

Stat 8311 – Estimating treatment means, unbalanced data

```
> searle <- data.frame(soil = rep(c("s1", "s2"), c(7, 8)),
+   var = c("v1", "v2", "v3")[c(1, 1, 1, 2, 2, 3, 3, 1, 1,
+   1, 1, 2, 3, 3, 3)], y = c(6, 10, 11, 13, 15, 14,
+   22, 12, 15, 19, 18, 31, 18, 9, 12))
```

Here are the observed row means, column means and cell means:

```
> mean(searle$y)
```

```
[1] 15
```

```
> tapply(searle$y, searle$soil, mean)
```

```
   s1    s2
13.00 16.75
```

```
> tapply(searle$y, searle$var, mean)
```

```
   v1     v2     v3
13.00000 19.66667 15.00000
```

```
> xtabs(y ~ soil + var, data = searle)/xtabs(~soil + var, data = searle)
```

```
   var
soil v1 v2 v3
s1   9 14 18
s2  16 31 13
```

Next, fit using two orders. I'll use the SAS contrasts, although *the result is invariant under reparameterization because only projections are used*, so the results can be compared to SAS.

```
> opt <- options(contrasts = c("contr.SAS", "contr.poly"))
> m1 <- aov(y ~ soil * var, data = searle)
> m2 <- update(m1, ~var * soil)
```

Fitted means are evidently based on the projections onto various subspaces. Consider first the projections for model m1.

```
> (p1 <- proj(m1)[, ])
```

```
 (Intercept)  soil      var  soil:var  Residuals
1             15 -2.00 -2.7659574 -1.234043 -3.0000e+00
2             15 -2.00 -2.7659574 -1.234043  1.0000e+00
3             15 -2.00 -2.7659574 -1.234043  2.0000e+00
4             15 -2.00  5.0531915 -4.053191 -1.0000e-00
5             15 -2.00  5.0531915 -4.053191  1.0000e-00
6             15 -2.00 -0.9042553  5.904255 -4.0000e+00
7             15 -2.00 -0.9042553  5.904255  4.0000e+00
8             15  1.75 -1.6755319  0.925532 -4.0000e+00
9             15  1.75 -1.6755319  0.925532 -1.0000e-00
10            15  1.75 -1.6755319  0.925532  3.0000e+00
```

```

11          15  1.75 -1.6755319  0.925532  2.0000e+00
12          15  1.75  6.1436170  8.106383 -4.1281e-16
13          15  1.75  0.1861702 -3.936170  5.0000e+00
14          15  1.75  0.1861702 -3.936170 -4.0000e+00
15          15  1.75  0.1861702 -3.936170 -1.0000e+00

```

In examining `p1`, the first column, the projection on the column of ones, we get one value for each observation, always equal to \bar{y}_{+++} . The second column is the projection of y on the part of the soil space that is orthogonal to the column of ones, relative to the correct inner product. Since soil has two levels, this is the projection on a one-dimensional subspace, and the projection is equal to -2 for observations with `soil = s1`, and 1.75 for `soil = s2`. It is therefore reasonable for the means for soils to be $15 - 2 = 13$ for `s1` and $15 + 1.75 = 16.75$ for `s2`.

Next consider the column for `var`. This is the projection on a two dimensional subspace for the three varieties, after adjusting for the overall mean and for `soil`. *The number of unique values for this projection is not equal to three, but rather there are six unique values, apparently corresponding to the six cells in the table.* Thus it is not clear how one would define a mean for each of the three varieties. What R appears to do is *compute a weighted average of the cell means*:

```
> tapply(p1[, 1] + p1[, 3], searle$var, mean)
```

```

      v1      v2      v3
12.85714 20.41667 14.75000

```

Finally the `(soil,var)` means are simply the cell means

```
> xtabs(apply(p1[, -5], 1, sum) ~ soil + var, data = searle)/xtabs(~soil +
+   var, data = searle)
```

```

      var
soil v1 v2 v3
s1   9 14 18
s2  16 31 13

```

```
> print(model.tables(m1, type = "mean"), digits = 6)
```

Tables of means

Grand mean

15

```

soil
  s1  s2
 13 16.75
rep  7  8.00

```

```

var
      v1      v2      v3
rep 7.0000 3.0000 5.00

```

```

soil:var
  var
soil  v1 v2 v3
  s1   9 14 18
  rep  3  2  2
  s2  16 31 13
  rep  4  1  3

