

Maloney_HW1

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Homework - Chapters 2 and 3

```
library(tidyverse)
```

```
## -- Attaching packages ----- tidyverse 1.2.1
## v ggplot2 3.1.0      v purrr  0.2.5
## v tibble  2.0.1      v dplyr  0.7.8
## v tidyr   0.8.2      v stringr 1.3.1
## v readr   1.3.1      v forcats 0.3.0

## -- Conflicts ----- tidyverse_conflicts()
## x dplyr::filter() masks stats::filter()
## x dplyr::lag()    masks stats::lag()
```

```
library(rethinking)
```

```
## Loading required package: rstan
## Loading required package: StanHeaders
## rstan (Version 2.18.2, GitRev: 2e1f913d3ca3)
## For execution on a local, multicore CPU with excess RAM we recommend calling
## options(mc.cores = parallel::detectCores()).
## To avoid recompilation of unchanged Stan programs, we recommend calling
## rstan_options(auto_write = TRUE)

## For improved execution time, we recommend calling
## Sys.setenv(LOCAL_CPPFLAGS = '-march=native')
## although this causes Stan to throw an error on a few processors.

##
## Attaching package: 'rstan'

## The following object is masked from 'package:tidyr':
##
##      extract

## Loading required package: parallel
## rethinking (Version 1.59)

##
## Attaching package: 'rethinking'

## The following object is masked from 'package:purrr':
##
##      map
```

```
library(tidybayes)
```

```
## NOTE: As of tidybayes version 1.0, several functions, arguments, and output column names
##      have undergone significant name changes in order to adopt a unified naming scheme.
```

```
## See help('tidybayes-deprecated') for more information.
```

