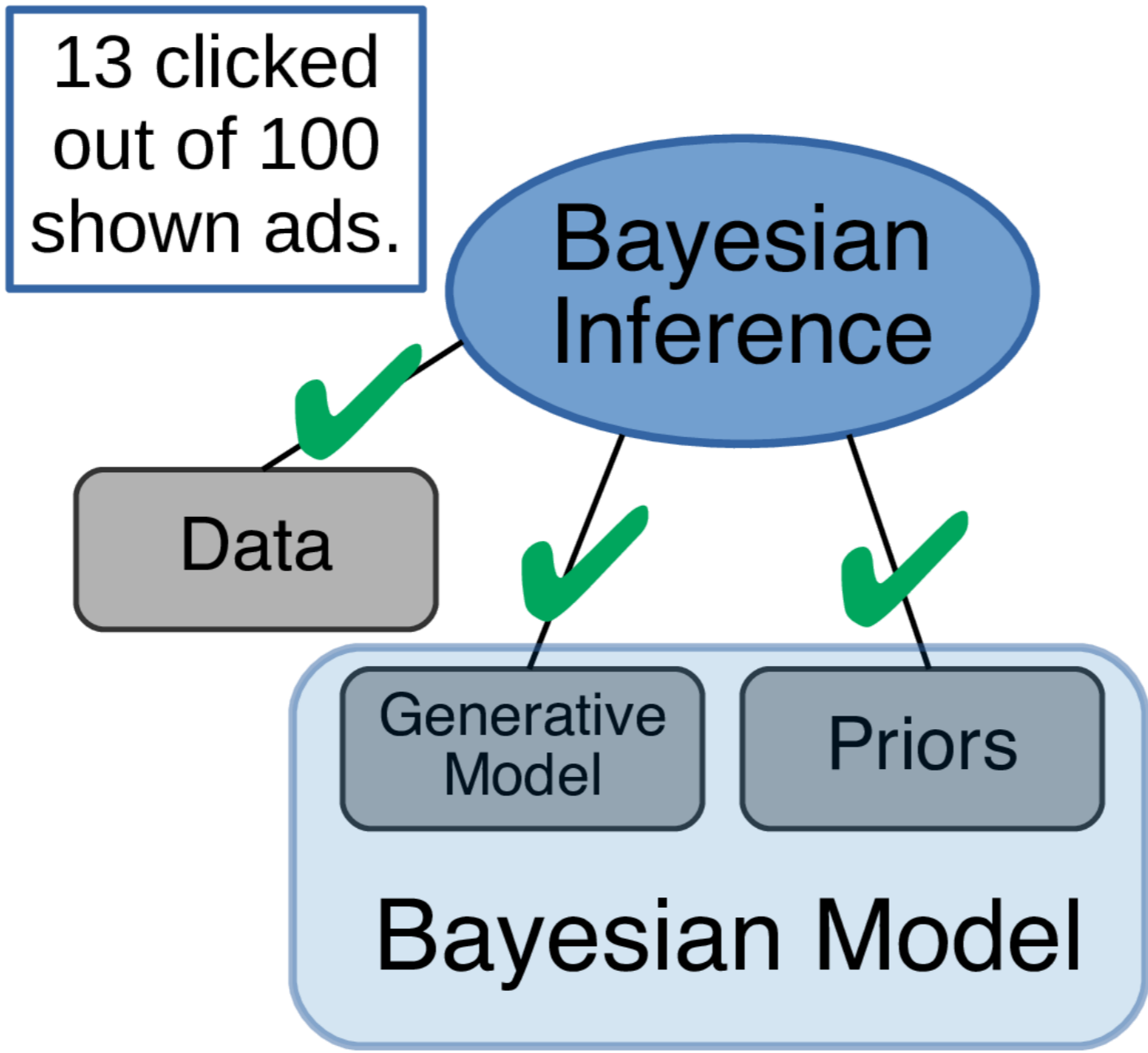


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Data Scientist









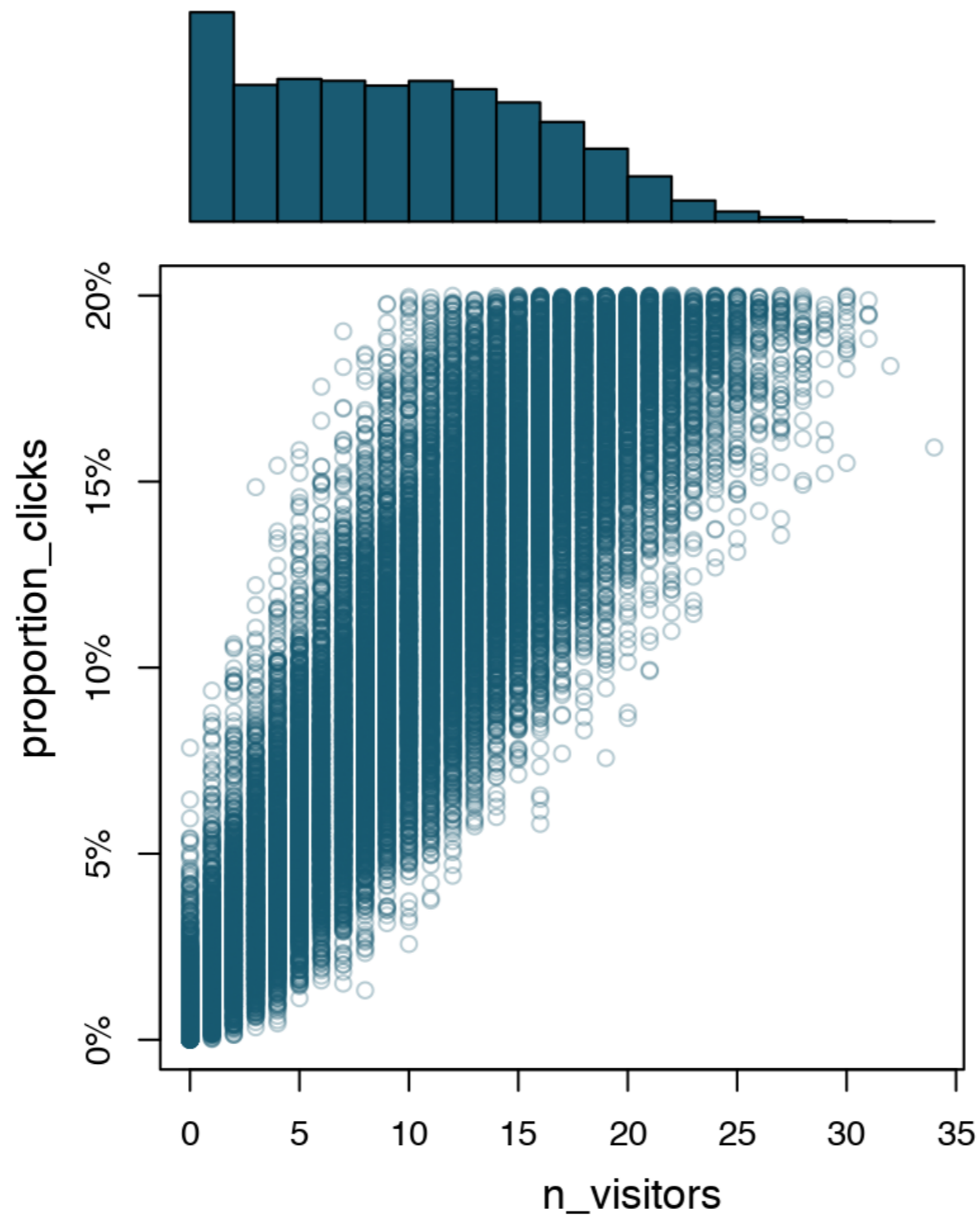
```
n_samples <- 1000000
n_ads_shown <- 100
proportion_clicks <- runif(n_samples, min = 0.0, max = 0.2)
n_visitors <- rbinom(n = n_samples, size = n_ads_shown,
                    prob = proportion_clicks)
```

