

Class 5: Data Visualization (ggplot)

AUTHOR

Jazz Zhang (A16149005)

Using GGPLOT

To use ggplot2 we first need to install it on our computers. To do this we will use the function

```
install.packages()
```

```
library(ggplot2)
```

Warning: package 'ggplot2' was built under R version 4.3.1

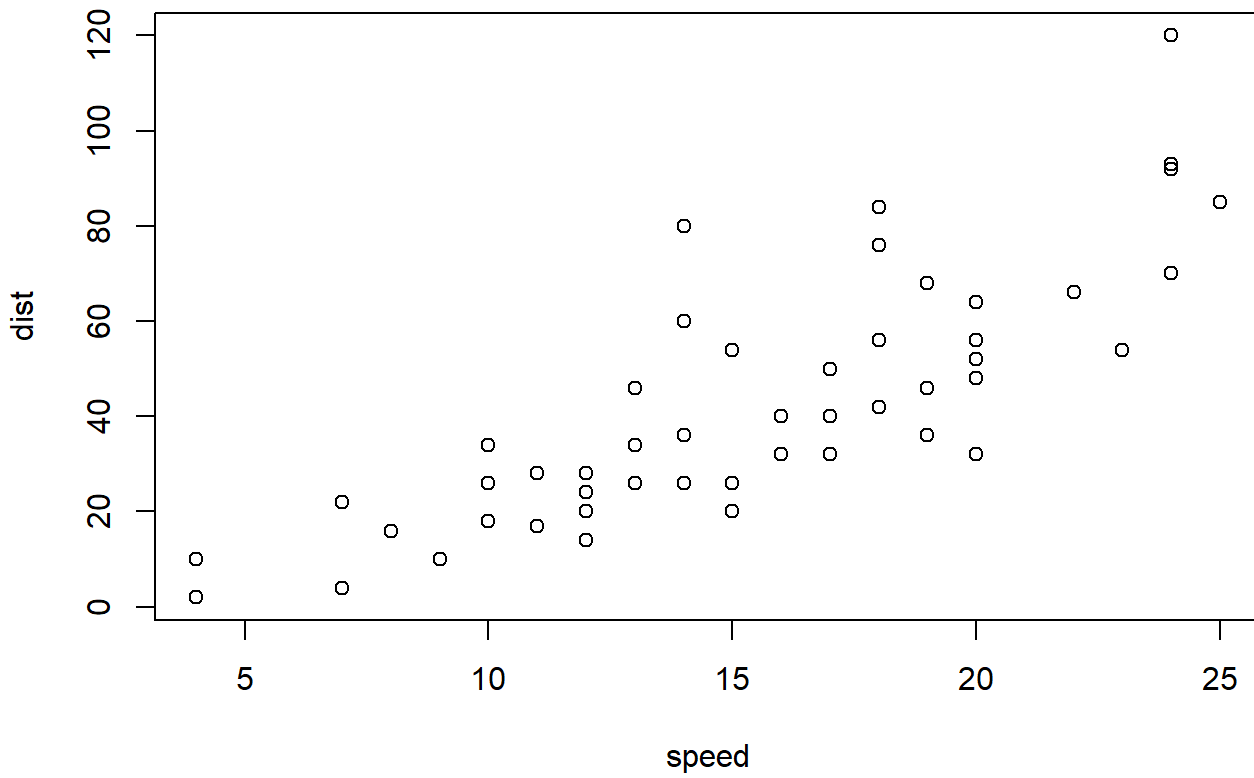
```
ggplot(cars)
```

```
head(cars)
```

```
  speed  dist  
1     4     2
```

2	4	10
3	7	4
4	7	22
5	8	16
6	9	10

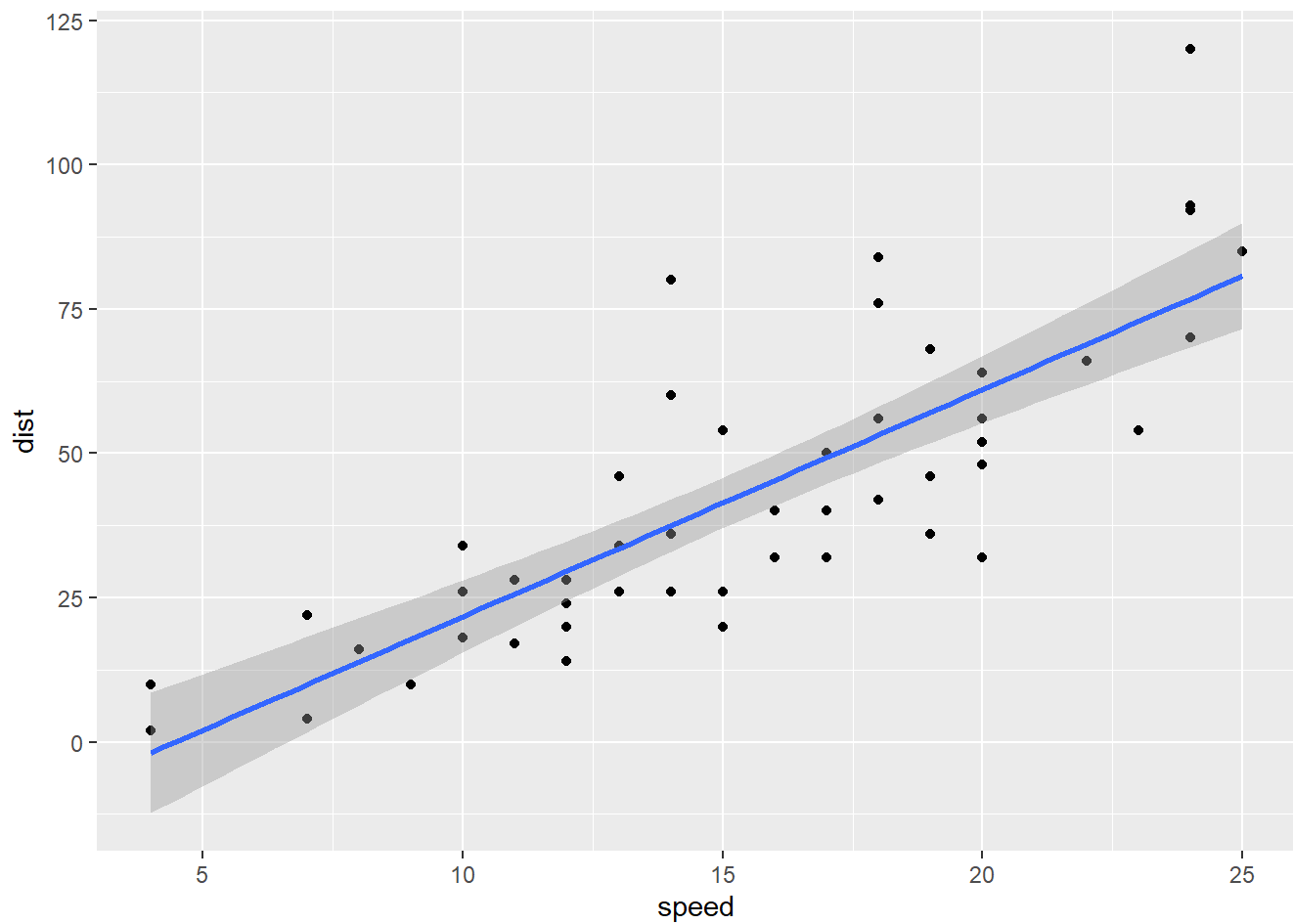
```
plot(cars)
```



ggplot: -data(data.frame for plotting) -aesthetics (aes() values - how to map data) -geom (type of plot)

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

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



```
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
```

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

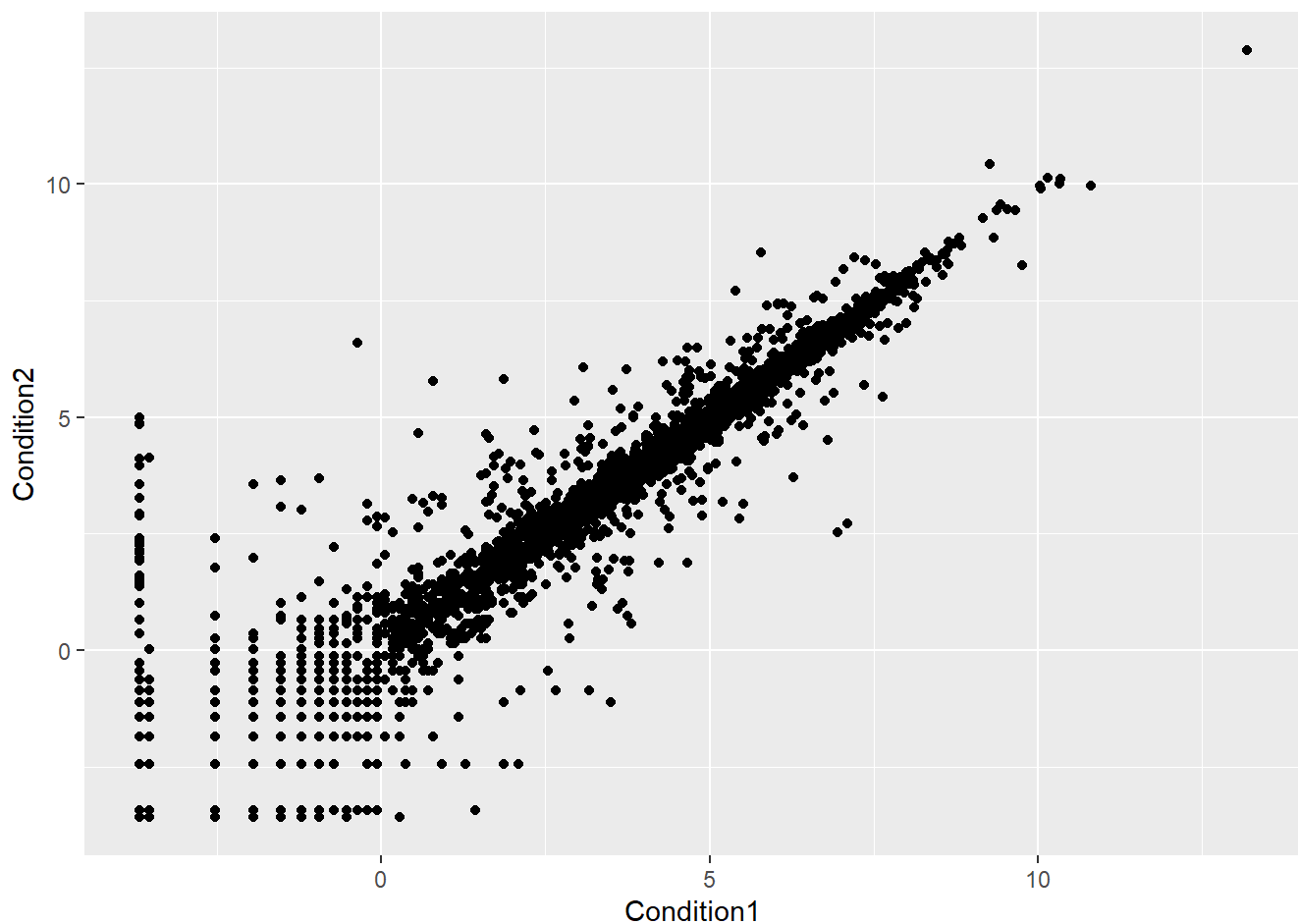
down	unchanging	up
72	4997	127

```
round( table(genes$State)/nrow(genes) * 100, 2 )
```

