

SC-Camp 2015 Introduction to R and Data Analysis

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SC-Camp 2015



- 1 Introduction to R
- 2 Crazy Examples
- 3 Practical Session
 Pre-requisites
 Objectives
 Practical Session Details





Summary

- 1 Introduction to R
- **2** Crazy Examples
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R?

R (pronounced aRrgh – pirate style) is a programming language and environment for statistical computing and graphics

- oriented towards data handling analysis and storage facility
- R Base
- Packages tools and functions (user contributed)
- R Base and most R packages are available from the Comprehensive R Archive Network (CRAN)
- Use R console or IDE: Rstudio, Deducer, vim/emacs...
- Comment is #, help is ? before a function name



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Using R

Installing/using packages

Install and load the ggplot2 package (even if already installed)

```
install.packages("ggplot2")
library(ggplot2)
```

Or in one step, install if not available then load:



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Using R

Usefull Functions

- List all objects in memory: 1s()
- Save an object: save(obj, file)
- Load an object: load(file)
- Set working directory: setwd(dir)





Data Structures

```
scalar:
```

$$s = 3.14$$

• vector:

$$v = c(1, 2, "ron")$$

list:

$$I = list(1:10, 'a', pi)$$

matrix:

$$m = matrix(seq(1:6), 2)$$

dataframe:

$$df = data.frame("col1" = seq(1:4), "col2" = c(5, 6, "cerveza", 6*7))$$

• . . .





Entering Data

Reading CSV or text files

```
# comma separated values
dat.csv <- read.csv(<file or url>)
# tab separated values
dat.tab <- read.table(<file or url>,
    header=TRUE, sep = "\t")
```





Entering Data

Reading data from other software: Excel, SPSS...

Excel Spreadsheets - need xlsx package

read.xlsx()

SPSS and Stata both need the foreign package

dat.dta <- read.dta(<file or url>)





Data Frames

Most easy structure to use, have a matrix structure.

- Observations are arranged as rows and variables, either numerical or categorical, are arranged as columns.
- Individual rows, columns, and cells in a data frame can be accessed through many methods of indexing.
- We most commonly use **object[row,column]** notation.





Accessing Items in a data.frame

Aside with R are provided example datasets, i.e. mtcars that can be used

```
data(mtcars)
head(mtcars)
colnames(mtcars)

# single cell value
mtcars[2,3]
# omitting row value implies all rows
mtcars[,3]
# omitting column values implies all columns
mtcars[2,]
```





Accessing Items in a data.frame

We can also access variables directly by using their names, either with **object[,"variable"**] notation or **object\$variable** notation.

```
# get first 10 rows of variable 'mpg' using two methods:
mtcars[1:10, "mpg"]
mtcars$mpg[1:10]
```





Exploring Data

Description Of Dataset

- Using dim, we get the number of observations(rows) and variables(columns) in the dataset.
- Using str, we get the structure of the dataset, including the class(type) of all variables. dim(mtcars) str(mtcars)
- summary when used on a dataset, returns distributional summaries of variables in the dataset. summary(mtcars)
- quantile function enables to get statistical metrics on the selected data quantile(mtcars\$mpg)





Exploring Data

Conditional Exploration

- subset enables to explore data conditionally subset(mtcars, cyl <= 5)
- **by** enables to call a particular function to sub-groups of data by(mtcars, mtcars\$cyl, summary)







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Random Text Generation

```
library(XML)
stem <- "http://www.5novels.com/classics/u5688"
hobbit <- NULL
for(i in 1:74) {
    if(i==1) { url <- paste0(stem, ".html") }</pre>
    else { url <- paste0(stem, "_", i, ".html") }
    x <- htmlTreeParse(url, useInternalNodes=TRUE)
    xx <- xpathApply(x, "//p", xmlValue)</pre>
    hobbit <- c(hobbit, gsub("\r", "", xx[-length(xx)]))
    Sys.sleep(0.5)
hobbit = paste(hobbit, collapse=' ')
```



Random Text Generation - 2

```
library(ngram)
ng2 <- ngram(hobbit, n=2)</pre>
```

babble(ng2, 128, seed=987654)





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Install and Run R

- On your local machine:
 - Find a release that fits your distribution at CRAN Archive
 - Install and launch R-Studio

```
http://cran.r-project.org/
https://www.rstudio.com/
```

On the cluster First connect to the cluster, then submit a job to run R.

```
\label{localhost} $$ ssh gaia-cluster $$ (frontend)$ oarsub -I -l core=1,walltime="00:30:00" (node)$ module load lang/R/3.2.0-ictce-7.3.5-bare (node)$ R
```

Install and Load a Package

```
(R-shell)$> install.packages("ggplot2")
(R-shell)$> library(ggplot2)
```





Objectives of this Practical Session

- Being able to plot data
 - histogram for data distribution
 - plot in different colors from different data sources
- Know some tips to organize your data
 - aggregate a dataset by column and apply an aggregation function
 - data.table package for binary search in datasets
 - performance in R operations
- R in parallel
 - on one machine
 - on a cluster with socket communications
 - MPI communications





Exercises

- Start the tutorial https://github.com/ULHPC/tutorials/tree/devel/advanced/R
 - Plot 2 graphs in section Simple Plotting
 - Answer 2 questions at the end of section Organizing your Data
 - Compare performance of aggregation operations w/wo parallelization
- Plot a speedup graph
 - with different number of cores and/or machines
 - needs: ggplot, parallel R





After the Practical Session: Problems

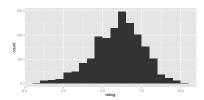
- regression models: example and exercise
- parallelization of K-means Clustering: example and exercise



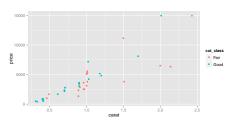


Simple Plotting

Movies Histogram:



Diamonds Plot with 2 colours:







PS Questions

Question: use ddply instead of tapply in the first example

```
ddply(DT, .(x), summarize, sum(v))
```

Question: return the min and max instead of the sum.

```
min max = function(data){
             c(min(data), max(data))
DT[,min max(v),by=x]
## or
DT[,c(min(v), max(v)),by=x]
```





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Usefull links

CRAN Archive

http://cran.r-project.org/

ggplot2 Documentation

http://docs.ggplot2.org/current/

 CRAN HPC Packages

http://cran.r-project.org/web/views/HighPerformanceComputing.html

Advanced R programming by Hadley Wickham

http://adv-r.had.co.nz/

