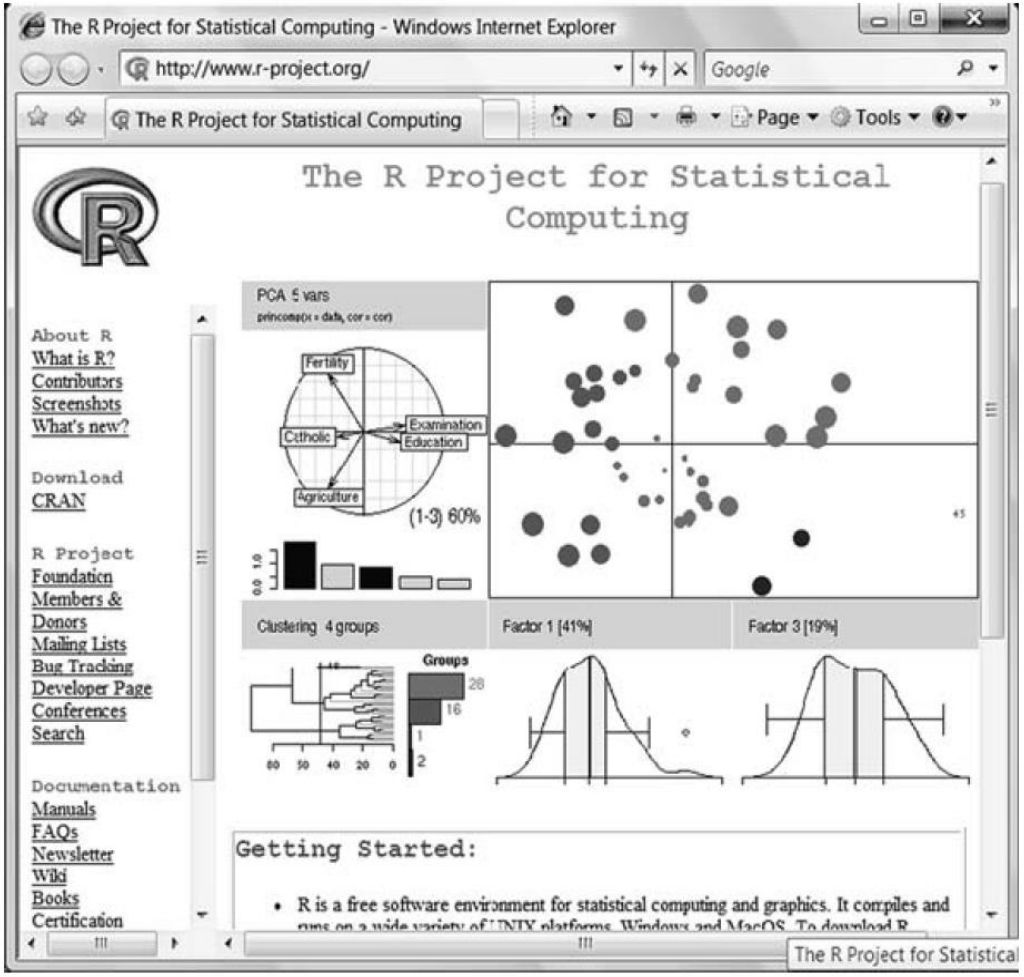
Introduction à R

L3PRO 2017

Evolution de S (années 70), souple, open source, complet, multi-plateforme, énorme communauté…



www.r-project.org

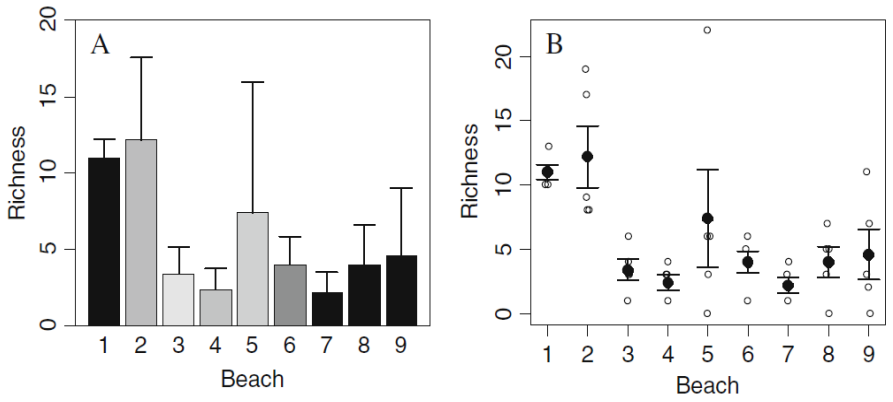


Fig. 7.3 **A**: Bar chart showing the benthic data. Mean values are represented by the *bars* with a *vertical line* showing standard deviations. The colours were changed to greyscale during the printing process. **B**: Strip chart for the raw data. The mean value per beach is plotted as a *filled dot*, and the *lines* represent the mean +/– the standard error

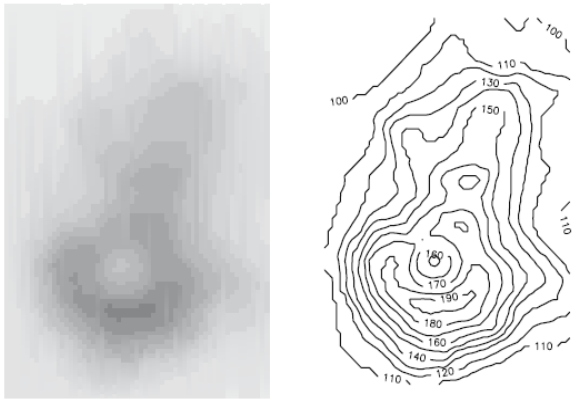


Fig. 5.1. Image plot and contour plot representations of Maunga Whau from the standard R `volcano` data set, for the same elevation class intervals (rotated to put north at the top)

