



Introduction to R

Yao Thibaut Kpegli

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Introduction

Purpose

- A quick and intensive reminder about R (4.5 hours)
 - to make you operational during your econometrics classes over the year

R: definition

R is a free software to conduct statistical and econometrics analysis, optimization, simulation, numerical computation, etc...

Installation

Installation

Download two distinct elements :

- download R from “comprehensive R archive network” (CRAN)
<https://cran.r-project.org/>
 - Execute the installation program
 - Keep default parameters
- Download RStudio, for example from
<https://posit.co/download/rstudio-desktop/>
 - Execute the installation program
 - Keep default parameters

Assign operator

Assign operator

create a scalar `a` that takes value 2

- `a = 2`
- `a <- 2`
- `2 -> a`

Common operations

Common operations

- addition : $2+2$
- subtraction $2-2$
- product : $2*2$
- division: $2/2$

Common math functions

Common math functions

- square/exponent : 2^2
- square root: `sqrt(2)` or `2**0.5`
- exponential : `exp(4)`
- logarithm : `log(4)`
- min: `min(2,3)`
- max: `max(2,3)`
- absolute val.: `abs(-4)`
- sign: `sign(-10)`
- round: `round(4.56789, 2)`
- ceiling: `ceiling(4.56789)`
- integer: `as.integer(4.56789)`

Vectors

Specific vectors

- repetition: `rep(1, 10)`
- sequence: `seq(from = 0, to = 20, by = 1)`
 - even sequence: `seq(from = 0, to = 20, by = 2)`
 - odd sequence: `seq(from = 1, to = 21, by = 2)`
- trend: `1980:2023`

Vectors

- `c(1,3,2,0,4,5)`
 $(1 \ 3 \ 4 \ 0 \ 4 \ 5)$
 - second element of the vector: `c(1,3,2,0,4,5)[2]`
 - standard-deviation : `sd(c(1,3,2,0,4,5))`
 - sum : `sum(c(1,3,2,0,4,5))`
 - median: `median(c(1,3,2,0,4,5))`
 - sort: `sort(c(1,3,2,0,4,5))`
 - rank: `rank(c(1,3,2,0,4,5))`
- vector of characters: `c("abcd", "a", "ab", "abc")`

Matrix operations

Matrix operations

- `matrix(1:6, nrow = 2)` or `matrix(1:6, ncol = 3)`

$$\begin{pmatrix} 1 & 3 & 5 \\ 2 & 4 & 6 \end{pmatrix}$$

- by default, R arranges elements by column "byrow = F"... Try `matrix(1:6, nrow = 2, byrow = T)`
- nb of row : `nrow(matrix(1:6, nrow = 2))`
- nb of col: `ncol(matrix(1:6, nrow = 2))`
- element at the line $i=1$ and column $j=3$: `matrix(1:6, nrow = 2)[1,3]`
- extract first column: `matrix(1:6, nrow = 2)[,1]`
- extract the third line: `matrix(1:6, nrow = 2)[2,]`
- extract the first and third columns: `matrix(1:6, nrow = 2)[1:2, c(1,3)]` or `matrix(1:6, nrow = 2)[1:2, c(1,3)]`

Matrix operations

Matrix operations

- transpose: `t(matrix(1:6, nrow = 2))`
- dimension: `dim(matrix(1:6, nrow = 2))`
- matrix product: `matrix(1:6, nrow = 2)%*%matrix(1:6, nrow = 3)`
- determinant: `det(matrix(1:6, nrow = 2)%*%matrix(1:6, nrow = 3))`
- eigenvalues : `eigen(matrix(1:6, nrow = 2)%*%matrix(1:6, nrow = 3))`
- inverse: `solve(matrix(1:6, nrow = 2)%*%matrix(1:6, nrow = 3))`

cross and Kronecker products

cross product

$$A = \begin{pmatrix} 0.5 \\ 2 \end{pmatrix} \quad \text{and} \quad B = \begin{pmatrix} 1 & 0 \\ 1 & 1 \end{pmatrix}$$

