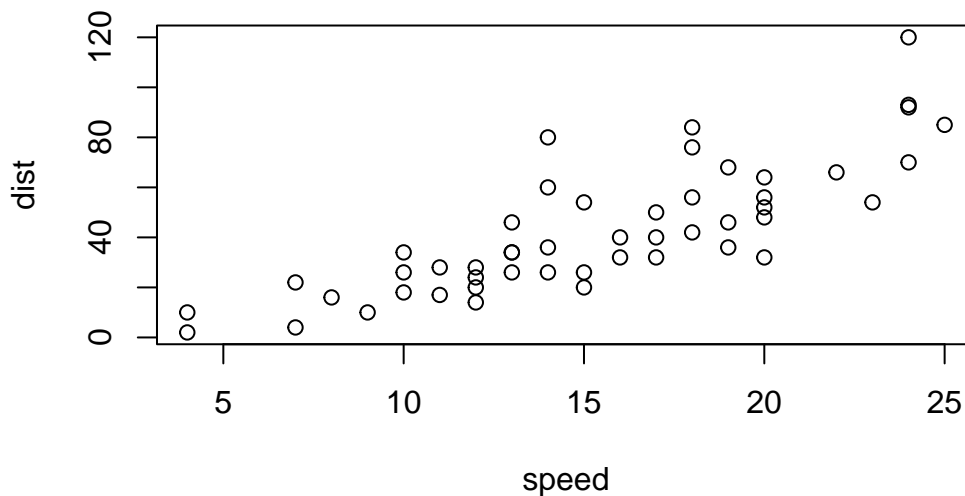


Class 5: Data Viz with ggplot

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R has many ways of plotting whatever crappy data you can throw at it. One that comes built-in is called **Base R** - the `plot()` function.

```
plot(cars)
```

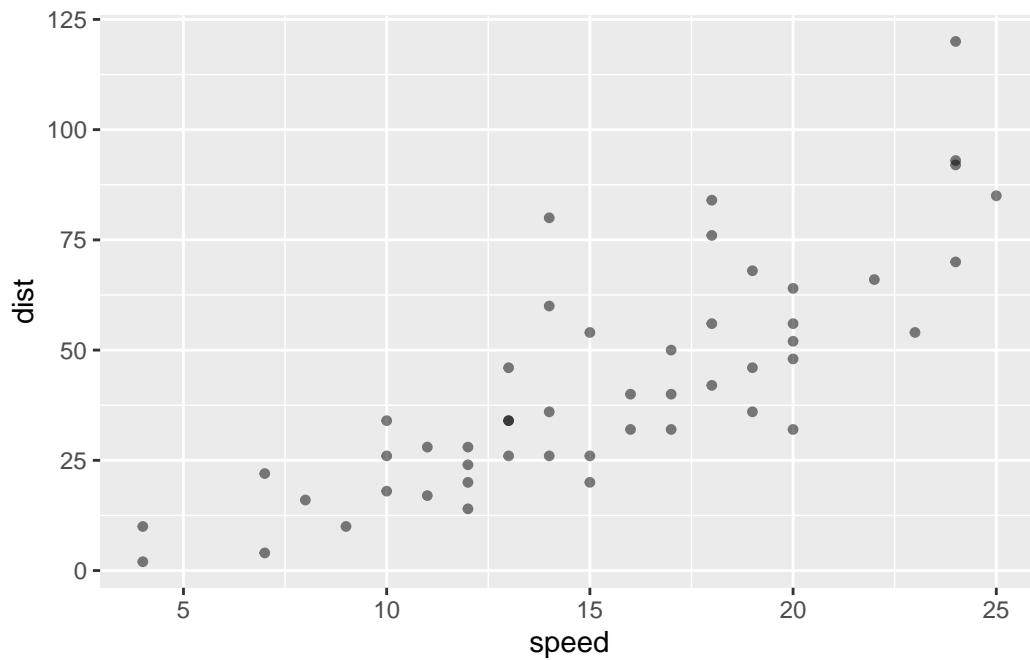


A very popular package in this particular avenue (haw!) is called **ggplot2**

Before using any add-on package it must be installed, you chowderhead! Using the `install.packages("ggplot2")` command/function. You can put this into the code, but it will reinstall it every time! So run it down in the console!

Then to use the package, it must be loaded up good with a `library(ggplot2)` call.

```
library(ggplot2)
ggplot(cars) + aes(x=speed, y=dist) + geom_point(shape = 20, size = 2, alpha = 0.5)
```

