

Lab4_Transformations

Nahom Ayele & Nick Huo

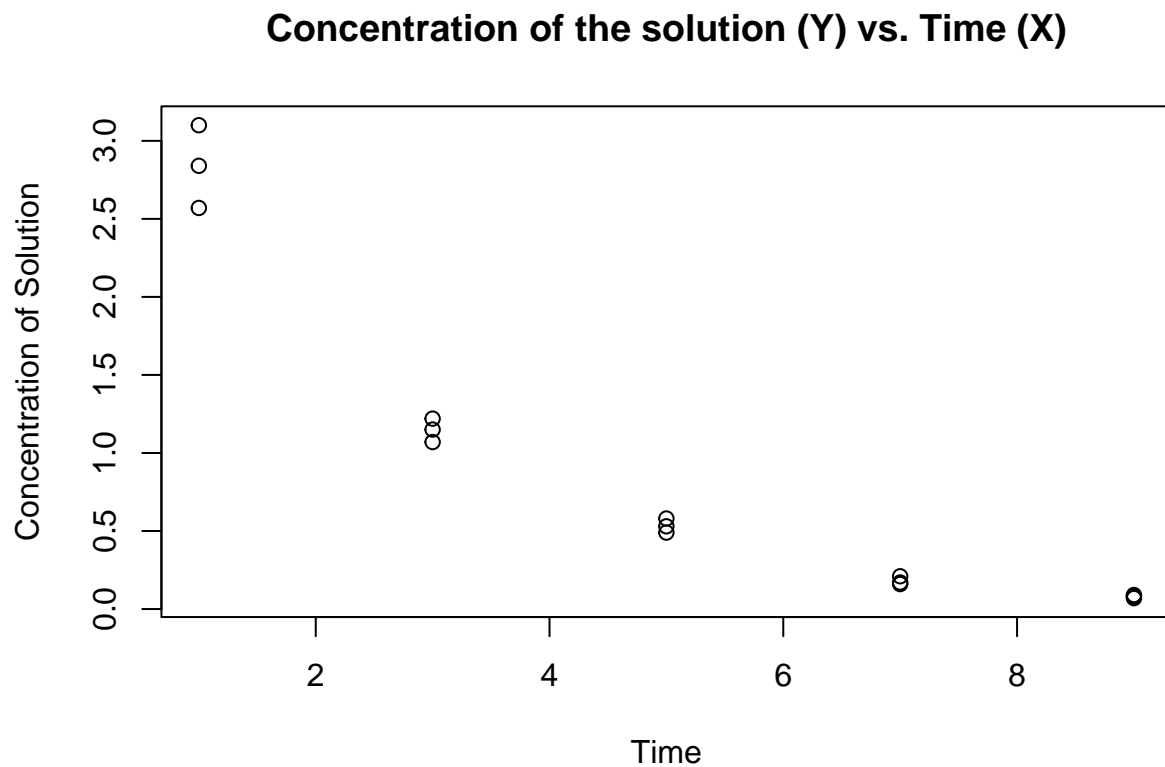
2022-09-30

```
# Import Data set  
sc <- read.csv("./SolutionConcentration.csv")  
x <- sc$x  
y <- sc$y
```

Question 1

Part (1)

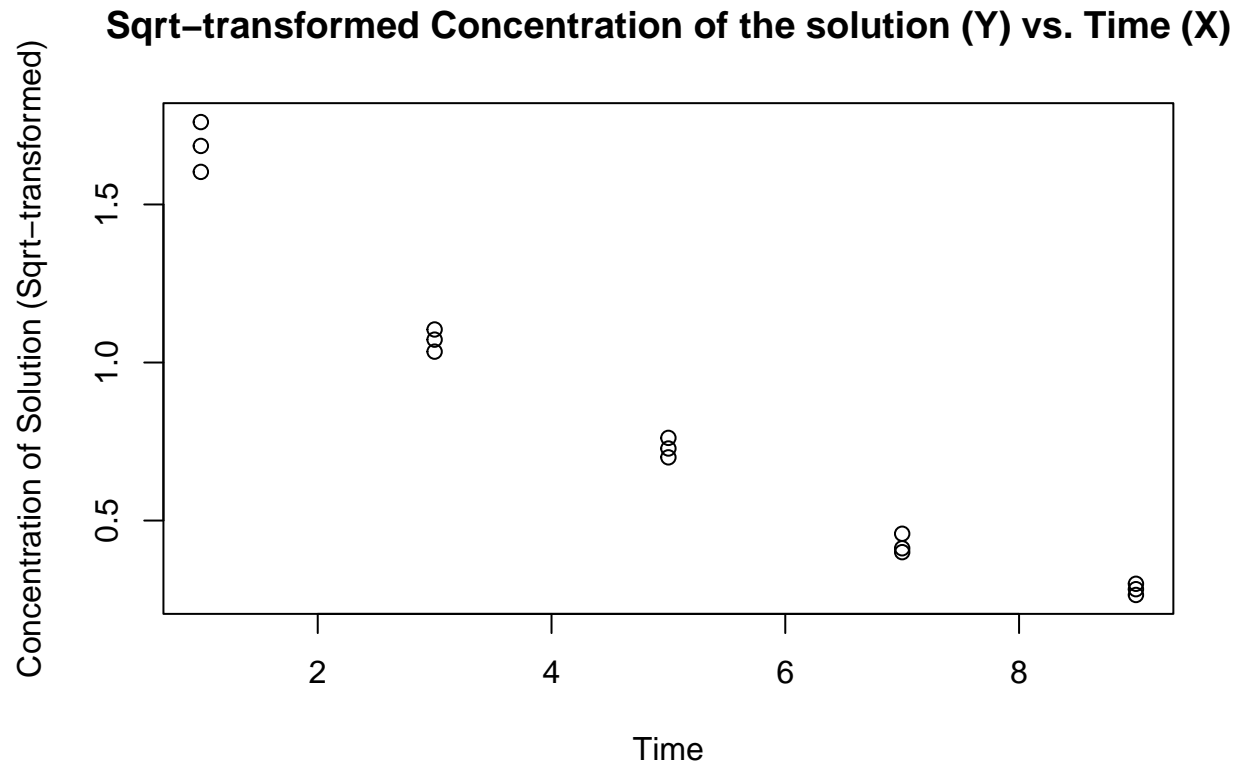
```
plot(x, y, main='Concentration of the solution (Y) vs. Time (X)',  
      xlab='Time', ylab='Concentration of Solution')
```



