## Randomized Complete Block Design (RCBD)

 $y_{ij} = \mu + \tau_i + \beta_j + \epsilon_{ij}$ , Where i = # of treatments, j = # of blocks. ( $\mu$  Grand Mean), ( $\tau$  Treatment effect), ( $\beta$  Block effect). Constraint:  $\sum \tau_i = 0$ ,  $\sum \beta_j = 0$ .

Estimates: Grand Mean:  $\hat{\mu} = \bar{y}_{..}$ , Treatment Effect:  $\hat{\tau}_i = \bar{y}_{i.} - \bar{y}_{..}$ , Block Effect:  $\hat{\beta}_i = \bar{y}_{.i} - \bar{y}_{..}$ 

Grand Smaple Average:  $\bar{y}_{..} = \frac{1}{ab} \sum_{i=1}^{a} \sum_{j=1}^{b} y_{ij}$ 

Sample Mean for  $i^{th}$  Treatment (averaged over all blocks):  $\bar{y}_{i.} = \frac{1}{b} \sum_{j=1}^{b} y_{ij}$ Sample Mean for  $j^{th}$  Block (averaged over all treatments):  $\bar{y}_{.j} = \frac{1}{a} \sum_{i=1}^{a} y_{ij}$ 

Fitted  $y_{ij}$  Values:  $\hat{y}_{ij} = \bar{y}_{i.} + \bar{y}_{.j} - \bar{y}_{..}$ 

### RCBD - ANOVA

Source	df	SS	MS	F
Treatments	a-1	SSTr	$MSTr = \frac{SSTr}{a-1}$	$F = \frac{MSTr}{MSE}$
Blocks	b-1	SSB	$MSB = \frac{\widetilde{SSB}}{b-1}$	$F = \frac{\widetilde{M}\widetilde{S}\widetilde{B}}{MSE}$
Error	(a-1)(b-1)	SSE	$MSE = \frac{\mathring{S}S\mathring{E}}{(a-1)(b-1)}$	
Total	ab-1	SST		

Pooled sample variance(est.  $\sigma^2$ ):  $s^2 = MSE$ . Test of Homogeneity of Treatment Effects:

> $H_0$ :  $\tau_1 = \tau_2 = \dots = \tau_a = 0$ .  $H_a$ : not all  $\tau_i = 0$ .

Reject  $H_0$  if  $F > F_{\alpha,a-1,(a-1)(b-1)}$ , or p-value  $< \alpha$ .

C.I for  $\mu_i$ :  $\bar{y}_{i}$ .  $\pm t_{\alpha/2,df_{error}} \frac{s}{\sqrt{h}}$ 

Fisher's LSD Multiple Comparison Test: Treatment i and j are significantly different if  $|\bar{y}_i - \bar{y}_j| > t_{\alpha/2, df_{error}} s_1 / \frac{2}{h}$ 

(Reject  $H_0: \tau_i = \tau_j$ ). **Fishers C.I.:**For all possible treatment differences,  $\bar{y}_i - \bar{y}_k \pm t_{\alpha/2, df_{error}} s \sqrt{\frac{2}{b}}$ .

**Tukey's Multiple Comparison Test:** Treatment i and j are significantly different if  $|\bar{y}_i - \bar{y}_j| > q_{\alpha,a,df_{error}} \frac{s}{\sqrt{\lambda}}$ .

**Tukey's C.I.:** For all possible treatment differences,  $\bar{y}_i - \bar{y}_k \pm q_{\alpha,a,df_{error}} \frac{s}{\sqrt{h}}$ .

## Two Factor Design

 $y_{ijk} = \mu + \tau_i + \beta_j + (\tau \beta)_{ij} + \epsilon_{ijk}$ . Constraint:  $\sum \tau_i = 0$ ,  $\sum \beta_j = 0$ , and  $\sum (\tau \beta)_{ij} = 0$  (for each fixed i,j). **Interaction:** if  $(\tau \beta)_{ij} = 0$  for every i,j then there is no interaction between factors A and B.

Estimates:  $\bar{y}_{...} = \frac{1}{abn} \sum_{i=1}^{a} \sum_{j=1}^{b} \sum_{k=1}^{n} y_{ijk}$ .  $\bar{y}_{i..} = \frac{1}{bn} \sum_{j=1}^{b} \sum_{k=1}^{n} y_{ijk}$ .  $\bar{y}_{.j} = \frac{1}{an} \sum_{i=1}^{a} \sum_{k=1}^{n} y_{ijk}$ .

Fitted Values:  $\hat{y}_{ijk} = \hat{\mu} + \hat{\tau}_i + \hat{\beta}_j + (\hat{\tau\beta})_{ij} = \bar{y}_{ij}$ .

