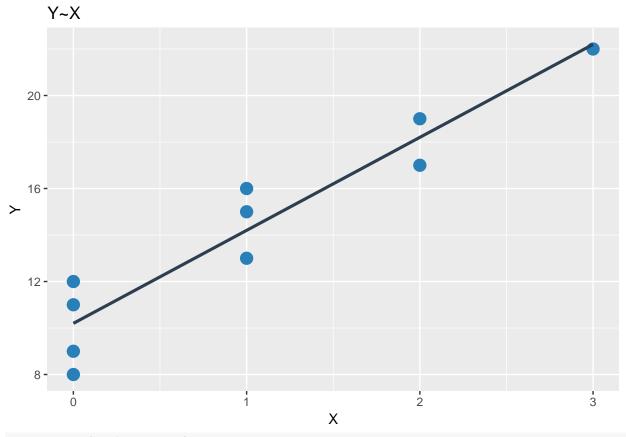
Homework Chapter 3

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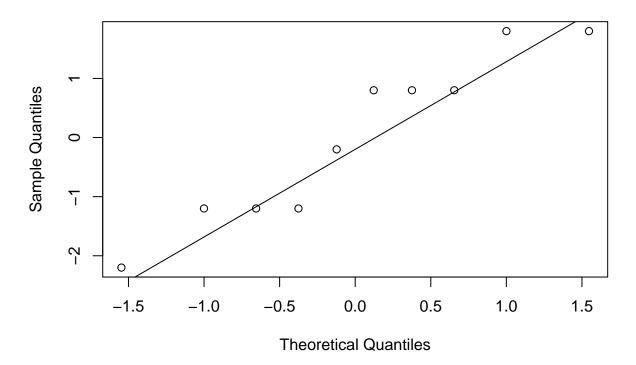
- 3.5 Refer to Airfreight breakage Problem 1.21.
- (e) Prepare a normal probability plot of the residuals. Also obtain the coefficient of correlation between the ordered residuals and their expected values under normality to ascertain whether the normality assumption is reasonable here. Use Table B.6 and $\alpha = .01$. What do you conclude?

```
library(ggplot2)
library(gridExtra)
data1 <- read.table("CH01PR21.txt",head=FALSE,col.names = c('Y','X'))</pre>
fit <- lm('Y~X',data1)</pre>
summary(fit)
##
## Call:
## lm(formula = "Y~X", data = data1)
## Residuals:
##
     Min
              1Q Median
                            3Q
                                  Max
##
     -2.2
            -1.2
                    0.3
                           0.8
                                  1.8
##
## Coefficients:
               Estimate Std. Error t value Pr(>|t|)
##
## (Intercept) 10.2000
                            0.6633 15.377 3.18e-07 ***
                 4.0000
                            0.4690
                                     8.528 2.75e-05 ***
## X
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
## Residual standard error: 1.483 on 8 degrees of freedom
## Multiple R-squared: 0.9009, Adjusted R-squared: 0.8885
## F-statistic: 72.73 on 1 and 8 DF, p-value: 2.749e-05
lm.scatter <- ggplot(data1, aes(x=X, y=Y)) +</pre>
  geom_point(color='#2980B9', size = 4) + xlim(c(0, 3)) +
  geom_smooth(method = lm, se=FALSE, fullrange=TRUE, color='#2C3E50', size=1.1) +
  labs(title='Y~X')
grid.arrange(lm.scatter)
```



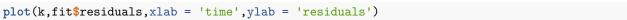
q <- qqnorm(fit\$residuals)
qqline(fit\$residuals)</pre>

