

Statistical Methods

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Topics

- 1 Introduction to Hypothesis Testings
- 2 One-Sample t Test for μ
- 3 Type I and II Errors and Their Probabilities

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Objectives

Objectives:

- Explain the meaning of the terms *hypothesis*, *test statistic*, *level of significance*, *rejection region*, *p-value*, and *decision rule*.
- Carry out a one-sample t test for a population mean.
- Distinguish between Type I and Type II errors.
- Know the relationship between the level of significance and the Type I error probability.

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Introduction to Hypothesis Testing

- A ***hypothesis*** is a claim about the value(s) of one or more population parameters, (e.g. μ).
- A ***hypothesis test*** is a statistical method for deciding between two hypotheses:
 - The ***null hypothesis*** (H_0) is the hypothesis we seek to **discredit**, but to which we give the **benefit of the doubt**.
 - The ***alternative hypothesis*** (H_a) is the hypothesis we seek to **substantiate**.

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- The **conclusion** in any hypothesis test will be to either
Reject H_0 or **Fail to Reject H_0** .
- The decision is based on whether a **test statistic** provides compelling evidence against H_0 , ...
... as determined by comparing its value to the sampling distribution it *would* follow if H_0 was true.
- A **decision rule** specifies when the evidence against H_0 is so compelling that H_0 should be rejected.

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- There are **two approaches** to developing a decision rule:
 1. The **rejection region approach**.
 2. The **p -value approach**.

In either case, we first choose a **level of significance** α , which indicates how strong the evidence against H_0 needs to be before we're willing to reject H_0 .

A **smaller** α requires **stronger evidence**.

The most commonly used values for α are **0.01**, **0.05**, and **0.10**.

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- A **rejection region** is the **set** all **test statistic values** for which H_0 should be rejected.
It's chosen in such a way that when H_0 is true, the test statistic will fall into that region just by chance with probability α .

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Decision Rule (RR approach):

Reject H_0 if the test statistic falls in the rejection region.
Fail to reject H_0 otherwise.

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- The **p-value** is a **probability** that answers the question:

"If H_0 was true, what's the chance we'd get a test statistic value that's as contradictory to H_0 (and consistent with H_a) as the one we got?"

Decision Rule (P-value approach):

Reject H_0 if p-value $< \alpha$.

Fail to reject H_0 if p-value $\geq \alpha$.

- We say that a result is **statistically significant** when we reject H_0 .

A statistically significant result is one that isn't likely just due to chance variation.

Steps in Performing a Hypothesis Test:

1. Identify and define the parameter(s) of interest.
2. State the null and alternative hypotheses.
3. Choose a level of significance α .
4. Check any assumptions required for the test.
5. Calculate the test statistic value.
6. Compute the p-value or determine the rejection region.
7. State the conclusion (using the decision rule).

One-Sample t Test for μ (8.3)

- Suppose X_1, X_2, \dots, X_n are a random sample from a population whose (unknown) mean is μ .
- We'll see how to use the sample to decide if μ is different from some **hypothesized value** μ_0 .

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- Because we're seeking to "disprove" the claim that μ is equal to μ_0 , the **null hypothesis** is that it *is* equal to μ_0 .

Null Hypothesis:

$$H_0 : \mu = \mu_0$$

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- The **alternative hypothesis** will depend on what we're trying to "prove":

Alternative Hypothesis: The alternative hypothesis will be one of

1. $H_a : \mu > \mu_0$ (one-sided, upper-tailed)
2. $H_a : \mu < \mu_0$ (one-sided, lower-tailed)
3. $H_a : \mu \neq \mu_0$ (two-sided, two-tailed)

depending on what we're trying to verify using the data.

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One-Sample t Test Statistic:

$$T = \frac{\bar{X} - \mu_0}{S/\sqrt{n}}$$

- T measures how many standard errors \bar{X} is away from μ_0 .
- \bar{X} is an estimator of the unknown population mean μ , so ...
 1. T will be approximately **zero** (most likely) if $\mu = \mu_0$.
 2. It will be **positive** (most likely) if $\mu > \mu_0$.
 3. It will be **negative** (most likely) if $\mu < \mu_0$.

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1. **Large positive** values of T provide **evidence against H_0 in favor of $H_a : \mu > \mu_0$.**
2. **Large negative** values of T provide **evidence against H_0 in favor of $H_a : \mu < \mu_0$.**
3. **Large positive and large negative** values of T provide **evidence against H_0 in favor of $H_a : \mu \neq \mu_0$.**

- If either:
 1. The sample is from a $N(\mu, \sigma)$ population, or
 2. The sample size n is large,

then

$$\frac{\bar{X} - \mu}{S/\sqrt{n}} \sim t(n-1).$$

- It follows that **if H_0 is true** (so $\mu = \mu_0$),

$$\frac{\bar{X} - \mu_0}{S/\sqrt{n}} \sim t(n-1).$$

Sampling Distribution of the Test Statistic Under H_0 :

If T is the one-sample t test statistic, then when

$$H_0 : \mu = \mu_0$$

is true,

$$T \sim t(n-1).$$

- The $t(n-1)$ curve gives us:
 - The **rejection region** as the **extreme $100\alpha\%$ of t values** (in the direction(s) specified by H_a).
 - The **p -value** as the **tail area(s) beyond the observed t value** (in the direction(s) specified by H_a).