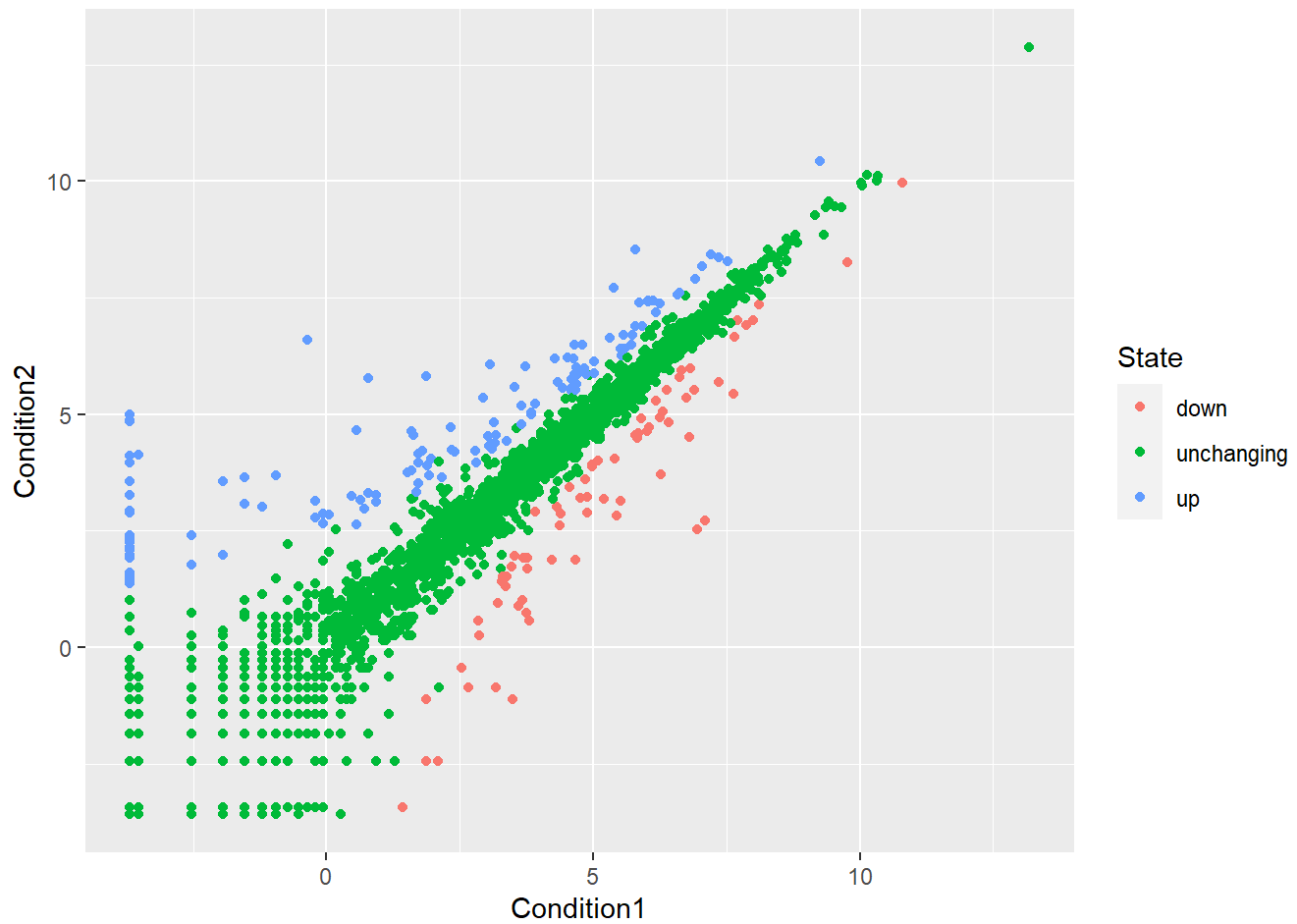
down	unchanging	up
1.39	96.17	2.44

5196 rows, 4 columns, 127 up-regulated genes, 2.44% up-regulated

```
ggplot(genes) +  
  aes(x=Condition1, y=Condition2) +  
  geom_point()
```

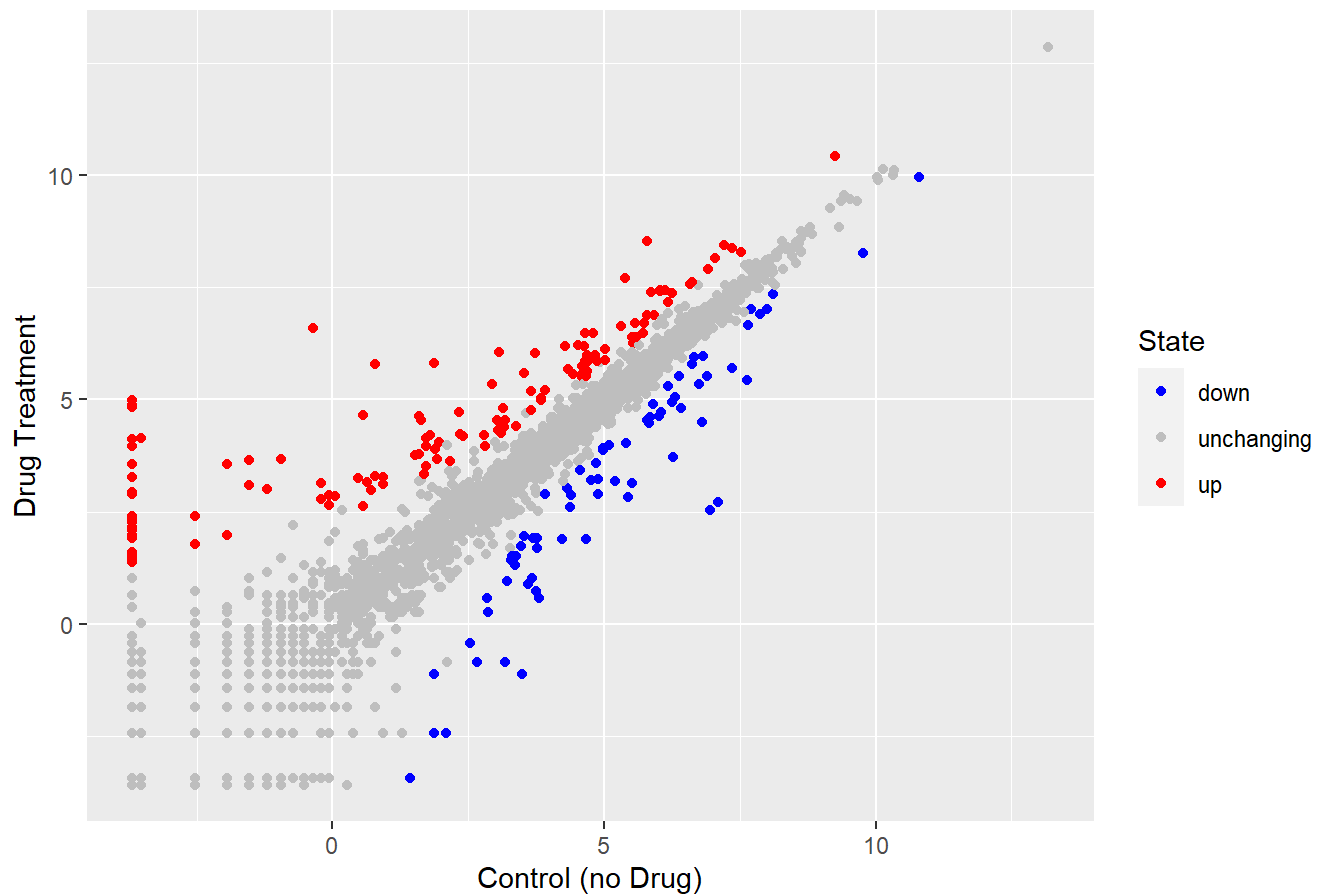


```
p <- ggplot(genes) +  
  aes(x=Condition1, y=Condition2, col=State) +  
  geom_point()  
p
```



```
p +  
  scale_colour_manual( values=c("blue","gray","red") )+  
  labs(title="Gene expression changes upon drug treatment", x="Control (no Drug)", y="Drug Treatment")
```