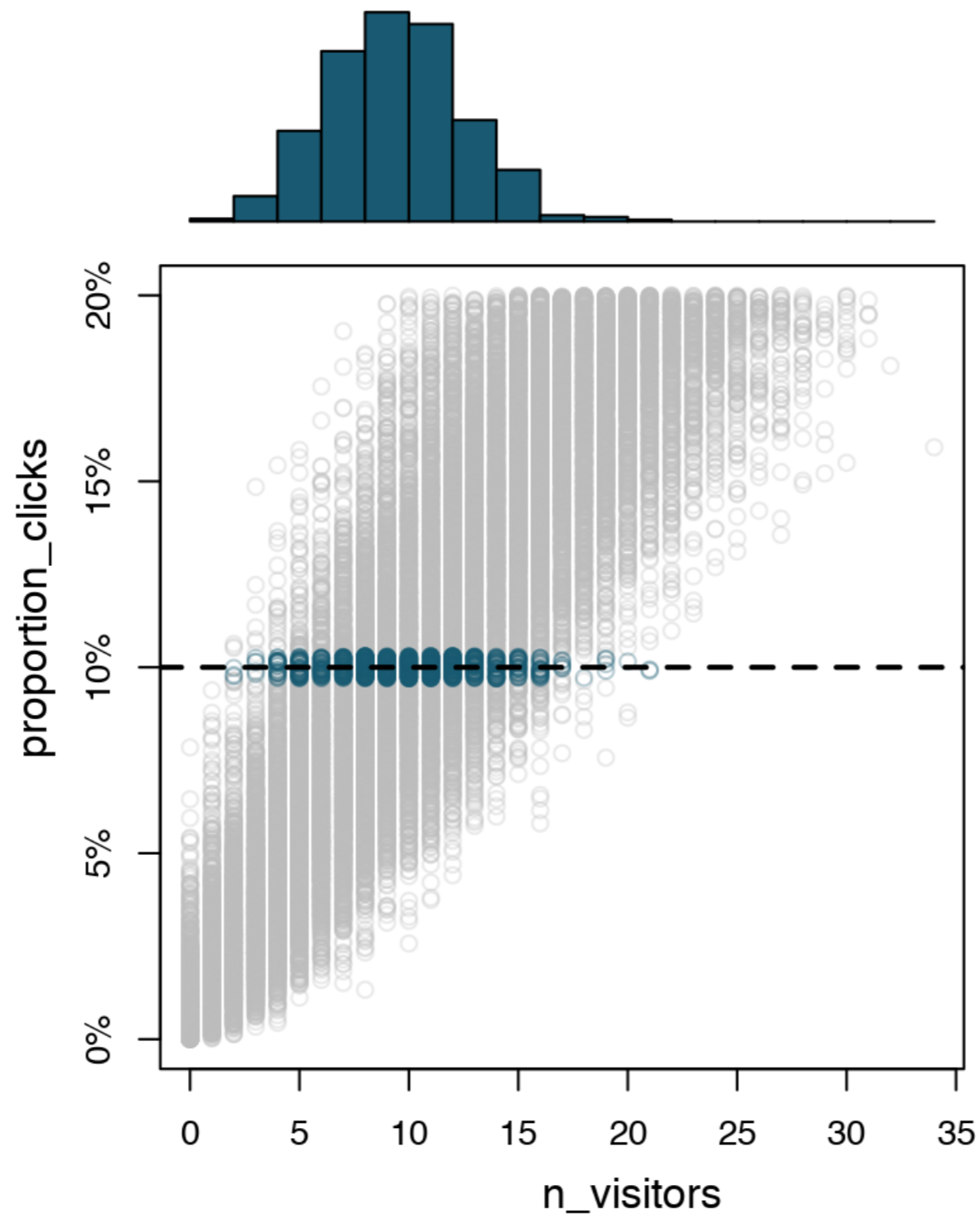


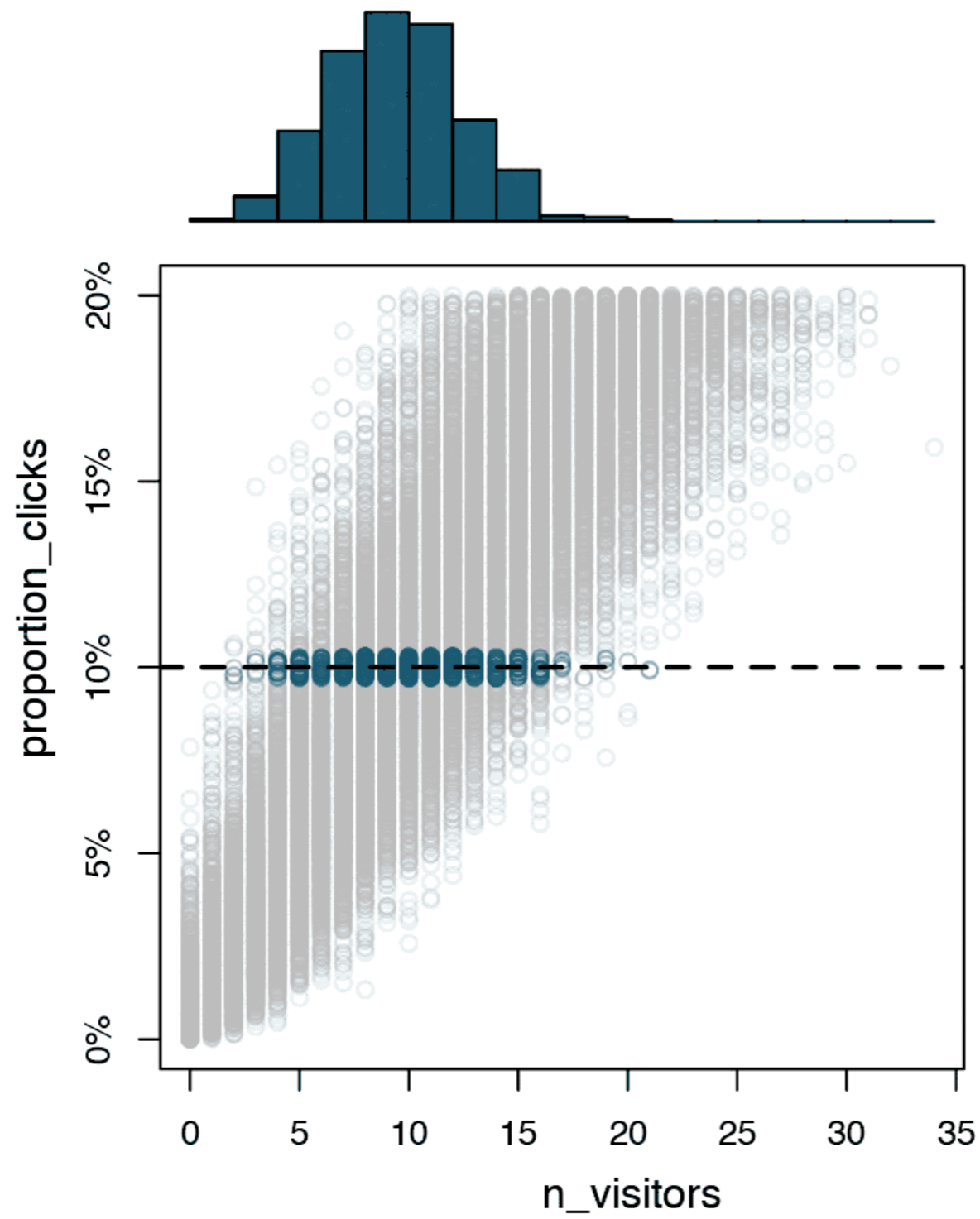
```
n_samples <- 100000
n_ads_shown <- 100
proportion_clicks <- runif(n_samples, min = 0.0, max = 0.2)
n_visitors <- rbinom(n = n_samples, size = n_ads_shown,
                    prob = proportion_clicks)
prior <- data.frame(proportion_clicks, n_visitors)
```

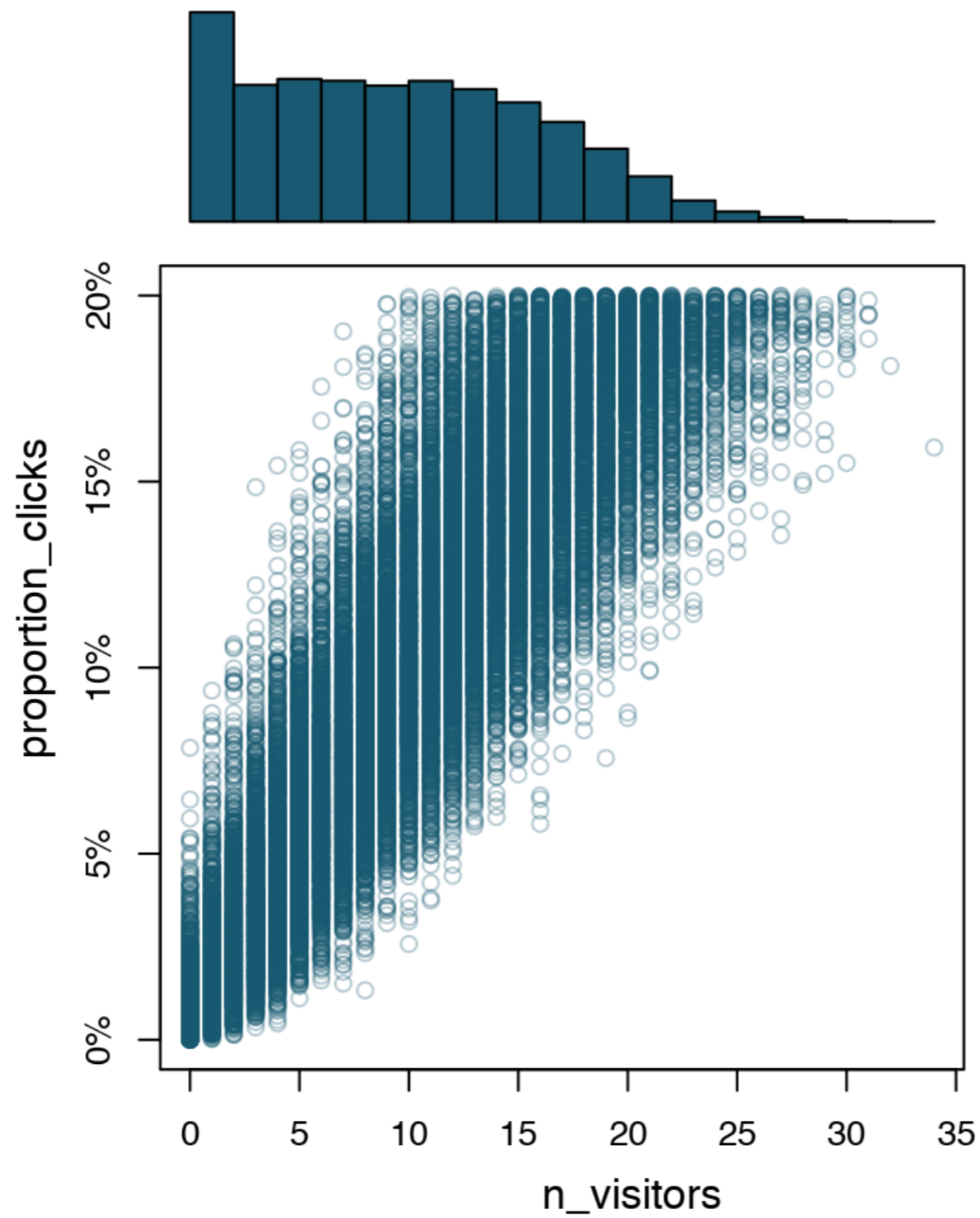
```
prior
```

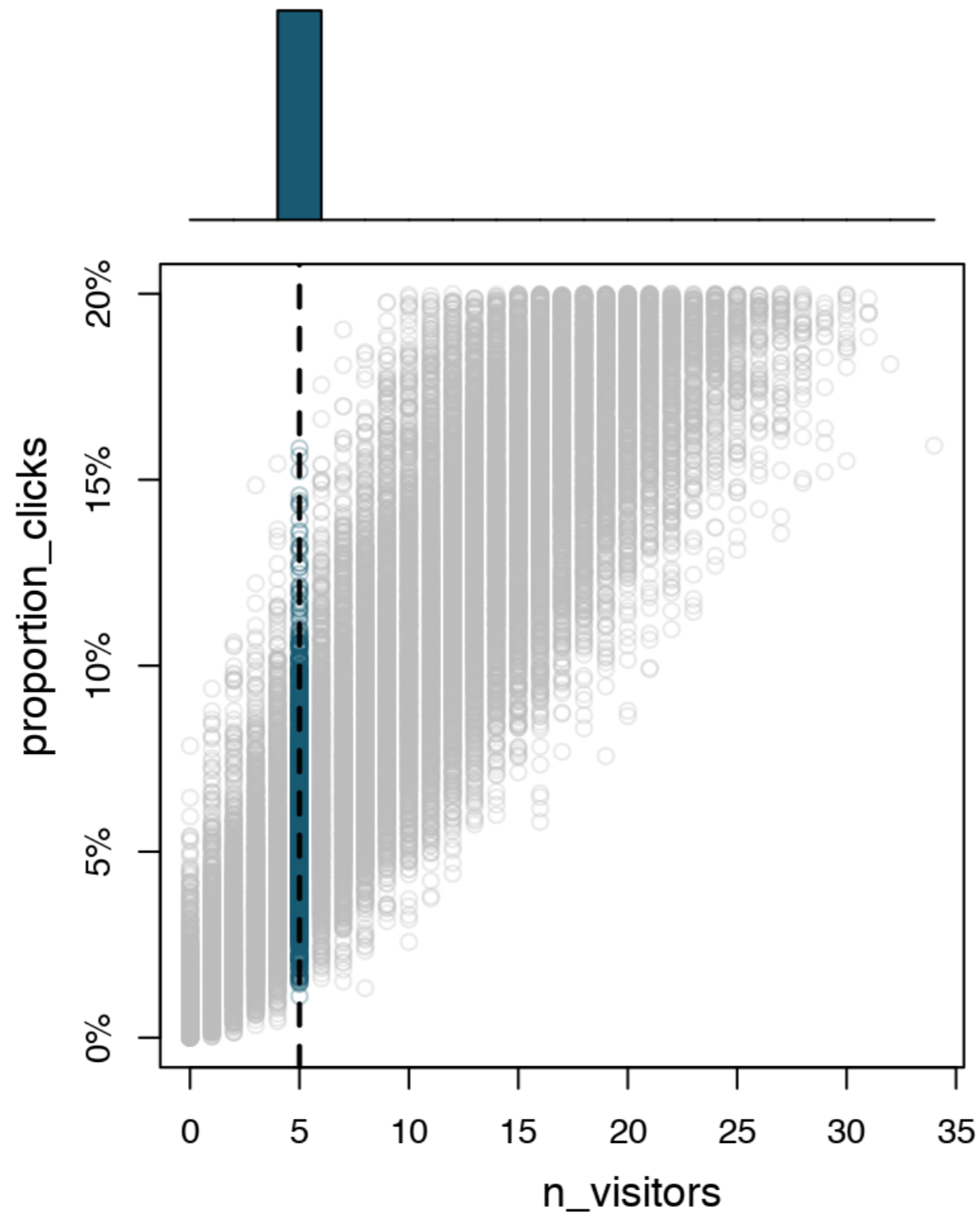
```
  proportion_clicks n_visitors
1          0.12         10
2          0.04          3
3          0.11         14
4          0.15         14
5          0.15         12
6          0.16         13
7          0.04          6
8          0.04          3
9          0.09         10
10         0.04          3
11         0.08          8
12         0.13         12
13         0.02          3
14         0.18         19
15         0.04          5
16         0.10         10
...         ...         ...
```

