

# Class05: Data Viz with GGPLOT

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Today we are playing and plotting graphics with R.

There are lots of ways to make cool figures in R. There is “base” R graphics (`plot()`, `hist()`, `boxplot()`, etc.)

There is also add-on packages, like `ggplot`

```
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)
```

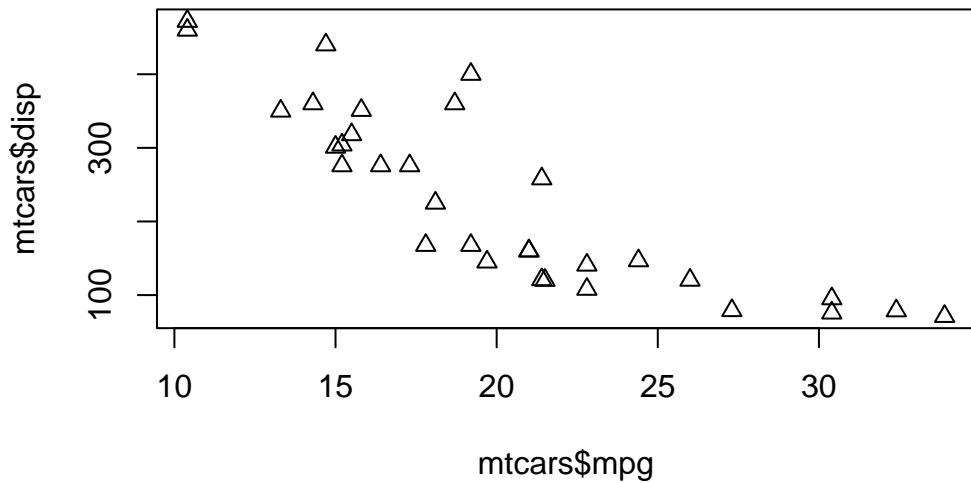


```
head(mtcars)
```

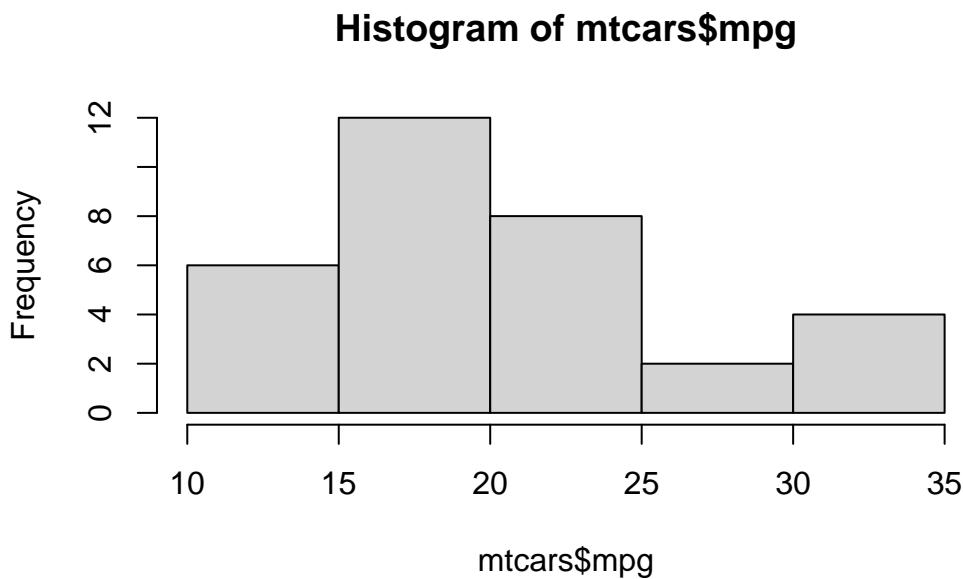
	mpg	cyl	disp	hp	drat	wt	qsec	vs	am	gear	carb
Mazda RX4	21.0	6	160	110	3.90	2.620	16.46	0	1	4	4
Mazda RX4 Wag	21.0	6	160	110	3.90	2.875	17.02	0	1	4	4
Datsun 710	22.8	4	108	93	3.85	2.320	18.61	1	1	4	1
Hornet 4 Drive	21.4	6	258	110	3.08	3.215	19.44	1	0	3	1
Hornet Sportabout	18.7	8	360	175	3.15	3.440	17.02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

```
plot(mtcars$mpg, mtcars$disp, pch=2)
```



