Problem Set #2

Anaya Hall & Christian Miller

Part 1: Theory

(for practice only)

Part 2: Applied - Returns to Scale in Electricity Supply

First, load our OLS function created in Problem Set #1. We're including a built in t-test this time around.

```
ols <- function(data, y_data, X_data, intercept = T, HO = 0, two_tail = T, alpha = 0.05) {
  # Function setup ----
    # Require the 'dplyr' package
   require(dplyr)
    # Function to convert tibble, data.frame, or tbl_df to matrix
   to_matrix <- function(the_df, vars) {</pre>
      # Create a matrix from variables in var
      new_mat <- the_df %>%
        #Select the columns given in 'vars'
        select_(.dots = vars) %>%
        # Convert to matrix
        as.matrix()
      # Return 'new_mat'
      return(new_mat)
   }
  # Create dependent and independent variable matrices ----
    # y matrix
   y <- to_matrix (the_df = data, vars = y_data)
    # X matrix
   X <- to_matrix (the_df = data, vars = X_data)</pre>
      # If 'intercept' is TRUE, then add a column of ones
      if (intercept == T) {
      X \leftarrow cbind(1,X)
      colnames(X) <- c("intercept", X_data)</pre>
      }
  # Calculate b, y_hat, and residuals ----
   b <- solve(t(X) %*% X) %*% t(X) %*% y
    y_hat <- X %*% b
    e <- y - y_hat
  # Useful ----
   n <- nrow(X) # number of observations</pre>
   k <- ncol(X) # number of independent variables
   dof <- n - k # degrees of freedom
```

```
i <- rep(1,n) # column of ones for demeaning matrix
 A \leftarrow diag(i) - (1 / n) * i %*% t(i) # demeaning matrix
 y_star <- A %*% y # for SST
 X_star <- A %*% X # for SSM
 SST <- drop(t(y_star) %*% y_star)</pre>
 SSM <- drop(t(b) %*% t(X_star) %*% X_star %*% b)
 SSR <- drop(t(e) %*% e)
# Measures of fit and estimated variance ----
 R2uc <- drop((t(y_hat) %*% y_hat)/(t(y) %*% y)) # Uncentered R^2
 R2 <- 1 - SSR/SST # Uncentered R^2
 R2adj \leftarrow 1 - (n-1)/dof * (1 - R2) # Adjusted R^2
 AIC \leftarrow log(SSR/n) + 2*k/n # AIC
 SIC \leftarrow log(SSR/n) + k/n*log(n) # SIC
 s2 <- SSR/dof # s~2
# Measures of fit table ----
 mof_table_df <- data.frame(R2uc, R2, R2adj, SIC, AIC, SSR, s2)</pre>
 mof_table_col_names \leftarrow c("$R^2_\text{uc}$", "$R^2$",
                            \$R^2_\text{text{adj}},
                            "SIC", "AIC", "SSR", "$s^2$")
 mof_table <- mof_table_df %>% knitr::kable(
   row.names = F,
    col.names = mof_table_col_names,
    format.args = list(scientific = F, digits = 4),
    booktabs = T,
    escape = F
 )
# t-test----
 # Standard error
 se <- as.vector(sqrt(s2 * diag(solve(t(X) %*% X))))</pre>
  # Vector of _t_ statistics
 t_stats <- (b - H0) / se
  # Calculate the p-values
 if (two tail == T) {
 p_values <- pt(q = abs(t_stats), df = dof, lower.tail = F) * 2