- cross product: `crossprod(A,B) = t(A)%*%B`
 - result: (2.5 , 2)
- Kronecker product: `kron(A, B)`
 - result:

$$\begin{pmatrix} 0.5 & 0 \\ 0.5 & 0.5 \\ 2 & 0 \\ 2 & 2 \end{pmatrix}$$

Matrix operations

Matrix operations

- `c(1,3,2,0,4,5)`
$$(\ 1 \ 3 \ 4 \ 0 \ 4 \ 5 \)$$
- `matrix(c(1,3,2,0,4,5), nrow = 2)`
- `matrix(c(1,3,2,0,4,5), nrow = 2, byrow=F)`
- `matrix(c(1,3,2,0,4,5), nrow = 2, byrow=T)`

identity matrix

`diag(3)`

$$\begin{pmatrix} 1 & 0 & 0 \\ 0 & 1 & 0 \\ 0 & 0 & 1 \end{pmatrix}$$

Matrix operations

rbind

- rbind with (1,3,2) and (0,4,5)
 - c(1,3,2) and c(0,4,5)
 - rbind(c(1,3,2),c(0,4,5))

$$\begin{pmatrix} 1 & 3 & 2 \\ 0 & 4 & 5 \end{pmatrix}$$

cbind

- cbind with (1,3,2) and (0,4,5)
 - c(1,3,2) and c(0,4,5)
 - cbind(c(1,3,2),c(0,4,5))

$$\begin{pmatrix} 1 & 0 \\ 3 & 4 \\ 2 & 5 \end{pmatrix}$$

Exercise

- Create a matrix X that takes the values

$$\begin{pmatrix} 1 & 1 \\ 1 & 4 \\ 1 & 2 \\ 1 & 5 \end{pmatrix}$$

- Compute $B = (X'X)^{-1}$
 - determine the dimension
 - extract the elements on the diagonal
- Create a matrix Y that takes value

$$\begin{pmatrix} 3 \\ 6 \\ 4 \\ 7 \end{pmatrix}$$

- Compute $(X'X)^{-1}X'Y$

Result

- `X <- cbind(1,c(1,4,2,5))`
- `B <- solve(crossprod(X,X))`
- `dim(B)`
- `diag(B)`
- `Y <- matrix(c(3,6,4,7),nrow=4)`
- `OLS <- B%*%crossprod(X,Y)`
- `OLS`

Working Directory

Working Directory

- Useful to import/export data and results
 - 1 set the work directory: `setwd("D:/ENS Paris Saclay/R2023_2024")`
 - 2 check what is work directory: `getwd()`

Definition

Definition

- dataframe = matrix of data
 - set of vectors with the same length
 - placed next to each other vertically
- Column = Variable
 - possible of different types: quantitative, numerical but also qualitative, characters, dates.
- Line = Observation

creation

Creation

- `example <- data.frame(one = 1:10, two = 11:20, three = 21:30)`
- or equivalently
 - `example <- as.data.frame(matrix(1:30, ncol=3))`
 - `names(example) <- c("one", "two", "three")`
- display data : `View(example)`

basic data manipulation

Useful symbols

- Strict inequality : $>$, $<$
- equal or unequal : $=<$, $>=$
- equal: $==$
- different equal: $!=$
- and : $\&$
- or : $|$

basic data manipulation

- mean: `mean(example$one)`
- stat des: `summary(example$one)`
- rename var.: `names(example)[names(example) == "ratio"] <- "new"`
- replace obs: `example$one[example$one==3] <- 5`

attach, detach, and with

attach, detach, and with

- “attach” allows to avoid referring to the dataframe at each line of code
 - e.g: attach(example), then directly write
 - mean(one)
 - min(two)
 - sd(two)
- “detach” is used to clear the “attach” command
- “attach” for a specific command: with(example, mean(two))

new variable

new variable

- new var.: `example$ratio <- (example$one)/(example$two)`
- dummy (binary) var.: `example$dumy_ch <- ifelse(example$one <= 5, "Low", "High")`
- dummy (binary) var.: `example$dumy_num <- ifelse(example$one <= 5, 1, 0)`

