

Bayesian Models 1

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Outline

This is a first short writing about Bayesian statistics. I am going to work through an example of performing posterior inference for data that have a binomial distribution.

1. Load in the data;
2. Perform some exploratory analysis;
3. Perform posterior inference using a discrete prior;
4. Perform posterior inference using a beta prior.

Suppose we want to estimate θ , the proportion of animals that can fly, from a random sample of animals. Before we start looking at the data, let's think about a prior for θ .

- (1) What values can it take on?
- (2) What's a reasonable prior mean?
- (3) How uncertain are we about our prior mean?

Suppose we want to estimate θ , the proportion of animals that can fly, from a random sample of animals.

We'll use the animals data set, which measures characteristics of 20 species.

It is available in the **cluster** package, one of the basic packages that comes with R.

```
library(cluster)
```

```
## Warning: package 'cluster' was built under R version 3.3.3
```

```
data(animals)
```

Description of the Data

The animals data includes indicators:

- (1) war, warm-blooded;
- (2) fly, able to fly;
- (3) ver, vertebrate;
- (4) end, endangered;
- (5) gro, live in groups;
- (6) hai, have hair.

```
help(animals)
```

```
## starting httpd help server ... done
```

Preparing the Data

The indicators are coded to take on: 2, if the animal has the characteristic; 1, otherwise.

We want to replace the 2's and 1's with 1's and 0's.

```
animals <- apply(animals, c(1, 2),
                function(x) {x - 1})
```

Describing the Data

Since we're interested in θ , the proportion of animals that can fly, we should look at the number of animals in the sample that can fly.

```
sum(animals[, "fly"])
```

```
## [1] 4
```

So we have $y=4$ from a $\text{Binomial}(n=20, \theta)$ distribution.

Inference Under a Discrete Prior

Suppose we believed:

- (1) $\theta \in \{0.01, \dots, 0.99\}$;
- (2) $p(\theta) = 1/99$ for each θ . This prior encodes the belief that it is impossible for: (a) All animals to fly, i.e. $\theta=1$; (b) No animals to fly, i.e. $\theta=0$. This is a uniform prior on θ . We can compute the posterior distribution using Bayes' rule.

```
y <- sum(animals[, "fly"])
n <- nrow(animals)

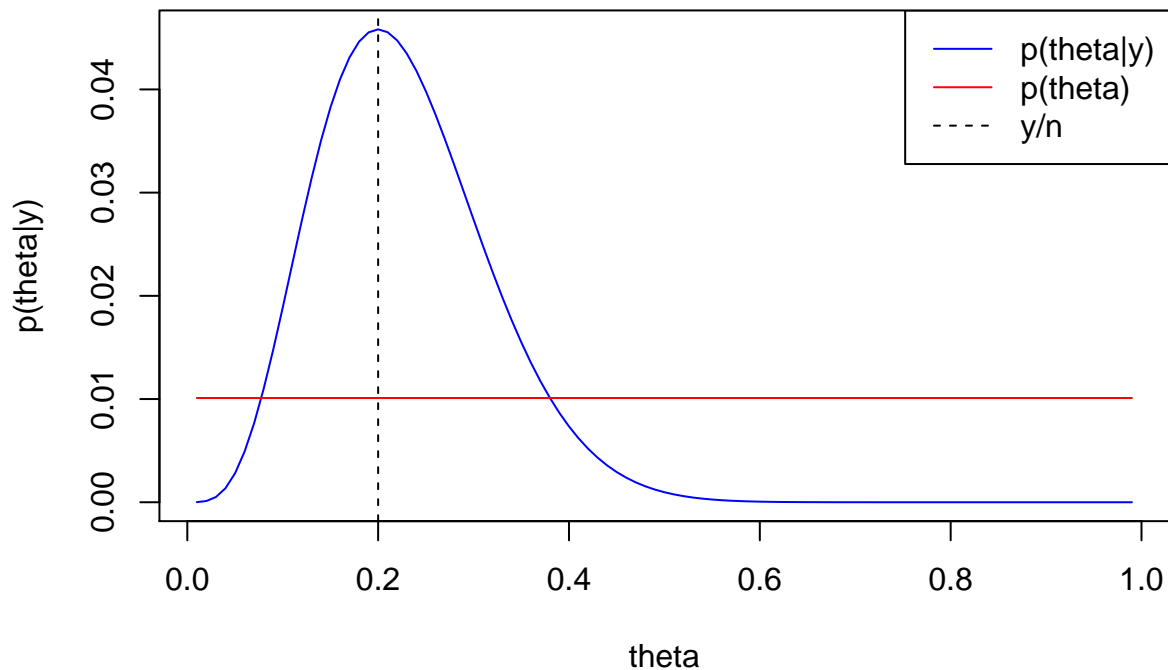
Theta.1 <- seq(0.01, 0.99, by = 0.01)
py.theta.1 <- dbinom(y, n, Theta.1)
ptheta.1 <- rep(1/length(Theta.1), length(Theta.1))
pytheta.1 <- py.theta.1*ptheta.1

ptheta.y.1 <- pytheta.1/(sum(pytheta.1))
plot(Theta.1, ptheta.y.1, type = "l",
     col = "blue",
     xlab = "theta",
     ylab = "p(theta|y)",
     cex.lab = 1, cex.axis = 1)

lines(Theta.1, ptheta.1, col = "red", lty = 1)

abline(v = mean(animals[, "fly"]), lty = 2)

legend("topright", lty = c(1, 1, 2),
      col = c("blue", "red", "black"),
      legend = c("p(theta|y)", "p(theta)",
                 "y/n"))
```