Chapter 2 Homework

Medium Problems

2M1

```
#WWW

#define grid
p_grid <- seq(from = 0, to = 1, length.out = 100)

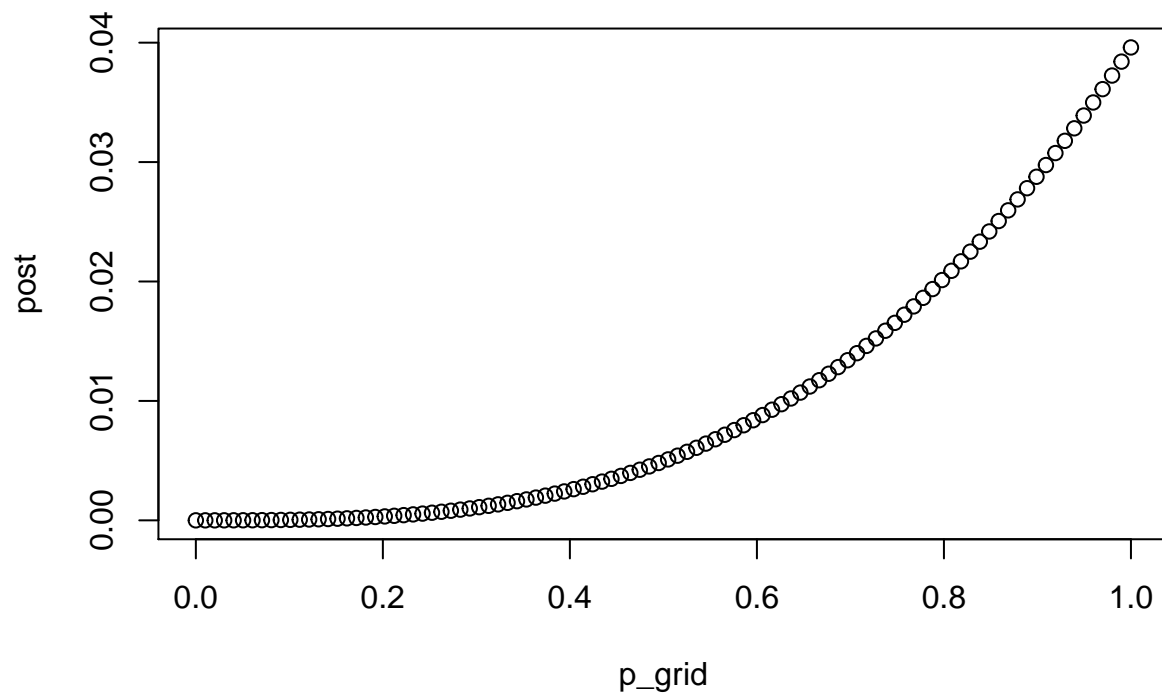
#define prior
prior <- rep(1, 100)
#prior <- exp(-5*abs(p_grid - 0.5))

#compute likelihood at each value in the grid
lh <- dbinom(3, size = 3, prob = p_grid)

#compute product of lh & prior
upost <- lh * prior

#standardize post
post <- upost/sum(upost)

plot(p_grid, post, type = "b")
```



```

#WWWL
#define grid
p_grid <- seq(from = 0, to = 1, length.out = 100)

#define prior
prior <- rep(1, 100)
#prior <- exp(-5*abs(p_grid - 0.5))

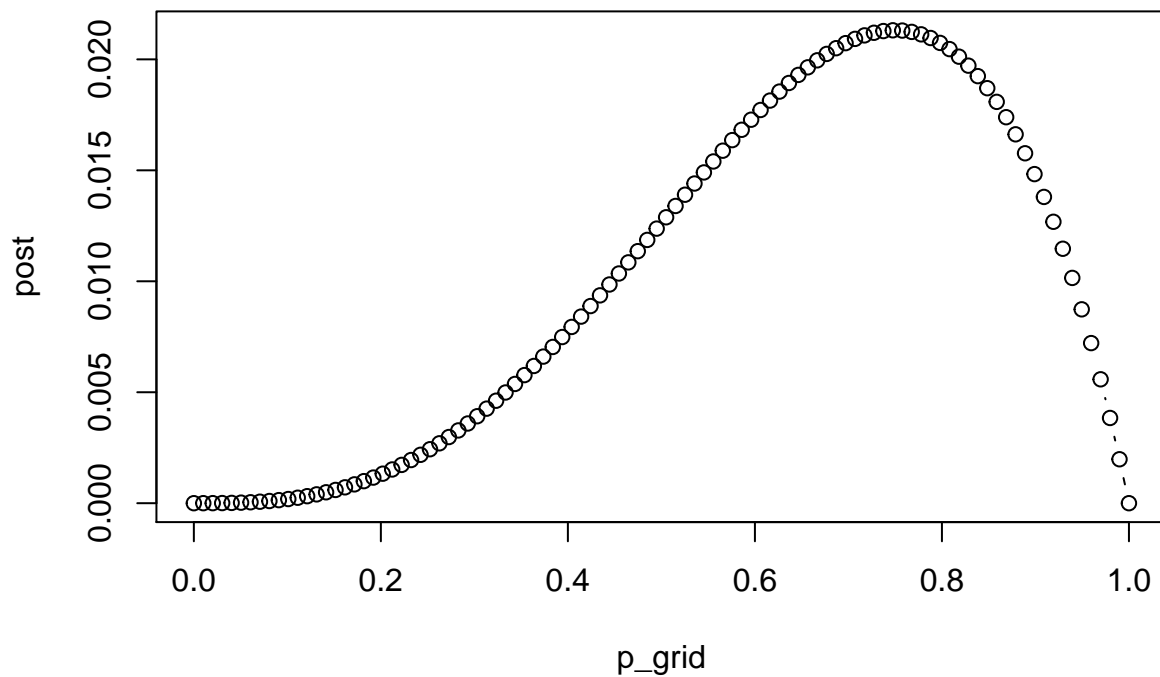
#compute likelihood at each value in the grid
lh <- dbinom(3, size = 4, prob = p_grid)

#compute product of lh & prior
upost <- lh * prior

#standardize post
post <- upost/sum(upost)

plot(p_grid, post, type = "b")

```



```

#LWWLWWW
#define grid
p_grid <- seq(from = 0, to = 1, length.out = 100)

#define prior
prior <- rep(1, 100)
#prior <- exp(-5*abs(p_grid - 0.5))