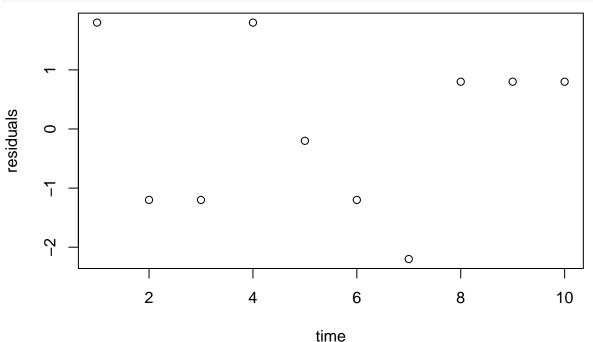
Normal Q-Q Plot



```
library(olsrr,warn.conflicts=FALSE)
resid=sort(fit$residuals)
n = length(data1$X)
k < -c(1:n)
z=qnorm((k-0.375)/(n+0.25))
MSE = sum(fit$residuals^2)/(fit$df.residual)
expect_residual = z* sqrt(MSE)
print(sprintf('The expected values of residuals is :'))
## [1] "The expected values of residuals is :"
expect_residual
   [1] -2.2940308 -1.4839673 -0.9721502 -0.5568998 -0.1818168 0.1818168
   [7] 0.5568998 0.9721502 1.4839673 2.2940308
ols_corr_test(fit)
## [1] 0.9609751
                               H_0: Normal
                                             H_a: not normal
r = 0.9609751 \ge 0.879, conclude H_0.
```

(f) Prepare a time plot of the residuals. What information is provided by your plot?





The residual versus time plot did not show any evidence that the error terms were correlated over time.

(g) Assume that (3.10) is applicable and conduct the Breusch-Pagan test to determine whether or not the error variance varies with the level of X. Use $\alpha = .10$. State the alternatives, decision rule, and conclusion. Does your conclusion support your preliminary findings in part (d)?

```
library(lmtest,warn.conflicts = FALSE )
## Loading required package: zoo
##
## Attaching package: 'zoo'
## The following objects are masked from 'package:base':
##
##
        as.Date, as.Date.numeric
library(latex2exp,warn.conflicts = FALSE )
data2 = data.frame(x=data1$X,y=fit$residuals^2)
fit2 = lm('y~x',data2)
SSR star = sum((fitted(fit2)-mean(data2$y))^2)
SSE = sum(fit$residuals^2)
XBP = (SSR_star/2) / (SSE/n)^2
print(sprintf('SSR* :%f',SSR_star))
## [1] "SSR* :6.400000"
print(sprintf('SSE :%f',SSE))
## [1] "SSE :17.600000"
print(sprintf('XBP^2:%f',XBP))
## [1] "XBP^2:1.033058"
bptest(data1$Y ~ data1$X,studentize=FALSE)
##
    Breusch-Pagan test
##
##
## data: data1$Y ~ data1$X
## BP = 1.0331, df = 1, p-value = 0.3094
    when H_0 holds,
                                            X_{BP}^2 \stackrel{\cdot}{\sim} \chi^2(1)
٠.
                                      H_0: \gamma_1 = 0 H_a: \gamma_1 \neq 0
where
                                         \log_e \sigma_i^2 = \gamma_0 + \gamma_1 X_i
If X_{BP}^2 < \chi^2(0.9;1) = 2.71, then conclude H_0; otherwise conclude H_a.
Here X_{RP}^2 = 1.0331 < 2.71, conclude H_0.
```

3.15 Solution concentration. A chermist studied the concentration of a solution (Y) over time (X). Fifteen identical solutions were prepared. The 15 solutions were randomly divided into five sets of three, and the five sets were measured, respectively, after 1, 3, 5, 7, and 9 hours. The results follow.

(a) Fit a linear regression function.

```
data3 <- read.table("CH03PR15.txt",head=FALSE,col.names = c('Y','X'))
fit3 <- lm('Y~X',data3)
summary(fit3)</pre>
```

```
##
## Call:
## lm(formula = "Y~X", data = data3)
##
## Residuals:
                                 3Q
##
       Min
                1Q Median
                                        Max
   -0.5333 -0.4043 -0.1373 0.4157
                                    0.8487
##
## Coefficients:
##
               Estimate Std. Error t value Pr(>|t|)
##
  (Intercept)
                 2.5753
                             0.2487 10.354 1.20e-07 ***
                -0.3240
                             0.0433 -7.483 4.61e-06 ***
## X
## ---
                  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
## Signif. codes:
##
## Residual standard error: 0.4743 on 13 degrees of freedom
## Multiple R-squared: 0.8116, Adjusted R-squared: 0.7971
## F-statistic: 55.99 on 1 and 13 DF, p-value: 4.611e-06
plot(data3$X,data3$Y)
abline(fit3)
             0
      0
      က
်
             0
      2
             0
     2
     2.0
data3$Y
     1.5
                              8
     1.0
      Ŋ
                                                8
      o.
                                                                 8
     0.0
                                                                                   0
                      2
                                                                          8
                                       4
                                                         6
                                            data3$X
```

