

ANOVA table (For $H_0: \gamma_1 = \dots = \gamma_t$) (CRD)

(Same as Lecture 4 Slides)

$$SST = SSE_0 - SSE$$

Source	df	Sum of Squares	Mean Squares	F
Treatment	t-1	SST	$mST = \frac{SST}{t-1}$	$F = \frac{mST}{mSE} = \frac{SST/t-1}{SSE/(n-t)}$
Error	n-t	SSE	$mSE = \frac{SSE}{n-t}$	
Total	n-1	SSTotal		Recall $t-1 = \text{rank}(H_S - H_{S0})$ $n-t = \text{rank}(I - I_{\bar{x}})$

Under $H_0: \gamma_1 = \gamma_2 = \dots = \gamma_t$

(Recall model)

$$y_{ij} = \mu + \gamma_i + \epsilon_{ij}$$

central F

$$SST \sim \chi^2_{(t-1, 0)}$$

$$\xrightarrow{\text{indep}} F = \frac{SST/t-1}{SSE/(n-t)} \sim F_{(t-1, n-t)}$$

(balanced design)

$$SSE \sim \chi^2_{(n-t, 0)}$$

$$\lambda = \frac{v}{\sum_{i=1}^t (m_i - \bar{m})^2}$$

In general

$$SST \sim \chi^2_{(t-1, \lambda)}$$

$$SSE \sim \chi^2_{(n-t, 0)}$$

$$\xrightarrow{\text{indep}} F = \frac{SST/(t-1)}{SSE/(n-t)} \sim F_{(t-1, n-t, \lambda)}$$

non central F

non centrality parameter

Determining the number of replicates (in a CRD)

Assume design is balanced!

Idea: Focus on hypothesis testing $H_0: \gamma_1 = \dots = \gamma_t = 0$ vs $H_a: \gamma_i \neq 0$ for some i

We have test statistic $F = \frac{\text{SST}_{t-1}}{\text{SSE}_{(n-t)}}$

For a given α we reject H_0 if $F_{\text{obs}} > F_c$ where F_c = critical value

such that $P(F > F_c | H_0 \text{ true}) = \alpha$ and we know $F \sim F_{(t-1, n-t, \alpha)}$ central F distribution
(in R: 'qt' function) (under H_0)

Recall:

Type I Error: Reject H_0 when H_0 true!

Prob (type I Error) = α (controlled)

Type II Error: Not Reject H_0 when H_0 false!

Since test statistic is fixed then r, σ^2 and

$P(\text{type II Error})$ depends on r and $\gamma_1, \dots, \gamma_t$ (since null hypothesis is not true!)

So we know that when H_0 is not true (H_0^c)

$$\Rightarrow F \sim F_{(t-1, n-t, \lambda)} \xrightarrow{\text{non-central } F} \text{ where } \lambda = \frac{r}{\sigma^2} \sum_{i=1}^t (\gamma_i - \bar{\gamma})^2, \quad \lambda > 0$$

$$\text{where } \lambda = \frac{\underline{\mu}^T (\mathbf{I} - H_{\perp}) \underline{\mu}}{\sigma^2} \text{ with } \underline{\mu} = \mathbf{x}\beta, \quad \beta = \begin{pmatrix} \gamma_1 \\ \vdots \\ \gamma_t \end{pmatrix}$$

$$\text{Recall that } \mu_i = \mu + \gamma_i, i=1, 2, \dots, t$$

$$\Rightarrow \underline{\mu} = \begin{pmatrix} \mu_1 \\ \mu_1 \\ \mu_2 \\ \mu_2 \\ \vdots \\ \mu_t \\ \mu_t \end{pmatrix}, \quad H_{\perp} = \frac{1}{n} \begin{pmatrix} 1 & 1 & \cdots & 1 \\ 1 & 1 & \cdots & 1 \\ 1 & 1 & \cdots & 1 \\ \vdots & \vdots & \ddots & \vdots \\ 1 & 1 & \cdots & 1 \end{pmatrix} \Rightarrow H_{\perp} \underline{\mu} = \bar{\mu} \underline{1}$$

