(A bit of) Advanced R

Part 1 - R-base programming

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Outline

- Control Statements
- 2 Functions
- 3 Functionals

References

Advanced R (Wickham, 2014), http://adv-r.had.co.nz/



A Language and Environment for Statistical Computing (R Core Team, 2017), https://www.R-project.org/



→ Many ideas/examples inspired/stolen there.

Prerequisites

Data Structure in base R

- 1 Atomic vector (integer, double, logical, character)
- Recursive vector (list)
- S Factors
- Matrices and array
- Data Frame
- --- Creation, Basic Operation, Manipulation, Representation

Resources

- Advanced R, chapters I.2, I.3 (Wickham, 2014, http://adv-r.had.co.nz/)
- An introduction to R programming http://julien.cremeriefamily.info/teachings_L3BI_ISV51.html

Developement environment I

The Rstudio API

- A full API with code, interpreter, workspace and plots
- Package developement and external code integration are easier
- Notebooks integration with Rmarkdown
- Interface with github → required tool for efficent development in R

My favorites shortcuts

- ctrl + return: execute current selection in console
- ctrl + 1/2/3/4: naviguate between panels
- ctrl + down/up: naviguate between tabs
- ctrl + shift + k: knit current doccument

Developement environment II

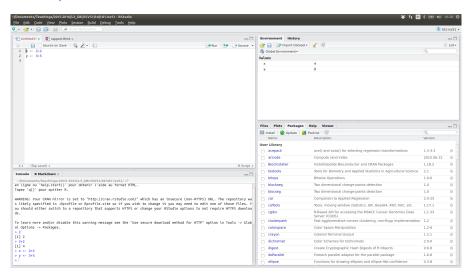


Figure 1: Screenshot of the Rstudio API

Outline

- 1 Control Statements
- 2 Functions
- 3 Functionals

Grouped expressions

Syntax

```
{expr_1; expr_2; ...; expr_n }
{
    expr_1
    ...
    expr_n
}
```

Remarks

- the last value is sent back
- un group statement can be passed to a function

Grouped expressions: examples

Example 1

```
expr1 <- {a<-3; b<-5; a*b}
expr1
## [1] 15</pre>
```

Example 2

```
tmp <- 12
expr2 <- {a<-3; b<-5; tmp<-a*b+tmp}</pre>
```

```
expr2
## [1] 27
tmp
## [1] 27
```

Grouped expressions: examples

Example 1

```
expr1 <- {a<-3; b<-5; a*b}
expr1
## [1] 15</pre>
```

Example 2

```
tmp <- 12
expr2 <- {a<-3; b<-5; tmp<-a*b+tmp}</pre>
```

expr2

[1] 27

tmp

[1] 27

Grouped expressions: examples

Example 1

```
expr1 <- {a<-3; b<-5; a*b}
expr1
## [1] 15</pre>
```

Example 2

```
tmp <- 12
expr2 <- {a<-3; b<-5; tmp<-a*b+tmp}

expr2
## [1] 27
tmp
## [1] 27</pre>
```

Conditional statements: if, if/else, if else

Standard syntax

```
if (condition) {
   expr_1
} else {
   expr_2
}
```

Vectorial form

```
ifelse(condition, a, b)
```

Remarks

- condition is logical, so use &, |, !, etc.
- else is optional
- elseif allows imbricating statements

Conditional statements: example

```
partiel <- 11
DS <- 14
if (partiel > 6 & mean(DS,partiel) > 10) {
    cat("\nregu(e).")
} else {
    cat("\nrecalé(e).")
}
```

reçu(e). Exercice

##

Use the vectorial ifelse to send the full vector of results

```
partiel <- c(11,5,6,12,9,8,14)
DS <- c(14,16,12,12,19,12,7)
```

```
ifelse(partiel > 6 & rowMeans(cbind(DS,partiel)) > 10, "reçu(e)", "recalê(e)")  
## [1] "reçu(e)" "recalê(e)" "recalê(e)" "reçu(e)" "reçu(e)" "reçu(e)" "reçu(e)"  
## [7] "reçu(e)"
```

Conditional statements: example

```
partiel <- 11
DS <- 14
if (partiel > 6 & mean(DS,partiel) > 10) {
    cat("\nrequ(e).")
} else {
    cat("\nrecalé(e).")
}
```

reçu(e).