    p_values <- pt(q = abs(t_stats), df = dof, lower.tail = F)</pre>
 }
  # Do we (fail to) reject?
 reject <- ifelse(p_values < alpha, reject <- "Reject", reject <- "Fail to Reject")
  # Nice table (data.frame) of results
 ttest_df <- data.frame(</pre>
    # The rows have the coef. names
    effect = rownames(b),
    # Estimated coefficients
    coef = as.vector(b) %>% round(3),
    # Standard errors
```

```
std_error = as.vector(se) %>% round(3),
      # t statistics
      t_stat = as.vector(t_stats) %>% round(3),
      # p-values
      p_value = as.vector(p_values) %>% round(4),
      # reject null?
      significance = as.character(reject)
   ttest_table <- ttest_df %>% knitr::kable(
      booktabs = T,
      format.args = list(scientific = F),
      escape = F
    )
  # Data frame for exporting for y, y_hat, X, and e vectors ----
    export_df <- data.frame(y, y_hat, e, X) %>% tbl_df()
    colnames(export_df) <- c("y","y_hat","e",colnames(X))</pre>
  # Return ----
    return(list(n=n, dof=dof, b=b, vars=export_df, R2uc=R2uc,R2=R2,
                R2adj=R2adj, AIC=AIC, SIC=SIC, s2=s2, SST=SST, SSR=SSR,
                mof_table=mof_table, ttest=ttest_table))
We'll also need a function to do an F-test for this Problem Set.
to_matrix <- function(the_df, vars) {</pre>
      # Create a matrix from variables in var
      new_mat <- the_df %>%
        #Select the columns given in 'vars'
        select (.dots = vars) %>%
        # Convert to matrix
        as.matrix()
      # Return 'new_mat'
```

```
# Return 'new_mat'
    return(new_mat)
}

# Joint test function (from Ed's notes). Could also write a more complex functions that takes R as
F_test <- function(data, y_var, X_vars) {
    # Turn data into matrices
    y <- to_matrix(data, y_var)
    X <- to_matrix(data, X_vars)
    # Add intercept
    X <- cbind(1, X)
    # Name the new column "intercept"
    colnames(X) <- c("intercept", X_vars)
    # Calculate n and k for degrees of freedom
    n <- nrow(X)</pre>
```

```
k \leftarrow ncol(X)
  # J is k-1
  J < - k - 1
  # Create the R matrix: bind a column of zeros
  \# onto a J-by-J identity matrix
  R <- cbind(0, diag(J))</pre>
  # Estimate coefficients
  b <- ols(data, y_var, X_vars)</pre>
  # Retrieve OLS residuals
  e <- b$vars$e
  # Retrieve s^2
  s2 <- b$s2 %<>% as.numeric()
  # Create the inner matrix R(X'X) \hat{(-1)}R'
  RXXR <- R \%*\% solve(t(X) \%*\% X) \%*\% t(R)
  # Calculate the F stat
  f_stat <- t(R %*% b$b) %*% solve(RXXR) %*% (R %*% b$b) / J / s2
  # Calculate the p-value;; why normal and not chi^2
  p_value <- pf(q = f_stat, df1 = J, df2 = n-k, lower.tail = F)</pre>
  # Create a data.frame of the f stat. and p-value
  results <- data.frame(
    f stat = f stat %>% as.vector(),
    p_value = p_value %>% as.vector())
  return(results)
}
```