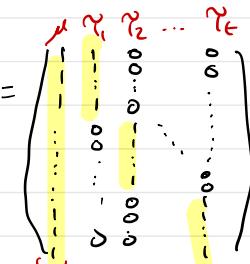
$\underline{1}^T \underline{1} = t$

and $\bar{\mu} = \frac{\mu_1 + \mu_2 + \dots + \mu_t}{t} = \frac{\mu + \gamma_1 + \gamma_2 + \dots + \gamma_t}{t} = \frac{\mu + \sum_{i=1}^t \gamma_i}{t} = \frac{\mu + \bar{\gamma}}{t}$

$$\Rightarrow \underline{\mu}^T \underline{\mu} = r\mu^2 + r\mu_1^2 + \dots + r\mu_t^2 = r \sum_{i=1}^t \mu_i^2 = \frac{\mu + \gamma_1 + \gamma_2 + \dots + \gamma_t}{t} \cdot \frac{\mu + \gamma_1 + \gamma_2 + \dots + \gamma_t}{t} = \frac{(\mu + \gamma_1 + \gamma_2 + \dots + \gamma_t)^2}{t^2} = \mu^2 + \frac{\sum_{i=1}^t \gamma_i^2}{t} = \mu^2 + \bar{\gamma}^2$$

$$\Rightarrow \underline{\mu}^T (\mathbf{I} - H_{\perp}) \underline{\mu} = \underline{\mu}^T \underline{\mu} - \underline{\mu}^T H_{\perp} \underline{\mu} = \underline{\mu}^T \underline{\mu} - n\bar{\mu} \underline{\mu}^T \underline{1} = \underline{\mu}^T \underline{\mu} - n\bar{\mu}^2 = r \sum_{i=1}^t \mu_i^2 - r t \bar{\mu}^2 = r \left(\sum_{i=1}^t \mu_i^2 - t \bar{\mu}^2 \right) = r \sum_{i=1}^t (\mu_i - \bar{\mu})^2 = r \sum_{i=1}^t (\mu + \gamma_i - \mu - \bar{\gamma})^2 = r \sum_{i=1}^t (\gamma_i - \bar{\gamma})^2$$

$$\Rightarrow \lambda = r \cdot \frac{r}{\sigma^2} \sum_{i=1}^t (\gamma_i - \bar{\gamma})^2 = r \cdot \frac{r}{\sigma^2} \sum_{i=1}^t (\mu_i - \bar{\mu})^2$$



$$\text{Prob}(\text{type II Error}) = P(\text{not Reject } H_0 | H_0^c) = P(F \leq F_c | H_0^c)$$

we would like this to be large!

$$\text{Power}(r, \sigma^2, \Sigma) = 1 - P(\text{type II Error}) = 1 - P(F \leq F_c | H_0^c) = P(F > F_c | H_0^c) = P(\text{Reject } H_0 | H_0 \text{ not true})$$

upper tail probability of
a non-central F distribution
in Lab 6 we showed it
was an increasing function
of λ .

MAIN IDEA

- ④ Fix σ^2 and Σ
- ④ Target $\boxed{\text{Power}(r, \sigma^2, \Sigma) = P_0}$ (0.9 say)

④ Determines r such that \star is true

- Notes
- ④ σ^2 is determined by experience or a pilot experiment.
 - ④ Σ will be fixed in a worst-case scenario (next page)

The hardest situation to detect is when:

- (A) The effects of two of the factor levels (say first and last) differ by Δ
- (B) All other effects are equal and midway

e.g.

(A)

$$\mu + \gamma_1 = k + \Delta/2$$

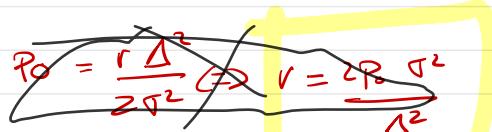
$$\mu + \gamma_t = k - \Delta/2$$

$$\mu + \gamma_2 = \mu + \gamma_3 = \dots = \mu + \gamma_{t-1} = k$$

$\frac{k-\Delta}{2}$ for some constant k (constant)

(B)

$$\Rightarrow \lambda = \frac{1}{\sigma^2} \sum_{i=1}^t (\mu_i - \bar{\mu})^2 = \frac{r \Delta^2}{2 \sigma^2} \Rightarrow P_0 = \frac{r \Delta^2}{2 \sigma^2}$$



$$P_0 = \text{power}^{\alpha} \left(r \sigma^2, \frac{r \Delta^2}{2 \sigma^2} \right)$$

(Same as Lecture ~~4~~ Slides)

