

# Homework #12

asm366

Things to complete:

- a. All examples completed by the TAs
- b. All of the homework problems assigned to you

Refer to **Homework & Discussion Submission Rules** on Canvas.

Refer to Gradescope for the Due Date and Due time.

If you needed to download a \*.RDATA file with this lab, make sure it was downloaded into the same folder as this file.

Additionally, run the first code chunk in this file before you start doing any work. By running the first code chunk, some variables will be loaded for you. You should use these variables in your work, when necessary.

Do not assign/select the cover page in GRADESCOPE.

Example 1:

```
# SHOW YOUR WORK HERE.
x.values <- seq(from = 0, to = 4, by = 0.05)
ln <- length(x.values)
errors <- rnorm(n = ln, sd = 1)

y.values <- (x.values - 1) * (x.values - 2) * (x.values - 3) + errors

simulated.dat <- data.frame(x.values, y.values)

plot(simulated.dat)

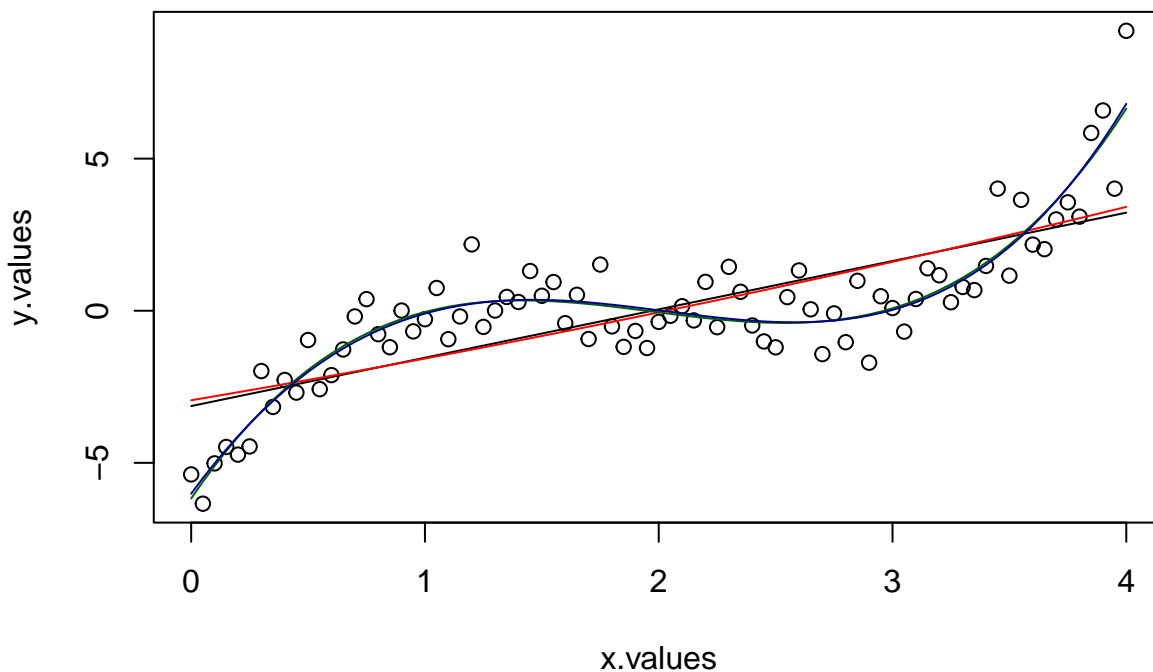
lm.fit <- lm(y.values ~ x.values, dat = simulated.dat)

quadratic.fit <- lm(y.values ~ x.values + I(x.values^2), dat = simulated.dat)

cubic.fit <- lm(y.values ~ x.values + I(x.values^2) + I(x.values^3), dat = simulated.dat)

quatric.fit <- lm(y.values ~ x.values + I(x.values^2) + I(x.values^3) +
  I(x.values^4), dat = simulated.dat)

points(x.values, predict(lm.fit), col = "black", type = "l")
points(x.values, predict(quadratic.fit), col = "red", type = "l")
points(x.values, predict(cubic.fit), col = "darkgreen", type = "l")
points(x.values, predict(quatric.fit), col = "darkblue", type = "l")
```



```
summary(quatric.fit)
##
## Call:
## lm(formula = y.values ~ x.values + I(x.values^2) + I(x.values^3) +
##     I(x.values^4), data = simulated.dat)
##
```

```
## Residuals:
##      Min       1Q   Median       3Q      Max
## -2.1705 -0.7139 -0.1297  0.6322  2.4007
##
## Coefficients:
##              Estimate Std. Error t value Pr(>|t|)
## (Intercept)   -6.01307    0.48857  -12.308 < 2e-16 ***
## x.values       10.45885    1.71478   6.099 4.13e-08 ***
## I(x.values^2)  -5.25804    1.75905  -2.989 0.00377 **
## I(x.values^3)   0.67418    0.66271   1.017 0.31223
## I(x.values^4)   0.04672    0.08217   0.569 0.57136
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 0.9453 on 76 degrees of freedom
## Multiple R-squared:  0.8609, Adjusted R-squared:  0.8536
## F-statistic: 117.6 on 4 and 76 DF,  p-value: < 2.2e-16

# FINISHED? KNIT YOUR WORK BEFORE GOING TO THE NEXT QUESTION.

## REVIEW Homework & Discussion Submission Rules ON CANVAS.

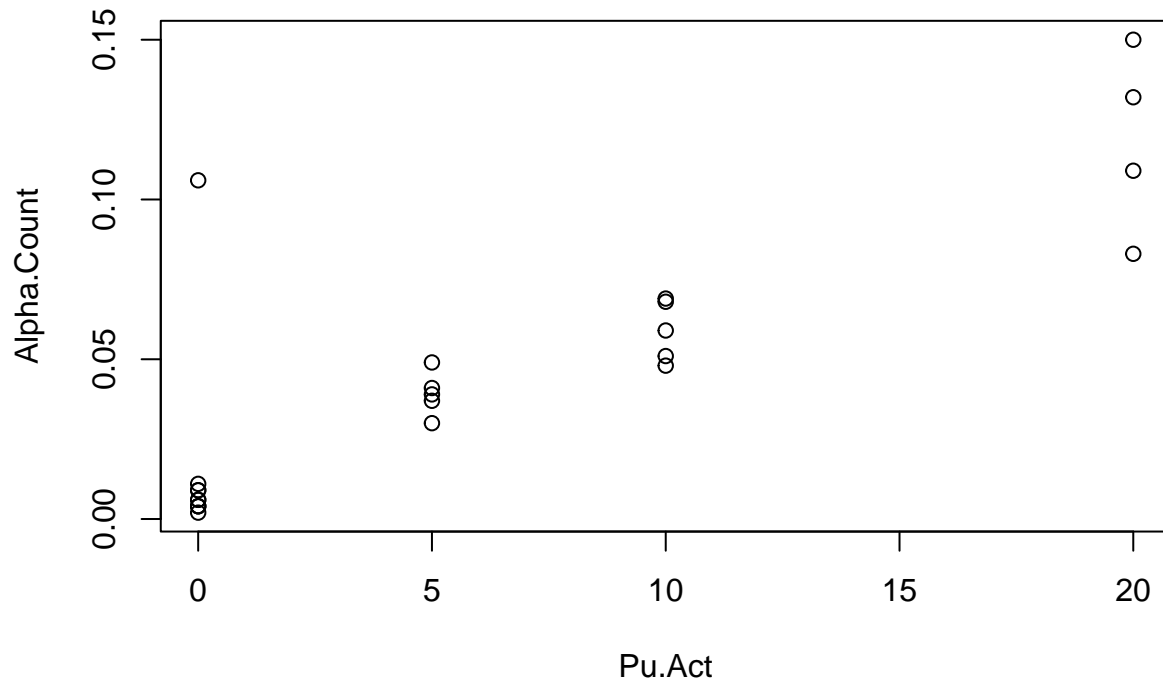
### THIS SHOULD BE SUBMITTED BEFORE 5:00PM ON THE DUE DATE LISTED ON
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```

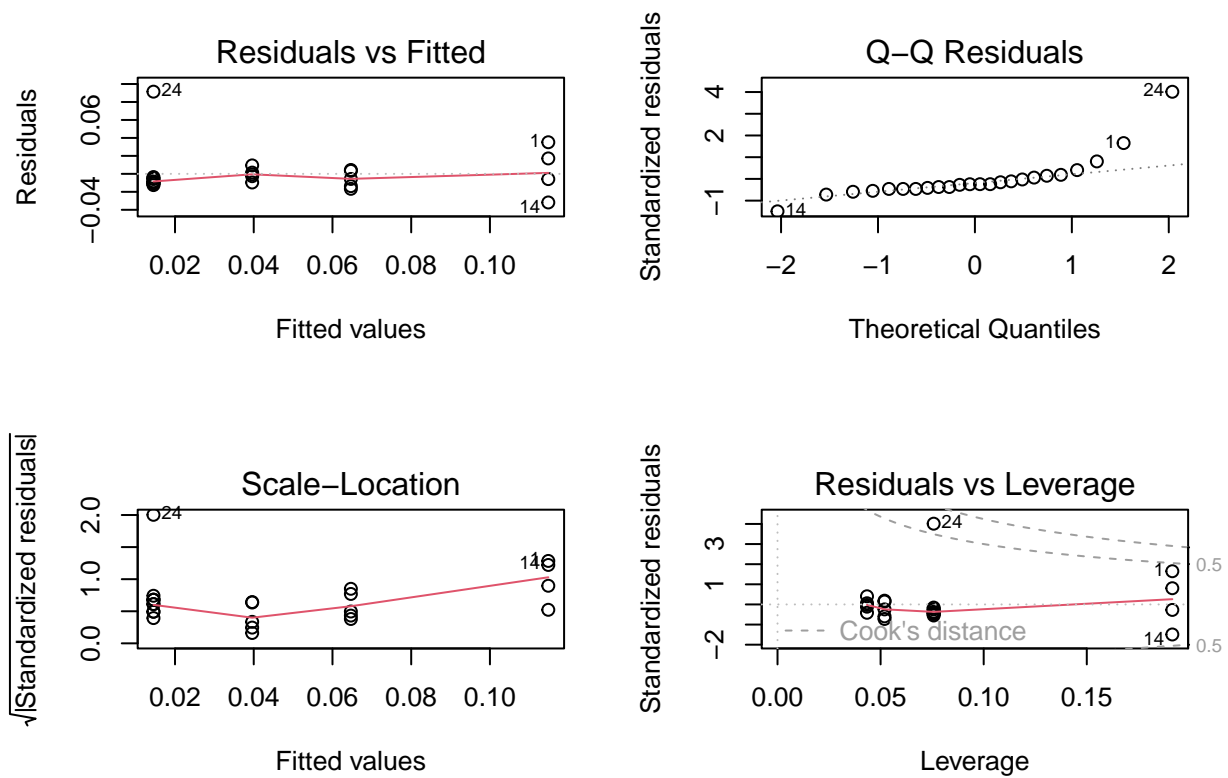
Example 2:

*# SHOW YOUR WORK HERE.*

