The Application of California School

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1 Introduction

This tutorial is to show how to estimate a multiple regression model and perform linear hypothesis testing. The application is about the test scores of the California school districts. We will use R to replicate the multiple regression with this data set in Chapter 6, and the hypothesis tests in Chapter 7, Stock and Watson. (2011).

Before running all R codes, we may first load all the packages.

library(AER)
library(foreign)
library(stargazer)

2 Reading the data and basic summary statistics

Let's first read the data into R and show some basic statistics.

Read the STATA file

Since the data is stored as an STATA file with the extension ".dta", we read the data using read.dta() in the library of foreign.

```
setwd("/Users/ztian/OneDrive/teaching/workshop/intro_org_RR/example")
classdata <- read.dta("./data/caschool.dta")</pre>
```

Summary

Upon reading the data, we often use summary() to see some basic statistics. Here we are not going to show summary statistics of all variables in the data set for the purpose of saving space, but only to select several variables of interest in Chapters 6 and 7, including test scores, testscr, student-teacher ratio, str, percentage of English learners, el_pct, expenditure per pupil, expn_stu, and percentage of students qualifying for free lunch, mean_pct.

```
summary(classdata[c("testscr", "str", "el_pct", "expn_stu", "meal_pct")])
```

testscr	str	${ m el_pct}$	$expn_stu$	$\mathrm{meal_pct}$
Min. :605.5	Min. :14.00	Min.: 0.000	Min. :3926	Min.: 0.00
1st Qu.:640.0	1st Qu.:18.58	1st Qu.: 1.941	1st Qu.:4906	1st Qu.: 23.28
Median $:654.5$	Median :19.72	Median: 8.778	Median: 5215	Median: 41.75
Mean : 654.2	Mean : 19.64	Mean $:15.768$	Mean :5312	Mean: 44.71
3rd Qu.:666.7	3rd Qu.:20.87	3rd Qu.:22.970	3rd Qu.:5601	3rd Qu.: 66.86
Max. :706.8	Max. $:25.80$	Max. :85.540	Max. :7712	Max. :100.00

3 Plots

par(oldpar)

Create a matrix of scatterplots using plot

We can create several scatterplots displayed in one graph with a matrix form.

```
oldpar <- par(mfrow = c(2, 3))
plot(classdata$str, classdata$testscr, col = "red",
     main = "Student-teacher ratio vs test scores",
     xlab = "Student-teacher ratio", ylab = "Test scores")
plot(classdata$el_pct, classdata$testscr, col = "blue",
     main = "English learners vs test scores",
     xlab = "Percentage of English learners",
     ylab = "Test scores")
plot(classdata$expn_stu, classdata$testscr, col = "green3",
     main = "Expenditure per pupil vs test scores",
     xlab = "Expenditure per pupil",
     ylab = "Test scores")
plot(classdata$meal_pct, classdata$testscr, col = "maroon",
     main = "Free lunch vs test scores",
     xlab = "Percentage of students with free lunch",
     ylab = "Test scores")
plot(classdata$calw_pct, classdata$testscr, col = "darkorange1",
    main = "Public assistance vs test scores",
    xlab = "Percentage of students in public assistance",
    ylab = "Test scores")
```

We can see that the codes above have some parts that are repeated in each plotting command. So these repetitive work can be concisely written in for a loop. The basic syntax of a for loop is for (var in seq) expr.

