

STAT 222 Lecture 20-21

Incomplete Block Designs

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Coverage

Chapter 11 Incomplete Block Designs

- ▶ Balanced Incomplete Block Designs (BIBD)
- ▶ Skip Section 11.3.2, 11.3.3, 11.4.5, 11.4.6 on Group Divisible Designs and Cyclic Designs

Incomplete Block Designs

Recall for a randomized complete block design (RCBD) of g treatments, the size k of each block has to be g (or multiples of g). Each treatment appear the same number of time(s) in a block.

In practice, the natural size of a block might not be equal to and is often smaller than the numbers of treatments ($k < g$).

We cannot include every treatment to every block.

We then have *Incomplete Block Design (IBD)*.

IBD is more difficult to analyze than complete block designs, but sometimes it's inevitable.

An Example We Must Use Incomplete Blocks

Eye irritation can be reduced with eyedrops. Three brands of eyedrops are to be compared for their ability to reduce eye irritation.

As there is a strong individual effect, subjects should be used as blocks.

If a subject can only be tested in one treatment period, the researchers can apply one brand of drop in the left eye and another brand in the right eye. The natural block size is limited to $k = 2$.

The study is force into incomplete blocks, with

$$\begin{array}{ccc} k = 2 & < & 3 = g \\ \text{(block size)} & < & \text{(number of treatments)} \end{array}$$

Example — A Marketing Psychology Experiment

- ▶ Goal: comparing 5 commercial ads: A, B, C, D, E
- ▶ Response: subjects' rating of a commercial after watching it
- ▶ A subject can watch multiple ads. A subject is a block
- ▶ Can use a RCBD if all subjects can watch all the 5 ads
- ▶ However, subjects may lose patience after watching too many ads, and they may forget the first few ads they see. Their response will be less accurate.
- ▶ To ensure the quality of the response of subjects, we may restrict the number of ads each subject watch to, say, 3. The block size is limited to $k = 3$.

Subject									
1	2	3	4	5	6	7	8	9	10
A	B	E	A	C	D	B	E	D	C
B	D	A	C	A	E	C	B	E	D
C	A	B	D	E	A	D	C	B	E

Some Poor Incomplete Block Designs (1)

Block				
1	2	3	4	5
A	B	C	D	E
A	B	C	D	E
A	B	C	D	E

- ▶ What's the drawback of the design above?

Some Poor Incomplete Block Designs (1)

Block				
1	2	3	4	5
A	B	C	D	E
A	B	C	D	E
A	B	C	D	E

- ▶ What's the drawback of the design above?
 - ▶ block effect and treatment effects are confounded

Some Poor Incomplete Block Designs (1)

Block				
1	2	3	4	5
A	B	C	D	E
A	B	C	D	E
A	B	C	D	E

- ▶ What's the drawback of the design above?
 - ▶ block effect and treatment effects are confounded
- ▶ To eliminate of block effects, better compare treatments **within a block**.

Some Poor Incomplete Block Designs (1)

Block				
1	2	3	4	5
A	B	C	D	E
A	B	C	D	E
A	B	C	D	E

- ▶ What's the drawback of the design above?
 - ▶ block effect and treatment effects are confounded
- ▶ To eliminate of block effects, better compare treatments **within a block**.
- ▶ **No treatment should appear twice in any block** as they contributes nothing no **within block comparisons**.

Some Poor Incomplete Block Designs (2)

Block			
1	2	3	4
A	B	E	E
B	C	F	F
C	D	G	G

Based on the model

$$y_{ij} = \mu + \underset{\text{(treatment)}}{\alpha_i} + \underset{\text{(block)}}{\beta_j} + \underset{\text{(i.i.d. } N(0, \sigma^2))}{\varepsilon_{ij}}$$

can one find an unbiased estimate for

- ▶ $\alpha_A - \alpha_B$?
- ▶ $\alpha_A - \alpha_D$?
- ▶ $\alpha_A - \alpha_E$?

Some Poor Incomplete Block Designs (2) — A v.s. B

Block			
1	2	3	4
A	B	E	E
B	C	F	F
C	D	G	G

Based on the model

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Some Poor Incomplete Block Designs (2) — A v.s. B

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A	B	E	E
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C	D	G	G

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can one find an unbiased estimate for $\alpha_A - \alpha_B$?

Yes, $y_{A1} - y_{B1}$ is an unbiased estimate for $\alpha_A - \alpha_B$ since

$$\begin{aligned} E[y_{A1} - y_{B1}] &= (\mu + \alpha_A + \beta_1) - (\mu + \alpha_B + \beta_1) \\ &= \alpha_A - \alpha_B \end{aligned}$$

Some Poor Incomplete Block Designs (2) — A v.s. D

Block			
1	2	3	4
A	B	E	E
B	C	F	F
C	D	G	G

Based on the model

$$y_{ij} = \mu + \underset{\text{(treatment)}}{\alpha_i} + \underset{\text{(block)}}{\beta_j} + \underset{\text{(i.i.d. } N(0, \sigma^2))}{\varepsilon_{ij}}$$

can one find an unbiased estimate for $\alpha_A - \alpha_D$?

Some Poor Incomplete Block Designs (2) — A v.s. D

Block			
1	2	3	4
A	B	E	E
B	C	F	F
C	D	G	G

Based on the model

$$y_{ij} = \mu + \underset{\text{(treatment)}}{\alpha_i} + \underset{\text{(block)}}{\beta_j} + \underset{\text{(i.i.d. } N(0, \sigma^2))}{\varepsilon_{ij}}$$

can one find an unbiased estimate for $\alpha_A - \alpha_D$?

Yes,

- ▶ $y_{A1} - y_{B1}$ is an unbiased estimate for $\alpha_A - \alpha_B$
- ▶ $y_{B2} - y_{D2}$ is an unbiased estimate for $\alpha_B - \alpha_D$
- ▶ Their sum $y_{A1} - y_{B1} + y_{B2} - y_{D2}$ would be an unbiased estimate for

$$(\alpha_A - \alpha_B) + (\alpha_B - \alpha_D) = \alpha_A - \alpha_D$$

Some Poor Incomplete Block Designs (2) — A v.s. E

Block			
1	2	3	4
A	B	E	E
B	C	F	F
C	D	G	G

Based on the model

$$y_{ij} = \mu + \underset{\text{(treatment)}}{\alpha_i} + \underset{\text{(block)}}{\beta_j} + \underset{\text{(i.i.d. } N(0, \sigma^2))}{\varepsilon_{ij}}$$

can one find an unbiased estimate for $\alpha_A - \alpha_E$?

Some Poor Incomplete Block Designs (2) — A v.s. E

Block			
1	2	3	4
A	B	E	E
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Based on the model

$$y_{ij} = \mu + \underset{\text{(treatment)}}{\alpha_i} + \underset{\text{(block)}}{\beta_j} + \underset{\text{(i.i.d. } N(0, \sigma^2))}{\varepsilon_{ij}}$$

can one find an unbiased estimate for $\alpha_A - \alpha_E$?

Incomplete block designs must be “connected”
or not all pairwise comparisons can be estimated.

Balanced Incomplete Block Designs (BIBD)

BIBD is not balanced in the general sense that all treatment-block combinations occur equally often. Rather they are balanced in the looser sense by the criteria described below.

A *balanced incomplete block design* with

g treatments,

b blocks,

k as the size of each block,

r replications of each treatment,

is a design satisfying the following:

Incomplete: ▶ $k < g$.

