Class 5: Data Visualization with ggplot

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##Using GGPLOT

To use ggplot2 we first need to install it on our computers. To do this we will use the function install.packages()

Before I use any package functions I have to load them up with a library() call, like so:

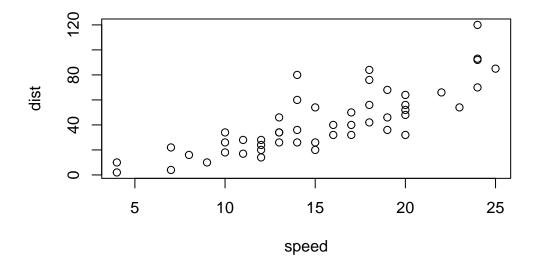
```
library(ggplot2)
head(cars)
```


cars

```
speed dist
1
       4
             2
2
       4
            10
3
       7
             4
       7
            22
5
       8
            16
6
       9
            10
7
      10
            18
8
      10
            26
9
      10
            34
            17
10
      11
```

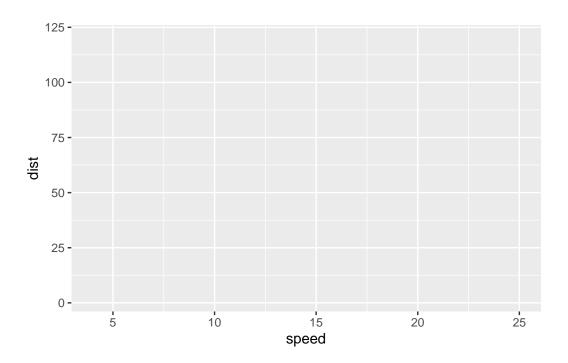
```
28
11
      11
12
      12
            14
13
      12
            20
14
      12
            24
15
       12
            28
16
      13
            26
17
      13
            34
18
      13
            34
19
      13
            46
20
      14
            26
21
      14
            36
22
      14
            60
23
            80
      14
24
      15
            20
25
      15
            26
26
      15
            54
27
      16
            32
28
      16
            40
29
      17
            32
      17
            40
30
            50
31
      17
32
      18
            42
33
      18
            56
34
            76
      18
35
      18
            84
      19
36
            36
37
      19
            46
38
       19
            68
39
      20
            32
40
      20
            48
      20
            52
41
42
      20
            56
43
      20
            64
44
      22
            66
45
      23
            54
      24
46
            70
      24
            92
47
48
      24
            93
49
      24
           120
50
      25
            85
```

There is always the "base R" graphics system, i.e. plot()

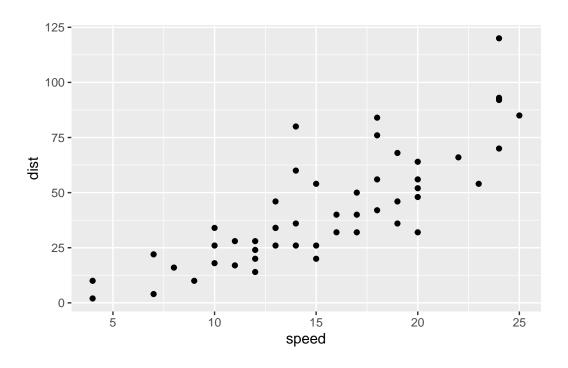


To use ggplot I need to spell out at least 3 sthing: -data(the stuff I want to plot as a data.frame) -aes (aes() values - how the data map to the plot) -geoms (how I want things drawn)

```
ggplot(cars) +
  aes(x=speed, y=dist)
```

