Statistical Learning Laboratory 1

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The dataset that we have used to solve the given task is part 12.

Part 2:

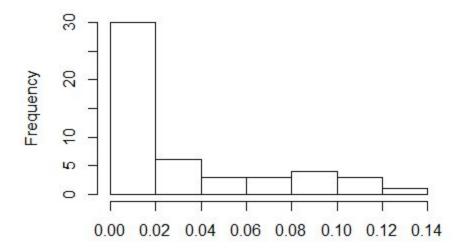
 Shapiro-wilk Normality test data: df\$density
 W = 0.75457, p-value = 9.117e-08
 Comment-- it is not a normal dist as p<0.005

data: df\$concentration

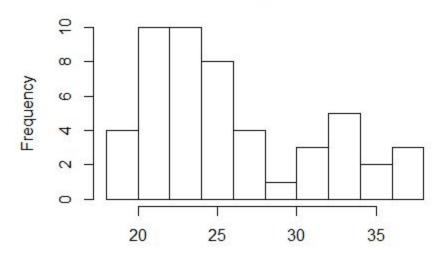
W = 0.89393, p-value = 0.0003046

Comment--it is not a normal dist as p<0.005

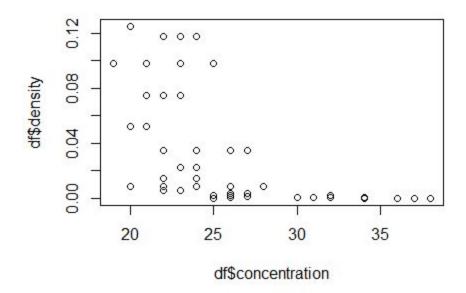
Histogram of .



Histogram of .



2. Scatterplot



3. because the output of p value is 7.737e-06 < 0.05 we can say that correlation exist correlation value is -0.5861335.

because the output of p value is 1.45e-12<0.05 we can say that correlation exist correlation value is 0.8071393.

4. comment on scatterplot and correlation as it can be seen from the scatterplot and the correlation values the variables are negatively correlated to each other and can be written as a negative linear relationship between the two variables.

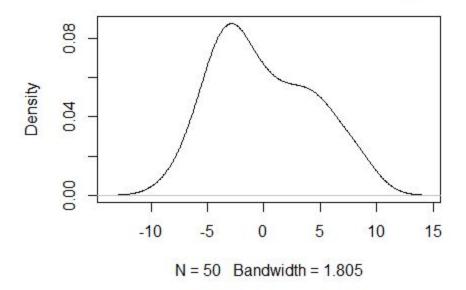
5. Coefficient of linear data

```
Coefficients:
(Intercept) df$density
28.87 -78.47
coeff =-78.47
```

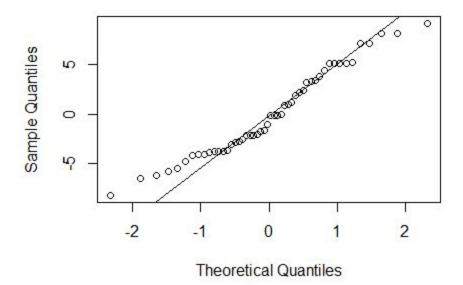
Residual Plot

```
> resid(model) #List of residuals
                                                               6
-2.1238092 -1.0226672 5.1614262 -0.1603812 -4.1018370 -1.7775256 -2.7152218 5.1473804
       10
                  11
                             12
                                        13
                                                   14
                                                              15
                                                                        16
                                                                                   17
                                                                                              18
-3.7549314 -6.1813241 -3.1018370 -0.0226672 -4.7549314 4.3458945 -6.4479591 -4.1238092 7.1337269
       19
                 20
                             21
                                                              24
                                                   23
 8.1337269 -2.1813241 3.2847782 1.1473804
                                           7.1337269 -5.4479591 -0.1238092 -3.8713735 -8.1813241
       28
                  29
                             30
                                        31
                                                   32
                                                              33
                                                                        34
                                                                                   35
 2.1844958 0.9342491 -3.7549314 -1.6134485 -2.8155042 3.3458945 -0.1813241
                                                                            3.1614262 -4.1813241
                             39
                                        40
                                                   41
                                                             42
                                                                        43
                                                                                   44
       37
                  38
                                                                                              45
 5.1473804 3.8396188 5.1258016 0.8761908 5.1275279 -5.7579097 -3.7579097 -2.0226672 9.1258016
       46
                 47
                            48
                                        49
 8.1268217 2.3458945 -3.7152218 -2.6134485 -2.1603812
```