Balanced: ▶ Each treatment appears at most once per block and has the same number of replicates r

▶ Each pair of treatments appear in a block the same number of times λ

Subject	1	2	3	4	5	6	7	8	9	10
	A	B	E	A	C	D	B	E	D	C
	B	D	A	C	A	E	C	B	E	D
	C	A	B	D	E	A	D	C	B	E

The design of the marketing psychology study is a BIBD with

$g =$ number of treatments $= 5$

$b =$ number of blocks $= 10$

$k =$ size of each block $= 3$

$r =$ number replicates per treatment $= 6$

The table below shows the blocks each treatment appears, verifying that each treatment appears $r = 6$ times.

Treatment	Block									
	1	2	3	4	5	6	7	8	9	10
A	✓	✓	✓	✓	✓	✓				
B	✓	✓	✓				✓	✓	✓	
C	✓			✓	✓		✓	✓		✓
D		✓		✓		✓	✓		✓	✓
E			✓		✓	✓		✓	✓	✓

Subject	1	2	3	4	5	6	7	8	9	10
	A	B	E	A	C	D	B	E	D	C
	B	D	A	C	A	E	C	B	E	D
	C	A	B	D	E	A	D	C	B	E

BIBD requires each pair of treatments appears in a block the same number (λ) of times. The table below verifies that, each treatment pair appears $\lambda = 3$ times for the design above.

Treatment-pair	Block									
	1	2	3	4	5	6	7	8	9	10
AB	✓	✓	✓							
AC	✓			✓	✓					
AD		✓		✓		✓				
AE			✓		✓	✓				
BC	✓						✓	✓		
BD		✓					✓		✓	
BE			✓					✓	✓	
CD				✓			✓			✓
CE					✓			✓		✓
DE						✓			✓	✓

First Balancing Condition of BIBD

The five numbers that describe a BIBD: g , b , k , r , and λ are not arbitrary.

There might not exist an allocation b blocks of k units to g treatments that is a BIBD.

- ▶ There are b blocks of size k each,
⇒ total number of experimental units is $N = bk$.
- ▶ There are g treatments, each appears r times in the design
⇒ total number of experimental units is $N = rg$.

Hence a BIBD must satisfy the **first balancing condition**:

$$N = bk = rg.$$

Second Balancing Condition of BIBD

In a BIBD, every pair of treatments must appear in a block the same number of times, say λ times.

Observe the total number of pairings involving treatment A equals

- ▶ $\lambda(g - 1)$, since A may be paired (appear in the same block) λ times with any of the other $g - 1$ treatments,
- ▶ $r(k - 1)$ since treatment A appears in r blocks. Within each of those blocks, there are $k - 1$ pairs including A as the block size is k

The **second balancing condition**

$$\boxed{r(k - 1) = \lambda(g - 1)}$$

Given g treatments and b blocks of size k , one can show that a BIBD that with r replicates per treatment and each pair of treatments show in a block λ times exists if and only if

$$bk = rg \text{ and } r(k - 1) = \lambda(g - 1).$$

Example (Eyedrop): $g = 3, k = 2$.

- ▶ Is it possible to find a BIBD w/ $b = 5$ subjects (blocks)?
- ▶ Is it possible to find a BIBD w/ $b = 6$ subjects (blocks)?

Example (Marketing Psychology): $g = 5, k = 3$.

- ▶ Is it possible to find a BIBD w/ $b = 5$ subjects (blocks)?
- ▶ Is it possible to find a BIBD w/ $b = 10$ subjects (blocks)?

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Example (Eyedrop): $g = 3, k = 2$.

- ▶ Is it possible to find a BIBD w/ $b = 5$ subjects (blocks)?
No. $r = bk/g = 2 \times 5/3 = 10/3$ is NOT an integer.
- ▶ Is it possible to find a BIBD w/ $b = 6$ subjects (blocks)?

Example (Marketing Psychology): $g = 5, k = 3$.

- ▶ Is it possible to find a BIBD w/ $b = 5$ subjects (blocks)?

- ▶ Is it possible to find a BIBD w/ $b = 10$ subjects (blocks)?

Given g treatments and b blocks of size k , one can show that a BIBD that with r replicates per treatment and each pair of treatments show in a block λ times exists if and only if

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No. $r = bk/g = 2 \times 5/3 = 10/3$ is NOT an integer.
- ▶ Is it possible to find a BIBD w/ $b = 6$ subjects (blocks)?
Yes, as $r = bk/g = 2 \times 6/3 = 4$ and $\lambda = \frac{r(k-1)}{(g-1)} = \frac{4(2-1)}{3-1} = 2$ are both integers

Example (Marketing Psychology): $g = 5, k = 3$.

- ▶ Is it possible to find a BIBD w/ $b = 5$ subjects (blocks)?

- ▶ Is it possible to find a BIBD w/ $b = 10$ subjects (blocks)?

Given g treatments and b blocks of size k , one can show that a BIBD that with r replicates per treatment and each pair of treatments show in a block λ times exists if and only if

$$bk = rg \text{ and } r(k - 1) = \lambda(g - 1).$$

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Example (Marketing Psychology): $g = 5, k = 3$.

- ▶ Is it possible to find a BIBD w/ $b = 5$ subjects (blocks)?
No. $r = bk/g = 3 \cdot 5/5 = 3$ is an integer, but $\lambda = \frac{r(k-1)}{(g-1)} = \frac{3(3-1)}{5-1} = 6/4$ is NOT an integer.
- ▶ Is it possible to find a BIBD w/ $b = 10$ subjects (blocks)?

Given g treatments and b blocks of size k , one can show that a BIBD that with r replicates per treatment and each pair of treatments show in a block λ times exists if and only if

$$bk = rg \text{ and } r(k - 1) = \lambda(g - 1).$$

Example (Eyedrop): $g = 3, k = 2$.

- ▶ Is it possible to find a BIBD w/ $b = 5$ subjects (blocks)?
No. $r = bk/g = 2 \times 5/3 = 10/3$ is NOT an integer.
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No. $r = bk/g = 3 \cdot 5/5 = 3$ is an integer, but $\lambda = \frac{r(k-1)}{(g-1)} = \frac{3(3-1)}{5-1} = 6/4$ is NOT an integer.
- ▶ Is it possible to find a BIBD w/ $b = 10$ subjects (blocks)?
Yes, as $r = bk/g = 10 \cdot 3/5 = 6$ and $\lambda = \frac{r(k-1)}{(g-1)} = \frac{6(3-1)}{5-1} = 3$ are both integers

Just like Latin Squares, it's not trivial to find a BIBD by oneself.

Appendix C.2 on p.609-615 of Oehlert's textbook gives a list of BIBD designs for $g \leq 9$.

► **A BIBD can be replicated to conduct a larger study.**

E.g., in the marketing psychology experiment, if we have $b = 20$ subjects (blocks) instead of 10, then we can do 2 repetitions of the BIBD below with $g = 5$, $k = 3$, $b = 10$, $r = 6$, $\lambda = 3$:

A	B	E	A	C	D	B	E	D	C
B	D	A	C	A	E	C	B	E	D
C	A	B	D	E	A	D	C	B	E

► **How to Do Randomization in BIBD?**

One obvious randomization is to randomize subjects to columns, then randomize the order of treatments in each block based on the above design.