```
pluto <- read.csv("Plutonium.csv")
pluto.fit <- lm(Alpha.Count ~ Pu.Act, dat = pluto)
plot(pluto[2:1])
```

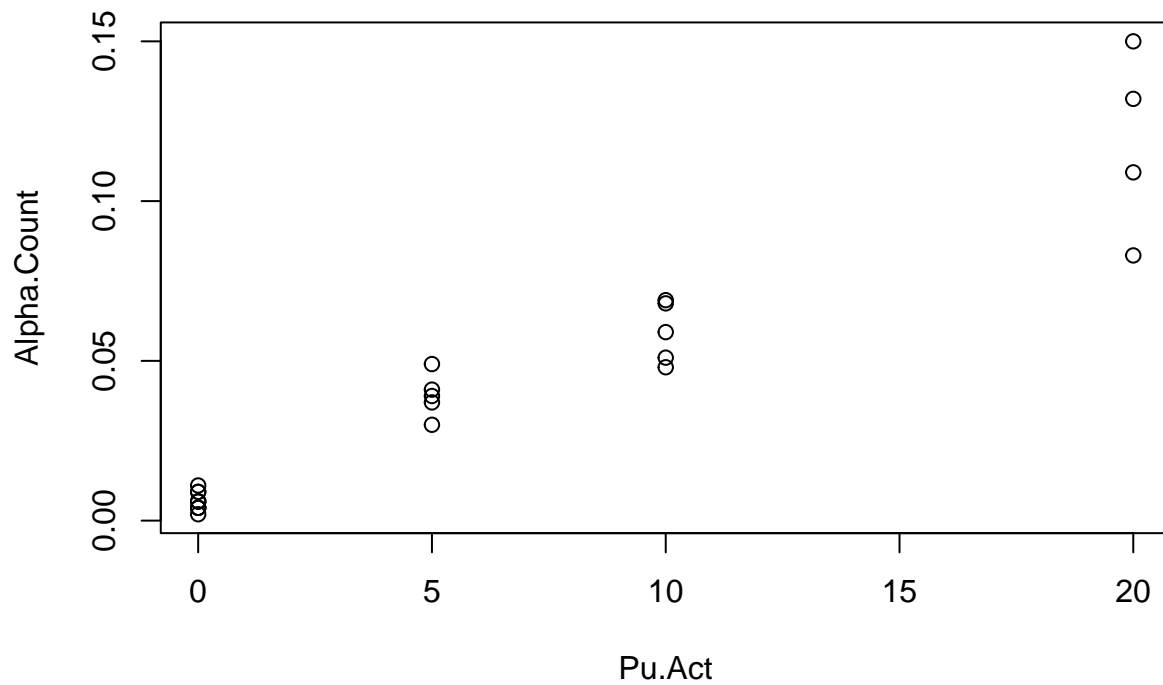


```
par(mfrow = c(2, 2))
plot(pluto.fit)
```

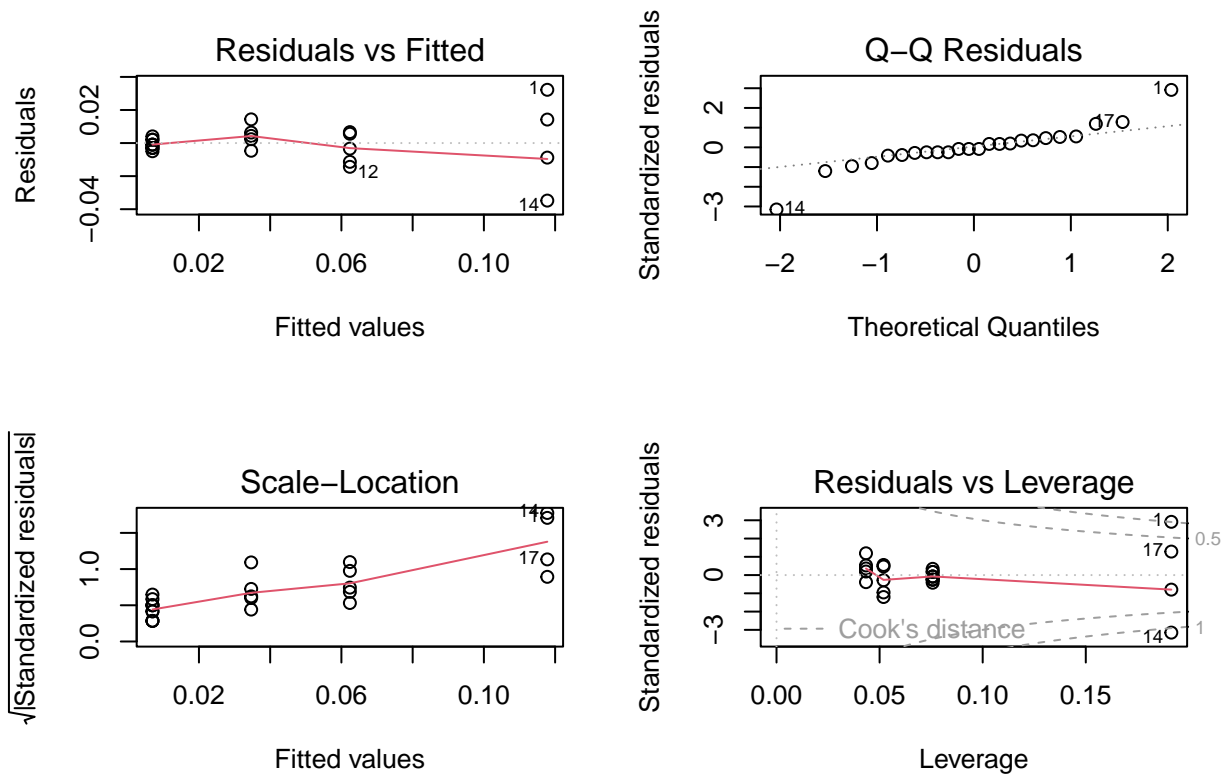


