

class05.R

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```
# Class 5: Data visualization Nicole Jacobson
```

```
#will learn how to use an array of plots using the ggplot2 package using prebuilt cars dataset
```

```
#looks at the top of the data set
```

```
head(cars)
```

```
##   speed dist
## 1     4    2
## 2     4   10
## 3     7    4
## 4     7   22
## 5     8   16
## 6     9   10
```

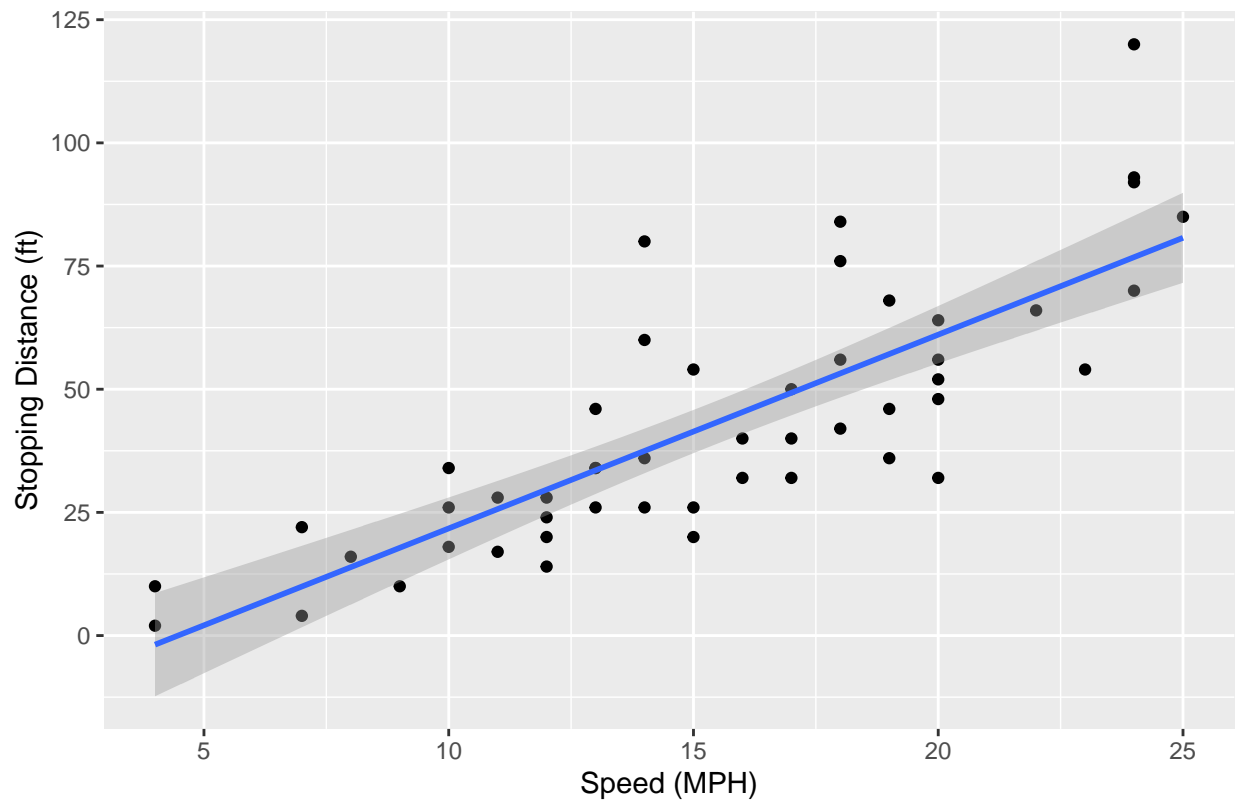
```
# all ggplots have at least 3 layers (data, ascetics -> x/y axis, labels ine color/thickness etc.,
# geometry -> type of graph ie scatter, bar, etc.)
```

```
#method=lm linear model will linearize the smooth line of best fit line, otherwise it will be curved
library(ggplot2)
```

```
ggplot(cars) + aes(x=speed, y=dist) + geom_point() + geom_smooth(method="lm") +
  ggtitle("Stopping Distance of Old Cars") + labs( x="Speed (MPH)", y="Stopping Distance (ft)")
```

```
## 'geom_smooth()' using formula 'y ~ x'
```

Stopping Distance of Old Cars



```
url <- "https://bioboot.github.io/bimm143_S20/class-material/up_down_expression.txt"
genes <- read.delim(url)
head(genes)
```