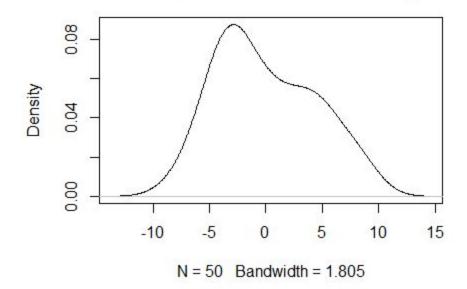
density.default(x = resid(model))



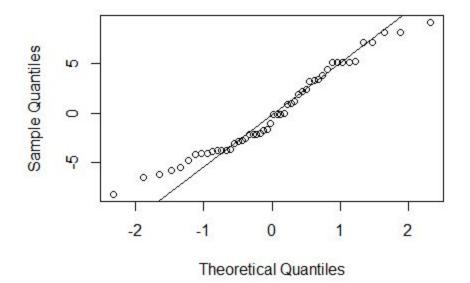
Normal Q-Q Plot



density.default(x = resid(model))



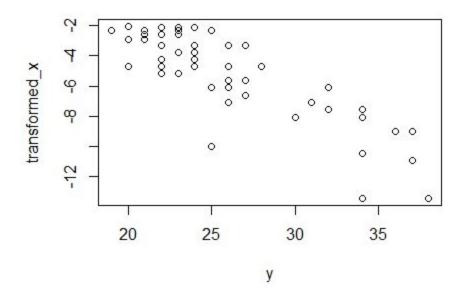
Normal Q-Q Plot



```
> y<-df$concentration
> x<-df$density
> x

[1] 0.035052 0.074573 0.000455 0.098306 0.022588 0.001233 0.002027 0.000276 0.098306 0.052497 0.008831
[12] 0.022588 0.074573 0.052497 0.117502 0.005433 0.035052 0.000102 0.000102 0.008831 0.002027 0.000276
[23] 0.000102 0.005433 0.035052 0.000037 0.008831 0.000749 0.125000 0.052497 0.003224 0.000749 0.117502
[34] 0.008831 0.000455 0.008831 0.000276 0.098306 0.00001 0.035052 0.000023 0.014227 0.014227 0.074573
[45] 0.000001 0.000014 0.117502 0.002027 0.003324 0.098306
> y

[1] 24 22 34 21 23 27 26 34 23 21 22 24 23 20 24 22 22 36 37 26 32 30 36 23 26 25 20 31 20 21 27 26 23
[34] 28 32 24 34 25 34 27 34 22 24 21 38 37 22 25 26 19
```



7. Pearson correlation coefficient

Because the output of p value is 6.695e-15 < 0.05 we can say that correlation exist correlation value is -0.849047.

```
> pearson_corr
```

Pearson's product-moment correlation

```
data: y and transformed_x
t = -11.134, df = 48, p-value = 6.695e-15
alternative hypothesis: true correlation is not equal to 0
95 percent confidence interval:
   -0.9118880 -0.7473117
sample estimates:
        cor
   -0.849047
```

