

ANALYSIS OF VARIANCE II

BIOS 662

Michael G. Hudgens, Ph.D.

mhudgens@bios.unc.edu

<http://www.bios.unc.edu/~mhudgens>

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Outline

- Multiple Comparisons
 - Scheffe
 - Tukey
 - Bonferroni

- Ch 12 of text

Multiple Comparisons

- Suppose we do n independent tests, each with probability α of making a type I error
- Suppose all n null hyp are true
- What is the probability of making at least one type I error?

$$1 - (1 - \alpha)^n$$

Multiple Comparisons

- Table 12.1: Probability of rejecting one or more null hyps when n independent tests are carried out at the α level and each null hyp is true

n	α		
	0.01	0.05	0.10
1	0.01	0.05	0.10
2	0.02	0.10	0.19
3	0.03	0.14	0.27
4	0.04	0.19	0.34
5	0.05	0.23	0.41
10	0.10	0.40	0.65
20	0.18	0.64	0.88
100	0.63	0.99	1.00

Multiple Comparisons

- Def 12.2 The probability of incorrectly rejecting at least one of the true null hyp in an experiment involving one or more tests or comparisons is called the *per experiment error rate (PEER)*
- PEER aka *family wise error rate (FWE)*

ANOVA and Multiple Comparisons

- Rejection of $H_0 : \mu_1 = \mu_2 = \cdots = \mu_K$ does not indicate where the \neq 's exist

- E.g.

$$H_A : \mu_1 = \mu_2 = \cdots = \mu_{K-1} \neq \mu_K$$

$$H_A : \mu_1 \neq \mu_2 \neq \cdots \neq \mu_{K-1} \neq \mu_K$$

- Usually we want to identify where the \neq 's exist

ANOVA

- Need multiple comparisons method to test the $\binom{K}{2}$ null hyps

$$H_0 : \mu_i = \mu_j \quad (i \neq j)$$

- Popular methods:
 - Scheffe
 - Tukey
 - Bonferroni (Sidak, Holm, Hochberg)

ANOVA: Scheffe

- For each pair of means, compute

$$t_{ij} = \frac{\bar{Y}_i - \bar{Y}_j}{\sqrt{MSE\left(\frac{1}{n_i} + \frac{1}{n_j}\right)}}$$

- Rejection region

$$C_\alpha = \left\{ t_{ij} : |t_{ij}| > \sqrt{(K - 1)F_{K-1, N-K, 1-\alpha}} \right\}$$

- Passive smoking example

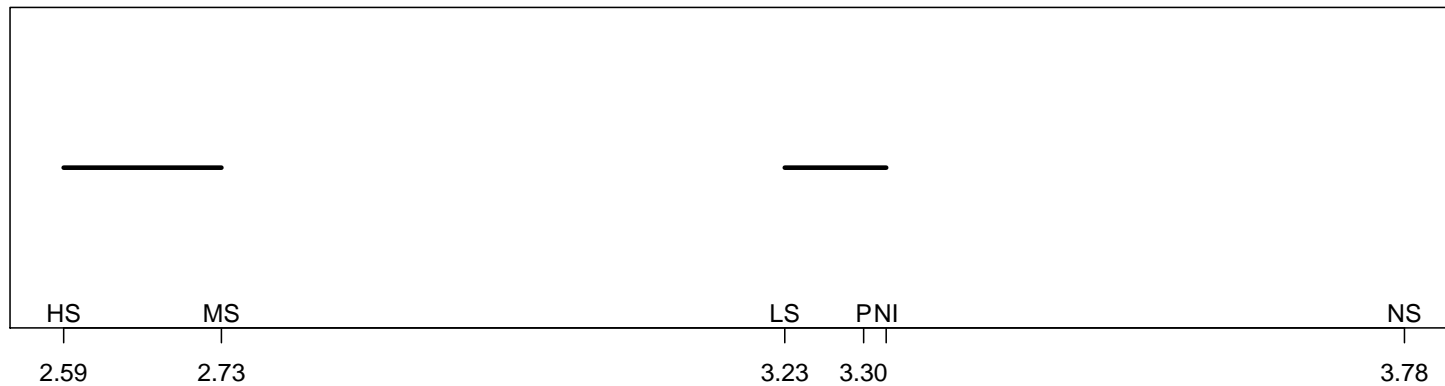
$$C_{0.05} = \{t_{ij} : |t_{ij}| > \sqrt{5F_{5, 1044, 0.95}} = \sqrt{5(2.21)} = 3.32\}$$

