

Introduction to R-part1

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Before we start,, why **R** you here? [Opening statement]

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What is R?

Program language, widely used among researchers used for statistical computing, big data processing, and elegant graphics

#Creator; Ross Ihaka



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Why R?

1. It is completely free
2. It is open-source
3. useful for large data “unlike spreadsheets”
4. Top quality graphics
5. big community of users among researchers and easy support
6. Superiority by far over current statistical and graphical softwares available such as SPSS , SAS, Matlab and others

http://www.burns-stat.com/pages/Tutor/R_relative_statpack.pdf (http://www.burns-stat.com/pages/Tutor/R_relative_statpack.pdf)

From my opinion, 3.5 reason to switch to R

1. Safety

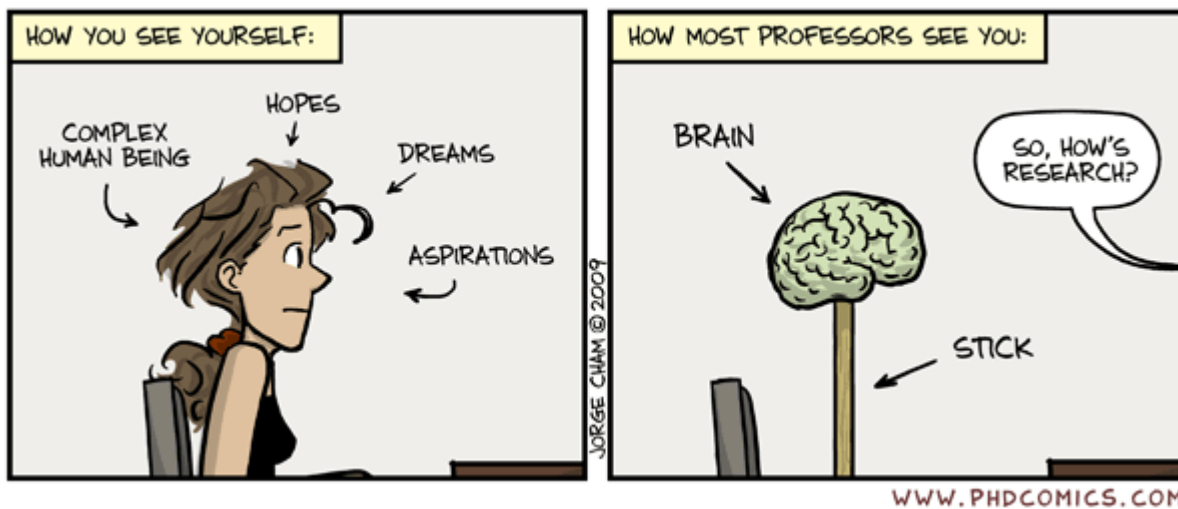
2. Speed

3. No reason spread sheet have poor graphics and statistics, but simply, good graphics and statistics haven't happened

0.5. I like it (its classy) like this cat:)



Additionally, R will solve this problem with your professor



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Researchers were asked in a marketing survey, tell me what you want? but guess what, majority of them (~73%) don't know their

