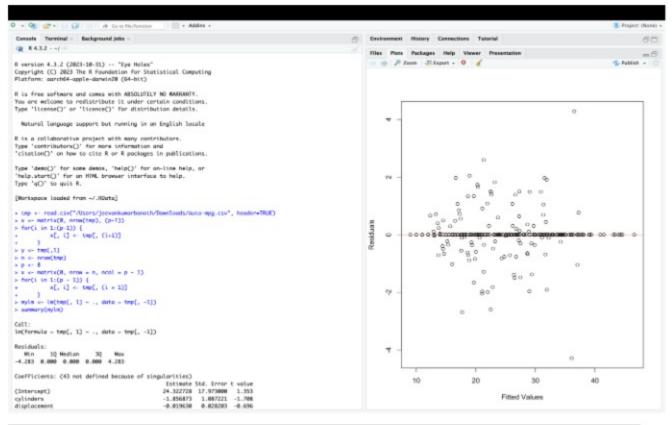
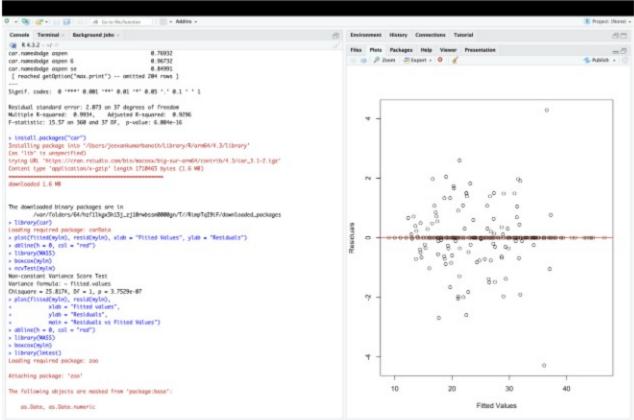
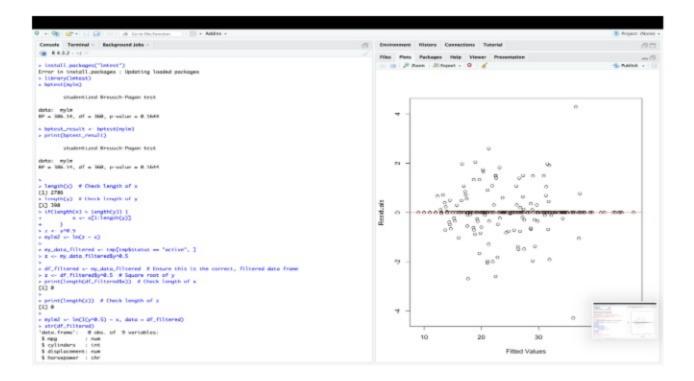
MTH-522_InClass

