

$$C = k_1\mu_1 + k_2\mu_2 + \dots + k_t\mu_t$$

$$\sum'_{i=1} k_i = 0$$

$$\underline{K} = \begin{pmatrix} k_1 \\ \vdots \\ k_t \end{pmatrix} \quad \bar{\underline{Y}} = \begin{pmatrix} \bar{Y}_{1*} \\ \vdots \\ \bar{Y}_{t*} \end{pmatrix}$$

$$\hat{C} = k_1\bar{Y}_{1*} + k_2\bar{Y}_{2*} + \dots + k_t\bar{Y}_{t*}$$

$$\hat{C} = (k_1 \dots k_t) \begin{pmatrix} \bar{Y}_{1*} \\ \vdots \\ \bar{Y}_{t*} \end{pmatrix} = \sum_{i=1}^t k_i \bar{Y}_{i*} = \underline{K} \cdot \bar{\underline{Y}} = \underline{K}^T \bar{\underline{Y}}$$

$$C_1 = \mu_1 - \mu_2$$

$$C_2 = \mu_1 - \mu_3$$

$$C_3 = \mu_2 - \mu_3$$

$$C_2 - \frac{1}{2}C_1 = (\mu_1 - \mu_3) - \frac{1}{2}(\mu_1 - \mu_2) = \frac{1}{2}\mu_1 + \frac{1}{2}\mu_2 - \mu_3 = C_4$$

$$C_4 = \frac{\mu_1 + \mu_2}{2} - \mu_3$$

A set of contrasts are linearly independent if they are orthogonal.

Consider two vectors : \underline{A} & \underline{B}

$$\underline{A} = \begin{pmatrix} a_1 \\ a_2 \\ a_3 \end{pmatrix} \quad \underline{B} = \begin{pmatrix} b_1 \\ b_2 \\ b_3 \end{pmatrix}$$

$$\underline{A} \bullet \underline{B} = \underline{A}^T \underline{B} = a_1b_1 + a_2b_2 + a_3b_3 = \|\underline{A}\| \|\underline{B}\| \cos \theta$$

- Two contrasts $\begin{cases} C = c_1\mu_1 + c_2\mu_2 + \dots + c_t\mu_t \\ D = d_1\mu_1 + d_2\mu_2 + \dots + d_t\mu_t \end{cases}$ are orthogonal if $\sum_{i=1}^t c_i d_i = 0$

$$\text{Model: } Y_{ij} = \mu + \alpha_i + \varepsilon_{ij}, \quad \varepsilon_{ij} \stackrel{iid}{\sim} N(0, \sigma_e^2)$$

$\rightarrow \hat{\mu} = \bar{Y}_{..}$ & $\hat{\alpha}_i = \bar{Y}_{i*} - \bar{Y}_{..}$ are Least Squares Estimators

- If $\hat{\mu}_i$ is the LSE of μ_i then $c_1\hat{\mu}_1 + c_2\hat{\mu}_2 + \dots + c_n\hat{\mu}_n$ is the LSE of $c_1\mu_1 + c_2\mu_2 + \dots + c_n\mu_n$

Density	Mean	Yield
Class	Levels	Values
		10 12
		20 16
DENSITY	5	10 20 30 40 50
		30 19
		40 18
		50 17

Dependent Variable: YIELD

Source	DF	Sum of Squares		Mean Square	F Value	Pr > F
		Model	Error			
	4	87.600000		21.900000	29.28	0.0001
	10		7.480000	0.748000		
Corrected Total	14	95.080000				

R-Square	C.V.	Root MSE	YIELD Mean
0.921329	5.273597	0.86487	16.4000

$$\begin{aligned} \hat{Y}_{ij} &= \hat{\beta}_0 + \hat{\beta}_1 X_1 + \hat{\beta}_2 X_2 + \dots + \hat{\beta}_k X_k \\ &= \hat{\beta}_0^* + \hat{\beta}_1^* P_{1i} + \hat{\beta}_2^* P_{2i} + \dots + \hat{\beta}_k^* P_{ki} \end{aligned}$$

$$\hat{\beta}_0^* = \bar{Y}_{..}$$

$$\hat{\beta}_1^* = \frac{\hat{C}_{lin}}{\sum_i c_{i,lin}^2} = \frac{\sum_{i=1}^t P_{1i} \bar{Y}_{i*}}{\sum_i P_{1*}^2}$$

$$\hat{\beta}_2^* = \frac{\hat{C}_{quad}}{\sum_i c_{i,quad}^2} = \frac{\sum_{i=1}^t P_{2i} \bar{Y}_{i*}}{\sum_i P_{2*}^2}$$

...

