

Viterbo_Giuseppe_rlab03

April 28, 2023

```
[1]: install.packages('tidyverse')
install.packages('gridExtra')
install.packages('emdbook')

library(tidyverse)
library(gridExtra)
library(emdbook)
```

Updating HTML index of packages in '.Library'

Making 'packages.html' ...
done

Updating HTML index of packages in '.Library'

Making 'packages.html' ...
done

also installing the dependencies 'bdsmatrix', 'mvtnorm', 'coda', 'bbmle'

Updating HTML index of packages in '.Library'

Making 'packages.html' ...
done

Attaching core tidyverse packages

```
tidyverse 2.0.0
dplyr      1.1.1    readr      2.1.4
forcats    1.0.0    stringr    1.5.0
ggplot2    3.4.2    tibble     3.2.1
lubridate  1.9.2    tidyr      1.3.0
purrr      1.0.1
Conflicts
```

```
tidyverse_conflicts()
dplyr::filter() masks stats::filter()
dplyr::lag() masks stats::lag()
Use the conflicted package
(<http://conflicted.r-lib.org/>) to force all conflicts to
become errors
```

Attaching package: ‘gridExtra’

The following object is masked from ‘package:dplyr’:

```
combine
```

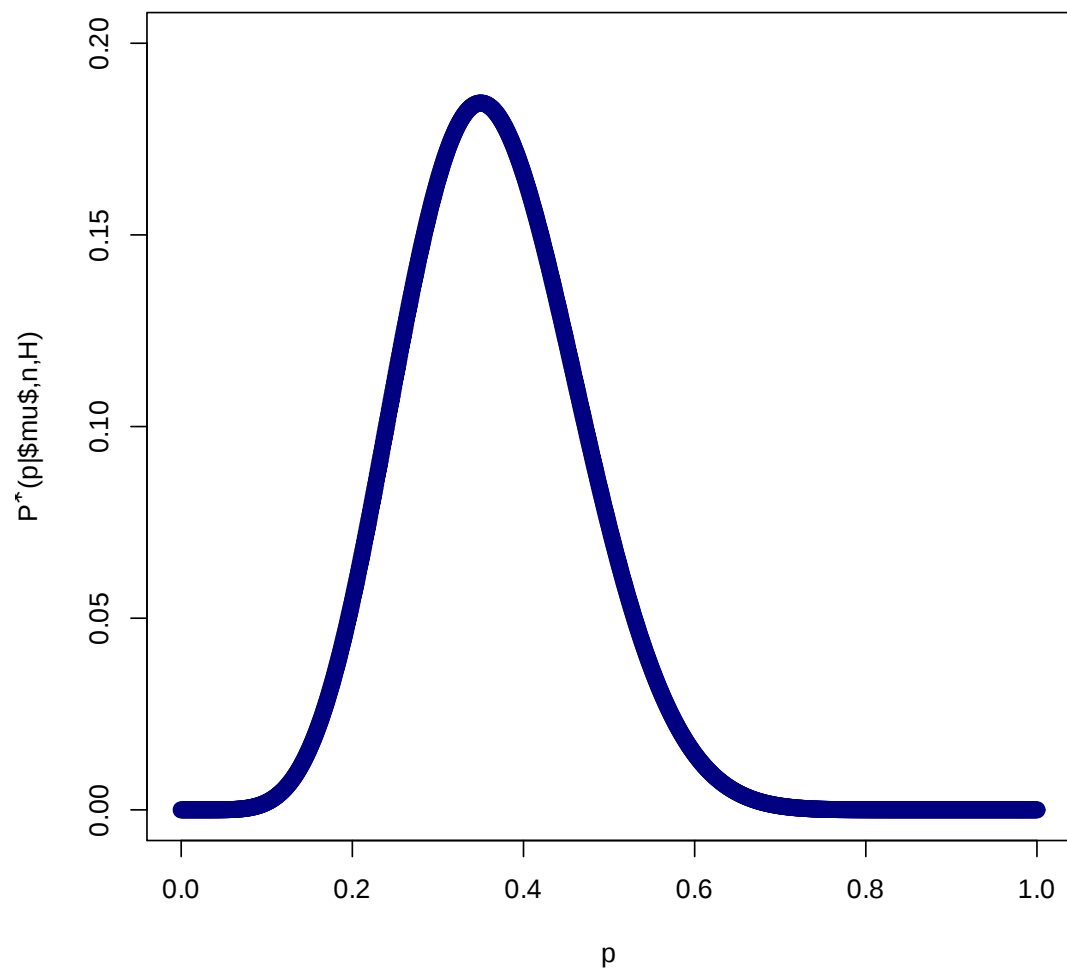
1 Ex.1

- study the binomial inference for a study that reports $y = 7$ successes in $n = 20$ independent trial. Assume the following priors:
 - a uniform distribution
 - a Jeffrey’s prior
 - a step function:
- plot the posterior distribution and summerize the results computing the first two moments
- compute a 95% credibility interval and give the results in a summary table
- draw the limits on the plot of the posterio distribution

1.0.1 Uniform distribution prior

the normalization $P(\pi|H)$ dosen’t depend on π

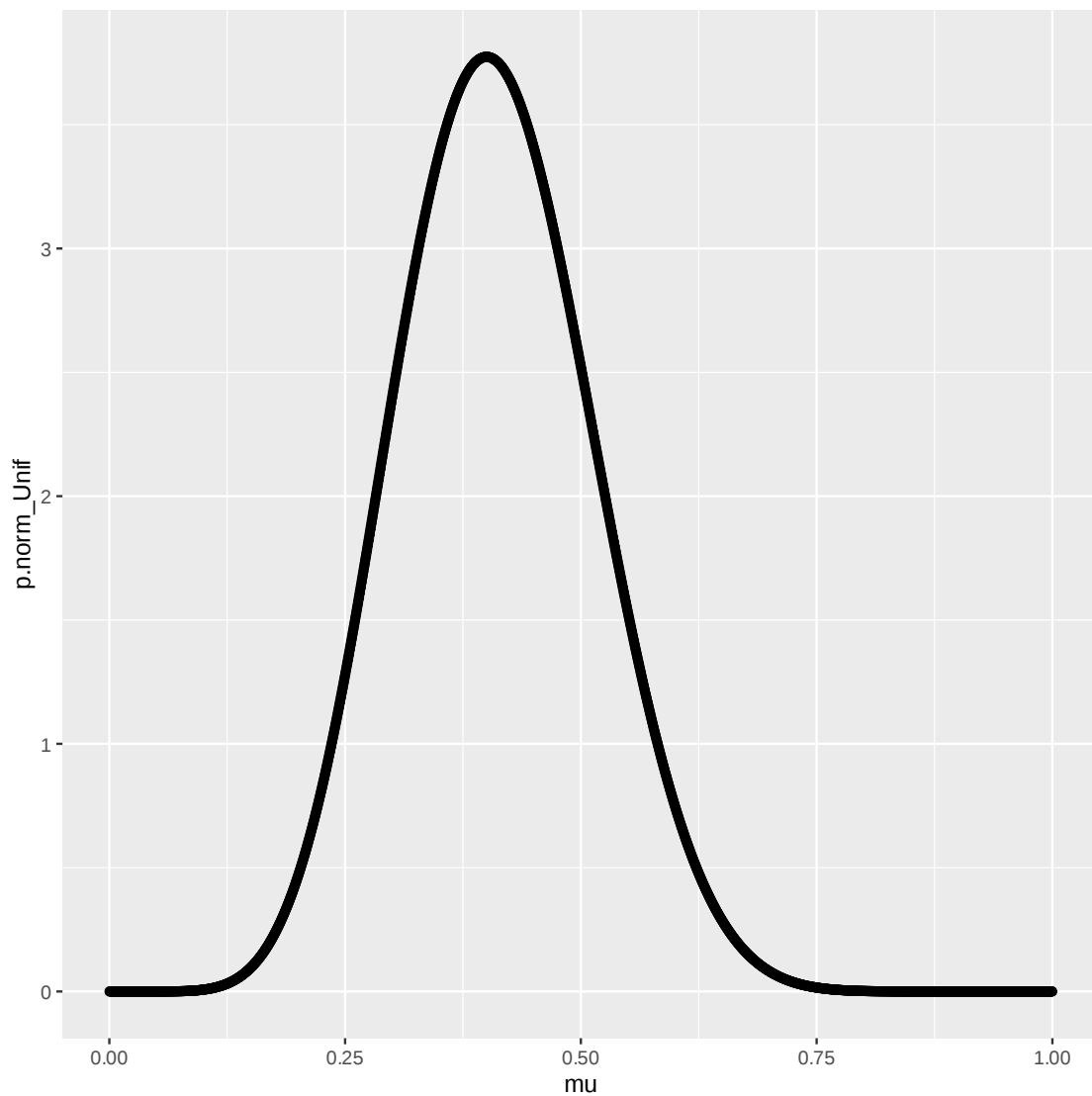
```
[5]: y <- 7    #success
n <- 20    #number of trial
mu <- seq(0,1, length.out=1000)
p.post_notnorm <- dbinom(x=y, size=n, prob=mu)
plot(mu, p.post_notnorm,
     col='navy', lty=1, lwd=3,
     ylim=c(0,0.2),
     xlab='p',
     ylab=expression(paste(P~symbol('*'), '(p|$mu$,n,H)')))
```



```
[7]: y <- 8
n <- 20;
n.sample <- 2000 #number of interval in which calculate the approximate
      →normalization function
delta.p <- 1/n.sample #length of the interval
mu <- seq(from = 1/(2*n.sample), by=1/n.sample, length.out = n.sample)
      →#possible chose of the p in the binomial distribution, called mu

p.star_Unif <- dbinom(x=y, size=n, prob=mu)
p.norm_Unif <- p.star_Unif/(delta.p * sum(p.star_Unif))
# plot(p, p.norm, col='red')
# points(p, p.star, col='navy')
ggplot() +
```

```
geom_point(aes(mu, p.norm_Unif))
```



1.0.2 Jeffrey's Prior

it is a prior invariant under any continuous transformation of the parameter: $g(\mu) \propto \frac{1}{\sqrt{\mu}}$ for $\mu > 0$

```
[8]: y <- 8
      n <- 20
      n.sample <- 2000
      delta.p <- 1/n.sample
      mu <- seq(from = 1/(2*n.sample), by=1/n.sample, length.out = n.sample)

      #p.star <- dgamma(mu, shape=(y+1/2), scale=n)
```