Gene expression changes upon drug treatment



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

library(dplyr)
```

Warning: package 'dplyr' was built under R version 4.3.1

Attaching package: 'dplyr'

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

filter, lag

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

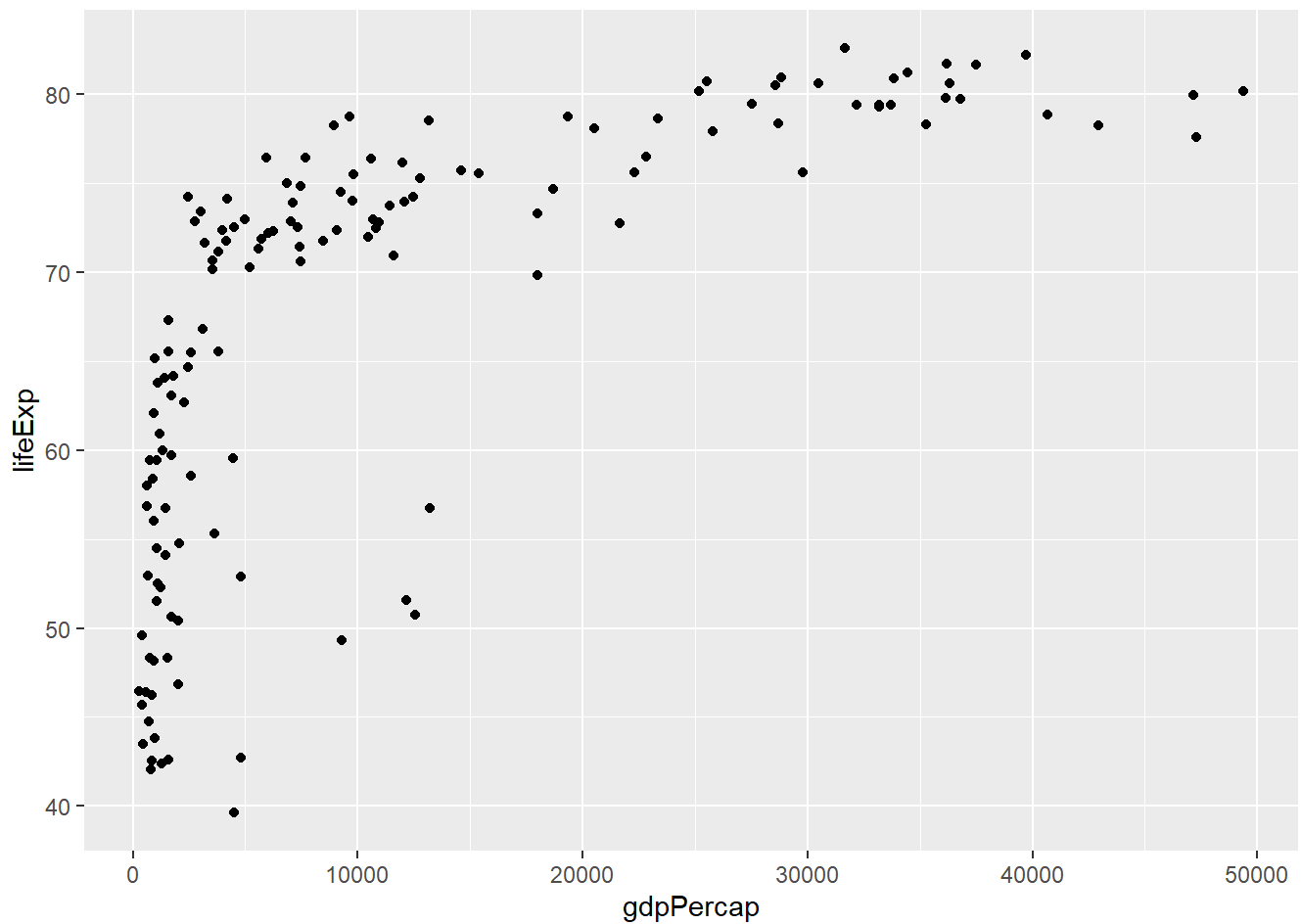
intersect, setdiff, setequal, union

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

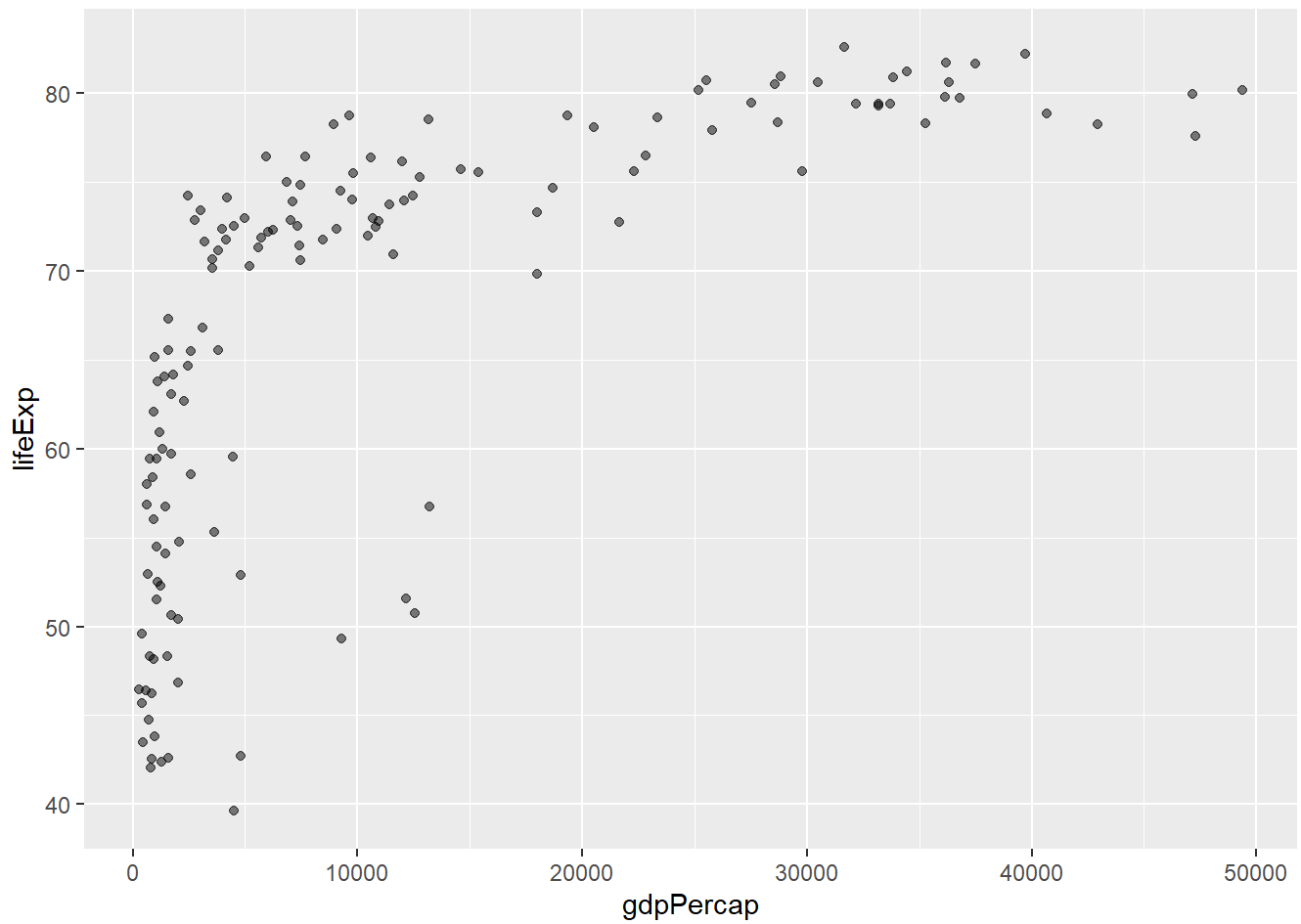
	country	continent	year	lifeExp	pop	gdpPercap
1	Afghanistan	Asia	2007	43.828	31889923	974.5803

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

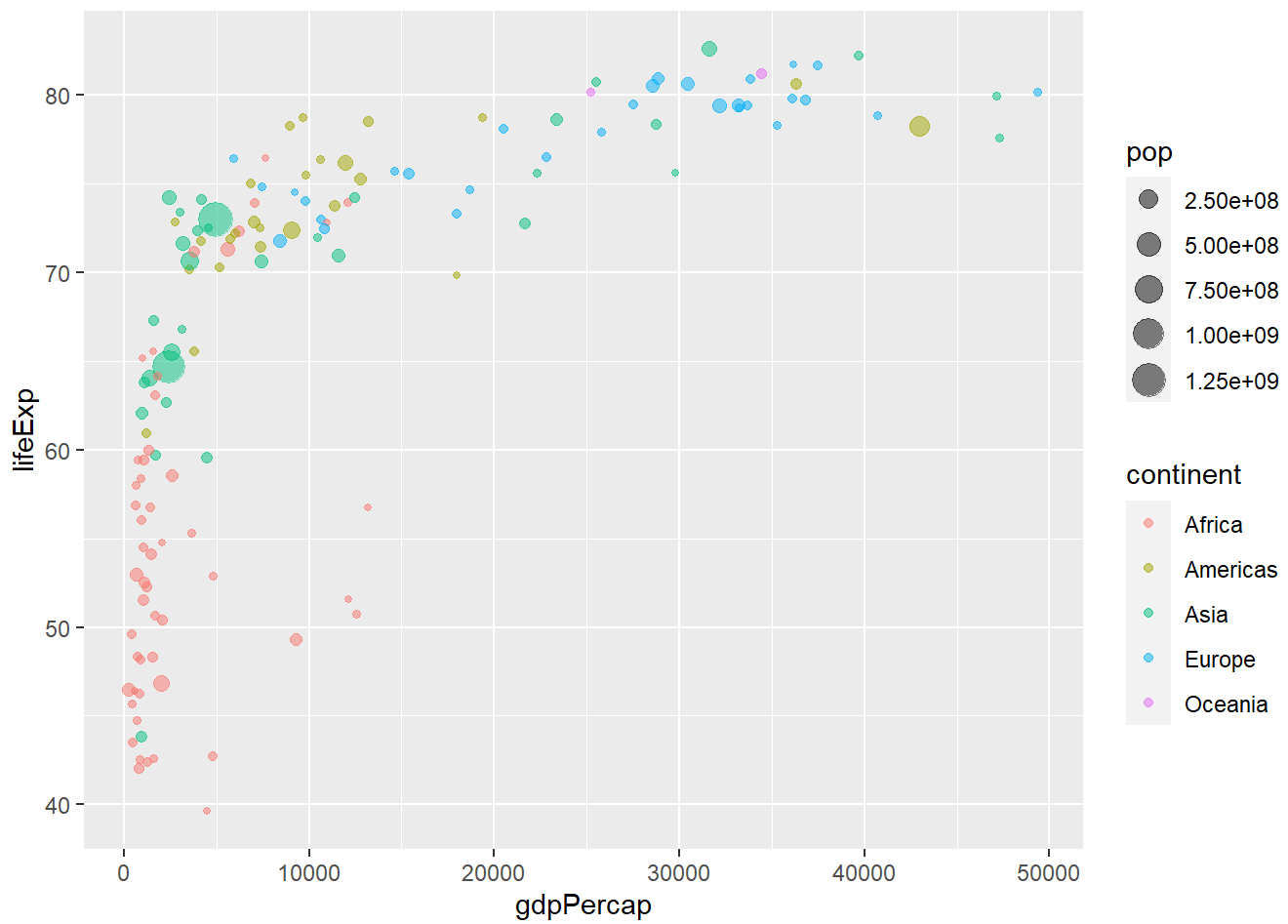
```
ggplot(gapminder_2007) +  
  aes(x=gdpPercap, y=lifeExp) +  
  geom_point()
```



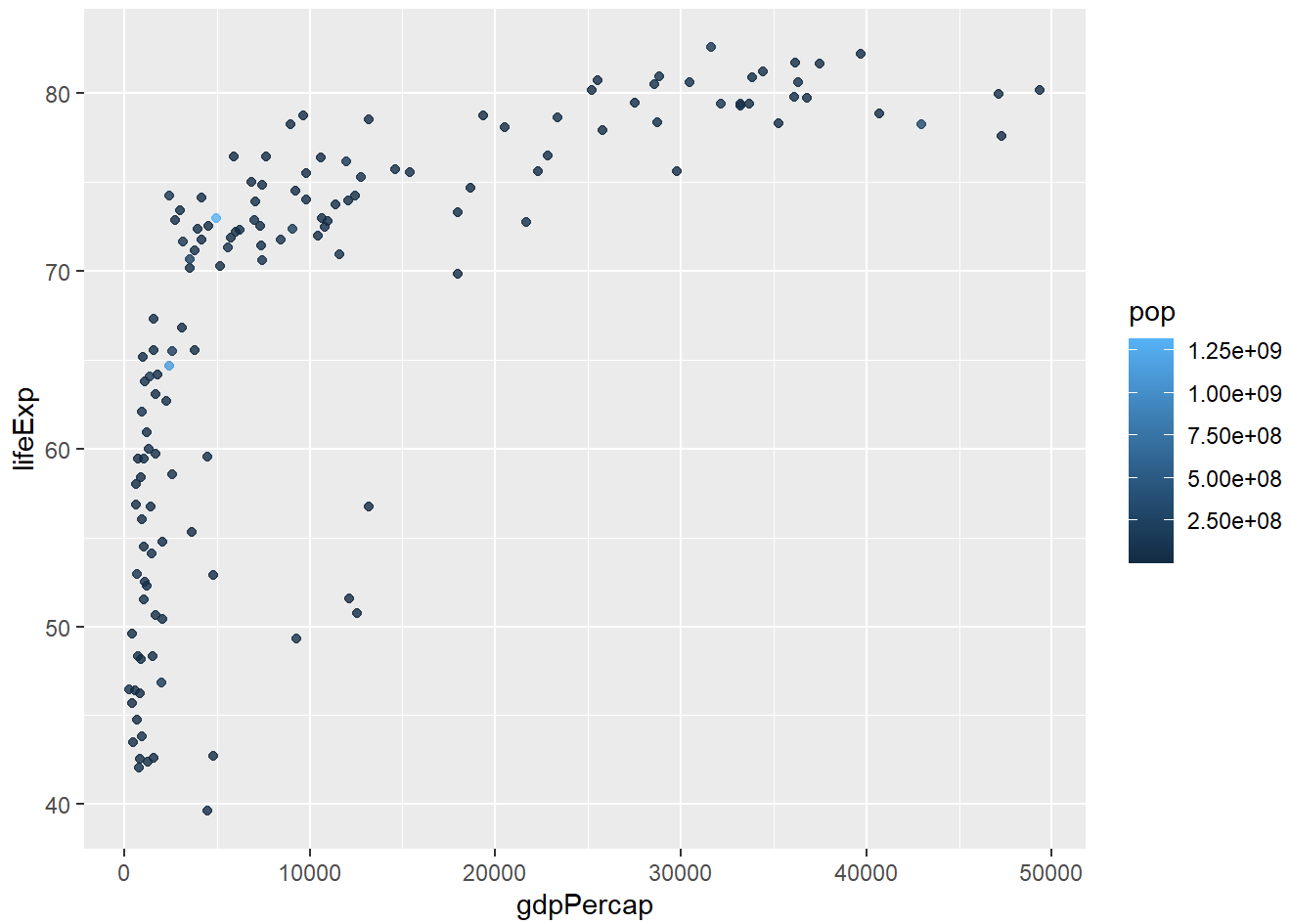
```
ggplot(gapminder_2007) +  
  aes(x=gdpPercap, y=lifeExp) +  
  geom_point(alpha=0.5)
```



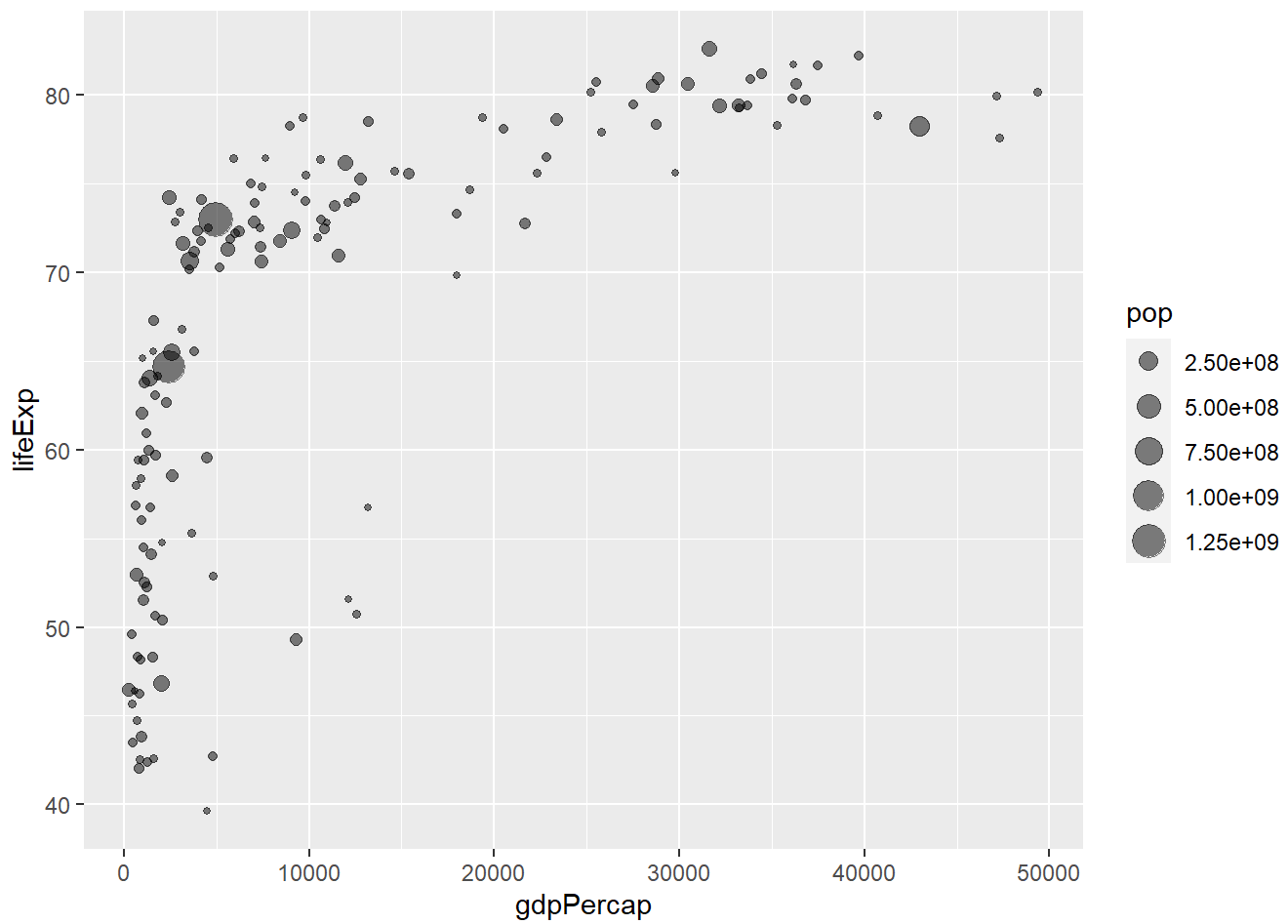
```
ggplot(gapminder_2007) +  
  aes(x=gdpPercap, y=lifeExp, color=continent, size=pop) +  
