

# R Programming - Part 1

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## Who am i?



@eugeneteo

## Week 1

Getting started and R nuts and bolts

## Install R on a Mac

Install R with [Homebrew](#) - the missing package manager for OS X

```
$ ruby -e "$(curl -fsSL https://raw.githubusercontent.com/Homebrew/install/master/install)"
$ brew tap homebrew/science
$ brew install gcc
$ brew install Caskroom/cask/xquartz
$ brew install R
```

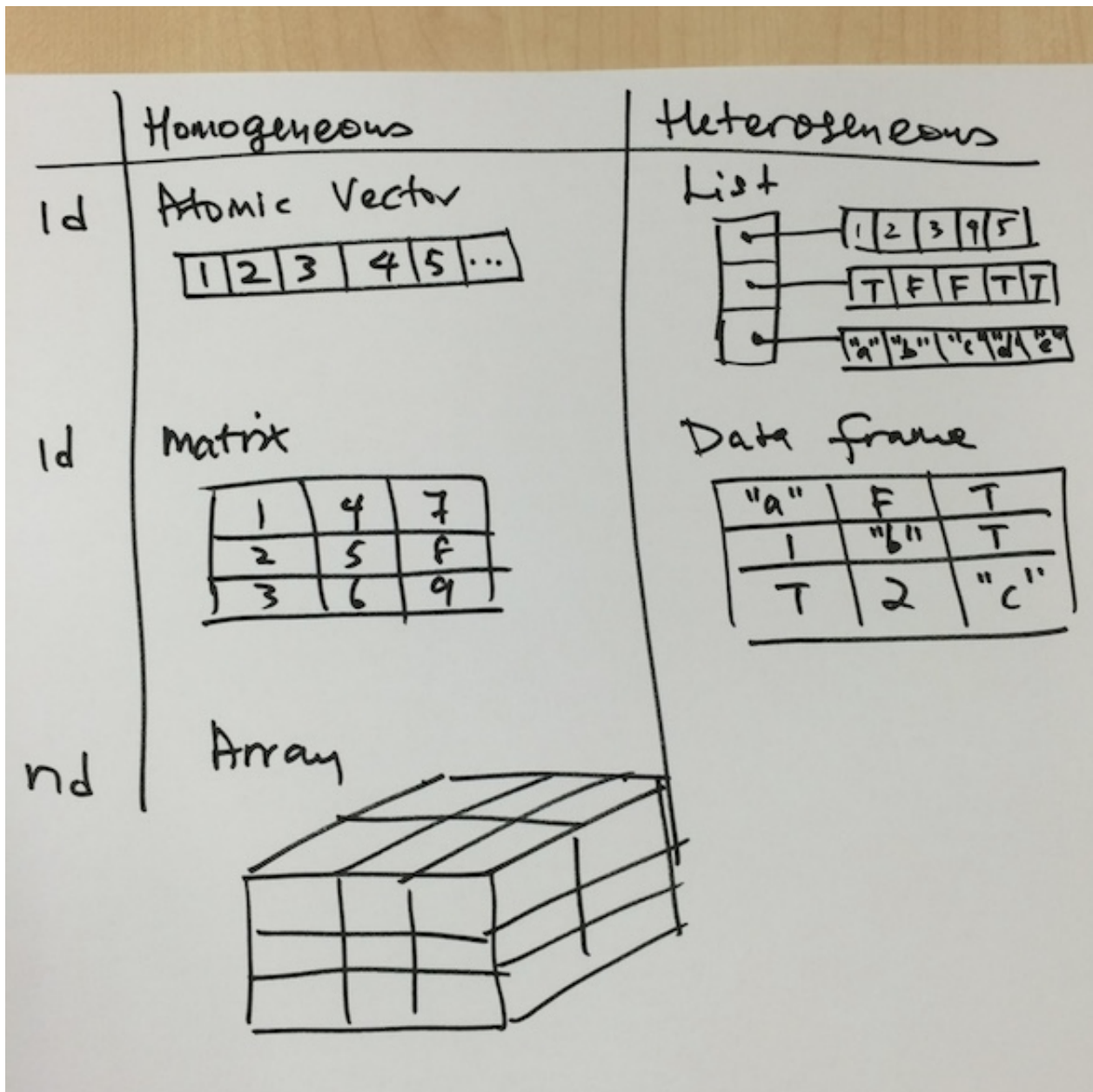
Then install the [RStudio Preview](#)