ANOVA Factorial design

$$y_{ijk} = \mu + \gamma_i + \alpha_j + \delta_{ij} + \epsilon_{ijk}$$

$$= (1 \bar{X}_1, \bar{X}_2, \bar{X}_{\text{int}})$$

$$SSE = \underline{y^T(I-H_x)y}$$
 stays the same! Just different \bar{X} and \Rightarrow different H_x

Test null hypothesis $H_0: \delta_{ij} = 0 \quad \forall i, j$ (no interactions)

$$\Rightarrow \bar{X}_0 = \left(1 \underbrace{\bar{X}_1}_{\substack{\text{Effect of } \gamma_i \\ \gamma_i}}, \underbrace{\bar{X}_2}_{\substack{\text{Effect of } \alpha_j \\ \alpha_j}} \right) \Rightarrow SSE_0 = \underline{y^T(I-H_{\bar{X}_0})y}$$
 since clearly $H_{\bar{X}_0}$ is also (dominant $\Rightarrow I - H_{\bar{X}_0}$ dominant too!)

$$\text{Also } SSE_0 - SSE > 0$$

$$\Rightarrow \boxed{SSE_0 - SSE} = \underline{y^T(H_x - H_{\bar{X}_0})y}$$
 $\text{rank}(H_x - H_{\bar{X}_0}) = \text{trace}(H_x - H_{\bar{X}_0})$

((sum of squares due to the interactions

SSint

$$= \text{trace}(H_x) - \text{trace}(H_{\bar{X}_0}) = \text{rank}(x) - \text{rank}(x_0)$$

$$= (t-1) + (s-1) + (t-1)(s-1) - (1 + (t-1) + (s-1))$$

$$= (t-1)(s-1)$$

(Same as Lecture 4 slides)

Theorem \Rightarrow

$$\frac{\underline{q}^T (Hx - Hx_0) \underline{q}}{s^2} \sim \chi^2_{((t-1)(s-1), \lambda)}$$

where $\lambda = \sum \underline{\beta}_j^T (Hx - Hx_0) \underline{\beta}_j$

under $H_0: \underline{\beta}_{ij} = 0$

$$\Rightarrow \underline{\beta}_0 = \underline{\beta}_0 \underline{\beta}_0$$

$$\Rightarrow \lambda = \sum \underline{\beta}_0^T \underline{\beta}_0^T (Hx - Hx_0) \underline{\beta}_0 \underline{\beta}_0$$

$$= \sum \underline{\beta}_0 \underline{\beta}_0^T \left(\underline{H_x} \underline{\beta}_0 - \underline{H_{\beta_0}} \underline{\beta}_0 \right) \underline{\beta}_0$$

$$\Rightarrow \lambda = 0$$

↳ since $\underline{\beta}_0$ is a subset of $\underline{\beta}$

otherwise

$$\lambda = \sum \underline{\beta}^T \underline{\beta}^T (Hx \underline{\beta} - Hx_0 \underline{\beta}) \underline{\beta}$$

$$= \sum \underline{\beta}^T \underline{\beta}^T (\underline{x} - \underline{H_{\beta_0}} \underline{x}) \underline{\beta}$$

$$= \sum \underline{\beta}^T \underline{\beta}^T (I - \underline{H_{\beta_0}}) \underline{x} \underline{\beta}$$

notes:
 $\underline{H_{\beta_0}} \underline{x} \neq \underline{x}$
or
 $\neq \underline{\beta_0}$

ANOVA Table for factorial $H_0: \gamma_{ij} = 0 \forall ij$ (Same as Lecture 4 Slides)

Balanced design

Source	df	Sum of Squares	Mean Squares	F
Interaction	$(t-1)(s-1)$	SS_{Int}	$SS_{Int}/(t-1)(s-1)$	$\frac{SS_{Int}/(t-1)(s-1)}{SSE/(st(v-1))}$
Error	$st(v-1)$	SS_E	$SSE/st(v-1)$	

Under H_0 :

$$F \sim F_{((s-1)(t-1), st(v-1), 0)}$$

In general:

$$F \sim F'_{((s-1)(t-1), st(v-1), \lambda)}$$

$$\begin{aligned} n - t &= str - ((t-1) + (s-1) + (t-1)(s-1)) \\ &\cancel{+ s-1} + t.s - \cancel{s-1} + t.s \\ &= str - st = st(v-1) \end{aligned}$$

λ in previous page!