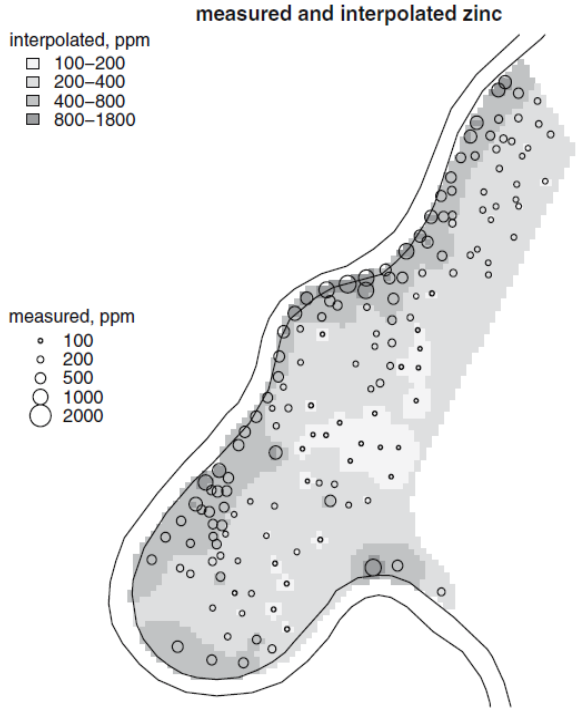
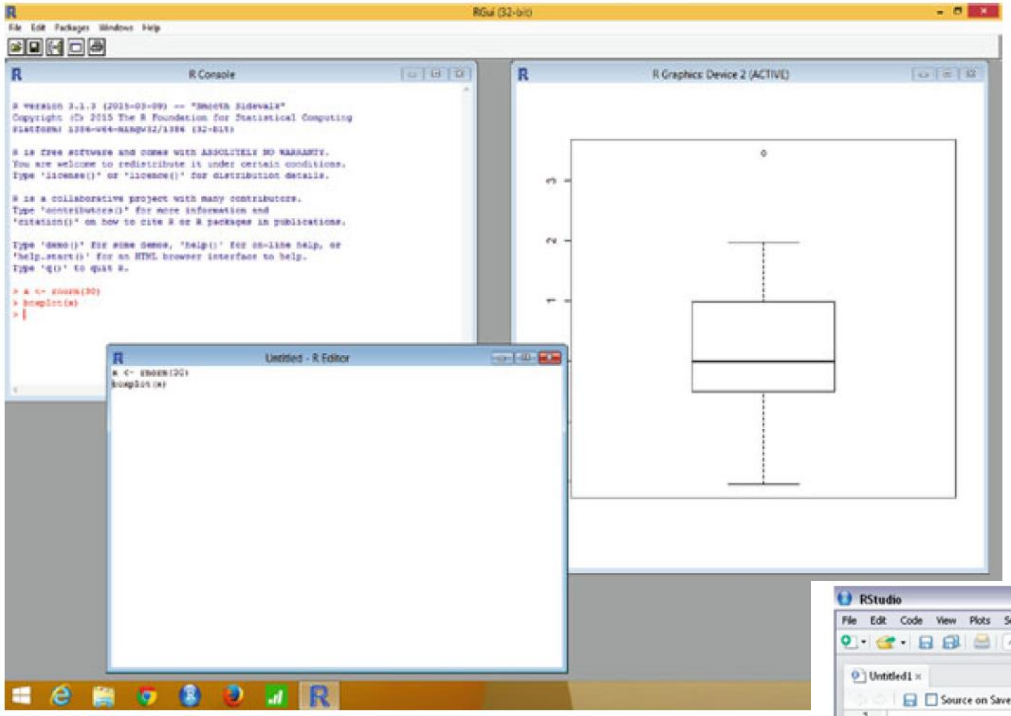
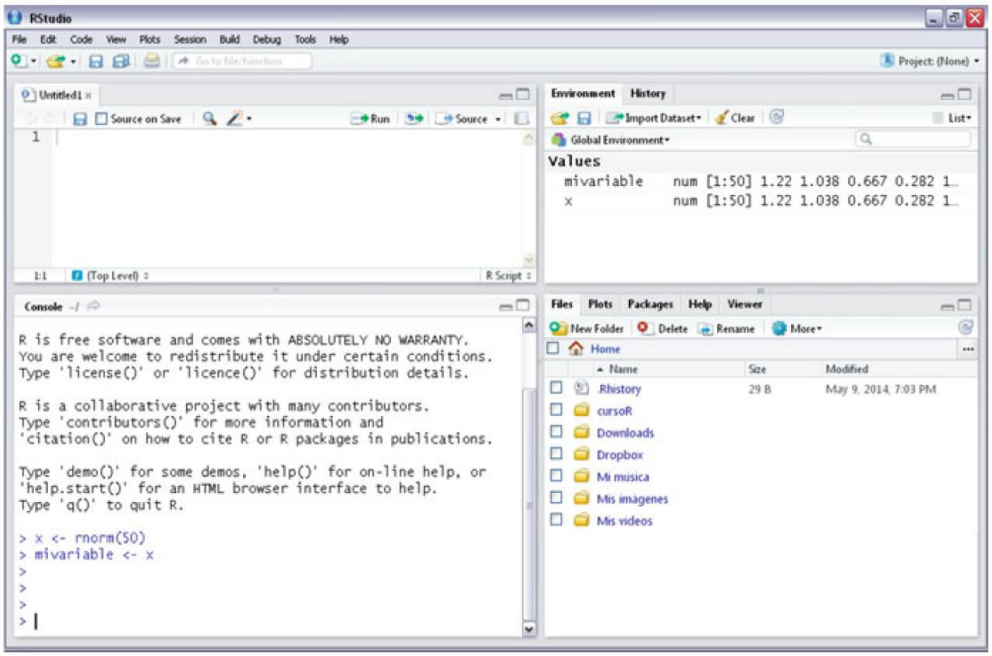


Fig. 3.8. Sample data points for zinc (ppm) plotted over an interpolated image, with symbol area proportional to measured concentration

Pourquoi R+Rstudio





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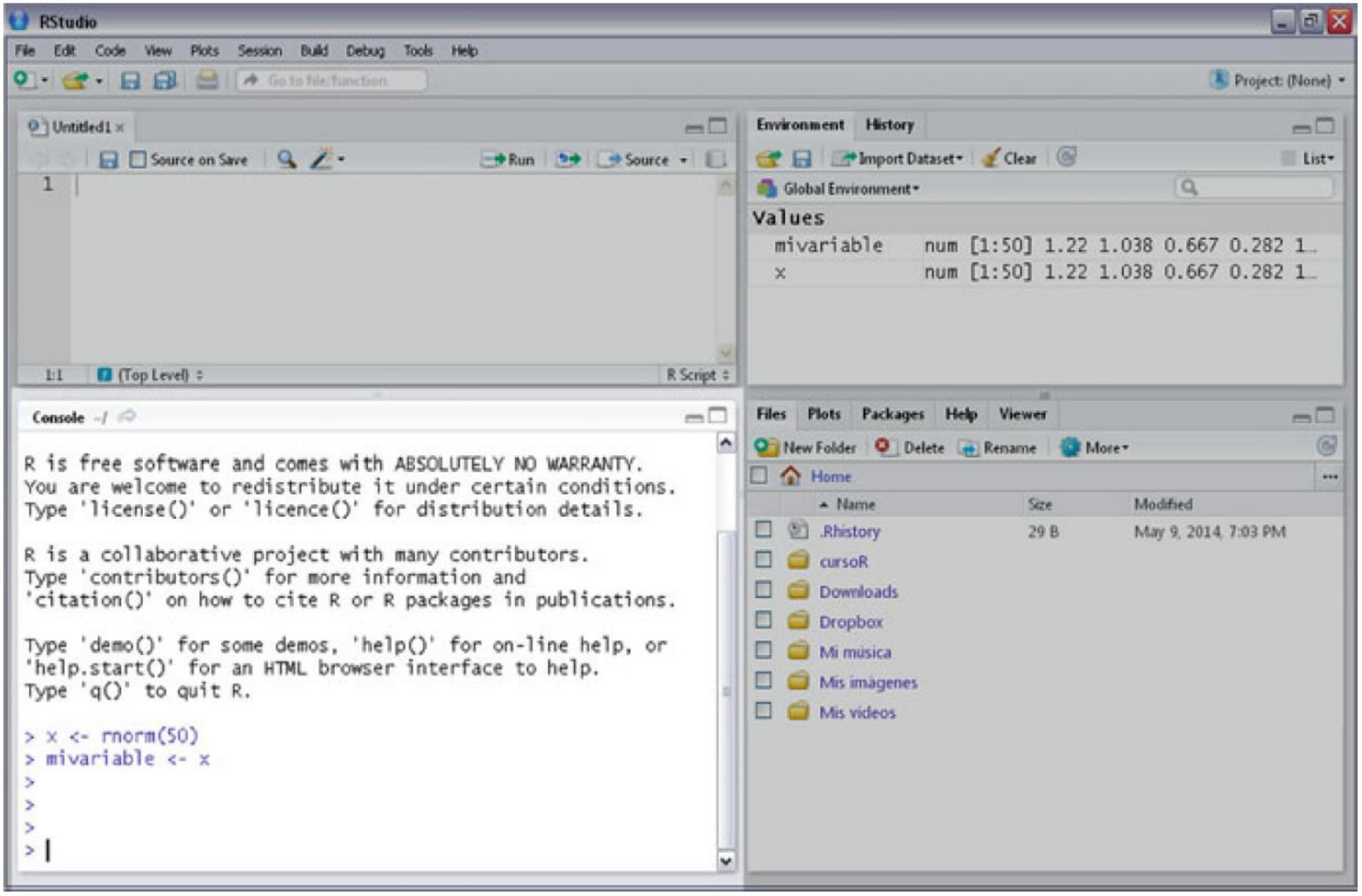
La console


