

# Lab 1

2024-03-29

```
library(ggplot2)
library(patchwork)
```

## 1 - Daniel Bernoulli

Let  $y_1, \dots, y_n | \theta \sim \text{Bern}(\theta)$ , and assume that you have obtained a sample with  $s = 22$  successes in  $n = 70$  trials. Assume a  $\text{Beta}(\alpha_0, \beta_0)$  prior for  $\theta$  and let  $\alpha_0 = \beta_0 = 8$ .

a)

Draw 10000 random values ( $n_{\text{Draws}} = 10000$ ) from the posterior  $\theta | y \sim \text{Beta}(\alpha_0 + s, \beta_0 + f)$ , where  $y = (y_1, \dots, y_n)$ , and verify graphically that the posterior mean  $E[\theta | y]$  and standard deviation  $\text{SD}[\theta | y]$  converges to the true values as the number of random draws grows large. [Hint: use `rbeta()` to draw random values and make graphs of the sample means and standard deviations of  $\theta$  as a function of the accumulating number of drawn values].

```
n <- 70      # Sample trials
s <- 22      # Sample successes
f <- n - s   # Sample failures

a0 <- b0 <- 8    # Prior parameters
nDraws <- 10000  # Number of random draws from posterior

an = a0 + s
bn = b0 + f
rdraw <- rbeta(nDraws, an, bn)

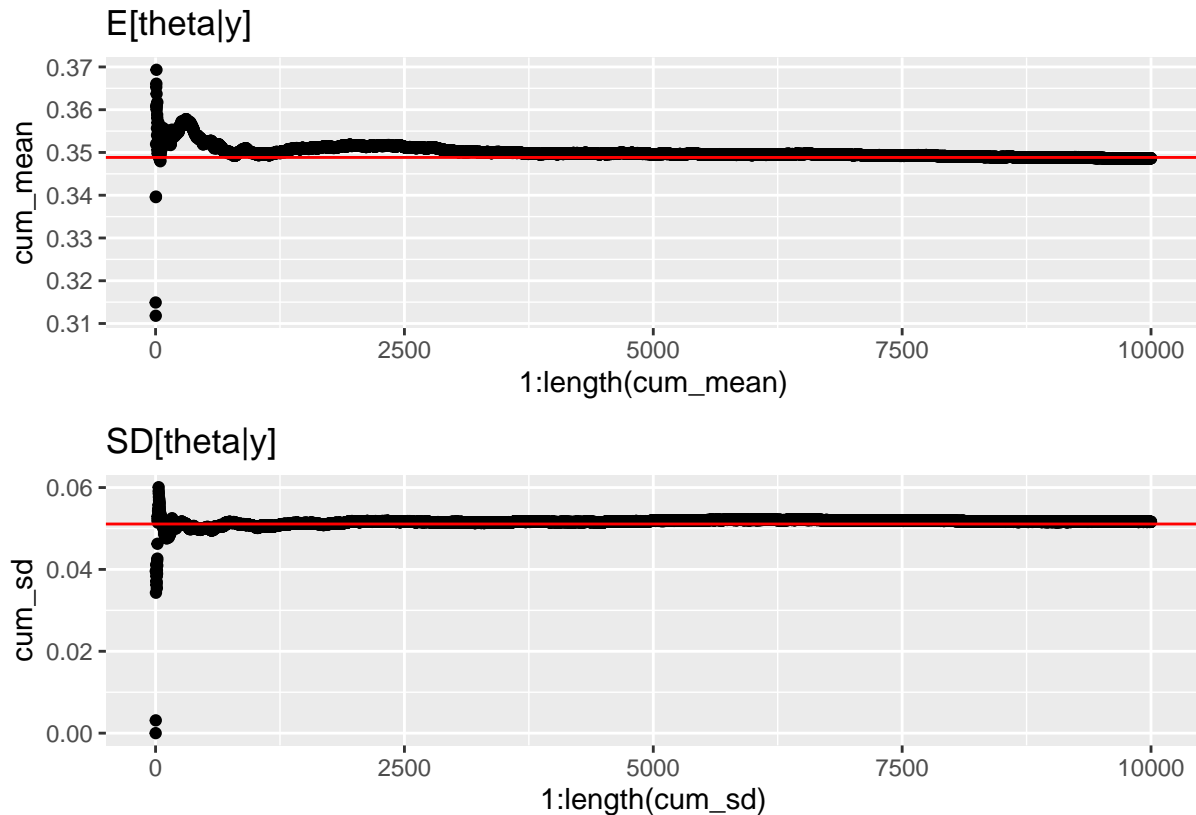
cum_n <- 1:nDraws
cum_mean <- cumsum(rdraw) / cum_n
cum_mean2 <- cumsum(rdraw ** 2) / cum_n
cum_var <- (cum_mean2 - cum_mean ** 2) * (n / (n - 1))
cum_sd <- sqrt(cum_var)

anlyt_mean <- an / (an + bn)
anlyt_sd <- sqrt(an * bn / ((an + bn) ** 2 * (an + bn + 1)))

p1 <- ggplot() +
  geom_point(aes(x=1:length(cum_mean), y=cum_mean)) +
  geom_hline(yintercept=anlyt_mean, color="red") + #or abline()
  ggtitle("E[ $\theta | y$ "])
```

```
p2 <- ggplot() +
  geom_point(aes(x=1:length(cum_sd), y=cum_sd)) +
  geom_hline(yintercept=anlyt_sd, color="red") +
  ggtitle("SD[theta|y]")
```

