

# Class 5: Data Vis with ggplot

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## Intro to ggplot

There are many graphics systems in R (ways to make plots and figures). These include “base” R plots. Today we will focus mostly on the **ggplot2** package.

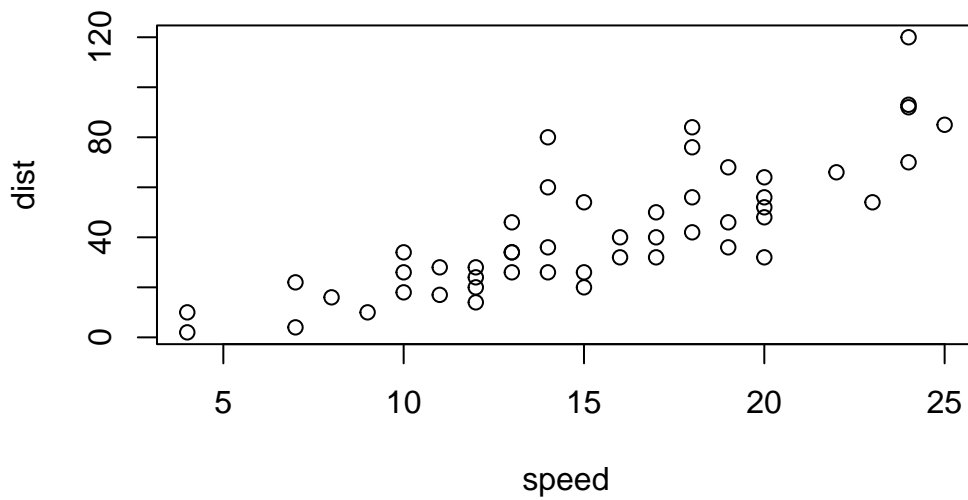
Let’s start with a plot of a simple in-built dataset called ‘cars’.

```
cars
```

	speed	dist
1	4	2
2	4	10
3	7	4
4	7	22
5	8	16
6	9	10
7	10	18
8	10	26
9	10	34
10	11	17
11	11	28
12	12	14
13	12	20
14	12	24
15	12	28
16	13	26
17	13	34
18	13	34
19	13	46
20	14	26
21	14	36
22	14	60

23	14	80
24	15	20
25	15	26
26	15	54
27	16	32
28	16	40
29	17	32
30	17	40
31	17	50
32	18	42
33	18	56
34	18	76
35	18	84
36	19	36
37	19	46
38	19	68
39	20	32
40	20	48
41	20	52
42	20	56
43	20	64
44	22	66
45	23	54
46	24	70
47	24	92
48	24	93
49	24	120
50	25	85

```
plot(cars)
```

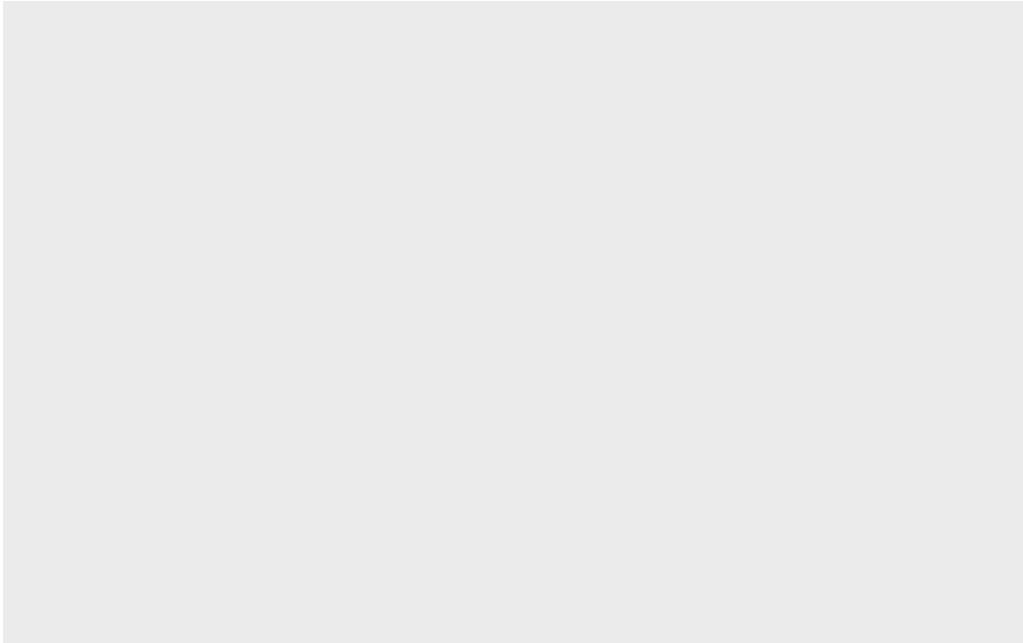


Let's see how we can make this figure using **ggplot**. First I need to install this package on my computer. To install any R package I use the function `install.packages()`

I will run `install.packages()` in my R console, not this Quarto so I don't have to reinstall every time that I render.

