# Advanced Programming with R

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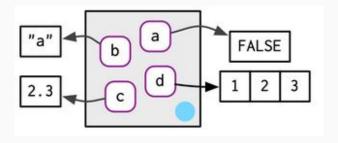
## Agenda

- Scoping & Environments
- Functionals & Split-Apply Paradigm
- Object Orientation (S3)
- Packages & Version Controlling

# Scoping & Environments

#### **Environments**

Like boxes, containing objects.



A bit simplified: If a function is called, its own environment is created with its parent being the environment from which it was called.

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#### **Environments**

```
simple_fun <- function(){</pre>
  a <- 1
  b <- "a"
  environment()
a <- simple_fun()
rlang::env_print(a)
# <environment: 0000000166B4598>
# parent: <environment: global>
 bindings:
# * b: <chr>
# * a: <dbl>
```

# Scoping

## Where does R find things?

- Argument matching (name, place...)
- Current environment
- Parent environment

## Programming advice

Keep it simple, this can create chaos!

# Scoping

```
add_things2 <- function(x) {
 x + 10 + y
add_things2(2)
# Error in add_things2(2): object 'y' not found
y <- 100
add_things2(2)
```

[1] 112

# **Functionals**

#### **Functionals**

A functional is a function that takes another function as an argument.

Focus on the apply-family. These functions apply a function repeatedly.

Can be seens as an abstraction of a for loop, with the following advantages

- requires less code to write
- does not store intermediate results
- no need to replace / grow

#### **Functionals**

The most commonly used functionals are:

- ullet lapply vector / list o list
- ullet sapply vector / list o vector (matrix)
- ullet apply matrix / array / data.frame o vector (matrix)
- tapply, by
- mapply, Map
- rapply, eapply, vapply

All of which have an argument that should be a function.

## lapply

Data frames are lists

```
lapply(iris, FUN = class)
# $Sepal.Length
# [1] "numeric"
  $Sepal.Width
# [1] "numeric"
#
# $Petal.Length
  [1] "numeric"
#
# $Petal.Width
# [1] "numeric"
#
# $Species
# [1] "factor"
```

## lapply

- an annonymous function can be used
- any type of element can be used
- other arguments can be passed through

```
lapply(c(min, median, max), FUN = function(fun, x) {
  fun(x)
, x = 2:8)
# [[1]]
# [1] 2
# [[2]]
# [1] 5
# [[3]]
# [1] 8
```

## sapply

- wrapper around lapply
- if possible, the ouput is combined into a atomic vector or matrix

```
sapply(airquality, FUN = sd)
    Ozone Solar.R Wind Temp Month
                                              Day
               NA 3.523001 9.465270 1.416522 8.864520
#
       NA
sapply(airquality, FUN = quantile, prob = c(.1, .9),
      na.rm = TRUE)
#
     Ozone Solar.R Wind Temp Month Day
# 10% 11 47.5 5.82 64.2 5 4
# 90% 87 288.5 14.90 90.0 9 28
```

- for objects with dimension (matrix, array, data.frame)
- apply over (a) chosen dimension(s)

```
my_matrix <- matrix(1:6, nrow = 2)
apply(my_matrix, 1, max)  # apply per row

# [1] 5 6
apply(my_matrix, 2, max)  # apply per column

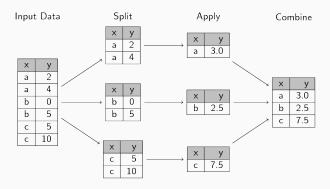
# [1] 2 4 6</pre>
```

```
my_matrix \leftarrow matrix(1:6, nrow = 2)
apply(my_matrix, c(1, 2),
     function(value) value^2) # apply per row and column
# [,1] [,2] [,3]
#[1,] 1 9 25
# [2,] 4 16 36
apply(airquality, 2, median)
#
   Ozone Solar.R
                  Wind
                                Month
                          Temp
                                        Day
      NΑ
             N A
                   9.7
                          79.0
                                7.0
                                         16.0
```

# Split & Apply & Combine

- split a data frame or vector (?split)
- apply computations on each split (lapply)
- combine the results (sapply, ?do.call)

## Split & Apply & Combine



# Exercises



**Object Oriented Programming** 

(S3)

# **Object Oriented Programming**

## Why?

- User-friendly: same function for different objects (summary())
- Coder-friendly: implementation can depends on object
- Coder-friendly: easier to maintain, extend

# **Object Oriented Programming**

#### **Basics**

- a *class* is a definition
- an object is an *instance* of a class.
- a method is a procedure that behaves differently depending of the class it is applied to.
- *inheritance*: classes are organized in hierarchy. "is-a-type-of"-relation.

# Object Oriented Programming in R

- S3
- S4
- RC
- R6
- ..

Compromise between interactive use, functional programming and object-oriented programming.

- "naming conventions"
- used in base R
- flexible: new classes, new methods

- class-attribute
- generics and methods
- inheritance and dispatch

A new class is made by adding an arbitrary class attribute to an object.

```
object <- 1:10
class(object) <- "myClass"
attributes(object)

# $class
# [1] "myClass"</pre>
```

Very flexible! Use it wisely!

```
class(iris)
# [1] "data.frame"
iris[1:4, 1:3]
#
  Sepal.Length Sepal.Width Petal.Length
# 1
               3.5
         5.1
                            1.4
# 2
         4.9 3.0 1.4
# 3
         4.7 3.2
                          1.3
         4.6
                3.1
                          1.5
# 4
```

Changing the class changes the behavior!

```
class(iris) <- "no data.frame"
class(iris)

# [1] "no data.frame"

iris[1:4, 1:3]

# Error in iris[1:4, 1:3]: incorrect number of dimensions</pre>
```

#### Good practice: constructor function

- defines the structure of the class
- should be used to create objects of that class

Write a separate validation-function for computationally intensive validation.

