Stats workshop

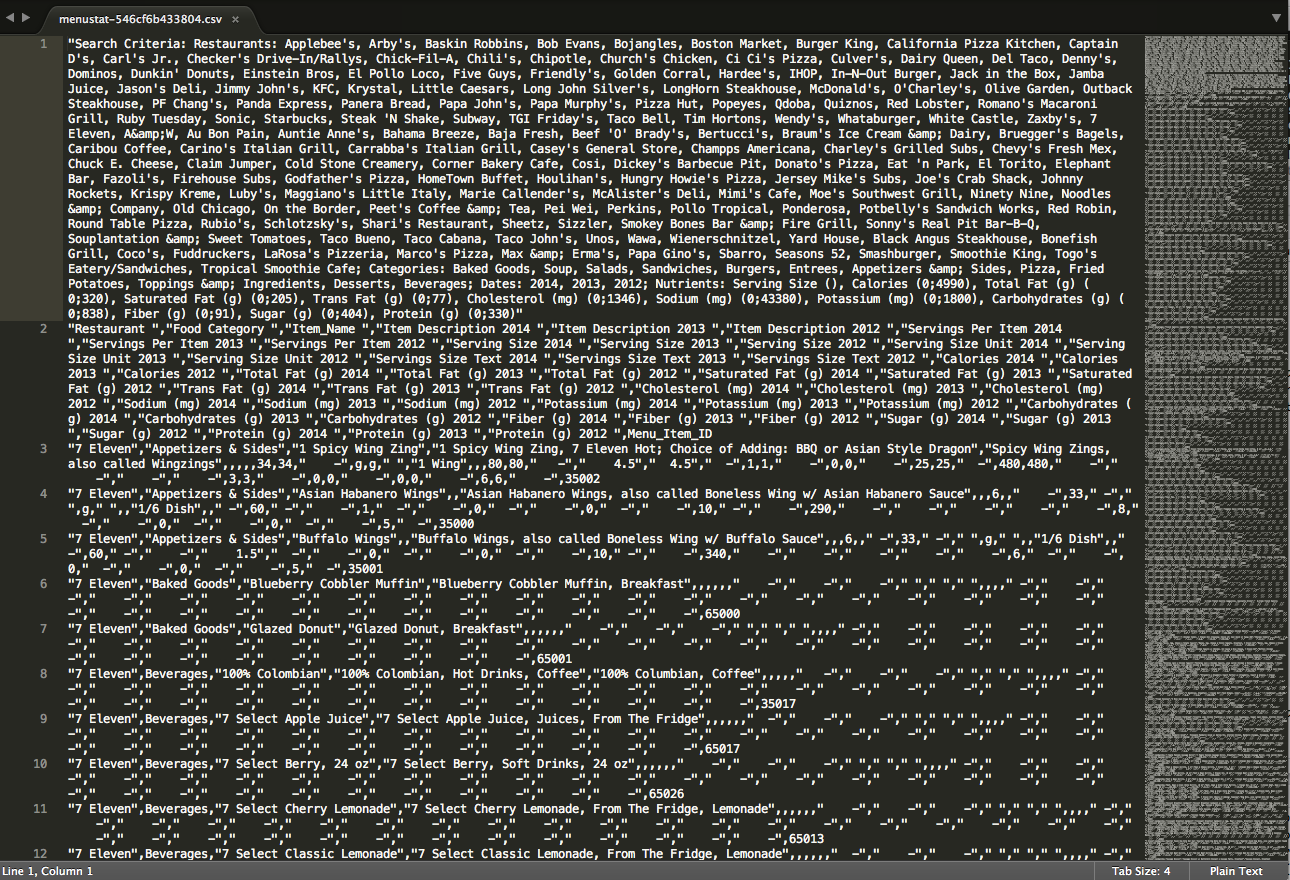
## Old Section

**I resisted deleting this. It isn't necessary for what I'm under the impression ya'll want. But maybe you'll still find it interesting or fun. I've commented out all the commands because some of them take a while to run. If you want to run it yourself, just remove the comment tags (#) on all lines with the exception of the first code chunk. The first one results in an error and will choke knitr. If you don't care, you can skip down to the part that says "Mark's Data"**

I've put together a document which should guide the work we're doing in our stats workshop. This is an applied statistics approach in which we will emphasize the day-to-day work of data analysis. First, we will need some data! Where do we get it? The typical approach in academic psychology is to run some kind of study. This isn't true for every discipline, and arguably, our own discipline doesn't take enough advantage of the data which can be found free for the taking in the real world. For example, I took the following data from [menustat.org](http://www.menustat.org). Let's try to read it in.