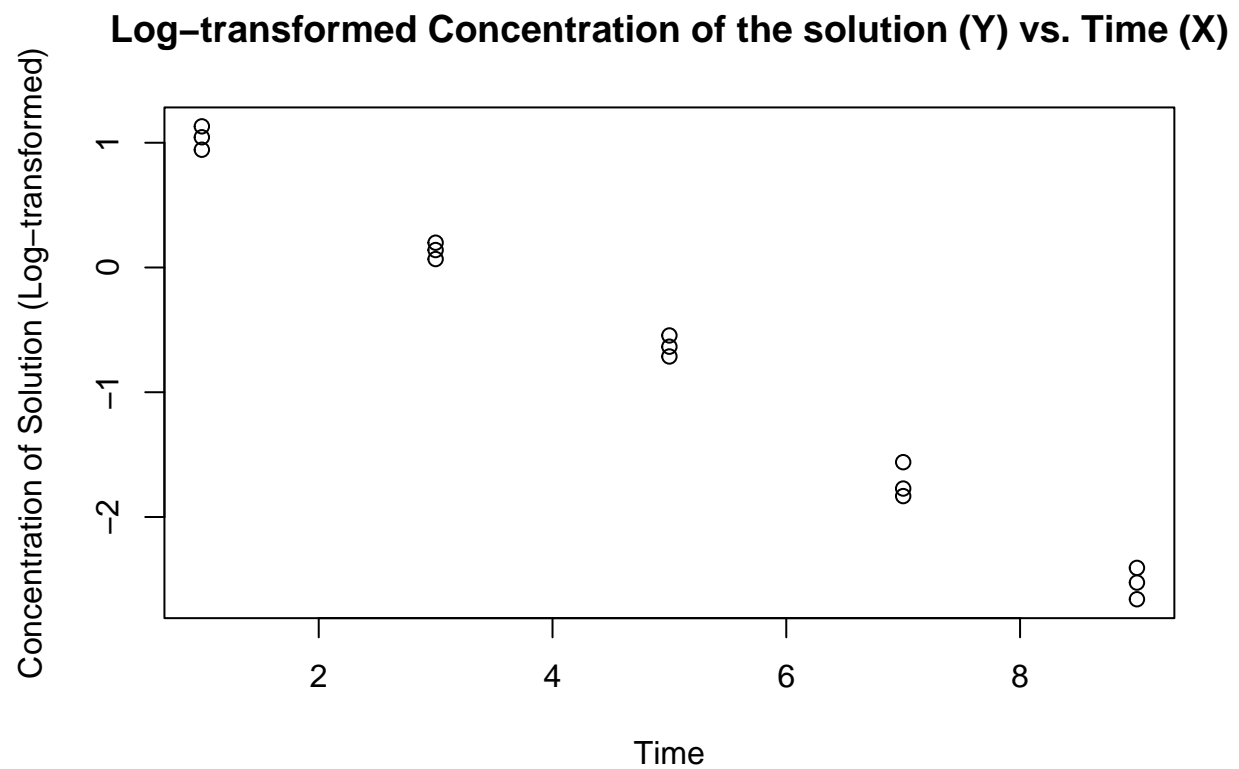
We observe that the relationship appears to be an exponential decay function. As time progresses, the concentration of the solution is exponentially decaying.