  geom_point(alpha=0.5)
```

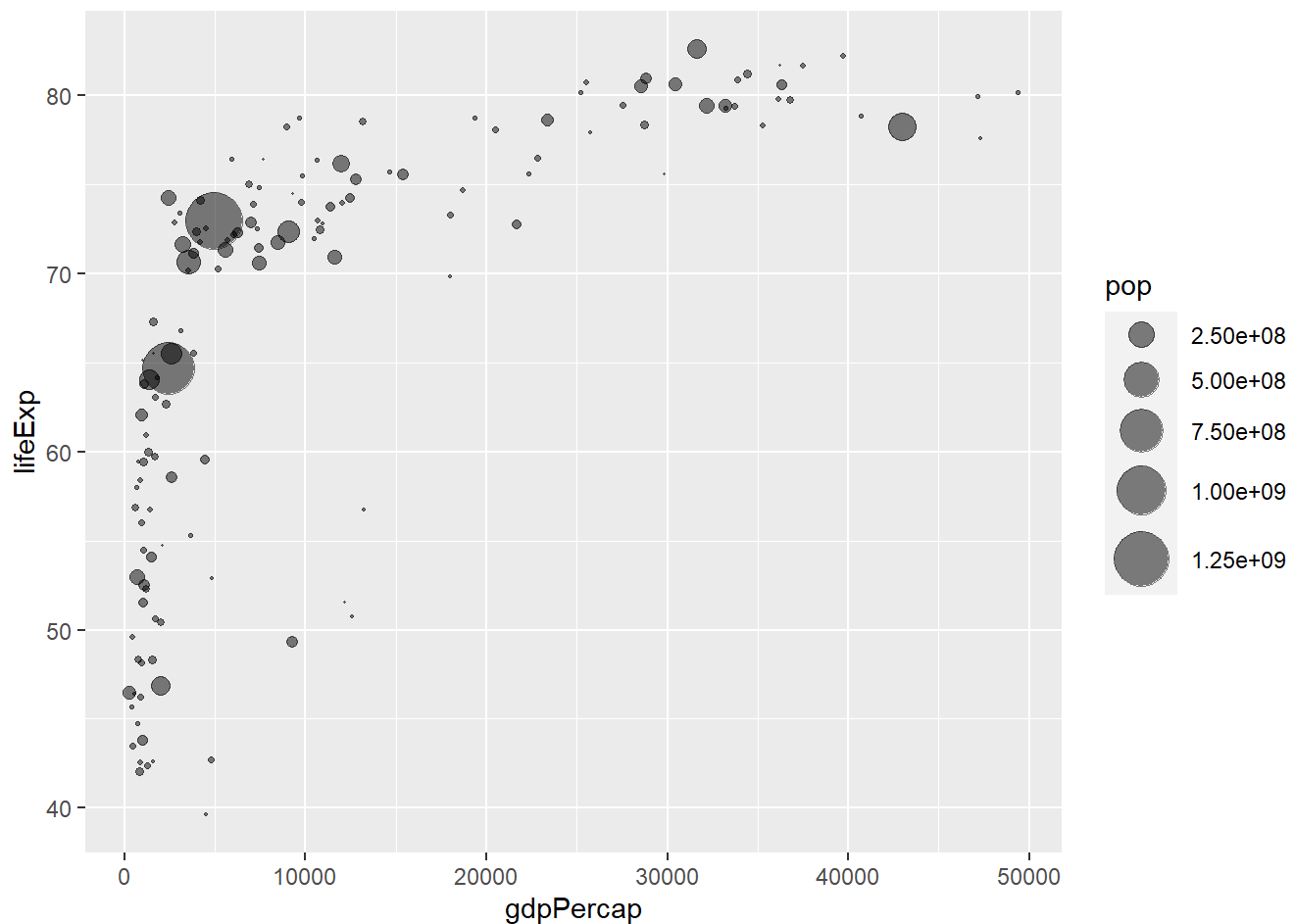
```
ggplot(gapminder_2007) +  
  aes(x = gdpPercap, y = lifeExp, color = pop) +  
  geom_point(alpha=0.8)
```



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



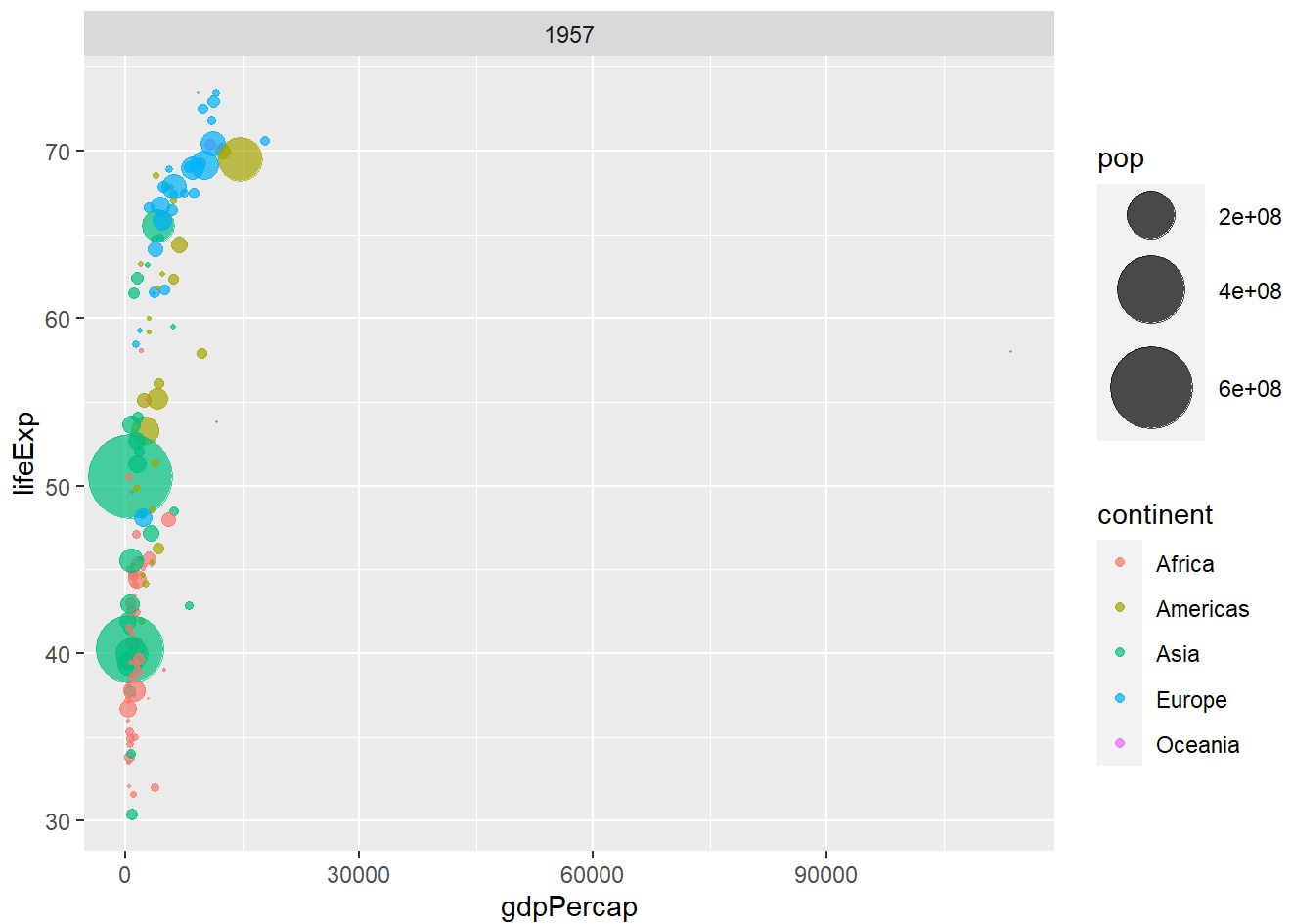
```
ggplot(gapminder_2007) +  
  geom_point(aes(x = gdpPercap, y = lifeExp,  
                 size = pop), alpha=0.5) +  
  scale_size_area(max_size = 10)
```



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

	country	continent	year	lifeExp	pop	gdpPercap
1	Afghanistan	Asia	1957	30.332	9240934	820.853
2	Albania	Europe	1957	59.280	1476505	1942.284
3	Algeria	Africa	1957	45.685	10270856	3013.976
4	Angola	Africa	1957	31.999	4561361	3827.940
5	Argentina	Americas	1957	64.399	19610538	6856.856
6	Australia	Oceania	1957	70.330	9712569	10949.650

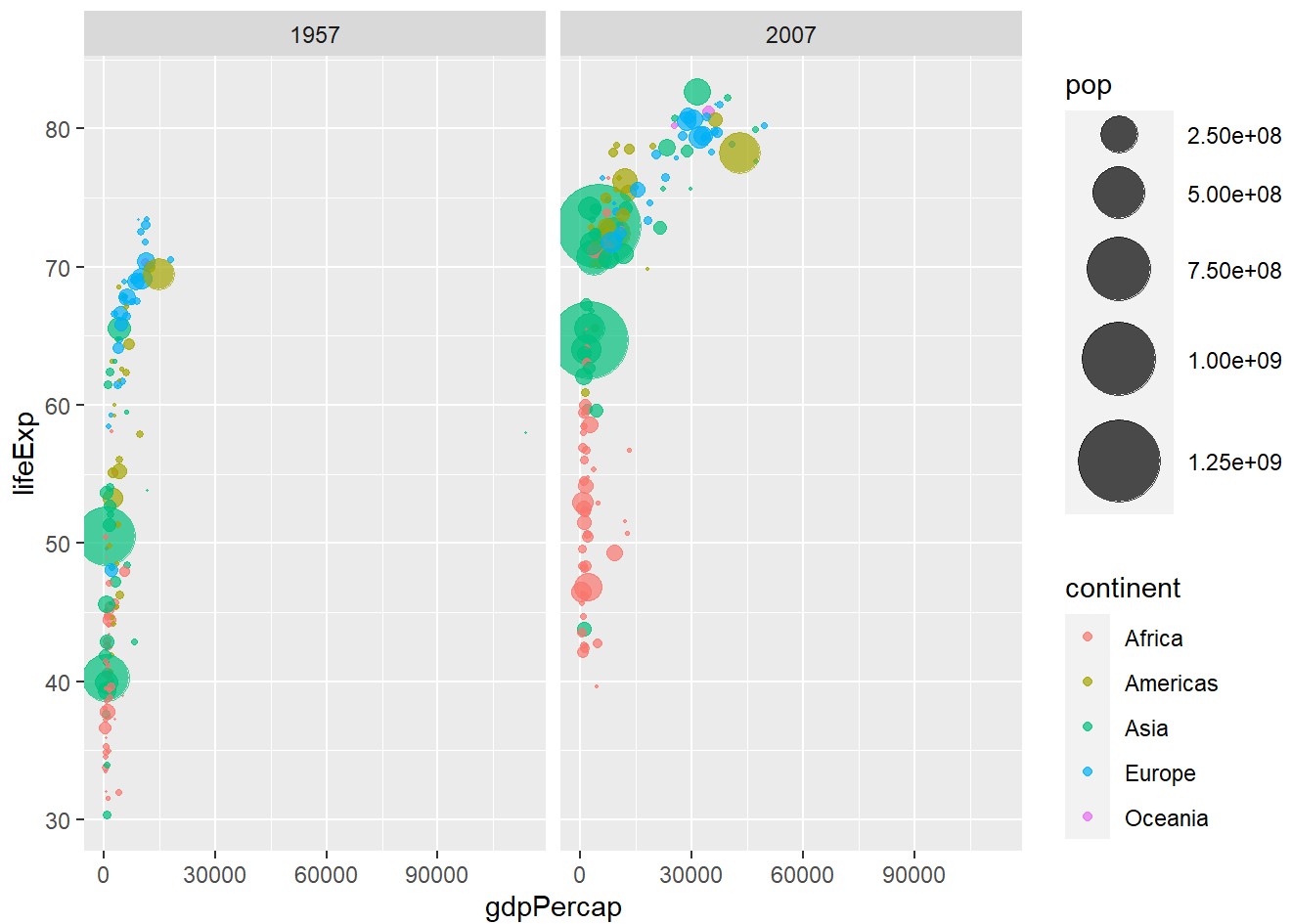
```
ggplot(gapminder_1957) +
  aes(x = gdpPercap, y = lifeExp, color = continent, size = pop) +
  geom_point(alpha=0.7) +
  scale_size_area(max_size = 15) +
  facet_wrap(~year)
```