We can compute the posterior mean easily.

```
etheta.y.1 <- sum(ptheta.y.1*Theta.1)
etheta.y.1
```

```
## [1] 0.2272727
```

We can also compute the posterior variance easily.

```
vtheta.y.1 <- sum(ptheta.y.1*(Theta.1 - etheta.y.1)^2)
vtheta.y.1
```

```
## [1] 0.007635645
```

Inference Under a Continuous Prior

Suppose we believed: (1) $\theta \in (0,1)$; (2) $p(\theta)=1$ for each $\theta \in (0,1)$. This is also uniform prior.

This prior corresponds to the $\text{Beta}(1, 1)$ distribution.

As you'll learn later, the beta distribution is conjugate for the binomial sampling model.

This means that we know that the posterior distribution of θ will be a beta distribution, with parameters we have closed form solutions for!

$p(\theta|y) \propto B(1+y, 1+n-y)$

```
atheta.y.2 <- 1 + y
btheta.y.2 <- 1 + n - y
Theta.2 <- seq(0.01, 0.99, by = 0.01)
```

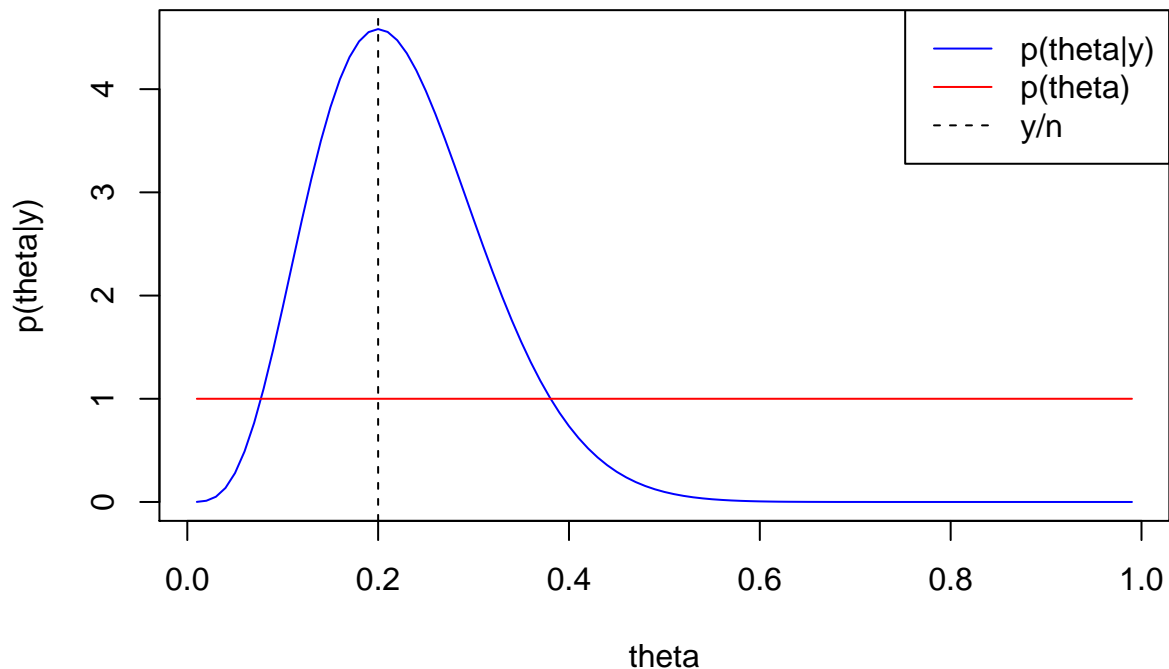
```

ptheta.2 <- dbeta(Theta.2, 1, 1)
ptheta.y.2 <- dbeta(Theta.2, atheta.y.2,
                    btheta.y.2)

plot(Theta.2, ptheta.y.2, type = "l",
     col = "blue",
     xlab = "theta",
     ylab = "p(theta|y)",
     cex.lab = 1, cex.axis = 1)

lines(Theta.2, ptheta.2, col = "red", lty = 1)
abline(v = mean(animals[, "fly"]), lty = 2)
legend("topright", lty = c(1, 1, 2),
     col = c("blue", "red", "black"),
     legend = c("p(theta|y)", "p(theta)",
                "y/n"))

```



We can directly compute the posterior mean from what we know about the beta distribution.

```

etheta.y.2 <- atheta.y.2/(atheta.y.2 + btheta.y.2)
etheta.y.2

```

```
## [1] 0.2272727
```

We can also compute the posterior variance.

```

vtheta.y.2 <- atheta.y.2*btheta.y.2/((atheta.y.2 + btheta.y.2)^2*(atheta.y.2 + btheta.y.2 + 1))
vtheta.y.2

```

```
## [1] 0.007635645
```

Note that we can compute the mode and any other feature of the Beta distribution that can be written as a function of its parameters.

