

class05.R

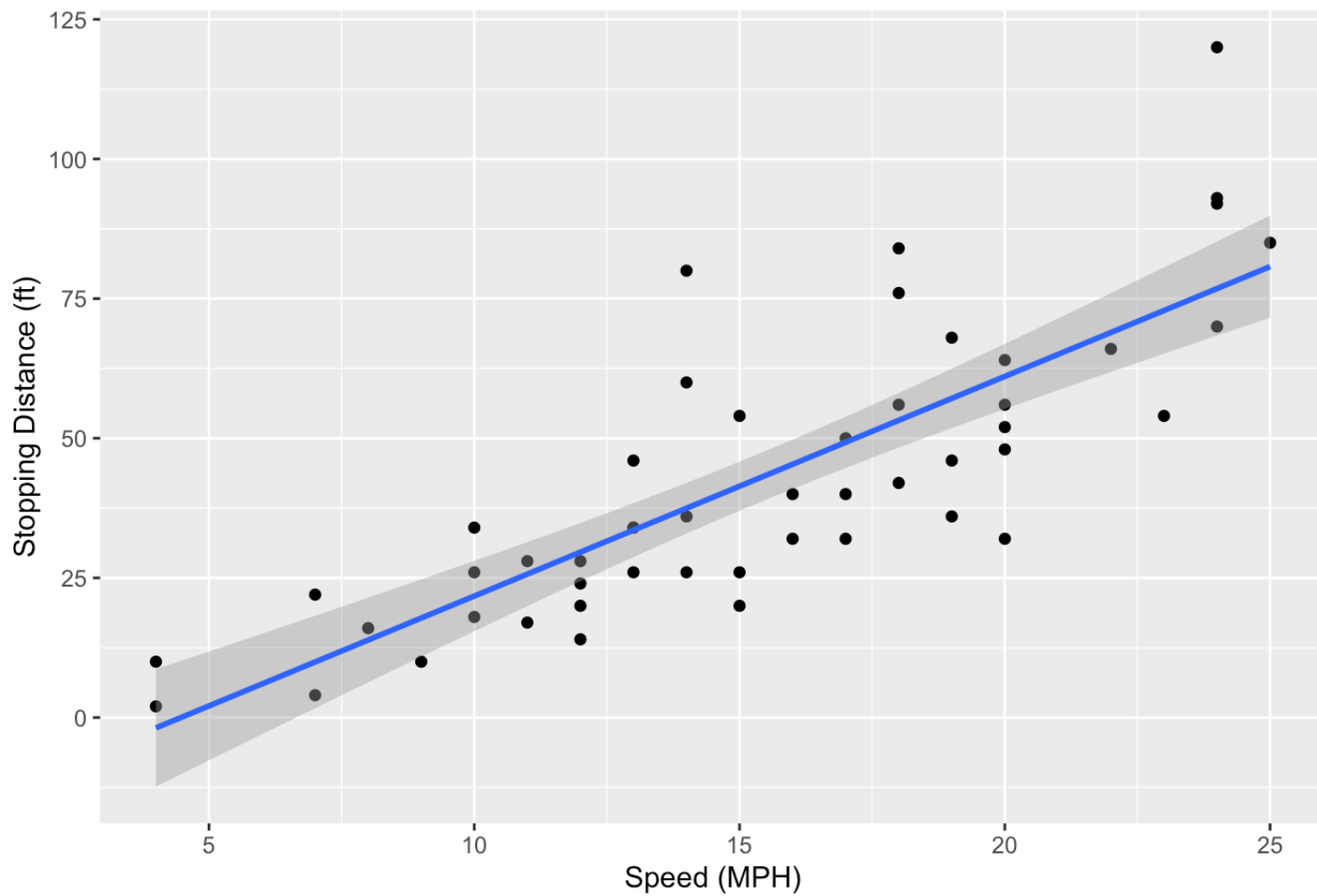
ShitianLi

2021-10-13

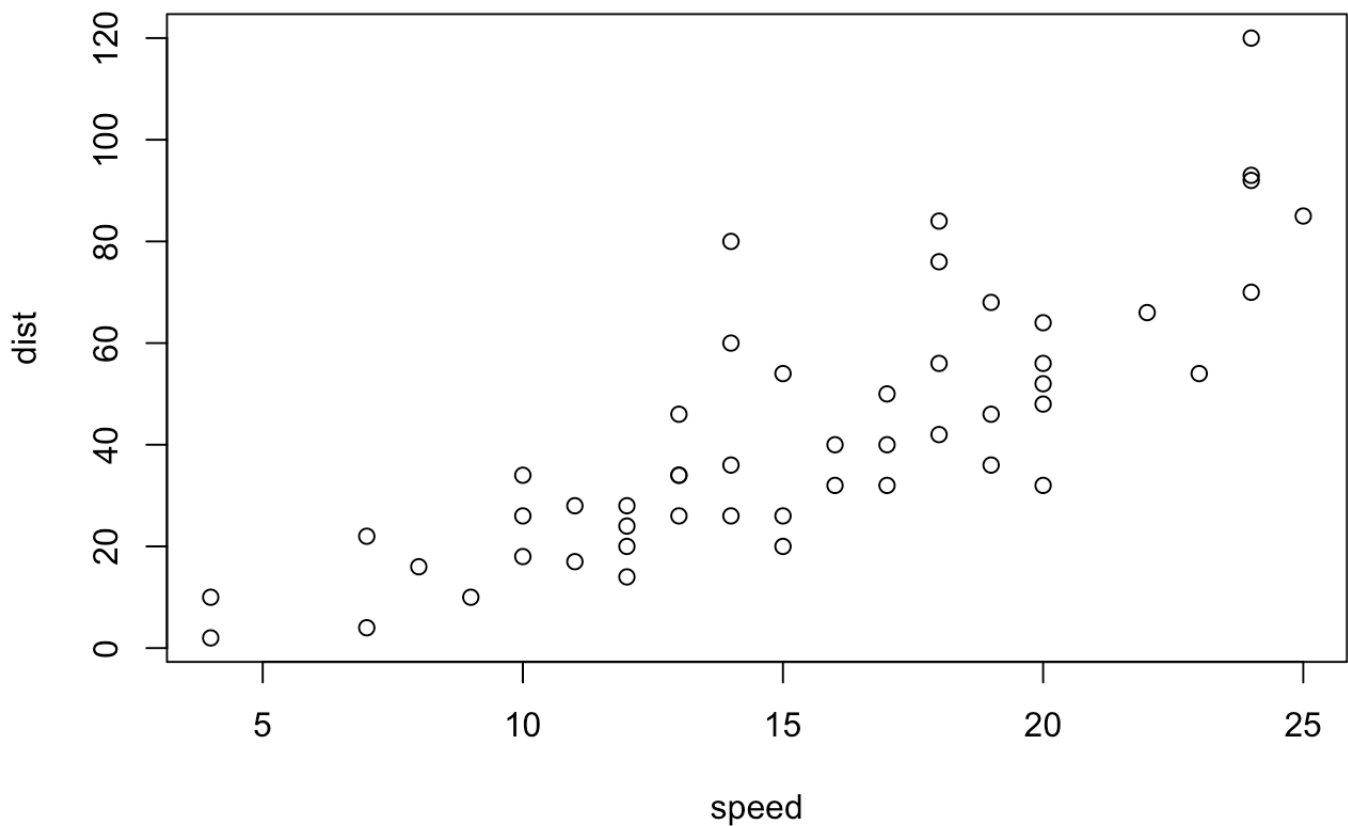
```
# Class 05: Data Visualization  
# Trying the ggplot2 package  
  
# First, install and load required packages:  
# install.packages("ggplot2")  
library(ggplot2)  
  
# we'll try with inbuilt cars dataset  
ggplot(cars) +  
  aes(x=speed, y=dist) +  
  geom_point() +  
  geom_smooth(method = "lm") +  
  labs(title = "Stopping Distance of Old Cars",  
        x = "Speed (MPH)",  
        y = "Stopping Distance (ft)")
```

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

Stopping Distance of Old Cars



```
# side note: R has in-built plotting  
plot(cars)
```



```
# import gene expression data
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
```

```
# Q. How many genes?
nrow(genes)
```

```
## [1] 5196
```

