

Chapter 12. Analysis of variance

Ch 11:	$I=2$	indep. samples	paired samples
Ch 12:	$I \geq 2$	one-way layout	two-way layout

1. One-way layout

One factor (factor A) with I levels (I treatments)

I independent IID samples $(Y_{i1}, \dots, Y_{iJ}), i = 1, \dots, I$

H_0 : all I treatments have the same effect

H_1 : there are systematic differences

Ex 1: seven labs

p. 444: data and boxplots, $I = 7, J = 10$

Normal theory model

Normally distributed observation $Y_{ij} \sim N(\mu_i, \sigma^2)$

$Y_{ij} = \mu + \alpha_i + \epsilon_{ij}, \sum \alpha_i = 0, \epsilon_{ij} \sim N(0, \sigma^2)$

obs = overall mean + differential effect + error

Maximum likelihood estimates

$$\begin{aligned} \hat{\mu} &= \bar{Y}_{..} && \text{pooled sample mean } \bar{Y}_{..} \\ \hat{\alpha}_i &= \bar{Y}_{i.} - \bar{Y}_{..} && \text{sample means } \bar{Y}_{1.}, \dots, \bar{Y}_{I.} \end{aligned}$$

Sums of squares: $SS_{\text{TOT}} = SS_{\text{A}} + SS_{\text{E}}$

$SS_{\text{TOT}} = \sum \sum (Y_{ij} - \bar{Y}_{..})^2$ total sum of squares

$SS_{\text{A}} = J \sum \hat{\alpha}_i^2$ between samples (factor A) sum of sq

$SS_{\text{E}} = \sum \sum \hat{\epsilon}_{ij}^2$ within samples (error) sum of squares
residuals $\hat{\epsilon}_{ij} = Y_{ij} - \bar{Y}_{i.}$

Degrees of freedom and mean squares:

$$df_A = I - 1, MS_A = \frac{SS_A}{df_A}, E(MS_A) = \sigma^2 + \frac{J}{I-1} \sum \alpha_i^2$$

$$df_E = I(J - 1), MS_E = \frac{SS_E}{df_E}, E(MS_E) = \sigma^2$$

$$\text{total df} = IJ - 1$$

Pooled sample variance $s_p^2 = MS_E$
 an unbiased estimate of σ^2

F-test

$$H_0 : \alpha_1 = \dots = \alpha_I = 0 \text{ against}$$

$$H_1 : \alpha_u \neq \alpha_v \text{ for some } (u, v)$$

Reject H_0 for large values of $F = \frac{MS_A}{MS_E}$

null distribution of F is $F_{I-1, I(J-1)}$

If $Z_i \sim N(0,1)$ indep., then $\frac{(Z_1^2 + \dots + Z_m^2)/m}{(Z_{m+1}^2 + \dots + Z_{m+n}^2)/n} \sim F_{m,n}$

Ex 1: seven labs

normal probability plot of residuals $\hat{\epsilon}_{ij}$, p. 450

Anova-1 table

Source	df	SS	MS	F	P-value
Labs	6	.125	.0210	5.66	.0001
Error	63	.231	.0037		
Total	69	.356			

Multiple comparisons: $\binom{7}{2} = 21$ pairwise comparisons

Lab	1	3	7	2	5	6	4
Mean	4.062	4.003	3.998	3.997	3.957	3.955	3.920

Bonferroni method

Take α as an overall level in k independent tests
if each done at significance level α/k

Proof: given that H_0 is true

number of significant results in k tests $X \sim \text{Bin}(k, \frac{\alpha}{k})$

$$P(X \geq 1 | H_0) = 1 - (1 - \frac{\alpha}{k})^k \approx \alpha$$

Warning: $k = \binom{I}{2}$ pairwise comparisons are not independent as required by the Bonferroni method

Simultaneous $100(1 - \alpha)\%$ CI

for $\binom{I}{2}$ pairwise differences $(\alpha_u - \alpha_v)$

$$(\bar{Y}_{u.} - \bar{Y}_{v.}) \pm t_{I(J-1)}(\frac{\alpha}{I(I-1)}) \cdot s_p \sqrt{\frac{2}{J}}$$