Before I can use any functions from add-on packages I need to load the package from my "library()" with the "library(ggplot2)" call.

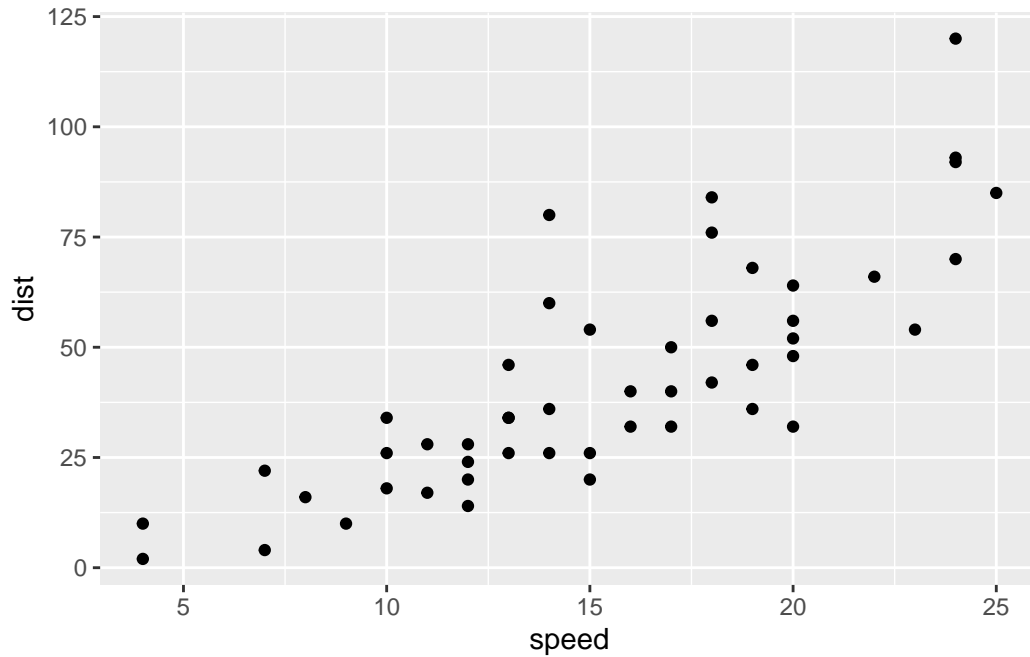
```
library(ggplot2)
ggplot(cars)
```



All ggplot figures have at least 3 things (called layers). These include:

- **data** (the input dataset I want to plot from)
- **aes** (the aesthetic mapping of the data to my plot)
- **geoms** (the `geom_point()`, `geom_line()`, etc that I want to draw from )

