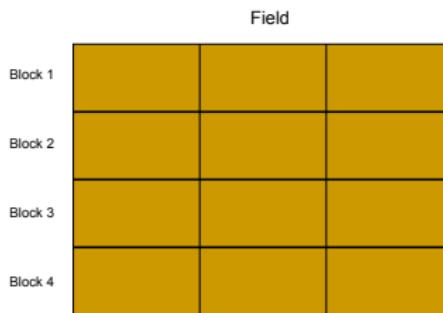
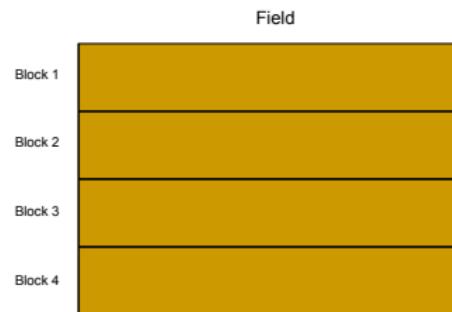
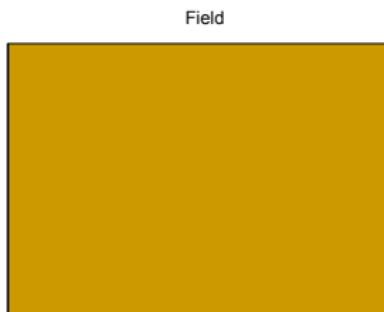


14. Linear Mixed-Effects Models for Data from Split-Plot Experiments

Start with a field, next partition field into blocks, then partition blocks into plots



Randomly Assign Genotypes to Plots within Blocks

Whole plot trt: genotype : A, B, C
Field



| | | | |
|---------|------------|------------|------------|
| Block 1 | Genotype C | Genotype A | Genotype B |
| Block 2 | Genotype B | Genotype A | Genotype C |
| Block 3 | Genotype A | Genotype B | Genotype C |
| Block 4 | Genotype B | Genotype C | Genotype A |

4 replic.
of each
genotype

Partition Each Whole Plot into Split Plots

Second factor: fertilizer – 4 levels 0, 50, 100, 150

| | Field | | |
|---------|------------|------------|------------|
| Block 1 | Genotype C | Genotype A | Genotype B |
| Block 2 | Genotype B | Genotype A | Genotype C |
| Block 3 | Genotype A | Genotype B | Genotype C |
| Block 4 | Genotype B | Genotype C | Genotype A |

Randomly Assign Fertilizer Amounts within Split Plots

randomly assigning fertilizer within each Field whole plot exp. unit

| | Genotype C | | | | Genotype A | | | | Genotype B | | | | |
|---------|------------|-----|-----|-----|------------|-----|-----|-----|------------|-----|-----|---|--|
| Block 1 | 0 | 100 | 150 | 50 | 50 | 100 | 150 | 0 | 150 | 100 | 50 | 0 | |
| Block 2 | Genotype B | | | | Genotype A | | | | Genotype C | | | | |
| | 150 | 100 | 50 | 0 | 0 | 50 | 150 | 100 | 100 | 50 | 150 | 0 | |
| Block 3 | Genotype A | | | | Genotype B | | | | Genotype C | | | | |
| | 100 | 50 | 0 | 150 | 0 | 100 | 150 | 50 | 50 | 100 | 150 | 0 | |
| Block 4 | Genotype B | | | | Genotype C | | | | Genotype A | | | | |
| | 0 | 50 | 100 | 150 | 150 | 100 | 50 | 0 | 50 | 150 | 100 | 0 | |

An Example Split-Plot Experiment

| Field | | | | | | | | | | | | <i>Whole Plot or Main Plot</i> | |
|---------|------------|-----|-----|-----|------------|-----|-----|-----|------------|-----|-----|------------------------------------|-----------------------------------|
| Block 1 | Genotype C | | | | Genotype A | | | | Genotype B | | | | <i>Split Plot or Sub Plot</i> |
| | 0 | 100 | 150 | 50 | 50 | 100 | 150 | 0 | 150 | 100 | 50 | 0 | |
| Block 2 | Genotype B | | | | Genotype A | | | | Genotype C | | | | |
| | 150 | 100 | 50 | 0 | 0 | 50 | 150 | 100 | 100 | 50 | 150 | 0 | |
| Block 3 | Genotype A | | | | Genotype B | | | | Genotype C | | | | |
| | 100 | 50 | 0 | 150 | 0 | 100 | 150 | 50 | 50 | 100 | 150 | 0 | |
| Block 4 | Genotype B | | | | Genotype C | | | | Genotype A | | | | |
| | 0 | 50 | 100 | 150 | 150 | 100 | 50 | 0 | 50 | 150 | 100 | 0 | |