```

```

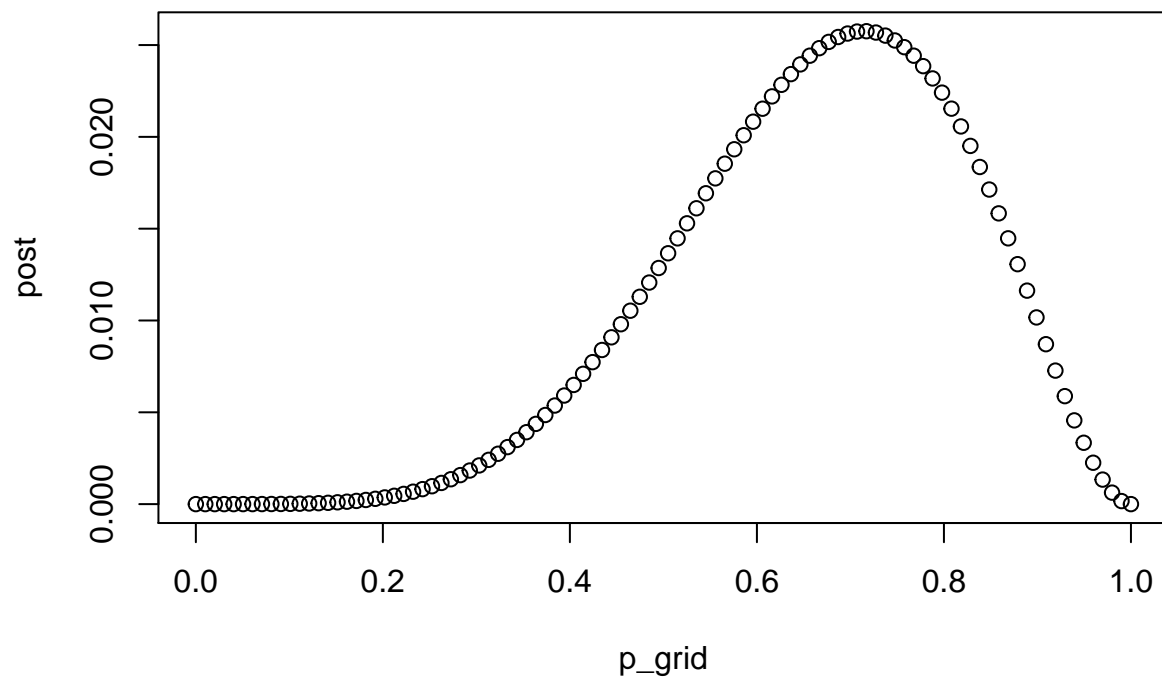
#compute likelihood at each value in the grid
lh <- dbinom(5, size = 7, prob = p_grid)

#compute product of lh & prior
upost <- lh * prior

#standardize post
post <- upost/sum(upost)

plot(p_grid, post, type = "b")

```



2M2

#WWW

```

#define grid
p_grid <- seq(from = 0, to = 1, length.out = 100)

#define prior
prior <- ifelse(p_grid < 0.5, 0, 2)

#compute likelihood at each value in the grid
lh <- dbinom(3, size = 3, prob = p_grid)

#compute product of lh & prior
upost <- lh * prior

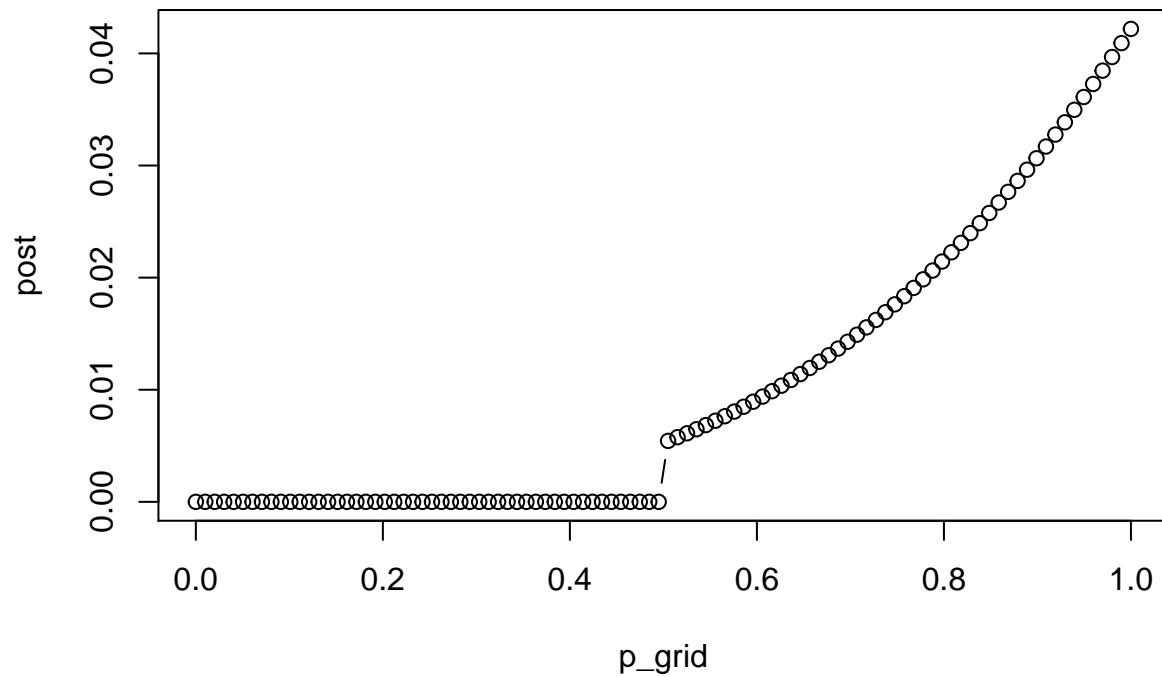
```

```

#standardize post
post <- upost/sum(upost)

plot(p_grid, post, type = "b")

