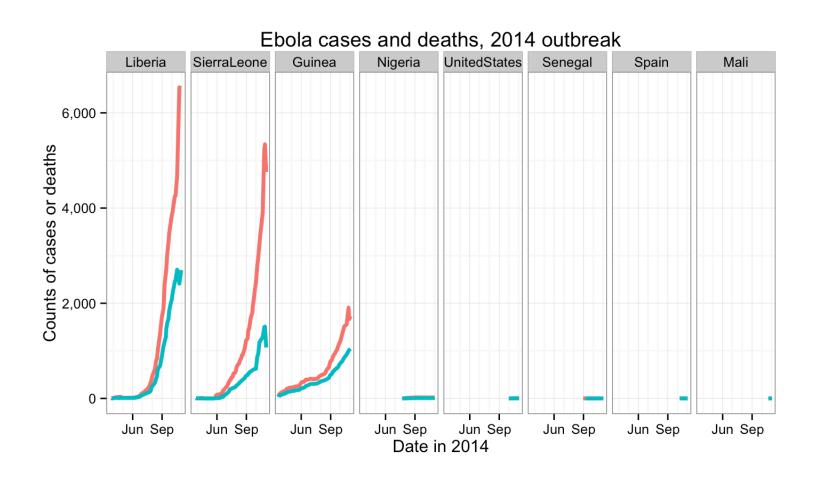
## Meeting 1: R and Ebola

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November 7, 2014

#### 2014 Ebola outbreak



#### First, catch your rabbit

GitHub

Caitlin River's repo of data for the 2014 Ebola outbreak:

Caitlin River's Ebola repo

#### First, catch your rabbit

#### Basic approach:

- Download data to your computer
- Make sure R is working in the directory with your data
- · Read data into R

#### Fancier approach:

Ask R to read data straight from GitHub

# Full details of basic approach for beginners

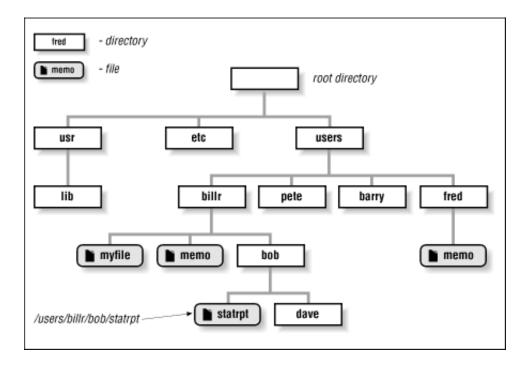
Step 1: Download the data to your computer.

To download a datafile from GitHub, check out the "Raw" button on the page...

Note: Make sure you save with the extension ".csv", and don't let your computer add on ".txt"

Step 2: Anytime you work in R, R will run from within a directory somewhere on your computer.

Let's review directories:



The default is usually your home directory (for example, mine is "/Users/brookeanderson"). You can easily change your working directory...

Once you open R, what is your working directory?

```
getwd()

## [1] "/Users/brookeanderson/FallRMeetings"

setwd("/Users/brookeanderson")
getwd()

## [1] "/Users/brookeanderson"
```

Check your working directory now:

```
getwd()
## [1] "/Users/brookeanderson/FallRMeetings"
```

If you're in the right directory, you should see our data file if you list the files in the directory:

```
list.files()
```

```
## [1] "country_timeseries.csv" "EbolaCenters.csv"
## [3] "ebolavirus_cds.nex" "Figures"
## [5] "Meeting1Code.r" "Meeting1Notes_files"
## [7] "Meeting1Notes.html" "Meeting1Notes.Rmd"
## [9] "README.md"
```

Step 3: Now we can read the data into R. It's a very basic command:

```
ebola <- read.table("country timeseries.csv", sep = ",",
                    header = TRUE)
ebola[1:3, 1:5]
##
          Date Day Cases Guinea Cases Liberia Cases SierraLeone
## 1 11/2/2014 225
                            1731
                                            NA
                                                            4759
## 2 10/31/2014 222
                              NA
                                          6525
                                                              NA
                         1667
## 3 10/29/2014 220
                                            NA
                                                            5338
```

But why does this work?

R can read in data from *a lot* of different formats. The only catch: you need to tell R how to do it.

