Statistical Computing with R Masters in Data Science 503 (S3) Third Batch, SMS, TU, 2024

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Review Preview

Basics of R

• Chapter from "R for Everyone" book

We will discuss today based on this chapter! • Basics of coding in R

 Chapter from "Hands-on Programming with R" book

 You must read this for the next class!

Basics of R

$$\bullet > 4 * 6 + 5$$

$$\bullet$$
 (4 * 6) + 5

•
$$(4+6)^2 * 5 / 10 + 9 - 1$$

• 5

• R can do Math!

• It follows PEMDAS rule

Parenthesis, Exponents,
 Multiplication, Division, Addition and Subtraction

• BODMAS rule?

Variables in R: assigning and removing

- x <- 2 (preferred)
- x = 2
- 2 -> x
- assign("x", 2)
- rm(x)

- Variable names can contain any combination of alphanumeric characters along with period(.) and underscore (_) e.g. age.group or age_group
- However, they cannot start with a number or an underscore e.g. _age or 5age
- Best practice is to use actual names, usually nouns for variables instead of single letter e.g. age, sex

R is case sensitive

• the Variable <- 17

• Age <- 50

• will give error if we type:

• will be different for:

• The Variable

• age

• THE VARIABLE

AGE

Data Types

• Numeric

• x < -c(1,2,3,4,5,6,7,8,9)

 Type of data can be checked using class() function

 For numeric "class" and "is.numeric" both works:

- class(x)
- is.numeric(x)

Data Types

Integer

• x <- c(1:9) or c(1L:9L)

• Or

• X <- c(1L,2L,3L,4L,5L,6L,7L,8L,9L)

 For integer "class" and "is.numeric" both works:

class(x)"integer"

• is.numeric(x)

TRUE (why?)

R promotes "integers" to "numeric" when needed

```
#Multiply integer by numeric in decimal values
```

- 4L * 2.8
- #Divide integer by integer giving decimal value
- 5L / 2L #Will not promote to numeric here
- 4L * 5L
- #Will also not promote here
- 2L + 4L + 5L

- class(4L)
- class(2.8)
- class(4L * 2.8)
- class(5L)
- class(2L)
- class(5L/2L)
- class(4L * 5L) (?)
- class(2L + 4L + 5L)

Data Types

Character

• x <- "data"

Factor

y <- factor("data")

• X

class(x)

nchar(x)

• y

class(y)

nchar(y)

Factors and attributes in R:

• Factor is used to create and store categorical variable in R like Sex (Male/Female), Blood group (A, B, AB, O) and Blood Rh factor (Positive/Negative) etc. We will need this for supervised learning!

- > gender <- factor(c("male", "female", "female", "male")
- > typeof(gender) #datatype
- > attributes(gender) #Levels and class
- > unclass(gender) #Check how it is stored in R

```
gender <- factor(c("male", "female", "female", "male"))</pre>
typeof(gender)
## "integer"
attributes(gender)
## $levels
## [1] "female" "male"
##
## $class
## [1] "factor"
```

You can see exactly how R is storing your factor with unclass:

```
unclass(gender)
## [1] 2 1 1 2
## attr(,"levels")
## [1] "female" "male"
```

https://rstudio-education.github.io/hopr/r-objects.html#attributes

Data Types

- Date
 - To store date
- POSIXct
 - To store date and time
- Easier manipulation of date and time objects can be accomplished using "lubridate" and "chron" packages

- date1 <- as.Date("2023-03-29")
- date1
- class(date1)
- as.numeric(date1)
- date2 <- as.POSIXct("2023-03-29 06:30")
- date2
- class(date2)
- as.numeric(date2)

Data Types

Logical

• TRUE (=1)

• FALSE (=0)

• TRUE * 5

• FALSE * 5

#Class and check:

• k <- TRUE

• class(k)

• is.logical(k)

Logical Data Types

• 2 == 3 (FALSE)

• 2 > 3 (FALSE)

• 2 != 3 (TRUE)

• 2 >= 3 (FALSE)

• 2 < 3 (TRUE)

"data" == "stats" (FALSE, why?)

• 2 <= 3 (TRUE)

"data" < "stats" (TRUE, why?)