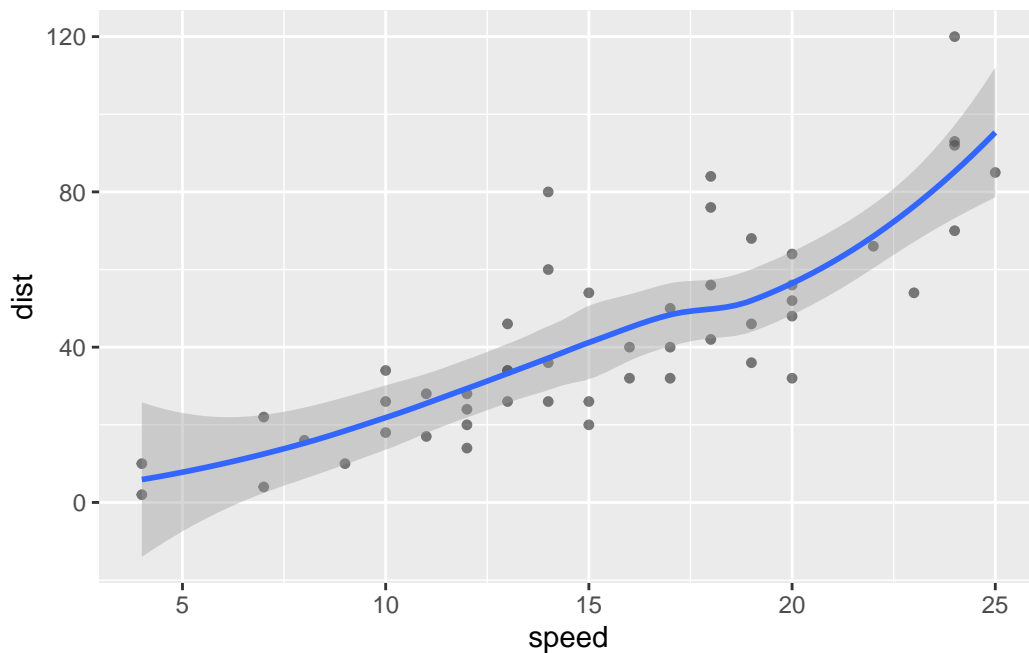


For “simple” plots like this one, base R code is fine, and in fact shorter, which in this case is great.

So to fit a model and show it on plot!

```
library(ggplot2)
ggplot(cars) + aes(x=speed, y=dist) + geom_point(shape = 20, size = 2, alpha = 0.5) + geom_smooth()
```

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



Every ggplot has at least 3 layers

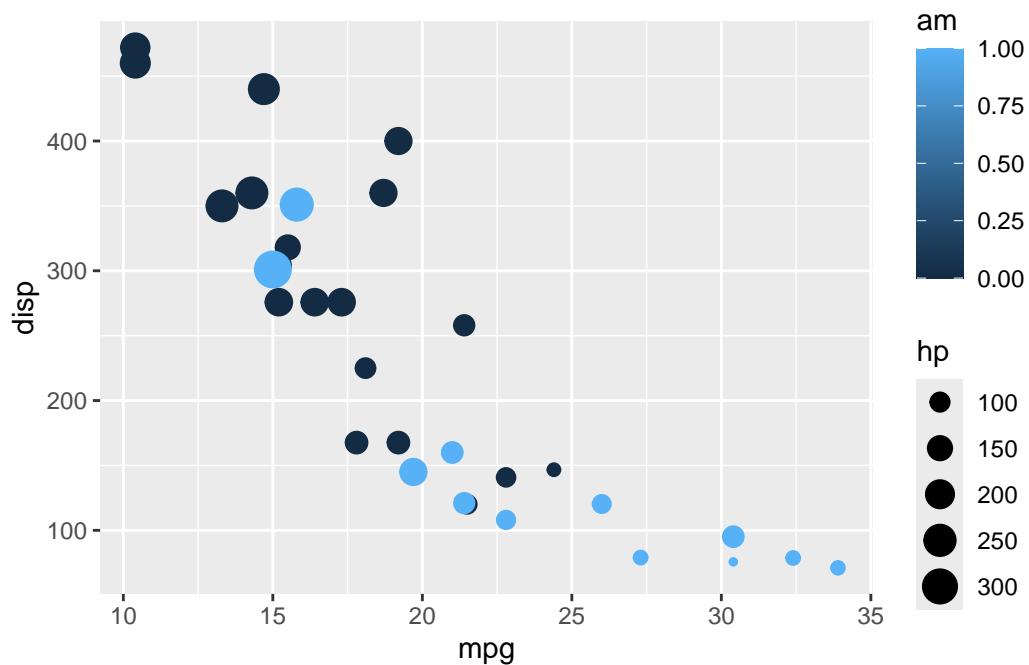
-**data** (data.frame with the numbers etc that you may possibly wish to plot) -**aesthetics** (mapping of your data columns to your plot) - **geoms**; there's a whole lot of these, (with basic types like `geom_point()`, `geom_line()`, `geom_col()`)

`mtcars`

	mpg	cyl	disp	hp	drat	wt	qsec	vs	am	gear	carb
Mazda RX4	21.0	6	160.0	110	3.90	2.620	16.46	0	1	4	4
Mazda RX4 Wag	21.0	6	160.0	110	3.90	2.875	17.02	0	1	4	4
Datsun 710	22.8	4	108.0	93	3.85	2.320	18.61	1	1	4	1
Hornet 4 Drive	21.4	6	258.0	110	3.08	3.215	19.44	1	0	3	1
Hornet Sportabout	18.7	8	360.0	175	3.15	3.440	17.02	0	0	3	2
Valiant	18.1	6	225.0	105	2.76	3.460	20.22	1	0	3	1
Duster 360	14.3	8	360.0	245	3.21	3.570	15.84	0	0	3	4
Merc 240D	24.4	4	146.7	62	3.69	3.190	20.00	1	0	4	2
Merc 230	22.8	4	140.8	95	3.92	3.150	22.90	1	0	4	2
Merc 280	19.2	6	167.6	123	3.92	3.440	18.30	1	0	4	4
Merc 280C	17.8	6	167.6	123	3.92	3.440	18.90	1	0	4	4
Merc 450SE	16.4	8	275.8	180	3.07	4.070	17.40	0	0	3	3
Merc 450SL	17.3	8	275.8	180	3.07	3.730	17.60	0	0	3	3
Merc 450SLC	15.2	8	275.8	180	3.07	3.780	18.00	0	0	3	3
Cadillac Fleetwood	10.4	8	472.0	205	2.93	5.250	17.98	0	0	3	4