#menus<-read.csv('menustat-546cf6b433804.csv')  
#this doesn't work. Throws an error:  
  
#Error in read.table(file = file, header = header, sep = sep, quote = quote, :   
# more columns than column names

Aw damn. Well, that didn't work. Why not? The error message says that there's a mismatch between the number of columns, and the number of names it has for the columns. Usually, this function reads the column names from the first line of the csv document, so I'm guessing that something is wrong with the first line of the csv. Opening it up in a text editor shows the following:



See the big chunk of text next to the 1 on the lefthand side? That's all the stuff that appears in the first line, and we can see that it's just kind of a description of the file. We can get rid of that. Line 2 seems to have the columns we want. So we're going to take a slightly different approach:

1. Read the file, line by line as one big vector
2. Remove the first line.
3. Turn the vector into a dataframe.

#menus<-readLines('menustat-546cf6b433804.csv')  
#menus<-menus[-1]  
#menus<-read.csv(textConnection(menus)) #this takes a minute...

Now we have a datafile for our menus data! This is a new datafile for all of us, so let's just get some information about it.

#dim(menus)

Serious data, ya'll. We've got 60,238 observations across 52 different variables. Let's get a little information about those variables.

#names(menus)

We can now see the names for our variables. Looks like there's one for restaurant, one for food category, one for the item, and then we have a bunch of variables which are repeated for each of 3 years: 2014, 2013, 2012. These variables describe the item, give us a serving size, and then the nutrient information (e.g. calories, fat, carbs, protein, etc.).

## Mark's Data

This wont just be a *lesson* about what to do when one has new data. This will be an actual example of what I would do when I have new data. Mark has sent me some of his data and I haven't done anything but download it yet. So, this is actually me playing around with new data. Let's see what I'm gonna do (even I don't know!). =)

#### A quick digression on organization

This doesn't matter much at first, but once you have multiple projects, and are collaborating with multiple people, and these projects drag out over months years, organization can really become difficult to maintain. There are ways to keep organized, however! I recently found a nice, short, readable [guide](http://faculty.chicagobooth.edu/matthew.gentzkow/research/CodeAndData.pdf) on all the stuff I've spent a while figuring out on my own. Some of what you will find in there you will find more or less relevant (check out chapters 3 and 4 in particular), but I'll emphasize a couple of things:

1. Use version control. [Github](https://github.com/) is easy to use, especially when you download their [gui](https://mac.github.com/). You'll never have to worry about having multiple, conflicting, unclear analysis files again for fear of losing something you wrote previously.
2. Write your analysis code so that you can run the whole thing from start to finish. It helps even more if you write it in [markdown](http://rmarkdown.rstudio.com/) (or something similar) with prose explaining what you're doing and what you're thinking. From there, it's only a short step to turn it into a methods/results section.

Of course, I should put caveats all around this thing because I often don't really know what I'm doing either. I've never had a data set where I was doing all the same old things. It's a constant process of learning something new. And while I've been fighting the organization war for years, it's only in the last 6 months or so that I've really decided that this was something I had to win. =)

### Get the data in and look at what we've got

First, let's get it in. I'm sure you're all familiar with how to do this. However, it helps to keep things organized. Everything related to this project should all be in the same folder. I've put Mark's data in the same folder I'm keeping this script. Before doing anything, you should click Session -> Set Working Directory -> To Source File Location.

I wish there were an automatic way to do that. Unfortunately, I don't think there is. Anyway, once that's done, we can move on to read in his data and get the dimensions.

breach <- read.csv('breachmeans.csv')  
dim(breach)

## [1] 20 32

Okay! We've got 20 observations for each of 32 variables. Let's look a little more closely at what those variables are. Next, I'm going to get the names of each variable:

names(breach)

## [1] "subj" "locomotion1" "locomotion2" "difflocomotion"  
## [5] "assessment1" "assessment2" "diffassessment" "promotion1"   
## [9] "promotion2" "diffpromotion" "prevention1" "prevention2"   
## [13] "diffprevention" "stress1" "stress2" "diffstress"   
## [17] "flourishing" "swls" "lie2" "lie1"   
## [21] "diffLie" "LxA1" "LxA2" "diffLxA"   
## [25] "age" "exp" "apride" "lpride"   
## [29] "eom" "rankindiv" "rank21" "averank"

Now, a bit of a closer look at what these things contain:

head(breach)

