### Alternatives to Classic Loop Structures in R

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#### Some Basics

- ▶ Repeating a process/task is part of our daily life and sciences isn't left out of this.
- Repeating a process in programming languages is referred to as looping, which forms one of the basic thing to learn when learning a new language
- The standard function for repeatition in programming are; repeat, do, for and while
- ▶ In R, this classic loop structures are not as fast as they are in other High Level Languages, hence the need for alternatives.
- ▶ However, they are not as slow as people make them seem.

#### The Apply set of Functions

- ► The Apply set of functions are the first set of alternatives to loops in R
- ► There are about 4 to 5 of then (including a multicore version, mclapply in the *parallel package*)
- ?apply, ?lapply and ?sapply opens up the help pages in R
- apply(x, MARGIN, FUN, ...) allows you apply a function over the rows (MARGIN = 1) or columns (MARGIN = 2) of a matrix or dataframe
- sapply(x, FUN, ...) and lapply(x, FUN, ...) allows you apply a function over a list, matix or dataframe as well. lapply() always returns a list, sapply() tries to simplify the results.

# Tidyverse Offering (purrr Package)

- tidyverse is an ecosystem of packages dedicated to making data analysis faster https://tidyverse.org/
- ► It thrives on a set of data structures, grammar and philosophy aimed at making data analysis faster and reproducible.
- purrr's map(.x, .f, ...), its variants and walk(.x, .f, ...) function makes it possible to apply a function that allows more than one input to a list, array or dataframe.
- ?purrr::map() gives you the relating help pages.

#### An Hello World Example

 Let's start with a toy example (Bootstrapping and Determinants)

```
boot_func <- function(dat) {</pre>
  # sample from data
  sample(dat, length(dat), replace = T)
  # mean and standard deviation of dat
  return(c(mean(dat), sd(dat)))
sdata <- rnorm(80000, 0.5, 1.5)
mdata <- vector("list", 80000)
mdata<- lapply(mdata, function(x) {</pre>
  matrix(runif(15, 0, 1), nrow = 3, ncol = 3)
})
```

# Hello World Example Cont: (Using for() loop)

```
#time for sdata
system.time(for(i in 1:1000) {
  boot func(dat = sdata)
})
##
     user system elapsed
##
     2.28 0.37 2.67
#time for mdata
system.time(for(i in 1:length(mdata)) {
  det(mdata[[i]])
})
##
     user system elapsed
     0.91
             0.00
                     0.91
##
```

# Hello World Example Cont: (Using sapply())

```
#time for sdata
system.time(sapply(1:1000, function(i) {
 boot func(dat = sdata)
 }))
##
     user system elapsed
     2.31 0.38 2.69
##
#time for mdata
system.time(sapply(mdata, det))
##
     user system elapsed
##
     0.95
             0.00
                    0.95
```

## Hello World Example Cont: (Using purrr::rerun())

```
#time for sdata
system.time(purrr::rerun(1000, boot_func(dat = sdata)
))
##
     user system elapsed
     2.45
             0.33
##
                    2.78
#time for mdata
system.time(purrr::map(mdata, det))
##
     user system elapsed
##
     0.95
             0.00
                    0.95
```

# A Data Reading Example

### A Model Fitting Example

- ► The chicken dataset contains birth information of 628 chickens obtained via in-breeding and cross-bredding.
- ► Let's try to fit a regression model to relate the Age and Birth Weight of the 628 chickens.

```
model_func <- function(ddata) {
  if( length(which(!is.na(ddata$BW))) >= 3 ) {
    model <- lm(BW ~ AGE, data = ddata)
    slope <- coef(model)[2]
  } else {
    slope <- NA
  }
  return(slope)
}</pre>
```

# Using for() loop

```
slopes <- c()
system.time( for(i in 1:length(bychick$data)) {
  result <- model_func(ddata = bychick$data[[i]])
  slopes <- c(slopes, result)
})</pre>
```

```
## user system elapsed
## 0.48 0.00 0.48
```

## Using lapply()

```
system.time(lapply(bychick$data, model_func))
```

```
## user system elapsed
## 0.49 0.00 0.48
```

```
Using purrr::map()
```

```
## user system elapsed
## 0.47 0.00 0.46
```

#### Still to Come

▶ Introduction to Parallel Programming in R