Blocking

- ▶ **Objective:** Reduce the variance of the experimental error (σ^2) and increase the power for detecting treatment factor effects so that results **generalise to whole population**
- ▶ Choose the experimental units for a study to be as homogeneous as possible. Sometimes difficult!
- ▶ Heterogeneous experimental units are grouped into homogeneous subgroups before they are randomly assigned to treatment factor levels
- ▶ The act of grouping the experimental units together in homogeneous groups is called blocking.
- ▶ In a **randomized block design**, a group of heterogeneous experimental units is used so that the conclusions can be more general

Examples of blocking

- ▶ Plots of land in agricultural experiments are usually **blocked by proximity** because plots in close proximity normally have similar soil characteristics
- ▶ When experimental units are animals, the **grouping (blocking) of genetically similar animals**, such as littermates, often reduces variability within groups
- ▶ When experimental units are trials, or points in time where treatments will be applied, they are often **blocked by time** since many lurking variables may change over time and trials in close temporal proximity are more alike

Randomized complete block design (RCB) with one treatment factor

- ▶ Treatment factor has t levels
- ▶ b blocks (or subgroups of homogeneous experimental units)
- ▶ Each block contains exactly t experimental units for a total of $t \times b$ experimental units
- ▶ The t experimental units within each block are as similar as possible
- ▶ The groups of experimental units vary enough from block to block to allow general conclusions to be drawn
- ▶ The randomization of experimental units to treatment factor levels is performed within each block.

Comparison between CRD and RCB $t = 3, b = 4$

```
levels<-c("level 1","level 2","level 3")
fac <- levels %>% rep(each = 4) %>% sample(12) %>% factor()
blocks <- factor(rep(c("block 1", "block 2", "block 3", "block 4"), each=3))
CRD <- data.frame( units=1:12,block=blocks,treatmentCRD=fac)
block1 <- sample(levels,3); block2 <- sample(levels,3)
block3 <- sample(levels,3); block4 <- sample(levels,3)
t<-c(block1,block2,block3,block4) %>% factor()
RCB<-data.frame(block = blocks, treatmentRCB = t)
cbind(CRD,RCB)

  units   block treatmentCRD   block treatmentRCB
1      1 block 1       level 1 block 1       level 1
2      2 block 1       level 1 block 1       level 2
3      3 block 1       level 3 block 1       level 3
4      4 block 2       level 3 block 2       level 3
5      5 block 2       level 1 block 2       level 1
6      6 block 2       level 3 block 2       level 2
7      7 block 3       level 3 block 3       level 1
8      8 block 3       level 2 block 3       level 3
9      9 block 3       level 2 block 3       level 2
10    10 block 4       level 2 block 4       level 3
11    11 block 4       level 1 block 4       level 2
12    12 block 4       level 2 block 4       level 1
```

Statistical model

$$y_{ij} = \mu + b_i + \tau_j + \epsilon_{ij}$$

where

- ▶ b_i is the effect of block $i \in \{1, \dots, b\}$
- ▶ τ_1, \dots, τ_t are the treatment effects
- ▶ All $\{\epsilon_{ij}\}$ are $N(0, \sigma^2)$ and independent
- ▶ Note: There is no interaction between block and treatment
- ▶ Only $t \times b$ experimental units, there would be zero degrees of freedom for the error term ssE if a block by treatment interaction term were included!

ANOVA for Randomised complete block design

Model $y_{ij} = \mu + b_i + \gamma_j + \epsilon_{ij}$

\Rightarrow in matrix form

$$\underline{y} = \underline{X}\underline{\beta} + \underline{\epsilon}$$

$$\underline{\beta} = \begin{pmatrix} \mu \\ b_1 \\ \vdots \\ b_b \\ \gamma_1 \\ \vdots \\ \gamma_k \end{pmatrix}$$

$$\underline{X} = \left(\begin{array}{cccccc|ccc} 1 & 1 & 0 & 0 & 0 & 0 & 0 & 1 & 0 & \dots & 0 \\ 1 & 1 & 0 & 0 & 0 & 0 & 0 & 0 & 1 & \dots & 0 \\ \vdots & \dots & \vdots \\ 1 & 0 & 0 & 0 & 0 & 0 & 0 & 0 & 0 & \dots & 0 \\ 1 & 0 & 0 & 0 & 0 & 0 & 0 & 0 & 0 & \dots & 0 \\ \vdots & \dots & \vdots \\ 1 & 0 & 1 & 0 & 0 & 0 & 0 & 0 & 0 & \dots & 0 \\ 1 & 0 & 1 & 0 & 0 & 0 & 0 & 0 & 0 & \dots & 0 \\ \vdots & \dots & \vdots \\ 1 & 0 & 0 & 0 & 0 & 0 & 0 & 1 & 0 & \dots & 0 \\ 1 & 0 & 0 & 0 & 0 & 0 & 0 & 0 & 1 & \dots & 0 \\ \vdots & \dots & \vdots \\ 1 & 0 & 0 & 0 & 0 & 0 & 0 & 0 & 0 & \dots & 1 \end{array} \right)$$