Use the vectorial ifelse to send the full vector of results

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partiel <- c(11,5,6,12,9,8,14)
DS <- c(14,16,12,12,19,12,7)

ifelse(partiel > 6 & rowMeans(cbind(DS,partiel)) > 10, "regu(e)", "recalé(e)")

## [1] "regu(e)" "recalé(e)" "recalé(e)" "reçu(e)" "reçu(e)" "recalé(e)"
## [7] "regu(e)"
```

Conditional statement: switch

Syntax

```
switch (expr,
    expr_1 = do_1,
    ...,
    expr_n = do_n,
    do_default
)
```

Remarks

- expr is either an integer or a character
- ullet if an integer with value i, the ith expression do_i is evaluated
- if a string the expression do_i so that expr == expr_i is evaluated

switch: examples

integer form

```
expr <- 2
switch(expr, cat("My value is 1"), cat("My value is 2"))
## My value is 2
expr <- 3
switch(expr, cat("My value is 2"), cat("My value is 2"))</pre>
```

character form

loop statement: for

Syntax

```
for (var in set) {
  expr(var)
}
for (var in set) # avoid this syntax!!
  expr(var)
for (var in set) expr(var)
```

Remarks

- var is the incremented variable
- set is a vector of the successive values
- generally slow compared to matricial/vectorize operation

for loop: examples

Example: C/C++ like

```
for (i in sample(1:5)) cat(i)
## 54231
v <- numeric(7)
for (i in seq_along(v)) v[i] <- i*3</pre>
```

Exercice:

Use a for loop to display the colnames of the data frame iris which are not a factor, by completing the follwoing piece of code

```
data(iris)
for (nom in colnames(iris)) {
   if (!is.factor(iris[,nom])) cat("",nom)
}
### Sepal.Length Sepal.Width Petal.Length Petal.Width
A more R-style way to do that is
cat(colnames(iris)[!sapply(iris, is.factor)])
```

for loop: examples

Example: C/C++ like

```
for (i in sample(1:5)) cat(i)
## 54231
v <- numeric(7)
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for (nom in colnames(iris)) {
   if (!is.factor(iris[,nom])) cat("",nom)
}
```

Sepal.Length Sepal.Width Petal.Length Petal.Width

A more R-style way to do that is

cat(colnames(iris)[!sapply(iris, is.factor)])

Sepal.Length Sepal.Width Petal.Length Petal.Width

for loop: examples

Example: C/C++ like

```
for (i in sample(1:5)) cat(i)
## 54231
v <- numeric(7)
for (i in seq_along(v)) v[i] <- i*3</pre>
```

Exercice:

Use a for loop to display the colnames of the data frame iris which are not a factor, by completing the following piece of code

```
data(iris)
for (nom in colnames(iris)) {
   if (!is.factor(iris[,nom])) cat("",nom)
}
```

Sepal.Length Sepal.Width Petal.Length Petal.Width

A more R-style way to do that is

```
cat(colnames(iris)[!sapply(iris, is.factor)])
```

```
## Sepal.Length Sepal.Width Petal.Length Petal.Width
```

Loop statement: while, repeat

Syntax

```
while (condition) {
  expr
}
repeat {
  expr
}
```

Remarks

- avoid imbrication (slow)
- can be interrupted/controlled with with break/next

```
repeat {
    expr
    if (condition) {break}
}
while (condition1) {
    expr_1
    if (condition2) {next}
    expr_2
}
```

Outline

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Functions definition

Syntax

```
my_func <- function(arg1,arg2, ...) {
  expression
}</pre>
```

Remarks

- The last value of the expression is returned
- One must use a list to send back several objects
- return() is used only when you need to send a value at an early stage in the expression
- In R, functions are object like any others and can manipulated as such

Function components I

\$use

Most of functions have three parts

- the body() (code inside the function)
- the formals() (list of arguments)
- the environment() (a set of bindings between symbols and objects, i.e, a place to store variables)

```
environment(var)

## <environment: namespace:stats>
formals(var)

## $x

## ##

## ## $y

## NULL

## ## $na.rm

## [1] FALSE
```

Function components II

body(var)

```
## {
##
       if (missing(use))
           use <- if (na.rm)
##
##
               "na.or.complete"
##
           else "everything"
##
       na.method <- pmatch(use, c("all.obs", "complete.obs", "pairwise.complete.obs",</pre>
           "everything", "na.or.complete"))
##
##
       if (is.na(na.method))
##
           stop("invalid 'use' argument")
       if (is.data.frame(x))
##
           x <- as.matrix(x)
##
##
       else stopifnot(is.atomic(x))
##
       if (is.data.frame(y))
##
           y <- as.matrix(y)
##
       else stopifnot(is.atomic(y))
##
       .Call(C cov, x, y, na.method, FALSE)
## }
```

Lexical Scoping I

Definition

Set of rule that governs how R looks up the value of a symbol

Name masking

If a name is not defined inside a function, R looks a level up