(b) Perform the F test to determine whether or not there is lack of fit of a linear regression function; use $\alpha = .025$. State the alternatives, decision rule, and conclusion.

```
c = length(unique(data3$X))
n = length(data3$X)
SSER = sum(fit3$residuals^2)
SSEF = 0
for (i in unique(data3$X)) {
    SSEF <- SSEF + sum((data3[data3$X==i,]$Y-mean(data3[data3$X==i,]$Y))^2)
}
Fvalue = (SSER-SSEF)/(c-2)/(SSEF/(n-c))</pre>
```

```
print(sprintf('SSE of Reduced Model :%f',SSER))

## [1] "SSE of Reduced Model :2.924653"

print(sprintf('SSE of Full Model :%f',SSEF))

## [1] "SSE of Full Model :0.157400"

print(sprintf('F-value :%f',Fvalue))

## [1] "F-value :58.603417"

print(sprintf('0.975 Quantile F(%d,%d) value:%f',c-2,n-c,qf(0.975,c-2,n-c)))

## [1] "0.975 Quantile F(3,10) value:4.825621"
```

 H_0 : the regression function is linear H_a : the regression function is not linear

$$F^* = \frac{\frac{SSE(R) - SSE(F)}{3}}{\frac{SSE(F)}{10}} \sim F(3, 10)$$

If $F^* \leq F(0.975; 3, 10)$, then conclude H_0 ; otherwise conclude H_a .

Here, $F^* = 58.603417 > 4.825621$, conclude H_a .

(c)

Does the test in part (b) indicate what regression function is appropriate when it leads to the conclusion that lack of fit of a linear regression function exists? Explain.

No, since the alternative H_a includes all regression functions other than a linear one.

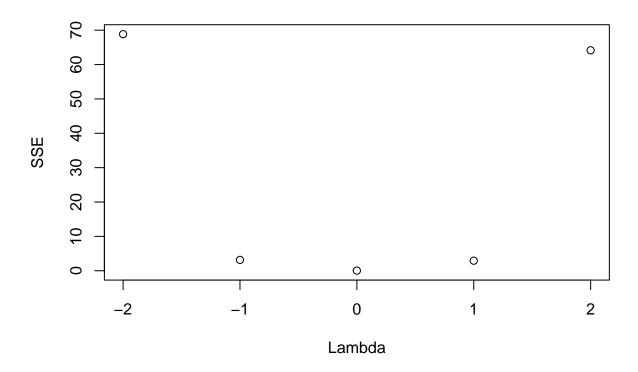
3.16

Refer to Solution concentration Problem 3.15.

(b) Use the Box—Cox procedure and standardization (3.36) to find an appropriate power transformation. Evaluate SSE for $\lambda = -.2, -.1, 0, .1, .2$. What transformation of Y is suggested?

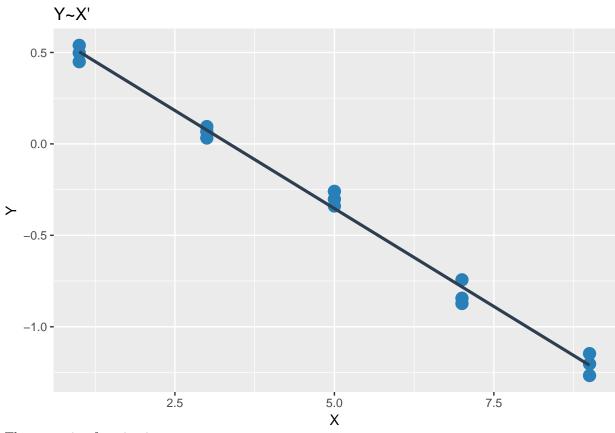
```
library(MASS,warn.conflicts = FALSE)
a = boxcox(fit3)
```

```
print(sprintf('The best power returned by Box-Cox is %f',a$x[which.max(a$y)]))
## [1] "The best power returned by Box-Cox is 0.020202"
SSE = c(0,0,0,0,0);
lam = c(-2,-1,0,1,2);
K2 = (prod(data3$Y))^(1/length(data3$Y));
print(K2)
## [1] 0.4762974
for (i in c(1:5)){
  data4 = data3
  if(lam[i]==0){
    data4\$Y = K2*log(data4\$Y)
  }
  else{
    K1 = \frac{1}{\text{lam}[i]} \frac{K2^{(lam[i]-1)}}{K2^{(lam[i]-1)}}
    data4\$Y = K1*(data4\$Y^lam[i] - 1)
  }
  SSE[i] = sum(lm('Y~X',data4)$residuals^2);
plot(c(-2:2),SSE,xlab='Lambda')
```



