STP530 HW4 Solution

```
2.25
See the R codes below to calculate SSR, SSE, and SSTO in the ANOVA table. Follow
the example in textbook Table 2.2 to construct the ANOVA table.
# Set the working directory to where you saved the data file
setwd("~/Documents/ASU/STP530-YiZheng/HW-HaozhenXu/HW4")
HW2_25.data <- read.table("CH01PR21.txt")</pre>
colnames(HW2_25.data) \leftarrow c("Y", "X")
head(HW2_25.data)
mod1 <- lm(Y ~ X, data=HW2_25.data)</pre>
summary(mod1)
# Calculate SSTO
SSR <- sum((predict(mod1) - mean(HW2_25.data$Y)) ^ 2)
SSE <- sum((HW2_25.data$Y - predict(mod1)) ^ 2)</pre>
SSTO \leftarrow sum((HW2_25.data\$Y - mean(HW2_25.data\$Y)) ^ 2)
# You can also use the anova() function in R to see a part of the ANOVA table
anova(mod1)
See R outputs below.
> SSR
[1] 160
> SSE
[1] 17.6
SST0
[1] 177.6
```

> anova(mod1)

Analysis of Variance Table

Response: Y

Df Sum Sq Mean Sq F value Pr(>F) X 1 160.0 160.0 72.727 2.749e-05 ***

Residuals 8 17.6 2.2

Now we can construct the ANOVA table as below.

Source of Variation	SS	df	MS
Regression	160.0	1	160.0
Error	17.6	8	2.2
Total	177.6	9	19.7

From the table above, we can see that the sums of square of regression and error and their corresponding degrees of freedom are additive.

b.

Five steps of F test:

• Assumptions:

$$\varepsilon i.i.d. \sim N(0, \sigma^2)$$

• Hypotheses:

$$H_0: \beta_1 = 0 \quad H_1: \beta_1 \neq 0$$

• Test-statistic:

$$F^* = \frac{MSR}{MSE} = \frac{160}{2.2} = 72.727$$

- P-value: Here the P-value is $P_{H_0}\{F^* > 72.727\} = 2.75 \times 10^{-5}$. Use R code 1 pf(q=72.727, df1=1, df2=8)
- Conclusion: Because the P-value = $2.75 \times 10^{-5} < \alpha = 0.05$, we reject H_0 at a significance level of 0.05. The population slope of this linear model is indeed non-zero.

c.

Using command summary() in R to view results of the fitted model, we have $t^* = 8.528$, and we can find $t^{*2} = 8.529^2 = 72.7 = F^*$ in (b). Also notice the p-values of these two tests are identical.

> summary(mod1)

Call:

 $lm(formula = Y \sim X, data = hwdata1)$

Residuals:

Coefficients:

Estimate Std. Error t value Pr(>|t|)
(Intercept) 10.2000 0.6633 15.377 3.18e-07 ***
X 4.0000 0.4690 8.528 2.75e-05 ***

Signif. codes: 0 '*** 0.001 '** 0.01 '* 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

d.

We can calculate \mathbb{R}^2 by using the values we entered in the ANOVA table:

$$R^2 = \frac{SSR}{SSTO} = 1 - \frac{SSE}{SSTO} = 1 - \frac{17.6}{177.6} = 0.9009$$

Also from the result in R, we have that $R^2 = 0.9009$, and thus $r = \sqrt{R^2} = 0.9492$. By the definition of R^2 , 90.09% of the variation in the response Y is explained by this fitted model.

3.25

We can use the code below to fit models and get residual plots and Q-Q plots respectively.

```
HW3_25.data <- read.table("APPENCO2.txt")</pre>
head(HW3_25.data)
# change column names
colnames(HW3_25.data)[8] <- "Y"
colnames(HW3_25.data)[5] <- "X1"</pre>
colnames(HW3_25.data)[9] <- "X2"</pre>
colnames(HW3_25.data)[16] <- "X3"
#fit three models respectively
fit1 < -lm(Y \sim X1, data = HW3_25.data)
fit2 < -lm(Y \sim X2, data = HW3_25.data)
fit3 < -lm(Y \sim X3, data = HW3_25.data)
# obtain residuals and residual plots and Q-Q plot respectively
plot(HW3_25.data$X1, fit1$residuals, xlab="Total Population")
gqnorm(fit1$residuals)
qqline(fit1$residuals, col='red')
plot(HW3_25.data$X2, fit2$residuals, xlab="Number of Hospital Beds")
qqnorm(fit2$residuals)
qqline(fit2$residuals, col='red')
plot(HW3_25.data$X3, fit3$residuals, xlab="Total Personal Income")
qqnorm(fit3$residuals)
qqline(fit3$residuals, col='red')
```

See the figures on the next page.

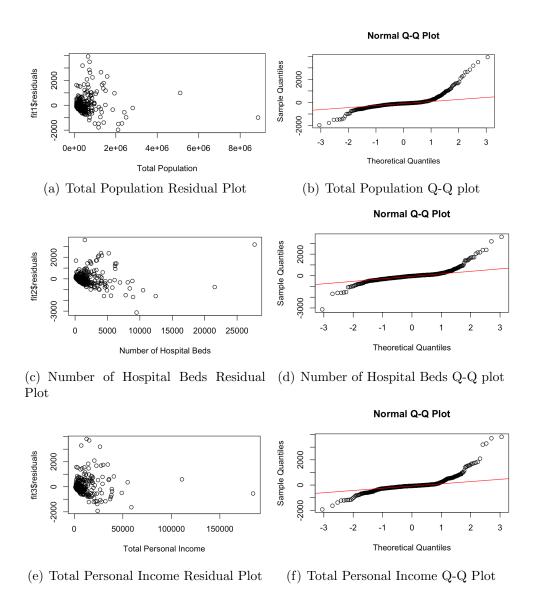


Figure 1: Residual Plots and Q-Q Plots

From the figures above, none of the three simple linear regression models seems a very good fit. For all three models, the residuals tend to distribute with a wider range when the value of X increases (i.e., the funnel shape), and the Q-Q plots are not very close to the straight reference lines on both ends.

None of the three models looks obviously more appropriate than the others.