11 Tests for the Effects of Two Factors (Cont'd)

MTH 3240 Environmental Statistics

Spring 2020

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MTH 3240 Environmental Statistics

Objectives

Objectives:

- Write out the group means and treatment effects versions of the two-factor ANOVA model, including any assumptions about the random error term *ε*. (Optional for Spring 2020)
- Interpret sums of squares, degrees of freedom, and mean squares in two-factor ANOVA.
- Carry out two-factor ANOVA *F* tests for the effects of two factors and their interaction effect.
- Interpret the interaction effect in a two-factor study.
- Obtain and interpret fitted values and residuals in two-factor ANOVA.
- Use plots to check the normality and common population standard deviation assumptions required by the ANOVA *F* tests.

Two-Factor ANOVA Models (Optional for Spring 2020)

• For two-factor ANOVA, we'll suppose each of the *ab* groups is a sample from a **normal** population.

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The mean of the population corresponding to the i, jth group is μ_{ij} .

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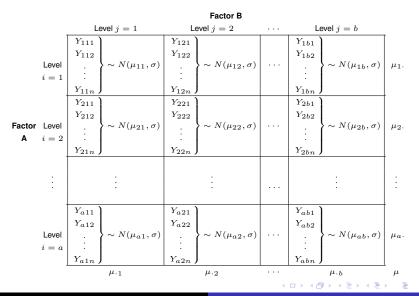
Two-Factor ANOVA Models (Optional for Spring 2020)

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The mean of the population corresponding to the i, jth group is μ_{ij} .

The population standard deviations are required to all the same, σ .

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- The value of μ_{ij} will depend on the level i of factor A and the level j of factor B.
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- We can describe the data using the statistical model called the additive effects two-factor ANOVA model.

In the model, each μ_{ij} is written in terms of an **effect** of level *i* of factor *A* and an **effect** of level *j* of factor *B*.

Two-Factor ANOVA (Cont'd)

Additive Two-Factor ANOVA Model (Optional for Spring 2020)

Additive Two-Factor ANOVA Model: One statistical model for describing data in a two-factor study is:

$$Y_{ijk} = \underbrace{\mu + \alpha_i + \beta_j}_{\text{This is } \mu_{ij}} + \epsilon_{ijk} ,$$

where

 Y_{ijk} is the *k*th observation at the *i*th level of factor A and *j*th level of factor B.

 μ is a constant called the **overall true mean**.

 α_i is the *effect* of the *i*th level of factor A.

 β_j is the *effect* of the *j*th level of factor *B*.

etitic is a N(0, σ) **random error** term. MTH 3240 Environmental Statistics

Example

For the soil phosphorus study, the **additive effects model** for describing a phosphorus concentration Y is of the form

Y =Overall Mean+Soil Type Effect+Topography Effect+Error

 But the additive effects model requires that the effects of the two factors be additive.

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Example

For the soil phosphorus study, the **additive effects model** for describing a phosphorus concentration Y is of the form

Y =Overall Mean+Soil Type Effect+Topography Effect+Error

- But the additive effects model requires that the effects of the two factors be additive.
- They're *additive* when the effect of factor *A* is the same regardless of the level of factor *B*, and the effect of factor *B* is the same regardless of the level of factor *A*.

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 When the effects of two factors aren't additive, we say that there's an *interaction effect* between them.

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Interactions

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- An *interaction effect* exists when the effect of factor A is different depending on the level of factor B, and the effect of factor B is different depending on the level of factor A.

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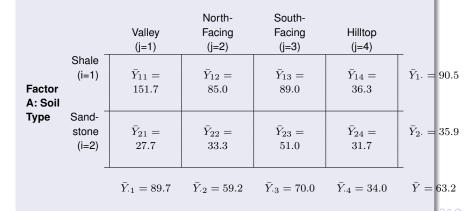
Interactions

- When the effects of two factors aren't additive, we say that there's an *interaction effect* between them.
- An *interaction effect* exists when the effect of factor A is different depending on the level of factor B, and the effect of factor B is different depending on the level of factor A.
- We can check for an interaction effect by graphing the group means either in a bar plot or in an interaction plot.

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Example

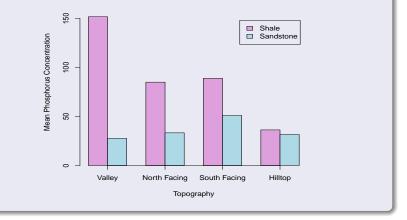
For the soil phosphorus study, the eight **group means** are shown in the **body** of the table below.



Factor B: Topography

A bar plot of the group means is shown (again) below.

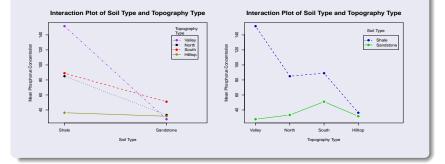
Mean Phosphorus Concentrations For Different Soil Type and Topography Combinations



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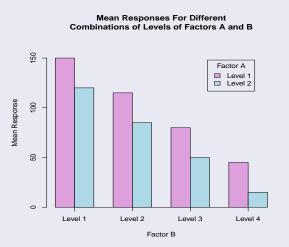
Here are interaction plots of the group means.



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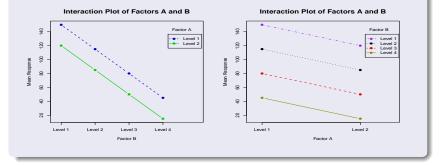
Example

Here's **bar plot** showing group means for two factors whose effects are **additive**.



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Here are **interaction plots** of those same **group means** (for two factors whose effects are **additive**).



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When the lines **aren't parallel**, the effects **aren't additive**, i.e. there's an **interaction effect**.

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 - We only have to look at one of the interaction plots or the other (not both).

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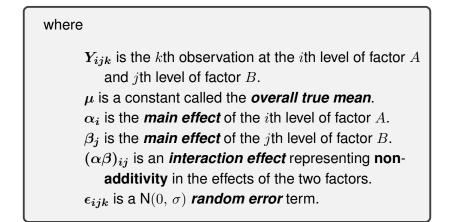
• The lines in an interaction plot won't ever be exactly parallel because of random *sampling error*.

Two-Factor ANOVA Model With Interaction (Optional for Spring 2020)

 The model we'll use when the effects of two factors aren't additive is the *two-factor ANOVA model with interaction effect*.

Two-Factor ANOVA Model With Interaction Effect: Another statistical model for describing data in a two-factor study is:

$$Y_{ijk} = \underbrace{\mu + \alpha_i + \beta_j + (\alpha\beta)_{ij}}_{\text{This is } \mu_{ij}} + \epsilon_{ijk} ,$$



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... and it can still be used even if they *are* additive.

So it's "safer" to include the interaction effect in the model when carrying out a two-factor ANOVA.

From here on, we'll *always* include the interaction effect in the model.

Example

For the soil phosphorus study, the interaction plots suggest that the effects of **soil type** and **topography** *aren't* **additive**, so the we'd definitely use the **ANOVA model** *with* the **interaction effect**.

For a phosphorus concentration *Y*, the model has the form:

Y =Overall Mean + Soil Type Effect + Topography Effect + Interaction Effect + Error

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Fitted Values and Residuals

• The group means $\bar{Y}_{11}, \bar{Y}_{12}, \ldots, \bar{Y}_{ab}$ are sometimes called *fitted values*.

Fitted Values:

Fitted Value for *ij*th Group = \bar{Y}_{ij}

• Comments:

• Statistical software reports **n** duplicates of the fitted value for each of the *ab* groups, one duplicate for each of the *n* individuals in the group.

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Residuals

• A *residual*, denoted e_{ijk} , is the **deviation** of an individual's observed Y_{ijk} value away from the **fitted value** for that individual.

Residuals:

$$e_{ijk} = Y_{ijk} - \bar{Y}_{ij}$$

• Statistical software reports the values of all N residuals, one for each individual in the study.

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Sums of Squares

 Recall that we'll test for a factor A main effect by comparing the between-rows variation to within-groups variation, ...

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and we'll test for a **factor B main effect** by comparing the **between-columns variation** to **within-groups** variation.

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and we'll test for a **factor B main effect** by comparing the **between-columns variation** to **within-groups** variation.

(We'll also test for an **interaction effect** by comparing **variation due to the interaction** to **within-groups** variation.)

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and we'll test for a **factor B main effect** by comparing the **between-columns variation** to **within-groups** variation.

(We'll also test for an **interaction effect** by comparing **variation due to the interaction** to **within-groups** variation.)

 We measure the different types of variation using sums of squares (and later mean squares).

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 We measure the between-rows variation by the factor A sum of squares, denoted SSA.

Factor A Sum of Squares:

$$\mathsf{SSA} = nb\sum_{i=1}^{a} (\bar{Y}_{i\cdot} - \bar{Y})^2.$$

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SSA =
$$nb \sum_{i=1}^{a} (\bar{Y}_{i\cdot} - \bar{Y})^2$$
.

SSA will be **large** when there's substantial variation among **row** means $\bar{Y}_{1.}, \bar{Y}_{2.}, \ldots, \bar{Y}_{a.}$, which would suggest **factor A** has an effect.

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• We measure the **between-columns variation** by the *factor B sum of squares*, denoted **SSB**.

Factor *B* Sum of Squares:

SSB =
$$na \sum_{j=1}^{b} (\bar{Y}_{j} - \bar{Y})^2$$
.

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• We measure the **between-columns variation** by the *factor B sum of squares*, denoted **SSB**.

Factor *B* Sum of Squares:

SSB =
$$na \sum_{j=1}^{b} (\bar{Y}_{j} - \bar{Y})^2$$
.

SSB will be **large** when there's substantial variation among **column means** $\bar{Y}_{1}, \bar{Y}_{2}, \ldots, \bar{Y}_{b}$, which would suggest **factor B** has an **effect**.

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• We measure variation due to an interaction effect by the *AB interaction sum of squares*, denoted SSAB.

AB Interaction Sum of Squares:

SSAB =
$$n \sum_{i=1}^{a} \sum_{j=1}^{b} (\bar{Y}_{ij} - \bar{Y}_{i} - \bar{Y}_{j} + \bar{Y})^2$$
.

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• We measure variation due to an interaction effect by the *AB interaction sum of squares*, denoted SSAB.

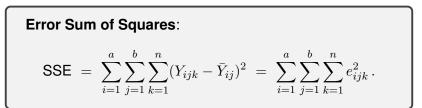
AB Interaction Sum of Squares:

SSAB =
$$n \sum_{i=1}^{a} \sum_{j=1}^{b} (\bar{Y}_{ij} - \bar{Y}_{i} - \bar{Y}_{j} + \bar{Y})^2$$
.

It can be shown that SSAB will be **large** when the **group means** $\bar{Y}_{11}, \bar{Y}_{12}, \ldots, \bar{Y}_{ab}$ aren't consistent with an additive effects model, which would suggest there's an **interaction effect** between **factors A** and **B** (i.e. their effects aren't additive).

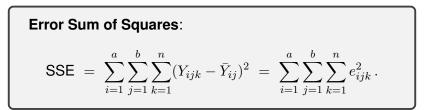
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 We measure within-groups variation by the error sum of squares, denoted SSE.



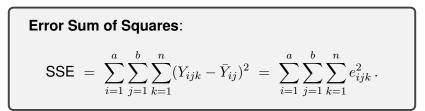
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The error sum of squares is just the *sum of squared residuals*, and reflects variation due to **random error**.

 We measure within-groups variation by the error sum of squares, denoted SSE.



The error sum of squares is just the *sum of squared residuals*, and reflects variation due to **random error**.

SSE will be **large** if there's **substantial** variation among individual observations (Y_{ijk} 's) within groups, and **small** otherwise.

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The ANOVA Partition

 The total sum of squares, denoted SSTo, measures total variation in the data.

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Total Sum of Squares: SSTo $= \sum_{i=1}^{a} \sum_{j=1}^{b} \sum_{k=1}^{n} (Y_{ijk} - \overline{Y})^2$.

The ANOVA Partition

 The total sum of squares, denoted SSTo, measures total variation in the data.

Total Sum of Squares:

SSTo =
$$\sum_{i=1}^{a} \sum_{j=1}^{b} \sum_{k=1}^{n} (Y_{ijk} - \bar{Y})^2$$
.

SSTo reflects **total variation** due both to effects of the factors (if they have any effects) **and** random variation among individual observations within groups.

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It can be shown that

Two-Factor ANOVA Partition:

SSTo = SSA + SSB + SSAB + SSE.

This splits the total variation in the data as:

Total Variation = Between-Rows Variation

- + Between-Columns Variation
- + Variation Due to Interaction

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+ Within-Groups Variation

Example

For the soil phosphorus study, statistical software reports the following **sums of squares**.

SSTo	=	51406.0	(Total variation)
SSA	=	17876.0	(Variation due to soil type)
SSB	=	9693.8	(Variation due to topography)
SSAB	=	11390.8	(Variation due to interaction)
SSE	=	12445.3	(Variation due to random error)

We see that the two-factor ANOVA partition holds since

51406.0 = 17876.0 + 9693.8 + 11390.8 + 12445.3.

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This indicates that a large portion of the total variation in phosphorus measurements (**17876.0** out of **51406.0**, or **35%**) is due to the difference between the row means (soil types).

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Degrees of Freedom

 Here are the *degrees of freedom* associated with the sums of squares in two-factor ANOVA. (These will determine which F distributions our p-values come from.)

Degrees of Freedom: For two-factor ANOVA, the degrees of freedom are:

$$df \text{ for SSTo} = N - 1$$

$$df \text{ for SSA} = a - 1$$

$$df \text{ for SSB} = b - 1$$

$$df \text{ for SSAB} = (a - 1)(b - 1)$$

$$df \text{ for SSE} = ab(n - 1) = N - ab$$

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• The degrees of freedom, like the associated sums of squares, are additive in the following sense.

Additivity of Degrees of Freedom:

df for SSTo = df for SSA+df for SSB+df for SSAB+df for \$SE.

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Example

For the soil phosphorus study, we have a = 2 soil types, b = 4 topographies, and n = 3 phosphorus observations per group. Thus the total number of phosphorus observations (overall sample size) is N = 24, and

df for SSTo = 23df for SSA = 1df for SSB = 3df for SSAB = 3

df for SSE = 16.

As expected,

23 = 1 + 3 + 3 + 16.

Mean Squares

- A mean square is a sum of squares divided by its degrees of freedom.
- The *factor A mean square*, *factor B mean square*, *AB interaction mean square*, and *mean squared error* are below.

Mean Squares: For two-factor ANOVA, the mean squares are					
$MSA = \frac{SSA}{a-1}$	$MSB = \frac{SSB}{b-1}$				
$MSAB = \frac{SSAB}{(a-1)(b-1)}$	$MSE = \frac{SSE}{ab(n-1)}$				

The ANOVA F Tests

• There are **three** sets of hypotheses for the **two-factor ANOVA** F **tests** tests.

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The ANOVA F Tests

• There are **three** sets of hypotheses for the **two-factor ANOVA** F **tests** tests.

In each case, H_0 says there's **no effect**, and H_a say's there's **an effect**.

