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.

```
# 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) {</pre>
  # Turn data into matrices
  y <- to_matrix(data, y_var)</pre>
  X <- to_matrix(data, X_vars)</pre>
  # Add intercept
  X \leftarrow cbind(1, X)
  # Name the new column "intercept"
  colnames(X) <- c("intercept", X_vars)</pre>
  # Calculate n and k for degrees of freedom
  n \leftarrow nrow(X)
  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 <- b_ols(data, y_var, X_vars)</pre>
  # Calculate OLS residuals
  e <- y - X %*% b
  # Calculate s^2
```

```
s2 <- (t(e) %*% e) / (n-k)
# Force s2 to numeric
s2 %<>% as.numeric()

# Create the inner matrix R(X'X)^(-1)R'
RXXR <- R %*% solve(t(X) %*% X) %*% t(R)
# Calculate the F stat
f_stat <- t(R %*% b) %*% solve(RXXR) %*% (R %*% b) / J / s2
# Calculate the p-value
p_value <- pf(q = f_stat, df1 = J, df2 = n-k, lower.tail = F)
# 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)</pre>

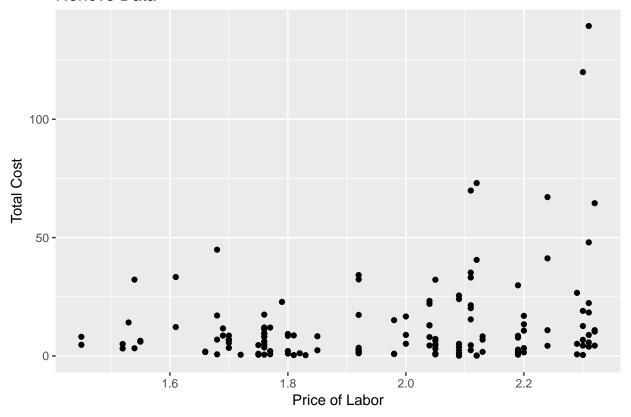
```
summary(nerlove)
         TC
                                           PL
                                                             PF
##
                           Q
   Min. : 0.082
                                          : 1.450
                           :
                                 2
                                                             :10.30
##
                     Min.
                                     Min.
                                                       Min.
   1st Qu.: 2.382
                     1st Qu.: 279
                                     1st Qu.: 1.760
                                                       1st Qu.:21.30
  Median : 6.754
                     Median: 1109
                                                       Median :26.90
                                     Median : 2.040
##
  Mean : 12.976
                     Mean : 2133
                                     Mean : 3.208
                                                       Mean
                                                             :26.18
##
   3rd Qu.: 14.132
                     3rd Qu.: 2507
                                     3rd Qu.: 2.190
                                                       3rd Qu.:32.20
##
   Max.
          :139.422
                     Max.
                           :16719
                                     Max.
                                            :181.000
                                                       Max.
                                                              :42.80
##
         PK
## 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
                        PL
##
           TC
                  Q
                               PF
                                     PK
##
       <dbl> <dbl> <dbl> <dbl> <dbl> <dbl>
                      2.09
                 2.
                            17.9
    1 0.0820
                                   183.
##
                      2.05
##
    2 0.661
                            35.1
                                   174.
                      2.05
                            35.1
##
    3 0.990
                 4.
                                   171.
##
    4 0.315
                 4.
                      1.83
                            32.2
                                   166.
    5 0.197
                      2.12
                            28.6
                                   233.
##
                 5.
##
    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
                      2.19
                13.
                            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=PL, y=TC)) +
  geom_point() +
  labs(title="Nerlove Data", x="Price of Labor", y="Total Cost")
```

Nerlove Data



plot rest of independent vars too.

Question 2:

Replicate regression I (page 176) in the paper.

