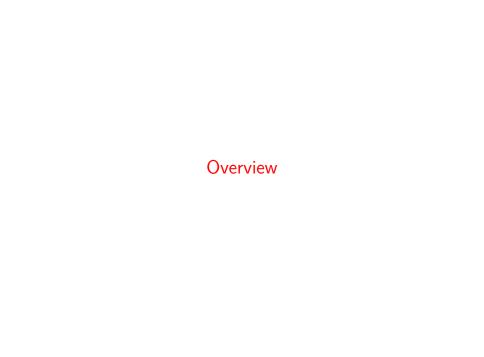
# MFE R Programming Workshop

Week 1

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Fall 2016



## 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.
  - You write fewer loops.
  - You write cleaner code.

## R vs C++

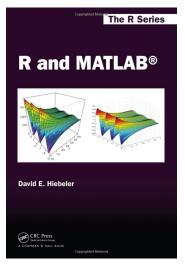
- Both are useful, and you will use both in the MFE program.
- R is an interpreted language.
  - ► Low programmer time.
  - ▶ A great tool for data munging, statistics, regressions, ect.
  - ▶ However, certain tasks in R can be slow (e.g. loops).
- ▶ C++ is very fast, but it takes longer to write programs.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

## Jack of All Trades, Master of None

- ▶ You are better served by learning R and C++ very well, rather than trying to learn R, C++, MATLAB, Python, Julia, SAS, ect.
- ▶ The MFE program is just too short.
  - ▶ You also need to learn finance!
- ➤ Once you are proficent 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:
  - You will see finance terms and math.
  - You may not understand all of the finance, but you will learn it throughout the program.
- ► 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

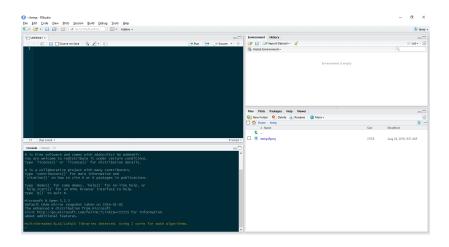
## Other Resources

- 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, ect.

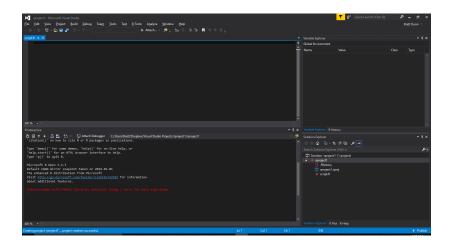
#### R Environment

- First, you need an R distribution.
  - ▶ I recommend Microsoft R Open.
  - ▶ https://mran.revolutionanalytics.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**

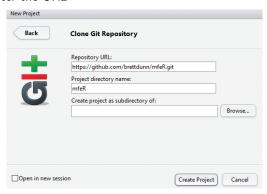


## R Tools for Visual Studio



## Course Materials

- ▶ https://github.com/brettdunn/mfeR
- ► 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) interups 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 cusor to the consule.
  - Ctrl+Shift+S runs the entite file.
  - ► Help/Keyboard Shortcuts lists all the available 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!

#### 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
- Identifiers: variable.name (Or variableName), FunctionName, kConstantName
- Line Length: maximum 80 characters
   Indentation: two spaces, no tabs
- 5 Spacing
- Curly Braces: first on same line, last on own line
- . Curry Braces
- else: Surround else with braces
   Assignment: use <-. not =
- Assignment: use <-, not =</li>
   Semicolons: don't use them
- General Layout and Ordering
   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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 ChemPhys
 Chemometrics and Computational Physics

 ClinicalTrials
 Clinical Trial Design, Monitoring, and Analysis

 Cluster
 Cluster Analysis & Finite Mixture Models

 Differential Equations
 Differential Equations

 Distributions
 Probability Distributions

 Econometrics
 Econometrics

Environmetrics Analysis of Ecological and Environmental Data

ExperimentalDesign Design of Experiments (DoE) & Analysis of Ex

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

Extreme Value Theory

Extreme Value Theory

Finance Empirical Finance

Genetics Statistical Genetics

Graphics Graphic Displays & Dynamic Graphics & Graphic Devices & Visualization

HighPerformanceComputing High-Performance and Parallel Computing with R

CRAN Task Views

MachineLearning Machine Learning & Statistical Learning

Medical Image Analysis
Meta-Analysis
Meta-Analysis

Meta-Analysis
Multivariate
Multivariate Statistics
NaturalLanguageProcessing
NumericalMathematics
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

#### **Variables**

- ▶ Unlike C++, R does not require variable types to be decleared.
- 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, ect.
- ▶ Variable assignment is done with <-.
  - = works, but there are reasons to prefer <-.</p>
- ▶ We can remove variables (e.g. to free up memory) with the rm function. gc() runs garbage collection.

```
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
  - 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
```

