## Homework Assignment 1

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Q1: (a) The predictor variable is the population. The response variable is the number of confirmed COVID-19 cases.

(b) Since 
$$\hat{\beta}_1 = \frac{\sum\limits_{i=1}^n (x_i - \bar{x})(y_i - \bar{y})}{\sum\limits_{i=1}^n (x_i - \bar{x})^2}$$
 and  $\hat{\beta}_0 = \bar{y} - \hat{\beta}_1 * \bar{x}$ , we can calculate that  $\hat{\beta}_0 = 2.85395$  and  $\hat{\beta}_1 = 0.07236$ .

```
cov.data <- data.frame(row.names = c("San Bernardino County", "Riverside County",</pre>
                    "Orange County", "San Diego County", "Santa Clara County", "Kern County", "Sacramento County",
                    "Fresno County", "Alameda County"))
cov.data$population <- c(2149, 2411, 3168, 3316, 1927, 887, 1525, 985, 1657)
cov.data$confirmed <- c(251, 239, 223, 212, 93, 85, 79, 81, 67)
x <- cov.data$population
y <- cov.data$confirmed
betaone <- ((x[1] - mean(x)) * (y[1] - mean(y)) + (x[2] - mean(x)) * (y[2] - mean(y)) +
                     (x[3] - mean(x)) * (y[3] - mean(y)) + (x[4] - mean(x)) * (y[4] - mean(y)) + (x[5] - mean(y))
                    mean(x)) * (y[5] - mean(y)) + (x[6] - mean(x)) * (y[6] - mean(y)) + (x[7] - mean(x)) *
                    (y[7] - mean(y)) + (x[8] - mean(x)) * (y[8] - mean(y)) + (x[9] - mean(x)) * (y[9] - mea
                   mean(y))/((x[1] - mean(x))^2 + (x[2] - mean(x))^2 + (x[3] - mean(x))^2 + (x[4] -
                   mean(x))^2 + (x[5] - mean(x))^2 + (x[6] - mean(x))^2 + (x[7] - mean(x))^2 + (x[8] - mean(x)
                   mean(x))^2 + (x[9] - mean(x))^2
betazero <- mean(y) - betaone * mean(x)
betaone
```

## [1] 0.07236

## betazero

## [1] 2.854

- $(c)\hat{\beta}_1$  is the average change of people confirmed for every 1000 increase in population.  $\hat{\beta}_0$  is the number of confirmed people when population is zero.
  - (d) Simple linear regression model: y = 2.854 + 0.07236\*x
  - (e) When x = 1000000, y = 72364

```
predicted <- betazero + 1e+06 * betaone
predicted</pre>
```

## [1] 72364

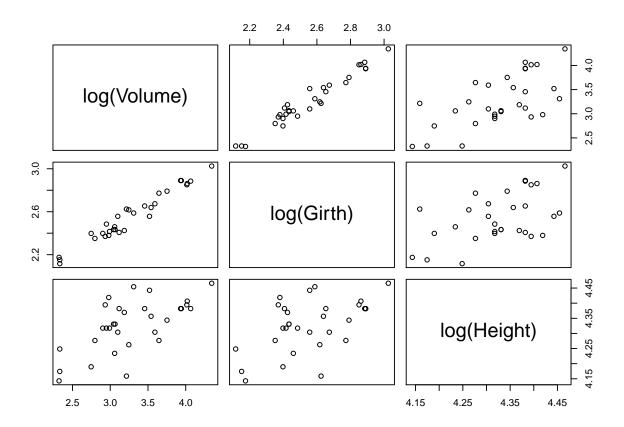
Q2

trees

```
##
      Girth Height Volume
## 1
                70
                      10.3
        8.3
## 2
        8.6
                65
                      10.3
## 3
        8.8
                63
                      10.2
## 4
       10.5
                72
                      16.4
       10.7
## 5
                81
                      18.8
## 6
       10.8
                83
                      19.7
## 7
       11.0
                66
                      15.6
## 8
       11.0
                75
                      18.2
## 9
       11.1
                      22.6
                80
## 10
       11.2
                75
                      19.9
## 11
      11.3
                79
                      24.2
## 12 11.4
                      21.0
                76
## 13 11.4
                76
                      21.4
## 14
      11.7
                69
                      21.3
## 15
      12.0
                75
                      19.1
## 16
      12.9
                74
                      22.2
## 17
       12.9
                      33.8
                85
## 18
      13.3
                86
                      27.4
## 19
       13.7
                71
                      25.7
## 20
      13.8
                      24.9
                64
## 21
       14.0
                78
                      34.5
## 22
       14.2
                80
                      31.7
## 23
       14.5
                74
                      36.3
## 24
       16.0
                72
                      38.3
## 25
       16.3
                77
                      42.6
## 26
      17.3
                81
                      55.4
## 27
      17.5
                82
                      55.7
## 28
      17.9
                80
                      58.3
## 29
       18.0
                80
                      51.5
## 30
       18.0
                80
                      51.0
## 31 20.6
                      77.0
                87
```

