

Practicals

Jumping Rivers

Practical 1

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1. Reproduce the timing results in chapter 2 using the `benchmark` function from the `rbenchmark` package. Remember to load the package using

```
library("rbenchmark")
```

2. **Case study** In this example, we are going to investigate loading a large data frame. First, we'll generate a large matrix of random numbers and save it as a csv file:¹

```
N = 1e+05
m = as.data.frame(matrix(runif(N), ncol = 1000))
write.csv(m, file = "example.csv", row.names = FALSE)
```

¹ If setting `N=1e6` is too large for your machine, reduce it a bit. For example, `N=50,000`.

We can read the file the back in again using `read.csv`

```
dd = read.csv("example.csv")
```

To get a baseline result, time the `read.csv` function call above.

```
system.time(read.csv("example.csv"))
```

```
##      user  system elapsed
##    0.114    0.000    0.114
```

We will now look ways of speeding up this step. + Explicitly define the classes of each column using `colClasses` in `read.csv`, for example, if we have 1000 columns that all have data type numeric, then:

```
```r
read.csv(file="example.csv",
 colClasses=rep("numeric", 1000))
```
```

- Use the `saveRDS` and `readRDS` functions:

```
saveRDS(m, file = "example.RData")
readRDS(file = "example.RData")
```

- Install the `readr` package via

```
install.packages("readr")
```

Then load the package in the usual way

```
library("readr")
```

This package contains the function `read_csv`; a replacement function for `read.csv`. Is this new function much better than the default?

Which of the above give the biggest speed-ups? Are there any downsides to using these techniques? Do your results depend on the number of columns or the number of rows?

```
## 1. Using RData files is the fastest -  
## although you have to read the data in first.  
## Set colClasses also produces an good  
## speed-up.
```

```
## 2. Setting colClasses R is no longer  
## checking your data types. If your data is  
## changing - for example it's coming from the  
## web or a database, this may be problem.
```

```
## 3. The results do depend on the number of  
## columns, as this code demonstrates  
N = c(1, 10, 100, 1000, 1000, 10000)  
l = numeric(5)  
for (i in seq_along(N)) {  
  m = as.data.frame(matrix(runif(N[6]), ncol = N[i]))  
  write.csv(m, file = "example.csv", row.names = FALSE)  
  cc = rep("numeric", N[i])  
  l[i] = system.time(read.csv("example.csv",  
    colClasses = cc))[3]  
}  
l
```

```
## Notice that when we have a large number of  
## columns, we get a slow down in reading in  
## data set (even though we have specified the  
## column classes). The reason for this slow  
## down is that we are creating a data frame  
## and each column has to be initialised with a  
## particular class.
```

Practical 2

1. In this question, we'll compare matrices and data frames. Suppose we have a matrix, `d_m`

```
## For fast computers d_m = matrix(1:1000000,
## ncol=1000) Slower computers
d_m = matrix(1:10000, ncol = 100)
dim(d_m)

## [1] 100 100
```

and a data frame `d_df`:

```
d_df = as.data.frame(d_m)
colnames(d_df) = paste0("c", 1:ncol(d_df))
```

- Using the following code, calculate the relative differences between selecting the first column/row of a data frame and matrix.

```
benchmark(replications=1000,
          d_m[1,], d_df[1,], d_m[,1], d_df[,1],
          columns=c("test", "elapsed", "relative"))
```

Can you explain the result? Try varying the number of replications.

```
## Two things are going on here 1. The very
## large difference when selecting columns and
## rows (in data frames) is because the data is
## stored in column major-order. Although the
## matrix is also stored in column major-order,
## because everything is the same type, we can
## efficiently select values.
```

```
## 2. Matrices are also more memory efficient:
```

```
m = matrix(runif(10000), ncol = 10000)
d = data.frame(m)
object.size(m)
```

```
## 80200 bytes
```

```
object.size(d)
```

```
## 1120568 bytes
```

- When selecting columns in a data frame, there are a few different methods. For example,

```
d_df$c10
d_df[, 10]
d_df[, "c10"]
d_df[, colnames(d_df) == "c10"]
```

Compare these four methods.

2. Consider the following piece of code:

```
a = NULL
for(i in 1:n)
  a = c(a, 2 * pi * sin(i))
```

This code calculates the values:

$$2\pi \sin(1), 2\pi \sin(2), 2\pi \sin(3), \dots, 2\pi \sin(n)$$

and stores them in a vector. Two obvious ways of speeding up this code are: - Pre-allocate the vector `a` for storing your results. - Remove $2 \times \pi$ from the loop, i.e. at the end of the loop have the statement: `2*pi*a`. Try the above techniques for speeding up the loop. Vary n and plot your results. 3. R is an interpreted language; this means that the interpreter executes the program source code directly, statement by statement. Therefore, every function call takes time.² Consider these three examples:

² This example is for illustrative purposes. Please don't start worrying about comments and brackets.

```
n = 1e6
## Example 1
I = 0
for(i in 1:n) {
  10
  I = I + 1
}
## Example 2
I = 0
for(i in 1:n){
  ((((((((((10))))))))))
  I = I + 1
}
## Example 3
I = 0
for(i in 1:n){
  ##This is a comment
  ##But it is still parsed
  ##So takes time
  ##But not a lot
```

```
##So don't worry!  
10  
I = I + 1  
}
```

Using the `benchmark` function, time these three examples.

Practical 3: parallel programming

1. To begin, load the `parallel` package and determine how many cores you have

```
library("parallel")
detectCores()
```

2. Run the parallel `apply` example in the notes.
 - On your machine, what value of `N` do you need to use to make the parallel code run quicker than the standard serial version?
 - When I ran the benchmarks, I didn't include the `makeCluster` and `stopCluster` functions calls. Include these calls in your timings. How does this affect your benchmarks?
3. Run the dice game Monte-Carlo example in the notes. Vary the parameter `M`.³

³ Try setting `M=50` and varying `N`.

Solutions

Solutions are contained within this package:

```
library("jrEfficient")
vignette("solutions1", package = "jrEfficient")
```