## Class05

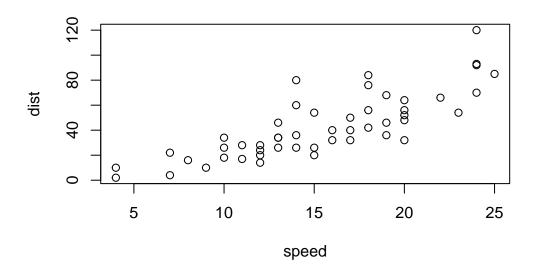
Liz

## Table of contents

#Our first plot

R has base graphics

plot(cars)



How would I plot this with 'ggplot2'? NO!

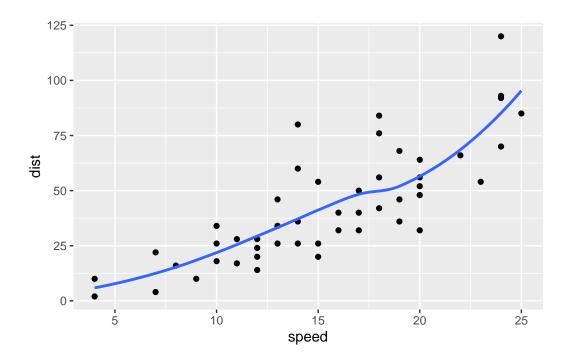
We need to install all load the ggplot package first. To install any package in R we use the 'install.packages()' function

```
library(ggplot2)
ggplot(cars)
```

Every ggplot needs at least 3 layers: **-Data** (i.e. the data.frame we have), **-Aes** (the aesthetic mapping of our data to what we want to plot) **Geoms** (How we want to plot this stuff!)

```
ggplot(data = cars) + aes(x = speed, y = dist) +
geom_point() + geom_smooth(se = FALSE)
```

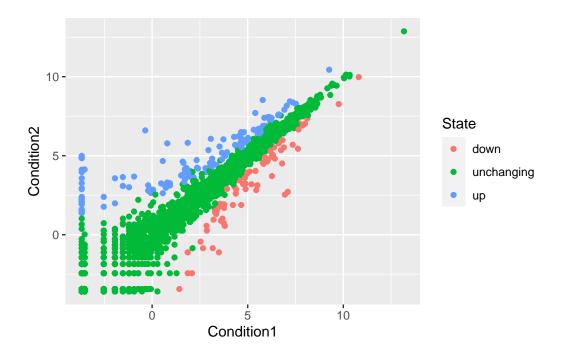
 $geom_smooth()$  using method = 'loess' and formula 'y ~ x'



url <- "https://bioboot.github.io/bimm143\_S20/class-material/up\_down\_expression.txt"
genes <- read.delim(url)
head(genes)</pre>

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

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

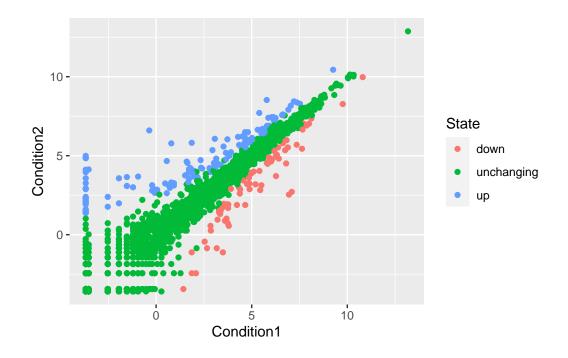


nrow(genes)

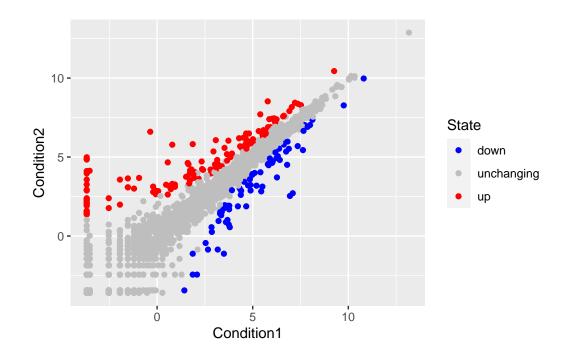
## [1] 5196

```
p <- ggplot(genes) +
   aes(x=Condition1, y=Condition2, col=State) +
   geom_point()

p</pre>
```

