

BestProject

May 10, 2023

```
[1]: #Cleaning the Dataset

# 1: begin by reading in the WLS dataset
WLS <- read.csv("WLS.csv")

# and have a look at it
summary(WLS)

# 2: delete all missing values. Look at the WLS Guide to see what indicates a
↳missing value
# Look at R Assignment for to see how to set values to NA

WLS[WLS < 0] <- NA

# 3: delete unreported values from the death-year vector; again, look at the
↳guide
# this will ONLY apply to the WLS$deatyr vector

WLS$deatyr[WLS$deatyr > 9000] <- NA

# 4: adjust income levels to 2020 and put them in thousands
# this will ONLY apply to the WLS$pi5760 and WLS$yrer74 vectors
WLS$pi5760 <- WLS$pi5760 * (10.16/10)
WLS$yrer74 <- WLS$yrer74 * (7.71/10)

# 5: reverse self-reported health vectors.
# suggestion: make them all go low-to-high

WLS$mx001rer <- 6 - WLS$mx001rer

# 6: some ridiculous outliers were entered into the socializing datasets
# no one can socialize 300 times in four weeks
# come up with a reasonable limit for these

WLS$q1z023rer[WLS$q1z023rer > 100] <- 100
WLS$z023rer[WLS$iz023rer > 100] <- 100
WLS$iz023rer[WLS$iz023rer > 100] <- 100
```

```
WLS$mz023rer[WLS$mz023rer > 100] <- 100
```

```
# 7: OPTIONAL BUT RECOMMENDED
```

```
# change the names of the vectors to something REASONABLE
```

```
# best to keep it short and meaningful
```

```
names(WLS) <- c("ID", "Birth", "Gender", "Death", "Income",  
               "Population", "Income75", "Health1", "Sad1", "Social1",  
               "Health2", "Sad2", "Social2", "Cog2", "Health3",  
               "Sad3", "Social3", "Cog3", "Health4", "Sad4",  
               "Social4", "Cog4", "Marriage", "Children", "Degree")
```

```
# check to see if you did it!
```

```
summary(WLS)
```

idpub	brdxdy	sexrsp	deatyr
Min. :900018	Min. : -3.00	Min. :1.000	Min. : -2.0
1st Qu.:908321	1st Qu.:39.00	1st Qu.:1.000	1st Qu.: -2.0
Median :916867	Median :39.00	Median :2.000	Median : -2.0
Mean :916879	Mean :36.47	Mean :1.516	Mean : 751.7
3rd Qu.:925388	3rd Qu.:39.00	3rd Qu.:2.000	3rd Qu.:2002.0
Max. :933957	Max. :40.00	Max. :2.000	Max. :9996.0

pi5760	pop57	yrer74	mx001rer
Min. : -3.00	Min. :2.000	Min. : -3.00	Min. : -3.000
1st Qu.: 24.00	1st Qu.:3.000	1st Qu.: 0.00	1st Qu.: 4.000
Median : 49.00	Median :5.000	Median : 50.00	Median : 4.000
Mean : 54.18	Mean :5.037	Mean : 78.04	Mean : 4.138
3rd Qu.: 71.00	3rd Qu.:6.000	3rd Qu.:130.00	3rd Qu.: 5.000
Max. :998.00	Max. :9.000	Max. :1650.00	Max. : 5.000
			NA's :3442

ru024rec	mz023rer	gx201re	gu024re
Min. : -3.0000	Min. : -3.000	Min. : -3.000	Min. : -5.0000
1st Qu.: -2.0000	1st Qu.: 1.000	1st Qu.: 2.000	1st Qu.: -2.0000
Median : 0.0000	Median : 3.000	Median : 2.000	Median : 0.0000
Mean : -0.0022	Mean : 3.906	Mean : 2.259	Mean : -0.1027
3rd Qu.: 0.0000	3rd Qu.: 5.000	3rd Qu.: 3.000	3rd Qu.: 0.0000
Max. : 7.0000	Max. :44.000	Max. : 5.000	Max. : 7.0000
NA's :1824	NA's :3442	NA's :2585	NA's :3052

iz023rer	gi503re	hx201re	hu024re
Min. : -3.000	Min. : -5.000	Min. : -30.000	Min. : -30.000
1st Qu.: 1.000	1st Qu.: -2.000	1st Qu.: 2.000	1st Qu.: -2.000
Median : 2.000	Median : 6.000	Median : 2.000	Median : -2.000
Mean : 3.363	Mean : 4.667	Mean : 1.409	Mean : -2.297
3rd Qu.: 4.000	3rd Qu.: 9.000	3rd Qu.: 3.000	3rd Qu.: -2.000
Max. :28.000	Max. :12.000	Max. : 5.000	Max. : 7.000

NA's :3472	NA's :3052	NA's :4165	NA's :4165
jz023rer	hi503rea	q1x001rer	q1u024re
Min. : -29.000	Min. : -30.000	Min. : -3.000	Min. : -2.000
1st Qu.: 1.000	1st Qu.: -2.000	1st Qu.: 3.000	1st Qu.: -2.000
Median : 2.000	Median : 6.000	Median : 4.000	Median : -2.000
Mean : 0.794	Mean : 3.616	Mean : 3.876	Mean : -0.996
3rd Qu.: 5.000	3rd Qu.: 8.000	3rd Qu.: 4.000	3rd Qu.: -2.000
Max. : 300.000	Max. : 12.000	Max. : 5.000	Max. : 7.000
NA's :4926	NA's :4165	NA's :6969	NA's :6969
q1z023rer	q1i503rea	q1c003re	hd201kd
Min. : -3.000	Min. : -5.000	Min. : -5.000	Min. : 0.00
1st Qu.: 0.000	1st Qu.: 4.000	1st Qu.: 1.000	1st Qu.: 2.00
Median : 2.000	Median : 6.000	Median : 1.000	Median : 3.00
Mean : 4.135	Mean : 5.381	Mean : 1.187	Mean : 3.24
3rd Qu.: 5.000	3rd Qu.: 8.000	3rd Qu.: 1.000	3rd Qu.: 4.00
Max. : 150.000	Max. : 12.000	Max. : 5.000	Max. : 10.00
NA's :6969	NA's :6969	NA's :6969	NA's :4165
dglev			
Min. : -3.000			
1st Qu.: -2.000			
Median : -2.000			
Mean : -1.078			
3rd Qu.: -2.000			
Max. : 4.000			

ID	Birth	Gender	Death
Min. :900018	Min. :37.00	Min. :1.000	Min. :1957
1st Qu.:908321	1st Qu.:39.00	1st Qu.:1.000	1st Qu.:1999
Median :916867	Median :39.00	Median :2.000	Median :2009
Mean :916879	Mean :38.84	Mean :1.516	Mean :2005
3rd Qu.:925388	3rd Qu.:39.00	3rd Qu.:2.000	3rd Qu.:2015
Max. :933957	Max. :40.00	Max. :2.000	Max. :2019
	NA's :586		NA's :6747
Income	Population	Income75	Health1
Min. : 0.00	Min. :2.000	Min. : 0.00	Min. :1.000
1st Qu.: 34.54	1st Qu.:3.000	1st Qu.: 0.00	1st Qu.:1.000
Median : 54.86	Median :5.000	Median : 55.51	Median :2.000
Mean : 63.76	Mean :5.037	Mean : 68.42	Mean :1.849
3rd Qu.: 75.18	3rd Qu.:6.000	3rd Qu.: 107.94	3rd Qu.:2.000
Max. :1013.97	Max. :9.000	Max. :1272.15	Max. :5.000
NA's :1346		NA's :1204	NA's :3455
Sad1	Social1	Health2	Sad2
Min. :0.000	Min. : 0.000	Min. :1.000	Min. :0.000
1st Qu.:0.000	1st Qu.: 1.000	1st Qu.:2.000	1st Qu.:0.000
Median :0.000	Median : 3.000	Median :2.000	Median :0.000
Mean :0.831	Mean : 3.945	Mean :2.268	Mean :0.752
3rd Qu.:0.000	3rd Qu.: 5.000	3rd Qu.:3.000	3rd Qu.:0.000

Max. :7.000	Max. :44.000	Max. :5.000	Max. :7.000
NA's :4269	NA's :3481	NA's :2598	NA's :5131
Social2	Cog2	Health3	Sad3
Min. : 0.000	Min. : 0.000	Min. :1.000	Min. :0.000
1st Qu.: 1.000	1st Qu.: 6.000	1st Qu.:2.000	1st Qu.:3.000
Median : 3.000	Median : 7.000	Median :2.000	Median :4.000
Mean : 3.665	Mean : 7.235	Mean :2.378	Mean :4.124
3rd Qu.: 4.000	3rd Qu.:10.000	3rd Qu.:3.000	3rd Qu.:5.000
Max. :28.000	Max. :12.000	Max. :5.000	Max. :7.000
NA's :3784	NA's :4988	NA's :4350	NA's :9680
Social3	Cog3	Health4	Sad4
Min. : 0.000	Min. : 0.000	Min. :1.000	Min. :0.000
1st Qu.: 1.000	1st Qu.: 5.000	1st Qu.:3.000	1st Qu.:2.000
Median : 3.000	Median : 6.000	Median :4.000	Median :4.000
Mean : 4.332	Mean : 6.754	Mean :3.891	Mean :3.622
3rd Qu.: 5.000	3rd Qu.: 9.000	3rd Qu.:4.000	3rd Qu.:5.000
Max. :300.000	Max. :12.000	Max. :5.000	Max. :7.000
NA's :5504	NA's :5719	NA's :6978	NA's :9719
Social4	Cog4	Marriage	Children
Min. : 0.000	Min. : 0.000	Min. :0.000	Min. : 0.00
1st Qu.: 0.000	1st Qu.: 4.000	1st Qu.:1.000	1st Qu.: 2.00
Median : 2.000	Median : 6.000	Median :1.000	Median : 3.00
Mean : 4.119	Mean : 5.458	Mean :1.221	Mean : 3.24
3rd Qu.: 5.000	3rd Qu.: 8.000	3rd Qu.:1.000	3rd Qu.: 4.00
Max. :100.000	Max. :12.000	Max. :5.000	Max. :10.00
NA's :6979	NA's :7002	NA's :6988	NA's :4165
Degree	NA		
Min. :1.000	Min. : NA		
1st Qu.:2.000	1st Qu.: NA		
Median :2.000	Median : NA		
Mean :2.345	Mean :NaN		
3rd Qu.:3.000	3rd Qu.: NA		
Max. :4.000	Max. : NA		
NA's :7965	NA's :10317		

```
[6]: install.packages("Hmisc")
library(Hmisc)

hist.data.frame(WLS)
```

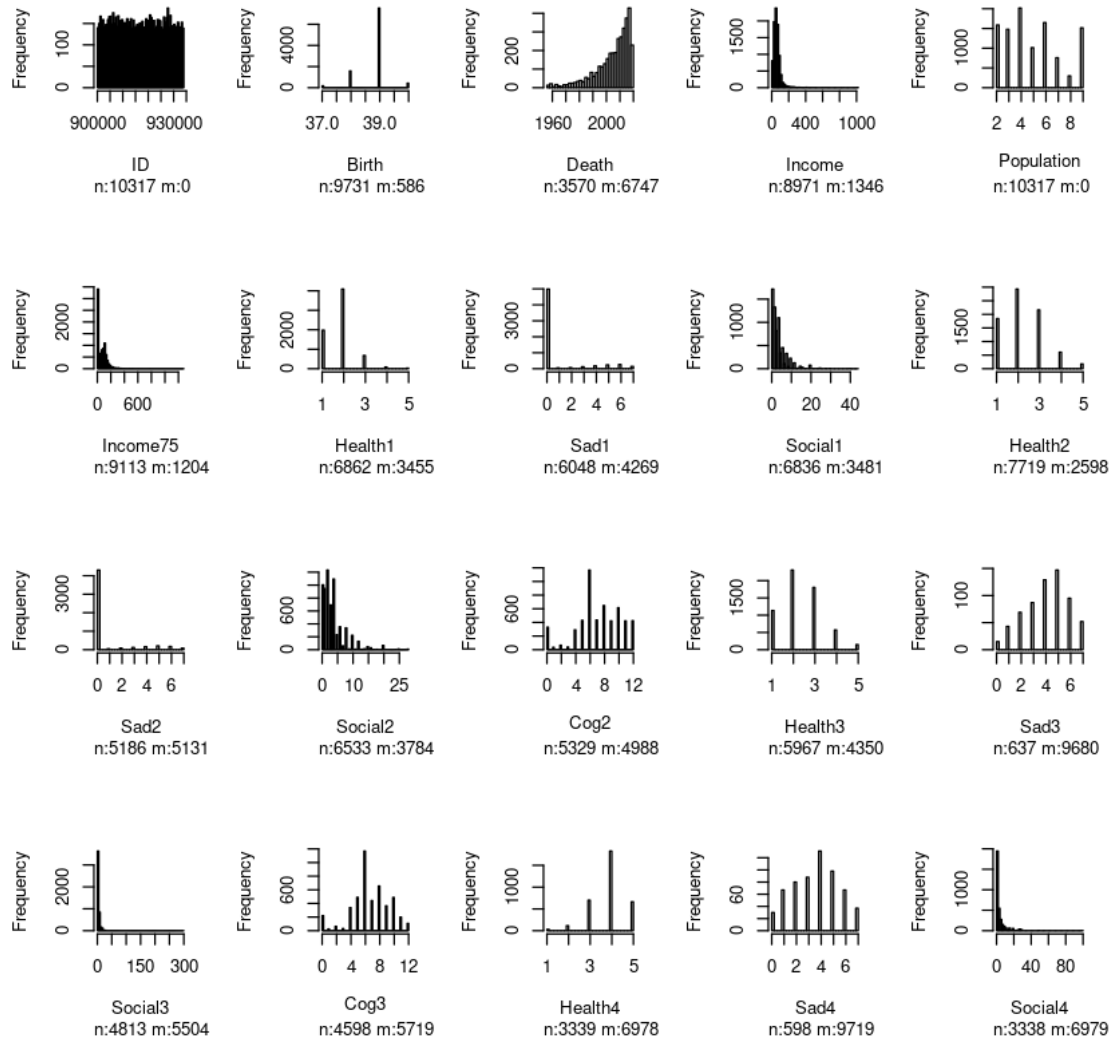
Installing package into '/opt/r'
(as 'lib' is unspecified)

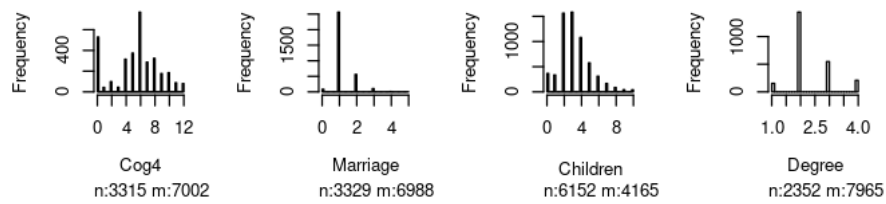
also installing the dependencies 'data.table', 'htmlTable', 'viridis', 'Formula'

Attaching package: 'Hmisc'

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

format.pval, units





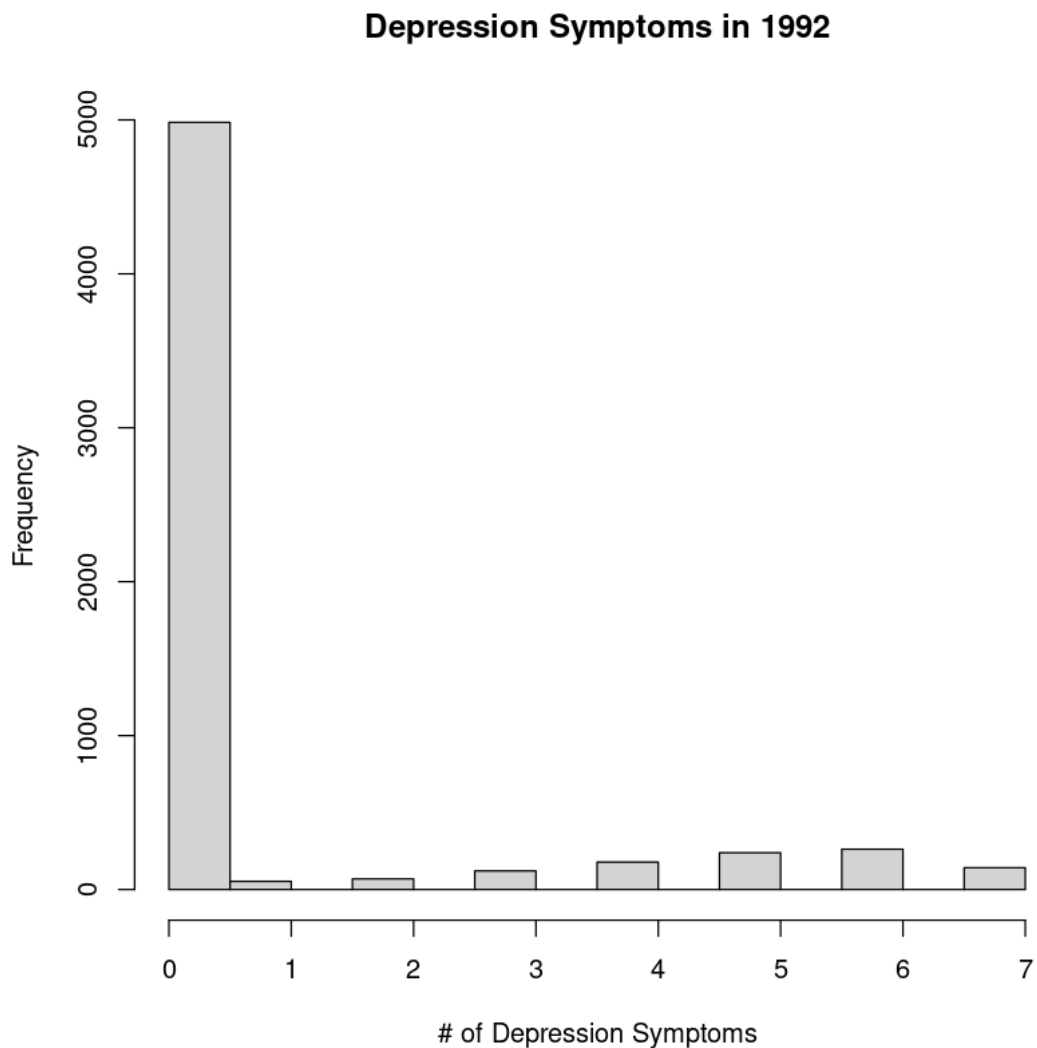
```
[ ]: #Interesting Comparison #1
```

Depression Symptoms with year 2020 (year of Covid). Also substed between male
 ↳ and female and create a hypothesis test if
 there is a signifincant amount of difference to say 2020 had a impact on
 ↳ the number of depression symptoms.

```
[61]: #Histogram Sad1 Dataset
```

```
hist(WLS$Sad1, # dataframe$vector
      breaks = 25, # breaks sets the approximate number of bars
      main = "Depression Symptoms in 1992", # graph title
      xlab = "# of Depression Symptoms", # x-axis title
```

```
ylab = "Frequency" # y-axis title
)
```



```
[11]: #Histogram Sad2 Dataset
```

```
hist(WLS$Sad2, # dataframe$vector
      breaks = 7, # breaks sets the approximate number of bars
      main = "Depression Symptoms in 2003", # graph title
      xlab = "# of Depression Symptoms", # x-axis title
      ylab = "Frequency",
      col = "gold" # y-axis title
    )
```

```

segments(0.751,0, # x and y coordinates for one end of the line segment
         0.751,4350, # x and y coordinates for the other end. Note that the x
         ↪value stays the same--a vertical line
         lwd = 3, # line width--make the line three times as thick for clarity
         col = "blue" # a favorite color of mine. YOU CAN'T USE IT
         )

text(1.3,4000, # x and y coordinates of where the text goes
     labels = "Mean", # what to print there
     cex = 1, # cex means Character Expansion, and can make the characters
     ↪larger or smaller. 1 is the default
     col = "blue"
     )

text(1.3,3800, # same x but y slightly lower
     labels = "4.124",
     cex = 1,
     col = "blue"
     )

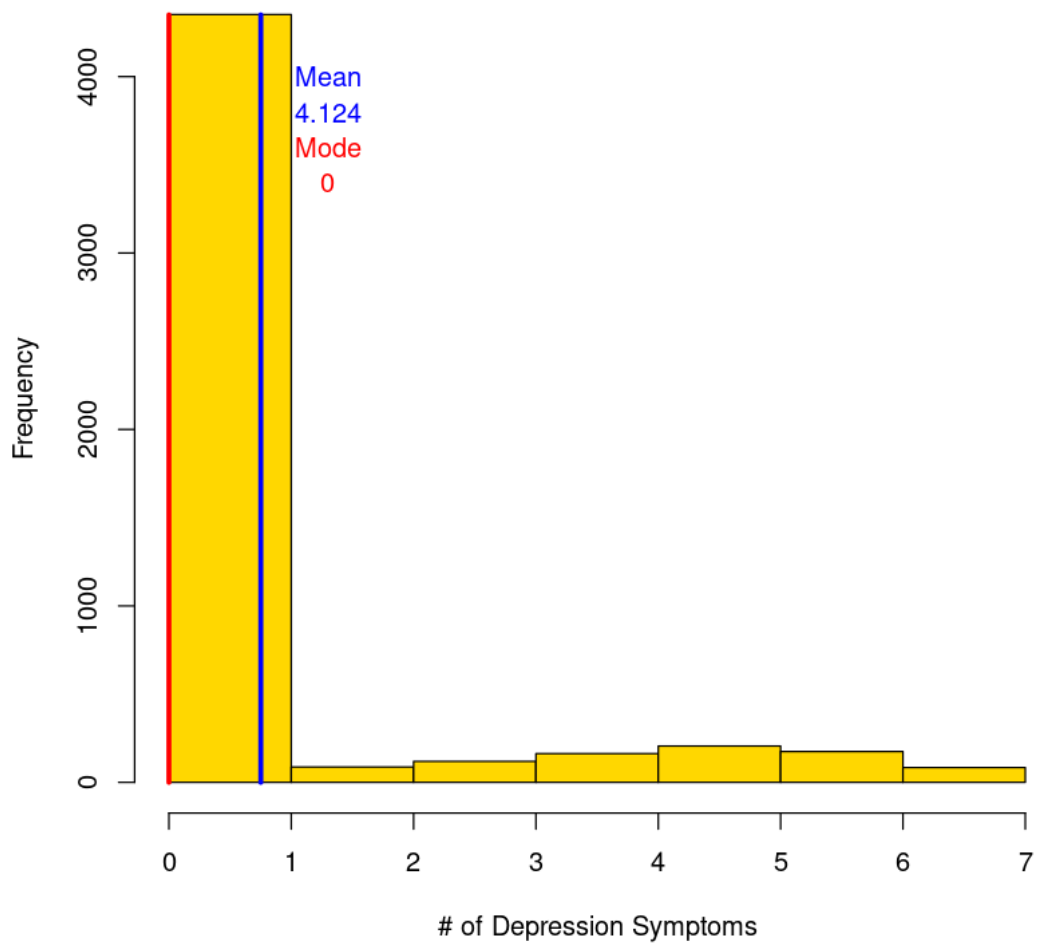
segments(0,0, # x and y coordinates for one end of the line segment
         0,4350, # x and y coordinates for the other end. Note that the x
         ↪value stays the same--a vertical line
         lwd = 3, # line width--make the line three times as thick for clarity
         col = "red" # a favorite color of mine. YOU CAN'T USE IT
         )

text(1.3,3600, # x and y coordinates of where the text goes
     labels = "Mode", # what to print there
     cex = 1, # cex means Character Expansion, and can make the characters
     ↪larger or smaller. 1 is the default
     col = "red"
     )

text(1.3,3400, # same x but y slightly lower
     labels = "0",
     cex = 1,
     col = "red"
     )

