

Homework 2

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Exercise 5

We want to generate pseudo-random numbers from a $\text{Beta}(3, 1)$ -distribution which has the density $f(x) = 3x^2$ using only `runif` implemented for random number generation in R.

Write a function implementing the inversion method.

```
library(GoFKernel)
```

```
## Loading required package: KernSmooth
```

```
## KernSmooth 2.23 loaded
```

```
## Copyright M. P. Wand 1997-2009
```

```
library(dplyr)
```

```
## Warning: package 'dplyr' was built under R version 4.2.3
```

```
##
```

```
## Attaching package: 'dplyr'
```

```
## The following objects are masked from 'package:stats':
```

```
##
```

```
##     filter, lag
```

```
## The following objects are masked from 'package:base':
```

```
##
```

```
##     intersect, setdiff, setequal, union
```

```
# Using the fact that a the cdf of the Beta(3, 1)
```

```
# distribution can be approximated by the binomial distribution with parameters n = 3, k = 0, and p =
```

```
# The cumulative distribution function:
```

```
cdf_binomial_alpha_3_beta_1 <- function(x) {  
  ifelse(x < 0, 0, ifelse(x > 1, 1, pbinom(0, 3, 1 - x)))  
}
```

```
cdf_inverse_cdf_binomial_alpha_3_beta_1 <- inverse(cdf_binomial_alpha_3_beta_1, lower = 0, upper = 1)
```

```
# Checking that it works:
```

```
cdf_inverse_cdf_binomial_alpha_3_beta_1(0.2)
```

```
## [1] 0.584804
```

```
cdf_binomial_alpha_3_beta_1(cdf_inverse_cdf_binomial_alpha_3_beta_1(0.2))
```

```
## [1] 0.2000005
```

```

# Generating random numbers using the inverse method:

gen_inv_beta_3_1 <- function(n) {
  sample <- runif(n)
  beta_3_1_sample <- numeric(n)
  for (i in 1:n) {
    beta_3_1_sample[i] <- cdf_inverse_cdf_binomial_alpha_3_beta_1(sample[i])
  }
  return(beta_3_1_sample)
}

# Measure execution time and save in time_1
time_1 <- system.time({
  gen_inv_beta_3_1(10000)
})

```

Write a function which implements a transformation method.

```

library(lmomco)
# From the class slides, we know that the random variable  $X = U / (U + V)$ 
# where  $U \sim \text{Gamma}(r, \text{lambda})$  and  $V \sim \text{Gamma}(s, \text{lambda})$  is distributed as  $\text{Beta}(r, s)$ 
# so that
n = 1000
sample <- runif(n)
lambda <- 1

cdf_gamma_3_lambda <- function(x) {
  pgamma(x,3,lambda)
}

cdf_inverse_gamma_3 <- function(p, shape = 3, rate = 1, lower = 0, upper = 100000) {
  uniroot(function(x) pgamma(x, shape = shape, rate = rate) - p, lower = lower, upper = upper)$root
}

cdf_inverse_gamma_1 <- function(p, shape = 1, rate = 1, lower = 0, upper = 100000) {
  uniroot(function(x) pgamma(x, shape = shape, rate = rate) - p, lower = lower, upper = upper)$root
}

# Checking that it works: (There is a small error in the last digit)
cdf_inverse_gamma_3(cdf_gamma_3_lambda(20))

## [1] 20.00001

# Function to generate samples from the inverse CDF
gen_transf_beta_3_1 <- function(n) {
  sample <- runif(n)
  sample_2 <- runif(n)
  U <- numeric(n)
  V <- numeric(n)
  B <- numeric(n)

  for (i in 1:n) {
    U[i] <- cdf_inverse_gamma_3(sample[i])
    V[i] <- cdf_inverse_gamma_1(sample_2[i])
  }
}

```

```

    B[i] <- U[i] / (U[i] + V[i])
  }

  return(B)
}

# Measure execution time and save in time_2
time_2 <- system.time({
  Beta_random <- gen_transf_beta_3_1(10000)
})

