

# Introduction to R programming

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# History of R

R is a dialect (an implementation) of S language.

...

Created in 1991 in new zealand.

# fragile

```
1 # The print function  
2 print('hello world')
```

Assignment operation,

```
1 price <- 30 # assignment operator
2 quantity <- 5
3 income <- price * quantity
```

vectorized

R objects as used before are really vectors.

```
1 output_msg <- "profit"
```

...

The object is character vector in R, the first is profit

[1] 10 a vector whose first element is the number ten  
We'll create a sequence with colon(:) operator.

```
1 ito <- 1:10
```

# R objects

atomic classes of objects:

- character
- integer
- numeric (float)
- complex
- logical ( booleans)

# keep information

vector - list

vector

each one of its elements are of the same class

list

can keep information of all kind



```
1 vector() # i don't have idea then;  
2 help("vector")
```

# Numbers

in R are generally treated as numeric objects ( double precision real numbers).

```
1 123L # explicit integer
```

## Special

- inf (infinity)
- NaN (Not A Number)

# Attributes

names, dimensions, class, lenght and another.

```
1 help("attributes")  
2 attributes(df) # rows and columns
```

# Vector - list

the `c()` (concatenate) function create **vectors** of objects.

```
1 df <- c(0.9, 0.87, 0.95) # numeric object
2 booleans <- c(TRUE, FALSE, T, F) # logical
```

## coercion

if you mix different kind of objects in a vector then occur coercion casting all the object to the same class, for instance `c('1', 2)` caste 2 to '2'.

# Type

In programming languages define the type of data is very important thus:

```
1 bool <- c(True, F)
2 class(bool)
3 # explicit conversion
4 as.numerical(bool)
5 as.character(bool)
6 binaries <- c(1,0,1)
7 as.logical(binaries)
8 vowels <- c('a','e','i')
9 as.logical(vowels)
```

```
1 data <- list('True', FALSE, 0 , 1.0)
```

Then the first element is a vector of characters, then second is a vector (logical) with the element FALSE, etc...

# Matrices

```
1 mat <- matrix(nrow=3, col= 2)
2 attributes(mat)
3 dim(mat)
```

```
1 matrix(1:4,nrow=2,ncol=2) # column-wise (left-upper
  )
2 data <- 1:4
3 dim(data) <- c(2,2)
4 print(data)
```

# cbind and rbind

```
1 x <- 5:10  
2 y <- 50:55  
3 cbind(x,y)  
4 rbind(x,y)
```



# Factors

Useful in categorical data, ordinal or cardinal, think in the problem itself that carry out pass categorical to numbers low=1, medium =2, high=3.

```
1 var <- factor(c('high', 'medium', 'low', 'low'))  
2 table(var)  
3 unclass(var) # how the factors are represented by R
```

# Baseline factor

```
1 var <- factor(c('cs','math','cs' ,'economics' ,  
2 'math','cs'),  
3 leves=c('math', 'cs', 'economics'))
```

The first level is used as baseline level in some statistical procedures.

# Missing values

## NA and NaN

- NA (Not Available) is general, a dont have information about missingness is a class!.
- NaN (Not a Number) arise from undefined mathematical operations as  $\frac{a}{0}$

```
1 is.nan()  
2 is.na()
```

a NaN (related to numerical) is also a NA but not viceversa.

The following is very important handled missing values.

```
1 data <- c(1,2,4, NA, NaN)
2 is.nan(data)
3 is.na(data)
4 data <- c('1', 2, NaN, NA)
5 is.nan(data)
6 is.na(data)
```

# DataFrame

Tabular data, important points:

- unlike matrices can handle different types of data (like list)
- could be converted to matrices as `df.matrix()`

Create a dataframe to test:

```
1 df <- data.frame(var1=c('red','red','blue','blue'),  
2   ,  
3   var2=c(T, T, F, F), var3=c(1,1,0,0))  
4 # Describe the data  
5 nrow(df)  
6 ncol(df)
```

# Names

```
1 booleans <- c(T,F,T,F)
2 #Give name to each element...
3 names booleans <- c("yes", 'not', 'si', 'no')
4 names booleans
```