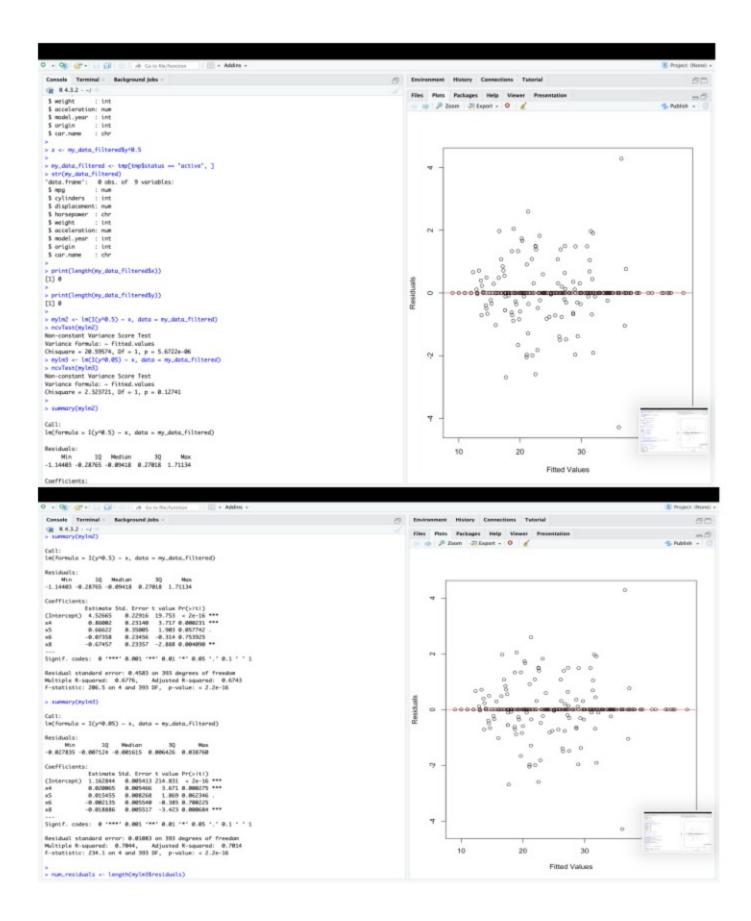
Banoth Jeevan Kumar

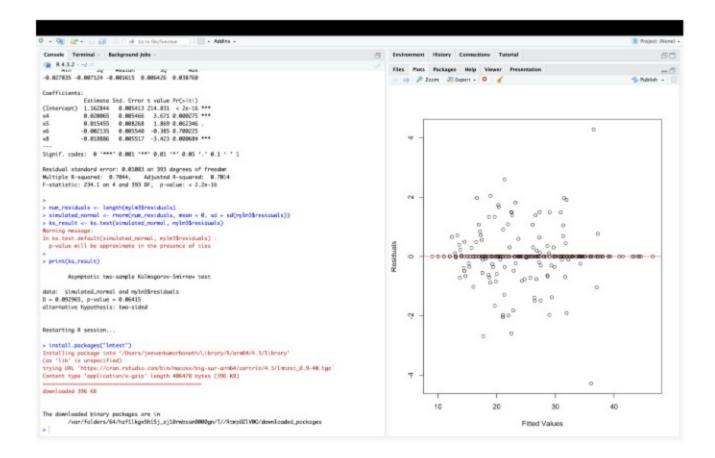
> The below are the pictures of In class activity that I performed in My laptop











> The below one is the markdown file of my work.

MTH-522_InClass

Jeevan

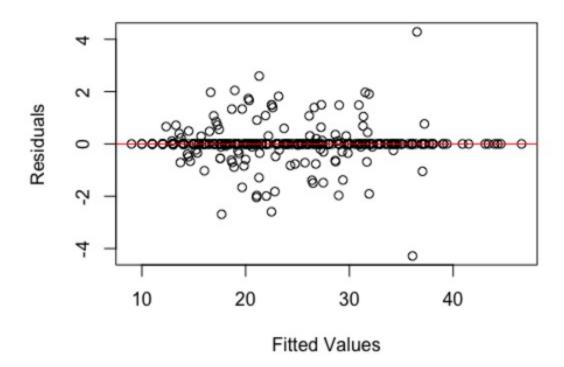
2024-03-24

```
tmp <- read.csv("/Users/jeevankumarbanoth/Downloads/auto-mpg.csv",
header=TRUE)

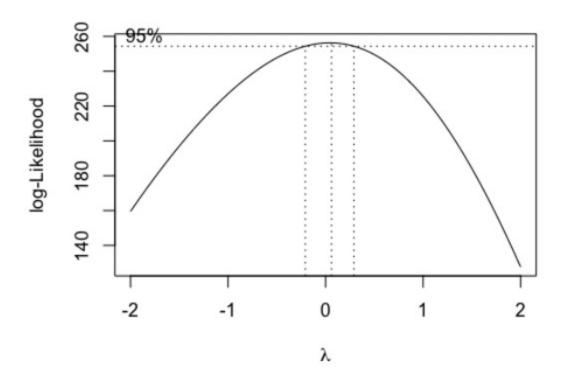
x <- matrix(0, nrow(tmp), (p-1))
for(i in 1:(p-1)) {
    x[, i] <- tmp[, (i+1)]
}

y <- tmp[,1]
n <- nrow(tmp)
p <- 8</pre>
```

