SC-HW1

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(1) Weibull distribution:

• Sampling by inverse CDF method:

 $F(x) = 1 - \exp(-(x/\theta)^{\beta}) \sim U(0,1)$, by Probability integral transformation.

Do some calculation, we have

$$F^{-1}(y) = \theta \times (\ln \frac{1}{1-y})^{1/\beta}, \text{where } Y \sim U(0,1).$$

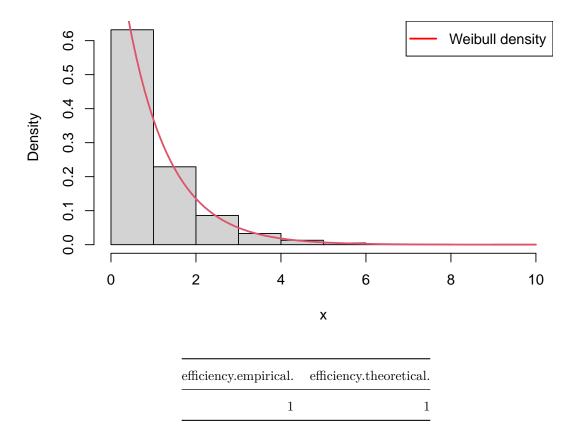
• Algorithm:

```
step 1: fixed (\theta, \beta), generate U \sim U(0, 1)
step 2: X = F^{-1}(U) \sim \text{Weibull}(\theta, \beta)
```

• $n=1000; (\theta, \beta) = (1,1)$ (i.e. Exp(1)):

```
Weibull(1000,1,1)
```

Histogram of Weibull by inverse CDF



由於這裡採用,Weibull random variable 由 Uniform random variable 藉由 inverse CDF method 轉換得來的,所以 empirical and theoretical efficiency 皆等於 1,都不受 parameters (θ,β) 所影響。由以上的例子驗證此結論。

(2) Pareto distribution:

 $\bullet\,$ Sampling by inverse CDF method:

$$F(x)=\int_0^x \frac{\beta}{\theta}(1+t/\theta)^{-(\beta+1)}dt=1-(1+x/\theta)^{-\beta}\sim U(0,1),$$
 by Probability intergral transformation.

Do some calculation, we have

$$F^{-1}(y) = \theta \times [(\frac{1}{1-y})^{1/\beta} - 1], \text{where } Y \sim U(0,1).$$

• Algorithm:

step 1: fixed
$$(\theta,\beta)$$
,
generate $U \sim U(0,1)$
step 2: $X = F^{-1}(U) \sim \operatorname{Pareto}(\theta,\beta)$

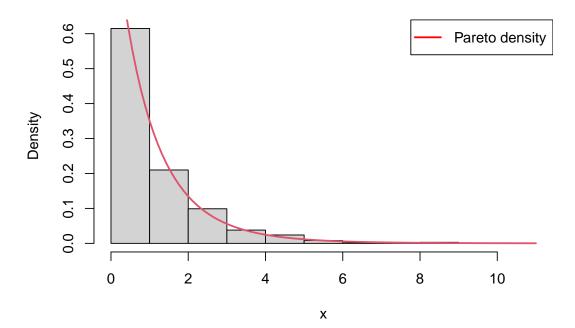
```
pareto <- function(n,scale,shape){
  x <- c()
  set.seed(2)</pre>
```

```
for(i in 1:n){
 x[i] <- scale*(1/(1-runif(1))^(1/shape)-1)
}
f = function(x,scale,shape){
 y <- c()
 for(i in 1:length(x)){
   y[i] <- shape/(scale*(1+x[i]/scale)^(shape+1))</pre>
 }
 return(y)
}
hist(x, freq = F,
     main="Histogram of Pareto by inverse CDF")
curve(f(x,scale,shape), col=2, lwd=2, add=T)
legend("topright",legend = c("Pareto density"),
     col=c("red"), lty=c(1), pch=c(-1), lwd=2)
table <- data.frame("efficiency(empirical)" = length(x)/n,
                  "efficiency(theoretical)" = 1)
return(kable(table,row.names = F))
```

• $n=1000; (\theta, \beta) = (10,10)$:

```
pareto(1000,10,10)
```

Histogram of Pareto by inverse CDF



efficiency.empirical.	efficiency.theoretical.
1	1

由於這裡採用,Pareto random variable 由 Uniform random variable 藉由 inverse CDF method 轉換得來的,所以 empirical and theoretical efficiency 皆等於 1,都不受 parameters (θ,β) 所影響。由以上的例子,驗證此結論。