# name and indexing

```
1 data <- list(age= 40, id = 0, ibm =23)
2 data$age
3
4 mat <- matrix(1:6, ncol =3, nrow=2)
5 dimnames(mat)<-list(c('A', 'B'),
6 c('col1', 'col2', 'col3'))
7 # try
8 rownames(mat) <- c("A", "B")
9 colnames(mat) <- c('col1', 'col2', 'col3')
10 # indexing
11 mat['A', 'col3']
```

# read and write common

- `read.csv()`
- `read.table()`
- `wite.table()`
- `save()`

Remeber that is very important get information in documentation  
`help("read.table")` also `read.csv()` is identical to `read.table` but if  
separator by deafulst is the comma(,) character. important also remark  
that R skip lines that begin with `#` character.



# subsetting

- [ hold the class
- [[ dont hold the class (think in lists)
- \$ extract elemens (named) in list or dataframe

```
1 data <- c('Blue', 'Green', 'Red')
2 data[1] # take the first element (numerical
  index)
3 data[1:2] # from first to second
4 data[c(T,F,T)] #logical index
5 data[data=='Green'] # two equals signs (=).
6 data[data=='Green' | data=='Blue']
7 data <- c(5,10,11,2, 7)
8 data[data>6]
```

# Research

- Research about the function paste
- Research about the function cat

# subsetting

## list

```
1 data <- list(var1= 5:10, var2 = 15:20)
2 data[1] + 1 # error is a list!
3 data[[1]] + 1 # numeric vector
4 class(data$var1)
5 class(data['var1'])
6 class(data[['var1']])
```

is important take care, given that could be produce errors!! Single Square Bracket always return an object of the same class of the original object. Double Square bracket with name is equal to dollar sign.

# extract multiple elements

Uses single bracket operator

```
1 data <- list(a=10:30)
2 data$b <- 50:80
3 data$c <- 100:130
4 data[c("a","b")] # try with numerical index
5 # dollar sign not allow computed index for instance
  ;
6 column <- 'b'
7 data$column # return NULL
8 data[[column]] #execute!
9 data$b # execute!
```

subsetting nested elements.

```
1 data <- list(col1=list(T,F,T),  
2 col2=list('Red','Blue'))  
3 data[[c(1,2)]]  
4 data[[1]][[3]]
```

# Subsetting matrix

$M_{ij} = M[i,j]$  where  $i$  is rows and  $j$  columnns.

```
1 mat <- matrix(1:10, 2,5)
2 mat[1,4]
3 mat[1,]
4 mat[,4]
5 class(mat[1,3])
6 class(mat[1,3, drop=F])
7 # A rule is broken [] only return the same class
8 class(mat[1,])
9 class(mat[1,,drop=F])
```

# Removing missing values

```
1 bools <- c(T,F)
2 !bools
3 data <- c(1, 0 , 0.5, NaN, NA)
4 data[!is.na(data)]
5 x <- c(NA, 3, NaN, 5, 10)
6 y <- c(10, 11, 4, Nan, 4)
7 complete.cases(x,y)
8 # assign to both arrays to extract the complete
   data.
```

# in DataFrame

```
1 data <- data.frame(  
2   age = c(32, NaN, 31, NaN),  
3   score = c(NaN, 70, 81, 90))  
4 index <- complete.cases(data)  
5 data[index,]
```



# Vectorized operations

```
1 x <- 10:15
2 y <- 0:5
3 x<y
4 x+y
5 x*y
6 x/y
```

```
1 mat <- matrix(1:9, 3,3)
2 mat2 <- matrix(5:13, 3, 3)
3 mat*mat2 # Element wise multiplication
4 mat %*% mat2 #Matrix multiplication
```

# Control structures

```
1  if (condition)
2  {
3  statement 1
4  } else{
5  statement 2 (when cond is FALSE)
6  }
```

# Example

```
1 price <- 30
2 quantity <- 50
3 income <- price * quantity
4 if (income>10){
5     print("Excellent!")
6 } else{
7     print(':(')
8 }
```