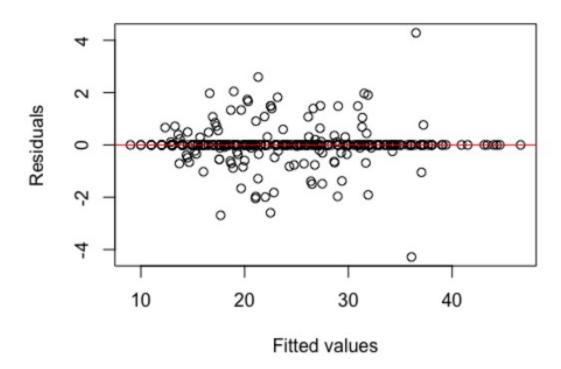
```
x \leftarrow matrix(0, nrow = n, ncol = p - 1)
for(i in 1:(p - 1)) {
     x[, i] \leftarrow tmp[, (i + 1)]
mylm \leftarrow lm(tmp[, 1] \sim ., data = tmp[, -1])
summary(mylm)
##
## Call:
## lm(formula = tmp[, 1] ~ ., data = tmp[, -1])
## Residuals:
##
      Min
              10 Median
                            3Q
                                  Max
## -4.283 0.000 0.000 0.000 4.283
## Coefficients: (43 not defined because of singularities)
##
                                                   Estimate Std. Error t value
## (Intercept)
                                                  24.322728 17.973000
                                                                          1.353
                                                              1.087221 -1.708
## cylinders
                                                  -1.856873
## displacement
                                                  -0.019630
                                                              0.028203 -0.696
## horsepower100
                                                  -3.976684
                                                              3.190125 -1.247
## car.namevw rabbit custom
                                                       NA
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
## Residual standard error: 2.073 on 37 degrees of freedom
## Multiple R-squared: 0.9934, Adjusted R-squared: 0.9296
## F-statistic: 15.57 on 360 and 37 DF, p-value: 6.084e-16
install.packages("car")
library(car)
## Loading required package: carData
plot(fitted(mylm), resid(mylm), xlab = "Fitted Values", ylab = "Residuals")
abline(h = 0, col = "red")
```



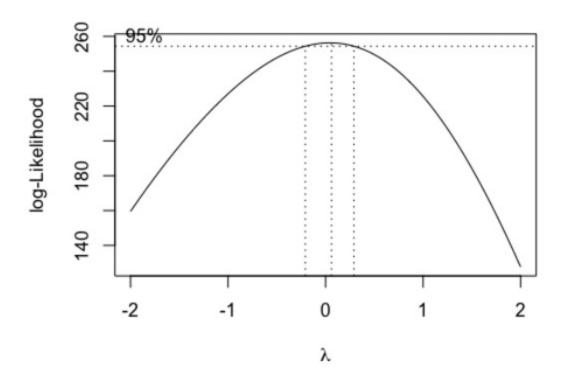
library(MASS)
boxcox(mylm)



