

RenR 480/711

Anova assumptions

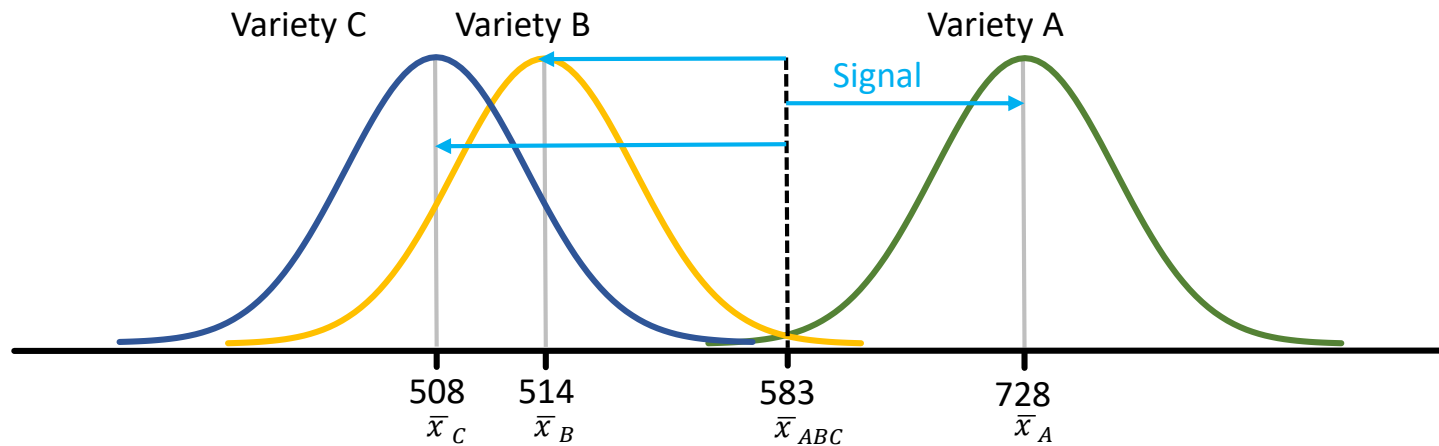
Multi-way Anova

Anova assumptions

1. Normal distribution of experimental errors/residuals
2. Equal variances between treatments
3. Independence of samples

Anova assumptions

1. Normal distribution of experimental errors/residuals

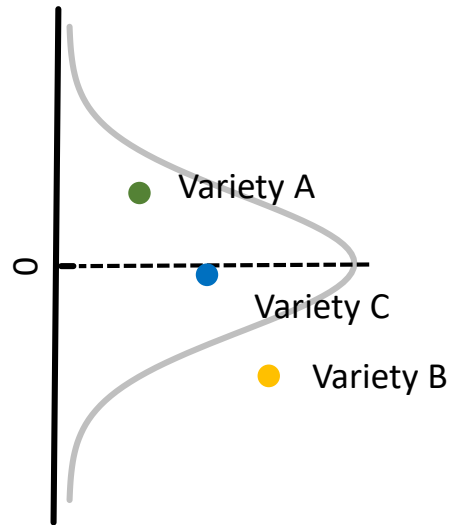


Assessment through residuals of the experiment

Residuals = difference between the experimental mean and treatment means

Anova assumptions

1. Normal distribution of experimental errors



Anova assumptions

1. Normal distribution of experimental errors

Testing for normality: **Shapiro-Wilk test**

H_0 Null Hypothesis: *Distribution of residuals = normal distribution*

H_1 Alternative Hypothesis: *Distribution of residuals \neq normal distribution*

$p > 0.05$ means that the distribution of residuals is normal

Problems with statistical tests for normality:

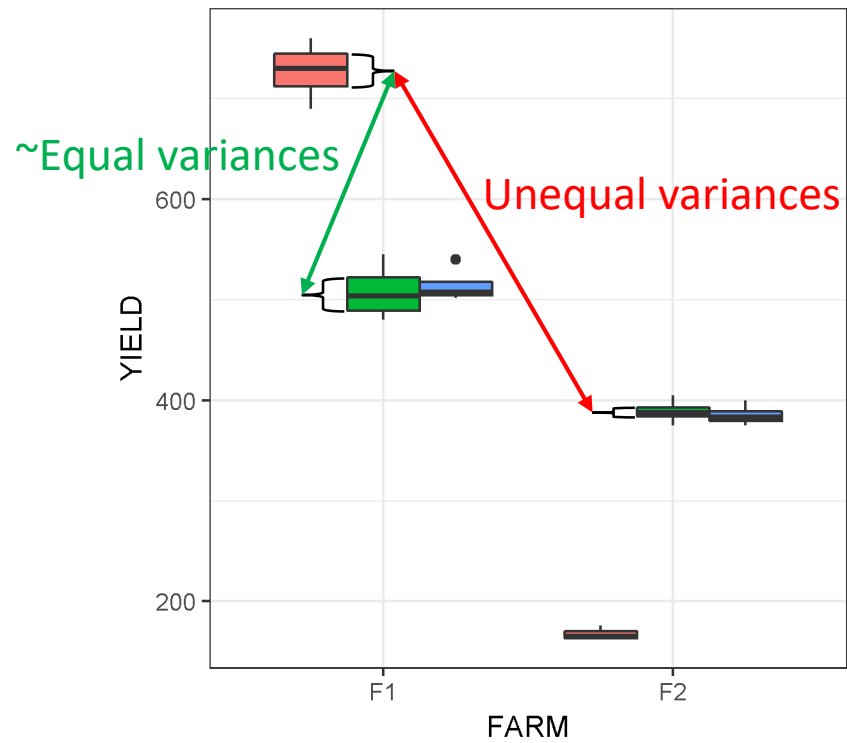
low n: **low power** \rightarrow test will fail to detect non-normality

high n: **high power** \rightarrow even tiny deviations from normality will be significant

