#### Efficient Coding in R

Adam Kaplan / Tomoya Sasaki

March 17, 2023

## Efficiency in R

- Why is some code "fast" and other code "slow" in R?
- Two main culprits:
  - Bad algorithms,
  - Inefficient memory usage.
- Does not suffice to use and understand apply() family of functions.
- Need to understand basics of computational complexity and memory allocation.

#### Roadmap

- Basic of computational complexity
- Memory allocation
- Memory prescriptions
- Vectorization: apply and purrr
- Parallelization: future and foreach
- Random number generator

#### Section 1

## Computational Complexity

# Basics of Computational Complexity

- A theoretically rigorous way to measure "speed" of an algorithm as a function of input size.
- Put simply: How "fast" can  $f(x_1, ..., x_n)$  be evaluated as a function of n.
- Before we go any deeper, what do we mean by speed?
  - How many inputs  $x_i$  does f need to go over.
- Why can't we just measure time from start to end?
  - Many other factors impact the speed with which your code runs; harder to pin down issues.
- What can you do?
  - Find more appropriate algorithms,
  - 2 Vectorize whenever possible (why? stay tuned),
  - If possible, parallelize.

## Computational Complexity Examples

```
# Block 1: Simple for loop
for (i in 1:n) {
 foo()
# Block 2: Sequential for loops
for (i in 1:n) {
 foo()
for (i in 1:n) {
  bar()
```

```
# Block 3: Nested for-loops
for (i in 1:n) {
 foo()
 for (i in 1:n) {
   bar()
# Block 4: Simple bootstrap
for (i in 1:b) {
  # Hint: mle with BFGS takes O(n^2)
  estimate <- mle(X_1, ..., X_n, method = BFGS)
```

#### Section 2

#### Memory Allocation in R

## Memory Allocation in R

- Now we are algorithm experts and will measure the complexity of every piece of code.
- Is this enough? Unfortunately no.
- Even if the algorithms are efficient, we can lose real-life performance due to inefficient memory usage.
- Will go over basics of memory (in R), data types, and vectorization to motivate efficiency prescriptions.

# What is memory?

- Computer consists of three main parts: CPU core(s) (different circuits),
   Memory (RAM, Hard drive), GPU (can ignore for this session)
- CPU is wicked fast.
- Many different types of memory: Volatile (RAM, Caches), Non-volatile (Hard drive, USB, CDs, ...).
- Every R session lives on RAM memory.
- RAM is actually very very fast for memory standards, but painstakingly slow compared to CPUs, particularly saving new things to it (writes).
- To write efficient code, we need to avoid unnecessary memory writes!

## Memory Usage in R

- What happens when I delete an object with rm()?
  - You tell R that this reference to an object is no longer needed.
  - If there are no more, it will be "garbage collected" automatically.
- Big culprit: Copying of objects.
  - R uses a "copy if modified" framework so if y <- x, and y[1] <- 0, y
    will no longer point to x, but will allocate a new chunk of memory and
    point to that.</li>
  - ullet Avoid implicit and explicit copying of objects whenever possible.
  - "Vectorizing" is mostly faster because it avoids this implicit copying.

# Implicit copying of objects

```
# Example from: http://adv-r.had.co.nz/memory.html
library(pryr)
x \leftarrow data.frame(matrix(runif(100 * 1e4), ncol = 100))
medians <- vapply(x, median, numeric(1))</pre>
for (i in seq along(medians) |> head(5)) {
  x[.i] \leftarrow x[.i] - medians[i]
  print(c(pryr::address(x), pryr::refs(x)))
}
## [1] "0x561fd08d9d70" "1"
## [1] "0x561fd481ce20" "1"
   [1] "0x561fd8d21e70" "1"
   [1] "0x561fcef6aa00" "1"
   [1] "0x561fd2055370" "1"
```

#### Explicit copying objects

- Define the size and type of object to avoid copying object.
- Example: creating a vector of ones.

```
f1 <- function() { # Adding ones one by one
  x < -1.0
  for (i in 2:10000) {
    x < -c(x, 1.0)
  }
  return(x)
f2 <- function() { # Define length + type of vector
  x <- numeric(10000) # Can be character(), integer(), etc.
  for (i in 1:10000) {
   x[i] <- 1.0
  }
  return(x)
f3 <- function() { # Vectorization
  x \leftarrow rep(1.0, 10000)
  return(x)
```

12.6us

## 3 f3()

#### Functions in R

- From last time, arguments are not copied until modified, yay!
- What if we return a function in a function?
- Turns out: Functions save their surrounding environment!
- So what? Imagine you create a large variable (x <- 1:1e+50) before
  you create your function, then your function will also copy the large
  variable into memory.</li>
- Now on to more applied examples of why data types matter and how to vectorize in R.