Flexibility: works for different sample sizes as well

replace $\sqrt{\frac{2}{J}}$ by $\sqrt{\frac{1}{J_u} + \frac{1}{J_v}}$

Ex 1: seven labs

95% CI for one difference $(\alpha_u - \alpha_v)$

$$(\bar{Y}_{u.} - \bar{Y}_{v.}) \pm t_{63}(0.025) \cdot \frac{s_p}{\sqrt{5}} = (\bar{Y}_{u.} - \bar{Y}_{v.}) \pm 0.055$$

where $t_{63}(0.025) = 2.00$, $s_p = \sqrt{0.0037} = 0.061$

Simultaneous 95% CI for $(\alpha_u - \alpha_v)$ by Bonferroni method

$$(\bar{Y}_{u.} - \bar{Y}_{v.}) \pm t_{63}(\frac{.05}{42}) \cdot \frac{s_p}{\sqrt{5}} = (\bar{Y}_{u.} - \bar{Y}_{v.}) \pm 0.086$$

Labs	1-4	1-6	1-5	3-4	7-4	2-4	1-2
Diff	0.142	0.107	0.105	0.083	0.078	0.077	0.065

significant differences between labs (1,4), (1,5), (1,6)

Tukey method

If the sample sizes are equal J , then

$\bar{Y}_i. \sim N(\mu + \alpha_i, \frac{\sigma^2}{J})$ are independent and

$\frac{\sqrt{J}}{s_p} \max_{u,v} |\bar{Y}_u. - \bar{Y}_v. - (\alpha_u - \alpha_v)| \sim \text{SR}(I, I(J - 1))$

Studentized range distribution $\text{SR}(t, \nu)$

t = number of samples, ν = df (the variance estimate)

Table 6, p. A14-A19

$q_{t,\nu}(\alpha) = 100(1 - \alpha)\%$ -percentile of $\text{SR}(t, \nu)$

$$\boxed{\text{Simultaneous CI} = (\bar{Y}_u. - \bar{Y}_v.) \pm q_{I, I(J-1)}(\alpha) \cdot \frac{s_p}{\sqrt{J}}}$$

Ex 1: seven labs

$$(\bar{Y}_u. - \bar{Y}_v.) \pm q_{7,63}(0.05) \cdot \frac{0.061}{\sqrt{10}} = (\bar{Y}_u. - \bar{Y}_v.) \pm 0.083$$

where $q_{7,60}(0.05) = 4.31$

significant differences (1,4), (1,5), (1,6), (3,4)

Kruskal-Wallis test

Nonparametric test for

H_0 : all observations are equal in distribution

when ϵ_{ij} are non-normal

Pooled sample size $N = J_1 + \dots + J_I$

pooled sample ranking: R_{ij} = ranks of Y_{ij}

$$\sum_{i,j} R_{ij} = \frac{N(N+1)}{2}, \bar{R}_{..} = \frac{N+1}{2}$$

$$\boxed{\text{Test statistic } K = \frac{12}{N \cdot (N+1)} \sum_{i=1}^I J_i \cdot (\bar{R}_i. - \frac{N+1}{2})^2}$$

Reject H_0 for large K

approximate null distribution $K \stackrel{a}{\sim} \chi_{I-1}^2$

Ex 1: seven labs

Actual measurements replaced by their ranks $1 \div 70$

Labs	1	2	3	4	5	6	7
	70	4	35	6	46	48	38
	63	3	45	7	21	5	50
	53	65	40	13	47	22	52
	64	69	41	20	8	28	58
	59	66	57	16	14	37	68
	54	39	32	26	42	2	1
	43	44	51	17	9	31	15
	61	56	25	11	10	34	23
	67	24	29	27	33	49	60
	55	19	30	12	36	18	62
Means	58.9	38.9	38.5	15.5	26.6	27.4	42.7

$K = 28.17$, $df = 6$, P-value ≈ 0.0001

2. Two-way layout

Two factors: factor A with I levels (levels = rows)

factor B with J levels (levels = columns)

Data $\{Y_{ijk}, 1 \leq i \leq I, 1 \leq j \leq J, 1 \leq k \leq K\}$

$I \cdot J$ cells with K observations per cell

total number of observations = $I \cdot J \cdot K$

Normal theory model

$$Y_{ijk} = \mu + \alpha_i + \beta_j + \delta_{ij} + \epsilon_{ijk}$$

grand mean + main effects + interaction

independent random errors $\epsilon_{ijk} \sim N(0, \sigma^2)$

Parameter constraints

$$\sum \alpha_i = 0, \text{df}_A = I - 1 \quad \sum \beta_j = 0, \text{df}_B = J - 1$$

$$\sum \delta_{i1} = 0, \dots, \sum \delta_{iJ} = 0 \quad \sum \delta_{1j} = 0, \dots, \sum \delta_{Ij} = 0$$

$$\text{df}_{AB} = IJ - I - (J - 1) = (I - 1)(J - 1)$$

MLE

$$\hat{\mu} = \bar{Y}_{...} \quad \hat{\alpha}_i = \bar{Y}_{i..} - \bar{Y}_{...} \quad \hat{\beta}_j = \bar{Y}_{.j.} - \bar{Y}_{...}$$

$$\hat{\delta}_{ij} = \bar{Y}_{ij.} - \bar{Y}_{...} - \hat{\alpha}_i - \hat{\beta}_j$$

Ex 2: iron retention

factor A: $I = 2$ different iron forms

factor B: $J = 3$ dosage levels, $K = 18$ obs per cell

Raw data, p. 396: $X_{ijk} = \%$ of iron retained

transformed data $Y_{ijk} = \ln(X_{ijk})$

p. 462-463: boxplots and plots of cell SDs vs cell means

MLE for the transformed data

$$\bar{Y}_{...} = 1.92 \quad \|\bar{Y}_{ij.}\| = \begin{pmatrix} 1.16 & 1.90 & 2.28 \\ 1.68 & 2.09 & 2.40 \end{pmatrix}$$

$$\hat{\alpha}_1 = -0.14, \hat{\alpha}_2 = 0.14$$

$$\hat{\beta}_1 = -0.50, \hat{\beta}_2 = 0.08, \hat{\beta}_3 = 0.42$$

$$\|\hat{\delta}_{ij}\| = \begin{pmatrix} -0.12 & 0.04 & 0.08 \\ 0.12 & -0.04 & -0.08 \end{pmatrix}$$

Sums of squares

$$SS_{\text{TOT}} = SS_A + SS_B + SS_{AB} + SS_E$$

$$SS_{\text{TOT}} = \sum_i \sum_j \sum_k (Y_{ijk} - \bar{Y}_{...})^2$$

$$SS_A = JK \sum_i \hat{\alpha}_i^2$$

$$SS_B = IK \sum_j \hat{\beta}_j^2$$

$$SS_{AB} = K \sum_i \sum_j \hat{\delta}_{ij}^2$$

$$SS_E = \sum_i \sum_j \sum_k (Y_{ijk} - \bar{Y}_{ij.})^2, \text{df}_E = IJ(K - 1)$$

Mean squares

$$MS_A = \frac{SS_A}{\text{df}_A} \quad E(MS_A) = \sigma^2 + \frac{JK}{I-1} \sum_i \alpha_i^2$$

$$MS_B = \frac{SS_B}{\text{df}_B} \quad E(MS_B) = \sigma^2 + \frac{IK}{J-1} \sum_j \beta_j^2$$

$$MS_{AB} = \frac{SS_{AB}}{\text{df}_{AB}} \quad E(MS_{AB}) = \sigma^2 + \frac{K}{(I-1)(J-1)} \sum_i \sum_j \delta_{ij}^2$$

$$MS_E = \frac{SS_E}{\text{df}_E} \quad E(MS_E) = \sigma^2$$

Three F -tests

$$H_A: \alpha_1 = \dots = \alpha_I = 0 \quad F_A = \frac{MS_A}{MS_E} \sim F_{\text{df}_A, \text{df}_E}$$

$$H_B: \beta_1 = \dots = \beta_J = 0 \quad F_B = \frac{MS_B}{MS_E} \sim F_{\text{df}_B, \text{df}_E}$$

$$H_{AB}: \text{all } \delta_{ij} = 0 \quad F_{AB} = \frac{MS_{AB}}{MS_E} \sim F_{\text{df}_{AB}, \text{df}_E}$$

Reject null hypothesis

for large values of the respective test statistic F

Inspect normal probability plot

for residuals $\hat{\epsilon}_{ijk} = Y_{ijk} - \bar{Y}_{ij.}$

Ex 2: iron retention

Anova-2 table for the transformed iron retention data

Source	df	SS	MS	F	P
Iron form	1	2.074	2.074	5.99	0.017
Dosage	2	15.588	7.794	22.53	0.000
Interaction	2	0.810	0.405	1.17	0.315
Error	102	35.296	0.346		
Total	107	53.768			