- methods for a class belong to generics
- when a generic is called for a specific class, the corresponing method for that class will be used. common generics are: print(), plot(), summary(), anova()
- same function, but different computation depending on object-class

```
body(print)
# UseMethod("print")
```

Creating a new method (for an existing generic):

```
print.myClass <- function(x, ...){</pre>
  cat("This is a myClass-print:\n")
  cat(round(c(nValues = length(x),
                 mean = mean(x),
                 SD = sd(x), 3, ...
my_object <- 1:5
class(my_object) <- "myClass"</pre>
print(my_object)
# This is a myClass-print:
# 5 3 1.581
```

## Good practices (enforced for packages on CRAN):

- A method must have all the arguments of the generic, including . . . if the generic does.
- A method must have arguments in exactly the same order as the generic.
- If the generic specifies defaults, all methods should use the same defaults.

Inspect the generic!

```
formalArgs(print)
# [1] "x" "..."

formalArgs(summary)
# [1] "object" "..."
```

## Creating a new generic:

```
center <- function(x, ...)</pre>
  UseMethod("center")
center.myClass <- function(x, ...){</pre>
  print("centering myClass:\n")
  return(x - mean(x))
center(my_object)
# [1] "centering myClass:\n"
# This is a myClass-print:
# 5 0 1.581
```

## S3 - Inheritance and Dispatch

Objects can have more than one class

```
my_model <- glm(as.factor(books) ~ pared, data = pisa,
                family = "binomial")
class(my_model)
# [1] "glm" "lm"
class(my_model) == "lm"
# [1] FALSE TRUE
inherits(my_model, "lm")
# [1] TRUE
```

Good practice: hierachical inheritance (subclass and superclass)

## S3 - Inheritance and Dispatch

When a method is not available for a (sub)class, the next available method (i.e., for the supperclass) will be used.

```
"variable.names.glm" %in% methods(class = "glm")
# [1] FALSE
"variable.names.lm" %in% methods(class = "lm")
# [1] TRUE
variable.names(my_model)
# [1] "(Intercept)" "pared"
```

# Exercises



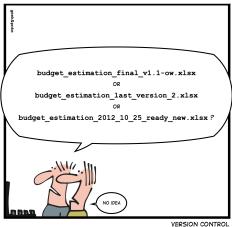
Version Controlling (Git + Github)

# Version Controlling

- Motivation
- Setup
- Work flow
- Recommendations
- Resources

## Motivation

### SIMPLY EXPLAINED



## Motivation

- Implementation of long term change history
  - No ridiculous file names
  - No archive subfolder
  - Always perfect overview of file history and changes
- Collaborations
  - What has changed?
  - Who has changed it?
  - Documentation of changes
  - Parallel working possible (merging)

#### But...



## Requirements

- Install git
- (optional) Install/Setup GUI for git (RStudio, Gitkraken, ...)
- Setup account for Github/Bitbucket/Gitlab/...
- Connect everything

### **RStudio**



## **RStudio**



#### Work flow

## Creating a repository

- Create an online repository (e.g. on Github)
  - Use an R specific .gitignore
  - Initialize with a short readme
- Clone the repository to your local machine
- (optional) Place an R project in the existing repository

## Work flow

## Working with a repository

- Before working: Synch your local repo (Pull)
- Perform changes in your local repository
- **Stage** your changes
- Commit your changes (aka new version)
- Push your changes

## Recommendations

- Keep it simple!
  - No branches/forks/pull requests
- Have meaningful commits
- Keep it lean (no big files)

#### Resources

## Git (+ R) Resources

- Small Intro (https://r-bio.github.io/intro-git-rstudio/)
- Happy Git with R (https://happygitwithr.com/)
- R Packages and Git (https://r-pkgs.org/git.html)
- Git Book (http://git-scm.com/book/en/v2)

Wrap Up

## **General Advice**

- Investing time in learning R pays off
- It's a steady learning curve
- Learn from masters
- Rewrite important code the first attempt is usually not the best approach

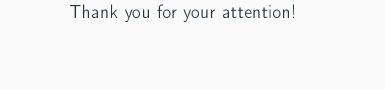
## General R Advice

- Document well
- Use a consistent style
- Write functions
- Split long functions in smaller ones
- Write wrappers
- Use Iteration (don't copy paste)
- Use matrix operations and vectorized functions instead of loops
- Use git

### Literature Recommendations

#### R Resources

- Avanced R Ed. 1 (http://adv-r.had.co.nz/)
- Avanced R Ed. 2 (https://adv-r.hadley.nz/)
- R Inferno (https: //www.burns-stat.com/pages/Tutor/R\_inferno.pdf)
- R Packages (https://r-pkgs.org/)
- Clean Code (https://enos.itcollege.ee/~jpoial/oop/ naited/Clean%20Code.pdf)



Thank you for your attention!

Questions? Remarks?