(3) skewed distribution I:

• Rejection sampling:

Do some simplification to f(x):

$$f(x) = \begin{cases} \frac{2r^2}{1+r^2}\phi_1(x) = f_1(x) & \text{if } x \ge 0\\ \frac{2}{1+r^2}\phi_2(x) = f_2(x) & \text{if } x < 0 \end{cases}, \text{ where } \begin{cases} \phi_1 : \text{pdf of } N(0, r^2)\\ \phi_2 : \text{pdf of } N(0, 1/r^2) \end{cases}$$

For any x:

$$\frac{f_1(x)}{f_2(x)} = r^2 \times \frac{1/r}{r} \times \exp(-\frac{x^2}{2}(\frac{1}{r^2} - r^2)) = \begin{cases} \geq 1 \text{ if } r \geq 1\\ \in (0,1) \text{ if } r \in (0,1) \end{cases}$$

Set:

proposal
$$g(x) = \begin{cases} \phi_1(x) \ , \ \text{if} \ r \geq 1 \\ \phi_2(x) \ , \ \text{if} \ r \in (0,1) \end{cases}$$
 , $x \in \mathbb{R}$

$$M = M(r) = \begin{cases} \frac{2r^2}{1+r^2} \ , \ \text{if} \ r \ge 1 \\ \frac{2}{1+r^2} \ , \ \text{if} \ r \in (0,1) \end{cases}$$

* Algorithm:

step 1: fixed r, generate $Y\sim g$ and $U\sim U(0,1)$, Y and U are indep; step 2: let X=Y if $U\leq \frac{f(y)}{M(r)\times g(y)}$,and reject the draw otherwise.

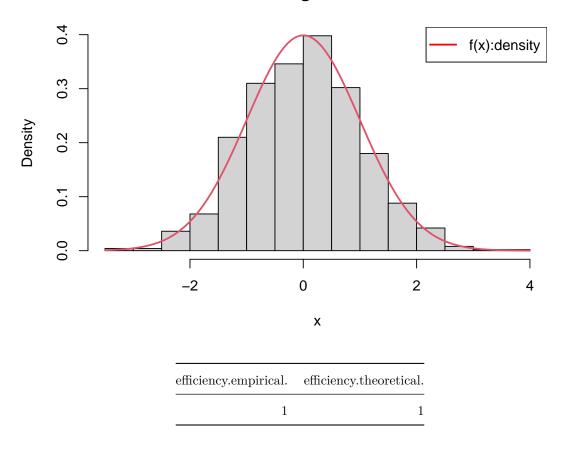
```
rej_3 <- function(n,r){
    M = function(r){
    z = 0
    if (r >=1){ z = 2*r^2/(1+r^2)}
    else { z = 2/(1+r^2)}
    return(z)
}
```

```
y = c()
  for (i in 1:length(x)){
    if (x[i] < 0) \{y[i] < -2/(1+r^2)*dnorm(x[i],mean = 0,sd = 1/r)\}
    else{y[i] \leftarrow 2*r^2/(1+r^2)*dnorm(x[i],mean = 0,sd = r)}
  }
  return(y)
g = function(x,r){
  y <- c()
 if (r \ge 1)\{y \leftarrow dnorm(x,mean = 0,sd = r)\}
  else{y \leftarrow dnorm(x,mean = 0,sd = 1/r)}
 return(y)
}
set.seed(3)
y <-c()
if (r >= 1){y \leftarrow rnorm(n, mean = 0, sd = r)}
else{y <- rnorm(n,mean = 0,sd = 1/r)}
u <- runif(n,0,1)
x = y[u \le f(y,r)/(M(r)*g(y,r))]
hist(x, 20, probability=T)
curve(f(x,r), col=2, lwd=2, add=T)
legend("topright",legend = c("f(x):density"),
     col=c("red"), lty=c(1), pch=c(-1), lwd=2)
table <- data.frame("efficiency(empirical)" = length(x)/n,</pre>
                     "efficiency(theoretical)" = 1/M(r))
return(kable(table,row.names = F))
```

• n=1000; r=1:

rej_3(1000,1)

