

Design

of

Experiment

(After 2^2 factorial Experiment)

Orthogonality of a Design and Confounding

Suppose, we have a random sample of n observations y_1, y_2, \dots, y_n which are independent from normal population with variance σ^2 .

Let us consider two orthogonal contrasts

$$A = \sum_{i=1}^m \lambda_i y_i \quad \text{with} \quad \sum_{i=1}^m \lambda_i = 0$$

$$\text{and} \quad B = \sum_{i=1}^m \mu_i y_i \quad \text{with} \quad \sum_{i=1}^m \mu_i = 0$$

$$\text{and} \quad \sum_{i=1}^m \lambda_i \mu_i = 0$$

Then, we have covariance (AB)

$$\text{Cov}(A, B) = E(AB) - E(A)E(B)$$

$$\text{Cov}(A, B) = \sigma^2 \sum_{i=1}^m \lambda_i \mu_i = 0$$

$$\left\{ \begin{aligned} E\left[\left(\sum_{i=1}^m \lambda_i x_i\right)\left(\sum_{i=1}^m \mu_i x_i\right)\right] \\ = \sum_{i=1}^m \lambda_i \mu_i E(x_i^2) \end{aligned} \right.$$

Therefore, A and B are independent and we can use A and B to estimate two different effects because the error in two estimates will not be related. These estimates are also said to be orthogonal.

The orthogonality of a design ensures that the different effect will be capable of separate estimations and testing without any difficulties. Hence, if data arise in an orthogonal design then there will be no difficulty of independent estimation and test of effects.

The designs C.R.D., R.B.D. and L.S.D. give an orthogonal design. The 2^2 and 2^3 etc. factorial experiment are conducted by using C.R.D., R.B.D. and L.S.D. However, the difficulty in conducting the factorial experiment in these designs is that as the

number of factors ^{no. of} or levels of factors as both increases, the no. of treatment combinations to be compared increases rapidly. This results the use of large sized blocks or squares to accommodate all the treatment combinations.

For Example - In a 2^4 factorial experiment there are 16 treatment combinations and it is advisable to adopt a R.B.D. for it. Because blocks of 16 plots are too big, ~~there~~ to ensure homogeneity within blocks, therefore, it is necessary to have a new device (technique) for designing experiment with a large number of treatments or treatment combinations. Once, such device is to take blocks of size less than the number of treatment combinations and have more than one block per replication. Then, the treatments are divided into as many groups as number of blocks per replication.

The different groups of treatment combination are allotted to blocks in such a way that the only unimportant treatment combinations get mixed up with the block comparison, these treatment combinations are said to be confounded or mixed up with block effects.

These effects can not be estimated separately. However, the remaining treatment effects which are not confounded with the block effect are still capable of separate estimate in testing. In a confounding design

we lose information on some treatment combinations (completely or partially) which are confounded. Therefore, their should be least important combination is confounded. usually we choose highest order interaction for confounding.

The device of confounding consist in subdividing the replicate into two or more equal sub groups (blocks) and the various treatment combinations into these two or more equal groups of a equal size following certain rules by which we sacrifice some information on certain higher order interaction and allocating the treatment combinations of any group to any block at random.

Confounding in factorial Experiments :->

Confounding in factorial experimental design is an arrangement of treatment combinations in blocks in which less important treatment effects are confounded (mixed up) with the blocks.

There are two types of confounding :-

- (1) Complete confounding
- (2) Partial confounding

when there are two or more replications then the question arises whether we confound some interactions in each replications

and different sets of interaction in different replications.

If ^{the} same set of interactions is confounded in all the replications then confounding is called complete confounding. In complete confounding all information of confounded interaction is lost and we lost all the information from all the replications.

If different sets of interaction are confounded in different replications then confounding is called partial confounding. In this confounding the confounded interactions can be recovered from those replications in which it is not confounded.

Confounding in a 2^3 experiment \rightarrow

In this experiment we have eight treatment combinations under comparison. Suppose, we decide to use two blocks to accommodate eight treatment combinations in a replicate. Thus, we have four plots in each ~~plot~~ block. Now, we are to decide the eight treatment combinations allocate into two groups of four treatment combinations in each group at random.

Remark \rightarrow The 2^n factorial experiment conducted using 2^k blocks. we have to confound $2^{n-k}-1$ treatment combinations in which 2^{n-k} will be generalised and they are automatically confounded.

with the blocks effect. If $k = 1$, then only one treatment combination will be confounding. If $k > 1$ then more than one treatment combination will be confounded according to the above rule.

In this case we have $k = 1$ and only one treatment combination will be confounded. Suppose, we decide the highest order interaction ABC to be confounded.

The interaction ABC is depend on
 $(a-1)(b-1)(c-1) = (abc) - (bc) - (ac) + (c) - (ab) + (b) + (a) - (1)$

Let we apply the four treatment combinations with plus sign in one block and the remaining four treatment combinations with negative sign in second block. Therefore, the block Ist and IInd contain treatment combinations as

+	+	+	+	
abc	a	b	c	→ Block I
1	ab	bc	ac	→ Block II

Here, the contrast measuring the interaction ABC also contain blocks effect i.e.

effect of block first - effect of block second

And we say that the interaction ABC is mixed up or confounded with block effect and we lose information of ABC. However, the other six contrast

of the treatment combinations A, B, C, AB, BC, CA still maintain their orthogonality to the block. 2: each two treatment combination of block Ist (block IInd) with plus sign and two treatment combination with negative sign of the remaining six treatment combinations. Therefore, they will contain no block effects and being orthogonal to ABC and blocks.

Thus, in this allocation of eight treatment combinations to two blocks, there is no difficulty in the estimation or testing of the remaining six treatment combinations.

The first two columns of ANOVA Table for 2^3 confounded experiment is given by

ANOVA TABLE

Source of Variation	Degrees of freedom	Sum of Squares	Mean Sum of Squares	Variance Ratio
Blocks	$2q - 1$	SS(Blocks)	MSS(Blocks)	
A	1	$[A]^2/8q$	MSSA	$F = MSSA/MSE$
B	1	$[B]^2/8q$	MSSB	$F = MSSB/MSE$
C	1	$[C]^2/8q$	MSSC	$F = MSSC/MSE$
AB	1	$[AB]^2/8q$	MSS(AB)	$F = MSS(AB)/MSE$
AC	1	$[AC]^2/8q$	MSS(AC)	$F = MSS(AC)/MSE$
BC	1	$[BC]^2/8q$	MSS(BC)	$F = \frac{MSS(BC)}{MSE}$
Error	$6(q-1)$	SSE	MSSE	
Total	$8q - 1$			

Comparison of Unconfounded and completely Confounded 2^m experiments \rightarrow

We know that the information of n effects contains in an experiment is the reciprocal of the variance of its estimate. In the case of unconfounded design the replicate is itself a block and in this case we denote the error variance by σ^2 . In a completely confounded design the replicate contains two blocks, i.e. a block is a half replicate. Therefore, in this case we denote the error variance by $\sigma^2/2$ and it is expected that $\sigma^2/2 < \sigma^2$, since the smaller block will have greater control over error as compare to the larger block.

The variance of the estimator of an effect (main effect or ~~an~~ interaction effect) of a 2^m eff. experiment in q replicates without confounding is $\frac{\sigma^2}{q 2^{m-2}}$ whereas the variance of the estimator of each unconfounded effect in a 2^m experiment in q replicates (where highest order interaction is completely confounded) is $\frac{\sigma^2/2}{q 2^{m-2}}$. Thus, the information about each effect in an unconfounded design is $\frac{q 2^{m-2}}{\sigma^2}$ whereas the information about each unconfounded effect in a completely confounded design is $\frac{q 2^{m-2}}{\sigma^2/2}$.

Since, $\sigma^2/2 < \sigma^2$

Therefore, the confounded design provides more information ~~of~~ ^{on} unconfounded effect as compared to the unconfounded design. But the confounded design provides no information as zero information about the effect has been completely confounded.

Remark \rightarrow when we are not sure which interactions are unimportant then we cannot sacrifice entire loss of information on that treatment combination. In such cases we distribute the loss of information among more than one treatment combination and we shall get ^{some} information on each of them. Then, this can be ^{assume} by a partially confounded design.

Partial Confounding in a 2^3 Experiment

Here, we have four ~~treatment~~ interactions AB, BC, AC, ABC. we take four replications and two blocks of size 4 in each replicate. The eight treatment combinations are allotted to a block of a replicate in such a way that the interaction AB is confounded in the first replicate, AC is confounded in the second replicate, BC is confounded in the third replicate and ABC is confounded in the fourth replicate. The layout before randomization will be as follows

		Replicate I			
B_1	(1)	(ab)	(c)	(abc)	
B_2	(a)	(b)	(bc)	(ac)	
		AB			

		Replicate II			
B_3	(1)	(a b)	(b)	(abc)	
B_4	(a)	(c)	(bc)	(ab)	
		AC			

B5	(1)	(bc)	(a)	(abc)
B6	(b)	(c)	(ac)	(ab)
	BC			

B7	(1)	(Ab)	(bc)	(ac)
B8	(a)	(b)	(c)	(abc)
	ABC			

The first two columns of Anova table in this case will be

S.S.	d.f.
Blocks	7
A	1
B	1
C	1
AB	1
AC	1
BC	1
ABC	1
Error	17
Total	31

The block sum of squares is computed from the eight blocks totals and the grand total. The sum of squares due to the main effects A, B and C (unconfounded effects) are computed using data from all the four replicates whereas the sum of squares due to any confounded interaction is obtained from those replicates where that particular interaction is not confounded.

