```
# Read data from florange.csv and input it into the dat vector.
dat <- read.csv("csv/florange.csv")</pre>
# Done to make the predict function work well.
x <- dat$acres
y <- dat$boxes
# Output the first 6 rows in dat.
head(dat)
##
        county boxes acres
## 1
       Brevard
                   51
## 2 Charlotte
                  821 13447
## 3
       Collier 2088 29351
## 4
        DeSoto 7688 66365
## 5
        Glades
                 368 5396
## 6
        Hardee 5306 43126
# Draw a scatterplot with x-axis as 'acres' and y-axis as 'boxes'.
plot(x, y)
                                                                                  0
                                                                           0
                                                                                     0
     2000 4000 6000
                                                            0
                                                                                0
                                             0
                            0
                     0
                           0
             0
                     10000
                                20000
                                           30000
                                                      40000
                                                                 50000
                                                                           60000
                                                 Χ
# Compute some common variables with common functions.
r <- cor(x, y)
xbar <- mean(x)
ybar <- mean(y)</pre>
cat("r:", r, "xbar:", xbar, "ybar:", ybar)
## r: 0.9635098 xbar: 16132.64 ybar: 1797.56
Therefore, r = 0.9635098, \bar{x} = 16132.64, and \bar{y} = 1797.56.
# Compute some common variables manually.
Sxx \leftarrow sum((x - xbar)^2)
Sxy \leftarrow sum((x - xbar) * (y - ybar))
cat("Sxx: ", Sxx, "Sxy: ", Sxy)
```

Sxx: 12450023404 Sxy: 1453128337

```
Therefore, S_{xx} = 12450023404 = 1.245 \times 10^{10} and S_{xy} = 1453128337 = 1.453 \times 10^{9}.
# R's lm function fits linear models
lm.1 < -lm(y ~x)
summary(lm.1)
##
## Call:
## lm(formula = y \sim x)
## Residuals:
         Min
                   1Q
                        Median
                                         3Q
## -2470.81
                -6.17
                           71.72 106.46 1677.32
## Coefficients:
                   Estimate Std. Error t value Pr(>|t|)
##
## (Intercept) -85.391989 186.178031 -0.459
## x
                   0.116717
                               0.006761 17.263 1.16e-14 ***
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
## Residual standard error: 754.4 on 23 degrees of freedom
## Multiple R-squared: 0.9284, Adjusted R-squared: 0.9252
## F-statistic: 298 on 1 and 23 DF, p-value: 1.164e-14
From the summary, we can see that \hat{\beta}_0 = -85.391989, \hat{\beta}_1 = 0.116717, Se(\hat{\beta}_1) = 0.006761, t = 17.263,
p-value = 1.64 \times 10^{-14}, and \hat{\sigma} = 754.4.
# Sum Squared Fitted Values
sum(lm.1$fitted.values ^ 2)
## [1] 250385207
# Sum Squared Residuals
sum(lm.1$residuals ^ 2)
## [1] 13089860
Therefore, SS(Res) = \sum_{i=1}^{n} e_i^2 = 13089860 = 1.31 \times 10^7.
# Manual calculation of sigma^2 estimate
sum(lm.1$residuals^2) / 23
## [1] 569124.3
Therefore, \hat{\sigma}^2 = 69124.3 = 5.7 \times 10^5.
# Manual calculation of sigma estimate
sqrt(sum(lm.1$residuals ^ 2) / 23)
## [1] 754.4033
Therefore, \hat{\sigma} = 754.4.
# t distribution values
qt(0.975, 23)
## [1] 2.068658
Therefore, c = 2.07.
```

```
# 95% confidence interval
confint(lm.1)
##
                      2.5 %
                                 97.5 %
## (Intercept) -470.5305905 299.7466119
                              0.1307034
## x
                  0.1027305
# 95% prediction interval with predicted boxes if we had 10000 acres
predict(lm.1, data.frame(x = 10000), interval = "prediction")
##
          fit
                    lwr
                             upr
## 1 1081.777 -512.0407 2675.595
```

Q: Is σ the same for all values of y?

A: It appears to not in the sense that the variance appears to be higher with respect to higher acres. Sigma will be smaller when there's less acres. Later, this will be testing equal variance or homoscedastic assumption. Later, when we talk about variable transformations we can consider taking the logarithm.

Q: Are the error terms plausibly independent? In other words, does knowing one e_i (residual) help predict e_j (another residual) for a different county?

A: There's diagnostics for checking this. However, intuitively there could be some common factors at play when two counties are geographically close.