

CLASS 5 DATA VIZ WITH GGPLOT

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Background

There are lots of ways to make plots in R. These include so-called “base R” (like the `plot()`) and add on packages like `ggplot2`

Lets make the same plot with these two graphics systems. We can use the inbuilt `cars` dataset:

```
head(cars)
```

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

With “base R” we can simply:

```
plot(cars)
```



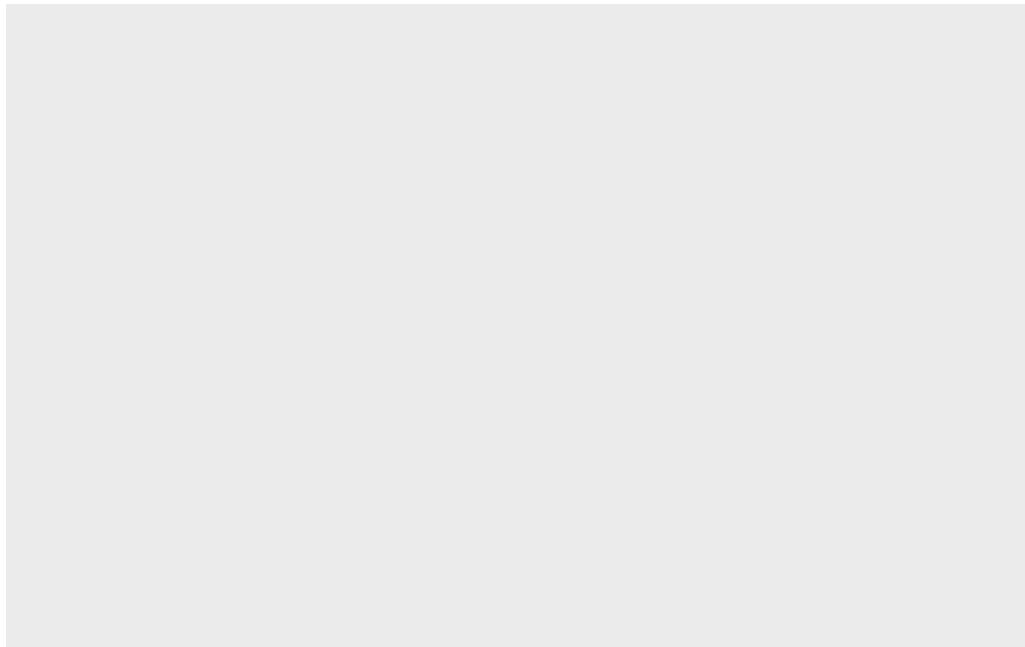
Now lets try ggplot. First I need to install the package using `install.packages("ggplot2")'`
>> ****N.B.** We never run `install.packages()`'** in a code chunk that otherwise we will re-install needlessly every time we render our document

every time we want to use an add-on package we need to load it up with a call to `'library()'`

```
library(ggplot2)
```

