

Practical R: Lecture 1

Introductions

- Abhijit Dasgupta (NIH/NIAMS)
 - PhD in Biostatistics
 - 15+ years using R
 - Runs the Statistical Programming DC meetup (formerly the R Users Group meetup)
 - Board member of Data Community DC
- Gene Buehler (NIH/NCATS)
 - PhD in Computer Science
 - 20 years in bioinformatics
 - R user for past ten years

Class Structure

- Weekly Lectures
- Suggested activities and online resources
- Semester Project

Semester Project

- Use R to import, manipulate, analyze and visualize a dataset.
- Ideally it will be your data, or your lab's data, or at least data relevant to your field
- Short presentations will be given on the last day of class for each project
- Grade will be based on your project presentation (although if you can't show up we can grade based on your presentation slides).

What this class isn't

- A programming class
- A bioinformatics class
- A workshop

Very Brief Course Outline

- Basics of using R (variables, functions, arguments)
- Getting Help (R help system, Google, command history)
- Importing data into R (from files, from URLs, and preloaded data)
- Packages: finding them, loading them, reading their documentation
- Visualizing Data: Graphs, graphs and more graphs
- Statistical Analysis: How and why
- Manipulating data: selecting, filtering, merging and more.
- Science specific analysis and visualization (eg. Heatmaps and dendograms for gene expression data)

There's more than one way

- Lots of ways to run R: command line, GUI, RStudio.
- Frequently more than one package to do what you want
- More than one statistical test you can use on a given type of data

Homework: Installing R and RStudio

- <https://cran.rstudio.com>
- <https://www.rstudio.com/products/rstudio/download/>
- Both are available for Windows, MacOSX, and Linux.

RStudio

Project: (None)

Go to file/function

Addins

Console

```
R version 3.3.1 (2016-06-21) -- "Bug in Your Hair"
Copyright (C) 2016 The R Foundation for Statistical Computing
Platform: x86_64-apple-darwin13.4.0 (64-bit)

R is free software and comes with ABSOLUTELY NO WARRANTY.
You are welcome to redistribute it under certain conditions.
Type 'license()' or 'licence()' for distribution details.

Natural language support but running in an English locale

R is a collaborative project with many contributors.
Type 'contributors()' for more information and
'citation()' on how to cite R or R packages in publications.

Type 'demo()' for some demos, 'help()' for on-line help, or
'help.start()' for an HTML browser interface to help.
Type 'q()' to quit R.

> x <- 1
> y <- 2
> x + y
[1] 3
> lilVector <- c(0,5,12,33,82,77,54,97)
> myMatrix <- matrix(1:100, nrow = 10, ncol=10)
> plot(myMatrix)
> |
```

Environment

Global Environment

Data	
myMatrix	int [1:10, 1:10] 1 2 3 4 5 6 7 8 9 10 ...
Values	
lilVector	num [1:8] 0 5 12 33 82 77 54 97
x	1
y	2

Files

New Folder Delete Rename More

Home

	Name	Size	Modified
<input type="checkbox"/>	Applications		
<input type="checkbox"/>	Desktop		
<input type="checkbox"/>	Development		
<input type="checkbox"/>	Documents		
<input type="checkbox"/>	Downloads		
<input type="checkbox"/>	Infocom		
<input type="checkbox"/>	Kaggle		
<input type="checkbox"/>	Library		
<input type="checkbox"/>	Movies		
<input type="checkbox"/>	Music		
<input type="checkbox"/>	perl5		
<input type="checkbox"/>	Pictures		
<input type="checkbox"/>	Public		

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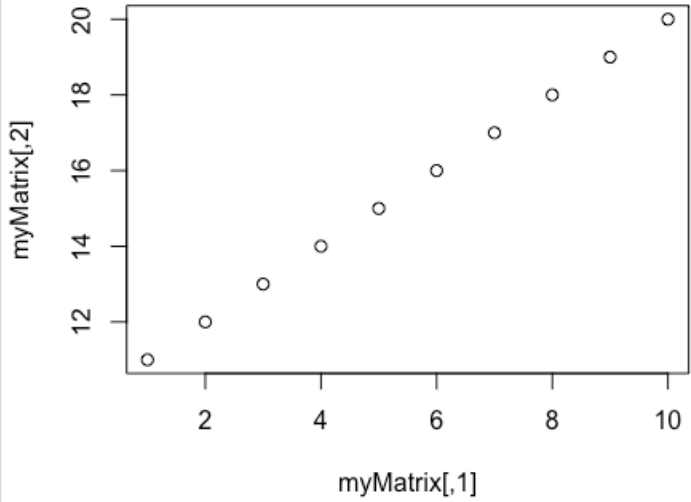
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Environment History