RStudio
File Edit Code View Plots Session Build Debug Tools Help
Project: (None)
Untitled1
Source on Save
Run
Source
1
1:1
(Top Level)
R Script
Environment History
Import Dataset
Clear
List
Global Environment
Values
mivariable num [1:50] 1.22 1.038 0.667 0.282 1..
x num [1:50] 1.22 1.038 0.667 0.282 1..
Console ~/
R is free software and comes with ABSOLUTELY NO WARRANTY.
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R is a collaborative project with many contributors.
Type 'contributors()' for more information and
'citation()' on how to cite R or R packages in publications.
Type 'demo()' for some demos, 'help()' for on-line help, or
'help.start()' for an HTML browser interface to help.
Type 'q()' to quit R.
> x <- rnorm(50)
> mivariable <- x
Files Plots Packages Help Viewer
New Folder
Delete
Rename
More
Home
Name
Size
Modified
.Rhistory 29 B May 9, 2014, 7:03 PM
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Downloads
Dropbox
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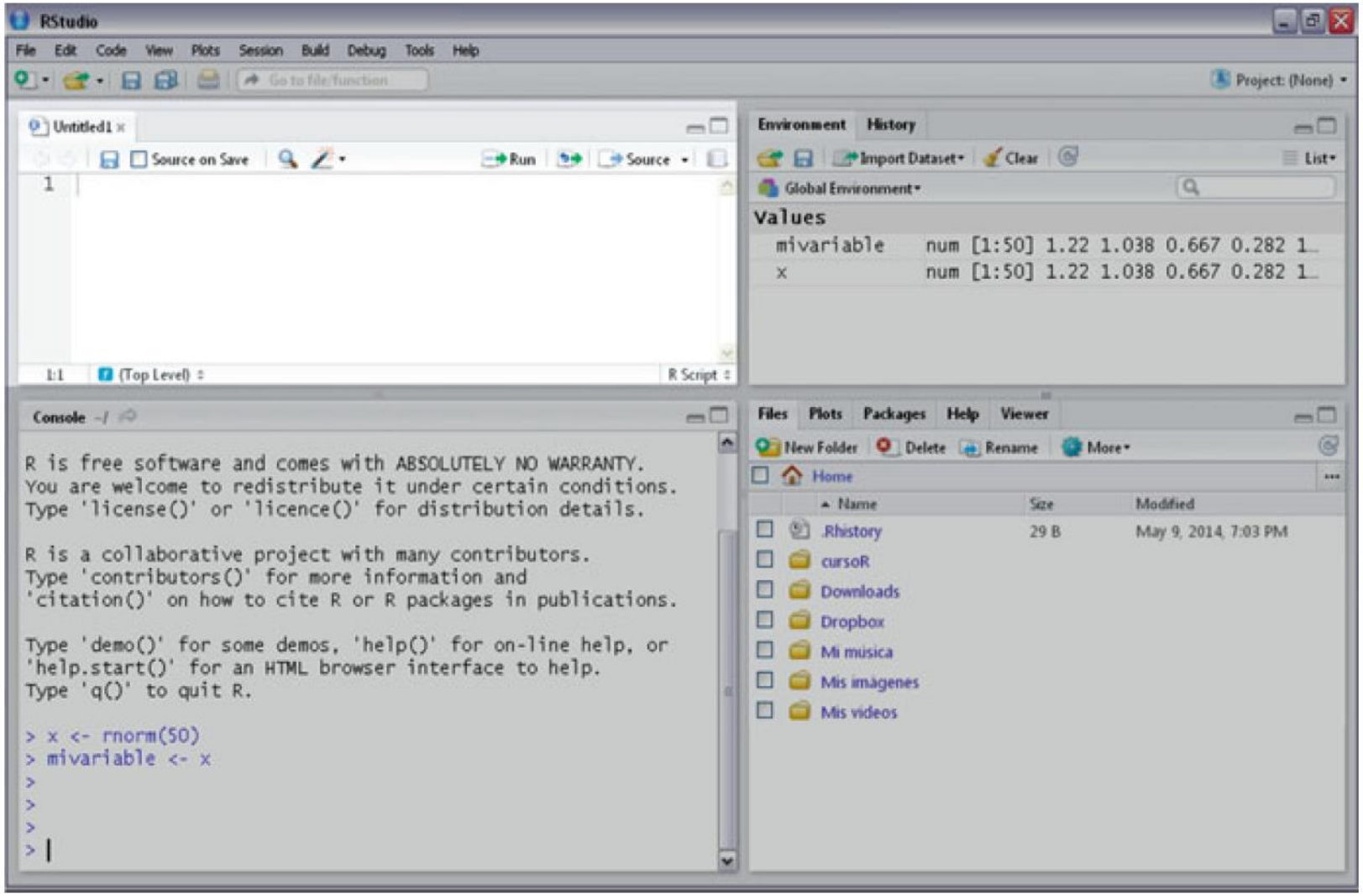
Mis imágenes
Mis videos