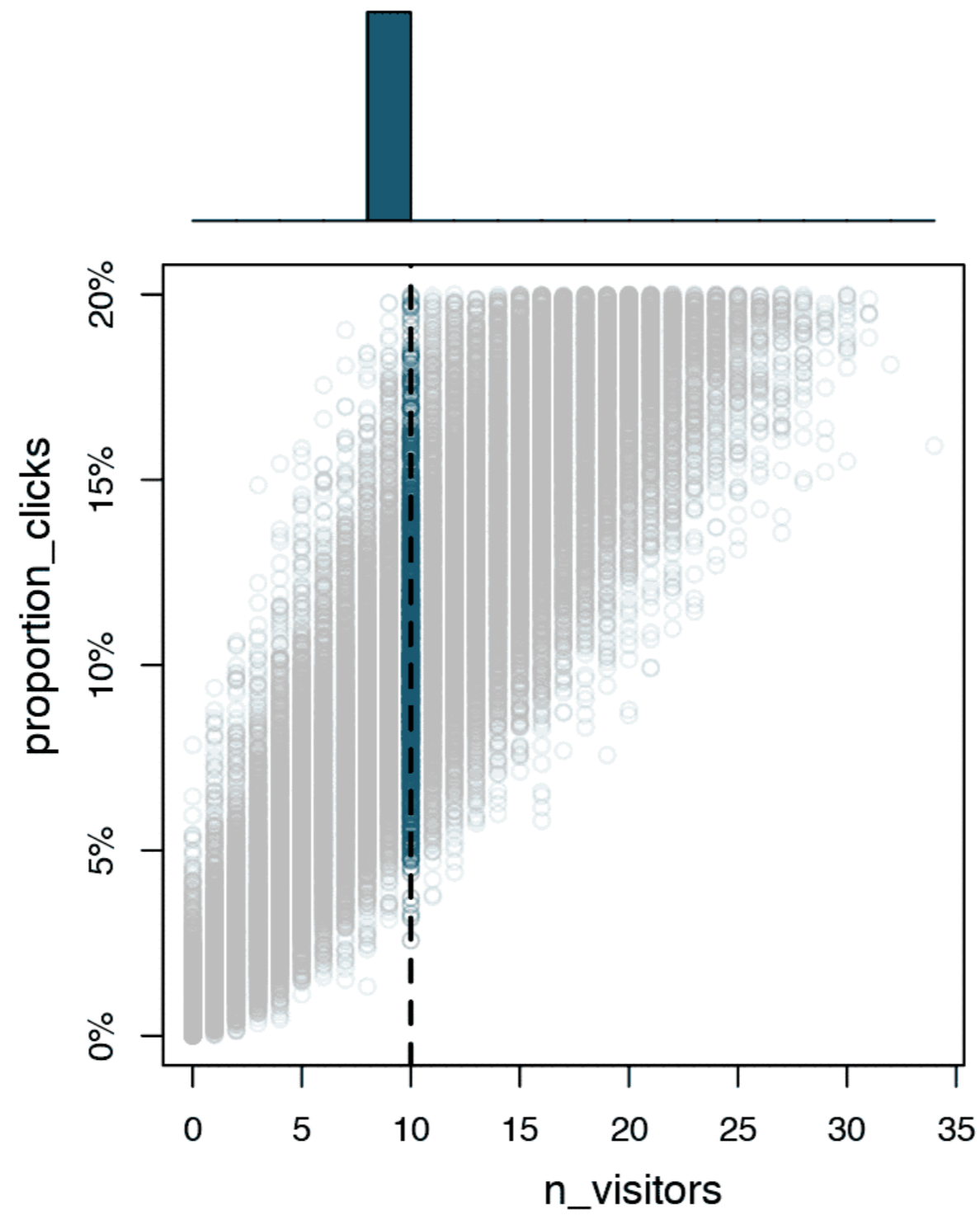












The essence of Bayesian inference

Bayesian inference is conditioning on data, in order to learn about parameter values.

Try some Bayesian inference yourself!

FUNDAMENTALS OF BAYESIAN DATA ANALYSIS IN R

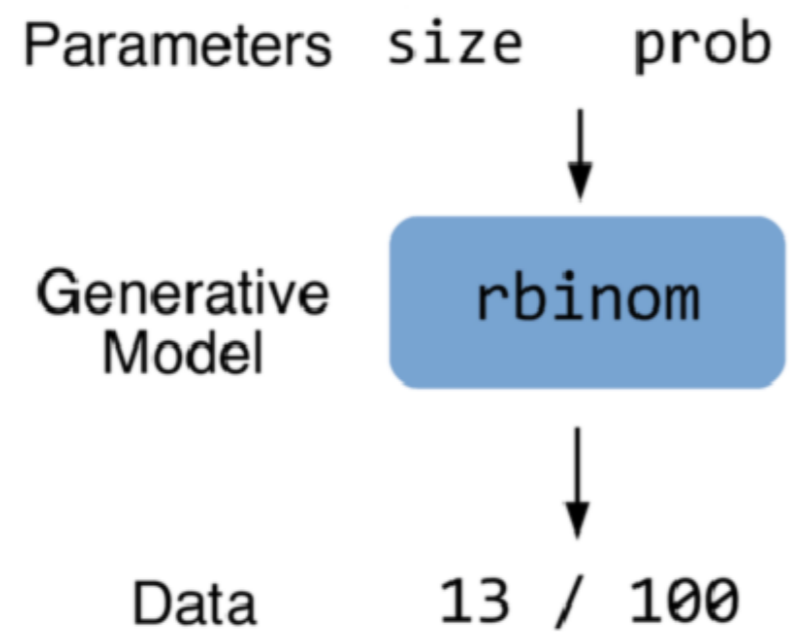
Bayesian inference, again!