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

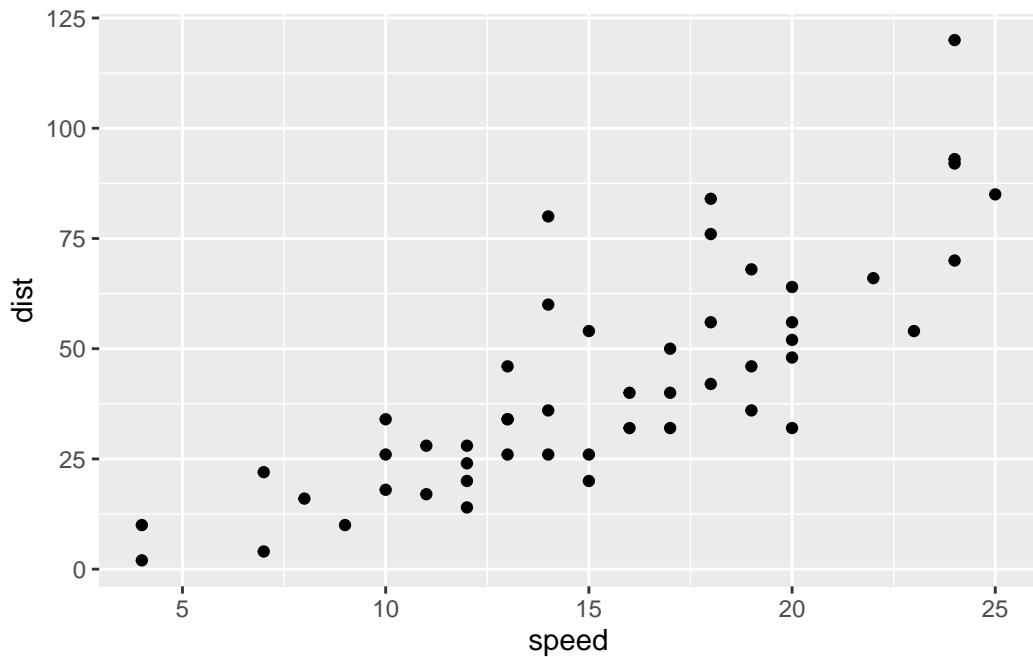


## GGPLOT

The main function in the ggplot2 package is `ggplot()`. First I need to install the `ggplot2` package. I can install any package with the function `install.packages()`.

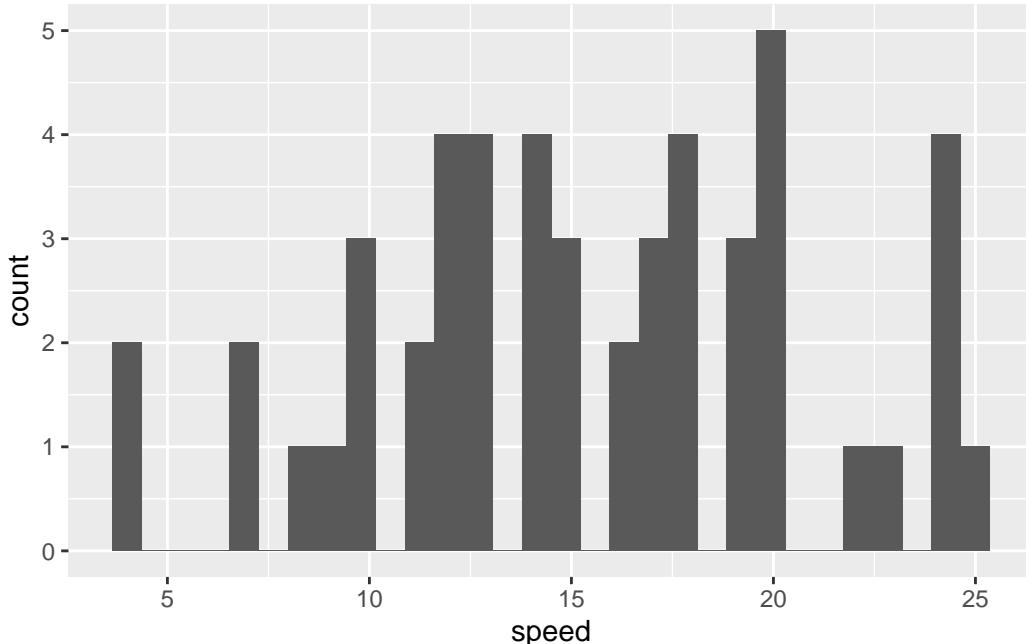
**N.B.** I never want to run `install.packages()` in my quarto source document!!!

```
library(ggplot2)
ggplot(cars) +
  aes(x=speed, y=dist) +
  geom_point()
```



```
ggplot(cars) +
  aes(speed) +
  geom_histogram()
```