\rightarrow Use residual plots to assess this assumption

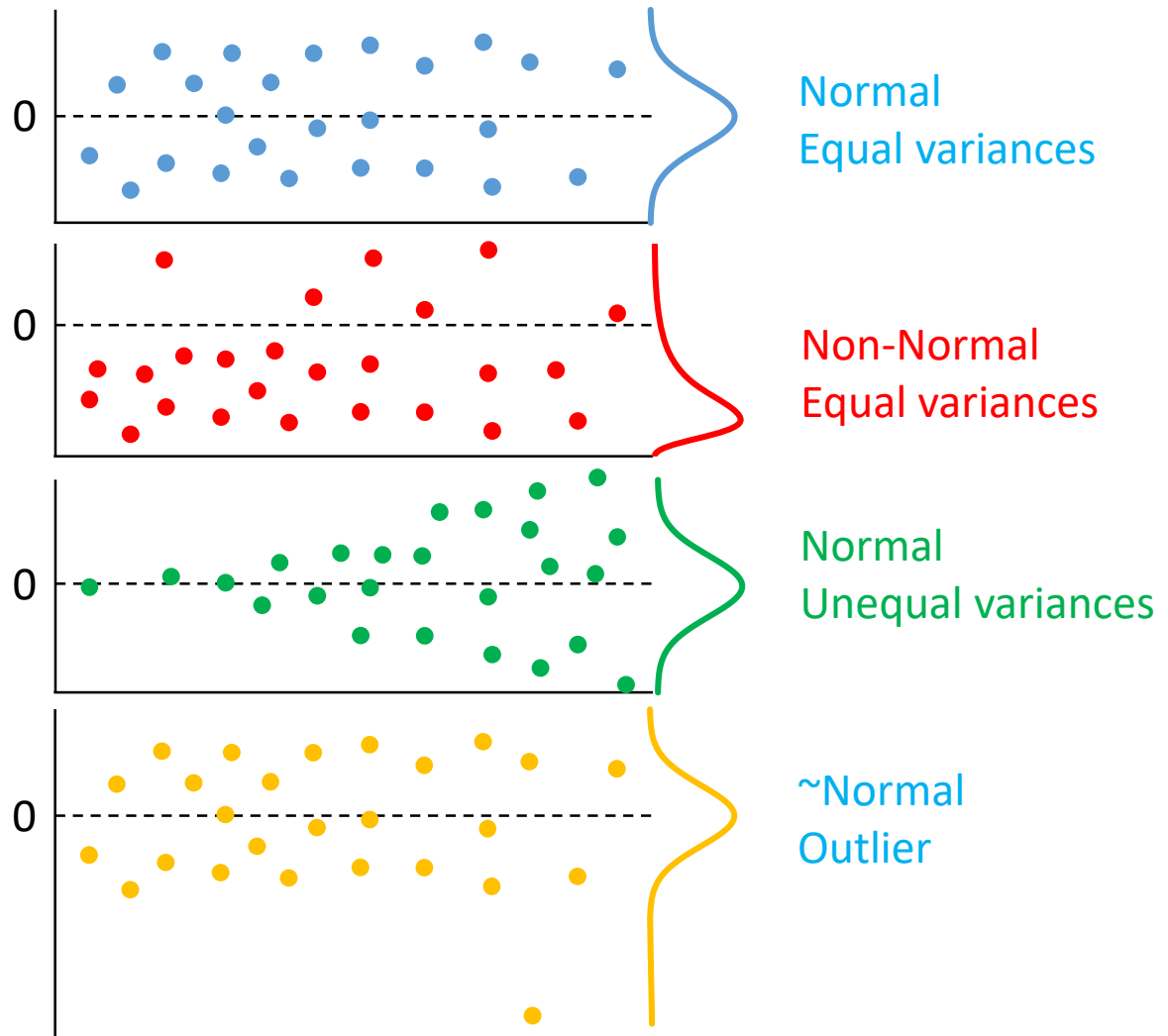
Anova assumptions

2. Equal variances between treatments



Anova assumptions

Residual plots



Transformations

Square root:

$$x_T = \sqrt{x + c}$$



All values have to be >1;
add a constant if needed

Log transformation:

$$x_T = \log(x)$$

Increasingly powerful

Inverse transformation

$$x_T = \frac{1}{x}$$



Anova assumptions

3. Independence of samples

Samples:

→ have been randomly samples *or*

→ come from a randomized design

This means: rows in your table do not influence each other

→ This assumption has to be addressed with the experimental design

Samples may not be independent if:

Multiple measurements on one subject (repeated measures)

