

Unit #7 (b): Hypothesis Testing

9.7.3-9.7.5, 9.9.4



Photo by alyssa teboda on [Unsplash](#)



At the end of this unit, students should be able to:

1. State and interpret hypotheses about differences across groups/populations.
2. Write and compute the test statistic for the a test comparing two population means with large samples (when the standard deviation is known and unknown). Use the test statistic to conduct the test and make a decision (using either a rejection region or a p-value).
3. Distinguish between lower-tail, upper-tail, and two-tailed tests. Reason about when each should be used.
4. Write and compute the test statistic for the two sample t-test. Recognize when this test applies. Use the test statistic to conduct the test and make a decision (using either a rejection region or a p-value).
5. Write and compute the test statistic for the test for proportion differences. Recognize when this test applies. Use the test statistic to conduct the test and make a decision (using either a rejection region or a p-value).
6. Describe the F distribution, including what it is parameterized by, how it arises, and what it is used to test.



Often, researchers are interested in differences with respect to the mean of a continuous variable across different populations or groups.

Example: A coffee company has two different methods for brewing coffee, and the company is interested in knowing whether there is a difference in caffeine levels across brewing methods.



Photo by Jan Huber on Unsplash

First, some theory.

$$\begin{array}{c} | \\ 1 \\ | \\ 1 \\ | \\ 2 \\ | \\ 2 \\ | \\ 2 \end{array} \quad \begin{array}{l} x_1 = 80 \text{ mg} \\ x_2 = 82 \text{ mg} \\ \vdots \\ \vdots \end{array}$$

Let X_1, \dots, X_{n_X} be a large random sample from a population with mean $\mu_X = E(X_i)$ and population variance $\sigma_X^2 = Var(X_i)$.

$$H_0 : \mu_X = \mu_Y \iff \mu_X - \mu_Y = 0$$

Let Y_1, \dots, Y_{n_Y} be a large random sample from a different, independent population with mean $\mu_Y = E(Y_i)$ and population variance $\sigma_Y^2 = Var(Y_i)$.

- $E(\bar{X} - \bar{Y}) = \underbrace{E(\bar{X}) - E(\bar{Y})}_{\text{unbiased est. of } \mu_X - \mu_Y} = \mu_X - \mu_Y$
- $Var(\bar{X} - \bar{Y}) = Var(\bar{X}) + Var(\bar{Y}) \quad \left[\begin{array}{l} \text{assuming} \\ x_i \text{ is ind. of } y_i \end{array} \right]$
$$= \frac{\sigma_X^2}{n_X} + \frac{\sigma_Y^2}{n_Y}$$



Photo by Jan Huber on Unsplash

Consider testing: $H_0 : \mu_X - \mu_Y = \delta_0$ against one of the three alternatives (e.g., $H_1 : \mu_X - \mu_Y \neq \delta_0$).

The test statistic

$$Z = \frac{\bar{X} - \bar{Y} - \delta_0}{\sqrt{\left(\frac{\sigma_X^2}{n_X} + \frac{\sigma_Y^2}{n_Y} \right)}} \sim N(0,1).$$

(Either $n_X > 30$
and $n_Y > 30$
or

$x_i \sim N$
&
 $y_i \sim N$

If $Z = z$ falls within the rejection region (defined by the alternative hypothesis), then we reject H_0 (or have some evidence against H_0). Otherwise, we fail to reject H_0 .



Photo by Jan Huber on Unsplash

Suppose that we didn't know the population variances. *If they are assumed to be roughly equal*, then the test statistic

$$T = \frac{\bar{X} - \bar{Y} - \delta_0}{\sqrt{S_p^2 \left(\frac{1}{n_X} + \frac{1}{n_Y} \right)}} \sim t_{n_X+n_Y-2}, \text{ where}$$

$$S_p^2 = \frac{(n_X - 1)S_X^2 + (n_Y - 1)S_Y^2}{n_x + n_Y - 2} \quad (\text{the "pooled" estimator of } \sigma_X^2 = \sigma_Y^2).$$

If $T = t$ falls within the rejection region (defined by the alternative hypothesis), then we reject H_0 (or have some evidence against H_0). Otherwise, we fail to reject H_0 .



Photo by Jan Huber on Unsplash

Suppose that we didn't know the population variances. *If they are not assumed to be roughly equal*, then the test statistic

$$T = \frac{\bar{X} - \bar{Y} - \delta_0}{\sqrt{\left(\frac{S_X^2}{n_X} + \frac{S_Y^2}{n_Y} \right)}} \text{ approx } t_v$$

Welch's
t-test

$$\left(\frac{S_X^2}{n_X} + \frac{S_Y^2}{n_Y} \right)^2$$

$$\text{where } v = \left(\frac{\left(S_X^2/n_X \right)^2}{n_X - 1} + \frac{\left(S_Y^2/n_Y \right)^2}{n_Y - 1} \right)$$

If $T = t$ falls within the rejection region (defined by the alternative hypothesis), then we reject H_0 (or have some evidence against H_0). Otherwise, we fail to reject H_0 .



Photo by Jan Huber on Unsplash

In R:

```
t.test(x, y, alternative = "two.sided",
var.equal = TRUE, conf.level = 0.9...)
```