```
pluto[24, 1]
## [1] 0.106

pluto[24, 1] <- 0.006
pluto.fit <- lm(Alpha.Count ~ Pu.Act, dat = pluto)
par(mfrow = c(1, 1))
plot(pluto[2:1])
```



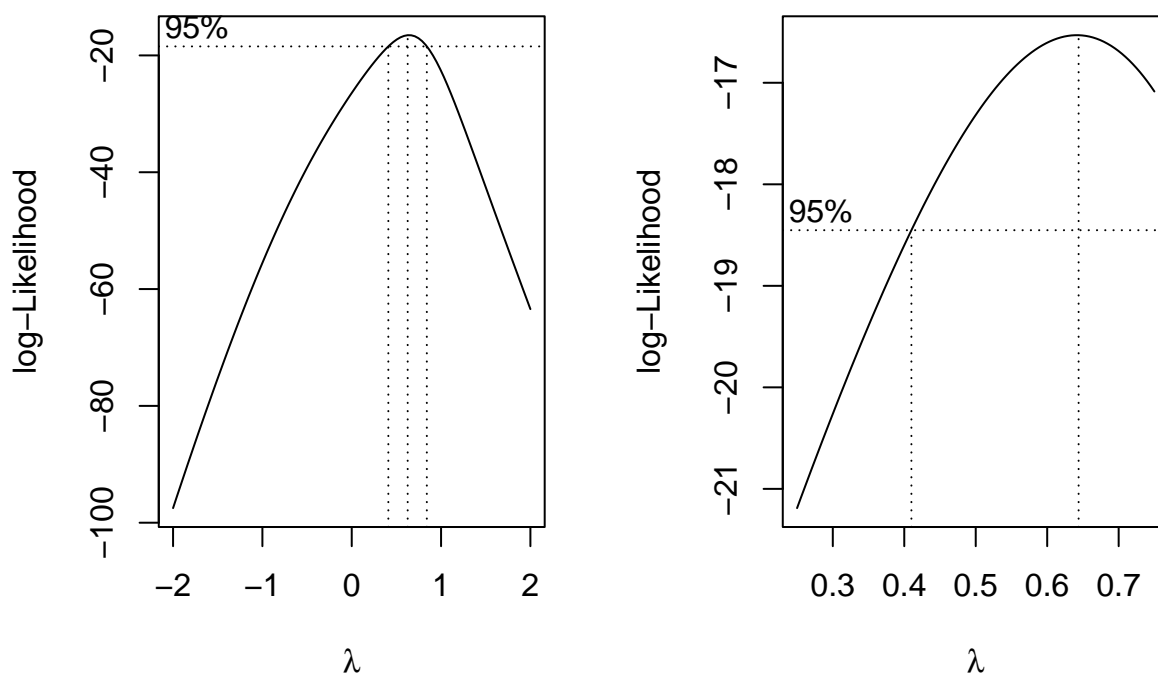
```
par(mfrow = c(2, 2))
plot(pluto.fit)
```



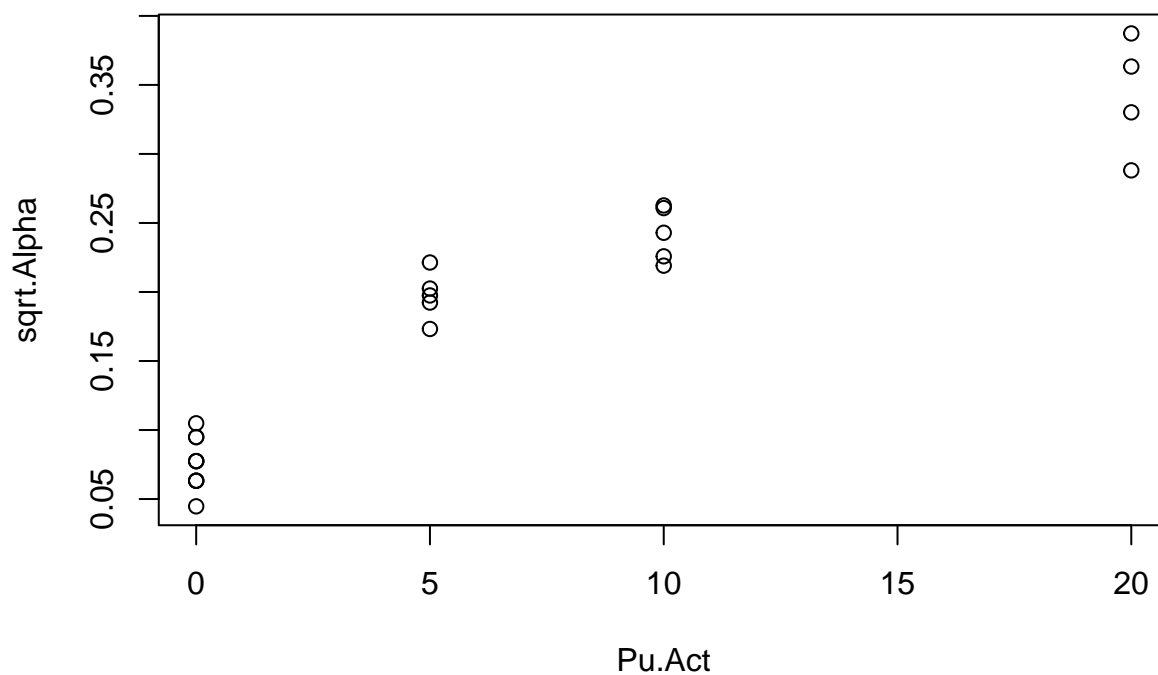
```
library(MASS)