Table XI:

$$k=1: (P_{11}, P_{12}, \dots, P_{15}) = (-2, -1, 0, 1, 2)$$

$$k=2: (P_{21}, P_{22}, \dots, P_{25}) = (2, -1, -2, -1, 2)$$

...

Density	P1	P2	...	Y_hat(degree 2 poly)
10	-2	2		12.0
20	-1	-1		16.2
30	0	-2		18.4
40	1	-1		18.6
50	2	2		16.8

t=3 Treatments, r=2 replicates

Cell Means Model:

$$\begin{aligned} Y_{11} &= \mu_1 + \varepsilon_{11} \\ Y_{12} &= \mu_1 + \varepsilon_{12} \\ Y_{21} &= \mu_2 + \varepsilon_{21} \\ Y_{22} &= \mu_2 + \varepsilon_{22} \\ Y_{31} &= \mu_3 + \varepsilon_{31} \\ Y_{32} &= \mu_3 + \varepsilon_{32} \end{aligned}$$

$$\begin{bmatrix} Y_{11} \\ Y_{12} \\ Y_{21} \\ Y_{22} \\ Y_{31} \\ Y_{32} \end{bmatrix} = \begin{bmatrix} 1 & 0 & 0 \\ 1 & 0 & 0 \\ 0 & 1 & 0 \\ 0 & 1 & 0 \\ 0 & 0 & 1 \\ 0 & 0 & 1 \end{bmatrix} \begin{bmatrix} \beta_1 \\ \beta_2 \\ \beta_3 \end{bmatrix} + \begin{bmatrix} \varepsilon_{11} \\ \varepsilon_{12} \\ \varepsilon_{21} \\ \varepsilon_{22} \\ \varepsilon_{31} \\ \varepsilon_{32} \end{bmatrix}$$

$$Y = X \beta + \varepsilon$$

Treatment Effects Model:

$$\begin{aligned} Y_{11} &= \mu + \alpha_1 + \varepsilon_{11} \\ Y_{12} &= \mu + \alpha_1 + \varepsilon_{12} \\ Y_{21} &= \mu + \alpha_2 + \varepsilon_{21} \\ Y_{22} &= \mu + \alpha_2 + \varepsilon_{22} \\ Y_{31} &= \mu + \alpha_3 + \varepsilon_{31} \\ Y_{32} &= \mu + \alpha_3 + \varepsilon_{32} \end{aligned}$$

$$\begin{bmatrix} Y_{11} \\ Y_{12} \\ Y_{21} \\ Y_{22} \\ Y_{31} \\ Y_{32} \end{bmatrix} = \begin{bmatrix} 1 & 1 & 0 \\ 1 & 1 & 0 \\ 1 & 0 & 1 \\ 1 & 0 & 1 \\ 1 & 0 & 0 \\ 1 & 0 & 0 \end{bmatrix} \begin{bmatrix} \beta_0 \\ \beta_1 \\ \beta_2 \end{bmatrix} + \begin{bmatrix} \varepsilon_{11} \\ \varepsilon_{12} \\ \varepsilon_{21} \\ \varepsilon_{22} \\ \varepsilon_{31} \\ \varepsilon_{32} \end{bmatrix}$$

$$Y = X \beta + \varepsilon$$

Bonferroni's 100(1- α)% SCI (for k planned contrasts)

$$\sum_{i=1}^t c_i \bar{Y}_i \pm t_{\frac{\alpha_{EW}/2}{k}, N-t} \sqrt{s^2 \sum_{i=1}^t c_i^2 / r_i} \quad (\text{Table 2})$$

$$\sum_{i=1}^t c_i \bar{Y}_i \pm t_{\alpha_{EW}/2, k, N-t} \sqrt{s^2 \sum_{i=1}^t c_i^2 / r_i} \quad (\text{Table 5})$$

