



## University of Strathclyde

### 53.483 DATA ANALYSIS I

*Tuesday 22nd January 2008  
10.00a.m. - 12noon*

*All questions may be attempted.*

*Credit will be given for the best THREE answers only.*

- 1.(a) The mathematical model for the response variable  $X_{ij}$  in a completely randomised design with  $k$  treatments and  $n$  replications per treatment can be described by

$$X_{ij} = \mu + \tau_j + \varepsilon_{ij}, \quad 1 \leq i \leq n, \quad 1 \leq j \leq k,$$

where  $\mu$  is the grand mean,  $\tau_j$  the effect of treatment  $j$  and  $\varepsilon_{ij}$  the experimental error. Assume that treatments are fixed effects,  $\sum \tau_j = 0$ . Assume also that  $\varepsilon_{ij}$  are independent and follow a  $N(0, \sigma_e^2)$  distribution. Give the definitions for  $SS_{treat}$  and  $MS_{treat}$ . Show that

$$E(MS_{treat}) = \sigma_e^2 + n\sigma_T^2,$$

where  $\sigma_T^2 = (\tau_1^2 + \dots + \tau_k^2)/(k-1)$ . (You may use the following results without proof. Let  $Y_i$  ( $1 \leq i \leq m$ ) be independent and normally distributed  $N(0, \sigma^2)$  variables. Define  $\bar{Y} = (Y_1 + \dots + Y_m)/m$ . Then  $\bar{Y} \sim N(0, \sigma^2/m)$  and

$$E\left[\frac{1}{m-1} \sum_{i=1}^m (Y_i - \bar{Y})^2\right] = \sigma^2. )$$

(13 marks)

- (b) The partially completed ANOVA table for a  $p \times q$  design is shown below:

Source	SS	d.f.	MS	F
A	3.06		1.02	
B	159.50	3		
$A \times B$	11.79			
Error				
Total	180.44	31		

Assume A is a fixed effect but B is a random effect so this is a mixed model. Using a suitable notation, give a clear description of the model, including any assumptions necessary to undertake the analysis of variance. Fill in the blanks in the ANOVA table. Carry out the F-tests for interaction  $A \times B$  and main effects A and B. Test using  $\alpha = 0.05$ .

(12 marks)

2. The following data were generated from a randomised block design with 4 treatments and 4 blocks:

	Block 1	Block 2	Block 3	Block 4
Treat 1	1.25	1.43	1.32	1.31
Treat 2	2.05	1.56	1.68	1.69
Treat 3	1.95	2.00	1.83	1.81
Treat 4	1.75	1.93	1.70	1.59

Assume that treatments and blocks are fixed effects. The classical mathematical model for the design is

$$X_{ij} = \mu + \beta_i + \tau_j + \varepsilon_{ij}, \quad 1 \leq j \leq 4, \quad 1 \leq i \leq 4,$$

where  $\mu$  is the grand mean,  $\tau_j$  is the effect of treatment  $j$ ,  $\beta_i$  is the effect of block  $i$  while  $\varepsilon_{ij}$  is the experimental error. The GLIM mathematical model for the design is

$$X_{ij} = \tilde{\mu} + \tilde{\beta}_i + \tilde{\tau}_j + \varepsilon_{ij}, \quad 1 \leq j \leq 4, \quad 1 \leq i \leq 4.$$

(a) State the classical assumptions and the GLIM assumptions. State the relationships between the classical parameters and the GLIM parameters by expressing  $\tilde{\mu}$ ,  $\tilde{\beta}_i$  and  $\tilde{\tau}_j$  in terms of  $\mu$ ,  $\beta_i$  and  $\tau_j$ .

(3 marks)

(b) State the GLIM commands you would use to input the data in order to produce the following results:

(3 marks)

```
[i] ? $fit$  
[o] deviance - 0.95584  
[o] residual df = 15  
  
[i] ? $fit +block$  
[o] deviance = 0.89167 (change = -0.06417)  
[o] residual df = 9 (change = -3)  
[o]  
[i] ? $fit + treat$  
[o] deviance = 0.17296 (change = -0.7187)  
[o] residual df = 9 (change = -3)  
[o]  
[i] ? $ disp e s v r$
```

```

[o]      estimate      s.e.   parameter
[o] 1     1.399  0.09169      1
[o] 2    -0.02000 0.09802  BLOCK(2)
[o] 3    -0.1175 0.09802  BLOCK(3)
[o] 4    -0.1500 0.09802  BLOCK(4)
[o] 5    -0.4175 0.09802  TREAT(2)
[o] 6     0.5700 0.09802  TREAT(3)
[o] 7     0.4150 0.09802  TREAT(4)
[o] scale parameter 0.01922
[i] ? $tab the yield mean for treat$
[o] TREAT      1      2      3
[o] MEAN    1.327  1.745  1.898  1.742

