# Experimental Design Note 5 Balanced Incomplete Block Design (BIBD)

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## Review: Nuisance Factor I

Nuisance Factor (may be present in experiment)

- has effect on response but its effect is not of interest.
- If  $unknown \rightarrow Protecting experiment through randomization.$
- If known (measurable) but uncontrollable → Analysis of Covariance.
- $lue{}$  If known and controllable ightarrow Blocking.

# Example: Penicillin Experiment I

In this experiment, four penicillin manufacturing processes (A, B, C, and D) were being investigated. Yield was the response. It was known that an important raw material, corn steep liquor, was quite variable. The experiment and its results were given below:

	blend 1	blend 2	blend 3	blend 4	blend 5
Α	89 <sub>1</sub>	844	812	87 <sub>1</sub>	79 <sub>3</sub>
В	883	77 <sub>2</sub>	87 <sub>1</sub>	92 <sub>3</sub>	814
C	97 <sub>2</sub>	92 <sub>3</sub>	874	89 <sub>2</sub>	80 <sub>1</sub>
D	944	$79_{1}$	85 <sub>3</sub>	844	882

Blend is a nuisance factor, treated as a block factor;

## Example: Penicillin Experiment II

- (Complete) Blocking: all the treatments are applied within each block, and they are compared within blocks.
- Advantage: Eliminate blend-to-blend (between-block) variation from experimental error variance when comparing treatments.
- Cost: degree of freedom.

# Catalyst Experiment I

Four catalysts are being investigated in an experiment. The experimental procedure consists of selecting a batch raw material, loading the pilot plant, applying each catalyst in a separate run and observing the reaction time. The batches of raw material are considered as blocks, however each batch is only large enough to permit three catalysts to be run.

catalyst	1	2	3	4	y <sub>i</sub> .
1	73	74	-	71	218
2	-	75	67	72	214
3	73	75	68	-	216
4	75	-	72	75	222
у. <sub>j</sub>	221	224	207	218	870= <i>y</i>

# Balanced Incomplete Block Design (BIBD) I

#### **BIBD** Properties

- There are a treatments and b blocks.
- Each block contains k (different) treatments.
- Each treatment appears in *r* blocks.
- **Each** pair of treatments appears together in  $\lambda$  blocks.
- a, b, k, r, and  $\lambda$  are not independent.
  - Arr N = ar = bk, where N is the total number of runs;
  - $\lambda(a-1)=r(k-1);$ 
    - for any fixed treatment  $i_0$
    - two different ways to count the total number of pairs including treatment  $i_0$  in the experiment.
    - a-1 possible pairs, each appears in  $\lambda$  blocks, so  $\lambda(a-1)$ ;

# Balanced Incomplete Block Design (BIBD) II

- treatment  $i_0$  appears in r blocks. Within each block, there are k-1 pairs including  $i_0$ , so r(k-1).
- $b \ge a$  (a brainteaser for math/stat students).
- Nonorthogonal design
- Reasoning for integer  $\lambda$ :
  - Each treatment is assigned to *r* blocks.
  - Each of those r blocks has k-1 remaining positions
  - Those r(k-1) positions must be evenly shared among the remaining a-1 treatments.
- Analyses are based on Intra- and Inter-Block Information

# Balanced Incomplete Block Design (BIBD) III

#### Example 1.

	ŀ	olock	<		$\mathcal{N}$	
treatment	1	2	3			
А	Α	-	Α	1	0	1
В	В	В	-	1	1	0
C	-	С	С	0	1	1

Incidence Matrix:  $\mathcal{N} = (n_{ij})_{a \times b}$  where  $n_{ij} = 1$ , if treatment i is run in block j; j otherwise.

In Example 1, a=3, b=3, k=2, r=2,  $\lambda=1$  where  $\lambda=$ the number of times each pair of treatments appears in the same block.

# Balanced Incomplete Block Design (BIBD) IV

#### Example 2

block												
Treatment	1	2	3	4	5	6			$\mathcal{N}$			
A	Α	Α	Α	-	-	-	1	1	1	0	0	0
В	В	-	-	В	В	_	1	0	0	1	1	0
С	-	C	-	C	-	C	0	1	0	1	0	1
D	-	-	D	-	D	D	0	0	1	0	1	1

$$a = 4$$
,  $b = 6$ ,  $k = 2$ ,  $r = 3$ , and  $\lambda = 1$ .

# Balanced Incomplete Block Design (BIBD) V

	block									
Treatment	1	2	3	4	5	6	7	8	9	10
A	Α	Α	Α			Α	Α	Α		
В	В	В		В		В			В	В
C	C			C	C		C	C	C	
D			D	D	D	D	D			D
Е		Ε	Ε		Ε			Ε	Ε	E

$$a = 5$$
,  $b = 10$ ,  $k = 3$ ,  $r = 6$ , and  $\lambda = 3$ .

