Presentation of R and R Studio

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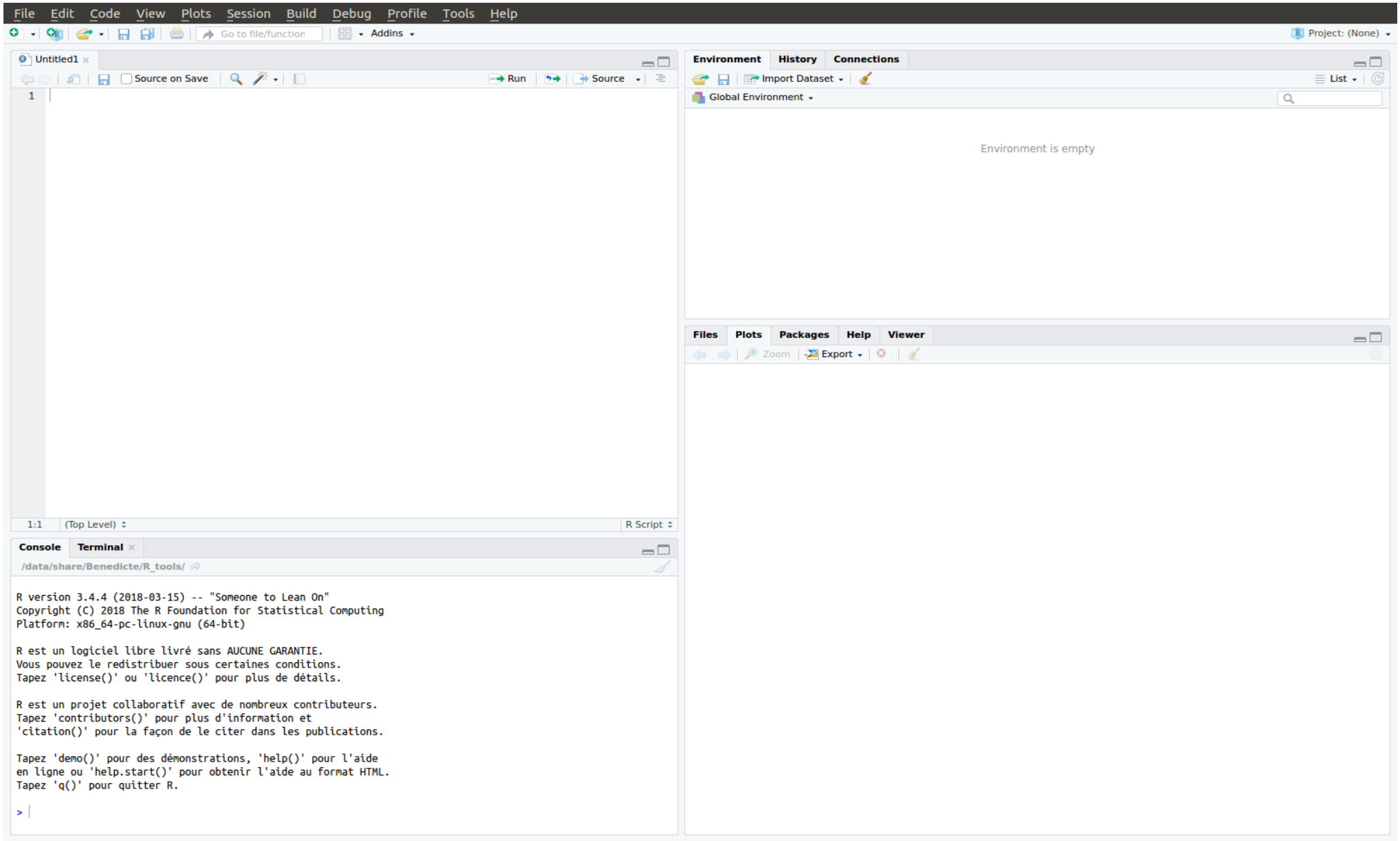
08/11/2018

What are R and R studio?

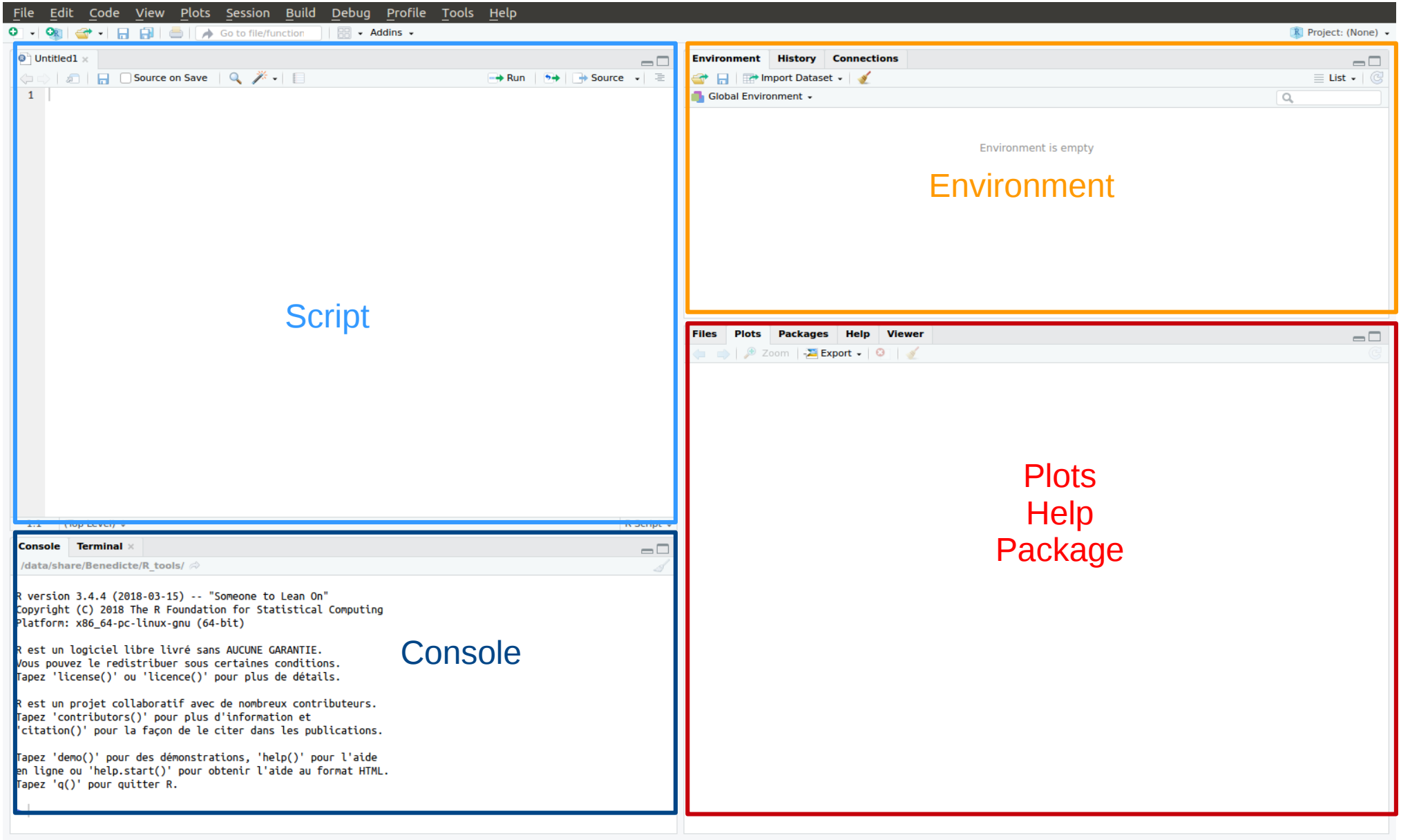
- R is a programming language created in 1993
- R and R studio are free, open source, software environment
- CRAN project: https://cran.r-project.org/

- R is useful to analyse data:
    - sorting complexe data frame
    - statisics
    - graphics
    - automatisation of repetitive tasks on datasets
    - and plenty of packages in function of your needs...

