

5 Regression Inference

```
library(tidyverse)
library(kableExtra)
library(sandwich)
library(lmtest)
```

5.1 Standardized coefficients

The j -th OLS coefficient has the conditional standard deviation

$$sd(\hat{\beta}_j|\mathbf{X}) = \sqrt{[(\mathbf{X}'\mathbf{X})^{-1}\mathbf{X}'\mathbf{D}\mathbf{X}(\mathbf{X}'\mathbf{X})^{-1}]_{jj}}.$$

Note that $[\mathbf{A}]_{jj}$ indicates the j -th diagonal element of the matrix \mathbf{A} .

Under the homoskedasticity assumption (A5), the standard deviation simplifies to

$$sd(\hat{\beta}_j|\mathbf{X}) = \sqrt{\sigma^2[(\mathbf{X}'\mathbf{X})^{-1}]_{jj}}.$$

The coefficient is unbiased with $E[\hat{\beta}_j|\mathbf{X}] = \beta_j$ and has the standardized representation

$$Z_j := \frac{\hat{\beta}_j - \beta_j}{sd(\hat{\beta}_j|\mathbf{X})}.$$

Under (A1)–(A4), $\sqrt{n}(\hat{\beta}_j - \beta_j)$ converges to a normal distribution, and therefore

$$Z_j \xrightarrow{D} \mathcal{N}(0, 1) \quad \text{as } n \rightarrow \infty.$$

A direct consequence is that

$$\lim_{n \rightarrow \infty} P\left(\hat{\beta}_j - z_{(1-\frac{\alpha}{2})}sd(\hat{\beta}_j|\mathbf{X}) \leq \beta_j \leq \hat{\beta}_j + z_{(1-\frac{\alpha}{2})}sd(\hat{\beta}_j|\mathbf{X})\right) = 1 - \alpha,$$

where $z_{(p)}$ is the p -quantile of the standard normal distribution. Thus, $\hat{\beta}_j \pm z_{(1-\frac{\alpha}{2})}sd(\hat{\beta}_j|\mathbf{X})$ defines an asymptotic $1 - \alpha$ confidence interval for β_j .

Under the normality assumption (A6), the OLS estimator $\hat{\beta}_j$ is normal conditional on \mathbf{X} , which implies that $Z_j \sim \mathcal{N}(0, 1)$ for any fixed sample size n . In this case, $\hat{\beta}_j \pm z_{(1-\frac{\alpha}{2})} sd(\hat{\beta}_j|\mathbf{X})$ is an exact confidence interval for β_j .

Note that \mathbf{D} is unknown and $sd(\hat{\beta}_j|\mathbf{X})$ is not computable in practice, so the confidence interval is not feasible.

5.2 Standard Errors

A standard error $se(\hat{\beta}_j)$ for an estimator $\hat{\beta}_j$ is an estimator of the standard deviation of the distribution of $\hat{\beta}_j$.

We say that the standard error is consistent if

$$\frac{se(\hat{\beta}_j)}{sd(\hat{\beta}_j|\mathbf{X})} \xrightarrow{p} 1.$$

This property ensures that, in practice, we can replace the unknown standard deviation with the standard error to apply inferential methods such as confidence intervals and t-tests.

To estimate the unknown standard deviation of the OLS estimator, the diagonal matrix $\mathbf{D} = \text{diag}(\sigma_1^2, \dots, \sigma_n^2)$ is replaced by some sample counterpart $\widehat{\mathbf{D}} = \text{diag}(\hat{\sigma}_1^2, \dots, \hat{\sigma}_n^2)$.

5.2.1 Robust standard errors

Various **heteroskedasticity-consistent (HC)** standard errors have been proposed in the literature:

HC type	weights
HC0	$\hat{\sigma}_i^2 = \hat{u}_i^2$
HC1	$\hat{\sigma}_i^2 = \frac{n}{n-k} \hat{u}_i^2$
HC2	$\hat{\sigma}_i^2 = \frac{\hat{u}_i^2}{1-h_{ii}}$
HC3	$\hat{\sigma}_i^2 = \frac{\hat{u}_i^2}{(1-h_{ii})^2}$

HC0 replaces the unknown variances with squared residuals, and HC1 is a bias-corrected version of HC0. HC2 and HC3 use the leverage values h_{ii} (the diagonal entries of the influence matrix \mathbf{P}) and give less weight to influential observations.