#### Section 3

#### Memory Prescriptions

## Memory Prescription 1: Data Types Matter

- List vs. Matrix vs. Dataframe (or Tibble)
  - Matrix algebra is heavily optimized!
  - Neither lists nor matrices have memory overhead.
  - Dataframes have many specialized functions such as grouping that are much faster than DIY approaches on lists or matrices.

```
x <- runif(10000)
y <- rnorm(10000)
DF <- data.frame(x = x, y = y) # Dataframe
TIB <- tibble::as_tibble(DF) # Tidyverse tibble
MAT <- cbind(x, y) # Matrix</pre>
```

```
bench::mark( # Sum two columns for each row
 apply(DF, 2, sum),
 apply(TIB, 2, sum),
 apply(MAT, 2, sum),
  colSums(MAT) # Implemented in C
) |> dplyr::select(expression, median)
## # A tibble: 4 x 2
##
                         median
    expression
## <bch:expr>
                       <bch:tm>
## 1 apply(DF, 2, sum)
                         221.9us
## 2 apply(TIB, 2, sum)
                        235.4us
## 3 apply(MAT, 2, sum)
                        149.4us
```

## 4 colSums(MAT)

23.50s

#### Memory Prescription 2: Vectorize

- "Vectorize" your codes as much as you can.
- Most R functions allow you to vectorize by default.

#### Section 4

Vectorization: apply and purrr

## Vectorization: apply and purrr

- apply, lapply, sapply, ... come built-in with R and allow you to apply any function in a vectorized way to a matrix/dataframe, list, or vector.
- purrr's map, map\_dbl, map\_dfc, ... are the tidyverse equivalents and extensions to the apply() family.

```
library(purrr)
d <- list(
  data.frame(quant = c("danny", "insong", "teppei")),
  data.frame(schools = c("MIT", "Harvard"))
# We want a list as an output
lapply(d, nrow)
# Default map returns a list
purrr::map(d, nrow)
# The tidy version
d |> purrr::map(nrow)
```

```
library(purrr)
v <- 1:10

# We want a list as an output
sapply(v, \(x) x * x)

# The tidy version
v |> purrr::map_dbl(\(x) x * x)
```

#### Vectorization using purrr

- map\_\* return different object type and will fail if it is not appropriate.
- Makes the code more predictable, and thus easier to debug.
- Of particular interest are map\_dfr and map\_dfc, which run a function on the input and then row/column bind the outputs together.
- Inspired popular parallelization package furrr which we will introduce you to shortly.

Parallelize: future and foreach

#### Section 5

Parallelize: future and foreach

## Why and when to parallelize

- Enable multiple computations to take place at the same time
- Useful when you have time-consuming, unordered tasks
  - Data cleaning
  - Bootstrap
  - Monte Carlo simulations
  - Any tasks with lots of loops, apply, or maps

## Common issues in parallel computing

- Often need two different codes for parallel and non-parallel computing
  - Increase potential bugs
- Methods for parallel computing differ across operating systems and different packages implements different methods
  - mclapply. parallel, doParallel etc
  - Parallelization with mclapply (forking) does not work in Windows
  - Your choice of package would decide how to run in parallel

# future package

- future provides a simple and uniform tool for async. parallel, and distributed processing in R
- Same coding style between sequential and parallel tasks
- Users decide how to parallelize: the code does not depend on how to run in parallel
  - Use the same code for parallel computing in different operating systems

```
f <- future(expr) # Evaluate in parallel
r <- resolved(f) # Check if done
v <- value(f) # Get result</pre>
```

# Same coding style

```
f3 <- function() {
  tmp <- rnorm(10000)
  for (i in 1:10000) {
    tmp <- tmp + i
  }
  sum(tmp)
}
library(future)
future_obj <- future::future(f3()) # Evaluate in parallel</pre>
```

# Choose how to parallelize

- Sequential: plan(sequential)
- Parallelize on local machine: plan(multisession)
- "Fork" your task (not fore Windows): plan(multicore)
- Multiple local or remote computers plan(cluster)

# Combine together

```
#' `multisession` using two cores:
#' Default setup uses the max # of cores
#' Print with future::availableCores()
future::plan(future::multisession, workers = 2)

future_obj <- future::future(f3()) # Evaluate in parallel
output2 <- future::value(future_obj) # Get result
output <- f3()
all.equal(output, output2)</pre>
```

future::plan() # Check current plan

```
## multisession:
## - args: function (..., workers = 2, envir = parent.frame())
## - tweaked: TRUE
```

