# MFE R Programming Workshop

Week 1

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## Overview

### Goals

- Learn to program in R.
- ▶ What does programming mean?
  - Language syntax.
  - Debugging.
  - Finding solutions.
  - Translating math to code.
- ► This is just the beginning; you'll develop these skills throughout the program.

## R as a language

- R is object oriented.
  - Everything is an object and functions operate differently when passed different types of objects.
- R is functional.
  - Everything that happens in R is a function call.
  - You write fewer loops.
  - You write cleaner code.
- R is extendable.
  - ▶ Interfaces to other software are part of R.

### R vs C++

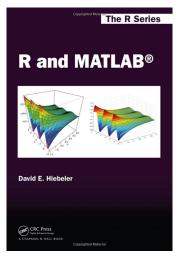
- ▶ Both are useful, and you will use both in the MFE program.
- R is an interpreted language.
  - Low programmer time.
  - ▶ A great tool for data munging, statistics, regressions, etc.
  - ▶ However, certain tasks in R can be slow (e.g. loops).
- ▶ C++ is very fast, but it takes longer to write programs.
- We can use both together!
- A good workflow:
  - 1. Write your program in R.
  - 2. If the program is too slow, benchmark your code.
  - 3. Try to speedup any bottlenecks in R.
  - 4. Convert any remaining bottlenecks to C++.

## R vs MATLAB, Python, ect

- ► Each language has its own set of strengths and weakness.
- ▶ You are better served by learning R and C++ very well, rather than trying to learn R, C++, MATLAB, Python, Julia, SAS, etc.
- The MFE program is just too short.
  - You also need to learn finance!
- Once you are proficient with R and C++, learning other languages is easy.
- ▶ Don't become a master of none!

### **MATLAB**

▶ If you want to learn MATLAB after learning R, take a look at R and MATLAB by David Hiebeler.



### Structure

- ▶ I will talk at the beginning of each class.
- ► For the remainder of the time you will break into your study groups and work on programming tasks.
- Tasks are designed to introduce you to the building blocks that will be used for course assignments throughout the MFE program.
- This course is a programming course with emphasis on methods for finance.
- ► The key skills will be translating mathematical algorithms into code and developing the ability to find helpful resources.

## Questions

Any questions before we start?

### R Resources: Books

- ► Introductory:
  - R for Everyone by Jared P. Lander
  - R Cookbook by Paul Teetor (free at UCLA LearnIT)
  - R for Data Science by Hadley Wickham (free as well)
- Intermediate:
  - ▶ The Art of R Programming by Norman Matloff
- Advanced:
  - Software for Data Analysis by John Chambers
  - Extending R by John Chambers
  - Advanced R by Hadley Wickham

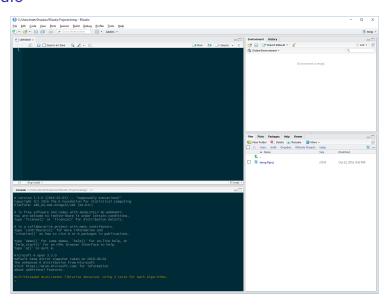
### Other Resources

- Book series:
  - Use R! Springer series
    - FYI: Many Springer textbooks are just \$25 through http://link.springer.com/. You need to be on campus or signed into the UCLA VPN. You can download the pdfs for free.
  - O'Reilly R Books (free at UCLA LearnIT)
- Built in documentation!
  - ?funcname
- ► Journal of Statistical Software
- Data science courses on Coursera
- ▶ Data Camp
- ▶ https://www.r-bloggers.com/
- ▶ https://twitter.com/rstudiotips
- ► Google, Stack Overflow, etc.

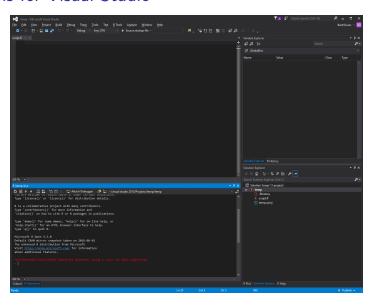
### R Environment

- First, you need an R distribution.
  - I recommend Microsoft R Open.
  - ▶ https://mran.microsoft.com/download/
- Second, you need an integrated development environment (IDE) for R.
  - R Studio is a fantastic environment to interact with R.
  - Other options:
    - R Tools for Visual Studio if you use Visual Studio.
    - ► Emacs Speaks Statistics (ESS) if you use Emacs.
- I am going to assume that you have a working installation of R Studio and that you have a basic understanding of how it works.
- I will show you some Visual Studio.
- My focus is going to be on R programming.