# Fix the warning messages on startup

```
$ defaults write org.R-project.R force.LANG en_US.UTF-8
$ export LC_ALL=en_US.UTF-8 \nextport LANG=en_US.UTF-8 >> ~/.bash_profile
$ . ~/.bash_profile
```

Read [Installing R on Mac - Warning messages: Setting LC\\_CTYPE failed, using "C"](#)

## Data structures



## Data structures

Data structures are organised by: \* their dimensionality (1d, 2d, or nd), and \* if they are homogeneous, or heterogeneous

R has no zero-dimensional, or scalar types

Individual numbers or strings are actually vectors of length one

```
length((x <- 1))
```

```
[1] 1
```

# Vectors

There are two types of 1d vectors: \* atomic vectors (homogeneous) \* lists (heterogeneous)

Both share three common properties: \* Type, `typeof()`, what it is \* Length, `length()`, how many elements it contains \* Attributes, `attributes()`, additional metadata

## Atomic vectors

There are four common types of atomic vectors: \* logical \* integer \* double \* character

```
log_var <- c(TRUE, FALSE, T, F)
char_var <- c(letters[1:3])
int_var <- c(1L, 2L, 3L)
dbl_var <- c(1, 2.1, 3.2)
```

## Atomic vectors

Atomic vectors are always flat

```
(x <- c(1, 2, 3, 4))
```

```
[1] 1 2 3 4
```

```
(y <- c(1, c(2, c(3, c(4)))))
```

```
[1] 1 2 3 4
```

## Coercion

All elements in an atomic vector must be of the same type. If not, coercion will occur from the least to the most flexible types: logical, integer, double, and character

```
(c("a", 1))
```

```
[1] "a" "1"
```

```
typeof(c(T, 1L, 2.0))
```

```
[1] "double"
```

# Lists

1d heterogeneous vector

```
str(x <- list(1:3, "a", c(T, F, T), c(1.2, 3.4)))
```

```
List of 4
 $ : int [1:3] 1 2 3
 $ : chr "a"
 $ : logi [1:3] TRUE FALSE TRUE
 $ : num [1:2] 1.2 3.4
```

# Lists

```
(x <- list(1:3, "a", c(T, F, T), c(1.2, 3.4)))
```

```
[[1]]
[1] 1 2 3

[[2]]
[1] "a"

[[3]]
[1] TRUE FALSE TRUE

[[4]]
[1] 1.2 3.4
```

# Lists

```
str(x <- list(a = 1:3, b= "a", c = c(T, F, T), d = c(1.2, 3.4)))
```

```
List of 4
 $ a: int [1:3] 1 2 3
 $ b: chr "a"
 $ c: logi [1:3] TRUE FALSE TRUE
 $ d: num [1:2] 1.2 3.4
```

# Lists

```
x <- list(a = 1:3, b= "a", c = c(T, F, T), d = c(1.2, 3.4))
x[[1]]
```

```
[1] 1 2 3
```

```
x$a
```

```
[1] 1 2 3
```

# Lists

Lists are sometimes called recursive vectors

A list can contain other lists

```
str(x <- list(list(list(list(list())))))
```

```
List of 1
 $ :List of 1
  ..$ :List of 1
   .. ..$ :List of 1
    .. .. ..$ : list()
```

# Matrices

2d homogeneous data structure

```
(x <- matrix(1:6, nrow = 2, ncol = 3))
```

```
      [,1] [,2] [,3]
[1,]    1    3    5
[2,]    2    4    6
```

```
dim(x)
```

```
[1] 2 3
```

# Matrices

```
x <- matrix(1:6, nrow = 2, ncol = 3)
# colnames(x) <- c("a", "b", "c")
# rownames(x) <- c("A", "B")
dimnames(x) <- list(c("A", "B"), # row
```

```
c("a", "b", "c"))  
x
```

```
  a b c  
A 1 3 5  
B 2 4 6
```

## cbind'ing and rbind'ing

```
x <- 1:3  
y <- 10:12  
cbind(x, y)
```

```
  x y  
[1,] 1 10  
[2,] 2 11  
[3,] 3 12
```

```
rbind(x, y)
```

```
  [,1] [,2] [,3]  
x     1     2     3  
y    10    11    12
```

## Factors

Used to represent categorical variables

May be represented by numbers, but does not make sense to add, subtract, average, etc

```
(x <- factor(c("A", "B", "AB", "O")))
```

```
[1] A  B  AB O  
Levels: A AB B O
```

```
str(x <- factor(c("A", "B", "AB", "O")))
```

```
Factor w/ 4 levels "A","AB","B","O": 1 3 2 4
```