## subj locomotion1 locomotion2 difflocomotion assessment1 assessment2  
## 1 1 5.00 4.58 -0.4167 4.17 3.67  
## 2 2 4.50 4.58 0.0833 4.25 3.92  
## 3 3 5.25 4.83 -0.4167 4.00 4.00  
## 4 4 4.83 5.00 0.1667 4.00 3.83  
## 5 5 5.08 4.92 -0.1667 4.50 3.83  
## 6 6 5.08 4.75 -0.3333 5.00 4.08  
## diffassessment promotion1 promotion2 diffpromotion prevention1  
## 1 -0.5000 3.50 4.00 0.500 1.8  
## 2 -0.3333 4.00 4.17 0.167 2.2  
## 3 0.0000 4.17 3.83 -0.333 1.0  
## 4 -0.1667 4.00 4.50 0.500 2.2  
## 5 -0.6667 4.83 4.83 0.000 3.4  
## 6 -0.9167 4.00 4.33 0.333 2.4  
## prevention2 diffprevention stress1 stress2 diffstress flourishing swls  
## 1 2.2 0.4 1.62 1.50 -0.125 6.50 6.0  
## 2 2.2 0.0 2.00 2.50 0.500 5.25 5.8  
## 3 1.4 0.4 2.38 2.25 -0.125 6.25 6.2  
## 4 2.0 -0.2 2.00 2.62 0.625 6.88 6.8  
## 5 3.0 -0.4 2.00 2.00 0.000 6.25 6.2  
## 6 2.2 -0.2 1.75 2.12 0.375 6.00 5.6  
## lie2 lie1 diffLie LxA1 LxA2 diffLxA age exp apride lpride eom rankindiv  
## 1 1.67 1.83 -0.167 25.0 21.0 -3.993 23 3 37 46 40 10  
## 2 2.67 2.33 0.333 20.2 21.0 0.757 26 6 38 38 35 4  
## 3 2.00 2.67 -0.667 27.6 23.4 -4.201 22 4 32 40 43 6  
## 4 2.33 3.00 -0.667 23.4 25.0 1.639 26 2 40 46 42 16  
## 5 1.83 1.83 0.000 25.8 24.2 -1.667 26 4 45 44 38 3  
## 6 3.00 3.17 -0.167 25.8 22.6 -3.278 29 3 38 44 34 5  
## rank21 averank  
## 1 13.0 11.50  
## 2 3.5 3.75  
## 3 5.0 5.50  
## 4 16.0 16.00  
## 5 3.5 3.25  
## 6 7.0 6.00

summary(breach)

## subj locomotion1 locomotion2 difflocomotion   
## Min. : 1.00 Min. :3.830 Min. :3.580 Min. :-1.7500   
## 1st Qu.: 5.75 1st Qu.:4.647 1st Qu.:4.560 1st Qu.:-0.3542   
## Median :10.50 Median :5.000 Median :4.920 Median :-0.2084   
## Mean :10.50 Mean :4.949 Mean :4.816 Mean :-0.1333   
## 3rd Qu.:15.25 3rd Qu.:5.122 3rd Qu.:5.270 3rd Qu.: 0.2708   
## Max. :20.00 Max. :5.750 Max. :5.420 Max. : 0.5000   
##   
## assessment1 assessment2 diffassessment promotion1   
## Min. :3.750 Min. :3.500 Min. :-1.3333 Min. :3.500   
## 1st Qu.:4.000 1st Qu.:3.830 1st Qu.:-0.5417 1st Qu.:3.830   
## Median :4.290 Median :3.920 Median :-0.3333 Median :4.000   
## Mean :4.351 Mean :4.017 Mean :-0.3333 Mean :4.058   
## 3rd Qu.:4.543 3rd Qu.:4.122 3rd Qu.:-0.1459 3rd Qu.:4.210   
## Max. :5.170 Max. :4.920 Max. : 1.1667 Max. :4.830   
##   
## promotion2 diffpromotion prevention1 prevention2   
## Min. :3.170 Min. :-0.83300 Min. :1.00 Min. :1.4   
## 1st Qu.:3.830 1st Qu.:-0.33300 1st Qu.:2.20 1st Qu.:2.2   
## Median :4.000 Median : 0.08350 Median :2.60 Median :2.7   
## Mean :4.041 Mean :-0.01665 Mean :2.53 Mean :2.6   
## 3rd Qu.:4.330 3rd Qu.: 0.33300 3rd Qu.:3.00 3rd Qu.:3.0   
## Max. :4.830 Max. : 0.50000 Max. :3.80 Max. :3.4   
##   
## diffprevention stress1 stress2 diffstress   
## Min. :-0.80 Min. :1.380 Min. :1.000 Min. :-0.75000   
## 1st Qu.:-0.25 1st Qu.:1.685 1st Qu.:1.718 1st Qu.:-0.12500   
## Median : 0.20 Median :1.880 Median :2.000 Median : 0.00000   
## Mean : 0.07 Mean :2.013 Mean :2.068 Mean : 0.08553   
## 3rd Qu.: 0.40 3rd Qu.:2.250 3rd Qu.:2.530 3rd Qu.: 0.37500   
## Max. : 0.80 Max. :3.500 Max. :3.120 Max. : 0.62500   
## NA's :1 NA's :1   
## flourishing swls lie2 lie1   
## Min. :4.880 Min. :4.40 Min. :1.000 Min. :1.000   
## 1st Qu.:5.750 1st Qu.:5.75 1st Qu.:2.290 1st Qu.:2.290   
## Median :6.000 Median :5.80 Median :2.670 Median :2.500   
## Mean :5.976 Mean :5.91 Mean :2.675 Mean :2.632   
## 3rd Qu.:6.250 3rd Qu.:6.20 3rd Qu.:3.042 3rd Qu.:3.000   
## Max. :7.000 Max. :7.00 Max. :4.670 Max. :4.500   
##   
## diffLie LxA1 LxA2 diffLxA   
## Min. :-0.6670 Min. :14.70 Min. :12.80 Min. :-16.479   
## 1st Qu.:-0.2085 1st Qu.:21.60 1st Qu.:20.80 1st Qu.: -3.790   
## Median : 0.0000 Median :25.00 Median :24.20 Median : -1.760   
## Mean : 0.0416 Mean :24.72 Mean :23.45 Mean : -1.258   
## 3rd Qu.: 0.2085 3rd Qu.:26.25 3rd Qu.:27.80 3rd Qu.: 2.675   
## Max. : 1.0000 Max. :33.10 Max. :29.30 Max. : 5.167   
##   
## age exp apride lpride   
## Min. :21.0 Min. :0.00 Min. :28.00 Min. :29.00   
## 1st Qu.:24.0 1st Qu.:2.00 1st Qu.:36.00 1st Qu.:38.00   
## Median :26.0 Median :3.00 Median :38.00 Median :41.50   
## Mean :25.9 Mean :3.25 Mean :37.45 Mean :41.35   
## 3rd Qu.:28.0 3rd Qu.:4.00 3rd Qu.:39.00 3rd Qu.:46.00   
## Max. :31.0 Max. :8.00 Max. :46.00 Max. :49.00   
##   
## eom rankindiv rank21 averank   
## Min. :34.00 Min. : 1.00 Min. : 1.00 Min. : 1.500   
## 1st Qu.:37.00 1st Qu.: 5.75 1st Qu.: 5.75 1st Qu.: 5.875   
## Median :38.50 Median :10.50 Median :11.00 Median :10.500   
## Mean :38.45 Mean :10.50 Mean :11.00 Mean :10.750   
## 3rd Qu.:40.00 3rd Qu.:15.25 3rd Qu.:16.25 3rd Qu.:16.500   
## Max. :43.00 Max. :20.00 Max. :21.00 Max. :20.000   
##