p1 / p2



b)

Draw 10000 random values from the posterior to compute the posterior probability  $\Pr(\theta > 0.3|y)$  and compare with the exact value from the Beta posterior. [Hint: use `pbeta()`].

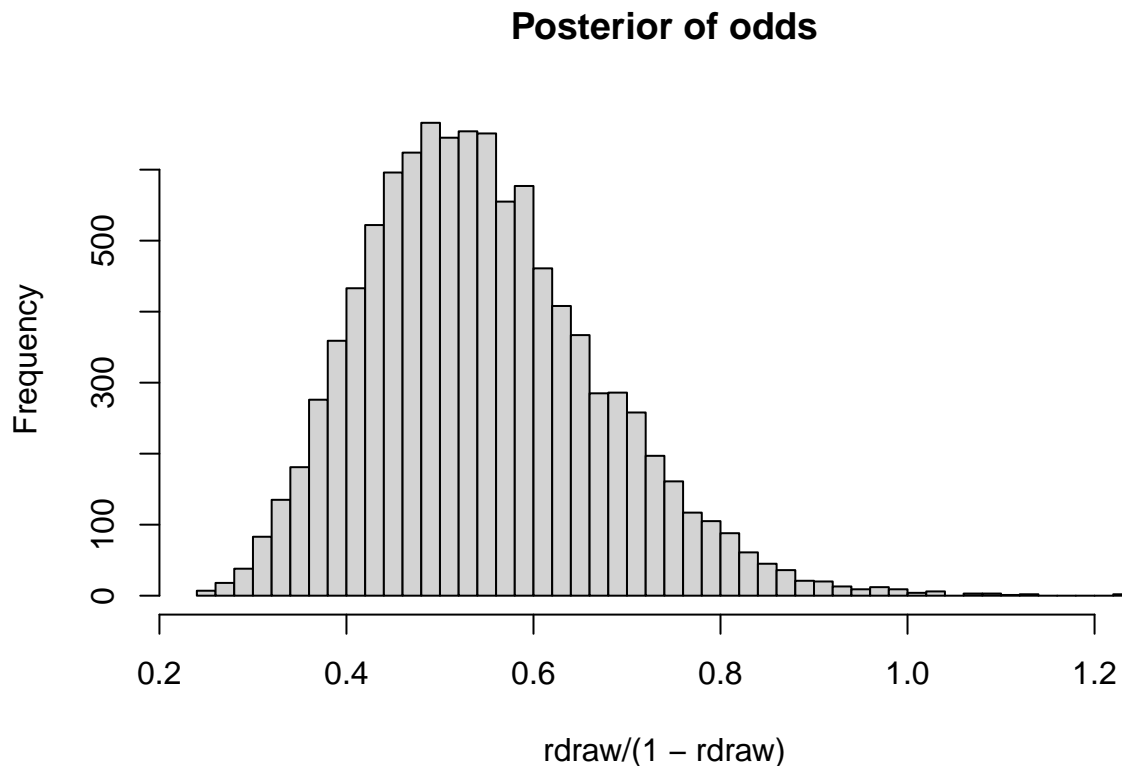
```
p_draw = sum(rdraw > 0.3) / length(rdraw)
p_anlyt = pbeta(0.3, an, bn, lower.tail=FALSE)
```

```
## Random draw from posterior:
##
##   Pr(theta > 0.3 | y) =  0.8242
##
## Exact value from Beta distribution:
##
##   Pr(theta > 0.3 | y) =  0.8285936
```