R studio


File Edit Code View Plots Session Build Debug Profile Tools Help
Go to file/function
Addins
Project: (None)
Untitled1
Source on Save
Run
Source
1
1:1
(Top Level)
R Script
Environment History Connections
Import Dataset
List
Global Environment
Environment is empty
Files Plots Packages Help Viewer
Zoom
Export
Console Terminal
/data/share/Benedicte/R_tools/
R version 3.4.4 (2018-03-15) -- "Someone to Lean On"
Copyright (C) 2018 The R Foundation for Statistical Computing
Platform: x86_64-pc-linux-gnu (64-bit)
R est un logiciel libre livré sans AUCUNE GARANTIE.
Vous pouvez le redistribuer sous certaines conditions.
Tapez 'license()' ou 'licence()' pour plus de détails.
R est un projet collaboratif avec de nombreux contributeurs.
Tapez 'contributors()' pour plus d'information et
'citation()' pour la façon de le citer dans les publications.
Tapez 'demo()' pour des démonstrations, 'help()' pour l'aide
en ligne ou 'help.start()' pour obtenir l'aide au format HTML.
Tapez 'q()' pour quitter R.
>

R studio


File Edit Code View Plots Session Build Debug Profile Tools Help
Go to file/function
Addins
Project: (None)
Untitled1
Source on Save
Run
Source
1
Script
Environment History Connections
Import Dataset
List
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Environment is empty
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Files Plots Packages Help Viewer
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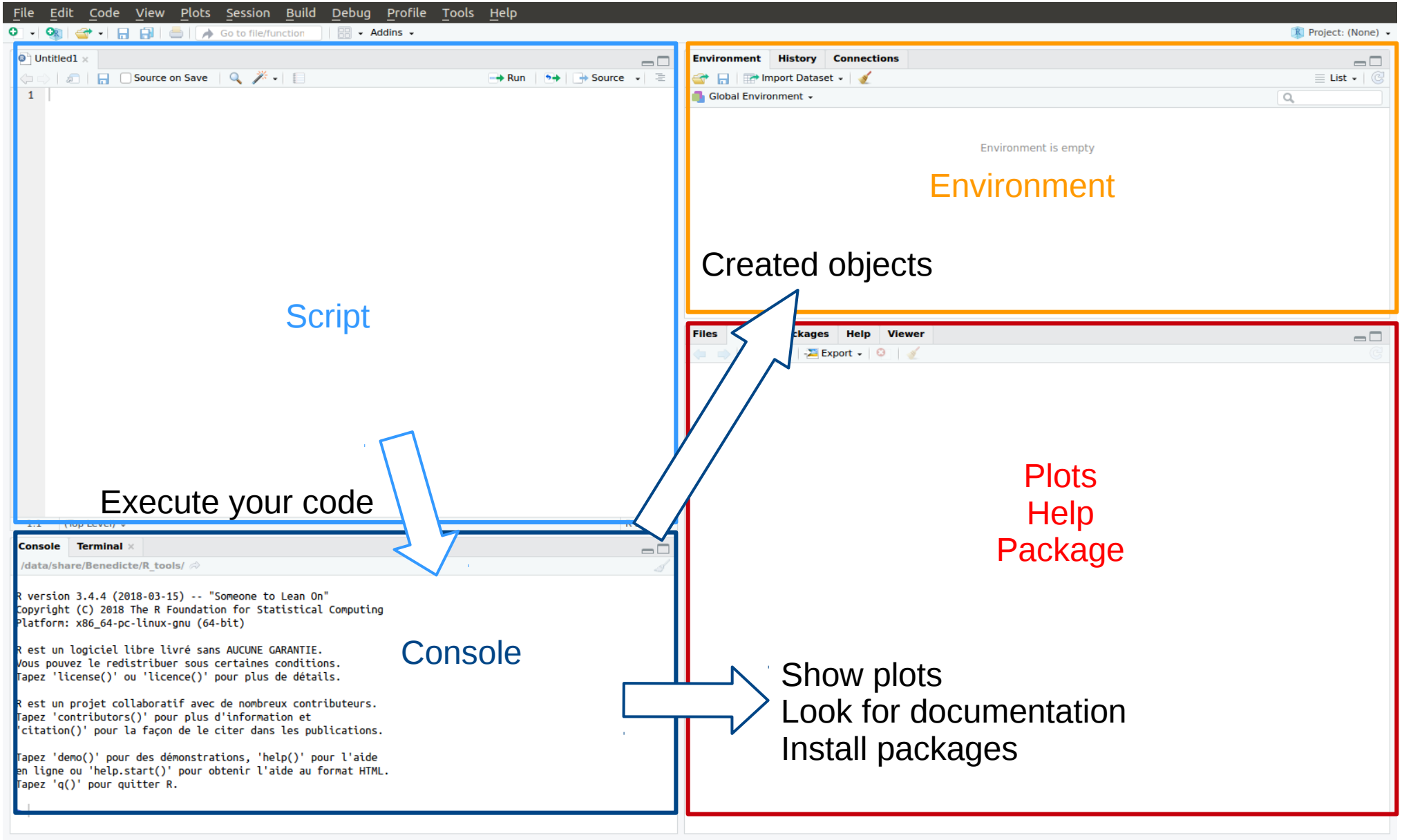
en ligne ou 'help.start()' pour obtenir l'aide au format HTML.
Tapez 'q()' pour quitter R.
Console