Most basically, we'll look at flat files:

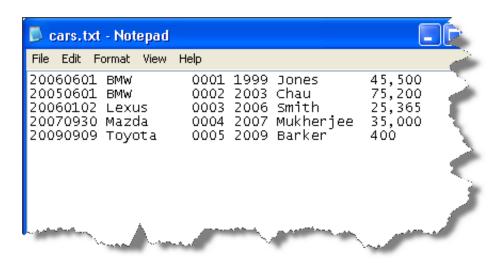
Fixed width files

Delimited files

- · ".csv": Comma-separated values
- ".tab", ".tsv": Tab-separated values
- Other possible delimiters: colon, semicolon, pipe (" | ")

See if you can identify what types of files the following files are...

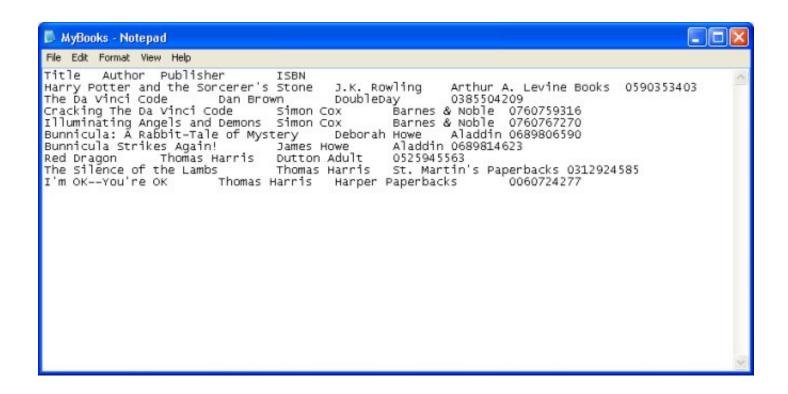
```
🌌 Bluestem_sample.txt - Note... 📮 🔲 🗶
File Edit Format Help
Species, Long, Lat
Bluestem, -94.6434,37.4556
Bluestem, -96.1527, 37.1606
Bluestem, -96.1493,37.0058
Bluestem,-94.8099,37.4403
Bluestem, -94.7353,37.0236
Bluestem, -94.6289,37.051
Bluestem, -94.7353,37.0381
Bluestem, -94.6108,37.0365
Bluestem, -94.672,37.0236
Purplestem, -94.8439,37.2779
Purplestem, -95.0871,37.1449
Purplestem, -95.217,37.4486
Purplestem, -94.6343,37.4556
Purplestem, -94.6525, 37.4556
Purplestem, -94.7859, 37.3685
```



```
H|20110606|pizza.txt|
D|10|chicken Pesto|20|23|30|5.5|7.4|9.9||
D|10|Meatball|10|53|60|6.5|8.4|10.9|
D|10|Fire Cracker|3|13|60|5.8|7.9|11.9|
D|10|Spinach|1|2|5|5.5|7.0|8.8|
D|10|BBQ Chicken|35|102|95|6.5|7.9|10.9|
D|10|Vegetarian|5|13|28|4.5|7.9|9.5|
D|10|Mexican|11|33|36|5.5|7.4|9.9|
D|10|The Monaco|22|53|7|5.5|7.5|8.9|
D|10|Chilli Prawn|5|5|6|5.5|7.4|9.9|
D|10|Chefs Special|8|18|40|5.8|7.8|9.8|
D|10|Marinara|3|17|41|5.5|7.4|9.0|
D|10|Supreme|50|52|58|5.5|7.4|9.2|
D|10|Margherita|9|19|87|5.0|7.0|8.0|
D|10|Napoli|60|85|66|5.2|7.2|9.2|
D|10|Ham and Pineapple|18|39|28|5.8|7.0|9.0|
```



```
File Edit Format View Help
Title, Subtitle, Larger Work, Contributor #1, Contributor #2, Contributor #3. Contributor
#4, Genre, Publisher, Published Location, Date
Published, Instrumentation, Key, Location, Indiana Connection, Sheet Music
Consortium, Notes, Complete
"""A"" You're Adorable", The alphabet song, , Buddy Kaye, Sidney Lippman, Fred Wise, , Popular
standard, Laurel Music Corporation, "New York, NY", 1948, Voice and
piano/guitar or ukulele,C Major,,None,Yes,Pérry Como pictured on cover,
"Aba Daba Honey Moon, The",,"""Two Weeks with Love"" Motion Picture",Arthur Fields,Walter
Donovan,,,"Popular Standard, Movie Selection",Leo Feist Inc.,"New
York, NY",1942, Voice and Piano, C Minor, None, Yes,
Abi Bezunt,,"""Mamele"" Motion Picture", Abraham Ellstein, Molly Picon,,, "Popular Standard,
Movie Selection", Metro Music Co., "New York, NY", 1939, Voice and
Piano, E Minor, , None, No, Molly Picon pictured on cover,
Abdul the Bulbul Ameer,,,Bob Kaai,Jim Smock,,,Popular Standard,Calumet Music Co.
,"Chicago, IL",1935,"Voice, Piano, Hawaiian Guitar, Ukulele",G
Major,,None,Yes,Ben Pollack pictured on cover,
About A Quarter to Nine,,""Go Into Your Dance" Motion Picture",Harry Warren,Al
Dubin,,,"Popular Standard, Movie Selection",M. Witmark & Sons, "New York,
NY",1935, "Voice, Piano, Guitar, Ukelele",E Minor,,None,No,Al Jolson and Ruby Keeler
pictured on cover.
Absent,,, John. W. Metcalf, Catherine Young Glen,,, Popular Standard, Arthur P.
Schmidt, "Boston, MA", 1899, Voice and Piano, G Major, None, Yes,,
The Academy Two-Step,,,Barclay Walker,,,,Popular Standard, Carlin & Lennox. "Indianapolis.
IN", Piano, F Major, Composer, No,,
Ac-cent-tchu-ate the Positive, Mister In Between, """Here Come the Waves"" Motion
Picture", Harold Arlen, Johnny Mercer, , , "Popular Standard, Movie
Selection", Edwin H. Morris & Co., "New York, NY", 1944, "Voice, Piano, Guitar", F
Major, None, Yes, Bing Crosby and Betty Hutton pictured on cover,
Across the Alley From the Alamo,,, Joe Greene,,,, Popular Standard, Leslie Music
```