```
##      Gene Condition1 Condition2      State
## 1    A4GNT -3.6808610 -3.4401355  unchanging
## 2     AAAS  4.5479580  4.3864126  unchanging
## 3    AASDH  3.7190695  3.4787276  unchanging
## 4     AATF  5.0784720  5.0151916  unchanging
## 5     AATK  0.4711421  0.5598642  unchanging
## 6 AB015752.4 -3.6808610 -3.5921390  unchanging
```

```
nrow(genes)
```

```
## [1] 5196
```

```
ncol(genes)
```

```
## [1] 4
```

calls a particular column from that dataset. Tells me how many ups, downs, and unchanging there are

```
table(genes$State)
```

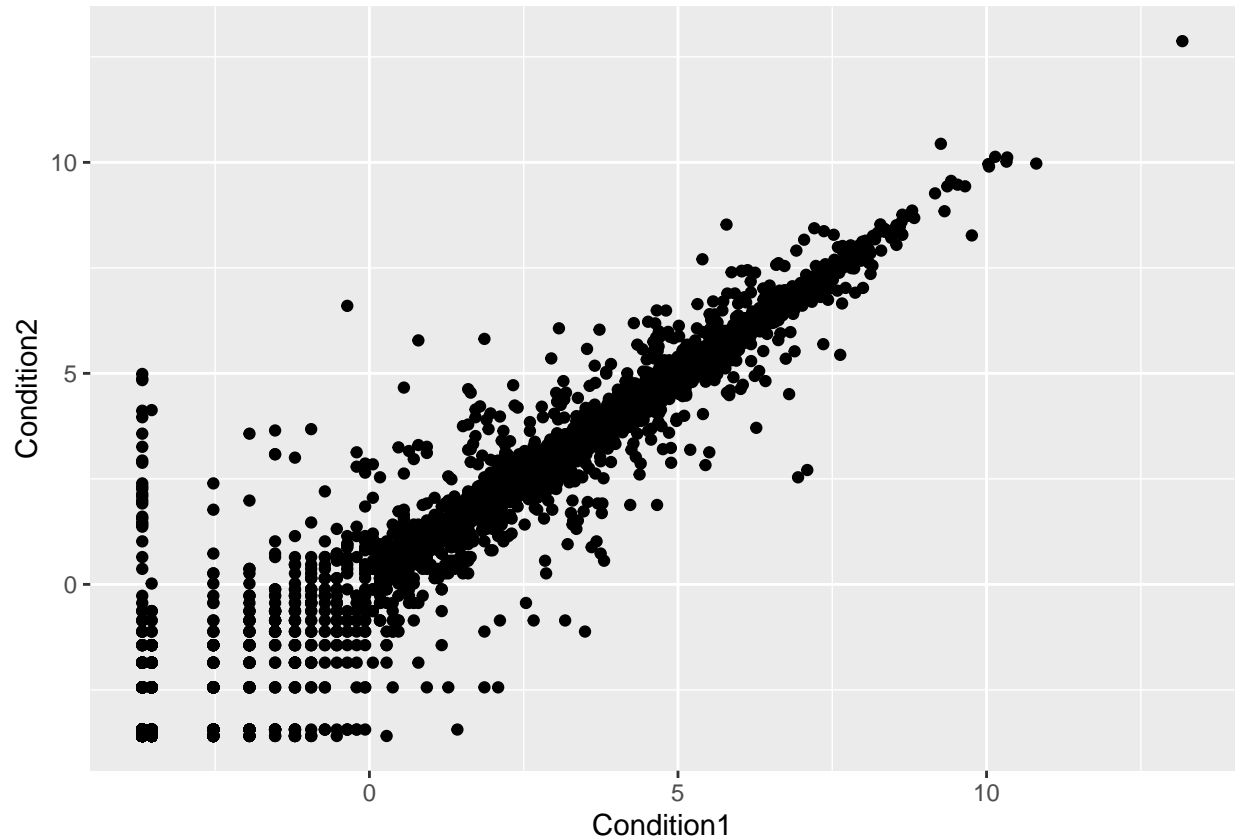
```
##
##      down  unchanged      up
##      72      4997      127
```

#what percent of total genes are up-regulated in this dataset? Here is the percent of genes that are up-regulated
by dividing table of the column State by the total number of entries which is nrow(genes)
x100 is to turn it into percent, 2 is how many significant digits
round will round the number to how many significant digits we want