R studio


File Edit Code View Plots Session Build Debug Profile Tools Help
Untitled1
Source on Save
Run
Source
Environment History Connections
Import Dataset
Global Environment
List
Environment is empty
Environment
Created objects
Script
Execute your code
Console Terminal
/data/share/Benedicte/R_tools/
R version 3.4.4 (2018-03-15) -- "Someone to Lean On"
Copyright (C) 2018 The R Foundation for Statistical Computing
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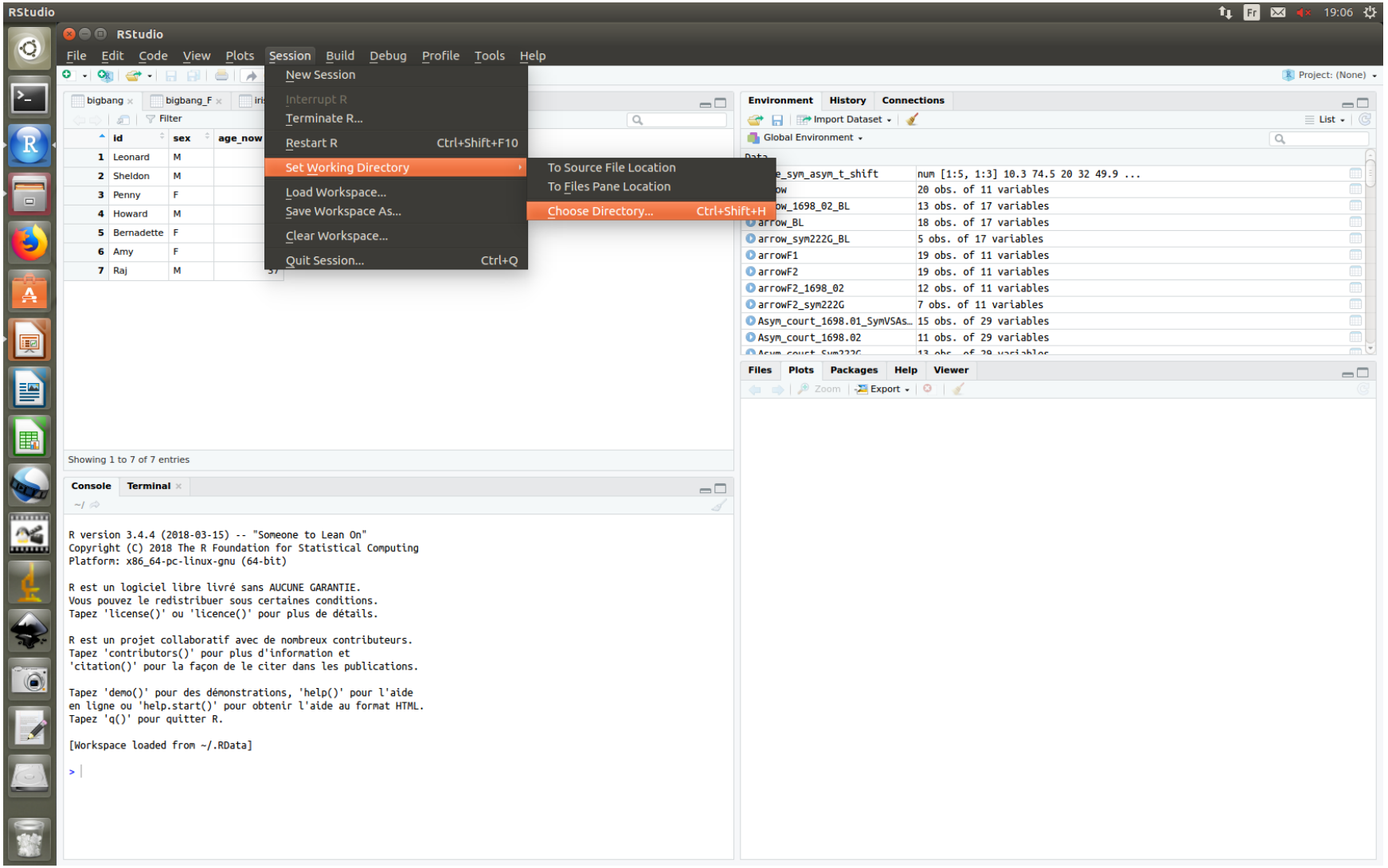
Tapez 'demo()' pour des démonstrations, 'help()' pour l'aide
en ligne ou 'help.start()' pour obtenir l'aide au format HTML.
Tapez 'q()' pour quitter R.
Console
Files
Help
Viewer
Export
Plots
Help
Package
Show plots
Look for documentation
Install packages

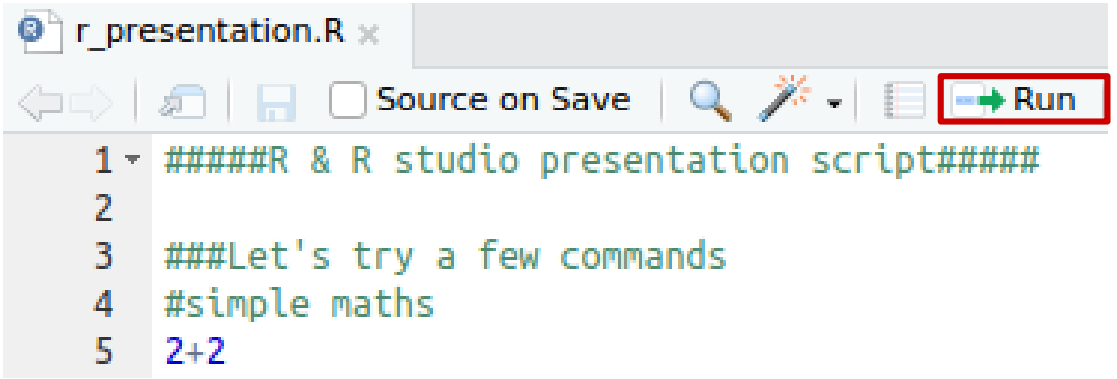
Before to start: let’s set our working directory



- Please, select your desktop as working directory and copy the iris.csv file in your desktop

How to write a command and run it?

- Let’s try a few commands:



To run a command:
- highlight it and click on run
or
- put your cursor in the line and press CTRL+enter

Lines that starts by # are not recognized as code but as **comments**

How to create an object?

- Let’s try a few commands:

r_presentation.R

Source on Save | Run

```
1  #####R & R studio presentation script#####
2
3  ###Let's try a few commands
4  #simple maths
5  2+2
6  #create objects
7  x=5
8  x
9  X=10
10 X
```

To run a command:
- highlight it and click on run
or
- put your cursor in the line and press CTRL+enter

If you reassign a value to an existing object, the previous one is removed

What can we do with the created objects?

- Let’s try a few commands:

r_presentation.R

Source on Save | Run

```
1  #####R & R studio presentation script#####
2
3  ###Let's try a few commands
4  #simple maths
5  2+2
6  #create objects
7  x=5
8  x
9  X=10
10 X
11 #logical test
12 x==X
13 x!=X
14 x<X
15 x>X
16 #simple maths with object
17 x+X
18 y=x+X
19 y
```

To run a command:
- highlight it and click on run
or
- put your cursor in the line and press CTRL+enter

R is case sensitive !
x ≠ X

A very important concept in R: vectors

- A vector is an object that contain a serie of variables of a same type, in a precise order
- c() is used to create vectors

object	modes	several modes possible in the same object?
vector	numeric, character, complex *or* logical	No
factor	numeric *or* character	No
array	numeric, character, complex *or* logical	No
matrix	numeric, character, complex *or* logical	No
data frame	numeric, character, complex *or* logical	Yes
ts	numeric, character, complex *or* logical	No
list	numeric, character, complex, logical, function, expression, ...	Yes

E.Paradis, R for beginners, 2005

Let’s create vectors and data frame: example with BigBang Theory

```
22  ###Different kinds of objects and data
23  #create vector
24  id=c("Leonard","Sheldon","Penny","Howard","Bernadette","Amy","Raj")
25  id
26  sex=c("M","M","F","M","F","F","M")
27  age_now=c(45,43,32,37,38,42,37)
28  #create dataframe
29  bigbang=data.frame(id,sex,age_now)
```

characters are written as ’’character’’ and appear in green

numbers appear in blue

Visualisation of bigbang with View()

```
22  ###Different kinds of objects and data
23  #create vector
24  id=c("Leonard","Sheldon","Penny","Howard","Bernadette","Amy","Raj")
25  id
26  sex=c("M","M","F","M","F","F","M")
27  age_now=c(45,43,32,37,38,42,37)
28  #create dataframe
29  bigbang=data.frame(id,sex,age_now)
30  View(bigbang)
```

r_presentation.R × bigbang ×

Filter

	id	sex	age_now
1	Leonard	M	45
2	Sheldon	M	43
3	Penny	F	32
4	Howard	M	37
5	Bernadette	F	38
6	Amy	F	42
7	Raj	M	37

Showing 1 to 7 of 7 entries

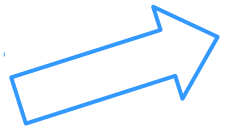
Different kinds of objects and data

```
22  ###Different kinds of objects and data
23  #create vector
24  id=c("Leonard","Sheldon","Penny","Howard","Bernadette","Amy","Raj")
25  id
26  sex=c("M","M","F","M","F","F","M")
27  age_now=c(45,43,32,37,38,42,37)
28  #create dataframe
29  bigbang=data.frame(id,sex,age_now)
30  View(bigbang)
31  #different type of data
32  class(bigbang)
33  class(id)
34  class(age_now)
35
```

```
Console   Terminal ×
~/
> #different type of data
> class(bigbang)
[1] "data.frame"
> class(id)
[1] "character"
> class(age_now)
[1] "numeric"
```

