Class 5: Data Visualization with ggplot

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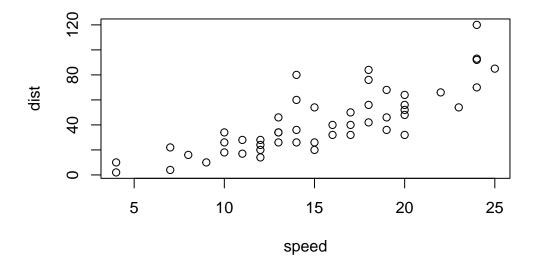
Background

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

Let's compare "base" and $\mathbf{ggplot2}$ briefly. We can use some example data that is built-in with R called \mathbf{cars} :

head(cars)

I can just call plot() in base R.



We can now make it pretty and we do this with ggplot! ggplot2

First we install ggplot. We do this with the code install.packages("ggplot2"). This only need to do this once and then it will be available on my computer any time I need to access ggplot.

Key Point: Packages can only be installed in the R console. Do not do it in quarto docs or R scripts.

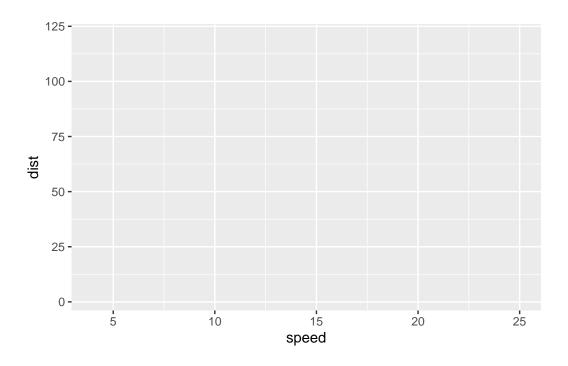
To use any add-on packages you have installed, you need to load it up with library ()

library(ggplot2)
ggplot(cars)

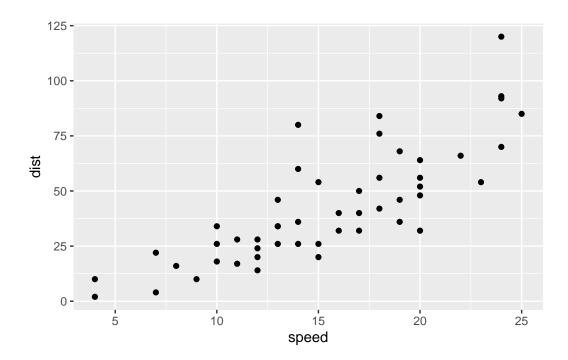
Every ggplot needs three different layers to make it show up:

- the data (and in our case it is cars)
- the **aes**thetics (how the data looks when mapped on the plot. Such as color and axis titles)
- the **geom**etry(what the lines/plots/data is drawn on the plot such as line width)

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



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

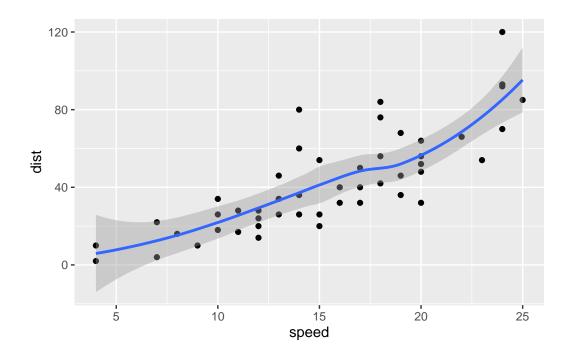


For more "simpler" plots, ggplot requires more words/code than base R. However, the defaults in ggplot are nicer and it is more efficient for complicated plots.

