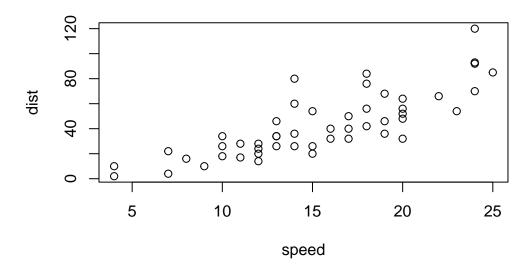
Class 5: Data Viz with ggplot

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R has lot's of ways to make figures and graphs in particular. One that comes with R out of the box is called "base" R - the plot() function.

plot(cars)

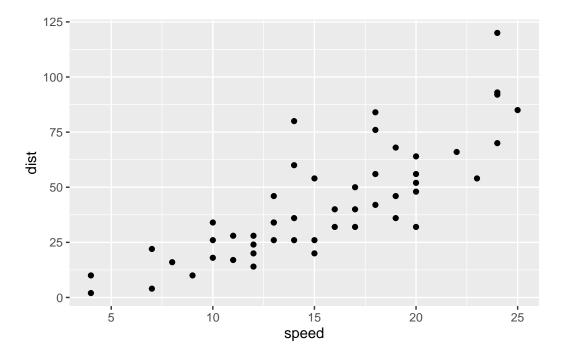


A very popular package in this area is called **ggplot2**.

Before I can use any add-on package like this I must install it with the install.packages("ggplot2") command/function.

Then to use the package I need to load it with a library(ggplot2) call.

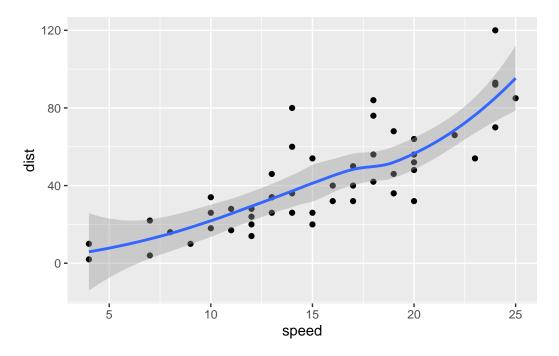
```
# install.packages("ggplot2")
library(ggplot2)
ggplot(cars) +
  aes(x=speed, y=dist) +
  geom_point()
```



For "simple" plots like this one base R code will be much shorter than ggplot code. Let's fit a model and show it on my plot:

```
library(ggplot2)
ggplot(cars) +
  aes(x=speed, y=dist) +
  geom_point() +
  geom_smooth()
```

 $[\]ensuremath{\text{`geom_smooth()`}}\ using method = 'loess' and formula = 'y ~ x'$



Every ggplot has at least 3 layers

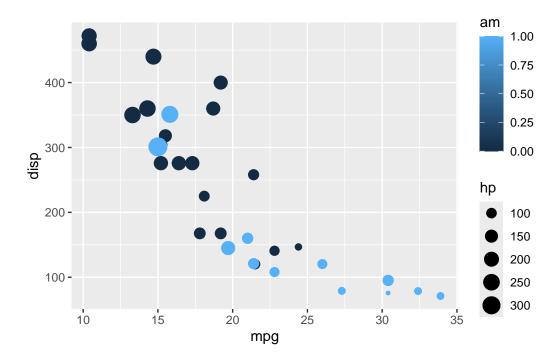
- data (data.frame with the numbers or stuff you want to plot)
- aesthetics (mapping of your data columns to your plot)
- **geom**etries (there are tones of these, basics are <code>geom_point()</code>, <code>geom_line()</code>, <code>geom_col())</code>

head(mtcars)