Models for BIBD

$$y_{ij} = \mu + \underset{\text{(treatment)}}{\alpha_i} + \underset{\text{(block)}}{\beta_j} + \underset{\text{(i.i.d. } N(0, \sigma^2))}{\varepsilon_{ij}}$$

for $i = 1, \dots, g$, and $j = 1, \dots, b$ with

$$\sum_{i=1}^g \alpha_i = \sum_{j=1}^b \beta_j = 0.$$

- ▶ **additive** model (no treatment-block interaction)
- ▶ Not all y_{ij} exist because of incompleteness

Parameter Estimates for BIBD

Let

$$I_{ij} = \begin{cases} 1, & \text{if treatment } i \text{ appears in block } j, \\ 0, & \text{otherwise.} \end{cases}$$

and define

$$Q_i = y_{i\bullet} - \frac{1}{k} \sum_j I_{ij} y_{\bullet j}, \quad Q'_j = y_{\bullet j} - \frac{1}{r} \sum_i I_{ij} y_{i\bullet}$$

the least square estimates for μ , α_i , β_j are

$$\hat{\mu} = \frac{y_{\bullet\bullet}}{N}, \quad \hat{\alpha}_i = \frac{kQ_i}{\lambda g}, \quad \hat{\beta}_j = \frac{kQ'_j}{\lambda b}$$

Remark: Can verify that $\sum_i Q_i = 0 \Rightarrow \sum_i \hat{\alpha}_i = 0$.

You won't be asked to estimate parameters manually for a BIBD.

Example of BIBD— Problem 14.3 on. p.381 in Oehlert

The State Board of Education has adopted basic skills tests for high school graduation. One of these is a writing test. The student writing samples are graded by professional graders, and the board is taking some care to be sure that the graders are grading to the same standard. We examine grader differences with the following experiment. There are 25 graders available. We select 30 writing samples at random, each writing sample will be graded by 5 graders. Thus each grader will grade $30 \times 5/25 = 6$ samples.

Data: <http://users.stat.umn.edu/~gary/book/fcdae.data/pr14.3>

Questions of Interest:

- ▶ Did the 25 graders grade consistently with each other?
- ▶ How to adjust the scores if graders didn't grade consistently?
- ▶ If graders didn't grade consistently, can we identify the graders that were inconsistent with others?

Exam	Grader					Score					Exam	Grader					Score				
1	1	2	3	4	5	60	59	51	64	53	16	1	9	12	20	23	61	67	69	68	65
2	6	7	8	9	10	64	69	63	63	71	17	2	10	13	16	24	78	75	76	75	72
3	11	12	13	14	15	84	85	86	85	83	18	3	6	14	17	25	67	72	72	75	76
4	16	17	18	19	20	72	76	77	74	77	19	4	7	15	18	21	84	81	76	79	77
5	21	22	23	24	25	65	73	70	71	70	20	5	8	11	19	22	81	84	85	84	81
6	1	6	11	16	21	52	54	62	54	55	21	1	8	15	17	24	70	65	61	66	66
7	2	7	12	17	22	56	51	52	57	51	22	2	9	11	18	25	84	82	86	85	86
8	3	8	13	18	23	55	60	59	60	61	23	3	10	12	19	21	72	85	77	82	79
9	4	9	14	19	24	88	76	77	77	74	24	4	6	13	20	22	85	75	78	82	83
10	5	10	15	20	25	65	68	72	74	77	25	5	7	14	16	23	58	64	58	57	58
11	1	10	14	18	22	79	77	77	77	79	26	1	7	13	19	25	66	71	73	70	70
12	2	6	15	19	23	70	66	63	62	66	27	2	8	14	20	21	73	67	63	70	66
13	3	7	11	20	24	48	49	51	48	50	28	3	9	15	16	22	58	70	69	61	71
14	4	8	12	16	25	75	64	75	68	65	29	4	10	11	17	23	95	84	88	88	87
15	5	9	13	17	21	79	77	81	79	83	30	5	6	12	18	24	47	47	51	49	56

Here an exam is a writing sample.

- ▶ Which factor is the treatment factor? Graders or Exams?
- ▶ Which factor is the block factor? Graders or writing samples?
- ▶ Is this a BIBD?

Exam	Grader					Score					Exam	Grader					Score				
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3	11	12	13	14	15	84	85	86	85	83	18	3	6	14	17	25	67	72	72	75	76
4	16	17	18	19	20	72	76	77	74	77	19	4	7	15	18	21	84	81	76	79	77
5	21	22	23	24	25	65	73	70	71	70	20	5	8	11	19	22	81	84	85	84	81
6	1	6	11	16	21	52	54	62	54	55	21	1	8	15	17	24	70	65	61	66	66
7	2	7	12	17	22	56	51	52	57	51	22	2	9	11	18	25	84	82	86	85	86
8	3	8	13	18	23	55	60	59	60	61	23	3	10	12	19	21	72	85	77	82	79
9	4	9	14	19	24	88	76	77	77	74	24	4	6	13	20	22	85	75	78	82	83
10	5	10	15	20	25	65	68	72	74	77	25	5	7	14	16	23	58	64	58	57	58
11	1	10	14	18	22	79	77	77	77	79	26	1	7	13	19	25	66	71	73	70	70
12	2	6	15	19	23	70	66	63	62	66	27	2	8	14	20	21	73	67	63	70	66
13	3	7	11	20	24	48	49	51	48	50	28	3	9	15	16	22	58	70	69	61	71
14	4	8	12	16	25	75	64	75	68	65	29	4	10	11	17	23	95	84	88	88	87
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Graders.
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3	11	12	13	14	15	84	85	86	85	83	18	3	6	14	17	25	67	72	72	75	76
4	16	17	18	19	20	72	76	77	74	77	19	4	7	15	18	21	84	81	76	79	77
5	21	22	23	24	25	65	73	70	71	70	20	5	8	11	19	22	81	84	85	84	81
6	1	6	11	16	21	52	54	62	54	55	21	1	8	15	17	24	70	65	61	66	66
7	2	7	12	17	22	56	51	52	57	51	22	2	9	11	18	25	84	82	86	85	86
8	3	8	13	18	23	55	60	59	60	61	23	3	10	12	19	21	72	85	77	82	79
9	4	9	14	19	24	88	76	77	77	74	24	4	6	13	20	22	85	75	78	82	83
10	5	10	15	20	25	65	68	72	74	77	25	5	7	14	16	23	58	64	58	57	58
11	1	10	14	18	22	79	77	77	77	79	26	1	7	13	19	25	66	71	73	70	70
12	2	6	15	19	23	70	66	63	62	66	27	2	8	14	20	21	73	67	63	70	66
13	3	7	11	20	24	48	49	51	48	50	28	3	9	15	16	22	58	70	69	61	71
14	4	8	12	16	25	75	64	75	68	65	29	4	10	11	17	23	95	84	88	88	87
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1	1 2 3 4 5	60 59 51 64 53	16	1 9 12 20 23	61 67 69 68 65
2	6 7 8 9 10	64 69 63 63 71	17	2 10 13 16 24	78 75 76 75 72
3	11 12 13 14 15	84 85 86 85 83	18	3 6 14 17 25	67 72 72 75 76
4	16 17 18 19 20	72 76 77 74 77	19	4 7 15 18 21	84 81 76 79 77
5	21 22 23 24 25	65 73 70 71 70	20	5 8 11 19 22	81 84 85 84 81
6	1 6 11 16 21	52 54 62 54 55	21	1 8 15 17 24	70 65 61 66 66
7	2 7 12 17 22	56 51 52 57 51	22	2 9 11 18 25	84 82 86 85 86
8	3 8 13 18 23	55 60 59 60 61	23	3 10 12 19 21	72 85 77 82 79
9	4 9 14 19 24	88 76 77 77 74	24	4 6 13 20 22	85 75 78 82 83
10	5 10 15 20 25	65 68 72 74 77	25	5 7 14 16 23	58 64 58 57 58
11	1 10 14 18 22	79 77 77 77 79	26	1 7 13 19 25	66 71 73 70 70
12	2 6 15 19 23	70 66 63 62 66	27	2 8 14 20 21	73 67 63 70 66
13	3 7 11 20 24	48 49 51 48 50	28	3 9 15 16 22	58 70 69 61 71
14	4 8 12 16 25	75 64 75 68 65	29	4 10 11 17 23	95 84 88 88 87
15	5 9 13 17 21	79 77 81 79 83	30	5 6 12 18 24	47 47 51 49 56

Here a exam is a writing sample.

