## Chapter 21: <u>Randomized Complete Block Design (RCBD)</u>

**Blocking:** 

• Consider a situation in which we have one factor of interest.

• If the experimental units are <u>not homogeneous</u>, there may be a great deal of variation in the response values, even among those having the same level of the factor.

• This leads to a large "experimental error variance" (measured by  $\sigma^2$ ).

**Example** (Chemistry experiment): Response: reaction rate of chemical agent Factor: Chemical agent (5 treatments)

- Only 5 chemical agents can be tested each day.
- A different technician conducts the experiment each day.
- Wide variability among technicians  $\rightarrow$

**Solution:** To account for variation due to technician, we treat technician as a "blocking factor".

• We separate the experimental units into groups called <u>blocks</u> and then assign each treatment

**Example 2**: Interested in the effect of three different teaching methods on student learning.

<u>Response</u>: Improvement on a test score <u>Treatments</u>: 3 methods (lecture; discussion; student presentation)

• But students will be taught be different instructors – we're not interested in the instructor effect, but it adds variability.

Solution:

**Example 3**: Lab animals of the same species given different diets and weight gain from birth is measured. **Experimental Units**: animals **Response**: Weight gain <u>Treatments</u>: diets

**Blocks:** 

<u>Goal</u>: Blocks should be chosen so that units in the same block are \_\_\_\_\_\_ but units in different blocks are

\_\_\_\_\_•

• When we use blocks, it reduces the unexplained variation – the

• This means there is more precision in estimating

#### **Model for RCBD**

• Assume there are *n*<sub>b</sub> blocks and *r* treatments.

**Equation:** 

• If the treatments are fixed,

• Often the blocks are random (they are randomly selected from a large population). In this case,

• If the blocks are fixed, then

• With this model, we separate the total variation in *Y* into three sources:

• The fitted values for the RCBD analysis are

and the residuals are

## ANOVA Table for RCBD

• Note that the experimental error variation is measured by the SS for block × treatment interaction. If this interaction is significant (we can check this using Tukey's additivity test), we could

(1)

or (2)

• In the case of (2), when blocks are random we could still test for treatment effects using (see p. 1063)

but we would not be able to test for block effects.

<u>Example</u> (Executive data): • Executives quantified a risk premium using 3 methods (utility, worry, comparison) <u>Subjects</u>: 15 executives <u>Response</u>: Degree of Confidence in Risk Premium <u>Treatments</u>: the 3 Methods (U, W, C) <u>Blocks</u>: Age Group of Executives (here, 5 \_\_\_\_\_ blocks) **Randomization scheme:** 

**ANOVA Table from SAS:** 

• We may test whether the mean confidence is equal for the three methods:

• If block effects are of interest, we could test for significant block effects by comparing

• SAS gives the P-value for this test.

## **Some Model Diagnostics**

• We may check model assumptions using

**Example** (Executive data):

# **Further Analysis of Treatment Effects**

• We may further analyze differences among the treatment means using contrast or multiple comparisons.

**Example** (Executive data):

**<u>Note</u>:** If the blocks are considered random, we still use: to test for significant variation across blocks.

- In this case we are testing:
- We may wish to do inference about

• When possible, more precise information in a RCBD can be attained (<u>even</u> if there is block × treatment interaction) if we have <u>replication within each block</u> (i.e., each treatment is repeated  $d \ge 2$  times within each block).

- This is called a generalized random block design.
- Each block contains *dr* units.
- The analysis is identical to the two-factor ANOVA.
- MSE can be calculated and serves as an estimate of

• The denominator MS for F-tests again depends on which factors (here, blocks and treatments) are fixed or random (Table 25.6, page 1053 is again useful).

 Example (Task Completion):

 <u>Response</u>: Time to Complete Task

 <u>Blocks</u>: Gender (2 levels, considered \_\_\_\_\_)

 <u>Treatments</u>: Distraction Levels (2 levels, considered \_\_\_\_\_)

#### **Balanced Incomplete Block Design (BIBD)**

• When our resources are too limited to use a RCBD, we may use a BIBD.

• This block design is called <u>incomplete</u> because not all the treatments appear in each block.

• It is called <u>balanced</u> because each treatment appears in the same block with every other treatment the same number of times.

**Example:** r = 4 treatments (A,B,C,D) and  $n_T = 12$  exper. units

• Suppose we can divide the units into 6 blocks.

