

Chapter 24

Multifactor studies - when three or more factors are studied simultaneously

Notation

$$\mu_{i\cdot\cdot} = \sum_c \mu_{ijk} \quad \mu_{i\cdot k} = \sum_b \mu_{ijk} \quad \mu_{i\cdot k} = \sum_a \mu_{ijk}$$

$$\mu_{i\cdot\cdot} = \frac{\sum_j \sum_k \mu_{ijk}}{bc} \quad \mu_{i\cdot k} = \frac{\sum_j \sum_k \mu_{ijk}}{ac} \quad \mu_{i\cdot k} = \frac{\sum_j \sum_k \mu_{ijk}}{ab}$$

$$\mu_{\dots} = \frac{\sum_i \sum_j \sum_k \mu_{ijk}}{abc}$$

Main effect of the i^{th} level of factor A: $\alpha_i = \mu_{i\cdot\cdot} - \mu_{\dots}$
 j^{th} level of factor B: $\beta_j = \mu_{\cdot j \cdot} - \mu_{\dots}$
 k^{th} level of factor C: $\gamma_k = \mu_{\cdot \cdot k} - \mu_{\dots}$

Two factor interactions

- between factor A at the i^{th} level and factor B at the j^{th} level: $(\alpha\beta)_{ij} = \mu_{ij\cdot} - \mu_{i\cdot\cdot} - \mu_{\cdot j \cdot} + \mu_{\dots}$
- between factor A at the i^{th} level and factor C at the k^{th} level: $(\alpha\gamma)_{ik} = \mu_{i\cdot k} - \mu_{i\cdot\cdot} - \mu_{\cdot \cdot k} + \mu_{\dots}$
- between factor B at the j^{th} level and factor C at the k^{th} level: $(\beta\gamma)_{jk} = \mu_{\cdot jk} - \mu_{\cdot j \cdot} - \mu_{\cdot \cdot k} + \mu_{\dots}$

NOTE:

$$\sum (\alpha\beta)_{ij} = 0 \quad \forall i \text{ and } j$$

$$\sum (\alpha\gamma)_{ik} = 0 \quad \forall k \text{ and } i$$

$$\sum (\beta\gamma)_{jk} = 0 \quad \forall k \text{ and } j$$

Three factor interaction $(\alpha\beta\gamma)_{ijk} = \mu_{ijk} - [\mu_{\dots} + \alpha_i + \beta_j + \gamma_k + (\alpha\beta)_{ij} + (\alpha\gamma)_{ik} + (\beta\gamma)_{jk}]$

NOTE:

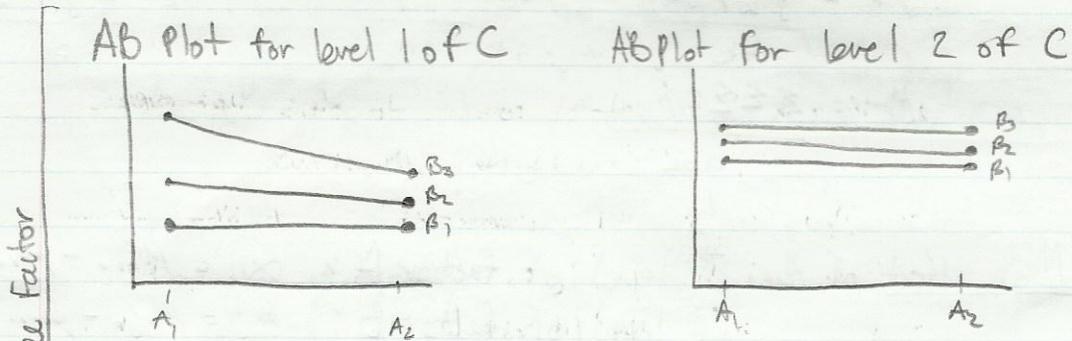
$$\sum_j (\alpha\beta\gamma)_{ijk} = 0 \quad \forall i, k$$

$$\sum_i (\alpha\beta\gamma)_{ijk} = 0 \quad \forall j, k$$

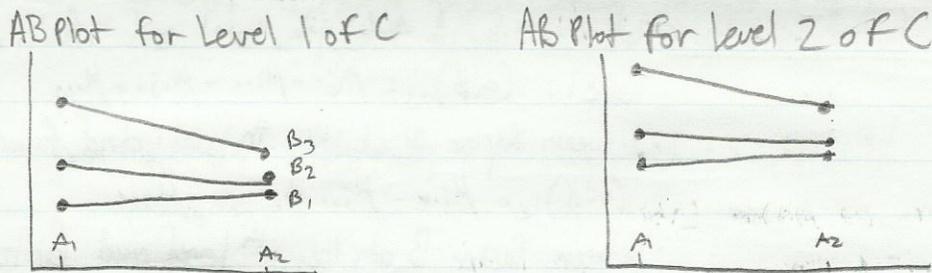
$$\sum_k (\alpha\beta\gamma)_{ijk} = 0 \quad \forall i, j$$

