

3.3

a)

Tumor counts: A cancer laboratory is estimating the rate of tumorigenesis in two strains of mice, A and B. They have tumor count data for 10 mice in strain A and 13 mice in strain B. Type A mice have been well studied, and information from other laboratories suggests that type A mice have tumor counts that are approximately Poisson-distributed with a mean of 12. Tumor count rates for type B mice are unknown, but type B mice are related to type A mice. The observed tumor counts for the two populations are

$$y_A = (12, 9, 12, 14, 13, 13, 15, 8, 15, 6);$$

$$y_B = (11, 11, 10, 9, 9, 8, 7, 10, 6, 8, 8, 9, 7).$$

a)

Find the posterior distributions, means, variances and 95% quantilebased confidence intervals for θ_A and θ_B , assuming a Poisson sampling distribution for each group and the following prior distribution:

$$\theta_A \sim \text{gamma}(120, 10),$$

$$\theta_B \sim \text{gamma}(12, 1),$$

$$p(\theta_A, \theta_B) = p(\theta_A) \times p(\theta_B)$$

Given

For Θ_A

`y_A=c(12, 9, 12, 14, 13, 13, 15, 8, 15, 6)`

`n_yA=length(y_A)`

`Prior_a_A=120`

`Prior_b_A=10`

For Θ_B

`y_B=c(11,11,10,9,9,8,7,10,6,8,8,9,7)`

`prior_a_B=12`

`prior_b_B=1`

`n_B=13`

Code block

```
#3.3 a
#For Theta A

y_A=c(12, 9, 12, 14, 13, 13, 15, 8, 15, 6)
n_yA=length(y_A)
Prior_a_A=120
Prior_b_A=10

posterior_mean_A=(Prior_a_A+sum(y_A))/(Prior_b_A+n_yA)
posterior_Var_A=(Prior_a_A+sum(y_A))/((Prior_b_A+n_yA)^2)
ConfIntervalA=qgamma( c (.025, .975),Prior_a_A+sum(y_A),Prior_b_A+n_yA)

#For Theta B
y_B=c(11,11,10,9,9,8,7,10,6,8,8,9,7)
prior_a_B=12
prior_b_B=1
n_B=13

#Applying the calculations

posterior_mean_B=(prior_a_B+sum(y_B))/(prior_b_B+n_B)
posterior_Var_B=(prior_a_B+sum(y_B))/((prior_b_B+n_B)^2)
```

```

ConfintervalB=qgamma( c (.025, .975),prior_a_B+sum(y_B),prior_b_B+n_B)

#Reporting results for theta A and theta B

cat(sprintf("theta A \n posterior_mean_A= %f \n posterior_Var_A= %f \n Confin
tervalA= %f \n" , posterior_mean_A,posterior_Var_A,ConfIntervalA ))

## theta A
## posterior_mean_A= 11.850000
## posterior_Var_A= 0.592500
## ConfintervalA= 10.389238
## theta A
## posterior_mean_A= 11.850000
## posterior_Var_A= 0.592500
## ConfintervalA= 13.405448

cat(sprintf(" theta B \n posterior_mean_B= %f \n posterior_Var_B= %f \n Confi
ntervalB= %f \n" , posterior_mean_B,posterior_Var_B,ConfintervalB ))

## theta B
## posterior_mean_B= 8.928571
## posterior_Var_B= 0.637755
## ConfintervalB= 7.432064
## theta B
## posterior_mean_B= 8.928571
## posterior_Var_B= 0.637755
## ConfintervalB= 10.560308

```

b)

Compute and plot the posterior expectation of Θ_B under the prior distribution

$\Theta_B \sim \text{gamma}(12 \times n_0, n_0)$

for each value of $n_0 \in \{1, 2, \dots, 50\}$.

Describe what sort of prior beliefs about Θ_B would be necessary in order for the posterior expectation of Θ_B to be close to that of Θ_A .

```

#For Theta A

y_A=c(12, 9, 12, 14, 13, 13, 15, 8, 15, 6)
n_yA=length(y_A)
Prior_a_A=120
Prior_b_A=10

posterior_mean_A=(Prior_a_A+sum(y_A))/(Prior_b_A+n_yA)

# n0 C {1,2,.....,50}

```

```

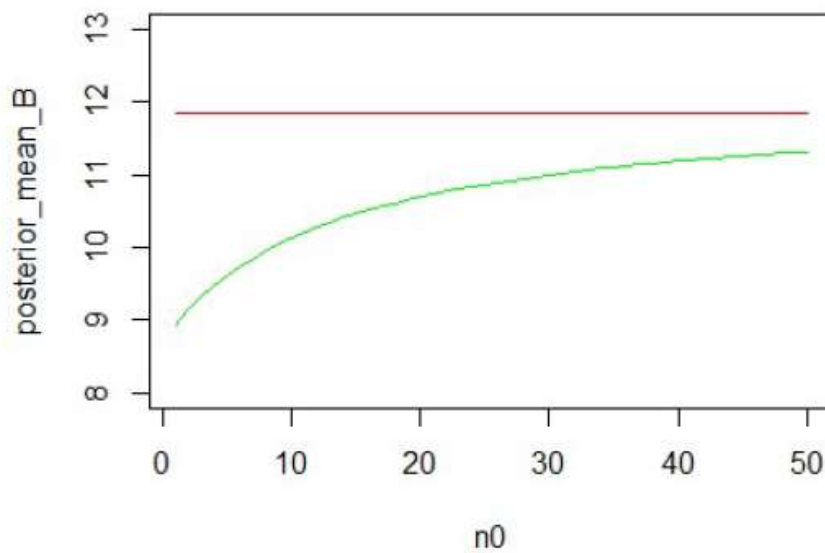
# For theta B
n0=seq(1,50)
prior_a_B=12*n0
prior_b_B=n0
y_B=c(11,11,10,9,9,8,7,10,6,8,8,9,7)
n_B=length(y_B)

sum_y_B=sum(y_B)

posterior_mean_B = (prior_a_B+sum_y_B)/(prior_b_B+n_B)

plot(posterior_mean_B, type = "l", col="green" , ylim = c(8,13) ,xlab="n0")
lines(n0,as.vector(rep(posterior_mean_A,50)),type = "l", col="red", xlab="n0"
)

```



The graph above shows the plot of posterior expectation of B. From the figure we see that the posterior expectation of B is approaching to that of A. For The posterior expectation of B to be close to that of A , n_0 should proceed further to 50. Or for the posterior expectation to approach to that of A it should have prior beliefs like A, when $n_0 = 10$ the prior belief of B is similar to that of A.

c)

Should knowledge about population A tell us anything about population B?