Different kinds of objects and data

```
22  ###Different kinds of objects and data
23  #create vector
24  id=c("Leonard","Sheldon","Penny","Howard","Bernadette","Amy","Raj")
25  id
26  sex=c("M","M","F","M","F","F","M")
27  age_now=c(45,43,32,37,38,42,37)
28  #create dataframe
29  bigbang=data.frame(id,sex,age_now)
30  View(bigbang)
31  #different type of data
32  class(bigbang)
33  class(id)
34  class(age_now)
35
40  class(45)
41  class("45")
```



```
> class(45)
[1] "numeric"
> class("45")
[1] "character"
```

characters are written as ''character'' so ''45'' is recognized as a character and not as a number

Sort your data

```
37  ###Sort your data
38  #names in function of sex
39  bigbang$id[bigbang$sex=="M"]
40  bigbang$id[bigbang$sex %in% "M"]
```

data$col refers to the column named col of the table named data

sorting is done on the elements between brackets [ ]

Sort your data

```
37  ###Sort your data
38  #names in function of sex
39  bigbang$id[bigbang$sex=="M"]
40  bigbang$id[bigbang$sex %in% "M"]
```

Console Terminal

```
> ###Sort your data
> #names in function of sex
> bigbang$id[bigbang$sex=="M"]
[1] Leonard Sheldon Howard  Raj
Levels: Amy Bernadette Howard Leonard Penny Raj Sheldon
> bigbang$id[bigbang$sex %in% "M"]
[1] Leonard Sheldon Howard  Raj
Levels: Amy Bernadette Howard Leonard Penny Raj Sheldon
>
```

Make a subtable

```
37  ###Sort your data
38  #names in function of sex
39  bigbang$id[bigbang$sex=="M"]
40  bigbang$id[bigbang$sex %in% "M"]
41  #make a subtable
42  bigbang_F=bigbang[bigbang$sex=="F",]
43  View(bigbang_F)
```

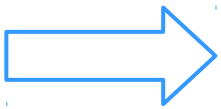
As we want a subtable with 2 dimensions, it is very important **not to forget the comma in the brackets**

r_presentation.R | bigbang

Filter

	id	sex	age_now
1	Leonard	M	45
2	Sheldon	M	43
3	Penny	F	32
4	Howard	M	37
5	Bernadette	F	38
6	Amy	F	42
7	Raj	M	37

Showing 1 to 7 of 7 entries



r_presentation.R | bigbang_F | bigba

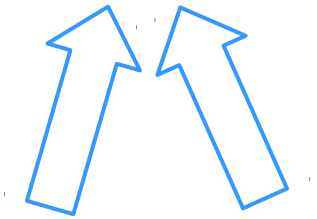
Filter

	id	sex	age_now
3	Penny	F	32
5	Bernadette	F	38
6	Amy	F	42

Showing 1 to 3 of 3 entries

Navigate into your table: subscript

```
44  #find a specific content: exemple row 5, column 1
45  bigbang[5,1]
```



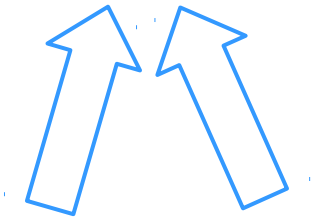
row number

column number

```
> #find a specific content: exemple row 5, column 1
> bigbang[5,1]
[1] Bernadette
Levels: Amy Bernadette Howard Leonard Penny Raj Sheldon
```

Navigate into your table

```
44  #find a specific content: exemple row 5, column 1
45  bigbang[5,1]
```



row number

column number

r_presentation.R | bigbang

Filter

	id	sex	age_now
1	Leonard	M	45
2	Sheldon	M	43
3	Penny	F	32
4	Howard	M	37
5	Bernadette	F	38
6	Amy	F	42
7	Raj	M	37

Showing 1 to 7 of 7 entries

Console | Terminal

~/

```
> #find a specific content:
> bigbang[5,1]
[1] Bernadette
Levels: Amy Bernadette Howa
```

Navigate into your table

```
44  #find a specific content: exemple row 5, column 1
45  bigbang[5,1]
46  bigbang[,1]
47  bigbang$id
48  bigbang[1,]
49
```

r_presentation.R × | bigbang ×

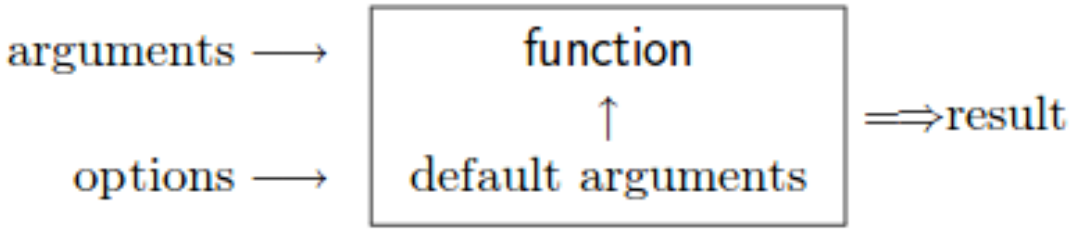
Filter

	id	sex	age_now
1	Leonard	M	45
2	Sheldon	M	43
3	Penny	F	32
4	Howard	M	37
5	Bernadette	F	38
6	Amy	F	42
7	Raj	M	37

Showing 1 to 7 of 7 entries

Let’s now use functions

- Functions in R are written as following: function()
- You can specify options and arguments in function of your needs
- The arguments you do not specify will be set by default



E.Paradis, R for beginners, 2005

Let’s now use functions

```
51  ###Functions
52  #use pre existing functions
53  mean(bigbang$age_now)
54  sum(bigbang$age_now)
```

You can also create your own function

- Let’s write a function to calculate the age of the actors at the beginning of the serie, so twelve years ago

```
55  #create your own function
56  age_season1=function(age){age-12}
57  age_season1(bigbang$age_now[bigbang$id=="Sheldon"])
58  age_season1(26)
```

Syntax to write your function:
myfunction=function(x) {what to do}

You can also create your own function

- Let’s write a function to calculate the age of the actors at the beginning of the serie, so twelve years ago

```
55  #create your own function
56  age_season1=function(age){age-12}
57  age_season1(bigbang$age_now[bigbang$id=="Sheldon"])
58  age_season1(26)
59  #add a new column to your data
60  bigbang$age_start=c(age_season1(bigbang$age_now))
```

You can also create your own function

- Let’s write a function to calculate the age of the actors at the beginning of the serie, so twelve years ago

```
55  #create your own function
56  age_season1=function(
57  age_season1(bigbang$a
58  age_season1(26)
59  #add a new column to
60  bigbang$age_start=c(a
61  View(bigbang)
```

r_presentation.R × bigbang ×

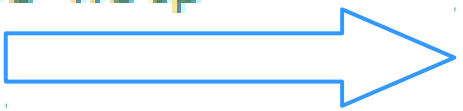
Filter

	id	sex	age_now	age_start
1	Leonard	M	45	33
2	Sheldon	M	43	31
3	Penny	F	32	20
4	Howard	M	37	25
5	Bernadette	F	38	26
6	Amy	F	42	30
7	Raj	M	37	25

Showing 1 to 7 of 7 entries

Finding help in R

```
64  ###Find help
65  ?mean
66  ??mean
```



Files | Plots | Packages | Help | Viewer

R: Arithmetic Mean | Find in Topic

mean {base} R Documentation

Arithmetic Mean

Description

Generic function for the (trimmed) arithmetic mean.

Usage

```
mean(x, ...)

## Default S3 method:
mean(x, trim = 0, na.rm = FALSE, ...)
```

Arguments

x	An R object. Currently there are methods for numeric/logical vectors and date, date-time and time interval objects. Complex vectors are allowed for `trim = 0`, only.
trim	the fraction (0 to 0.5) of observations to be trimmed from each end of x before the mean is computed. Values of trim outside that range are taken as the nearest endpoint.
na.rm	a logical value indicating whether NA values should be stripped before the computation proceeds.
...	further arguments passed to or from other methods.