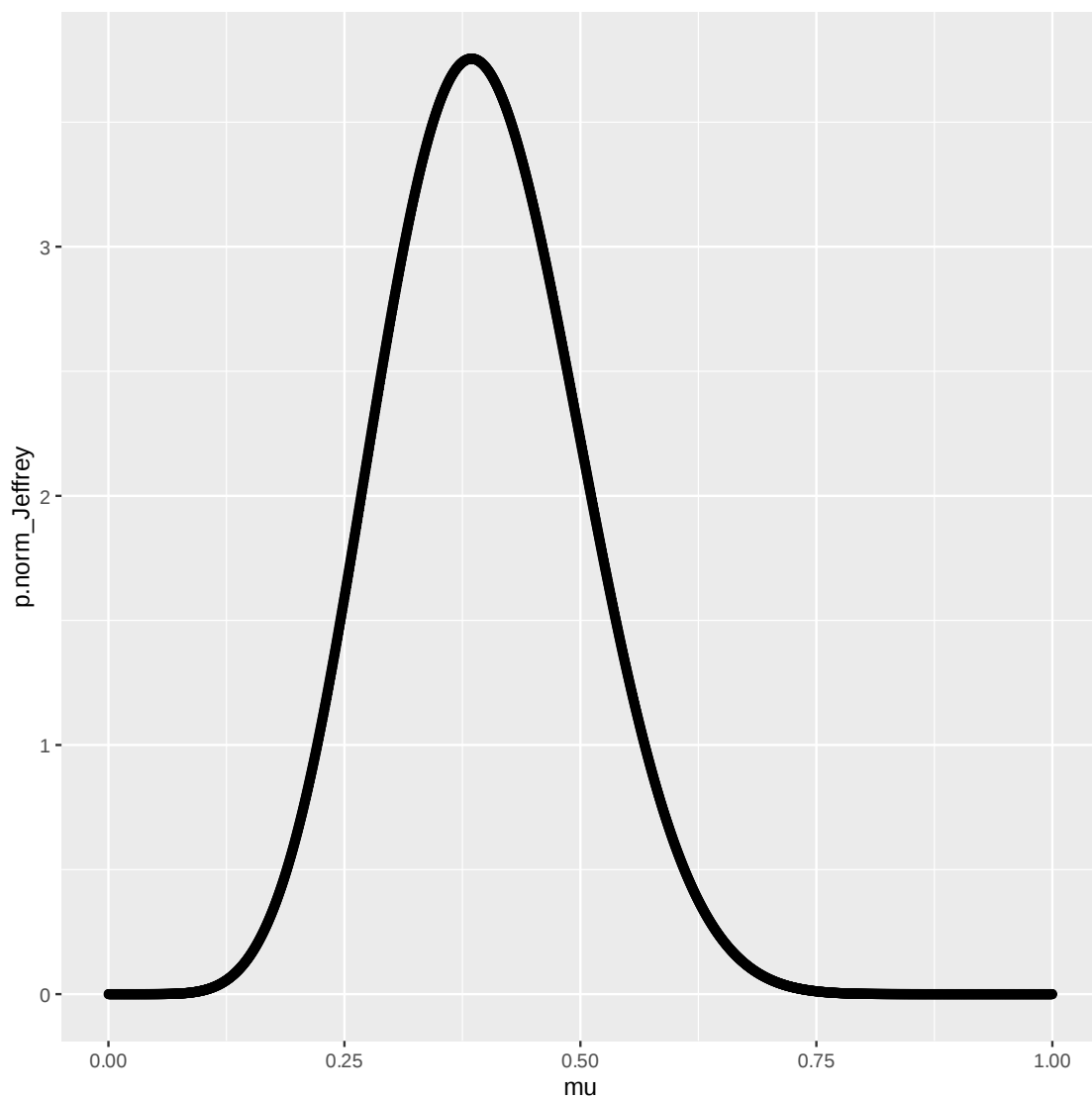
```

# p.norm <- p.star / (delta.p * sum(p.star))
#p.norm_Jeffrey <- p.star / (integrate(function(k) {dgamma(k, shape=(y+1/2),
↪scale=n)}, lower = 0, upper = 1))$value #return the same as the approximated
↪version

p.star_Jeffrey <- dbinom(x=y, size=n, prob=mu) * 1/((mu)**(1/2))
p.norm_Jeffrey <- p.star_Jeffrey / integrate(function(k) {dbinom(x=y, size=n,
↪prob=k) * 1/((k)**(1/2)) }, lower = 0, upper = 1)$value

ggplot() +
geom_point(aes(mu, p.norm_Jeffrey))

