

Estimating σ^2

- We now know a lot about estimating β .
- There's another parameter in the regression model - σ^2 - the variance of each ε_i .
- Note that $\sigma^2 = \text{var.}(\varepsilon_i) = E[(\varepsilon_i - E(\varepsilon_i))^2] = E(\varepsilon_i^2)$.
- The *sample* counterpart to this *population* parameter is the *sample* average of the "residuals": $\hat{\sigma}^2 = \frac{1}{n} \sum_{i=1}^n e_i^2 = \frac{1}{n} \mathbf{e}'\mathbf{e}$.
- However, there is a *distortion* in this estimator of σ^2 .
- Although mean of e_i 's is zero (if intercept in model), not all of e_i 's are independent of each other - only $(n - k)$ of them are.

It can be shown that:

$$E(\hat{\sigma}^2) = E\left(\frac{1}{n} \mathbf{e}'\mathbf{e}\right) = \frac{1}{n}(n - k)\sigma^2 < \sigma^2 \quad ; \quad \mathbf{BIASED}$$

Easy to convert this to an *Unbiased estimator* -

$$s^2 = \frac{1}{(n - k)} \mathbf{e}'\mathbf{e}$$

- " $(n - k)$ " is the "*degrees of freedom*" - number of independent sources of information in the " n " residuals (e_i 's).
- We can use " s " as an estimator of σ , but it is a *biased estimator*.
- Call " s " the "*standard error of the regression*", or the "*standard error of estimate*".
- s^2 is a *statistic* - has its own sampling distribution, *etc.* More on this to come.
- Let's see one immediate *application* of s^2 and s .
- Recall sampling distribution for LS estimator, \mathbf{b} :

$$\mathbf{b} \sim N[\boldsymbol{\beta}, \sigma^2(\mathbf{X}'\mathbf{X})^{-1}]$$

- So, $\text{var.}(b_i) = \sigma^2[(\mathbf{X}'\mathbf{X})^{-1}]_{ii}$; σ^2 is *unobservable*.
- If we want to report variability associated with b_i as an estimator of β_i , we need to use an estimator of σ^2 .

- $est. var. (b_i) = s^2[(X'X)^{-1}]_{ii}$.
- $\sqrt{est. var. (b_i)} = s.d. (b_i) = s\{[(X'X)^{-1}]_{ii}\}^{1/2}$.
- We call this the “*standard error*” of b_i .
- This quantity will be very important when it comes to constructing *interval estimates* of our regression coefficients; and when we construct *tests of hypotheses* about these coefficients.

Confidence Intervals & Hypothesis Testing

- So far, we’ve concentrated on “*point*” estimation.
- We will make use of the assumption of *Normally distributed* errors.
- Recall that:

$$\mathbf{b} \sim N[\boldsymbol{\beta}, \sigma^2(X'X)^{-1}]$$

$$b_i \sim N[\beta_i, \sigma^2((X'X)^{-1})_{ii}] \quad ; \quad \text{why still } \textit{Normal}?$$

- So, we can *standardize*:

$$z_i = (b_i - \beta_i) / \sqrt{\sigma^2[(X'X)^{-1}]_{ii}}$$

- But σ^2 is *unknown*, so we can’t use z_i directly to draw inferences about b_i . We must replace σ^2 with an estimator, e.g. s^2

When we replace σ^2 with s^2 in the formula for z_i , z_i is no longer Normally distributed. Instead, the statistic follows a Student-t distribution, and we call it a *t* statistic. That is:

$$\left[\frac{b_i - \beta_i}{s.e. (b_i)} \right] \sim t_{(n-k)}$$

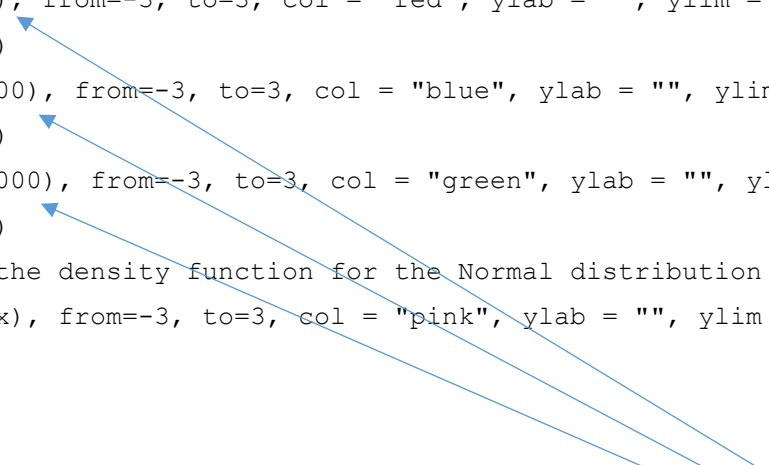
We can use this to construct *confidence intervals* and *test hypotheses* about β_i .

Note: This result used all of our assumptions about the linear regression model – including the assumption of *Normality for the errors*.

Note: The t-distribution becomes the Normal distribution as the sample size grows.

Try running the following R code one line at a time:

```
par(lwd = 3)
#"dt" is the density function for the t-distribution
curve(dt(x,5), from=-3, to=3, col = "red", ylab = "", ylim = c(0, 0.4))
par(new=TRUE)
curve(dt(x,100), from=-3, to=3, col = "blue", ylab = "", ylim = c(0, 0.4))
par(new=TRUE)
curve(dt(x,1000), from=-3, to=3, col = "green", ylab = "", ylim = c(0, 0.4))
par(new=TRUE)
#"dnorm" is the density function for the Normal distribution
curve(dnorm(x), from=-3, to=3, col = "pink", ylab = "", ylim = c(0, 0.4))
```

A diagram consisting of three blue arrows pointing from a rectangular box to the '5', '100', and '1000' values in the R code. The box is located at the bottom right of the code block and contains the text 'Degrees of freedom for the t-distribution (n-k)'. The arrows point from the box to the '5' in 'dt(x,5)', the '100' in 'dt(x,100)', and the '1000' in 'dt(x,1000)'.

Degrees of freedom for the t-distribution (n-k)

```
#Area under the curve, to the left of "-2", t-distribution for various d.o.f.
pt(-2, 5)
pt(-2, 100)
pt(-2, 1000)
#Area under the curve, to the left of "-2", standard Normal distribution
pnorm(-2)
```

```
#To get the 10th percentile from a t-distribution with 15 d.o.f., for example
qt(0.1, 15)
```

Example 1:

$$\hat{y} = 1.4 + 0.2x_2 + 0.6x_3$$

(0.7) (0.05) (1.4)

$$H_0: \beta_2 = 0 \quad \text{vs.} \quad H_A: \beta_2 > 0$$

$$t = \left[\frac{b_2 - \beta_2}{s.e.(b_2)} \right] = \left[\frac{0.2 - 0}{0.05} \right] = 4 \quad ; \quad \text{suppose } n = 20$$

$$t_c(5\%) = 1.74 \quad ; \quad t_c(1\%) = 2.567 \quad ; \quad \text{d.o.f.} = 17$$

$t > t_c \Rightarrow$ *Reject H_0 .*

Degrees of Freedom	90th Percentile	95th Percentile	97.5th Percentile	99th Percentile	99.5th Percentile
1	3.078	6.314	12.706	31.821	63.657
2	1.886	2.920	4.303	6.965	9.925
:	:	:	:	:	:
15	1.341	1.753	2.131	2.602	2.947
16	1.337	1.746	2.120	2.583	2.921
17	1.333	1.740	2.110	2.567	2.898

Example 2:

$$\hat{y} = 1.4 + 0.2x_2 + 0.6x_3$$

(0.7) (0.05) (1.4)

$$H_0: \beta_1 = 1.5 \quad \text{vs.} \quad H_A: \beta_1 \neq 1.5$$

$$t = \left[\frac{b_1 - \beta_1}{s.e.(b_1)} \right] = \left[\frac{1.4 - 1.5}{0.7} \right] = -0.1429 \quad ; \text{ d.o.f.} = 17$$

$$t_c(5\%) = \pm 2.11$$

$$|t| < t_c \Rightarrow \text{Do Not Reject } H_0$$

(Against H_A , at the 5% significance level.)

Example 3:

$$\hat{y} = 1.4 + 0.2x_2 + 0.6x_3$$

(0.7) (0.05) (1.4)

$$H_0: \beta_1 = 1.5 \quad \text{vs.} \quad H_A: \beta_1 < 1.5$$

$$t = \left[\frac{b_1 - \beta_1}{s.e.(b_1)} \right] = \left[\frac{1.4 - 1.5}{0.7} \right] = -0.1429 \quad ; \text{ d.o.f.} = 17$$

$$p\text{-value} = Pr. [t < -0.1429 | H_0 \text{ is True}]$$

$$\text{in R:} \quad \text{pt}(-0.1429, 17)$$

$$p = 0.444$$

What do you conclude?

Some Properties of Tests:

Null Hypothesis (H_0) Alternative Hypothesis (H_A)

Classical hypothesis testing –

- Assume that H_0 is *TRUE*
- Compute value of test statistic using random sample of data
- Determine *distribution* of the test statistic (*when H_0 is true*)
- Check if observed value of test statistic is likely to occur, *if H_0 is true*
- If this event is sufficiently *unlikely*, then **REJECT H_0** (in favour of H_A)

Note:

1. Can never **accept** H_0 . Why not?
2. What constitutes “*unlikely*” – subjective?
3. Two types of errors we might incur with this process

Type I Error: **Reject H_0** when in fact it is **True**

Type II Error: **Do Not Reject H_0** when in fact it is **False**

- $\text{Pr.}[\text{I}] = \alpha =$ Significance level of test = “size” of test
- $\text{Pr.}[\text{II}] = \beta$; say
- Value of β will depend on how H_0 is **False**. Usually, many ways.

Definition:

The “**Power**” of a test is $\text{Pr.}[\text{Reject } H_0 \text{ when it is } \text{False}]$.

So, $\text{Power} = 1 - \text{Pr.}[\text{Do Not Reject } H_0 \mid H_0 \text{ is } \text{False}] = 1 - \beta$.

- As β typically changes, depending on the way that H_0 is false, we usually have a **Power Curve**.
- For a fixed value of α , this curve plots Power against parameter value(s).
- We want our tests to have *high power*.
- We want the power of our tests to *increase* as H_0 becomes *increasingly false*.

Confidence Intervals

We can also use our t-statistic to construct a confidence interval for β_i .

$$Pr. [-t_c \leq t \leq t_c] = (1 - \alpha)$$

$$\Rightarrow Pr. \left[-t_c \leq \left[\frac{b_i - \beta_i}{s.e.(b_i)} \right] \leq t_c \right] = (1 - \alpha)$$

$$\Rightarrow Pr. [-t_c s.e.(b_i) \leq (b_i - \beta_i) \leq t_c s.e.(b_i)] = (1 - \alpha)$$

$$\Rightarrow Pr. [-b_i - t_c s.e.(b_i) \leq (-\beta_i) \leq -b_i + t_c s.e.(b_i)]$$

$$= (1 - \alpha)$$

$$\Rightarrow Pr. [b_i + t_c s.e.(b_i) \geq \beta_i \geq b_i - t_c s.e.(b_i)] = (1 - \alpha)$$

$$\Rightarrow Pr. [b_i - t_c s.e.(b_i) \leq \beta_i \leq b_i + t_c s.e.(b_i)] = (1 - \alpha)$$

Interpretation –

The interval, $[b_i - t_c s.e.(b_i), b_i + t_c s.e.(b_i)]$ is *random*.

The parameter, β_i , is *fixed* (but unknown).

If we were to take a sample of n observations, and construct such an interval, and then repeat this exercise many, many, times, then $100(1 - \alpha)\%$ of such intervals would cover the true value of β_i .

If we just construct an interval, for our *given* sample of data, we'll never know if *this particular* interval covers β_i , or not.

Example 1

$$\hat{y} = 0.3 - 1.4x_2 + 0.7x_3$$

$$(0.1) \quad (1.1) \quad (0.2)$$

Construct a 95% confidence interval for β_1 when $n = 30$.

$$\text{d.o.f.} = (n - k) = 27 \quad ; \quad (\alpha/2) = 0.025$$

$$t_c = \pm 2.052 \quad ; \quad b_1 = 0.3 \quad ; \quad \text{s.e.}(b_1) = 0.1$$

The 95% Confidence Interval is:

$$[b_1 - t_c \text{ s.e.}(b_1) \quad , \quad b_1 + t_c \text{ s.e.}(b_1)]$$

$$\Rightarrow [0.3 - (2.052)(0.1) \quad , \quad 0.3 + (2.052)(0.1)]$$

$$\Rightarrow [0.0948 \quad , \quad 0.5052]$$

Don't forget the units of measurement!

Example 2

$$\hat{y} = 0.3 - 1.4x_2 + 0.7x_3$$

$$(0.1) \quad (1.1) \quad (0.2)$$

Construct a 90% confidence interval for β_2 when $n = 16$.

$$\text{d.o.f.} = (n - k) = 13 \quad ; \quad (\alpha/2) = 0.05$$

$$t_c = \pm 1.771 \quad ; \quad b_2 = -1.4 \quad ; \quad \text{s.e.}(b_2) = 1.1$$

The 95% Confidence Interval is:

$$[b_2 - t_c \text{ s.e.}(b_2) \quad , \quad b_2 + t_c \text{ s.e.}(b_2)]$$

$$\Rightarrow [-1.4 - (1.771)(1.1) \quad , \quad -1.4 + (1.771)(1.1)]$$

$$\Rightarrow [-3.3481 \quad , \quad 0.5481]$$

Questions:

- Why do we construct the interval *symmetrically* about point estimate, b_i ?
- How can we use a Confidence Interval to test hypotheses?
- For instance, in the last Example, can we reject $H_0: \beta_2 = 0$, against a 2-sided alternative hypothesis?