Subset of dataframe

Subset of dataframe

- keep lines with $two \leq 16$: `set_obs <- subset(example, two <= 16)`
- delete column two: `set_var <- subset(example, select = -two)`
- keep columns one and three : `set_varA <- subset(example, select = c("one", "three"))`
- combination of conditions: `set_obs_var <- subset(example, two <= 16, select = -two)`

Exportation

exportation in text file

```
write.table(example, file="example.txt", col.names=TRUE)
```

exportation in csv file

```
write.csv(example, file="example.csv")
```

exportation in excel file

- 1 install package: `install.packages("writexl")`
- 2 call library: `library("writexl")`
- 3 export: `write_xlsx(example, "example.xlsx")`

Importation

Importation of text file

```
text_file <- read.table("example.txt", header=TRUE)
```

exportation of csv file

```
read.csv(example, file="example.csv")
```

Importation of Excel file

- 1 install package: `install.packages("xlsx")`
- 2 `library(xlsx)`
- 3 `Excel_file <- read.xlsx("example.xlsx")`

Exercise

- Use data "Journals" within the package AER of R:
`data("Journals", package = "AER")`
- Create a new variable `citeprice = price/citations`
- Attach the dataframe Journal
- Use the codes from this class to compute:
 - OLS estimator of the regression
 $\log(subs) = a \times \log(price/citations) + b + \epsilon :$

$$(X'X)^{-1}X'Y$$

- matrix of variance-covariance:

$$\hat{\sigma}_{\epsilon}^2(X'X)^{-1}$$

- t-stat
- compare the results with those obtained from "lm(.)"
command

Exercise: result

- `install.packages("AER")`
- `data("Journals", package = "AER")`
- `Journals$citeprice <- Journals$price/Journals$ Citations`
- `names(Journals)`
- `View(Journals)`
- `attach(Journals)`
- `X<-cbind(log(citeprice),1)`
- `Y<-log(subs)`
- `beta<-solve(crossprod(X,X))%*%t(X)%*%Y ; beta`
- `e<-Y-X%*%beta ; e`
- `N<-nrow(Journals) ; N`
- `vcm<- (kronecker(crossprod(e,e)/(N-2),solve(crossprod(X,X)))) ;
vcm`
- `sd<-sqrt(diag(vcm))`
- `sd`
- `tstat<-beta/sd`
- `tstat`
- `lms= lm(logsubs logpc, data = Journals)`
- `summary(lms)`

Vertical merge of data

Vertical merge of data

- `v1=data.frame(Num =c(1,-1,0,3,0,2),
Str=c("L1","L2","L3","M1","M2","D1"))`
- `v2=data.frame(Num=c(10,20,30,40),
Str=c("L1","L2","M1","D3"))`
- `v <- rbind(d1,d2)`

Horizontal merge

Horizontal merge

- `h1=data.frame(id =c(1,2,3,4,5,6), Num =c(1,-1,0,3,0,2), Str=c("L1","L2","L3","M1","M2","D1"))`
- `h2=data.frame(id =c(1,2,3,4,5,7), Name=c("Rac","Elo","Fra","Hon","Hor","Ben"))`
- inner join: `merge_inner <- merge(x=h1,y=h2,by="id")`
- left join: `merge_left <- merge(x=h1,y=h2,by="id", all.x=T)`
- right join: `merge_right <- merge(x=h1,y=h2,by="id", all.y=T)`
- outer join: `merge_outer <- merge(x=h1,y=h2,by="id", all=T)`

Scatter plot

ggplot2

- install package: `install.packages("ggplot2")`
- scatter plot:
`ggplot(Journals, aes(log(citeprice),log(subs))) +
geom_point()`
- add colors (as a third dimension): `ggplot(Journals,
aes(log(citeprice),log(subs))) + geom_point(aes(color =
pages))`
- add linear fit:
`ggplot(Journals, aes(log(citeprice),log(subs))) +
geom_point(aes(color = pages)) + geom_smooth(method =
"lm", se=F)`
- add linear fit+confidence interval:
`ggplot(Journals, aes(log(citeprice),log(subs))) +
geom_point(aes(color = pages)) + geom_smooth(method =
"lm", se=T)`

