# (A bit of) Advanced R

Part 1 - towards better R-base programming

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http://github/jchiquet/CourseAdvancedR

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### Outline

1 Function, Functionals

#### References

- R Core Team (2017): A Language and Environment for Statistical Computing https://www.R-project.org/
- Wickham (2014): Advanced R, retrieved from http://adv-r.had.co.nz/
- Gillespie & Lovelace (2016): efficient R programming https://bookdown.org/csgillespie/efficientR/

### Prerequisites

#### Data Structure in base R

- 1 Atomic vector (integer, double, logical, character)
- Recursive vector (list)
- § Factors
- Matrices and array
- 6 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 programmming http://julien.cremeriefamily.info/teachings\_L3BI\_ISV51.html

### Outline

1 Function, Functionals

# The [a-z]\*pply family I

#### Example with factors (tapply)

```
data <- rnorm(100)
sexe <- factor( sample(c("H","F"), 100, replace = TRUE))
mean.1 <- tapply(data, sexe, mean) ## good
mean.2 <- c() ## complicated
for (1 in levels(sexe))
    mean.2 <- c(mean.2, mean(data[sexe == 1]))</pre>
```

Example with list or data.frame (sapply/lapply)

# The [a-z]\*pply family II

[1] TRUE [1] TRUE [1] TRUE [1] FALSE

### The do.call function I

## 3 bayes 0.9477400 141.573739

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

Suppose you have the outputs of 100 simulations at your disposable, stored in a list like that

### The do.call function II

## [1] 300

```
## [1] 100
How would you store them in a single data frame?
all.res <- do.call(rbind, res)
dim(all.res)</pre>
```

#### The Reduce function

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

→ can be use to post-process your list of simulations obtained via mclapply just like do.call

Say more... (map, Reduce)

# A Reduce 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)
y <- diabetes$y
x <- diabetes$x
n <- length(y)
lasso <- glmnet(x,y)
plot(lasso)</pre>
```

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

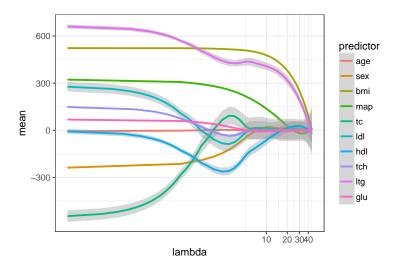
Compute the regularization paths for all subsets, removing one individual at once

```
paths <- parallel::mclapply(1:n, function(i) {
    glmnet(x[-i, ], y[-i], lambda = lasso$lambda)$beta
}, mc.cores = 4)</pre>
```

Computing the envelop around the average regularization path with Reduce

mean.path <- Reduce("+", paths)/n

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



### References

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