### **RStudio**



### R Tools for Visual Studio



### Course Materials

- ▶ https://github.com/brettdunn/mfeR2017
- ► The materials for this course were created in RStudio, using R Markdown.
- ► To create your own RStudio project:
  - ► File / New Project / Version Control / Git
  - Enter the URL



## R Basics

### Command Line Interface

➤ To run a command in R, type it into the console next to the > symbol and press the Enter key.

$$2 + 3$$

```
## [1] 5
```

- ▶ Up Arrow + Enter repeats the line of code.
- ► Esc (Windows/Mac) or Ctrl-C (Linux) interrupts a command.

### **RStudio**

- ► To start, create a new R Script file.
  - ► File/New File/R Script
- ➤ You can type your commands in the R Script file and run them on the Console.
  - Easy way to save your work.
  - Ctrl+Enter sends the line at the cursor to the console.
  - Ctrl+Shift+S runs the entire file.
  - Help/Keyboard Shortcuts lists all the available shortcuts.
    - RStudio Keyboard Shortcuts.
    - ▶ Check out the multiple cursors.
- ▶ For larger tasks with many files, create an R project.
- Visual Studio is similar.

### General Comments

- Make your code easy to read.
- ► Check out Google's R Style Guide
- Comment your code!
  - # indicates a comment in R.
  - Ctrl+Shift+c comments the line.

#### Google's R Style Guide

R is a high-level programming language used primarily for statistical computing and graphics. The goal of the R Programming Style Guide is to make our R code easier to read, share, and verify. The rules below were designed in collaboration with the entire R user community at Google.

#### Summary: R Style Rules

- 1. File Names: end in .R
- 2. Identifiers: variable.name (Or variableName), FunctionName, kConstantName
- 3. Line Length: maximum 80 characters 4. Indentation: two spaces, no tabs
- Curly Braces: first on same line, last on own line
- 7. else: Surround else with braces
- 8. Assignment: use < -, not =
- 9. Semicolons: don't use them 10. General Layout and Ordering
- 11. Commenting Guidelines: all comments begin with # followed by a space; inline comments need two spaces before the #
- 12. Function Definitions and Calls
- 13. Function Documentation
- 14. Example Function
- 15. TODO Style: TODO(username)

## R Packages

- ▶ A package is essentially a library of prewritten code designed to accomplish some task or a collection of tasks.
- R has a huge collection of user-contributed packages.
  - Warning: Not all packages are of the same quality.



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CRAN Task Views

Differential Equations Probability Distributions

**Rayesian Inference** 

Distributions Econometrics Econometrics Analysis of Ecological and Environmental Data Environmetrics

ExperimentalDesign Design of Experiments (DoE) & Analysis of Experimental Data

ExtremeValueTheory Extreme Value Theory Empirical Finance Finance Genetics Statistical Genetics

Graphics Graphic Displays & Dynamic Graphics & Graphic Devices & Visualization HighPerformanceComputing High-Performance and Parallel Computing with R

Machine Learning Machine Learning & Statistical Learning

MedicalImaging Medical Image Analysis

MetaAnalysis Meta-Analysis Multivariate Multivariate Statistics Natural Language Processing Natural Language Processing NumericalMathematics Numerical Mathematics Official Statistics

Official Statistics & Survey Methodology

### R Packages

- Installing a packages:
  - Ctrl+7 in RStudio accesses the packages pane
  - You can also type install.packages("packageName")
- Uninstalling a package:
  - remove.packages("packageName")
- Loading packages:
  - require(packageName) or library(packageName) loads a package into R
  - ► The difference is that require returns TRUE if the package loads or FALSE if it doesn't.
- Unloading packages
  - detach(package:packageName)
- ▶ If two packages have the same function name use two colons: -package1::func or package2::func

## Getting Help in R

- ▶ To get help on a function, use ?.
- ► The example function runs the examples contained in the help file.
- ▶ To run a search through R's documentation, use ??.
- To get help on a package, type help(package="packageName")

```
?seq # pulls up the help page
example(seq) # runs the examples in R
??"normal distibution" # runs a search
help(package = "xts") # gets help on the xts package
?'+' # gets help on the + function
```

