Exercise set 3

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Exercise 3.3

The Pareto(a, b) distribution has cdf

$$F(x) = 1 - \frac{b^a}{x^a} \tag{1}$$

Derive the probability inverse transformation $F^{-1}(U)$ and use the inverse transform method to to simulate a random sample from the Pareto(2, 2) dist.

The function F^{-1} is derived by setting F(X) = U and solving for X which gives us.

$$U = 1 - \frac{4}{x^2} \implies X = \frac{2}{\sqrt{(1-u)}}$$
 (2)

and this we have $F^{-1}(U) = \frac{2}{\sqrt{(1-u)}}$

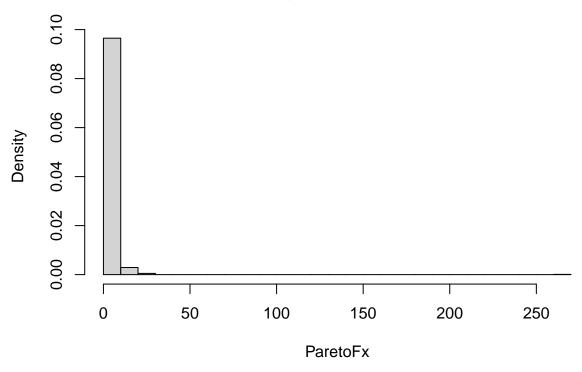
and in general terms $F^{-1}(U) = \frac{2}{\sqrt{(1-u)}}$

Calculate using R

```
simPareto <- function(nSim, a, b) {
  u <- runif(nSim)
  return(b*(1-u)^(-1/a))
}

nSim = 1000
ParetoFx <- simPareto(nSim = nSim, a = 2, b = 2)
hist(ParetoFx, breaks=30, prob=TRUE)</pre>
```

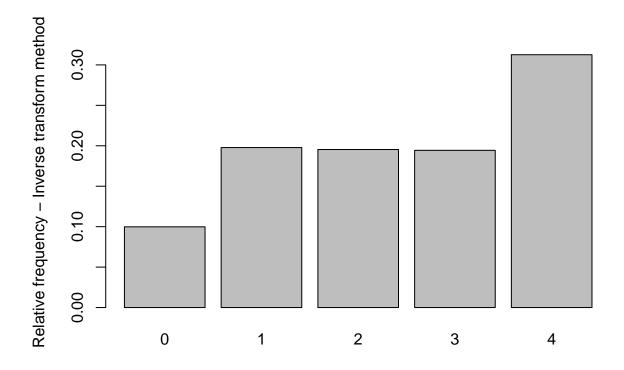




Exercise 3.5

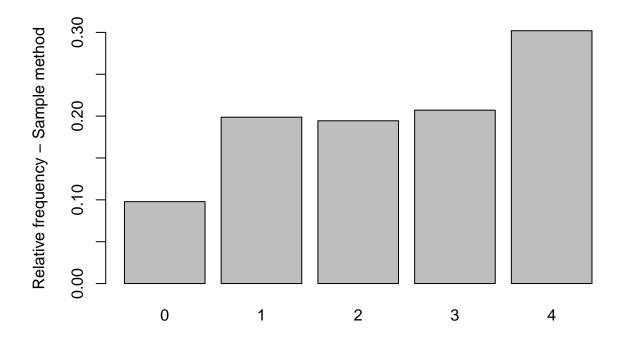
Use the inverse transform method to generate a random sample of size from distribution

```
# Alternatively the built in sample function can be used
## Generating data from the number of heads example where
## f(0)=1/8, f(1)=3/8, f(2)=3/8 and f(3)=1/8
\# Function for simulating number of heads according to the distribution above
discrv <- function(Nsim){</pre>
   U <- runif(Nsim)</pre>
   X <- rep(0,Nsim)</pre>
   X[(U>0.1) & (U<=0.3)] <-1
   X[(U>0.3) & (U<=0.5)] <- 2
   X[(U>0.5) & (U<=0.7)] <- 3
   X[U>0.7] <- 4
   return(X)
}
Nsim <- 10000
dfx <- discrv(Nsim)</pre>
relfreq <- table(dfx)/Nsim
barplot(relfreq,ylab="Relative frequency - Inverse transform method")
```



```
# Alternatively the built in sample function can be used

dfx2 <- sample(0:4,size=Nsim,replace=TRUE,prob=c(0.1,0.2,0.2,0.2,0.3))
relfreq <- table(dfx2)/Nsim
barplot(relfreq,ylab="Relative frequency - Sample method")</pre>
```