(c) Use the transformation $Y' = \log_{10} Y$ and obtain the estimated linear regression function for the transformed data.

```
data4 = data3
data4\$Y = K2*log(data4\$Y)
fit4 =lm('Y~X',data4)
summary(fit4)
##
## Call:
## lm(formula = "Y~X", data = data4)
##
## Residuals:
##
        Min
                   1Q
                          Median
                                        ЗQ
## -0.090981 -0.048717 0.007472 0.036753 0.093824
##
## Coefficients:
##
                Estimate Std. Error t value Pr(>|t|)
                                     25.01 2.22e-12 ***
## (Intercept) 0.718217
                           0.028713
## X
               -0.214299
                           0.004998 -42.88 2.19e-15 ***
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
## Residual standard error: 0.05475 on 13 degrees of freedom
## Multiple R-squared: 0.993, Adjusted R-squared: 0.9924
## F-statistic: 1838 on 1 and 13 DF, p-value: 2.188e-15
lm.scatter <- ggplot(data4, aes(x=X, y=Y)) +</pre>
  geom_point(color='#2980B9', size = 4) +
  geom_smooth(method = lm, se=FALSE, fullrange=TRUE, color='#2C3E50', size=1.1) +
  labs(title='Y~X\'')
grid.arrange(lm.scatter)
```



The regression function is

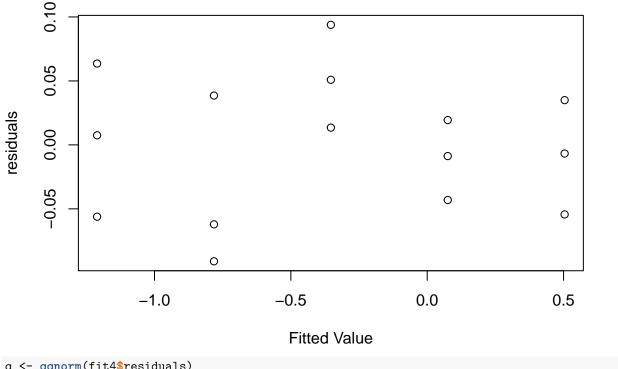
$$Y' = -0.214299 + 0.718217$$

i.e.

$$Y = e^{\frac{-0.44993X + 1.50799}{0.4762974}}$$

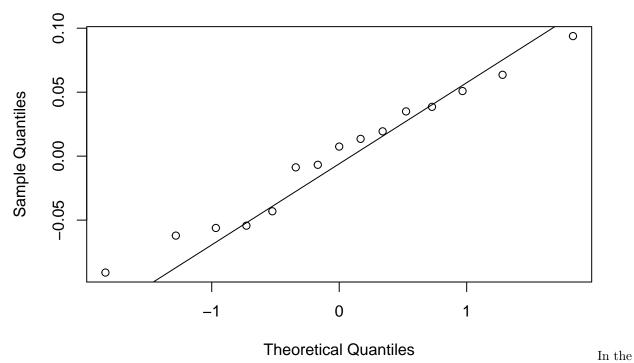
(e) Obtain the residuals and plot them against the fitted values. Also prepare a normal probability plot. What do your plots show?

plot(fit4\$fitted.values,fit4\$residuals,xlab = 'Fitted Value',ylab = 'residuals')



q <- qqnorm(fit4\$residuals)
qqline(fit4\$residuals)</pre>





first plot, residuals lie randomly near x-axis. In the second plot, the points lie near a straight line. Therefore, the plots show that the transformed model is linear and normal.

(f) Express the estimated regression function in the original units.

```
d = data.frame('X'=seq(0,10,0.1))
tyh = predict(fit4,d)
plot(d$X,exp(tyh/K2),xlab = 'X',ylab = 'Y',type = 'l')
points(data3$X,data3$Y)
```