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

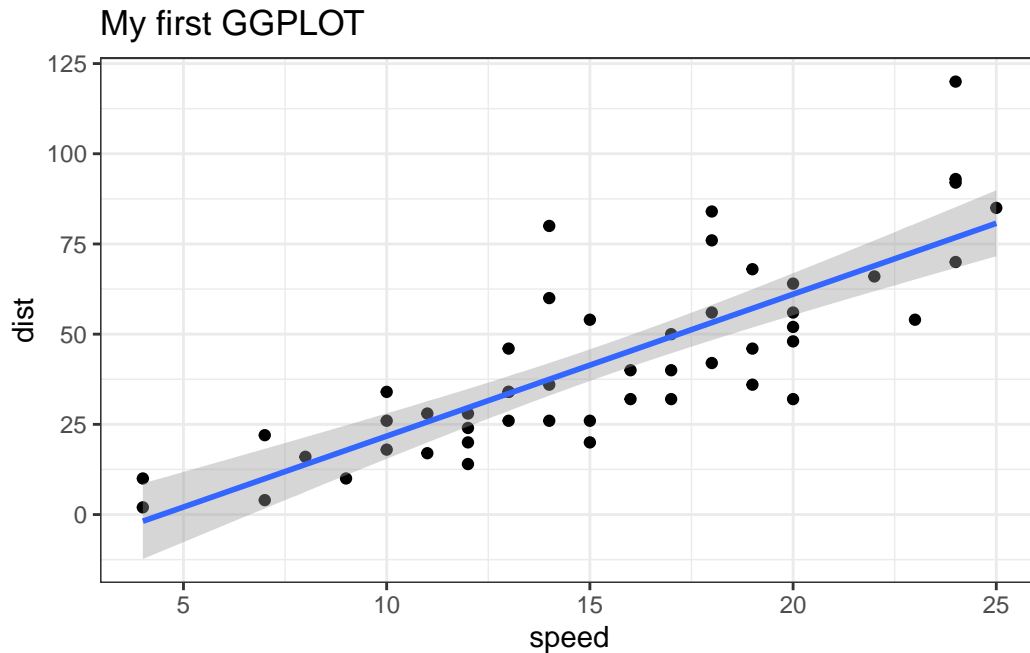


Use base R plots for data that you want to see but don't need a nice, polished graph. `ggplot` is more work, but more editable for nicer representations.

Let's add a line to show the relationship here:

```
ggplot(cars) +  
  aes(x= speed, y= dist) +  
  geom_point() +  
  geom_smooth (method="lm") +  
  theme_bw() +  
  labs(title= "My first GGPlot")
```

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



1. Which geometric layer should be used to create scatter plots in ggplot2?

`geom_point()`

## Gene expression figure

The code to read the dataset, retrieving online from class dataset

```
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

How many genes are in this dataset?

```
nrow(genes)
```

```
[1] 5196
```

A first plot of this dataset

```
p <- ggplot(genes) +  
  aes(x=Condition1, y=Condition2, col= State) +  
  geom_point()  
  
colnames(genes)
```

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

```
ncol(genes)
```

```
[1] 4
```

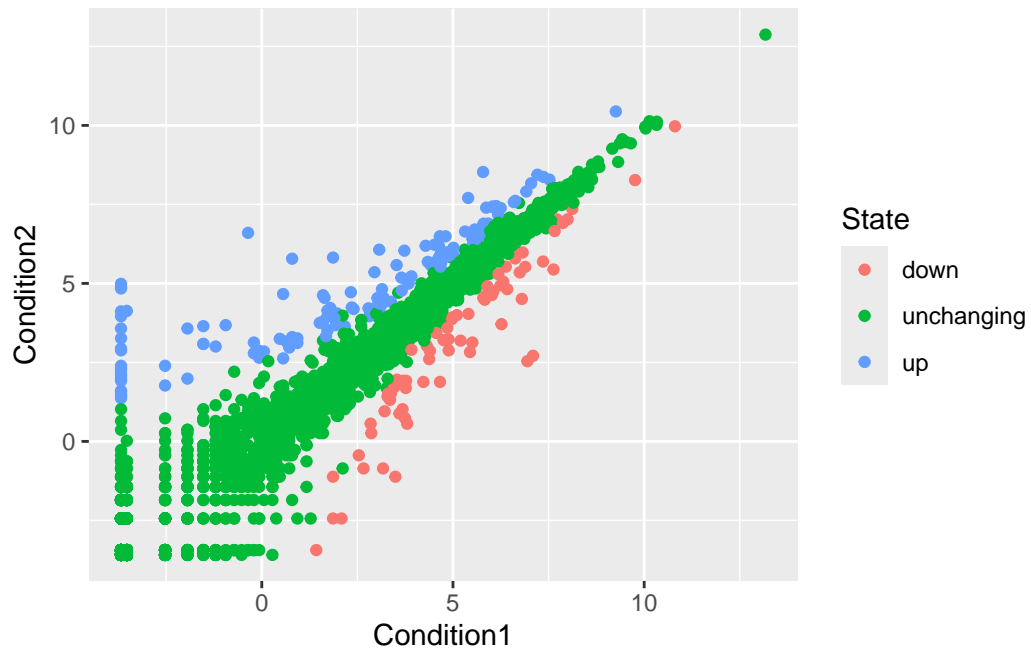
```
## Graphical representation of the State values  
table(genes$State)
```

down	unchanging	up
72	4997	127

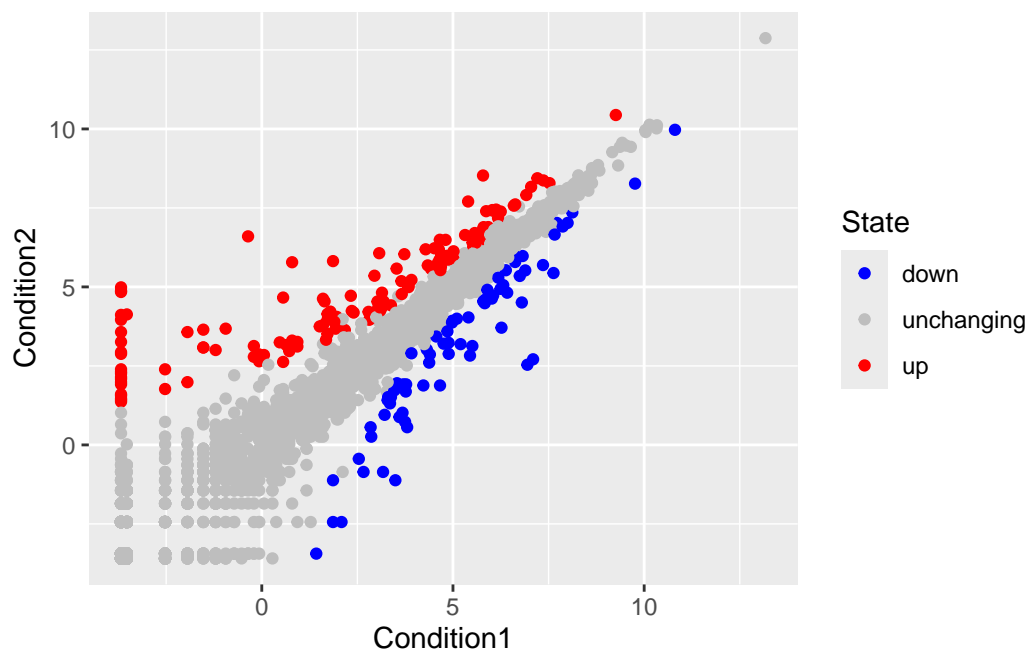
```
## Graphical representation of the State values div by total # of genes  
round(table(genes$State)/nrow(genes) *100, 2)
```

down	unchanging	up
1.39	96.17	2.44