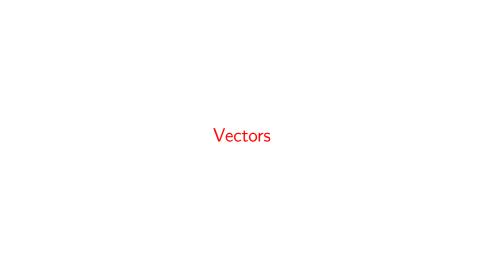
# **Even More Casting**

```
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 and 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

- ▶ A vector is a collection of elements, all of the same type.
- 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 <-</p>
- Create vectors of numbers using the function c()

```
myvector <- c(1,2,3,4)
samevector <- 1:4
vectorsubset <- myvector[1:3]
vectorsubset</pre>
```

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

# c() function

▶ c(), which stands for concatenate is a very flexible function.

```
myvec1 <- 2:4
myvec2 <- c(1,3,5,myvec1)
myvec2</pre>
```

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

## Accesing elements of vectors

▶ Elements can be accessed using []

```
myvec <- c(2,4,6,8)
myvec[4]

## [1] 8

myvec[c(1,3)]

## [1] 2 6</pre>
```

# Length

▶ length() returns the vector length

```
myvec <- 1:23
length(myvec)</pre>
```

```
## [1] 23
```

## Recycling

## [1] 2 4 4 6

- Vectors are recycled when an operation acts elementwise
- Be careful with and aware of this behavior!
- ▶ In some cases it is useful, others confusing

```
vec1 <- 1
vec2 <- 1:4
vec3 <- 1:2
vec2+vec1
## [1] 2 3 4 5
vec2+vec3</pre>
```

## seq and rep

- Useful functions for generating vectors
- ► See ?seq and ?rep for details

```
myvec1 <- seq(1,10,2)
myvec1</pre>
```

```
## [1] 1 3 5 7 9
```

```
myvec2 <- rep(c(1,2),3)
myvec2</pre>
```

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

```
rep(c(1,2), each=2)
```

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

## **NULL** and NA

## [1] 0

- NULL is the non-existent value in R
- ► NA is the missing place hold

```
myvec1 <- 5:8
myvec1[2] <- NA
myvec1

## [1] 5 NA 7 8

myvec2 <- NULL
length(myvec2)</pre>
```

# Filtering (1)

Select subsets using vectors of logicals

```
vec1 <- 1:5
vec2 <- c(TRUE, FALSE, TRUE, FALSE, TRUE)
vec1[vec2]</pre>
```

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

# Filtering (2)

Select subsets using vectors of logical

```
vec1 <- 1:6
vec2 \leftarrow vec1 > 3
vec2
## [1] FALSE FALSE FALSE TRUE TRUE TRUE
vec1[vec2]
## [1] 4 5 6
vec1[vec1>3]
## [1] 4 5 6
```

# Assigning to filter

You can assign to the subsets

```
vec1 <- 1:6
vec1[vec1<2] <- NA
vec1</pre>
```

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

#### Functions on vectors

Functions typically operate on vectors

```
x <- 1:1000
mean(x)
```

```
## [1] 500.5
```

#### names

You can give names to elements of vectors

```
myvec <- 1:3
names(myvec) <- c("A","B","C")
myvec["B"]</pre>
```

## B

## 2

# Matrices

#### Intro

 Matrices are vectors with a number of rows and number of columns attribute

```
myvec <- 1:10
mymat <- matrix(myvec, nrow=2, ncol=5)
mymat</pre>
```

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

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

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

#### Matrix operations

► Many matrix operations are surrounded by % signs

```
mymat1 <- matrix(1:4, nrow=2)</pre>
mymat2 <- matrix(5:8, nrow=2)</pre>
mymat1 %*% mymat2
## [,1] [,2]
## [1,] 23 31
## [2,] 34 46
mymat1 + mymat2
## [,1] [,2]
## [1,] 6 10
## [2,] 8 12
```

#### apply

- apply allows you to apply a function across a dimension of a matrix
- The third argument is a function!