- This experiment has two factors: genotype and fertilizer amount.
- Genotype has levels A, B, and C.
- Fertilizer has levels 0, 50, 100, 150 lbs. N / acre.
- Genotype is called the *whole-plot* (or *main-plot*) factor because its levels are randomly assigned to whole plots (main plots).
- Fertilizer is called the *split-plot* factor because its levels are randomly assigned to split plots within each whole plot.

Experimental Units in Split-Plot Designs

- Whole plots are the *whole-plot experimental units* because the levels of the whole-plot factor (genotype) are randomly assigned to whole plots.
- The split-plots are the *split-plot experimental units* because the levels of the split-plot factor (amount of fertilizer) are randomly assigned to split plots within each whole plot.
- Thus, we have two different sizes of experimental units in split-plot experimental designs.

Same Treatment Structure in an RCBD

| | Field | | | | | | | | | | | | | |
|---------|-------|-----|-----|-----|-----|----|-----|-----|-----|-----|-----|-----|-----|--|
| Block 1 | B | B | A | C | B | C | A | A | C | B | C | A | | |
| | 100 | 0 | 0 | 100 | 150 | 50 | 50 | 150 | 150 | 50 | 0 | 100 | 100 | |
| Block 2 | A | A | C | A | B | B | C | C | A | C | B | B | | |
| | 150 | 0 | 50 | 50 | 100 | 50 | 100 | 0 | 100 | 150 | 150 | 0 | 150 | |
| Block 3 | C | A | A | B | B | B | A | C | A | C | C | B | | |
| | 0 | 0 | 100 | 100 | 50 | 0 | 150 | 50 | 50 | 150 | 100 | 150 | | |
| Block 4 | B | C | B | A | C | A | B | C | B | C | A | A | | |
| | 0 | 150 | 50 | 150 | 100 | 0 | 150 | 50 | 100 | 0 | 100 | 50 | | |

Same Treatment Structure in an CRD

Field

| | | | | | | | | | | | |
|---------|----------|----------|----------|----------|----------|----------|----------|----------|----------|----------|----------|
| B 50 | B 0 | A 150 | B 100 | A 100 | C 150 | A 50 | B 0 | A 50 | C 100 | C 0 | C 100 |
| A 50 | A 0 | C 50 | B 50 | B 150 | B 50 | A 0 | C 0 | A 100 | C 50 | B 150 | B 0 |
| C 0 | A 0 | A 100 | A 150 | A 0 | B 0 | A 150 | B 150 | A 50 | B 150 | C 100 | A 100 |
| B 50 | B 100 | B 100 | C 150 | C 100 | C 50 | A 150 | C 50 | C 150 | C 0 | C 150 | B 100 |

Why Use a Split-Plot Design?

- Split-plot designs usually arise because logistical constraints make a CRD or RCBD impractical.
- For example, it may be easier to change from one fertilizer level to another as a tractor drives through a field, while it may be more difficult to change from planting one genotype to planting another.
- In the engineering literature, split-plot designs are sometimes called designs with hard-to-change factors.