Warning: package 'ggplot2' was built under R version 4.4.3

```
ggplot(cars)
```

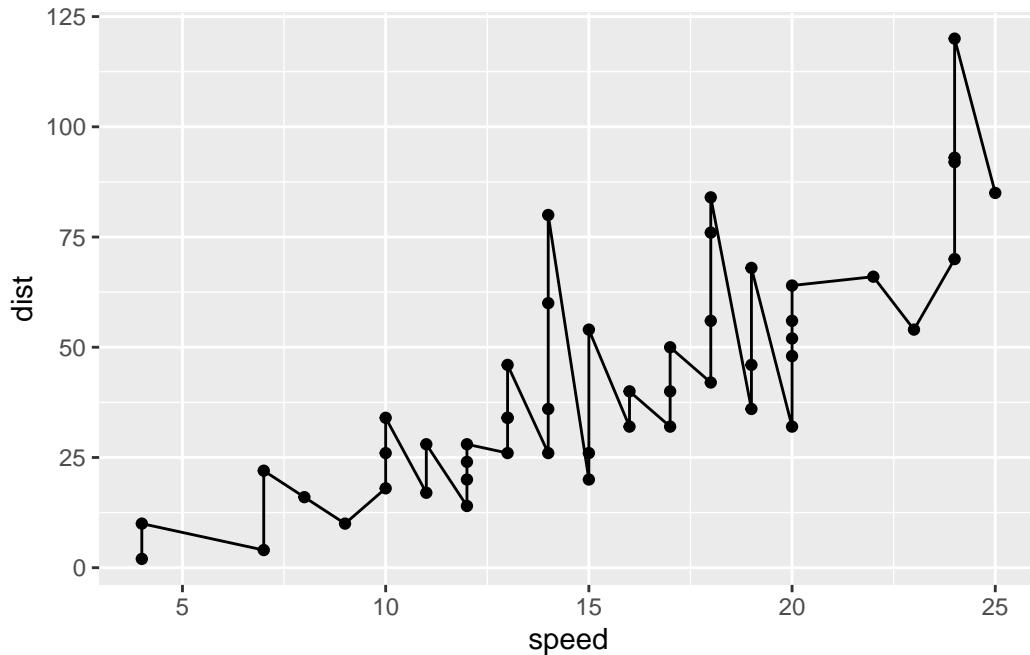


Every ggplot needs at least three things:

1. the **data** i.e. stuff to plot as a data.frame
2. the **aes** or aesthetics that map the data to the plot
3. the **geom_** or geometry i.e. the plot type such as points, lines etc.

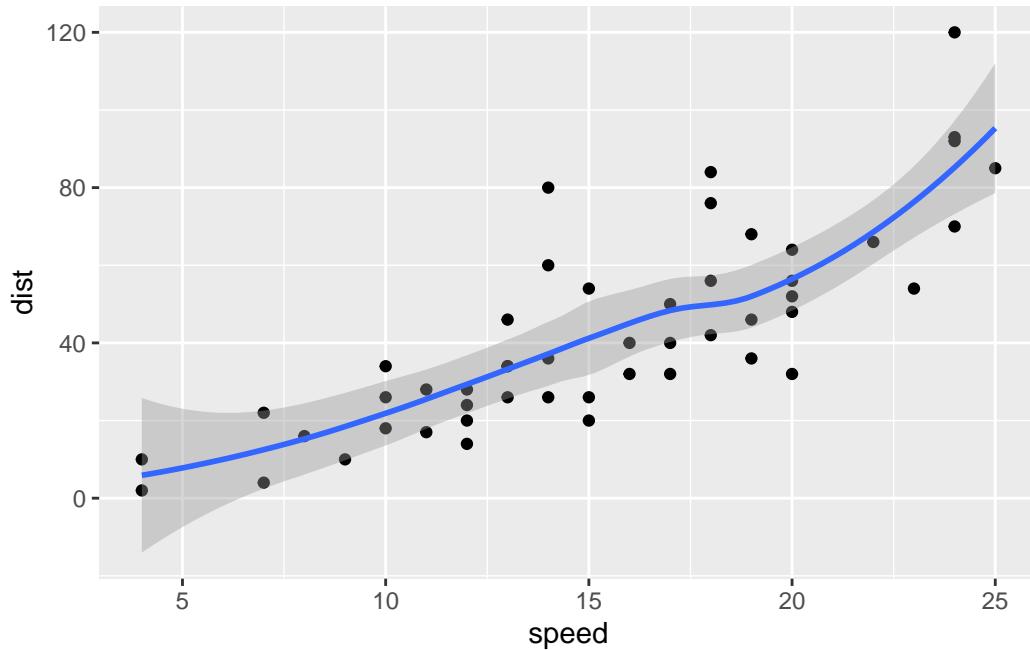
```
install.packages("ggplot2")
```

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



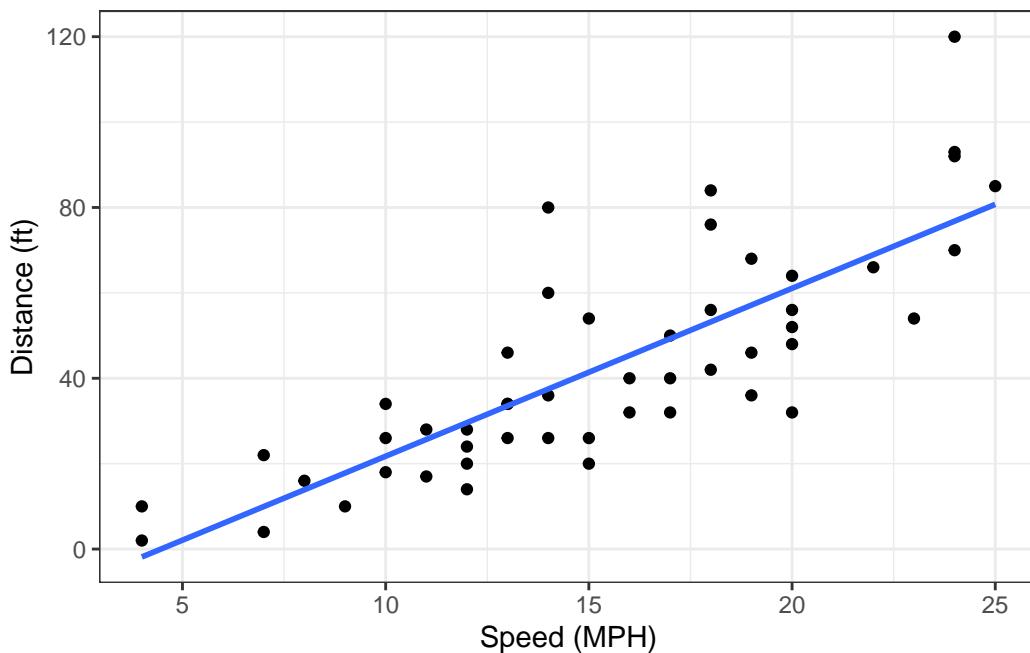
```
ggplot(cars) + aes(x=speed, y=dist) + geom_point() +geom_smooth ()
```