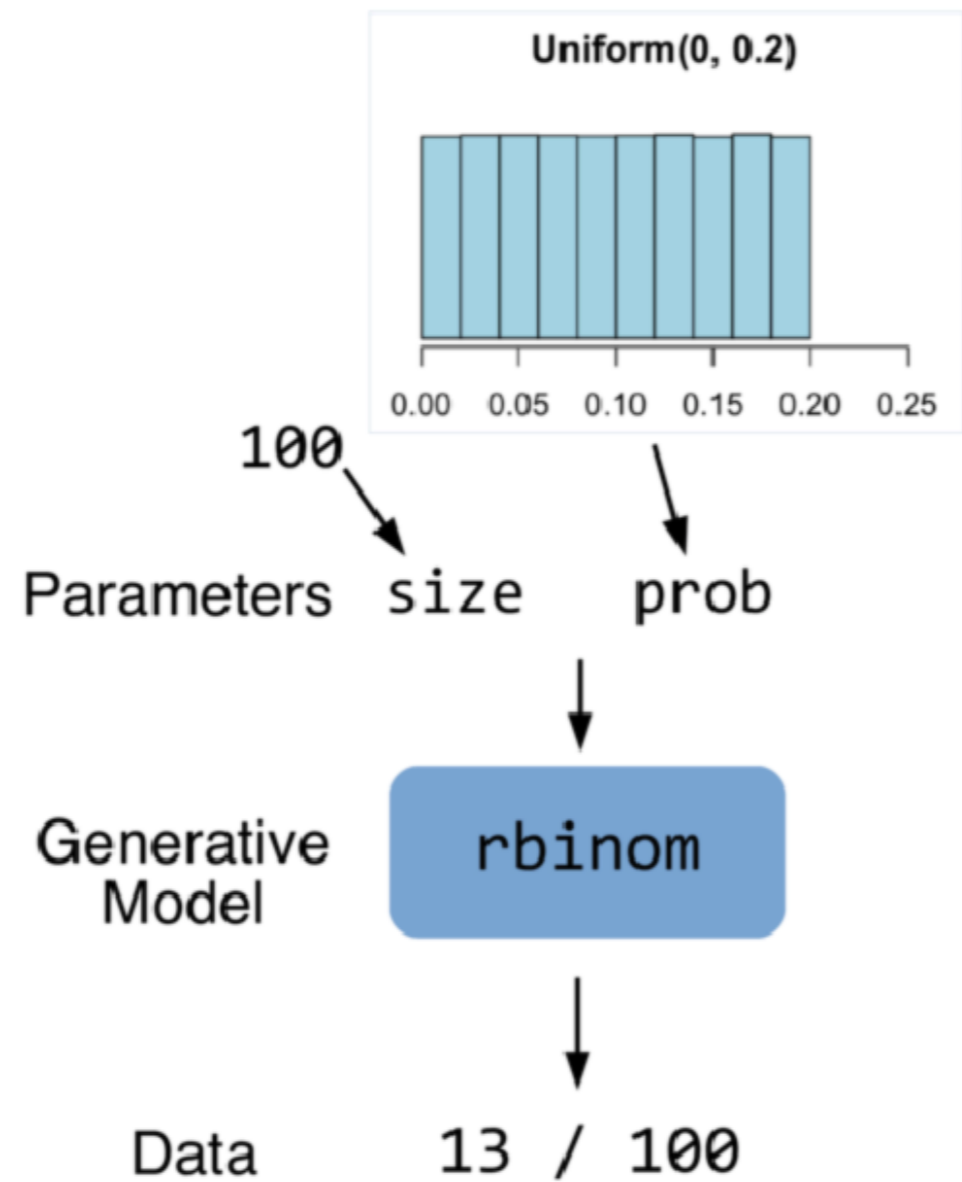
FUNDAMENTALS OF BAYESIAN DATA ANALYSIS IN R

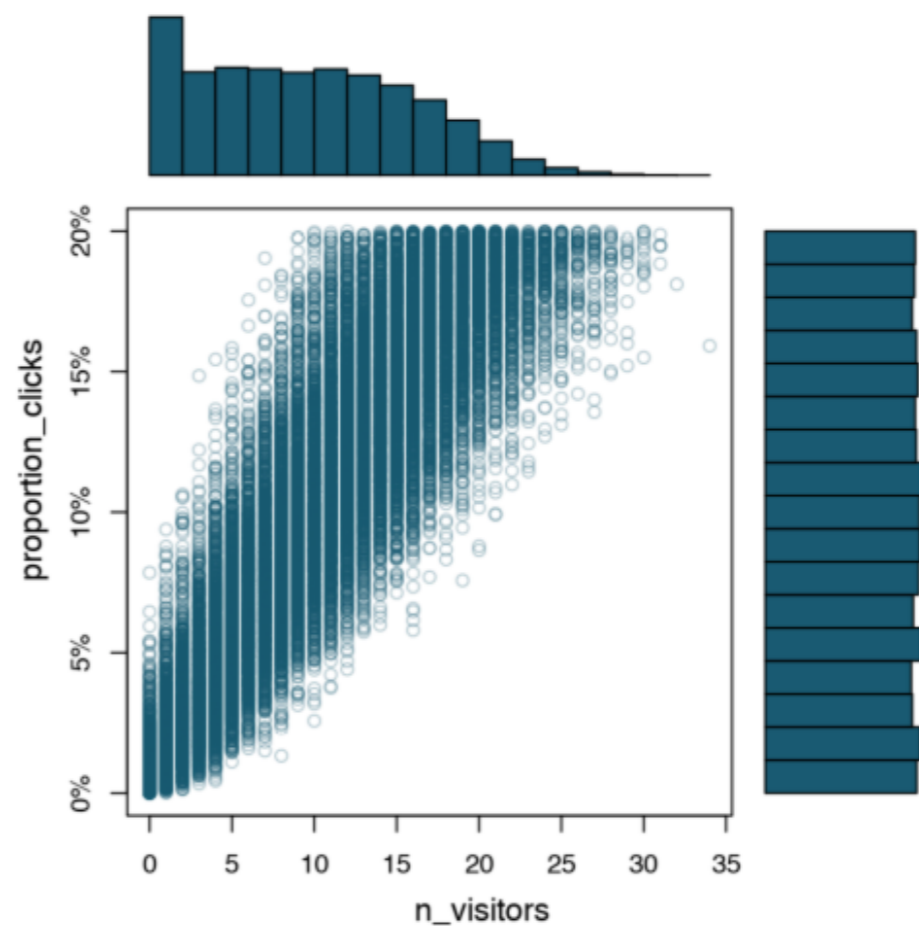
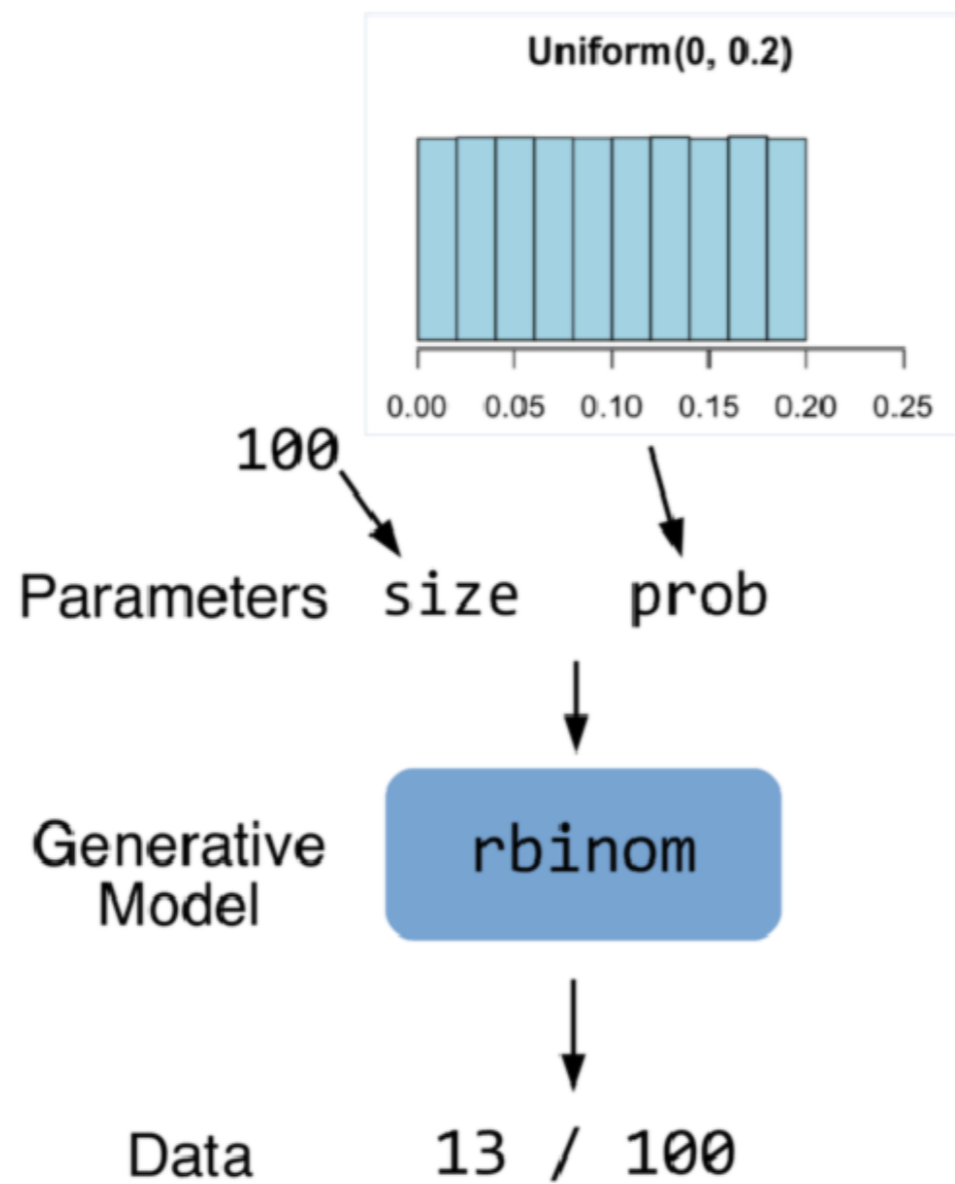


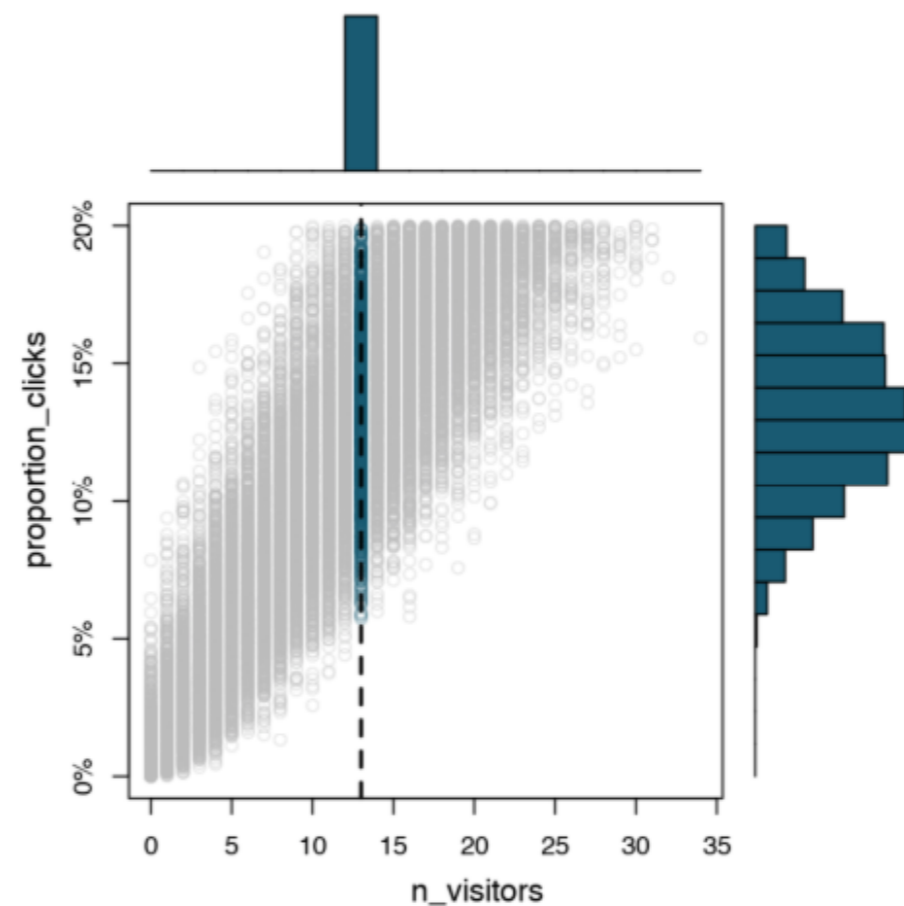
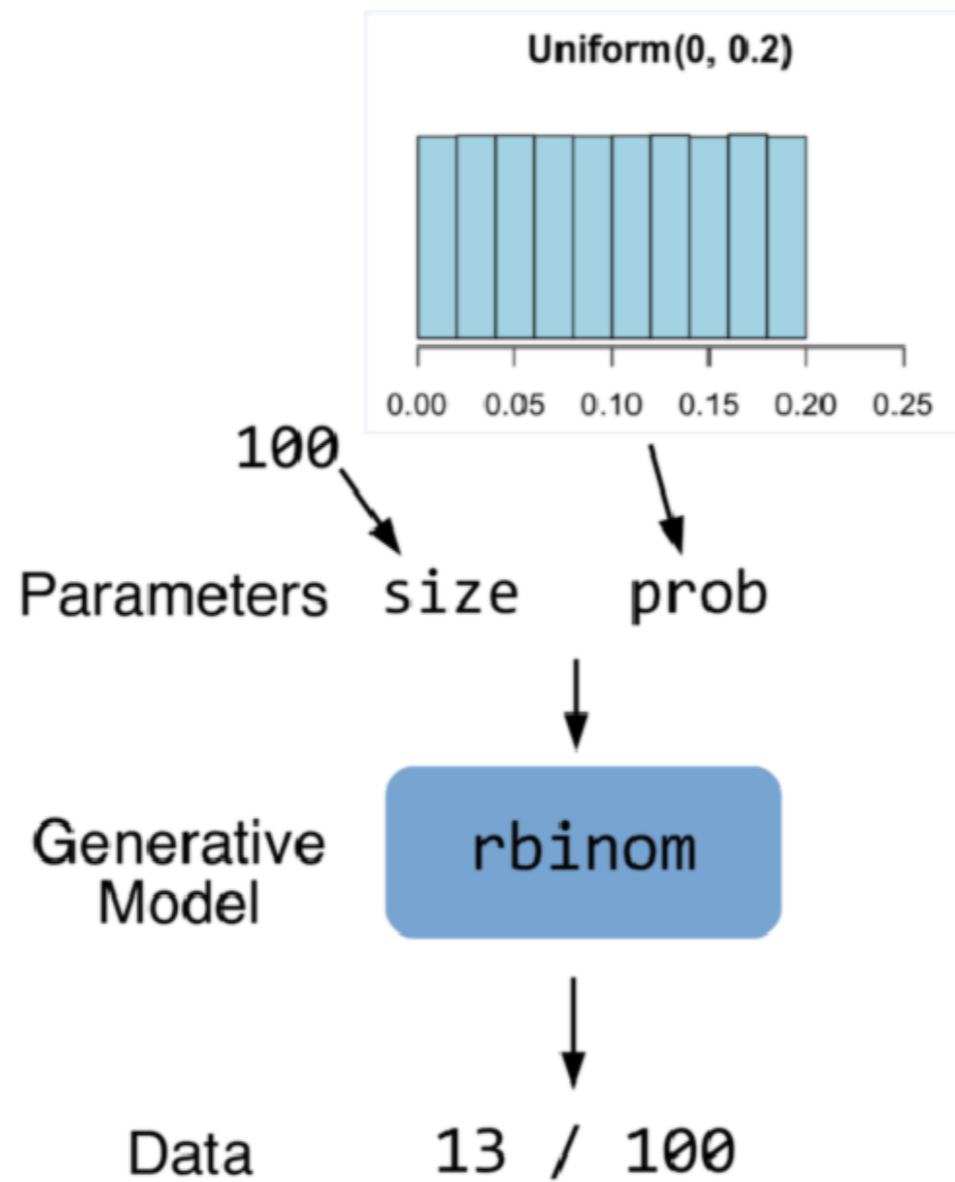
Rasmus Bååth
Data Scientist

