

Source of Variation	df	SS	MS	F _{Obs}	T=total	M=Model	E=Error
Intercept	1	SSI	SSI	SSI/MSE	Source	Formula	Alternate
Model(uncorrected)	q	USS(model)	USSM/q	(USSM/q)/MSE	CSS(M)	$\sum \hat{y}_i^2 - N\bar{y}^2 = \sum (\hat{y}_i - \bar{y})^2$	USS(M) - SSI
Model(corrected)	q-1	CSS(model)	CSSM/q-1	(CSSM/q-1)/MSE	CSS(T)	$\sum y_i^2 - N\bar{y}^2 = \sum (y_i - \bar{y})^2$	USS(T) - SSI & CSS(M) + SSE
Error(residual)	N-q	SSE	SSE/(N-q)=MSE	-	CSS(E)/USS(E)/SSE	CSS(T) - CSS(M)	USS(T) - USS(M) = USS(E)
Total(uncorrected)	N	USS(total)	-	-	SSE	$\sum (\hat{y} - y_i)^2$	$\sum \hat{\epsilon}^2$
Total(corrected)	N-1	CSS(total)	-	-	USS(M)	$\sum \hat{y}_i^2$	USS(T) - SSE
					USS(T)	$\sum y_i^2$	USS(M) + SSE

(8 points) Calculate the corrected R^2_c , interpret its value, and test the hypothesis that its corresponding population value is zero, that is, $H_0 : \rho^2_c = 0$.

Solution: Since $\bar{y} = 3.4$, we have $CSS(regression) = \sum_i \hat{y}_i^2 - 5\bar{y}^2 = 11.2$
 $CSS(total) = \sum_i y_i^2 - 5\bar{y}^2 = 17.2$ and corrected $R^2_c = 11.2/17.2 = 0.65$.

$H_0 : \rho^2_c = 0$ is equivalent to $H_0 : \beta_1 = \beta_2 = 0$. Thus we have $F - test = \frac{11.2/2}{\frac{5.0}{2}} = 2.25 \sim F_{2,2}$.

Two-way ANOVA coding schemes

Reference Cell: The total number of columns of **X**: 1 col of 1's for reference cell mean, (a-1) for factor A, (b-1) for factor B and (a-1)(b-1) for interaction.

Cell Means:

$\begin{bmatrix} 1 & 0 & 0 \\ 1 & 1 & 0 \\ 1 & 0 & 1 \\ 0 & 0 & 1 \end{bmatrix}$

Classic ANOVA: The total number of columns of **X** is 1 + a + b + a*b

$\begin{bmatrix} 1 & 1 & 0 & 0 \\ 1 & 1 & 0 & 0 \\ 1 & 0 & 0 & 1 \\ 1 & 0 & 0 & 1 \end{bmatrix}$

Effect: same as reference cell but you replace the zeros for the reference row and range with -1s

$\begin{bmatrix} 1 & -1 & -1 \\ 1 & 1 & 0 \\ 1 & 0 & 1 \end{bmatrix}$

Cell Means Per Coding 2 drugs A and B (α) and 3 dosages 1, 2, and 3 (β). The model is $y_{ijk} = \mu + \alpha_1 I(drugA) + \beta_1 I(dose1) + \beta_2 I(dose2) + e_{ijk}$

Drug	Dose	Mean	With interactions
A	1	$\mu + \alpha_1 + \beta_1$	+ γ_{A1}
A	2	$\mu + \alpha_1 + \beta_2$	+ γ_{A2}
A	3	$\mu + \alpha_1$	+ γ_{A3}
B	1	$\mu + \beta_1$	
B	2	$\mu + \beta_2$	
B	3	μ	

Cell Means For 2-way ANOVA can only have max a*b terms, which are the interaction terms.

Drug	Dose	Mean
A	1	γ_{A1}
A	2	γ_{A2}
A	3	γ_{A2}
B	1	γ_{B1}
B	2	γ_{B2}
B	3	γ_{B3}

$$(X'X)^{-1} = \begin{bmatrix} .308 & -.06 & -.017 \\ -.06 & -.025 & -.004 \\ -.017 & -.004 & .006 \end{bmatrix} X'y = \begin{bmatrix} 405 \\ 1402 \\ 2350 \end{bmatrix}$$

Source	DF	SS	MS	Fval	P > F
Model	2	79	239.5	349.6	<.001
Error	97	11	.113		
Ctotal	99	90			
param	est	se	tval	p > t	
x0	.67	.187	3.58	.009	
x1	1.35	.053	25.47	< .001	
x2	1.61	.026	61.81	< .001	