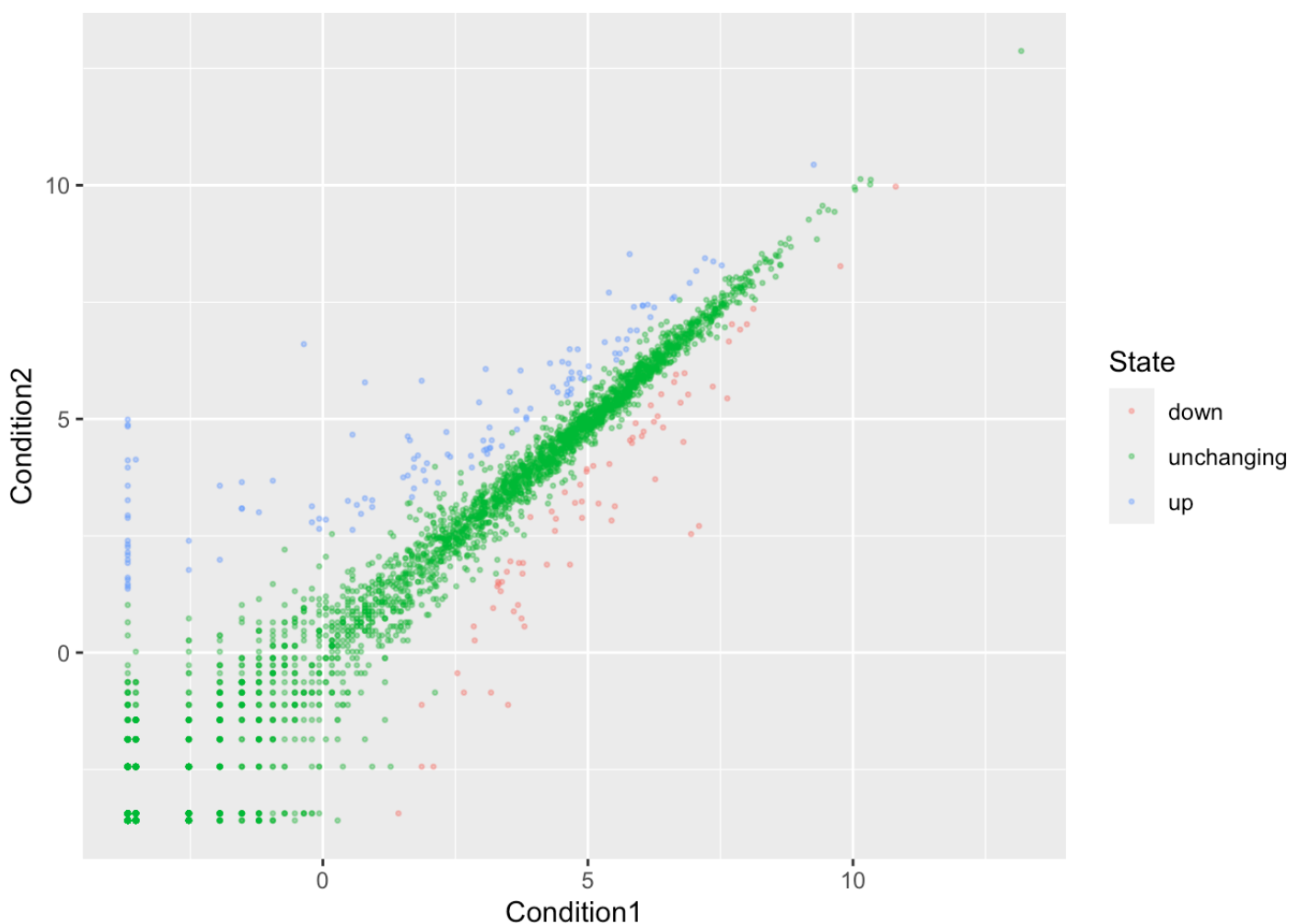
```
# Q. How many genes are up-regulated?  
table(genes$State)
```

```
##  
##      down  unchanging      up  
##      72    4997      127
```