Lincoln Continental	10.4	8	460.0	215	3.00	5.424	17.82	0	0	3	4
Chrysler Imperial	14.7	8	440.0	230	3.23	5.345	17.42	0	0	3	4
Fiat 128	32.4	4	78.7	66	4.08	2.200	19.47	1	1	4	1
Honda Civic	30.4	4	75.7	52	4.93	1.615	18.52	1	1	4	2
Toyota Corolla	33.9	4	71.1	65	4.22	1.835	19.90	1	1	4	1
Toyota Corona	21.5	4	120.1	97	3.70	2.465	20.01	1	0	3	1
Dodge Challenger	15.5	8	318.0	150	2.76	3.520	16.87	0	0	3	2
AMC Javelin	15.2	8	304.0	150	3.15	3.435	17.30	0	0	3	2
Camaro Z28	13.3	8	350.0	245	3.73	3.840	15.41	0	0	3	4
Pontiac Firebird	19.2	8	400.0	175	3.08	3.845	17.05	0	0	3	2
Fiat X1-9	27.3	4	79.0	66	4.08	1.935	18.90	1	1	4	1
Porsche 914-2	26.0	4	120.3	91	4.43	2.140	16.70	0	1	5	2
Lotus Europa	30.4	4	95.1	113	3.77	1.513	16.90	1	1	5	2
Ford Pantera L	15.8	8	351.0	264	4.22	3.170	14.50	0	1	5	4
Ferrari Dino	19.7	6	145.0	175	3.62	2.770	15.50	0	1	5	6
Maserati Bora	15.0	8	301.0	335	3.54	3.570	14.60	0	1	5	8
Volvo 142E	21.4	4	121.0	109	4.11	2.780	18.60	1	1	4	2

```
library(ggplot2)
ggplot(mtcars) + aes(x=mpg, y=disp, size = hp, color = am) + geom_point()
```



```
library(ggplot2)
ggplot(mtcars) + aes(x=mpg, y=disp, size = hp, color = am, label = rownames(mtcars)) + geom
```

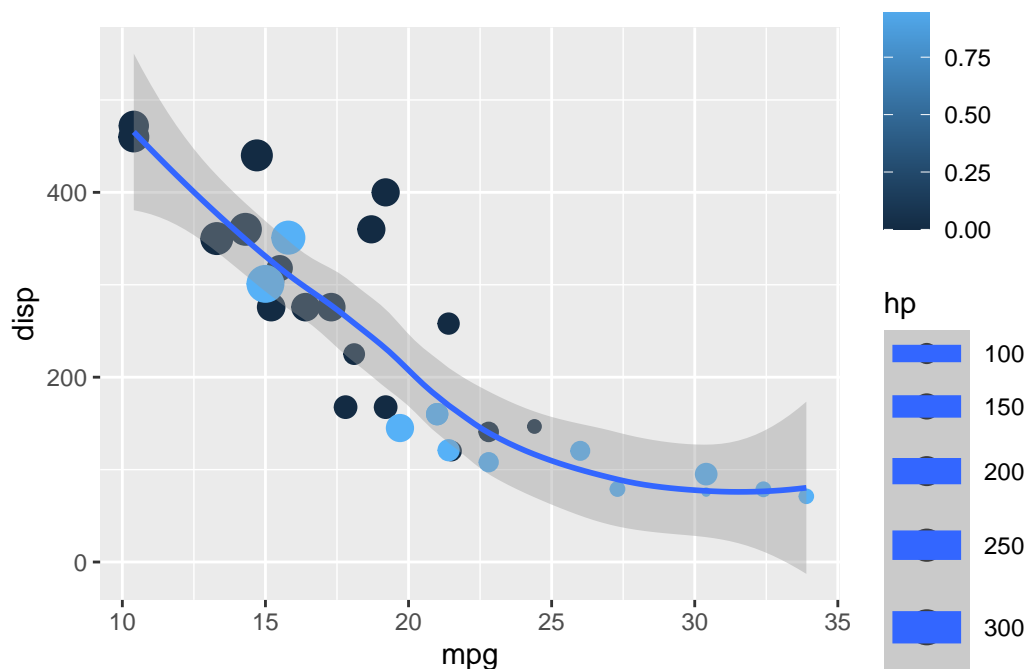
Warning: Using `size` aesthetic for lines was deprecated in ggplot2 3.4.0.
i Please use `linewidth` instead.

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

Warning: The following aesthetics were dropped during statistical transformation: size, colour, and label.

i This can happen when ggplot fails to infer the correct grouping structure in the data.

i Did you forget to specify a `group` aesthetic or to convert a numerical variable into a factor?



```
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