par(mfrow = c(1, 2))
boxcox(pluto.fit, lambda = seq(-2, 2, 1/10))
```

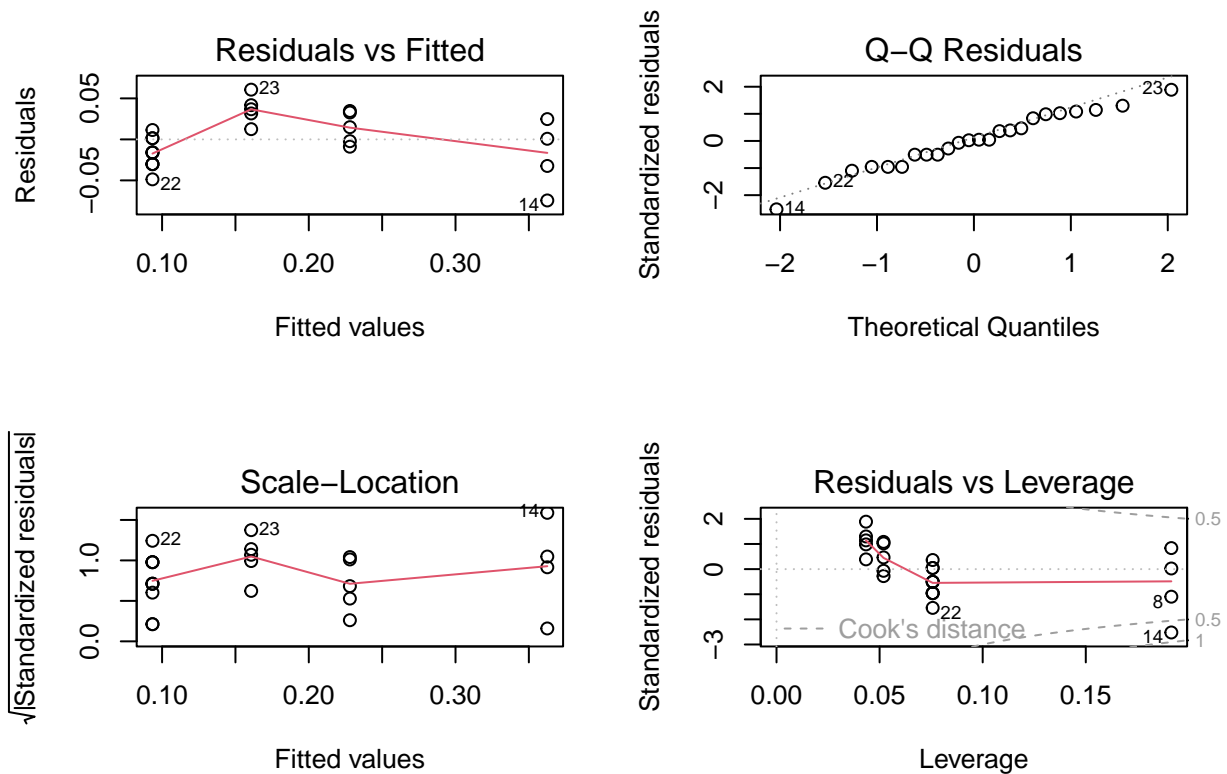
```
boxcox(pluto.fit, lambda = seq(0.25, 0.75, 1/20))
```



```
pluto$sqrt.Alpha <- sqrt(pluto$Alpha.Count)
pluto.fit.sqrt <- lm(sqrt.Alpha ~ Pu.Act, data = pluto)
par(mfrow = c(1, 1))
plot(pluto[2:3])
```



```
par(mfrow = c(2, 2))
plot(pluto.fit.sqrt)
```



```
pluto$sqrt.Pu <- sqrt(pluto$Pu.Act)
pluto.fit.both <- lm(sqrt.Alpha ~ sqrt.Pu, data = pluto)
par(mfrow = c(1, 1))
pluto.fit.both
##
## Call:
## lm(formula = sqrt.Alpha ~ sqrt.Pu, data = pluto)
##
## Coefficients:
## (Intercept)      sqrt.Pu
##      0.07341      0.05719

summary(pluto.fit.both)
##
## Call:
## lm(formula = sqrt.Alpha ~ sqrt.Pu, data = pluto)
##
## Residuals:
##      Min       1Q   Median       3Q      Max
## -0.041067 -0.010463  0.001092  0.011335  0.058134
##
## Coefficients:
##              Estimate Std. Error t value Pr(>|t|)
## (Intercept)  0.073410   0.007300   10.06 1.09e-09 ***
## sqrt.Pu      0.057188   0.002872   19.91 1.47e-15 ***
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 0.02422 on 22 degrees of freedom
```



```
## Multiple R-squared:  0.9474, Adjusted R-squared:  0.945  
## F-statistic: 396.4 on 1 and 22 DF,  p-value: 1.465e-15
```

```
# FINISHED? KNIT YOUR WORK BEFORE GOING TO THE NEXT QUESTION.
```

```
## REVIEW Homework & Discussion Submission Rules ON CANVAS.
```

```
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```

```
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```

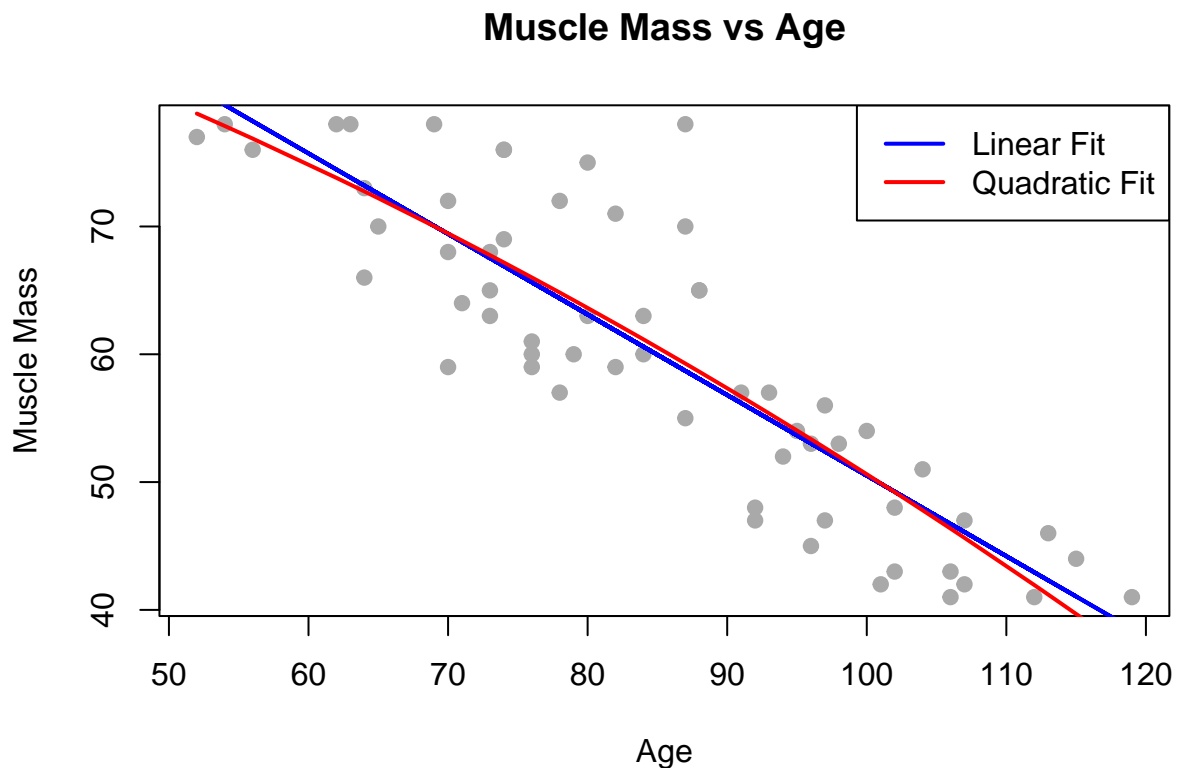
Problem 3:

```
data <- read.csv("muscleMass.csv")
model1 <- lm(mass ~ age, data = data)

model2 <- lm(mass ~ age + I(age^2), data = data)

plot(data$age, data$mass, main = "Muscle Mass vs Age", xlab = "Age", ylab = "Muscle Mass",
     pch = 19, col = "darkgray")

lines(data$age, fitted(model1), col = "blue", lwd = 2)
lines(sort(data$age), fitted(model2)[order(data$age)], col = "red", lwd = 2)
legend("topright", legend = c("Linear Fit", "Quadratic Fit"), col = c("blue",
    "red"), lwd = 2)
```



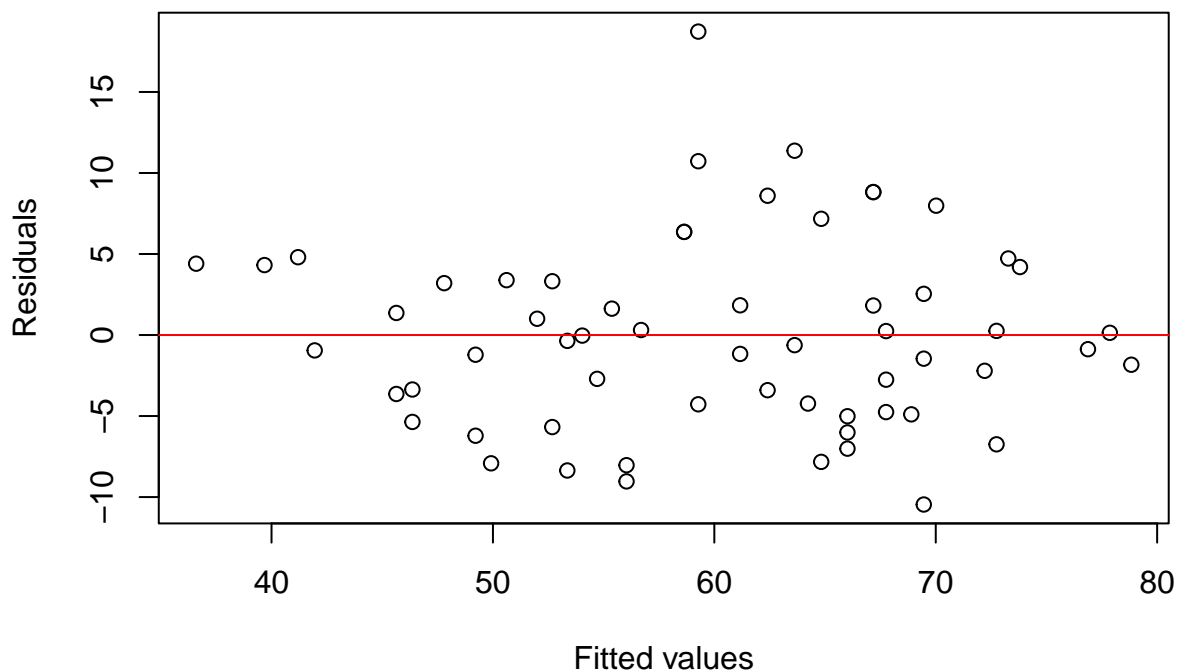
```
summary(model2)
##
## Call:
## lm(formula = mass ~ age + I(age^2), data = data)
##
## Residuals:
##      Min       1Q   Median       3Q      Max
## -10.4593  -4.3954  -0.4927   3.5878  18.7248
##
## Coefficients:
##              Estimate Std. Error t value Pr(>|t|)
## (Intercept)  97.768763   19.364306   5.049 4.86e-06 ***
## age         -0.247811    0.461302  -0.537   0.593
## I(age^2)     -0.002237    0.002684  -0.834   0.408
## ---
```

```
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 5.964 on 57 degrees of freedom
## Multiple R-squared:  0.7531, Adjusted R-squared:  0.7444
## F-statistic: 86.92 on 2 and 57 DF,  p-value: < 2.2e-16

anova(model1, model2)
## Analysis of Variance Table
##
## Model 1: mass ~ age
## Model 2: mass ~ age + I(age^2)
##   Res.Df    RSS Df Sum of Sq   F Pr(>F)
## 1      58 2052.2
## 2      57 2027.5  1    24.722 0.695 0.4079

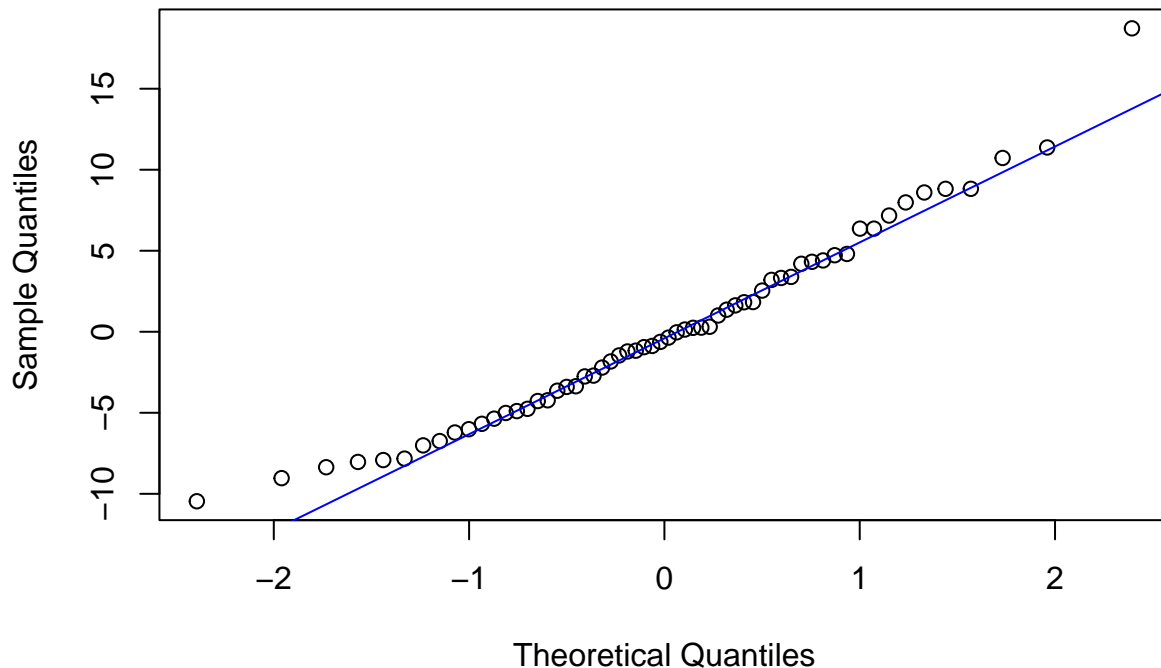
plot(model2$fitted.values, model2$residuals, xlab = "Fitted values", ylab = "Residuals",
      main = "Residuals vs Fitted (Quadratic Model)",
      abline(h = 0, col = "red"))
```

### Residuals vs Fitted (Quadratic Model)



```
qqnorm(model2$residuals)
qqline(model2$residuals, col = "blue")
```

## Normal Q-Q Plot



```
new_data <- data.frame(age = 48)

predict(model1, newdata = new_data, interval = "confidence", level = 0.99)
##      fit      lwr      upr
## 1 83.28382 78.1548 88.41283

predict(model2, newdata = new_data, interval = "confidence", level = 0.99)
##      fit      lwr      upr
## 1 80.71912 71.03998 90.39827
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```

Null Hypothesis:  $B_1 = B_2 = 0$  (the terms are not significant, so the relationship is not significant) Alternate Hypothesis  $B_1 \neq 0$  or  $B_2 \neq 0$  ((at least one of) the terms are significant, so the relationship is significant) F-statistic p-value:  $2.2e-16$ , because  $p < \alpha$  at the 0.05 level, there is a significant regression relationship using the quadratic model.

Null Hypothesis:  $B_2 = 0$  (the quadratic term is not significant) Alternate Hypothesis  $B_2 \neq 0$  (the quadratic term is significant) F-statistic p-value: 0.4079, because  $p > \alpha$  at the 0.05 level, there is an insignificant difference between the two so the quadratic term is unnecessary.

Based on the residual vs. fitted values, I see a general random shape in the plot, satisfying the condition of linearity. The variances seem to be quite evenly spread out too, satisfying the condition of homoskedasticity. Finally, the plot of the residuals follows the line on the Q-Q plot really well, so the residuals should be

normally distributed.