c)

Draw 10000 random values from the posterior of the odds  $\phi = \theta/(1-\theta)$  by using the previous random draws from the Beta posterior for  $\theta$  and plot the posterior distribution of  $\phi$ . [Hint: `hist()` and `density()` can be utilized].

```
hist(rdraw / (1 - rdraw), breaks=50, main="Posterior of odds")
```



## 2 - Log-normal distribution and the Gini coefficient

A common model for non-negative continuous variables is the log-normal distribution.

$$y \sim \text{logNormal}(\mu, \sigma^2) \Rightarrow \log(y) \sim \text{Normal}(\mu, \sigma^2)$$

Let  $y_1, \dots, y_n$ ,  $\log N(\mu, \sigma^2)$ , where  $\mu = 3.6$  is assumed to be known but  $\sigma$  is unknown with non-informative prior  $p(\sigma^2) \propto 1/\sigma^2$ . The posterior for  $\sigma^2$  is the  $\text{Inv-}\chi^2(n, \text{to}^2)$  distribution.

a)

Draw 10000 random values from the posterior of  $\sigma^2$  by assuming  $\mu = 3.6$  and plot the posterior distribution.

Normal model with unknown variance:

model:

$$y_1, \dots, y_n | \mu, \sigma \sim \text{logN}(\mu, \sigma^2)$$

Non-informative prior

$$p(\sigma^2) \propto 1/\sigma^2$$

Posterior:

mu is given here i.e 3.6 (Only in the below formula we take it as n no matter if it is “n” or “n-1”)

$$\mu | \sigma^2, x \sim N(x, \sigma^2/n),$$

(Note, in the slides it was given “n-1” , but here we taken “n” since that is what is given in the question)

$$\sigma^2 | x \sim \text{Inv-}\chi^2(n, \tau^2)$$

where (in the slides it was given “n-1” , but here we taken “n” since that is what is given in the question))

$$\tau^2 = (\sum (\log y_i - \mu)^2) / n$$

To simulate from the posterior:

(Note, in the slides it was given “n-1” , but here we taken “n” since that is what is given in the question)

$$X \sim \chi^2(n)$$

A draw from Inv- $\chi^2(n, \tau^2)$  distribution((Note, in the slides it was given “n-1” , but here we taken “n” since that is what is given in the question))

