

KEY POINTS

chapter 10

Type I error (false positive)

occurs when you incorrectly reject a true null hypothesis,

Type II error (false negative)

happens when you fail to reject a false null hypothesis.

P = population proportion

σ = population standard deviation

μ = mean

$p < \alpha$ = "is sufficient" = you reject

$p > \alpha$ = "is NOT sufficient" = you DON'T reject

What parameter is addressed in the hypothesis?

Proportion, p

Mean, μ

Provided $np_0(1 - p_0) \geq 10$, use the normal distribution with

$$z_0 = \frac{\hat{p} - p_0}{\sqrt{\frac{p_0(1 - p_0)}{n}}} \text{ where } \hat{p} = \frac{x}{n} \quad \text{Section 10.2}$$

Provided the sample size is greater than 30 or the data come from a population that is normally distributed, use Student's t -distribution with $n - 1$ degrees of freedom with

$$t_0 = \frac{\bar{x} - \mu_0}{\frac{s}{\sqrt{n}}} \quad \text{Section 10.3}$$

Population Proportion

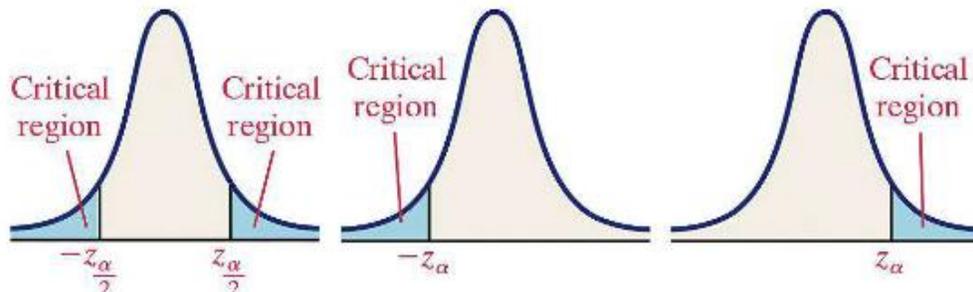
Two-Tailed	Left-Tailed	Right-Tailed
$H_0: p = p_0$	$H_0: p = p_0$	$H_0: p = p_0$
$H_1: p \neq p_0$	$H_1: p < p_0$	$H_1: p > p_0$

Critical value

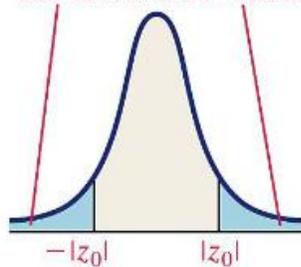
$$-z_{\frac{\alpha}{2}} \text{ and } z_{\frac{\alpha}{2}}$$

$$-z_{\alpha}$$

$$z_{\alpha}$$

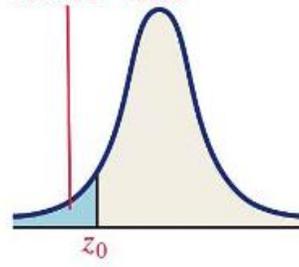


The sum of the area in the tails is the P -value



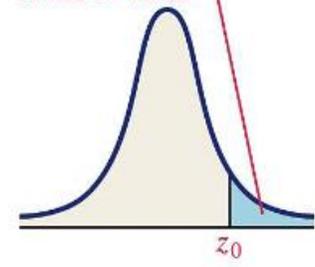
$$P\text{-value} = 2P(z > |z_0|)$$

The area left of z_0 is the P -value



$$P\text{-value} = P(z < z_0)$$

The area right of z_0 is the P -value



$$P\text{-value} = P(z > z_0)$$

- the sample is obtained by simple random sampling or the data result from a randomized experiment.
- $np_0(1 - p_0) \geq 10$.
- the sampled values are independent of each other. That is, the sample size is less than 5% of the population size.