HC1 and HC3 are the most common choices and can be written as

$$se_{hc1}(\hat{\beta}_j) = \sqrt{\left[(\mathbf{X}'\mathbf{X})^{-1} \left(\frac{n}{n-k} \sum_{i=1}^n \hat{u}_i^2 \mathbf{X}_i \mathbf{X}_i' \right) (\mathbf{X}'\mathbf{X})^{-1} \right]_{jj}},$$

$$se_{hc3}(\hat{\beta}_j) = \sqrt{\left[(\mathbf{X}'\mathbf{X})^{-1} \left(\sum_{i=1}^n \frac{\hat{u}_i^2}{(1-h_{ii})^2} \mathbf{X}_i \mathbf{X}_i' \right) (\mathbf{X}'\mathbf{X})^{-1} \right]_{jj}}.$$

All versions perform similarly well in large samples, but HC3 performs best in small samples and is the preferred choice.

HC standard errors are also known as **heteroskedasticity-robust standard errors** or simply **robust standard errors**.

Estimators for the full covariance matrix of $\hat{\beta}$ have the form

$$\widehat{\mathbf{V}} = (\mathbf{X}'\mathbf{X})^{-1} \mathbf{X}' \widehat{\mathbf{D}} \mathbf{X} (\mathbf{X}'\mathbf{X})^{-1}.$$

The HC3 covariance estimator can be written as

$$\widehat{\mathbf{V}}_{hc3} = (\mathbf{X}'\mathbf{X})^{-1} \left(\sum_{i=1}^n \frac{\hat{u}_i^2}{(1-h_{ii})^2} \mathbf{X}_i \mathbf{X}_i' \right) (\mathbf{X}'\mathbf{X})^{-1}.$$

5.2.2 Classical standard errors

Classical standard errors put equal weights on all observations:

$$\hat{\sigma}_i^2 = s_u^2 = \frac{1}{n-k} \sum_{j=1}^n \hat{u}_j^2.$$

This implies $\widehat{\mathbf{D}} = s_u^2 \mathbf{I}_n$ and $\mathbf{X}' \widehat{\mathbf{D}} \mathbf{X} = s_u^2 \mathbf{X}' \mathbf{X}$. Therefore, the classical covariance matrix estimator reduces to

$$\widehat{\mathbf{V}}_{hom} = s_u^2 (\mathbf{X}'\mathbf{X})^{-1}.$$

The classical standard errors are

$$se_{hom}(\hat{\beta}_j) = \sqrt{s_u^2 [(\mathbf{X}'\mathbf{X})^{-1}]_{jj}}.$$

Classical standard errors are only valid under (A5) and are also known as **constant variance standard errors** or **homoskedasticity-only standard errors**. Classical standard errors should only be used if there are very good reasons for the error terms to be homoskedastic.

5.2.3 Standard Errors in R

The covariance matrix estimates can be computed using the `vcovHC()` function from the `sandwich` package. HC3 is the default version. The standard errors are the square roots of their diagonal entries.

```
fit = lm(wage ~ education + experience + black + female, data = cps)
hom = sqrt(diag(vcovHC(fit, "const")))
HC1 = sqrt(diag(vcovHC(fit, "HC1")))
HC3 = sqrt(diag(vcovHC(fit)))
tibble("Variable" = names(coefficients(fit)), hom, HC1, HC3) |>
  mutate_if(is.numeric, round, digits = 4) |>
  kbl(align = 'c')
```

Variable	hom	HC1	HC3
(Intercept)	0.4910	0.5666	0.5667
education	0.0305	0.0408	0.0409
experience	0.0072	0.0067	0.0067
black	0.2684	0.2243	0.2243
female	0.1670	0.1603	0.1604

5.3 Interval estimates

5.3.1 Asymptotic Intervals

A **confidence interval** $I_{1-\alpha}$ for β_j with coverage probability $1 - \alpha$ is asymptotically valid if

$$\lim_{n \rightarrow \infty} P(\beta_j \in I_{1-\alpha}) = 1 - \alpha.$$

Under (A1)–(A4), we can use

$$I_{1-\alpha} = [\hat{\beta}_j - z_{(1-\frac{\alpha}{2})} se_{hc}(\hat{\beta}_j); \hat{\beta}_j + z_{(1-\frac{\alpha}{2})} se_{hc}(\hat{\beta}_j)],$$

where $se_{hc}(\hat{\beta}_j)$ is any HC-type standard error. $z_{(p)}$ can be returned using `qnorm(p)`.