Value

If `trim` is zero (the default), the arithmetic mean of the values in x is computed, as a numeric or complex vector of length one. If x is not logical (coerced to numeric), numeric (including integer) or complex, `NA_real_` is returned, with a warning.

If `trim` is non-zero, a symmetrically trimmed mean is computed with a fraction of `trim` observations deleted from each end before the mean is computed.

References

Becker, R. A., Chambers, J. M. and Wilks, A. R. (1988) *The New S Language*. Wadsworth & Brooks/Cole.

See Also

`weighted.mean`, `mean.POSIXct`, `colMeans` for row and column means.

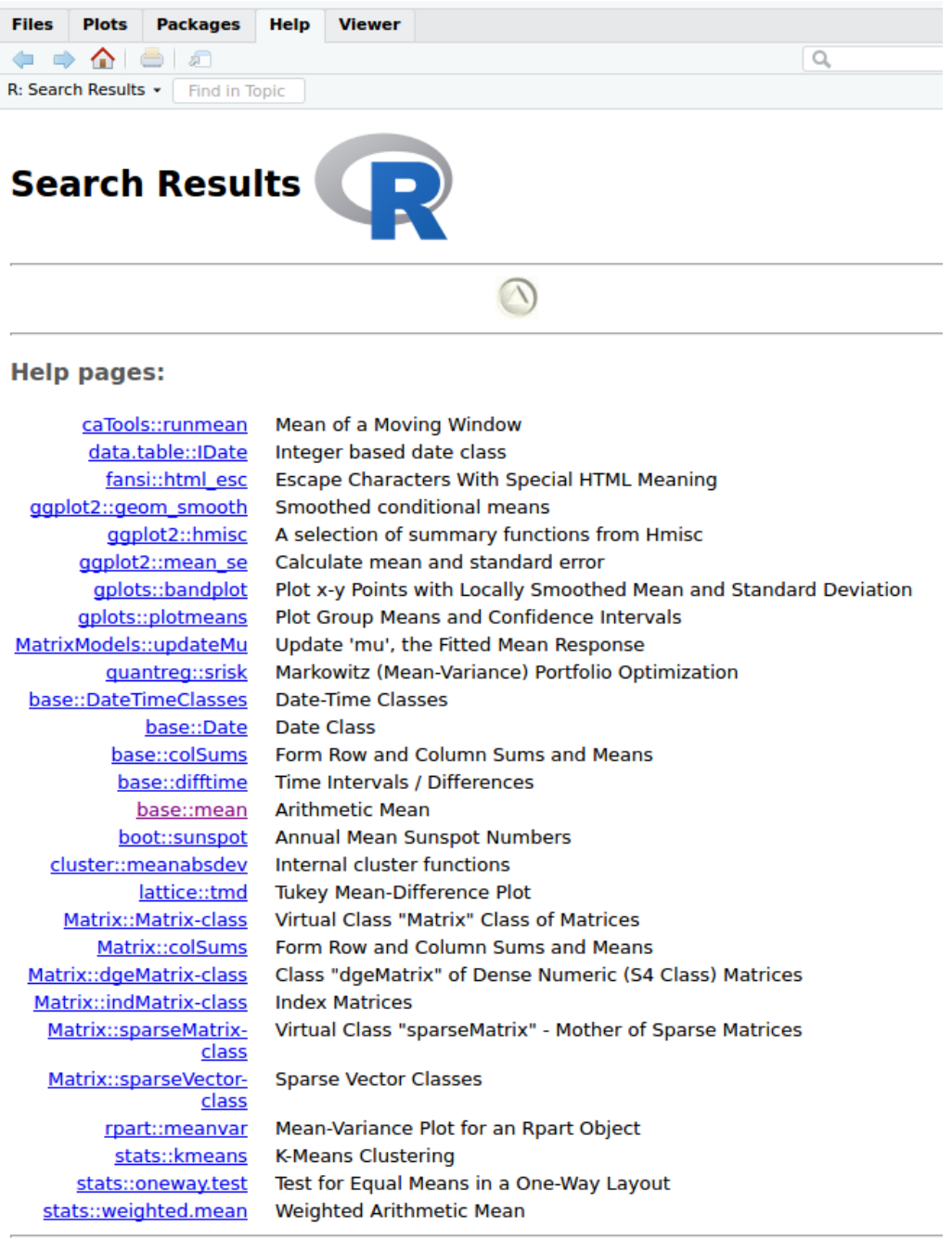
Examples

```
x <- c(0:10, 50)
xm <- mean(x)
c(xm, mean(x, trim = 0.10))
```

Finding help in R

```
64  ###Find help
65  ?mean
66  ??mean
```

If you do not know the exact name of the function, use ??



Finding help on the web

- Books/ebooks
  - Emmanuel Paradis – R for beginners
  - Michael Crawley - The R book
  - Andrie de Vries & Joris Meys - R for dummies
- Websites
  - Statistical Tools for High-Throughput Data Analysis (www.sthda.com)
  - Stackoverflow (http://stackoverflow.com)
- Forums
  - R-bloggers (www.r-bloggers.com)

Do not ask questions on forums without doing preliminary research

Give a sample of data and script to illustrate your problem

Export the data created on R in csv file

```
76  ###Export your data
77  #write.csv(bigbang, "~/Bureau/bigbang.csv")
78  write.csv(bigbang, "bigbang.csv")
```

To easily find the pathway, you can right-clic on a document located in the place of interest and clic on properties, the pathway will be indicated

Alternatively to write the pathway, you can set your working directory manually:
Session>Set working directory>choose location

Import your own data in R studio

- A few precautions before to import a document
  - Decimal numbers have to be written using a dot and not a comma: 2.5 instead of 2,5
  - Do not mix empty cases and cases with NA in a same column
  - Replace #VALUE ! (that happens in excell sheet when you apply function on Nas) by NA
  - Convert your excell sheet in .csv format

Import your own data in R studio

- Iris dataset is a free dataset avaiable in R and regularly used in courses and examples

```
80  ###Import your data
81  #iris=read.csv("~/Bureau/iris.csv",header=T)
82  iris=read.csv("iris.csv",header=T)
83  View(iris)
```

To easily find the pathway, you can right-clic on a document located in the place of interest and clic on properties, the pathway will be indicated

If you specified that the column separator of your csv file is tab, add the following argument:
read.csv(‘’path/filename.csv’’, sep= ‘’\t’’)

Import your own data in R studio

```
80  ###Import your data
81  #iris=read.csv("~/Bureau/iris.csv",header=T)
82  iris=read.csv("iris.csv",header=T)
83  View(iris)
```

r_presentation.R × | iris × | iris_vir × | bigbang_F × | bigbang ×

Filter

	Sepal.Length	Sepal.Width	Petal.Length	Petal.Width	Species
1	5.1	3.5	1.4	0.2	setosa
2	4.9	3.0	1.4	0.2	setosa
3	4.7	3.2	1.3	0.2	setosa
4	4.6	3.1	1.5	0.2	setosa
5	5.0	3.6	1.4	0.2	setosa
6	5.4	3.9	1.7	0.4	setosa
7	4.6	3.4	1.4	0.3	setosa
8	5.0	3.4	1.5	0.2	setosa
9	4.4	2.9	1.4	0.2	setosa
10	4.9	3.1	1.5	0.1	setosa
11	5.4	3.7	1.5	0.2	setosa
12	4.8	3.4	1.6	0.2	setosa
13	4.8	3.0	1.4	0.1	setosa
14	4.3	3.0	1.1	0.1	setosa

