Chapter-03-assignment.R

Ruijuan

February 23, 2016

install packages

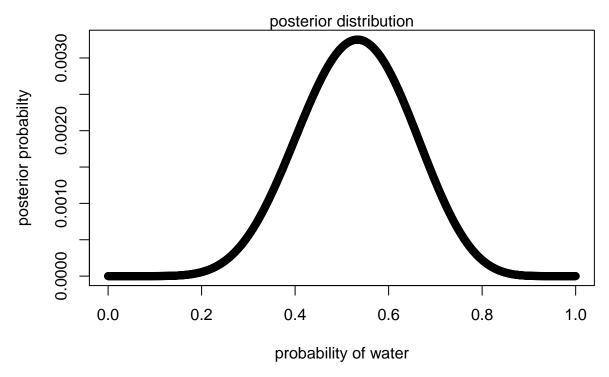
```
library(rmarkdown)
library(rstan)
## Loading required package: ggplot2
## rstan (Version 2.9.0-3, packaged: 2016-02-11 15:54:41 UTC, GitRev: 05c3d0058b6a)
## For execution on a local, multicore CPU with excess RAM we recommend calling
## rstan_options(auto_write = TRUE)
## options(mc.cores = parallel::detectCores())
library(rethinking)
## Loading required package: parallel
## rethinking (Version 1.58)
R code 3.27
p_grid <- seq(0, 1, length.out = 1000)</pre>
prior <- rep(1, 1000)</pre>
likelihood <- dbinom(6, size = 9, prob = p_grid)</pre>
posterior <- likelihood * prior</pre>
posterior <- posterior/sum(posterior)</pre>
set.seed(100)
samples <- sample(p_grid, prob = posterior, size = 1e4, replace = TRUE)</pre>
use the values in samples to answer the questions
3E1-3E3 the intervals of defined boundary
3E1. How much posterior probability lies below p=0.2
sum(posterior[p_grid < 0.2])</pre>
## [1] 0.0008560951
sum(samples<0.2)/1e4</pre>
## [1] 5e-04
3E2. How much posterior proabbility lies above p=0.8
```

```
sum(posterior[p_grid > 0.8])
## [1] 0.1203449
sum(samples>0.8)/1e4
## [1] 0.1117
3E3. How much posterior probability lies between p=0.2 and p=0.8
sum(posterior[p_grid > 0.2 & p_grid<0.8])</pre>
## [1] 0.878799
sum(samples>0.2 & samples<0.8)/1e4</pre>
## [1] 0.8878
The intervals of defined mass
3E4. 20% of the posterior probability lies below which value of p?
quantile(samples, 0.2)
##
          20%
## 0.5195195
3E5. 20% of the posterior probability lies above which value of p?
quantile(samples, 0.8)
          80%
## 0.7567568
3E6. Which values of p contain the narrowest interval equal to 66% of the posterior probability?
HPDI(samples, prob = 0.66)
        10.66
##
                   0.661
## 0.5205205 0.7847848
3E7. Which values of p contain 66% of the posterior probability, assuming equal posterior
proability both below and above the interval?
PI(samples, prob = 0.66)
          17%
                     83%
## 0.5005005 0.7687688
```

3M1. Suppose the globe tossing data had turned out to be 8 water in 15 tosses.

Construct the posterior distribution, using grid approximation. Use the same flat prior as before

```
p_grid <- seq(0, 1, length.out = 1000)
prior <- rep(1, 1000)
likelihood <- dbinom(8, size = 15, prob = p_grid)
posterior <- likelihood * prior
posterior <- posterior/sum(posterior)</pre>
```



3M2. Draw 10,000 samples from the grid approximation from above.

Then use the samples to calculate the 90% HPDI() for p.

```
set.seed(8808)
samples <- sample(p_grid, prob = posterior, size = 1e4, replace = TRUE)
# the 90% highest posterior density (narrowest) interval
HPDI(samples, prob = 0.9)</pre>
```

```
## |0.9 0.9|
## 0.3413413 0.7317317
```

STOP AFTER 3M2 FOR 02/25 ASSIGNMENT

3M3

3M4

3M5

3H1

3H2

3H3

3H4

3H5