### **Variables**

- ▶ Unlike C++, R does not require variable types to be declared.
- A variable can take on any data type.
- ▶ A variable can also hold any R object such as a function, the result of an analysis, a plot, etc.
- ▶ Variable assignment is done with <- (Alt+- in RStudio).</p>
  - ▶ = works, but there are reasons to prefer <-.
- We can remove variables (e.g. to free up memory) with the rm function.
  - gc() runs garbage collection.
  - rm(list=ls()) clears the workspace.

```
x <- 2  # x is a pointer
x  # the same output as print(x)
```

```
## [1] 2
```

```
rm(x) # removes x
```

## Data Types

- ▶ There are many different data types in R.
- ▶ The four main types of data most likely to be used are:
  - 1. numeric
  - character (string)
  - Date/POSIXct (time-based)
  - 4. logical (TRUE/FALSE)
- ► The data type can be checked with the class function

```
x <- as.Date("2010-12-21")
class(x)
```

```
## [1] "Date"
```

## Casting

```
x <- "2010-12-21"
class(x)
## [1] "character"
Х
## [1] "2010-12-21"
x <- as.Date(x)
class(x)
## [1] "Date"
Х
## [1] "2010-12-21"
```

## More Casting

```
x <- as.numeric(x)
class(x)
## [1] "numeric"
is.numeric(x)
## [1] TRUE
x # number of days since Jan 1, 1970
## [1] 14964
```

## **Even More Casting**

## [1] 0.8

```
x \leftarrow as.integer(x) \# x \leftarrow 14964L assigns an integer
class(x)
## [1] "integer"
is.integer(x)
## [1] TRUE
is.numeric(x) # R promotes int to numeric as needed
## [1] TRUE
4L / 5L
```

## Logicals

```
\# TRUE == 1 a.n.d. FALSE == 0
x <- TRUE # TRUE, FALSE, T, F are logicals
is.logical(x)
## [1] TRUE
5 == 5 # != tests for inequality
## [1] TRUE
"a" < "b" # works on characters as well
## [1] TRUE
```

## Vectors

### **Vectors**

- ▶ A vector is a collection of elements, all of the *same* type.
- ▶ In R, a vector does not have a dimension attribute.
  - There is no difference between a row vector and a column vector.
- We will learn about:
  - Recycling
    - The automatic lengthening of vectors.
  - Filtering
    - The extraction of subsets of vectors.
  - Vectorization
    - Where functions are applied element-wise to vectors.

## Vectors and Assignment

## [1] 1 2 3 4 5 6

- ► Assigning values to variables can be done with <-.
- ▶ Often, we create vectors using the c() function.
  - ► The "c" stands for combine because the arguments into a vector.

```
x <- c(1, 2, 3, 4)
x
## [1] 1 2 3 4
y <- c(x, 5, 6)
y
```

## Creating Vectors with seq and rep

- ▶ Both seq and rep are useful functions for generating vectors.
- ► See ?seq and ?rep for details
- seq is also useful in loops
- ▶ 1:10 is the same as seq(1,10,1)

```
x \leftarrow seq(from = 1, to = 10, by = 2)
X
## [1] 1 3 5 7 9
y \leftarrow rep(c(1, 2), times = 3)
## [1] 1 2 1 2 1 2
rep(c(1,2), each=2)
```

## Obtaining the Length of a Vector

## integer(0)

▶ length() returns the vector length

```
x <- c(TRUE, FALSE, TRUE, FALSE)
length(x)
## [1] 4
x \leftarrow c() # x is NULL
1:length(x) # that could mess you up in a for loop
## [1] 1 0
seq(x)
             # a safe way to loop through a vector
```

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## Accessing Elements of Vectors

- ► Elements can be accessed using []
  - ▶ Help on the [ function can be found by typing ?'['
- ▶ Unlike C/C++, R indexing starts at 1, not 0.
- ▶ The [ function can take a vector as an arguments.

```
x <- c("a", "b", "c", "d")
x[1]
                        # access the first element
## [1] "a"
x[c(1, 3)]
                         # access elements 1 and 3
## [1] "a" "c"
x[c(TRUE, FALSE, TRUE, FALSE)] # second way
## [1] "a" "c"
```

### **NULL** and NA

- ▶ NULL is the non-existent value in R.
- ▶ NA is the missing place holder.

```
x <- 5:8
x[2] <- NA
x
```

```
## [1] 5 NA 7 8
```

```
y <- NULL length(y)
```

```
## [1] 0
```