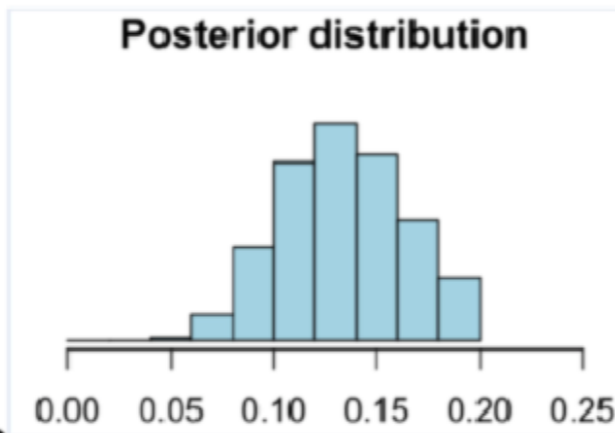
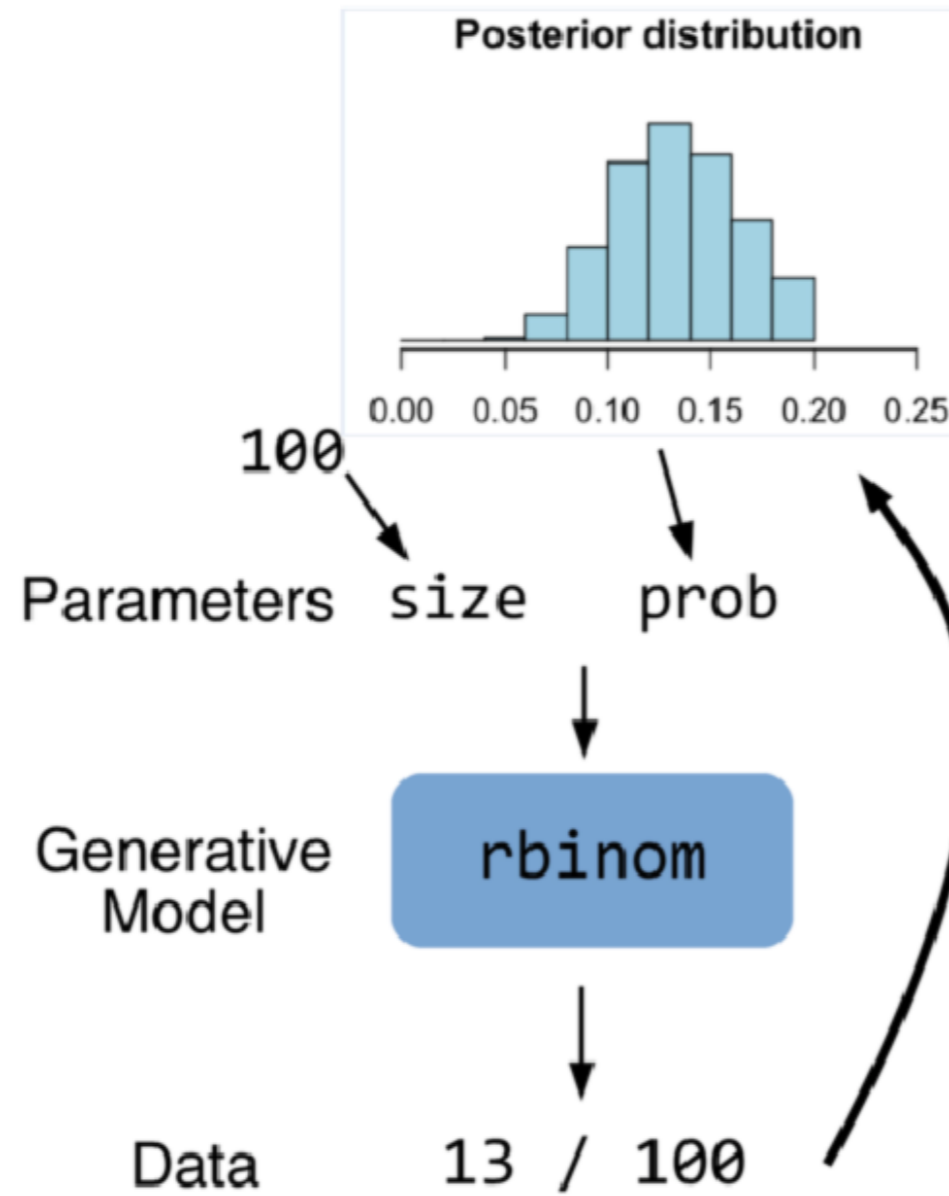
```
prop_success <- 0.1
size <- 100
# Simulating data
data <- c()
for(i in 1:size) {
  data[i] <-
    runif(1, min = 0, max = 1) <
      prop_success
}
data <- as.numeric(data)
```