``stat_bin()`` using ``bins = 30``. Pick better value ``binwidth``.



Every ggplot needs at least 3 things:

- The **data** (given with `ggplot(cars)`)
- The **aesthetics** mapping (given with `aes()`)
- The **geom** (given with `geom_point()`)

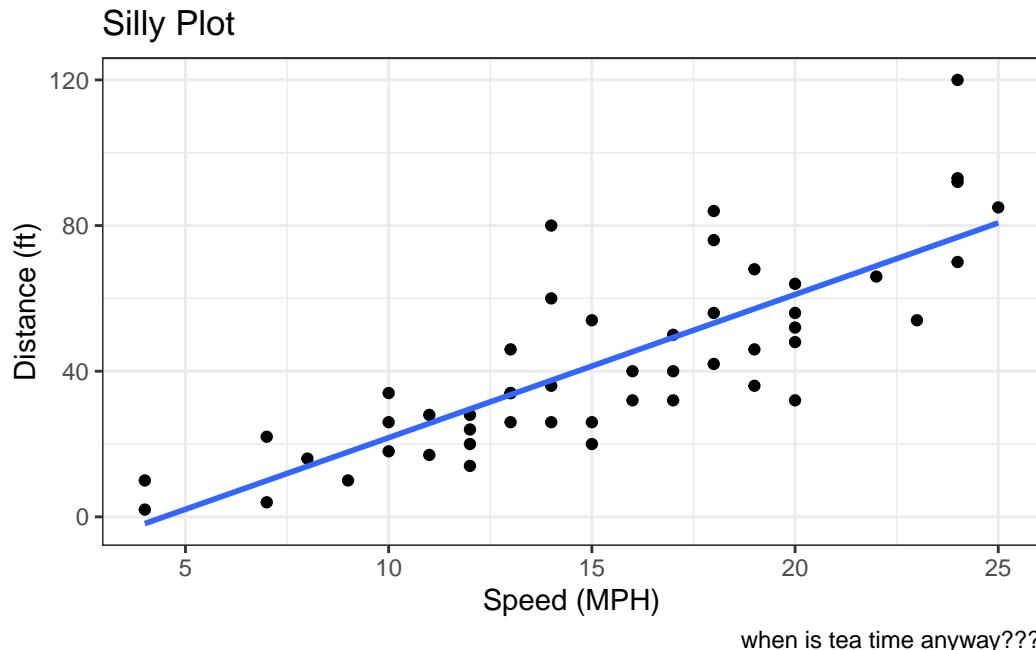
For simple canned graphs, “base” R is nearly always faster

### Adding more layers

Let’s add a line and a title, subtitle and caption as well as custom axis labels

```
ggplot(cars) +
  aes(x=speed, y=dist) +
  geom_point() +
  geom_smooth(method="lm", se=FALSE) +
  labs(title = "Silly Plot",
       x = "Speed (MPH)",
       y = "Distance (ft)",
       caption="when is tea time anyway??") +
  theme_bw()

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



### Plot some expression data

Read data file from online URL

```
url <- "https://bioboot.github.io/bimml43_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 are in this wee dataset?