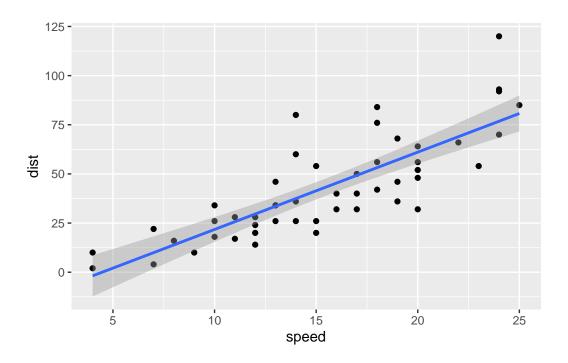


```
ggplot(cars) +
  aes(x=speed, y=dist) +
  geom_point()
```



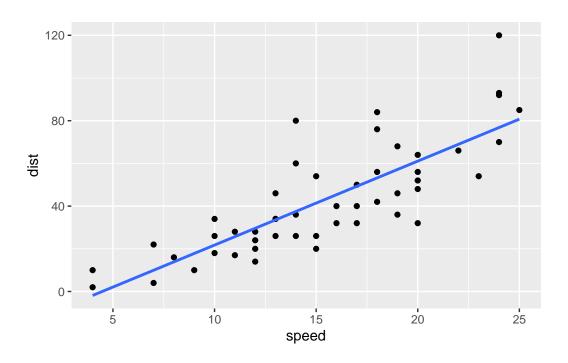
```
ggplot(cars) +
  aes(x=speed, y=dist) +
  geom_point() +
  geom_smooth(method="lm")
```

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



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

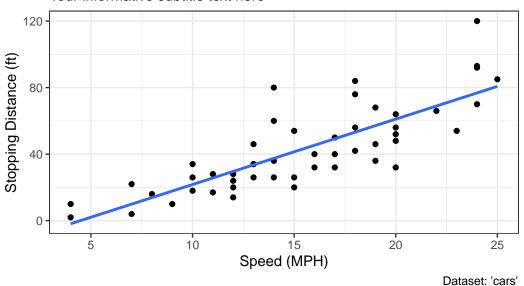
`geom_smooth()` using formula = 'y ~ x'



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

Speed and Stopping Distances of Cars

Your informative subtitle text here



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

```
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
```

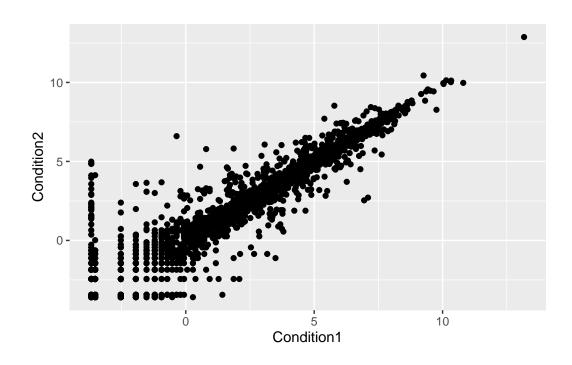
nrow(genes)

[1] 5196

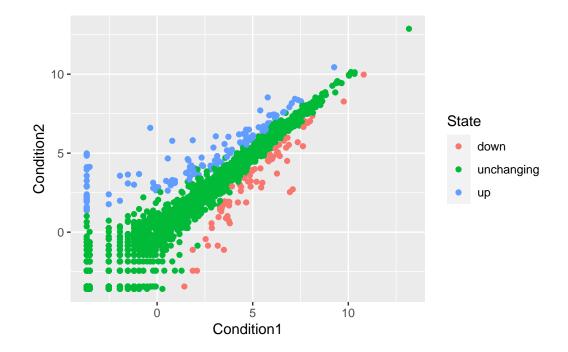
colnames(genes)

[1] "Gene" "Condition1" "Condition2" "State"