$$\sigma^2 = (n * \tau^2) / X$$

Then draw mu from the above formula but in this case we assume it to be 3.6

```
obs <- c(33, 24, 48, 32, 55, 74, 23, 17)
mu <- 3.6

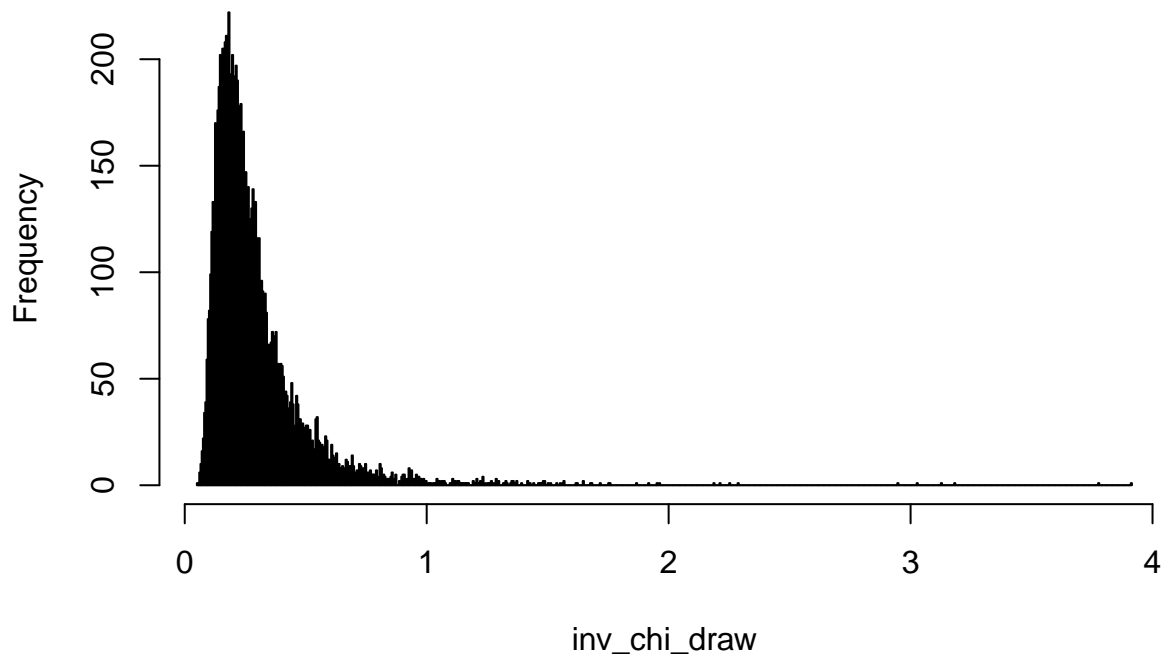
#Sample variance calculation
values <- c()
for (i in 1:length(obs)){
  values[i] <- (log(obs[i]) - mu)^2
}

sample_var <- sum(values)/8

# Simulation from posterior:
inv_chi_draw <- c()