These are quick ways to examine what your dataframe looks like. head will return the first 5 rows of the dataframe (you can also use tail to get the last 5 rows). summary, when called on a dataframe, will return a description of each variable. Mark's dataframe is a little unusual for psych studies in that it doesn't seem to have any factor variables. We can see this in another way by asking for the structure, using the str command.

str(breach)

## 'data.frame': 20 obs. of 32 variables:  
## $ subj : int 1 2 3 4 5 6 7 8 9 10 ...  
## $ locomotion1 : num 5 4.5 5.25 4.83 5.08 5.08 4.67 5 5.58 5.08 ...  
## $ locomotion2 : num 4.58 4.58 4.83 5 4.92 4.75 5.08 4.33 5.25 5.33 ...  
## $ difflocomotion: num -0.4167 0.0833 -0.4167 0.1667 -0.1667 ...  
## $ assessment1 : num 4.17 4.25 4 4 4.5 5 4 4.5 4.42 4.25 ...  
## $ assessment2 : num 3.67 3.92 4 3.83 3.83 4.08 3.5 4 3.67 3.92 ...  
## $ diffassessment: num -0.5 -0.333 0 -0.167 -0.667 ...  
## $ promotion1 : num 3.5 4 4.17 4 4.83 4 3.83 3.83 4 4.67 ...  
## $ promotion2 : num 4 4.17 3.83 4.5 4.83 4.33 4.17 3.5 3.83 4 ...  
## $ diffpromotion : num 0.5 0.167 -0.333 0.5 0 0.333 0.333 -0.333 -0.167 -0.667 ...  
## $ prevention1 : num 1.8 2.2 1 2.2 3.4 2.4 2.8 3 2.8 2.6 ...  
## $ prevention2 : num 2.2 2.2 1.4 2 3 2.2 3 2.8 3.2 2.8 ...  
## $ diffprevention: num 0.4 0 0.4 -0.2 -0.4 -0.2 0.2 -0.2 0.4 0.2 ...  
## $ stress1 : num 1.62 2 2.38 2 2 1.75 3.5 2.25 1.75 1.75 ...  
## $ stress2 : num 1.5 2.5 2.25 2.62 2 2.12 3.12 2.88 1.75 1.75 ...  
## $ diffstress : num -0.125 0.5 -0.125 0.625 0 0.375 -0.375 0.625 0 0 ...  
## $ flourishing : num 6.5 5.25 6.25 6.88 6.25 6 7 6 5.75 6 ...  
## $ swls : num 6 5.8 6.2 6.8 6.2 5.6 7 5.8 5.8 5.8 ...  
## $ lie2 : num 1.67 2.67 2 2.33 1.83 3 4.67 2.33 1 3.17 ...  
## $ lie1 : num 1.83 2.33 2.67 3 1.83 3.17 3.83 2.33 1 2.83 ...  
## $ diffLie : num -0.167 0.333 -0.667 -0.667 0 -0.167 0.833 0 0 0.333 ...  
## $ LxA1 : num 25 20.2 27.6 23.4 25.8 25.8 21.8 25 31.2 25.8 ...  
## $ LxA2 : num 21 21 23.4 25 24.2 22.6 25.8 18.8 27.6 28.4 ...  
## $ diffLxA : num -3.993 0.757 -4.201 1.639 -1.667 ...  
## $ age : int 23 26 22 26 26 29 28 24 27 22 ...  
## $ exp : int 3 6 4 2 4 3 0 3 4 1 ...  
## $ apride : int 37 38 32 40 45 38 38 33 36 37 ...  
## $ lpride : int 46 38 40 46 44 44 49 40 38 38 ...  
## $ eom : int 40 35 43 42 38 34 39 41 38 37 ...  
## $ rankindiv : int 10 4 6 16 3 5 12 9 11 19 ...  
## $ rank21 : num 13 3.5 5 16 3.5 7 21 10 8 15 ...  
## $ averank : num 11.5 3.75 5.5 16 3.25 6 16.5 9.5 9.5 17 ...