```
gapminder_1957_2007 <- gapminder %>% filter(year==1957 | year==2007)
head(gapminder_1957_2007)
```

	country	continent	year	lifeExp	pop	gdpPercap
1	Afghanistan	Asia	1957	30.332	9240934	820.8530
2	Afghanistan	Asia	2007	43.828	31889923	974.5803
3	Albania	Europe	1957	59.280	1476505	1942.2842
4	Albania	Europe	2007	76.423	3600523	5937.0295
5	Algeria	Africa	1957	45.685	10270856	3013.9760
6	Algeria	Africa	2007	72.301	33333216	6223.3675

```
ggplot(gapminder_1957_2007) +
  aes(x = gdpPercap, y = lifeExp, color = continent, size = pop) +
  geom_point(alpha=0.7) +
  scale_size_area(max_size = 15) +
  facet_wrap(~year)
```

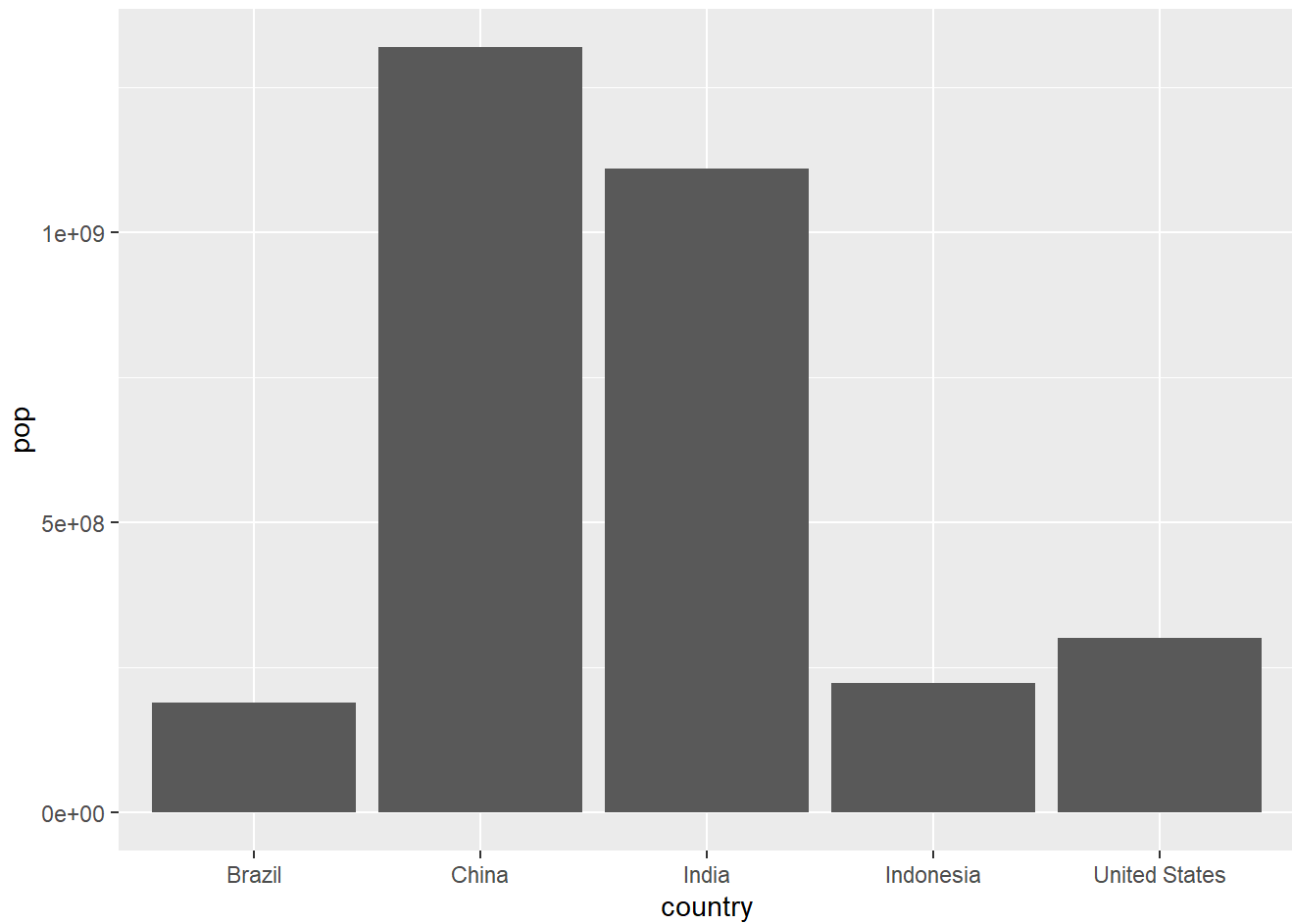