```


Depression Symptoms in 2003



[17]: *#Histograms Sad3 Dataset*

```
hist(WLS$Sad3, # dataframe$vector
      breaks = 7, # breaks sets the approximate number of bars
      main = "Depression Symptoms in 2011", # graph title
      xlab = "# of Depression Symptoms", # x-axis title
      ylab = "Frequency",
      col = "lightcoral" # y-axis title
)

segments(4.124,0, # x and y coordinates for one end of the line segment
         4.124,155, # x and y coordinates for the other end. Note that the x
         ↪ value stays the same--a vertical line
```

```

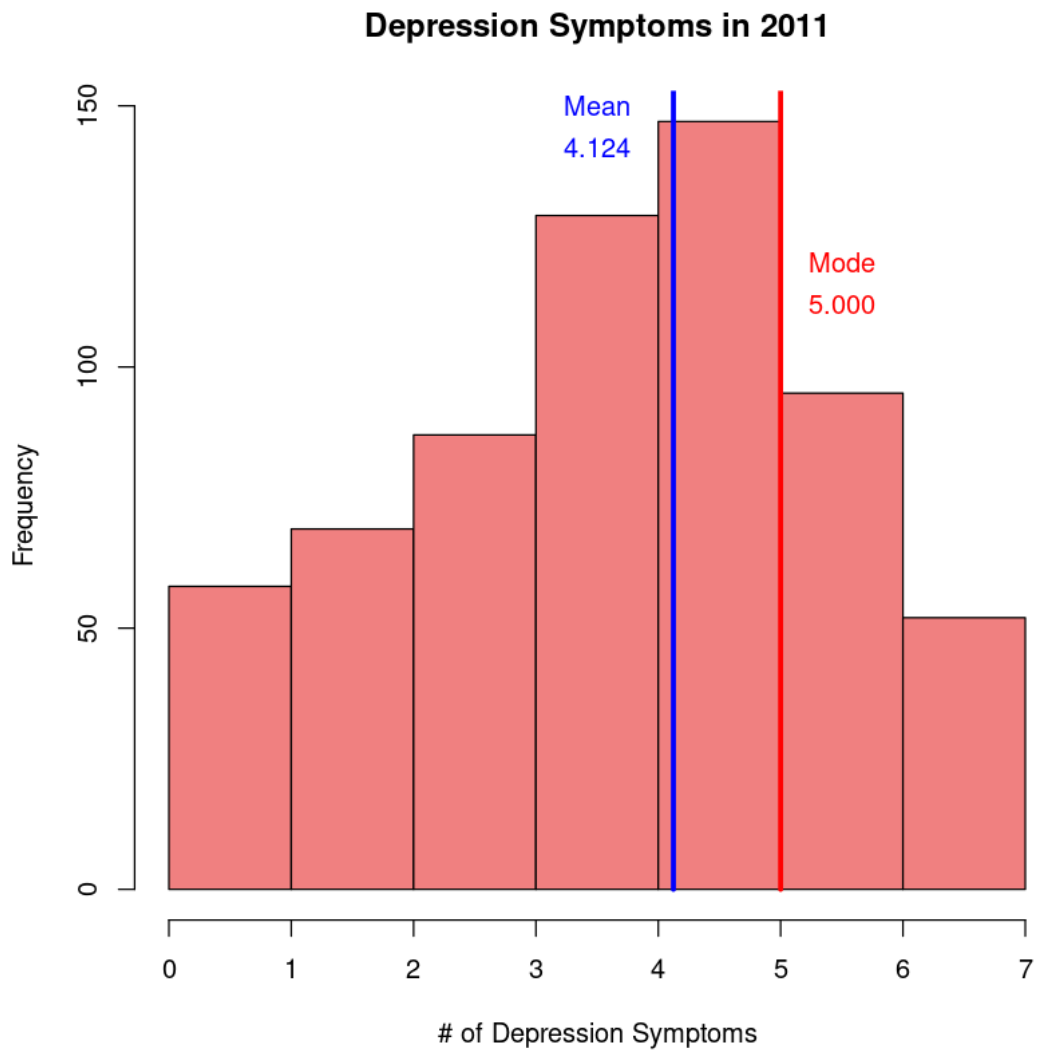
        lwd = 3, # line width--make the line three times as thick for clarity
        col = "blue" # a favorite color of mine. YOU CAN'T USE IT
    )

text(3.5,150, # x and y coordinates of where the text goes
     labels = "Mean", # what to print there
     cex = 1, # cex means Character Expansion, and can make the characters
     ↪larger or smaller. 1 is the default
     col = "blue"
)
text(3.5,142, # same x but y slightly lower
     labels = "4.124",
     cex = 1,
     col = "blue"
)

segments(5,0, # x and y coordinates for one end of the line segment
         5,155, # x and y coordinates for the other end. Note that the x value
         ↪stays the same--a vertical line
         lwd = 3, # line width--make the line three times as thick for clarity
         col = "red" # a favorite color of mine. YOU CAN'T USE IT
)

text(5.5, 120, # x and y coordinates of where the text goes
     labels = "Mode", # what to print there
     cex = 1, # cex means Character Expansion, and can make the characters
     ↪larger or smaller. 1 is the default
     col = "red"
)
text(5.5,112, # same x but y slightly lower
     labels = "5.000",
     cex = 1,
     col = "red"
)

```



[4]: *#Histogram for Depression Symptoms in 2020*

```
hist(WLS$Sad4, # dataframe$vector
      breaks = 7, # breaks sets the approximate number of bars
      main = "Depression Symptoms in 2020", # graph title
      xlab = "# of Depression Symptoms", # x-axis title
      ylab = "Frequency",
      col = "lightblue"
    )

segments(3.620,0, # x and y coordinates for one end of the line segment
```

```

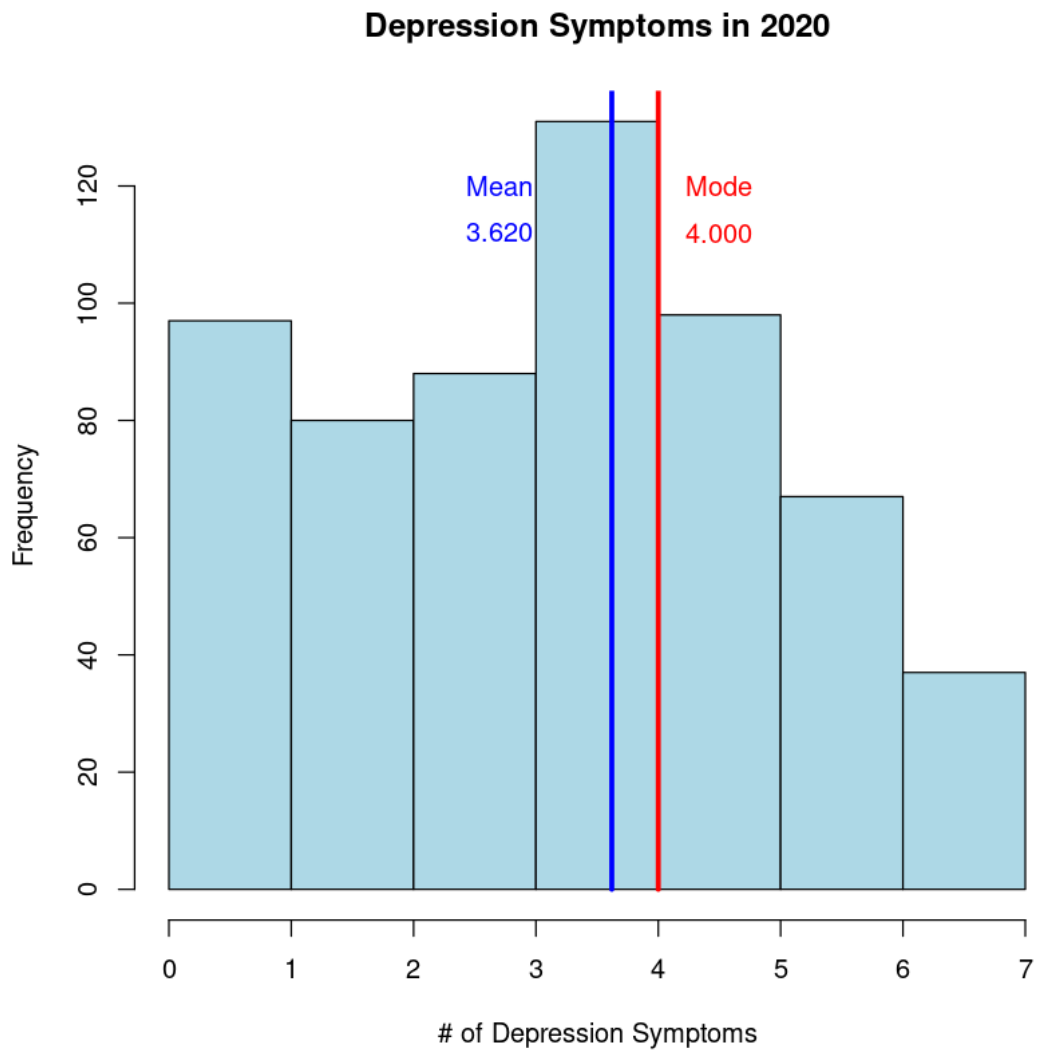
    3.620,155, # x and y coordinates for the other end. Note that the x
    ↪value stays the same--a vertical line
    lwd = 3, # line width--make the line three times as thick for clarity
    col = "blue" # a favorite color of mine. YOU CAN'T USE IT
  )

text(2.7,120, # x and y coordinates of where the text goes
     labels = "Mean", # what to print there
     cex = 1, # cex means Character Expansion, and can make the characters
    ↪larger or smaller. 1 is the default
     col = "blue"
  )
text(2.7,112, # same x but y slightly lower
     labels = "3.620",
     cex = 1,
     col = "blue"
  )

segments(4,0, # x and y coordinates for one end of the line segment
         4,155, # x and y coordinates for the other end. Note that the x value
    ↪stays the same--a vertical line
         lwd = 3, # line width--make the line three times as thick for clarity
         col = "red" # a favorite color of mine. YOU CAN'T USE IT
       )

text(4.5, 120, # x and y coordinates of where the text goes
     labels = "Mode", # what to print there
     cex = 1, # cex means Character Expansion, and can make the characters
    ↪larger or smaller. 1 is the default
     col = "red"
  )
text(4.5,112, # same x but y slightly lower
     labels = "4.000",
     cex = 1,
     col = "red"
  )

```



[15]: *#Significant difference in means between 2003, 2011, and 2020 bulk samples*

```
var.test(WLS$Sad3, WLS$Sad4)
```

```
t.test(WLS$Sad3, WLS$Sad4,  
       conf.level = 0.99,  
       alternative = "g",  
       var.equal = TRUE)
```

```
prop.test(x = c(4304, 15),  
         n = c(5186, 637),  
         conf.level = 0.99,  
         alternative = "t")
```

```

    )

prop.test(x = c(4304, 15),
          n = c(5186, 637),
          conf.level = 0.99,
          alternative = "g"
        )

zerosad2 <- subset(WLS, Sad2 < 1)
zerosad3 <- subset(WLS, Sad3 < 1)

zerosad22 <- subset(WLS, Sad2 < 8)
zerosad33 <- subset(WLS, Sad3 < 8)

nrow(zerosad2)
nrow(zerosad3)

nrow(zerosad22)
nrow(zerosad33)

zerosad5 <- subset(WLS, Sad3 < 1)
zerosad6 <- subset(WLS, Sad4 < 1)

zerosad55 <- subset(WLS, Sad3 < 8)
zerosad66 <- subset(WLS, Sad4 < 8)

nrow(zerosad5)
nrow(zerosad6)

nrow(zerosad55)
nrow(zerosad66)

#Results: There is not a significant different in their variances, but there is
↪ a significant different in their means. (Very significant)

```

F test to compare two variances

```

data:  WLS$Sad3 and WLS$Sad4
F = 0.88097, num df = 636, denom df = 597, p-value = 0.1157
alternative hypothesis: true ratio of variances is not equal to 1
95 percent confidence interval:
 0.7519912 1.0316598
sample estimates:

```

ratio of variances
0.880971

Two Sample t-test

data: WLS\$Sad3 and WLS\$Sad4
t = 4.8462, df = 1233, p-value = 7.091e-07
alternative hypothesis: true difference in means is greater than 0
99 percent confidence interval:
0.2606826 Inf
sample estimates:
mean of x mean of y
4.124019 3.622074

2-sample test for equality of proportions with continuity correction

data: c(4304, 15) out of c(5186, 637)
X-squared = 1921.4, df = 1, p-value < 2.2e-16
alternative hypothesis: two.sided
99 percent confidence interval:
0.7850017 0.8277560
sample estimates:
prop 1 prop 2
0.82992673 0.02354788

2-sample test for equality of proportions with continuity correction

data: c(4304, 15) out of c(5186, 637)
X-squared = 1921.4, df = 1, p-value < 2.2e-16
alternative hypothesis: greater
99 percent confidence interval:
0.7869868 1.0000000
sample estimates:
prop 1 prop 2
0.82992673 0.02354788

4304

15

5186

637

15

30

637

598

```
[34]: prop.test(x = c(15, 30),
               n = c(637, 598),
               conf.level = 0.99,
               alternative = "t"
               )

prop.test(x = c(15, 30),
          n = c(637, 598),
          conf.level = 0.99,
          alternative = "l"
          )

zerosad5 <- subset(WLS, Sad3 < 1)
zerosad6 <- subset(WLS, Sad4 < 1)

zerosad55 <- subset(WLS, Sad3 < 8)
zerosad66 <- subset(WLS, Sad4 < 8)

nrow(zerosad5)
nrow(zerosad6)

nrow(zerosad55)
nrow(zerosad66)
```

2-sample test for equality of proportions with continuity correction

```
data:  c(15, 30) out of c(637, 598)
X-squared = 5.49, df = 1, p-value = 0.01913
alternative hypothesis: two.sided
99 percent confidence interval:
 -0.055956534  0.002717848
sample estimates:
      prop 1      prop 2 
0.02354788 0.05016722
```

2-sample test for equality of proportions with continuity correction

```
data:  c(15, 30) out of c(637, 598)
X-squared = 5.49, df = 1, p-value = 0.009563
alternative hypothesis: less
```


99 percent confidence interval:

-1.000000e+00 3.340637e-05

sample estimates:

prop 1 prop 2
0.02354788 0.05016722

15

30

637

598

```
[8]: #Significant difference in means between 2011 and 2020 in two samples (male and female)
```

```
male <- subset(WLS, Gender == 1)
female <- subset(WLS, Gender == 2)
```

```
var.test(male$Sad4,female$Sad4
)
```

```
t.test(male$Sad4,female$Sad4,
conf.level = 0.99,
var.equal = TRUE)
```

F test to compare two variances

data: male\$Sad4 and female\$Sad4

F = 1.0931, num df = 200, denom df = 396, p-value = 0.4583

alternative hypothesis: true ratio of variances is not equal to 1

95 percent confidence interval:

0.8631481 1.3977221

sample estimates:

ratio of variances
1.093082

Two Sample t-test

data: male\$Sad4 and female\$Sad4

t = 0.45906, df = 596, p-value = 0.6464

alternative hypothesis: true difference in means is not equal to 0

99 percent confidence interval:

-0.3456273 0.4949563

sample estimates:

```
mean of x mean of y
3.671642 3.596977
```

```
[41]: #It seems that depression symptoms increased from the 1900's to around 2011,
      ↪ then decreased significantly during Covid
```

```
mean(WLS$Sad1, na.rm = TRUE)
mean(WLS$Sad2, na.rm = TRUE)
mean(WLS$Sad3, na.rm = TRUE)
mean(WLS$Sad4, na.rm = TRUE)
```

```
#Medians
```

```
median(WLS$Sad1, na.rm = TRUE)
median(WLS$Sad2, na.rm = TRUE)
median(WLS$Sad3, na.rm = TRUE)
median(WLS$Sad4, na.rm = TRUE)
```

```
#Modes
```

```
library(gtools)
```

```
stat_mode(WLS$Sad1)
stat_mode(WLS$Sad2)
stat_mode(WLS$Sad3)
stat_mode(WLS$Sad4)
```

```
0.831183862433862
```

```
0.751831854994215
```

```
4.12401883830455
```

```
3.62207357859532
```

```
[3]: #Does your education level have an effect on the number of depression symptoms?
```

```
summary(aov(                                # summarize ANOVA
          Sad4 ~ factor(Degree),             # the formula. factor() means treat this as a
          ↪ factor, not numeric
          data = WLS                          # the dataframe
          ))
```

```
#This is not concrete evidence that there is no relationship with the
↪ relatively high p-value. There are four factors within the Degree Dataframe.
```

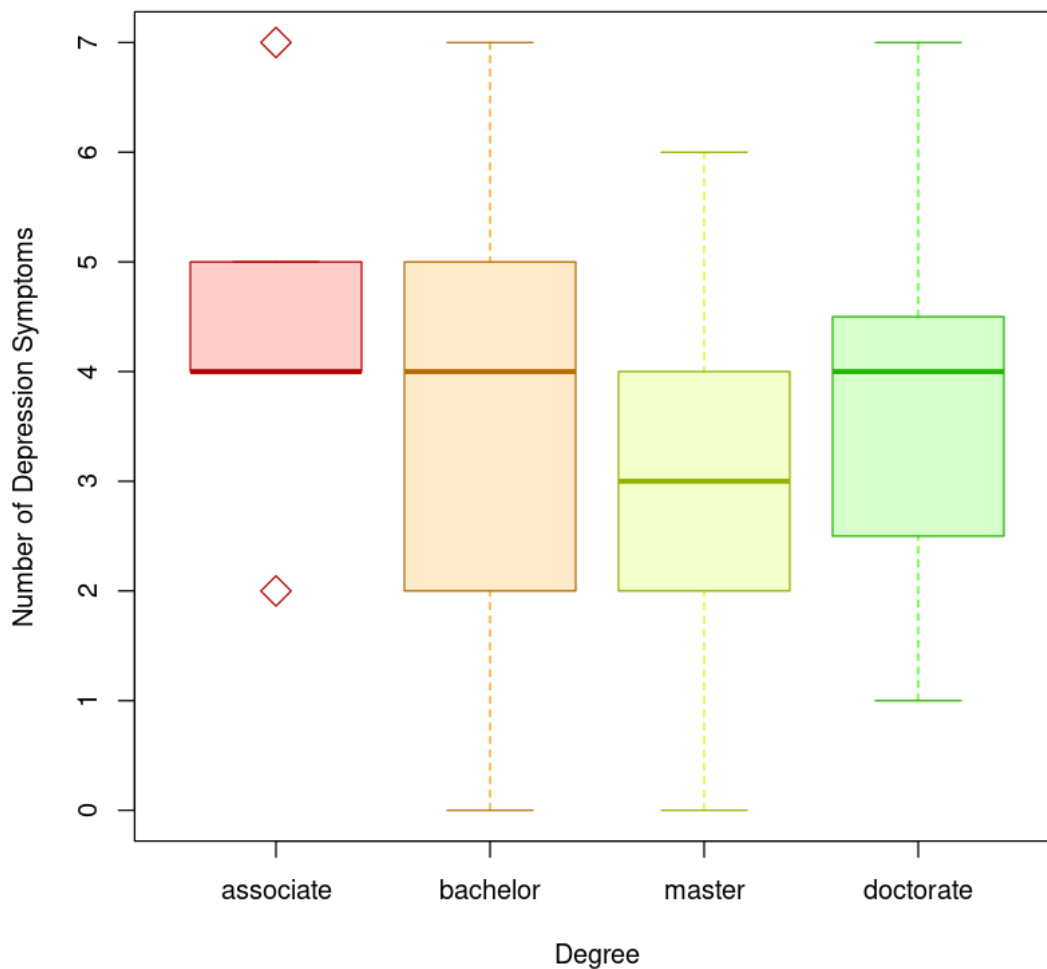
```
              Df Sum Sq Mean Sq F value Pr(>F)
factor(Degree)  3   17.1    5.705    1.661  0.179
```

```
Residuals      119  408.7   3.435
10194 observations deleted due to missingness
```

```
[4]: c1 <- rainbow(10)
      c2 <- rainbow(10, alpha=0.2)
      c3 <- rainbow(10, v=0.7)

      boxplot(Sad4~Degree,
               data = WLS,
               names = c("associate", "bachelor", "master", "doctorate"),
               xlab = "Degree",
               ylab = "Number of Depression Symptoms",
               main = "The Relationship between Depression and Degree",
               col = c2,
               medcol = c3,
               whiskcol = c1,
               staplecol = c3,
               boxcol = c3,
               outcol = c3,
               pch = 23,
               cex = 2)
```