We think that the most appropriate transformation would be a squared root or a log transform, because we need to bring the high Y values at small X down and not change the low Y values as much.

```
## plotting Log transform
plot(x, sqrt(y), main='Sqrt-transformed Concentration of the solution (Y) vs. Time (X)',
      xlab='Time', ylab='Concentration of Solution (Sqrt-transformed)')
```



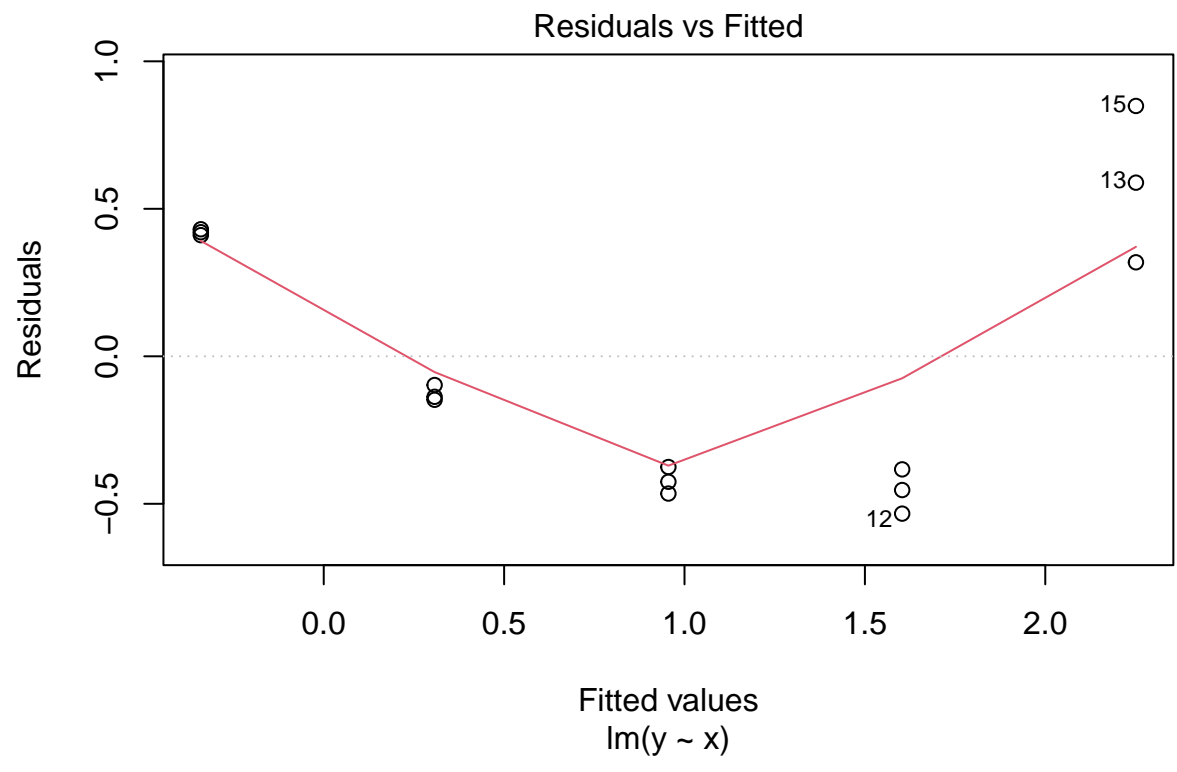
```
plot(x, log(y), main='Log-transformed Concentration of the solution (Y) vs. Time (X)',
      xlab='Time', ylab='Concentration of Solution (Log-transformed)')
```

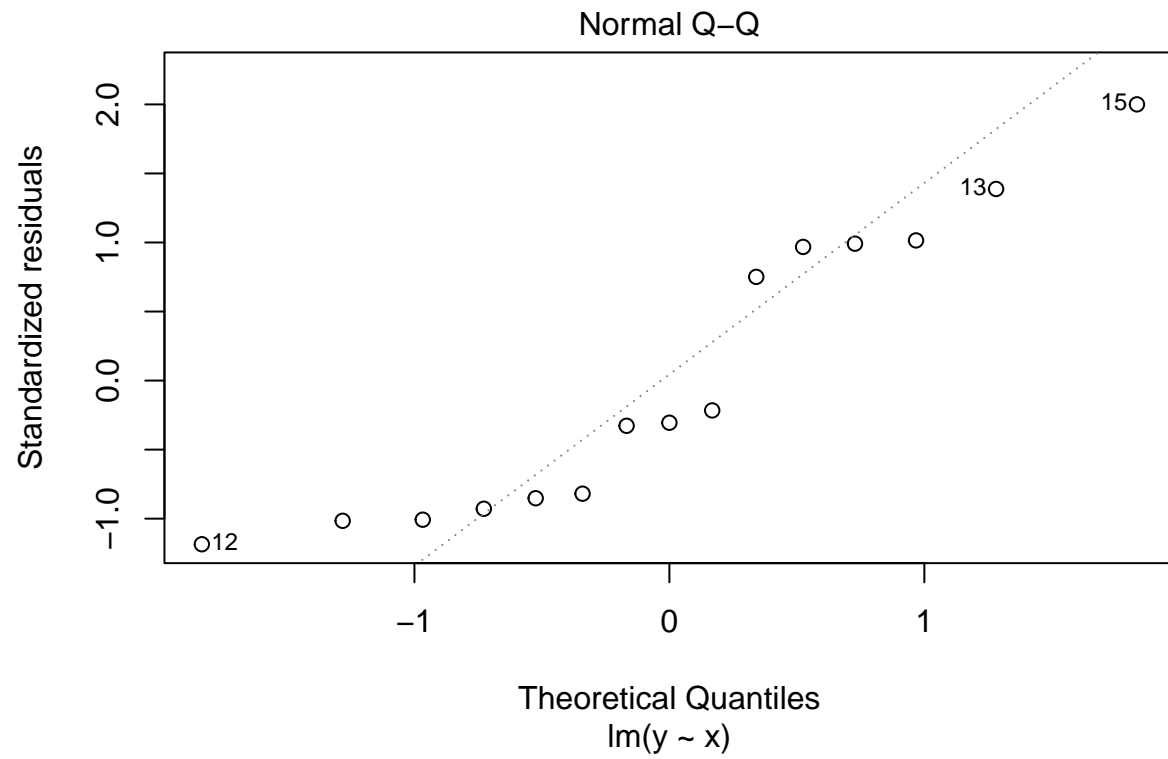


It appears that the log transformation is better as it makes the relationship between time (X) and log-concentration ($\ln[Y]$) quite linear.