This highlights that within R, the dataframe is really just a collection of vectors. Mark has two types: numeric and integer.

##### Question:

**1. We wont always be fortunate enough to have .csv files. How can we read in excel files? Spss? STATA? Word?**

library(xlsx)  
read.xlsx()  
library(foreign)  
read.spss()  
read.dta()

Microsoft word is possible (everything is *possible*), but a bit more complicated (see tm package)

### Data types

From an R standpoint, there are few differences between numeric and integer types. It's good to know they exist, because the type of object you pass to functions can be a major source of headaches (e.g. if a function expects a factor and you try to pass it a numeric), but for now, we can generally treat numeric and integers as the same thing - quantitative variables. This can be contrasted with two other types that we do not see here: factors and characters. A factor is a categorical variable, and a character is just a collection of text strings. Let's go ahead and make a couple of these. First, we'll make a variable indicating whether someone is happy or not:

breach$happy <- rep(c("yes", "no"), each=10)  
breach$happy #there's only 20 observations, so we can just print out all of them.

## [1] "yes" "yes" "yes" "yes" "yes" "yes" "yes" "yes" "yes" "yes" "no"   
## [12] "no" "no" "no" "no" "no" "no" "no" "no" "no"

str(breach$happy)

## chr [1:20] "yes" "yes" "yes" "yes" "yes" "yes" "yes" ...

summary(breach$happy)

## Length Class Mode   
## 20 character character

Ah! A character variable! But that's just because we fed it some text strings, isn't it? As psychologists, we know that this is actually a categorical variable. People fall into one of two categories: Those who are happy, and those who have failed research projects. ;)

Anyway, since this is a categorical variable, we want this to be a factor.

breach$happy <- as.factor(breach$happy)  
breach$happy

## [1] yes yes yes yes yes yes yes yes yes yes no no no no no no no   
## [18] no no no   
## Levels: no yes

str(breach$happy)

## Factor w/ 2 levels "no","yes": 2 2 2 2 2 2 2 2 2 2 ...

summary(breach$happy)

## no yes   
## 10 10

You can see that calling some of the same functions on data which is identical in appearance but *stored* in two different ways can lead to some very different output. It's key to keep this in mind no matter what you're doing in R. If something doesn't work the way you expect it to, maybe it's because you're feeding in an unexpected data type. I think I spent about a month in my third year figuring this out. A long, painful month.

##### Questions:

1. Let's say Mark ran a study in which his participants recieved 2, 4, or 8 'happy' primes. We create a factor variable which distinguishes these groups:

* a <- as.factor(rep(c(2,4,8), each=10))
* For some purposes, this will be fine. But let's say for some reason we want to treat this as a numeric value rather than a factor. We run the following command:
* a <- as.numeric(a)
* What happened? Did it work as you expected?

1. How are factor variables represented internally in R (*hint*: Why, when we call str(breach$happy), do we get a bunch of numbers)? What's the logic behind their representation?

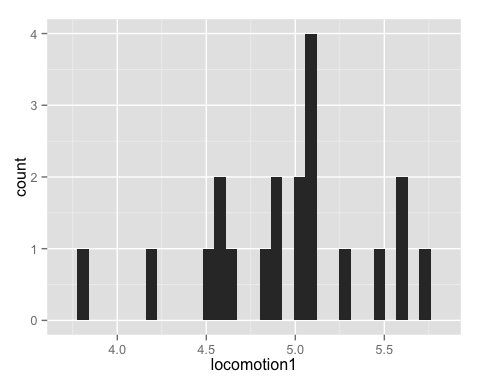
Now that we know a little bit about data types, let's think of what types of things we might be interested in looking at. It's time to know a bit about ggplot, don't you think?