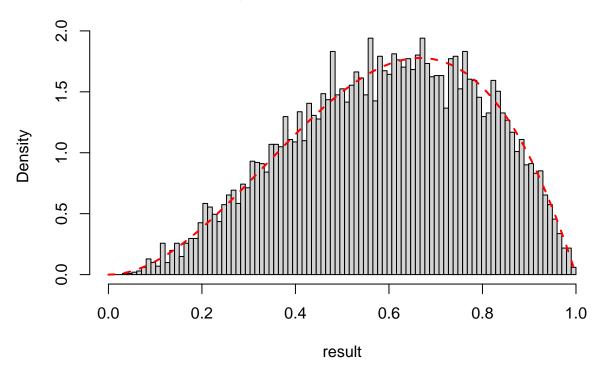


Exercise 3.7

Write a function to generate a random sample of size n from the beta (a,b) distribution by the acceptance-rejection method. Sample size should be 1000 from the beta (3,2) distribution. Graph the sample with the theoretical Beta (3,2) density.

```
## Function for acceptance rejection generation
rejectionBeta <- function(a, b, x, n) {
   f \leftarrow function(x) (x^(a-1)*(1-x)^(b-1) / beta(a, b))
   g <- function(x) 1
   C \leftarrow \max(f(x)/g(x))
   naccepts <- 0
   result.sample <- rep(NA, n)
   while (naccepts < n) {</pre>
    y <- runif(1)
    u <- runif(1)
    if (u \le f(y) / (C*g(y)))  {
      naccepts <- naccepts + 1</pre>
      result.sample[naccepts] = y
    }
   }
   result.sample
```

Histogram of data and true density



Exercise 3.17

```
a = 3, b = 2

Niter <- 1000
Nsim <- 5000

system.time(for (i in 1:Niter)
   rejectionBeta(3, 2, x, Nsim))</pre>
```

```
##
      user system elapsed
##
     38.22
              0.01
                     38.23
system.time(for (i in 1:Niter)
   rbeta(Nsim, shape1=3, shape2=2))
##
            system elapsed
##
      0.83
              0.00
                      0.83
a = 1, b = 1
Niter <- 1000
Nsim <- 5000
system.time(for (i in 1:Niter)
  rejectionBeta(1, 1, x, Nsim))
##
            system elapsed
      user
##
     21.00
              0.02
                     21.03
system.time(for (i in 1:Niter)
   rbeta(Nsim, shape1=1, shape2=1))
##
            system elapsed
      user
##
      0.84
              0.00
```

Exercise 1

a. Simulate the probability that component 1 and component 2 is working.

b. The system works if and only if at least one of components 1 or 2 works.

```
library("Rlab")
                                                     # Load Rlab package
## Rlab 2.15.1 attached.
##
## Attaching package: 'Rlab'
## The following objects are masked from 'package:stats':
##
##
       dexp, dgamma, dweibull, pexp, pgamma, pweibull, qexp, qgamma,
##
       qweibull, rexp, rgamma, rweibull
## The following object is masked from 'package:datasets':
##
##
       precip
n < -10
s <- 10000
mat \leftarrow matrix(rbern(n*s, c(0.7,0.95,0.95,0.95,0.99,0.99,0.92,0.92,0.92,0.7)), n, s)
p <- numeric(4)</pre>
p[1] \leftarrow (sum(mat[1,]) / s) * (sum(mat[2,]) / s)
```

p[2] <- sum(mat[1,] / s) + sum(mat[2,] / s) - p[1]

c. The system works if and only if both component 1 and component 2 and at least one of components 3 or 4 work.

```
p[3] <- p[1] * (sum(mat[3,] / s) + sum(mat[4,] / s) - (sum(mat[3,]) / s) * (sum(mat[4,]) / s))
d. The system works if and only if at least 7 of the 10 components work.
p[4] <- sum(colSums(mat) < 7) / s</pre>
```

Calculated probabilities

```
p
## [1] 0.6634800 0.9849200 0.6618118 0.0069000
```

Exercise 2

a. Find conditional and unconditional probability of find larger than 8

Simulate $P(X > 8 \mid \text{ field found})$ using inverse transform method

```
triangle <- function(n, a, b, c) {</pre>
  f <- function (x) {
    return((c-a)/(b-a))
  }
  u <- runif(n)
  sample <- ifelse(u < f(c),</pre>
          a + sqrt(u*(b-a)*(c-a)),
          b - sqrt((1-u)*(b-a)*(b-c))
  return(sample)
}
sims <- 10000
a <- 2
b <- 10
c <- 6
vals <- triangle(sims, 2, 10, 6)</pre>
p1 <- sum(vals > 8) / sims
p2 \leftarrow p1 * 0.4
```