```

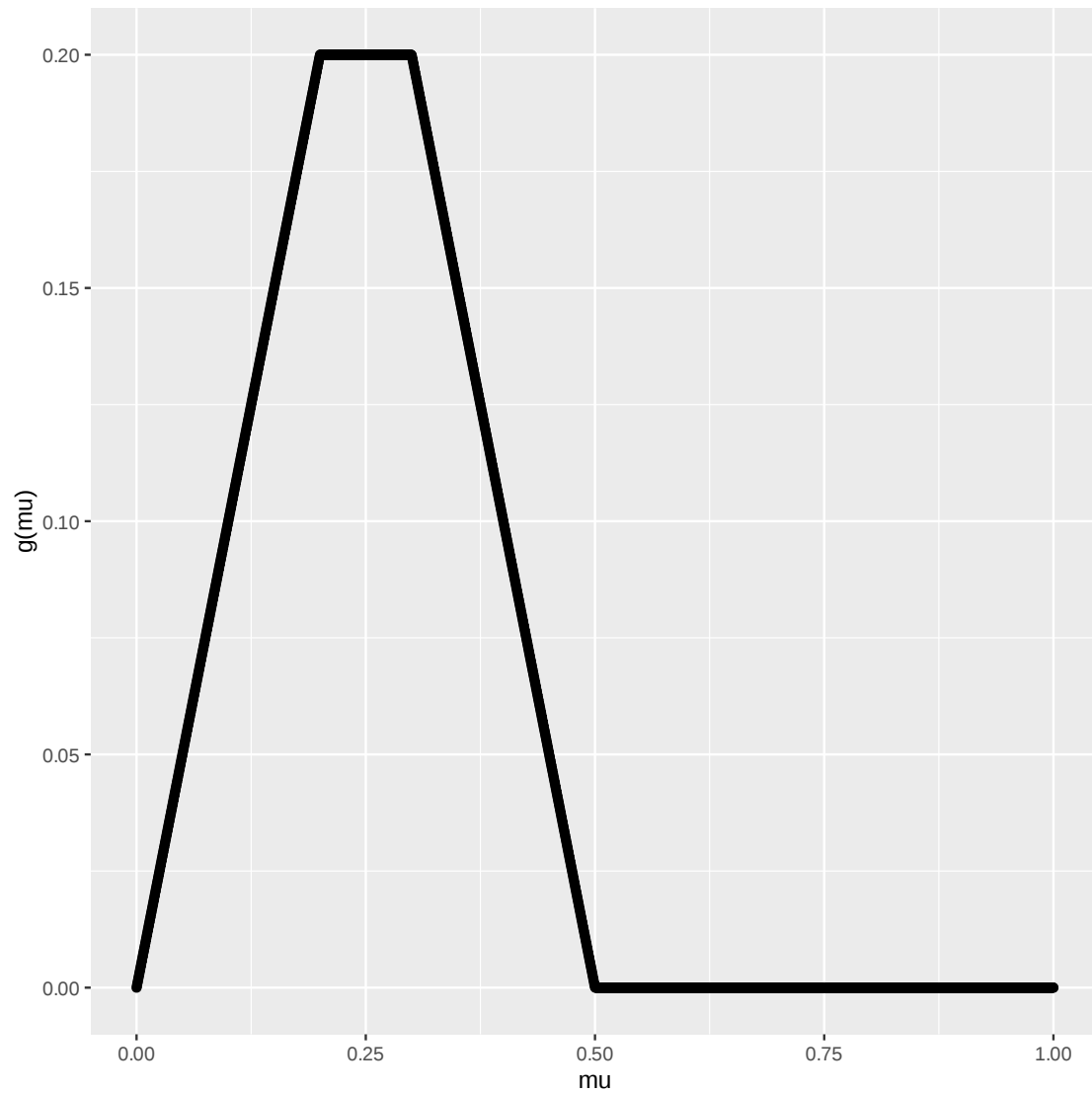


1.0.3 Step function

```
[9]: y <- 8
n <- 20
mu <- seq(0,1,length.out = 2000)

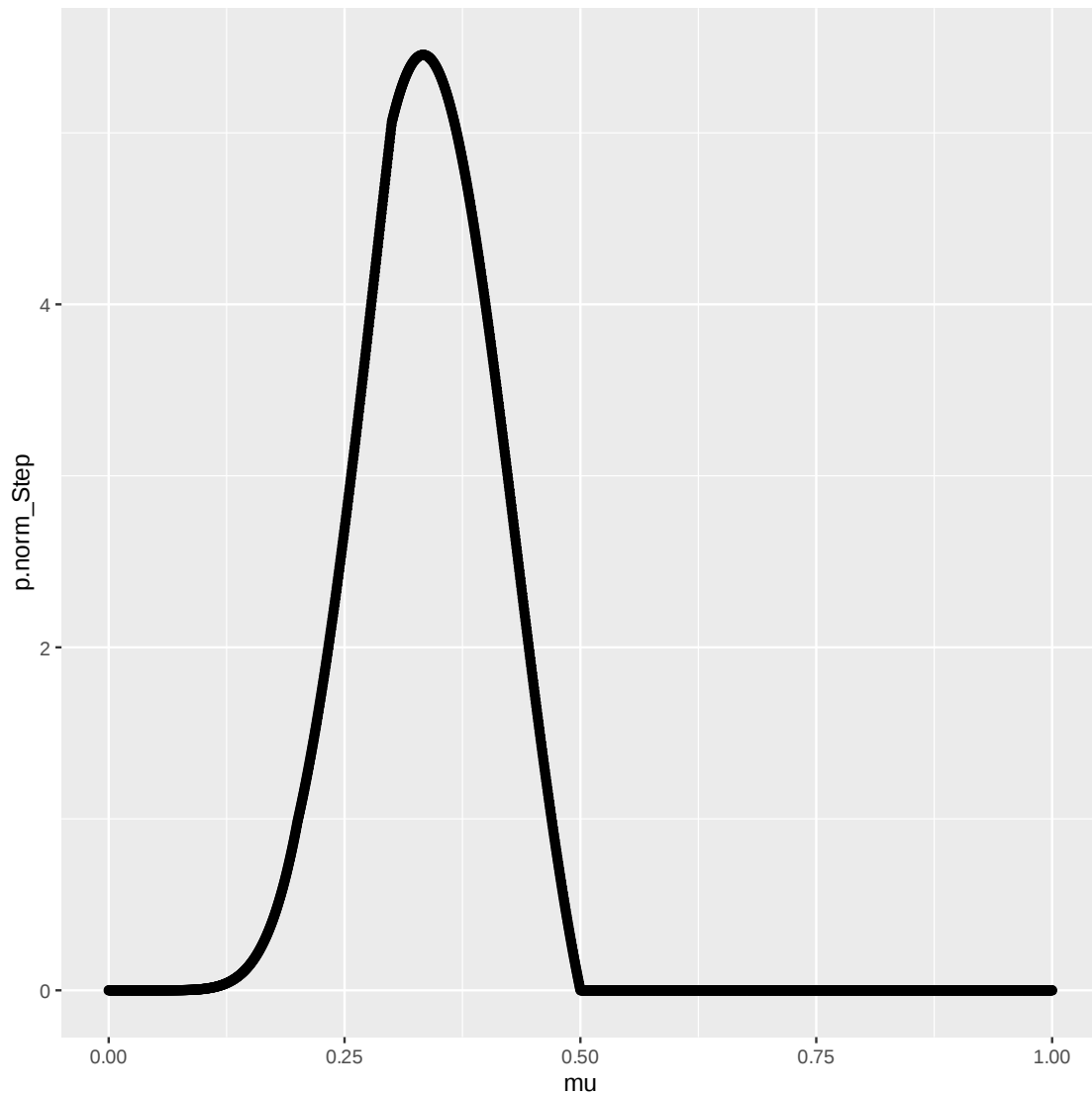
g <- function(k) {
  g.val <- ifelse(k <= 0.2,
    k,
    ifelse(k > 0.2 & k <= 0.3,
      0.2,
      ifelse(k > 0.3 & k <= 0.5,
        0.5 - k,
        ifelse(k > 0.5,
          0,
          0)
        )
      )
    )
  return(g.val)
}
```

```
[10]: ggplot()+
  geom_point(aes(mu, g(mu)))
```



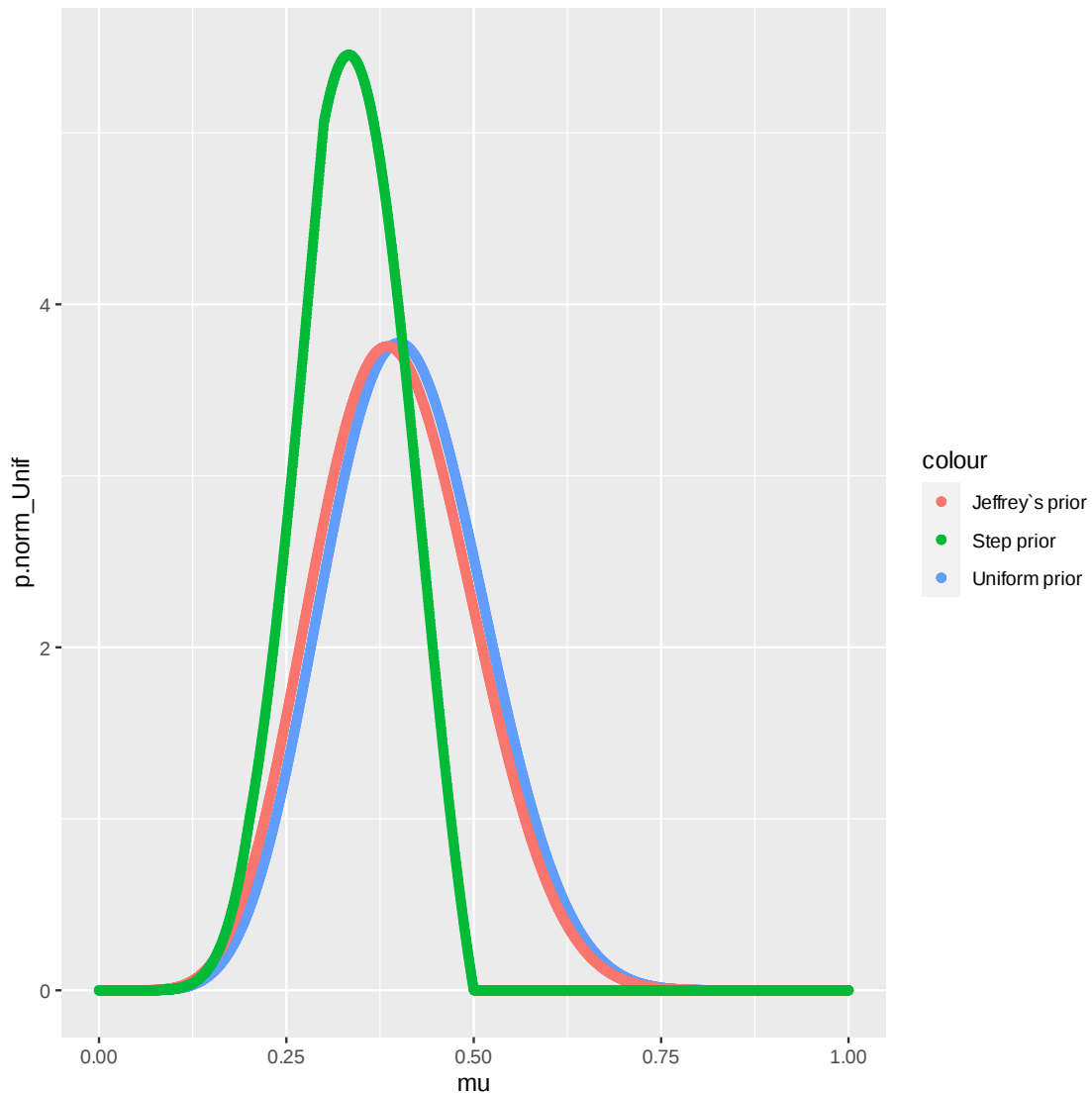
```
[11]: p.star_Step <- dbinom(x=y, size=n, prob=mu) * g(mu)
p.norm_Step <- p.star_Step / (integrate(function(k) {dbinom(x=y, size=n, prob=k)}
↳ * g(k)}, lower = 0, upper = 1))$value

ggplot() +
geom_point(aes(mu, p.norm_Step))
```



2 Plot Posterior distribution

```
[12]: ggplot() +  
  geom_point(aes(mu, p.norm_Unif, color='Uniform prior')) +  
  geom_point(aes(mu, p.norm_Jeffrey, color='Jeffrey`s prior')) +  
  geom_point(aes(mu, p.norm_Step, color='Step prior')) -> posterior_plot  
posterior_plot
```

2.1 First and Second Momenta

```
[172]: y <- 8; n <- 20; n.sample <- 2000; delta.mu <- 1/n.sample;
mu <- seq(from=1/(2*n.sample), by=1/n.sample, length.out = n.sample)

#Uniform prior
p.star_Unif<- dbinom(x=y, size=n, prob=mu) * dunif(mu, min=0, max=1)
p.norm_Unif <- p.star_Unif/(delta.mu * sum(p.star_Unif))
first_moment_Unif <- delta.mu * sum(mu * p.norm_Unif)
second_moment_Unif <- delta.mu * sum(((first_moment_Unif-mu)**2)*p.norm_Unif)

#Jeffrey prior
```

```

p.star_Jeffrey <- dbinom(x=y, size=n, prob=mu) * 1/(mu ** (1/2))
p.norm_Jeffrey <- p.star_Jeffrey / (delta.mu*sum(p.star_Jeffrey))
first_moment_Jeffrey <- delta.mu * sum(mu * p.norm_Jeffrey)
second_moment_Jeffrey <- delta.mu * sum(((first_moment_Jeffrey-mu)**2) * p.
  ↪norm_Jeffrey)

#Step prior
p.star_Step <- dbinom(x=y, size=n, prob=mu) * g(mu)
p.norm_Step <- p.star_Step / (delta.mu * sum(p.star_Step))
first_moment_Step <- delta.mu * sum(mu * p.norm_Step)
second_moment_Step <- delta.mu * sum(((first_moment_Step-mu)**2) * p.norm_Step)

#PLot
ggplot()+
geom_point(aes(mu, p.norm_Unif, color='Uniform Prior')) +
geom_vline(aes(xintercept=first_moment_Unif, color='Uniform Prior'),
  ↪linetype='dashed')+

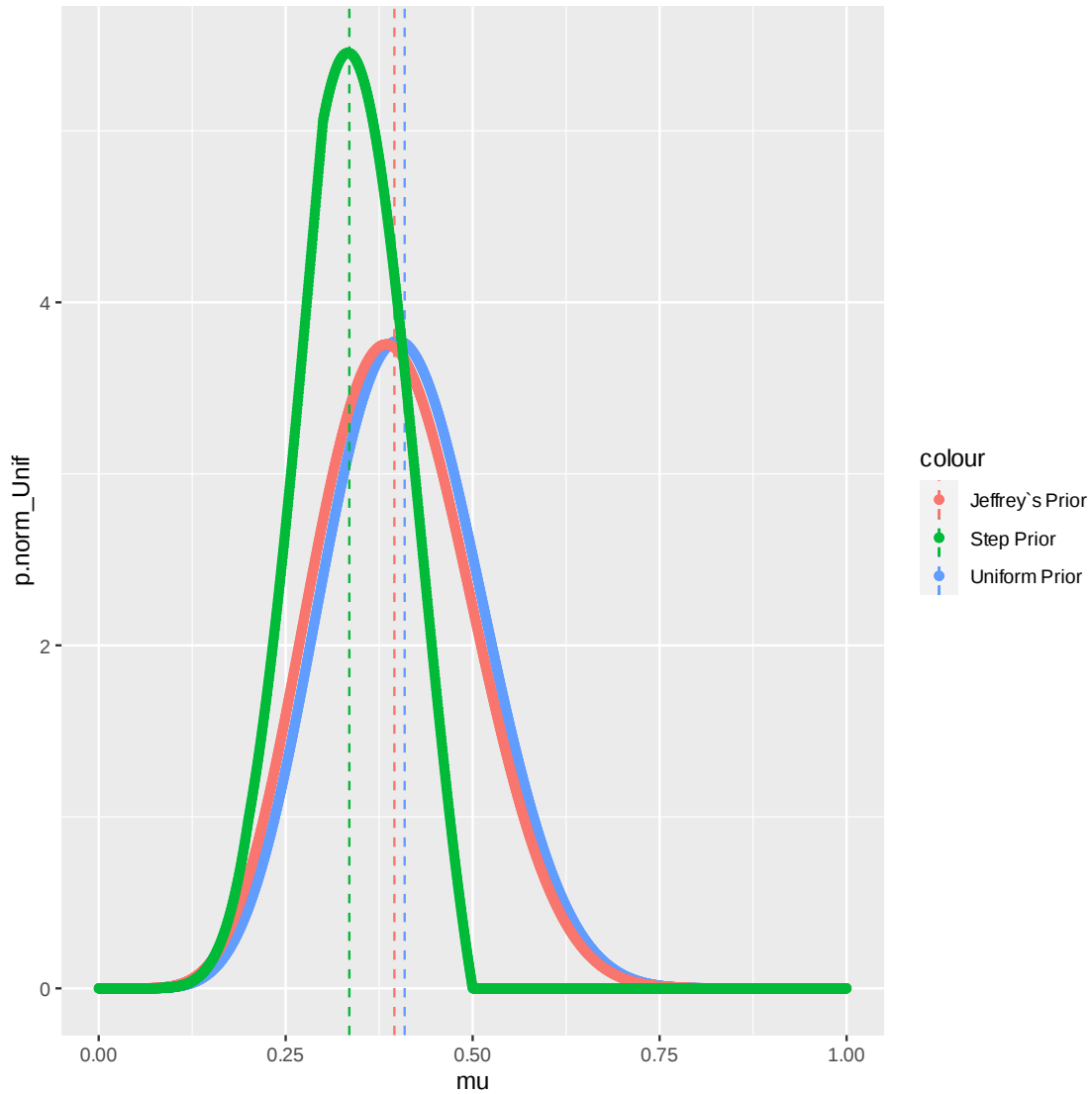
geom_point(aes(mu, p.norm_Jeffrey, color='Jeffrey`s Prior'))+
geom_vline(aes(xintercept=first_moment_Jeffrey, color='Jeffrey`s Prior'),
  ↪linetype='dashed') +

geom_point(aes(mu, p.norm_Step, color='Step Prior'))+
geom_vline(aes(xintercept=first_moment_Step, color='Step Prior'),
  ↪linetype='dashed')

Momenta <- tibble(
  Prior = c('Uniform', 'Jeffrey`s', 'Step'),
  First_momenta = c(first_moment_Unif, first_moment_Jeffrey,
  ↪first_moment_Step),
  Second_momenta = c(second_moment_Unif, second_moment_Jeffrey,
  ↪second_moment_Step),
)
Momenta