```
## Default color scheme  
p
```



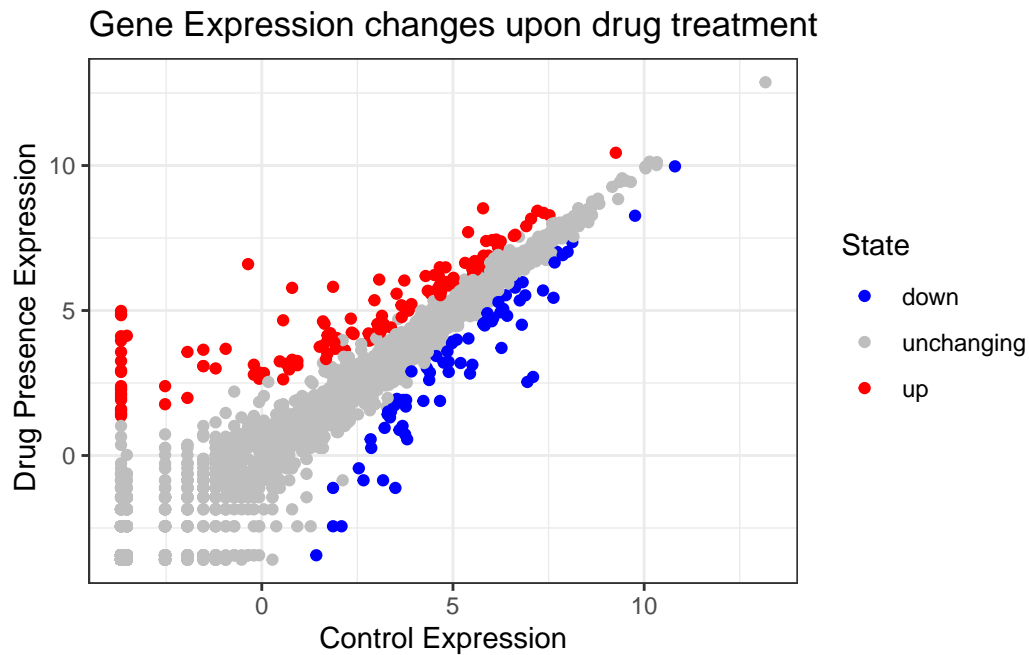
```
## Personalized color scheme
p + scale_color_manual(values= c("blue", "gray", "red"))
```



Lets add some labels and a title

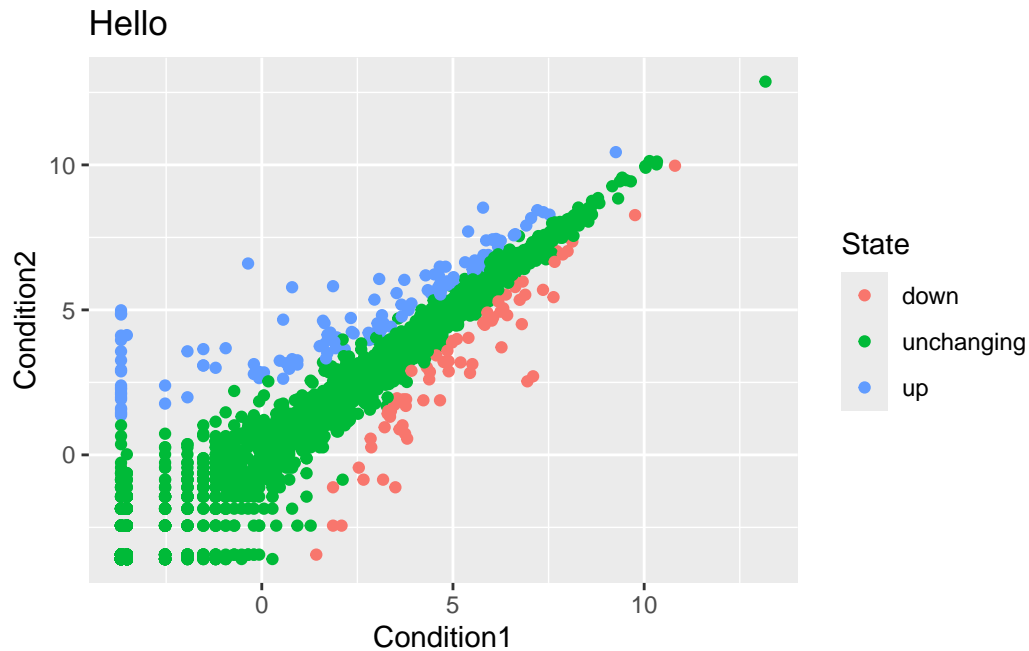