b. By simulation and the risk-weighted and the non risk-weighted expectation for the total resource R

```
tenTriangles <- function(Nsim) {
    n1 <- sum(triangle(Nsim, 2, 6, 4) / sims)
    n2 <- sum(triangle(Nsim, 3, 11, 7) / sims)
    n3 <- sum(triangle(Nsim, 2, 6, 4) / sims)
    n4 <- sum(triangle(Nsim, 1, 9, 5) / sims)
    n5 <- sum(triangle(Nsim, 8, 10, 9) / sims)
    n6 <- sum(triangle(Nsim, 5, 9, 7) / sims)
    n7 <- sum(triangle(Nsim, 2, 6, 4) / sims)
    n8 <- sum(triangle(Nsim, 3, 5, 4) / sims)
    n9 <- sum(triangle(Nsim, 8, 12, 10) / sims)
    n10 <- sum(triangle(Nsim, 3, 7, 5) / sims)

mat <- matrix(0, 10, 2)</pre>
```

```
mat[1,] <- c(n1, n1*0.8)
mat[2,] <- c(n2, n2*0.3)
mat[3,] <- c(n3, n3*0.6)
mat[4,] <- c(n4, n4*0.6)
mat[5,] <- c(n5, n5*0.5)
mat[6,] <- c(n6, n6*0.9)
mat[7,] <- c(n7, n7*0.5)
mat[8,] <- c(n8, n8*0.8)
mat[9,] <- c(n9, n9*0.4)
mat[10,] <- c(n10, n10*0.4)

return(c(sum(mat[,1]), sum(mat[,2])))
}
tenTriangles(10000)</pre>
```

[1] 58.99372 32.70854

d. By simulation and an estimate of the accompanying standard deviation for the two expectations. Find how many replications are necessary to be 95% certain that the errors in the estimates of the means are not exceeding 0.2.

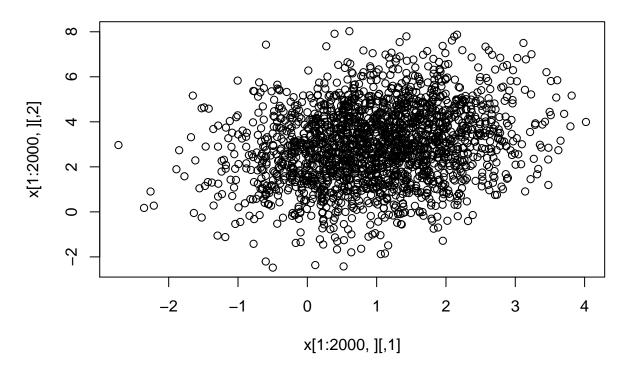
```
sample <- 100
nsim <- 10000
matrix <- matrix(nrow = 0, ncol = 2)
for (i in 1:sample) {
    matrix <- rbind(matrix, tenTriangles(nsim))
}
sdr1 <- sd(matrix[,1])
sdr2 <- sd(matrix[,2])
n1 <- (4*sdr1^2)/(0.2^2)
n2 <- (4*sdr2^2)/(0.2^2)
n1
## [1] 0.1088601
n2
## [1] 0.03579746</pre>
```

Exercise 3

a.

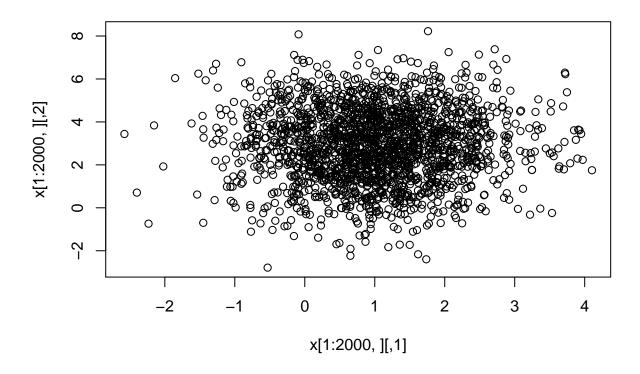
```
library(mvtnorm)

sims <- 10000
sigma <- matrix(c(1,0.4041,0.4041,3), ncol=2)
x <- rmvnorm(n=sims, mean=c(1, 3), sigma=sigma)
p <- sum((x[,1] + x[,2]) >= 3) / sims
plot(x[1:2000,])
```



```
p
## [1] 0.6751
    b.
library(mvtnorm)

sims <- 10000
sigma <- matrix(c(1,0,0,3), ncol=2)
x <- rmvnorm(n=sims, mean=c(1, 3), sigma=sigma)
p <- sum((x[,1] + x[,2]) >= 3) / sims
plot(x[1:2000,])
```



р

[1] 0.6789