

# 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()`.

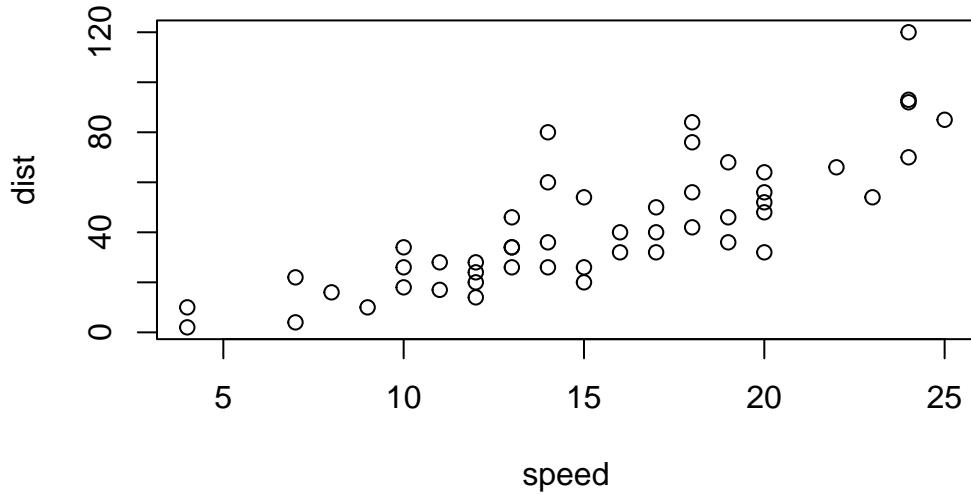
`ggplot` is a popular add-on package.

```
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)
```