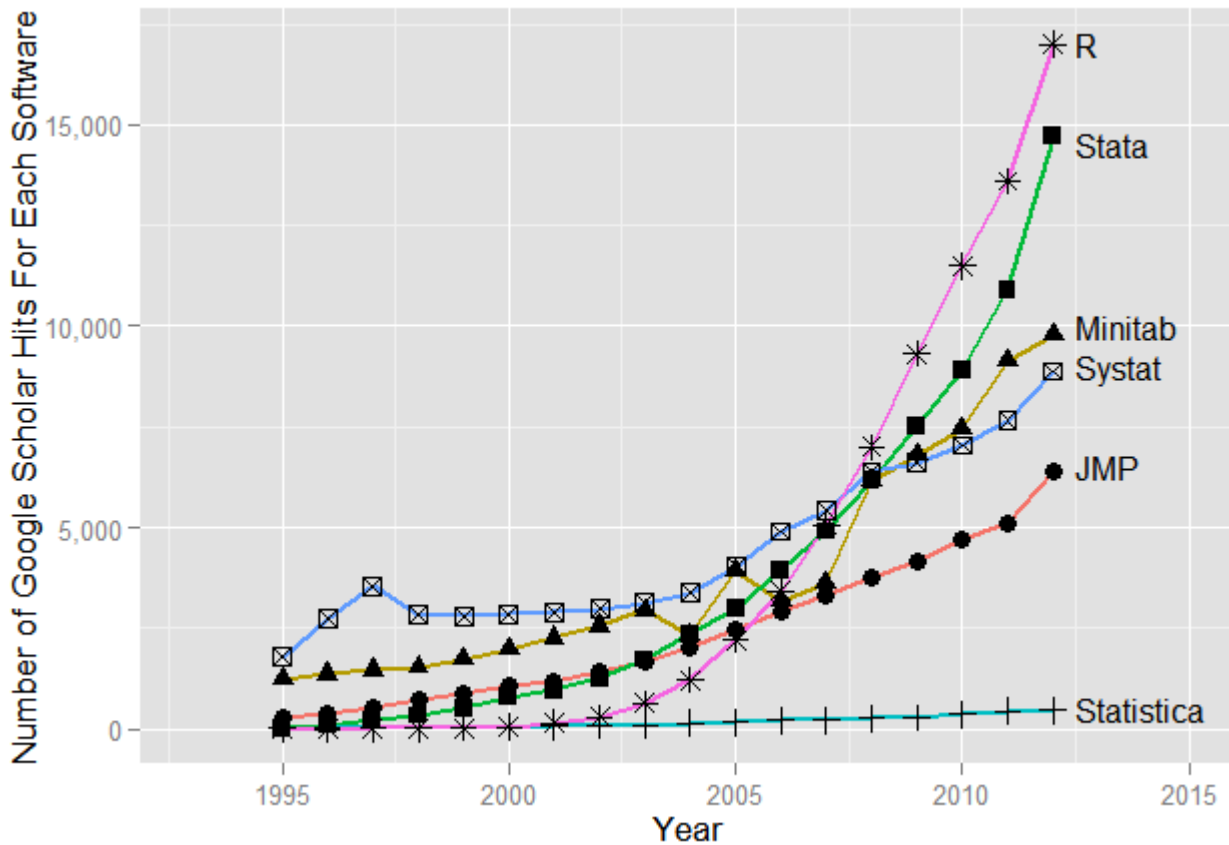
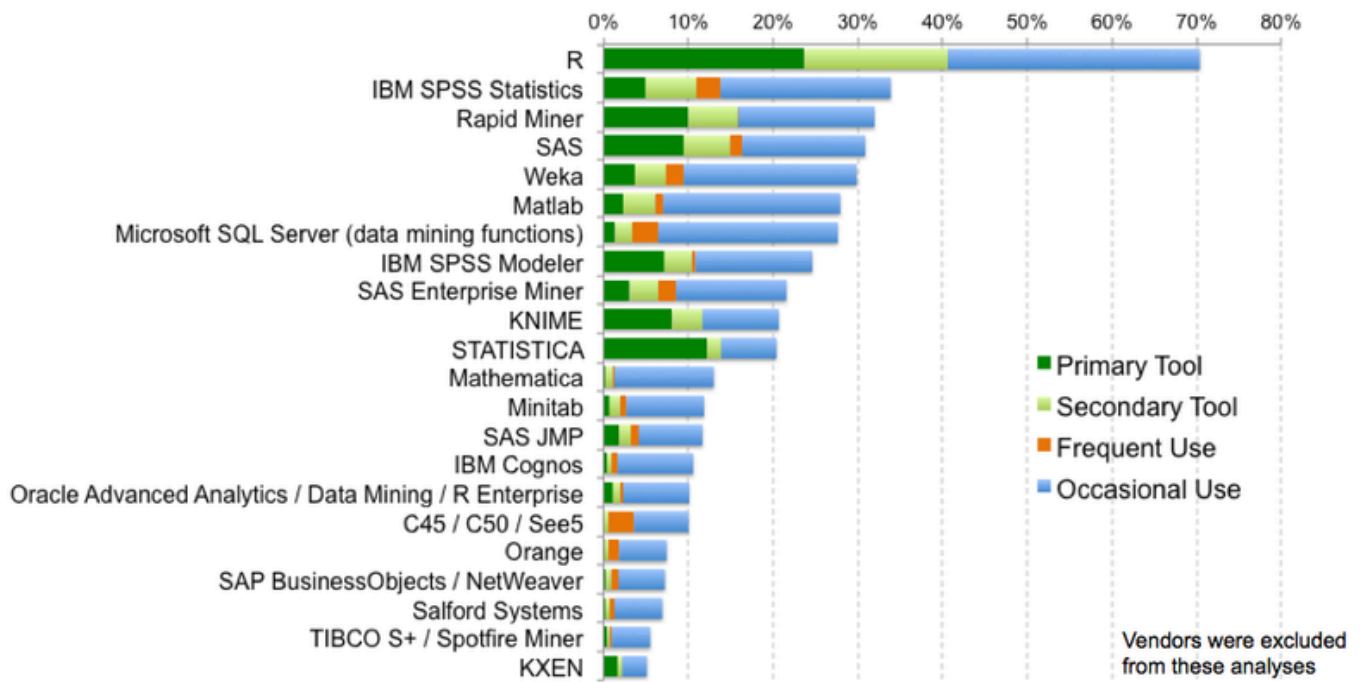
needs!!

**"People don't know what they want
until you show it to them"**

Steve Jobs

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R users statistics



** You can find solution packages for your professional needs

R in science

Researchers have used R to devise software packages in all kinds of disciplines. A few are listed below; there are thousands more at the [Comprehensive R Archive Network \(CRAN\)](#).