Scheffe: Passive smoking example

Comparison	t_{ij}	Significant
NS-PS	6.01	yes
NS-NI	3.65	yes
NS-LS	6.27	yes
NS-MS	13.17	yes
NS-HS	14.92	yes
PS-NI	-0.16	no
PS-LS	0.88	no
PS-MS	7.14	yes
PS-HS	8.90	yes
NI-LS	0.71	no
NI-MS	4.67	yes
NI-HS	5.79	yes
LS-MS	6.27	yes
LS-HS	8.03	yes
MS-HS	1.76	no

Scheffe: Example

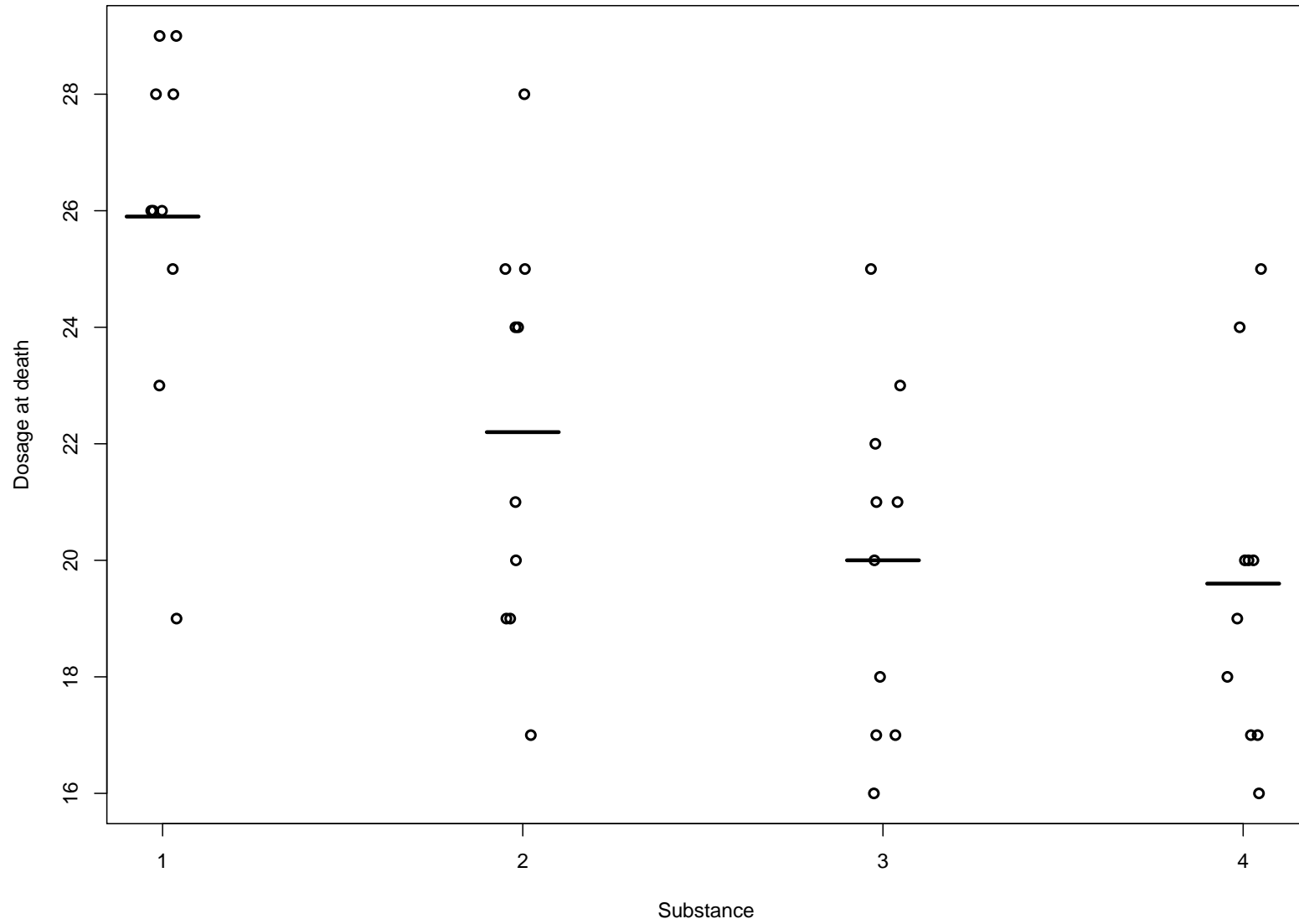
- Overall conclusions about similarities and difference among the population means using schematic diagram
- Use overbars to connect means that do not significantly differ