```
round( table(genes$State)/nrow(genes) * 100, 2)
```

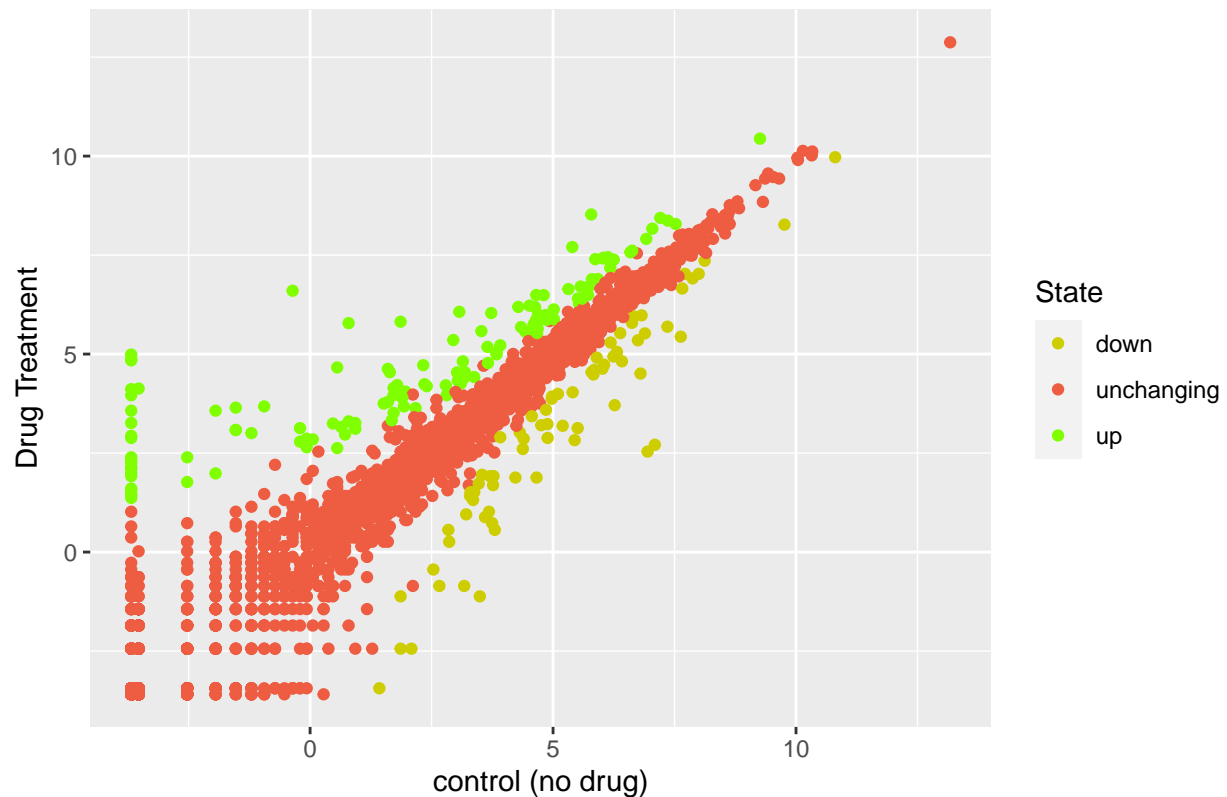
```
##
##      down  unchanged      up
##      1.39      96.17      2.44
```