Astrophysics

The [solarR](#) package provides functions to determine the solar radiation that falls on Earth.

Carbon dating

[Bchron](#) creates chronologies based on radiocarbon- and non-radiocarbon-dated depths of sediments.

Climate science

[raincpc](#) allows researchers to obtain and analyse daily global rainfall data from the US National Oceanic and Atmospheric Administration's Climate Prediction Center.

Epidemiology

[DCluster](#) is a package for the detection of spatial clusters of diseases.

Chemistry

[ChemmineR](#) is a cheminformatics toolkit for analysing small molecules in R.

Genetics

[Bioconductor](#) provides tools for the analysis of high-throughput genomic data.

Pharmacokinetics

The [PKfit](#) package can model the half-life and dose absorption of drugs.

Palaeoecology

[Neotoma](#) provides access to data on pollen, fossil mammals and everything else on the Neotoma palaeoecology database.

Oceanography

[deSolve](#) is a package for solving differential equations.

Graphics

[ggplot2](#) one of the most popular visualization packages in R.

Phylogeny

[dendextend](#) compares trees of evolutionary relationships.

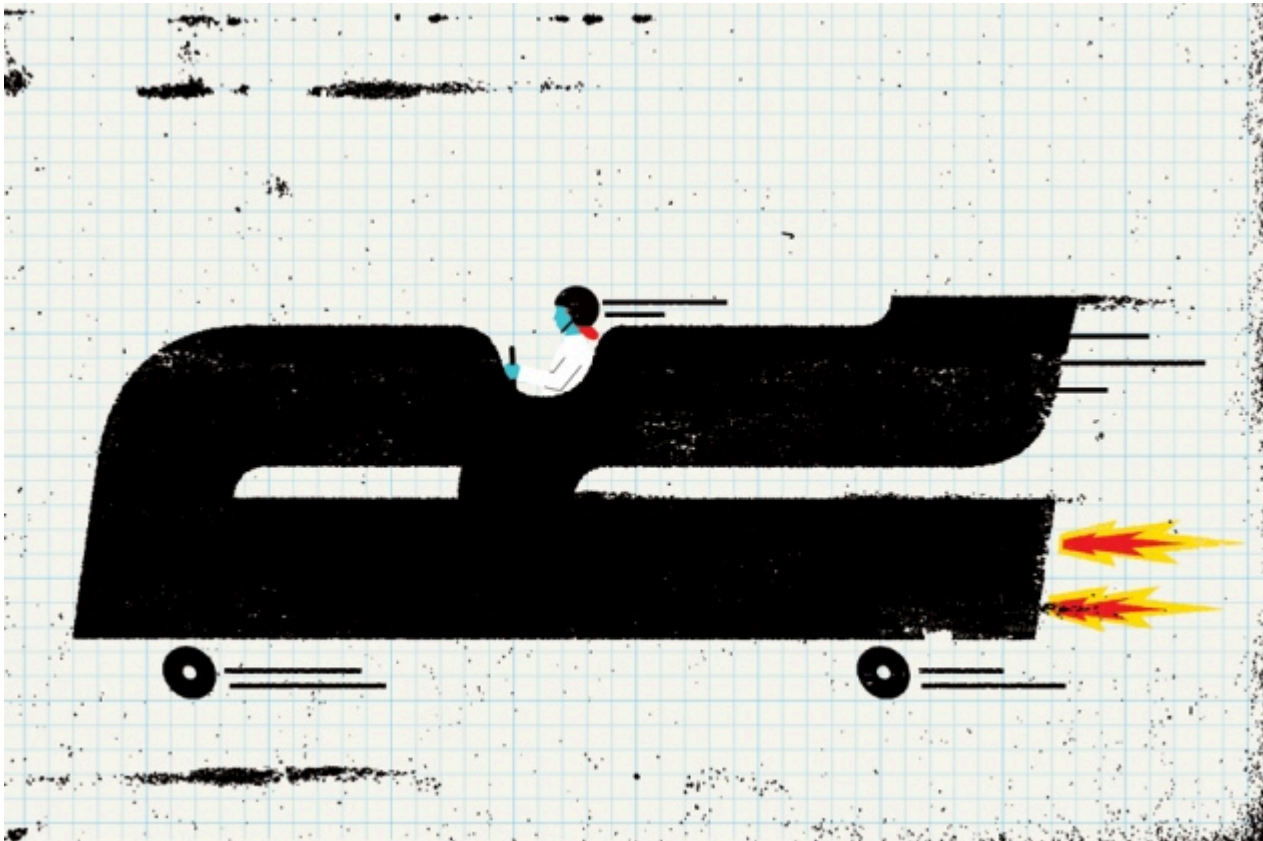
Genomics

The [QuasR](#) package lets researchers quantify and annotate short reads from sequencing experiments.