2) c) Test $H_0 : \beta_1 = 1$

t-test= $\frac{\hat{\beta}_1 - 1}{se(\hat{\beta}_1)} = 6.6 \sim t_{97}$ $6.6 > 1.96$ reject H_0

d) Test $H_0 : \beta_1 = \beta_2 = 1$

$B_1 = 1 \quad B_2 = 1$

$C = \begin{bmatrix} 0 & 1 & 0 \\ 0 & 0 & 1 \end{bmatrix} \theta_0 = \begin{bmatrix} 1 \\ 1 \end{bmatrix} H_0 : C\theta = \theta_0 \quad \hat{\theta} = \begin{bmatrix} 1.35 \\ 1.61 \end{bmatrix}$

calc M \quad calc M^{-1}

F-test= $\frac{43.83}{113} = 387.8 \sim F_{2,97}$

e) 95% CI of $\beta_1 + \beta_2$ (θ)

$\hat{\theta} = \hat{\beta}_1 + \hat{\beta}_2 = 2.96 \quad C = [0, 1, 1]$

$se(\hat{\theta}) = \sigma^2_{\hat{\theta}} = \sqrt{.0026}$

$2.96 \pm 1.96\sqrt{.0026} = [2.86, 3.06]$

f) Transform x_1, x_2 to $z_1 = x_1 - 2, z_2 = x_2 - 4$

Refit with $y^* = \beta_0^* + \beta_1^* z_1 + \beta_2^* z_2$

$= (\beta_0^* - 2\beta_1^* - 4\beta_2^*) + \beta_1^* x_1 + \beta_2^* x_2$

$\beta_0^* = \beta_0 + 2\beta_1 + 4\beta_2 = 9.8 \quad C = [1, 2, 4]$

$\sigma^2 = \hat{\sigma}^2 C(X'X)^{-1}C^{-1} = .085 \quad t = 115.3$

3) f) is $H_0 : \beta_0 - \beta_2 = 0$ and $\beta_1 + 2\beta_2 = 2$ and $2\beta_0 + \beta_1 = 2$ testable? Reduce to ETH?

$$y = \begin{bmatrix} 0 \\ 2 \\ 6 \\ 10 \end{bmatrix} X = \begin{bmatrix} 1 & 6 & 11 \\ 1 & 7 & 13 \\ 1 & 8 & 15 \\ 1 & 9 & 17 \end{bmatrix} \beta = \begin{bmatrix} \beta_0 \\ \beta_1 \\ \beta_2 \end{bmatrix}$$

$$C = \begin{bmatrix} 1 & 0 & -1 \\ 0 & 1 & 2 \\ 2 & 1 & 0 \end{bmatrix} \text{ reduces to } \begin{bmatrix} 1 & 0 & -1 \\ 0 & 1 & 2 \\ 0 & 0 & 0 \end{bmatrix} \text{ 2pivots } \theta_0 = \begin{bmatrix} 0 \\ 2 \end{bmatrix}$$

not FR, not testable, reduces to C^* testable

$$C^* = \begin{bmatrix} 1 & 0 & -1 \\ 0 & 1 & 2 \end{bmatrix} \theta_0^* = \begin{bmatrix} 0 \\ 2 \end{bmatrix} H_0 : \begin{bmatrix} \beta_0 - \beta_2 = 0 \\ \beta_1 + 2\beta_2 = 2 \end{bmatrix}$$

$$\mathbf{1) a) } x = \begin{bmatrix} x_1 \\ x_2 \\ x_3 \end{bmatrix} \sim N(0, \Sigma) \quad \Sigma = \begin{bmatrix} 1 & 0 & .6 \\ 0 & 1 & .5 \\ .6 & .5 & 1 \end{bmatrix}$$

$$c = \begin{bmatrix} 2 & 1 & -1 \end{bmatrix}$$

$$2x_1 + x_2 - x_3 \sim N(c\mu, c\Sigma c') = N(0, 2.6)$$

$$\mathbf{b) } Cov(x_1 - x_2, 2x_2 + x_3) = c_1 \Sigma c_2' = -1.9$$

$$c_1 = \begin{bmatrix} 1 & -1 & 0 \end{bmatrix} \quad c_2 = \begin{bmatrix} 0 & 2 & 1 \end{bmatrix}$$

$$\mathbf{2011 1) } \hat{\beta} = \begin{bmatrix} 24.877 \\ -.231 \\ -.185 \end{bmatrix} \quad \hat{\sigma}^2 = 5.59 \quad dfE = 197$$