Ques \rightarrow Given a block how to find the interaction confounded

First of all find the key block. If key block is not given then key block may be obtained by selecting any treatment combination from the given block and multiplying all the treatment combinations in the block by that treatment combination. From the key block we can find the number of factors as well as the block size ($2^5; 2^3$). Now, find out the unit matrix of order k .

Example \rightarrow

		A	B	C	D	E
a c d e	a c d	1	0	1	1	0
b c d	b e d e	0	1	1	1	1
e	1	0	0	0	0	0
a b c e e	a b c	1	1	1	0	0
a d	a d e	1	0	0	1	1
b d e	b d	0	1	0	1	0
a b	a b e	1	1	0	0	1
c	c e	0	0	1	0	1

A	B	C	D	E
1	0	0	1	1
0	1	0	1	0
0	0	1	0	1

$$D A' B' C' = A B C$$

$$E A' B' C' = A C E$$

$\Rightarrow B C D E$
it is confounded.

In the similar way we can find out other two unit matrices

General rule for Confounding in 2^m series \rightarrow

let the design is $(2^m, 2^k)$. Therefore,

$$\text{Treatment Combination} = 2^k$$

Block size = 2^k

No. of block per replication = $\frac{2^n}{2^k} = 2^{n-k}$

Total no. of interaction confounded = $2^{n-k} - 1$

No. of independent interaction confounded = $n-k$

Generalised interaction confounded = $2^{n-k} - 1$ (n-k)

Example \rightarrow

		A	B	C
a	1	0	0	0
b	ab	1	1	0
ac	c	0	0	1
bc	abc	1	1	1

A	B	C
1	1	0
0	0	1

AB^1C^0

$\Rightarrow AB$ is confounded.

Ques.

Comparison of the information about unconfounded effect and confounded effect in

Partial Confounding Design \rightarrow

In partial confounded design the main effects A, B, C are not confounded in any replicate. So, that they are estimated from all four replicates and the experiment contains $8/2 = 4$ information about each of the main effects. But each interaction is confounded in one replicate and unconfounded in the remaining three repli.

cates. Thus, we can estimate the interaction
from those replicates where it is not confou-
 nded. For Example if the interaction AB is
 confounded from the replicates I, Σ , this can
 be estimated using replicates II, III and IV. Only
 these replicates contains information about
 the interaction AB , and the amount of the
 information is $6/\sigma^2/2$. Thus, relative informa-
 tion of each partially confounded interaction
 with respect to the unconfounded main
 effect is

$$\frac{\frac{6}{\sigma^2/2}}{\frac{6}{\sigma^2}} = \frac{3}{4}$$

and this is the same as the proportion of
 replicates given information about the confou-
 nded interaction.

Table for amount of information about
different confounded interaction in 2^3 ex-
periments \rightarrow

effect	Amount of information		
	Unconfounded design	ABC completely confounded	AB, BC, AC, ABC partially confou- unded
A	$6/\sigma^2$	$6/\sigma^2/2$	$6/\sigma^2/2$
B	$6/\sigma^2$	$6/\sigma^2/2$	$6/\sigma^2/2$
AB	$6/\sigma^2$	$6/\sigma^2/2$	$6/\sigma^2/2$
C	$6/\sigma^2$	$6/\sigma^2/2$	$6/\sigma^2/2$
AC	$6/\sigma^2$	$6/\sigma^2/2$	$6/\sigma^2/2$
BC	$6/\sigma^2$	$6/\sigma^2/2$	$6/\sigma^2/2$
ABC	$6/\sigma^2$	0	$6/\sigma^2/2$

$$\text{Plot Size} = \frac{2^n}{2^k} = 2^{n-k}$$

Confounding in 2^4 experiment in 2^2 blocks

In this case we have 16 treatment combinations and these treatment combinations are allotted for the blocks in a replicate. The total no. of confounded effects are $2^{4-2} - 1 = 3$ in which $2^2 - 1 - 2 = 1$ will be generalised interaction. Suppose, we select ABC and ABD for confounding then

$$\begin{aligned} ABCABD &= A^2B^2CD \\ &= A^0B^0CD = CD \end{aligned}$$

will also be confounded.

we can make first the key blocks of ABC and ABD then we make confounded blocks

The 16 treatment combinations can be written in a systematic order as follows: -
 1, a, b, ab, c, ac, bc, abc,
 d, ad, bd, abd, cd, acd, bcd, abcd

B₁

(1)
(ab)
(acd)
(bcd)

B₂

(a)
(b)
(cd)
(abcd)

B₃

(c)
(abc)
(ad)
(bd)

B₄

(d)
(abd)
(ac)
(bc)

Confounding in more than two blocks

For a 2^4 experiment 2 blocks per replication are reasonable in experiment with a large number of factors. We however, use blocks (72) per replicate confounding with two groups one interaction is per annum, more than one interaction is confounded in such cases the key block contains ~~after~~ are obtained from the 2^{4-1} more than one homogeneous equations simultaneously.

For example is the key block of size 2^3 in 2^5 factorial experiment can be obtained from the solutions of the following equations

$$x_1 + x_2 + x_3 = 0$$

$$x_1 + x_4 + x_5 = 0$$

These equations indicate that the interactions ABC and ADE are confounded simultaneously in the same replication. Any solution of the above two homogeneous equations is also a solution of the equation which is obtained from the linear combination of the equations. This shows that some other interaction is also confounded. On adding the above two equations we get

$$x_2 + x_3 + x_4 + x_5 = 0$$

No other equation is possible from their linear combination. Therefore, the interaction BCDE is also confounded which is called generalised interaction of ABC and ADE. The key block can be obtained by first obtaining three independent

Solutions of the homogeneous equations and then taking all linear combinations. In this way we get the key block

Key block					
0	0	0	0	0	1
0	1	1	1	1	b c d e
1	0	1	0	1	a c e
1	0	1	1	0	a c d
1	1	0	1	0	a b d
1	1	0	0	1	a b e
0	0	0	1	1	d e
0	1	1	0	0	b c

There are three more blocks in a replicate which are obtained from the solutions of ^{the following} three sets of equations

$$\left. \begin{aligned} x_1 + x_2 + x_3 &= 0 \\ x_1 + x_4 + x_5 &= 1 \end{aligned} \right\}$$

$$\left. \begin{aligned} x_1 + x_2 + x_3 &= 1 \\ x_1 + x_4 + x_5 &= 0 \end{aligned} \right\}$$

$$\left. \begin{aligned} x_1 + x_2 + x_3 &= 1 \\ x_1 + x_4 + x_5 &= 1 \end{aligned} \right\}$$

3^m Experiments \rightarrow when factors are taken at three levels instead of ~~2~~ the experiment becomes more informative. Let m factors be denoted by A, B, C, \dots so on and each having three levels. The levels of three factors are denoted by 0, 1 and 2. The possible combinations of these three levels of each factor will give rise to 3^m treatment combinations. Here, we use the number system reduced modulo 3, i.e.

$$0 = 3 = 6 = 9 = \dots$$

$$1 = 4 = 7 = 10 = \dots$$

$$2 = 5 = 8 = 11 = \dots$$

In this system we divide a number greater than or equal to 3 by 3 and take the remainder to be equal to the original number.

3^2 - factorial Experiments \rightarrow

In this experiment there are two factors (let) A and B each having three levels 0, 1 and 2. Therefore, there are 9 treatment combinations of type (x_1, x_2) where, x_1 and x_2 take any of the values 0, 1 and 2. Thus, 9 treatment combinations are $(0,0), (1,0), (0,1), (1,1), (2,0), (0,2), (1,2), (2,1), (2,2)$ or $(00, 10, 01, 11, 20, 02, 12, 21, 22)$. Among these 9 treatment combinations there will be 8 comparisons i.e. degrees of freedom = 8 can be partitioned as 2 d.f. for each of main effects

A and B and four d.f. for the interaction AB. These components can be obtained by a two-way table as follows.

		level of A			
level of B	0	00	10	20	Total for B [B] ₀
	1	01	11	21	[B] ₁
	2	02	12	22	[B] ₂
Total for A		[A] ₀	[A] ₁	[A] ₂	M

The sum of squares can be obtained as

$$SSA = \frac{[A]_0^2 + [A]_1^2 + [A]_2^2}{3q} - \frac{M^2}{9q}$$

where, q is the number of replicate.

$$SSB = \frac{[B]_0^2 + [B]_1^2 + [B]_2^2}{3q} - \frac{M^2}{9q}$$

$$SS(A \times B) = \frac{[00]^2 + [10]^2 + [20]^2 + [01]^2 + \dots + [22]^2}{q} - \frac{M^2}{9q}$$

The four degrees of freedom of A x B can be further partition into two more orthogonal contrast each having two degrees of freedom. These components are very useful in confounding in the 3rd experiment. Thus, the interaction A x B carrying four degrees of freedom can be partition as AB and AB².