### Names of Vector Elements

- ▶ You can give names to elements of vectors, and you can access elements by their name.
- ▶ The function as.vector removes the names from a vector.

```
x <- 1:3
names(x) <- c("A","B","C")
x <- c(A=1, B=2, C=3) # another way
x["B"]</pre>
```

```
## B
## 2
```

```
as.vector(x) # the names are removed
```

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

#### Recycling

- When applying an operation to two vectors that requires them to be the same length, R automatically recycles the shorter one, until it is long enough to match the longer one.
- Be careful with and aware of this behavior!
- ▶ In some cases it is useful, others confusing.

```
# the shorter vector will be recycled c(2, 4, 6) + c(1, 1, 1, 2, 2, 2)
```

```
## [1] 3 5 7 4 6 8
```

```
# this is the same as
rep(c(2, 4, 6), 2) + c(1, 1, 1, 2, 2, 2)
```

```
## [1] 3 5 7 4 6 8
```

### **Logical Operators**

- R has several logical operations that act on vectors.
- ▶ !, ==, !=, &, &&, |, ||, xor(), any(), all(), >, >=, <=, <

```
x <- c(TRUE, FALSE, TRUE)
y <- c(TRUE, FALSE, FALSE)
x == y</pre>
```

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

!x

## [1] FALSE TRUE FALSE

# Logical Operations (2)

- ▶ &&, ||, any(), and all() return a length-one vector.
- ► The shorter forms are vectorized, meaning they can return a vector:

```
x <- c(FALSE, TRUE, TRUE)
y <- c(TRUE, TRUE, FALSE)
x & y</pre>
```

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

► The longer form evaluates left to right examining only the first element of each vector:

```
x && y
```

```
## [1] FALSE
```

## **Filtering**

We select subsets of vectors with vectors of logicals.

```
x <- 1:5
y <- c(TRUE, FALSE, TRUE, FALSE, TRUE)
x[y]</pre>
```

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

# Filtering (2)

Filtering amounts to generating filtering indices (i.e. vectors of logicals).

```
x \leftarrow c(5, 2, -3, 8)

idx \leftarrow x*x > 8

idx
```

## [1] TRUE FALSE TRUE TRUE

```
# another way
">"(x*x, 8)
```

## [1] TRUE FALSE TRUE TRUE

# Assigning to a Filter

- You can assign elements to the subsets.
  - ▶ This allows you change elements that meet certain criteria.

```
x <- 1:6

x[x <= 2] <- NA

x
```

```
## [1] NA NA 3 4 5 6
```

#### Filtering with subset()

## [1] 5 79

► The subset function filters and removes any NAs.

```
x \leftarrow c(3, 1:5, NA, 79)
Х
## [1] 3 1 2 3 4 5 NA 79
x[x > 4]
## [1] 5 NA 79
subset(x, x > 4)
```

## Filtering with Indices

- Select using the row/column number.
- ▶ Exclude entries with a negative vector.

```
x <- 10:20
x[c(1:2, 6:7)]

## [1] 10 11 15 16

x[-c(1,3)] # removes 1st and 3rd obs

## [1] 11 13 14 15 16 17 18 19 20</pre>
```

#### The Selection Function which()

which() gives us the position in a vector where a condition occurs.

```
x < -c(3, 1:5, NA, 79)
 x > 4
```

## [1] FALSE FALSE FALSE FALSE TRUE NA TRUE

```
which(x > 4)
```

## [1] 6 8

► See https://stackoverflow.com/questions/6918657/whats-the-use-of-which.

%in%

%in% returns a logical vector indicating if there is a match or not for its left operand.

```
1:5 %in% c(1,3)
```

## [1] TRUE FALSE TRUE FALSE FALSE

#### Vectorization: Functions on Vectors

- R functions typically operate on vectors.
- ▶ Often, there is an argument to ignore missing data.

```
x \leftarrow c(1:1000, NA)
mean(x)
## [1] NA
mean(x, na.rm = TRUE)
## [1] 500.5
log(x)[998:1001]
```

**##** [1] 6.905753 6.906755 6.907755

## Lists

#### **Creating Lists**

- A list is a structure that combines objects of different type and length.
- ▶ You can create a list where the elements are of type list.

```
element1 <- 1:5
element2 <- matrix(1:6, nrow=2)
mylist <- list(el1=element1, el2=element2)
mylist</pre>
```

```
## $el1
## [1] 1 2 3 4 5
##
## $el2
## [1,] [,2] [,3]
## [1,] 1 3 5
## [2,] 2 4 6
```

