INF 511 Assignment 2

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Question 1

(a)

```
sapflow.df<- readRDS(file="sapflow.rds")
str(sapflow.df)

## 'data.frame': 16 obs. of 5 variables:
## $ V1: num 1 2 1 2 1 2 1 2 1 2 ...
## $ V2: Factor w/ 2 levels "lo", "hi": 2 2 1 1 2 2 2 2 1 1 ...
## $ V3: Factor w/ 2 levels "lo", "hi": 1 1 1 1 2 2 2 2 2 1 1 ...
## $ V4: Factor w/ 2 levels "lo", "hi": 2 2 2 2 2 2 2 1 1 ...
## $ V5: num 192 248 168 204 303 ...</pre>
```

V2 seems to be a binary variable that would most likely take on a binomial distribution because there are only two options.

(b)

```
names(sapflow.df) = c("light", "fertilizer", "temperature", "moisture", "sapflow")
head(sapflow.df)
```

```
##
     light fertilizer temperature moisture sapflow
## 1
        1
                                             192.5
                  hi
                               10
                                       hi
## 2
        2
                  hi
                              lo
                                       hi
                                             248.3
## 3
        1
                                             168.4
                  lo
                              lo
                                       hi
        2
## 4
                  lo
                              lo
                                       hi
                                             204.2
## 5
        1
                  hi
                              hi
                                       hi
                                             302.8
## 6
                  hi
                              hi
                                       hi
                                             341.2
```

(c)

```
sapflow.df$light = factor(sapflow.df$light, labels = c("lo","hi"))
head(sapflow.df)
```

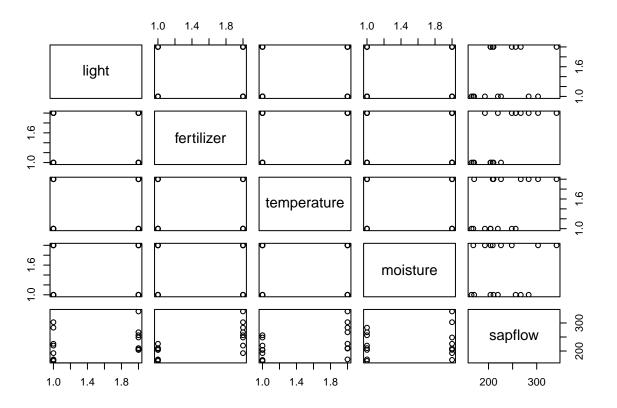
```
##
     light fertilizer temperature moisture sapflow
## 1
        10
                    hi
                                 10
                                           hi
                                                 192.5
## 2
        hi
                    hi
                                                 248.3
                                 10
                                           hi
## 3
                    10
                                 10
                                                 168.4
        10
                                           hi
## 4
        hi
                    10
                                 10
                                           hi
                                                 204.2
## 5
                    hi
                                           hi
                                                 302.8
        10
                                 hi
## 6
        hi
                                 hi
                                           hi
                                                 341.2
```

str(sapflow.df)

```
## 'data.frame': 16 obs. of 5 variables:
## $ light : Factor w/ 2 levels "lo", "hi": 1 2 1 2 1 2 1 2 1 2 1 2 ...
## $ fertilizer : Factor w/ 2 levels "lo", "hi": 2 2 1 1 2 2 1 1 2 2 ...
## $ temperature: Factor w/ 2 levels "lo", "hi": 1 1 1 1 2 2 2 2 2 1 1 ...
## $ moisture : Factor w/ 2 levels "lo", "hi": 2 2 2 2 2 2 2 1 1 ...
## $ sapflow : num 192 248 168 204 303 ...
```

(d)

```
pairs(sapflow.df)
```



The pairs function reports the factors as their numeric value of 1 or 2.

(e)

A single histogram may not be enough to assess if sapflow is normally distributed because it is just a visual assessment of normality which is inherently subjective. Creating a histogram is not formally or statistically assessing normality.