Vectors

• A vector is collection of elements, all of the same type.

Vectors do not have dimension

 Vector in R is like a <u>set</u> with different types of data Vectors in R are not like the mathematical vector

R is a vectorized language

 Column and row vectors can be represented as one-dimensional matrices, however!

Vectors and its operation in R

- x <- c(1,2,3,4,5,6,7,8,9,10)
- x is a vector containing 10 elements
- c stands for "combine"
- It combines multiple elements into a vector
- Shortcut is: 1:10 or 10:1 or -2:3 or 5:-7

- x * 3 #Multiplication by a scalar
- x + 2 #Addition with a scalar
- x 3 #Subtraction with a scalar
- x / 4 # Division by a scalar
- x^2 #Exponentiation by a scalar
- sqrt(x) #Square root

#Two vector of equal length

- x <- 1:10
- y <- -5:4
- x+y
- x-y
- x*y
- x/y
- x^y

- #Check length of the vector
- length(x)
- length(y)
- length(x+y)

#Two vectors of unequal length

- x <- 1:10
- z < -c(1,2)
- X+Z
- Shorter vector get recycled i.e. its elements are repeated, in order, until they have been matched up with every element of the longer vector

#Two vectors of unequal length

- x <- 1:10
- w < -c(1,2,3)
- X+W
- If the longer one is not a multiple of shorter one, warning is given

#Comparing vectors

$$x <= 5$$

- # Using "any" and "all"
- x <- 10:1
- y <- -4:5
- any(x<y)
- all(x<y)

- #Using "nchar"
- nchar(y)

#Assessing individual elements of a vector

#Giving names to a vector

• x[1] retrieves first element of x

#Name value pair

c(One="a", Two="y", Last="r")

 x[1,2] retrieves first and second elements of x

#Create vector then name it

w <- 1:3

names(w) <- c("a", "b", "c")

W

• x[c(1,4)] retrieves?

Calling in-built functions in R

We have already used

- nchar
- length
- as.Date
- as.POSIXct

#We can also use

- mean, var, sd
- round
- factorial

#Getting details of a sensed function aproos("mea")

Missing data in R

R has two types of missing data

• NA

• NULL

 Statistical programs use various techniques to represent missing data such as dash, a period or even the number 99

R uses NA

 NA is represented as just another elements of a vector

NA type missing data in R

- zchar <- c("Hockey", NA, "Cricket")
- nchar(z)
- z <- c(1,2,NA,8,3,NA,3)
- mean(z)
- Missing data can be handled using multiple imputation with mi, mice and Amelia packages

#The "is.na" function tests each element of vector for missingness

• is.na(z)

#The na.rm function with =TRUE argument will remove NA so that we can get values for:

- mean(z, na.rm=TRUE)
- var(z, na.rm=TRUE)
- sd(z, na.rm=TRUE)

NULL type missing data in R

- NULL is the absence of anything
- It is "nothingness"
- Functions can sometimes return NULL and their arguments can be NULL

 NULL is atomical and cannot exist with a vector

- z <- c(1, NULL, 3)
- Z
- [1] 1 3
- is.null(z)
- d <- NULL
- is.null(d)
- is.null(7)

Pipes in R

- A new convention for calling functions in R is the pipe
- The pipe comes from the "magrittr" package BUT starting R 4.0.0 it has in-built pipe now
- x <- 1:10
- mean(x)

- Mean of x with pipe:
- library(magrittr)
- x %>% mean

 It works by taking the value or object on the left hand side of the pipe and inserting it into the first argument of the function that is on the right-hand side of the pipe

Chained pipes in R

 Pipes are most useful when used in a pipeline to chain together a series of function calls

 Given a vector z that contains numbers and NAs, we want to find out how may NAs are present

 Pipes is negligible slower than nesting; but not a bottleneck #Traditionally we do it by nesting

• z <-c(1,2,NA,8,3,NA,3)

sum(is.na(z))

#Pipes, without nesting

• z %>% is.na %>% sum

#Additional argument z %>% mean(na.rm=TRUE)

Advanced data structures in R

Data Frame (data.frame)

- In R data.frame, each column is a vector, each of which is has the same length and same type
- It lets each column holds a different type of data