Scheffe: Example II

- Four cardiac substances tested for relative potencies
- For each substance, ten guinea pigs anesthetized
- Outcome: dosage at death
- Data display with group means on following slide

Scheffe: Example II



Scheffe: Example II

- Global F-test strongly rejects null of equality of four population means ($p = 0.0002$)

- Critical region

$$C_{0.05} = \{t_{ij} : |t_{ij}| > \sqrt{3F_{3,36,0.95}} = \sqrt{3 * 2.866} = 2.93\}$$

- Note denominator of t_{ij} is always $\sqrt{MSE/5} = 1.396$

- So could also write critical region in terms of *minimum significant difference*

$$C_{0.05} = \{|\bar{Y}_{i.} - \bar{Y}_{j.}| > 2.93 * 1.396 = 4.09\}$$

Scheffe: SAS

```
proc glm; class group; model dose=group; means group/scheffe;
```

Scheffe's Test for dose

NOTE: This test controls the Type I experimentwise error rate.

Alpha	0.05
Error Degrees of Freedom	36
Error Mean Square	9.747222
Critical Value of F	2.86627
Minimum Significant Difference	4.0942

Means with the same letter are not significantly different.

Scheffe Grouping	Mean	N	group
A	25.900	10	1
A			
B A	22.200	10	2
B			
B	20.000	10	3
B			
B	19.600	10	4

ANOVA: Scheffe

- For each pair of means, can compute multiplicity adjusted confidence intervals using Scheffe's method also

$$\bar{Y}_i - \bar{Y}_j \pm \sqrt{MSE \left(\frac{1}{n_i} + \frac{1}{n_j} \right)} \times \sqrt{(K - 1) F_{K-1, N-K, 1-\alpha}}$$

- The probability is at least $1 - \alpha$ these intervals simultaneously straddle the associated population mean differences
- What happens when $K = 2$?
- For cardiac substance example,

$$\bar{Y}_i - \bar{Y}_j \pm 4.09$$

Scheffe: SAS

```
proc glm; class group; model dose=group; means group/scheffe cldiff;
```

Scheffe's Test for dose

Comparisons significant at the 0.05 level are indicated by ***.