Table 2 vs. Table 5 (e.g., $k=5$ contrasts, $\alpha_{EW}=0.05$, $df=10$)

$$t_{\frac{\alpha_{EW}/2}{k}, N-t} = t_{\frac{0.05/2}{5}, 10} = t_{0.005, 10} = 3.169 \quad (\text{Table 2})$$

$$t_{\alpha_{EW}/2, k, N-t} = t_{0.05/2, 5, 10} = t_{0.025, 5, 10} = 3.17 \quad (\text{Table 5})$$

Rejection Region (Bonferroni correction)

$$|t_0| > t_{\frac{\alpha_{EW}/2}{k}, N-t} \quad (\text{Table 2})$$

$$|t_0| > t_{\alpha_{EW}/2, k, N-t} \quad (\text{Table 5})$$

$$C = \sum_{i=1}^t c_i \mu_i \quad \hat{C} = \sum_{i=1}^t c_i \bar{Y}_{i\bullet} \quad t_0 = \frac{\hat{C}}{SE(\hat{C})} = \frac{\sum_{i=1}^t c_i \bar{Y}_{i\bullet}}{\sqrt{s^2 \sum_{i=1}^t c_i^2 / r_i}}$$

100(1- α)% SCI for several C's

$$\hat{C} \pm w SE(\hat{C})$$

$$\hat{C} \pm MSD$$

$$\sum_{i=1}^t c_i \bar{Y}_{i\bullet} \pm w \sqrt{s^2 \sum_{i=1}^t c_i^2 / r_i}$$

Bonferroni's Method

* k planned contrasts

$$* MSD = t_{\frac{\alpha_{EW}/2}{k}, N-t} \sqrt{s^2 \sum_{i=1}^t c_i^2 / r_i}$$

$$\hat{C} = \sum_{i=1}^t c_i \bar{Y}_{i\bullet}$$

Scheffé's Method

* all possible contrasts

$$* MSD = \sqrt{(t-1)F_{\alpha, t-1, N-t}} \sqrt{s^2 \sum_{i=1}^t c_i^2 / r_i}$$

$$\hat{C} = \sum_{i=1}^t c_i \bar{Y}_{i\bullet}$$

Dunnett's Method

* comparing k treatments with a control ($r_i=r$)

$$* MSD = d_{\alpha, k, N-t} \sqrt{2s^2 / r}$$

* Reject Ho if $|\bar{Y}_{i\bullet} - \bar{Y}_{Control}| > MSD$

* Table VI for 1-sided and 2-sided critical values for $d_{\alpha, k, N-t}$

$$\hat{C} = \bar{Y}_{i\bullet} - \bar{Y}_{Control}$$

Tukey's HSD Method

* all pairwise comparisons ($r_i=r$)

$$* MSD = q_{\alpha, k, N-t} \sqrt{s^2 / r}$$

* Reject Ho if $|\bar{Y}_{i\bullet} - \bar{Y}_{j\bullet}| > MSD$

$$\hat{C} = \bar{Y}_{i\bullet} - \bar{Y}_{j\bullet}$$

* Unequal replications \rightarrow Tukey-Kramer Method with $MSD = q_{\alpha, k, N-t} \sqrt{\frac{s^2}{2} \left(\frac{1}{r_i} + \frac{1}{r_j} \right)}$

Fisher's LSD Method (Protected T-tests)

* If the omnibus F-test from the ANOVA is not significant then STOP
Otherwise, test each pair with an ordinary T-test

$$* \text{Reject } H_0 \text{ if } \frac{\bar{Y}_{i\bullet} - \bar{Y}_{j\bullet}}{\sqrt{s^2 \left(\frac{1}{r_i} + \frac{1}{r_j} \right)}} > t_{\alpha/2, N-t}$$

* Controls α_{EW} in the weak sense through the overall F-test

Student Newmann Keuls (SNK) Method

* One of several multiple range tests

* "step-down procedure" since you are looking at different ranges of means

Ryan Einot Gabriel Welsch (REGWQ) Method

* Another multiple range test

Multiple Comparisons with Best (MCB) Method

* Similar to Dunnett's procedure

uses 1-sided critical values for $d_{\alpha, k, N-t}$

* less commonly used in biostatistics, more common in engineering applications