- x <- 10:1
- y <- -4:5
- q <-c("Hockey", "Football", "Baseball", Kabaddi", "Rugby", "Pingpong", "Basketball", "Tennis", "Cricket", "Volleyball")
- theDF <-data.frame(x, y, q)
- theDF

Advanced data structures in R

theDF <-data.frame(First=x, Second=y, Sport=q)

names(theDF)

names(theDF)[3]

rownames(theDF)

rownames(theDF) <- c("One",
 "Two", "Three", "Four", "Five",
 "Six", "Seven", "Eight", "Nice",
 "Ten")

- Setting them back to generic index
- rownames(theDF) <- NULL
- rownames(theDF)

Advanced data structures in R

#Printing first few rows

head(theDF)

#Printing first seven rows

head(theDF, n=7)

Printing last few rows

tail(theDF)

class(theDF)
 #Structure of data frame by variables

str(theDF)

- theDF[3,2]; theDF[3, 2:3]
- theDF[, 3]; theDF[3,]
- theDF[, c("First", "Sport")]
- theDF[, "Sport", drop=FALSE]

Lists in R

- Often a container is needed to hold arbitrary objects of either the same type or varying types
- R accomplishes this through lists
- They store any number of items of any type; it will be helpful hold large texts with numbers and symbols and mining it

- A list can contain all numerics or characters or a mix of two or data.frame or, recursively, other lists
- Lists are created with list function where each argument to the function becomes an element of the list

Lists in R

#Three element list

• list1 <- list(1,2,3)

#Single element list

• list2 <- list(c(1,2,3)

#Two vector list

• list3 <- list(c(1,2,3), 3:7))

#List with data.frame and vector

list4 <- list(theDF, 1:10)

#Three element list

list5 <- list(theDF, 1:10, list3)

#Names of the list

- names(list5)
- names(list5) <-c("data.frame", "vector", "list")
- names(list5)
- list5
- list6 <- list(TheDataFrame=theDF, TheVector=1:10, TheList=list3)
- names(list6)

Access elements of list

- Use double square brackets
- Specify either the element number or name
- list5[[1]]
- list5[["data.frame"]]
- This allows access to only one element at a time

- #Accessed element manipulation
- lists5[[1]]\$Sport #Sport variable
- lists5[[1]][, "Second"]
- lists5[[1]][, "Second", drop=F]
- length(list5)
- #Adding new element
- list5[[4]] <- 2
- list5[["NewElement"]] <-3:6
- names(list5) & list5

Matrices in R

- This is a similar to a data.frame
- It is rectangular with rows and columns except that every single element must be the same type, most commonly all numerics
- They also act similarly to vectors with elements to element addition, multiplication etc.

- A <- matrix(1:10, nrow=5)
- B <- matrix(21:30, nrow=5)
- C <- matrix(21:40, nrow=2)
- nrow(A)
- ncol(B)
- dim(C)
- A + B; A * B; A B; A = B

Matrix multiplication and names in R

- Matrix multiplication of A and B matrices?
- Number of columns of the left hand matrix to be same as number of rows of right hand matrix
 - A %*% C will work
 - A %*% B will not work
- Both A and B are 5 x 2 matrices so we will transpose B
 - A %*% t(B)

#Column/row names of matrix:

- colnames(A)
- colnames(A) <- c("Left", "Right")
- rownames(A) <- c("1st", "2nd", "3rd", "4th", "5th")
- t(A)
- colnames(B) <- c("First", "Second")
- rownames(B) <- c("One", "Two", "Three", "Four", "Five")

Arrays in R

 An array is essentially a multidimensional vector

- It must be of the same type, and individual elements are accessed in a similar fashion using square brackets
- Very useful for creating and/or replicating multi-way tables in R

- Array: first element is the row index, the second is the column index and remaining elements are for outer dimensions
- theArray <- array(1:12, dim=c(2,3,2))
 - 2 dimensional matrices both with 2 rows and 3 columns
- theArray [1, ,] 1st row of both
- theArray[1, ,1] 1st row of first
- theArray[,1,] 1st column of both

Questions/queries?

• Next session:

R Studio and use for coding, data manipulation and analysis

Thank you!

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