

Class05: Data vis with ggplot2

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Today we will be playing with plotting and graphics in R!

There are many ways to make cool figures in R.

There are base R graphs like `plot()`, `hist()`, and `boxplot()`.

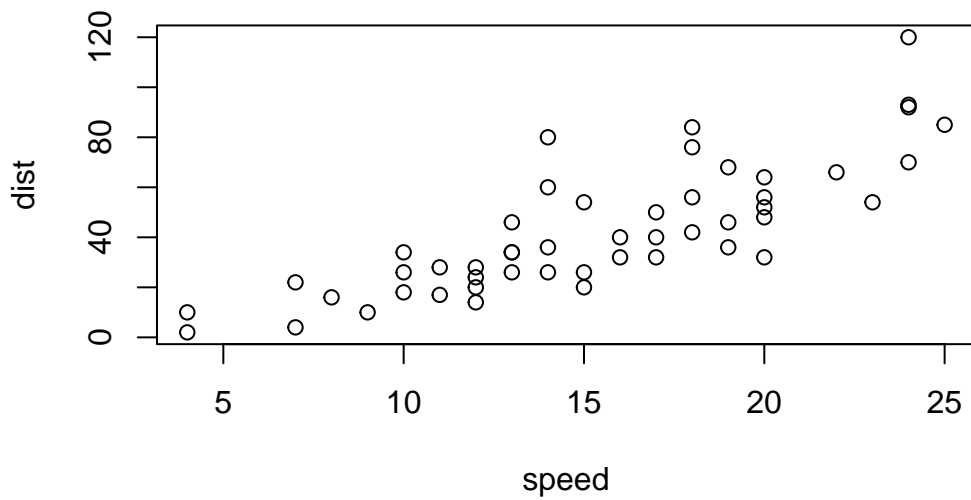
ggplot is a popular add-on package.

```
head(cars)
```