Population Mean

$p < \alpha$ = "is sufficient" = you reject

$p > \alpha$ = "is NOT sufficient" = you DON'T reject

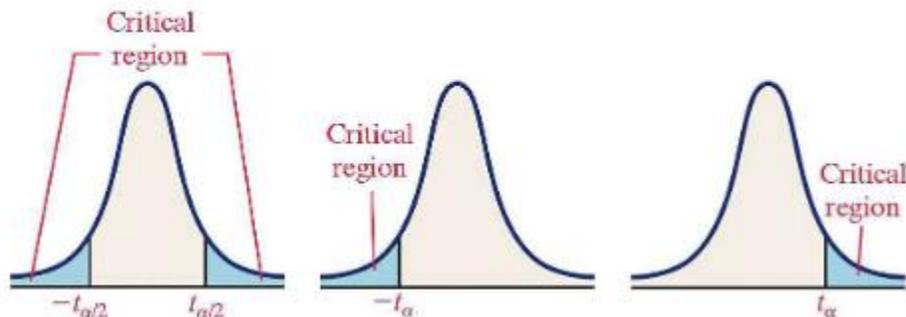
Two-Tailed	Left-Tailed	Right-Tailed
$H_0: \mu = \mu_0$	$H_0: \mu = \mu_0$	$H_0: \mu = \mu_0$
$H_1: \mu \neq \mu_0$	$H_1: \mu < \mu_0$	$H_1: \mu > \mu_0$

Critical value(s)

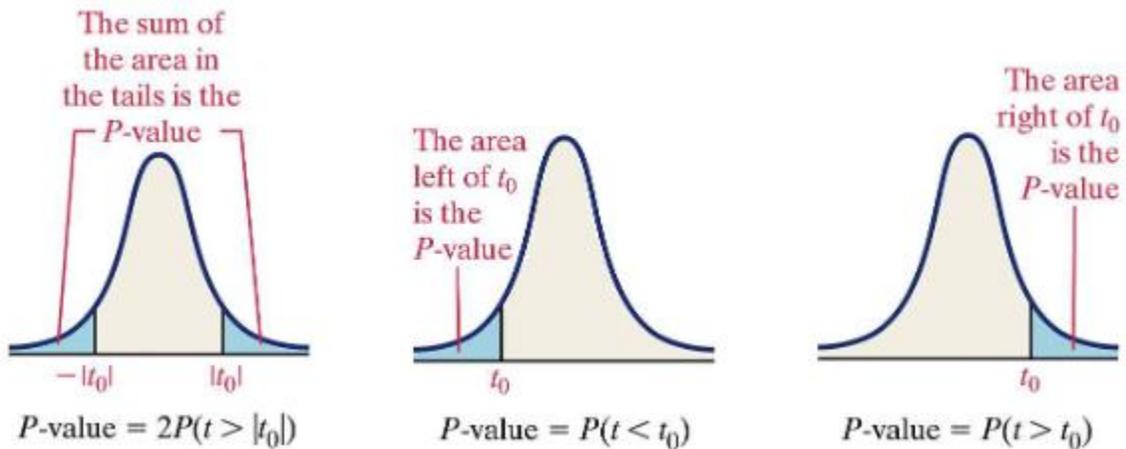
$-t_{\frac{\alpha}{2}}$ and $t_{\frac{\alpha}{2}}$

$-t_{\alpha}$

t_{α}



If **t-value** falls in between the critical values on a number line, **REJECT & IS SUFFICIENT**



- the sample is obtained using simple random sampling or from a randomized experiment.
- the sample has no outliers and the population from which the sample is drawn is normally distributed, or the sample size, n , is large ($n \geq 30$).
- the sampled values are independent of each other. That is, the sample size is less than 5% of the population size.

P-value is the likelihood or probability that a sample will result in a statistic such as the one obtained if the statement in the null hypothesis is true.

BOTTOM LINE IS the lower the P-value, the stronger the evidence against the statement in the null hypothesis.