[Less](#) ▲

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So working with R is an adventure



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#Getting started (Installation)

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We need to download R from “Comprehensive R Archive Network- CRAN”

<http://cran.r-project.org/> (<http://cran.r-project.org/>)

Install it !

Hint1. During R installation, use default setting, such as C as destination folder.

Hint2. a list of components to install will appear, choose, core files, message translation and select one of 32 bit or 64 bit depend on your operating system.

Hint3. For startup option, select “No (Accept defaults)”

Hint4. In additional tasks select “create desktop icon, save version number, associate R with .RData”

Now you have R on your PC machine - navigate for the R logo

You can work with R as it is, this is called the naked version, however, another software called R studio makes life more easy in term of viewing and dealing with R. Note that u need to install R first then R studio, R studio can not work alone but uses R as a core.

you can download R studio from their webpage

<https://www.rstudio.com/products/rstudio/download/>
(<https://www.rstudio.com/products/rstudio/download/>)

Install it !

navigate for R studio “Blue circle with R”

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Working with R studio

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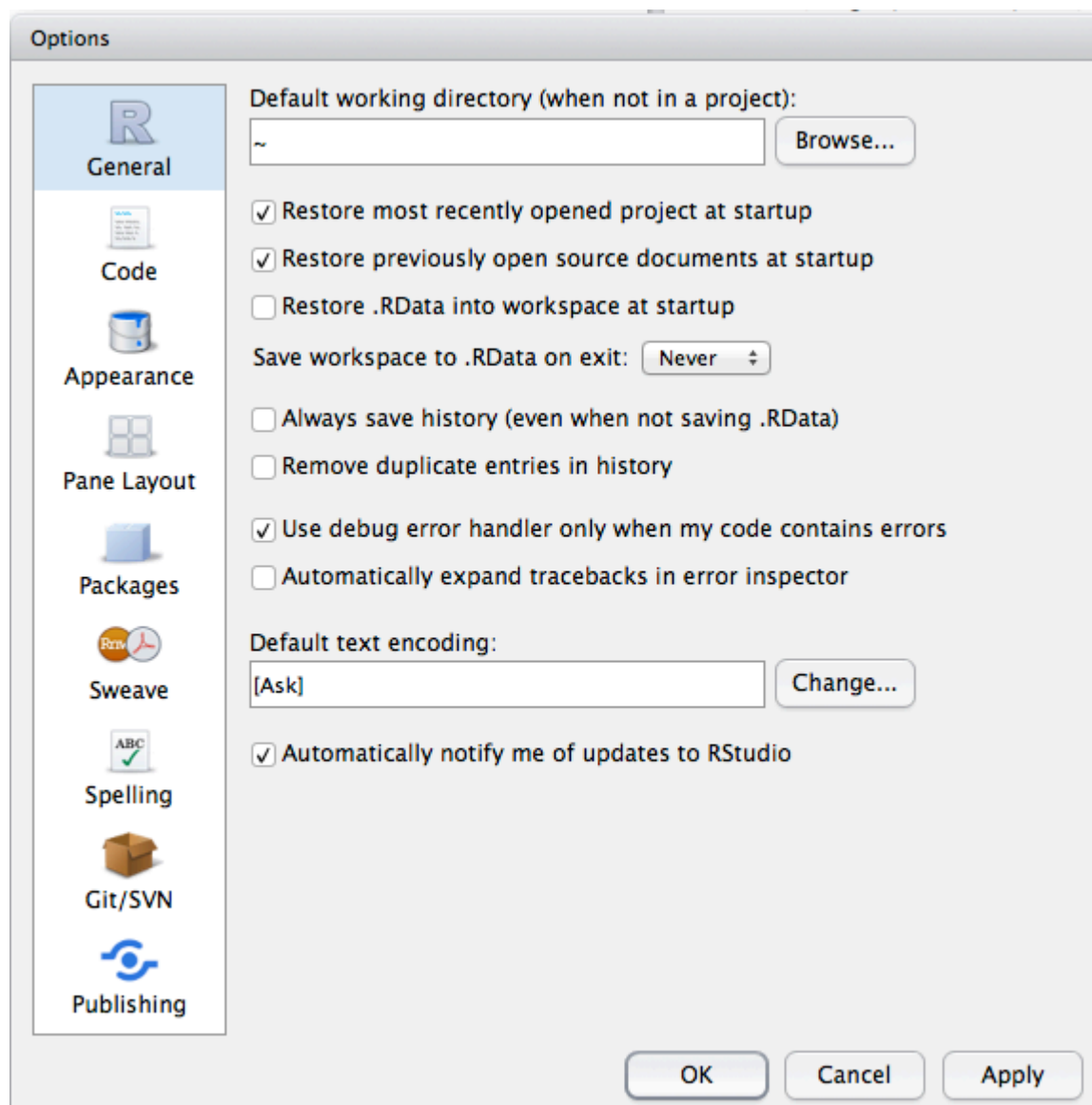
##Basic setting