```
nrow(genes)
```

```
[1] 5196
```

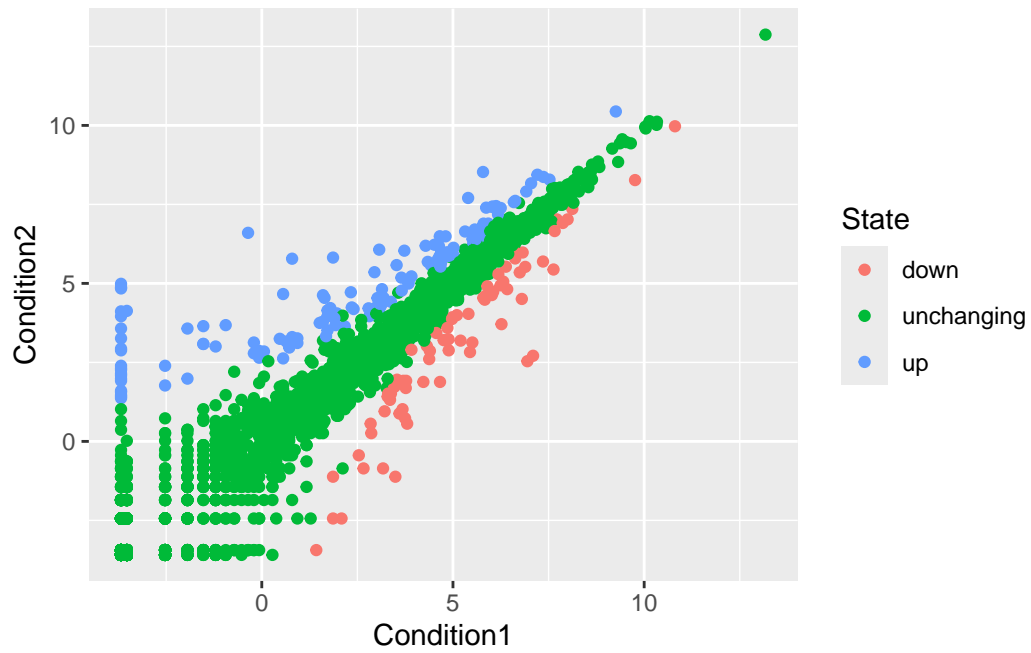
```
ncol(genes)
```

```
[1] 4
```

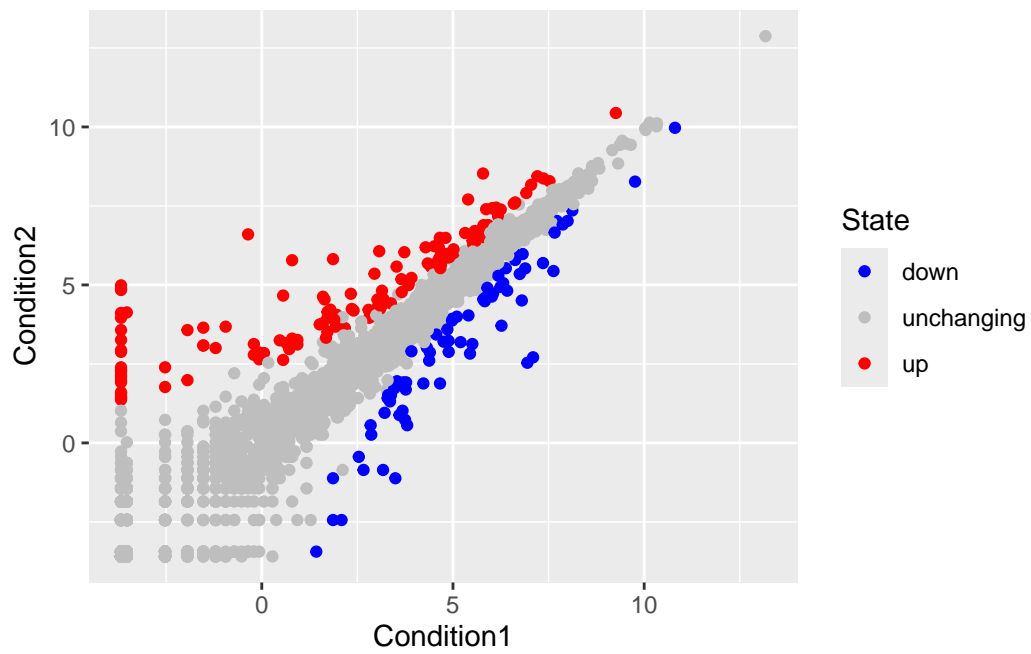
```
round(table(genes$State) / nrow(genes), 2)
```

down	unchanging	up
0.01	0.96	0.02

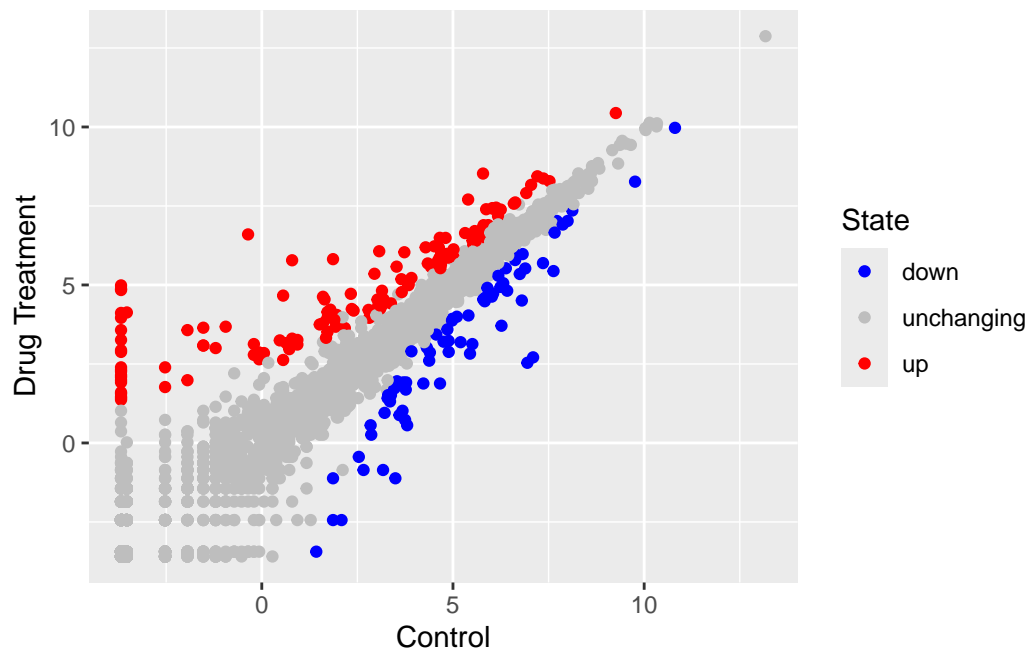
```
p <- ggplot(genes) + aes(Condition1, Condition2, color = State) + geom_point()
p
```