Cell Means $Y_{ijkm} = \mu_{ijk} + \epsilon_{ijkm}$

Factor Effects $Y_{ijkm} = \mu_{...} + \alpha_i + \beta_j + \delta_k + (\alpha\beta)_{ij} + (\alpha\delta)_{ik} + (\beta\delta)_{jk} + (\alpha\beta\delta)_{ijk} + \epsilon_{ijkm}$



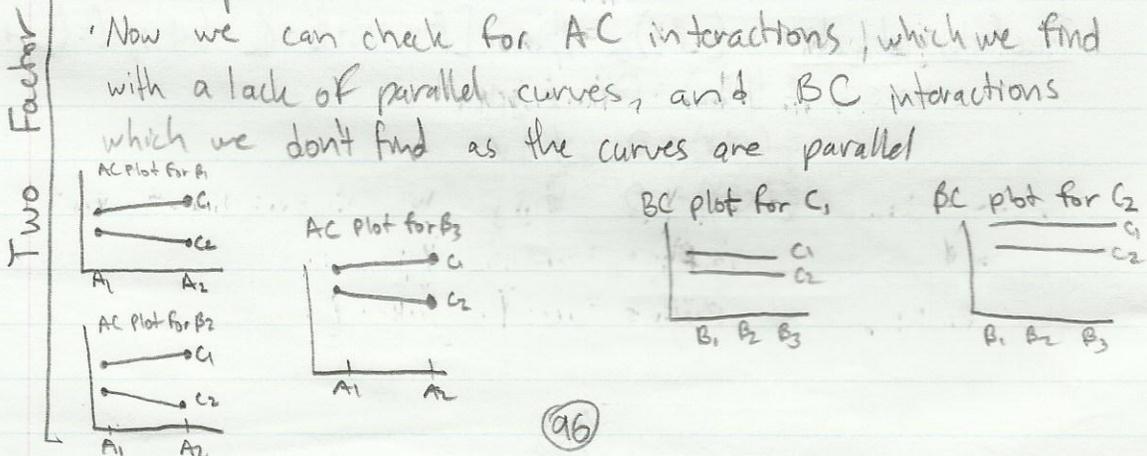
Here, as the two graphs won't be approximately parallel a three factor interaction will exist,



Here, as the two graphs will be approximately parallel a three factor interaction will not exist

We can also check these for two factor interactions which we find for AB because of the the curves aren't parallel in each panel

Now we can check for AC interactions, which we find with a lack of parallel curves, and BC interactions which we don't find as the curves are parallel



```

SAS proc sgscatter noautolegend; /* Two factor interactions */
      series x=A y=Y / group=B
      scatter x=A y=Y / group=B markerchar=judge
run;
/* You'd have to get separate graphs for each level of C
a clever data step could do this but spanel is easier */
proc spanel; /* three factor interactions
panelby c / rows=1 columns=k; /* Note: we can get interaction
scatter x=A y=Y / group=B; plots by fitting the
reg x=A y=Y / group=B; appropriate model in
run; glm w/ plots=all; */
proc sgscatter; /* Main effects Plot */
plot y*A / loess;
run;

```

Fitting the ANOVA Model

$$\begin{array}{l}
 \bar{Y}_{ijk.} = \sum_m Y_{ijkm} \\
 \bar{Y}_{ij..} = \sum_k \sum_m Y_{ijkm} \\
 \bar{Y}_{i..k.} = \sum_j \sum_m Y_{ijkm} \\
 \bar{Y}_{.j.k.} = \sum_i \sum_m Y_{ijkm} \\
 \bar{Y}_{i...} = \sum_j \sum_k \sum_m Y_{ijkm} \\
 \bar{Y}_{.j..} = \sum_i \sum_k \sum_m Y_{ijkm} \\
 \bar{Y}_{..k.} = \sum_i \sum_j \sum_m Y_{ijkm} \\
 \bar{Y}_{....} = \sum_i \sum_j \sum_k \sum_m Y_{ijkm}
 \end{array}
 \quad
 \begin{array}{l}
 \bar{Y}_{ijk.} = Y_{ijk.}/n \\
 \bar{Y}_{ij..} = Y_{ij..}/cn \\
 \bar{Y}_{i..k.} = Y_{i..k.}/bn \\
 \bar{Y}_{.j.k.} = Y_{.j.k.}/an \\
 \bar{Y}_{i...} = Y_{i...}/bcn \\
 \bar{Y}_{.j..} = Y_{.j..}/acn \\
 \bar{Y}_{..k.} = Y_{..k.}/abn \\
 \bar{Y}_{....} = \bar{Y}_{....}/abcn
 \end{array}$$

