MFE R Programming Workshop

Week 1

Brett Dunn and Mahyar Kargar

Fall 2017

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++

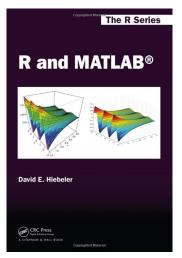
- ▶ Both are useful, and you will use both in the MFE program.
- R is an interpreted language.
 - ▶ Low programmer time.
 - ▶ A great tool for data munging, statistics, regressions, etc.
 - ▶ However, certain tasks in R can be slow (e.g. loops).
- ▶ C++ is very fast, but it takes longer to write programs.
- 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

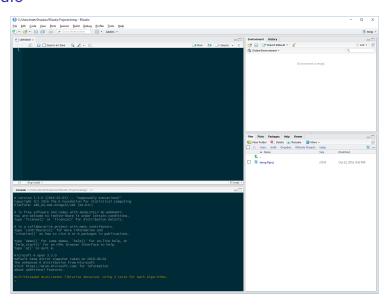
Other Resources

- 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.

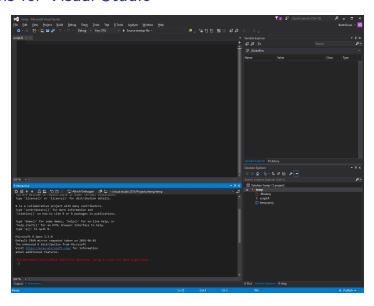
R Environment

- 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

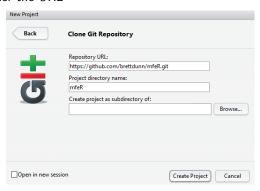


R Tools for Visual Studio



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 URI



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.



CRAN Mirrors What's new? Task Views Search

About R R Homepage The R Journal

Software R Sources R Binaries Packages Other

Documentation Manuals FAOs Contributed

Bayesian

ChemPhys

Chemometrics and Computational Physics

Finance Genetics Graphics

MetaAnalysis

ClinicalTrials Cluster

DifferentialEquations Distributions

Econometrics Environmetrics

ExperimentalDesign ExtremeValueTheory

Machine Learning MedicalImaging

Multivariate

NumericalMathematics Official Statistics

Rayesian Inference Clinical Trial Design, Monitoring, and Analysis

Cluster Analysis & Finite Mixture Models Differential Equations Probability Distributions

Econometrics Analysis of Ecological and Environmental Data Design of Experiments (DoE) & Analysis of Experimental Data

Extreme Value Theory Empirical Finance Statistical Genetics

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

CRAN Task Views

Machine Learning & Statistical Learning Medical Image Analysis

Meta-Analysis Multivariate Statistics Natural Language Processing Natural Language Processing Numerical Mathematics

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).
 - ▶ = 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)</pre>
```

```
## [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
 - 2. 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

```
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
## [1] 0.8
```

Logicals

[1] TRUE

```
\# 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
```

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

- ► 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
```

[1] 1 2 3 4 5 6

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
```

33 / 66

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

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
```

[1] 1

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

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

Everything that happens in R is a function call

[1] 15

```
`<-`(mynumber, 3)</pre>
print(mynumber)
## [1] 3
^+ (3,4)
## [1] 7
a <- : (11,20)
[(a,5)]
```

Function Definitions

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

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

- ▶ 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.

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:
 - 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.

Lab 1

Let's work on Lab 1.