post_draw <- function(n){
  for(i in 1:n){
    chi_draw <- rchisq(1,8)
    inv_chi_draw[i] <- ((8)*sample_var)/(chi_draw)
  }
  hist(inv_chi_draw, breaks=1000)
}

post_draw(10000)
```

## Histogram of inv\_chi\_draw



b)

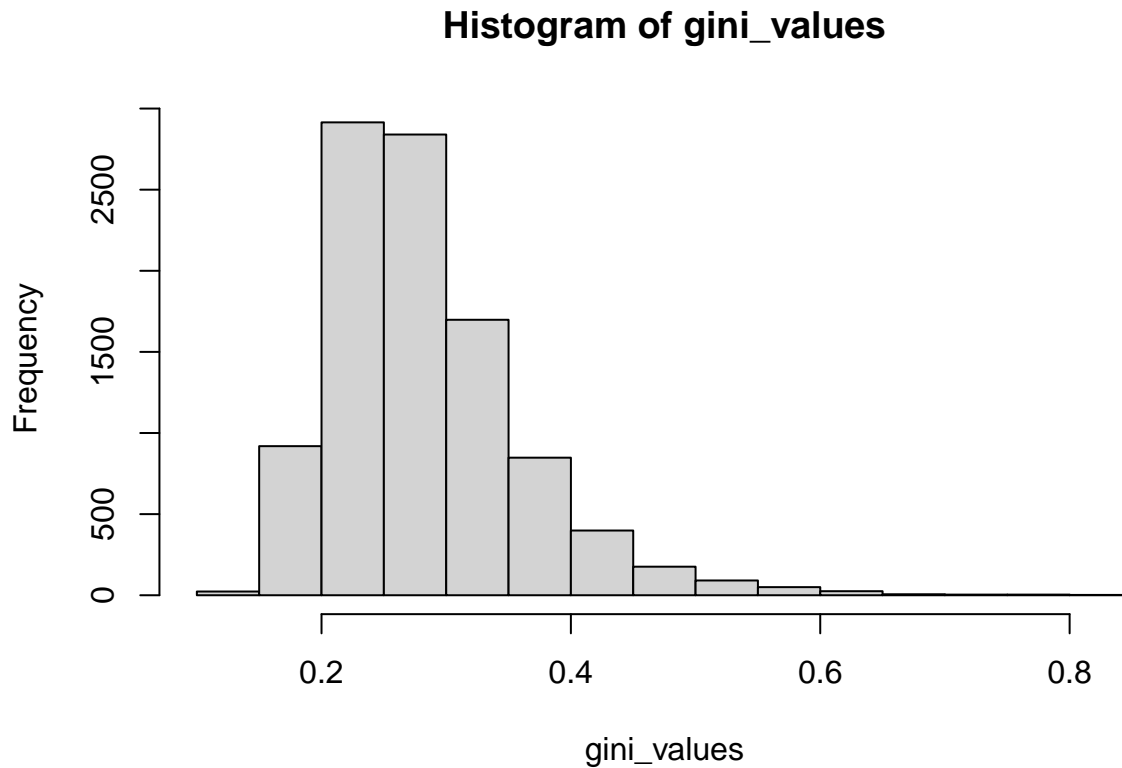
We need to plot the posterior distribution of the gini coefficient which follows a log N ( $\mu$ ,  $\sigma^2$ ) distribution. ( $G|y$ )