group Comparison	Difference	Simultaneous		
	Between Means	95% Confidence Limits		
1 - 2	3.700	-0.394	7.794	
1 - 3	5.900	1.806	9.994	***
1 - 4	6.300	2.206	10.394	***
2 - 1	-3.700	-7.794	0.394	
2 - 3	2.200	-1.894	6.294	
2 - 4	2.600	-1.494	6.694	
3 - 1	-5.900	-9.994	-1.806	***
3 - 2	-2.200	-6.294	1.894	
3 - 4	0.400	-3.694	4.494	
4 - 1	-6.300	-10.394	-2.206	***
4 - 2	-2.600	-6.694	1.494	
4 - 3	-0.400	-4.494	3.694	

ANOVA: Tukey

- Alternative multiple comparisons approach to Scheffe
- Critical region

$$C_\alpha = \{t_{ij} : |t_{ij}| > (q_{K,N-K,1-\alpha})/\sqrt{2}\}$$

where $q_{k,m,1-\alpha}$ is the $1 - \alpha$ quantile of the *studentized range*; see `qtukey` in R and `probmc("Range",....)` in SAS

- Multiplicity adjusted CIs

$$\bar{Y}_i - \bar{Y}_j \pm \sqrt{MSE * 2/n} \times (q_{K,N-K,1-\alpha})/\sqrt{2}$$

ANOVA: Tukey

- What is studentized range?
- Suppose Y_1, \dots, Y_k iid $N(\mu, \sigma^2)$
- Let s be an estimator of σ with m degrees of freedom,
 $s \perp Y_1, \dots, Y_k$

- Then

$$\frac{Y_{(k)} - Y_{(1)}}{s}$$

has a studentized range distribution with parameters k
and m

ANOVA: Tukey

- Cardiac substance example with $\alpha = 0.05$

$$q_{K,N-K,1-\alpha}/\sqrt{2} = q_{4,36,.95}/\sqrt{2} = 2.69$$

- Compared with Scheffe critical value (2.93), easier to reject; equivalently, Tukey confidence intervals will be narrower
- For this reason, Tukey is preferred to Scheffe in balanced designs where all pairwise comparisons are being considered
- Otherwise, use Scheffe or Bonferroni-type method

Tukey: SAS

```
proc glm; class group; model dose=group; means group/tukey cldiff;
```

Tukey's Studentized Range (HSD) Test for dose
Comparisons significant at the 0.05 level are indicated by ***.

group Comparison	Difference	Simultaneous		
	Between Means	95% Confidence Limits		
1 - 2	3.700	-0.060	7.460	
1 - 3	5.900	2.140	9.660	***
1 - 4	6.300	2.540	10.060	***
2 - 1	-3.700	-7.460	0.060	
2 - 3	2.200	-1.560	5.960	
2 - 4	2.600	-1.160	6.360	
3 - 1	-5.900	-9.660	-2.140	***
3 - 2	-2.200	-5.960	1.560	
3 - 4	0.400	-3.360	4.160	
4 - 1	-6.300	-10.060	-2.540	***
4 - 2	-2.600	-6.360	1.160	
4 - 3	-0.400	-4.160	3.360	

Tukey: R

```
> fit <- aov(dose ~ group)
> TukeyHSD(fit,"group")
```

Tukey multiple comparisons of means
95% family-wise confidence level

Fit: aov(formula = dose ~ group)

```
$group
      diff      lwr      upr      p adj
2-1 -3.7  -7.460351  0.06035128 0.0551754
3-1 -5.9  -9.660351 -2.13964872 0.0008587
4-1 -6.3 -10.060351 -2.53964872 0.0003701
3-2 -2.2  -5.960351  1.56035128 0.4048758
4-2 -2.6  -6.360351  1.16035128 0.2621133
4-3 -0.4  -4.160351  3.36035128 0.9916615
```