```



```

#WWWL
#define grid
p_grid <- seq(from = 0, to = 1, length.out = 100)

#define prior
prior <- ifelse(p_grid < 0.5, 0, 2)

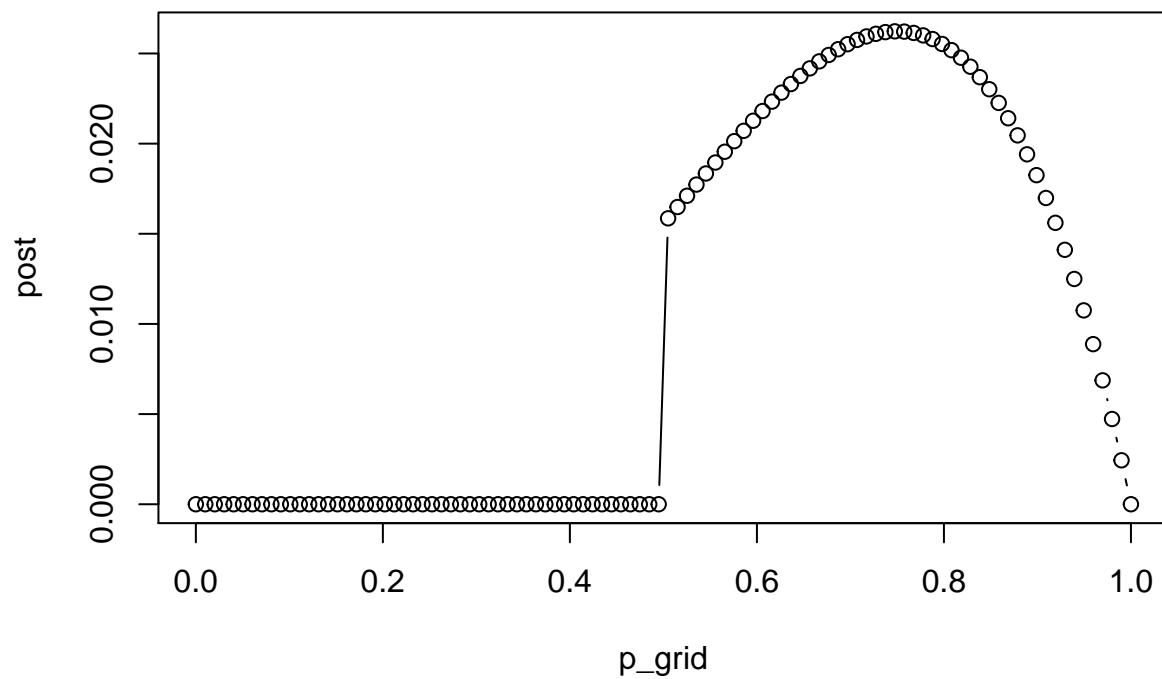
#compute likelihood at each value in the grid
lh <- dbinom(3, size = 4, prob = p_grid)

#compute product of lh & prior
upost <- lh * prior

#standardize post
post <- upost/sum(upost)

plot(p_grid, post, type = "b")

