Brooks-Lab1

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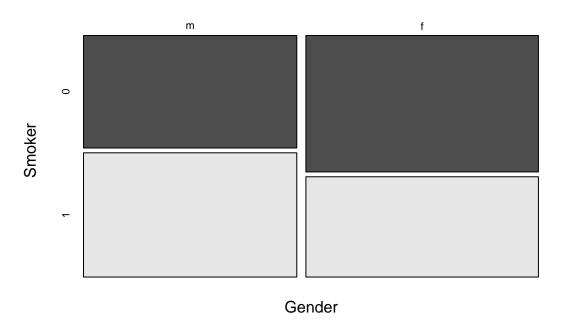
February 6, 2016

```
#Show the top few rows of the matrix
head(cdc)
      genhlth exerany hlthplan smoke100 height weight wtdesire age gender
## 1
                                                       175 77
        good
                 0
                                         70
                                               175
                          1
                                   0
## 2
        good
                  0
                          1
                                   1
                                         64
                                              125
                                                       115 33
                                  1 60 105
## 3
                  1
                                                                   f
        good
                          1
                                                      105 49
## 4
        good
                  1
                          1
                                  0 66 132
                                                      124 42
                      1
                                 0 61
                                               150
                                                       130 55
                                                                   f
## 5 very good
                  0
## 6 very good
                                  0
                                         64
                                               114
                                                       114 55
#Show the names of each column in the matrix
names(cdc)
## [1] "genhlth" "exerany" "hlthplan" "smoke100" "height"
                                                         "weight"
## [7] "wtdesire" "age"
                          "gender"
#Show the length (Num of cases) of an individual column
length(cdc$genhlth)
## [1] 20000
#Show the width (Num of Variables) of the overall matrix
length(cdc)
## [1] 9
#There are 20,000 cases within this data set.
#There are 9 different variables in this data set
#five number summary of height
summary(cdc$height)
##
     Min. 1st Qu. Median
                         Mean 3rd Qu.
                                          Max.
    48.00 64.00 67.00
                          67.18 70.00
                                         93.00
##
#IQR for height
IQRH <- 70.00 - 64.00
IQRH
```

[1] 6

```
#five number summary of age
summary(cdc$age)
##
     Min. 1st Qu. Median
                              Mean 3rd Qu.
                                              Max.
##
     18.00 31.00 43.00
                             45.07 57.00
                                             99.00
#IQR for age
IQRA <- 57.00 - 31.00
IQRA
## [1] 26
#Relative Distrisution of gender and health rating
table(cdc$gender, cdc$exerany)
##
##
          0
               1
    m 2149 7420
##
    f 2937 7494
##
#total number of males in the study
TotMales <- 2149 + 7420
TotMales
## [1] 9569
#proportion of sample in excellent health
TotalPeople \leftarrow 2149 + 7420 + 2937 + 7494
TotalPeople
## [1] 20000
#Excellent Health
ExHealth < -7420 + 7494
#Proportion of participants in excellent health
ExProp <- ExHealth / TotalPeople</pre>
ExProp
## [1] 0.7457
#Create a mosaic plot of smoking habits Vs gender
mosaicplot(table(cdc$gender,cdc$smoke100), xlab = 'Gender', ylab = 'Smoker', color = TRUE)
```

table(cdc\$gender, cdc\$smoke100)



#It appears that males have more of a smoking habit than females.

#Subset the CDC matrix to only people uder the age of 23 and smoke more than 100 cigarettes
under23_and_smoke <- subset(cdc, age < 23 & smoke100 == 1)
head(under23_and_smoke)</pre>

```
genhlth exerany hlthplan smoke100 height weight wtdesire age gender
## 13 excellent
                       1
                                0
                                         1
                                                     185
                                                              220
                                                                   21
                                0
## 37 very good
                       1
                                         1
                                               70
                                                     160
                                                              140 18
## 96 excellent
                       1
                                1
                                         1
                                                     175
                                                              200 22
                                               64
                                                     190
                                                                           f
## 180
            good
                       1
                                1
                                         1
                                                              140 20
## 182 very good
                       1
                                1
                                         1
                                               62
                                                      92
                                                               92 21
                                                                           f
                                0
                                               64
## 240 very good
                                                     125
                                                              115 22
```

dim(under23_and_smoke)

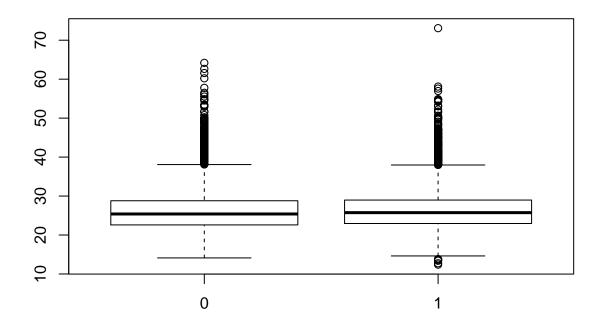
```
## [1] 620 9
```

```
bmi <- (cdc$weight / cdc$height^2) * 703
boxplot(bmi ~ cdc$genhlth)</pre>
```



 $\#This\ boxplot\ is\ showing\ the\ BMI\ (Body\ Mass\ Index)\ of\ the\ participants\ in\ the\ case\ study\ \#compared\ to\ their\ general\ overall\ health.$ Each box plot\ shows the BMI\ of\ the\ participants\ \#but\ it\ is\ serparated\ out\ by\ their\ general\ health.