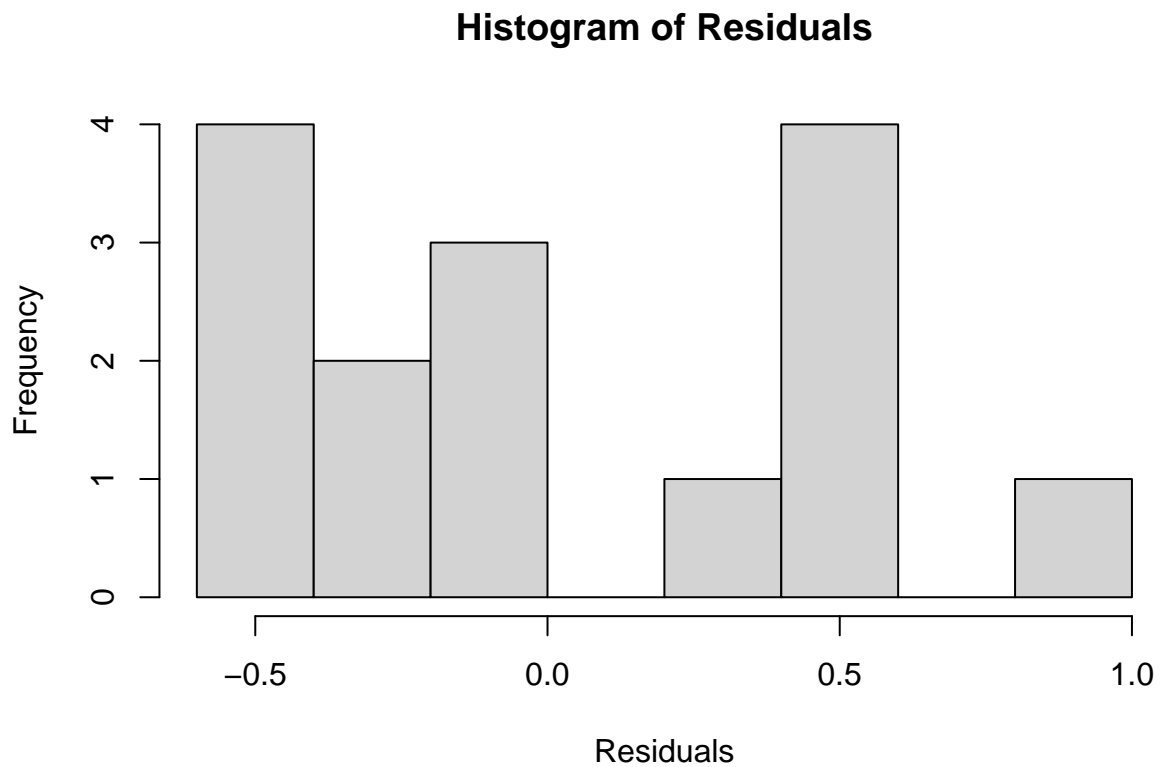
Part (2)

```
og_model <- lm(y ~ x)
plot(og_model, which=c(1,2))
```





```
hist(resid(og_model), main='Histogram of Residuals', xlab='Residuals')
```



CONCLUSIONS FROM CONDITIONS??

Part (3)

```
std_obs_wi <- function(y, lbd) {
  n <- length(y)
  k2 <- prod(y)^(1/n) # geometric mean of the Yi observations

  if (lbd==0) {
    return (k2*log(y))
  } else {
    k1 <- 1/(lbd*k2^(lbd-1))
    return (k1*((y^lbd)-1))
  }
}
```

Part (4)

```
sse <- function(x, w) {
  stdmodel <- lm(w~x)
  return (sum((fitted(stdmodel)-w)^2))
}
```