Le script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File Edit Code View Plots Session Build Debug Tools Help
Project: (None)
Untitled1
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1
1:1
(Top Level)
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.Rhistory 29 B May 9, 2014, 7:03 PM
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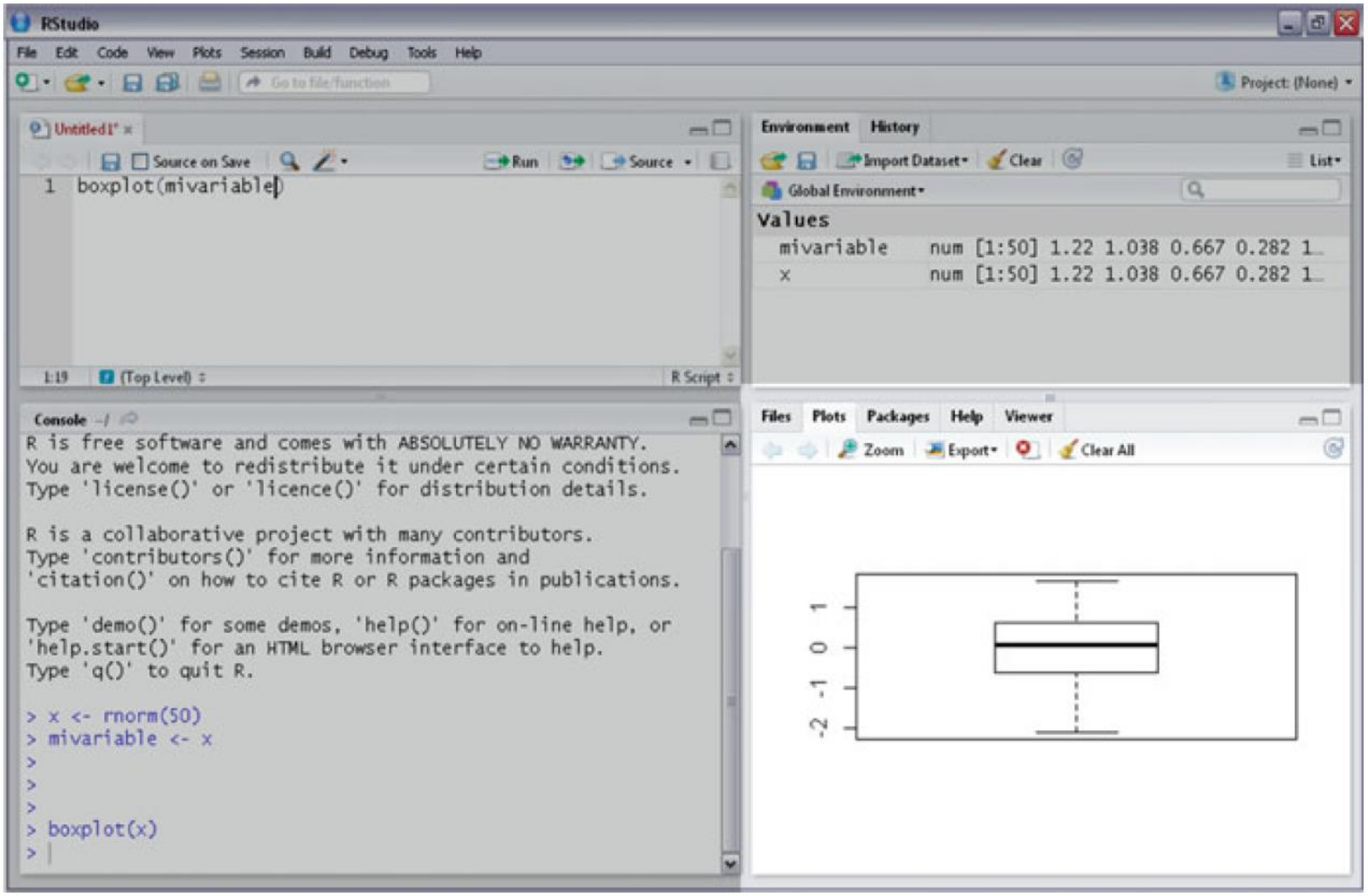
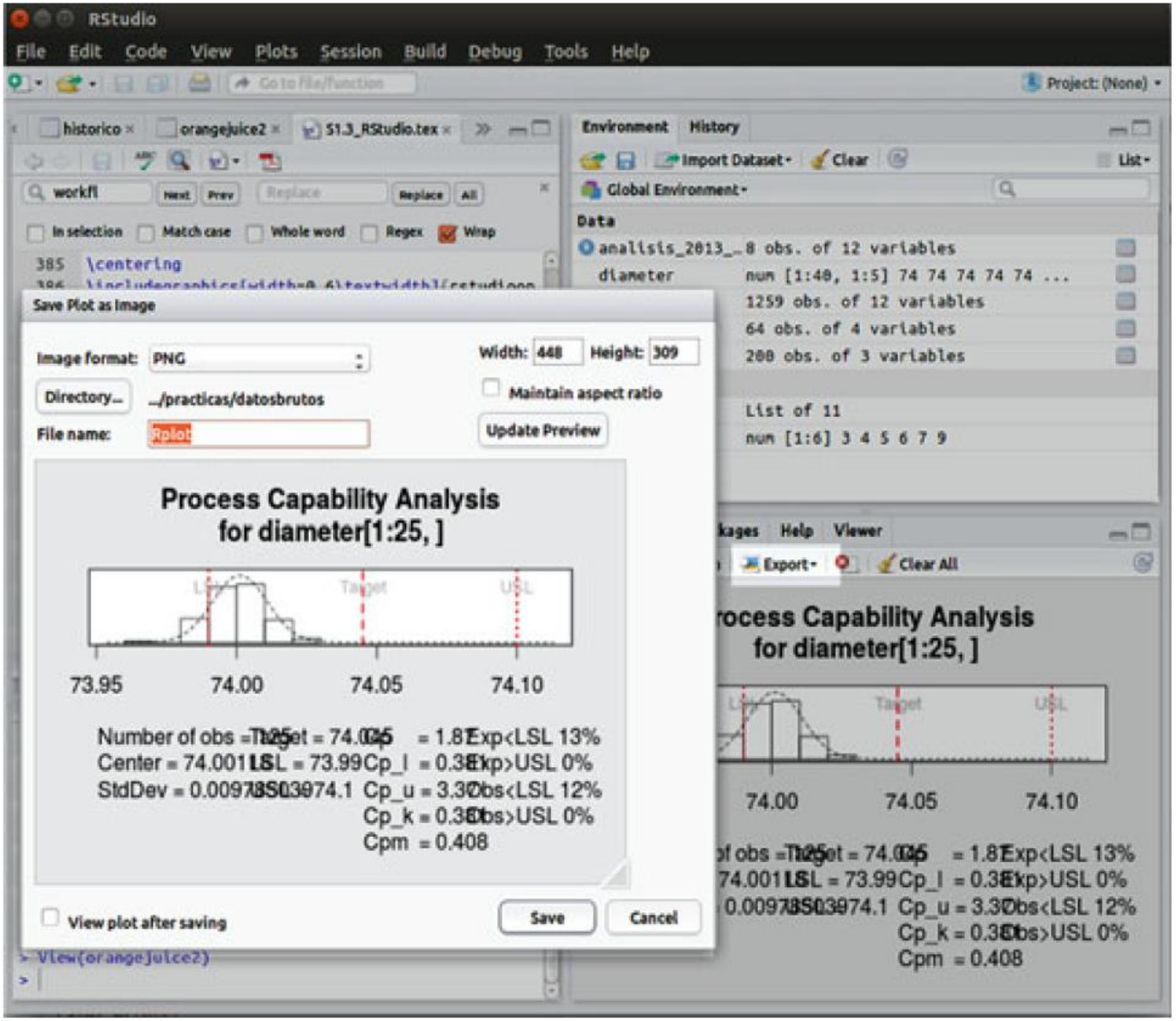
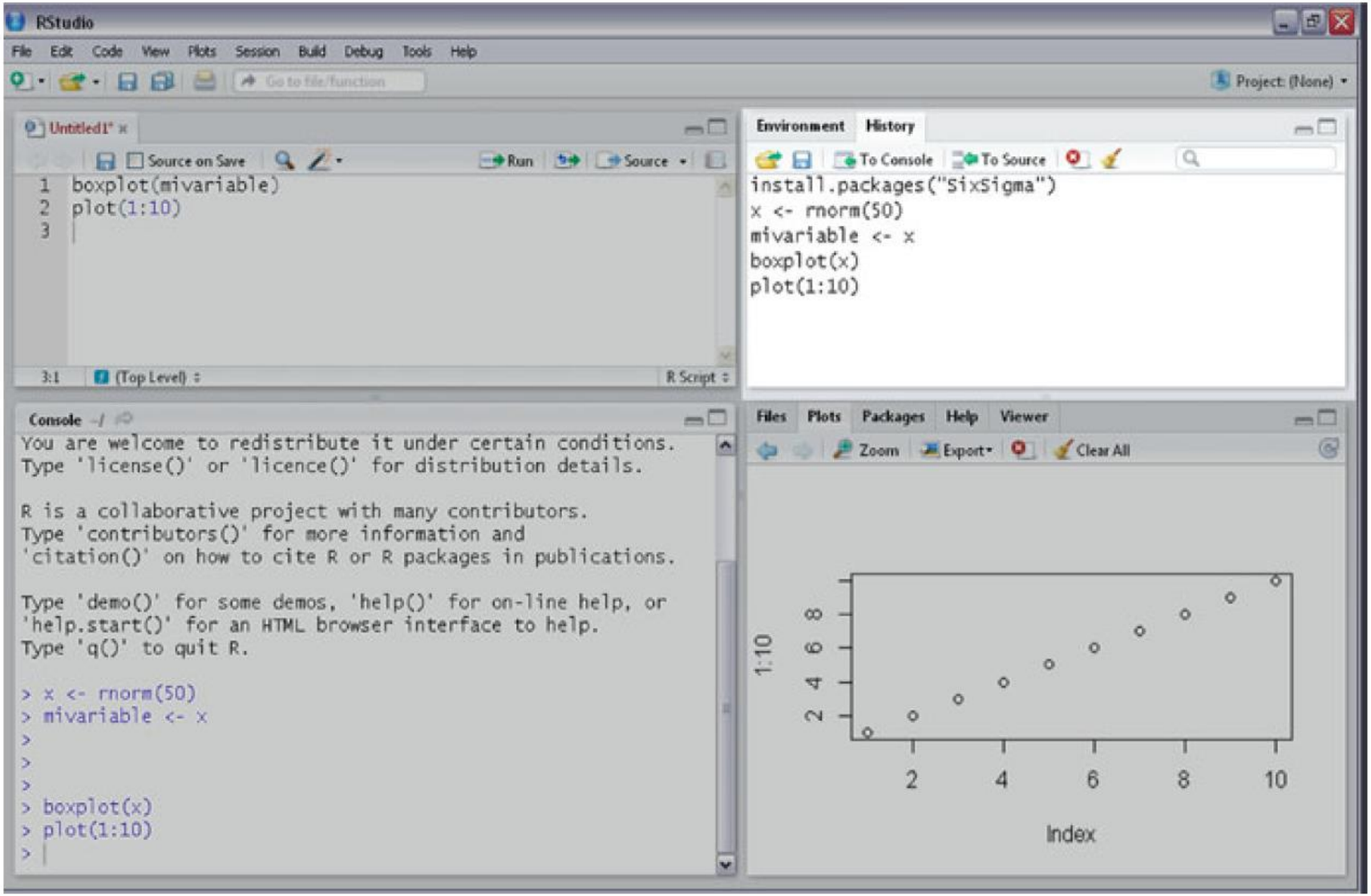
La fenêtre Plot



