lab05: Data vis with ggplot

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Background

There are many graphic systems available in R. These include "base" R and tons of add-on packages like **ggplot2**.

Let's compare "base" and **ggplot2** briefly: We can use some example data that is built-in with R called **cars**:

head(cars)

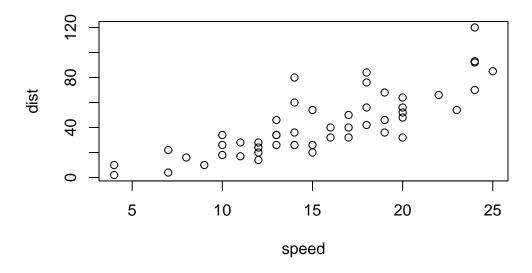
	speed	dist
1	4	2
2	4	10
3	7	4
4	7	22

5 8 16 6 9 10

Plotting in Base

In base R I can just call plot()

plot(cars)



Importing ggplot

How we can do this with ggplot2

First we need to install the package. We do this install.packages("ggplot2"). I only need to do this once and then it will be available on my computer from then on.

Key point: I only install packages in the R console not within quarto docs or R scripts.

Before using any add-on package, loading it up with a call to library()

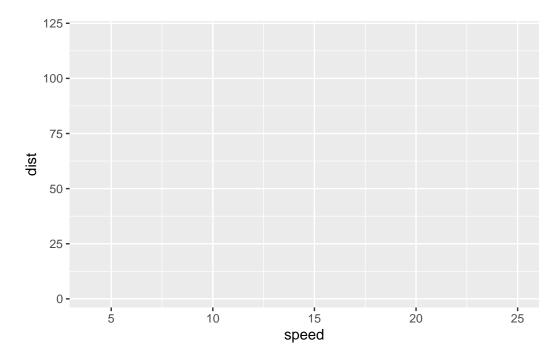
```
library(ggplot2)
ggplot(cars)
```

Every ggplot has at least 3 things:

- the data (in our case cars)
- the **aes**thetics (how the data map to the plot)
- the **geom**s that determine how the plot is drawn (lines, points, column, boxplots, densities, etc)

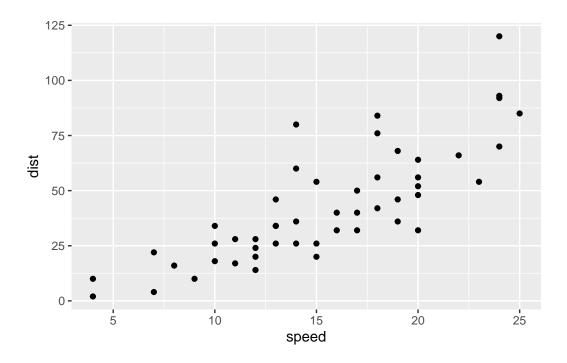
Two of three elements: introduced the aesthetics such as specifying the x and y axes.

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



All three elements: For "simple" plots, ggplot is much more verbose than base R but the defaults are nicer and for complicated plots it becomes much more efficient and structured.

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



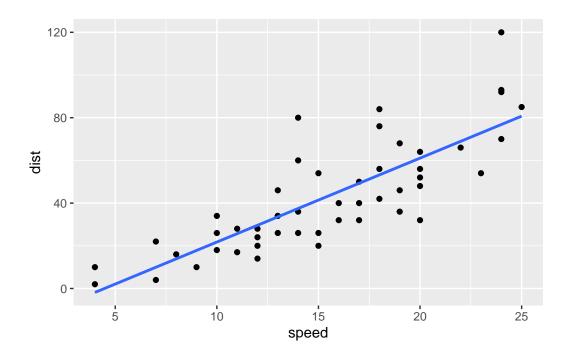
Q. Add a line to show the relationship of speed and stopping distance (i.e add another "layer")

```
p <- ggplot(cars) +
  aes(x = speed, y = dist) +
  geom_point() +
  geom_smooth(se = FALSE, method = "lm")</pre>
```

I can always save any ggplot object (i.e plot) and then use it later for adding more layers.