```
Regression I:
```

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:
$$log(\frac{TC}{P_F}) = \beta_0 + \beta_1 Q + \beta_2 \Big(log(\frac{P_L}{P_F})\Big) + \beta_3 \Big(log(\frac{P_K}{P_F})\Big)$$
 Where:
$$TC = \text{total production cost},$$

$$P_L = \text{wage rate},$$

$$P_K = \text{"price" of capital},$$

$$P_F = \text{price of fuel},$$

$$Q = \text{output (measured in kWh)}$$
 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.
$$\# \text{Create log variables}$$

$$\text{nerlove } \text{$\ensuremath{$\ensur$$

Variable names:

)

PLscaled = PLlog - PFlog, PKscaled = PKlog - PFlog

```
log(\frac{TC}{P_F}) = "TC
scaled"
log(\frac{P_L}{P_F}) = "PL<br/>scaled"
log(\frac{P_K}{P_F}) = "PK
scaled"
# 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"),
                intercept = T, HO = 0, alpha = 0.05)
```

reg_I\$ttest

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

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. \mathbb{R}^2 matches!

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

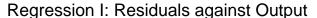
What is the p- value associated with you test statistic? What is your point estimate of returns to scale? Constant? Increasing? Decreasing?

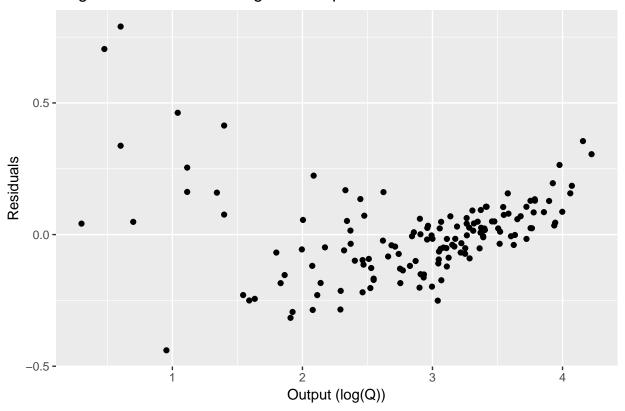
The p-value is 0.000. We estimate that returns to scale is......

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. 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?

Correlation coefficient of residuals with output: what does this mean?

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

effect	coef	std_error	t_stat	p_value	significance
PLscaled	0.615	0.729	-0.528	0.6024	Fail to Reject
PKscaled	-0.081	0.706	-1.531	0.1384	Fail to Reject

Regression IIIB

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

Regression IIIC

effect	coef	std_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

Regression IIID

reg_IIID\$ttest

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

```
# Regression IIIE
```

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

```
## May want to clean this up and run as for loop???
```

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?

[need to answer this still!]

Coefficients roughly match the results of the paper!

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))
df %<>% select(-gvar)
# interact with output variable
df %<>%
   mutate(
      1Q_1 = Qlog*g1,
      1Q_2 = Qlog*g2,
      1Q_3 = Qlog*g3,
      1Q_4 = Qlog*g4,
      1Q_5 = 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.

reg_IV\$ttest

effect	coef	std_error	t_stat	p_value	significance
lQ_1	0.397	0.043	-14.002	0.0000	Reject
lQ_2	0.648	0.147	-2.389	0.0183	Reject
lQ_3	0.885	0.297	-0.388	0.6989	Fail to Reject
lQ_4	0.909	0.274	-0.333	0.7393	Fail to Reject
lQ_5	1.063	0.131	0.478	0.6336	Fail to Reject
g1	-1.815	0.305	-9.231	0.0000	Reject
g2	-2.194	0.489	-6.538	0.0000	Reject
g3	-2.879	0.972	-3.993	0.0001	Reject
g4	-2.922	0.966	-4.059	0.0001	Reject
g5	-3.511	0.599	-7.525	0.0000	Reject
PLscaled	0.426	0.163	-3.520	0.0006	Reject
PKscaled	0.104	0.152	-5.888	0.0000	Reject

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

Question 7:

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?

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.