The defining equations divide the 9 treatment total into three groups, a comparison between which give the corresponding components thus $A \times B$ can be divided as AB and AB^2 .

$$x_1 + x_2 = 0 \text{ gives } [00] + [12] + [21] = [AB]_0$$

$$x_1 + x_2 = 1 \text{ gives } [01] + [10] + [22] = [AB]_1$$

$$x_1 + x_2 = 2 \text{ gives } [02] + [20] + [11] = [AB]_2$$

$$x_1 + 2x_2 = 0 \text{ gives } [00] + [11] + [22] = [AB^2]_0$$

$$x_1 + 2x_2 = 1 \text{ gives } [02] + [10] + [21] = [AB^2]_1$$

$$x_1 + 2x_2 = 2 \text{ gives } [01] + [12] + [20] = [AB^2]_2$$

Therefore, the sum of squares due to component AB is given by

$$SS(AB) = \frac{[AB]_0^2 + [AB]_1^2 + [AB]_2^2}{3q} = \frac{M^2}{9q}$$

and

$$SS(AB^2) = \frac{[AB^2]_0^2 + [AB^2]_1^2 + [AB^2]_2^2}{3q} = \frac{M^2}{9q}$$

The first two columns of anova table are

Source of variation	Degrees of freedom
Replication	$(q-1)$
A	2
B	2
$A \times B \rightarrow AB$	2
$A \times B \rightarrow AB^2$	2
Error	$3(q-1)$
Total	$9q-1$

3³ - Experiment

In this experiment we have 27 treatment combinations are of the form (x_1, x_2, x_3) where x_1, x_2 and x_3 are the levels of 3 factors A, B and C respectively, all the treatment combinations can systematically be written as

000	001	002
100	101	102
010	011	012
110	111	112
200	201	202
210	211	212
020	021	022
120	121	122
220	221	222

The 27 treatment combinations will have a sum of squares carrying 26 degrees of freedom. The treatment sum of squares can be calculated from the 27 treatment totals taking q replicates. In this experiment the treatment combinations can be subdivided into the main effects and interaction effects having degrees of freedom as follows

Replicates $-(q-1)$, T.C. $- 26$,

A $- 2$, B $- 2$, C $- 2$, A \times B $- 4$,

A \times C $- 4$, B \times C $- 4$, A \times B \times C $- 8$

Errors $- 26(q-1)$, Total $-(27q-1)$

The sum of squares due to main effect and two factor interaction AXB, AXC, BXC calculated from the three, two-way tables in the usual manner. However, the three factor interaction sum of squares is obtained by subtraction of these components from the treatment sum of squares.

Here, each set of four or eight degrees of freedom can also be partitioned into their orthogonal contrast (components) carrying 2 degrees of freedom as follows

Components of SS	Defining Equation	Mod
$AXB \begin{matrix} \nearrow AB \\ \searrow AB^2 \end{matrix}$	$x_1 + x_2 = 0, z_1 = 2$	3
	$x_1 + 2x_2 = 0, z_1 = 2$	3
$AXC \begin{matrix} \nearrow AC \\ \searrow AC^2 \end{matrix}$	$x_1 + x_3 = 0, z_1 = 2$	3
	$x_1 + 2x_3 = 0, z_1 = 2$	3
$BXC \begin{matrix} \nearrow BC \\ \searrow BC^2 \end{matrix}$	$x_2 + x_3 = 0, z_1 = 2$	3
	$x_2 + 2x_3 = 0, z_1 = 2$	3
$ABC \begin{matrix} \nearrow ABC \\ \searrow ABC^2 \\ \searrow AB^2C \\ \searrow AB^2C^2 \end{matrix}$	$x_1 + x_2 + x_3 = 0, z_1 = 2$	3
	$x_1 + x_2 + 2x_3 = 0, z_1 = 2$	3
	$x_1 + 2x_2 + x_3 = 0, z_1 = 2$	3
	$x_1 + 2x_2 + 2x_3 = 0, z_1 = 2$	3

Using each of the defining equations we divide the 27 treatment combinations into three groups and a comparison among these three combinations totals is the correct sum of squares carrying 2 d.f.

as Example: The sum of squares due to components AB^2C can be obtained as follows

$$SS(AB^2C) = \frac{[AB^2C]_0^2 + [AB^2C]_1^2 + [AB^2C]_2^2}{99} - \frac{\mu^2}{279}$$

where,

$$[AB^2C]_0 = [000] + [011] + [110] + [201] + [212] + [102] + [121] + [220]$$

$$[AB^2C]_1 = [001] + [100] + [012] + [111] + [202] + [210] + [020] + [122] + [221]$$

$$[AB^2C]_2 = [002] + [101] + [010] + [112] + [200] + [211] + [021] + [120] + [222]$$

Similarly the other sum of squares can be obtained by using their defining equations. The each mean sum of squares can be obtained by dividing the corresponding sum of squares by its degrees of freedom.

ANOVA table first two columns

SV	degrees of freedom
Replication	(9-1)
T.C.	26
A	2
B	2
AxB → AB	2
→ AB ²	2
AxC → AC	2
→ AC ²	2

C	2
B x C	4
A x B x C	8
Error	$26(q-1)$
Total	$(27q-1)$

Split Plot Design - In conducting experiments sometimes some factors have to be applied in larger experimental units while some other factors can be applied in comparatively smaller experimental units.

For example ~~if~~ when we wish to compare the different methods of irrigation and ploughing. In such cases it is possible to introduce a second factor which does not require large plot, with a small number of levels into the same experiment at a little extra cost. This can be done by splitting the plot of the first factor into as many sub plots as there are levels of the second factor.

A split plot design is a design with atleast one blocking factor where the experimental units within each block are assigned to the treatment factor levels as usual, and in addition the blocks are assigned at random to the levels of a further treatment factor. The designs have a nested blocking structure. In a split plot design the experimental units are called split plots or sub plots and are nested within the whole plots or main plots.

Layout \rightarrow And Randomization \rightarrow

There are two separate randomization processes in a split plot design.

- (1) For the main plot
- (2) For the subplots.

In each replication main plot treatments are first randomly assign to the main plots followed by a random assignment on the ~~same~~ ^{sub} plot treatments within each main plot treatment. This procedure is applied for all replications. A possible layout of a split plot experiment with four main plot treatments, three sub plot treatments and two replications is given below

Replicate - I

b ₂	b ₁	b ₂	b ₁
b ₁	b ₃	b ₃	b ₂
b ₃	b ₂	b ₁	b ₃
a ₁	a ₄	a ₃	a ₂

Replicate - II

b ₁	b ₃	b ₂	b ₃
b ₂	b ₁	b ₃	b ₂
b ₃	b ₂	b ₁	b ₁
a ₂	a ₄	a ₁	a ₃

The above layout has the following important features: →

- (1) The size of the main plot is q times the size of the sub plot.
- (2) Each main plot treatment is tested q times whereas each sub plot treatment is tested $p \cdot q$ times, thus number of times a subplot treatment

as tested will always be larger than that for the main plot. Therefore, the subplot treatment have the more precision as compare to the main plot treatment.

Analysis \rightarrow let p levels of factor A be randomized by using the layout of RBD and LSD and the q levels of factor B are then randomized within each whole plot of factor A by dividing each whole plot into q subplots.

Thus, for split plot design we use the following model.

$$Y_{ijk} = \mu + b_i + \tau_j + e_{ij} + \gamma_k + \delta_{jk} + e'_{ijk}$$

$$i=1, 2, \dots, p, \quad j=1, 2, \dots, q, \quad k=1, 2, \dots, q$$

Here,

τ_j , γ_k and δ_{jk} are the fixed effects due to the j th level of factor A, k th level of factor B and interaction between them respectively.

Thus, obviously,

$$\sum_j \tau_j = \sum_k \gamma_k = \sum_j \delta_{jk} = \sum_k \delta_{jk} = 0$$

b_i , e_{ij} , e'_{ijk} are random components which are independently normally distributed with mean zero and respective variance such that

$$b_i \text{ i.i.d. } N(0, \sigma_b^2)$$

$$e_{ij} \text{ i.i.d. } N(0, \sigma_e^2)$$

$$e'_{ijk} \text{ i.i.d. } N(0, \sigma_{e'}^2)$$

In this design the analysis can be done into two stages, at the 1st stage we use the analysis of

RBD with p treatment in r blocks, the whole plot analysis will be as follows

S.V:	d.f.	Sum of Squares
Blocks	$r-1$	$bq \sum_i (\bar{y}_{i..} - \bar{y}_{000})^2$ = SS _B
whole plot treatment (A)	$(p-1)$	$rq \sum_j (\bar{y}_{.j0} - \bar{y}_{000})^2$ = SS _A
whole plot error E_I	$(r-1)(p-1)$	$q \sum_{i,j} (\bar{y}_{ij0} - \bar{y}_{i..} - \bar{y}_{.j0} + \bar{y}_{000})^2$ = SS _{E_I}
Total	$rp-1$	$q \sum_{i,j} (\bar{y}_{ij0} - \bar{y}_{000})^2$

Here, it can be shown that

$$E(MSSA) = \sigma_e^2 + q\sigma_e^2 + \phi_1(\tau_j)$$

$$E(MSSE_I) = \sigma_e^2 + q\sigma_e^2$$

Under H_{01} : All the effects due to the j th levels of A are same, i.e.

$$H_{01}: \tau_j = 0 \forall j$$

$$\phi_1 = 0$$

otherwise, $\phi_1 > 0$

Thus, a test for testing H_{01} is provided by

$$F = \frac{MSSA}{MSSE_I} \sim F(\alpha) \left\{ (p-1), (r-1)(p-1) \right\}$$

The second stage will be the analysis of the sub plot analysis within whole blocks which is as follows :-