- ▶ Which factor is the treatment factor? Graders or Exams?
Graders.
- ▶ Which factor is the block factor? Graders or writing samples?
Exams.
- ▶ Is this a BIBD?

Yes, $g = 25$, $b = 30$, $k = 5$, $r = \frac{bk}{g} = 6$, $\lambda = \frac{r(k-1)}{g-1} = 1$.

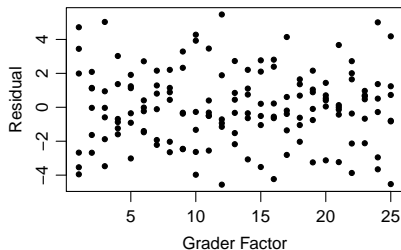
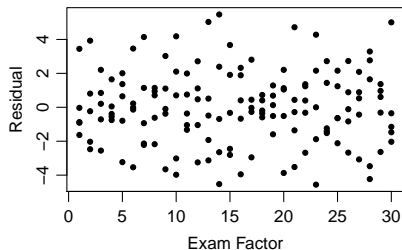
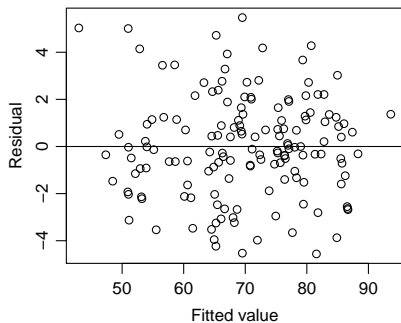
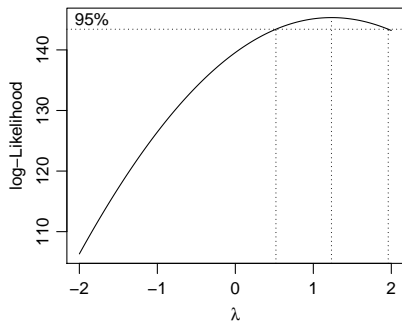
$$\begin{array}{ccccccc} y_{ij} & = & \mu & + & \alpha_i & + & \beta_j & + & \varepsilon_{ij} \\ \text{(score)} & & & & \text{(grader)} & & \text{(exam)} & & \end{array}$$

As writing samples differ in levels, we expect β_j not all equal.

If graders were consistent, they should give the same score to the same writing sample, i.e., $\alpha_1 = \alpha_2 = \dots = \alpha_{25}$

```
pr14.3 = read.table(  
  "http://users.stat.umn.edu/~gary/book/fcdae.data/pr14.3", h=T)  
pr14.3$EXAM = as.factor(pr14.3$exam)  
pr14.3$GRADER = as.factor(pr14.3$grader)  
lm1 = lm(score ~ EXAM + GRADER, data=pr14.3)
```

Always Check Model Assumptions First



How to Adjust Scores as Graders Were Inconsistent?

Based on the model $y_{ij} = \mu + \alpha_i + \beta_j + \varepsilon_{ij}$, the score of the i th writing sample is $\mu + \beta_j$, which is estimated by $\hat{\mu} + \hat{\beta}_j$.

How to get $\hat{\beta}_j$ in R? Recall R by default estimates parameters using the baseline constraints $\alpha_1 = \beta_1 = 0$, not the zero-sum constraints $\sum_{i=1}^g \alpha_i = \sum_{j=1}^b \beta_j = 0$.

One can use `contrasts()` and `contr.sum()` to force R using the zero-sum constraints.

```
contrasts(pr14.3$EXAM) = contr.sum(30)
contrasts(pr14.3$GRADER) = contr.sum(25)
lm1 = lm(score ~ EXAM + GRADER, data=pr14.3)
```

```
lm1$coef
(Intercept)          EXAM1          EXAM2      .... (omitted)
    69.960         -12.568         -3.368
EXAM28          EXAM29          GRADER1      .... (omitted)  GRADER24
   -2.128          16.192         -0.840                          0.160
```

Why is there no estimate for exam #30, nor for grader #25?

```

muhat = lm1$coef[1]
betahat = vector("numeric",length=30)
betahat[1:29] = lm1$coef[2:30]
betahat[30] = -sum(betahat[1:29])
adjustedscore = muhat + betahat; adjustedscore
 [1] 57.39 66.59 84.39 75.15 69.47 56.38 51.62 60.42 77.50 71.50
[11] 77.85 65.65 49.33 68.21 80.57 65.79 74.79 73.95 78.11 83.35
[21] 66.12 83.44 80.24 78.76 60.24 69.51 67.67 67.83 86.15 50.83
names(adjustedscore) = 1:30
adjustedscore
   1    2    3    4    5    6    7    8    9   10
57.39 66.59 84.39 75.15 69.47 56.38 51.62 60.42 77.50 71.50
  11   12   13   14   15   16   17   18   19   20
77.85 65.65 49.33 68.21 80.57 65.79 74.79 73.95 78.11 83.35
  21   22   23   24   25   26   27   28   29   30
66.12 83.44 80.24 78.76 60.24 69.51 67.67 67.83 86.15 50.83

```

Compare adjusted scores with unadjusted scores (average of the 5 raw scores per exam).

```
library(mosaic)
unadjustedscore = mean(score ~ EXAM, data=pr14.3)
unadjustedscore
  1    2    3    4    5    6    7    8    9   10   11   12   13
57.4 66.0 84.6 75.2 69.8 55.4 53.4 59.0 78.4 71.2 77.8 65.4 49.2
 14   15   16   17   18   19   20   21   22   23   24   25   26
69.4 79.8 66.0 75.2 72.4 79.4 83.0 65.6 84.6 79.0 80.6 59.0 70.0
 27   28   29   30
67.8 65.8 88.4 50.0
```

Difference of unadjusted and adjusted scores:

```
sort(unadjustedscore - adjustedscore)
 28    18     8    23    25     6    30    15     2
-2.032 -1.552 -1.416 -1.240 -1.240 -0.976 -0.832 -0.768 -0.592
 21    20    10    12    13    11     1     4    27
-0.520 -0.352 -0.296 -0.248 -0.128 -0.048  0.008  0.048  0.128
  3    16     5    17    26     9    22    14    19
 0.208  0.208  0.328  0.408  0.488  0.904  1.160  1.192  1.288
  7    24    29
 1.784  1.840  2.248
```

ANOVA Table for BIBD

```
anova(lm(score ~ GRADER + EXAM, data=pr14.3))
```

Analysis of Variance Table

Response: score

	Df	Sum Sq	Mean Sq	F value	Pr(>F)
GRADER	24	4073.1	169.71	23.659	< 2.2e-16
EXAM	29	13342.0	460.07	64.138	< 2.2e-16
Residuals	96	688.6	7.17		

```
anova(lm(score ~ EXAM + GRADER, data=pr14.3))
```

Analysis of Variance Table

Response: score

	Df	Sum Sq	Mean Sq	F value	Pr(>F)
EXAM	29	16609.0	572.72	79.8424	< 2.2e-16
GRADER	24	806.2	33.59	4.6828	0.00000002694
Residuals	96	688.6	7.17		

The two ANOVA tables have identical SSE
but *different SS* for EXAM and GRADER. Why?

- ▶ As a BIBD doesn't include all treatment-block combination, it does NOT have a *balanced* factorial structure of treatment \times block.
- ▶ For unbalanced factorial data, there are 3 types of sum of squares
 - ▶ the `anova()` command gives the *Type I sum of squares*
 - ▶ What's a Type I Sum of Square?