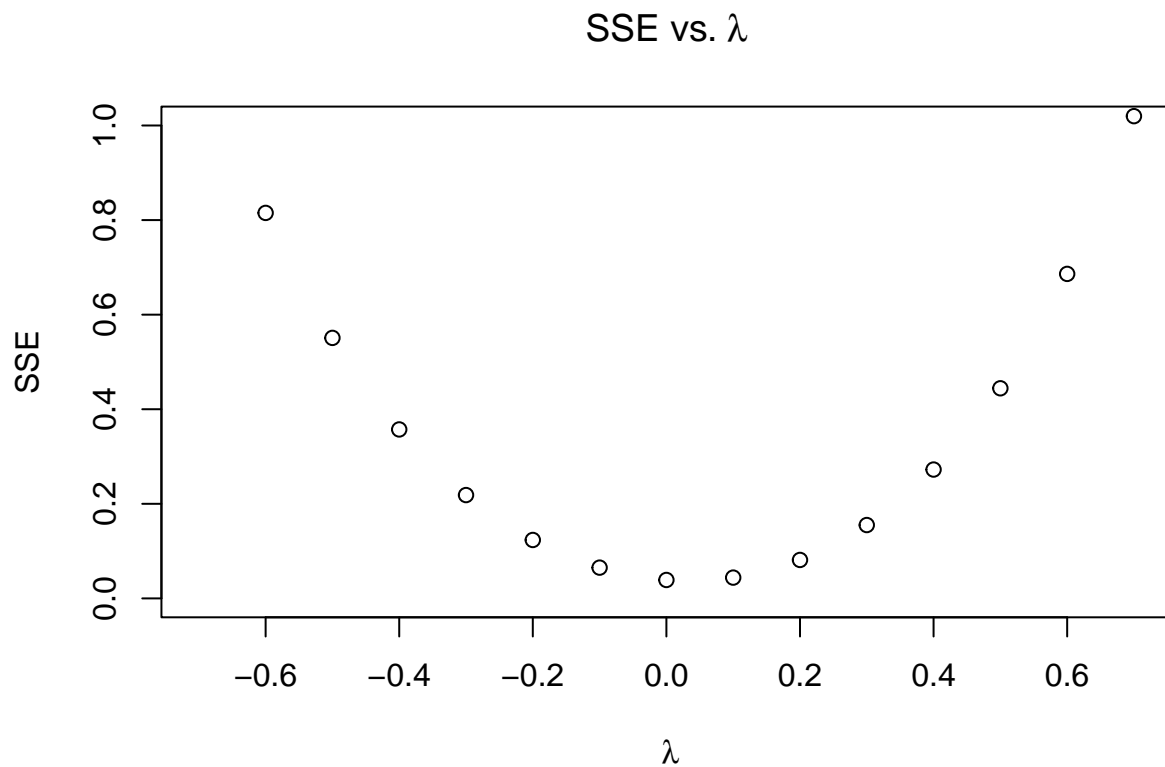
Part (5)

```
lambda_set <- seq(from=-3, to=3, by=.1)
sse_calc <- rep(NA, length(lambda_set))

for(i in 1:length(lambda_set)){
  Wi_calc <- std_obs_wi(y, lambda_set[i])
  sse_calc[i] <- sse(x, Wi_calc)
}
```

Part (6)

```
Q3_plot <- plot(lambda_set, sse_calc, xlab = expression(~lambda),
  ylab= "SSE", main= expression(paste("SSE vs.",~lambda)),
  xlim=c(-.7,.7), ylim=c(0,1))
```



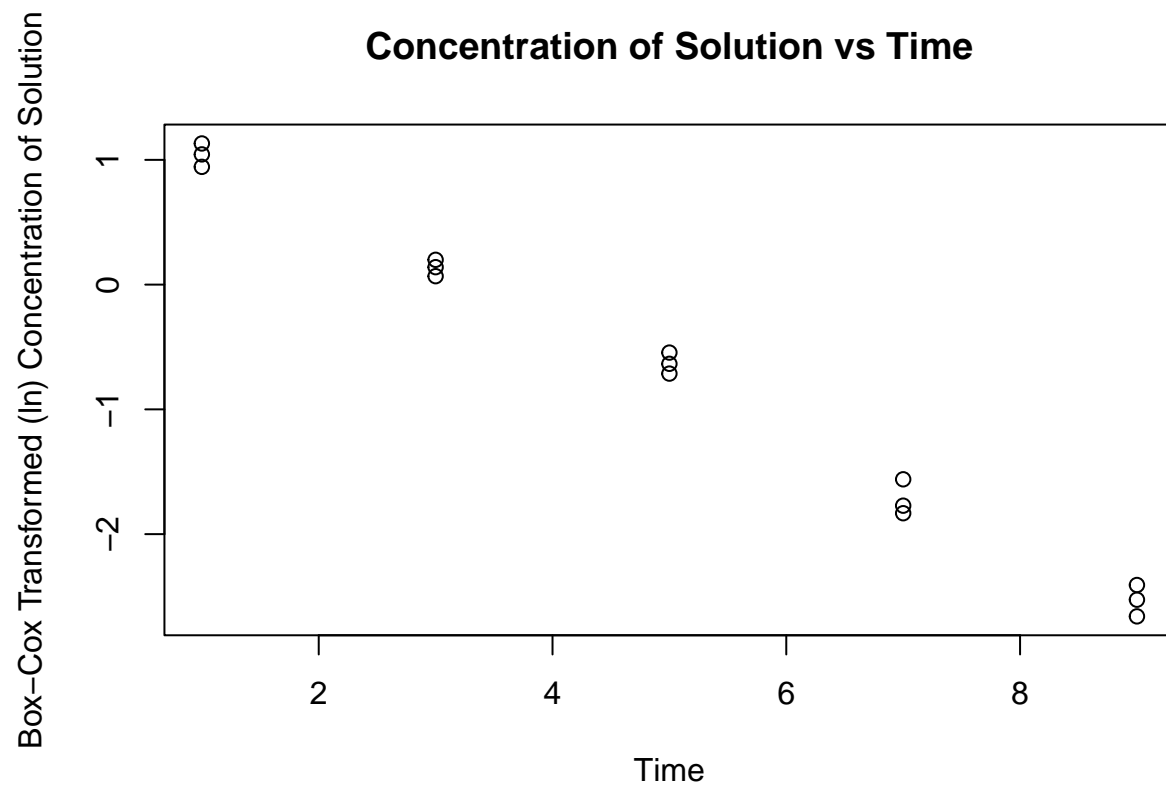
```
## Calculating the min lambda
lambda_set[which.min(sse_calc)]
```

```
## [1] 0
```