Significant effect due to iron form

log scale difference $\hat{\alpha}_2 - \hat{\alpha}_1 = \bar{Y}_{2..} - \bar{Y}_{1..} = 0.28$

multiplicative effect of $e^{0.28} = 1.32$ on a linear scale

interaction is not significant

Additive model

If $K = 1$ we cannot estimate interaction

$$Y_{ij} = \mu + \alpha_i + \beta_j + \epsilon_{ij}$$

$$\hat{\mu} = \bar{Y}_{..} \quad \hat{\alpha}_i = \bar{Y}_{i.} - \bar{Y}_{..} \quad \hat{\beta}_j = \bar{Y}_{.j} - \bar{Y}_{..}$$

$$\hat{\epsilon}_{ij} = Y_{ij} - \bar{Y}_{..} - \hat{\alpha}_i - \hat{\beta}_j$$

Sums of squares

$$SS_{\text{TOT}} = SS_{\text{A}} + SS_{\text{B}} + SS_{\text{E}}$$

$$SS_{\text{A}} = J \sum_i \hat{\alpha}_i^2 \quad df_{\text{A}} = I - 1$$

$$SS_{\text{B}} = I \sum_j \hat{\beta}_j^2 \quad df_{\text{B}} = J - 1$$

$$SS_{\text{E}} = \sum_i \sum_j \hat{\epsilon}_{ij}^2 \quad df_{\text{E}} = (I - 1)(J - 1)$$

If $\epsilon_{ij} \sim N(0, \sigma^2)$, apply F -tests to test H_{A} and H_{B}

Randomized block design

Experimental design with I treatments

randomly assigned within each of J blocks

To test $H_0: \alpha_1 = \dots = \alpha_I = 0$, no treatment effects

use two-way layout ANOVA

The block effect is anticipated and is not of major interest

Block	Treatments	Observation
Homogen. plot of land divided into I subplots	I fertilizers	The yield on the subplot (i, j)
A four-wheel car	4 tire types	tire's life-length
A litter of I animals	I diets	the weight gain

Ex. 3: experiment on itching

Data, p. 467

$I = 7$ treatments to relieve itching

$J = 10$ blocks (male volunteers aged 20-30)

$K = 1$ observation per cell

Y_{ij} = the duration of the itching in seconds

Boxplots and normal prob. plot of residuals, p. 468-469

placebo cell variance: different response to placebo

Anova-2 table:

Source	df	SS	MS	F	P
Drugs	6	53013	8835	2.85	0.018
Subjects	9	103280	11476	3.71	0.001
Error	54	167130	3096		
Total	69	323422			

$$\boxed{\text{Simultaneous CI} = (\bar{Y}_{u.} - \bar{Y}_{v.}) \pm q_{I,(I-1)(J-1)}(\alpha) \cdot \frac{s_p}{\sqrt{J}}}$$

Tukey's method of multiple comparison reveals

$$q_{I,(I-1)(J-1)}(\alpha) \cdot \frac{s_p}{\sqrt{J}} = q_{7,54}(0.05) \cdot \sqrt{\frac{3096}{10}} = 75.8$$

only one significant difference: papaverine vs placebo

Treat	2	1	6	7	4	5	3
Mean	208.4	191.0	176.5	167.2	148.0	144.3	118.2

Friedman's test

Nonparametric test, when ϵ_{ij} are non-normal, to test

H_0 : no treatment effects

Ranking within the block number j

$(R_{1j}, \dots, R_{Ij}) = \text{ranks of } (Y_{1j}, \dots, Y_{Ij})$

$$R_{1j} + \dots + R_{Ij} = \frac{I(I+1)}{2}$$

$$\frac{1}{I}(R_{1j} + \dots + R_{Ij}) = \frac{I+1}{2} \text{ and } \bar{R}_{.j} = \frac{I+1}{2}$$

$$\boxed{\text{Test statistic } Q = \frac{12J}{I(I+1)} \sum_{i=1}^I (\bar{R}_{i.} - \frac{I+1}{2})^2}$$

approximate null distribution $Q \stackrel{a}{\sim} \chi_{I-1}^2$

Q is a measure of agreement between J rankings

reject H_0 for large values of Q

Ex. 3: experiment on itching

R_{ij} and $\bar{R}_{i.}$ are given on p. 470

$$\frac{I+1}{2} = 4, Q = 14.86, \text{ df} = 6, \text{ P-value} \approx 0.0214$$