

Speaking Science

W241 Final Project

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```
d <- fread('speaking_science_data_03-24_clean.csv')
head(d)
```

```
##      start_date      end_date      ip_address duration_in_seconds
## 1: 3/23/2020 20:59 3/23/2020 21:00 196.17.67.134             85
## 2: 3/23/2020 21:00 3/23/2020 21:03 68.105.189.229            185
## 3: 3/23/2020 20:59 3/23/2020 21:03 76.176.54.192            284
## 4: 3/23/2020 21:00 3/23/2020 21:06 107.185.127.204           379
## 5: 3/23/2020 21:04 3/23/2020 21:06 71.6.87.50              163
## 6: 3/23/2020 21:02 3/23/2020 21:06 72.216.72.106            275
##      recorded_date      response_id latitude longitude      mturk_id
## 1: 3/23/2020 21:00 R_3si5y4qwGLzKYPh 34.05440 -118.2440 A1201H00DXBGC3
## 2: 3/23/2020 21:03 R_1jqhwR0mmrLPaoy 36.05881 -115.3104 A12ATVBE1I4567
## 3: 3/23/2020 21:03 R_2ASHk9ILabLrZCB 33.02870 -117.0846 A900V3976AFYF
## 4: 3/23/2020 21:06 R_3n6PLtv1K8spEfC 33.96750 -118.1464 A2C73Y1COWCA51
## 5: 3/23/2020 21:06 R_3kbIZqjBa0kb1pG 37.76880 -122.2620 A10ROXYXMV5MBO
## 6: 3/23/2020 21:06 R_sRUrOCfBuUjTAuB 32.89461 -111.7493 A83OLM1ZQC083
##      browser_type browser_version      browser_os browser_resolution
## 1:      Chrome      80.0.3987.149 Windows NT 6.3      1366x768
## 2:      Chrome      80.0.3987.149 Windows NT 10.0     1536x864
## 3:      Chrome      79.0.3945.136      Android 7.0      360x640
## 4:      Chrome      80.0.3987.149 Windows NT 10.0     1920x1080
## 5:      Chrome      80.0.3987.132      Macintosh      1440x900
## 6:      Chrome      80.0.3987.149 Windows NT 10.0     1536x864
##      time_read_intro time_read_article credibility importance q1 q1_correct q2
## 1:          2.607          17.922             5             4 1             0 3
## 2:          7.588          29.649             5             4 3             1 5
## 3:         13.345         166.381             6             7 1             0 3
## 4:          2.854          67.943             7             6 4             0 3
## 5:         18.607          47.366             6             6 2             0 1
## 6:         10.128         163.601             7             7 3             1 1
##      q2_correct q3 q3_correct q4 q4_correct q5 q5_correct q6 q6_correct
## 1:          1 2             0 1             0 3             0 NA             0
```

