

MATH 3307

Lesson 31

Inference for the Mean of a Population

In chapter 7, we discussed one type of inference about a population – the confidence interval. In this chapter we will begin discussion the **significance test**.

H_0 : is the **null hypothesis**. The **null hypothesis** states that there is no effect or change in the population. It is the statement being tested in a test of significance.

This is the accepted value that we are trying to confirm or deny

H_a : is the **alternate hypothesis**. The **alternative hypothesis** describes the effect we suspect is true, in other words, it is the alternative to the “no effect” of the null hypothesis.

Three possible Alternate Hypothesis: Null hypothesis is too large, too small, or incorrect.

Since there are only two hypotheses, there are only two possible decisions: *reject the null hypothesis in favor of the alternative* or *don't reject the null hypothesis*. We will never say that we accept the null hypothesis.

RH_0
 FRH_0

For inference about a population mean:

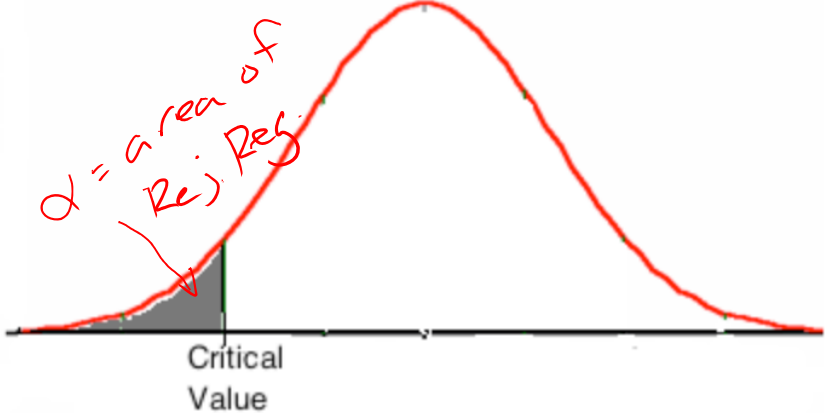
$H_0 : \mu = \mu_0$ where μ_0 represents the given population mean.

$$H_0: \mu = 150$$

↳ Accepted Value

Current Value

For inference about a population mean:

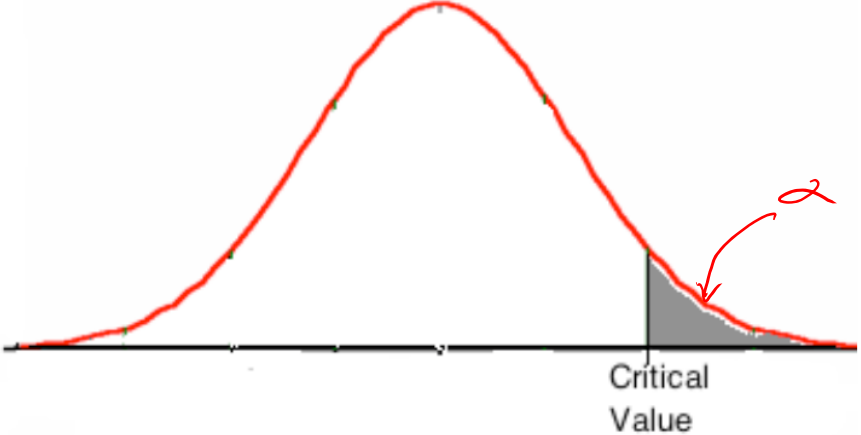
Alternate Hypothesis	Rejection Region
$H_a : \mu < \mu_0$ <p>To find a critical value: qnorm(a) [z test] qt(a,df) [t test]</p>	

If you are given a Significance Level, α , this can be used to determine the critical value and the rejection region. The total area of the rejection region will have the value of α .