```



```
#LWWLWWW
#define grid
p_grid <- seq(from = 0, to = 1, length.out = 100)

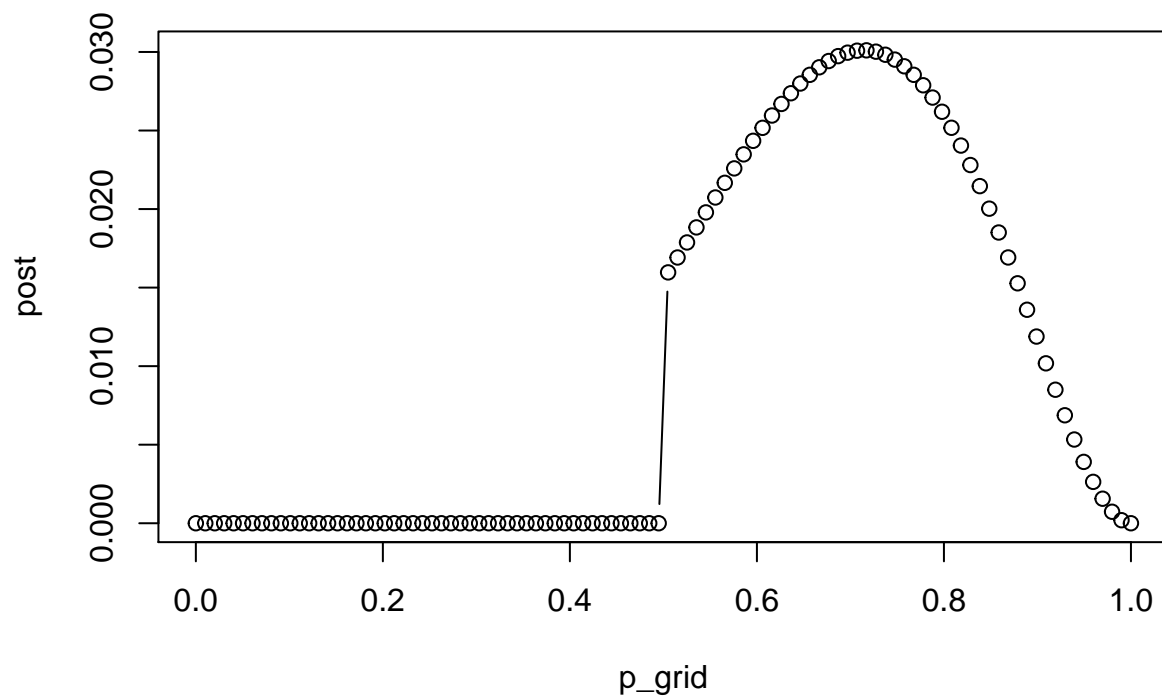
#define prior
prior <- ifelse(p_grid < 0.5, 0, 2)
#prior <- exp(-5*abs(p_grid - 0.5))

#compute likelihood at each value in the grid
lh <- dbinom(5, size = 7, prob = p_grid)

#compute product of lh & prior
upost <- lh * prior

#standardize post
post <- upost/sum(upost)

plot(p_grid, post, type = "b")
```



2M3

```
pwe <- 0.7
plm <- 1
lh <- 0.3/1
priorodds <- 1
postodds <- priorodds*lh
(post <- postodds/(postodds + 1))
```

```
## [1] 0.2307692
```

The probability of the globe was Earth, given seeing land is 0.2307692.

2M4

```
ww <- 0
bw <- 1
bb <- 2

(p <- bb/(bb + bw + ww))
```

```
## [1] 0.6666667
```

Given the fact that a card with two white sides cannot produce a black side facing up, a black and white card can produce a black side facing up 1 way, and a card with two black sides can produce a black side facing up two ways, the probability that the other side of a card with a black face up is $2/3$.

2M5

```
ww <- 0
bw <- 1
bb <- 2*2

(p <- bb/(bb + bw + ww))
```

```
## [1] 0.8
```

If there are two cards that have black on both sides, the probability that a card with a black side facing up also has black on the other side is now $4/5$.