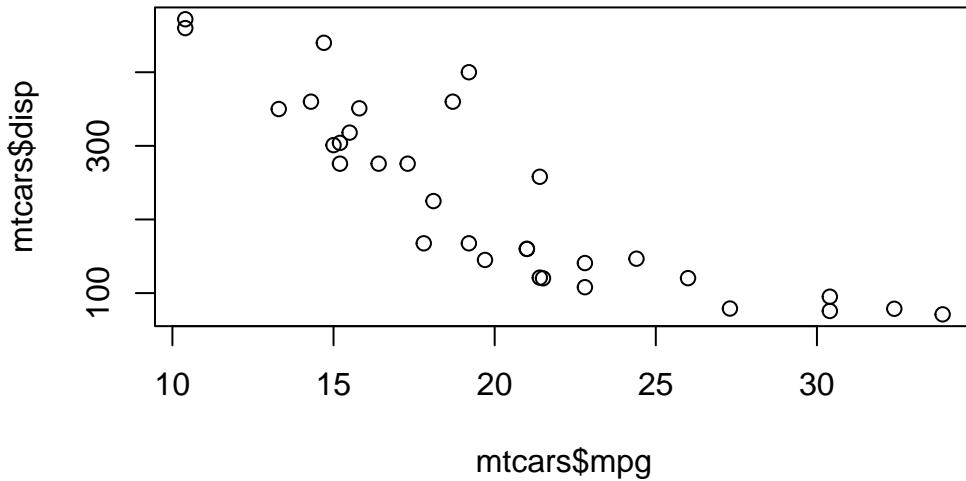


```
head(mtcars)
```

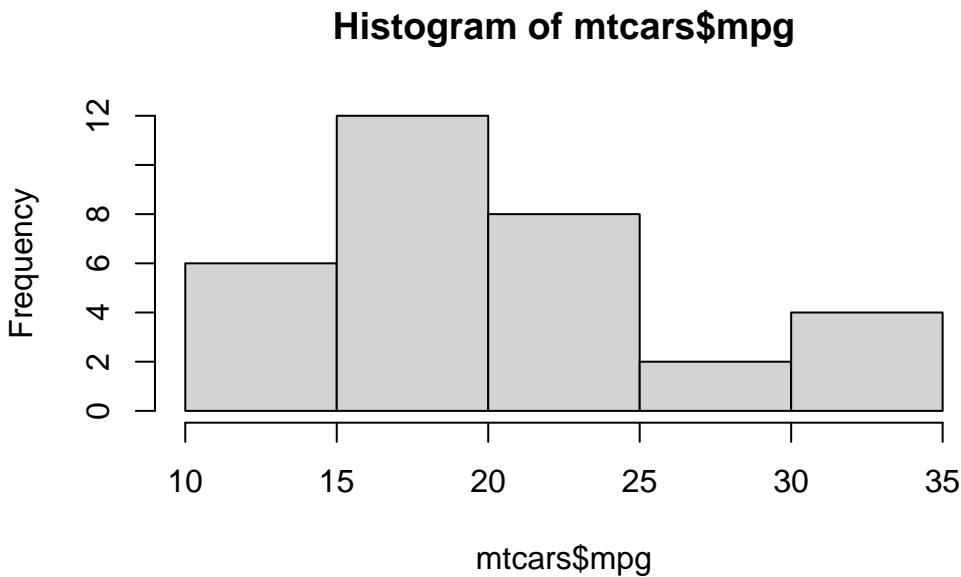
	mpg	cyl	disp	hp	drat	wt	qsec	vs	am	gear	carb
Mazda RX4	21.0	6	160	110	3.90	2.620	16.46	0	1	4	4
Mazda RX4 Wag	21.0	6	160	110	3.90	2.875	17.02	0	1	4	4
Datsun 710	22.8	4	108	93	3.85	2.320	18.61	1	1	4	1
Hornet 4 Drive	21.4	6	258	110	3.08	3.215	19.44	1	0	3	1
Hornet Sportabout	18.7	8	360	175	3.15	3.440	17.02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



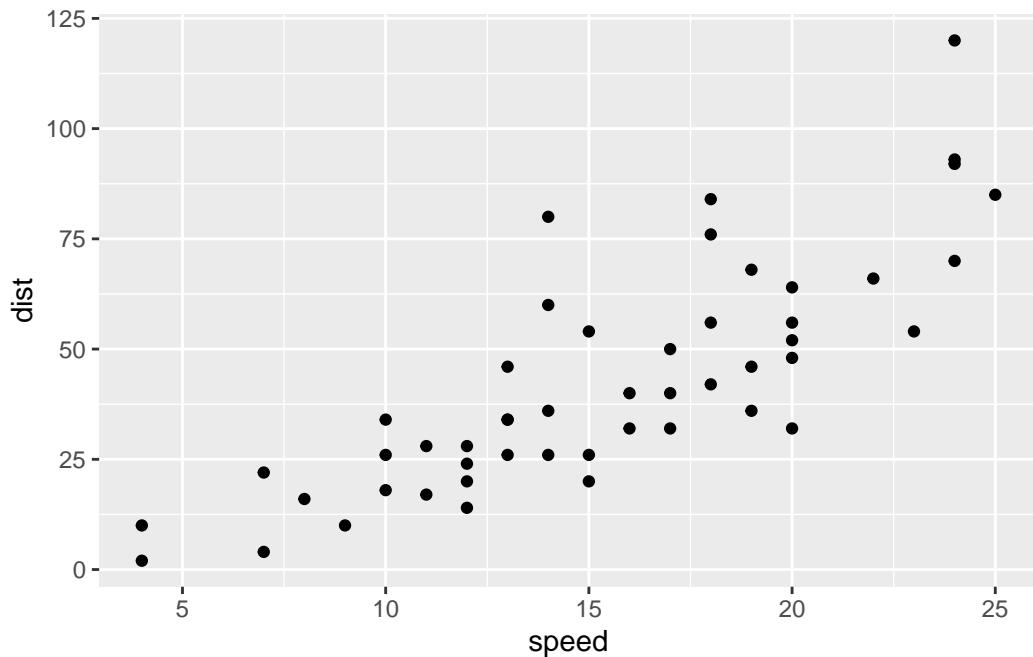
The main function on the ggplot2 package is `ggplot()`. To install the package, use the