```

	Prior <chr>	First_momenta <dbl>	Second_momenta <dbl>
A tibble: 3 × 3	Uniform	0.4090909	0.010510241
	Jeffrey`s	0.3953488	0.010624362
	Step	0.3350930	0.004673032



3 Compute 95% credibility interval and Plot the limits

The function `ncredit` is used in order to compute the credibility interval

```
[147]: # Uniform prior
lower_bound_Unif <- ncredit(mu, p.norm_Unif, level=0.95)[['lower']]
upper_bound_Unif <- ncredit(mu, p.norm_Unif, level=0.95)[['upper']]

# Jeffrey prior
lower_bound_Jeffrey <- ncredit(mu, p.norm_Jeffrey, level=0.95)[['lower']]
upper_bound_Jeffrey <- ncredit(mu, p.norm_Jeffrey, level=0.95)[['upper']]
```

```

# Step prior
lower_bound_Step <- ncredint(mu, p.norm_Step, level=0.95)[['lower']]
upper_bound_Step <- ncredint(mu, p.norm_Step, level=0.95)[['upper']]

#Summary table
Credibility_Interval <- tibble(
  Prior = c('Uniform', 'Jeffrey', 'Step'),
  Lower_bound = c(lower_bound_Unif, lower_bound_Jeffrey,
    ↪lower_bound_Step),
  Upper_bound = c(upper_bound_Unif, upper_bound_Jeffrey,
    ↪upper_bound_Step)
)
Credibility_Interval

#Plot the limits
ggplot()+
geom_point(aes(mu, p.norm_Unif, color='Uniform Prior')) +
geom_vline(aes(xintercept=lower_bound_Unif,color='Uniform Prior'),
  ↪linetype='dashed')+
geom_vline(aes(xintercept=upper_bound_Unif, color='Uniform Prior'),
  ↪linetype='dashed')+

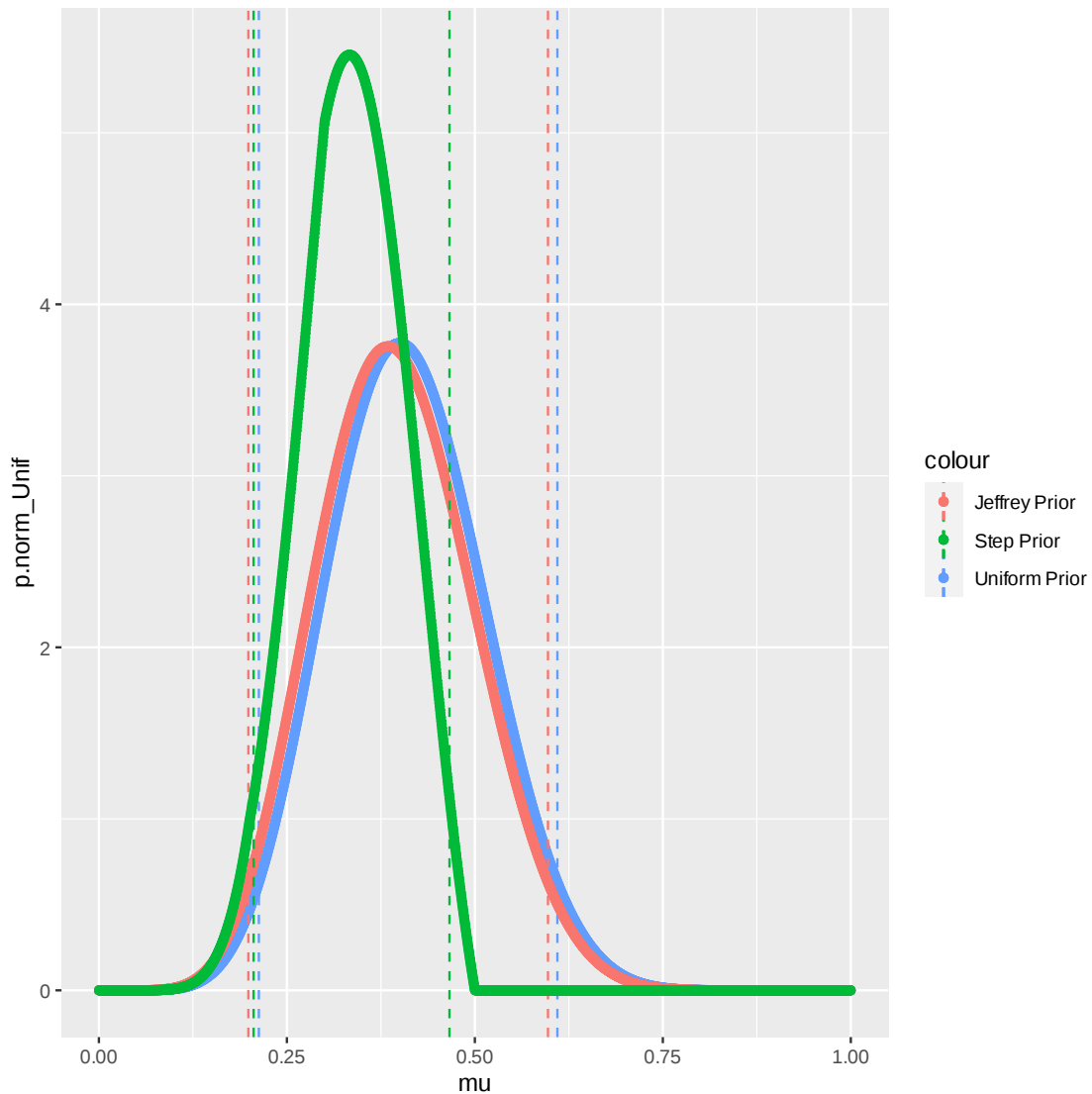
geom_point(aes(mu, p.norm_Jeffrey, color='Jeffrey Prior')) +
geom_vline(aes(xintercept=lower_bound_Jeffrey,color='Jeffrey Prior'),
  ↪linetype='dashed')+
geom_vline(aes(xintercept=upper_bound_Jeffrey, color='Jeffrey Prior'),
  ↪linetype='dashed') +

geom_point(aes(mu, p.norm_Step, color='Step Prior')) +
geom_vline(aes(xintercept=lower_bound_Step,color='Step Prior'),
  ↪linetype='dashed')+
geom_vline(aes(xintercept=upper_bound_Step, color='Step Prior'),
  ↪linetype='dashed')

```

A tibble: 3 × 3

	Prior <chr>	Lower_bound <dbl>	Upper_bound <dbl>
	Uniform	0.21275	0.60975
	Jeffrey	0.19875	0.59725
	Step	0.20575	0.46625