The Relationship between Depression and Degree



```
[2]: #It seems that people with a master's degree have significantly less depression
      ↳ symptoms than the rest of the degrees
```

```
associate <- subset(WLS, Degree == 1)
bachelor <- subset(WLS, Degree == 2)
master <- subset(WLS, Degree == 3)
doctor <- subset(WLS, Degree == 4)
```

```
[33]: sum(!is.na(associate))
      sum(!is.na(bachelor))
      sum(!is.na(master))
      sum(!is.na(doctor))
```

```
jesus <- 2476 + 23825 + 9180 + 3584
```

```
jesus
```

2476

23825

9180

3584

39065

```
[6]: #Conduct a t.test to see if the different between the master mean is
      ↪significantly different from any other mean since the rest seem to have a
      ↪similar mean.

      #Perform t.test master with associate
      t.test(master$Sad4,associate$Sad4,
              conf.level = 0.95,
              var.equal = TRUE)

      #Perform t.test master with bachelor
      t.test(master$Sad4,bachelor$Sad4,
              conf.level = 0.95,
              alternative = "g",
              var.equal = TRUE)

      #Perform t.test master with doctorate
      t.test(master$Sad4,doctor$Sad4,
              conf.level = 0.95,
              var.equal = TRUE)

      t.test(master$Sad4,associate$Sad4,
              conf.level = 0.99,
              alternative = "l",
              var.equal = TRUE)

      #The only significant difference was between masters and associates. People with
      ↪a master's degree have a significantly lower amount of depression symptoms
      ↪than people with only associate's degrees.
```

Two Sample t-test

data: master\$Sad4 and associate\$Sad4

t = -2.1773, df = 34, p-value = 0.03649

```

alternative hypothesis: true difference in means is not equal to 0
95 percent confidence interval:
 -2.8108392 -0.0968531
sample estimates:
mean of x mean of y
 2.846154  4.300000

```

Two Sample t-test

```

data: master$Sad4 and bachelor$Sad4
t = -1.3953, df = 99, p-value = 0.917
alternative hypothesis: true difference in means is greater than 0
95 percent confidence interval:
 -1.329738      Inf
sample estimates:
mean of x mean of y
 2.846154  3.453333

```

Two Sample t-test

```

data: master$Sad4 and doctor$Sad4
t = -1.2466, df = 36, p-value = 0.2206
alternative hypothesis: true difference in means is not equal to 0
95 percent confidence interval:
 -2.1553961  0.5143705
sample estimates:
mean of x mean of y
 2.846154  3.666667

```

Two Sample t-test

```

data: master$Sad4 and associate$Sad4
t = -2.1773, df = 34, p-value = 0.01825
alternative hypothesis: true difference in means is less than 0
99 percent confidence interval:
 -Inf 0.1761856
sample estimates:
mean of x mean of y
 2.846154  4.300000

```

[16]: *#Lets dive deep into the sample of people who have masters and people who have associates and see what other*

```

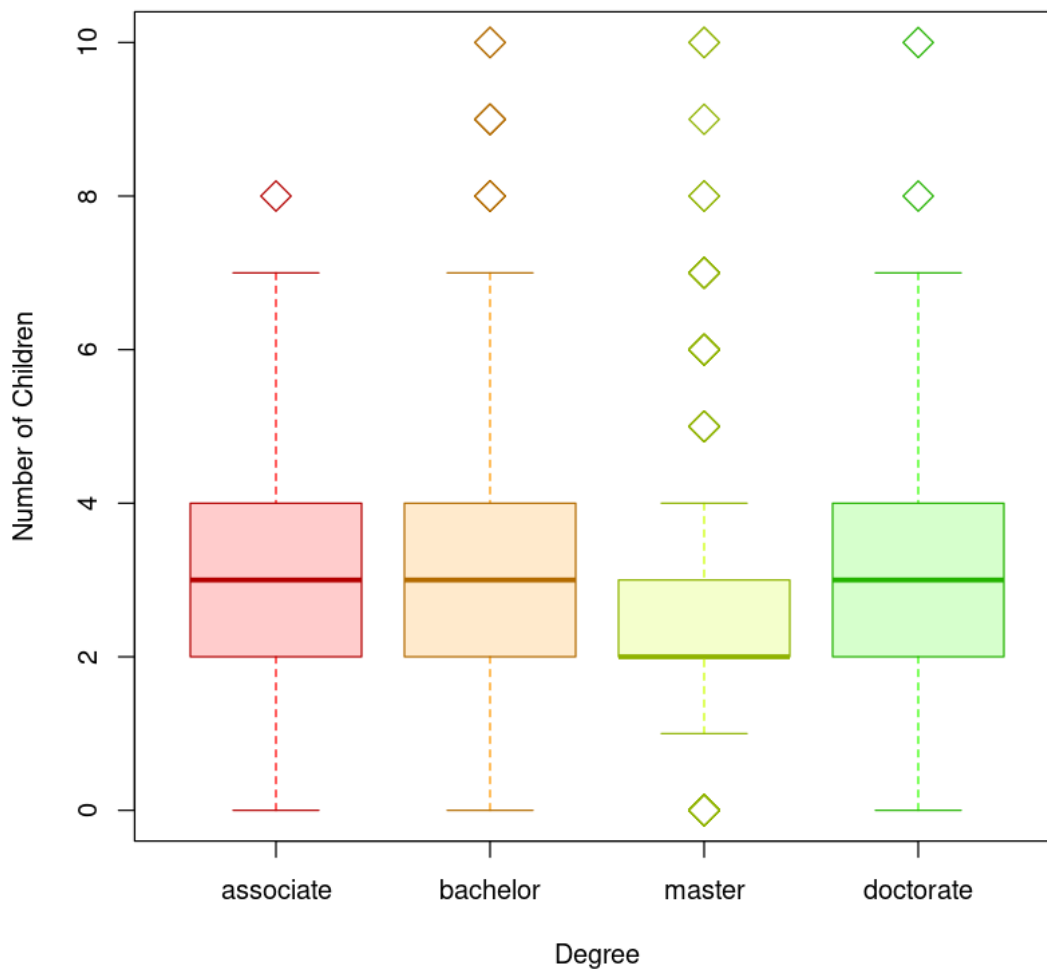
#significantly different factors there are between them. FIRST, LET's SEE IF
↳THE NUMBER OF CHILDREN IS CAUSING
#THIS DIFFERENCE!

c1 <- rainbow(10)
c2 <- rainbow(10, alpha=0.2)
c3 <- rainbow(10, v=0.7)

boxplot(Children~Degree,
        data = WLS,
        names = c("associate", "bachelor", "master", "doctorate"),
        xlab = "Degree",
        ylab = "Number of Children",
        main = "The Relationship between Children and Degree",
        col = c2,
        medcol = c3,
        whiskcol = c1,
        staplecol= c3,
        boxcol = c3,
        outcol = c3,
        pch = 23,
        cex = 2)

```

The Relationship between Children and Degree



[8]: *#Wow, looking at the boxplot only, it seems like people with master's degrees have on average less children!*

```
master1 <- subset(WLS, Degree == 3)
associate1 <- subset(WLS, Degree == 1)

#Perform t.test master with associate
t.test(master1$Children, associate1$Children,
       conf.level = 0.99,
       alternative = "1",
       var.equal = TRUE)
```


Two Sample t-test

```
data: master1$Children and associate1$Children
t = -3.5965, df = 514, p-value = 0.0001768
alternative hypothesis: true difference in means is less than 0
99 percent confidence interval:
    -Inf -0.2424976
sample estimates:
mean of x mean of y
 2.523002  3.213592
```

```
[ ]: #There is an extremely significant difference in means between the number of
      ↳ children with associates and the number of children with masters.
```

```
[8]: #Scatter plot with Sad4/master + Sad4/associate + Children/master + Children/
      ↳ associate
```

```
#Plot the means to emphasize difference in means
```

```
mean(master1$Children, na.rm = TRUE)
mean(associate1$Children, na.rm = TRUE)

mean(master$Sad4, na.rm = TRUE)
mean(associate$Sad4, na.rm = TRUE)
```

2.52300242130751

3.21359223300971

2.84615384615385

4.3

```
[6]: #With the masters subset, plot the Sad4 and Children Against each other
```

```
plot(master$Sad4, master1$Children, # x vector, y vector
      main = "Master's Degree Relationship with Depression and Children",
      xlab = "# of Depression Symptoms for Masters Degree",
      ylab = "# of Children for Masters Degree"
)

boxplot(master$Sad4 ~ master1$Children,
        data = WLS,
        xlab = "# of Depression Symptoms for Masters Degree",
        ylab = "# of Children for Masters Degree",
        main = "Masters Degree Relationship with Deperssion and Children")

plot(associate$Sad4, associate$Children,
```

```

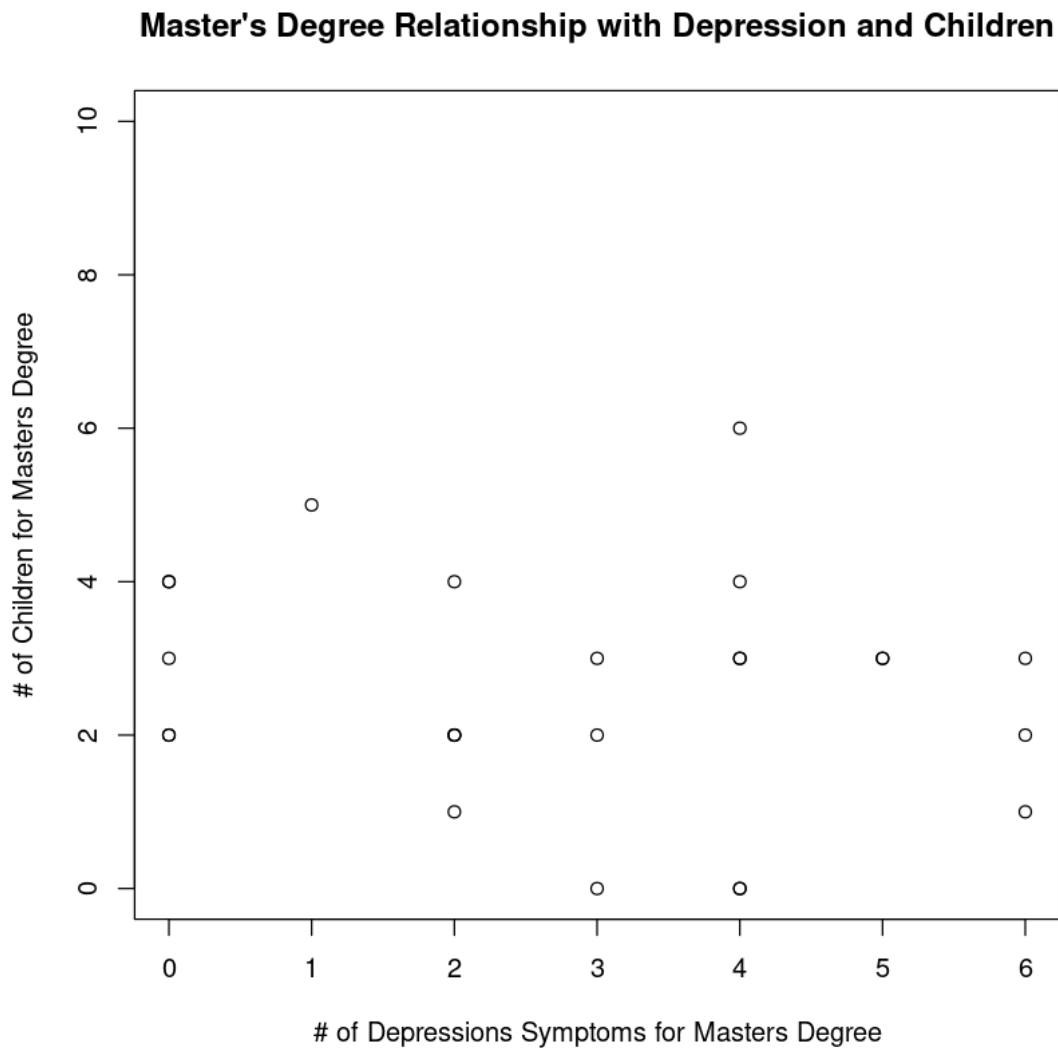
main = "Associate's Degree Relationship with Depression and Children",
xlab = "# of Depression Symptoms for Associates Degree",
ylab = "# of Children for Associates Degree")

```

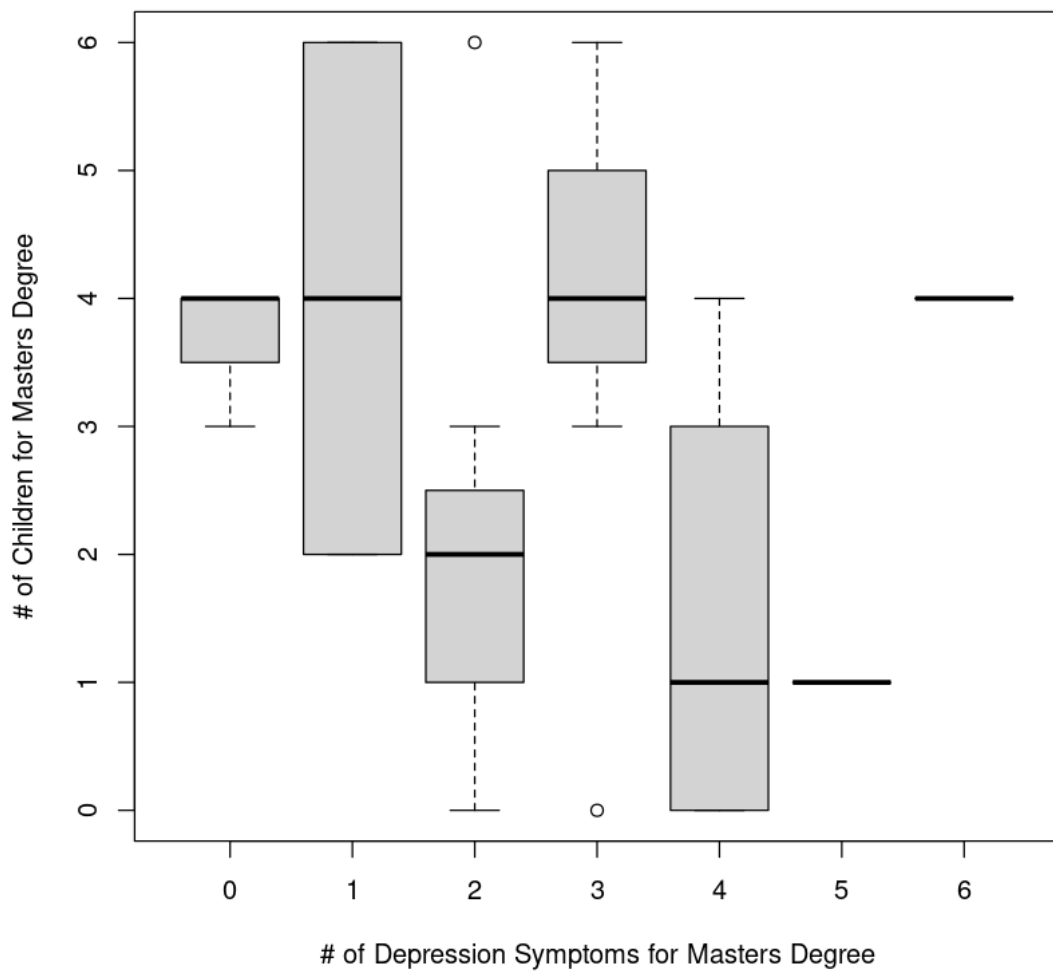
```

boxplot(associate$Sad4~associate1$Children,
  data = WLS,
  xlab = "# of Depression Symptoms for Masters Degree",
  ylab = "# of Children for Masters Degree",
  main = "Masters Degree Relationship with Deperssion and Children")

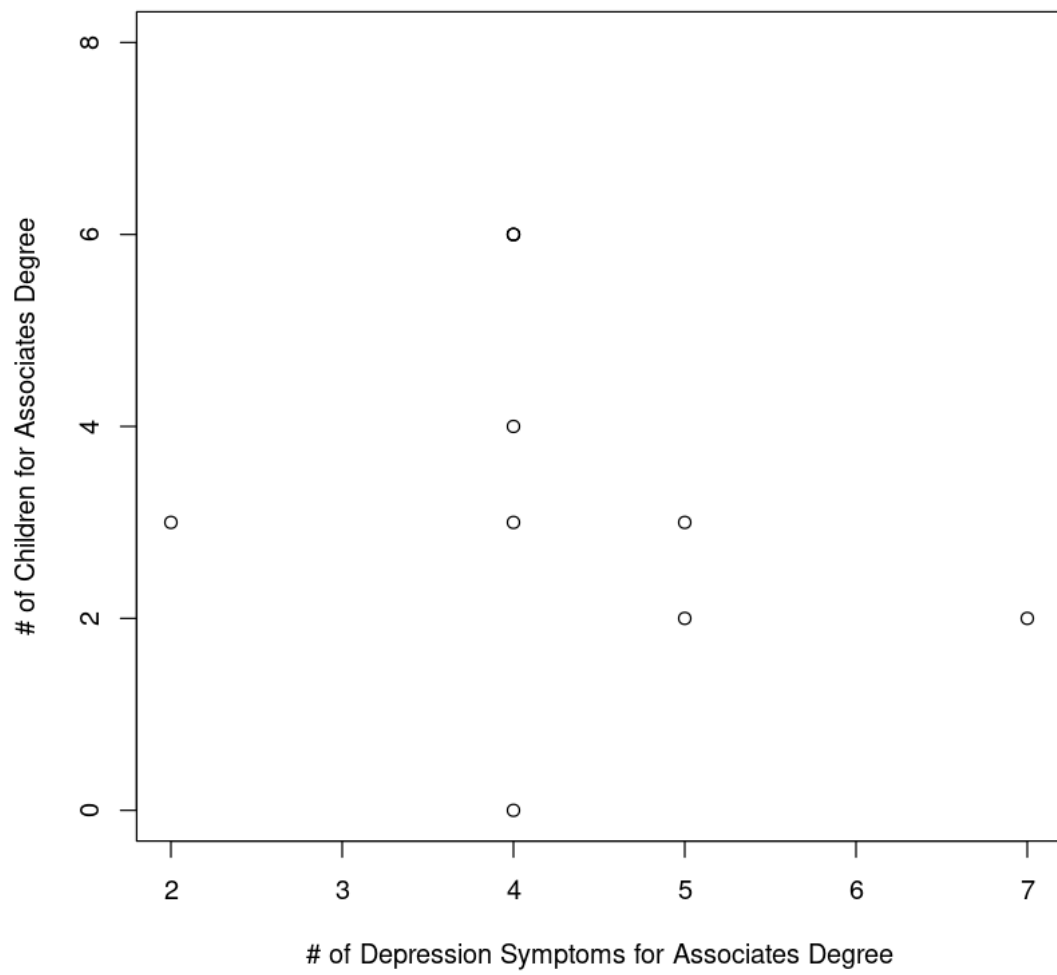
```



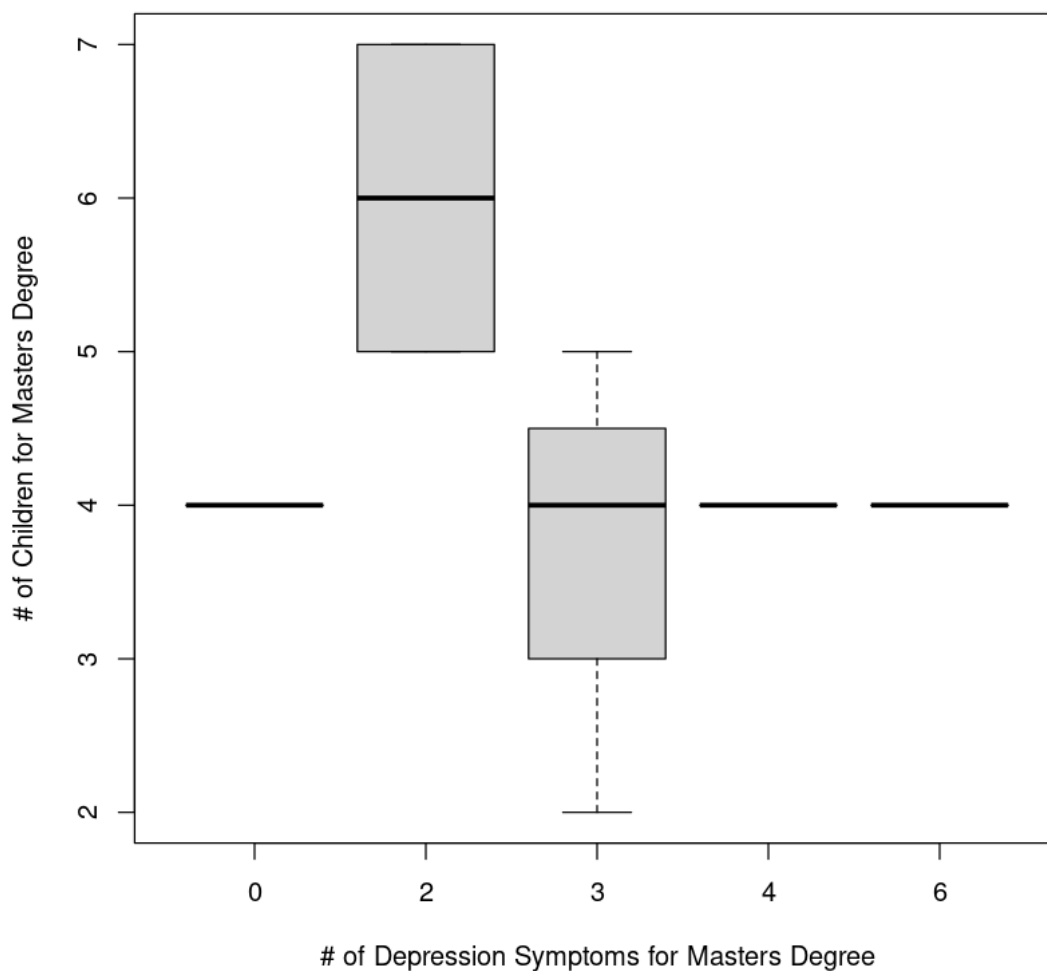
Masters Degree Relationship with Deperssion and Children