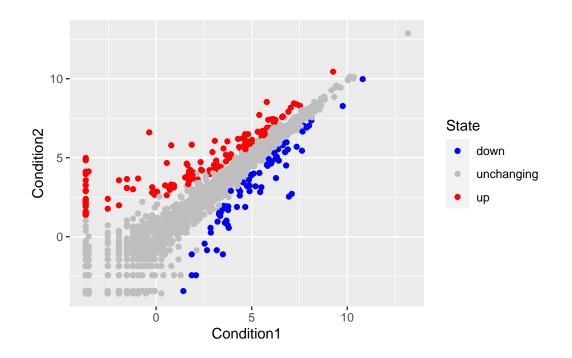
```
ncol(genes)
[1] 4
  table(genes$State)
      down unchanging
                              up
        72
                 4997
                             127
  round ( table(genes$State)/nrow(genes) * 100, 2)
      down unchanging
                              up
                            2.44
      1.39
                96.17
  ggplot(genes) +
    aes(Condition1, Condition2) +
    geom_point()
```



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

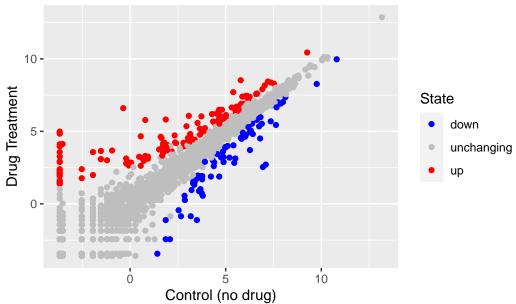


```
p + scale_colour_manual( values=c("blue", "gray", "red") )
```



```
p + scale_colour_manual(values=c("blue", "gray", "red")) +
    labs(title="Gene Expression Changes Upon Drug Development", x="Control (no drug)", y="Drug")
```

Gene Expression Changes Upon Drug Development



```
url <- "https://raw.githubusercontent.com/jennybc/gapminder/master/inst/extdata/gapminder.
gapminder <- read.delim(url)

# install.packages("dplyr") ## un-comment to install if needed
library(dplyr)

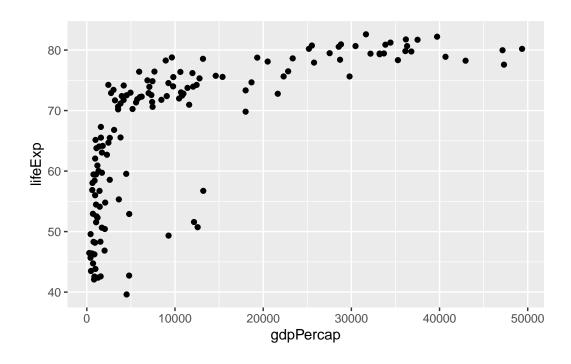
Attaching package: 'dplyr'

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

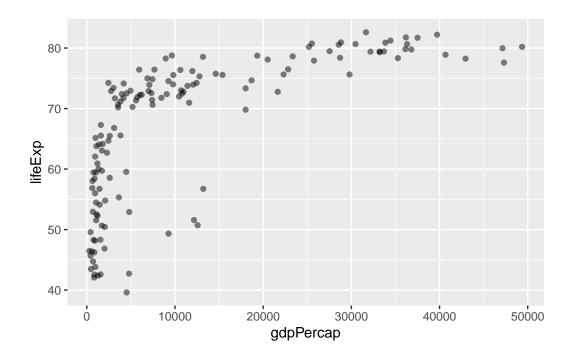
The following objects are masked from 'package:base':
    intersect, setdiff, setequal, union

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

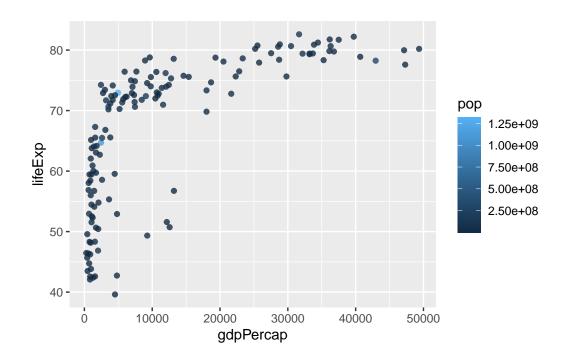
ggplot(gapminder_2007) +
    aes(x=gdpPercap, y=lifeExp) +
    geom_point()
```



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