S.V.	d.d.	Sum of Squares
Sub plot treatment (B)	$(q-1)$	$rp \sum_k (\bar{y}_{00k} - \bar{y}_{000})^2 = SS_B$
Interaction (AB)	$(p-1)(q-1)$	$rs \sum_{j,k} (\bar{y}_{0jk} - \bar{y}_{0j0} - \bar{y}_{00k} + \bar{y}_{000})^2 = SS(AB)$
Sub Plot Error (EII)	$p(q-1)$ $(p-1)$	$\sum_{i,j,k} (y_{ijk} - \bar{y}_{0jk} - \bar{y}_{i00} + \bar{y}_{000})^2 = SS_{EII}$
Total between Sub Plots within whole Plots	$(rp-1)$ $rp(q-1)$	$\sum_{i,j,k} (y_{ijk} - \bar{y}_{i00})^2$

Complete analysis of variance table for split plot design

S.V.	d.f.	SS	MSS	F [MSSE]	F
Blocks	$(r-1)$	$SS(\text{blocks})$	$MSS(\text{blocks})$	$\frac{SS(\text{blocks})}{r-1}$ $\frac{SS(\text{blocks})}{r-1}$	$\frac{MSSA - MSSA}{MSS_{E_I}}$
Treat (A)	$(p-1)$	SSA	$MSSA$	$\sigma_e^2 + \rho \sigma_c^2 + \rho_I (\tau_j^2)$	
Error I	$(r-1)(p-1)$	SSE_I	$MSSE_I$	$\sigma_e^2 + \rho \sigma_c^2$	
Treat (B)	$(q-1)$	SSB	$MSSB$	$\sigma_e^2 + \rho_{II} (\tau_k^2)$	$\frac{MSSB}{MSSE_{II}}$
Interaction (AB)	$(p-1)(q-1)$	$SS(AB)$	$MSS(AB)$	$\sigma_e^2 + \rho_{III} (\delta_{jk}^2)$	$\frac{MSS(AB)}{MSS_{E_{II}}}$
Error II	$p(q-1)$ $p(q-1)$	SSE_{II}	$MSSE_{II}$	σ_e^2	$\frac{MSS_{E_{II}}}{MSS_{E_{II}}}$
Total	$rpq-1$	TSS			

Merits and Demerits of Split Plot Design

Merits →

- (1) In this design there are two errors type I and type II where $E_I > E_{II}$. Therefore, the main effect B and interaction AB will be estimated and tested.
- (2) In this design we can introduce the second factor B which requires small experimental material along A at a little extra cost.
- (3) If we have a choice of allocation and the factors A and B then we shall apply most important factors to the sub plot rather than whole plots.

Demerits →

The presence of the two errors make the analysis difficult. Also, sometimes the error I may be too large.

Strip Plot Design → In split plot design one factor requires smaller unit as compare to the other, and we can increase the precision on the factor B and factor AB as such.

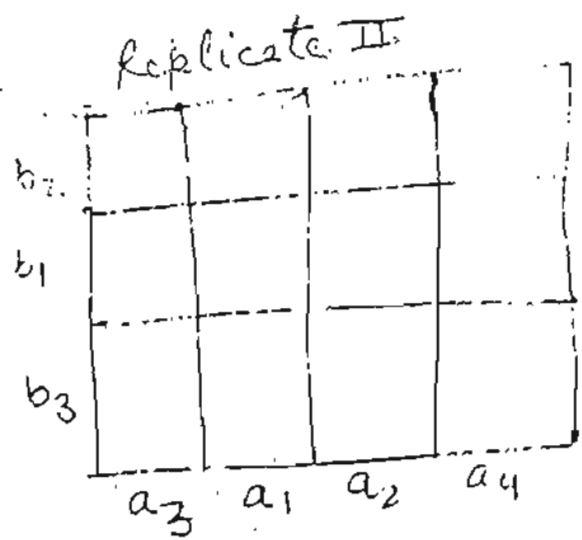
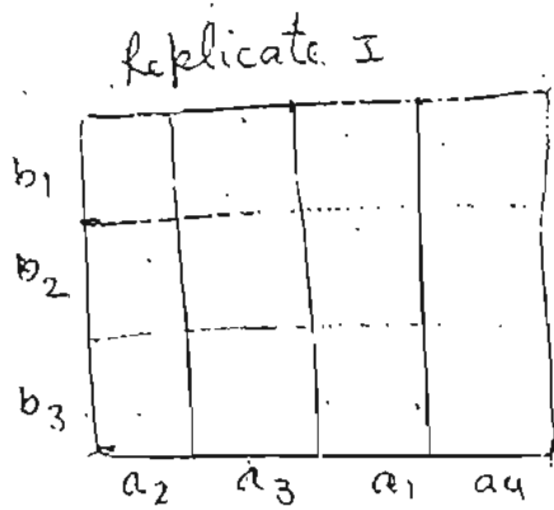
sacrifices some precision on A. In that design the factor A is of less important than factor B. However, sometimes we may have factors A and B each requiring large units.

For Example \rightarrow Suppose, four levels of spacing and three levels of method of ploughing are to be tested in the same experiment. Therefore, to accommodate both the factors in the large units we use strip plot design.

Randomization and Layout \rightarrow

In this design each block (replicate) is divided into number of vertical and horizontal strips depending on the levels of respective factors. Let factor A represent the vertical factor with p levels, factor B represent the horizontal factor with q levels and r represent the number of replication. To layout the experiment, the experimental area is divided into r blocks. Each block is divided into q horizontal strips and q treatments are randomly assign to these strips in each of the r blocks separately and independently. Then, each block is divided into p vertical strips and p treatments are randomly assign to these strips in each of the r replicates separately and independently.

A possible layout of a strip plot design with $p = 4, (a_1, a_2, a_3, a_4), q = 3, (b_1, b_2, b_3)$ and $r = 2$.



~~Analysis~~ → The strip plot design sacrifices precision on the main effects of both the factors in order to provide higher precision on the interaction which will usually be more accurately determined ~~than~~ than split plot design.

Analysis → In this design we have three errors for different effects. The analysis of variance for a strip plot design will be based on the model

$$\sqrt{y_{ijk}} = \mu + \alpha_i + \alpha_j + (\alpha\alpha)_{ij} + \beta_k + (\beta\beta)_{j'k} + (\alpha\beta)_{i'k} + (\alpha\alpha\beta)_{i'jk}$$

where y_{ijk} denote the response or output of the yield receiving j th level of factor A, k th level of factor B in the i th replicate. Thus, $i = 1, 2, \dots, r$; $j = 1, 2, \dots, p$; $k = 1, 2, \dots, q$.

$(\alpha\alpha)_{ij}$, $(\beta\beta)_{j'k}$ and $(\alpha\alpha\beta)_{i'jk}$ are three errors which are such that

$$(\alpha\alpha)_{ij} \stackrel{i.i.d.}{\sim} N(0, \sigma_1^2)$$

$$(\beta\beta)_{j'k} \stackrel{i.i.d.}{\sim} N(0, \sigma_2^2)$$

$(\sigma^2 \alpha^2)_{ijk}$ ~~is~~ $N(\sigma^2, \sigma^2)$

$\alpha_j, \beta_k, (\alpha\beta)_{jk}$ are fixed effects.

after doing some mathematical calculations we obtain the following analysis of variance table.

S.V.	d.f.	SS	MSS	E(MSS)	F
Replicate	$(r-1)$	SSR	MSSR	$\frac{\sigma^2 + \rho\sigma_1^2}{r}$	MSSA
Treat (A)	$(p-1)$	SSA	MSSA	$\frac{\sigma^2 + \rho\sigma_1^2}{(p-1)}$	
Errors E _I	$(b-1)(p-1)$	SS E _I	MSSE _I	$\frac{\sigma^2 + \rho\sigma_1^2}{(b-1)}$	
Treat (B)	$(q-1)$	SSB	MSSB	$\frac{\sigma^2 + \rho\sigma_2^2}{q}$	MSSB
Errors E _{II}	$(b-1)(q-1)$	SS E _{II}	MSSE _{II}	$\frac{\sigma^2 + \rho\sigma_2^2}{(b-1)}$	MSSE _{II}
Treat (AB)	$(p-1)(q-1)$	SS(AB)	MSS(AB)	$\frac{\sigma^2 + \rho\sigma_3^2}{(p-1)}$	MSS(AB)
Errors E _{III}	$(b-1)(p-1)(q-1)$	SS E _{III}	MSSE _{III}	$\frac{\sigma^2 + \rho\sigma_3^2}{(b-1)}$	MSSE _{III}
Total	$bpq-1$	TSS			

Balanced Incomplete Block Design (BIBD)

In a confounded design we sacrifice some or total information on any of the treatment combinations to maintain the homogeneity within block. Sometimes we come across the situation

where all the treatments have an equal importance then we can not afford to sacrifice information on any of them by confounding. In such situations we use BIBD. The BIBD was first proposed by Yates in 1936.

Here incomplete block means a block which do not contain complete set of treatments. The use of incomplete block becomes necessary because as the number of treatments increases the block size increases which results and increase in the heterogeneity. When the number of replications of all pairs of treatments in an incomplete blocks is same then an important series of design known as balanced incomplete block design is obtained. In this design all the treatment effects are estimated with equal precision.