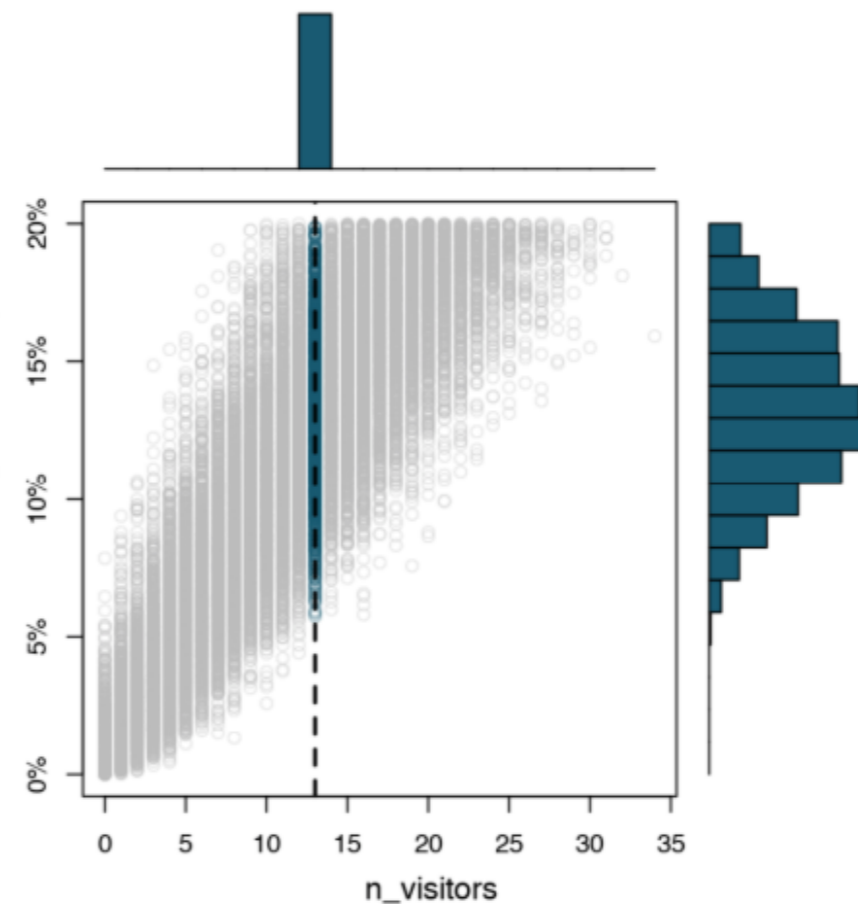


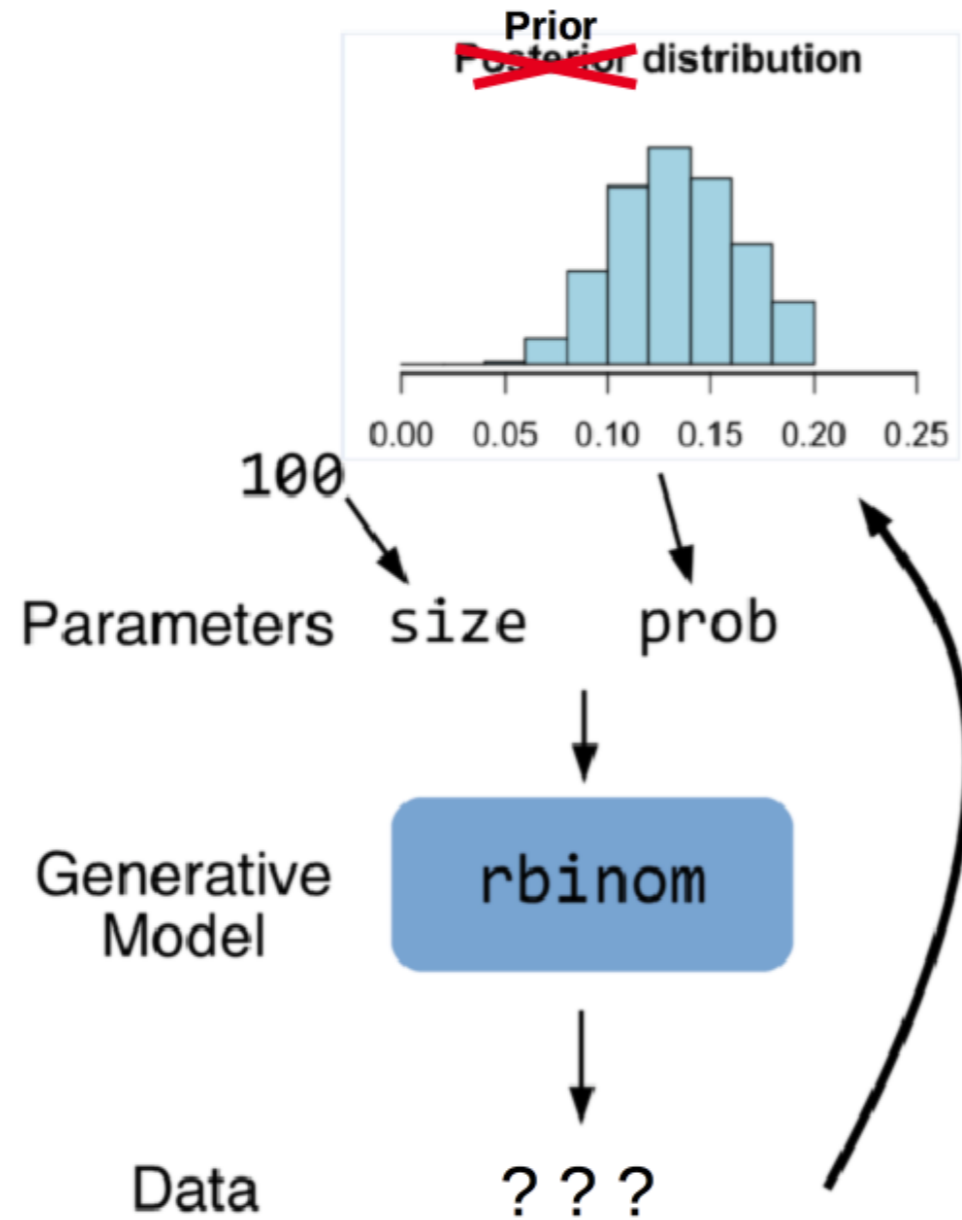




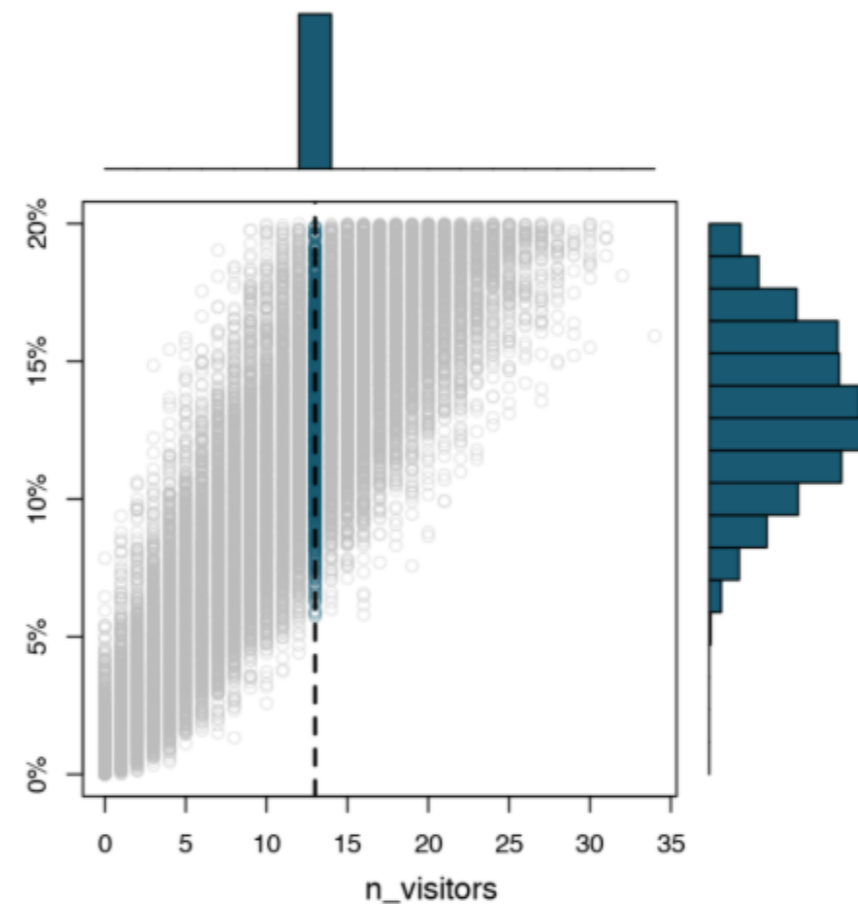


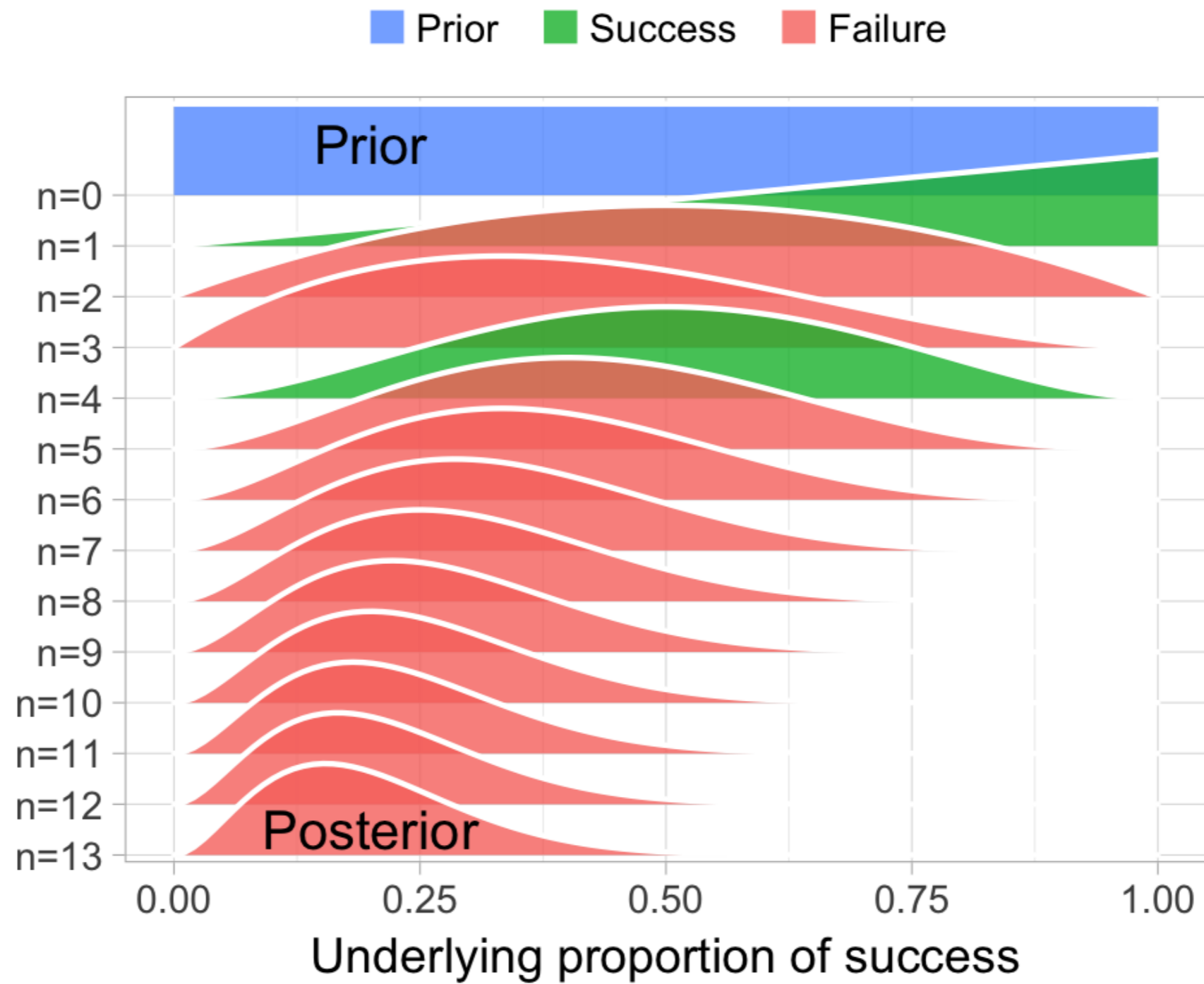
Bayesian Inference

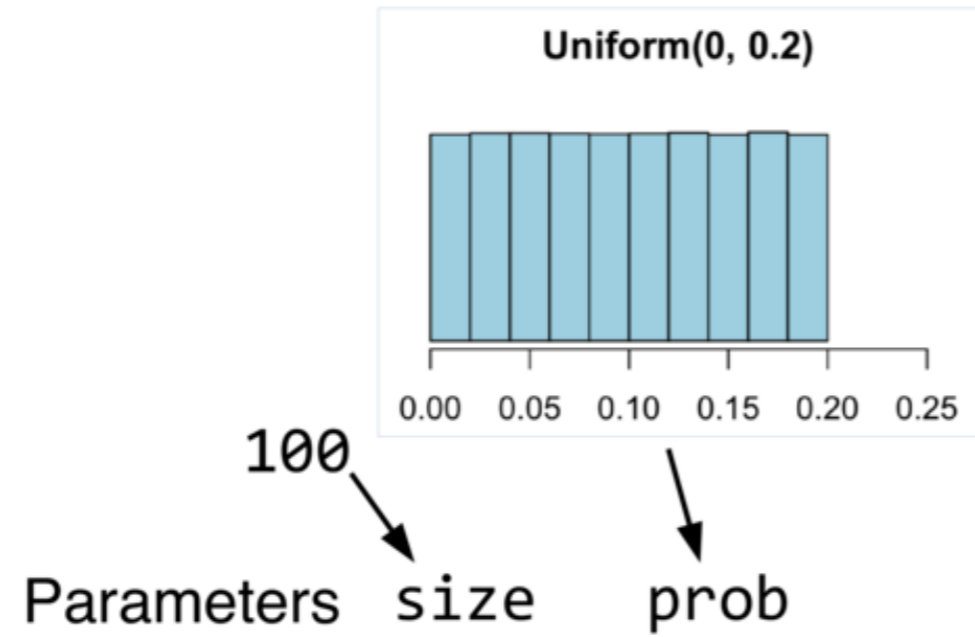