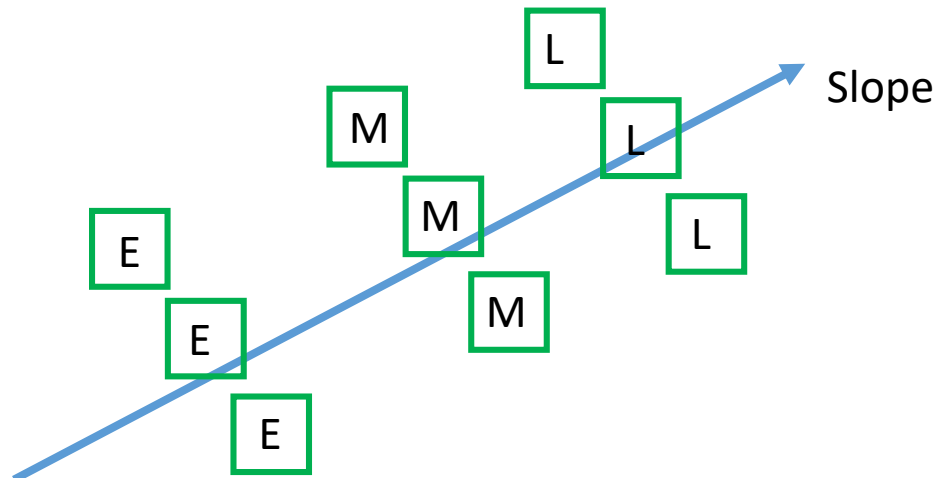
Observations are correlated in space

Observations are correlated in time

Anova assumptions

Spatial correlation

Example: Species richness in forest stands



Successional stage

E= Early

M= Mid

L= Late

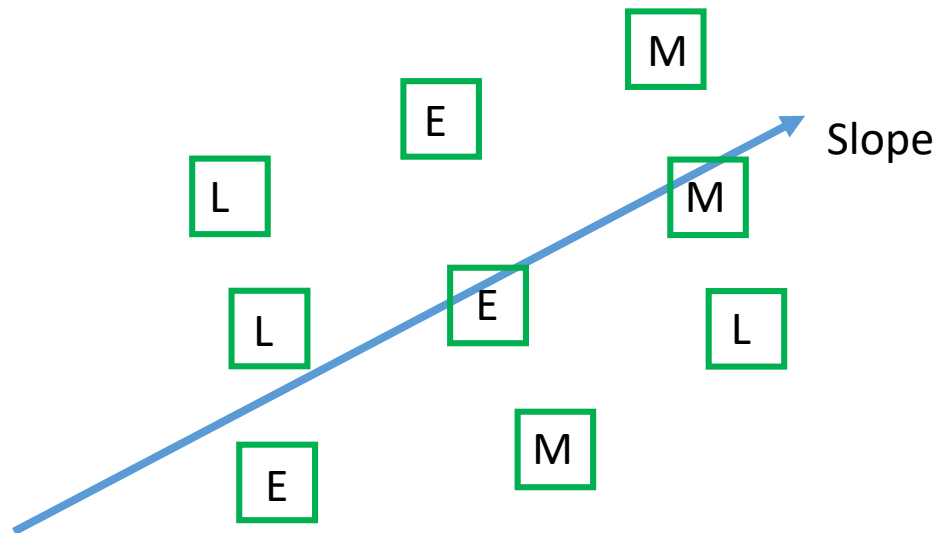
Stage	Richness
E	24
E	20
E	22
M	15
M	14
M	17
L	9
L	12
L	11

- Pseudoreplication because samples are spatially not independent
- Observed differences **cannot** be attributed to stage because slope is confounding & and there is no way of teasing this apart

Anova assumptions

Spatial correlation

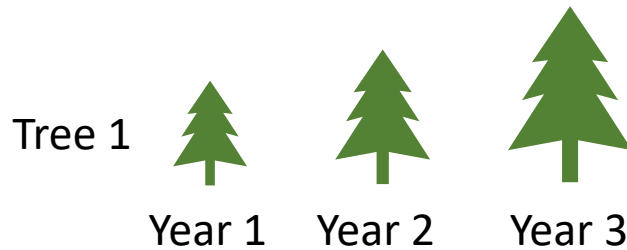
Better:



Creating a randomized design avoids pseudoreplication

Anova assumptions

Temporal correlation



Tree	Year	Height
T1	1	5
T1	2	6
T1	3	7
T2	1	6
T2	2	6
T2	3	7
T3	1	15
T3	2	16
T3	3	17

Rows are not independent – standard ANOVA not applicable
→ repeated measures ANOVA or regression analysis

Multi-way ANOVA

Multi-way-ANOVA

Same as One-way-Anova but with more than one treatment

```
anova(lm(YIELD~VARIETY*FARM))
```

```
anova(lm(YIELD~VARIETY+FARM+FARM:VARIETY))
```