- (a) There are 31 rows and 3 columns in the dataset "trees". The variable names are "Girth", "Height" and "Volume".
- (b) Here is the scatterplot:

```
pairs(formula = log(Volume) ~ log(Girth) + log(Height), data = trees)
```



```
(c)
cor(log(trees), method = c("pearson", "kendall", "spearman"))

## Girth Height Volume
## Girth 1.0000 0.5302 0.9767
## Height 0.5302 1.0000 0.6486
## Volume 0.9767 0.6486 1.0000

(d) No, there are no missing values.

is.na(cor(log(trees)))
```

```
## Girth Height Volume
## Girth FALSE FALSE FALSE
## Height FALSE FALSE FALSE
## Volume FALSE FALSE FALSE

(e)

mod <- lm(log(Volume) ~ log(Girth) + log(Height), data = trees)
summary(mod)</pre>
```

```
##
## Call:
## lm(formula = log(Volume) ~ log(Girth) + log(Height), data = trees)
## Residuals:
##
       \mathtt{Min}
                  1Q Median
                                    3Q
                                            Max
## -0.16856 -0.04849 0.00243 0.06364 0.12922
##
## Coefficients:
##
               Estimate Std. Error t value Pr(>|t|)
## (Intercept)
                -6.632
                           0.800
                                    -8.29 5.1e-09 ***
                  1.983
                             0.075
                                     26.43 < 2e-16 ***
## log(Girth)
## log(Height)
                             0.204
                                      5.46 7.8e-06 ***
                  1.117
## ---
## Signif. codes: 0 '*** 0.001 '** 0.01 '* 0.05 '.' 0.1 ' 1
## Residual standard error: 0.0814 on 28 degrees of freedom
## Multiple R-squared: 0.978, Adjusted R-squared: 0.976
## F-statistic: 613 on 2 and 28 DF, p-value: <2e-16
 (f)
mod$coefficients
## (Intercept) log(Girth) log(Height)
##
       -6.632
                     1.983
                                 1.117
X = model.matrix(mod)
Y = log(trees) $Volume
(beta_hat = solve(t(X) %*% X) %*% t(X) %*% Y)
##
                 [,1]
## (Intercept) -6.632
## log(Girth)
               1.983
## log(Height) 1.117
These two results are the same above.
 (g)
hat_y <- X %*% beta_hat
head(hat_y)
      [,1]
##
## 1 2.310
## 2 2.298
## 3 2.309
## 4 2.808
## 5 2.977
## 6 3.023
```

head(mod\$residuals)

head((t(mod\$residuals) %\*% mod\$residuals/28))

Q3 The procedure is below:

$$SSR = \sum_{i=1}^{n} (y_i - \beta_0 - \beta_1 x_i)^2$$

The derivative of SSR with respect to  $\beta_0$  is

$$\frac{d}{d\hat{\beta}_0} \sum_{i=1}^n (y_i - \beta_0 - \beta_1 x_i)^2 = -2 \sum_{i=1}^n (y_i - \beta_0 - \beta_1 x_i) = -2 \sum_{i=1}^n y_i + 2n\beta_0 + 2\beta_1 \sum_{i=1}^n x_i = -2n\bar{y} + 2n\hat{\beta}_0 + 2n\hat{\beta}_1 \bar{x}$$

Set this derivative = 0, we can get

$$-2n\bar{y} + 2n\hat{\beta}_0 + 2n\hat{\beta}_1\bar{x} = 0.$$

So

$$\hat{\beta_0} = \bar{y} - \hat{\beta_1} * \bar{x}$$

The derivative of SSR with respect to  $\beta_1$  is

$$\frac{d}{d\hat{\beta}_1} \sum_{i=1}^n \left( y_i - \beta_0 - \beta_1 x_i \right)^2 = -2x_i \sum_{i=1}^n \left( y_i - \hat{\beta}_0 - \hat{\beta}_1 x_i \right) = -2\sum_{i=1}^n x_i y_i + 2\hat{\beta}_0 \sum_{i=1}^n x_i + 2\hat{\beta}_1 \sum_{i=1}^n x_i^2 + 2\hat{\beta}_0 \sum_{i=1}^n x_i + 2\hat{\beta}_0 \sum_{i=1}^n x_$$

Set this derivative = 0, we can get

$$-2\sum_{i=1}^{n} x_i y_i + 2\hat{\beta}_0 \sum_{i=1}^{n} x_i + 2\hat{\beta}_1 \sum_{i=1}^{n} x_i^2 = 0.$$

So

$$-\sum_{i=1}^{n} x_i y_i + (\bar{y} - \hat{\beta}_1 \bar{x}) \sum_{i=1}^{n} x_i + \hat{\beta}_1 \sum_{i=1}^{n} x_i^2 = 0$$

$$\hat{\beta}_1 = \frac{\bar{y} \sum_{i=1}^{n} x_i - \sum_{i=1}^{n} x_i y_i}{\bar{x} \sum_{i=1}^{n} x_i - \sum_{i=1}^{n} x_i^2} = \frac{n \bar{x} \bar{y} - \sum_{i=1}^{n} x_i y_i}{n \bar{x}^2 - \sum_{i=1}^{n} x_i^2} = \frac{\sum_{i=1}^{n} (x_i - \bar{x})(y_i - \bar{y})}{\sum_{i=1}^{n} (x_i - \bar{x})^2}$$

Proved