

Randomized Complete Block Design (RCBD)

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Introduction

Blocking is used to handle nuisance factors (effects on responses but not of interest) that are known and controllable. If all treatments are applied within each block and are compared within blocks, the design is (complete) blocking. Randomization is restricted because only occurs within blocks.

Pros: Eliminate between-block variation from experimental error variance.

Cons: Degrees of freedom

Statistical Model: Two-Way ANOVA

$y_{ij} = \mu + \tau_i + \beta_j + \epsilon_{ij}$, $i = 1, 2, \dots, a$ and $j = 1, 2, \dots, b$ where a is the number of treatments and b is the number of experimental units.

μ = grandmean, τ_i = i th treatment effect, β_j = j th block effect, $\epsilon_{ij} \sim N(0, \sigma^2)$

$$\sum_{i=1}^a \tau_i = 0; \sum_{j=1}^b \beta_j = 0$$

Parameter Estimates

$$\hat{\mu} = \bar{y}_{..}; \hat{\tau}_i = \bar{y}_{i.} - \bar{y}_{..}; \hat{\beta}_j = \bar{y}_{.j} - \bar{y}_{..}; \hat{\epsilon}_{ij} = y_{ij} - \bar{y}_{i.} - \bar{y}_{.j} + \bar{y}_{..}$$

Sum of Squares

$$SS_T = SS_{Treatment} + SS_{Block} + SS_E = \sum \sum (y_{ij} - \bar{y}_{..})^2 = (\sum \sum y_{ij}^2) - y_{..}^2/N \text{ with df} = ab - 1$$

$$SS_{Treatment} = b \sum (\bar{y}_{i.} - \bar{y}_{..})^2 = b \sum \hat{\tau}_i^2 = \frac{1}{b} (\sum y_{i.}^2) - y_{..}^2/N \text{ with df} = a - 1$$

$$SS_{Block} = a \sum (\bar{y}_{.j} - \bar{y}_{..})^2 = a \sum \hat{\beta}_j^2 = \frac{1}{a} (\sum y_{.j}^2) - y_{..}^2/N \text{ with df} = b - 1$$

$$SS_E = \sum \sum (y_{ij} - \bar{y}_{i.} - \bar{y}_{.j} + \bar{y}_{..})^2 = \sum \sum \hat{\epsilon}_{ij}^2 \text{ with df} = (a-1)(b-1)$$

Mean Squared

$$MS_{Treatment} = SS_{Treatment}/(a-1) \text{ and expected value is } \sigma^2 + \sigma_{\tau\beta}^2 + b \sum_{i=1}^a \tau_i^2/(a-1)$$

$$MS_{Block} = SS_{Block}/(b-1) \text{ and expected value is } \sigma^2 + a \sum_{j=1}^b \beta_j^2/(b-1)$$

$$MS_E = SS_E/((a-1)(b-1)) = \sigma^2 + \sigma_{\tau\beta}^2 \text{ and expected value is } \sigma^2$$

$$\sigma_{\tau\beta}^2 = 0 \text{ if there is no interaction.}$$

Hypothesis Testing

Usually tests treatment effect, block effects are not usually of interest.

$H_0 : \tau_1, \tau_2, \dots, \tau_a = 0$ vs. H_1 : at least one is not

$$F_0 = \frac{MS_{Treatment}}{MS_E} = \frac{SS_{Treatment}/(a-1)}{SS_E/((a-1)(b-1))} \sim F_{\alpha, a-1, (a-1)(b-1)}$$

ANOVA Table

■ **TABLE 4.2**

Analysis of Variance for a Randomized Complete Block Design

Source of Variation	Sum of Squares	Degrees of Freedom	Mean Square	F_0
Treatments	$SS_{Treatments}$	$a - 1$	$\frac{SS_{Treatments}}{a - 1}$	$\frac{MS_{Treatments}}{MS_E}$
Blocks	SS_{Blocks}	$b - 1$	$\frac{SS_{Blocks}}{b - 1}$	
Error	SS_E	$(a - 1)(b - 1)$	$\frac{SS_E}{(a - 1)(b - 1)}$	
Total	SS_T	$N - 1$		

Figure 1: RCBD Table

Checking Assumptions

Checking Normality

Histogram, QQ plot of residuals, Shapiro-Wilk Test

Checking Constant Variance

Residual plots: residuals vs (\hat{y}_{ij} , blocks, treatments)

Checking Additivity (no interaction between treatment and block effects)

Residuals vs \hat{y}_{ij} , Tukey's One-degreee Freedom Test of Non-additivity

Tukey's Test for Non-additivity

To check for interaction fully needs $(a - 1)(b - 1)$ degrees of freedom (needs replicates).

$$y_{ij} = \mu + \tau_i + \beta_j + \gamma\tau_i\beta_j + \epsilon_{ij}$$

$$H_0 : \gamma = 0 \text{ vs. } H_1 : \gamma \neq 0$$

$$SS_N = \frac{[\sum_i \sum_j y_{ij} y_{i.} y_{.j} - y_{..} (SS_{T_{rt}} + SS_{B_{lk}} + y_{..}^2 / ab)]^2}{ab SS_{T_{rt}} SS_{B_{lk}}}, \text{ with df} = 1$$

$$SS'_E = SS_E - SS_N, \text{ with df} = (a-1)(b-1) - 1$$

$$F_0 = \frac{SS_N}{SS'_E / [(a-1)(b-1)-1]} \sim F_{\alpha, 1, (a-1)(b-1)-1}$$

Note: Convenient Procedure Fit additive model, acquire \hat{y}_{ij} , $q_{ij} = \hat{y}_{ij}^2$, then fit the model with q_{ij} , and then check type III SS for q_{ij}

Post-ANOVA Treatments Comparison

Similar to Completely Randomized Design (CRD) procedure. Replace n by b and degrees of freedom for error is $(b-1)(a-1)$