#### 2 Factor - ANOVA

Source	df	SS	MS	F	
Factor A	a-1	SSA	$MSA = \frac{SSA}{a-1}$	$F = \frac{MSA}{MSE}$	
Factor B	b-1	SSB	$MSB = \frac{\tilde{S}S\tilde{B}}{h-1}$	$F = \frac{\widetilde{MSB}}{\widetilde{MSE}}$	
AB Interaction	(a-1)(b-1)	SSAB	$MSAB = \frac{\ddot{S}S\ddot{A}B}{(a-1)(b-1)}$	$F = \frac{MSAB}{MSE}$	
Error	N-ab	SSE	$MSE = \frac{SSE}{N-ab}$		
Total	N-1	SST			

Note: N = abnIf n = 1 assume no interaction. Test of significance of AB Interaction should be done first. If there is interaction it is not usesful

to test for significance of A and B.

Test of Significance of AB Interaction:  $H_0$ :  $(\tau\beta)_{11} = (\tau\beta)_{12} = \cdots = (\tau\beta)_{ab} = 0$ .  $H_a$ : not all  $(\tau\beta)_{ij}$  are the same. Reject  $H_0$  if  $F > F_{\alpha,(a-1)(b-1),N-ab}$ , or p-value  $< \alpha$ .

If  $H_0$  is rejected, conclude that there exists interaction between Factors A and B at significance level  $\alpha$ .

#### Case 1: No Interaction between A and B

It is meaningful to test for the significance of the main effect of A and the main effect of B.

May combine the interaction sum of squares with the error sum of squares.

Anova: df:  $df_{error} = abn - a - b + 1$ , SSE: SSE' = SSAB + SSE, MSE:  $MSE' = \frac{SSE'}{abn - a - b + 1}$ .

Note that F values must be updated with the new MSE before testing.

**Tukeys C.I.:** For  $\tau_i - \tau_{i'}$  (or  $\beta$ ). Equivalent to FWE =  $\alpha$  for hypothesis test.

Factor A:  $\bar{y}_{i\cdots} - \bar{y}_{i^{\prime}\cdots} \pm q_{\alpha,a,df_{error}} \sqrt{\frac{MSE}{bn}}$ . Factor B:  $\bar{y}_{\cdot j\cdot} - \bar{y}_{\cdot j^{\prime}\cdot} \pm q_{\alpha,b,df_{error}} \sqrt{\frac{MSE}{an}}$ . Fisher's C.I.: For  $\tau_i - \tau_{i^{\prime}}$  (or  $\beta$ ). Equivalent to  $\alpha$  for hypothesis test.

Factor A:  $\bar{y}_{i\cdots} = t_{\alpha/2,df_{error}} \sqrt{\frac{2MSE}{bn}}$ . Factor B:  $\bar{y}_{\cdot j\cdot} = \bar{y}_{\cdot j\cdot} \pm t_{\alpha/2,df_{error}} \sqrt{\frac{2MSE}{an}}$ 

#### Case 2: Interaction between A and B

Not meaningful to test significance of main effect of A and B. Treat each individual factor-level combonation of  $i^{th}$  level of A, and  $j^{th}$  level of B, as a treatment. So there are ab treatments. Test every pair of treatments using Tukey's or Fisher's. **Test:**  $H_0$ :  $\mu_{11} = \mu_{12} = \cdots = \mu_{ab}$  using the F test for CRD.

Tukey's C.I.: 
$$\bar{y}_{ij} - \bar{y}_{i'j'} \pm q_{\alpha,ab,abn-ab} \sqrt{\frac{MSE}{n}}$$
  
Fisher's C.I.:  $\bar{y}_{ij} - \bar{y}_{i'j'} \pm t_{\alpha/2,abn-ab} \sqrt{\frac{2MSE}{n}}$ 

# $2^k$ Factorial Design Using the [-1, +1] Notation

#### 2 Factor - ANOVA

Std Order	A	В	AB	$\mathbf{C}$	AC	BC	ABC
(1)	_	_	+	_	+	+	_
a	+	_	_	_	_	+	+
b	_	+	_	_	+	_	+
ab	+	+	+	_	_	_	_
c	_	_	+	+	_	_	+
ac	+	—	_	+	+	_	_
bc	_	+	_	+	_	+	_
abc	+	+	+	+	+	+	+

Note:  $A \times B = AB$  and so on. Factor interactions: There will be k main effects and  $\binom{k}{2}$  two-factor interactions,  $\binom{k}{3}$  three-factor interactions. Contrasts:

Contrasts: Estimated main effect:  $\frac{1}{2^{k-1}} \cdot \sum_{i=1}^{2^k} c_i \bar{y}_i$ 

**Model:** Standard error:  $se(\hat{\beta}) = \sqrt{\frac{\sigma^2}{n2^k}} = \sqrt{\frac{MSE}{n2^k}}$ . Confidence interval:  $\hat{\beta} \pm t_{\alpha/2,df_{error}} \cdot se(\hat{\beta})$