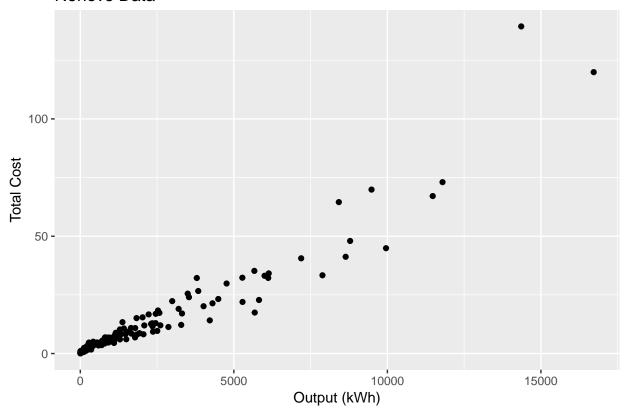
Question 1:

Read the data into R. Inspect it. Sort by size (Q (kwh)).

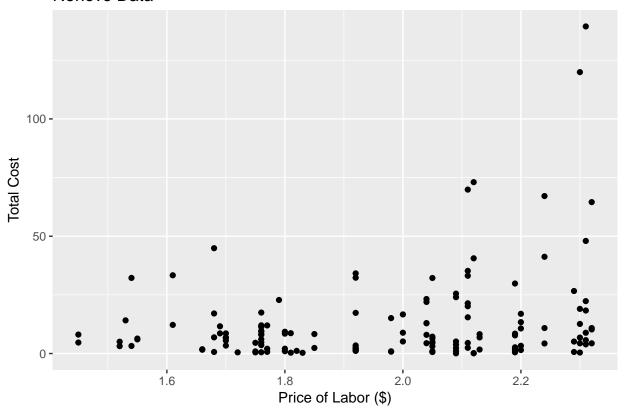
```
nerlove <- readxl::read_excel("nerlove.xls", col_names=T)
summary(nerlove)</pre>
```

```
##
          TC
                                            PL
                                                              PF
          : 0.082
                                  2
                                               1.450
                                                               :10.30
##
   Min.
                      Min.
                                      Min.
                                                        Min.
                      1st Qu.: 279
##
   1st Qu.: 2.382
                                      1st Qu.: 1.760
                                                        1st Qu.:21.30
   Median : 6.754
                      Median: 1109
                                      Median : 2.040
                                                        Median :26.90
##
         : 12.976
##
   Mean
                      Mean
                           : 2133
                                      Mean : 3.208
                                                        Mean
                                                               :26.18
##
   3rd Qu.: 14.132
                      3rd Qu.: 2507
                                      3rd Qu.: 2.190
                                                        3rd Qu.:32.20
           :139.422
##
   Max.
                      Max. :16719
                                      Max.
                                             :181.000
                                                               :42.80
                                                        Max.
          PΚ
##
##
   {	t Min.}
          :138.0
   1st Qu.:162.0
##
##
  Median :170.0
## Mean
           :174.5
## 3rd Qu.:183.0
##
   Max.
           :233.0
```

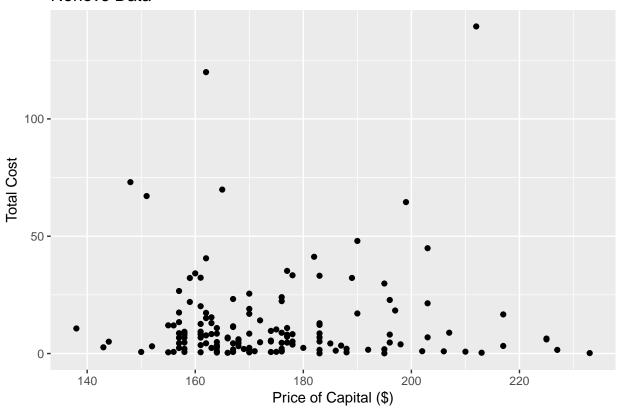
```
# Fix typo in 13th row (missing a decimal!)
# DO THIS MORE ELEGANTLY!
nerlove[13, "PL"] <- 1.81
# nerlove %>%
   filter(PL > 100) %>%
     mutate(PL = PL/100)
nerlove %>% arrange(Q)
## # A tibble: 145 x 5
##
          TC
                 Q
                     PL
                           PF
                                 PK
##
       <dbl> <dbl> <dbl> <dbl> <dbl> <dbl>
##
   1 0.0820
               2. 2.09
                         17.9 183.
               3. 2.05 35.1 174.
##
   2 0.661
  3 0.990
               4.
                   2.05 35.1 171.
##
   4 0.315
               4. 1.83 32.2 166.
##
##
   5 0.197
               5.
                   2.12 28.6 233.
   6 0.0980
               9. 2.12 28.6 195.
##
##
  7 0.949
              11.
                   1.98 35.5 206.
##
  8 0.675
              13. 2.05 35.1 150.
## 9 0.525
              13.
                   2.19 29.1 155.
## 10 0.501
               22. 1.72 15.0 188.
## # ... with 135 more rows
Plot the series and make sure your data are read in correctly.
ggplot(nerlove, aes(x=Q, y=TC)) +
  geom_point() +
  labs(title="Nerlove Data", x="Output (kWh)", y="Total Cost")
```