Associate's Degree Relationship with Depression and Children



Masters Degree Relationship with Deperssion and Children



[13]: *#Also, you can maybe do that "even though associates had more meeting's with
 ↪ friends they had more depression symptoms
 #than people with masters.*

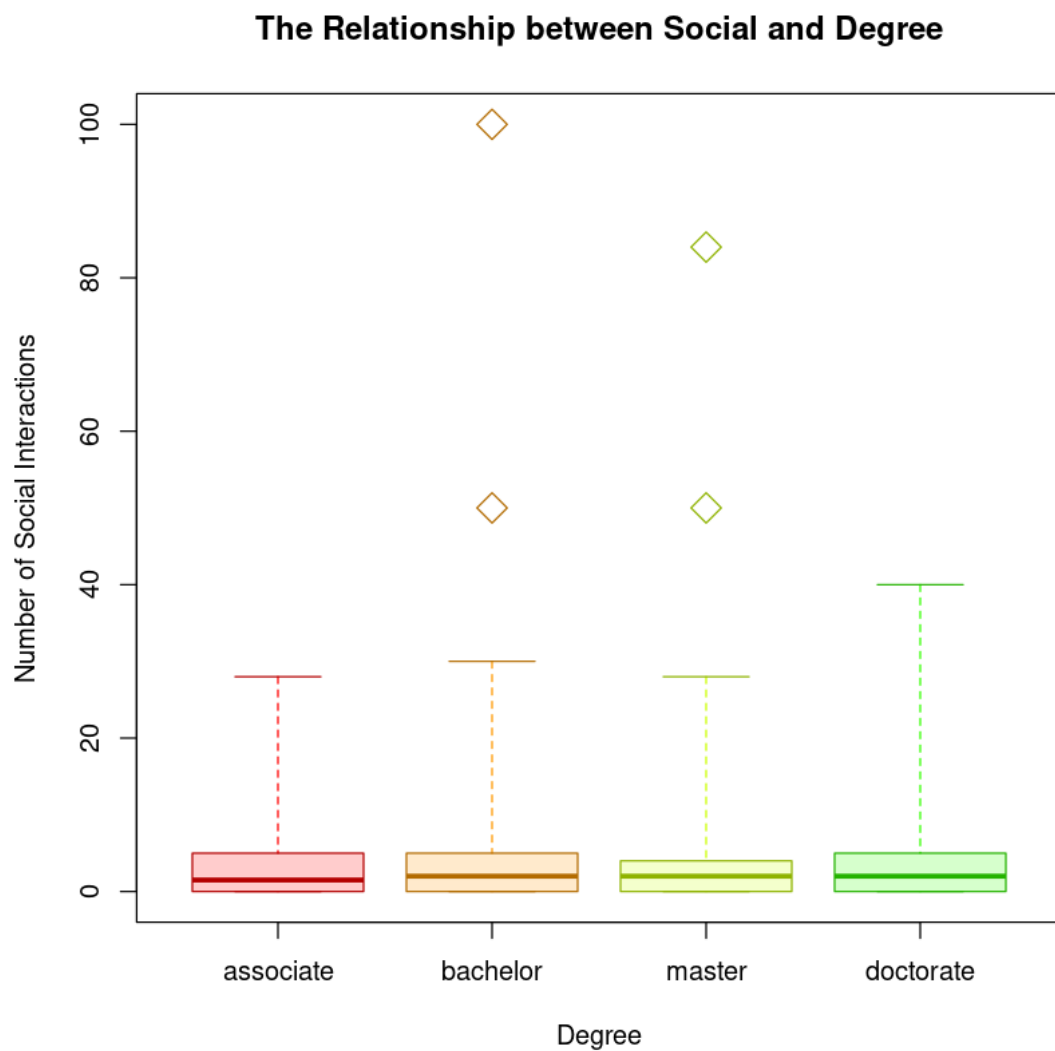
```
c1 <- rainbow(10)
c2 <- rainbow(10, alpha=0.2)
c3 <- rainbow(10, v=0.7)

boxplot(Social4~Degree,
  data = WLS,
  names = c("associate", "bachelor", "master", "doctorate"),
  xlab = "Degree",
  ylab = "Number of Social Interactions",
```

```

main = "The Relationship between Social and Degree",
range = 8,
col = c2,
medcol = c3,
whiskcol = c1,
staplecol= c3,
boxcol = c3,
outcol = c3,
pch = 23,
cex = 2)

```



```
[38]: master2 <- subset(WLS, Degree == 3)
associate2 <- subset(WLS, Degree == 1)

t.test(master2$Social4, associate2$Social4,
        conf.level = 0.99,
        var.equal = TRUE)

var.test(associate$Social, master$Social)
```

Two Sample t-test

```
data: master2$Social4 and associate2$Social4
t = 0.49644, df = 302, p-value = 0.6199
alternative hypothesis: true difference in means is not equal to 0
99 percent confidence interval:
 -2.452257  3.614040
sample estimates:
mean of x mean of y
 4.253968  3.673077
```

```
Error in var.test.default(associate$Social, master$Social): not enough 'x'
↳ observations
Traceback:

1. var.test(associate$Social, master$Social)
2. var.test.default(associate$Social, master$Social)
3. stop("not enough 'x' observations")
```

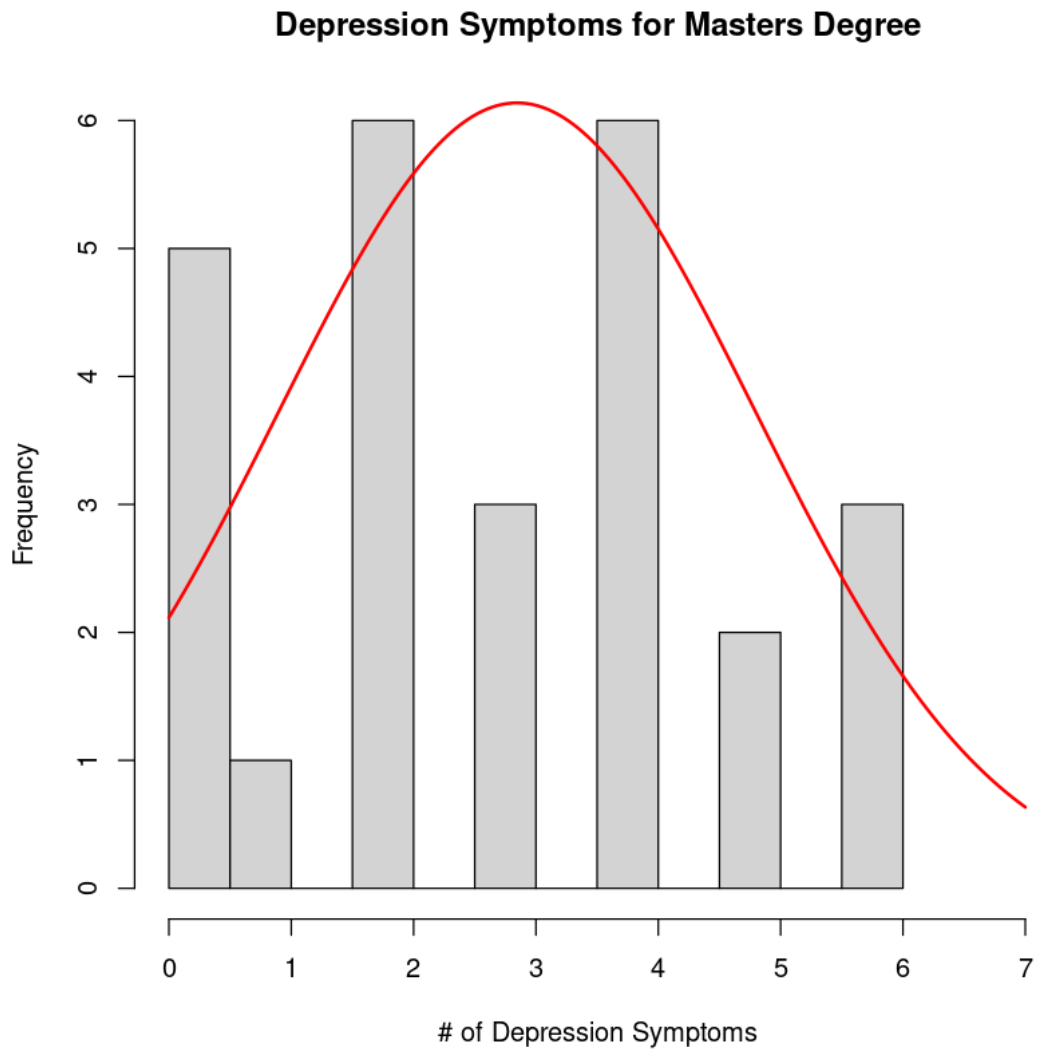
```
[ ]: #There is no significance between the number of interactions and degree. If
↳ there is no significance here, there is no point
# in searching if depression symptoms are related to number of interactions for
↳ the subsets associate and masters.
```

```
[45]: #Use normal distributions in order to compare the mean values of the subsets
↳ for depression and children

hist(master$Sad4, # dataframe$vector
      breaks = 20, # breaks sets the approximate number of bars
      main = "Depression Symptoms for Masters Degree", # graph title
      xlab = "# of Depression Symptoms", # x-axis title
      ylab = "Frequency",
      xlim = c(0,7) # y-axis title
    )
```

```
x <- seq(0,7,0.01)

lines(x,
      # x coordinate
      30*dnorm(x, mean = 2.846, sd = 1.95), # first peak
      col = "red",
      lwd = 2
    )
```



```
[44]: hist(associate$Sad4, # dataframe$vector
  breaks = 20, # breaks sets the approximate number of bars
  main = "Depression Symptoms for Associates Degree", # graph title
  xlab = "# of Depression Symptoms", # x-axis title
```



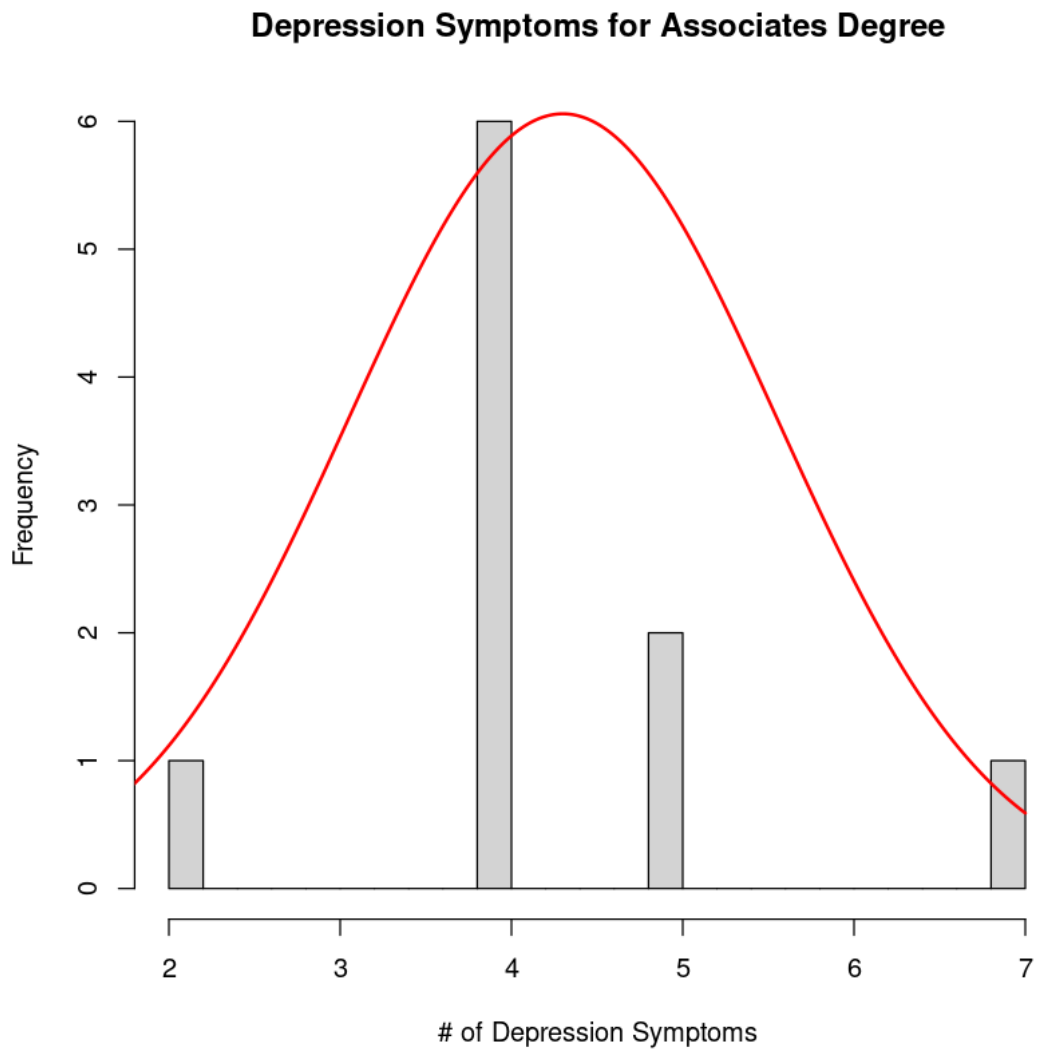
```

ylab = "Frequency" # y-axis title
)

x <- seq(0,7,0.01)

lines(x,
      # x coordinate
      19*dnorm(x, mean = 4.3, sd = 1.251), # first peak
      col = "red",
      lwd = 2
    )

```



```
[16]: #Many people who had an associates degree did not report depression symptoms.
```

```
length(associate$Sad4)
```

```
length(master$Sad4)
```

```
#All I am trying to do is convert these histograms into distributions so that I  
  ↳ can plot them all into one and it will  
# have 4 distributions, 2 for children, 2 for depression symptoms, and all for  
  ↳ the masters and associates degree.
```

154

548

```
[18]: #Also add another conditions where you only include the
```

```
associate$Sad4
```

```
1. <NA> 2. <NA> 3. <NA> 4. <NA> 5. 5 6. <NA> 7. 5 8. <NA> 9. <NA> 10. <NA> 11. <NA>  
12. <NA> 13. <NA> 14. <NA> 15. <NA> 16. <NA> 17. <NA> 18. <NA> 19. <NA> 20. 7  
21. <NA> 22. <NA> 23. <NA> 24. <NA> 25. <NA> 26. 2 27. <NA> 28. <NA> 29. <NA>  
30. <NA> 31. <NA> 32. <NA> 33. <NA> 34. <NA> 35. <NA> 36. <NA> 37. <NA> 38. <NA>  
39. <NA> 40. <NA> 41. <NA> 42. <NA> 43. <NA> 44. <NA> 45. <NA> 46. <NA> 47. <NA>  
48. <NA> 49. <NA> 50. <NA> 51. <NA> 52. <NA> 53. <NA> 54. <NA> 55. <NA> 56. <NA>  
57. <NA> 58. <NA> 59. <NA> 60. <NA> 61. <NA> 62. <NA> 63. <NA> 64. <NA> 65. <NA>  
66. <NA> 67. <NA> 68. <NA> 69. <NA> 70. <NA> 71. <NA> 72. <NA> 73. <NA> 74. <NA>  
75. <NA> 76. 4 77. <NA> 78. <NA> 79. <NA> 80. <NA> 81. <NA> 82. <NA> 83. <NA>  
84. <NA> 85. <NA> 86. <NA> 87. <NA> 88. <NA> 89. <NA> 90. <NA> 91. <NA> 92. 4  
93. <NA> 94. <NA> 95. <NA> 96. <NA> 97. <NA> 98. 4 99. <NA> 100. <NA> 101. <NA>  
102. <NA> 103. <NA> 104. <NA> 105. <NA> 106. <NA> 107. <NA> 108. <NA> 109. <NA>  
110. <NA> 111. <NA> 112. <NA> 113. 4 114. <NA> 115. <NA> 116. <NA> 117. <NA>  
118. <NA> 119. <NA> 120. <NA> 121. <NA> 122. <NA> 123. <NA> 124. <NA> 125. <NA>  
126. <NA> 127. <NA> 128. <NA> 129. <NA> 130. <NA> 131. <NA> 132. <NA> 133. <NA>  
134. <NA> 135. 4 136. 4 137. <NA> 138. <NA> 139. <NA> 140. <NA> 141. <NA> 142. <NA>  
143. <NA> 144. <NA> 145. <NA> 146. <NA> 147. <NA> 148. <NA> 149. <NA> 150. <NA>  
151. <NA> 152. <NA> 153. <NA> 154. <NA>
```

```
[39]: sd(associate$Sad4, na.rm = TRUE)
```

1.25166555703457

```
[ ]: hist(master1$Children, # dataframe$vector  
  breaks = 20, # breaks sets the approximate number of bars  
  main = "# of Children for Masters Degree", # graph title  
  xlab = "# of Depression Symptoms", # x-axis title  
  ylab = "Frequency" # y-axis title  
)
```

```
hist(associate$Sad4, # dataframe$vector
```

```

breaks = 20, # breaks sets the approximate number of bars
main = "Depression Symptoms for Associates Degree", # graph title
xlab = "# of Depression Symptoms", # x-axis title
ylab = "Frequency" # y-axis title
)

hist(associate1$Children, # dataframe$vector
breaks = 20, # breaks sets the approximate number of bars
main = "# of Children for Associates Degree", # graph title
xlab = "# of Depression Symptoms", # x-axis title
ylab = "Frequency" # y-axis title
)

```

```

[ ]: #RESET FOR HISTOGRAMS WITH LINES. CREATE A PLOT with 4 means and alternate
     ↪ colors to show means.

```

```

[14]: associate <- subset(WLS, Degree == 1)
      bachelor <- subset(WLS, Degree == 2)
      master <- subset(WLS, Degree == 3)
      doctor <- subset(WLS, Degree == 4)

      associate_less <- subset(associate, Children < 3)
      bachelor_less <- subset(bachelor, Children < 3)
      master_less <- subset(master, Children < 3)
      doctorate_less <- subset(doctor, Children < 3)

      t.test(associate_less$Children, master_less$Children,
             conf.level = 0.99,
             var.equal = TRUE)

```

Two Sample t-test

```

data: associate_less$Children and master_less$Children
t = 2.2558, df = 265, p-value = 0.0249
alternative hypothesis: true difference in means is not equal to 0
99 percent confidence interval:
 -0.05014971  0.71816590
sample estimates:
mean of x mean of y
 1.641026  1.307018

```

```

[6]: WLS99 <- subset(WLS, Children < 4)

associate99 <- subset(WLS99, Degree == 1)
bachelor99 <- subset(WLS99, Degree == 2)
master99 <- subset(WLS99, Degree == 3)
doctor99 <- subset(WLS99, Degree == 4)

c1 <- rainbow(4)
c2 <- rainbow(4, alpha=0.2)
c3 <- rainbow(4, v=0.7)

boxplot(Sad4~Degree,
        data = WLS99,
        names = c("associate", "bachelor", "master", "doctorate"),
        xlab = "Degree",
        ylab = "Number of Depression Symptoms",
        main = "Depression vs. Degree Under Four Children",
        range = 8,
        col = c2,
        medcol = c3,
        whiskcol = c1,
        staplecol= c3,
        boxcol = c3,
        outcol = c3,
        pch = 23,
        cex = 2)

text(1,4.3, # same x but y slightly lower
     labels = "4p",
     cex = 1,
     col = "black"
    )

text(2,3.8, # same x but y slightly lower
     labels = "29p",
     cex = 1,
     col = "black"
    )

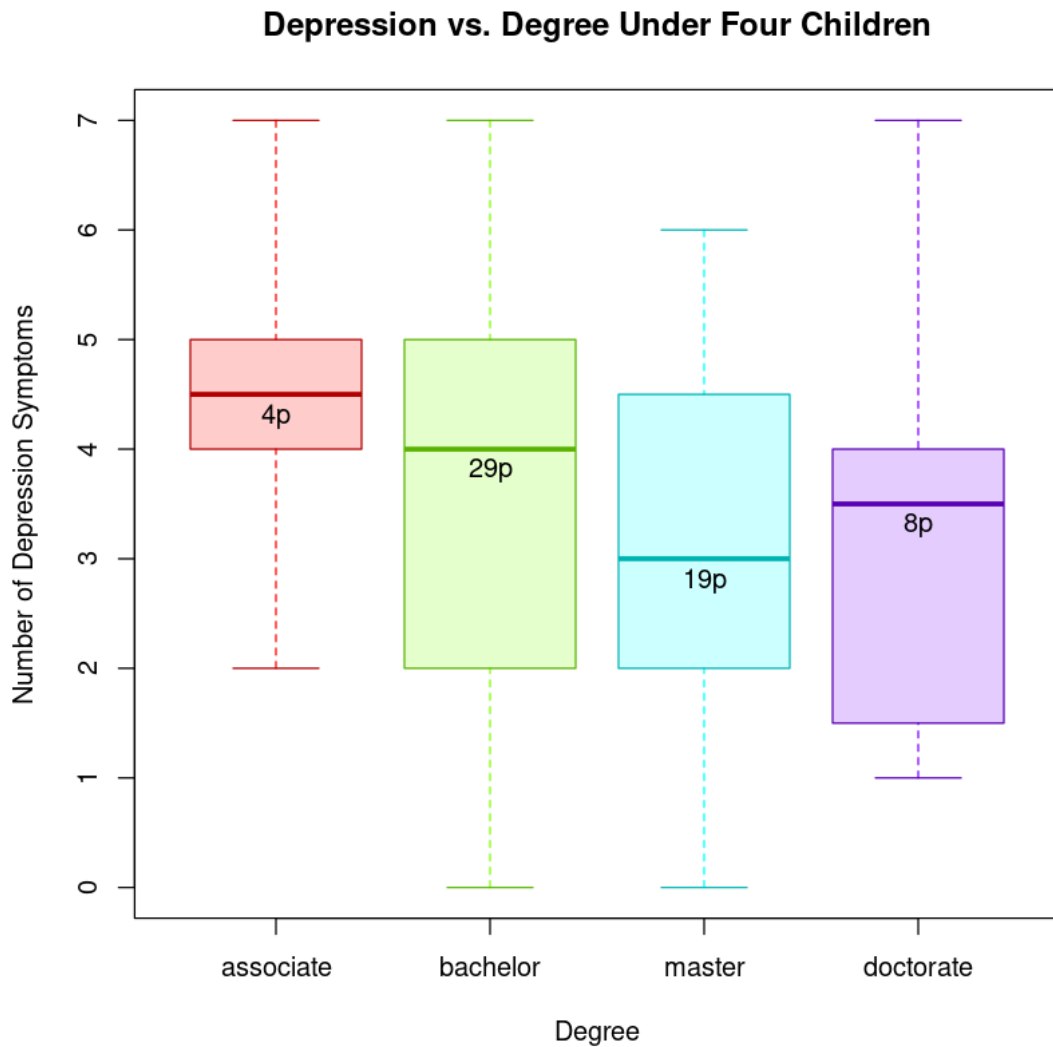
text(3,2.8, # same x but y slightly lower
     labels = "19p",
     cex = 1,
     col = "black"
    )

