

The first Homework of Statistical Computing

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1 Exercise 3.5

1.1 R code

Listing 1: R code of Exercise 3.5

```
1 Inv_trans <- function(n){  
2   rs = runif(n);  
3   for(i in 1:n){  
4     temp = rs[i]  
5     if (temp <= 0.1){  
6       rs[i] = 0;  
7     }else if(temp <= 0.3){  
8       rs[i] = 1;  
9     }else if(temp <= 0.5){  
10      rs[i] = 2;  
11     }else if(temp <= 0.7){
```

```

12     rs[i] = 3;
13 }else{
14     rs[i] = 4;
15 }
16 }
17 rs
18 }
19
20 RVs = Inv_trans(1000);
21 mySize = 400;
22 theoretical = c(.1, .2, .2, .2, .3)
23 T = matrix(rep(0,5*5),5,5)
24 for (i in 1:5){
25     mySample = sample(RVs,size=mySize, replace=F)
26     empirical = table(mySample)/mySize
27     T[i,] = empirical
28 }
29 T = rbind(theoretical,T)
30 rownames(T) = c('theoretical prob','empirical prob 1',
31                'empirical prob 2','empirical prob 3',
32                'empirical prob 4','empirical prob 5')
33 colnames(T) = c(0,1,2,3,4)
34 print(T)
35 write.csv(T,file='3.5_result.csv')

```

1.2 experiment result

I have repeated the experiment for 5 times. Compared with the theoretical probability, the five relative frequency tables are as follows:

表 1: experiment results of Exercise 3.5					
	0	1	2	3	4
theoretical prob	0.1000	0.2000	0.2000	0.2000	0.3000
empirical prob 1	0.1100	0.2275	0.1875	0.2025	0.2725
empirical prob 2	0.1200	0.2125	0.1900	0.1825	0.2950
empirical prob 3	0.1375	0.1925	0.1650	0.2125	0.2925
empirical prob 4	0.1200	0.1950	0.1825	0.2025	0.3000
empirical prob 5	0.1075	0.2275	0.2250	0.1725	0.2675

2 Exercise 3.6

Suppose Y is a random variable generated from density function $g(y)$, and U from Uniform(0, 1). According to the procedure of the Acceptance-Rejection method, we have

$$P(\text{accepted}|Y = y) = P(U < \frac{f(y)}{cg(y)}|Y = y) = \frac{f(y)}{cg(y)}$$

$$P(Y \text{ is accepted for one iteration})$$

$$= \int_{-\infty}^{+\infty} P(\text{accepted}|Y = y)g(y) dy$$

$$\begin{aligned}
&= \int_{-\infty}^{+\infty} \frac{f(y)}{cg(y)} g(y) dy \\
&= \frac{1}{c} \int_{-\infty}^{+\infty} f(y) \\
&= \frac{1}{c}
\end{aligned}$$

And finally we have

$$\begin{aligned}
f_{Y|accepted}(y) &= \frac{f(Y = y \text{ and accepted})}{P(Y \text{ is accepted for one iteration})} \\
&= \frac{P(\text{accepted} | Y = y)g(y)}{P(Y \text{ is accepted for one iteration})} \\
&= \frac{\frac{f(y)}{cg(y)}g(y)}{\frac{1}{c}} \\
&= f(y)
\end{aligned}$$

3 Exercise 3.9

3.1 R code

Listing 2: R code of Exercise 3.9

```

1 fe <- function(n){
2   U1 = -1 + 2*runif(n)
3   U2 = -1 + 2*runif(n)
4   U3 = -1 + 2*runif(n)
5   for(i in 1:n){
6     if (abs(U3[i])>=abs(U2[i]) && abs(U3[i])>=abs(U1[i])){
7       U3[i] = U2[i]
8     }
9   }
10  U3
11 }
12
13 hist(fe(10000),col='orange',breaks=20,freq = F,xlab = 'x',ylab = 'density estimate',main='
    histogram density estimate by 10000 samples ')
14 xx = seq(-1,1,length = 1000)
15 yy = 0.75*(1-xx^2)
16 lines(xx,yy,col='blue',lwd=2)

```

3.2 histogram density estimate

The histogram (orange bars) density estimate by 10000 samples is as follows:

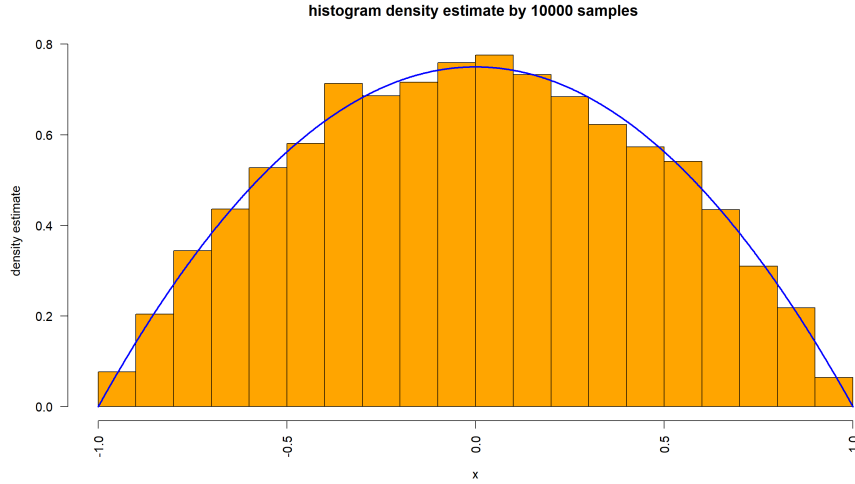


图 1: histogram density estimate of f_e

The blue line is f_e density function, which is consistent with the histogram.

4 Exercise 3.10

Assume that the variates generated by the algorithm in Exercise 3.9 is from a density $f(x)$. From the Law of Total probability, we have

$$\begin{aligned}
 f(x) &= f_{U_3}(x)P(|U_3| \text{ is not maximum}) + f_{U_2}(x)P(|U_3| \text{ is maximum}) \\
 &= f_{U_3}(x)P(|U_3| \text{ is not maximum} | U_3 = x) + f_{U_2}(x)P(|U_3| \text{ is maximum} | U_2 = x) \\
 &= \frac{1}{2}[1 - P(|U_3| \text{ is maximum} | U_3 = x)] + \frac{1}{2}P(|U_3| \text{ is maximum} | U_2 = x) \\
 &= \frac{1}{2}[1 - P(|U_1| < |x|, |U_2| < |x| | U_3 = x)] + \frac{1}{2}P(|U_3| > |U_1| | U_2 = x, |U_3| > |x|)P(|U_3| > |x| | U_2 = x) \\
 &= \frac{1}{2}(1 - x^2) + \frac{1}{2} \frac{\int_{|x|}^1 t dt}{(1 - |x|)}(1 - |x|) \\
 &= \frac{3}{4}(1 - x^2)
 \end{aligned}$$

Therefore, the algorithm given in Exercise 3.9 generates variates from the density f_e .

5 Exercise 3.12 and 3.13

5.1 R code

Listing 3: R code of Exercise 3.12 and 3.13

```

1 Exp_Gamma_mixture <- function(r,beta,n){
2   U1 = rgamma(n,r,beta)
3   U2 = rexp(n,U1)
4 }
5
6 r = 4

```

```

7 beta = 2
8 results = Exp_Gamma_mixture(r,beta,1000)
9 hist(results,col='orange',probability = T,breaks=30,xlab='x',ylab = 'density estimate',
10      main='density histogram of the sample from Exponential-Gamma mixture and Pareto density
      curve')
11 xx = seq(min(results),max(results),length=1000)
12 yy = beta^r * r * (beta+xx)^(-r-1)
13 lines(xx,yy,col='blue',lwd=2)

```

5.2 density histogram

The empirical and theoretical (Pareto) distributions are as follows:

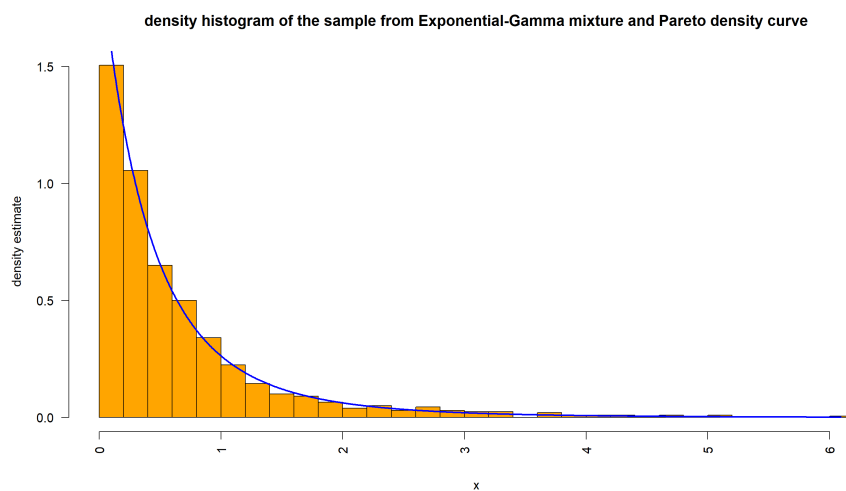


图 2: Histogram of sample from Exp-Gamma mixture and the Pareto density Curve

We can observe that the density histogram of the sample from Exponential-Gamma mixture is consistent with the Pareto density curve.