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Files Plots Packages Help Viewer

Zoom Export Publish



myMatrix[,1]	myMatrix[,2]
1	1
2	4
3	9
4	16
5	25
6	36
7	49
8	64
9	81
10	100

RStudio

Go to file/function Addins Project: (None)

Console ~/ ↻

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Files **Plots** **Packages** **Help** **Viewer**

Install Update

Name	Description	Version	
System Library			
<input type="checkbox"/> boot	Bootstrap Functions (Originally by Angelo Canty for S)	1.3-18	⊗
<input type="checkbox"/> class	Functions for Classification	7.3-14	⊗
<input type="checkbox"/> cluster	"Finding Groups in Data": Cluster Analysis Extended Rousseeuw et al.	2.0.4	⊗
<input type="checkbox"/> codetools	Code Analysis Tools for R	0.2-14	⊗
<input type="checkbox"/> compiler	The R Compiler Package	3.3.1	⊗
<input checked="" type="checkbox"/> datasets	The R Datasets Package	3.3.1	⊗
<input type="checkbox"/> foreign	Read Data Stored by Minitab, S, SAS, SPSS, Stata, Systat, Weka, dBase, ...	0.8-66	⊗
<input checked="" type="checkbox"/> graphics	The R Graphics Package	3.3.1	⊗
<input checked="" type="checkbox"/> grDevices	The R Graphics Devices and Support for Colours and Fonts	3.3.1	⊗
<input type="checkbox"/> grid	The Grid Graphics Package	3.3.1	⊗
<input type="checkbox"/> KernSmooth	Functions for Kernel Smoothing Supporting Wand & Jones (1995)	2.23-15	⊗
<input type="checkbox"/> lattice	Trellis Graphics for R	0.20-33	⊗
<input type="checkbox"/> MASS	Support Functions and Datasets for Venables and Ripley's MASS	7.3-45	⊗

RStudio

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Files **Plots** **Packages** **Help** **Viewer**

Home Find in Topic

R Resources

- [Learning R Online](#)
- [CRAN Task Views](#)
- [R on StackOverflow](#)
- [Getting Help with R](#)

RStudio

- [RStudio IDE Support](#)
- [RStudio Cheat Sheets](#)
- [RStudio Tip of the Day](#)
- [RStudio Packages](#)
- [RStudio Products](#)

Manuals

- [An Introduction to R](#)
- [Writing R Extensions](#)
- [R Data Import/Export](#)
- [The R Language Definition](#)
- [R Installation and Administration](#)
- [R Internals](#)

Reference

- [Packages](#)
- [Search Engine & Keywords](#)

R Start-up Message

R version 3.2.0 (2015-04-16) -- "Full of Ingredients"
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Platform: x86_64-apple-darwin13.4.0 (64-bit)

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[Workspace loaded from ~/.RData]

>

Unpacking the Startup

Version Number

Often silly version name

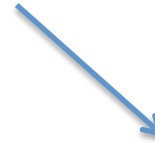
```
R version 3.2.0 (2015-04-16) -- "Full of Ingredients"  
Copyright (C) 2015 The R Foundation for Statistical Computing  
Platform: x86_64-apple-darwin13.4.0 (64-bit)
```

Operating system this version is made for

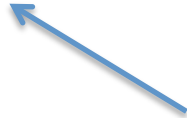
32 or 62 bit

Unpacking the Startup

“Don’t sue us”



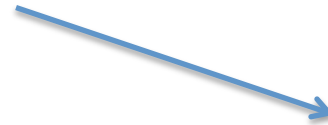
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Terms under which you can distribute the software

Unpacking the Startup

R has support for different local conventions. For example, you can get it to use “,” instead of “.” for decimal notation



Natural language support but running in an English locale

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R is made by real people who deserve credit (citations) for their work as much as you do. Please give it to them.