$$\begin{array}{l}
 \hat{\mu}_{ijk.} = \bar{Y}_{ijk.} \rightarrow \hat{Y}_{ijkm} = \bar{Y}_{ijk.} \text{ w/ } e_{ijkm} = Y_{ijkm} - \hat{Y}_{ijkm} = Y_{ijkm} - \bar{Y}_{ijk.} \\
 \hat{\mu}_{i...} = \bar{Y}_{i...} \\
 \hat{\alpha}_i = \bar{Y}_{i...} + \bar{Y}_{....} \\
 \hat{\beta}_j = \bar{Y}_{.j..} + \bar{Y}_{....} \\
 \hat{\gamma}_k = \bar{Y}_{..k.} + \bar{Y}_{....} \\
 (\alpha\beta)_{ij} = \bar{Y}_{ij..} + \bar{Y}_{i...} + \bar{Y}_{.j..} + \bar{Y}_{....} \\
 (\alpha\gamma)_{ik} = \bar{Y}_{i..k.} + \bar{Y}_{i...} + \bar{Y}_{..k.} + \bar{Y}_{....} \\
 (\beta\gamma)_{jk} = \bar{Y}_{.j.k.} + \bar{Y}_{.j..} + \bar{Y}_{..k.} + \bar{Y}_{....} \\
 (\alpha\beta\gamma)_{ijk.} = \bar{Y}_{ijk.} - \bar{Y}_{ij..} - \bar{Y}_{i..k.} - \bar{Y}_{.j.k.} + \bar{Y}_{i...} + \bar{Y}_{.j..} + \bar{Y}_{..k.} - \bar{Y}_{....}
 \end{array}$$

SAS to fit the ANOVA MODEL

```
proc glm outstat=full; /* Check which interactions
class A B C; we can drop (Type III) */
model Y = A|B|C /solution;
run;
proc glm outstat=reduced; /* fit the reduced model */
class A B C;
model Y = A B|C /solution;
run
/* We can test that it's appropriate to use the reduced */
data test1;
set full reduced;
if _SOURCE_="ERROR";
proc means data=test1 noprint;
var ss df;
output out=test2 min=minss mindf max=maxss maxdf;
data nested;
set test2;
fstar = ((maxss - minss) / (maxdf - mindf)) / (minss / mindf);
pvalue = 1 - cdf('F', fstar, maxdf - mindf, mindf);
proc print;
run; /* Look at P-value */
```

- Sum of Squares and Mean squared errors are outlined in table 24.5 on pg. 1006
- Check Residuals the same way:
 - * Residuals vs each treatment
 - * Normal Probability Plot
 - * Residuals vs predicted Val

More sums of squares

$$SSTO = SSTreatment + SSE$$

$$\equiv \sum_i \sum_j \sum_k \sum_m (Y_{ijkm} - \bar{Y} \dots)^2 = n \sum_i \sum_j \sum_k (\bar{Y}_{ijk.} - \bar{Y} \dots)^2 + \sum_i \sum_j \sum_k \sum_m e_{ijkm}^2$$

$$SSTreatment = SSA + SSB + SSC + SSAB + SSAC + SSBC + SSABC$$

where $SSA = nbc \sum_i (\bar{Y}_{i...} - \bar{Y} \dots)^2$

$$SSB = nac \sum_j (\bar{Y}_{.j..} - \bar{Y} \dots)^2$$

$$SSC = nab \sum_k (\bar{Y}_{...k.} - \bar{Y} \dots)^2$$

$$SSAB = nc \sum_i \sum_j (\bar{Y}_{ij..} - \bar{Y}_{i...} - \bar{Y}_{.j..} + \bar{Y} \dots)^2$$

$$SSAC = nb \sum_i \sum_k (\bar{Y}_{i.k.} - \bar{Y}_{i...} - \bar{Y}_{...k.} + \bar{Y} \dots)^2$$

$$SSBC = na \sum_j \sum_k (\bar{Y}_{.jk.} - \bar{Y}_{.j..} - \bar{Y}_{...k.} + \bar{Y} \dots)^2$$

$$SSABC = n \sum_i \sum_j \sum_k (\bar{Y}_{ijk.} - \bar{Y}_{ij..} - \bar{Y}_{i.k.} - \bar{Y}_{.jk.} + \bar{Y}_{i...} + \bar{Y}_{.j..} + \bar{Y}_{...k.} - \bar{Y} \dots)^2$$

degrees of freedom

1 a-1

1 b-1

1 c-1

1 (a-1)(b-1)

