Chapter 8

8.1

For the following data, use R to verify that the least squares regresion line is $\hat{Y}=1.8X-8.5$

```
X: 5, 8, 9, 7, 14 Y: 3, 1, 6, 7, 19
```

```
dat <- data.table(
    X = c(5, 8, 9, 7, 14),
    Y = c(3, 1, 6, 7, 19))

lm(Y ~ X, data = dat)</pre>
```

```
Call:
```

```
lm(formula = Y ~ X, data = dat)
```

Coefficients:

```
(Intercept) X
-8.478 1.823
```

-5.730159 1.746032

Also verify that the Theil-Sen estimator, the slope is estimated to be 1.746 and the intercept is estimated to be -7.968.

```
tsreg(dat$X, dat$Y)$coef
Intercept
```

8.2

Using the R function *Isfit*, compute the residuals using the data in E1,

Verify that if you square and sum the residusls, you get 46.585.

```
res <- lsfit(dat$X, dat$Y)$resid
sum(res^2)
[1] 46.58407
```

8.3

Verify that for the data in E1, if you use $\hat{Y}=2X-9$, the sum of the squared residuals is greater than 46.584.

```
Yhat <- 2*dat$X - 9
res <- sum( (dat$Y - Yhat)^2 )
```

```
stopifnot(res > 46.583)
res
```

[1] 53

Why would you expect a value greater than 46.584?

The x coefficent increased.

8.4

Suppose that based on n=25 values, $s_x^2=12$ and $\sum (X_i-\bar{X})(Y_i-\bar{Y})=144.$

What is the slope of the least squares regression?

$$A=144, C=(n-1)s_x^2=288, b_1=A/C=144/288=.5$$

8.5

The following table reports breast cancer rates plus levels of solar radiation (in calories per day) for various cities in the United States. The data are stored in the file cancer rate dat.txt.

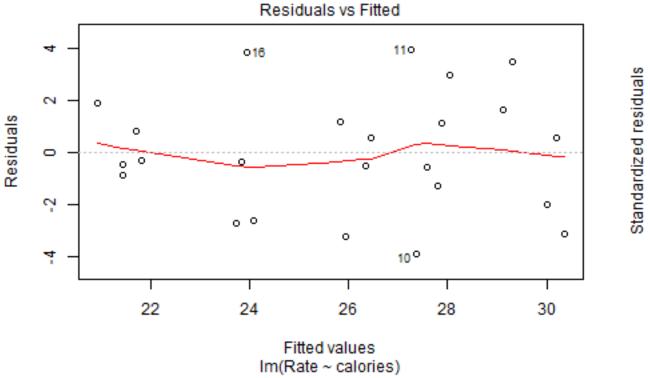
```
dat <- data.table::fread(paste0(data.dir, "cancer_rate_dat.txt"), fill = T, sep = "&")
dat</pre>
```

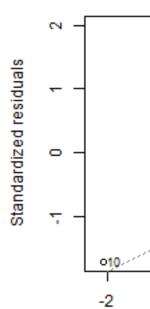
	City	Rate	calories
1:	New York	32.75	300
2:	Chicago	30.75	275
3:	Pittsburgh	28.00	280
4:	Seattle	27.25	270
5:	Boston	30.75	305
6:	Cleveland	31.00	335
7:	Columbus	29.00	340
8:	Indianapolis	26.50	342
9:	New Orleans	27.00	348
10:	Nashville	23.50	354
11:	Washington, DC	31.20	357
12:	Salt Lake City	22.70	394
13:	Omaha	27.00	380
14:	San Diego	25.80	383
15:	Atlanta	27.00	397
16:	Los Angeles	27.80	450
17:	Miami	23.50	453
18:	Fort Worth	21.50	446
19:	Tampa	21.00	456
20:	Albuquerque	22.50	513

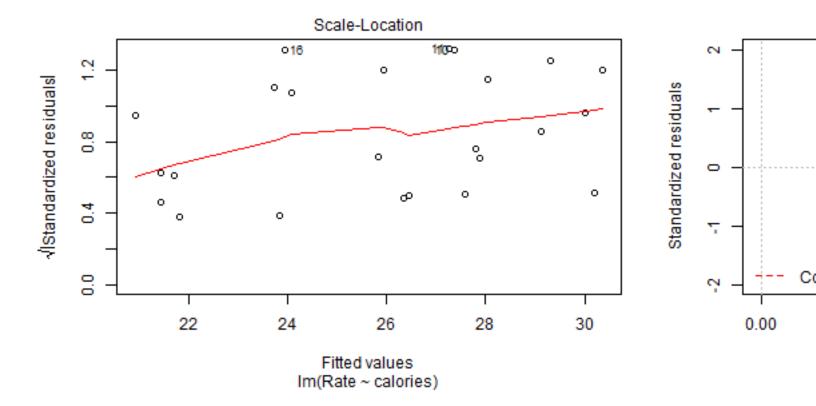
```
21: Las Vegas 21.50 510
22: Honolulu 20.60 520
23: El Paso 22.80 535
24: Phoenix 21.00 520
City Rate calories
```

Fit a OLS regression to predict cancer rates and comment on what this suggests.

```
fit <- lm(Rate ~ calories, data = dat)
plot(fit)</pre>
```







For the following data, use R to compute the least squares regresion line for predicting GPA given SAT.