```

```

text(4,3.3, # same x but y slightly lower
     labels = "8p",
     cex = 1,
     col = "black"
)

```



```

[10]: WLSA <- subset(WLS, Children < 3)

associate99A <- subset(WLSA, Degree == 1)
bachelor99A <- subset(WLSA, Degree == 2)
master99A <- subset(WLSA, Degree == 3)

```

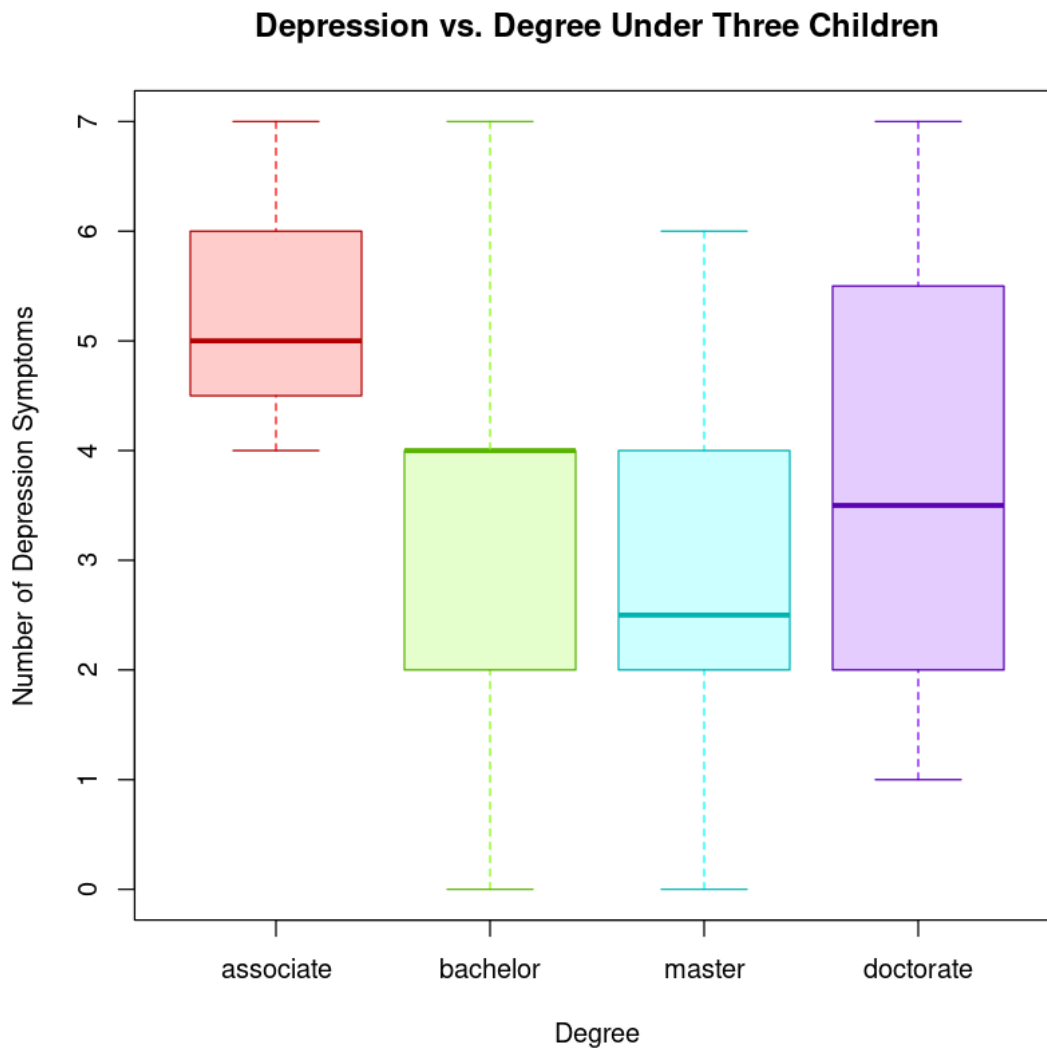
```

doctor99A <- subset(WLSA, Degree == 4)

c1 <- rainbow(4)
c2 <- rainbow(4, alpha=0.2)
c3 <- rainbow(4, v=0.7)

boxplot(Sad4~Degree,
        data = WLSA,
        names = c("associate", "bachelor", "master", "doctorate"),
        xlab = "Degree",
        ylab = "Number of Depression Symptoms",
        main = "Depression vs. Degree Under Three Children",
        range = 8,
        col = c2,
        medcol = c3,
        whiskcol = c1,
        staplecol= c3,
        boxcol = c3,
        outcol = c3,
        pch = 23,
        cex = 2)

```



```
[9]: WLSB <- subset(WLS, Children < 5)

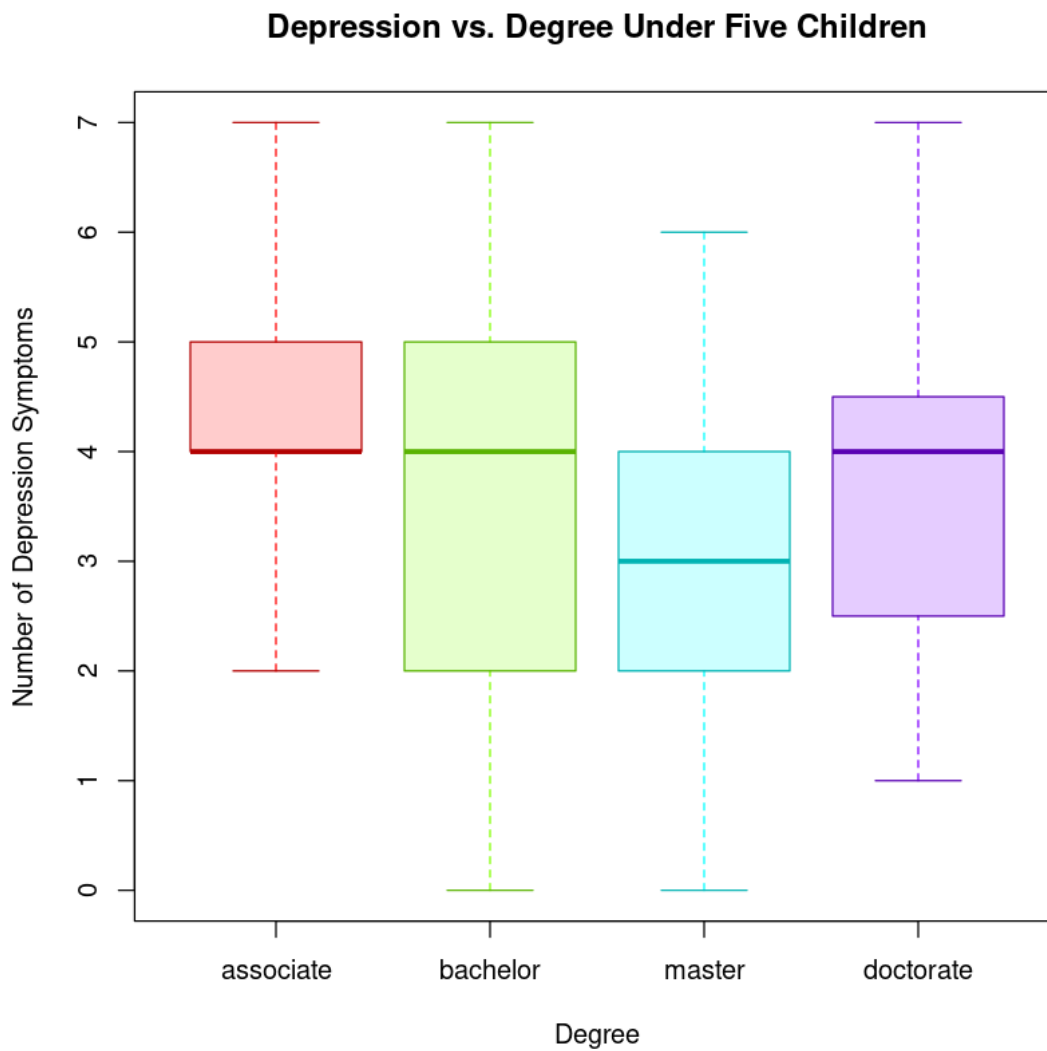
associate99B <- subset(WLSB, Degree == 1)
bachelor99B <- subset(WLSB, Degree == 2)
master99B <- subset(WLSB, Degree == 3)
doctor99B <- subset(WLSB, Degree == 4)

c1 <- rainbow(4)
c2 <- rainbow(4, alpha=0.2)
c3 <- rainbow(4, v=0.7)
```

```

boxplot(Sad4~Degree,
  data = WLSB,
  names = c("associate", "bachelor", "master", "doctorate"),
  xlab = "Degree",
  ylab = "Number of Depression Symptoms",
  main = "Depression vs. Degree Under Five Children",
  range = 8,
  col = c2,
  medcol = c3,
  whiskcol = c1,
  staplecol= c3,
  boxcol = c3,
  outcol = c3,
  pch = 23,
  cex = 2)

```




```

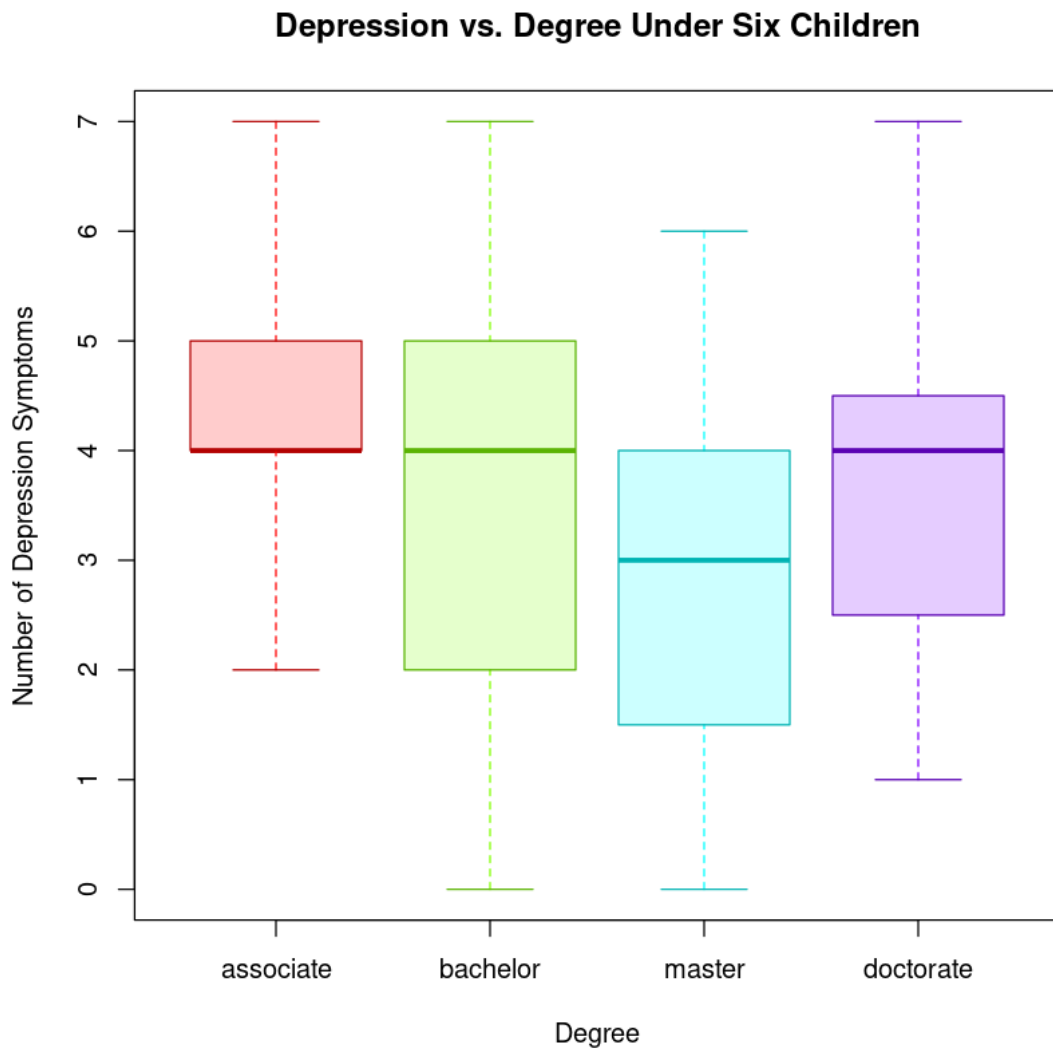
[11]: WLSC <- subset(WLS, Children < 6)

associate99C <- subset(WLSC, Degree == 1)
bachelor99C <- subset(WLSC, Degree == 2)
master99C <- subset(WLSC, Degree == 3)
doctor99C <- subset(WLSC, Degree == 4)

c1 <- rainbow(4)
c2 <- rainbow(4, alpha=0.2)
c3 <- rainbow(4, v=0.7)

boxplot(Sad4~Degree,
        data = WLSC,
        names = c("associate", "bachelor", "master", "doctorate"),
        xlab = "Degree",
        ylab = "Number of Depression Symptoms",
        main = "Depression vs. Degree Under Six Children",
        range = 8,
        col = c2,
        medcol = c3,
        whiskcol = c1,
        staplecol = c3,
        boxcol = c3,
        outcol = c3,
        pch = 23,
        cex = 2)

```



```
[12]: WLS D <- subset(WLS, Children < 7)

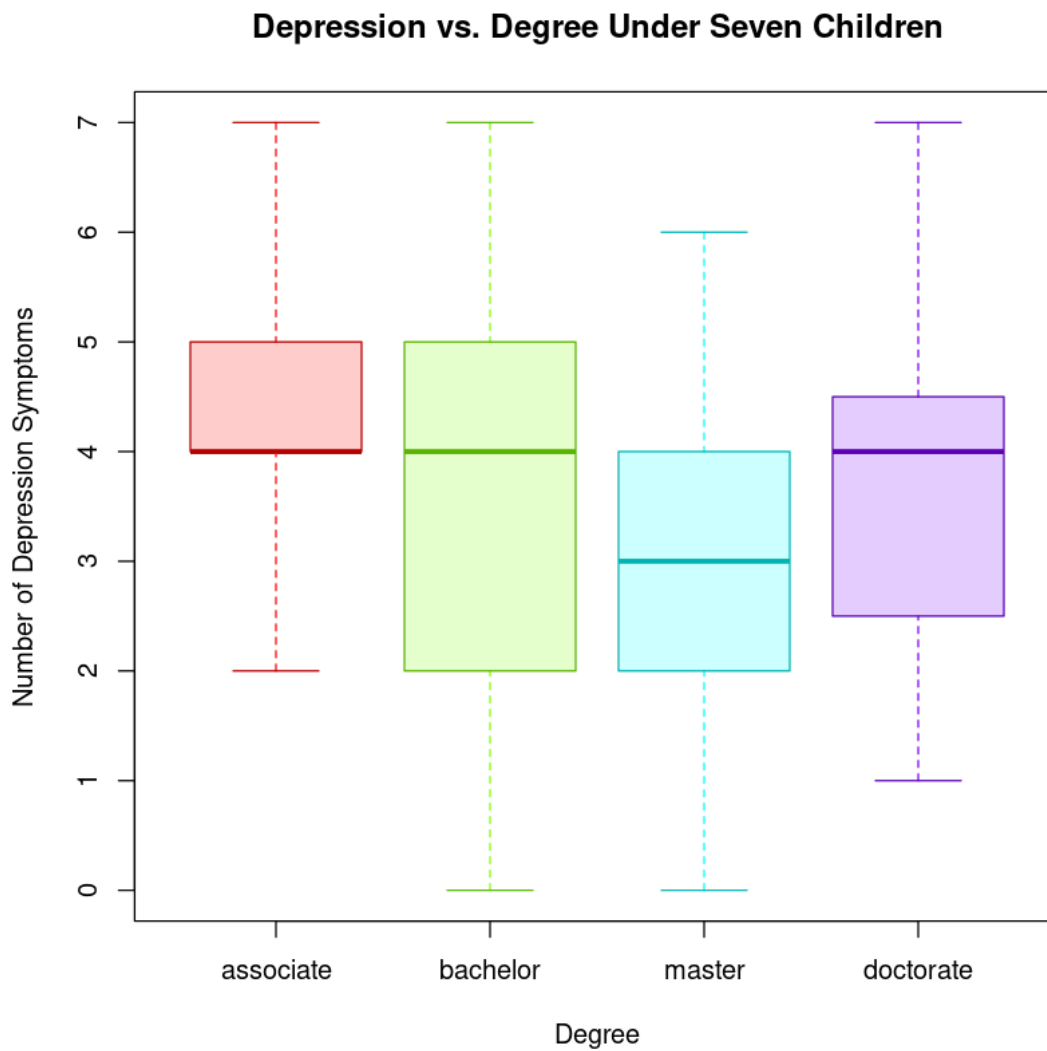
associate99D <- subset(WLS D, Degree == 1)
bachelor99D <- subset(WLS D, Degree == 2)
master99D <- subset(WLS D, Degree == 3)
doctor99D <- subset(WLS D, Degree == 4)

c1 <- rainbow(4)
c2 <- rainbow(4, alpha=0.2)
c3 <- rainbow(4, v=0.7)
```

```

boxplot(Sad4~Degree,
  data = WLSData,
  names = c("associate", "bachelor", "master", "doctorate"),
  xlab = "Degree",
  ylab = "Number of Depression Symptoms",
  main = "Depression vs. Degree Under Seven Children",
  range = 8,
  col = c2,
  medcol = c3,
  whiskcol = c1,
  staplecol= c3,
  boxcol = c3,
  outcol = c3,
  pch = 23,
  cex = 2)

```



```

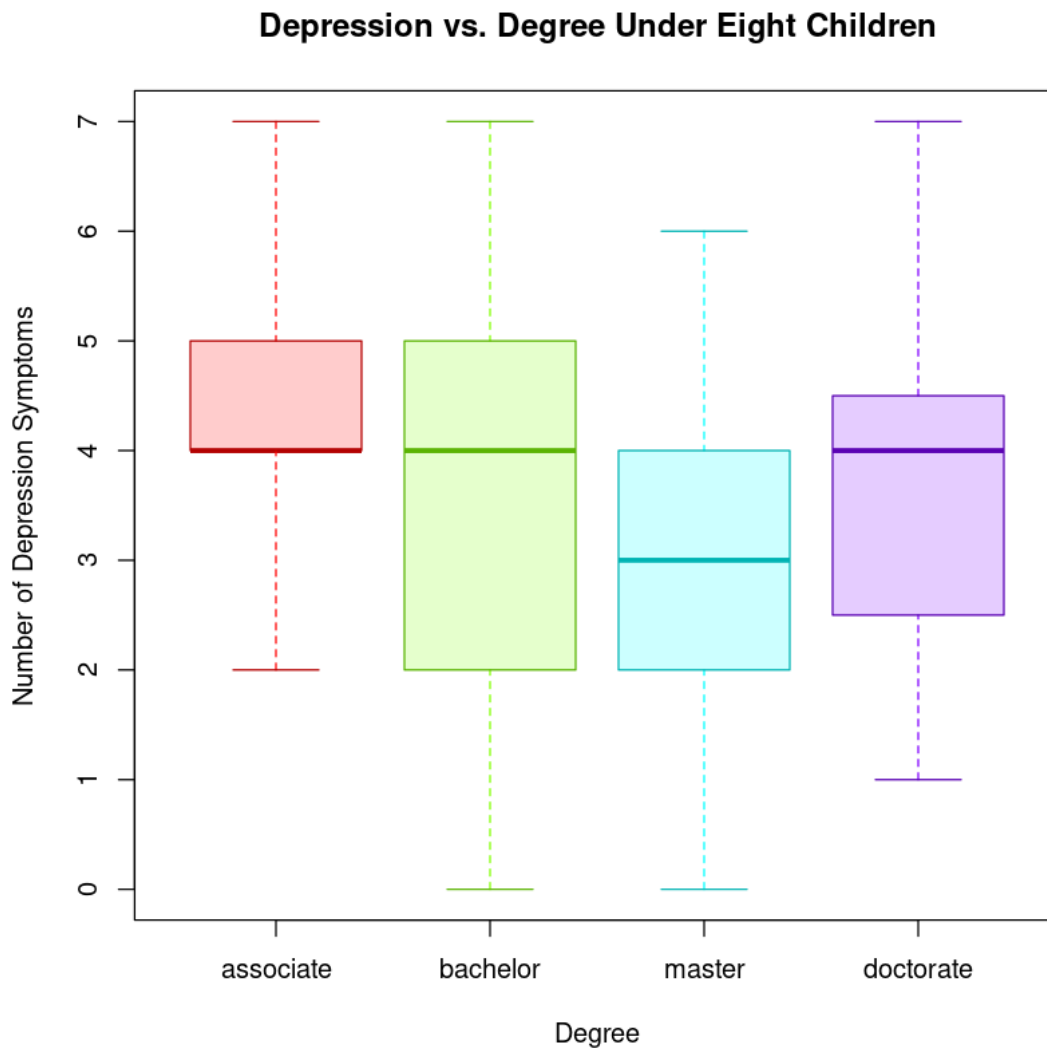
[13]: WLSE <- subset(WLS, Children < 8)

associate99E <- subset(WLSE, Degree == 1)
bachelor99E <- subset(WLSE, Degree == 2)
master99E <- subset(WLSE, Degree == 3)
doctor99E <- subset(WLSE, Degree == 4)

c1 <- rainbow(4)
c2 <- rainbow(4, alpha=0.2)
c3 <- rainbow(4, v=0.7)

boxplot(Sad4~Degree,
        data = WLSE,
        names = c("associate", "bachelor", "master", "doctorate"),
        xlab = "Degree",
        ylab = "Number of Depression Symptoms",
        main = "Depression vs. Degree Under Eight Children",
        range = 8,
        col = c2,
        medcol = c3,
        whiskcol = c1,
        staplecol = c3,
        boxcol = c3,
        outcol = c3,
        pch = 23,
        cex = 2)