## - call: future::plan(future::multisession, workers = 2)

# User friendly functions

- Add future in front of the function
- Evaluate, check, get result within one function

```
# apply / future.apply
lapply(x, your_fun)

future::plan(future::multisession, workers = 2)
future.apply::future_lapply(x, your_fun)
```

• furrr package: integrate with purrr

```
# purrr / furrr
x |> purrr::map(your_fun)
library(furrr)
future::plan(future::sequential)
x |> furrr::future_map(your_fun)
```

# foreach for parallel computing

- A parallel/distributed computing framework for the R language
   A tool for "what to parallelize"
- Enables computation across multiple CPU cores and computers
- foreach runs sequentially in the absence of a parallel adapter
- doFuture from future framework provides a useful adapter for foreach

### Basic structure

```
library(foreach)
#' By default return a list where each element
#' is an output from each iteration
res <- foreach::foreach (i = 1:3) %do% {
  i + 1
}
res
## [[1]]
## [1] 2
##
## [[2]]
## [1] 3
##
## [[3]]
## [1] 4
```

## Basic structure 2

```
# You can iterate over two variables with the same length
res <- foreach::foreach(i = 1:3, j = 11:13) %do% {
  i + j
}
res
## [[1]]
## [1] 12
##
## [[2]]
## [1] 14
##
## [[3]]
## [1] 16
```

## Basic structure 3

```
# The output can be a vector
res <- foreach::foreach(
  i = 1:3, j = 11:13, .combine = "c"
) %do% {
  i + j
res
## [1] 12 14 16
# The output can be pretty flexible
res <- foreach::foreach(
  i = 1:3, j = 11:13, .combine = "+"
) %do% {
  i + j
res
```

## Parallelization with foreach and future

```
library(doFuture)
doFuture::registerDoFuture() # Register cores
# `multisession` with 2 cores
future::plan(future::multisession, workers = 2)
#' Compute mean while trimming values less than 0.1
#' `dopar` automatically detects parallel adapter
cutoff \leftarrow 0.1
y <- foreach::foreach(x = mtcars, .combine = "c") %dopar% {
  mean(x, trim = cutoff)
}
head(y, 4)
## [1] 19.696154
                    6.230769 222.523077 141.192308
```

#### Some notes

```
library(dplyr)
doFuture::registerDoFuture() # Register cores
# `multisession` with 2 cores
future::plan(future::multisession, workers = 2)
# Call functions explictly
y <- foreach::foreach(
  x = mtcars, .packages = c("dplyr")
) %dopar% {
  data |> dplyr::filter(x > 0)
}
```

#### xvii and foreach

- Your computer usually does not have enough cores for parallel computing
  - # of your cores = # of processors you can run simultaneously
- Use xvii, which has more than 20 cores
- The above codes work with xvii
- Do not use too many cores

## Section 6

## Random number generators

# Random number generation with Futures

- Motivating example: Monte Carlo simulations.
- To ensure replicability we would usually set a random seed using set.seed(02135) before running the simulations.
  Unfortunately, when we parallelize, since each process is independent of
- Unfortunately, when we parallelize, since each process is independent of each other, it is very likely that the random state across processes is no longer pseudo-random. In fact, it will likely be correlated.
- Even the naive solution of setting a different seed for each iteration does not guarantee the desired level of randomness.
- Thankfully, there are random number generation algorithms specifically designed for parallel computing. The default for future and most commonly used one is L'Ecuyer CMRG algorithm.
- Using it is as easy as in a sequential program.

# Random number generation in future.apply

```
library(future.apply)
# Set the seed
set.seed(1234)
# Specify that you want to use random numbers
out1 <- future.apply::future_sapply(1:10, \(x) runif(1),
   future.seed = TRUE
)</pre>
```

# Random number generation in furrr

```
library(furrr)
# Set the seed
set.seed(1234)
# Specify that you want to use random numbers
out2 <- furrr::future_map_dbl(1:10, \(x) runif(1),
    .options = furrr::furrr_options(seed = TRUE)
)
# Check that they are the same
all(out1 == out2)</pre>
```

## [1] TRUE

## Section 7

## Conclusion

# Key Takeaways

- Choose your data types appropriately.
- Use canned functions whenever possible and otherwise try to avoid nested for loops if possible.
- Vectorize whenever you can.
- Be on the lookout for explicit or implicit copying of objects.

## Section 8

Thank you for listening!