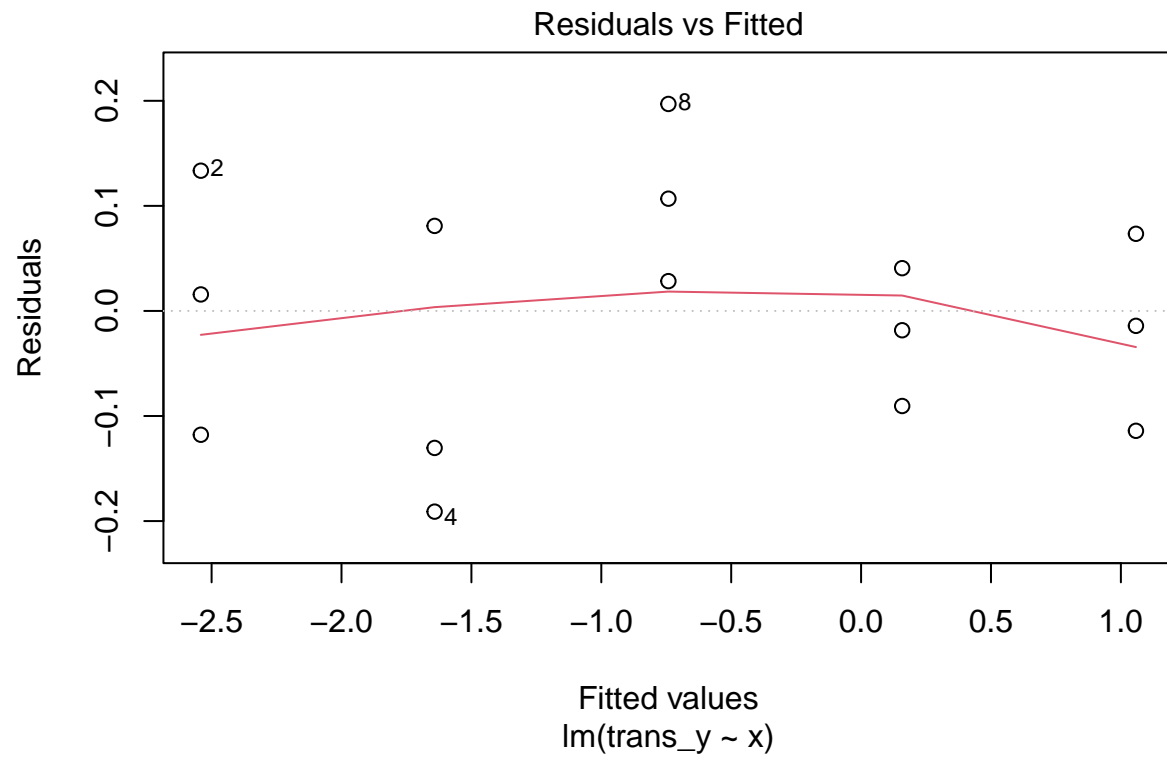
The value of λ that minimizes our *SSE* is 0.

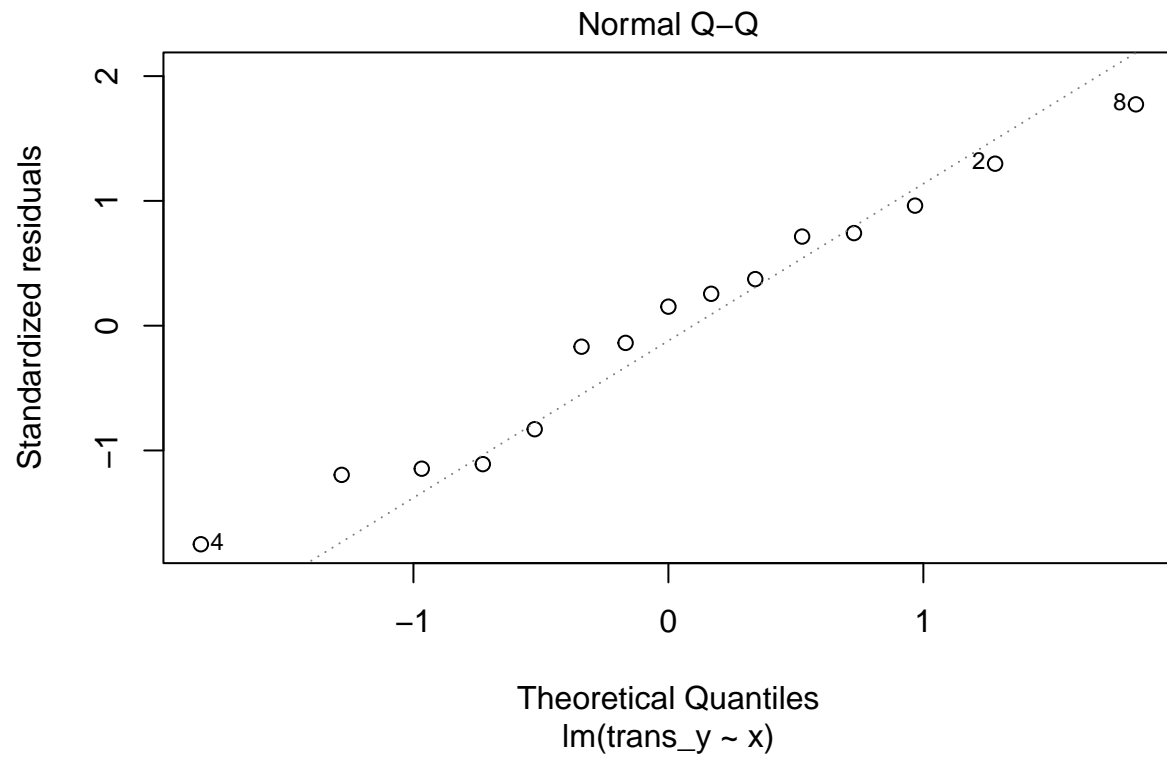
Part (7)

```
## plotting the Suggested Box-Cox transformation
trans_y <- log(y)
plot(x, trans_y, main = "Concentration of Solution vs Time",
     xlab="Time", ylab="Box-Cox Transformed (ln) Concentration of Solution")
```