2M6

```
ww <- 0*3
bw <- 1*2
bb <- 2

(p <- bb/(bb + bw + ww))
```

```
## [1] 0.5
```

If there are two ways to pull out a black and white card and 3 ways to pull out a white and white card for every way to pull out a black and black card, the probability that the other side is black is now $1/2$.

2M7

```
ww <- 0 * (1 + 0)
bb <- 2 * (2 + 1)
bw <- 1 * (0 + 2)

(p <- bb/(bb + bw + ww))
```

```
## [1] 0.75
```

If a second card is drawn with a white side face up, the probability that the first card with the black side facing up has black on the other side, is now 0.75.

Chapter 3 Homework

Easy Problems

```
p_grid <- seq(from = 0, to = 1, length.out = 1000)
prior <- rep(1, 1000)
lh <- dbinom(6, size = 9, prob = p_grid)
post <- lh * prior
post <- post/sum(post)
set.seed(100)
samples <- sample(p_grid, prob = post, size = 1e4, replace = T)
samples <- as_tibble(samples)
```

```
## Warning: Calling `as_tibble()` on a vector is discouraged, because the behavior is likely to change :
## This warning is displayed once per session.
```


3E1

```
less20 <- samples %>% filter(value < 0.2) %>%  
  summarise(sum = n()/1e4)
```

There is .0005 posterior probability that lies below $p = 0.2$.

3E2

```
more80 <- samples %>% filter(value > 0.8) %>%  
  summarise(sum = n()/1e4)
```

There is 0.1117 posterior probability that lies above $p = 0.8$.

3E3

```
bw2080 <- samples %>% filter(value > 0.2 & samples < 0.8) %>%  
  summarise(sum = n()/1e4)
```

There is 0.8878 posterior probability that lies between $p = 0.2$ and $p = 0.8$.

3E4

```
quantile(samples$value, p = 0.2)
```

```
##      20%  
## 0.5195195
```

20% of the posterior probability lies below $p = 0.52$.

3E5

```
quantile(samples$value, p = 0.8)
```

```
##      80%  
## 0.7567568
```

20% of the posterior probability lies above $p = 0.757$.

3E6

```
HPDI(samples$value, p = 0.66)
```

```
## |0.66      0.66|  
## 0.5205205 0.7847848
```

The values of p which contain the narrowest interval equal to 66% of the posterior probability are 0.521 and 0.785.

3E7

```
quantile(samples$value, p = 0.83)
```

```
##      83%  
## 0.7687688
```

```
quantile(samples$value, p = 0.17)
```

```
##          17%  
## 0.5005005
```