Discuss whether or not it makes sense to have

$$p(\Theta_A, \Theta_B) = p(\Theta_A) \times p(\Theta_B).$$

From the problem we understand that

A and B are independent, and that prior knowledge about the population A will have nothing to influence to that of B. Hence it makes sense for us to have

$$p(\Theta_A, \Theta_B) = p(\Theta_A) \times p(\Theta_B).$$

3.4)

Mixtures of beta priors: Estimate the probability of teen recidivism based on a study in which there were $n = 43$ individuals released from incarceration and $y = 15$ re-offenders within 36 months.

- a) Using a $\text{beta}(2,8)$ prior for Θ , plot $p(\Theta)$, $p(y|\Theta)$ and $p(\Theta|y)$ as functions of Θ . Find the posterior mean, mode, and standard deviation of Θ . Find a 95% quantile-based confidence interval.

Given

$$a=2, b=8, y=15 \text{ and } n=43$$

Objective

To plot $p(\Theta)$, $p(y|\Theta)$ and $p(\Theta|y)$ as functions of Θ

The beta distribution

$$p(\Theta) = \text{dbeta}(\Theta; a; b) = \text{dbeta}(\Theta; 2; 8)$$

The binomial distribution

A random variable Y has a $\text{binomial}(n, \Theta)$ distribution if

$$\Pr(Y = y|\Theta) = \text{dbinom}(y; n; \Theta) \quad p(y|\Theta) = \text{dbinom}(15; 43; \Theta)$$

The posterior distribution

$$p(\Theta | y) = \text{dbeta}(\Theta; a + y; b + ny) = \text{dbeta}(\Theta; 17; 36)$$

b)

Now for beta(8,2) prior for Θ

So by changing the values to a=8, b=2, we get

$$p(\Theta) = \text{dbeta}(\Theta, a, b) = \text{dbeta}(\Theta; 8; 2)$$

$$p(y | \Theta) = \text{dbinom}(n, y, \Theta) = \text{dbinom}(15, 43, \Theta)$$

$$p(\Theta | y) = \text{dbeta}(\Theta, a + y, b + ny) = \text{dbeta}(\Theta, 17, 36)$$

c)

We consider the following prior distribution

$$P(\Theta) = \frac{1}{4} \frac{\Gamma(10)}{\Gamma(2)\Gamma(8)} [3\Theta (1-\Theta)^7 + \Theta^7 (1-\Theta)] \quad \text{-----}(1)$$

Considering that (1) is a 75-25 % mixture of a beta(2,8) and beta (8,2) prior distribution

We can write (1) as

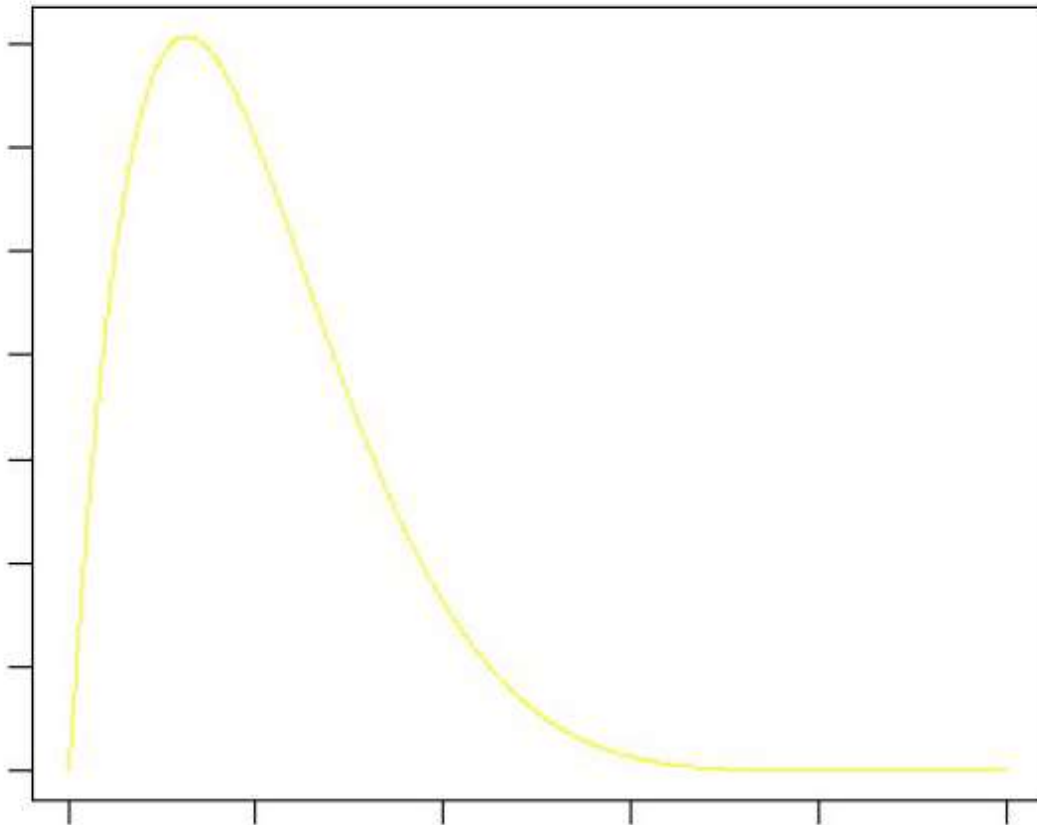
$$\begin{aligned} P(\Theta) &= \frac{3}{4} \frac{\Gamma(10)}{\Gamma(2)\Gamma(8)} [3\Theta (1-\Theta)^7] + \frac{1}{4} \frac{\Gamma(10)}{\Gamma(2)\Gamma(8)} [\Theta^7 (1-\Theta)] \\ &= 75 \% \text{ of beta } (2,8) \text{ prior} + 25 \% \text{ of beta}(8,2) \text{ prior} \end{aligned}$$

