

**CONFIDENCE INTERVALS:****Define parameter(s):**

<b>Proportions</b>	<b>Means</b>
<ul style="list-style-type: none"> <li>➤ One sample: <math>p</math> = true <b>proportion</b> of...</li> <li>➤ Two sample: <math>p_1</math> = true <b>proportion</b> of... <math>p_2</math> = true <b>proportion</b> of...</li> </ul>	<ul style="list-style-type: none"> <li>➤ One sample: <math>\mu</math> = true <b>mean</b> of...</li> <li>➤ Two sample: <math>\mu_1</math> = true <b>mean</b> of... <math>\mu_2</math> = true <b>mean</b> of...</li> </ul>

**Name the procedure:**

<b>Proportions</b>	<b>Means</b>
<ul style="list-style-type: none"> <li>➤ One-sample z-interval for <math>p</math></li> <li>➤ Two-sample z-interval for <math>p_1 - p_2</math></li> </ul>	<ul style="list-style-type: none"> <li>➤ One-sample t-interval for <math>\mu</math></li> <li>➤ Two-sample t-interval for <math>\mu_1 - \mu_2</math></li> </ul>

**Conditions:**

<b>Proportions</b>	<b>Means</b>
<ul style="list-style-type: none"> <li>➤ One sample:               <ol style="list-style-type: none"> <li>1) Random sample</li> <li>2) Independent if <math>n \leq 10\%N</math></li> <li>3) App Normal if <math>n\hat{p} \geq 10</math> and <math>n(1 - \hat{p}) \geq 10</math></li> </ol> </li> <li>➤ Two sample:               <ol style="list-style-type: none"> <li>1) Random samples or Assignment</li> <li>2) Independent if <math>n_1 \leq 10\%N</math> &amp; <math>n_2 \leq 10\%N</math></li> <li>3) Approximately Normal if BOTH:                   <ul style="list-style-type: none"> <li><math>n_1\hat{p}_1 \geq 10</math> and <math>n_1(1 - \hat{p}_1) \geq 10</math> &amp;</li> <li><math>n_2\hat{p}_2 \geq 10</math> and <math>n_2(1 - \hat{p}_2) \geq 10</math></li> </ul> </li> </ol> </li> </ul>	<ul style="list-style-type: none"> <li>➤ One sample:               <ol style="list-style-type: none"> <li>1) Random sample</li> <li>2) Independent if <math>n \leq 10\%N</math></li> <li>3) Approximately Normal if:                   <ul style="list-style-type: none"> <li>➤ Population Normal</li> <li>➤ <math>n \geq 30</math> by CLT</li> <li>➤ Sample shows no strong skewness and outliers</li> </ul> </li> </ol> </li> <li>➤ Two sample:               <ol style="list-style-type: none"> <li>1) Random samples or Assignment</li> <li>2) Independent if <math>n_1 \leq 10\%N</math> &amp; <math>n_2 \leq 10\%N</math></li> <li>3) Approximately Normal if:                   <ul style="list-style-type: none"> <li>➤ BOTH Populations Normal</li> <li>➤ BOTH <math>n_1 \geq 30</math> &amp; <math>n_2 \geq 30</math> by CLT</li> <li>➤ BOTH Samples shows no strong skewness and outliers</li> </ul> </li> </ol> </li> </ul>

**Interpret a confidence interval:**

<b>Proportions</b>	<b>Means</b>
<ul style="list-style-type: none"> <li>➤ One sample: We are _____% confident that the interval from _____ to _____ captures the true <b>proportion</b> of...</li> <li>➤ Two sample: We are _____% confident that the interval from _____ to _____ captures the true <b>difference in proportions</b> of...</li> </ul>	<ul style="list-style-type: none"> <li>➤ One sample: We are _____% confident that the interval from _____ to _____ captures the true <b>mean</b> of...</li> <li>➤ Two sample: We are _____% confident that the interval from _____ to _____ captures the true <b>difference in means</b> of...</li> </ul>

**Interpret a confidence level:** In the long run, \_\_\_\_\_ % of intervals generated capture the population proportion/mean of...