Digression: Sum of Squares for Unbalanced Factorial Data

What Happens If Factorial Data Become Unbalanced

- ▶ no simple formulae for parameter estimates and SS.
- ▶ the parameter estimates and SS of a term will depend on the presence of other terms in the model, e.g., the estimates for α_i 's might be different in the following 3 models

$$y_{ijk} = \mu + \alpha_i + \beta_j + \alpha\beta_{ij} + \varepsilon_{ijk}$$

$$y_{ijk} = \mu + \alpha_i + \beta_j + \varepsilon_{ijk}$$

$$y_{ijk} = \mu + \alpha_i + \varepsilon_{ijk}$$

- ▶ need to rely on statistical software for computation
- ▶ there are 3 variations of SS

Notation for Models

In the following, we denote various models by listing the included effect. For example,

- ▶ $(1, A, B, AB)$ denotes the model $y_{ijk} = \mu + \alpha_i + \beta_j + \alpha\beta_{ij} + \varepsilon_{ijk}$
- ▶ $(1, A, B)$ denotes the model $y_{ijk} = \mu + \alpha_i + \beta_j + \varepsilon_{ijk}$
- ▶ $(1, A, B, C, AB, AC)$ denotes the model

$$y_{ijkl} = \mu + \alpha_i + \beta_j + \gamma_k + \alpha\beta_{ij} + \alpha\gamma_{ik} + \varepsilon_{ijkl}$$

Here the “**1**” stands for the grand mean μ .

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$$y_{ijkl} = \mu + \alpha_i + \beta_j + \gamma_k + \alpha\beta_{ij} + \alpha\gamma_{ik} + \varepsilon_{ijkl}$$

Here the “1” stands for the grand mean μ .

In the following $SSE(\text{model})$ denotes the SSE of that model, e.g., $SSE(1, A, B, AB)$ means the SSE of the model

$$y_{ijk} = \mu + \alpha_i + \beta_j + \alpha\beta_{ij} + \varepsilon_{ijk}.$$

For unbalanced data, there is no simple formula to compute the SSE. One must write the model as a regression model and use statistical software to compute the SSE.

Adjusted Sum of Squares (1)

The **adjusted sum of squares** for main effects B adjusted for A is defined as

$$SS(B|1, A) = SSE(1, A) - SSE(1, A, B).$$

- ▶ $SS(B|1, A) \geq 0$ since the model $(1, A)$ is *included* (*nested*) in the model $(1, A, B)$ and hence the latter always has a smaller SSE
- ▶ $SS(B|1, A)$ is the reduction in SSE after B is included in the model
- ▶ $SS(B|1, A)$ describes the effect of B adjusted for A since with consider two models that A is present in both and the two models only differ by B

Adjusted Sum of Squares (2)

Likewise, the **adjusted sum of squares** for main effects B adjusted for A, C, and AC is

$$SS(B|1, A, C, AC) = SSE(1, A, C, AC) - SSE(1, A, B, C, AC).$$

Adjusted Sum of Squares (2)

Likewise, the **adjusted sum of squares** for main effects B adjusted for A, C, and AC is

$$SS(B|1, A, C, AC) = SSE(1, A, C, AC) - SSE(1, A, B, C, AC).$$

In general, the **adjusted sum of squares** for a term adjusted for some other terms is

$$\begin{aligned} &SS(\text{a term}|\text{some other terms}) \\ &= SSE(\text{some other terms}) - SSE(\text{a term}, \text{some other terms}) \end{aligned}$$

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$$\begin{aligned} &SS(\text{a term} | \text{some other terms}) \\ &= SSE(\text{some other terms}) - SSE(\text{a term}, \text{some other terms}) \end{aligned}$$

For balanced data, adjusted SS = unadjusted SS

$$SS(A|1, B) = SS(A|1, B, C) = SS(A|1, B, C, BC) = SS(A|1).$$

Type I Sum of Squares

For a specified model, the *Type I Sum of Squares* (aka. *Sequential Sum of Squares*) for any term is adjusted for those terms that precede it in the model.

- ▶ E.g, the Type I SS's for the model (1, A, B, AB, C) are

Source	d.f.	Type I SS
A	$a - 1$	$SS(A 1)$
B	$b - 1$	$SS(B 1, A)$
AB	$(a - 1)(b - 1)$	$SS(AB 1, A, B)$
C	$c - 1$	$SS(C 1, A, B, AB)$

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C	$c - 1$	$SS(C 1, A, B, AB)$

Type I SS's depend on how the terms are **ordered** in a model:

- ▶ E.g, if the terms in the model (1, A, B, AB, C) is reshuffled as (1, C, A, B, AB), then the Type I SS's become

Source	d.f.	Type I SS
C	$c - 1$	$SS(C 1)$
A	$a - 1$	$SS(A 1, C)$
B	$b - 1$	$SS(B 1, A, C)$
AB	$(a - 1)(b - 1)$	$SS(AB 1, A, B, C)$

Type I Sum of Squares and SSE Add Up to SST (1)

Source	Type I SS
<i>A</i>	$SS(A 1) = SSE(1) - SSE(1, A)$
<i>B</i>	$SS(B 1, A) = SSE(1, A) - SSE(1, A, B)$
<i>AB</i>	$SS(AB 1, A, B) = SSE(1, A, B) - SSE(1, A, B, AB)$
<i>C</i>	$SS(C 1, A, B, AB) = SSE(1, A, B, AB) - SSE(1, A, B, AB, C)$
Error	$SSE(1, A, B, AB, C)$
Sum	

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C	$SS(C 1, A, B, AB) = SSE(1, A, B, AB) - \cancel{SSE(1, A, B, AB, C)}$
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Sum	

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Sum	

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Sum	

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C	$SS(C 1, A, B, AB) = \cancel{\text{SSE}(1, A, B, AB)} - \cancel{\text{SSE}(1, A, B, AB, C)}$
Error	$\cancel{\text{SSE}(1, A, B, AB, C)}$
Sum	$\text{SSE}(1) = \text{SST}$

SSE(1) is the SSE for the model $y_{ijkl} = \mu + \varepsilon_{ijkl}$, of which the optimal (least square) estimate for μ is the overall mean $\bar{y}_{\dots\dots}$. Hence,

$$\text{SSE}(1) = \sum_{ijkl} (y_{ijkl} - \bar{y}_{\dots\dots})^2 = \text{SST}.$$

Type I Sum of Squares and SSE Add Up to SST (2)

If the order of terms in the model $(1, A, B, AB, C)$ is changed to $(1, C, A, B, AB)$,

- ▶ the Type I SS's are changed;
- ▶ $SSE(1, A, B, AB, C) = SSE(1, C, A, B, AB)$ is not affected by the order of terms;
- ▶ the Type I SS's and the SSE always add up to SST.

Source	Type I SS
C	$SS(C 1) = SSE(1) - SSE(1, C)$
A	$SS(A 1, C) = SSE(1, C) - SSE(1, C, A)$
B	$SS(B 1, C, A) = SSE(1, C, A) - SSE(1, C, A, B)$
AB	$SS(AB 1, A, B, C) = SSE(1, C, A, B) - SSE(1, A, B, C, AB)$
Error	$SSE(1, C, A, B, AB)$
Sum	

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Sum	

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Sum	

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- ▶ the Type I SS's are changed;
- ▶ $SSE(1, A, B, AB, C) = SSE(1, C, A, B, AB)$ is not affected by the order of terms;
- ▶ the Type I SS's and the SSE always add up to SST.

