

Test : - assumptions

- Hypo :

- Test statistic & its null distn.

$\overline{\overline{Y}}$

$\overline{Y_i}$

Y_{ij} ^{ith}
 $\underbrace{Y_{i1}, \dots, Y_{iJ}}$

Var within a group: i : $(Y_{ij} - \overline{Y_i})^2$
 $X_1 \dots X_n \Rightarrow \frac{1}{n-1} \sum_{i=1}^n (X_i - \bar{X})^2$

Var between groups: $\overline{Y_1}, \overline{Y_2}, \dots, \overline{Y_I}$
 $\overline{\overline{Y}}$

^{1th}
 Y_{11}
 Y_{12}
 \vdots
 Y_{1J}

nd
 \vdots

^{Ith}
 Y_{I1}
 \vdots
 Y_{IJ}

$(Y_{ij} - \overline{\overline{Y}})^2 \rightarrow \text{var of whole}$

Anova :

Hypo ?

How to form test statistic

null distn of $F \rightarrow$ slide 19.

search, F distn & def of F-distn.

$\left. \begin{matrix} SS_B \\ SS_W \end{matrix} \right\} \rightarrow F \left| \begin{matrix} \text{slide} \\ 15-18 \end{matrix} \right.$

Anova: test to compare means *

H_0 : all means are equal *

\Rightarrow all α_i are the same, $i = 1, I$.

(I groups.

$i = 1, \dots, I$

$j = 1, \dots, J_i$

$I = 7$

$J = 10$

$$Y_{ij} = \mu + \alpha_i + \underbrace{e_{ij}}_{\text{random error} \sim N(0; \sigma^2)}$$

group 1: $\mu + \alpha_1$

— 2: $\mu + \alpha_2$

\vdots
 I : $\mu + \alpha_I$

α_i same

$$\sum_{i=1}^I \alpha_i = 0$$

Test statistic \Rightarrow variance analysis:

within group var $\leftarrow SS_W$

between group var $\leftarrow SS_B$

$\Rightarrow F = \text{formular by } SS_B \text{ \& } SS_W$

F-distrib ? 2 df: df_1 & df_2 .

$\chi^2_{df_1}$ has $df = df_1$

$\chi^2_{df_2}$ has $df = df_2$

are indep.

$$t_{df} \sim Z$$

$\frac{\chi^2_{df_1} / df_1}{\chi^2_{df_2} / df_2}$ has F distrib with df_1 and df_2 .

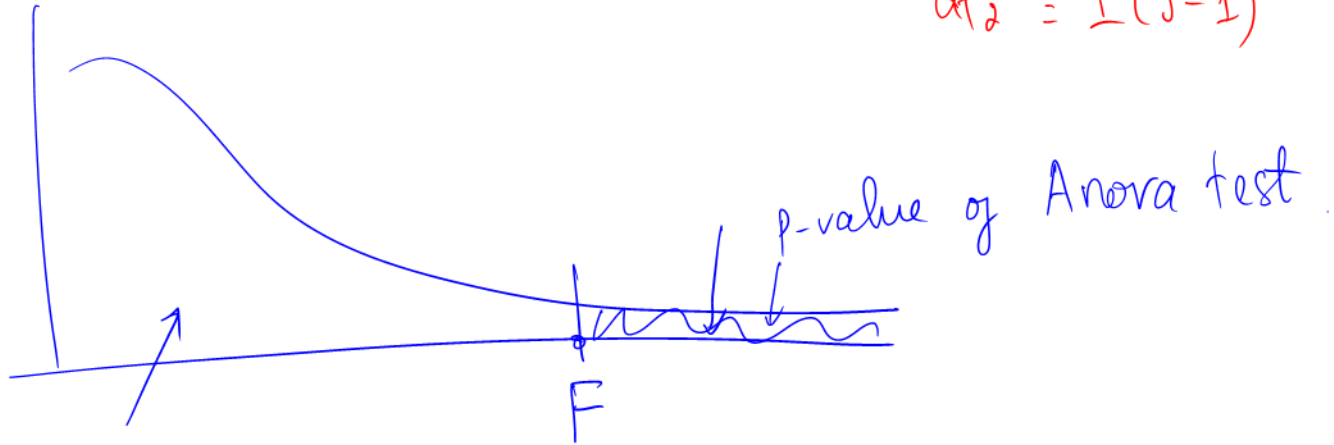
Using the definition of F distn, we can get the distn of the test statistic