There are 5196 in this dataset

Q2. How many “up” regulated genes are there?

```
sum(genes$State == "up")
```

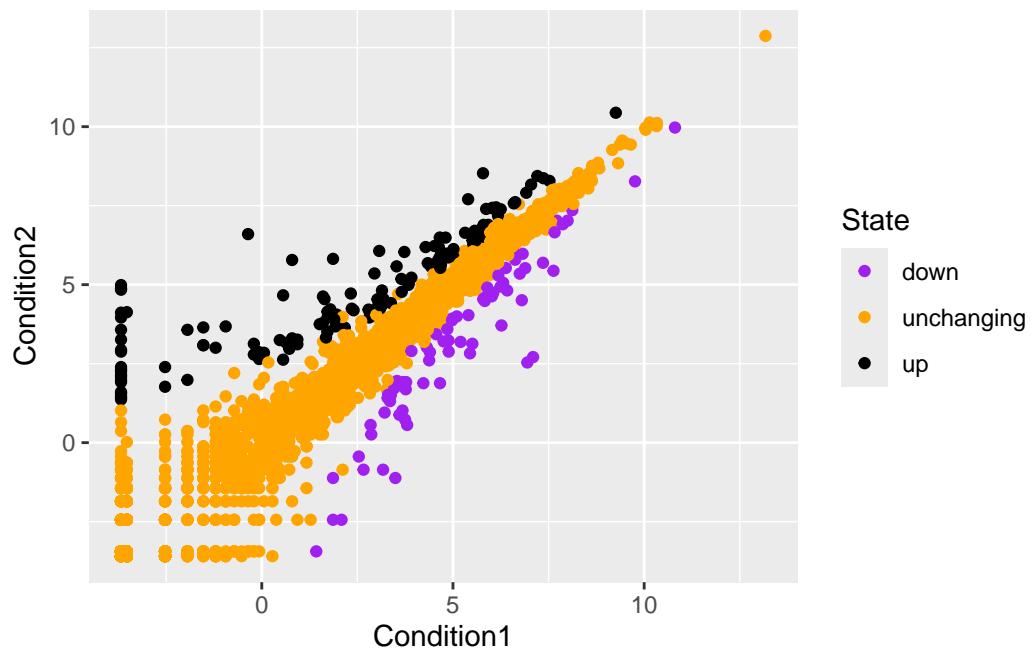
```
[1] 127
```

There are 127 “up” regulated genes.

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

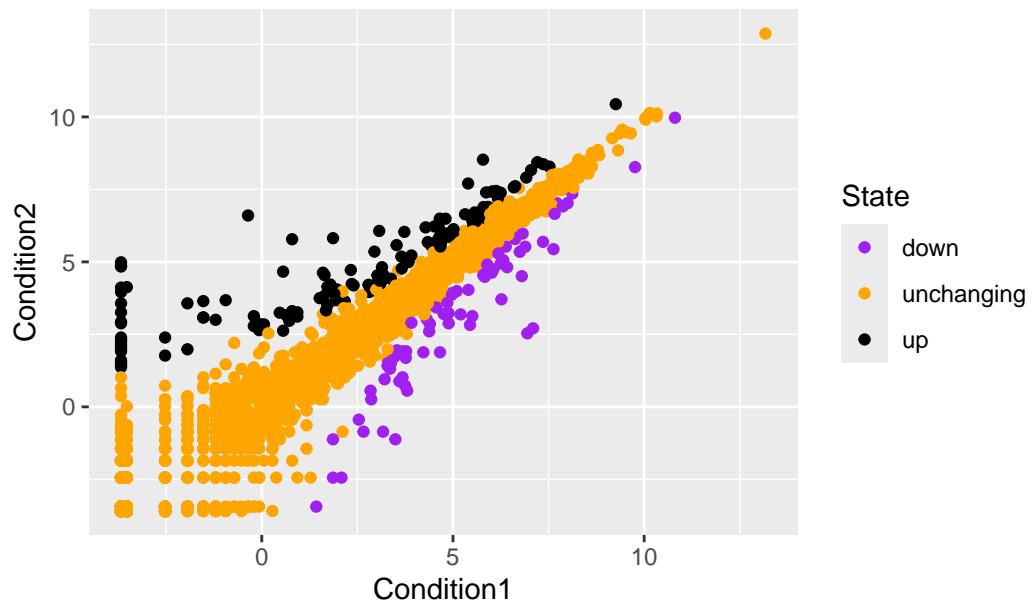
	down	unchanging	up
	72	4997	127

```
p<-ggplot(genes) +  
  aes(x=Condition1, y=Condition2, col=State) +  
  geom_point() +  
  scale_color_manual(values=c("purple","orange","black"))  
p
```



```
p + labs(title="look at me")
```