```

(c) Based on the GLIM output, draw up the ANOVA table and state what conclusions can be drawn from it (tests using  $\alpha = 0.05$ ).

(5 marks)

(d) Carry out the Newman-Keuls MRT for the 4 treatments (test using  $\alpha = 0.05$ ) and state your conclusions.

(7 marks)

(e) Based on the GLIM output, the standard error (s.e.) for the GLIM parameter  $\tilde{\tau}_2$  is 0.09802. State (without proof) the least-squares estimator for the  $\tilde{\tau}_2$  in terms of the observations, and show its variance from first principles and hence verify its s.e. = 0.09802.

(7 marks)

3. Data are provided on content uniformity of film-coated tablets produced for a cardiovascular drug used to lower blood pressure. A random sample of 3 batches was chosen at each of two manufacturing sites, and from each batch 5 tablets were examined. We wish to know if there are differences between the two sites (fixed effects) and between batches within sites (random effects) i.e. mixed model.

		Site		
		1		2
		Batch		
		1	2	3
Rep1		5.03	4.64	5.10
		5.10	4.73	5.15
		5.25	4.82	5.20
		4.98	4.95	5.08
		5.05	5.06	5.14
		5.05	5.46	4.90
		4.96	5.15	4.95
		5.12	5.18	4.86
		5.12	5.18	4.86
		5.05	5.11	5.07

(a) State a linear model that may be used to represent the data and state the assumptions of the model.

(6 marks)

(b) State the GLIM commands you would use to input the data in order to produce the following results.

[i] ? \$fit\$

[o] deviance = 0.76247

[o] d.f. = 29

[o]

[i] ? \$fit + site\$

[o] deviance = 0.74421 (change = -0.01825)

[o] d.f. = 28 (change = -1)

[o]

[i] ? \$fit + site.batch\$

[o] deviance = 0.29020 (change = -0.4540)

[o] d.f. = 24 (change = -4)

(5 marks)

(c) Using the GLIM results provided, give the ANOVA table and undertake hypothesis tests to see if there are significant differences between the two sites and between batches within sites. Test using  $\alpha = 0.05$ .

(7 marks)

(d) Compute the 90% confidence interval for  $\sigma_e^2$  (the variance of the experimental error).

(7 marks)

4.(a) In a  $2^4$  design with treatment  $A, B, C, D$  and 8 blocks and 3 replicates, the design is arranged so that  $AB$ ,  $BC$  and  $CD$  are confounded with block effects in 3 replicates. Specify the units to go into each block and identify the composition of the ANOVA table.

(12 marks)

(b) In a  $2^4$ -design, 3 replications are carried out for each of the following half of the 16 treatment combinations:

(1)  $b \ c \ bc \ ad \ abd \ acd \ abcd$

Identify the defining contrast and indicate aliases.

(9 marks)

(c) Identify the composition of the ANOVA table for part (b).

(4 marks)

(1)

## Solutions to DA1 Exam 2007/8

$$1@ \quad T_j = \sum_{i=1}^n x_{ij}, \quad \bar{T}_j = \frac{T_j}{n}$$

$$G = \sum_{i=1}^n \sum_{j=1}^k x_{ij}, \quad \bar{G} = \frac{G}{nk}$$

$$SS_{\text{treat}} = n \sum_{j=1}^k (\bar{T}_j - \bar{G})^2$$

$$MS_{\text{treat}} = \frac{SS_{\text{treat}}}{k-1}$$

$$\text{Note } \bar{T}_j = \frac{1}{n} \sum_{i=1}^n (\mu + \tau_j + \epsilon_{ij})$$

$$= \mu + \tau_j + \bar{\epsilon}_j$$

$$\text{where } \bar{\epsilon}_j = \frac{1}{n} \sum_{i=1}^n \epsilon_{ij} \sim N(0, \frac{\sigma_e^2}{n})$$