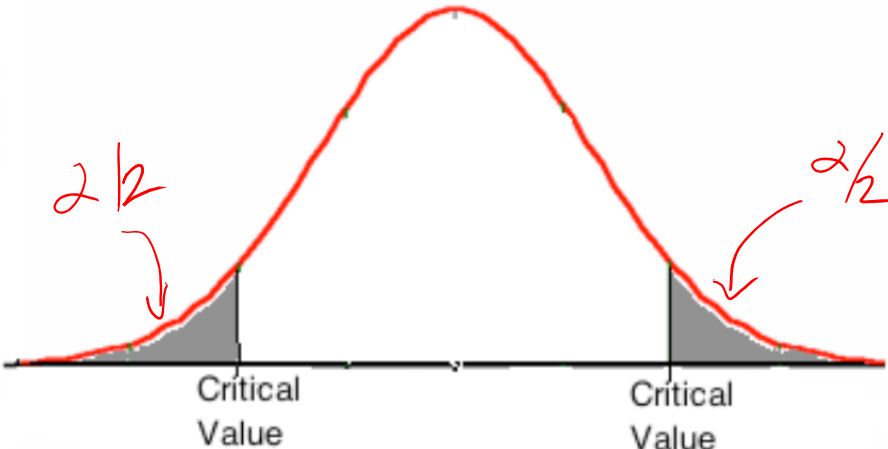
The rejection region is the set of values of the test statistic that will lead to a rejection of the null hypothesis.

The critical value is the boundary of the rejection region.

For inference about a population mean:

Alternate Hypothesis	Rejection Region
<p data-bbox="472 797 701 850">$H_a : \mu > \mu_0$</p> <p data-bbox="279 894 674 932">To find the critical value:</p> <p data-bbox="279 943 600 980">qnorm(1-a) [z test]</p> <p data-bbox="279 992 600 1029">qt(1-a,df) [t test]</p>	 <p data-bbox="1661 1052 1766 1127">Critical Value</p>

For inference about a population mean:

Alternate Hypothesis	Rejection Region
<p data-bbox="472 641 703 698">$H_a : \mu \neq \mu_0$</p> <p data-bbox="237 730 583 771">To find critical values:</p> <ul data-bbox="237 779 457 868" style="list-style-type: none"><li data-bbox="237 779 457 820">$\pm \text{qnorm}(a/2)$<li data-bbox="237 820 457 868">$\pm \text{qt}(a/2, df)$	 <p data-bbox="1092 673 1207 755">$\alpha/2$</p> <p data-bbox="1837 673 1942 755">$\alpha/2$</p> <p data-bbox="1234 901 1344 982">Critical Value</p> <p data-bbox="1669 901 1778 982">Critical Value</p>

The probability, computed assuming that the null hypothesis is true, that the test statistic would take a value as extreme or more extreme than that actually observed is called the ***p*-value** of the test. A result with a small *p*-value is called **statistically significant**. This means that chance alone would rarely produce so extreme a result. We say that a value is **statistically significant** when the *p*-value is as small as, or smaller than, the given significance level, α . If we are not given α , we can interpret the results like this:

- If the *p*-value is less than 1%, we say that there is overwhelming evidence to infer that the alternative hypothesis is true. (We also say that the test is highly significant)
- If the *p*-value is between 1% and 5%, we say that there is strong evidence to infer that the alternative hypothesis is true. (We also say that the test is significant)
- If the *p*-value is between 5% and 10%, we say that there is weak evidence to infer that the alternative hypothesis is true. (We also say that the test not statistically significant)
- If the *p*-value is exceeds 10%, we say that there is no evidence to infer that the alternative hypothesis is true.

To summarize:

if α is given:

$p < \alpha$ means to RH_0

$p > \alpha$ means to FRH_0

if α is not given:

$p < 10\%$ means RH_0 (with varying certainty)

$p > 10\%$ means FRH_0

To find *p*-value:

H_a less than: $\text{pnorm}(z)$ or $\text{pt}(t,df)$

H_a greater than: $1-\text{pnorm}(z)$
or $1-\text{pt}(t,df)$

H_a is not equal: $2*\text{pnorm}(z)$ or
 $2*\text{pt}(t,df)$ {must use negative z,t }

Steps to follow:

When performing a significance test, we follow these steps:

1. Check assumptions.
2. State the null and alternate hypotheses.
3. Graph the rejection region, labeling the critical values.
4. Calculate the test statistic.
5. Find the p -value. If this answer is less than the significance level, α , we can reject the null hypothesis in favor of the alternate.
6. Give your conclusion using the context of the problem. When stating the conclusion you can give results with a confidence of $(1 - \alpha)(100)\%$.

Check the casa calendar: Help with Hypothesis testing PDF. This is an interactive PDF that will guide you through all the steps you need to do to complete your hypothesis test.

Z Test (to calculate the test statistic):

z – test

Assumptions:

1. An SRS of size n from the population.
2. Known population standard deviation, σ .
3. Either a normal population or a large sample ($n \geq 30$).