CodeBlock

```
par("mar") #Checking margins
## [1] 5.1 4.1 4.1 2.1

par(mar=c(1,1,1,1)) #change margins

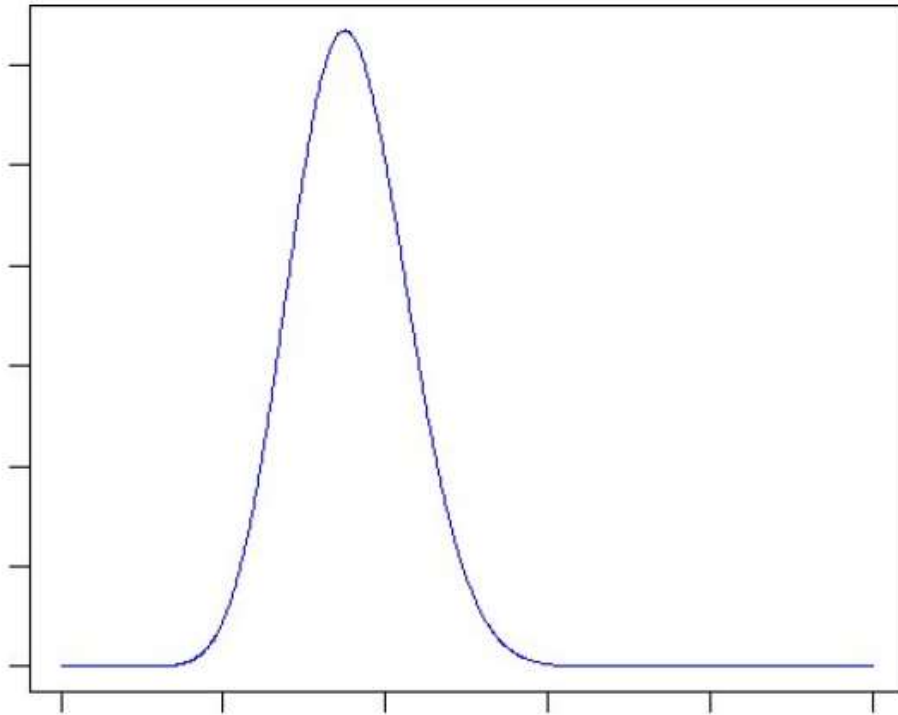
a<-2
b<-8
y<-15
n<-43
theta<-seq(0,1,.001)
pr <- dbeta(theta, a, b)
plot(pr,type='l', col="yellow")
```



x- label – Theta

y – label - Density

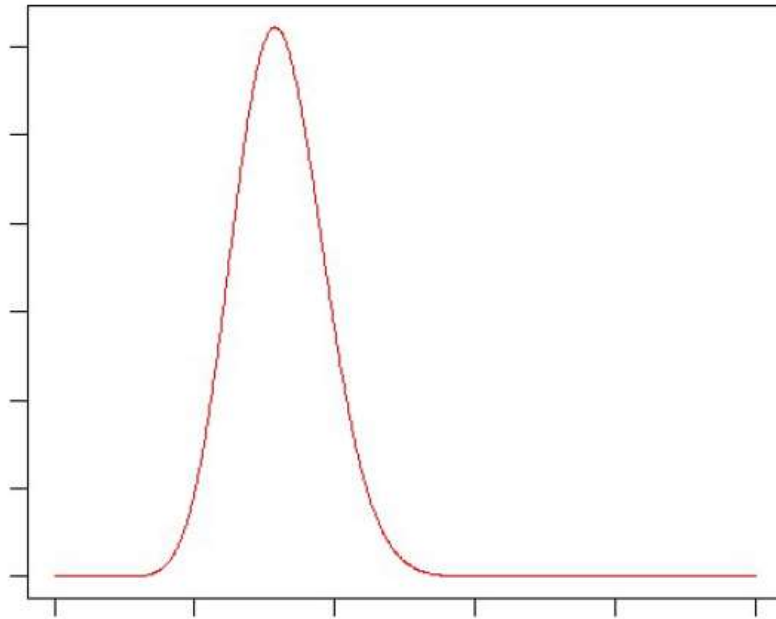
```
lh <- dbinom(y,n,theta)
plot(lh,type='l', col="blue")
```