$$diag((X'X)^{-1}) = [.03752, .0017, .00056]$$

$$y = \beta_0 + \beta_1 x_1 + \beta_2 x_2 + \epsilon$$

$$se\beta_1 = .032 \quad se\beta_2 = .05605$$

$$95\% \text{ CI } x_1 = 2 \quad x_2 = 6 \quad C = \begin{bmatrix} 1 & 2 & 6 \end{bmatrix}$$

$$\hat{\beta}^* \pm 1.96 * se(\hat{\beta}^*) \quad \hat{\beta}^* = \hat{\beta}_0 + 2\hat{\beta}_1 + 6\hat{\beta}_2 = 23.3$$

$$se = \sqrt{Var(\hat{\beta}^*)} = \sqrt{\hat{\sigma}^2 M} = .1946$$

$$\text{Test } H_0 : \beta_1 = 3\beta_2$$

$$\theta = \beta_1 - 3\beta_2 \quad H_0 : \theta = 0 \quad C = \begin{bmatrix} 0 & 1 & -3 \end{bmatrix} \quad t =$$

$$.327/.1868 = 1.75 \sim t_{97} \approx Z = 1.96 \quad \text{FTR } H_0$$

$$\text{Center } x_1 \quad x_2 \text{ at means } 1.5 \quad z_1 = x_1 - 1$$

$$z_2 = x_2 - 5 \quad y^* = \beta_0^* + \beta_1^* z_1 + \beta_2^* z_2$$

$$\beta_0^* + \beta_1^* (x_1 - 1) + \beta_2^* (x_2 - 5)$$

$$y^* = (\beta_0^* - \beta_1^* - 5\beta_2^*) + \beta_1^* x_1 + \beta_2^* x_2$$

$$\beta_0^* = \beta_0 + \beta_1 + 5\beta_2 = -.9116 \quad \beta_1^* = \beta_1 \quad \beta_2^* = \beta_2$$

$$C = \begin{bmatrix} 1 & 1 & 5 \end{bmatrix} \quad se(\beta_0^*) = \sqrt{M * \hat{\sigma}^2} = .254$$

2) 4 dose levels (1,2,3,4) 2 methods (oj, aa)

n=800 ref cell \Rightarrow ref group: aa dose 1

$$y_i = \mu + \alpha_1 x_{1i} + \alpha_2 x_{2i} + \alpha_3 x_{3i} + \beta x_{4i} + \epsilon_i$$

dummy variables:

$$x_{1i} = \begin{cases} 1 & \text{if dose}=2 \\ 0 & \text{otherwise} \end{cases} \quad \text{all xs defined similarly}$$

$$x_{4i} = I(\text{method} = oj)$$

cell mean of each group

Method	Dose	Mean
oj	1	$\mu + \beta$
oj	2	$\mu + \alpha_1 + \beta$
oj	3	$\mu + \alpha_2 + \beta$
oj	4	$\mu + \alpha_3 + \beta$
aa	1	μ
aa	2	$\mu + \alpha_1$
aa	3	$\mu + \alpha_2$
aa	4	$\mu + \alpha_3$

add interaction btwn method, dose $\Rightarrow \theta : 8 \times 1$

let μ_o and μ_a be overall means for the two

methods

write them for models w/ and w/o interaction

Derive 2 C matrices for testing $H_0 : \mu_o = 2\mu_a$

under the 2 models

$$\text{w/o: } \mu_o = \mu + \frac{\alpha_1 + \alpha_2 + \alpha_3}{4} + \beta$$

$$\mu_a = \mu + \frac{\alpha_1 + \alpha_2 + \alpha_3}{4}$$

$$H_0 : \mu_o = 2\mu_a \Leftrightarrow \mu + \frac{\alpha_1 + \alpha_2 + \alpha_3}{4} - \beta = 0$$

$$C = (1 \ 1/4 \ 1/4 \ 1/41) \text{ for } \theta = (\mu, \alpha_1, \alpha_2, \alpha_3, \beta)$$

$$\text{w/: } H_0 : \mu_o = 2\mu_a \Leftrightarrow$$

$$\mu + \frac{\alpha_1 + \alpha_2 + \alpha_3}{4} - \frac{\gamma_{[11]} + \gamma_{12} + \gamma_{13}}{4} - \beta = 0$$

$$C = (1 \ 1/4 \ 1/4 \ 1/411/4 \ -1/4 \ -1/4) \text{ for}$$

$$\theta = (\mu, \alpha_1, \alpha_2, \alpha_3, \beta, ga_{11}, \gamma_{12}, \gamma_{13})$$

treat dose as a continuous variable and fit

model with additive effects of method and dose level with no interaction

nested w/in

$$y_i = \mu + \alpha_1 x_{1i} + \alpha_2 x_{2i} + \alpha_3 x_{3i} + \beta x_{4i} + \epsilon_i? \text{ yes}$$