SAT: 500, 530, 590, 660, 610, 700, 570, 640 GPA: 2.3, 3.1, 2.6, 3.0, 2.4, 3.3, 2.6, 3.5

```
dat <- data.table(
    SAT = c(500, 530, 590, 660, 610, 700, 570, 640),
    GPA = c(2.3, 3.1, 2.6, 3.0, 2.4, 3.3, 2.6, 3.5)
)

fit <- lm(GPA ~ SAT, data = dat)
coef(fit)

(Intercept)    SAT
0.484615385 0.003942308</pre>
```

8.7

Compute the residuals for the data used in the previous problem and verify that sum to zero.

```
round(sum((dat$GPA - fit$fitted.values)), 5)
[1] 0
```

For the following data, use R to compute the least squares regression line for predicting Y from X.

X: 40, 41, 42, 43, 44, 45, 46 Y: 1.62, 1.63, 1.90, 2.64, 2.05, 2.13, 1.94

```
dat <- data.table(
    X = c(40, 41, 42, 43, 44, 45, 46),
    Y = c(1.62, 1.63, 1.90, 2.64, 2.05, 2.13, 1.94)
)
summary(fit <- lm(Y ~ X, data = dat))</pre>
```

```
Call:
```

lm(formula = Y ~ X, data = dat)

Residuals:

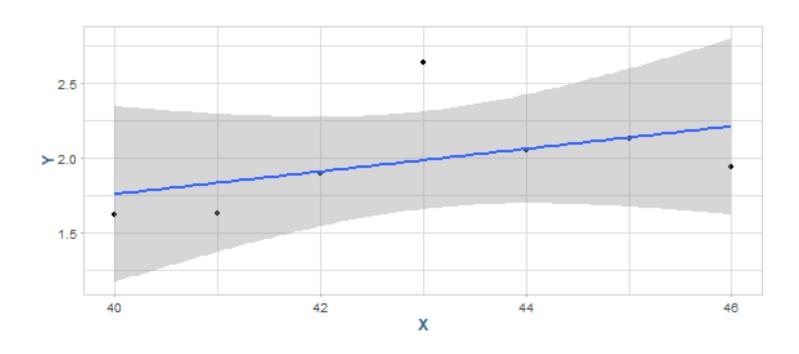
```
1 2 3 4 5 6 7
-0.141071 -0.206429 -0.011786 0.652857 -0.012500 -0.007857 -0.273214
```

Coefficients:

```
Estimate Std. Error t value Pr(>|t|)
(Intercept) -1.25321 2.73157 -0.459 0.666
X 0.07536 0.06346 1.188 0.288
```

Residual standard error: 0.3358 on 5 degrees of freedom Multiple R-squared: 0.22, Adjusted R-squared: 0.064 F-statistic: 1.41 on 1 and 5 DF, p-value: 0.2883

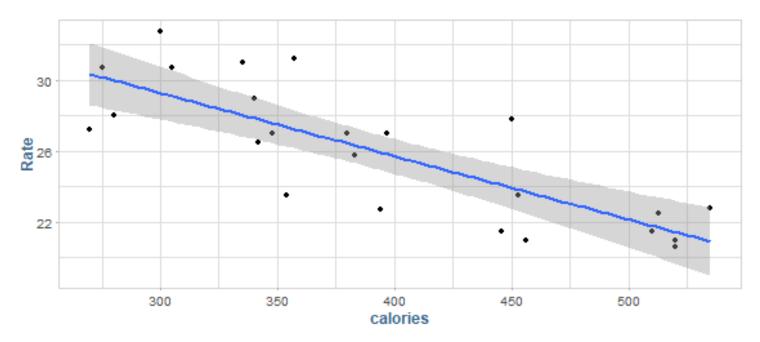
```
ggplot(dat, aes(X, Y)) +
  geom_point() +
  geom_smooth(method = "lm")
```



In exercise 5, what would be the least squares estimate of the cancer rate given a solar radiation of 600?

```
dat <- data.table::fread(paste0(data.dir, "cancer_rate_dat.txt"), fill = T, sep = "&")

ggplot(dat, aes(calories, Rate)) +
    geom_point() +
    geom_smooth(method = "lm")</pre>
```



```
fit <- lm(Rate ~ calories, data = dat)

coef(fit)

(Intercept) calories
39.99094634 -0.03565283

39.99 -0.037*600

[1] 17.79</pre>
```

December 24, 2019

Why might this be unreasonable?