## BIBD: Statistical Model I

Statistical Model

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

for 
$$i = 1, 2, \dots, a$$
;  $j = 1, 2, \dots, b$ .

- additive model (without interaction).
- Not all  $y_{ii}$  exist because of incompleteness.
- Usual treatment and block restrictions:  $\sum_i \tau_i = 0$ ;  $\sum_i \beta_i = 0$ .
- Nonorthogonality of treatments and blocks.

## Use Type III Sums of Squres and Ismeans

## Model Estimates I

Least squares estimates for parameters

$$\hat{\mu} = \frac{y..}{N}; \ \hat{\tau}_i = \frac{kQ_i}{\lambda a}; \ \hat{\beta}_j = \frac{rQ_j'}{\lambda b}$$

where

$$\begin{aligned} Q_i &= y_{i\cdot} - \frac{1}{k} \sum_j n_{ij} y_{\cdot j}; \quad Q_j^{'} &= y_{\cdot j} - \frac{1}{r} \sum_i n_{ij} y_{i\cdot} \\ var(Q_i) &= var(y_{i\cdot}) + var\left(\frac{1}{k} \sum_j n_{ij} y_{\cdot j}\right) - 2cov\left(y_{i\cdot}, \frac{1}{k} \sum_j n_{ij} y_{\cdot j}\right) \\ &= r\sigma^2 + \frac{r}{k^2} k\sigma^2 - \frac{2}{k} r\sigma^2 = \frac{(k-1)r}{k}\sigma^2 \end{aligned}$$

■ 
$$var(\hat{\tau}_i) = \left(\frac{k}{\lambda a}\right)^2 var(Q_i) = \left(\frac{k}{\lambda a}\right)^2 \frac{(k-1)r}{k} \sigma^2 = \frac{k(a-1)}{\lambda a^2} \sigma^2$$
;  
 $var(\hat{\tau}_i - \hat{\tau}_j) = \frac{2k\sigma^2}{\lambda a}$ ;

## Model Estimates I

Source of	Sum of	Degrees of	Mean	F
Variation	Squares	Freedom	Square	
Block	SS <sub>Block</sub>	b - 1	MS <sub>Block</sub>	
Treatment(Adj.)	SS <sub>Treatment(Adj.)</sub>	a-1	$MS_{Treatment(Adj.)}$	$MS_{Treatment(Adj.)}/MSE$
Error	SSE	N - a - b + 1	MSE `	, , ,
Total	$SS_T$	N - 1		

where 
$$SS_T = \sum_{i} \sum_{j} y_{ij}^2 - y_{..}^2 / N$$
;  $SS_{Block} = \frac{1}{k} \sum_{j} y_{.j}^2 - y_{..}^2 / N$ 

■ *SS*<sub>Treatment</sub> needs adjustment for incompleteness

$$Q_i = y_i - \frac{1}{k} \sum_j n_{ij} y_{\cdot j}$$

where  $n_{ij} = 1$  if trt i in block j; = 0 otherwise.

trt i's total minus trt i's block averages.

## Model Estimates II

- *SSE* by subtraction
- If  $F_0 > F_{\alpha,a-1,N-a-b+1}$ , then reject  $H_0$ .

## Mean Tests and Contrasts

- Must compute adjusted means (Ismeans)
- Adjusted mean is  $\hat{\mu} + \hat{\tau}_i$
- Standard error of adjusted mean is  $\sqrt{MSE\left(\frac{k(a-1)}{\lambda a^2} + \frac{1}{N}\right)}$
- Contrasts based on adjusted treatment totals For a contrast:  $\sum_i c_i \mu_i$ Its estimate:  $\sum_i c_i \hat{\tau}_i = \frac{k}{\lambda a} \sum_i c_i Q_i$ Contrast sum of squares:

$$SS_C = \frac{k \left(\sum_{i=1}^a c_i Q_i\right)^2}{\lambda a \sum_{i=1}^a c_i^2}$$



# Pairwise Comparison

- Pairwise comparison  $\tau_i \tau_i$ :
  - Bonferroni:

$$CD = t_{lpha/2m, ar-a-b+1} \sqrt{MSE rac{2k}{\lambda a}}$$

Tukey:

$$CD = rac{q_{lpha}(a, ar - a - b + 1)}{\sqrt{2}} \sqrt{MSE} rac{2k}{\lambda a}$$

In SAS, there are two procedures: **PROC ANOVA** and **PROC GLM**. PROC ANOVA is for balanced data and PROC GLM is for both balanced or unbalanced data.

See BIBD SAS.