### Preliminary visualizations

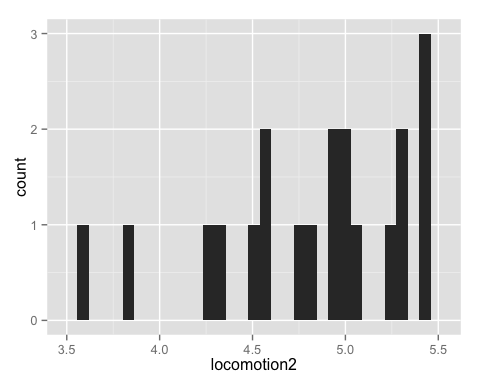
ggplot is a package built by the incredibly productive [Hadley Wickham](http://had.co.nz). It follows a general plotting philosophy published and described elsewhere called *Grammer of Graphics*. To create a ggplot object, you must first decide what you're plotting. Since I see that we have two scores for most of our measurements (plus I've spoken to Mark about this work a few times), I'm going to treat these as two measurements of the same concept on the same individual. One thing we obviously might be interested in is how these measurements changed between session one and session two.

Please note that there's no hypothesis testing going on yet! We're just looking at data! You should spend quite a lot of time doing this. Anyway, on with it. Let's make a couple of histograms for the locomotion variable.

library(ggplot2)  
plot <- ggplot(breach, aes(x=locomotion1))  
plot + geom\_histogram()



plot <- ggplot(breach, aes(x=locomotion2))  
plot + geom\_histogram()



Before we pull apart the figures, let's look at what we've done here. For each plot, there are two lines of code. The first line from the first plot:

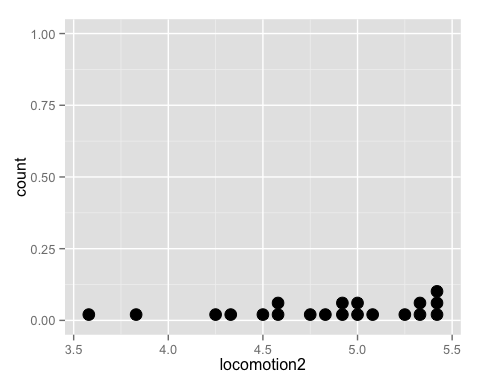
plot <- ggplot(breach, aes(x=locomotion1))

This makes an object we've called plot. This object is stored internally as a list, but we don't really care about that. What we care about is that this is a ggplot object made from the breach dataframe (1st argument). We've also assigned some aesthetics (via aes) for this particular plot. In particular, we've told it that we want the variable locomotion1 to appear on the x axis (2nd argument). This will create an empty plot, onto which we can begin placing layers. Each layer added to the plot will do something to change the superficial appearance. However, in general, after the first line, we've already given R **all** the data we need to make the figure we're going for. Now, all we need to do is specify exactly what we want the data to look like. The next line:

plot + geom\_histogram()

Here, we've taken the empty plot object, and added a layer. This layer takes the form of a histogram, but it could have just as easily been something else. There aren't very many layers which will work without some additional information (like a second variable), but just for kicks, here's an example of something else we *could* have done:

plot + geom\_dotplot()



We've taken the exact same underlying object (plot), and added a different layer. It's clear that the underlying data is the same, but this lets us visualize it in a slightly different way.

The original objective, however, was to compare the first locomotion score to the second. While we can do that with these plots, it isn't ideal for this purpose for a couple of reasons:

1. Having two separate plots forces us to shift our eyes back and forth between the two. It would be much better if they were plotted together.
2. The sparsity of our data leads our histogram to be not-very-smooth. Kind of chunky looking. I love chunky peanut butter, but histograms, not so much.

So, we can fix this by plotting both of these variables on the same object, letting them overlap somehow, and using a different kind of layer. But wait! How can we do this? Look back at the creation of the plotting object:

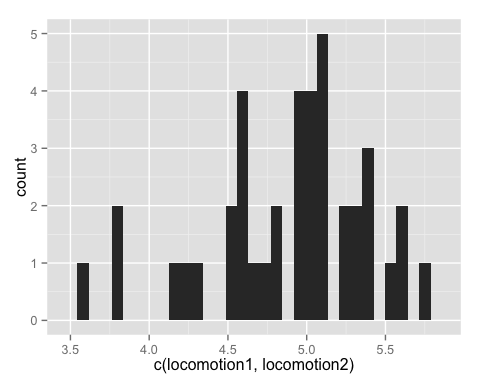
plot <- ggplot(breach, aes(x=locomotion1))

We've assigned a variable to x, but what we want is to have TWO variables on x. Your intuition may say to do this:

plot <- ggplot(breach, aes(x=c(locomotion1, locomotion2)))

Your intuition would be wrong. This will plot just fine, but it isn't what we want. Here, R has no way of knowing which scores are from locomotion1 and locomotion2, and so will just plot them all together. Observe:

plot <- ggplot(breach, aes(x=c(locomotion1, locomotion2)))  
plot + geom\_histogram()



*Side note: using c() will combine things into a vector. As soon as you perform c() on something, all the component parts are treated as coming from the same thing.*

We can get around this issue, but it requires some wrangling of our data. To preview, we will eventually aim to have two vectors in our dataframe: one of the values to be plotted along the x axis, and another indicating which of two groups each value belongs to.