## SIGNIFICANCE TESTS:

**Write hypotheses (including defining parameters):**

Proportions	Means
<ul style="list-style-type: none"> <li>➤ One sample:  <math>H_0: p =</math>  <math>H_a: p \neq, &lt;, \text{ or } &gt;</math>  <math>p =</math> true <b>proportion</b> of...</li> <li>➤ Two sample:  <math>H_0: p_1 = p_2</math>  <math>H_a: \neq, &lt;, \text{ or } &gt;</math>  <math>p_1 =</math> true <b>proportion</b> of...  <math>p_2 =</math> true <b>proportion</b> of...</li> </ul>	<ul style="list-style-type: none"> <li>➤ One sample:  <math>H_0: \mu =</math>  <math>H_a: \mu \neq, &lt;, \text{ or } &gt;</math>  <math>\mu =</math> true <b>mean</b> of...</li> <li>➤ Two sample:  <math>H_0: \mu_1 = \mu_2</math>  <math>H_a: \mu \neq, &lt;, \text{ or } &gt;</math>  <math>\mu_1 =</math> true <b>mean</b> of...  <math>\mu_2 =</math> true <b>mean</b> of...</li> </ul>

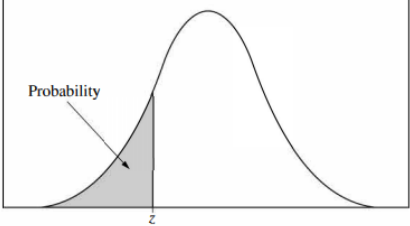
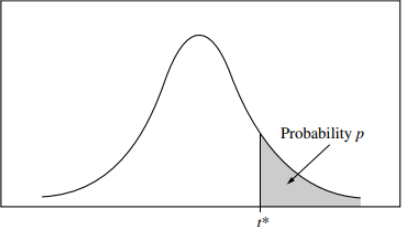
**Name the procedure:**

Proportions	Means
<ul style="list-style-type: none"> <li>➤ One-sample z-test for <math>p</math></li> <li>➤ Two-sample z-test for <math>p_1 - p_2</math></li> </ul>	<ul style="list-style-type: none"> <li>➤ One-sample t-test for <math>\mu</math></li> <li>➤ Two-sample t-test for <math>\mu_1 - \mu_2</math></li> <li>➤ Paired t-test for <math>\mu_d</math></li> </ul>

**Conditions:**

Proportions	Means
<ul style="list-style-type: none"> <li>➤ One sample:               <ol style="list-style-type: none"> <li>1) Random sample</li> <li>2) Independent if <math>n \leq 10\%N</math></li> <li>3) App Normal if <math>np_0 \geq 10</math> and <math>n(1 - p_0) \geq 10</math></li> </ol> </li> <li>➤ Two sample:               <ol style="list-style-type: none"> <li>1) Random samples or Assignment</li> <li>2) Independent if <math>n_1 \leq 10\%N</math> &amp; <math>n_2 \leq 10\%N</math></li> <li>3) Approximately Normal if BOTH:  <math>n_1 \hat{p}_c \geq 10</math> and <math>n_1(1 - \hat{p}_c) \geq 10</math> &amp;  <math>n_2 \hat{p}_c \geq 10</math> and <math>n_2(1 - \hat{p}_c) \geq 10</math></li> </ol> </li> </ul>	<ul style="list-style-type: none"> <li>➤ One sample:               <ol style="list-style-type: none"> <li>1) Random sample</li> <li>2) Independent if <math>n \leq 10\%N</math></li> <li>3) Approximately Normal if:                   <ul style="list-style-type: none"> <li>➤ Population Normal</li> <li>➤ <math>n \geq 30</math> by CLT</li> <li>➤ Sample shows no strong skewness and outliers</li> </ul> </li> </ol> </li> <li>➤ Two sample:               <ol style="list-style-type: none"> <li>1) Random samples or Assignment</li> <li>2) Independent if <math>n_1 \leq 10\%N</math> &amp; <math>n_2 \leq 10\%N</math></li> <li>3) Approximately Normal if:                   <ul style="list-style-type: none"> <li>➤ BOTH Populations Normal</li> <li>➤ BOTH <math>n_1 \geq 30</math> &amp; <math>n_2 \geq 30</math> by CLT</li> <li>➤ BOTH Samples shows no strong skewness and outliers</li> </ul> </li> </ol> </li> </ul>