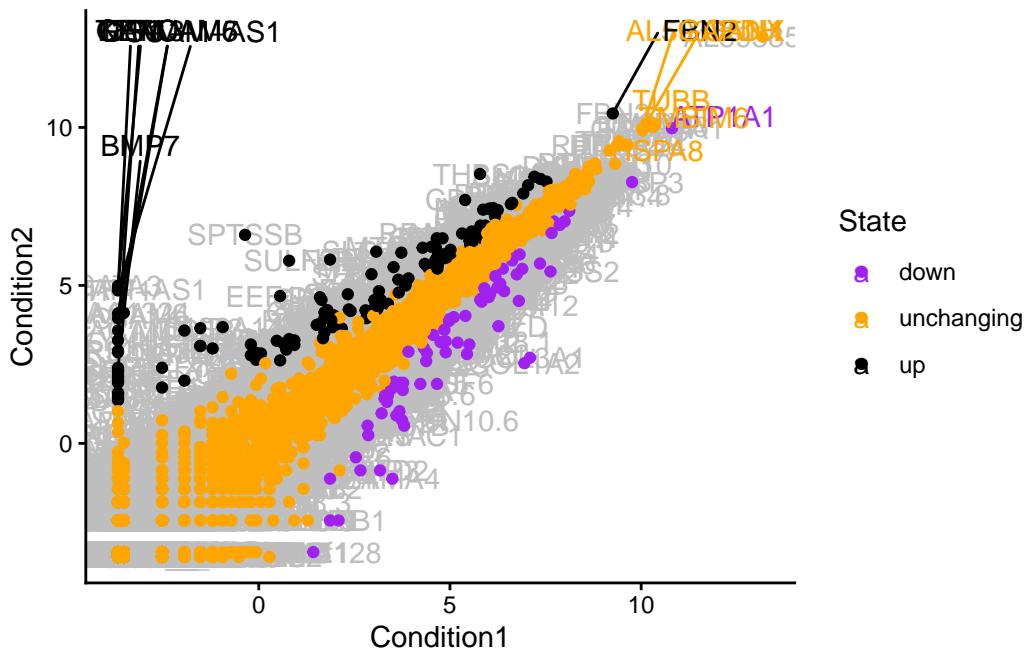
look at me



Silly example of adding labels

```
library(ggrepel)
pg<-ggplot(genes) +
  aes(x=Condition1, y=Condition2, col=State, label=Gene) +
  geom_text(color="gray") +
  geom_point() +
  scale_color_manual(values=c("purple","orange","black")) +
  geom_text_repel(max.overlaps=100) +
  theme_classic()
pg
```

Warning: ggrepel: 5181 unlabeled data points (too many overlaps). Consider increasing max.overlaps



## Going Further

Playing with different dataset to make something animated

```
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

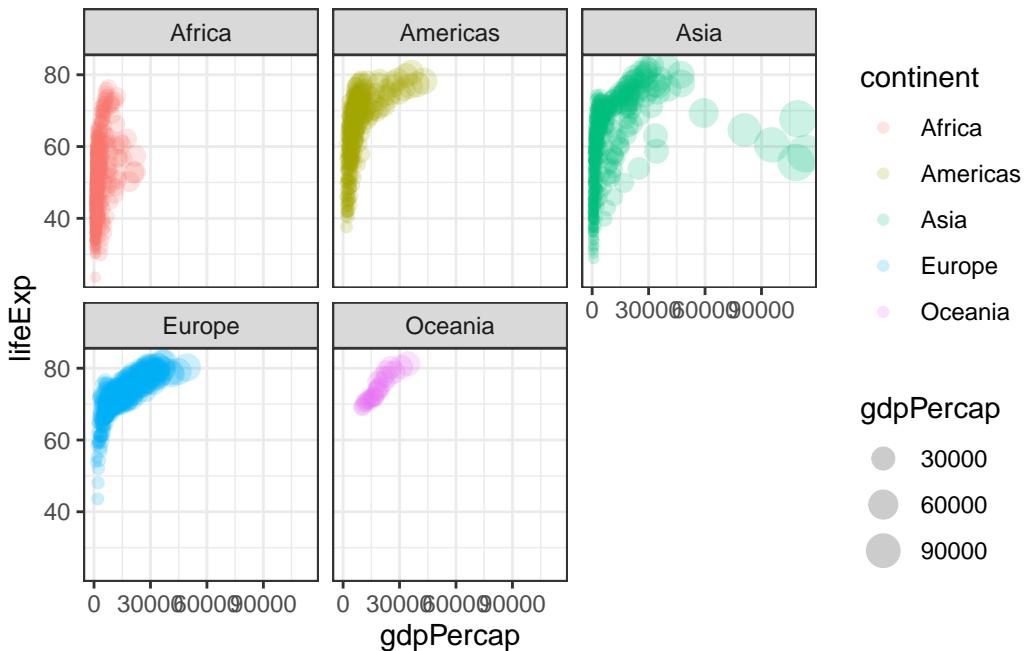
```