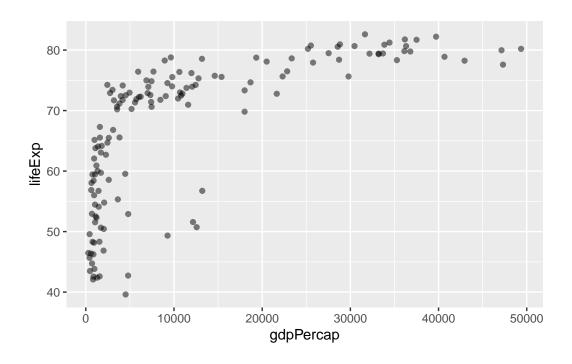


p + scale\_colour\_manual( values=c("blue", "gray", "red") )

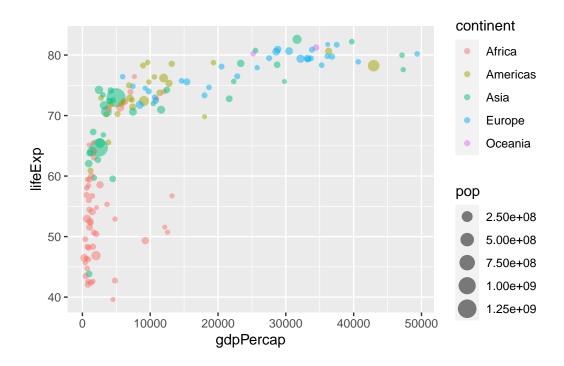


```
library(gapminder)
  library(gapminder)
  url <- "https://raw.githubusercontent.com/jennybc/gapminder/master/inst/extdata/gapminder.
  gapminder <- read.delim(url)</pre>
  library(dplyr)
Attaching package: 'dplyr'
The following objects are masked from 'package:stats':
    filter, lag
The following objects are masked from 'package:base':
    intersect, setdiff, setequal, union
  gapminder_2007 <- gapminder %>% filter(year==2007)
  library("gapminder")
  library(dplyr)
  gapminder_2007 <- gapminder %>% filter(year==2007)
  library(ggplot2)
  ggplot(gapminder_2007)
```

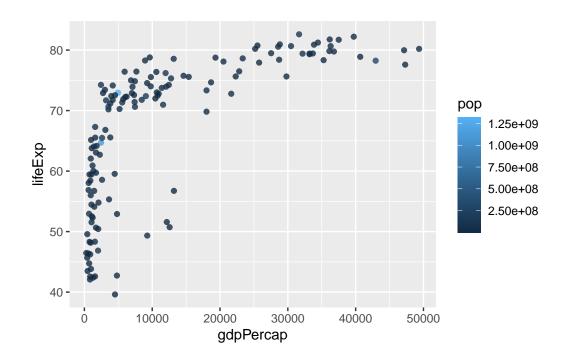
```
ggplot(gapminder_2007) +
aes(x=gdpPercap, y=lifeExp) +
geom_point(alpha=0.5)
```



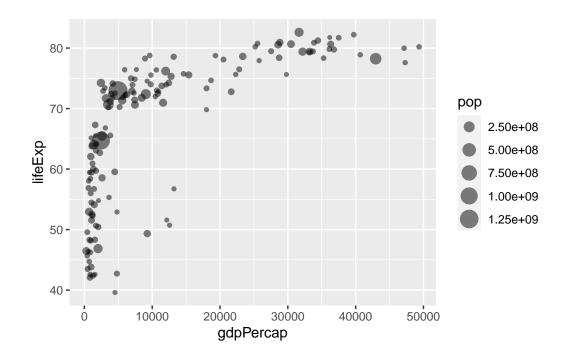
```
ggplot(gapminder_2007) +
  aes(x=gdpPercap, y=lifeExp, color=continent, size=pop) +
  geom_point(alpha=0.5)
```

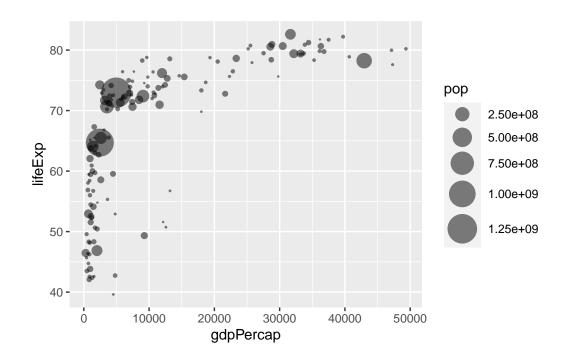


```
ggplot(gapminder_2007) +
  aes(x = gdpPercap, y = lifeExp, color = pop) +
  geom_point(alpha=0.8)
```