**Use a test statistic to calculate a P-value:**

Proportions	Means
<p>Test statistic: <math>z</math></p> <p><math>2^{\text{nd}} \rightarrow \text{VARS} \rightarrow</math></p> <div style="border: 1px solid black; padding: 5px; width: fit-content;"> <code>normalcdf</code>            lower:            upper:  <math>\mu: 0</math>  <math>\sigma: 1</math>            Paste         </div> 	<p>Test statistic: <math>t^*</math></p> <p><math>2^{\text{nd}} \rightarrow \text{VARS} \rightarrow</math></p> <div style="border: 1px solid black; padding: 5px; width: fit-content;"> <code>tcdf</code>            lower:            upper:            df:            Paste         </div> <p>*df = <math>n - 1</math></p> 

## Make a conclusion:

Proportions	Means
<p>➤ One sample: p-value <math>\leq \alpha</math>, reject <math>H_0</math>. We have convincing evidence that the true <b>proportion</b> of...(H<sub>a</sub>)</p> <p>p-value <math>&gt; \alpha</math>, fail to reject <math>H_0</math>. We do not have convincing evidence that the true <b>proportion</b> of...(H<sub>a</sub>)</p> <p>➤ Two sample: p-value <math>\leq \alpha</math>, reject <math>H_0</math>. We have convincing evidence that the true <b>difference in proportions</b> of...(H<sub>a</sub>)</p> <p>p-value <math>&gt; \alpha</math>, fail to reject <math>H_0</math>. We do not have convincing evidence that the true <b>difference in proportions</b> of...(H<sub>a</sub>)</p>	<p>➤ One sample: p-value <math>\leq \alpha</math>, reject <math>H_0</math>. We have convincing evidence that the true <b>mean</b> of...(H<sub>a</sub>)</p> <p>p-value <math>&gt; \alpha</math>, fail to reject <math>H_0</math>. We do not have convincing evidence that the true <b>mean</b> of...(H<sub>a</sub>)</p> <p>➤ Two sample: p-value <math>\leq \alpha</math>, reject <math>H_0</math>. We have convincing evidence that the true <b>difference in means</b> of...(H<sub>a</sub>)</p> <p>p-value <math>&gt; \alpha</math>, fail to reject <math>H_0</math>. We do not have convincing evidence that the true <b>difference in means</b> of...(H<sub>a</sub>)</p>

**Interpret a P-value:** Assuming the  $H_0$  (in context) is true, there is a (*P-value*) probability of getting a sample proportion/mean as extreme or more extreme than (*sample proportion/mean*) by chance alone.

## Based on a confidence interval, do we have convincing evidence for a claim?

- One sample:
- \*Compare the claim to the interval. Is the value of the claim inside or outside the interval?
  - Example: claim: different than 14, interval (11, 13) → *Because the value 14 is not a plausible value found within the confidence interval, we have convincing evidence that...is different than 14.*
  - Example: claim: less than 12, interval (11, 13) → *Because the value 12 is a plausible value found within the confidence interval, we do not have convincing evidence that...less than 12.*
- Two sample:
- Because the interval includes 0 as a plausible value for  $p_1 - p_2 / \mu_1 - \mu_2$ , we do not have convincing evidence that...
  - Because the interval does not contain 0, we have convincing evidence that...

## WHY do we CHECK Conditions?

- **Random:**
- **Random sample:** Generalize the result to the population from which the sample was selected.
  - **Random assignment:** Create two groups that are roughly equivalent to allow for a cause-and-effect conclusion between the two groups.
- **10% Condition:** (Only used if sampling without replacement)
- The sampling is without replacement and thus “dependent.” With less than 10% of the population ( $n < 10\%N$ ), it creates an “essentially independent” environment and makes our standard error valid.
- **Normal/Large Counts:**
- Guarantees that the sampling distribution will be approximately normal.
    - **Proportions:**  $n\hat{p} \geq 10$  and  $n(1 - \hat{p}) \geq 10$   
\*This allows us to use the normal approximation to estimate the p-value of the test.
    - **Means:** Population is normal,  $n \geq 30$  by CLT, or sample shows no strong skewness and outliers  
\*This allows us to use a t distribution to estimate the p-value of the test.