```
gapminder_top5 <- gapminder %>%
  filter(year==2007) %>%
  arrange(desc(pop)) %>%
  top_n(5, pop)
```

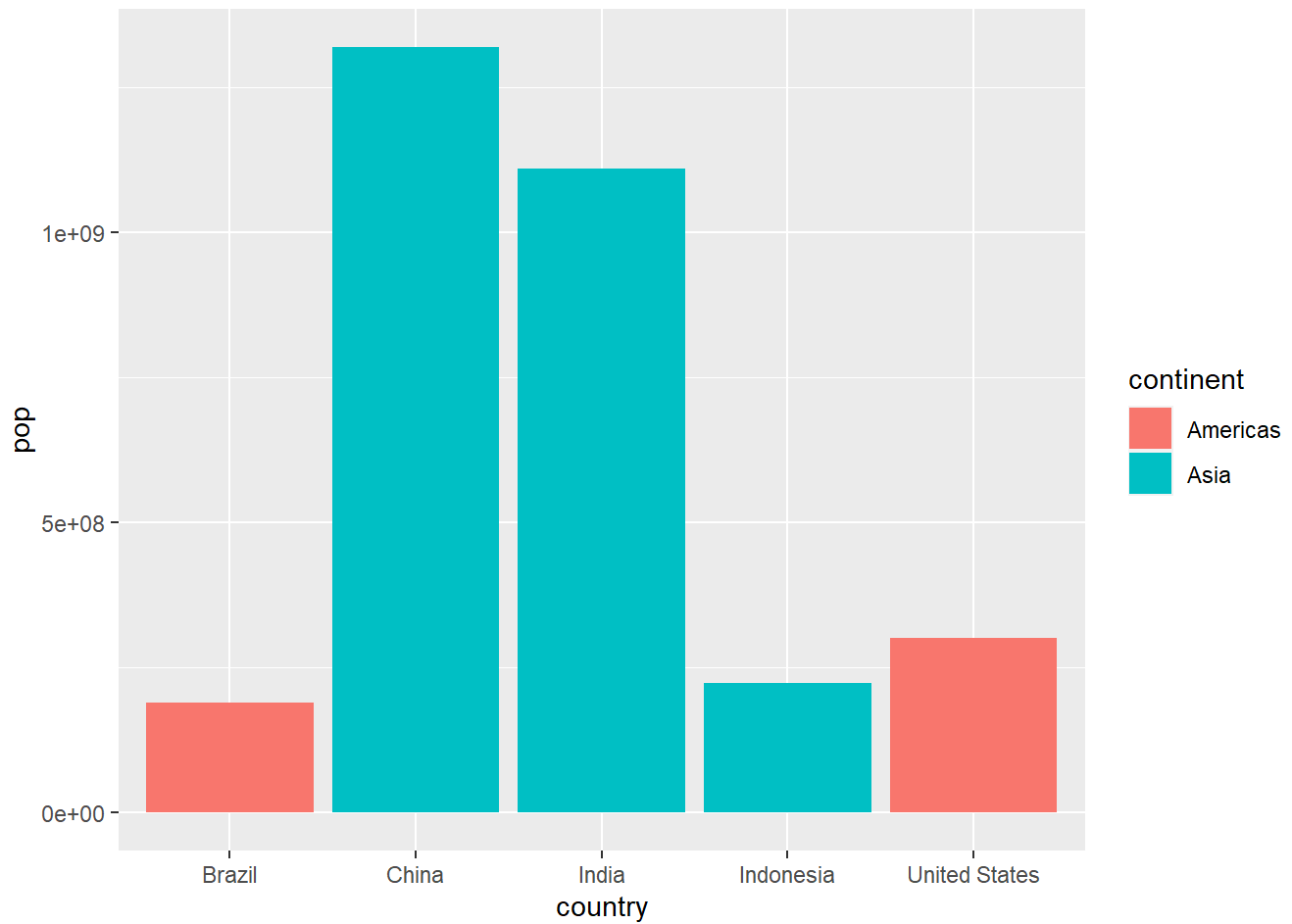
```
gapminder_top5
```

	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

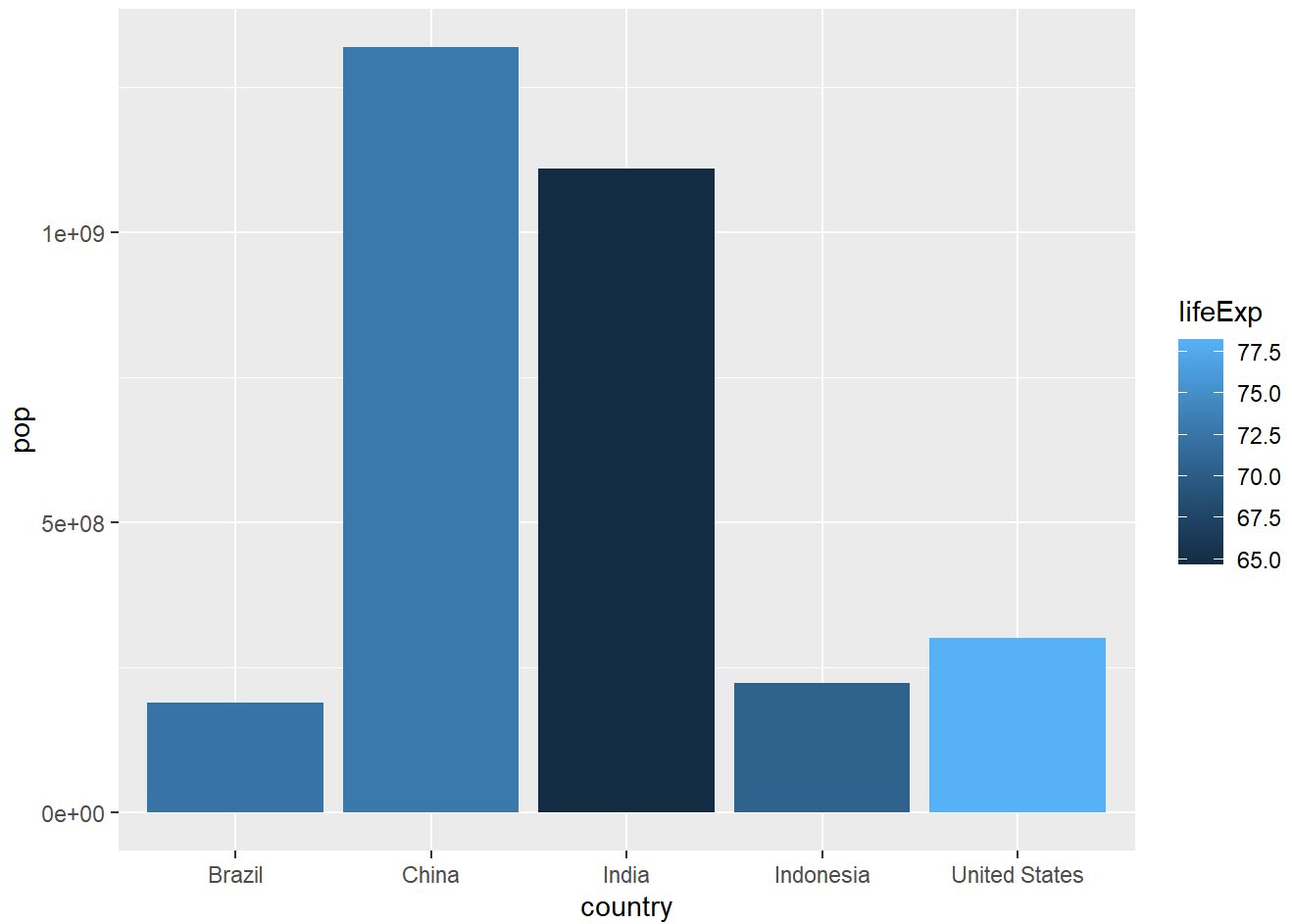
```
ggplot(gapminder_top5) +
  geom_col(aes(x = country, y = pop))
```



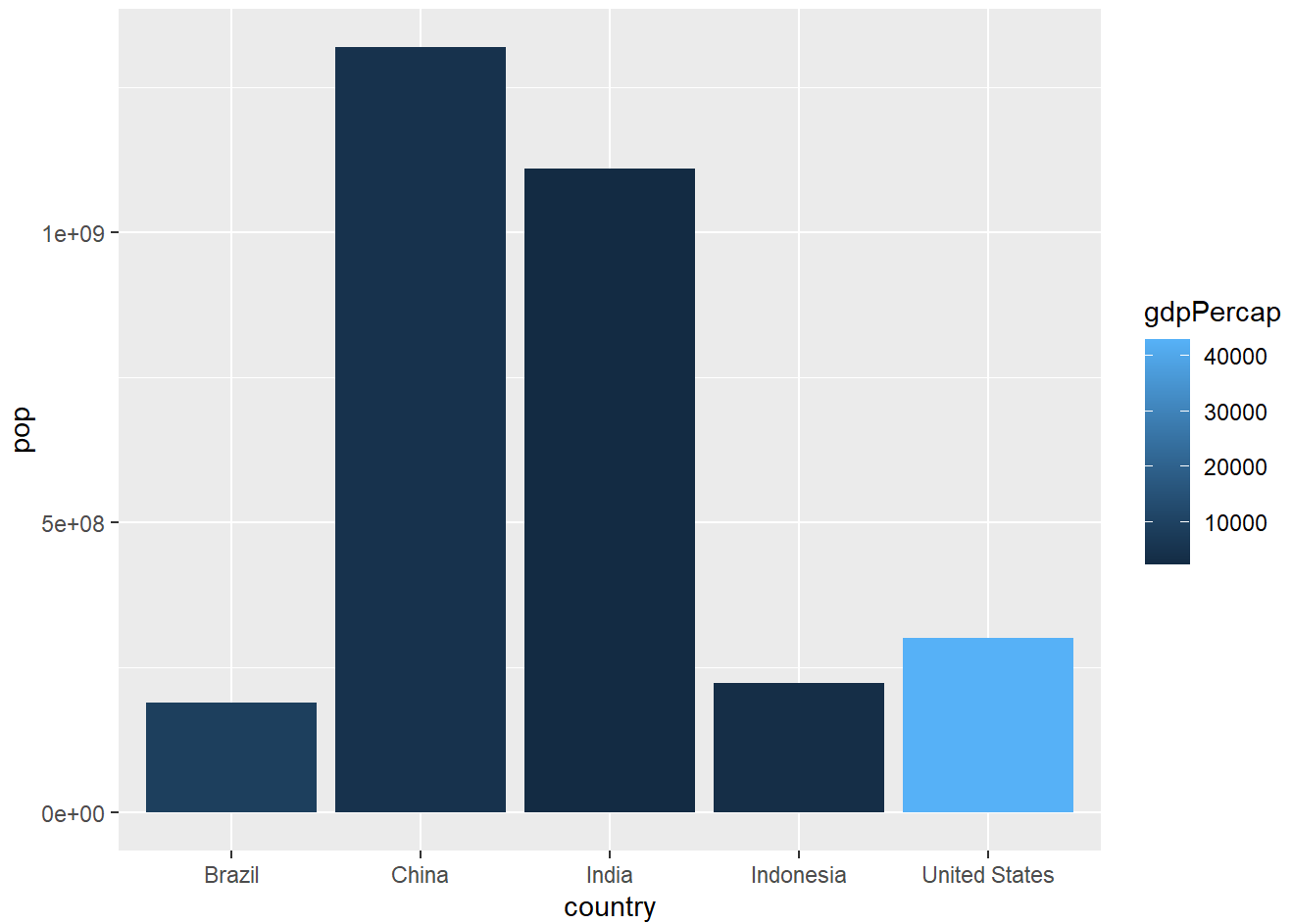
```
ggplot(gapminder_top5) +  
  geom_col(aes(x = country, y = pop, fill = continent))
```



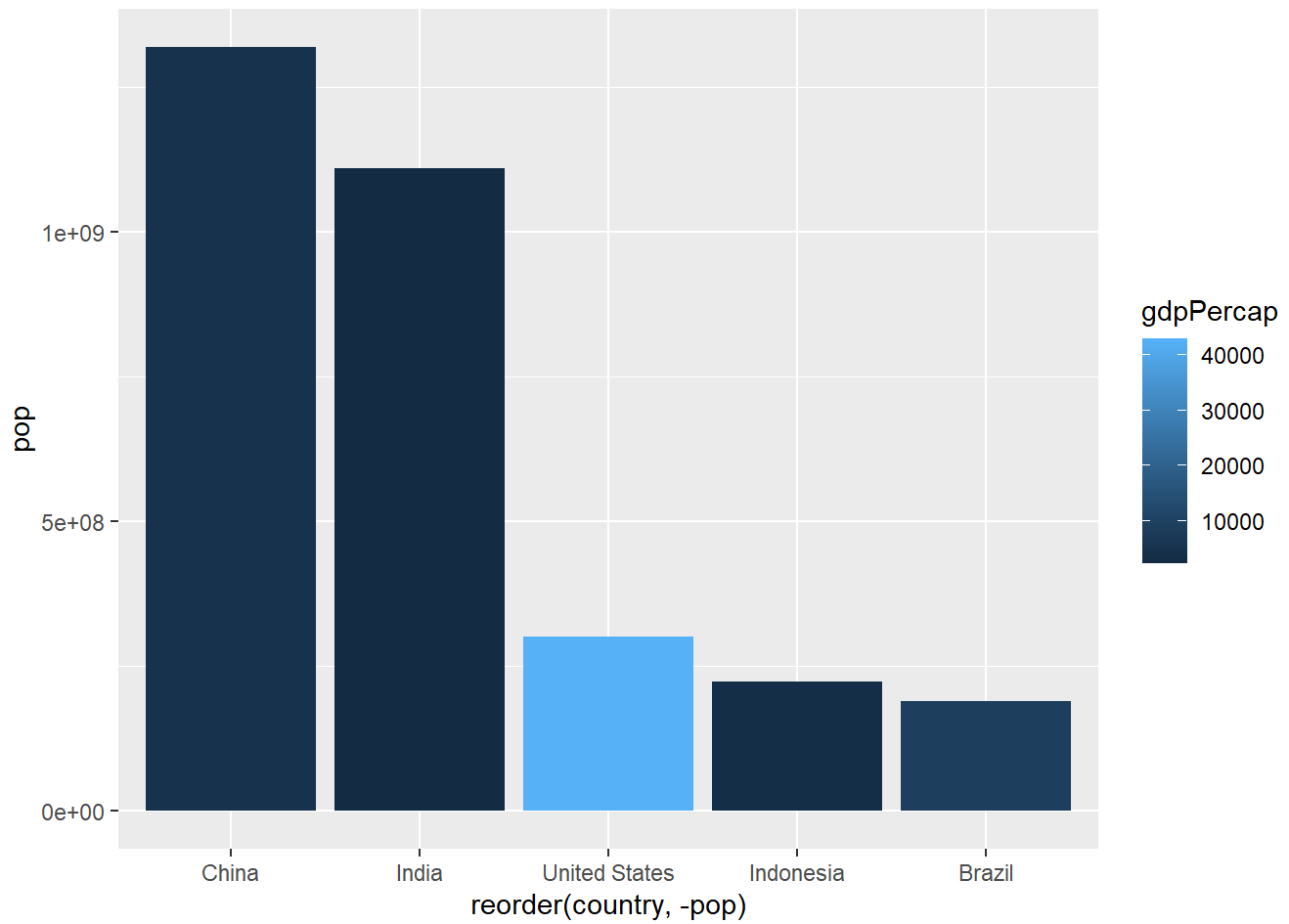
```
ggplot(gapminder_top5) +  
  geom_col(aes(x = country, y = pop, fill = lifeExp))
```

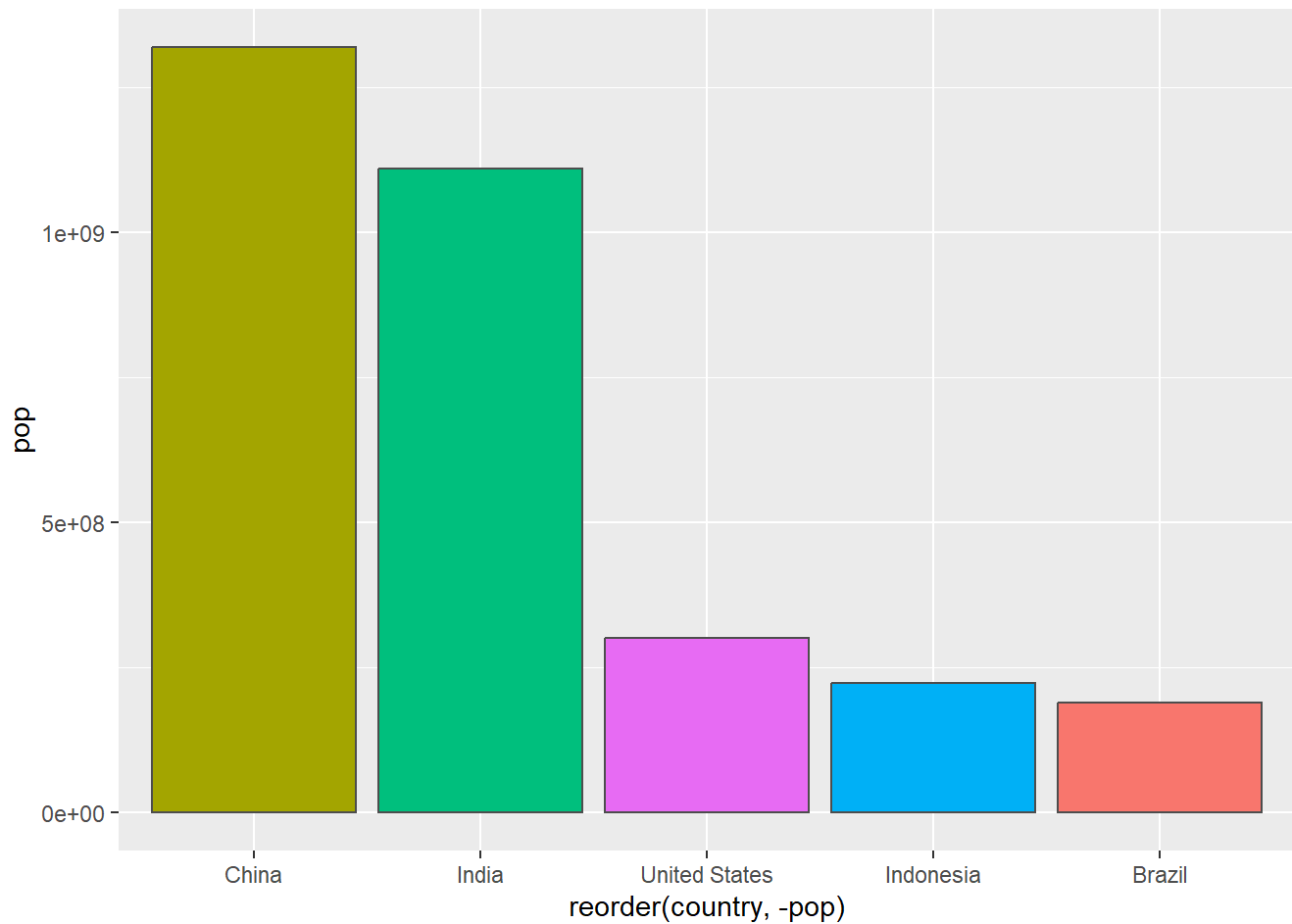
```
ggplot(gapminder_top5) +  
  aes(x=country, y=pop, fill=gdpPercap) +  
  geom_col()
```



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