```



```
[14]: #Under 3 Children
t.test(associate99$Children, master99$Children,
       conf.level = 0.99,
       alternative = "g",
       var.equal = TRUE)

#Under 4 Children
t.test(associate99A$Children, master99A$Children,
       conf.level = 0.99,
       alternative = "g",
       var.equal = TRUE)

#Under 5 Children
```

```

t.test(associate99B$Children, master99B$Children,
       conf.level = 0.99,
       alternative = "g",
       var.equal = TRUE)

#Under 6 Children
t.test(associate99C$Children, master99C$Children,
       conf.level = 0.99,
       alternative = "g",
       var.equal = TRUE)

#Under 7 Children
t.test(associate99D$Children, master99D$Children,
       conf.level = 0.99,
       alternative = "g",
       var.equal = TRUE)

#Under 8 Children
t.test(associate99E$Children, master99E$Children,
       conf.level = 0.99,
       alternative = "g",
       var.equal = TRUE)

```

Two Sample t-test

```

data: associate99$Children and master99$Children
t = 2.6658, df = 377, p-value = 0.004005
alternative hypothesis: true difference in means is greater than 0
99 percent confidence interval:
 0.0470054      Inf
sample estimates:
mean of x mean of y
 2.158730  1.778481

```

Two Sample t-test

```

data: associate99A$Children and master99A$Children
t = 2.2558, df = 265, p-value = 0.01245
alternative hypothesis: true difference in means is greater than 0
99 percent confidence interval:
-0.01253996      Inf
sample estimates:
mean of x mean of y
 1.641026  1.307018

```

Two Sample t-test

```
data: associate99B$Children and master99B$Children
t = 3.6382, df = 452, p-value = 0.0001531
alternative hypothesis: true difference in means is greater than 0
99 percent confidence interval:
 0.1926661      Inf
sample estimates:
mean of x mean of y
 2.635294  2.097561
```

Two Sample t-test

```
data: associate99C$Children and master99C$Children
t = 3.6776, df = 478, p-value = 0.0001311
alternative hypothesis: true difference in means is greater than 0
99 percent confidence interval:
 0.210242      Inf
sample estimates:
mean of x mean of y
 2.815217  2.239691
```

Two Sample t-test

```
data: associate99D$Children and master99D$Children
t = 3.9322, df = 496, p-value = 4.808e-05
alternative hypothesis: true difference in means is greater than 0
99 percent confidence interval:
 0.2673313      Inf
sample estimates:
mean of x mean of y
 3.010204  2.352500
```

Two Sample t-test

```
data: associate99E$Children and master99E$Children
t = 4.0474, df = 508, p-value = 2.993e-05
alternative hypothesis: true difference in means is greater than 0
99 percent confidence interval:
 0.3061423      Inf
sample estimates:
mean of x mean of y
```

3.166667 2.443627

```
[15]: #Under 4 Children
t.test(bachelor99$Children, master99$Children,
       conf.level = 0.99,
       alternative = "g",
       var.equal = TRUE)

#Under 3 Children
t.test(bachelor99A$Children, master99A$Children,
       conf.level = 0.99,
       alternative = "g",
       var.equal = TRUE)

#Under 5 Children
t.test(bachelor99B$Children, master99B$Children,
       conf.level = 0.99,
       alternative = "g",
       var.equal = TRUE)

#Under 6 Children
t.test(bachelor99C$Children, master99C$Children,
       conf.level = 0.99,
       alternative = "g",
       var.equal = TRUE)

#Under 7 Children
t.test(bachelor99D$Children, master99D$Children,
       conf.level = 0.99,
       alternative = "g",
       var.equal = TRUE)

#Under 8 Children
t.test(bachelor99E$Children, master99E$Children,
       conf.level = 0.99,
       alternative = "g",
       var.equal = TRUE)
```

Two Sample t-test

```
data: bachelor99$Children and master99$Children
t = 3.7873, df = 1073, p-value = 8.038e-05
alternative hypothesis: true difference in means is greater than 0
99 percent confidence interval:
 0.09792243      Inf
sample estimates:
```


mean of x mean of y
2.032938 1.778481

Two Sample t-test

data: bachelor99A\$Children and master99A\$Children
t = 2.3652, df = 704, p-value = 0.009146
alternative hypothesis: true difference in means is greater than 0
99 percent confidence interval:
0.002231035 Inf
sample estimates:
mean of x mean of y
1.464435 1.307018

Two Sample t-test

data: bachelor99B\$Children and master99B\$Children
t = 3.2646, df = 1264, p-value = 0.0005629
alternative hypothesis: true difference in means is greater than 0
99 percent confidence interval:
0.06818482 Inf
sample estimates:
mean of x mean of y
2.335563 2.097561

Two Sample t-test

data: bachelor99C\$Children and master99C\$Children
t = 4.0043, df = 1366, p-value = 3.277e-05
alternative hypothesis: true difference in means is greater than 0
99 percent confidence interval:
0.1345163 Inf
sample estimates:
mean of x mean of y
2.561224 2.239691

Two Sample t-test

data: bachelor99D\$Children and master99D\$Children
t = 3.8627, df = 1415, p-value = 5.861e-05
alternative hypothesis: true difference in means is greater than 0
99 percent confidence interval:

```
0.132549      Inf
sample estimates:
mean of x mean of y
2.686332 2.352500
```

Two Sample t-test

```
data: bachelor99E$Children and master99E$Children
t = 3.3576, df = 1438, p-value = 0.0004033
alternative hypothesis: true difference in means is greater than 0
99 percent confidence interval:
0.09356767      Inf
sample estimates:
mean of x mean of y
2.749031 2.443627
```

```
[16]: #Under 4 Children
t.test(doctor99$Children, master99$Children,
       conf.level = 0.99,
       alternative = "g",
       var.equal = TRUE)

#Under 3 Children
t.test(doctor99A$Children, master99A$Children,
       conf.level = 0.99,
       alternative = "g",
       var.equal = TRUE)

#Under 5 Children
t.test(doctor99B$Children, master99B$Children,
       conf.level = 0.99,
       alternative = "g",
       var.equal = TRUE)

#Under 6 Children
t.test(doctor99C$Children, master99C$Children,
       conf.level = 0.99,
       alternative = "g",
       var.equal = TRUE)

#Under 7 Children
t.test(doctor99D$Children, master99D$Children,
       conf.level = 0.99,
       alternative = "g",
       var.equal = TRUE)
```

```
#Under 8 Children
t.test(doctor99E$Children, master99E$Children,
       conf.level = 0.99,
       alternative = "g",
       var.equal = TRUE)
```

Two Sample t-test

```
data: doctor99$Children and master99$Children
t = 2.6402, df = 423, p-value = 0.004295
alternative hypothesis: true difference in means is greater than 0
99 percent confidence interval:
 0.03513418      Inf
sample estimates:
mean of x mean of y
 2.082569  1.778481
```

Two Sample t-test

```
data: doctor99A$Children and master99A$Children
t = 1.4896, df = 292, p-value = 0.0687
alternative hypothesis: true difference in means is greater than 0
99 percent confidence interval:
-0.1014176      Inf
sample estimates:
mean of x mean of y
 1.484848  1.307018
```

Two Sample t-test

```
data: doctor99B$Children and master99B$Children
t = 3.0665, df = 504, p-value = 0.001141
alternative hypothesis: true difference in means is greater than 0
99 percent confidence interval:
 0.09005374      Inf
sample estimates:
mean of x mean of y
 2.474453  2.097561
```

Two Sample t-test

```

data: doctor99C$Children and master99C$Children
t = 2.5997, df = 529, p-value = 0.004796
alternative hypothesis: true difference in means is greater than 0
99 percent confidence interval:
 0.03489662      Inf
sample estimates:
mean of x mean of y
 2.580420  2.239691

```

Two Sample t-test

```

data: doctor99D$Children and master99D$Children
t = 3.0414, df = 550, p-value = 0.001233
alternative hypothesis: true difference in means is greater than 0
99 percent confidence interval:
 0.1002285      Inf
sample estimates:
mean of x mean of y
 2.782895  2.352500

```

Two Sample t-test

```

data: doctor99E$Children and master99E$Children
t = 2.7892, df = 561, p-value = 0.002731
alternative hypothesis: true difference in means is greater than 0
99 percent confidence interval:
 0.06884338      Inf
sample estimates:
mean of x mean of y
 2.864516  2.443627

```

```
[84]: WLS3 <- subset(WLS, Children < 3)
```

```

associate <- subset(WLS3, Degree == 1)
bachelor <- subset(WLS3, Degree == 2)
master <- subset(WLS3, Degree == 3)
doctor <- subset(WLS3, Degree == 4)

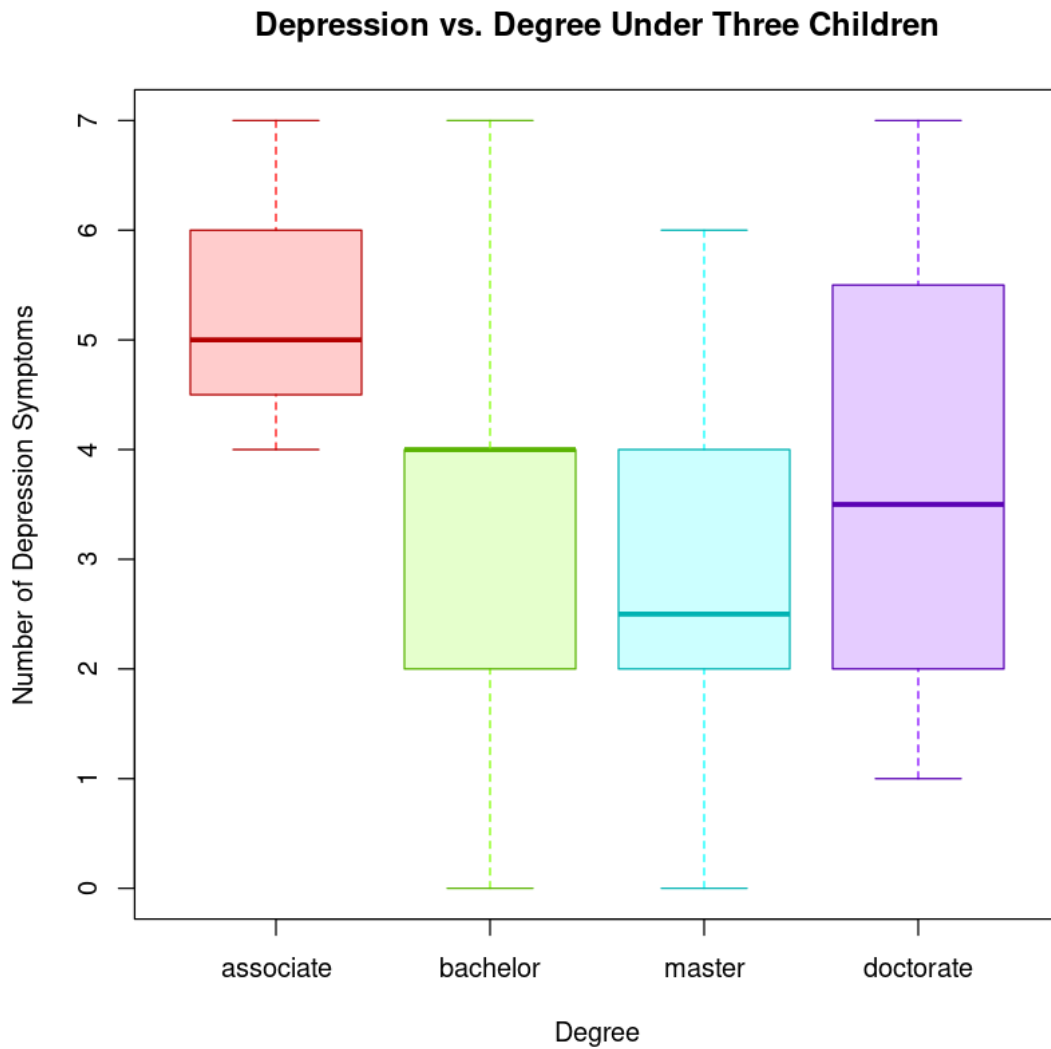
c1 <- rainbow(4)
c2 <- rainbow(4, alpha=0.2)
c3 <- rainbow(4, v=0.7)

```

```

boxplot(Sad4~Degree,
  data = WLS3,
  names = c("associate", "bachelor", "master", "doctorate"),
  xlab = "Degree",
  ylab = "Number of Depression Symptoms",
  main = "Depression vs. Degree Under Three Children",
  range = 8,
  col = c2,
  medcol = c3,
  whiskcol = c1,
  staplecol= c3,
  boxcol = c3,
  outcol = c3,
  pch = 23,
  cex = 2)

```



```
[87]: associate$Sad4 #3
      bachelor$Sad4 #28
      master$Sad4 #12
      doctor$Sad4 #4
```

1. <NA> 2. <NA> 3. 5 4. <NA> 5. <NA> 6. <NA> 7. 7 8. <NA> 9. <NA> 10. <NA> 11. <NA>
 12. <NA> 13. <NA> 14. <NA> 15. <NA> 16. <NA> 17. <NA> 18. <NA> 19. <NA> 20. <NA>
 21. <NA> 22. <NA> 23. <NA> 24. <NA> 25. <NA> 26. <NA> 27. <NA> 28. <NA> 29. <NA>
 30. 4 31. <NA> 32. <NA> 33. <NA> 34. <NA> 35. <NA> 36. <NA> 37. <NA> 38. <NA>
 39. <NA>

1. <NA> 2. <NA> 3. <NA> 4. <NA> 5. <NA> 6. <NA> 7. <NA> 8. <NA> 9. <NA>
 10. <NA> 11. <NA> 12. <NA> 13. <NA> 14. <NA> 15. <NA> 16. <NA> 17. <NA> 18. 3
 19. <NA> 20. 4 21. <NA> 22. <NA> 23. <NA> 24. <NA> 25. <NA> 26. <NA> 27. <NA>
 28. <NA> 29. <NA> 30. <NA> 31. <NA> 32. <NA> 33. <NA> 34. <NA> 35. <NA> 36. <NA>
 37. <NA> 38. <NA> 39. 1 40. <NA> 41. <NA> 42. <NA> 43. <NA> 44. <NA> 45. <NA>
 46. <NA> 47. <NA> 48. <NA> 49. <NA> 50. <NA> 51. <NA> 52. <NA> 53. <NA> 54. <NA>
 55. <NA> 56. <NA> 57. 5 58. <NA> 59. <NA> 60. 4 61. <NA> 62. <NA> 63. <NA> 64. <NA>
 65. <NA> 66. <NA> 67. <NA> 68. <NA> 69. <NA> 70. <NA> 71. <NA> 72. <NA> 73. <NA>
 74. <NA> 75. <NA> 76. <NA> 77. <NA> 78. <NA> 79. <NA> 80. <NA> 81. <NA> 82. <NA>
 83. <NA> 84. <NA> 85. <NA> 86. <NA> 87. <NA> 88. <NA> 89. <NA> 90. <NA> 91. <NA>
 92. <NA> 93. <NA> 94. 2 95. <NA> 96. <NA> 97. <NA> 98. <NA> 99. <NA> 100. <NA>
 101. <NA> 102. <NA> 103. <NA> 104. 3 105. <NA> 106. <NA> 107. <NA> 108. 6 109. <NA>
 110. 5 111. <NA> 112. <NA> 113. <NA> 114. <NA> 115. <NA> 116. <NA> 117. <NA>
 118. <NA> 119. <NA> 120. <NA> 121. <NA> 122. <NA> 123. <NA> 124. <NA> 125. <NA>
 126. <NA> 127. <NA> 128. <NA> 129. <NA> 130. <NA> 131. <NA> 132. <NA> 133. 5
 134. <NA> 135. <NA> 136. <NA> 137. <NA> 138. <NA> 139. <NA> 140. <NA> 141. <NA>
 142. <NA> 143. <NA> 144. <NA> 145. <NA> 146. <NA> 147. <NA> 148. <NA> 149. <NA>
 150. <NA> 151. <NA> 152. <NA> 153. <NA> 154. <NA> 155. <NA> 156. <NA> 157. <NA>
 158. <NA> 159. <NA> 160. 4 161. <NA> 162. 3 163. <NA> 164. <NA> 165. 1 166. <NA>
 167. <NA> 168. <NA> 169. <NA> 170. <NA> 171. <NA> 172. 4 173. <NA> 174. <NA>
 175. <NA> 176. <NA> 177. 4 178. <NA> 179. <NA> 180. <NA> 181. <NA> 182. <NA>
 183. <NA> 184. <NA> 185. 7 186. <NA> 187. <NA> 188. <NA> 189. <NA> 190. <NA>
 191. <NA> 192. <NA> 193. <NA> 194. <NA> 195. <NA> 196. <NA> 197. <NA> 198. <NA>
 199. <NA> 200. <NA> 201. 202. <NA> 203. <NA> 204. <NA> 205. <NA> 206. <NA>
 207. <NA> 208. <NA> 209. <NA> 210. <NA> 211. <NA> 212. <NA> 213. <NA> 214. <NA>
 215. <NA> 216. <NA> 217. <NA> 218. <NA> 219. <NA> 220. <NA> 221. <NA> 222. <NA>
 223. <NA> 224. <NA> 225. <NA> 226. <NA> 227. <NA> 228. <NA> 229. <NA> 230. <NA>
 231. <NA> 232. <NA> 233. <NA> 234. <NA> 235. <NA> 236. <NA> 237. <NA> 238. <NA>
 239. <NA> 240. <NA> 241. 2 242. <NA> 243. <NA> 244. <NA> 245. <NA> 246. <NA>
 247. <NA> 248. <NA> 249. <NA> 250. 3 251. <NA> 252. <NA> 253. <NA> 254. <NA>
 255. <NA> 256. <NA> 257. 1 258. <NA> 259. <NA> 260. <NA> 261. <NA> 262. <NA>
 263. 0 264. <NA> 265. <NA> 266. <NA> 267. <NA> 268. <NA> 269. <NA> 270. <NA>
 271. <NA> 272. 2 273. <NA> 274. <NA> 275. <NA> 276. <NA> 277. <NA> 278. <NA>
 279. <NA> 280. <NA> 281. <NA> 282. <NA> 283. <NA> 284. <NA> 285. <NA> 286. <NA>

48. <NA> 49. <NA> 50. <NA> 51. <NA> 52. <NA> 53. <NA> 54. <NA> 55. <NA> 56. <NA>
57. <NA> 58. <NA> 59. <NA> 60. <NA> 61. <NA> 62. <NA> 63. <NA> 64. <NA> 65. 1
66. <NA>

```
[89]: t.test(associate$Sad4,doctor$Sad4,
          conf.level = 0.99,
          var.equal = TRUE)

var.test(associate$Sad4, doctor$Sad4)

prop.test(x = c(12, 18),
          n = c(47, 60),
          conf.level = 0.99,
          alternative = "t"
          )
```

Two Sample t-test

```
data: associate$Sad4 and doctor$Sad4
t = 0.95794, df = 5, p-value = 0.3821
alternative hypothesis: true difference in means is not equal to 0
99 percent confidence interval:
 -5.081232  8.247899
sample estimates:
mean of x mean of y
 5.333333  3.750000
```

F test to compare two variances

```
data: associate$Sad4 and doctor$Sad4
F = 0.37333, num df = 2, denom df = 3, p-value = 0.567
alternative hypothesis: true ratio of variances is not equal to 1
95 percent confidence interval:
 0.02326919 14.62178464
sample estimates:
ratio of variances
 0.3733333
```