p

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

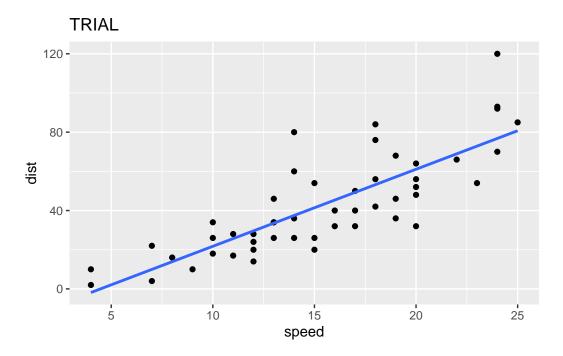


Q. Add a title and subtitle to the plot

Option 1

p + ggtitle("TRIAL")

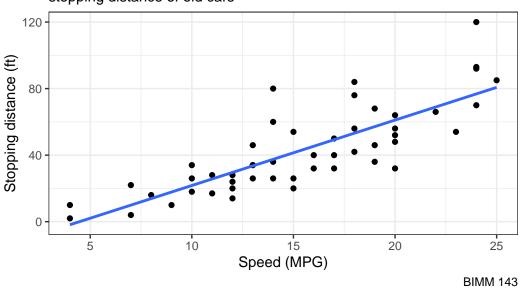
[`]geom_smooth()` using formula = 'y ~ x'



Option 2

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

My first ggplot stopping distance of old cars



Gene expression

Read input data into R

```
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
```

Q. How many genes are in this dataset?

```
nrow(genes)
```

[1] 5196

Q. How many columns are in the dataset?

ncol(genes)

[1] 4

Q. What are the column names?

colnames(genes)

- [1] "Gene" "Condition1" "Condition2" "State"
 - Q. How many up and down regulated genes are there?

table(genes\$State)

```
down unchanging up
72 4997 127
```

Q. What fraction of total genes are upregulated?

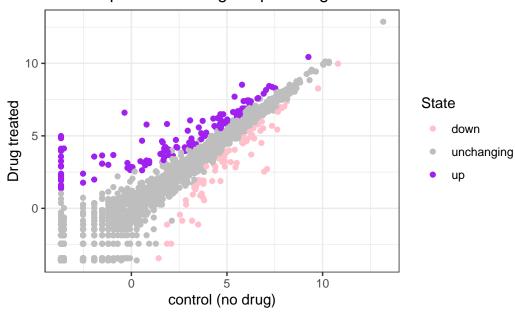
```
round( table(genes$State)/nrow(genes) * 100, 2 )
```

```
down unchanging up
1.39 96.17 2.44
```

Q. Make a first plot of this data

Color scatterplot

Gene expression changes upon drug treatment



Using different geoms

Let's plot some aspects of the in-built mtcars dataset.

head(mtcars)

```
      mpg cyl disp
      hp drat
      wt qsec vs am gear carb

      Mazda RX4
      21.0
      6
      160
      110
      3.90
      2.620
      16.46
      0
      1
      4
      4

      Mazda RX4 Wag
      21.0
      6
      160
      110
      3.90
      2.875
      17.02
      0
      1
      4
      4

      Datsun 710
      22.8
      4
      108
      93
      3.85
      2.320
      18.61
      1
      1
      4
      1

      Hornet 4 Drive
      21.4
      6
      258
      110
      3.08
      3.215
      19.44
      1
      0
      3
      1
```

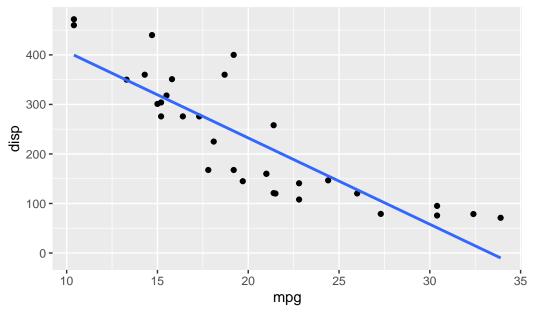
```
Hornet Sportabout 18.7 8 360 175 3.15 3.440 17.02 0 0 3 2 Valiant 18.1 6 225 105 2.76 3.460 20.22 1 0 3 1
```

