

Why does $SS_{Blocks \times Fert} + SS_{Blocks \times Geno \times Fert} = SS_{Error}$?
End lecture 29
04-07-25

- There are no terms in our model corresponding to $Block \times Fert$ combinations; thus, there is no reason to devote a separate line of our ANOVA table to $Block \times Fert$.
- Also, it can be shown that

$$E(MS_{Blocks \times Fert}) = E(MS_{Blocks \times Geno \times Fert}) = \sigma_e^2$$

Thus, it makes sense to estimate σ_e^2 with an inverse variance weighted average of independent unbiased estimators:

For this slide only, let

is needed to calculate the weights in the weight average

① = Blocks × Fert and ② = Blocks × Geno × Fert.

For $\ell = \underline{1}, \underline{2}$, $MS_{\ell} \sim \frac{E(MS_{\ell})}{df_{\ell}} \chi^2_{df_{\ell}} \Rightarrow \text{Var}(MS_{\ell}) = 2\sigma_e^4 / \underline{df_{\ell}}$.

$$\begin{aligned} \frac{\text{Var}^{-1}(MS_1)MS_1 + \text{Var}^{-1}(MS_2)MS_2}{\text{Var}^{-1}(MS_1) + \text{Var}^{-1}(MS_2)} &= \frac{\frac{df_1}{2\sigma_e^4}MS_1 + \frac{df_2}{2\sigma_e^4}MS_2}{\frac{df_1}{2\sigma_e^4} + \frac{df_2}{2\sigma_e^4}} \\ &= \frac{df_1MS_1 + df_2MS_2}{df_1 + df_2} \\ &= \frac{SS_1 + SS_2}{df_1 + df_2} \end{aligned}$$

MSE

Thus, we combine the $Blocks \times Fert$ and $Blocks \times Geno \times Fert$ lines of the ANOVA table and label the resulting line as $Error$.

$$SS_{Blocks \times Fert} + SS_{Blocks \times Geno \times Fert} = SS_{Error}$$

$$df_{Blocks \times Fert} + df_{Blocks \times Geno \times Fert} = df_{Error}$$

$$MS_{Error} = SS_{Error} / df_{Error}$$

$$E(MS_{Error}) = \sigma_e^2$$

Now let's look at the ANOVA table and the analyses that can be done with it in more detail.

For greater generality, let

- w = the number of levels of the whole-plot treatment factor,
here $w = 3$ Genotype: A, B, C
- s = the number of levels of the split-plot treatment factor, and
here $s = 4$ 0, 50, 100, 150
- b = the number of blocks. *$b = 4$*

ANOVA Table for the Traditional Split-Plot Design

Source	DF
<i>Blocks</i>	$b - 1$
<i>Genotypes</i>	$w - 1$
<i>Blocks</i> \times <i>Geno</i>	$(b - 1)(w - 1)$
<i>Fert</i>	$s - 1$
<i>Geno</i> \times <i>Fert</i>	$(w - 1)(s - 1)$
<i>Blocks</i> \times <i>Fert</i>	$(b - 1)(s - 1)$
+ <i>Blocks</i> \times <i>Geno</i> \times <i>Fert</i>	+ $(b - 1)(w - 1)(s - 1)$
<i>C.Total</i>	$bws - 1$

combine

ANOVA Table for the Traditional Split-Plot Design

Source	DF
<i>Blocks</i>	$b - 1$
<i>Genotypes</i>	$w - 1$
<i>Blocks</i> \times <i>Geno</i>	$(b - 1)(w - 1)$
<i>Fert</i>	$s - 1$
<i>Geno</i> \times <i>Fert</i>	$(w - 1)(s - 1)$
<i>Error</i>	$w(b - 1)(s - 1)$
<i>C.Total</i>	$bws - 1$