```
p + scale_color_manual( values = c("blue", "gray", "red"))
```



```
p + scale_color_manual( values = c("blue", "gray", "red")) + labs(x = "Control", y = "Drug Treatment")
```



```
ggsave("druggeneratedblue.pdf")
```

Saving 5.5 x 3.5 in image

```
url <- "https://raw.githubusercontent.com/jennybc/gapminder/master/inst/extdata/gapminder.tsv"
gapminder <- read.delim(url)
library(dplyr)
```

Attaching package: 'dplyr'

The following objects are masked from 'package:stats':

filter, lag

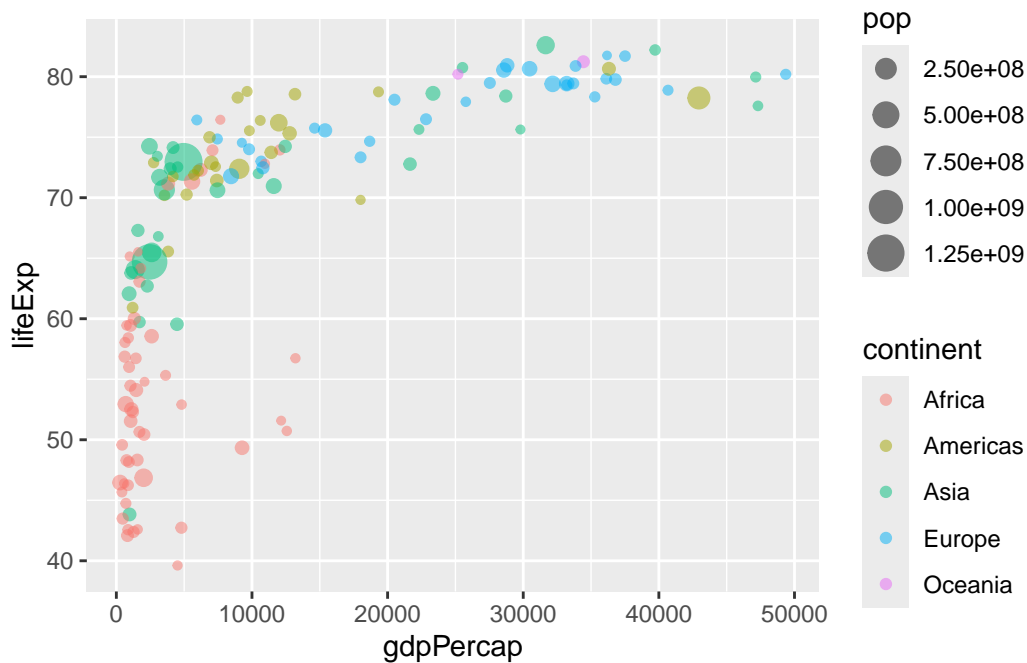
The following objects are masked from 'package:base':

intersect, setdiff, setequal, union

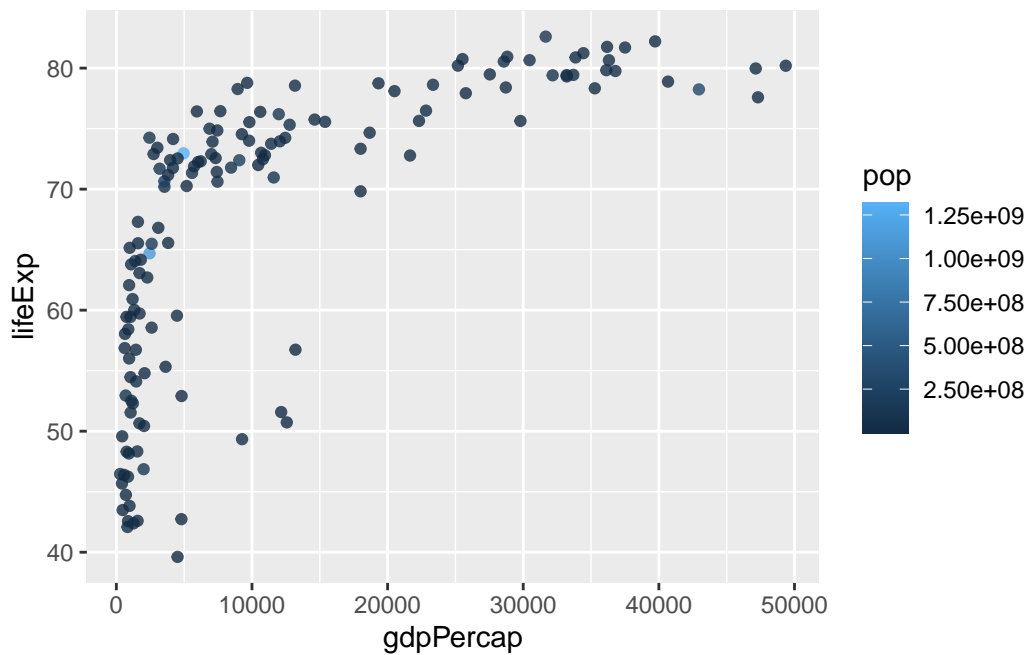