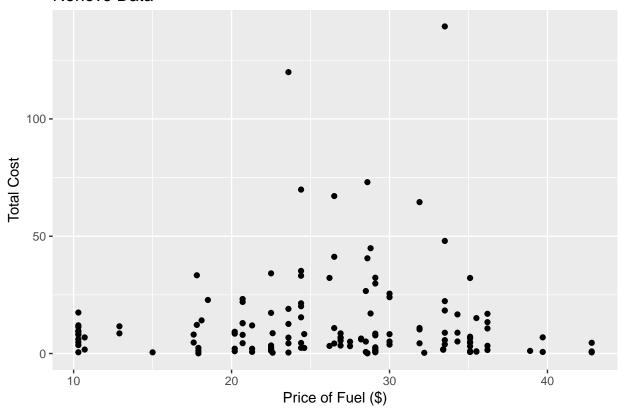
```
ggplot(nerlove, aes(x=PL, y=TC)) +
geom_point() +
labs(title="Nerlove Data", x="Price of Labor ($)", y="Total Cost")
```



```
ggplot(nerlove, aes(x=PK, y=TC)) +
  geom_point() +
  labs(title="Nerlove Data", x="Price of Capital ($)", y="Total Cost")
```



```
ggplot(nerlove, aes(x=PF, y=TC)) +
geom_point() +
labs(title="Nerlove Data", x="Price of Fuel ($)", y="Total Cost")
```



Question 2:

Replicate regression I (page 176) in the paper.

Regression I:

$$log(TC) - log(P_F) = \beta_0 + \beta_1 Q + \beta_2 \Big(log(P_L) - log(P_F)\Big) + \beta_3 \Big(log(P_K) - log(P_F)\Big)$$

Equivalent to:

$$log(\frac{TC}{P_F}) = \beta_0 + \beta_1 Q + \beta_2 log(\frac{P_L}{P_F}) + \beta_3 log(\frac{P_K}{P_F})$$

Where:

TC = total production cost,

 P_L = wage rate,

 P_K = "price" of capital,

 P_F = price of fuel,

Q = output (measured in kWh)

In generalized Cobb-Douglas form:

$$\beta_1 = \frac{1}{r}$$

$$\beta_2 = \frac{a_L}{r},$$

$$\beta_3 = \frac{a_K}{r}$$

Prepare variables for Regression I.

```
# Create log variables
nerlove %<>% mutate(
  TClog = log(TC),
  Qlog = log(Q),
  PLlog = log(PL),
  PKlog = log(PK),
  PFlog = log(PF)
)
# Create PF scaled variables
nerlove %<>% mutate(
  TCscaled = TClog - PFlog,
  PLscaled = PLlog - PFlog,
  PKscaled = PKlog - PFlog
Variable names:
log(\frac{TC}{P_F}) = "TCscaled"
log(\frac{P_L}{P_E}) = "PLscaled"
log(\frac{P_K}{P_F}) = "PKscaled"
# Regression I:
# dep var = (log costs - log fuel price) = TCscaled
reg_I <- ols(data = nerlove,y_data = "TCscaled",</pre>
              X_data = c("Qlog","PLscaled","PKscaled"),
```

effect	coef	std_error	t_stat	p_value	significance
intercept	-2.037	0.384	-5.301	0.0000	Reject
Qlog	0.721	0.017	41.334	0.0000	Reject
PLscaled	0.593	0.205	2.898	0.0044	Reject
PKscaled	-0.007	0.191	-0.039	0.9692	Fail to Reject

reg_I\$mof_table

reg_I\$ttest

R_{uc}^2	R^2	$R_{\rm adj}^2$	SIC	AIC	SSR	s^2
0.966	0.9316	0.9301	-3.433	-3.515	4.082	0.02895

Coefficients are pretty close to those in the paper. R^2 matches!

intercept = T, HO = 0, alpha = 0.05)

Question 3:

Conduct the hypothesis test using constant returns to scale ($\beta_1 = 1$) as your null hypothesis.

effect	coef	std_error	t_stat	p_value	significance
intercept	-2.037	0.384	-7.903	0.0000	Reject
Qlog	0.721	0.017	-16.020	0.0000	Reject
PLscaled	0.593	0.205	-1.990	0.0485	Reject
PKscaled	-0.007	0.191	-5.282	0.0000	Reject