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.

```

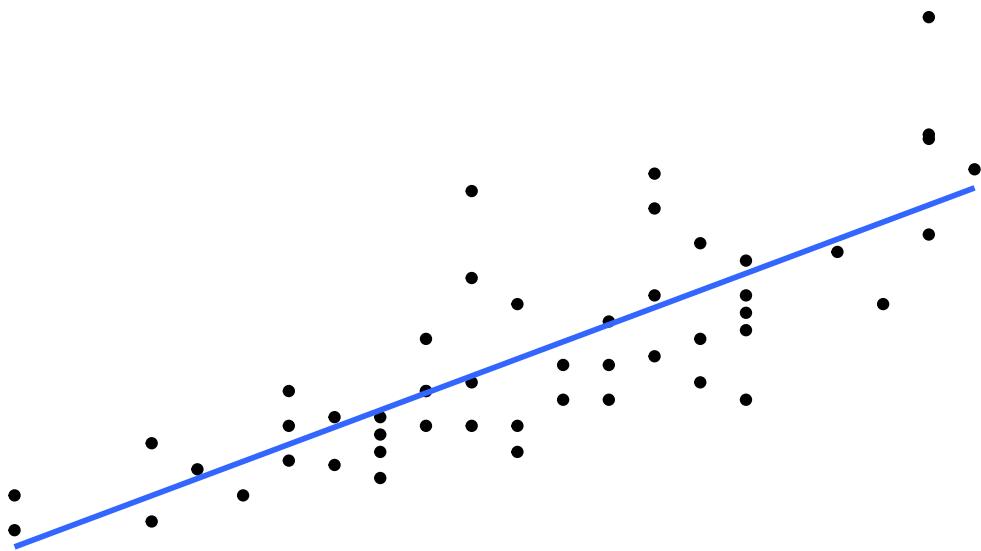
ggplot(cars) + aes(speed, dist) + geom_point() + geom_smooth(method="lm", se=FALSE) +
  labs(
    title = "Distance vs Speed",
    subtitle = "Fit to a linear model line",
    x = "Speed (mph)",
    y = "Stopping Distance (ft)") +
  theme_void()

`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

```

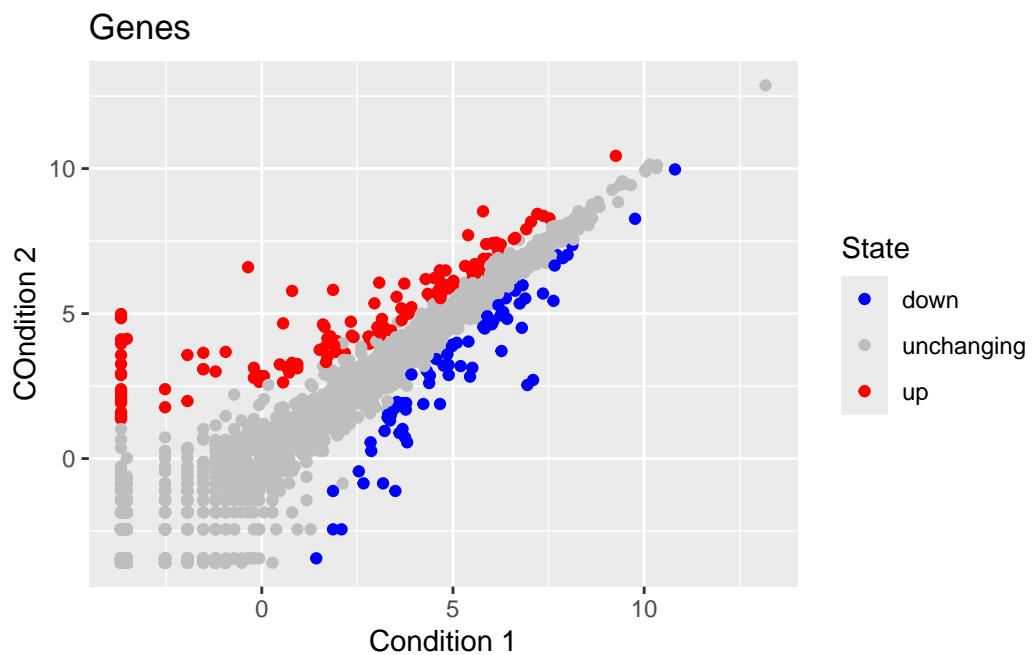
Q1. How many genes in this dataset?

```
dim(genes)
```

```
[1] 5196     4
```