#### Accessing Elements of Lists

## [1] 1

We can access a list component in several different ways.

```
mylist <- list(A=1, univ=c("UCLA", "USC"),</pre>
               mymat=matrix(1:4, nrow=2))
mylist[[1]] # first way
## [1] 1
mylist[["A"]] # second way
## [1] 1
mylist$A
               # third way
```

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## Removing Components of Lists

▶ We can delete a component of a list by setting it to NULL.

```
mylist <- list(A=1)</pre>
mylist$B <- c(1, 2) # adds a component to a list
mylist
## $A
## [1] 1
##
## $B
## [1] 1 2
mylist$A <- NULL
mylist
```

```
## $B
## [1] 1 2
```

### Subsetting Lists

- Subsets of lists are done with single [].
  - ► A single [] returns a sublist of the original list

```
## $A
## [1] 1
##
## $mymat
## [,1] [,2]
## [1,] 1 3
## [2,] 2 4
```

## Lists of Objects

Lists can store all types of objects.

```
1 \leftarrow list(a = matrix(1:4, nrow=2),
         b = list(A=1:10,B=20:30))
1[["a"]]
## [,1] [,2]
## [1,] 1 3
## [2,] 2 4
1[[2]][[1]]
   [1] 1 2 3 4 5 6 7 8 9 10
##
```

# Simplfying vs Preserving Subsetting

Note that [[ simplifies the result, returning a vector (str displays the structure of an R object):

```
str(mylist[[2]])
## chr [1:2] "UCLA" "USC"
```

▶ But [ is preserving subsetting and returns a list:

```
str(mylist[2])

## List of 1
## $ univ: chr [1:2] "UCLA" "USC"
```

# Applying Functions to a List with lapply

lapply implicitly loops over each list element and applies a function.

```
mylist <- list(A=1:10,B=2:17,C=745:791)
lapply(mylist,mean)</pre>
```

```
## $A
## [1] 5.5
##
## $B
## [1] 9.5
##
## $C
## [1] 768
```

## An Example of lapply

From ?lapply: lapply(X, FUN, ...) returns a list of the same length as X, each element of which is the result of applying FUN to the corresponding element of X.

```
1 <- c("A","B","B","A","A","B")
lapply(c("A","B"), function(letter) which(l==letter))

## [[1]]
## [1] 1 4 5
##
## [[2]]
## [1] 2 3 6</pre>
```

#### Investments Problem Set 1

You will submit a list of answers.

```
hw <- source("./lecture1p.R")$value
str(hw)</pre>
```

#### **Functions**

#### **Function Definitions**

- ▶ The last value evaluated is what is returned by the function.
- You can also write return(x^2).
  - This is useful if you want to break out of the function early.

```
myfunc <- function(x) x^2
myfunc(10)</pre>
```

```
## [1] 100
```

### Scope Rules for Functions

# You can't access N out here

▶ Variables defined inside a function are local to that function.

```
myfunc <- function(x) {
    N <- 10
    return(N*x^2) # return is optional
}
myfunc(10)
## [1] 1000</pre>
```



# The Pipe Operator %>%

- ► The magnittr package provides a pipe operator.
- ► See vignette("magrittr").
- Basic piping:
  - x %>% f is equivalent to f(x)
  - x %>% f(y) is equivalent to f(x, y)
  - x %>% f %>% g %>% h is equivalent to h(g(f(x)))
- ▶ The argument placeholder:
  - $\rightarrow$  x %>% f(y, .) is equivalent to f(y, x)
  - $\rightarrow$  x %>% f(y, z = .) is equivalent to f(y, z = x)

# Expose the variables with %\$%

► The %\$% allows variable names (e.g. column names) to be used in a function.

```
library(magrittr)
iris %>%
  subset(Sepal.Length > mean(Sepal.Length)) %$%
  cor(Sepal.Length, Sepal.Width)
```

```
## [1] 0.3361992
```

# Compound assignment pipe operations with %<>%

There is also a pipe operator which can be used as shorthand notation in situations where the left-hand side is being "overwritten":

```
iris$Sepal.Length <-
iris$Sepal.Length %>%
sqrt()
```

Use the %<>% operator to avoid the repetition:

```
iris$Sepal.Length %<>% sqrt
```

► This operator works exactly like %>%, except the pipeline assigns the result rather than returning it.