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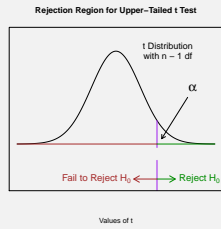
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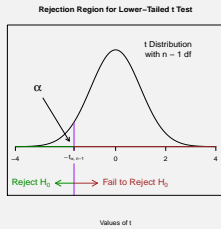
Rejection Region: The **rejection region** is the **set of t values** in the tail of the $t(n - 1)$ curve:

1. To the **right of $t_{\alpha, n - 1}$** if the alternative hypothesis is $H_a : \mu > \mu_0$:



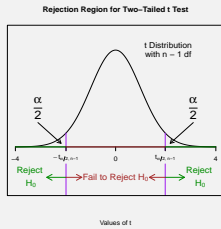
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2. To the **left of $-t_{\alpha, n - 1}$** if the alternative hypothesis is $H_a : \mu < \mu_0$:



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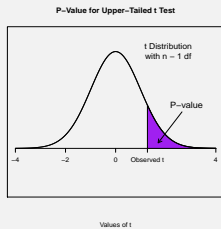
3. To the **left of $-t_{\alpha/2, n - 1}$** and **right of $t_{\alpha/2, n - 1}$** if the alternative hypothesis is $H_a : \mu \neq \mu_0$:



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P-Value: The **p-value** is the **tail area** under the $t(n - 1)$ curve:

1. To the **right of the observed t** if the alternative hypothesis is $H_a : \mu > \mu_0$:



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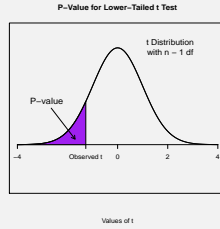
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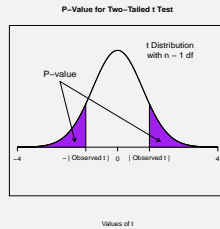
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2. To the **left** of the **observed t** if the alternative hypothesis is $H_a : \mu < \mu_0$:



3. To the **left of $-|t|$ and right of $|t|$** if the alternative hypothesis is $H_a : \mu \neq \mu_0$:



- The rejection region and p-value approaches **always reach the same conclusion.**

(The **p-value** will be less than α if and only if t is in the **rejection region**).

Example

A quality control engineer monitors a machine that puts cereal into boxes.

According to the label, each box is supposed to contain **16 oz** of cereal.

The machine will need to be adjusted if the boxes are systematically being **under-filled** or **over-filled**.

From past experience, the engineer knows that the weight (ounces) of the cereal in a box follows a **normal** distribution.

To decide if the boxes are being **under-filled or overfilled**, the engineer will test the **hypotheses**

$$H_0: \mu = 16$$

$$H_a: \mu \neq 16$$

where μ is the true (unknown) population mean weight.

A random sample of **ten** boxes gives

$$\bar{x} = 16.6 \quad \text{and} \quad s = 0.9.$$

The observed **test statistic** is

$$\begin{aligned} t &= \frac{\bar{x} - \mu_0}{s/\sqrt{n}} \\ &= \frac{16.6 - 16}{0.9/\sqrt{10}} \\ &= \mathbf{2.11}. \end{aligned}$$

Thus the **sample mean** weight, $\bar{x} = 16.6$, is about **2.11 standard errors above 16 ounces**.

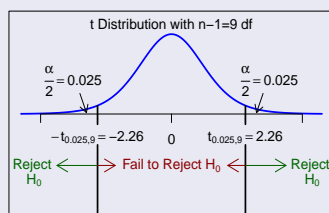
For the **rejection region**, using a **level of significance $\alpha = 0.05$** , the **t critical value** is

$$t_{0.025, 9} = \mathbf{2.262},$$

and so the decision rule is

Reject H_0 if $t < -2.262$ or $t > 2.262$.
 Fail to reject H_0 otherwise.

Rejection Region for Two-Sided t Test



Values of t

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Because the test statistic, $t = 2.11$, is **not** in the rejection region, we **fail to reject** H_0 .