4 Ex.2

4.1 Plot the posterior distribution and compute first and second moments

```
[178]: n <- 116
y <- 17
n.sample <- 4000; delta.mu <- 1/n.sample
mu <- seq(from=1/(2*n.sample), by = delta.mu, length.out = n.sample)

# Uniform prior
p.star_Unif <- dbinom(x=y, size=n, prob=mu) * dunif(mu, min = 0, max=1)
p.norm_Unif <- p.star_Unif/(delta.mu * sum(p.star_Unif))
```

```

first_moment_Unif <- delta.mu * sum(mu * p.norm_Unif)
second_moment_Unif <- delta.mu * sum(((first_moment_Unif-mu)**2) * p.norm_Unif)

#Beta prior
p.star_Beta <- dbinom(x=y, size=n, prob=mu) * dbeta(mu,1,4)
p.norm_Beta <- p.star_Beta / (delta.mu * sum(p.star_Beta))
first_moment_Beta <- delta.mu * sum(mu * p.norm_Beta)
second_moment_Beta <- delta.mu * sum(((first_moment_Beta - mu)**2) * p.
  ↪norm_Beta)

Momenta <- tibble(
  Prior = c('Unif', 'Beta'),
  First_moment = c(first_moment_Unif, first_moment_Beta),
  Second_moment = c(second_moment_Unif, second_moment_Beta)
)

Momenta

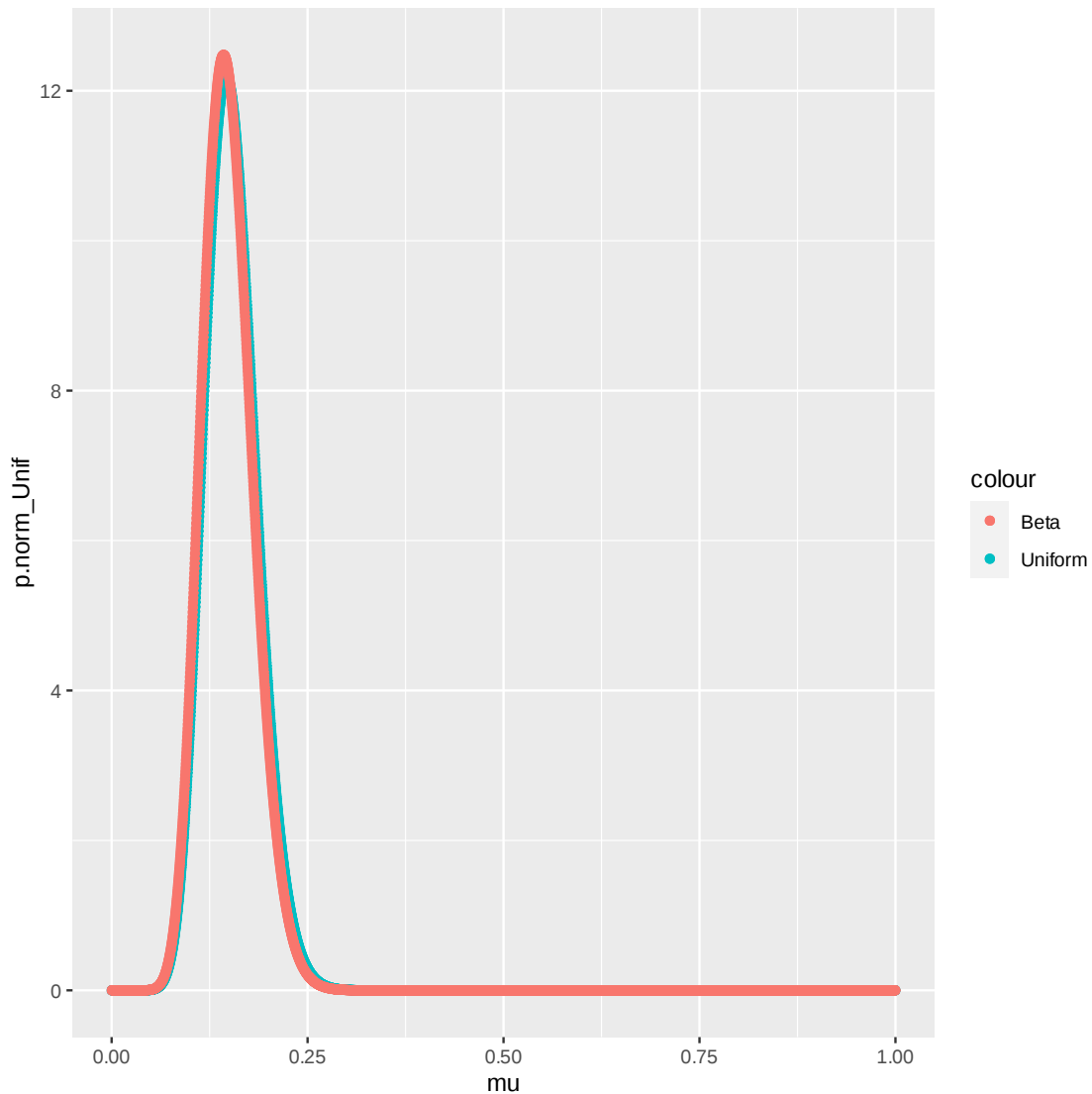
ggplot() +
geom_point(aes(mu, p.norm_Unif, color='Uniform')) +
geom_point(aes(mu, p.norm_Beta, color='Beta'))

```

```

A tibble: 2 × 3
  Prior    First_moment Second_moment
  <chr>    <dbl>         <dbl>
1 Unif    0.1525424    0.001086329
2 Beta    0.1487603    0.001037957

```



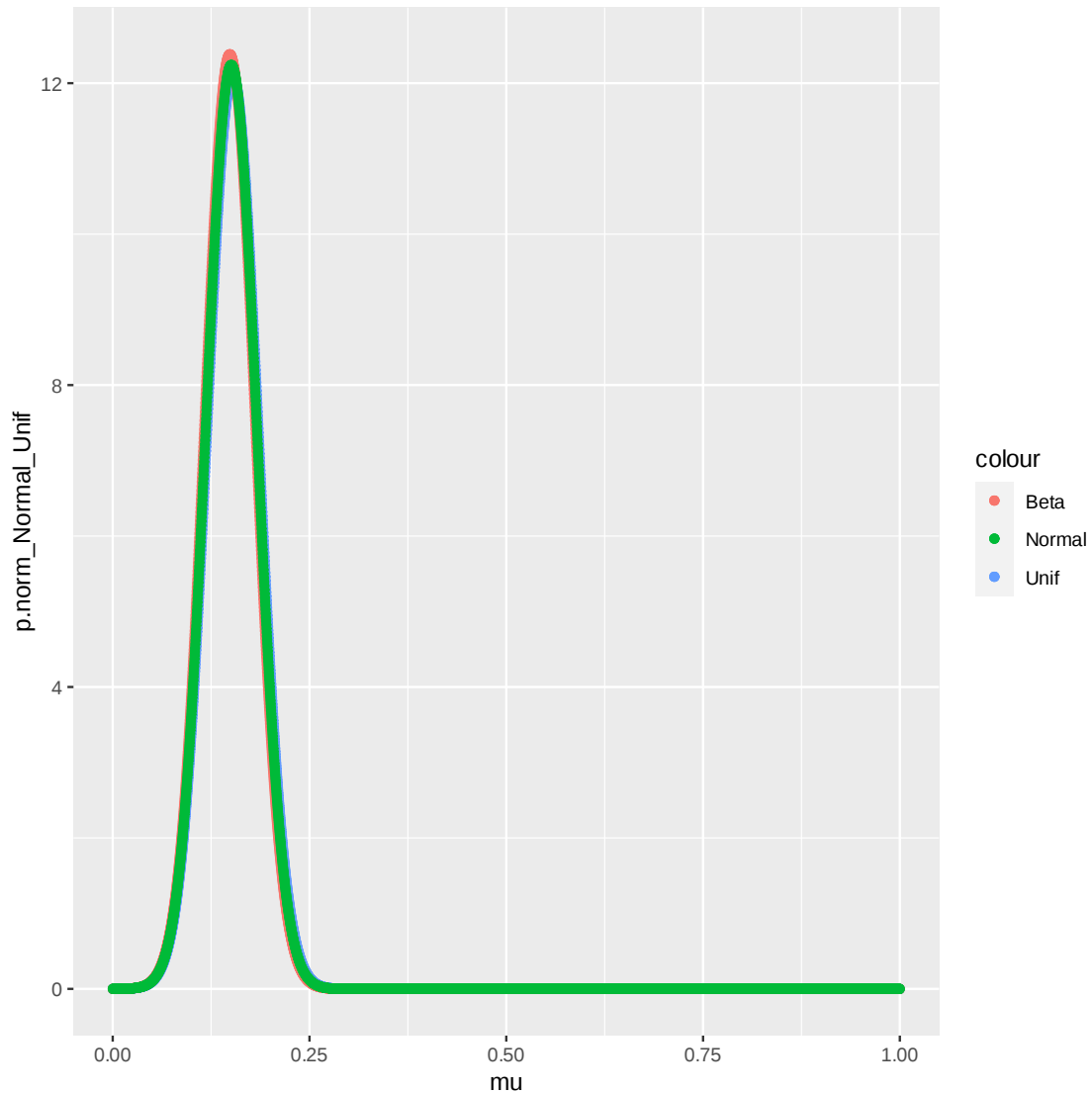
4.2 Find a normal approximation for the posterior

For the normal posterior distribution I'm gonna use a normal distribution with mean equal to the first moments of the posterior distribution found in the last ex, same for the std

```
[180]: p.norm_Normal_Unif <- dnorm(mu, mean=first_moment_Unif,
  ↪sd=sqrt(second_moment_Unif))
p.norm_Normal_Beta <- dnorm(mu, mean=first_moment_Beta,
  ↪sd=sqrt(second_moment_Beta))

p.norm_Normal <- dnorm(mu, mean=(first_moment_Unif + first_moment_Beta)/2,
  sd=sqrt((second_moment_Unif + second_moment_Beta)/2))
```

```
ggplot() +
  geom_point(aes(mu, p.norm_Normal_Unif, color='Unif'))+
  geom_point(aes(mu, p.norm_Normal_Beta, color='Beta')) +
  geom_point(aes(mu, p.norm_Normal, color='Normal'))
```



4.3 Compute a 95% credibility interval both for the original posterior and for the normal approximation, giving the results in a summary table

```
[186]: lower_bound_Unif <- ncredint(mu, p.norm_Unif, level = 0.95)[['lower']]
       upper_bound_Unif <- ncredint(mu, p.norm_Unif, level = 0.95)[['upper']]

       lower_bound_Beta <- ncredint(mu, p.norm_Beta, level = 0.95)[['lower']]
```



```

upper_bound_Beta <- ncredint(mu, p.norm_Beta, level = 0.95)[['upper']]

lower_bound_Normal <- ncredint(mu, p.norm_Normal, level = 0.95)[['lower']]
upper_bound_Normal <- ncredint(mu, p.norm_Normal, level = 0.95)[['upper']]

Credibility_Interval <- tibble(
  Prior = c('Unif', 'Beta', 'Normal approximation'),
  lower_bound = c(lower_bound_Unif, lower_bound_Beta,
    ↪ lower_bound_Normal),
  upper_bound = c(upper_bound_Unif, upper_bound_Beta,
    ↪ upper_bound_Normal)
)
Credibility_Interval