```
## 2:      0 3      1 1      0 4      1 1      1
## 3:      1 1      0 4      0 4      1 1      1
## 4:      1 3      1 2      0 2      0 NA      0
## 5:      0 1      0 2      0 1      0 1      1
## 6:      0 3      1 1      0 3      0 1      1
##      questions_correct time_answering_questions donation time_donation
## 1:      1      22.689      50      7.384
## 2:      4     125.852      1      4.044
## 3:      3      67.334      1      7.657
## 4:      2     287.135      0      5.991
## 5:      1      49.709     50      7.145
## 6:      3      72.550      0     14.611
##      city state   zip treatment
## 1: Los Angeles   CA 90009      1
## 2:  Las Vegas    NV 89113      0
## 3:  San Diego    CA 92127      0
## 4: Bell Gardens   CA 90201      1
## 5:  Vallejo      CA 94589      1
## 6: Casa Grande   AZ 85122      1
```

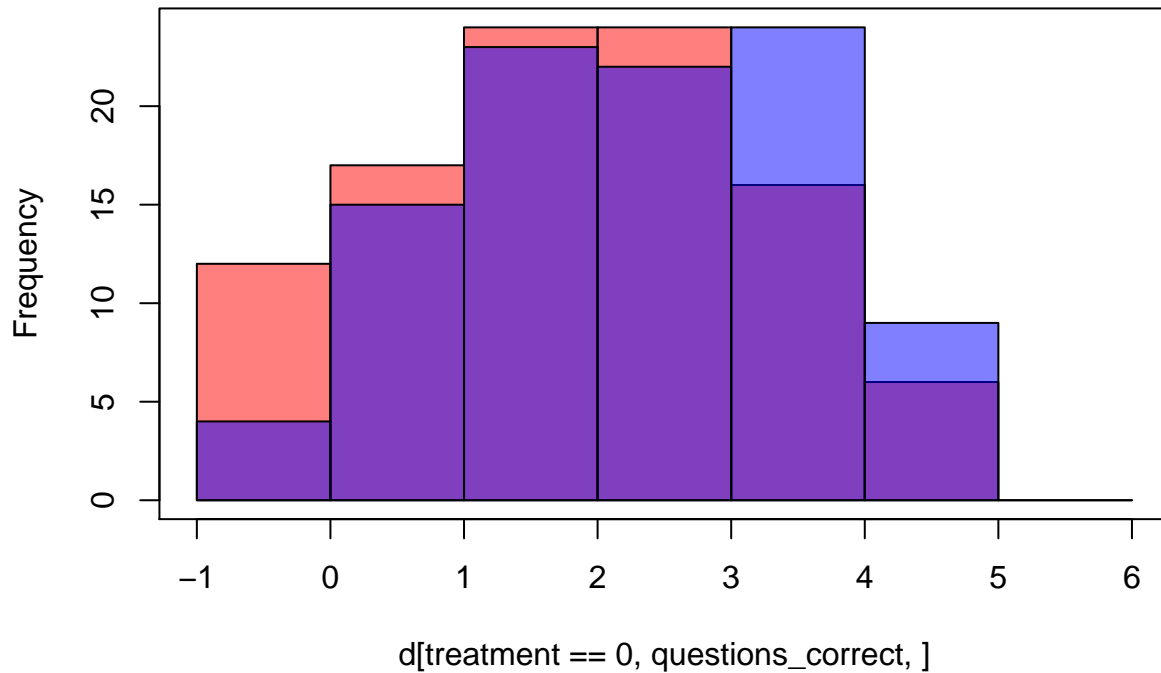
Functions

Simple Linear Regression

```
mod <- lm(questions_correct ~ treatment + time_read_article, data=d)
summary(mod)

##
## Call:
## lm(formula = questions_correct ~ treatment + time_read_article,
##     data = d)
##
## Residuals:
##      Min       1Q   Median       3Q      Max
## -2.69093 -0.94764  0.06458  0.91228  2.80838
##
## Coefficients:
##              Estimate Std. Error t value Pr(>|t|)
## (Intercept)   1.6241406  0.1641966   9.891  < 2e-16 ***
## treatment      0.4009627  0.1781902   2.250   0.0256 *
## time_read_article 0.0037659  0.0005634   6.685 2.42e-10 ***
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 1.247 on 193 degrees of freedom
## Multiple R-squared:  0.2074, Adjusted R-squared:  0.1992
## F-statistic: 25.25 on 2 and 193 DF, p-value: 1.817e-10
hist(d[treatment == 0, questions_correct,], col=rgb(1,0,0,0.5), breaks=seq(-1,6, by=1))
hist(d[treatment == 1, questions_correct,], col=rgb(0,0,1,0.5), breaks=seq(-1,6, by=1), add = T)
box()
```

Histogram of d[treatment == 0, questions_correct,]