• If we divide the units into 4 blocks, then:

• In practice, once we pick a design, we would randomly arrange the treatments within each block.

Advantages of BIBD:

• Can use block design when block size is smaller than the number of treatments.

• Same precision in estimating each treatment effect

• Scheffe and Tukey procedures can be used when the design is balanced.

#### **Disadvantages of BIBD:**

• BIBDs don't exist for every combination of number of treatments, number of blocks, and block size.

[A list of BIBDs for selected combinations of r,  $n_b$ , and  $r_b$  (block size) is given in Table B.15 (p. 1345-1347).]

• Must assume no treatment × block interaction.

• Analysis is more complicated than for RCBD.

• With BIBDs, out F-tests require the Reduced vs. Full Model approach with indicators for blocks and treatments (see p. 1177-1179).

• Can do this correctly in SAS (with PROC MIXED) or in R (see examples).

#### Example (Table 28.2 data):

• We wish to compare mean consumer ratings (the response) across five formulations of a cereal (the treatments).

• Since there is probably variation across consumers, we use the <u>ten</u> consumers as the blocks.

• For quality purposes, each consumer can only rate <u>three</u> formulations (must use <u>incomplete</u> design).

# **<u>SAS example</u>**: If blocks are considered a random sample from a population of consumers, we use a

**Example:** 

• If blocks are considered fixed, we can include them in the MODEL statement and omit the RANDOM statement.

#### Latin Square Designs

• These are efficient designs for situations in which we have <u>two</u> blocking factors.

**Example:** (Drug study)

<u>Subjects</u>: Patients <u>Response</u>: Cholesterol Reduction <u>Treatments</u>: 4 different drugs <u>Blocking Factors</u>: Age Group (4 levels), Blood Pressure Status (Low, Medium, High, Extreme)

• For a RCBD, even if we only have 1 observation per cell, we'd need

• May be infeasible to obtain that many patients.

• A Latin Square Design is arranged so that one blocking factor is the <u>row factor</u> and the other blocking factor is the <u>column factor</u>.

In a Latin Square:

• In the example above, a Latin Square would require

# Advantages: (1)

(2)

# Disadvantages:

(1)

(2)

(3)

# **<u>Randomization Scheme</u>**: For your particular value of *r*:

- (1)
- (2)
- (3)

Example (r = 4):

**Example:** Assessing effect of background music on bank tellers' productivity:

**Experimental Units:** Working Days for Bank Crew **<u>Response</u>:** Productivity Rating <u>Treatments</u>: Type of Music (A = Slow Vocal, B= Medium Vocal, C = Fast Vocal, D = Medium Instrumental, E= Fast Instrumental) <u>Row Factor</u>: Week (1, 2, 3, 4, 5) <u>Column Factor</u>: Weekday (M, Tu, W, Th, F)

• A Latin Square design can complete the experiment in

After Randomization, the Design is:

**Model for Latin Square Design** 

#### **ANOVA table for Latin Square Model**

• Formulas for the sums of squares given on p. 1189.

• To test for significant differences in mean response among the treatments, we test:

by comparing

SAS example (music data): Using  $\alpha = 0.01$ ,

• Specific treatment comparisons can be investigating using contrasts or multiple comparisons.

**Music example:** 

• If desired, effects of blocking factors can be tested using appropriate F-statistics from the ANOVA table.

Model assumptions may be checked via:

- (1)
- (2)
- (3)

• If the treatments are random or if either blocking factor has random levels, the usual adjustments are made to the model.

• Section 28.7 discusses Latin Square designs with replication – more than one observation per cell.

• In some cases, multiple experimental units can be given the same treatment-row-column combination.

• In that case, we have replication within each cell, and we can formally test the fit of the additive (no-interaction) model using a lack-of-fit F-test (see example with Table 28.8 data).

• In other cases (such as the music/productivity experiment?), we cannot obtain multiple observations in each treatment-rowcolumn combination.

• In that case, we can augment the experiment with more data by using multiple Latin Squares (each arrangement is selected independently).

<u>Note</u>: If the treatments in a Latin Square are factorial with two factors A and B, then the treatment SS can be decomposed as usual (SSTR = SSA + SSB + SSAB), to assess separately the effects of factors A and B and their interaction.