```

	Prior <chr>	lower_bound <dbl>	upper_bound <dbl>
A tibble: 3 × 3	Unif	0.090375	0.217875
	Beta	0.088125	0.212625
	Normal approximation	0.086875	0.214375

4.4 Add the limits on the plot of the posterior distributions

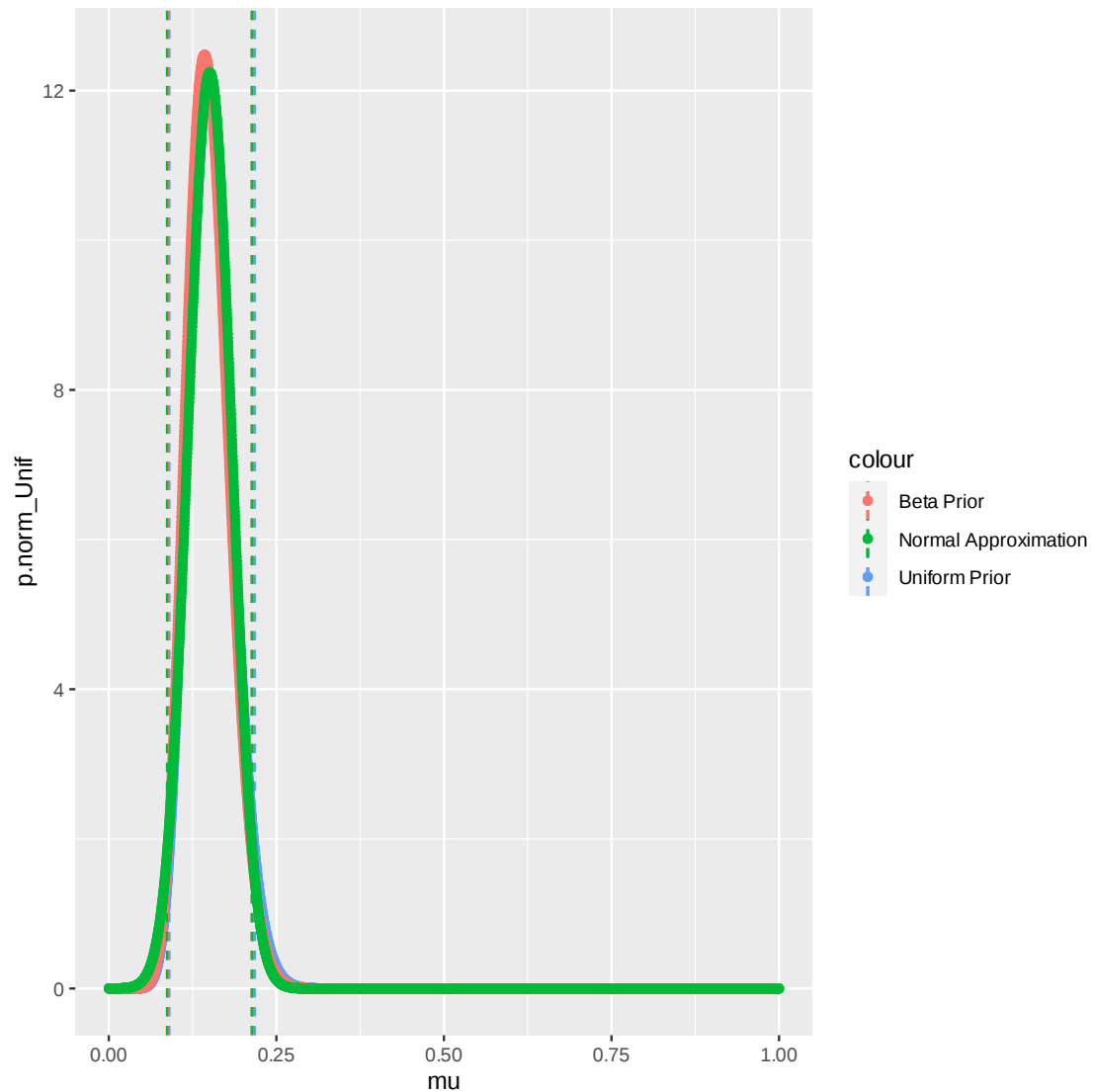
```

[188]: ggplot() +
  geom_point(aes(mu, p.norm_Unif, color='Uniform Prior')) +
  geom_vline(aes(xintercept=lower_bound_Unif,color='Uniform Prior'),
    ↪ linetype='dashed')+
  geom_vline(aes(xintercept=upper_bound_Unif, color='Uniform Prior'),
    ↪ linetype='dashed')+

  geom_point(aes(mu, p.norm_Beta, color='Beta Prior')) +
  geom_vline(aes(xintercept=lower_bound_Beta,color='Beta Prior'),
    ↪ linetype='dashed')+
  geom_vline(aes(xintercept=upper_bound_Beta, color='Beta Prior'),
    ↪ linetype='dashed') +

  geom_point(aes(mu, p.norm_Normal, color='Normal Approximation')) +
  geom_vline(aes(xintercept=lower_bound_Normal,color='Normal Approximation'),
    ↪ linetype='dashed')+
  geom_vline(aes(xintercept=upper_bound_Normal, color='Normal Approximation'),
    ↪ linetype='dashed')

```



5 Ex. 3

5.1 Assuming a flat prior, and a beta prior, plot the likelihood, prior and posterior distributions for the data set

```
[214]: n <- 30
y <- 15

n.sample <- 4001; delta.mu = 1/n.sample
mu <- seq(from=1/(2*n.sample), by = delta.mu, length.out = n.sample)

# Uniform prior
```

```

p.star_Unif <- dbinom(x=y, size=n, prob=mu) * dunif(mu, min = 0, max=1)
p.norm_Unif <- p.star_Unif/(delta.mu * sum(p.star_Unif))
first_moment_Unif <- delta.mu * sum(mu * p.norm_Unif)
second_moment_Unif <- delta.mu * sum(((first_moment_Unif-mu)**2) * p.norm_Unif)

#Beta prior
p.star_Beta <- dbinom(x=y, size=n, prob=mu) * dbeta(mu,2, 2)
p.norm_Beta <- p.star_Beta / (delta.mu * sum(p.star_Beta))
first_moment_Beta <- delta.mu * sum(mu * p.norm_Beta)
second_moment_Beta <- delta.mu * sum(((first_moment_Beta - mu)**2) * p.
  ↪norm_Beta)

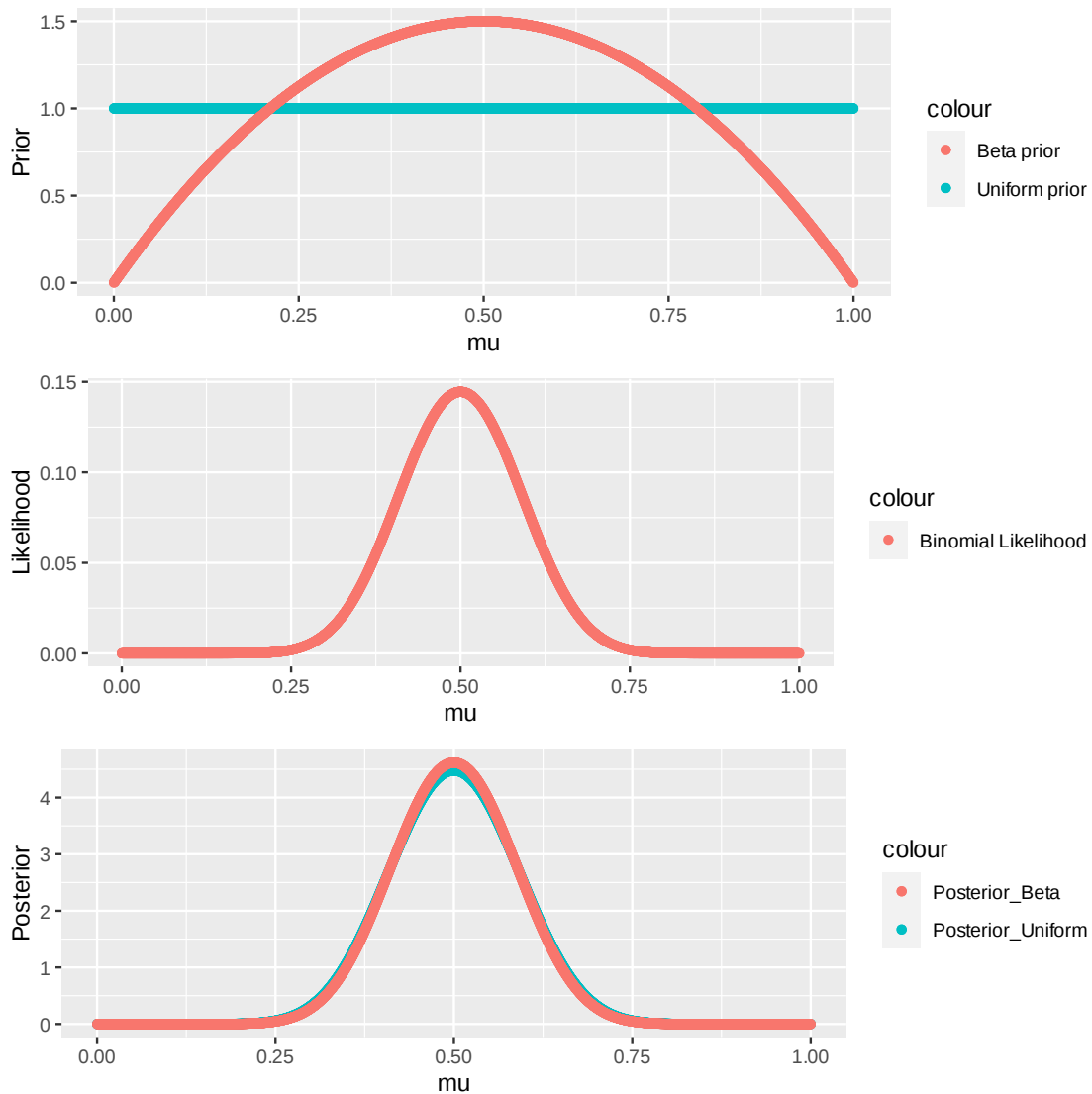
#Prior plot
ggplot() +
geom_point(aes(mu, dunif(mu, min=0, max=1), color='Uniform prior')) +
geom_point(aes(mu, dbeta(mu, 2,2), colour='Beta prior')) +
labs(y='Prior')-> prior_plot

#Likelihood plot
ggplot() +
geom_point(aes(mu, dbinom(x=y, size=n, prob=mu), color='Binomial Likelihood')) +
labs(y='Likelihood')-> Likelihood_plot

#Posterior plot
ggplot() +
geom_point(aes(mu, p.norm_Unif, color='Posterior_Uniform')) +
geom_point(aes(mu, p.norm_Beta, color='Posterior_Beta'))+
labs(y='Posterior')-> posterior_plot

grid.arrange(prior_plot, Likelihood_plot, posterior_plot, nrow=3)