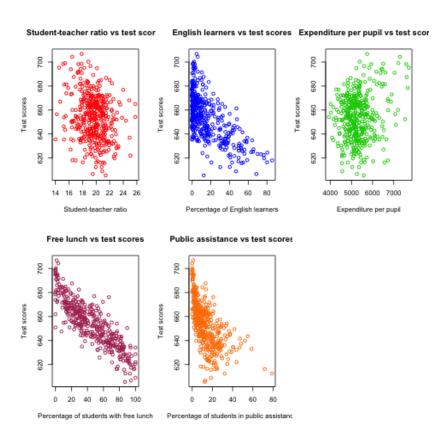


Figure 1: The scatterplots between several variables and test scores

```
xvars <- c("str", "el_pct", "expn_stu", "meal_pct", "calw_pct")</pre>
yvars <- c("testscr")</pre>
xlabels <- c("Student-teacher ratio", "Percentage of English learners",
              "Expenditure per pupil", "Percentage of students with free lunch",
              "Percentage of students in the public assistant program")
ylabels <- "Test scores"
titles <- c("student-teacher ratio vs test scores",
            "English learners vs test scores",
            "Expenditure per pupil vs test scores",
            "Free lunch vs test scores vs test scores",
             "public assistance vs test scores")
colors <- c("red", "green3", "blue", "maroon", "darkorange1")</pre>
op \leftarrow par(mfrow = c(2, 3))
for (i in seq(along=xvars)) {
    fm <- formula(paste(yvars, "~", xvars[i]))</pre>
    plot(fm, data = classdata, col = colors[i], main = titles[i],
         xlab = xlabels[i], ylab = ylabels)
}
par(op)
```

4 Estimation

Let us first replicate the regression results in Equation (7.19). The unit of the expenditure per pupil is dollars in the data set but it is in thousand dollars in regression. So we need to convert the unit in the data set by dividing expn_stu by 1000, which is done directly in the formula.

The OLS estimation

```
model.76 <- testscr ~ str + I(expn_stu / 1000) + el_pct</pre>
```

Notice the function I() in the formula. The arithmetic operations, +, *, :, /, and $\hat{}$, have special meanings in R's formula. Using the function I() protects the original arithmetic meanings of these operations from being interpreted in terms of a formula.

The regression estimation can be done with lm() and use summary() afterwards.

```
res.model.76 <- lm(model.76, data = classdata)
summary(res.model.76)
Call:</pre>
```

```
lm(formula = model.76, data = classdata)
```

Residuals:

```
Min 1Q Median 3Q Max -51.340 -10.111 0.293 10.318 43.181
```

Coefficients:

```
Estimate Std. Error t value Pr(>|t|)
(Intercept)
                649.57795
                            15.20572 42.719 < 2e-16 ***
                  -0.28640
                                      -0.596 0.55149
str
                             0.48052
I(expn_stu/1000)
                  3.86790
                             1.41212
                                       2.739 0.00643 **
el_pct
                 -0.65602
                             0.03911 -16.776 < 2e-16 ***
Signif. codes: 0 '*** 0.001 '** 0.01 '* 0.05 '.' 0.1 ' 1
```

Residual standard error: 14.35 on 416 degrees of freedom Multiple R-squared: 0.4366, Adjusted R-squared: 0.4325 F-statistic: 107.5 on 3 and 416 DF, p-value: < 2.2e-16

We can extract some components in the reported results. Use coef() to get the coefficient estimates, resid() to get the residuals, and fitted() or predict() to get the fitted values. Alternatively, we can think the lm and summary.lm objects returned by lm() and summary() are the list object so that we can use the "\$" operator to get each component of the lists. Below are some examples of extracting regression results.

```
# get some components of the results
bhat <- coef(res.model.76)
rsq <- summary(res.model.76)$r.squared
adj.rsq <- summary(res.model.76)$adj.r.squared</pre>
```

• Interpretation of the results

As for the coefficients

- 1. The intercept is 649.5779, which is significant at 1% significance level. It does not have real meaning in this application, just determining the position of the sample regression line crossing the vertical axis.
- 2. The coefficient on str is -0.2864, implying that increasing one more student per teacher would decrease test scores by 0.2864 unit. However, this estimated coefficient is not significant at the 10% level.
- 3. The coefficient on expenditure per pupil is 3.8679, significantly positive at the 5% level, implying that an increase in expenditure per pupil by one thousand dollars lead to an increase in test scores by 3.8679 unit.
- 4. The coefficient on the percentage of English learners is -0.656, significantly negative at the 1% level, implying that an increase in the percentage of English learners by one percent results in a decrease of test scores by 0.656.

Besides, the R^2 and \bar{R}^2 are 0.4366 and 0.4325, respectively. Overall, the model explains about 43% variation of test scores with the included explanatory variables, which is modest in the sense that a little more than half of the variation of test scores is not accounted for in the model.