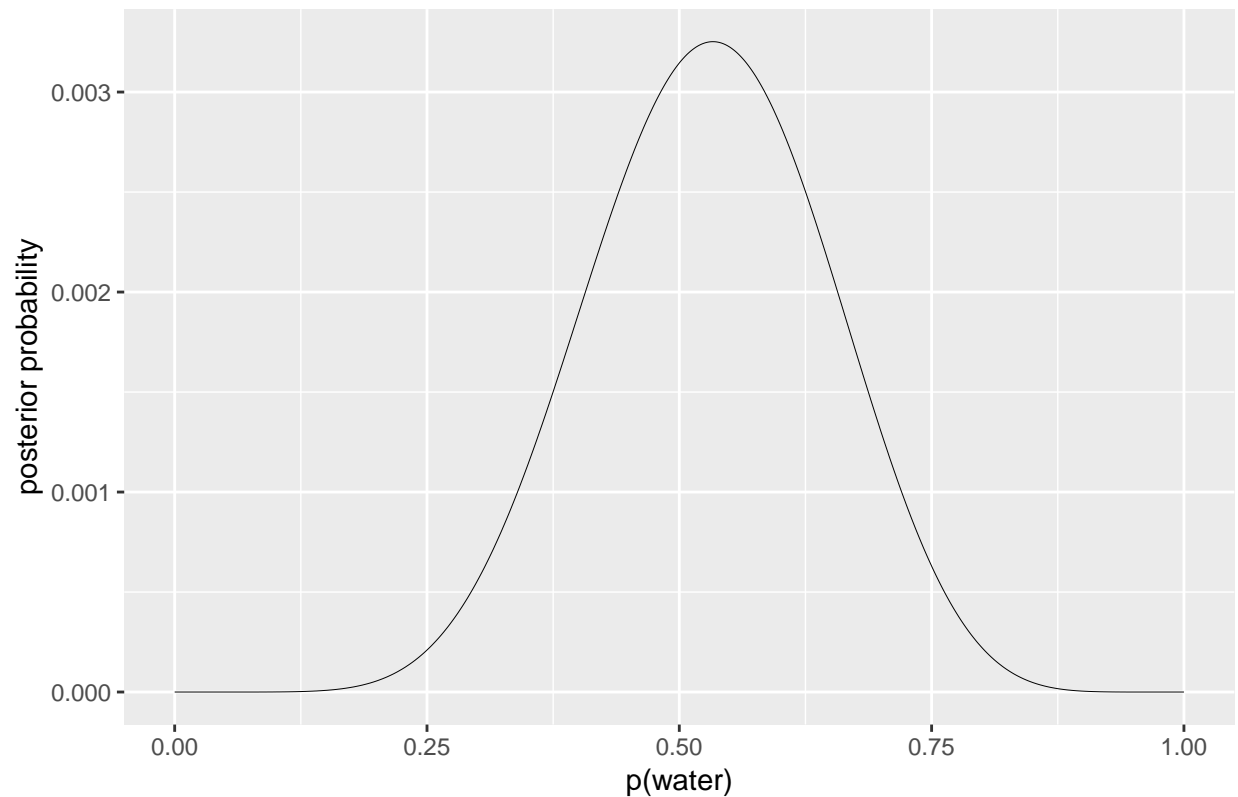
The values of p which contain 66% of the posterior probability, assuming equal posterior probability both below and above the interval are 0.500 and 0.769.

Medium Problems

3M1

```
n <- 1000  
n_success <- 8  
n_trials <- 15  
  
d <- tibble(p_grid = seq(from = 0, to = 1, length.out = n),  
            prior = 1) %>%  
  mutate(lh = dbinom(n_success, size = n_trials, prob = p_grid),  
         post = lh * prior,  
         post = post/sum(post))  
  
d %>%  
  ggplot(aes(x = p_grid, y = post)) +  
  geom_line(size = 1/10) +  
  labs(x = "p(water)",  
       y = "posterior probability") + ggtitle("8 Water in 15 Tosses")
```

8 Water in 15 Tosses



3M2

```
#drawing samples
samples <- tibble(samples = sample(d$p_grid, prob = d$post, size = 10000, replace = T)) %>%
  mutate(sample_n = 1:n())
head(samples)
```

```
## # A tibble: 6 x 2
##   samples sample_n
##   <dbl>     <int>
## 1  0.666         1
## 2  0.703         2
## 3  0.376         3
## 4  0.381         4
## 5  0.456         5
## 6  0.521         6
```

```
#90% HPDI
HPDI(samples$samples, p = 0.9)
```

```
##      |0.9      0.9|
## 0.3383383 0.7317317
```

The 90% HPDI for p is 0.338-0.731.

3M3

```
ppc <- tibble(sample = rbinom(1e4, size = 15, prob = samples$samples))
p8 <- ppc %>% filter(sample == 8) %>%
  summarise(sum = n()/1e4)
```

There is a p8 probability of getting 8 tosses out of 15.

3M4

```
newsim <- tibble(sample = rbinom(1e4, size = 9, prob = samples$samples))
p69 <- newsim %>% filter(sample == 6) %>%
  summarise(sum = n()/1e4)
```

There is a p69 probability of getting 6 tosses out of 9.