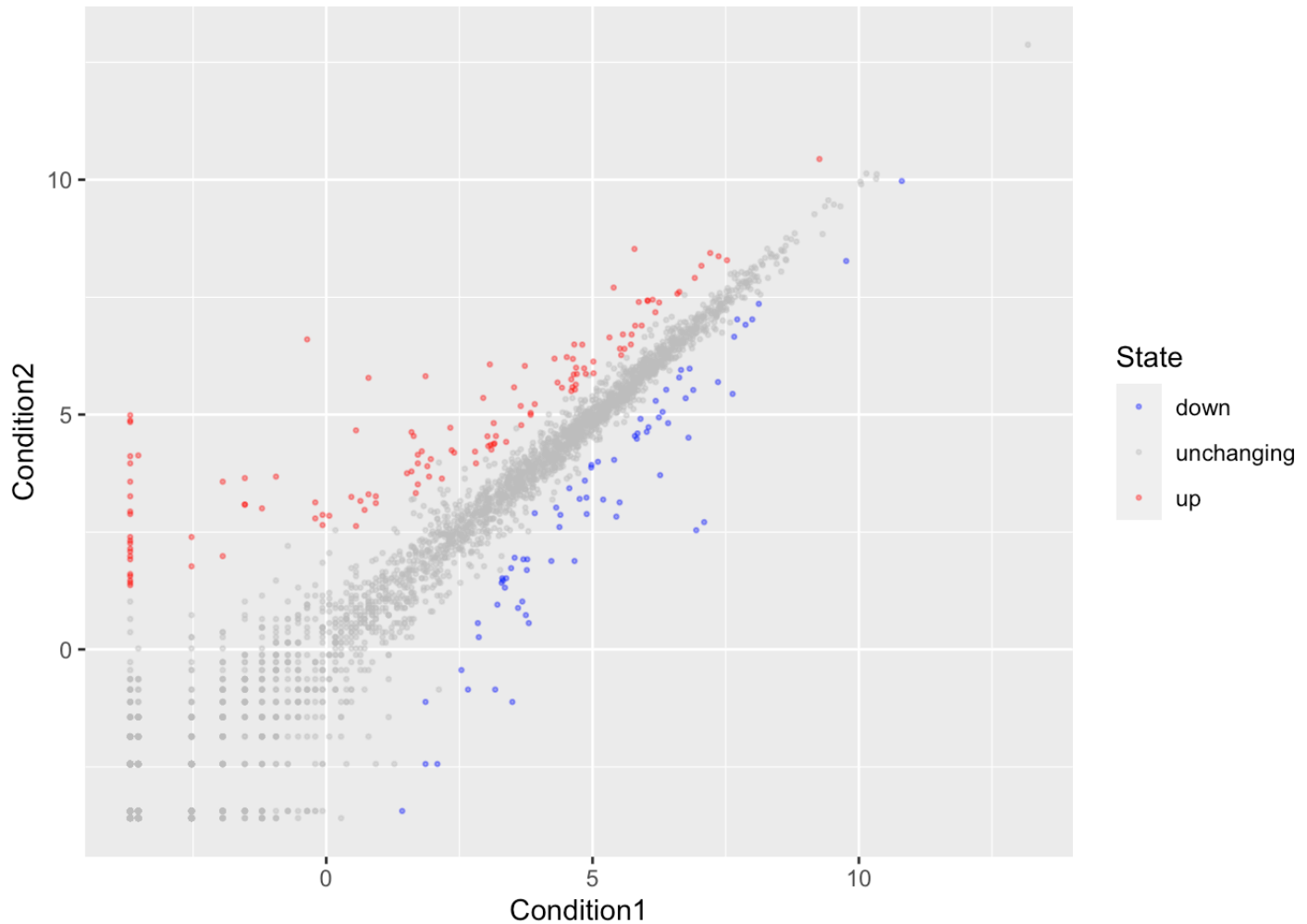
```
# Q. What percentage is up?  
round(table(genes$State) / nrow(genes) * 100, 2)
```

```
##  
##      down  unchanging      up  
##      1.39    96.17      2.44
```

```
# Let's make a figure  
p <- ggplot(genes, aes(Condition1, Condition2, col=State)) +  
  geom_point(alpha = 0.4, size = 0.5)  
p
```