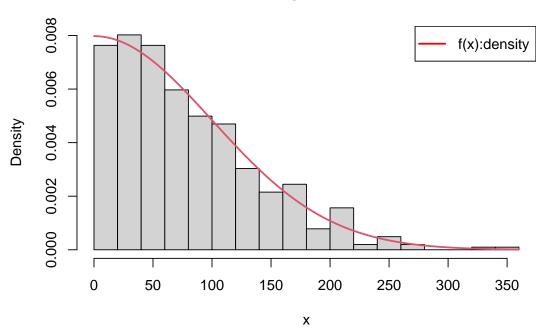
Histogram of x



• n=1000; r = 100:

rej_3(1000,100)

Histogram of x

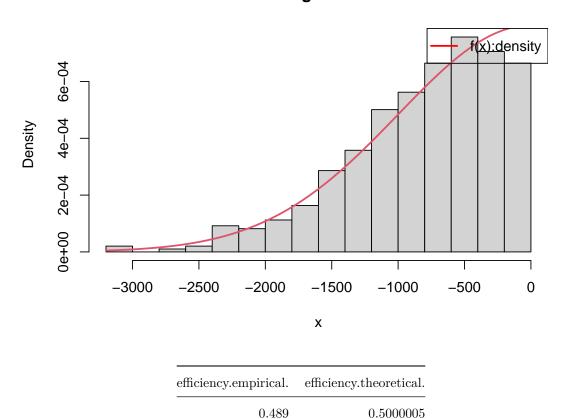


efficiency.empirical.	efficiency.theoretical.
0.511	0.50005

• n=1000; r=0.001:

rej_3(1000,0.001)

Histogram of x



由於 M=M(r) ,為一個 v 的函數,故 theoretical efficiency =1/M=1/M(r),其 efficiency 會受 parameter r 影響。另外,以上任取三個不同的 r ,從所得到的模擬結果,可以觀察出其 empirical

(4) skewed distribution II:

efficiency 一致於 theoretical efficiency。由以上的例子,驗證此結論。

• Rejection sampling:

$$\begin{split} f(x) &= 2h(x)G(\alpha x) \\ &\leq 2h(x) \times 1 \ (\because G(\alpha x) \leq 1, \forall x) \end{split}$$

```
Set: M=2 \label{eq:monoposition} proposal g(x)=h(x): pdf of the t_v distribution
```

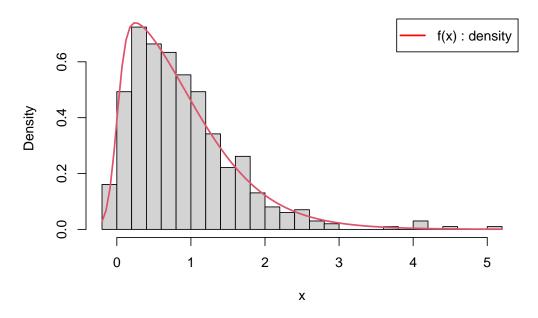
• Algorithm:

```
step 1: generate Y \sim t_v and U \sim U(0,1), Y and U are indep; step 2: let X=Y if U \leq \frac{f(y)}{2 \times g(y)},and reject the draw otherwise.
```

```
rej_4 <- function(n,alpha,v){</pre>
  M=2
  f = function(x,alpha,v){
    y = c()
    for(i in 1:length(x)){
      y[i] \leftarrow 2*dt(x[i], df = v)*pt(alpha*x[i], df = v)
    }
    return(y)
  }
  #g = dt(x,v)
  set.seed(4)
  u = runif(n,0,1)
  y \leftarrow rt(n, df = v)
  x = y[u \le f(y,alpha,v)/(M*dt(x = y,df = v))]
  hist(x, 20, probability=T)
  curve(f(x,alpha,v), col=2, lwd=2, add=T)
  legend("topright",legend = c("f(x) : density"),
       col=c("red"), lty=c(1), pch=c(-1), lwd=2)
  table <- data.frame("efficiency(empirical)" = length(x)/n,
                       "efficiency(theoretical)" = 1/M)
  return(kable(table,row.names = F))
```

• n=1000; $(\alpha, v) = (10, 10)$:

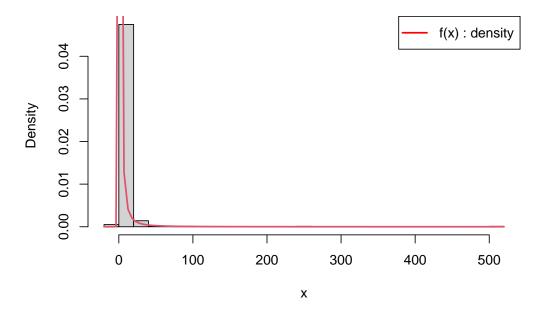
Histogram of x



efficiency.empirical.	efficiency.theoretical.
0.497	0.5

• n=1000 ; $(\alpha, v) = (99, 1)$:

Histogram of x



efficiency.empirical.	efficiency.theoretical.
0.495	0.5

由於 M=2 ,為一個常數,故 theoretical efficiency =1/M=0.5,其 efficiency 不受 parameter (α,v) 影響。另外,以上是任取 (α,v) 所得到的模擬結果,可以觀察出其 empirical efficiency 一致於 theoretical efficiency。由以上的例子,驗證此結論。

(5) 2-dimension:

• Gibbs sampling:

We firstly calculate f(x|y) and f(y|x):

$$\begin{split} f(x) &= \int_0^1 f(x,y) dy = 2(1-x) \int_0^1 (1-y)(1-xy)^{-3} dy = 1. \\ f(y) &= \text{similar to above} = 1. \\ f(x|y) &= 2(1-x)(1-y)(1-xy)^{-3}, \forall x \in (0,1) \text{ for fixed } y. \\ f(y|x) &= 2(1-x)(1-y)(1-xy)^{-3}, \forall y \in (0,1) \text{ for fixed } x. \end{split}$$

Consider the rejection sampling:

By probability integral transformation,

$$\begin{split} Z_X &= 1 - F_{X|y}(x) = \int_x^1 f(t|y) dt \\ &= 2(1-y) [\frac{1}{2y}(1-t)(1-ty)^{-2}]_x^1 + \frac{1}{2y} \int_x^1 (1-ty)^{-2} dt] \\ &= 2(1-y) [\frac{-(1-x)}{2y(1-xy)^2} + \frac{(1-x)}{2y(1-y)(1-xy)}] \\ &= \frac{-(1-x)(1-y)}{y(1-xy)^2} + \frac{1-x}{y(1-xy)} \\ &= (\frac{1-x}{1-xy})^2 \sim U(0,1). \end{split}$$
 Similarly, $Z_Y = 1 - F_{Y|x}(y) = (\frac{1-y}{1-xy})^2 \sim U(0,1).$

Do some calculation, we have

$$X|y = \frac{\sqrt{Z_X} - 1}{y\sqrt{Z_X} - 1}.$$

$$Y|x = \frac{\sqrt{Z_Y} - 1}{x\sqrt{Z_Y} - 1}.$$

• Algorithm:

step 1: Set $x^{(0)}, y^{(0)}$ be the initial value , both $\in (0, 1)$.

step 2: t=t+1, by inverse CDF method,draw

```
X \sim f(x|y^{(t-1)}), \text{denoted as } x^{(t)} Y \sim f(y|x^{(t-1)}), \text{denoted as } y^{(t)}
```

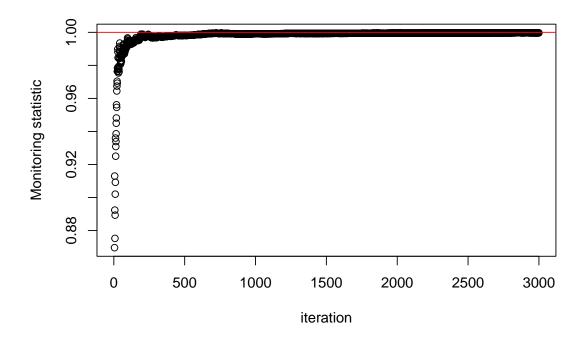
step 3: repeat step 2 until converge.