Showing 1 to 14 of 150 entries

Remove the eventual NAs

```
73  ###Import your data
74  iris=read.csv("~/Bureau/iris.csv",header=T)
75  View(iris)
76  #remove eventual NA
77  iris=iris[!is.na(iris$Species),]
```

is.na() return the cases that contain Nas

!is.na() make the contrary: it gives you the cases where there is something else than NA

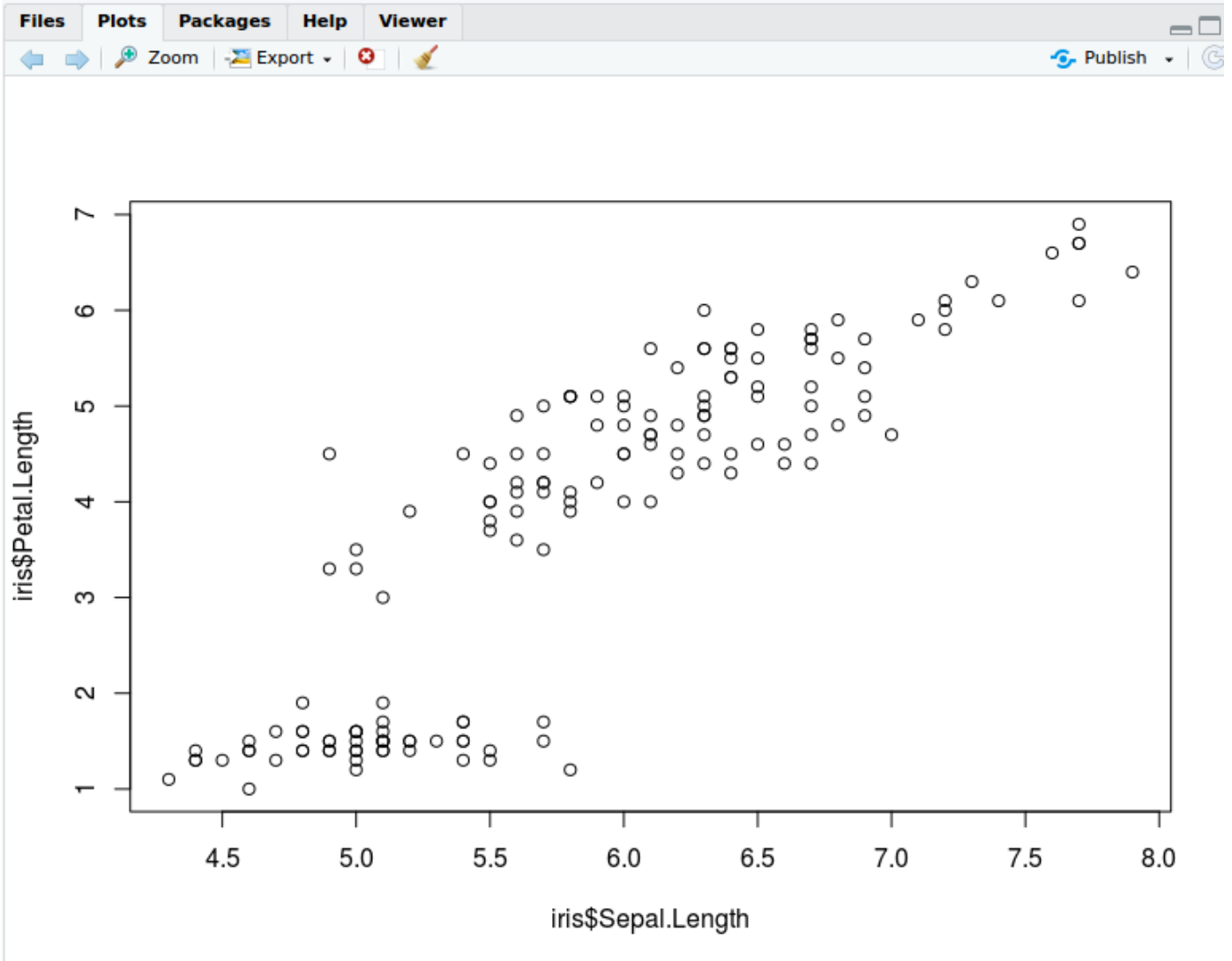
Create sub-tables in function of the specie

```
73  ###Import your data
74  iris=read.csv("~/Bureau/iris.csv",header=T)
75  View(iris)
76  #remove eventual NA
77  iris=iris[!is.na(iris$Species),]
78  #make subtables
79  levels(iris$Species)
80  iris_set=iris[iris$Species=="setosa",]
81  iris_vers=iris[iris$Species=="versicolor",]
82  iris_vir=iris[iris$Species=="virginica",]
83  View(iris_vir)
```

As you want a subtable and not a vector, do not forget the comma after the the specie

Draw a plot

```
86  ###Make plots
87  #simple plot
88  plot(iris$Sepal.Length,iris$Petal.Length)
```



Draw a nicer plot

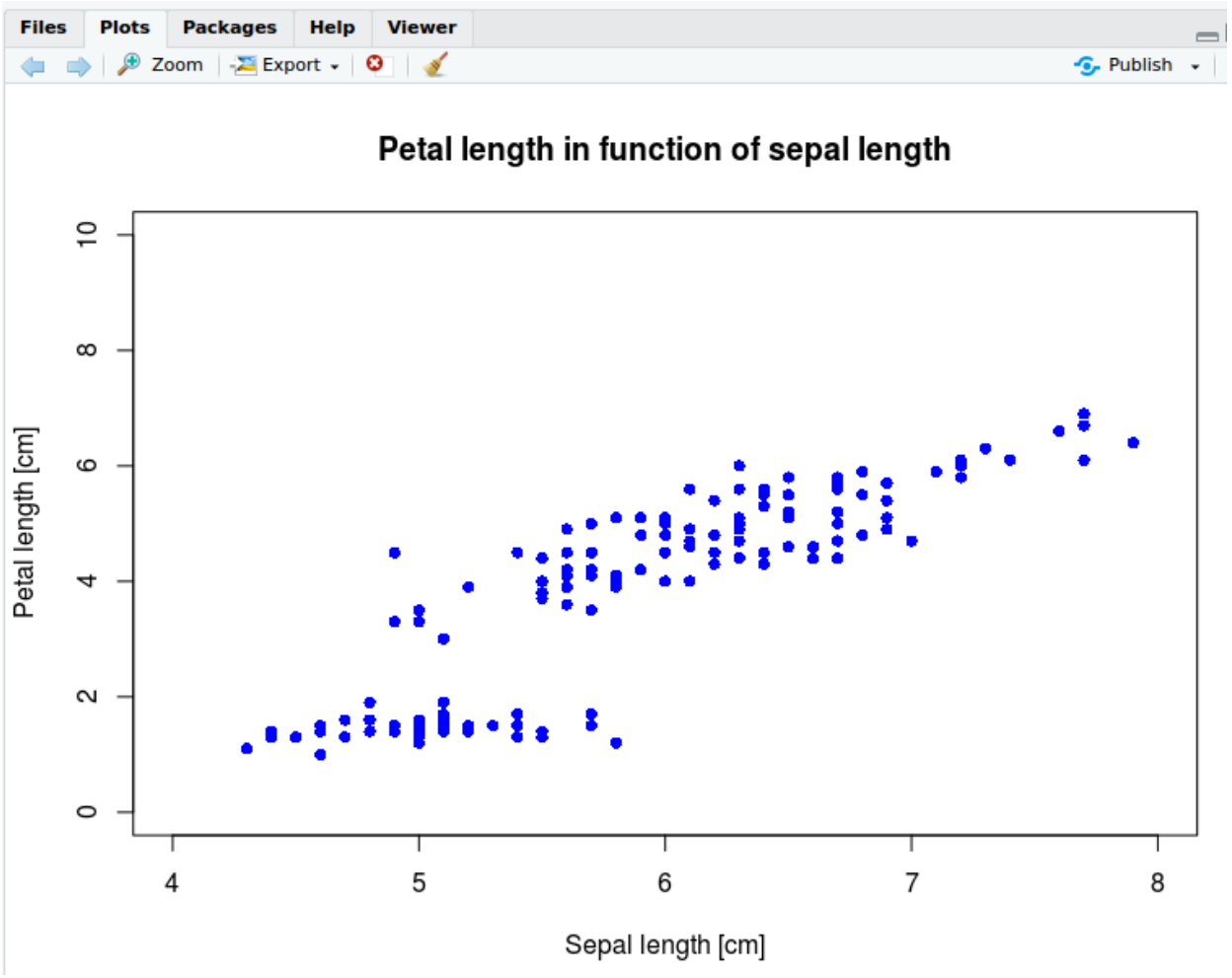
```
89  #add argmuents
90  plot(iris$Sepal.Length,iris$Petal.Length,
91       main="Petal length in function of sepal length",
92       xlab="Sepal length [cm]",ylab="Petal length [cm]",
93       col="blue",pch=16,
94       ylim=c(0,10),xlim=c(4,8))
```

