

Module 2: Completely Randomized Designs

Pairwise Comparisons

What Are Pairwise Comparisons?

Which treatment means differ from each other?

Pairwise comparisons test:

- μ_1 VS μ_2
- μ_1 VS μ_3
- μ_2 VS μ_3

For t treatments, there are: $\binom{t}{2} = \frac{t(t-1)}{2}$ possible pairwise comparisons.

Example 2.1: Running Shoes

Response: Lap time (seconds)

Treatment structure:

- One-way
- Factor: Shoe type
- 3 Levels: control, lightweight, and stability
- $t = 3$

Experimental structure:

- CRD
- Experimental Unit: Individual ($r = 2$)
- Measurement Unit: Individual ($N = 6$)

Example 2.1: Shoes

With $t = 3$ shoe types:

- Control vs Lightweight ($H_0 : \mu_1 - \mu_2 = 0$)
- Control vs Stability ($H_0 : \mu_1 - \mu_3 = 0$)
- Lightweight vs Stability ($H_0 : \mu_2 - \mu_3 = 0$)

That's _____ comparisons.

Pairwise Comparisons as Contrasts

Each pairwise comparison is a contrast:

- Control vs Lightweight: $C = (1, -1, 0)$
- Control vs Stability: $C = (1, 0, -1)$
- Lightweight vs Stability: $C = (0, 1, -1)$

The Multiple Testing Problem

Each comparison is a valid t-test.

But performing *many tests* increases the chance of a:

- Type I error (false positive)

This is called the *family-wise error rate*.

Why Error Rates Inflate

Suppose each test uses $\alpha = 0.05$. Then:

- Probability of *no* false positives (i.e., correctly fail to reject the null):
 $1 - \alpha = 1 - 0.05 = 0.95$
- Assuming independence, the probability of all tests leading to *no* false positives: $(1 - \alpha)^{\# \text{ tests}} = (0.95)^3 = 0.857$
- Probability of *at least one* false positive (i.e., a Type I error):
 $1 - [(1 - \alpha)^{\# \text{ tests}}] = 1 - 0.857 = 0.143.$

Note this increases with the number of tests.

What We Need

When doing many comparisons, we want to:

- Control the overall Type I error rate
- While still detecting real differences

Post Hoc Tests: adjustments to the p-values (making them larger) and/or confidence intervals (making them wider).

Fisher's Protected LSD

1. First perform the overall ANOVA F-test
2. Only if F is significant, perform all pairwise t-tests

Note: From the ANOVA on Example 2.1 Shoe Types: ($F = 21.5$; $df = 2,3$; $p = 0.017$)

```
1 library(emmeans)
2 shoe_lsmeans <- emmeans(shoe_mod, specs = ~ Shoe)
3 pairs(shoe_lsmeans, infer = c(TRUE, TRUE), adjust = "none")
```

contrast	estimate	SE	df	lower.CL	upper.CL	t.ratio	p.value
Control - Lightweight	-8	2	3	-14.36	-1.64	-4.000	0.0280
Control - Stability	5	2	3	-1.36	11.36	2.500	0.0877
Lightweight - Stability	13	2	3	6.64	19.36	6.500	0.0074

Confidence level used: 0.95

```
1 library(multcomp)
2 cld(shoe_lsmeans, adjust = "none", Letters = LETTERS, decreasing = T)
```

Shoe	emmean	SE	df	lower.CL	upper.CL	.group
Lightweight	69	1.41	3	64.5	73.5	A
Control	61	1.41	3	56.5	65.5	B
Stability	56	1.41	3	51.5	60.5	B

Confidence level used: 0.95

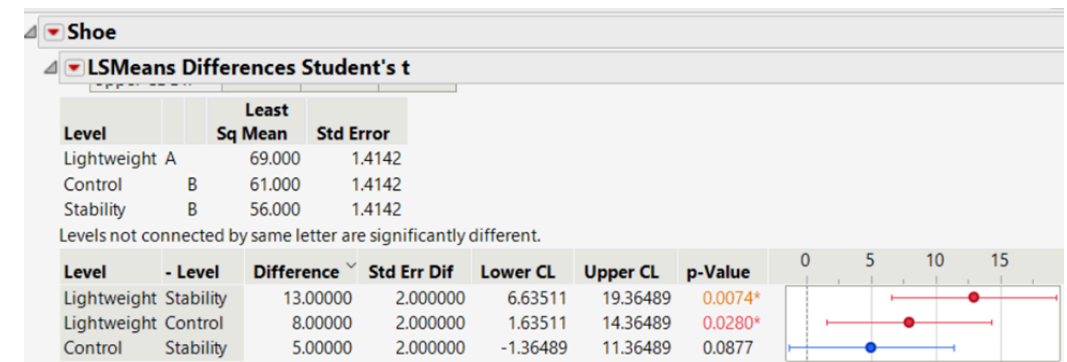
significance level used: $\alpha = 0.05$

NOTE: If two or more means share the same grouping symbol, then we cannot show them to be different.