The heteroskedasticity-consistent covariance matrix

Note that standard errors and t statistics reported by summary() are the homoskedasticity-only s.e. and t's. The heteroskedasticity-robust covariance matrix can be obtained by vcovHC() in the package of sandwich.

```
htvarm <- vcovHC(res.model.76, type = "HC1")
```

5 Hypothesis tests

Testing a single coefficient

Running summary(res.model.76) can give you t-statistics for all coefficients. However, as noted above, these t-statistics are the homoskedasticity-only t-statistics. We should use the heteroskedasticity-robust ones.

```
# homoskedasticity-only
coeftest(res.model.76)

# heteroskedasticity-robust, t distribution
cftest.t <- coeftest(res.model.76, vcov = htvarm)
cftest.t

# heteroskedasticity-robust, normal distribution
cftest.n <- coeftest(res.model.76, vcov = htvarm, df = Inf)
cftest.n

(Intercept)
   649.5779

   str
-0.2864

   str
0.2864</pre>
```

```
I(expn_stu/1000)
         3.8679
I(expn_stu/1000)
         3.8679
el_pct
-0.656
el_pct
0.656
[1] 0.4366
[1] 0.4325
t test of coefficients:
                  Estimate Std. Error t value Pr(>|t|)
(Intercept)
                649.577947 15.205719 42.7193 < 2.2e-16 ***
                 -0.286399
                            0.480523 -0.5960 0.551489
str
I(expn_stu/1000)
                  3.867902
                             1.412122
                                        2.7391 0.006426 **
                             0.039106 -16.7755 < 2.2e-16 ***
el_pct
                 -0.656023
Signif. codes: 0 '*** 0.001 '** 0.01 '* 0.05 '.' 0.1 ' 1
t test of coefficients:
                  Estimate Std. Error t value Pr(>|t|)
(Intercept)
                649.577947 15.458343 42.0212 < 2e-16 ***
                 -0.286399
                            0.482073 -0.5941 0.55277
str
I(expn_stu/1000)
                  3.867902 1.580722
                                        2.4469 0.01482 *
                 -0.656023
                             0.031784 -20.6397 < 2e-16 ***
el_pct
Signif. codes: 0 '*** 0.001 '** 0.01 '* 0.05 '.' 0.1 ' 1
z test of coefficients:
                  Estimate Std. Error z value Pr(>|z|)
(Intercept)
                649.577947 15.458343 42.0212 < 2e-16 ***
                 -0.286399 0.482073 -0.5941 0.55245
I(expn_stu/1000)
                  3.867902
                             1.580722
                                        2.4469 0.01441 *
                 -0.656023
                             0.031784 -20.6397 < 2e-16 ***
el_pct
Signif. codes: 0 '*** 0.001 '** 0.01 '* 0.05 '.' 0.1 ' 1
```

We can see from the results above that

1. whether we use the homoskedasticity-only or heteroskedasticity-robust variance matrices does not affect the coefficient estimates because the calculation of these estimates does not involve the variance matrices.

- 2. Using the homoskedasticity-only or heteroskedasticity-robust variance matrices yields different standard errors and t-statistics. Even though the homoskedasticity-only standard errors of student-teacher ratios seems smaller than the heteroskedasticity-robust ones, we cannot say that the estimates with the homoskedasticity-only standard errors are more efficient or precise because we are using a wrong variance matrix in this case.
- 3. The p-values from t distribution and standard normal distribution are slightly different, given the corresponding t-statistics are identical in the two cases.

Testing joint hypotheses

• Zero restrictions Let's first test the joint zero restrictions.

test1 <- linearHypothesis(res.model.76,

$$H_0: \beta_1 = 0, \beta_2 = 0 \text{ vs. } H_1: \beta_1 \neq 0 \text{ or } \beta_2 \neq 0$$

We can use the function linear Hypothesis () to test this and any linear hypotheses.

```
c("str = 0", "I(expn_stu/1000) = 0"),
    vcov = htvarm, test = "F")
test1
test1.F <- test1$F[2]
test1.p <- test1$"Pr(>F)"[2]
Linear hypothesis test
Hypothesis:
str = 0
I(expn_stu/1000) = 0
Model 1: restricted model
Model 2: testscr ~ str + I(expn_stu/1000) + el_pct
Note: Coefficient covariance matrix supplied.
  Res.Df Df
                 F
                     Pr(>F)
     418
     416 2 5.4337 0.004682 **
               0 '*** 0.001 '** 0.01 '* 0.05 '. ' 0.1 ' ' 1
Signif. codes:
```

The F-statistic is 5.4337 with the p-value as 0.0047, which is less than 1%. Therefore, we can reject the null hypothesis at the 1% level.

