

## Hypothesis Testing Two Sample

### Null Hypothesis

$$H_0: \mu_1 = \mu_2$$

$$H_0: \mu_1 - \mu_2 = 0 \quad \text{Not Different}$$

$$H_0: \delta = 0$$

### Alternative Hypothesis

$$H_A: \mu_1 \neq \mu_2$$

$$H_A: \mu_1 - \mu_2 \neq 0 \quad \text{Different}$$

$$H_A: \delta \neq 0$$

## Hypothesis Testing Error Types

	Reject $H_0$	Accept $H_0$
H <sub>0</sub> True	Type I Error (False Positive)	Correct
H <sub>0</sub> False	Correct	Type II Error (False Negative)

## Hypothesis Testing

### Error Types

Decision	Truth		
	$\delta < 0$	$\delta = 0$	$\delta > 0$
$\delta < 0$	<b>Correct</b>	Type I	Type III
$\delta = 0$	Type II	<b>Correct</b>	Type II
$\delta > 0$	Type III	Type I	<b>Correct</b>

$$\delta = \mu_1 - \mu_2$$

## Hypothesis Testing

### Probabilities

#### Alpha

$$\alpha = P(\text{Type I Error})$$

$$\alpha = P(\text{Reject } H_0 \mid H_0 \text{ is True})$$

#### Beta

$$\beta = P(\text{Type II Error})$$

$$\beta = P(\text{Accept } H_0 \mid H_0 \text{ is False})$$

#### Power

$$= 1 - \beta$$

## Comparing Two Sample Means

*t*-tests

$$t = \frac{\bar{x}_1 - \bar{x}_2}{s_{\bar{x}_1 - \bar{x}_2}} = \frac{\bar{d}}{s_{\bar{d}}}$$

$$s_{\bar{d}} = \sqrt{\frac{2s_p^2}{n}}$$

$$df = (n_1 - 1) + (n_2 - 1)$$

$$H_0 : \bar{x}_1 - \bar{x}_2 = \bar{d} = 0$$

## Switchgrass

Variety Trial (Yield, Mg/ha)

Replication	Cultivar	
	A	B
1	10.8	8.9
2	12.7	8.9
3	11.4	8.6
4	13.6	8.8
5	13.9	8.5
6	13.0	6.1
<b>Mean</b>	12.6	8.3
<b>Variance</b>	1.46	1.24

## Switchgrass Comparison

*t*-test

$$H_0 : \mu_A = \mu_B$$

$$t = \frac{12.571 - 8.300}{\sqrt{\frac{2(1.349)}{6}}} = \frac{4.271}{0.671} = 6.37$$

- The probability that  $t_{1,10\text{ df}} > 6.37$  is 0.000081.
- Therefore, the probability of the two means being from the same population is very low.

## Comparing Two Sample Means Standard Errors (of the difference)

if  $s^2_1 = s^2_2$  and  $n_1 = n_2$ , then:

$$S_{\bar{d}} = \sqrt{\frac{2s_p^2}{n}}$$

if  $s^2_1 = s^2_2$ , then:

$$S_{\bar{d}} = \sqrt{s_p^2 \left( \frac{1}{n_1} + \frac{1}{n_2} \right)}$$

## Comparing Two Sample Means Unequal Variances

if  $s^2_1 \neq s^2_2$ , then:

$$S_{\bar{d}} = \sqrt{\frac{s_1^2}{n_1} + \frac{s_2^2}{n_2}}$$

Satterthwaite's Approximation:

$$df = \frac{(S_{\bar{x}_1}^2 + S_{\bar{x}_2}^2)^2}{\frac{(S_{\bar{x}_1}^2)^2}{n_1 - 1} + \frac{(S_{\bar{x}_2}^2)^2}{n_2 - 1}}$$

## Hypothesis Testing

### Relative Consequences of Type I & II Errors

	Low probability of false negative ↓ $\beta < 0.05$ Power high ↑	High probability of false negative ↑ $\beta > 0.05$ Power low ↓
Low probability of false positive $\alpha < 0.05$ ↓	<b>Ideal situation</b> Frequently not achievable, but can be done with large effect sizes	Typical of many published experiments in agronomy
High probability of false positive $\alpha > 0.05$ ↑	Exploratory research Positive effects are often re-evaluated	<b>Not an ideal situation</b> Precision may be improved by increasing sample size or reducing experimental complexity

Source: Kimberly Garland Campbell. 2018. Chapter 1 - Errors in Statistical Decision Making. Applied Statistics in Agricultural, Biological, and Environmental Sciences. ASA-CSSA-SSSA.