# else if

```
1  if (condition){  
2    Statement 1  
3  } else if (condition) {  
4    Statement alternative  
5  } else{  
6    else statement  
7  }
```

```
1 price <- 10
2 quantity <- 1
3 income <- price * quantity
4 if (income>10){
5     print("Excellent!")
6 } else if (income == 10){
7     print('equilibrium')
8 } else {
9     print(':(')
10 }
```

# loop

for

```
1 for (num in 1:10){  
2   print(num)  
3 }
```

# Examples

```
1 x <- c(10, 0, 5)
2 for (i in x){
3   print(i)
4 }
5
6 for (i in seq_along(x) ){
7   print(i)
8 }
9
10 for (i in 1:length(x)){
11   print(x[i] + 1)
12 }
```



# Nested loop

```
1 for (i in 1:2) {  
2   for (j in 3:5){  
3     print(c(i,j))  
4   }}
```

As exercise nested a matrix! `ncol` and `nrow`.

# Example

...

See simulation mean

See simulation estimator counter

See simulation

# while loop

```
1 acum <- 1
2 while (acum < 5) {
3   print(acum)
4   acum <- acum + 1
5 }
```

# OR and AND

- `&&` logical AND operator
- `&` element wise logical operator used in vectors to test AND
- `||` logical OR operator
- `|` element wise logical operator used in vectors to test OR

```
1 T && F
2 T || F
3 x <- c(T, T, F, F)
4 y <- c(T, F, T, F)
5 x&y
6 x|y
```

```
1 # Search a number in the vector!
2 array <- c(10,40, 50, 1)
3 flag = FALSE
4 to_search = 1
5 index <- 1
6 while (index <= length(array) && flag==FALSE){
7   print(array[[index]])
8   if (array[[index]] == to_search){
9     flag <- TRUE
10  }
11  index <- index + 1
12 }
13 flag
```

The conditions are evaluated from left to right.

# Repeat, Next, Break

This topics are to research...

- `repeat` is a infinite loop whose exit condition is `break` statement.
- `next` skip a loop iteration (make an example with `NAN`)

# Functions

```
1 function_name <- function(arguments){  
2   function statement  
3 }
```

```
1 future_val <- function(initial_val, r, n){  
2   initial_val*(1 + r)^n #you could uses return  
3 }
```

# Excercise

...

write a function that compute the number of periods ( $n$ )



# Exercise

...

write a function that compute the number of periods ( $n$ )

```
1 periods <- function(lambda, r){  
2   return(log(lambda)/log(r +1 ))  
3 }
```

Pi simulation

(Click here)

# Practical

See lab

# notes

Default arguments works like in python the `return` statement is necessary R return the last statement in the function body. The arguments work as in python by name or position.  
always check the arguments

```
args(print)
```

Important **lazy evaluation** evaluate arguments if they are needed.

```
1 # lazy evaluation
2 power <- function(a,b,c){
3   a^(b)
4 }
5 power(10,2)
6 # error not is arised
```

# Important things in functions

## ... argument (ellipsis)

is important be careful using ellipsis (a undefined number of parameters).

```
1 suma <- function(...) {  
2   args <- list(...)  
3   acum <- 0  
4   for (arg in args){  
5     acum <- acum + arg}  
6   acum  
7 }  
8 suma(1,2,3,4,5,6,7,8,9,10)  
9 print(10*11/2)
```

ellipsis working as wrapped!

```
1 mean <- function(..., message='the mean is'){
2   print(message)
3   suma(...) / length(list(...))
4 }
5 mean(1,3,4) # arguments appear after of ellipses
   then they must be named explicitly.
```

# Scoping rules

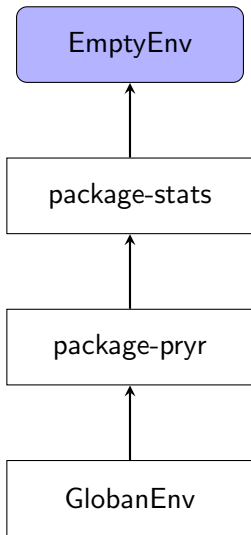
The scope is the place where a variables is defined and its visibility.

- Lexical scoping
- Dynamic scoping

in R are exist *free variables*.