In practice, t-quantiles are often used instead of z-quantiles:

$$I_{1-\alpha} = [\hat{\beta}_j - t_{(1-\frac{\alpha}{2}, n-k)} se_{hc}(\hat{\beta}_j); \hat{\beta}_j + t_{(1-\frac{\alpha}{2}, n-k)} se_{hc}(\hat{\beta}_j)],$$

where $t_{(p,m)}$ is the p -quantile of the t-distribution with m degrees of freedom. $t_{(p,m)}$ can be returned using `qt(p,m)`.

Asymptotically, it makes no difference whether t- or z-quantiles are used. We have

$$t_{(1-\frac{\alpha}{2}, n-k)} > z_{(1-\frac{\alpha}{2})}$$

for any fixed n , which makes the t-based confidence intervals a little wider (conservative), but asymptotically they coincide because

$$\lim_{n \rightarrow \infty} t_{(1-\frac{\alpha}{2}, n-k)} = z_{(1-\frac{\alpha}{2})}.$$

You can use the `coefci()` function from the `lmtest` package. `coefci(fit)` calculates classical confidence intervals, `coefci(fit, vcov. = vcovHC)` uses HC3 standard errors, and `coefci(fit, vcov. = vcovHC, df=Inf)` considers z-quantiles instead of t-quantiles.

```
coefci(fit, vcov. = vcovHC)
```

	2.5 %	97.5 %
(Intercept)	-22.8201704	-20.5988645
education	3.0549552	3.2151008
experience	0.2311859	0.2574641
black	-3.2951083	-2.4157606
female	-7.7505755	-7.1219793

You can use `qt(p, df = nu)` and `qnorm(p)` to get the t- and z-quantiles, where p is the probability and nu is the degrees of freedom. The CDF values for the standard normal and t-distributions can be calculated using `pt()` and `pnorm()`.

5.3.2 Exact Intervals

An exact confidence interval $I_{1-\alpha}$ for β_j satisfies

$$P(\beta_j \in I_{1-\alpha}) = 1 - \alpha$$

for any sample size n .

Exact confidence intervals for the regression coefficients are only available if the homoskedasticity and normality assumptions (A5) and (A6) hold. In this case,

$$\frac{(n-k)s_u^2}{\sigma^2} \sim \chi_{n-k}^2,$$

which implies that

$$\frac{se_{hom}(\hat{\beta}_j)}{sd(\hat{\beta}_j|\mathbf{X})} \sim \sqrt{\chi_{n-k}^2 / (n-k)}.$$

Replacing the true standard deviation with the classical standard error in the standardized OLS coefficient Z_j yields

$$\frac{\hat{\beta}_j - \beta_j}{se_{hom}(\hat{\beta}_j)} = \frac{Z_j}{se_{hom}(\hat{\beta}_j)/sd(\hat{\beta}_j|\mathbf{X})} \sim \frac{\mathcal{N}(0, 1)}{\sqrt{\chi_{n-k}^2/(n-k)}} = t_{n-k}.$$

Therefore,

$$I_{1-\alpha, hom} = [\hat{\beta}_j - t_{(1-\frac{\alpha}{2}, n-k)} se_{hom}(\hat{\beta}_j); \hat{\beta}_j + t_{(1-\frac{\alpha}{2}, n-k)} se_{hom}(\hat{\beta}_j)]$$

is an exact confidence interval for β_j under (A1)–(A6).

5.4 t-Tests

The **t-statistic** is the OLS estimator standardized with the standard error. Under (A1)–(A4) we have

$$T = \frac{\hat{\beta}_j - \beta_j}{se_{hc}(\hat{\beta}_j)} \xrightarrow{D} \mathcal{N}(0, 1).$$

This result can be used to test the hypothesis $H_0 : \beta_j = \beta_j^0$. The t-statistic for this hypothesis is

$$T_0 = \frac{\hat{\beta}_j - \beta_j^0}{se_{hc}(\hat{\beta}_j)},$$

which satisfies $T_0 = T \xrightarrow{D} \mathcal{N}(0, 1)$ under H_0 .