- Check covergence:
- 1. Set iteration:3000, $x^{(0)} = 0.5, y^{(0)} = 0.5$, generating the samples (X,Y)

```
gibb <- function(init, n.iter){</pre>
  xx <- matrix(init,1,2)</pre>
  x \leftarrow xx[1,1]
  for (i in 1:(2*n.iter)){
    zy <- runif(1,0,1)</pre>
    zx <- runif(1,0,1)</pre>
    y \leftarrow (sqrt(zy)-1)/(sqrt(zy)*x-1)
    xx \leftarrow rbind(xx,c(x,y))
    x \leftarrow (sqrt(zx)-1)/(sqrt(zx)*y-1)
    xx \leftarrow rbind(xx,c(x,y))
  }
  return(xx[2*(1:n.iter)-1,])
}
set.seed(231)
n.iter <- 3000
init <-c(0.5,0.5)
test <- gibb(init=init, n.iter=n.iter)</pre>
```

2. Check Monitoring statistic $\sqrt{\hat{R_n}}$:

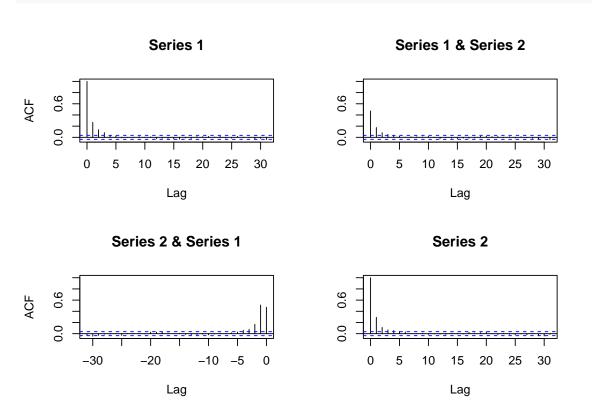
```
gelman.Rhat <- function(mc.draw){
    n.iter <- dim(mc.draw)[1]
    b1 <- w <-rep(0, n.iter)
    for (i in 5:n.iter){
        b1[i] <-var(apply(mc.draw[1:i,],2,mean)) # b1 = b_n/n
        w[i] <-mean(apply(mc.draw[1:i,],2,var)) # within variance: w_n
    }
    Rhat = b1/w+((1:n.iter)-1)/(1:n.iter)
    return(Rhat)
}
plot(gelman.Rhat(test),xlab ="iteration",ylab = "Monitoring statistic")
abline(h=1,col=c("red"))</pre>
```



由上可觀察出, $\sqrt{\hat{R_n}} \xrightarrow{\text{n is large}} 1$,根據理論,所有的抽樣服從一個穩健分布。

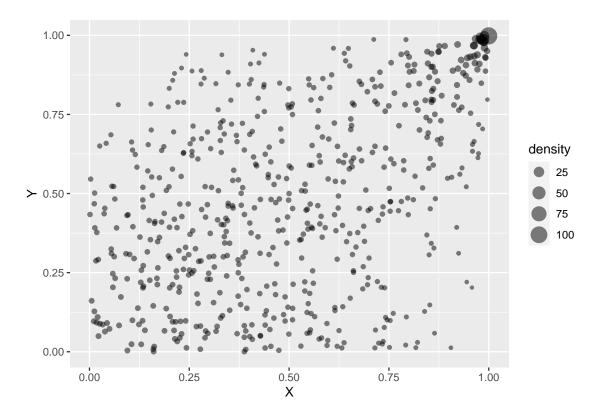
• Check serial correlations:





由上可以看出,大約每 5 個 lag 其 autocorrelation 趨於 0,於是這裡從 3000 個模擬出來的樣本中,抽取 $(X_1,Y_1),(X_6,Y_6),....,(X_{2996},Y_{2996})$ 來觀察其 Bubble plot 與原始的 f(x,y) plot 做個對比:

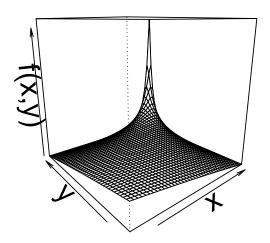
(i) Bubble plot



可觀察出,以"f(x,y)"的大小作為點的形狀大小,可以發現點座標越靠近"X=1 和 Y=1" 時,其 f(x,y) 會非常大。

(ii) density plot

```
f <- function(x){
   2*(1-x[1])*(1-x[2])*(1-x[1]*x[2])^(-3)
}
grid1 <- seq(0.05,1,length=40)</pre>
```



可觀察出模擬結果 (Bubble plot) 與原始的 f(x,y) plot 有一定程度上的相似性。 最後,由於這裡採用的抽樣樣本是用 inverse CDF method 轉換得來的,所以 empirical and theoretical efficiency 皆等於 1。