Note that the F-statistic is computed with the heteroskedasticity-robust variance matrix and tested against a F distribution of (2, 416) degree of freedom.

• linear restrictions Let's test the following restriction,

$$H_0: \beta_1 + \beta_2 = 0, H_1: \beta_1 + \beta_2 \neq 0$$

We still use linearHypothesis(). But this time we use the argument white.adjust for which we specify "hc1" and test against a Chi-squared distribution with one degree of freedom. Therefore, what we get is a Wald statistic.

```
# b1 + b2 = 0
test2 <- linearHypothesis(res.model.76,
    c("str + I(expn_stu/1000) = 0"),
    white.adjust = "hc1", test = "Chisq")
test2
test2.x <- test2$Chisq[2]
test2.p <- test2$"Pr(>Chisq)"[2]
[1] 5.4337
[1] 0.0047
Linear hypothesis test
Hypothesis:
str + I(expn_stu/1000) = 0
Model 1: restricted model
Model 2: testscr ~ str + I(expn_stu/1000) + el_pct
Note: Coefficient covariance matrix supplied.
  Res.Df Df Chisq Pr(>Chisq)
     417
     416 1 3.6319
                      0.05668 .
2
              0 '*** 0.001 '** 0.01 '* 0.05 '. '0.1 ' 1
Signif. codes:
```

The Wald statistic is 3.6319 and the p-value is 0.0567, which is less than 10% and greater than 5%. That means that the null hypothesis can be rejected at the 10% level but not at the 5% level. This result implies that the effects of hiring more teachers on test scores could be to some extent similar to increasing more expenditure per pupil.

The homoskedasticity-only F statistic can be computed without specifying vcov or white.adjust.

```
[1] 3.6319
[1] 0.0567
Linear hypothesis test
Hypothesis:
str + I(expn_stu/1000) = 0
Model 1: restricted model
Model 2: testscr ~ str + I(expn_stu/1000) + el_pct
  Res.Df
           RSS Df Sum of Sq
                                 F Pr(>F)
     417 86562
     416 85700 1
                     862.09 4.1847 0.04142 *
2
               0 '*** 0.001 '** 0.01 '* 0.05 '.' 0.1 ' 1
Signif. codes:
```

The homoskedasticity-only F test points to rejecting the null hypothesis at both 5% and 10% levels.

6 Control variables and model specifications

In this section we estimate different models for the application of test scores. The variable of interest is student-teacher ratios, STR. In the base specification, we include the percentage of students who are English learners, PctEL, and the percentage of students who are eligible for free or subsidized lunch, LchPct, as control variables. An alternative control variable is the percentage of students who receive public assistance.

```
model1 <- lm(testscr ~ str, data = classdata)
model2 <- lm(testscr ~ str + el_pct, data = classdata)
model3 <- lm(testscr ~ str + el_pct + meal_pct, data = classdata)
model4 <- lm(testscr ~ str + el_pct + calw_pct, data = classdata)
model5 <- lm(testscr ~ str + el_pct + meal_pct + calw_pct, data = classdata)</pre>
```

We compute the heteroskedasticity-robust standard errors for the coefficients in all model specifications. The function vcovHC is used to get the heteroskedasticity-consistent covariance matrix (HCCM), in which we set the argument type to be HC1. The heteroskedasticity-robust standard errors of coefficients are the square roots of the diagonal elements of these HCCMs.

```
hccm1 <- vcovHC(model1, type = "HC1")
se1 <- sqrt(diag(hccm1))
hccm2 <- vcovHC(model2, type = "HC1")
se2 <- sqrt(diag(hccm2))
hccm3 <- vcovHC(model3, type = "HC1")
se3 <- sqrt(diag(hccm3))</pre>
```

```
hccm4 <- vcovHC(model4, type = "HC1")</pre>
se4 <- sqrt(diag(hccm4))</pre>
hccm5 <- vcovHC(model5, type = "HC1")</pre>
se5 <- sqrt(diag(hccm5))</pre>
Finally, the results for all models are displayed in Table (1) that replicates Table 7.1 in Chapter
7. To create a LATEX table, we use the function stargazer.
stargazer(model1, model2, model3, model4, model5,
  title = "Results of regressions of test scores and class size",
  covariate.labels = c("student-teacher ratio",
       "percent English learners",
       "percent eligible for subsidized lunch",
       "percent on public assistance"),
  dep.var.caption = "average test scores in the district",
  se = list(se1, se2, se3, se4, se5), df = FALSE,
  font.size = "small",
  header = FALSE,
  label = "table:tbl71")
```