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



```
ggplot(cars) +
  aes(x=speed, y=dist) + geom_point()+
  geom_smooth(method = "lm", se=FALSE) +
  labs(x="Speed (MPH)", y ="Distance (ft)", Title = "stopping Distance of old Cars") + theme
```

Ignoring unknown labels:
 * Title : "stopping Distance of old Cars"
`geom_smooth()` using formula = 'y ~ x'



gene expression plot

read some data on the effects of GLP-1 inhibitor (drug) on gene expression values:

```
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

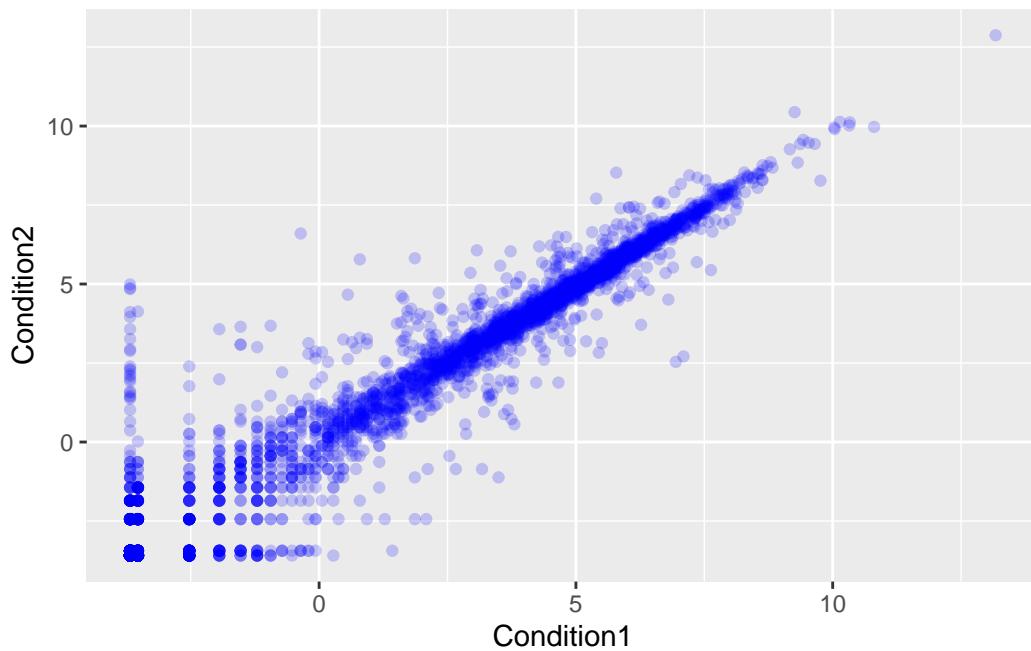
```

Version 1 plot - start simple by getting some ink on the page

```

url <- "https://bioboot.github.io/bimm143_S20/class-material/up_down_expression.txt"
genes <- read.delim(url)
ggplot(genes) +
  aes(x= Condition1, y=Condition2) +
  geom_point(col="blue", alpha = 0.2)