```



5.2 Evaluate the most probable value for the coin probability p and, integrating the posterior probability distribution, give an estimate for a 95% credibility interval

```
[215]: cat('The most probable value is:', mu[which.max(p.norm_Beta)])
```

The most probable value is: 0.5

```
[216]: lower_bound <- ncredint(mu, p.norm_Beta, level=0.95)[['lower']]
upper_bound <- ncredint(mu, p.norm_Beta, level=0.95)[['upper']]
cat('The 95% credibility interval is:', '[', lower_bound, ',', upper_bound, ']')
```

The 95% credibility interval is: [0.3355411 , 0.6644589]

5.3 Repeat the same analysis assuming a sequential analysis of the data 1. Show how the most probable value and the credibility interval change as a function of the number of coin tosses (i.e. from 1 to 30)

Head H is 1, Tail is 0

```
[221]: trial = c(0,0,0,0,0,1,0,0,1,1,0,0,1,1,1,0,1,0,1,0,1,1,0,1,0,1,0,1,1,1)

[241]: t <- c()

most_prob_Unif <- c()
most_prob_Beta <- c()

t_lower_bound_Unif <- c()
t_upper_bound_Unif <- c()
t_lower_bound_Beta <- c()
t_upper_bound_Beta <- c()

for(i in trial){
  t <- c(t,i)
  n <- length(t)
  y <- sum(t)

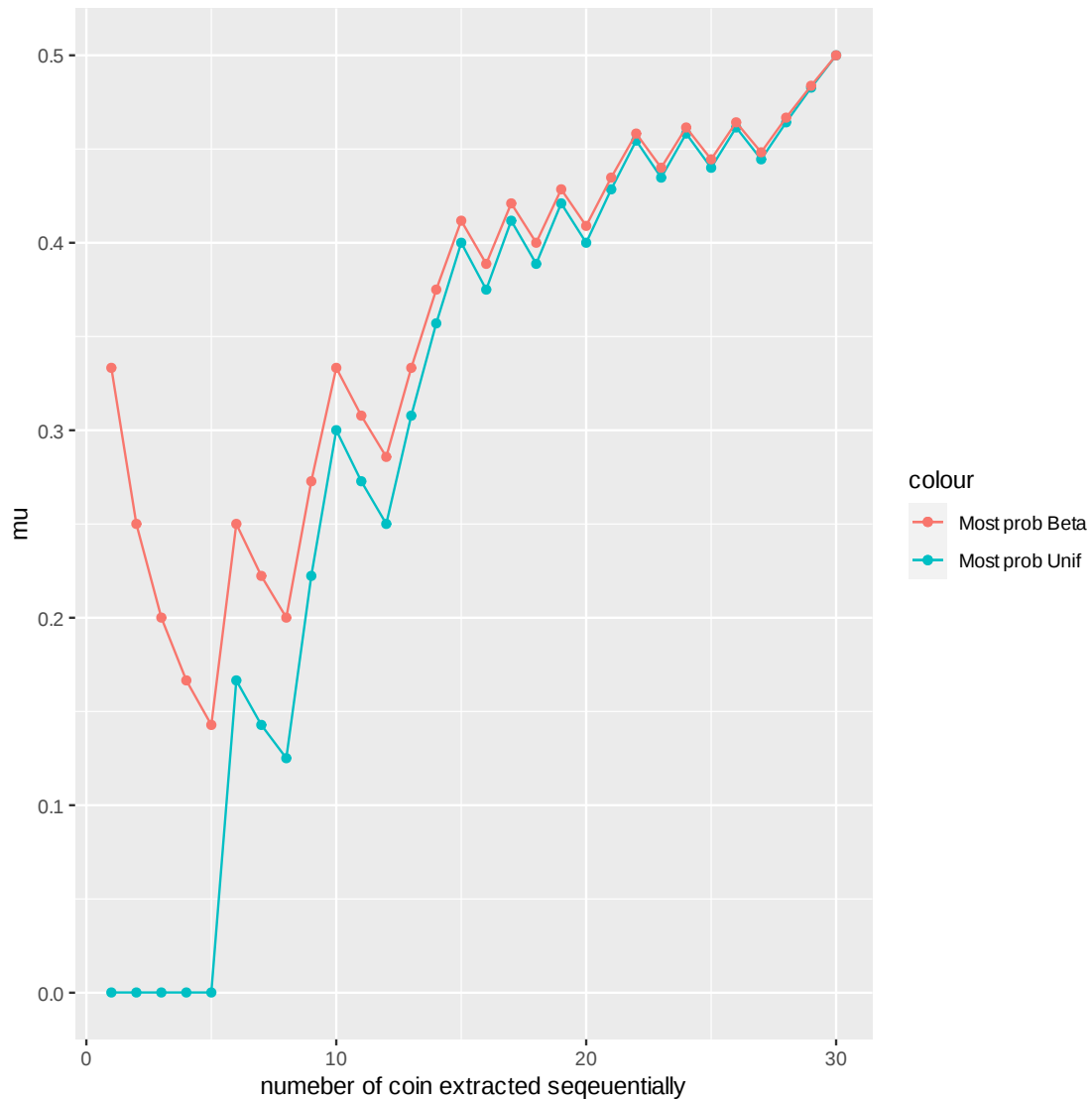
  n.sample <- 4001; delta.mu = 1/n.sample
  mu <- seq(from=1/(2*n.sample), by = delta.mu, length.out = n.sample)

  # Uniform prior
  p.star_Unif <- dbinom(x=y, size=n, prob=mu) * dunif(mu, min = 0, max=1)
  p.norm_Unif <- p.star_Unif/(delta.mu * sum(p.star_Unif))
  most_prob_Unif <- c(most_prob_Unif, mu[which.max(p.norm_Unif)])
  t_lower_bound_Unif <- c(t_lower_bound_Unif, ncredint(mu, p.norm_Unif,
↪level=0.95)[['lower']])
  t_upper_bound_Unif <- c(t_upper_bound_Unif, ncredint(mu, p.norm_Unif,
↪level=0.95)[['upper']])

  #Beta prior
  p.star_Beta <- dbinom(x=y, size=n, prob=mu) * dbeta(mu,2, 2)
  p.norm_Beta <- p.star_Beta / (delta.mu * sum(p.star_Beta))
  most_prob_Beta <- c(most_prob_Beta, mu[which.max(p.norm_Beta)])
  t_lower_bound_Beta <- c(t_lower_bound_Beta, ncredint(mu, p.norm_Beta,
↪level=0.95)[['lower']])
  t_upper_bound_Beta <- c(t_upper_bound_Beta, ncredint(mu, p.norm_Beta,
↪level=0.95)[['upper']])
}
```

```
[258]: ggplot() +
geom_point(aes(seq(1,30), most_prob_Unif, color='Most prob Unif')) +
geom_line(aes(seq(1,30), most_prob_Unif, color='Most prob Unif')) +
```

```
geom_point(aes(seq(1,30), most_prob_Beta, color='Most prob Beta')) +
geom_line(aes(seq(1,30), most_prob_Beta, color='Most prob Beta')) +
labs(x = 'numeber of coin extracted sequentially', y='mu')
```