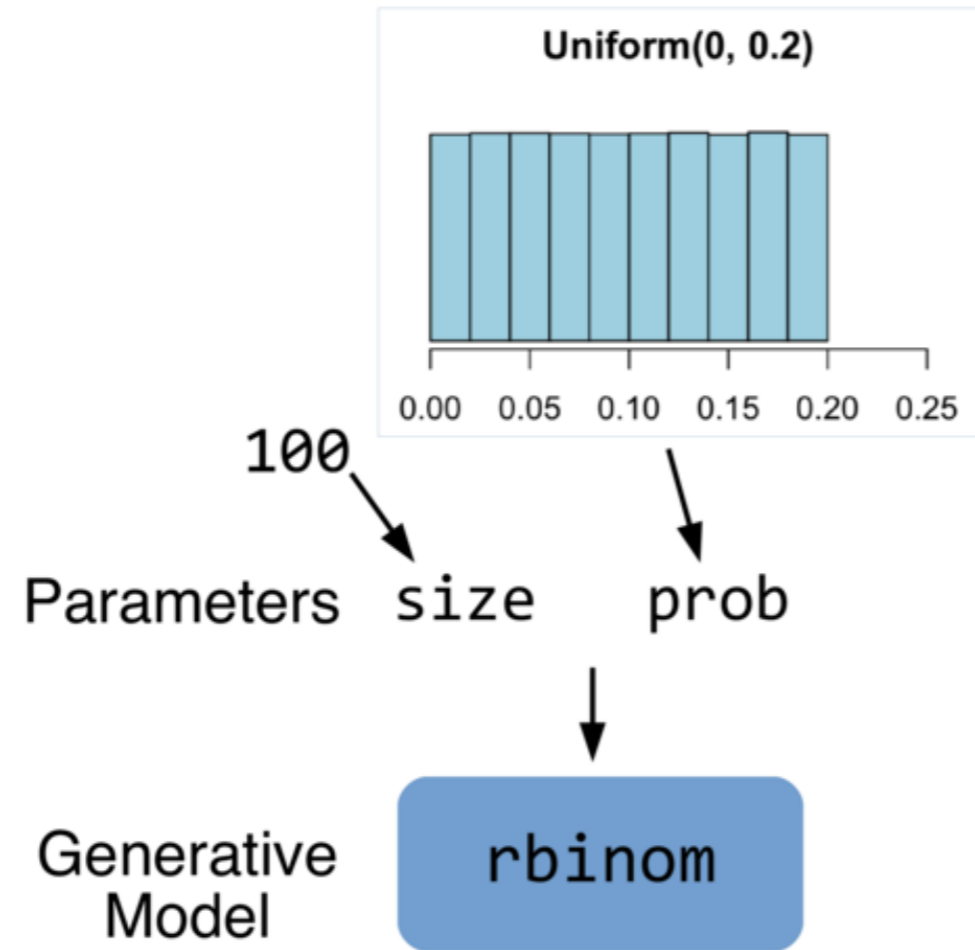


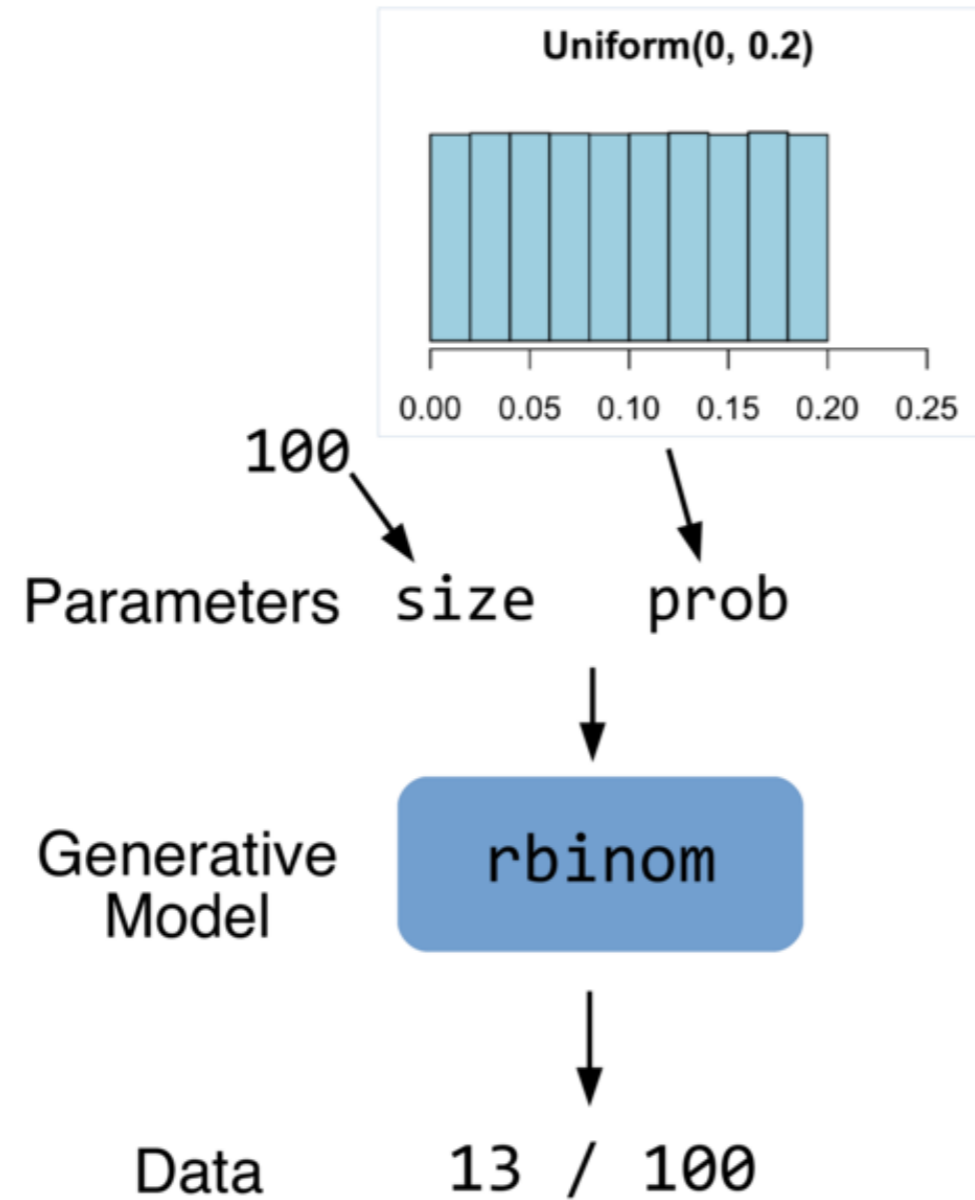
Bayesian Inference

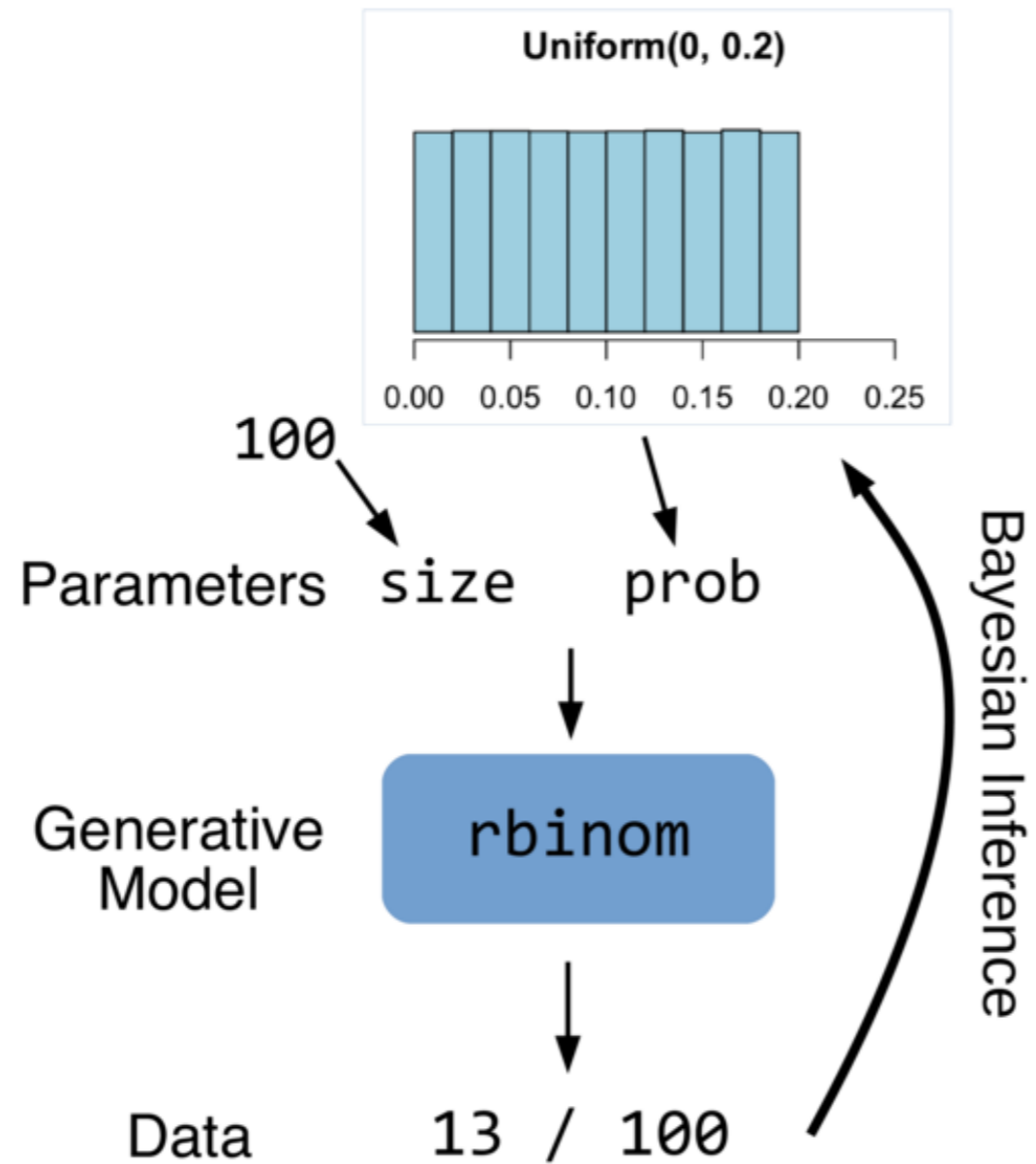


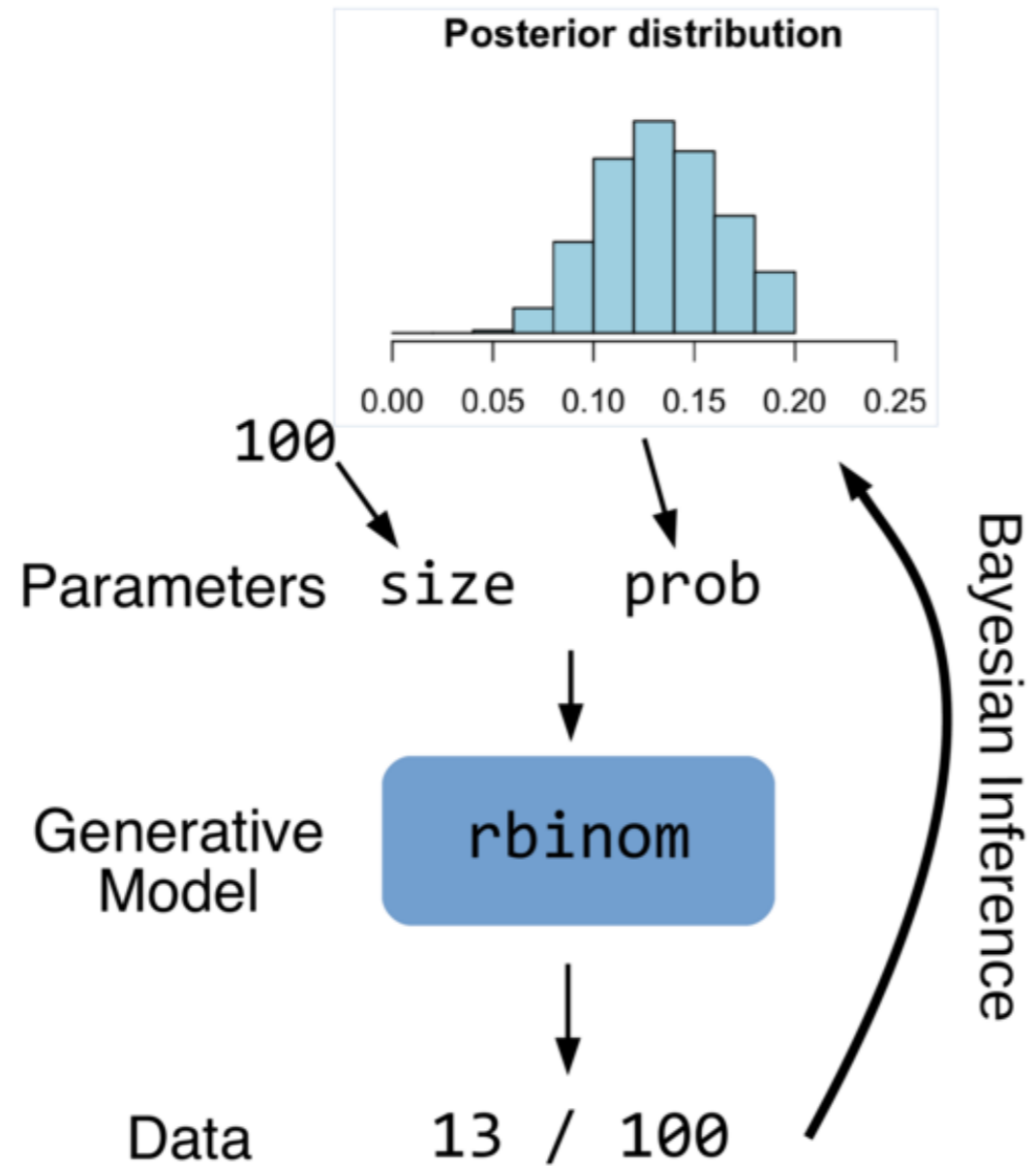


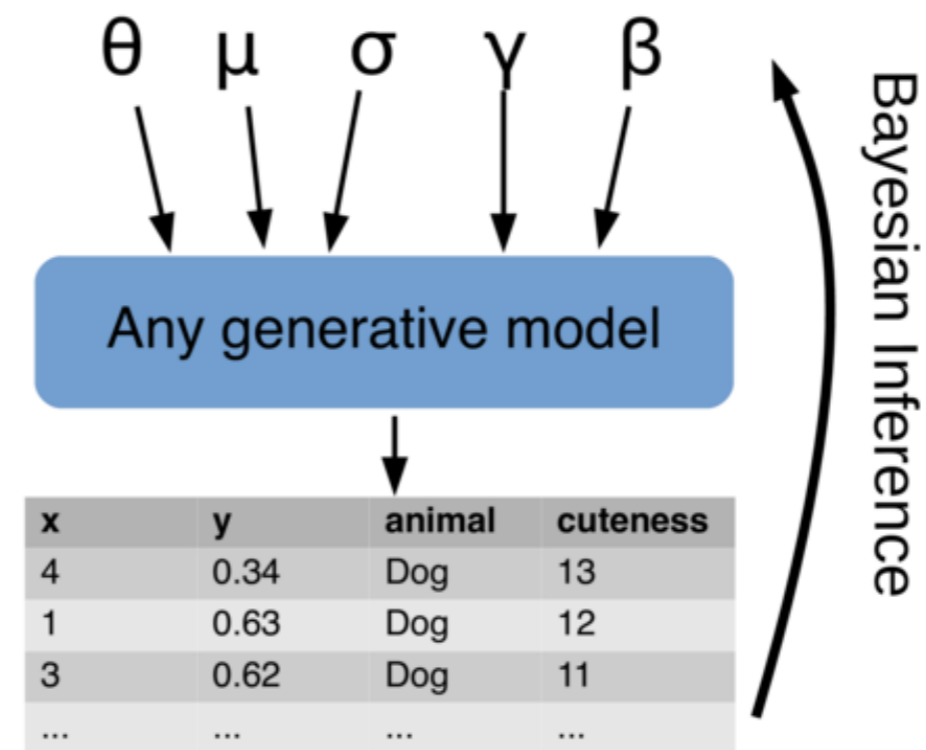