1000233 1000234 1000235 1000236 1000237 1000238 1000240 1000241 1000242 1000244 1000245 1000246 1000247 1000248 1000249 1000250	Miralda Faley Baylog Gallardo Christian Baufield Frazier Garrido Williams Morel Padilla Rosenberg Blanchard Wiggins Miller Coon Chretien	John Nick Cathy Mike Daniel Robert Edward Zachary David Damian Wayne Phong S David Jeffrey Terry Walter Timothy
1000233 1000234 1000235	Miralda Faley Baylog	John Nick Cathy

R can read any of these types of files using one of the read.table and read.fwf functions. Find out more about those functions with:

?read.table
?read.fwf

Now let's read the data in and assign it (<-) to an object named ebola:

Notice that the function is read.table, and we've specified a value for the options sep and header. We'll talk about functions and options more later...

Question for you: What would have happened if we hadn't assigned the data we were reading in to ebola?

If this worked, you should have an object in your R session named ebola. You can check out the beginning of it:

```
ebola[1:3, 1:5]
```

```
## Date Day Cases_Guinea Cases_Liberia Cases_SierraLeone
## 1 11/2/2014 225 1731 NA 4759
## 2 10/31/2014 222 NA 6525 NA
## 3 10/29/2014 220 1667 NA 5338
```

There are a number of other functions you can use to check out your data. For example, try:

```
head(ebola)
tail(ebola)
summary(ebola)
str(ebola)
ebola[1,]
```

### Details of fancier approach

But, wait. **Everyone** is using GitHub for sharing data now. Surely there's a simpler way??

This is R. Of course there is!

If you don't know how to do it, try Googling:

"read data into r github"

Check out the first result:

"data- Read a CSV from github into R - Stack Overflow"

It turns out that you can read it straight from GitHub using the RCurl package.

If you don't have the package yet, you'll need to install it:

```
install.packages("RCurl")
```

Then, to use it in your R session, you'll need to call it:

```
library(RCurl)
```

Now you have all the RCurl tools available in your session.