→ $\sum_{k=1}^m \lambda = 3$

Incomplete Block Designs →

A block design which consist v treatments in b blocks each of size k such that each of the treatment is replicated r times and each pair of treatments occurs once and only once in the same block is called incomplete block designs.

Example → Suppose, $v = b = 5$

$$r = k = 4$$

	I	1	2	3	4
Blocks	II	1	2	3	5
	III	1	2	4	5
	IV	1	3	4	5
	V	2	3	4	5

Parameters of BIBD \rightarrow BIBD has five parameters which are given as

v = no. of treatments

b = no. of blocks

k = block size

r = no. of replicates for each treatment

λ = no. of ~~blocks~~ replication of each treatment pair or no. of blocks in which any pair of treatments occurs together.

Definition of BIBD \rightarrow A BIBD is defined as,

"An arrangement of v treatments in b blocks of each of size k ($k < b$) such that each treatment is replicated r times. No treatment occurs more than one in a block and each pair of treatment occurs together in λ blocks

Incidence Matrix \rightarrow A matrix $N = (n_{ij})$ $\begin{matrix} i=1,2,\dots,v \\ j=1,2,\dots,b \end{matrix}$

associated with any design D , where n_{ij} denotes

The no. of times the i th treatment occurs in the j th block, is called incidence matrix. Thus, by the definition of the BIBD

$$n_{ij} = \begin{cases} 1 & \text{if } i\text{th treatment occurs in the } j\text{th block} \\ 0 & \text{otherwise} \end{cases}$$

For the existence of a BIBD for the parameteric relation of BIBD \rightarrow

(1) $rv = bk$

(2) $\lambda(v-1) = r(k-1)$

(3) $b \geq v$

(iv) $b > r > \lambda$

(v) $r \geq k$ and $v > k$

(vi) $b \geq v + r - k$

(1) $rv = bk$

Proof \rightarrow Here, v is the no. of treatment in a replication therefore, rv will be total no. of plots in r replicates.

b is the no. of blocks in r replicates and k is the no. of plots in a block therefore bk is the total no. of plots in r replicate.

Hence, $rv = bk$. Proved

$$(2) \lambda(u-1) = r(k-1)$$

Proof \rightarrow Since v -treatments give rise to $\binom{v}{2}$ pairs and since each pair occurs λ times. So, total no. of times of the pairs occurs in the design is $\lambda \binom{v}{2}$.

Further, since the size of each block is k . So, each block give rise to $\binom{k}{2}$ pairs and total no. of treatment pairs in all the b blocks is $b \binom{k}{2}$.

$$\text{Therefore, } \lambda \binom{v}{2} = b \binom{k}{2}$$

$$\lambda \frac{v!}{2!(v-2)!} = b \frac{k!}{2!(k-2)!}$$

$$\Rightarrow \lambda v(v-1) = b k(k-1) \quad ; \text{ since } rv = bk$$

$$\Rightarrow \lambda r(v-1) = r k(k-1)$$

$$\Rightarrow \lambda(v-1) = r(k-1) \quad \underline{\underline{\text{Proved}}}$$

(3) $b \geq v$ (Fisher inequality)

Proof \rightarrow let us consider an incidence matrix

$$N = (n_{ij})_{v \times b} \quad ; \quad i = 1, 2, \dots, v, \quad j = 1, 2, \dots, b$$

where,

$$n_{ij} = \begin{cases} 1 & ; \text{ if } i\text{th treatment occurs in the } j\text{th block} \\ 0 & ; \text{ otherwise} \end{cases}$$

$$\left\{ \text{let } NN' = \begin{bmatrix} n_{11} & n_{12} & \dots & n_{1b} \\ \vdots & \vdots & \ddots & \vdots \\ n_{v1} & n_{v2} & \dots & n_{vb} \end{bmatrix} \begin{bmatrix} n_{11} & \dots & n_{1b} \\ n_{12} & \dots & n_{2b} \\ \vdots & \ddots & \vdots \\ n_{v1} & \dots & n_{vb} \end{bmatrix} \right. \left. \begin{matrix} n_{11}^2 + n_{12}^2 + \dots + n_{1b}^2 \\ \vdots \\ n_{v1}^2 + n_{v2}^2 + \dots + n_{vb}^2 \end{matrix} \right.$$

$$\text{Let } NN^T = \begin{bmatrix} \sum_j m_{1j}^2 & \sum_j m_{1j} m_{2j} & \dots & \sum_j m_{1j} m_{uj} \\ \sum_j m_{2j} m_{1j} & \sum_j m_{2j}^2 & \dots & \sum_j m_{2j} m_{uj} \\ \vdots & \vdots & \ddots & \vdots \\ \sum_j m_{uj} m_{1j} & \sum_j m_{uj} m_{2j} & \dots & \sum_j m_{uj}^2 \end{bmatrix}_{u \times b}$$

where, for BIBC $\sum_{j=1}^b m_{ij} = m_{i1} + m_{i2} + \dots + m_{ib} = r_i$
 $= r_i = \sum_{j=1}^b m_{ij}^2$

Similarly,

$$\sum_{i=1}^u m_{ij} = m_{1j} + m_{2j} + \dots + m_{uj} = k_j$$

$$= k_j = \sum_{i=1}^u m_{ij}^2$$

Remark

$$\begin{bmatrix} m_{11} & m_{12} & \dots & m_{1b} \\ m_{21} & m_{22} & \dots & m_{2b} \\ \vdots & \vdots & \ddots & \vdots \\ m_{u1} & m_{u2} & \dots & m_{ub} \end{bmatrix}_{u \times b}$$

$\rightarrow r_i$
 $\rightarrow k_j$

Sum of the i th row of a incidence matrix $= r_i$
 Sum of the j th column of a incidence matrix $= k_j$

$$\sum_{j=1}^b m_{ij} m_{ij} = \lambda (r_i + k_j)$$

Thus

$$NN^T = \begin{bmatrix} r_1 & \lambda & \dots & \lambda \\ \lambda & r_1 & \dots & \lambda \\ \vdots & \vdots & \ddots & \vdots \\ \lambda & \lambda & \dots & r_1 \end{bmatrix}_{u \times u}$$

$$NN' = \begin{bmatrix} \alpha + (\alpha-1)\lambda & \lambda & - & - & \lambda \\ \alpha + (\alpha-1)\lambda & \alpha & - & - & \lambda \\ \vdots & \vdots & & & \\ \alpha + (\alpha-1)\lambda & \lambda & - & - & \alpha \end{bmatrix} \quad C_1 \rightarrow C_1 + C_2 - C_3$$

$$= \begin{bmatrix} \alpha + (\alpha-1)\lambda & \lambda & - & - & \lambda \\ 0 & (\alpha-\lambda) & - & - & 0 \\ \vdots & \vdots & & & \\ 0 & - & - & 0 & - & - & - & (\alpha-\lambda) \end{bmatrix}$$

$R_2 \rightarrow R_2 - R_1$
 $R_3 \rightarrow R_3 - R_1$
 $R_i \rightarrow R_i - R_1 \quad (i = 3, \dots, \alpha)$

$$= [\alpha + (\alpha-1)\lambda] \begin{bmatrix} 1 & \lambda & - & - & \lambda \\ 0 & (\alpha-\lambda) & - & - & 0 \\ \vdots & \vdots & & & \\ 0 & - & - & 0 & - & - & - & (\alpha-\lambda) \end{bmatrix}$$

$$\Rightarrow [\alpha + (\alpha-1)\lambda] \begin{bmatrix} 1 & - & - & - & 0 \\ \vdots & (\alpha-\lambda) & - & - & \\ \vdots & \vdots & & & \\ 0 & - & - & - & (\alpha-\lambda) \end{bmatrix}$$

$$|NN'| = [\alpha + (\alpha-1)\lambda] (\alpha-\lambda)^{\alpha-1} \neq 0$$

Therefore, Rank of $NN' = \alpha$ (because the order of $NN' = \alpha$)

$\Rightarrow \rho(NN') \leq \rho(N) \leq \min(a, \alpha)$

But we know that

$$n \geq \text{Rank of } NN'$$

$$\Rightarrow b \geq \alpha$$

Proved

$$(iv) \quad b > r > \lambda$$

Proof \rightarrow Since, r is the no. of times i th treatment is replicated and λ is the no. of times, i th treatment occurs with j th treatment and since due to incomplete blocks the i th treatment cannot occur r times with j th treatment. Therefore.

$$\lambda < r \quad \text{--- (1)}$$

Again, since $v > k$ (For BIBD)

$$\Rightarrow v > \frac{rv}{b} \quad \because bk = rv$$

$$\Rightarrow b > r \quad \text{--- (2)}$$

from eqⁿ (1) and (2)

$$b > r > \lambda$$

Proved

$$(v) \quad r \geq k \text{ and } v > k$$

Proof \rightarrow we know that

$$\begin{aligned} bk &= rv \\ \Rightarrow \frac{b}{v} &= \frac{r}{k} \quad \text{--- (1)} \end{aligned}$$

But we have $b \geq v$

$$\Rightarrow \frac{b}{v} \geq 1$$

on putting in eqⁿ (1) we get

$$\Rightarrow \frac{r}{k} \geq 1$$

$$\Rightarrow r \geq k$$

So, $v > k$

Proved

(vi) $b \geq u + r - k$
 Proof: we know that
 $r \geq k$ and $u > k$ (for BIBD)

Therefore, $(r - k) \geq 0$ and $(u - k) > 0$

$$(r - k)(u - k) \geq 0$$

$$ru - kv - rk + k^2 \geq 0$$

$$bk - kv - rk + k^2 \geq 0$$

$$k(b - v - r + k) \geq 0$$

$$b - v - r + k \geq 0$$

$$b \geq v + r - k$$

Proved

Symmetrical BIBD \rightarrow

A BIBD is said to be symmetrical if $b = u$ and $r = k$. In this case the incidence matrix N is a square matrix.