Question 2

(a)

```
X<- model.matrix(sapflow ~ light + fertilizer + temperature + moisture, data=sapflow.df)
y<- sapflow.df$sapflow</pre>
```

$$E(Y_1 \mid x_1) = \beta_0 * X_{1,0} + \beta_1 * X_{1,1} + \beta_2 * X_{1,2} + \beta_3 * X_{1,3} + \beta_4 * X_{1,4}$$

$$E(Y_1 \mid x_1) = \beta_0 * 1 + \beta_1 * 0 + \beta_2 * 1 + \beta_3 * 0 + \beta_4 * 1$$

$$E(Y_1 \mid x_1) = \beta_0 + \beta_2 + \beta_4$$

(b)

$$\begin{split} & E(Y_5 \mid x_5) = \beta_0 * X_{5,0} + \beta_1 * X_{5,1} + \beta_2 * X_{5,2} + \beta_3 * X_{5,3} + \beta_4 * X_{5,4} \\ & E(Y_5 \mid x_5) = \beta_0 * 1 + \beta_1 * 0 + \beta_2 * 1 + \beta_3 * 1 + \beta_4 * 1 \\ & E(Y_5 \mid x_5) = \beta_0 + \beta_2 + \beta_3 + \beta_4 \\ & E(Y_5 \mid x_5) - E(Y_1 \mid x_1) = \beta_0 + \beta_2 + \beta_3 + \beta_4 - \beta_0 - \beta_2 - \beta_4 \\ & E(Y_5 \mid x_5) - E(Y_1 \mid x_1) = \beta_3 \end{split}$$

The difference mean sapflow between high and low temperature and other inputs is equal to β_3 .

(c)

```
model = lm(sapflow ~ light + fertilizer + temperature + moisture, data = sapflow.df)
summary(model)
```

```
##
## Call:
## lm(formula = sapflow ~ light + fertilizer + temperature + moisture,
       data = sapflow.df)
##
##
## Residuals:
##
       Min
                1Q
                                 3Q
                    Median
                                        Max
##
  -43.387 -16.863
                     5.125
                             16.069
                                     34.787
##
## Coefficients:
                 Estimate Std. Error t value Pr(>|t|)
##
## (Intercept)
                   151.96
                                14.73 10.318 5.4e-07 ***
## lighthi
                    26.60
                                        2.019 0.068511 .
                                13.17
## fertilizerhi
                    69.30
                                13.17
                                        5.261 0.000268 ***
```

```
## temperaturehi 43.92 13.17 3.334 0.006660 **
## moisturehi 14.63 13.17 1.110 0.290602
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 26.35 on 11 degrees of freedom
## Multiple R-squared: 0.8004, Adjusted R-squared: 0.7278
## F-statistic: 11.03 on 4 and 11 DF, p-value: 0.0007654
```

The model tells us that when moving from low to high temperature there is a 43.92 mL/h increase in sapflow.

(d)

- i. $\hat{\beta}_3 = 43.92$
- ii. SE $\hat{\beta}_3 = 13.17$
- iii. t-value $\hat{\beta}_3 = 3.334$
- iv. 11 degrees of freedom
- \mathbf{v} . p-value = 0.006660
- vi. Yes, the effect of temperature is statistically significant because the t-value is far into the tails of the t distribution, therefore the p-value is small and less than alpha.

(e)

```
xtxi = solve(t(X) %*% X)
betahat = xtxi %*% t(X) %*% y
betahat
## [,1]
```

```
## (Intercept) 151.9625
## lighthi 26.6000
## fertilizerhi 69.3000
## temperaturehi 43.9250
## moisturehi 14.6250
```

coef(model)

```
## (Intercept) lighthi fertilizerhi temperaturehi moisturehi
## 151.9625 26.6000 69.3000 43.9250 14.6250
```

The beta hats are very similar from the manual calculations and the model created with lm.