Now all it takes to read the data in is:

```
github.page <- getURL("https://raw.githubusercontent.com/cmrivers/ebola/master/com/cmrivers/ebola/master/com/cmrivers/ebola/master/com/cmrivers/ebola/master/com/cmrivers/ebola/master/com/cmrivers/ebola/master/com/cmrivers/ebola/master/com/cmrivers/ebola/master/com/cmrivers/ebola/master/com/cmrivers/ebola/master/com/cmrivers/ebola/master/com/cmrivers/ebola/master/com/cmrivers/ebola/master/com/cmrivers/ebola/master/com/cmrivers/ebola/master/com/cmrivers/ebola/master/com/cmrivers/ebola/master/com/cmrivers/ebola/master/com/cmrivers/ebola/master/com/cmrivers/ebola/master/com/cmrivers/ebola/master/com/cmrivers/ebola/master/com/cmrivers/ebola/master/com/cmrivers/ebola/master/com/cmrivers/ebola/master/com/cmrivers/ebola/master/com/cmrivers/ebola/master/com/cmrivers/ebola/master/com/cmrivers/ebola/master/com/cmrivers/ebola/master/com/cmrivers/ebola/master/com/cmrivers/ebola/master/com/cmrivers/ebola/master/com/cmrivers/ebola/master/com/cmrivers/ebola/master/com/cmrivers/ebola/master/com/cmrivers/ebola/master/com/cmrivers/ebola/master/com/cmrivers/ebola/master/com/cmrivers/ebola/master/com/cmrivers/ebola/master/com/cmrivers/ebola/master/com/cmrivers/ebola/master/com/cmrivers/ebola/master/com/cmrivers/ebola/master/com/cmrivers/ebola/master/com/cmrivers/ebola/master/com/cmrivers/ebola/master/com/cmrivers/ebola/master/com/cmrivers/ebola/master/com/cmrivers/ebola/master/com/cmrivers/ebola/master/com/cmrivers/ebola/master/com/cmrivers/ebola/master/com/cmrivers/ebola/master/com/cmrivers/ebola/master/com/cmrivers/ebola/master/com/cmrivers/ebola/master/com/cmrivers/ebola/master/com/cmrivers/ebola/master/com/cmrivers/ebola/master/com/cmrivers/ebola/master/com/cmrivers/ebola/master/com/cmrivers/ebola/master/com/cmrivers/ebola/master/com/cmrivers/ebola/master/com/cmrivers/ebola/master/com/cmrivers/ebola/master/com/cmrivers/ebola/master/com/cmrivers/ebola/master/com/cmrivers/ebola/master/com/cmrivers/ebola/master/com/cmrivers/ebola/master/com/cmrivers/ebola/master/com/cmrivers/ebola/master/com/cmrivers/ebola/master/com/cmriver/
```

Note: It runs off the page, but the full "https" for the geturl function is just the web address of the raw data we want from GitHub:

ebola.2[1:3, 1:5]

```
## Date Day Cases_Guinea Cases_Liberia Cases_SierraLeone
## 1 11/2/2014 225 1731 NA 4759
## 2 10/31/2014 222 NA 6525 NA
## 3 10/29/2014 220 1667 NA 5338
```

#### What kind of data can you get into R?

The sky is the limit...

- Tables on webpages (e.g., the table near the end of this page)
- Files from other statistical packages (SAS, Excel, Stata, SPSS)
- Data in a database (e.g., SQL)
- Really crazy data formats used in other disciplines (e.g., netCDF files from climate folks, MRI data stored in Analyze, NIfTI, and DICOM formats)
- Data through APIs (e.g., GoogleMaps, Twitter)
- Incrediably messy data using scan and readLines

Find out more in Chapter 3 of The R Book.

## Challenges for the more advanced

#### Challenge 1

GitHub user BrcMapsTeam has geojson data on the locations of Ebola medical centers in West Africa as well as a link to a GoogleDocs dataset with the same information. See if you can get this data into R from one of these two sources.

#### Challenge 2

GitHub user evogytis has a repo called ebolaGuinea2014 with .nex files with protein coding sequences from Ebola genomes. He ultimately is using this data for phylogenetic analysis of the Ebola outbreak in Guinea. See if you can read one of these files into R. Also, see if you can figure out what tools exist for doing phylogenetic analysis in R.