x- label - Theta

y - label - Density

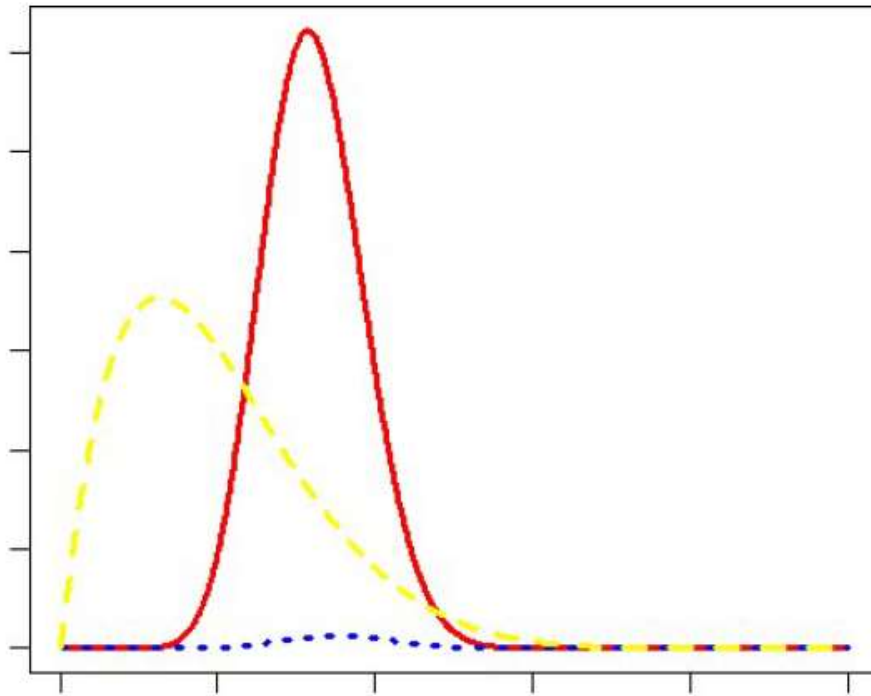
```
pos <- dbeta(theta, a+y, b+n-y)
plot(pos,type='l', col="red")
```

x- label - Theta

y - label - Density

```
#Plotting against  
plot(pos, col = "red", type='l', ylab = "Density", xlab = "theta", lty = 1, lwd  
= 3)  
lines(lh, type='l', col="blue", lty = 3, lwd = 3)  
lines(pr, col = "yellow", lty = 2, lwd = 3)
```