Inference Under Another Continuous Prior

Suppose we believed:

- (1) $\theta \sim (0,1)$;
- (2) $E[\theta] = 0.43$, i.e. about 2/5 of animals fly;
- (3) $V[\theta] = 0.03$, i.e. we're pretty certain θ 's around 0.43.

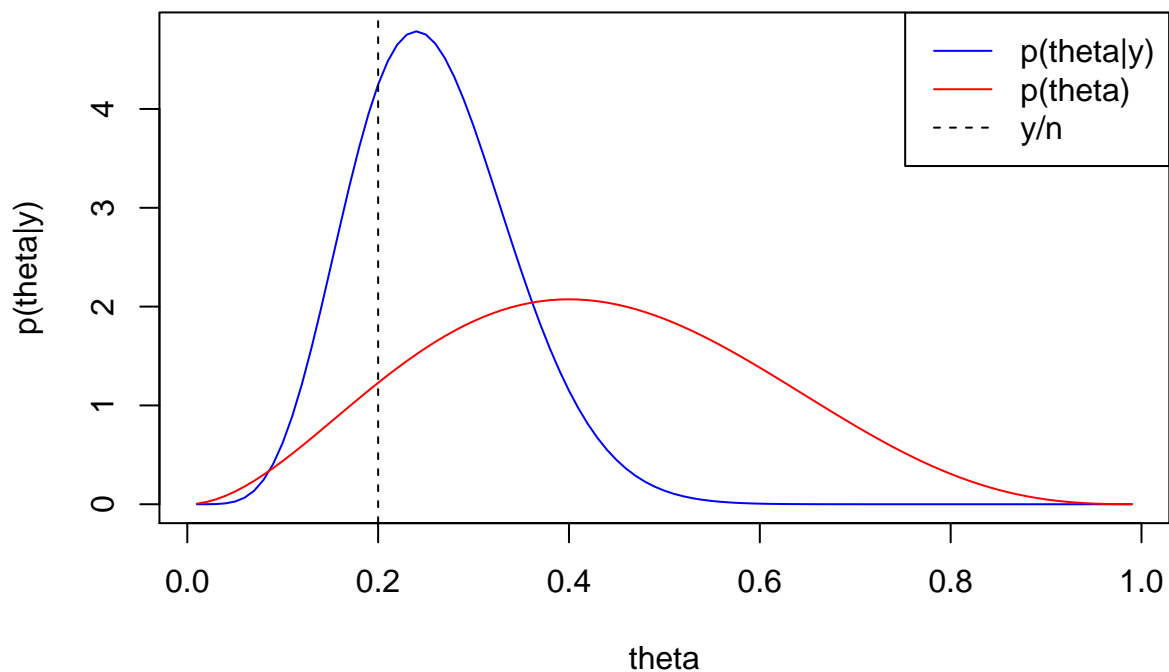
This corresponds to a beta distribution with $a=3$ and $b=4$.

Again, we can use the fact that the beta distribution is conjugate for the binomial sampling model to compute the parameters of the posterior distribution.

```
atheta.y.3 <- 3 + y
btheta.y.3 <- 4 + n - y
Theta.3 <- seq(0.01, 0.99, by = 0.01)
ptheta.3 <- dbeta(Theta.3, 3, 4)
ptheta.y.3 <- dbeta(Theta.3, atheta.y.3,
                   btheta.y.3)

plot(Theta.3, ptheta.y.3, type = "l",
     col = "blue",
     xlab = "theta",
     ylab = "p(theta|y)",
     cex.lab = 1, cex.axis = 1)

lines(Theta.3, ptheta.3, col = "red", lty = 1)
abline(v = mean(animals[, "fly"]), lty = 2)
legend("topright", lty = c(1, 1, 2),
     col = c("blue", "red", "black"),
     legend = c("p(theta|y)", "p(theta)",
                "y/n"))
```



Again, we can directly compute the posterior mean from what we know about the beta distribution.

```
etheta.y.3 <- atheta.y.3/(atheta.y.3 + btheta.y.3)
etheta.y.3
```

```
## [1] 0.2592593
```

We can also compute the posterior variance.

```
vtheta.y.3 <- atheta.y.3*btheta.y.3/((atheta.y.3 + btheta.y.3)^2*(atheta.y.3 + btheta.y.3 + 1))
vtheta.y.3
```

```
## [1] 0.006858711
```

Again, we can compute the mode and any other feature of the Beta distribution that can be written as a function of its parameters.