

ORIGIN = 1

Nested Two-way ANOVA Model for Balanced Data

Nested ANOVA designs occur when, either by efficiency of design or by necessity, not all combinations of factors are studied. Nested design in a two-way ANOVA (involving two factors) permits analysis of the independent factor (factor A below) in the same way as a standard (crossed) two-way ANOVA. The difference lies in factor B nested within A. Here effects of different cell blocks within single treatment levels of A are assessed much as if in single factor ANOVA.

Data Structure:

Data are structured such that different levels of Nested Factor B occur only within single levels of independent Factor A. Other possible combinations are not studied.

Let index i, j indicate the i th row (treatment classes of Variable A) and j th column (treatment classes of Variable B)

Treatment Classes of Variable A:	Nested Two-Way ANOVA					
	Treatment Classes of Variable B Nested Within A:					
	#1	#2	#3	#4	#5	#6
#1	n	n				
#2			n	n		
#3					n	n
Each cell consists of n replicates with means \bar{Y}_{ij}						

Also let: $\bar{Y}_{.i}$ = mean over all columns for A_i .
 \bar{Y}_{ij} = mean within data blocks.
 $\bar{Y}_{..}$ = overall mean.

Model:

$$Y_{ijk} = \mu + \alpha_i + \beta_{j(i)} + \epsilon_{ijk}$$

where:

Note: this model has no associated test for interactions because of the nested design.

Restrictions:

$$\sum_i \alpha_i = 0 \quad \sum_j \beta_{j \text{ in } i} = 0$$

μ is a constant = grand mean of all objects.
 α_i are effect coefficients for classes i in Variable A.
 β_j are nested coefficients for classes j of Variable B in A.
 ϵ_{ijk} is the error term specific to each object i, j, k
 $i = 1$ to a , $j = 1$ to b , $k = 1$ to n , $N = abn$

Assumptions:

- ϵ_{ijk} are a random sample $\sim N(0, \sigma^2)$
- variance is homogeneous across cells

Variable Assignment:

ZAR := READPRN("c:/DATA/Biostatistics/ZarEX15.1aR.txt")

X := ZAR⁽²⁾ < response (dependent) variable Y

		B₁	B₂	
A₁	$Y_{11} :=$	$\begin{pmatrix} 102 \\ 104 \end{pmatrix}$	$Y_{12} :=$	$\begin{pmatrix} 103 \\ 104 \end{pmatrix}$
		B₃	B₄	
A₂	$Y_{21} :=$	$\begin{pmatrix} 108 \\ 110 \end{pmatrix}$	$Y_{22} :=$	$\begin{pmatrix} 109 \\ 108 \end{pmatrix}$
		B₅	B₆	
A₃	$Y_{31} :=$	$\begin{pmatrix} 104 \\ 106 \end{pmatrix}$	$Y_{32} :=$	$\begin{pmatrix} 105 \\ 107 \end{pmatrix}$

< response blocks

a := 3 b := 2 n := 2 N := 12

Zar Example 15.1

	conc	drug	source
1	102	1	A
2	104	1	A
3	103	1	Q
4	104	1	Q
5	108	2	D
6	110	2	D
7	109	2	B
8	108	2	B
9	104	3	L
10	106	3	L
11	105	3	S
12	107	3	S

Means:

$GM := \text{mean}(X)$

$GM = 105.8333$ < **grand mean**

$$B := \begin{pmatrix} \text{mean}(Y_{11}) & \text{mean}(Y_{12}) \\ \text{mean}(Y_{21}) & \text{mean}(Y_{22}) \\ \text{mean}(Y_{31}) & \text{mean}(Y_{32}) \end{pmatrix}$$

$$B = \begin{pmatrix} 103 & 103.5 \\ 109 & 108.5 \\ 105 & 106 \end{pmatrix}$$
 < **block means**

$$A := \begin{pmatrix} \text{mean}(Y_{11}, Y_{12}) \\ \text{mean}(Y_{21}, Y_{22}) \\ \text{mean}(Y_{31}, Y_{32}) \end{pmatrix}$$

$$A = \begin{pmatrix} 103.25 \\ 108.75 \\ 105.5 \end{pmatrix}$$
 < **Independent Factor A means (row means)**

Sums of Squares:

$k := 1 \dots N$

$$SSTO := \sum_k (X_k - GM)^2$$

$SSTO = 71.6667$ < **total SS**

$i := 1 \dots a$

$$SSA := b \cdot n \cdot \sum_{i=1}^a (A_i - GM)^2$$

$SSA = 61.1667$ < **among groups SS**

$i := 1 \dots b$

$$SSB := n \cdot \sum_{i=1}^a \sum_{j=1}^b (B_{i,j} - GM)^2$$

$SSB = 62.6667$ < **among all subgroups SS**

$SSBinA := SSB - SSA$

$SSBinA = 1.5$ < **subgroups SS**

$SSE := SSTO - SSB$

$SSE = 9$ < **error SS**

ANOVA Table for Nested design:

Sum of Squares:

Degrees of Freedom:

Mean Squares:

$SSA = 61.1667$

$df_A := a - 1$

$df_A = 2$

$MSA := \frac{SSA}{df_A}$

$MSA = 30.5833$

$SSBinA = 1.5$

$df_{BinA} := a \cdot (b - 1)$

$df_{BinA} = 3$

$MSBinA := \frac{SSBinA}{df_{BinA}}$

$MSBinA = 0.5$

$SSE = 9$

$df_E := a \cdot b \cdot (n - 1)$

$df_E = 6$

$MSE := \frac{SSE}{df_E}$

$MSE = 1.5$

$SSTO = 71.6667$

$df_T := N - 1$

$df_T = 11$

F-Test for Independent Factor Effects:**Hypotheses:**

$$H_0: \alpha_i = 0 \text{ for all } i$$

$$H_1: \text{At least one } \alpha_i \neq 0$$

Test Statistic:

$$F := \frac{MSA}{MSE} \quad F = 20.3889$$

Sampling Distribution of the Test Statistic F:

If Assumptions hold and H_0 is true then $F \sim F_{(df_A, df_E)}$

Critical Value of the Test:

$$\alpha := 0.05 \quad < \text{Probability of Type I error must be explicitly set}$$

$$C := qF(1 - \alpha, df_A, df_E) \quad C = 5.1433$$

Decision Rule:

IF $F > C$, THEN REJECT H_0 OTHERWISE ACCEPT H_0

Probability Value:

$$P := 1 - pF(F, df_A, df_E) \quad P = 0.00211$$

F-Test for Nested Factor Specific Effects:**Hypotheses:**

$$H_0: \beta_{j \text{ in } i} = 0 \text{ for all values of } j \text{ within a specific level } i$$

$$H_1: \text{At least one } \beta_{j \text{ in } i} < 0$$

Test Statistic:

$$F := \frac{MS_{\text{BinA}}}{MSE} \quad F = 0.3333$$

Sampling Distribution of the Test Statistic F:

If Assumptions hold and H_0 is true then $F \sim F_{(df_{\text{BinA}}, df_E)}$

Critical Value of the Test:

$$\alpha := 0.05 \quad < \text{Probability of Type I error must be explicitly set}$$

$$C := qF(1 - \alpha, df_{\text{BinA}}, df_E) \quad C = 4.7571$$

Decision Rule:

IF $F > C$, THEN REJECT H_0 OTHERWISE ACCEPT H_0

Probability Value:

$$P := 1 - pF(F, df_{\text{BinA}}, df_E) \quad P = 0.802202$$

Prototype in R:

```
#BALANCED NESTED ANOVA
#FIXED EFFECTS (ANOVA MODEL I)
#ZAR EXAMPLE 15.1a
ZAR=read.table("c:/DATA/Biostatistics/ZarEX15.1aR.txt")
ZAR
attach(ZAR)
Y=conc
A=factor(drug)
B=factor(source)

anova(lm(Y~A+B%in%A))
```

Analysis of Variance Table

Response: Y

	Df	Sum Sq	Mean Sq	F value	Pr(>F)
A	2	61.167	30.583	20.3889	0.002110 **
A:B	3	1.500	0.500	0.3333	0.802202
Residuals	6	9.000	1.500		

Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

Confidence Intervals for Fixed Factor Means:

Critical Value:

$$C := \left| qt\left(\frac{\alpha}{2}, df_E\right) \right| \quad C = 2.4469$$

For Factor A:

$$CI_A := \left(A - C \cdot \sqrt{\frac{MSE}{b \cdot n}} \quad A + C \cdot \sqrt{\frac{MSE}{b \cdot n}} \right)$$

$$CI_A = \left[\begin{pmatrix} 101.7516 \\ 107.2516 \\ 104.0016 \end{pmatrix} \quad \begin{pmatrix} 104.7484 \\ 110.2484 \\ 106.9984 \end{pmatrix} \right]$$

From calculations above:

$$A = \begin{pmatrix} 103.25 \\ 108.75 \\ 105.5 \end{pmatrix} < \text{factor A means}$$

$$a = 3$$

$$b = 2$$