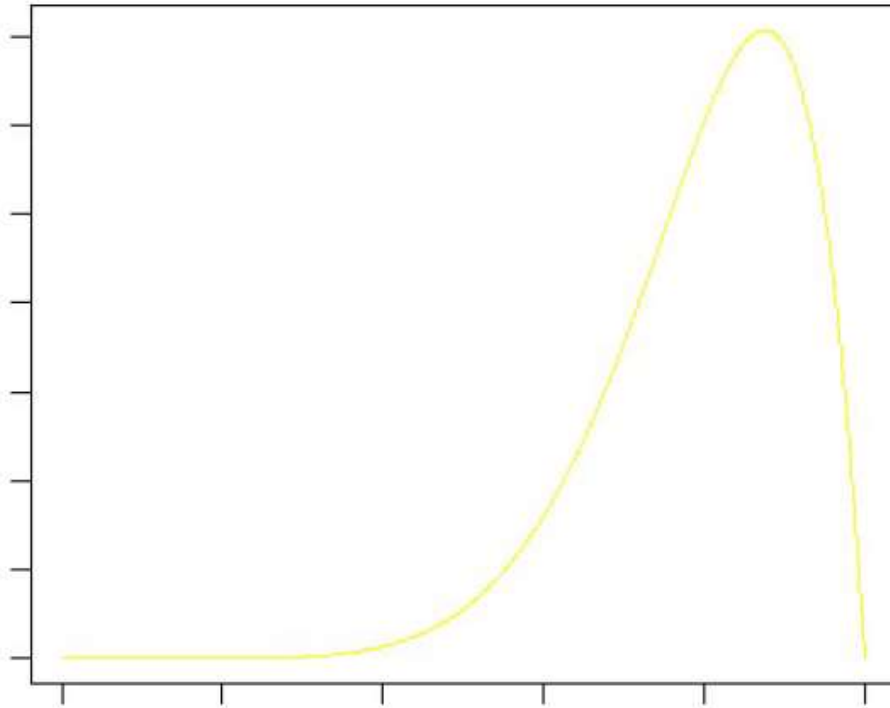


x- label - Theta

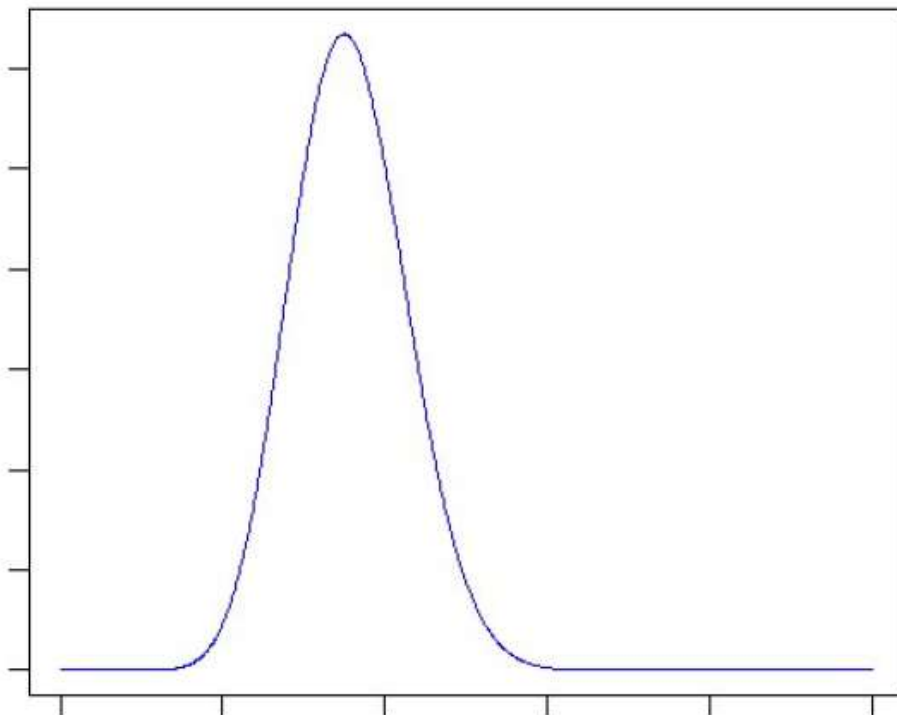
y - label - Density

```
a<-8
b<-2
y<-15
n<-43
theta<-seq(0,1,.001)
pr <- dbeta(theta, a, b)
lh <- dbinom(y,n,theta)
pos <- dbeta(theta, a+y, b+n-y)

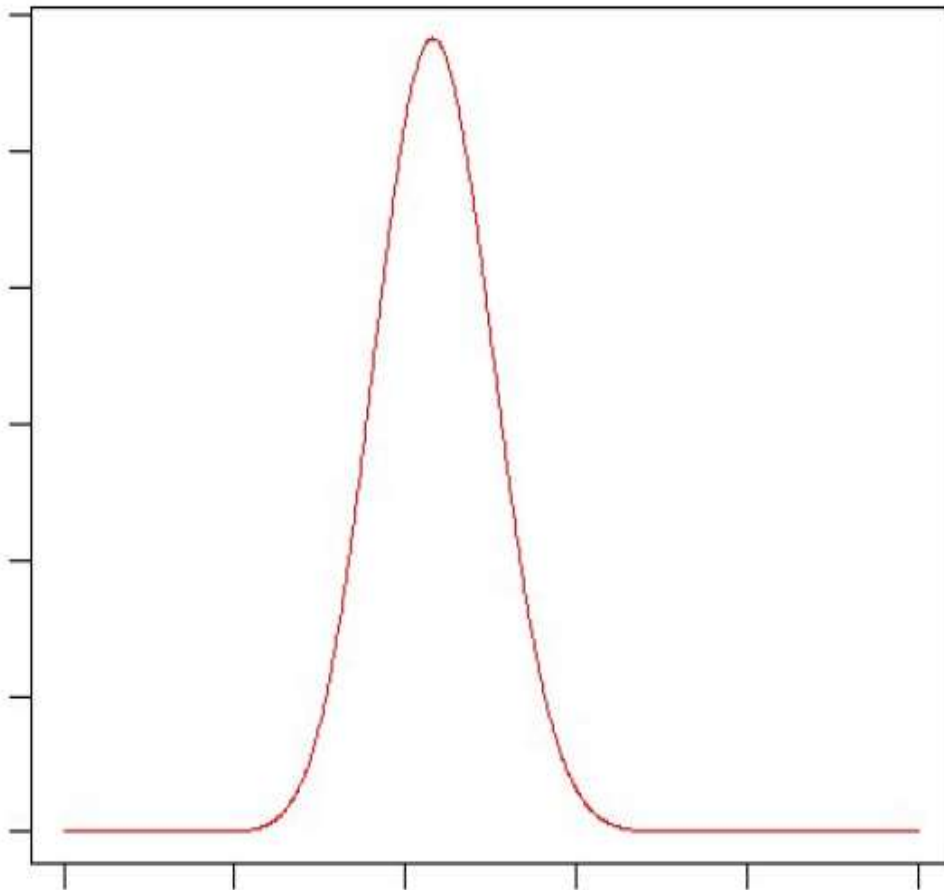
plot(pr,type='l', col="yellow")
```



```
plot(lh,type='l', col="blue")
```