There are 5196 genes in this dataset

```
ggplot(genes) + aes(Condition1, Condition2, col=State, label=Gene) + geom_point() + scale_co
```



```
##going further
```

```

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

gapminder <- read.delim(url)

head(gapminder)

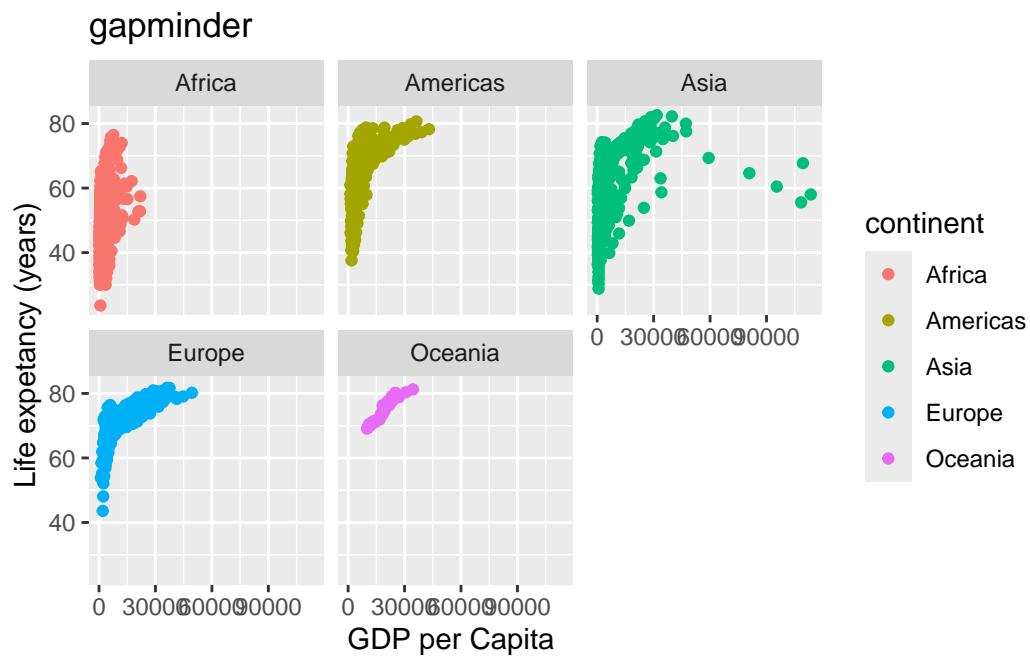
  country continent year lifeExp      pop gdpPercap
1 Afghanistan Asia 1952 28.801 8425333 779.4453
2 Afghanistan Asia 1957 30.332 9240934 820.8530
3 Afghanistan Asia 1962 31.997 10267083 853.1007
4 Afghanistan Asia 1967 34.020 11537966 836.1971
5 Afghanistan Asia 1972 36.088 13079460 739.9811
6 Afghanistan 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 expectancy (years)") + facet_wrap(~continent)

```



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