1 U V

$H_0:$ $b_i = 0$ vs $H_a:$ $b_i \neq 0$ for some i

Under $H_0:$ $y_{ij} = \mu + \gamma_j + \epsilon_{ij}$

\Rightarrow in matrix form

$$\underline{y} = \underline{X}_{ob} \underline{\alpha} + \underline{\epsilon} \quad \underline{\alpha} = \begin{pmatrix} \mu \\ \gamma_1 \\ \vdots \\ \gamma_k \end{pmatrix}$$

$$\underline{X}_{ob} := (\underline{1}, \nabla), \underline{X}_{op} = (\underline{1}, \nabla)$$

$H_0:$ $\gamma_i = 0$ vs $H_a:$ $\gamma_i \neq 0$ for some i

under H_0

$$y_{ij} = \mu + b_i + \epsilon_{ij}$$

$$\underline{\alpha} = \begin{pmatrix} \mu \\ b_1 \\ \vdots \\ b_b \end{pmatrix}$$

$$\Rightarrow \underline{y} = \underline{X}_{or} \underline{\alpha} + \underline{\epsilon}$$

Now, we can do F tests

$$\begin{aligned} SSE &= \underline{y}^T (\mathbf{I} - H_{\bar{x}}) \underline{y}, \text{ rank}(H_{\bar{x}}) = n - \text{rank}(\bar{x}) \Rightarrow \frac{SSE}{\sigma^2} \sim \chi^2_{((t-1)(b-1), d)} \\ &= t \cdot b - (1 + t - 1 + b - 1) \\ &= t \cdot b - t - b + 1 = t(b-1) - (b-1) \\ &= (t-1)(b-1) \end{aligned}$$

$$SS_{B_{ob}} = SSE_{(H_0^{(b)})} - SSE \quad \text{rank}(H_{\bar{x}} - H_{\bar{x}_{ob}}) = \text{rank}(\bar{x}) - \text{rank}(\bar{x}_{ob}) \Rightarrow \frac{SS_{B_{ob}}}{\sigma^2} \sim \chi^2_{(b-1, \lambda_b)}$$
$$\underline{y}^T (H_{\bar{x}} - H_{\bar{x}_{ob}}) \underline{y} = 1 + t - 1 + b - 1 - (1 + t - 1) = b - 1 \quad \lambda_b = \underline{\mu}^T (\mathbf{I} - H_{\bar{x}_{ob}}) \underline{\mu} / \sigma^2$$

$$\begin{aligned} SST &= SSE_{(H_0^{(r)})} - SSE \\ &= \underline{y}^T (H_{\bar{x}} - H_{\bar{x}_{or}}) \underline{y} \quad \text{rank}(H_{\bar{x}} - H_{\bar{x}_{or}}) = \\ &\quad = 1 + t - 1 - b - 1 - (1 + b - 1) = t - 1 \quad \Rightarrow \frac{SST}{\sigma^2} \sim \chi^2_{(t-1, \lambda_r)} \\ &\quad \lambda_t = \underline{\mu}^T (\mathbf{I} - H_{\bar{x}_{or}}) \underline{\mu} / \sigma^2 \end{aligned}$$

ANOVA Table for RCB

Source	df	Sum of Squares	Mean Squares	F ratio
Blocks	b-1	SS _{Blocks}	$\frac{SS_{\text{Blocks}}}{b-1}$	
Treatments	t-1	SST	$\frac{SST}{t-1}$	$\frac{SST/t-1}{\frac{SSE}{(t-1)(b-1)}}$
Error	(t-1)(b-1)	SSE	$\frac{SSE}{(t-1)(b-1)}$	

$$F = \frac{\frac{SST/t-1}{SSE/(t-1)(b-1)}}{t-1, (t-1)(b-1), \lambda}$$