## Hypothesis Testing

### Relative Consequences of Type I & II Errors

Agronomy Example	Low probability of false negative ↓ $\beta < 0.05$ Power high ↑	High probability of false negative ↑ $\beta > 0.05$ Power low ↓
Low probability of false positive $\alpha < 0.05$ ↓	Growers adopt new variety and are happy.	Growers don't adopt new variety. Cost is associated with unrealized potential profit from increased yields of new variety.
High probability of false positive $\alpha > 0.05$ ↑	Growers adopt new variety, but it doesn't perform better than old variety. Cost differential depends on relative cost of seed. Or Type 3 error occurs, and new variety actually performs worse than old variety.	Experiment is too variable to make a decision. Resources for trialing wasted.

Source: Kimberly Garland Campbell. 2018. Chapter 1 - Errors in Statistical Decision Making. Applied Statistics in Agricultural, Biological, and Environmental Sciences. ASA-CSSA-SSSA.

## Switchgrass Comparison

### t-test

$$\bar{d}_{.05} = 2.008 \sqrt{\frac{2(1.349)}{6}} = 3.41$$

- The lowest difference that can be detected at  $\alpha=0.05$  is 3.41.
- You can be confident that you have the power to detect larger differences since there is no possibility of making a Type II error.
- What if you wanted to be able to detect smaller differences?