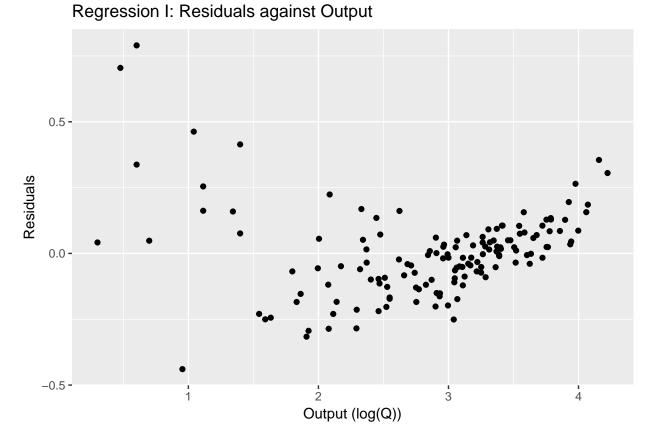
What is the p- value associated with you test statistic? What is your point estimate of returns to scale? Constant? Increasing? Decreasing?

The p-value is 0.000. The point estimate of returns to scale is $\frac{1}{\beta} = r = 1.3875639$, hence returns to scale is increasing.

Question 4:

Plot residuals against output.

```
ggplot(reg_I$vars, aes(y=e, x=Qlog)) + geom_point() + labs(title="Regression I: Residuals against
```



What do you notice? What does this potentially tell you from an economic perspective?

Evidence of heteroskedasticity: residuals seem to track the log output through parabola. We may want to rethink our specification!

Compute the correlation coefficient of the residuals with output for the entire sample? What does this tell you?

```
# R = cov(xy)/var(x)var(y)
R_I <- (cov(x=reg_I$vars$e, y=reg_I$vars$Qlog)/(var(reg_I$vars$e)*var(reg_I$vars$Qlog))) %>% sign:
```

The correlation coefficient is extremely small: 7.01e-14. This tells us that, conversely, there is very little correlation between the output of the firm and the undetermined part of the model. This supports our strict exogeneity assumption.

Question 5:

Divide your sample into 5 subgroups of 29 firms each according to the level of output. Estimate the regression model again for each group separately.

effect	coef	std_error	t_stat	p_value	significance
intercept	-1.452	1.366	-1.795	0.0848	Fail to Reject
Qlog	0.400	0.084	-7.101	0.0000	Reject
PLscaled	0.615	0.729	-0.528	0.6024	Fail to Reject
PKscaled	-0.081	0.706	-1.531	0.1384	Fail to Reject

effect	coef	std_error	t_stat	p_value	significance
intercept	-2.818	0.614	-6.222	0.0000	Reject
Qlog	0.658	0.116	-2.939	0.0070	Reject
PLscaled	0.094	0.274	-3.304	0.0029	Reject
PKscaled	0.378	0.277	-2.250	0.0335	Reject

effect	coef	$\operatorname{std}\operatorname{_error}$	t_stat	p_value	significance
intercept	-3.185	0.734	-5.705	0.0000	Reject
Qlog	0.938	0.198	-0.312	0.7578	Fail to Reject
PLscaled	0.402	0.199	-2.997	0.0061	Reject
PKscaled	0.250	0.187	-4.010	0.0005	Reject

effect	coef	std_error	t_stat	p_value	significance
intercept	-2.843	0.506	-7.596	0.0000	Reject
Qlog	0.912	0.107	-0.818	0.4210	Fail to Reject
PLscaled	0.507	0.187	-2.630	0.0144	Reject
PKscaled	0.093	0.164	-5.525	0.0000	Reject

effect	coef	std_error	t_stat	p_value	significance
intercept	-2.916	0.454	-8.618	0.0000	Reject
Qlog	1.044	0.065	0.683	0.5008	Fail to Reject
PLscaled	0.603	0.197	-2.014	0.0549	Fail to Reject
PKscaled	-0.289	0.175	-7.374	0.0000	Reject

Can you replicate Equations IIIA - IIIE? Calculate the point estimates for returns to scale for each sample. Is there a pattern relating to size of output?

```
reg_III_table <- reg_III_df %>% knitr::kable(
    col.names = c("Regression", "Returns to Scale"),
    format.args = list(scientific = F, digits = 4),
    booktabs = T,
    escape = F
)
reg_III_table
```

Regression	Returns to Scale
IIIA	2.4982
IIIB	1.5194
IIIC	1.0658
IIID	1.0964
IIIE	0.9575

Coefficients roughly match the results of the paper! As the size (output) of the firms get larger they see decreasing, increasing returns to scale. The largest bucket even sees decreasing returns to scale.