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 - H_{OB} : There's no factor B effect H_{aB} : There is a factor B effect

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- 1. Test for a *factor* A *main effect*:
 - H_{OA} : There's no factor A effect H_{aA} : There is a factor A effect
- 2. Test for a *factor* B *main effect*:
 - H_{OB} : There's no factor B effect H_{aB} : There is a factor B effect
- 3. Test for an *AB* interaction effect:
 - H_{OAB} : There's no factor A and B interaction effect
 - H_{aAB} : There is a factor A and B interaction effect

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(Optional for Spring 2020)

In terms of the **ANOVA model parameters** these are written as:

1. Test for a *factor* A *main effect*:

 H_{OA} : $\alpha_1 = \alpha_2 = \cdots = \alpha_a = 0$ H_{aA} : The α_i 's don't all equal 0

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2. Test for a *factor* B main effect:

 $\begin{aligned} H_{OB}: & \beta_1 = \beta_2 = \cdots = \beta_b = 0 \\ H_{aB}: & \text{The } \beta_j \text{'s don't all equal 0} \end{aligned}$

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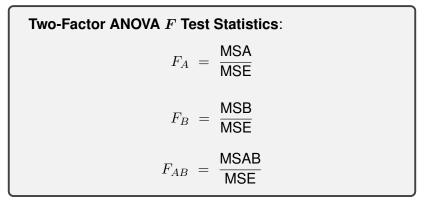
2. Test for a *factor* B *main effect*:

 $\begin{aligned} H_{OB}: & \beta_1 = \beta_2 = \dots = \beta_b = 0 \\ H_{aB}: & \text{The } \beta_j \text{'s don't all equal 0} \end{aligned}$

3. Test for an *AB* interaction effect:

 $H_{OAB}: \quad (\alpha\beta)_{11} = (\alpha\beta)_{12} = \dots = (\alpha\beta)_{ab} = 0$ $H_{aAB}: \quad \text{The } (\alpha\beta)_{ij}\text{'s don't all equal } 0$

 Here are the corresponding two-factor ANOVA F test statistics:



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Not that we can think of

$$F_A = \frac{\text{Between-Rows Variation}}{\text{Within-Groups Variation}}.$$

$$F_B = \frac{\text{Between-Columns Variation}}{\text{Within-Groups Variation}}.$$

$$F_{AB} = \frac{\text{Variation Due To Interaction}}{\text{Within-Groups Variation}}.$$

 In each case, if H₀ was true, it can be shown, F would be approximately equal to 1.

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- But if if H_0 was true was true, F would be greater than 1.

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In each case, *large* values of F (larger than about 1) provide evidence in favor of H_a .

Now suppose the *ab* groups are samples from normal populations that all have the same standard deviation *σ* (or that they have the same *σ* and the common sample size *n* is large).

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In this case, the **null distributions** are as follows.

Sampling Distribution of *F* **Under** H_0 : If F_A , F_B , and F_{AB} are the two-factor ANOVA *F* test statistics, then:

• When H_{OA} is true,

$$F_A = \frac{\mathsf{MSA}}{\mathsf{MSE}} \sim F(a-1, N-ab).$$

2 When H_{OB} is true,

$$F_B = \frac{\mathsf{MSB}}{\mathsf{MSE}} \sim F(b-1, N-ab).$$

3 When H_{OAB} is true,

$$F_{AB} = \frac{\mathsf{MSAB}}{\mathsf{MSE}} \sim F((a-1)(b-1), N-ab).$$

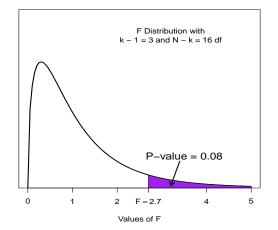
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• **P-values** and **rejection regions** are obtained from the *upper tail* of the **appropriate** *F* **distribution**.

- **P-values** and **rejection regions** are obtained from the *upper tail* of the **appropriate** *F* **distribution**.
- The next slide shows the **p-value** when the observed test statistic value is F = 2.7.

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P-Value for ANOVA F Test



The ANOVA Table

The results of an analysis of variance are summarized in a two-factor ANOVA table having the form shown below.