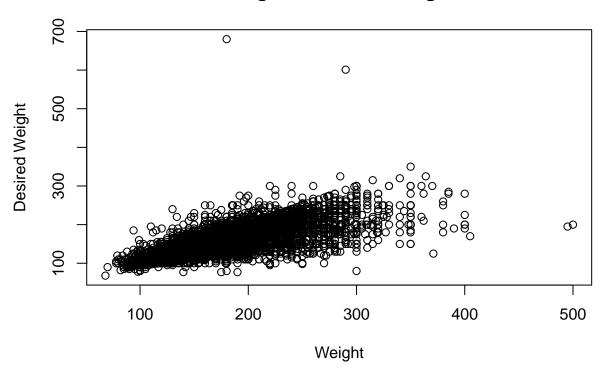
boxplot(bmi ~ cdc\$smoke100)



#I chose to compare the BMI and amount of cigraettes in a liftime
#I chose that because I have heard that smoking can cause mass amounts of weight
#loss. Poeple loo sickly and malnurished. I thought I was going to see the mean
#very close to zero. Instead I am seeing that both of the graphs look almost
#indentical. That was not what I was expecting.

```
plot(cdc$weight, cdc$wtdesire, title(main = 'Weight Vs Desired Weight'), xlab = 'Weight',
    ylab = 'Desired Weight')
```

Weight Vs Desired Weight



#The two variables seem to have a linear relationship. Most of the people look to be about 100 pounds #less than their current weight. It appears that most of the people would like to lose the same #amount.

```
wdiff <- (cdc$weight - cdc$wtdesire)
str(wdiff)</pre>
```

int [1:20000] 0 10 0 8 20 0 9 10 20 10 ...

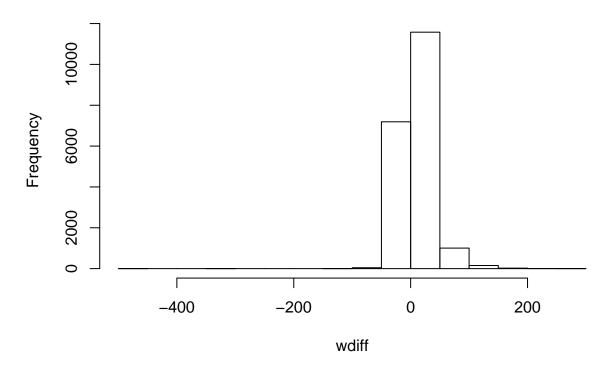
```
#The new object wdiff is and integer data type
#If widff is 0 then the person is already at their desired weight
#If widff is negative then their desired weight is more than their current weight
#If widff is positive than their current wieght is larger than their desired weight

summary(wdiff)
```

```
## Min. 1st Qu. Median Mean 3rd Qu. Max.
## -500.00 0.00 10.00 14.59 21.00 300.00
```

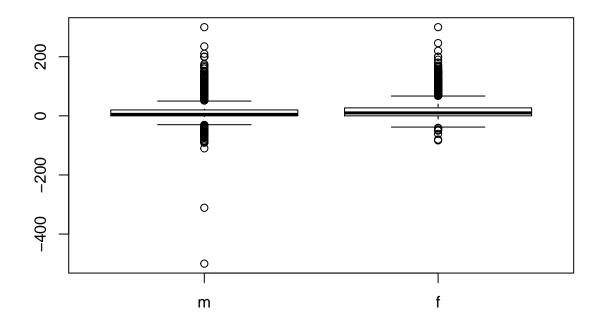
```
hist(wdiff)
```

Histogram of wdiff



#It actually appears to be a pretty normal distribution. The graph is centered around 0 #which means that a lot of the people are happy with their current weight. There are a #few outliers, but overall people are happy with their weight or want to make very #minute changes.

boxplot(wdiff ~ cdc\$gender)



#It appears than me actually want to gain some weight, while women are more about losing it.
#I am guessing women want to lose weight to keep a sloim figure, while men probably want
#to gain more muscle.

mean(cdc\$weight)

[1] 169.683

sd(cdc\$weight)

[1] 40.08097

```
PosSd <- sd(cdc$weight) + mean(cdc$weight)
NegSd <- -sd(cdc$weight) + mean(cdc$weight)
OneSDTotal <- subset(cdc, cdc$weight > NegSd & cdc$weight < PosSd)
dim(OneSDTotal)</pre>
```

[1] 14152 9

PropOneSD <- 14152/20000 PropOneSD

[1] 0.7076