write down H_0 for comparing the 2 models and

derive C matrix. df of F test under H_0 ? Let

$$\alpha_2 = 2\alpha_1, \alpha_3 = 3\alpha_1 \text{ (treating dose as}$$

continuous)

$$C = \begin{bmatrix} 0 & -2 & 1 & 0 & 0 \\ 0 & -3 & 0 & 1 & 0 \end{bmatrix}$$

$$\text{assuming } \theta = (\mu, \alpha_1, \alpha_2, \alpha_3, \beta)'$$

$$\text{df}=2, \ 7955$$

2010 1) n = 1000 drugs = A,B doses=1,2,3,4,5

ref cell $y = \mu + \alpha_i + \beta_j + \epsilon$ (w/out interaction)

Drug B, Dose 5 ref

α_1 models effect of drug A

Source	DF	Type I SS	MS	F-value
x_i	1	Type I SS	T1SS/df = T1SS	MS/MSE(model)
$\sum_{i=1}^{q-1} T1SS_{x_i} = CSS(M)$				

β_j models effect for dose j

dims: $y = 1000 \times 1 \quad X = 1000 \times 6 \quad \beta = 6 \times 1$

$$\epsilon = 1000x1 \quad \epsilon \sim N(0, \sigma^2 I)$$

Cell mean for each combo of drug and dose:

Drug	Dose	Mean
A	1	$\mu + \alpha_1 + \beta_1$
A	2	$\mu + \alpha_1 + \beta_2$
A	3	$\mu + \alpha_1 + \beta_3$
A	4	$\mu + \alpha_1 + \beta_4$
A	5	$\mu + \alpha_1$
B	1	$\mu + \beta_1$
B	2	$\mu + \beta_2$
B	3	$\mu + \beta_3$
B	4	$\mu + \beta_4$
B	5	μ

for ref cell model p=6 df model = 6 - 1 = 5

with interaction $y_{ijk} = \mu + \alpha_i + \beta_j + \gamma_{ij} + \epsilon$

$\mu + \alpha_i + \beta_4 + \beta_3 + \beta_2 + \beta_1 + \gamma_{14} + \gamma_{13} + \gamma_{12} + \gamma_{11} + \epsilon$

dim ref model: $\beta = 10 \times 1$ everything else same

Drug	Dose	Mean
A	1	$\mu + \alpha_1 + \beta_1 + \gamma_{11}$
A	2	$\mu + \alpha_1 + \beta_2 + \gamma_{12}$
A	3	$\mu + \alpha_1 + \beta_3 + \gamma_{13}$
A	4	$\mu + \alpha_1 + \beta_4 + \gamma_{14}$
A	5	$\mu + \alpha_1$
B	1	$\mu + \beta_1$
B	2	$\mu + \beta_2$
B	3	$\mu + \beta_3$
B	4	$\mu + \beta_4$
B	5	μ

γ_{11} is the diff btwn the diff of dose1 and dose5

given drug A and b respectively

$$= [E(y|A, 1) - E(y|A, 5)] - [E(y|B, 1) - E(y|B, 5)]$$

Let μ_A and μ_B be the overall mean values of

cholesterol level for drug A and B

$$\mu_A = [(\mu + \alpha_1 + \beta_1 + \gamma_{11}) + (\mu + \alpha_1 + \beta_2 + \gamma_{12}) +$$

$$(\mu + \alpha_1 + \beta_3 + \gamma_{13}) + (\mu + \alpha_1 + \beta_4 + \gamma_{14}) + (\mu + \alpha_1)]/5$$

$$\mu_B = [(\mu + \beta_1) + (\mu + \beta_2) + (\mu + \beta_3) + (\mu + \beta_4) + \mu]/5$$

$$H_0 : \mu_A = \mu_B \Leftrightarrow H_0 \frac{\alpha_1 + \gamma_{11} + \gamma_{12} + \gamma_{13} + \gamma_{14}}{5} = 0$$

2) full model in each cell, dose is treated as

interval variable

$$\hat{y} = 9.82 + 3.55 * I(drug = A) + 2.14 dose + .296 age$$

$$+ 2.21 I(drug = A) * dose + .3 I(drug = A) * age$$

fitted model when drug B is used (ref level)

$$\hat{y} = 98.82 + 2.145 dose + 2.96 age$$

Write down fitted model when drug A is used

$$\hat{y} = 98.82 + 3.55 * 1 + 2.14 dose + .296 age +$$

$$2.21 * 1 * dose + .3 * 1 * age$$

$$\Rightarrow \hat{y} = 102.37 + 4.35 dose + .596 age$$