```



Let's color by **State** up, down or no change

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

	down	unchanging	up
72	4997	127	

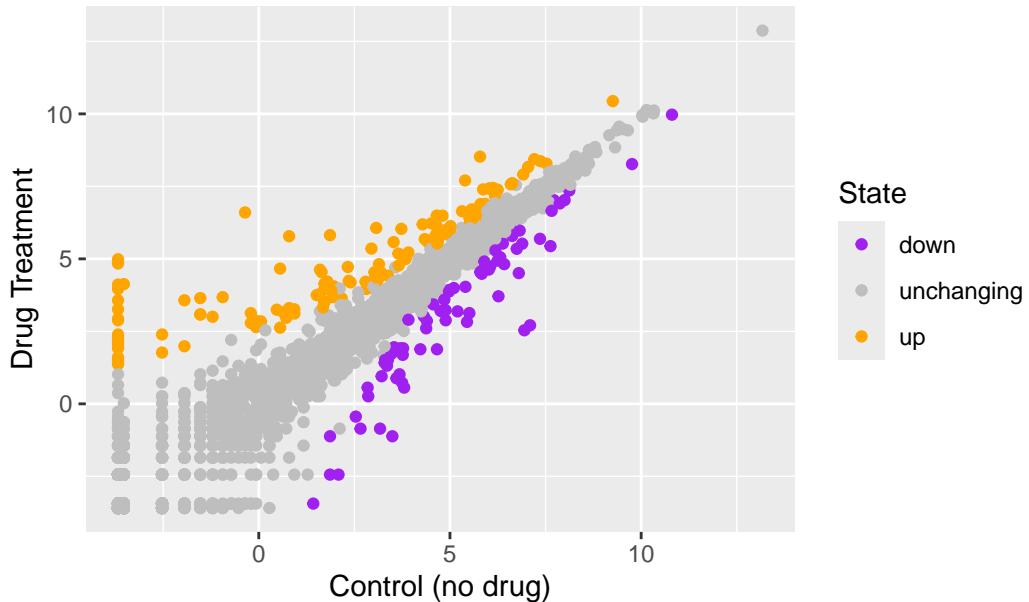
```
library(ggplot2)
```

```

url <- "https://bioboot.github.io/bimm143_S20/class-material/up_down_expression.txt"
genes <- read.delim(url)
ggplot(genes) + aes(x=Condition1, y=Condition2, col=State) + geom_point() + scale_color_manual

```

Gene Expression Changes Upon Drug Treatment



going further with gapminder

here we explore the famous gapminder

```

# File location online
url <- "https://raw.githubusercontent.com/jennybc/gapminder/master/inst/extdata/gapminder.csv"
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

Q. How many rows does this dataset have?

```
nrow(gapminder)
```

```
[1] 1704
```

How many different continents are in this dataset

```
table(gapminder$continent)
```

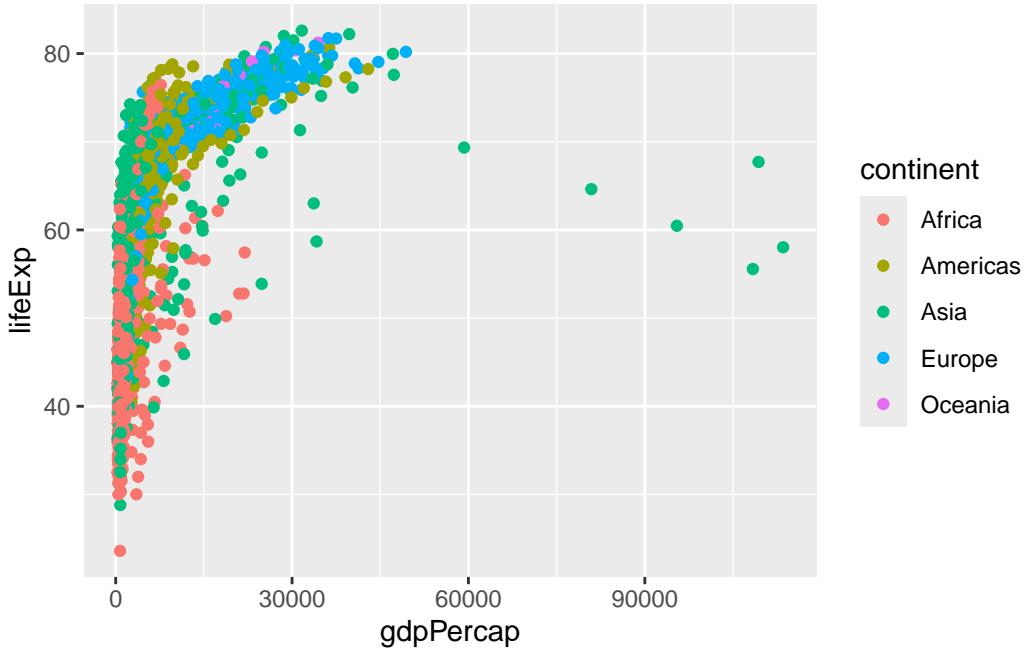
	Africa	Americas	Asia	Europe	Oceania
	624	300	396	360	24

Version 1 plot gdpPercap vs. LifeExp for all rows

