

# 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 repetition 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 them (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, matrix 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

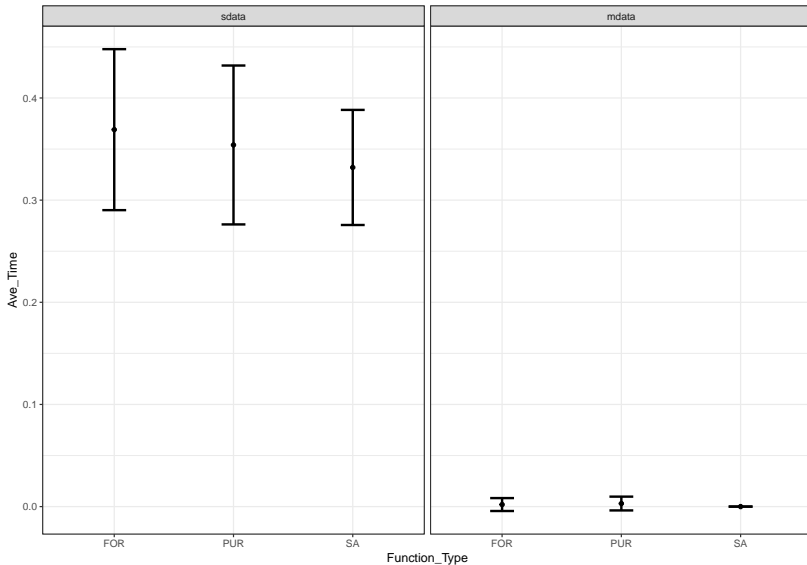
- ▶ Let's start with a toy example
- ▶ Bootstrapping and Determinants

```
#bootstrap function
boot_func <- function(dat) {
  # sample from data
  sample(dat, length(dat), replace = T)
  # mean and standard deviation of dat
  return(mean(dat))
}

sdata <- rnorm(80000, 0.5, 1.5)
# 3 X 3 matrices from uniform distribution
mdata <- vector("list", 80000)
mdata<- lapply(mdata, function(x) {
  matrix(runif(15, 0, 1), nrow = 3, ncol = 3)
})
```

- ▶ To be used for, `sapply`, `purrr::rerun()` & `purrr::map()`

## How They fared



► Considerably less time for the determinants, why?

## A Model Fitting Example

- ▶ The chicken dataset contains birth information of 628 chickens obtained via in-breeding or cross-breeding.
- ▶ 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)  
}
```

## 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)
})
```

>	user	system	elapsed
>	0.83	0.00	1.06



## Using lapply()

```
system.time(lapply(bychick$data, model_func))  
>    user  system elapsed  
>    0.72    0.00    0.72
```

## Using `purrr::map()`

```
system.time(map(bychick$data, model_func))  
>    user  system elapsed  
>    0.85    0.00    0.97
```

# Why Use these Alternatives?

- ▶ Cleaner and more readable codes
- ▶ Cleaner environment
- ▶ Dedicated outputs

## Still to Come

- ▶ Introduction to Parallel Programming in R
- ▶ Tidyverse, A language within a Language
- ▶ Check <https://github.com/fomotis/AvoidingLoops> for updates on this tutorial and associated files