```
gapminder_2007 <- gapminder %>% filter(year==2007)
head(gapminder_2007)
```

	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

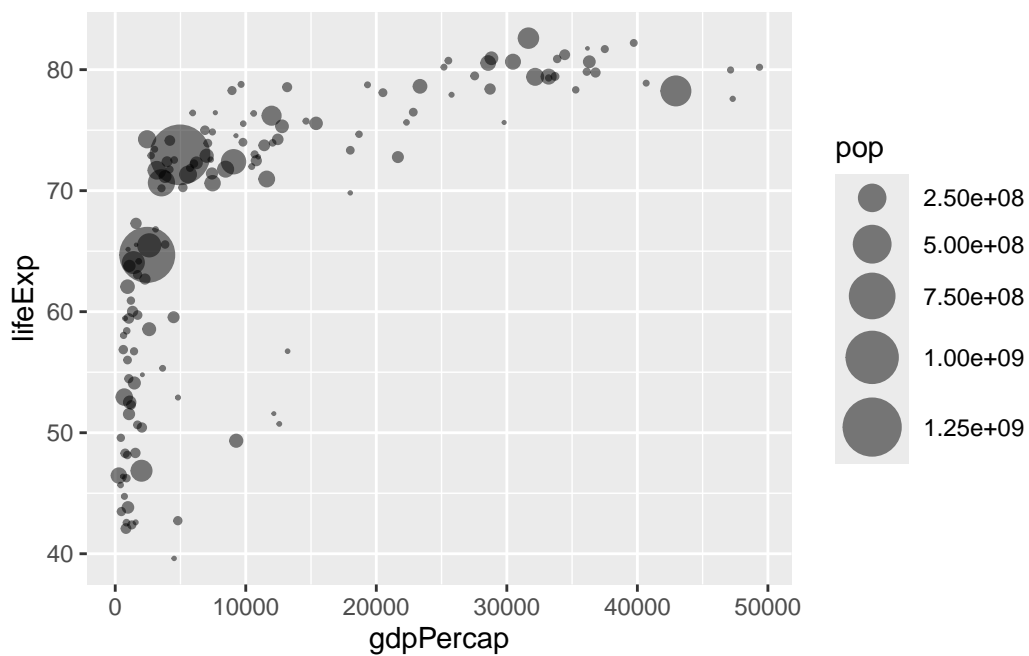
```
ggplot(gapminder_2007) + aes(x= gdpPercap, y= lifeExp, color = continent, size = pop) + geom.
```



```
ggplot(gapminder_2007) + aes(x= gdpPercap, y= lifeExp, color = pop) + geom_point(alpha = 0.8)
```

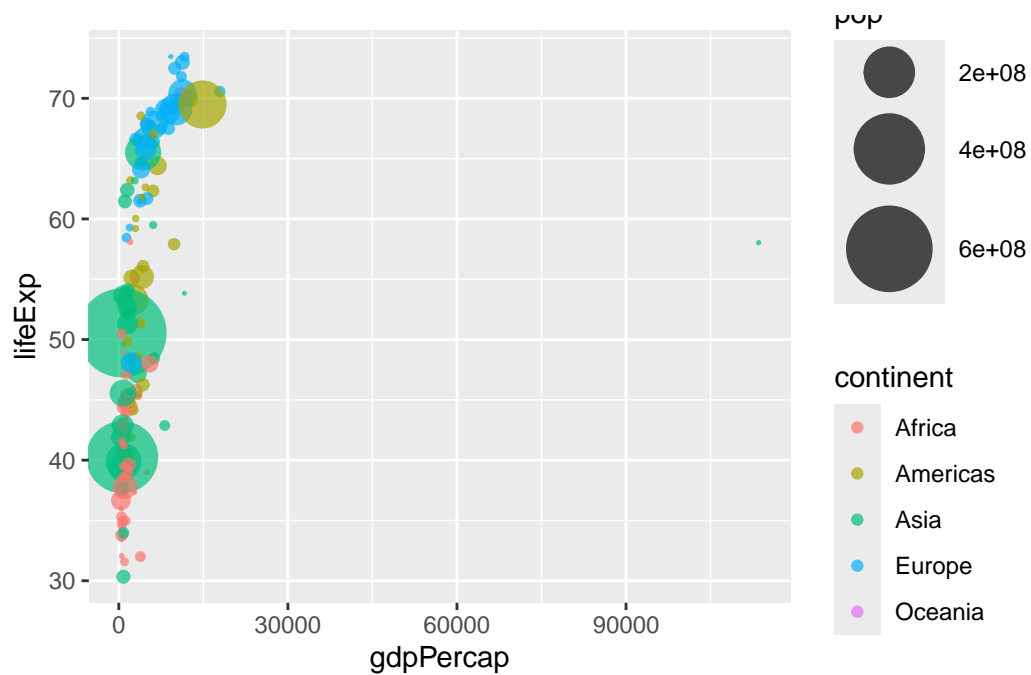


```
ggplot(gapminder_2007) + aes(x= gdpPercap, y= lifeExp, size = pop) + geom_point(alpha = 0.5)
```

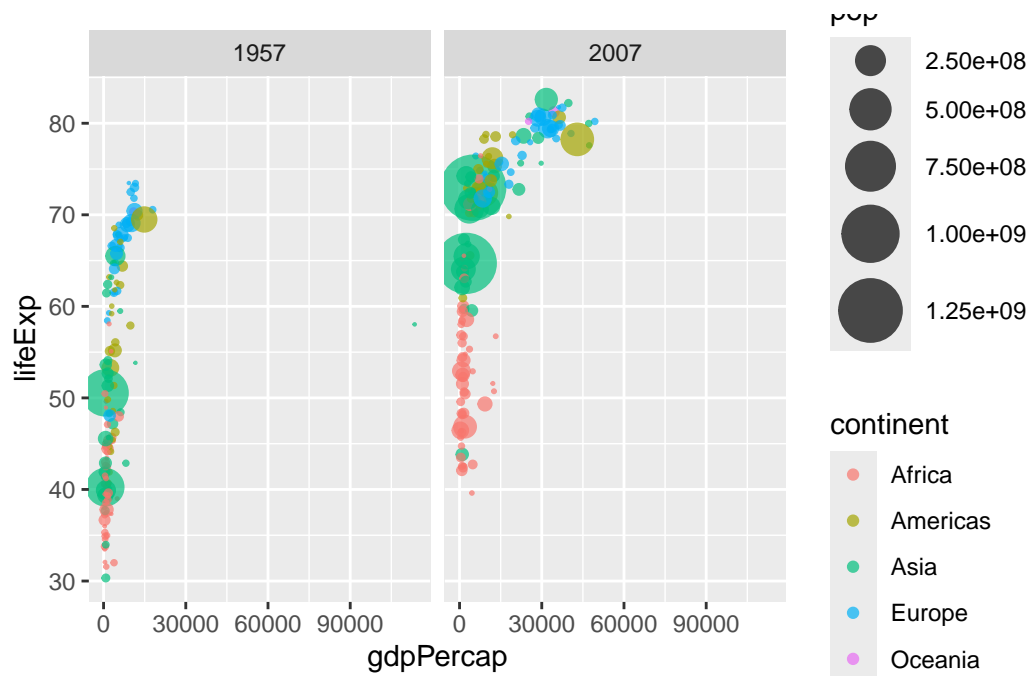


The lab text is wrong here, places alpha outside central parentheticals.