```
stargazer(mod, type = "text")
```

```
##
## =====
##                               Dependent variable:
##                               -----
##                               questions_correct
##                               -----
## treatment                      0.401**
##                               (0.178)
##
## time_read_article              0.004***
##                               (0.001)
##
## Constant                      1.624***
##                               (0.164)
##
## -----
## Observations                   196
## R2                             0.207
## Adjusted R2                   0.199
## Residual Std. Error          1.247 (df = 193)
## F Statistic                   25.249*** (df = 2; 193)
## =====
## Note:                          *p<0.1; **p<0.05; ***p<0.01
```

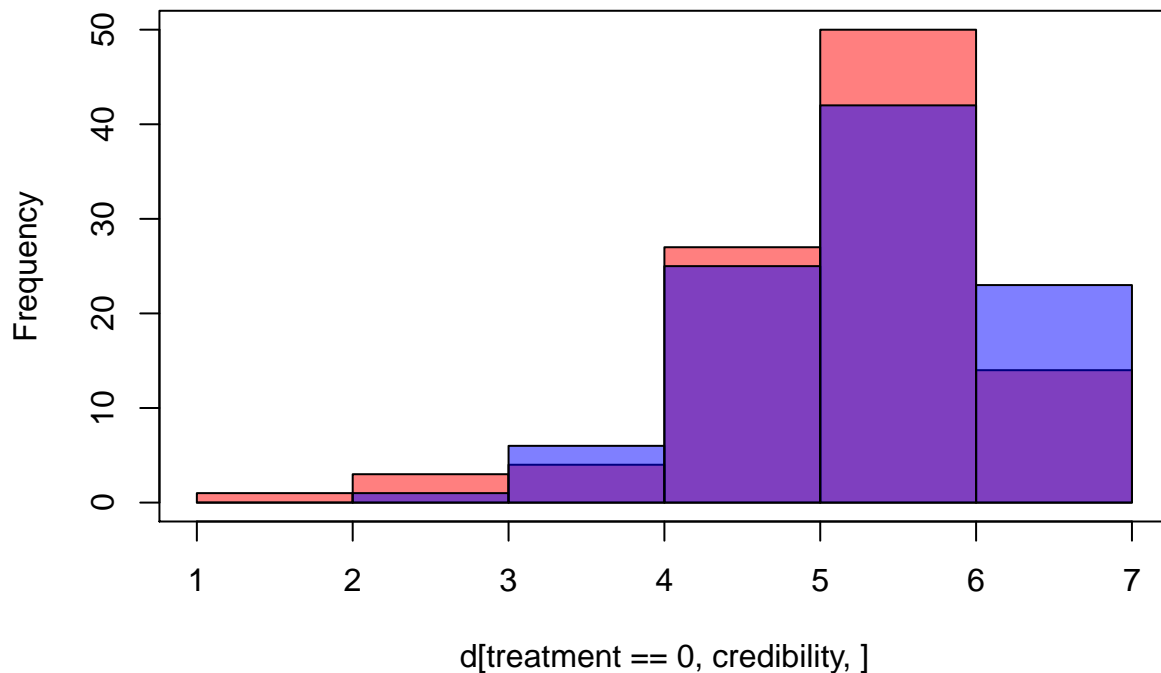
```

mod <- lm(credibility ~ treatment, data=d)
summary(mod)

