

UCBadmit

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UCBadmit

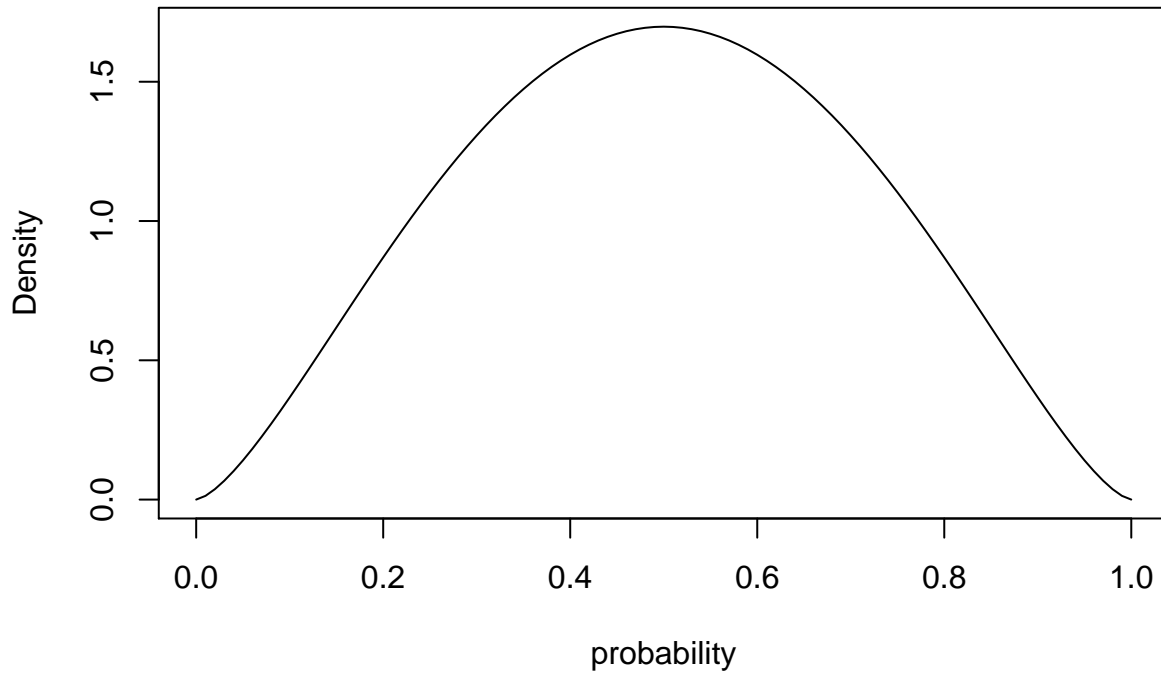
reference: McElreath, Statistical Rethinking, chap 11, p.347

```
library(rethinking)
library(ggplot2)
```

admissions

beta distribution

```
## R code 11.25
pbar <- 0.5
theta <- 5
curve( dbeta2(x,pbar,theta) , from=0 , to=1 ,
       xlab="probability" , ylab="Density" )
```



data

```
## R code 11.26
data(UCBadmit)
d <- UCBadmit
str(d)

## 'data.frame':  12 obs. of  5 variables:
## $ dept          : Factor w/ 6 levels "A","B","C","D",...: 1 1 2 2 3 3 4 4 5 5 ...
## $ applicant.gender: Factor w/ 2 levels "female","male": 2 1 2 1 2 1 2 1 2 1 ...
## $ admit          : int  512 89 353 17 120 202 138 131 53 94 ...
## $ reject         : int  313 19 207 8 205 391 279 244 138 299 ...
## $ applications   : int  825 108 560 25 325 593 417 375 191 393 ...

d <- d[ , c("admit", "applications")]
```

model

$$A_i \sim \text{BetaBinomial}(n_i, \bar{p}_i, \theta)$$

$$\text{logit}(\bar{p}_i) = \alpha$$

$$\alpha \sim \text{Normal}(0, 10)$$

$$\theta \sim \text{HalfCauchy}(0, 1)$$

stan

```

m11.5 <- map2stan(
  alist(
    admit ~ dbetabinom(applications,pbar,theta),
    logit(pbar) <- a,
    a ~ dnorm(0,2),
    theta ~ dexp(1)
  ),
  data=d,
  constraints=list(theta="lower=0"),
  start=list(theta=3),
  iter=4000 , warmup=1000 , chains=2 , cores=2 )