But we also did not show them to be the same.

▼ Shoe > LSMeans Student's t

▼ LSMeans Differences Student's t > Ordered Differences Report



Tukey's Honestly Significant Difference (HSD)

- Designed specifically for *all* pairwise comparisons
- Controls family-wise error rate

```
1 library(emmeans)
2 shoe_lsmeans <- emmeans(shoe_mod, specs = ~ Shoe)
3 pairs(shoe_lsmeans, infer = c(TRUE, TRUE), adjust = "tukey")
```

contrast	estimate	SE	df	lower.CL	upper.CL	t.ratio	p.value
Control - Lightweight	-8	2	3	-16.36	0.358	-4.000	0.0560
Control - Stability	5	2	3	-3.36	13.358	2.500	0.1681
Lightweight - Stability	13	2	3	4.64	21.358	6.500	0.0151

Confidence level used: 0.95

Conf-level adjustment: tukey method for comparing a family of 3 estimates

P value adjustment: tukey method for comparing a family of 3 estimates

```
1 cld(shoe_lsmeans, adjust = "tukey", Letters = LETTERS, decreasing = T)
```

Shoe	emmean	SE	df	lower.CL	upper.CL	.group
Lightweight	69	1.41	3	62.2	75.8	A
Control	61	1.41	3	54.2	67.8	AB
Stability	56	1.41	3	49.2	62.8	B

Confidence level used: 0.95

Conf-level adjustment: sidak method for 3 estimates

P value adjustment: tukey method for comparing a family of 3 estimates

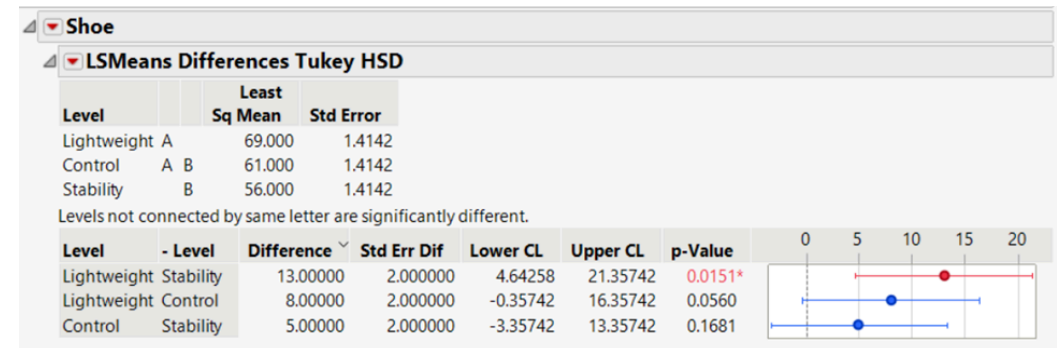
significance level used: alpha = 0.05

NOTE: If two or more means share the same grouping symbol,
then we cannot show them to be different.

But we also did not show them to be the same.

♥ Shoe > LSMeans Tukey HSD

♥ LSMeans Differences Tukey HSD > Ordered Differences Report



Bonferroni Adjustment

Bonferroni controls error by:

- Using a smaller significance level

$$\alpha^* = \frac{\alpha}{\text{number of tests}}$$

Simple, but often conservative.

Dunnet's Test

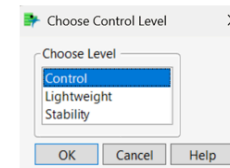
Useful when interest lies in comparing all treatments to a control.

```
1 library(emmeans)
2 shoe_lsmeans <- emmeans(shoe_mod, specs = ~ Shoe)
3 contrast(shoe_lsmeans,
4   method = "trt.vs.ctrl", # Treatment vs. control contrast
5   adjust = "dunnet", # Dunnett adjustment
6   infer = c(TRUE, TRUE),
7   ref = "Control"
8 )
```

contrast	estimate	SE	df	lower.CL	upper.CL	t.ratio	p.value
Lightweight - Control	8	2	3	0.0241	15.98	4.000	0.0496
Stability - Control	-5	2	3	-12.9759	2.98	-2.500	0.1511

Confidence level used: 0.95
Conf-level adjustment: dunnett method for 2 estimates
P value adjustment: dunnett method for 2 tests

▼ Shoe > LSMeans Dunnet > Choose Level (Control)



Shoe						
LSMeans Differences Dunnett						
α= 0.050 Q= 3.86634 Control=Control Adjustment = Dunnett						
Level	- Level	Difference	Std Err Dif	Lower CL	Upper CL	p-Value
Lightweight	Control	8.00000	2.000000	0.2673	15.73268	0.0458*
Stability	Control	-5.00000	2.000000	-12.7327	2.73268	0.1405