```
y <- 2
func <- function(x) c(x,y)
func(4)
## [1] 4 2</pre>
```

This applies to funciton defined in another function

```
x <- 2
func <- function(y) {
   sub_func <- function(z) c(x,y,z)
   sub_func(5)
}
func(3)
## [1] 2 3 5</pre>
```

Lexical Scoping II

function vs variable

R makes the distinction between variable and function names

```
n <- function(x) x/2
f <- function() {n <- 10 ; n(n)}
f()
## [1] 5</pre>
```

Fresh star

[1] E

An environement is created at each time a function is called

```
f <- function() {
    a <- ifelse(exists("a"), a + 1, 1)
    print(a)
}
f()
## [1] 1
f()
## [1] 1
a <- 4
f()
## [1] 5
f()</pre>
```

Function arguments

Calling function

Arguments can be specified 1. by name 2. by partial name 3. by position Here are some stupid (but correct) call to mean(x=,trim=,na.rm=)

```
mean(1:10, n = T)
## [1] 5.5
mean(1:10, , FALSE)
## [1] 5.5
mean(1:10, 0.05, FALSE)
## [1] 5.5
mean(, TRUE, x = c(1:10, NA))
## [1] 5.5
```

Exercice

Clarify the following function calls

```
x <- sample(replace = TRUE, 20, x = c(1:10, NA))
y <- runif(min = 0, max = 1, 20)
cor(m = "k", y = y, u = "p", x = x)</pre>
```

Default arguments

Arguments can have default values in R

```
f <- function(a = 1, b = 2) c(a,b)
f()
## [1] 1 2</pre>
```

The missing function

```
You can check wether an argument was passed or not with missing f <- function(a = 1, b = 2) c(missing(a),missing(b)) f(a)

## [1] FALSE TRUE

Hence, you can assign a default value a posteriori

f <- function(a, b = 2) {
  if (missing(a)) a <- 3
  c(a, b)
  }
  f(a)
```

Default arguments

Arguments can have default values in R

```
f <- function(a = 1, b = 2) c(a,b)
f()
## [1] 1 2</pre>
```

The missing function

You can check wether an argument was passed or not with missing:

```
f <- function(a = 1, b = 2) c(missing(a),missing(b))
f(a)
## [1] FALSE TRUE</pre>
```

Hence, you can assign a default value a posteriori

```
f <- function(a, b = 2) {
   if (missing(a)) a <- 3
   c(a, b)
}
f(a)
## [1] 4 2</pre>
```

Lazy evaluation

[1] 4 12

```
Arguments are evaluated only if they are used, which is known as "lazy evaluation"
f \leftarrow function(a = 1, b = 4*a) c(a,b)
f()
## [1] 1 4
f(43)
## [1] 43 172
Even better (or worse...)
f <- function(a = 1, b = d) {
  d \leftarrow 4 + 2 * a; c(a,b)
f()
## [1] 1 6
f(4)
```

The ... argument

The argument ... matches any argument not otherwise matched

- useful when collecting argument to call another function
- do not need to precify the name ofd required argument
- the counterpart is that any misspelled argument is passed to ... and show no warning

Example: plot

Many argument in plot are passed to the par function that manages the graphical paramters:

```
plot(1:5, col = "red")
plot(1:5, llty = "dotted")
```

Capturing ...

list() can be used to easily captured arguments passed with ...

```
f <- function(...) names(list(...))
f(a = 1, b = 2)
## [1] "a" "b"
```

Calling function with a list

The do.call function constructs and executes a function call from a name or a function and a list of arguments to be passed to it:

```
do.call(mean, list(x = 1:10, trim = 0.05, na.rm = FALSE))
## [1] 5.5

Exercice
Suppose the outputs of 100 simulations are stored in a list like that class(res)
## [1] *list*
res[[1]]
## method mse timing
## 1 lasso 0.8222197 0.06823225
## 2 ridge 0.8912516 0.91282404
## 3 bayes 0.9756149 123 53501949
```

length(res

[1] 100

How would you store them in a single data frame?