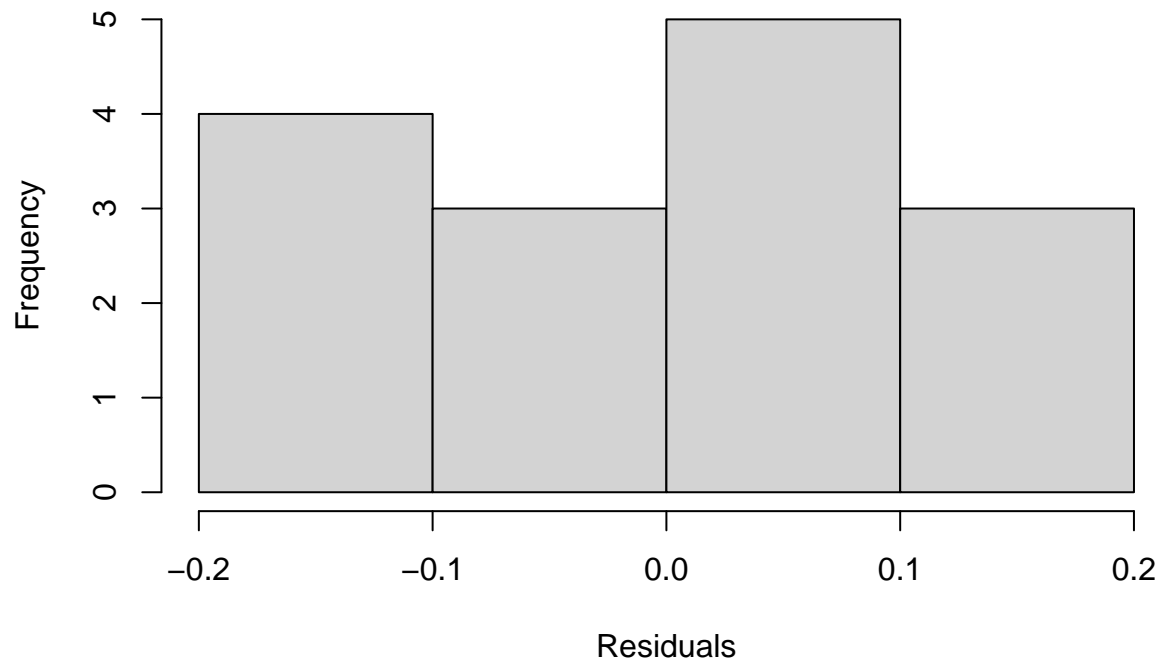
```
bc_model <- lm(trans_y ~ x)
plot(bc_model, which=c(1,2))
```



```
hist(resid(bc_model), main='Transformed Model: Histogram of Residuals', xlab='Residuals')
```

