Replication of Examples in Chapter 4

Zheng Tian 3/21/2017

Introduction

This document is to show how to estimate a simple regression model and perform linear hypothesis testing. The application concerns the test scores of the California school districts. We will use R to estimate the simple regression model with the data set in Chapter 4.

Before running all R codes, we may first load the package AER by running library (AER).

Reading the data and basic summary statistics

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

Read the data file

The textbook comes with two files for the California test score data set, caschool.xlsx and caschool.dta, the former of which is an Excel file and the latter is a Stata file. We need to read either one of these two files into R so that we can use the data set.

R has several built-in functions that can read an ASCII data file, which can have such an extension as .txt, .csv, dat, to name a few. However, these built-in functions cannot handle an Excel file or a Stata file. So, in order to read caschool.xlsx or caschool.dta in to R, we need additional packages.

To read a State file with an extension of .dta, we use the function, read.dta, in the library of foreign.

```
library(foreign)
classdata <- read.dta("caschool.dta")</pre>
```

classdata is a dataframe object in R. If you want to check whether your reading is correct, you can type head(classdata), which, by default, displays the first six observations of all variables in the data frame object.

```
head(classdata[c("observat", "district", "testscr", "str")])
```

```
##
     observat
                                     district testscr
## 1
            1
                           Sunol Glen Unified 690.80 17.88991
            2
## 2
                         Manzanita Elementary 661.20 21.52466
## 3
                  Thermalito Union Elementary 643.60 18.69723
## 4
            4 Golden Feather Union Elementary 647.70 17.35714
## 5
            5
                     Palermo Union Elementary 640.85 18.67133
## 6
                      Burrel Union Elementary 605.55 21.40625
```

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 4, including test scores, testscr, student-teacher ratio, str.

```
df <- classdata[c("testscr", "str")]
summary(df)</pre>
```

```
##
       testscr
                          str
##
    Min.
           :605.5
                     Min.
                            :14.00
##
    1st Qu.:640.0
                     1st Qu.:18.58
##
   Median :654.5
                     Median :19.72
##
   Mean
           :654.2
                     Mean
                            :19.64
##
  3rd Qu.:666.7
                     3rd Qu.:20.87
   Max.
           :706.8
                     Max.
                             :25.80
```

Formally, we can create a table showing important summary statistics, like the following table (Table 4.1 in the book).

```
# Replicate the summary statistics in Table 4.1
summary4.1 <- function(df) {
   ave <- sapply(df, mean)
   std <- sapply(df, sd)
   perctile <- sapply(df, function(x)
   quantile(x, probs = c(0.1, 0.25, 0.4, 0.5, 0.6, 0.75, 0.9)))
   return(rbind(ave, std, perctile))
}
library(xtable)
sumtab <- xtable(t(summary4.1(df)))</pre>
```

In the above code, I defined a function called summary4.1. In R, we use function() to define a custom function. In this case, the function summary4.1 takes one argument df. The code within the function is enclosed with the curly braces, { }, and what the function yields is controlled by return() in the last line.

Within the function summary4.1, I use a special function in R, sapply(). It takes each component in a list object, which is df in this case, and impose a function on this component, and return a simplified object. Check the help information for apply(), lapply(), sapply(), mapply(), and tapply().

Finally, the function xtable() writes the matrix that contains the summary statistics into a table in either a LaTeX or an HTML file.

```
# print as a latex table
print(sumtab, type = "latex")
```

% latex table generated in R 3.3.2 by xtable 1.8-2 package % Sun Mar 26 15:29:33 2017

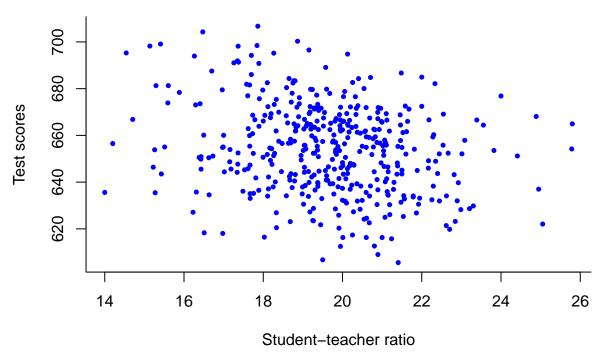
	ave	std	10%	25%	40%	50%	60%	75%	90%
testscr	654.16	19.05	630.40	640.05	649.07	654.45	659.40	666.66	678.86
str	19.64	1.89	17.35	18.58	19.27	19.72	20.08	20.87	21.87

Create a scatterplot using plot()

It is always a good practice to make a scatterplot between an independent variable and a dependent variable before setting up a regression model. Let's draw a scatterplot of student-teacher ratios and test scores. plot is the most basic function in R to draw figures. Here I use it to draw the scatterplot as follows.

```
# generate a scatterplot
plot(df$str, df$testscr, col = "blue", pch =16, cex = 0.7, bty = "l",
    main = "Scatterplot of Test Score vs. Student-Teacher Ratio",
    xlab = "Student-teacher ratio", ylab = "Test scores")
```

Scatterplot of Test Score vs. Student-Teacher Ratio



We can compute the correlation coefficient between the two variables, using the function cor.

```
# calculate correlation coefficient
cor(df$str, df$testscr)
```

[1] -0.2263628

The OLS estimation

Set up the linear regression model

We establish the following linear regression model for the relationship between test scores and class sizes

$$TestScore_i = \beta_0 + \beta_1 STR_i + u_i$$

Estimate in R

The OLS estimation can be implemented in R with the function 1m. The most important argument in this function is the model to be estimated, which is called a **formula** object in R. A formula is defined using the format $y \sim x1 + x2$, in which the symbol of \sim links the left-hand side variable, y, and the right-hand side variables, x1, x2.

We can add more independent variables in the right-hand side with each being appended to the formula by the symbol of +. The constant term is by default included in the model. After estimation, we use summary to see the results.

```
mod1 <- lm(testscr ~ str, data = df)
summary(mod1)</pre>
```

```
##
## Call:
## lm(formula = testscr ~ str, data = df)
##
## Residuals:
##
      Min
               1Q Median
                               3Q
                                      Max
## -47.727 -14.251
                    0.483 12.822 48.540
##
## Coefficients:
##
              Estimate Std. Error t value Pr(>|t|)
## (Intercept) 698.9330
                           9.4675 73.825 < 2e-16 ***
                           0.4798 -4.751 2.78e-06 ***
               -2.2798
## str
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 18.58 on 418 degrees of freedom
## Multiple R-squared: 0.05124,
                                   Adjusted R-squared: 0.04897
## F-statistic: 22.58 on 1 and 418 DF, p-value: 2.783e-06
```

For now, we just pay attention to the estimates of the two coefficients, which is 698.93 for the intercept, β , and -2.28 for the slope.

$$\widehat{TestScore} = 698.93 - 2.28 \times STR$$

Plot the sample regression line

The sample regression line can be added to the scatterplot by using the function abline. And an annotation can be added by using the function text