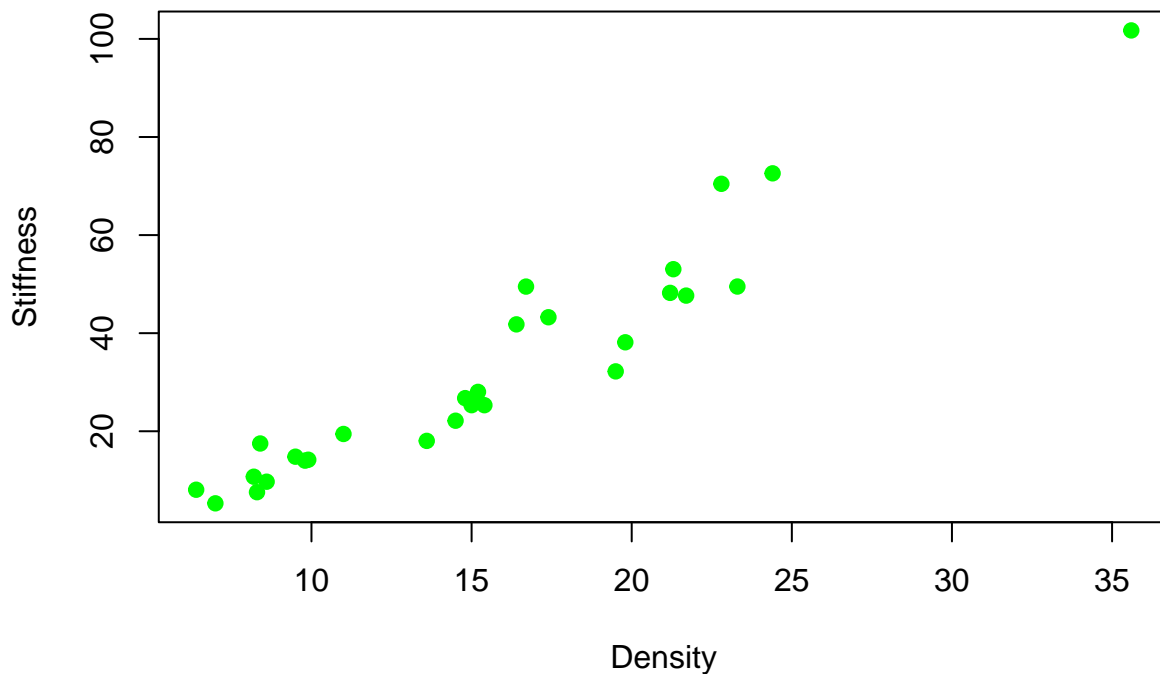
Problem 4:

```
data <- read.csv("ParticleBoard.csv")
head(data)
```

```
##   Density Stiffness   Temp
## 1    9.5    14.814 70.61056
## 2    8.4    17.502 72.30000
## 3    9.8    14.007 66.15377
## 4   11.0    19.443 70.05781
## 5    8.3    7.573 69.33919
## 6    9.9    14.191 69.12882
```

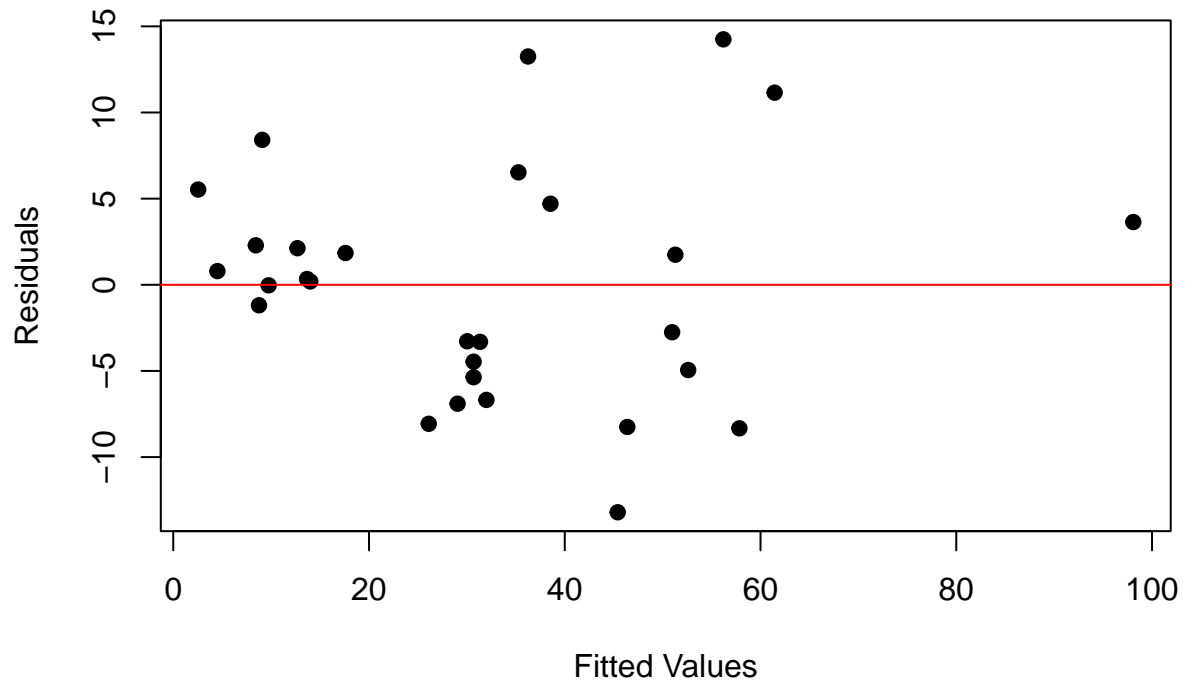
```
plot(data$Density, data$Stiffness, main = "Scatterplot of Stiffness vs Density",
      xlab = "Density", ylab = "Stiffness", pch = 19, col = "green")
```

**Scatterplot of Stiffness vs Density**



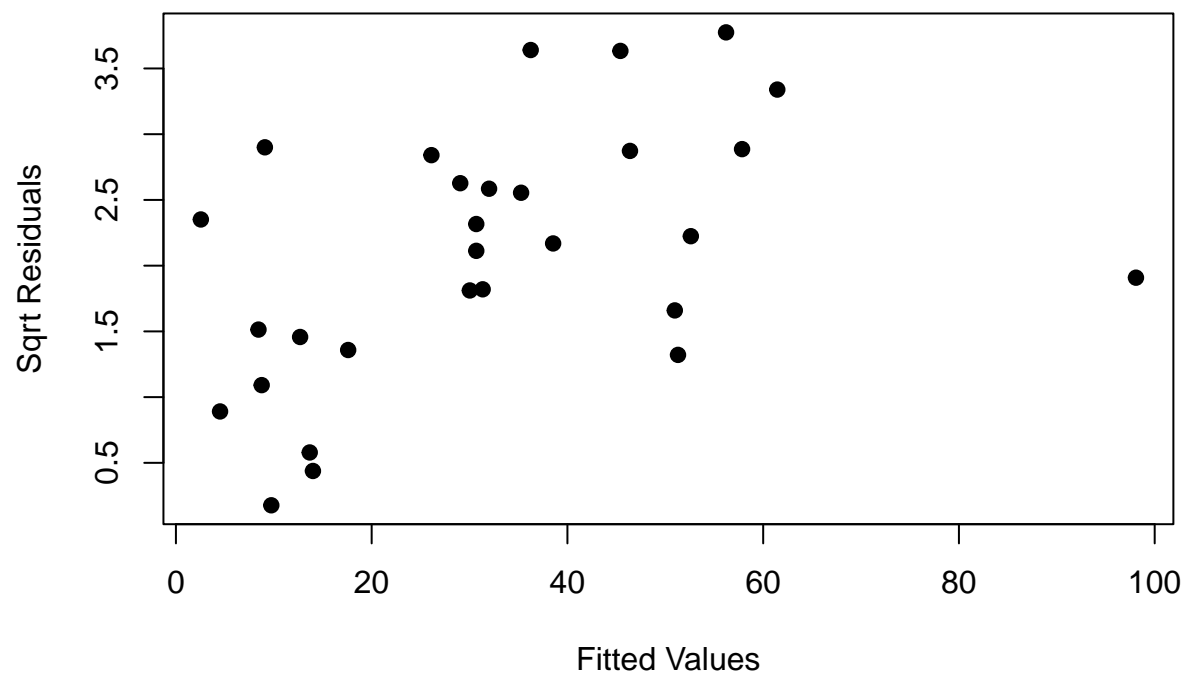
```
model <- lm(Stiffness ~ Density, data = data)
plot(model$fitted.values, model$residuals, main = "Residuals vs Fitted",
      xlab = "Fitted Values", ylab = "Residuals", pch = 19)
abline(h = 0, col = "red")
```