1 (a-1)(c-1)

1 (b-1)(c-1)

1 (a-1)(b-1)(c-1)

$$SSTO = SSA + SSB + SSC + SSAB + SSAC + SSBC + SSABC + SSE$$

F-Test for factor effect

H_0 : All of this factor level = 0 i.e. $(\alpha\beta\gamma)_{ijk} = 0$

H_a : Not H_0

$$F^* = \frac{\text{Associated Mean Square}}{\text{Mean Square Error}} \quad \text{i.e.} \quad F^* = \frac{MSABC}{MSE}$$

Reject if $F^* > F[1-\alpha, \text{degrees of freedom}, (n-1)/abc]$ i.e. Reject if $F^* > F[1-\alpha, (a-1)(b-1)(c-1), (n-1)abc]$

Strategy for analysis

- 1.) Do three factor interactions exist?
- 2.) If not do two factor interactions exist?
- 3.) If not can we drop single main effect?
- 4.) If yes in 1, can we make a transformation to make it unimportant?
- 5.) If not analyze three factors jointly in terms of M_{ijk}
- 6.) If theres just one important two factor analyze the effects jointly in terms of $M_{i.}, M_{.jk.}, M_{i.k.}$. Analyze third effect separately

7.) If there are two or three important two-factor interactions analyze the three factors jointly in terms of μ_{ijk} .

** There are exceptions to this strategy **

Estimation of factor level Mean $\hat{\mu}_{i...} = \bar{Y}_{i...}$

w/ variance $s^2(\bar{Y}_{i...}) = \frac{MSE}{nbc}$

Confidence intervals for factor level mean: $\bar{Y}_{i...} \pm t [1-\alpha/2, (n-1)abc] s(\bar{Y}_{i...})$

Contrast of factor level means: $L = \sum c_i \mu_{i...}$ where $\sum c_i = 0$

Confidence intervals for contrasts: $\hat{L} \pm t [1-\alpha/2, (n-1)abc] s(\hat{L})$

w/ $\hat{L} = \sum c_i \bar{Y}_{i...}$

$s^2(\hat{L}) = \frac{MSE}{nbc} \sum c_i^2$

Hypothesis Test for contrasts $H_0: L = 0$

$H_a: L \neq 0$

$t^* = \frac{\hat{L}}{s(\hat{L})}$; reject when $> t [1-\alpha/2, (n-1)abc]$

Multiple Contrasts Conf. Interval

$\hat{L} \pm \frac{1}{2} q [1-\alpha, a, (n-1)abc]$ TUKEY

$\hat{L} \pm (a-1) F [1-\alpha, a-1, (n-1)abc]$ SCHEFFE

$\hat{L} \pm t [1-\alpha/2g, (n-1)abc]$ BONFERRONI

Multiple Contrasts Hypothesis Test

$q^* = \frac{\sqrt{2} \hat{\sigma}}{s(\hat{L})}$ reject if $> q [1-\alpha, a, (n-1)abc]$

$F^* = \frac{\hat{L}^2}{(a-1)s^2(\hat{L})}$ reject if $> F [1-\alpha, a-1, (n-1)abc]$

$t^* = \frac{\hat{L}}{s(\hat{L})}$ reject if $> t [1-\alpha/2g, (n-1)abc]$

SAS proc glm; /* All pairwise differences */

class A B C;

model Y = A B C;

lsmeans A / pdiff adjust = tukey alpha = .05 cl;

lsmeans B*C / pdiff adjust = tukey

run;

```
proc glimmix; /* Contrasts */
```

```
class ABC;
```

```
model y = A B C B*C;
```

```
lsestimate A "A" 1 -1 / adjust = +schiffé bonferroni alpha = .05 cl;
```

```
lsestimate B*C "d1" -1 1 0 0,  
"d2" 0 0 -1 1 / adjust = +schiffé bonferroni alpha = .05 cl;
```

```
run;
```

skip 24.6 & 24.7