##
## Call:
## lm(formula = credibility ~ treatment, data = d)
##
## Residuals:
##      Min       1Q   Median       3Q      Max
## -3.6566 -0.6566  0.1753  0.3434  1.3434
##
## Coefficients:
##              Estimate Std. Error t value Pr(>|t|)
## (Intercept)  5.65657    0.09306  60.781  <2e-16 ***
## treatment    0.16818    0.13229   1.271   0.205
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 0.926 on 194 degrees of freedom
## Multiple R-squared:  0.008262,    Adjusted R-squared:  0.00315
## F-statistic: 1.616 on 1 and 194 DF,  p-value: 0.2052
hist(d[treatment == 0, credibility,], col=rgb(1,0,0,0.5), breaks=seq(1,7, by=1))
hist(d[treatment == 1, credibility,], col=rgb(0,0,1,0.5), breaks=seq(1,7, by=1), add = T)
box()

```

Histogram of d[treatment == 0, credibility,]



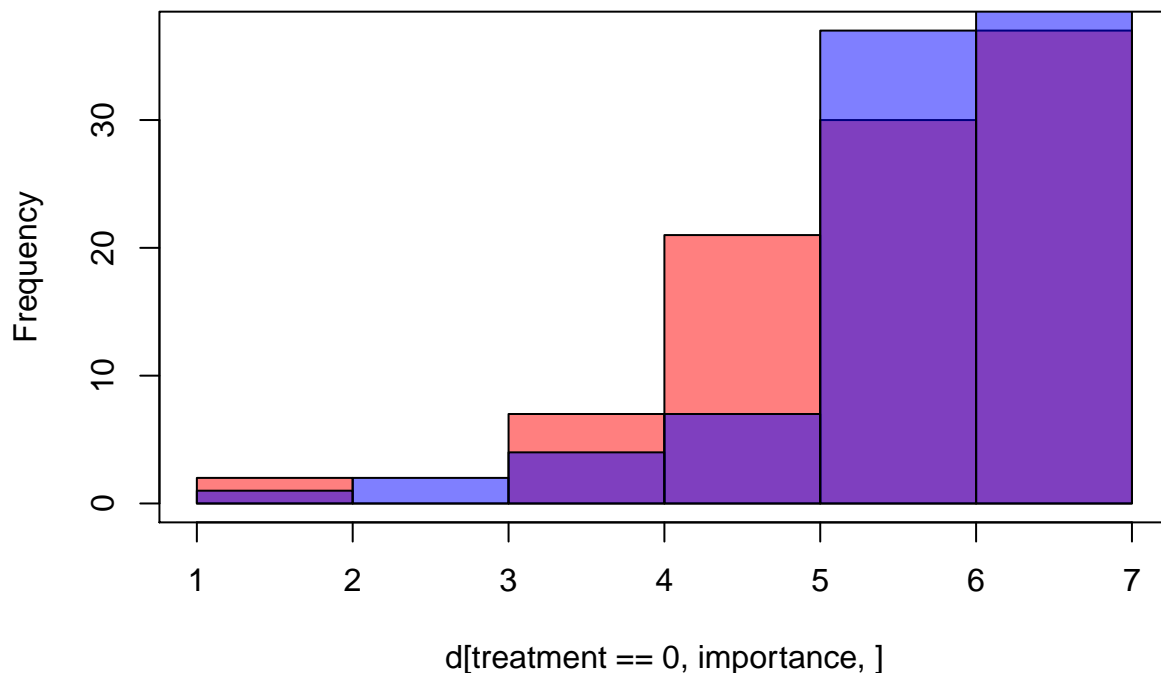
```

mod <- lm(importance ~ treatment, data=d)
summary(mod)

##
## Call:
## lm(formula = importance ~ treatment, data = d)
##
## Residuals:
##      Min       1Q   Median       3Q      Max
## -5.2062 -0.2062  0.0825  0.7938  1.0825
##
## Coefficients:
##              Estimate Std. Error t value Pr(>|t|)
## (Intercept)   5.9175     0.1143  51.782  <2e-16 ***
## treatment     0.2887     0.1616   1.786   0.0757 .
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 1.126 on 192 degrees of freedom
## (2 observations deleted due to missingness)
## Multiple R-squared:  0.01634,    Adjusted R-squared:  0.01122
## F-statistic:  3.19 on 1 and 192 DF,  p-value: 0.07566
hist(d[treatment == 0, importance,], col=rgb(1,0,0,0.5), breaks=seq(1,7, by=1))
hist(d[treatment == 1, importance,], col=rgb(0,0,1,0.5), breaks=seq(1,7, by=1), add = T)
box()

```

Histogram of d[treatment == 0, importance,]