```
ggplot(genes) +
  aes(x=Condition1, y=Condition2, col= State) +
  geom_point() +
  theme_bw() +
  labs(title= "Gene Expression changes upon drug treatment",
        x= "Control Expression",
        y= " Drug Presence Expression") +
  scale_color_manual(values= c("blue", "gray", "red"))
```



Defining the graph as an object makes it easier to edit or have multiple versions without having to rewrite the entire code again

```
p + labs(title="Hello")
```



## 7. Going further

```
# File location online
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

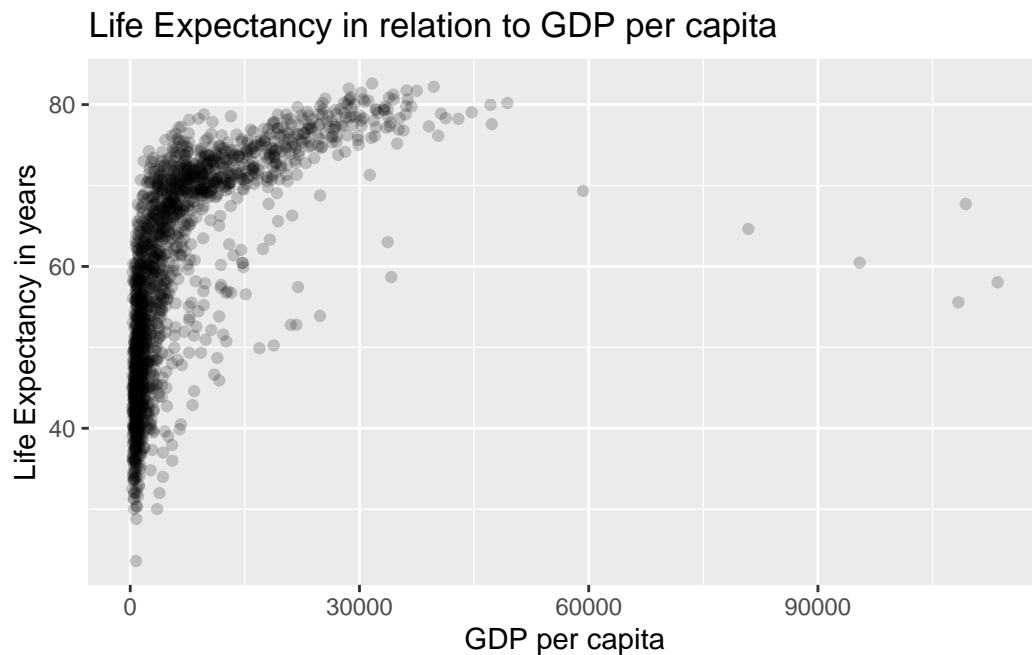
```
#install.packages("dplyr") ## un-comment to install if needed
#library(dplyr)

## alpha function edits transparency of points, on a scale for 0 - transparent, to 1 - full
ggplot(gapminder) +
```

```

aes(x=gdpPercap, y=lifeExp) +
geom_point(alpha= 0.2) +
labs (title= "Life Expectancy in relation to GDP per capita", x= "GDP per capita", y= "Life Expectancy in years")

