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# TD - Exercises in R on uncertainty propagation
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# Bertrand Iooss - Fabrice Gamboa - Barranquilla - 2015
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```
# SOLUTIONS
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#####
```

```
rm(list=ls())
```

```
graphics.off()
```

```
library(triangle)
```

```
library(mistral)
```

```
#####
```

```
#2.1
```

```
library(triangle)
```

```
# 2 solutions for visualizing the probability densities
```

```
# with plot and density
```

```
N <- 1e6
```

```
x11()
```

```
par(mfcol=c(2,2))
```

```
F1 <- rnorm(N,3e4,9e3)
```

```

plot(density(F1))
E1 <- rtriangle(N,2.8e7,4.8e7,3e7)
plot(density(E1))
L1 <- runif(N,250,260)
plot(density(L1))
l1 <- rtriangle(N,310,450,400)
plot(density(l1))

# or with curve
par(mfcol=c(2,2))
curve(dnorm(x,3e4,9e3),from = -10000 , to = 72000, col='blue',xlab='F',ylab='pdf',lwd=2)
curve(dtriangle(x,2.8e7,4.8e7,3e7),from = 2.7e7 , to = 4.9e7,col='red',xlab='E',ylab='pdf',lwd=2)
curve(dunif(x,250,260),from = 248 , to = 262,col=rgb(0.3,0.7,0.3),xlab='L',ylab='pdf',lwd=2)
curve(dtriangle(x,310,450,400),from = 305 , to = 455,col='orange',xlab='I',ylab='pdf',lwd=2)

```

#2.2

```

beam <- function(x){
  # parameters in the following order: F, E, L, I
  # this function takes as inputs a n x d matrix and returns a n-size vector as output
  if (is.vector(x)){x <- matrix(x,nrow=1)} # turning a vector to a matrix
  result <- (x[,1]*x[,3]^3)/(3*x[,2]*x[,4])
  return(result)
}

```

#2.3

```

N <- 1000

```

```

F2 <- rnorm(N,3e4,9e3)
E2 <- rtriangle(N,2.8e7,4.8e7,3e7)
L2 <- runif(N,250,260)
I2 <- rtriangle(N,310,450,400)
data = cbind(F2,E2,L2,I2)
z = beam(data)
mean(z) ; var(z)

summary(z)

x11() ; hist(z)
?hist
hist(z,probability=TRUE,col='lightblue',main='histogram of the beam')
lines(density(z),col='blue',lwd=2)

# convergence :

m_hat_cv <- c()      # initialisation
var_hat_cv <- c()

seqn = c(seq(10,5000,by=50))# sample size in the loop

for (k in seqn){ # loop
  Fk <- rnorm(k,3e4,9e3)
  Ek <- rtriangle(k,2.8e7,4.8e7,3e7)
  Lk <- runif(k,250,260)
  Ik <- rtriangle(k,310,450,400)

```

```

datak = cbind(Fk,Ek,Lk,lk)

Yk = beam(datak)

m_hat_cv <- c(m_hat_cv,mean(Yk))

var_hat_cv <- c(var_hat_cv,var(Yk))

}

# graph

x11()

par(mfrow = c(2, 1))

plot(seqn, m_hat_cv,col='blue',type='l',xlab='n',ylab='mean estimation',main='mean convergence')

plot(seqn, var_hat_cv,col='red',type='l',xlab='n',ylab='variance estimation',main='variance
convergence')

```

#2.4

```
F <- 3e4 ; E <- 3e7 ; L <- 250 ; l <- 400 # nominal point
```

```
Moycum <- beam(cbind(F,E,L,l))
```

```
print(Moycum)
```

```
help(deriv) # Formal differentiation
```

```
DE <- deriv(y ~ ( F * L^3 ) / ( 3 * E * l ),c("F","E","L","l"))
```

```
print(DE)
```

```
#print(eval(DE))
```

```
grad <- eval(DE)
```

```
grad <- attributes(grad)$gradient[1,]
```

```
print("Dérivées exactes :")
```

```
print(grad)
```

```
#Quadratic summation
```

```
Varcum <- grad[1]^2*var(F1)+grad[2]^2*var(E1)+grad[3]^2*var(L1)+grad[4]^2*var(I1)
print(Varcum)
```

```
#2.5
```

```
quantile(z,0.95) # empirical quantile
```

```
#####
```

```
#3.1
```

```
failure <- function(x){
  # there is a failure when this function is negative
  # parameters are in the following order: F, E, L, I
  # this function takes as inputs a n x d matrix and returns a n-size vector as output

  if (is.vector(x)){x <- matrix(x,nrow=1)}
  result <- 30 - (x[,1]*x[,3]^3)/(3*x[,2]*x[,4])
  return(result)
}
```

```
#3.2
```

```
n <- 10000
```

```
# radom sampling of inputs
```

```
F1 <- rnorm(n,3e4,9e3)
```

```
E1 <- rtriangle(n,2.8e7,4.8e7,3e7)
```

```
L1 <- runif(n,250,260)
```

```
I1 <- rtriangle(n,310,450,400)
```

```
data <- cbind(F1,E1,L1,I1)
```

```
fail_Y <- failure(data)
```

```
# use function sum
```

```
pf = sum(fail_Y < 0) / n
```

```
cv <- sqrt((1-pf)/(n*pf))
```

```
#3.3
```

```
cvlim <- 0.015
```

```
Nlim <- 1e9
```

```
N = 1e6
```

```
Nrun = N
```

```
cv = 1
```

```
z = NULL
```

```
res = NULL
```

```
while ((Nrun <= Nlim) & (cv > cvlim)){
```

```
  F2 <- rnorm(N,3e4,9e3)
```

```
  E2 <- rtriangle(N,2.8e7,4.8e7,3e7)
```

```
  L2 <- runif(N,250,260)
```

```
  I2 <- rtriangle(N,310,450,400)
```

```
  data = cbind(F2,E2,L2,I2)
```

```
  z = c(z,failure(data))
```

```
  pf <- sum(z < 0) / Nrun
```

```
cv <- sqrt((1-pf)/(Nrun*pf))
Nrun <- length(z)
res <- rbind(res,c(Nrun,pf,cv))
print(c(Nrun,pf,cv))
}
x11()
plot(res[,1],res[,2])
```

#3.4

```
eps = 1e-7
Method = "AR"
IS = FALSE
N.calls = 2000
q = 1
u.dep = c(0,0,0,0)

choice.law = list()
choice.law[[1]] = list("norm",c(3e4,9e3))
choice.law[[2]] = list("triangle",c(2.8e7,4.8e7,3e7))
choice.law[[3]] = list("unif",c(250,260))
choice.law[[4]] = list("triangle",c(310,450,400))

res=FORM(f=failure,u.dep=u.dep,choice.law=choice.law,
        N.calls=N.calls,eps=eps,Method=Method,IS=IS,q=q)
print(res)
```

#3.5

eps = 1e-7

Method = "AR"

IS = TRUE

N.calls = 2000

q = 0.1

u.dep = c(0.1,0,0,0)

choice.law = list()

choice.law[[1]] = list("norm",c(3e4,9e3))

choice.law[[2]] = list("triangle",c(2.8e7,4.8e7,3e7))

choice.law[[3]] = list("unif",c(250,260))

choice.law[[4]] = list("triangle",c(310,450,400))

res=FORM(f=failure,u.dep=u.dep,choice.law=choice.law,

N.calls=N.calls,eps=eps,Method=Method,IS=IS,q=q)

print(res)