Because 600 is outside of the bounds of seen values (extrapolation).

8.10

Maximal oxygen uptake (MOU) is a measure of an individual's physical fitness. You want to know how MOU is related to how fast someone can run a mile. Suppose you randomly sample six athletes and get:

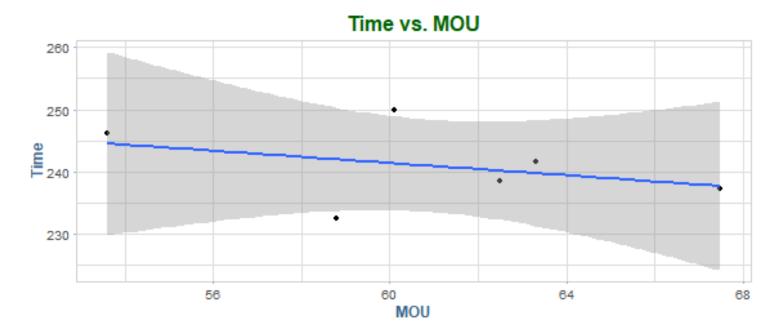
MOU (kl/kg): 63.3, 60.1, 53.6, 58.8, 67.5, 62.5 Time (seconds): 241.5, 249.8, 246.1, 232.4, 237.2, 238.4

Compute the least squres regression line and comment on what the results suggest.

```
dat <- data.table(
    MOU = c(63.3, 60.1, 53.6, 58.8, 67.5, 62.5),
    Time = c(241.5, 249.8, 246.1, 232.4, 237.2, 238.4)
)

fit <- lm(Time ~ MOU, data = dat)

ggplot(dat, aes(MOU, Time)) +
    geom_point() +
    geom_smooth(method = "lm") +
    labs(title = "Time vs. MOU")</pre>
```



Generally, time decreases as MOU increases.

8.11

Verify that for the following pairs of points, the least squares regression line has a slope of zero. Plot the points and comment on the assumption that the regression line is straight.

```
X: 1, 2, 3, 4, 5, 6 Y: 1, 4, 7, 7, 4, 1
```

8.12

Repeat the last exercise, only for points:

```
X: 1, 2, 3, 4, 5, 6 Y: 4, 5, 6, 7, 8, 2
```

```
dat <- data.table(
    X = c(1, 2, 3, 4, 5, 6),</pre>
```

Vitamin A is required for good health. However, one bite of polar bear liver results in death becasue it contains a high concentration of vitamin A.

Comment on this fact in terms of extrapolation.

Vitaim A as a function of health is a bounded variable. Predicting a response when the independent variables are far from what has been observed can lead to undefined results (death).

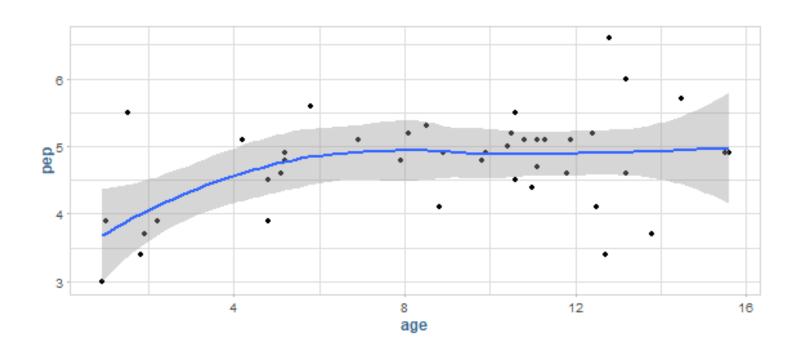
8.14

Socket et al. (1987) report data related to patterns of residual insulin secretion in children. A portion of the study was concerned with whether age can be used to predict the logarithm of C-peptide concentrations at diagnosis. The observed values are (data file):

Replicate the LOESS smoothed curve in fig 8.4.1.

```
dat <- data.table::fread(paste0(data.dir, "diabetes_sockett_dat.txt"), fill = T)

ggplot(dat, aes(age, pep)) +
    geom_point() +
    geom_smooth(method = "loess")</pre>
```



- 8.15
- 8.16
- 8.17
- 8.18
- 8.19
- 8.20
- 8.21
- 8.22
- 8.23
- 8.24
- 8.25
- 8.26
- 8.27
- 8.28
- 8.29
- 8.30
- 8.31
- 8.32
- 8.33
- 8.34
- 8.35
- 8.36