```
ggplot(gapminder_top5) +  
  aes(x=reorder(country, -pop), y=pop, fill=country) +  
  geom_col(col="gray30") +  
  guides(fill="none")
```

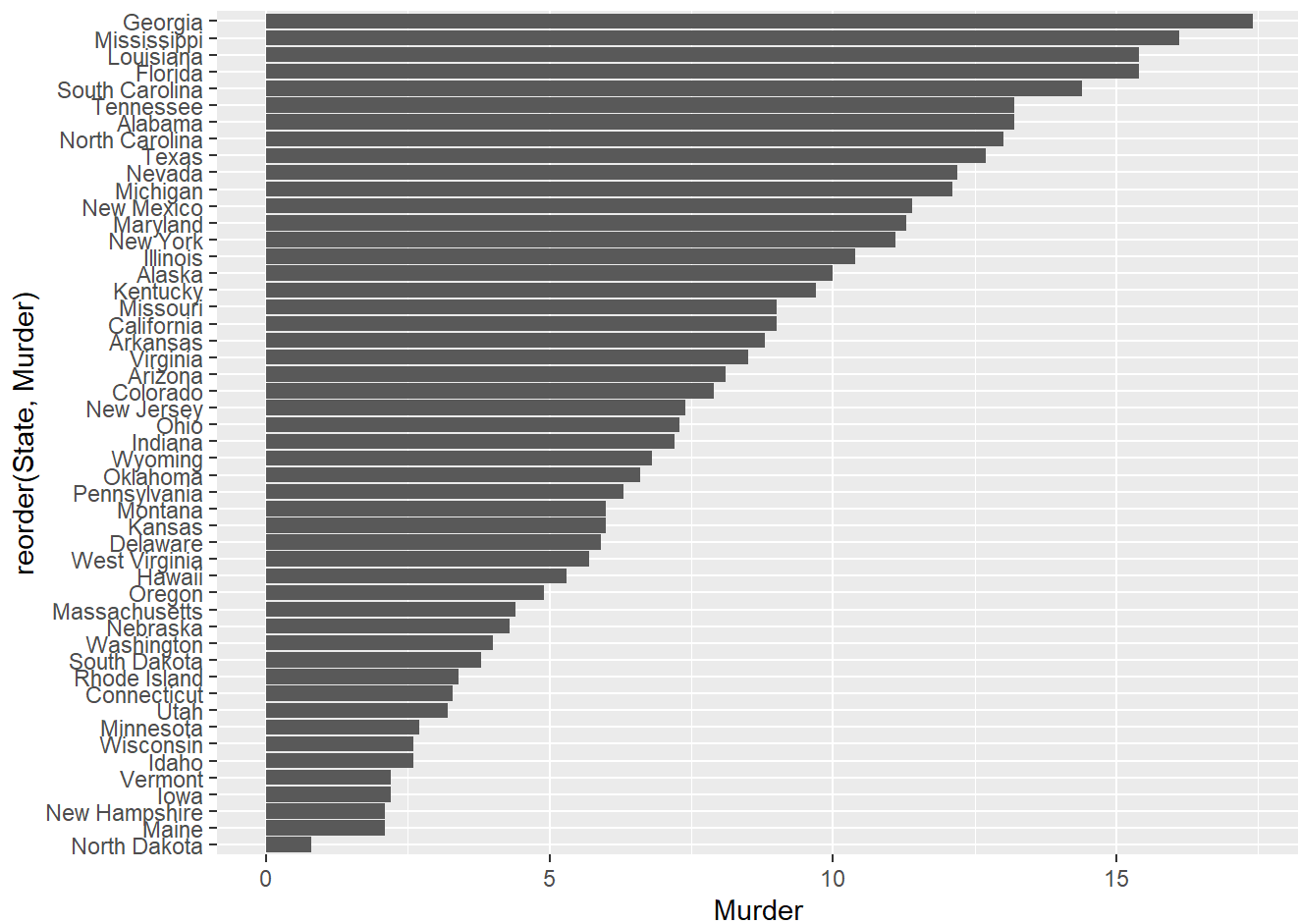


