Class05: Data Visualization w/ ggplot

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Graphics Systems in R

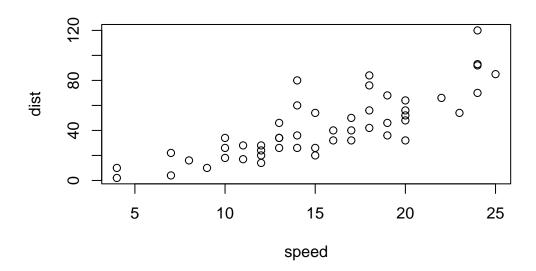
There are many graphics systems in R for making plots and figures.

We have already played around with "base R" graphics and the plot() function

Today we will start learning about a popular graphics pkg called ggplot2().

This is an add on package - i.e. we need to install it. I install it like I install any package: install.packages()

plot(cars)



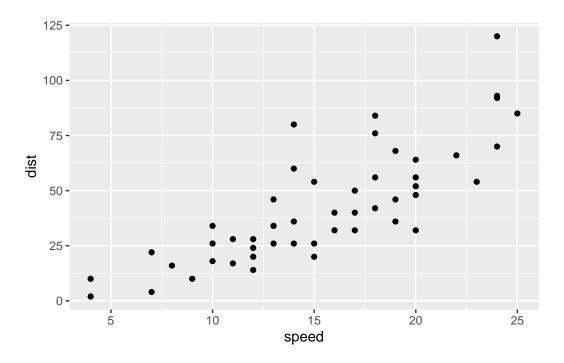
Before I can use the functions from a package, I have to load up the package from my library. We use the library(ggplot2) command to load it up.

```
#install.packages(ggplot2)
library(ggplot2)
ggplot(cars)
```

Every ggplot is made up of at least 3 things:

- data: the values that will go into your plot
- aes (aesthetics): how the columns of data map to the plot aesthetics (specifies x and y, color, etc.)
- geoms (**geometries**): how the plot actually looks; graph type (points, lines, or columns, etc.)

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



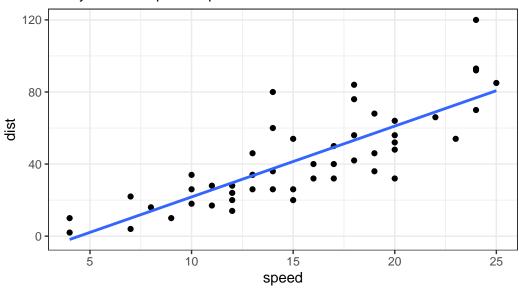
For simple plots, ggplot is more verbose than the base R plot.

Add more layers to our ggplot:

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

Stopping distance of Old Cars

A silly little example of a plot



Lab5 Gene Scatterplot Portion

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

[1] 5196

ncol(genes)

[1] 4

```
num_upreg = sum(genes$State == 'up') #number of upregulated genes
num_upreg
```

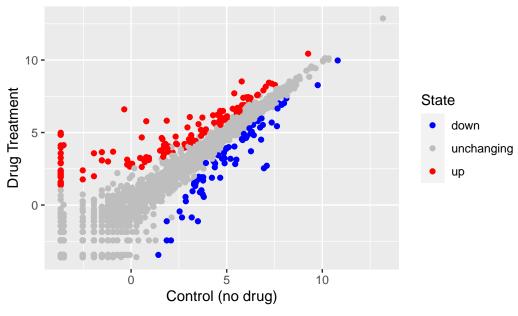
[1] 127

```
frac\_upreg = round(num\_upreg / nrow(genes) * 100, 2) #percentage of upregulated genes frac\_upreg
```

[1] 2.44

```
ggplot(genes) +
   aes(x=Condition1, y=Condition2, col =State) +
   scale_colour_manual( values=c("blue", "gray", "red") ) +
   geom_point() +
   labs(title="Gene Expression Changes Upon Drug Treatment", x="Control (no drug)", y="Drug")
```