### Residuals vs Fitted



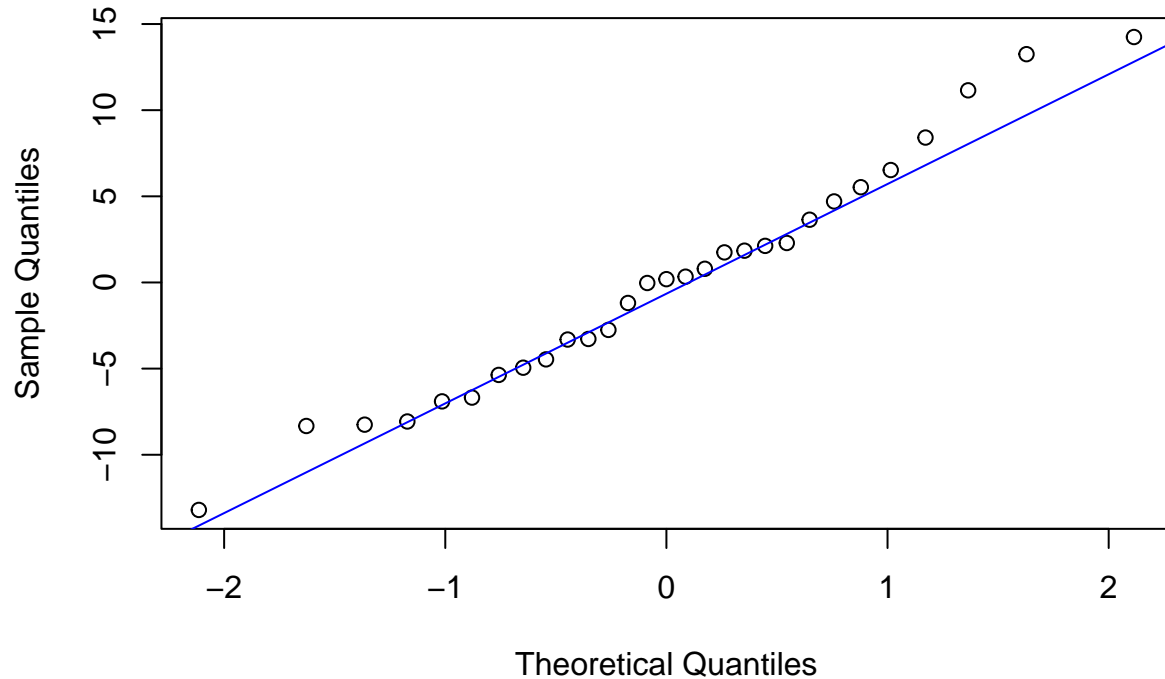
```
sqrt_abs_resid <- sqrt(abs(model$residuals))  
plot(model$fitted.values, sqrt_abs_resid, main = "Scale-Location Plot",  
      xlab = "Fitted Values", ylab = "Sqrt Residuals", pch = 19)
```