##### Questions:

1. There are other reasons we might not like the plots we just produced. For example, maybe we want the bins into which the observations are sorted to be a bit larger. Can you do this? What setting do you think produces the best visualization for the data? (*hint*: ?geom\_histogram)
2. In the last plot, the x axis label is horrible. Change it to something more appropriate. You might find [this](http://docs.ggplot2.org/current/) helpful.

### Reshaping

We're going to reshape our data so that we can feed it into ggplot. Mark's data is currently organized in the 'wide' format. In psychology studies, this usually means that each row is an individual, and everything we know or have collected from that individual is on that row. For many statistical computing purposes, it is common to use a 'long' format. Here, the rows take the form of observations on an individual. Since we're observing many of our variables twice on the same individual, this implies that each individual will have two rows. If we had done an experiment with, let's say, 200 reaction time trials, then we would have 200 rows for each individual in a 'long' format.

In order to get this data from wide to long, we can use the reshape2 package, created by....yep, that's right. Our dear friend [Hadley Wickham](http://had.co.nz). (I know! This guy! Jeez! Everytime I load one of his packages I think about sending him one of the books on his [amazon wishlist](http://www.amazon.com/gp/registry/1Y65N3VFA613B)).

There are two main functions in the reshape2 package: melt and cast. The former takes wide data and converts it to long, and the latter does the converse. So we need to use melt.

library(reshape2)  
breach <- melt(breach,   
 id.vars = c('subj', 'flourishing', 'swls', 'age', 'exp', 'apride',   
 'lpride', 'eom', 'rankindiv', 'rank21', 'averank'),  
 measure.vars = c('locomotion1', 'locomotion2', 'assessment1', 'assessment2',   
 'promotion1', 'promotion2', 'prevention1', 'prevention2',   
 'stress1', 'stress2', 'lie2', 'lie1', 'LxA1', 'LxA2'))  
str(breach)

## 'data.frame': 280 obs. of 13 variables:  
## $ subj : int 1 2 3 4 5 6 7 8 9 10 ...  
## $ flourishing: num 6.5 5.25 6.25 6.88 6.25 6 7 6 5.75 6 ...  
## $ swls : num 6 5.8 6.2 6.8 6.2 5.6 7 5.8 5.8 5.8 ...  
## $ age : int 23 26 22 26 26 29 28 24 27 22 ...  
## $ exp : int 3 6 4 2 4 3 0 3 4 1 ...  
## $ apride : int 37 38 32 40 45 38 38 33 36 37 ...  
## $ lpride : int 46 38 40 46 44 44 49 40 38 38 ...  
## $ eom : int 40 35 43 42 38 34 39 41 38 37 ...  
## $ rankindiv : int 10 4 6 16 3 5 12 9 11 19 ...  
## $ rank21 : num 13 3.5 5 16 3.5 7 21 10 8 15 ...  
## $ averank : num 11.5 3.75 5.5 16 3.25 6 16.5 9.5 9.5 17 ...  
## $ variable : Factor w/ 14 levels "locomotion1",..: 1 1 1 1 1 1 1 1 1 1 ...  
## $ value : num 5 4.5 5.25 4.83 5.08 5.08 4.67 5 5.58 5.08 ...

summary(breach)

## subj flourishing swls age   
## Min. : 1.00 Min. :4.880 Min. :4.40 Min. :21.0   
## 1st Qu.: 5.75 1st Qu.:5.750 1st Qu.:5.75 1st Qu.:24.0   
## Median :10.50 Median :6.000 Median :5.80 Median :26.0   
## Mean :10.50 Mean :5.976 Mean :5.91 Mean :25.9   
## 3rd Qu.:15.25 3rd Qu.:6.250 3rd Qu.:6.20 3rd Qu.:28.0   
## Max. :20.00 Max. :7.000 Max. :7.00 Max. :31.0   
##   
## exp apride lpride eom   
## Min. :0.00 Min. :28.00 Min. :29.00 Min. :34.00   
## 1st Qu.:2.00 1st Qu.:36.00 1st Qu.:38.00 1st Qu.:37.00   
## Median :3.00 Median :38.00 Median :41.50 Median :38.50   
## Mean :3.25 Mean :37.45 Mean :41.35 Mean :38.45   
## 3rd Qu.:4.00 3rd Qu.:39.00 3rd Qu.:46.00 3rd Qu.:40.00   
## Max. :8.00 Max. :46.00 Max. :49.00 Max. :43.00   
##   
## rankindiv rank21 averank variable   
## Min. : 1.00 Min. : 1.00 Min. : 1.500 locomotion1: 20   
## 1st Qu.: 5.75 1st Qu.: 5.75 1st Qu.: 5.875 locomotion2: 20   
## Median :10.50 Median :11.00 Median :10.500 assessment1: 20   
## Mean :10.50 Mean :11.00 Mean :10.750 assessment2: 20   
## 3rd Qu.:15.25 3rd Qu.:16.25 3rd Qu.:16.500 promotion1 : 20   
## Max. :20.00 Max. :21.00 Max. :20.000 promotion2 : 20   
## (Other) :160   
## value   
## Min. : 1.000   
## 1st Qu.: 2.600   
## Median : 3.830   
## Mean : 6.367   
## 3rd Qu.: 4.830   
## Max. :33.100   
## NA's :1