The **two-sided t-test** for H_0 against $H_1 : \beta_j \neq \beta_j^0$ is given by the test decision

$$\begin{aligned} \text{do not reject } H_0 & \quad \text{if } |T_0| \leq t_{(1-\frac{\alpha}{2}, n-k)}, \\ \text{reject } H_0 & \quad \text{if } |T_0| > t_{(1-\frac{\alpha}{2}, n-k)}. \end{aligned}$$

The value $t_{(1-\frac{\alpha}{2}, n-k)}$ is called the **critical value**.

This test is asymptotically of size α :

$$\lim_{n \rightarrow \infty} P(\text{we reject } H_0 | H_0 \text{ is true}) = \alpha.$$

We can also use the critical value $z_{(1-\frac{\alpha}{2})}$ instead of $t_{(1-\frac{\alpha}{2}, n-k)}$ to get an asymptotically valid test of size α .

If (A5)–(A6) hold, and $se_{hom}(\hat{\beta}_j)$ is used instead of $se_{hc}(\hat{\beta}_j)$, then the t-quantile based t-test is of exact size α .

p-values provide a quick alternative way to make the test decision. The t-test decision rule is equivalent to

$$\begin{aligned} &\text{reject } H_0 && \text{if p-value} < \alpha \\ &\text{do not reject } H_0 && \text{if p-value} \geq \alpha, \end{aligned}$$

where

$$p\text{-value} = 2(1 - F(|T_0|)),$$

and F is the CDF of t_{n-k} or $\mathcal{N}(0, 1)$, depending on whether the t- or z-quantile critical values are used.

The p-values can be calculated using $2*(1-pt(abs(T0), n-k))$ and $2*(1-pnorm(abs(T0), n-k))$, where T_0 is the t-statistic for H_0 .

```
coeftest(fit, vcov. = vcovHC)
```

t test of coefficients:

	Estimate	Std. Error	t value	Pr(> t)
(Intercept)	-21.7095175	0.5666566	-38.312	< 2.2e-16 ***
education	3.1350280	0.0408533	76.739	< 2.2e-16 ***
experience	0.2443250	0.0067036	36.447	< 2.2e-16 ***
black	-2.8554345	0.2243222	-12.729	< 2.2e-16 ***
female	-7.4362774	0.1603553	-46.374	< 2.2e-16 ***

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

`coeftest()` is another function from the `lmtest` package and works similarly to `coefci()`. You can specify different standard errors: `coeftest(fit, vcov. = vcovHC, type = "HC1")`. `coeftest(fit)` returns the t-test results for classical standard errors which is identical to the output of the base-R command `summary(fit)`.

To represent very small numbers where there are n zero digits before the first nonzero digit after the decimal point, R uses scientific notation in the form $e-n$. For example, $2.2e-16$ means 0.00000000000000022 .

5.5 Joint Testing

When multiple hypotheses are to be tested, repeated t-tests will not yield valid inferences.

Each t-test has a probability of falsely rejecting H_0 (type I error) of α , but if multiple t-tests are used on different coefficients, then the probability of falsely rejecting at least once (joint type I error probability) is greater than α (multiple testing problem).

5.5.1 Joint Hypotheses

Consider the general hypothesis

$$H_0 : \mathbf{R}\boldsymbol{\beta} = \mathbf{r},$$

where \mathbf{R} is a $q \times k$ matrix with $\text{rank}(\mathbf{R}) = q$ and \mathbf{r} is a $q \times 1$ vector.

Let's look at a linear regression with $k = 3$:

$$Y_i = \beta_1 + \beta_2 X_{i2} + \beta_3 X_{i3} + u_i$$

- Example 1: The hypothesis $H_0 : (\beta_2 = 0 \text{ and } \beta_3 = 0)$ implies $q = 2$ constraints and is translated to $H_0 : \mathbf{R}\boldsymbol{\beta} = \mathbf{r}$ with

$$\mathbf{R} = \begin{pmatrix} 0 & 1 & 0 \\ 0 & 0 & 1 \end{pmatrix}, \quad \mathbf{r} = \begin{pmatrix} 0 \\ 0 \end{pmatrix}.$$

- Example 2: The hypothesis $H_0 : \beta_2 + \beta_3 = 1$ implies $q = 1$ constraint and is translated to $H_0 : \mathbf{R}\boldsymbol{\beta} = \mathbf{r}$ with

$$\mathbf{R} = (0 \quad 1 \quad 1), \quad \mathbf{r} = (1).$$

In practice, the most common multiple hypothesis tests are tests of whether multiple coefficients are equal to zero, which is a test of whether those regressors should be included in the model.