common plots

bar plot

```
ggplot(Journals) + geom_bar(aes(society))
```

histogram

```
ggplot(Journals) + geom_histogram(aes(log(subs)))
```

empirical cumulative density function

```
ggplot(Journals) + stat_ecdf(aes(log(subs)))
```

empirical density

```
ggplot(Journals) + geom_density(aes(log(subs)))
```

box plot

```
ggplot(Journals) + geom_boxplot(aes(subs))
```

Function

- create function $f(\alpha) = \alpha^2$

```
f <- function(alpha){  
  y <- alpha**2  
  return(y)  
}
```
- computation:
 - at a specific value: $f(0)$
 - on a range: $f(-10:10)$
 - on a range: $f(\text{seq}(\text{from}=-2, \text{to}=2, \text{by}=0.1))$

Function

- create function $f(x) = x^2 + y^2$

```
f <- function(x,y){  
  z <- x**2+y**2  
  return(z)  
}
```
- computation:
 - at a specific value: `f(1,1)`
 - on a range: `f(0:2,-1:1)`
 - on a range: `f(seq(from=0,to=2, by=0.1),seq(from=-,to=1, by=0.1))`

Function

- create function $f(x) = x^2 + y^2$

```
f <- function(x){  
  z <- x[1]**2 + x[2]**2  
  return(z)  
}
```
- computation:
 - at a specific value: $f(c(1,1))$

grid

grid

- ① Two vectors: v1 and v2
- ② To get all possible combinations of elements from v1 and v2
 - use “`expand.grid(v1,v2)`”
 - valid also for more than two vectors: `expand.grid(v1,v2,...,vn)`

grid

- vector: `v1<- c(-1:1)`
- vector 2: `v2 <- c(0:2)`
- grid: `expand.grid(v1,v2)`

Exercise

- create a vector x taking values from -1 to 1, by a step of 0.01
- create a vector y taking values from -1 to 1, by a step of 0.01
- create a grid of values of x and y
- compute $f(x, y)$ over the grid
- plot the function over the grid
 - define the color option of ggplot on the $\log(f)$
 - use the 3D plot

Result

- vector x : `x<- seq(from=-1, to=1,by=0.01)`
- vector y : `y<- seq(from=-1, to=1,by=0.01)`
- grid : `data <- data.frame(expand.grid(x,y))`
- `names(data) <- c("x","y")`
- `data$z <- (data$x)**2 + (data$y)**2`
- plot:
 - `ggplot ggplot(data, aes(x,y)) + geom_point(aes(color = log(z)))+ scale_color_gradientn(colours = rainbow(4))`
 - 3D plot
 - `library("plotly")`
 - `plot_ly(x=data$x, y=data$y, z=data$z, type="scatter3d", color=data$z, mode="markers")`

Optimization

$$\min_x f(x) = x^2$$

- create function $f(x) = x^2$

```
f <- function(x){  
  y= x**2  
  return(y)  
}
```
- initial values:

```
initial_value <- 5
```
- use “optim(...)” function that **minimizes** functions:

```
optim(initial_value, f, method="BFGS")
```

Optimization

$$\max_x f(x) = x - 0.5x^2$$

- create function $f(x) = x - 0.5x^2$

```
f <- function(x){  
  y= x - 0.5*x**2  
  return(-y)  
}
```
- initial values:

```
initial_value <- 5
```
- use “optim(...)” function that **minimizes** functions:

```
optim(initial_value, f, method="BFGS")
```

Optimization

$$\min_{x,y} f(x,y) = x^2 + y^2$$

- create function $f(x) = x^2 + y^2$

```
f <- function(x){  
  y = x[1]**2+x[2]**2  
  return(y)  
}
```
- initial values:

```
initial_value <- c(10, 10)
```
- use “optim(...)” function:

```
optim(initial_value, f, method="BFGS")
```