Source of variation	df	Sum of squares	Mean Squares	F-value	p-value
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Variety	2				
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Farm	1				
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Variety:Farm	2				
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Residuals	18				
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Analog to One-way Anova

$$df_{\text{variety}} = n_{\text{levels of Variety}} - 1$$

$$df_{\text{Farm}} = n_{\text{levels of Farm}} - 1$$

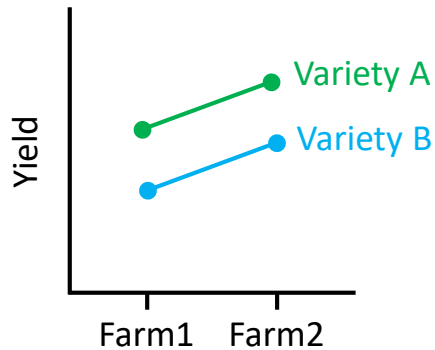
$$df_{\text{Variety:Farm}} = (n_{\text{levels of Variety}} - 1)(n_{\text{levels of Farm}} - 1)$$

$$df_{\text{Residuals}} = n_{\text{observations}} - n_{\text{levels of Variety}} \times n_{\text{levels of Farm}}$$

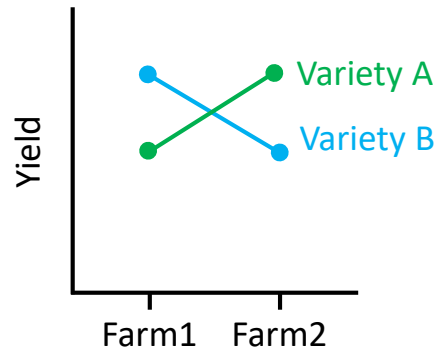
Interaction

Multi-way-ANOVA

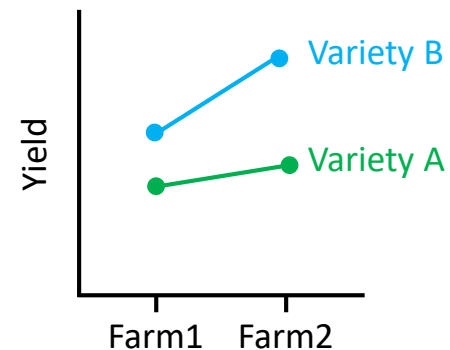
Interaction: Different story under different conditions



Variety A better than B
Farm 2 better than Farm 1
→ no interaction



Main effects n.s.
(Variety and Farm)
However, VarA is better on
Farm 2 and VarB is better
on Farm 1
→ significant interaction



Main effects significant
(but difficult to interpret
with overall means)
→ significant interaction

In case of significant interactions:

- Disregard n.s. main effects
- Avoid to interpret significant main effects
- Show the different stories for different conditions with graphs
- Follow up with separate analyses for subsets (e.g. Farm1 and Farm2)

Pairwise comparisons

Pairwise comparisons

Experiment-wise Type I-Error

Number of possible comparisons (3 varieties: A,B,C)

A-B

A-C

B-C

$$c = \frac{\text{treatments}(\text{treatments} - 1)}{2}$$

$$c = \frac{3 \times (3 - 1)}{2} = 3$$

Probability of making a Type-I error in at least one comparison

=1-probability of making no type-I error at all

$$=1 - 0.95^c$$

$$=1 - 0.95^3$$

$$=0.86$$

Need for adjustment!

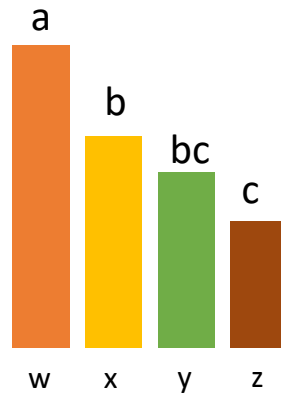
Pairwise comparisons

Bonferroni-Adjustment

$$\frac{\alpha}{c} = \frac{0.05}{3} = 0.017$$

Reporting pairwise comparisons

Graphics



	w	x	y	z
w		*	*	*
x			n.s.	*
y				n.s.
z				

Tables

	Var A	Var B	Var C
Farm1	728 a A	508 b A	514 b A
Farm2	168 a B	389 b B	385 b B