

General principle: Memory allocation

WRITING EFFICIENT R CODE



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If we programmed in C...

WE'RE IN CHARGE OF MEMORY ALLOCATION

```
// C code: request memory for a number
x = (double *) malloc(sizeof(double));

// Free the memory
free(x);
```

- In R, memory allocation happens automatically
- R allocates memory in RAM to store variables
- Minimize variable assignment for speed

Example: Sequence of integers

$1, 2, \dots, n$

The obvious and best way

```
## Method 1  
x <- 1:n
```

Not so bad

```
## Method 2  
x <- vector("numeric", n) # length n  
for(i in 1:n)  
  x[i] <- i
```

Don't ever do this!

```
## Method 3  
x <- NULL # Length zero  
for(i in 1:n)  
  x <- c(x, i)
```

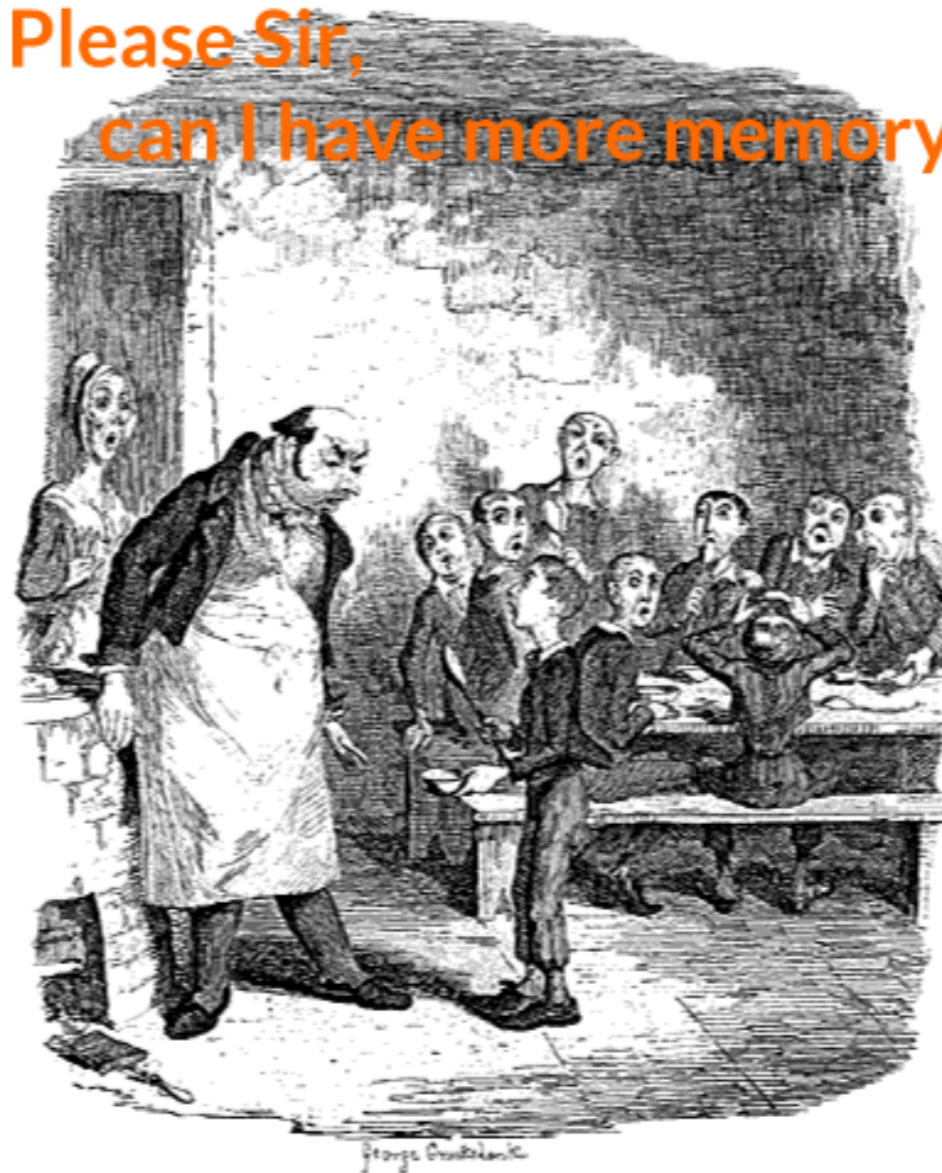
Benchmarking

- Method 1: 1:n
- Method 2: Preallocate
- Method 3: Growing

TIME IN SECONDS

n	1	2	3
10^5	0.00	0.02	0.2
10^6	0.00	0.2	30
10^7	0.00	2	3800

Please Sir,
can I have more memory?



Welcome to R club!

The first rule of R club: never, ever grow a vector.

Let's practice!

WRITING EFFICIENT R CODE

The importance of vectorizing your code

WRITING EFFICIENT R CODE



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General rule

- Calling an R function eventually leads to C or FORTRAN code
 - This code is *very* heavily optimized

Goal

- Access the underlying C or FORTRAN code as quickly as possible; the fewer functions call the better.