```

```

##
## SAMPLING FOR MODEL 'admit ~ dbetabinom(applications, pbar, theta)' NOW (CHAIN 1).
## WARNING: No variance estimation is
##           performed for num_warmup < 20
##
##
## Chain 1, Iteration: 1 / 1 [100%] (Sampling)
## Elapsed Time: 6e-06 seconds (Warm-up)
##               7e-05 seconds (Sampling)
##               7.6e-05 seconds (Total)
## Computing WAIC
## Constructing posterior predictions
## [ 600 / 6000 ]
## [ 1200 / 6000 ]
## [ 1800 / 6000 ]
## [ 2400 / 6000 ]
## [ 3000 / 6000 ]
## [ 3600 / 6000 ]
## [ 4200 / 6000 ]
## [ 4800 / 6000 ]
## [ 5400 / 6000 ]
## [ 6000 / 6000 ]

```

```

## R code 11.27
precis(m11.5)

```

```

##           Mean StdDev lower 0.89 upper 0.89 n_eff Rhat
## theta  2.76  0.99      1.20      4.19 4610  1
## a      -0.37  0.31     -0.89      0.10 4437  1

```

analysis

```

## R code 11.28
post <- extract.samples(m11.5)
quantile( logistic(post$a) , c(0.025,0.5,0.975) )

```

```

##           2.5%           50%           97.5%
## 0.2731602 0.4079590 0.5583337

```

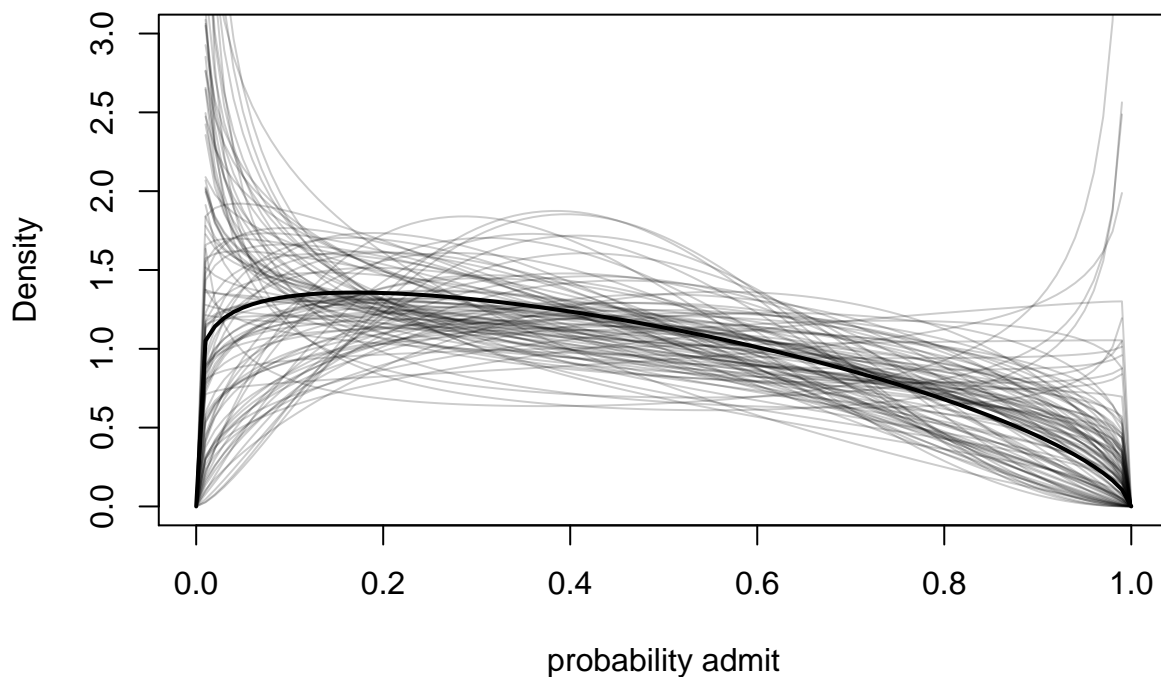
```

## R code 11.29
post <- extract.samples(m11.5)

# draw posterior mean beta distribution
curve( dbeta2(x,mean(logistic(post$a)),mean(post$theta)) , from=0 , to=1 ,
       ylab="Density" , xlab="probability admit", ylim=c(0,3) , lwd=2 )

# draw 100 beta distributions sampled from posterior
for ( i in 1:100 ) {
  p <- logistic( post$a[i] )
  theta <- post$theta[i]
  curve( dbeta2(x,p,theta) , add=TRUE , col=col.alpha("black",0.2) )
}

```



postcheck

```

## R code 11.30
postcheck(m11.5)

## [ 100 / 1000 ]
[ 200 / 1000 ]
[ 300 / 1000 ]
[ 400 / 1000 ]
[ 500 / 1000 ]
[ 600 / 1000 ]
[ 700 / 1000 ]
[ 800 / 1000 ]
[ 900 / 1000 ]
[ 1000 / 1000 ]
## [ 100 / 1000 ]
[ 200 / 1000 ]

```

[300 / 1000]
[400 / 1000]
[500 / 1000]
[600 / 1000]
[700 / 1000]
[800 / 1000]
[900 / 1000]
[1000 / 1000]