(f)

```
S = t(X)\% * \%X
Xinv = solve(S)
H = X \% *\% Xinv \% *\% t(X)
Н
                                              7
##
                     3
                                  5
                                        6
                                                    8
                       ## 1
     0.3125
           0.1875
                 0.1875
           0.3125
                 0.0625
                       0.1875
                              0.0625
                                    0.1875 -0.0625
                                                0.0625
## 2
     0.1875
## 3
          0.0625
                       0.1875 0.0625 -0.0625
                                          0.1875
     0.1875
                 0.3125
                                                0.0625
                                                      0.0625
                 0.0625
     0.0625
           0.1875
                                                0.1875 -0.0625
                 0.0625 -0.0625
## 5
     0.1875
           0.0625
                             0.3125 0.1875
                                          0.1875
                                                0.0625 0.0625
                             0.1875 0.3125
                                          0.0625
## 6
     0.0625
           0.1875 -0.0625 0.0625
                                                0.1875 -0.0625
## 7
     0.0625 -0.0625 0.1875 0.0625 0.1875 0.0625
                                         0.3125
                                                0.1875 -0.0625
                0.0625  0.1875  0.0625  0.1875  0.1875  0.3125 -0.1875
    -0.0625
          0.0625
## 9
     0.1875
           ## 10
     0.1875
## 11 0.0625 -0.0625 0.1875 0.0625 -0.0625 -0.1875 0.0625 -0.0625
                                                      0.1875
0.0625
## 13 0.0625 -0.0625 -0.0625 -0.1875 0.1875 0.0625 0.0625 -0.0625
                                                      0.1875
0.0625
                                                      0.0625
## 15 -0.0625 -0.1875  0.0625 -0.0625  0.0625 -0.0625
                                         0.1875
                                                0.0625 0.0625
## 16 -0.1875 -0.0625 -0.0625 0.0625 -0.0625 0.0625
                                         0.0625
                                                0.1875 -0.0625
##
        10
              11
                    12
                           13
                                 14
                                       15
                                             16
## 1
     0.0625
           0.0625 -0.0625 0.0625 -0.0625 -0.0625 -0.1875
     0.1875 -0.0625
                0.0625 -0.0625 0.0625 -0.1875 -0.0625
## 3
    -0.0625
          -0.0625 -0.0625 -0.1875 0.1875 0.0625 0.0625 -0.0625
## 5
     -0.1875   0.0625   -0.0625   0.0625   -0.0625   0.1875
                                         0.0625
    -0.0625 -0.0625 0.0625 -0.0625
                             0.0625 0.0625
## 8
                                          0.1875
## 9
     0.1875   0.1875   0.0625   0.1875   0.0625   0.0625   -0.0625
## 10 0.3125 0.0625
                 0.1875 0.0625
                             0.1875 -0.0625
                                          0.0625
## 11
    0.0625 0.3125
                 0.1875  0.0625  -0.0625  0.1875
                                          0.0625
## 12 0.1875 0.1875
                0.3125 -0.0625
                             0.0625 0.0625
                                          0.1875
     0.0625
          0.0625 -0.0625 0.3125
                             0.1875 0.1875
                                          0.0625
     0.1875 -0.0625
                 0.0625
                       0.1875
                              0.3125
                                    0.0625
                                          0.1875
## 15 -0.0625
           0.1875
                 0.0625
                       0.1875
                              0.0625
                                    0.3125
                                          0.1875
## 16  0.0625  0.0625  0.1875  0.0625  0.1875  0.1875
                                         0.3125
(g)
```

```
yhat = H%*%y
yhat
```

```
##
           [,1]
## 1
      235.8875
## 2
      262.4875
```

```
## 3
      166.5875
## 4
      193.1875
      279.8125
## 6
      306.4125
## 7
      210.5125
## 8
      237.1125
## 9
      221.2625
## 10 247.8625
## 11 151.9625
## 12 178.5625
## 13 265.1875
## 14 291.7875
## 15 195.8875
## 16 222.4875
```

fitted(model)

```
##
                    2
                             3
                                      4
                                                5
                                                         6
                                                                   7
                                                                            8
          1
## 235.8875 262.4875 166.5875 193.1875 279.8125 306.4125 210.5125 237.1125
##
                   10
                                                        14
                            11
                                      12
                                               13
                                                                  15
## 221.2625 247.8625 151.9625 178.5625 265.1875 291.7875 195.8875 222.4875
all(round(yhat - fitted(model), 10)==0)
```

[1] TRUE

Yes, the yhats calculated from the matrix-vector computations match the yhats from the model created with lm().

(h)

```
resids = y-yhat
all(round(resids - residuals(model), 10)==0)
```

[1] TRUE

Yes, the residuals from the matrix-vector computation match the residuals from the model created with lm().

(i)

```
sd(resids)
```

[1] 22.56238

From the model the residual standard error: 26.35

The standard deviation of errors from our matrix-vector computation is slightly different from the model made with lm(). The difference in the estimates is reasonable considering there are only 16 observations.

(j)

```
xtxi <- solve(t(X) %*% X)</pre>
varHat = sum(((y-yhat)^2) / 11)
varB = xtxi * varHat
all(round(varB - vcov(model), 10)==0)
## [1] TRUE
Yes, our matrix computation of Var(\hat{\beta}) is the same as vcov(model).
sigHat = sqrt(varHat)
(seHatB3 = sqrt(diag(xtxi)) * sigHat)[4]
## temperaturehi
         13.17359
##
summary(model)$coefficients[4,2]
## [1] 13.17359
Yes, the matrix computation of the \hat{se}(\hat{\beta}_3) matches the \hat{se}(\hat{\beta}_3) from the lm() object.
(k)
(R2 = cor(y, yhat)^2)
```

```
(R2 = cor(y, yhat)^2)

## [,1]
## [1,] 0.8003646

summary(model)$r.squared
```

[1] 0.8003646

Yes, our computed value of R2 agrees with the output from our previously computed lm() model.