```
head(do.call(rbind, res), 2)
```

```
# method mse timing
# 1 lasso 0.8222197 0.05823235
```

Calling function with a list

The do.call function constructs and executes a function call from a name or a function and a list of arguments to be passed to it:

```
do.call(mean, list(x = 1:10, trim = 0.05, na.rm = FALSE))
## [1] 5.5
```

Exercice

Suppose the outputs of 100 simulations are stored in a list like that

```
class(res)
## [1] "list"
res[[1]]
## method mse timing
## 1 lasso 0.8222197  0.05823235
## 2 ridge 0.6912516  0.91282404
## 3 bayes 0.9756149 123.53501949
length(res)
## [1] 100
```

How would you store them in a single data frame?

```
## method mse timing
## 1 lasso 0.8222197 0.05823235
## 2 ridge 0.6912516 0.91282404
```

Calling function with a list

The do.call function constructs and executes a function call from a name or a function and a list of arguments to be passed to it:

```
do.call(mean, list(x = 1:10, trim = 0.05, na.rm = FALSE))
## [1] 5.5
```

Exercice

```
Suppose the outputs of 100 simulations are stored in a list like that

class(res)

## [1] "list"

res[[1]]

## method mse timing
## 1 lasso 0.8222197 0.05823235

## 2 ridge 0.6912516 0.91282404
## 3 bayes 0.9756149 123.53501949

length(res)

## [1] 100
```

How would you store them in a single data frame?

```
head(do.call(rbind, res), 2)
## method mse timing
```

```
## 1 lasso 0.8222197 0.05823235
## 2 ridge 0.6912516 0.91282404
```

Infix functions

Definition

Infix function (contrary to prefix functions) are function where the name comes between the argument (like "-" or "+").

```
R comes with the following infix functions predefined: \%, \%*\%, \%*\%, \%in\%, \%o\%, \%x\%, :, ::, $::, $, @, ^, *, /, +, -, >, >=, <, <=, ==, !=, !, &, &&, |, ||, ~, <-, <<-
```

Example

Can be use to define operator

```
"%+%" <- function(x,y) paste(x,y)
"Université" %+% "Paris" %+% "Dauphine"
## [1] "Université Paris Dauphine"</pre>
```

Exercice

Create infix functions for intersection, union and setdiff and test it on simple vectors.

Primitive functions: definition

NULL.

- Primitive functions are functions from the base package that call C code directly
- Primitive functions do not contain R code, as so

```
## function (..., na.rm = FALSE) .Primitive("sum")
formals(sum)

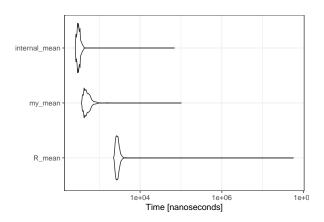
## NULL
body(sum)

## NULL
environment(sum)
```

Primitive functions: performance

Function defined internally (either with .Primitive either called via .Internal) are sometimes incredibly faster (written in C), but cannot by called directly in packages submitted to CRAN.

```
x <- rnorm(100)
R_mean <- function(x) mean(x)
my_mean <- function(x) sum(x)/length(x)
internal_mean <- function(x) .Internal(mean(x))</pre>
```



Compile your functions with base::compiler I

The R byte code compiler

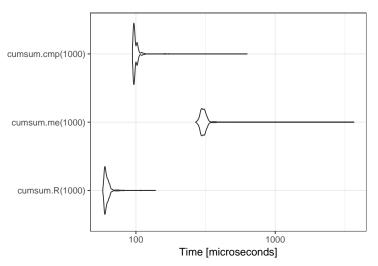
The function cmpfun compiles the body of your function and returns a new function with the same formals and the body replaced by the compiled body expression.

```
cumsum.R <- function(n) {
    x <- rnorm(n)
    cumsum.x
}

cumsum.me <- function(n) {
    x <- rnorm(n)
    res <- 0
    for (i in 1:length(x))
    res <- res + x[i]
    res
}

cumsum.cmp <- compiler::cmpfun(cumsum.me)</pre>
```

Compile your functions with base::compiler II



- If you cannot avoid a loop, you will save some time
- Can be set automatically with compiler::enableJIT(3)

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Functional for matrix and arrays: apply

Definition

Applies a function along a dimension of an array (row/columns of matrix).