```
gapminder_1957 <- gapminder %>% filter(year==1957)
ggplot(gapminder_1957) + aes(x= gdpPercap, y= lifeExp, size = pop, color = continent) + geom.
```



```
nottwodiffeentyyearsforunclearreasons <- gapminder %>% filter(year==1957 | year==2007)
ggplot(nottwodiffeentyyearsforunclearreasons) + aes(x= gdpPercap, y= lifeExp, size = pop, col
```

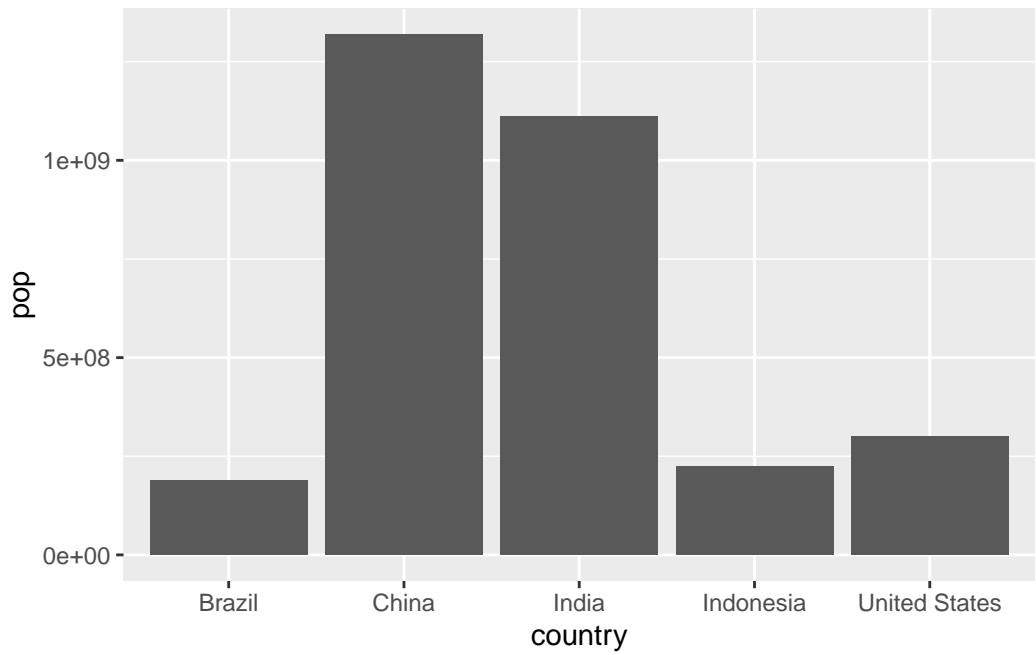


```
gapminder_top5 <- gapminder %>%
  filter(year==2007) %>%
  arrange(desc(pop)) %>%
  top_n(5, pop)