6 Exercise 3.14

6.1 R code

Listing 4: R code of Exercise 3.14

```

1 rmvn.Choleski <-
2   function(n, mu, Sigma) {
3     # generate n random vectors from MVN(mu, Sigma)
4     # dimension is inferred from mu and Sigma
5     d <- length(mu)
6     Q <- chol(Sigma) # Choleski factorization of Sigma
7     Z <- matrix(rnorm(n*d), nrow=n, ncol=d)
8     X <- Z %*% Q + matrix(mu, n, d, byrow=TRUE)
9     X
10  }
11
12 Sigma = c(1,-.5,.5,-.5,1,-.5,.5,-.5,1)

```

```

13 Sigma = matrix(Sigma,nrow=3)
14 miu = c(0,1,2)
15 samples = rmvn.Choleski(200,miu,Sigma)
16
17 X1 = samples[,1]
18 X2 = samples[,2]
19 X3 = samples[,3]
20 pairs(samples)
21 apply(samples,MARGIN = 2,FUN = mean)
22 cor(X1,X2)
23 cor(X1,X3)
24 cor(X2,X3)

```

6.2 scatter plots for each pair of variables

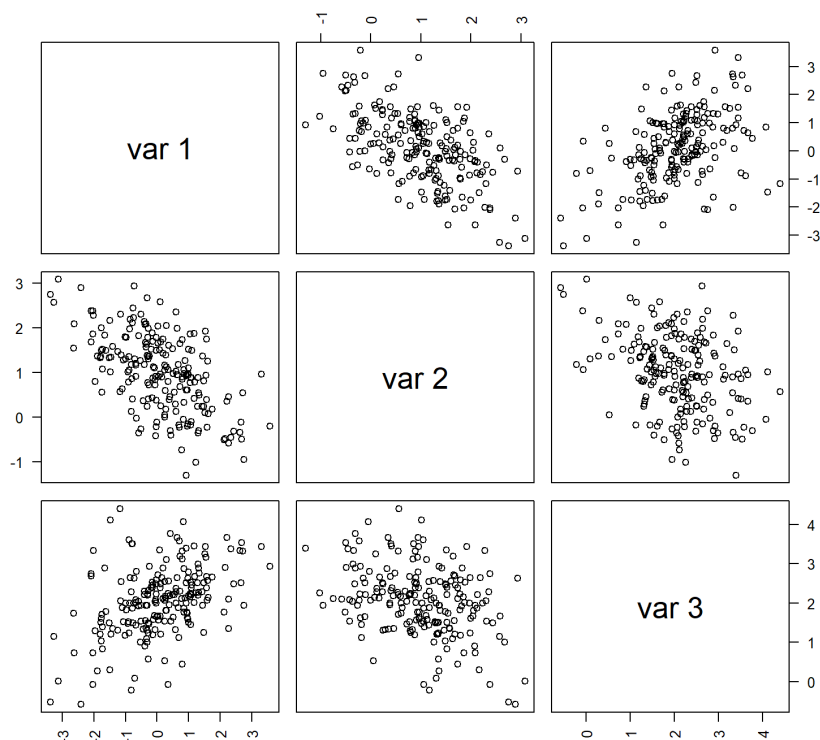


图 3: scatter plots for each pair of variables

From fig3, we can see that the scatter plot of var1 and var2 (Plot-12 for short) is centered at approximately (0,1), scatter plot of var1 and var3 (Plot-13 for short) is centered at approximately (0,2), and scatter plot of var2 and var3 (Plot-23 for short) is centered at (1,2), which correspond to the parameters $\mu = (0, 1, 2)$. On the other hand, Plot-12 and Plot-23 are distributed around the main diagonal, while Plot-13 is distributed around the secondary diagonal, which are consistent with the parameters $\rho_{12} = \rho_{23} = -0.5$, $\rho_{13} = 0.5$.