ANOVA Table Sums of Squares

$i = 1, \dots, w$

i : genotype

y_{ijk}

j = fertilizer $j = 1, \dots, s$

Block: $k = 1, \dots, b$

Source	Sum of Squares
Block $(b-1)$	$\sum_{i=1}^w \sum_{j=1}^s \sum_{k=1}^b (\bar{y}_{..k} - \bar{y}_{...})^2$
Geno $(w-1)$	$\sum_{i=1}^w \sum_{j=1}^s \sum_{k=1}^b (\bar{y}_{i..} - \bar{y}_{...})^2$
Block \times Geno $(b-1)(w-1)$	$\sum_{i=1}^w \sum_{j=1}^s \sum_{k=1}^b (\bar{y}_{i.k} - \bar{y}_{i..} - \bar{y}_{..k} + \bar{y}_{...})^2$
Fert	$\sum_{i=1}^w \sum_{j=1}^s \sum_{k=1}^b (\bar{y}_{.j.} - \bar{y}_{...})^2$
Geno \times Fert	$\sum_{i=1}^w \sum_{j=1}^s \sum_{k=1}^b (\bar{y}_{ij.} - \bar{y}_{i..} - \bar{y}_{.j.} + \bar{y}_{...})^2$
Error	$\sum_{i=1}^w \sum_{j=1}^s \sum_{k=1}^b (y_{ijk} - \bar{y}_{i.k} - \bar{y}_{ij.} + \bar{y}_{i..})^2$
C.Total	$\sum_{i=1}^w \sum_{j=1}^s \sum_{k=1}^b (y_{ijk} - \bar{y}_{...})^2$

Simplified ANOVA Table Sums of Squares

$\frac{\text{Sum of squares}}{df} = MS$
 next derive $E(MS)$

Source	Sum of Squares
<i>Block</i>	$ws \sum_{k=1}^b (\bar{y}_{..k} - \bar{y}_{...})^2$
<i>Geno</i>	$sb \sum_{i=1}^w (\bar{y}_{i..} - \bar{y}_{...})^2$
<i>Block</i> \times <i>Geno</i>	$s \sum_{i=1}^w \sum_{k=1}^b (\bar{y}_{i.k} - \bar{y}_{i..} - \bar{y}_{..k} + \bar{y}_{...})^2$
<i>Fert</i>	$wb \sum_{j=1}^s (\bar{y}_{.j.} - \bar{y}_{...})^2$
<i>Geno</i> \times <i>Fert</i>	$b \sum_{i=1}^w \sum_{j=1}^s (\bar{y}_{ij.} - \bar{y}_{i..} - \bar{y}_{.j.} + \bar{y}_{...})^2$
<i>Error</i>	$\sum_{i=1}^w \sum_{j=1}^s \sum_{k=1}^b (y_{ijk} - \bar{y}_{i.k} - \bar{y}_{ij.} + \bar{y}_{i..})^2$
<i>C.Total</i>	$\sum_{i=1}^w \sum_{j=1}^s \sum_{k=1}^b (y_{ijk} - \bar{y}_{...})^2$

$$E(MS_{Geno}) = \frac{sb}{w-1} \sum_{i=1}^w E(\bar{y}_{i..} - \bar{y}_{...})^2$$

Handwritten notes: A red 'X' is next to the fraction. The denominator $w-1$ is circled in red. The entire fraction and sum are enclosed in a blue box.

$$\bar{y}_{i..} = \bar{\mu}_{i.} + \bar{w}_{i.} + \bar{e}_{i..}$$

Handwritten notes: "same $\bar{y}_{...}$ " and "note that b_k will cancel itself".

$$= \frac{sb}{w-1} \sum_{i=1}^w E(\bar{\mu}_{i.} - \bar{\mu}_{..} + \bar{w}_{i.} - \bar{w}_{..} + \bar{e}_{i..} - \bar{e}_{...})^2$$

due to $E(w_{ijk}) = E(e_{ijk}) = 0$ all crossproducts

$$= sb \left\{ \frac{\sum_{i=1}^w (\bar{\mu}_{i.} - \bar{\mu}_{..})^2}{w-1} + E \left[\frac{\sum_{i=1}^w (\bar{w}_{i.} - \bar{w}_{..})^2}{w-1} \right] + E \left[\frac{\sum_{i=1}^w (\bar{e}_{i..} - \bar{e}_{...})^2}{w-1} \right] \right\}$$

$$= sb \frac{w^2}{b}$$

Handwritten notes: This term is highlighted in yellow.

$$= sb \frac{\sum_{i=1}^w (\bar{\mu}_{i.} - \bar{\mu}_{..})^2}{w-1} + sb \frac{\sigma_w^2}{b} + sb \frac{\sigma_e^2}{sb}$$