```
mymat1 <- matrix(1:10, nrow=2)
# mean across rows
apply(mymat,1,mean)</pre>
```

```
## [1] 5 6
```

#### cbind and rbind

Column bind and Row bind

```
mymat1 <- matrix(1:4,nrow=2)
mymat2 <- matrix(6:9,nrow=2)
mymat3 <- matrix(10:11,ncol=2)
cbind(mymat1,mymat2)</pre>
```

```
## [,1] [,2] [,3] [,4]
## [1,] 1 3 6 8
## [2,] 2 4 7 9
```

rbind(mymat1,mymat3)

```
## [,1] [,2]
## [1,] 1 3
## [2,] 2 4
## [3,] 10 11
```

# Lists

#### Intro

► Lists are where you really start to see the advantages of R as a statistics and data manipulation language

```
element1 <- 1:5
element2 <- matrix(1:6,nrow=3)
mylist <- list(el1=element1,el2=element2)
mylist[["el1"]]

## [1] 1 2 3 4 5

mylist[["el2"]]</pre>
```

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

### Subsetting lists

Subsets of lists are with single []

```
mylist <- list(A=1,B=2,C=3,D=4)
# this returns a list because of the single []
mylist[c(1,3)]</pre>
```

```
## $A
## [1] 1
##
## $C
## [1] 3
```

#### 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
```

### lapply example

```
g <- c("M", "F", "F", "I", "M", "M", "F")
lapply(c("M", "F", "I"), function(gender) which(g==gender))
## [[1]]
## [1] 1 5 6
##
## [[2]]
## [1] 2 3 7
##
## [[3]]
## [1] 4
```

Data Frames

#### Intro

- In my mind, data.frame is the core data type in R
- The nicest part is that they can hold different types
- ► Each column must be the same length

```
dat1 <- 1:4
dat2 <- rep(c("A","B"),each=2)
myframe <- data.frame(col1=dat1,col2=dat2)
myframe</pre>
```

```
## col1 col2
## 1 1 A
## 2 2 A
## 3 3 B
## 4 4 B
```

#### Subsets

## [1] A A B B ## Levels: A B

Subsetting is just like a matrix

```
myframe[1,2]
## [1] A
## Levels: A B
myframe[1,]
## col1 col2
## 1 1 A
myframe[,2]
```

### Reading in data

- Reading in data typically gives you a data.frame
- read.table is the basic function to read in tabular data
- read.csv is a special cast of read.table
- ► As usual see ?read.table
- Often you want to set stringsAsFactors = FALSE

### Adding columns

```
dat1 <- 1:4
dat2 <- rep(c("A","B"),each=2)
myframe <- data.frame(col1=dat1,col2=dat2)
myframe$col3 <- 5:8
myframe</pre>
```

```
## 1 1 A 5
## 2 2 A 6
## 3 3 B 7
## 4 4 B 8
```

#### names

- column names in data.frames are specificed by names()
- this is because data.frames are actually lists with special attributes
- that means that the usual list functions work on data.frames
- lapply, etc

### Long example

```
all2006 <- read.csv("2006.csv",header=TRUE,as.is=TRUE)
# exclude hourly-wagers
all2006 <- all2006[all2006$Wage Per=="Year", ]
# exclude weird cases
all2006 <- all2006[all2006$Wage_Offered_From > 20000,]
all2006$rat <- all2006$Wage_Offered_From
                  / all2006$Prevailing_Wage_Amount
se2006 <- all2006[grep("Software Engineer", all2006),]
```



# For loops (1)

## [1] 3 4 5 6 7

```
x <- c(1:5)
y <- NULL
for(i in 1:length(x)) {
    y[i] <- x[i] + 2
}
y</pre>
```

# For loops (2)

or another nice way to make a for loop

```
x <- c(1:3)
for(element in x) {
   print(element + 2)
}</pre>
```

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

### While loops

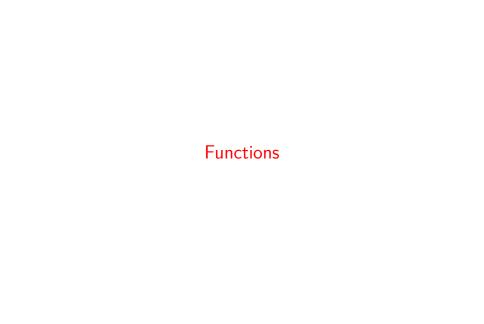
## [1] 3 4 5 6 7

```
x <- c(1:5)
y <- NULL
i <- 1
while(i<=length(x)) {
    y[i] <- x[i] + 2
    i <- i+1
}
y</pre>
```

#### **Conditional Statements**

## [1] 10

```
x <- -10
myabs <- x
if(x<0) {
    myabs <- -x
}
myabs</pre>
```



#### Function definitions

► Note that the last value evaluated is what is returned by the function

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

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

### Scope rules

▶ Variables defined inside a function are local to that function

```
myfunc <- function(x) {
    N <- 10
    N*x^2
}
myfunc(10)</pre>
```

## [1] 1000

```
# You can't access N out here
```

Lab 1

Let's work on Lab 1.