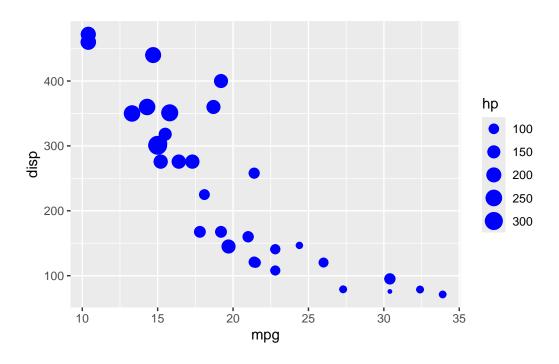
	mpg	cyl	disp	hp	${\tt drat}$	wt	qsec	٧s	\mathtt{am}	gear	carb
Mazda RX4	21.0	6	160	110	3.90	2.620	16.46	0	1	4	4
Mazda RX4 Wag	21.0	6	160	110	3.90	2.875	17.02	0	1	4	4
Datsun 710	22.8	4	108	93	3.85	2.320	18.61	1	1	4	1
Hornet 4 Drive	21.4	6	258	110	3.08	3.215	19.44	1	0	3	1
Hornet Sportabout	18.7	8	360	175	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

Make me a ggplot of the 'mtcars' data set using mpg vs disp and set the size of the points to the hp and set the color to am

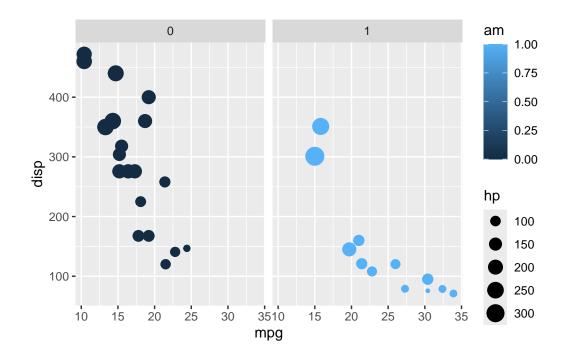
```
library(ggplot2)
ggplot(mtcars) +
  aes(x=mpg, y=disp, size=hp, col=am) +
  geom_point()
```



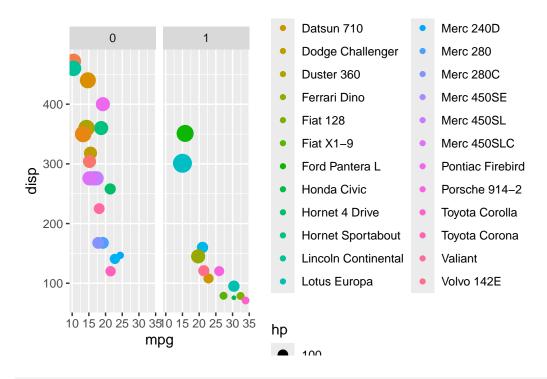
```
library(ggplot2)
ggplot(mtcars) +
  aes(x=mpg, y=disp, size=hp, col=am) +
  geom_point(col='blue')
```



```
library(ggplot2)
ggplot(mtcars) +
  aes(x=mpg, y=disp, size=hp, col=am) +
  geom_point() +
  facet_wrap(~am)
```



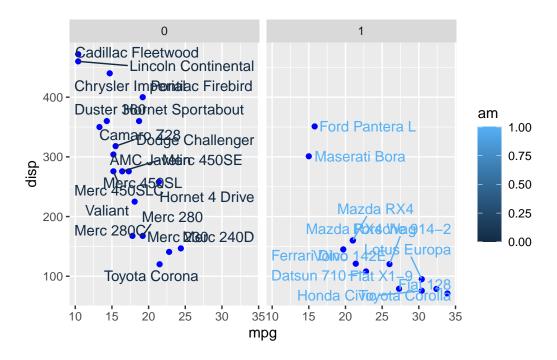
```
library(ggplot2)
ggplot(mtcars) +
  aes(x=mpg, y=disp, size=hp, col=rownames(mtcars)) +
  geom_point() +
  facet_wrap(~am)
```