$$F = \frac{SS_B / (I - 1)}{SS_W / [I(J - 1)]}$$

$H_0 \sim F_{df_1, df_2}$ where

$$df_1 = I - 1 \quad \&$$

$$df_2 = I(J - 1)$$



Anova assumptions checking:

- equal var of I groups.

normality of I groups \rightarrow Bar or Levene

normality of residuals, indep hist, QQ plot, Shapiro test.

if each group is not normal but has large size \rightarrow can accept.

21 pairwise tests \Rightarrow prob of having at least 1 type 1 error: 66%

\Rightarrow too large!

\Rightarrow need correct it. to reduce from 66% \downarrow 10% only (for eg)

\Rightarrow instead of controlling $\alpha = 0.05$ (5%) for each test, we'll control $\alpha = 0.1$ for all 21 tests. \rightarrow goal.

\Rightarrow we are using "correction"! \Rightarrow Bonferroni \leftarrow
 \Rightarrow Tukey \leftarrow
 \Rightarrow Dunnett \leftarrow

For Bonf: have total k pairwise tests ($k = 21$) \Rightarrow we allow

$\alpha = \underline{0.1}$ type 1 error for all k tests, then

for each pairwise test, we use a sig. level $\alpha = \frac{\alpha}{k}$.

$$\alpha = \frac{0.1}{21}$$

\Rightarrow this idea works quite well when k is small.

For Tukey correction:

for each pair of 2 groups, they form a CI for the difference of the means.

$$\begin{cases} (\mu_1 - \mu_2) \\ (\mu_1 - \mu_3) \\ \vdots \\ \mu_1 - \mu_k \\ \vdots \end{cases}$$

$$\alpha = 0.1$$

not conf. level for each interval but a conf. level of $(1 - \alpha)$ for all the k intervals.

Recal: CI and 2-sided test: 95%

$$H_0: \mu_1 - \mu_2 = 0$$

$$\downarrow$$
$$\text{p-value} < \underline{0.05}$$

get CI for $(\mu_1 - \mu_2)$:

$$(-0.5; -0.1)$$

$$0 \neq \underline{\mu_1 - \mu_2}$$

in general: CI for $\mu_1 - \mu_2$:

$$\downarrow$$
$$(\bar{Y}_1 - \bar{Y}_2) \pm \boxed{} * SE(\bar{Y}_1 - \bar{Y}_2)$$

point estimate \pm margin of error.

For each pairwise comparison test: lab 1 vs lab 2: μ_1 vs μ_2 .

\Rightarrow alpha $\underline{\alpha = 0.05} \Rightarrow$ compare the real p-value vs α .

For the correction (Bonf or Tukey), we do not use α for each test, but we use α (FERate) \Rightarrow compare the adjusted p-value of each test vs α .

For eg: Bonf: $k = 21$ tests

lab1 vs lab2 \Rightarrow p-value: 0.01. \leftarrow

With Bonf correction, $\alpha = 0.1$ then \Rightarrow adjusted p-value vs $\alpha = 0.1$

$$\text{or: } \frac{\text{p-value}}{0.01} \text{ vs } \frac{\alpha}{k} = \frac{0.1}{21}$$

Usually, output of Bonf correction or Tukey will provide adjusted-p-value \Rightarrow vs α (F Error).

Comparing btw Bonf vs Tukey:

if the samples are under Anova \Rightarrow Tukey $>$ Bonf.

if the samples are not under Anova. (under Kruskal Wallis)
then Tukey cannot be used \Rightarrow only use Bonf.