## Factors

```
x <- factor(c("A", "B", "AB", "O"))  
nlevels(x)
```

```
[1] 4
```

```
attributes(x)
```

```
$levels  
[1] "A"  "AB" "B"  "O"  
  
$class  
[1] "factor"
```

## Factors

```
x <- factor(c("A", "B", "AB", "O"))  
attr(x, "levels") # levels(x)
```

```
[1] "A"  "AB" "B"  "O"
```

```
attr(x, "class") # class(x)
```

```
[1] "factor"
```

## Missing values

```
x <- c(1, 2, NA, 4, 5)  
is.na(x)
```

```
[1] FALSE FALSE  TRUE FALSE FALSE
```

```
table(is.na(x))
```

```
FALSE  TRUE  
    4     1
```

## Data frames

2d heterogeneous data structure

Can be converted into matrix using `data.matrix()`

```
(x <- data.frame(a = 1:3, b = c(T, F, T)))
```

```
      a      b
1 1  TRUE
2 2 FALSE
3 3  TRUE
```

```
dim(x)
```

```
[1] 3 2
```

## Data frames

Remember that all elements in a matrix must be of the same type

```
(x <- data.matrix(data.frame(a = 1:3, b = c(T, F, T))))
```

```
      a b
[1,] 1 1
[2,] 2 0
[3,] 3 1
```

## Reading data

```
(x <- read.csv(text = "value\n12\n1\n.\n9"))
```

```
      value
1        12
2         1
3          .
4         9
```

```
typeof(x$value)
```

```
[1] "integer"
```

## Reading data

```
x <- read.csv(text = "value\n12\n1\n.\n9")
as.double(x$value)
```

```
[1] 3 2 1 4
```

```
# ??
```



```
str(x$value)
```

```
Factor w/ 4 levels ".","1","12","9": 3 2 1 4
```

That's not what we want...

## Reading data

Second try

```
(x <- read.csv(text = "value\n12\n1\n.\n9", na.strings = "."))
```

```
  value
1    12
2     1
3    NA
4     9
```

```
as.double(x$value)
```

```
[1] 12  1 NA  9
```

## Reading data

Third try

```
str(x <- read.csv(text = "value\n12\n1\n.\n9", na.strings = ".",
                      stringsAsFactors = FALSE))
```

```
'data.frame':  4 obs. of  1 variable:
 $ value: int  12 1 NA 9
```

## Subsetting

```
x <- c(letters) # built-in constant
x[1]
```

```
[1] "a"
```

```
x[1:5]
```

```
[1] "a" "b" "c" "d" "e"
```

```
x[x > "s"] # return the TRUEs
```

```
[1] "t" "u" "v" "w" "x" "y" "z"
```

## Subsetting vectors

```
x <- c(letters)
table(y <- x > "w")
```

```
FALSE  TRUE
    23     3
```

```
x[y] # return the TRUEs
```

```
[1] "x" "y" "z"
```

## Subsetting lists

```
x <- list(foo = 1:4, bar = 0.6)
x[1]
```

```
$foo
[1] 1 2 3 4
```

```
x[[1]]
```

```
[1] 1 2 3 4
```

```
x[["foo"]]
```

```
[1] 1 2 3 4
```

## Subsetting lists

```
x <- list(foo = 1:4, bar = 0.6, baz = "hello")
x[c(1, 3)] # select 1st and 3rd vectors
```

```
$foo
[1] 1 2 3 4
```

```
$baz  
[1] "hello"
```

## Subsetting lists

```
x <- list(foo = 1:4, bar = 0.6, baz = "hello")  
x[[c(1, 4)]]
```

```
[1] 4
```

```
x[[c(2, 1)]]
```

```
[1] 0.6
```

```
x[[c(3, 1)]]
```

```
[1] "hello"
```

## Subsetting matrices

```
(x <- matrix(1:6, 2, 3))
```

```
      [,1] [,2] [,3]  
[1,]    1    3    5  
[2,]    2    4    6
```

```
x[1,] # returns a vector
```

```
[1] 1 3 5
```

```
x[,1] # returns a vector
```

```
[1] 1 2
```