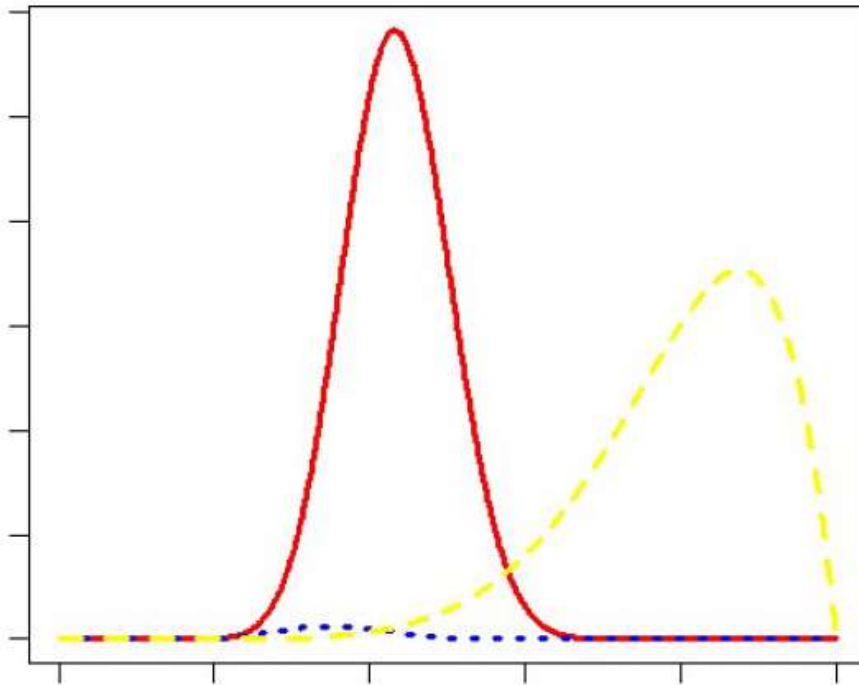
```
plot(pos,type='l', col="red")
```



x-label - Theta

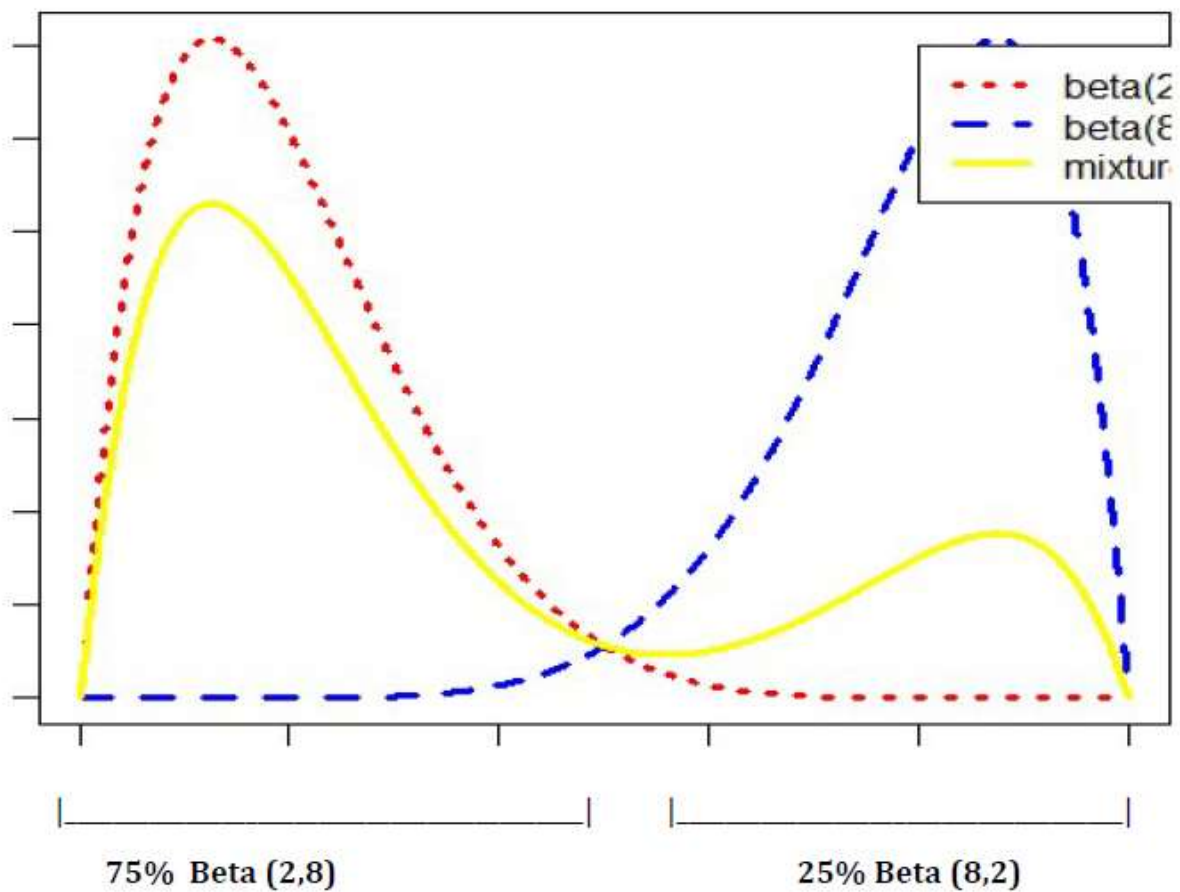
y-label - Density

```
plot(pos, col = "red", type='l', ylab = "Density", xlab = "theta", lty = 1, lwd
= 3)
lines(lh, type='l', col="blue", lty = 3, lwd = 3)
lines(pr, col = "yellow", lty = 2, lwd = 3)
legend(x=1500,y=5.0, c("Prior", "Likelihood", "Posterior"), lty = c(2, 3, 1),
      lwd = c(3, 3, 3), col = c("yellow", "blue", "red"))
```



```
a<-2
b<-8
y<-15
n<-43
theta<-seq(0,1,.001)
pr1 <- dbeta(theta, a, b)
a<-8
b<-2
y<-15
n<-43
theta<-seq(0,1,.001)
pr2 <- dbeta(theta, a, b)

plot(pr1, type='l', col="red", lty = 3, lwd = 3, ylab = "Density", xlab = "theta")
lines(pr2, col = "blue", lty = 2, lwd = 3)
lines((((3/4)*pr1)+((1/4)*pr2)),type='l', col="yellow", lty = 1, lwd = 3)
legend(x=800,y=3.5, c("beta(2,8) prior", "beta(8,2) prior", "mixture prior"),
lty = c(3, 2, 1),
      lwd = c(3, 3, 3), col = c("red","blue","yellow"))
```



x- label - Theta

y - label - Density

From the above plot we can see that there is a clear inclination in the mixture curve towards 75% Beta(2,8). This represents that the density is higher on the side with the greater concentration of theta.

d)

From equation (1) we know that

$$P(\Theta) = \frac{1}{4} \frac{\Gamma(10)}{\Gamma(2)\Gamma(8)} [3\Theta (1-\Theta)^7 + \Theta^7 (1-\Theta)] \text{ -----(3)}$$

$$P(y|\Theta) = \Theta^y (1-\Theta)^{n-y}$$

$$P(y|\Theta) = \Theta^7 (1-\Theta)^{28} \text{ -----(2)}$$

From (3) and (2)

$$P(\Theta) \times P(y|\Theta) = \frac{1}{4} \frac{\Gamma(10)}{\Gamma(2)\Gamma(8)} [3\Theta^{16} (1-\Theta)^{35} + \Theta^{22} (1-\Theta)^{29}]$$

ii)

We know

$Y \in \text{binomial}(n, \Theta)$

Hence the binomial distribution $\rightarrow P(y|\Theta) = \Theta^y (1-\Theta)^{n-y}$

And also the beta prior $P(\Theta) = d \text{ beta}(\Theta, a, b)$. These identify the distributions

iii)