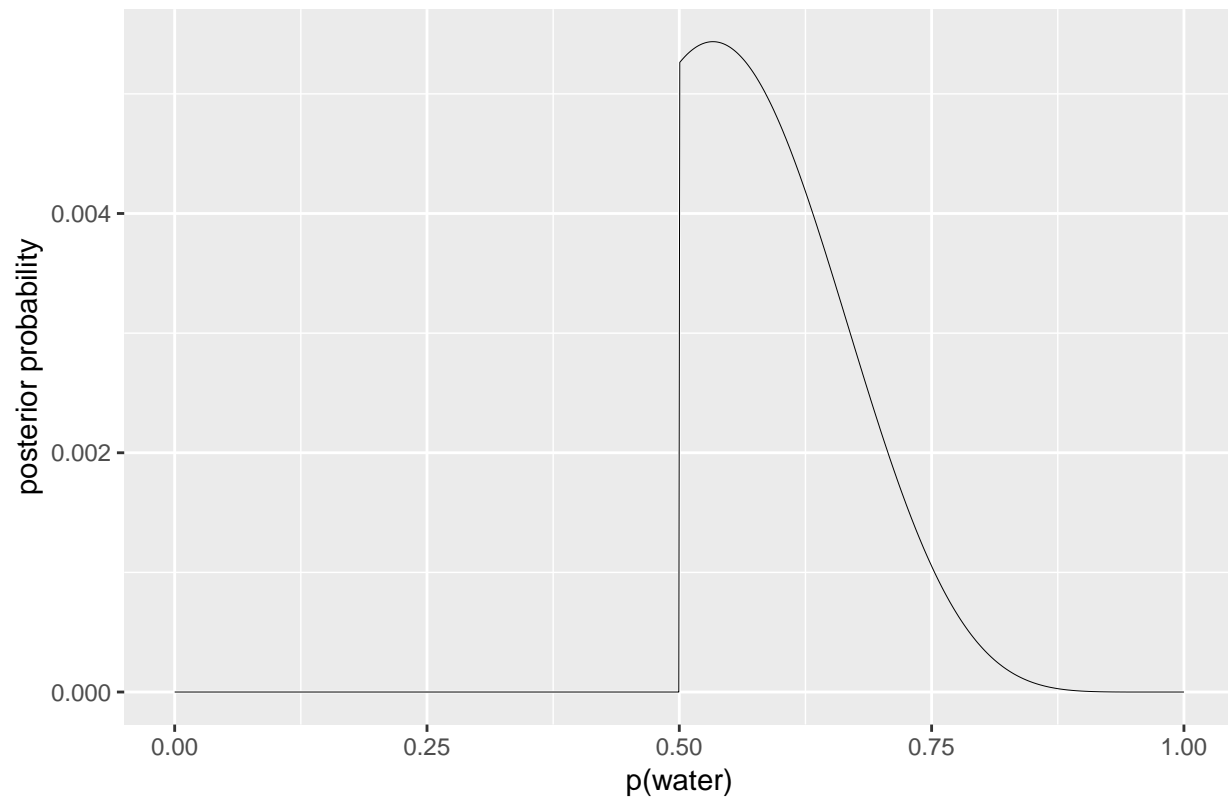
3M5

```
n <- 1000
n_success <- 8
n_trials <- 15

d <- tibble(p_grid = seq(from = 0, to = 1, length.out = n),
  prior = ifelse(p_grid < 0.5, 0, 2)) %>%
  mutate(lh = dbinom(n_success, size = n_trials, prob = p_grid),
    post = lh * prior,
    post = post/sum(post))

d %>%
  ggplot(aes(x = p_grid, y = post)) +
  geom_line(size = 1/10) +
  labs(x = "p(water)",
    y = "posterior probability") + ggtitle("8 Water in 15 Tosses, new Prior")
```

8 Water in 15 Tosses, new Prior



```
#drawing samples
samples <- tibble(samples = sample(d$p_grid, prob = d$post, size = 10000, replace = T)) %>%
  mutate(sample_n = 1:n())
```

```
#90% HPDI
HPDI(samples$samples, p = 0.9)
```

```
##      |0.9      0.9|
## 0.5005005 0.7097097
```

```
#posterior predictive check
ppc <- tibble(sample = rbinom(1e4, size = 15, prob = samples$samples))
p8 <- ppc %>% filter(sample == 8) %>%
  summarise(sum = n()/1e4)
print(p8)
```

```
## # A tibble: 1 x 1
##       sum
##   <dbl>
## 1 0.159
```

```
#probability of observing 6 water in 9 tosses
newsim <- tibble(sample = rbinom(1e4, size = 9, prob = samples$samples))
p69 <- newsim %>% filter(sample == 6) %>%
  summarise(sum = n()/1e4)
print(p69)
```

```
## # A tibble: 1 x 1
```

```
##      sum
##    <dbl>
## 1 0.236
```