Sauver une image




RStudio
File Edit Code View Plots Session Build Debug Tools Help
Project: (None)
Untitled1*
Source on Save
Run
Source
1 boxplot(mivariable)
2 plot(1:10)
3
3:1 (Top Level)
R Script
Environment History
To Console
To Source
install.packages("SixSigma")
x <- rnorm(50)
mivariable <- x
boxplot(x)
plot(1:10)
Console
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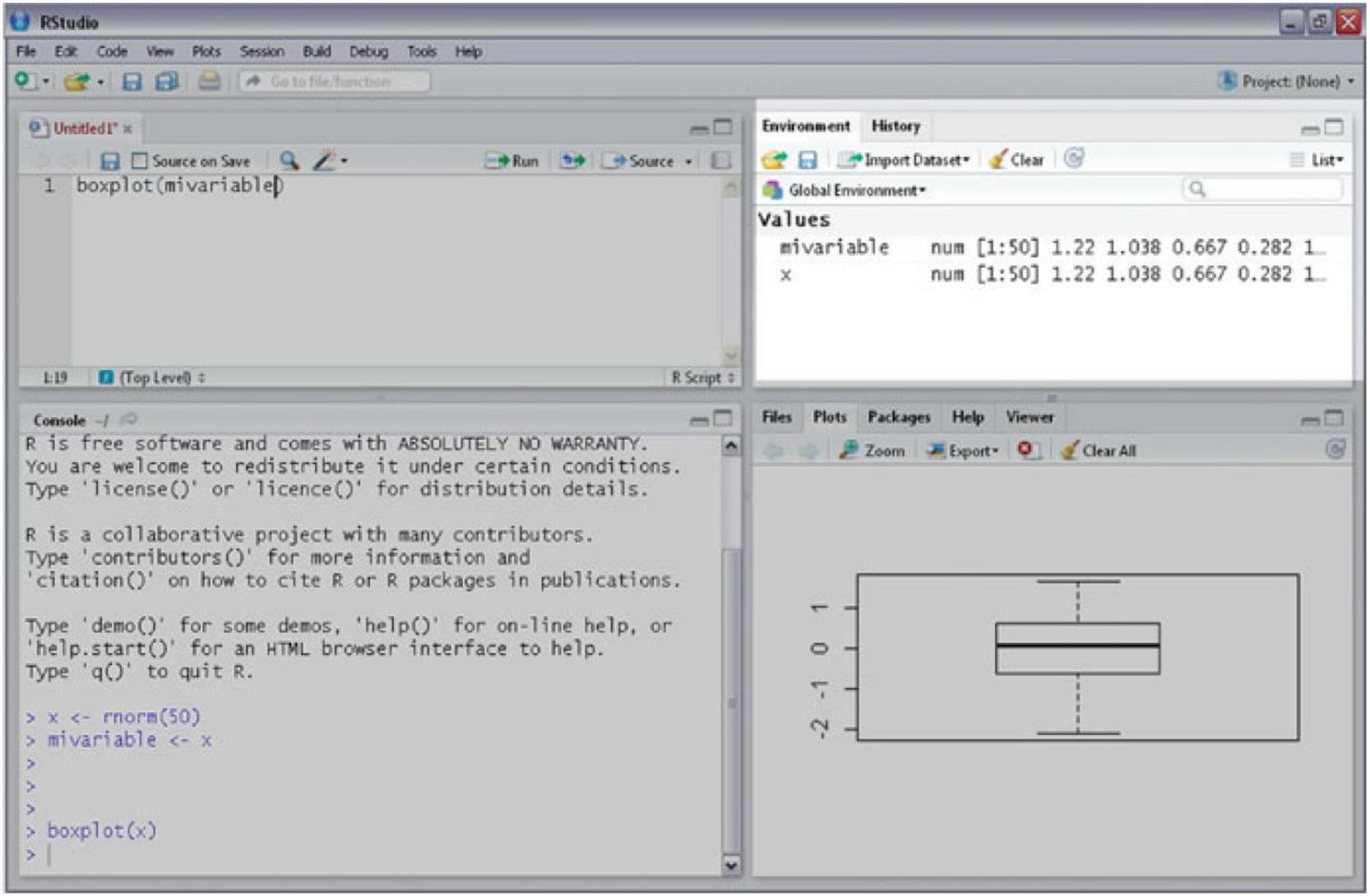
Type 'q()' to quit R.
> x <- rnorm(50)
> mivariable <- x
> boxplot(x)
> plot(1:10)
Files Plots Packages Help Viewer
Zoom
Export
Clear All
1:10
2 4 6 8
2 4 6 8 10
Index