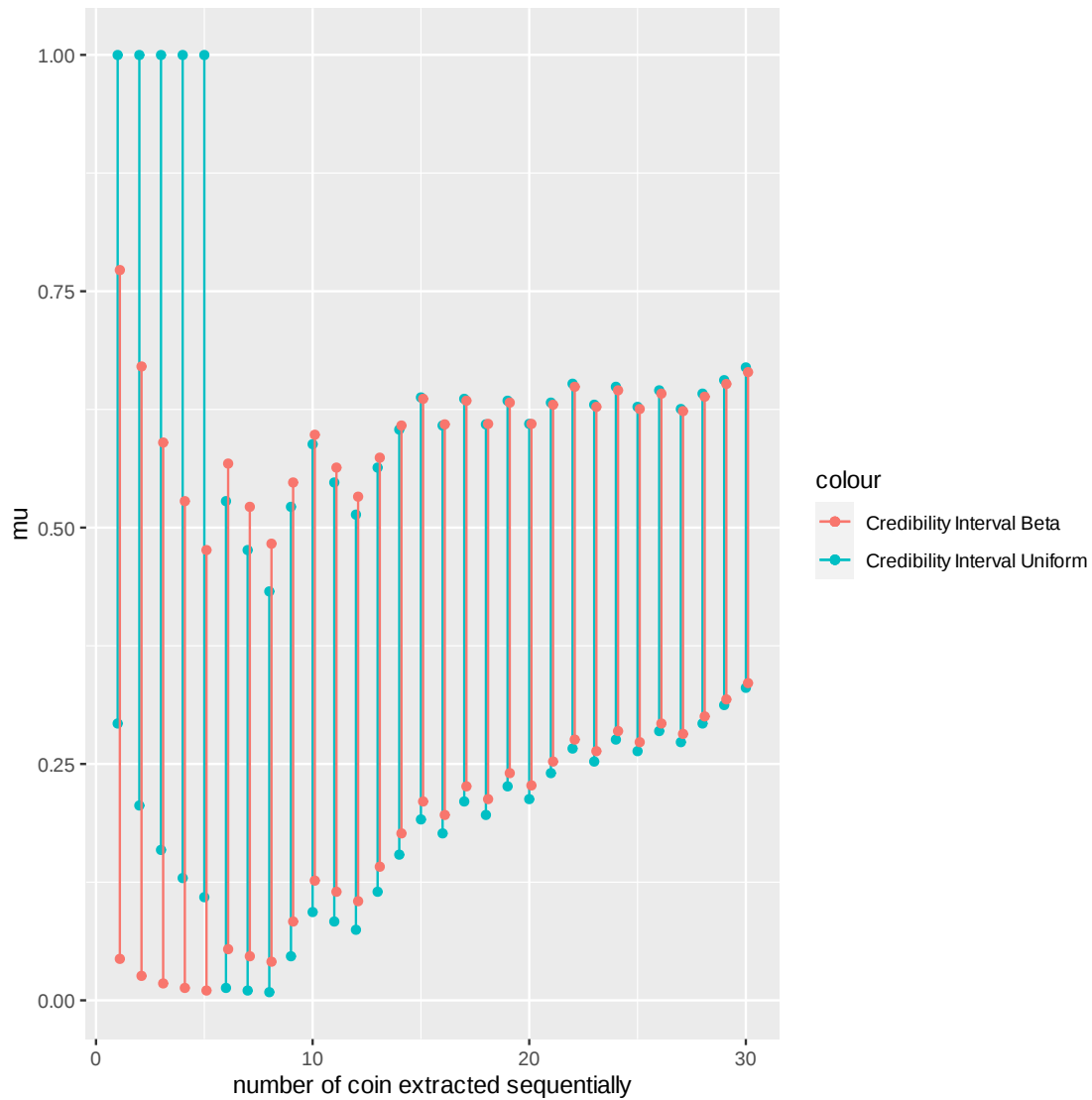
[252]: t_upper_bound_Unif

1. 0.99987503124219	2. 0.99987503124219	3. 0.99987503124219	4. 0.99987503124219
5. 0.99987503124219	6. 0.527993001749563	7. 0.476255936015996	8. 0.432516870782304
9. 0.521994501374656	10. 0.588227943014247	11. 0.547738065483629	12. 0.51374656335916
13. 0.563484128967758	14. 0.603724068982754	15. 0.637715571107223	16. 0.607973006748313
17. 0.636215946013497	18. 0.609222694326418	19. 0.634216445888528	20. 0.609722569357661
21. 0.632216945763559	22. 0.652211947013247	23. 0.629967508122969	24. 0.648962759310173
25. 0.627718070482379	26. 0.645213696575856	27. 0.62546863284179	28. 0.641714571357161

29. 0.655961009747563 30. 0.669457635591102

[]: t

```
[259]: ggplot() +  
  geom_segment(aes(x = seq(1,30), y= t_lower_bound_Unif, xend=seq(1,30),  
    ↪yend=t_upper_bound_Unif, colour='Credibility Interval Uniform')) +  
  geom_point(aes(x=seq(1,30), y = t_lower_bound_Unif, colour='Credibility_  
    ↪Interval Uniform'))+  
  geom_point(aes(x=seq(1,30), y = t_upper_bound_Unif, colour='Credibility_  
    ↪Interval Uniform')) +  
  geom_segment(aes(x = seq(from=1.1, to=30.1, by=1), y= t_lower_bound_Beta,  
    ↪xend=seq(from=1.1, to=30.1, by=1), yend=t_upper_bound_Beta,  
    ↪colour='Credibility Interval Beta')) +  
  geom_point(aes(x = seq(from=1.1, to=30.1), y = t_lower_bound_Beta,  
    ↪colour='Credibility Interval Beta'))+  
  geom_point(aes(x = seq(from=1.1, to=30.1), y = t_upper_bound_Beta,  
    ↪colour='Credibility Interval Beta'))+  
  labs(y='mu', x='number of coin extracted sequentially')
```



5.4 Do you get a different result, by analyzing the data sequentially with respect to a one-step analysis (i.e. considering all the data as a whole)?

No, in the end the most probable value and the credibility interval are evaluated in the same way when the last coin result is considered

6 Ex.4

```
[109]: #1 is white
        #0 is black
```



```

#box k
k = sample(seq(0,5,1), 1)
cat('the Box is the :', k, '\n')

#n is number of extraction
n=100
extraction <- rbinom(n, size=1, prob=k/5)
cat('Sequence of balls:', extraction)

#function to obtain the  $P(E_w/H_j)$  and  $p(E_b/H_j)$ 
p.star <- function(i,j){
  if(i==1){
    return(j/5)
  }else{
    return((5-j)/5)
  }
}

#Posterior before extracting the first ball is the flat distribution
P_0 <- c(1/5)
P_1 <- c(1/5)
P_2 <- c(1/5)
P_3 <- c(1/5)
P_4 <- c(1/5)
P_5 <- c(1/5)

P_posterior <- c(rep(1/5, 6))

#Function that for each extraction update the value of the Posterior_
↪distribution
for(i in extraction){
  P_posterior_star <- c()
  for(j in seq(1,6,1)){
    P_posterior_star <- c(P_posterior_star, p.star(i,(j-1))*P_posterior[j])
  }
  P_posterior <- P_posterior_star/(sum(P_posterior_star))

  P_0 <- c(P_0, P_posterior[1])
  P_1 <- c(P_1, P_posterior[2])
  P_2 <- c(P_2, P_posterior[3])
  P_3 <- c(P_3, P_posterior[4])
  P_4 <- c(P_4, P_posterior[5])
  P_5 <- c(P_5, P_posterior[6])
}

#Plot

```

```

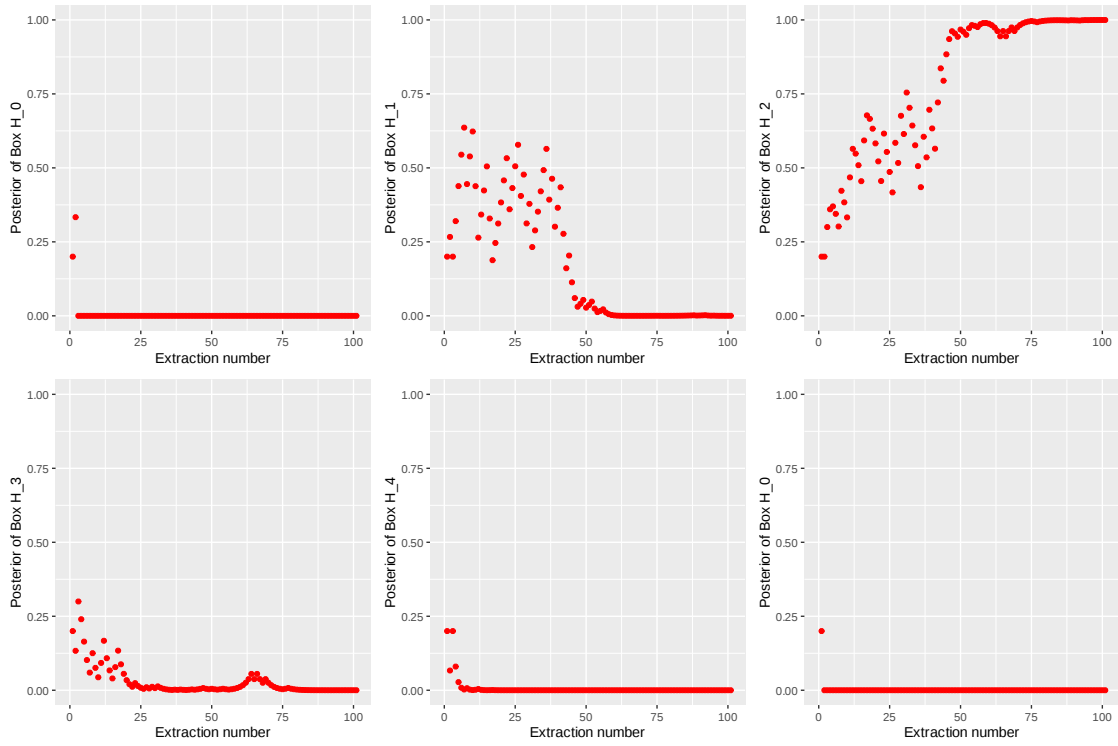
plot_P_0 <- ggplot() + geom_point(aes(seq(1, n+1), P_0), color='red') +
  ↪labs(x='Extraction number', y='Posterior of Box H_0') + xlim(0, n+1) +
  ↪ylim(0,1)
plot_P_1 <- ggplot() + geom_point(aes(seq(1, n+1), P_1), color='red') +
  ↪labs(x='Extraction number', y='Posterior of Box H_1') + xlim(0, n+1) +
  ↪ylim(0,1)
plot_P_2 <- ggplot() + geom_point(aes(seq(1, n+1), P_2), color='red') +
  ↪labs(x='Extraction number', y='Posterior of Box H_2') + xlim(0, n+1) +
  ↪ylim(0,1)
plot_P_3 <- ggplot() + geom_point(aes(seq(1, n+1), P_3), color='red') +
  ↪labs(x='Extraction number', y='Posterior of Box H_3') + xlim(0, n+1) +
  ↪ylim(0,1)
plot_P_4 <- ggplot() + geom_point(aes(seq(1, n+1), P_4), color='red') +
  ↪labs(x='Extraction number', y='Posterior of Box H_4') + xlim(0, n+1) +
  ↪ylim(0,1)
plot_P_5 <- ggplot() + geom_point(aes(seq(1, n+1), P_5), color='red') +
  ↪labs(x='Extraction number', y='Posterior of Box H_0') + xlim(0, n+1) +
  ↪ylim(0,1)

options(repr.plot.width=12, repr.plot.height=8)
grid.arrange(plot_P_0, plot_P_1, plot_P_2, plot_P_3, plot_P_4, plot_P_5,
  ↪nrow=2, ncol=3)

```

the Box is the : 2

Sequence of balls: 0 1 0 0 0 0 1 0 0 1 1 0 0 0 1 1 0 0 0 0 0 1 0 0 0 1 0 1 0 1 0
 0 0 0 0 1 0 1 0 0 1 1 0 1 1 1 0 0 1 0 0 1 1 0 0 1 1 1 1 1 1 0 1 0 0 1 0 0 0
 0 0 0 1 1 0 0 0 0 0 0 0 0 0 0 0 1 0 0 0 1 1 0 1 1 0 1 0 0



[]: