

# Lecture 20

## Review

*STAT 8010 Statistical Methods I*  
March 31, 2020

Inference for One  
Population Mean

Inference for Two  
Population Means

Inferences for Matched  
Pairs

ANOVA

Multiple Comparisons  
& Linear Contrasts

RCBD

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# Agenda

- 1 Inference for One Population Mean
- 2 Inference for Two Population Means
- 3 Inferences for Matched Pairs
- 4 ANOVA
- 5 Multiple Comparisons & Linear Contrasts
- 6 RCBD

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# Inference for One Population Mean

## Inferences for One Population Mean $\mu$

**Goal:** To infer  $\mu = \mathbb{E}(X)$  from a random sample  $\{X_1, X_2, \dots, X_n\}$

- Point estimation:

$$\bar{X}_n = \frac{\sum_{i=1}^n X_i}{n}$$

- Interval Estimation:  $100 \times (1 - \alpha)\%$  Confidence Interval (CI)

- $\sigma = \sqrt{\text{Var}(X)}$  is known:

$$\left( \bar{X}_n - z_{\alpha/2} \frac{\sigma}{\sqrt{n}}, \bar{X}_n + z_{\alpha/2} \frac{\sigma}{\sqrt{n}} \right)$$

- $\sigma$  is unknown:

$$\left( \bar{X}_n - t_{\alpha/2, df=n-1} \frac{s}{\sqrt{n}}, \bar{X}_n + t_{\alpha/2, df=n-1} \frac{s}{\sqrt{n}} \right)$$

## Assumptions

- Data is a random sample from the population
- i) sample size  $n$  is sufficiently large (e.g.,  $n > 30$ ) or the population  $X$  follows a normal distribution

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# Margin of Error & Sample Size Calculation

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- Margin of error:

$$z_{\alpha/2} \frac{\sigma}{\sqrt{n}} \quad \text{if } \sigma \text{ known}$$

$$t_{\alpha/2, df=n-1} \frac{s}{\sqrt{n}} \quad \text{if } \sigma \text{ unknown}$$

$\Rightarrow$  CI for  $\mu = \bar{X}_n \pm \text{margin of error}$

- Sample size determination:

$$n = \left( \frac{z_{\alpha/2} \times \sigma}{\text{margin of error}} \right)^2,$$

if  $\sigma$  is given

- 1 State the null and alternative hypotheses:

$$H_0 : \mu = \mu_0 \text{ vs. } H_a : \mu > \text{ or } \neq \text{ or } < \mu_0$$

- 2 Compute the test statistic:

$$z_{obs} = \frac{\bar{X}_n - \mu_0}{\sigma/\sqrt{n}}, \quad \sigma \text{ known}; \quad t_{obs} = \frac{\bar{X}_n - \mu_0}{s/\sqrt{n}}, \quad \sigma \text{ unknown}$$

- 3 Make the decision of the test:

Rejection Region/ P-Value Methods

- 4 Draw the conclusion of the test:

We (do/do not) have enough statistical evidence to conclude that ( $H_a$  in words) at  $\alpha$  significant level.

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## Type I, II Error & Power

True State	Decision	
	Reject $H_0$	Fail to reject $H_0$
$H_0$ is true	Type I error	Correct
$H_0$ is false	Correct	Type II error

- Type I error:  $\mathbb{P}(\text{Reject } H_0 | H_0 \text{ is true}) = \alpha$
- Type II error:  $\mathbb{P}(\text{Fail to reject } H_0 | H_0 \text{ is false}) = \beta$
- The power (PWR):  $\mathbb{P}(\text{Reject } H_0 | H_0 \text{ is false}) = 1 - \beta$ .

$$\Rightarrow \text{PWR}(\mu_a) = 1 - \beta(\mu_a) = 1 - \mathbb{P}\left(z^* \leq z_\alpha - \frac{|\mu_0 - \mu_a|}{\sigma/\sqrt{n}}\right)$$

# Duality of Hypothesis Test with Confidence Interval

There is an interesting relationship between CIs and hypothesis tests. If  $H_0 = \mu_0$  is rejected with significance level  $\alpha$  then the corresponding confidence interval does not contain  $\mu_0$  with the confidence level  $(1 - \alpha)$ , and vice versa

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Hypothesis testing at $\alpha$ level	$(1 - \alpha) \times 100\%$ CI
$H_0 : \mu = \mu_0$ vs. $H_a : \mu \neq \mu_0$	$\bar{X} \pm t_{\alpha/2, n-1} s / \sqrt{n}$
$H_0 : \mu = \mu_0$ vs. $H_a : \mu > \mu_0$	$(\bar{X} - t_{\alpha, n-1} s / \sqrt{n}, \infty)$
$H_0 : \mu = \mu_0$ vs. $H_a : \mu < \mu_0$	$(-\infty, \bar{X} + t_{\alpha, n-1} s / \sqrt{n})$



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# Inferences for Two Population Means

## Statistical Inference for $\mu_1 - \mu_2$

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- **Assumptions:** Populations  $X_1$  and  $X_2$  both follow a normal distribution (or their sample sizes are large enough); Data are random sample from their population
- Point estimation:  $\bar{X}_1 - \bar{X}_2$
- Interval estimation:

$$\bar{X}_1 - \bar{X}_2 \pm \text{margin of error},$$

where margin of error =

$$t_{\alpha/2, df^*} \sqrt{\frac{s_1^2}{n_1} + \frac{s_2^2}{n_2}}, \quad df^* = \frac{\left(\frac{s_1^2}{n_1} + \frac{s_2^2}{n_2}\right)^2}{\frac{(s_1^2/n_1)^2}{n_1-1} + \frac{(s_2^2/n_2)^2}{n_2-1}}$$

When  $s_1$  and  $s_2$  “similar enough”, we replace  $\sqrt{\frac{s_1^2}{n_1} + \frac{s_2^2}{n_2}}$  by

$$s_p \sqrt{\frac{1}{n_1} + \frac{1}{n_2}}, \text{ where } s_p = \sqrt{\frac{(n_1-1)s_1^2 + (n_2-1)s_2^2}{n_1+n_2-2}}, df = n_1 + n_2 - 2$$

## Hypothesis Testing for $\mu_1 - \mu_2$

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- State the null and alternative hypotheses:
  - Upper-tailed test:  $H_0 : \mu_1 - \mu_2 = 0$  vs.  $H_a : \mu_1 - \mu_2 > 0$
  - Lower-tailed test:  $H_0 : \mu_1 - \mu_2 = 0$  vs.  $H_a : \mu_1 - \mu_2 < 0$
  - Two-tailed test:  $H_0 : \mu_1 - \mu_2 = 0$  vs.  $H_a : \mu_1 - \mu_2 \neq 0$
- Compute the test statistic:

$$t_{obs} = \frac{\frac{\bar{X}_1 - \bar{X}_2}{s_p \sqrt{\frac{1}{n_1} + \frac{1}{n_2}}}, \quad \sigma_1 = \sigma_2}{\frac{\bar{X}_1 - \bar{X}_2}{\sqrt{\frac{s_1^2}{n_1} + \frac{s_2^2}{n_2}}}, \quad \sigma_1 \neq \sigma_2}$$

- Make the decision of the test:  
Rejection Region/ P-Value Methods
- Draw the conclusion of the test

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# Inferences for Matched Pairs

- **When to use:** before/after study, pairing subjects, study on twins, etc
- $H_0 : \mu_{diff} = 0$  vs.  $H_a : \mu_{diff} > 0$  or  $\mu_{diff} < 0$  or  $\mu_{diff} \neq 0$ , where  $\mu_{diff}$  is the population mean of the paired difference
- Test statistic:  $t_{obs} = \frac{\bar{X}_{diff} - 0}{\frac{s_{diff}}{\sqrt{n}}}$

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# ANOVA

## Overall F-Test

- $H_0 : \mu_1 = \mu_2 = \cdots = \mu_J$   
 $H_a : \text{At least one mean is different}$

- ANOVA Table:

Source	df	SS	MS	F statistic
Treatment	$J - 1$	$SSTr$	$MSTr = \frac{SSTr}{J-1}$	$F = \frac{MSTr}{MSE}$
Error	$N - J$	$SSE$	$MSE = \frac{SSE}{N-J}$	
Total	$N - 1$	$SSTo$		

- Test Statistic:  $F_{obs} = \frac{MSTr}{MSE}$ . Under  $H_0$ ,  $F^* \sim F_{df_1=J-1, df_2=N-J}$

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# Multiple Comparisons & Linear Contrasts



# Family-Wise Error Rate (FWER) and Multiple Comparisons

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- **Family-Wise Error Rate (FWER)  $\bar{\alpha}$** : the probability of making 1 or more type I errors in a set of hypothesis tests
- **Bonferroni Correction**: Adjust the significant level for each of the  $m$  tests to be  $\frac{\alpha}{m}$  to control the **FWER**
- **Fisher's LSD and Tukey's HSD**

- **Definition:** Let  $c_1, c_2, \dots, c_J$  are constants where  $\sum_{j=1}^J c_j = 0$ , then  $L = \sum_{j=1}^J c_j \mu_j$  is called a **linear contrast** of the population means.
- Point Estimation:

$$\hat{L} = \sum_{j=1}^J c_j \bar{X}_j$$

- Interval Estimation:

$$(\hat{L} - t_{\alpha/2, df=N-J} \hat{se}_{\hat{L}}, \hat{L} + t_{\alpha/2, df=N-J} \hat{se}_{\hat{L}}),$$

$$\text{where } \hat{se}_{\hat{L}} = \sqrt{\text{MSE} \left( \frac{c_1^2}{n_1} + \dots + \frac{c_J^2}{n_J} \right)}$$

- Hypothesis Testing for linear contrasts

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- ANOVA table

Source	df	SS	MS	F statistic
Treatment	$J - 1$	$SS_{trt}$	$MS_{trt} = \frac{SS_{trt}}{J-1}$	$F_{trt} = \frac{MS_{trt}}{MS_{err}}$
Block	$B - 1$	$SS_{blk}$	$MS_{blk} = \frac{SS_{blk}}{B-1}$	
Error	$(B - 1)(J - 1)$	$SS_{err}$	$MS_{err} = \frac{SS_{err}}{(B-1)(J-1)}$	
Total	$N - 1$	$SStot$		

- Why we may want to do blocking (See the example in Lecture 17)?
- Use interaction plot to assess the additivity assumption (i.e., treatment effects are consistent across blocks)