Handwritten notes: The terms $\frac{\sigma_w^2}{b}$ and $\frac{\sigma_e^2}{sb}$ are crossed out with blue lines.

$$= sb \frac{\sum_{i=1}^w (\bar{\mu}_{i.} - \bar{\mu}_{..})^2}{w-1} + s\sigma_w^2 + \sigma_e^2$$

Handwritten notes: The final result is underlined twice.

involving either $\bar{w}_{i.}$, $\bar{w}_{..}$, $\bar{e}_{i..}$, $\bar{e}_{...}$ will disappear

$$\begin{aligned}
E(MS_{Block \times Geno}) &= \frac{s}{(w-1)(b-1)} \sum_{i=1}^w \sum_{k=1}^b E(\bar{y}_{i \cdot k} - \bar{y}_{i \cdot \cdot} - \bar{y}_{\cdot \cdot k} + \bar{y}_{\cdot \cdot \cdot})^2 \\
&= \frac{s}{(w-1)(b-1)} \sum_{i=1}^w \sum_{k=1}^b E(w_{ik} - \bar{w}_{i \cdot} - \bar{w}_{\cdot k} + \bar{w}_{\cdot \cdot} + \bar{e}_{i \cdot k} - \bar{e}_{i \cdot \cdot} - \bar{e}_{\cdot \cdot k} + \bar{e}_{\cdot \cdot \cdot})^2 \\
&= \frac{s}{(w-1)(b-1)} E \left[\sum_{i=1}^w \sum_{k=1}^b (w_{ik} - \bar{w}_{i \cdot})^2 - 2 \sum_{i=1}^w \sum_{k=1}^b (w_{ik} - \bar{w}_{i \cdot})(\bar{w}_{\cdot k} - \bar{w}_{\cdot \cdot}) \right. \\
&\quad \left. + \sum_{i=1}^w \sum_{k=1}^b (\bar{w}_{\cdot k} - \bar{w}_{\cdot \cdot})^2 + e^2 \text{ sum} \right] \\
&= \frac{s}{(w-1)(b-1)} E \left[\sum_{i=1}^w \sum_{k=1}^b (w_{ik} - \bar{w}_{i \cdot})^2 - w \sum_{k=1}^b (\bar{w}_{\cdot k} - \bar{w}_{\cdot \cdot})^2 + e^2 \text{ sum} \right] \\
&= \frac{s}{(w-1)(b-1)} [w(b-1)\sigma_w^2 - w(b-1)\sigma_w^2/w + E(e^2 \text{ sum})]
\end{aligned}$$

It can be shown that

$$\begin{aligned} E(\text{e}^2 \text{ sum}) &= E \left[\sum_{i=1}^w \sum_{k=1}^b (\bar{e}_{i \cdot k} - \bar{e}_{i \cdot \cdot} - \bar{e}_{\cdot \cdot k} + \bar{e}_{\cdot \cdot \cdot})^2 \right] \\ &= \frac{(w-1)(b-1)}{s} \sigma_e^2. \end{aligned}$$

Putting it all together yields

$$E(\text{MS}_{Block \times Geno}) = \underline{s\sigma_w^2 + \sigma_e^2}.$$

Source	Expected Mean Squares
<i>Block</i>	
<i>Geno</i>	$s\sigma_w^2 + \sigma_e^2 + \frac{sb}{w-1} \sum_{i=1}^w (\bar{\mu}_{i.} - \bar{\mu}_{..})^2$ ✓
<i>Block</i> × <i>Geno</i>	$s\sigma_w^2 + \sigma_e^2$ ✓
<i>Fert</i>	
<i>Geno</i> × <i>Fert</i>	
<i>Error</i>	

The Test for Whole-Plot Factor Main Effects

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~~end lecture 29~~
4-9-25

To test for genotype main effects, i.e.,

$$H_0 : \underline{\bar{\mu}_{1.} = \cdots = \bar{\mu}_{w.}} \iff H_0 : \frac{sb}{w-1} \sum_{i=1}^w (\bar{\mu}_{i.} - \bar{\mu}_{..})^2 = 0,$$

compare $\frac{MS_{Geno}}{MS_{Block \times Geno}}$ to a central F distribution with $\underline{w-1}$ and $\underline{(w-1)(b-1)}$ degrees of freedom.