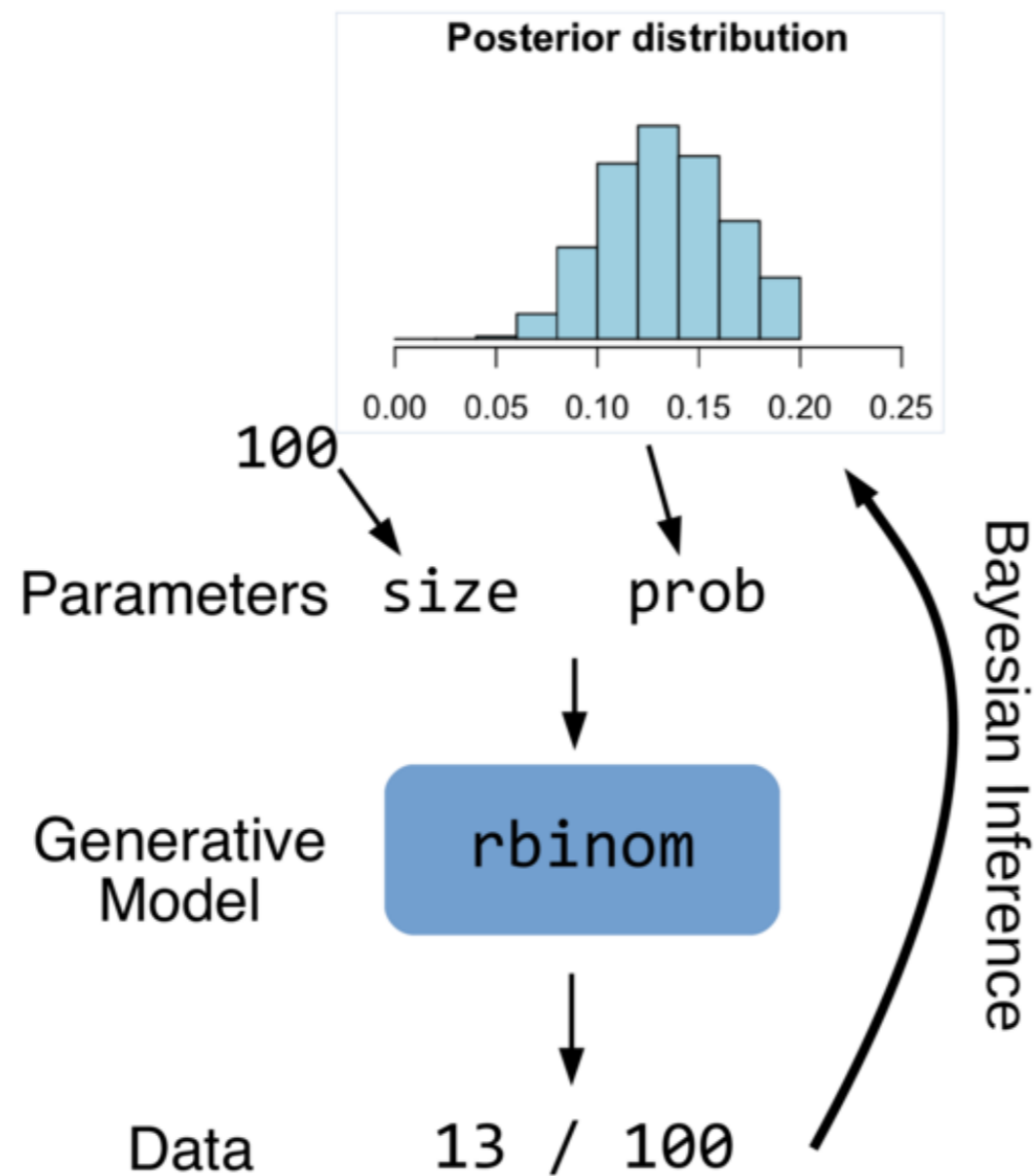


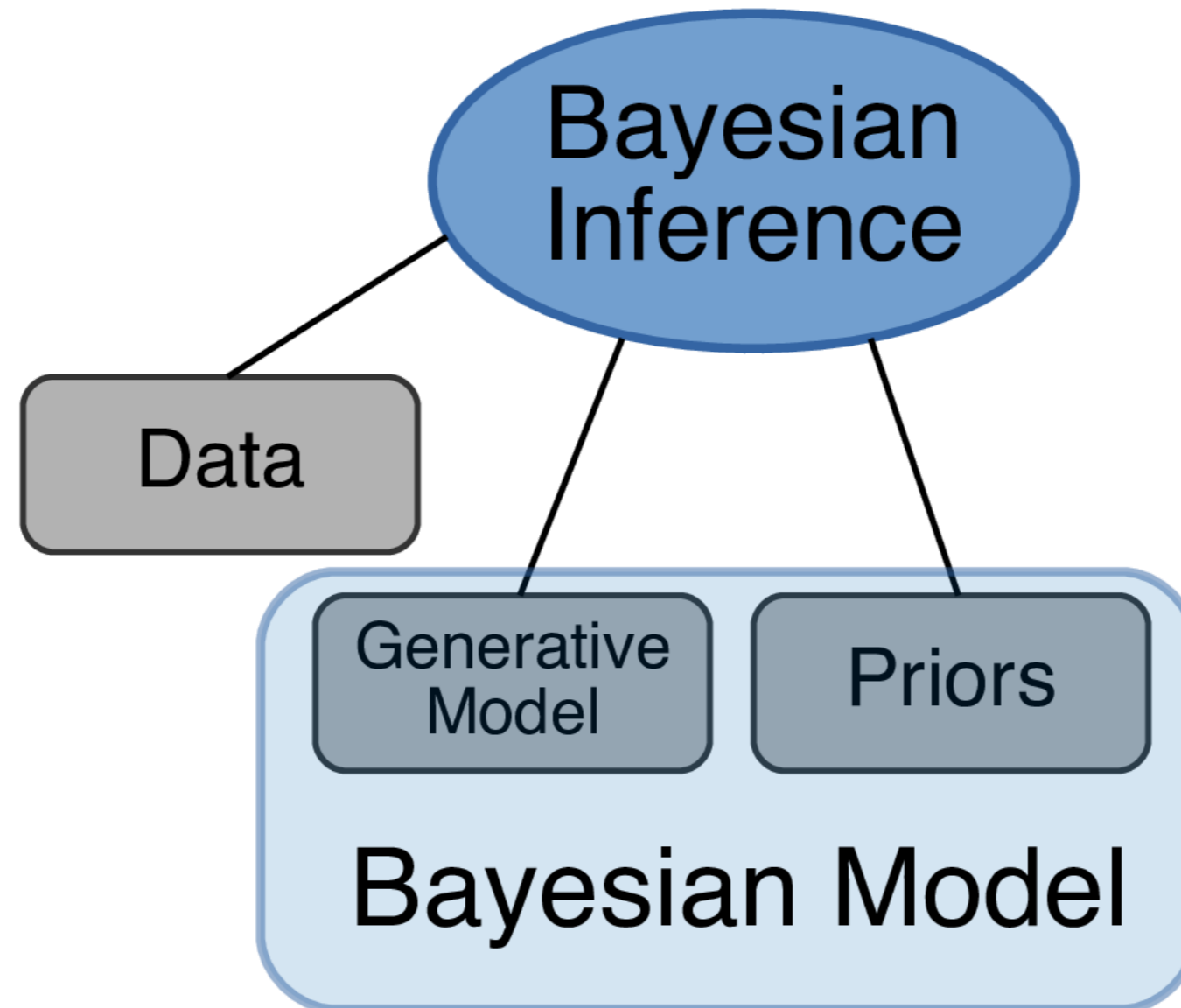


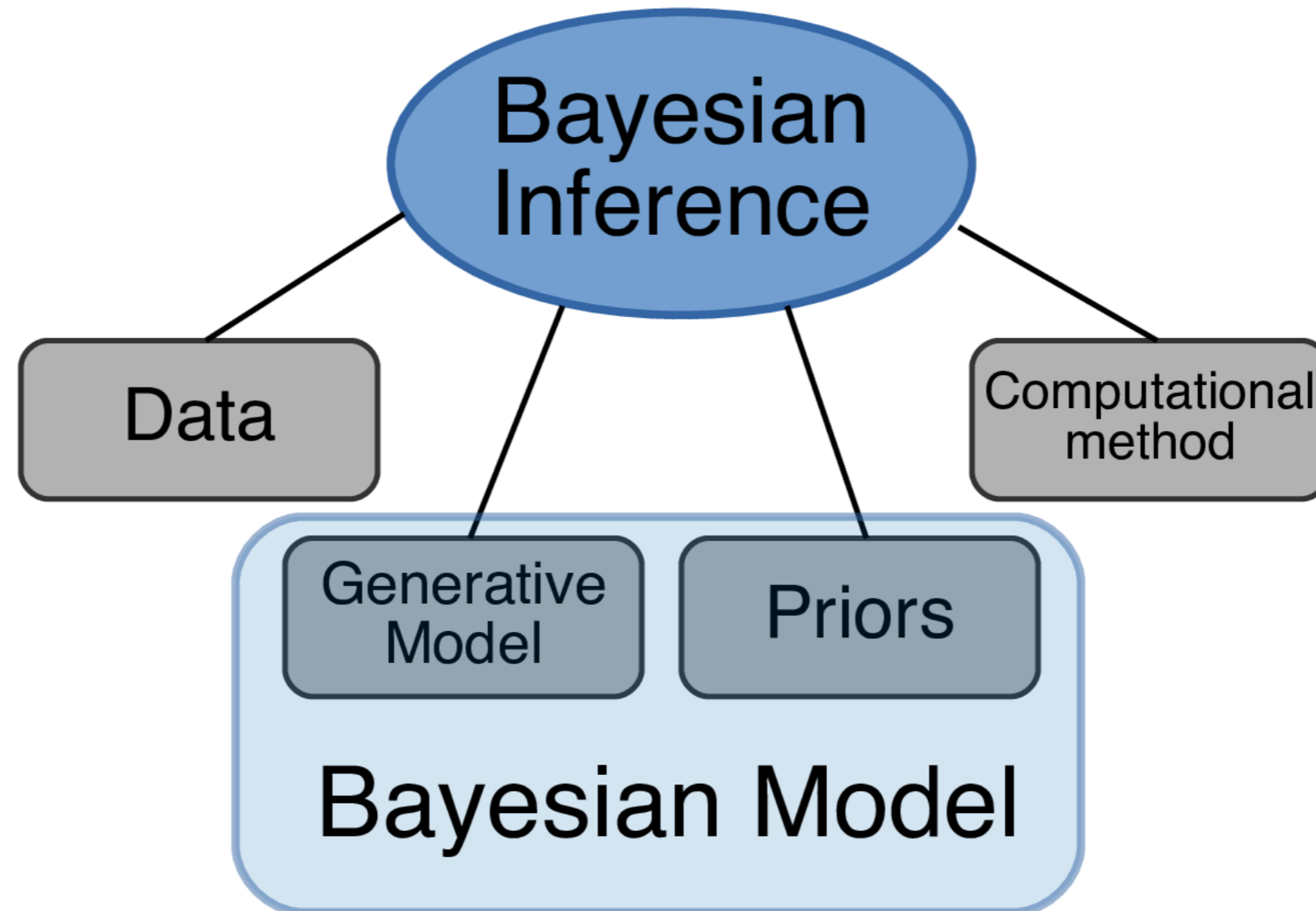












Next up: Why use Bayes?

FUNDAMENTALS OF BAYESIAN DATA ANALYSIS IN R