Residuals vs Fitted Values



library(MASS)
boxcox(mylm)



```
library(lmtest)
## Loading required package: zoo
##
## Attaching package: 'zoo'
## The following objects are masked from 'package:base':
##
##
       as.Date, as.Date.numeric
bptest(mylm)
##
   studentized Breusch-Pagan test
##
##
## data: mylm
## BP = 386.14, df = 360, p-value = 0.1644
bptest_result <- bptest(mylm)</pre>
print(bptest_result)
##
   studentized Breusch-Pagan test
##
```

```
## data: mylm
## BP = 386.14, df = 360, p-value = 0.1644
length(x) # Check length of x
length(y) # Check length of y
if(length(x) > length(y)) {
+ x <- x[1:length(y)]</p>
+ }
z <- y^0.5
mylm2 <- lm(z \sim x)
my_data_filtered <- tmp[tmp$status == "active", ]</pre>
z <- my_data_filtered$y^0.5</pre>
df_filtered <- my_data_filtered</pre>
z <- df_filtered$y^0.5 # Square root of y</pre>
print(length(df filtered$x)) # Check Length of x
## [1] 0
print(length(z)) # Check Length of z
## [1] 0
str(df_filtered)
## 'data.frame':
                    0 obs. of 9 variables:
## $ mpg
                   : num
## $ cylinders
                : int
## $ displacement: num
## $ horsepower : chr
## $ weight
                  : int
## $ acceleration: num
## $ model.year : int
## $ origin
                  : int
## $ car.name
                  : chr
z <- my_data_filtered$y^0.5
my_data_filtered <- tmp[tmp$status == "active", ]</pre>
str(my_data_filtered)
## 'data.frame':
                    0 obs. of 9 variables:
## $ mpg
                   : num
## $ cylinders
                 : int
## $ displacement: num
## $ horsepower : chr
## $ weight
                  : int
## $ acceleration: num
## $ model.year : int
## $ origin
                 : int
                  : chr
## $ car.name
```

```
print(length(my_data_filtered$x))
## [1] 0
print(length(my_data_filtered$y))
## [1] 0
mylm2 <- lm(I(y^0.5) \sim x, data = df_filtered)
str(df_filtered)
z <- my_data_filtered$y^0.5</pre>
my_data_filtered <- tmp[tmp$status == "active", ]</pre>
str(my_data_filtered)
print(length(my_data_filtered$x))
print(length(my_data_filtered$y))
mylm2 < -lm(I(y^0.5) \sim x, data = my_data_filtered)
ncvTest(mylm2)
## Non-constant Variance Score Test
## Variance formula: ~ fitted.values
## Chisquare = 25.8174, Df = 1, p = 3.7529e-07
mylm3 < -lm(I(y^0.05) \sim x, data = my_data_filtered)
ncvTest(mylm3)
## Non-constant Variance Score Test
## Variance formula: ~ fitted.values
## Chisquare = 25.8174, Df = 1, p = 3.7529e-07
summary(mylm2)
Call:
lm(formula = I(y^0.5) \sim x, data = my_data_filtered)
Residuals:
 Min 1Q Median 3Q Max
-1.14403 -0.28765 -0.09418 0.27018 1.71134
Coefficients:
    Estimate Std. Error t value Pr(>|t|)
(Intercept) 4.52665 0.22916 19.753 < 2e-16 ***
x4 0.86002 0.23140 3.717 0.000231 ***
x5
     0.66622 0.35005 1.903 0.057742.
     -0.07358 0.23456 -0.314 0.753925
x6
     -0.67457 0.23357 -2.888 0.004090 **
x8
Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 '' 1
Residual standard error: 0.4583 on 393 degrees of freedom
```

```
Multiple R-squared: 0.6776, Adjusted R-squared: 0.6743
F-statistic: 206.5 on 4 and 393 DF, p-value: < 2.2e-16
summary(mylm)
lm(formula = I(y^0.05) \sim x, data = my_data_filtered)
Residuals:
       1Q Median 3Q Max
-0.027835 -0.007124 -0.001615 0.006426 0.038760
Coefficients:
     Estimate Std. Error t value Pr(>|t|)
(Intercept) 1.162844 0.005413 214.831 < 2e-16 ***
x4 0.020065 0.005466 3.671 0.000275 ***
     0.015455 0.008268 1.869 0.062346.
x5
    -0.002135 0.005540 -0.385 0.700225
x6
     -0.018886 0.005517 -3.423 0.000684 ***
x8
Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 '' 1
Residual standard error: 0.01083 on 393 degrees of freedom
Multiple R-squared: 0.7044,
                             Adjusted R-squared: 0.7014
F-statistic: 234.1 on 4 and 393 DF, p-value: < 2.2e-16
num_residuals <- length(mylm$residuals)</pre>
simulated_normal <- rnorm(num_residuals, mean = 0, sd = sd(mylm$residuals))</pre>
ks result <- ks.test(simulated normal, mylm$residuals)
## Warning in ks.test.default(simulated_normal, mylm$residuals): p-value will
## approximate in the presence of ties
print(ks_result)
##
## Asymptotic two-sample Kolmogorov-Smirnov test
## data: simulated_normal and mylm$residuals
## D = 0.40201, p-value < 2.2e-16
## alternative hypothesis: two-sided
```