```
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

```
ggplot(gapminder) + aes(x = gdpPercap, lifeExp, col=continent) + geom_point()
```

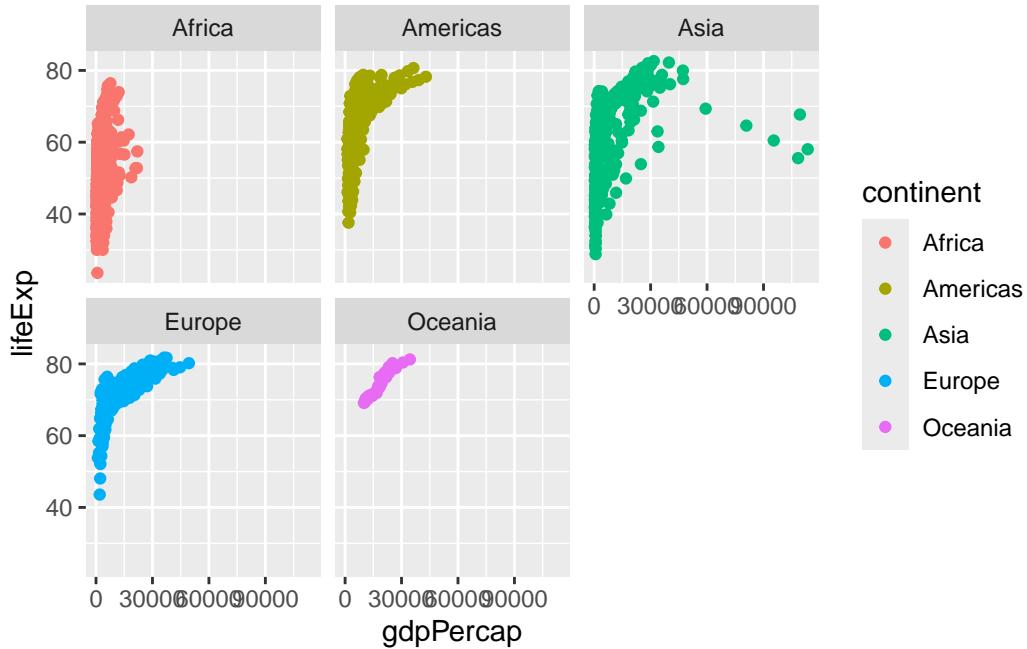


I want to see a plot for each continent – in ggplot language this is called “faceting”

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

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



```
##First look at the dplyr package
```

Another add on package with a function called ‘filter()’ that we want to use

```
install.packages("dplyr")
```

```
library(dplyr)
```

```
Warning: package 'dplyr' was built under R version 4.4.3
```

```
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
```

```
url <- "https://raw.githubusercontent.com/jennybc/gapminder/master/inst/extdata/gapminder.ts"
gapminder <- read.delim(url)
filter(gapminder, year == 2007, country == "Ireland")
```

```
country continent year lifeExp      pop gdpPercap
1 Ireland     Europe 2007  78.885 4109086      40676
```

```
url <- "https://raw.githubusercontent.com/jennybc/gapminder/master/inst/extdata/gapminder.ts"
gapminder <- read.delim(url)
filter(gapminder, year == 2007, country == "United States")
```

```
country continent year lifeExp      pop gdpPercap
1 United States   Americas 2007  78.242 301139947  42951.65
```

```
url <- "https://raw.githubusercontent.com/jennybc/gapminder/master/inst/extdata/gapminder.ts"
gapminder <- read.delim(url)
input <- filter(gapminder, year == 2007 | year == 1977)
```

```
library(ggplot2)
url <- "https://raw.githubusercontent.com/jennybc/gapminder/master/inst/extdata/gapminder.ts"
gapminder <- read.delim(url)
input <- filter(gapminder, year == 2007 | year == 1977)
ggplot(input) +
  geom_point(aes(x = gdpPercap, y= lifeExp, color=continent)) +
  facet_wrap(~year)
```