###Before we dig deep in R , i would like to pass through some setting in R studio

###1. open R studio -> Tools-> global setting-General

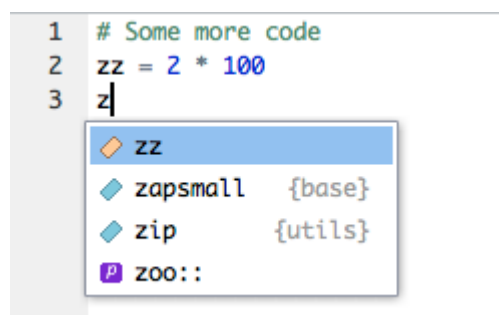
If you would like R studio to store the session and load it again, u need to check “restore .RData on exit”. some ppl like to start on a clean base, in that case “unckeck it and also set”save

workspace on exit to NEVER”



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R studio can do auto complete whenever you start typing then press “Tab” key a pop up menu will appear for you.



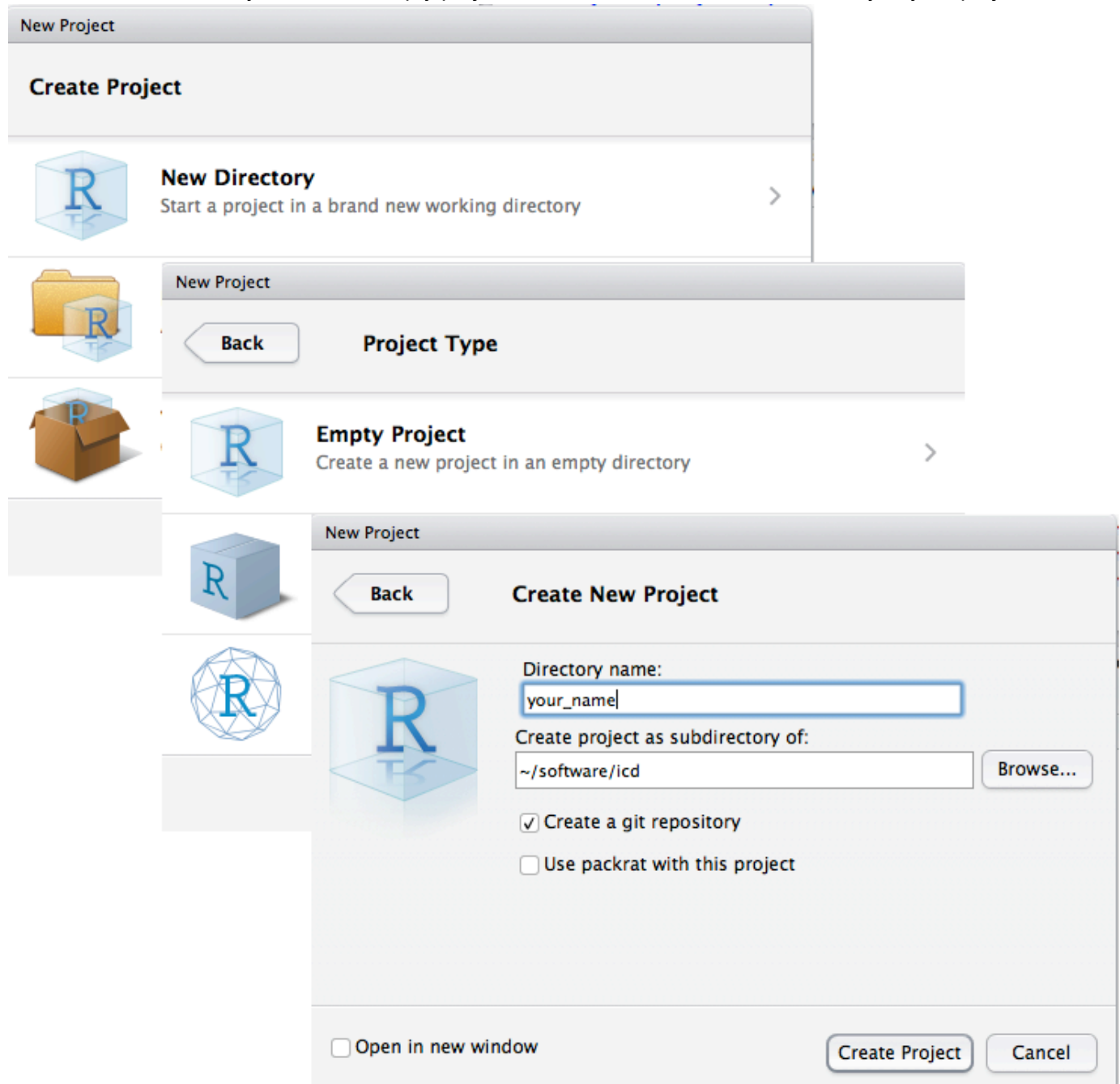
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###R studio is very powerfull, using projects management, you can control several project, working with several project at once.

