Meeting 2: R and Ebola

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Last time, we read in our data

##	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

Today's plan

- Dataframes and vectors
- Subsetting
- Functions
- · Example of a function the plot function

A vector is a string of values:

Example 1: Start of the vector of the dates when Ebola cases were reported

```
## [1] "11/2/2014" "10/31/2014" "10/29/2014" "10/27/2014" "10/25/2014"
## [6] "10/22/2014" "10/21/2014" "10/19/2014" "10/18/2014" "10/14/2014"
```

Example 2: Start of the vector of number of cases reported in Guinea

```
## [1] 1731 NA 1667 1906 NA NA 1553 1540 NA 1519
```

You can make a new vector using the concatenation vector, c(...):

```
x <- c(1, 5, 7, 9, 10)
x

## [1] 1 5 7 9 10

class.names <- c("Taylor", "Maggie", "Mimi", "Brianna", "Jon")
class.names

## [1] "Taylor" "Maggie" "Mimi" "Brianna" "Jon"</pre>
```

A dataframe is made up of a lot of vectors stuck together (Notice how each column is a vector)

##	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
## 4	10/27/2014	218	1906	NA	5235
## 5	10/25/2014	216	NA	6535	NA
## 6	10/22/2014	214	NA	NA	3896
## 7	10/21/2014	213	1553	NA	NA
## 8	10/19/2014	211	1540	NA	3706
## 9	10/18/2014	210	NA	4665	NA
## 10	10/14/2014	206	1519	NA	3410

You can make a new dataframe using the function data.frame():

```
## name number
## 1 Taylor 1
## 2 Maggie 5
## 3 Mimi 7
## 4 Brianna 9
## 5 Jon 10
```

You can use indexing ([...], [..., ...]) to subset from a vector or dataframe, like:

```
vector[locations] ## Generic code
dataframe[row locations, column locations] ## Generic code
```

A vector has one dimension, so you index without a comma (i.e., in one dimension):

```
class.names[1]

## [1] "Taylor"

class.names[c(2, 3, 4)] ## Equivalent: class.names[2:4]

## [1] "Maggie" "Mimi" "Brianna"
```

A dataframe has two dimensions (rows and columns), so you index with a column:

```
class.data[1,1]

## [1] Taylor
## Levels: Brianna Jon Maggie Mimi Taylor

class.data[1:3, 1:2]

## name number
## 1 Taylor 1
## 2 Maggie 5
## 3 Mimi 7
```

To get all values in a dimension (row or column), leave that part of the index blank:

```
class.data[1, ]

## name number

## 1 Taylor 1

class.data[ , 1]

## [1] Taylor Maggie Mimi Brianna Jon

## Levels: Brianna Jon Maggie Mimi Taylor
```

For columns, you can use column names instead of location:

```
class.data[3:4, "number"]

## [1] 7 9

class.data[3:4, c("name", "number")]

## name number

## 3 Mimi 7

## 4 Brianna 9
```

You can also pull a column (vector) from a dataframe using \$, like:

```
dataframe$column.name ## Generic code
```

For example, to get the column of ebola with cases from Guinea:

```
head(ebola$Cases_Guinea)
```

```
## [1] 1731 NA 1667 1906 NA NA
```

Note: I've used head to look at just the start of the vector since the whole thing would be really long.

Now you try...

Try to get the following vectors from the dataset:

- · Date
- The ten most recent counts of cases in the US
- The earliest twenty counts of deaths in Liberia
- A dataframe of the first five observations of date, cases in Mali and deaths in Mali

Hint: Try using colnames (ebola) to find out the names of all the columns in ebola. Also, use dim(ebola) to find out the dimensions of the dataframe so you can get the index numbers right for the latest ten data points.

Functions

Functions

In general, functions in R take the following structure:

```
function.name(required information, options) ## Generic code
```

The result of the function will be output to your R session, unless you choose to save the output in an object:

```
new.object <- function.name(required information, options) ## Generic code</pre>
```

Functions

Examples of this structure:

Find out more about a function by using ?function.name. This will take you to the help page for the function, where you can find out all the possible arguments for the function, required and optional.

Example of a function

The plot function has two required arguments: the x coordinates of points in the plot, and the y coordinates of points in the plot. The generic structure is:

```
plot(x = x coordinates, y = y coordinate) ## Generic code
```

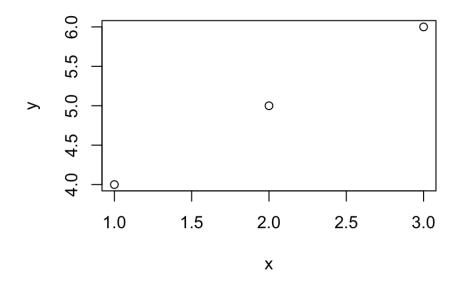
As long as you put the x coordinates first and the y coordinates second, you can leave out the x =and y =:

```
plot(x coordinates, y coordinate) ## Generic code
```

```
x <- c(1, 2, 3)

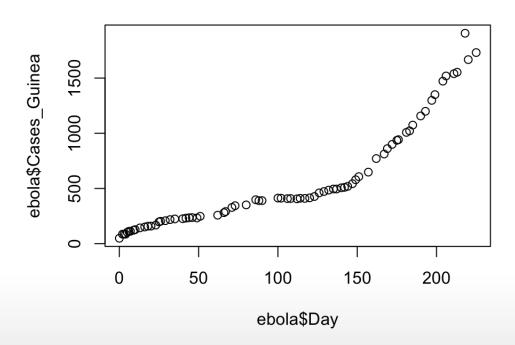
y <- c(4, 5, 6)

plot(x, y)
```



Now that you know how to pull out two vectors you want from the ebola dataset, you can plot them:

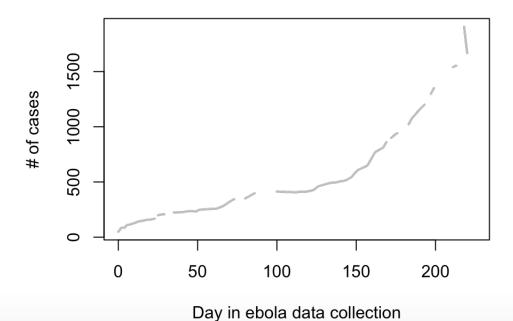
plot(ebola\$Day, ebola\$Cases_Guinea)



The plot function also has *many* optional arguments (check ?plot, ? plot.default). For example,

- type: What do you want to plot? Points ("p")? Lines ("1")?
- main: Give a title to your plot ("My title")
- * xlab, ylab: Give nicer labels to your x- and y-axes (xlab = "Day
 in Ebola data collection")
- * xlim, ylim: Specify the range of your x- and y- axes (xlim =
 c(0, 100))

Guinea ebola cases



Now you try...

Try plotting:

- Deaths in Liberia by day
- Mortality rate in Liberia by day
- Deaths in Liberia by date

Experiment with options like type; col, pch and cex (when you're plotting points, type = "p"), lwd (when you're plotting lines, type = "l"), main, sub, xlim, and ylim.

Hint: Try using colnames (ebola) to find out the names of all the columns in ebola.