Q. Scatterplot of mpg vs disp

```
p1 <- ggplot(mtcars) +
  aes(x = mpg, y = disp) +
  geom_point() +
  geom_smooth(se = FALSE, method = "lm") +
  labs(title = "Scatterplot of mpg vs disp", x = "mpg", y = "disp")
p1</pre>
```

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

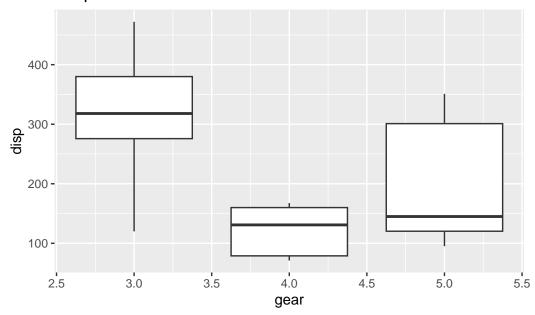
Scatterplot of mpg vs disp



Q. Boxplot of gear vs disp

```
p2 <- ggplot(mtcars) +
  aes(x = gear, y = disp, group = gear) +
  geom_boxplot() +
  labs(title = "Boxplot of carb", x = "gear", y = "disp")
p2</pre>
```

Boxplot of carb



Q. Barplot of carb

```
p3 <- ggplot(mtcars) +
  aes(x = carb) +
  geom_bar() +
  labs(title = "Barplot of carb")
p3</pre>
```

Times 10.0 - 10.

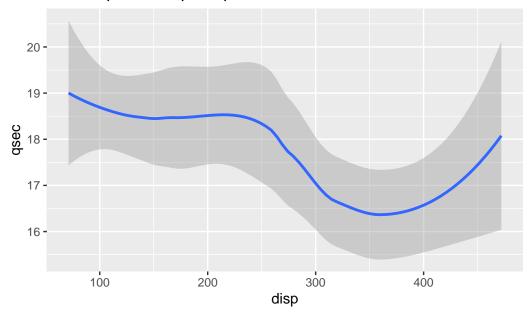
Q. Smooth of disp vs qsec

```
p4 <- ggplot(mtcars) +
  aes ( x = disp, y = qsec) +
  geom_smooth() +
  labs(title = "Smoothplot of disp vs qsec", x = "disp", y = "qsec")
p4</pre>
```

carb

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

Smoothplot of disp vs qsec



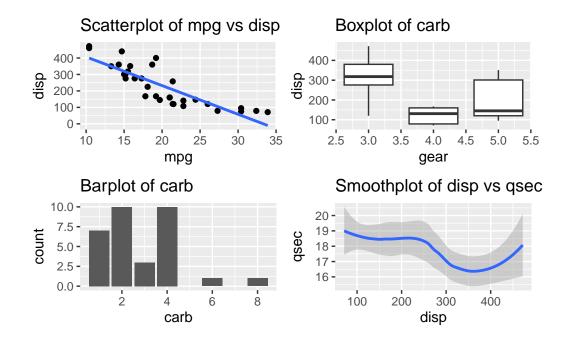
Panel of plots

I want to combine all these plots into one figure with multiple pannels.

We can use the **pathcwork** package to do this.

```
library(patchwork)
(p1 + p2 + p3 + p4)
```

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



```
ggsave(filename ="myplot.png", width = 10, height = 10)
```

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

Life Expectancy by country plot

Importing/Reading the file

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

```
country continent year lifeExp
                                         pop gdpPercap
1 Afghanistan
                  Asia 1952
                             28.801
                                              779.4453
                                     8425333
                                     9240934
2 Afghanistan
                  Asia 1957
                             30.332
                                              820.8530
3 Afghanistan
                  Asia 1962 31.997 10267083
                                              853.1007
4 Afghanistan
                  Asia 1967 34.020 11537966 836.1971
```

```
5 Afghanistan Asia 1972 36.088 13079460 739.9811
6 Afghanistan Asia 1977 38.438 14880372 786.1134
```

Q. How many countries are in this dataset?

```
length(table(gapminder$country))
```

[1] 142

Gapminder plot

```
ggplot(gapminder) +
  aes(x = gdpPercap, y = lifeExp, col = continent, size = pop) +
  geom_point(alpha = 0.3) +
  facet_wrap(~continent) +
  theme_bw()
```