Source	Type I SS
C	$SS(C 1) = SSE(1) - \cancel{SSE(1, C)}$
A	$SS(A 1, C) = \cancel{SSE(1, C)} - \cancel{SSE(1, C, A)}$
B	$SS(B 1, C, A) = \cancel{SSE(1, C, A)} - \cancel{SSE(1, C, A, B)}$
AB	$SS(AB 1, A, B, C) = \cancel{SSE(1, C, A, B)} - \cancel{SSE(1, A, B, C, AB)}$
Error	$\cancel{SSE(1, C, A, B, AB)}$
Sum	

Type I Sum of Squares and SSE Add Up to SST (2)

If the order of terms in the model (1, A, B, AB, C) is changed to (1, C, A, B, AB),

- ▶ the Type I SS's are changed;
- ▶ $SSE(1, A, B, AB, C) = SSE(1, C, A, B, AB)$ is not affected by the order of terms;
- ▶ the Type I SS's and the SSE always add up to SST.

Source	Type I SS
C	$SS(C 1) = \text{SSE}(1) - \cancel{SSE(1, C)}$
A	$SS(A 1, C) = \cancel{SSE(1, C)} - \cancel{SSE(1, C, A)}$
B	$SS(B 1, C, A) = \cancel{SSE(1, C, A)} - \cancel{SSE(1, C, A, B)}$
AB	$SS(AB 1, A, B, C) = \cancel{SSE(1, C, A, B)} - \cancel{SSE(1, A, B, C, AB)}$
Error	$\cancel{SSE(1, C, A, B, AB)}$
Sum	$\text{SSE}(1) = \text{SST}$

Example: Popcorn Microwave Data Revisit

$3 \times 2 \times 3$ factorial design with 2 replicates.

Brand (<i>i</i>)	Power (<i>j</i>)	Time (<i>k</i>)		
		1 (4 min)	2 (4.5 min)	3 (5 min)
1	1 (500 W)	73.8, 65.5	70.3, 91.0	72.7, 81.9
1	2 (625 W)	70.8, 75.3	78.7, 88.7	74.1, 72.1
2	1 (500 W)	73.7, 65.8	93.4, 76.3	45.3, 47.6
2	2 (625 W)	79.3, 86.5	92.2, 84.7	66.3, 45.7
3	1 (500 W)	62.5, 65.0	50.1, 81.5	51.4, 67.7
3	2 (625 W)	82.1, 74.5	71.5, 80.0	64.0, 77.0

```
popcorn = read.table(  
  "http://www.stat.uchicago.edu/~yibi/s222/popcorn.txt", h=T)  
popcorn$brand = as.factor(popcorn$brand)  
popcorn$power = as.factor(popcorn$power)  
popcorn$time = as.factor(popcorn$time)
```

For balanced data, SS's are not affected by the order of the terms in the model

```
anova(lm(y ~ brand*time+power, data=popcorn))
```

Analysis of Variance Table

Response: y

	Df	Sum Sq	Mean Sq	F value	Pr(>F)
brand	2	331.10	165.55	2.3031	0.1199906
time	2	1554.58	777.29	10.8133	0.0003825
power	1	455.11	455.11	6.3313	0.0183703
brand:time	4	1433.86	358.46	4.9868	0.0040523
Residuals	26	1868.95	71.88		

```
anova(lm(y ~ power+brand*time, data=popcorn))
```

Analysis of Variance Table

Response: y

	Df	Sum Sq	Mean Sq	F value	Pr(>F)
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brand:time	4	1433.86	358.46	4.9868	0.0040523
Residuals	26	1868.95	71.88		

If the first observation (73.8) is removed `popcorn[-1,]` is removed, the data become unbalanced.

```
anova(lm(y ~ brand*time+power, data=popcorn[-1,]))
```

Analysis of Variance Table

Response: y

	Df	Sum Sq	Mean Sq	F value	Pr(>F)
brand	2	334.95	167.48	2.3017	0.1209104
time	2	1559.57	779.79	10.7172	0.0004353
power	1	443.81	443.81	6.0996	0.0206998
brand:time	4	1483.60	370.90	5.0975	0.0038263
Residuals	25	1819.02	72.76		

```
anova(lm(y ~ power+brand*time, data=popcorn[-1,]))
```

Analysis of Variance Table

Response: y

	Df	Sum Sq	Mean Sq	F value	Pr(>F)
power	1	480.66	480.66	6.6061	0.0165065
brand	2	304.29	152.15	2.0911	0.1446414
time	2	1553.38	776.69	10.6746	0.0004454
brand:time	4	1483.60	370.90	5.0975	0.0038263
Residuals	25	1819.02	72.76		

Type I ANOVA table

The Type I ANOVA table for unbalanced data are identical to the ANOVA table for balanced data in every aspect except the SS's are replaced by the Type I SS.

Source	d.f.	Type I SS	MS	F-value
A	$a - 1$	$SS(A 1)$	SS_A/df_A	MS_A/MSE
B	$b - 1$	$SS(B 1, A)$	SS_B/df_B	MS_B/MSE
C	$c - 1$	$SS(C 1, A, B)$	SS_C/df_C	MS_C/MSE
AB	$(a - 1)(b - 1)$	$SS(AB 1, A, B, C)$	SS_{AB}/df_{AB}	MS_{AB}/MSE
AC	$(a - 1)(c - 1)$	$SS(AC 1, A, B, C, AB)$	SS_{AC}/df_{AC}	MS_{AC}/MSE
BC	$(a - 1)(c - 1)$	$SS(BC 1, A, B, C, AB, AC)$	SS_{BC}/df_{BC}	MS_{BC}/MSE
ABC	$(a - 1)(b - 1)(c - 1)$	$SS(ABC 1, A, B, C, AB, AC, BC)$	SS_{ABC}/df_{ABC}	MS_{ABC}/MSE
Error	$N - abc$	SSE	SSE/df_E	
Total	$N - 1$	SST		

Type I SS's and the SSE always add up to SST.

Why Type I SS's Are Not Ideal?

Look the 3 F -statistic for the 3 main effects in the previous page.

- ▶ The F -statistic for A is unadjusted
- ▶ The F -statistic for B is adjusted with A
- ▶ The F -statistic for C is adjusted with both A and B

When considering whether a term, say A, is needed in a model, one should look at the *net effect* of A after adjusting for the effect of other terms.

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1, B, C, BC.

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- ▶ Thus, a more sensible adjusted SS for A is $SS(A|1, B, C, BC)$.

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Look the 3 F -statistic for the 3 main effects in the previous page.

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When considering whether a term, say A, is needed in a model, one should look at the *net effect* of A after adjusting for the effect of other terms.

- ▶ What are the terms that should be accounted for before considering A?

1, B, C, BC.

- ▶ Why not adjusting for AB, AC and ABC?
- ▶ Thus, a more sensible adjusted SS for A is $SS(A|1, B, C, BC)$.
- ▶ Such adjusted SS's are called the *Type II Sum of Squares*.

Type II Sum of Squares

The Type II SS_U of an effect U (U can be a main effect or an interaction) is computed as follows:

- ▶ take the biggest hierarchical model without effect U , and then compare it to the model with U added.

Here “largest hierarchical model” means all the effects that don't include term U . E.g., for the model (1, A, B, C, AB, AC, BC, ABC),

- ▶ the Type II SS for AB is $SS(AB|1, A, B, C, AC, BC)$
- ▶ the Type II SS for C is $SS(C|1, A, B, AB)$ but not $SS(C|1, A)$ or $SS(C|1, A, AB)$

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- ▶ the Type II SS for C is $SS(C|1, A, B, AB)$ but not $SS(C|1, A)$ or $SS(C|1, A, AB)$

Unlike Type I SS, Type II SS does NOT depend on the *order* of terms in a model.

Type II ANOVA table for 3-Way Data

Source	d.f.	Type II SS	MS	F-value
A	$a - 1$	$SS(A 1, B, C, BC)$	SS_A/df_A	MS_A/MSE
B	$b - 1$	$SS(B 1, A, C, AC)$	SS_B/df_B	MS_B/MSE
C	$c - 1$	$SS(C 1, A, B, AB)$	SS_C/df_C	MS_C/MSE
AB	$(a - 1)(b - 1)$	$SS(AB 1, A, B, C, AC, BC)$	SS_{AB}/df_{AB}	MS_{AB}/MSE
AC	$(a - 1)(c - 1)$	$SS(AC 1, A, B, C, AB, BC)$	SS_{AC}/df_{AC}	MS_{AC}/MSE
BC	$(a - 1)(c - 1)$	$SS(BC 1, A, B, C, AB, AC)$	SS_{BC}/df_{BC}	MS_{BC}/MSE
ABC	$(a - 1)(b - 1)(c - 1)$	$SS(ABC 1, A, B, C, AB, AC, BC)$	SS_{ABC}/df_{ABC}	MS_{ABC}/MSE
Error	$N - abc$	SSE	SSE/df_E	

Type II SS of terms in a model will NOT sum to SST

Computing Type II ANOVA Table in R

The build-in function `anova()` in R gives Type I SS's only. To get the Type II SS's, first load the library `car` (which is the short for "Companion to Applied Regression"), and then use the function `Anova()` as follows.

```
library(car)
Anova(yourmodel, type=2)
```

Note the first letter `A` in `Anova()` is a **capital letter A**.

Type II ANOVA table:

```
library(car)
lm2 = lm(y ~ brand*power*time, data=popcorn[-1,])
Anova(lm2, type=2)
Anova Table (Type II tests)
```

Response: y

	Sum Sq	Df	F value	Pr(>F)
brand	292.02	2	1.6082	0.229256
power	497.05	1	5.4747	0.031753
time	1559.33	2	8.5876	0.002644
brand:power	141.08	2	0.7770	0.475450
brand:time	1464.49	4	4.0326	0.017689
power:time	68.18	2	0.3755	0.692505
brand:power:time	49.33	4	0.1358	0.966830
Residuals	1543.43	17		

```
lm2 = lm(y ~ brand*power*time, data=popcorn[-1,])
```

```
anova(lm2)
```

```
Analysis of Variance Table
```

```
Response: y
```

	Df	Sum Sq	Mean Sq	F value	Pr(>F)
brand	2	334.95	167.48	1.8447	0.188353
power	1	450.00	450.00	4.9565	0.039804
time	2	1553.38	776.69	8.5548	0.002688
brand:power	2	205.72	102.86	1.1330	0.345227
brand:time	4	1435.95	358.99	3.9540	0.019026
power:time	2	68.18	34.09	0.3755	0.692505
brand:power:time	4	49.33	12.33	0.1358	0.966830
Residuals	17	1543.43	90.79		

Note the Type I ANOVA table given by build-in `anova()` command is different from the Type II table given by the `Anova()` in the `car` library.

Back to BIBD

ANOVA for BIBD (Type I Sum of Squares!)

Source	d.f.	SS	MS	F-value
Block	$b - 1$	SS_{block}	MS_{block}	(MS_{block}/MSE)
Treatment	$g - 1$	SS_{trt}	MS_{trt}	MS_{trt}/MSE
Error	$N - g - b + 1$	SSE	MSE	
Total	$N - 1$	SS_{total}		

Let $I_{ij} = \begin{cases} 1, & \text{if treatment } i \text{ appears in block } j, \\ 0, & \text{otherwise.} \end{cases}$

$$\text{Then } SS_{total} = \sum_{i=1}^g \sum_{j=1}^b I_{ij} (y_{ij} - \bar{y}_{\bullet\bullet})^2$$

$$SS_{block} = k \sum_{j=1}^b (\bar{y}_{\bullet j} - \bar{y}_{\bullet\bullet})^2 \quad (\text{unadjusted, Type I})$$

$$SS_{trt} = \frac{k}{\lambda g} \sum_i Q_i^2 = \frac{\lambda g}{k} \sum_i \hat{\alpha}_i^2 \quad (\text{adjusted for block, Type I \& II})$$

$$SSE = SS_{total} - SS_{block} - SS_{trt}$$

ANOVA for BIBD (Type I Sum of Squares!)

Source	d.f.	SS	MS	F-value
Block	$b - 1$	SS_{block}	MS_{block}	(MS_{block}/MSE)
Treatment	$g - 1$	SS_{trt}	MS_{trt}	MS_{trt}/MSE
Error	$N - g - b + 1$	SSE	MSE	
Total	$N - 1$	SS_{total}		

Let $I_{ij} = \begin{cases} 1, & \text{if treatment } i \text{ appears in block } j, \\ 0, & \text{otherwise.} \end{cases}$

$$\text{Then } SS_{total} = \sum_{i=1}^g \sum_{j=1}^b I_{ij} (y_{ij} - \bar{y}_{\bullet\bullet})^2$$

$$SS_{block} = k \sum_{j=1}^b (\bar{y}_{\bullet j} - \bar{y}_{\bullet\bullet})^2 \quad (\text{unadjusted, Type I})$$

$$SS_{trt} = \frac{k}{\lambda g} \sum_i Q_i^2 = \frac{\lambda g}{k} \sum_i \hat{\alpha}_i^2 \quad (\text{adjusted for block, Type I \& II})$$

$$SSE = SS_{total} - SS_{block} - SS_{trt}$$

For incomplete block designs, **always place *Block ahead of Treatment* in the ANOVA table.** The SS_{trt} will then be adjusted for Block and hence is Type II.

```
anova(lm(score ~ GRADER + EXAM, data=pr14.3))
```

Analysis of Variance Table

Response: score

	Df	Sum Sq	Mean Sq	F value	Pr(>F)
GRADER	24	4073.09	169.712	23.6593	< 2.22e-16
EXAM	29	13342.04	460.070	64.1377	< 2.22e-16
Residuals	96	688.62	7.173		

```
anova(lm(score ~ EXAM + GRADER, data=pr14.3))
```

Analysis of Variance Table

Response: score

	Df	Sum Sq	Mean Sq	F value	Pr(>F)
EXAM	29	16608.96	572.723	79.84239	< 2.22e-16
GRADER	24	806.18	33.591	4.68282	0.00000002694
Residuals	96	688.62	7.173		

Which ANOVA table should we look at to determine the significance of treatment (GRADER)?

Pairwise Comparisons

Estimate of $\alpha_{i_1} - \alpha_{i_2}$ is

$$\hat{\alpha}_{i_1} - \hat{\alpha}_{i_2} = \frac{k}{\lambda g} (Q_{i_1} - Q_{i_2})$$

- ▶ $SE(\hat{\alpha}_{i_1} - \hat{\alpha}_{i_2}) = \sqrt{MSE \left(\frac{2k}{\lambda g} \right)}$
- ▶ t -statistic = $\frac{\hat{\alpha}_{i_1} - \hat{\alpha}_{i_2}}{SE}$ with $df = df$ of MSE
- ▶ Tukey's HSD controlling FWER at α is

$$HSD = \frac{q_{\alpha}(g, \text{df of MSE})}{\sqrt{2}} \times SE.$$

How to Identify Inconsistent Graders?

We can do pairwise comparisons for the grader effects $\alpha_{i_1} - \alpha_{i_2}$ using the t -statistic $= \frac{\hat{\alpha}_{i_1} - \hat{\alpha}_{i_2}}{SE}$ where

$$SE = \sqrt{\text{MSE} \left(\frac{2k}{\lambda g} \right)} = \sqrt{7.173 \left(\frac{2 \times 5}{1 \times 25} \right)} \approx 1.6939$$

with $df = (\text{df of MSE}) = 96$.

Tukey's critical value for FWER = 0.05 is

```
qtukey(0.95, 25, df = 96)/sqrt(2)
[1] 3.768
```

Tukey's HSD $= \frac{q_{0.05}(25, 96)}{\sqrt{2}} SE = 3.7676 \times 1.6939 \approx 6.382$.

We have obtained $\hat{\alpha}_1, \hat{\alpha}_2, \dots, \hat{\alpha}_{24}$ in R on page 21.

```
lm1$coef[31:54]
```

GRADER1	GRADER2	GRADER3	GRADER4	GRADER5	GRADER6	GRADER7
-0.84	3.24	-6.36	7.48	-3.48	-2.36	1.60
GRADER8	GRADER9	GRADER10	GRADER11	GRADER12	GRADER13	GRADER14
-1.56	-1.12	0.48	2.16	1.32	0.76	-1.60
GRADER15	GRADER16	GRADER17	GRADER18	GRADER19	GRADER20	GRADER21
-1.60	-2.60	1.24	0.20	-0.40	1.80	-1.24
GRADER22	GRADER23	GRADER24				
1.52	-0.12	0.16				

The last one can be computed as $\hat{\alpha}_{25} = -\sum_{i=1}^{24} \hat{\alpha}_i = 1.32$ as $\sum_{i=1}^{25} \hat{\alpha}_i = 0$.

```
alphahat25 = -sum(lm1$coef[31:54]); alphahat25  
[1] 1.32  
names(alphahat25) = "GRADER25"  
alphahat = c(lm1$coef[31:54], alphahat25)
```

sort(alphahat)

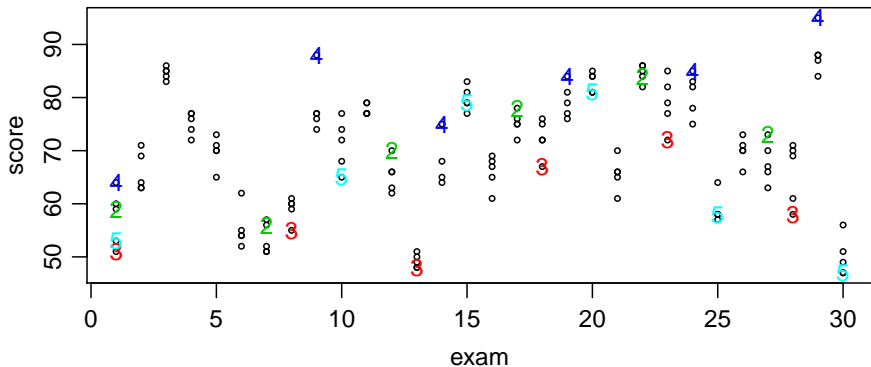
GRADER3	GRADER5	GRADER16	GRADER6	GRADER15	GRADER14	GRADER8
-6.36	-3.48	-2.60	-2.36	-1.60	-1.60	-1.56
GRADER21	GRADER9	GRADER1	GRADER19	GRADER23	GRADER24	GRADER18
-1.24	-1.12	-0.84	-0.40	-0.12	0.16	0.20
GRADER10	GRADER13	GRADER17	GRADER25	GRADER12	GRADER22	GRADER7
0.48	0.76	1.24	1.32	1.32	1.52	1.60
GRADER20	GRADER11	GRADER2	GRADER4			
1.80	2.16	3.24	7.48			

Underline Diagram for pairwise comparison between graders:
(at FWER = 5%, Tukey's HSD = 6.38)

3 5 16 6 15 14 8 21 9 1 19 23 24 18 10 13 17 25 12 22 7 20 11 2 4

After Tukey's adjustment, only Grader #3 and #4 are significantly inconsistent with most other graders.

Grader #2 and #5 were consistent with all the rest except #3 and #4.



- ▶ Grader #3 always gave the lowest score among the 5 graders grading the same exam
- ▶ Grader #4 always gave scores that substantially higher than the scores given by the other graders for the same exam.
- ▶ Grader #2 tends to give higher scores, Grader #5 tended to give lower scores, but not as much as Grader #3 and #4.

Tukey's HSD in emmeans

The `emmeans` library also works for incomplete block designs.

```
library(emmeans)
lm1 = lm(score ~ EXAM + GRADER, data=pr14.3)
lm1em = emmeans(lm1, "GRADER")
summary(pairs(lm1em, infer=c(T,T), level=0.95, adjust="tukey"))
```

Output on the next page.

Observe the CI's all equals to their respective estimate \pm HSD.
e.g., the CI for Grade #1- Grade #2 is

$$-4.08 \pm 6.382 = (2.302, -10.462).$$

contrast	estimate	SE	df	lower.CL	upper.CL	t.ratio	p.value
1 - 2	-4.08	1.69	96	-10.462	2.302	-2.409	0.7545
1 - 3	5.52	1.69	96	-0.862	11.902	3.259	0.1919
1 - 4	-8.32	1.69	96	-14.702	-1.938	-4.912	0.0009
1 - 5	2.64	1.69	96	-3.742	9.022	1.559	0.9973
1 - 6	1.52	1.69	96	-4.862	7.902	0.897	1.0000
1 - 7	-2.44	1.69	96	-8.822	3.942	-1.440	0.9991
1 - 8	0.72	1.69	96	-5.662	7.102	0.425	1.0000
1 - 9	0.28	1.69	96	-6.102	6.662	0.165	1.0000
1 - 10	-1.32	1.69	96	-7.702	5.062	-0.779	1.0000
1 - 11	-3.00	1.69	96	-9.382	3.382	-1.771	0.9857
1 - 12	-2.16	1.69	96	-8.542	4.222	-1.275	0.9999
1 - 13	-1.60	1.69	96	-7.982	4.782	-0.945	1.0000
1 - 14	0.76	1.69	96	-5.622	7.142	0.449	1.0000
1 - 15	0.76	1.69	96	-5.622	7.142	0.449	1.0000
1 - 16	1.76	1.69	96	-4.622	8.142	1.039	1.0000

[reachedgetOption("max.print") -- omitted 285 rows]

Results are averaged over the levels of: EXAM

Confidence level used: 0.95

Conf-level adjustment: tukey method for comparing a family of 25 estimates

P value adjustment: tukey method for comparing a family of 25 estimates

```
out = summary(pairs(lm1em, infer=c(F,T), level=0.95, adjust="tukey"))
subset(out, out$p.value < 0.05)
```

	contrast	estimate	SE	df	t.ratio	p.value
3	1 - 4	-8.32	1.6939	96	-4.9118	0.00093757984520
25	2 - 3	9.60	1.6939	96	5.6674	0.00004203552076
27	2 - 5	6.72	1.6939	96	3.9672	0.02714856237458
48	3 - 4	-13.84	1.6939	96	-8.1705	0.00000000079632
51	3 - 7	-7.96	1.6939	96	-4.6992	0.00211890975287
54	3 - 10	-6.84	1.6939	96	-4.0380	0.02164538214065
55	3 - 11	-8.52	1.6939	96	-5.0298	0.00058886856553
56	3 - 12	-7.68	1.6939	96	-4.5339	0.00391182008602
57	3 - 13	-7.12	1.6939	96	-4.2033	0.01252639251896
61	3 - 17	-7.60	1.6939	96	-4.4867	0.00464383583599
62	3 - 18	-6.56	1.6939	96	-3.8727	0.03643635567587
64	3 - 20	-8.16	1.6939	96	-4.8173	0.00135190254111
66	3 - 22	-7.88	1.6939	96	-4.6520	0.00252953732975
68	3 - 24	-6.52	1.6939	96	-3.8491	0.03916121121781
69	3 - 25	-7.68	1.6939	96	-4.5339	0.00391182008602
70	4 - 5	10.96	1.6939	96	6.4703	0.00000118997242

```
[ reached 'max' /getOption("max.print") -- omitted 13 rows ]
```