rownames(mtcars)

```
[1] "Mazda RX4"
                            "Mazda RX4 Wag"
                                                   "Datsun 710"
 [4] "Hornet 4 Drive"
                            "Hornet Sportabout"
                                                   "Valiant"
 [7] "Duster 360"
                                                   "Merc 230"
                            "Merc 240D"
[10] "Merc 280"
                            "Merc 280C"
                                                   "Merc 450SE"
                            "Merc 450SLC"
[13] "Merc 450SL"
                                                   "Cadillac Fleetwood"
[16] "Lincoln Continental"
                            "Chrysler Imperial"
                                                   "Fiat 128"
[19] "Honda Civic"
                            "Toyota Corolla"
                                                   "Toyota Corona"
[22] "Dodge Challenger"
                            "AMC Javelin"
                                                   "Camaro Z28"
                            "Fiat X1-9"
[25] "Pontiac Firebird"
                                                   "Porsche 914-2"
[28] "Lotus Europa"
                            "Ford Pantera L"
                                                   "Ferrari Dino"
[31] "Maserati Bora"
                            "Volvo 142E"
```

```
library(ggrepel)
ggplot(mtcars) +
  aes(x=mpg, y=disp, label=rownames(mtcars), col=am) +
  geom_point(col='blue') +
  facet_wrap(~am) +
  geom_text_repel()
```



Gene expression plot

```
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
```

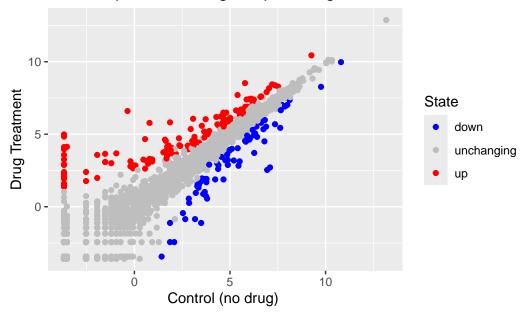
colnames(genes)

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

ncol(genes)

[1] 4

Gene Expresion Changes Upon Drug Treatment



nrow(genes)

[1] 5196

There are 5196 genes in this dataset

```
unique(genes$State)
```

```
[1] "unchanging" "up" "down"
```

The table() function is a super useful utility to tell me how many entries of each type there are.

```
round(table (genes$State) / nrow(genes), 4)
```

```
down unchanging up 0.0139 0.9617 0.0244
```

The functions nrow(), ncol(), and table() are ones I want you to know.

```
library(dplyr)
```

```
Attaching package: 'dplyr'

The following objects are masked from 'package:stats':

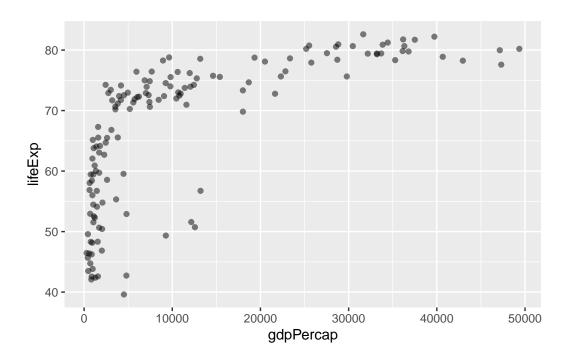
filter, lag

The following objects are masked from 'package:base':

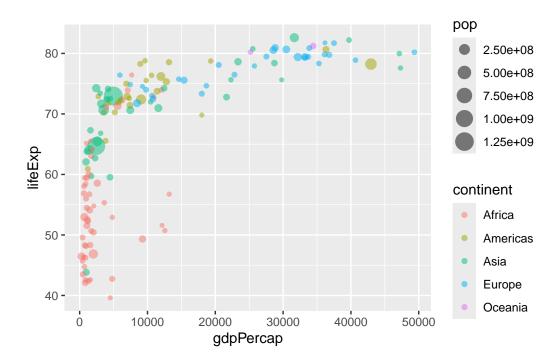
intersect, setdiff, setequal, union
```