Vectorized functions

- Many R functions are vectorized
 - Single number but return a vector

```
rnorm(4)
```

```
-0.7247  0.2502  0.3510  0.6919
```

- Vector as input

```
mean(c(36, 48))
```

```
42
```

Generating random numbers

```
library(microbenchmark)
n <- 1e6
x <- vector("numeric", n)
microbenchmark(
  x <- rnorm(n),
  {
    for(i in seq_along(x))
      x[i] <- rnorm(1)
  },
  times = 10
)
```

```
# Unit: milliseconds
# expr      lq mean   uq  cld
# rnorm(n)   60  70   80  a
# Looping  2600 2700 2800  b

## Output trimmed for presentation
```

Compare

```
x <- vector("numeric", n)
for(i in seq_along(x))
  x[i] <- rnorm(1)
```

to

```
x <- rnorm(n)
```

Why is the loop slow?

LOOPING

```
x <- vector("numeric", n)
for(i in seq_along(x))
  x[i] <- rnorm(1)
```

GENERATION

- *Loop*: one million calls to `rnorm()`
- *Vectorized*: a single call to `rnorm()`

ALLOCATION

```
x <- vector("numeric", n)
```

- *Loop*: One-off cost
- *Vectorized*: Comparable

ASSIGNMENT

- *Loop*: One million calls to the assignment method
- *Vectorized*: a single assignment

R club

The second rule of R club: use a vectorized solution wherever possible.

Let's practice!

WRITING EFFICIENT R CODE

Data frames and matrices

WRITING EFFICIENT R CODE



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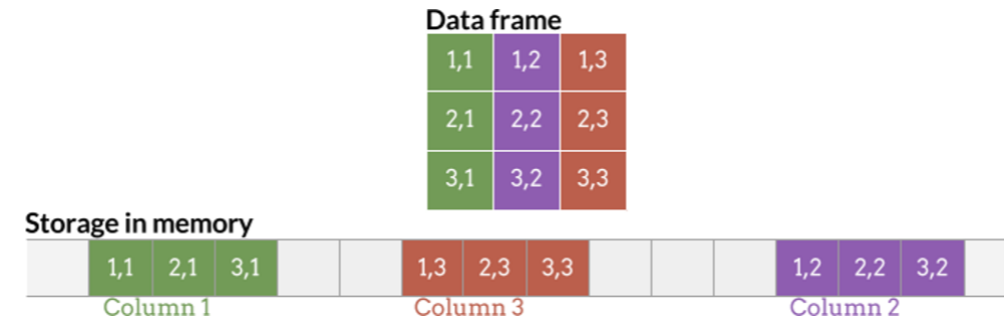
The data frame

- Key data structure in R
 - Copied in other languages
 - Python: pandas data frame
 - If you can't beat them, join them!

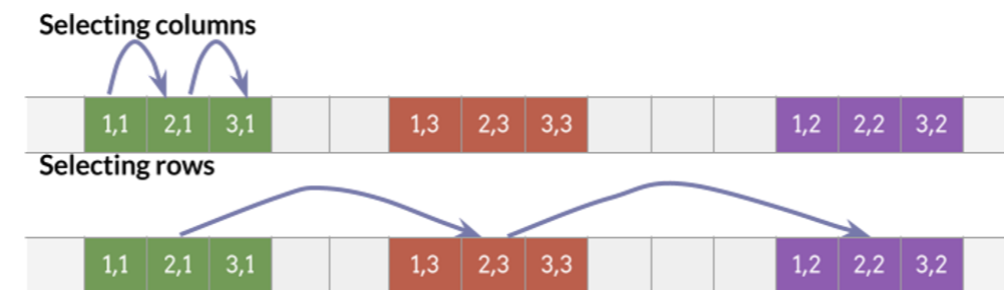
Data frames

- Tabular structure: rows and columns
 - `read.csv()` and friends returns a data frame
 - Columns
 - Data must be the same type
 - Rows
 - Different type

STORAGE



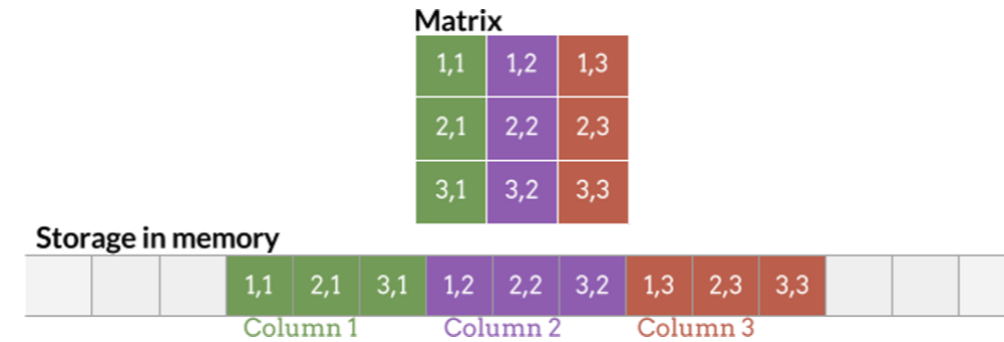
SELECTION



Matrices

- It's a rectangular data structure
 - You can perform usual subsetting and extracting operations
 - BUT - every element must be the same data type

STORAGE



SELECTION



R club

The third rule of R club: Use a matrix whenever appropriate.

Let's practice!

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