Spearman rank correlation coefficient

Because the output of p value is 1.45e-12 < 0.05 we can say that correlation exist correlation value is -0.8071393.

```
> spearman_corr
         Spearman's rank correlation rho
 data: y and transformed_x
 S = 37634, p-value = 1.45e-12
 alternative hypothesis: true rho is not equal to 0
 sample estimates:
        rho
 -0.8071393
     8.
a.
> cor.diff.test(x,transformed_x,y,y,method="pearson") #no sign difference
cor1: r=0.726, p=2.49e-09, n=50
cor2: r=1, p=0, n=50
diffence: p(one-sided)=0, p(two-sided)=0
$cor1
$cor1$estimate
[1] 0.7256493
$cor1$p.value
[1] 2.494812e-09
$cor1$n
[1] 50
$cor2
$cor2$estimate
[1] 1
$cor2$p.value
[1] 0
$cor2$n
[1] 50
$p.value.twosided
[1] 0
$p.value.onesided
[1] 0
```

B.

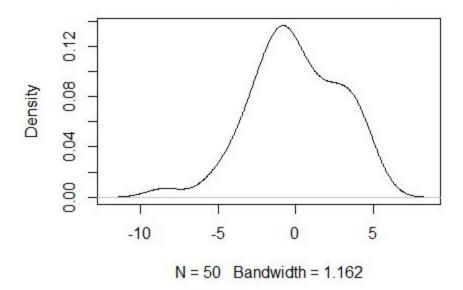
```
> cor.diff.test(x,transformed_x,y,y,method = 'spearman')
 cor1: r=1, p=0, n=50
 cor2: r=1, p=0, n=50
 diffence: p(one-sided)=0, p(two-sided)=0
 $cor1$estimate
 [1] 1
 $cor1$p.value
 [1] 0
 $cor1$n
 [1] 50
 $cor2
 $cor2$estimate
 [1] 1
 $cor2$p.value
 [1] 0
 $cor2$n
 [1] 50
 $p. value. twosided
 [1] 0
 $p.value.onesided
 [1] 0
9.
Linear model after transformation
call:
 lm(formula = y \sim transformed_x)
 Coefficients:
   (Intercept) transformed_x
        18.068
                        -1.542
```

```
call:
lm(formula = y \sim transformed_x)
Residuals:
           1Q Median
   Min
                           3Q
-8.4661 -1.5631 -0.3267 2.2199 5.0346
Coefficients:
            Estimate Std. Error t value Pr(>|t|)
             18.0676 0.8574 21.07 < 2e-16 ***
(Intercept)
                        0.1385 -11.13 6.7e-15 ***
transformed_x -1.5417
Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
Residual standard error: 2.89 on 48 degrees of freedom
Multiple R-squared: 0.7209, Adjusted R-squared: 0.7151
F-statistic: 124 on 1 and 48 DF, p-value: 6.695e-15
call:
lm(formula = df$concentration ~ df$density)
Residuals:
   Min
          10 Median
                      3Q
-8.181 -3.745 -0.602 3.331 9.126
Coefficients:
           Estimate Std. Error t value Pr(>|t|)
(Intercept) 28.8743 0.7905 36.526 < 2e-16 ***
                      15.6558 -5.012 7.74e-06 ***
df$density -78.4682
Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
Residual standard error: 4.432 on 48 degrees of freedom
Multiple R-squared: 0.3436, Adjusted R-squared: 0.3299
F-statistic: 25.12 on 1 and 48 DF, p-value: 7.737e-06
```

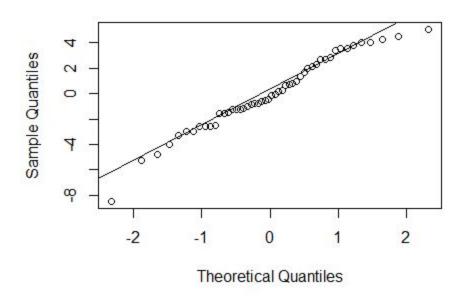
Comment: both p value and f value suggest the model have become better.