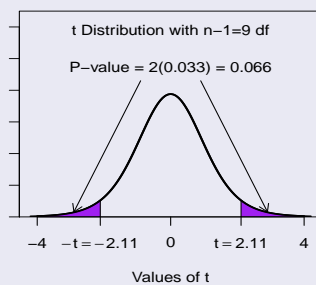
Thus the t value we got is **not** among the most extreme **5%** of values we'd get if the **population mean μ was 16 ounces**.

There's **no statistically significant evidence** that the population mean cereal box weight μ is different from 16 ounces.

The result that the engineer got (by taking a random sample) can be explained by chance variation (sampling error).

The **p-value** is the **probability** that by chance we'd get a t value as far away from zero (in either direction) as $t = 2.11$ if the **population mean μ was 16 oz**.

P-Value for Two-Sided t Test



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From the **two tail** areas of the **sampling distribution** that the test statistic would follow under H_0 (the $t(9)$ distribution), to the **right of 2.11** and **left of -2.11**,

$$\text{p-value} = 2(0.033) = 0.066.$$

Thus we'd get a result like the one we got **6.6%** of the time **even if the population mean μ was 16 ounces.**

Using $\alpha = 0.05$, the **decision rule** is

Reject H_0 if $\text{p-value} < 0.05$.
Fail to reject H_0 if $\text{p-value} \geq 0.05$.

Because $0.066 \geq 0.05$, we **fail to reject H_0 .**

- The next exercise illustrates the fact that using a **smaller α** means we require **stronger evidence** against H_0 before we're willing to reject H_0 .

Exercise

In the last example, if the engineer had used a level of significance $\alpha = 0.10$ instead, would his **conclusion** be any **different**?

What if he used $\alpha = 0.01$?

Data Snooping: Don't Do It

- Choosing a **direction** for a **one-sided H_a** is intended to be a **prediction** of what the data will indicate.
- **Data snooping** refers to waiting until **after you've looked at the data** to decide on a direction for H_a , and then choosing the direction for H_a that best fits what you **already see in the data.**
- Data snooping is "**cheating**" because it results in an **artificially small p-value**, which can lead to mistakenly declaring a spurious result to be real.

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- A **one-sided H_a** should only be used if you have a specific direction in mind **prior** to looking at the data.
 Otherwise, use a **two-sided H_a** .
- The next example shows that **data snooping** can lead to a **p-value** that's **half as large as it should be**.

Exercise

Suppose the engineer who monitors cereal box weights was to "cheat" by **data snooping**, and deciding, **after** and noticing that the sample mean, $\bar{x} = 16.6$, is above the target value **16 oz**, to do a **one-sided, upper-tailed** test of

$$H_0 : \mu = 16$$

$$H_a : \mu > 16$$

- What would the (artificially small) **p-value** be?
- Using $\alpha = 0.05$, as before, would the **conclusion** be **different**?

Type I and II Errors and Their Probabilities

Type I and II Errors

- A **Type I error** occurs when H_0 is **mistakenly rejected** (even though H_0 is true).
- A **Type II error** occurs when H_0 is **mistakenly not rejected** (even though H_a true).
- These are analogous to **false positives** and **false negatives** in medical tests.

| | | True State of Nature | |
|---------------|----------------------|----------------------|------------------|
| | | H_0 | H_a |
| Your Decision | Reject H_0 | Type I Error | Correct Decision |
| | Fail to Reject H_0 | Correct Decision | Type II Error |

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Type I Error Probabilities and the Level of Significance

- It turns out that the **chance** of making a **Type I error** (when H_0 is true) is α , the **level of significance**.
- To see why, consider the **rejection region** approach.
 - The **rejection region** is the most extreme **100 α %** of the sampling distribution that the test statistic would follow if H_0 was true.
 - A **Type I error** occurs when the test statistic falls into the **rejection region** even though H_0 is true.

- Takeaway:
 - In order to reject H_0 when $\alpha = 0.05$, we require that the evidence against H_0 be so strong that it would occur by chance only 5% of the time if H_0 was true.
 - In order to reject H_0 when $\alpha = 0.01$, we require even stronger evidence. We require evidence that would occur by chance only 1% of the time.

- The choice of what value to use for α will depend on the consequences of making a Type I error: if they're serious, choose α to be very small.

Exercise

Let μ denote the true mean radioactivity level (pCi/L) in a certain lake.

The value 5 pCi/L is considered the dividing line between **safe** and **unsafe** water.

To decide whether the water is safe, 50 water specimens are sampled from the lake, and the radioactivity level measured in each specimen.

a) Describe what the **Type I** and **Type II errors** would be (in the context of this problem) for each of the following sets of hypotheses.

$$H_0 : \mu = 5$$

$$H_a : \mu > 5$$

$$H_0 : \mu = 5$$

$$H_a : \mu < 5$$

b) If we were to test the second set of hypotheses, which **level of significance** would you recommend, $\alpha = 0.10$, $\alpha = 0.05$, or $\alpha = 0.01$?

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