2-sample test for equality of proportions with continuity correction

```
data: c(12, 18) out of c(47, 60)
X-squared = 0.086337, df = 1, p-value = 0.7689
alternative hypothesis: two.sided
99 percent confidence interval:
```



```
-0.2873992 0.1980375
sample estimates:
  prop 1    prop 2 
0.2553191 0.3000000
```

```
[3]: WLS3 <- subset(WLS, Social4 > 3.5)
     WLS4 <- subset(WLS, Social4 < 3.5)
```

```
[19]: quantile(WLS$Social4, 0.75, na.rm = TRUE)
```

```
75\%: 5
```

```
[6]: associate7 <- subset(WLS3, Degree == 1)
     bachelor7 <- subset(WLS3, Degree == 2)
     master7 <- subset(WLS3, Degree == 3)
     doctor7 <- subset(WLS3, Degree == 4)

     c1 <- rainbow(4)
     c2 <- rainbow(4, alpha=0.2)
     c3 <- rainbow(4, v=0.7)

     boxplot(Sad4~Degree,
             data = WLS3,
             names = c("associate", "bachelor", "master", "doctorate"),
             xlab = "Degree",
             ylab = "Number of Depression Symptoms",
             main = "Depression vs. Degree with Upper 50% Social",
             range = 8,
             col = c2,
             medcol = c3,
             whiskcol = c1,
             staplecol = c3,
             boxcol = c3,
             outcol = c3,
             pch = 23,
             cex = 2)

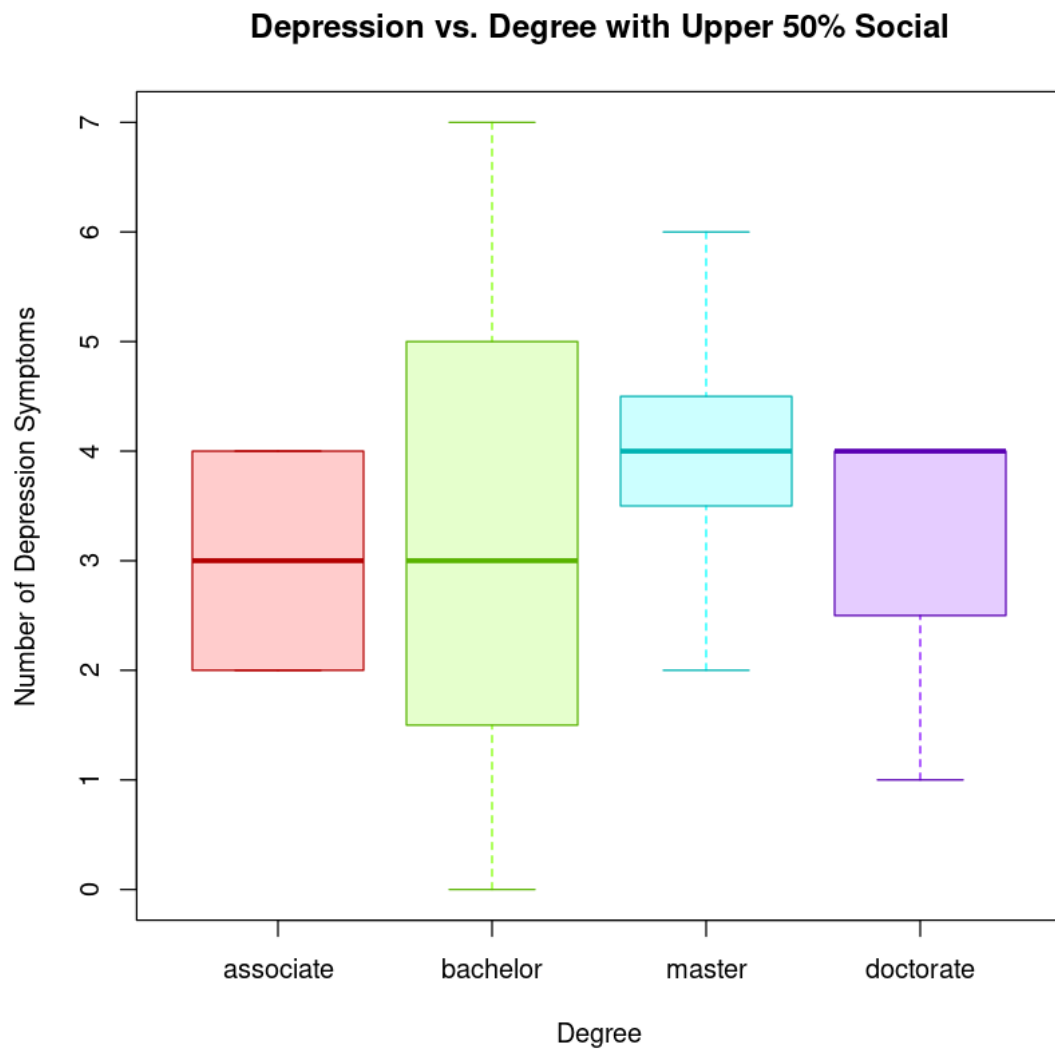
     mean(associate7$Children, na.rm = TRUE)
     mean(bachelor7$Children, na.rm = TRUE)
     mean(master7$Children, na.rm = TRUE)
     mean(doctor7$Children, na.rm = TRUE)
```

```
3.17647058823529
```

```
2.78139534883721
```

```
2.63333333333333
```

3.07894736842105



```
[7]: associate8 <- subset(WLS4, Degree == 1)
bachelor8 <- subset(WLS4, Degree == 2)
master8 <- subset(WLS4, Degree == 3)
doctor8 <- subset(WLS4, Degree == 4)

c1 <- rainbow(4)
c2 <- rainbow(4, alpha=0.2)
c3 <- rainbow(4, v=0.7)

boxplot(Sad4~Degree,
        data = WLS4,
```

```

names = c("associate", "bachelor", "master", "doctorate"),
xlab = "Degree",
ylab = "Number of Depression Symptoms",
main = "Depression vs. Degree with Lower 50% Social",
range = 8,
col = c2,
medcol = c3,
whiskcol = c1,
staplecol= c3,
boxcol = c3,
outcol = c3,
pch = 23,
cex = 2)

mean(associate8$Children, na.rm = TRUE)
mean(bachelor8$Children, na.rm = TRUE)
mean(master8$Children, na.rm = TRUE)
mean(doctor8$Children, na.rm = TRUE)

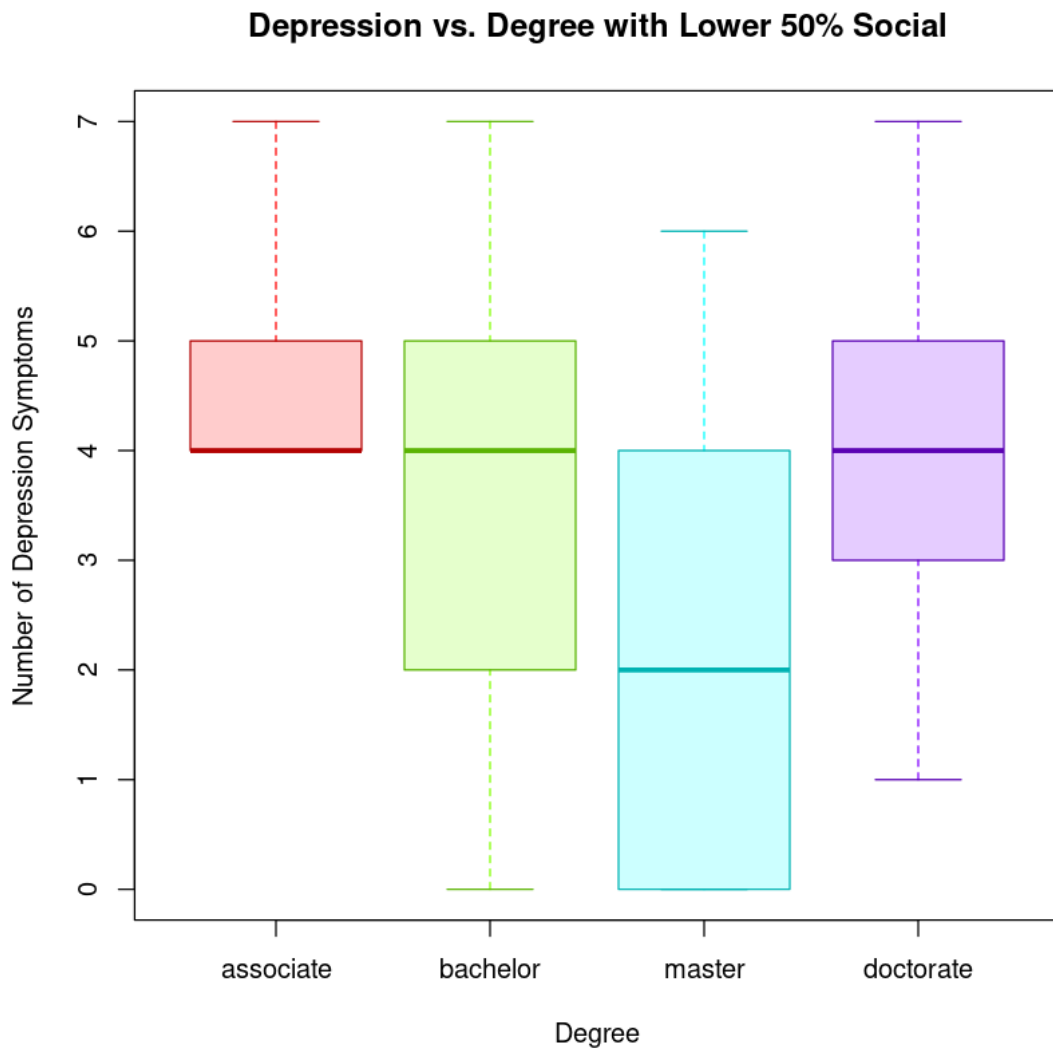
```

3.4

2.80366492146597

2.72258064516129

3.02985074626866



```
[17]: t.test(associate7$Sad4,associate8$Sad4,  
        conf.level = 0.99,  
        var.equal = TRUE)  
  
t.test(master7$Sad4,master8$Sad4,  
        conf.level = 0.99,  
        var.equal = TRUE)  
  
t.test(doctor7$Sad4,doctor8$Sad4,  
        conf.level = 0.99,  
        var.equal = TRUE)  
  
t.test(associate8$Sad4,master8$Sad4,
```

```

      conf.level = 0.99,
      alternative = "g",
      var.equal = TRUE)

t.test(associate7$Sad4, master7$Sad4,
      conf.level = 0.99,
      var.equal = TRUE)

```

Two Sample t-test

```

data: associate7$Sad4 and associate8$Sad4
t = -1.8501, df = 8, p-value = 0.1015
alternative hypothesis: true difference in means is not equal to 0
99 percent confidence interval:
 -4.572177  1.322177
sample estimates:
mean of x mean of y
  3.000    4.625

```

Two Sample t-test

```

data: master7$Sad4 and master8$Sad4
t = 2.1483, df = 24, p-value = 0.04199
alternative hypothesis: true difference in means is not equal to 0
99 percent confidence interval:
 -0.5031741  3.8365075
sample estimates:
mean of x mean of y
 4.000000  2.333333

```

Two Sample t-test

```

data: doctor7$Sad4 and doctor8$Sad4
t = -0.75864, df = 10, p-value = 0.4656
alternative hypothesis: true difference in means is not equal to 0
99 percent confidence interval:
 -4.602270  2.824492
sample estimates:
mean of x mean of y
 3.000000  3.888889

```

Two Sample t-test

```

data: associate8$Sad4 and master8$Sad4
t = 2.994, df = 24, p-value = 0.003147
alternative hypothesis: true difference in means is greater than 0
99 percent confidence interval:
 0.3841222      Inf
sample estimates:
mean of x mean of y
 4.625000  2.333333

```

Two Sample t-test

```

data: associate7$Sad4 and master7$Sad4
t = -1.0328, df = 8, p-value = 0.3319
alternative hypothesis: true difference in means is not equal to 0
99 percent confidence interval:
 -4.24884  2.24884
sample estimates:
mean of x mean of y
      3      4

```

```

[36]: prop.test(x = c(2, 3584),
               n = c(20, 39065),
               conf.level = 0.99,
               alternative = "g"
               )

```

```

Warning message in prop.test(x = c(2, 3584), n = c(20, 39065), conf.level =
0.99, :
"Chi-squared approximation may be incorrect"

```

2-sample test for equality of proportions with continuity correction

```

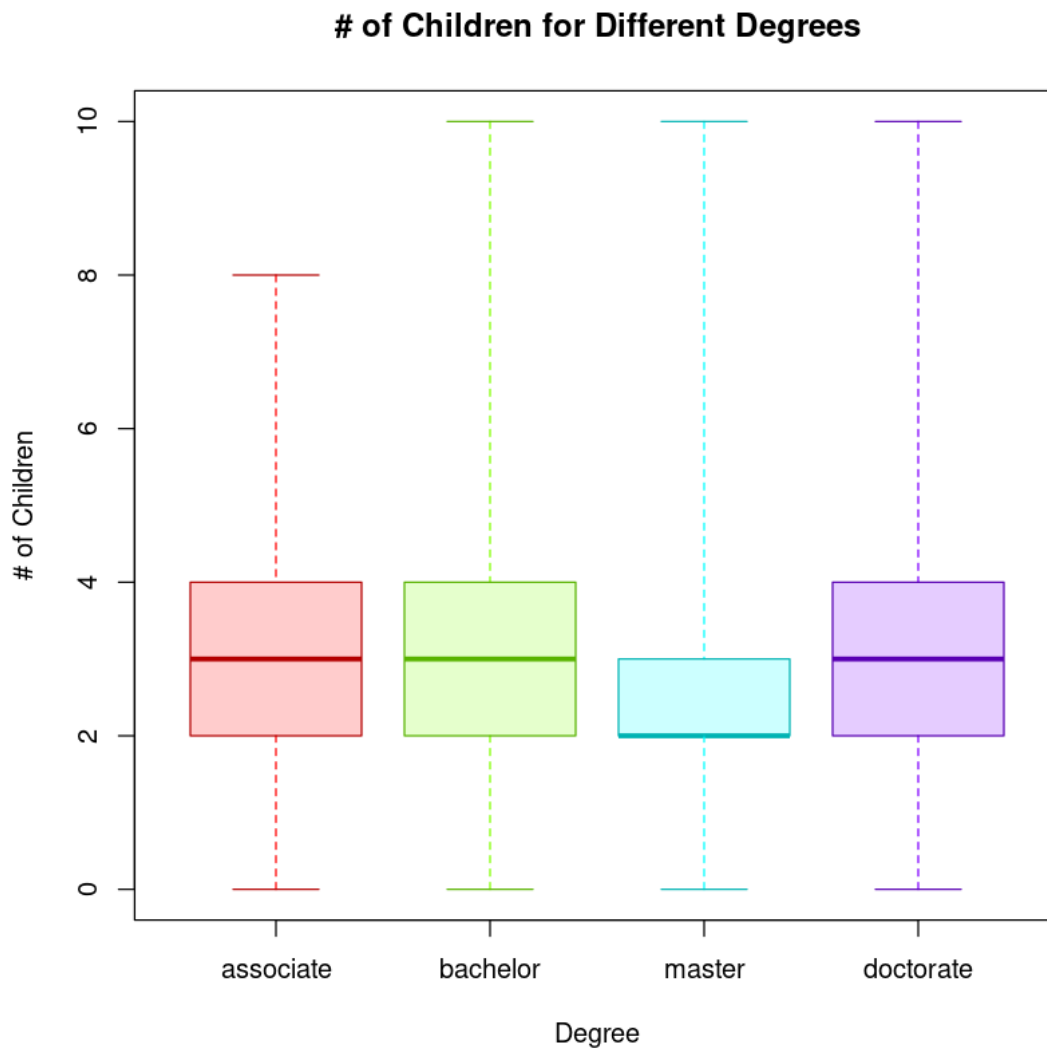
data: c(2, 3584) out of c(20, 39065)
X-squared = 1.8703e-28, df = 1, p-value = 0.5
alternative hypothesis: greater
99 percent confidence interval:
 -0.1560931  1.0000000
sample estimates:
 prop 1      prop 2
0.1000000 0.09174453

```

```
[27]: WL4MALE <- subset(WLS, Gender = 1)
      WL4FMALE <- subset(WLS, Gender = 2)
```

```
[29]: c1 <- rainbow(4)
      c2 <- rainbow(4, alpha=0.2)
      c3 <- rainbow(4, v=0.7)

      boxplot(Children~Degree,
              data = WL4MALE,
              names = c("associate", "bachelor", "master", "doctorate"),
              xlab = "Degree",
              ylab = "# of Children",
              main = "# of Children for Different Degrees",
              range = 8,
              col = c2,
              medcol = c3,
              whiskcol = c1,
              staplecol = c3,
              boxcol = c3,
              outcol = c3,
              pch = 23,
              cex = 2)
```

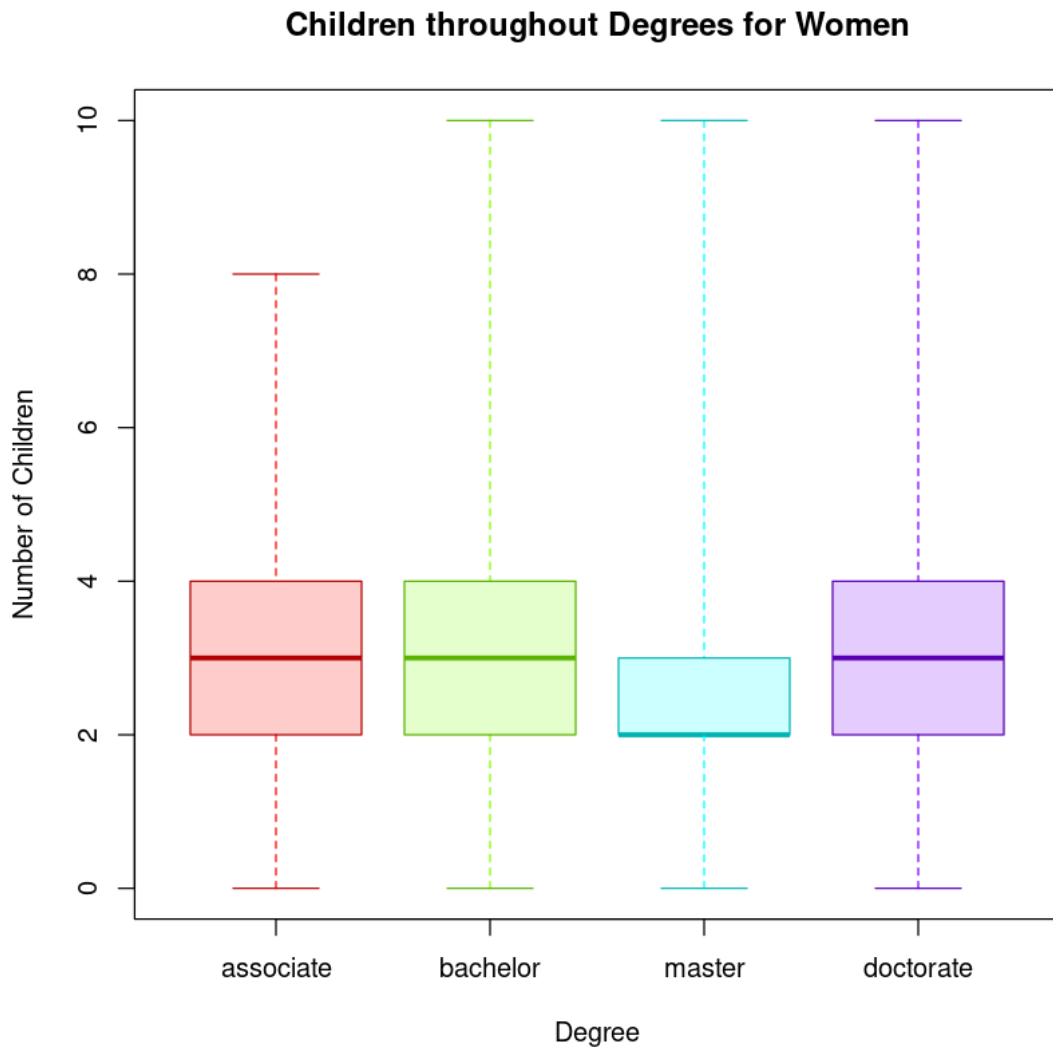


```
[31]: c1 <- rainbow(4)
c2 <- rainbow(4, alpha=0.2)
c3 <- rainbow(4, v=0.7)

boxplot(Children~Degree,
  data = WL4FMALE,
  names = c("associate", "bachelor", "master", "doctorate"),
  xlab = "Degree",
  ylab = "Number of Children",
  main = "Children throughout Degrees for Women",
  range = 8,
```



```
col = c2,  
medcol = c3,  
whiskcol = c1,  
staplecol= c3,  
boxcol = c3,  
outcol = c3,  
pch = 23,  
cex = 2)
```



```
[15]: mean(female$Children, na.rm = TRUE)  
mean(male$Children, na.rm = TRUE)
```