7 Appendix: R codes

```
library(AER)
# read the data files into R
# read the dta file
library(foreign)
classdata <- read.dta("caschool.dta")</pre>
head(classdata)
str(classdata)
summary(classdata[, c("testscr", "str", "el_pct", "expn_stu", "meal_pct")])
model.76 <- testscr ~ str + I(expn_stu / 1000) + el_pct</pre>
res.model.76 <- lm(model.76, data = classdata)
summary(res.model.76)
# scatterplot
oldpar \leftarrow par(mfrow = c(2, 2))
plot(classdata$str, classdata$testscr, col = "red",
     main = "student-teacher ratio vs test scores",
     xlab = "Student-teacher ratio", ylab = "Test scores")
```

Table 1: Results of regressions of test scores and class size

	average test scores in the district						
			testscr				
	(1)	(2)	(3)	(4)	(5)		
student-teacher ratio	-2.280^{***} (0.519)	-1.101** (0.433)	-0.998^{***} (0.270)	-1.308^{***} (0.339)	-1.014^{***} (0.269)		
percent English learners		-0.650^{***} (0.031)	-0.122^{***} (0.033)	-0.488^{***} (0.030)	-0.130^{***} (0.036)		
percent eligible for subsidized lunch			-0.547^{***} (0.024)		-0.529^{***} (0.038)		
percent on public assistance				-0.790^{***} (0.068)	-0.048 (0.059)		
Constant	698.933*** (10.364)	686.032*** (8.728)	700.150*** (5.568)	697.999*** (6.920)	700.392*** (5.537)		
Observations	420	420	420	420	420		
\mathbb{R}^2	0.051	0.426	0.775	0.629	0.775		
Adjusted R ²	0.049	0.424	0.773	0.626	0.773		
Residual Std. Error	18.581	14.464	9.080	11.654	9.084		
F Statistic	22.575***	155.014***	476.306***	234.638***	357.054***		

Note:

*p<0.1; **p<0.05; ***p<0.01

```
plot(classdata$el_pct, classdata$testscr, col = "blue",
     main = "English learners vs test scores",
     xlab = "Percentage of English learners",
     ylab = "Test scores")
plot(classdata$expn_stu, classdata$testscr, col = "green",
     main = "Expenditure per pupil vs test scores",
     xlab = "Expenditure per pupil",
     ylab = "Test scores")
plot(classdata$meal_pct, classdata$testscr, col = "black",
     main = "Free lunch vs test scores",
     xlab = "Percentage of students with free lunch",
     ylab = "Test scores")
par(oldpar)
# do this in a loop
xvars <- c("str", "el_pct", "expn_stu", "meal_pct")</pre>
yvars <- c("testscr")</pre>
xlabels <- c("Student-teacher ratio", "Percentage of English learners",</pre>
             "Expenditure per pupil", "Percentage of students with free lunch")
ylabels <- "Test scores"</pre>
titles <- c("student-teacher ratio vs test scores",
            "English learners vs test scores",
            "Expenditure per pupil vs test scores",
            "Free lunch vs test scores")
colors <- c("red", "green3", "blue", "black")</pre>
op \leftarrow par(mfrow = c(2, 2))
for (i in seq(along=xvars)) {
    fm <- formula(paste(yvars, "~", xvars[i]))</pre>
    plot(fm, data = classdata, col = colors[i], main = titles[i],
         xlab = xlabels[i], ylab = ylabels)
par(op)
## # use pairs() or scatterplot.matrix() in car
## pairs(~ testscr +str + el_pct + expn_stu + meal_pct,
##
         data = classdata)
```

```
## scatterplot.matrix(~ testscr +str + el_pct + expn_stu + meal_pct,
##
                    data = classdata)
###
                estimation
                                         ###
model.76 <- testscr ~ str + I(expn_stu / 1000) + el_pct</pre>
res.model.76 <- lm(model.76, data = classdata)
summary(res.model.76)
# get some components of the results
bhat <- coef(res.model.76)</pre>
rsq <- summary(res.model.76)$r.squared
adj.rsq <- summary(res.model.76)$adj.r.squared</pre>
htvarm <- vcovHC(res.model.76, type = "HC1")</pre>
# application of the FWL theorem
# purge the effect of English learners and expenditure per pupil
mod.fwl1 <- lm(testscr ~ el_pct + I(expn_stu/1000), data = classdata)</pre>
mod.fwl2 <- lm(str ~ el_pct + I(expn_stu/1000), data = classdata)</pre>
mod.fwl3 <- lm(resid(mod.fwl1) ~ resid(mod.fwl2))</pre>
# test whether the FWL theorm works
# first check for the equality of the coefficient on str
if (abs(coef(res.model.76)[2] - coef(mod.fwl3)[2]) < 1.0e-10) {
   cat("The coefficient on str is the same.\n")
   } else {
   cat("The FWL theorem fails? Check your step!\n")
}
# second check for the equality of residuals
if (all(abs(resid(mod.fwl3) - resid(res.model.76)) < 1.0e-10)) {</pre>
   cat("The residuals are all the same.\n")
   } else {
   cat("The FWL theorem fails? Check your step!\n")
}
###
                                         ###
            hypothesis testing
```

```
# single coefficient test
# homoskedasticity-only
coeftest(res.model.76)
# heteroskedasticity-robust, t distribution
cftest.t <- coeftest(res.model.76, vcov = htvarm)</pre>
# heteroskedasticity-robust, normal distribution
cftest.n <- coeftest(res.model.76, vcov = htvarm, df = Inf)</pre>
# joint hypothesis
# b1 = 0, b2 = 0
test1 <- linearHypothesis(res.model.76,
           c("str = 0", "I(expn_stu/1000) = 0"),
           vcov = htvarm, test = "F")
test1
test1.F <- test1$F[2]
test1.p <- test1$"Pr(>F)"[2]
# b1 + b2 = 0
test2 <- linearHypothesis(res.model.76,
           c("str + I(expn_stu/1000) = 0"),
           white.adjust = "hc1", test = "Chisq")
test2
test2.x <- test2$Chisq[2]
test2.p <- test2$"Pr(>Chisq)"[2]
# homoskedasticity-only F
test2.hm <- linearHypothesis(res.model.76,</pre>
           c("str + I(expn_stu/1000) = 0"),
           test = "Chisq")
test2.hm
Control variables and model specifications
# replicate Table 7.1
model1 <- lm(testscr ~ str, data = classdata)</pre>
model2 <- lm(testscr ~ str + el_pct, data = classdata)</pre>
model3 <- lm(testscr ~ str + el_pct + meal_pct, data = classdata)</pre>
model4 <- lm(testscr ~ str + el_pct + calw_pct, data = classdata)</pre>
model5 <- lm(testscr ~ str + el_pct + meal_pct + calw_pct, data = classdata)</pre>
```

```
hccm1 <- vcovHC(model1, type = "HC1")</pre>
se1 <- sqrt(diag(hccm1))</pre>
hccm2 <- vcovHC(model2, type = "HC1")</pre>
se2 <- sqrt(diag(hccm2))</pre>
hccm3 <- vcovHC(model3, type = "HC1")</pre>
se3 <- sqrt(diag(hccm3))</pre>
hccm4 <- vcovHC(model4, type = "HC1")</pre>
se4 <- sqrt(diag(hccm4))</pre>
hccm5 <- vcovHC(model5, type = "HC1")</pre>
se5 <- sqrt(diag(hccm5))</pre>
library(stargazer)
stargazer(model1, model2, model3, model4, model5,
           title = "Results of regressions of test scores and class size",
           covariate.labels = c("student-teacher ratio",
                                  "percent English learners",
                                  "percent eligible for subsidized lunch",
                                  "percent on public assistance"),
           dep.var.caption = "average test scores in the district",
           se = list(se1, se2, se3, se4, se5), df = FALSE,
           font.size = "small",
           label = "table:tbl71")
```

References

James H Stock and Mark W. Watson. *Introduction to Econometrics*. Addison Wesley Longman, Boston, 3rd edition, 2011.