Figure: Enviroment tree



First the symbol: value is searched in the global environment, and after in each one of the packages listed in the following command.

```
1 search()
```

This loads all packages that are loaded currently in R. Global environment is always first and base the last, when you load a package the namespace occupies the second place and all others shift down a position.

- R functions are stored and searched in the same way of other object, therefore are searched in the environment tree.

# Anonymous functions

```
1 sapply(c(1,2,3,4), function(x) x^2)
```

## << – operator

assign the variable to the global environment (not local where was defined).

```
1 local_var <- function(a){
2     result <- a^2
3     result
4 }
5 local_var(10)
6 result # object 'result' not found!
7
8 global_var <- function(a){
9     result <<- a^2
10    result
11 }
12 global_var(10)
13 result
```

# Closures

## Analyze

```
1 counter <- function(){
2   counter <- 0
3   function(){
4     counter <-< counter + 1
5     counter
6   }
7 }
8 acum <- counter()
9 acum()
10 acum()
11 acum()
```

# Closures

- A function defined inside of another captures information from the outer function (its environment)
- The inner function can access and modify the variable count even though the function has already finished execution.

# closures

```
1 experiment <- function(sample_size){  
2   mean <- 0  
3   function(){  
4     mean <- mean(rnorm(sample_size))  
5     mean  
6   }  
7 }  
8 low_size <- experiment(10)  
9 medium_size <- experiment(35)  
10 for (i in 1:5){  
11   print(c(low_size(), medium_size()))  
12 }
```

Extend the function! to confidence intervals.



# Closure

```
1 pi_estimator <- function(trials){
2   result <- function(){
3     acum <- 0
4     for (i in 1:trials){
5       if (sum(runif(2,-1,1)**2) <= 1){
6         acum <- acum + 1
7       }}
8     acum/trials*4
9   }}
10 small_trials <- pi_estimator(1000)
11 big_trials <- pi_estimator(1000000)
12 print(c(small_trials(), big_trials()))
```

# Closures

Given lexical scoping `low_size` and `medium_size` can reference its environment definition!! and therefore the sample size defined 10 and 35 respectively.

```
1 ls(environment(low_size))  
2 get('trials', environment(small_trials)) # knowing  
  its environment!
```

# Exercise

```
1 var <- 1
2 call <- function() var
3 change <- function() {
4     var <- var + 100
5     #var <-< var + 100
6     call()
7 }
8 change() # the value is?
```

# Important point!

define and after call a function in global environment resemble to dynamic scoping given that calling and defining environment are the same (add an example).

Analyze

```
1 y <- 3
2 f <- function(x){
3   y <- 2
4   y^2 + eval(expression(g(x)),envir = parent.frame())
      # comment this line
5 #y^2 + g(x) # uncomment this line
6 }
7 g <- function(x){
8   x^y
9 }
10 f(2)
```

# Dates and time

- Dates are stores as the number of days since 01-01-1970
- Times as the number of seconds from 01-01-1970

```
1 date_ <- as.Date('1971-01-01')  
2 unclass(date_)
```

represented by POSIXct or POSIXlt

```
1 now <- as.POSIXlt(Sys.Date())
2 names(unclass(now))
3 now$mday
4 Sys.Date() # is in POSIXct format
5 # The number of seconds since 01-01-1970.
```

# strptime function

Data is written in a different format

```
1 date_str <- c("December 29, 2024 9:10")
2 date_ <- strptime(date_str, "%B %d, %Y %H:%M")
3 date_
4 start <- as.Date("2021-01-03")
5 finish <- as.Date("2022-02-01")
6 finish - start # take care with the compatibility
```

# Research time

- apply
- sapply
- lapply
- tapply
- mapply



# Excercise Write standard deviation function

# Excercise Write standard deviation function

```
1 std_ <- function(..., ddof=0){
2   args <- as.numeric(list(...))
3   acum <- 0
4   mean_ <- mean(args)
5   for (arg in args){
6     acum <- acum + (arg - mean_)**2
7   }
8   (acum / (length(args) - ddof))^0.5
9 }
10 print(std_(10,3,1,4, ddof=1))
11 print(sd(c(10,3,1,4)))
```