Optimization over dataframe (e.g. OLS)

```
Y <- log(Journals$subs)
X <- log(Journals$price/Journals$citations)
ols <- function(z){
  sse <- sum((Y - z[1]*X-z[2])^2)
  return(sse)
}
initial_value <- c(1, 1)
optim(initial_value, ols, method="BFGS")
```

for

for

```
n <- nrow(Journals)
count <- 0
for (i in 1:n) {
  if(Journals$society[i] == "yes"){count = count+1}
}
print(count)
```


while

while

```
i <- 1
while (Journals$society[i] == "no") {
  i = i+1
}
print(i)
```

Random values and re-allocation

useful random values

- normal: `rnorm(n, mu, sigma)`
 - vector of 10 random values of $N(0, 1)$: `rnorm(10, 0, 1)`
- uniform: `runif(n, a, b)`
 - vector of 10 random values of $U[0, 1]$: `runif(10, 0, 1)`

random re-allocation

- `sample(vector)`
 - `v <- 1:10`
 - `sample(v)`

Exercise

- ① First replication
 - simulates a vector ϵ of $n = 1000$ random values of $N(0, 2)$
 - simulates a vector x of $n = 1000$ random values of $U(-5, 5)$
 - generate a vector $y = 2 \times x + 1 + \epsilon$
 - compute OLS estimator of the regression of y on x
 - save the OLS estimates
- ② 1000 replications
 - Repeat the first question 1000 times
- ③ Compute the mean of the OLS estimates over the 1000 replications
- ④ Compare with the true values of parameters: 2 and 1

Result

- nb of replication: `rep <- 10000`
- sample size: `n <- 1000`
- true coef: `coef <- c(2,1)`
- stock vector: `stock <- matrix(rep(NA,2*rep), ncol=2)`
- loop “for” to make replications:
 for (i in 1:rep){
 sim <- data.frame(x=runif(n,-5,5),er=rnorm(n,0,2))
 sim\$y = coef[1]*sim\$x + coef[2] + sim\$er
 lms= lm(y ~ x, data = sim)
 stock[i,] <- lms\$coefficients
 }
- compute the mean: `Mean <-
 c(mean(stock[,1]),mean(stock[,2]))`
- show the result: `Mean`

Exercise

- provides an approximated value for $F(x)$

$$F(x) = \int_{-10}^{10} f(x) dx$$

with $f(\cdot)$ the density function of the standard normal distribution, and $\pi \simeq 3.14$

- hint: $F(x) = (10 - (-10)) \times E[f(x)]$, with $E[\cdot]$ the expectation computed with $U[-10, 10]$
 - this value is almost 1
- 1 simulates a vector x of $n = 100000000$ random values of $U(-10, 10)$
 - 2 compute $f(\cdot)$ over the simulated values of x
 - 3 Compute the mean of the \bar{f} over the n simulated values
 - rmk (central limit theorem): \bar{f} converges in proba towards $E[f(x)]$
 - 4 compute the approximated values as $20 \times \bar{f}$

Result

- uniform : $x \leftarrow \text{runif}(1000000000, -10, 10)$
- compute f: $f \leftarrow (1/(\sqrt{2 \cdot 3.14})) \cdot \exp(-0.5 \cdot x^2)$
- compute mean of f: $f_bar \leftarrow \text{mean}(f)$
- compute approximated value: $\text{approx} \leftarrow f_bar \cdot 20$
- show the approximated value: `approx`

Exercise

Provide an approximation for the quantity π

- hint: leverage on (i) the area of a circle $x^2 + y^2 \leq 1$, (ii) the area of a square centered at $(0,0)$ and whose side is 2, and (ii) random draws of x and y from $U[-1, 1]$.

Result

- simulate random values on the square whose side is 2: `pid <- data.frame(x=runif(100000000,-1,1),y=runif(100000000,-1,1))`
- identify random values that belong to the area of a circle
 $x^2 + y^2 \leq 1$
 - `pid$area <- (pid$x)**2+(pid$y)**2`
 - `pid$dum <- ifelse(pid$area <= 1, 1, 0)`
- compute the approximation: `pi_approx <- 4*mean(pid$dum)`
 - note that “4” corresponds to the area of the square whose side is 2
- `pi_approx`

Next: STATA !