

Exercises Chapter 6

1. An alternative, equivalent representation of the F -test statistic is the following:

$$F = \frac{(\sum_{i=1}^n \hat{\varepsilon}_{iR}^2 - \sum_{i=1}^n \hat{\varepsilon}_{iU}^2) / q}{(\sum_{i=1}^n \hat{\varepsilon}_{iU}^2) / (n - K)} = \frac{(SS_R - SS_U) / q}{SS_U / (n - K)},$$

where $\hat{\varepsilon}_{iU}$ are the residuals from the *unrestricted* (i.e., the usual) regression of Y on X , and where $\hat{\varepsilon}_{iR}$ are the residuals from the *restricted* ordinary least squares regression which minimizes the following *restricted* version of the OLS-objective function

$$\min_{\tilde{\beta}} S_n(\tilde{\beta}) = (Y - X\tilde{\beta})'(Y - X\tilde{\beta}) \quad \text{such that} \quad R\tilde{\beta} - r = 0$$

where the restriction is just the null hypothesis.

The standard F -test. The standard F -test for a linear regression tests the hypothesis that all coefficients except the intercept are equal to zero. In this case, $\hat{\varepsilon}_{iR}$ are simply the residuals from regressing Y on only the intercept. In this standard case we have,

$$F_1 = \frac{(\sum_{i=1}^n (Y_i - \bar{Y})^2 - \sum_{i=1}^n \hat{\varepsilon}_{iU}^2) / (K - 1)}{(\sum_{i=1}^n \hat{\varepsilon}_{iU}^2) / (n - K)}$$

since here $\hat{\varepsilon}_{iR}^2 = (Y_i - \bar{Y})^2$.

Show that F_1 is equal to F_2 with

$$F_2 = \frac{R_U^2 / (K - 1)}{(1 - R_U^2) / (n - K)},$$

where R_U^2 denotes the coefficient of determination of the unrestricted regression model.

2. Install the R package AER and load the package. The ARE-package contains the data set Journals. Check ?Journals to learn more about the data. Create the variables citeprice (journal price per citations) and age (journal age) as following:

```
> # install.packages("AER")
> suppressMessages(library("AER"))
> ## attach the data-set Journals to the current R-session
> data("Journals", package = "AER")
> ## ?Journals # Check the help file
> ##
> ## Select variables "subs" and "price"
> journals <- Journals[, c("subs", "price")]
> ## Define variable 'journal-price per citation'
> journals$citeprice <- Journals$price/Journals$citations
> ## Define variable 'journal-age'
> journals$age <- 2020 - Journals$foundingyear
> ## Check variable names in 'journals'
> names(journals)

[1] "subs"      "price"     "citeprice" "age"
```

Estimate the coefficients β_1 and β_2 of the following linear regression model

$$\log(Y_i) = \beta_1 + \beta_2 \log(X_i) + \varepsilon_i, \quad i = 1, \dots, n$$

with $\log(Y) = \log(\text{subs})$ (i.e., logarithm of the number of library subscriptions) and $\log(X) = \log(\text{citeprice})$ (i.e., logarithm of the journal price per citations).

- (a) Do you have heteroscedastic error-term variances? Explain your answer by discussing a diagnostic plot showing the residuals against the fitted values.
- (b) Estimate the standard error of the OLS estimator $\hat{\beta}_2$ using an appropriate variance estimator.

3. Consider the following multiple linear regression model:

$$Y_i = \beta_1 + \beta_2 X_{2i} + \beta_3 X_{3i} + \varepsilon_i, \quad i = 1, \dots, n$$

(in matrix notation) $Y = X\beta + \varepsilon$

where $\beta = (1, -5, 5)'$, ε_i is a heteroscedastic error term

$$\varepsilon_i \sim N(0, \sigma_i^2) \quad \text{with} \quad \sigma_i = |X_{3i}|,$$

and where for all $i = 1, \dots, n = 100$:

- $X_{2i} \sim N(10, 1.5^2)$
- $X_{3i} \sim U[0.2, 8]$

You're given the following data generated from this regression model:

```
> set.seed(109) # Sets the "seed" of the random number generators:
> n <- 100      # Number of observations
> ## Generate two explanatory variables plus an intercept-variable:
> X_1 <- rep(1, n)          # Intercept
> X_2 <- rnorm(n, mean=10, sd=1.5) # Draw realizations from a normal distr.
> X_3 <- runif(n, min = 0.2, max = 8) # Draw realizations from a t-distr.
> X <- cbind(X_1, X_2, X_3)   # Save as a Nx3-dimensional data matrix.
> beta <- c(1, -5, 5)
> ## Generate realizations from the heteroscedastic error term
> eps <- rnorm(n, mean=0, sd=abs(X_3))
> ## Dependent variable:
> Y <- X %*% beta + eps
```

- (a) Compute the theoretical covariance matrix variance $\text{Var}(\hat{\beta})$ of the OLS estimator $\hat{\beta}$ for the given data generating process and the given data.
- (b) Use a Monte-Carlo simulation to generate 10000 variance estimates

$$\widehat{\text{Var}}_{\text{HC3},1}(\hat{\beta}_2), \dots, \widehat{\text{Var}}_{\text{HC3},10000}(\hat{\beta}_2)$$

and 10000 variance estimates

$$\widehat{\text{Var}}_{\text{HC3},1}(\hat{\beta}_3), \dots, \widehat{\text{Var}}_{\text{HC3},10000}(\hat{\beta}_3).$$

These estimates represent typical estimation results. (Of course, in practice you observe only one variance estimation result $\widehat{\text{Var}}_{\text{HC3}}(\hat{\beta}_2)$ for $\text{Var}(\hat{\beta}_2)$ and one $\widehat{\text{Var}}_{\text{HC3}}(\hat{\beta}_3)$ for $\text{Var}(\hat{\beta}_3)$.)

- (i) Visualize the Monte Carlo realizations for the variance estimates. Add points displaying the sample mean of the Monte Carlo realizations and points displaying the true variance values.
- (ii) Do the Monte Carlo realizations $\widehat{\text{Var}}_{\text{HC3},r}(\hat{\beta}_2)$ and $\widehat{\text{Var}}_{\text{HC3},r}(\hat{\beta}_3)$, $r = 1, \dots, 10000$ estimate the true variances $\text{Var}(\hat{\beta}_2)$ and $\text{Var}(\hat{\beta}_3)$ well on average?
- (iii) Are there large estimation uncertainties?

4. The `Boston` housing data set (contained in the R package `MASS`) contains observations on housing values in suburbs of Boston. Let's consider the following regression model

$$\text{medv}_i = \beta_1 + \beta_2 \text{ptratio}_i + \beta_3 \text{lstat}_i + \beta_4 \text{age}_i + \beta_5 \text{crim}_i + \beta_6 \text{nox}_i + \varepsilon_i$$

where $i = 1, \dots, n$ indexes the suburbs. Check `?Boston` in R to get an overview about the variables. You can assume that the assumptions of Chapter 6 hold. The following R code computes the regression estimates:

```
> library("lmtest") # for coeftest()
> library("sandwich") # for robust se
> library("MASS") # for Boston housing data
> data("Boston") # Check: ?Boston; names(Boston)
> lm_obj <- lm(medv ~ ptratio + lstat + age + crim + nox, data = Boston)
> vcovHC3_mat <- vcovHC(lm_obj, type = "HC3")
> round(coeftest(lm_obj, vcov = vcovHC3_mat), 3)
```

t test of coefficients:

	Estimate	Std. Error	t value	Pr(> t)
(Intercept)	55.746	3.013	18.504	<2e-16 ***
ptratio	-1.181	0.147	-8.046	<2e-16 ***
lstat	-0.868	0.081	-10.662	<2e-16 ***
age	0.060	0.017	3.527	<2e-16 ***
crim	-0.024	0.036	-0.674	0.501
nox	-8.059	3.318	-2.429	0.016 *

Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

- Use R to test $H_0 : \beta_6 = 0$ versus $H_A : \beta_6 < 0$ by means of an t -test. What is the correct p -value and what is the test decision when using a significance level of $\alpha = 0.01$?
- Use R to test $H_0 : \beta_5 = \beta_6 = 0$ versus $H_A : \beta_5 \neq 0$ and/or $\beta_6 \neq 0$ by means of an F -test. What is the marginal significance value in this case?
- What is the maximal probability of a type I error if you test the null hypothesis in (b) by means of two separate t -tests instead of one F -test? How does this compare to the probability of a type I error for the F test in (b).