$$G = 2\Phi(\sigma/\sqrt{2}) - 1$$

and  $\Phi$  is a cumulative distribution function (CDF) for the standard normal distribution with mean zero and unit variance. So we use `pnorm()` to calculate this since `pnorm()` gives the distribution function while `dnorm()` gives the density function.

```
gini_values <- c()
gini <- function(n){
  for(i in 1:n){
    gini_values[i] <- 2*pnorm(sqrt(inv_chi_draw[i])/sqrt(2))-1
  }
  hist(gini_values)
}

gini(10000)
```



c)

Use the posterior draws from b) to compute a 95% equal tail credible interval for G. A 95% equal tail credible interval (a,b) cuts off 2.5% percent of the posterior probability mass to the left of a, and 2.5% to the right of b.

Approximate 95% credible interval for G

$$E(G|y) \pm 1.96 \cdot SD(G|y)$$

```
upper <- mean(gini_values) + 1.96*sd(gini_values)
lower <- mean(gini_values) - 1.96*sd(gini_values)
```

```
## [1] "Upper limit:"
```

```
## [1] 0.4340629
```

```
## [1] "lower limit:"
```

```
## [1] 0.1315154
```

d)

Use the posterior draws from b) to compute a 95% Highest Posterior Density Interval (HPDI) for G.

```
gini_den<-density(gini_values)
gini_den_df <- data.frame(x = gini_den$x, y = gini_den$y)

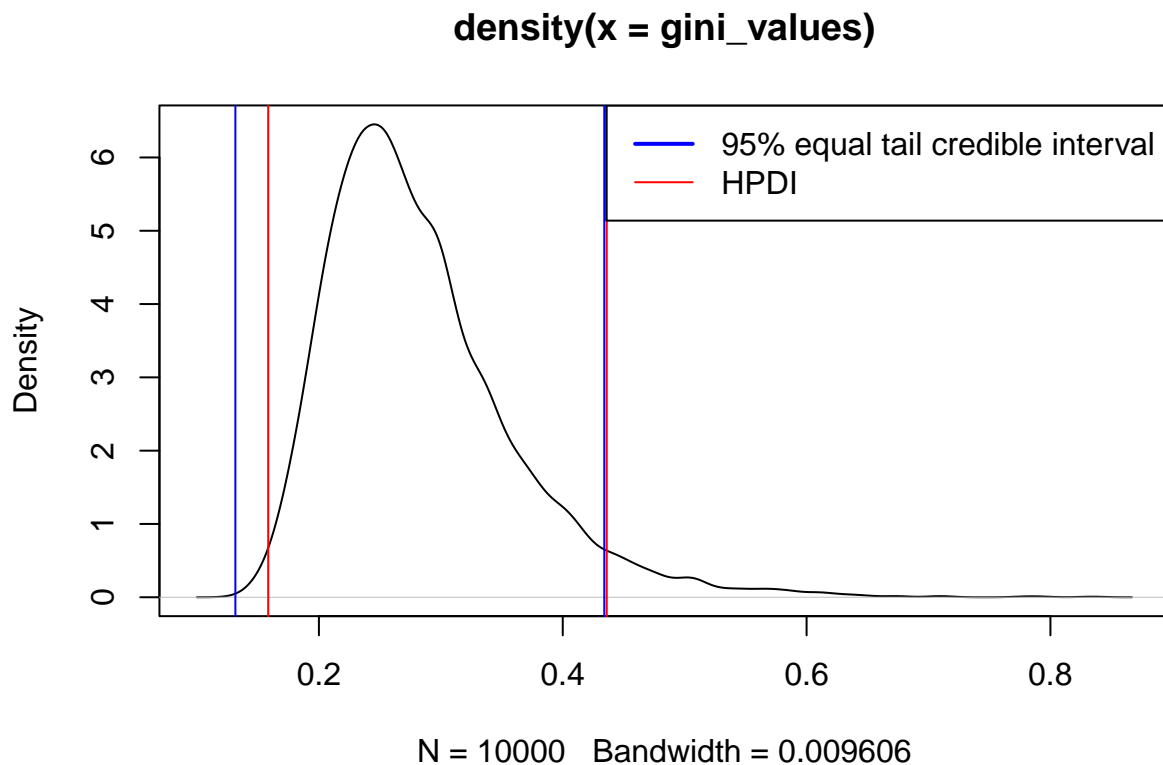
#x gives the corrdinates of the points where the density is estimated and y gives the estimated
#density values.

gini_den_df <- gini_den_df[order(gini_den_df$y),]
gini_den_df<- data.frame(x = gini_den_df$x, y = gini_den_df$y)
gini_den_cum = cumsum(gini_den_df$y)/sum(gini_den_df$y) # we divide so that it sums to 1.
value = which(gini_den_cum >= 0.05)[1]
gini_den_df = gini_den_df[(value+1):length(gini_den$y),]
gini_den_interval <- c(min(gini_den_df$x),max(gini_den_df$x))

plot(gini_den)
abline(v=gini_den_interval[1],col ="red")
abline(v=gini_den_interval[2], col = "red")

abline(v=upper,col ="blue")
abline(v=lower, col = "blue")

legend("topright", legend = c("95% equal tail credible interval", "HPDI"),
      col = c("blue", "red"), lty = 1, lwd = c(2, 1))
```



### 3 - Bayesian inference for the concentration parameter in the von Mises distribution

a)

Derive the expression for what the posterior  $p(x|y, \kappa)$  is proportional to. Hence, derive the function  $f(\kappa)$  such that  $p(x|y, \mu) = f(\kappa)$ . Then, plot the posterior distribution of  $\kappa$  for the wind direction data over a fine grid of  $\kappa$  values.

Model:

$$p(y_i|\mu, \kappa) = \frac{\exp(\kappa \cos(y_i - \mu))}{2\pi I_0(\kappa)}$$

Likelihood:

$$\begin{aligned} p(y|\mu, \kappa) &= \prod_{i=1}^{10} \frac{\exp(\kappa \cos(y_i - \mu))}{2\pi I_0(\kappa)} \\ &= \frac{\exp(\kappa \sum_{i=1}^{10} \cos(y_i - \mu))}{(2\pi I_0(\kappa))^{10}} \end{aligned}$$

