Introduction to R

Dr. Arabin Kumar Dey

Assistant Professor
Department of Mathematics
Indian Institute of Technology Guwahati

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Outline

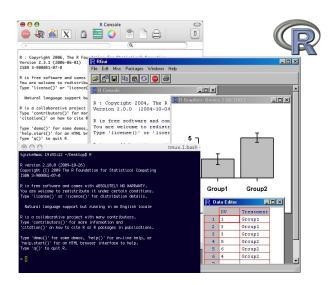
- Installing R
- 2 Why using R?
- Working with Data Set
- 4 Using Data Available in R

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Installing R on Linux/Windows

- Go to http://cran.r-project.org
- Select to download the latest version:
- Install and Open. The R window should look like :



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- Efficient functions and data structures for data analysis
- Powerful graphics
- Access to fast growing number of analysis packages
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- Starting R The R GUI versions under Windows and Mac OS X can be opened by double-clicking their icons.
- Alternatively, one can start it by typing 'R' in a terminal (default under Linux).
- The R environment is controlled by hidden files in the startup directory:.RData, .Rhistory and .Rprofile (optional).
- ## Closing R > q()Save workspace image? [y/n/c]:
- Note When responding with 'y', then the entire R workspace will be written to the .RData file which can become very large. Often it is sufficient to just save an analysis protocol in an R source file. This way one can quickly regenerate all data sets and objects.

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Installation of Cran package

- Linux install.packages("mypackage_1.0.tar.gz", repos=NULL)
- Hands-on Experience · · ·

- ## Create an object with the assignment operator '< –' (or '=')
 - > object $<-\cdots$
- ## List objects in current R session> Is()
- ## Return content of current working directory> dir()
- ## Return path of current working directory > getwd()
- ## Change current working directorysetwd("/home/user")

- ## General R command syntax
 - > object < function(arguments)
 - > object < object[arguments]
- ## Execute an R script
 - > source("my script.R")
- ## Execute an R script from command-line
 - > R CMD BATCH my_script.R
 - > R -slave < my_script.R
- ## Finding help
 - >?function
- ## Load a library
 - > library("my_library")
- ## Summary of all functions within a library
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- ## Numeric data: 1, 2, 3 > x < -c(1, 2, 3); x; is.numeric(x); as.character(x)
- ## Character data: "a", "b", "c" > x < -c("1", "2", "3");
 x; is.character(x); as.numeric(x)
- ## Logical data: TRUE, FALSE, TRUE > x < -1: 10 < 5; x >!x
- ## Return indices for the 'TRUEs' in logical vectorwhich(x)

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Caution !!

On the other hand, if you give a variable the same name as an existing function, R will treat the identifier as a variable if used as a variable, and will treat it as a function when it is used as a function:

c < -2 #typing c yields "2"

c(c,c) #yields a vector containing two 2s.

Better to avoid using 'c' as a variable



Data Objects

```
## Vectors (1D)
> myVec < - 1:10; names(myVec) < - letters[1:10]</p>
> myVec[1:5]; myVec[c(2,4,6,8)]; myVec[c("b", "d", "f")]

## Matrices (2D), Data Frames (2D) and Arrays (≥2D)
> myMA <- matrix(1:30, 3, 10, byrow = T)</li>
> myDF <- data.frame(Col1=1:10, Col2=10:1)</li>
> myDF[1:4, ]; myDF[,c("Col2", "Col1", "Col1")]
## Lists: containers for any object type
> myl < - list(name="Fred", wife="Mary", no childrens</li>
```

- > myL < list(name="Fred", wife="Mary", no.children=3, child.ages=c(4,7,9))
 > myL[[4]][1:2]
- ## Functions: piece of code > myfct < function(arg1, arg2, ...)

General Subsetting Rules

- ## Subsetting by indices > myVec < -1:26; names(myVec) < - LETTERS > myVec[1:4]
- ## Subsetting by same length logical vectors
 myLog < myVec > 10
 - > myVec[myLog]
- ## Subsetting by field names
 myVec[c("R" "K" "M")]
 - > myVec[c("B", "K", "M")]
- ## Special case> iris\$Species

Basic Operators and Calculations

- Comparison operators: ==, !=, <, >, <=, >= ## Example: > 1 == 1
- Logical operators: AND : &, OR : |, NOT : !## Example: > x < -1 : 10; y < -10 : 1> x > y & x > 5
- Calculations: ## Example:
 x + y; sum(x); mean(x), sd(x); sqrt(x)
 apply(matrix(c(1,2,3,4,4,5,6,7,8,9,9,4),3,4), 1, mean)

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- ## Import Data into Rread.delim("myData.csv") ??a < -read.table("myData.txt");
- Extract the portion you want to use, for example, x < -a[,2]
- ## Export Data from R to File
 write.table(myframe, file="myfile.csv", sep="\t", quote=F)

- The unique() function to make vector entries unique > unique(iris\$Sepal.Length); length(unique(iris\$Sepal.Length))
- The table() function counts the occurrences of entries
 table(iris\$Species)
- The aggregate() function computes statistics of data aggregates
 > aggregate(iris[,1:4], by=list(iris\$Species), FUN=mear na rm=T)
- The %in% function returns the intersect between two vectors
 month.name[month.name %in% c("May", "July")]
- The merge() function joins data frames based on a common key column
 - > merge(frame1, frame2, by.x=1, by.y=1, all = TRUE)

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To use a data set available in one of the R packages, install that package (if needed). Load the package into R, using the library() function.

> library(MASS)

Extract the data set you want from that package, using the data() function. In our case, the data set is called JohnsonJohnson.

> data(JohnsonJohnson)

Working with Datasets in R

To use the variable names when working with data, use attach():

- > data(JohnsonJohnson)
- > attach(JohnsonJohnson)

After the variable names have been "attached", to see the variable names, use names():

- > names(JohnsonJohnson) To see the descriptions of the variables, use ?:
- > ?JohnsonJohnson

```
After modifying variables, use detach() and attach() to save the results: # Make a copy of the data set johnson.copy < — JohnsonJohnson; detach(JohnsonJohnson) attach (johnson.copy) # Change the 10 th observation for JohnsonJohnson johnson.copy[10,1] < — 999
```

Caution!!

Avoid using attach() if possible. Many strange things can occur if you accidentally attach the same data frame multiple times, or forget to detach. Instead, you can refer to a variable using \$.

Moral of the story

"attach at your own risk!"