```

mod <- lm(time_read_article ~ treatment, data=d)
summary(mod)

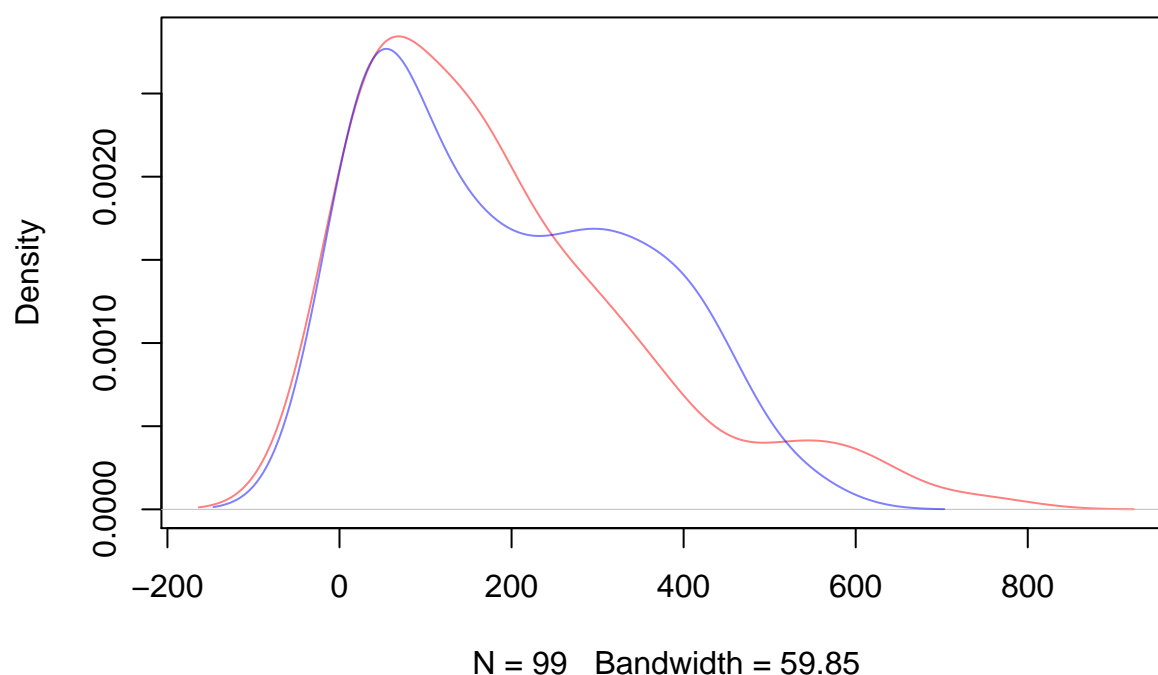
##
## Call:
## lm(formula = time_read_article ~ treatment, data = d)
##
## Residuals:
##      Min       1Q   Median       3Q      Max
## -179.58 -138.20  -32.26   98.13  555.57
##
## Coefficients:
##              Estimate Std. Error t value Pr(>|t|)
## (Intercept)  188.318     15.971   11.792 <2e-16 ***
## treatment      7.592     22.702    0.334   0.738
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 158.9 on 194 degrees of freedom
## Multiple R-squared:  0.0005761, Adjusted R-squared:  -0.004576
## F-statistic: 0.1118 on 1 and 194 DF,  p-value: 0.7384

d1 <- density(d[treatment == 0, time_read_article,])
d2 <- density(d[treatment == 1, time_read_article,])