$$\bar{G} = \frac{1}{nk} \sum_{i=1}^n \sum_{j=1}^k (\mu + \tau_j + \epsilon_{ij})$$

$$= \mu + \frac{1}{nk} \sum_{i=1}^n \sum_{j=1}^k \epsilon_{ij}$$

$$= \mu + \bar{\epsilon}$$

$$\text{where } \bar{\epsilon} = \frac{1}{k} \sum_{j=1}^k \left( \frac{1}{n} \sum_{i=1}^n \epsilon_{ij} \right) = \frac{1}{k} \sum_{j=1}^k \bar{\epsilon}_j \sim N(0, \frac{\sigma_e^2}{nk})$$

Compute

$$E(MS_{\text{treat}}) = \frac{n}{k-1} E \left[ \sum_{j=1}^k (\mu + \tau_j + \bar{\epsilon}_j - \mu - \bar{\epsilon})^2 \right]$$

$$= \frac{n}{k-1} E \left[ \sum_{j=1}^k (\tau_j^2 + 2\tau_j(\epsilon_j - \bar{\epsilon}) + (\epsilon_j - \bar{\epsilon})^2) \right]$$

$$\begin{aligned}
 &= \frac{n}{k-1} \left[ \sum_{j=1}^k \bar{\tau}_j^2 + \sum_{j=1}^k 2\bar{\tau}_j (Ee_j - E\bar{e}) + E \sum_{j=1}^k (e_j - \bar{e})^2 \right] \\
 &= n \frac{\bar{\tau}_1^2 + \dots + \bar{\tau}_k^2}{k-1} + 0 + n E \left[ \frac{1}{k-1} \sum_{j=1}^k (e_j - \bar{e})^2 \right] \\
 &= n \bar{\sigma}_T^2 + n \cdot \frac{\bar{\sigma}_e^2}{n} = n \bar{\sigma}_T^2 + \bar{\sigma}_e^2
 \end{aligned}$$

1(b) This is a  $p \times q$  design with  $p = 4$  (A) and  $q = 4$  (B)

$$X_{ijk} = \mu + \alpha_i + \beta_j + \gamma_{ij} + \epsilon_{ijk}$$

$$1 \leq i \leq 4, \quad 1 \leq j \leq 4, \quad 1 \leq k \leq 2$$

where  $\mu$  - grand mean

$\alpha_i$  - effect of factor A level  $i$

$\beta_j = \dots - - - - - 13 - - - j$

$\beta_{ij}$  - effect of interaction between factor A level  $i$  & factor B level  $j$

# Eijk - experimental error

$\therefore A$  is a fix effect while  $B$  is a random effect

This is a mixed-effects model

## Assumptions:

$\alpha_i$  is constants,  $\sum_{i=1}^n \alpha_i = 0$

$$\beta_j \sim N(0, \sigma_{\beta}^2)$$

$$\alpha \beta_{ij} \sim N(0, \sigma_{AB}^2)$$

$$E_{ijk} \sim N(0, \sigma_e^2)$$

ANOVA

<u>Source</u>	<u>SS</u>	<u>d.f.</u>	<u>ms</u>	<u>F</u>
A	3.06	3	1.02	$\frac{1.02}{1.31} = 0.77$
B	159.50	3	53.17	$\frac{53.17}{0.38} = 139.92$
AB	11.79	9 (=3x3)	1.31	$\frac{1.31}{0.38} = 3.44$
Error	6.09	16	0.38	
Total	180.44	31		

Hypothesis tests

$$AB: H_0: \sigma_{AB}^2 = 0$$

$$H_a: \sigma_{AB}^2 \neq 0$$

$$F = \frac{ms_{AB}}{ms_{\text{error}}} = \frac{1.31}{0.38} = 3.44$$

$$F_{0.05}(9, 16) = 2.55 < F$$

$\therefore$  reject  $H_0$ , i.e. AB is significant

$$A: H_0: \alpha_i = 0$$

$$H_a: \alpha_i's \text{ differ}$$

$$F = \frac{ms_A}{ms_{AB}} = \frac{1.02}{1.31} = 0.77$$

$$F_{0.05}(3, 9) = 3.86 > 0.77$$

$\therefore$  don't reject  $H_0$ , i.e. A is not sign.

$$B: H_0: \sigma_B^2 = 0$$

$$H_a: \sigma_B^2 > 0$$

$$F = \frac{ms_B}{ms_{\text{error}}} = \frac{53.17}{0.38} = 139.92$$

$$F_{0.05}(3, 16) = 3.24 < 139.92$$

$\therefore$  reject  $H_0$ , i.e. B is significant

(4)

2 @ The classical assumptions.