	speed	dist
1	4	2
2	4	10
3	7	4
4	7	22
5	8	16
6	9	10

Let's plot this with base R.

```
plot(cars)
```

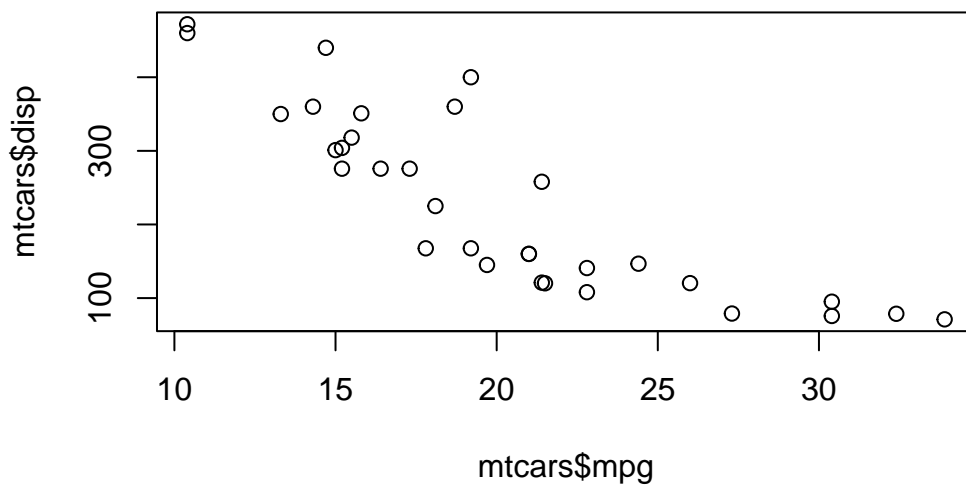


```
head(mtcars)
```

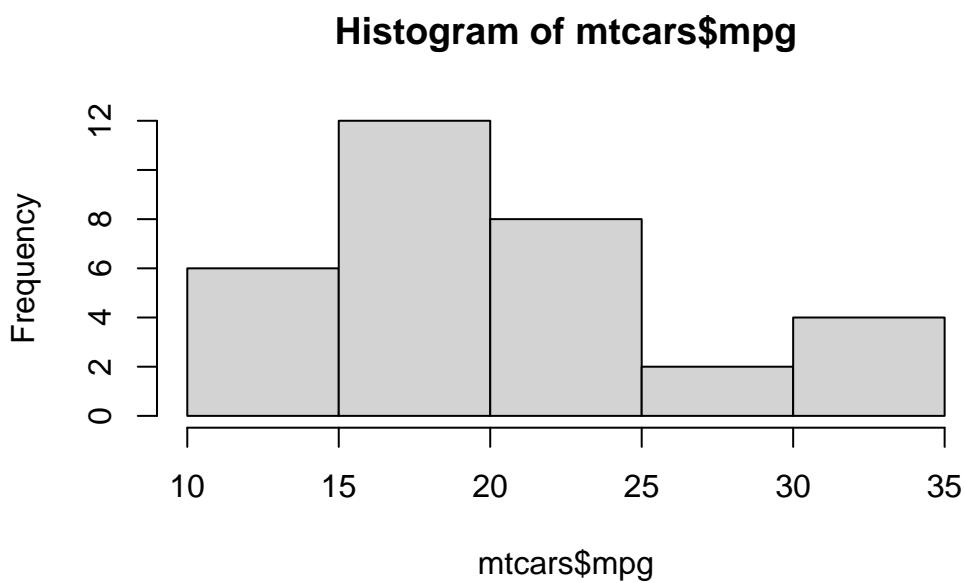
	mpg	cyl	disp	hp	drat	wt	qsec	vs	am	gear	carb
Mazda RX4	21.0	6	160	110	3.90	2.620	16.46	0	1	4	4
Mazda RX4 Wag	21.0	6	160	110	3.90	2.875	17.02	0	1	4	4
Datsun 710	22.8	4	108	93	3.85	2.320	18.61	1	1	4	1
Hornet 4 Drive	21.4	6	258	110	3.08	3.215	19.44	1	0	3	1
Hornet Sportabout	18.7	8	360	175	3.15	3.440	17.02	0	0	3	2
Valiant	18.1	6	225	105	2.76	3.460	20.22	1	0	3	1

Let's plot mpg vs displacement

```
plot(mtcars$mpg, mtcars$disp)
```



```
hist(mtcars$mpg)
```



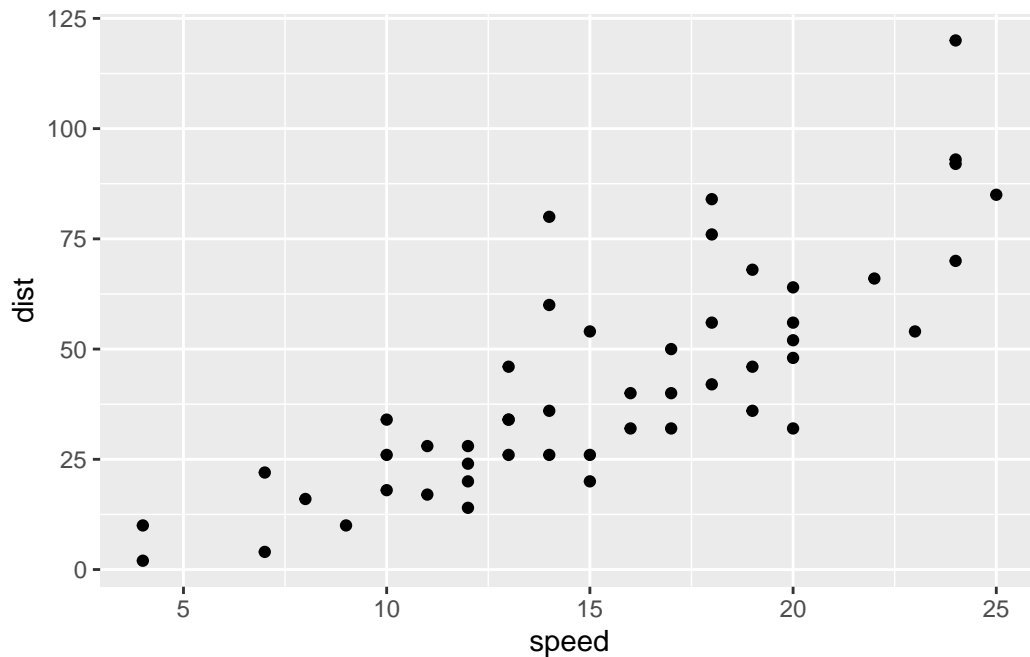
The main function on the ggplot2 package is **ggplot()**. To install the package, use the

function `install.package`. I already had `ggplot2` installed. You have to call it with `library(ggplot)`.

```
library(ggplot2)
```

Warning: package 'ggplot2' was built under R version 4.3.3

```
ggplot(cars) + aes(speed, dist) + geom_point()
```



ggplot needs: - **data**, given with `ggplot(x)`, defines the source of the data -**aesthetic**, given with `aes(x,y)`, defines the plotted data and axes -**geom**, given with `geom_(point)` for example, defines how it is visualized.