Lets create a new R Studio project

Select File > New project

###Select New Directory ###Select Empty project ###Select a name and sub directory of your project.



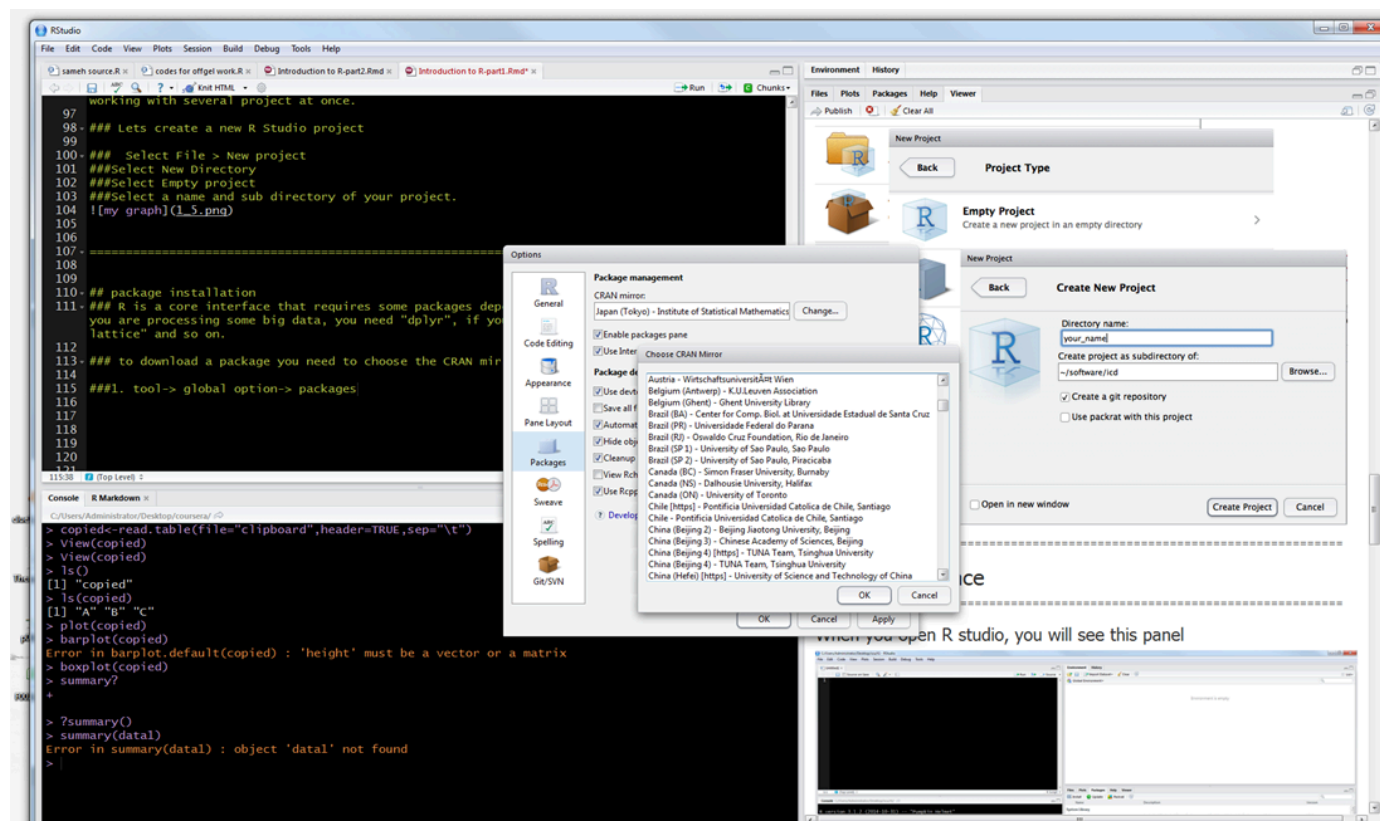
package installation

R is a core interface that requires some packages depending on what you do. For example if you are processing some big data, you need “dplyr”, if you want graphics you need “ggplot2 or

lattice" and so on.