5.5.2 Wald Test

The Wald distance is the vector $\mathbf{d} = \mathbf{R}\hat{\boldsymbol{\beta}} - \mathbf{r}$, and the Wald statistic is the squared standardized Wald distance vector:

$$W = (\mathbf{R}\hat{\boldsymbol{\beta}} - \mathbf{r})' (\mathbf{R}\widehat{\mathbf{V}}\mathbf{R}')^{-1} (\mathbf{R}\hat{\boldsymbol{\beta}} - \mathbf{r}).$$

Under H_0 we have

$$W \xrightarrow{D} \chi_q^2.$$

The test decision for the **Wald test**:

$$\begin{aligned} \text{do not reject } H_0 & \quad \text{if } W \leq \chi_{(1-\alpha, q)}^2, \\ \text{reject } H_0 & \quad \text{if } W > \chi_{(1-\alpha, q)}^2, \end{aligned}$$

where $\chi_{(p, m)}^2$ is the p -quantile of the chi-squared distribution with m degrees of freedom. $\chi_{(p, m)}^2$ can be returned using `qchisq(p, m)`.

5.5.3 F-Test

The F statistic is the Wald statistic scaled by by the number of constraints:

$$F = \frac{W}{q} = \frac{1}{q}(\mathbf{R}\hat{\boldsymbol{\beta}} - \mathbf{r})'(\mathbf{R}\widehat{\mathbf{V}}\mathbf{R}')^{-1}(\mathbf{R}\hat{\boldsymbol{\beta}} - \mathbf{r}).$$

The test decision for the **F-test**:

$$\begin{aligned} \text{do not reject } H_0 & \quad \text{if } F \leq F_{(1-\alpha, q, n-k)}, \\ \text{reject } H_0 & \quad \text{if } F > F_{(1-\alpha, q, n-k)}, \end{aligned}$$

where $F_{(p, m_1, m_2)}$ is the p -quantile of the F distribution with m_1 degrees of freedom in the numerator and m_2 degrees of freedom in the denominator. $F_{(p, m_1, m_2)}$ can be returned using `qf(p, m1, m2)`.

For single constraint ($q = 1$) hypotheses of the form $H_0 : \beta_j = \beta_j^0$, the Wald test is equivalent to a t-test using the z-quantile, and the F-test is equivalent to a t-test using the t-quantile.

The Wald and the F-test are asymptotically equivalent and have asymptotic sizes α under (A1)–(A4) when a HC version of the covariance matrix estimator $\widehat{\mathbf{V}}$ is used. The F test is slightly more conservative for small samples.

In the special case of homoscedastic and normal errors (A5)–(A6), the F test has exact size α when $\widehat{\mathbf{V}}_{hom}$ is used, similar to the exact t-test.

5.5.4 Testing in R

In our regression from above, we can test whether the two coefficients for `experience` and `female` are both zero. The `waldtest()` function from the `lmtest` package allows you to specify the names of the variables directly.

```
waldtest(fit, c("experience", "female"), vcov = vcovHC)
```

```
Wald test
```

```
Model 1: wage ~ education + experience + black + female
```

```
Model 2: wage ~ education + black
```

```
  Res.Df Df      F    Pr(>F)
1    50737
2    50739  -2 1490.9 < 2.2e-16 ***
```

```
---
```

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

```
waldtest(fit, c("experience", "female"), vcov = vcovHC, test = "Chisq")
```

Wald test

Model 1: wage ~ education + experience + black + female

Model 2: wage ~ education + black

	Res.Df	Df	Chisq	Pr(>Chisq)
1	50737			
2	50739	-2	2981.8	< 2.2e-16 ***

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

An alternative is to fit a nested model and apply the function to the fitted models. The following command will produce the same output as above:

```
fit2 = lm(wage ~ education + black, data = cps)
waldtest(fit, fit2, vcov = vcovHC)
```

User-specified constraints of the general form $R\beta = r$ can be tested with the `linearHypothesis()` function from the `car` package.

5.6 R-codes

[methods-sec05.R](#)