Base R is nearly always faster, but ggplot is more flexible and customized

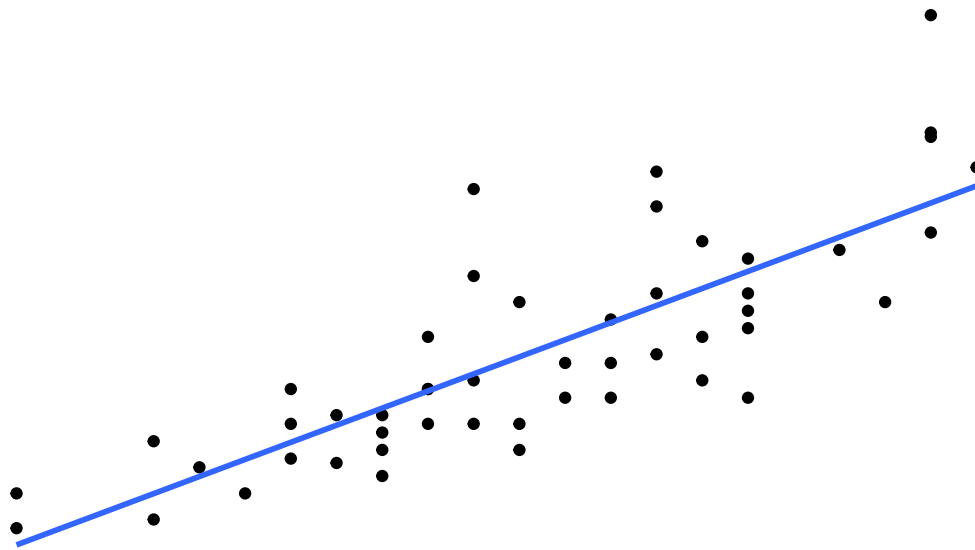
Let's add more layers

Let's add a line, a title, a subtitle, a caption, and custom axis labels.

```
ggplot(cars) + aes(speed, dist) + geom_point() + geom_smooth(method="lm", se=FALSE) +
  labs(
    title = "Distance vs Speed",
    subtitle = "Fit to a linear model line",
    x = "Speed (mph)",
    y = "Stopping Distance (ft)") +
  theme_void()
```

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

Distance vs Speed
Fit to a linear model line



Let's plot expression data!

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

	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

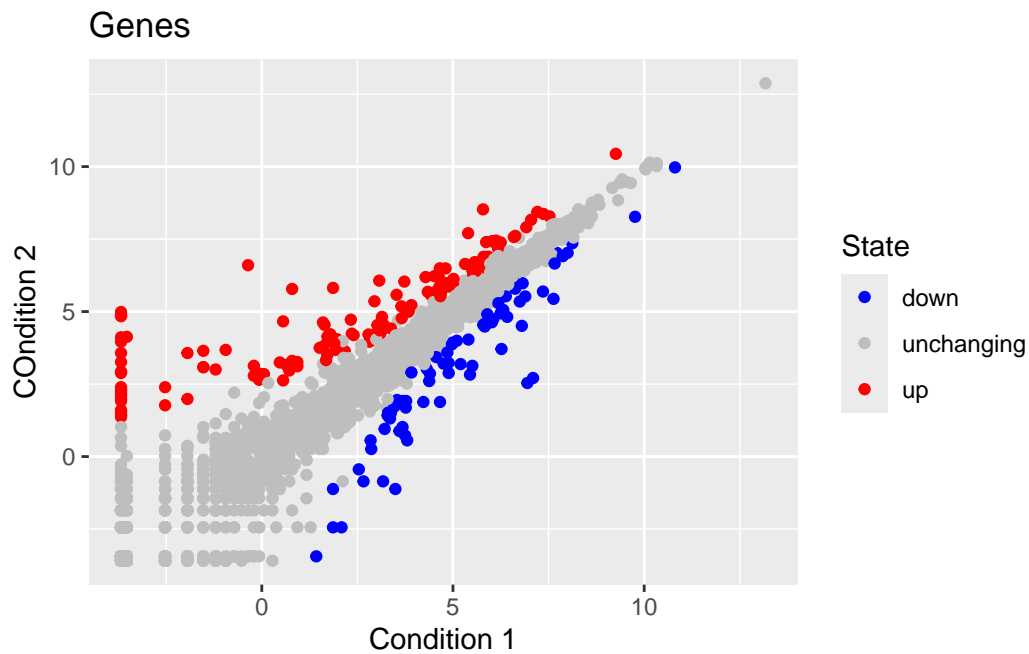
Q1. How many genes in this dataset?

```
dim(genes)
```

```
[1] 5196    4
```