3.23959687906372

3.23959687906372

```
[17]: summary(WLS$Gender)
```

Min.	1st Qu.	Median	Mean	3rd Qu.	Max.
1.000	1.000	2.000	1.516	2.000	2.000

```
[74]: b <- cor(WLS$Degree,WLS$Sad4,use="p")
c <- cor(WLS$Degree,WLS$Social4,use="p")
d <- cor(WLS$Degree,WLS$Cog4,use="p")
e <- cor(WLS$Degree,WLS$Marriage,use="p")
f <- cor(WLS$Degree,WLS$Children,use="p")

# Finally, make a plot of these correlation values. The x-axis simply goes
  ↳ from 1 to 13,
# one number per month. Relabel these with dates.

plot(c(1,2,3,4,5), # x values, made into a vector
     c(b,c,d,e,f), # y values, made into a vector
     type = "b", # this means plot (b)oth lines and points
     ylim = c(-0.5,0.5), # make the y-axis go from 0WLS$.5 to 1, you might
  ↳ adjust this
     xlab = "Degree (Associate to Doctorate)",
     ylab = "Correlation Value (r)",
     main = "Correlation of Degree with All",
     xaxt = "n" # this means (n)o x-axis, because we will add our own with the
  ↳ names of the months
     )

text(1.05,-0.14, # same x but y slightly lower
     labels = "-0.101",
     cex = 1,
     col = "purple"
     )

text(2,-0.05, # same x but y slightly lower
     labels = "0.025",
     cex = 1,
     col = "purple"
     )

text(3,-0.04, # same x but y slightly lower
     labels = "0.040",
     cex = 1,
     col = "purple"
     )
```

```

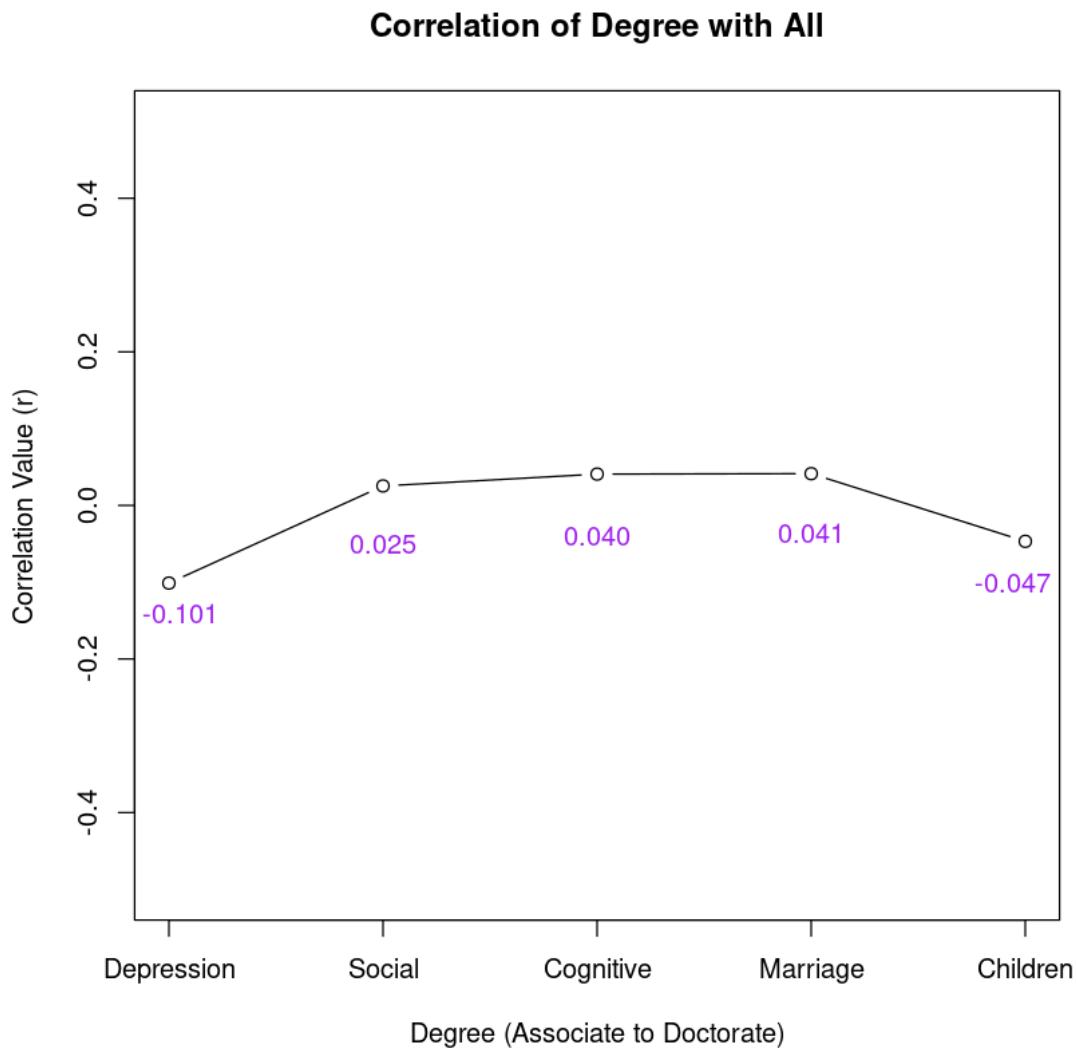
text(4,-0.035, # same x but y slightly lower
     labels = "0.041",
     cex = 1,
     col = "purple"
)

text(4.94,-0.1, # same x but y slightly lower
     labels = "-0.047",
     cex = 1,
     col = "purple"
)

axis(1, # 1 is the bottom axis
     at = c(1,2,3,4,5), # for space reasons, plot just 5 months at intervals
     of three
     label = c("Depression","Social","Cognitive","Marriage","Children")
)

#With more education, you do not get happier are sadder, you do not make more
  children, you do not get smarter,

```



```
[75]: cor(WLS$Sad4,WLS$Children,use="p")
```

```
0.00853522569344083
```

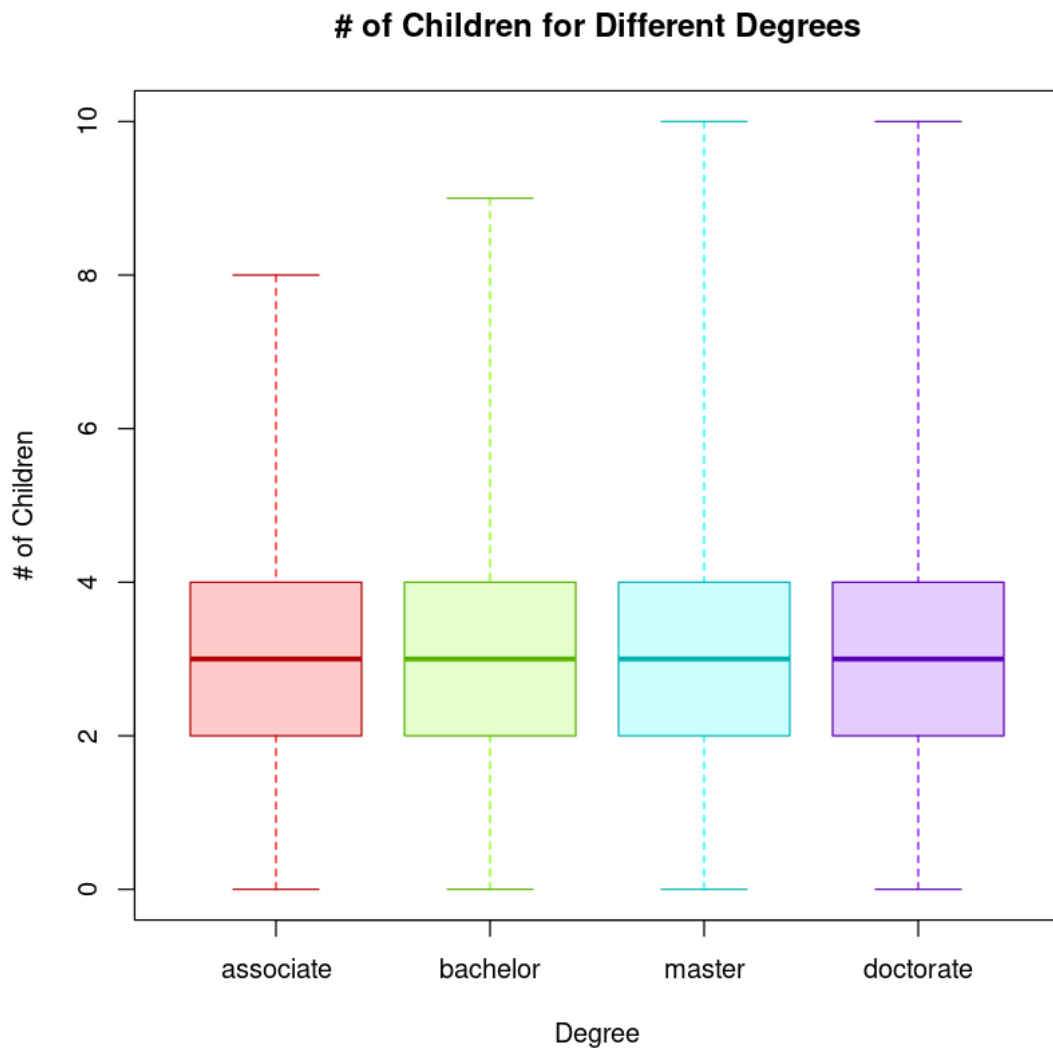
```
[48]: #Just Male Degree vs. Children
```

```
WLS10 <- subset(WLS, Gender == 1)

associate11 <- subset(WLS10, Degree == 1)
bachelor11 <- subset(WLS10, Degree == 2)
master11 <- subset(WLS10, Degree == 3)
doctor11 <- subset(WLS10, Degree == 4)
```

```
c1 <- rainbow(4)
c2 <- rainbow(4, alpha=0.2)
c3 <- rainbow(4, v=0.7)

boxplot(Children~Degree,
        data = WLS10,
        names = c("associate", "bachelor", "master", "doctorate"),
        xlab = "Degree",
        ylab = "# of Children",
        main = "# of Children for Different Degrees",
        range = 8,
        col = c2,
        medcol = c3,
        whiskcol = c1,
        staplecol= c3,
        boxcol = c3,
        outcol = c3,
        pch = 23,
        cex = 2)
```



```
[49]: mean(associate11$Children, na.rm = TRUE)
mean(bachelor11$Children, na.rm = TRUE)
mean(master11$Children, na.rm = TRUE)
mean(doctor11$Children, na.rm = TRUE)

median(associate11$Children, na.rm = TRUE)
median(bachelor11$Children, na.rm = TRUE)
median(master11$Children, na.rm = TRUE)
median(doctor11$Children, na.rm = TRUE)
```

3.14

2.8263358778626

2.78798586572438

2.96575342465753

3

3

3

3

[10]: *#Just Female Degree vs. Children*

```
WLS9 <- subset(WLS, Gender == 2)
```

```
associate6 <- subset(WLS9, Degree == 1)
```

```
bachelor6 <- subset(WLS9, Degree == 2)
```

```
master6 <- subset(WLS9, Degree == 3)
```

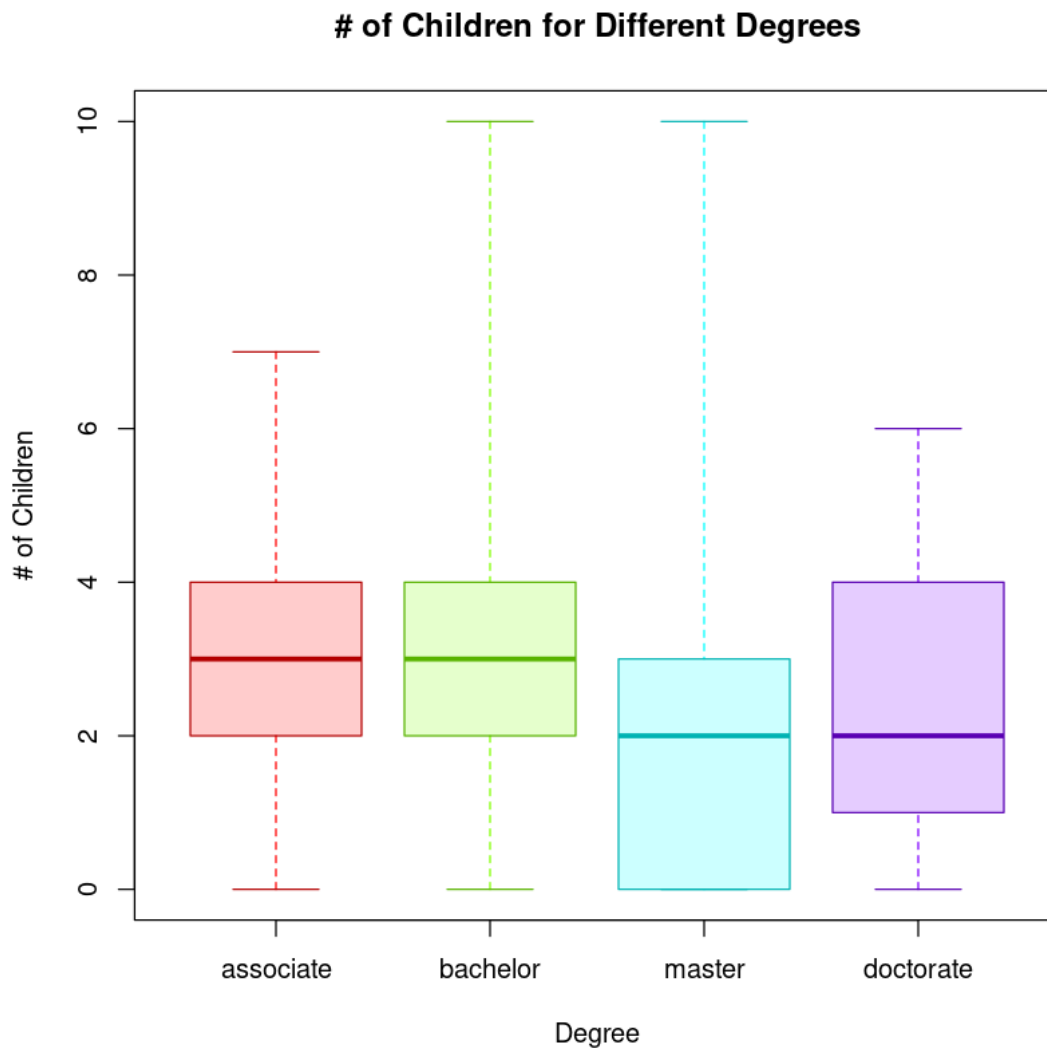
```
doctor6 <- subset(WLS9, Degree == 4)
```

```
c1 <- rainbow(4)
```

```
c2 <- rainbow(4, alpha=0.2)
```

```
c3 <- rainbow(4, v=0.7)
```

```
boxplot(Children~Degree,
        data = WLS9,
        names = c("associate", "bachelor", "master", "doctorate"),
        xlab = "Degree",
        ylab = "# of Children",
        main = "# of Children for Different Degrees",
        range = 8,
        col = c2,
        medcol = c3,
        whiskcol = c1,
        staplecol = c3,
        boxcol = c3,
        outcol = c3,
        pch = 23,
        cex = 2)
```



```
[11]: t.test(master6$Children,bachelor6$Children,
        conf.level = 0.99,
        alternative = "1",
        var.equal = TRUE)

t.test(doctor6$Children,bachelor6$Children,
        conf.level = 0.99,
        var.equal = TRUE)

mean(associate6$Children, na.rm = TRUE)
mean(bachelor6$Children, na.rm = TRUE)
mean(master6$Children, na.rm = TRUE)
mean(doctor6$Children, na.rm = TRUE)
```



```

median(associate6$Children, na.rm = TRUE)
median(bachelor6$Children, na.rm = TRUE)
median(master6$Children, na.rm = TRUE)
median(doctor6$Children, na.rm = TRUE)

```

Two Sample t-test

```

data: master6$Children and bachelor6$Children
t = -5.112, df = 647, p-value = 2.102e-07
alternative hypothesis: true difference in means is less than 0
99 percent confidence interval:
 -Inf -0.4630557
sample estimates:
mean of x mean of y
 1.946154  2.797688

```

Two Sample t-test

```

data: doctor6$Children and bachelor6$Children
t = -0.31664, df = 528, p-value = 0.7516
alternative hypothesis: true difference in means is not equal to 0
99 percent confidence interval:
 -1.478423  1.155774
sample estimates:
mean of x mean of y
 2.636364  2.797688

```

3.28301886792453

2.79768786127168

1.94615384615385

2.63636363636364

3

3

2

2

```

[19]: mean_values <- c(3.283, 2.797, 1.946, 2.636)
      median_values <- c(3,3,2,2)

      mean_values1 <- c(3.14,2.826,2.787,2.965)

```

```

median_values1 <- c(2,2,2,2)

degree_labels <- c("Associate", "Bachelor", "Master", "Doctorate")

plot(c(1, 2, 3, 4), # x values, made into a vector
     mean_values, # y values, the mean values for each degree
     type = "b",   # this means plot (b)oth lines and points
     ylim = c(0, 6), # make the y-axis go from 0 to 10, adjust as needed
     xlim = c(0, 5), # make the x-axis go from 0 to 5
     xlab = "Degree (Associate to Doctorate)",
     ylab = "Mean Number of Children",
     main = "Comparison of Mean # of Children for Men and Women on Degree",
     xaxt = "n",
     col = "purple" # this means (n)o x-axis, because we will add our own with
     ↪ the names of the degrees
)

# Add custom x-axis labels
axis(1, at = c(1, 2, 3, 4), labels = degree_labels)

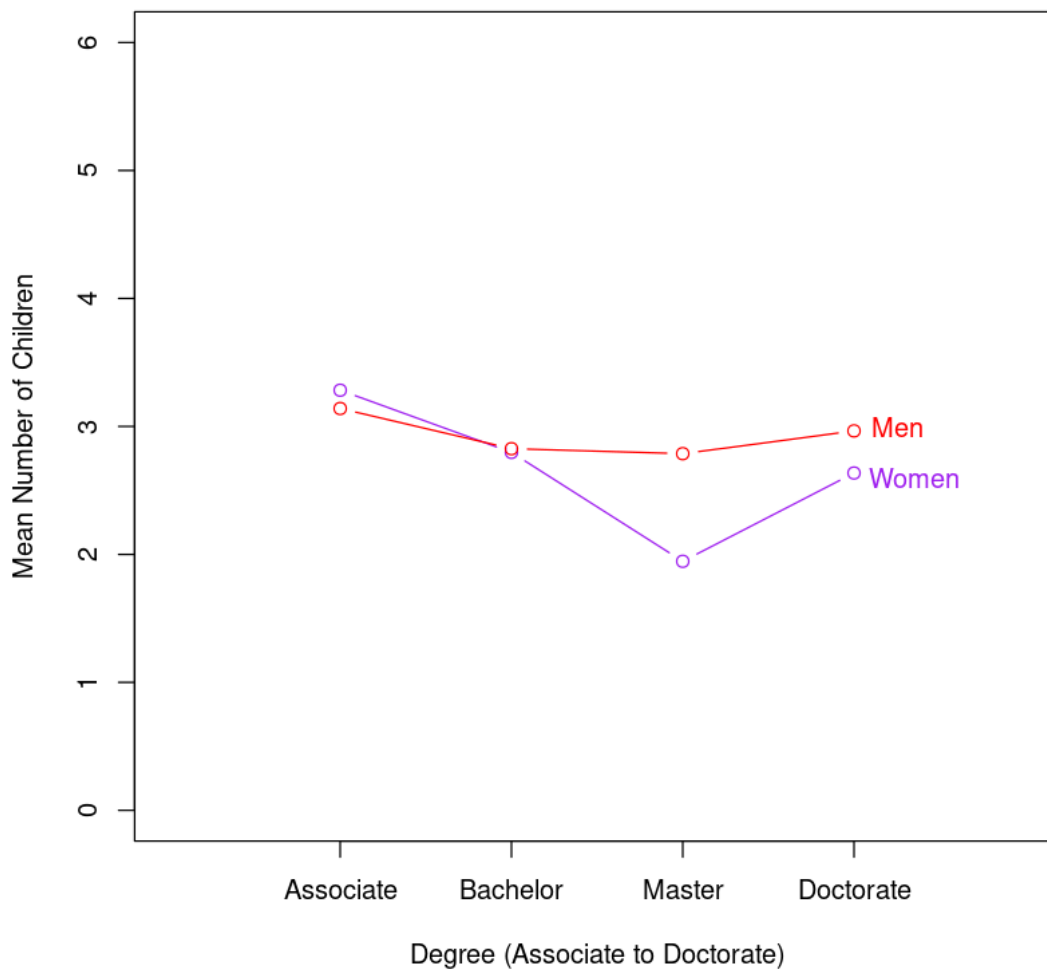
lines(c(1, 2, 3, 4), mean_values1, col = "red", type = "b")
points(c(1, 2, 3, 4), mean_values1, col = "white", pch = 20)

text(4.25,3, # same x but y slightly lower
     labels = "Men",
     cex = 1,
     col = "red"
)

text(4.35,2.6, # same x but y slightly lower
     labels = "Women",
     cex = 1,
     col = "purple"
)

```

Comparison of Mean # of Children for Men and Women on Degree



```
[62]: prop.test(x = c(1.946, 2.767),  
               n = c(10.662, 11.72),  
               conf.level = 0.99,  
               alternative = "1"  
               )
```

Warning message in `prop.test(x = c(1.946, 2.767), n = c(10.662, 11.72),
conf.level = 0.99, :`
"Chi-squared approximation may be incorrect"

2-sample test for equality of proportions with continuity correction

data: c(1.946, 2.767) out of c(10.662, 11.72)

```

X-squared = 8.2279e-31, df = 1, p-value = 0.5
alternative hypothesis: less
99 percent confidence interval:
 -1.000000  0.398767
sample estimates:
   prop 1    prop 2 
0.1825174 0.2360922

```

```

[14]: mean_values3 <- c(3.17, 2.78, 2.63, 3.08)

mean_values4 <- c(3.4,2.8,2.72,3.03)

degree_labels1 <- c("Associate", "Bachelor", "Master", "Doctorate")

plot(c(1, 2, 3, 4), # x values, made into a vector
     mean_values3, # y values, the mean values for each degree
     type = "b",   # this means plot (b)oth lines and points
     ylim = c(0, 6), # make the y-axis go from 0 to 10, adjust as needed
     xlim = c(0, 5), # make the x-axis go from 0 to 5
     xlab = "Degree (Associate to Doctorate)",
     ylab = "Mean Number of Depression Symptoms",
     main = "Upper and Lower Social Interaction with Degree on Depression",
     xaxt = "n",
     col = "green" # this means (n)o x-axis, because we will add our own with
     ↪the names of the degrees
)

# Add custom x-axis labels
axis(1, at = c(1, 2, 3, 4), labels = degree_labels1)

lines(c(1, 2, 3, 4), mean_values4, col = "red", type = "b")
points(c(1, 2, 3, 4), mean_values4, col = "white", pch = 20)

text(4.25,3, # same x but y slightly lower
     labels = "Men",
     cex = 1,
     col = "green"
)

text(4.35,2.6, # same x but y slightly lower
     labels = "Women",
     cex = 1,
     col = "red"
)

```

)

Upper and Lower Social Interaction with Degree on Depression