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

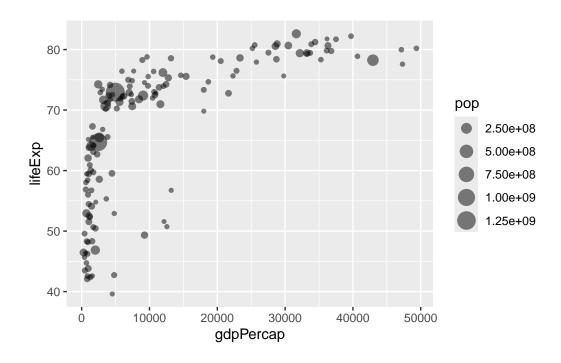
ggplot(gapminder_2007) +
  aes(x=gdpPercap, y=lifeExp) +
  geom_point(alpha=0.5)
```

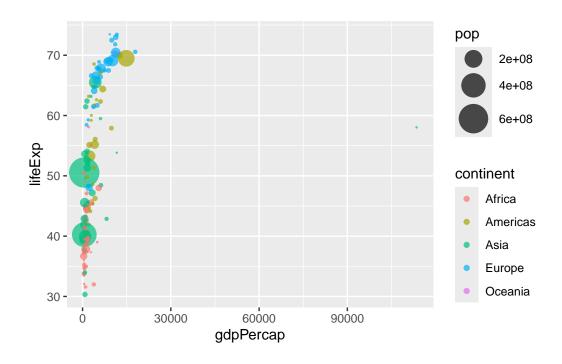


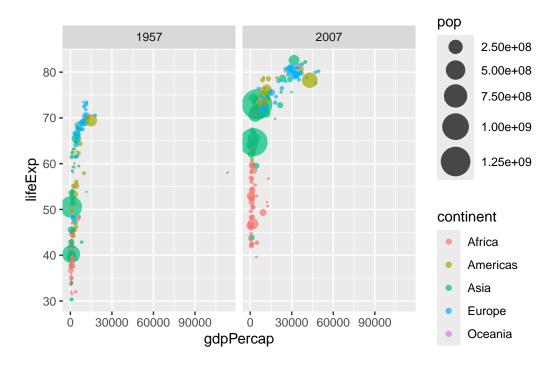
```
ggplot(gapminder_2007) +
  aes(x=gdpPercap, y=lifeExp, color=continent, size=pop) +
  geom_point(alpha=0.5)
```



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

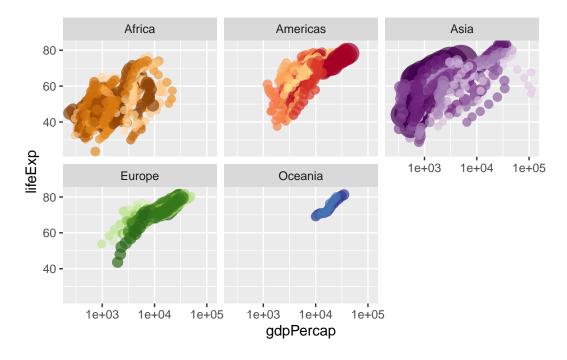






```
library(gapminder)
library(gganimate)

# Setup nice regular ggplot of the gapminder data
ggplot(gapminder, aes(gdpPercap, lifeExp, size = pop, colour = country)) +
    geom_point(alpha = 0.7, show.legend = FALSE) +
    scale_colour_manual(values = country_colors) +
    scale_size(range = c(2, 12)) +
    scale_x_log10() +
    # Facet by continent
    facet_wrap(~continent)
```



```
# Here comes the gganimate specific bits
# labs(title = 'Year: {frame_time}', x = 'GDP per capita', y = 'life expectancy') +
# transition_time(year) +
# shadow_wake(wake_length = 0.1, alpha = FALSE)
```

head(gapminder)

```
# A tibble: 6 x 6
 country
              continent year lifeExp
                                            pop gdpPercap
  <fct>
              <fct>
                                 <dbl>
                                                    <dbl>
                        <int>
                                          <int>
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.
```