RStudio
File Edit Code View Plots Session Build Debug Tools Help
Project: (None)
Untitled1*
Source on Save
Run
Source
1 boxplot(mivariable)
1:19
(Top Level)
R Script
Environment
History
Import Dataset
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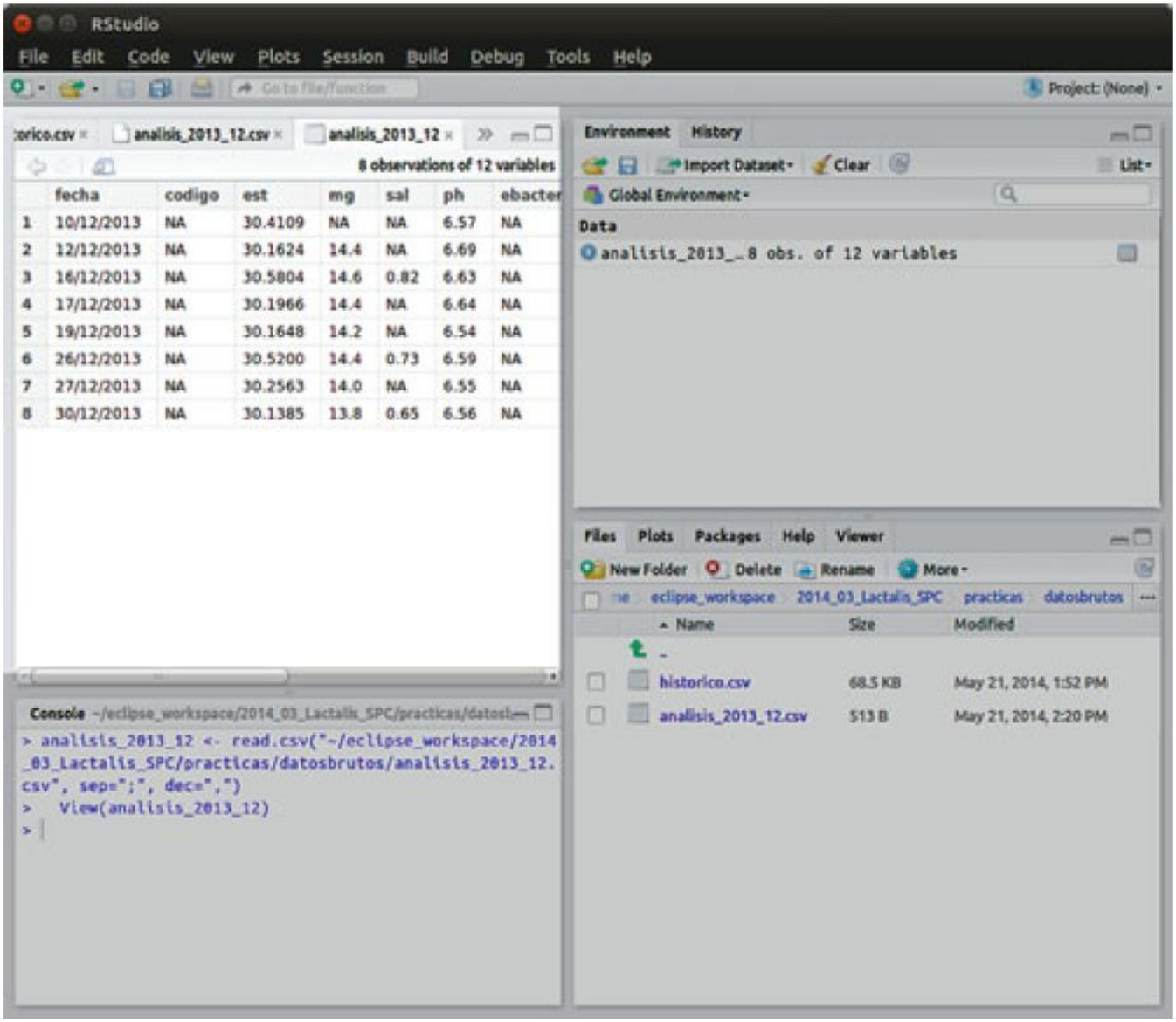
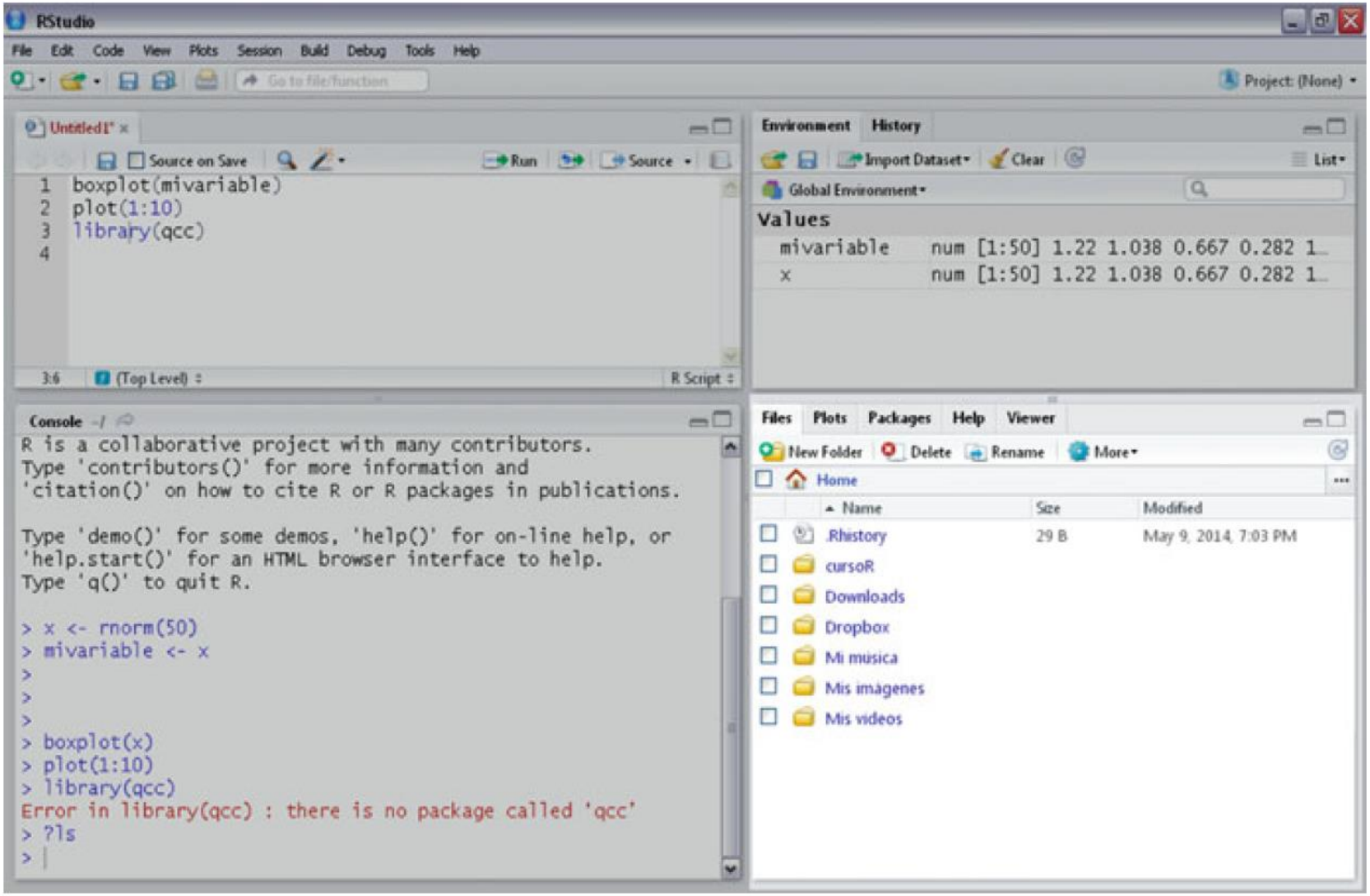
> x <- rnorm(50)
> mivariable <- x
> boxplot(x)
Files
Plots
Packages
Help
Viewer
Zoom
Export
Clear All
-2 -1 0 1