```



More specific, one year only

```

## even though we installed dplyr in the brain below, we need to pull it from the library to
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)
```

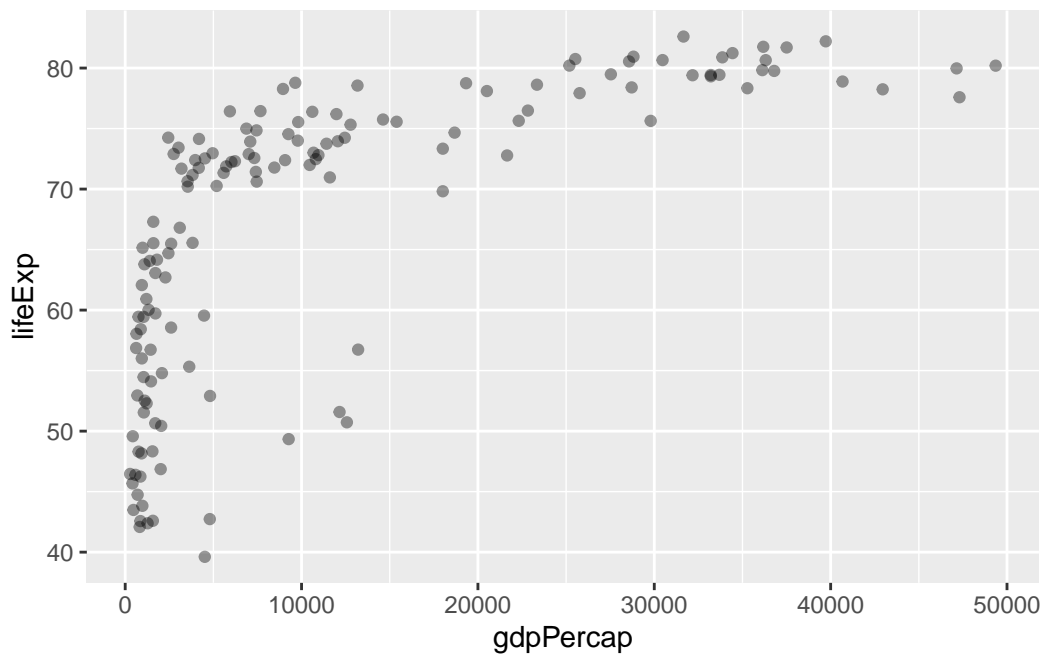
*## There are a few new functions that have the same name as other ones in R, replacing it wi*

Now we can see values only from 2007

```
head(gapminder_2007)
```

	country	continent	year	lifeExp	pop	gdpPercap
1	Afghanistan	Asia	2007	43.828	31889923	974.5803
2	Albania	Europe	2007	76.423	3600523	5937.0295
3	Algeria	Africa	2007	72.301	33333216	6223.3675
4	Angola	Africa	2007	42.731	12420476	4797.2313
5	Argentina	Americas	2007	75.320	40301927	12779.3796
6	Australia	Oceania	2007	81.235	20434176	34435.3674

```
ggplot(gapminder_2007) +  
  aes(x=gdpPercap, y=lifeExp) +  
  geom_point(alpha=0.4)
```



Adding more variables to aes()

```
ggplot(gapminder_2007) +
  aes(x=gdpPercap, y=lifeExp, color=continent, size=pop) +
  geom_point(alpha=0.4) +
  labs(title= "2007 Life Exp vs GDP/capita")
```