Try adding a line to show the relationship between speed and stopping distance. Tip: by adding another layer

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

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



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

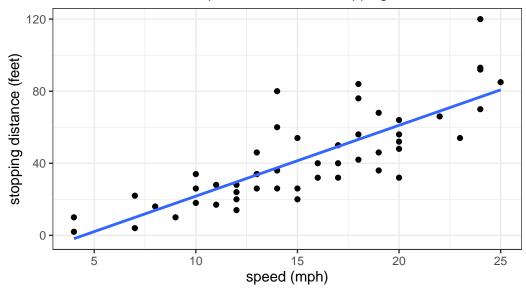
I can save an object from ggplot for future use using x <- x

Try adding axis titles

```
p + labs(
    title="speed vs. distance",
    subtitle="cars that have a faster speed have a faster stopping distance",
    x="speed (mph)",
    y="stopping distance (feet)"
)+
    theme_bw()
```

speed vs. distance

cars that have a faster speed have a faster stopping distance



Gene expression plot

Read input data

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

Gene Condition1 Condition2 State

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

```
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
```

How many genes are in this dataset? How many columns and what are their names?

```
nrow(genes)
```

[1] 5196

```
ncol(genes)
```

Γ1 4

```
colnames(genes)
```

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

How many "up" and "down" regulated genes are there?

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

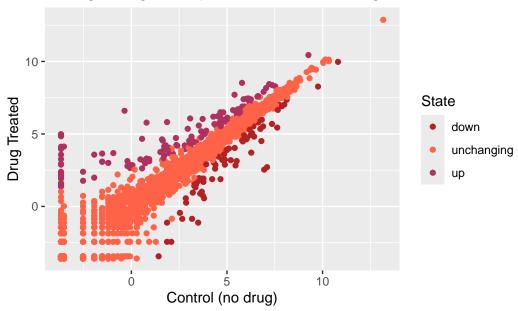
```
down unchanging up
72 4997 127
```

Custom Color Plot

Make a plot with the data

```
ggplot(genes)+
  aes(x=Condition1, y=Condition2, col=State)+
  scale_color_manual(
    values=c("firebrick","tomato","maroon")
)+
  geom_point()+
  labs(title="Changes of gene expression based on drug treatments", x="Control (no drug)", y=
```

Changes of gene expression based on drug treatments



Using Different Geoms

This will be plotted using the built-in mtcars dataset

head(mtcars)

```
wt qsec vs am gear carb
                  mpg cyl disp hp drat
Mazda RX4
                 21.0
                       6 160 110 3.90 2.620 16.46
Mazda RX4 Wag
                 21.0
                        6 160 110 3.90 2.875 17.02
                                                                4
Datsun 710
                 22.8 4 108 93 3.85 2.320 18.61
                                                                1
Hornet 4 Drive
                 21.4
                       6 258 110 3.08 3.215 19.44
                                                           3
                                                               1
                                                           3
                                                                2
Hornet Sportabout 18.7
                          360 175 3.15 3.440 17.02 0 0
Valiant
                 18.1
                          225 105 2.76 3.460 20.22 1 0
                                                                1
```

Scatterplot of mpg vs disp

Boxplot of gear vs disp

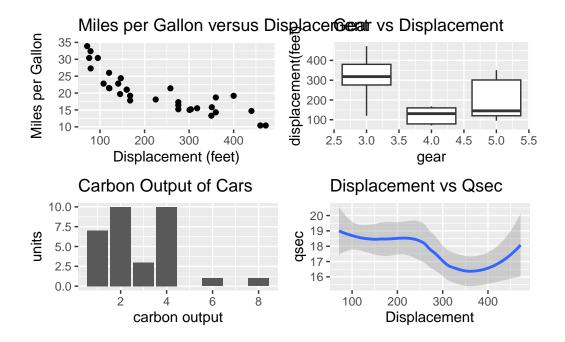
Barplot of carb

Smooth of disp vs qsec

I want to now combine all of these plots into one figure with a panel!

Use the code library(patchwork) to get this.

^{&#}x27;geom_smooth()' using method = 'loess' and formula = 'y ~ x'



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

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

More Aesthetic Features!

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

To just see the table

head(gapminder)

```
      country
      continent
      year
      lifeExp
      pop
      gdpPercap

      1
      Afghanistan
      Asia
      1952
      28.801
      8425333
      779.4453

      2
      Afghanistan
      Asia
      1957
      30.332
      9240934
      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

How many countries are in this dataset?

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

[1] 142

Plot GDP vs Life Expectancy and color by continent

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

