

## The Design and Analysis of Experiments

### Completely Randomised Design (CRD)

*Aim:* To find if there is a significant difference between the population means of  $c$  “treatments” ( $c \geq 3$ ).

*Data Pattern:*  $n$  observations are taken on each of  $c$  treatments and the resulting data can be tabulated as follows:

	Population mean	Observations	Total	Sample mean
Treatment 1	$\mu_1$	$x_{11}, x_{12}, \dots, x_{1n}$	$T_1$	$\bar{x}_1$
Treatment 2	$\mu_2$	$x_{21}, x_{22}, \dots, x_{2n}$	$T_2$	$\bar{x}_2$
$\vdots$	$\vdots$	$\vdots$	$\vdots$	$\vdots$
Treatment $c$	$\mu_c$	$x_{c1}, x_{c2}, \dots, x_{cn}$	$T_c$	$\bar{x}_c$

where

$$T_i = \sum_{j=1}^n x_{ij}, \quad \bar{x}_i = \frac{T_i}{n}.$$

*Analysis of Variance (ANOVA):* Compute

$$T = \sum_{j=1}^c T_j,$$

$$SS_{total} = \sum_{i=1}^c \sum_{j=1}^n x_{ij}^2 - \frac{T^2}{cn},$$

$$SS_{treat} = \frac{1}{n} \sum_{i=1}^c T_i^2 - \frac{T^2}{cn},$$

$$SS_{error} = SS_{total} - SS_{treat},$$

$$MS_{treat} = \frac{SS_{treat}}{c - 1},$$

$$MS_{error} = \frac{SS_{error}}{c(n - 1)}.$$

Arrange them as the one-way ANOVA table:

Source of variation	Sum of squares	d.f.	Mean square	$F$ -ratio
Treatments	$SS_{treat}$	$c - 1$	$MS_{treat}$	$\frac{MS_{treat}}{MS_{error}}$
Error	$SS_{error}$	$c(n - 1)$	$MS_{error}$	
Total variation	$SS_{total}$	$cn - 1$		

*Hypothesis Test:*

$H_0: \mu_1 = \mu_2 = \dots = \mu_c$  (i.e. treatment means are the same).

$H_1: \mu_i$  differ (i.e. treatment means are different).

Given the significance level  $\alpha$ , find  $F_{\alpha, c-1, c(n-1)}$  from the  $F$ -distribution table with  $\nu_1 = c - 1$  and  $\nu_2 = c(n - 1)$ . If

$$F = \frac{MS_{treat}}{MS_{error}} > F_{\alpha, c-1, c(n-1)}$$

reject  $H_0$  and conclude that the treatment means are significantly different at  $\alpha$  level of significance; otherwise do not reject  $H_0$  and conclude that the treatment means are not significantly different at  $\alpha$  level of significance.

*Confidence Intervals:* Compute

$$s = \sqrt{MS_{error}}.$$

The  $(1 - \alpha)100\%$  confidence interval for  $\mu_i$ , the population mean of treatment  $i$  is given by

$$\bar{x}_i \pm t_{\alpha/2, c(n-1)} s \sqrt{\frac{1}{n}}.$$

The  $(1 - \alpha)100\%$  confidence interval for  $\mu_i - \mu_j$ , the difference between the population means of treatments  $i$  and  $j$  is given by

$$\bar{x}_i - \bar{x}_j \pm t_{\alpha/2, c(n-1)} s \sqrt{\frac{2}{n}}.$$

### **Randomised Block Design (RBD)**

*Aim:* To find (i) if there is a significant difference between the means of  $c$  “treatments”, and (ii) if there is a significant difference between the means of  $r$  “blocks”.