### Scale-Location Plot



```
qqnorm(model$residuals)
qqline(model$residuals, col = "blue")
```

Normal Q-Q Plot



```
data$sqrt_Stiffness <- sqrt(data$Stiffness)

powers <- seq

# FINISHED? KNIT YOUR WORK BEFORE GOING TO THE NEXT QUESTION.

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```

They appear to have a positive linear relationship.

The residuals appear to have a pattern however. They start near zero then have a sort of downward dip and then go back up again. This means we might need a quadratic or non linear term.

In the scale-location plot there seems to be a sort of pattern, kind of curved maybe square root curved. This is interesting and likely means the data violates the principles of heteroskedasticity.

The dots generally stick to the line but its definitely not a perfect fit to the normal distribution.

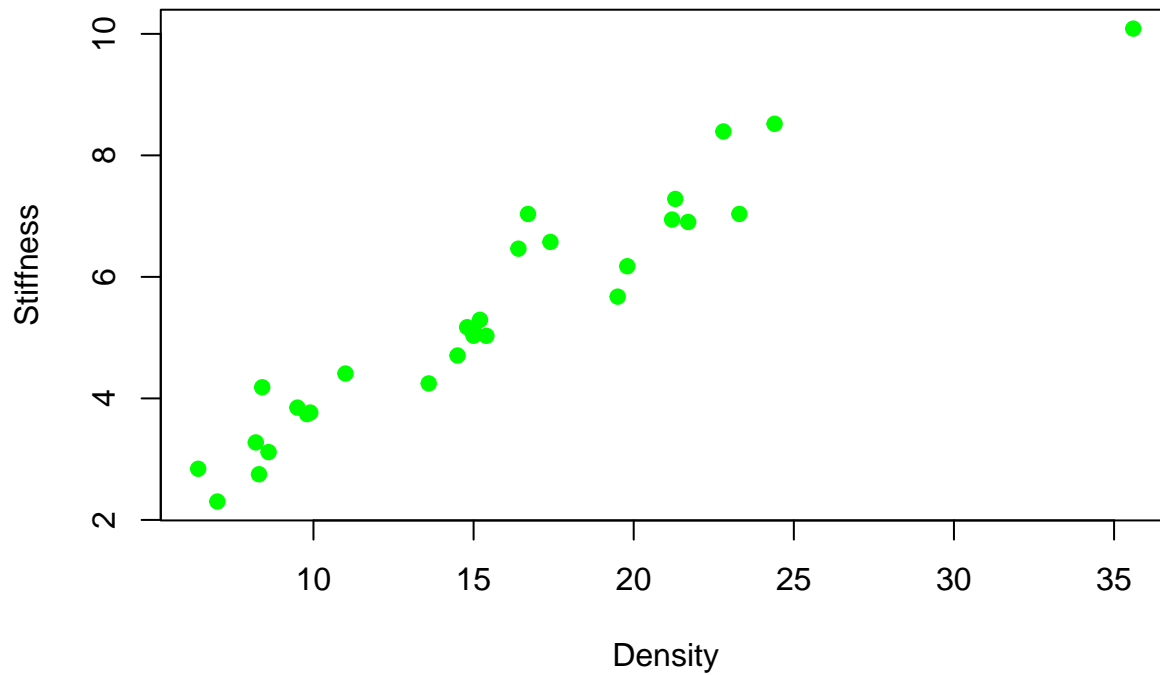
I think I would do Stiffness to the square root and then density based on the graphs. The residuals sort of fan out at higher values and Stiffness looks slightly right skewed.

I would transform Stiffness for sure because as we see in the residual plot there seems to be some sort of non-constant variance and transforming the response (Stiffness) and help solve this.



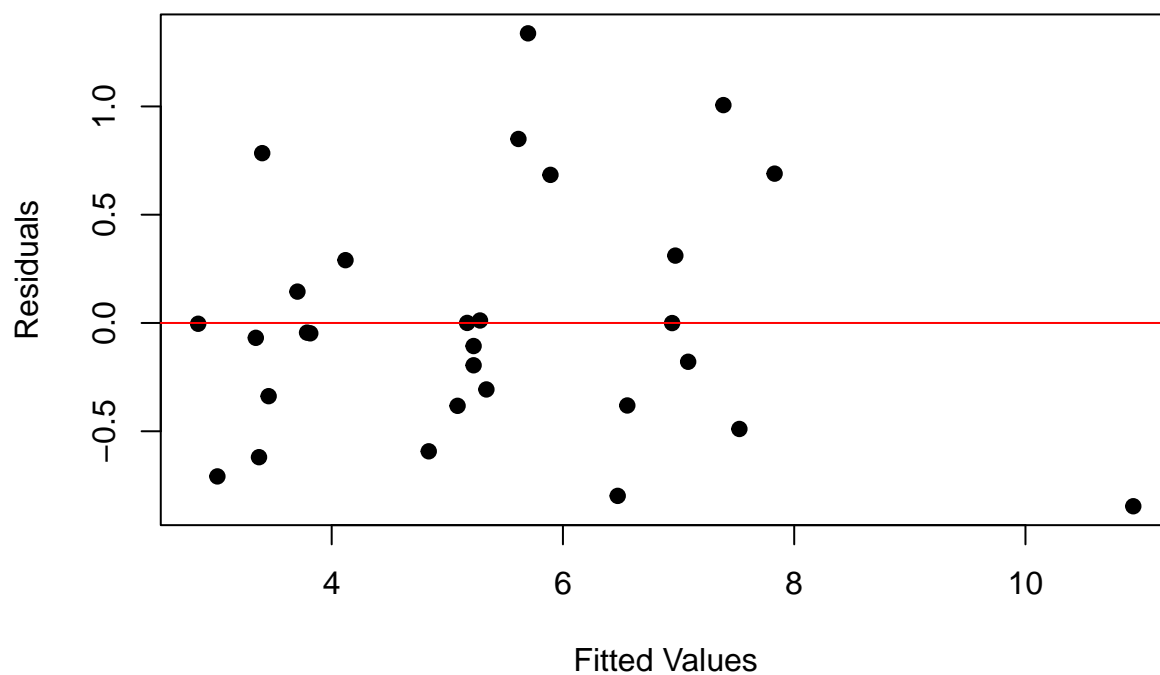
```
data$Stiffness = sqrt(data$Stiffness)
plot(data$Density, data$Stiffness, main = "Scatterplot of Stiffness vs Density",
      xlab = "Density", ylab = "Stiffness", pch = 19, col = "green")
```

**Scatterplot of Stiffness vs Density**



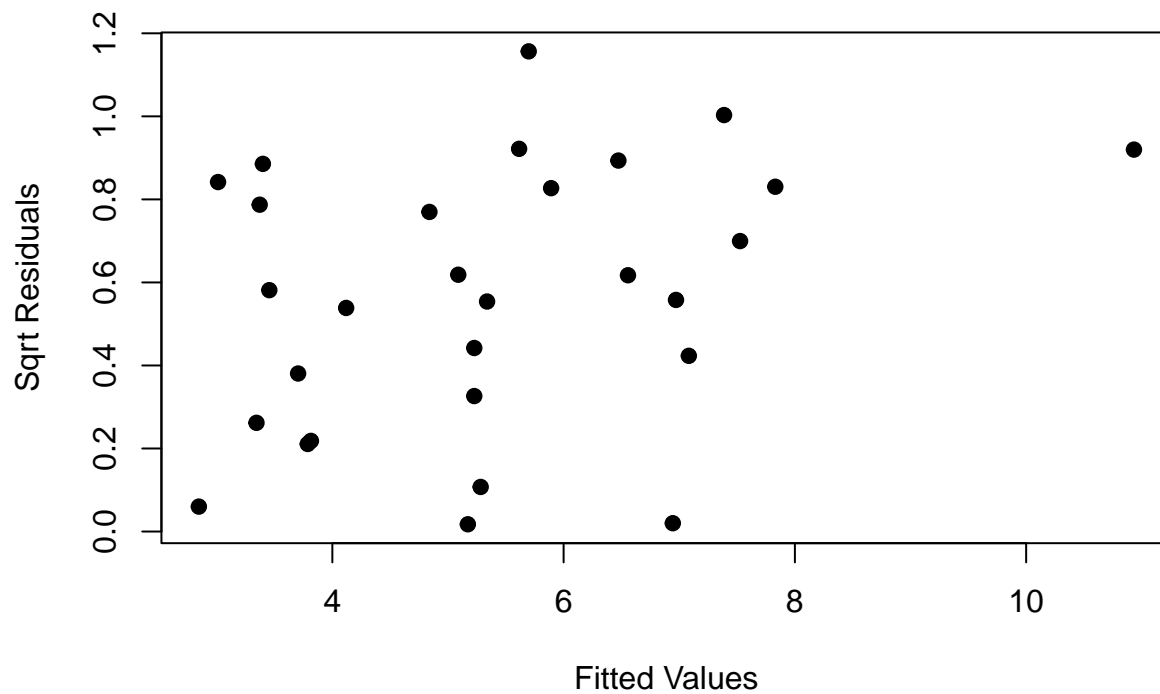
```
model <- lm(Stiffness ~ Density, data = data)
plot(model$fitted.values, model$residuals, main = "Residuals vs Fitted",
      xlab = "Fitted Values", ylab = "Residuals", pch = 19)
abline(h = 0, col = "red")
```

## Residuals vs Fitted



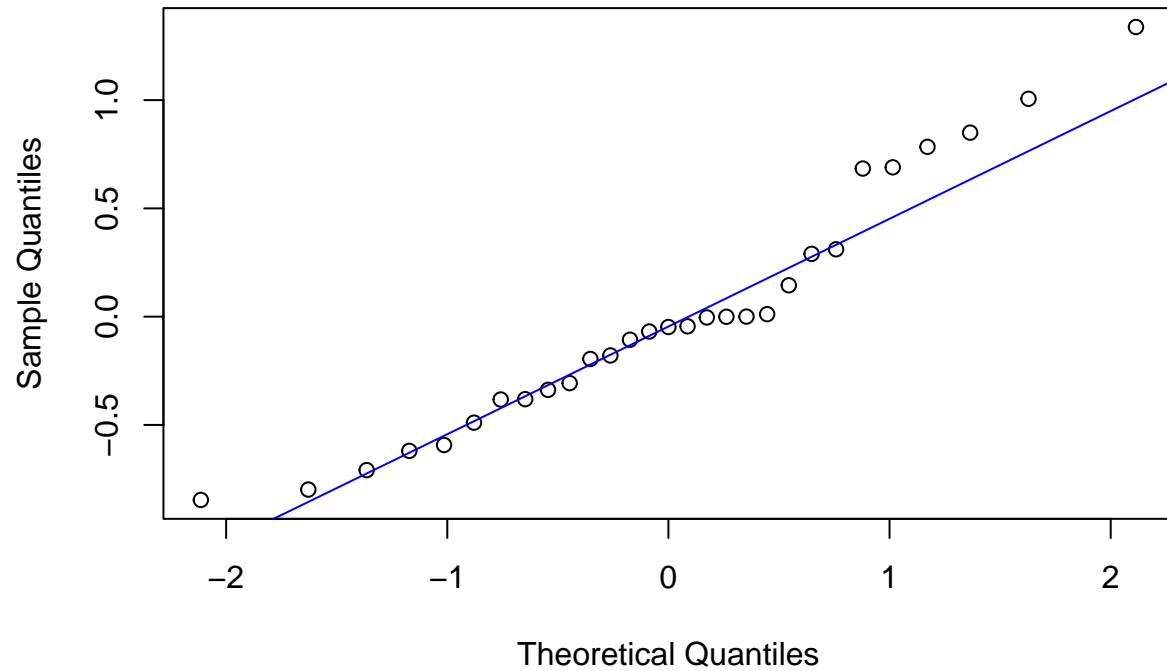
```
sqrt_abs_resid <- sqrt(abs(model$residuals))  
plot(model$fitted.values, sqrt_abs_resid, main = "Scale-Location Plot",  
      xlab = "Fitted Values", ylab = "Sqrt Residuals", pch = 19)
```

## Scale-Location Plot



```
qqnorm(model$residuals)
qqline(model$residuals, col = "blue")
```

**Normal Q-Q Plot**



```
data$sqrt_Stiffness <- sqrt(data$Stiffness)
powers <- seq
```