There are 5196 genes in this dataset

```
ggplot(genes) + aes(Condition1, Condition2, col=State, label=Gene) + geom_point() + scale_color_manual(
  labs(
    title = "Genes",
    x = "Condition 1",
    y = "COndition 2")
  )
```



##going further

```
url <- "https://raw.githubusercontent.com/jennybc/gapminder/master/inst/extdata/gapminder.tsv"

gapminder <- read.delim(url)
```

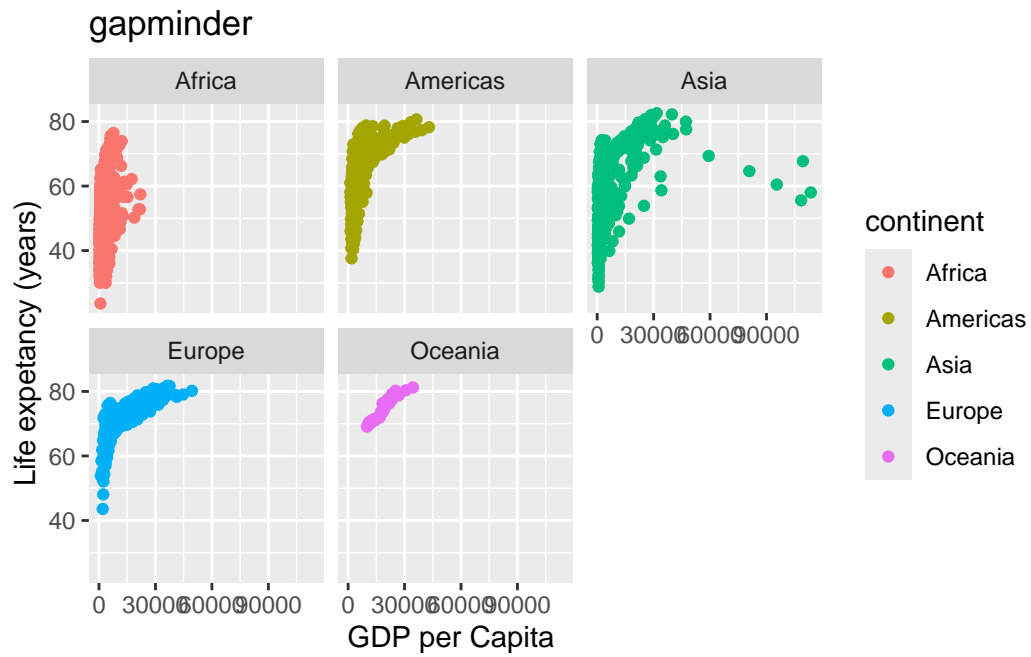
```
head(gapminder)
```

	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

```
tail(gapminder)
```

	country	continent	year	lifeExp	pop	gdpPercap
1699	Zimbabwe	Africa	1982	60.363	7636524	788.8550
1700	Zimbabwe	Africa	1987	62.351	9216418	706.1573
1701	Zimbabwe	Africa	1992	60.377	10704340	693.4208
1702	Zimbabwe	Africa	1997	46.809	11404948	792.4500
1703	Zimbabwe	Africa	2002	39.989	11926563	672.0386
1704	Zimbabwe	Africa	2007	43.487	12311143	469.7093

```
ggplot(gapminder) + aes(gdpPercap, lifeExp, col=continent, label=country) + geom_point() +
  labs(
    title = "gapminder",
    x = "GDP per Capita ",
    y = "Life expetancy (years)" + facet_wrap(~continent)
```



Seems lower GDP per capita can correlate with lower life expectancies. Capitalism...