$\mu, \beta_i, \tau_j$  are all constants,

$$\sum_{i=1}^4 \beta_i = 0, \quad \sum_{j=1}^4 \tau_j = 0$$

$\epsilon_{ij} \sim N(0, \sigma^2)$  independently

The GLIM assumptions.

$\tilde{\mu}, \tilde{\beta}_i, \tilde{\tau}_j$  are all constants,

$$\tilde{\beta}_1 = 0, \quad \tilde{\tau}_1 = 0$$

$$\tilde{\mu} = \mu + \beta_1 + \tau_1$$

$$\tilde{\beta}_i = \beta_i - \beta_1 \quad (2 \leq i \leq 4)$$

$$\tilde{\tau}_j = \tau_j - \tau_1 \quad (2 \leq j \leq 4)$$

(a)

```

[i] ? $uni 16$
[i] ? $data yield$
[i] ? $read
[i] $REA? 1.25 1.43 1.32 1.31
[i] $REA? 2.05 1.56 1.68 1.69
[i] $REA? 1.95 2.00 1.83 1.81
[i] $REA? 1.75 1.93 1.70 1.59
[i] ? $ca block=%gl(4,1)$
[i] ? $ca treat=%gl(4,4)$
[i] ? $factor block 4 treat 4$
[i] ? $look yield block treat$
[i] ? $yvar yield$
```

	YIELD	BLOCK	TREAT
[o] 1	1.250	1.000	1.000
[o] 2	1.430	2.000	1.000
[o] 3	1.320	3.000	1.000
[o] 4	1.310	4.000	1.000
[o] 5	2.050	1.000	2.000
[o] 6	1.560	2.000	2.000
[o] 7	1.680	3.000	2.000
[o] 8	1.690	4.000	2.000
[o] 9	1.950	1.000	3.000
[o] 10	2.000	2.000	3.000
[o] 11	1.830	3.000	3.000
[o] 12	1.810	4.000	3.000
[o] 13	1.750	1.000	4.000
[o] 14	1.930	2.000	4.000
[o] 15	1.700	3.000	4.000
[o] 16	1.590	4.000	4.000

(3)

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## ANOVA

Source	SS	df	MS	F
Block	0.06417	3	0.02139	1.11290
Treat	0.7187	3	0.23957	12.4646
Error	0.17296	9	0.01922	
Total	0.95583	12		

Hypothesis test for treatments.

$$H_0: \tau_1 = \tau_2 = \tau_3 = \tau_4 = 0$$

$$H_a: \tau_j's \text{ differ}$$

$$\text{Given } \alpha = 0.05, F_{3,9}(0.05) = 3.86$$

$$\therefore F = 12.4646 > F_{3,9}(0.05)$$

∴ reject  $H_0$  and accept  $H_1$ , i.e. there is a significant difference among treat. means.

Hypothesis test for blocks.

$$H_0: \beta_1 = \beta_2 = \beta_3 = \beta_4 = 0$$

$$H_a: \beta_i's \text{ differ}$$

$$\therefore F = 1.1129 < F_{3,9}(0.05)$$

∴ don't reject  $H_0$ 

3. M.R.T.

Treat	1	4	2	3
Mean	1.327	1.742	1.745	1.898

$$MSerror = 0.01922, d.f. = 9, s.e. \text{ of mean} = \sqrt{0.01922/4} = 0.06932$$

P	2	3	4
Range	3.2	3.95	4.42

P	2	3	4
LS range	0.2218	0.2738	0.3064

(6)

## Tests

- 3 vs 1:  $1.898 - 1.327 = 0.571 > 0.3064 \Rightarrow$  significant  
 3 vs 4:  $1.898 - 1.742 = 0.156 < 0.2738 \Rightarrow$  not significant  
 3 vs 2:  $1.898 - 1.745 = 0.153 < 0.2218 \Rightarrow$  not significant  
 2 vs 1:  $1.745 - 1.327 = 0.418 > 0.2738 \Rightarrow$  significant  
 2 vs 4:  $1.745 - 1.742 = 0.003 < 0.2218 \Rightarrow$  not significant  
 4 vs 1:  $1.742 - 1.327 = 0.415 > 0.2218 \Rightarrow$  significant