*Data Pattern:*

	Treat 1	...	Treat $c$	Block total	Block mean
Block 1	$x_{11}$	...	$x_{1c}$	$B_1$	$\bar{B}_1$
Block 2	$x_{21}$	...	$x_{2c}$	$B_2$	$\bar{B}_2$
...	...		...	...	...
Block $r$	$x_{r1}$	...	$x_{rc}$	$B_r$	$\bar{B}_r$
Treat total	$T_1$	...	$T_c$		
Treat mean	$\bar{T}_1$	...	$\bar{T}_c$		

where

$$B_i = \sum_{j=1}^c x_{ij}, \quad \bar{B}_i = \frac{B_i}{c},$$

$$T_j = \sum_{i=1}^r x_{ij}, \quad \bar{x}_j = \frac{T_j}{r}.$$

Analysis of Variance (ANOVA): Compute

$$G = \sum_{i=1}^c T_i, \quad \bar{G} = \frac{G}{rc},$$

$$SS_{total} = \sum_{i=1}^r \sum_{j=1}^c x_{ij}^2 - \frac{G^2}{rc},$$

$$SS_{treat} = \frac{1}{r} \sum_{j=1}^c T_j^2 - \frac{G^2}{rc},$$

$$SS_{block} = \frac{1}{c} \sum_{i=1}^r B_i^2 - \frac{G^2}{rc},$$

$$SS_{error} = SS_{total} - SS_{treat} - SS_{block},$$

$$MS_{treat} = \frac{SS_{treat}}{c-1},$$

$$MS_{block} = \frac{SS_{block}}{r-1},$$

$$MS_{error} = \frac{SS_{error}}{(r-1)(c-1)}.$$

Arrange them as the ANOVA table:

Source of variation	Sum of squares	d.f.	Mean square	F-ratio
Treatments	$SS_{treat}$	$c-1$	$MS_{treat}$	$\frac{MS_{treat}}{MS_{error}}$
Blocks	$SS_{block}$	$r-1$	$MS_{block}$	$\frac{MS_{block}}{MS_{error}}$
Error	$SS_{error}$	$(r-1)(c-1)$	$MS_{error}$	
Total variation	$SS_{total}$	$rc-1$		

Hypothesis Tests:

$H_0$ : treatment means are the same.

$H_1$ : treatment means are different.

Given the significance level  $\alpha$ , find  $F_{\alpha, c-1, (r-1)(c-1)}$  from the  $F$ -distribution table with  $\nu_1 = c-1$  and  $\nu_2 = (r-1)(c-1)$ . If

$$F = \frac{MS_{treat}}{MS_{error}} > F_{\alpha, c-1, (r-1)(c-1)}$$

reject  $H_0$  and conclude that the treatment means are significantly different at  $\alpha$  level of significance; otherwise do not reject  $H_0$  and conclude that the treatment means are not significantly different at  $\alpha$  level of significance.

$H_0$ : block means are the same.

$H_1$ : block means are different.

Given the significance level  $\alpha$ , find  $F_{\alpha, r-1, (r-1)(c-1)}$  from the  $F$ -distribution table with  $\nu_1 = r - 1$  and  $\nu_2 = (r - 1)(c - 1)$ . If

$$F = \frac{MS_{block}}{MS_{error}} > F_{\alpha, r-1, (r-1)(c-1)}$$

reject  $H_0$  and conclude that the block means are significantly different at  $\alpha$  level of significance; otherwise do not reject  $H_0$  and conclude that the block means are not significantly different at  $\alpha$  level of significance.

*Confidence Intervals:* Compute

$$s = \sqrt{MS_{error}}.$$

The  $(1 - \alpha)100\%$  confidence interval for  $\mu_j$ , the population mean of treatment  $j$ , is given by

$$\bar{T}_j \pm t_{\alpha/2, (r-1)(c-1)} s \sqrt{\frac{1}{r}}.$$

The  $(1 - \alpha)100\%$  confidence interval for  $\mu_j - \mu_k$ , the difference between the population means of treatments  $j$  and  $k$ , is given by

$$\bar{T}_j - \bar{T}_k \pm t_{\alpha/2, (r-1)(c-1)} s \sqrt{\frac{2}{r}}.$$