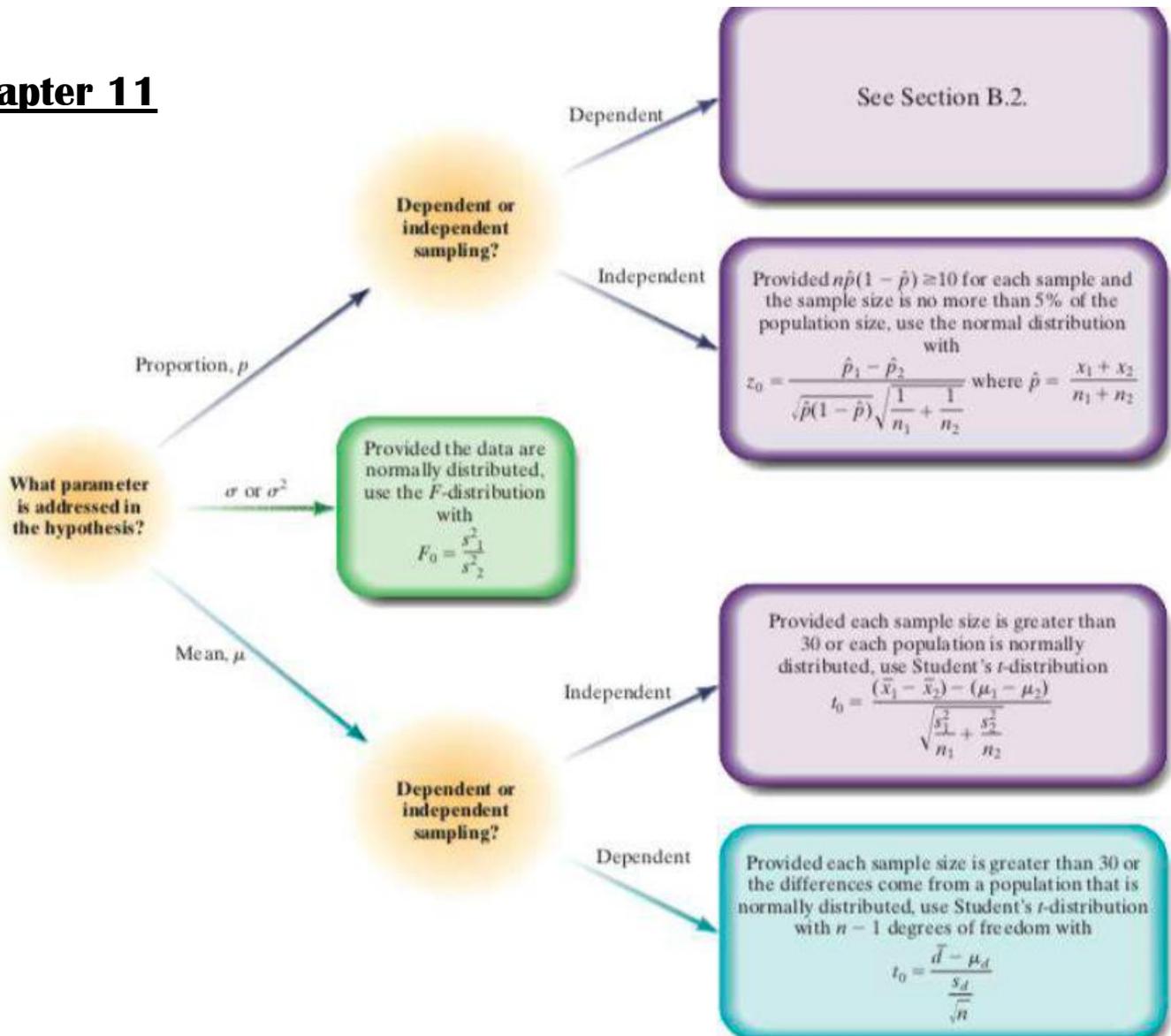
Confidence Interval Critical Values, $z_{\alpha/2}$

