

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

End lecture 29
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- 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 \times Fert and 2 = Blocks \times Geno \times Fert.

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

$$\frac{\text{Var}^{-1}(MS_1)MS_1 + \text{Var}^{-1}(MS_2)MS_2}{\text{Var}^{-1}(MS_1) + \text{Var}^{-1}(MS_2)} = \frac{\frac{df_1}{2\sigma_e^4}MS_1 + \frac{df_2}{2\sigma_e^4}MS_2}{\frac{df_1}{2\sigma_e^4} + \frac{df_2}{2\sigma_e^4}}$$

$$= \frac{df_1 MS_1 + df_2 MS_2}{df_1 + df_2}$$

$$= \frac{SS_1 + SS_2}{df_1 + df_2}$$

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

here $w = 3$

Genotype : A, B, C

- w = the number of levels of the whole-plot treatment factor,
- $s = 4$ $0, 50, 100, 150$
- s = the number of levels of the split-plot treatment factor, and
- $b = 4$

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

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

$i = 1, \dots, w$

i : genotype

y_{ijk}

$j = \text{fertilizer}$ $j = 1, \dots, s$

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

Simplified ANOVA Table Sums of Squares

$$\frac{\text{Sum of squares}}{df} = MS$$

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

next derive

$E(MS)$

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

~~if~~

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

same $\bar{y}...$

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 $E(\omega_{ik}) = E(e_{ijk}) = 0$ all crossproducts

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

$$= \sigma_w^2 / b$$

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

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

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

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

It can be shown that

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

Putting it all together yields

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

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

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

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.