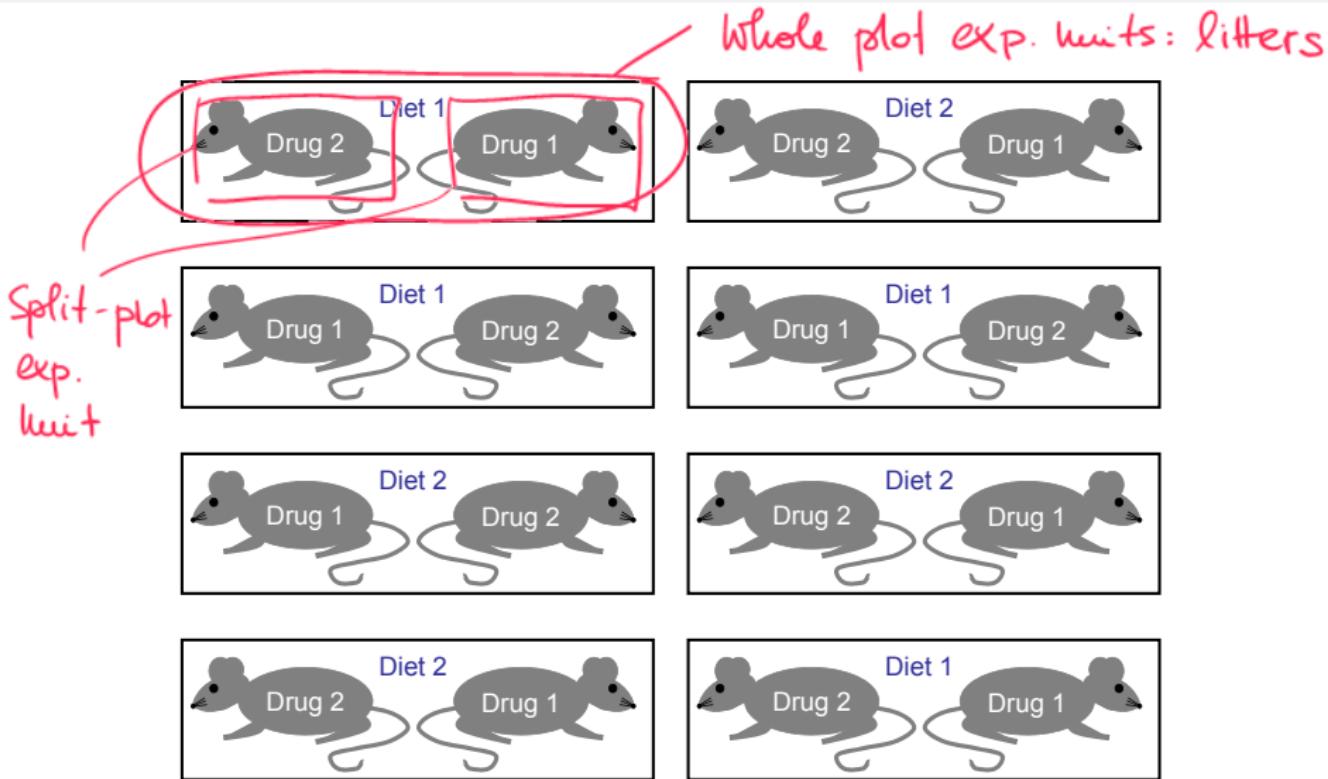
Recognizing Designs with Split-Plot Structures

- Many variations on split-plot designs are used for practical reasons.
- Examples include split-split-plot designs and split-block designs, but the names of these designs are not so important.
- Pay close attention to the experimental unit to which the levels of each factor are randomly assigned to recognize split-plot-like design structures.

Split-plot designs may not involve plots of land.

- Suppose eight pairs of mice from eight litters are housed in eight cages so that each cage holds two mice from the same litter.
- Suppose diets 1 and 2 are randomly assigned to the litters with four litters per diet.
- Within each cage, suppose drugs 1 and 2 are randomly assigned to the mice with one mouse per drug.

A Split-Plot Experimental Design



- Diet is the whole-plot treatment factor.
 - Litters are the whole-plot experiment units.
-
- Drug is the split-plot treatment factor.
 - Mice are the split-plot experiment units.
-

Mouse example:

^{indep.}
 $\sum 8$ litters

Diet $i = 1, 2$, Drug $j = 1, 2$, Litter $k = 1, 2, 3, 4$ (within each Diet i)

$$y_{ijk} = \mu + \alpha_i + \beta_j + \gamma_{ij} + \ell_{ik} + e_{ijk} \quad (i = 1, 2; j = 1, 2; k = 1, \dots, 4)$$