Question 6:

Create "dummy variables" for each industry. Interact them with the output variable to create five "slope coefficients".

```
# create group categorical var
for (i in 1:5) {
  d[[i]] %<>% mutate(gvar=i)
}
# unsplit
df <- rbind(d[[1]], d[[2]], d[[3]], d[[4]], d[[5]])
# create dummies
df %<>%
   mutate(
      g1 = ifelse(gvar==1, 1, 0),
      g2 = ifelse(gvar==2, 1, 0),
      g3 = ifelse(gvar==3, 1, 0),
      g4 = ifelse(gvar==4, 1, 0),
      g5 = ifelse(gvar==5, 1, 0)) %>% select(-gvar)
# interact with output variable
df %<>%
   mutate(
      1Q_A = Qlog*g1,
     1Q_B = Qlog*g2,
     1Q_C = Qlog*g3,
```

```
1Q_D = Qlog*g4,
1Q_E = Qlog*g5)
```

Run a model, letting the intercept and slope coefficient on output differ across plants, but let the remainder of the coefficients be pooled across plants.

Can also omit one of the dummie vars (g) and include an intercept >> get the same coefs reg_IV\$ttest

effect	coef	std_error	t_stat	p_value	significance
lQ_A	0.397	0.043	9.214	0.0000	Reject
lQ_B	0.648	0.147	4.402	0.0000	Reject
lQ_C	0.885	0.297	2.976	0.0035	Reject
lQ_D	0.909	0.274	3.321	0.0012	Reject
lQ_E	1.063	0.131	8.091	0.0000	Reject
PLscaled	0.426	0.163	2.608	0.0101	Reject
PKscaled	0.104	0.152	0.681	0.4967	Fail to Reject
g1	-1.815	0.305	-5.952	0.0000	Reject
g2	-2.194	0.489	-4.491	0.0000	Reject
g3	-2.879	0.972	-2.963	0.0036	Reject
g4	-2.922	0.966	-3.024	0.0030	Reject
g5	-3.511	0.599	-5.857	0.0000	Reject

reg_IV\$mof_table

R_{uc}^2	R^2	R_{adj}^2	SIC	AIC	SSR	s^2
0.9802	0.9602	0.9569	-3.701	-3.947	2.372	0.01784

```
booktabs = T,
  escape = F
)

reg_IV_table
```

Regression	Returns to Scale
IVA	2.520
IVB	1.543
IVC	1.130
IVD	1.100
IVE	0.941

Are there any noticeable changes in returns to scale from the previous part?

We got roughly the same point estimates on returns to scale as the previous part! There is a slight bit of variation which is from the pooled labor and capital price effect. I would imagine we'd get even closer to the previous result if we interacted our industry dummy variable with each of them.

Question 7:

1

11.9797 8.901198e-13

Conduct a statistical test comparing the first model you estimate to the last model you estimated. (Hint: Is one model a restricted version of the other?). Would separate t-test have given you the same results?

Regression I is the restricted model, Regression IV is unrestricted model. This being, Regression I says that all five industries all have the same slope on output and all have the same intercept, thus placing restrictions.

```
F = \frac{(SSR_R - SSR_U)/J}{GSR_U}
      SSR_U/(n-k)
# Calculate F_stat using SSR approach
ssr_u <- reg_IV$SSR
dof <- reg_IV$dof</pre>
ssr_r <- reg_I$SSR
j <- 8
f_statistic <- ((ssr_r - ssr_u) / j) / (ssr_u / dof)
# Calculate the p-value
p_value \leftarrow pf(q = f_statistic, df1 = j, df2 = dof, lower.tail = F)
# Create a data.frame of the f stat. and p-value
f_test <- data.frame(</pre>
  f_statistic = f_statistic %>% as.vector(),
  p_value = p_value %>% as.vector())
f_{test}
##
     f_statistic
                        p_value
```

```
F_test(df, "TCscaled", c("lQ_A", "lQ_B", "lQ_C", "lQ_D", "lQ_E", "PLscaled", "PKscaled", "g1", "g2"
## f stat p value
```

```
## f_stat p_value
## 1 291.9844 2.238126e-87
```