Classic plot of increasing GDP overtime.

```

ggplot(gapminder) +
  aes(x=gdpPercap, y=lifeExp, col=continent, size=gdpPercap) +
  geom_point(alpha=0.2) +
  facet_wrap(~continent) +
  theme_bw()

```

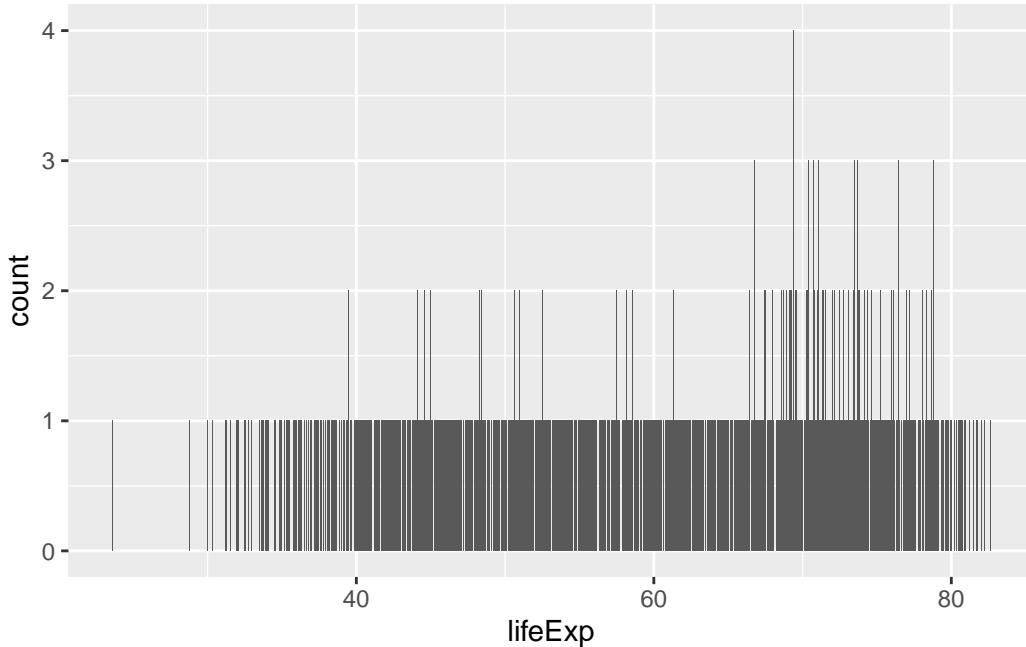


Wanted to try spitting more bars... couldn't

```

ggplot(gapminder) +
  aes(x=lifeExp) +
  geom_bar()

```



### Bonus gganimate code that never rendered lol

```
library(gapminder) library(gganimate)
```

### Setup nice regular ggplot of the gapminder data

```
ggplot(gapminder, aes(gdpPercap, lifeExp, size = pop, colour = country)) + geom_point(alpha = 0.7, show.legend = FALSE) + scale_colour_manual(values = country_colors) + scale_size(range = c(2, 12)) + scale_x_log10() + # Facet by continent facet_wrap(~continent) + # Here comes the gganimate specific bits labs(title = 'Year: {frame_time}', x = 'GDP per capita', y = 'life expectancy') + transition_time(year) + shadow_wake(wake_length = 0.1, alpha = FALSE) ""
```