"An arrangement of v treatments in b blocks of each of size k ($k < b$) such that each treatment is replicated r times, no treatment occurs more than one in a block and each pair of treatment occurs together in λ blocks."

For $n=2$

Theorem \rightarrow For a symmetrical BIBD (v, λ) will be a perfect square when v is even.

Proof \rightarrow we have

$$|NN'| = r k (r - \lambda) v - 1$$

$$\Rightarrow |N| |N'| = r^2 (r - \lambda) v - 1 \quad (\text{for } r = k)$$

$$\Rightarrow |N|^2 = r^2(r-1)^{u-1}$$

$$\Rightarrow |N| = \pm r(r-1)^{(u-1)/2}$$

Since, the determinant of an incidence matrix N is an integer.

Therefore, $r(r-1)^{(u-1)/2}$ is an integer. Also, r is the integer. Therefore $(r-1)^{(u-1)/2}$ is integer. Thus, $(r-1)^{(u-1)/2}$ will be a perfect square, when it is ~~even~~ even. Proved

Theorem 2 \rightarrow In a symmetrical BIBD the no. of treatment common between any two block is λ . OR

In a symmetrical design every block different from the first block has exactly λ treatments in common with the first block.

Proof \rightarrow let a_i be the no. of treatments common to the first and i th ~~block~~ block and $i=1, 2, \dots, b$. Since, each of the k treatment of the 1st block occurs $(r-1)$ times in the other block. Then

$$\sum_{i=2}^b a_i = k(r-1) \quad \text{--- (1)}$$

Again, Since each of the $\binom{k}{2}$ pairs of treatments of the first block occurs $(\lambda-1)$ times in the remaining blocks.

Hence, $\sum_{i=2}^b \binom{a_i}{2} p a_i^2 = (\lambda - 1) \binom{k}{2}$

$$\Rightarrow \sum_{i=2}^b a_i (a_i - 1) = (\lambda - 1) k (k - 1)$$

$$\Rightarrow \sum_{i=2}^b a_i^2 = (\lambda - 1) k (k - 1) + \sum_{i=2}^b a_i \quad \text{--- (2)}$$

from eqⁿ (1) $(\sum a_i)$ is

$$\sum_{i=2}^b a_i^2 = (\lambda - 1) k (k - 1) + k(b - 1) \quad \text{--- (3)}$$

let $\sum_{i=2}^b (a_i - \lambda)^2 = \sum_{i=2}^b (a_i^2 + \lambda^2 - 2a_i\lambda)$

$$= \sum_{i=2}^b a_i^2 + \sum_{i=2}^b \lambda^2 - 2 \sum_{i=2}^b a_i \lambda$$

$$= (\lambda - 1) k (k - 1) + k(b - 1) + (b - 1)\lambda^2 - 2k(\lambda - 1)\lambda$$

If, $\lambda = k, b = u$ then

$$= (\lambda - 1) k (k - 1) + k(k - 1) + (u - 1)\lambda^2 - 2k(\lambda - 1)\lambda$$

$$= k(k - 1) [\lambda - \lambda + \lambda - 2\lambda] + (u - 1)\lambda^2$$

$$= k(k - 1) (-\lambda) + (u - 1)\lambda^2$$

$$= -\lambda k(k - 1) + \lambda^2(u - 1)$$

If $\lambda(u - 1) = \lambda(k - 1)$ then and $\lambda = k$

$$= -\lambda k(k - 1) + \lambda \cdot k(k - 1)$$

$$= 0$$

$$\Rightarrow \sum_{i=2}^b (a_i - \lambda)^2 = 0$$

$$\Rightarrow \boxed{a_i = \lambda}$$

Proved

Resolvable BIBD \longrightarrow

A BIBD with parameters v, b, r, λ, k is said to be resolvable BIBD if b blocks can be divided into n groups for the sets of $\frac{b}{n}$ blocks each. $\frac{b}{n}$ is an integer such that $\frac{b}{n}$ blocks forming any of the sets gives a complete replication of all the v treatments.

Example \rightarrow let us consider a BIBD with parameters $v=4, b=6, r=3, k=2, \lambda=1$

1	1	2	} I st group
2	3	4	
3	1	3	} II nd group
4	2	4	
5	1	4	} III rd group
6	2	3	

Thm. 9.10

Theorem \rightarrow Base Inequality

For a resolvable BIBD prove that $b \geq v + r - 1$

Proof \rightarrow Suppose, $n = \frac{b}{r}$

$$\Rightarrow b = nr$$

We know that
 $b \cdot k = 3 \cdot 2$

Therefore,

$$n \cdot k = 3 \cdot 2$$

$$\Rightarrow v = nk \quad \text{--- (1)}$$

Again, consider

$$\lambda(v-1) = s(k-1)$$

$$\Rightarrow \lambda(nk-1) = s(k-1) \quad \left\{ \begin{array}{l} \text{from eq. (1)} \\ \text{from eq. (1)} \end{array} \right.$$

$$\Rightarrow s = \frac{\lambda(nk-1)}{k-1}$$

$$\Rightarrow s = \lambda n + \frac{\lambda(n-1)}{(k-1)}$$

$$\Rightarrow s - \lambda n = \frac{\lambda(n-1)}{(k-1)} \quad \text{--- (2)}$$

Since, s is an integer, λ and n are also integers.
Therefore, from eq. (2) we conclude that $\frac{\lambda(n-1)}{(k-1)}$
must be an integer.

Let us consider,

$$b \leq v + s - 1 \quad \text{--- (3)}$$

$$\Rightarrow b - s < v - 1$$

$$\Rightarrow nk - s < v - 1 \quad \because b = nk$$

$$\Rightarrow (n-1)s < v - 1$$

$$\Rightarrow \lambda s(n-1) < \lambda(v-1)$$

$$\Rightarrow \lambda s(n-1) < s(k-1) \quad \left\{ \begin{array}{l} \because \lambda(v-1) = s(k-1) \end{array} \right.$$

$$\Rightarrow \lambda(n-1) < (k-1)$$

$$\Rightarrow \frac{\lambda(n-1)}{(k-1)} < 1$$

which is a contradiction of the fact that
 $\frac{\lambda(n-1)}{(k-1)}$ is an integer.

Hence, our assumption is wrong and we must have,

$$n \geq v + r - 1$$

Proved

Statistical Analysis for BIBD (Intra Block Analysis) (recovery) \rightarrow

This analysis is known as intra block analysis because it is carried out without recovery of interblock information. Let

y_{ij} be the response of j^{th} block receiving i^{th} treatment if it occurs. Here, we consider the model

$$y_{ij} = \mu + \alpha_i + \beta_j + e_{ij} \quad \text{--- (1)} \quad \begin{array}{l} i = 1, 2, \dots, t \\ j = 1, 2, \dots, b \end{array}$$

where, μ = general mean effect

α_i = additional effect due to i^{th} treatment

β_j = additional effect due to j^{th} block

e_{ij} = intrablock error. assume to be i.i.d.

d.e. $e_{ij} \sim N(0, \sigma^2)$

$$\sum_i \alpha_i = 0, \quad \sum_j \beta_j = 0$$

The least square estimates of μ , α_i and β_j can be obtained by minimizing residual sum of squares

$$S = \sum_i \sum_j n_{ij} (y_{ij} - \mu - \alpha_i - \beta_j)^2 \quad \text{--- (2)}$$

where, $n_{ij} = \begin{cases} 1 & ; \text{ when } i^{\text{th}} \text{ treatment occurs in } j^{\text{th}} \text{ block} \\ 0 & ; \text{ otherwise} \end{cases}$

$$T_i = n\hat{\mu} + g\alpha_i + \sum_j m_{ij}\hat{\beta}_j \quad \text{--- (9)}$$

$$B_j = K\hat{\mu} + \sum_i m_{ij}\alpha_i + k\hat{\beta}_j \quad \text{--- (10)}$$

Let us consider a quantity

$$Q_i = \left(T_i - \sum_j \frac{m_{ij}\hat{\beta}_j}{k} \right)$$

On putting the values of T_i and B_j

$$= \cancel{n\hat{\mu}} + g\alpha_i + \sum_j \cancel{m_{ij}\hat{\beta}_j} - \sum_j \cancel{m_{ij}\hat{\mu}} - \sum_j m_{ij} \left(\frac{\sum_i m_{ij}\alpha_i}{k} \right) - \sum_j \cancel{m_{ij}\hat{\beta}_j}$$

$$= g\alpha_i - \frac{1}{k} \sum_j m_{ij} [m_{1j}\alpha_1 + m_{2j}\alpha_2 + \dots + m_{ij}\alpha_i + \dots + \alpha_u \lambda]$$

$$= g\alpha_i - \frac{1}{k} [\alpha_1 \lambda + \alpha_2 \lambda + \dots + \alpha_i \lambda + \dots + \alpha_u \lambda]$$

$$= g\alpha_i - \frac{\lambda}{k} \sum_{i=1}^u \alpha_i - \frac{g\alpha_i}{k} + \frac{\lambda\alpha_i}{k}$$

$$= g\alpha_i - \frac{g\alpha_i}{k} + \frac{\lambda\alpha_i}{k}$$

$$= \frac{1}{k} [gk - g + \lambda] \alpha_i$$

$$= \frac{1}{k} [\lambda(k-1) + \lambda] \alpha_i$$

$$= \frac{1}{k} [\lambda(k-1+1)] \alpha_i$$

$$Q_i = \frac{\lambda}{k} \alpha_i \quad \text{--- (11)}$$

The quantity Q_i is called adjusted treatment total.