Conclusions: treat 1 differs from treatments 2, 3, 4  
 while treatments 2, 3, 4 do not differ

- (e) From 2(a), we know  $\tilde{\epsilon}_2 = \epsilon_2 - \bar{t}_1$ , while  
 the l.s. estimators for  $\epsilon_2$  and  $\bar{t}_1$  are  
 $\bar{t}_2$  and  $\bar{t}_1$ , respectively, so the l.s. estimator  
 for  $\tilde{\epsilon}_2$  is

$$\bar{t}_2 - \bar{t}_1$$

$$\text{where } \bar{t}_2 = \frac{1}{4} \sum_{i=1}^4 X_{i2}, \quad \bar{t}_1 = \frac{1}{4} \sum_{i=1}^4 X_{i1}$$

Since  $X_{ij}$ 's are independent random variable  
 with variance  $\sigma_e^2$ ,

$$\begin{aligned}\text{Var}(\bar{t}_2 - \bar{t}_1) &= \text{Var}(\bar{t}_2) + \text{Var}(\bar{t}_1) \\ &= \frac{1}{16} \sum_{i=1}^4 \text{Var}(X_{i2}) + \frac{1}{16} \sum_{i=1}^4 \text{Var}(X_{i1}) \\ &= \frac{1}{16} \sum_{i=1}^4 \sigma_e^2 + \frac{1}{16} \sum_{i=1}^4 \sigma_e^2 \\ &= \frac{\sigma_e^2}{2}\end{aligned}$$

The s.e. of the l.s. estimator for  $\tilde{\epsilon}_2$  is then  
 $\sigma_e / \sqrt{2}$ . Estimate  $\sigma_e$  by  $s_e = \sqrt{MS_{\text{error}}} = 0.1386$ .  
 Then the s.e. of the l.s. estimator for  $\tilde{\epsilon}_2$  is

$$\frac{0.1386}{\sqrt{2}} = 0.09802$$

which is the same as the GLM output

(7)

3 a) This is a nested designs, since different batches are used at each site. The mathematical model is

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

$$1 \leq i \leq 2, \quad 1 \leq j \leq 3, \quad 1 \leq k \leq 5$$

where  $\mu$  - grand mean

$\alpha_i$  - effect of site  $i$

$\beta_{j(i)}$  - effect of batch  $j$  nested  
in ~~treatment~~ site  $i$

$\epsilon_{ijk}$  - experimental error.

From the given conditions, we know that sites are fixed effects but batches are random effects. The corresponding assumptions are:

$\alpha_1$  and  $\alpha_2$  are constants,  $\alpha_1 + \alpha_2 = 0$

$$\beta_{j(i)} \sim N(0, \sigma_B^2) \quad \left. \right\} \text{independently.}$$

$$\epsilon_{ijk} \sim N(0, \sigma_e^2)$$

(6)

(8)

b)

```

0] ? $units 30
i] ? $data content$ 
i] ? $read
i] $REA? 5.03 5.10 5.25 4.98 5.05
i] $REA? 4.64 4.73 4.82 4.95 5.06
i] $REA? 5.10 5.15 5.20 5.08 5.14
i] $REA? 5.05 4.96 5.12 5.12 5.05
i] $REA? 5.46 5.15 5.18 5.18 5.11
i] $REA? 4.90 4.95 4.86 4.86 5.07
i] ? $ca site=%gl(2,15) :batch=%gl(3,5)$ } 
i] ? $fac site 2 batch 3$ 
i] ? $yvar contents$ 

```



may use

\$ ca site=%gl(2,15) :batch=%gl(3,5)\$

\$ fac site 2 batch 6 \$

(5)

c) ANOVA

Bounce	SS	d.f.	MS	F
Sites	0.01825	1	0.01825	$\frac{0.01825}{0.11350} = 0.16$
Batches within sites	0.4540	4	0.11350	$\frac{0.11350}{0.01209} = 9.39$
Error	0.02902	24	0.01209	
Total	0.76247	29		

(7)

Tests from

Sites:  $F_{0.05}(1, 4) = 7.71 > 0.16$  $\Rightarrow$  InsignificantBatches within sites:  $F_{0.05}(4, 24) = 2.75 < 9.39$  $\Rightarrow$  Significant.