to download a package you need to choose the CRAN mirror

###1. tool-> global option-> packages



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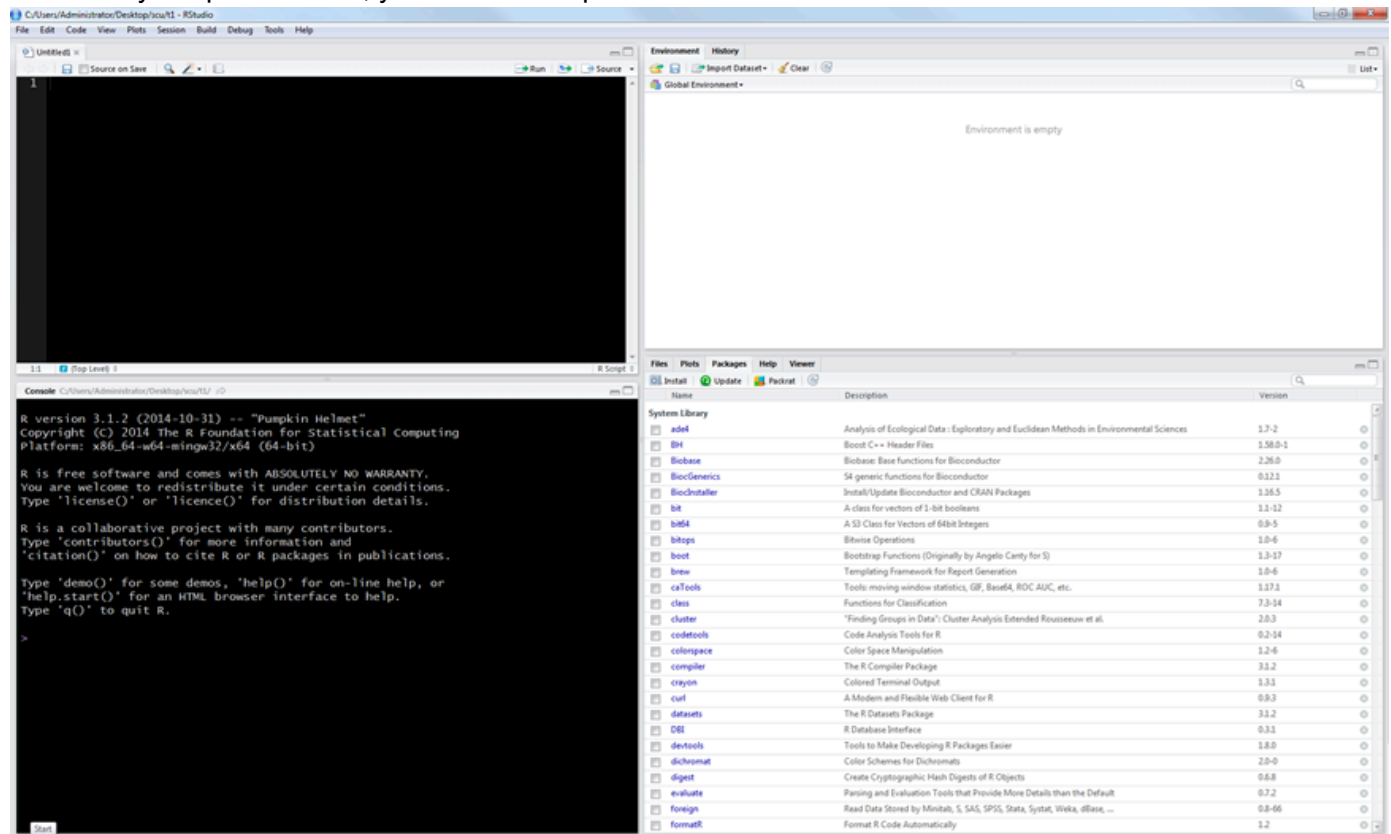
alternatively, you can search the package name (ex: ggplot2 download) download the zip or tar.gz file to your pc, then load it

packages -> Install-> Install from package archive-> browse your file-> install

###[in some cases, some packages require dependencies 'other prior packages to be installed, please read the error messages if its the case']

R studio interface

###When you open R studio, you will see this panel



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##Basic method to load your data

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The standard way to load your sample is to load it as comma separated files (.csv).

###There is some points you need to take care about ###1. from spreadsheet-> save your file as (.csv), use only 1 sheet in the excel workbook ###2. copy and paste your csv file in the same folder where you are working ###3. always try to use simple file names without any complex tags

here is your first code:)

```
data <- read.csv("test1.csv",header=T)
```

###lets try it !

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The golden code

This R code accept the data in the clipboard of your PC. so you copy your data and submit the code, the code gets what in the clipboard.

Here is the code

```
copied <- read.table(file="clipboard",header=TRUE,sep="\t")
```

The R built-in function

The function `file.choose()` allows you to browse files in your computer

```
Copied <- read.csv(file.choose())
```

###after loading your file to the R, you can check if your data is properly loaded by several ways

1. type `ls()` in the console

2. check global environment

##Now, you can start !