Now, suppose we want to test

$$H_0: \alpha_i = 0 \quad \forall i = 1, 2, \dots, u$$

for this purpose we first calculate sum of squares

due to regression when all the parameters are taken into consideration.

$$\text{Thus, } R(\hat{\mu}, \hat{\alpha}_i, \hat{\beta}_j) = \hat{\mu}^2 (1 + \sum_i \hat{\alpha}_i T_i) + \sum_j \hat{\beta}_j B_j$$

$$= \hat{\mu}^2 (1 + \sum_i \hat{\alpha}_i T_i) + \sum_j \hat{\beta}_j \left[\frac{B_j}{K} - \hat{\mu} - \frac{1}{K} \sum_i n_{ij} \hat{\alpha}_i \right]$$

from eqⁿ (10)

$$= \hat{\mu}^2 (1 + \sum_i \hat{\alpha}_i T_i) + \sum_j \frac{\hat{\beta}_j^2}{K} - \hat{\mu} \sum_j \hat{\beta}_j - \frac{1}{K} \sum_j \hat{\beta}_j \sum_i n_{ij} \hat{\alpha}_i$$

$$= \sum_i \hat{\alpha}_i T_i + \sum_j \frac{\hat{\beta}_j^2}{K} - \frac{1}{K} \sum_i \hat{\alpha}_i \sum_j n_{ij} \hat{\beta}_j$$

$$= \sum_j \frac{\hat{\beta}_j^2}{K} + \sum_i \hat{\alpha}_i \left[T_i - \frac{\sum_j n_{ij} \hat{\beta}_j}{K} \right]$$

$$= \sum_j \frac{\hat{\beta}_j^2}{K} + \sum_i \hat{\alpha}_i Q_i \quad \left[\hat{\alpha}_i = \frac{K Q_i}{\lambda \alpha} \right]$$

$$= \sum_j \frac{\hat{\beta}_j^2}{K} + \sum_i \frac{K}{\lambda \alpha} Q_i^2 \quad \text{--- (12)}$$

Sum of square due to regression under null hypothesis i.e. when only two parameters are taken into consideration

$$R(\mu, \beta_j) = \hat{\mu}^2 (1 + \sum_j \hat{\beta}_j B_j)$$

$$= \hat{\mu}^2 (1 + \sum_j \left[\frac{B_j}{K} - \hat{\mu} \right] B_j)$$

From eqⁿ (10) on putting $\hat{\alpha}_i = 0$; $\hat{\mu}^2$

$$= \sum_j \frac{\hat{\beta}_j^2}{K} \quad \text{--- (13)}$$

Thus, using eqⁿs (12) & (13) the adjusted treatment sum of squares is given by

$$R(\mu, \hat{\alpha}_i, \hat{\beta}_j) - R(\mu, \beta_j) = \frac{K}{\lambda \alpha} \sum_i Q_i^2$$

The sum of squares due to blocks ignoring treatments i.e. unadjusted block sum of squares is given by

$$\sum_j \frac{B_j^2}{k} - \frac{G^2}{9v} \quad \left[\frac{G^2}{9v} = \text{correction factor} \right]$$

Thus the ANOVA table for intra block analysis of BIBD is given by

ANOVA TABLE

Source of Variation	Degree of freedom	Sum of Squares
Blocks (unadjusted)	$(b-1)$	$\sum_j \frac{B_j^2}{k} - \frac{G^2}{9v}$
Treatments (adjusted)	$(v-1)$	$\frac{k}{\lambda v} \sum \theta_i^2$
Intra block error	$bk - b - v + 1$	by difference
Total	$bk - 1$	$\sum_{i,j} (m_{ij} \theta_{ij}^2) - \frac{G^2}{9v}$

Graeco - Latin Square \rightarrow

This is another name for pair of orthogonal Latin Squares superimposed on one another. Here one square is represented by Greek letters and other by Latin letters. In this arrangement Greek (or Latin) letters occurs once

in each row and once in each column, and once with each latin letter and once with each greek letter.

A $m \times m$ Graeco latin square is an incomplete four-way layout with all the four factors at the same level m and observations are taken on only m^2 of the possible m^4 of treatment combination.

For Ex \rightarrow

$$\begin{bmatrix} A & B & C \\ B & C & A \\ C & A & B \end{bmatrix}_{3 \times 3} \cdot \begin{bmatrix} \gamma & \beta & \alpha \\ \alpha & \gamma & \beta \\ \beta & \alpha & \gamma \end{bmatrix}$$

$$\begin{bmatrix} A\gamma & B\beta & C\alpha \\ B\alpha & C\gamma & A\beta \\ C\beta & A\alpha & B\gamma \end{bmatrix}_{3 \times 3}$$

In Graeco-latin square the anova table will have five components rows, columns, latin letters, Greek letters and error. The degrees of freedom for each of the first four components will be $(m-1)$ and ~~four~~ ^{for error} will be $(m-1)(m-3)$.

Remark: \rightarrow A field can consist of either finite or infinite number of elements. A field with infinite number of elements is called infinite field. If the number of elements i.e. marks

If field is finite then the field is called finite or Galois field.

Finite Field and the Design of Experiment

The properties of the finite field or Galois field makes construction and analysis of several types of design very convenient.

These properties are as follows :-

(1) A rule is made that any positive number is equal to the remainder R when N is divided by any integer number p and R is written as

$$R = N \text{ mod } p$$

(2) If p is a prime number then all the four operations addition, subtraction, multiplication and division of the elements are possible.

(3) when any element of prime modulo is multiplied in turn by all its non-zero elements each time a different product is obtained. This ensures that all possible divisions occurs. This is not true when p is not prime. All divisions are also not possible in this case, when division is possible the elements are said to form a Galois field.

(ii) There is at least one element in every field, different powers of which give the different non-zero elements of the field, such an element is called the primitive root of the field.

Ex: \rightarrow when $p = 7$
the primitive root is 3 because
 $3^0 = 1, 3^1 = 3, 3^2 = 2, 3^3 = 6, 3^4 = 4, 3^5 = 5, 3^6 = 1$

$$\text{Thus, } x^{p-1} = 1$$

where, x is any element of the field.

Though the above device ensures that an infinite sets of numbers could be reduced to only p elements, they can be manipulated just like the infinite set of numbers. However, it is not true when p is not prime. But, when p is a power of prime then such treatments are again possible by defining the elements in the following modified ways.

(i) let the number of elements be $S = p^m$ (S is power of prime) where p is a prime and m is integer.

let us now define S elements consisting of zero and $(S-1)$ polynomials up to $(m-1)$ degree and they can be written by using any symbol α (let). The coefficients in the polynomials are the elements ⁱⁿ the modulo p , with these elements as coefficients the total number of such polynomials is $(S-1)$.

has $\alpha \rightarrow S = 3^2$ then the elements of the modulo 3 has the coefficients $0, 1, 2, \alpha, \alpha+1, \alpha+2, 2\alpha, 2\alpha+1, 2\alpha+2$.

(ii) These elements can be added and subtracted, under modulo p but they cannot be multiplied and divided unless a device for reducing the polynomials is obtained. Such a device consist of choosing an irreducible polynomials of degree n or more is equated to the remainder, when it is divided by the irreducible polynomial. Such reduced polynomials are called minimum functions.

These functions are not easy to obtain. However, some of them which may be required are given below

Galois Field $GF(S)$	Minimum function
2^2	$\alpha^2 + \alpha + 1$
2^3	$\alpha^3 + \alpha + 1$
3^2	$\alpha^2 + \alpha + 2$

There may be more than one minimum function in a field

(iii) The ϕ non-zero elements can be obtained from different powers of α up to $\alpha^{p^n - 2}$

and α^{p^n-1} is always 1,

Ex: \rightarrow In the ~~the~~ 3^2 field with minimum function $(\alpha^2 + \alpha + 2)$ is

$$\alpha^0 = 1, \alpha^1 = \alpha, \alpha^2 = 2\alpha + 1$$

$$\alpha^3 = 2\alpha + 2, \alpha^4 = 2, \alpha^5 = 2\alpha, \alpha^6 = \alpha + 2, \alpha^7 = \alpha + 1, \alpha^8 = 1$$

Multiplication^{and} division become very easy by using such powers as elements.