Two-Factor ANOVA Table:											
Source	DF	SS	MS	F	P-value						
Factor A	a-1	SSA	MSA = SSA/(a-1)	MSA/MSE	р						
Factor B	b-1	SSB	MSB = SSB/(b-1)	MSB/MSE	р						
Interaction ((a-1)(b-1)	SSAB	MSAB = SSAB/((a-1)(b-1))	MSAB/MSE	р						
Error	ab(n-1)	SSE	MSE = SSE/(ab(n-1))								
Total	N-1	SSTo									

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• First look at the results for the *AB* interaction effect.

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- First look at the results for the *AB* interaction effect.
 - If it's statistically significant, there's no need proceed to the results of the tests for main effects since we already know that both factors have effects.

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(But the effect of each factor is different depending on the level of the other factor.)

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If the AB interaction isn't significant, proceed to the results of the tests for factor A and B main effects.

Example

For the soil phosphorus study, the **ANOVA table** (obtained using software) is below.

Source of		Sum of	Mean		
Variation	df	Squares	Square	f	P-value
Soil Type	1	17876.0	17876.0	22.98	0.000
Topography Type	3	9693.8	3231.3	4.15	0.024
Interaction	3	11390.8	3796.9	4.88	0.013
Error	16	12445.3	777.8		
Total	23	51406.0			

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What **conclusions** are appropriate (using $\alpha = 0.05$)?

The p-value for the interaction is **0.013**, so we conclude there's a **statistically significant** interaction effect between **soil type** and **topography**.

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In this case, there's **no need to proceed** to look at the p-values for the **soil type** and **topography main effects**.

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In this case, there's **no need to proceed** to look at the p-values for the **soil type** and **topography main effects**.

We **already know** (because the interaction is significant) that each has an effect on phosphorus, **regardless** of what their p-values are.

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Each factor has a **different effect** depending on the **level** of the **other factor**.

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Two-Factor ANOVA (Cont'd)

Checking the ANOVA Assumptions

• The **ANOVA** *F* tests require that the *ab* groups (samples) are from **normal** populations (or that their sample sizes are **large**) whose **standard deviations** are all **equal**.

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• Two ways to check the normality assumption:

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 - Make *ab separate* histograms or normal probability plots, one for each of the *ab* groups.

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- Two ways to check the **normality assumption**:
 - Make *ab separate* histograms or normal probability plots, one for each of the *ab* groups.
 - Make a single histogram or normal probability plot plot of the N residuals e_{ijk} .

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• A few ways to check the equal population standard deviation assumption:

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• An *individual value plot* of the *ab* groups.

- A few ways to check the equal population standard deviation assumption:
 - An *individual value plot* of the *ab* groups.
 - A plot of the *residuals* (*y*-axis) versus **fitted values** (group means, *x*-axis).

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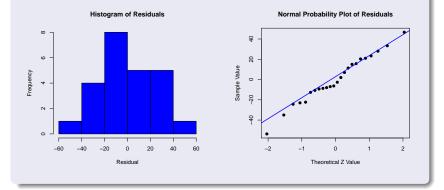
- A few ways to check the equal population standard deviation assumption:
 - An *individual value plot* of the *ab* groups.
 - A plot of the *residuals* (*y*-axis) versus **fitted values** (group means, *x*-axis).

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 In both plots, we look for roughly equal amounts of within-group (vertical) spread across the *ab* groups.

Example

For the soil phosphorus study, a **histogram** and **normal probability plot** of the **residuals** are below.



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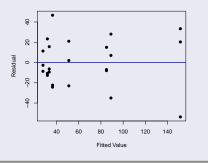
The plots show that the **normality assumption** appears to be met.

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The plots show that the **normality assumption** appears to be met.

A plot the **residuals** versus the **fitted values** (**group means**) is below.



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Residuals Versus Fitted Values

The amount of (vertical) spread of the points is roughly the same from left to right, so the **equal standard deviation assumption** appears to be met.

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The amount of (vertical) spread of the points is roughly the same from left to right, so the **equal standard deviation assumption** appears to be met.

Because the **normality** and **equal standard deviation assumptions** are met, the results of the F tests are valid.

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