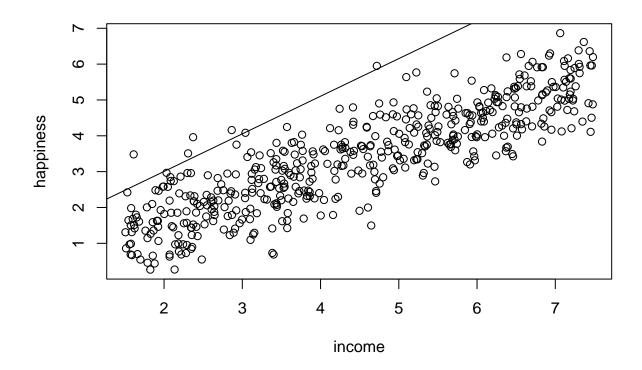
HW 03

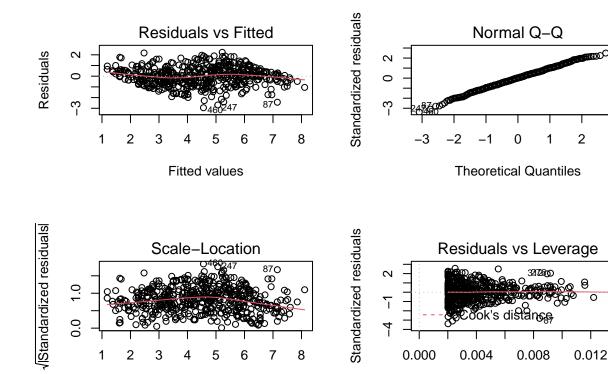
Tuan Bui

Question 01:

```
income_data <- read.csv('~/OneDrive - Stony Brook University/SBU/MAT + AMS/Fall 2021/AMS 380/hw/03/income_data</pre>
attach(income_data)
fit <- lm(income ~ happiness)</pre>
##
## Call:
## lm(formula = income ~ happiness)
##
## Coefficients:
## (Intercept)
                  happiness
        0.9053
                     1.0497
# a. The least square regression line equation: income = 0.9053 + 1.0497 * happiness
# b. Plot the points and regression line in the same figure
plot(income, happiness)
abline(fit)
```



```
# c. Check assumptions:
par(mfrow = c(2,2))
plot(fit)
```



Fitted values

Sample correlation coefficient is 0.8656337

```
## 2. Homoscedasticity: it is satisfied because the square root of standardized residuals is symmetrical
## 3. Independence: assume it is satisfied

## 4. Normality:
shapiro.test(residuals(fit))

##

## Shapiro-Wilk normality test
##

## data: residuals(fit)
## W = 0.99682, p-value = 0.4377

### p-value is 0.4377 greater than the significance level 0.05, so residuals is normal distributed, normal d
```

1. Linearity: it is satisfied because the residuals are symmetrically distributed around the 0-line

3

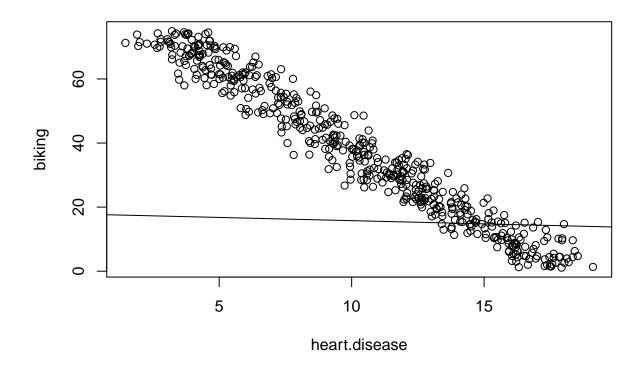
Leverage

```
## The corresponding population correlation test:
cor.test(income, happiness)
##
##
  Pearson's product-moment correlation
## data: income and happiness
## t = 38.505, df = 496, p-value < 2.2e-16
## alternative hypothesis: true correlation is not equal to 0
## 95 percent confidence interval:
## 0.8417942 0.8861031
## sample estimates:
        cor
## 0.8656337
### p-value is less than 2.2e-16, which is less than the significance level 0.05, reject HO. The correl
summary(fit)
##
## Call:
## lm(formula = income ~ happiness)
## Residuals:
       Min
                 1Q Median
                                   3Q
## -2.94796 -0.57730 0.02277 0.55661 2.23185
## Coefficients:
              Estimate Std. Error t value Pr(>|t|)
## (Intercept) 0.90533
                          0.10039
                                   9.018 <2e-16 ***
                          0.02726 38.505
## happiness
               1.04973
                                            <2e-16 ***
## Signif. codes: 0 '*** 0.001 '** 0.01 '* 0.05 '.' 0.1 ' ' 1
## Residual standard error: 0.8708 on 496 degrees of freedom
## Multiple R-squared: 0.7493, Adjusted R-squared: 0.7488
## F-statistic: 1483 on 1 and 496 DF, p-value: < 2.2e-16
## The coefficient of determination is 0.7493
## p-value for coefficient of happiness is less than 2.2e-16, which is less than the significance level
# f. ANOVA table of the regression
anova(fit)
## Analysis of Variance Table
##
## Response: income
             Df Sum Sq Mean Sq F value
## happiness 1 1124.32 1124.32 1482.6 < 2.2e-16 ***
## Residuals 496 376.13
                           0.76
## Signif. codes: 0 '*** 0.001 '** 0.01 '* 0.05 '.' 0.1 ' ' 1
```