Level of Confidence	Critical Value, $z_{\alpha/2}$
0.90 or 90%	1.645
0.95 or 95%	1.96
0.98 or 98%	2.33
0.99 or 99%	2.575

(c) Hypothesis Testing Critical Values

Level of Significance, α	Left-Tailed	Right-Tailed	Two-Tailed
0.10	-1.28	1.28	± 1.645
0.05	-1.645	1.645	± 1.96
0.01	-2.33	2.33	± 2.575

chapter 11



A sampling method is **independent** when an individual selected for one sample does not dictate which individual is to be in a second sample. A sampling method is **dependent** when an individual selected to be in one sample is used to determine the individual in the second sample. Dependent samples are often referred to as **matched-pairs** samples. It is possible for an individual to be matched against him or herself.

- the samples are independently obtained using simple random sampling or through a completely randomized experiment with two levels of treatment,
- $n_1 \hat{p}_1 (1 - \hat{p}_1) \geq 10$ and $n_2 \hat{p}_2 (1 - \hat{p}_2) \geq 10$, and
- $n_1 \leq 0.05N_1$ and $n_2 \leq 0.05N_2$ (the sample size is no more than 5% of the population size); this requirement ensures the independence necessary for a binomial experiment.

Two-Tailed	Left-Tailed	Right-Tailed
$H_0: p_1 = p_2$	$H_0: p_1 = p_2$	$H_0: p_1 = p_2$
$H_1: p_1 \neq p_2$	$H_1: p_1 < p_2$	$H_1: p_1 > p_2$

Note: p_1 is the population proportion for population 1, and p_2 is the population proportion for population 2.

Two-Tailed	Left-Tailed	Right-Tailed
If $z_0 < -z_{\frac{\alpha}{2}}$ or $z_0 > z_{\frac{\alpha}{2}}$, reject the null hypothesis.	If $z_0 < -z_{\alpha}$, reject the null hypothesis.	If $z_0 > z_{\alpha}$, reject the null hypothesis.

for large sample size

- the sample is obtained using simple random sampling or the data result from a matched-pairs design experiment.
- the sample data are dependent (matched pairs).
- the differences are normally distributed with no outliers or the sample size, n , is large ($n \geq 30$).
- the sampled values are independent (sample size is no more than 5% of population size).

Two-Tailed	Left-Tailed	Right-Tailed
$H_0: \mu_d = 0$	$H_0: \mu_d = 0$	$H_0: \mu_d = 0$
$H_1: \mu_d \neq 0$	$H_1: \mu_d < 0$	$H_1: \mu_d > 0$

Two-Tailed	Left-Tailed	Right-Tailed
If $t_0 < -t_\alpha$ or	If $t_0 < -t_\alpha$, reject	If $t_0 > t_\alpha$, reject the
t_0		nu
<p>Suppose that a simple random sample of size n_1 is taken from a population with unknown mean μ_1 and unknown standard deviation σ_1. In addition, a simple random sample of size n_2 is taken from a population with unknown mean μ_2 and unknown standard deviation σ_2.</p>		

Two-Tailed	Left-Tailed	Right-Tailed
$H_0: \mu_1 = \mu_2$	$H_0: \mu_1 = \mu_2$	$H_0: \mu_1 = \mu_2$
$H_1: \mu_1 \neq \mu_2$	$H_1: \mu_1 < \mu_2$	$H_1: \mu_1 > \mu_2$

Two-Tailed	Left-Tailed	Right-Tailed
If $t_0 < -t_{\frac{\alpha}{2}}$ or $t_0 > t_{\frac{\alpha}{2}}$, reject the null hypothesis.	If $t_0 < -t_\alpha$, reject the null hypothesis.	If $t_0 > t_\alpha$, reject the null hypothesis.