```

```
> proj(m2 <- update(m1, ~var * soil))[, ]
```

	(Intercept)	var	soil	var:soil	Residuals
1	15	-2.000000e+00	-2.765957	-1.234043	-3.000000e+00
2	15	-2.000000e+00	-2.765957	-1.234043	1.000000e+00
3	15	-2.000000e+00	-2.765957	-1.234043	2.000000e+00
4	15	4.666667e+00	-1.613475	-4.053191	-1.000000e+00
5	15	4.666667e+00	-1.613475	-4.053191	1.000000e+00
6	15	8.881784e-16	-2.904255	5.904255	-4.000000e+00
7	15	8.881784e-16	-2.904255	5.904255	4.000000e+00
8	15	-2.000000e+00	2.074468	0.925532	-4.000000e+00
9	15	-2.000000e+00	2.074468	0.925532	-1.000000e-00
10	15	-2.000000e+00	2.074468	0.925532	3.000000e+00
11	15	-2.000000e+00	2.074468	0.925532	2.000000e+00
12	15	4.666667e+00	3.226950	8.106383	-1.761829e-16
13	15	8.881784e-16	1.936170	-3.936170	5.000000e+00
14	15	8.881784e-16	1.936170	-3.936170	-4.000000e+00
15	15	8.881784e-16	1.936170	-3.936170	-1.000000e+00

```
> model.tables(m2, type = "mean")
```

Tables of means

Grand mean

15

```

var
  v1    v2 v3
  13 19.67 15
rep  7  3.00  5

```

```

soil
  s1    s2
  12.52 17.17
rep  7.00  8.00

```

```

var:soil
  soil
var  s1 s2
  v1  9 16
  rep  3  4
  v2 14 31

```

```

rep 2 1
v3 18 13
rep 2 3

```

```

options nocenter ls=75;
proc glm data=work.searle;
class var soil;
model days = var soil var*soil/ss1 ss2 ss3 solution;
lsmeans var soil var*soil;
run;

```

Dependent Variable: days

Source	DF	Sum of Squares	Mean Square	F Value	Pr > F
Model	5	400.0000000	80.0000000	6.00	0.0103
Error	9	120.0000000	13.3333333		
Corrected Total	14	520.0000000			
	R-Square	Coeff Var	Root MSE	days Mean	
	0.769231	24.34322	3.651484	15.00000	

Source	DF	Type I SS	Mean Square	F Value	Pr > F
var	2	93.3333333	46.6666667	3.50	0.0751
soil	1	83.9007092	83.9007092	6.29	0.0334
var*soil	2	222.7659574	111.3829787	8.35	0.0089

Source	DF	Type II SS	Mean Square	F Value	Pr > F
var	2	124.7340426	62.3670213	4.68	0.0405
soil	1	83.9007092	83.9007092	6.29	0.0334
var*soil	2	222.7659574	111.3829787	8.35	0.0089

Source	DF	Type III SS	Mean Square	F Value	Pr > F
var	2	192.1276596	96.0638298	7.20	0.0135
soil	1	123.7714286	123.7714286	9.28	0.0139
var*soil	2	222.7659574	111.3829787	8.35	0.0089

Parameter estimates

Parameter	Estimate	Error	t Value	Pr > t
Intercept	13.0000000 B	2.10818511	6.17	0.0002
var 1	3.0000000 B	2.78886676	1.08	0.3100
var 2	18.0000000 B	4.21637021	4.27	0.0021
var 3	0.0000000 B	.	.	.
soil 1	5.0000000 B	3.33333333	1.50	0.1679
soil 2	0.0000000 B	.	.	.
var*soil 1 1	-12.0000000 B	4.34613494	-2.76	0.0221
var*soil 1 2	0.0000000 B	.	.	.

The GLM Procedure Least Squares Means

```

var    days LSMEAN
1      12.5000000

```

```
2      22.5000000
3      15.5000000
```

```
soil    days LSMEAN
1      13.6666667
2      20.0000000
```

```
var    soil    days LSMEAN
1      1      9.0000000
1      2     16.0000000
2      1     14.0000000
2      2     31.0000000
3      1     18.0000000
3      2     13.0000000
```

Disconnected data

The Big Idea with disconnected data is that the dimensions of subspaces may change depending on the order of fitting.

```
> xtabs(y ~ row + col, data = d1 <- data.frame(row = factor(c(1:4,
+      4)), col = factor(c(1:4, 3)), y = 1:5))
```

```
      col
row 1 2 3 4
  1 1 0 0 0
  2 0 2 0 0
  3 0 0 3 0
  4 0 0 5 4
```

```
> (a1 <- aov(y ~ row + col, data = d1))
```

Call:

```
aov(formula = y ~ row + col, data = d1)
```

Terms:

```
              row col
Sum of Squares  9.5 0.5
Deg. of Freedom  3  1
```

2 out of 7 effects not estimable
Estimated effects may be unbalanced

```
> (a2 <- update(a1, ~col + row))
```

Call:

```
aov(formula = y ~ col + row, data = d1)
```

Terms:

```

                col row
Sum of Squares    8  2
Deg. of Freedom   3  1

```

2 out of 7 effects not estimable
 Estimated effects may be unbalanced

```
> model.tables(a1, type = "m")
```

Tables of means
 Grand mean

3

```

row
  1 2 3  4
  1 2 3 4.5
rep 1 1 1 2.0

```

```

col
  1 2  3  4
  3 3 3.25 2.5
rep 1 1 2.00 1.0

```

The `Anova` command in R fails with empty cells because the dimensions of subspaces change, depending on the order of fitting. This could be avoided if `Anova` refit models rather than using the Wald approach to testing.

SAS recognizes that the disconnected data has row and column effects that are not estimable.