$\mu + \alpha_i + \beta_j + \gamma_{ij}$ = mean for Diet i and Drug j

ℓ_{ik} = random litter effect = whole-plot exp. unit random effect

e_{ijk} = random error effect = split-plot exp. unit random effect

$$\begin{array}{c}
 \mathbf{y} = \begin{bmatrix} y_{111} \\ y_{121} \\ y_{112} \\ y_{122} \\ y_{113} \\ y_{123} \\ y_{114} \\ y_{124} \\ y_{211} \\ y_{221} \\ y_{212} \\ y_{222} \\ y_{213} \\ y_{223} \\ y_{214} \\ y_{224} \end{bmatrix} & \beta = \begin{bmatrix} \mu \\ \alpha_1 \\ \alpha_2 \\ \beta_1 \\ \beta_2 \\ \gamma_{11} \\ \gamma_{12} \\ \gamma_{21} \\ \gamma_{22} \end{bmatrix} & \mathbf{u} = \begin{bmatrix} \ell_{11} \\ \ell_{12} \\ \ell_{13} \\ \ell_{14} \\ \ell_{21} \\ \ell_{22} \\ \ell_{23} \\ \ell_{24} \end{bmatrix} & \mathbf{e} = \begin{bmatrix} e_{111} \\ e_{121} \\ e_{112} \\ e_{122} \\ e_{113} \\ e_{123} \\ e_{114} \\ e_{124} \\ e_{211} \\ e_{221} \\ e_{212} \\ e_{222} \\ e_{213} \\ e_{223} \\ e_{214} \\ e_{224} \end{bmatrix}
 \end{array}$$

fixed random
 / /
 } interactions

Overall mean \bar{M}

$$\mathbf{X} = \left[\begin{matrix} \mathbf{1}_{16 \times 1}, & \boxed{\mathbf{I}_{2 \times 2} \otimes \mathbf{1}_{8 \times 1}}, & \boxed{\mathbf{1}_{8 \times 1} \otimes \mathbf{I}_{2 \times 2}}, & \mathbf{I}_{2 \times 2} \otimes \mathbf{1}_{4 \times 1} \otimes \mathbf{I}_{2 \times 2} \end{matrix} \right]$$

reflects two diets

reflects drug

interactions

$$\mathbf{Z} = \mathbf{I}_{8 \times 8} \otimes \mathbf{1}_{2 \times 1} = \begin{pmatrix} 1 \\ 1 \end{pmatrix} \text{ two mice within each litter}$$

8 indep. litters

$$\mathbf{y} = \mathbf{X}\boldsymbol{\beta} + \mathbf{Z}\boldsymbol{u} + \boldsymbol{\epsilon}$$

$$\left\{ \begin{bmatrix} u \\ e \end{bmatrix} \sim \mathcal{N} \left(\begin{bmatrix} 0 \\ 0 \end{bmatrix}, \begin{bmatrix} \sigma_\ell^2 \mathbf{I} & \mathbf{0} \\ \mathbf{0} & \sigma_e^2 \mathbf{I} \end{bmatrix} = \begin{bmatrix} G & \mathbf{0} \\ \mathbf{0} & R \end{bmatrix} \right) \right.$$

$$\begin{aligned}
 \text{Var}(\mathbf{Z}\mathbf{u}) &= \mathbf{Z}\mathbf{G}\mathbf{Z}^\top = \sigma_\ell^2 \mathbf{Z}\mathbf{Z}^\top \\
 &= \sigma_\ell^2 \left[\underset{8 \times 8}{\mathbf{I}} \otimes \underset{2 \times 1}{\mathbf{1}} \right] \left[\underset{8 \times 8}{\mathbf{I}} \otimes \underset{2 \times 1}{\mathbf{1}} \right]^\top \\
 &= \sigma_\ell^2 \left[\underset{8 \times 8}{\mathbf{I}} \otimes \underset{2 \times 2}{\mathbf{1}\mathbf{1}^\top} \right] \\
 &= \text{Block Diagonal with blocks: } \begin{bmatrix} \sigma_\ell^2 & \sigma_\ell^2 \\ \sigma_\ell^2 & \sigma_\ell^2 \end{bmatrix}
 \end{aligned}$$

$$\underline{\text{Var}(\mathbf{y})} = \mathbf{ZGZ}^\top + \mathbf{R} = \sigma_\ell^2 \mathbf{I}_{8 \times 8} \otimes \mathbf{11}^\top_{2 \times 2} + \sigma_e^2 \mathbf{I}$$

= Block Diagonal with blocks:

$$\begin{bmatrix} \frac{\sigma_\ell^2 + \sigma_e^2}{\sigma_\ell^2} & \sigma_\ell^2 \\ \sigma_\ell^2 & \underline{\sigma_\ell^2 + \sigma_e^2} \end{bmatrix}$$

end lecture 28

4-4-25