

# Tips and Tricks for R Code Optimisation

## ➤ Optimising Code

Find biggest bottleneck and eliminate it;

➔ Profile code

```
Rprof("profiling.out")  
## CODE ##  
Rprof() summaryRprof("profiling.out")
```

When the bottleneck is found, try some of the following solutions;

1. **Look for existing solutions (online etc.)**
2. **Do less work**
  - Make functions do less work by tailoring functions to specific inputs/outputs
  - E.g. if possible avoid using `apply()` on a data frame (`apply()` first turns its input into a matrix)
3. **Vectorise**
  - Avoid loops
  - `rowSums()`, `colSums()`, `rowMeans()`, and `colMeans()`
4. **Parallelise**
  - Doesn't reduce computing time, but saves time by using more of a computer's resources
5. **Avoid copies**
  - If using `c()`, `append()`, `cbind()`, `rbind()`, or `paste()` to create a bigger object, R must first allocate space for the new object and then copy the old object to its new home. If repeating this many times, e.g. in a for loop, it can prove quite time-consuming
6. **Byte-code compile**
  - `compiler::cmpfun(function)`
  - Generally increases speed by 5 – 10% (base R already byte-code compiled)

## ➤ Function Speed

`microbenchmark(function, function, unit = "eps")` – compares speed of functions, be sure to take note of the units used, unit = "eps" notes the number of evaluations needed to take 1 second.

`system.time()` – less precise than above

Avoid unnecessary arguments – additional arguments slow down the function



### ➤ Memory

`pryr::object_size()` – accounts for shared elements within an object and includes size of environment

`pryr::mem_used()` – reports total size of all objects in memory

`gc()` – releases objects which are no longer used (R runs garbage collection automatically)

### ➤ Loops

- Avoid appending to a vector/list with each pass of a loop. Instead, first create an empty vector/list of the correct and then fill its elements.
- Taking statements which check for conditions outside the loop greatly speeds up code.
- Run the loop only for TRUE conditions
- For example,

```
output <- character(nrow(df))
condition <- (df$col1 + df$col2 + df$col3 + df$col4) > 4
for (i in (1:nrow(df))[condition]) { # run loop only for true conditions
  if (condition[i]) {
    output[i] <- "greater_than_4"
  } else {
    output[i] <- "lesser_than_4"
  }
}
df$output
```

- Use `ifelse()` when possible
- Use `which()` to select rows
- Use `apply()` rather than for loops e.g. `apply(data, 1, FUN = function)`

### ➤ Modification in place

Loops are notoriously slow as they modify a copy rather than modifying in place

Increase efficiency by using a list in this case, rather than a data frame. (lists modify in place)

### ➤ Data Structures

Simpler data structures which only store one data type can be manipulated faster e.g. representing data in a matrix rather than a data frame speeds up code considerably.

### ➤ data.table

This package speeds up operations on large data frames (particularly when sub-setting and indexing)



### ➤ Parentheses

Avoid using unnecessary brackets – the more ( ) used the longer code takes to run

{ } run quicker than ( ) - { } is treated as a special operator whose arguments aren't automatically evaluated, unlike ( ) which is an inbuilt operator whose arguments are automatically evaluated

**Note:** *time saved minimal*

### ➤ Readable Code

Code which is readable and reproducible can greatly save time

[Google's R Style Guide](#) is standard practice when coding in R

### ➤ Further Reading

[How to Speed Up R Code: An Introduction](#)

[Burns' R Inferno](#)