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###In general, and as routine work, i would like you to always save your codes in a script file. the extension is usually `.R` to do that

###1. File -> new file-> **new R script** # i recommend to create a script store file for your own use. ###2. type your code, select it (or stop the cursor on the line) and press run ###3. the code will be moved to the console and evaluated by R

lets try this on a file you loaded to R

1. open spreadsheet and create a table from 3 column named C, A, B
2. add 10 random values in each column
3. save the sheet as .csv (save as-> CSV comma delimited)
4. copy and paste the sheet in the folder you are working in in R (extension written below console) [most users forget this!]
5. Load your file into R (copy and paste the code in the R script then open the spread sheet, marke the table, copy it, then push run in R studio “top right of your script window”)
6. Ok, lets check if its loaded approbriatly or not

```
# ignore this part
A <- c(2,4,5,6,4,5,6,42,2)
B <- c(34,5,6,4,3,5,98,5,3)
C <- c(43,23,32,23,52,23,54,23,54)
copied <- data.frame(A,B,C)
copied
```

```
##      A  B  C
## 1   2 34 43
## 2   4  5 23
## 3   5  6 32
## 4   6  4 23
## 5   4  3 52
## 6   5  5 23
## 7   6 98 54
## 8  42  5 23
## 9   2  3 54
```

```
str(copied)
```

```
## 'data.frame':   9 obs. of  3 variables:
## $ A: num  2 4 5 6 4 5 6 42 2
## $ B: num 34 5 6 4 3 5 98 5 3
## $ C: num 43 23 32 23 52 23 54 23 54
```

```
class(copied)
```

```
## [1] "data.frame"
```

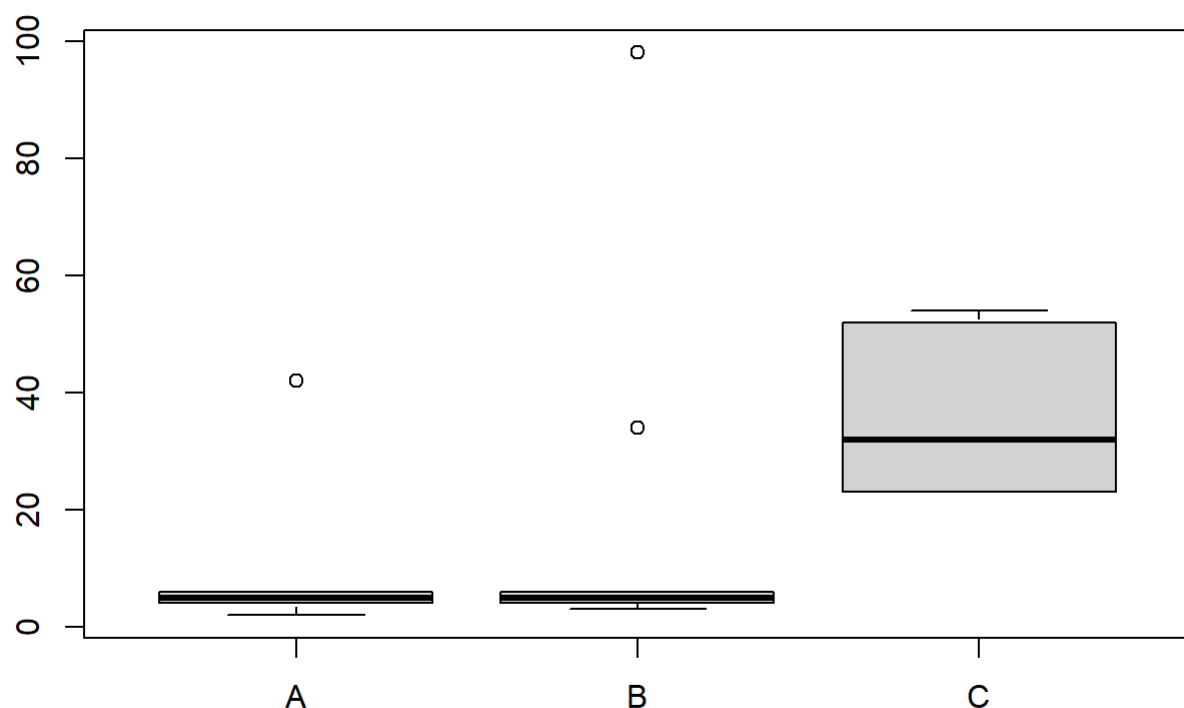
Lets see summary statistics for your data by passing this code

```
summary(copied)
```

```
##           A           B           C
## Min.      : 2.000   Min.      : 3.00   Min.      :23.00
## 1st Qu.:  4.000   1st Qu.:  4.00   1st Qu.:23.00
## Median :  5.000   Median :  5.00   Median :32.00
## Mean      :  8.444   Mean      :18.11   Mean      :36.33
## 3rd Qu.:  6.000   3rd Qu.:  6.00   3rd Qu.:52.00
## Max.      :42.000   Max.      :98.00   Max.      :54.00
```

after loading your data lets draw a boxplot by passing this simple code

```
boxplot(copied)
```



The help system

R includes a help system. for example if you are using a function like `summary` and you want to know its options

type

`?summary`

`help(summary)`

or ??summary

Congratulations!