Gene Expression Changes Upon Drug Treatment

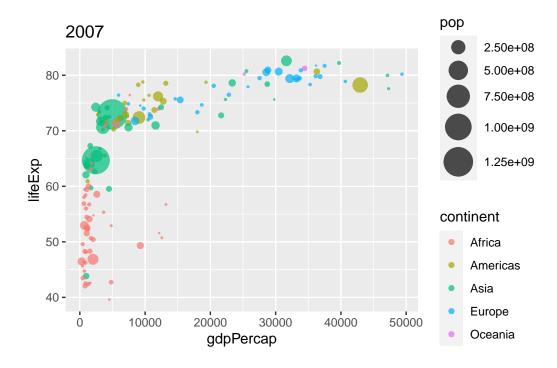


Part 7

```
#install.packages(gapminder)
#install.packages(dplyr)

library(gapminder)
library(dplyr)
```

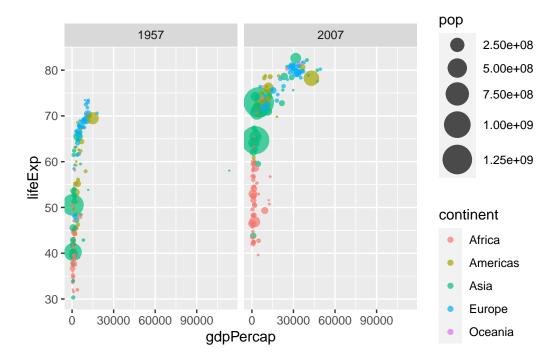
```
Attaching package: 'dplyr'
The following objects are masked from 'package:stats':
    filter, lag
The following objects are masked from 'package:base':
    intersect, setdiff, setequal, union
  # File location online
  url <- "https://raw.githubusercontent.com/jennybc/gapminder/master/inst/extdata/gapminder.
  gapminder <- read.delim(url)</pre>
  gapminder_2007 <- gapminder %>% filter(year==2007)
2007 dataset
  p <- ggplot(gapminder_2007) +</pre>
    aes(x=gdpPercap, y=lifeExp, color=continent, size=pop) +
    geom_point(alpha=0.7) +
    scale_size_area(max_size = 10) +
    labs(title=2007)
  p
```



and 2007 dataset

```
gapminder_1957 <- gapminder %>% filter(year==1957 | year==2007)

q <- ggplot(gapminder_1957) +
   aes(x=gdpPercap, y=lifeExp, color=continent, size=pop) +
   geom_point(alpha=0.7) +
   scale_size_area(max_size = 10) +
   facet_wrap(~year)
q</pre>
```



Part 8 Bar Charts

5 Largest Population Countries

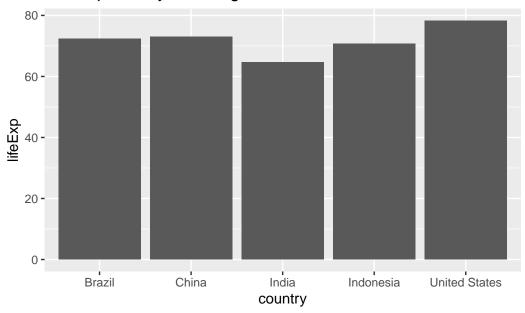
```
gapminder_top5 <- gapminder %>%
  filter(year==2007) %>%
  arrange(desc(pop)) %>%
  top_n(5, pop)

gapminder_top5
```

```
country continent year lifeExp
                                             pop gdpPercap
1
         China
                    Asia 2007 72.961 1318683096
                                                  4959.115
2
         India
                    Asia 2007 64.698 1110396331
                                                  2452.210
3 United States Americas 2007 78.242
                                       301139947 42951.653
4
     Indonesia
                    Asia 2007
                              70.650
                                       223547000
                                                  3540.652
5
        Brazil Americas 2007 72.390
                                      190010647
                                                  9065.801
```

```
ggplot(gapminder_top5) +
  geom_col(aes(x = country, y = lifeExp)) +
  labs(title="Life Expectancy of 5 Largest Countries in 2007")
```

Life Expectancy of 5 Largest Countries in 2007



```
r <- ggplot(gapminder_top5) +
  aes(x=reorder(country, -pop), y=pop, fill=gdpPercap) +
  geom_col() +
  labs(title="Population of 5 Largest Countries in 2007 With GDP per Capita")
r</pre>
```

Population of 5 Largest Countries in 2007 With GDP per Cap

