

# SAS Lab 2

Hao Zhang

## Objective of the Lab

- Run SAS for the two-way models
- Test hypothesis for the interaction effects
- Test hypothesis for the mean effects
- Multiple comparisons for the main effects
- Use estimate statement for any contrast
- Know the difference of the ordering of factors in the `class` statement
- Know the difference between lsmeans and means statements

An experiment was conducted to determine the effects of three different pesticides of the yield of fruit from two different varieties of a citrus tree. Six trees of each variety were randomly selected from an orchard. The four pesticides were then randomly assigned to two trees of each variety, and applications were made according to recommended levels. Yields of fruits (in bushels per tree) were obtained after the test period.

Variety	Pesticide		
	1	2	3
1	49	50	38
	39	55	*Missing: results in unequal sample sizes
2	55	67	53
	41	58	42

\* Note: This tree was accidentally hit by a vehicle and has no yield data.

## Two-way Complete Model

1. This is a completely randomized design. Without any knowledge about the interaction, we will employ the two way complete model with Pesticide as Factor A (with 3 levels) and Variety as Factor B (with two levels).

```
data pest;
input variety pesticide yield;
lines;
1 1 49
1 1 39
1 2 50
1 2 55
1 3 38
2 1 55
2 1 41
2 2 58
2 3 53
2 3 42
2 2 67
;
proc print;
run;
```

variety ←

pesticide

skip missing

Default  $\alpha=0.05$

proc glm data=pest; ~~class~~ pesticide variety; ~~model~~ yield=variety pesticide variety\*pesticide; run;

(1) (2) (3)

interaction

2. Is there a significant interaction effect? (3)

The p-values 0.82. Fail to reject the null hypothesis. No significant interaction effects.

**Always use the p-value that corresponds to the Type III SS. (Sum Squares)**

3. Are there significant differences in the mean yields among the three pesticides? Use  $\alpha = 0.10$

The p-values is 0.089. We reject the null hypothesis at  $\alpha = 0.1$  level of significance and believe there is a significance among the three main effects of pesticide.

4. Are there significant differences in the mean yields between the two varieties? Use  $\alpha = 0.10$ .

The p-value is 0.1998 and we fail to reject the null hypothesis which says no difference between the main effects of variety.

5. Find 90% simultaneous confidence intervals for the pairwise comparisons of the main effects of pesticide.

```
proc glm data=pest;
class variety pesticide;
model yield=variety pesticide variety*pesticide;
lsmeans pesticide/cl pdiff adjust=Tukey;
run;
```

Default  $\alpha=0.05$

Least Squares Means for Effect pesticide				
i	j	Difference Between Means	Simultaneous 90% Confidence Limits for LSMean(i)-LSMean(j)	
1	2	-11.500000	-24.940808	1.940808
1	3	3.250000	-11.777280	18.277280
2	3	14.750000	-0.277280	29.777280

Default  $\alpha=0.05$   
otherwise specify

```
lsmeans pesticide / cl pdiff adjust=tukey(alpha=0.10;
```

Difference Between		Simultaneous 95% Confidence Limits
j	j	Means for LSMean(i)-LSMean(j)
1	2	-11.500000 -28.139482, 5.139482
1	3	3.250000 -15.353507, 21.853507
2	3	14.750000 -3.853507, 33.353507

"Confidence + least"

- **ci**: To produce the least squares estimates and the confidence interval for each treatment means (these are not simultaneous confidence intervals).
- **pdiff**: To provide confidence intervals for pairwise differences. If a method for simultaneous confidence intervals are provided through **adjust =**, then simultaneous confidence intervals are provided. Compare the difference if you use the following statement:

```
lsmeans pesticide/cl pdiff;
```

6. Find a 95% confidence interval for the difference between the mean yield for the two varieties (variety 2-variety 1) when pesticide 1 is applied.

First we must know how SAS coded the treatments. Since `pesticide` is entered into the `class` statement before `variety`, `pesticide` is Factor A and `variety` is Factor B. We are therefore comparing  $\mu_{12}$  with  $\mu_{11}$ .

Factor A and variety is factor B, we are therefore comparing  $\mu_{12}$  with  $\mu_{11}$

$\mu_{12} - \mu_{11} = (\mu + \alpha_1 + \beta_2 + \text{interaction}) - (\mu + \alpha_1 + \beta_1 + (\alpha\beta)_{11})$

$= (-\beta_1 + \beta_2) + (-\alpha\beta)_{11} + (\alpha\beta)_{12}$

$\begin{matrix} 11 & 21 \\ 12 & 22 \\ 13 & 23 \end{matrix} \}$  6 total interaction

```
estimate "v2-v1|pesticide=1" variety -1 10 variety*pesticide -1 1 0 0 0 0;
```

SAS output provides estimate=4.0000000 and standard error=7.23187389. We will need to find the  $t$  critical value  $t_{5,0.025} = 2.5706$ . We therefore have the 95% confidence interval  $4.0 \pm 18.59$ . Estimate  $\pm (t_{df, \alpha/2})(SE)$

Here we are estimating  $\mu_{21} - \mu_{31}$ , for which we must first express it in terms of main-effects and interaction effects.

$$\begin{aligned} \mu_{21} - \mu_{31} &= (\mu + \alpha_2 + \beta_1 + (\alpha\beta)_{21}) - (\mu + \alpha_3 + \beta_1 + (\alpha\beta)_{31}) \\ &= (\alpha_2 - \alpha_3) + ((\alpha\beta)_{21} - (\alpha\beta)_{31}). \end{aligned}$$

estimate "P2-P3|v=1" pesticide 0 1 -1 variety\*pesticide 0 0 1 0 -1 0;  
 variety=1

code follows AxB format  
depend which variables comes before "class"

7. In this problem, I will show you the order that the factors appear in the `class` statement is important. Compare the output from the following program:

```
proc glm data=pest;
class pesticide variety;
model yield=variety pesticide variety*pesticide/solution;
run;
```

*Shows detailed parameter*

with those from

```
proc glm data=pest;
class variety pesticide;
model yield=variety pesticide variety*pesticide/solution ;
run;
```

We see that the interaction plots are different and the treatments are coded differently.

## Two-way Main-Effects Model

Suppose the experimenter has knowledge that no interaction effects exist. Then the two-way main effects model can be used. Let us see how to get the simultaneous confidence intervals in Problem 5 and the confidence interval in Problem 6.

```
proc glm data=pest;
class pesticide variety;
model yield=pesticide variety;
lsmeans pesticide/cl pdiff adjust=Tukey;
estimate 'v2-v1' -1 1;
run;
```

Note how the p-values changed! The confidence intervals for the pairwise comparison become shorter.

 Download

 Print

 

### Activity Details

Task: View this topic