Les variables



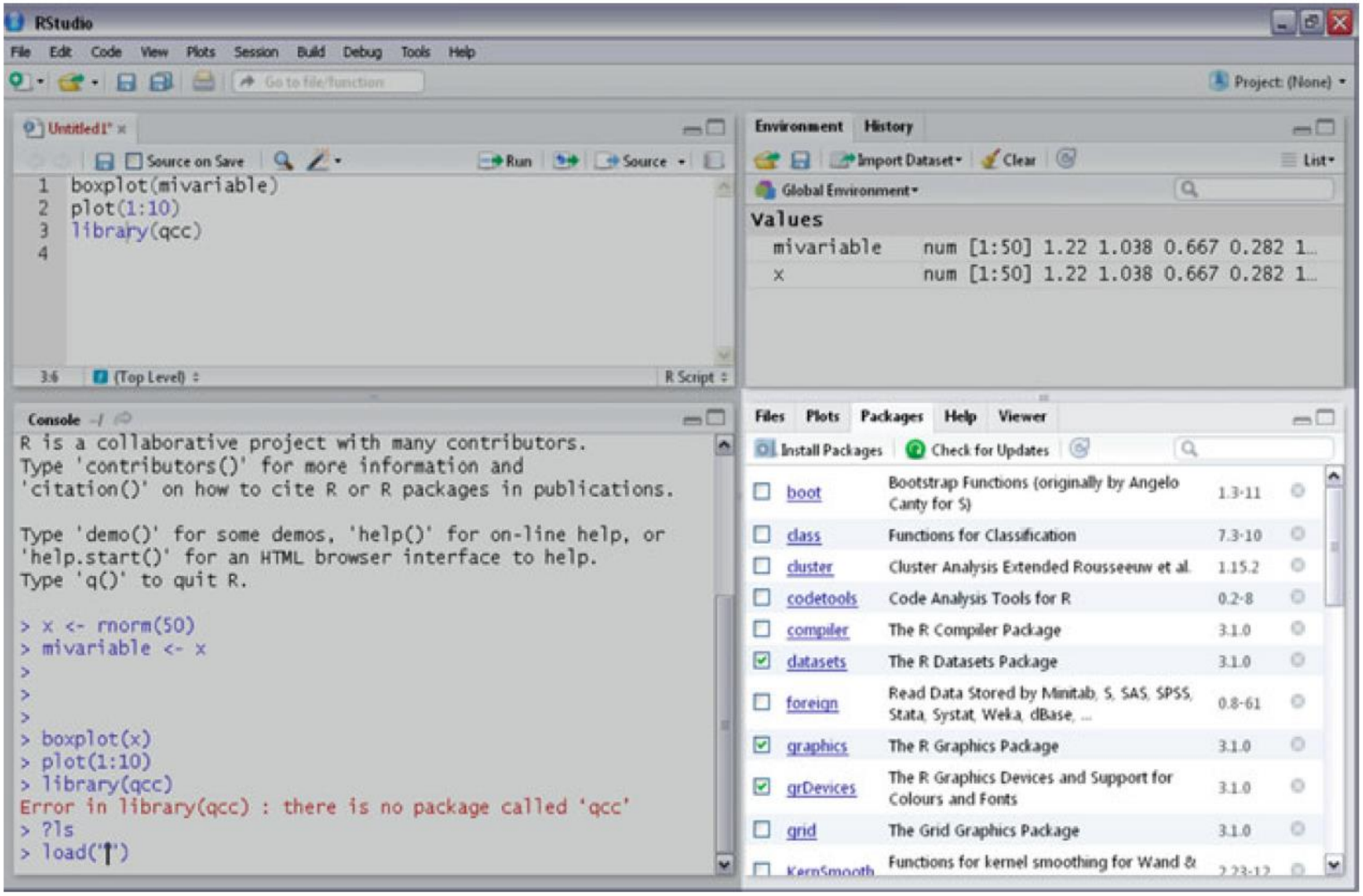
Le répertoire de travail



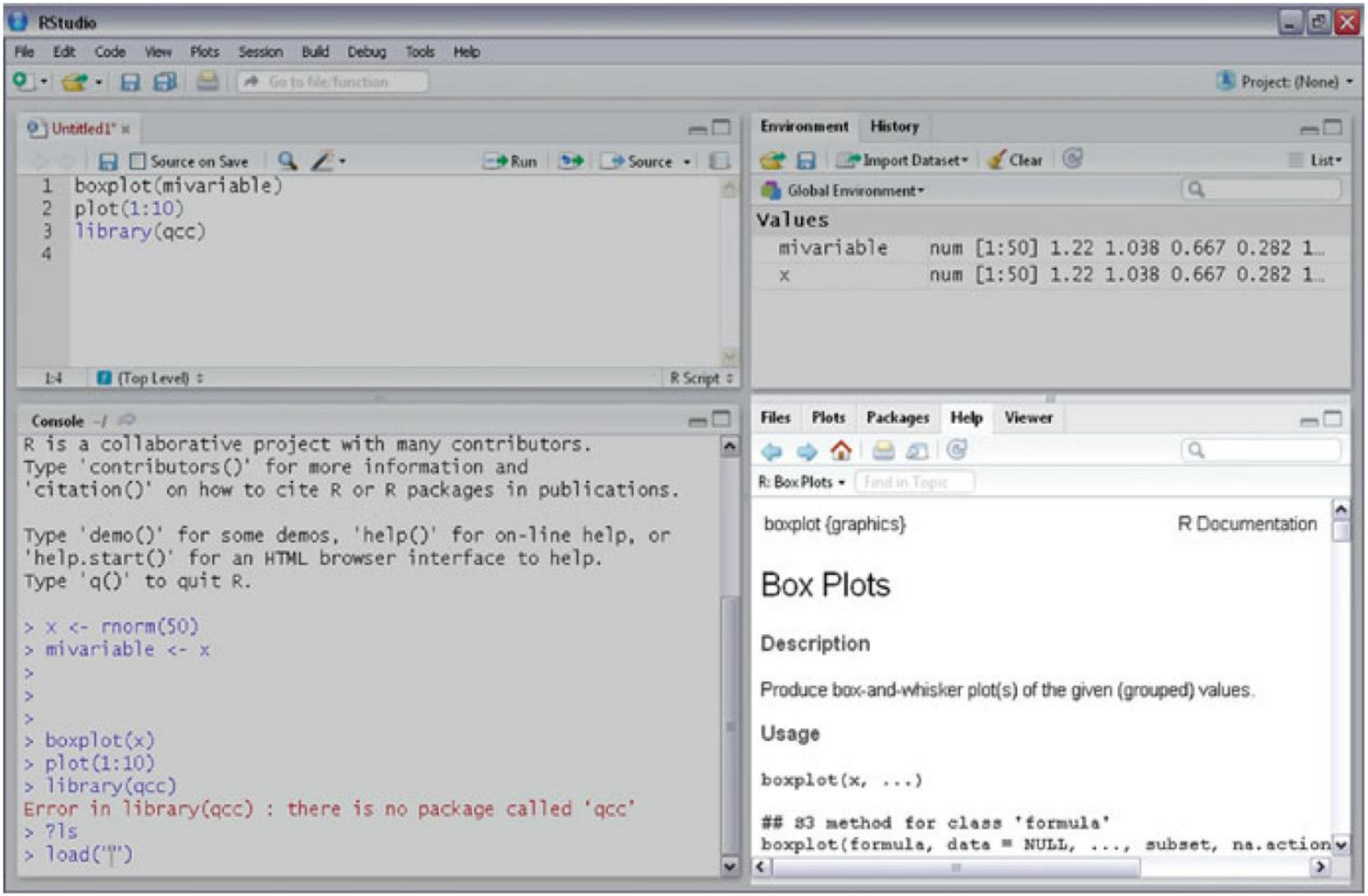

```
## Correct:
setwd("C:/myscripts")
setwd("C:\\myscripts")

## Incorrect:
setwd("C:\myscripts")
```

Les packages



Les packages



Les expressions – quelques exemples

```
> 1
[1] 1

> 1+1
[1] 2

> 1+
+ |

> 5>6
[1] FALSE

> log(pi)
[1] 1.14473

> pi
[1] 3.141593
```

```
> str(log)
function (x, base = exp(1))
> exp(1)
[1] 2.718282

> log(1000,base=10)
[1] 3

> log()
Error: argument "x" is missing, with
no default

> ?seq
> help("seq")

> rnorm(20)
 [1] -0.3373624346  0.8834426105
 [3] -0.0005967386 -0.2083691171
 [5] -0.2893600292  1.5701507665
 [7] -0.0290318028  0.3608562449
 [9] -0.6914431117  0.3347585358
[11] -2.1484833764  1.2388822292
[13] -1.4536453004  1.8971513103
[15]  0.6304689200 -1.3084480885
[17]  0.6844385933  1.7017605077
[19]  1.5247046231  2.2565003356
```

Les expressions – quelques exemples

```
> x <- 1 # Affectation.
> x # Affichage.
[1] 1

> 2 -> x
> x
[1] 2

> 2*8*10+exp(1)
[1] 162.7183


> 2*8*
+ 10+
+ exp(1)
[1] 162.7183
```

```
> # Il est possible d'écrire 2 instructions
> # sur la même ligne grâce au signe ;
> Mon.Poids <- 75 ; Ma.Taille <- 1.90
> Mon.IMC <- Mon.Poids/Ma.Taille^2
> Mon.IMC
[1] 20.77562

> source("http://www.biostatisticien.eu/springeR/IMC.R",
+        encoding="utf8")
> affiche.IMC(Mon.IMC)
```

Type numerique (*numeric*)

Il y a deux types numeriques : les entiers (integer) et les reels (real ou double).

```
> a <- 1
> b <- 3.4
> c <- as.integer(a)
> typeof(c)
[1] "integer"
```