To compute the z – test statistic, we use the formula:

$$z = \frac{\bar{x} - \mu_0}{\sigma / \sqrt{n}}$$

T Test (to calculate the test statistic):

t – test

Assumptions:

1. An SRS of size n from the population.
2. Unknown population standard deviation.
3. Either a normal population or large sample ($n \geq 30$).

To compute the t – test statistic, we use the formula $t = \frac{\bar{x} - \mu_0}{s / \sqrt{n}}$, where s is the sample standard deviation. The t – test will use $n - 1$ degrees of freedom.

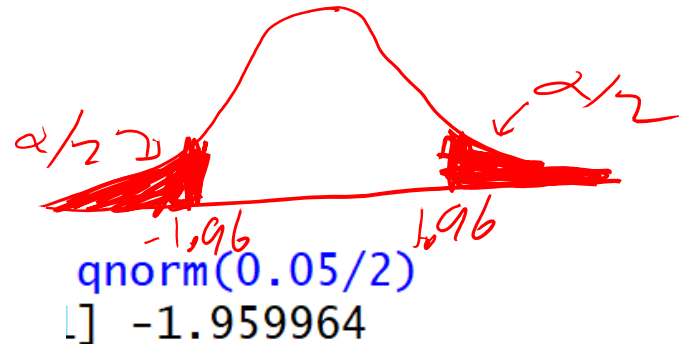
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Example: $H_0: \mu = 18.2$ Population: $\mu = 18.2$
 $H_a: \mu \neq 18.2$ $\sigma = 1.38$ Z-Test

Example 1: A laboratory is asked to evaluate the claim that the amount of the active ingredient in a bug spray is 18.2 grams for a 70-gram bottle with a standard deviation of 1.38 grams. The mean amount of the active ingredient in 40 randomly selected 70-gram bottles of the bug spray is $\bar{x} = 16.828$ grams. Do these analyses indicate that the amount of the active ingredient is different than the original claim at an $\alpha = 0.05$ significance level?

Sample: $n = 40$
 $\bar{x} = 16.828$

$\alpha = .05$



Rej. H_0 : $Z < -1.96$ or $Z > 1.96$

Example:

Example 1: A laboratory is asked to evaluate the claim that the amount of the active ingredient in a bug spray is 18.2 grams for a 70-gram bottle with a standard deviation of 1.38 grams. The mean amount of the active ingredient in 40 randomly selected 70-gram bottles of the bug spray is $\bar{x} = 16.828$ grams. Do these analyses indicate that the amount of the active ingredient is different than the original claim at an $\alpha = 0.05$ significance level?

$$H_0: \mu = 18.2$$

$$H_a: \mu \neq 18.2$$

Population:

$$\mu = 18.2$$

$$\sigma = 1.38$$

$$\bar{x} = 16.828$$

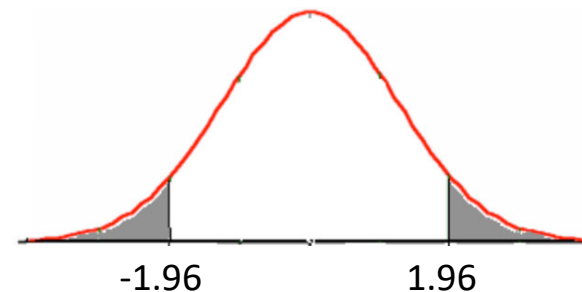
$$\alpha = 0.05$$

Since the population standard deviation is given, use a z-test.

Rejection Region is 2-sides, so the area of one tail is $0.05/2 = 0.025$.

$$qnorm(1-0.025) = 1.96$$

$$qnorm(0.025) = -1.96$$



Example (Continued):

$$\frac{(16.828 - 18.2)}{(1.38 / \sqrt{40})} = -6.287891$$

$$z = \frac{\bar{x} - \mu_0}{\sigma / \sqrt{n}} = \frac{16.828 - 18.2}{1.38 / \sqrt{40}} = -6.288$$