Okay, we're getting closer. We can see that we've now got 280 observations across 13 different variables. How did this happen and what are we left with? Let's look at the melt function, line-by-line.

breach <- melt(breach,

Nothing mind-blowing here. We've just called the melt function, and then fed it our dataframe as the first argument. The next line is the id.vars argument:

id.vars = c('subj', 'flourishing', 'swls', 'age', 'exp', 'apride', 'lpride', 'eom', 'rankindiv', 'rank21', 'averank'),

If we look closely at the updated breach dataframe, it's clear that these are the variables which get repeated. That is, while there are only 20 subjects, each subject now has their data spread across 14 rows. All the other variables listed in this argument are similarly repeated. Finally, we have the measure.vars argument:

measure.vars = c('locomotion1', 'locomotion2', 'assessment1', 'assessment2', 'promotion1', 'promotion2', 'prevention1', 'prevention2', 'stress1', 'stress2', 'lie2', 'lie1', 'LxA1', 'LxA2'))

What happened to all of these in our new dataframe? it seems they've all been dumped into two new variables. One has the name variables and the other has the name value. Let's examine the data for subject 2.

breach[breach$subj == 2,] #read: from the dataframe breach, give me the rows where subject = 2. After the comma is where one would select columns. Since I wanted all the columns, I just left it blank.

## subj flourishing swls age exp apride lpride eom rankindiv rank21  
## 2 2 5.25 5.8 26 6 38 38 35 4 3.5  
## 22 2 5.25 5.8 26 6 38 38 35 4 3.5  
## 42 2 5.25 5.8 26 6 38 38 35 4 3.5  
## 62 2 5.25 5.8 26 6 38 38 35 4 3.5  
## 82 2 5.25 5.8 26 6 38 38 35 4 3.5  
## 102 2 5.25 5.8 26 6 38 38 35 4 3.5  
## 122 2 5.25 5.8 26 6 38 38 35 4 3.5  
## 142 2 5.25 5.8 26 6 38 38 35 4 3.5  
## 162 2 5.25 5.8 26 6 38 38 35 4 3.5  
## 182 2 5.25 5.8 26 6 38 38 35 4 3.5  
## 202 2 5.25 5.8 26 6 38 38 35 4 3.5  
## 222 2 5.25 5.8 26 6 38 38 35 4 3.5  
## 242 2 5.25 5.8 26 6 38 38 35 4 3.5  
## 262 2 5.25 5.8 26 6 38 38 35 4 3.5  
## averank variable value  
## 2 3.75 locomotion1 4.50  
## 22 3.75 locomotion2 4.58  
## 42 3.75 assessment1 4.25  
## 62 3.75 assessment2 3.92  
## 82 3.75 promotion1 4.00  
## 102 3.75 promotion2 4.17  
## 122 3.75 prevention1 2.20  
## 142 3.75 prevention2 2.20  
## 162 3.75 stress1 2.00  
## 182 3.75 stress2 2.50  
## 202 3.75 lie2 2.67  
## 222 3.75 lie1 2.33  
## 242 3.75 LxA1 20.20  
## 262 3.75 LxA2 21.00

This (I think) really clearly illustrates what's going on here. Two things happen with all the variables listed in the measure.vars argument. First, the name of each variable gets put into a new vector called, unfortunately variable. Second, the corresponding value for that variable is listed in a second new vector labeled 'value'. Thus, since we had 14 variables listed in measure.vars, we now have 14 lines for each subject.

But, like I pointed out before, this isn't quite what we wanted. Remember our goal: We want a histogram where we can plot two variables on the x axis. And remember how I said we would do it:

**2 vectors** - *Vector 1: values to be plotted* - *Vector 2: indicates which of two groups each value belongs to.*

I know we just turned our 'wide' data into 'long', but now we need to go back. Our data is long, but it's *too* long... (yeah, yeah... har har)

#breach <- dcast(breach, )