## Subsetting matrices

```
(x <- matrix(1:6, 2, 3))
```

```
      [,1] [,2] [,3]  
[1,]    1    3    5
```

```
[2,]    2    4    6
```

```
x[1, 2] # return vector of length 1
```

```
[1] 3
```

```
x[1, 2, drop = FALSE] # return 1x1 matrix
```

```
      [,1]  
[1,]    3
```

## Removing NA values

```
x <- c(1, 2, NA, 4, NA, 5)  
(bad <- is.na(x))
```

```
[1] FALSE FALSE  TRUE FALSE  TRUE FALSE
```

```
(x[!bad])
```

```
[1] 1 2 4 5
```

```
(x[bad])
```

```
[1] NA NA
```

## Removing NA values

```
x <- c(1, 2, NA, 4, NA, 5)  
y <- c("a", "b", NA, "d", NA, NA)  
(good <- complete.cases(x, y))
```

```
[1]  TRUE  TRUE FALSE  TRUE FALSE FALSE
```

```
(x[good]) # show all good cases
```

```
[1] 1 2 4
```

```
(y[good][1:2]) # show 2 of 3 good cases
```

```
[1] "a" "b"
```

# Removing NA values

```
data("airquality")  
class(airquality)
```

```
[1] "data.frame"
```

```
good <- complete.cases(airquality)  
(airquality[good, ][1:6, ])
```

	Ozone	Solar.R	Wind	Temp	Month	Day
1	41	190	7.4	67	5	1
2	36	118	8.0	72	5	2
3	12	149	12.6	74	5	3
4	18	313	11.5	62	5	4
7	23	299	8.6	65	5	7
8	19	99	13.8	59	5	8

## Week 2

Programming with R

## Control structures - if/else

```
x <- 5  
if (x > 3) {  
  y <- 10  
} else {  
  y <- 0  
}  
y
```

```
[1] 10
```

## Control structures - if/else

```
x <- 5  
y <- if (x > 3) {  
  10  
} else {  
  0  
}
```

```
y
```

```
[1] 10
```

## Control structures - if/else

```
x <- 5  
y <- ifelse(x > 3, 10, 0)  
y
```

```
[1] 10
```

## Control structures - for

```
for (a in 1:10)  
  print(a)
```

```
[1] 1  
[1] 2  
[1] 3  
[1] 4  
[1] 5  
[1] 6  
[1] 7  
[1] 8  
[1] 9  
[1] 10
```

## Control structures - for

```
(x <- c(letters[1:3])) # built-in constant
```

```
[1] "a" "b" "c"
```

```
for (a in 1:length(x))  
  print(x[a])
```

```
[1] "a"  
[1] "b"  
[1] "c"
```

## Control structures - for

```
x <- c(letters[1:3])  
for (a in seq_along(x)) # element indices of x  
  print(x[a])
```

```
[1] "a"  
[1] "b"  
[1] "c"
```

```
for (letter in x)  
  print(letter)
```

```
[1] "a"  
[1] "b"  
[1] "c"
```

## Control structures - for

```
(x <- matrix(1:6, nrow = 2, ncol = 3))
```

```
      [,1] [,2] [,3]  
[1,]    1    3    5  
[2,]    2    4    6
```

```
for (a in 1:nrow(x))  
  for (b in 1:ncol(x))  
    print(x[a, b])
```

```
[1] 1  
[1] 3  
[1] 5  
[1] 2  
[1] 4  
[1] 6
```

## Control structures - for

```
x <- matrix(1:6, nrow = 2, ncol = 3)  
for (a in seq_len(nrow(x))) # sh's `seq 1 n`  
  for (b in seq_len(ncol(x)))  
    print(x[a, b])
```

```
[1] 1  
[1] 3  
[1] 5
```

```
[1] 2
[1] 4
[1] 6
```

## Control structures - for

```
x <- c(letters) # 26
for (letter in seq_along(x)) {
  if (letter == 20) break # stop at 20th
  if (letter %% 2 == 0) next # skip even
  print(x[letter])
}
```

```
[1] "a"
[1] "c"
[1] "e"
[1] "g"
[1] "i"
[1] "k"
[1] "m"
[1] "o"
[1] "q"
[1] "s"
```

## seq vs seq\_along

```
a <- c(8, 9, 10)
b <- 10

seq(a)
```

```
[1] 1 2 3
```

```
seq_along(a) # element indices of x
```

```
[1] 1 2 3
```

## seq vs seq\_along

```
a <- c(8, 9, 10)
b <- 10

seq(b) # became seq_len? :-(
```



```
[1] 1 2 3 4 5 6 7 8 9 10
```

```
seq_along(b) # element indices of x
```

```
[1] 1
```

Read [seq vs seq\\_along](#). When will using seq cause unintended results?