(9)

d) Note from the ANOVA that

$$S_e^2 = MS_{\text{error}} = 0.01209$$

d.f. for error = 24

$\therefore$  the 90% C.I. for  $\sigma_e^2$  is

$$\frac{24 S_e^2}{\chi^2_{24}(0.05)} < \sigma_e^2 < \frac{24 S_e^2}{\chi^2_{24}(0.95)}$$

From the  $\chi^2$ -table,

$$\chi^2_{24}(0.05) = 36.42$$

$$\chi^2_{24}(0.95) = 13.85$$

$\therefore$  The C.I. is

$$\frac{24 \times 0.01209}{36.42} < \sigma_e^2 < \frac{24 \times 0.01209}{13.85}$$

$$0.00796 < \sigma_e^2 < 0.0209$$

(7)

(10)

4(a) From the def. contrasts  $AB, BC, CD$  we derive  
the following interactions are also confounded.

$$AB \times BC \pmod{2} = AC$$

$$AB \times CD \pmod{2} = ABCD$$

$$BC \times CD \pmod{2} = BD$$

$$AB \times BC \times CD \pmod{2} = AD$$

For all 3 replicates - the 16 treat' combin's  
are assigned into 8 blocks as follows.

$$(1) \quad a \ b \ ab \ c \ ac \ bc \ abc \\ d \ ad \ bd \ abd \ cd \ acd \ bcd \ abcd$$

↓ By AB

$$\left\{ \begin{array}{l} Gp\ 1: (1) \ ab \ c \ abc \ d \ abd \ cd \ abcd \\ Gp\ 2: a \ b \ ac \ bc \ ad \ bd \ acd \ bcd \end{array} \right.$$

↓ By BC

$$\left\{ \begin{array}{l} Gp\ 1: (1) \ abc \ d \ abcd \\ Gp\ 2: ab \ c \ abd \ cd \end{array} \right.$$

$$\left\{ \begin{array}{l} Gp\ 3: a \ bc \ ad \ bcd \\ Gp\ 4: b \ ac \ bd \ acd \end{array} \right.$$

↓ By CD

$$\text{Block 1: } (1) \ abcd$$

$$2: abc \ d$$

$$3: ab \ cd$$

$$4: c \ abd$$

$$5: a \ bcd$$

$$6: bc \ ad$$

$$7: b \ acd$$

$$8: ac \ bd$$

(10)

ANOVA

<u>Source</u>	<u>d.f.</u>
A	1
B	1
C	1
D	1
ABC	1
ABD	1
ACD	1
BCD	1
Blocks	7
Replicates	2
Blocks $\times$ Repl	14
Error	26
Total	47

(12)

40) the  $\frac{1}{2}$ -replicate is

(1) b c be ad abd acd abcd

 $A \times (\text{the } \frac{1}{2}\text{-rep})$  (mod. 2) yields

a ab ac abc d bd cd bed

= the other  $\frac{1}{2}$ -rep.so A  $\in$  the def. contrast $B \times (\text{the } \frac{1}{2}\text{-rep})$  (mod. 2) yields

b (1) bc b abd ad abcd acd

= the same  $\frac{1}{2}$ -rep.so B  $\notin$  the def. contrast $C \times (\text{the } \frac{1}{2}\text{-rep})$  (mod. 2) yields

(12)

$c' bc \text{ or } b' acd abcd ad' abd$   
 $= \text{the same } \frac{1}{2} - \text{rep}.$

So C & the def. contrast

$D \times (\text{the } \frac{1}{2} - \text{rep}) \text{ (mod. 2) yields}$

$d' bd' cd' bcd' a' ab' ac' abc'$   
 $= \text{the other } \frac{1}{2} - \text{rep}.$

So D & the defining contrast

Hence the def. contrast is AD

Factor/interaction	Aliase
A	D
B	ABD
C	ACD
AB	BD
AC	CD
BC	ABCD
ABC	BCD

#### 4(c) ANOVA

Source	d.f.
Rep'l.	2
A or D	1
B or ABD	1
C or ACD	1
AB or BD	1
AC or CD	1
BC or ABCD	1
ABC or BCD	1
error	14
Total	23

(9)

(4)