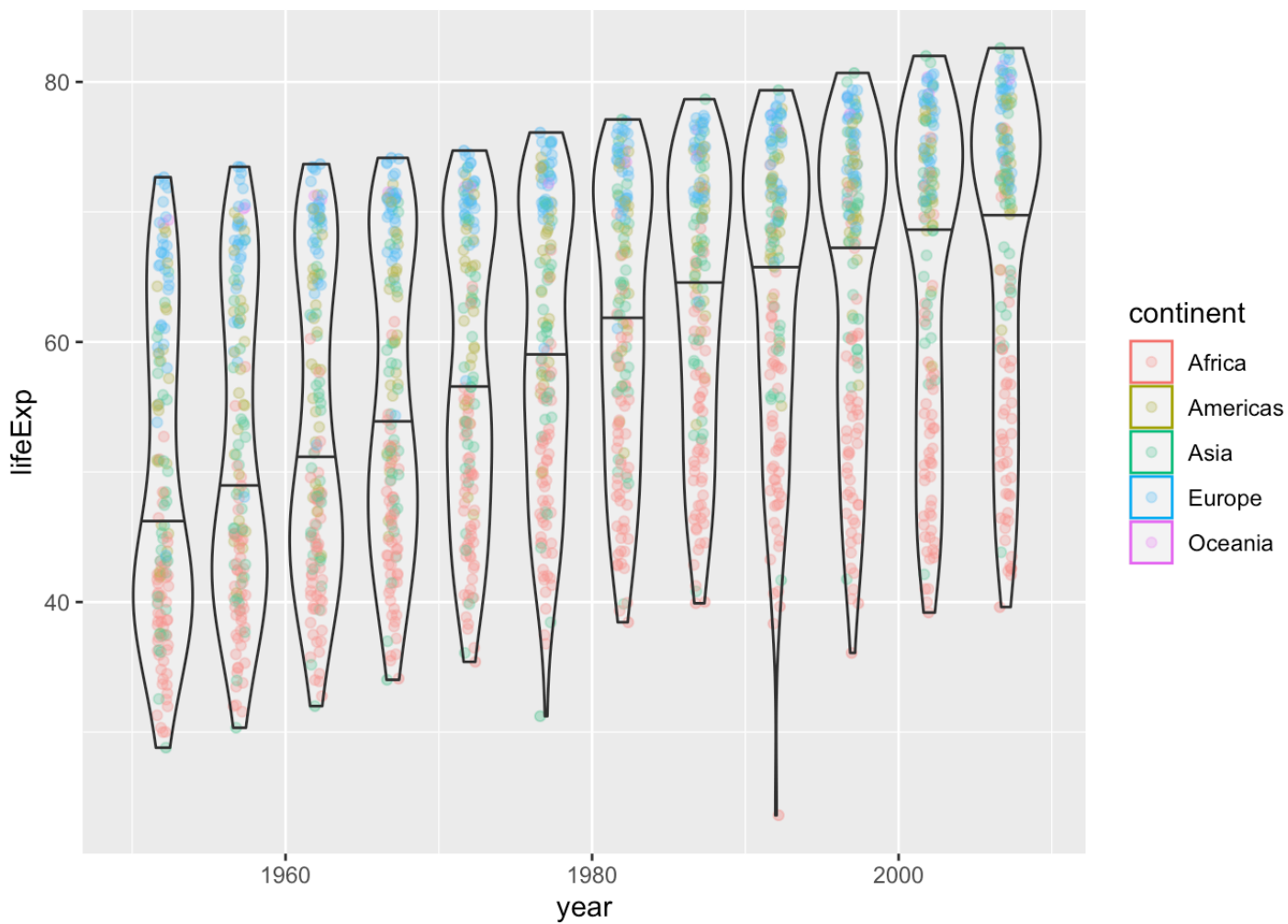
```
# change color scheme
p + scale_color_manual(values = c("blue", "grey", "red"))
```



```
# Let's explore thegapminder dataset
# install.packages("gapminder")
library(gapminder)
head(gapminder)
```

```
## # A tibble: 6 × 6
##   country    continent  year lifeExp      pop gdpPercap
##   <fct>      <fct>    <int>  <dbl>    <int>    <dbl>
## 1 Afghanistan Asia      1952   28.8  8425333    779.
## 2 Afghanistan Asia      1957   30.3  9240934    821.
## 3 Afghanistan Asia      1962   32.0 10267083    853.
## 4 Afghanistan Asia      1967   34.0 11537966    836.
## 5 Afghanistan Asia      1972   36.1 13079460    740.
## 6 Afghanistan Asia      1977   38.4 14880372    786.
```

```
# Let's make a new plot of year vs. lif exp
ggplot(gapminder, aes(year, lifeExp, col=continent)) +
  geom_jitter(alpha=0.3, width=0.4) +
  geom_violin(aes(group=year), alpha = 0.2,
             draw_quantiles = 0.5)
```



```
# install the plotly
# install.packages("plotly")
library(plotly)
```

```
##
## Attaching package: 'plotly'
```

```
## The following object is masked from 'package:ggplot2':
##
## last_plot
```

```
## The following object is masked from 'package:stats':
```

```
##
```

```
## filter
```

```
## The following object is masked from 'package:graphics':
```

```
##
```

```
## layout
```

```
ggplotly()
```