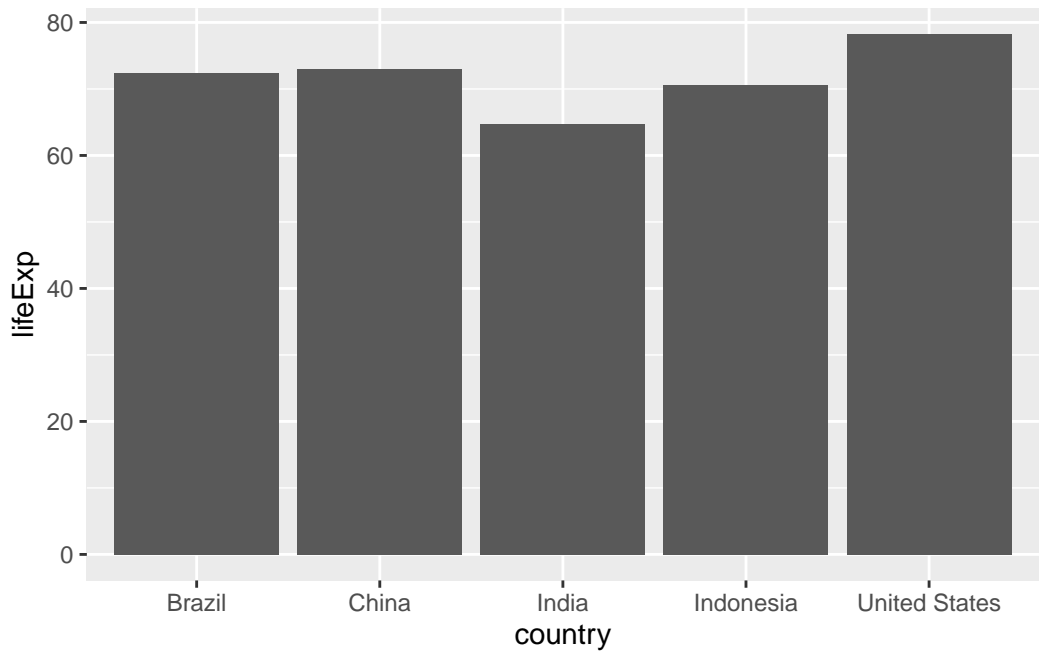
gapminder_top5
```

	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

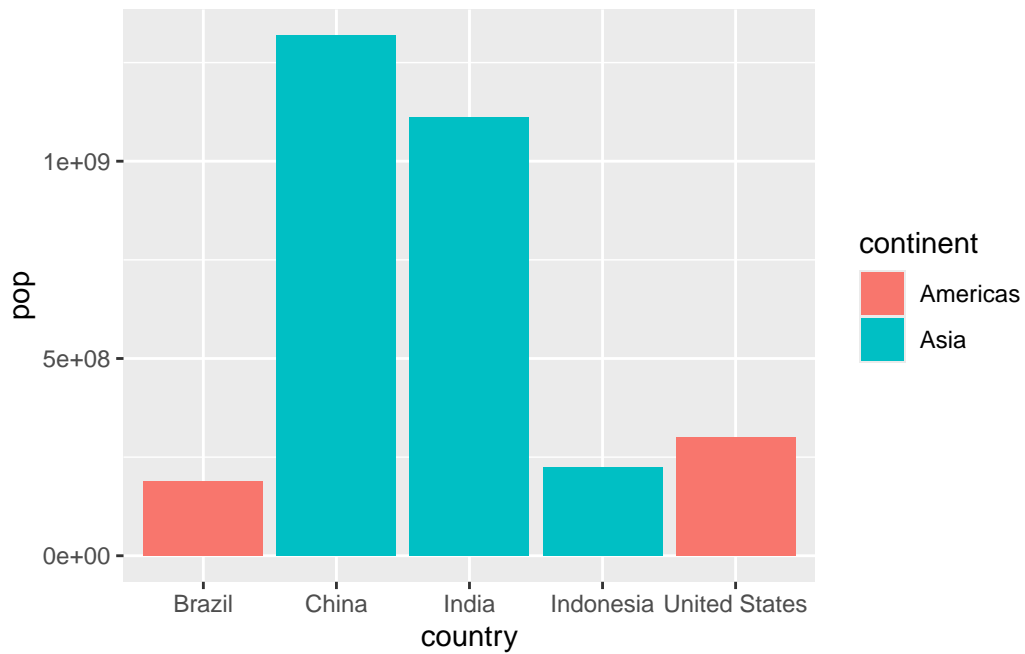
```
ggplot(gapminder_top5) + geom_col(aes(x= country, y = pop))
```



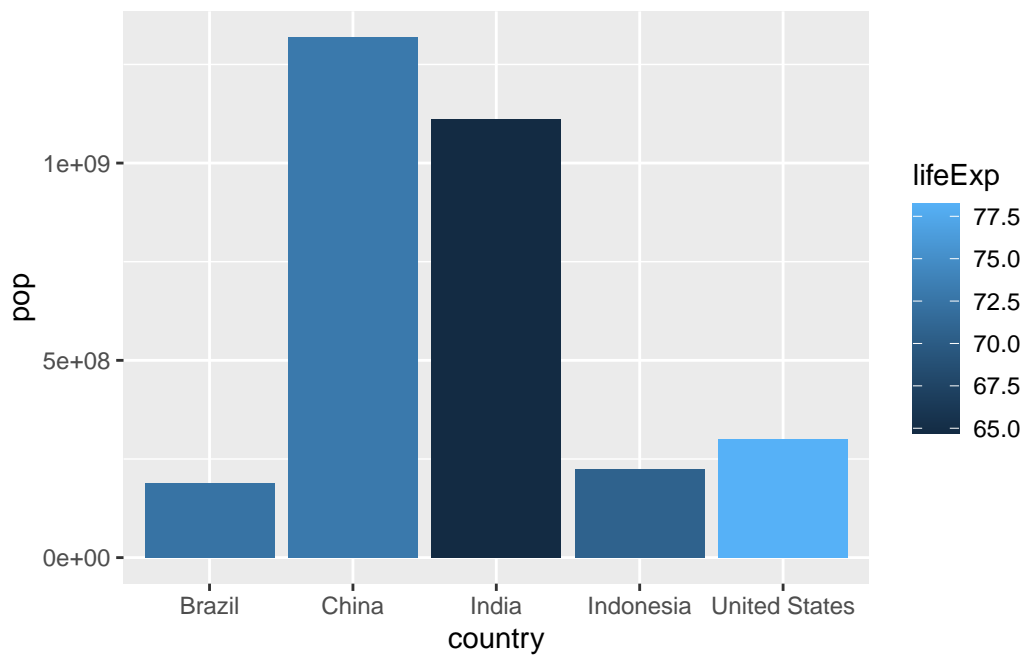
```
ggplot(gapminder_top5) + geom_col(aes(x= country, y = lifeExp))
```



```
ggplot(gapminder_top5) + geom_col(aes(x= country, y = pop, fill = continent))
```



```
ggplot(gapminder_top5) + geom_col(aes(x= country, y = pop, fill = lifeExp))
```



```
ggplot(gapminder_top5) + aes(x=reorder(country, -pop), y = pop, fill = gdpPercap) + geom_col
```