plot(d1, col=rgb(1,0,0,0.5))
lines(d2, col=rgb(0,0,1,0.5))

```

```
density.default(x = d[treatment == 0, time_read_article, ])
```

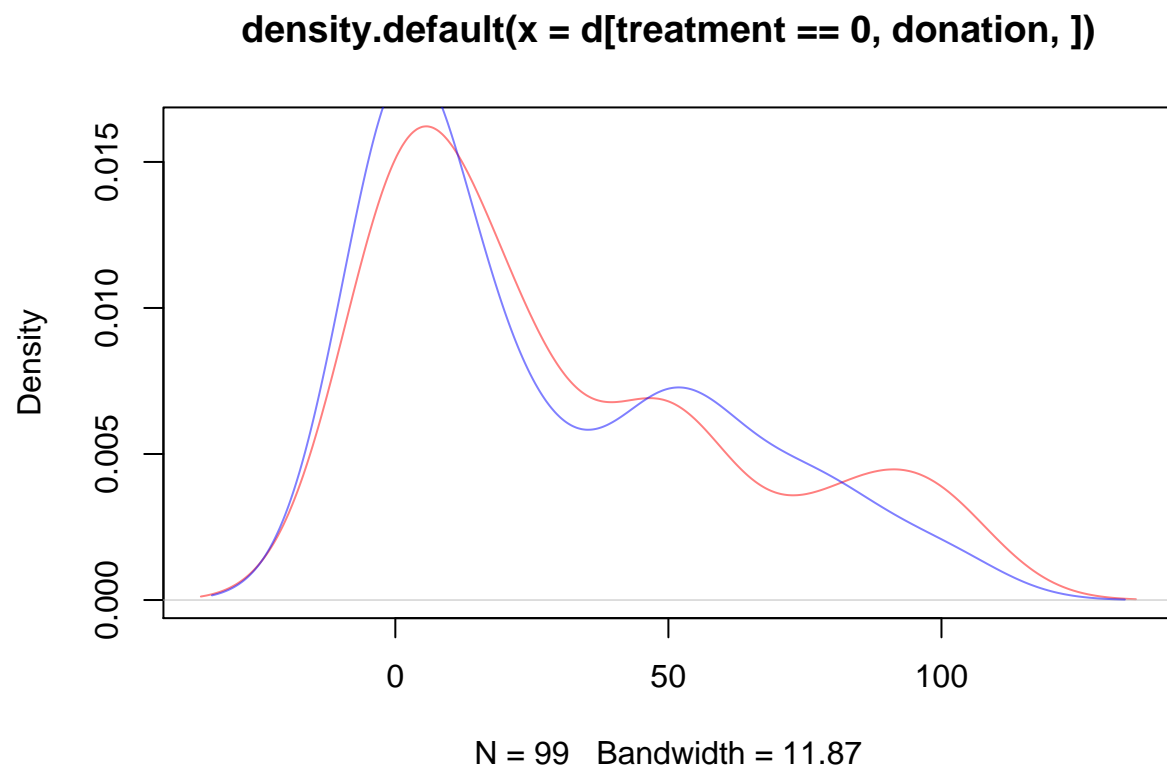


```
mod <- lm(donation ~ treatment, data=d)
summary(mod)

##
## Call:
## lm(formula = donation ~ treatment, data = d)
##
## Residuals:
##      Min       1Q   Median       3Q      Max
## -30.74  -27.16  -11.45   22.84   72.84
##
## Coefficients:
##              Estimate Std. Error t value Pr(>|t|)
## (Intercept)   30.737     3.224   9.533  <2e-16 ***
## treatment     -3.583     4.583  -0.782   0.435
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 32.08 on 194 degrees of freedom
## Multiple R-squared:  0.00314,    Adjusted R-squared:  -0.001998
## F-statistic: 0.6111 on 1 and 194 DF,  p-value: 0.4353

d1 <- density(d[treatment == 0, donation,])
d2 <- density(d[treatment == 1, donation,])

plot(d1, col=rgb(1,0,0,0.5))
lines(d2, col=rgb(0,0,1,0.5))
```



Randomization Inference

Testing the sharp null hypothesis that the treatment has no effect for anyone.

#TODO