# **Functionals in R**

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Note: This is from *Functional Programming in R* https://github.com/Emaasit/Functional-Programming-in-R by Daniel Emaasit

### **Functionals**

Functions that take a function as input and returns a vector as output. Examples:

- lapply
- sapply + vapply (vector output)
- mapply + Map (multiple inputs)
- apply (matrices & arrays)
- tapply

These functionals are already implemented in base R.

#### **Uses of Functionals**

- Commonly used as an alternative to **For Loops**.
- For encapsulating common data manipulation tasks like split-apply-combine

#### **Advantages using Functionals**

- Reduce bugs
- Functionals implemented in base R are efficient & fast

# lapply (for lists)

lapply() takes a function, applies it to each element in a list, & returns a list.

```
input_list <- as.list(mtcars)</pre>
output_list <- lapply(input_list, length)</pre>
unlist(output list)
## mpg cyl disp
                   hp drat
                             wt gsec
                                            am gear carb
                                       VS
              32
                   32 32
##
   32 32
                             32 32
                                       32
                                            32
                                                 32
                                                      32
mtcars[] <- lapply(mtcars, function(x) x / mean(x))</pre>
```

#### **Looping Pattern 1: Over elements**

Looping over the elements in a list

```
lapply(xs, function(x) {})
```

#### **Looping Pattern 2: Over numeric indices**

Looping over the numeric indices in a list

```
lapply(seq_along(xs), function(x) {})
```

**Looping Pattern 3: Over the names** 

Looping over the names in a list

```
lapply(names(xs), function(nm) {})
```

## sapply & vapply (vector outputs)

Functionals that take a function, apply it to every element in a list, and return an atomic vector.

- sapply() guesses while vapply() takes an additional argument for the output type
- vapply() is better suited for use inside functions.

```
df <- data.frame(x = 1:10, y = Sys.time() + 1:10)
sapply(df, class)

## $x
## [1] "integer"
##
## $y
## [1] "POSIXct" "POSIXt"

# vapply(df, class, character(2))</pre>
```

## mapply & Map (multiple inputs)

Used when you have two or more lists (or data frames) that you need to process in parallel.

```
# Generate some sample data
xs <- replicate(n = 5, expr = runif(10), simplify = FALSE)</pre>
ws <- replicate(n = 5, expr = rpois(10, 5) + 1, simplify = FALSE)
Map(function(xs, ws) {weighted.mean(xs, ws)}, xs, ws)
## [[1]]
## [1] 0.5896866
##
## [[2]]
## [1] 0.3445823
##
## [[3]]
## [1] 0.5223156
##
## [[4]]
## [1] 0.429464
##
```

```
## [[5]]
## [1] 0.5215169
```

### Parallelisation (mclapply + mcMap)

Since we can compute each element in any order, it's easy to dispatch tasks to different cores, and compute them in parallel using the **parallel** package.

```
library(parallel)
system.time(mclapply(1:1000, sqrt, mc.cores = 4))

## user system elapsed
## 0.007 0.013 0.012

system.time(lapply(1:1000, sqrt))

## user system elapsed
## 0.000 0.000 0.001
```

## apply (for matrices & arrays)

apply() is a variant of lapply for working hig-order dimensional data objects

```
m <- matrix(1:100, nrow = 10)
apply(m, 1, mean) ## MARGINS, 1 = rows & 2 = columns
## [1] 46 47 48 49 50 51 52 53 54 55
apply(m, 2, mean)
## [1] 5.5 15.5 25.5 35.5 45.5 55.5 65.5 75.5 85.5 95.5</pre>
```

# tapply (for ragged arrays)

Ragged arrays are arrays where each row can have a different number of columns. apply() is useful for sumarizing a data set.

```
## Generate some ragged data
pulse <- round(rnorm(n = 22, mean = 70, sd = 10 / 3) + rep(c(0, 5), c(10,
12)))
group <- rep(c("A", "B"), c(10, 12))
split(pulse, group)

## $A
## [1] 75 66 67 72 63 66 71 76 68 74
##
## $B
## [1] 75 77 70 81 82 81 75 74 78 76 77 75

tapply(pulse, group, length)

## A B
## 10 12</pre>
```

```
tapply(pulse, group , mean)
## A B
## 69.80 76.75
```

# the plyr package

#### Reduce

Reduce a vector to a single value

```
Reduce(sum, 1:100)
## [1] 5050
## Find the values that occur in each element in this list
1 <- replicate(n = 5, sample(x = 1:10, size = 15, replace = TRUE), simplify = FALSE)
Reduce(intersect, 1)
## [1] 2 10 3 9</pre>
```

## **Predicate Functionals (Filter, Find, Position)**

- Predicates are functions that return a single TRUE or FALSE (e.g. is.character)
- Predicate Functionals applies a predicate to each element of a list or data frame.

### **Mathematical Functionals**