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

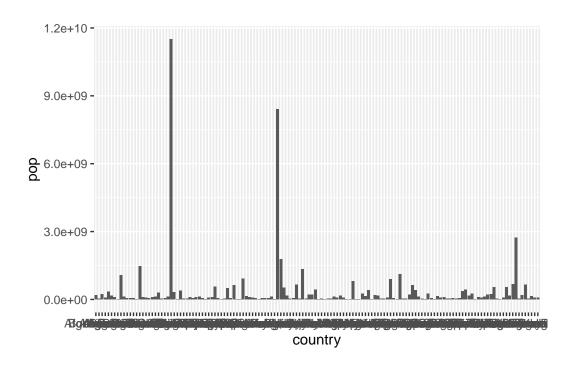




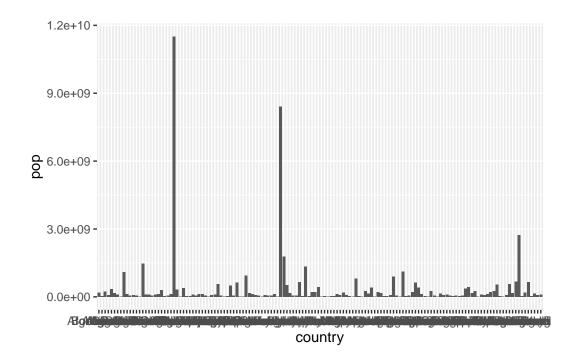
```
library("gapminder")

gapminder_top5 <- gapminder

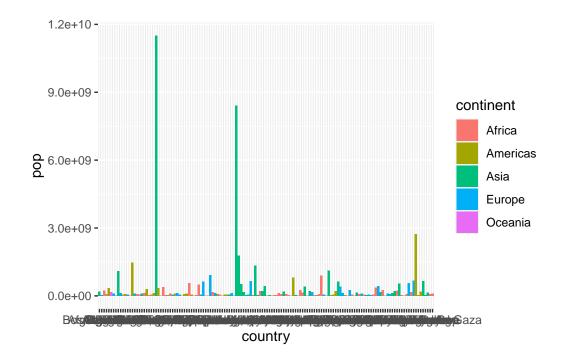
ggplot(gapminder_top5) +
  geom_col(aes(x = country, y = pop))</pre>
```



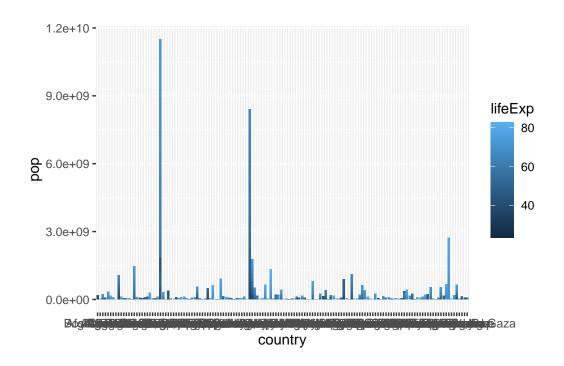
```
ggplot(gapminder_top5) +
geom_col(aes(x = country, y = pop))
```



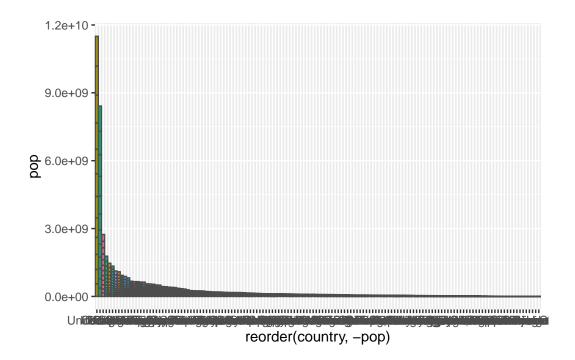
```
ggplot(gapminder_top5) +
geom_col(aes(x = country, y = pop, fill = continent))
```



```
ggplot(gapminder_top5) +
geom_col(aes(x = country, y = pop, fill = lifeExp))
```



```
ggplot(gapminder_top5) +
  aes(x=reorder(country, -pop), y=pop, fill=country) +
  geom_col(col="gray30") +
  guides(fill="none")
```



```
USArrests$State <- rownames(USArrests)
ggplot(USArrests) +
  aes(x=reorder(State,Murder), y=Murder) +
  geom_col() +
  coord_flip()</pre>
```