```
ggplot(genes) + aes(x=Condition1, y=Condition2) + geom_point()
```



```
p <- ggplot(genes) + aes(x=Condition1, y=Condition2, col=State) + geom_point() + labs(x="control (no drug)", y="treatment (drug)")
p + scale_colour_manual(values=c("yellow3", "tomato2", "chartreuse"))
```

Gene Expression Changes Upon Drug Treatment

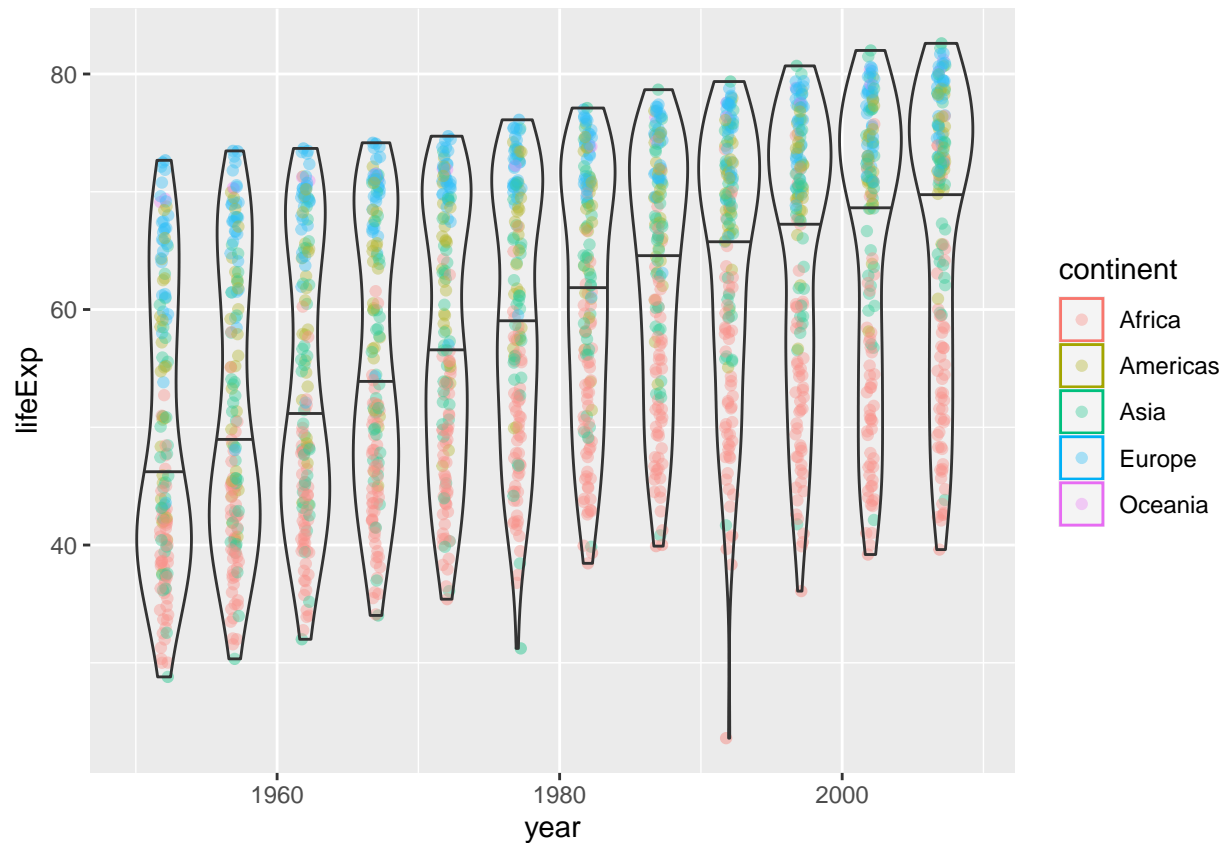


#now let's explore the gapminder dataset

```
#install.packages("gapminder")
library(gapminder)
head(gapminder)
```

```
## # A tibble: 6 x 6
##   country    continent  year lifeExp      pop gdpPercap
##   <fct>      <fct>    <int>  <dbl>   <int>   <dbl>
## 1 Afghanistan Asia      1952   28.8  8425333    779.
## 2 Afghanistan Asia      1957   30.3  9240934    821.
## 3 Afghanistan Asia      1962   32.0 10267083    853.
## 4 Afghanistan Asia      1967   34.0 11537966    836.
## 5 Afghanistan Asia      1972   36.1 13079460    740.
## 6 Afghanistan Asia      1977   38.4 14880372    786.
```

```
ggplot(gapminder) + aes(x=year, y=lifeExp, col=continent) + geom_jitter(width=0.3, alpha=0.4) +
  geom_violin(aes(group=year), alpha=0.2, draw_quantiles=0.5)
```



```
#install.packages("dplyr")
library(dplyr)
```

```
##
## Attaching package: 'dplyr'

## The following objects are masked from 'package:stats':
##
##   filter, lag

## The following objects are masked from 'package:base':
##
##   intersect, setdiff, setequal, union
```

```
gapminder_2007 <- gapminder %>% filter(year==2007)

gapminder_2007
```

```
## # A tibble: 142 x 6
##   country    continent  year lifeExp      pop gdpPercap
##   <fct>      <fct>    <int>  <dbl>    <int>    <dbl>
## 1 Afghanistan Asia      2007   43.8  31889923    975.
## 2 Albania    Europe    2007   76.4   3600523   5937.
## 3 Algeria    Africa    2007   72.3  33333216   6223.
```

```
## 4 Angola      Africa      2007      42.7 12420476    4797.
## 5 Argentina   Americas    2007      75.3 40301927   12779.
## 6 Australia   Oceania     2007      81.2 20434176   34435.
## 7 Austria     Europe      2007      79.8  8199783   36126.
## 8 Bahrain     Asia       2007      75.6  708573    29796.
## 9 Bangladesh  Asia       2007      64.1 150448339   1391.
## 10 Belgium    Europe      2007      79.4 10392226   33693.
## # ... with 132 more rows
```

```
ggplot(gapminder_2007) + aes(x=gdpPercap, y=lifeExp) + geom_point(alpha=0.5)
```