Unpacking the Startup

Some R packages have “demos” or their capabilities. For example, `demo(graphics)` will show you some of the cool graphs R can make.

R’s help function. You will probably use this a lot.

Type `'demo()'` for some demos, `'help()'` for on-line help, or `'help.start()'` for an HTML browser interface to help. Type `'q()'` to quit R.

[Workspace loaded from ~/.RData]

How you quit R from the console.

A “workspace” is where the stuff you do in R gets stored when you quit.

Start of RMarkdown Presentation

Expressions

Typing “1+1” on the console will cause R to evaluate the expression and tell us the results

```
> 1+1
```

```
[1] 2
```

The “[1]” before the answer is R’s line numbering scheme for reading long sets of numbers.

For example, if we were to type “letters” which is R’s built in variable with the letters of the alphabet:

```
> letters
```

```
[1] "a" "b" "c" "d" "e" "f" "g" "h" "i" "j" "k" "l"  
[13] "m" "n" "o" "p" "q" "r" "s" "t" "u" "v" "w" "x"  
[25] "y" "z"
```

Standard Math Stuff

Addition and subtraction symbols are what you would expect (+ and -). Multiplication and division are * and / respectively.

```
> 5-3
```

```
[1] 2
```

```
> 5*3
```

```
[1] 15
```

Order of Operations

Order of operations is as you might remember from math class. Multiplication and division before addition and subtraction. You can force the order of operations to be what you want by enclosing things you want done first in parentheses.

```
> 5 * 3 + 2
```

```
[1] 17
```

```
> 5 * (3 + 2)
```

```
[1] 25
```

Functions

Functions are how things get done in R. They take “arguments” as their input and output something useful (hopefully). R has a lot of built in functions. You can download and install packages to get access to more functions. You can easily write your own functions, but that won't be the focus of this course.

```
> sqrt(25)
```

```
[1] 5
```

Arguments, optional and required

```
log()
```

Error: argument "x" is missing, with no default

```
> log(100)
```

```
[1] 4.60517
```

```
> log(100, base=10)
```

```
[1] 2
```

Variable assignment

```
> x <- 2  
> x * 5
```

```
[1] 10
```


Variable Naming

If you name your variables things like “x” and “y”, chances are excellent you will soon forget what they mean. R variable names can be long, so make them as long as you need to explain what they are.

```
> plate_map.csv <- read.csv("plate_map.csv")  
> plate_map.csv.colnames <- colnames(plate_map.csv)
```

Basic “Atomic” Data types

For the purposes of this class, R has three base “types” of data (there are a few more we won’t concern ourselves with, like for complex numbers).

```
> class(12)
```

```
[1] "numeric"
```

```
> class("Susan")
```

```
[1] "character"
```

```
> class(TRUE)
```

```
[1] "logical"
```

Vectors

A vector is an ordered list of values in R, all of the same type.

```
> childAges <- c(8,10)
> childNames <- c("Bronwyn","Margot")
> childLikesMacAndCheese <- c(FALSE, TRUE)
> c(TRUE, "Bronwyn", 8)
```

```
[1] "TRUE"      "Bronwyn" "8"
```

```
> c(8,12,35,"no data",16,82)
```

```
[1] "8"          "12"         "35"         "no data" "16"
[6] "82"
```

Describing Data

Data can be continuous or discrete. Continuous data takes on values in a range. Discrete data can take on only a limited number of values. Its important to think about the meaning of the data, so that the way in which R stores it does not confuse the meaning of the data.

```
> myChromosomes <- c(1,3,8)
> myChromosomes
```

```
[1] 1 3 8
```

```
> factor(c(1,3,8), levels=1:22)
```

```
[1] 1 3 8
```

```
22 Levels: 1 2 3 4 5 6 7 8 9 10 11 12 13 14 15 ... 22
```

Ordinal Data

An ordinal variable is one that has a natural ordering to it. Whereas there is no natural ordering to eye colors (a nominal variable), there would be a natural ordering to grades.

```
> classGrades <- factor(c("B","D","A"),  
+                       levels=c("F","D","C","B","A"),  
+                       ordered=TRUE)  
> classGrades
```

```
[1] B D A
```

```
Levels: F < D < C < B < A
```

Telling R that our data is ordinal will help it do the right statistical tests and build models correctly.