Ex: $\rightarrow \alpha^3 = 2\alpha + 2, \alpha^6 = \alpha + 2$

$$\begin{aligned}(2\alpha + 2)(\alpha + 2) &= \alpha^3 \cdot \alpha^6 \\ &= \alpha^9 \\ &= \alpha^8 \cdot \alpha \\ &= 1 \cdot \alpha = \alpha\end{aligned}$$

Construction of Mutually Orthogonal Latin Squares (MOLS) and BIBD \rightarrow

construction of orthogonal series (OSS)

when S is a prime or power of a prime the following series of a design can be constructed GF(S)

$$v = S^2$$

$$b = (S+1)S$$

$$r = (S+1)$$

$$k = S$$

$$\lambda = 1$$

Ex: \rightarrow let us consider $S = 4 = 2^2$

$$v = S^2 = 16$$

$$b = 5 \times 4 = 20$$

$$r = 5$$

$$k = 4$$

$$\lambda = 1$$

Thus design with these parameters can be constructed as follows:

Remark - If we are given the prime or power of prime then we get those primitive roots

$3 \rightarrow 2$ $17 \rightarrow 3$
 $5 \rightarrow 2$ $19 \rightarrow 2$
 $7 \rightarrow 3$ $23 \rightarrow 2$
 $11 \rightarrow 2$
 $13 \rightarrow 2$

Step-1 \rightarrow Construct complete set of MOLES of order 4. Since, $5 = 4 = 2^2$ is a power of prime therefore we have $4-1 = 3$ Latin Squares in complete sets of MOLES.

First we construct three Latin Squares for this we write the elements of Galois field as $\{0, 1, \alpha, 1+\alpha\}$.

Now, we follow the following addition modulo rule

	0	1	α	$1+\alpha$
0	0	1	α	$1+\alpha$
1	1	0	$1+\alpha$	α
α	α	$1+\alpha$	0	1
$1+\alpha$	$1+\alpha$	α	1	0

let us assume $0 \rightarrow A, 1 \rightarrow B, \alpha \rightarrow C, 1+\alpha \rightarrow D$

then we get the Latin Square

(1)

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

1	ABCD	ABCD
2	COAB	COAB
3	BADC	DCBA
4	DCBA	BADC

are the cyclic rotation

To get the complete set of ~~rows~~ MOLES we fixed the first row and give the cyclic rotation to the other rows. Therefore, the complete set of

(2)	α	β	γ	δ	(1)
	γ	δ	α	β	2
	δ	γ	β	α	3
	β	α	δ	γ	(2)

(3)	P	Q	R	S	1
	S	R	Q	P	4
	Q	P	S	R	2
	R	S	P	Q	3

Step-2 \rightarrow To write down the design, we write rows as blocks.

Step-3 \rightarrow Then columns as blocks

Step-4 \rightarrow By superimposing (1) on (4) we write first row with those numbers coming along with first letter, second row with those numbers coming along with second letter and so on.

Step-5 \rightarrow Same as Step-4 by superimposing (2) on (4).

(4)

1	1	2	3	4
2	5	6	7	8
3	9	10	11	12
4	13	14	15	16

step - 2

5	1	5	9	13
6	2	6	10	14
7	3	7	11	15
8	4	8	12	16

step - 3

9	1	6	11	16
10	2	5	12	15
11	3	8	9	14
12	4	7	10	13

→ A position of A on OSI (1)

step - 4

13	1	7	12	14
14	2	8	11	13
15	3	5	10	16
16	4	6	9	15

(10) = (4)

(5)

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

(6)

comp

Construction of Orthogonal Series 2 (OS₂)

This is also equivalent to BIBD when S is a power of prime. The following series of design can be constructed

$$v = b = s^2 + s + 1$$

$$k = r = s + 1$$

$$\lambda = 1$$

This series can also be constructed with the help of complete set of HOLS. To construct OS₂ we need of OS₁ which we describe in the following example: - let us suppose

$$s = 4 = 2^2$$

$$v = b = 4^2 + 4 + 1 = 21$$

$$k = r = 4 + 1 = 5$$

$$\lambda = 1$$

Step-1: write down OS₁.

Step-2: Here, number of columns $k = 5$ whereas in OS₁ $k = 4$ also number of treatments in OS₂ is 21 whereas in OS₁ $v = 16$. Therefore, we have to add five more treatments namely 17, 18, 19, 20, 21. Now, in the new fifth column we add treatment 17 four times, treatment 18 four times, - - - treatment 21 four times such that we have $k = 5$ and $v = 21$.

Step-3: In OS₂ $b = 21$ but in OS₁ $b = 20$. Therefore, to make $b = 21$ we add a block of 5 new treatments 17, 18, 19, 20, 21 such that $b = 21$ and we can check that $r = 5$ and $\lambda = 1$

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

Construction of BIBD through Difference

Method \rightarrow

Let $\alpha_0, \alpha_1, \dots, \alpha_{k-1}$ be the elements of the Galois field $GF(s)$ of order s where s is the power or power of prime. If we find k elements out of s which are $\alpha_{i_1}, \alpha_{i_2}, \dots, \alpha_{i_k}$ such that $k(k-1)$ differences $\{(\alpha_{i_j} - \alpha_{i_m}), j \neq m\}$ are unknown null elements of the Galois field each occurring exactly λ .

$$\lambda = \frac{k(k-1)}{s-1}$$

Then, we can construct a symmetrical BIBD with parameters $v = b = s$ and $r = k$ and

$$\lambda = \frac{k(k-1)}{s-1}$$

The blocks can be obtained by adding to the initial block the non-null elements of $GF(s)$ in succession.

Remark \rightarrow Here, number of differences will be $kC_2 = \frac{k(k-1)}{2}$ but we can take both side differences therefore, no. of differences will be $k(k-1)$.

Example \rightarrow let $k=3, v=7$ then

The primitive root of 7 is 3. Then, we know that first row of BIBD will be $\{3^0, 3^1, 3^2\}$

$$= 1, 2, 4$$

\equiv under modulo 7

Here, it is notable that we take always even power in increasing order (ascending order) of primitive root. Now, the differences among elements are as 1, 2, 4 of GF(7).

$$r - 2 = -1 = 6 \quad \text{under mod 7}$$

$$1 - 4 = -3 = 4$$

$$\checkmark 2 - 1 = 1$$

$$2 - 4 = -2 = 5$$

$$4 - 1 = 3$$

$$4 - 2 = 2$$

$$7 - 1 = 6$$

$$7 - 3 = 4$$

$$7 - 2 = 5$$

All the elements are non-null elements of GF(7) and each occurs only once because

$$\text{here, } \lambda = \frac{k(k-1)}{s-1} = \frac{3 \times 2}{6-1} = 1$$

Hence, the BIBD will be

1	2	4
2	3	5
3	4	6
4	5	0
5	6	1
6	0	2
0	1	3

Ques 4 Construct a BIBD with $v=16$, $k=5$ with method of difference.

Then the primitive root of 11 is 5. Therefore,
we know that first row of $B^{-1}C^T$ will be
 $5^0, 5^2, 5^4, 5^6, 5^8$

Complementary BIBD \rightarrow

A complementary BIBD can be obtained by a given BIBD by writing those treatments which have not appeared with a block. If the parameters of ^{original} BIBD are v, b, k, λ, r , then the parameters of the complementary BIBD will be

$$v' = v, b' = b, r' = b - r, k' = v - k, \lambda' = b - 2r + \lambda$$

Ex \rightarrow $k = 3, v = 7$

initial block

1	2	4	3	5	6	0
2	3	5	1	4	6	0
3	4	6	1	2	5	0
4	5	0	1	2	3	6
5	6	1	2	3	4	0
6	0	2	2	3	4	0
0	1	3	1	3	4	5
			2	4	5	6

Residual BIBD \rightarrow

Given a symmetrical BIBD we get another BIBD called residual BIBD by deleting the first block and deleting the treatments belonging to the initial block from the remaining $(b-1)$ blocks. The parameters of the residual BIBD

will be $v' = v - k, b' = b - 1, r' = r, k' = k - \lambda, \lambda' = \lambda$

3	5
3	6
5	0

5	6
6	5
6	3

Derived BIBD \rightarrow from a symmetrical BIBD we can get another BIBD called derived BIBD by ^(+ab) omitting the initial block and retaining only those treatments in the remaining $(b-1)$ block which have appeared in the initial block.

2
4
4
1
2
1

The parameters of the derived BIBD will be $k' = k, b' = b-1, r' = r-1, k' = \lambda, \lambda' = \lambda - 1$

Remark \rightarrow

The complementary BIBD exist for all BIBD but residual and derived BIBD exist only for symmetrical BIBD.

Association Scheme \rightarrow Given v treatments, a relation satisfying the following conditions is said to be an association scheme with m groups.

(1) Any two treatments α and β are either I, II, ... or m^{th} associates and their relationship is symmetrical. we denote $(\alpha, \beta) = i$ when α and β are i^{th} associate.

(2) Each treatment α has n_i i^{th} associates, the number n_i being independent of α .

(3) If $(\alpha, \beta) = i$, the number of treatments γ that satisfy simultaneously $(\alpha, \gamma) = j$ and $(\beta, \gamma) = j'$ is $p_{jj'}^i$, and this number is independent of α and β . Further,

$$p_{jj'}^i = p_{j'j}^i$$

The numbers

The number $v, n_i, p_{jj'}^i$ are called parameters of the association scheme

Partially Balanced Incomplete Block

Design (PBIBD) \rightarrow

A PBIBD with m associate groups is an arrangement of v treatments in b blocks of size k ($k < v$) such that

(1) Every treatment occurs at most once in a set.

(2) Every treatment is replicated g_i times.

(3) Two treatments α and β replicated $h_{i\alpha\beta}$ times. If $(\alpha, \beta) = i$ and $h_{i\alpha\beta}$ is independent of treatments α and β .

$v, b, g_i, k, h_{i\alpha\beta}$ are called parameters of R.B.I.B.D.