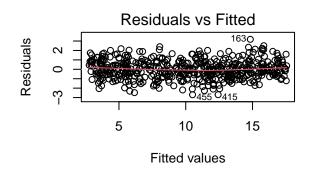
```
### The p-value for ANOVA F-test is less than 2.2e-16, which is less than the significance level 0.05, detach(income_data)
```

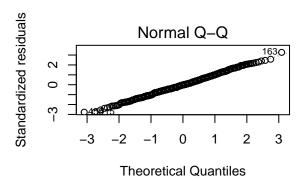
Question 02:

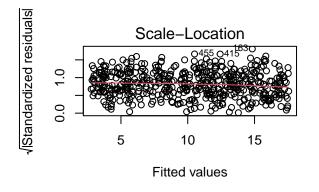
```
heart_data <- read.csv('~/OneDrive - Stony Brook University/SBU/MAT + AMS/Fall 2021/AMS 380/hw/03/heart
attach(heart_data)
fit <- lm (heart.disease ~ biking)</pre>
##
## Call:
## lm(formula = heart.disease ~ biking)
## Coefficients:
## (Intercept)
                     biking
       17.7779
                    -0.2003
##
# a. The least square regression line equation:
## heart.disease = 17.7779 - 0.2003 * biking
# b. Plot
plot(heart.disease, biking)
abline(fit)
```

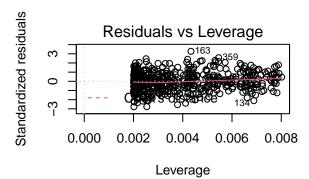


```
# c. Check assumptions:
par(mfrow = c(2,2))
plot(fit)
```









```
## 1. Linearity: it is satisfied because the residuals are symmetrically distributed around the O-line
## 2. Homoscedasticity: it is satisfied because the square root of standardized residuals is symmetrica
## 3. Independence: assume it is satisfied
## 4. Normality:
```

```
##
## Shapiro-Wilk normality test
##
## data: residuals(fit)
## W = 0.99801, p-value = 0.8351
```

shapiro.test(residuals(fit))

p-value is 0.8351 greater than the significance level 0.10, so residuals is normal distributed, nor
d. Sample correlation coefficient between the 2 variables:
cor(heart.disease, biking)

[1] -0.9753352

Sample correlation coefficient is -0.9753352

```
## The corresponding population correlation test:
cor.test(heart.disease, biking)
##
## Pearson's product-moment correlation
##
## data: heart.disease and biking
## t = -98.409, df = 496, p-value < 2.2e-16
## alternative hypothesis: true correlation is not equal to 0
## 95 percent confidence interval:
## -0.9792783 -0.9706530
## sample estimates:
##
         cor
## -0.9753352
### p-value is less than 2.2e-16, which is less than the significance level 0.10, reject HO. The correl
summary(fit)
##
## Call:
## lm(formula = heart.disease ~ biking)
## Residuals:
##
      Min
               1Q Median
                               3Q
## -2.6975 -0.6277 -0.0205 0.6482 3.1787
##
## Coefficients:
               Estimate Std. Error t value Pr(>|t|)
## (Intercept) 17.777880
                         0.088450 200.99 <2e-16 ***
## biking
              -0.200297
                          0.002035 -98.41
                                             <2e-16 ***
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
## Residual standard error: 0.9747 on 496 degrees of freedom
## Multiple R-squared: 0.9513, Adjusted R-squared: 0.9512
## F-statistic: 9684 on 1 and 496 DF, p-value: < 2.2e-16
## The coefficient of determination is 0.9513
## p-value for coefficient of biking is less than 2.2e-16, which is less than the significance level 0.
# f. The percentage of people in the town who have heart disease if the percentage of people who bike t
heart.disease_rate <- 17.7779 - 0.2003 * 65
heart.disease_rate
## [1] 4.7584
## There are 4.7584% people in the town who have heart disease if the percentage of people who bike to
# g. The 90% confidence interval:
confint(fit, level = 0.90)
```

```
## 5 % 95 %
## (Intercept) 17.6321212 17.9236392
## biking -0.2036511 -0.1969429
```

detach(heart_data)