title

y and x subtitles

color and type of dot

limits of the y and x axis



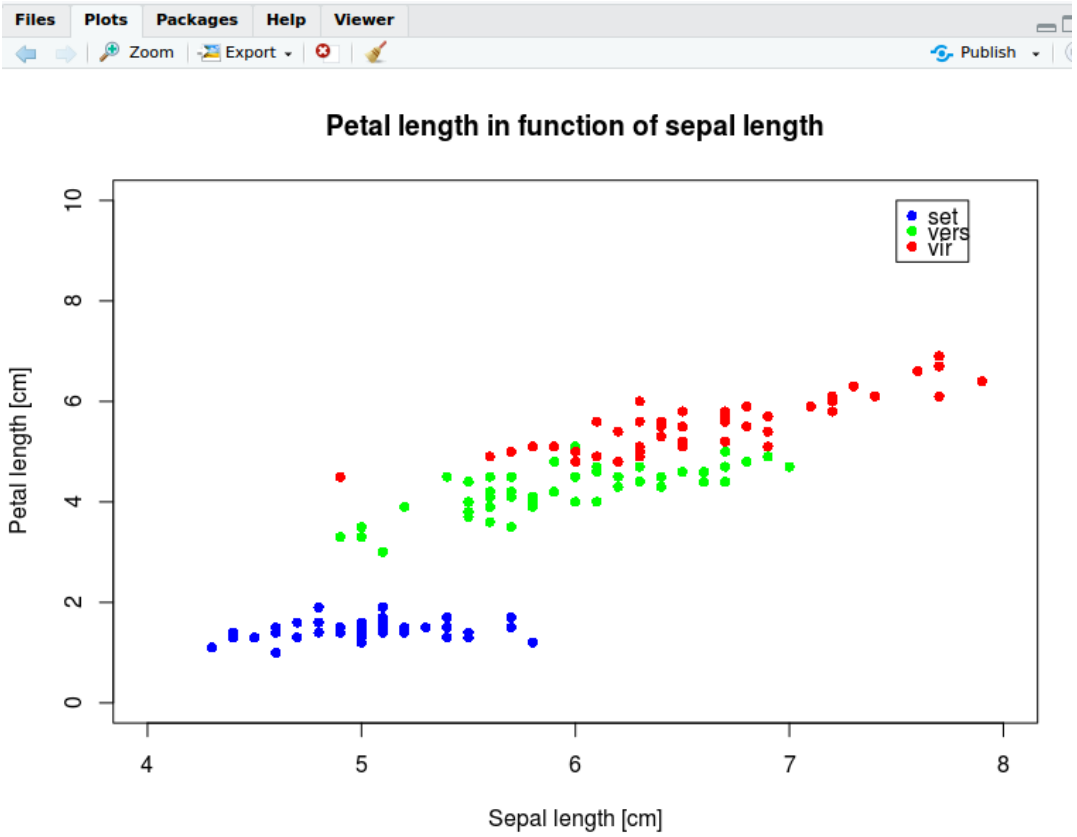
Draw the nicest plot

use the subtable

use points() to add data from other subtables

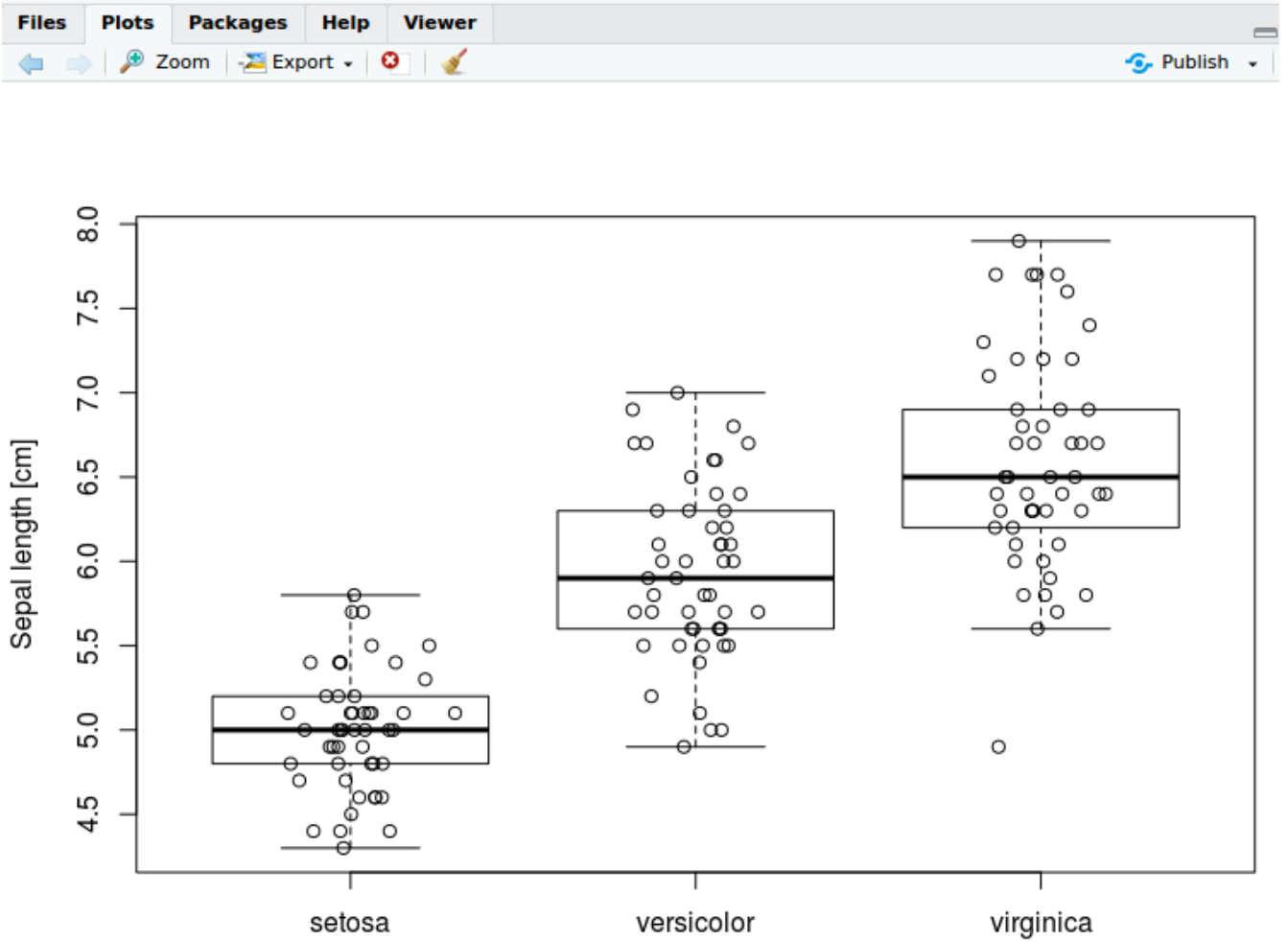
```
95  #add groups of dots of different colors
96  plot(iris_set$Sepal.Length,iris_set$Petal.Length,
97       main="Petal length in function of sepal length",
98       xlab="Sepal length [cm]",ylab="Petal length [cm]",
99       col="blue",pch=16,
100      ylim=c(0,10),xlim=c(4,8))
101 points(iris_vers$Sepal.Length,iris_vers$Petal.Length,col="green",pch=16)
102 points(iris_vir$Sepal.Length,iris_vir$Petal.Length,col="red",pch=16)
103 legend(7.5,10,legend=c("set","vers","vir"),col=c("blue","green","red"),pch=16)
```