Bonferroni method

- Let A_1, A_2, \dots, A_n be a series of events
- Bonferroni inequality

$$\Pr(A_1 \cup A_2 \cup \dots \cup A_n) \leq \sum_{i=1}^n \Pr(A_i)$$

- Let A_i be the event we reject H_{0i} when H_{0i} is true for $i = 1, 2, \dots, n$

$$\Pr(A_i) = \alpha_i$$

Bonferroni method

- Probability of at least one type I error

$$\Pr(A_1 \cup A_2 \cup \cdots \cup A_n) \leq \sum_{i=1}^n \alpha_i$$

- If $\alpha_i = \alpha^*$ for all i ,

$$\sum_{i=1}^n \alpha_i = n\alpha^*$$

- If we want $\Pr(A_1 \cup \cdots \cup A_n) \leq \alpha$, choose $\alpha^* = \alpha/n$
- For ANOVA with K groups, there are $\binom{K}{2}$ tests; therefore

$$\alpha^* = \frac{\alpha}{\binom{K}{2}}$$

Bonferroni Method: Passive Smoking Example

- $K = 6; \binom{6}{2} = 15$
- $\alpha^* = 0.05/15 = 0.0033$
- Two-sided test,

$$\alpha^*/2 = 0.00167$$

- Rejection region

$$C_\alpha = \{|t_{ij}| > t_{N-k, 1-\alpha^*/2} = t_{1044, .9983} = 2.94\}$$

Bonferroni Method

- In SAS Proc GLM, **means group/bon;**
- In R, `pairwise.t.test(...,p.adj="bonf")`
- Sometimes called *least-significant difference (LSD)* method (Kleinbaum et al *Applied Regression Analysis* 3rd edition)
- Applicable well beyond ANOVA
- Choice of $\alpha_i = \alpha / \binom{K}{2}$ for all i is standard, but not necessary

Bonferroni Method

- Def 12.1 The significance level at which each test or comparison is carried out in an experiment is call the *per comparison error rate (PCER)*
- Bonferroni uses

$$PCER = \frac{\alpha}{\binom{K}{2}}$$

to insure

$$PEER \leq \alpha$$

- Bonferroni-type improvements (Sidak, Holm, Hochberg, Westfall and Young) available; Procs GLM and Multtest; beware dependencies in test statistics

Generalizations

- Up to this point we have considered all pairwise comparisons of means
- Other parameter combinations may be of interest
- For instance ...

Factor level means

- Single factor level mean

$$\frac{\bar{Y}_{i.} - \mu_i}{\sqrt{MSE/n_i}} \sim t_{N-K}$$

- $(1 - \alpha) \times 100\%$ CI for μ_i

$$\bar{Y}_{i.} \pm t_{N-K;1-\alpha/2} \sqrt{MSE/n_i}$$

- Testing $H_0 : \mu_i = c$ vs $H_A : \mu_i \neq c$

$$t_i = \frac{\bar{Y}_{i.} - c}{\sqrt{MSE/n_i}} \sim t_{N-K}$$

$$C_\alpha = \{t_i : |t_i| > t_{N-K;1-\alpha/2}\}$$

Linear combinations and contrasts

- *Linear combination*

$$L = \sum_{i=1}^K c_i \mu_i$$

- *Contrast* if $\sum_i c_i = 0$

- Estimator

$$\hat{L} = \sum_{i=1}^K c_i \bar{Y}_i.$$

- Compute CIs and test statistics using

$$\frac{\hat{L} - L}{\sqrt{MSE \sum_i c_i^2 / n_i}} \sim t_{N-K}$$

Conclusion

- Factor level means, i.e, μ_1, μ_2, \dots : Bonferroni
In SAS, Proc GLM/ANOVA: **means group/bon clm;**
- Pairwise comparisons: Tukey (if balanced); o/w Bonferroni if number of comparisons not too large and planned *a priori*

Conclusion

- Contrasts: Scheffe or Bonferroni

eg, multiplicity adjusted CIs for a family of contrasts of the form

$$\hat{L} \pm \sqrt{MSE \sum_i c_i^2 / n_i} \times \sqrt{(K - 1) F_{K-1, N-K; 1-\alpha}}$$

- Linear combinations: Bonferroni