```
head(USArrests)
```

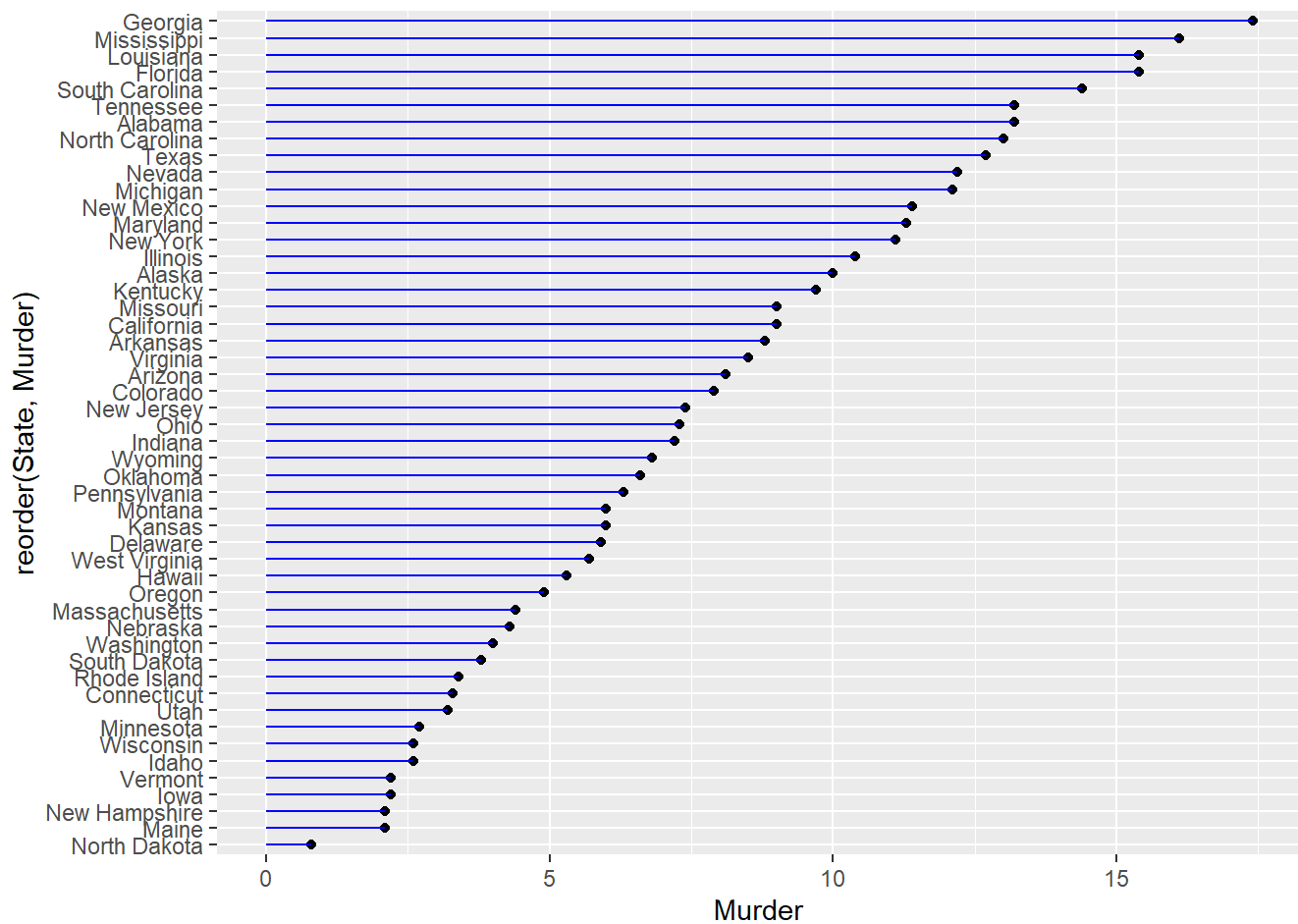
	Murder	Assault	UrbanPop	Rape
Alabama	13.2	236	58	21.2
Alaska	10.0	263	48	44.5
Arizona	8.1	294	80	31.0
Arkansas	8.8	190	50	19.5
California	9.0	276	91	40.6
Colorado	7.9	204	78	38.7

```
USArrests$State <- rownames(USArrests)
```

```
ggplot(USArrests) +
  aes(x=reorder(State,Murder), y=Murder) +
  geom_col() +
  coord_flip()
```



```
ggplot(USArrests) +
  aes(x=reorder(State,Murder), y=Murder) +
  geom_point() +
  geom_segment(aes(x=State,
                  xend=State,
                  y=0,
                  yend=Murder), color="blue") +
  coord_flip()
```



```
library(gapminder)
```

Warning: package 'gapminder' was built under R version 4.3.1

Attaching package: 'gapminder'

The following object is masked _by_ '.GlobalEnv':

gapminder

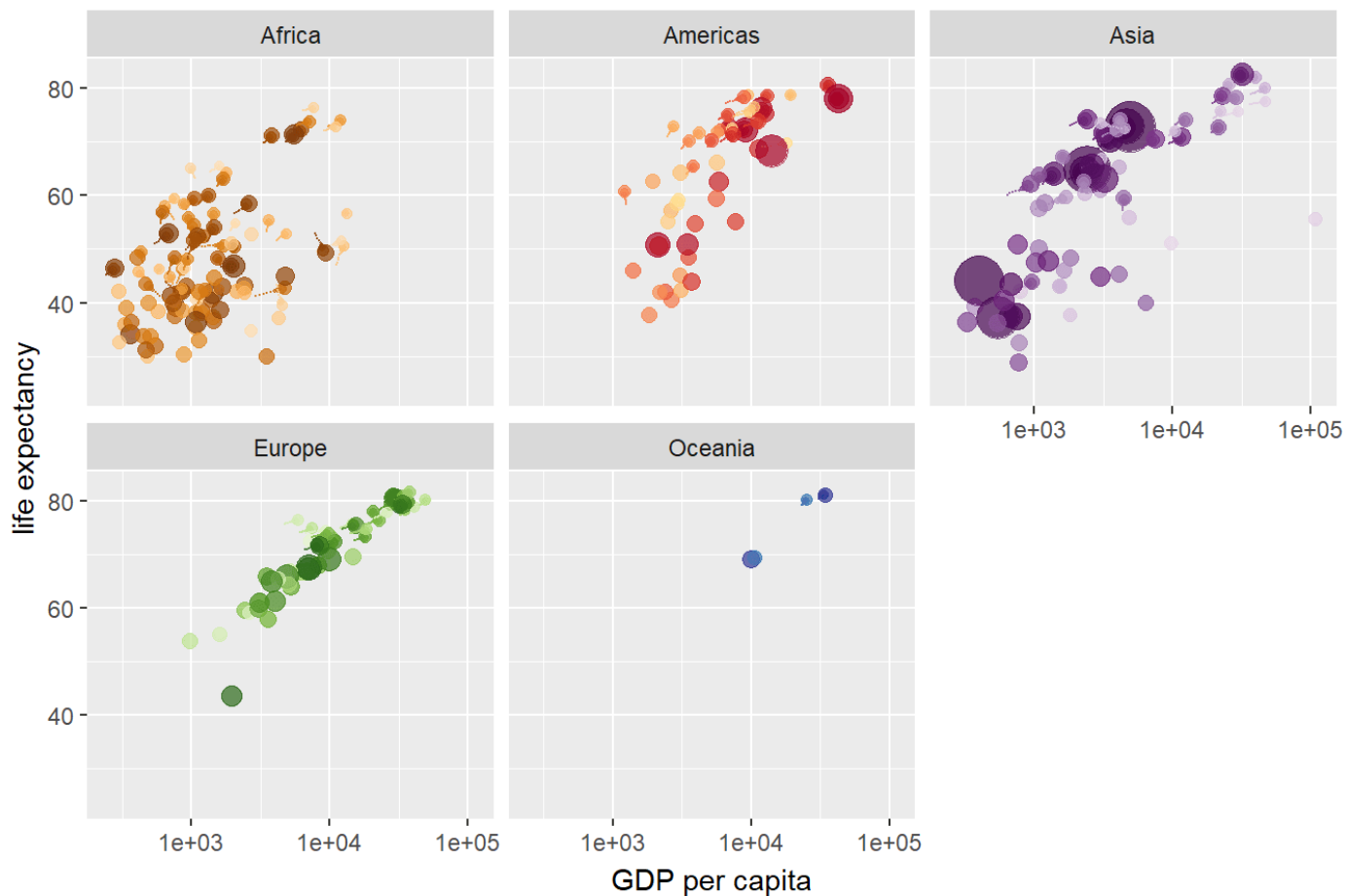
```
library(gganimate)
```

Warning: package 'gganimate' was built under R version 4.3.1

```
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_wrap(~continent) +
  # gganimate
  labs(title = 'Year: {frame_time}', x = 'GDP per capita', y = 'life expectancy') +
```

```
transition_time(year) +
shadow_wake(wake_length = 0.1, alpha = FALSE)
```

Year: 1952



```
library(patchwork)
```

Warning: package 'patchwork' was built under R version 4.3.1

```
p1 <- ggplot(mtcars) + geom_point(aes(mpg, disp))
p2 <- ggplot(mtcars) + geom_boxplot(aes(gear, disp, group = gear))
p3 <- ggplot(mtcars) + geom_smooth(aes(dis, qsec))
p4 <- ggplot(mtcars) + geom_bar(aes(carb))
```

```
(p1 | p2 | p3) / p4
```

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