eventually add legend

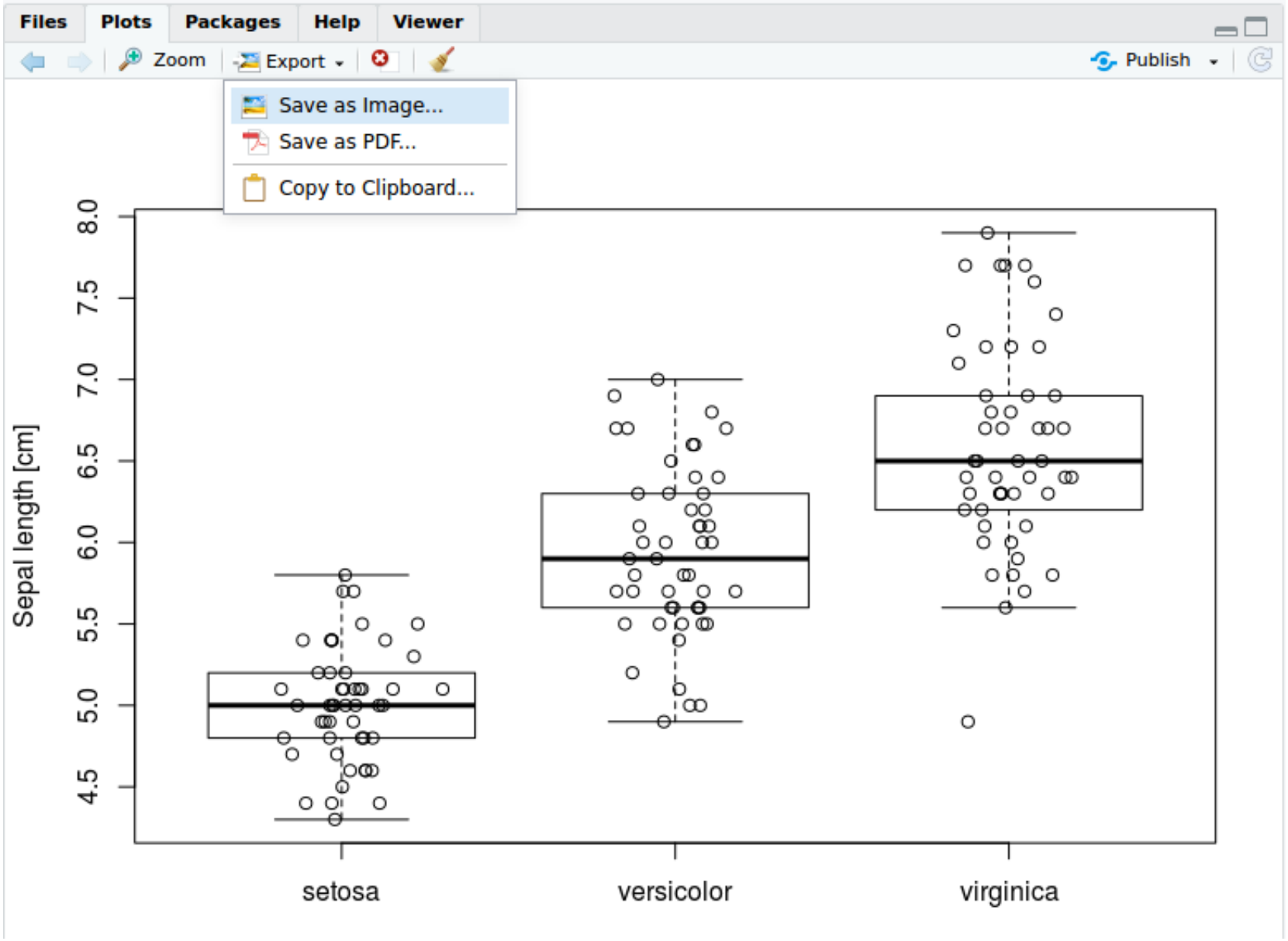


What about a boxplot?

```
112  #add dots plus barplot
113  boxplot(iris_set$Sepal.Length,iris_vers$Sepal.Length,iris_vir$Sepal.Length,
114          names=c("setosa","versicolor","virginica"),
115          ylab="Sepal length [cm]",outline=F)
116  points(rnorm(length(iris_set$Sepal.Length),1,0.1),iris_set$Sepal.Length)
117  points(rnorm(length(iris_vers$Sepal.Length),2,0.1),iris_vers$Sepal.Length)
118  points(rnorm(length(iris_vir$Sepal.Length),3,0.1),iris_vir$Sepal.Length)
```



Save your plot



Other usefull function linked to plots

```
129  #other functions to do plots
130  hist()
131  barplot()
132  pie()
133
134  #other usefull functions
135  par(mfrow=c(2,2))
136  arrows(x0,y0,x1,y1,angle)
137  text(x,y,"mytext")
```

other kinds of plots

split your plot window to see several plots (here 2x2)

add arrows or lines (angle=0) on your plots

add text on your plot

When you add elements to your plot, that are not between the axis, add the argument xpd=T to see it

A few lines of statistics: is the sepal length different between species?

```
129  ###Statistics: example
130  #Is the sepal length different between species?
131  wilcox.test(iris_set$Sepal.Length,iris_vers$Sepal.Length)
132  wilcox.test(iris_vers$Sepal.Length,iris_vir$Sepal.Length)
133  wilcox.test(iris_set$Sepal.Length,iris_vir$Sepal.Length)
134
```

130:17 R & R studio presentation script

Console Terminal

~/

```
> #Is the sepal length different between species?
> wilcox.test(iris_set$Sepal.Length,iris_vers$Sepal.Length)

        Wilcoxon rank sum test with continuity correction

data:  iris_set$Sepal.Length and iris_vers$Sepal.Length
W = 168.5, p-value = 8.346e-14
alternative hypothesis: true location shift is not equal to 0

> wilcox.test(iris_vers$Sepal.Length,iris_vir$Sepal.Length)

        Wilcoxon rank sum test with continuity correction

data:  iris_vers$Sepal.Length and iris_vir$Sepal.Length
W = 526, p-value = 5.869e-07
alternative hypothesis: true location shift is not equal to 0

> wilcox.test(iris_set$Sepal.Length,iris_vir$Sepal.Length)

        Wilcoxon rank sum test with continuity correction

data:  iris_set$Sepal.Length and iris_vir$Sepal.Length
W = 38.5, p-value < 2.2e-16
alternative hypothesis: true location shift is not equal to 0
```

To conclude...

- Basic tools
  - Create, use, visualise vectors and data frames
  - Use and create functions
  - Find help
  - Draw and save plots and boxplots
  - Import and export your data
  - A few lines of statistics
- Many more possibilities in function of your needs
  - automatisation of repetitive tasks: for boucle and apply()
  - specific packages
- Practise !