Type booleen ou logique (*logical*)

```
> b>a
[1] TRUE
> a==b
[1] FALSE
> is.numeric(a)
[1] TRUE
> is.integer(a)
[1] FALSE
> x <- TRUE
> is.logical(x)
[1] TRUE
```

```
> TRUE + T + FALSE*F + T*FALSE + F
[1] 2
```

Données manquantes (*NA*)

```
> x <- c(3,NA,6)
> is.na(x)
[1] FALSE  TRUE FALSE
> mean(x) # Tentative de calcul de la moyenne de x.
[1] NA


> mean(x,na.rm=TRUE)
[1] 4.5
```

Type chaînes de caracteres (*character*)

```
> a <- "R est mon ami"
> is.character(a)
[1] TRUE
```

Les vecteurs (*vector*)

Cette structure de données est la plus simple. Elle représente une suite de données de même type.

```
> c(3,1,7)
[1] 3 1 7
> seq(from=0,to=1,by=0.1)
 [1] 0.0 0.1 0.2 0.3 0.4 0.5 0.6 0.7 0.8 0.9 1.0
> seq(from=0,to=20,length=5)
[1]  0  5 10 15 20
> vec <- 2:36
> vec
 [1]  2  3  4  5  6  7  8  9 10 11 12 13 14 15 16 17 18 19 20 21 22
 23 24 25 26 27 28 29 30 31 32 33 34 35 36
```

```
> x <- 1:3
> x
[1] 1 2 3
```

Les matrices (*matrix*) et les tableaux (*arrays*)

Généralisation de la notion de vecteur

```
> X <- matrix(1:12,nrow=4,ncol=3,byrow=TRUE)
> X
     [,1] [,2] [,3]
[1,]    1    2    3
[2,]    4    5    6
[3,]    7    8    9
[4,]   10   11   12
```

```
> Y <- matrix(1:12,nrow=4,ncol=3,byrow=FALSE)
> Y
     [,1] [,2] [,3]
[1,]    1    5    9
[2,]    2    6   10
[3,]    3    7   11
[4,]    4    8   12
```

```
> X <- array(1:12,dim=c(2,2,3))
> X
, , 1

     [,1] [,2]
[1,]    1    3
[2,]    2    4

, , 2

     [,1] [,2]
[1,]    5    7
[2,]    6    8

, , 3

     [,1] [,2]
[1,]    9   11
[2,]   10   12
```

Les listes (*list*)

les listes permettent de regrouper dans une même structure des données de types différents

```
> A <- list(TRUE,-1:3,matrix(1:4,nrow=2),c(1,3),"Une chaîne de caractères")
> A
[[1]]
[1] TRUE

[[2]]
[1] -1  0  1  2  3

[[3]]
     [,1] [,2]
[1,]    1    3
[2,]    2    4

[[4]]
[1] 1 3

[[5]]
[1] "Une chaîne de caractères"
```

Le tableau individus / variables (*data.frame*)

Chaque colonne représente une variable particulière dont tous les éléments sont du même type

```
> IMC <- data.frame(Sexe=c("H","F","H","F","H","F"),
+ Taille=c(1.83,1.76,1.82,1.60,1.90,1.66),
+ Poids=c(67,58,66,48,75,55),
+ row.names=c("Rémy","Lol","Pierre","Domi","Ben","Cécile"))
> IMC
       Sexe Taille Poids
Rémy      H   1.83    67
Lol       F   1.76    58
Pierre    H   1.82    66
Domi      F   1.60    48
Ben       H   1.90    75
Cécile    F   1.66    55
```

Les facteurs (*factor*) et les variables ordinales (*ordered*)

```
> x <- factor(c("bleu","vert","bleu","rouge",
+               "bleu","vert","vert"))
> x
[1] bleu  vert  bleu  rouge bleu  vert  vert
Levels: bleu rouge vert

> levels(x)
[1] "bleu"  "rouge" "vert"
```

```
> z <- ordered(c("Petit","Grand","Moyen","Grand","Moyen",
+ "Petit","Petit"),levels=c("Petit","Moyen","Grand"))
> z
[1] Petit Grand Moyen Grand Moyen Petit Petit
Levels: Petit < Moyen < Grand
```

Les erreurs à ne pas commettre.

Attention à la syntaxe!

Fermer les parenthèses

Chaînes de caractères avec « »

Manque de parenthèses dans les fonctions

Mauvais arguments dans les fonctions

Longueurs incompatibles

Mauvaise syntaxe

- Les arguments doivent être séparés par une virgule
- Les conditions sont entre parenthèses
- Le point est le séparateur numérique
- Expressions sur une ligne ou séparées par un point virgule

Utilisons le jeu de données cars du package `dataset`

```
> str(cars)
'data.frame':   50 obs. of  2 variables:
 $ speed: num  4 4 7 7 8 9 10 10 10 11 ...
 $ dist : num  2 10 4 22 16 10 18 26 34 17 ...
```

```
> is.data.frame(cars)
[1] TRUE
```

```
> head(cars)
  speed dist
1     4    2
2     4   10
3     7    4
4     7   22
5     8   16
6     9   10
```

```
> cars2 <- cars
> cars2$dist * 0.3048
 [1]  0.6096  3.0480  1.2192  6.7056  4.8768  3.0480  5.4864  7.9248 10.3632
[10]  5.1816  8.5344  4.2672  6.0960  7.3152  8.5344  7.9248 10.3632 10.3632
[19] 14.0208  7.9248 10.9728 18.2880 24.3840  6.0960  7.9248 16.4592  9.7536
[28] 12.1920  9.7536 12.1920 15.2400 12.8016 17.0688 23.1648 25.6032 10.9728
[37] 14.0208 20.7264  9.7536 14.6304 15.8496 17.0688 19.5072 20.1168 16.4592
[46] 21.3360 28.0416 28.3464 36.5760 25.9080
```

cars2$dist équivalent à
— cars2[["dist"]]
— cars2[, "dist"]
— cars2[,2]

```
> cars2[2,3]
[1] 3.048
> cars2$dist_m[2]
[1] 3.048

> cars2[2:4,]
  speed dist dist_m
2     4   10 3.0480
3     7    4 1.2192
4     7   22 6.7056

> dim(cars2)
[1] 50  3
```

```
> dist_m=cars2$dist * 0.3048
> cars2=cbind(cars2,dist_m)
> head(cars2)
  speed dist dist_m
1     4    2 0.6096
2     4   10 3.0480
3     7    4 1.2192
4     7   22 6.7056
5     8   16 4.8768
6     9   10 3.0480

> dim(cars2)
[1] 50  4

> cars2=rbind(cars2,cars2)
> dim(cars2)
[1] 100   4
```

```
> cars2$speed_km <- ifelse(cars$speed < 10, NA, cars$speed*1.60934)
> head(cars2, n = 10)
   speed dist speed_km
1      4    2       NA
2      4   10       NA
3      7    4       NA
4      7   22       NA
5      8   16       NA
6      9   10       NA
7     10   18 16.09340
8     10   26 16.09340
9     10   34 16.09340
10    11   17 17.70274
```

Les conditions

```
if (test_expression) {
 statement
}
```

```
x <- 5
if(x > 0){
     print("Positive number")
}
```

```
x <- -5
if(x > 0){
      print("Non-negative number")
} else {
      print("Negative number")
}
```

```
x <- 0
if (x < 0) {
    print("Negative number")
} else if (x > 0) {
    print("Positive number")
} else print("Zero")
```

Les boucles (for)

```
for (val in sequence)
{
     statement
}
```

```
x <- c(2,5,3,9,8,11,6)
count <- 0
for (val in x) {
     if(val %% 2 == 0)   count = count+1
}
print(count)
```

Les boucles (while)

```
while (test_expression)
{
    statement
}
```

```
i <- 1

while (i < 6) {
    print(i)
    i = i+1
}
```