As you can see, we got an incredibly low p_value between the first and last model, menaing the differentiated slope on output and intercept coefficients are joint, statistically significant. This SSR approach is different than just doing an F-Test on the last regression because it is saying, are all the variables (including the pooled price on labor and capital) jointly, statistically significant from just the mean of the scaled log of total cost. I don't even want to begin thinking about running a myriad of t-test on the 8 individual restrictions of our first model. Theory tells us we are likely to get different results because the F-test correctly ajusts the joint confidence intervals while seperate t-test would be a ridgid overlapping of confidence intervals that may(not) correctly approximate the F-test.

Question 8:

To see whether returns to scale declined with output, Nerlove tested a nonlinear specification by including $\ln(y)^2$ as a regressor. Conduct a statistical test you feel is appropriate to test this hypothesis.

It follows that returns to scale (r) = $\frac{1}{\alpha + \beta \ln(y)}$, where α is the coefficient on $\ln(y)$ and β is the coefficient on $\ln(y)^2$ in the below regression.

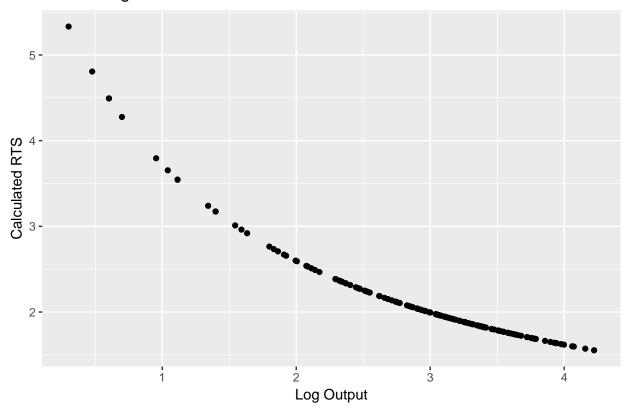
effect	coef	std_error	t_stat	p_value	significance
intercept	-1.635	0.305	-5.365	0.0000	Reject
Qlog	0.153	0.062	2.466	0.0149	Reject
$Qlog_sq$	0.116	0.012	9.418	0.0000	Reject
PLscaled	0.481	0.161	2.984	0.0034	Reject
PKscaled	0.074	0.150	0.494	0.6218	Fail to Reject

We're going to save these results and first plot our calculated returns to scale variable against log output to see if there is a trend.

```
df %<>% mutate(rts_nl = 1/(reg_nl$b[2] + reg_nl$b[3]*df$Qlog))

ggplot(df, aes(x=Qlog, y=rts_nl)) +
   geom_point() +
   labs(title="Decreasing Returns to Scale?", x="Log Output", y="Calculated RTS")
```

Decreasing Returns to Scale?



It would seem that our Returns to Scale Decreases as output increases. Whether this is statistically significant is the question. We'll run a regression with these two variables and our pooled labor and capital price in a t-test. Dut to the non-linear decline of returns to scale, we'll also include the squared log output in the regression.

effect	coef	std_error	t_stat	p_value	significance
intercept	5.484	0.151	36.224	0.0000	Reject
Qlog	-1.842	0.031	-59.947	0.0000	Reject
$Qlog_sq$	0.227	0.006	36.941	0.0000	Reject
PLscaled	0.042	0.080	0.524	0.6010	Fail to Reject
PKscaled	0.003	0.075	0.034	0.9728	Fail to Reject

Here, it would seem that for every one percent increase of output for firms, we see a -0.018423 decrease in the firms returns to scale. The results is statistically significant, though, we did not take the robust variance of the original paper throughout, so I don't know how much we can rely on that.