```
apply(array, dim, FUN, ...)
```

Example

```
mat <- matrix(1:6, 2, 3)
apply(mat, 2, max)

## [1] 2 4 6

arr <- array(1:12, c(2,3,2))
apply(arr, 3, colMeans)

## [,1] [,2]
## [1,] 1.5 7.5
## [2,] 3.5 9.5
## [2,] 5.5 11.5</pre>
```

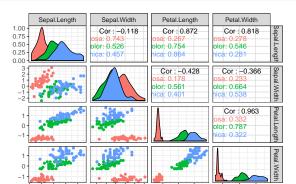
Other array functionals: sweep

Definition

'Sweep out' a summary statistic along a dimension of an array for a given function sweep(array, dim, stat, FUN, ...)

Example

```
out <- sweep(iris[, 1:4], 2, colMeans(iris[, 1:4]), "-") ## center
out <- sweep(out, 2, sqrt(colSums(out^2)/(nrow(out) - 1)), "/") ## scale
iris_sc <- data.frame(out, Species = iris$Species)
GGally::ggpairs(iris_sc, columns = 1:4, ggplot2::aes(colour = Species))</pre>
```



Other array functionals: outer

Definition

```
outer(array1, array2, FUN, ...)
```

Example

The more basic example is the kronecker product, but FUN can be anything!

```
outer(1:3, 1:4, "*")
## [,1] [,2] [,3] [,4]
## [2,] 2 4 6 8
## [3,] 3 6 9 12

1:3 %% 1:4
## [,1] [,2] [,3] [,4]
## [1,] 1 2 3 4
## [2,] 2 4 6 8
## [3,] 3 6 9 12
```

Grouped functional: tapply

Definition

Applies a function on a vector partioned by a factor: combine a split + lapply operation

Example

```
with(iris, tapply(Sepal.Length, Species, mean)) # readable
##
       setosa versicolor virginica
       5.006
                   5.936
                              6.588
##
with(iris, lapply(split(Sepal.Length, Species), mean)) # still ok (I think)
## $setosa
## [1] 5.006
##
## $versicolor
  Γ1] 5.936
##
## $virginica
## [1] 6.588
stat <- c() # less readable (and naming is lost)
for (1 in levels(iris$Species))
  stat <- c(stat, mean(iris$Sepal.Length[iris$Species == 1]))</pre>
stat
  [1] 5.006 5.936 6.588
```

Functionals for list: Reduce and co

Definition

'Reduce' uses a binary function to successively combine the elements of a given vector

```
Reduce(FUN, vector, init, right = FALSE, accumulate = FALSE)
Reduce(f, 1:3) <-> f(f(1,2))
```

Exercise

Consider a list of vectors of integer.

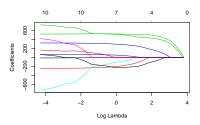
```
my_list <- replicate(10, sample.int(20, 10, replace = TRUE), simplify = FALSE)</pre>
```

Find which values occur in every element of the list visited so far, starting from the end.

Example: "jacknifing" a lasso solution path

A single Lasso fit of the diabete data set

```
library(glmnet)
library(lars) # the diabetes data set (part of the lars package)
data(diabetes)
x <- diabetes$x; y <- diabetes$y; n <- length(y)
lasso <- glmnet(x,y)
plot(lasso, xvar = "lambda")</pre>
```

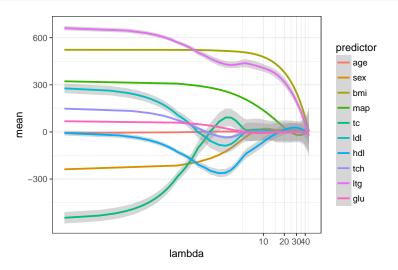


Compute the regularization paths for all subsets, removing one individual at once

```
paths <- lapply(1:n, function(i) {
    glmnet(x[-i, ], y[-i], lambda = lasso$lambda)$beta
})</pre>
```

A Reduce example: "jacknifing" a lasso solution path II

Computing the envelop around the average regularization path with Reduce



References I

R Core Team. (2017). *R: A language and environment for statistical computing*. Vienna, Austria: R Foundation for Statistical Computing. Retrieved from https://www.R-project.org/

Wickham, H. (2014). *Advanced r.* CRC Press. Retrieved from http://adv-r.had.co.nz/