## Hypothesis Testing Power

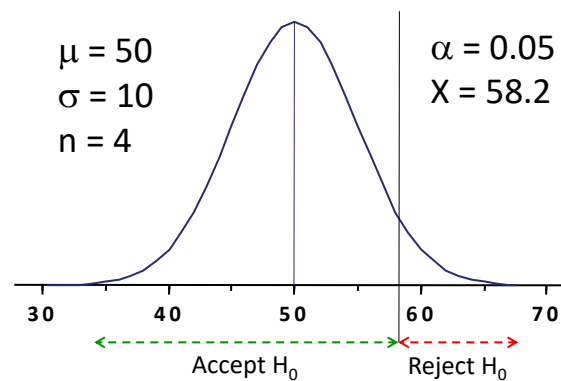
$$\text{Power} = 1 - \beta$$

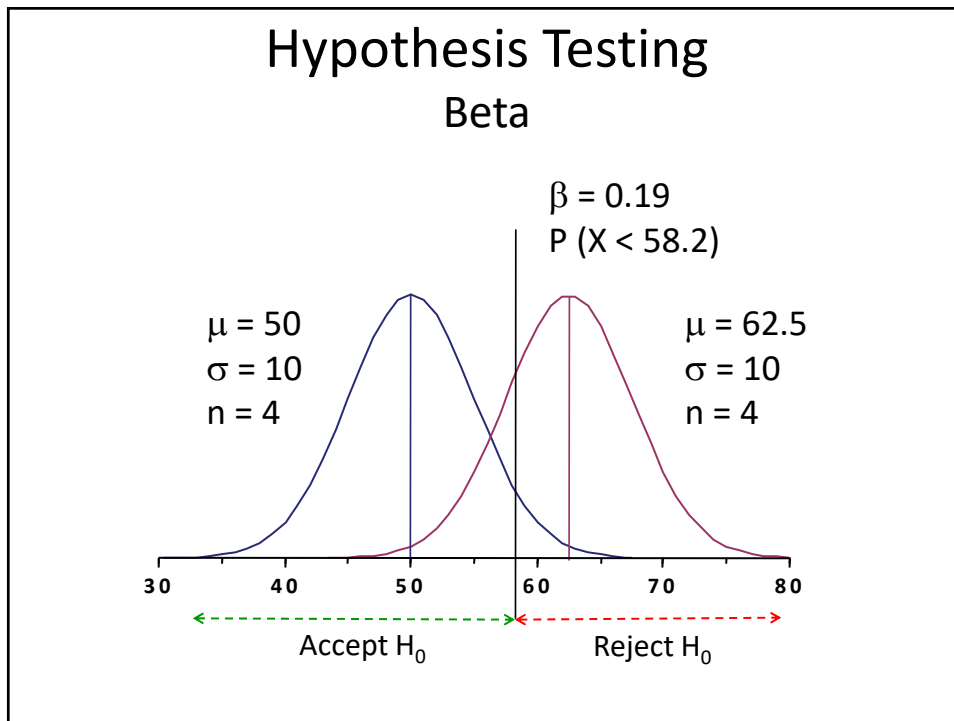
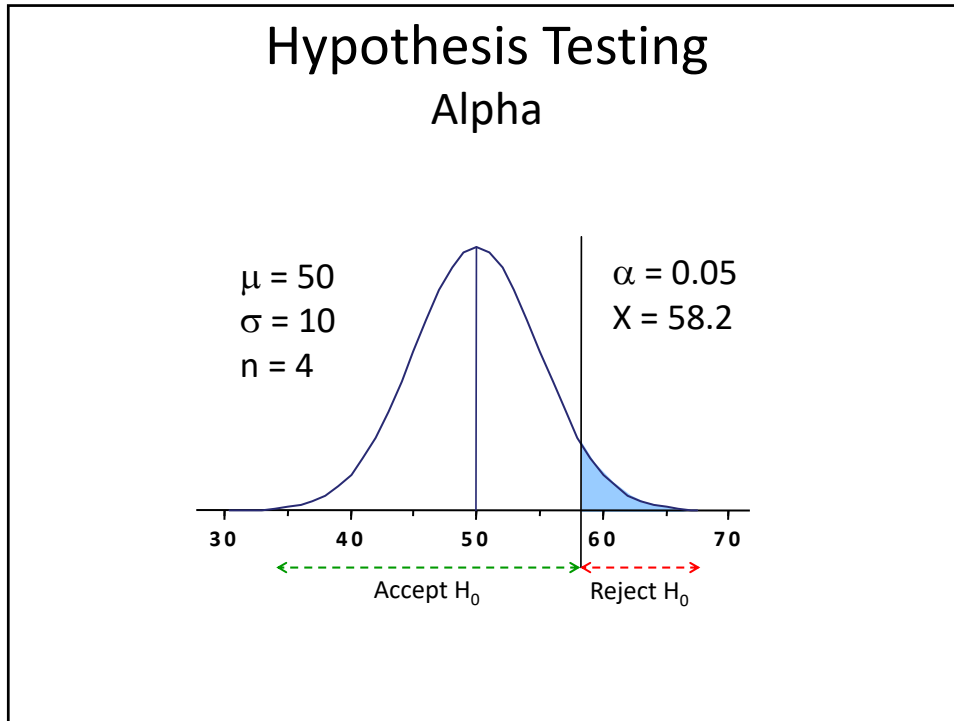
The ability to detect true differences

### Factors:

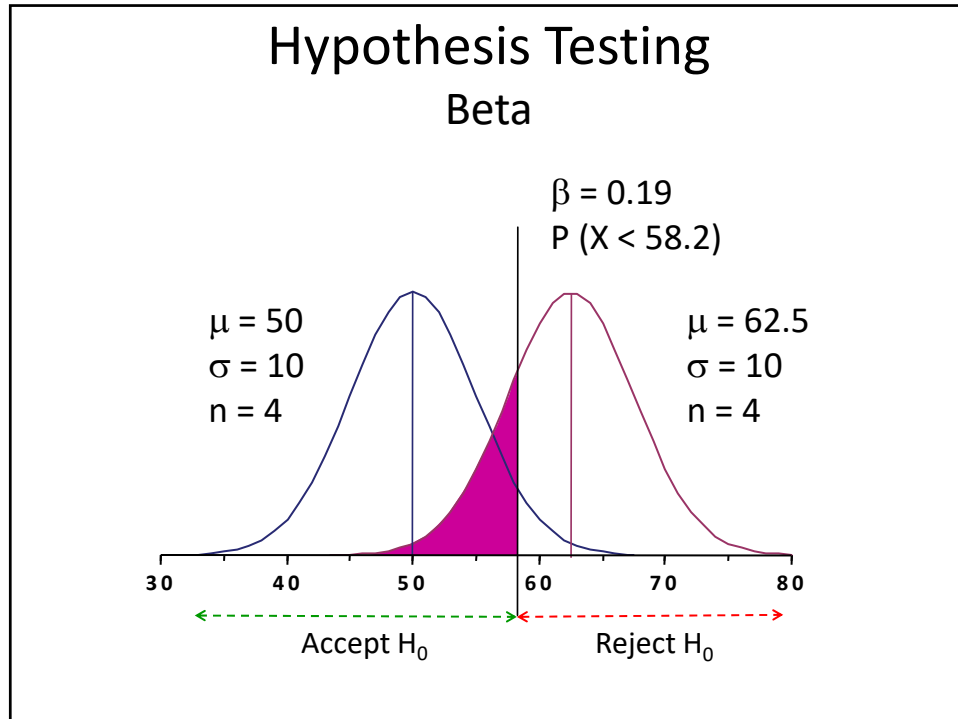
- $H_A$  is true
- Magnitude of mean difference
- Alpha – inversely related
- Variance
- Sample size

## Hypothesis Testing Alpha









### Improving Power

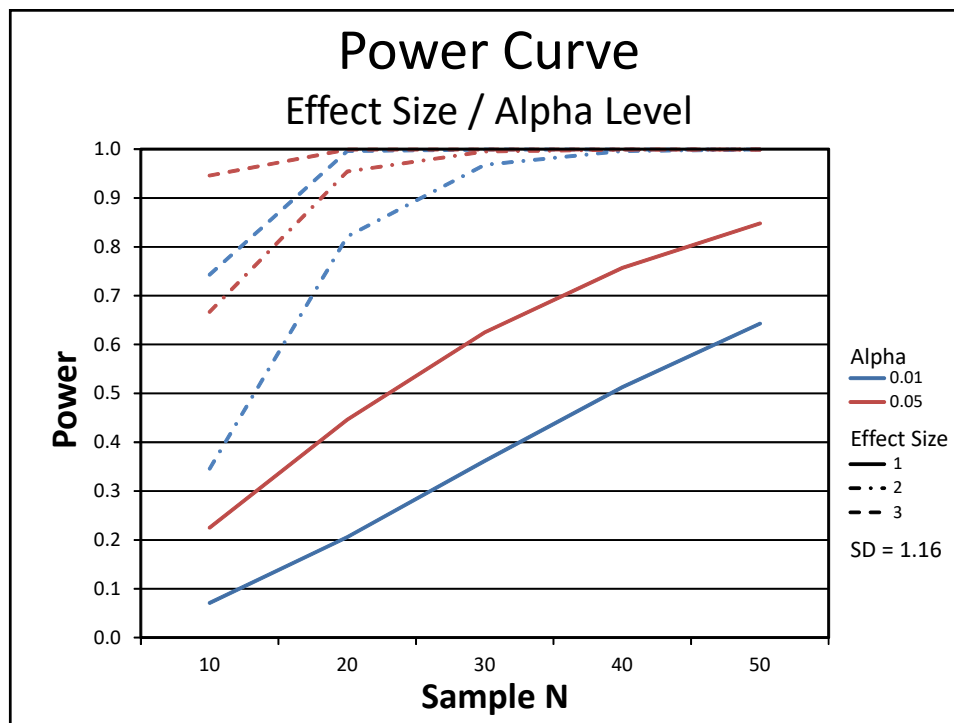
The power to detect a mean difference increases when:

- The magnitude of the difference increases
- The alpha level is relaxed (increased)
- The number of replicates increases
- The precision ( $1/\sigma^2$ ) increases (i.e.  $\sigma^2$  decreases)

## Effect Size

Statistical significance does not equate to biological or economic significance.

- It means that the ratio of the treatment variance to error variance is large enough to be considered unlikely to occur randomly.
- A difference (effect size) that is considered statistically important may not be practically so.
- Conversely, a difference that cannot be detected statistically may be quite important biologically or economically.
- ∴ It is important to think about what effect sizes you need to be able to detect in the design process.



## Hypothesis Testing Power and Replication

$$n = \frac{(t_{\alpha/2} + t_{\beta})^2 \sigma_D^2}{\delta^2}$$

Where:

n = minimum number of replicates

$t_{\alpha/2}$  = alpha

$t_{\beta}$  = beta

$\sigma_D^2$  = variance of differences ( $2\sigma^2$ )

$\delta^2$  = minimum difference

## Power and Replication

Number of reps required to detect  
mean differences

$\delta$ % Mean	CV				
	5	10	15	20	25
5	16				
10	4	16			
15	2	7	16		
20	1	4	9	16	
25	1	3	6	10	16

$\alpha = 0.05, \beta = 0.2, n = 6^*$

\* Conservative

## Mean Comparisons Improving Sensitivity

$$t = \frac{\bar{X}_1 - \bar{X}_2}{S_{\bar{X}_1 - \bar{X}_2}} = \frac{\bar{d}}{S_{\bar{d}}} \qquad S_{\bar{d}} = \sqrt{\frac{2s_p^2}{n}}$$

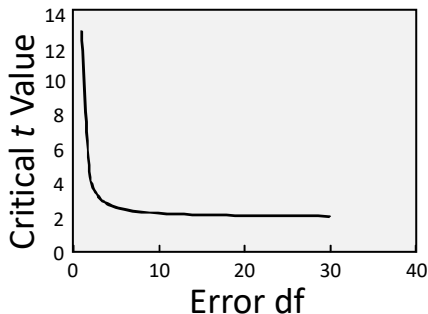
The sensitivity of an experiment for detecting treatment differences depends on the magnitude of the variance ( $S^2$ ) and the number of replications ( $n$ ).

There are two approaches for improving sensitivity:

1. Increase the number of reps used to estimate  $S^2$
2. Decrease the  $S^2$  by using design control

## Mean Comparisons Improving Sensitivity - Replication

$$LSD = t_{.05} \times S_{\bar{d}} = t_{.05} \times \sqrt{\frac{2MS_E}{r}} = t_{.05} \times \sqrt{2} \times \frac{RMSE}{\sqrt{r}}$$



**General Rule** – the sensitivity of an experiment doubles for each 4-fold increase in the number of replications.

## Mean Comparisons

### Improving Sensitivity – Plot Size

Smith's Formula:

$$V_s = \frac{V}{s^b} \quad \ln(V_n) = \ln(V_1) - b \ln(n)$$

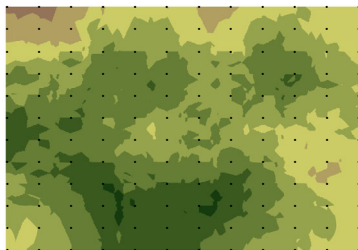
where:

$V_s$  = variance among plots of size  $s$

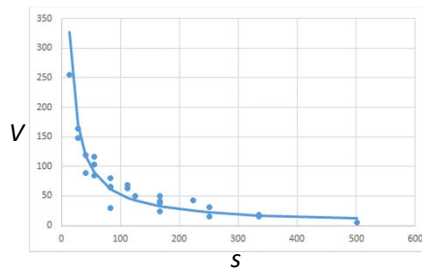
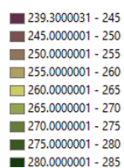
$b$  = soil heterogeneity index

## Mean Comparisons

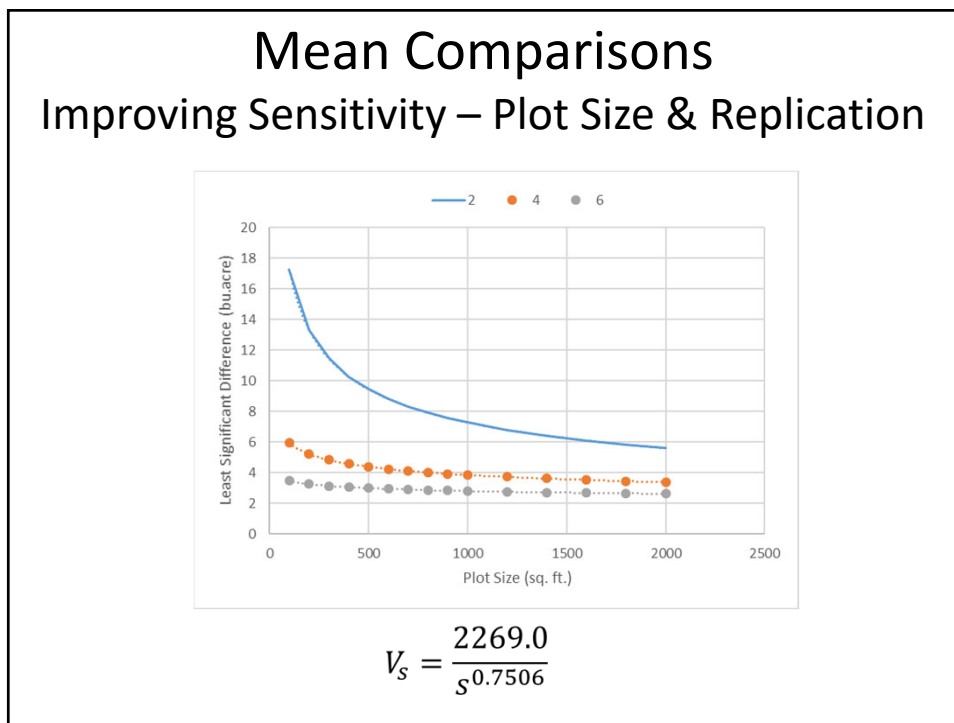
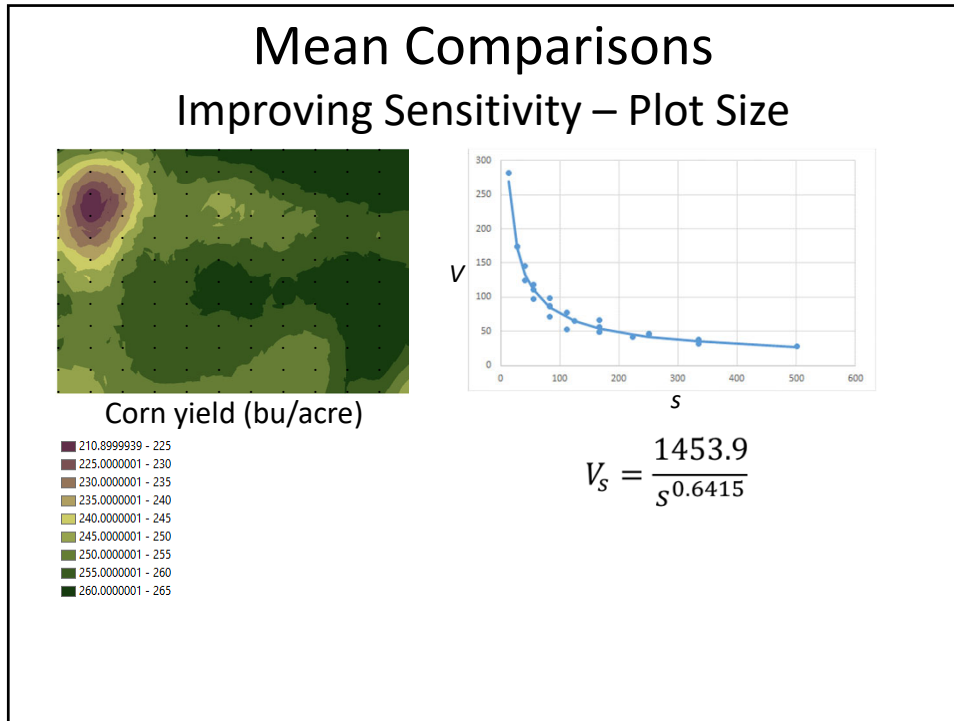
### Improving Sensitivity – Plot Size



Corn yield (bu/acre)



$$V_s = \frac{3733.1}{s^{0.9257}}$$



## Mean Comparisons

### Factors Effecting Plot Size

Adapted from Petersen, 1994

Factor	Small plots	→→→	Large Plots
Soil variability	Uniform	→→→→→→	Heterogeneous
Crop	Turf	→ Cereals →	Row crops → Pasture
Research phase	Basic	→→	Developmental →→ Adaptive
Experiment type	Breeding	→ Fertility →	Tillage → Irrigation
Machinery	None	→→→	Research →→→ Farm scale

## Mean Comparisons

### Improving Sensitivity – Design Control

Design controls are used to decrease the value of  $S^2$  by accounting for some of the unexplained variation in the measured response and partitioning it out of the error mean square ( $S^2$ ). The net effect is to make  $S^2$  smaller.

There are two common approaches for doing this:

- Blocking methods that partition some of the error SS to a blocking factor
- Using covariates to explain some of the variation not accounted for by treatments

## One-Factor ANOVA Switchgrass Example

Source	DF	Sum of Squares	Mean Square	F Value	Pr > F
Cultivar	9	355.5271	39.50301	4.57	0.0002
Error	50	432.5709	8.651419		
Total	59	788.098			

Cultivar	Mean Yield (Mg/ha)
A	12.6
B	8.3
C	14.2
D	12.7
E	9.4
F	6.7
G	13.9
H	9.6
I	9.4
J	9.1

We know from the ANOVA that there is at least one significant difference among pairs of means. How do we find out which pairs actually differ?