11.

density.default(x = resid(model_new))



Normal Q-Q Plot



A quantile normal plot - good for checking normality

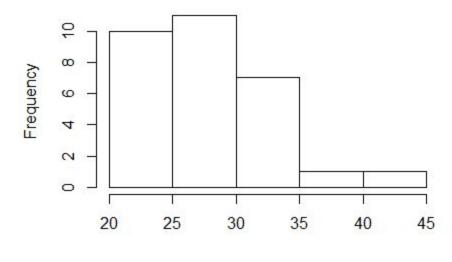
12. coefficient of determination ---- 0.7208808 coefficient of determination of old model --- 0.3435525

13. model performance has become better after the transfromation as can be seen form Q12.

Part 1:

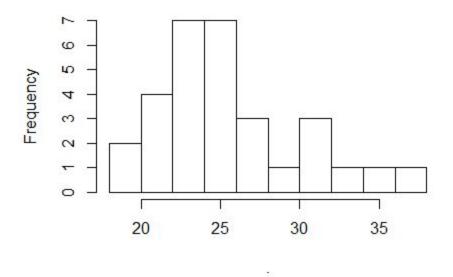
1. Shapiro-Wilk Test

Histogram of .



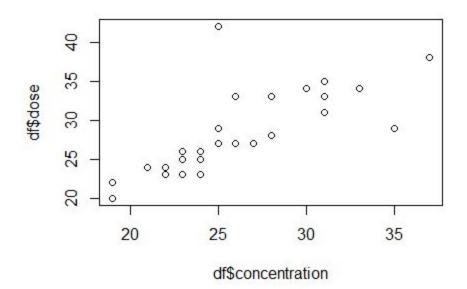
DOSE

Histogram of .



CONCENTRATION

2. Scatterplot



```
3.
```

> pearson_corr

```
Pearson's product-moment correlation
```

because the

output of p value is 4.53e-0.6<0.05 we can say that correlation exist. Cor value -- 0.7307623

> spearman_corr

Spearman's rank correlation rho

```
data: df$dose and df$concentration
S = 553.11, p-value = 2.074e-10
alternative hypothesis: true rho is not equal to 0
sample estimates:
    rho
0.876949
```

output of p value is 2.074e-10<0.05 we can say that correlation exist. Cor value -- 0.876949

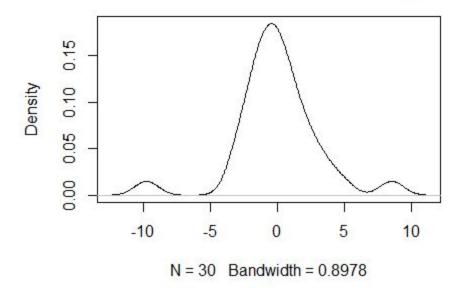
4.

as it can be seen from the scatterplot and the correlation values the variables are negatively correlated to each other and can be written as a negative linear relationship between the two variables.

5.

```
7.
 > summary(model_new) r.squared
 [1] 0.7208808
 > mmary(model) r. squared
 Error in mmary(model) : could not
 > summary(model)$r.squared
 [1] 0.5340135
8.
 > resid(model) #List of residuals
  3.26143390 -3.00429730 4.80144705 -0.19000219 -1.55286121 -0.64143828 -0.19000219 0.80999781
                     10
                                           12
                                                       13
                                                                  14
                                                                             15
                                11
 -9.74711686 3.35001096 0.35001096 1.80999781
                                               0.08427976
                                                          8.53571585 -0.55286121 -2.27857926
         17
                     18
                                19
                                            20
                                                       21
                                                                  22
                                                                             23
 -2.27857926
             1.35856172 -1.46428415 -1.01284806
                                               0.71286998 -0.91572024
                                                                      0.35856172 2.17285683
                                                       29
                                                                  30
                     26
                                27
                                           28
  0.80999781 1.98715194 -0.55286121 -1.73001535 -3.01284806 -1.27857926
```

density.default(x = resid(model))



9.

10. model performance has become better after the transfromation.