Plotting $p(\Theta) \times p(y|\Theta)$ for a variety of values

Code Block

```
attach(mtcars)
par(mfrow=c(2,2))
```

```
a<-2
b<-8
y<-15
n<-43
theta<-seq(0,1,.001)
pr1 <- dbeta(theta, a, b)
```

```
a<-8
b<-2
y<-15
n<-43
theta<-seq(0,1,.001)
pr2 <- dbeta(theta, a, b)
```

```
lh <- dbinom(y,n,theta)
plot( ( (lh)*(((3/4)*pr1)+((1/4)*pr2)) ),type='l', col="yellow", lty = 1, lwd
= 3,ylab = "Density", xlab = "theta")
```

```
a<-2
b<-8
y<-15
n<-43
theta<-seq(0,10,.005)
pr1 <- dbeta(theta, a, b)
a<-8
b<-2
y<-15
n<-43
theta<-seq(0,10,.005)
pr2 <- dbeta(theta, a, b)
lh <- dbinom(y,n,theta)
```

```
## Warning in dbinom(y, n, theta): NaNs produced
```

```
plot( ( (lh)*(((3/4)*pr1)+((1/4)*pr2)) ),type='l', col="yellow", lty = 1, lwd
= 3,ylab = "Density", xlab = "theta")
```



```

a<-2
b<-8
y<-15
n<-43
theta<-seq(0,1,.004)
pr1 <- dbeta(theta, a, b)
a<-8
b<-2
y<-15

```

```

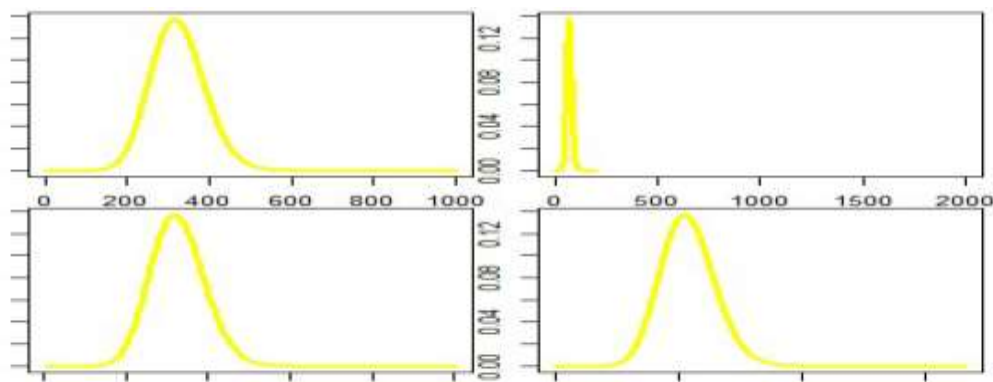
n<-43
theta<-seq(0,1,.004)
pr2 <- dbeta(theta, a, b)
lh <- dbinom(y,n,theta)
plot( ( (lh)*(((3/4)*pr1)+((1/4)*pr2)) ),type='l', col="yellow", lty = 1, lwd
= 3,ylab = "Density", xlab = "theta")

```

```

a<-2
b<-8
y<-15
n<-43
theta<-seq(0,1,.00006)
pr1 <- dbeta(theta, a, b)
a<-8
b<-2
y<-15
n<-43
theta<-seq(0,1,.00006)
pr2 <- dbeta(theta, a, b)
lh <- dbinom(y,n,theta)
plot( ( (lh)*(((3/4)*pr1)+((1/4)*pr2)) ),type='l', col="yellow", lty = 1, lwd
= 3,ylab = "Density", xlab = "theta")

```



X label - Theta
Y label- Density

e)

The weights of the mixture distribution from ii) is given by their sums

Interpreting their values shows that the value for density is higher for greater concentration of theta.

Problem 3

Hoff 3.7 3.7 Posterior prediction: Consider a pilot study in which $n_1 = 15$ children enrolled in special education classes were randomly selected and tested for a certain type of learning disability. In the pilot study, $y_1 = 2$ children tested positive for the disability.

a) Using a uniform prior distribution, find the posterior distribution of θ , the fraction of students in special education classes who have the disability.

Find the posterior mean, mode and standard deviation of θ , and plot the posterior density.

[Solution : Given: $n_1 = 15$ & $y_1 = 2$

and we know that for $\theta \sim \text{beta}(1, 1)(\text{uniform})$,

$$(\theta|Y = y) \sim \text{beta}(1 + y, 1 + n - y)$$

to get the likelihood, assuming this is an i.i.d

$$Pr(Y = y|\theta) = \theta^y (1 - \theta)^{n-y} = \text{dbinom}(y, n, \theta) = \text{dbinom}(2, 15, \theta) \text{ which is a binomial distribution that gives the}$$

From the prior inference condition of the uniform distribution, we know that

$$p(\theta|Y) = \text{beta}(1 + y, 1 + n - y) = \text{beta}(3, 14)$$

The above equations demonstrates that for any i.i.d prior, the $\sum y_i$ is the sufficient data to predict the posterior distribution.

Posterior Mean: It is given by the formula

$$E[\theta|y] = \frac{a + y}{a + b + n} = \frac{3}{17} = 0.1764$$

Posterior Mode:

$$\text{mode}[\theta|y] = \frac{a + y - 1}{a + b + n - 2} = \frac{2}{15} = 0.1333$$

Posterior Standard Deviation:

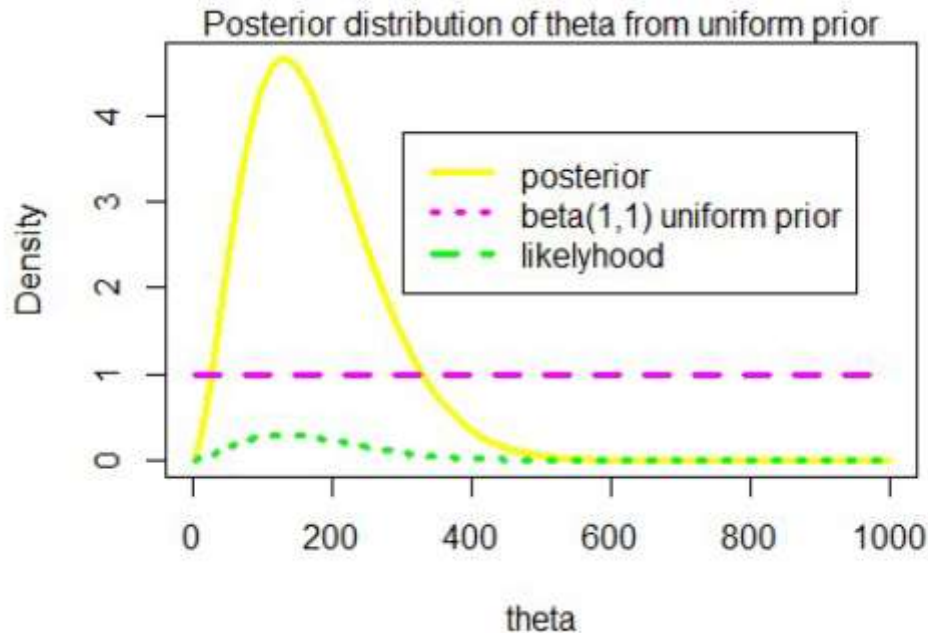
$$Var[\theta|y] = \frac{E[\theta|y] E[1 - \theta|y]}{a + b + n + 1} =$$

$$[0.176 * (1 - (3/17))] / 18 = 0.0080568$$

$$Sd = 0.0049$$

Codeblock

```
#3.7 a)
a=3
b=14
theta = seq(0.005, 0.995, length = 1000)
posterior = dbeta(theta, a, b)
prior = dbeta(theta,1,1)
likelihood = dbinom(2,15,theta)
plot(posterior, col = "Yellow",typ='l', ylab = "Density", xlab = "theta",lty
= 1, lwd = 3)
lines(prior, col = "magenta", lty = 2, lwd = 3)
lines(likelihood, col = "green", lty = 3, lwd = 3)
legend(x=300,y=3.8, c("posterior","beta(1,1) uniform prior","likelihood"), lt
y = c(1, 3, 2),
      lwd = c(3, 3, 3), col = c( "Yellow","Magenta","green"))
mtext("Posterior distribution of theta from uniform prior")
```



Researchers would like to recruit students with the disability to participate in a long-term study, but first they need to make sure they can recruit enough students. Let $n_2 = 278$ be the number of children in special education classes in this particular school district, and let Y_2 be the number of students with the disability.

- b) Find $Pr(Y_2 = y_2 | Y_1 = 2)$, the posterior predictive distribution of Y_2 , as follows:
- Discuss what assumptions are needed about the joint distribution of (Y_1, Y_2) such that the following is true:

[Solution :

$$Pr(Y_2 = y_2 | Y_1 = 2) = \int_0^1 Pr(Y_2 = y_2 | \theta) p(\theta | Y_1 = 2) d\theta$$

Using class room Notations:

For this to be true, in general when $Y = y_1, \dots, y_n$ are considered in the denominator of $p(y \sim / Y)$, then $Y = y_1, \dots, y_n$ should follow iid. And y_1, \dots, y_n are conditionally independent.

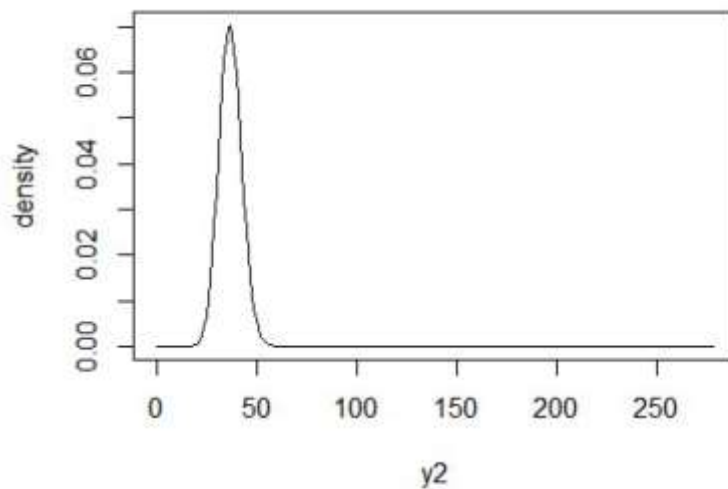
But here we are asked about, $y_2 = y \sim$ and $Y = y_i$. SO this makes y_1 and y_2 are dependent. Because, y_2 dependent is dependent on θ and θ is dependent on y_1 .

Based on what text book says below: y_1 and y_2 are dependent.

1. The predictive distribution does not depend on any unknown quantities. If it did, we would not be able to use it to make predictions.
2. The predictive distribution depends on our observed data. In this distribution, $y \sim$ is not independent of Y_1, \dots, Y_n (recall Section 2.7). This is because observing Y_1, \dots, Y_n gives information about θ , which in turn gives information about $y \sim$. It would be bad if $y \sim$ were independent of Y_1, \dots, Y_n - it would mean that we could never infer anything about the un-sampled population from the sample cases.

d)

```
mean<-278 * 2 /15
sd=sqrt(mean * (1 - (2/15) ))
y2=seq(0,278)
plot(y2,dbinom(x = y2, size = 278, prob = (2/15)), ylab = "density", type="l"
)
```



```
print(paste0("mean = ",mean," sd = ",sd ))
## [1] "mean = 37.0666666666667 sd = 5.66784301515528"
```

inference: I will use the posterior calculated in c for this. At lower sample size, the MLE estimates are not good in predictions. With increase in sample size