Strength of Evidence Revisited

Grinnell College

April 9, 2025

Suppose we are interested in testing the hypothesis that the true average hallux length for male sharp-shinned hawks is 11mm. To this end, we collected three samples:

- **Sample 1:** $\overline{x}_1 = 12.3178$, $\hat{\sigma} = 5$, n = 25
- **Sample 2:** $\overline{x}_2 = 12.7109$, $\hat{\sigma} = 5$, n = 25
- Sample 2: $\overline{x}_3 = 12.5109$, $\hat{\sigma} = 5$, n = 25

Provide the following additional information:

- 1. What are the critical values associated with 80% and 90% confidence?
- 2. What are the error rates associated with 80% and 90% confidence
- 3. Draw a t-distribution on a sheet of paper and mark where each of these critical values lay (leave plenty of room)

$\mathsf{Sample}\ 1$

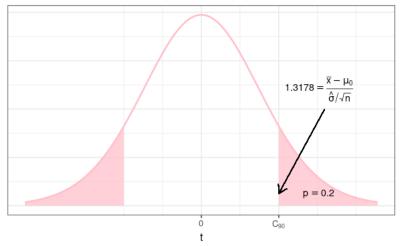
From Sample 1, we find $\overline{x}_1 = 12.3178$, $\hat{\sigma} = 5$, n = 25. The associated *t*-statistic is

$$t = \frac{12.3178 - 11}{5/\sqrt{25}} = 1.3178$$

- ▶ What decision do we make regarding *H*⁰ at 80% confidence?
- What decision do we make regarding H₀ at 90% confidence?
- What is the greatest amount of confidence we could obtain while still rejecting H₀?
- Where does this t-statistic fall relative to the CV you drew in the warm-up?

$\mathsf{Sample}\ 1$

Sample 1

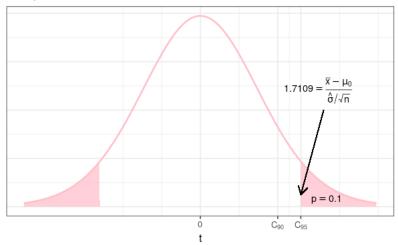


From Sample 1, we find $\overline{x}_2 = 12.7109$, $\hat{\sigma} = 5$, n = 25. The associated *t*-statistic is

$$t = \frac{12.7109 - 11}{5/\sqrt{25}} = 1.7109$$

- ▶ What decision do we make regarding *H*⁰ at 80% confidence?
- What decision do we make regarding H₀ at 90% confidence?
- What is the greatest amount of confidence we could obtain while still rejecting H₀?
- Where does this t-statistic fall relative to the CV you drew in the warm-up?

Sample 2

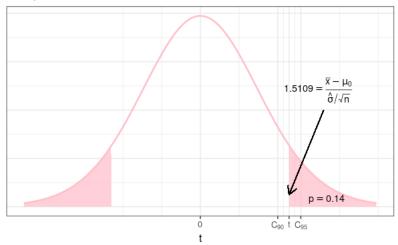


From Sample 1, we find $\overline{x}_3 = 12.5109$, $\hat{\sigma} = 5$, n = 25. The associated *t*-statistic is

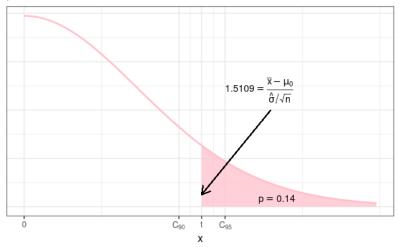
$$t = \frac{12.7109 - 11}{5/\sqrt{25}} = 1.5109$$

- ▶ What decision do we make regarding *H*⁰ at 80% confidence?
- ▶ What decision do we make regarding *H*⁰ at 90% confidence?
- What is the greatest amount of confidence we could obtain while still rejecting H₀?
- Where does this t-statistic fall relative to the CV you drew in the warm-up?

Sample 3



p-value with df = 24



t-distribution

When the null hypothesis is true,

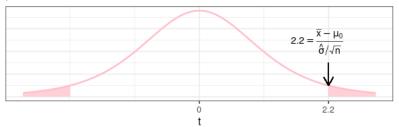
$$t = \frac{\overline{x} - \mu_0}{\hat{\sigma}/\sqrt{n}}$$

follows a *t*-distribution with n-1 degrees of freedom

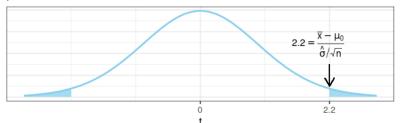
The degrees of freedom tells us, relatively speaking, what values are considered "large"

t = 2.2 may be considered "large" when df = 30 but not when df = 5

p-value with df = 5

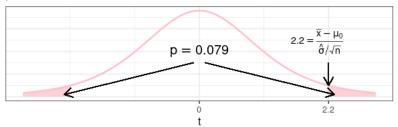


p-value with df = 30

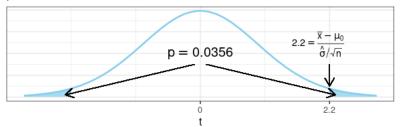


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p-value with df = 5



p-value with df = 30



- Quantifies t-statistic without reference to critical values or degrees of freedom
- Instead allows me to specify an error rate and come to conclusions accordingly

Make note that each value of C corresponds to an error rate, α , which is equal to 1 minus the confidence i.e., a 90% confidence corresponds to a 10% error rate

Likewise, each t statistic corresponds to a p-value. When t>C, it will follow that $p<\alpha$

Suppose we are interested in testing the hypothesis that the true average hallux length for male sharp-shinned hawks is 11mm. To this end, we collected two samples:

- Sample 1: $\overline{x}_1 = 15.61$, $\hat{\sigma} = 6.72$, n = 10
- Sample 2: $\overline{x}_2 = 13.61$, $\hat{\sigma} = 6.12$, n = 25

We might notice in passing that:

- 1. Sample 1 has an observed sample mean that is further away from μ_0
- 2. Sample 2 has more than double the observations as Sample 1
- 3. The observed variability in both samples is about the same

t-statistics and p-values

We can start by constructing *t*-statistics for each of our samples **Sample 1**:

$$t = \frac{15.61 - 11}{6.72/\sqrt{10}} = 2.17$$

Sample 2:

$$t = \frac{13.61 - 11}{6.12/\sqrt{25}} = 2.13$$

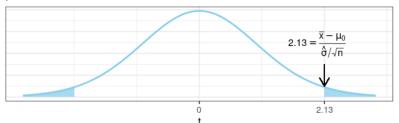
We might conclude that, having the larger *t*-statistic that Sample 1 provides more evidence against the null; however, the *null distribution* for each statistic is different according to its degrees of freedom

p-values

p-value with df = 9

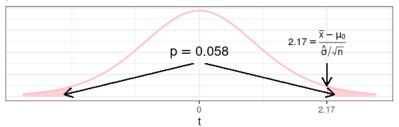


p-value with df = 24

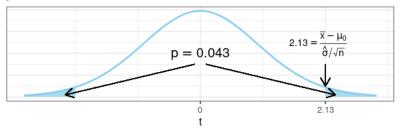


p-values

p-value with df = 9



p-value with df = 24



Suppose we were wishing to test our hypothesis $H_0: \mu = 11$ with each of our two samples with a Type I error rate of $\alpha = 0.05$

Sample 1:

With a *t* statistic of t = 2.17 following a null distribution with df = 9, we find a *p*-value of p = 0.058. Since $p > \alpha$, we fail to reject our null hypothesis

Sample 2:

With a *t* statistic of t = 2.13 following a null distribution with df = 24, we find a *p*-value of p = 0.043. Since $p < \alpha$, we reject our null hypothesis

If we were to construct 95% confidence intervals with our observed data, we would first find critical values for each of our null distributions, according to sample size (note: C_{df} refers to critical value with df degrees of freedom):

$$C_9 = 2.262$$
 $C_{24} = 2.063$

Immediately, we see that for our first sample, $t = 2.17 < C_9$, telling us that our observed data is within the middle 95% and we would fail to reject

Likewise for our second sample, $t = 2.13 > C_{24}$, indicating that we *would* reject

Relationship between CI and α

Now consider the confidence intervals themselves:

Sample 1:

$$15.61 \pm 2.262 \times \left(6.72/\sqrt{10}\right) = (10.8, 20.4)$$

Sample 2:

$$13.61 \pm 2.063 \times \left(6.12/\sqrt{25}\right) = (11.1, 16.2)$$

Here, we see that the null hypothesis $H_0: \mu = 11$ is contained within the 95% confidence interval of Sample 1, indicating that we fail to reject, while it is *not* within the interval for Sample 2, indicating rejection

Hypothesis testing involves formulating statements about our population and then checking the consistency of our hypothesis with observed data

Rather than getting a binary yes/no answer by checking *t*-statistics with a specific critical value, a **p-value** allows us to *quantify* to what extent our observed data is consistent with a null hypothesis

There is a one-to-one relationship between critical values and our Type I error rate, α

Checking that t < C is equivalent to checking if $p < \alpha$