# lapply

As an argument, receive a list (or cast it to one) and a function that receive as input each element of the list.

```
1 lapply(1:4, function(x){x^2})
```

```
1 lapply # ellipses represent arguments in the  
  function.  
2 data <- list(list(10,3,1),  
3 list(10,3,1,4),  
4 list(10,3))  
5 lapply(data, function(x){sd(unlist(x))})  
6 lapply(data, function(x){do.call(std_, x)}) #unpack  
7 lapply(data, function(x) { do.call(std_,  
8 c(x, ddof = 1)) })
```

List in R are "similar" to dictionary

```
1 lapply(list(col1=1:5, col2=10:15),  
2 mean, trim=0, na.rm=T) # note that are passed mean  
    function arguments!
```

# sapply

simplify the result of lapply

```
1 lapply(1:4, function(x) x**2)
2 sapply(1:4, function(x) x**2) # a vector is
   returned!
```

# apply

apply usually used to apply a function over a row or column of a matrix. The first argument is data, the second is the axis(1 or 2) and the third the function.

```
1 mat <- matrix(c(0, 1, 10,  
2                 1, 0, 3,  
3                 5, 0, 0),  
4                 nrow = 3, byrow = TRUE)  
5 apply(mat, 1, sum) # sum all values in each row  
6 apply(mat, 2, sum) # sum all values in each column
```

# Research

- rowSums
- colMeans
- ...

how works

- quantile

- mapply
- tapply

# split

Split objects into groups determined by factors.

```
1 data <- c(rnorm(5, 1), rnorm(5, 45), rnorm(5, 1000)
  )
2 f <- gl(3, 5) # three groups with five observations
  each one
3 split(data, f) # three groups
4 lapply(split(data,f), mean)
```



# apply - split

```
1 df <- data.frame(income=c(10, 14, 15, 23.2, 21.3),  
2 exp = c(6,8,3, 2, 4),  
3 group=c('math','math','math','cs','cs'))  
4 split(df, df$group)  
5 lapply(split(df, df$group), function(x) colMeans(x  
    [, c('income', 'exp')]))
```

## more than one factor

```
1 df <- data.frame(income=c(10, 14, 15, 23.2, 21.3,
2   21.4),
3   exp = c(6,8,3, 2, 4,5),
4   group=c('math','math','math','cs','cs', 'cs'),
5   sex = c('female', 'male', 'male', 'female', 'female',
6     'male'))
7
8 lapply(split(df, interaction(df$group, df$sex)),
9   function(x) colMeans(x[, c('income', 'exp')]))
10
11 lapply(split(df, list(df$group, df$sex)), function(
12   x) colMeans(x[, c('income', 'exp')])) #split
13   allow us uses list
```

# Debug

research about

- `traceback`
- `debug`
- `browser`
- `trace`
- `recover`

# str

`str()` help us to visualize the data!

```
1 data <- list(1,2,3,4,5,6,7,8)
2 str(data)
3 str(df)
```

# Simulation

```
1 set.seed(1) # uses seeds!  
2 par(mar = c(4, 4, 1, 1))  
3 domain <- seq(-3, 3 , length.out = 100)  
4 range <- dnorm(domain)  
5 plot(domain, range, type='l')  
6 cumulative_range <- pnorm(domain)  
7 plot(domain, cumulative_range, type='l')
```

# Research statistical distributions

```
help("distribution")
```

- rpois
- dt
- dnbinom
- ...

## prefixes

- d(ensity)
- r(andom number generator)
- p(robability - cumulative distribution)
- q(uantile function)

# linear reg

```
1 linear <- function(b0, b1, x, u){  
2   b0 + b1*x + u  
3 }  
4 sample_size <- 500  
5 b0 = 1  
6 b1 = 3.4  
7 x <- rnorm(sample_size, 10, 3)  
8 u <- rnorm(sample_size, 0, 1)  
9 y <- linear(b0, b1, x, u)  
10 lm(y ~ x )
```

# Sample

```
1 data <- c(100, 20, 5, 71, 89, 32, 121)
2 sample(data,2)
3 sample(data, 2, replace=T)
```