## Control structures - while

```
count <- 0
while (count < 10) {
  print(count)
  count <- count + 1
}
```

```
[1] 0
[1] 1
[1] 2
[1] 3
[1] 4
[1] 5
[1] 6
[1] 7
[1] 8
[1] 9
```

## Control structures - while

```
x <- 5
while (x > 2 && x < 6) {
  print(x)
  coin <- rbinom(1, 1, 0.5) # n, size, prob
  if (coin == 1) { # random walk
    x <- x + 1
  } else {
    x <- x - 1
  }
}
```

```
[1] 5
[1] 4
[1] 5
```

## Functions

```
args(lm)
```

```
function (formula, data, subset, weights, na.action, method = "qr",  
  model = TRUE, x = FALSE, y = FALSE, qr = TRUE, singular.ok = TRUE,  
  contrasts = NULL, offset, ...)  
NULL
```

Unnamed arguments are matched in order they are listed in the function definition

```
lm(data = mydata, y ~ x, model = FALSE, 1:100)  
lm(y ~ x, mydata, 1:100, model = FALSE)
```

## Functions

```
f <- function(a, b = 1, c = 2, d = NULL) {  
  
}
```

If no default value, you can set the argument value to NULL

## Functions - lazy evaluation

Arguments to functions are evaluated only as needed

```
f <- function(a, b) {  
  a ^ 2  
}  
f(10)
```

```
[1] 100
```

## Functions - lazy evaluation

```
f <- function(a, b) {  
  print(a)  
  print(b)  
}  
f(10)
```

```
[1] 10
```

```
Error in print(b) : argument "b" is missing, with no default
```

# Functions

The `...` argument indicates a variable number of arguments that are usually passed on to other functions

Also necessary if the number of arguments passed to the function cannot be known in advance

```
args(plot)
```

```
function (x, y, ...)  
NULL
```

```
myplot <- function(x, y, type = "l", ...) {  
  plot(x, y, type, ...)  
}
```

# Functions

Any arguments appear after `...` cannot be partially matched

```
args(paste)
```

```
function (... , sep = " ", collapse = NULL)  
NULL
```

```
paste("a", "b", sep = ":")
```

```
[1] "a:b"
```

```
paste("a", "b", se = ":")
```

```
[1] "a b :"
```

# Dates and times

Dates are stored internally as the # of days since 1970-01-01

Times are stored internally as the # of seconds since 1970-01-01

```
(date <- as.Date("1970-01-01"))
```

```
[1] "1970-01-01"
```

# Dates and times

Times are represented using: \* the `POSIXct` class (useful if you want to store times in a data frame) \* the `POSIXlt` class (useful if you need information like the day of the week, day of the year, month, day of the month)

# Dates and times

```
(x <- Sys.Date())
```

```
[1] "2015-01-27"
```

```
(y <- as.POSIXct(x))
```

```
[1] "2015-01-27 08:00:00 SGT"
```

```
unclass(y)
```

```
[1] 1.422e+09
```

# Dates and times

```
x <- Sys.Date()  
(y <- as.POSIXlt(x))
```

```
[1] "2015-01-27 UTC"
```

```
names(unclass(y))
```

```
[1] "sec" "min" "hour" "mday" "mon" "year" "wday" "yday" "isdst"
```

```
y$mday
```

```
[1] 27
```

# Dates and times

```
datestring <- c("January 10, 2015 10:40", "December 9, 2014 9:10")  
(x <- strptime(datestring, "%B %d, %Y %H:%M"))
```

```
[1] "2015-01-10 10:40:00 SGT" "2014-12-09 09:10:00 SGT"
```

```
class(x)
```

```
[1] "POSIXlt" "POSIXt"
```

Read `?strptime` for more details

## Dates and times

```
(x <- as.POSIXlt(as.Date("2015-01-01")))
```

```
[1] "2015-01-01 UTC"
```

```
(y <- strptime("9 Jan 2014 11:34:21", "%d %b %Y %H:%M:%S"))
```

```
[1] "2014-01-09 11:34:21 SGT"
```

```
x - y
```

```
Time difference of 356.9 days
```

## References

[R Programming](#) by Roger D. Peng, Jeff Leek and Brian Caffo

[Advanced R](#) by Hadley Wickham

## Thanks

Join our [iDA Data Sci MOOC](#) Facebook group!