Transformed Model: Histogram of Residuals



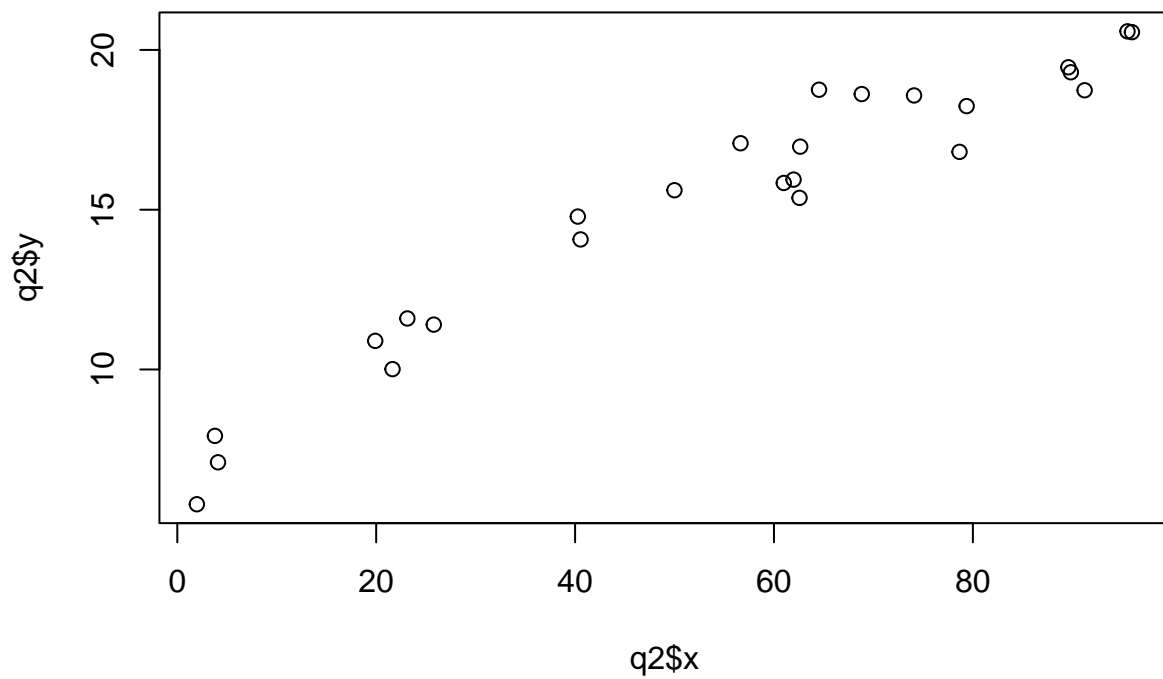
CONDITIONS

Section 2 - Transforming X - The Box-Tidwell Transformation

```
q2 <- read.csv("./Lab4q2.csv")
```

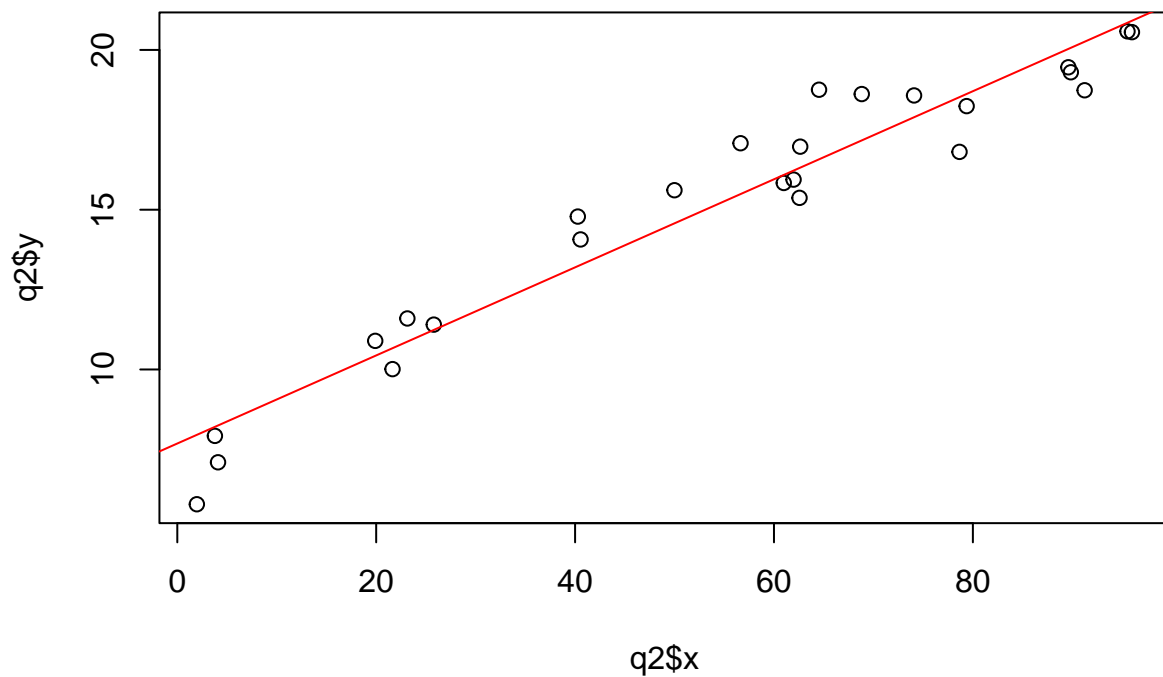
Part (1)

```
#plotting the data  
plot(q2$x, q2$y)
```



Part (2)

```
plot(q2$x, q2$y)
abline(lin_model<-lm(q2$y~q2$x), col = "red")
```



Part (3)

```
alpha_bt <- function(xi, yi, aj) {
  x.a <- xi^aj
  b <- x.a*log(xi)
  model1 <- lm(yi~x.a)
  model2 <- lm(yi~x.a+b)
  gamma <- coef(model2)[3]
  new_a <- gamma/coef(model1)[2] + aj
  return(new_a)
}
```

Part (4)

```
a_vec <- NULL
a_vec[1] <- 1
for (i in 1:9) {
  next_a <- alpha_bt(q2$x, q2$y, a_vec[i])
  a_vec[i+1] <- next_a
}

#model_bt <- lm()
```

fit the regression model and conduct a residual analysis