```

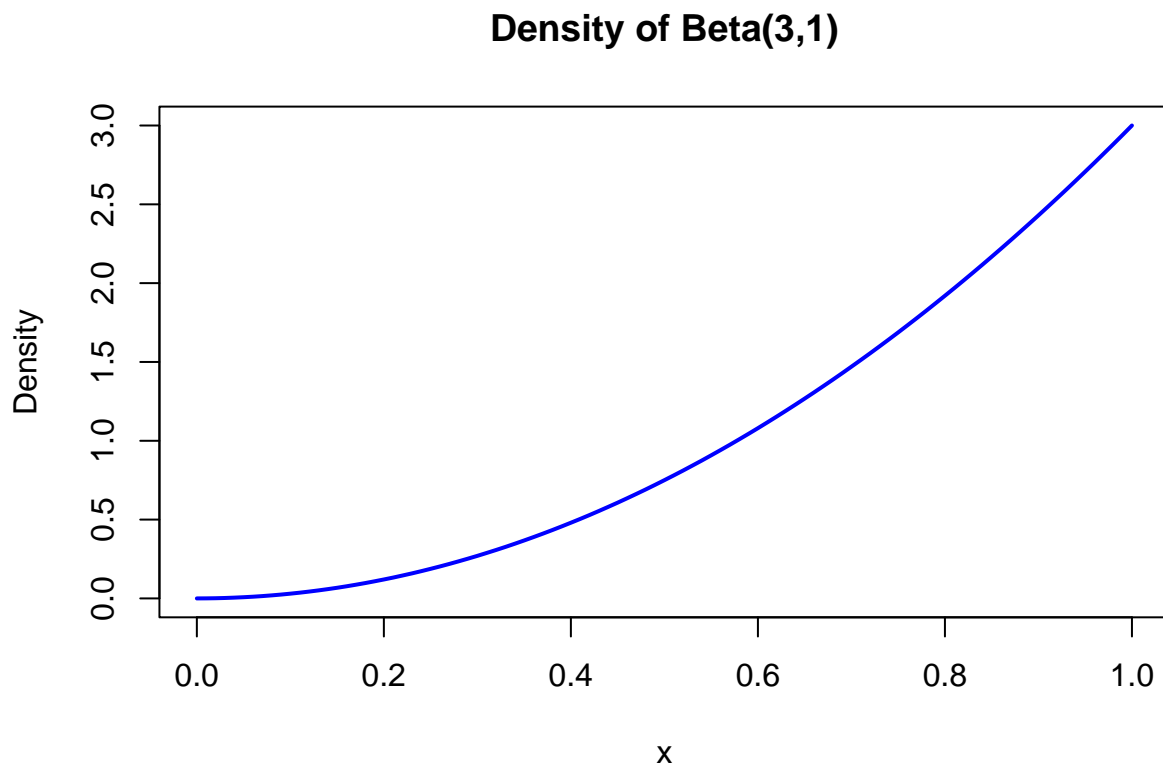
Write a function which implements rejection sampling using a suitable proposal density.

*# Following the example in the class, we know that the highest
value of the density function provided is at*

```

# Plot the density function of Beta(3,1)
curve(dbeta(x, 3, 1), from = 0, to = 1, lwd = 2, col = "blue",
      ylab = "Density", xlab = "x", main = "Density of Beta(3,1)")

```



*# The density function of the proposal distribution is such
that it is monotonically increasing and has a maximum at 1 with
a value of 3.*

*# We establish the minimal possible M equal to 3
And choose as distribution for Y the uniform distribution in 0,1*

```

M <- 3

rbeta31_rs <- function(n) {
  x <- numeric(n)
  i <- 1
  while (i <= n) {
    y <- runif(1)
    u <- runif(1)
    if (u <= y^2) {
      x[i] <- y
      i <- i + 1
    }
  }
  return(x)
}

time_3 <- system.time({
  rbeta31_rs(10000)
})

print(time_1)

##      user      system elapsed
##    0.28      0.00      0.72

print(time_2)

##      user      system elapsed
##    0.95      0.00      1.55

print(time_3)

##      user      system elapsed
##    0.03      0.01      0.11

# The 'user time' is the CPU time charged for the execution of user instructions of the calling process

```

- Compare the computational performance of these implementations.

The Rejection Sampling Method seems to be the fastest, followed by the Inversion Method and then the Transformation Method.