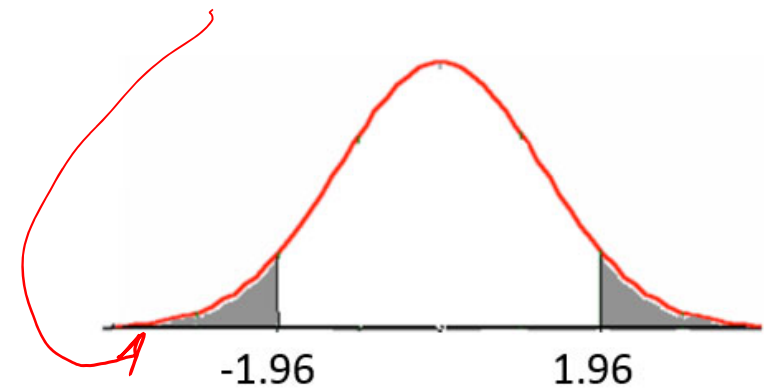
$z = -6.288$ falls within our rejection region

$$(-6.288 < -1.96)$$

$$2 * \text{pnorm}(-6.287891)$$

$$1] 3.218077e-10$$

$R H_0$



P-Value: $P(Z < -6.288) + P(Z > 6.288) = 2P(Z < -6.288) \approx 0$

$$p = 0 < \alpha = .05$$

(Since $p < \alpha$, we can conclude that we can reject the null hypothesis.)

Conclusion: Based on 95% certainty, we can reject the null hypothesis, in favor of saying that the amount of active ingredient is not 18.2 grams per can.

Example: Popper 27

$$\left. \begin{array}{l} \mu = 200 \\ \sigma = 9 \end{array} \right\} z\text{-test}$$

Mr. Murphy is an avid golfer. Suppose he has been using the same golf clubs for quite some time. Based on this experience, he knows that his average distance when hitting a ball with his current driver (the longest-hitting club) under ideal conditions is 200 yards with a standard deviation of 9. After some preliminary swings with a new driver, he obtained the following sample of driving distances:

205	198	220	210	194	201	213	191	211	203
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$$n = 10$$

He feels that the new club does a better job. Do you agree?

$$H_0: \mu = 200$$

$$H_a: \mu > 200$$

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assign("x",c(205,198,220,210,194,201,213,191,211,203))
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```
mean(x)
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[1] 204.6
```

1. What is the sample mean?

a. 200

b. 204.6

c. 190

d. 215

2. What is the population mean? *Given*

a. 200

b. 204.6

c. 190

d. 215

3. Which test statistic can be used?

a. z-test

b. t-test

4. What is our null hypothesis?

a. $\mu = 200$

b. $\mu < 200$

c. $\mu > 200$

d. $\mu \neq 200$

5. What is our alternate hypothesis?

a. $\mu = 200$

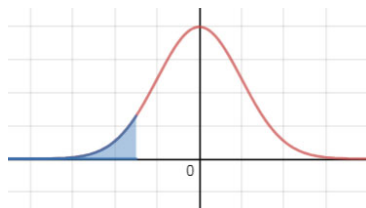
b. $\mu < 200$

c. $\mu > 200$

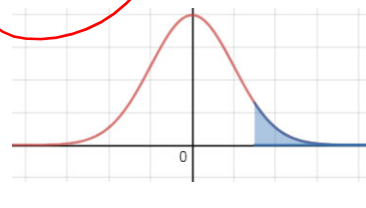
d. $\mu \neq 200$

6. Which of the following is a correct representation of our rejection region?

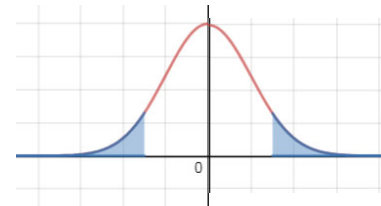
a.



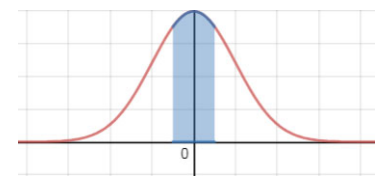
b.



c.



d.



Since we were not provided a significance level (α) in this question, we cannot provide a numeric rejection region. Our conclusions will be based entirely on the p-value.

$$Z = \frac{\bar{X} - \mu}{s/\sqrt{n}}$$

$$\frac{(\text{mean}(x) - 200) / (9 / \sqrt{10})}{1} = 1.616275$$

7. What is the value of the test statistic?

- a. 5.11
- b. 4.85
- c. -1.616
- d. 1.616

8. What is the p-value?

$$1 - \text{pnorm}(1.616275) = 0.05301743$$

- a. 0.0530
- b. 0.9474
- c. 0.016
- d. 0.9804

9. Do we reject the null hypothesis in favor of the alternate hypothesis?

- a. Yes, with overwhelming evidence.
- b. Yes, with strong evidence.
- c. Yes, with weak evidence.
- d. No, we fail to reject the null hypothesis.

$P = 5.3\%$
 $0.05 < P < 0.10$
R_H (Weak Evidence)