Prior: (exponential distribution)

$$p(\kappa) = 0.5e^{-0.5\kappa}$$

Posterior:

$$\begin{aligned} p(\kappa|y, \mu) &\propto \frac{\exp(\kappa \sum_{i=1}^{10} \cos(y_i - \mu))}{2\pi I_0(\kappa)} * \exp(-0.5\kappa) \\ &\propto \frac{\exp[\kappa(-0.5 + \sum_{i=1}^{10} \cos(y_i - \mu))]}{I_0(\kappa)^{10}} \end{aligned}$$

```
y <- c(-2.79, 2.33, 1.83, -2.44, 2.23, 2.33, 2.07, 2.02, 2.14, 2.54) # Data points
mu <- 2.4 # Given parameter mu

post <- function(k, y, mu) {
  num <- exp(k * (-0.5 + sum(cos(y - mu))))
  denom <- besseli(k, 0) ** 10

  res = num / denom
  res[is.na(res)] <- 0

  return (res)
}

post_norm <- function(k, y, mu, c_norm) {
  res = post(k, y, mu) / c_norm

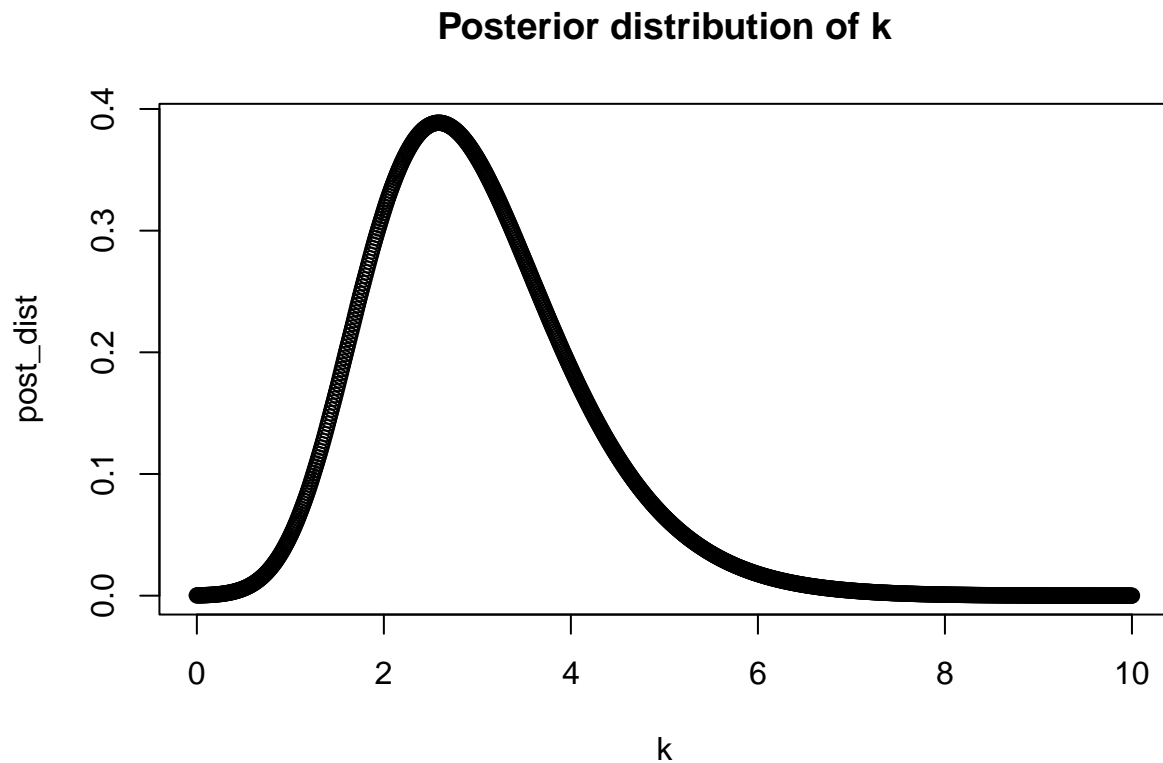
  return (res)
}

#[Hint: you need to normalize the posterior distribution of  so that it integrates to one.]
c_norm <- integrate(post, 0, Inf, y=y, mu=mu)$value
```



```
k <- seq(from=0, to=10, length.out=1000)
post_dist <- post_norm(k, y, mu, c_norm)

plot(k, post_dist, main="Posterior distribution of k")
```



Find the (approximate) posterior mode of  $k$  from the information in a). ## b)

```
paste0("Approx. mode of distribution of k: ", round(k[which.max(post_dist)], 3))
```

```
## [1] "Approx. mode of distribution of k: 2.583"
```